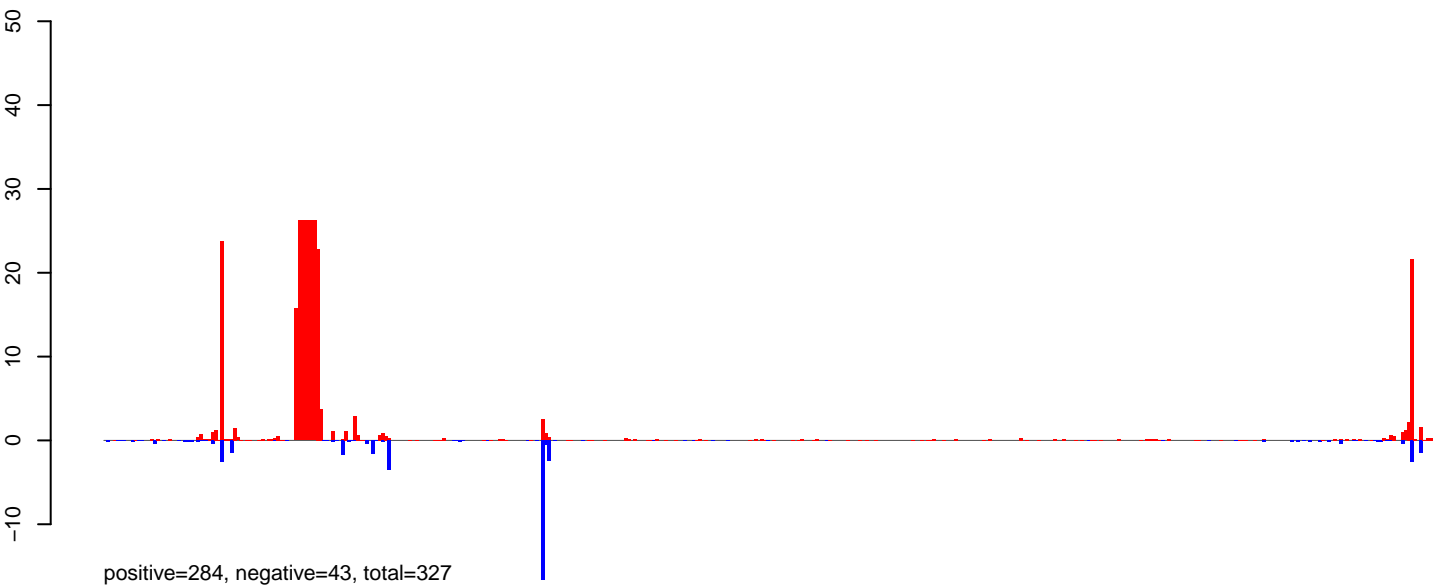
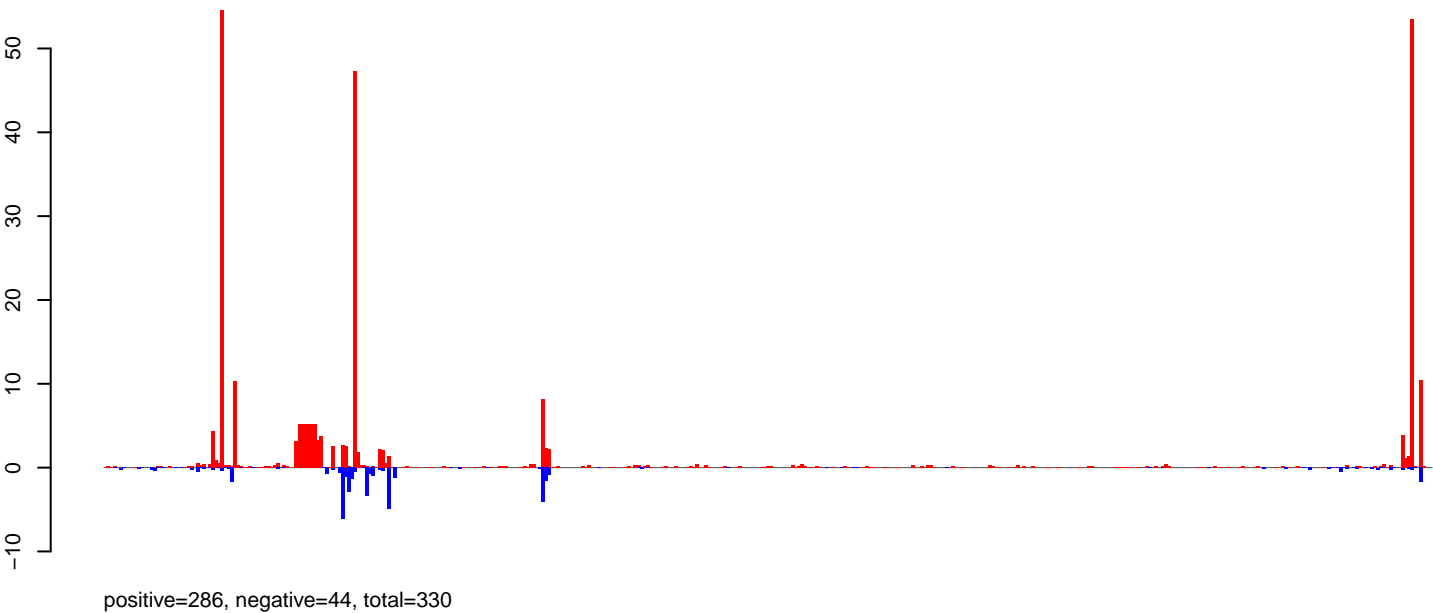


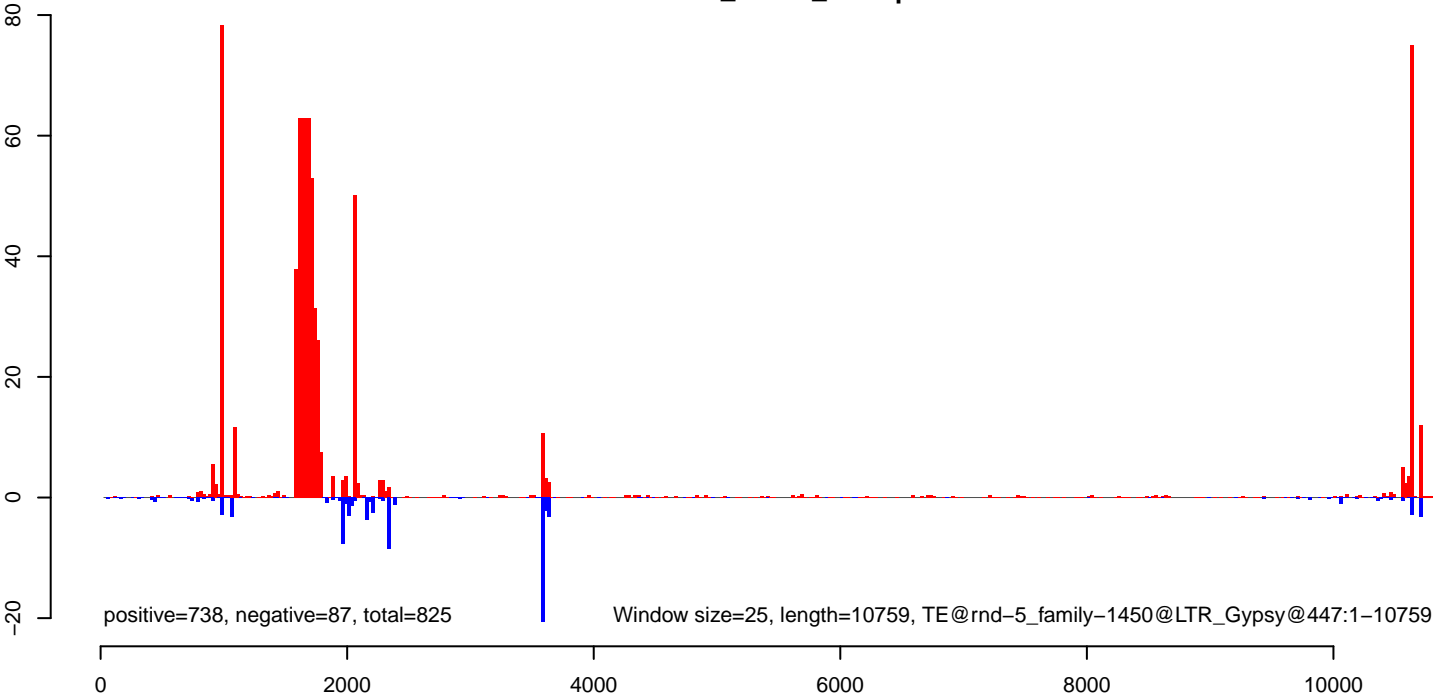
CuQuin_Mcarc_NL.18_23.rep



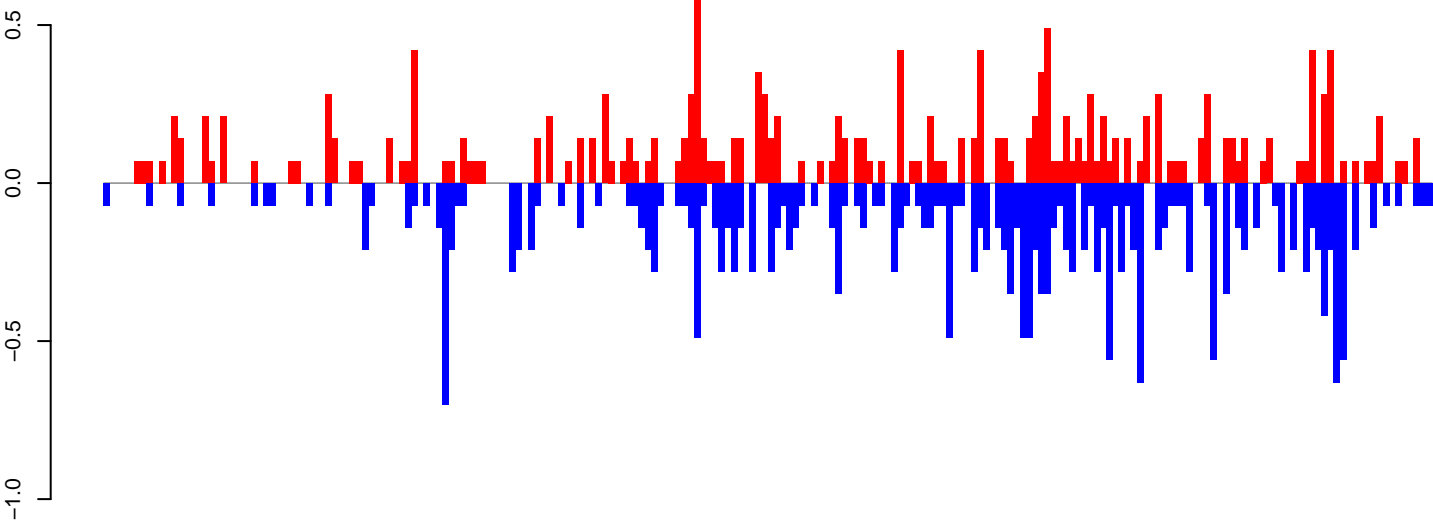
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

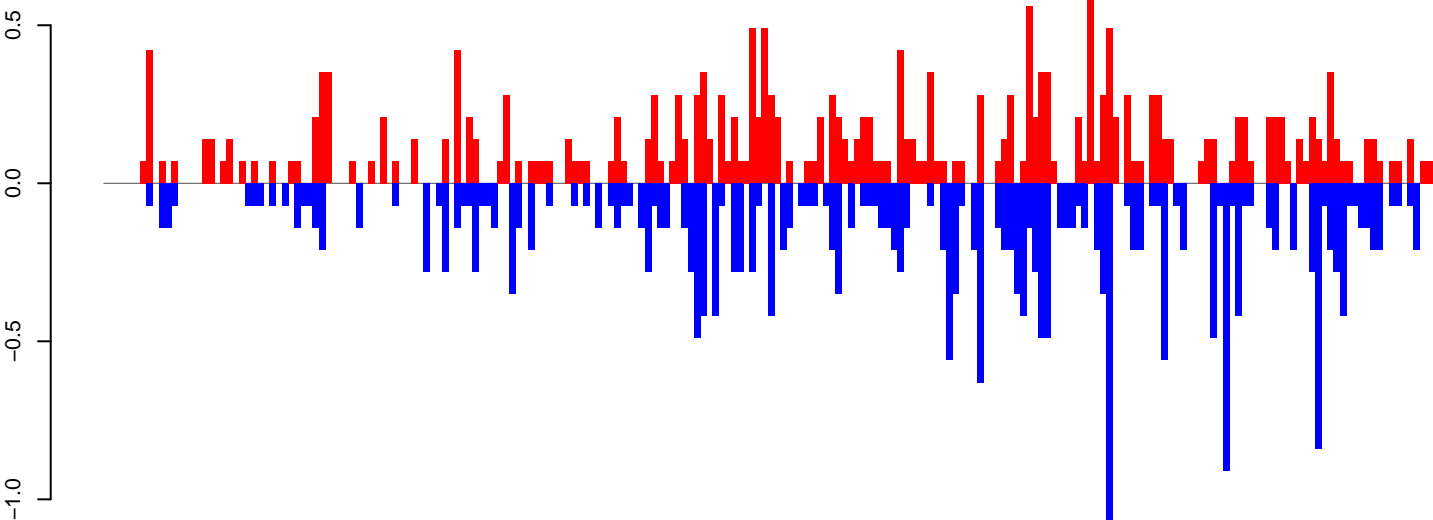


CuQuin_Mcarc_NL.18_23.rep



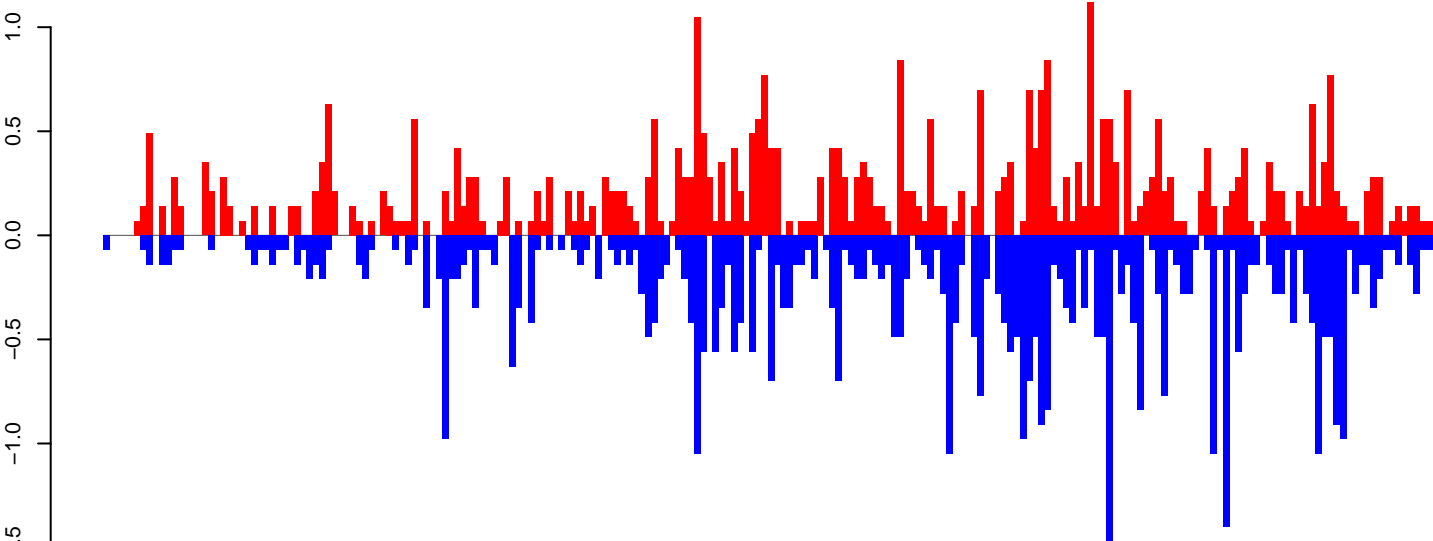
positive=17, negative=22, total=39

CuQuin_Mcarc_NL.24_35.rep



positive=22, negative=26, total=48

CuQuin_Mcarc_NL.rep

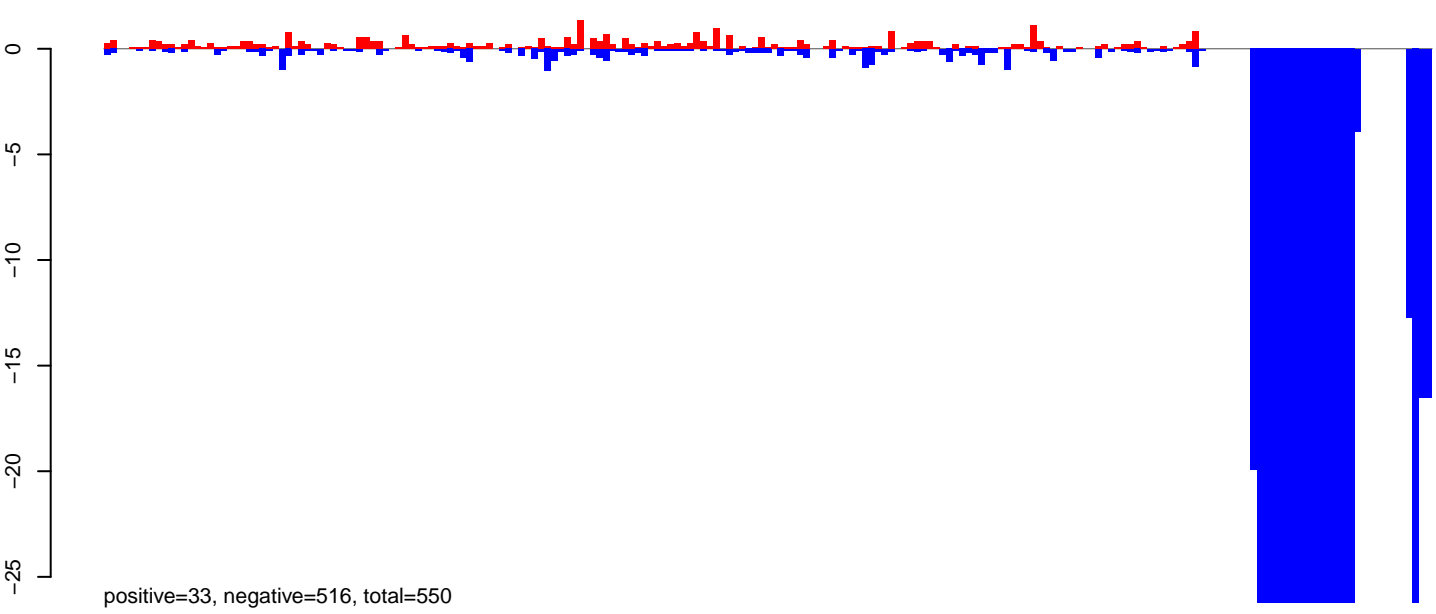


positive=41, negative=50, total=91

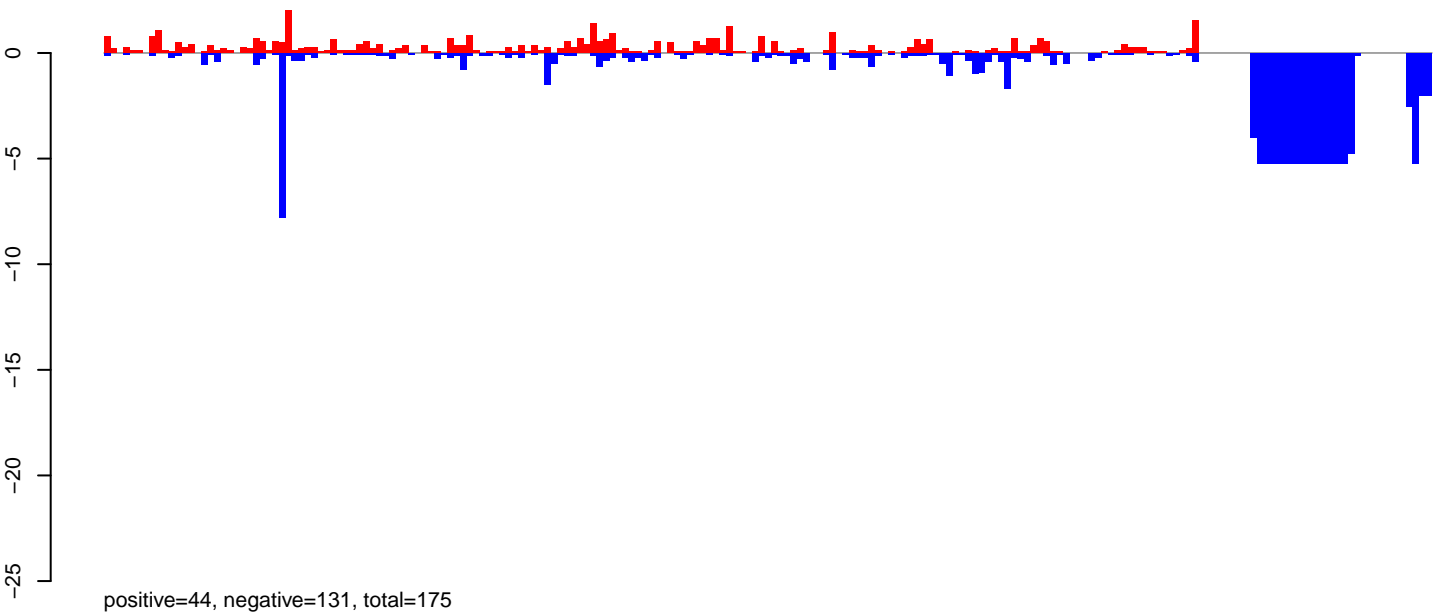
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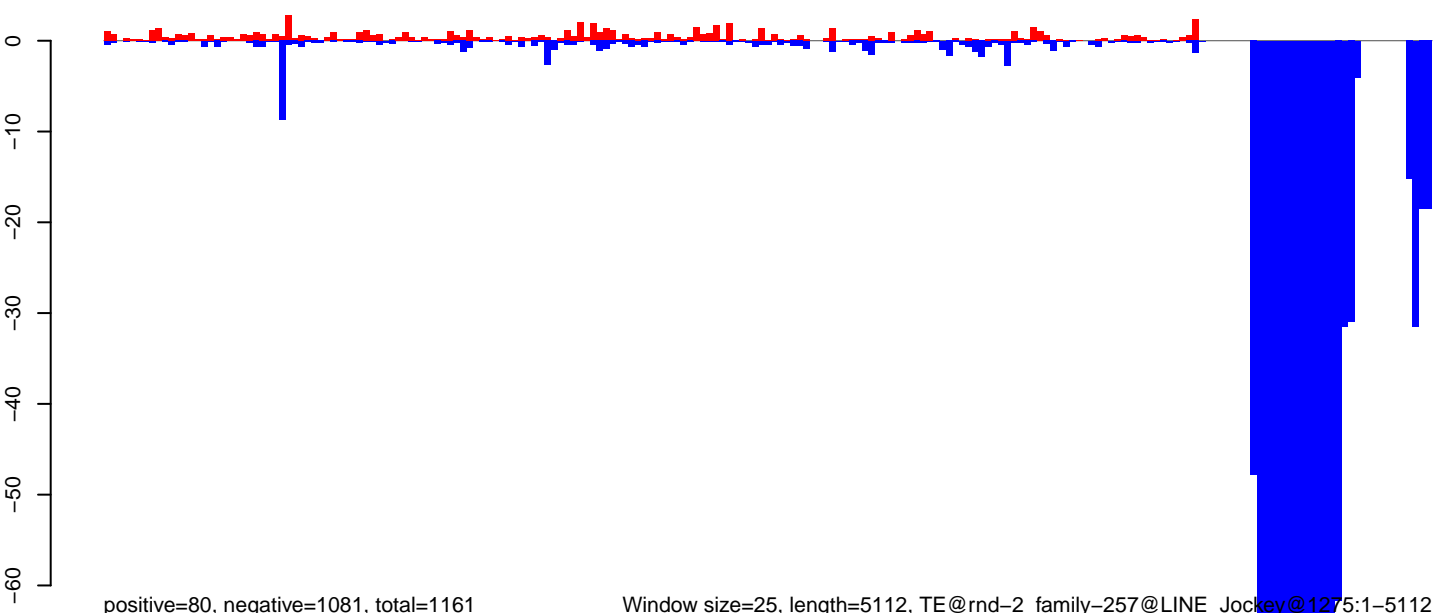
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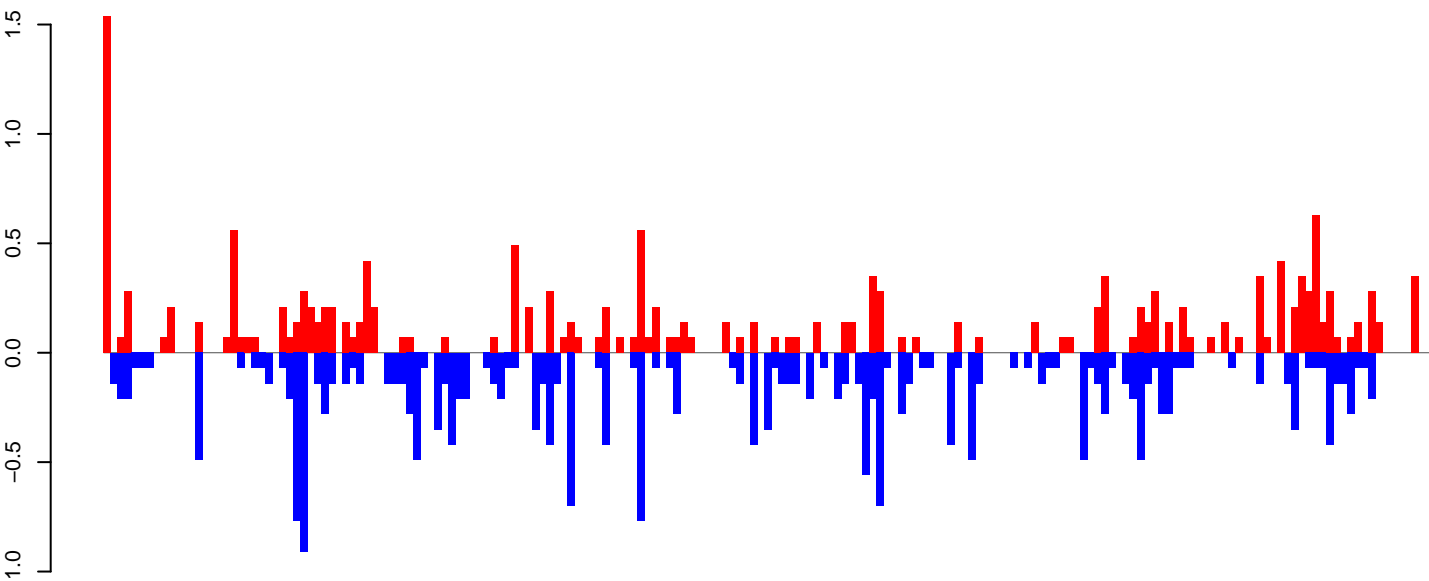
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

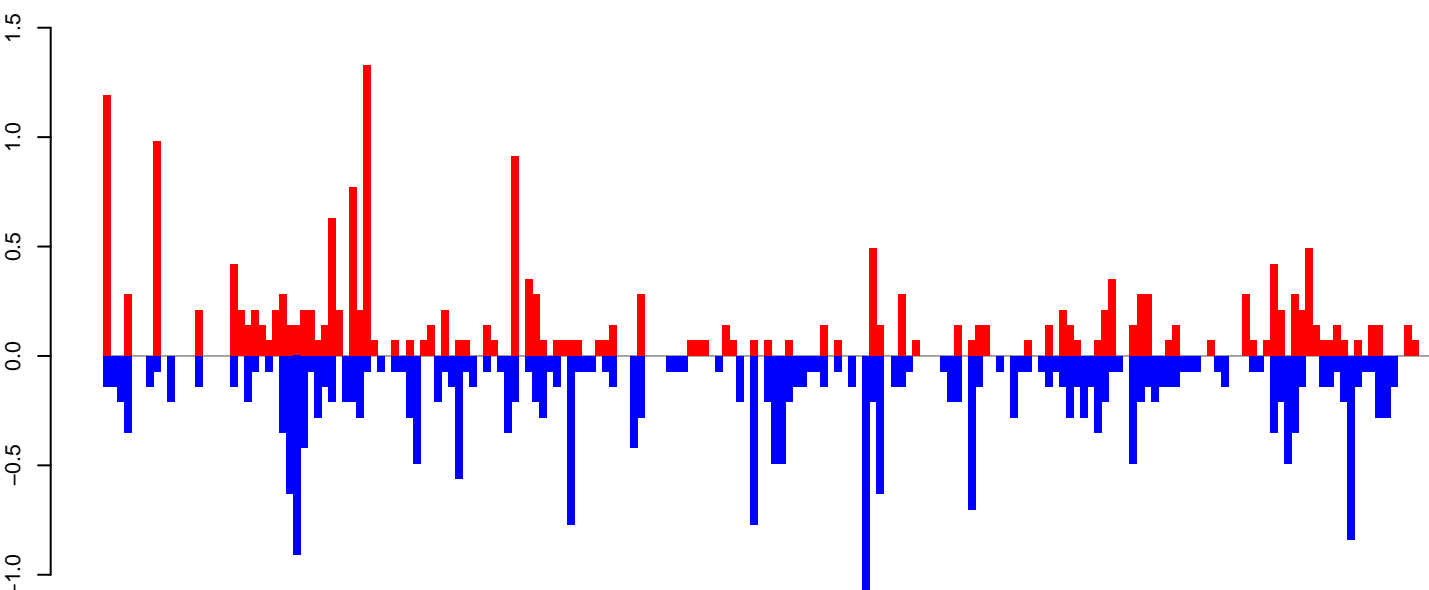


CuQuin_Mcarc_NL.18_23.rep



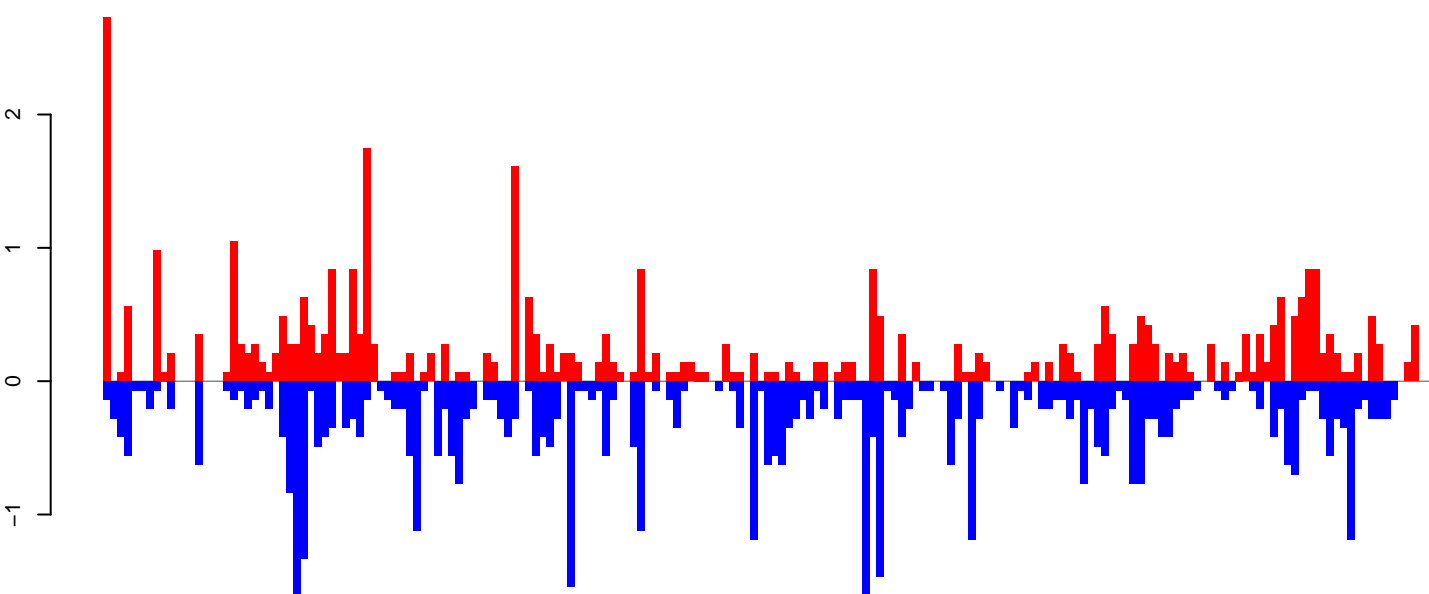
positive=17, negative=22, total=39

CuQuin_Mcarc_NL.24_35.rep



positive=19, negative=27, total=46

CuQuin_Mcarc_NL.rep

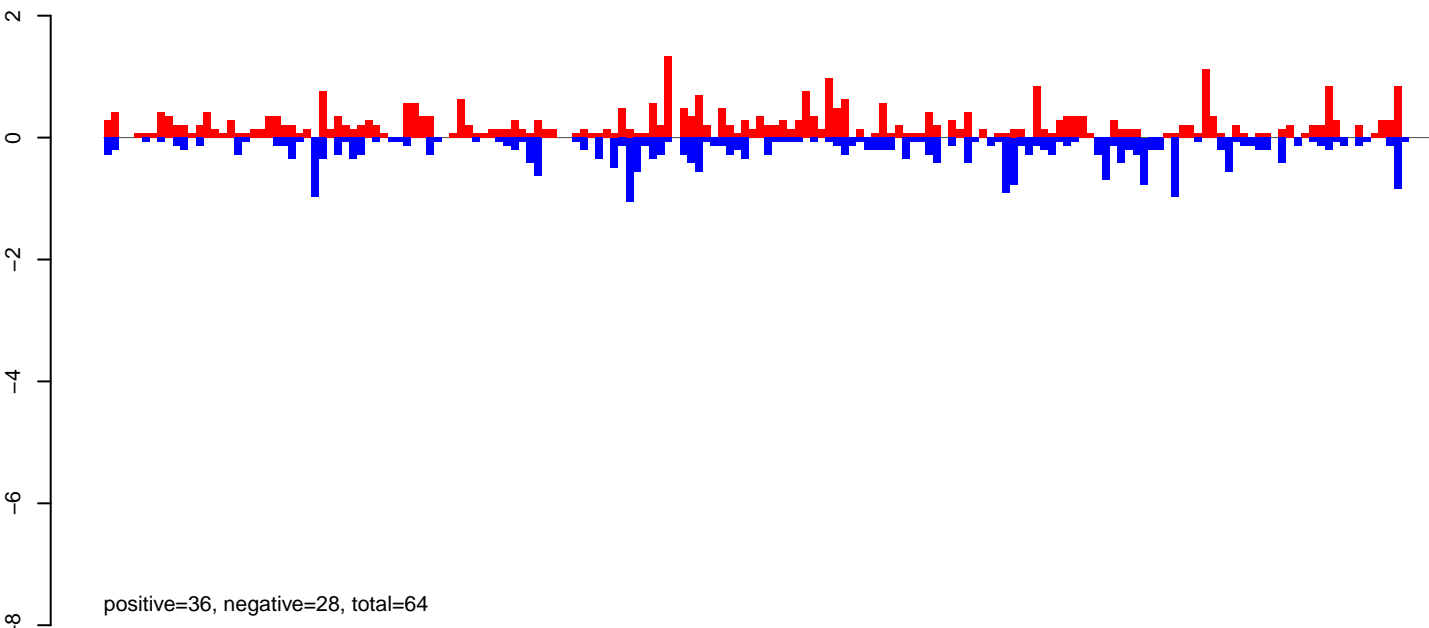


positive=38, negative=50, total=88

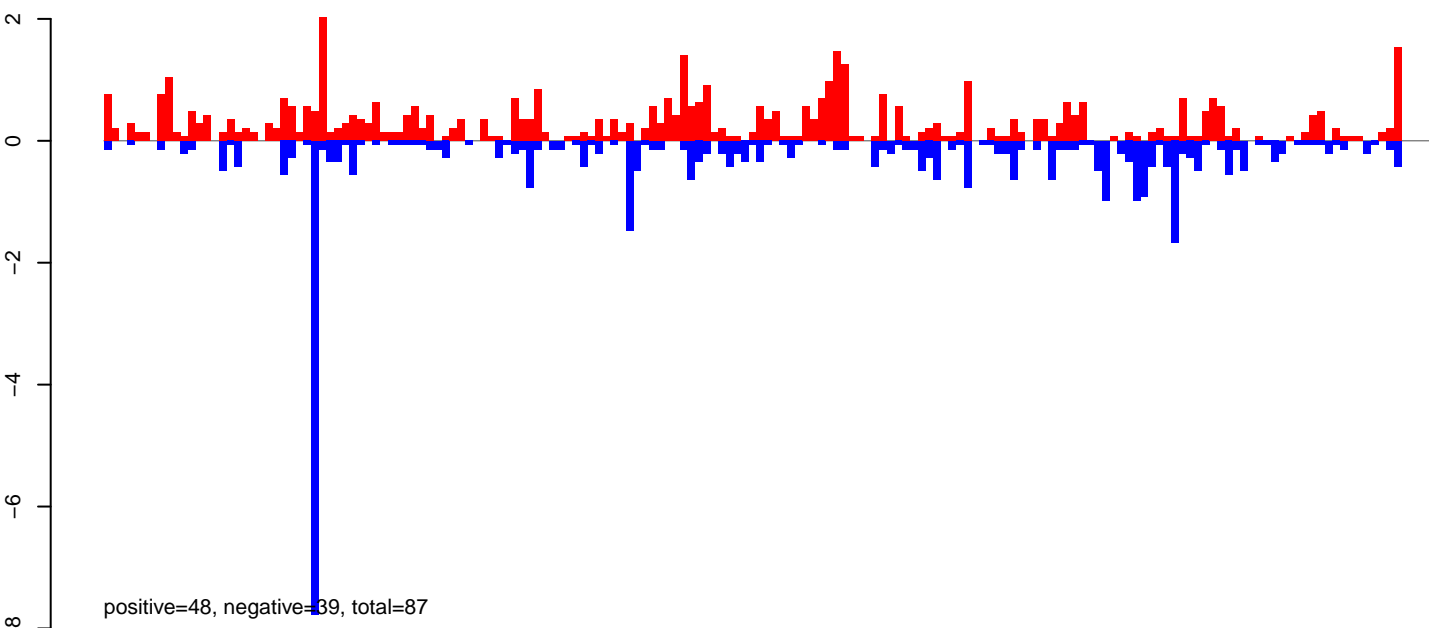
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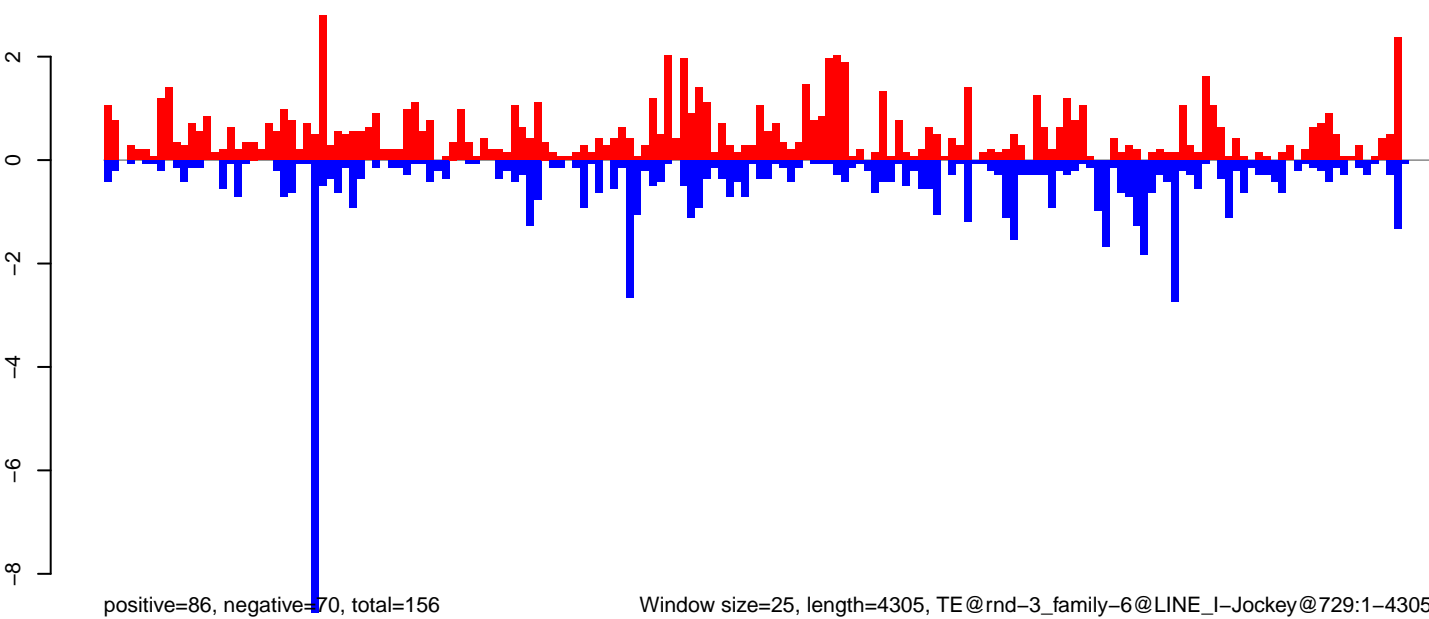
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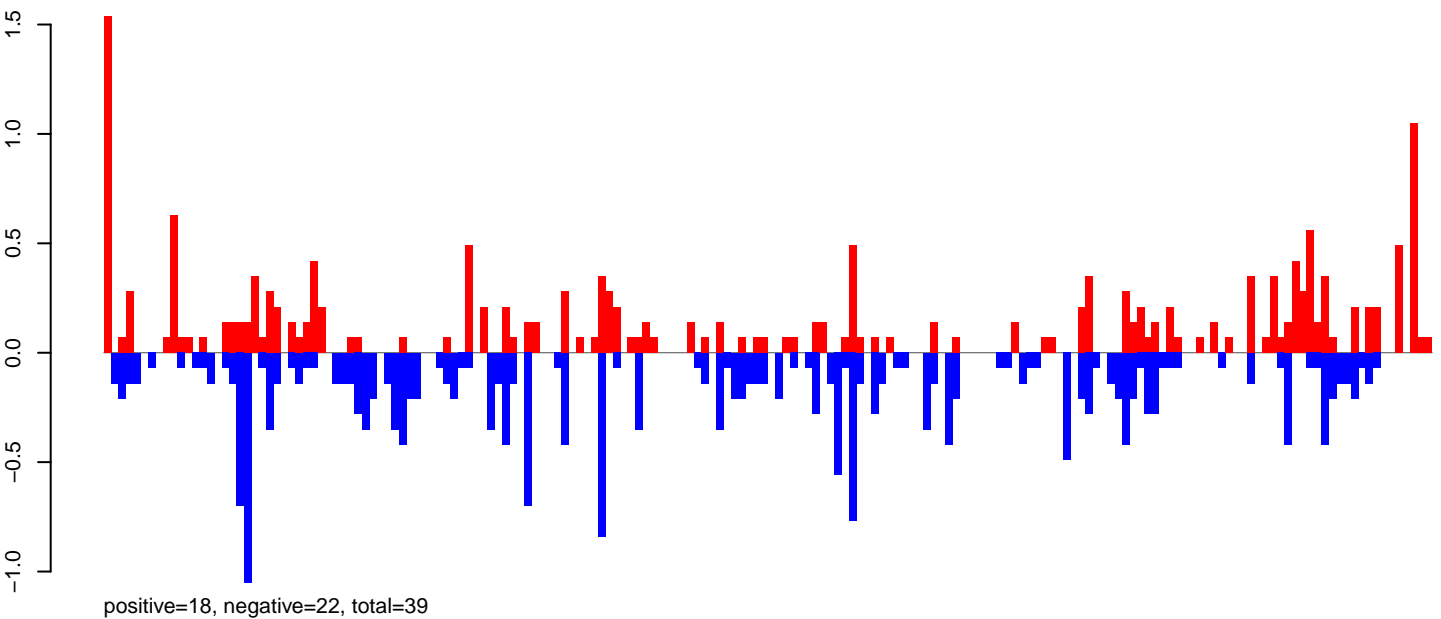
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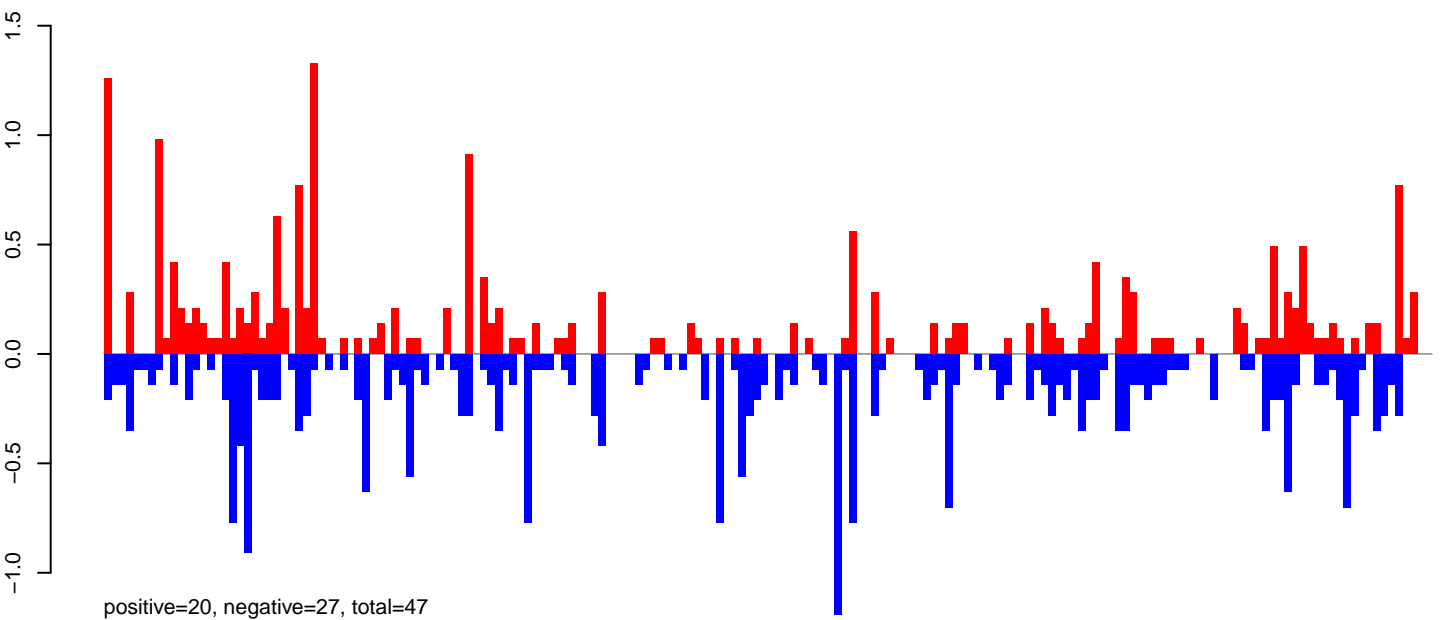
CuQuin_Mcarc_NL.rep



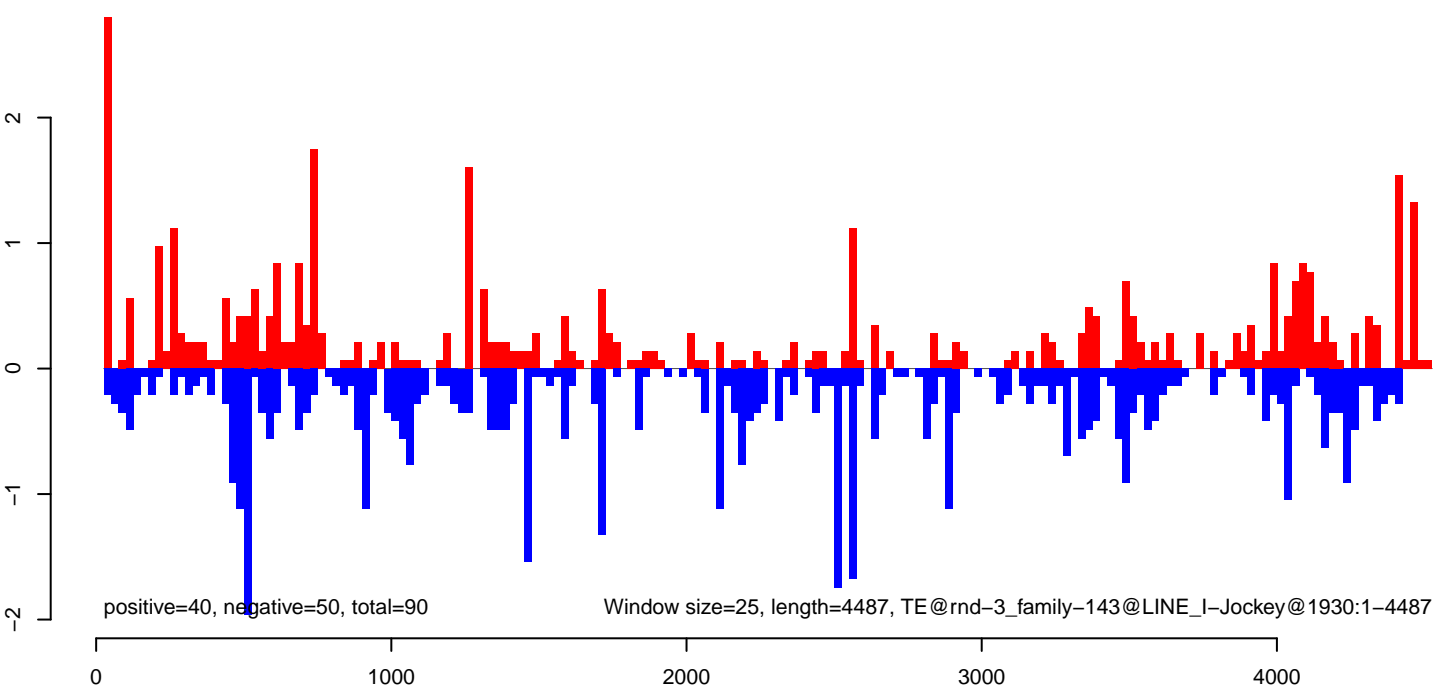
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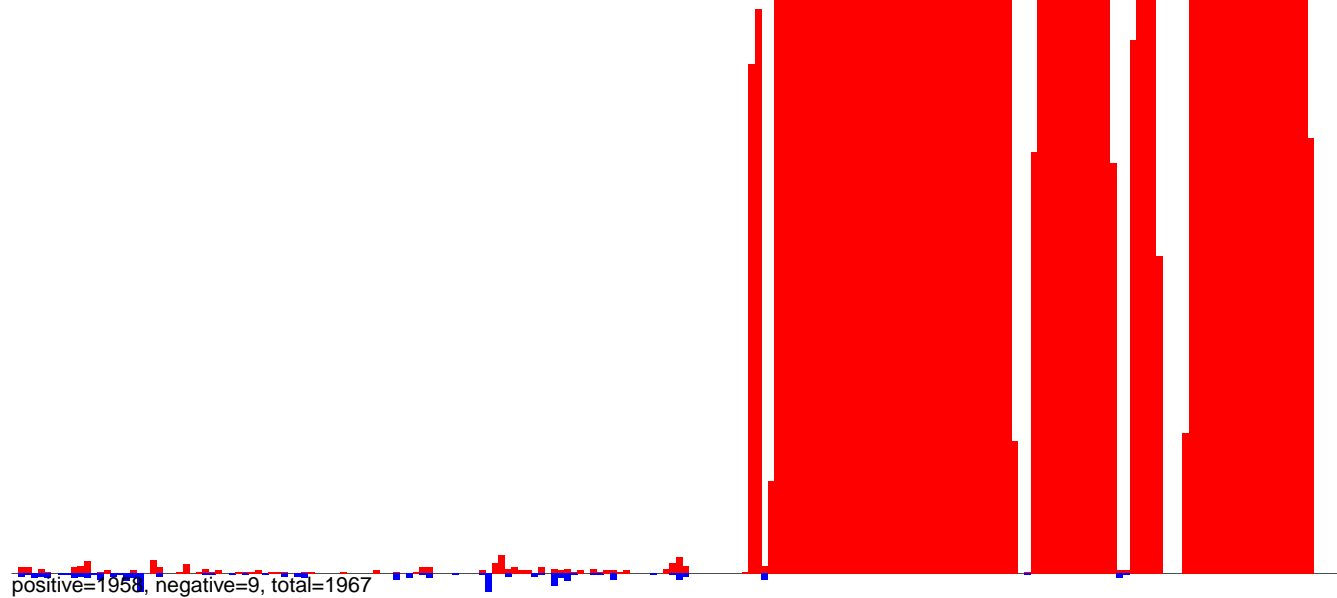
CuQuin_Mcarc_NL.24_35.rep



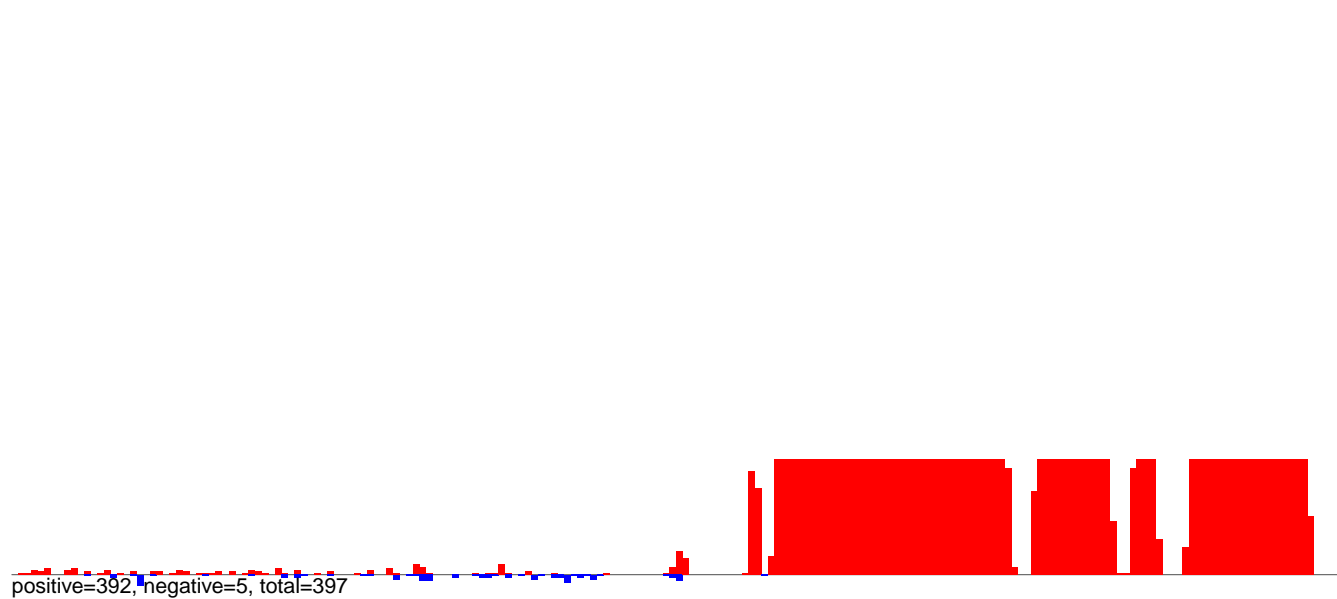
CuQuin_Mcarc_NL.rep



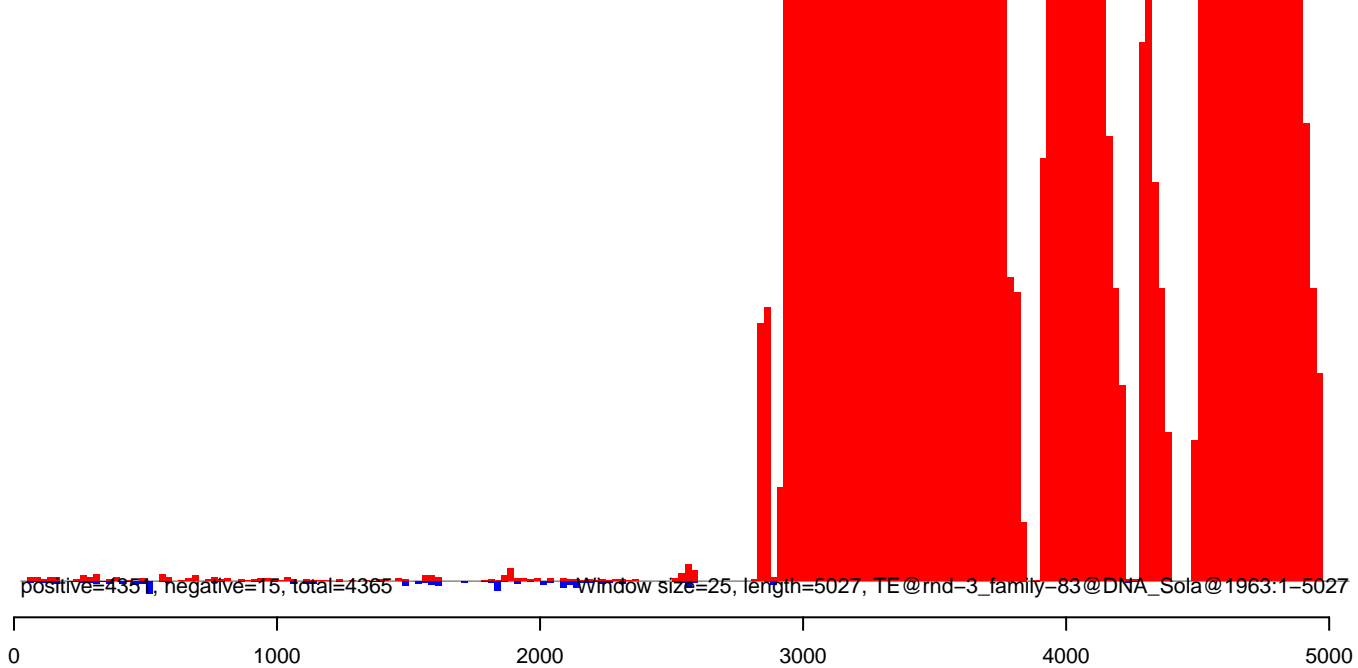
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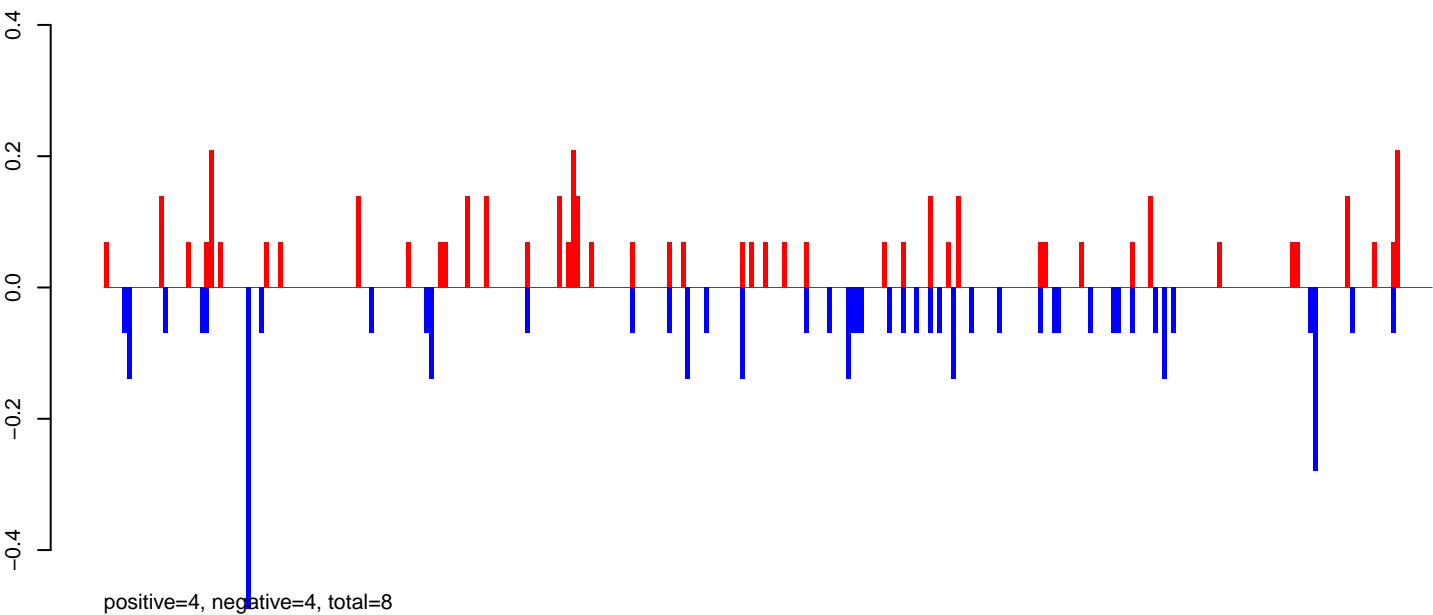
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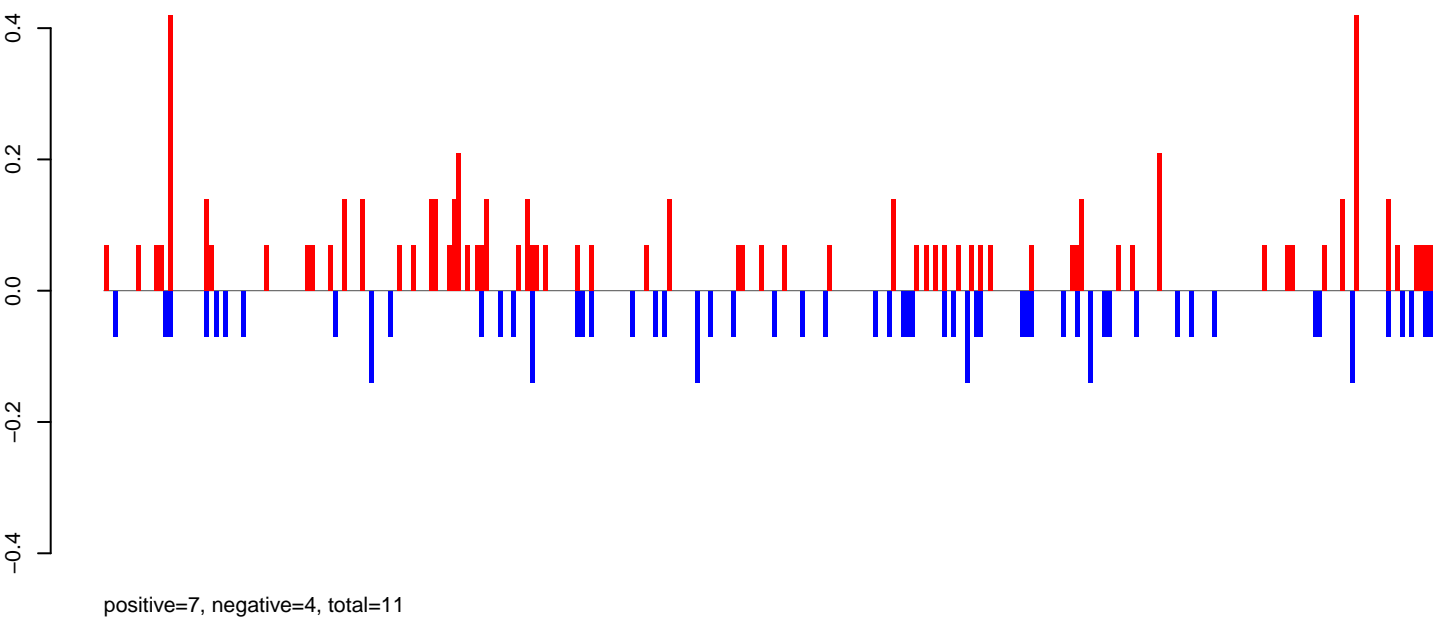
CuQuin_Mcarc_NL.rep



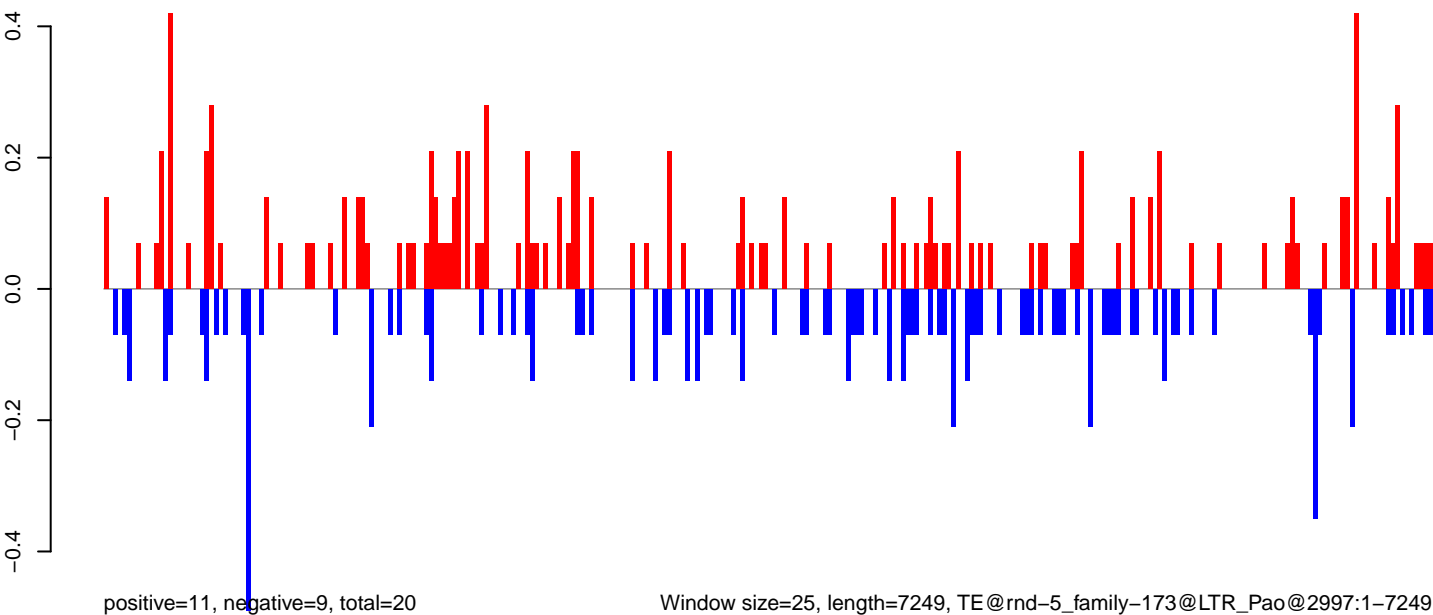
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CuQuin_Mcarc_NL.24_35.rep



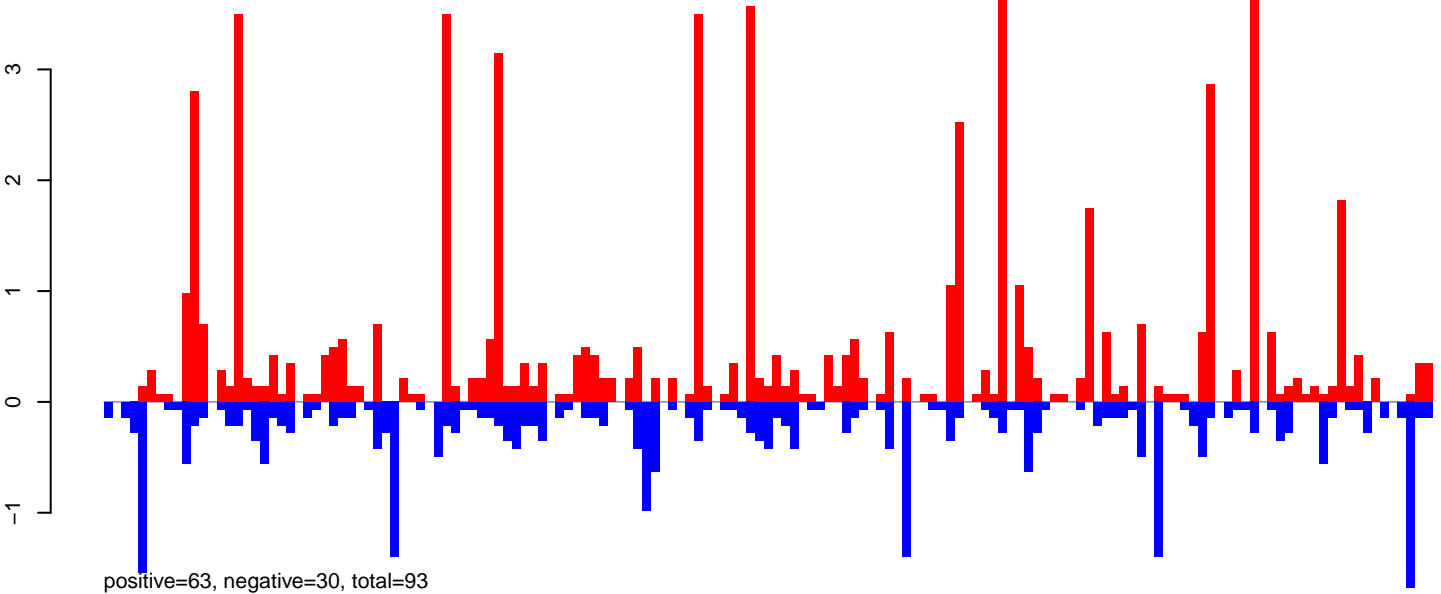
CuQuin_Mcarc_NL.rep



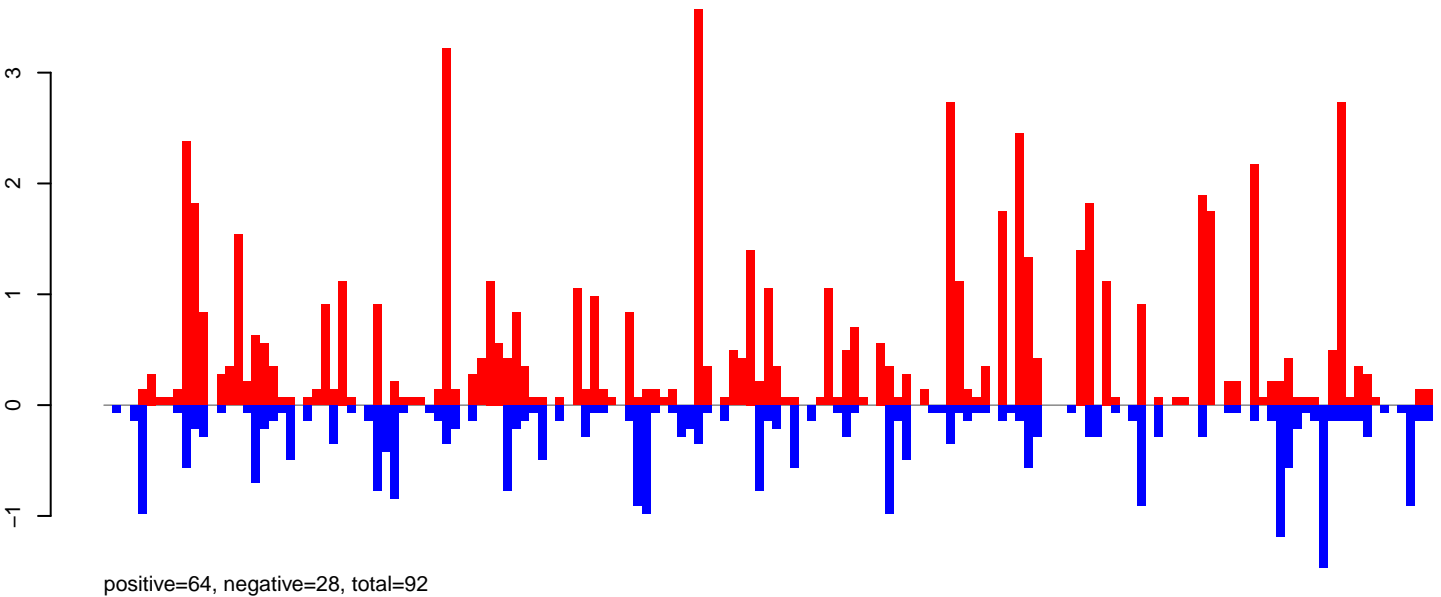
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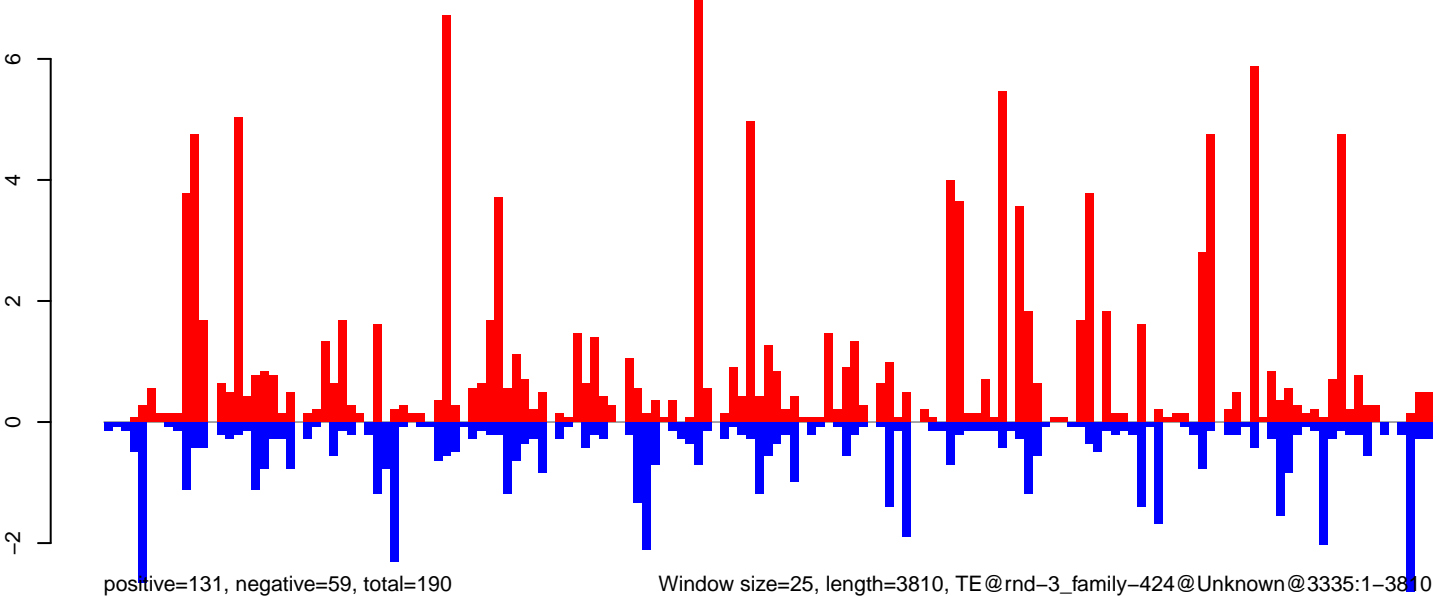
CuQuin_Mcarc_NL.18_23.rep



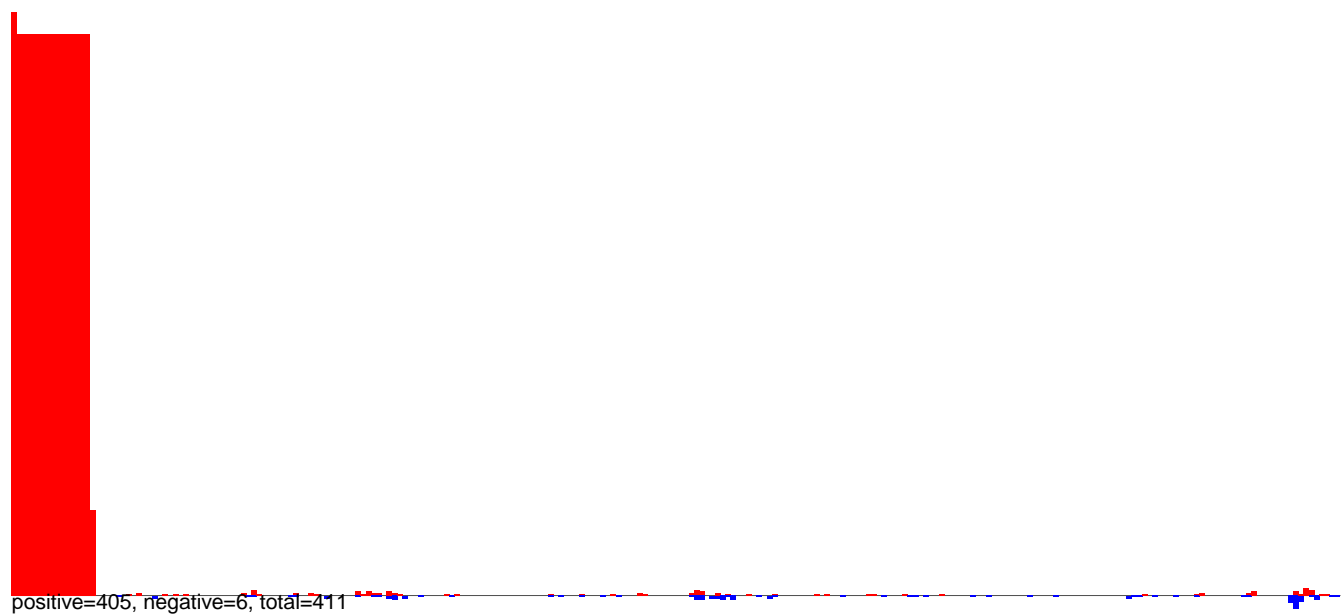
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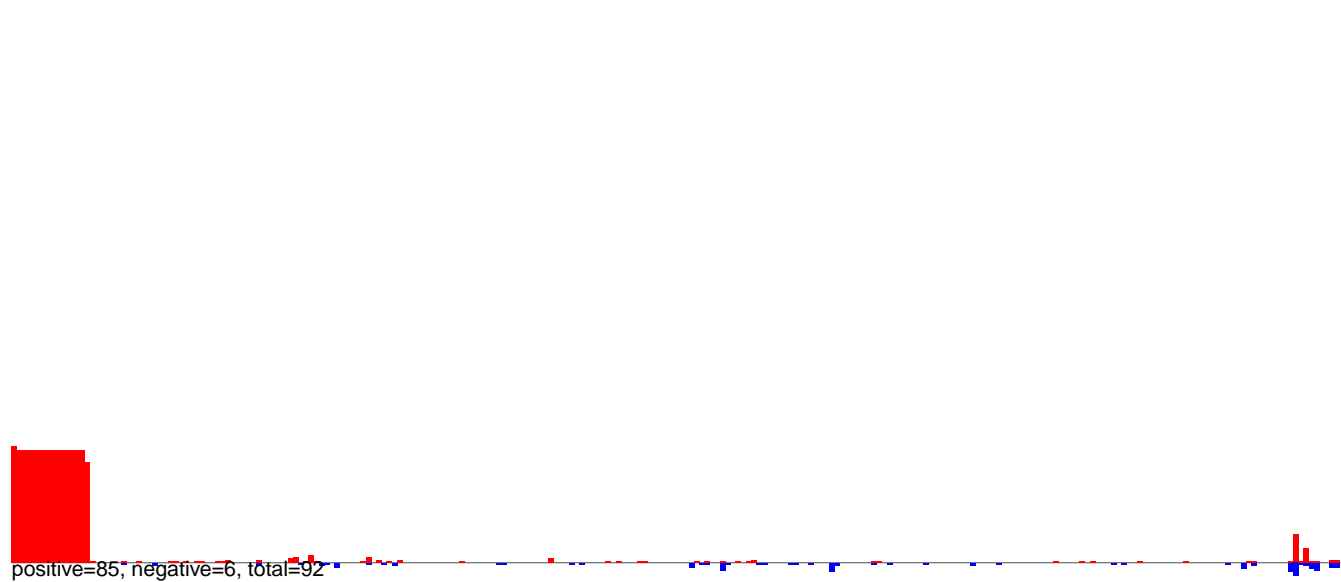
CuQuin_Mcarc_NL.rep



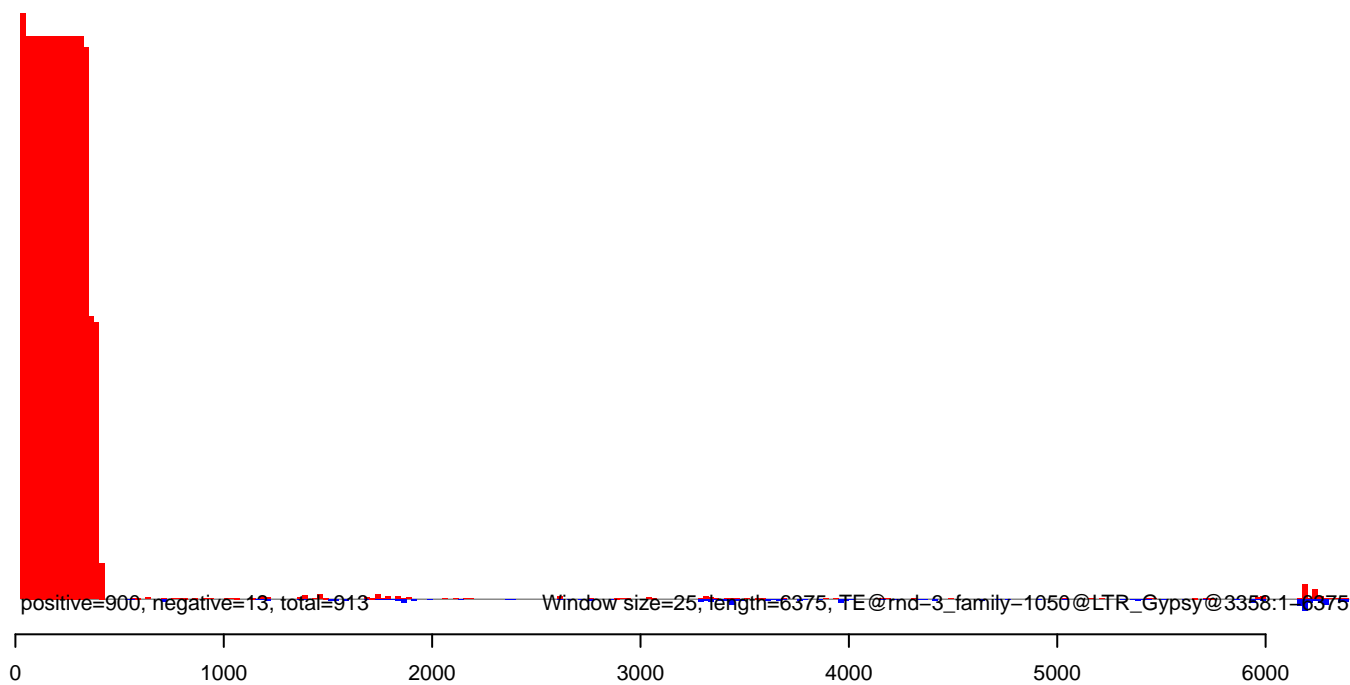
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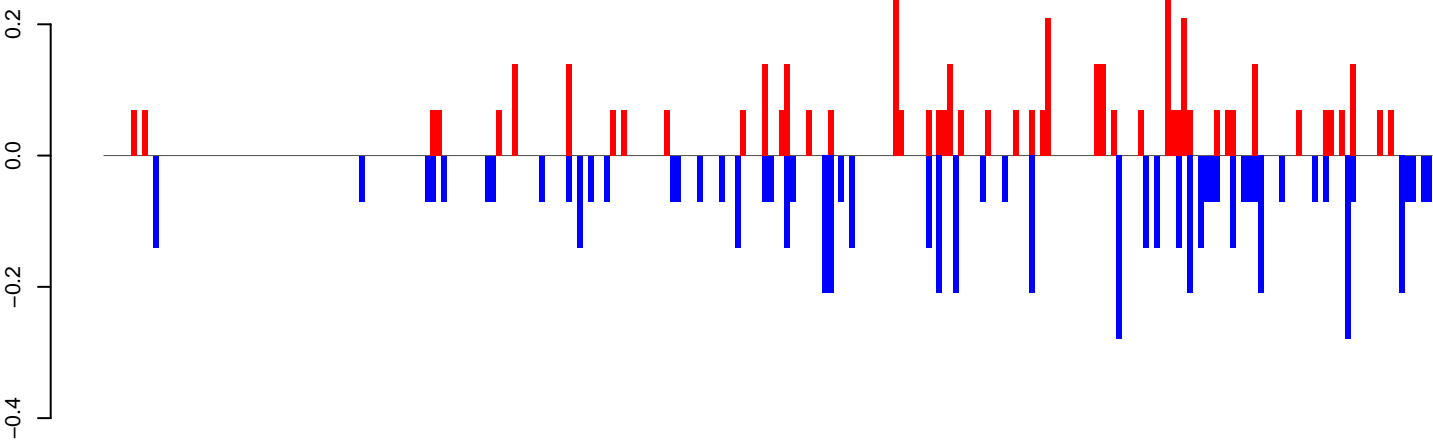
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CuQuin_Mcarc_NL.rep

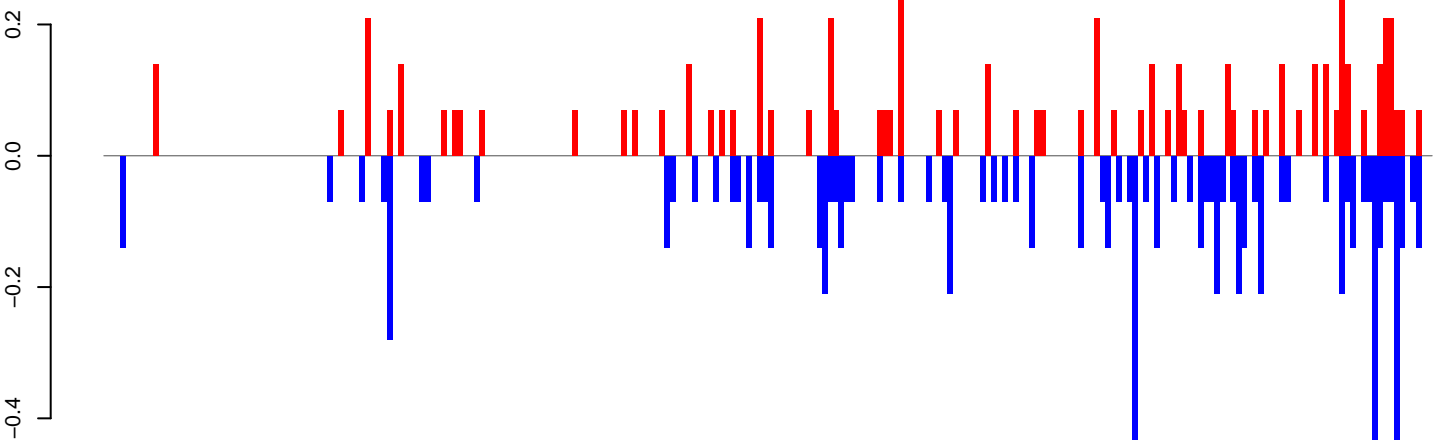


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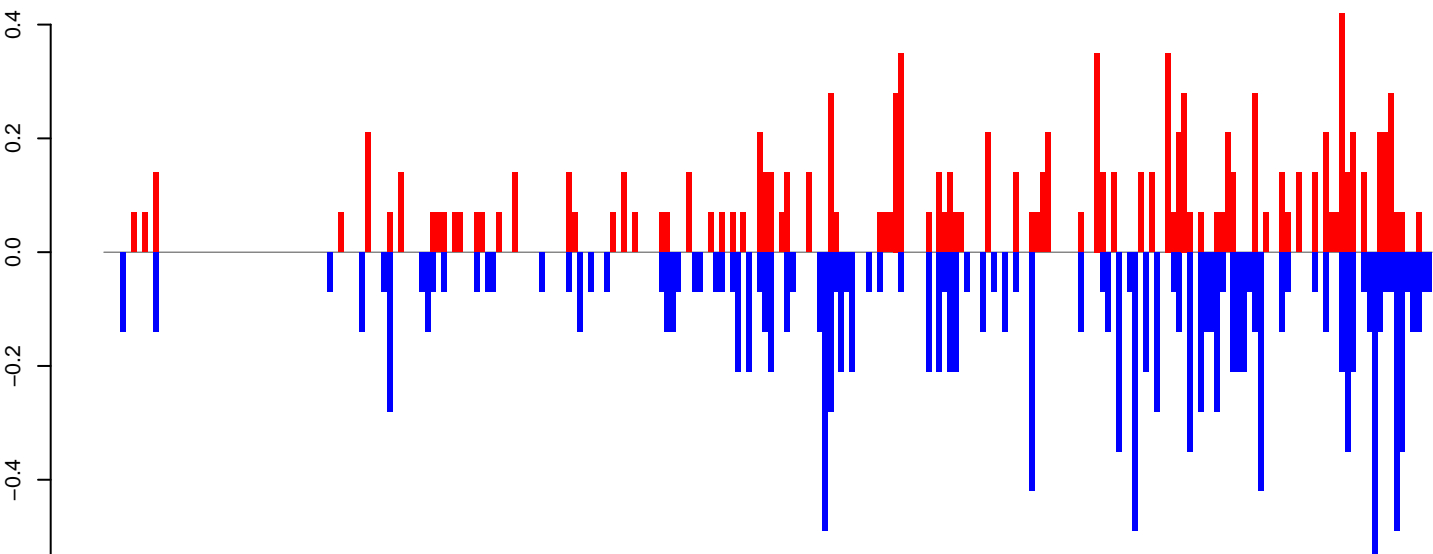
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CuQuin_Mcarc_NL.24_35.rep



positive=6, negative=8, total=15

CuQuin_Mcarc_NL.rep

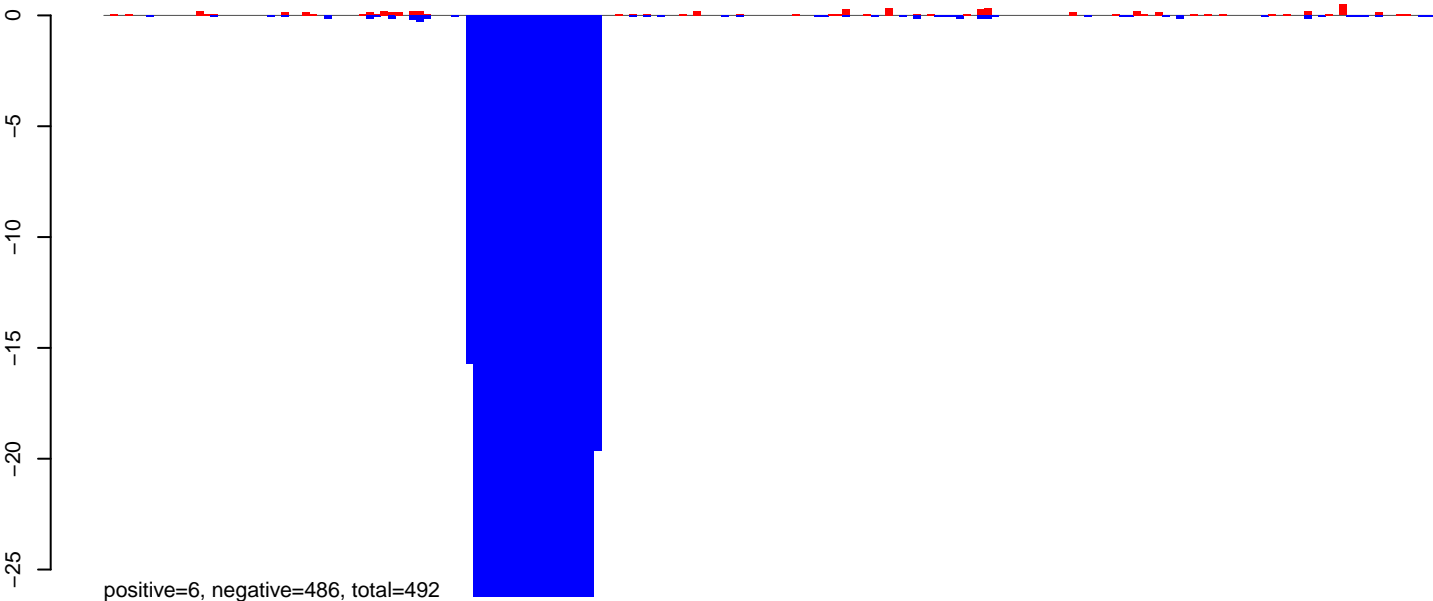


positive=11, negative=15, total=27

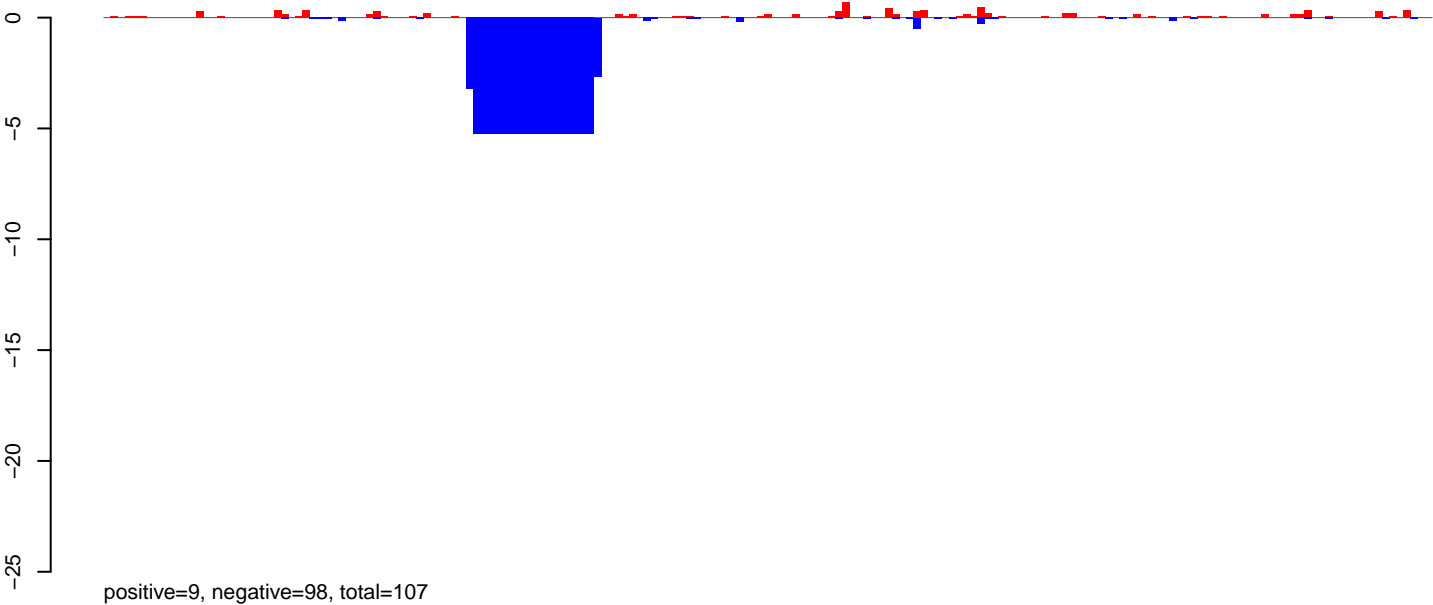
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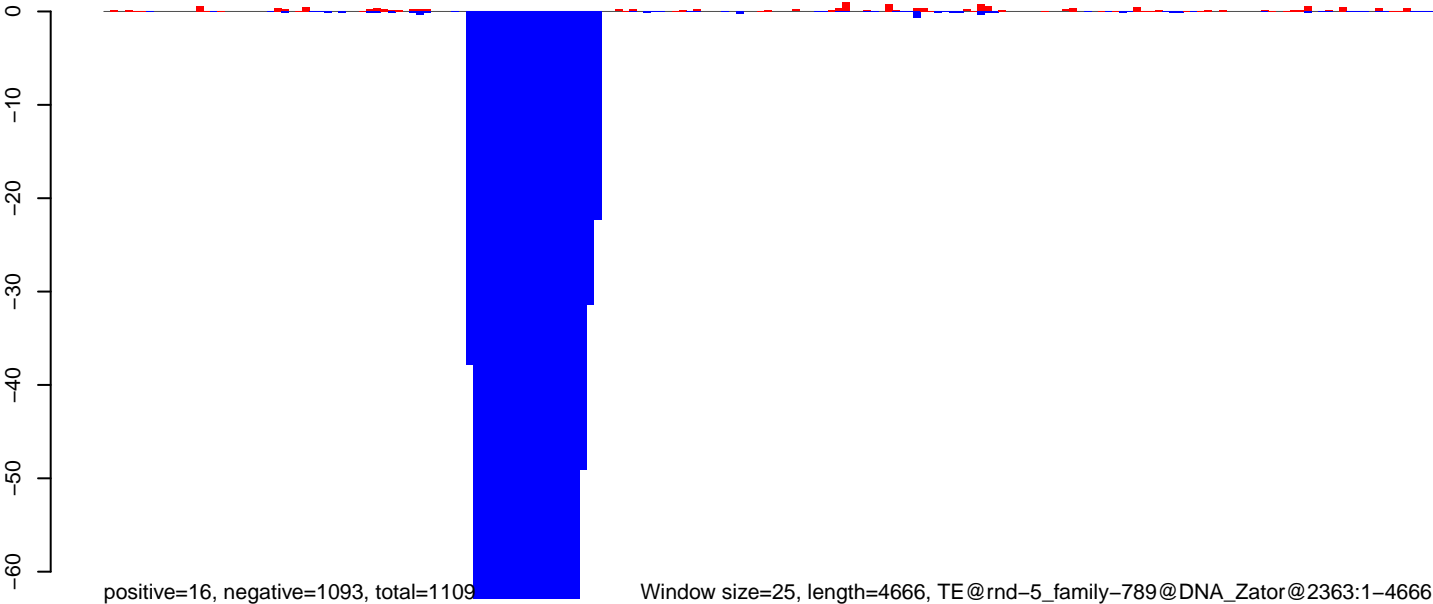
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



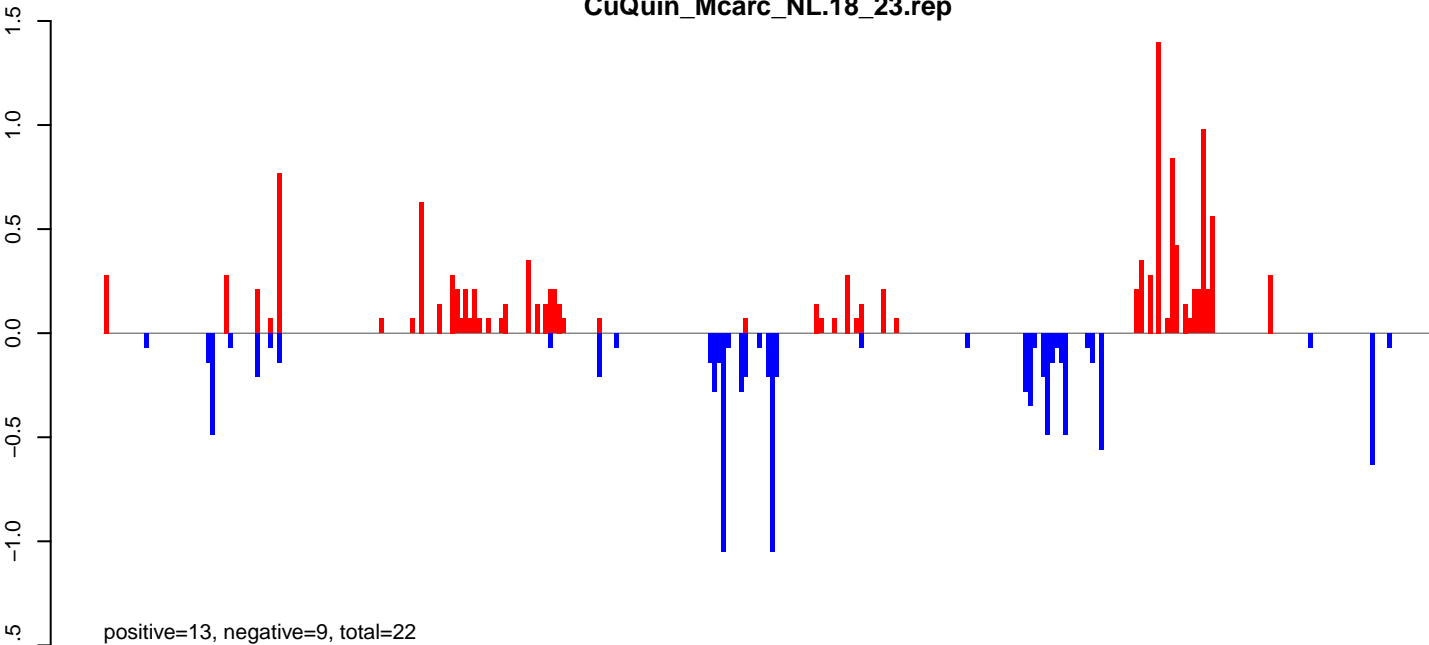
CuQuin_Mcarc_NL.rep



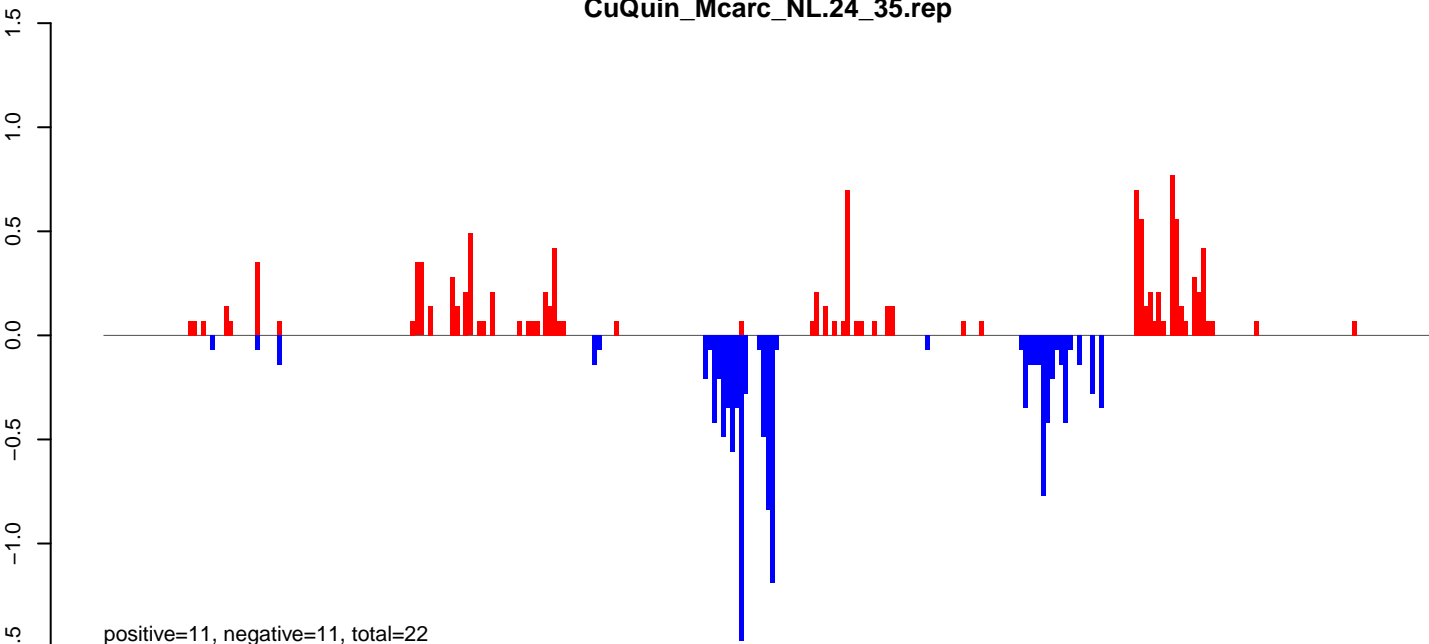
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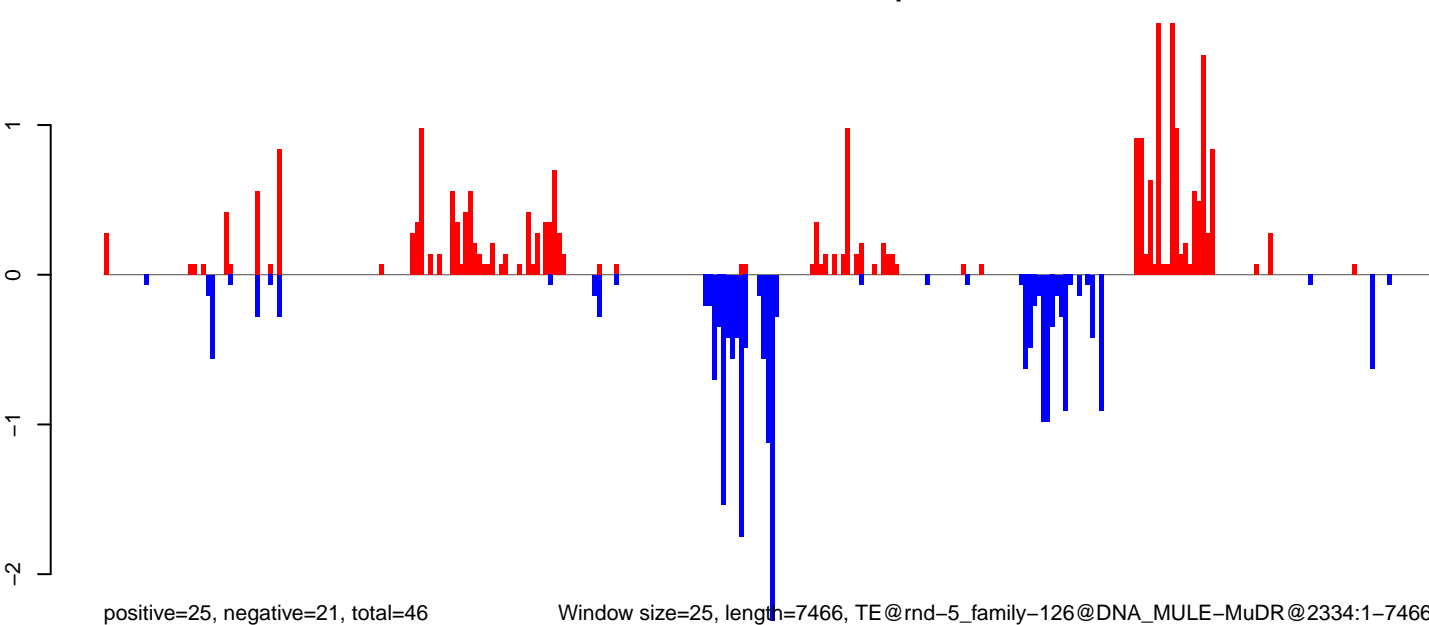
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

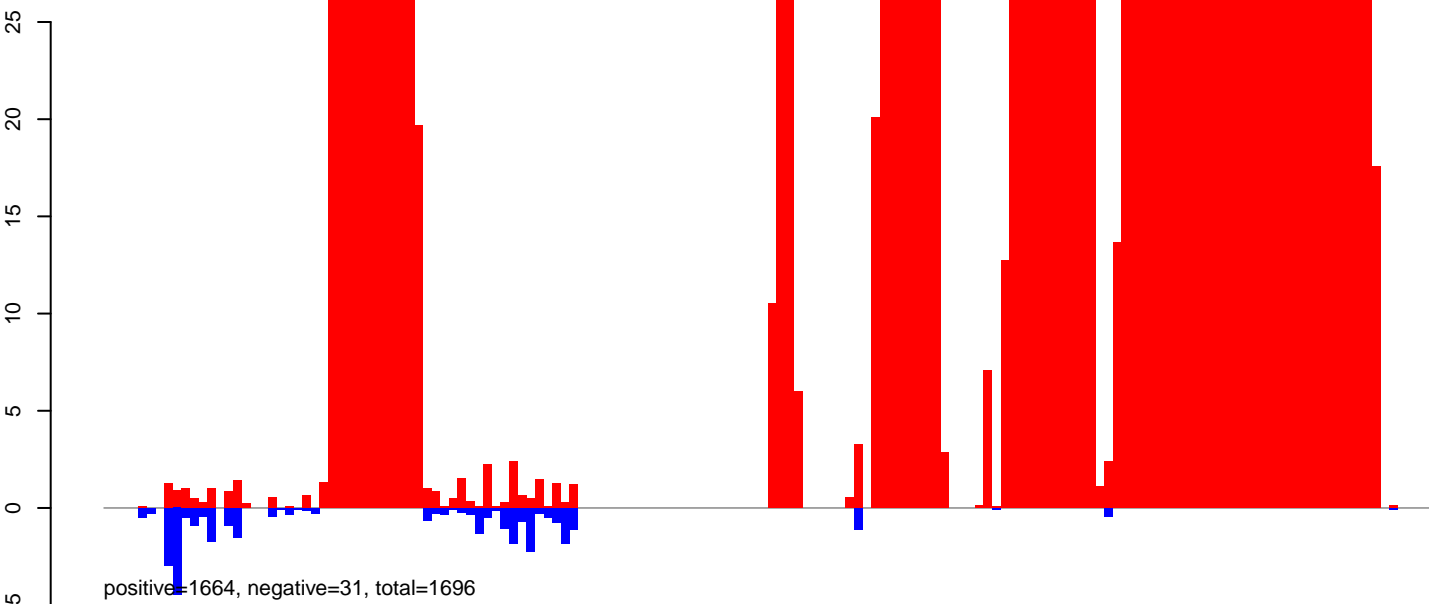


CuQuin_Mcarc_NL.rep

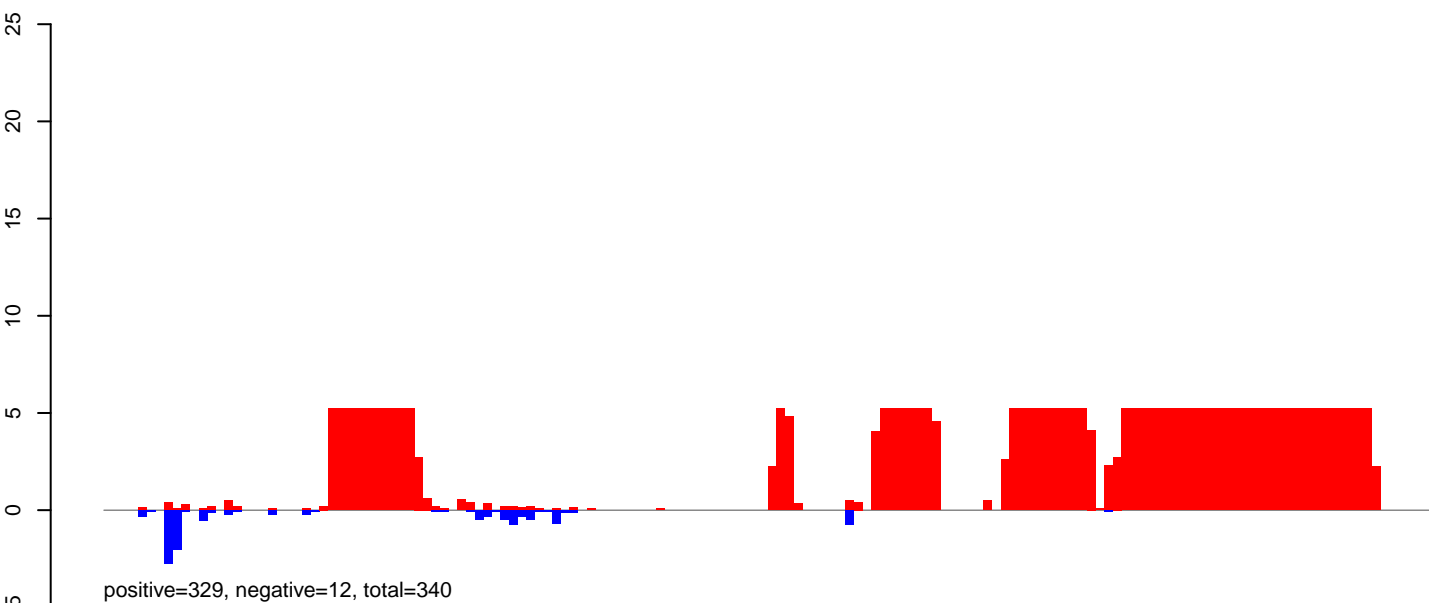


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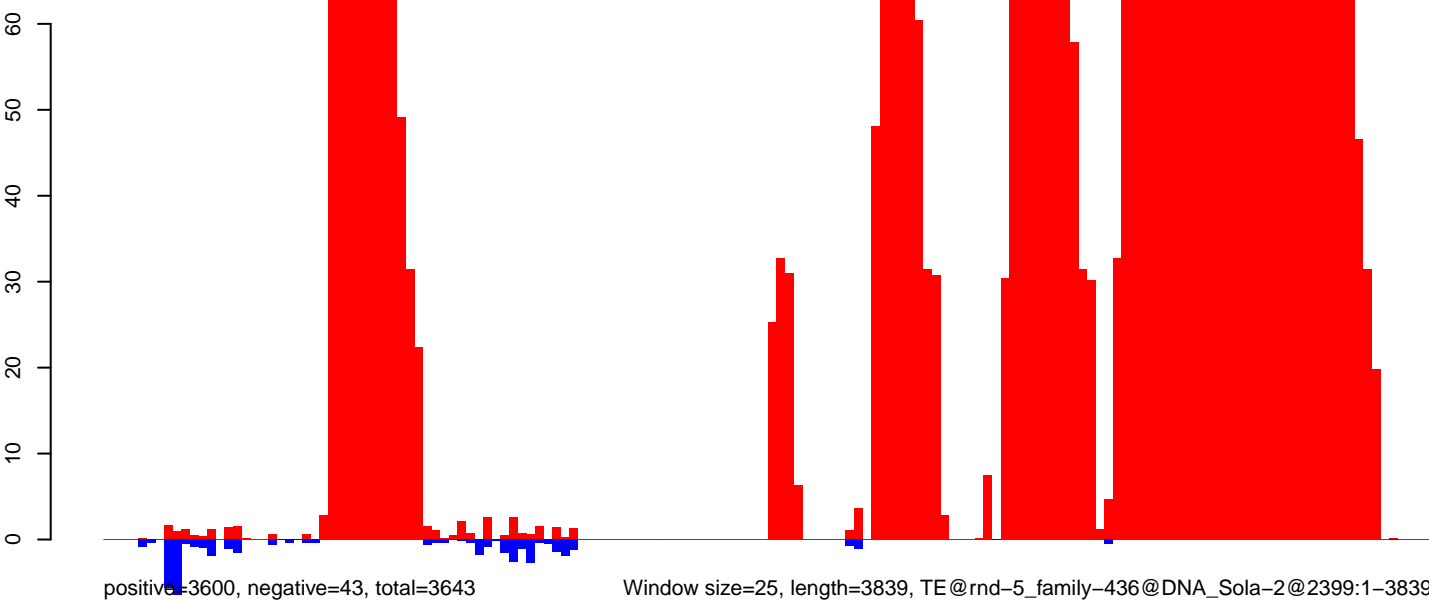
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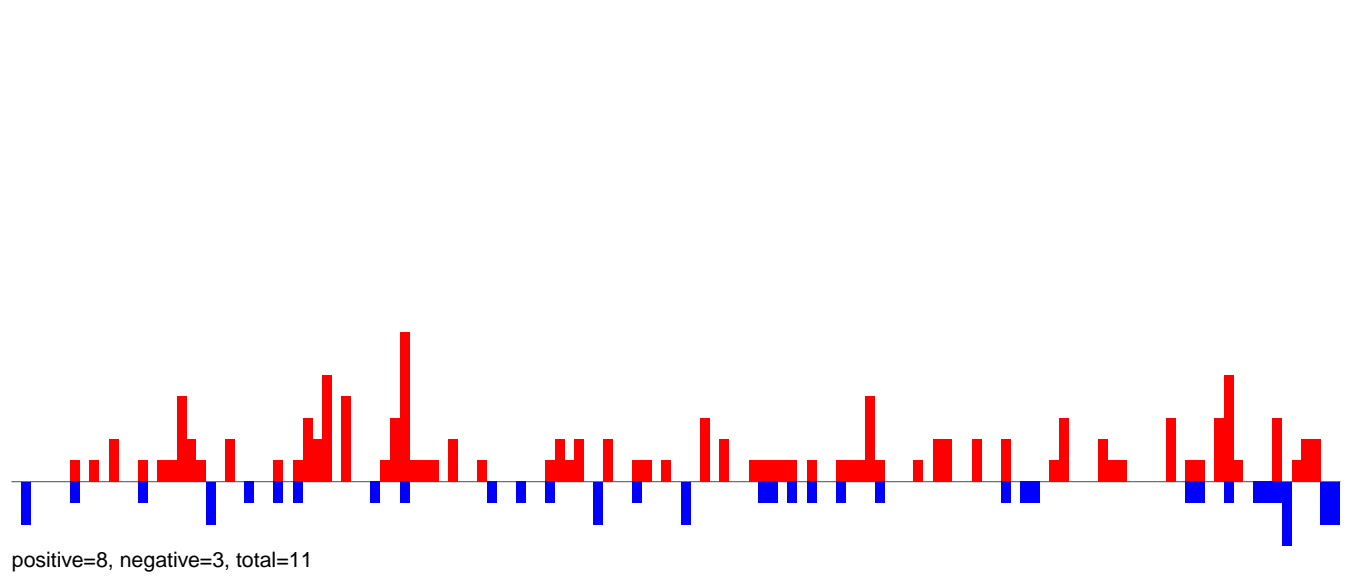
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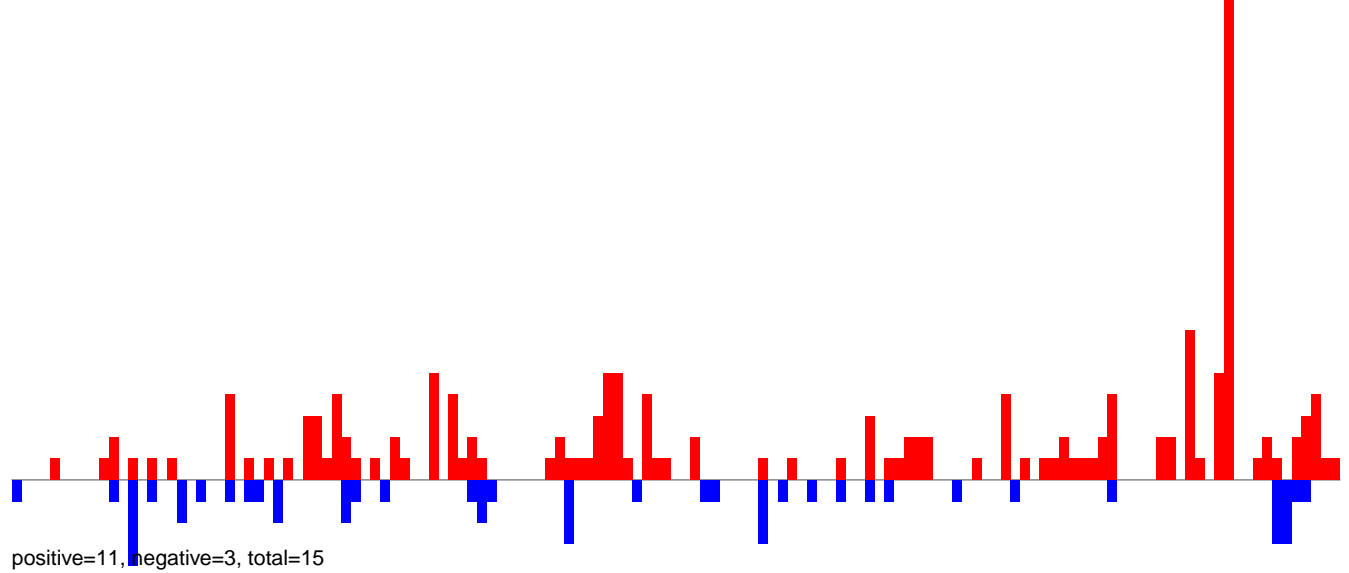
CuQuin_Mcarc_NL.rep



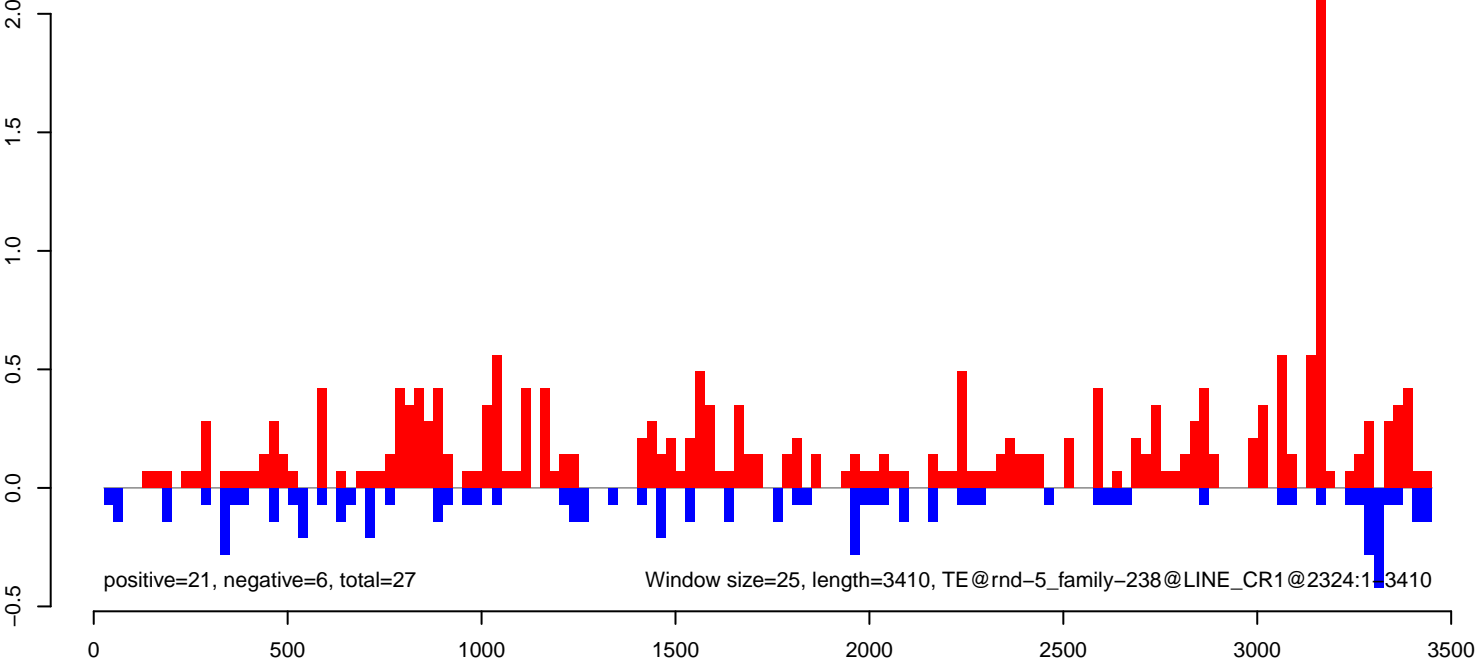
CuQuin_Mcarc_NL.18_23.rep



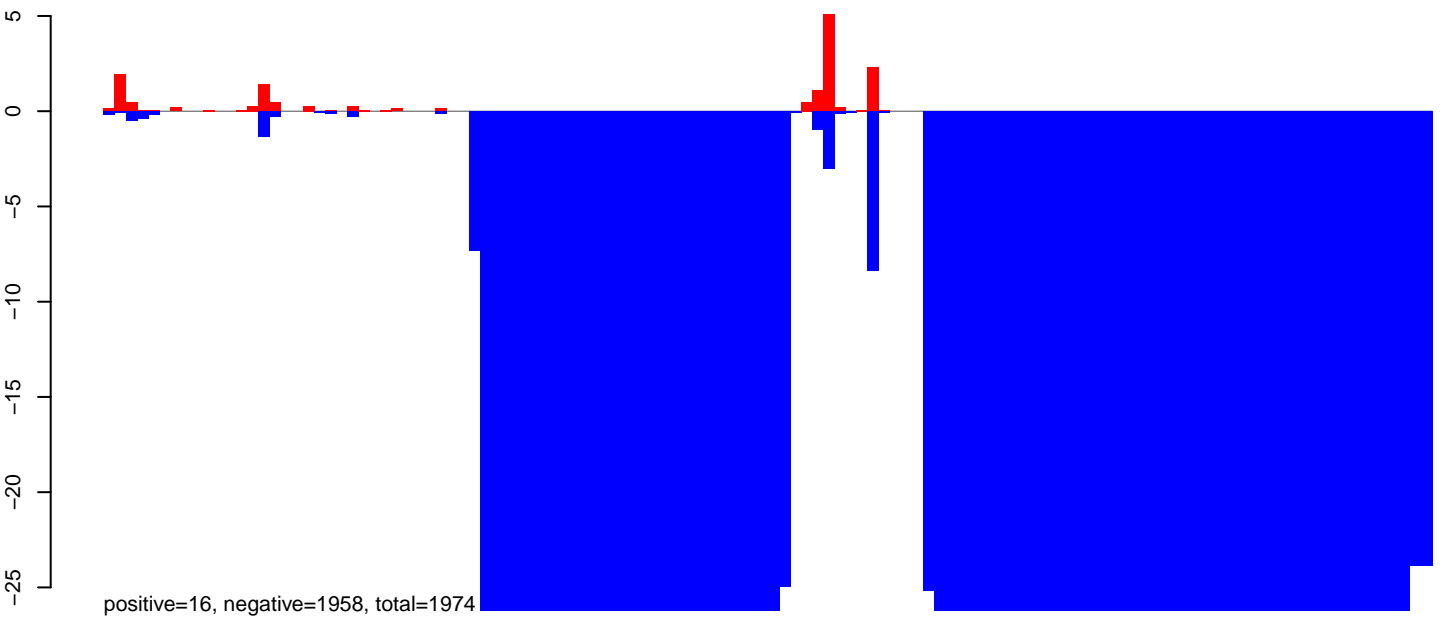
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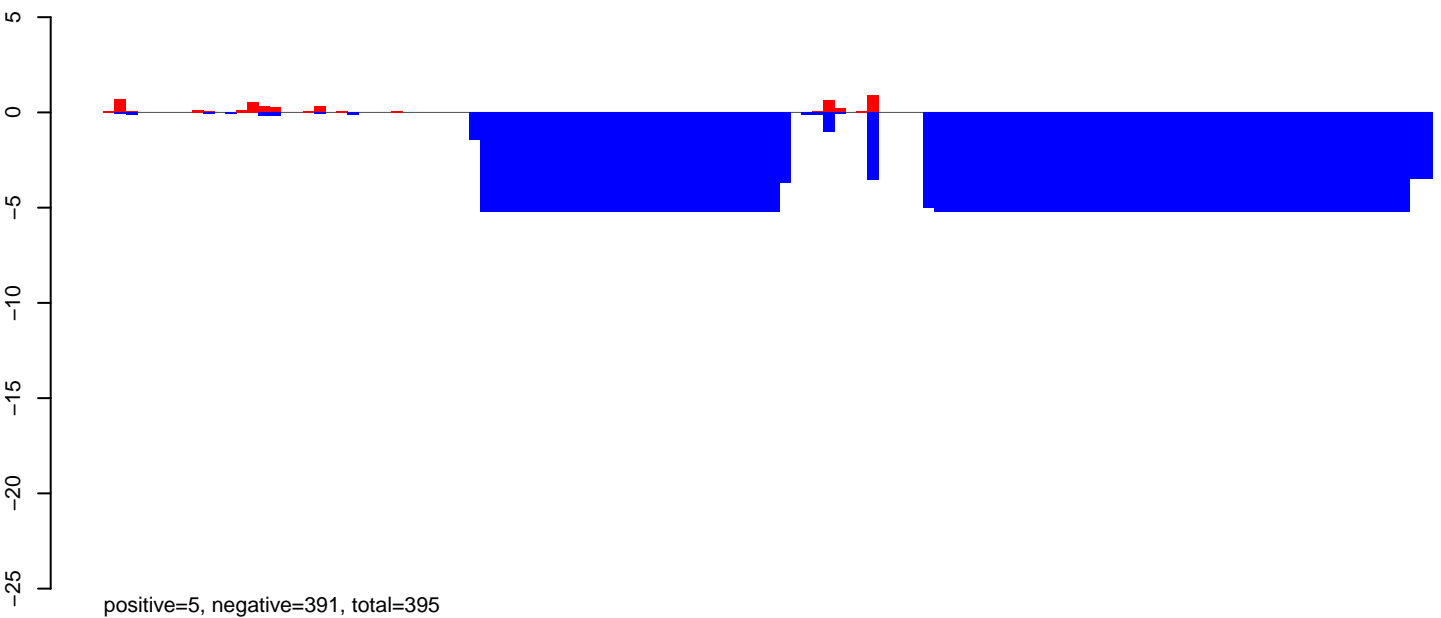
CuQuin_Mcarc_NL.rep



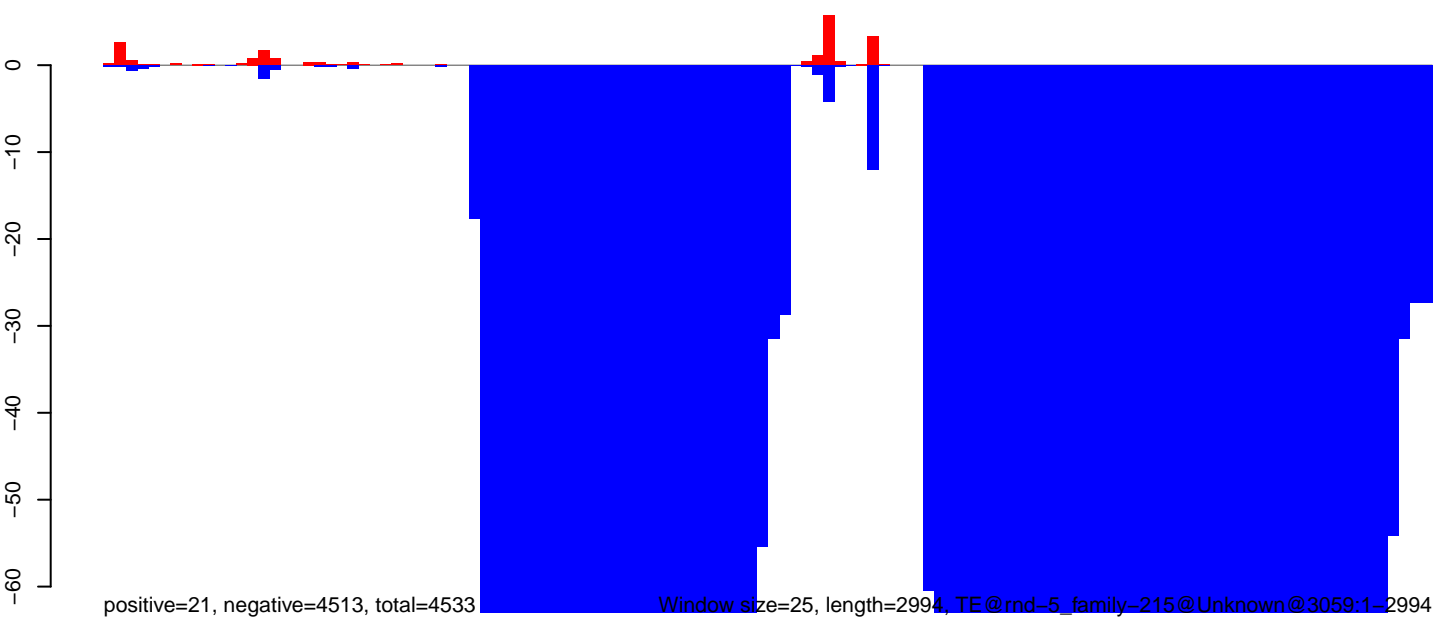
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

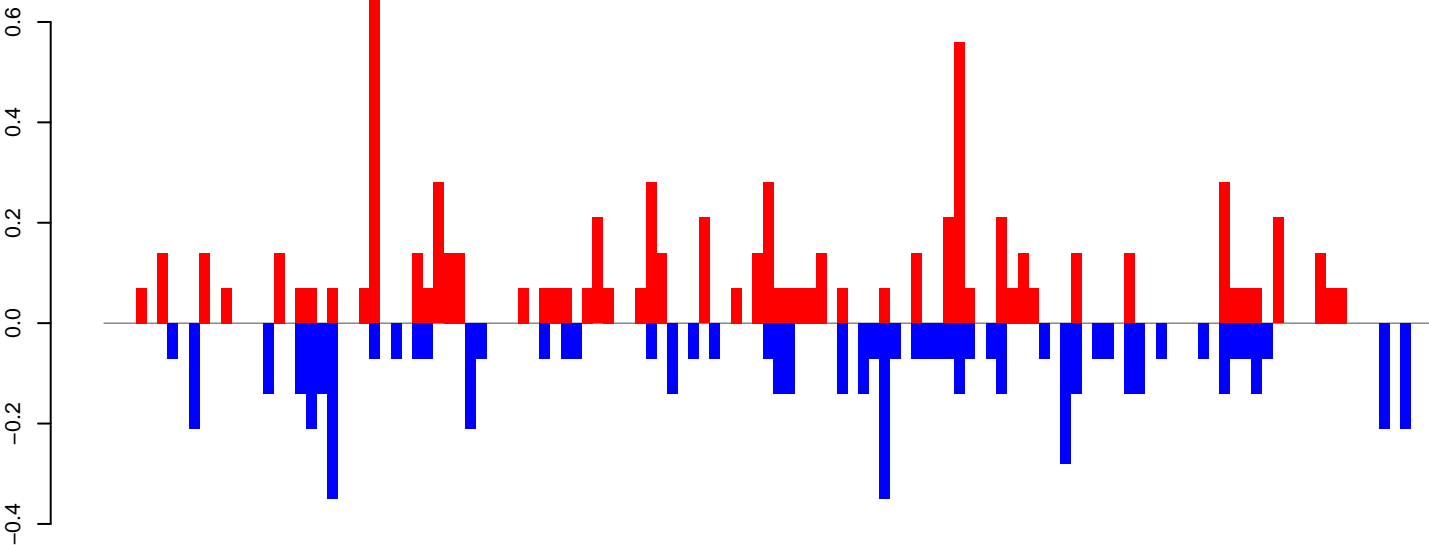


CuQuin_Mcarc_NL.rep



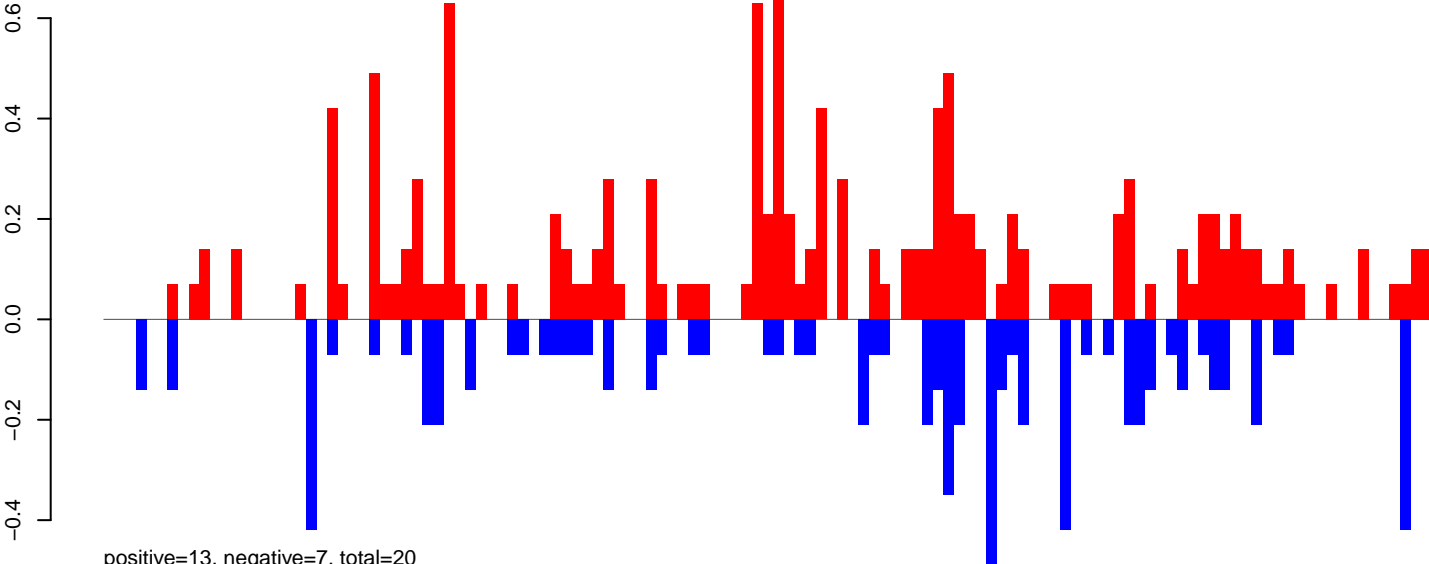
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CuQuin_Mcarc_NL.18_23.rep



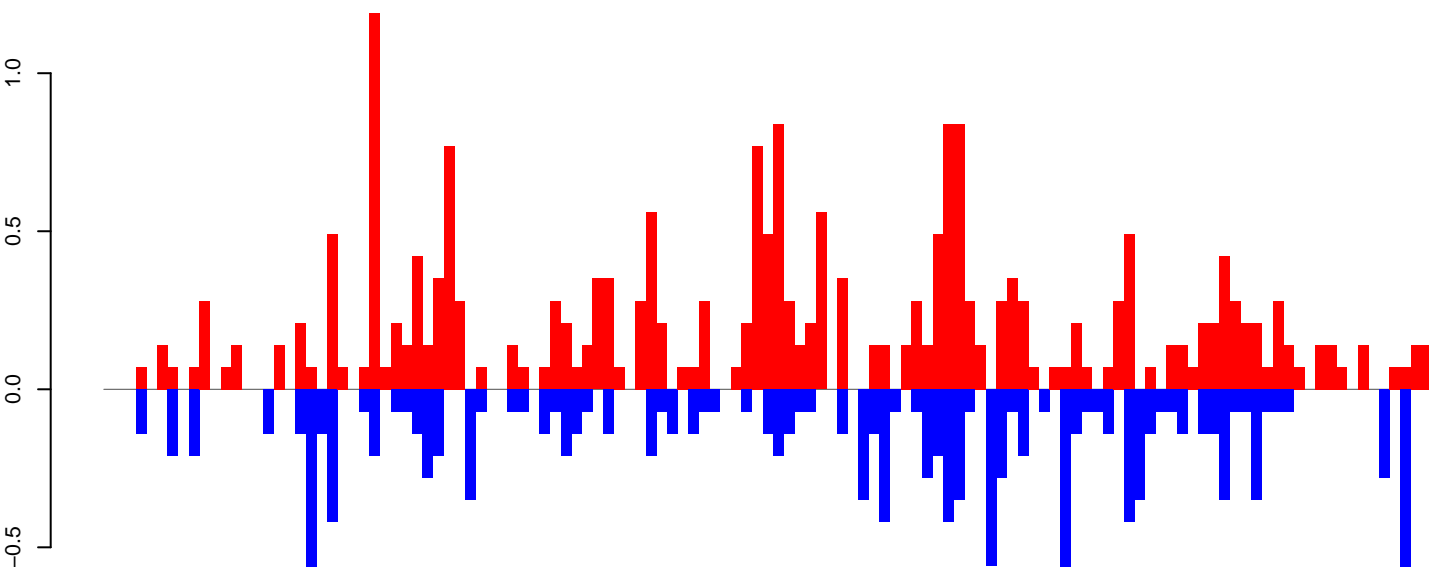
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CuQuin_Mcarc_NL.24_35.rep



positive=13, negative=7, total=20

CuQuin_Mcarc_NL.rep

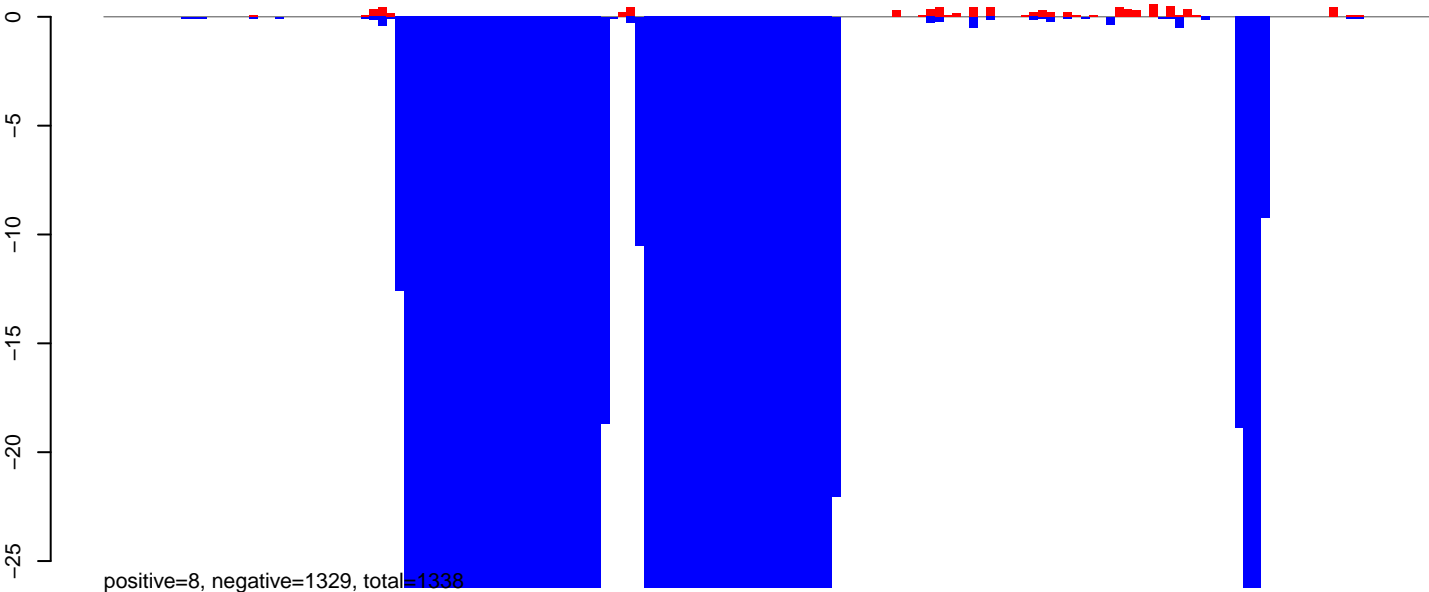


positive=22, negative=14, total=36

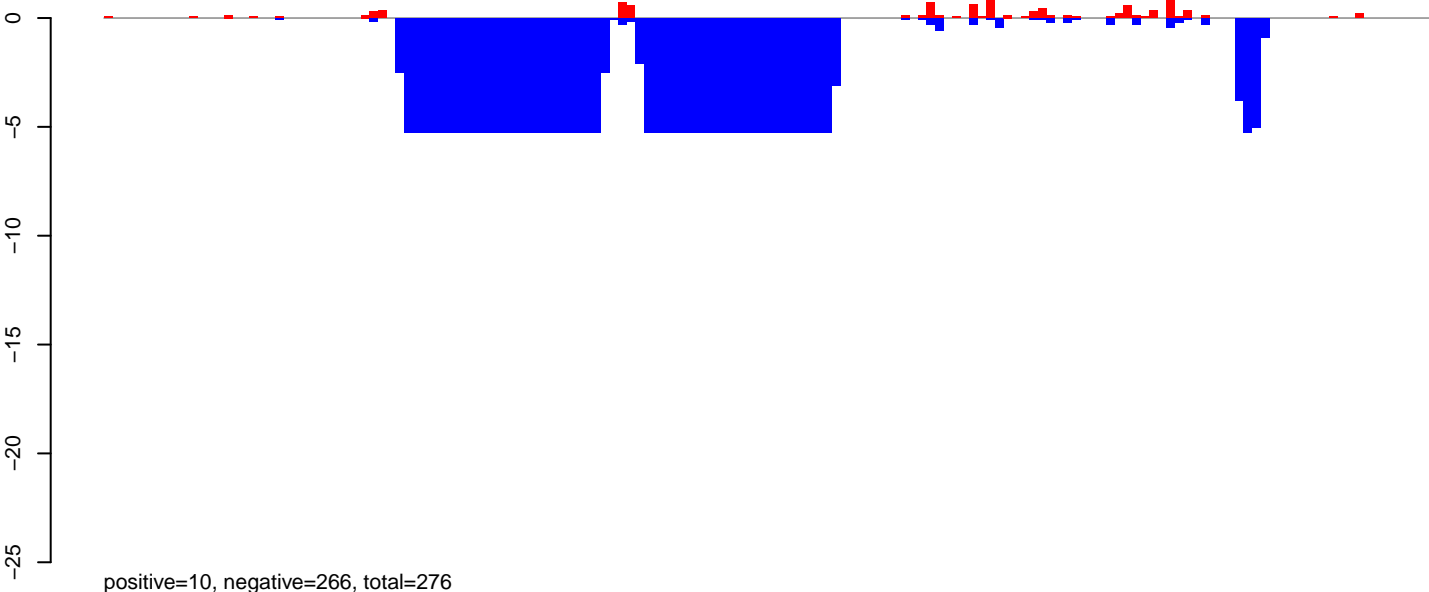
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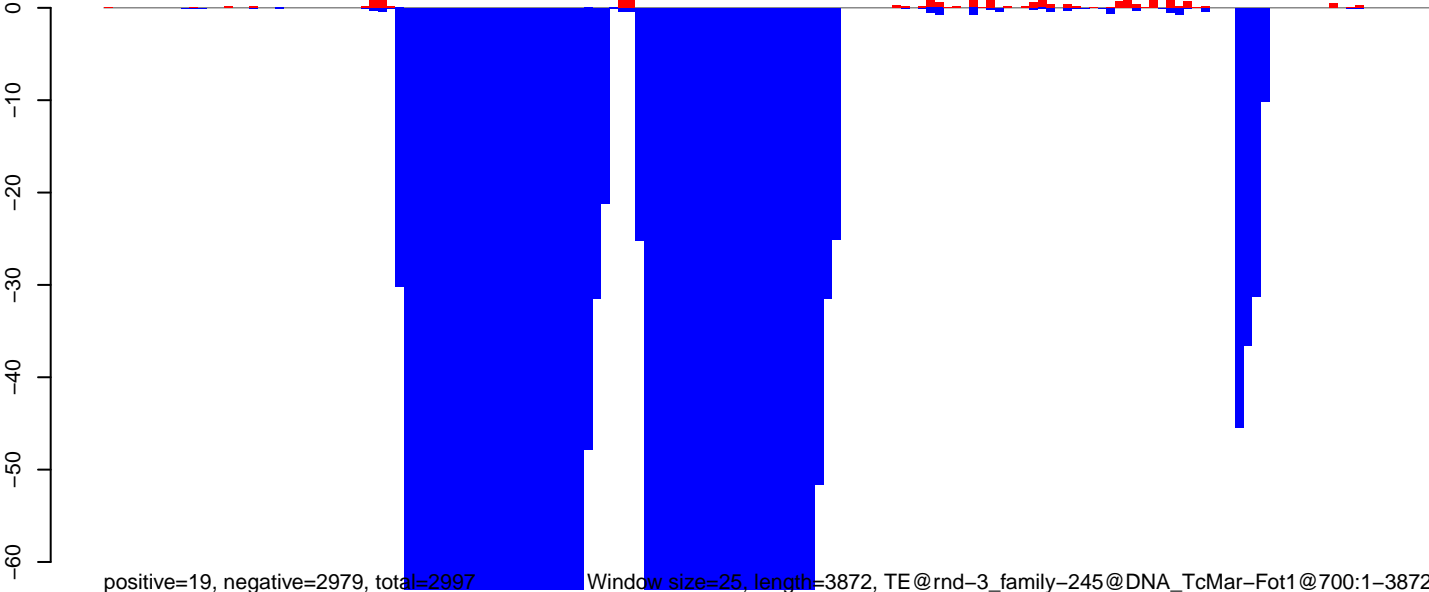
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CuQuin_Mcarc_NL.24_35.rep

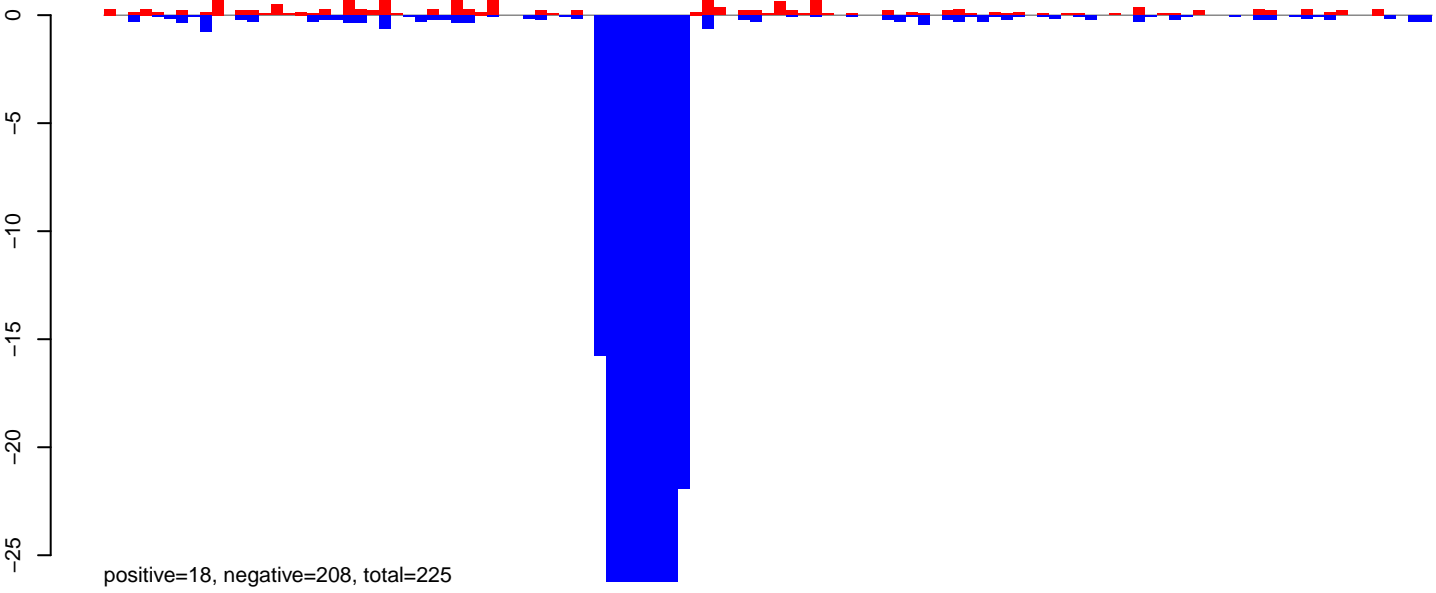


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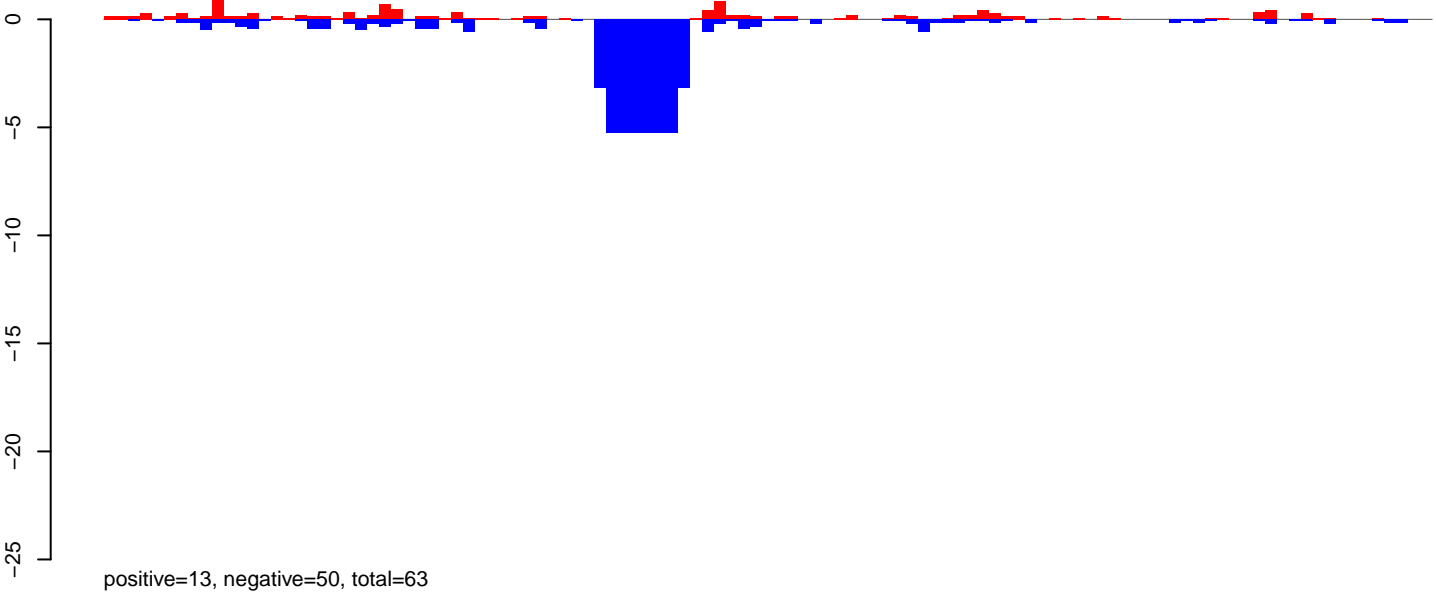


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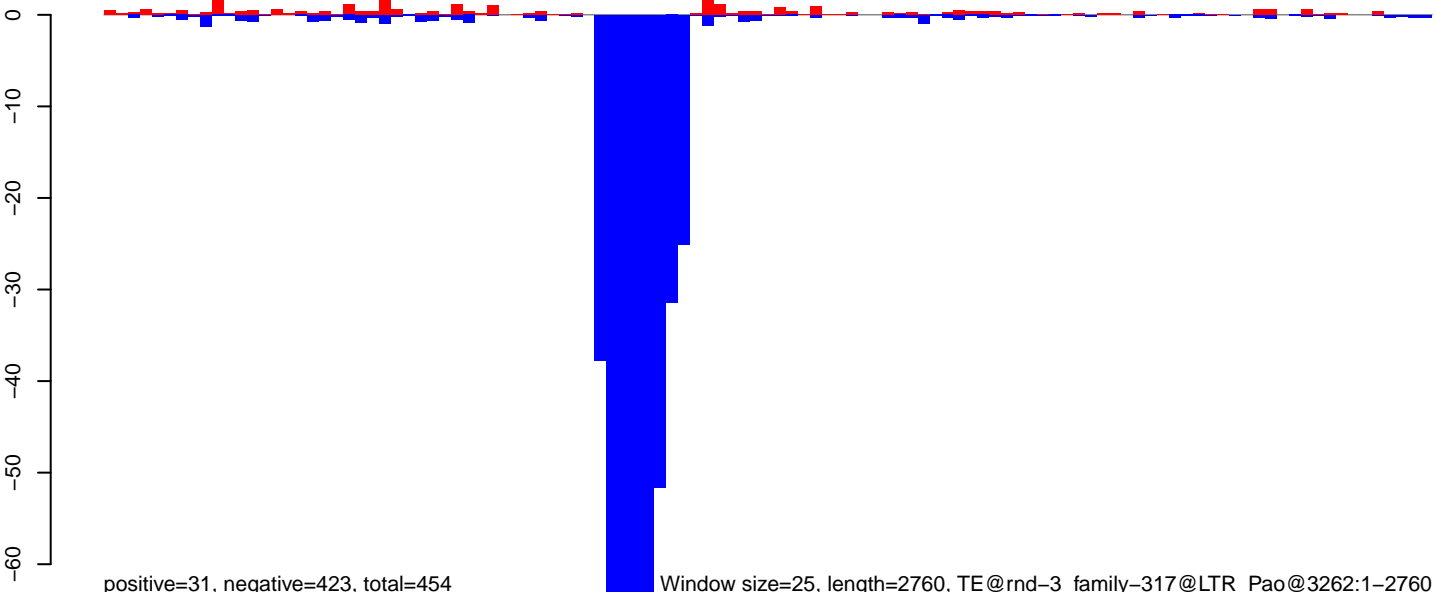
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CuQuin_Mcarc_NL.24_35.rep

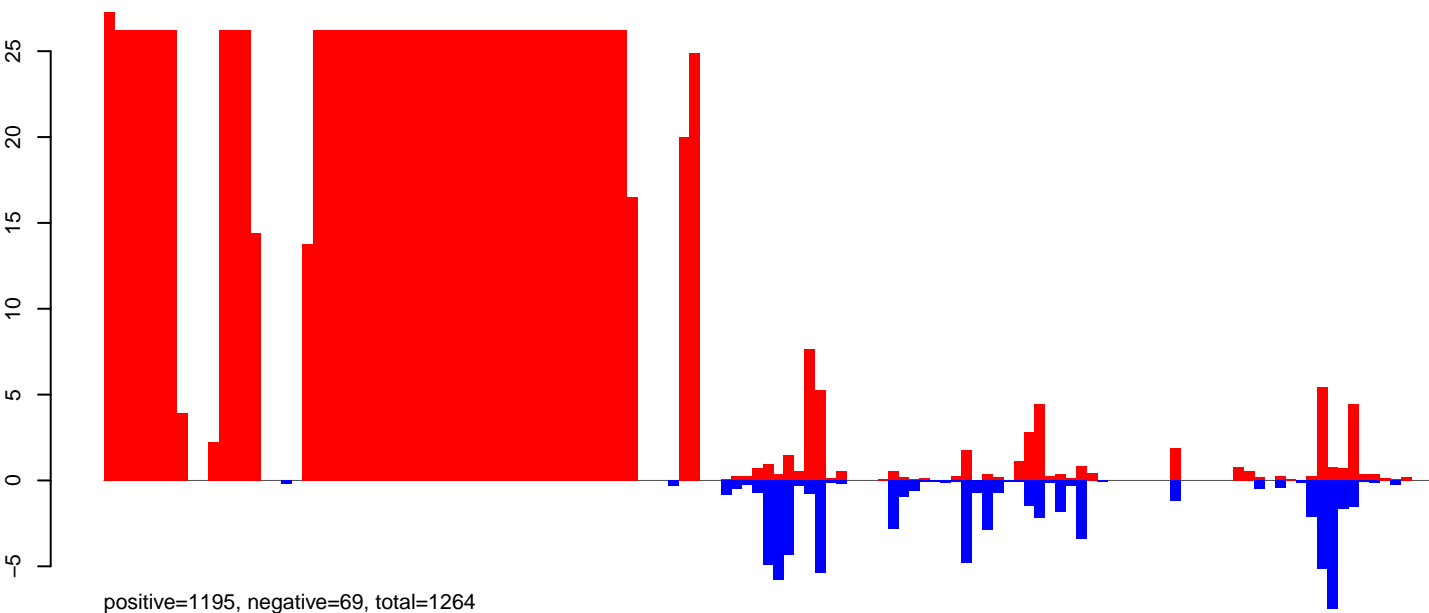


CuQuin_Mcarc_NL.rep

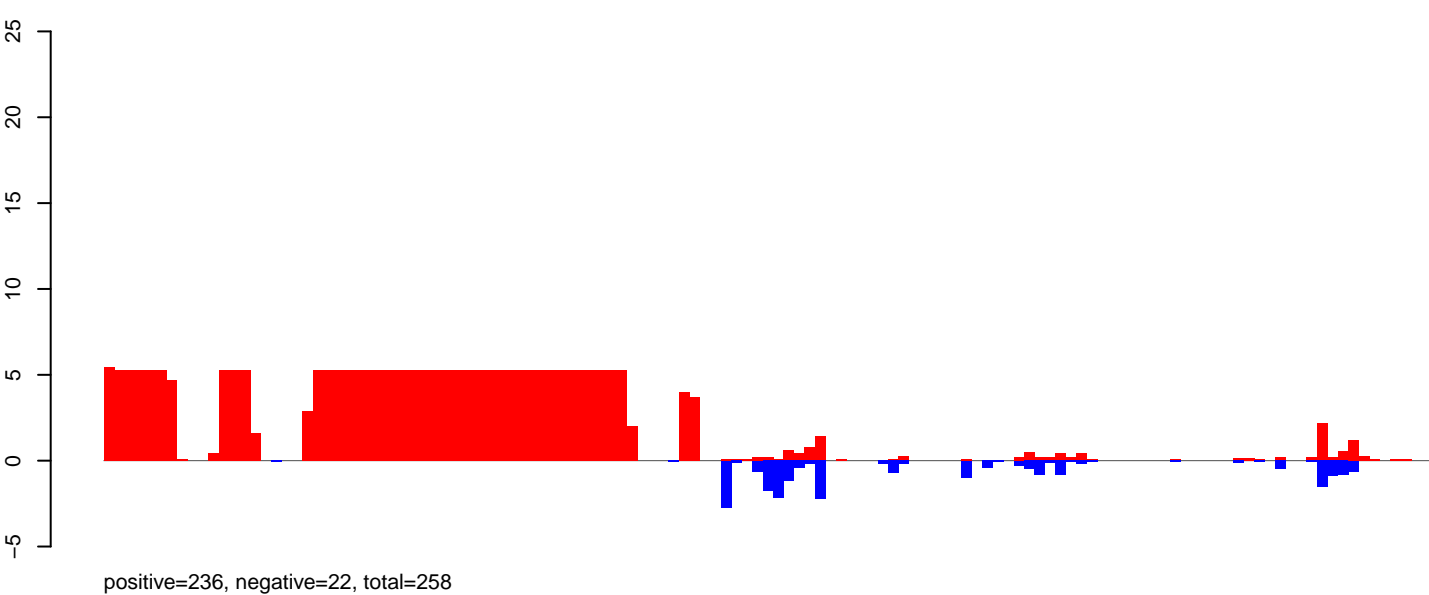


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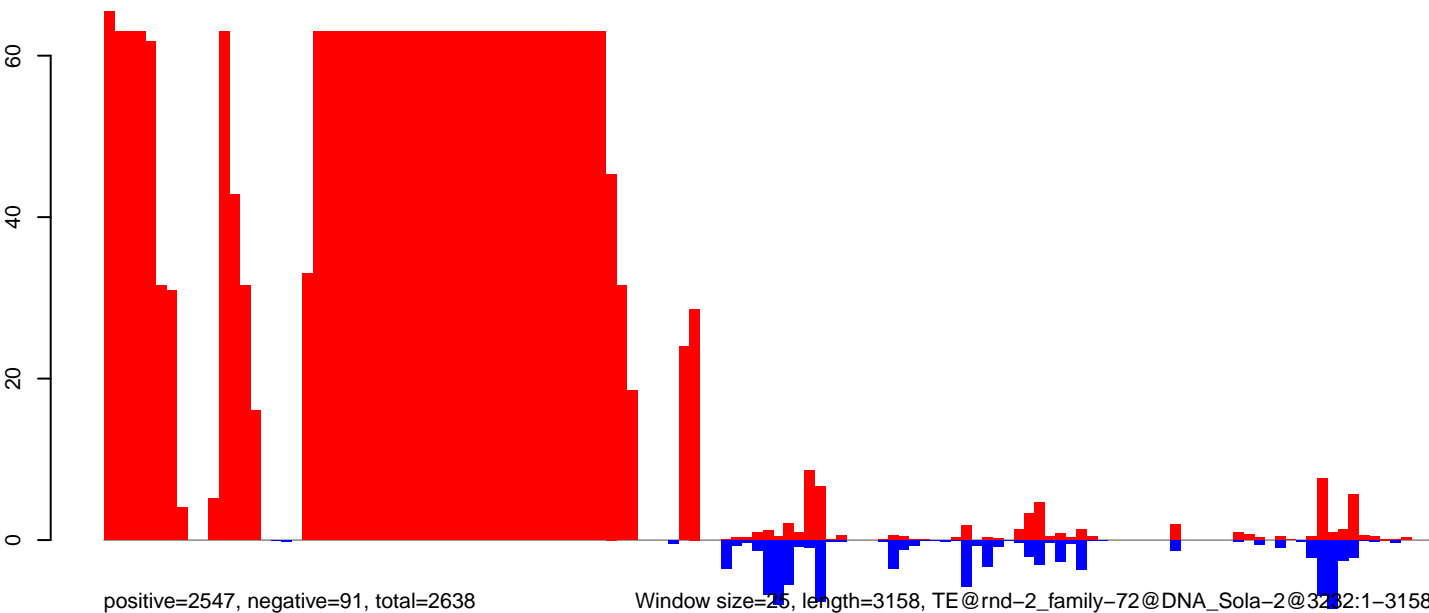
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

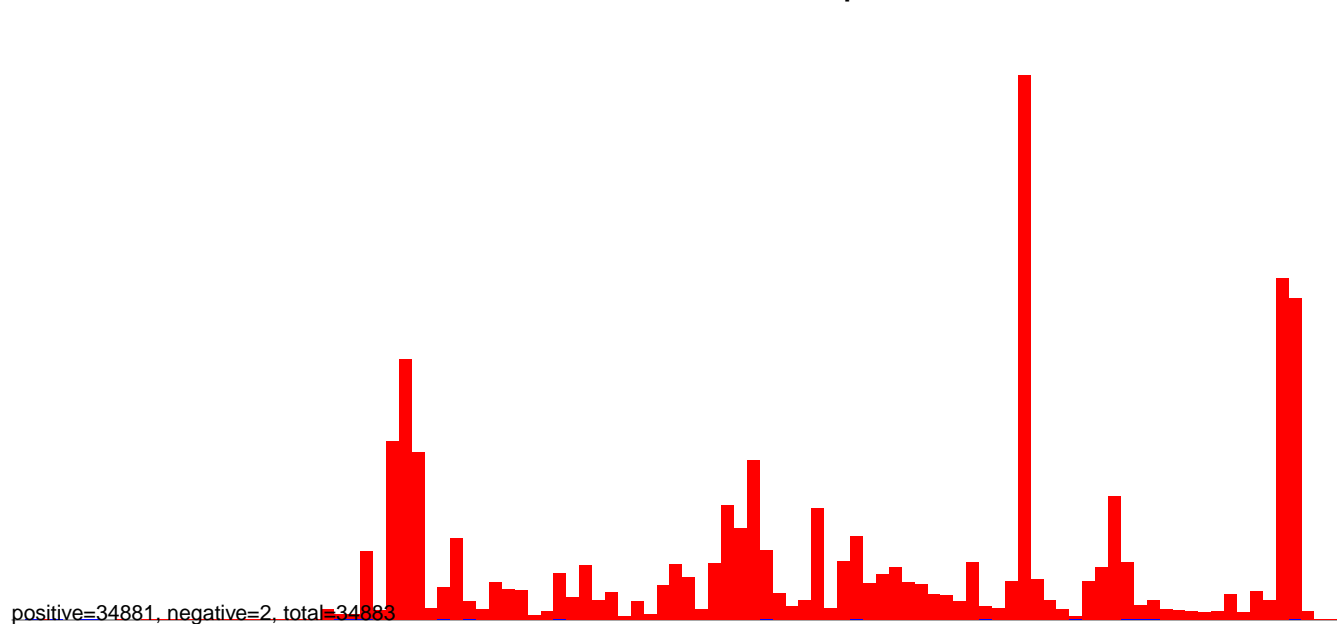


CuQuin_Mcarc_NL.rep

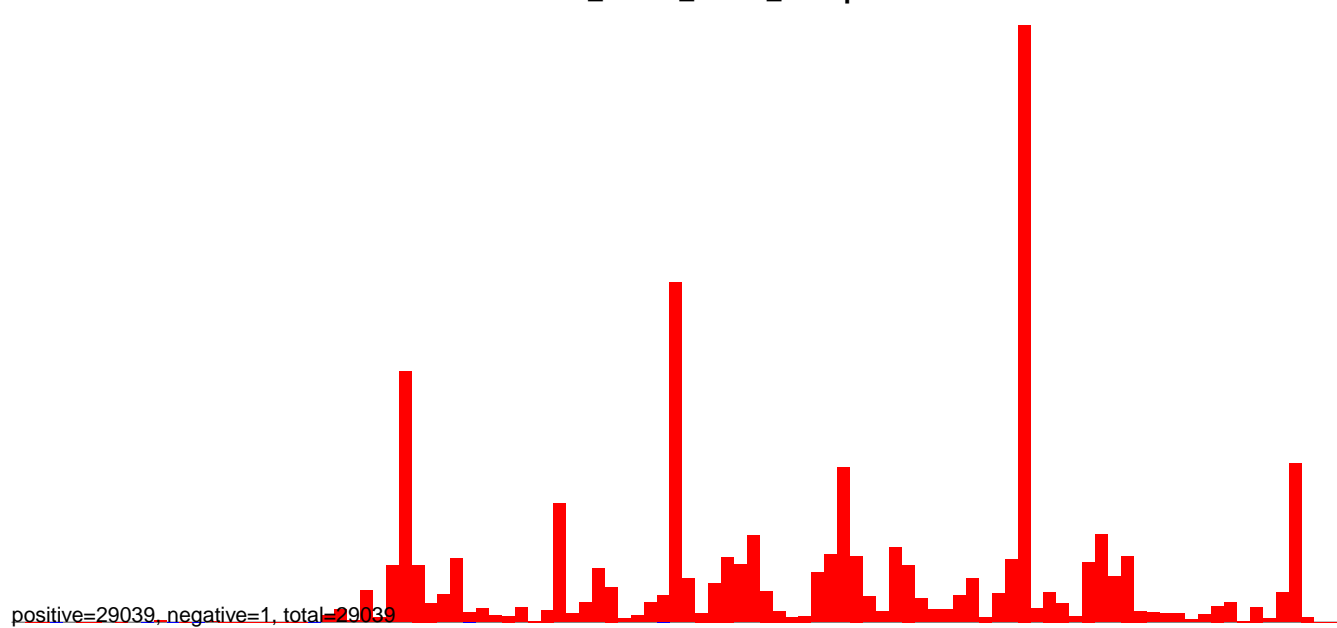


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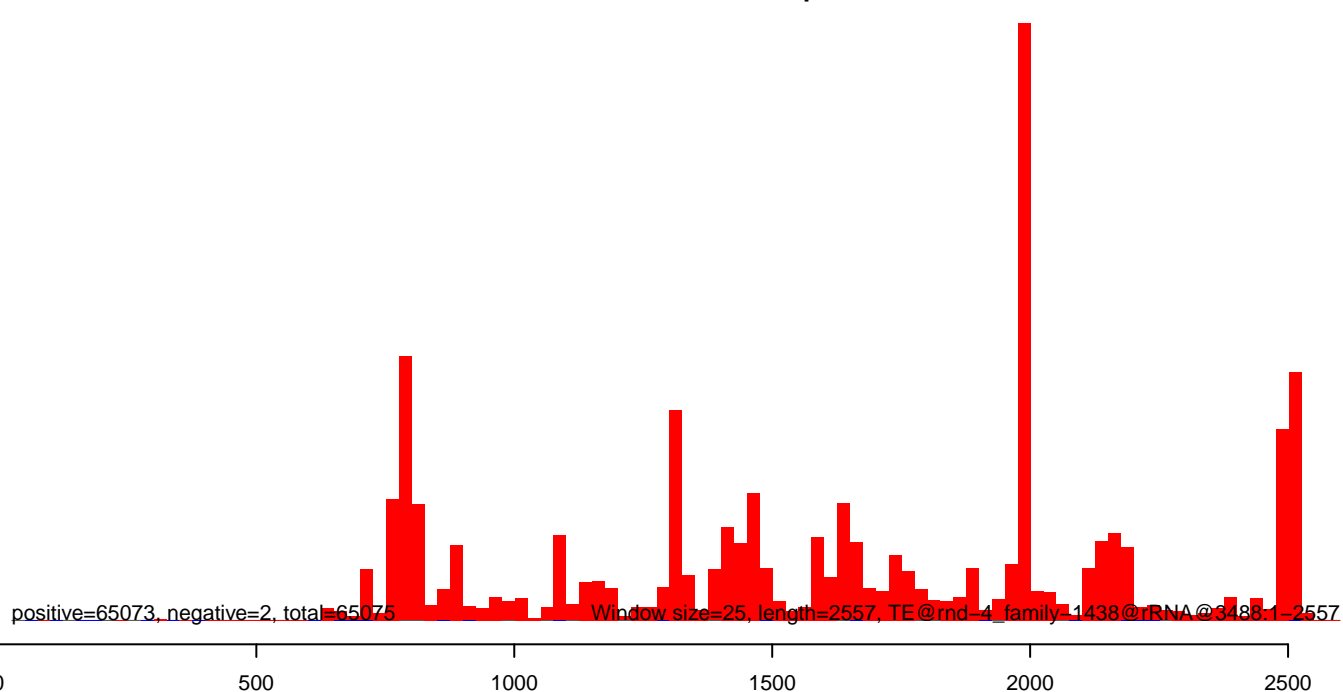
CuQuin_Mcarc_NL.18_23.rep



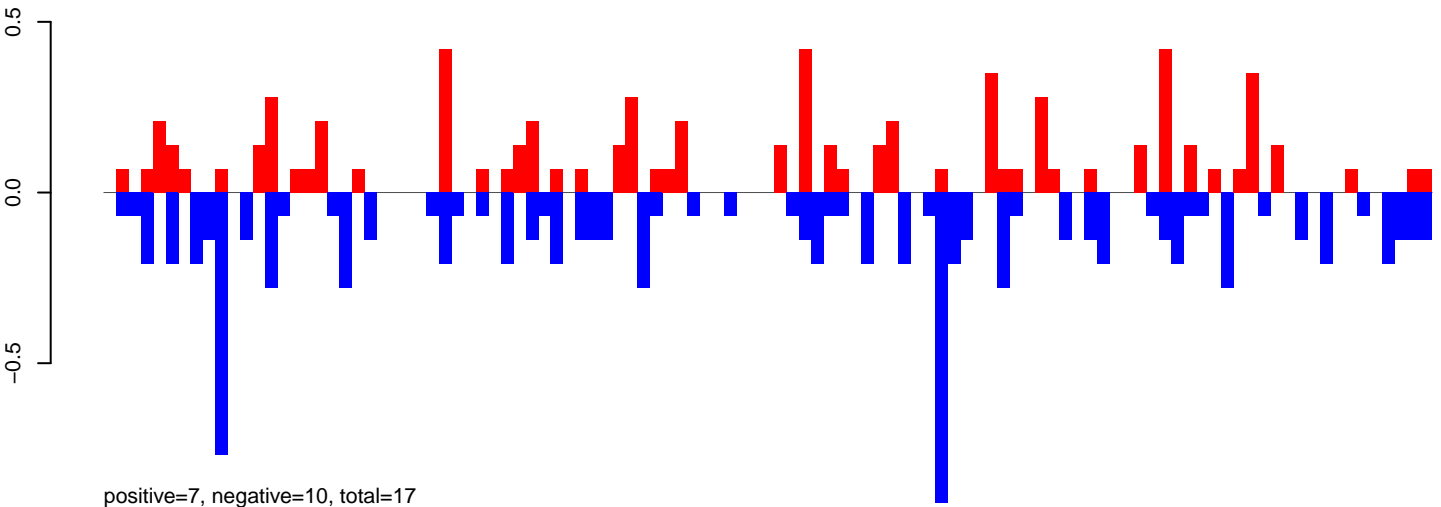
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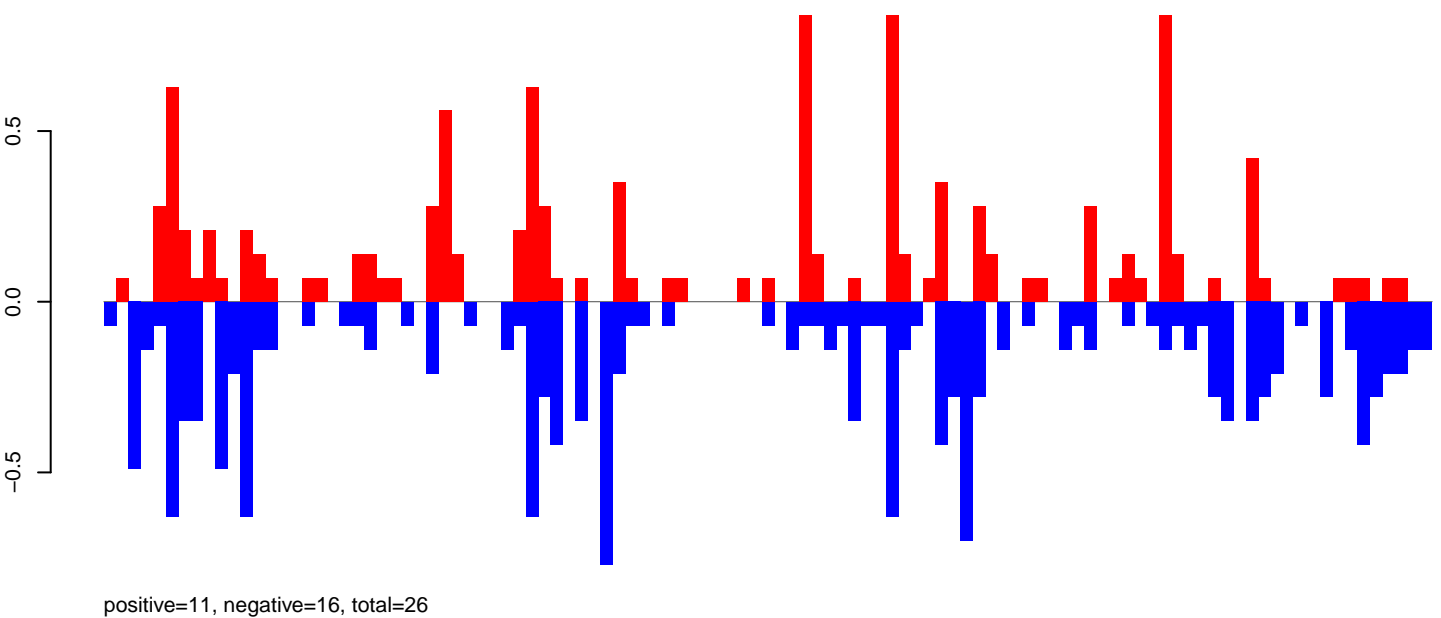
CuQuin_Mcarc_NL.rep



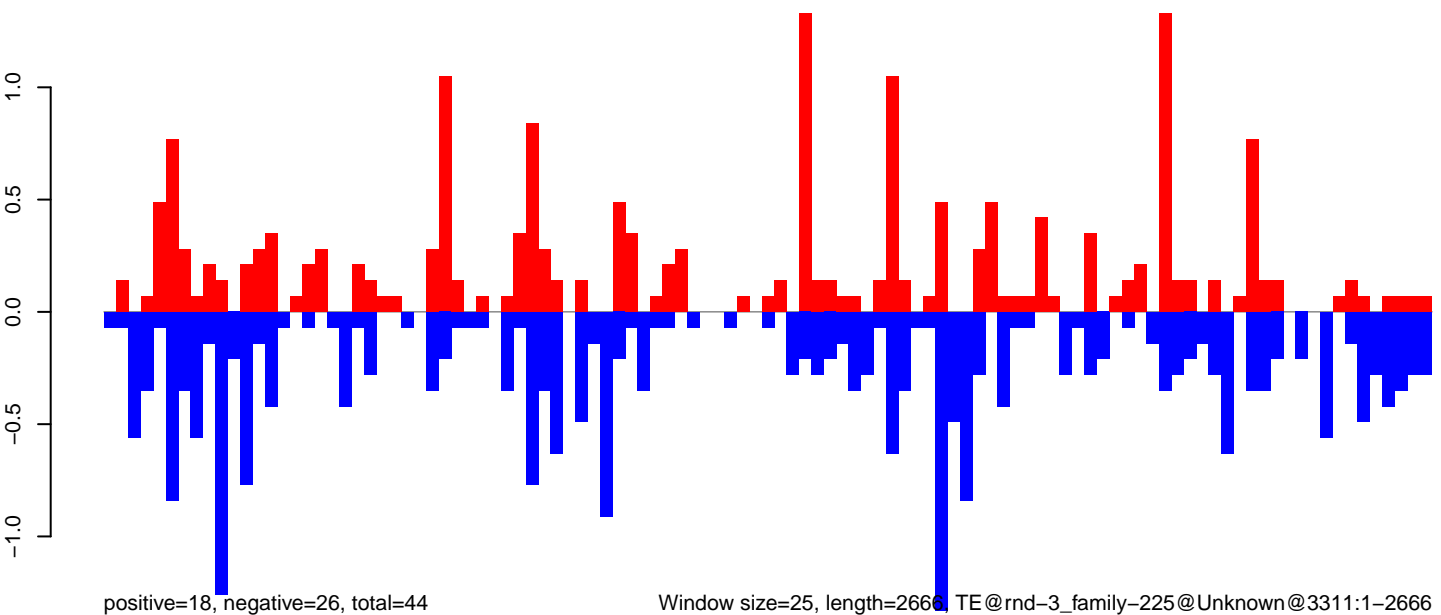
CuQuin_Mcarc_NL.18_23.rep



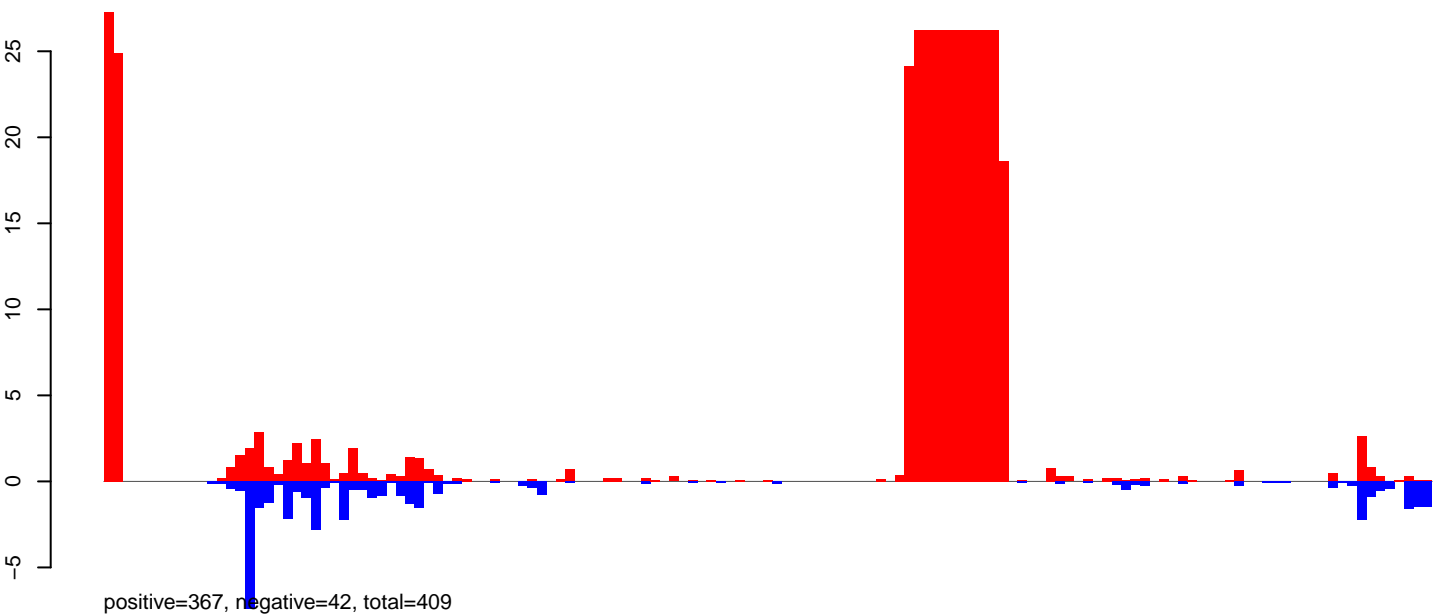
CuQuin_Mcarc_NL.24_35.rep



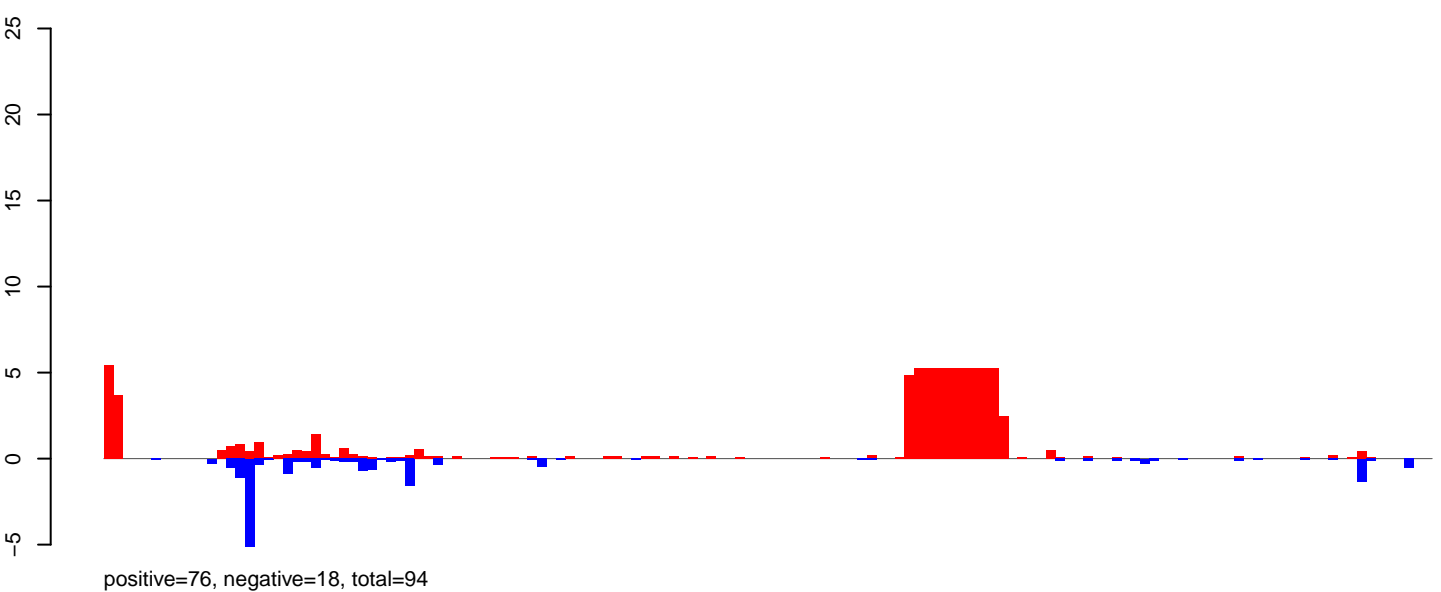
CuQuin_Mcarc_NL.rep



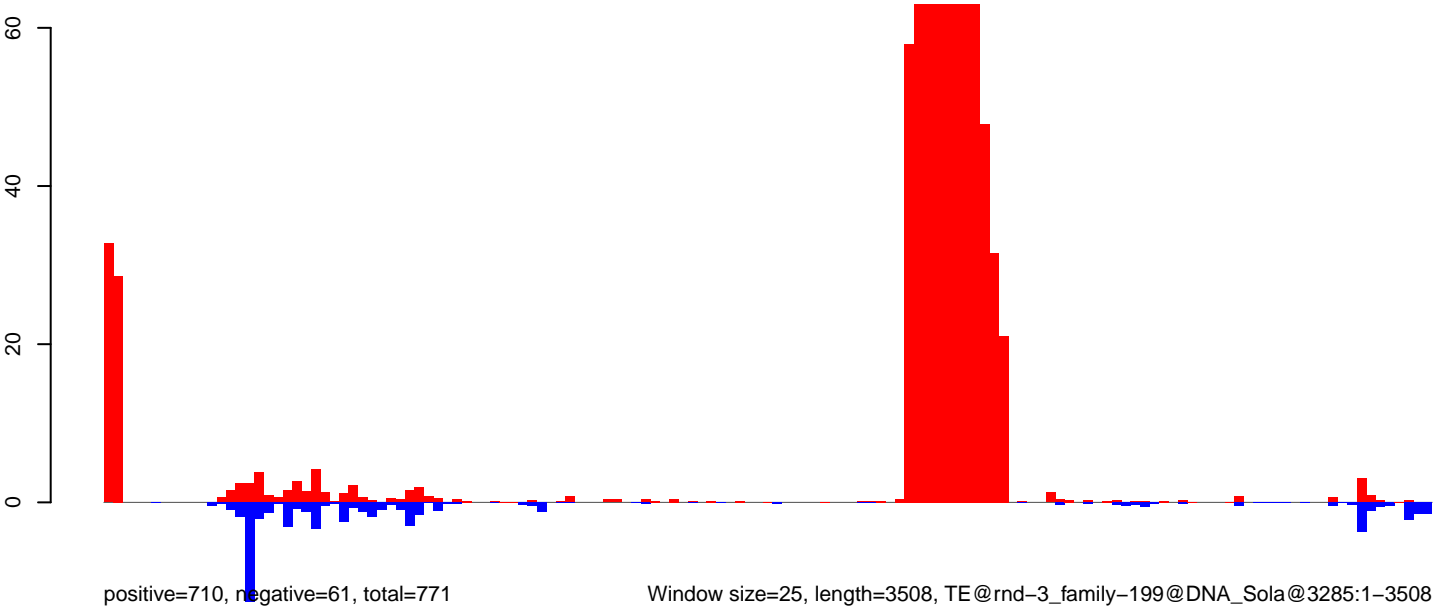
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



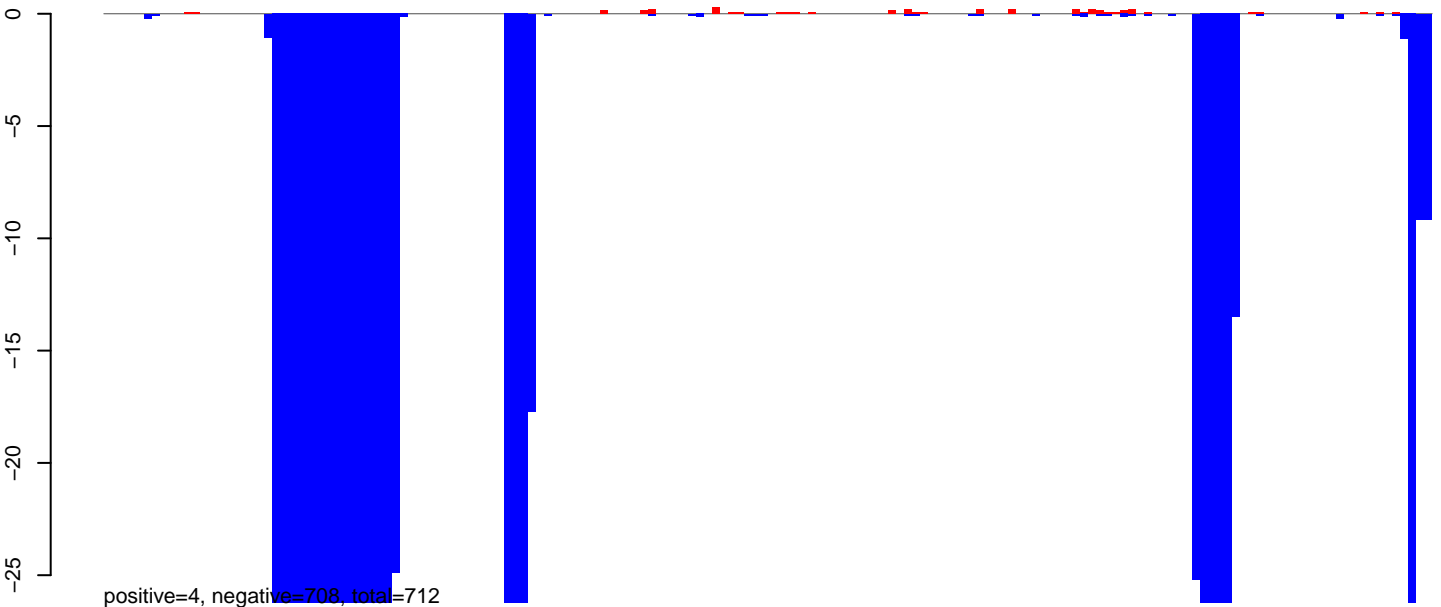
CuQuin_Mcarc_NL.rep



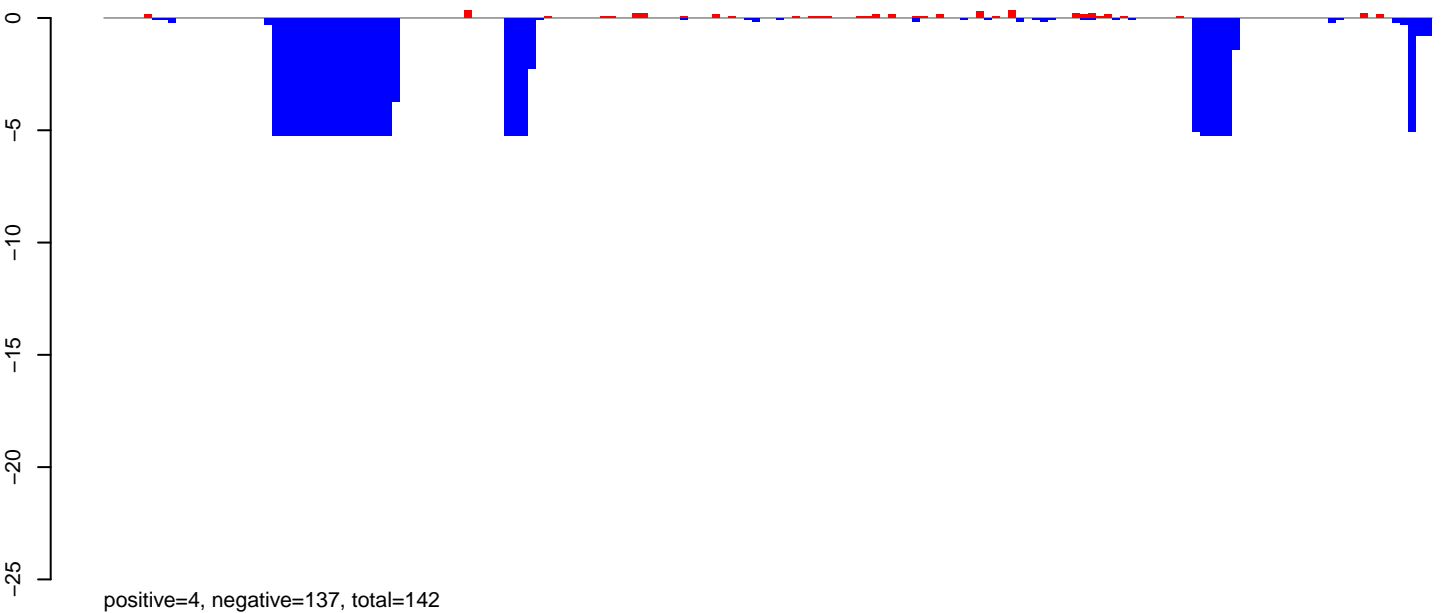
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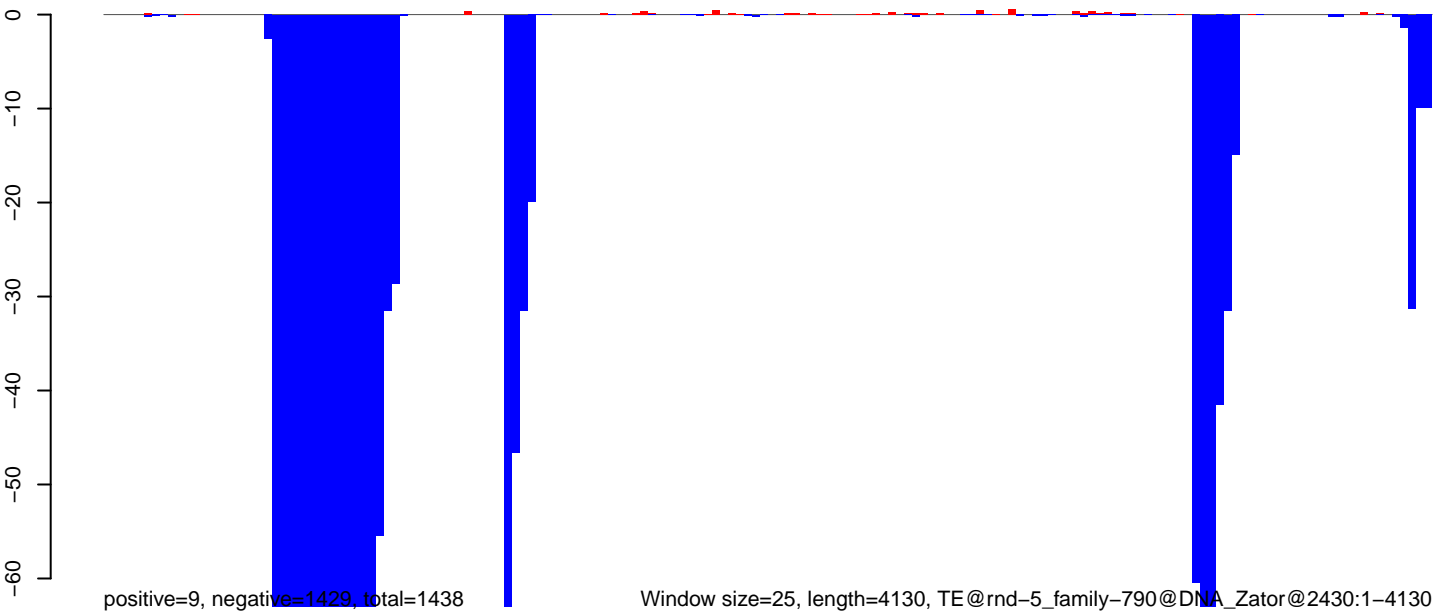
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



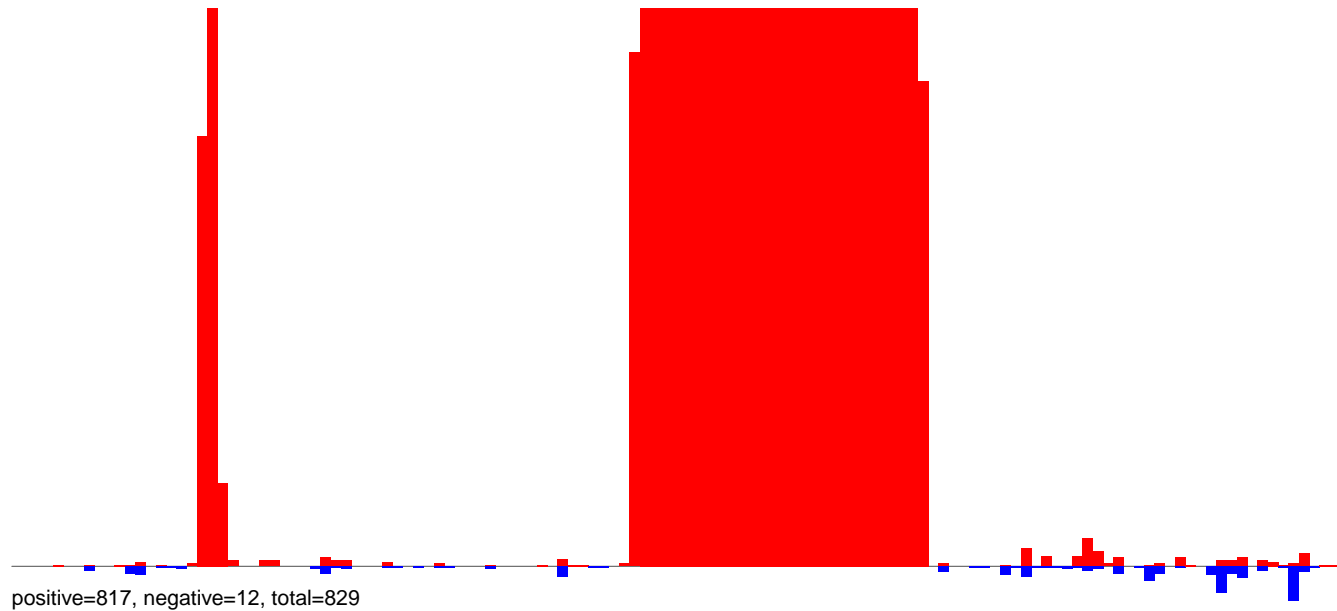
CuQuin_Mcarc_NL.rep



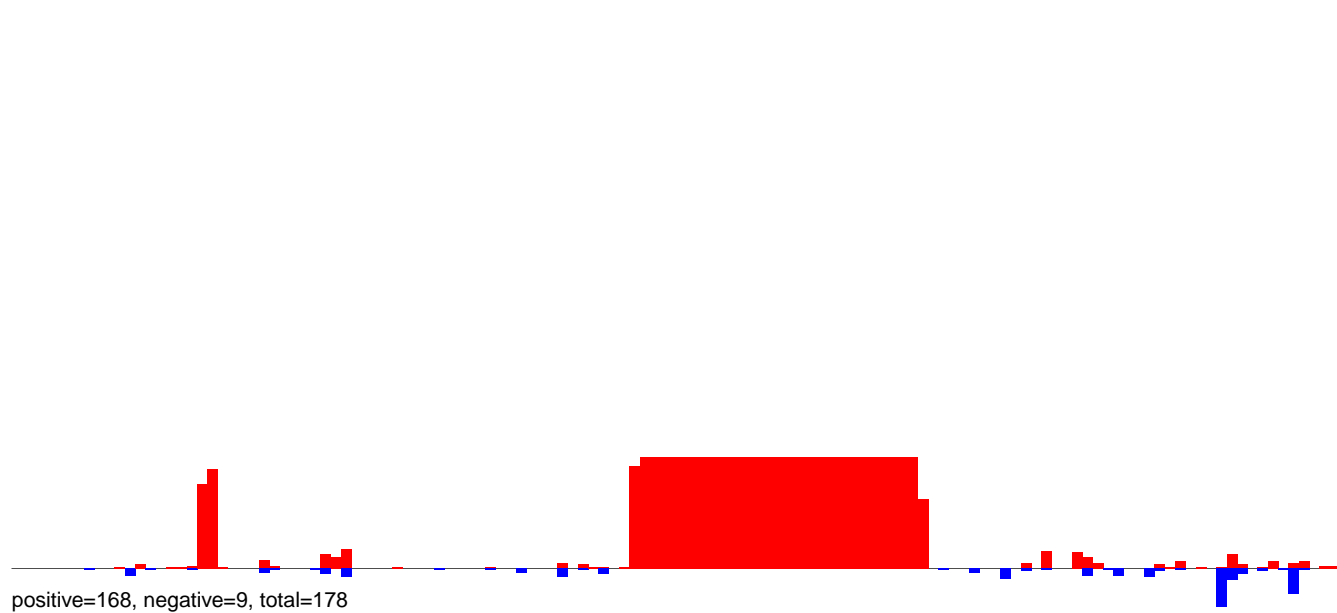
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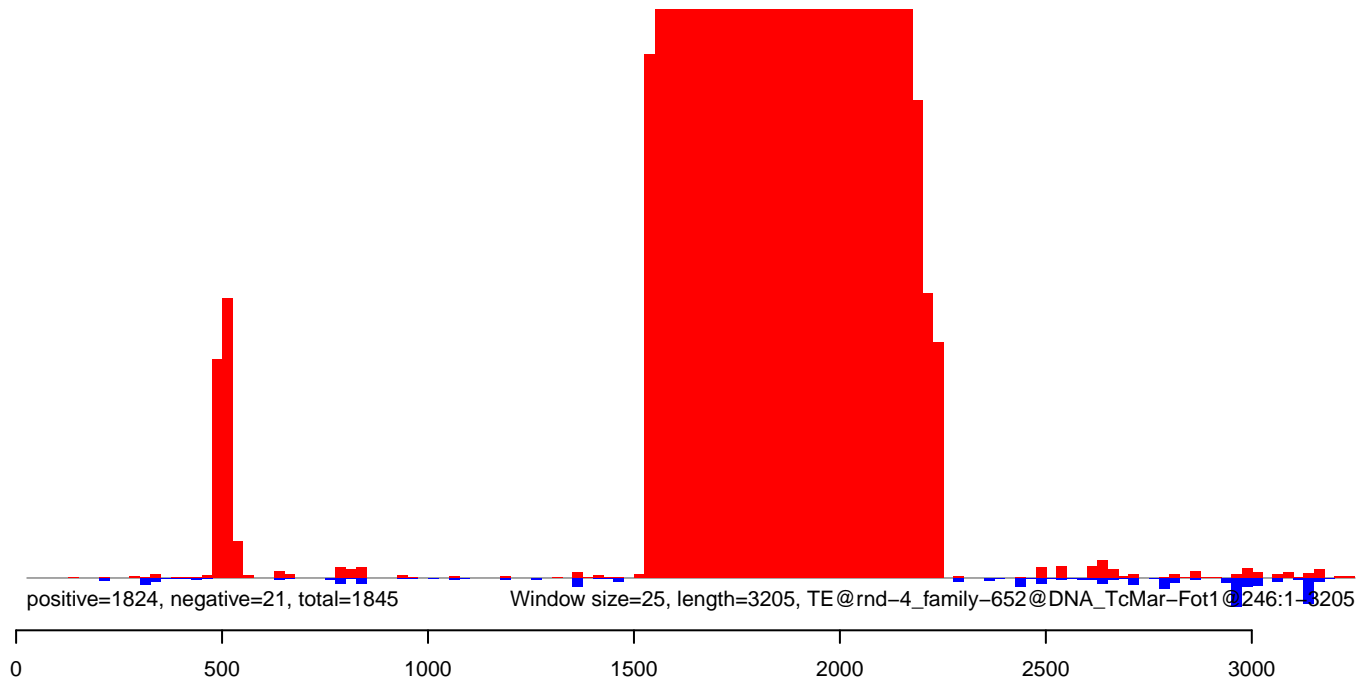
CuQuin_Mcarc_NL.18_23.rep



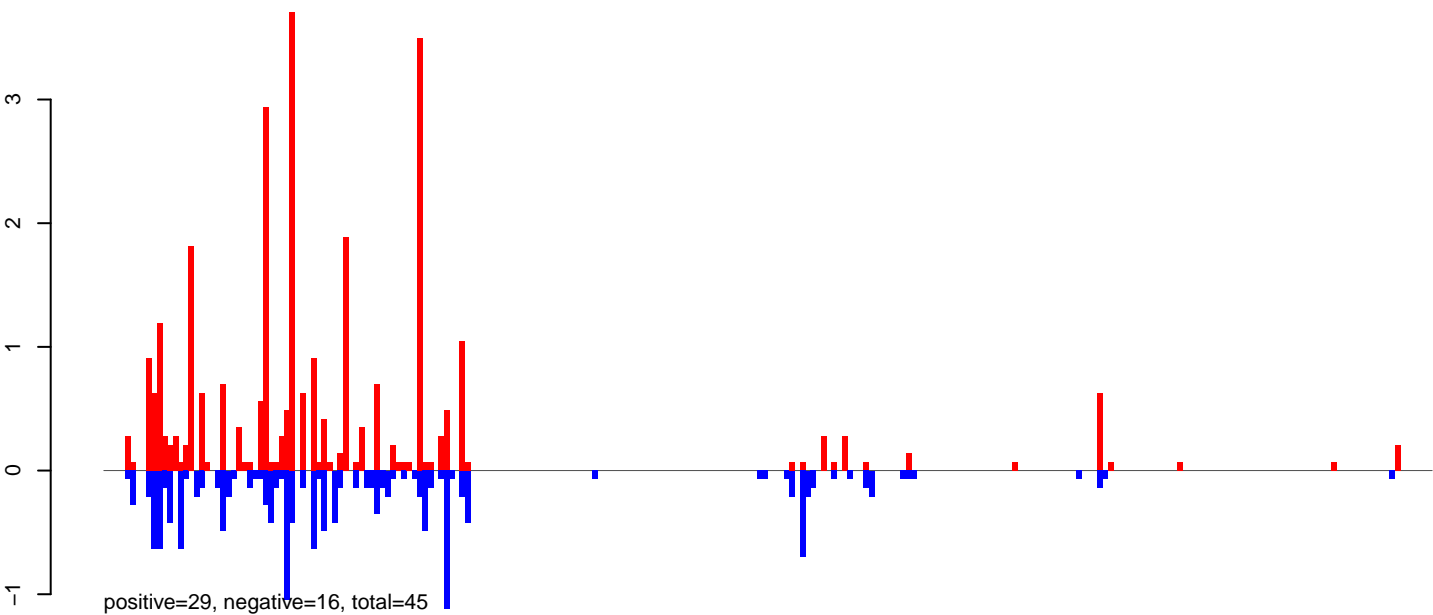
CuQuin_Mcarc_NL.24_35.rep



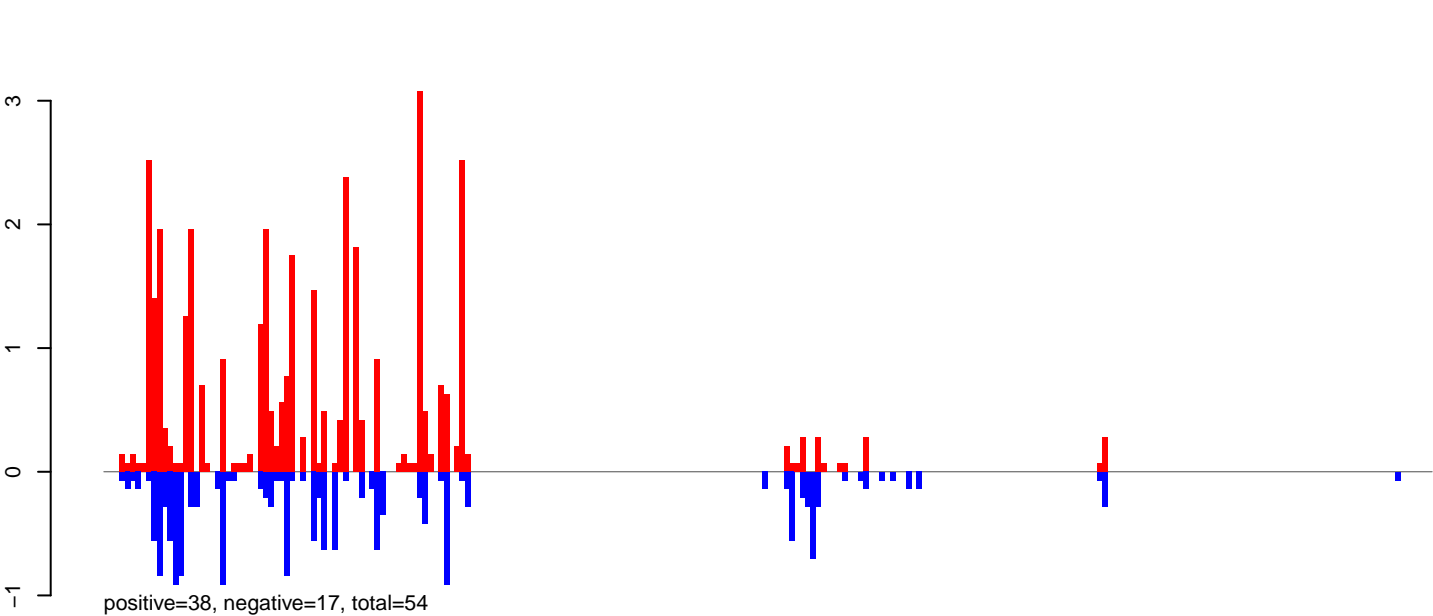
CuQuin_Mcarc_NL.rep



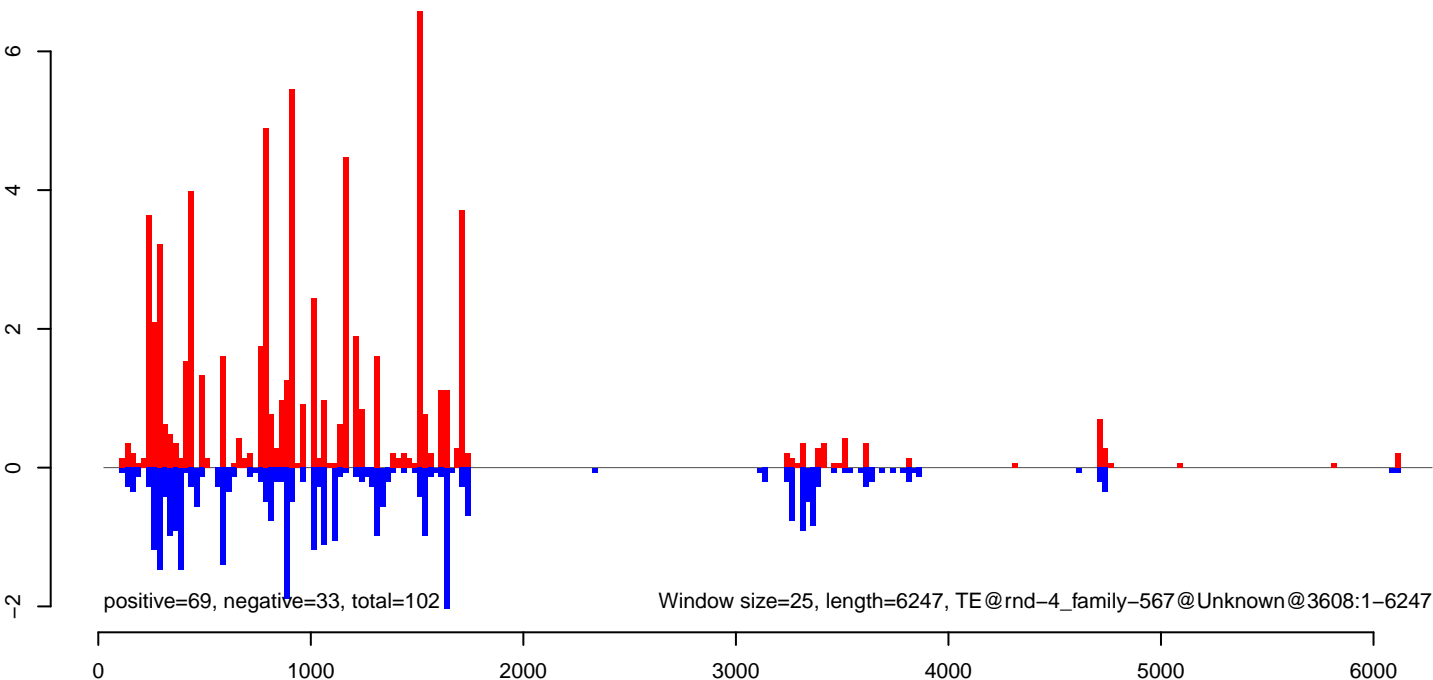
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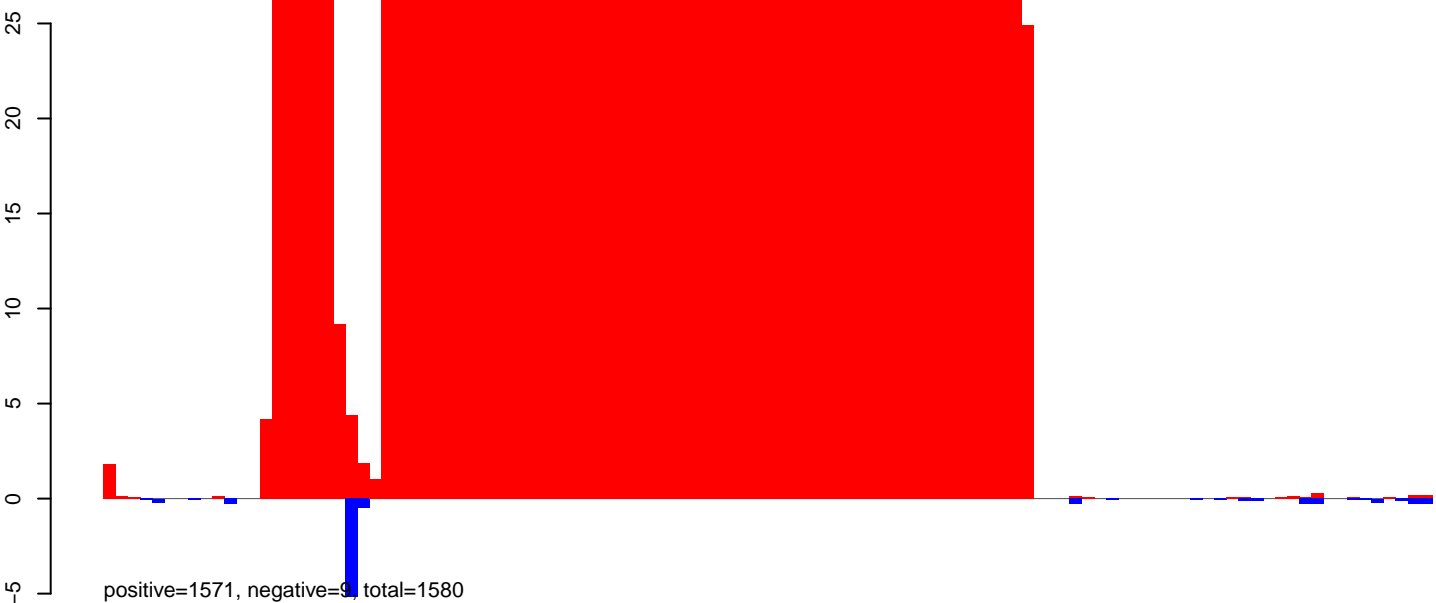
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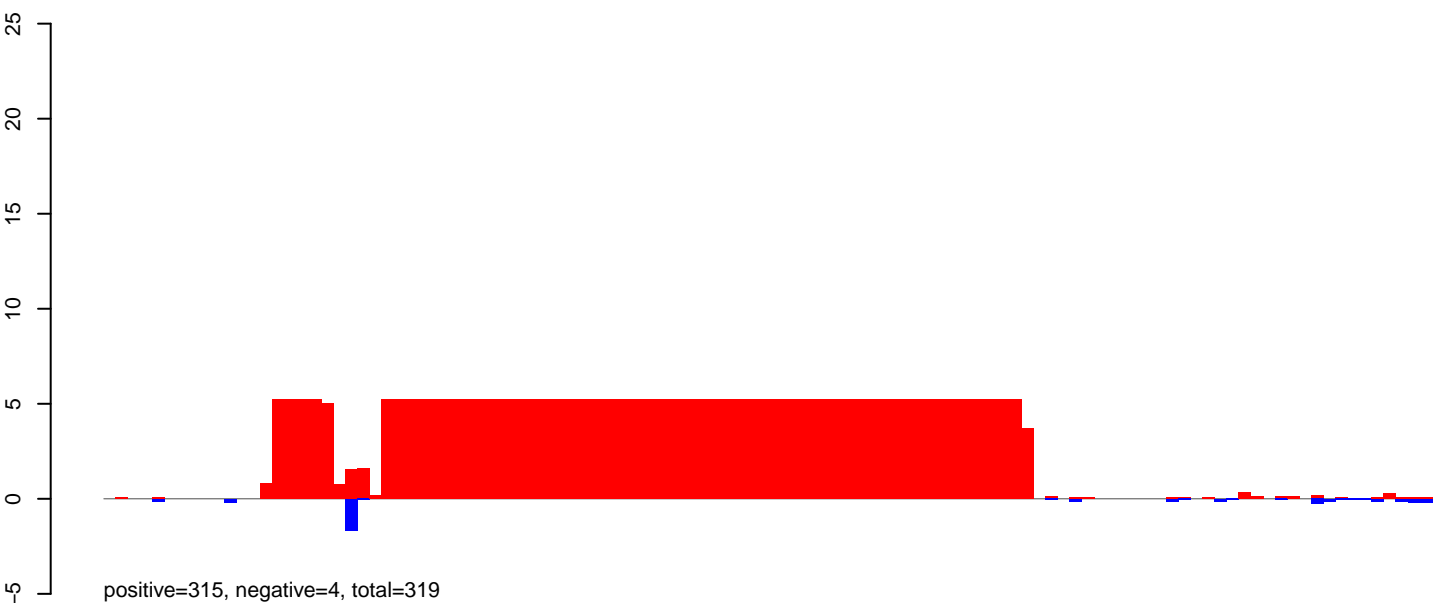
CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep



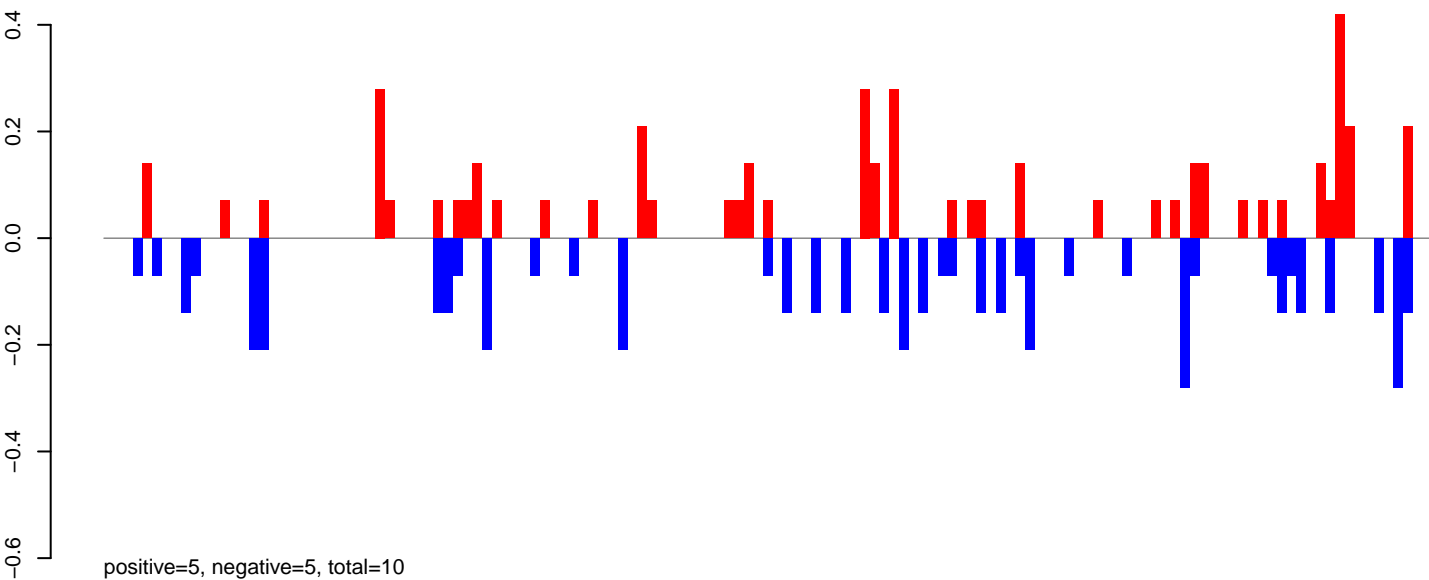
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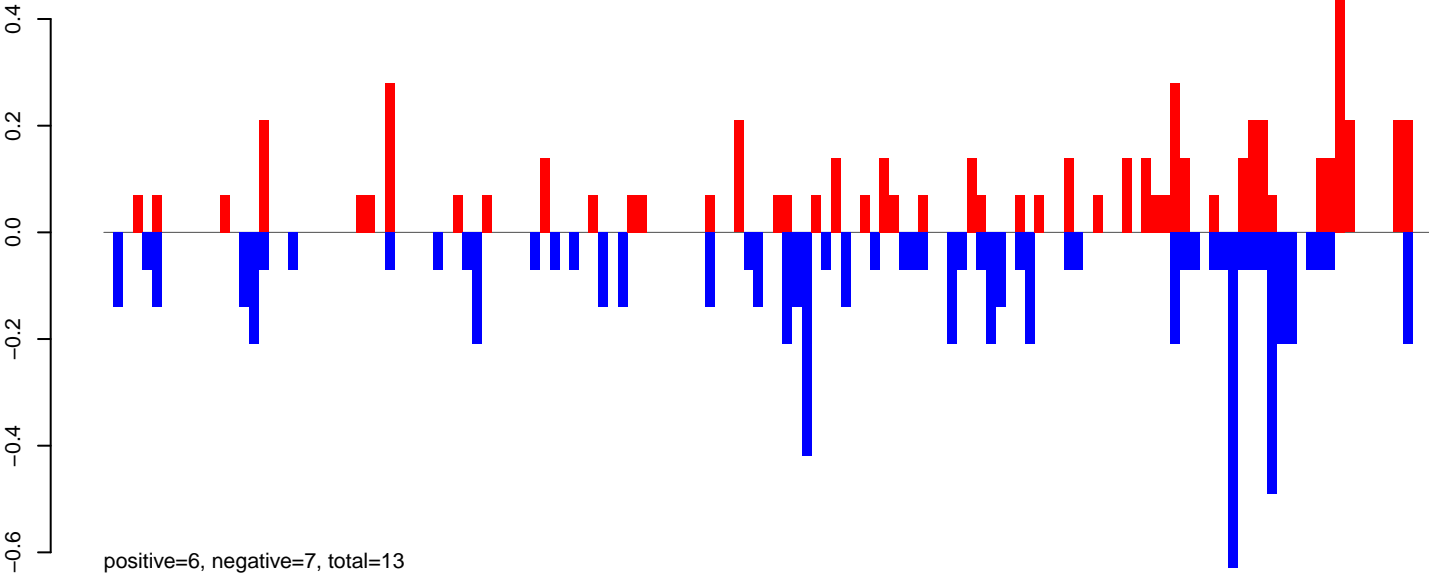
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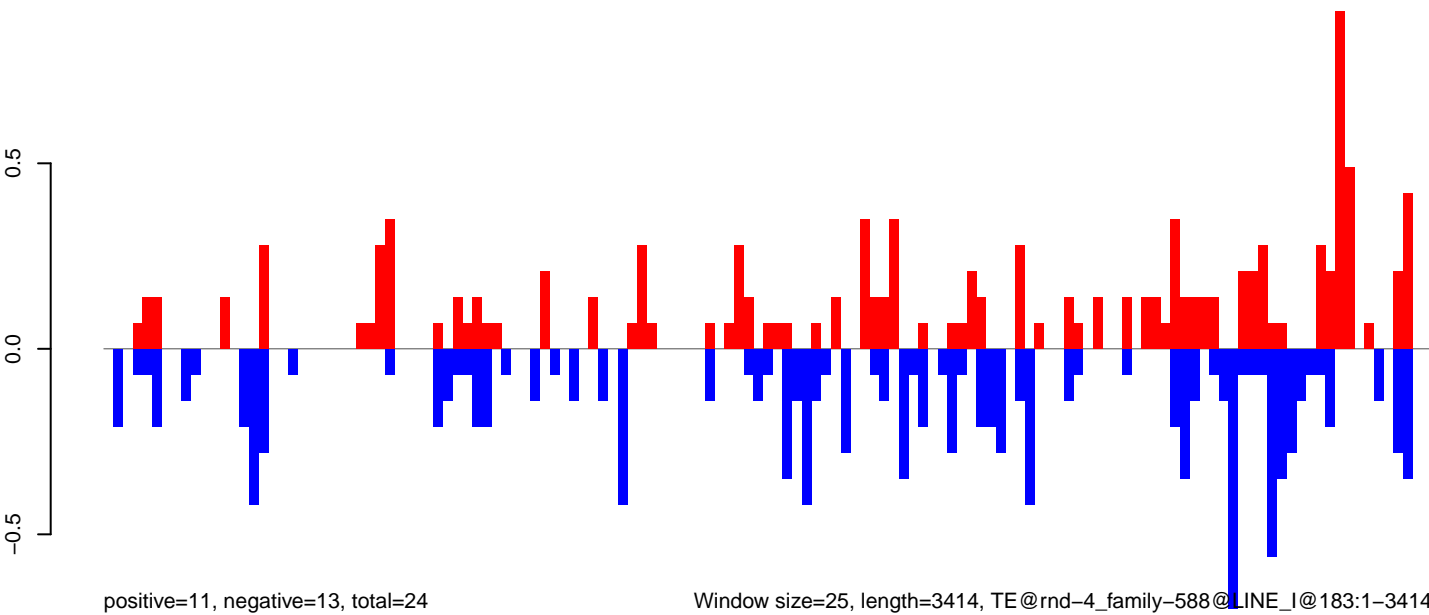
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CuQuin_Mcarc_NL.24_35.rep



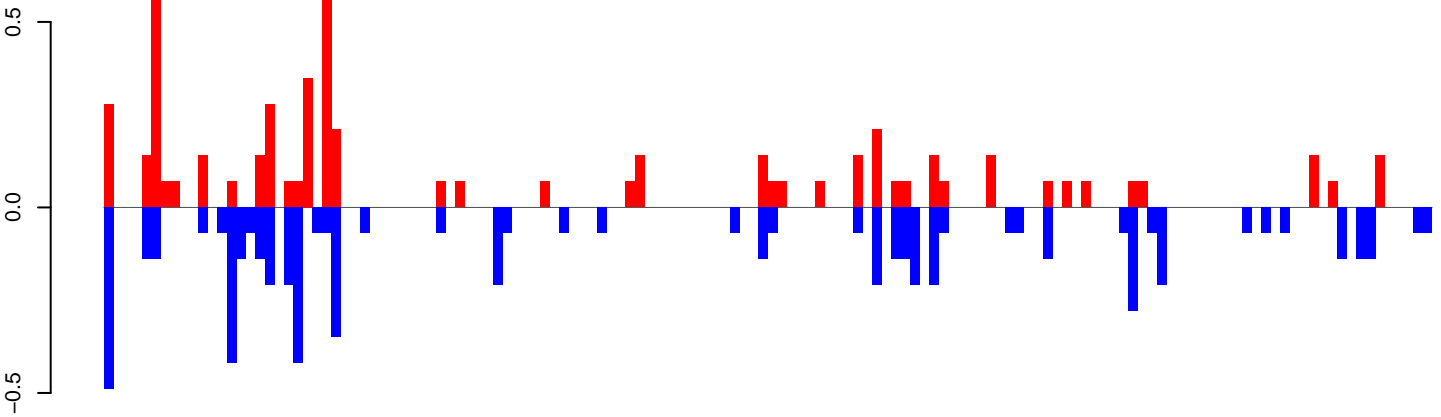
CuQuin_Mcarc_NL.rep



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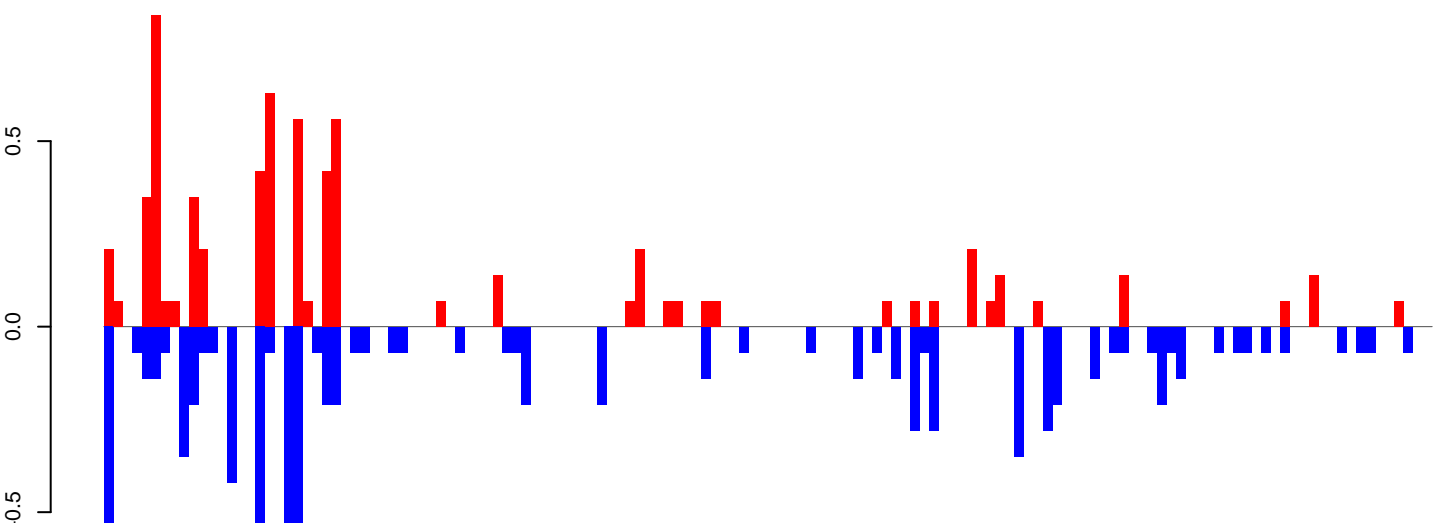
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CuQuin_Mcarc_NL.18_23.rep



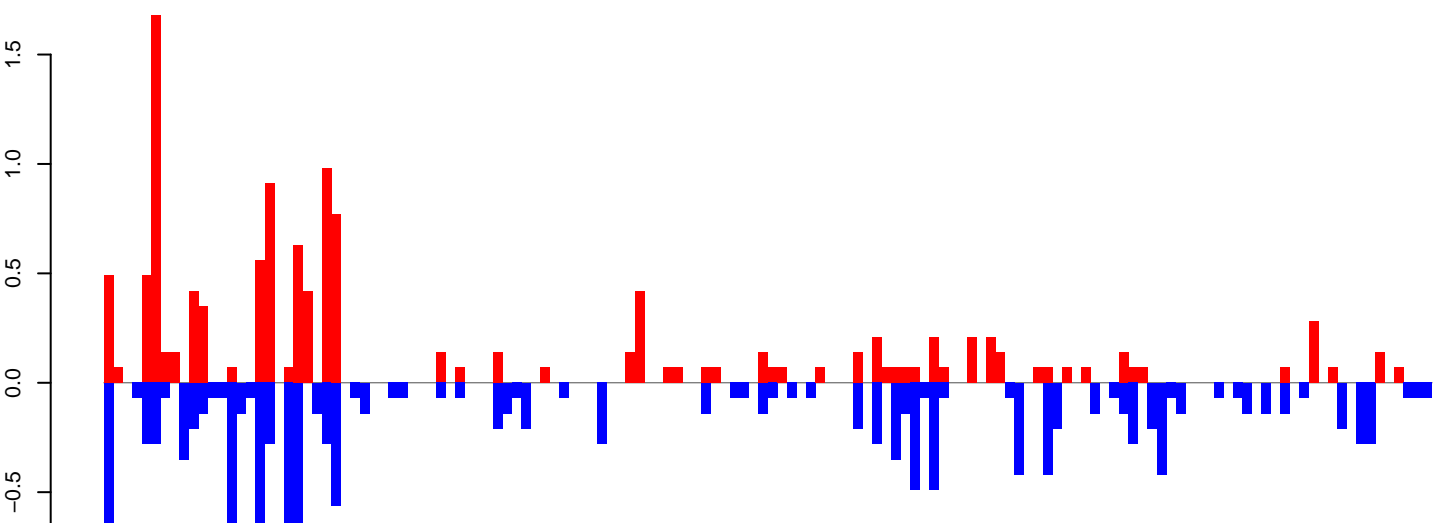
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CuQuin_Mcarc_NL.24_35.rep



positive=7, negative=9, total=16

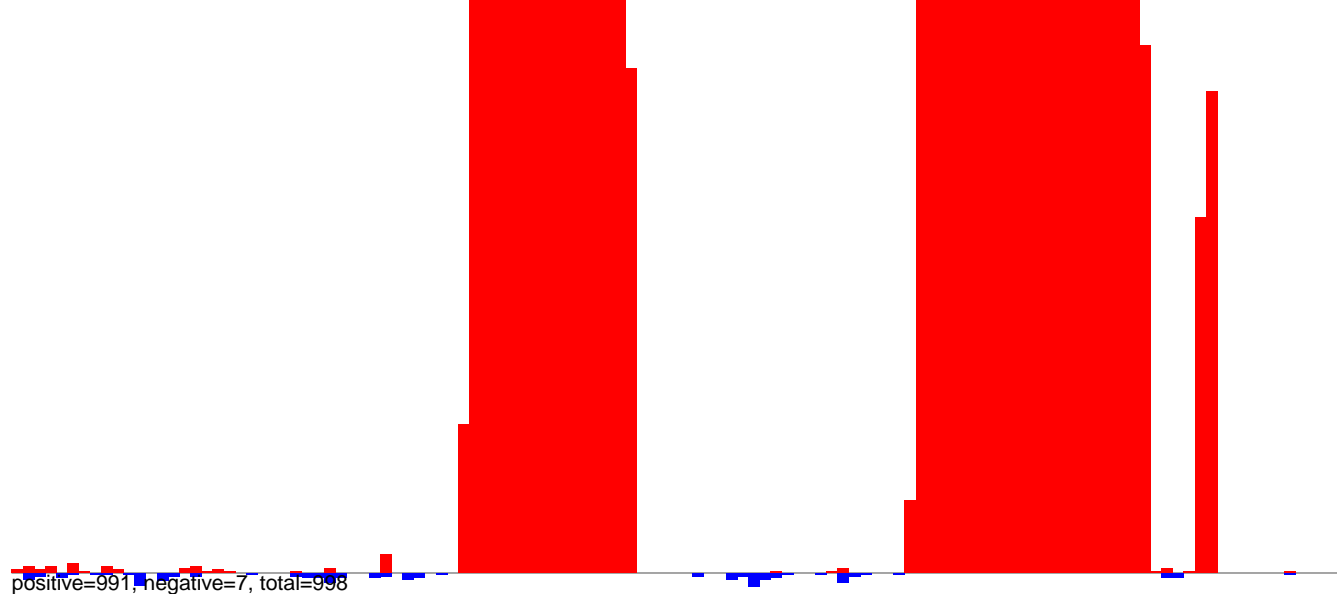
CuQuin_Mcarc_NL.rep



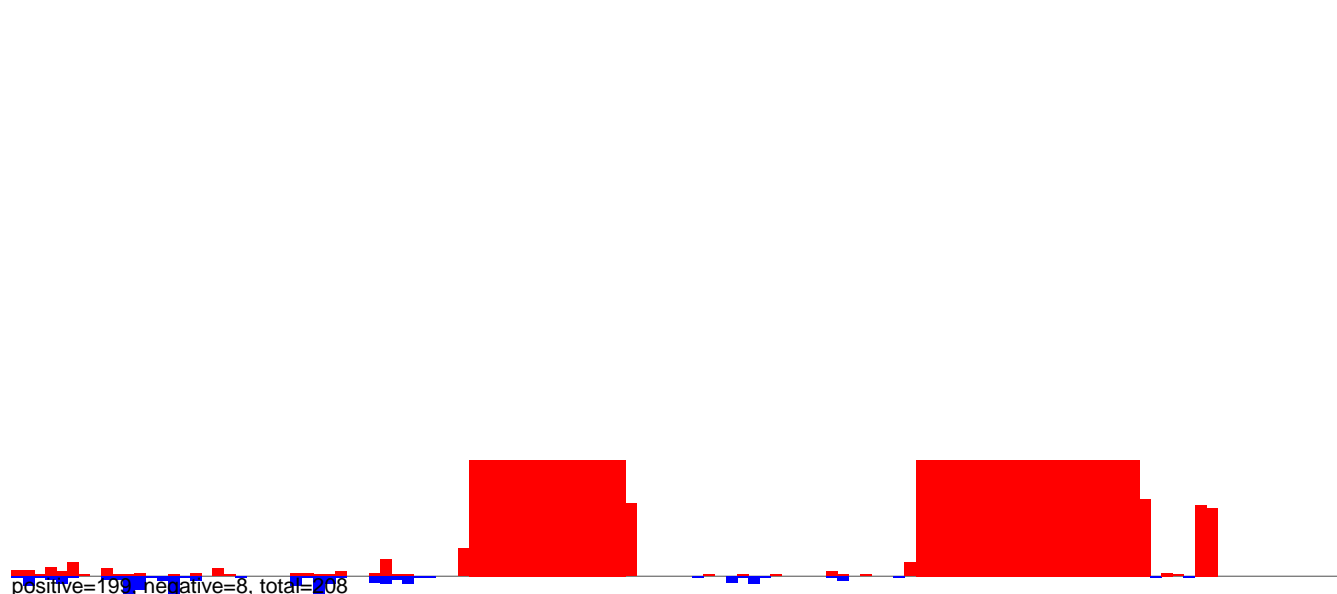
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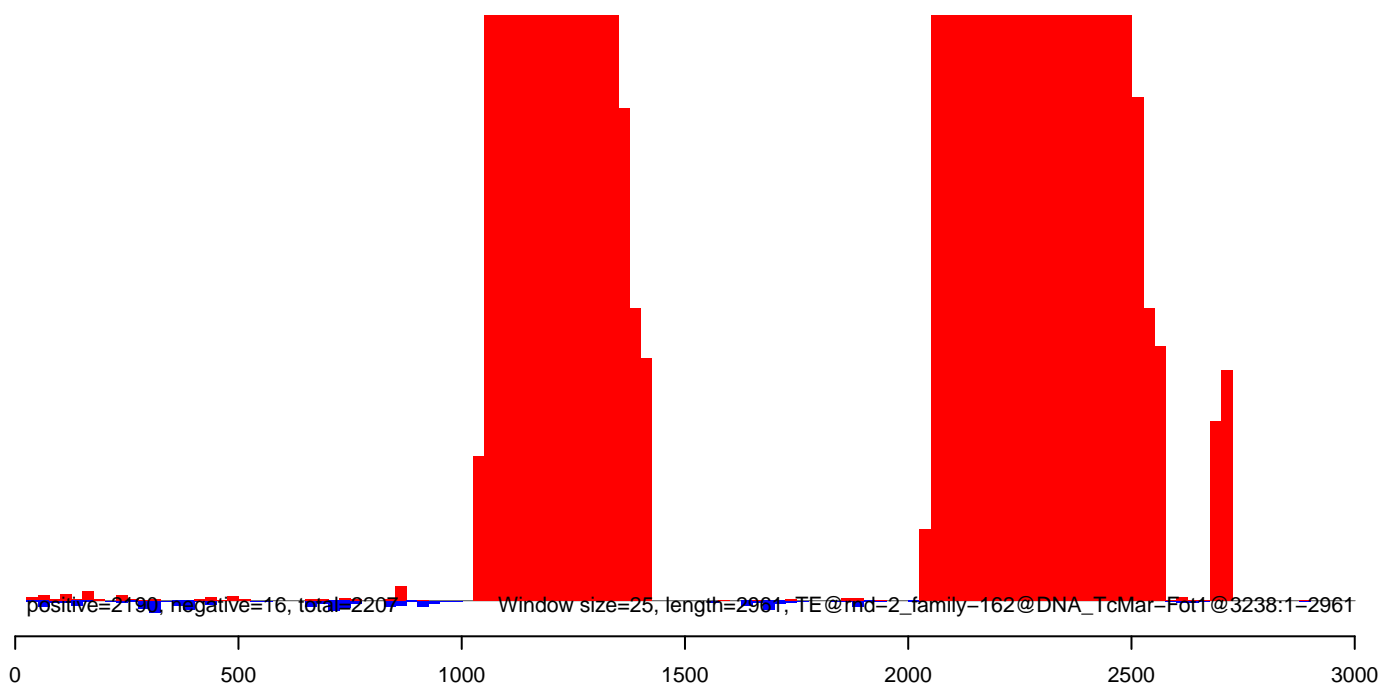
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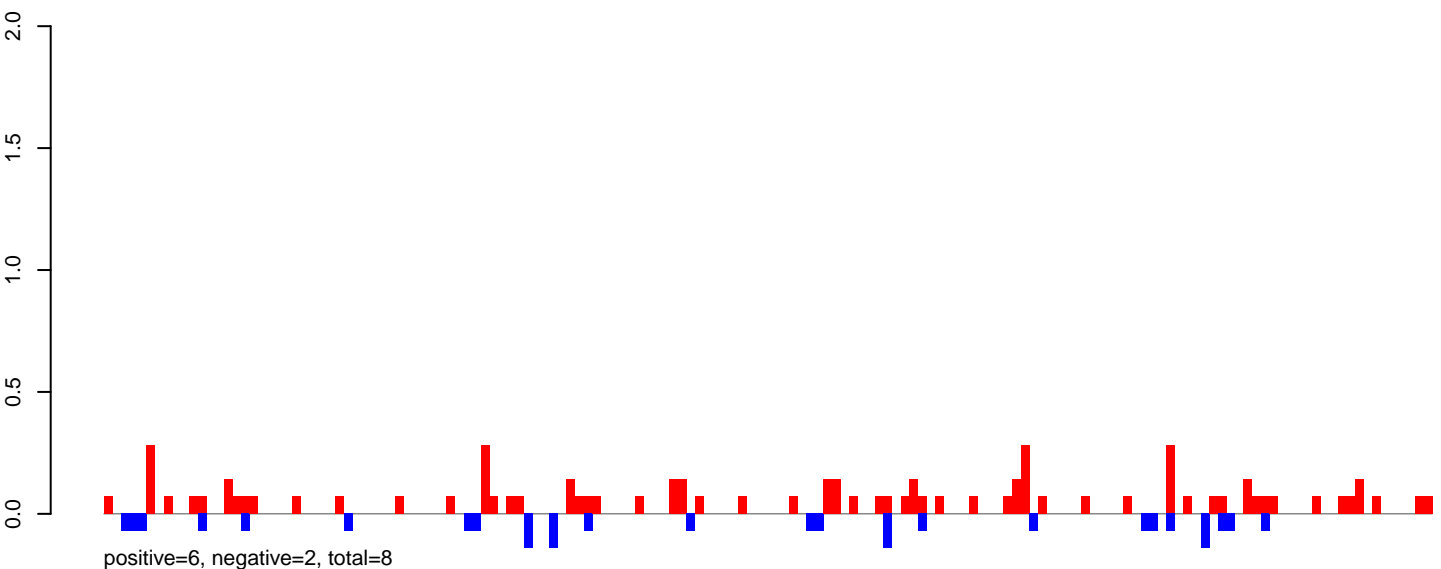
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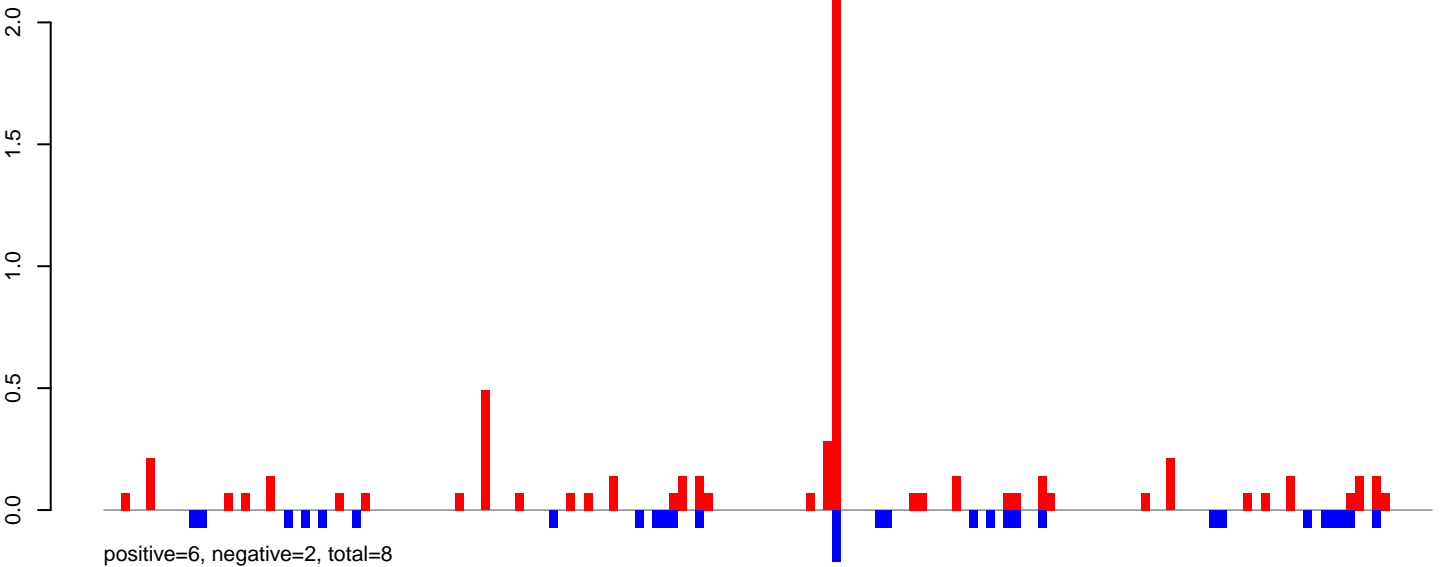
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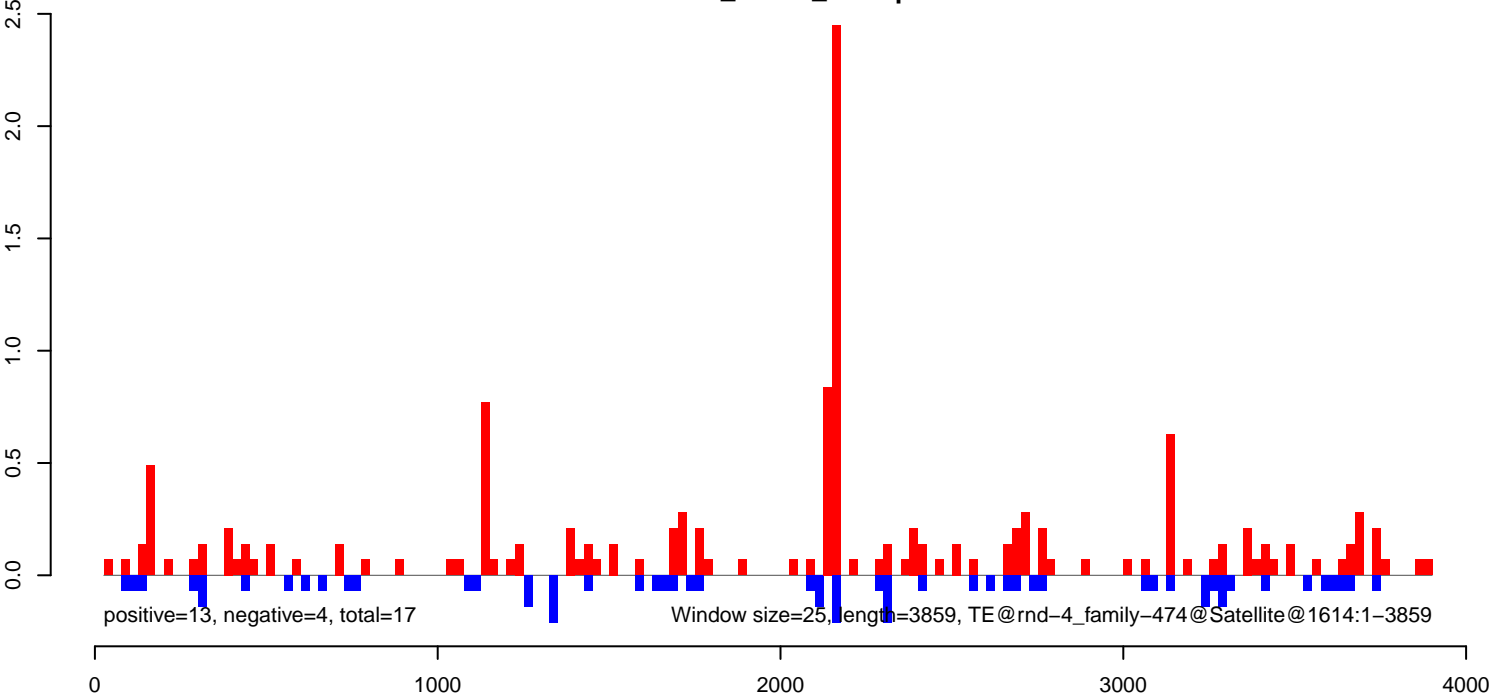
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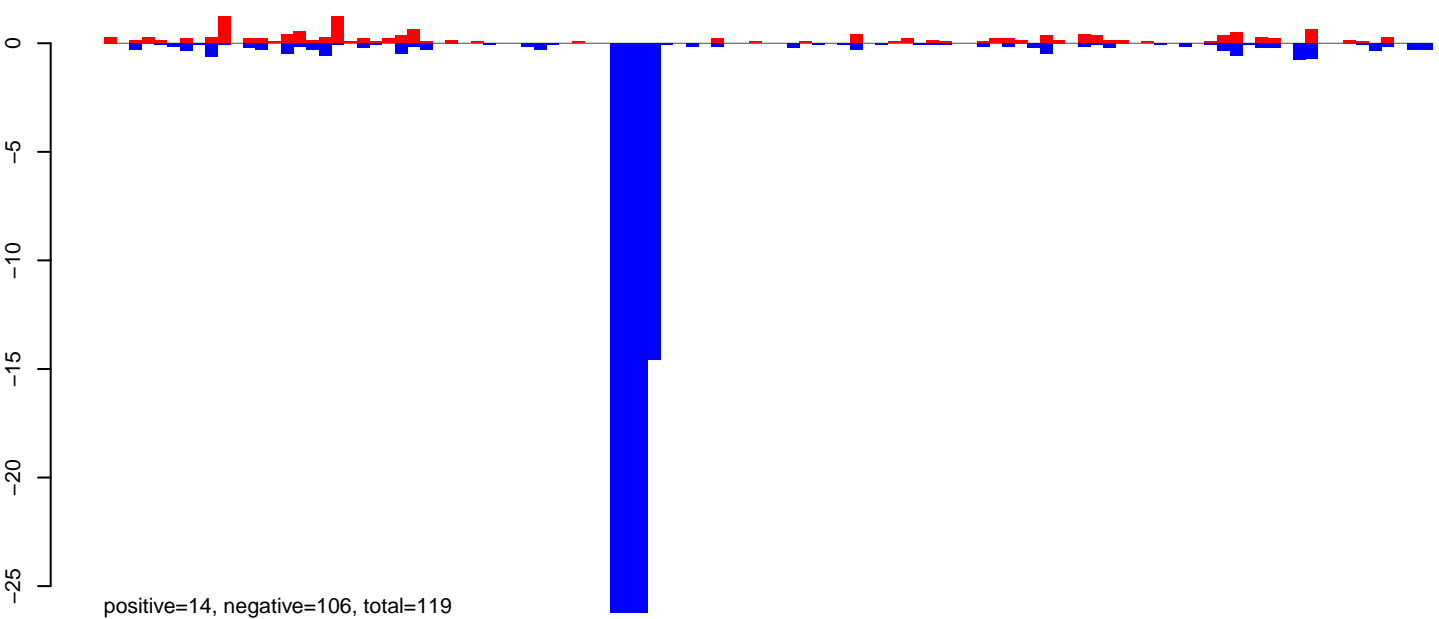
CuQuin_Mcarc_NL.24_35.rep



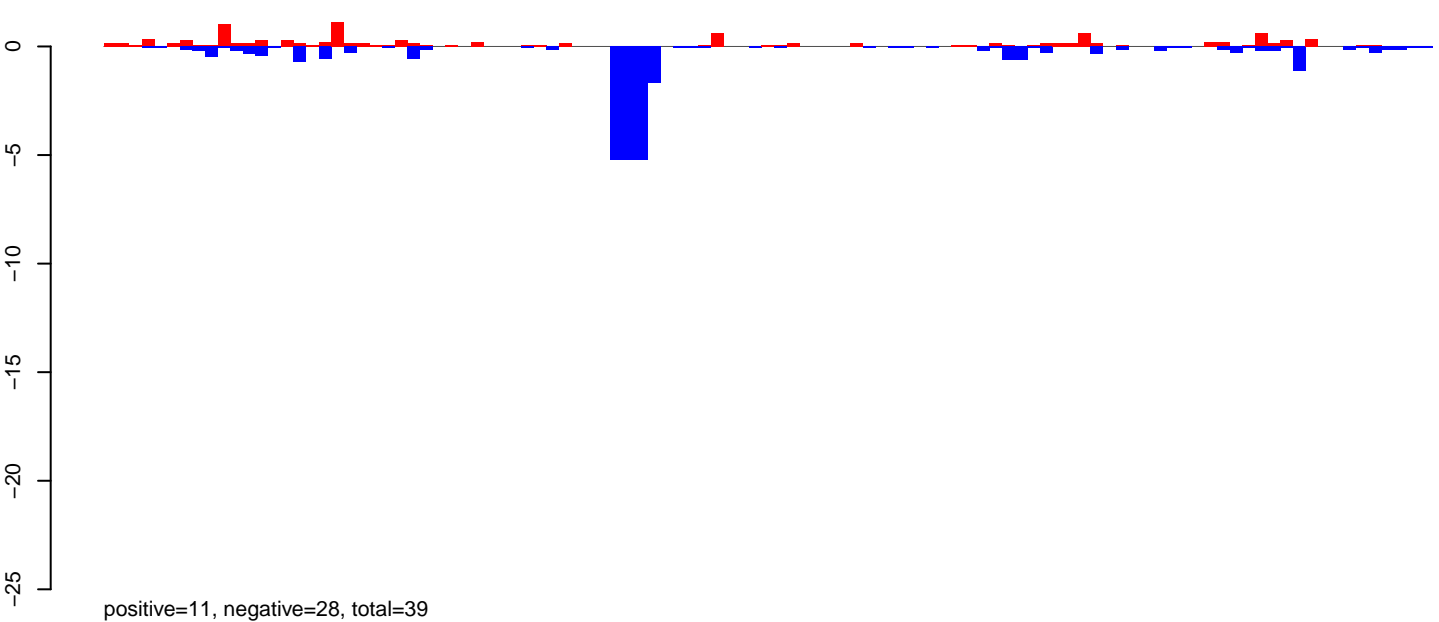
CuQuin_Mcarc_NL.rep



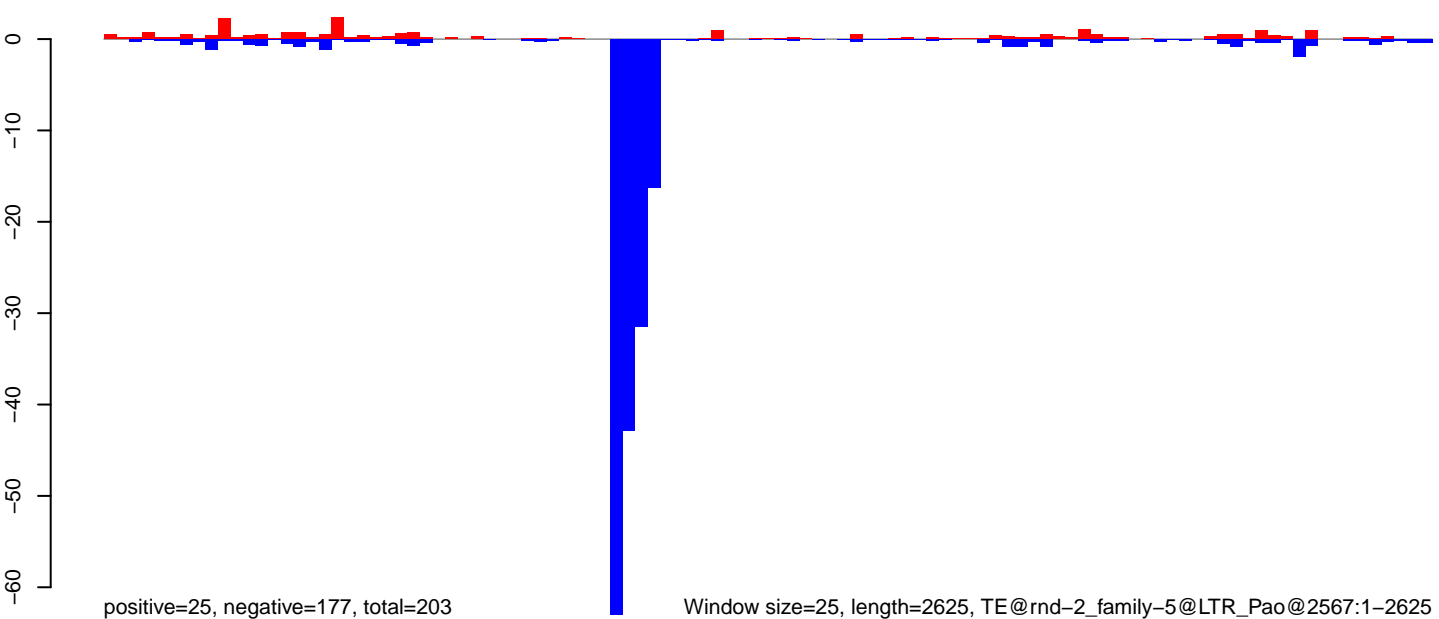
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CuQuin_Mcarc_NL.24_35.rep

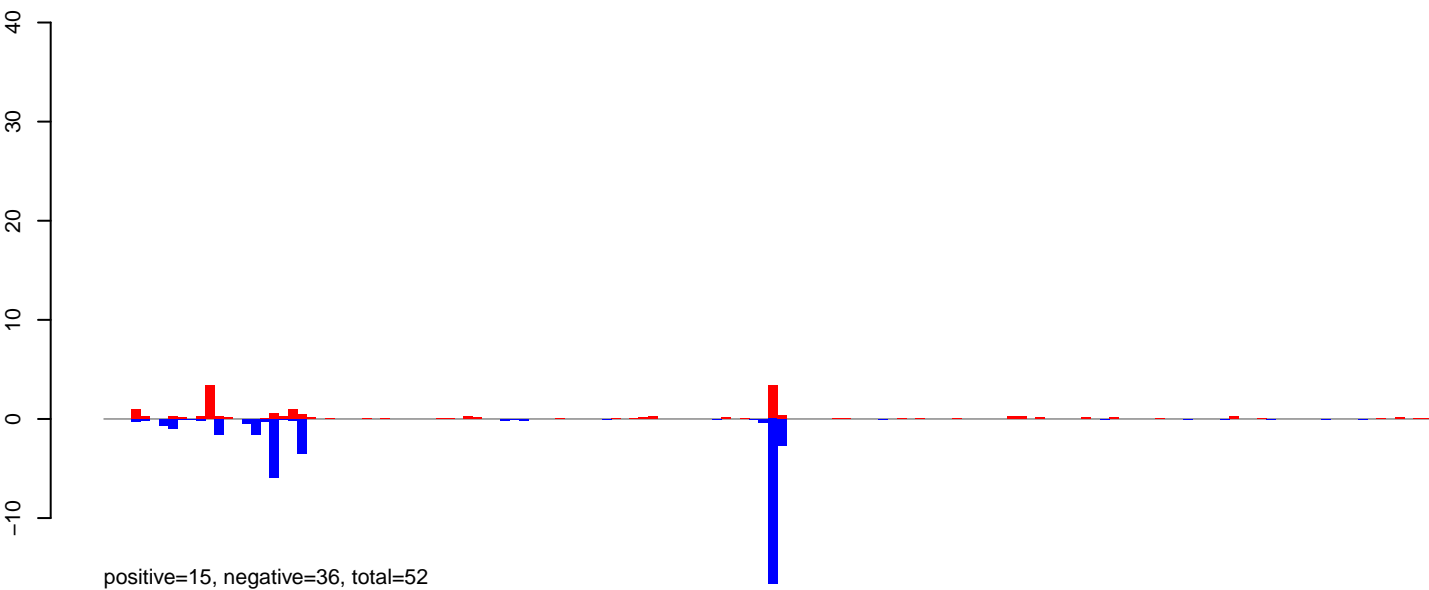


CuQuin_Mcarc_NL.rep

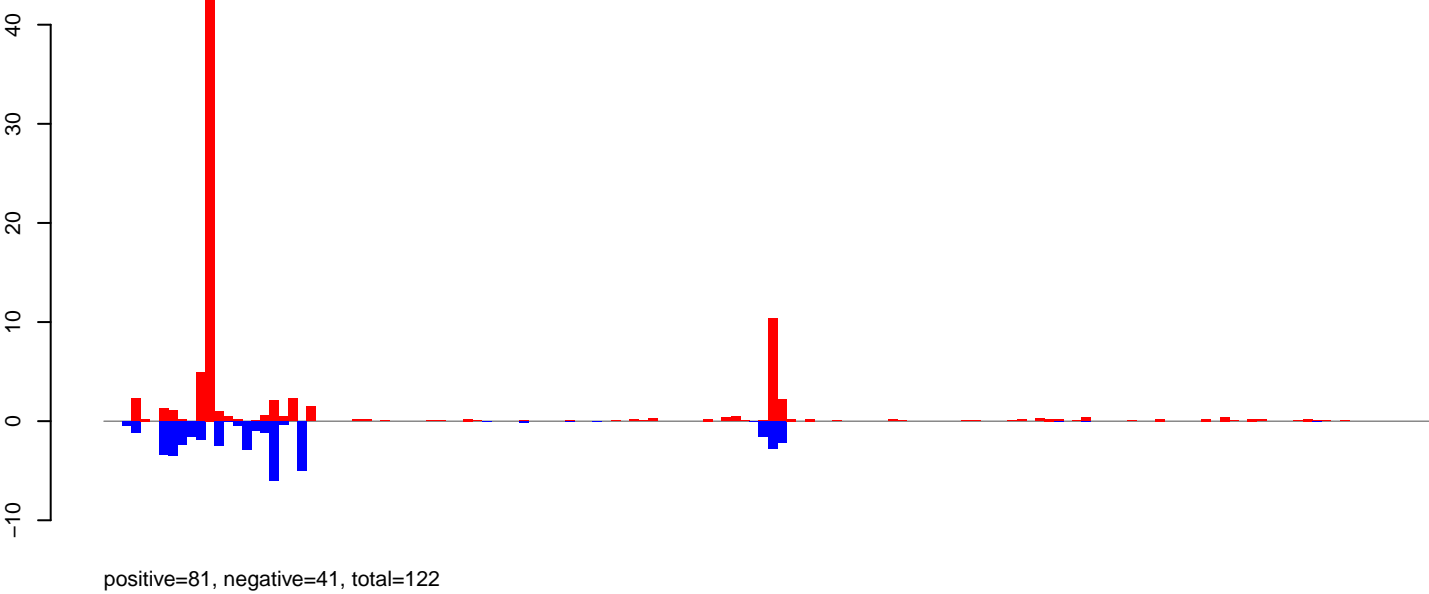


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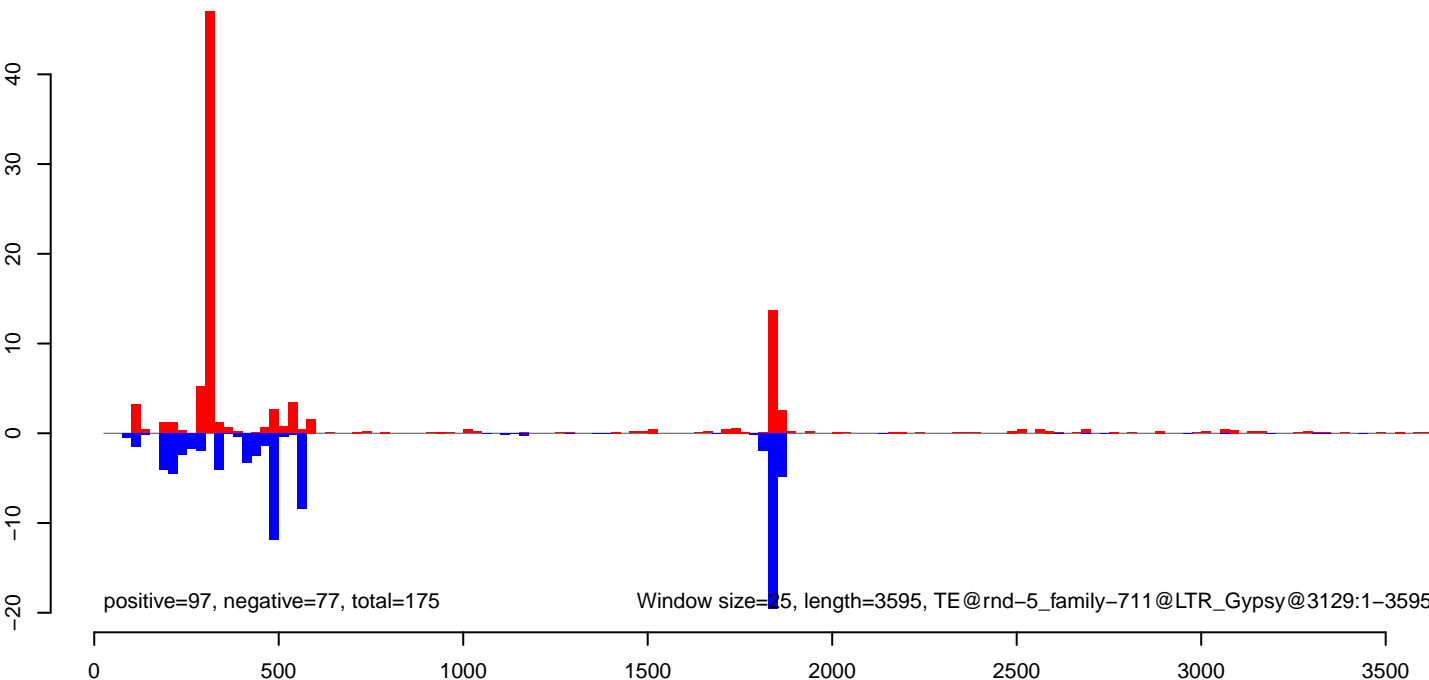
CuQuin_Mcarc_NL.18_23.rep



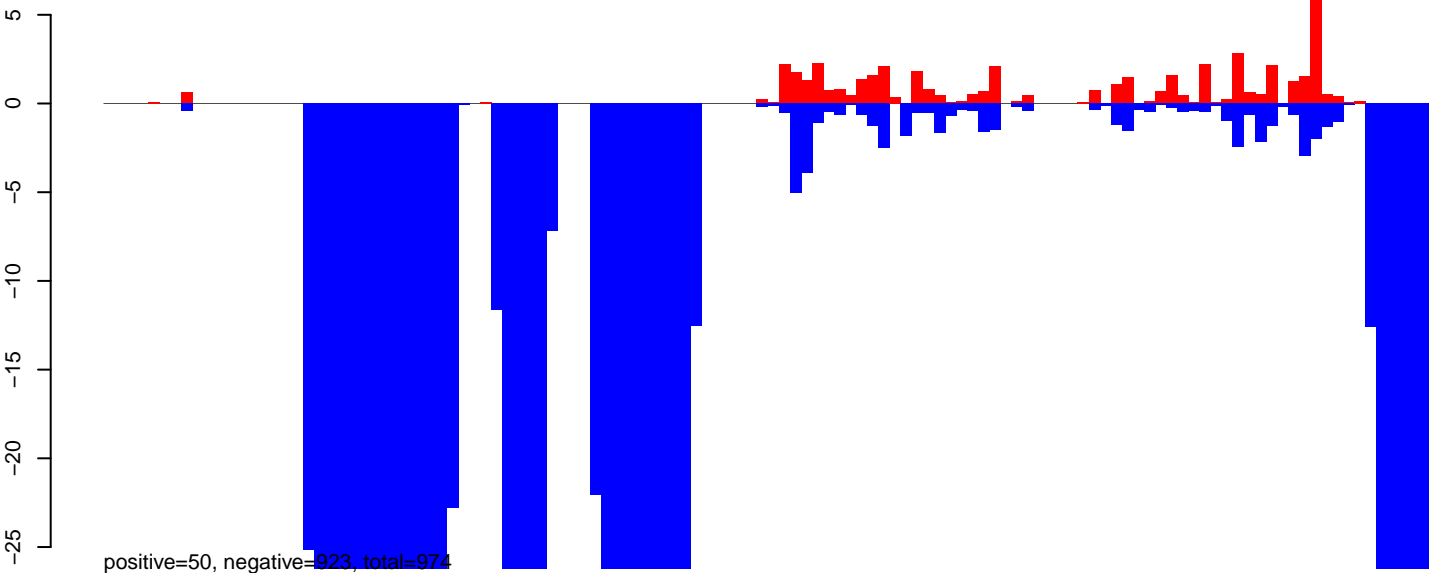
CuQuin_Mcarc_NL.24_35.rep



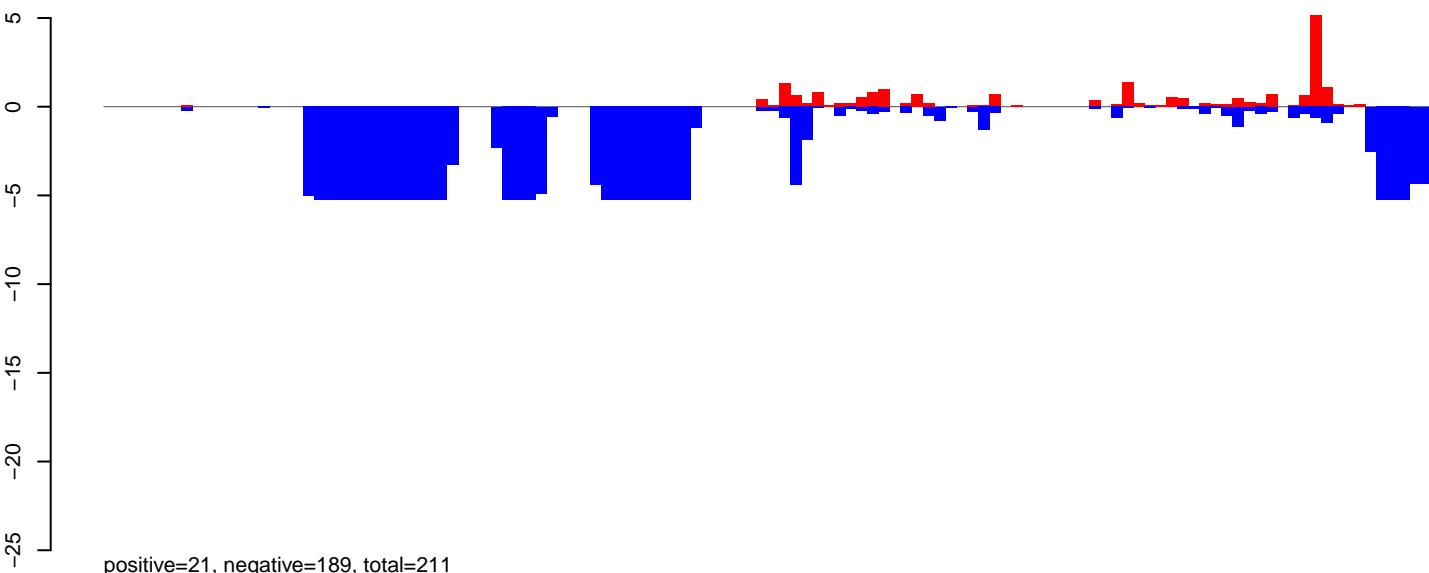
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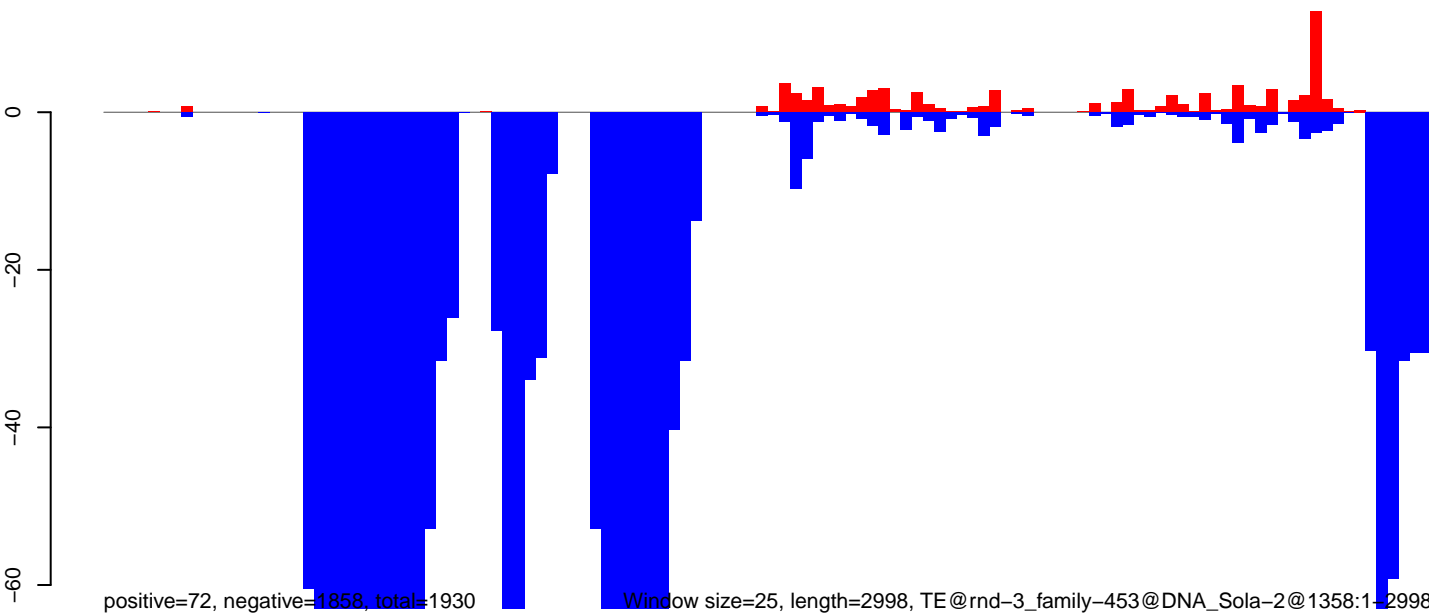
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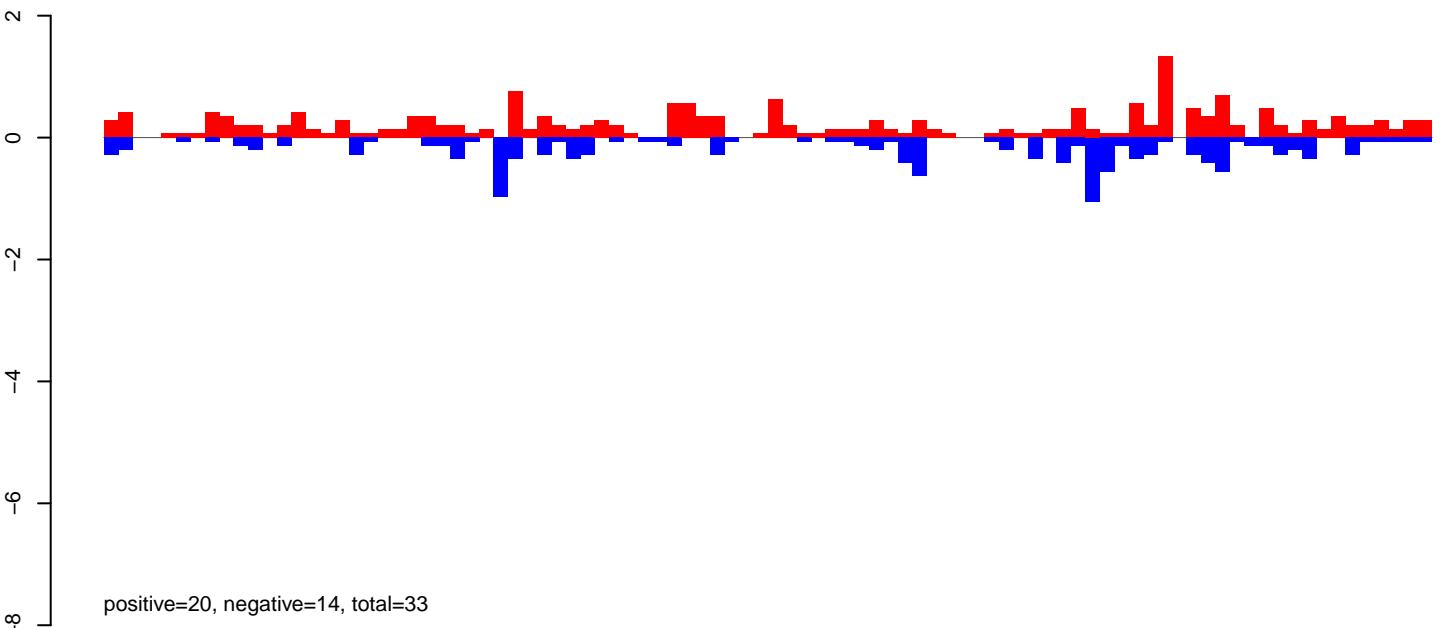
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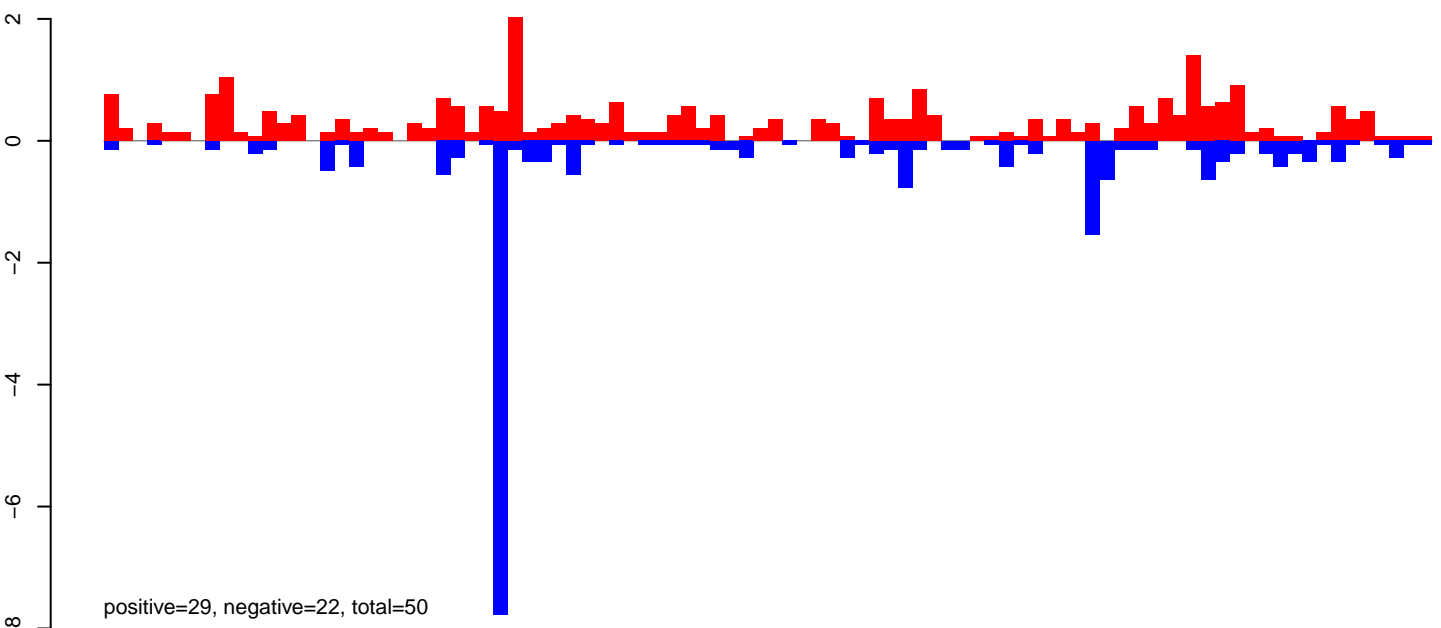
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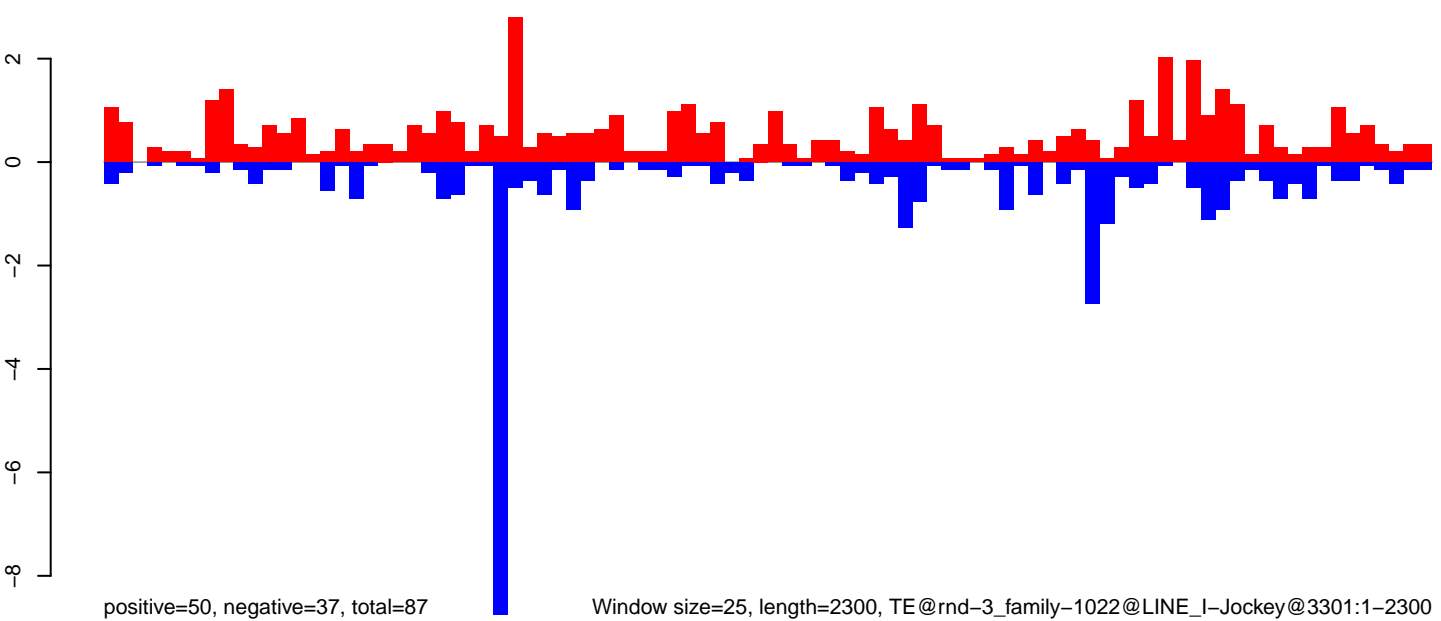
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CuQuin_Mcarc_NL.24_35.rep

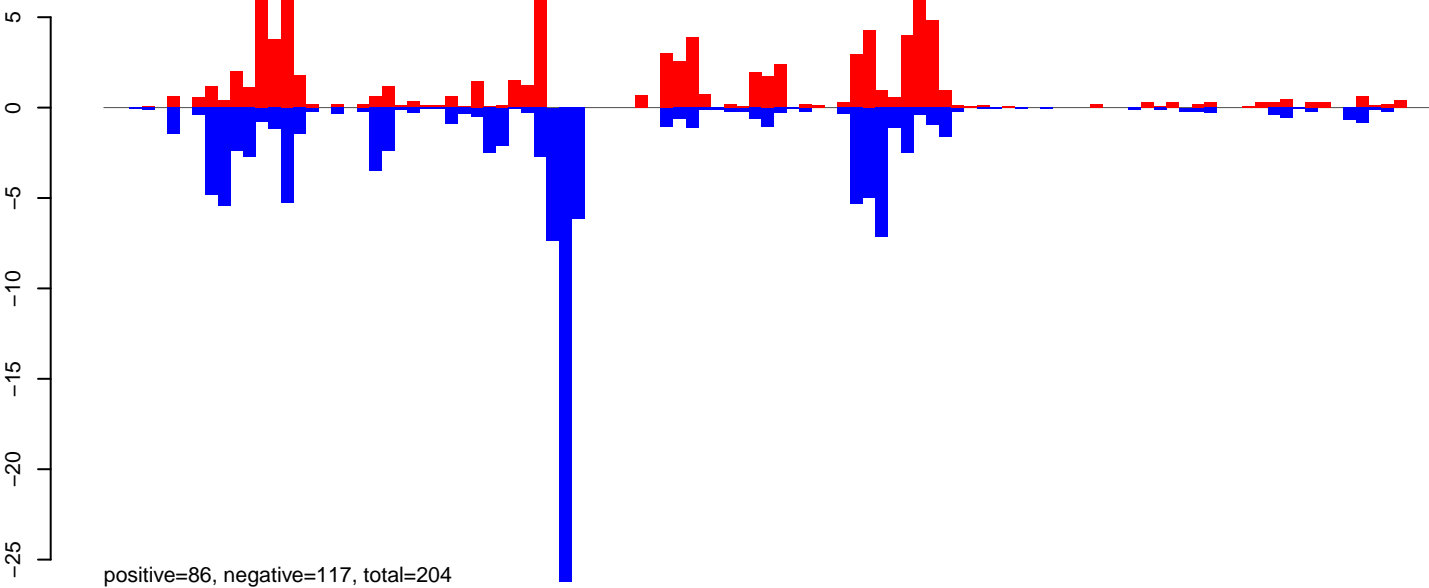


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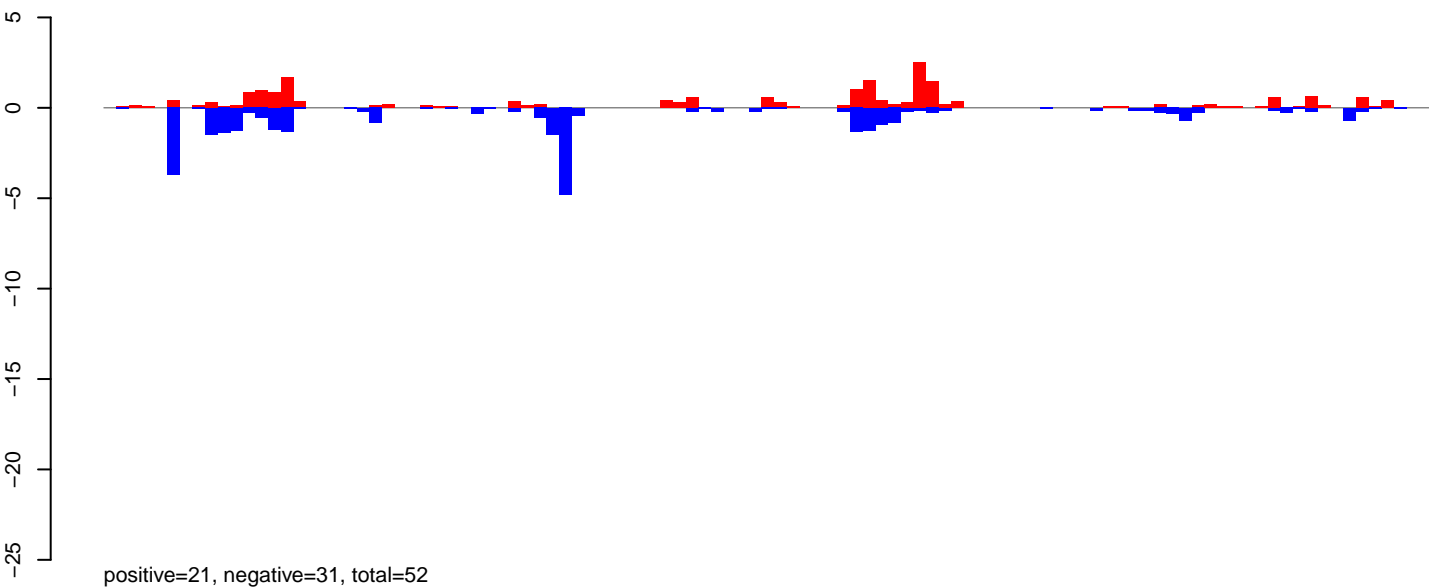


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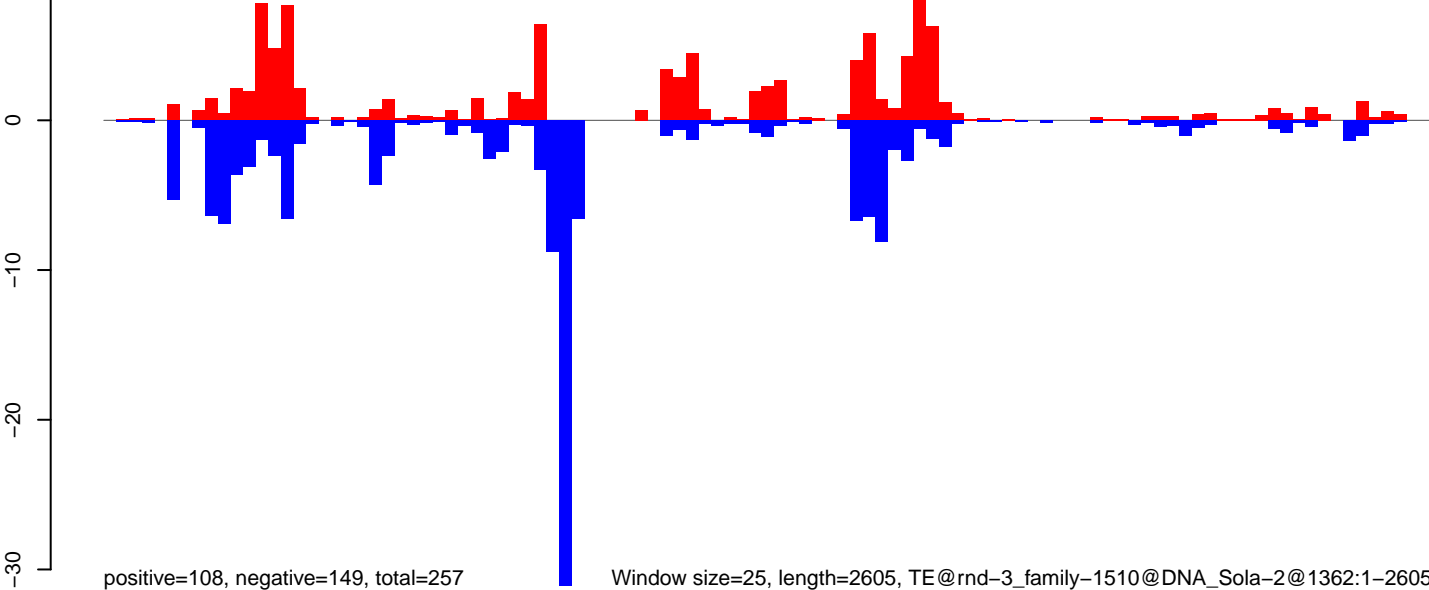
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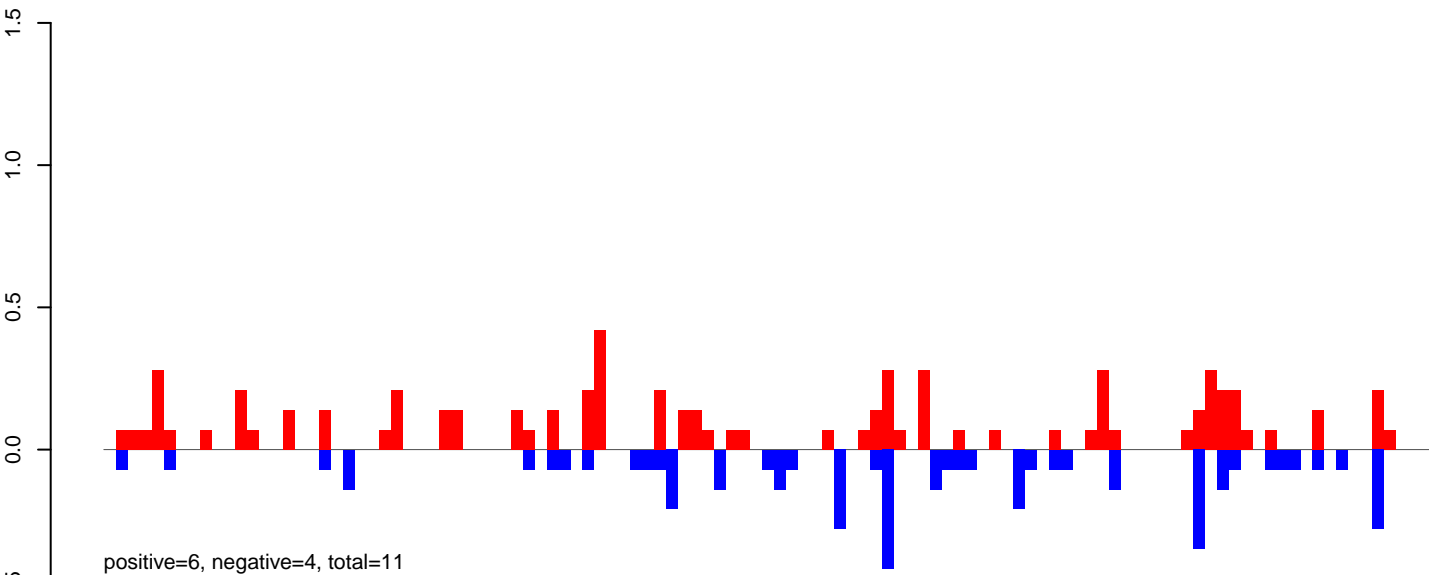
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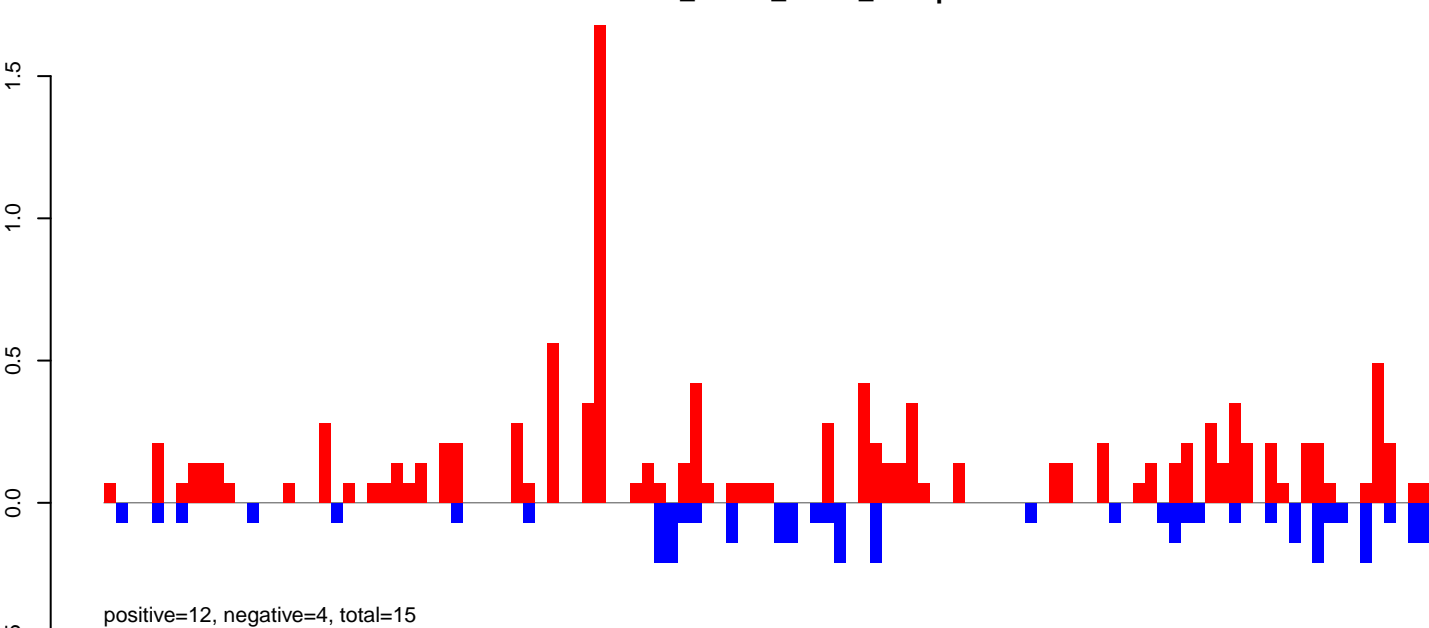
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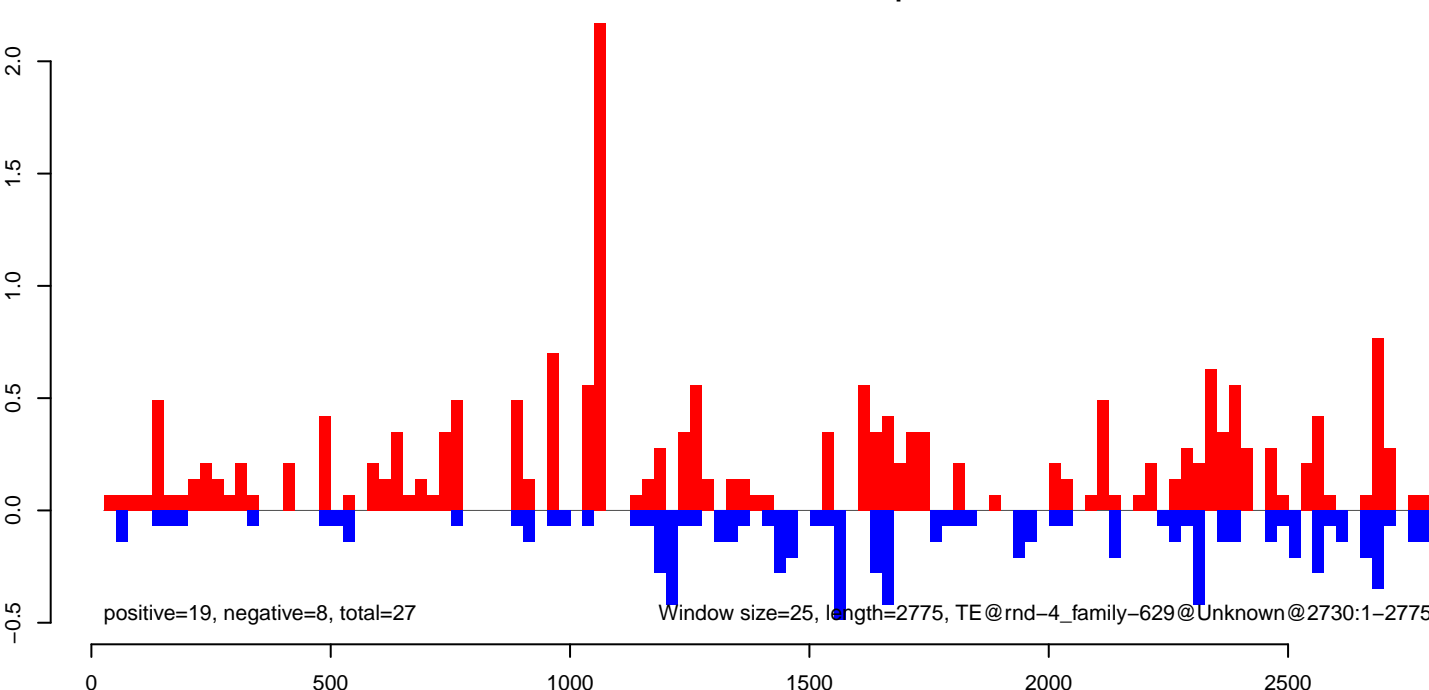
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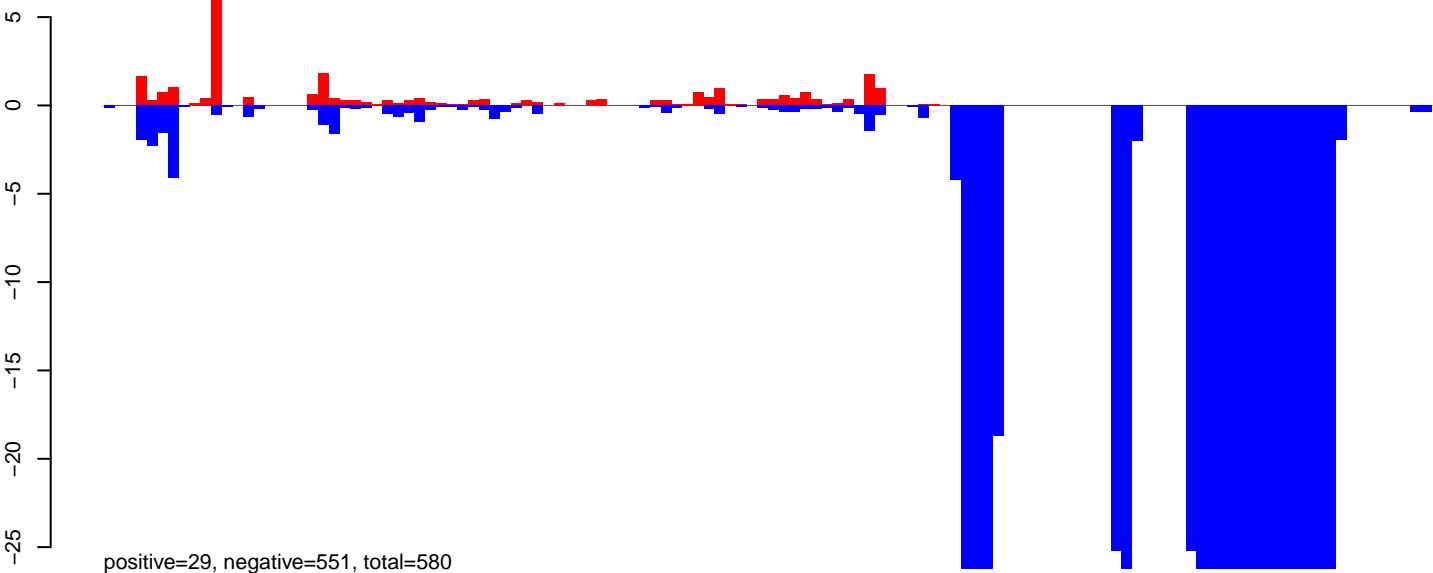
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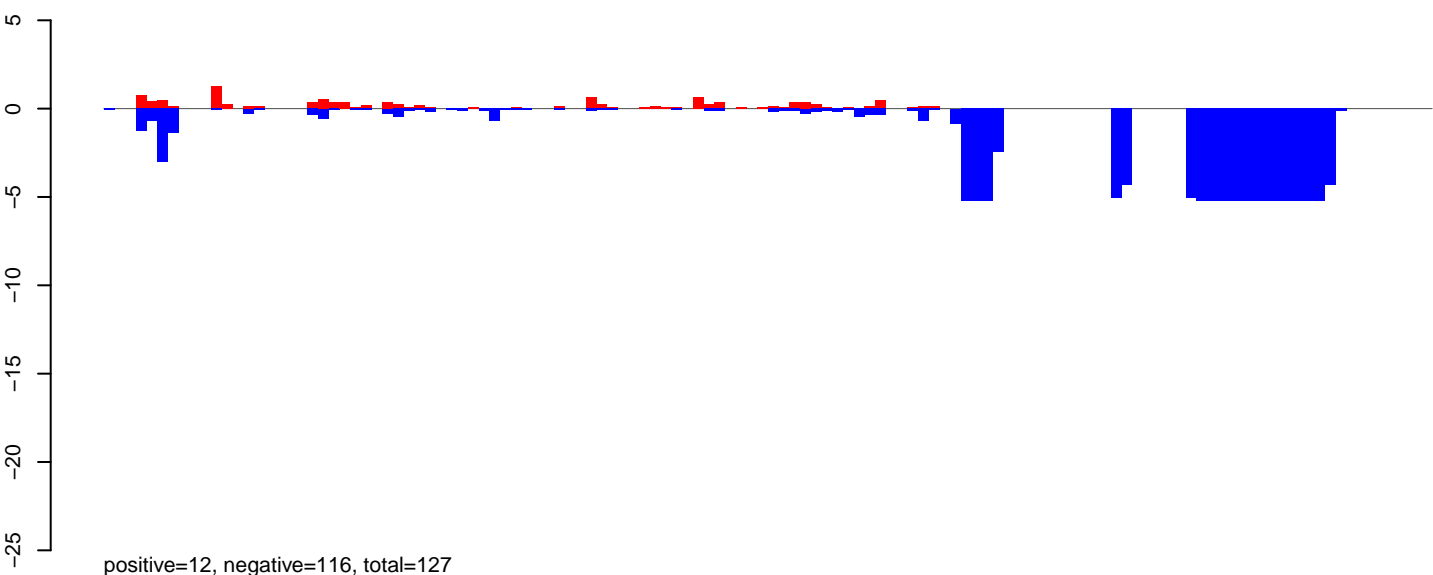
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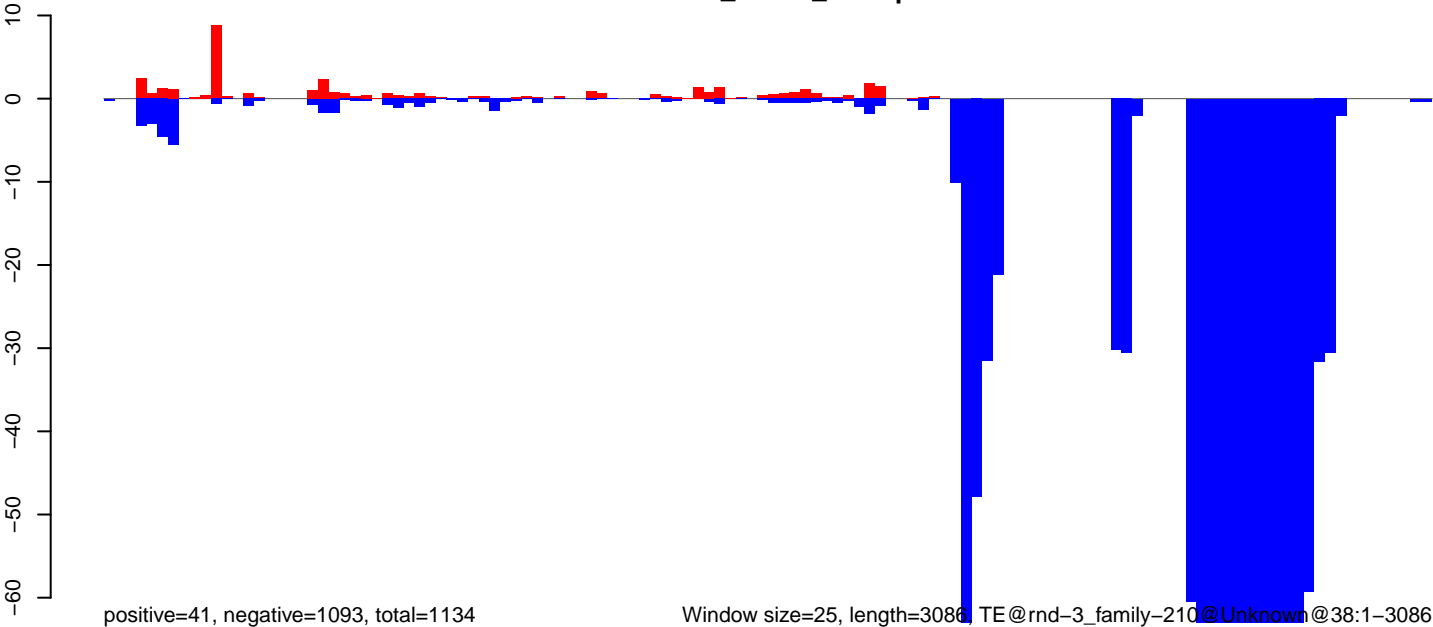
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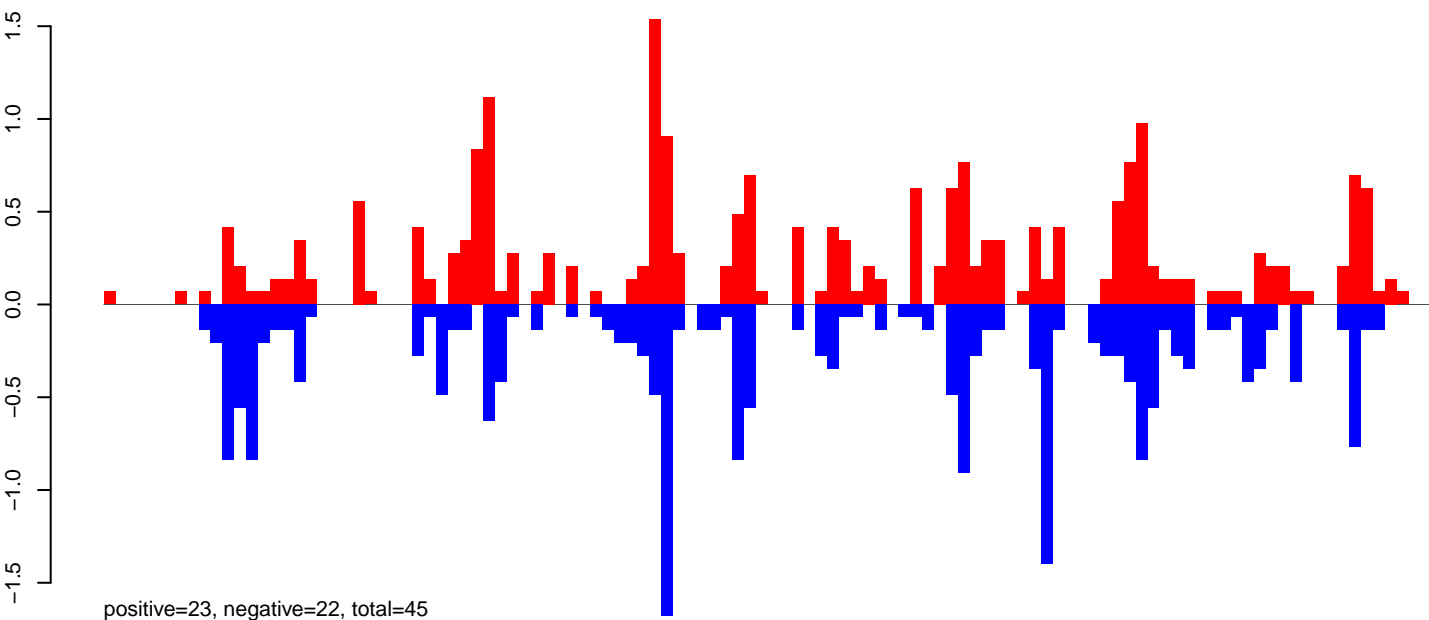
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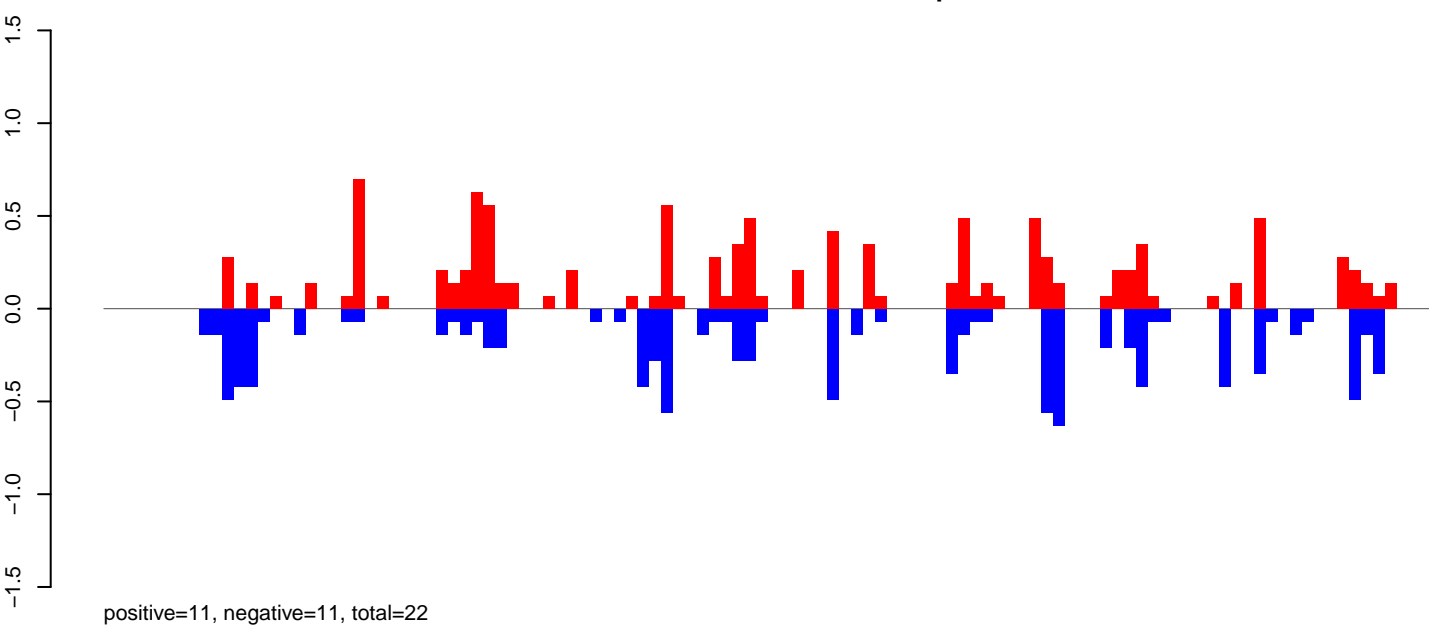
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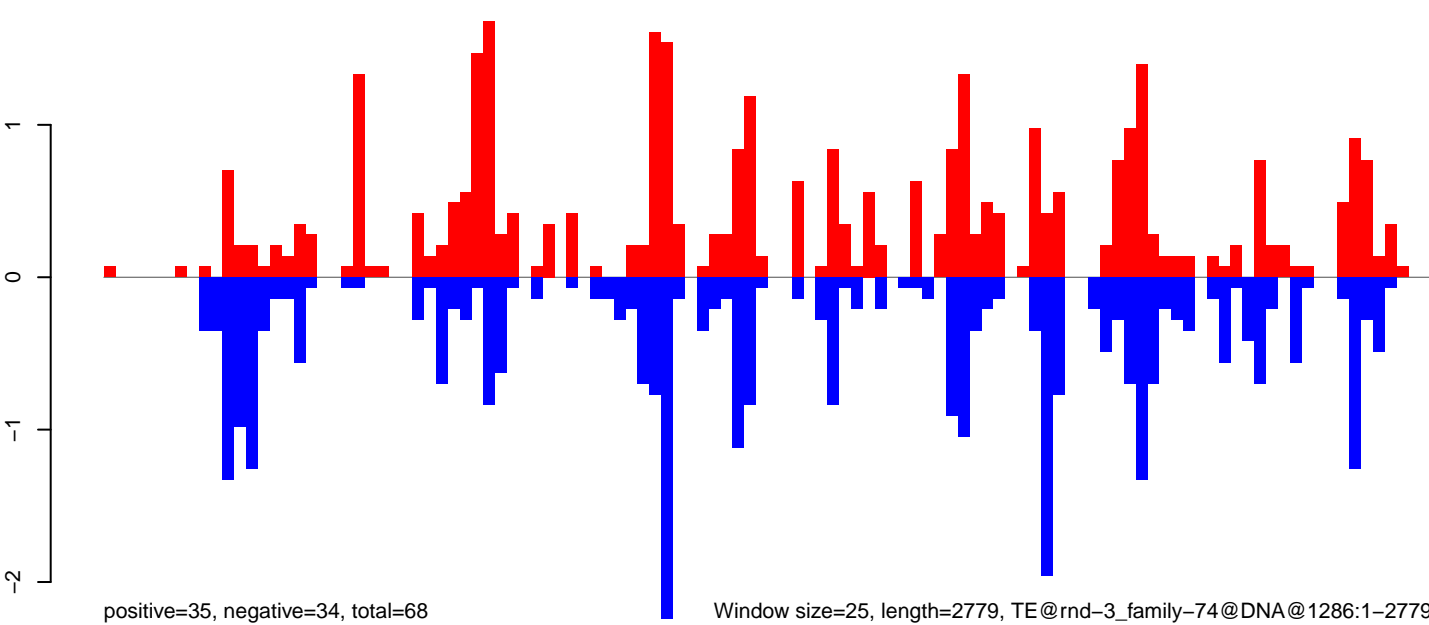
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CuQuin_Mcarc_NL.24_35.rep

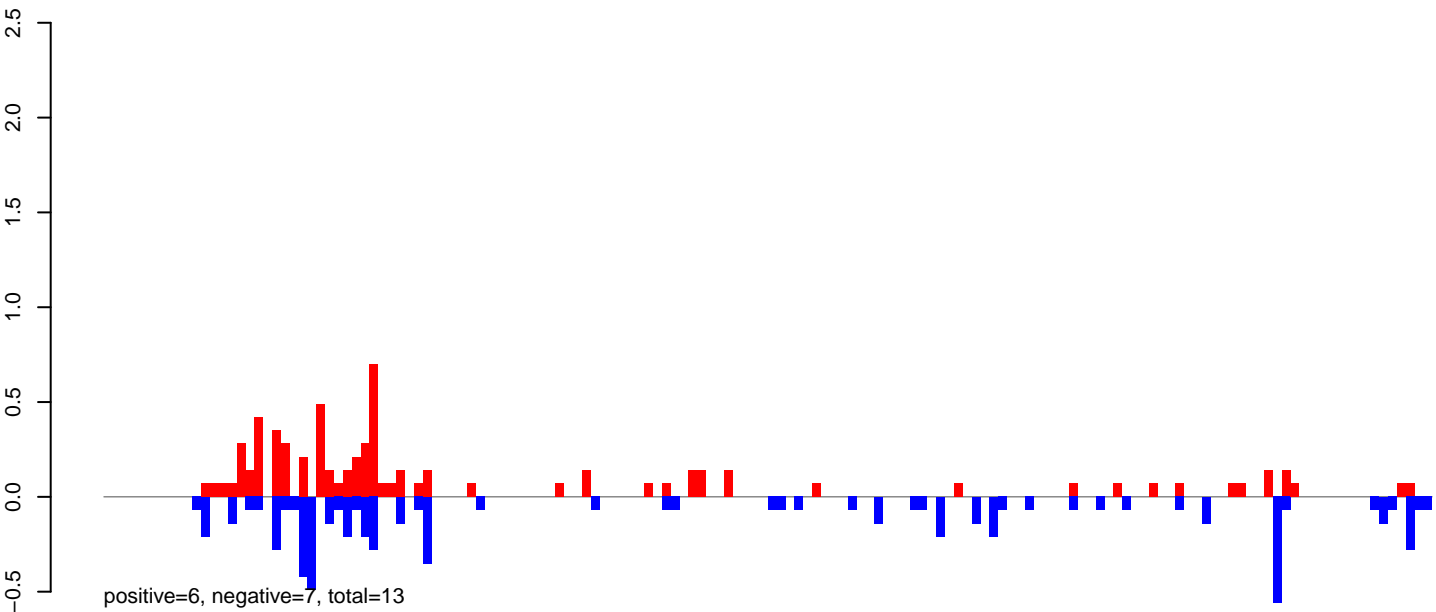


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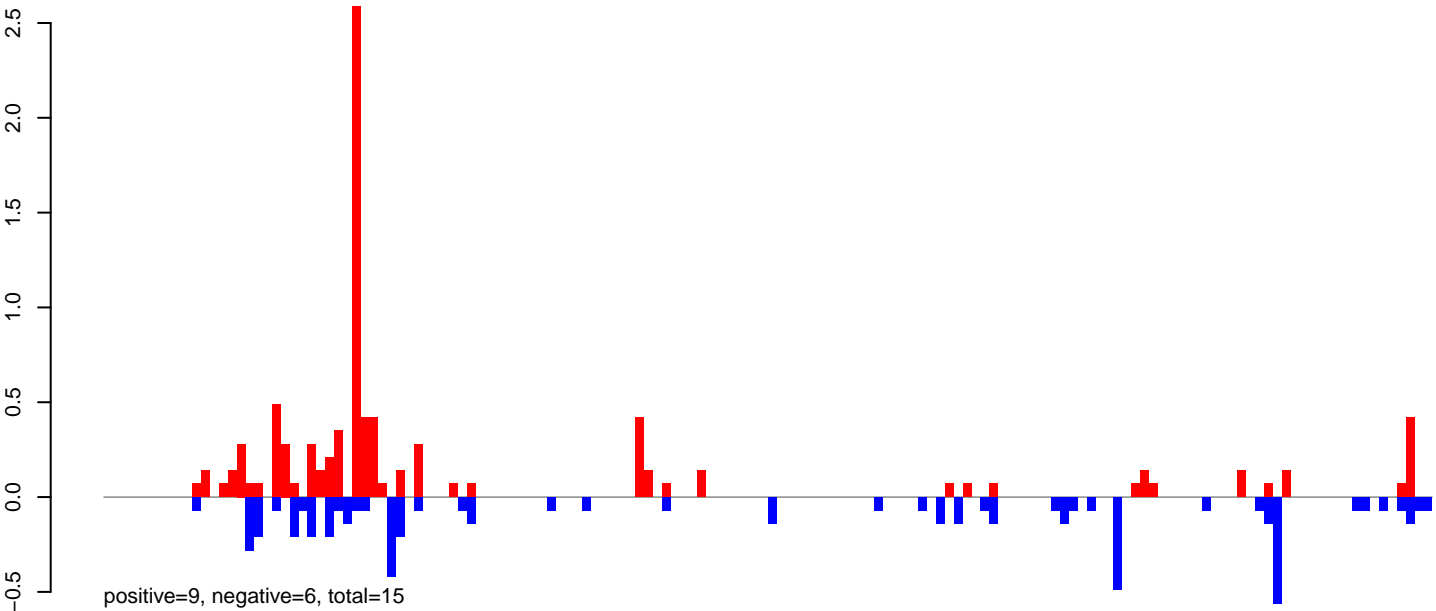


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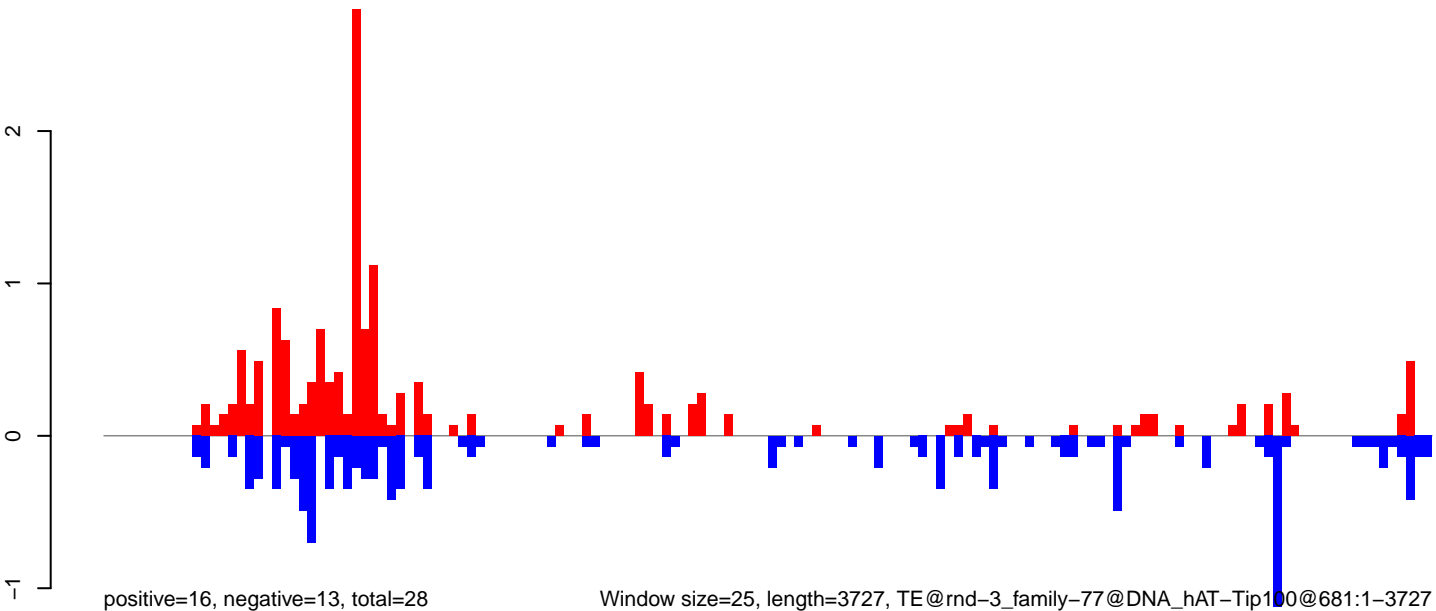
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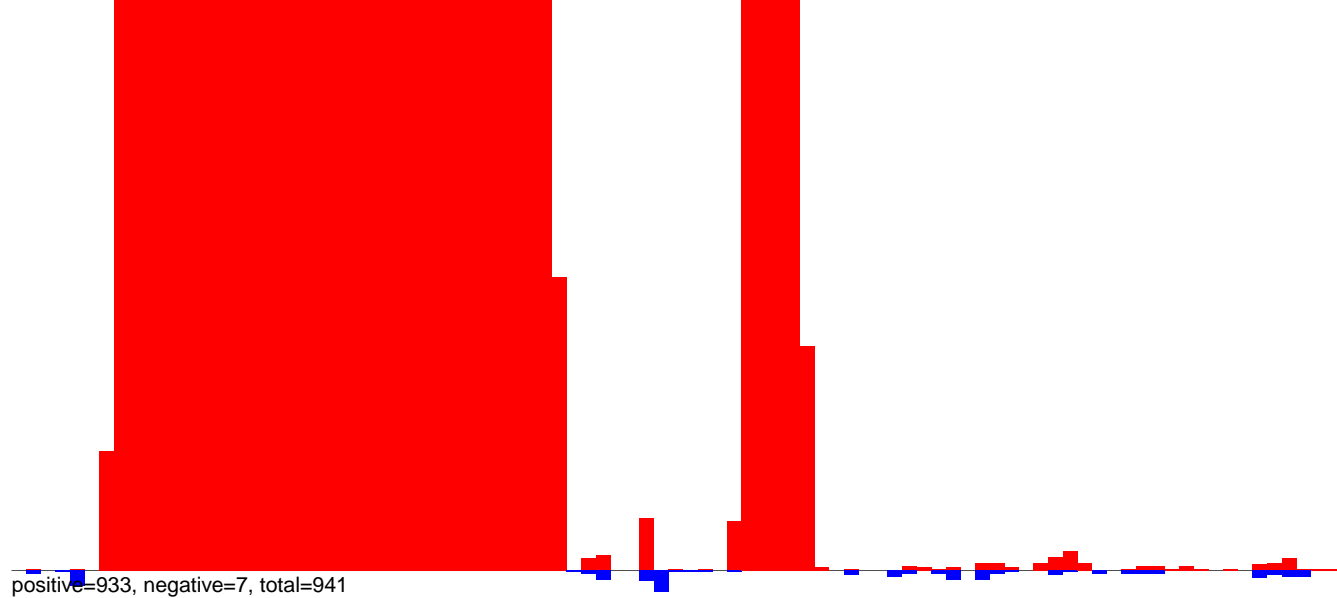
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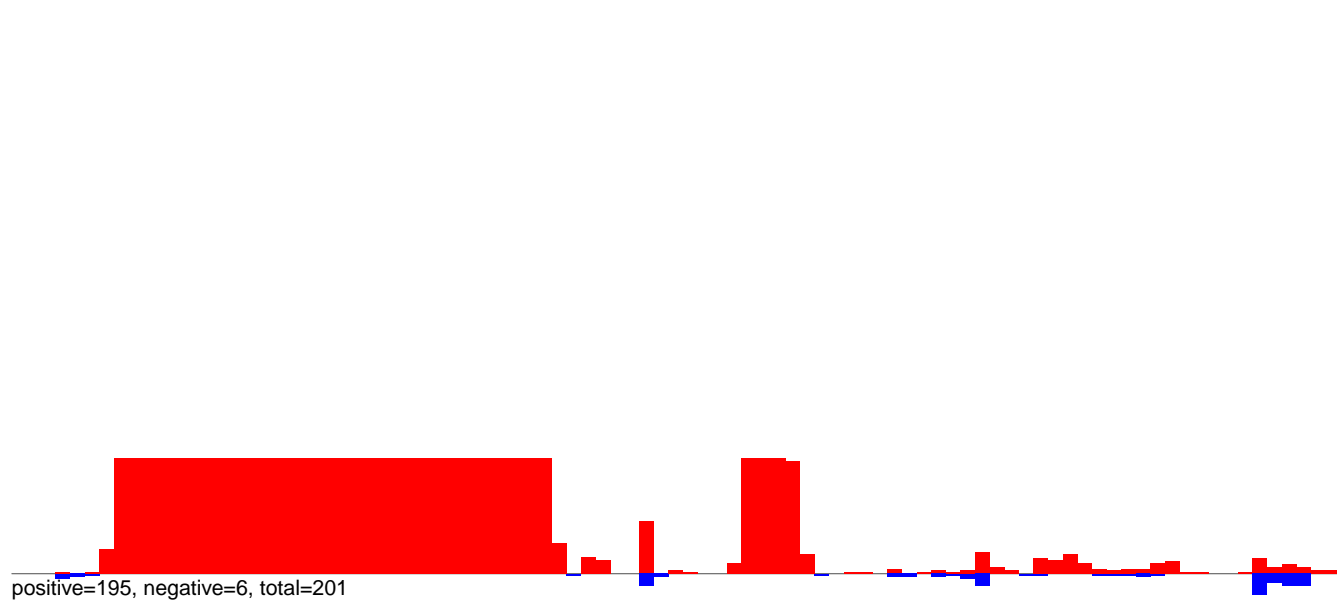
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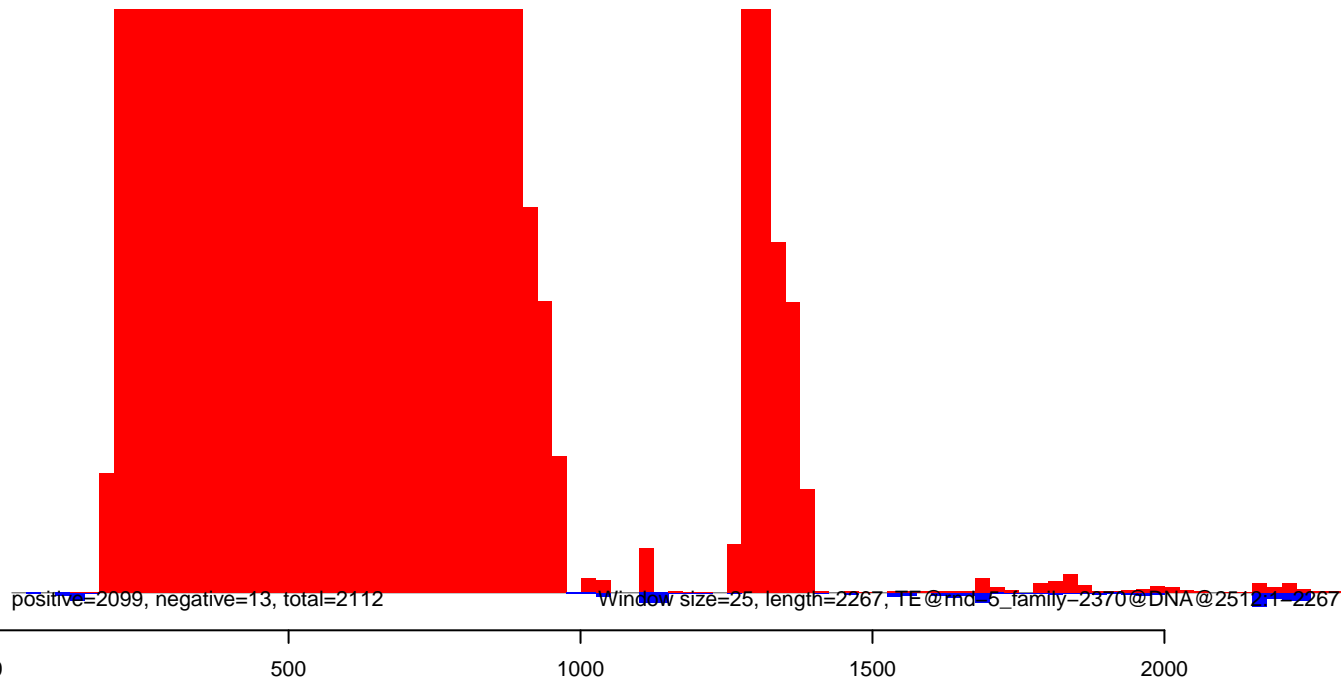
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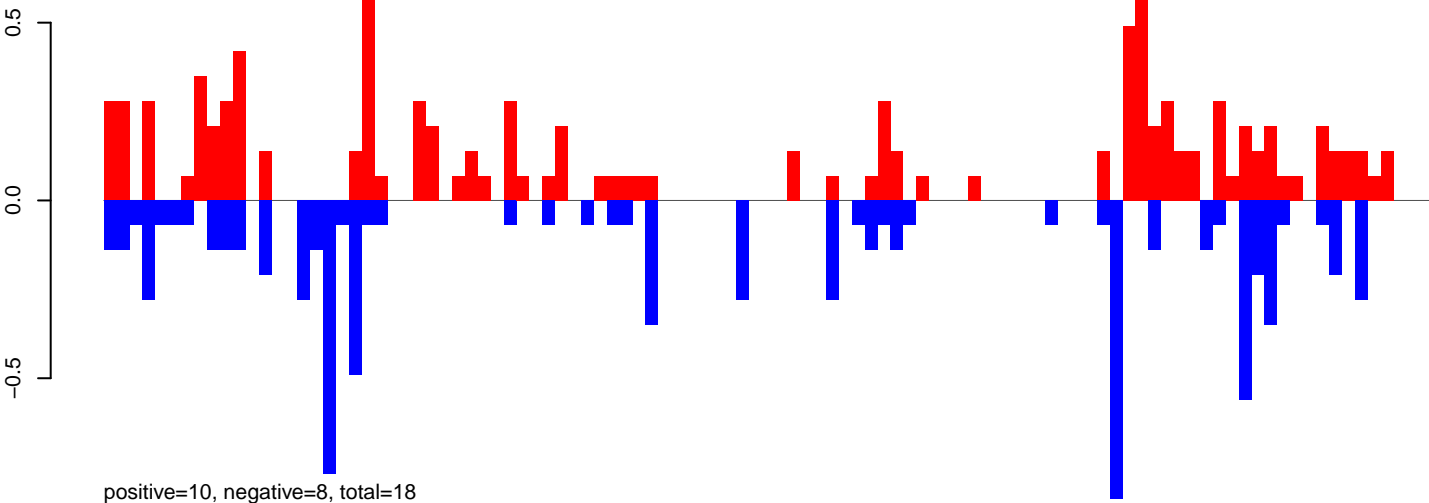
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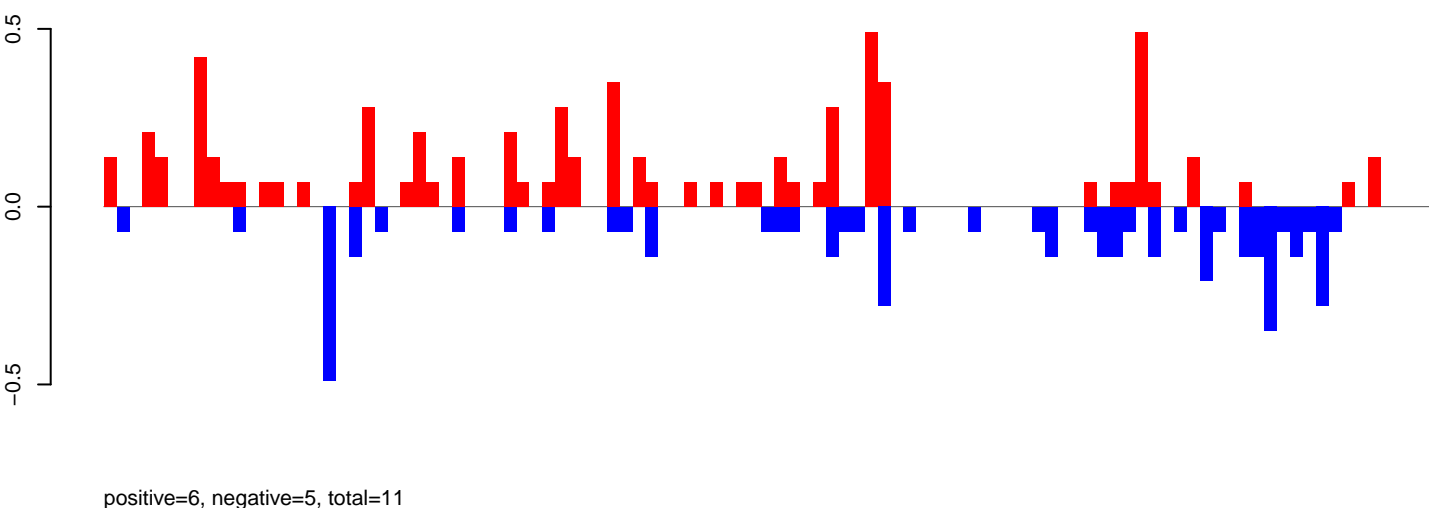
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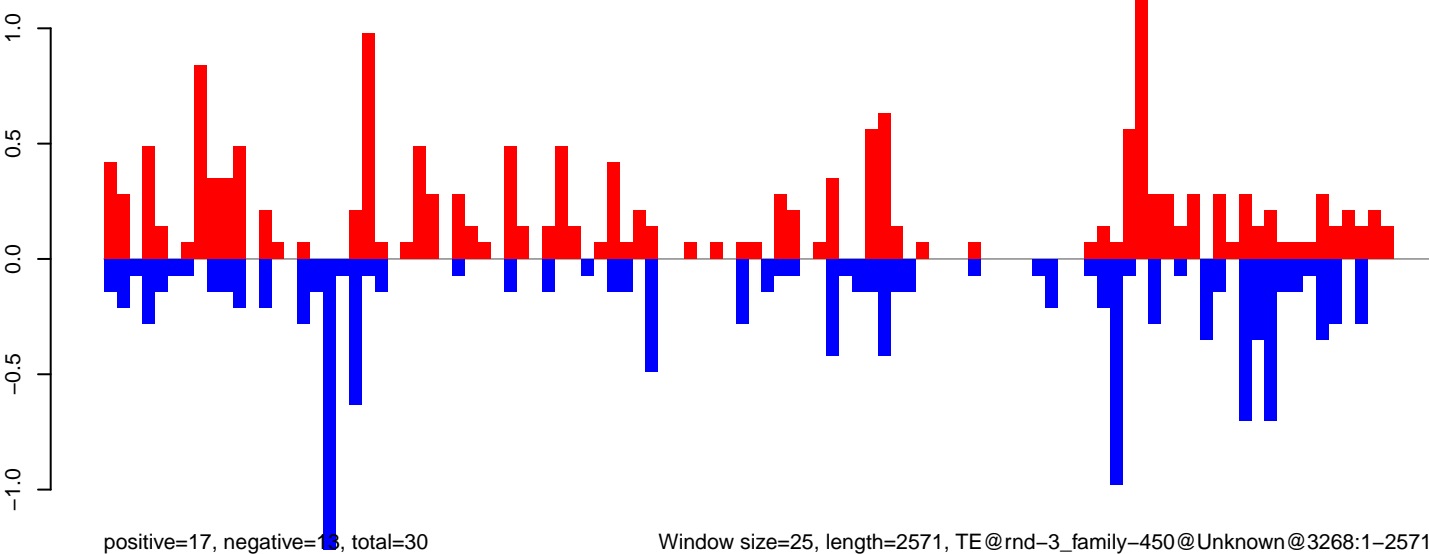
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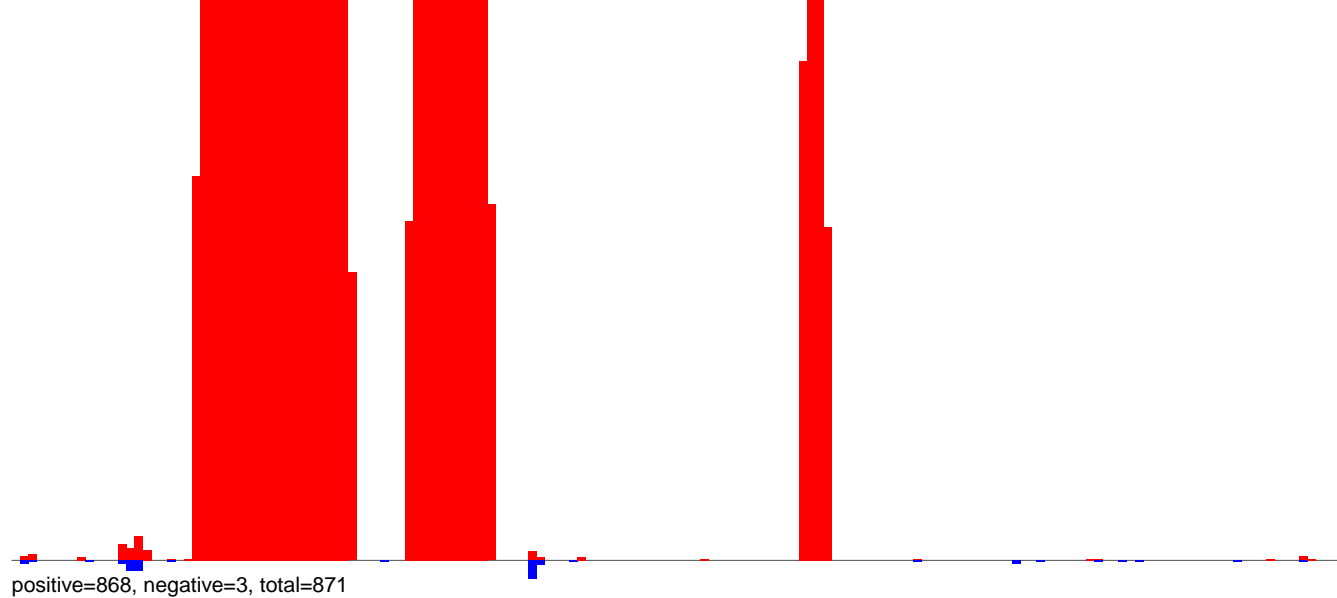
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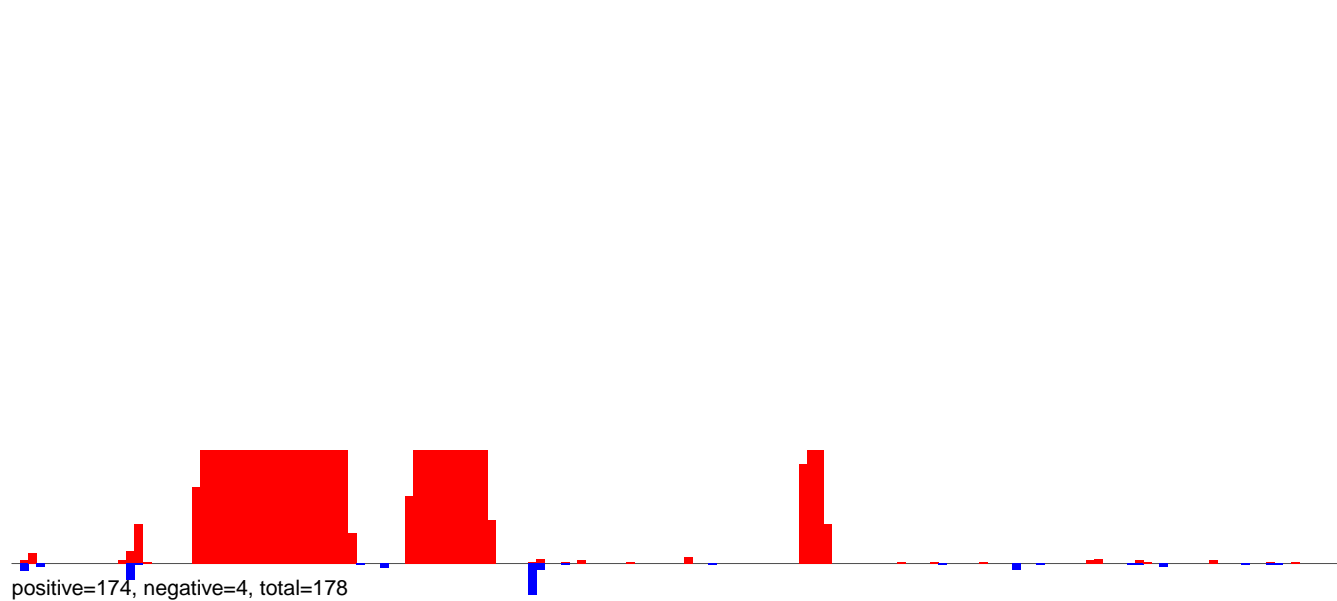
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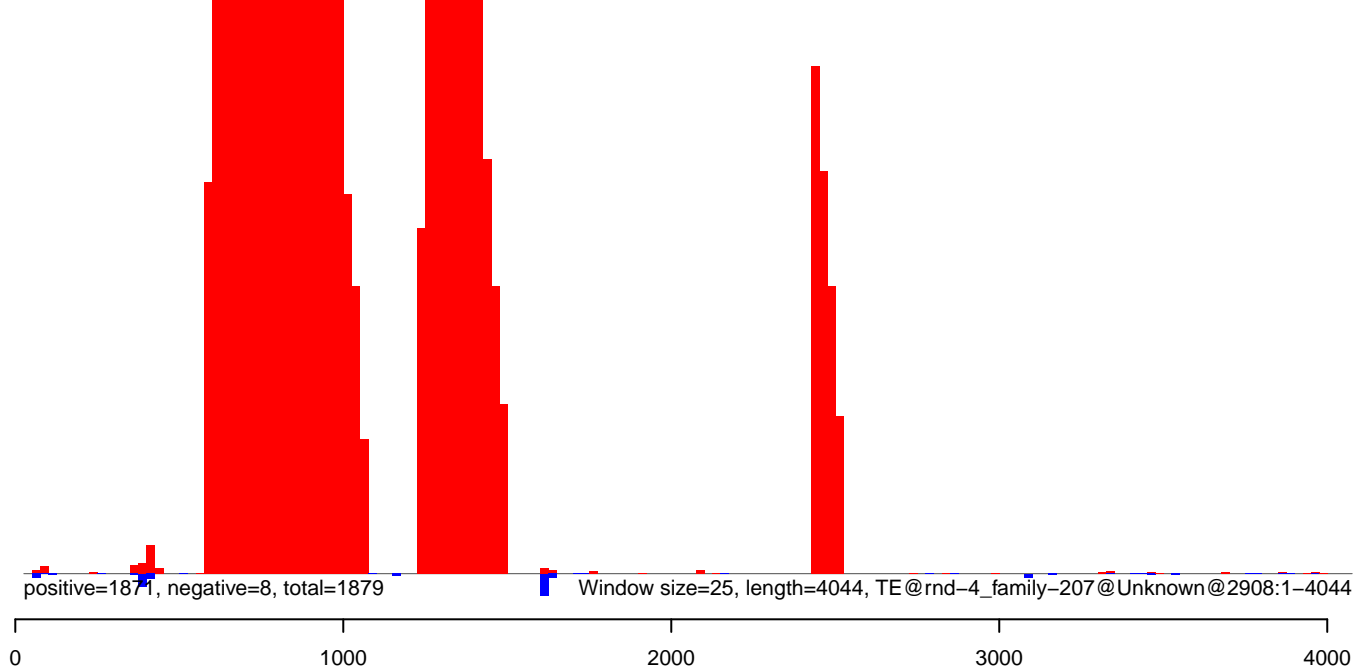
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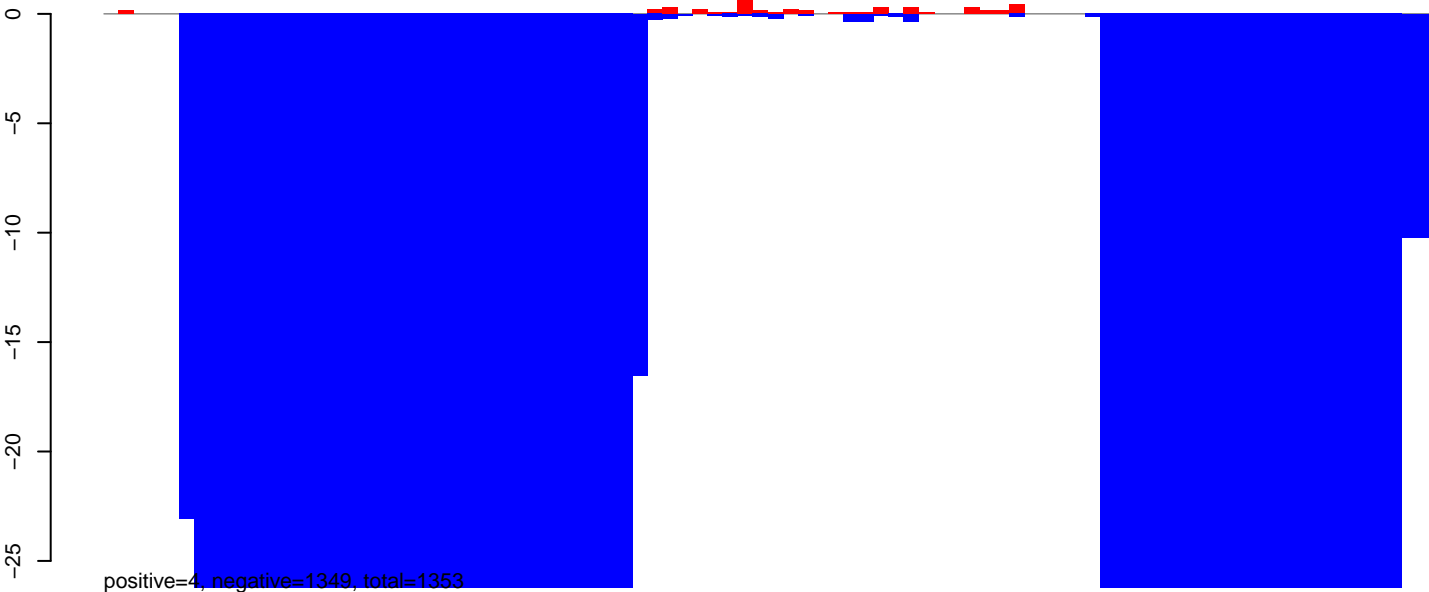
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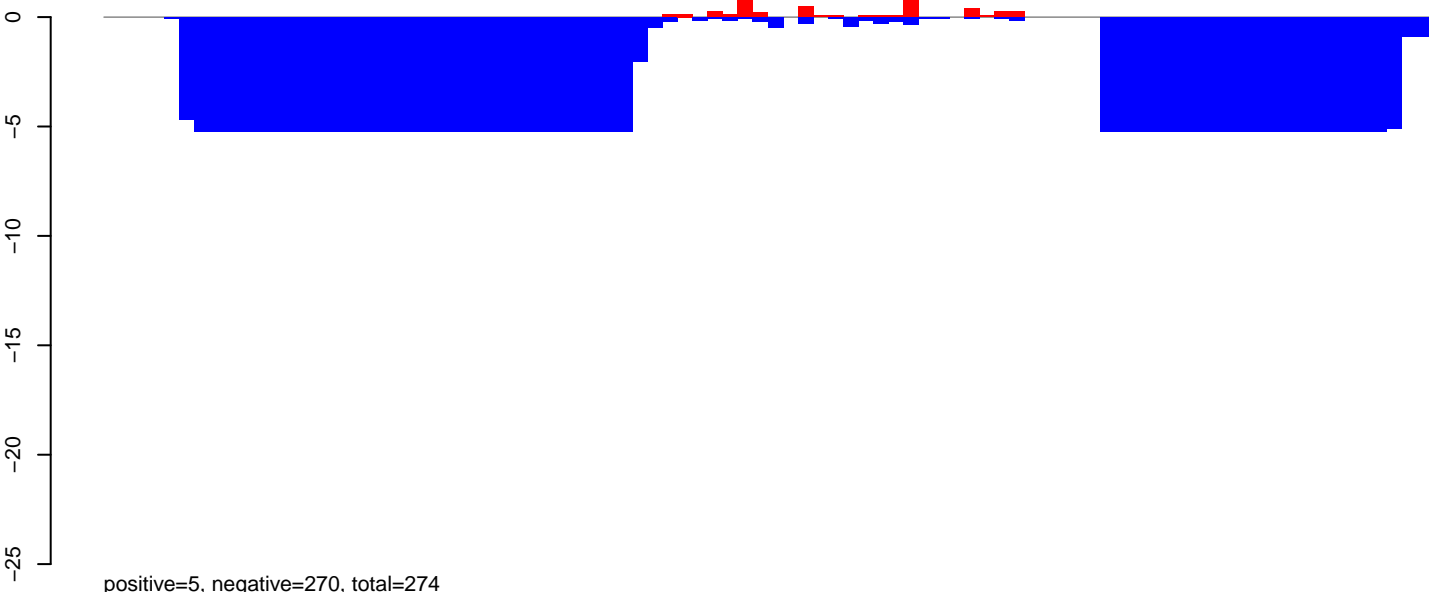
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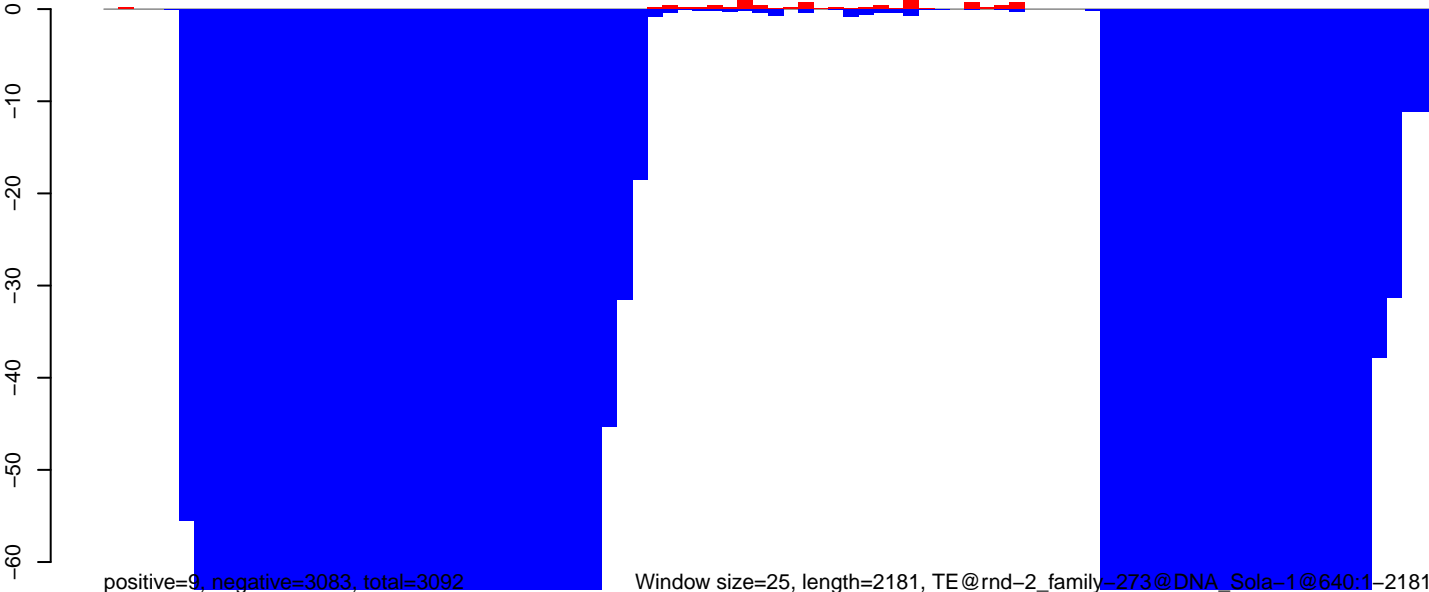
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CuQuin_Mcarc_NL.24_35.rep

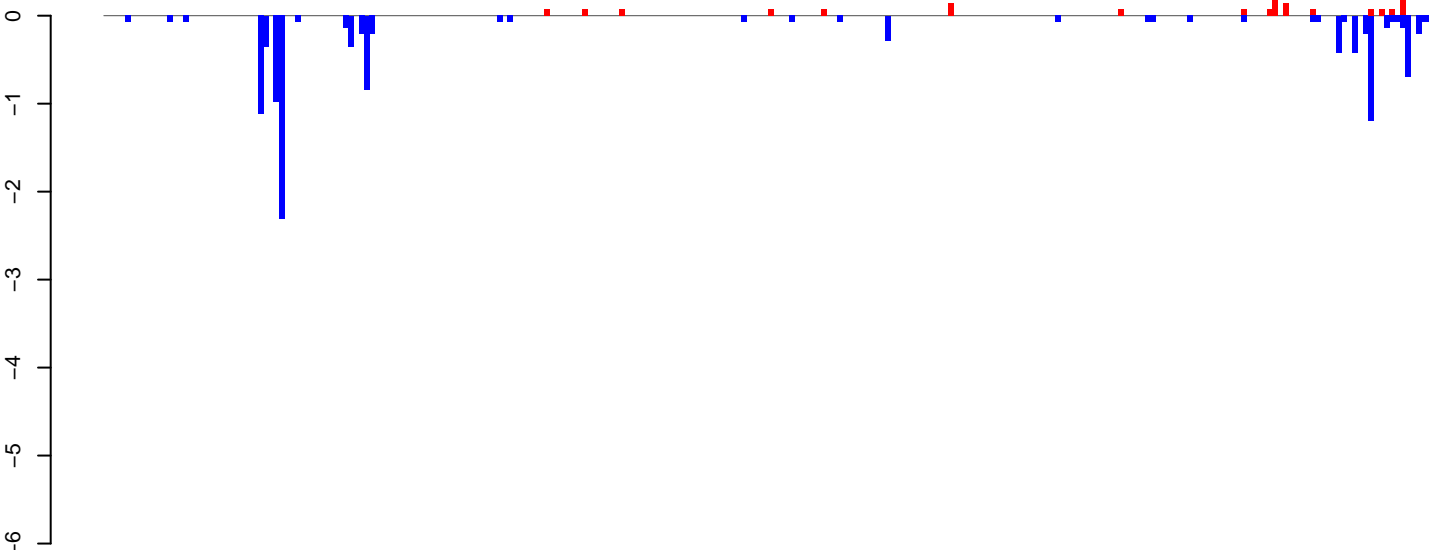


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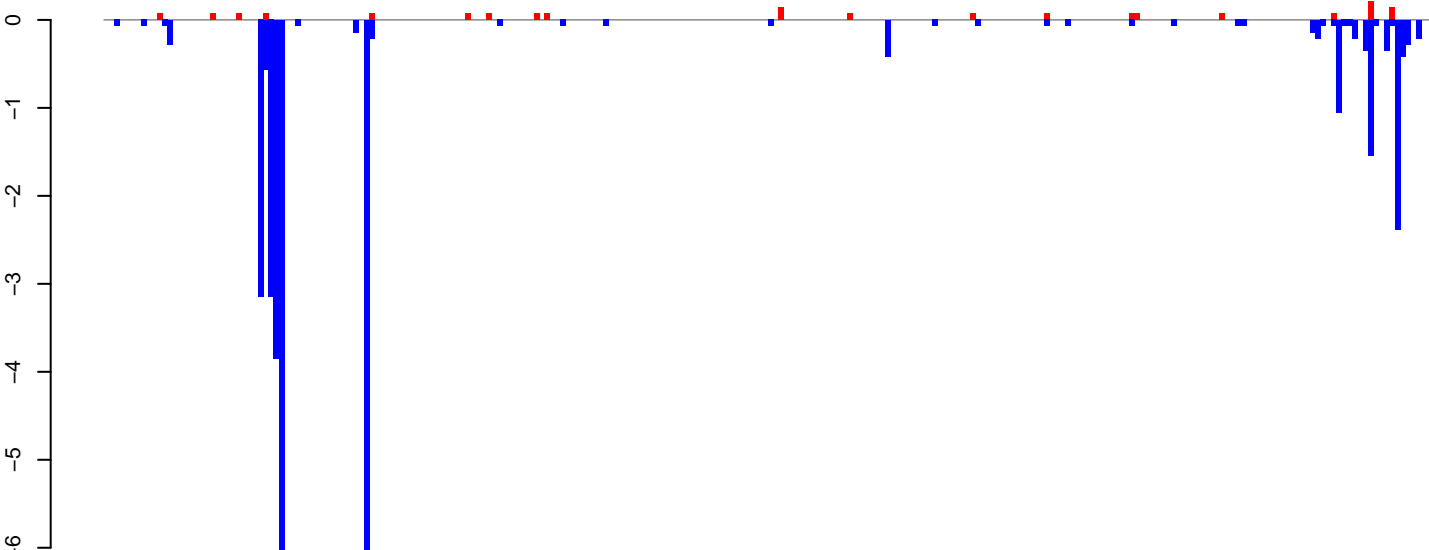
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CuQuin_Mcarc_NL.18_23.rep



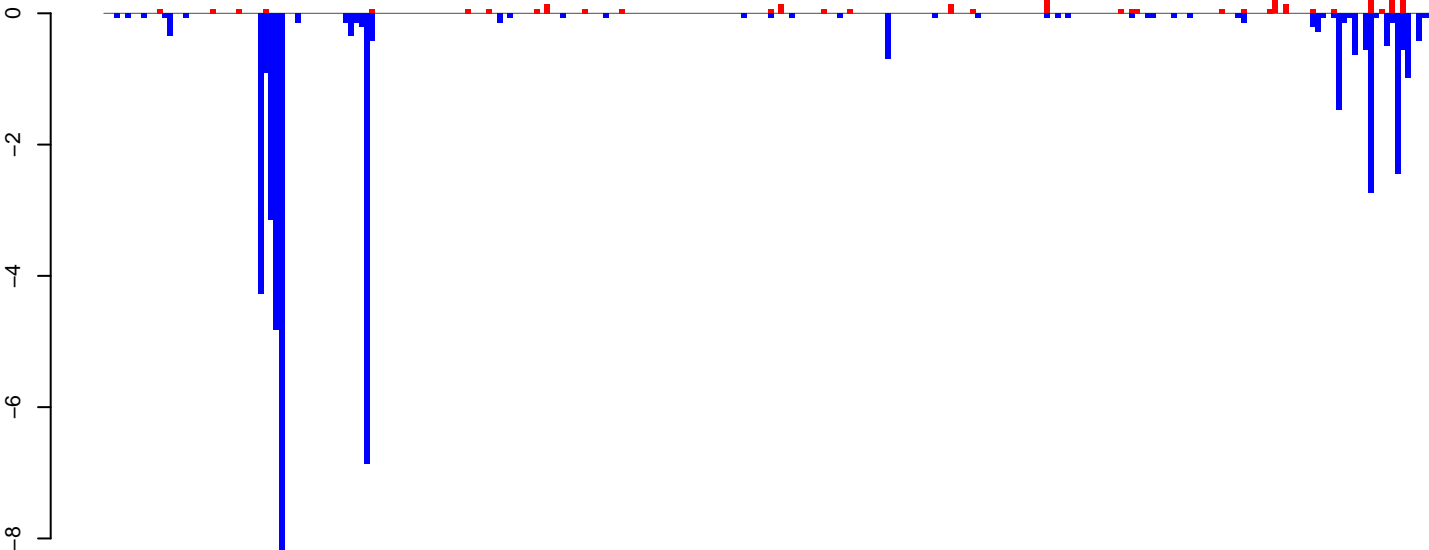
positive=2, negative=12, total=13

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=33, total=34

CuQuin_Mcarc_NL.rep

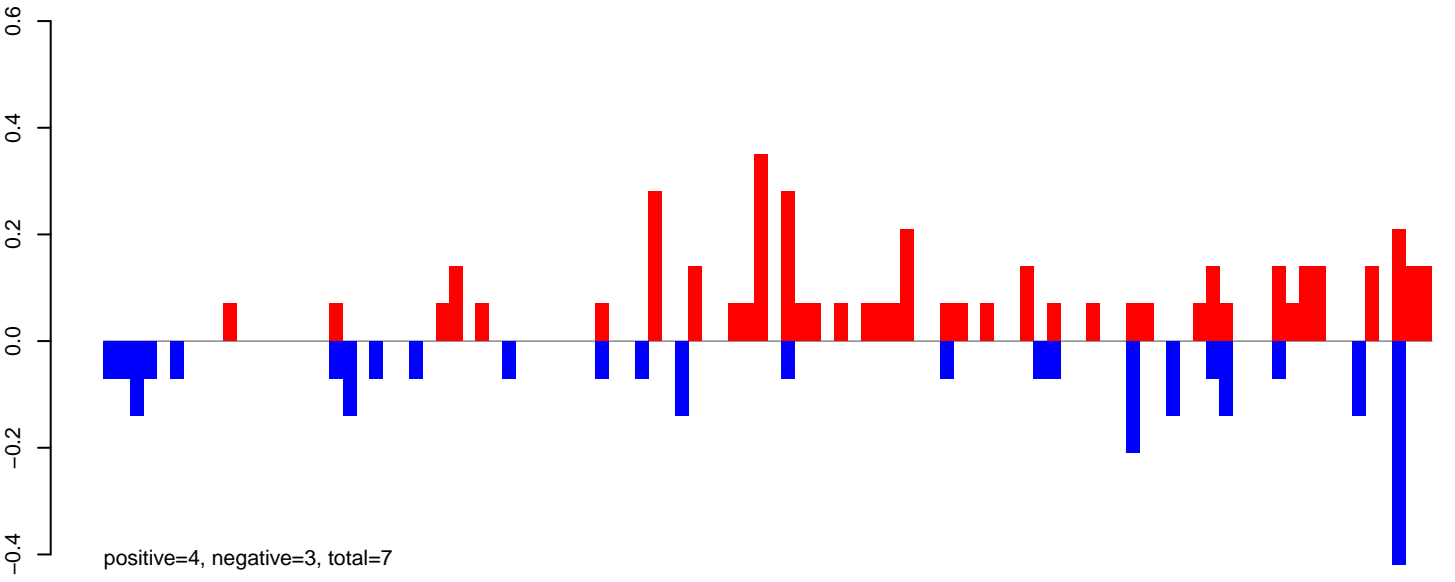


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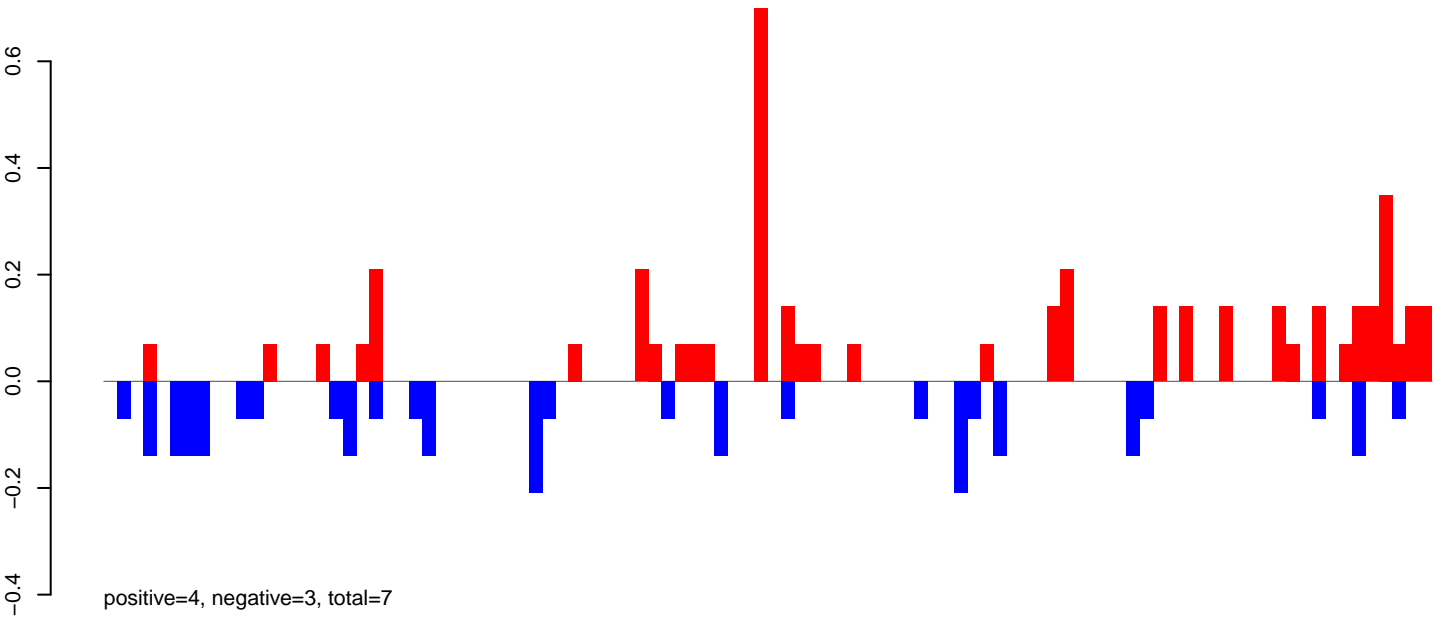
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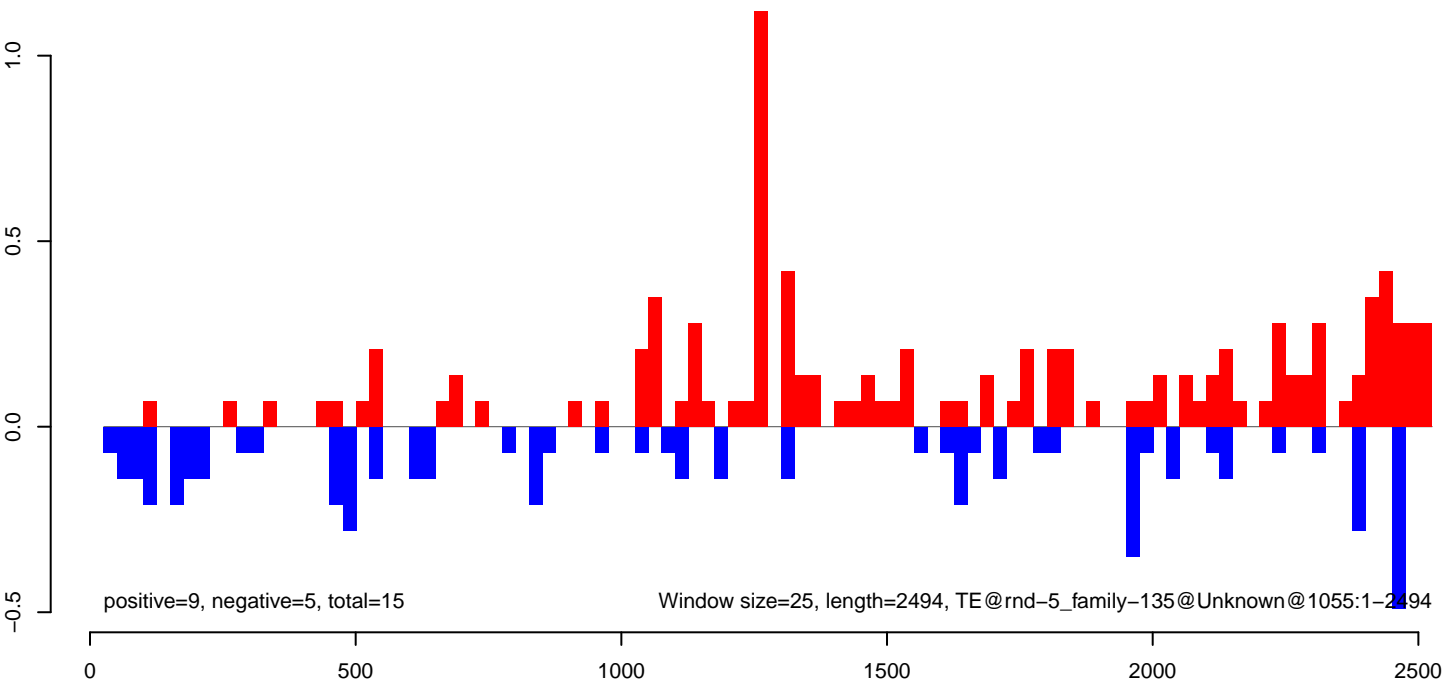
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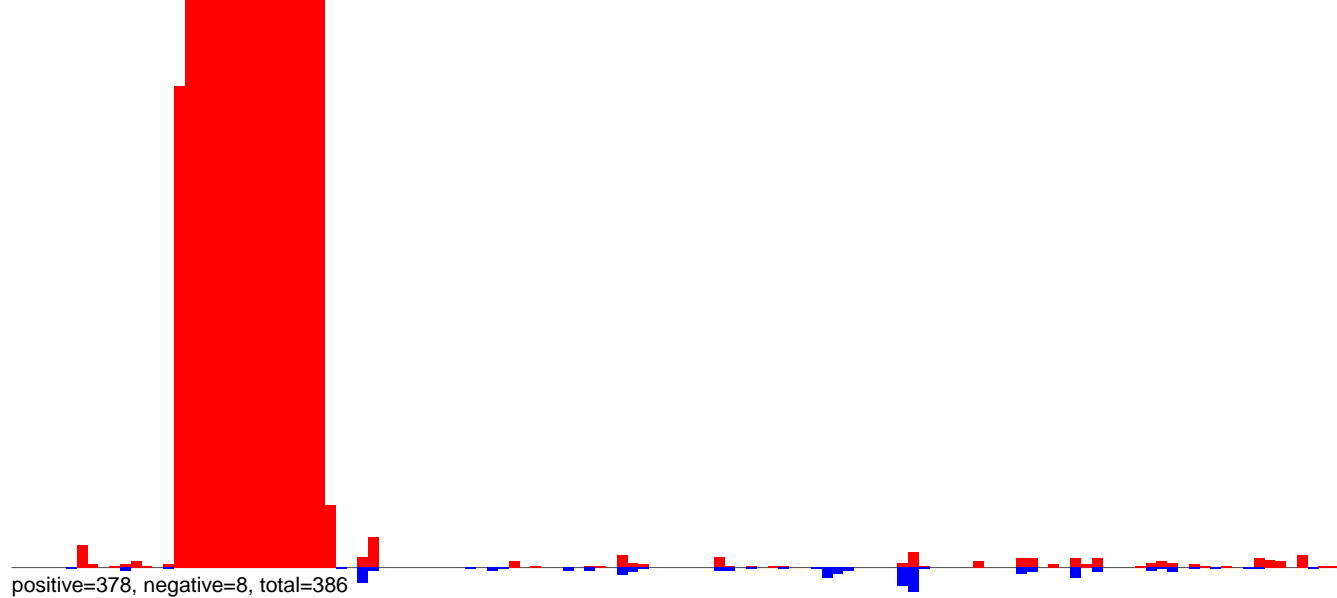
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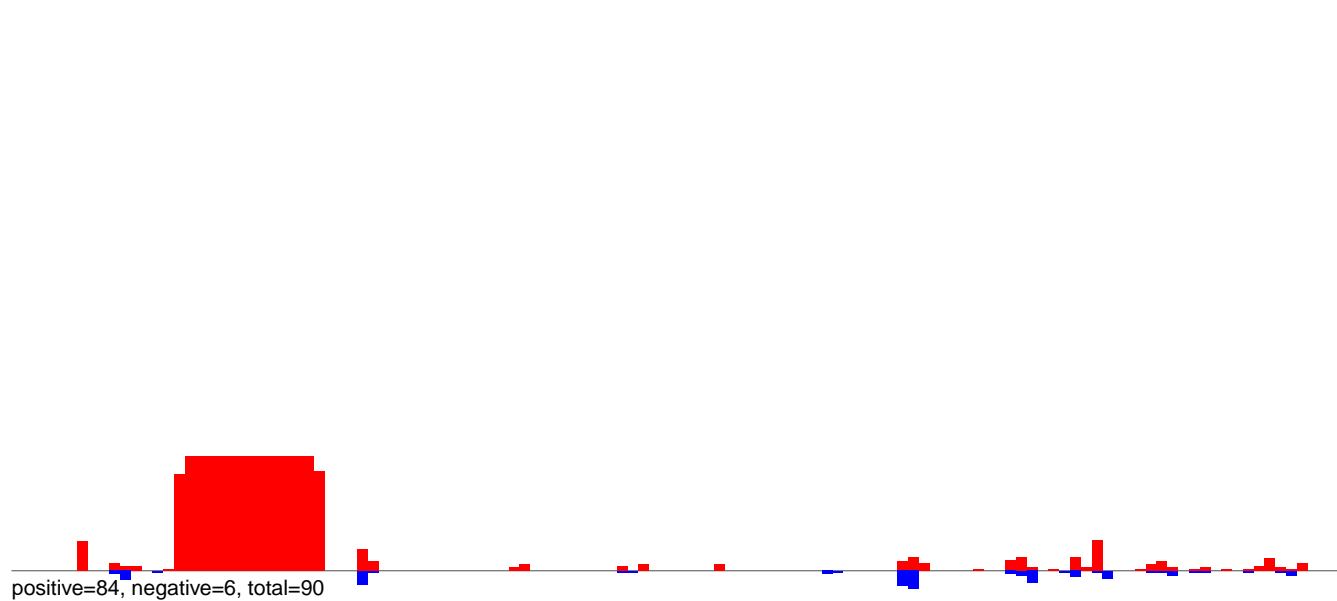
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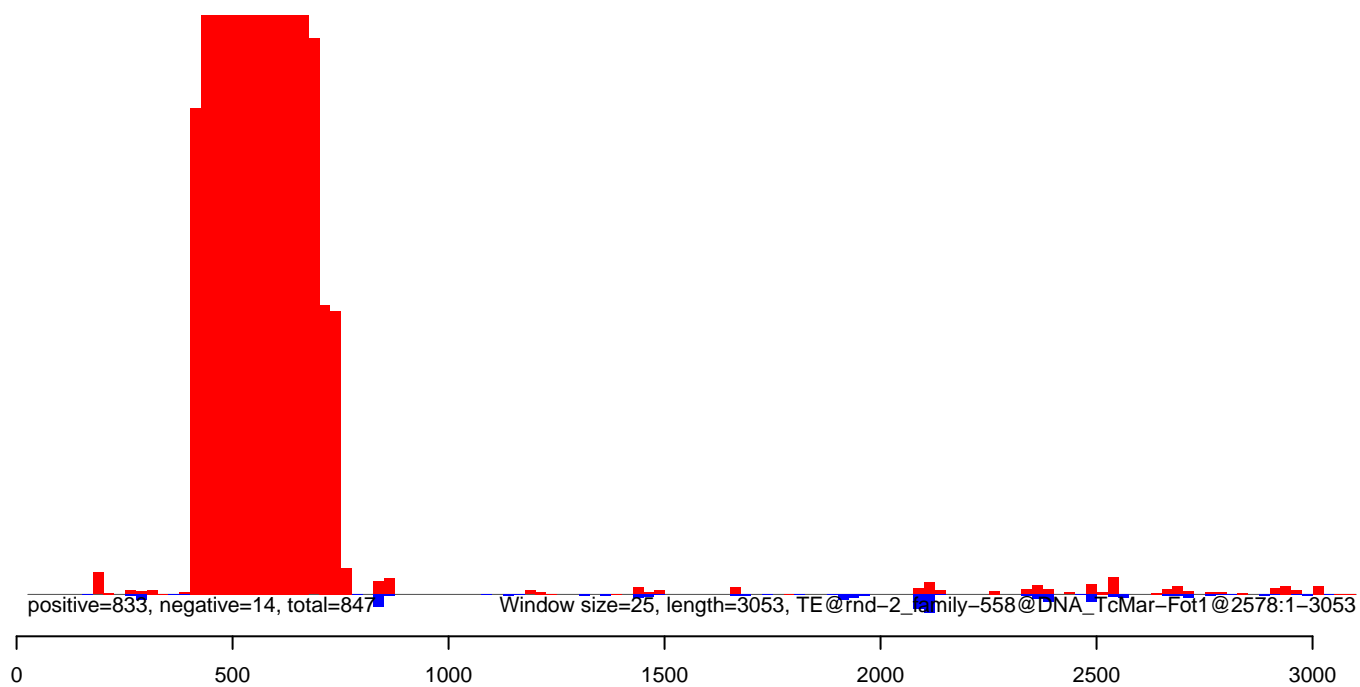
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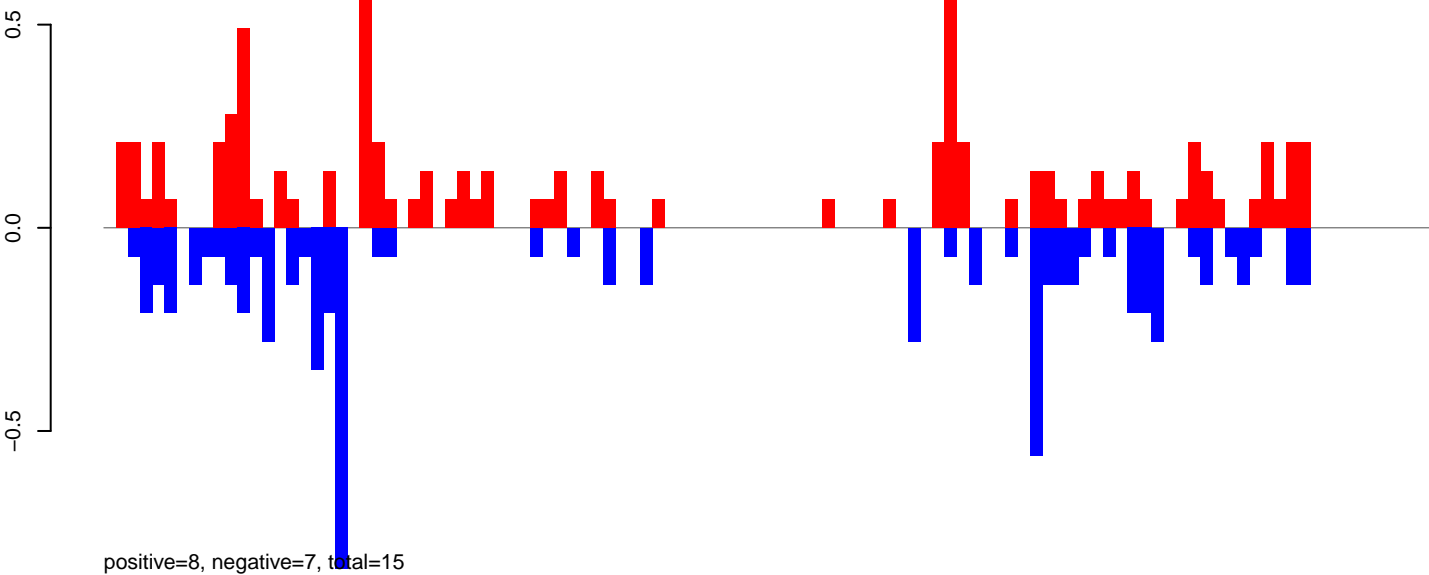
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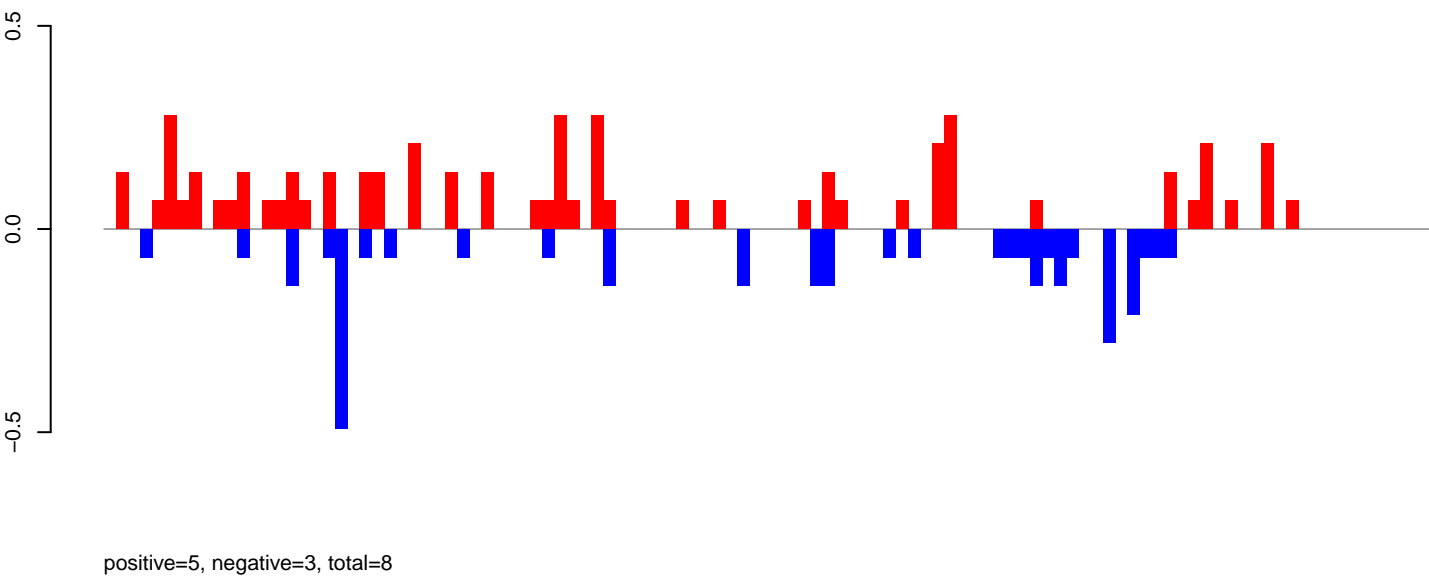
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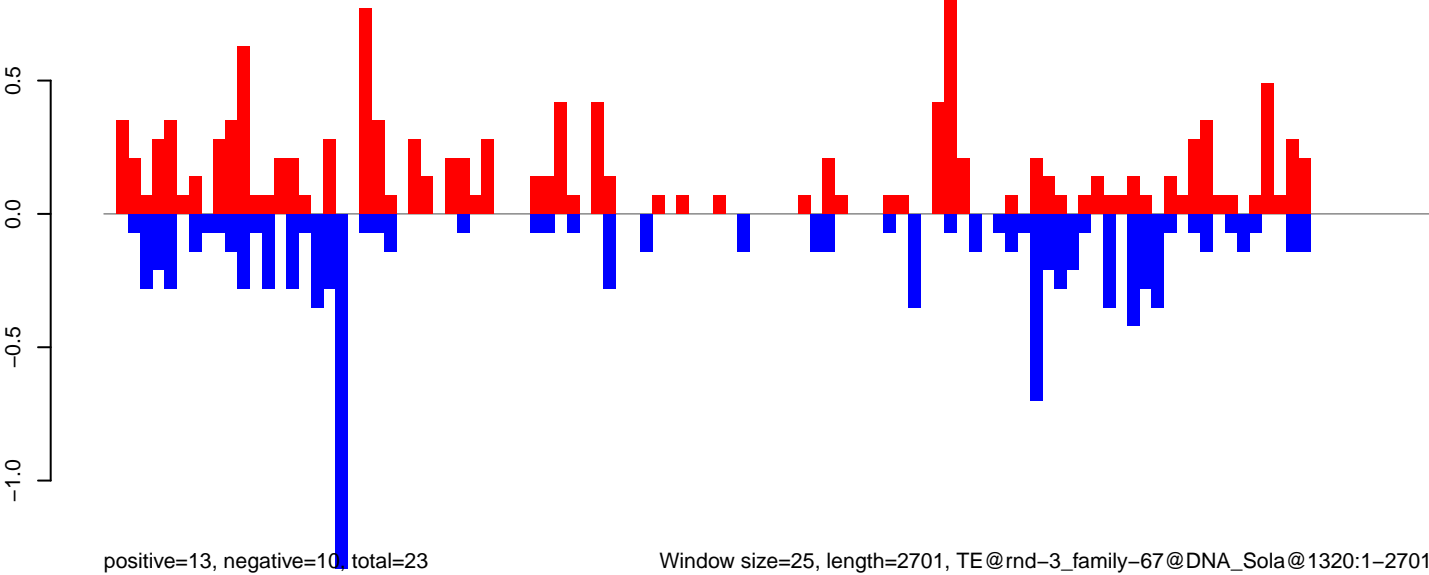
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CuQuin_Mcarc_NL.24_35.rep



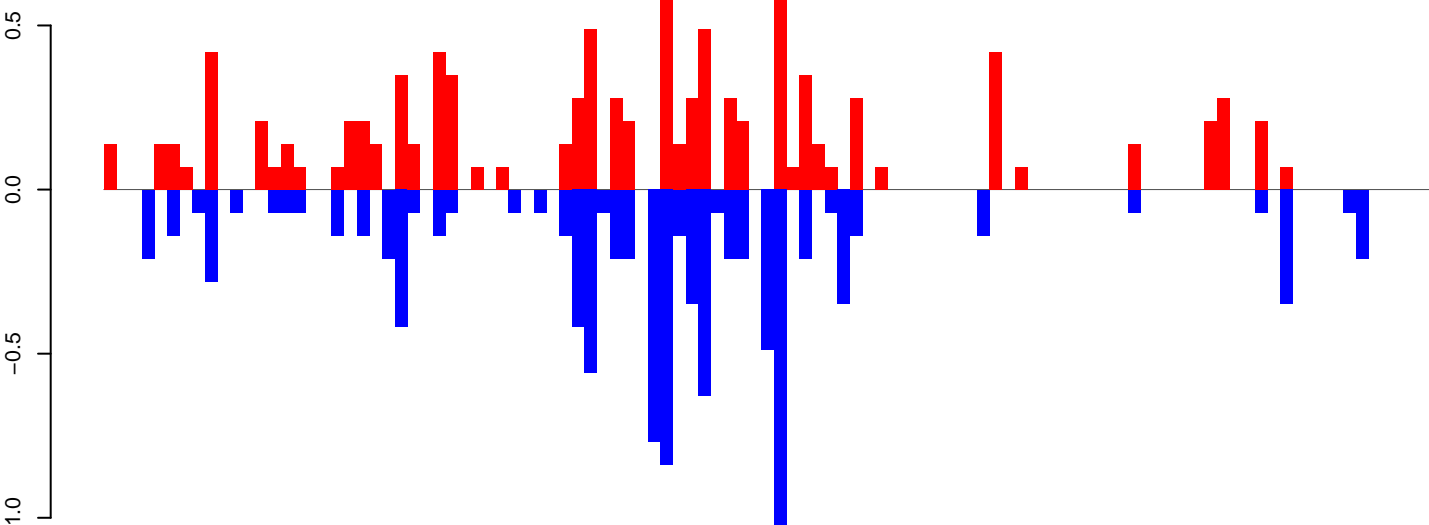
CuQuin_Mcarc_NL.rep



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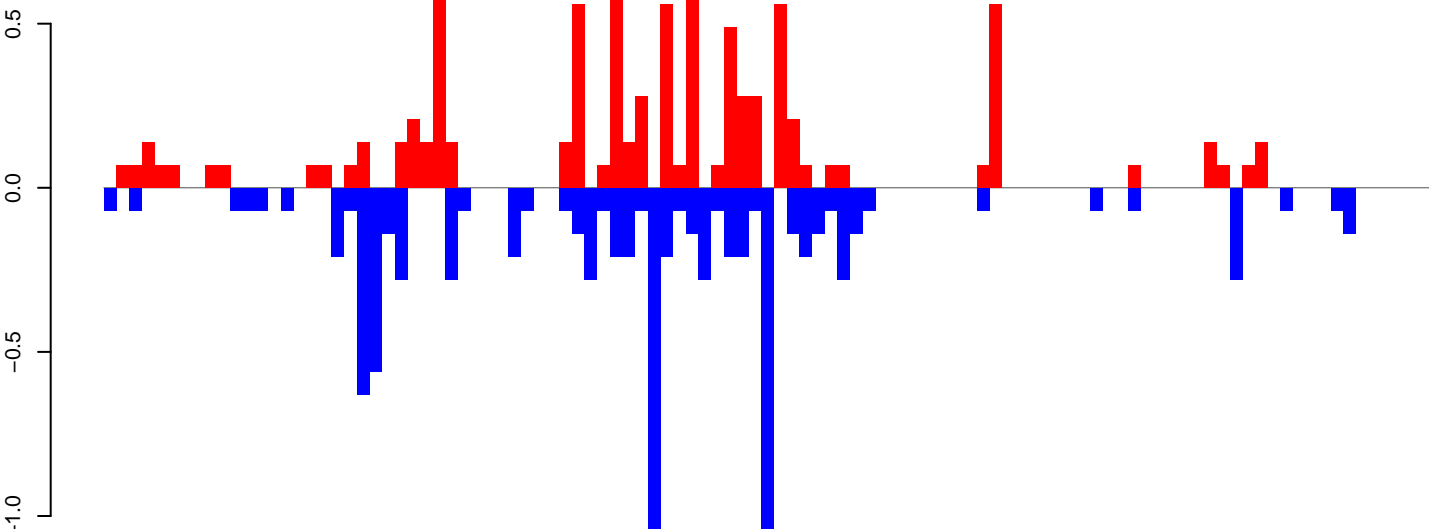
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CuQuin_Mcarc_NL.18_23.rep



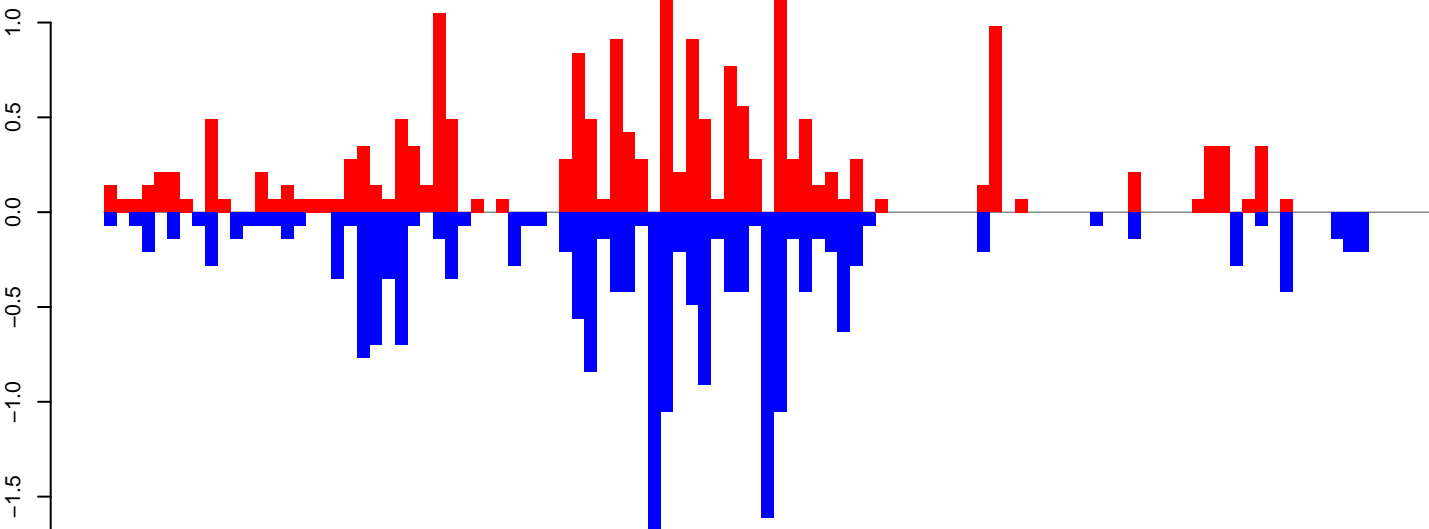
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CuQuin_Mcarc_NL.24_35.rep



positive=8, negative=9, total=18

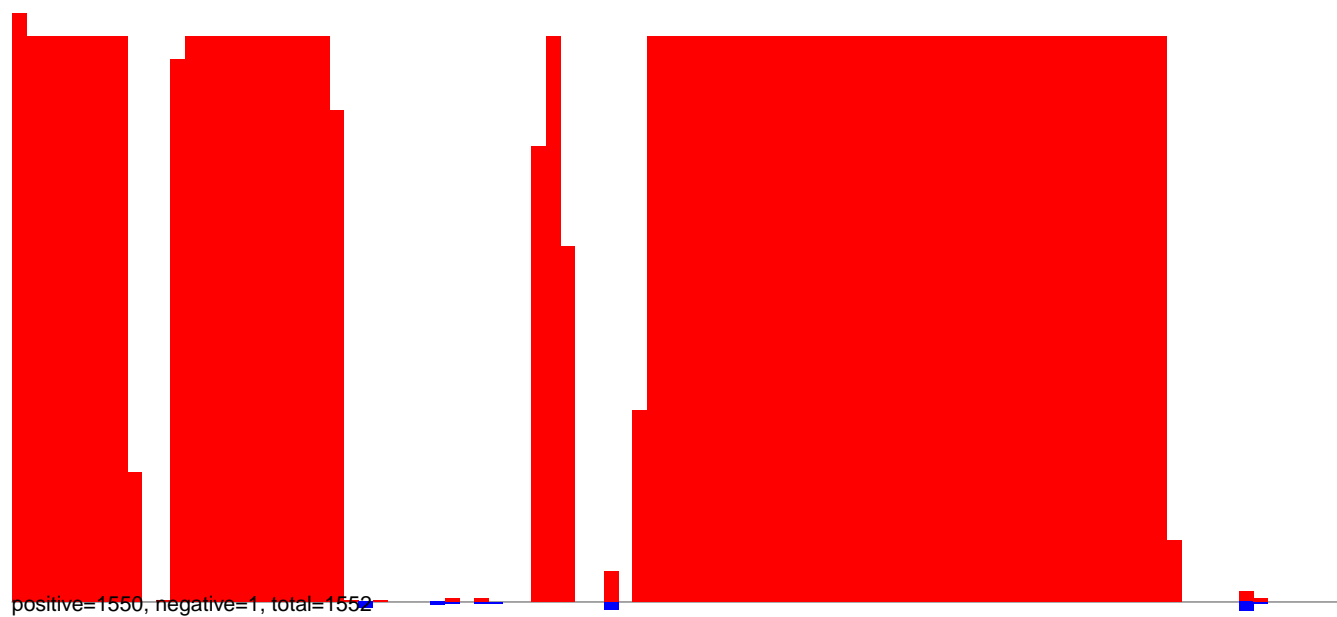
CuQuin_Mcarc_NL.rep



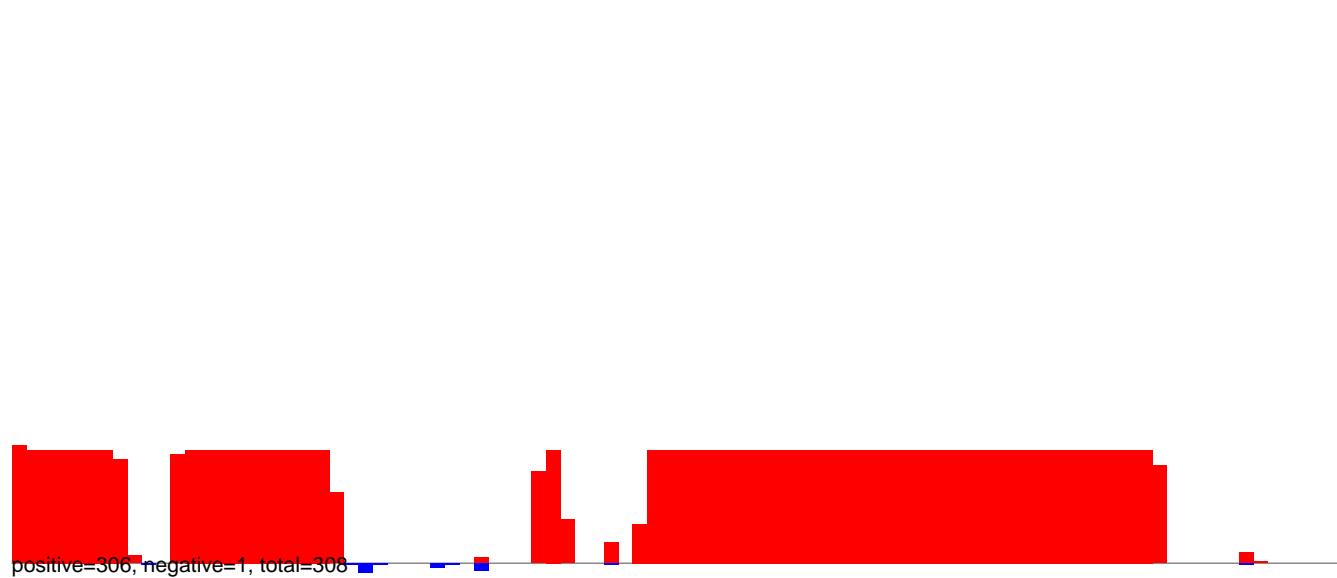
positive=19, negative=20, total=39

Window size=25, length=2612, TE@rnd-5_family-112@DNA_TcMar-Fot1@3046:1-2612

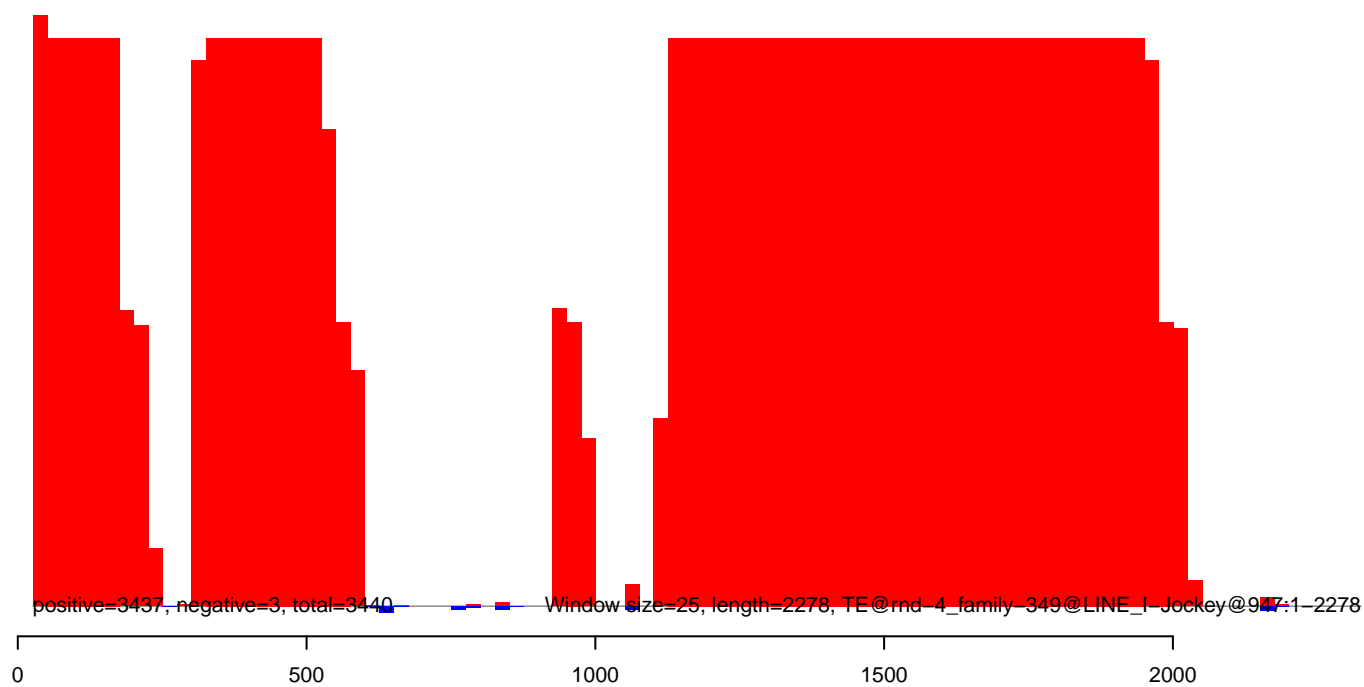
CuQuin_Mcarc_NL.18_23.rep



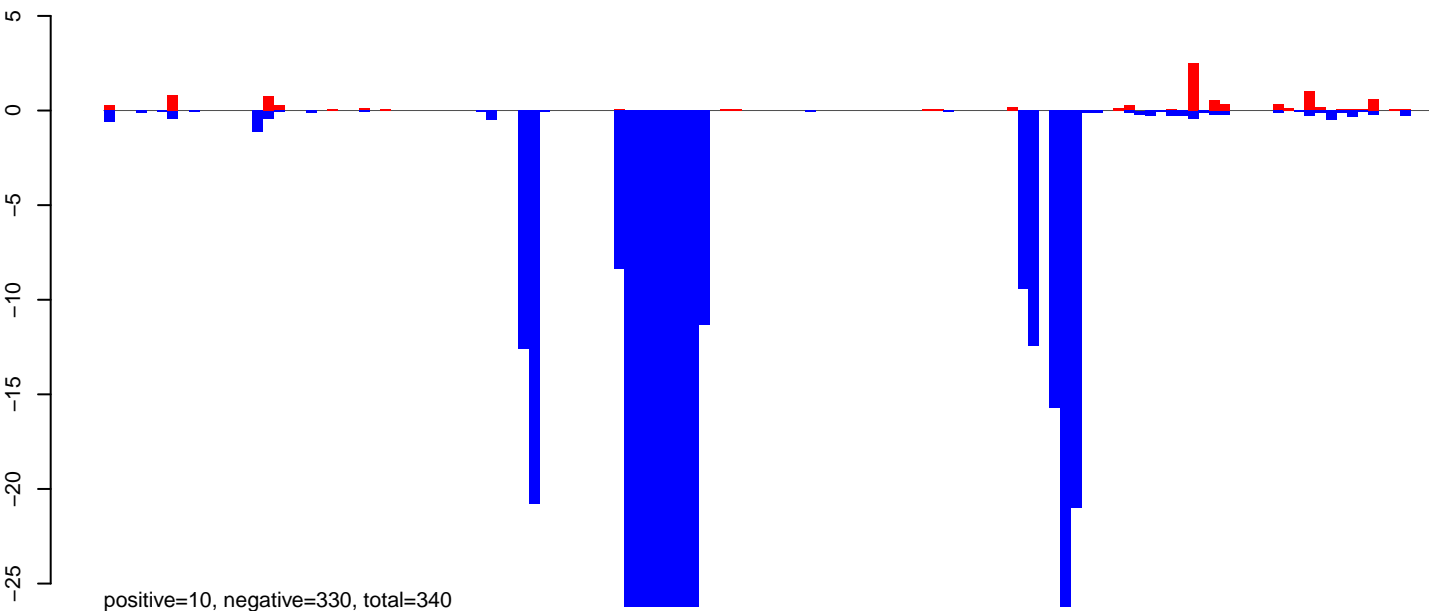
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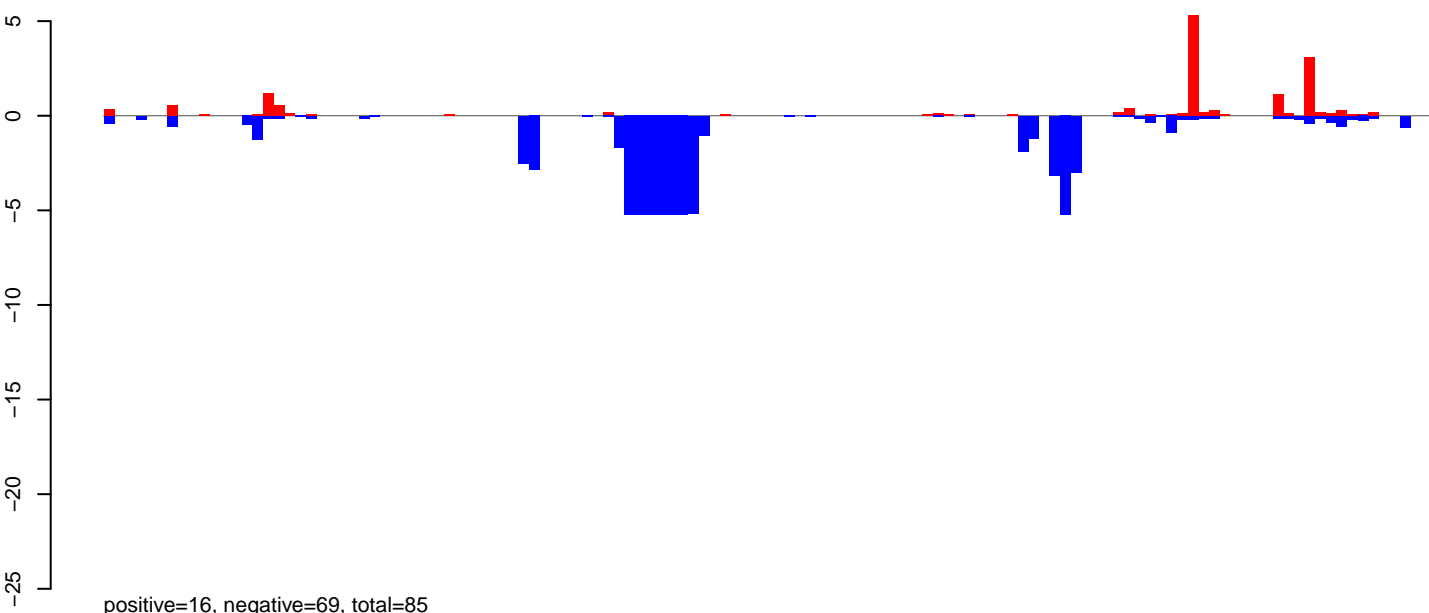
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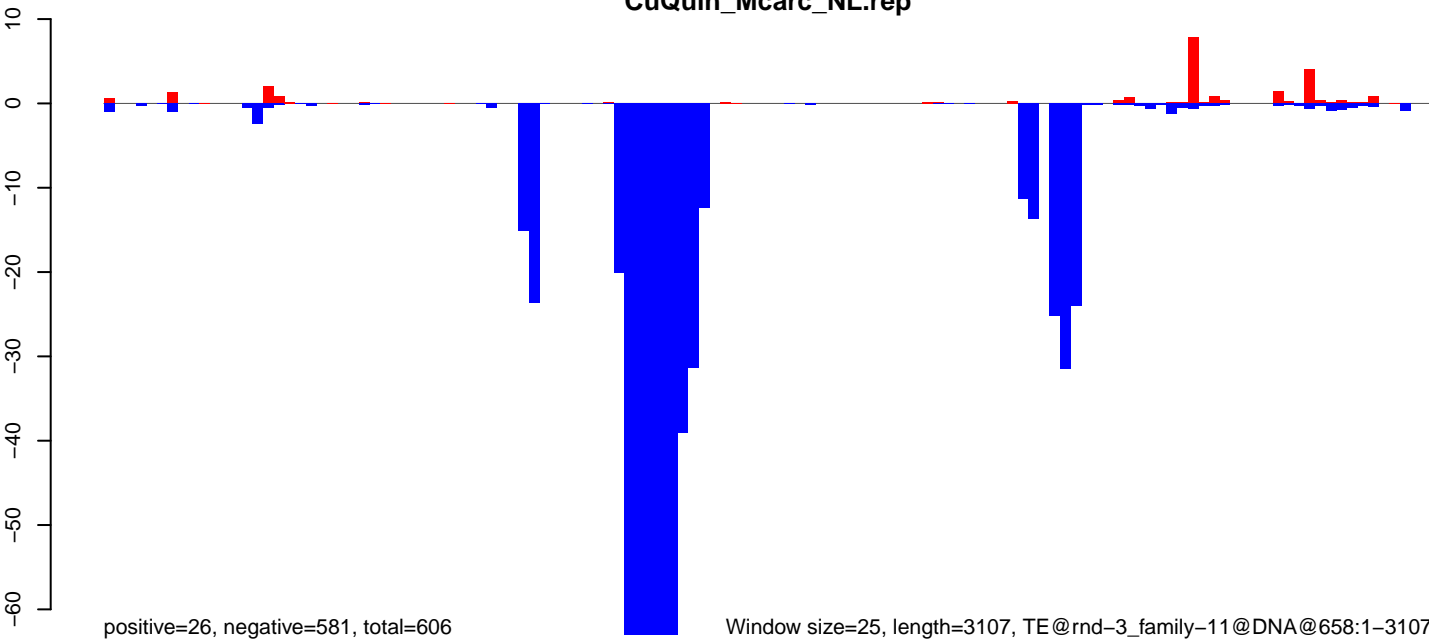
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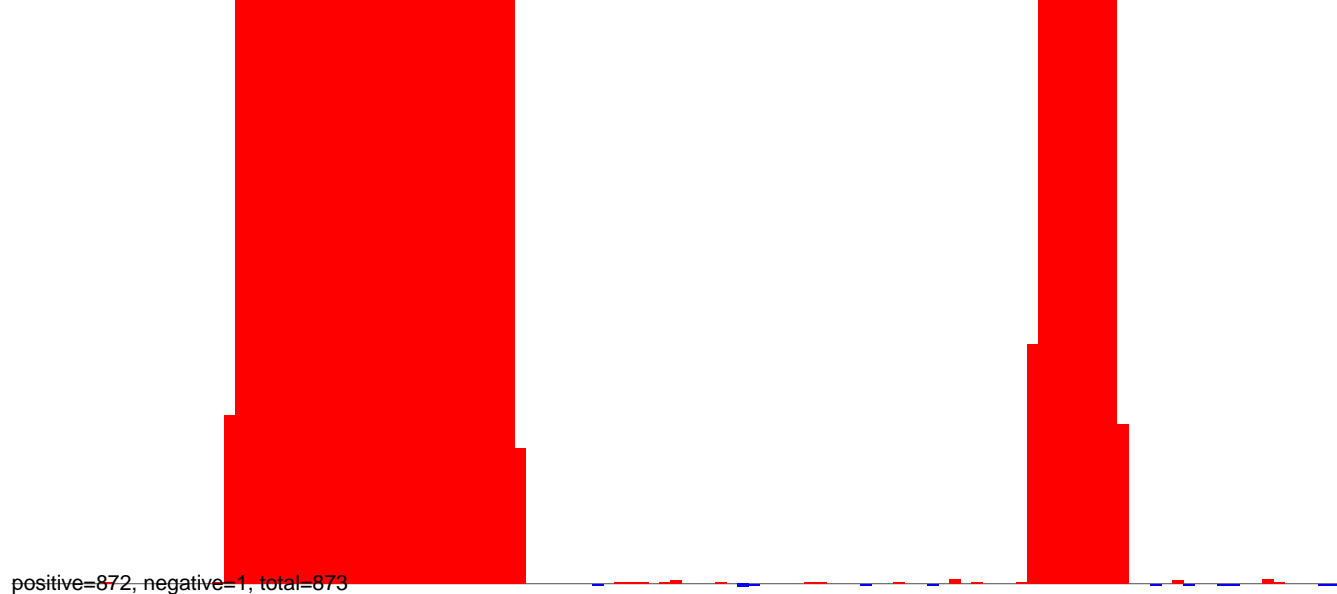
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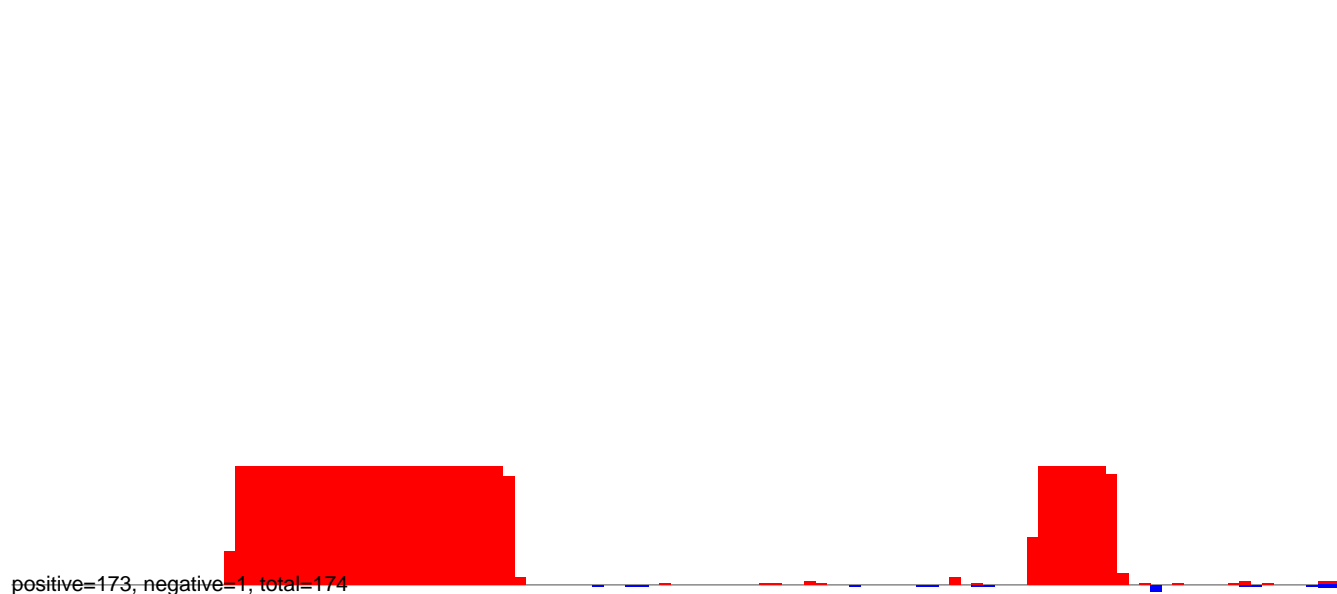
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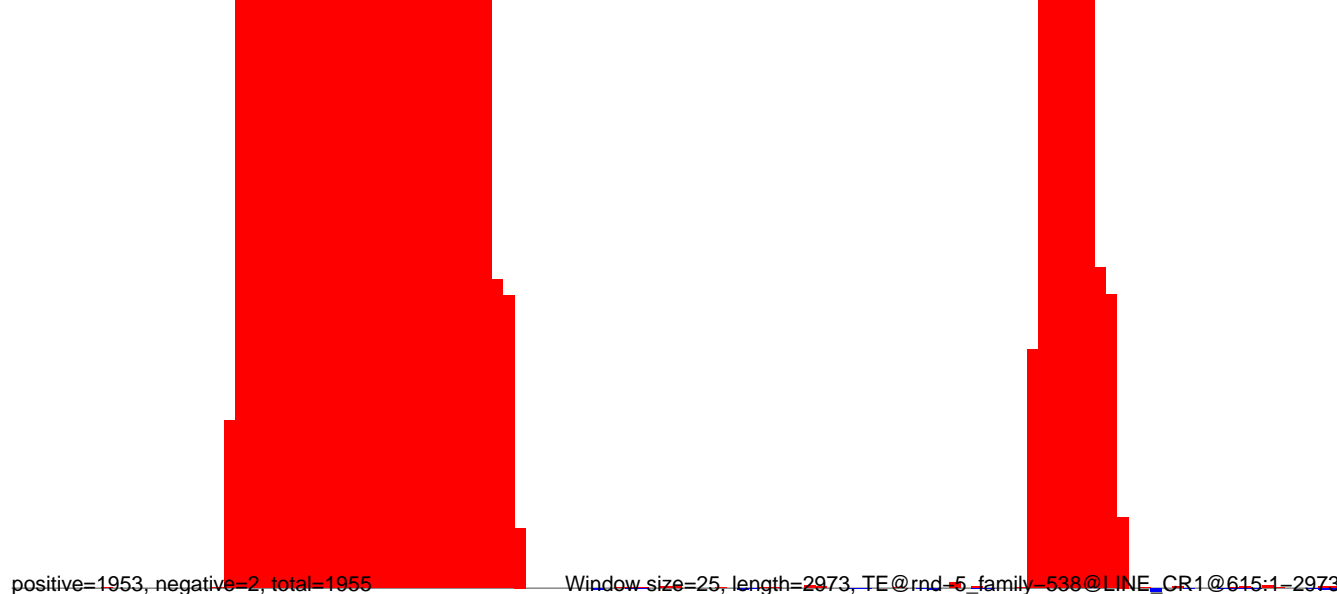
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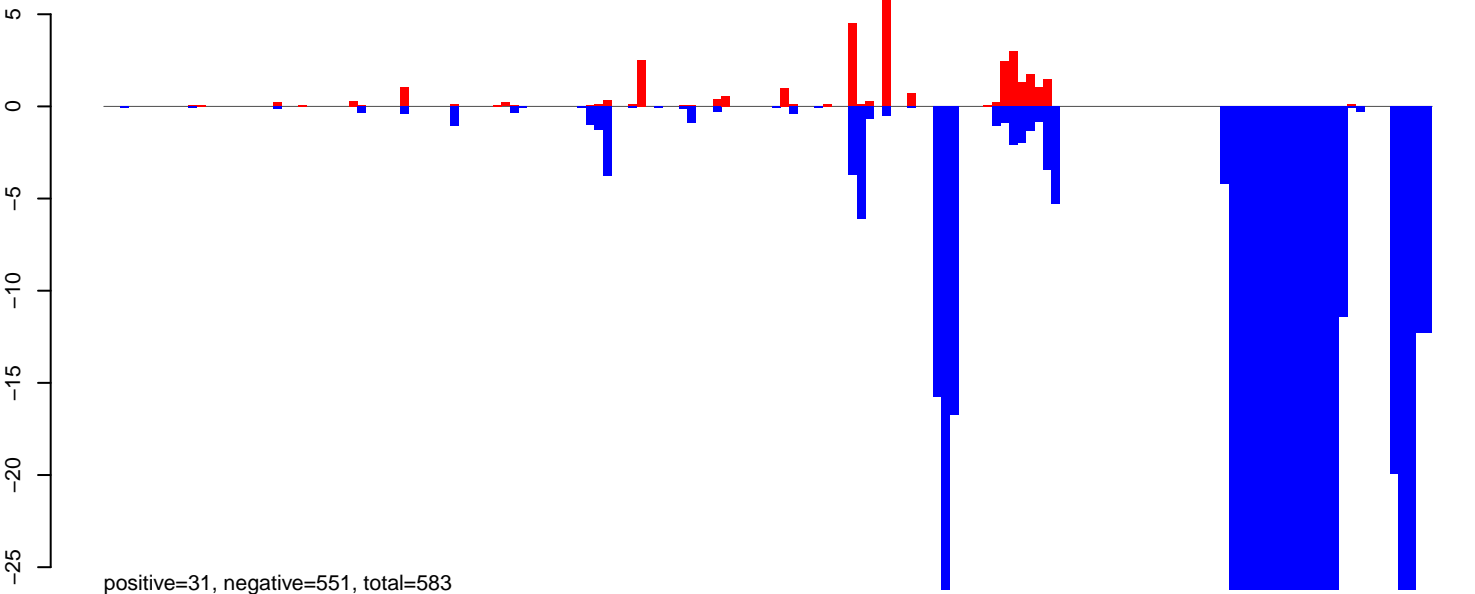
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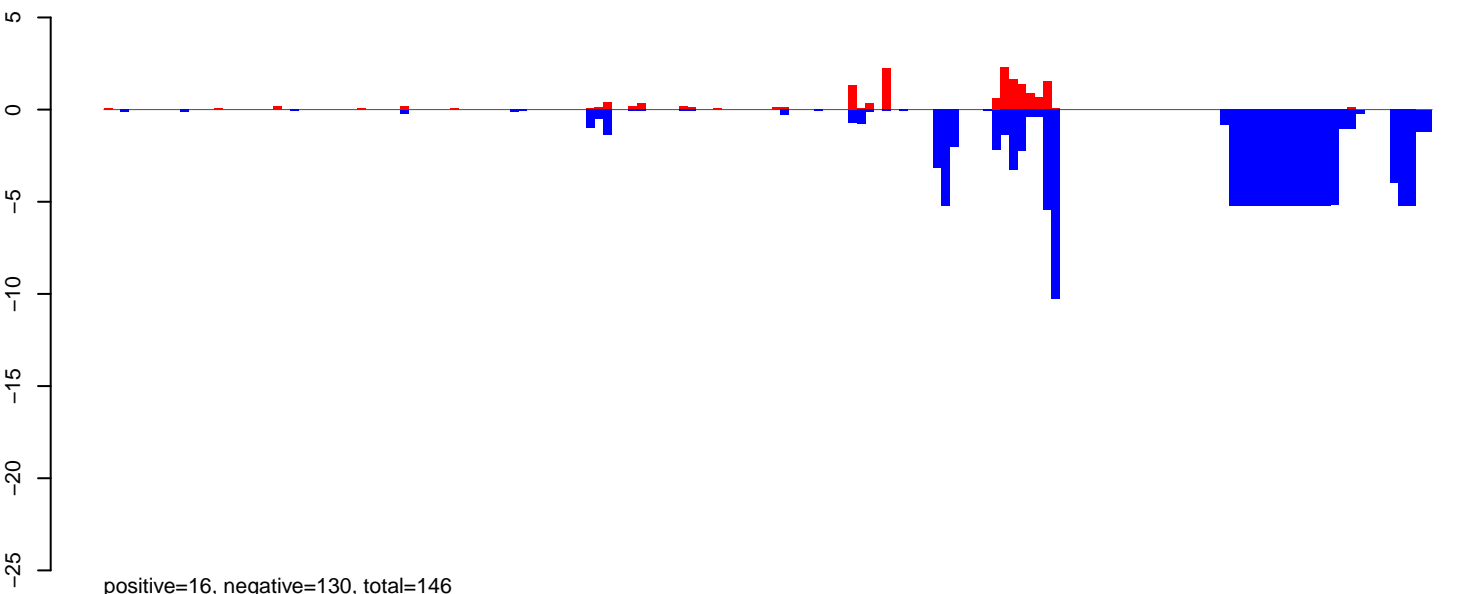
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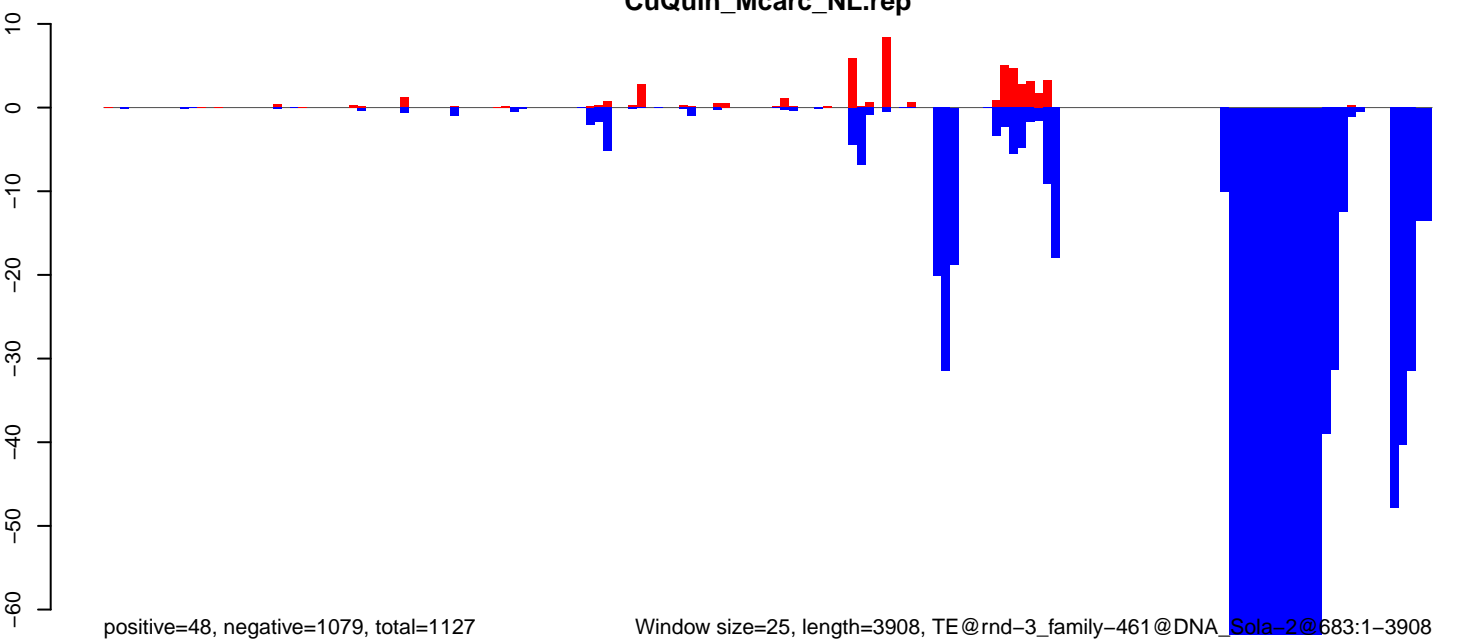
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CuQuin_Mcarc_NL.24_35.rep

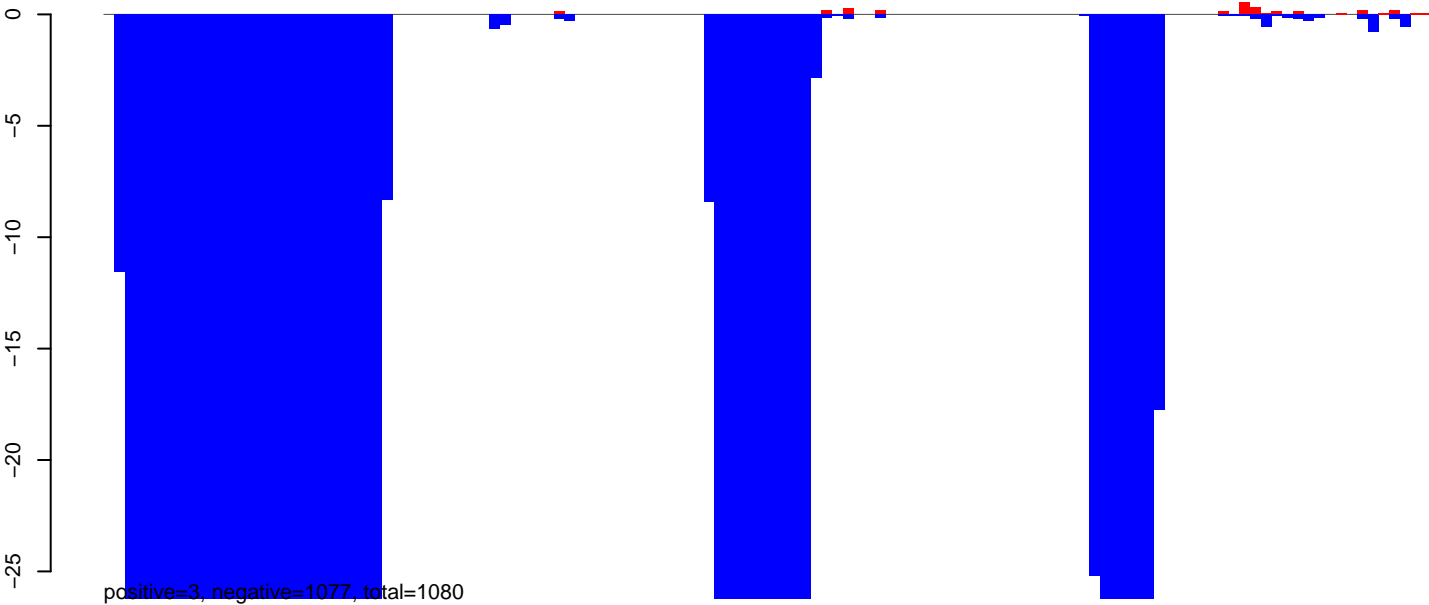


CuQuin_Mcarc_NL.rep

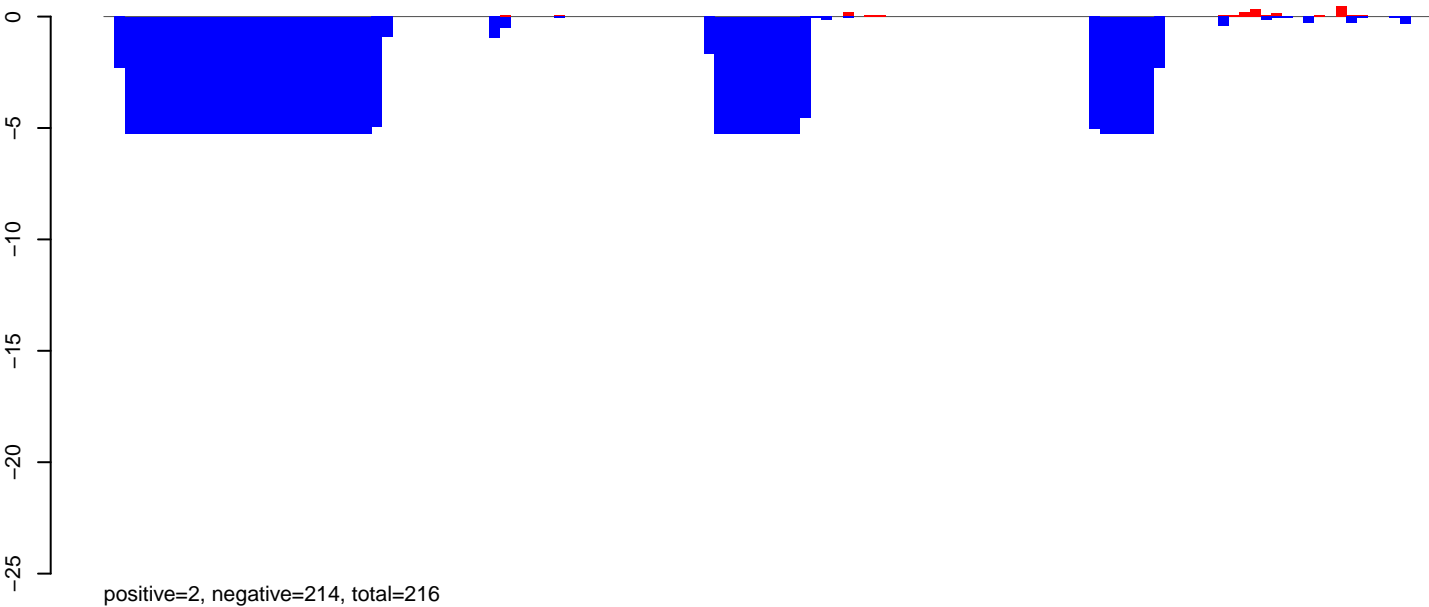


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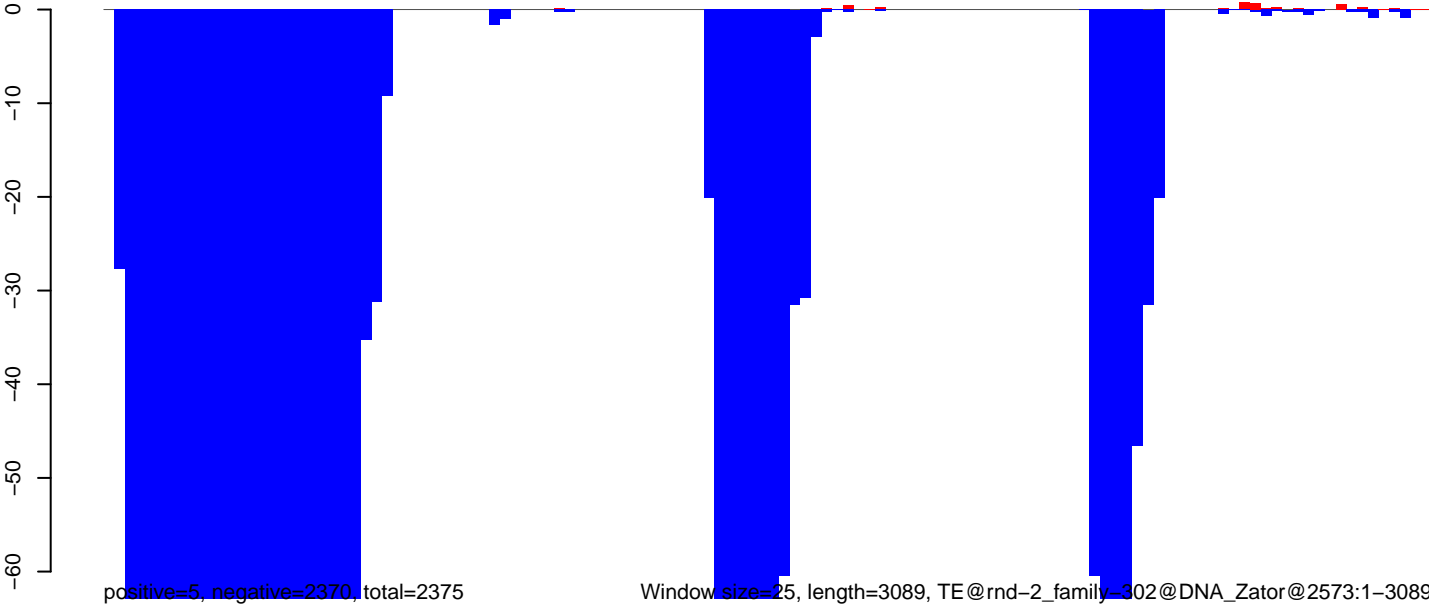
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CuQuin_Mcarc_NL.24_35.rep

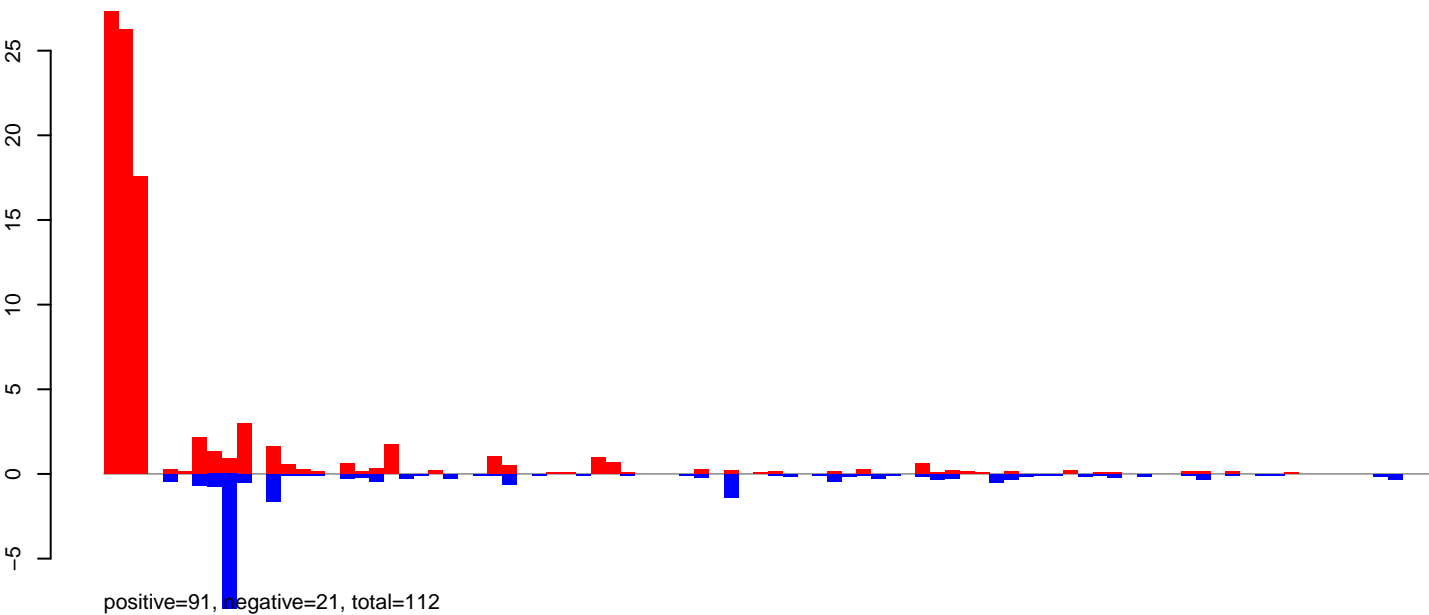


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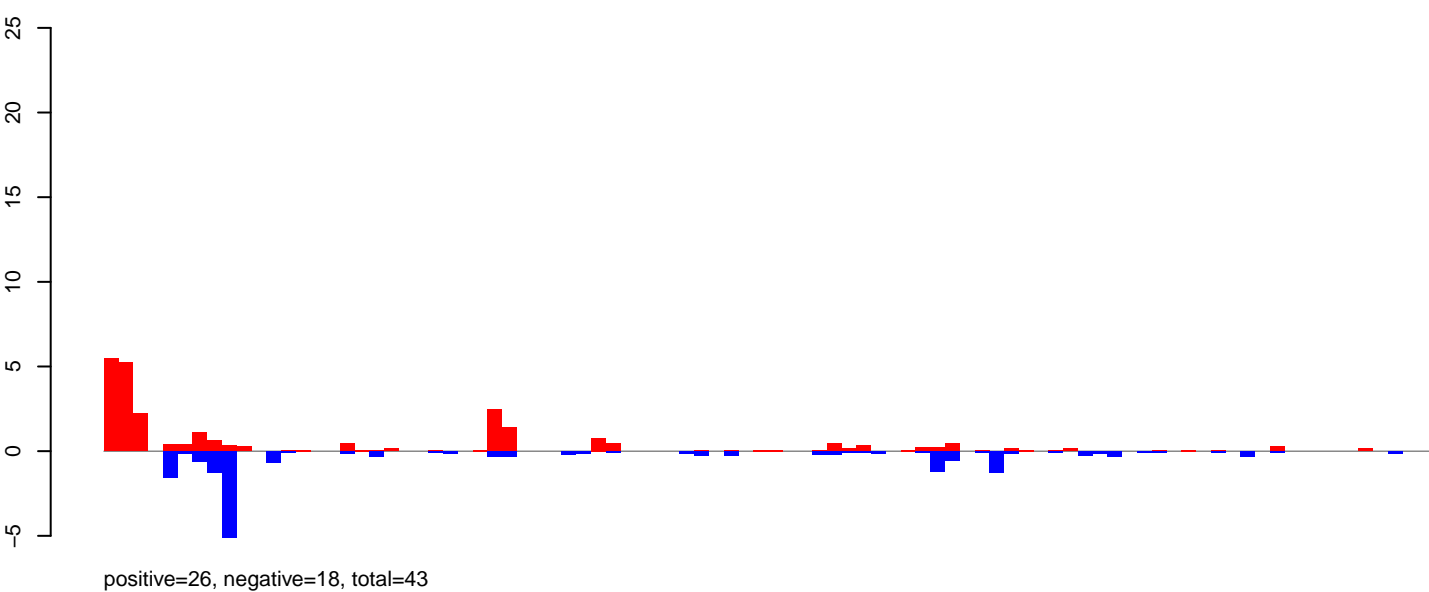


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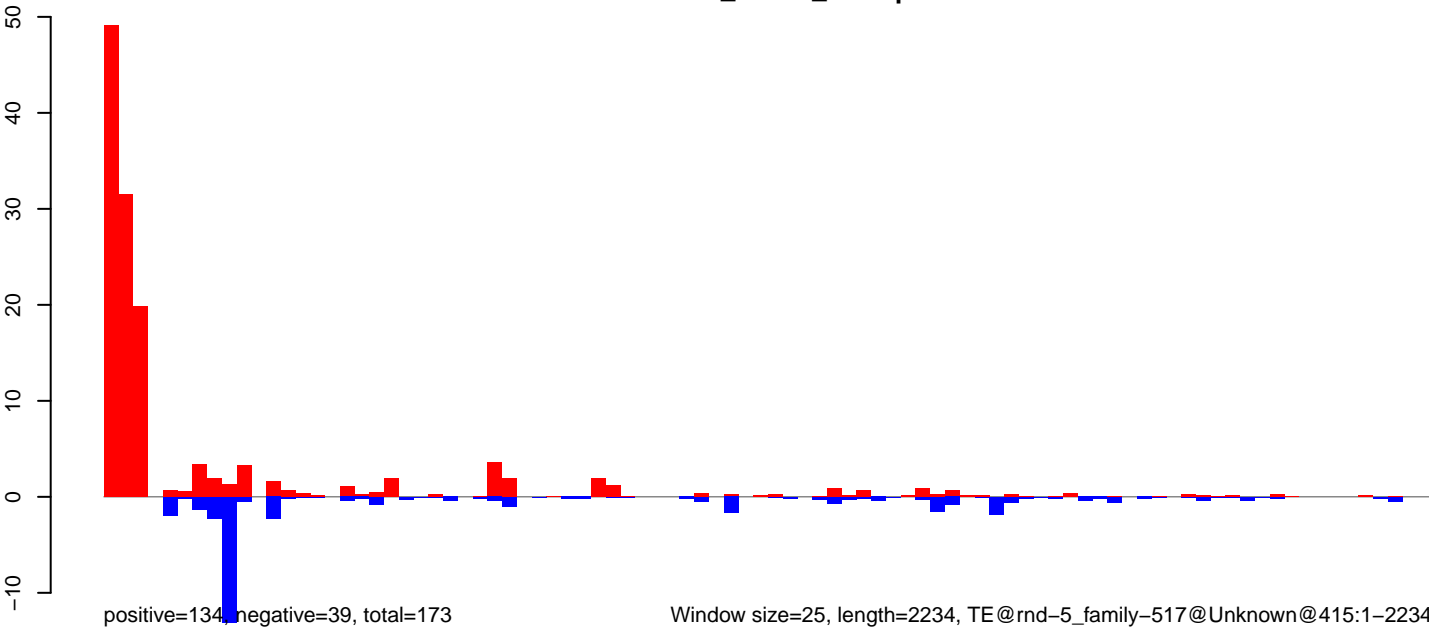
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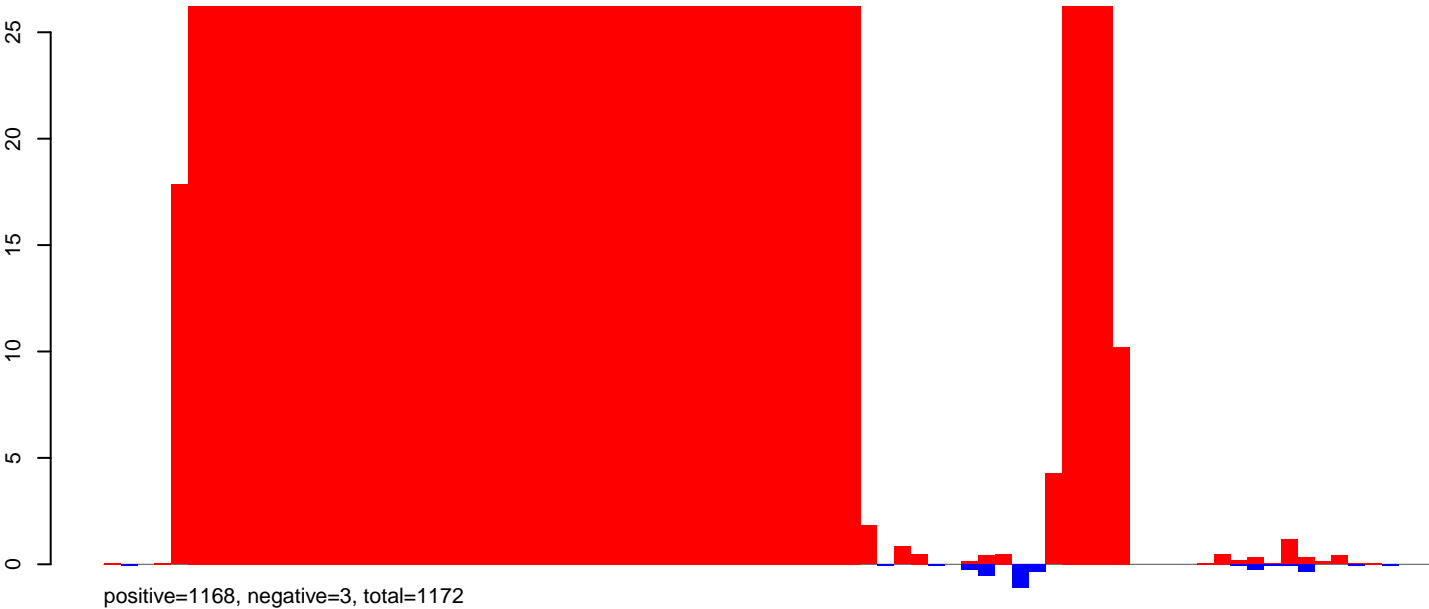


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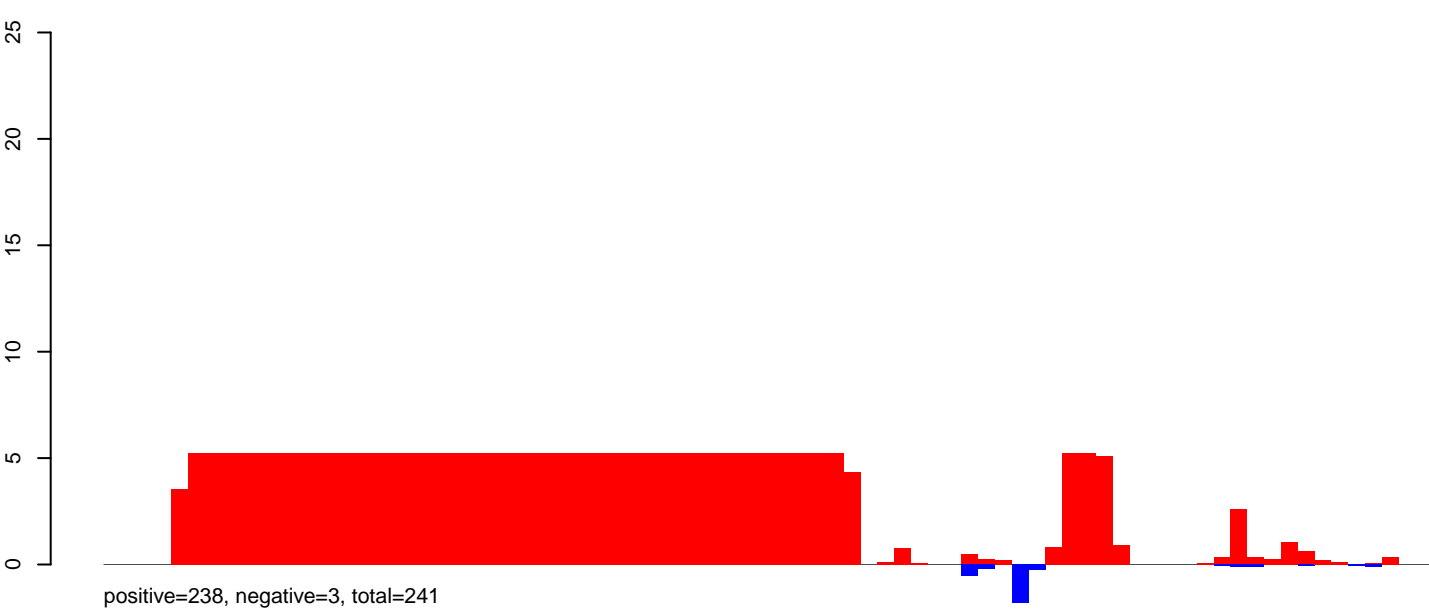


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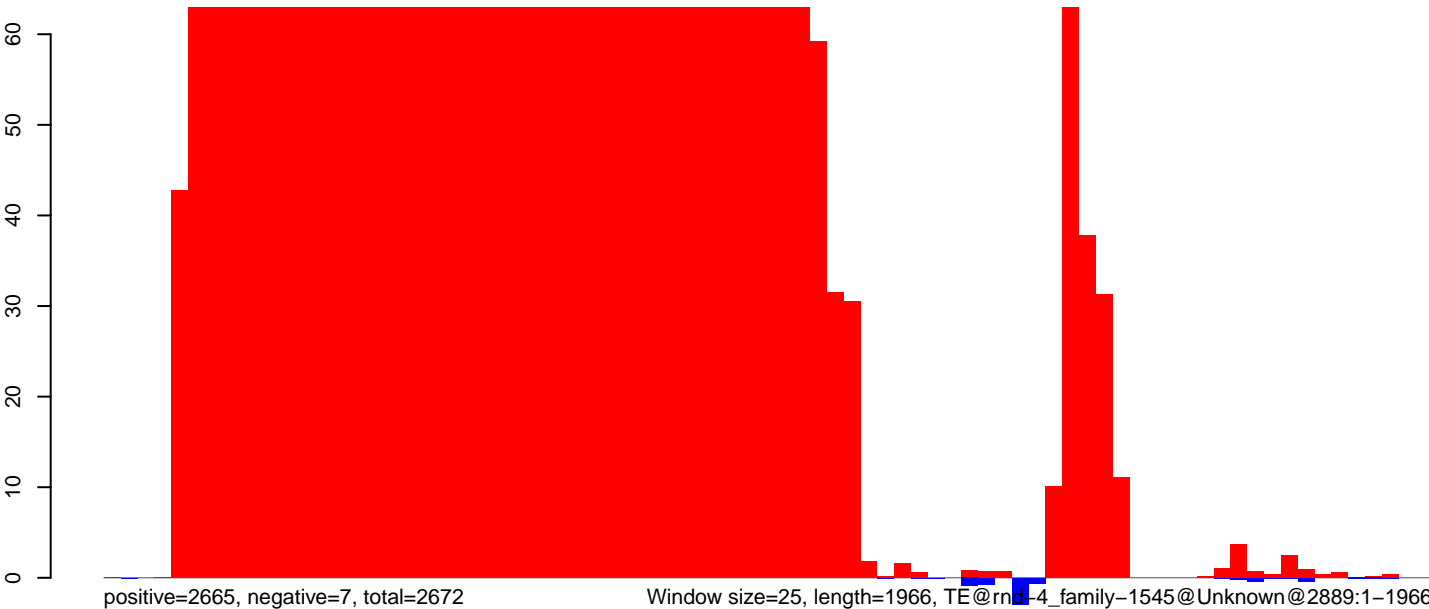
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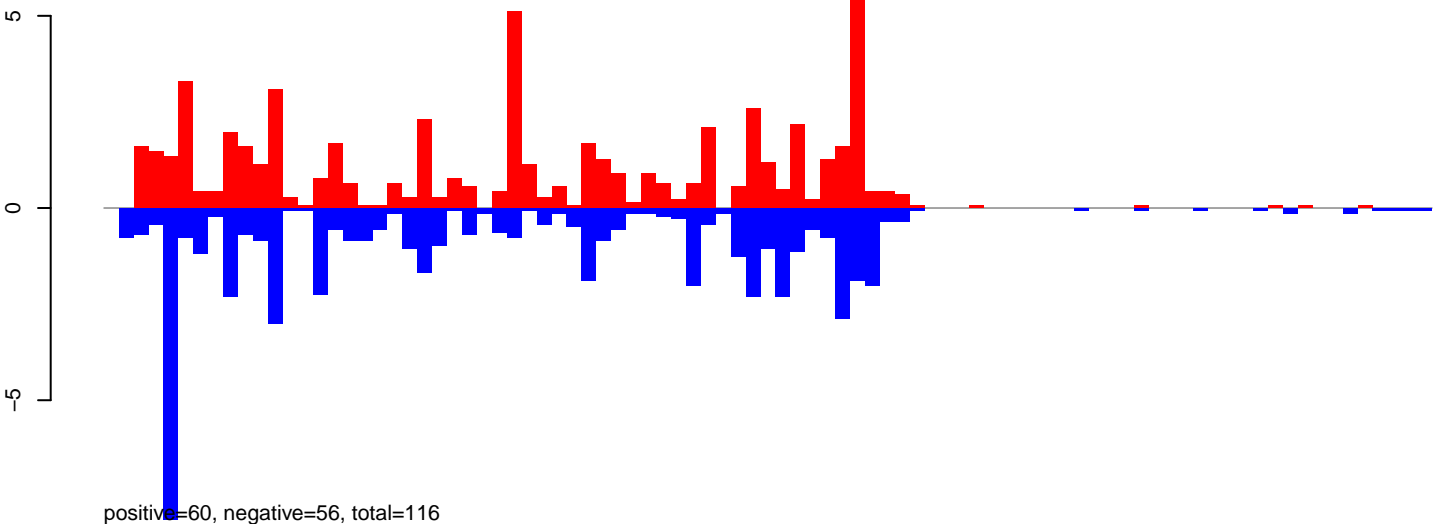


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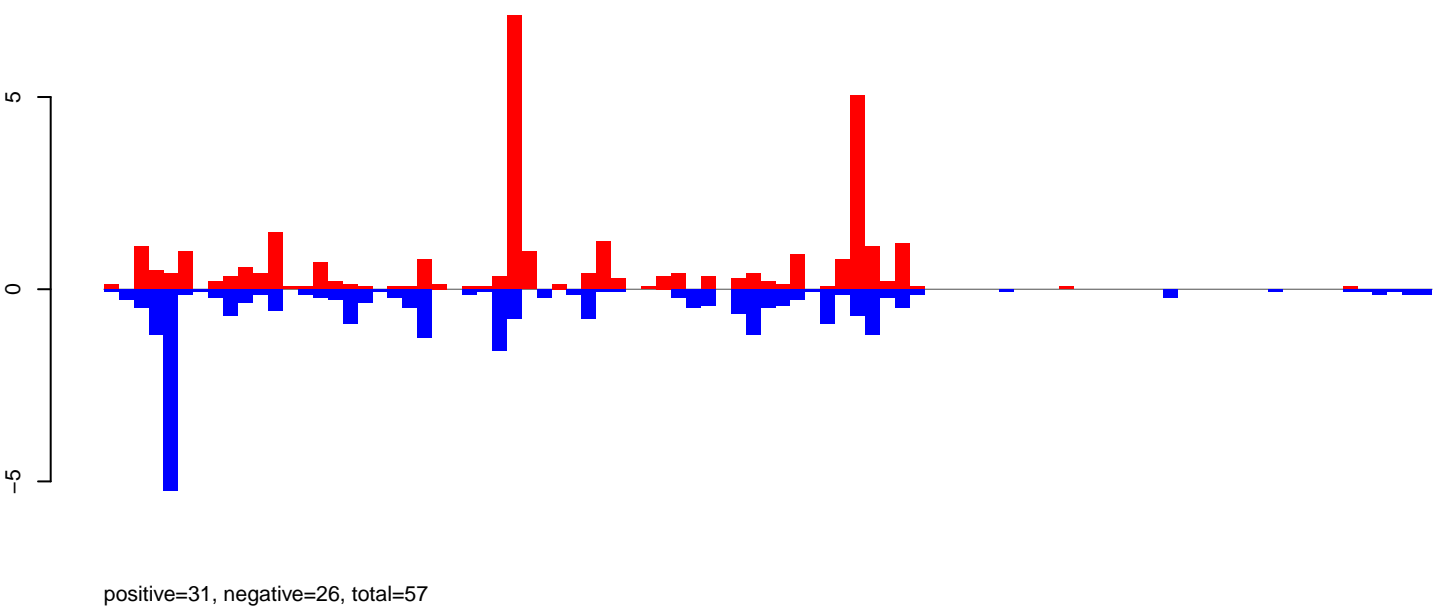


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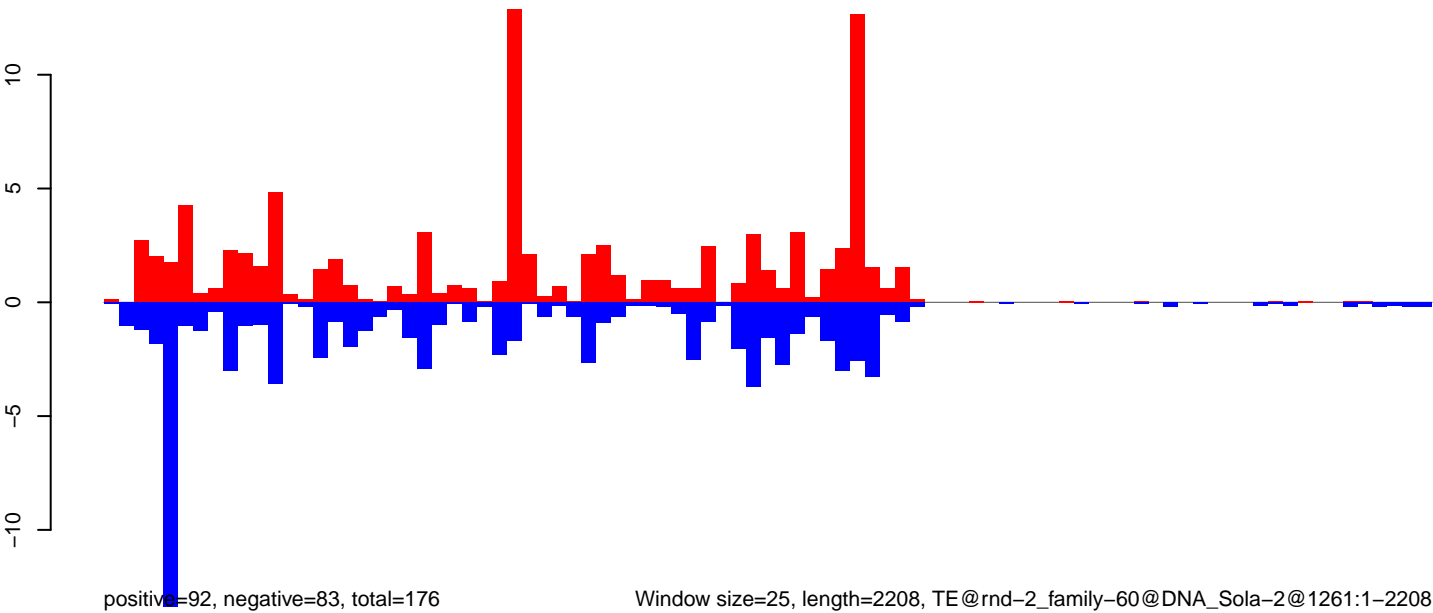
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CuQuin_Mcarc_NL.24_35.rep



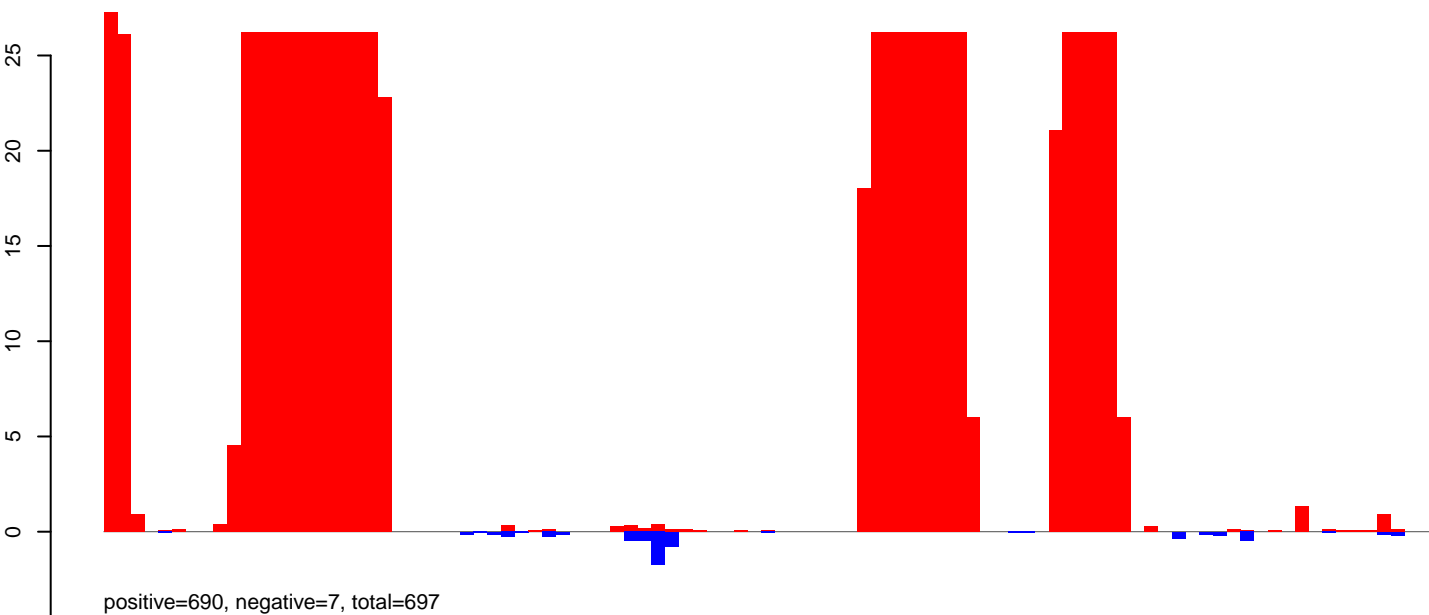
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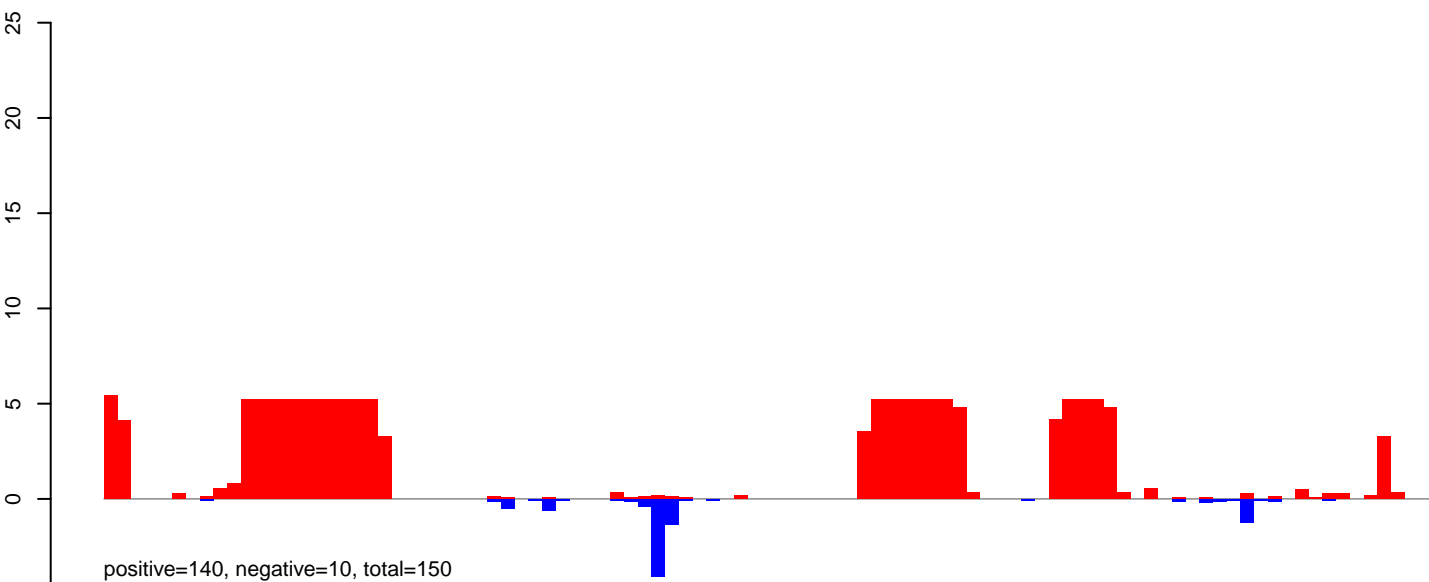
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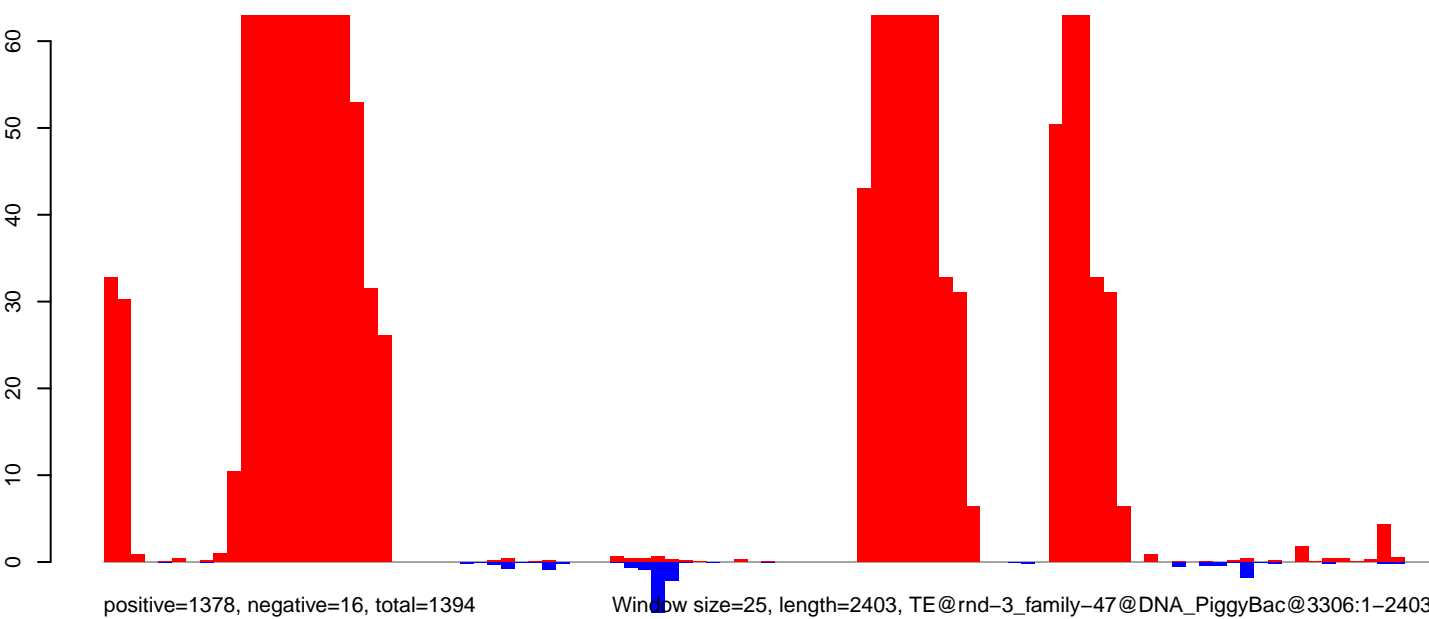
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CuQuin_Mcarc_NL.24_35.rep



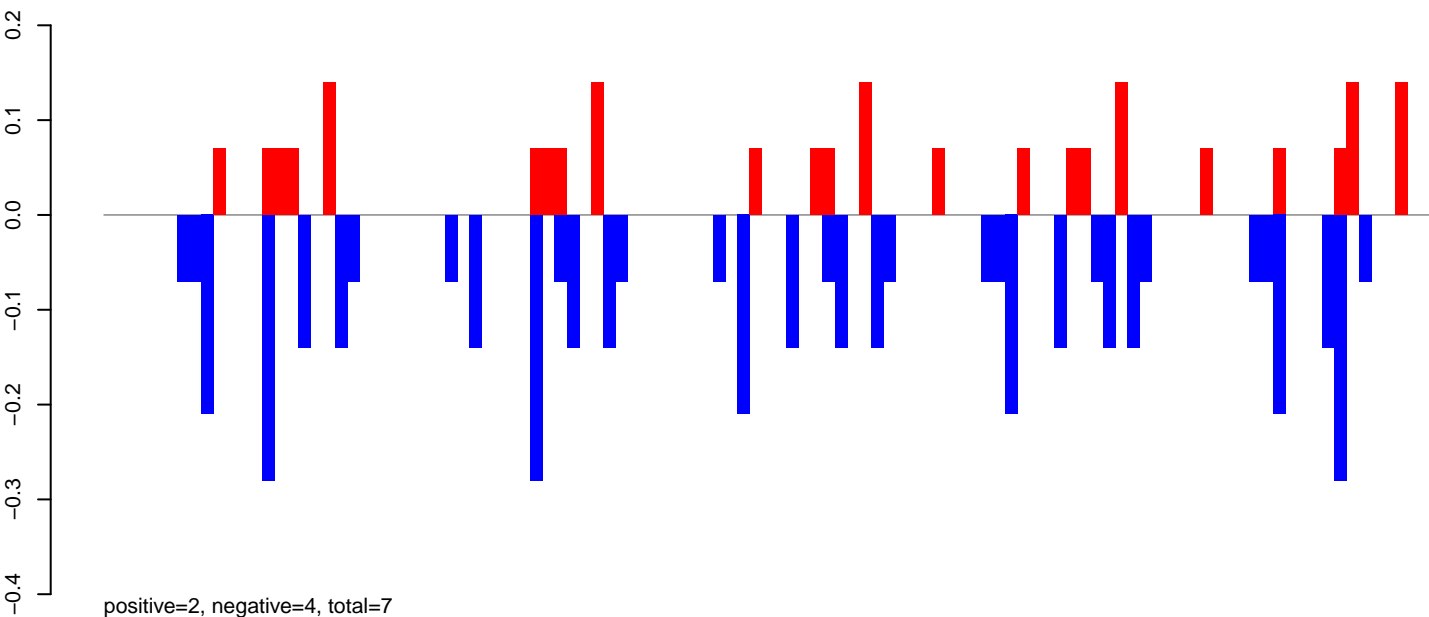
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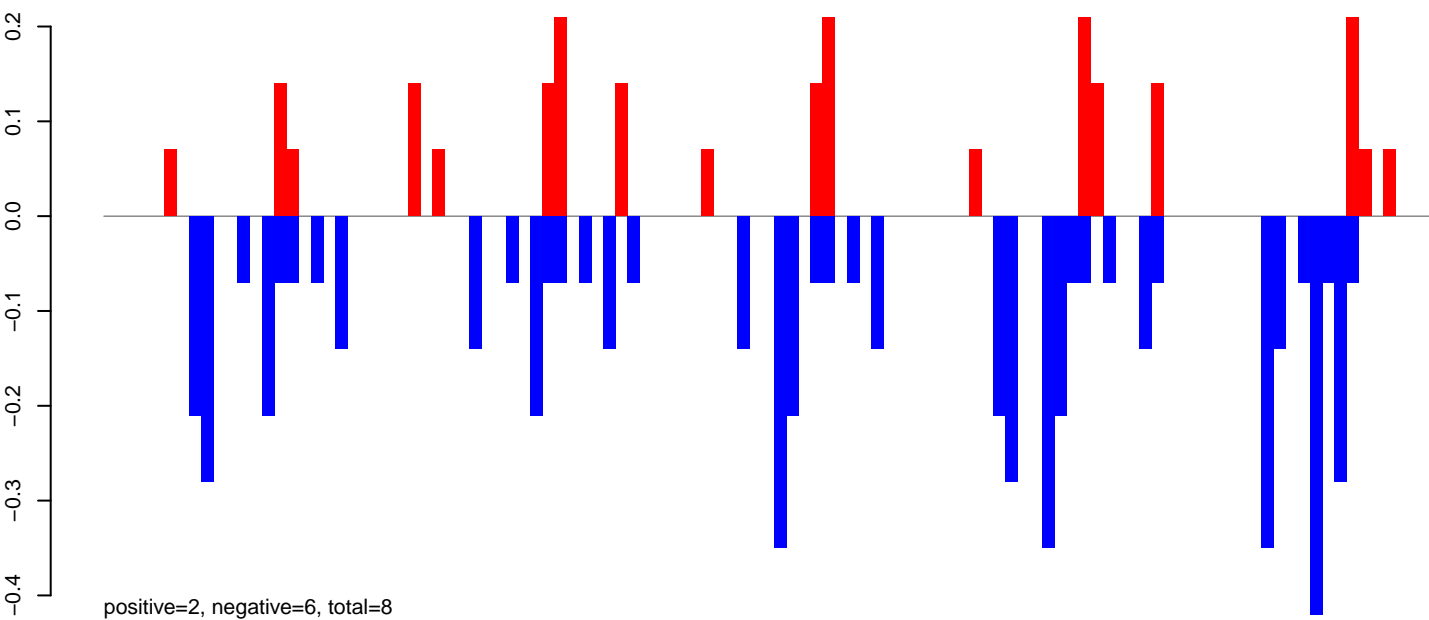
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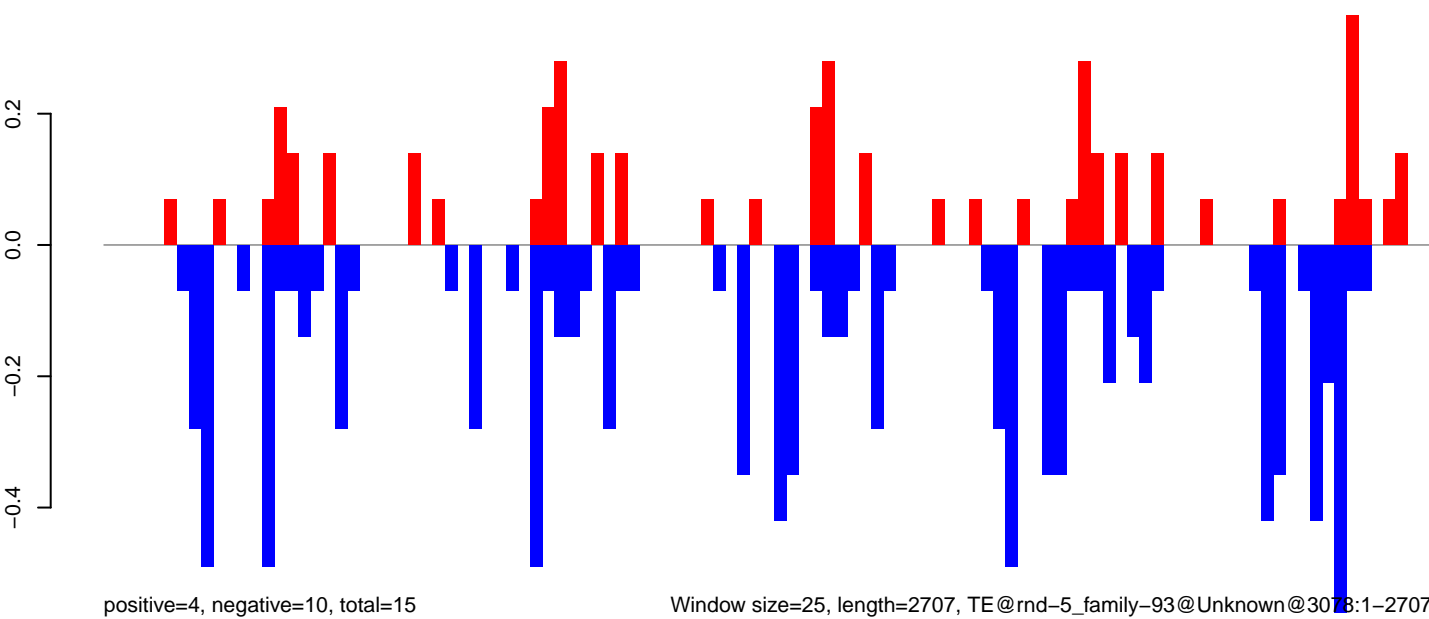
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CuQuin_Mcarc_NL.24_35.rep



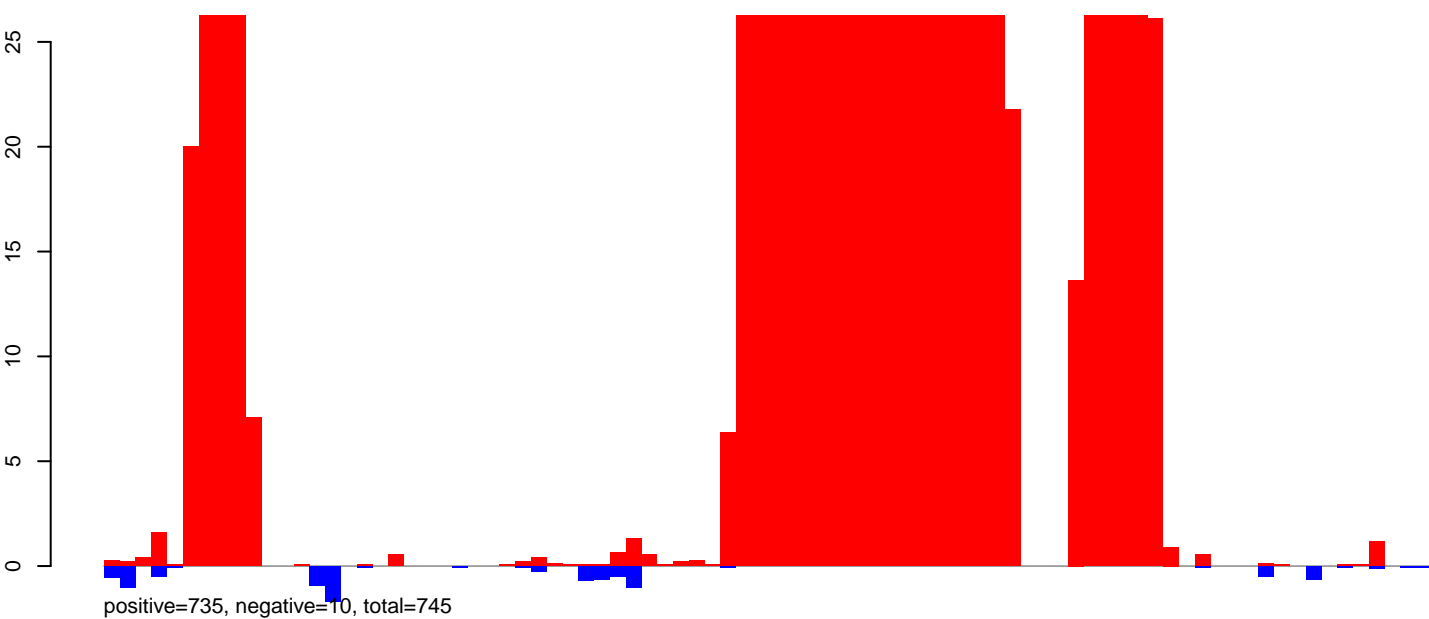
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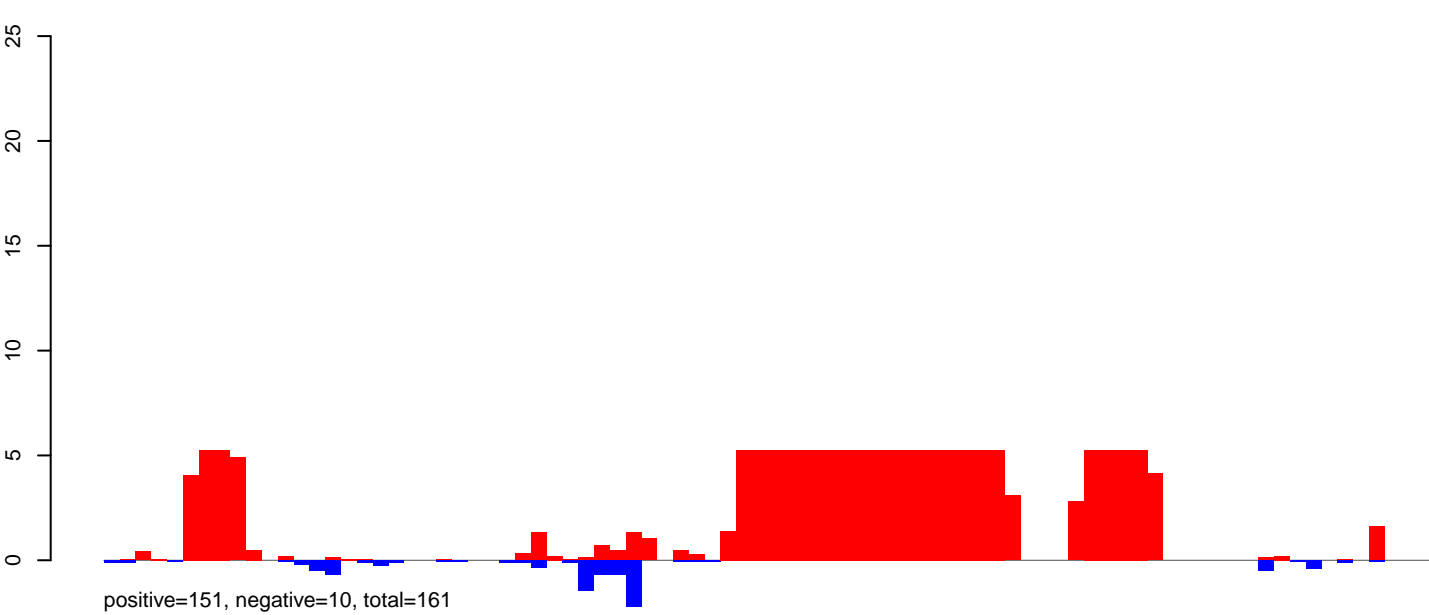
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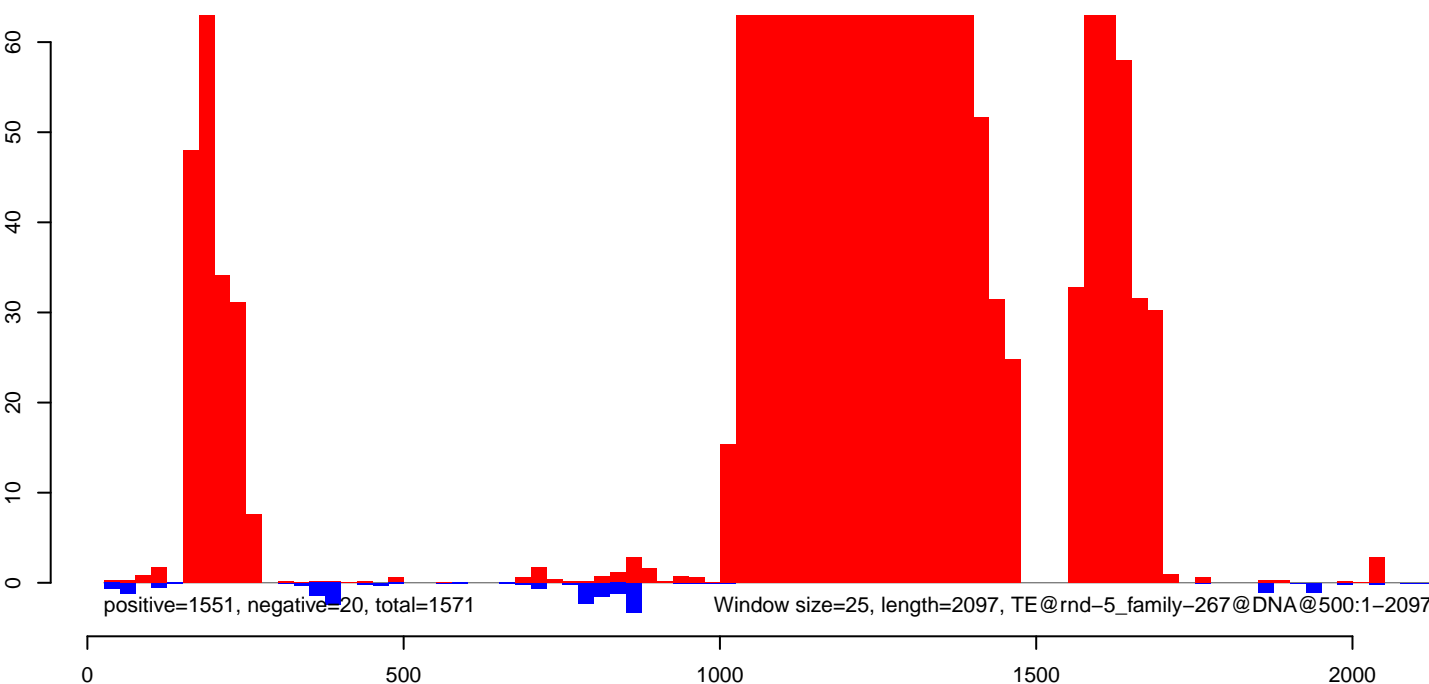
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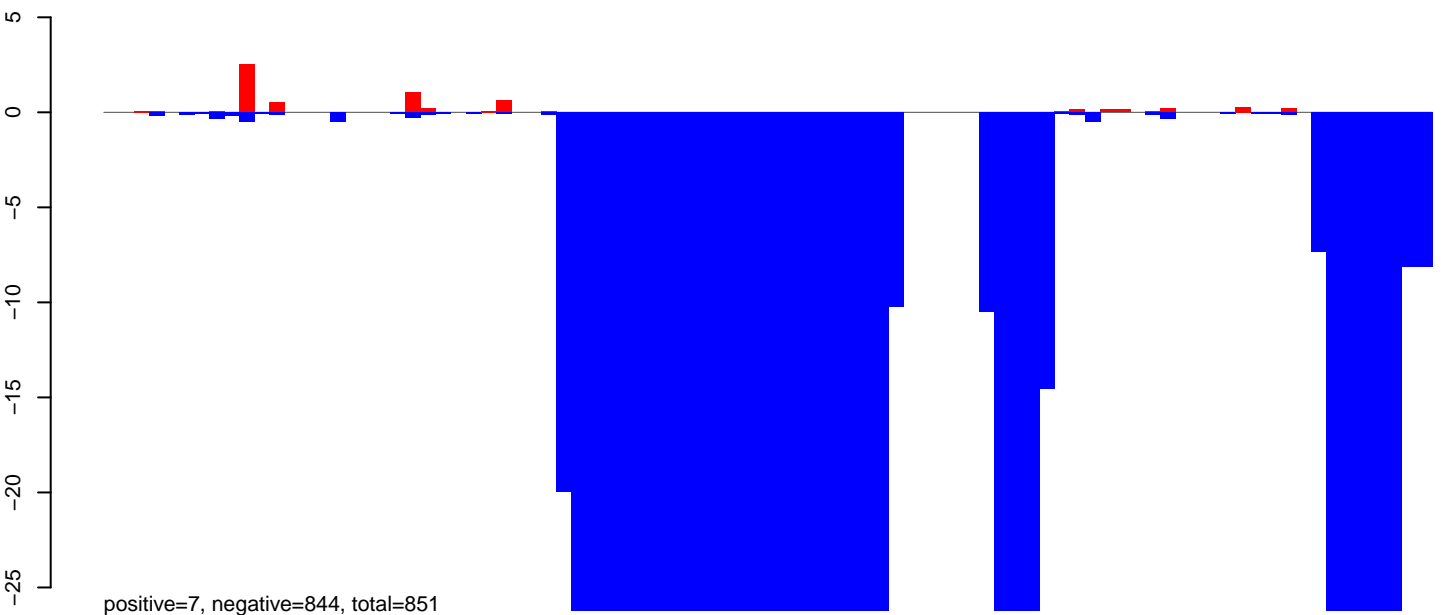
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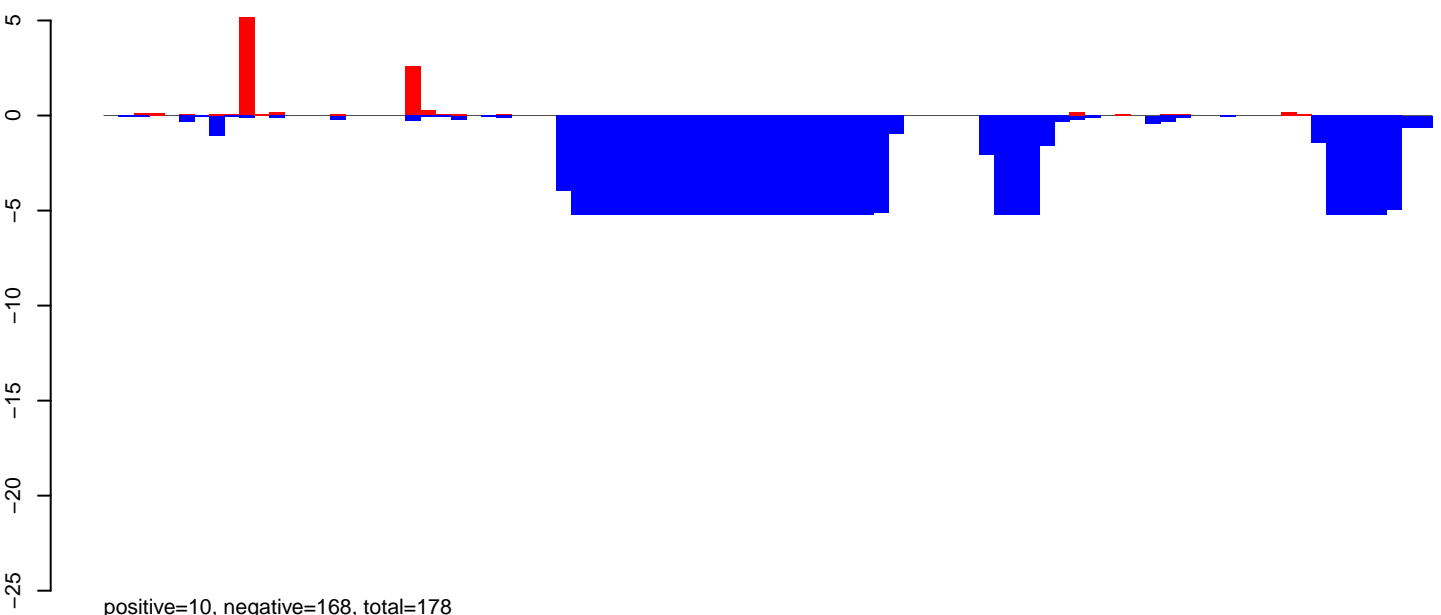
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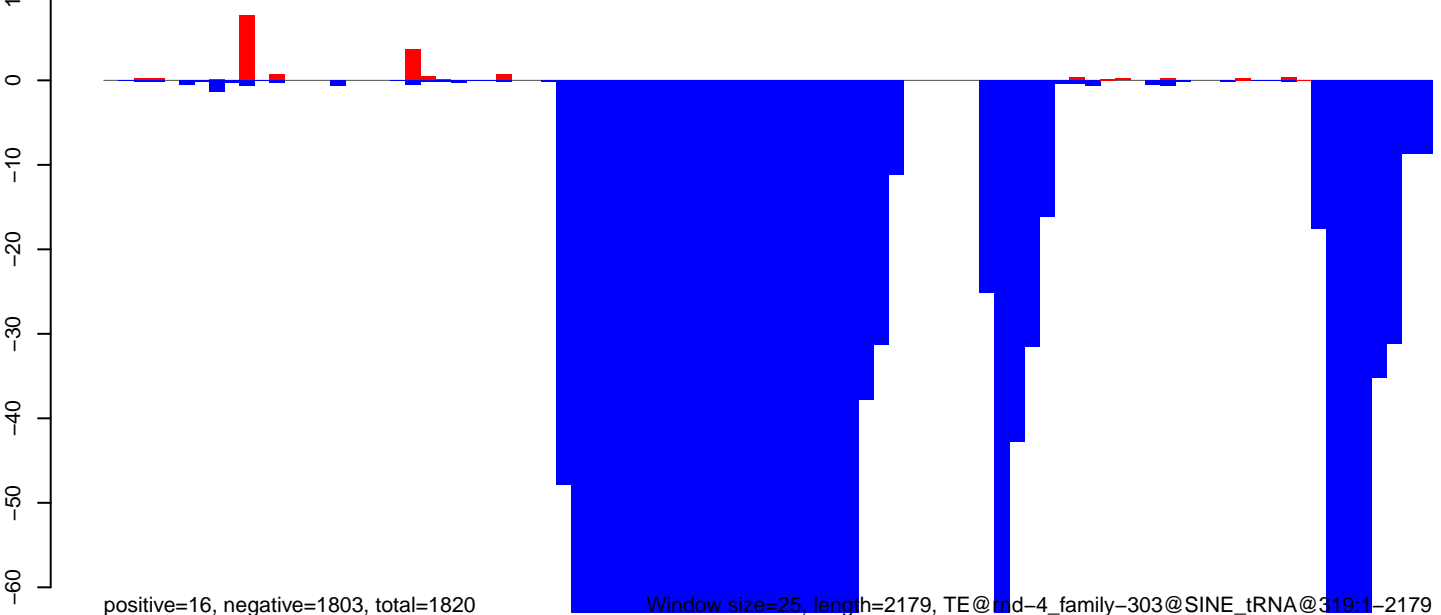
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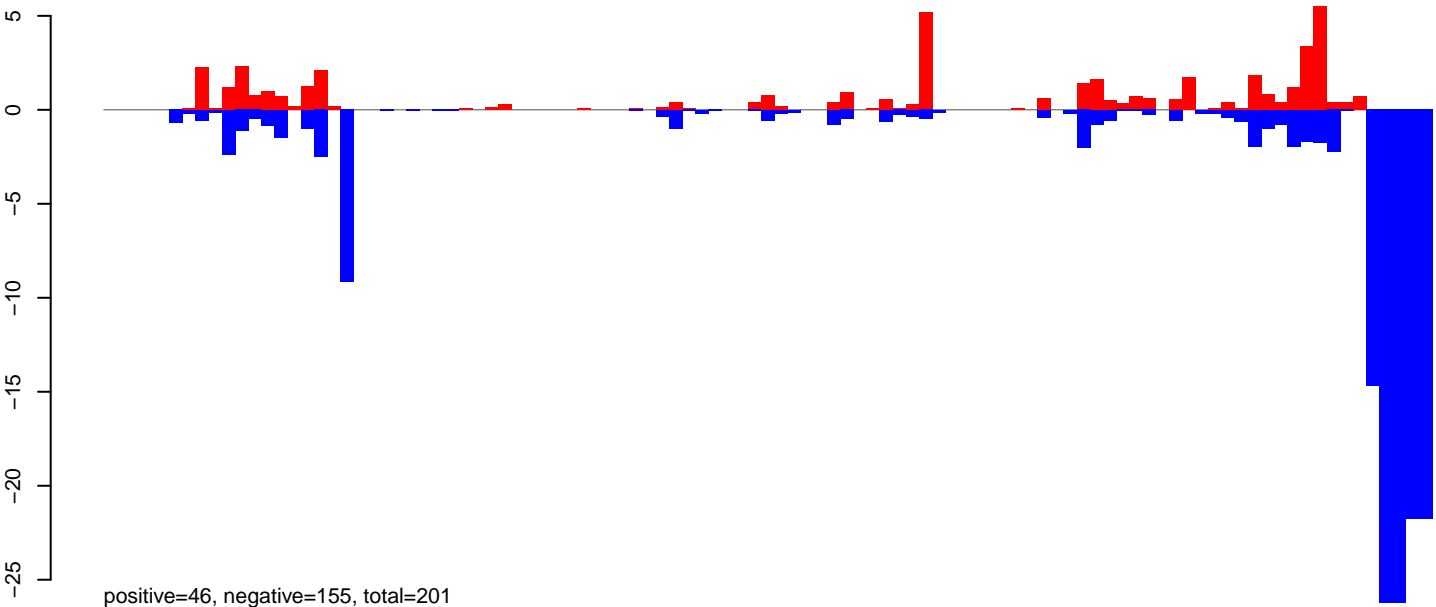


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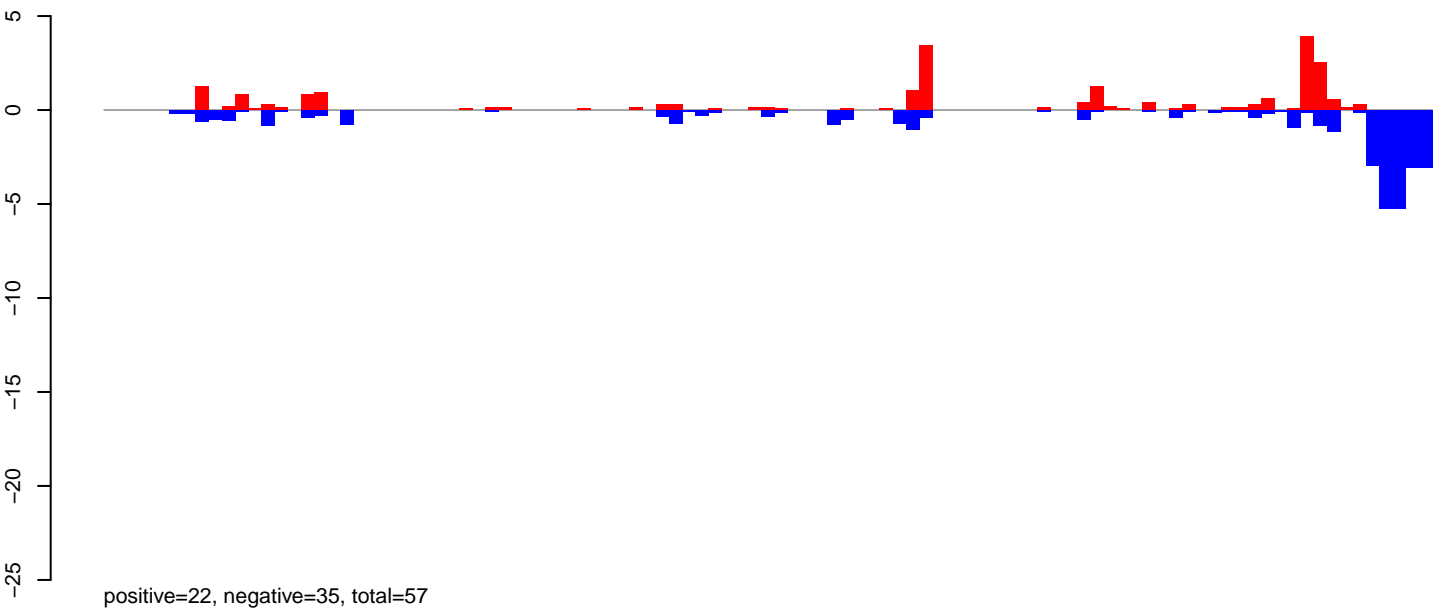


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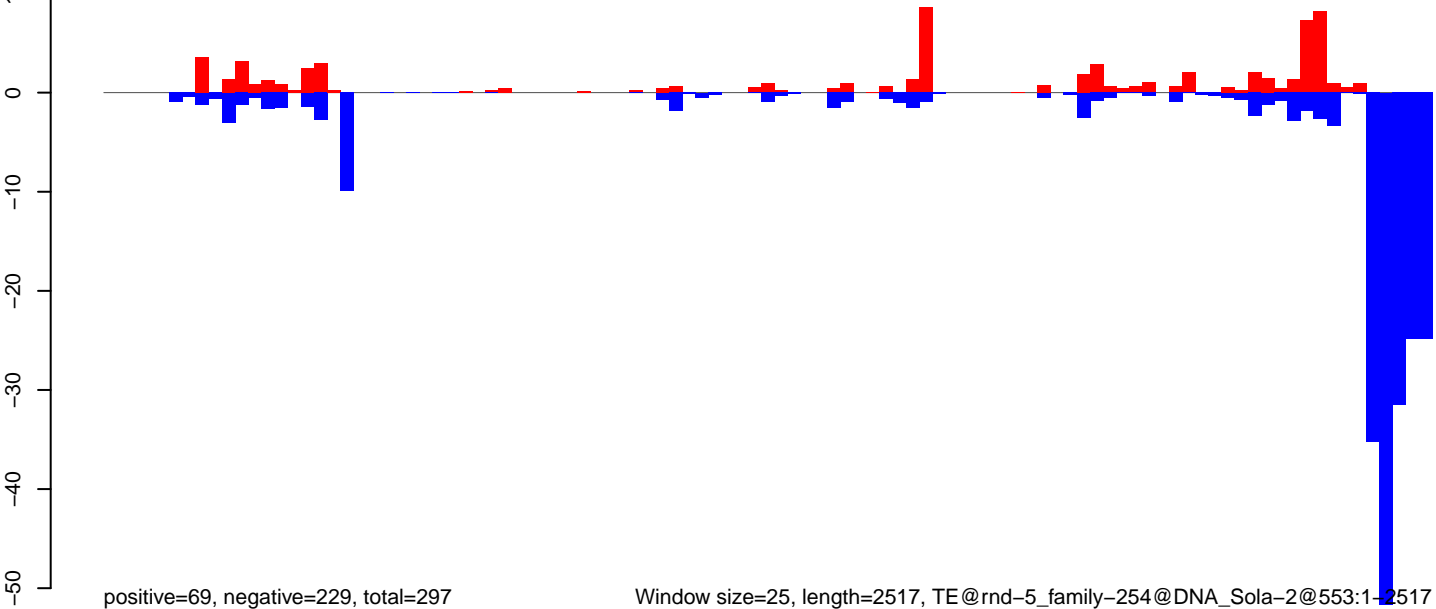
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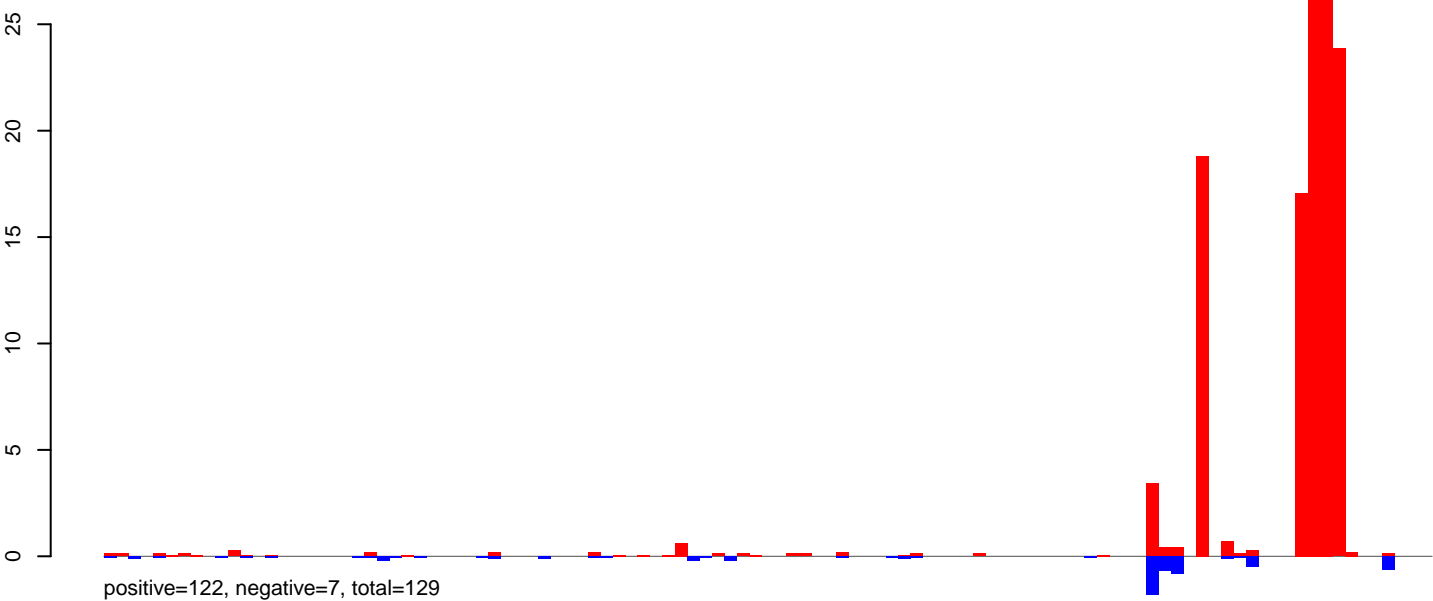
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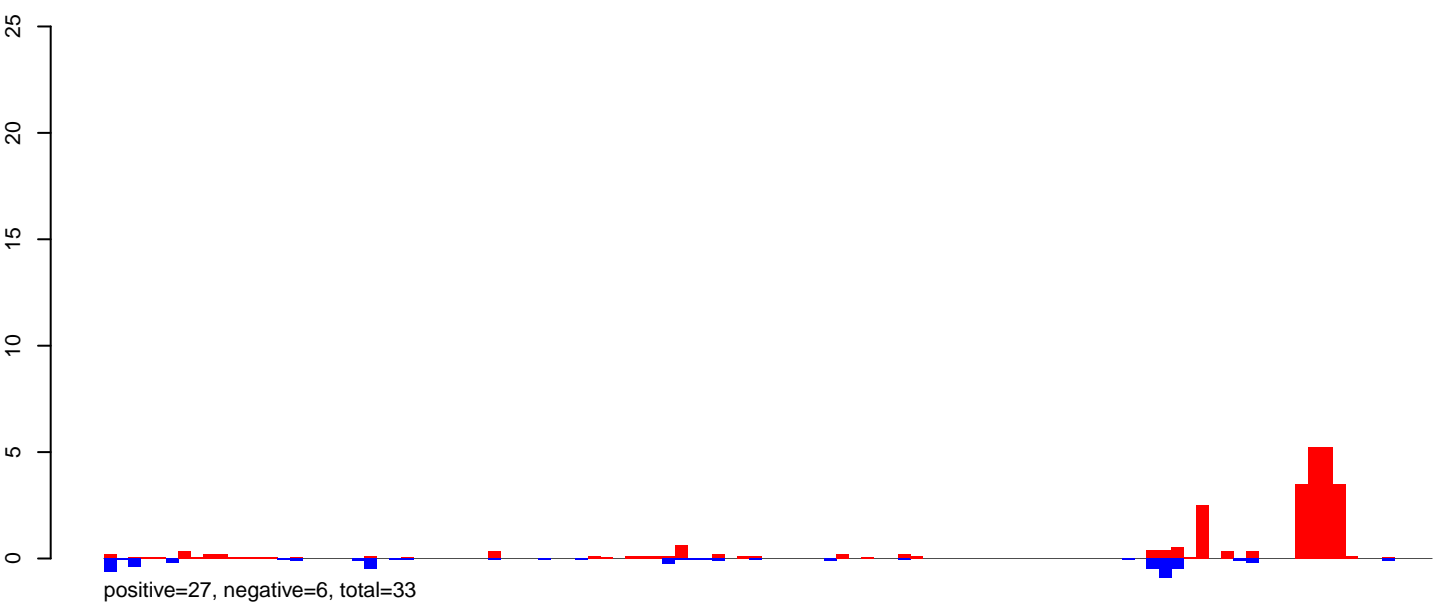
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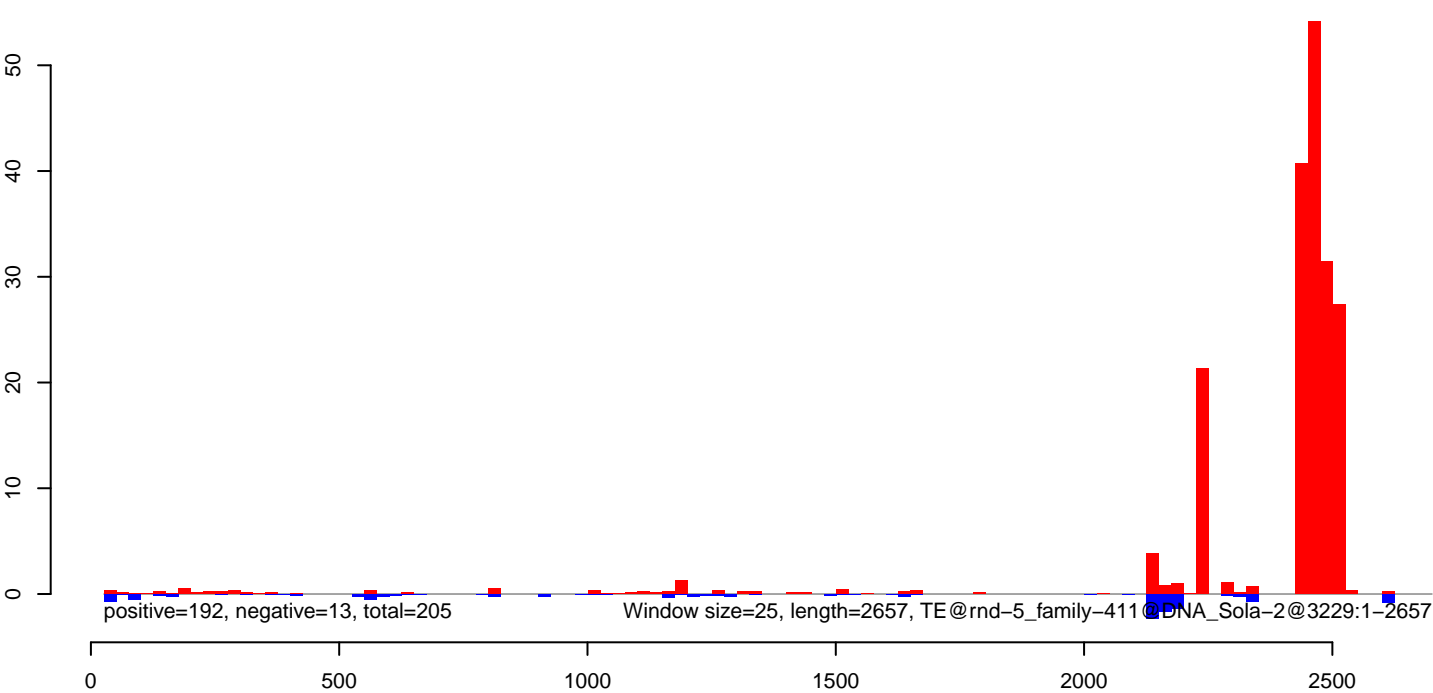
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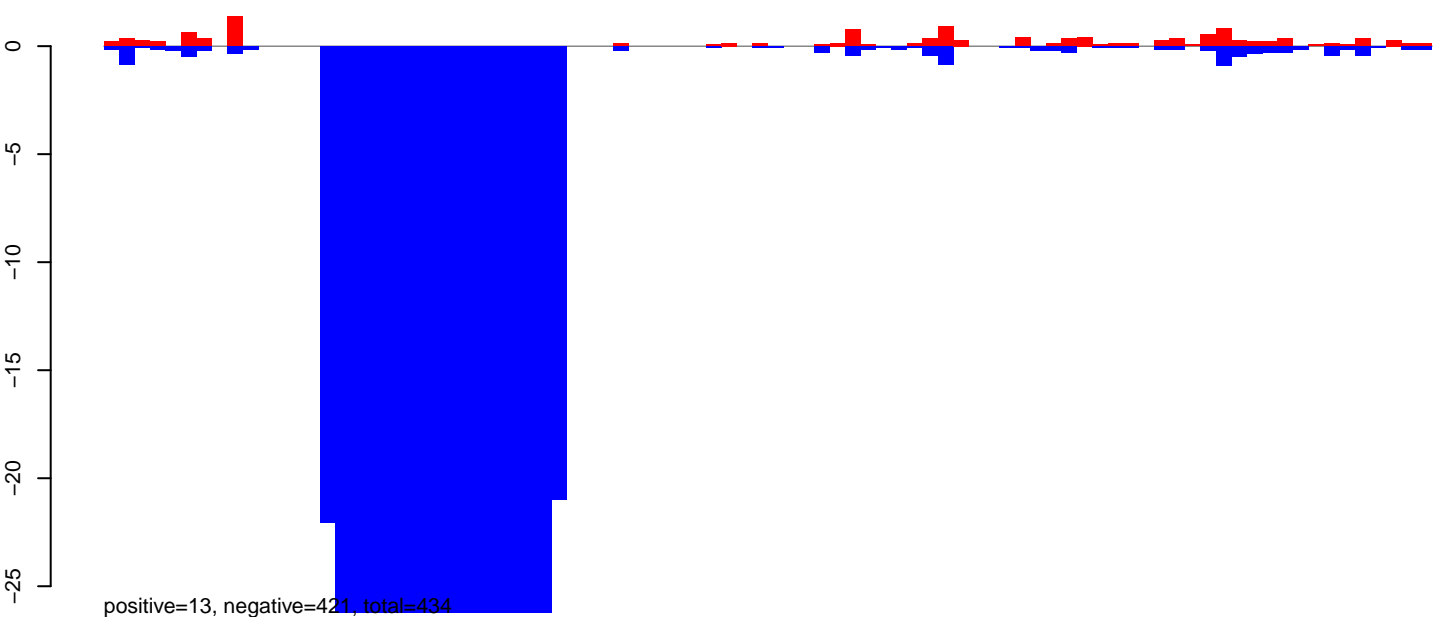
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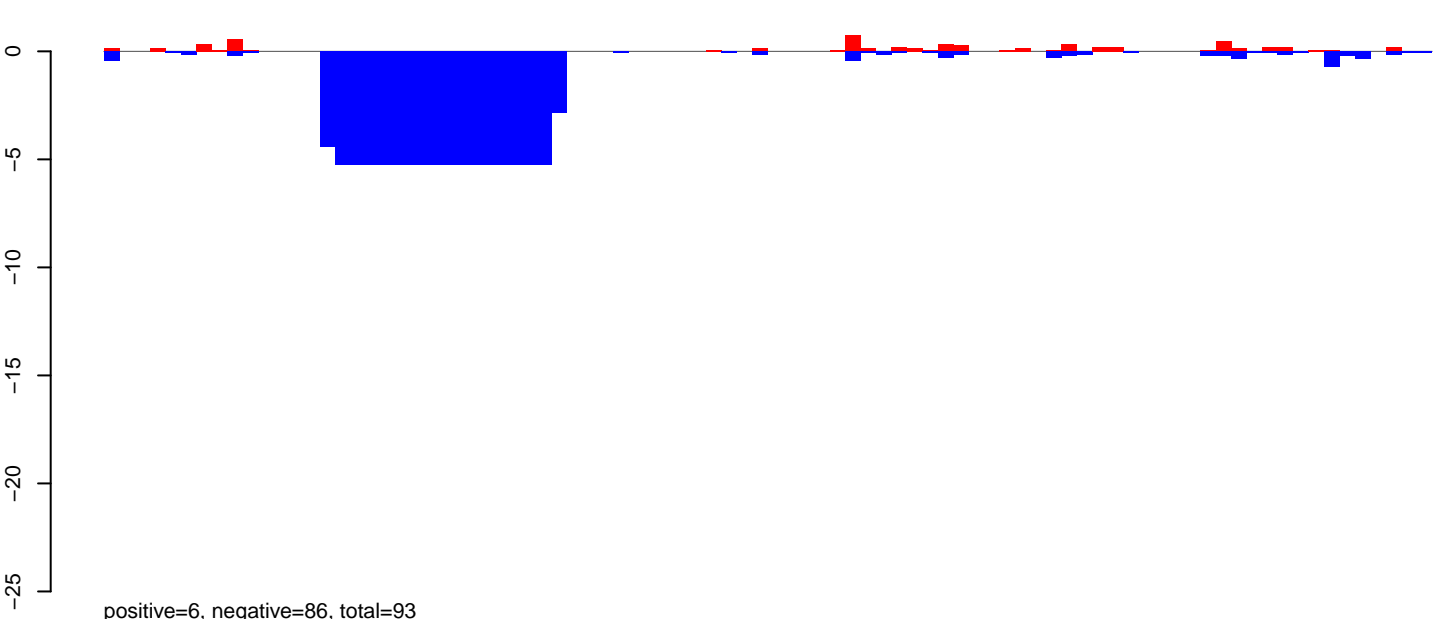
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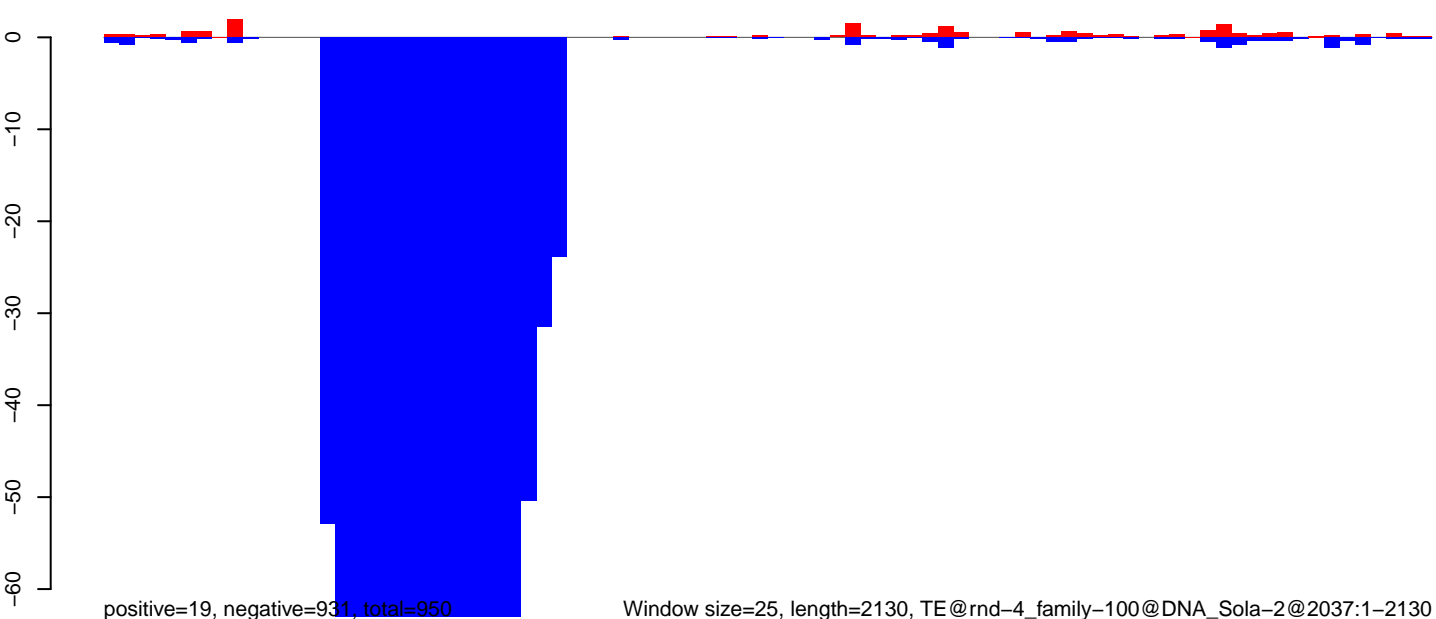
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CuQuin_Mcarc_NL.24_35.rep

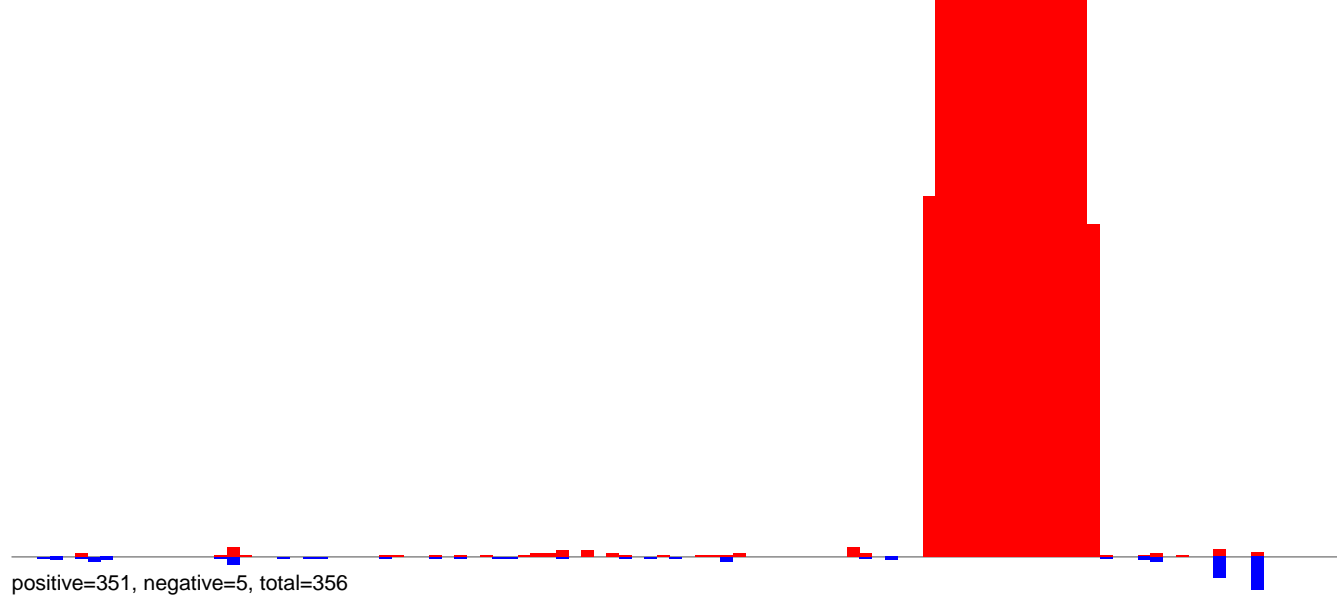


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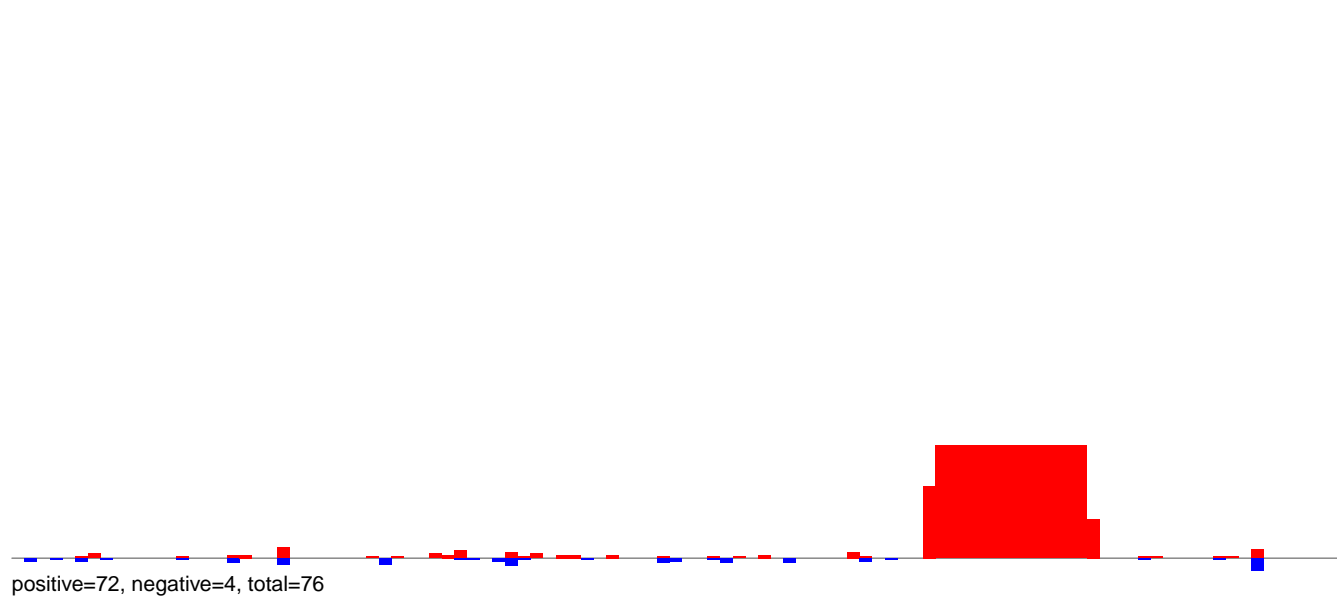


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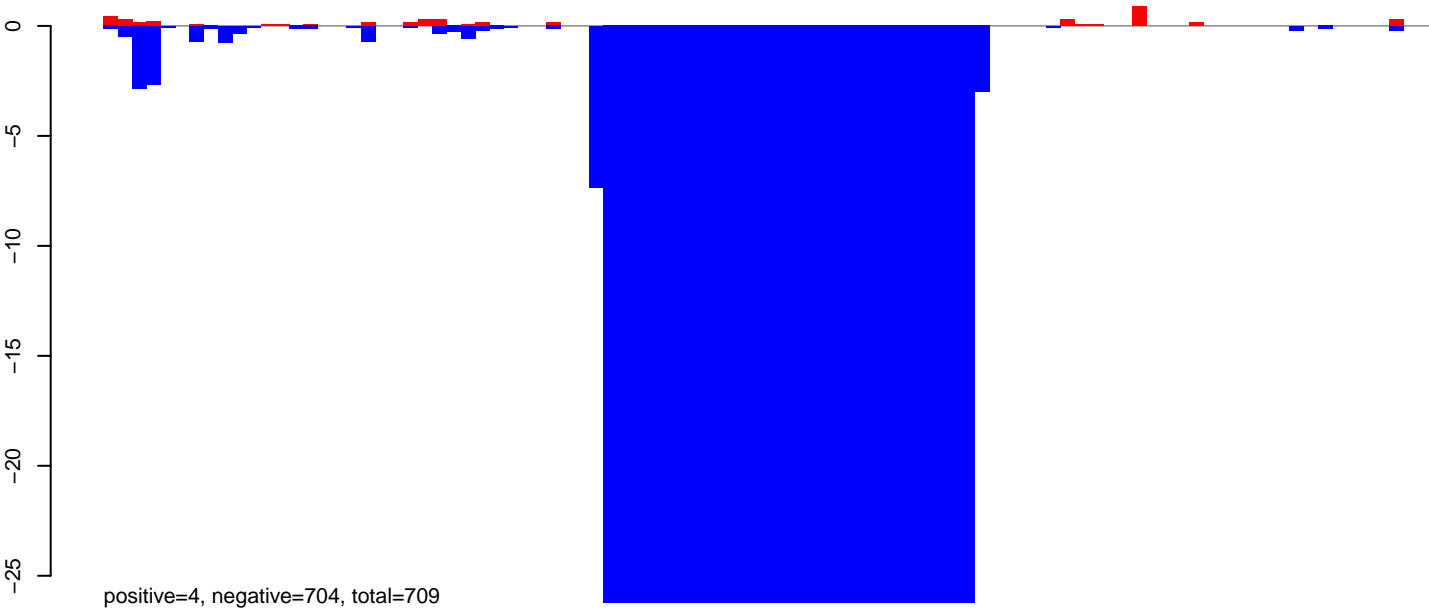
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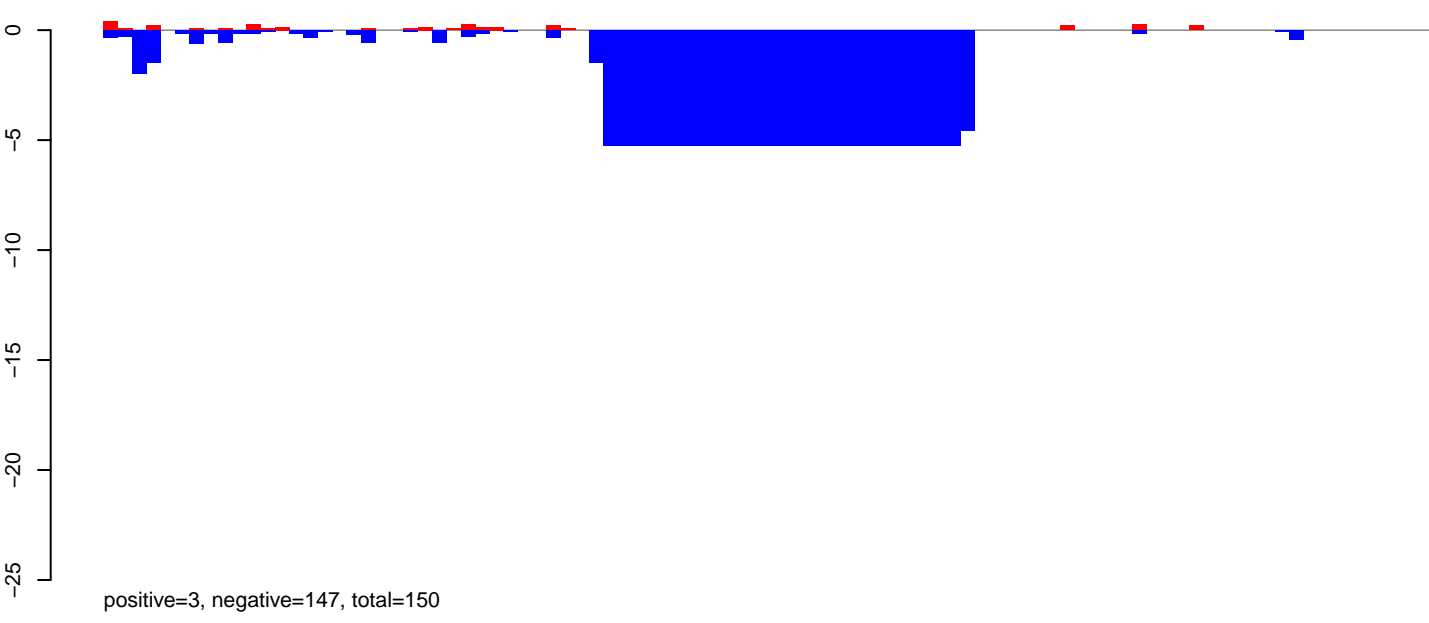
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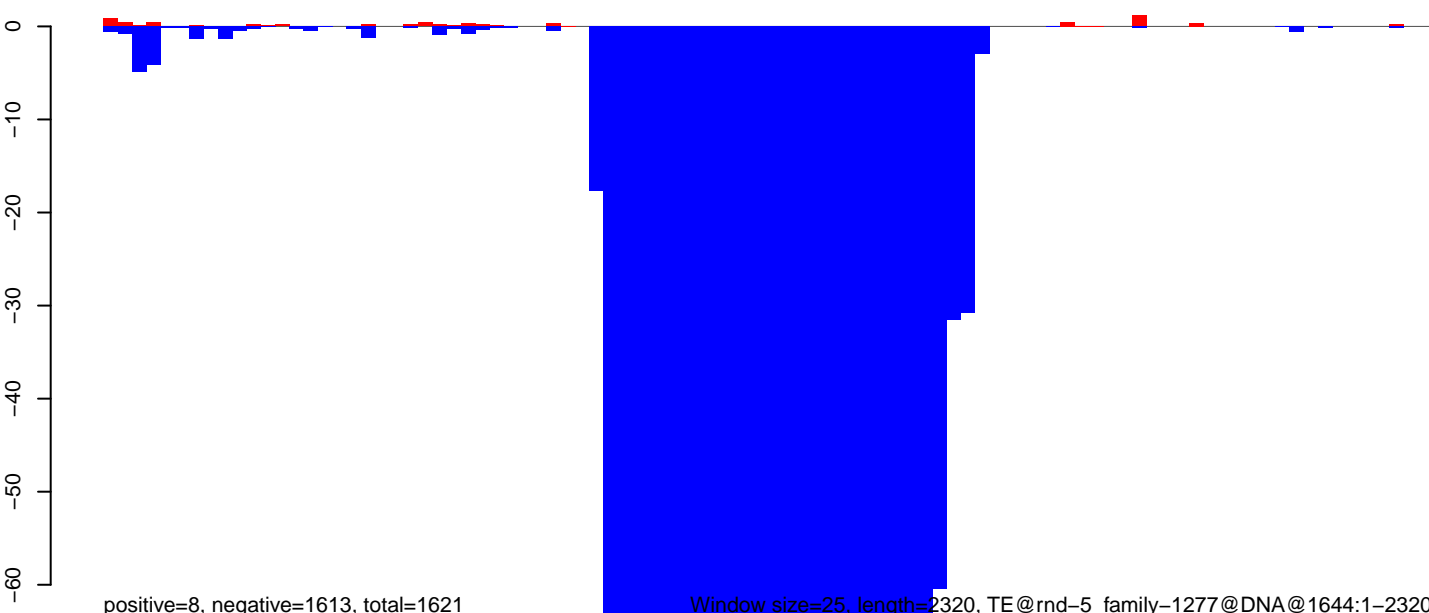
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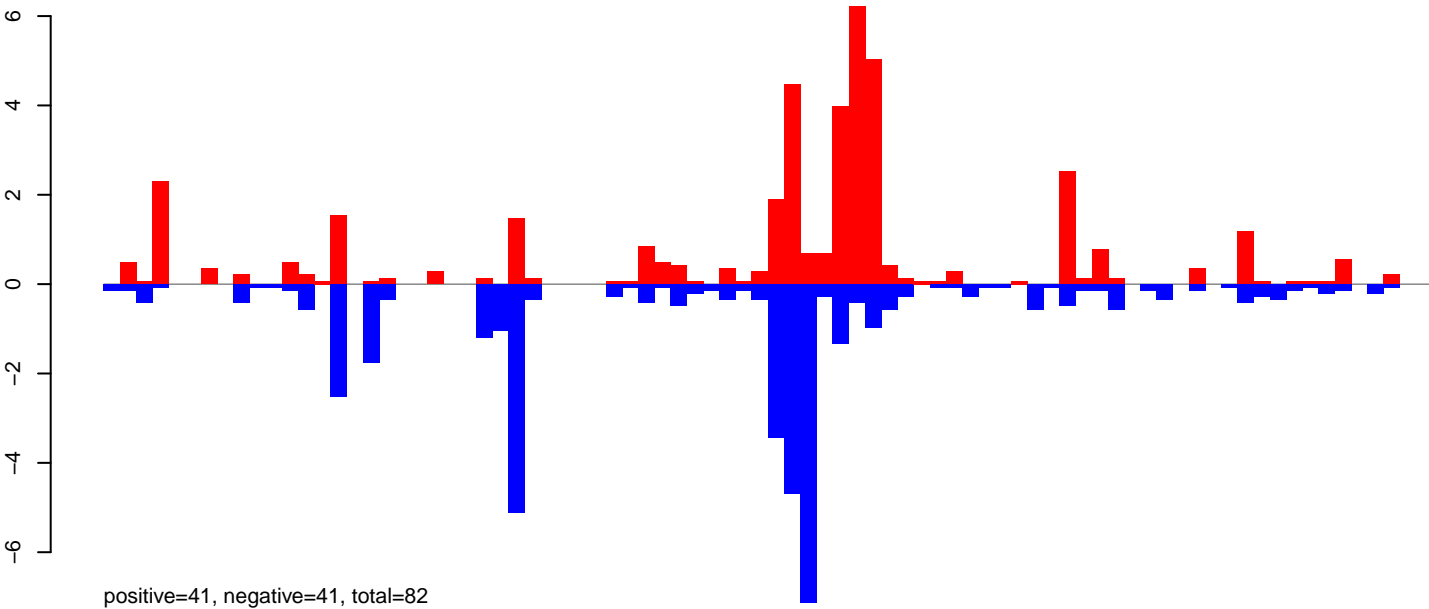
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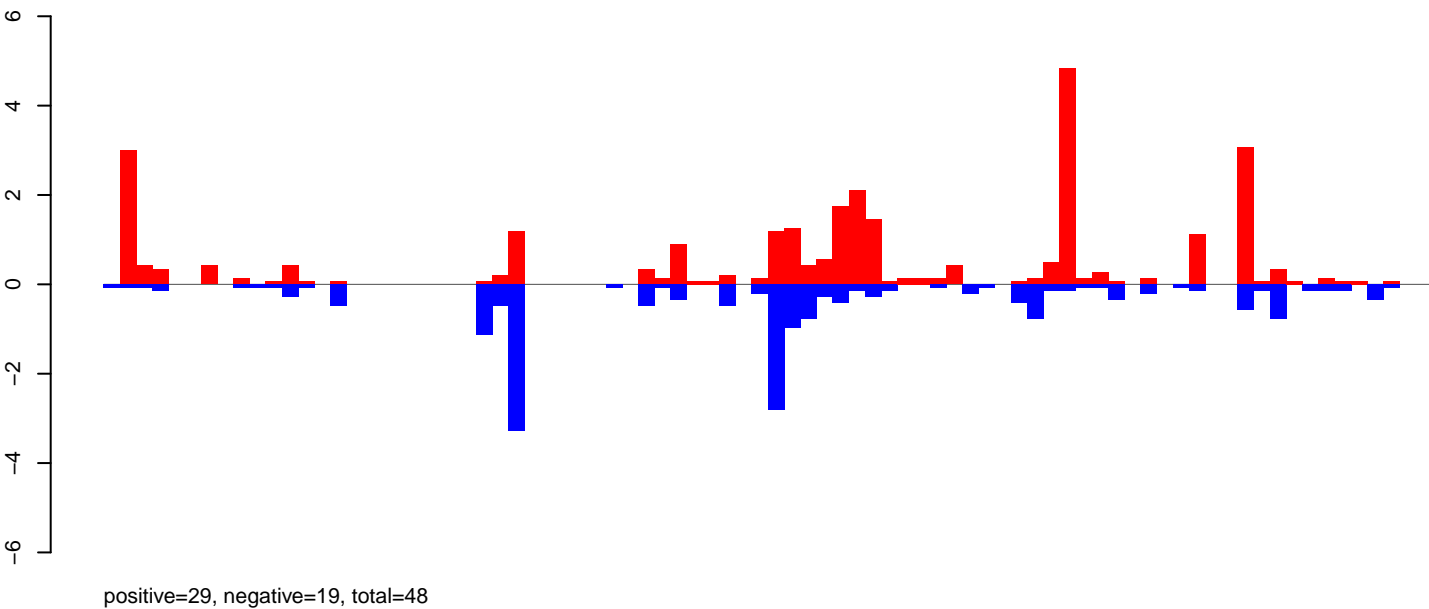
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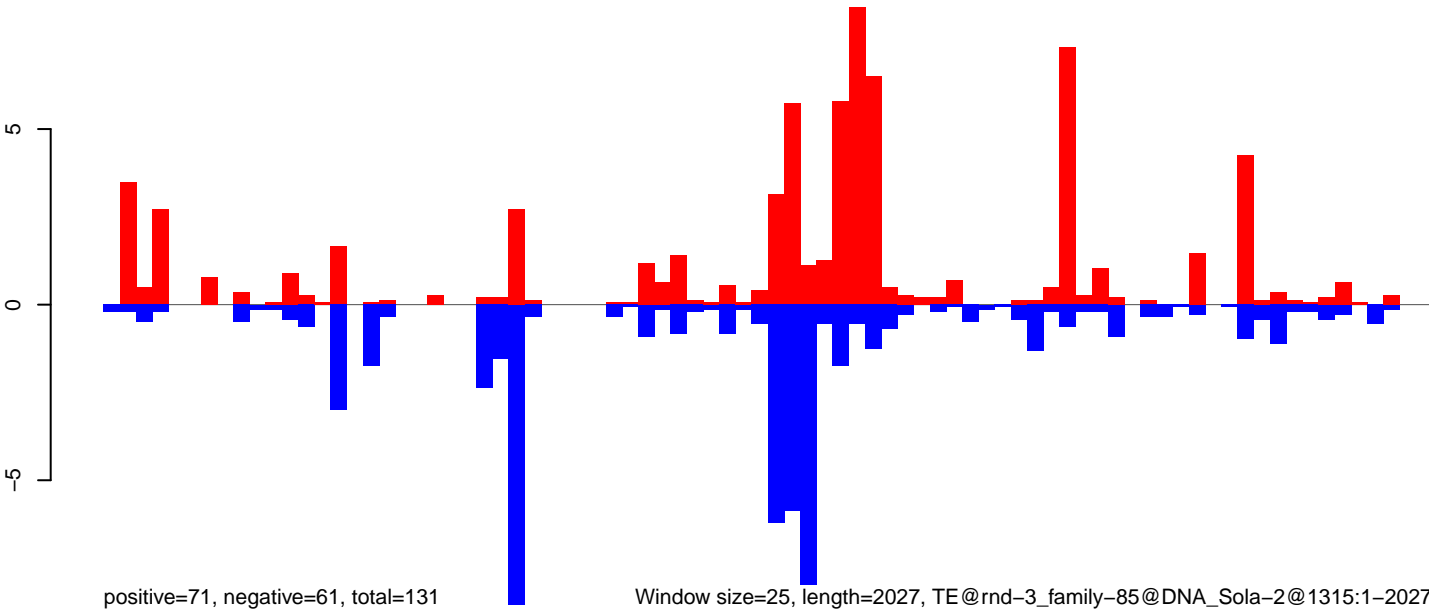
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CuQuin_Mcarc_NL.24_35.rep



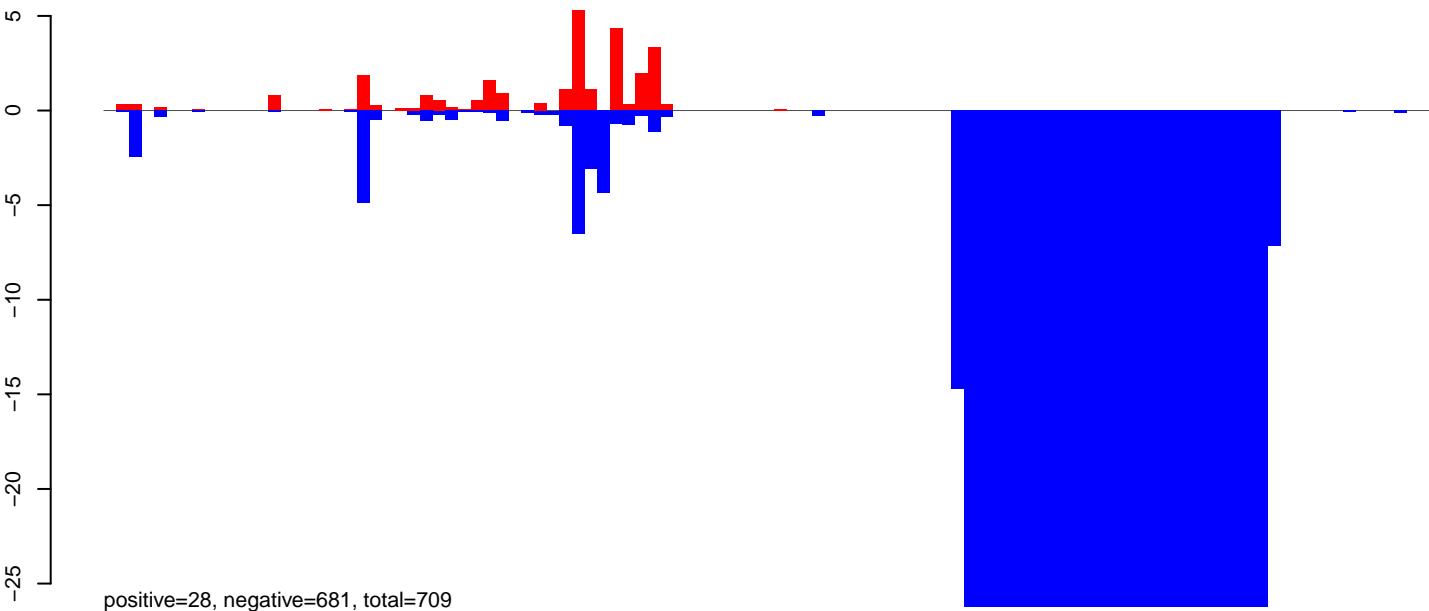
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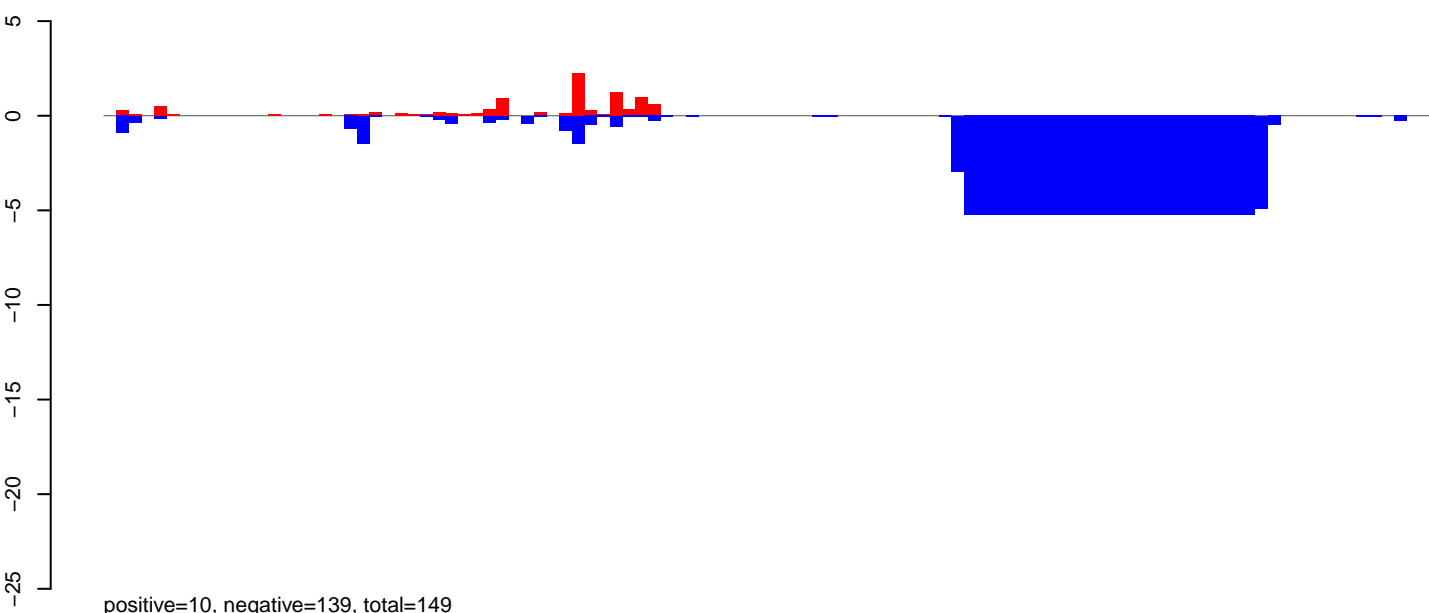
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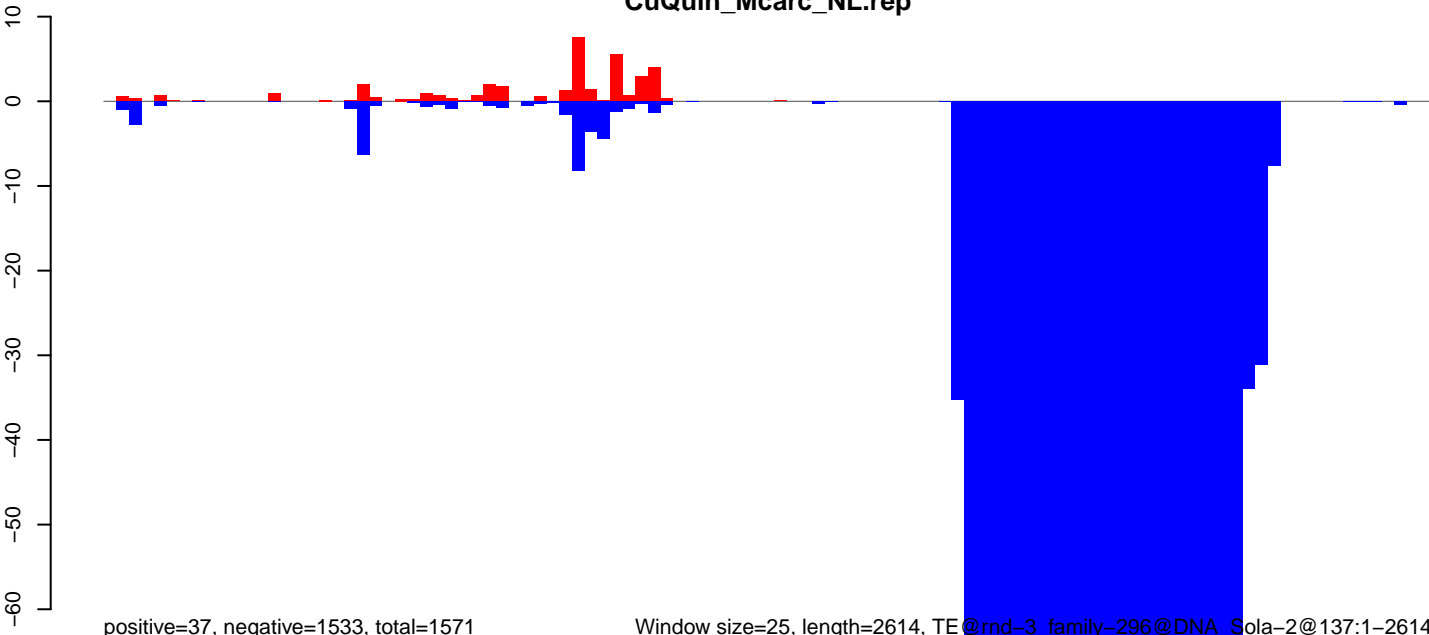
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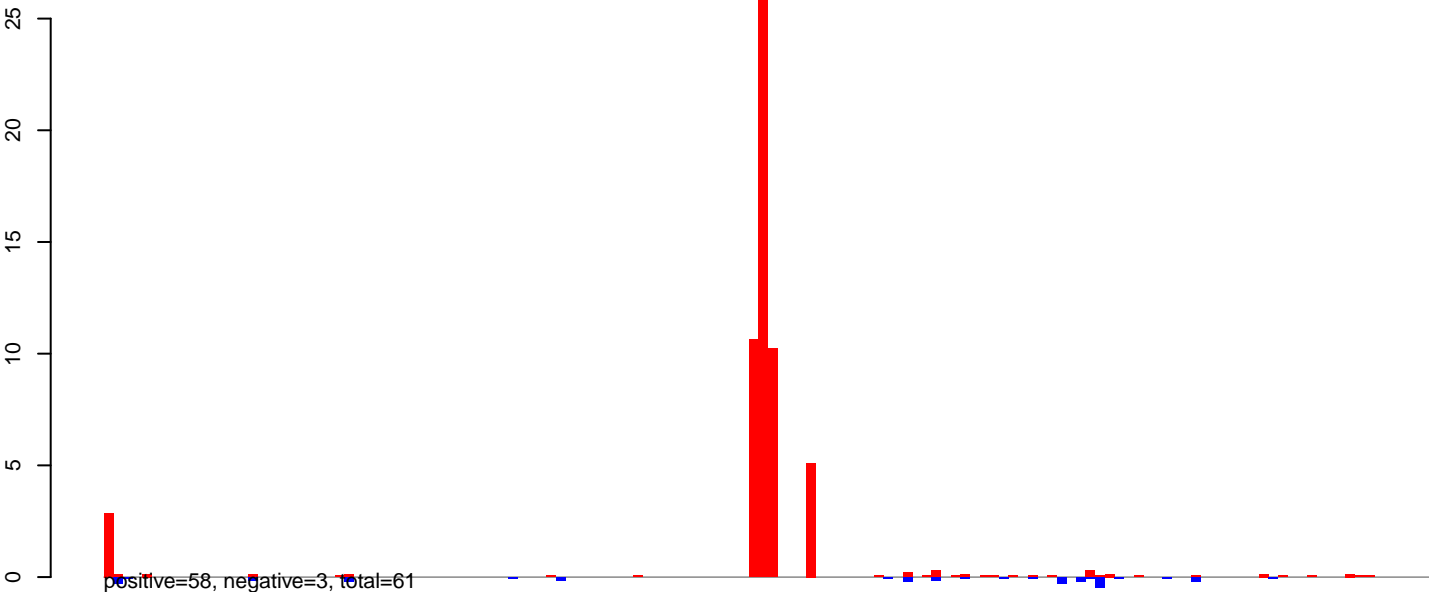


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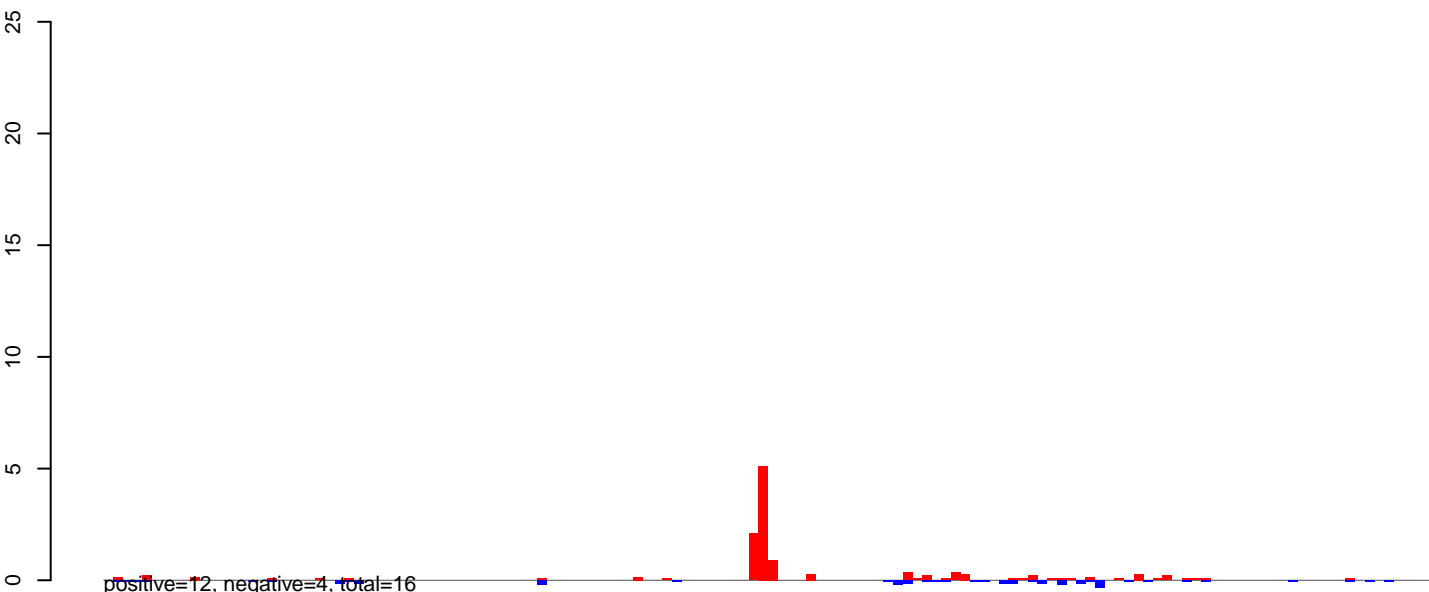


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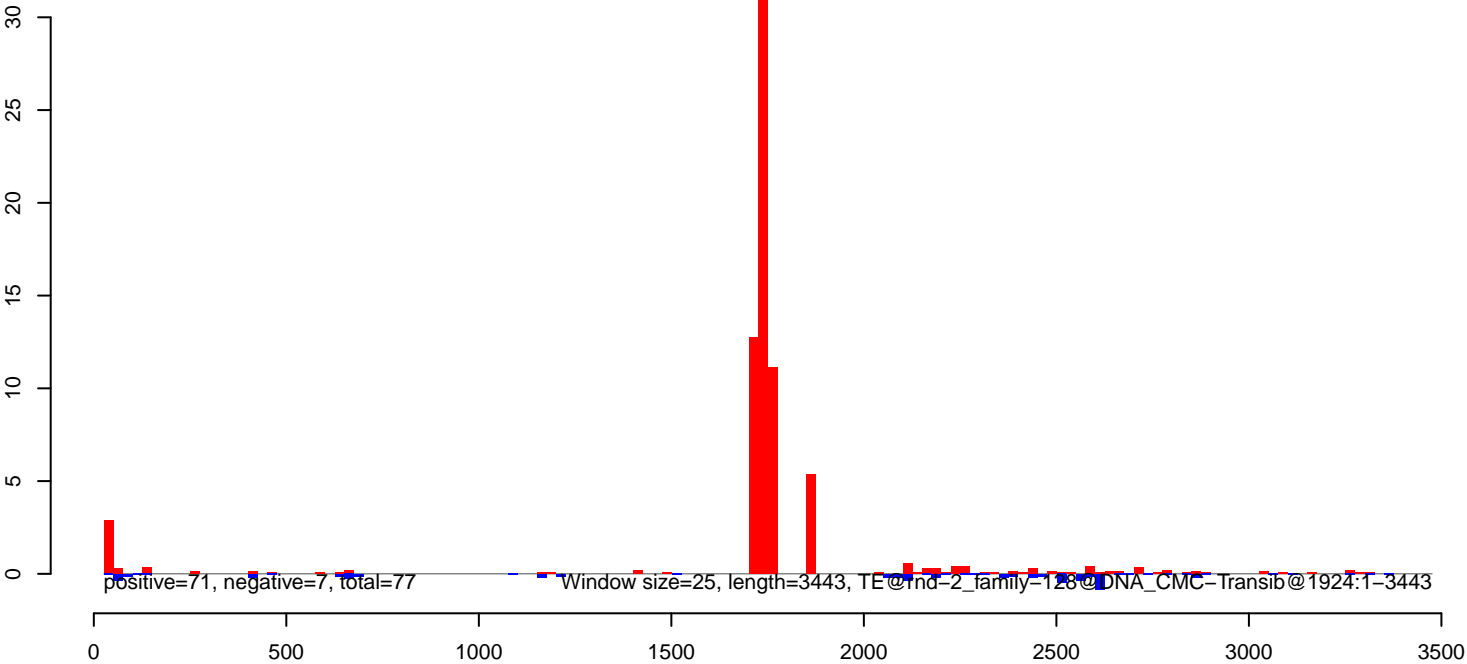
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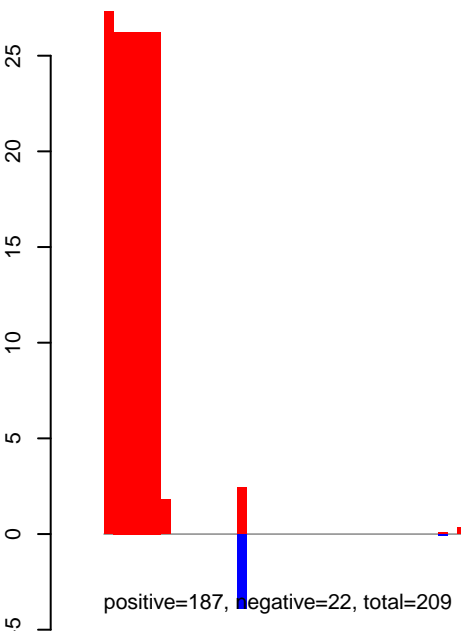
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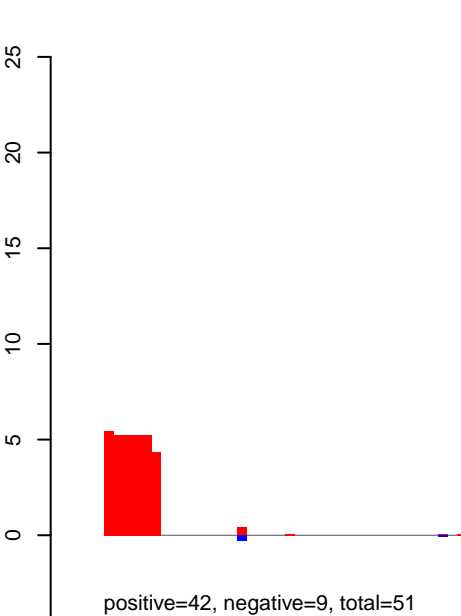
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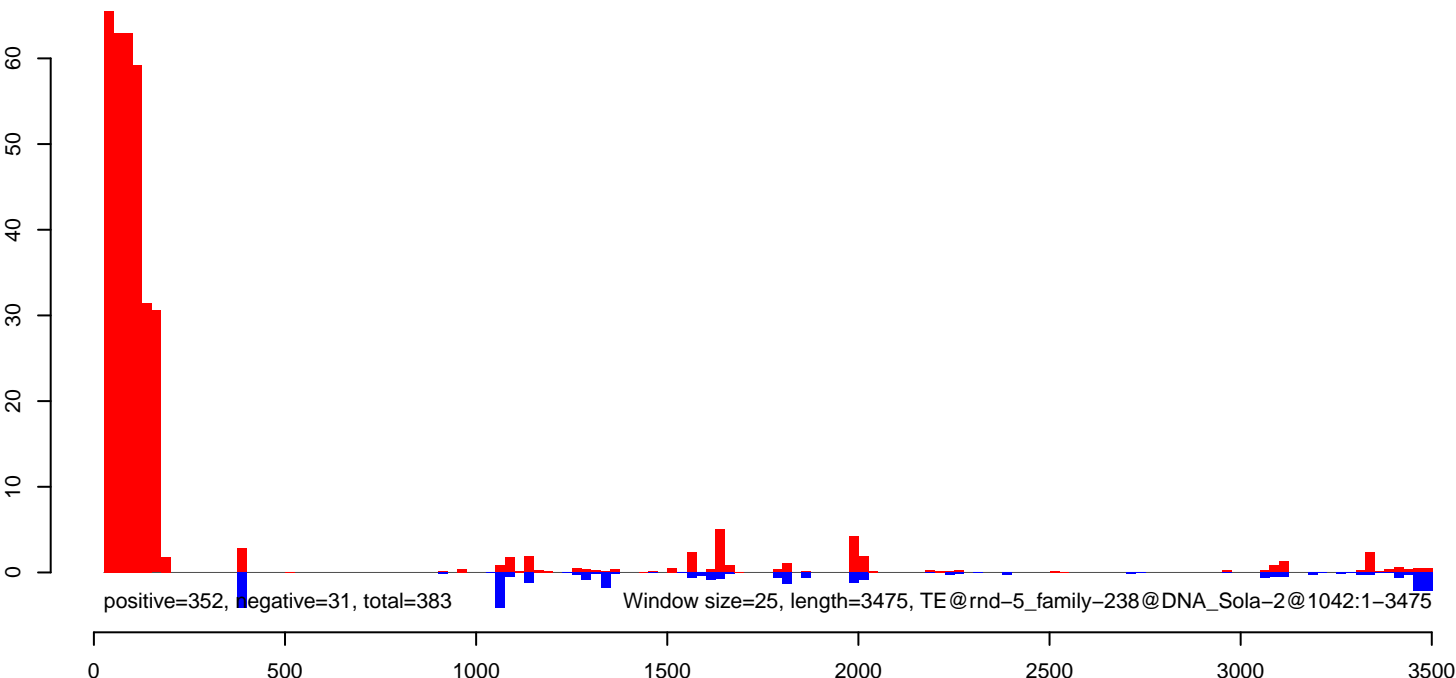
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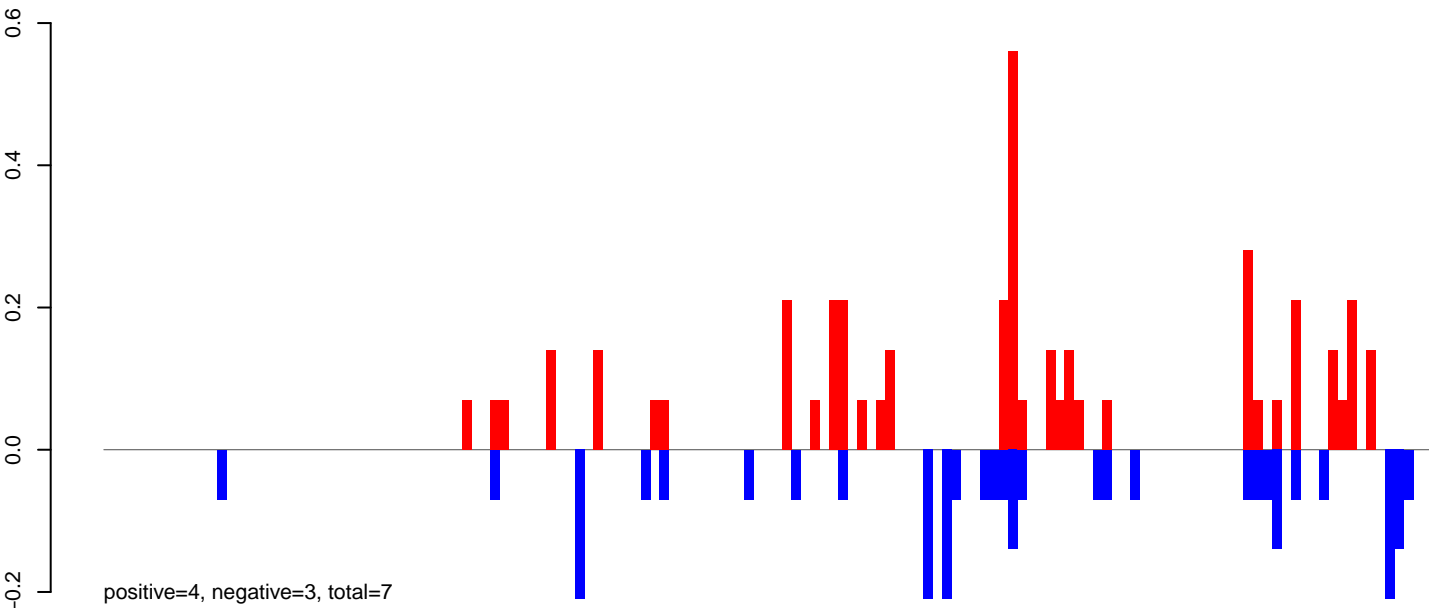
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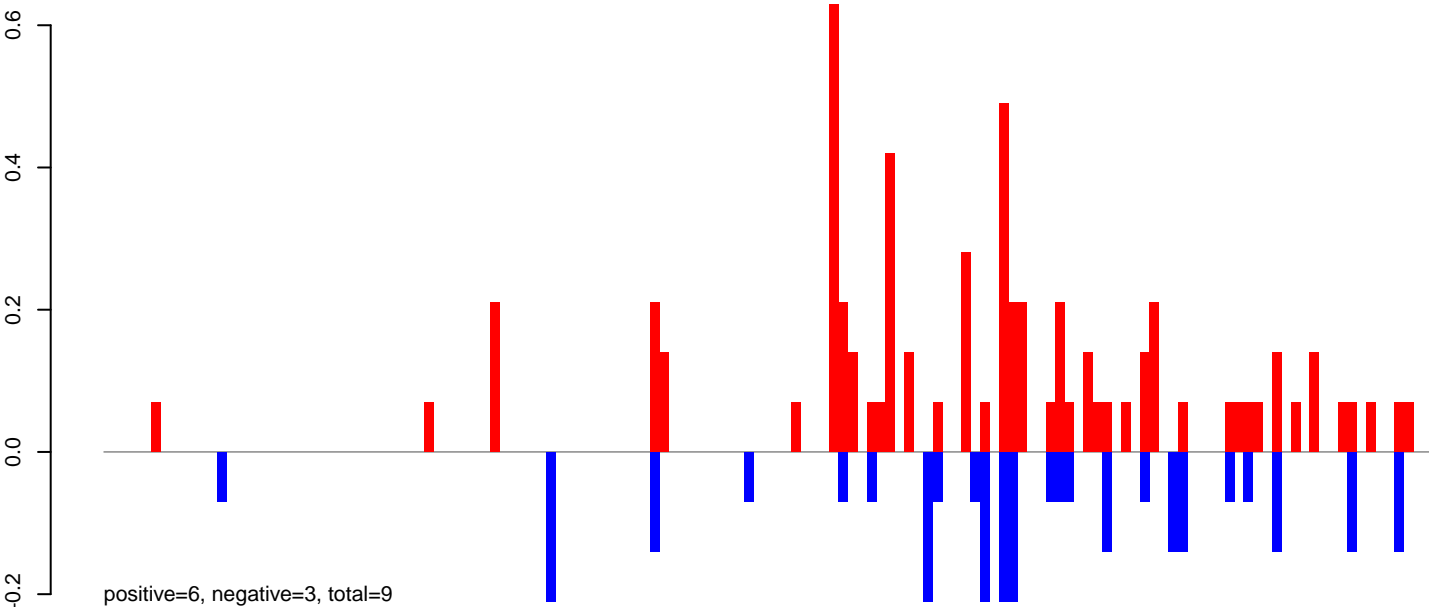
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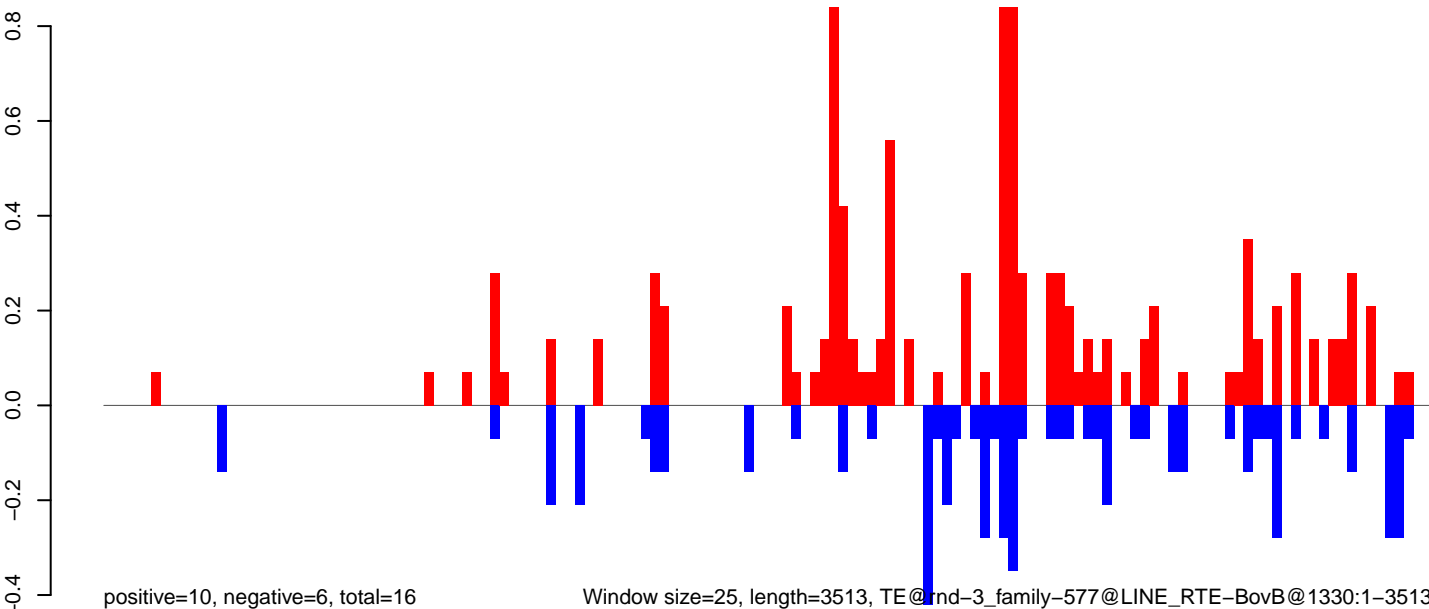
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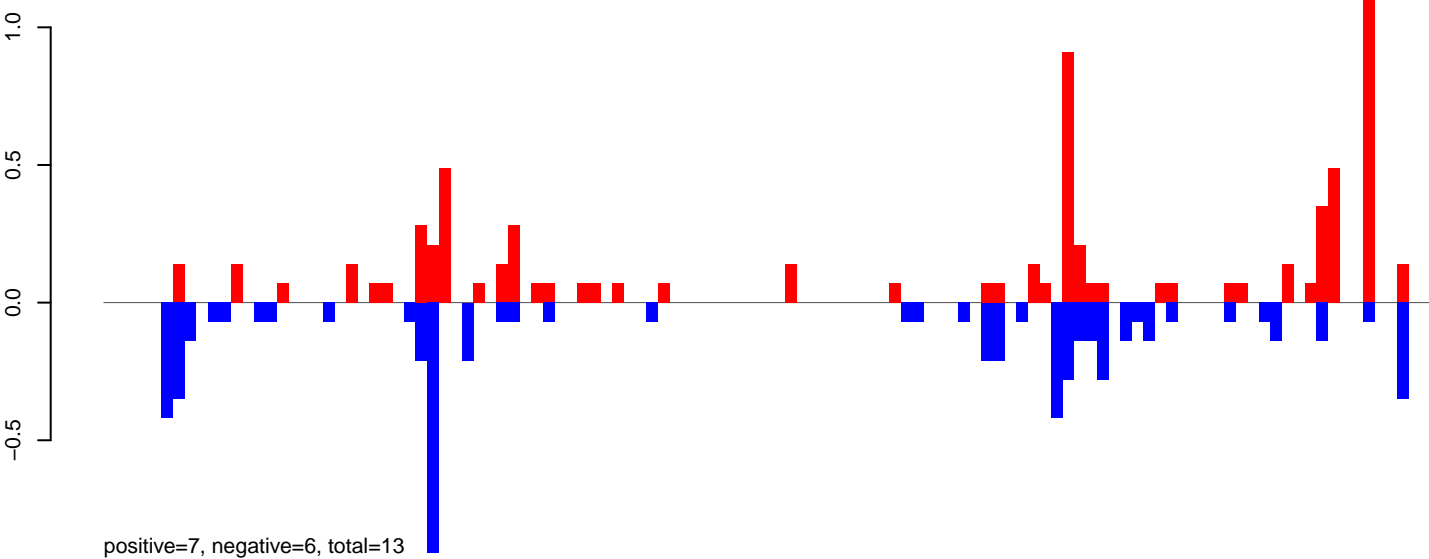
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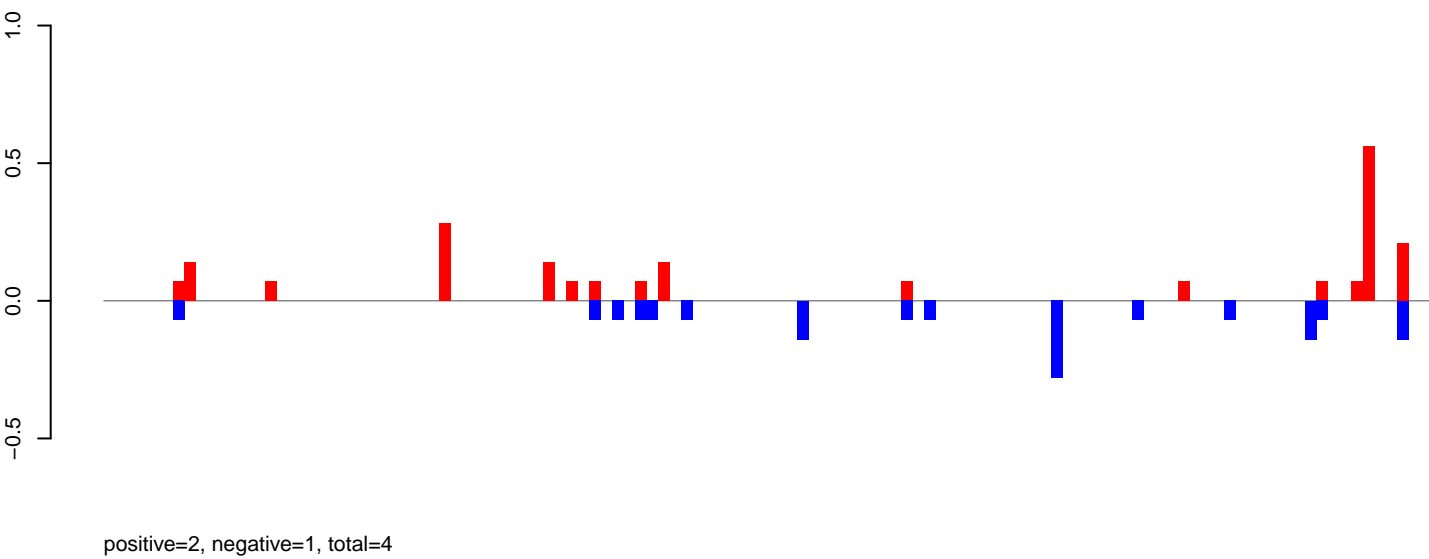
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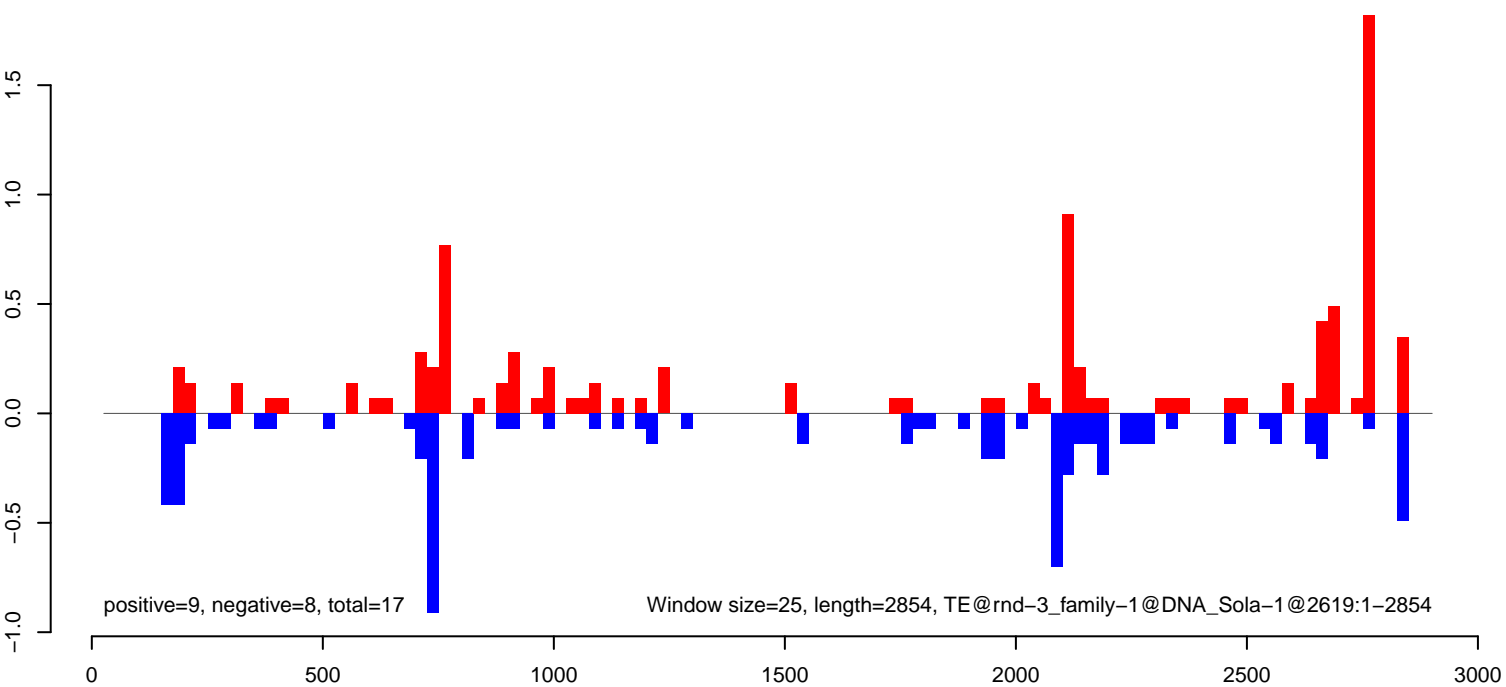
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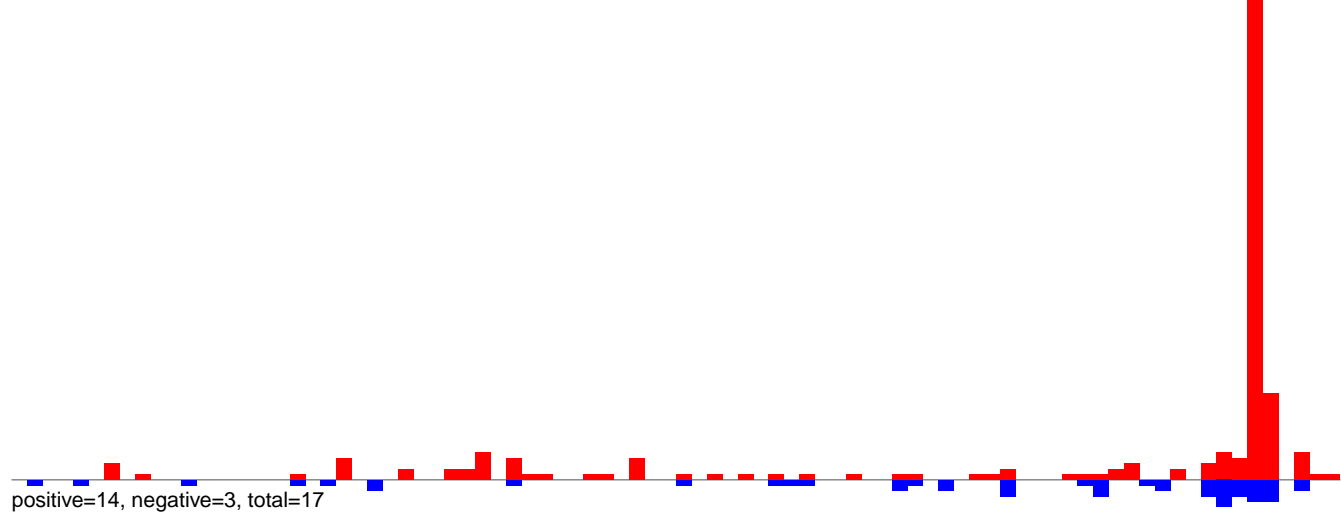
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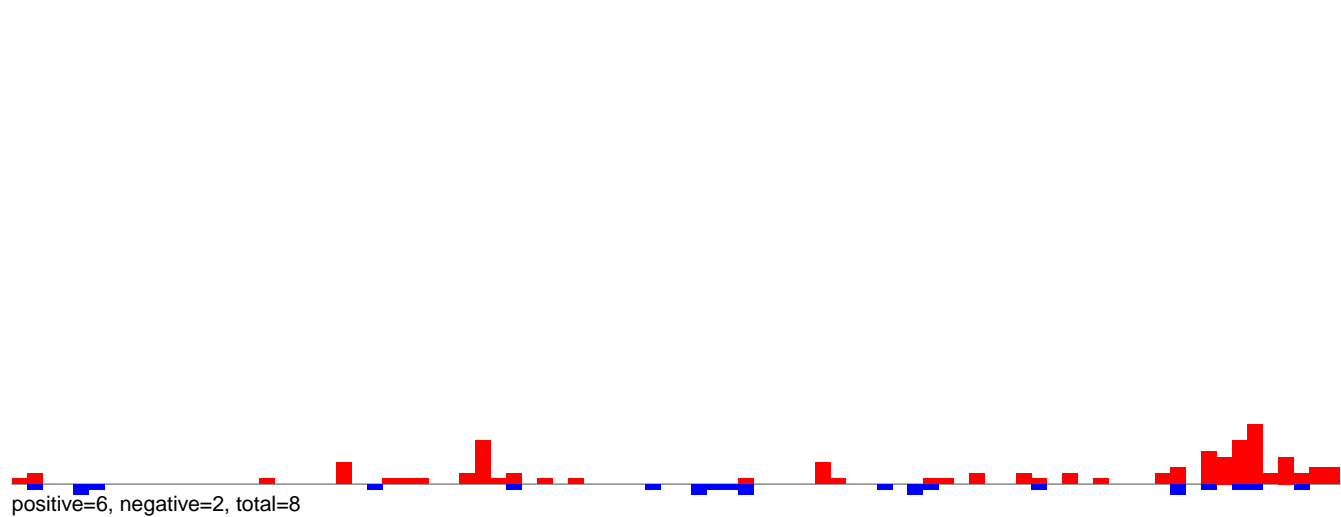
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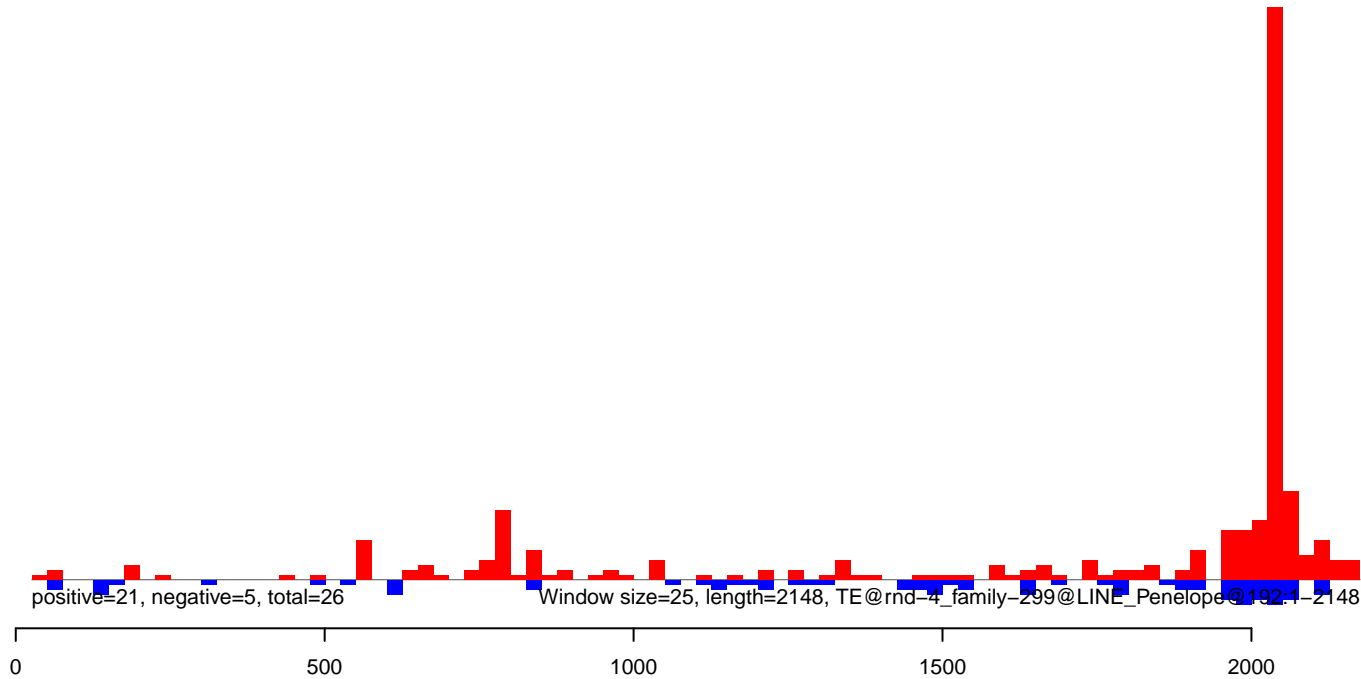
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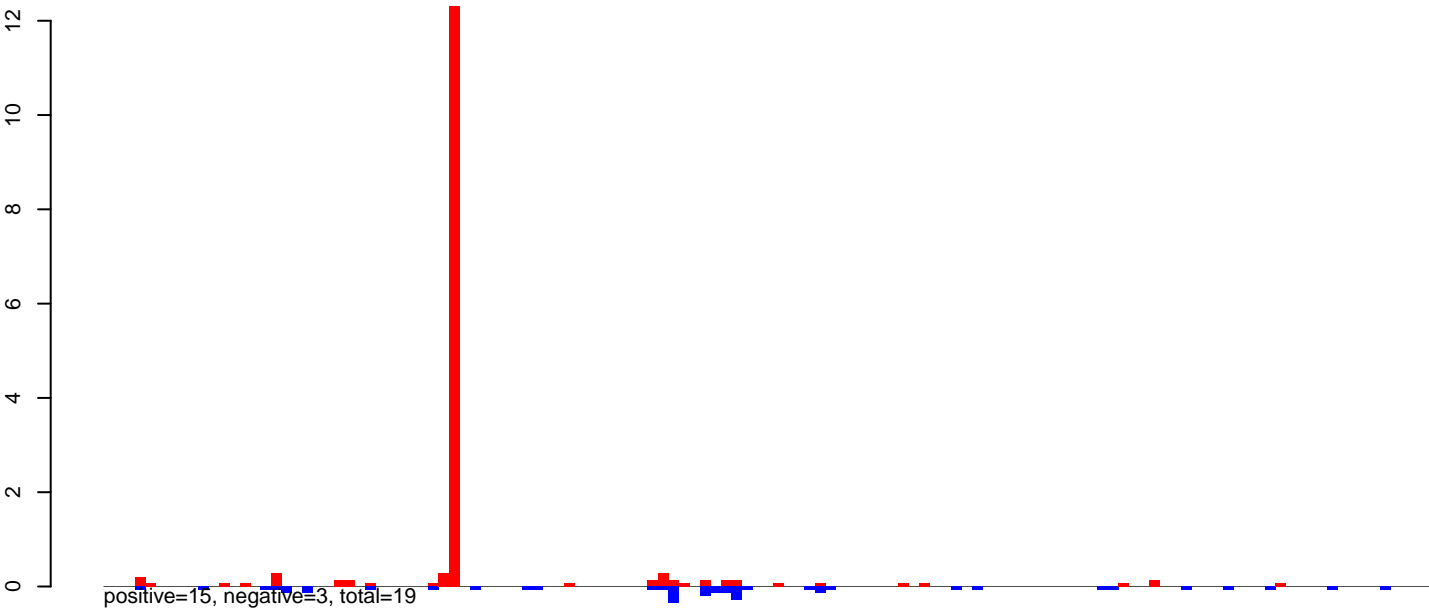
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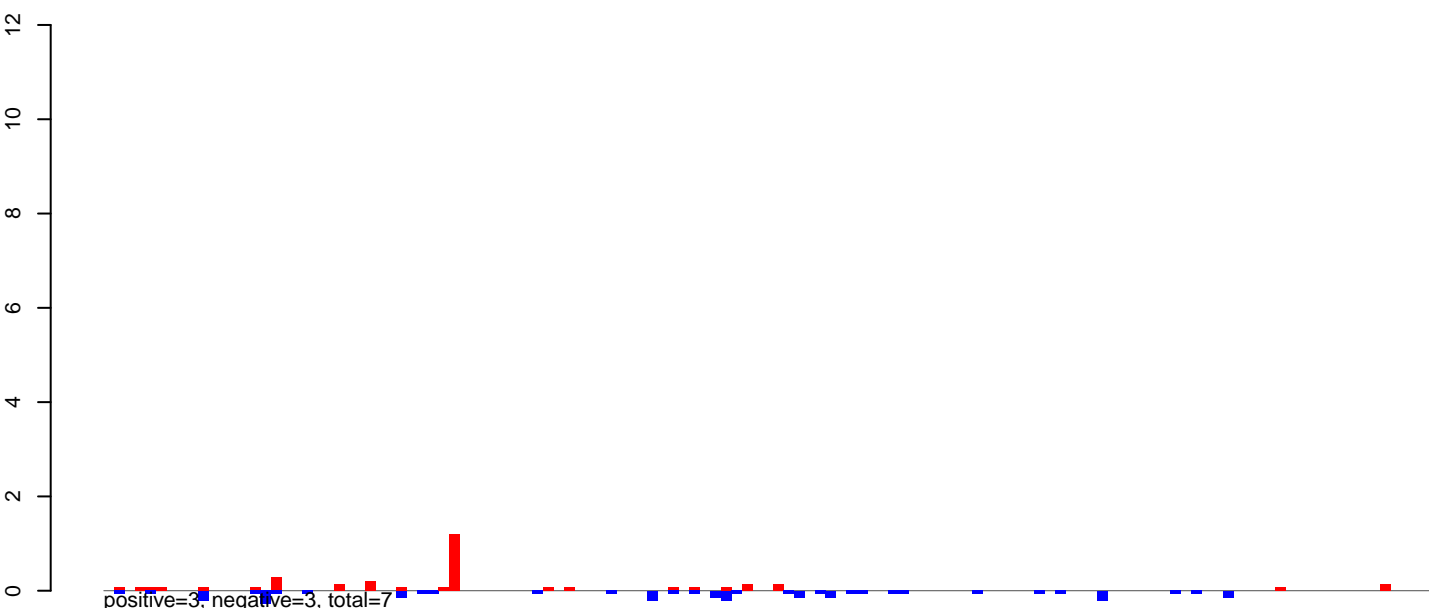
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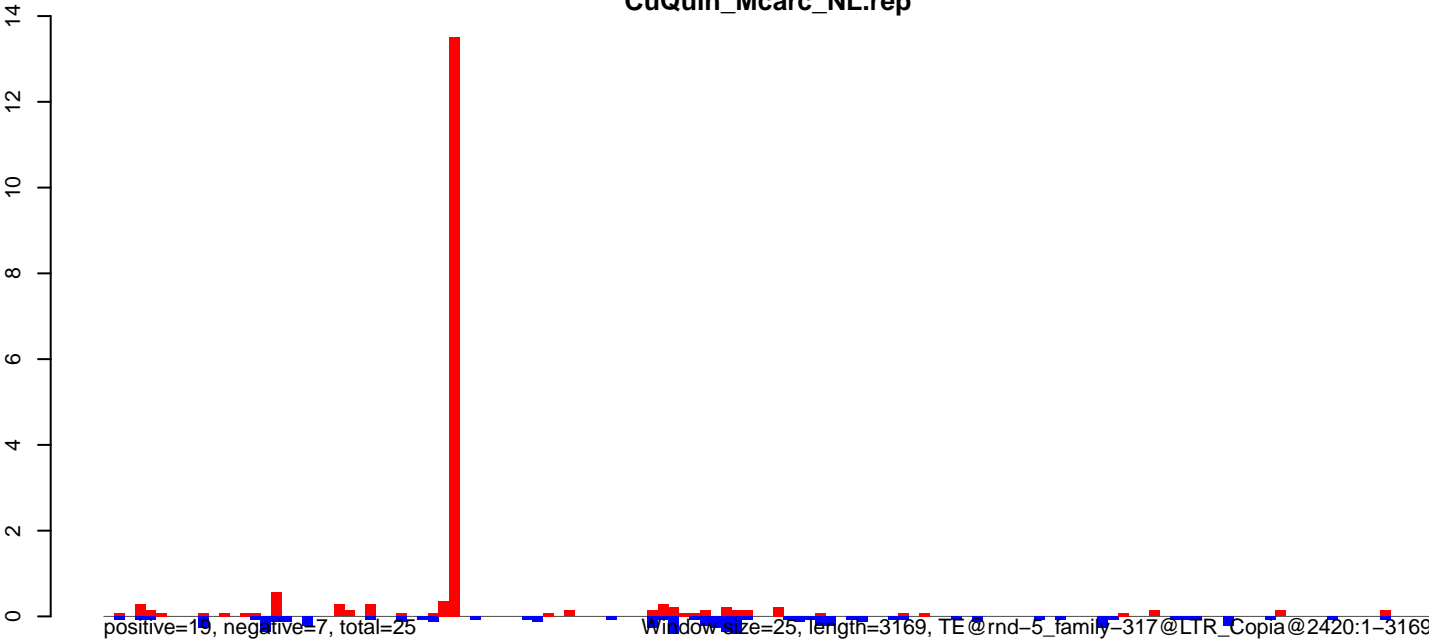
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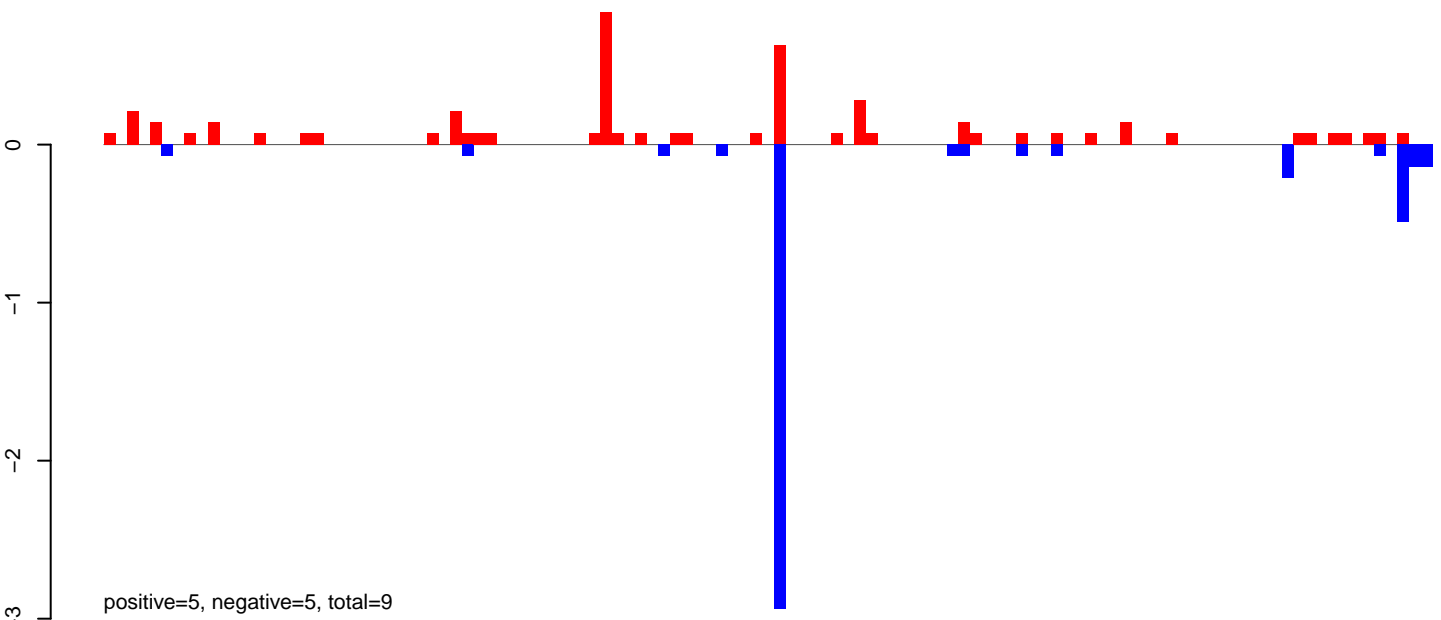


CuQuin_Mcarc_NL.rep

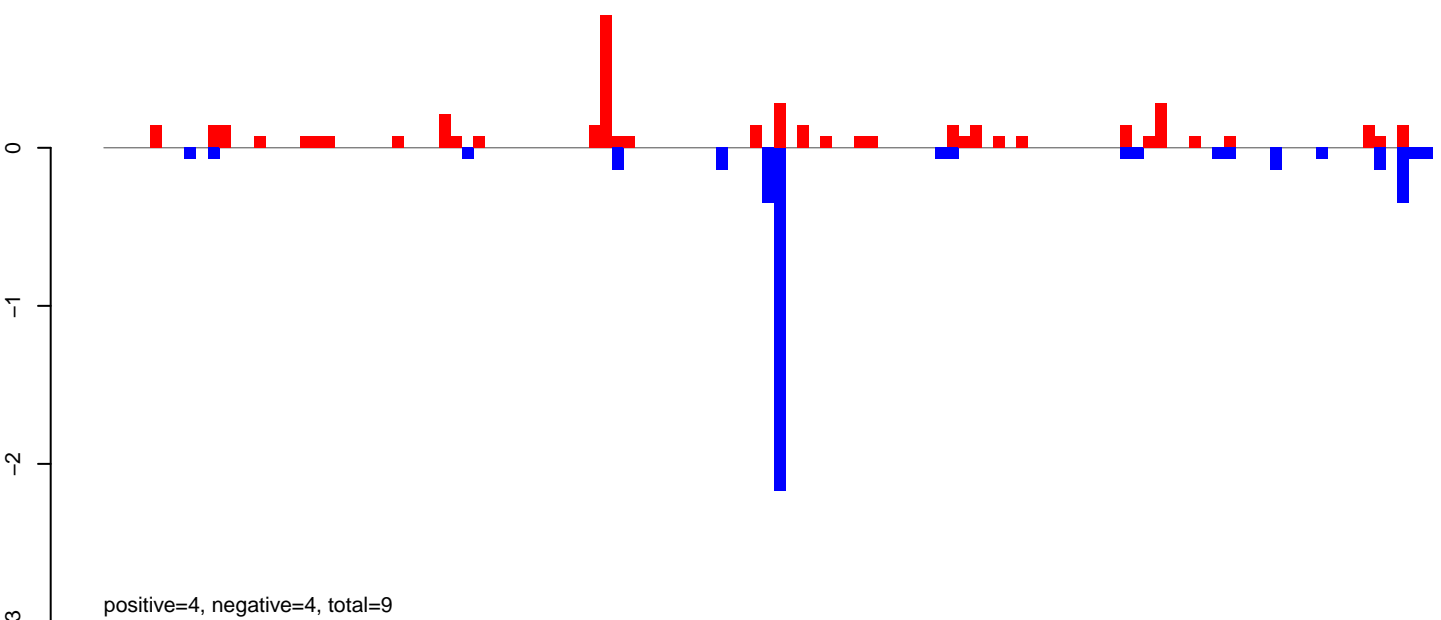


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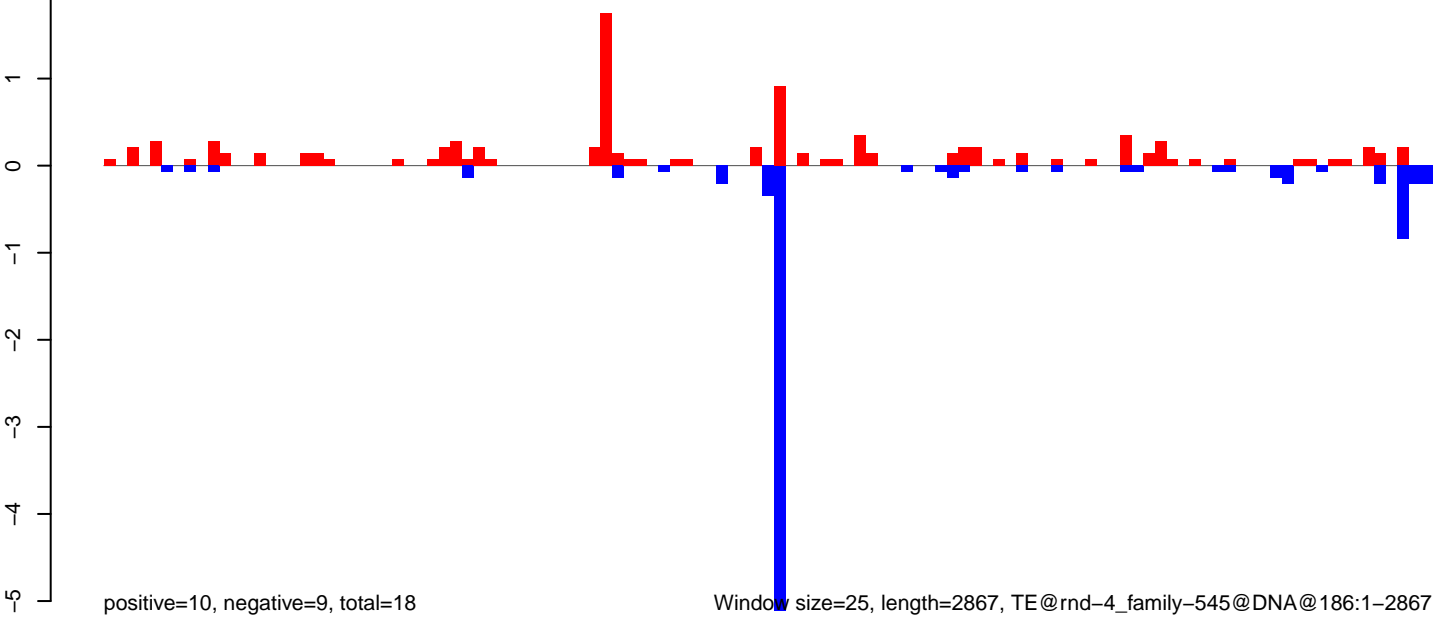
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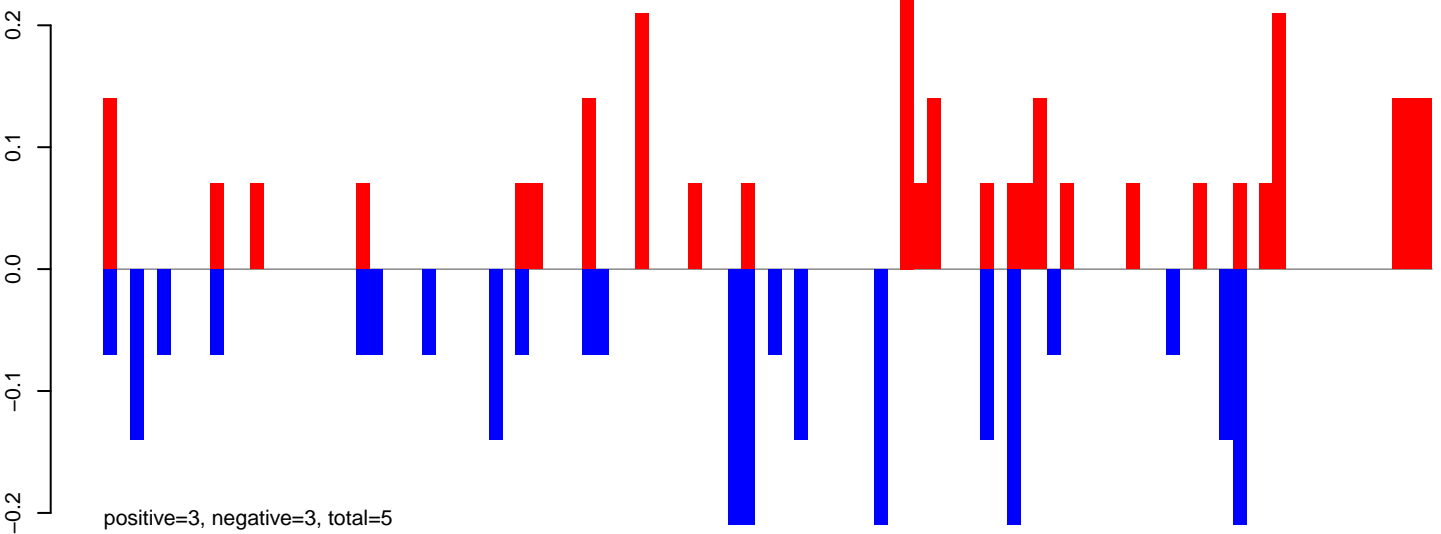
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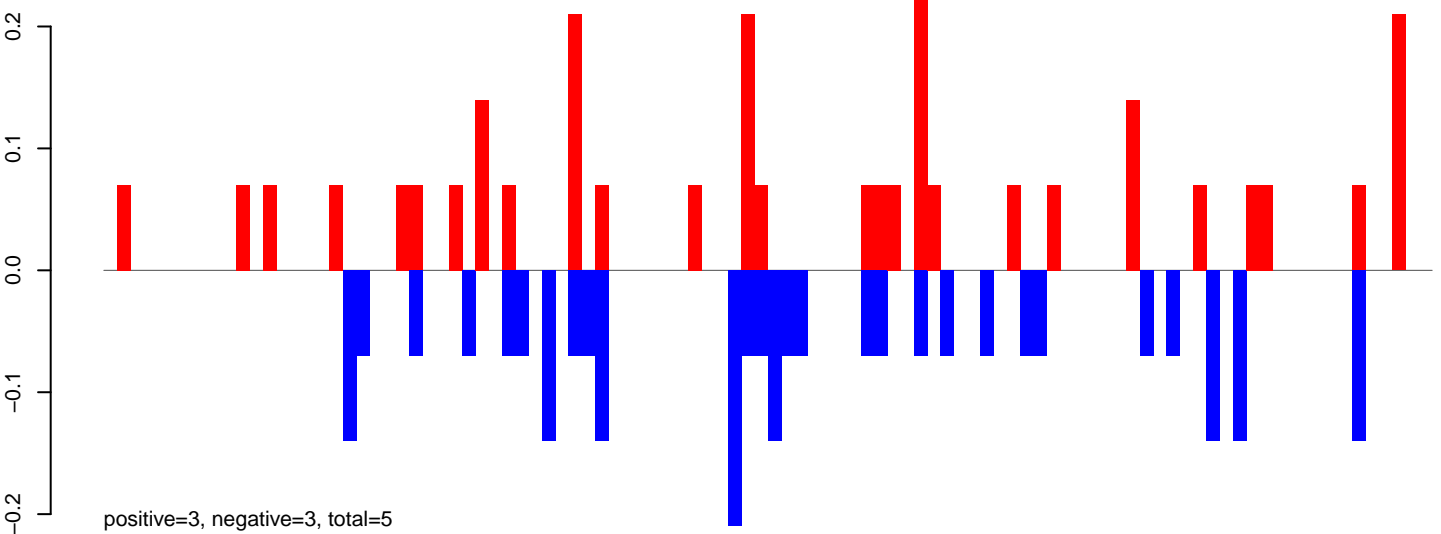
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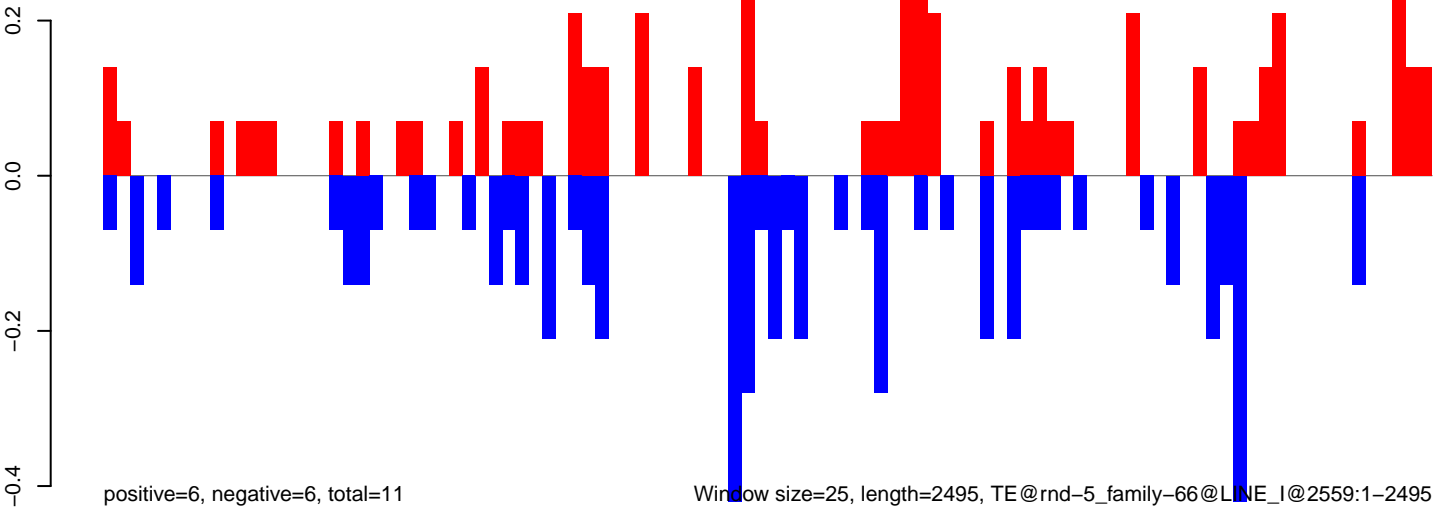
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CuQuin_Mcarc_NL.24_35.rep

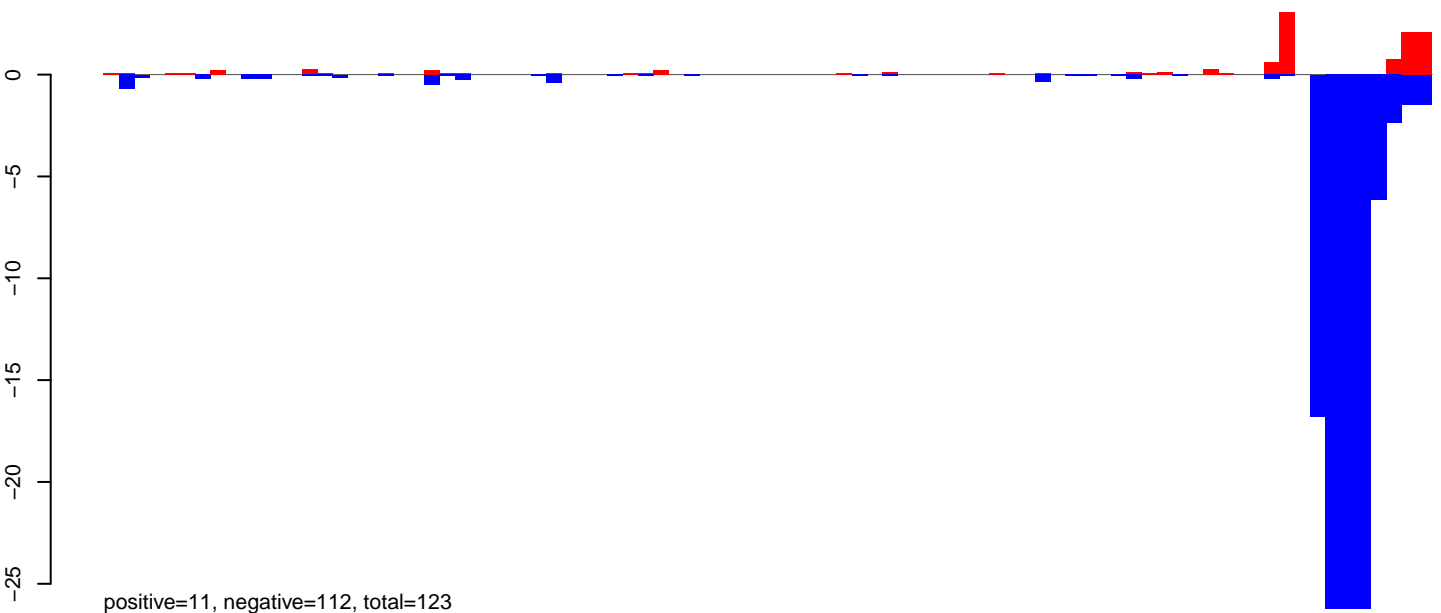


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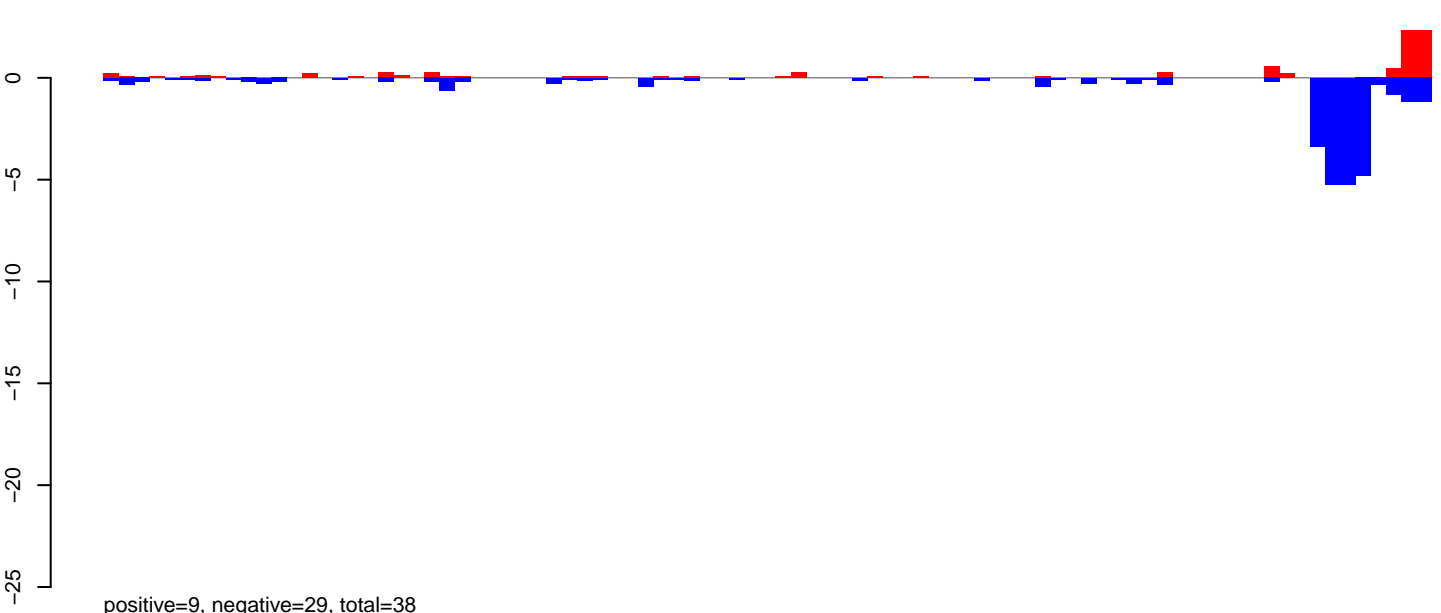


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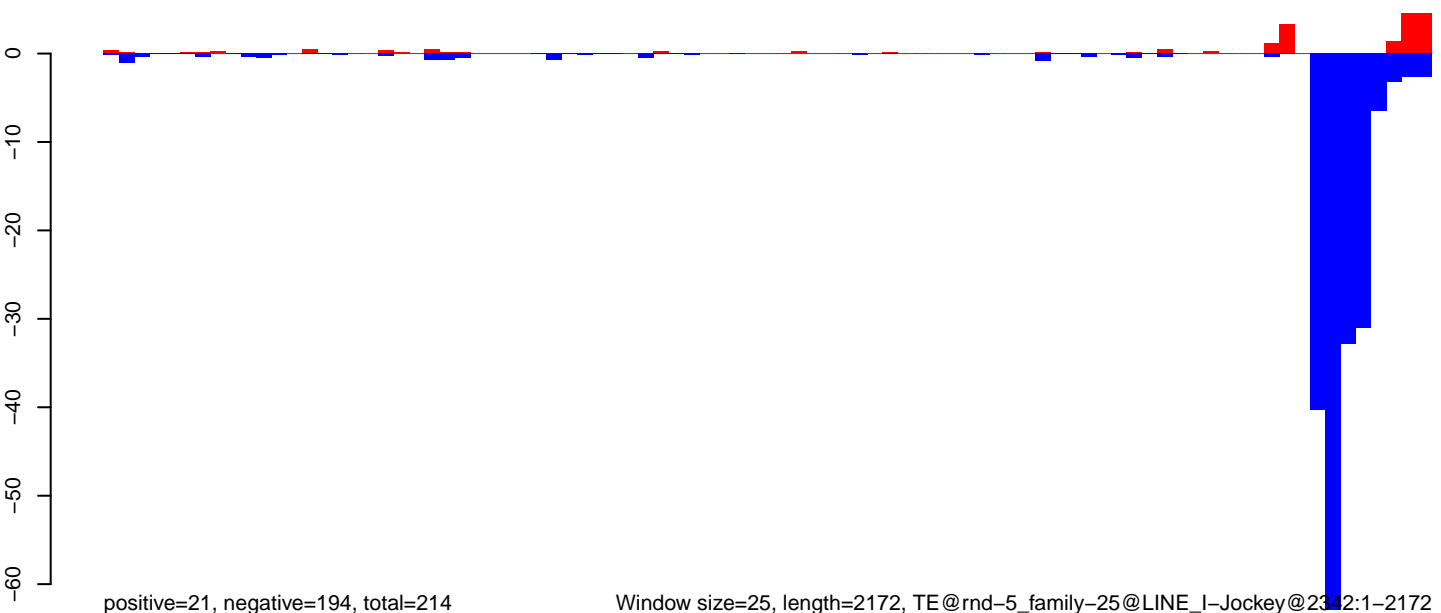
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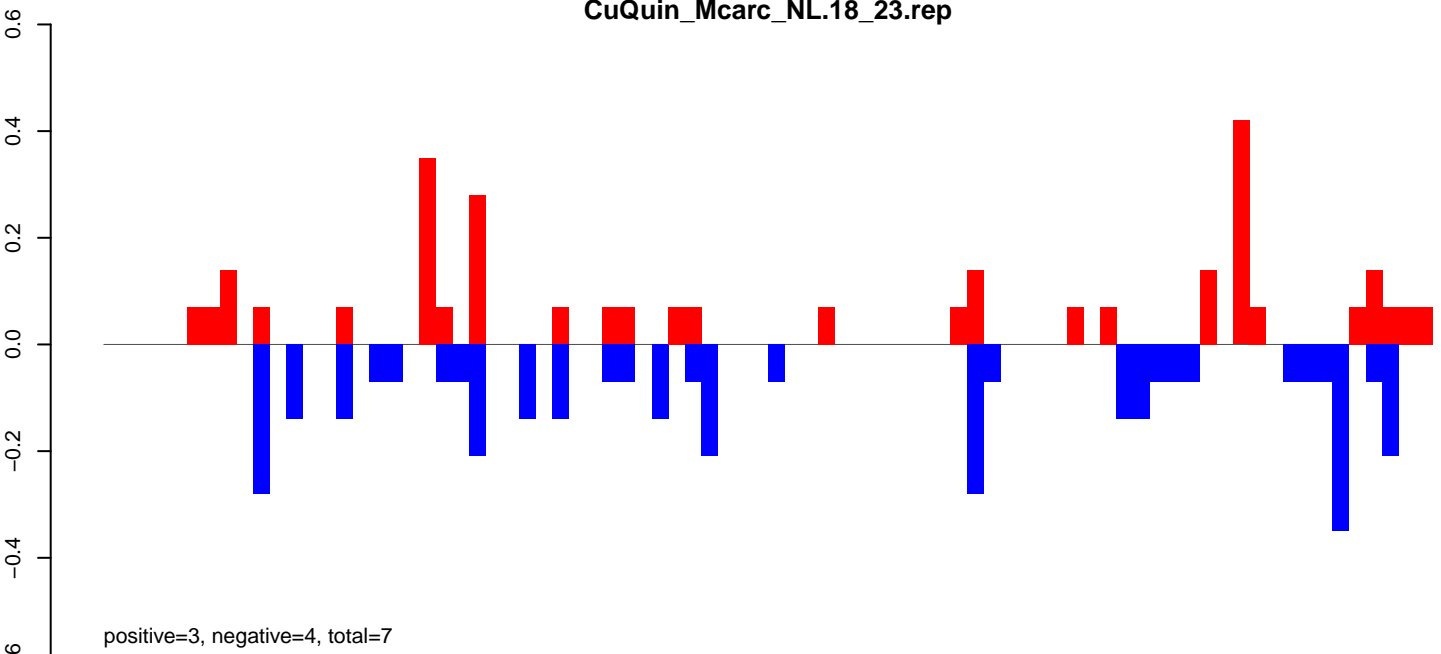
CuQuin_Mcarc_NL.24_35.rep



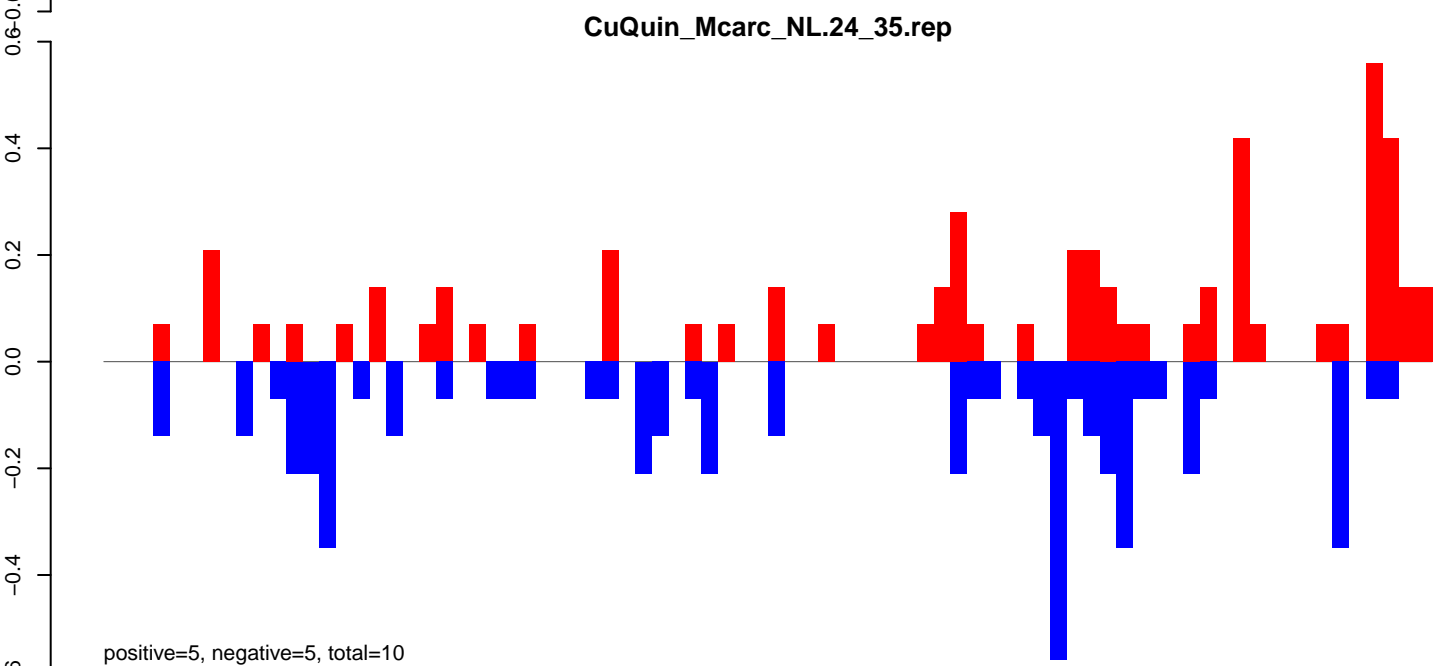
CuQuin_Mcarc_NL.rep



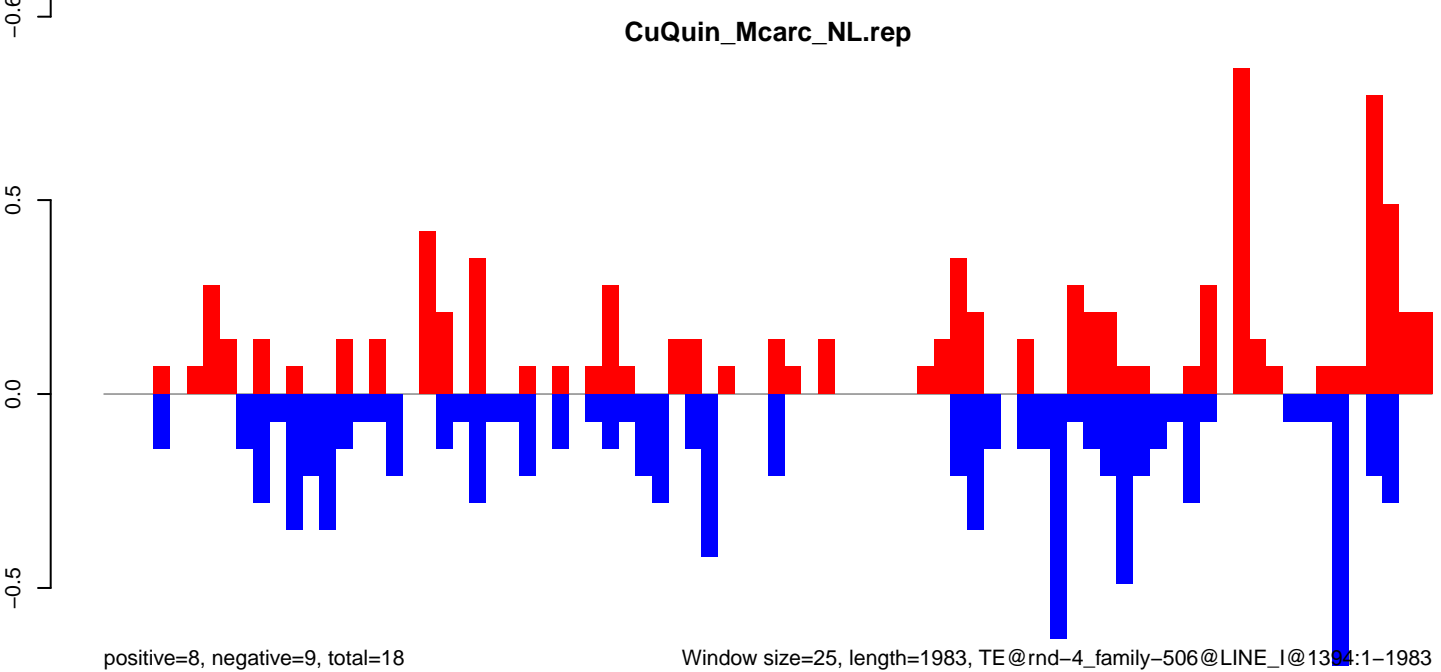
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



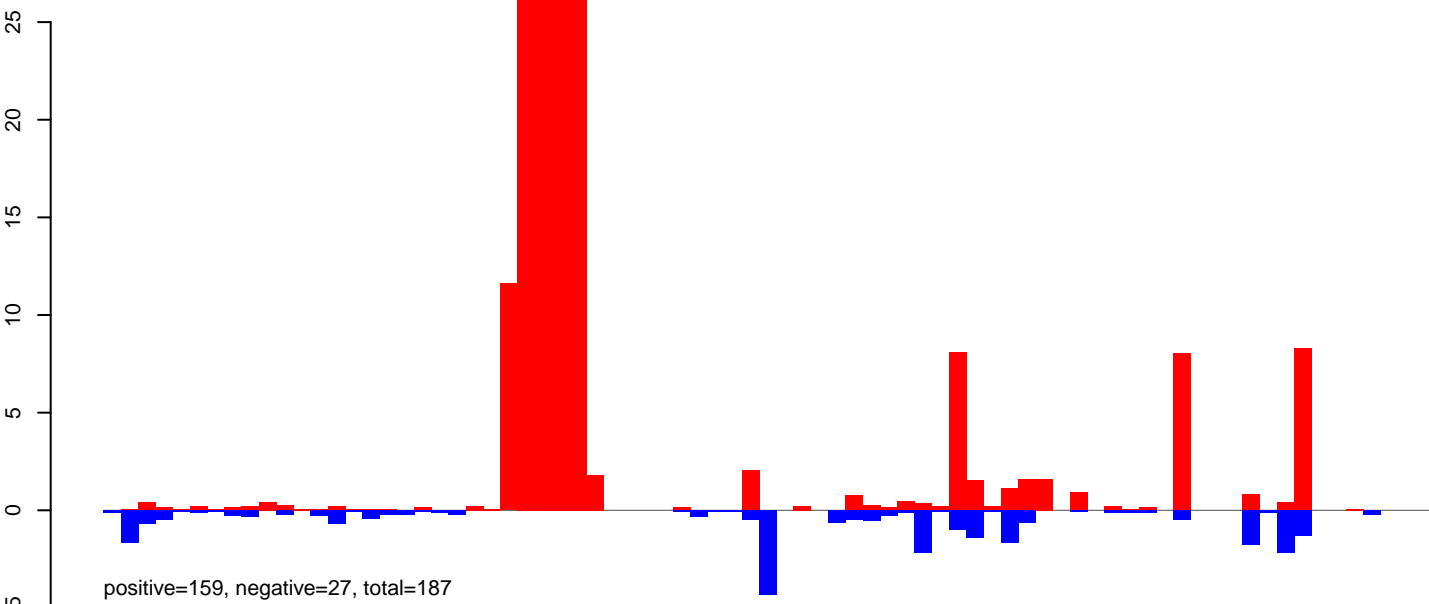
CuQuin_Mcarc_NL.rep



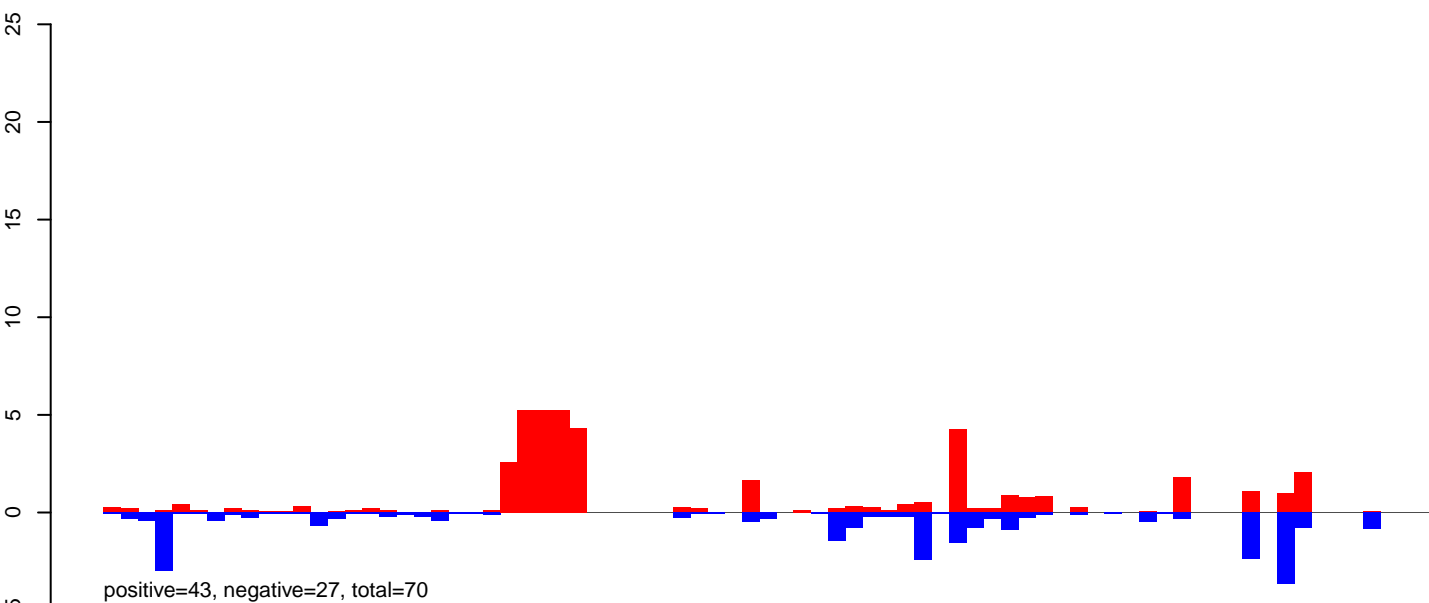
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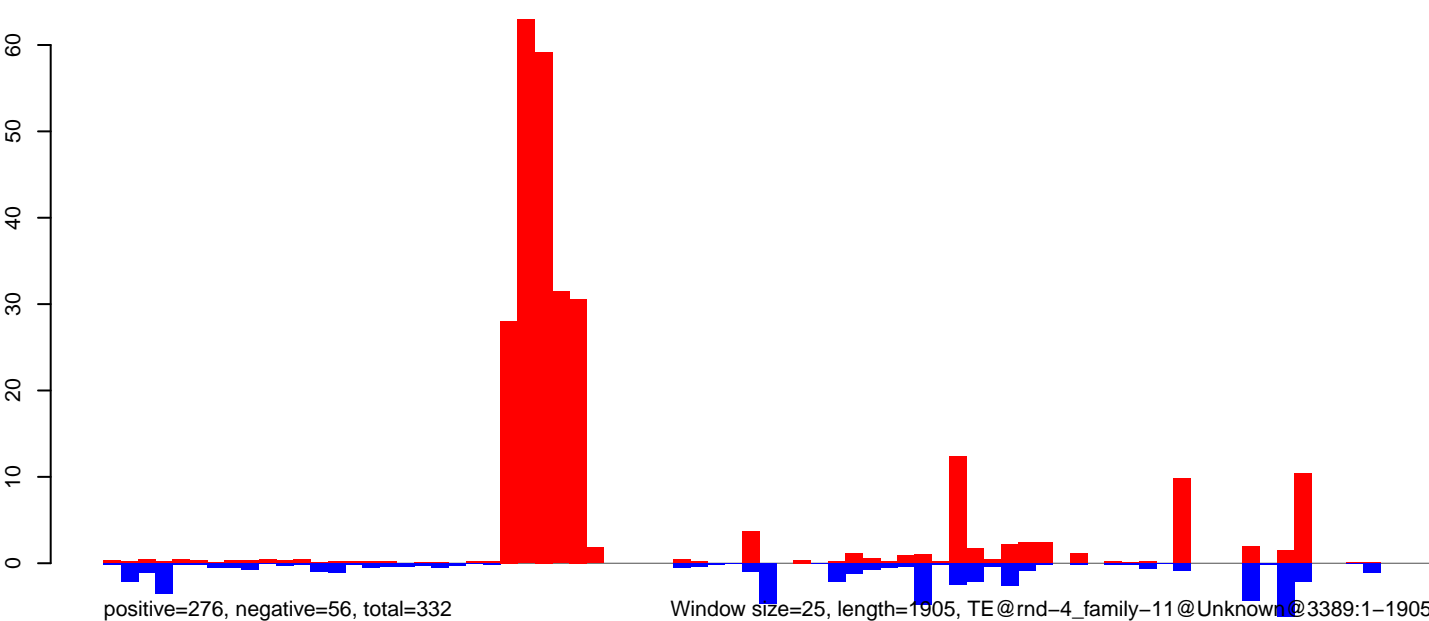
CuQuin_Mcarc_NL.18_23.rep



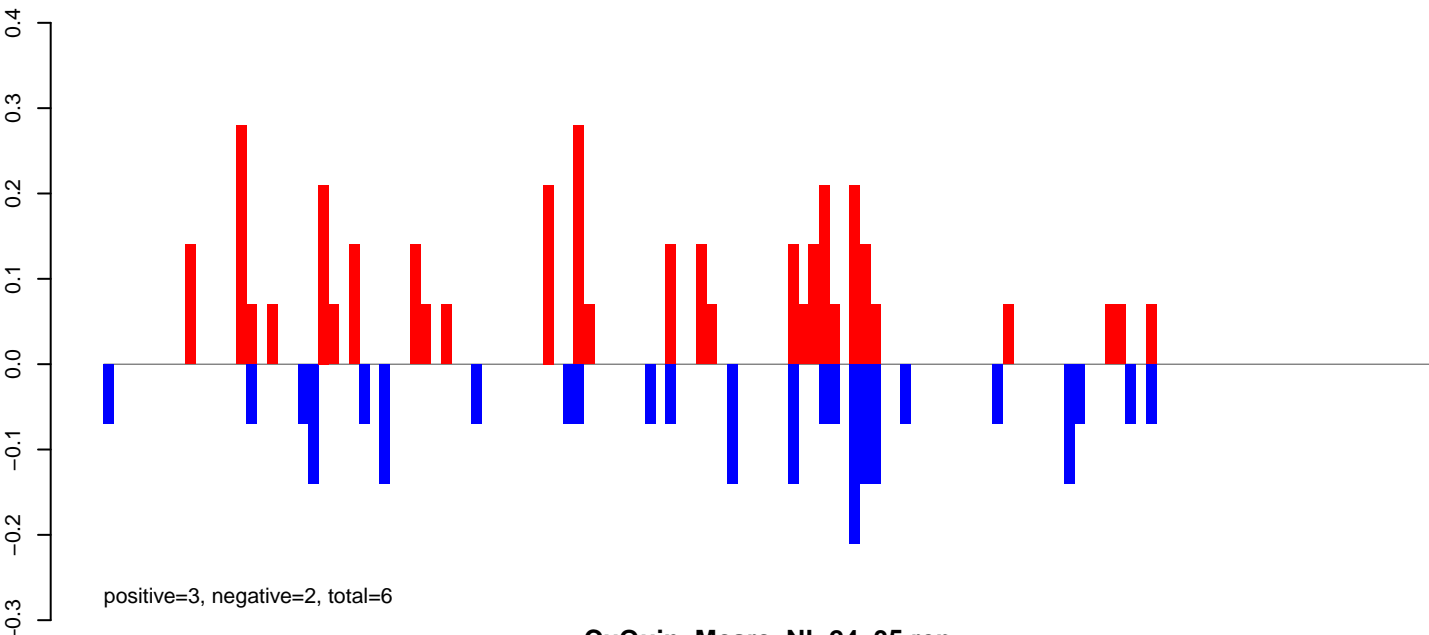
CuQuin_Mcarc_NL.24_35.rep



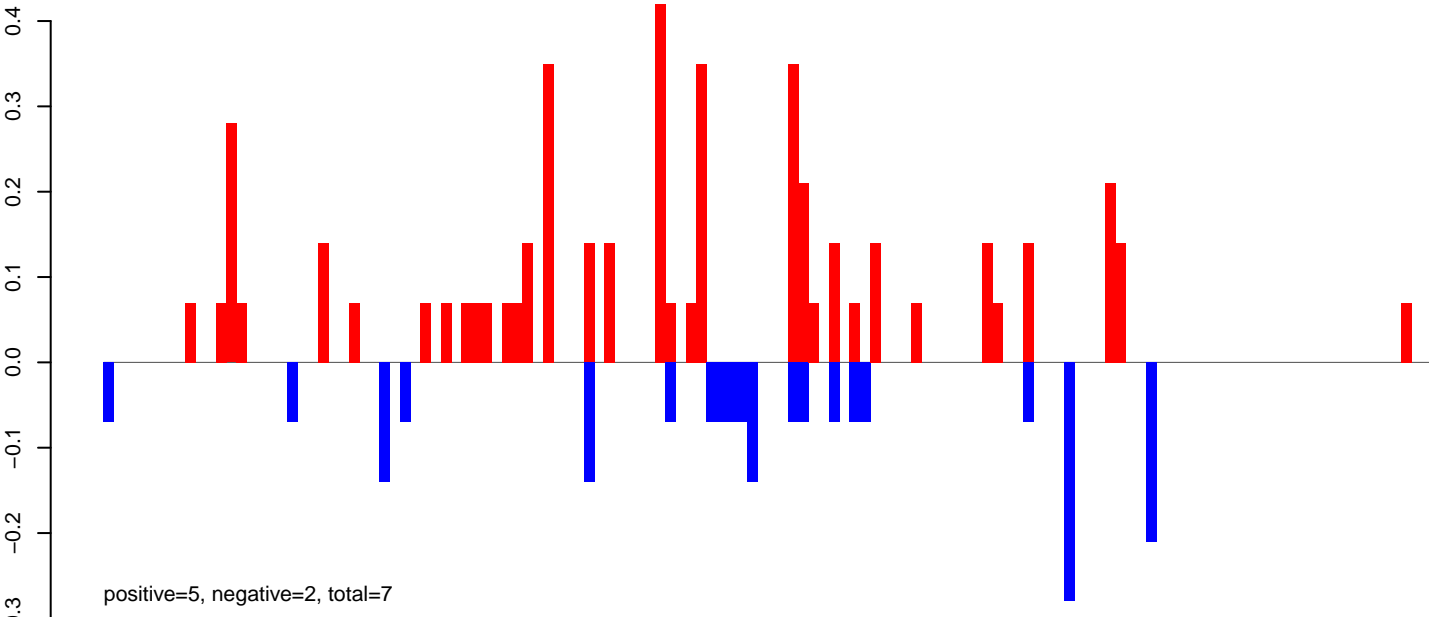
CuQuin_Mcarc_NL.rep



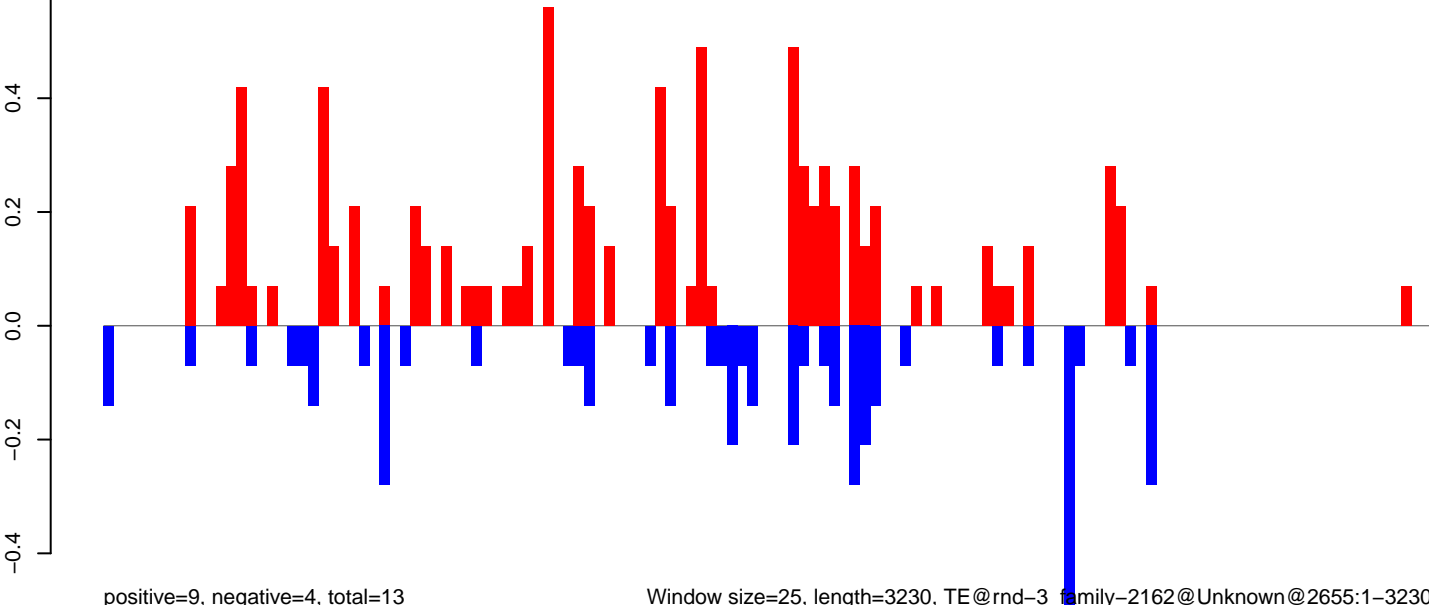
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

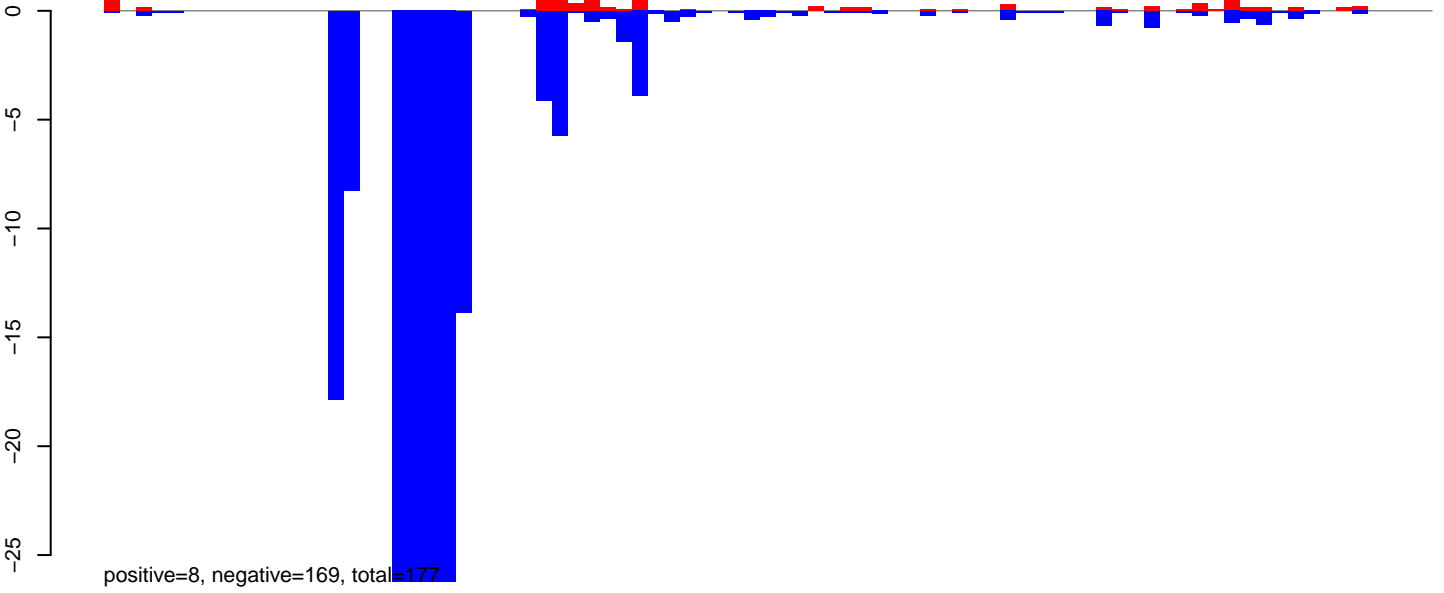


CuQuin_Mcarc_NL.rep

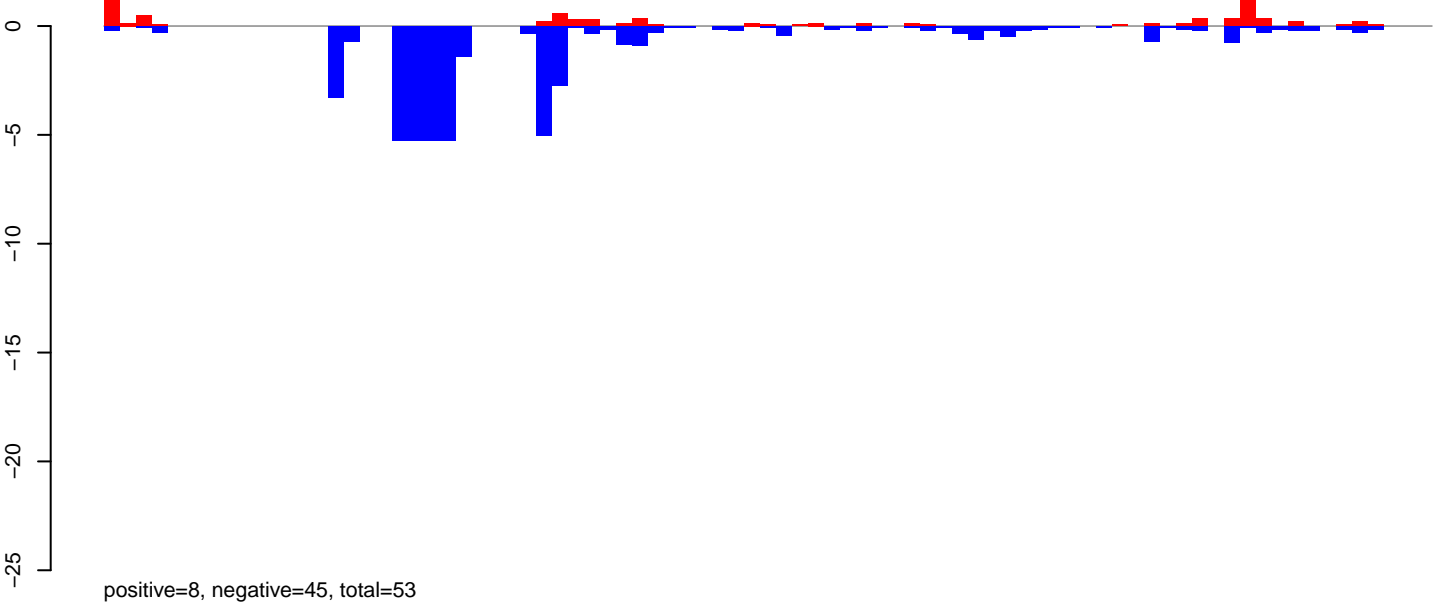


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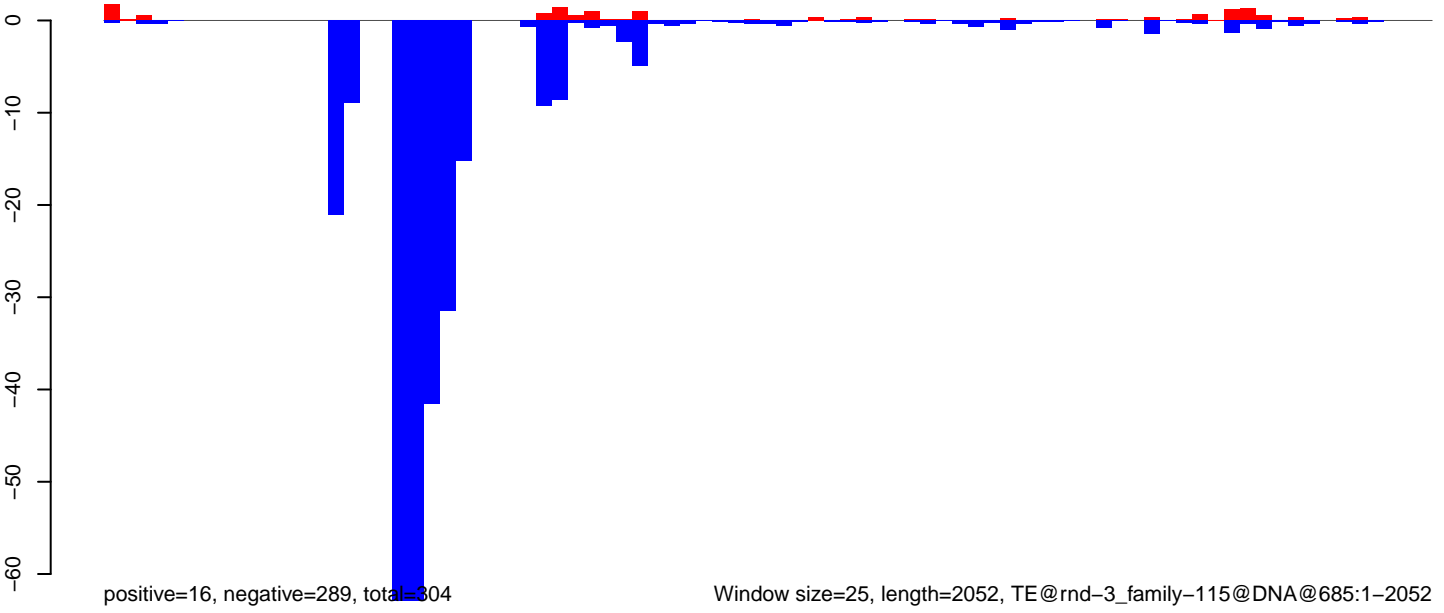
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CuQuin_Mcarc_NL.24_35.rep



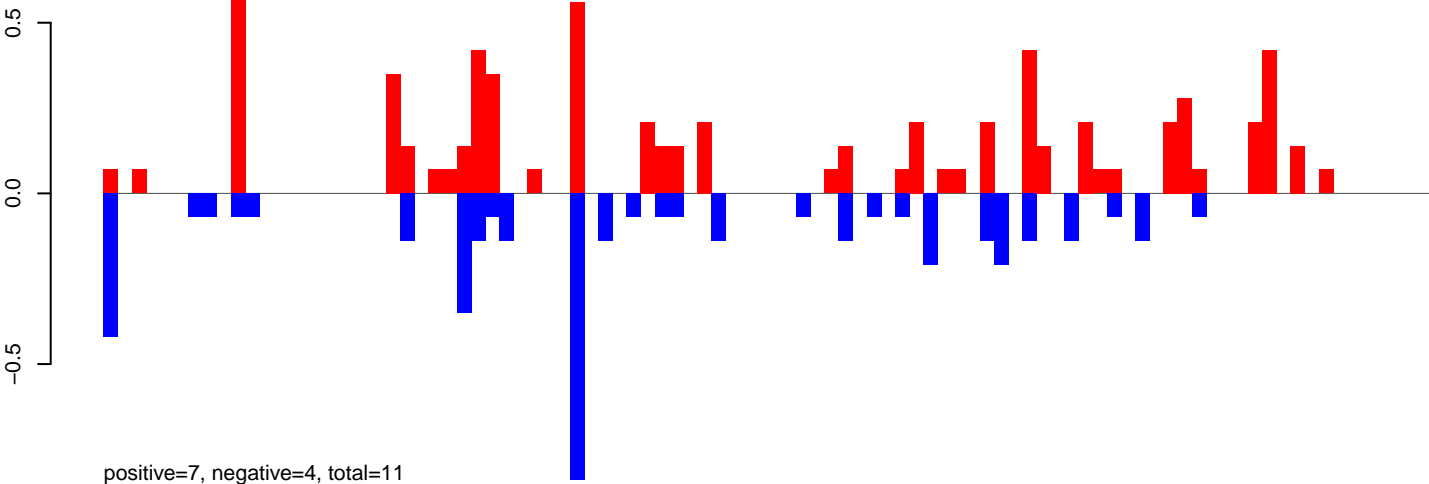
CuQuin_Mcarc_NL.rep



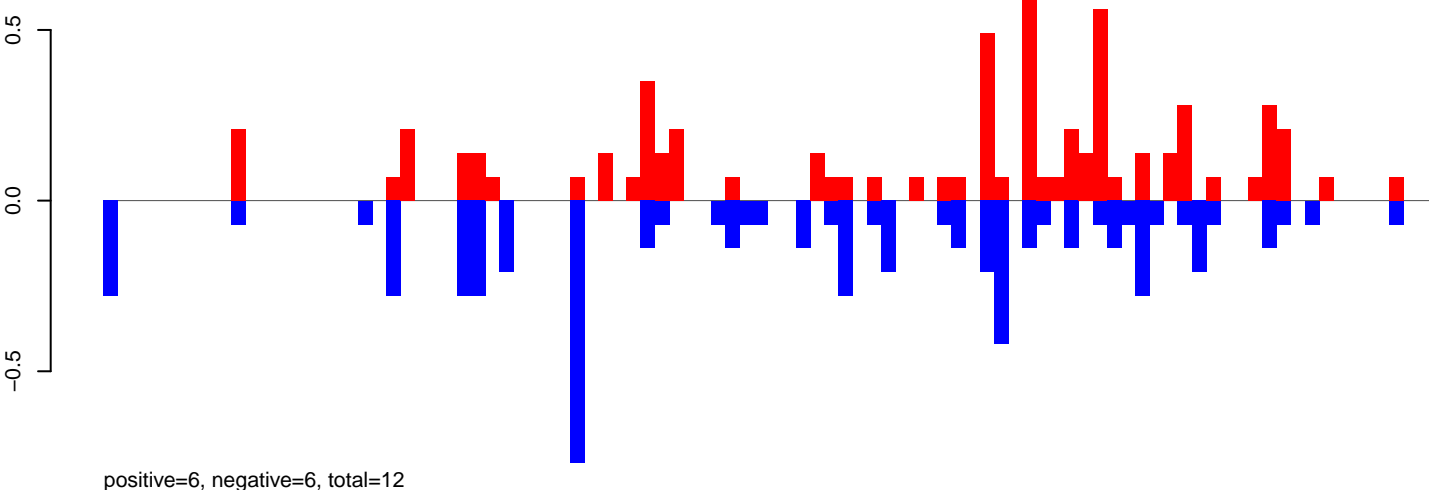
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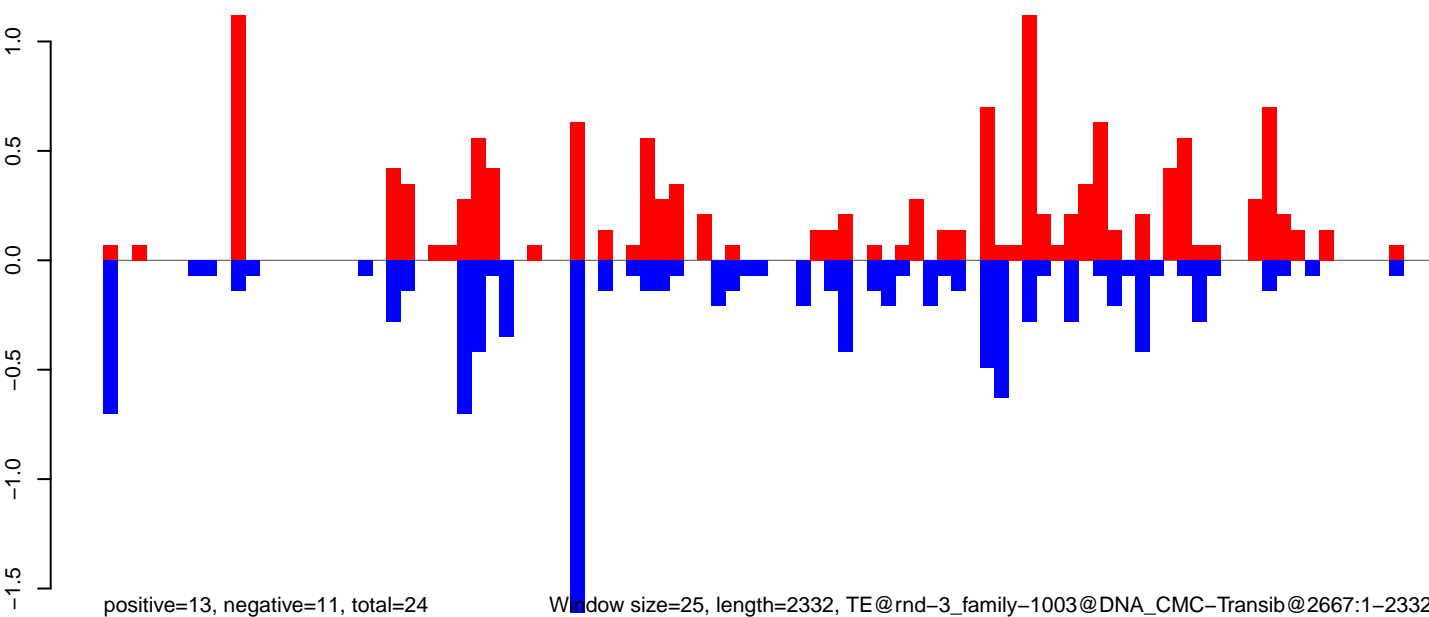
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

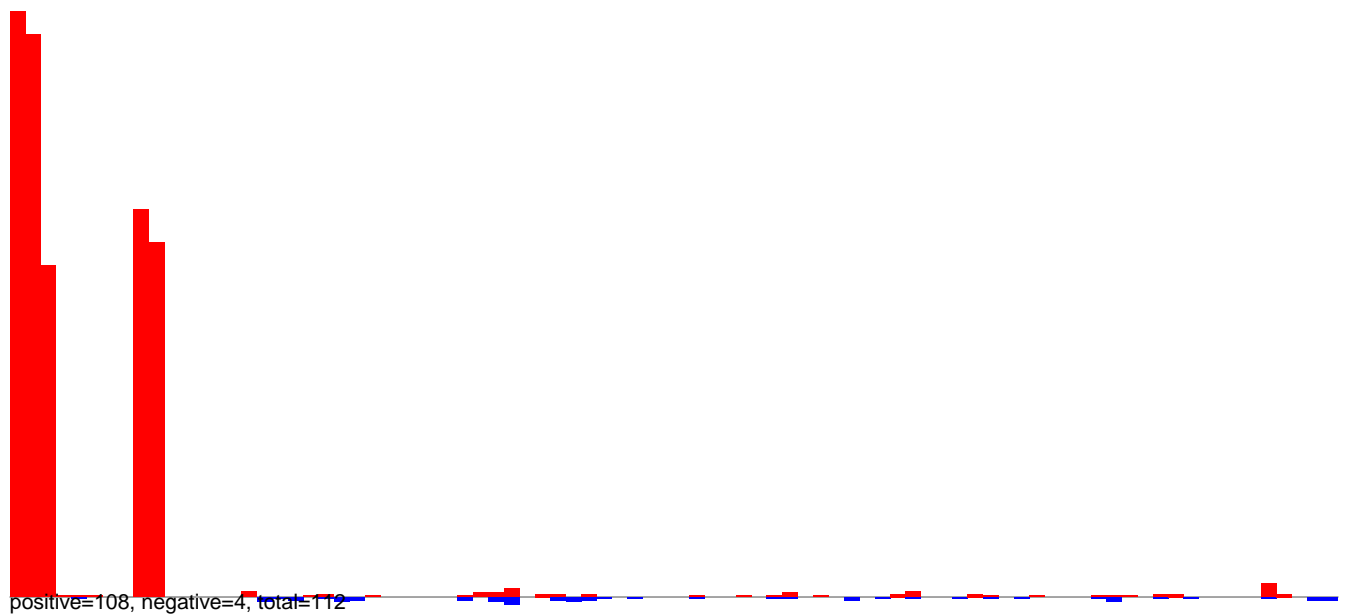


CuQuin_Mcarc_NL.rep

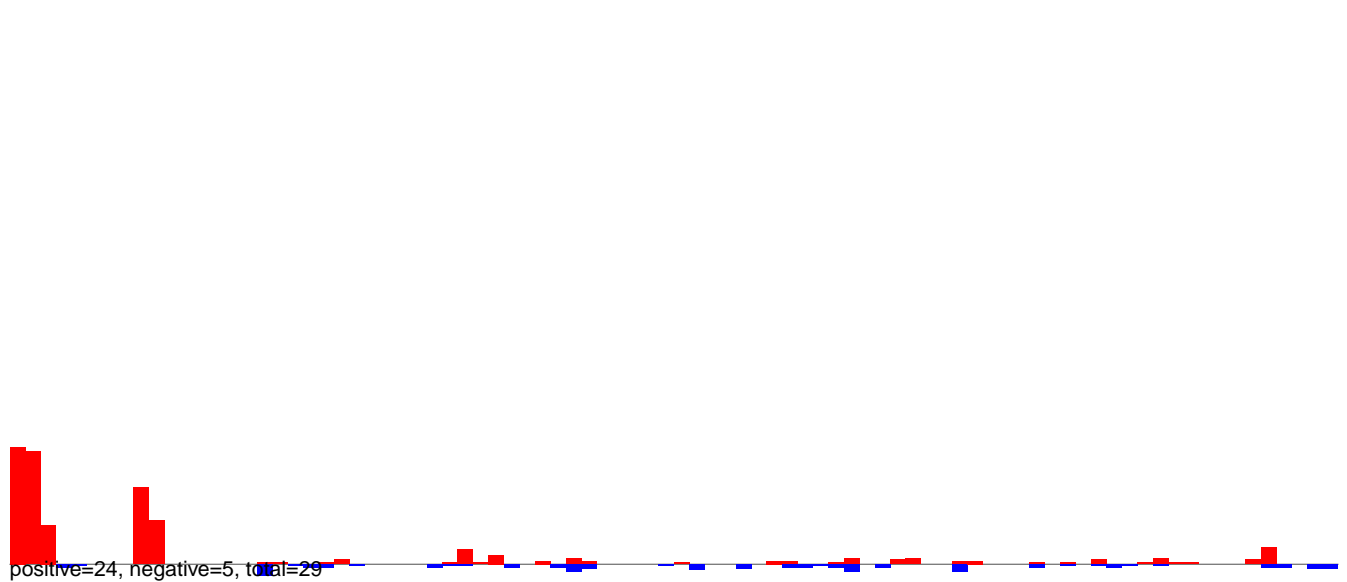


Window size=25, length=2332, TE@rnd-3_family-1003@DNA_CMC-Transib@2667:1-2332

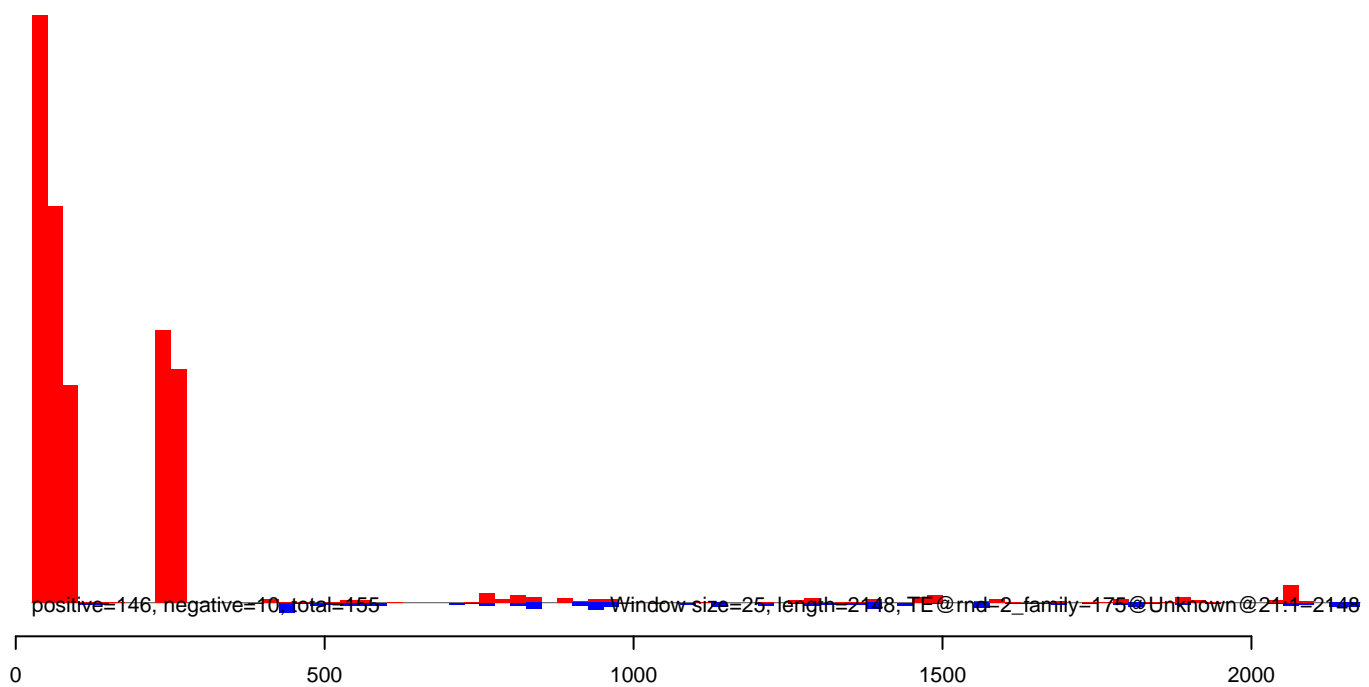
CuQuin_Mcarc_NL.18_23.rep



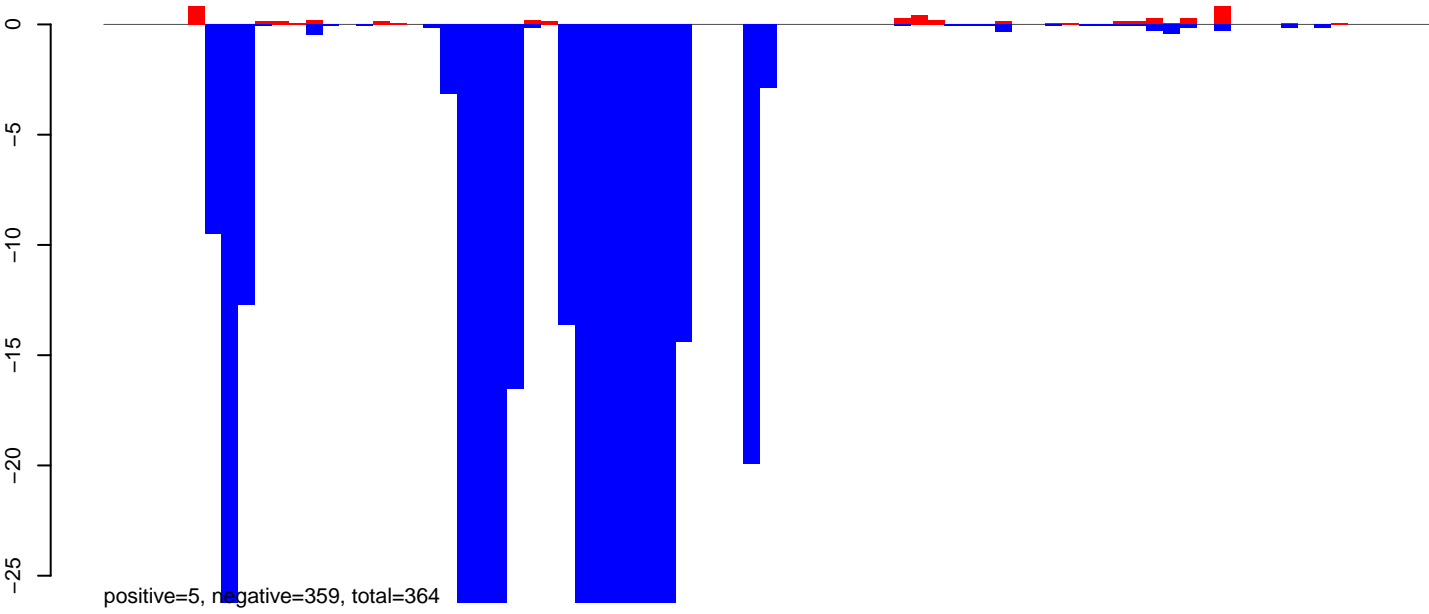
CuQuin_Mcarc_NL.24_35.rep



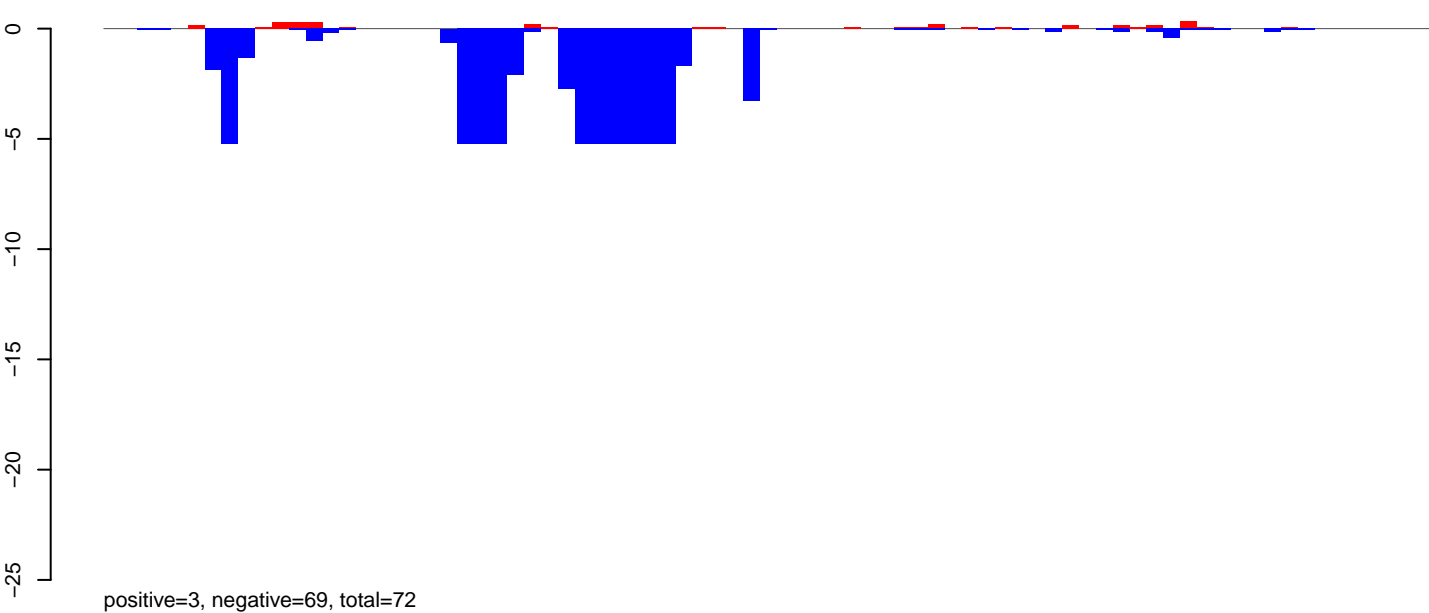
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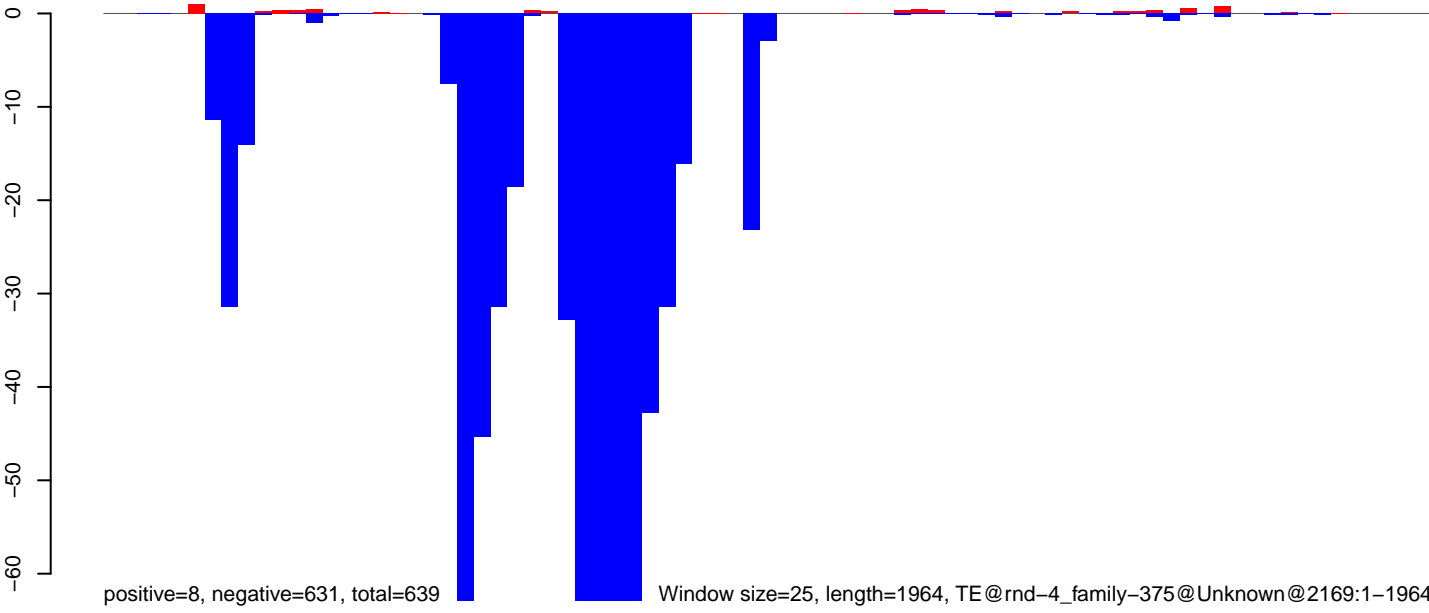
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CuQuin_Mcarc_NL.24_35.rep



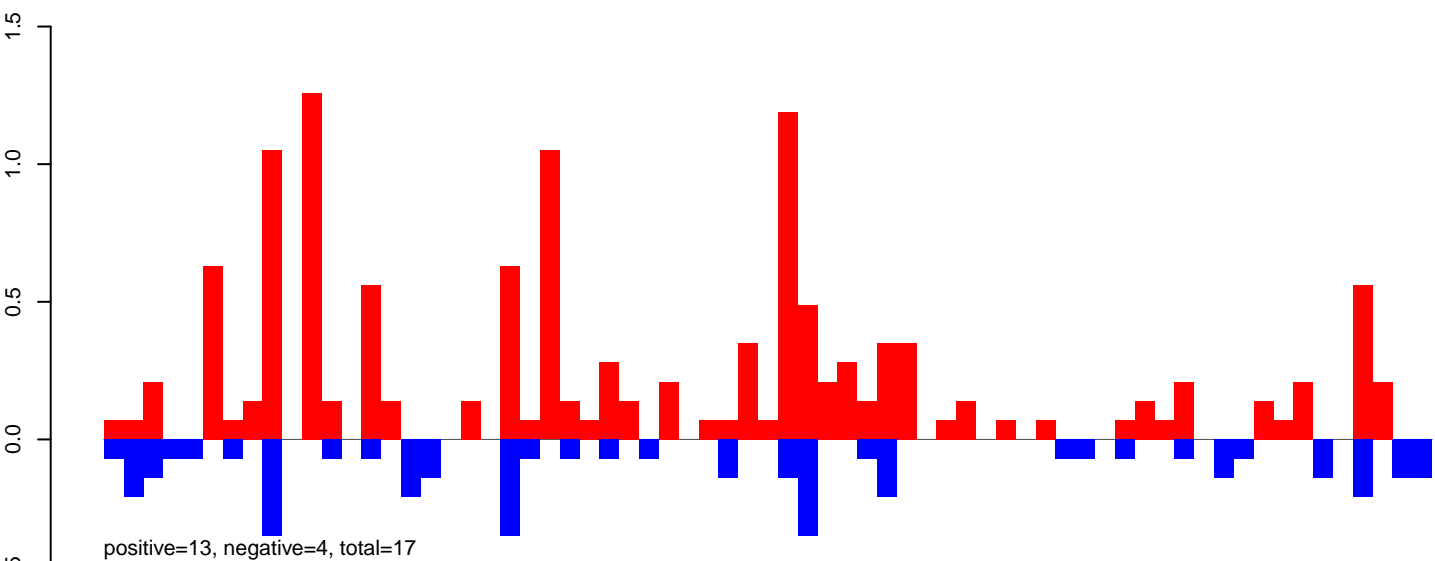
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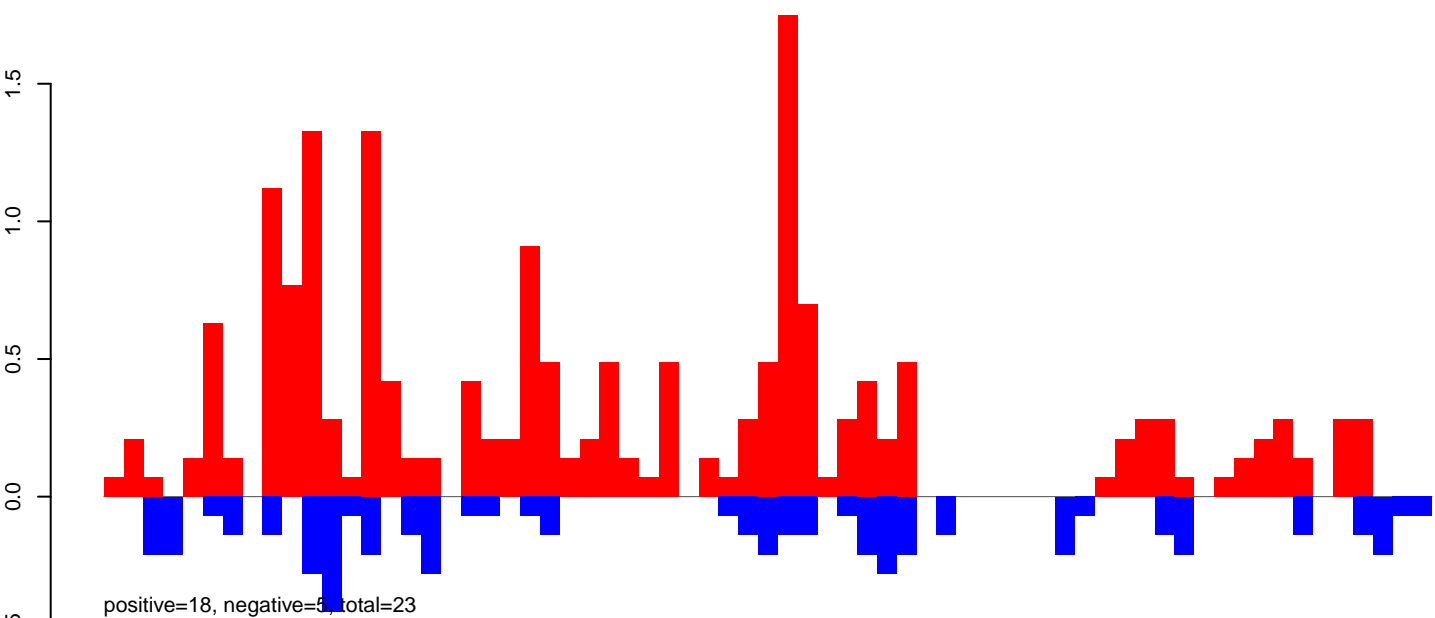
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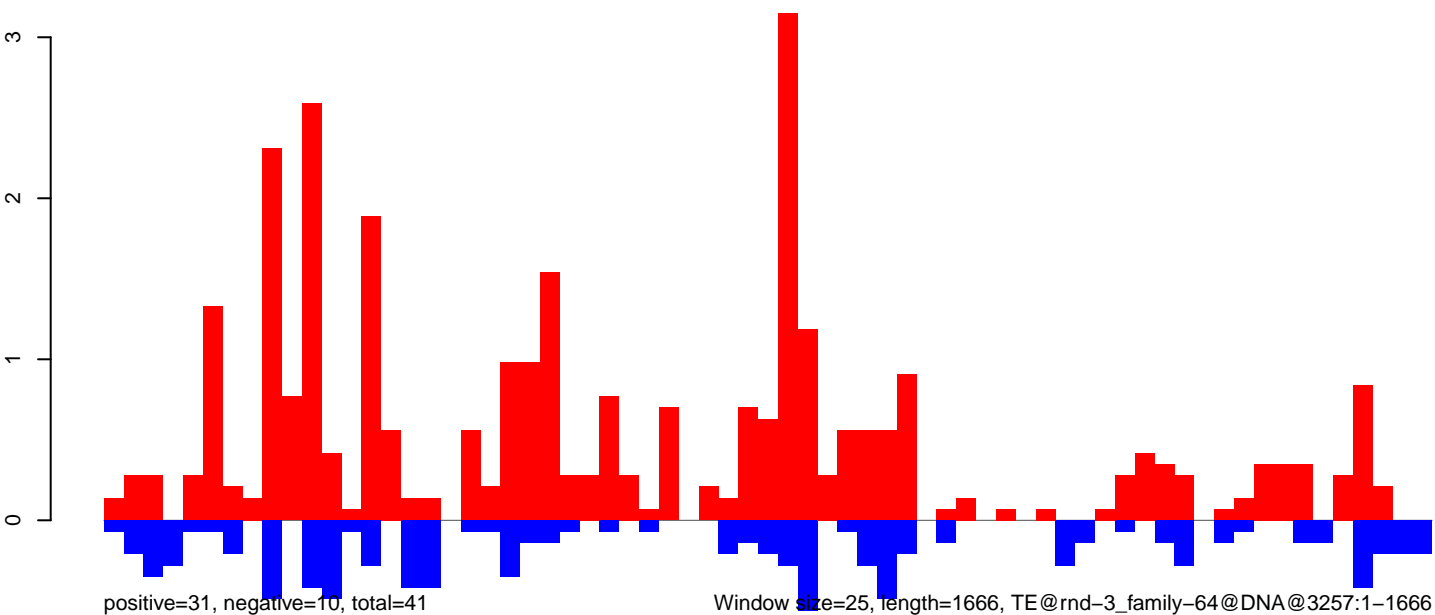
CuQuin_Mcarc_NL.18_23.rep



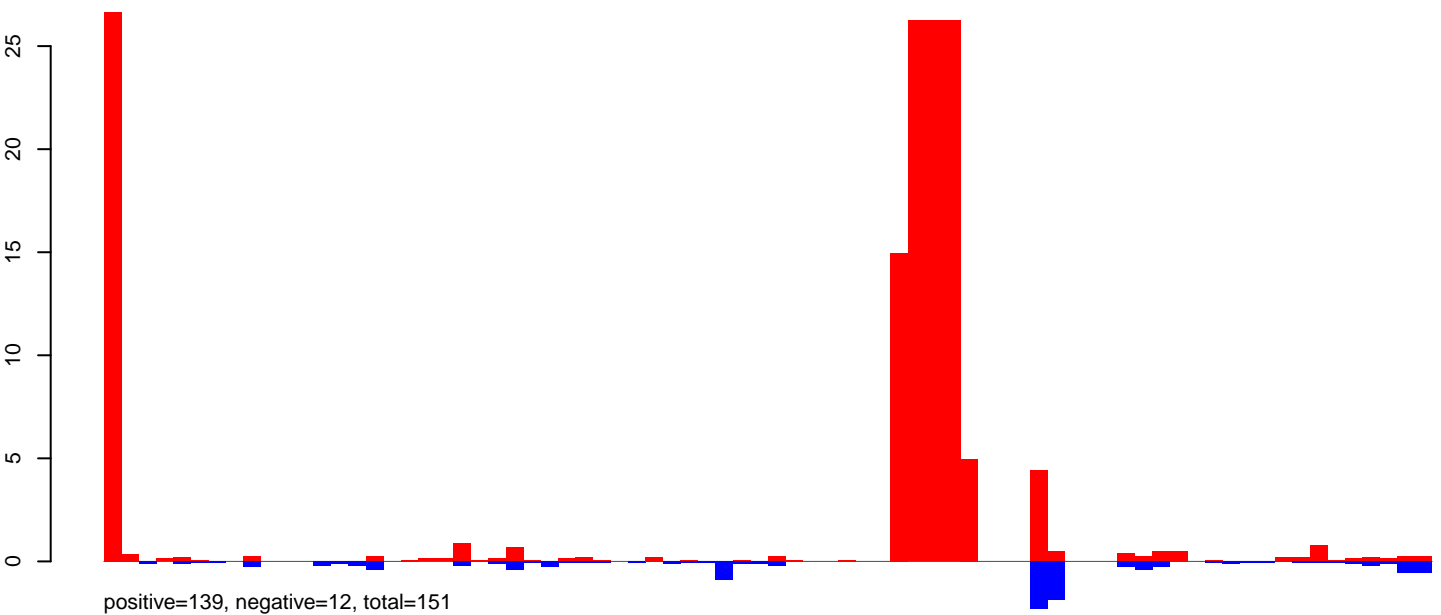
CuQuin_Mcarc_NL.24_35.rep



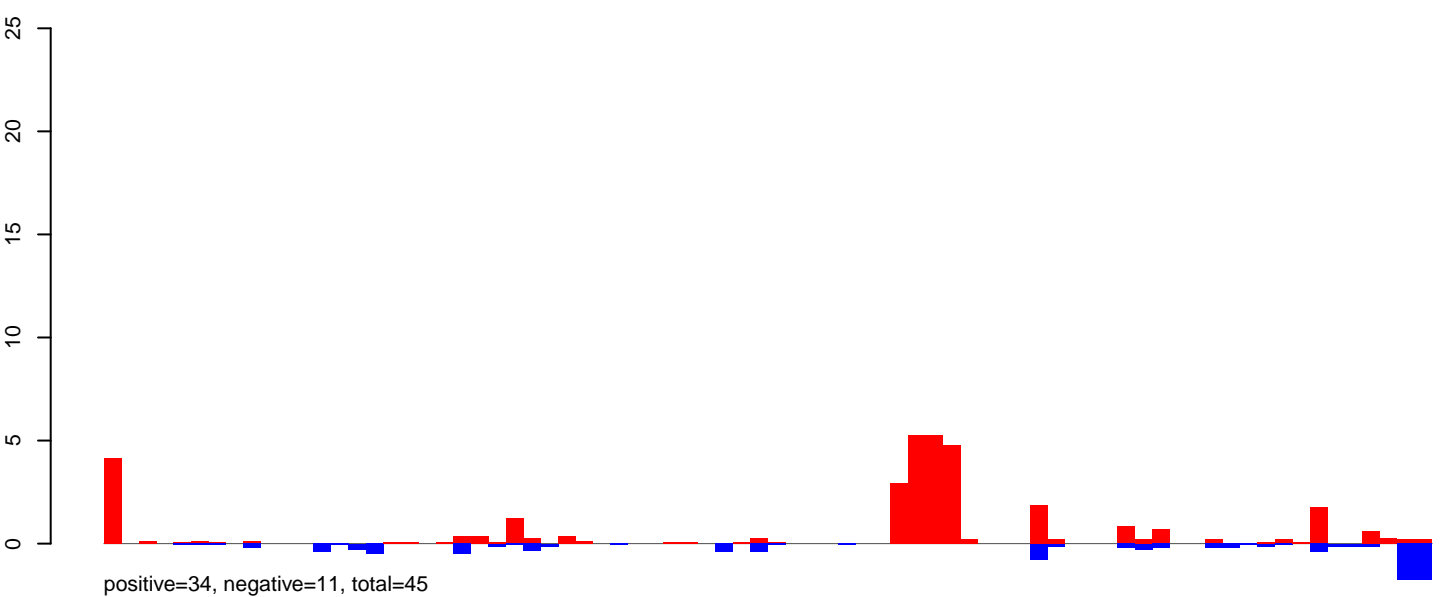
CuQuin_Mcarc_NL.rep



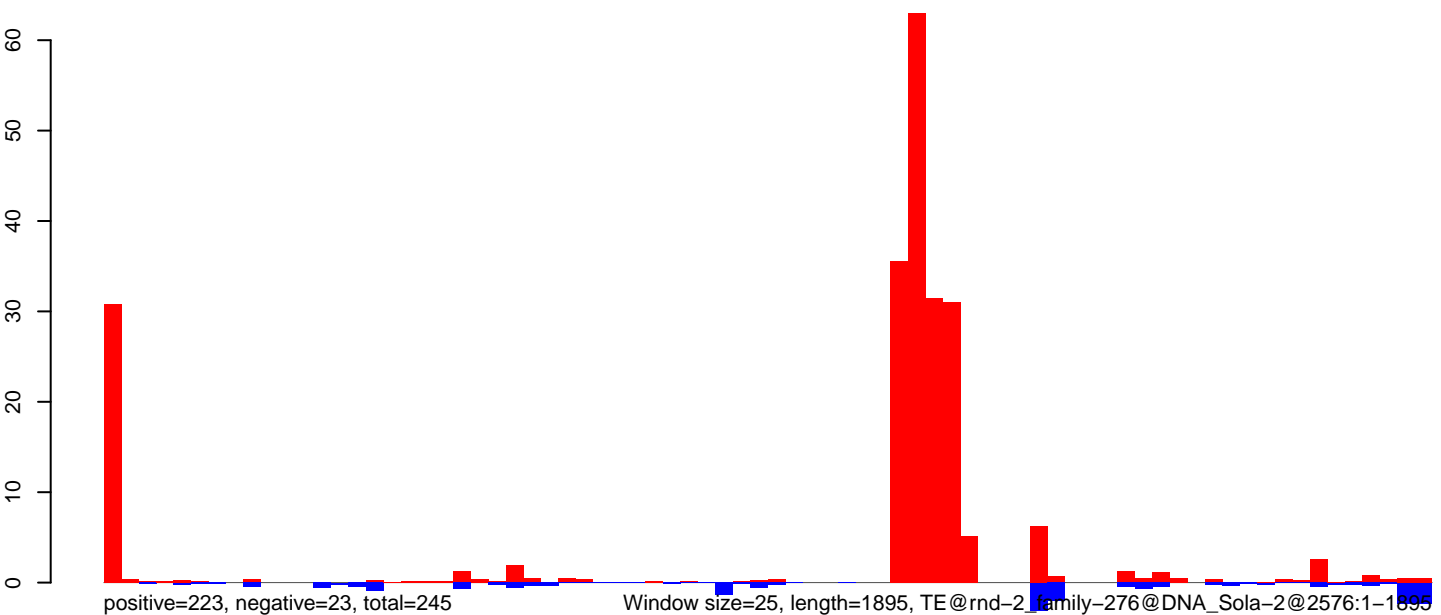
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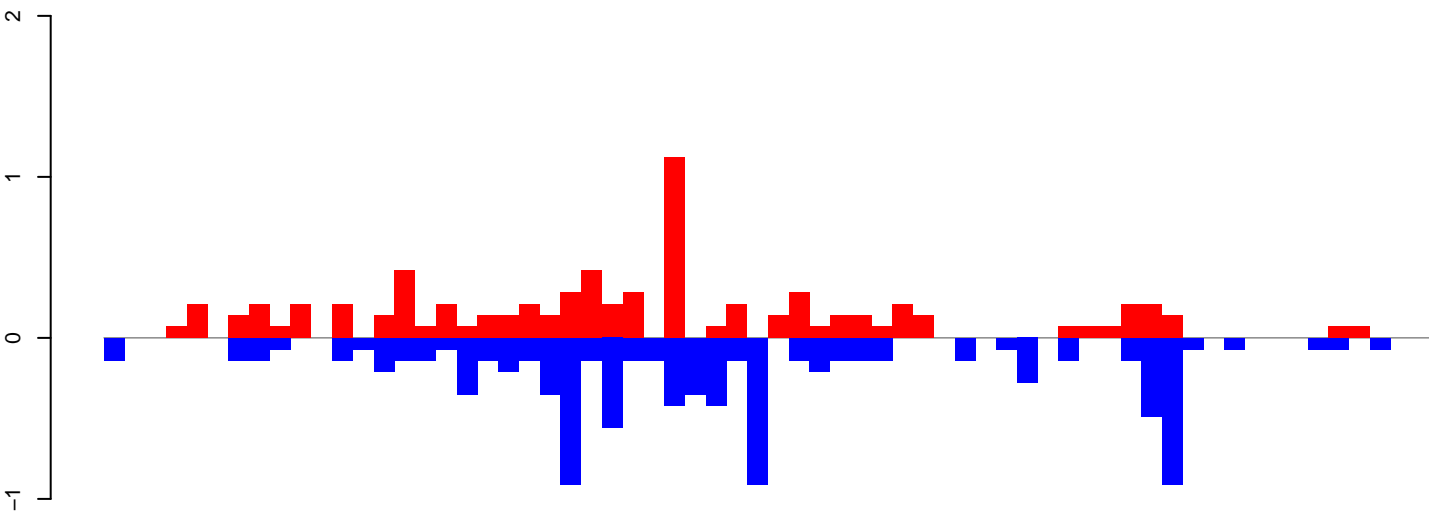
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CuQuin_Mcarc_NL.rep

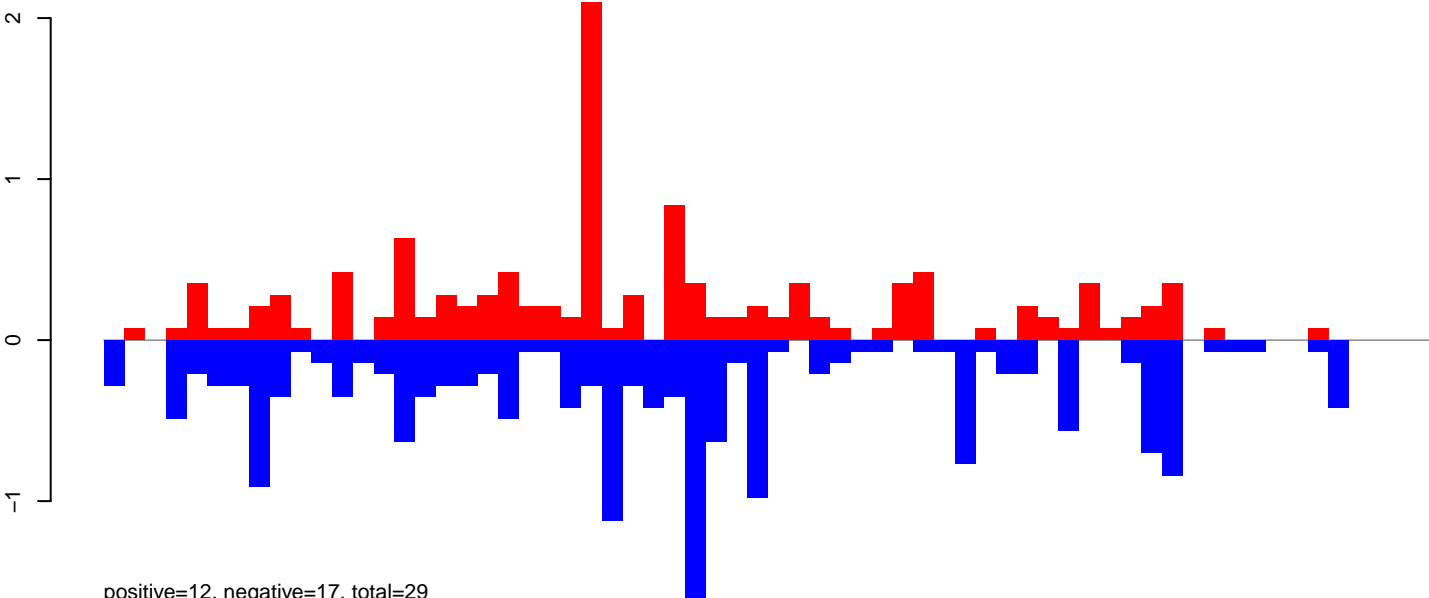


CuQuin_Mcarc_NL.18_23.rep



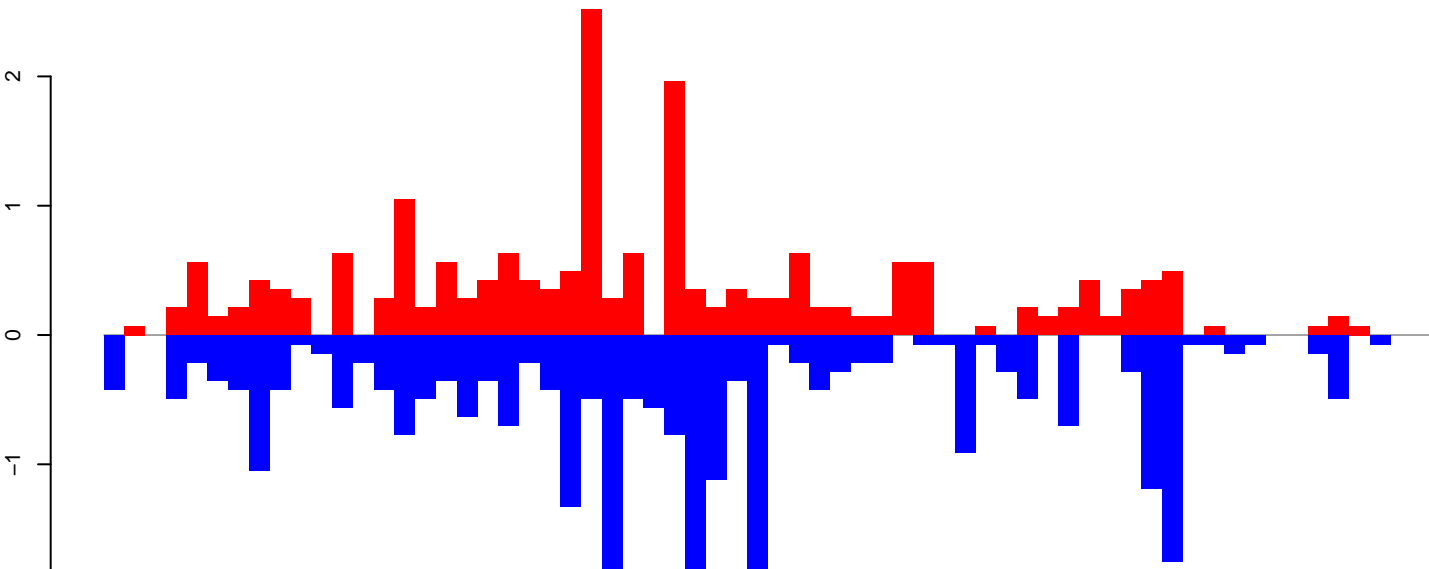
positive=7, negative=10, total=17

CuQuin_Mcarc_NL.24_35.rep



positive=12, negative=17, total=29

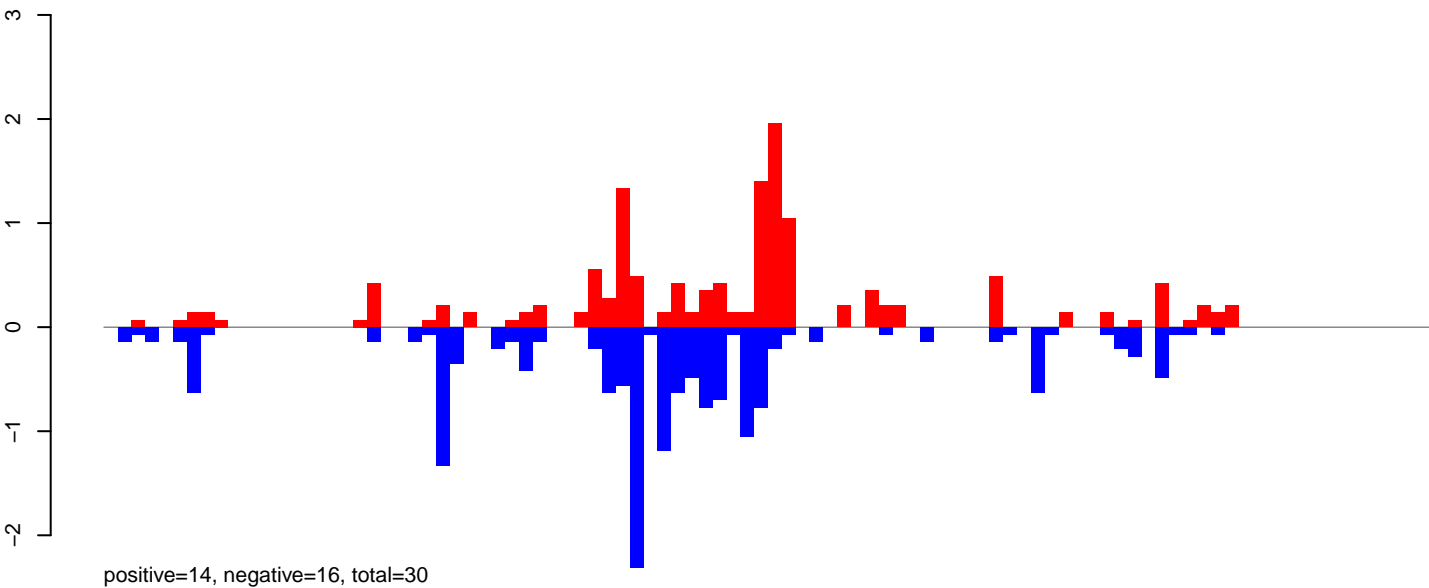
CuQuin_Mcarc_NL.rep



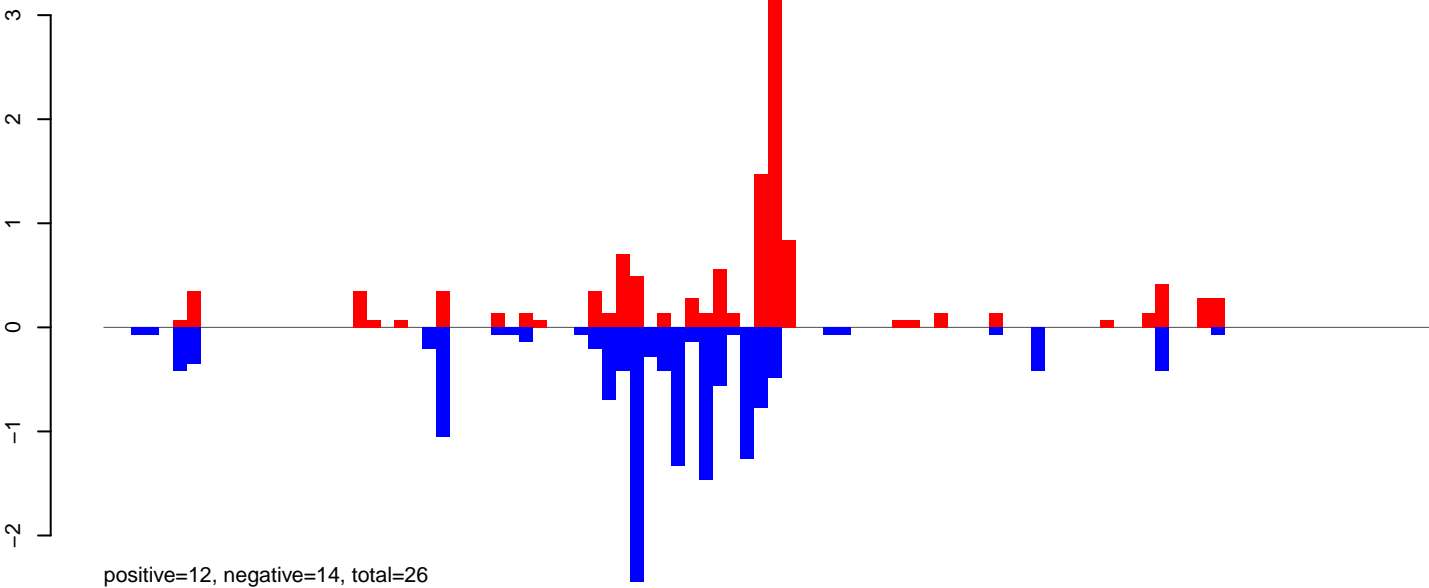
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Window size=25, length=1577, TE@rnd-5_family-4105@Unknown@2485:1-1577

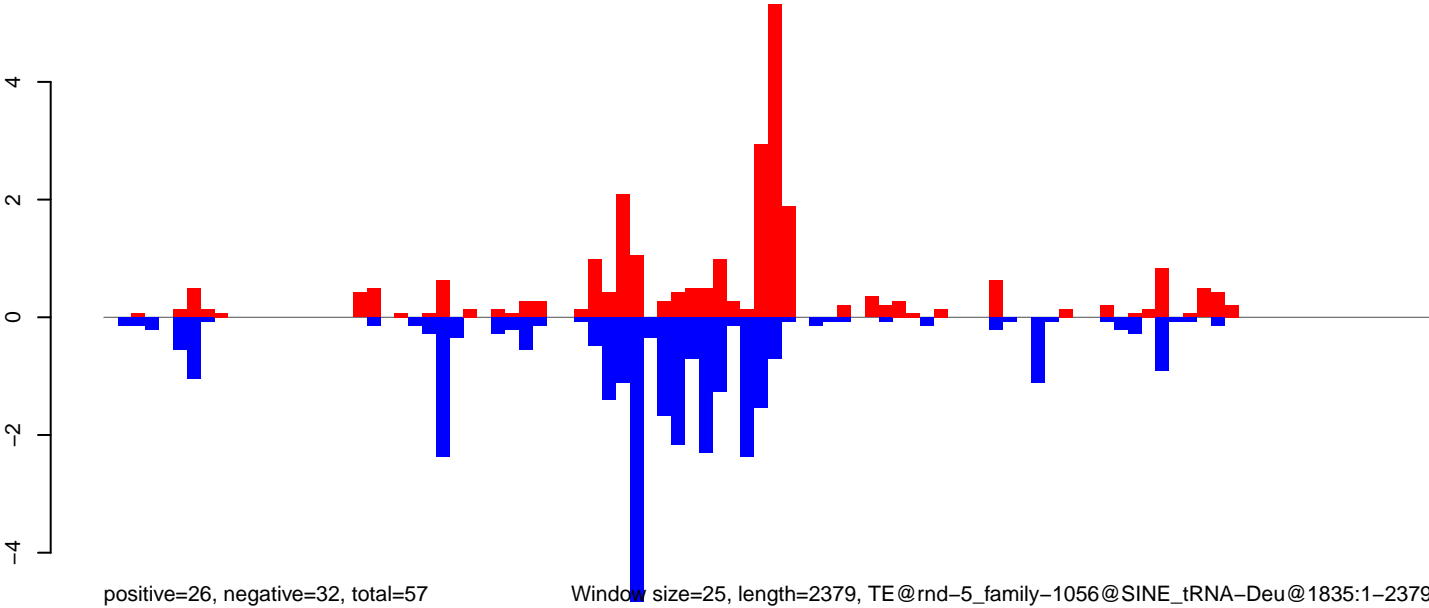
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CuQuin_Mcarc_NL.24_35.rep

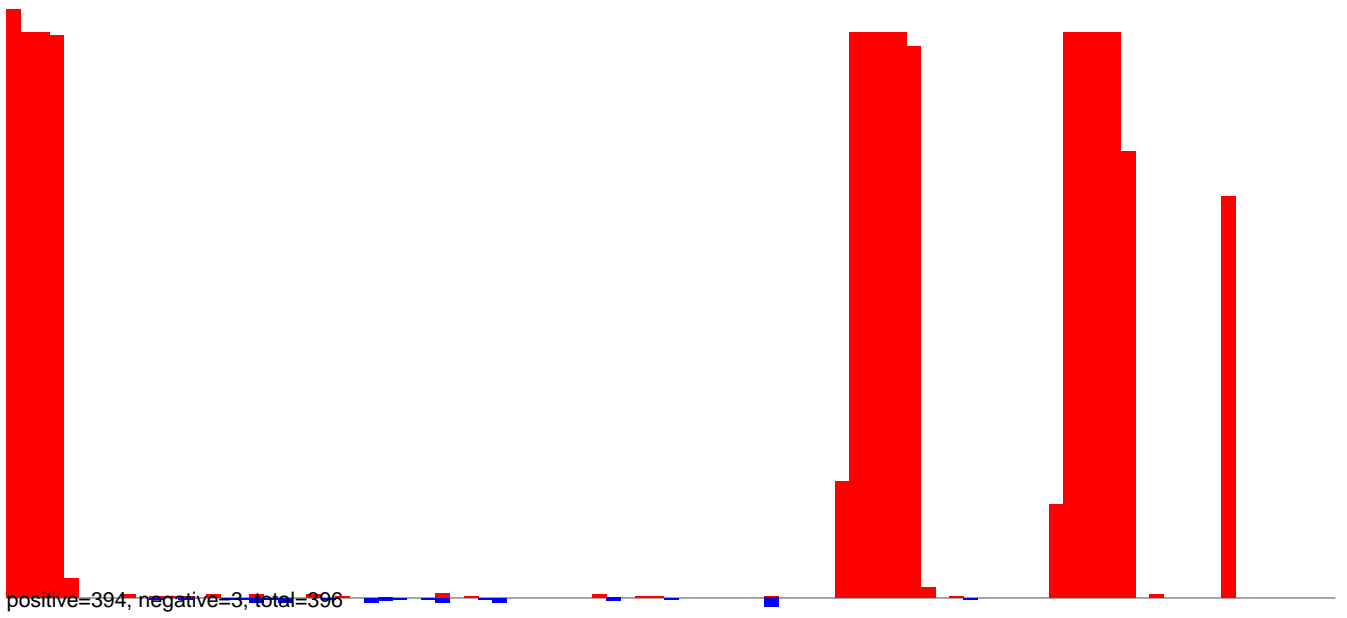


CuQuin_Mcarc_NL.rep

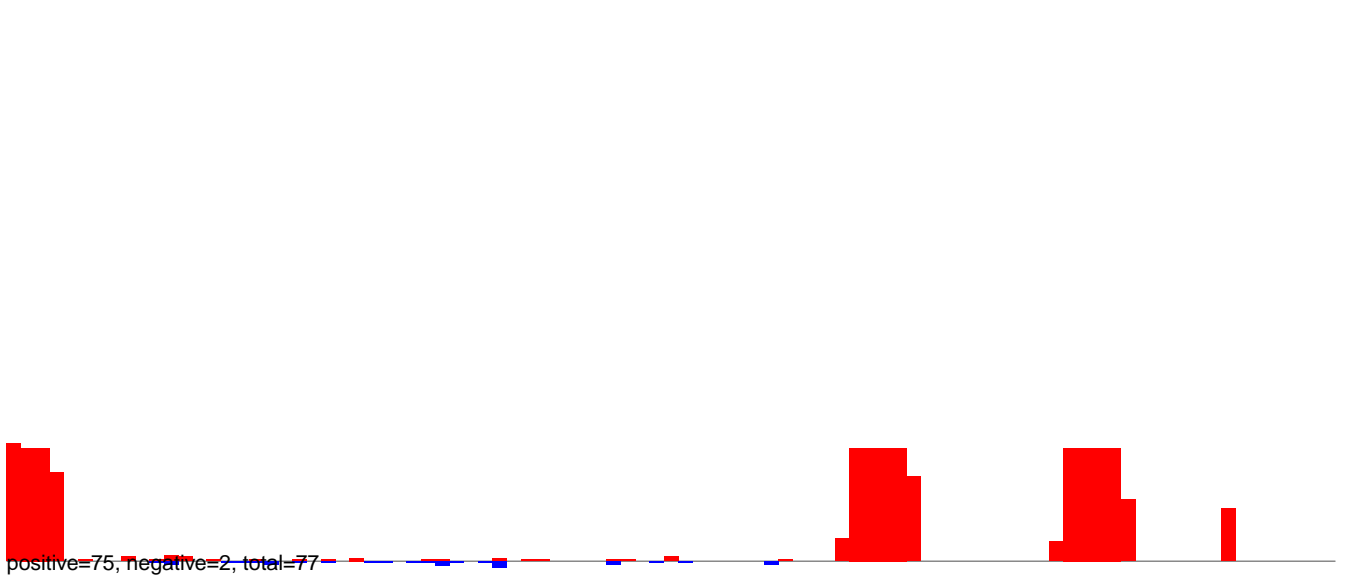


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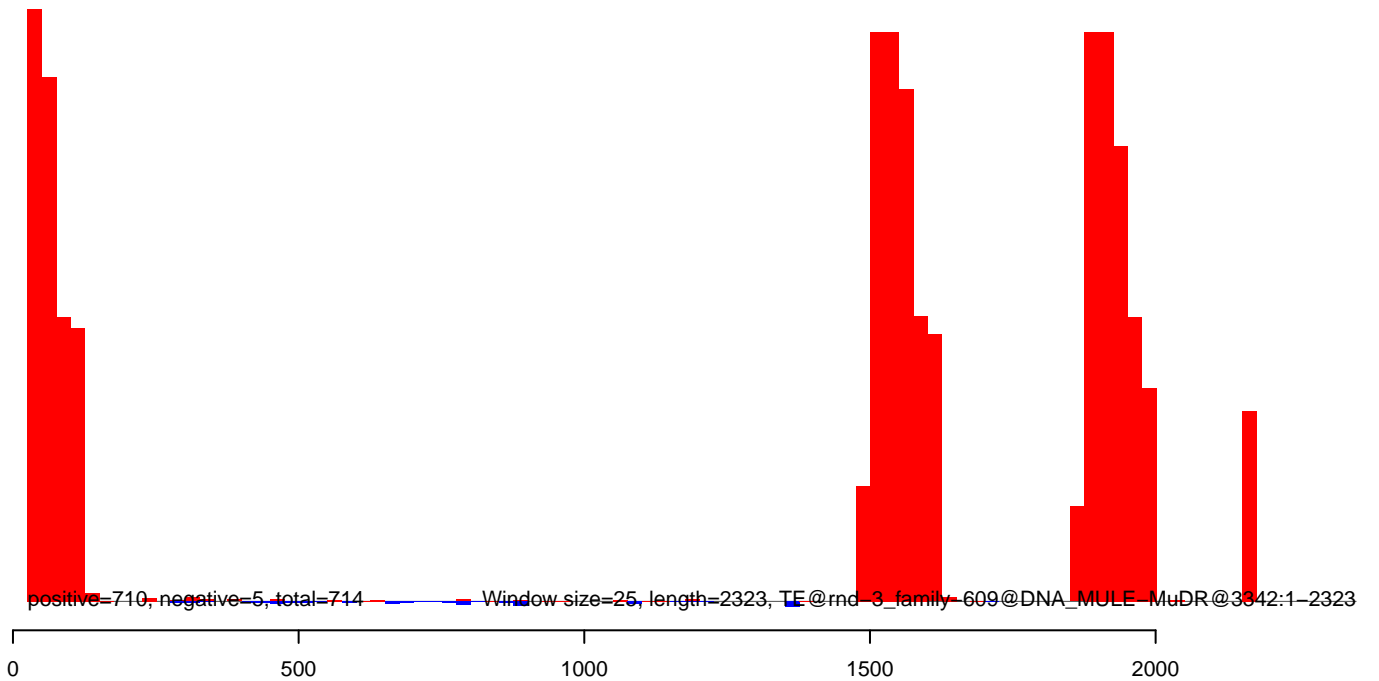
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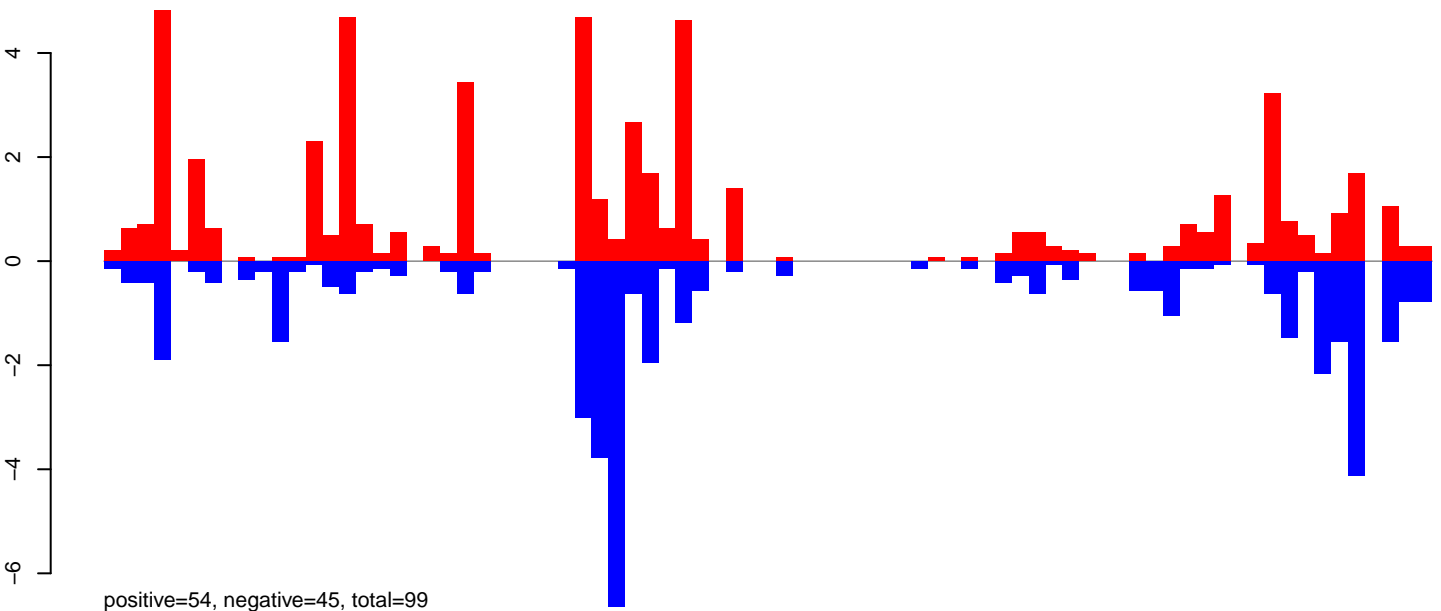
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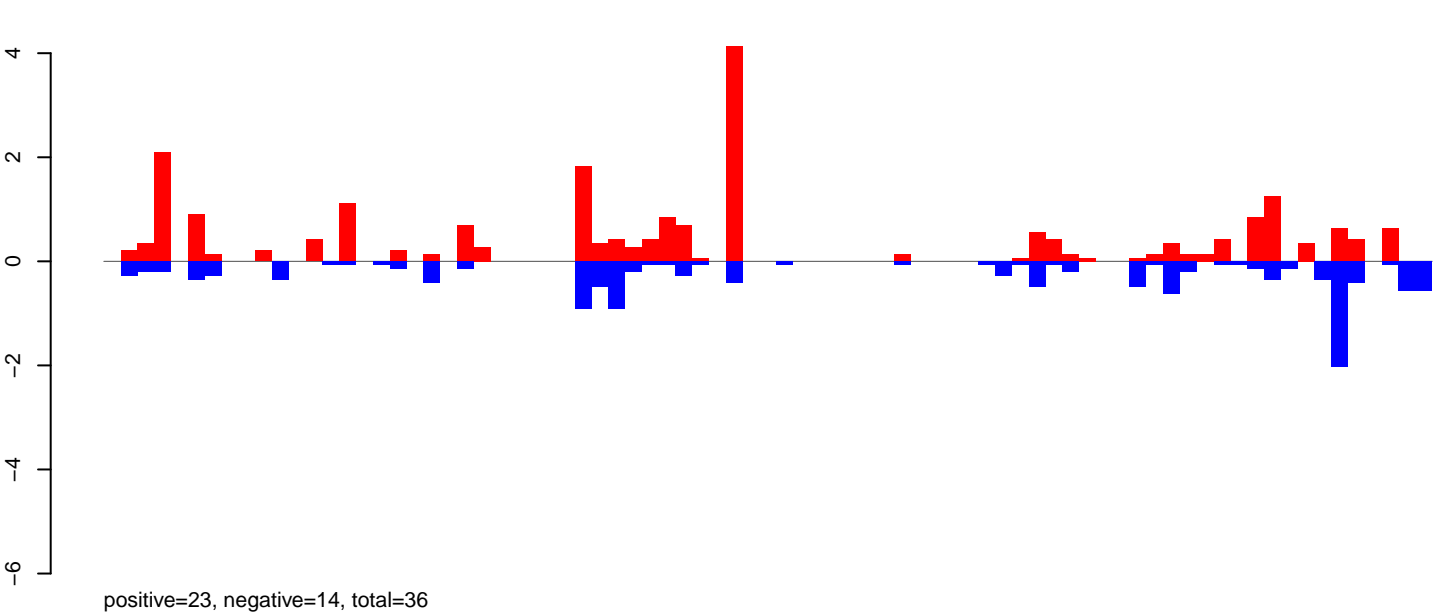
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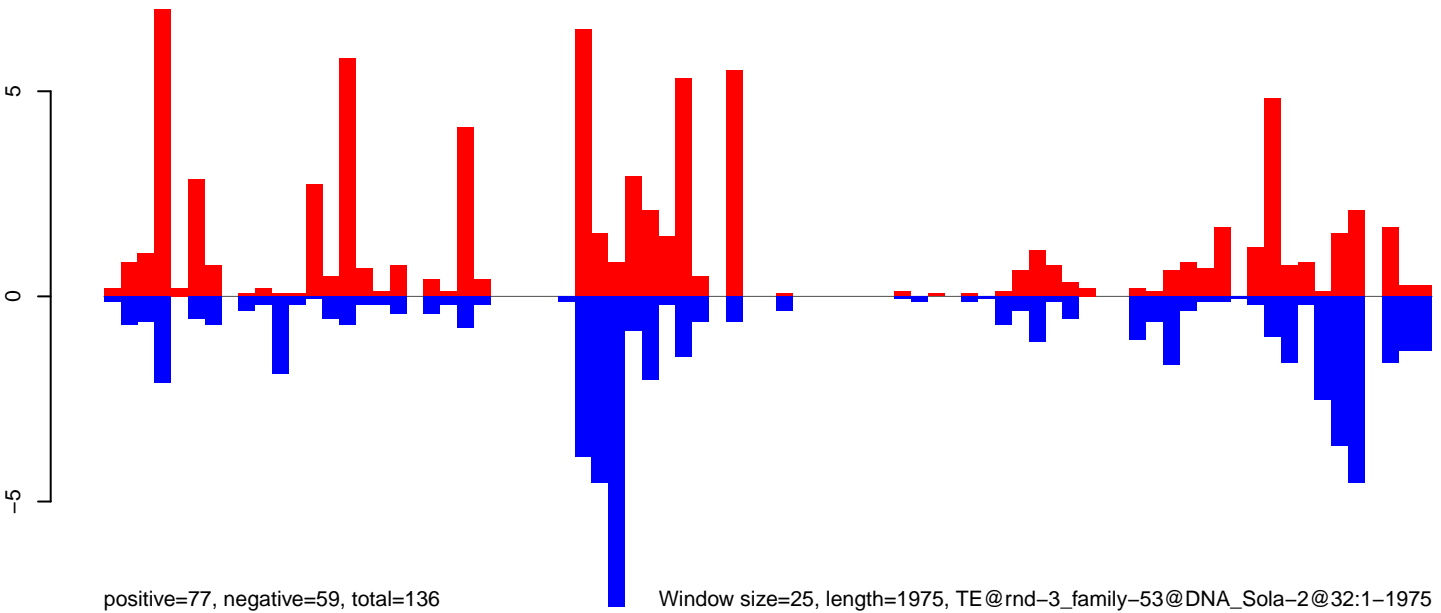
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CuQuin_Mcarc_NL.24_35.rep



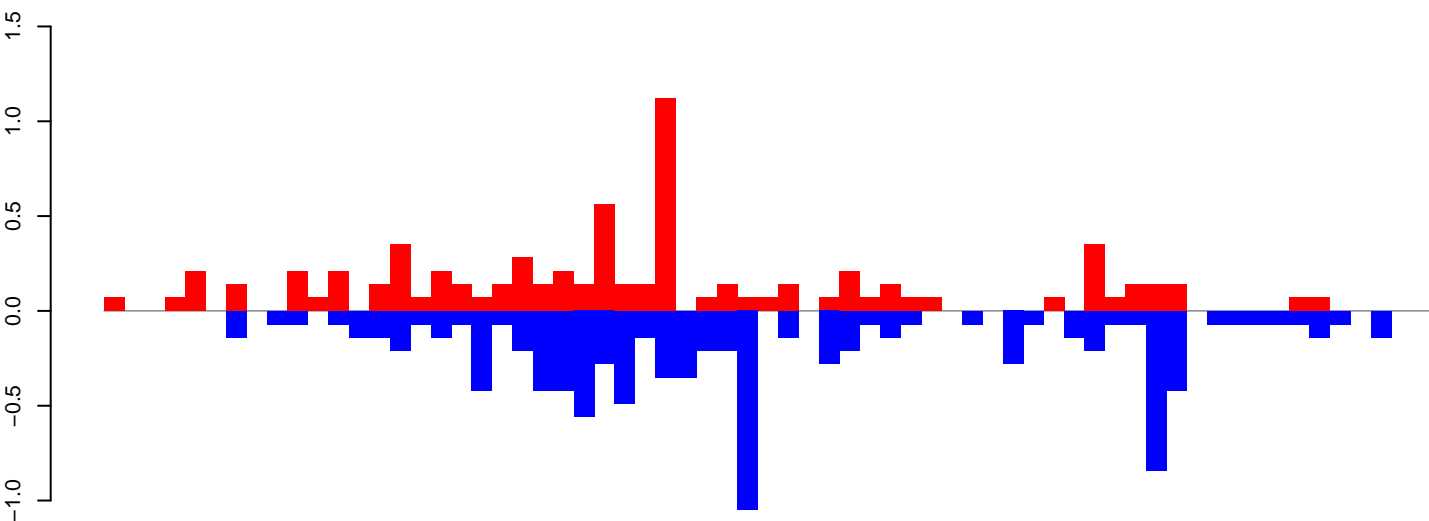
CuQuin_Mcarc_NL.rep



Window size=25, length=1975, TE@rnd-3_family-53@DNA_Sola-2@32:1-1975

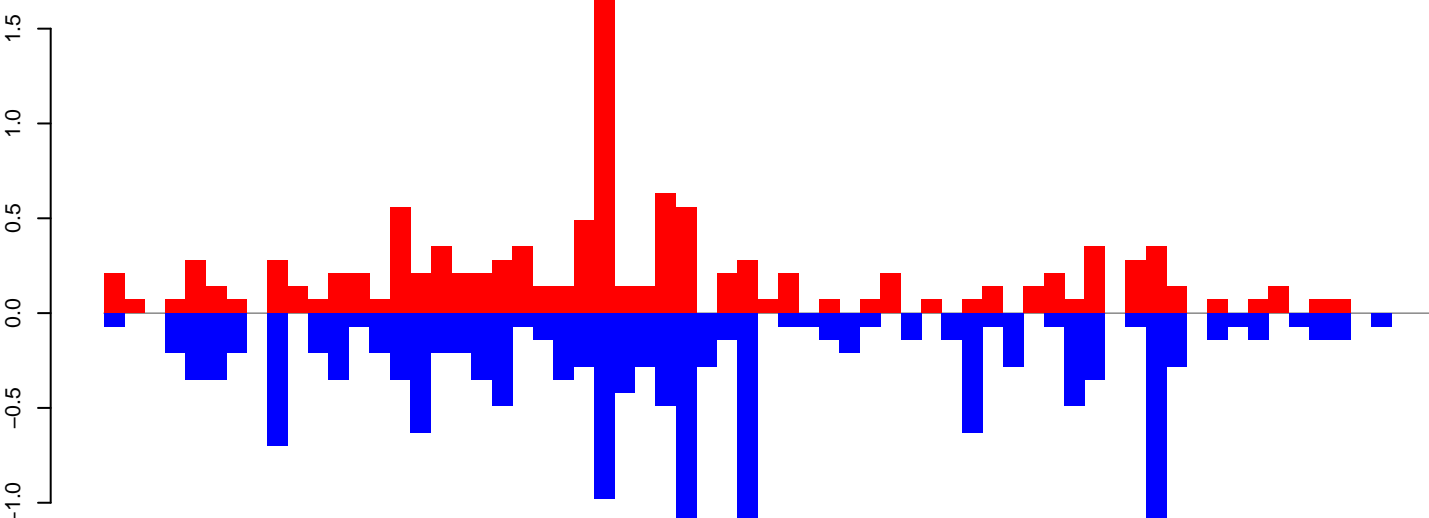
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CuQuin_Mcarc_NL.18_23.rep



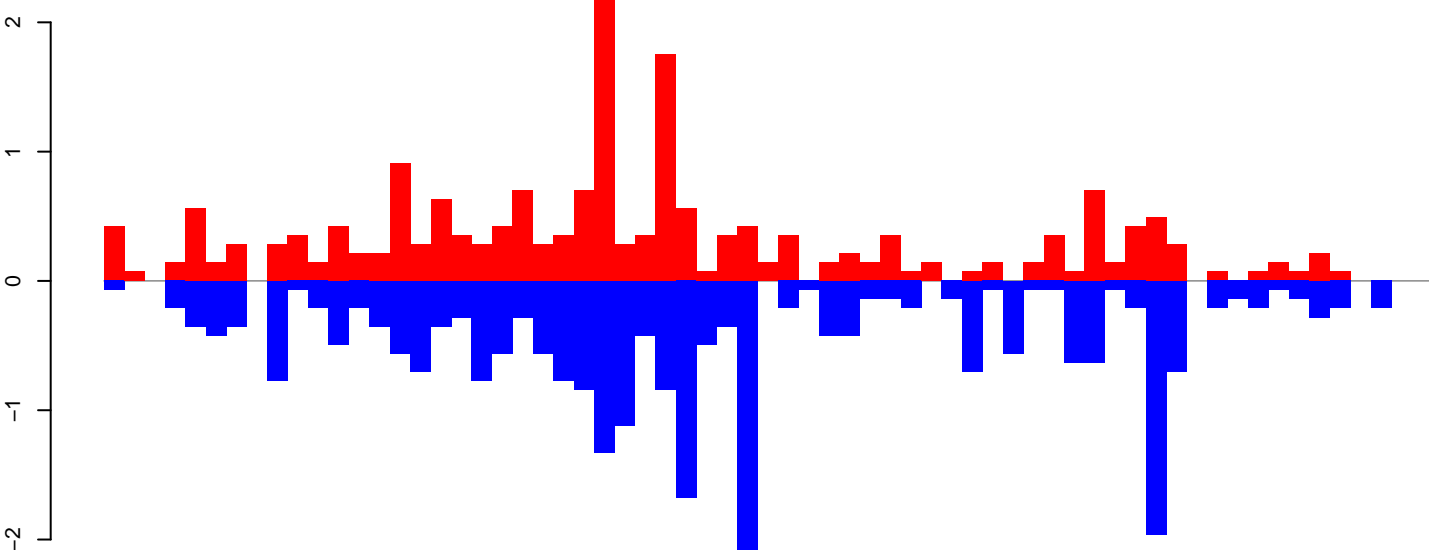
positive=7, negative=10, total=17

CuQuin_Mcarc_NL.24_35.rep



positive=11, negative=16, total=27

CuQuin_Mcarc_NL.rep

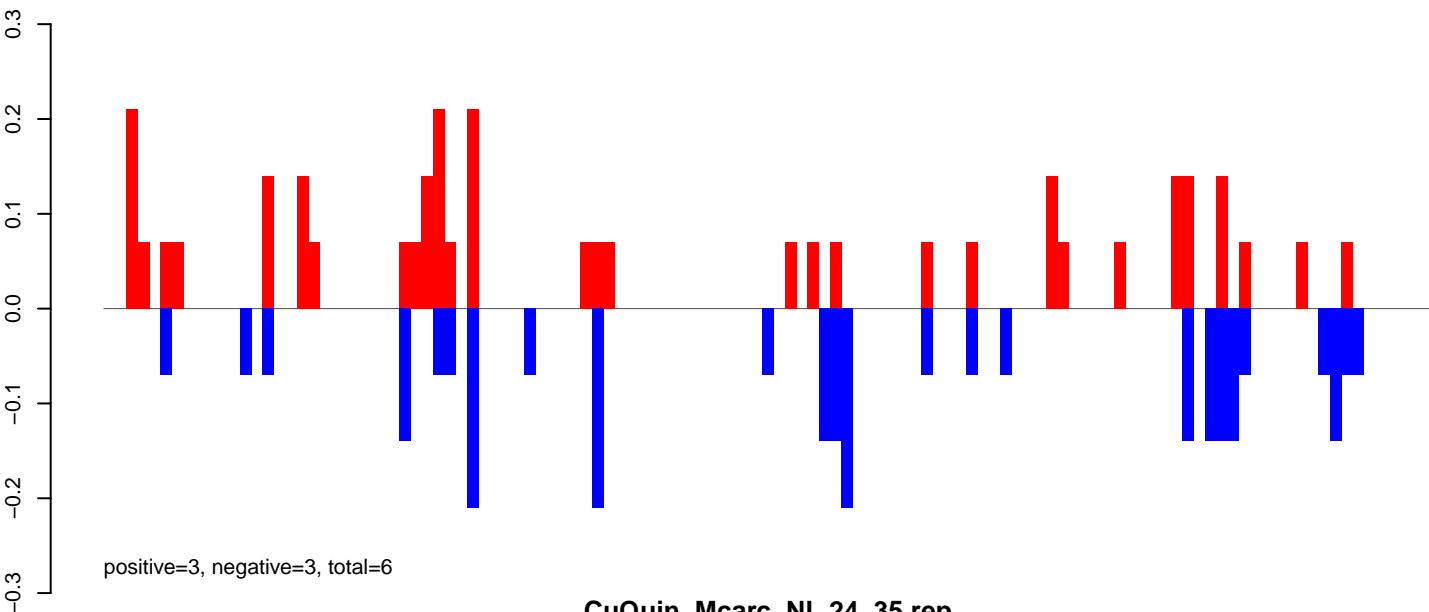


positive=19, negative=27, total=46

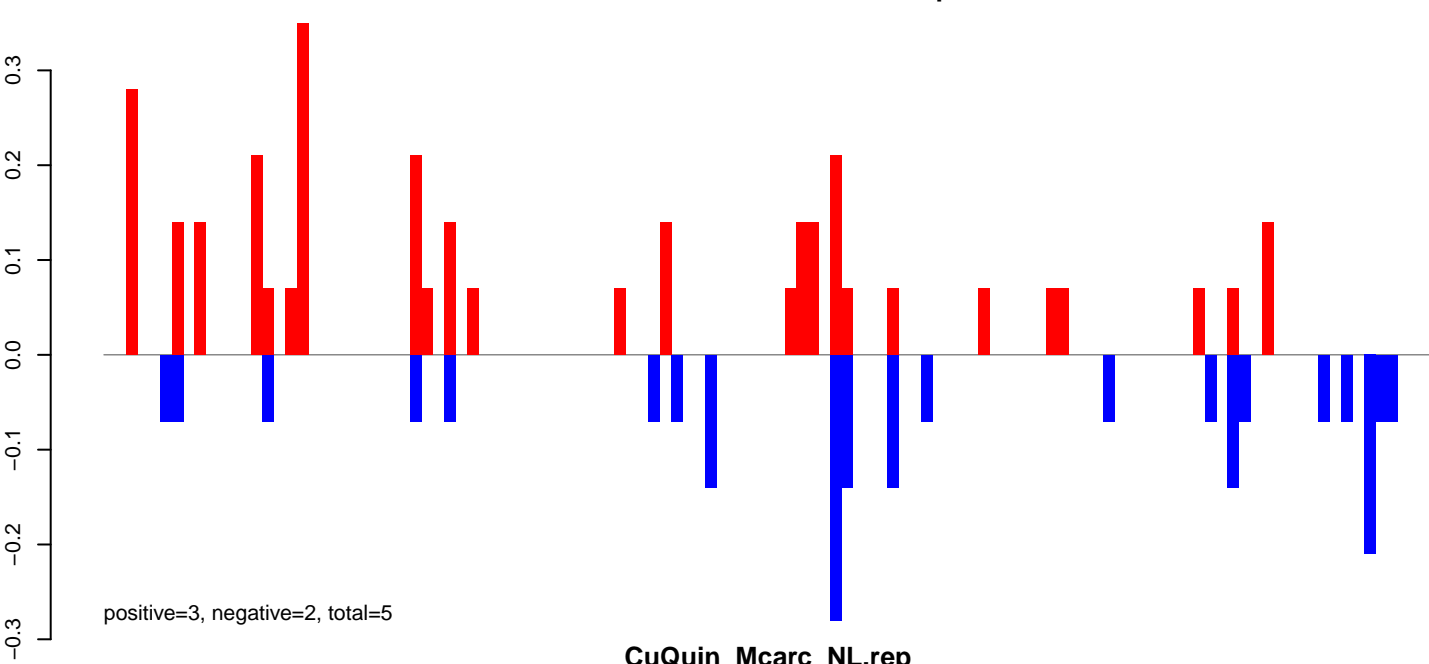
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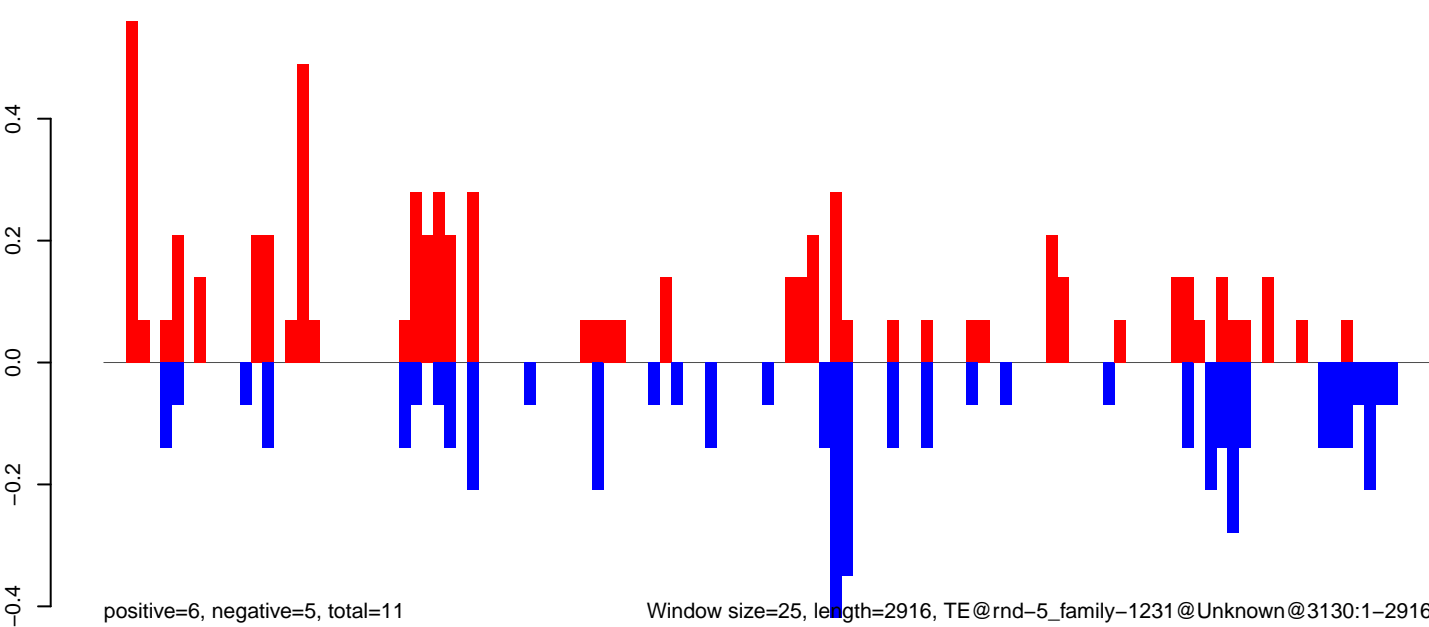
CuQuin_Mcarc_NL.18_23.rep



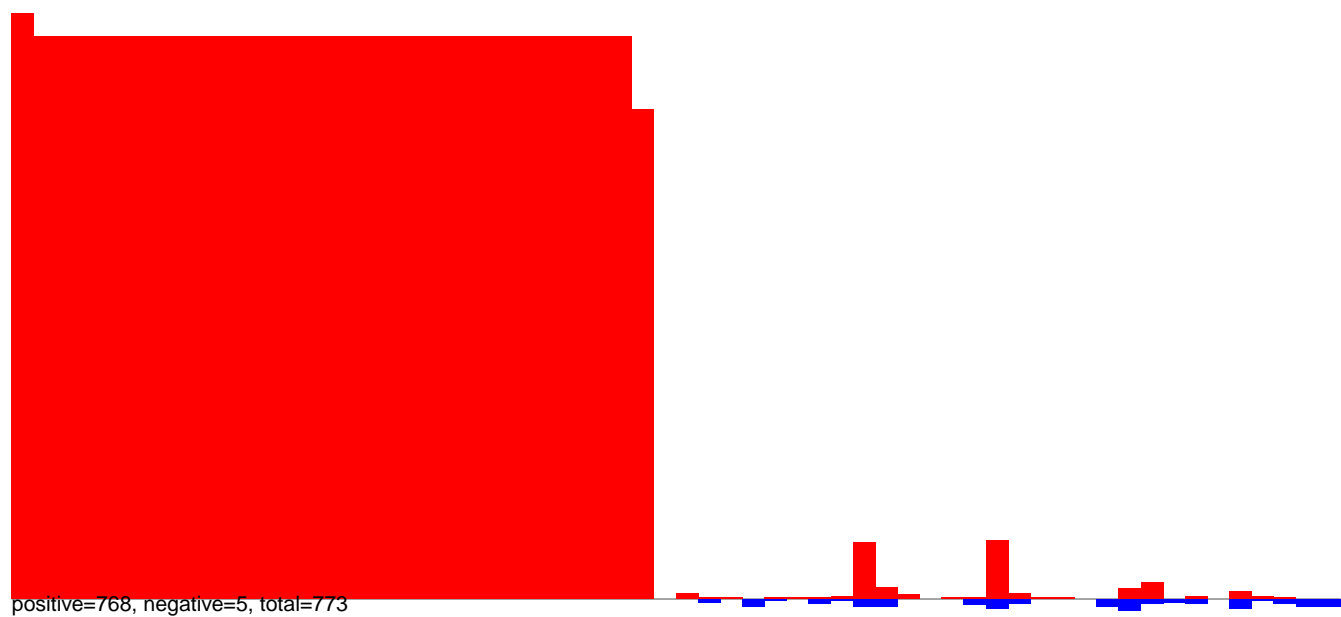
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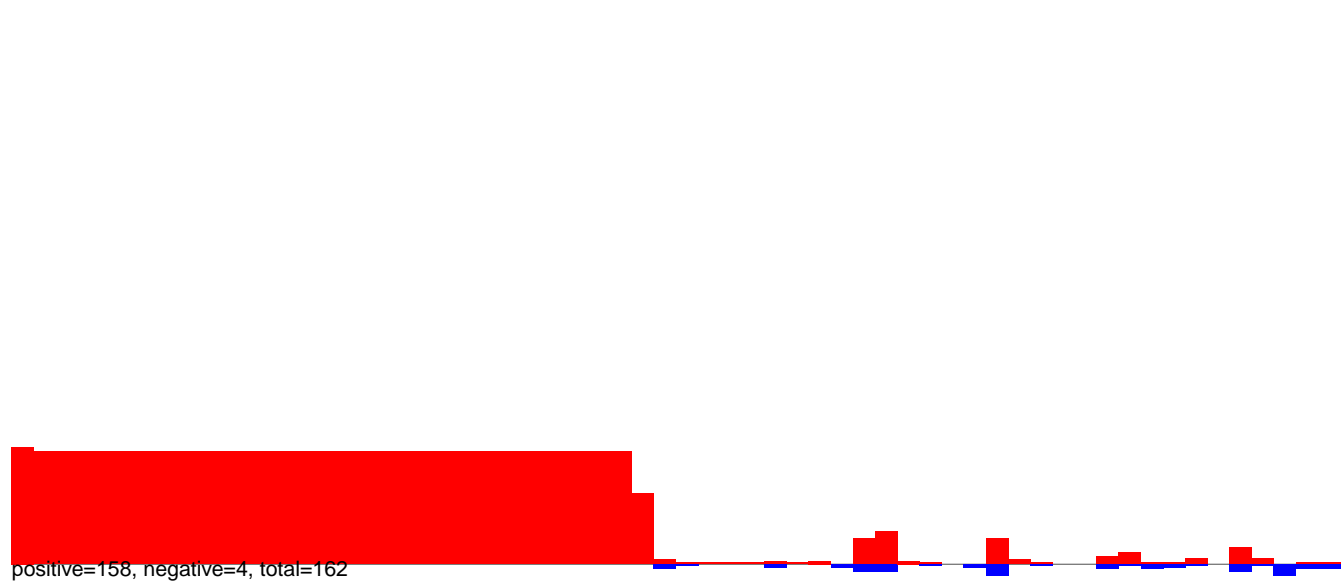
CuQuin_Mcarc_NL.rep



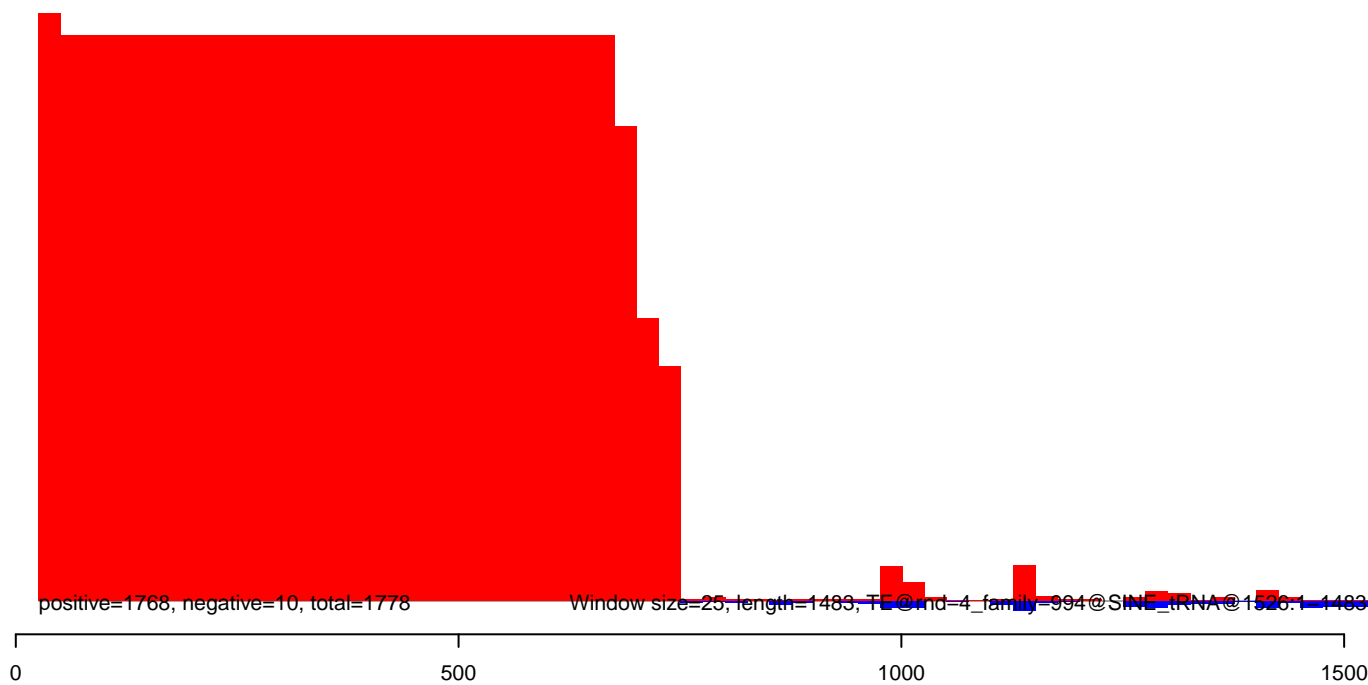
CuQuin_Mcarc_NL.18_23.rep



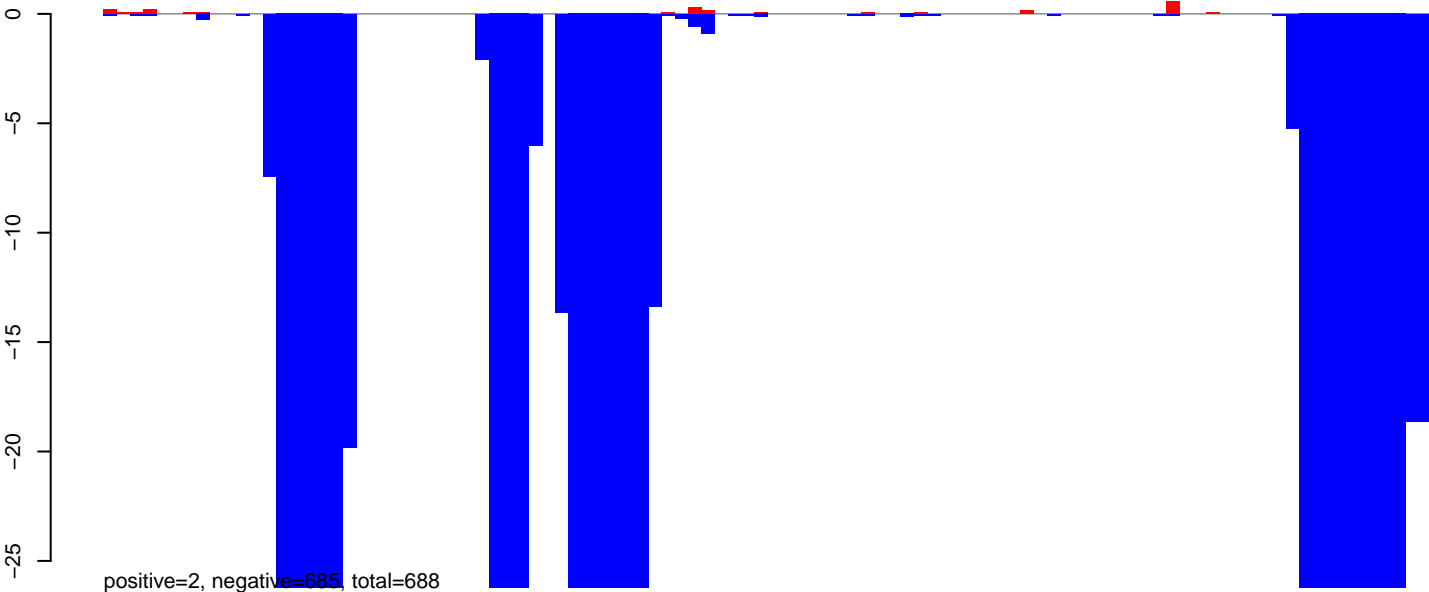
CuQuin_Mcarc_NL.24_35.rep



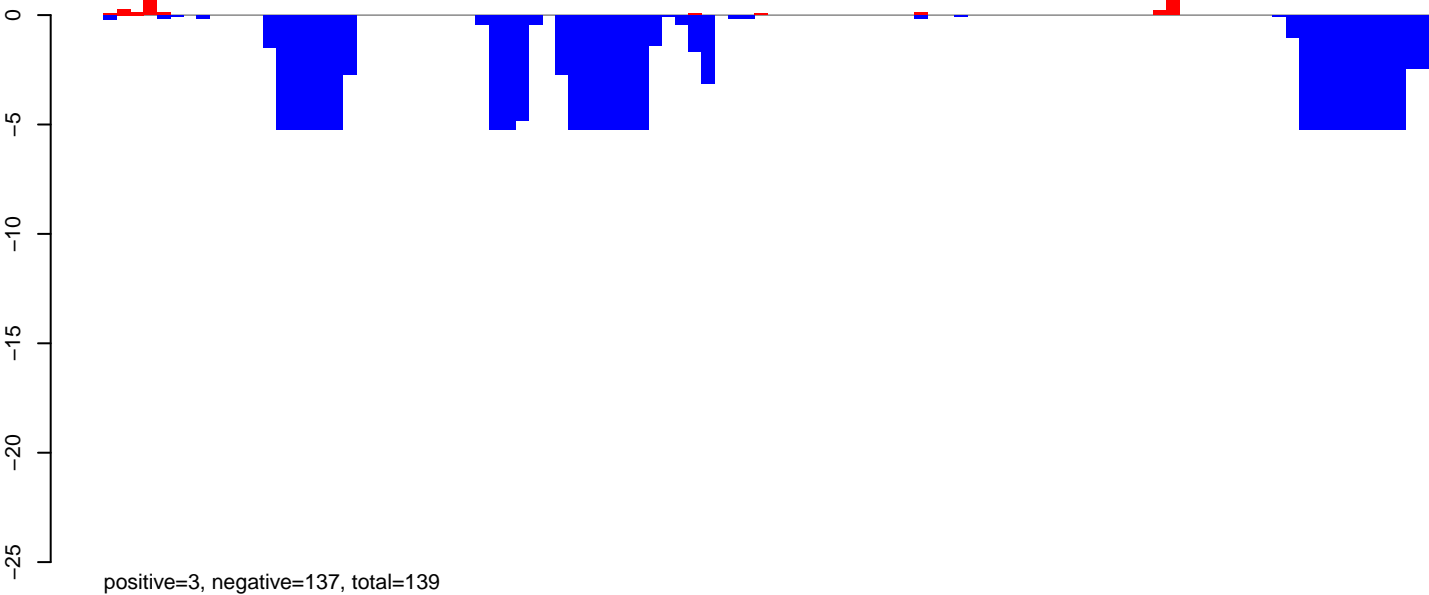
CuQuin_Mcarc_NL.rep



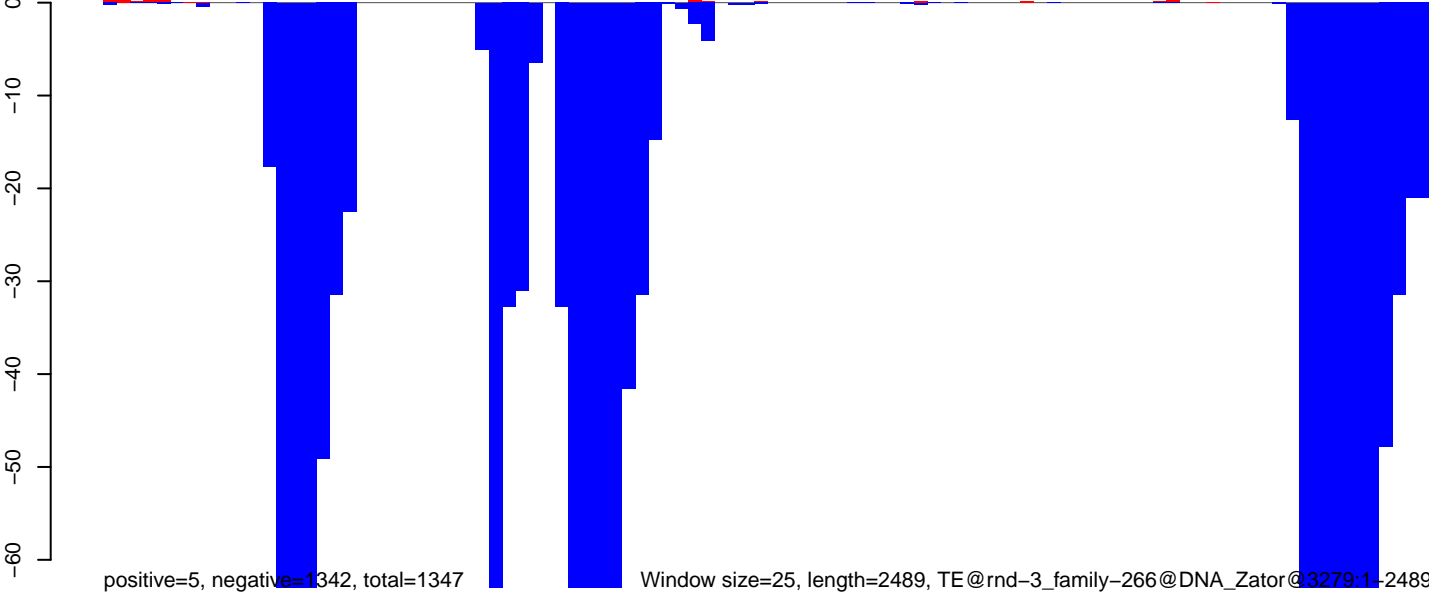
CuQuin_Mcarc_NL.18_23.rep



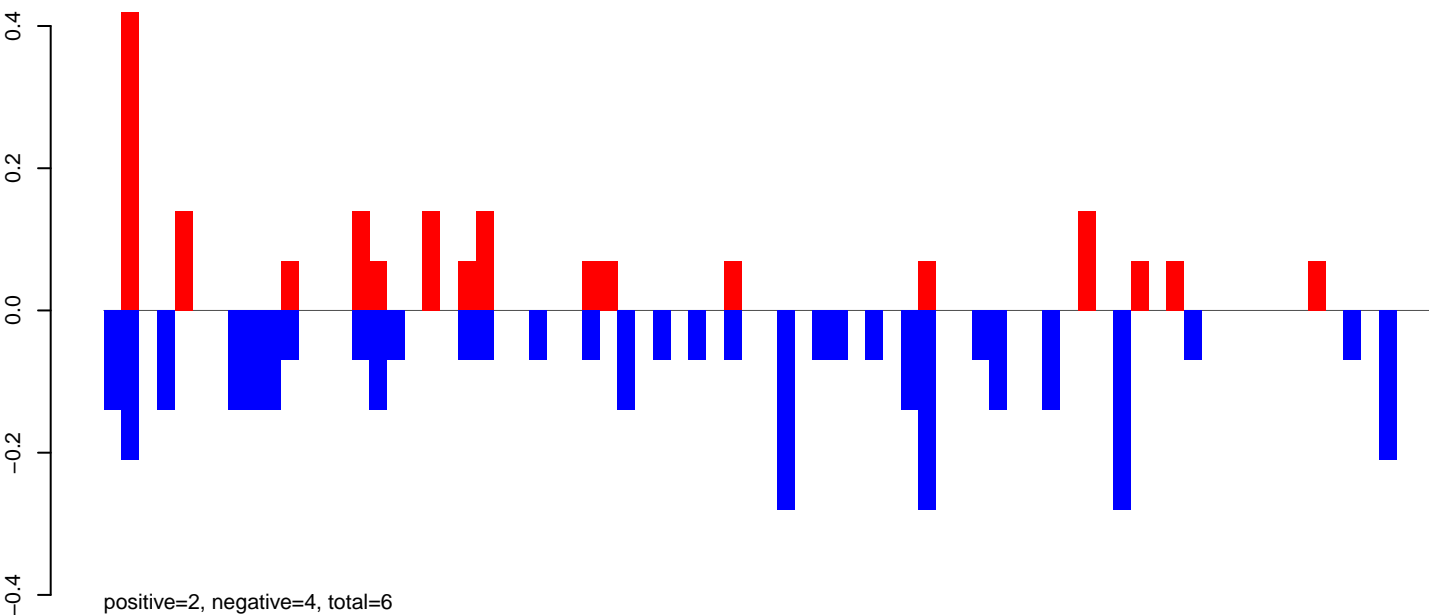
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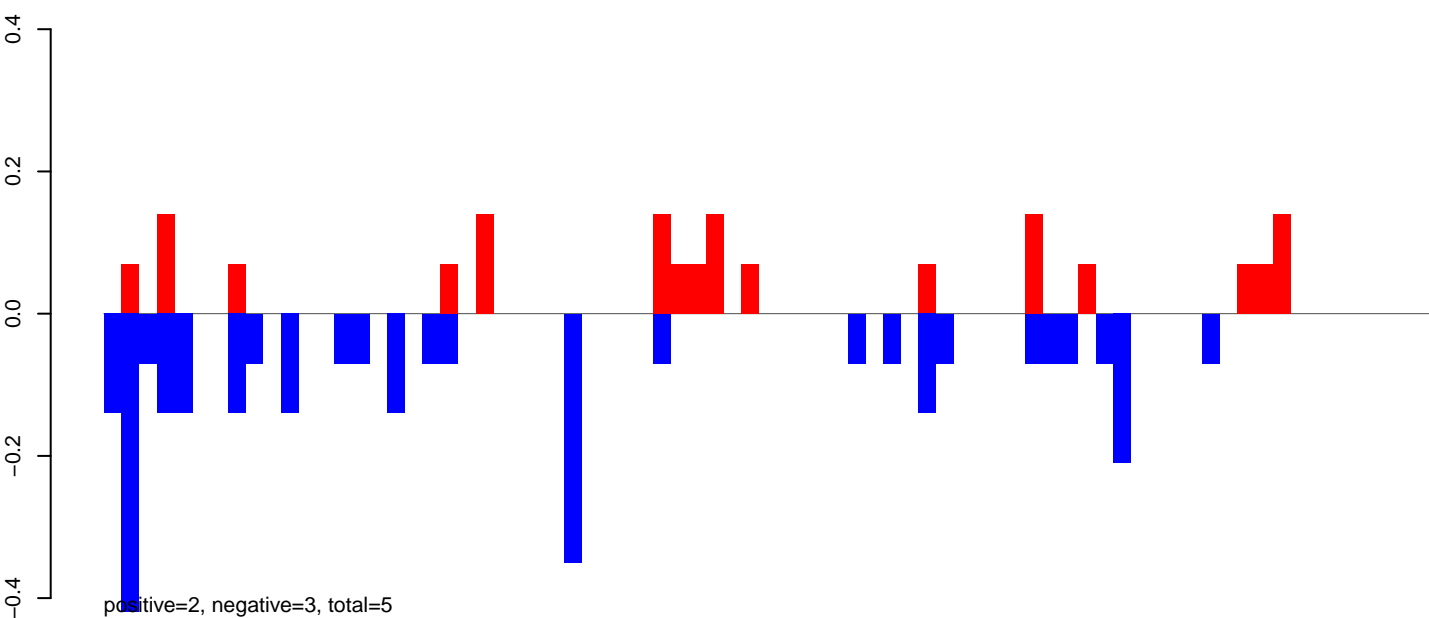
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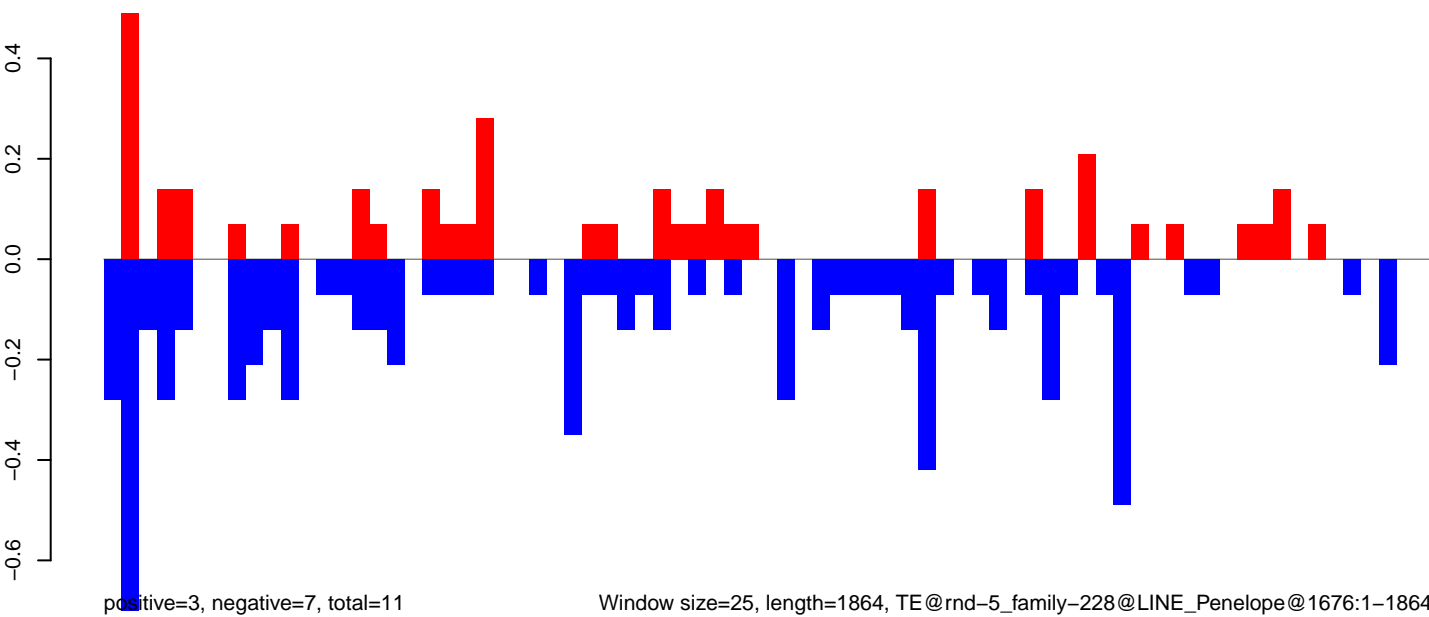
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CuQuin_Mcarc_NL.24_35.rep



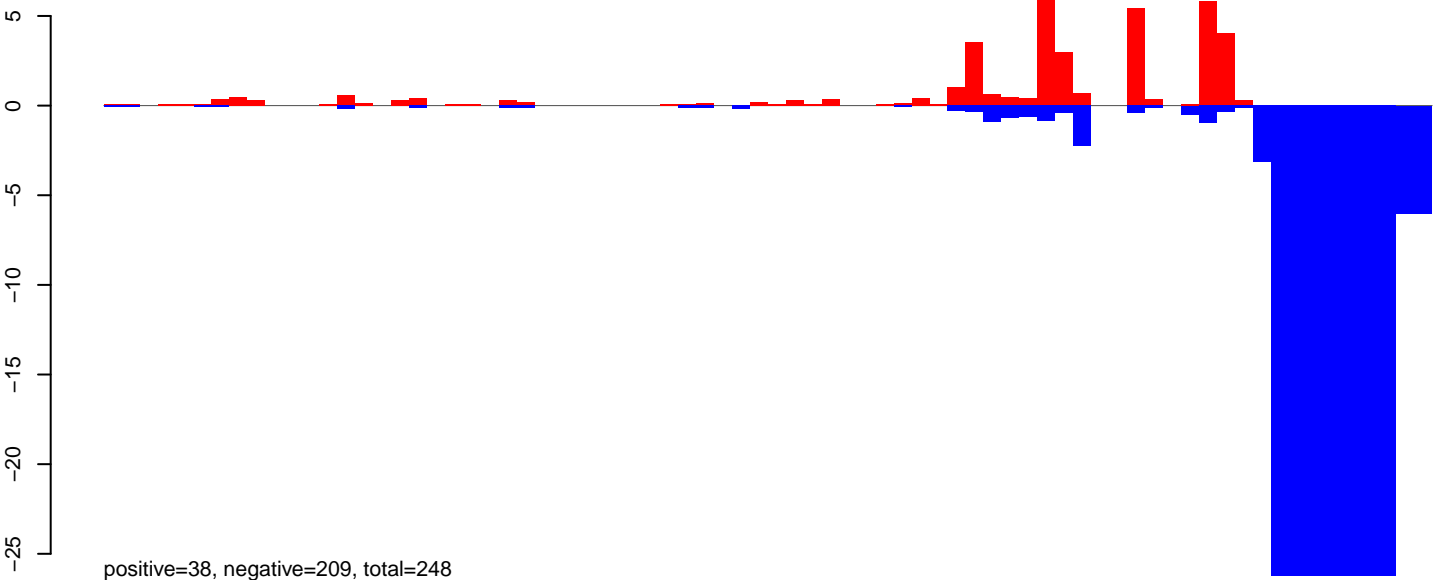
CuQuin_Mcarc_NL.rep



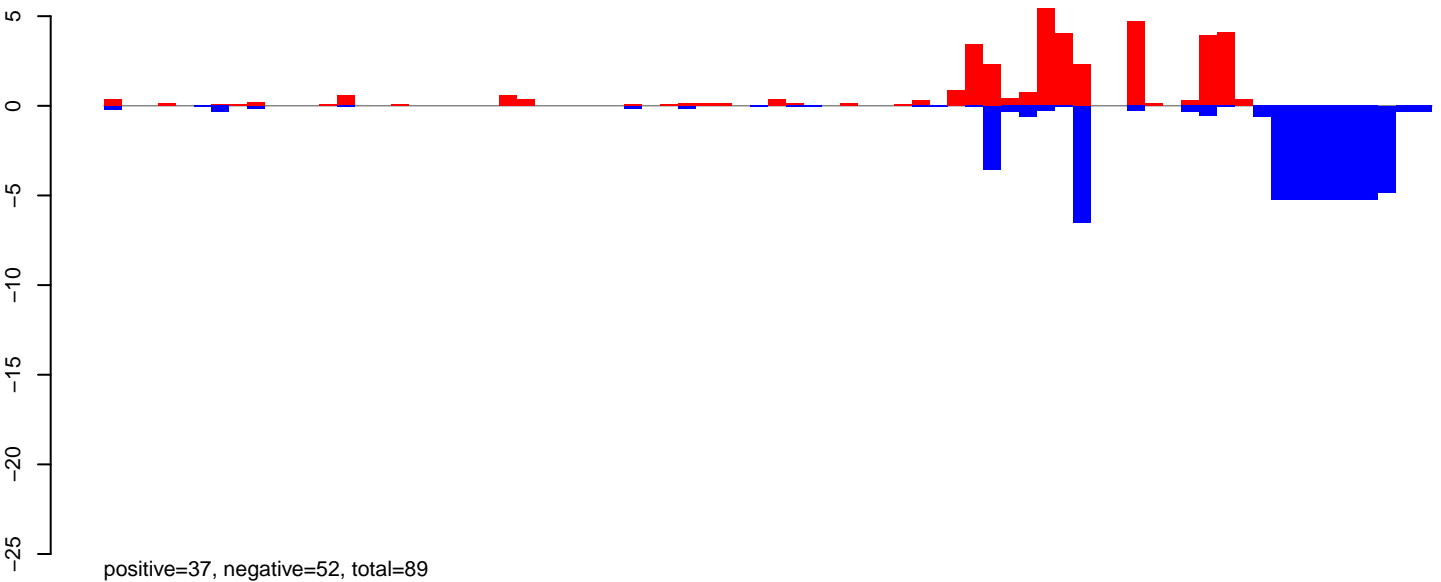
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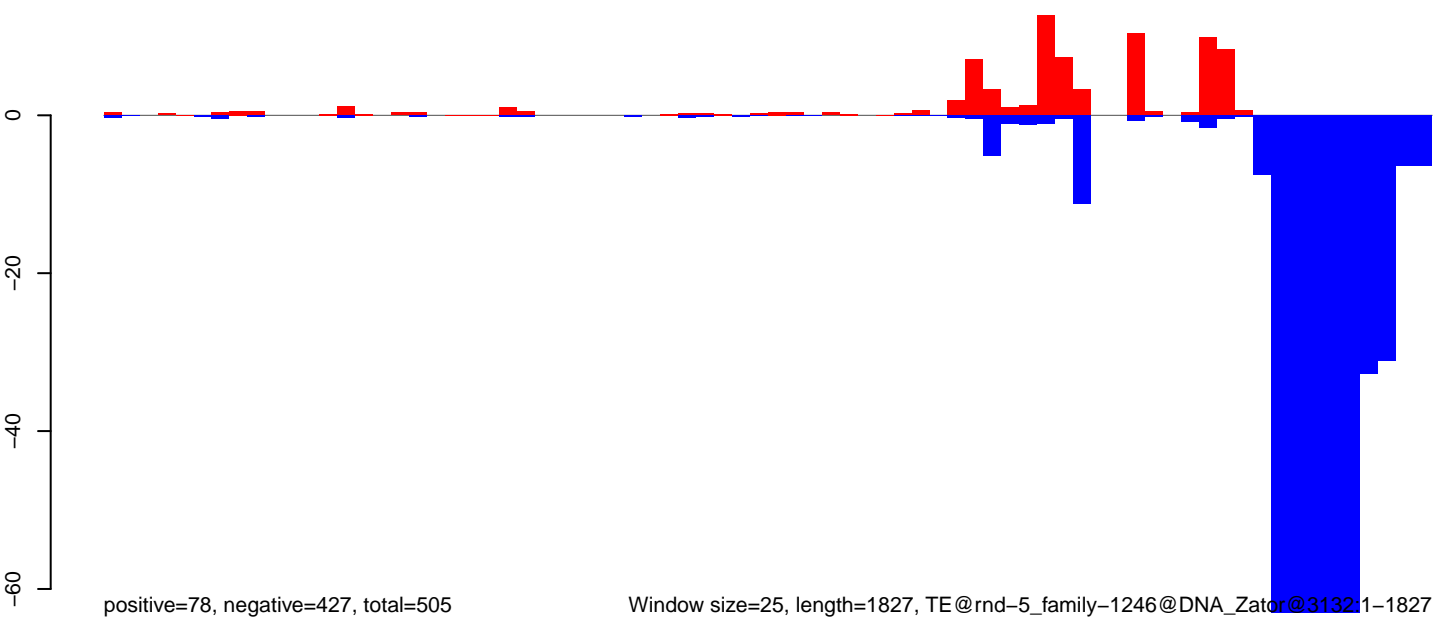
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



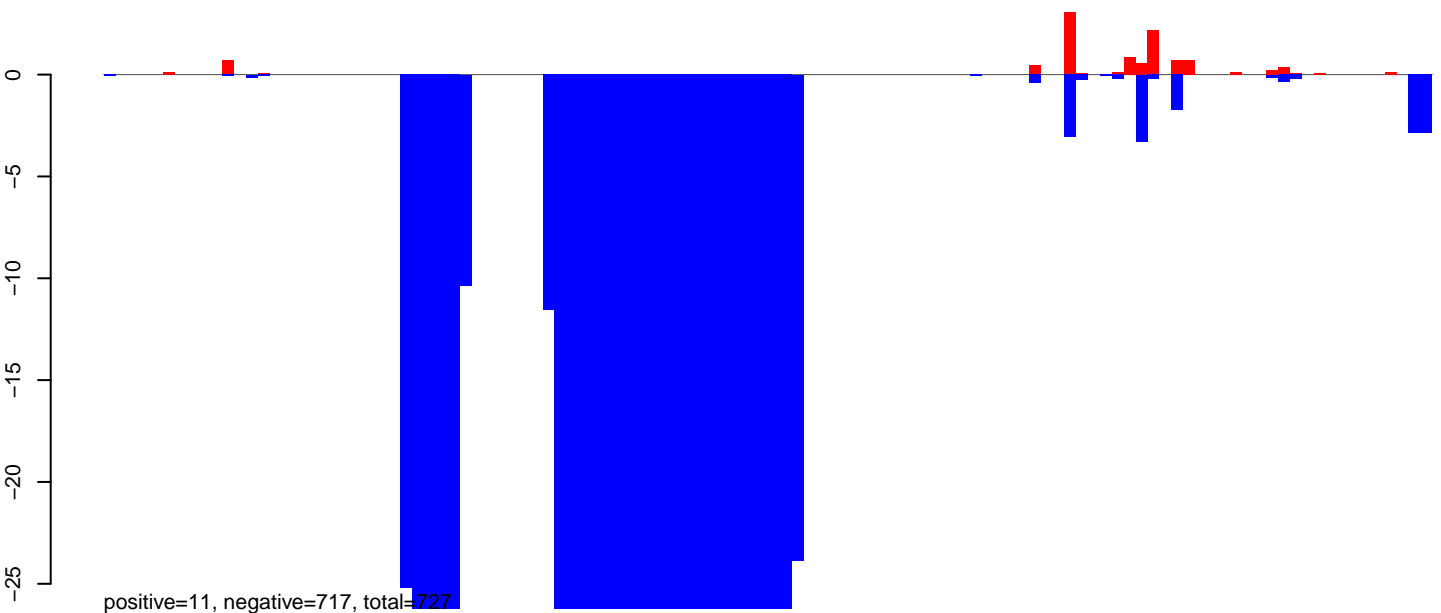
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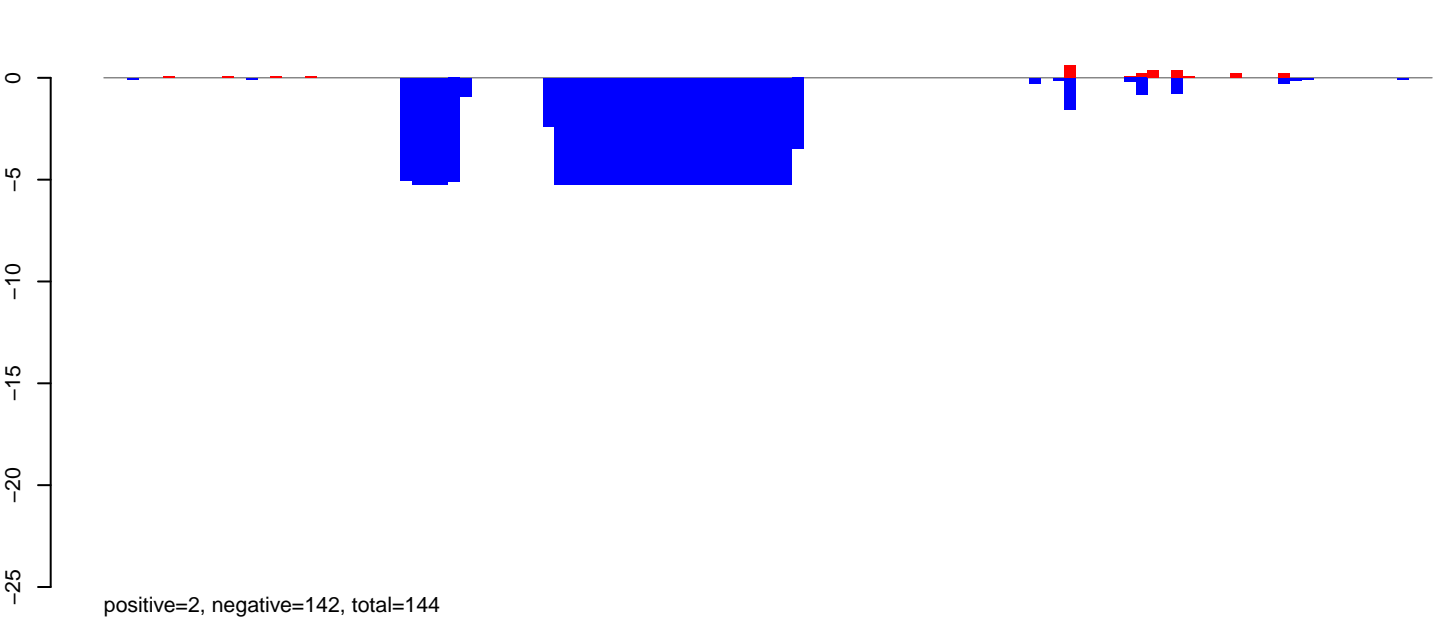
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0 500 1000 1500

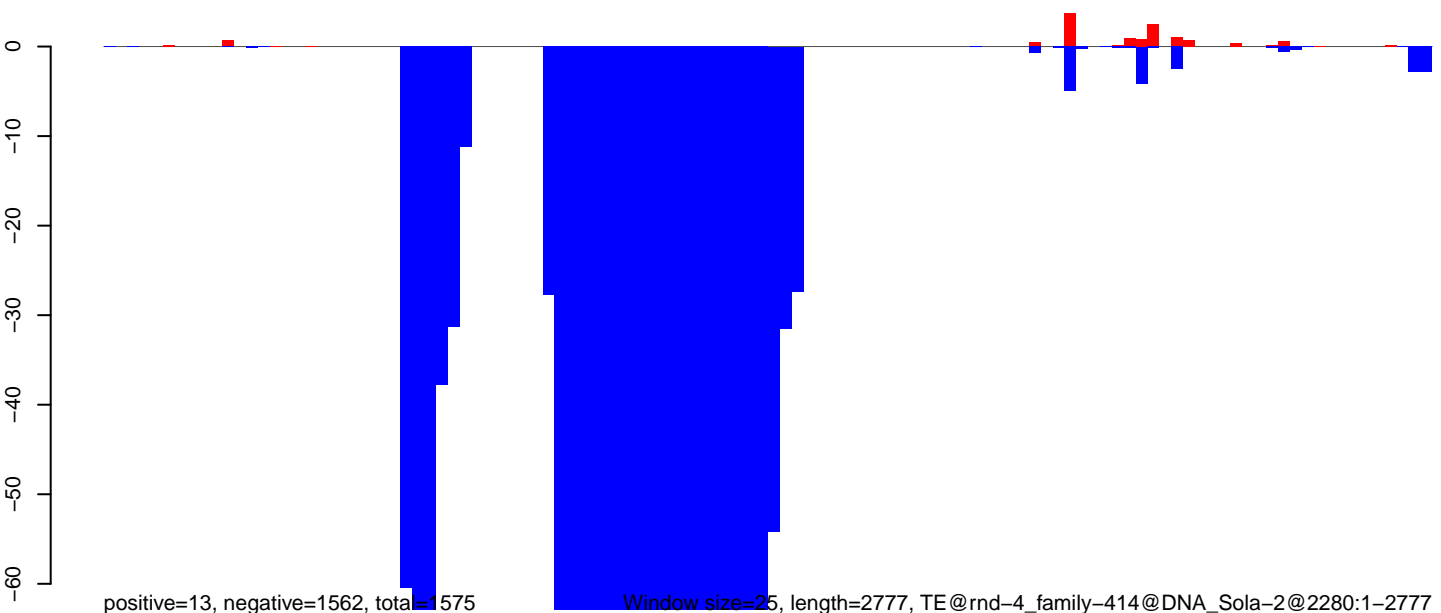
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CuQuin_Mcarc_NL.24_35.rep

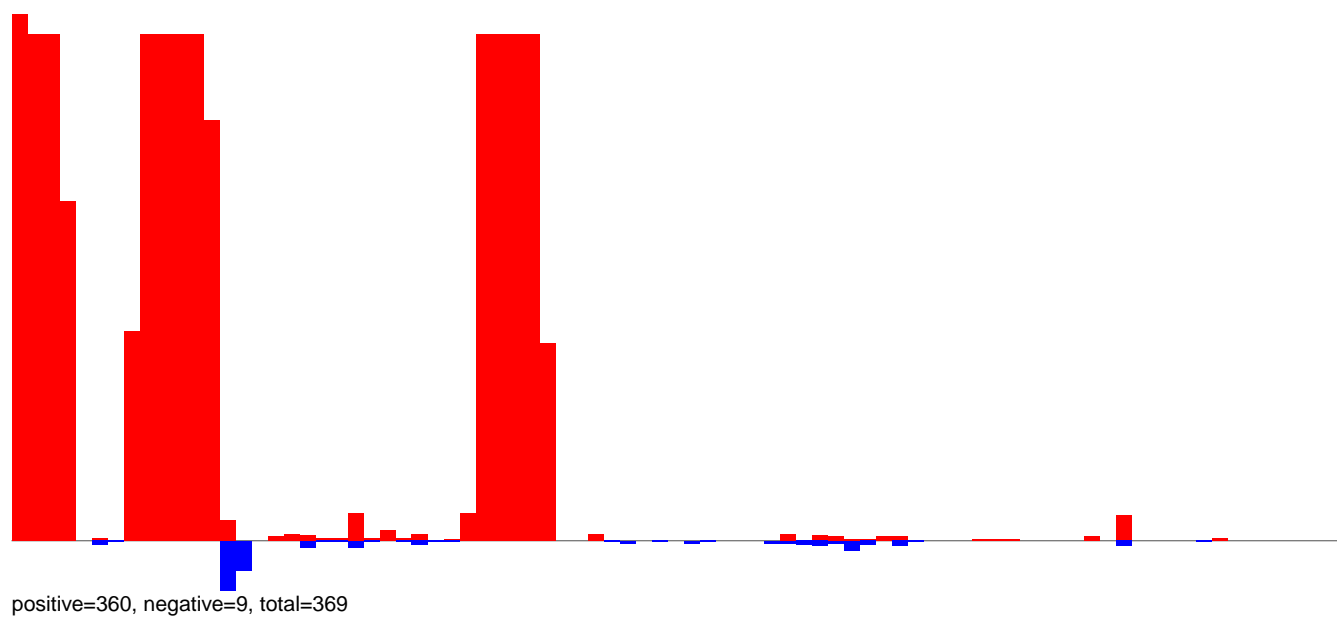


CuQuin_Mcarc_NL.rep

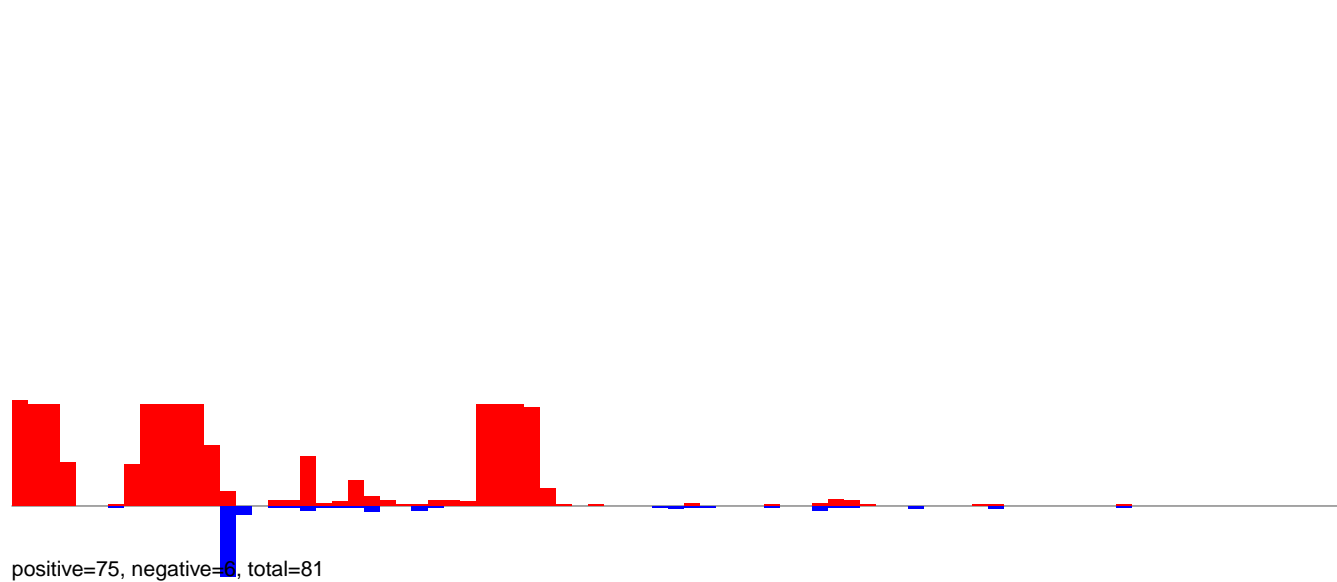


Window size=25, length=2777, TE@rnd-4_family-414@DNA_Sola-2@2280:1-2777

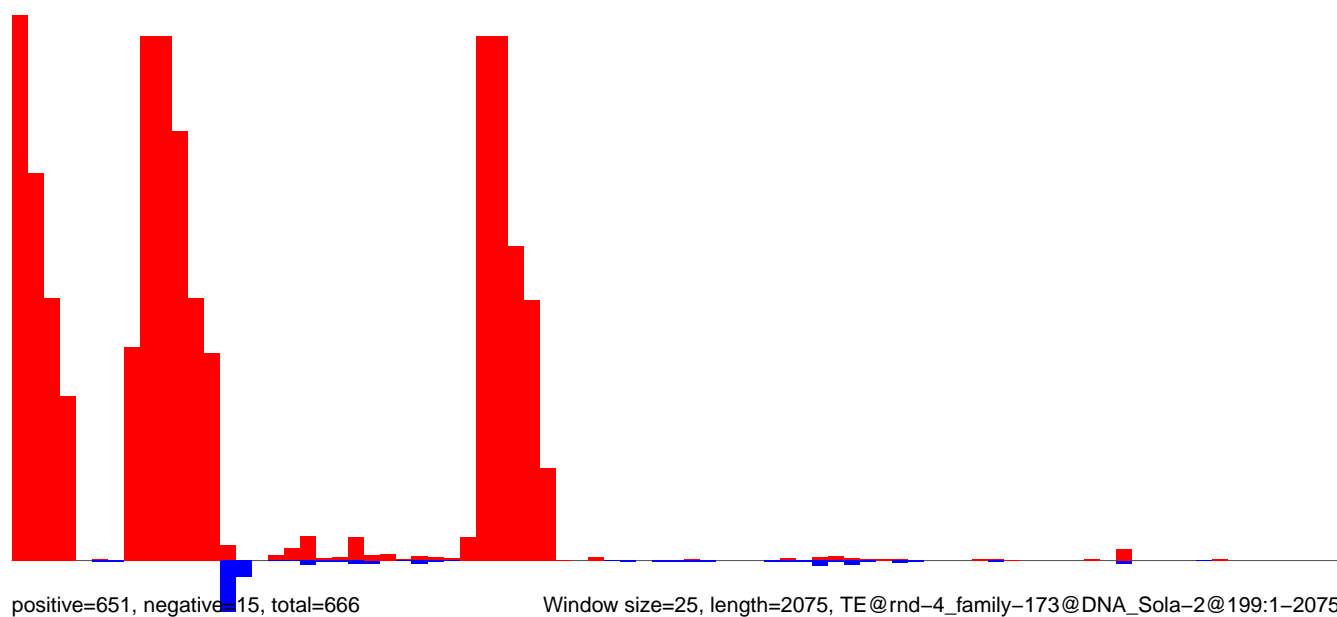
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CuQuin_Mcarc_NL.24_35.rep

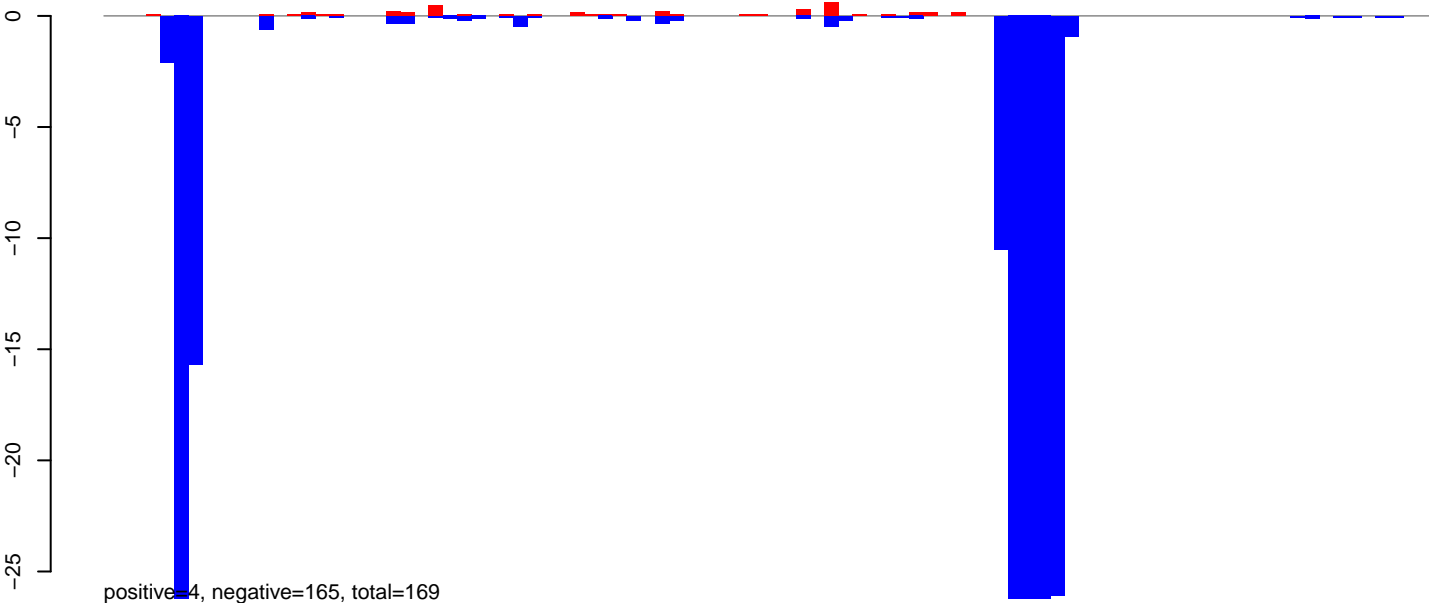


CuQuin_Mcarc_NL.rep

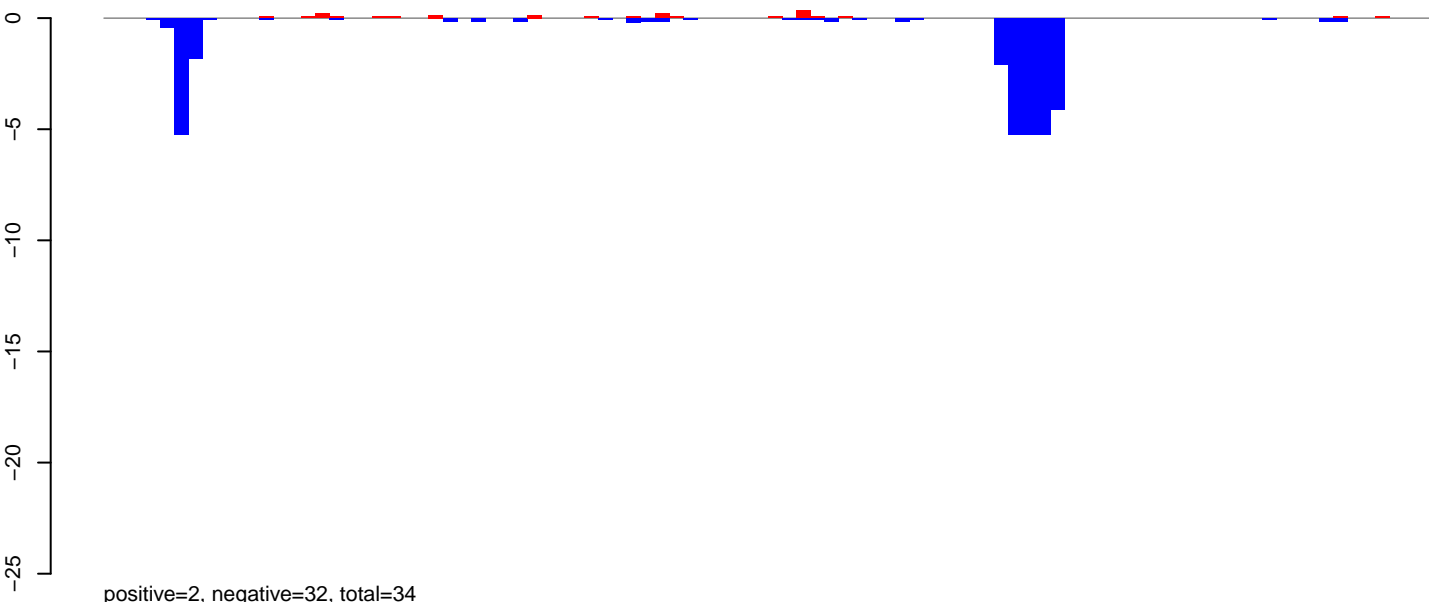


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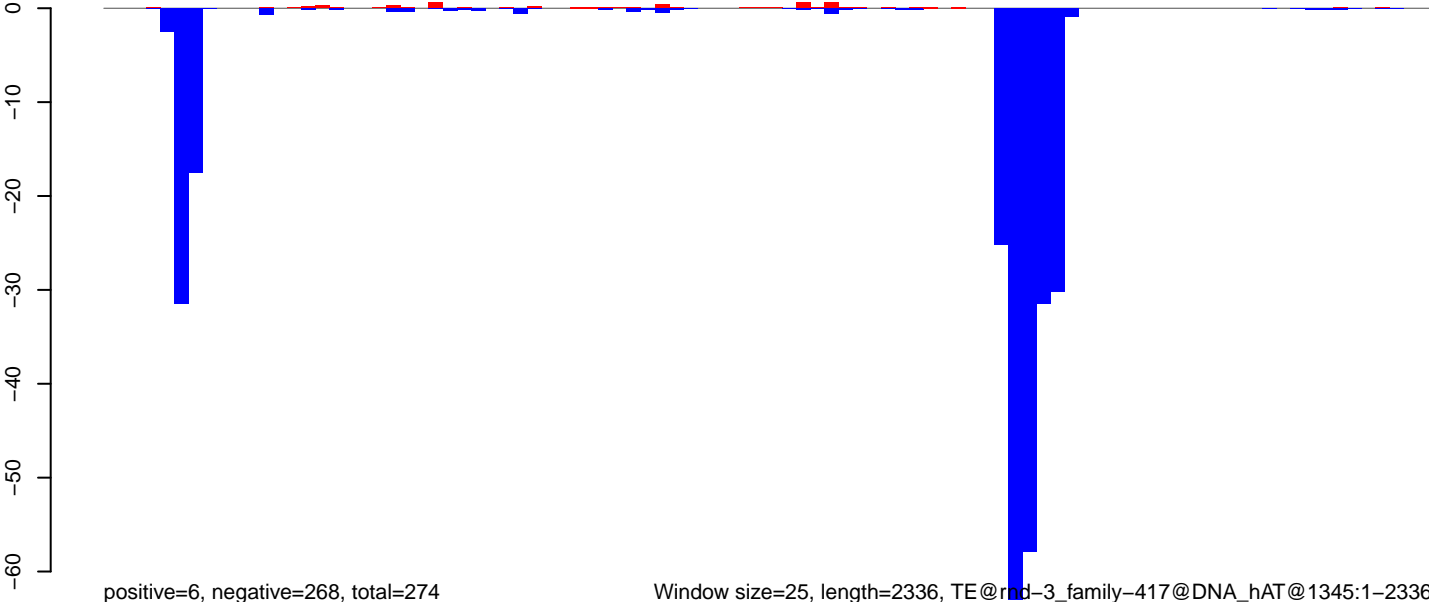
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



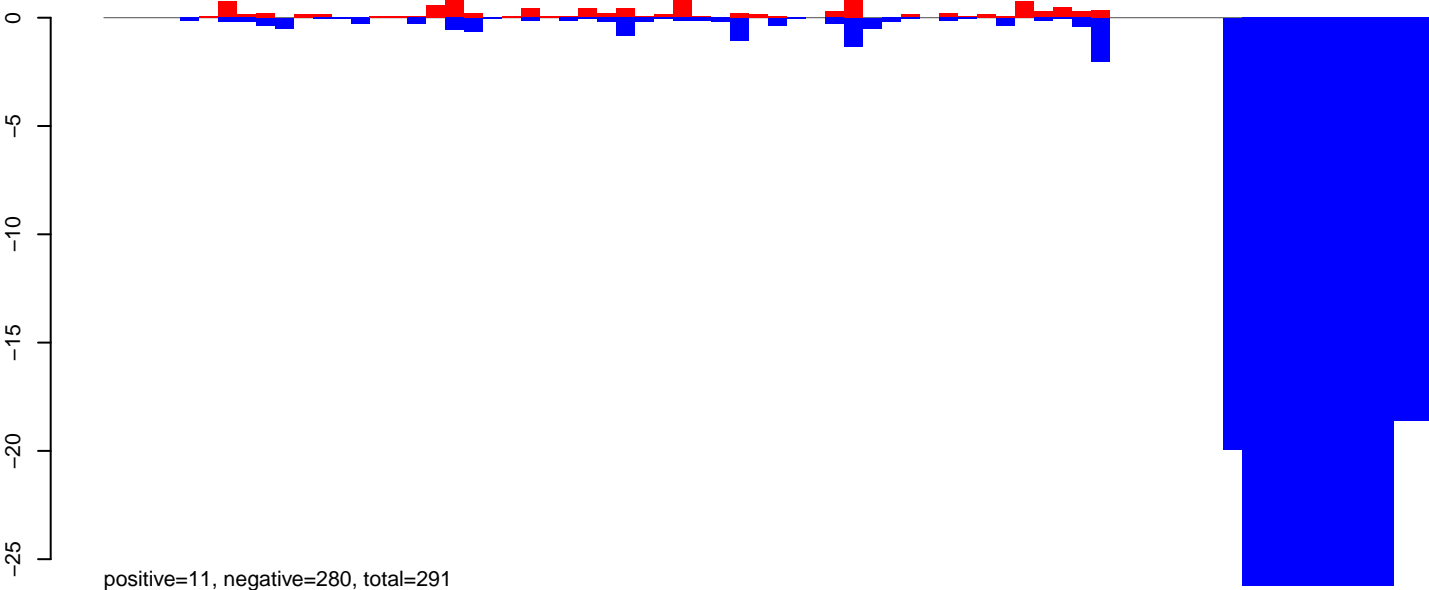
CuQuin_Mcarc_NL.rep



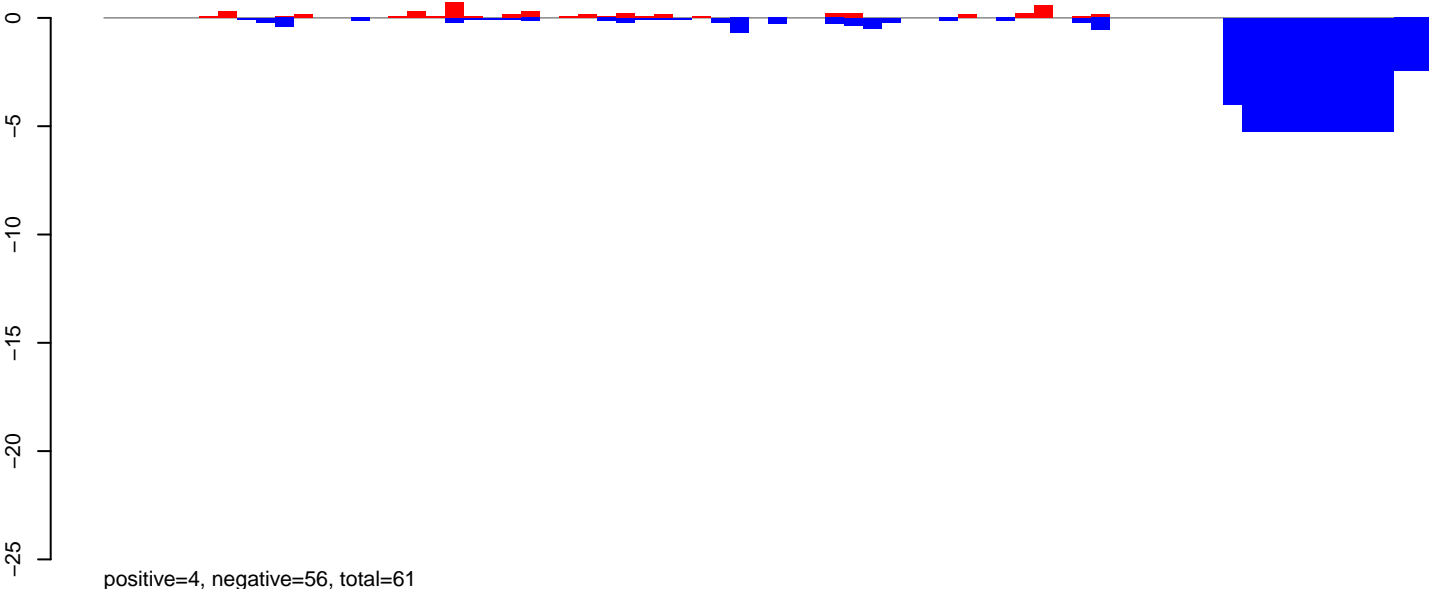
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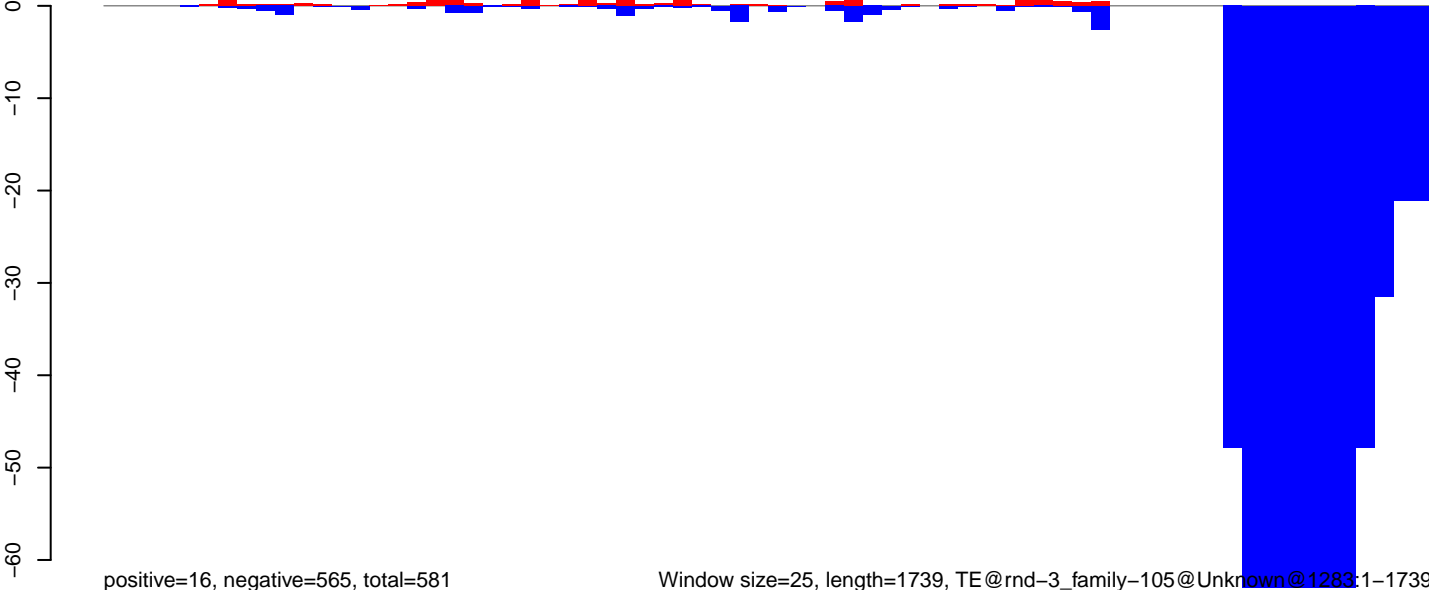
CuQuin_Mcarc_NL.18_23.rep



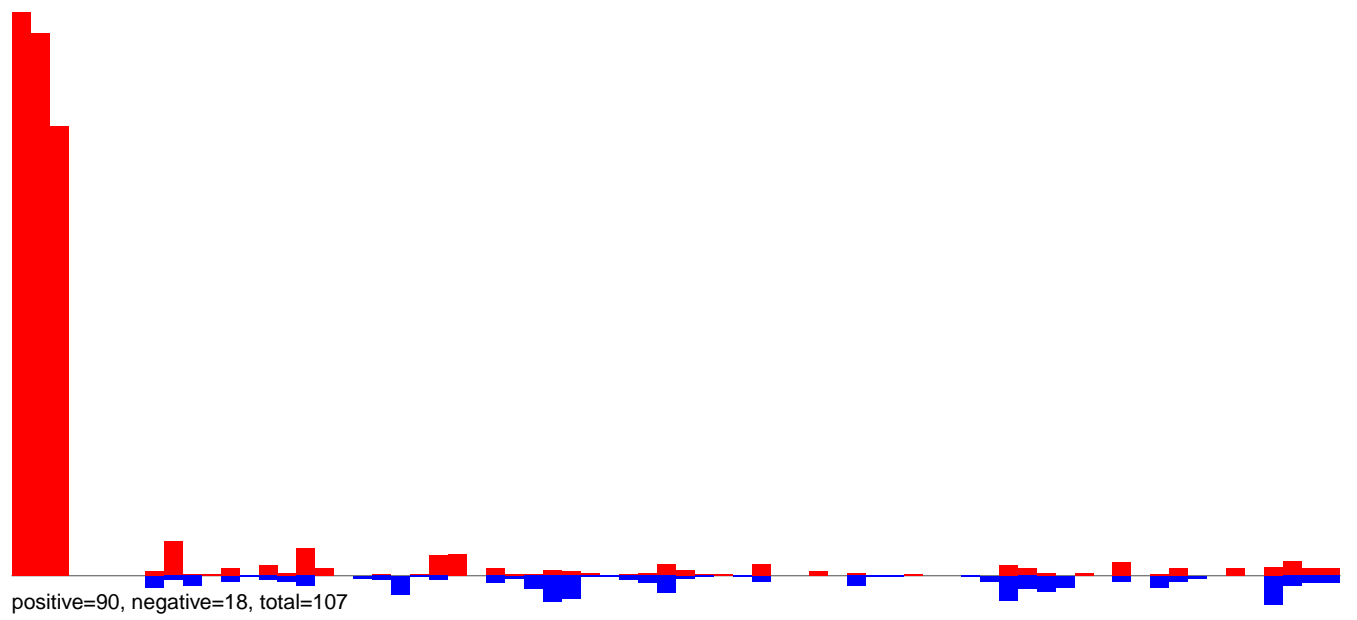
CuQuin_Mcarc_NL.24_35.rep



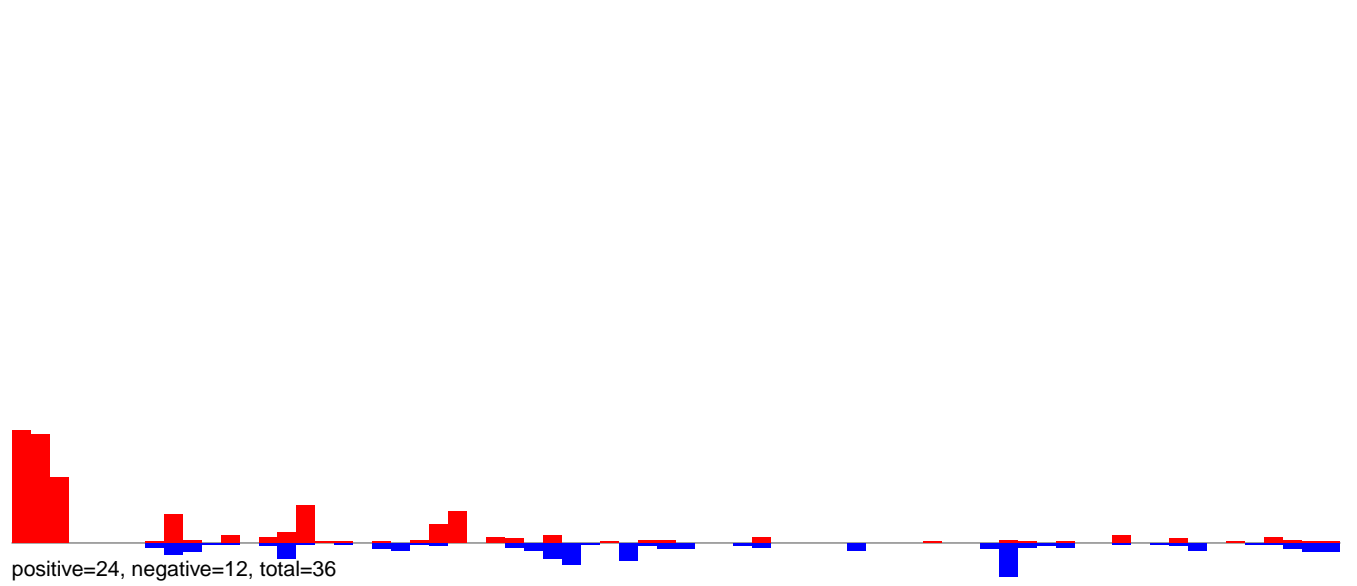
CuQuin_Mcarc_NL.rep



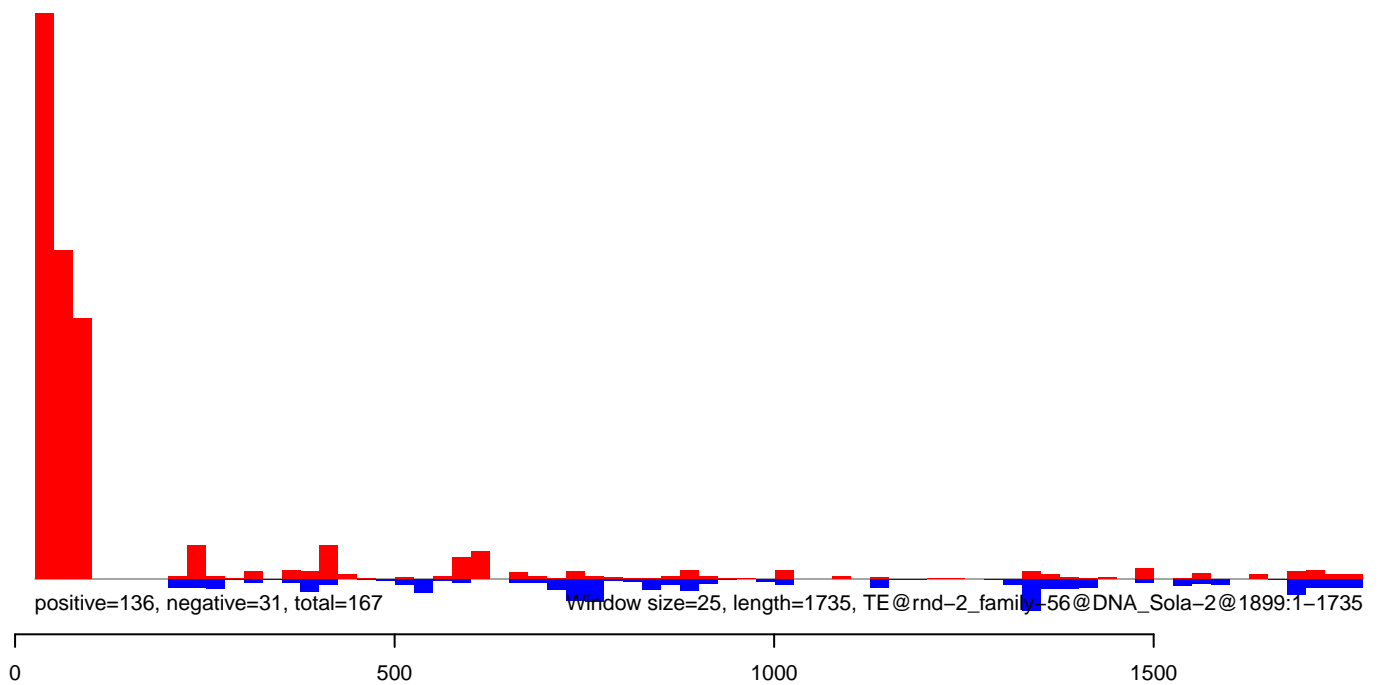
CuQuin_Mcarc_NL.18_23.rep



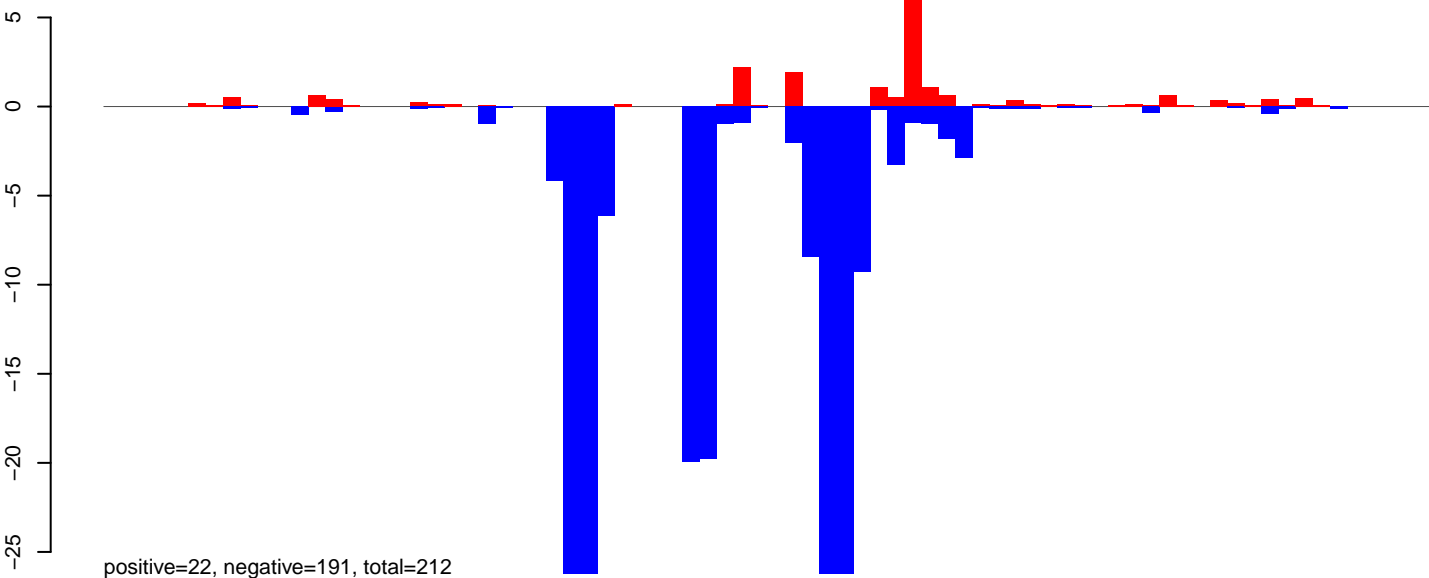
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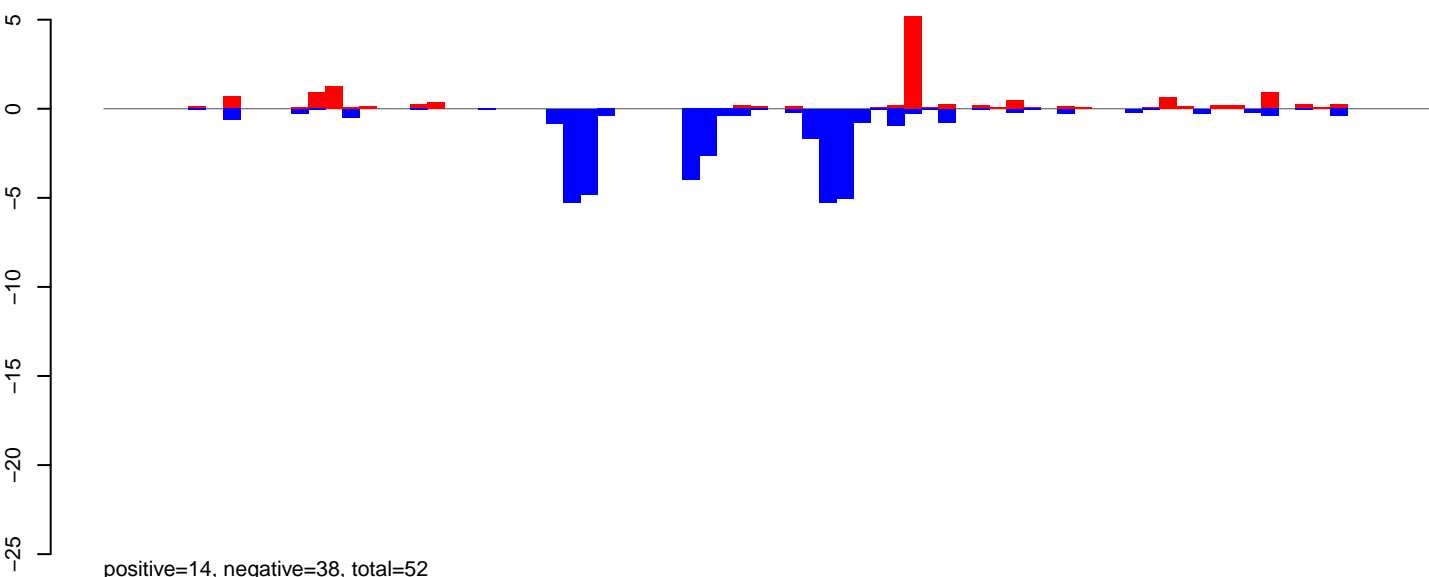
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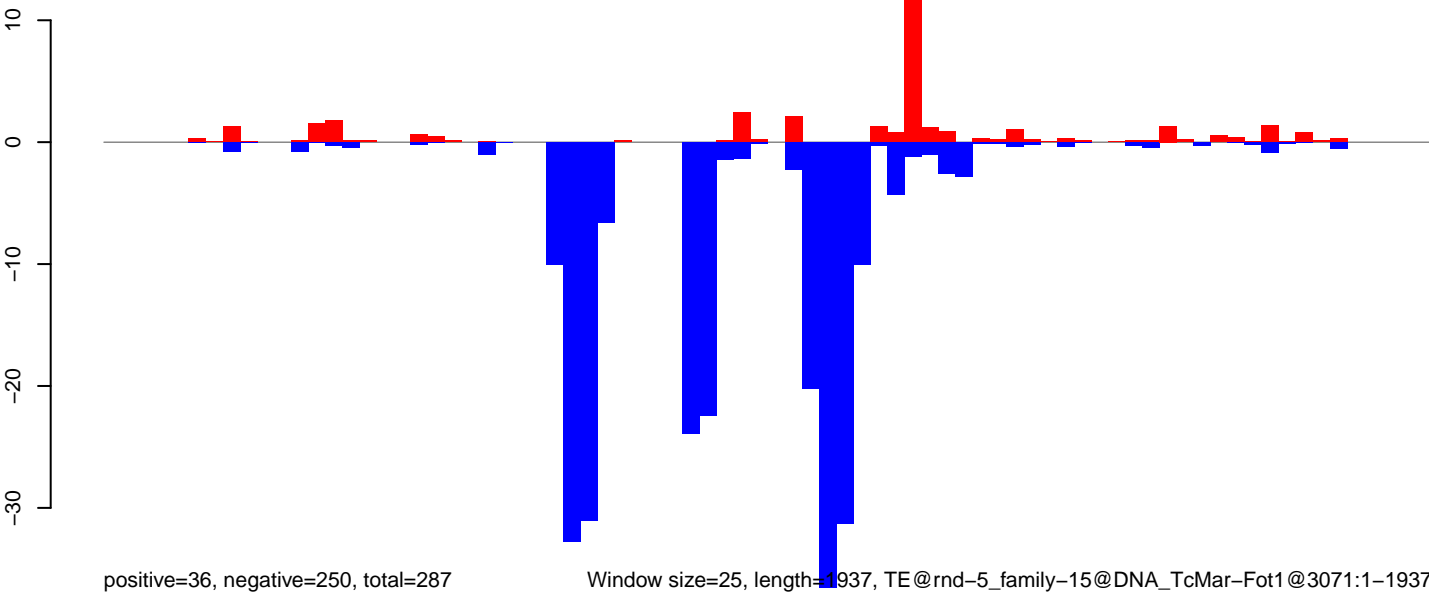
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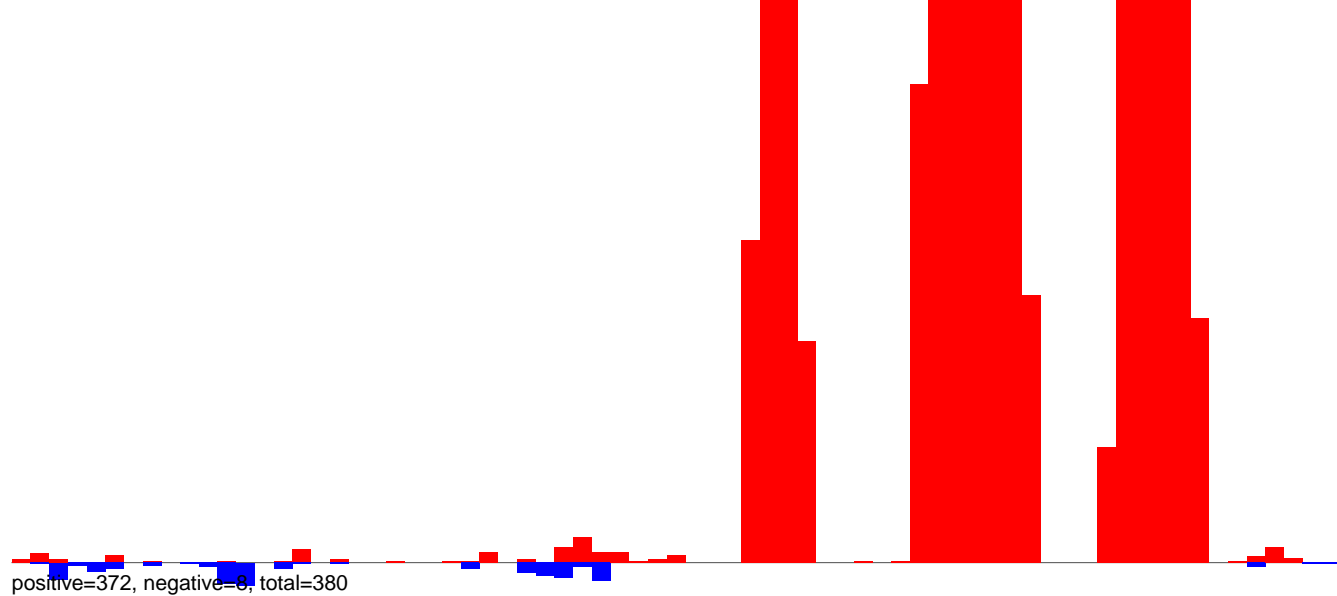
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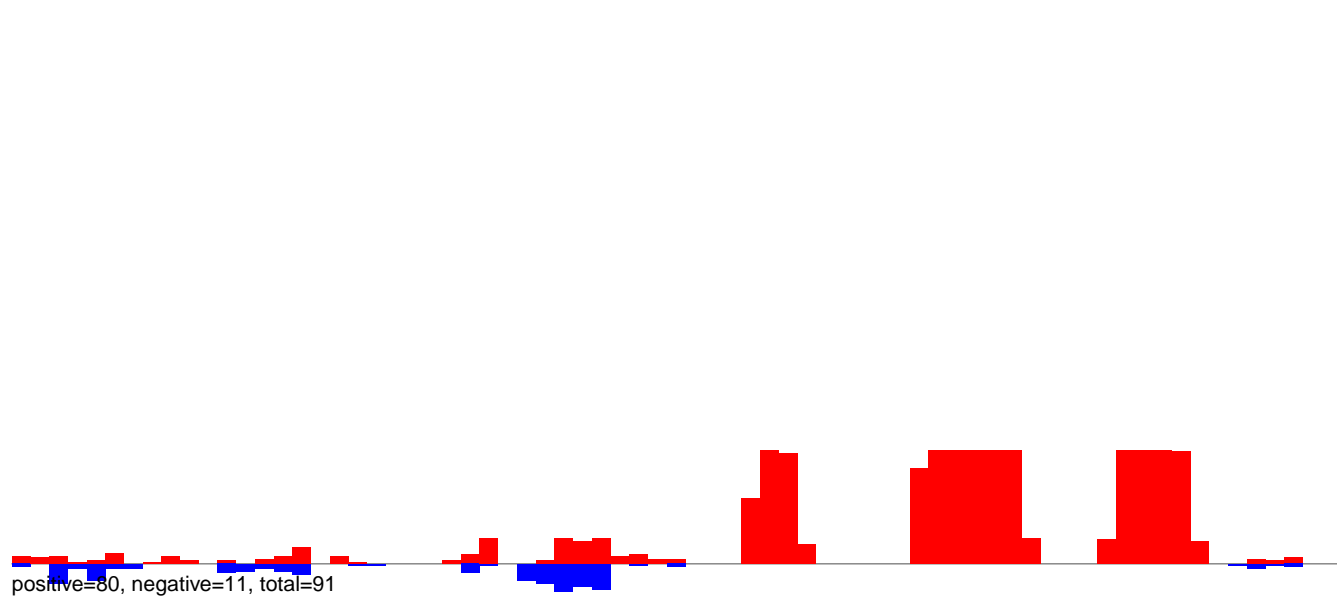
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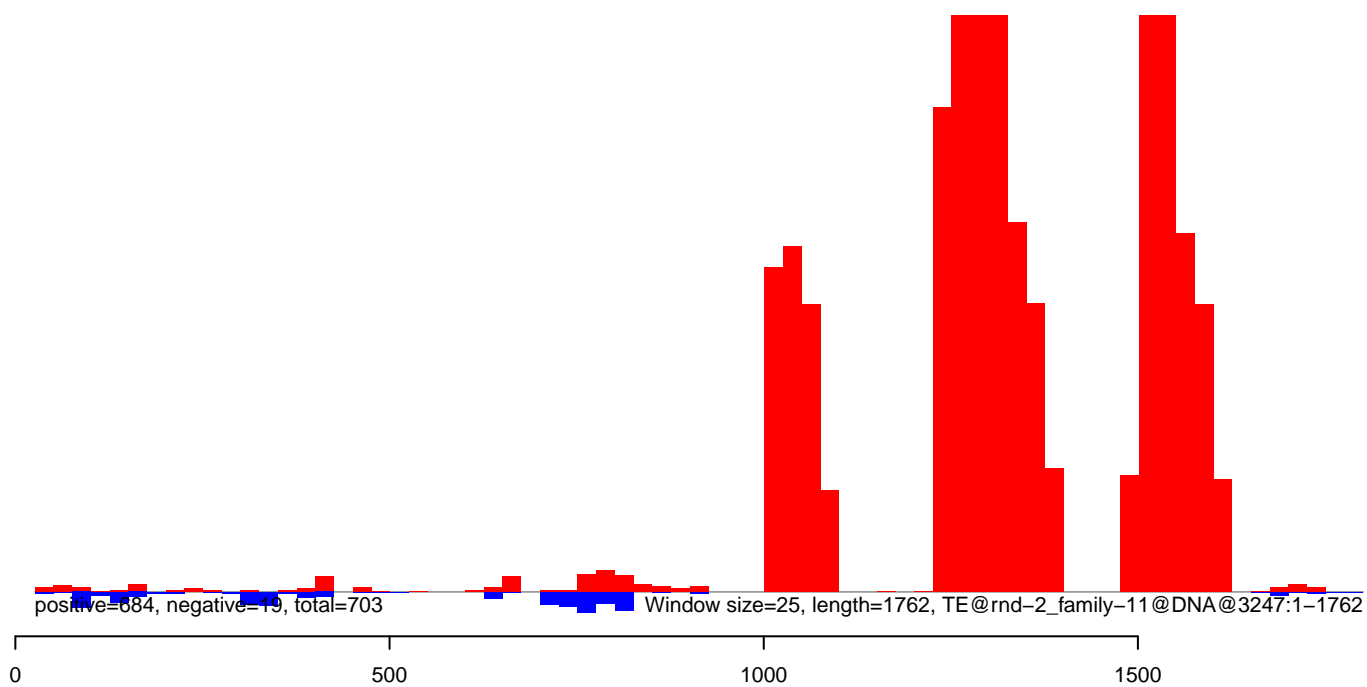
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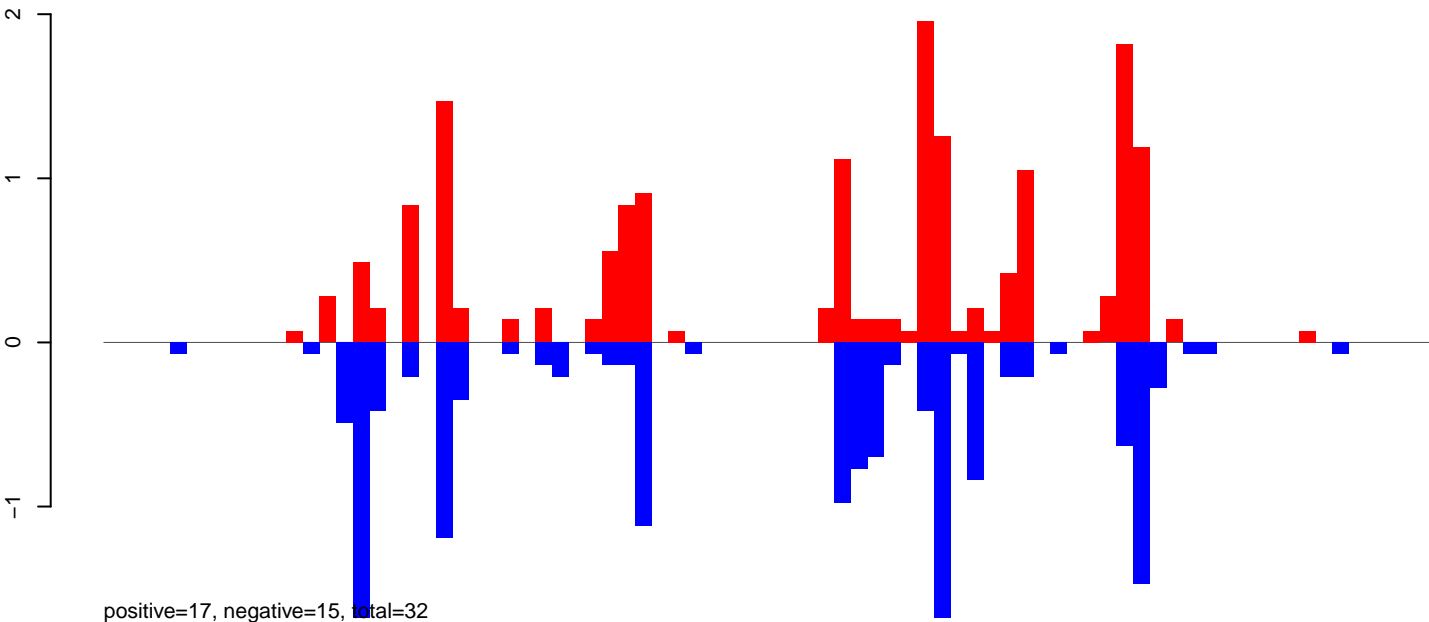
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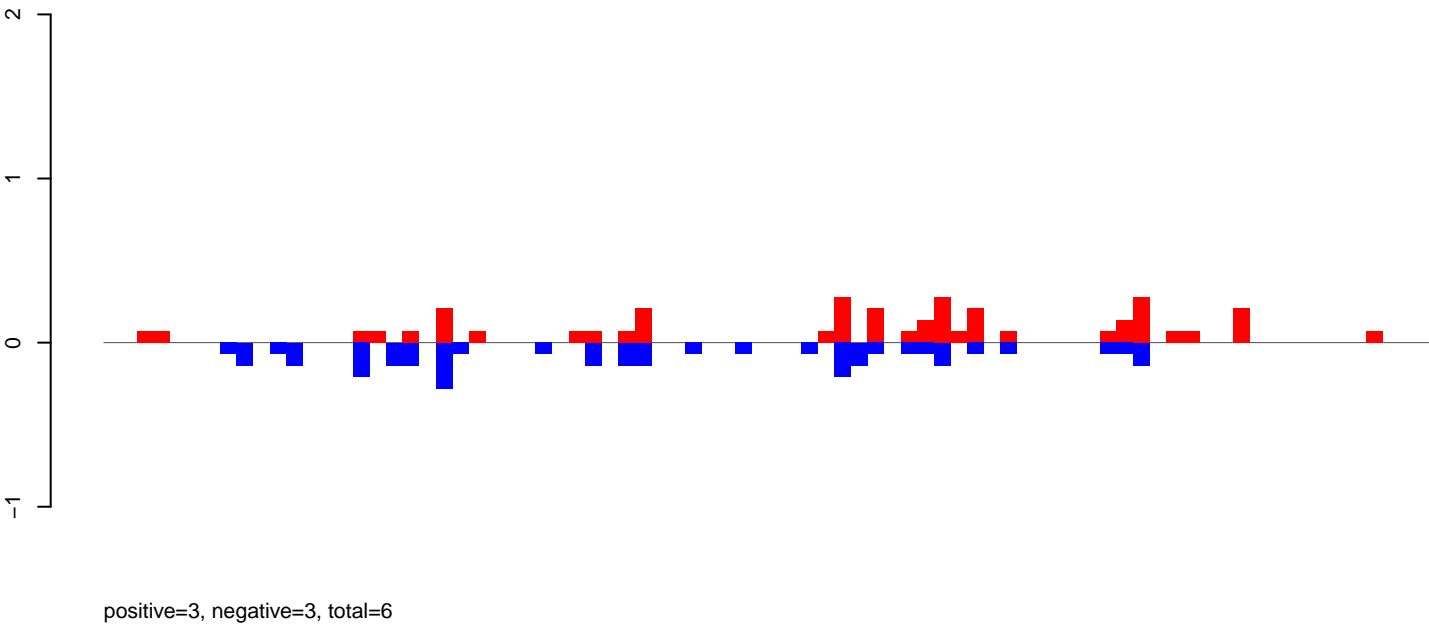
CuQuin_Mcarc_NL.rep



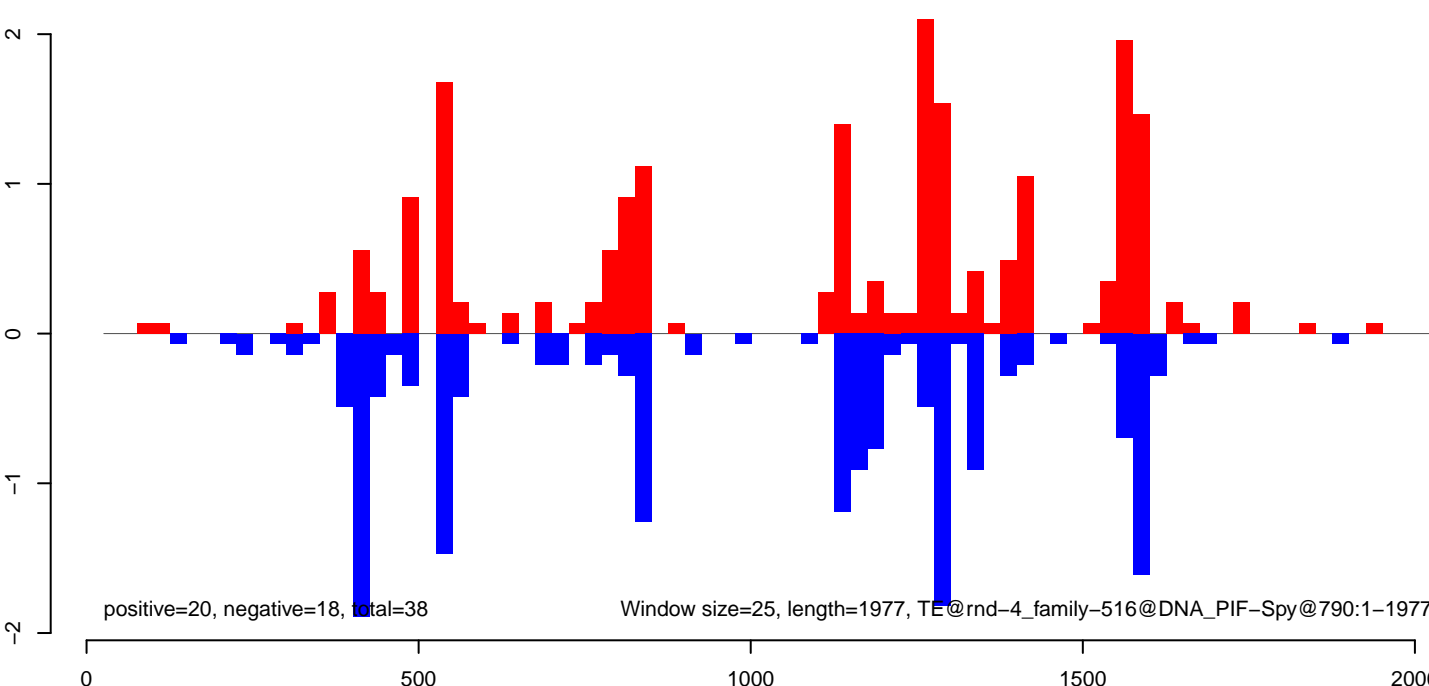
CuQuin_Mcarc_NL.18_23.rep



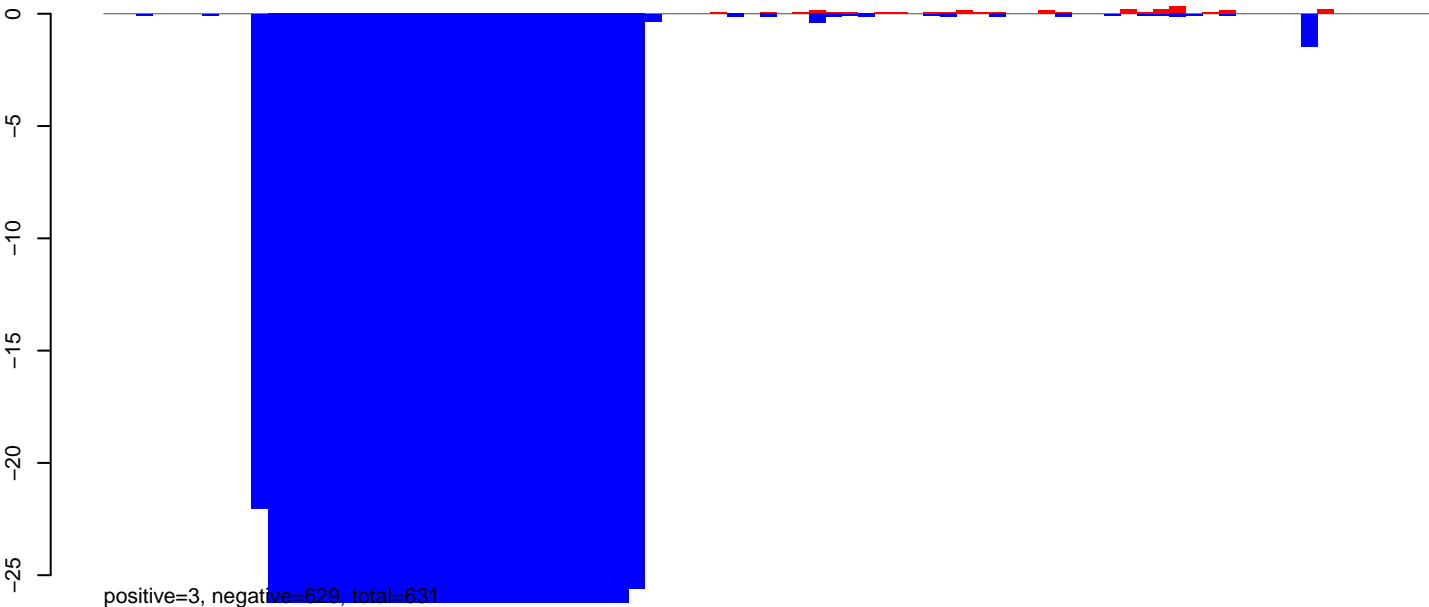
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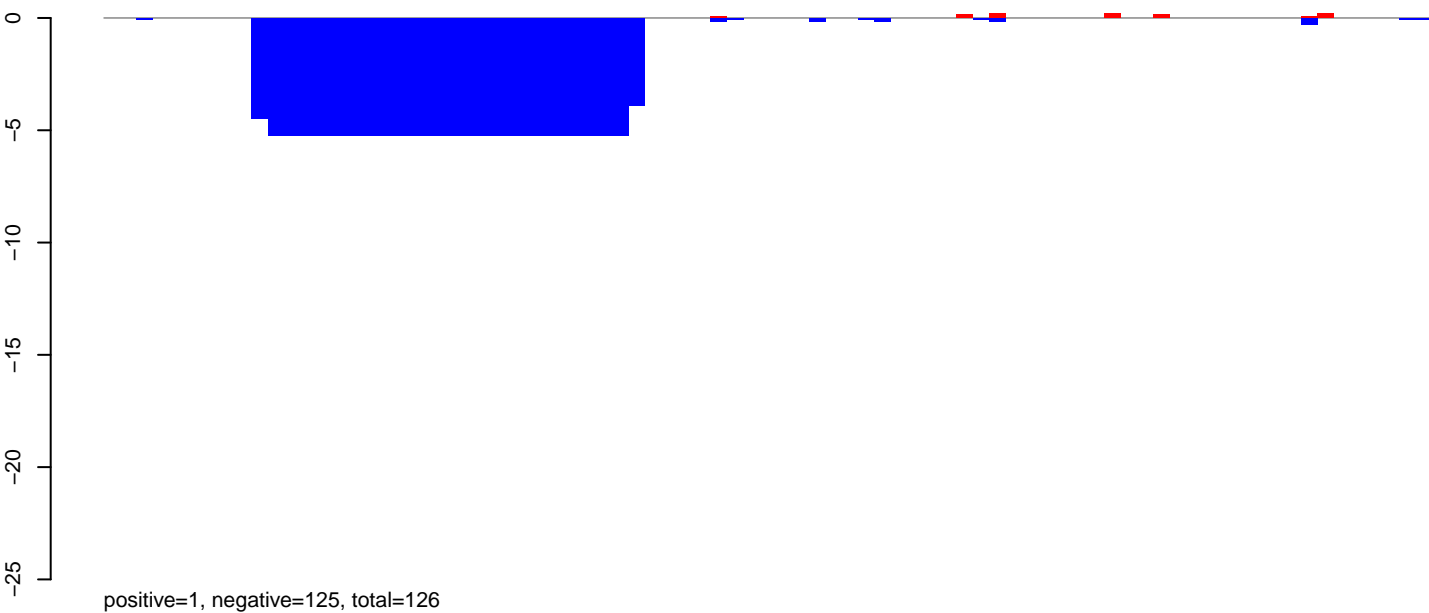
CuQuin_Mcarc_NL.rep



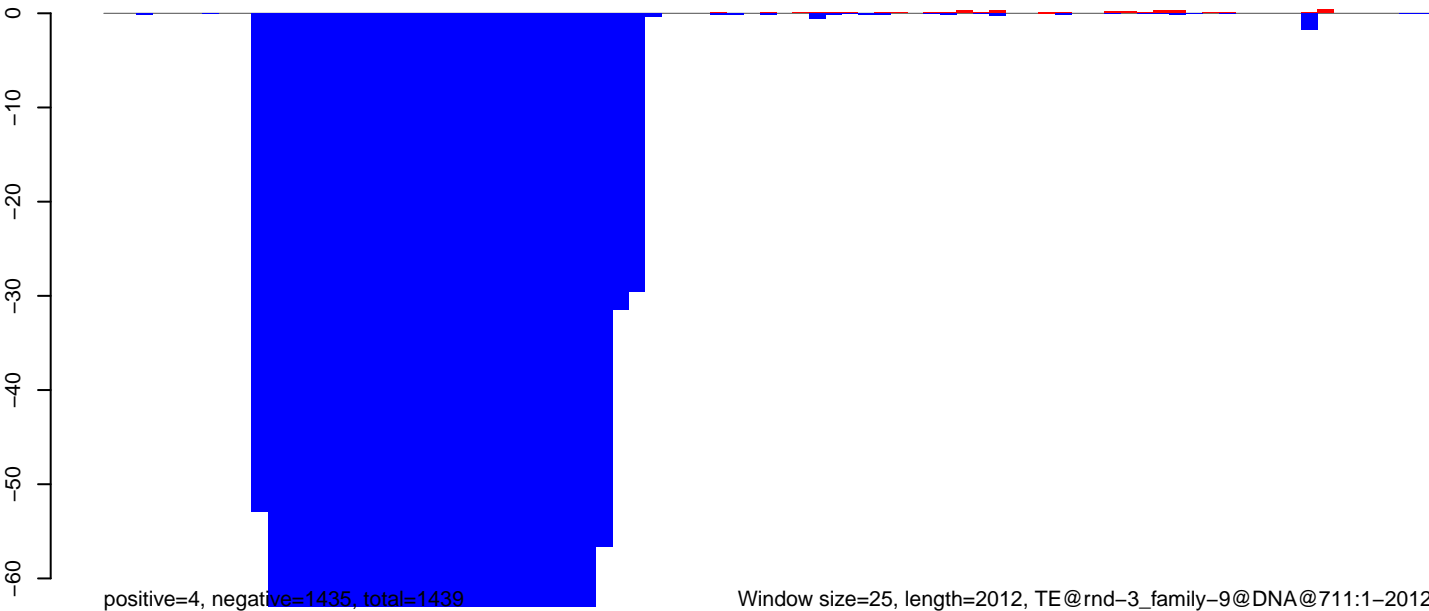
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



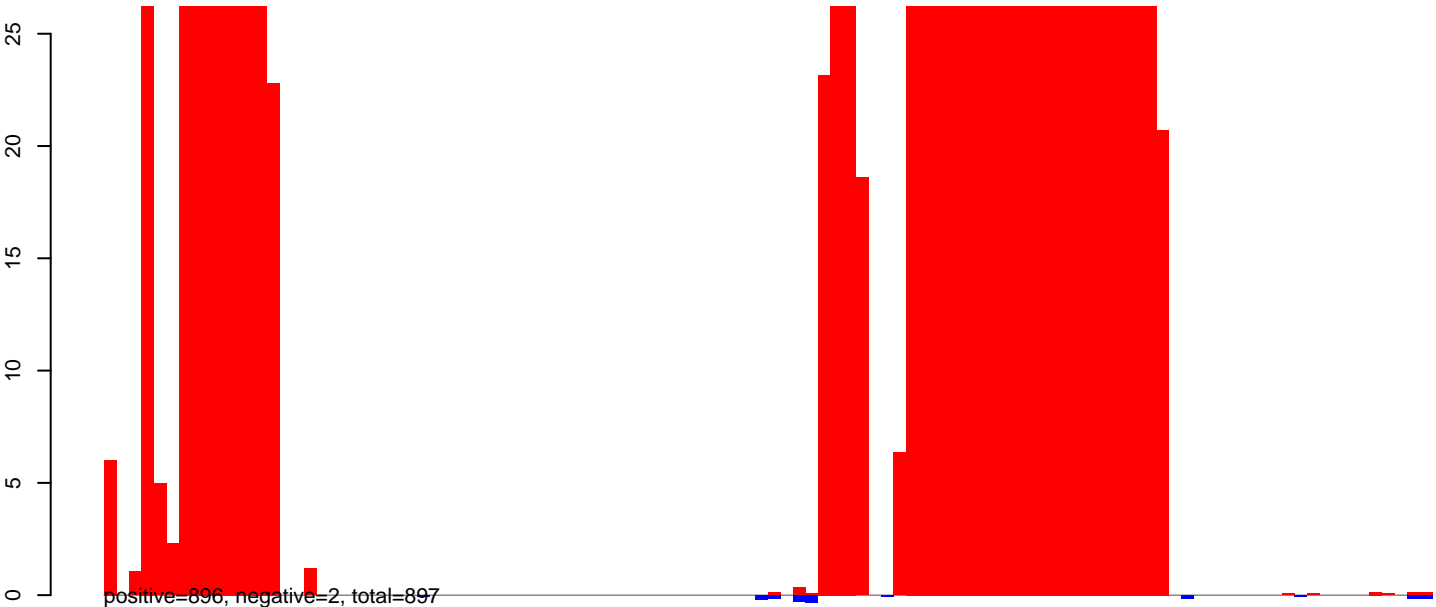
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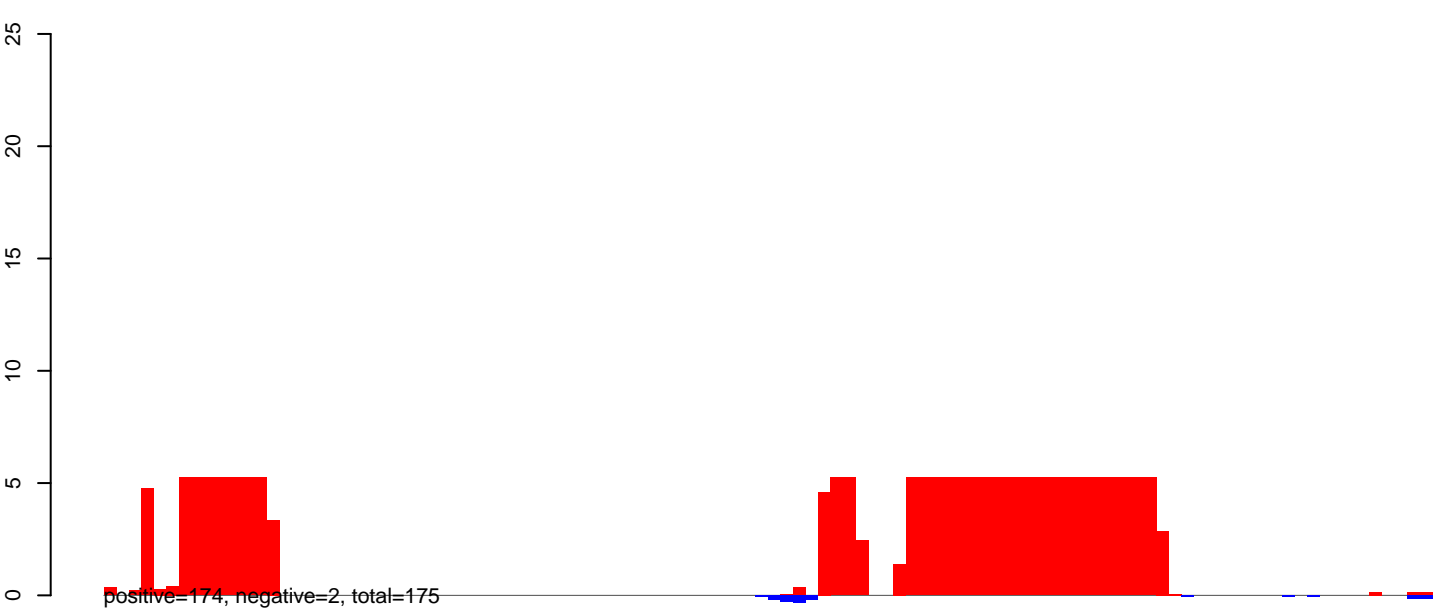
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0 500 1000 1500 2000

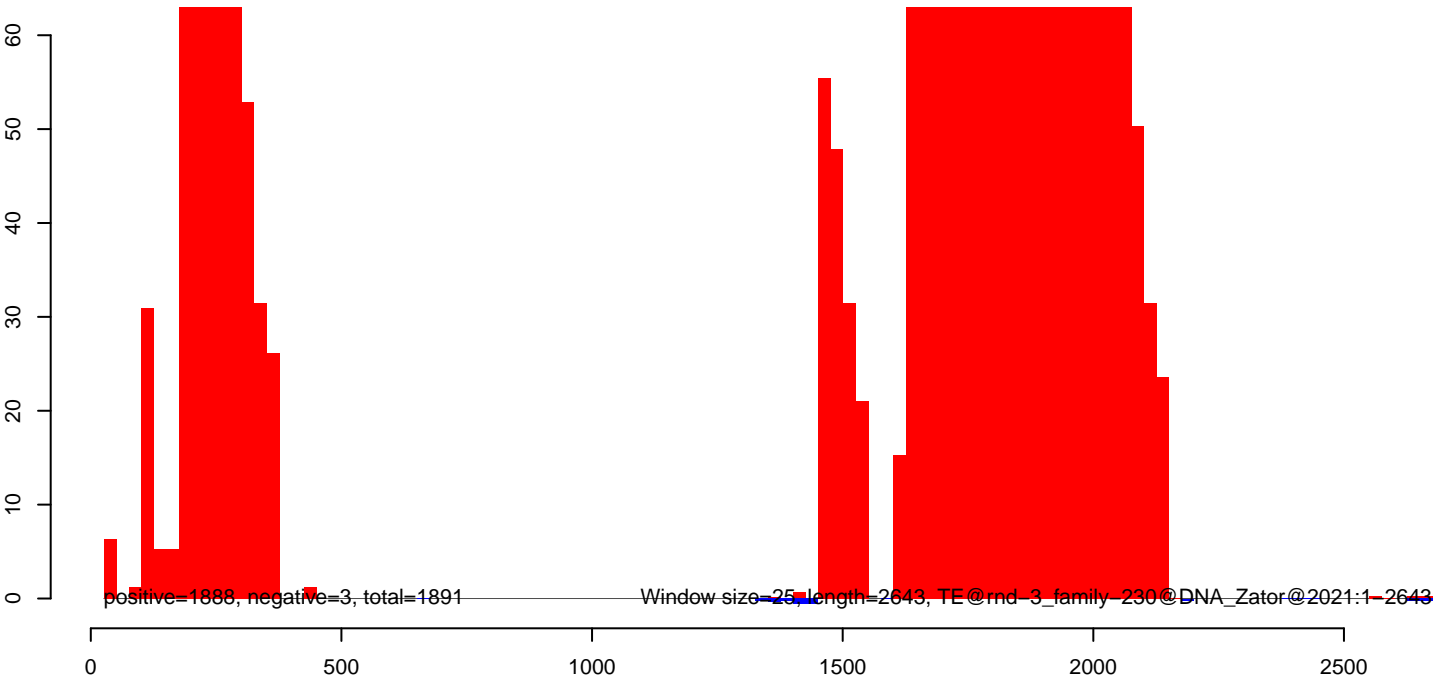
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CuQuin_Mcarc_NL.24_35.rep

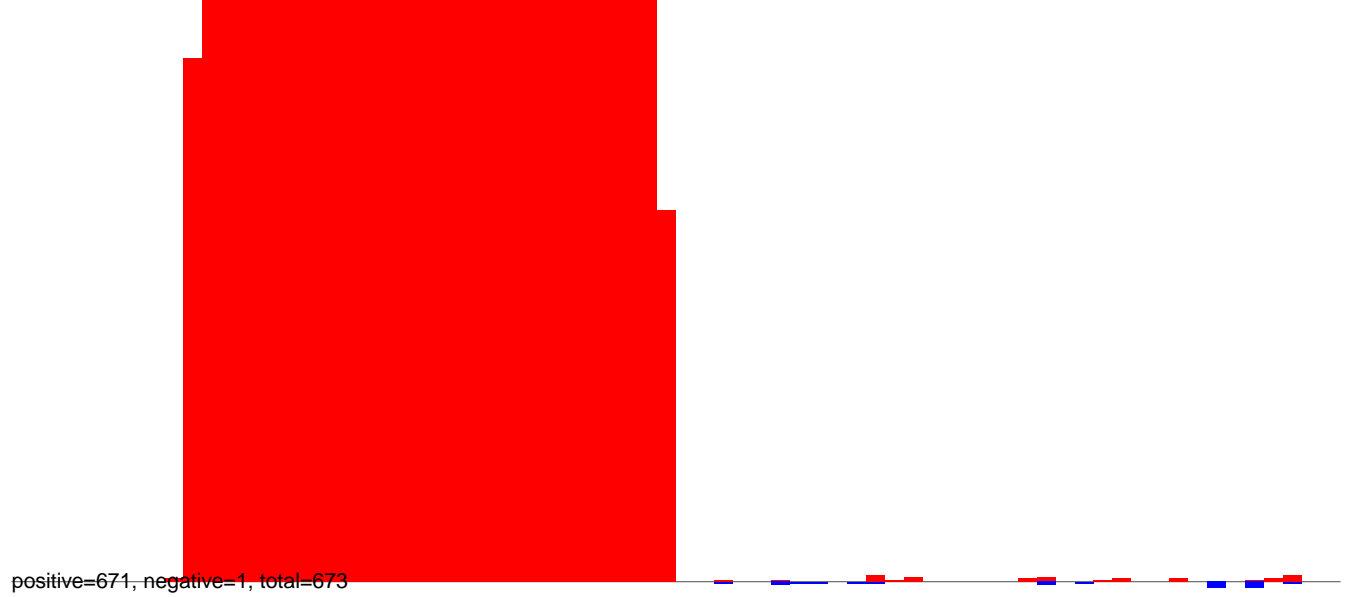


CuQuin_Mcarc_NL.rep

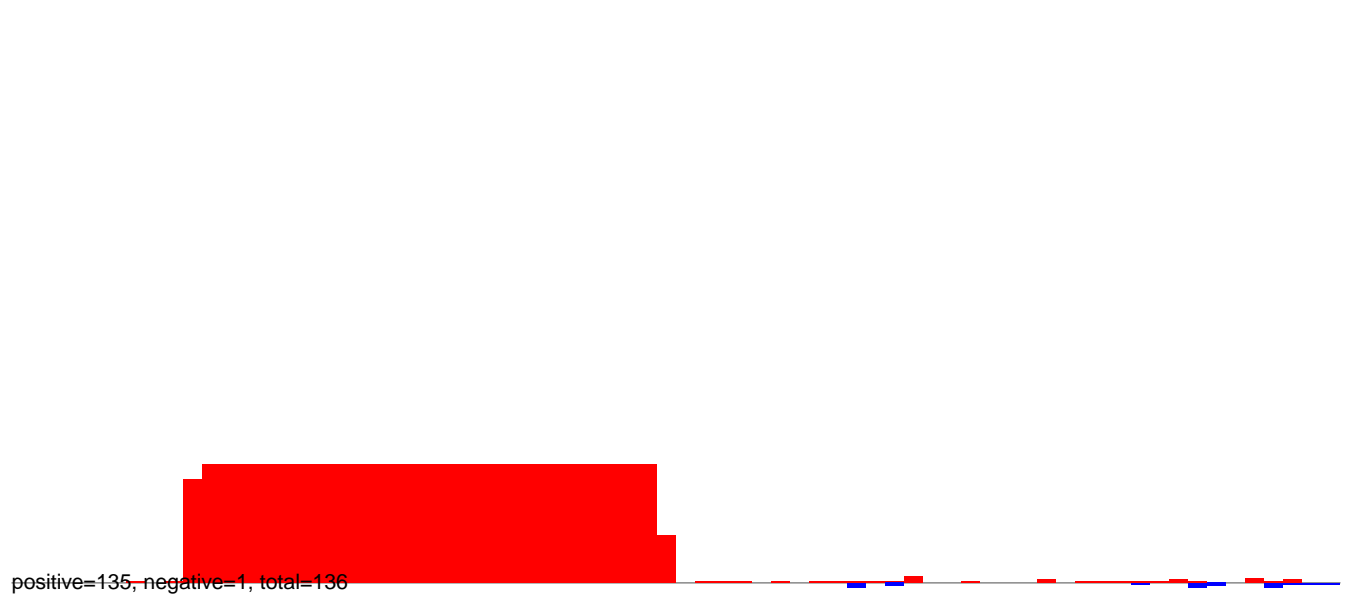


Window size=25, length=2643, TE@rnd-3_family-230@DNA_Zator@2021:1-2643

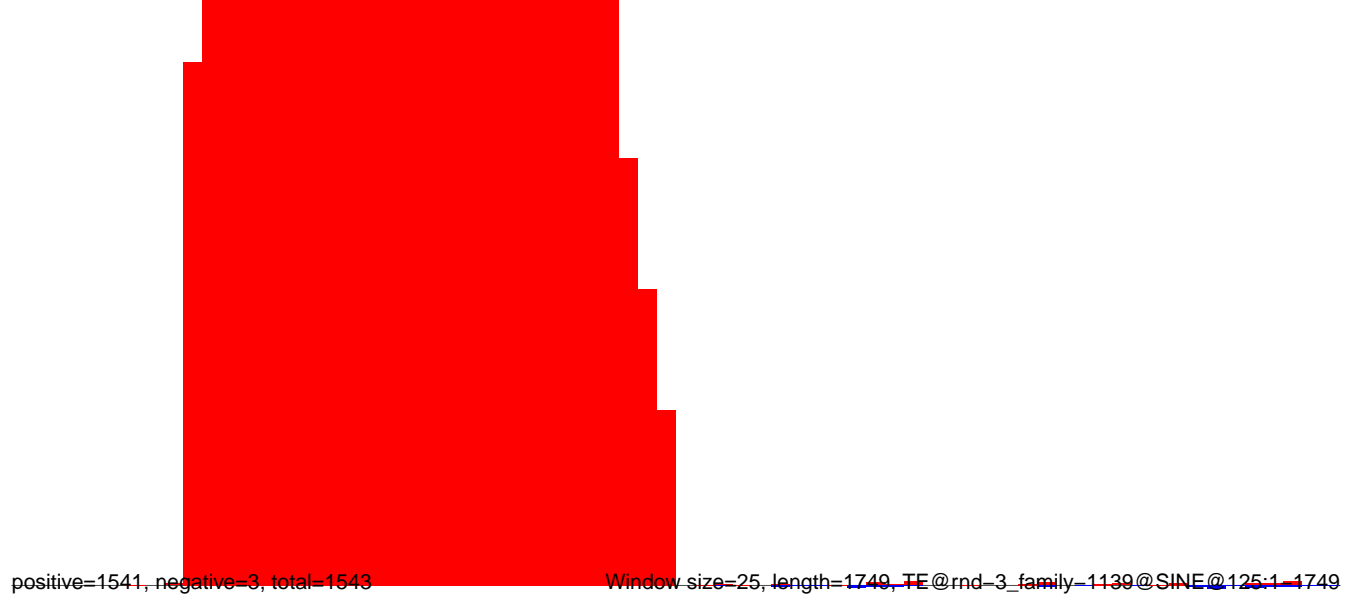
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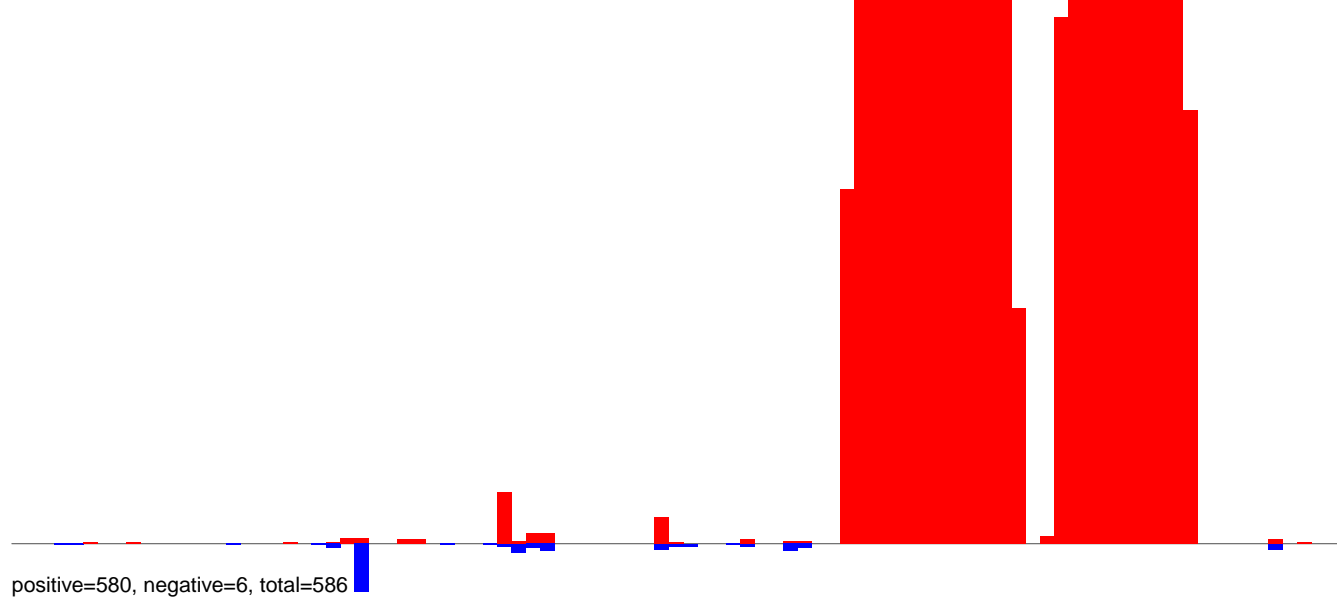
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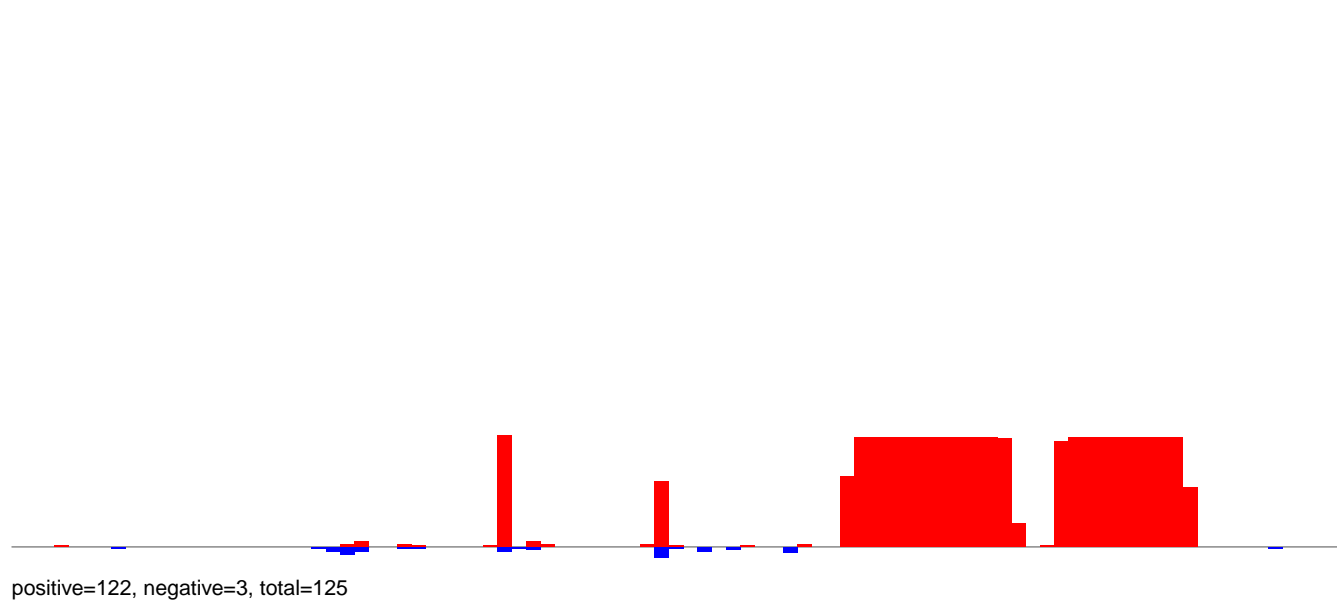
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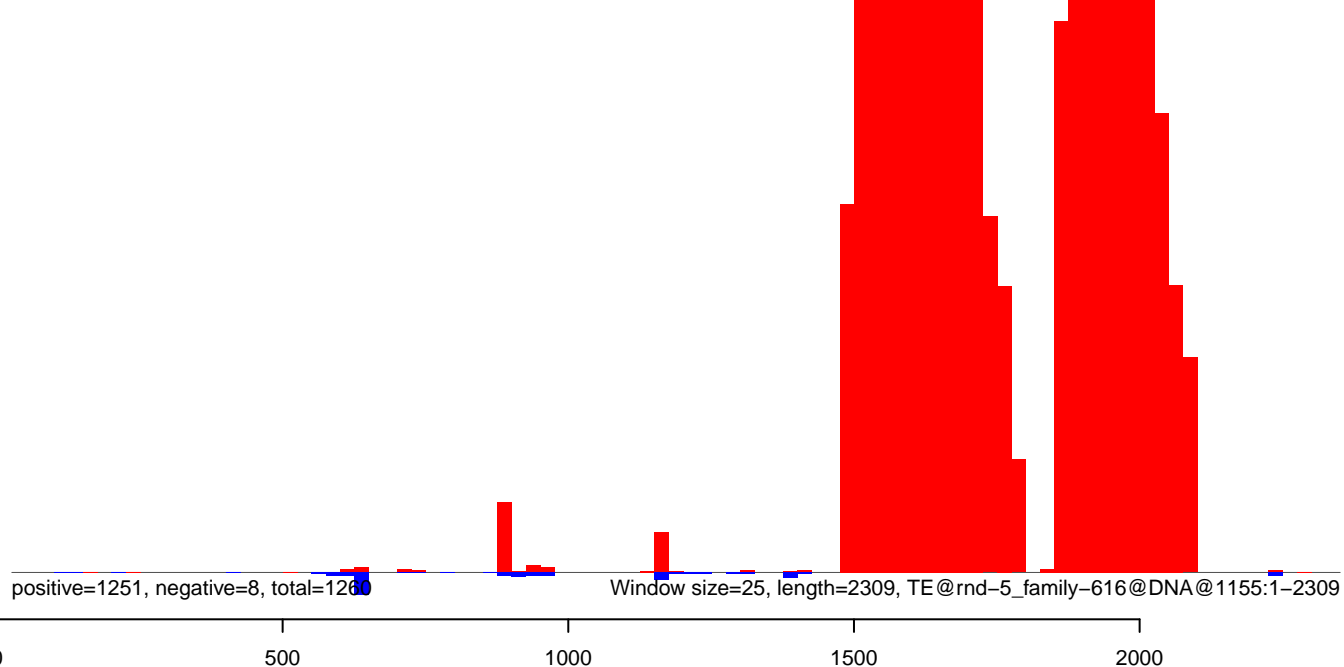
CuQuin_Mcarc_NL.18_23.rep



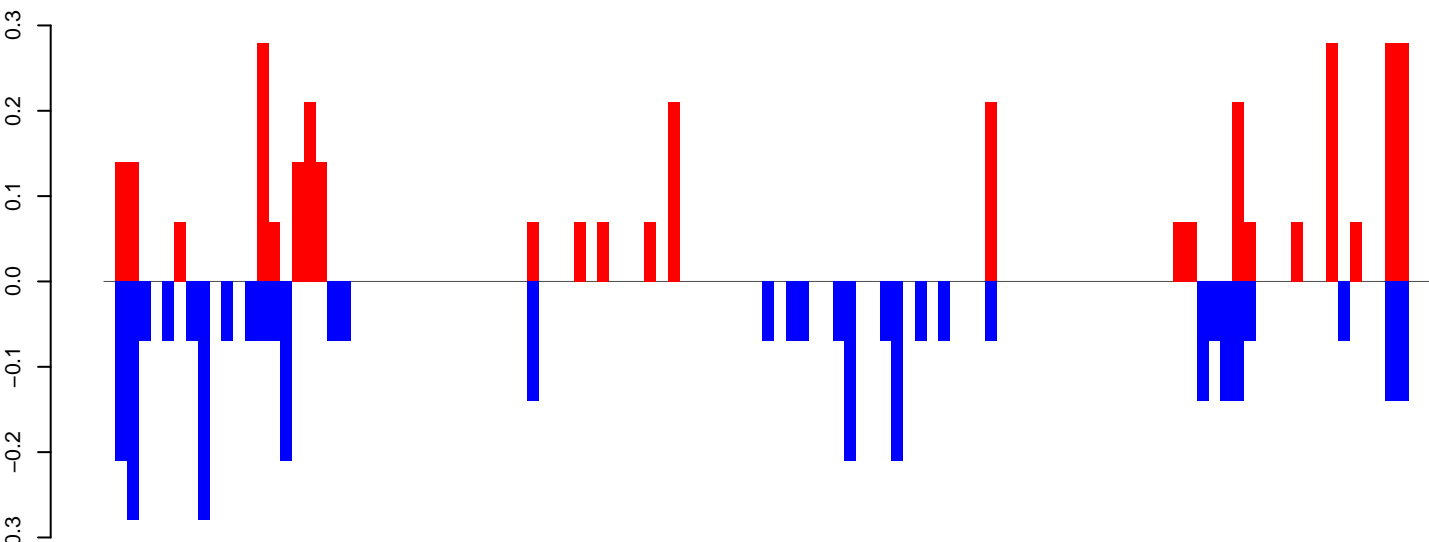
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CuQuin_Mcarc_NL.rep

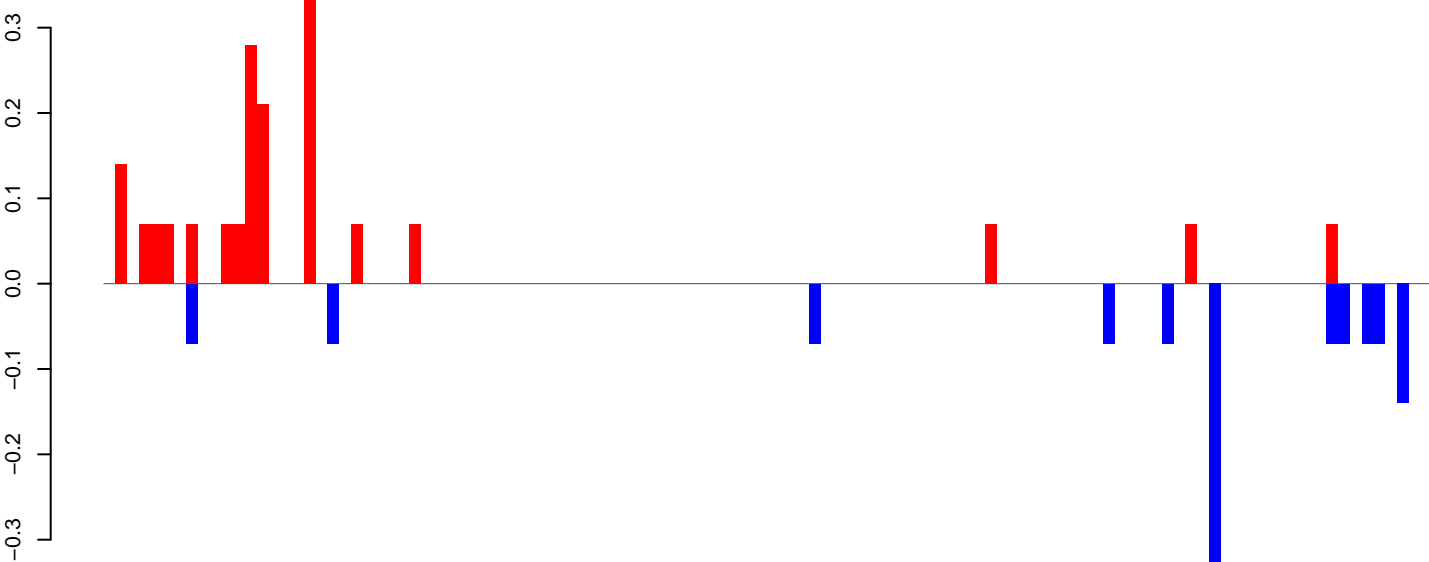


CuQuin_Mcarc_NL.18_23.rep



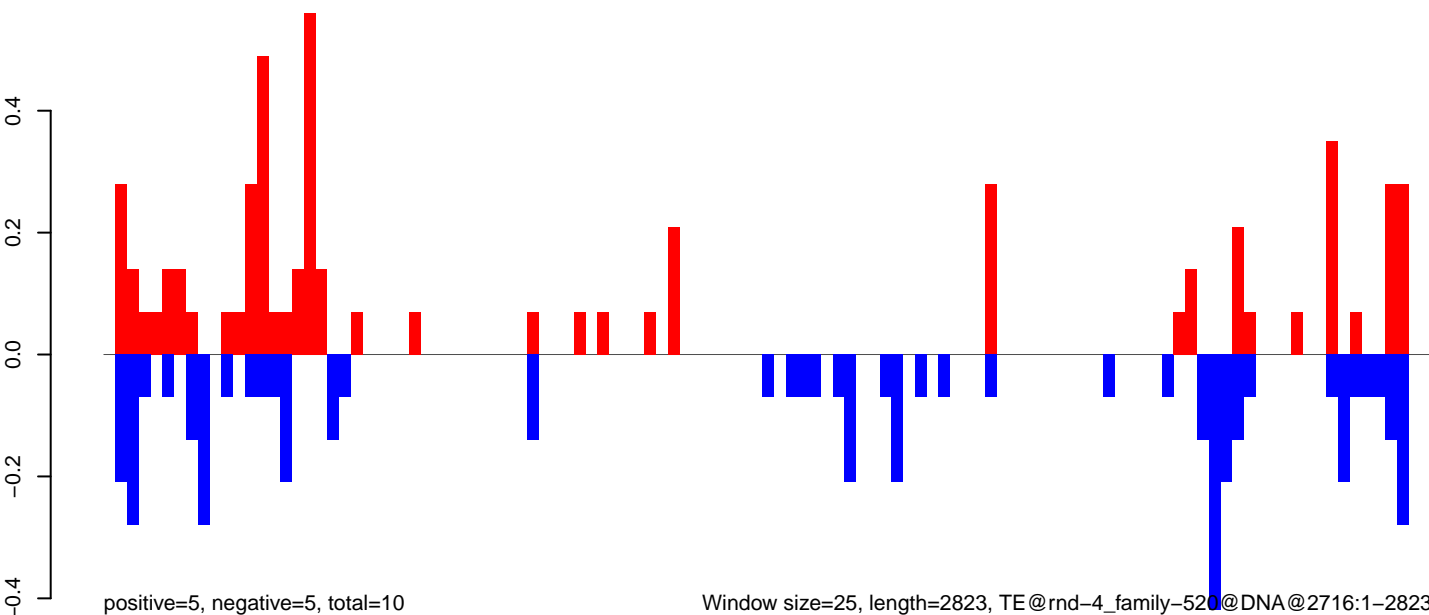
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CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=1, total=3

CuQuin_Mcarc_NL.rep



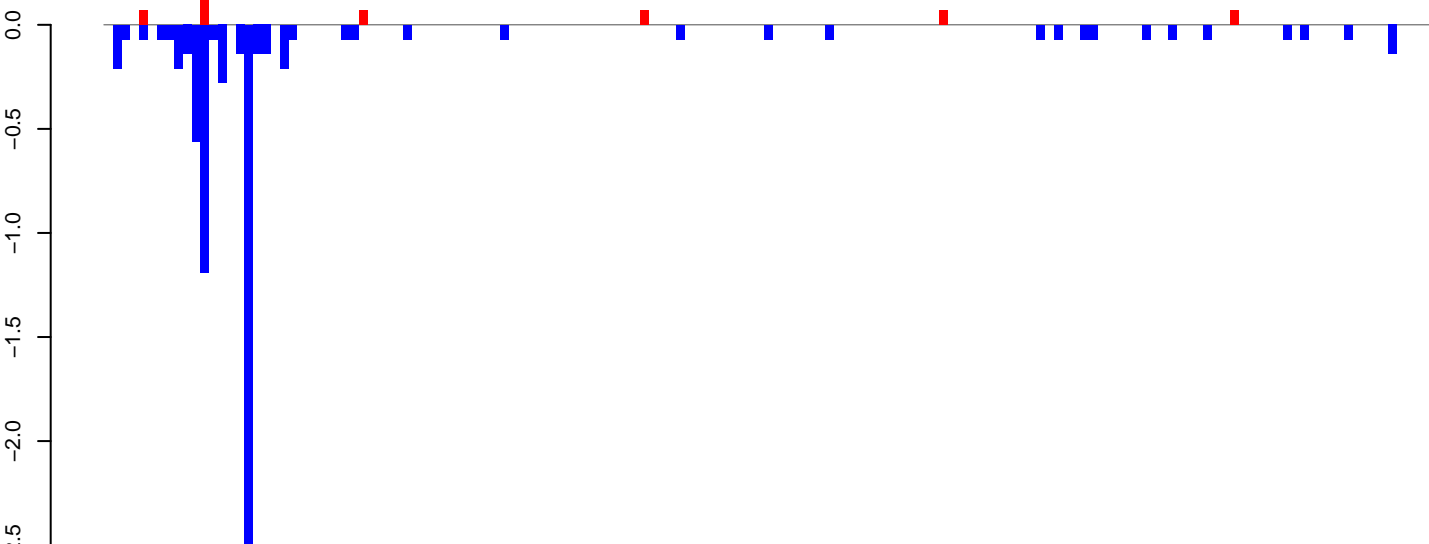
positive=5, negative=5, total=10

Window size=25, length=2823, TE@rnd-4_family-520@DNA@2716:1-2823

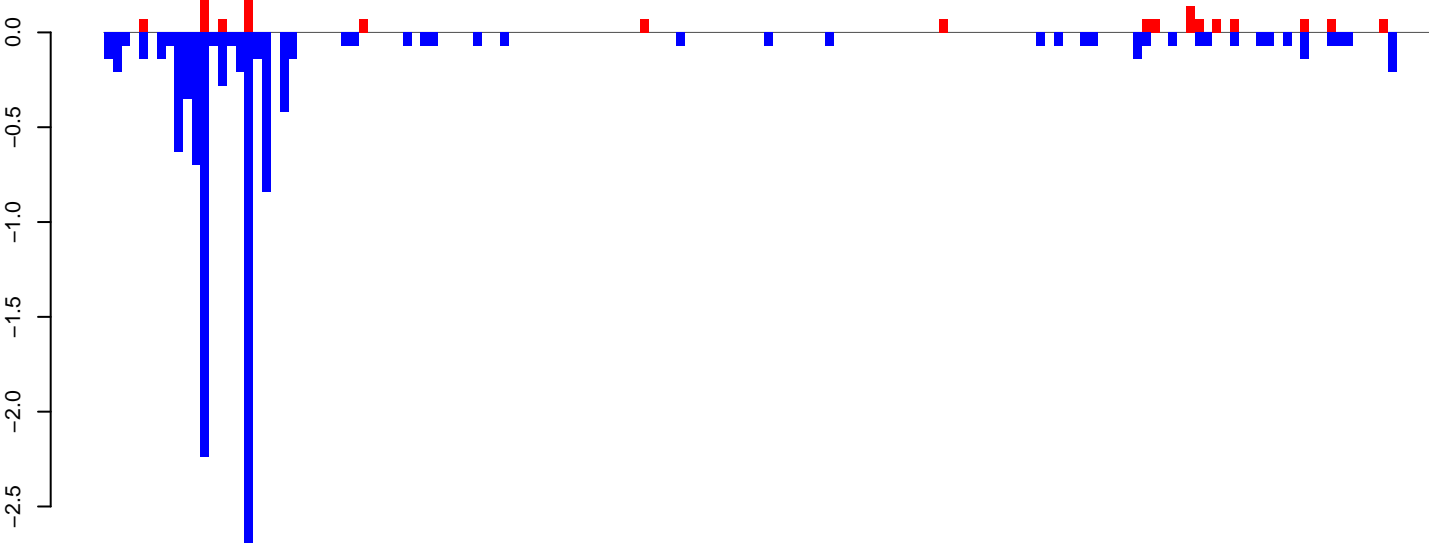
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

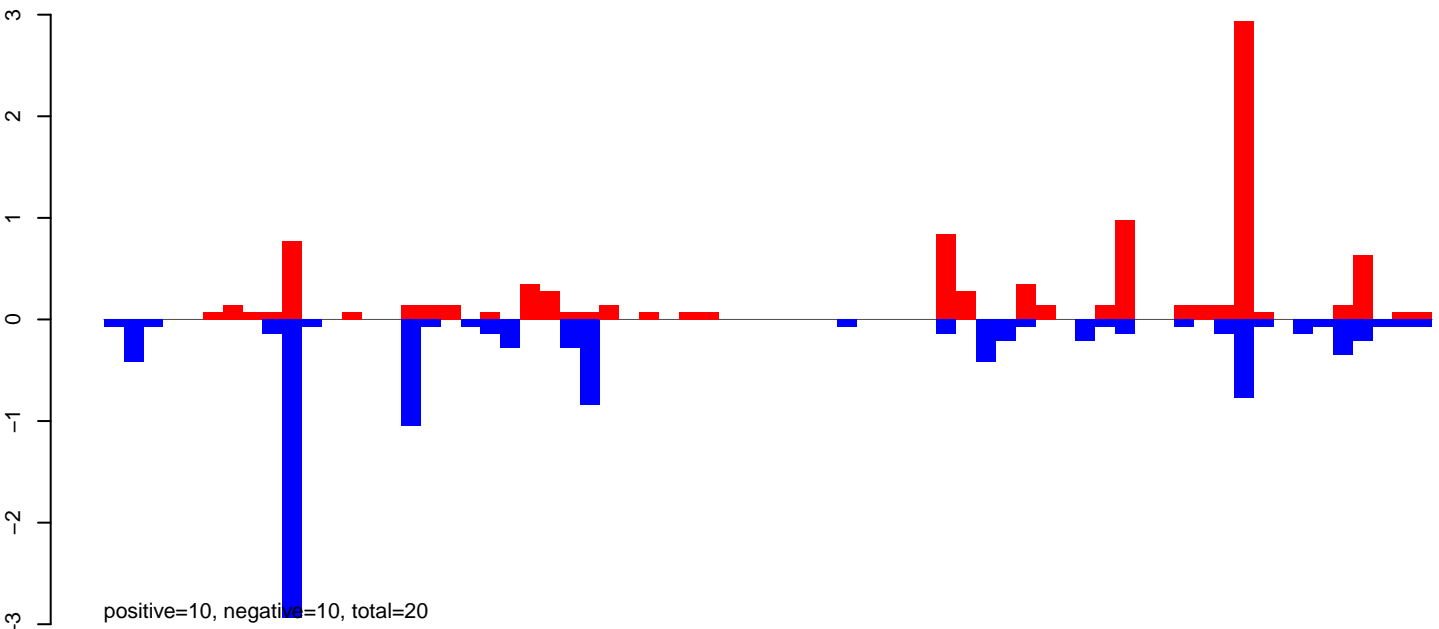


CuQuin_Mcarc_NL.rep

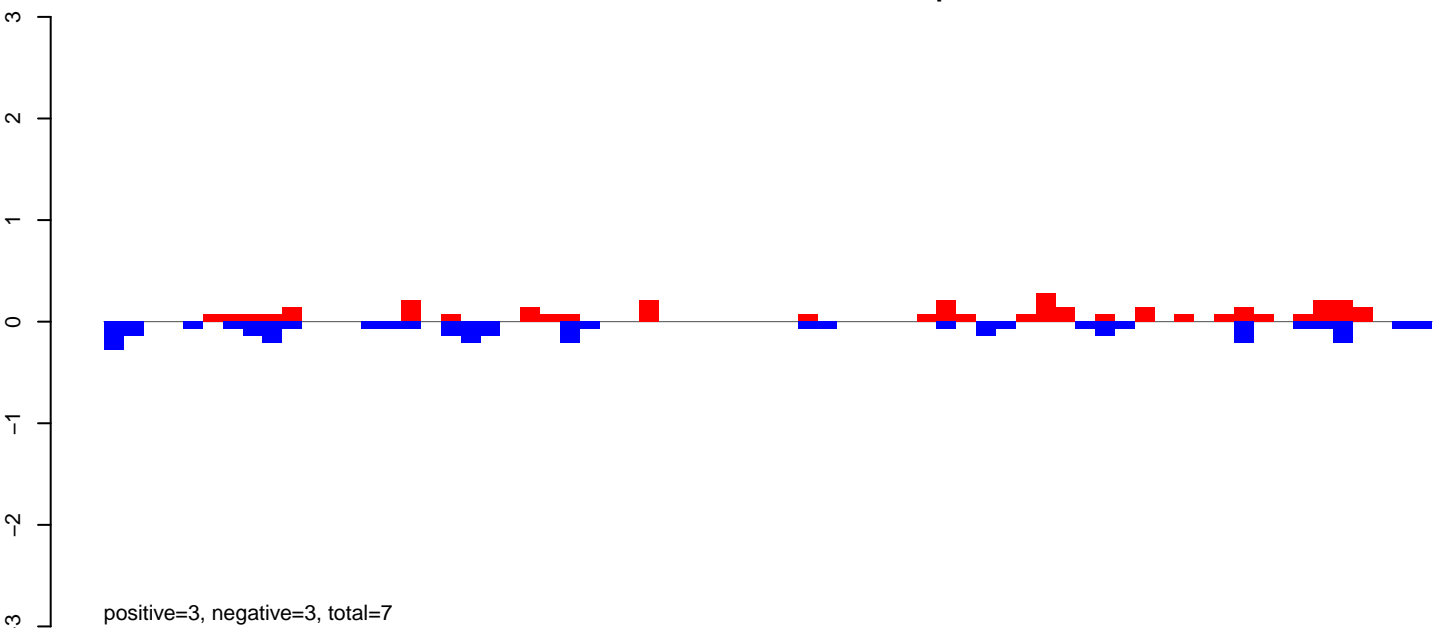


Window size=25, length=3771, TE@rnd-4_family-1142@LTR_Gypsy@2791:1-3771

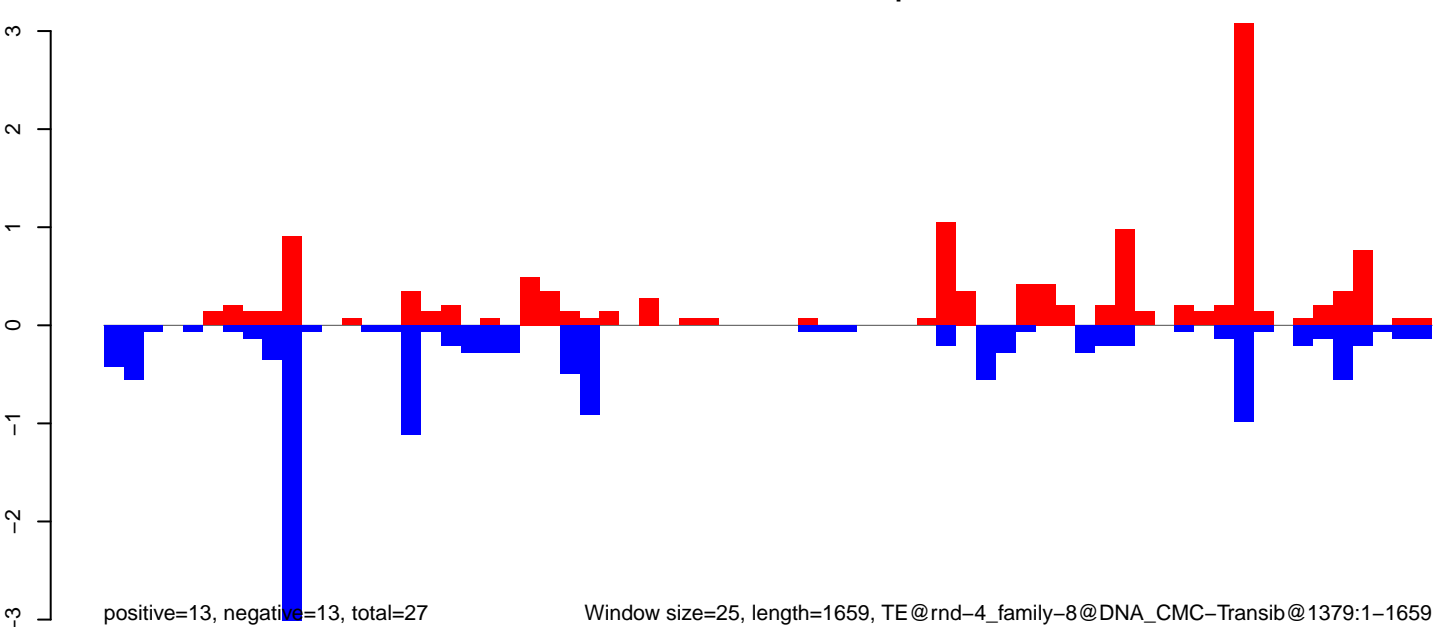
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CuQuin_Mcarc_NL.24_35.rep



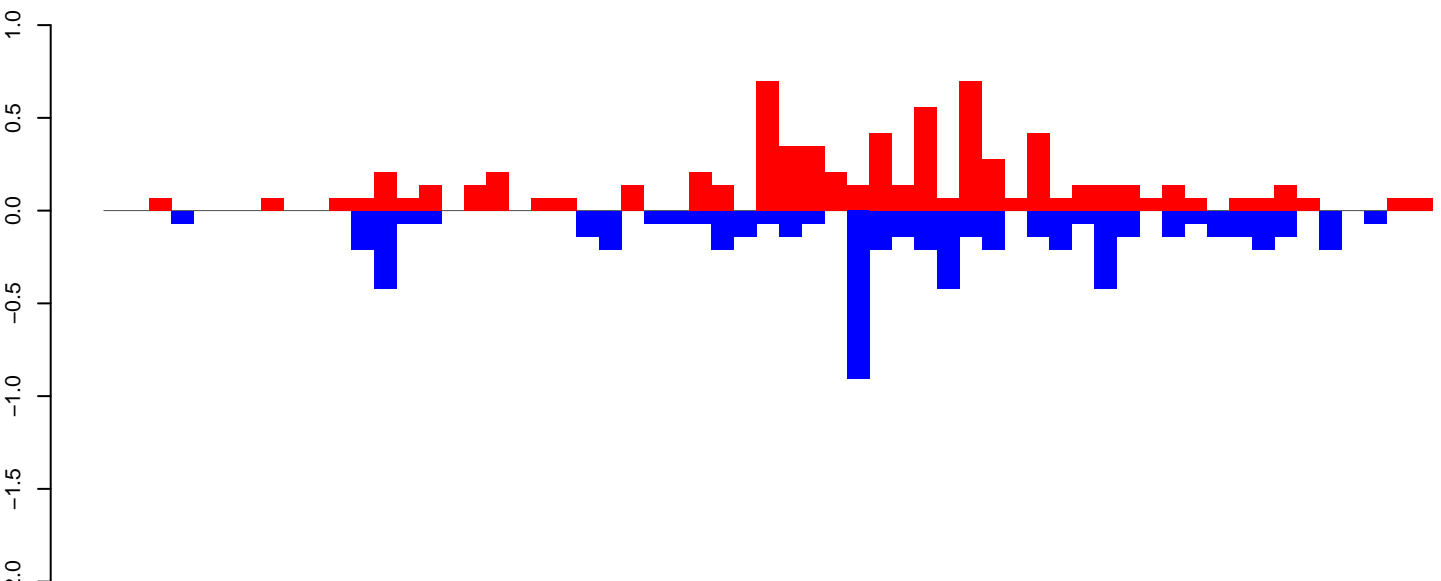
CuQuin_Mcarc_NL.rep



Window size=25, length=1659, TE@rnd-4_family-8@DNA_CMC-Transib@1379:1-1659

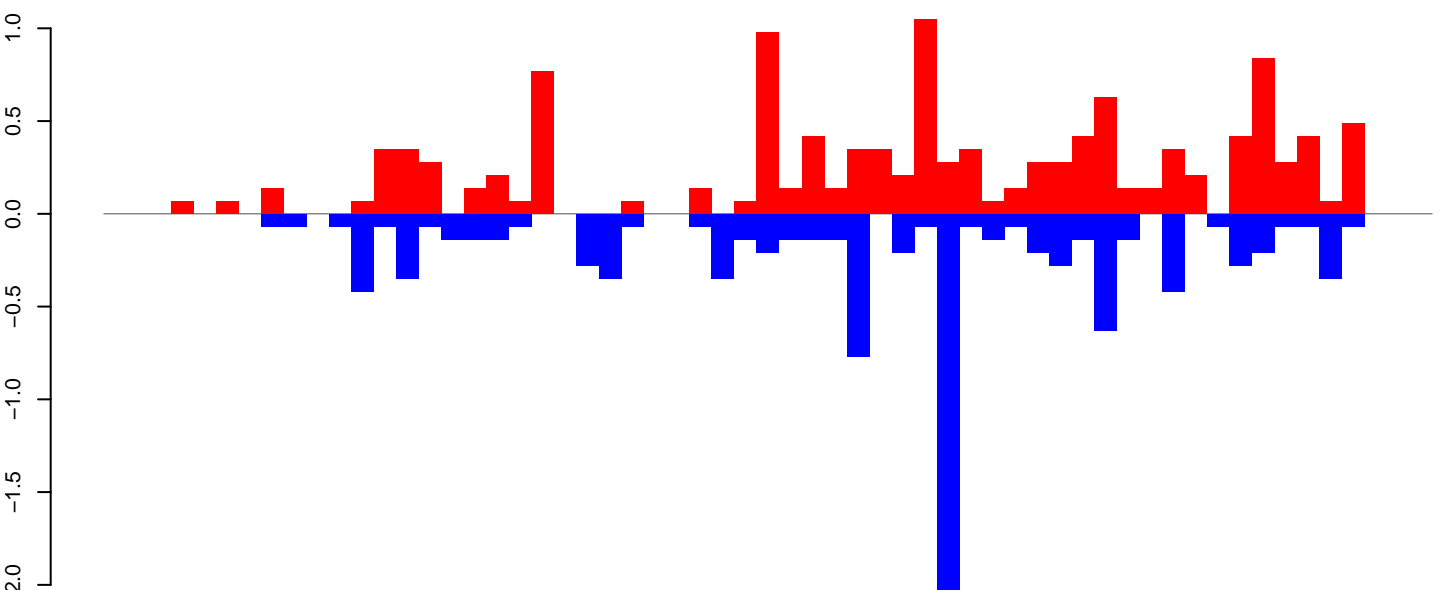
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CuQuin_Mcarc_NL.18_23.rep



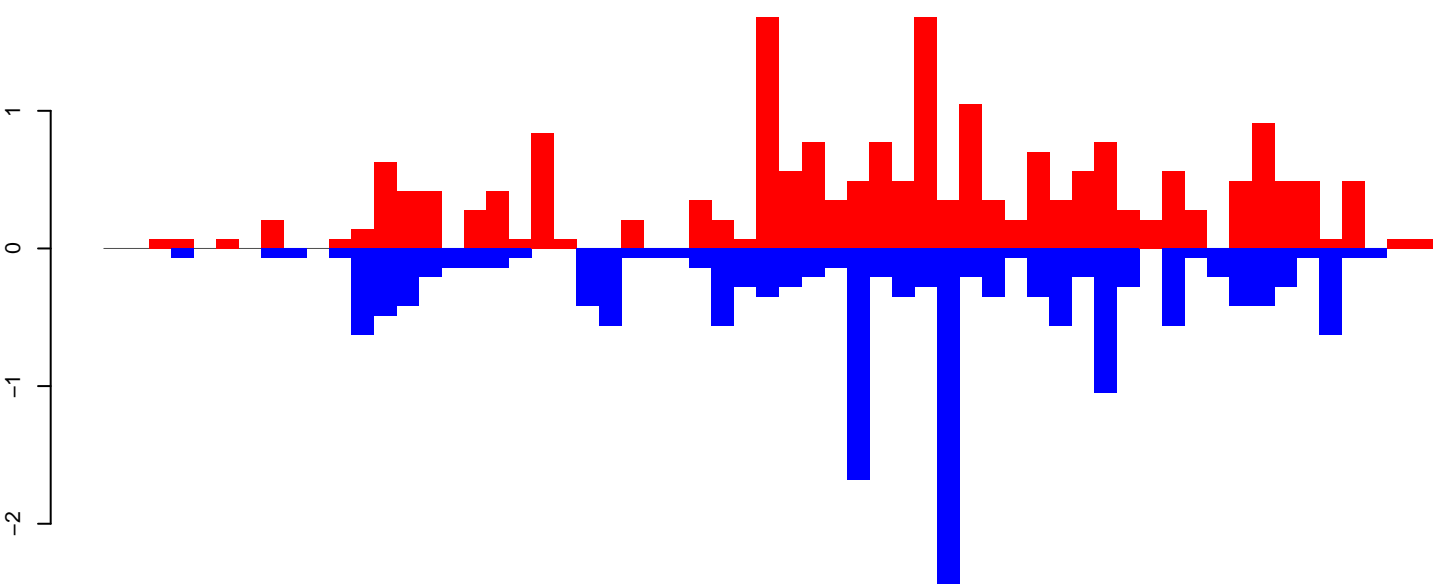
positive=7, negative=6, total=14

CuQuin_Mcarc_NL.24_35.rep



positive=12, negative=10, total=22

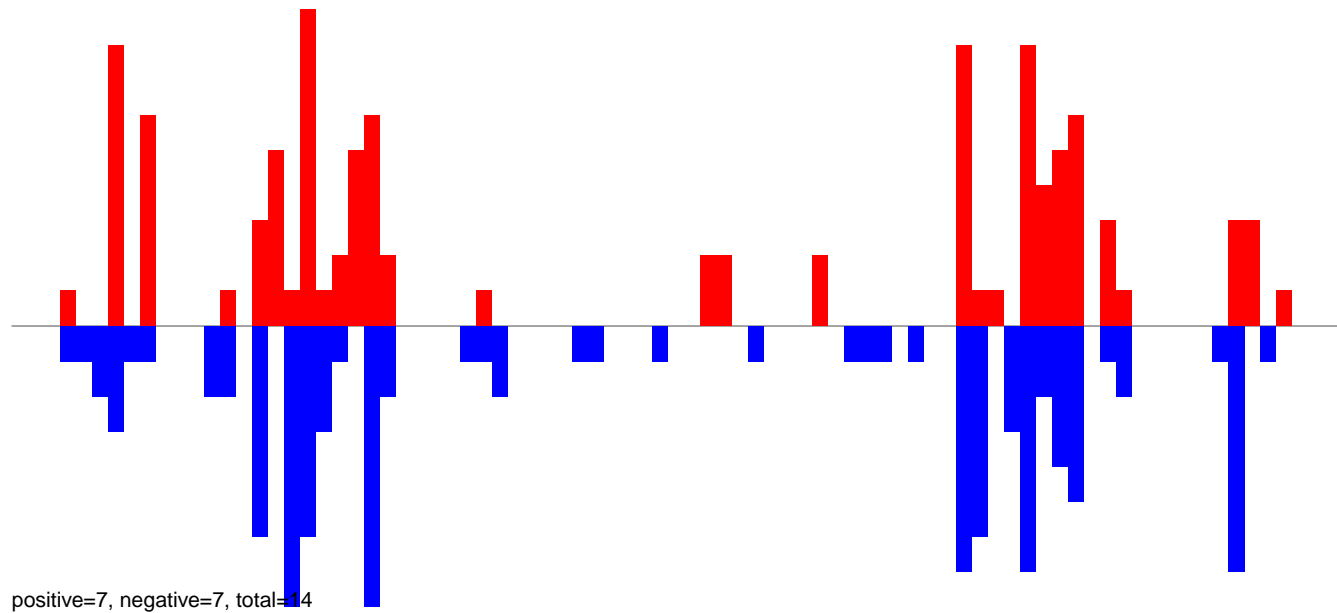
CuQuin_Mcarc_NL.rep



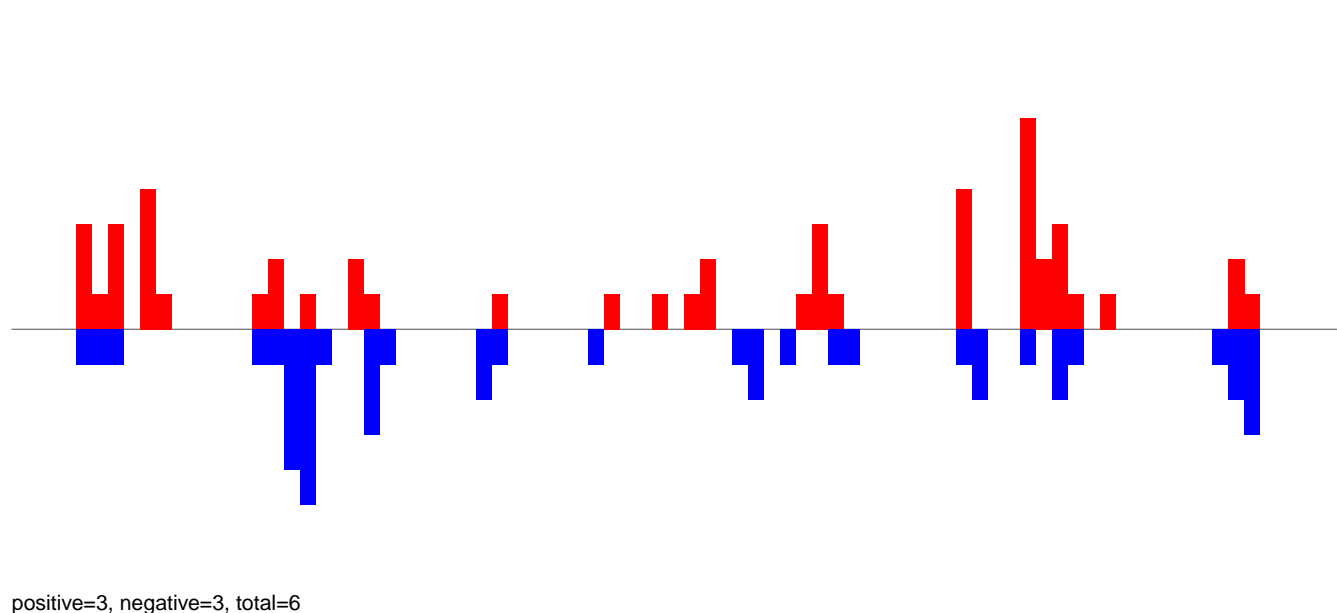
positive=20, negative=17, total=37

Window size=25, length=1471 TE@rnd-4_family-354@Unknown@2137:1-1471

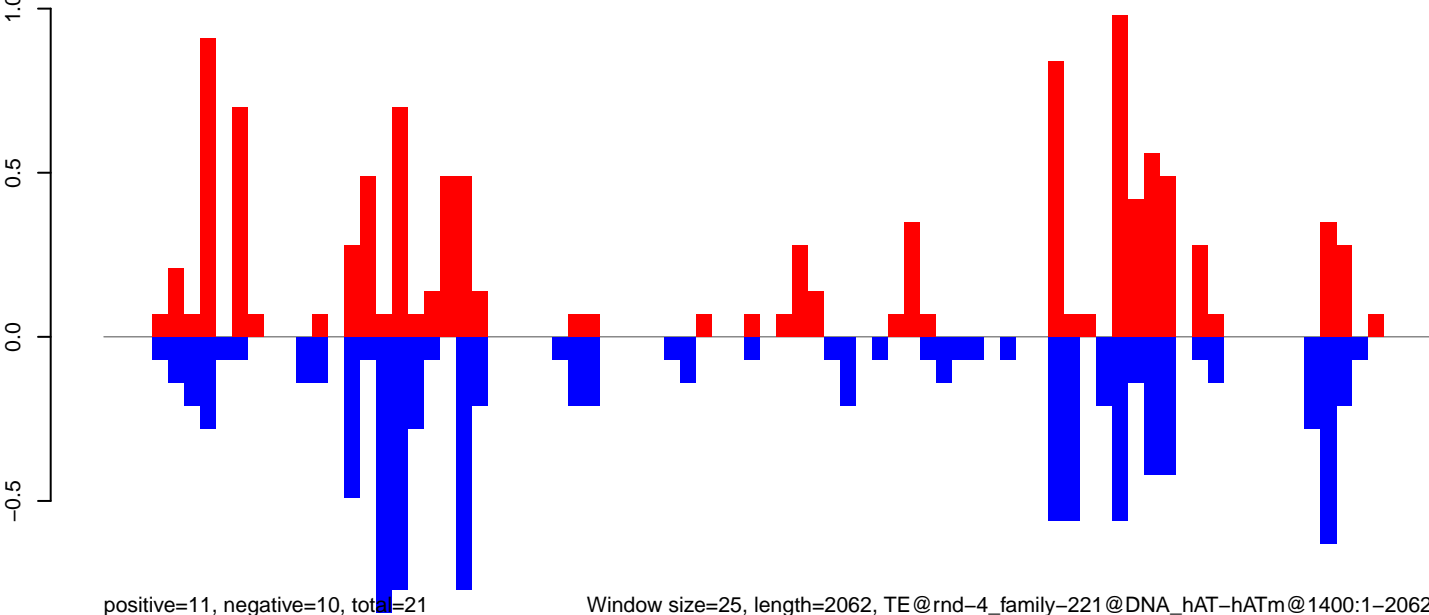
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CuQuin_Mcarc_NL.24_35.rep



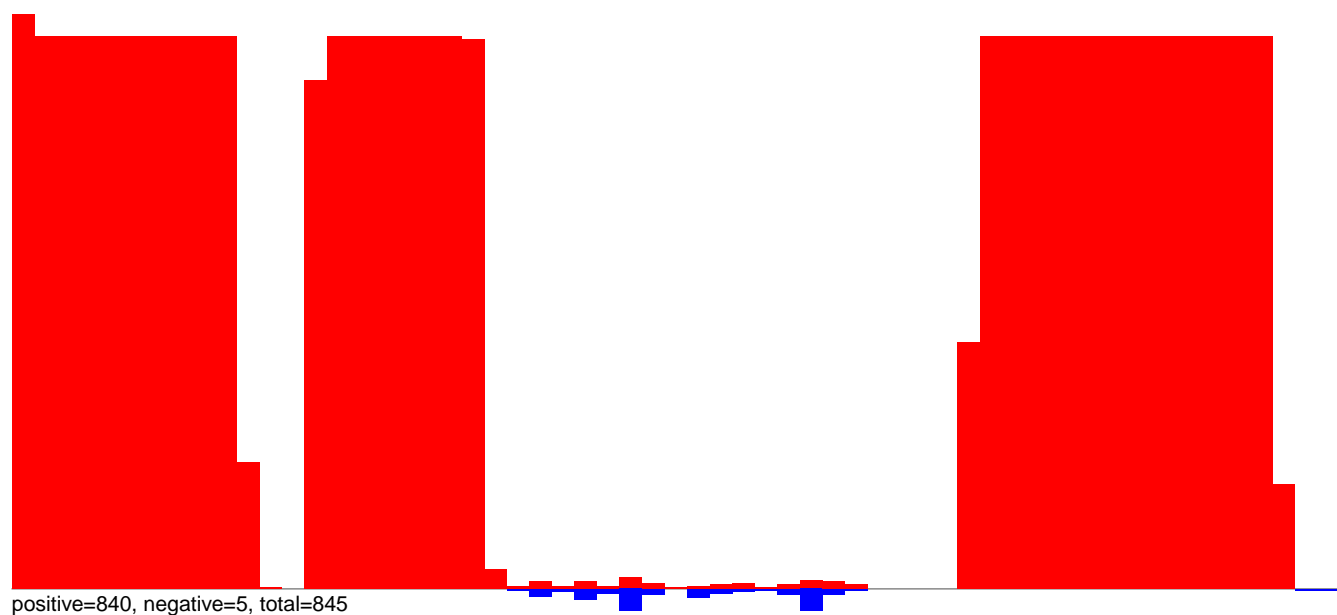
CuQuin_Mcarc_NL.rep



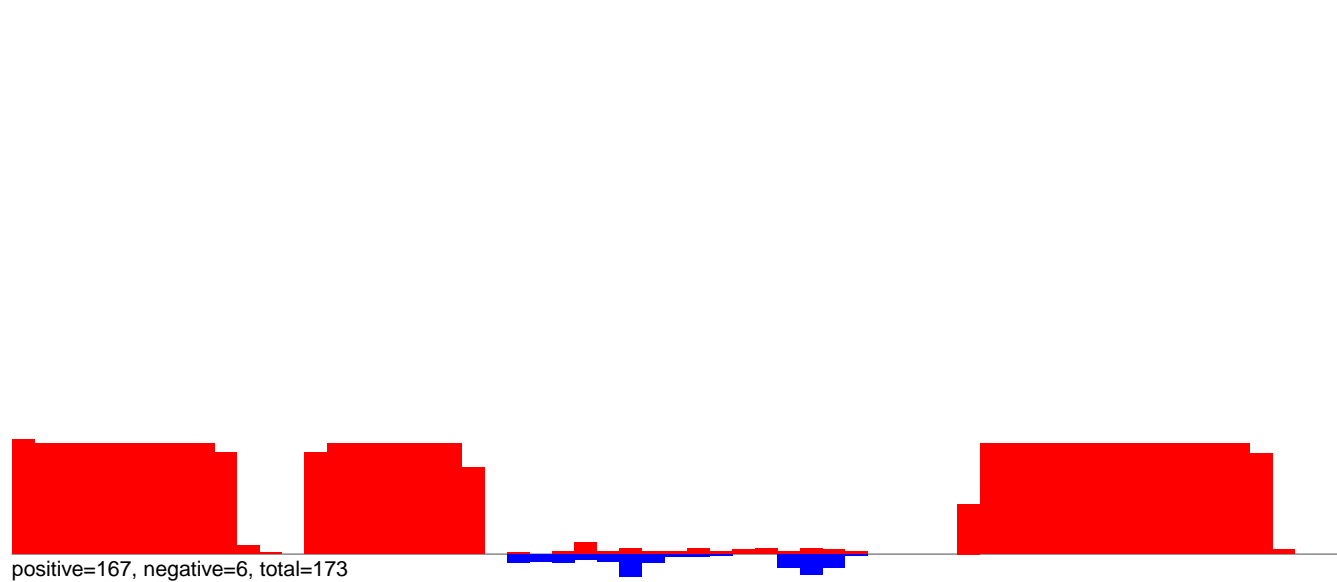
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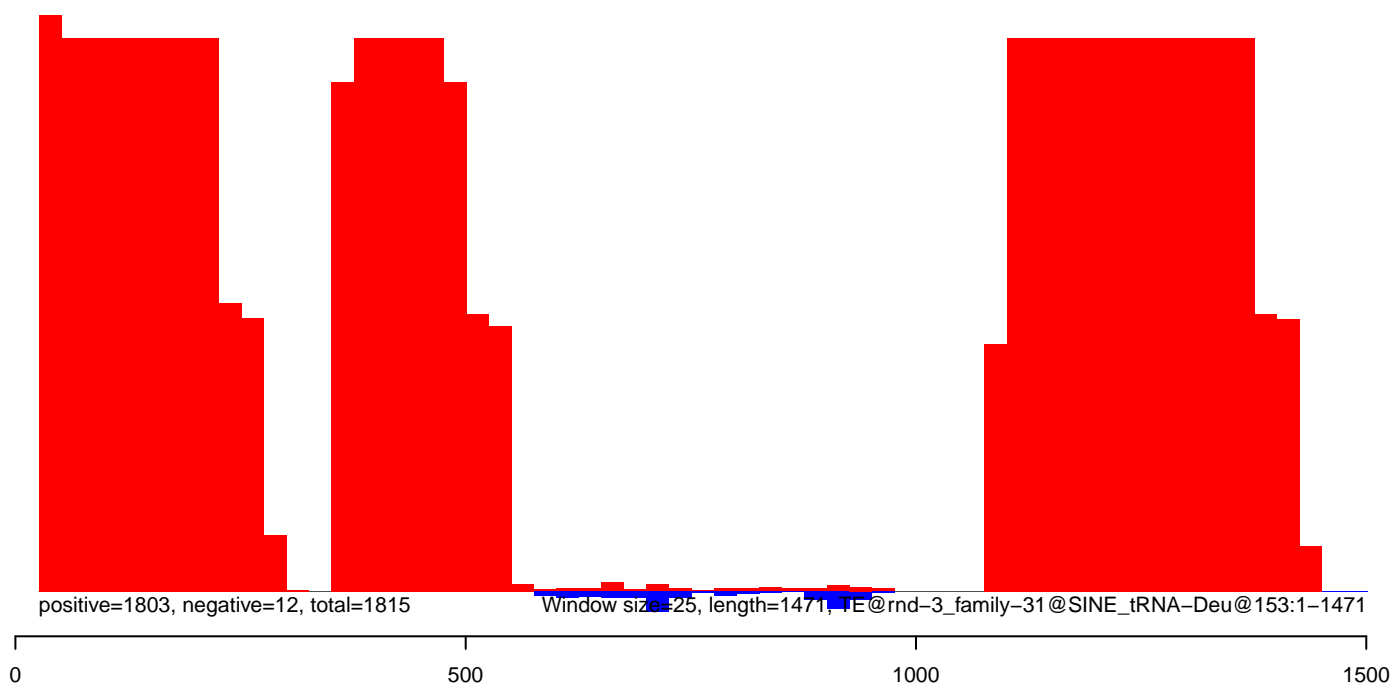
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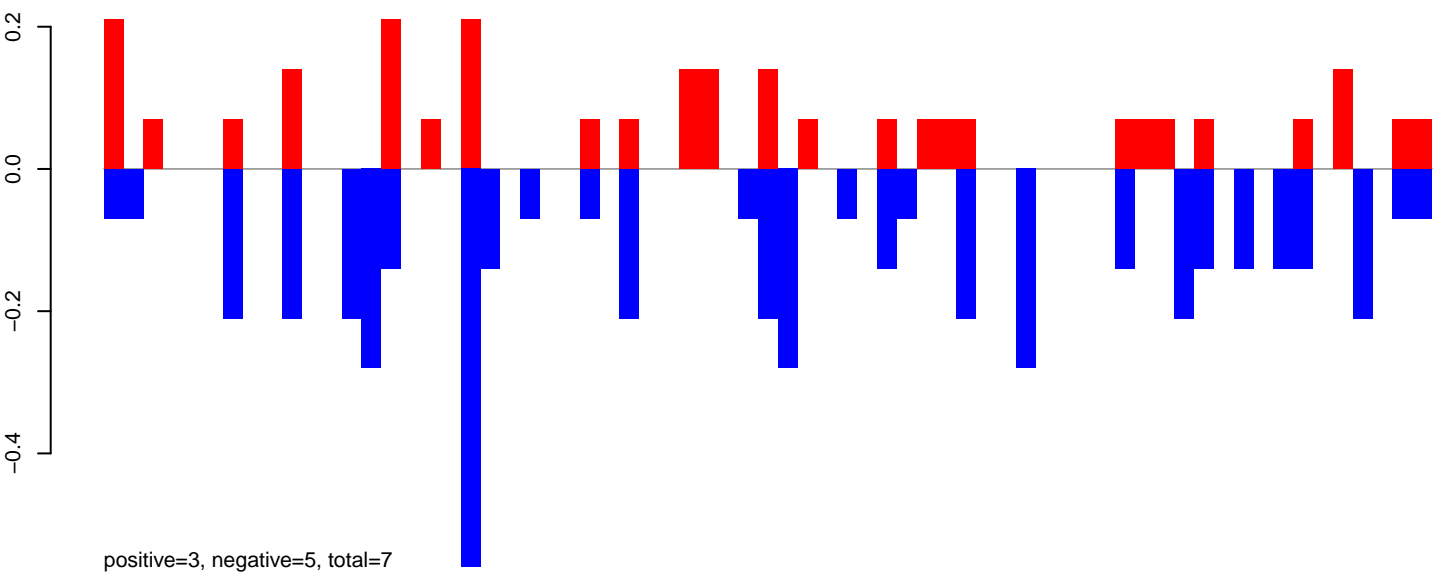
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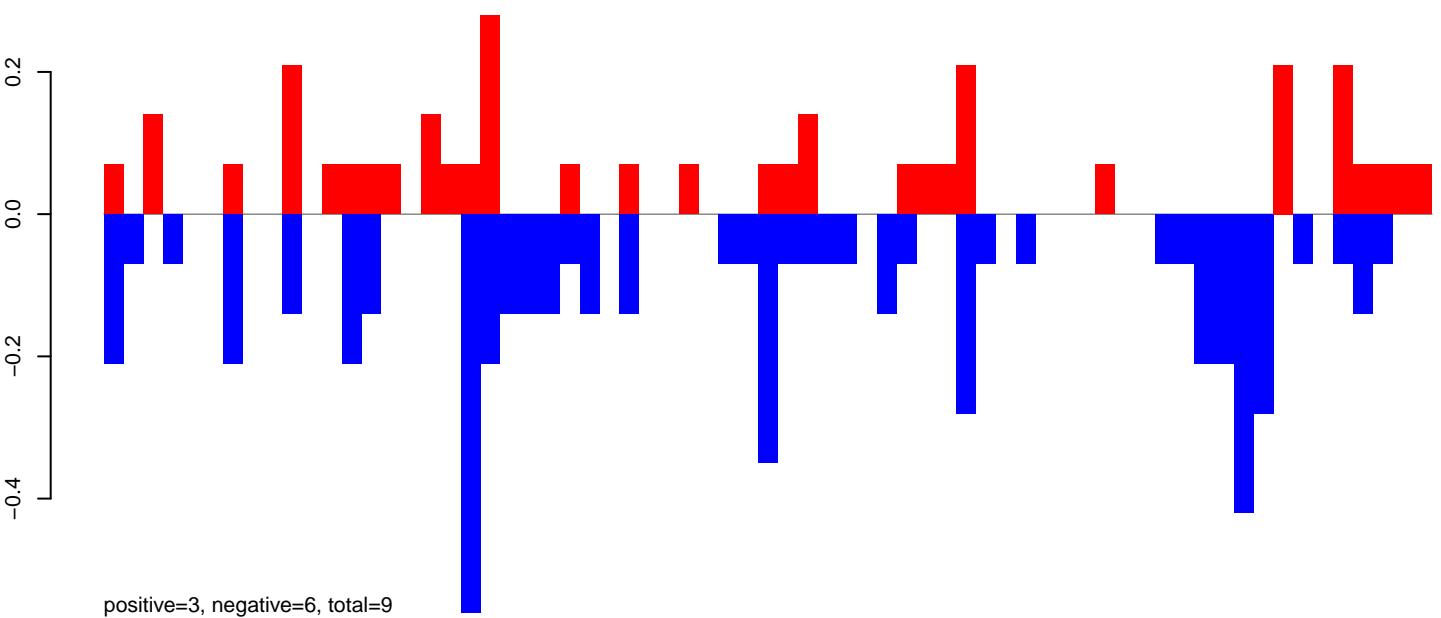
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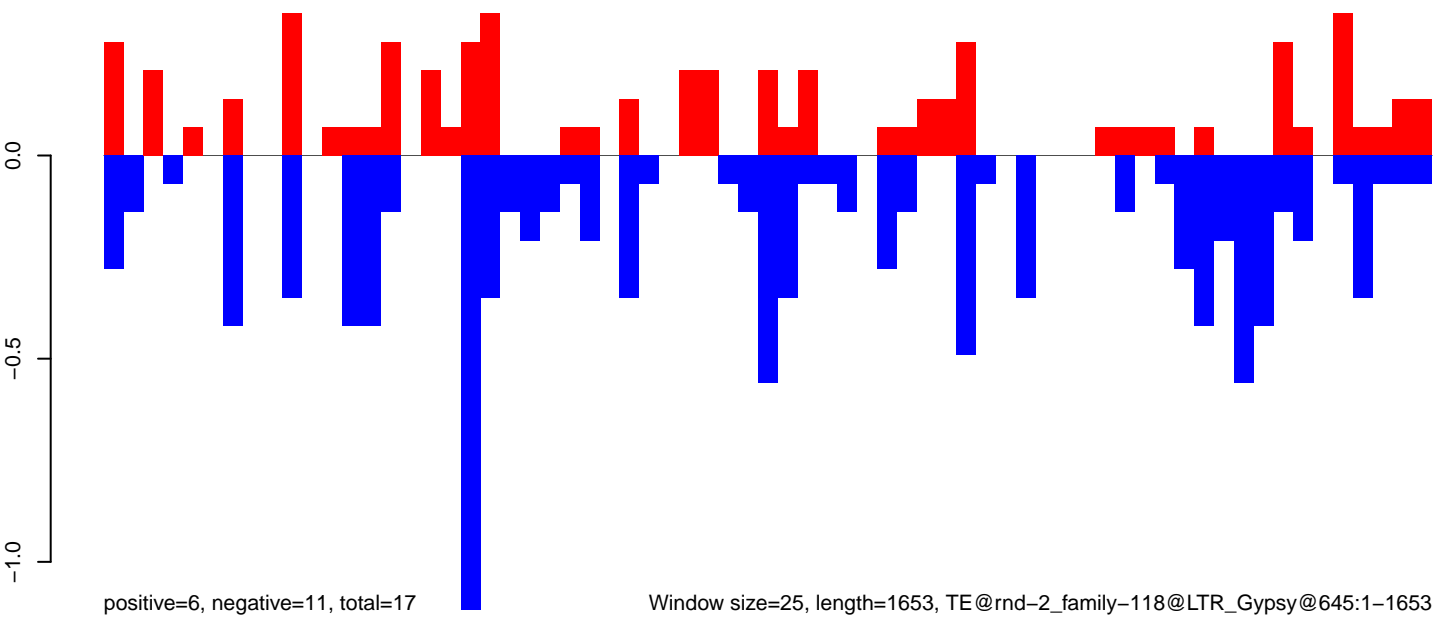
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



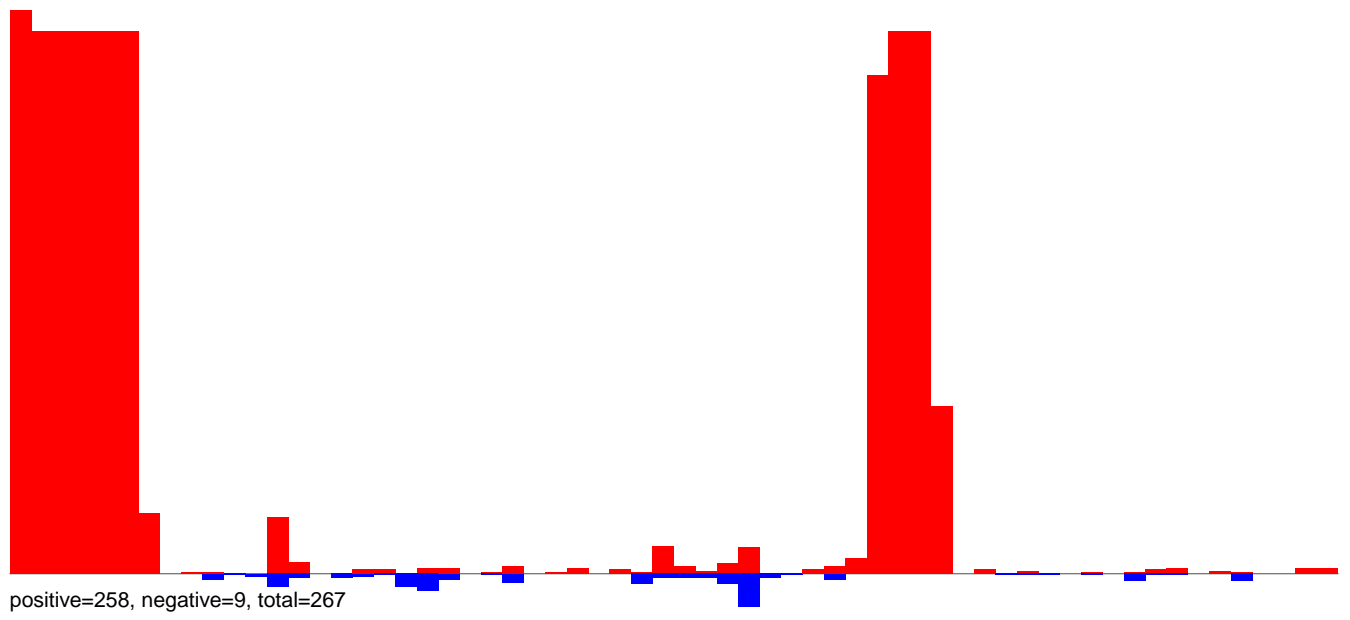
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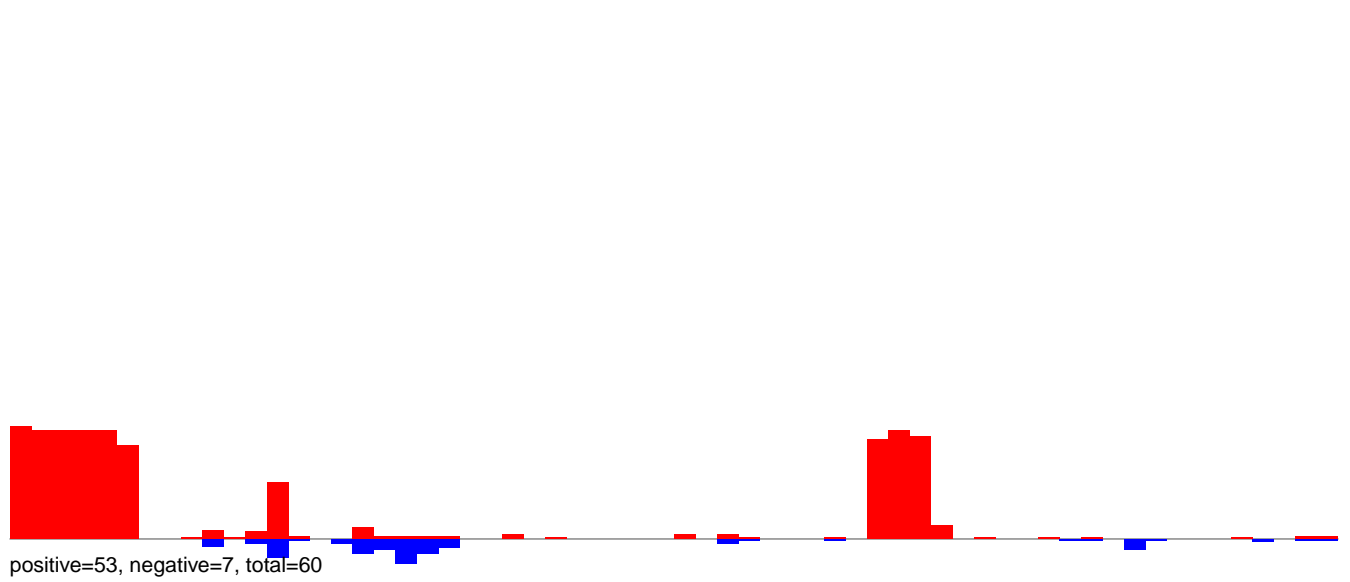
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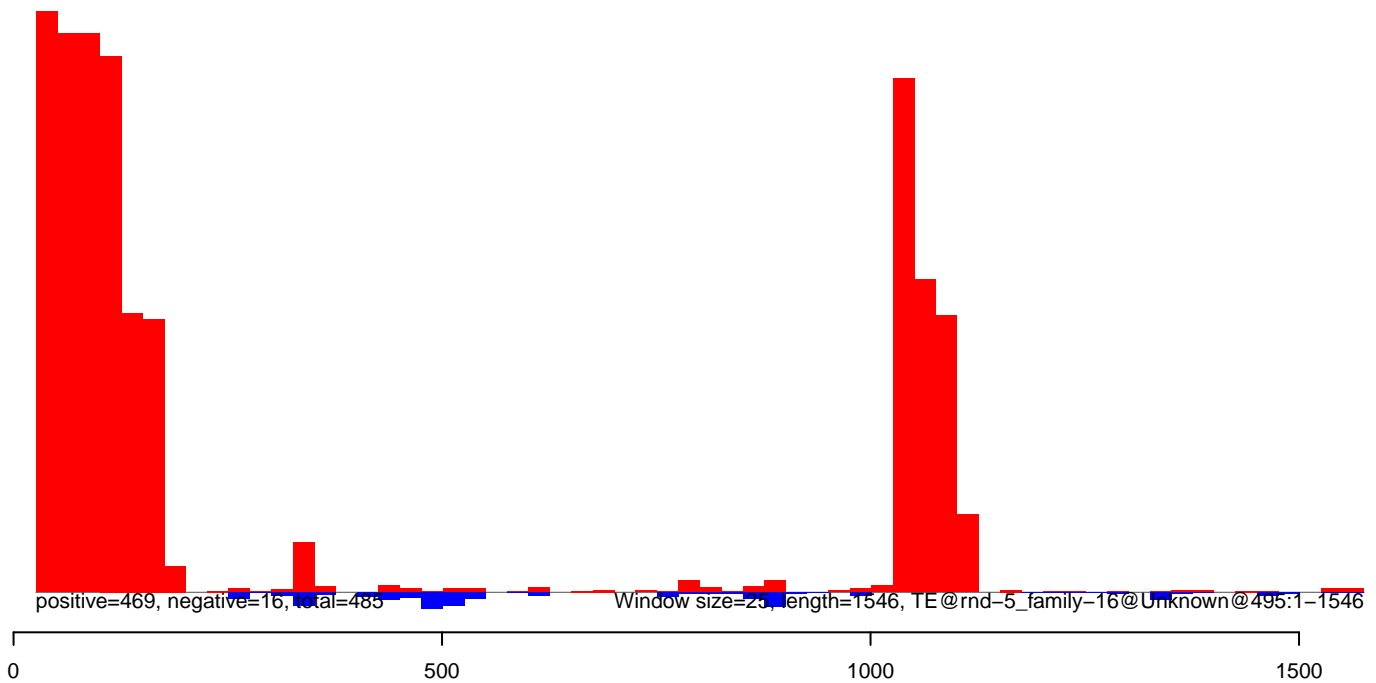
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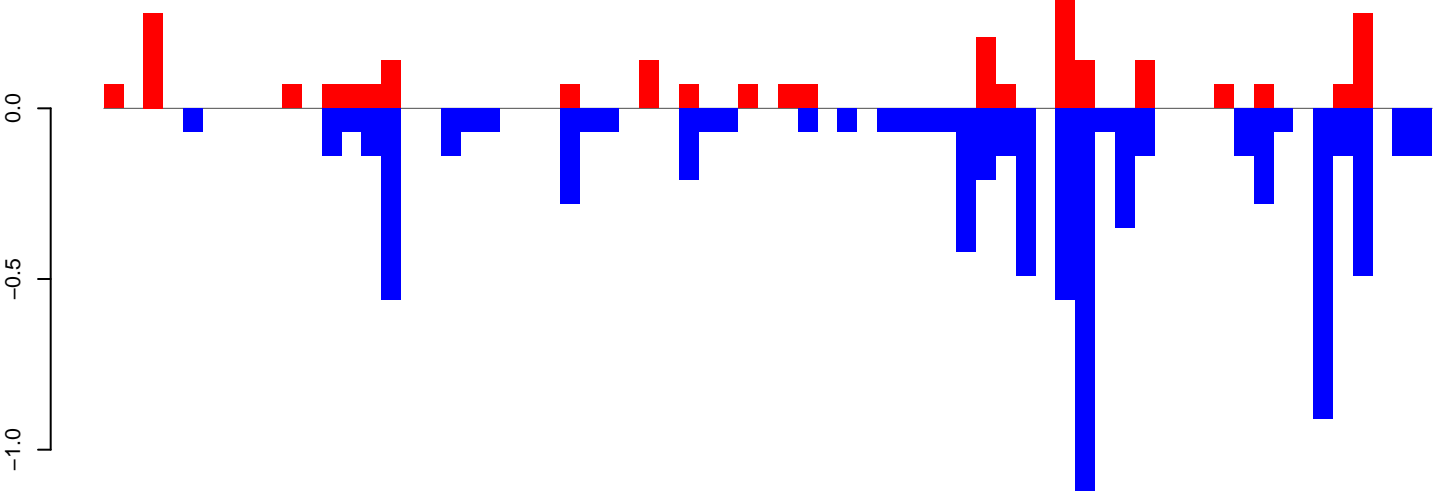
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CuQuin_Mcarc_NL.rep

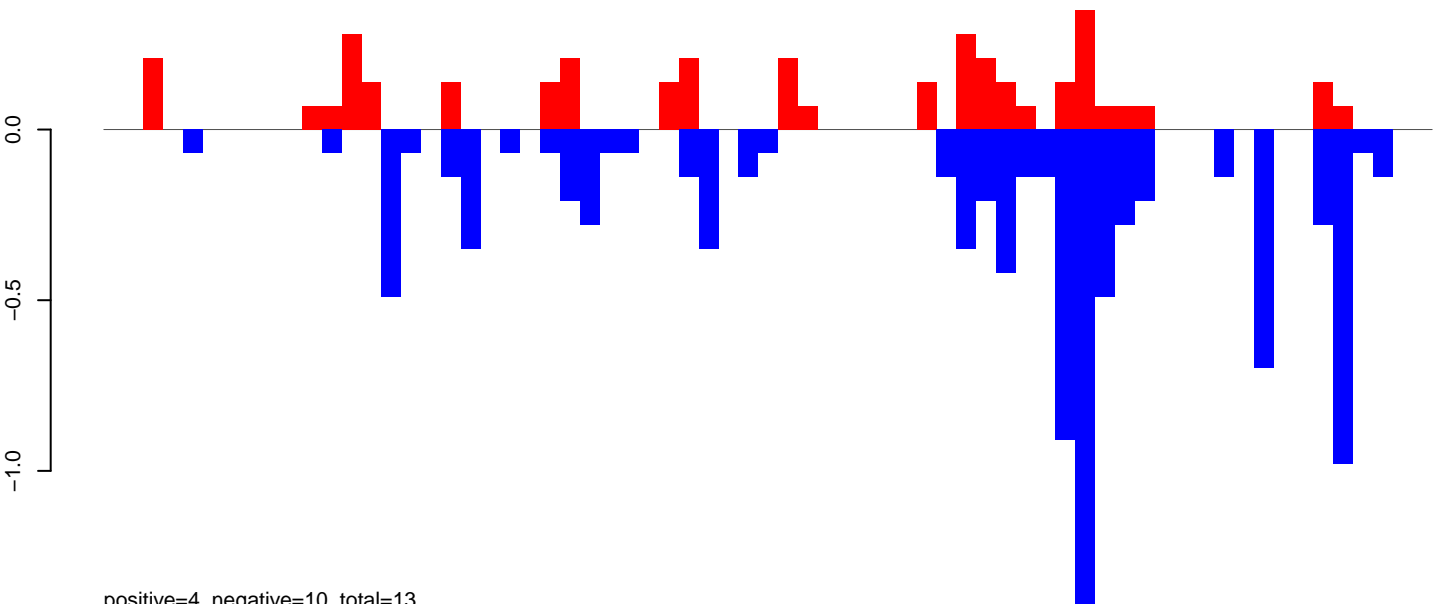


CuQuin_Mcarc_NL.18_23.rep



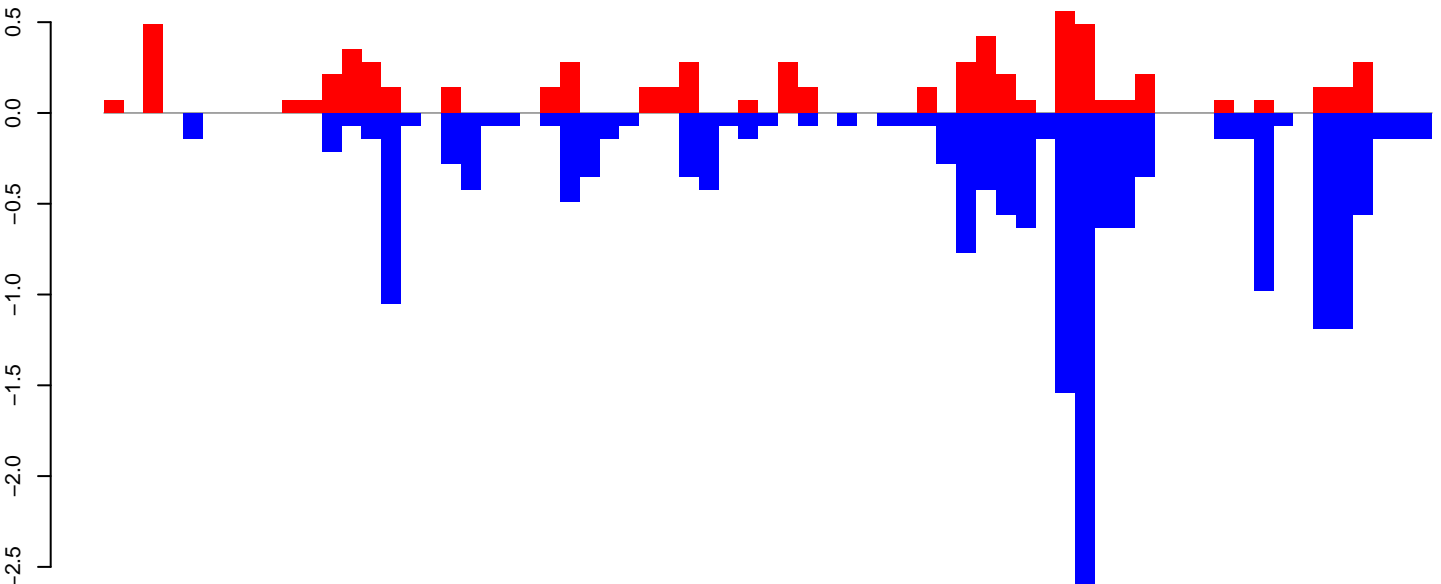
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CuQuin_Mcarc_NL.24_35.rep



positive=4, negative=10, total=13

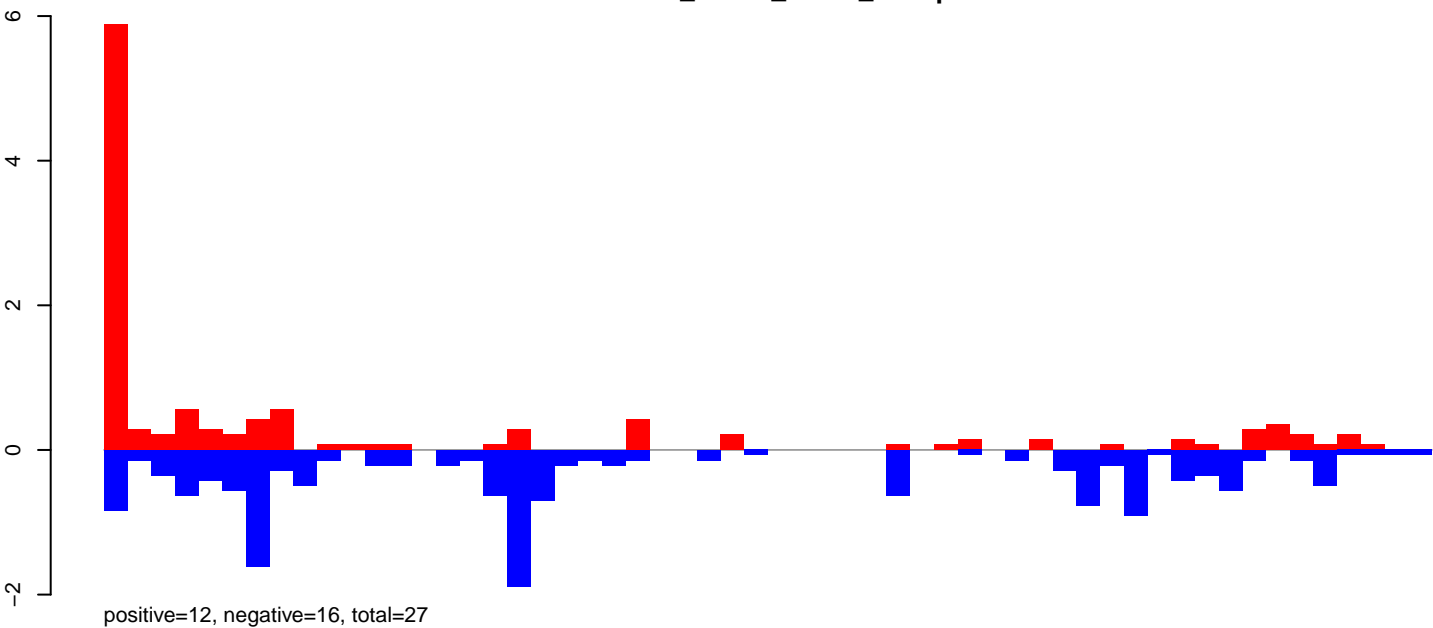
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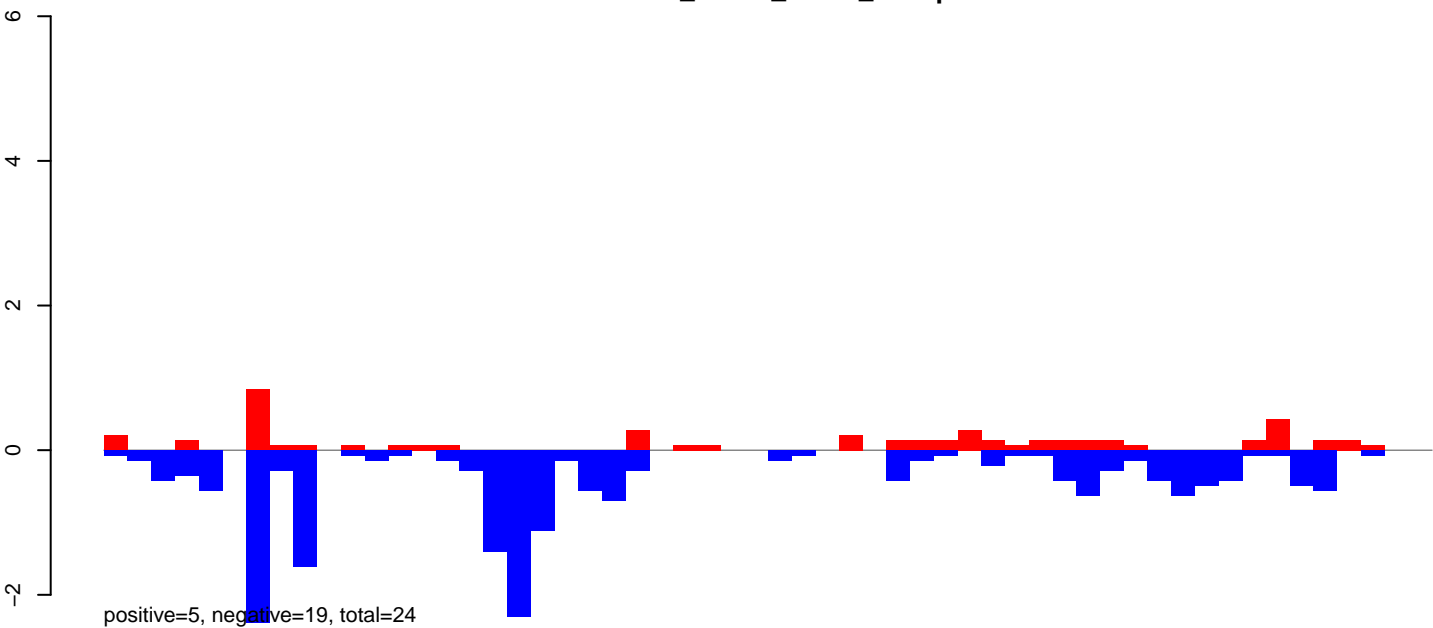
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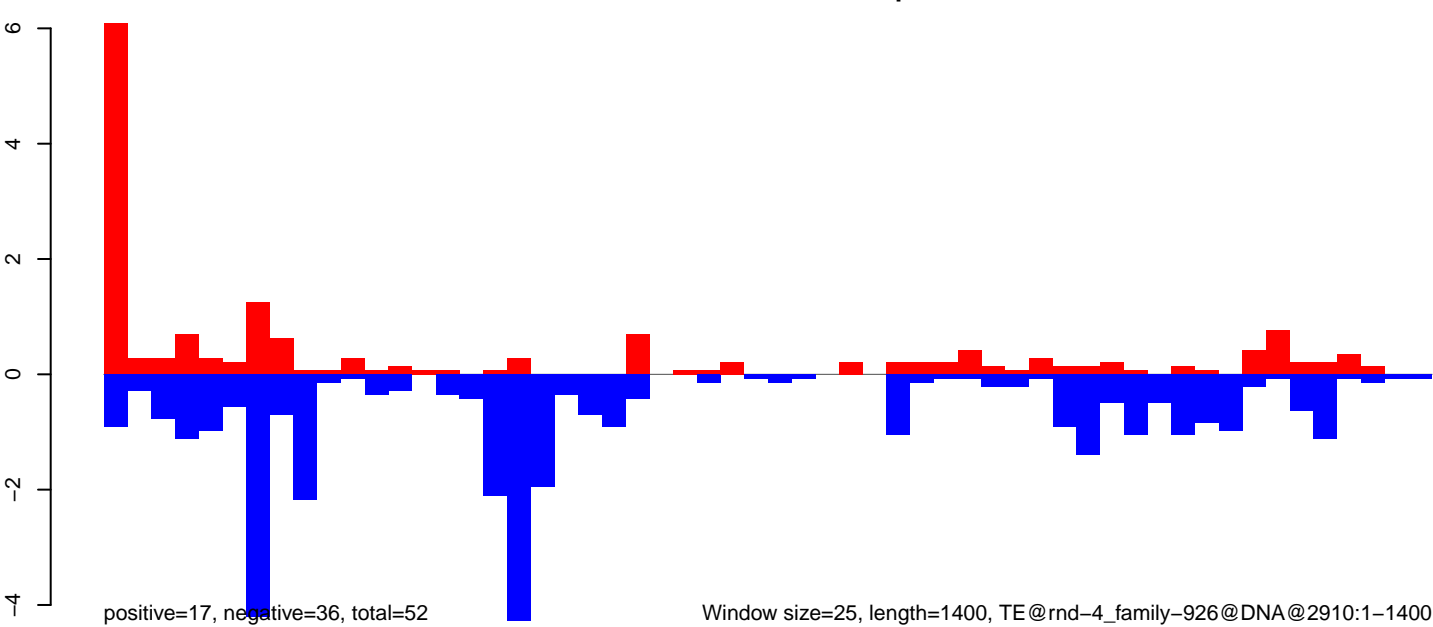
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

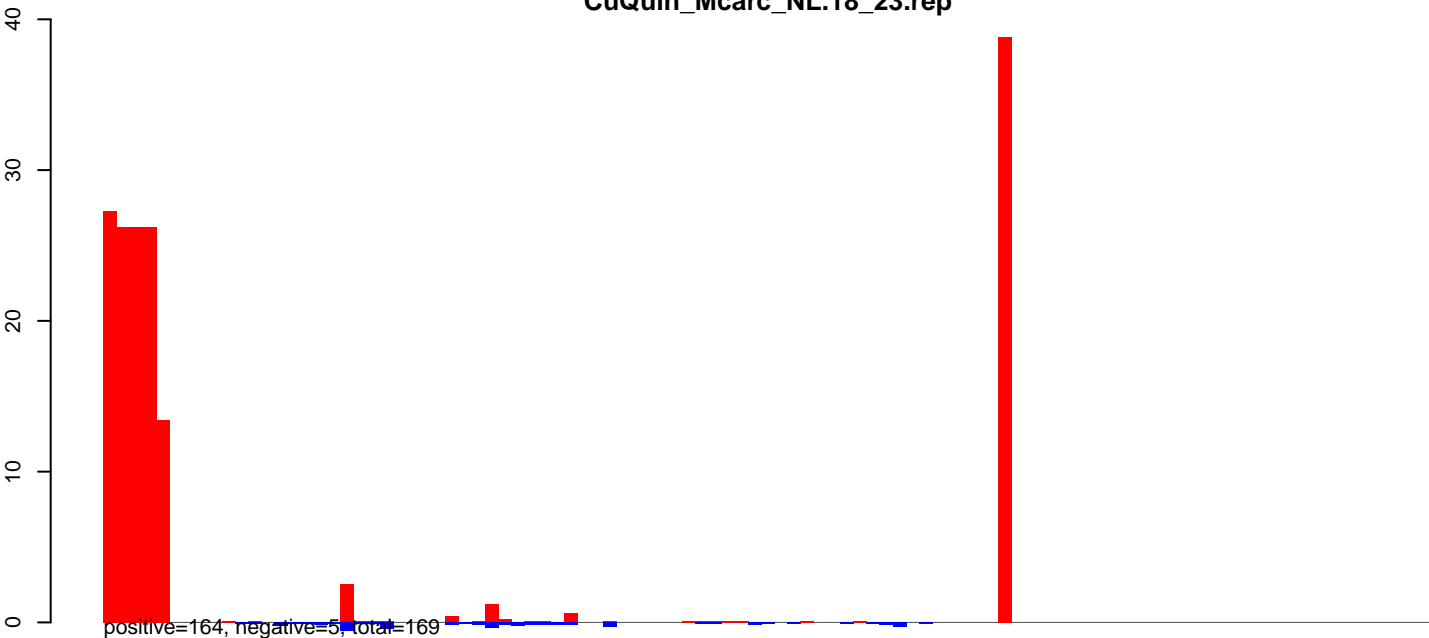


CuQuin_Mcarc_NL.rep

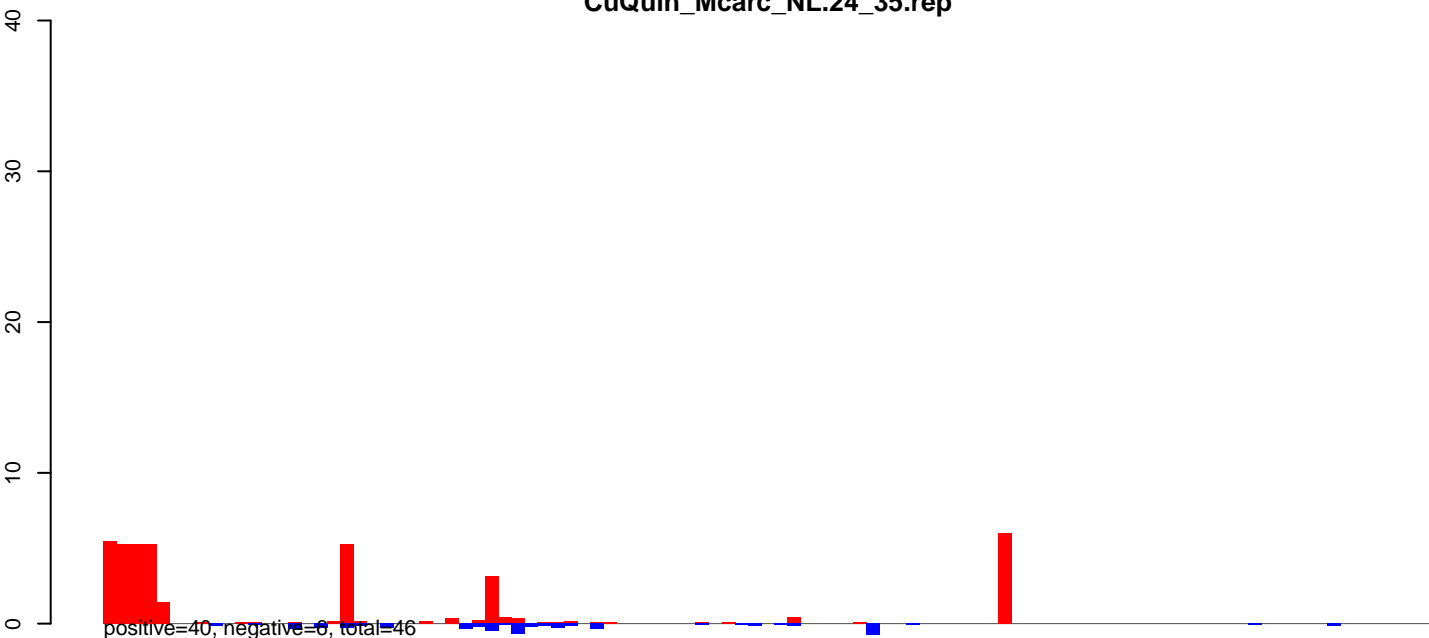


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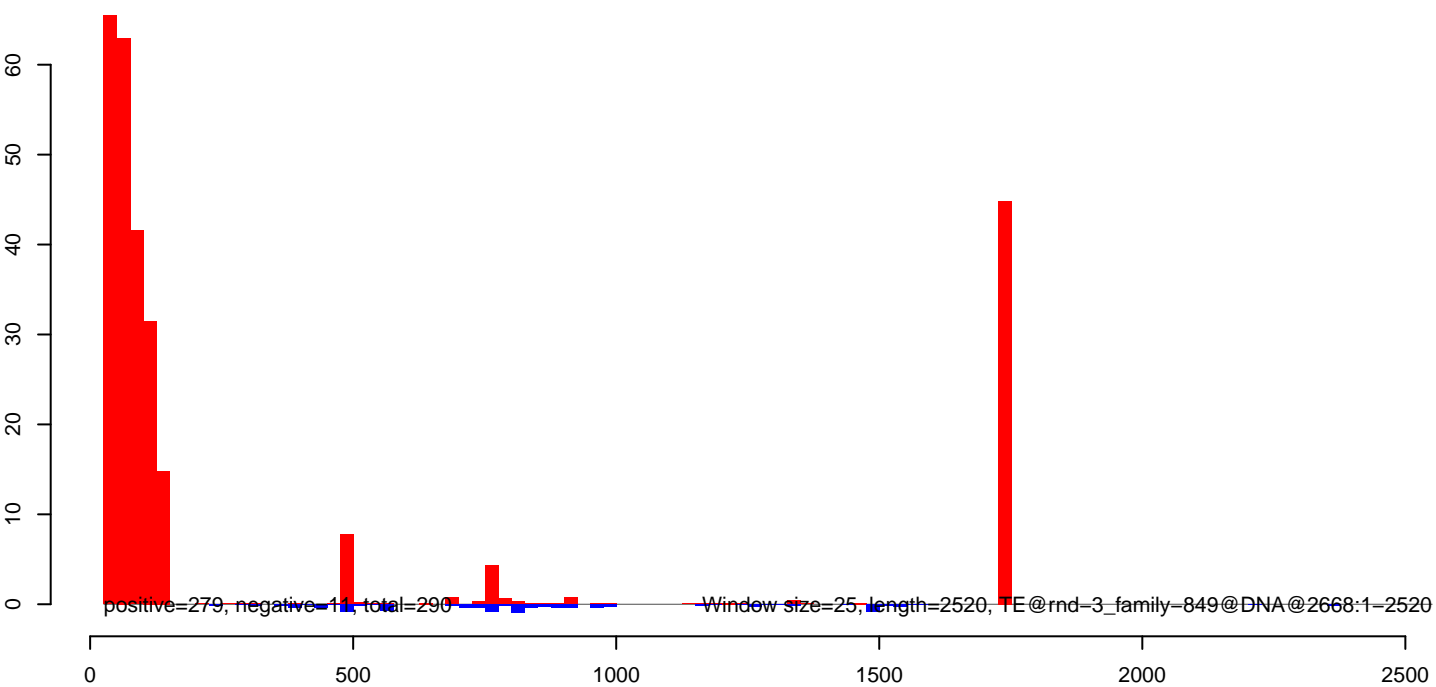
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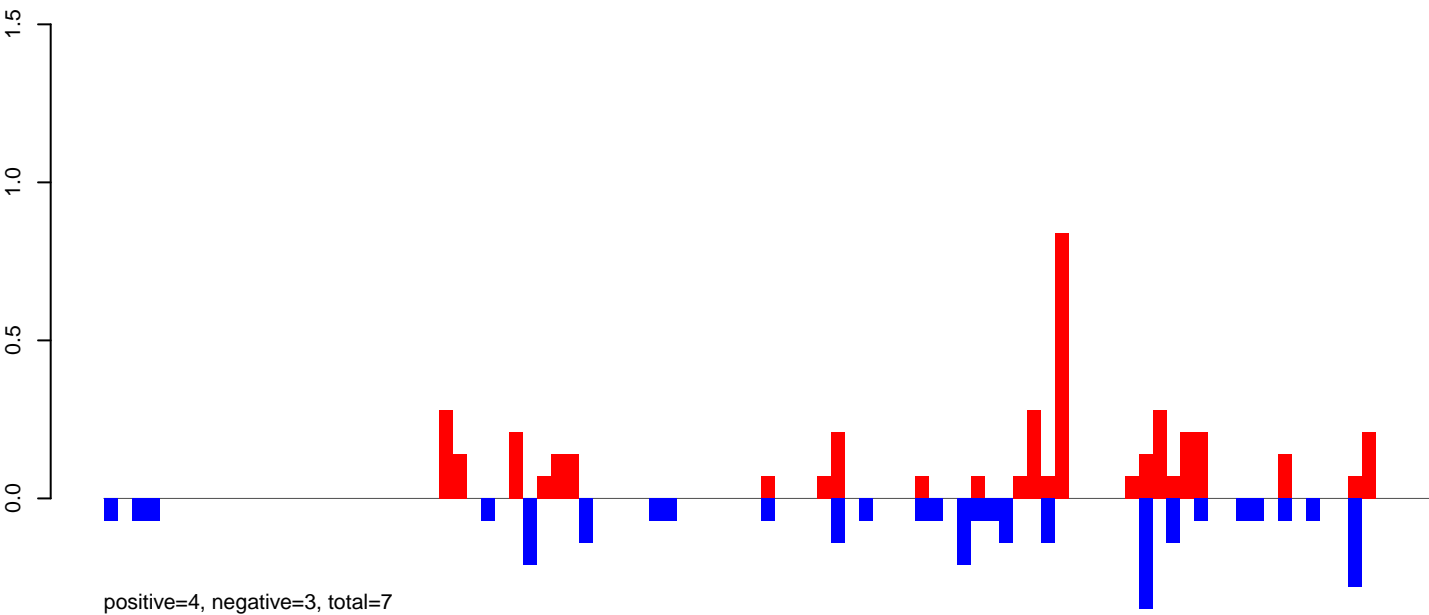
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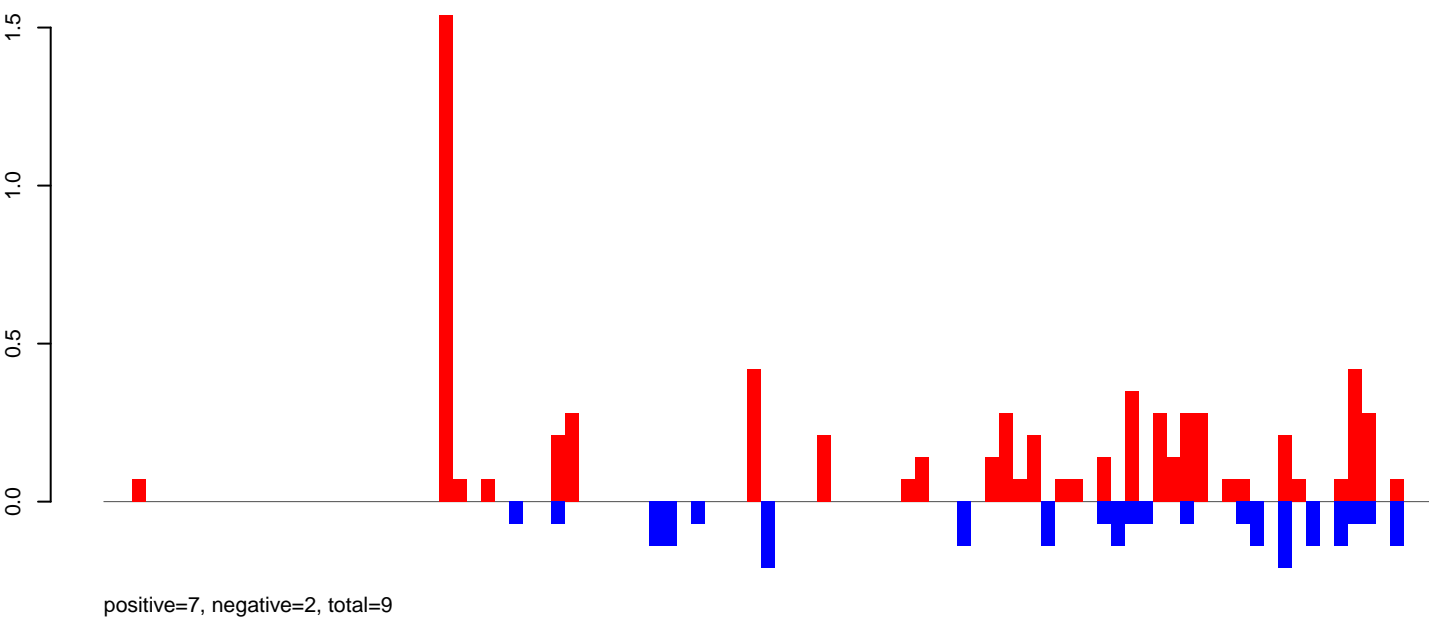
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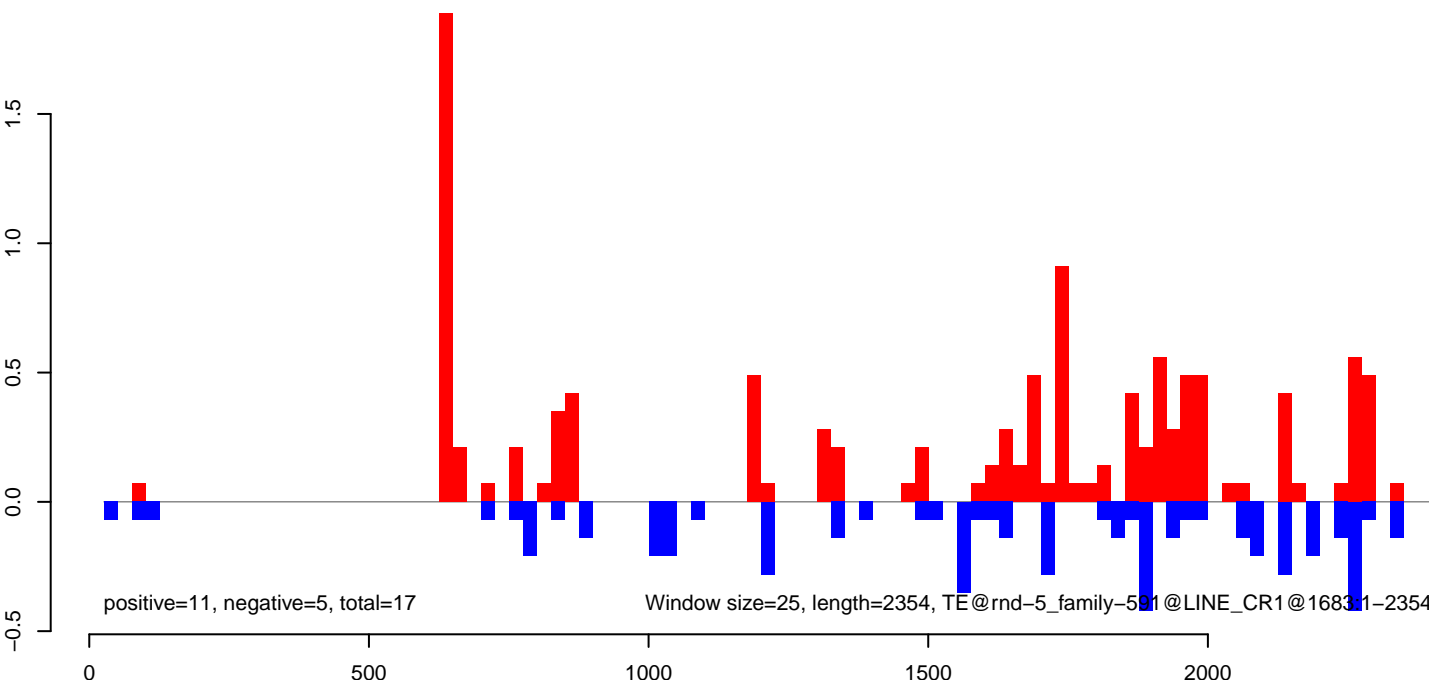
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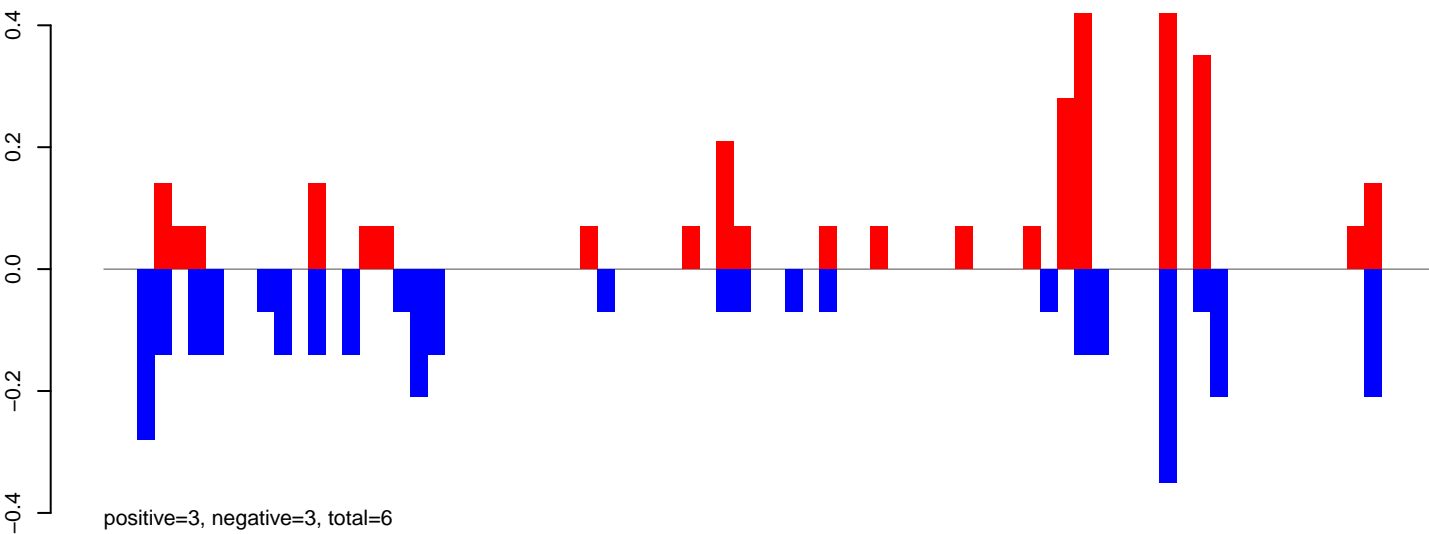
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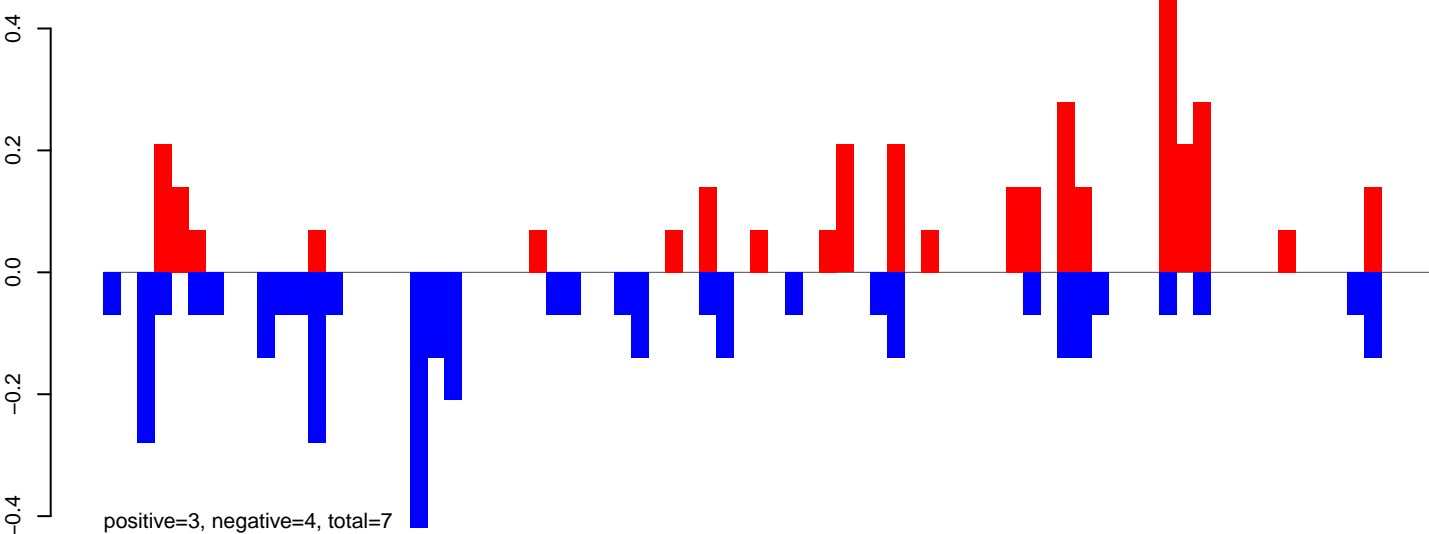
CuQuin_Mcarc_NL.rep



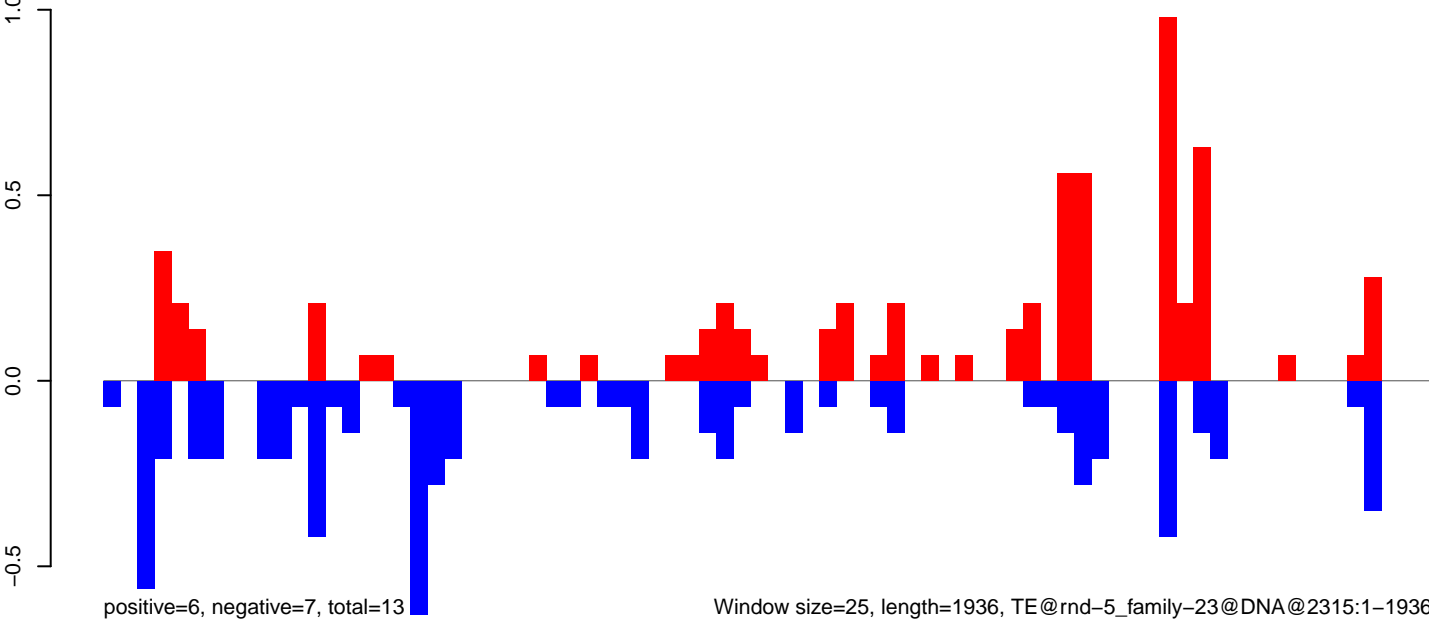
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CuQuin_Mcarc_NL.24_35.rep



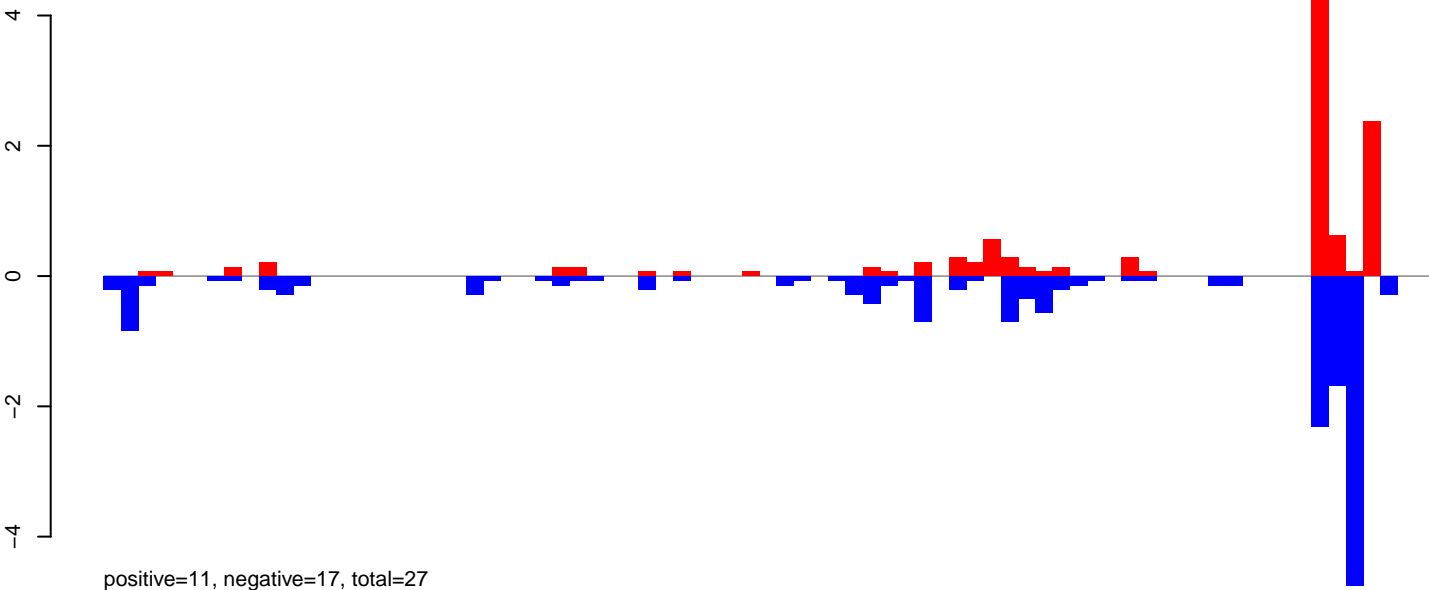
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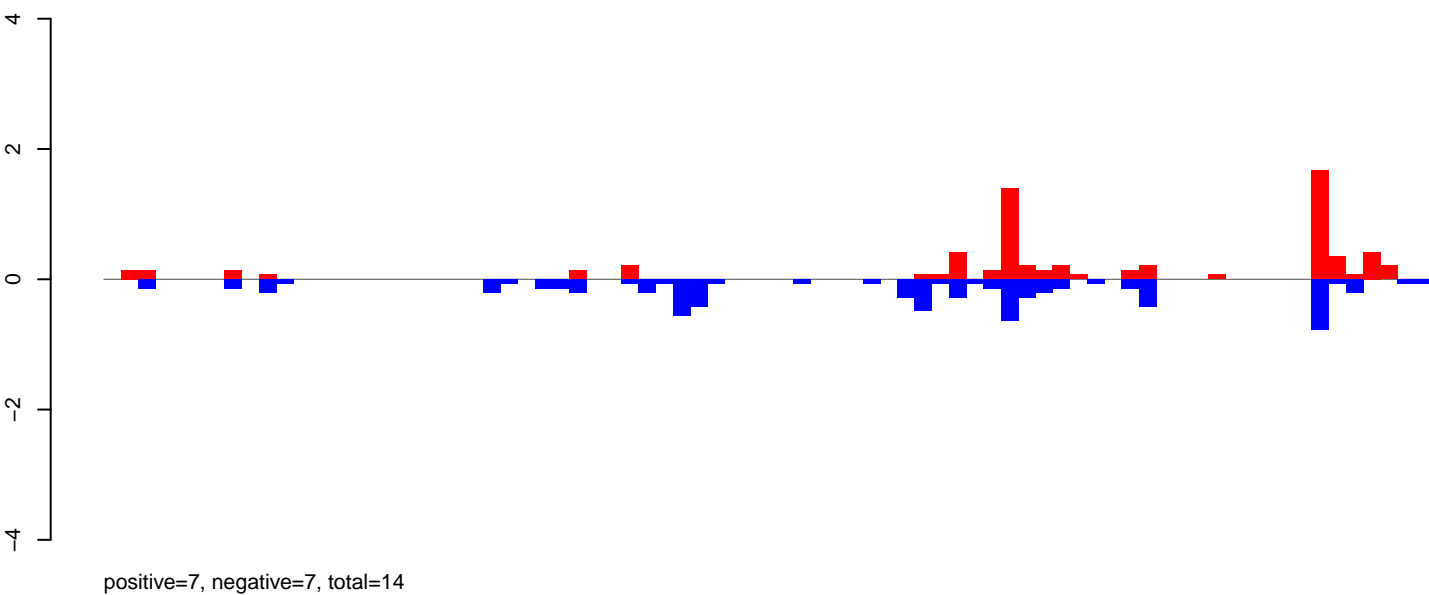
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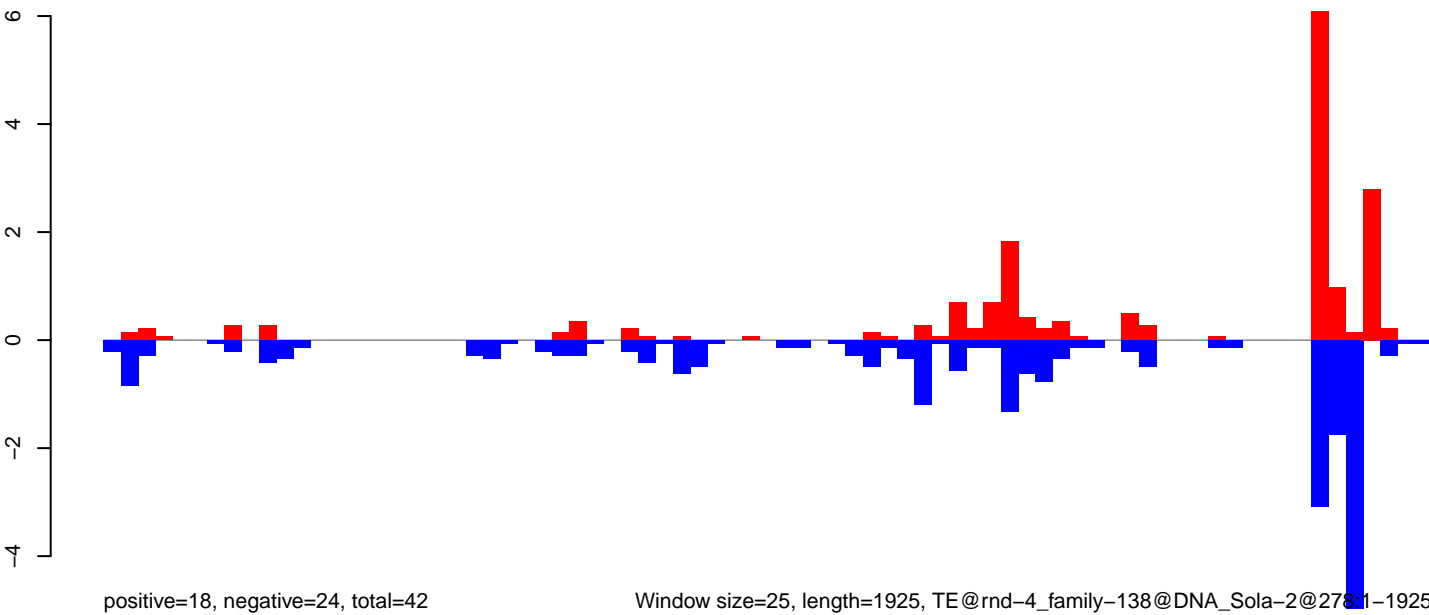
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CuQuin_Mcarc_NL.24_35.rep

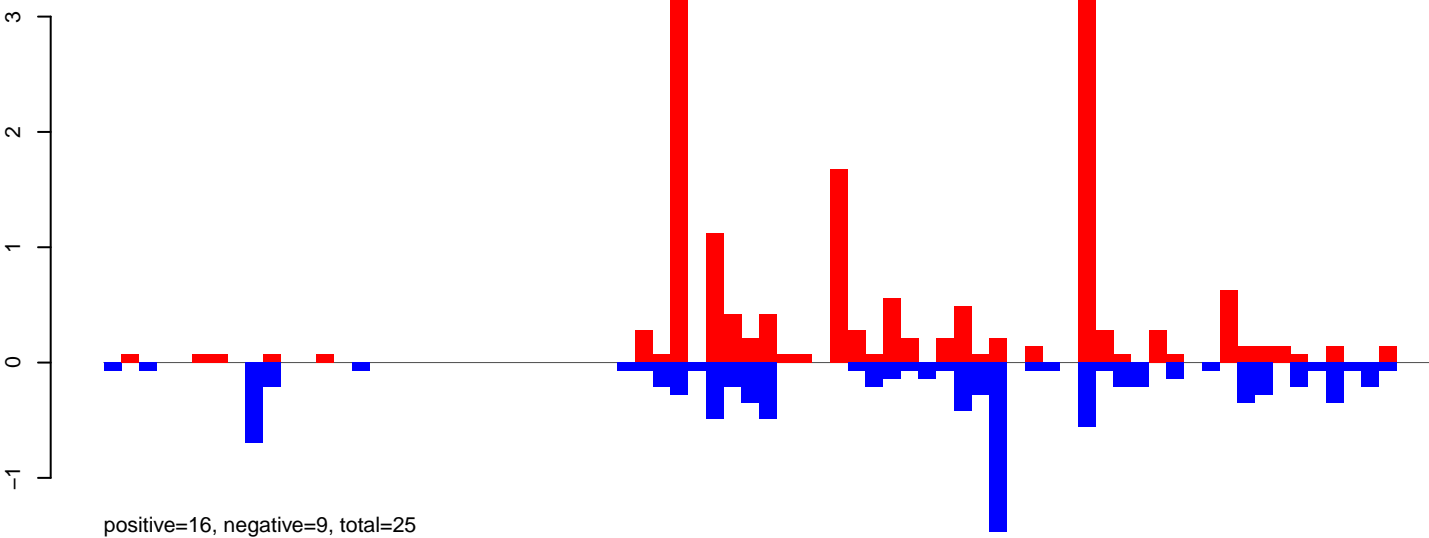


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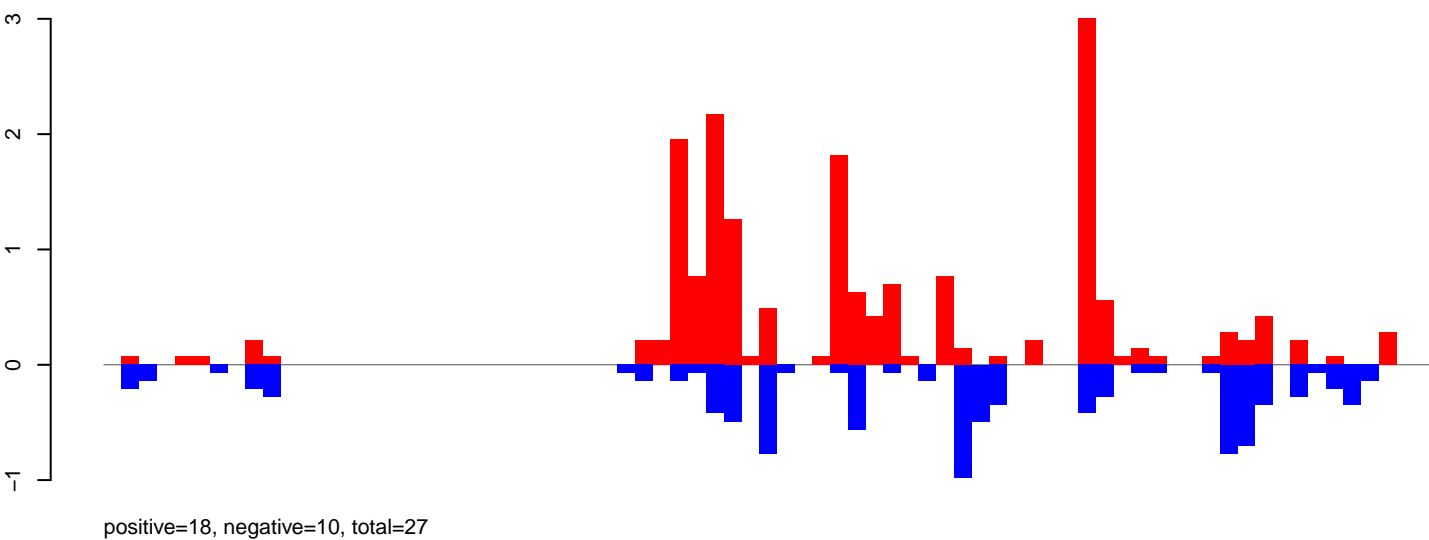


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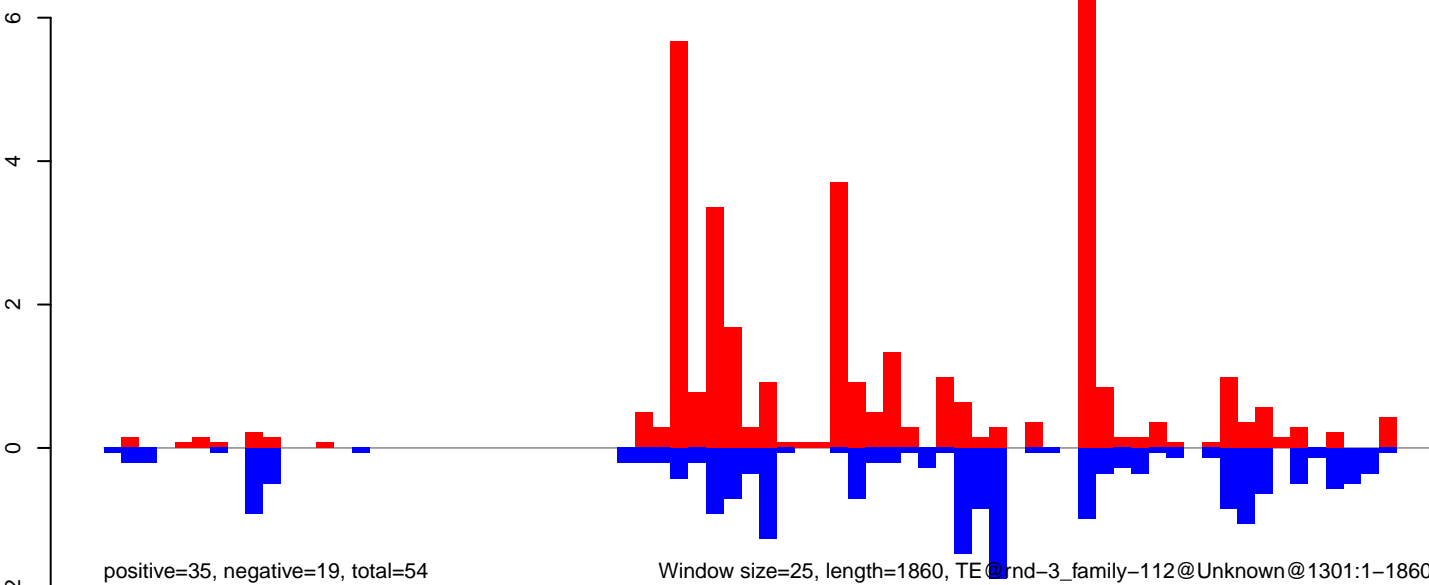
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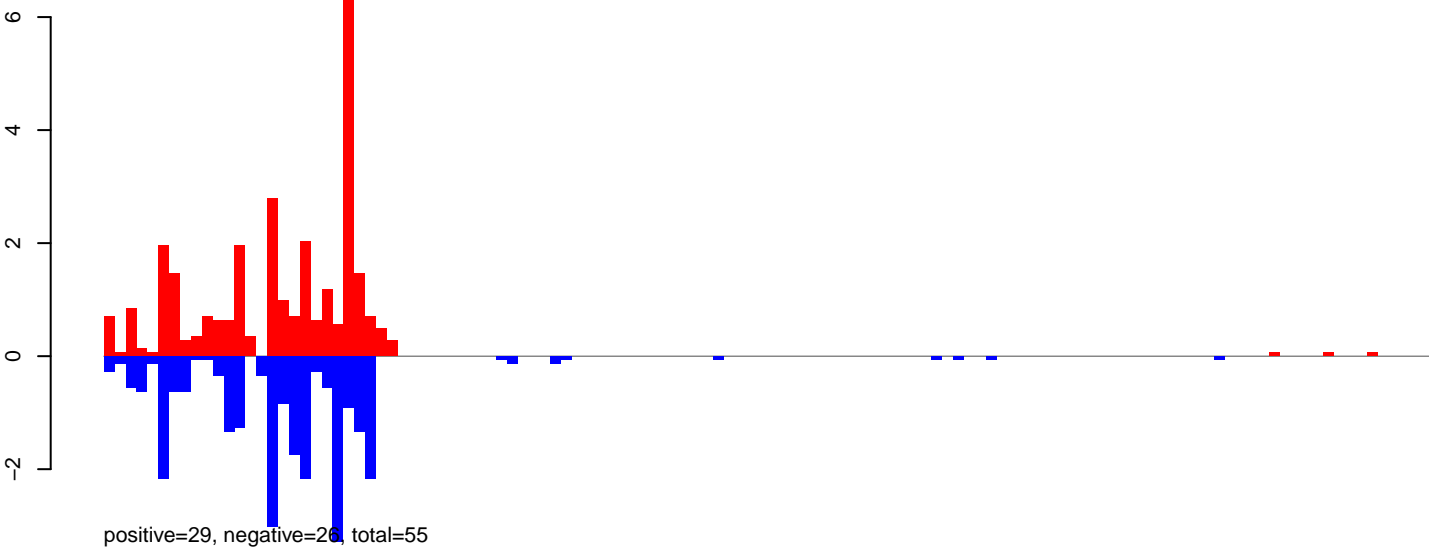
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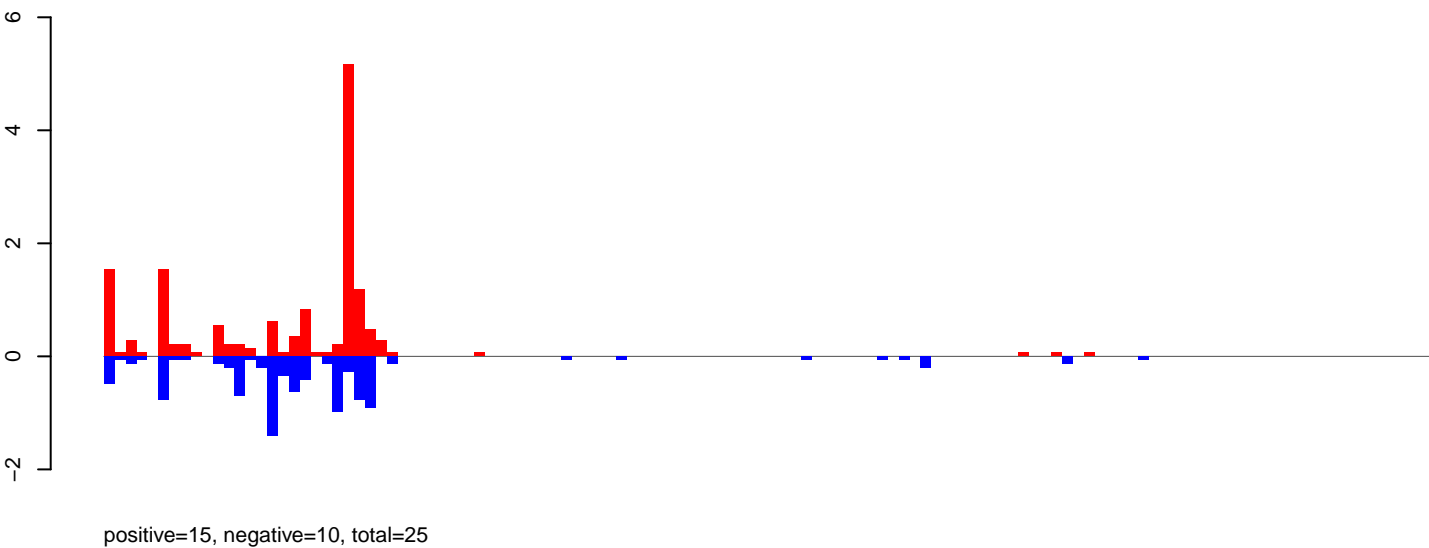
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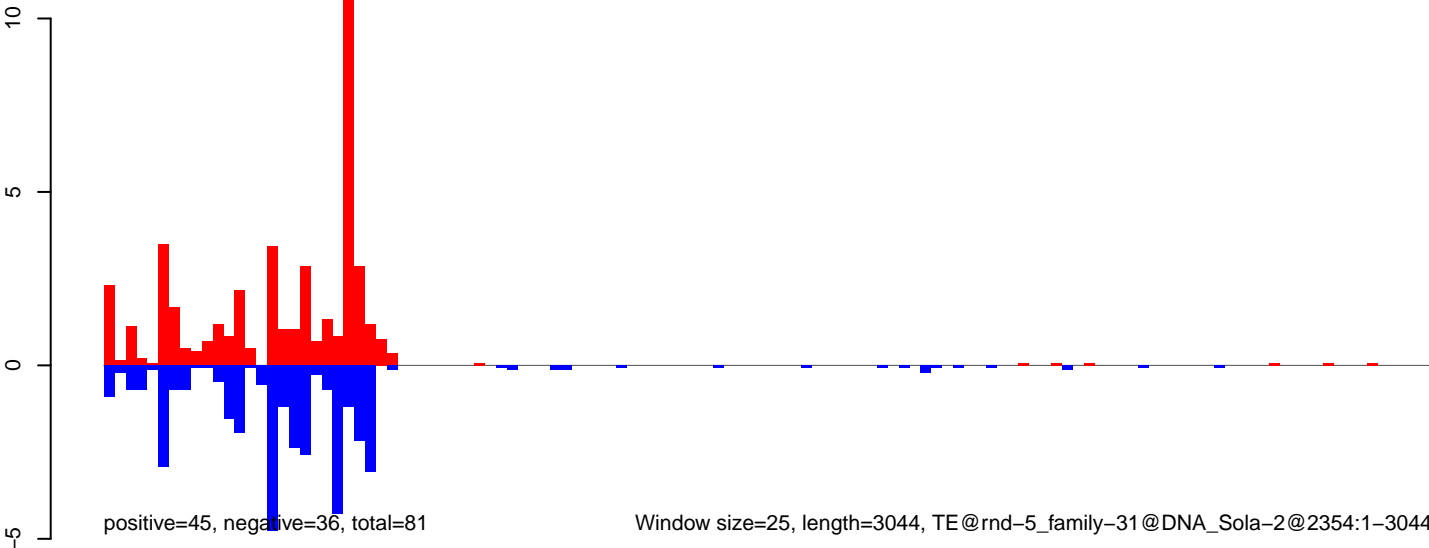
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CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



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CuQuin_Mcarc_NL.24_35.rep

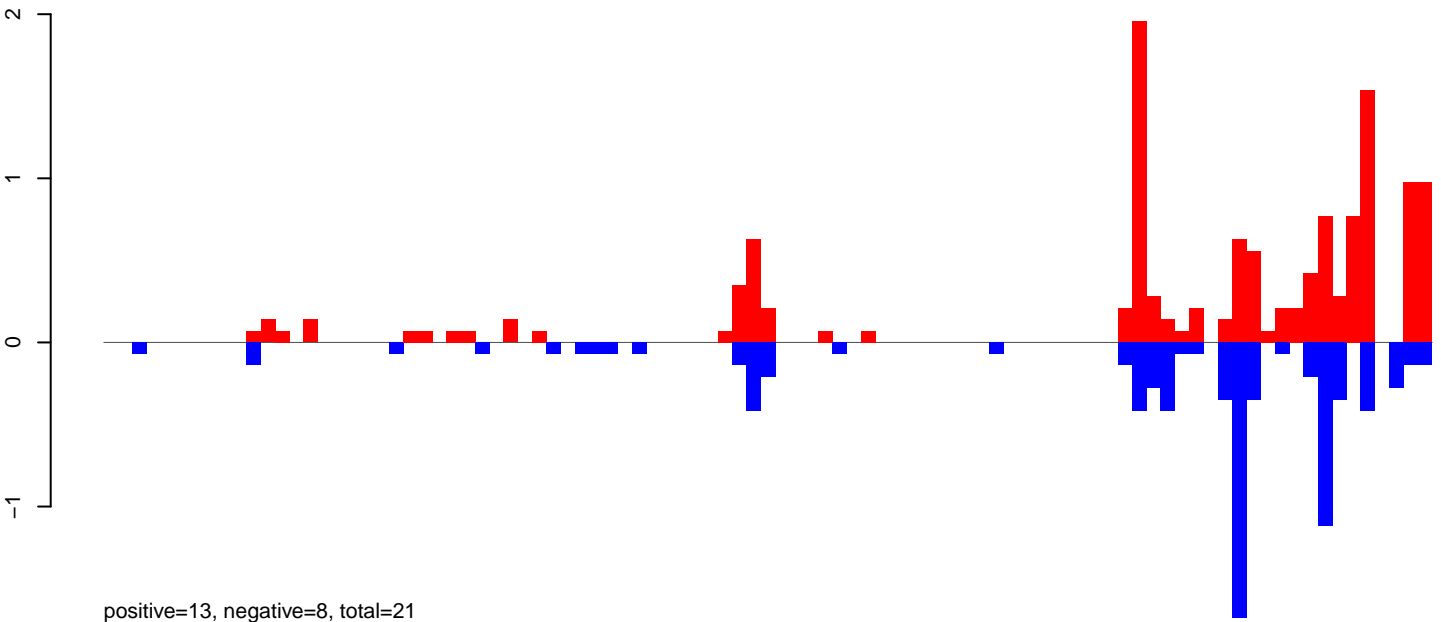


CuQuin_Mcarc_NL.rep

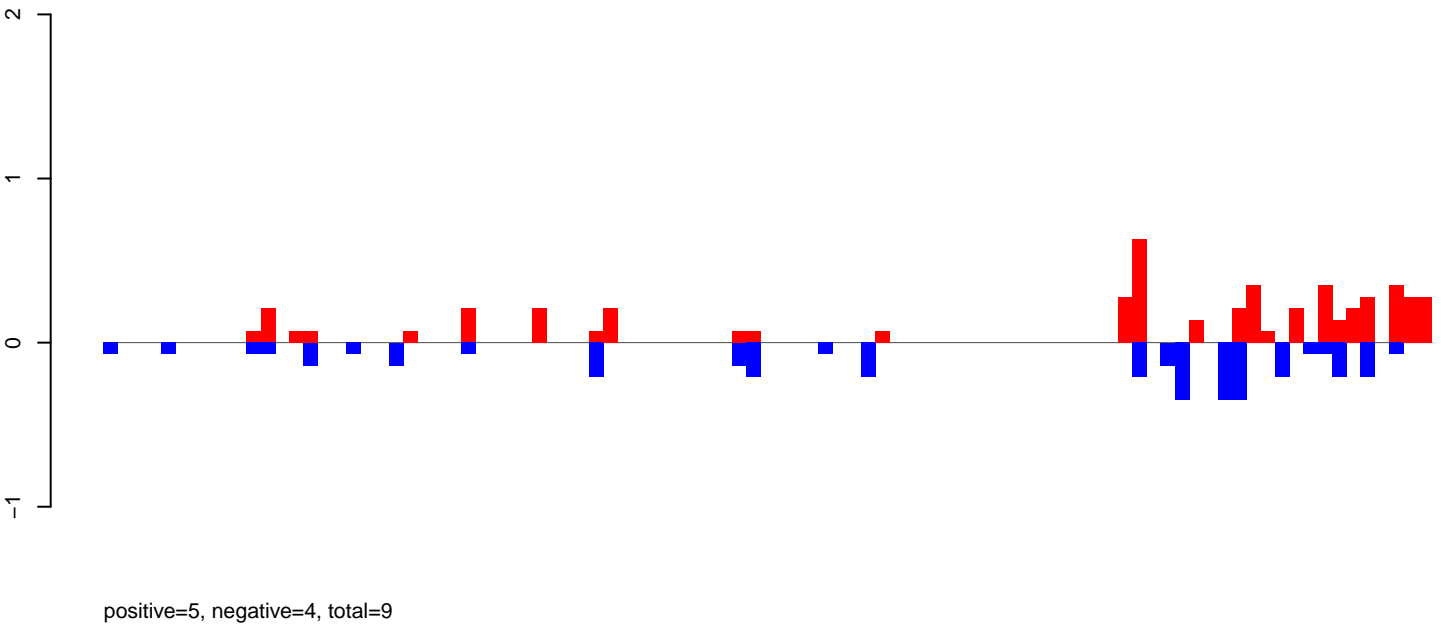


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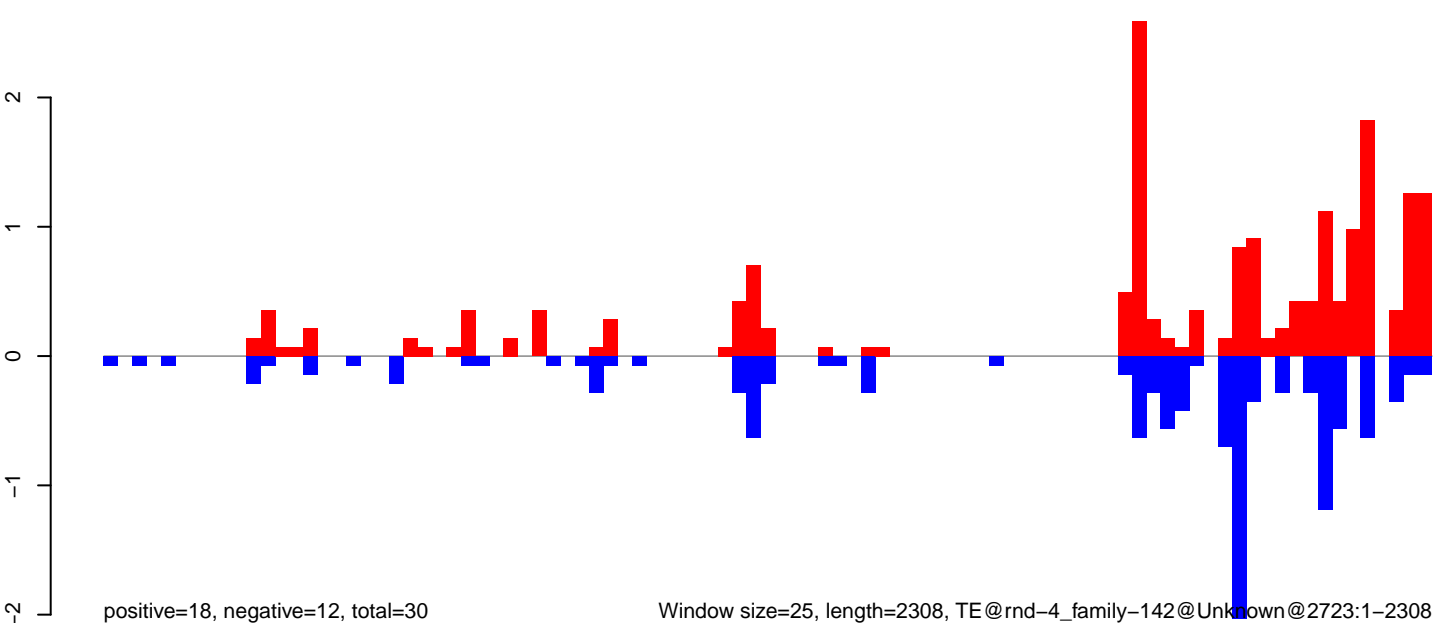
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CuQuin_Mcarc_NL.24_35.rep

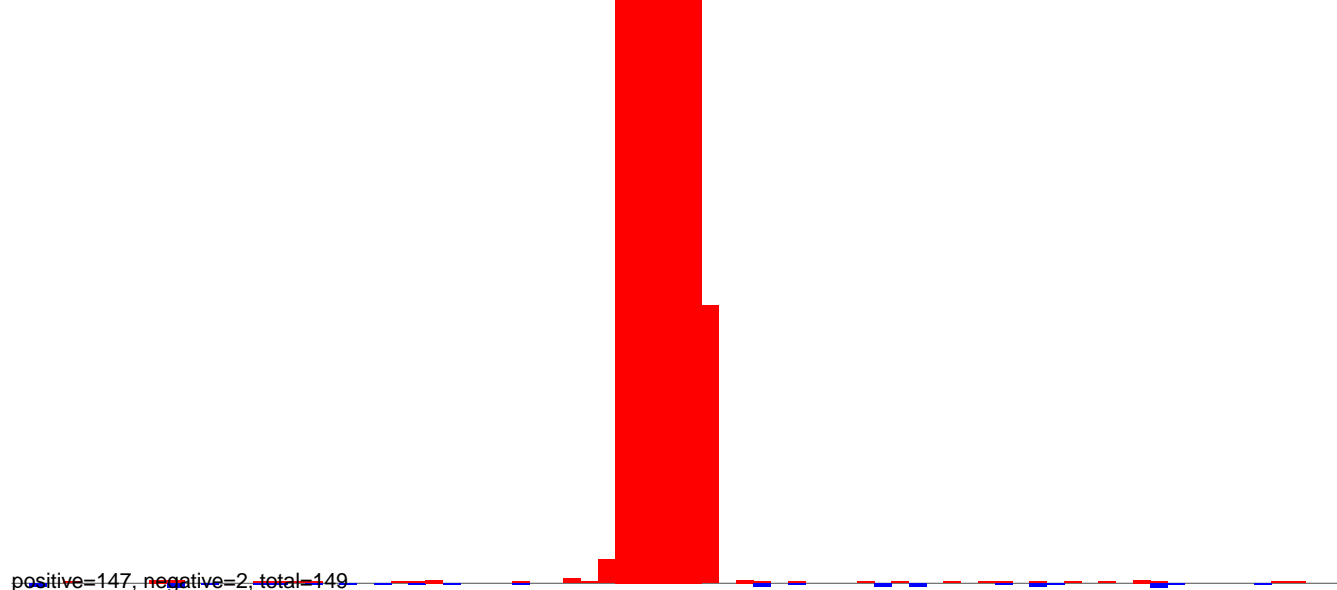


CuQuin_Mcarc_NL.rep

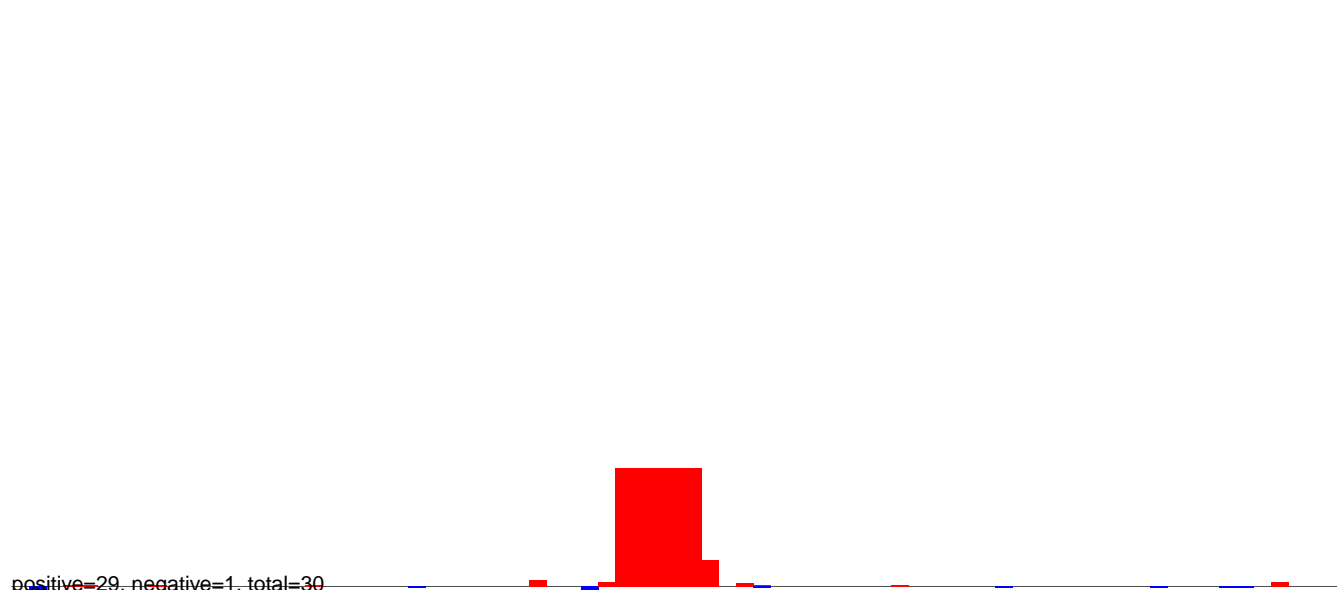


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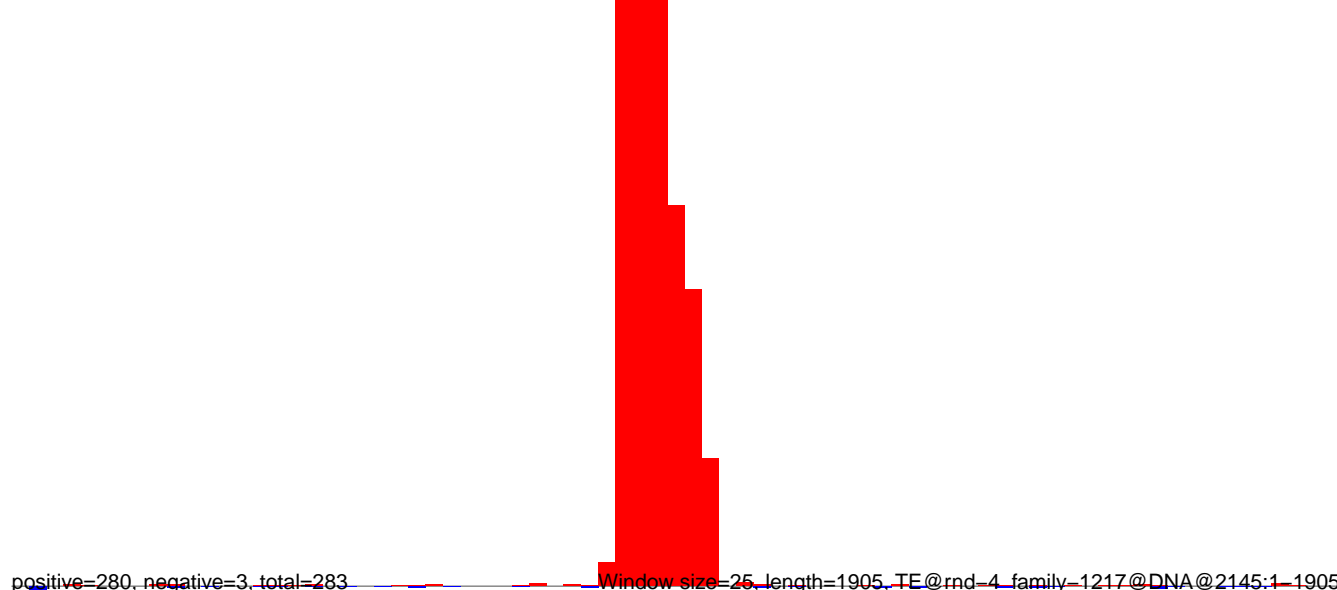
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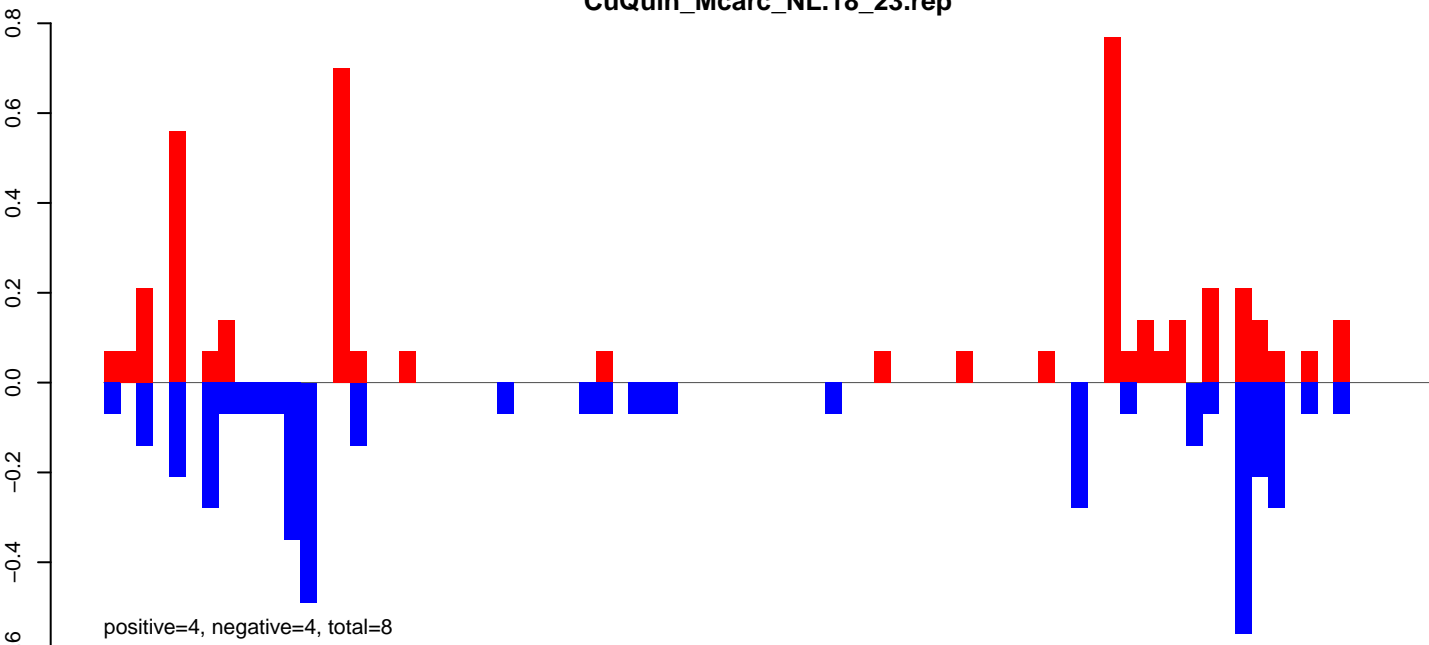
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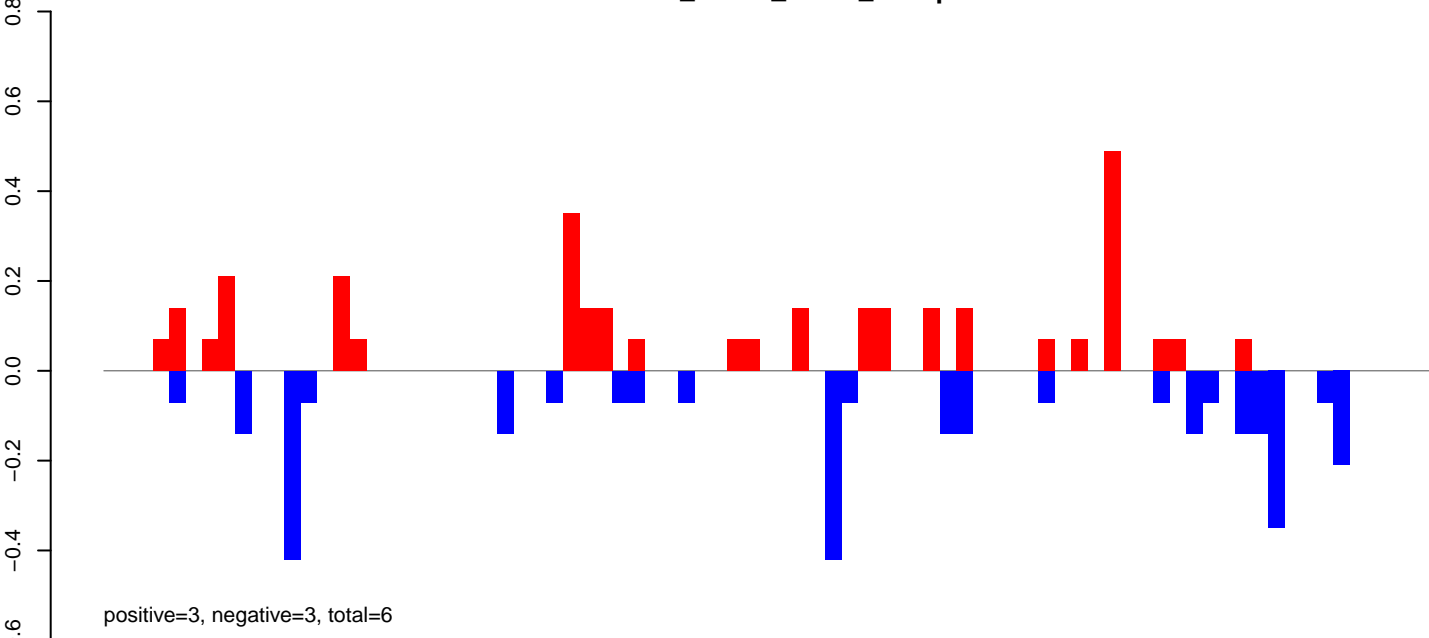
CuQuin_Mcarc_NL.rep



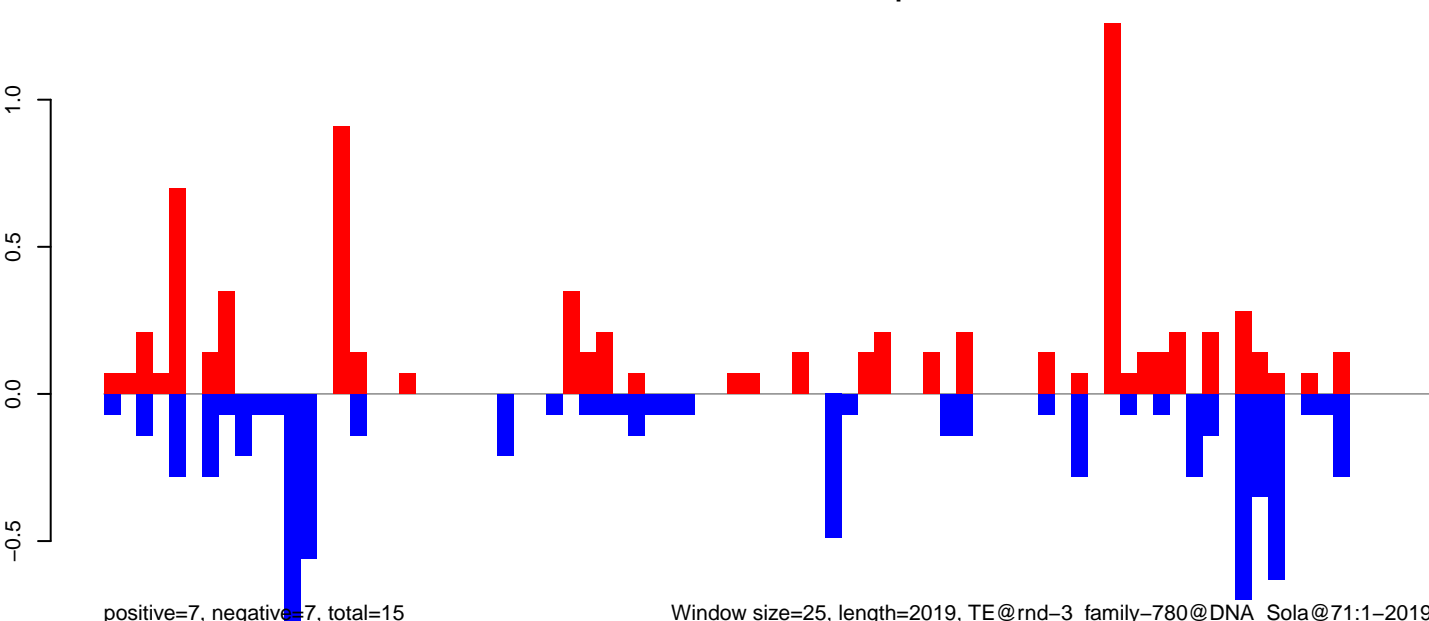
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CuQuin_Mcarc_NL.24_35.rep



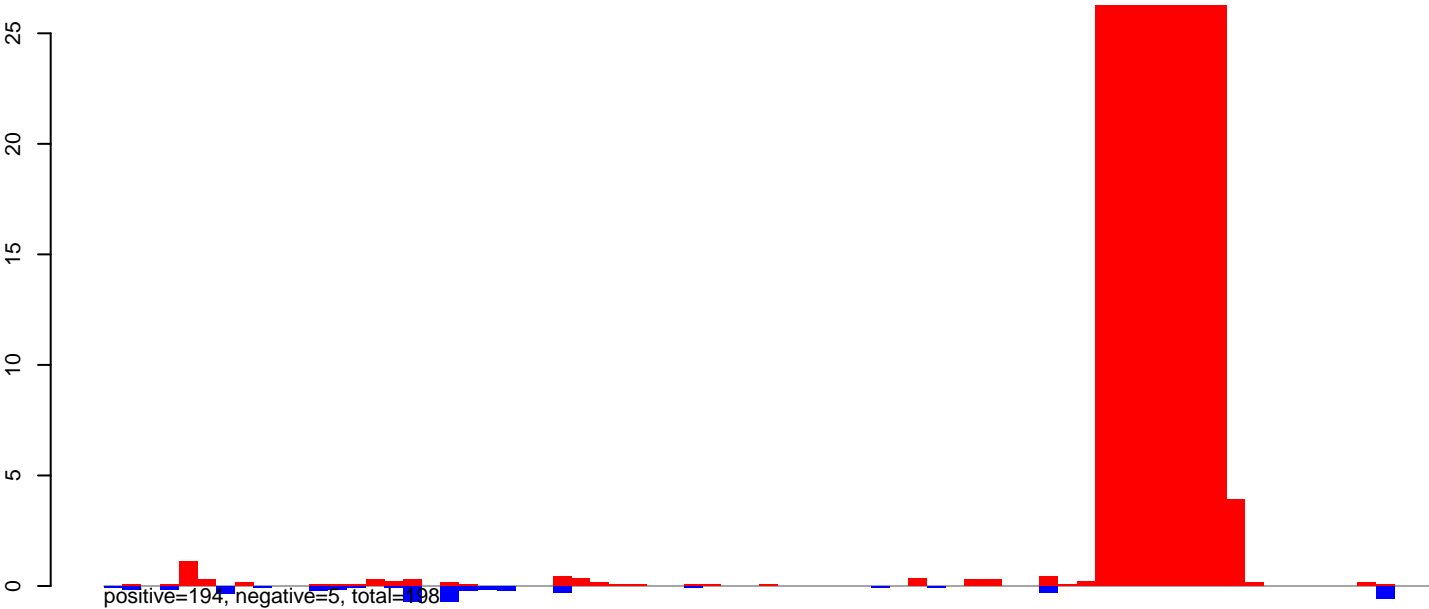
CuQuin_Mcarc_NL.rep



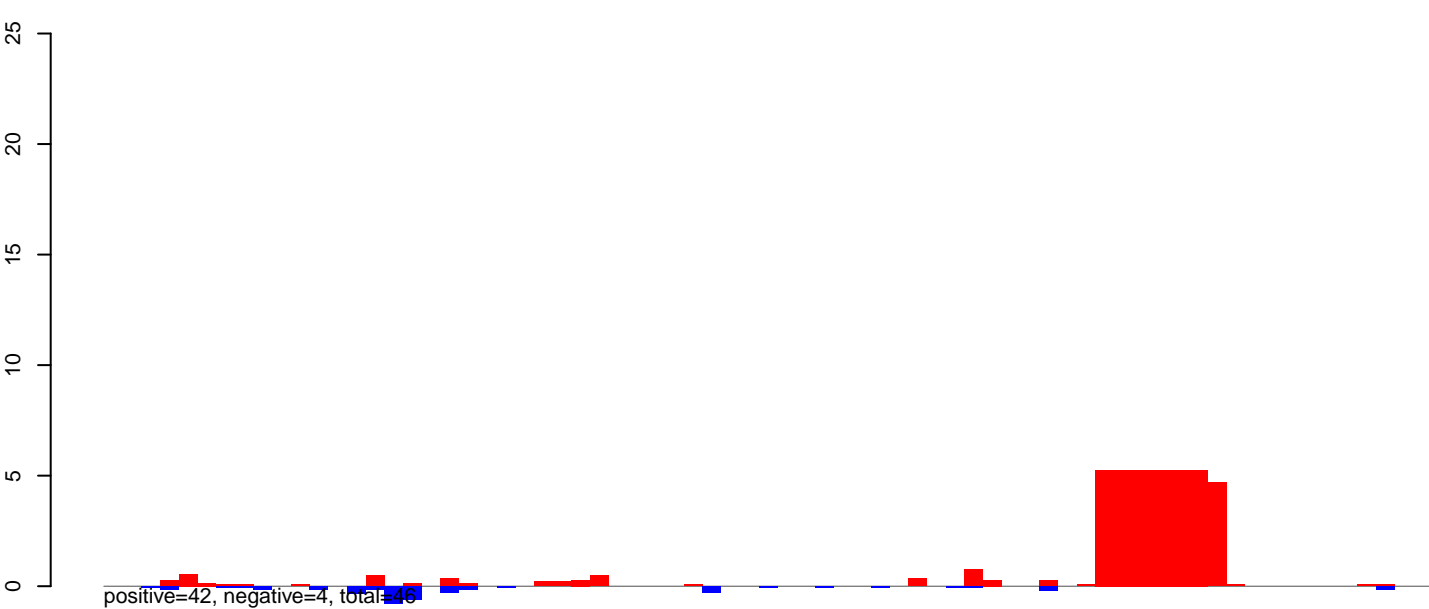
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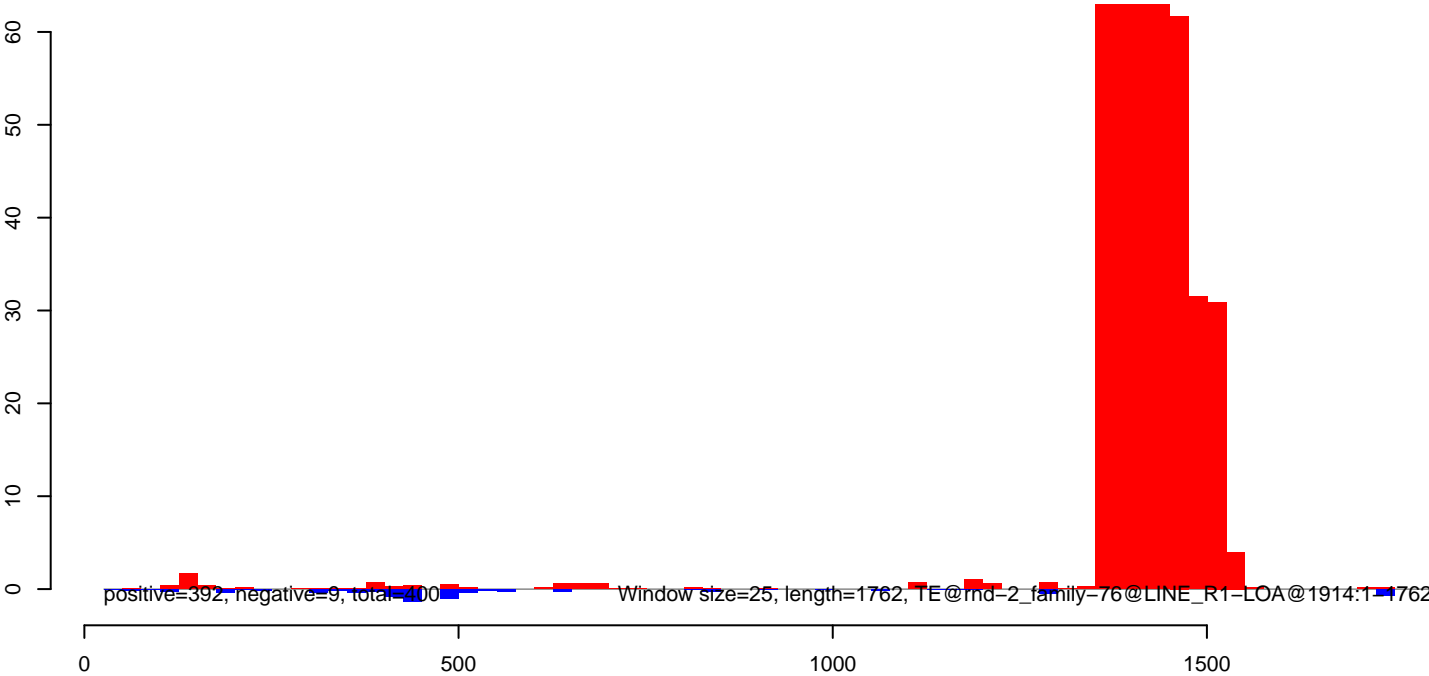
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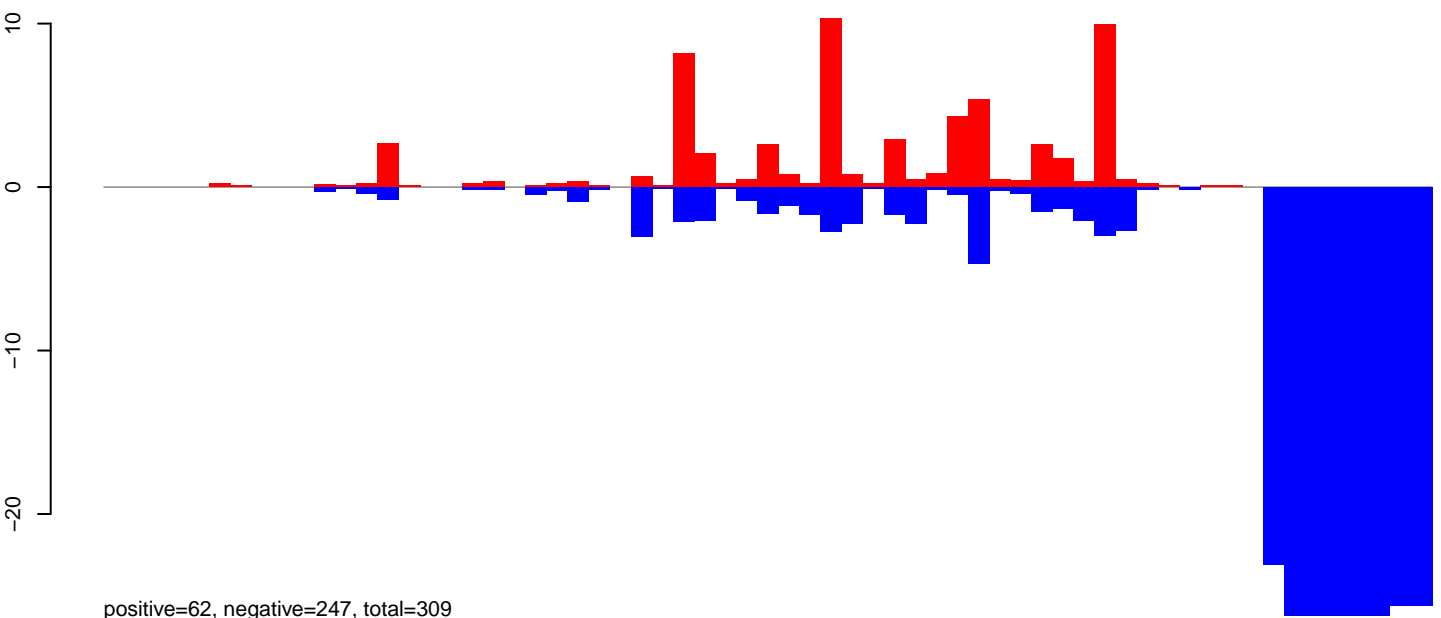
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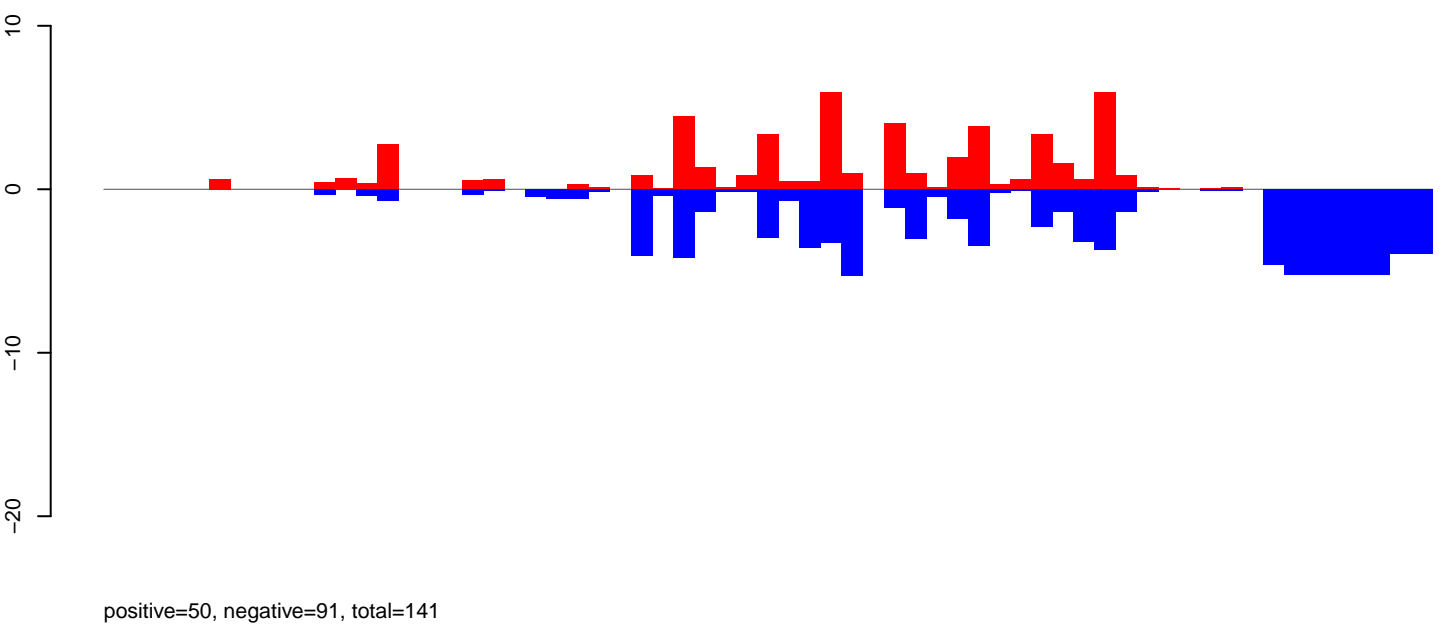
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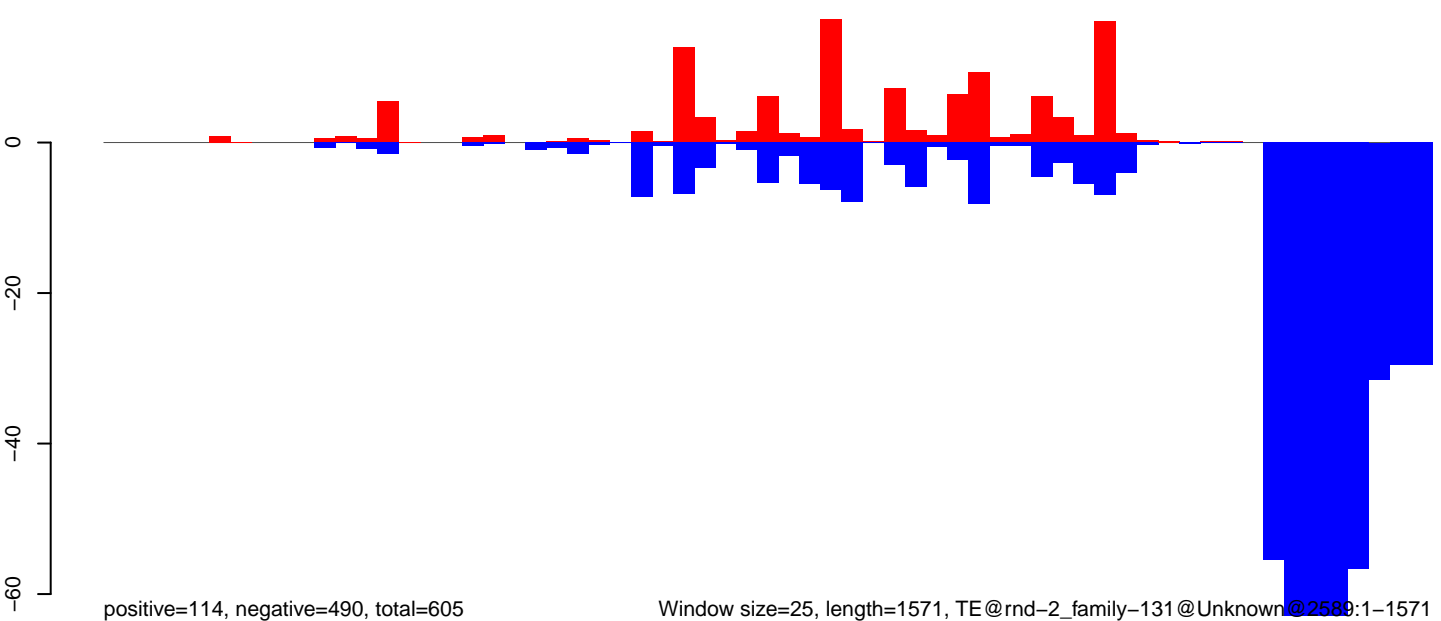
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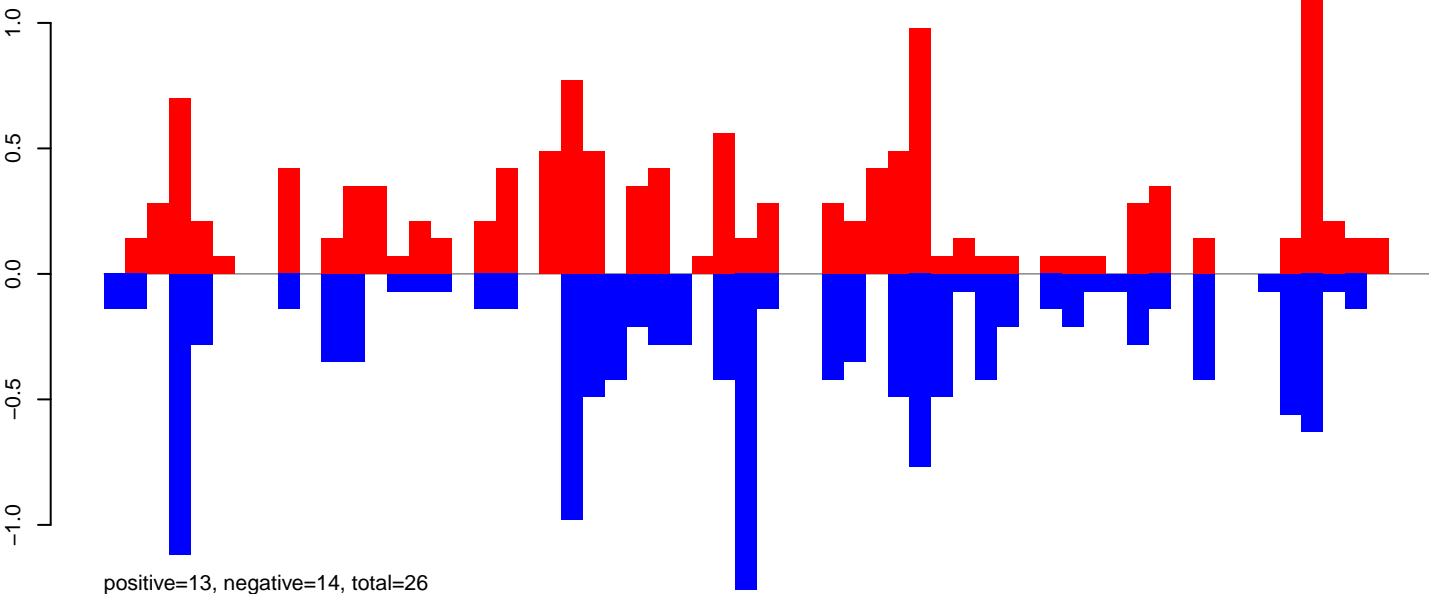
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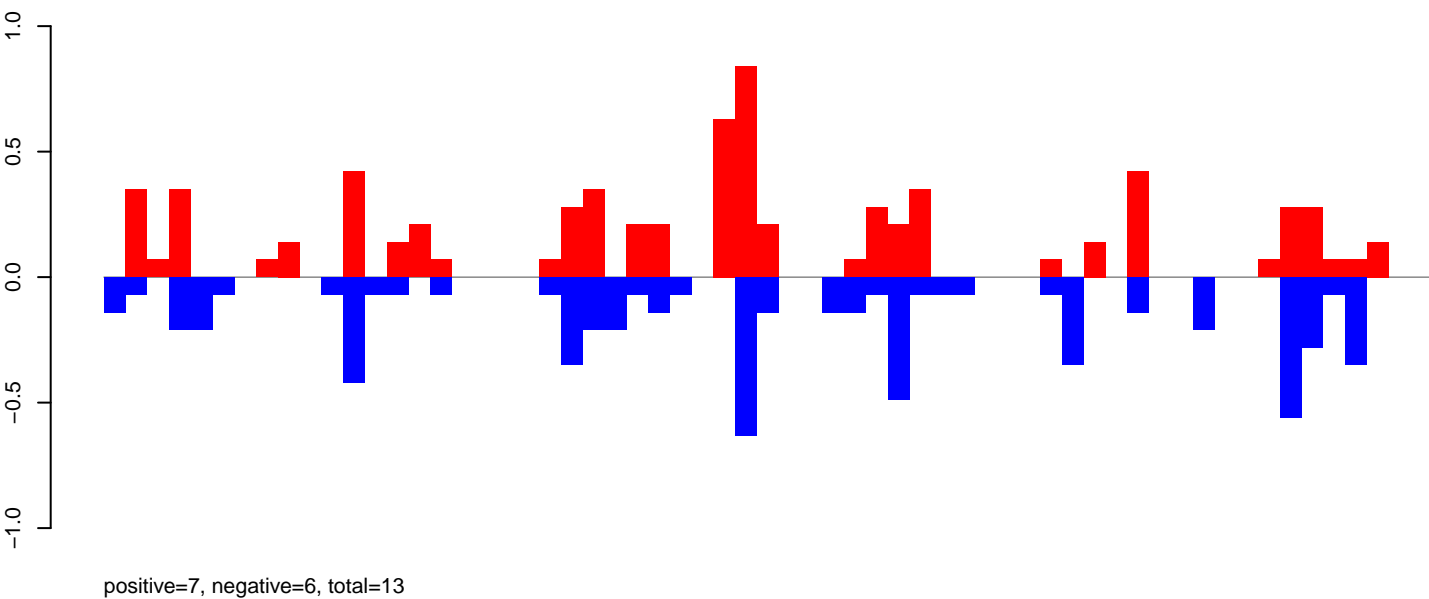
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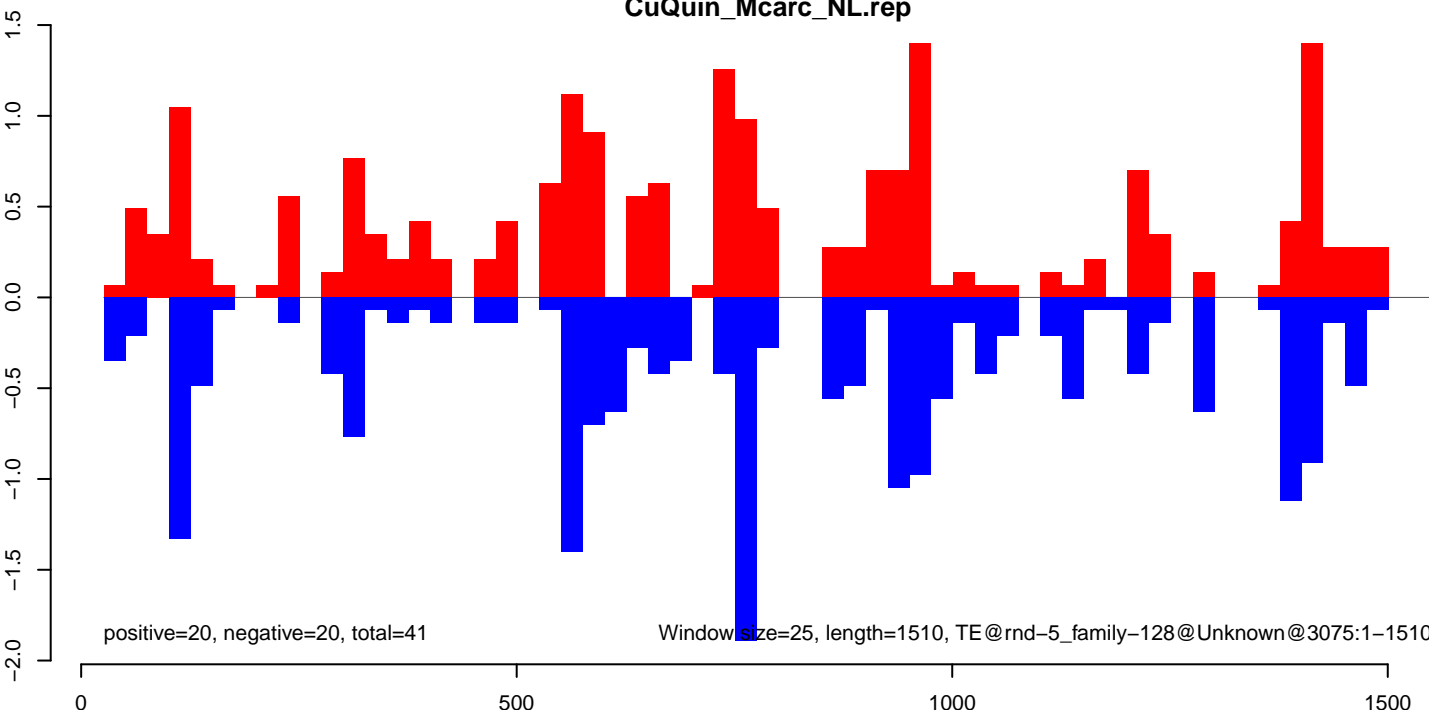
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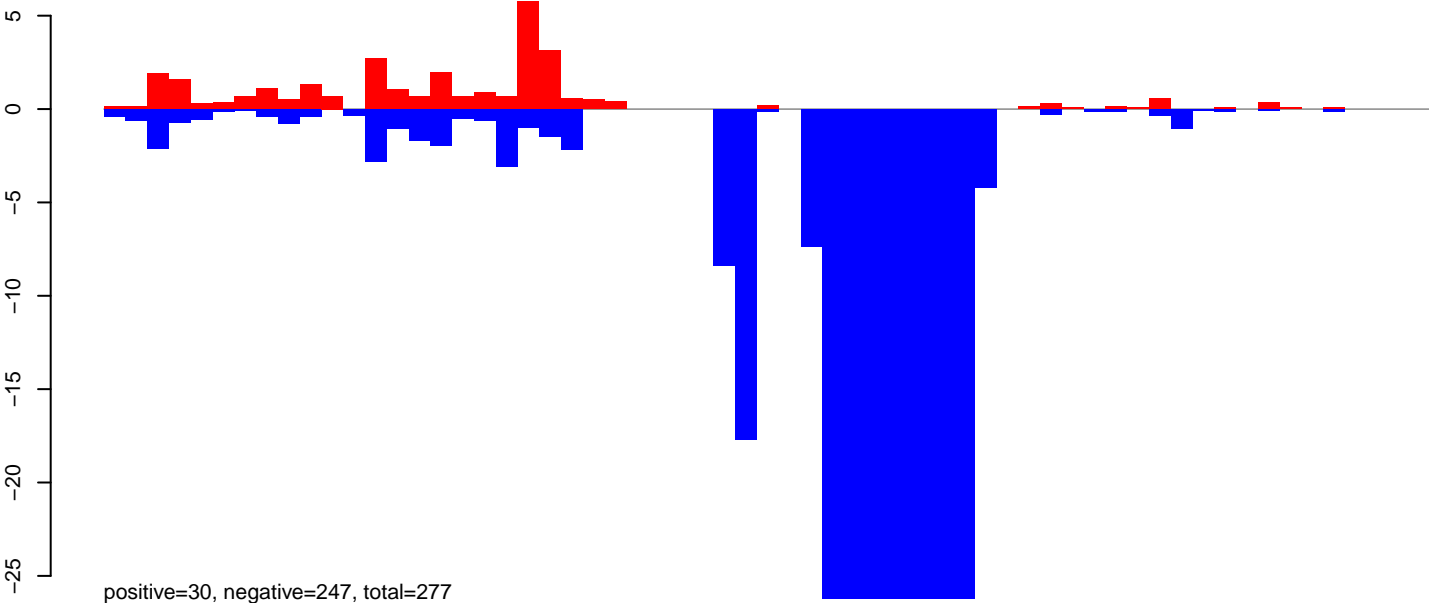
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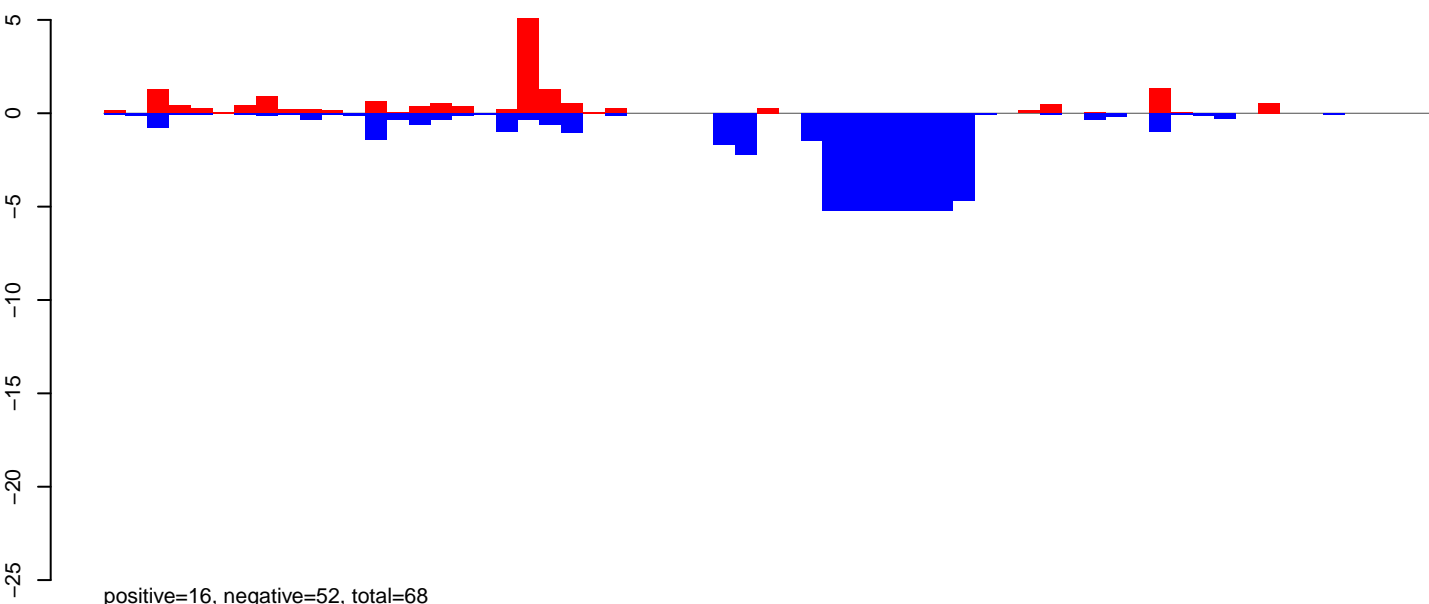
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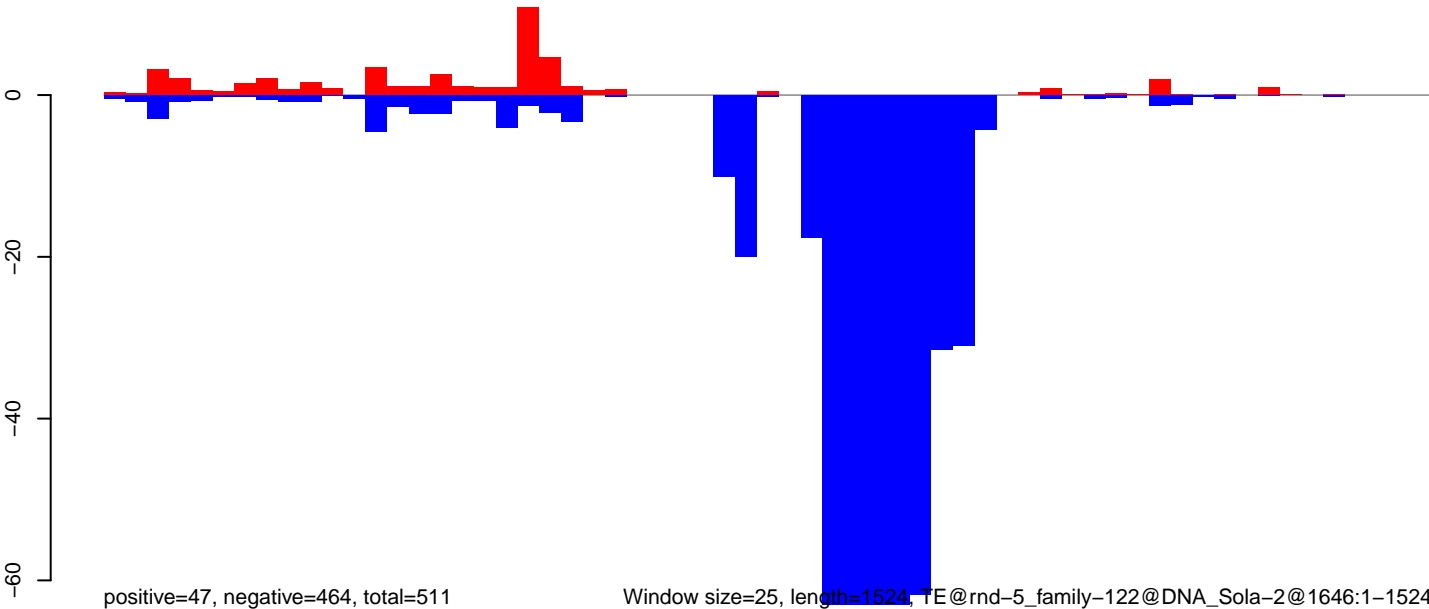
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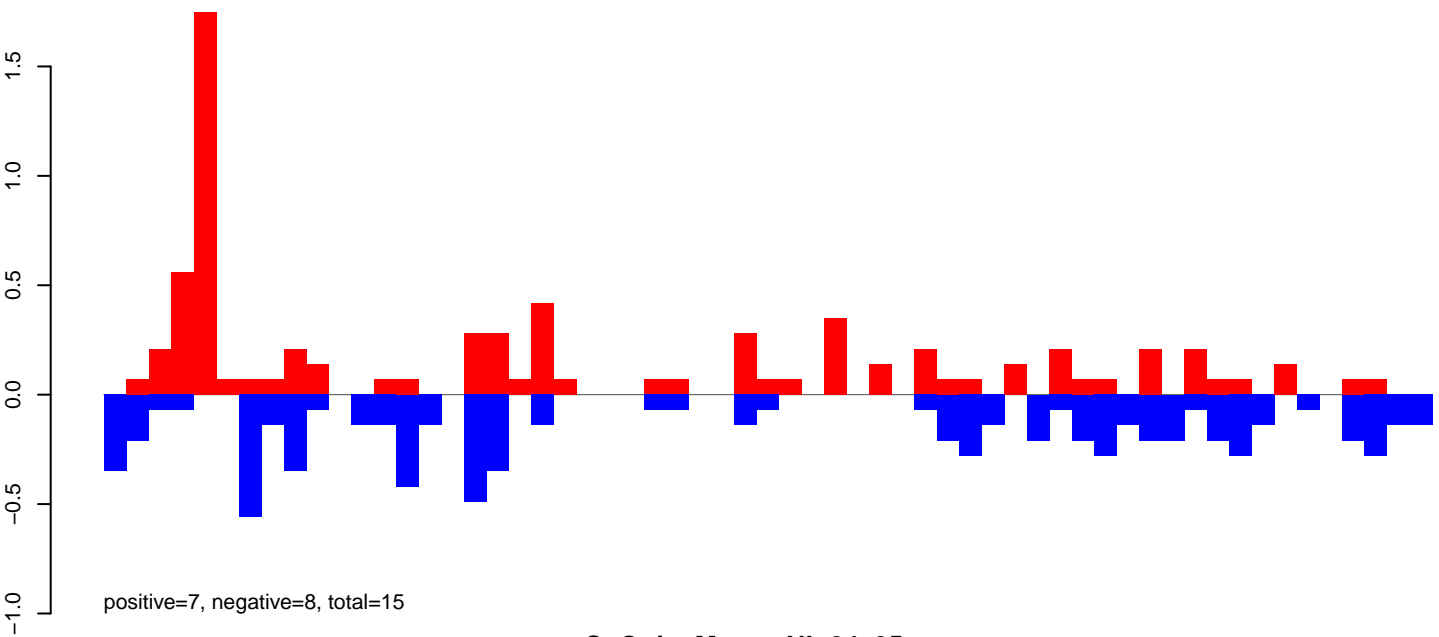
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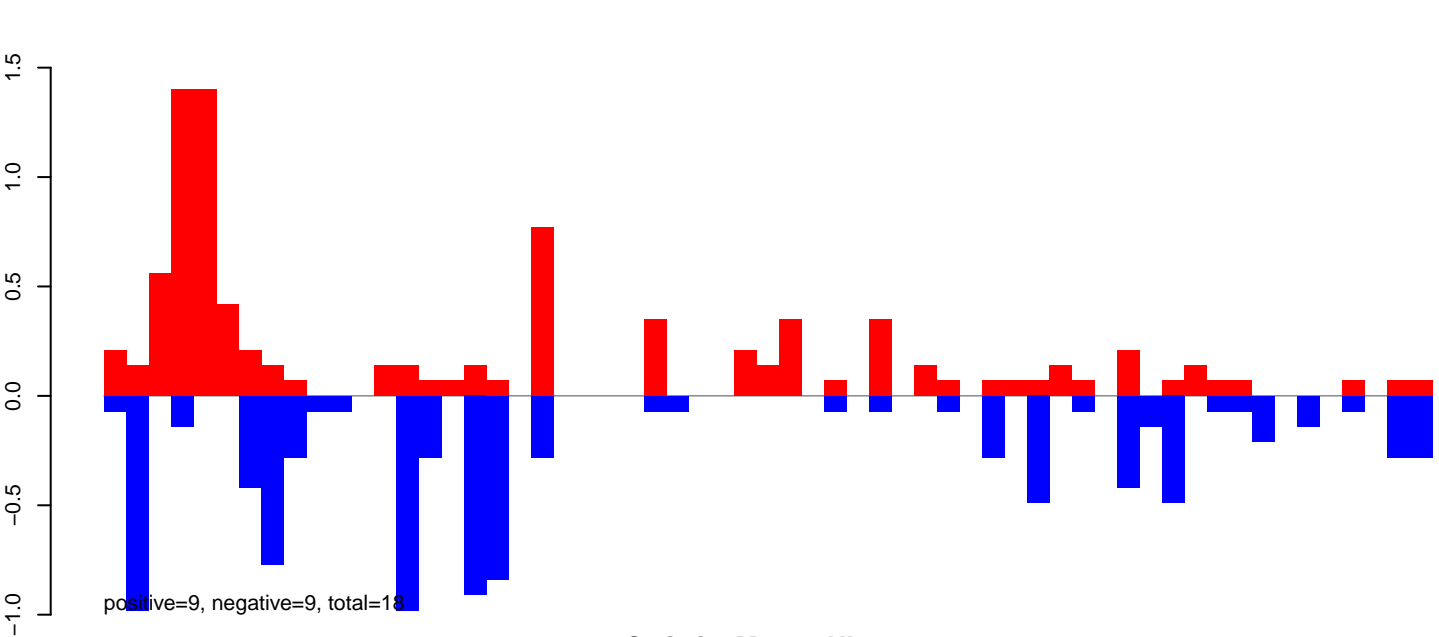
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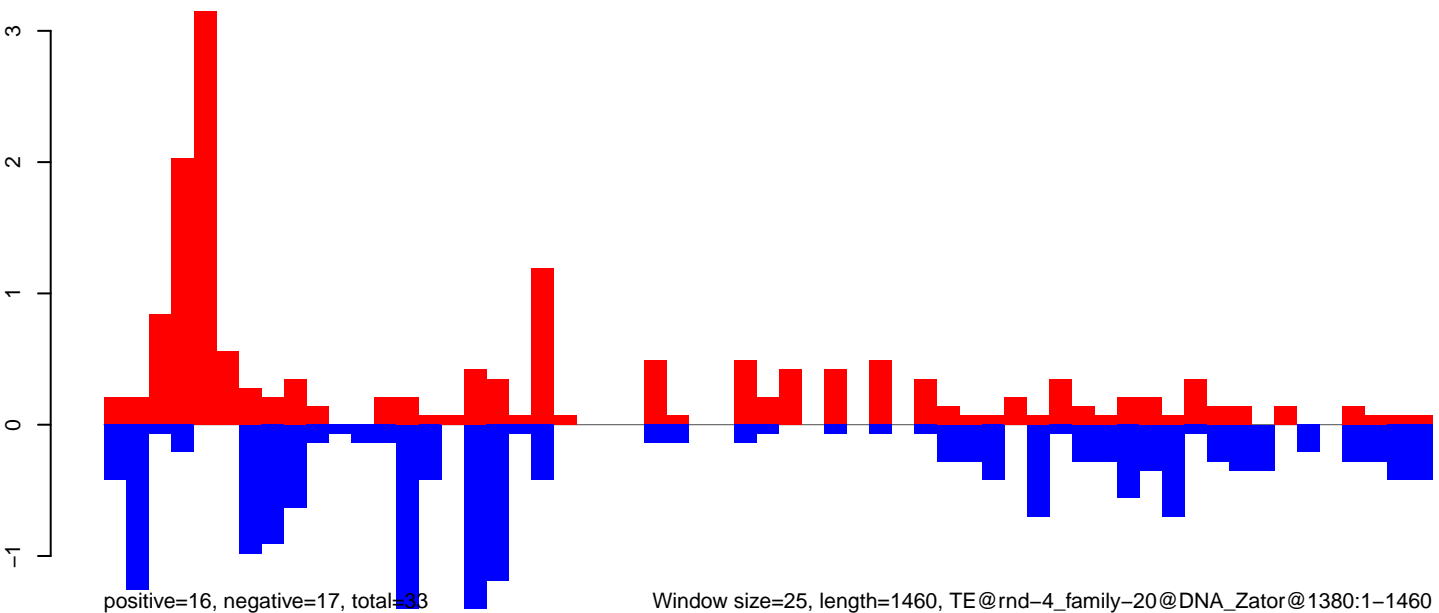
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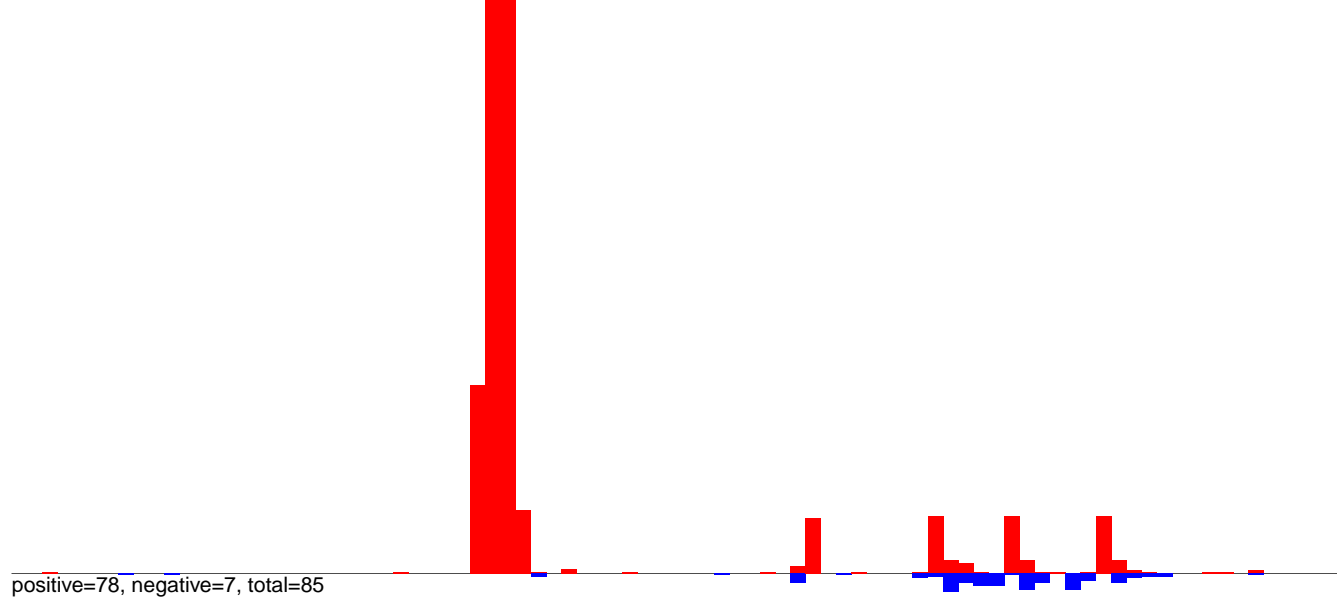
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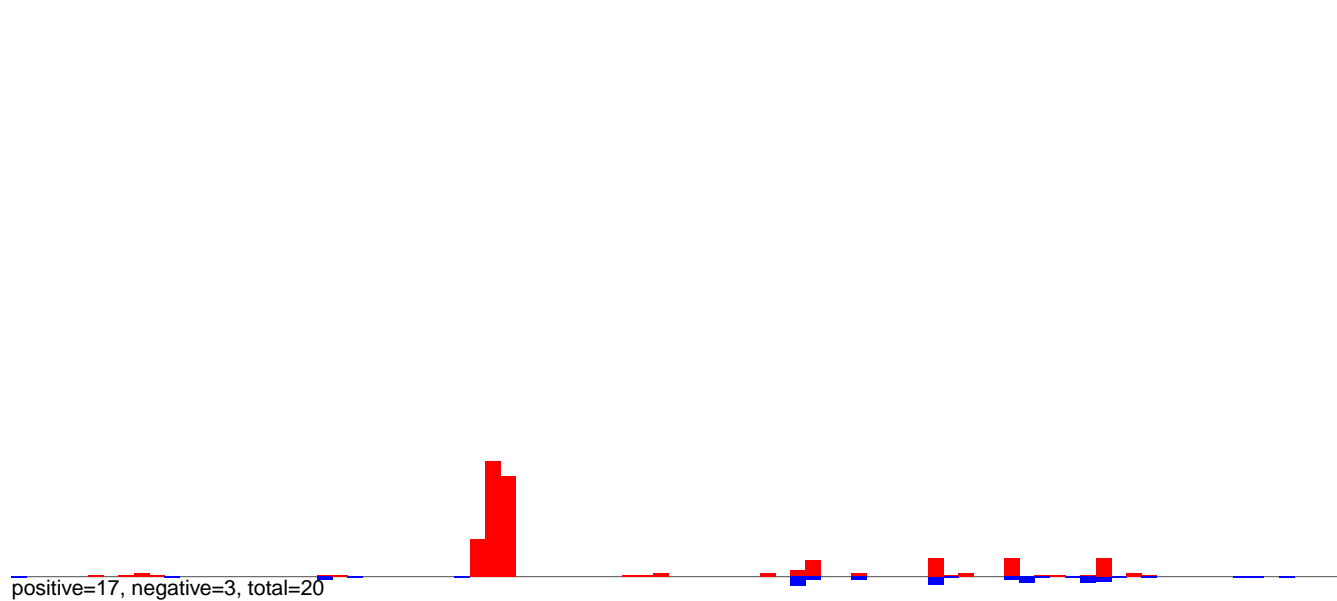
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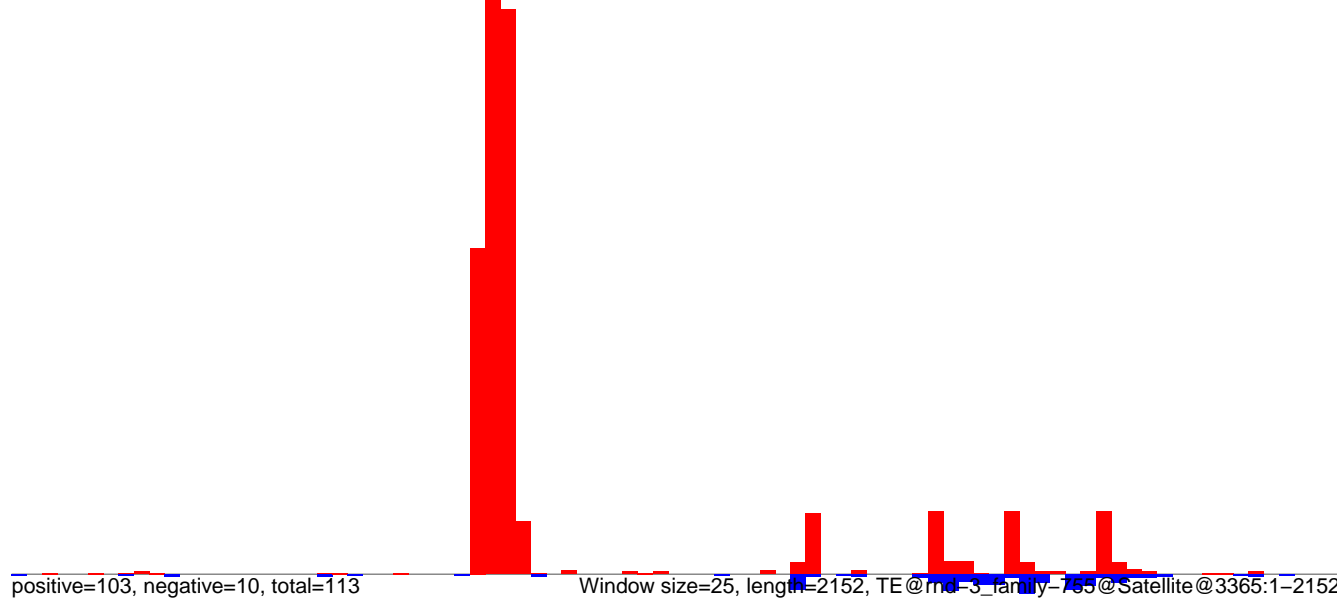
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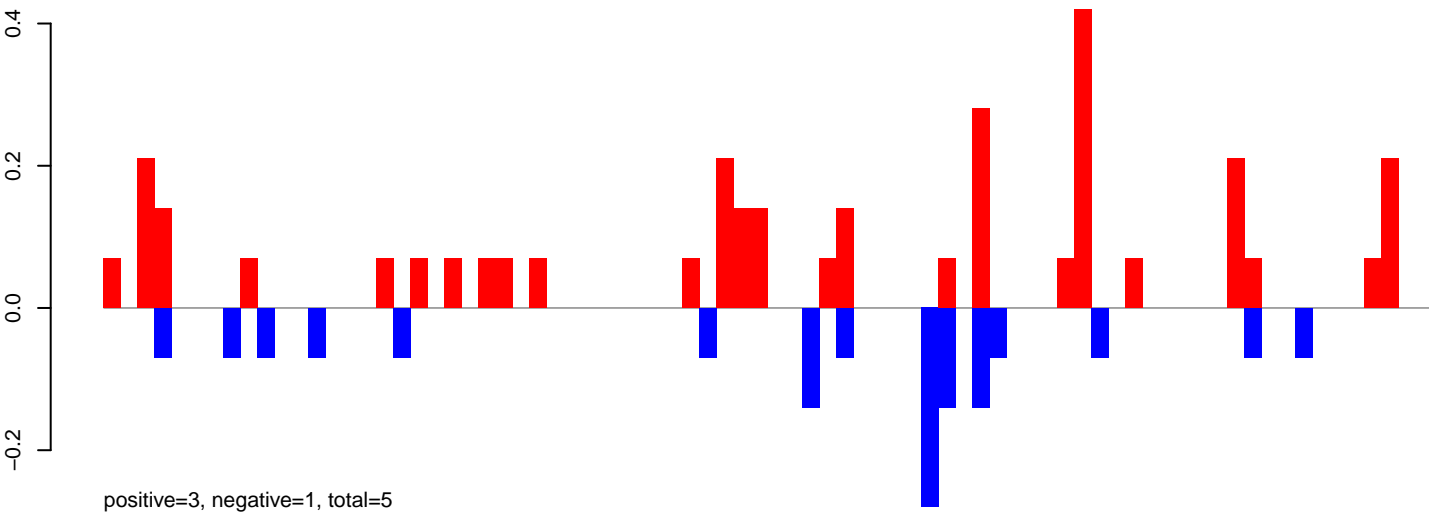
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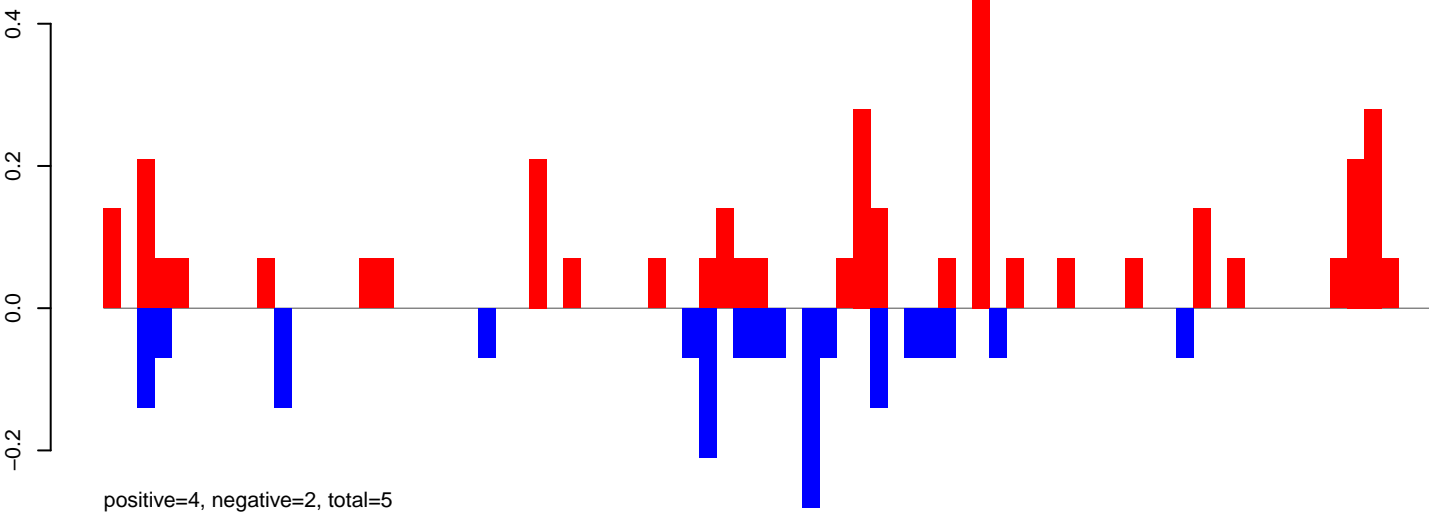
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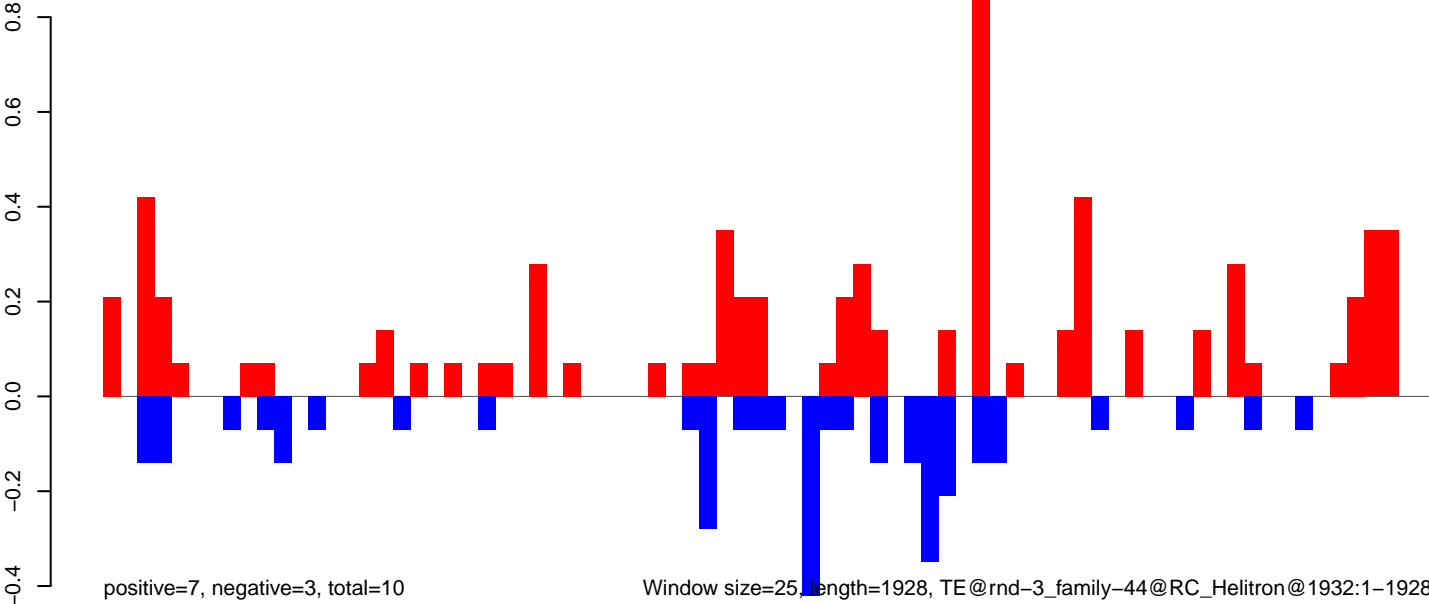
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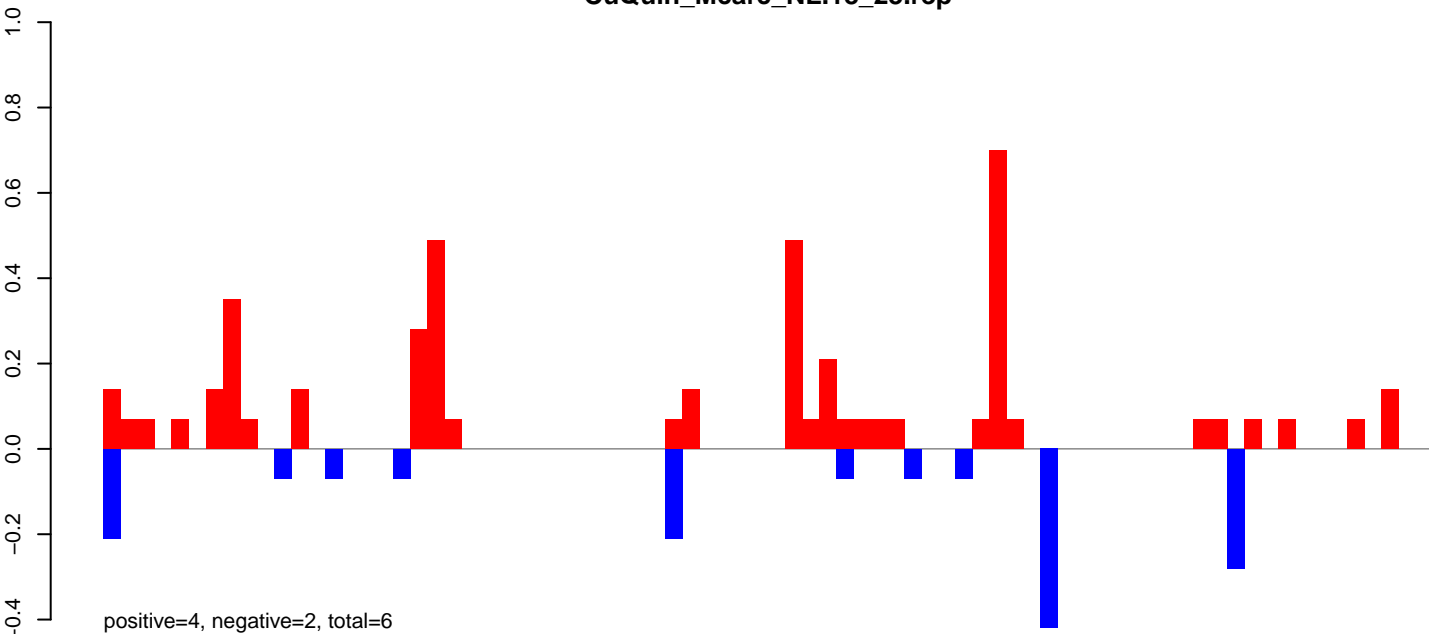
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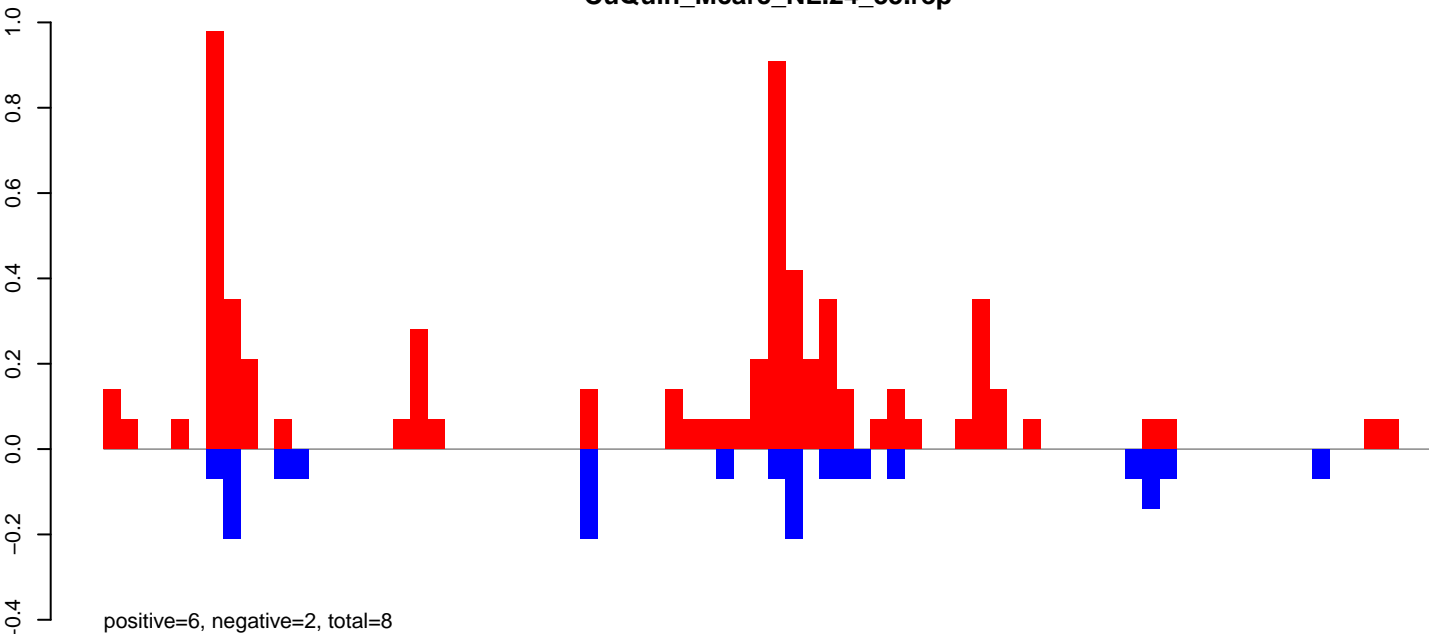
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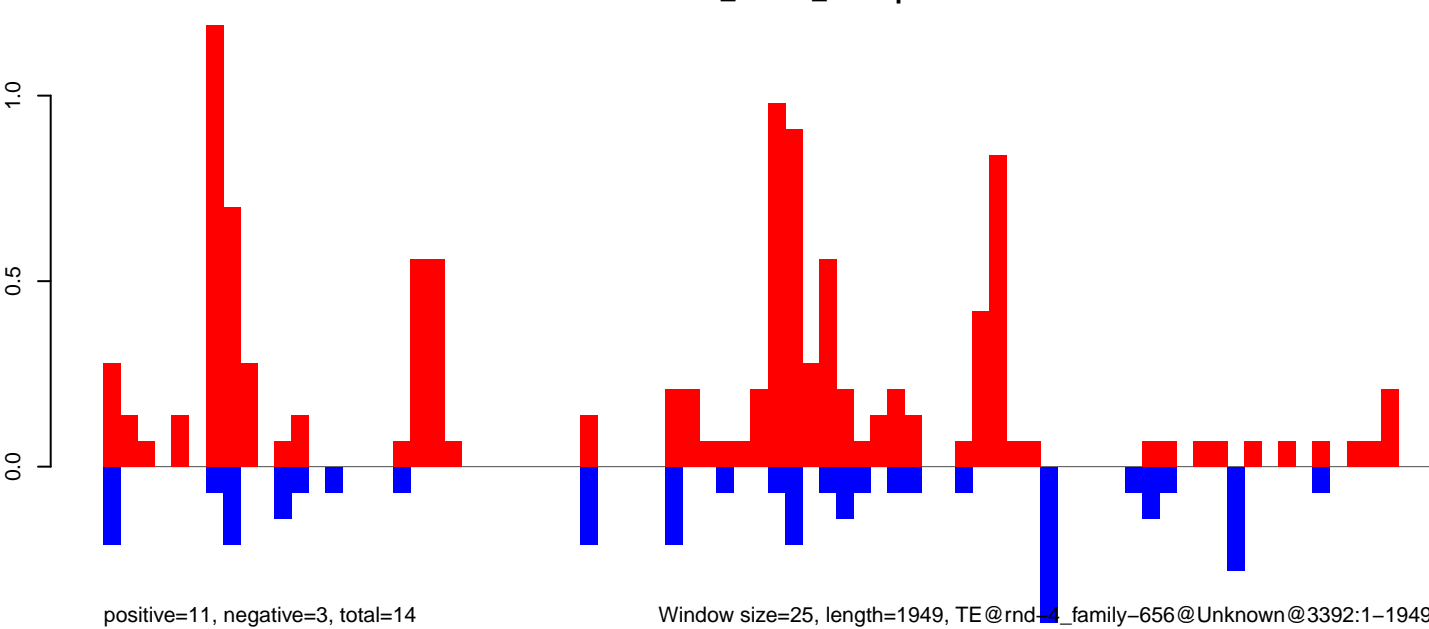
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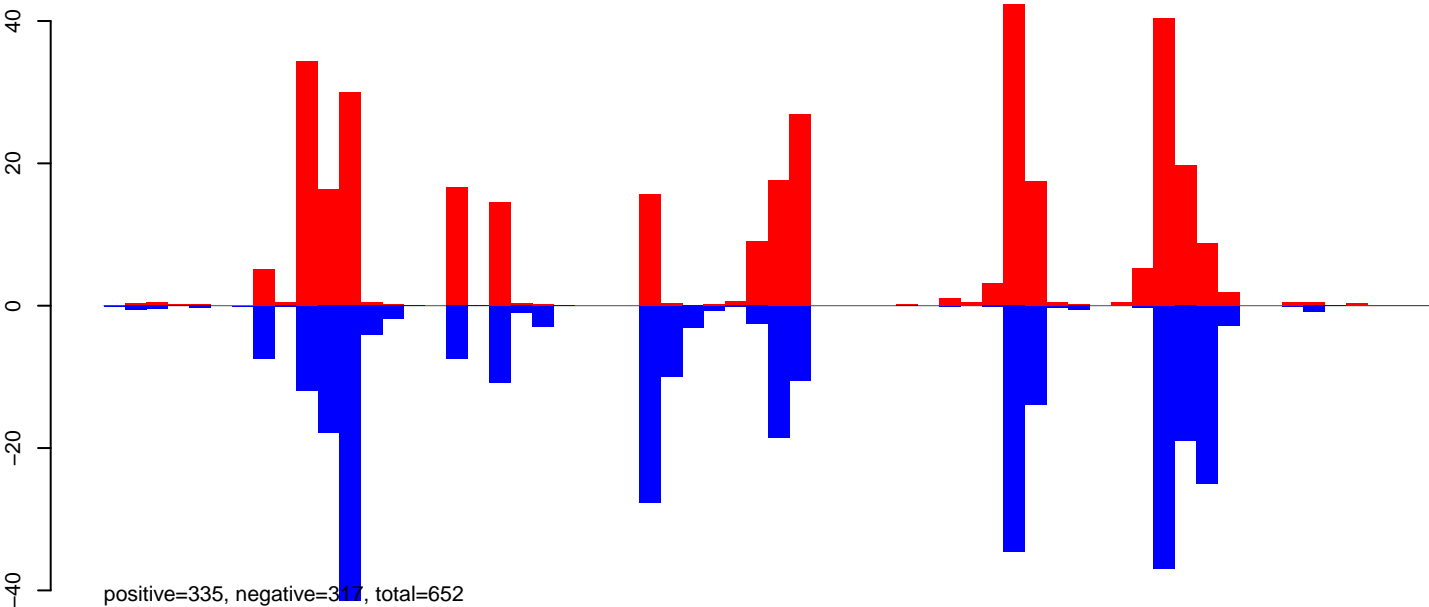
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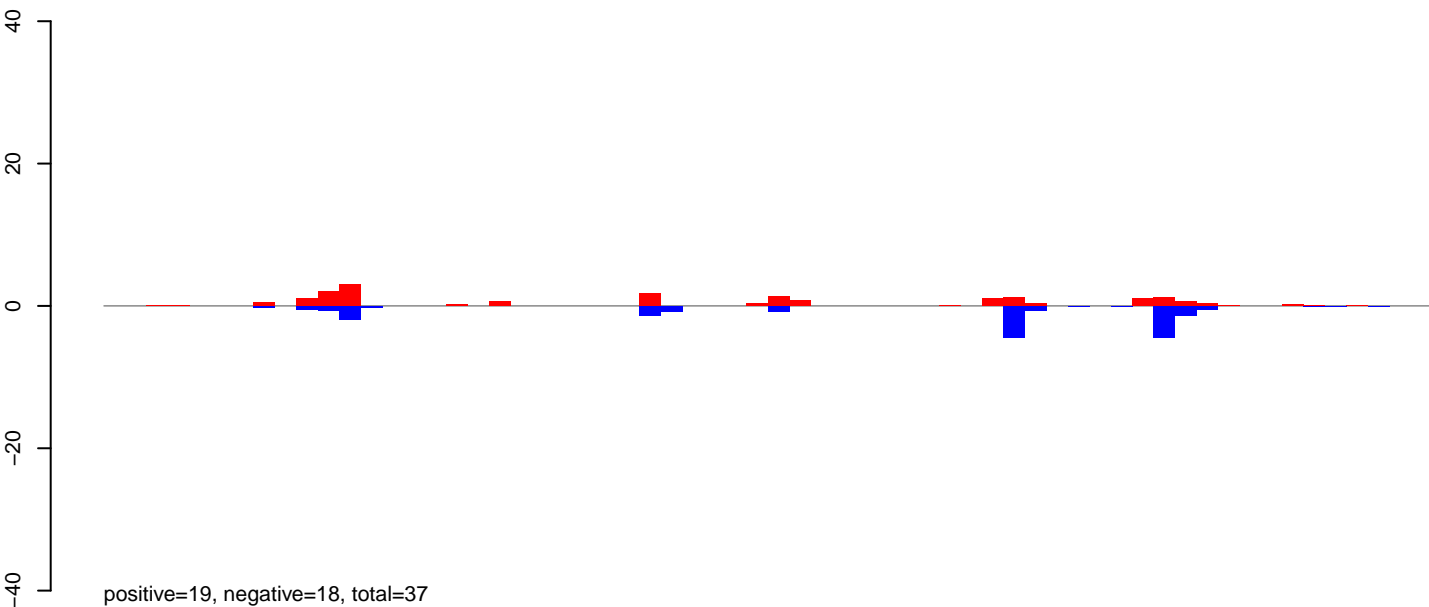
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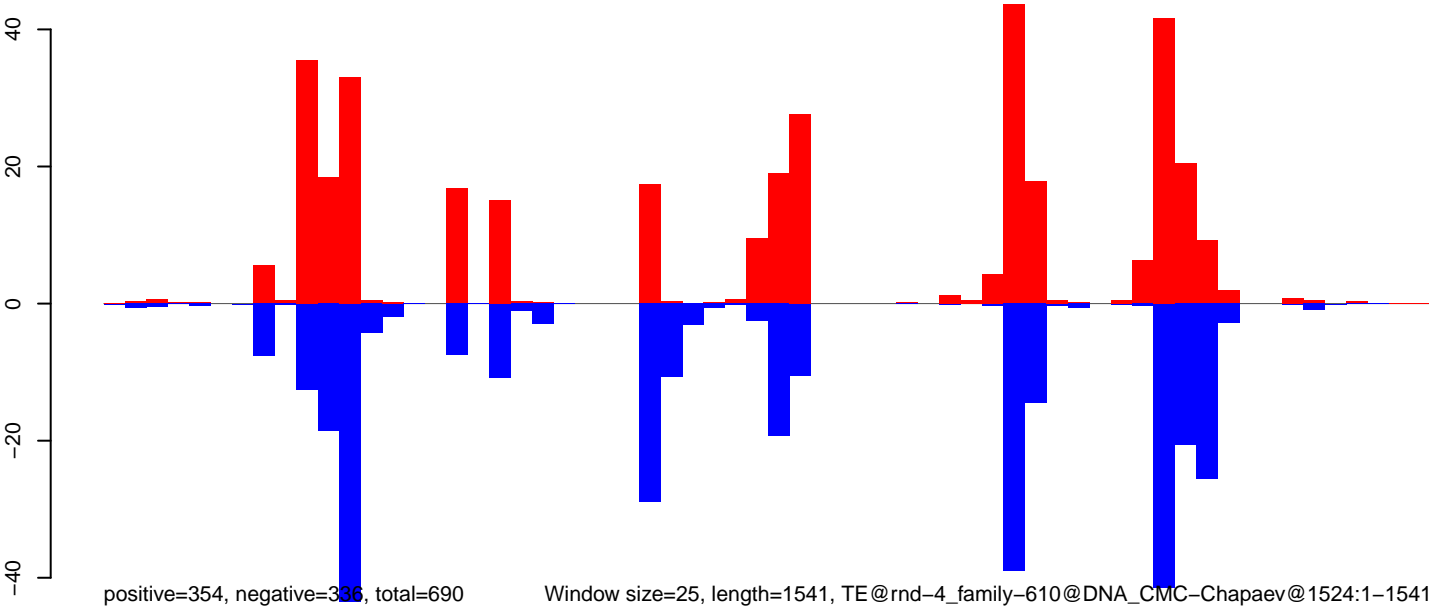
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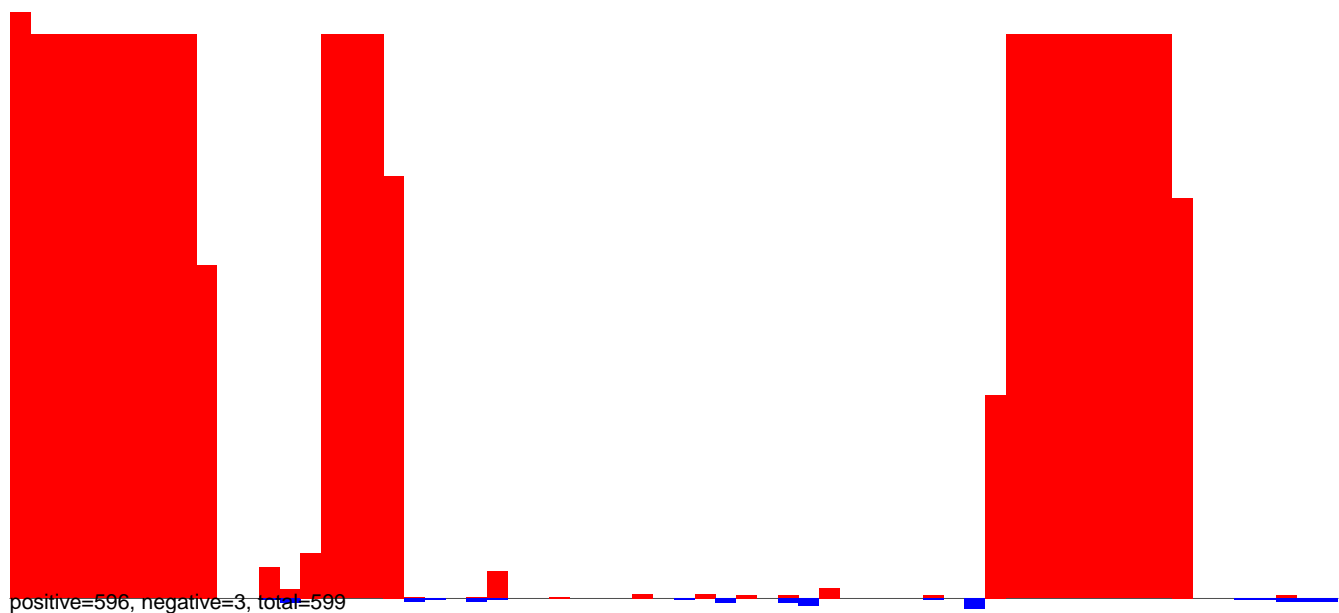
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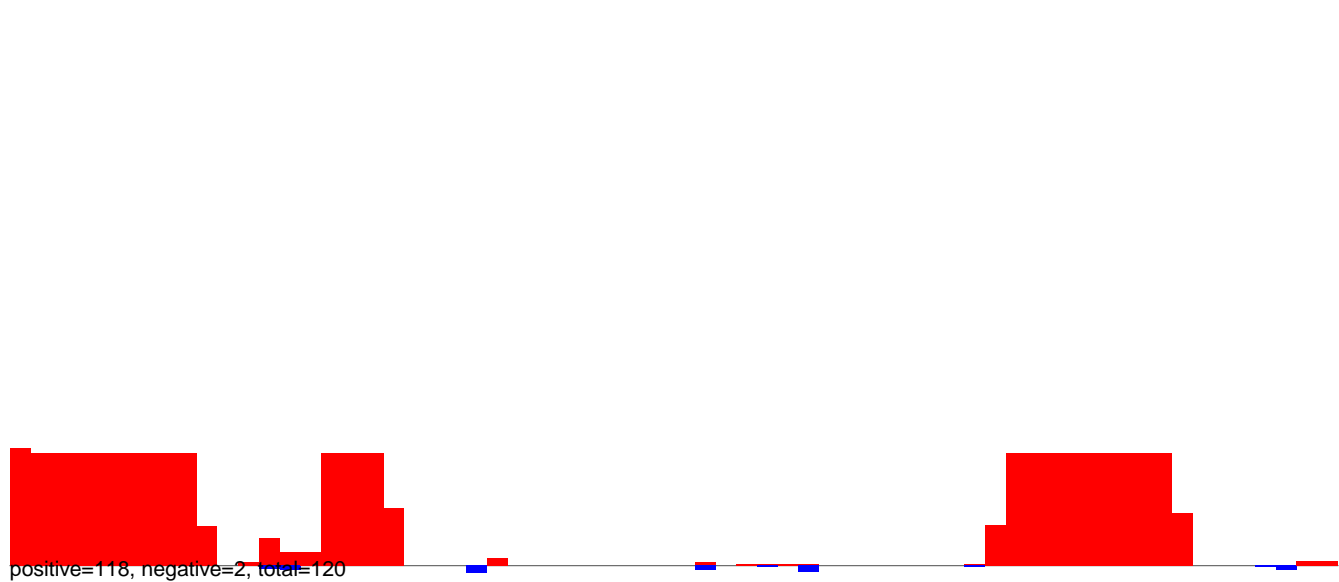
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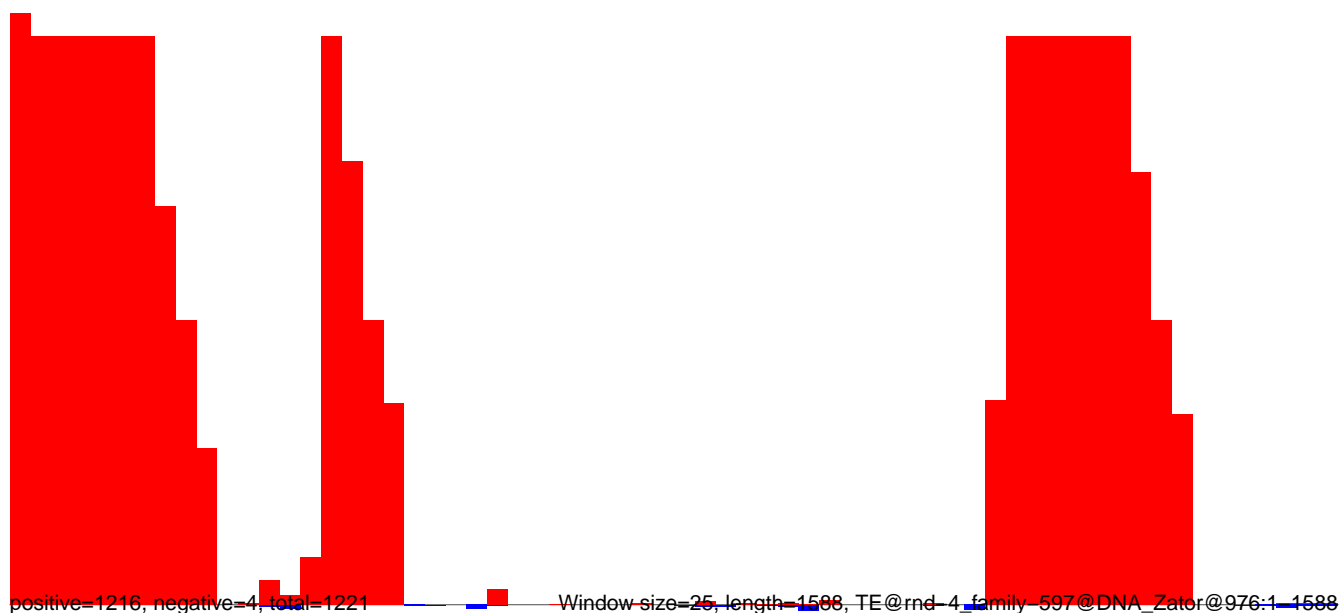
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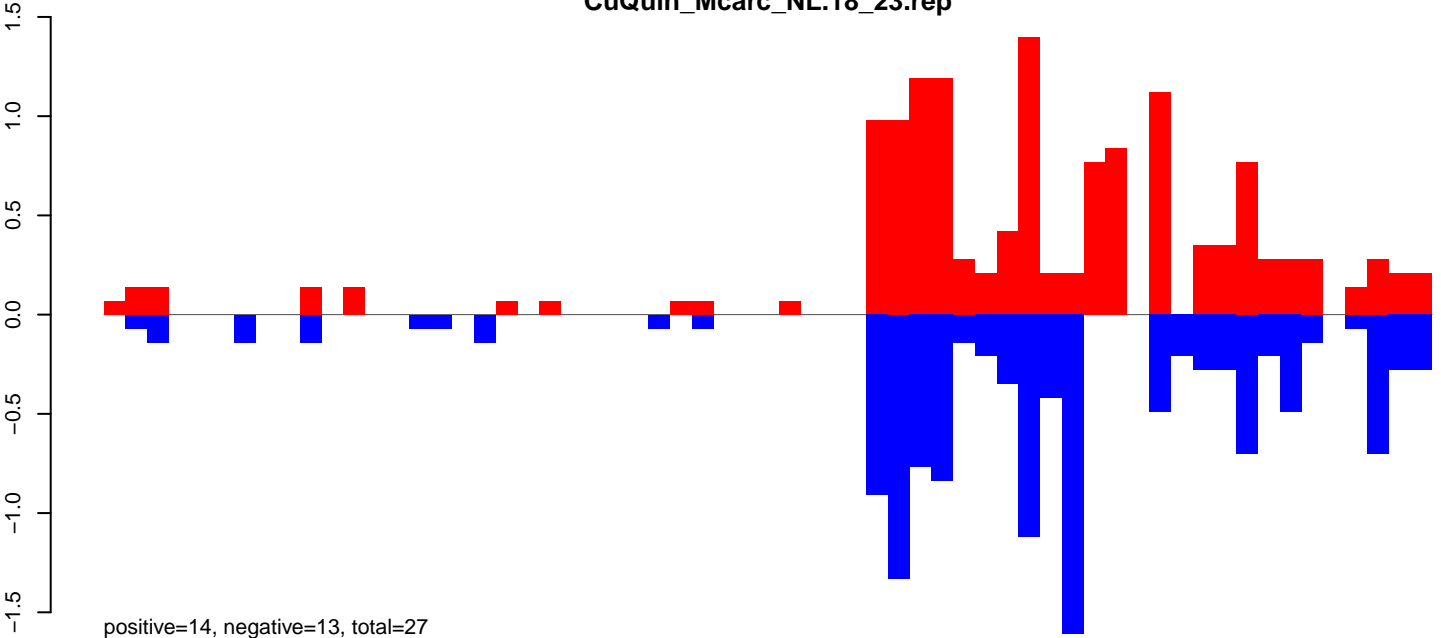
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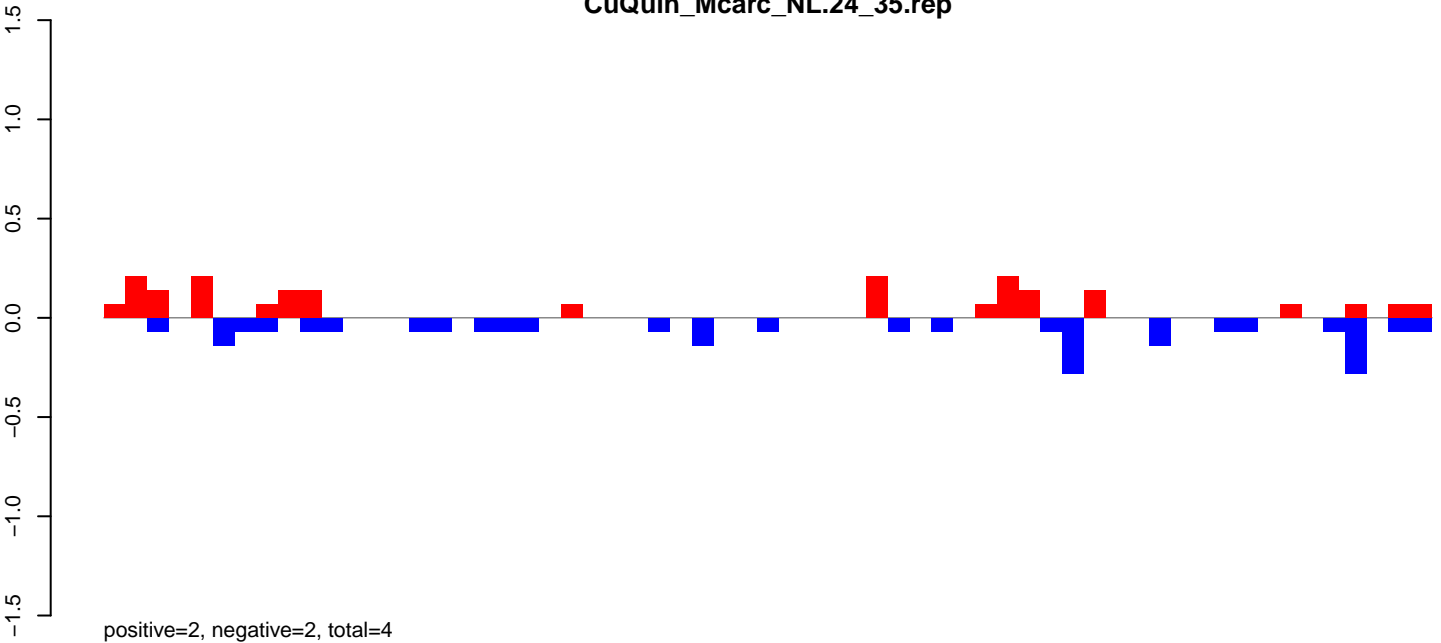
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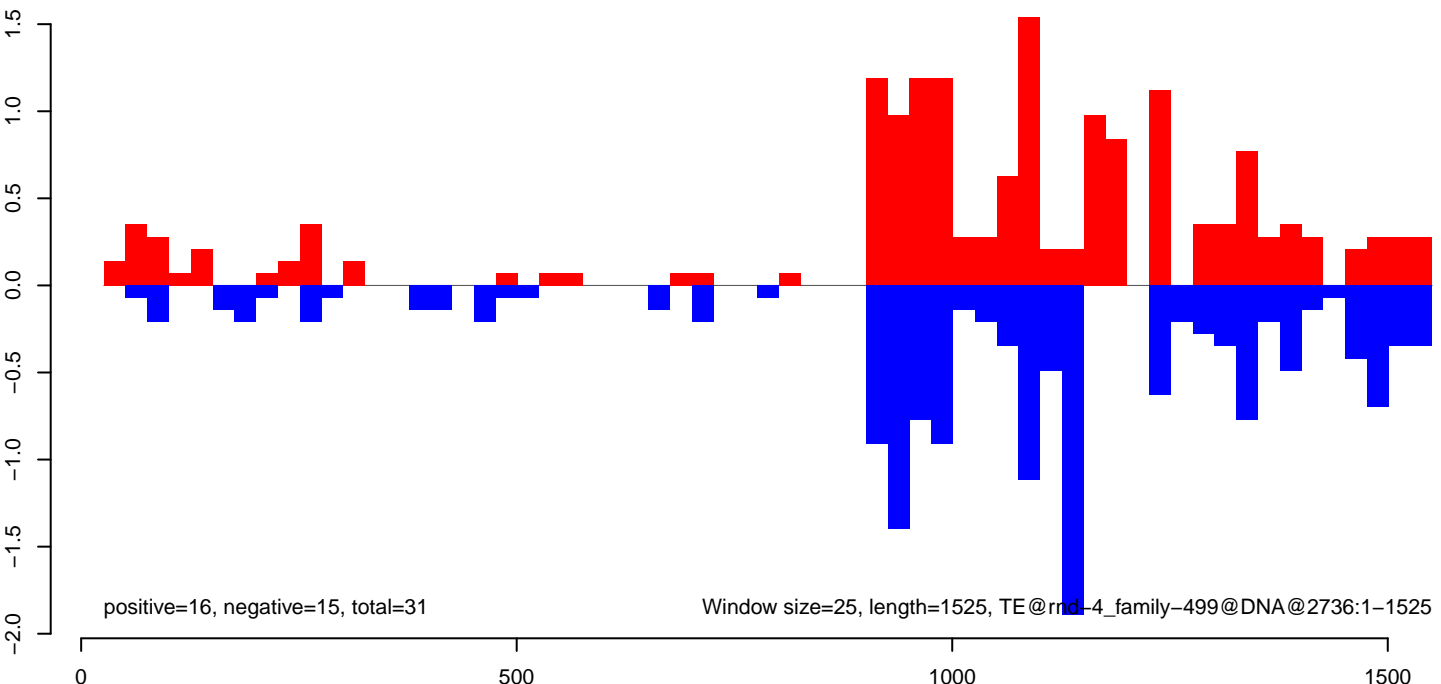
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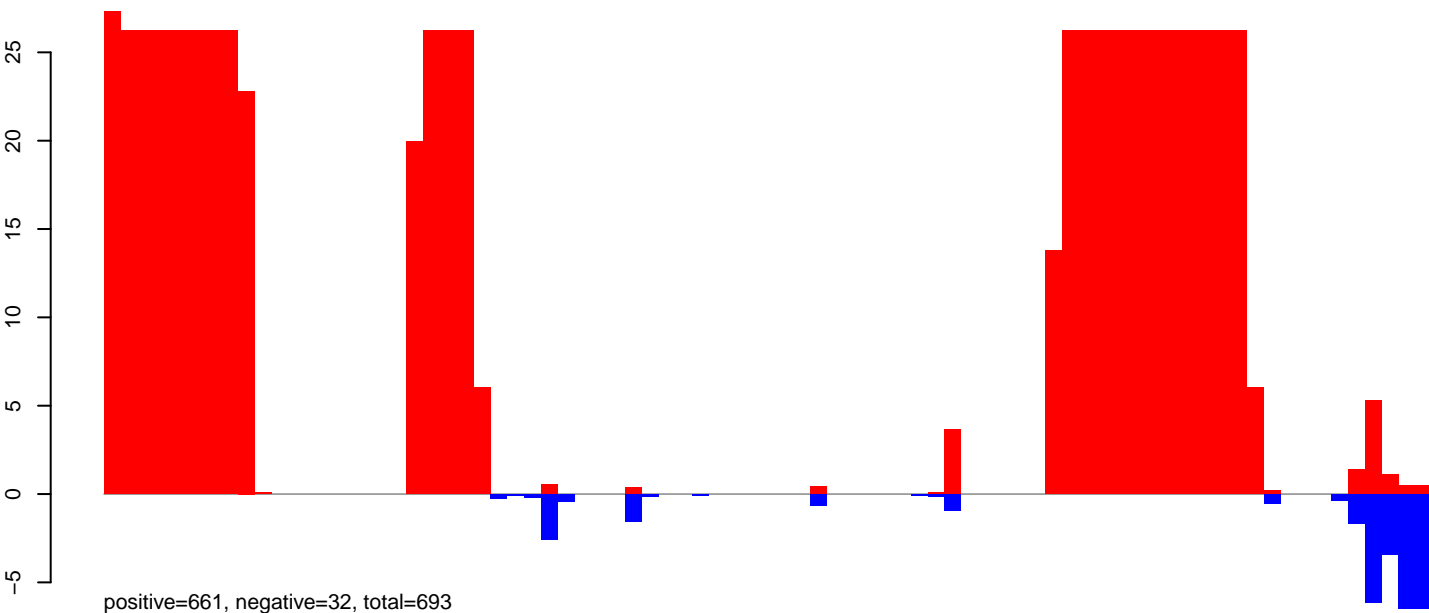
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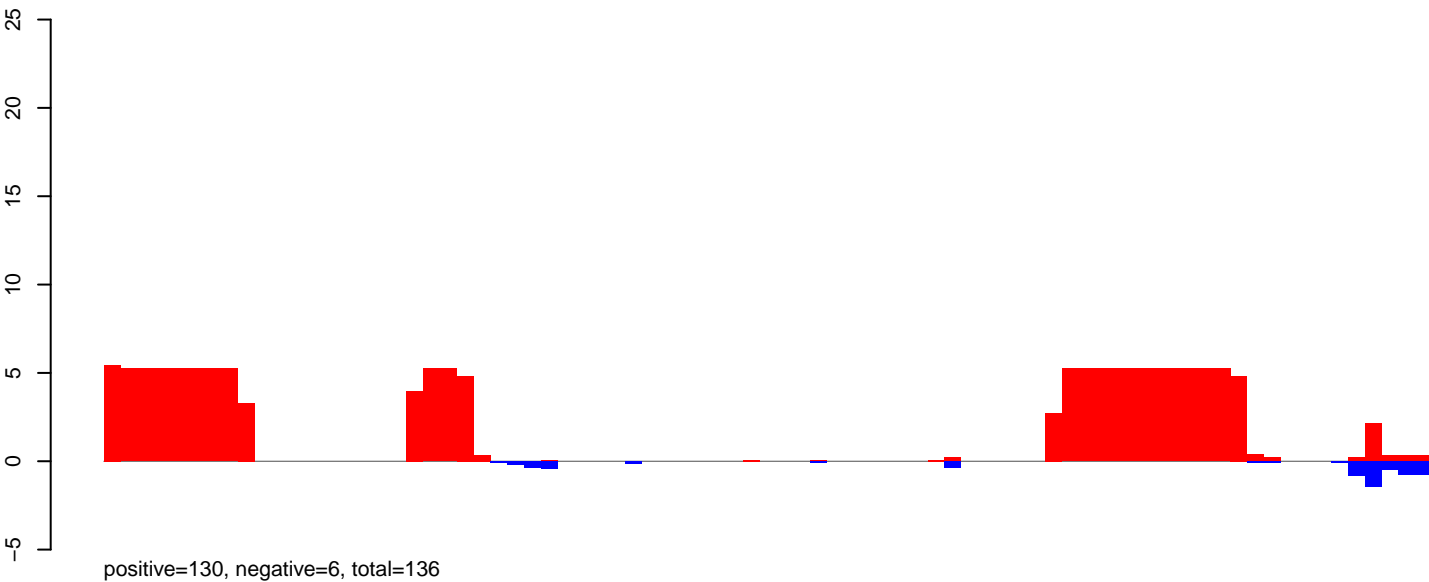
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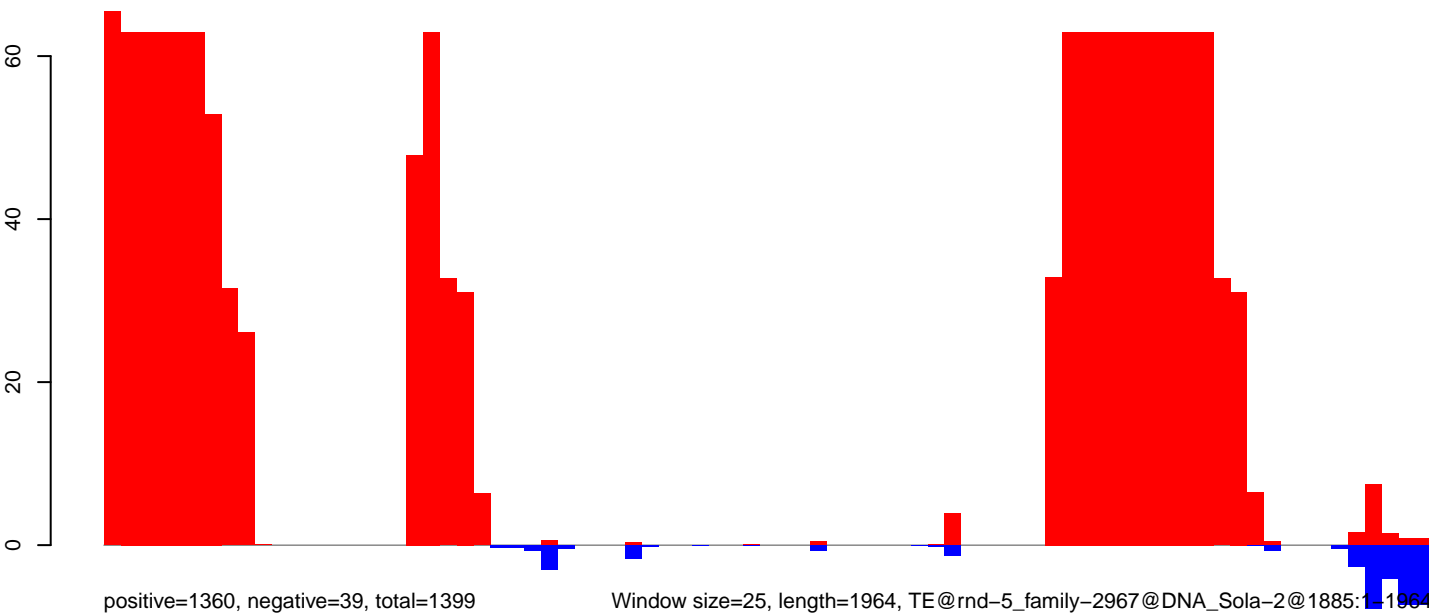
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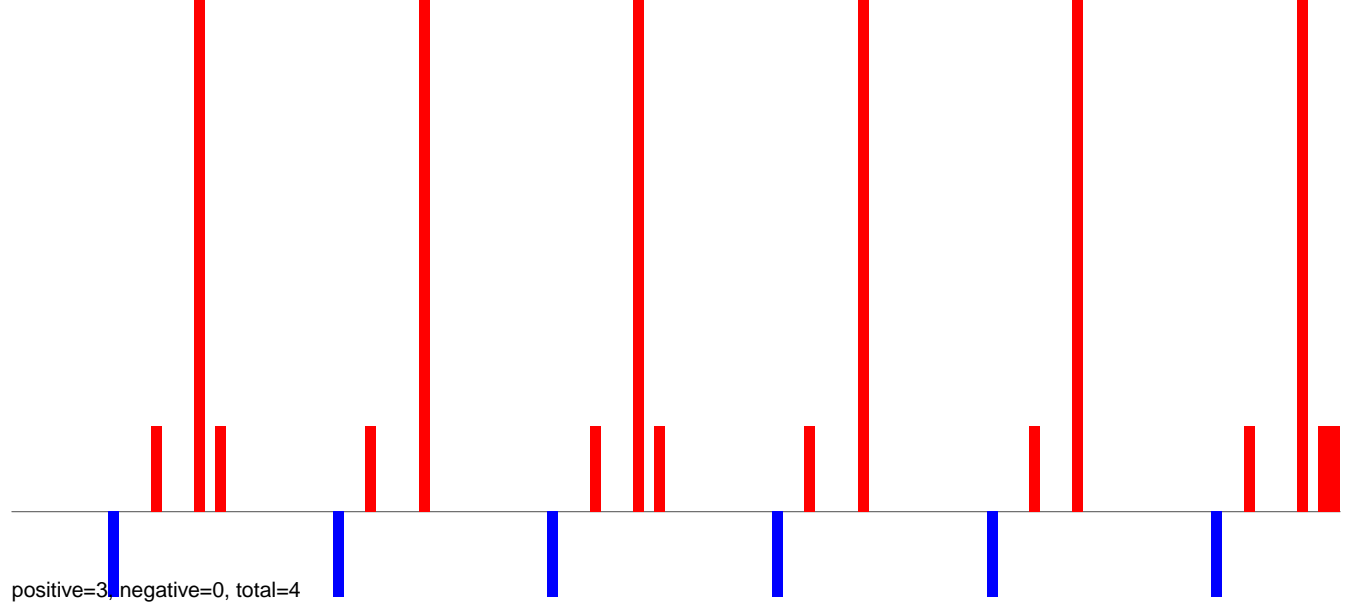
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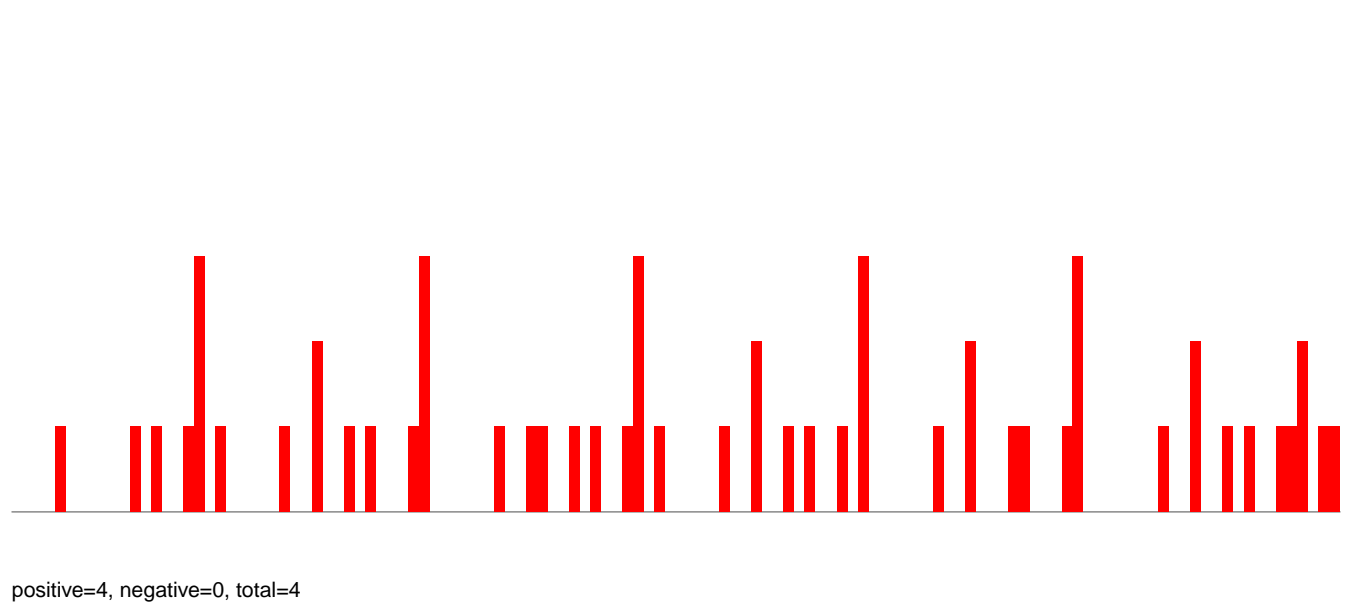
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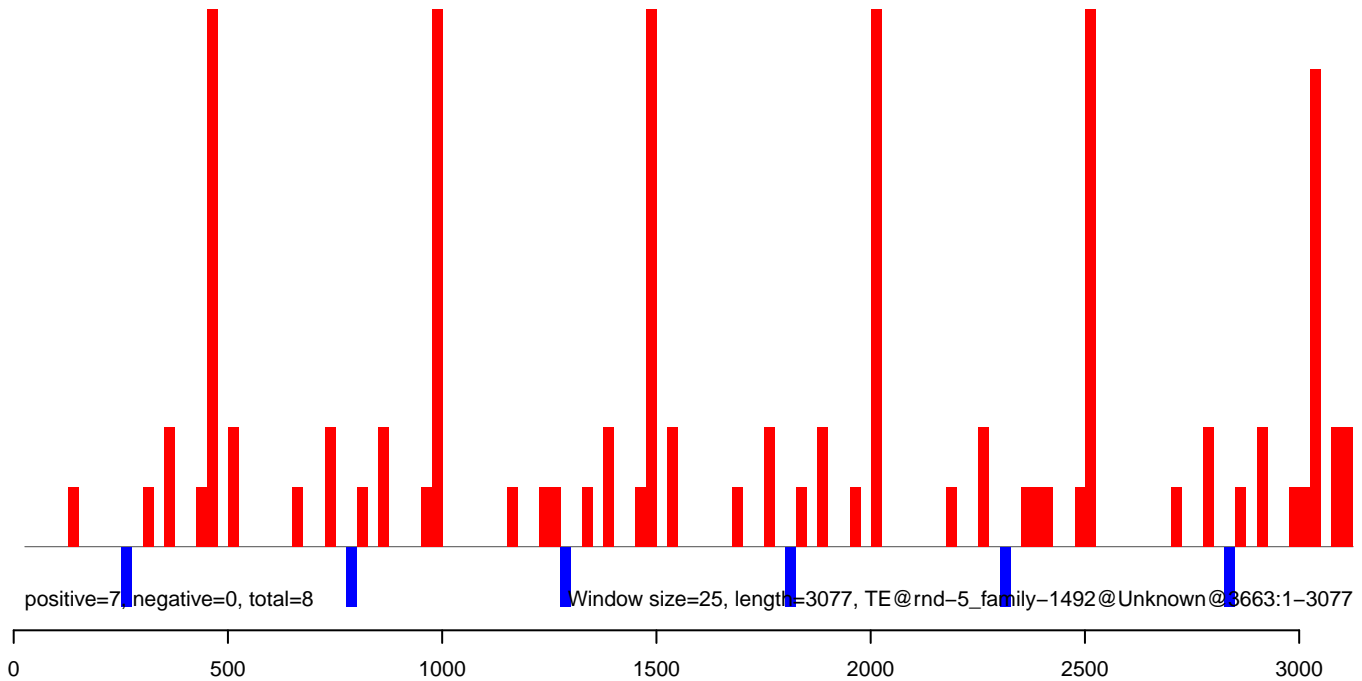
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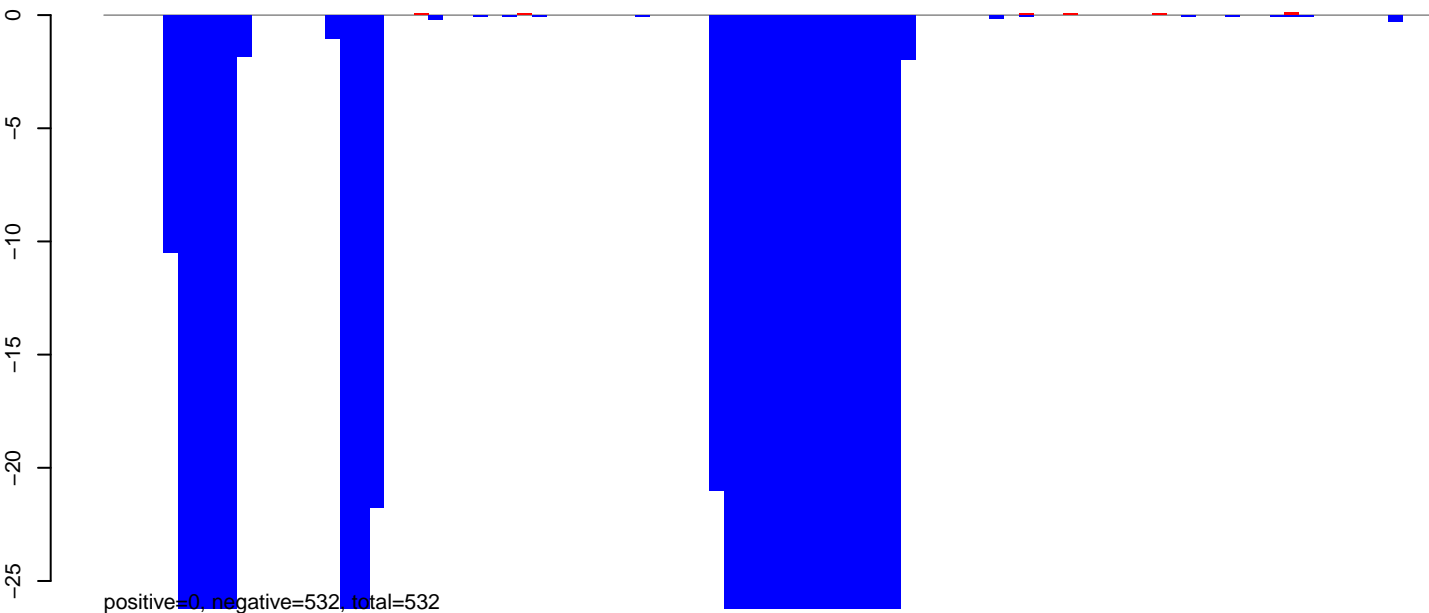
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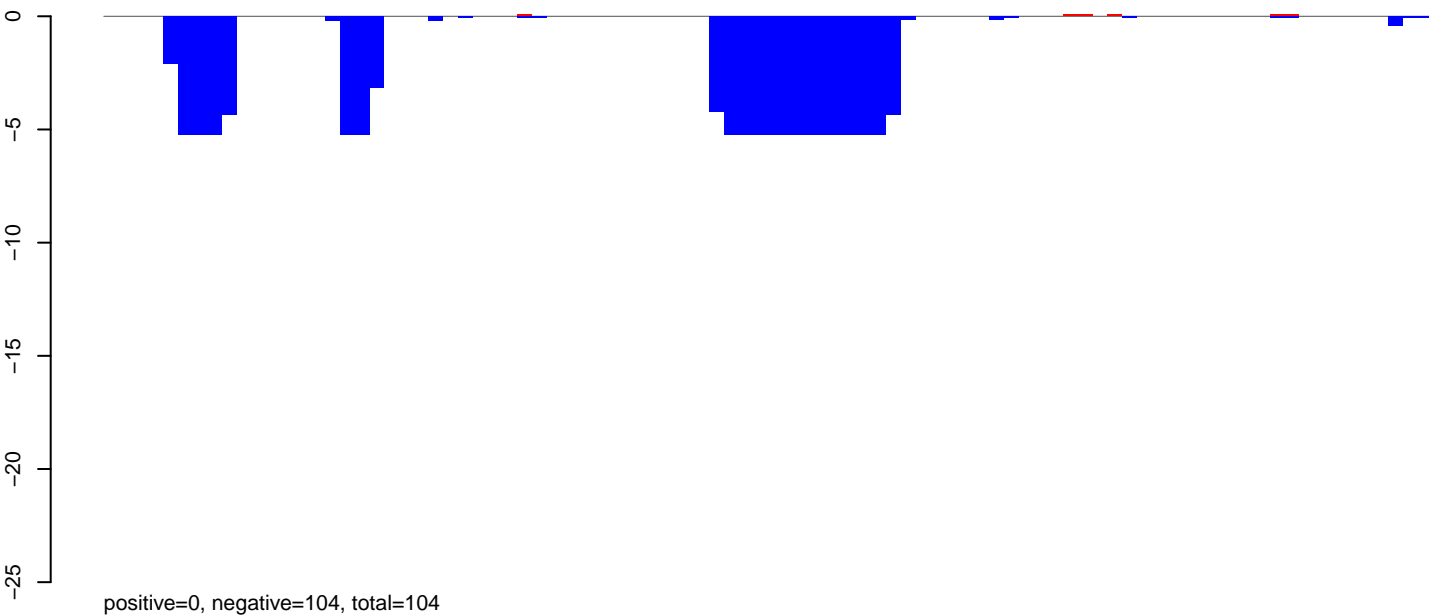
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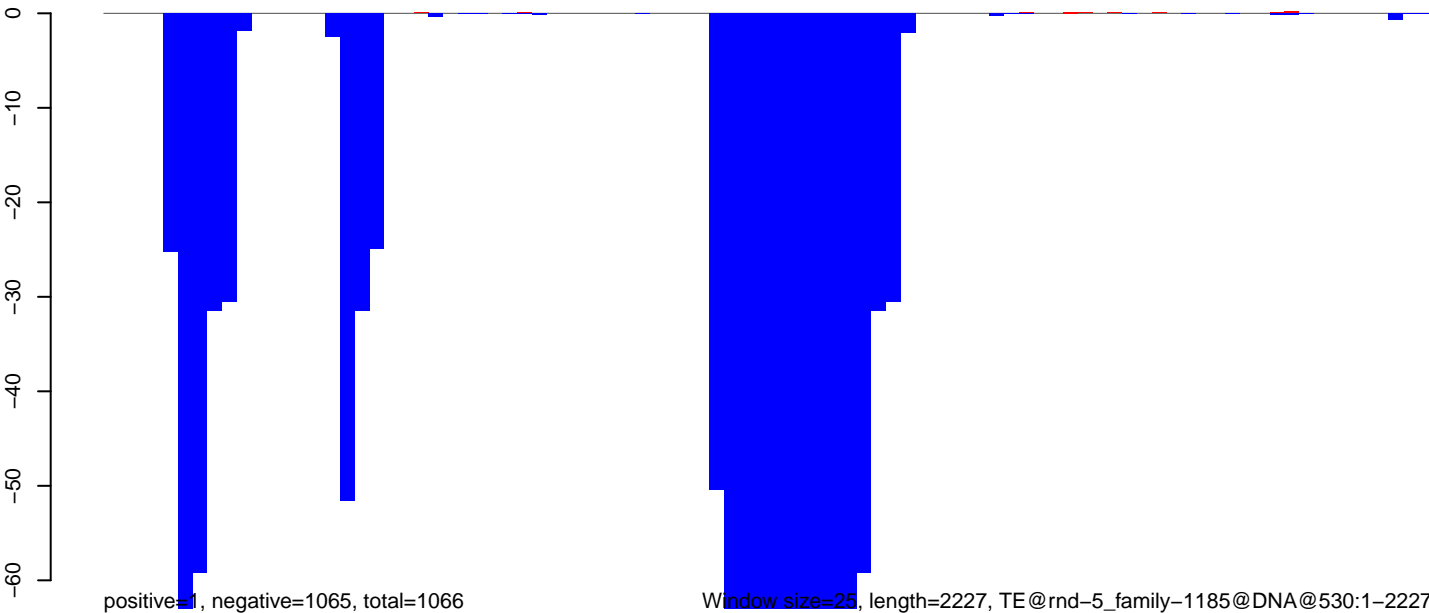
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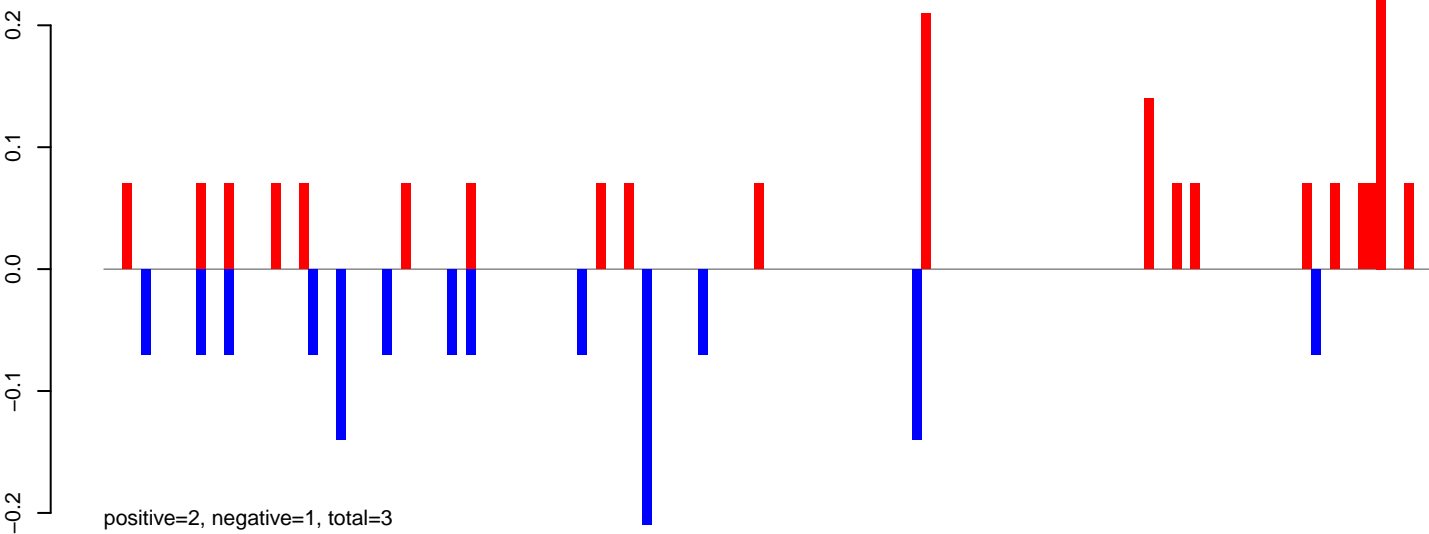
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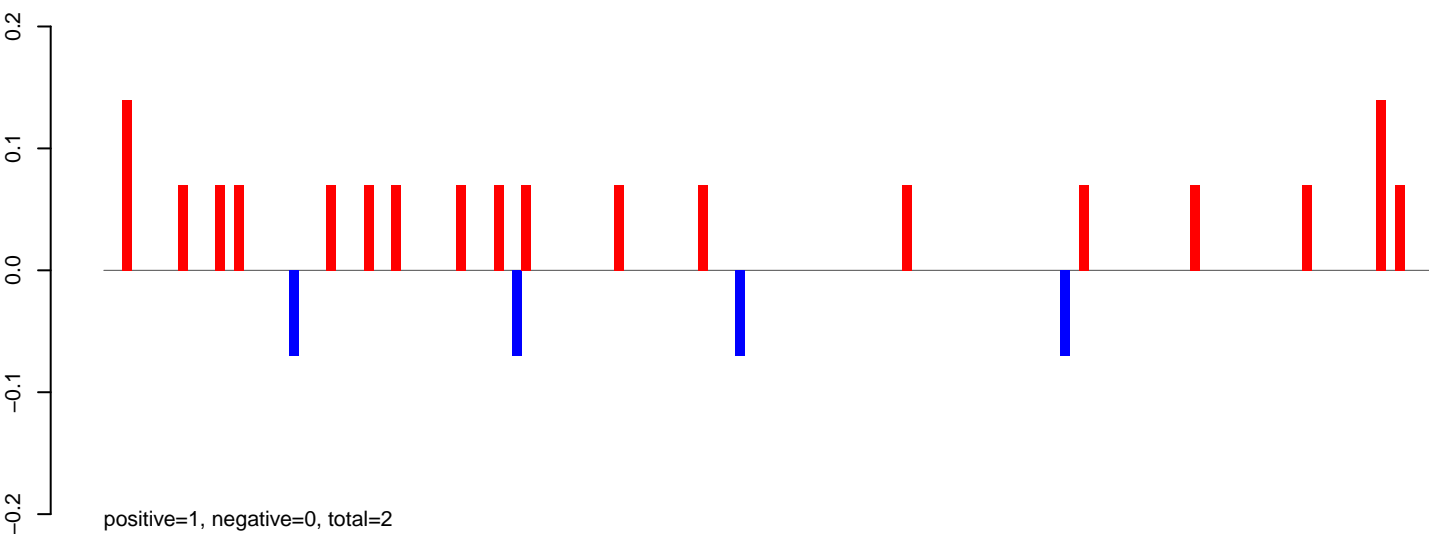
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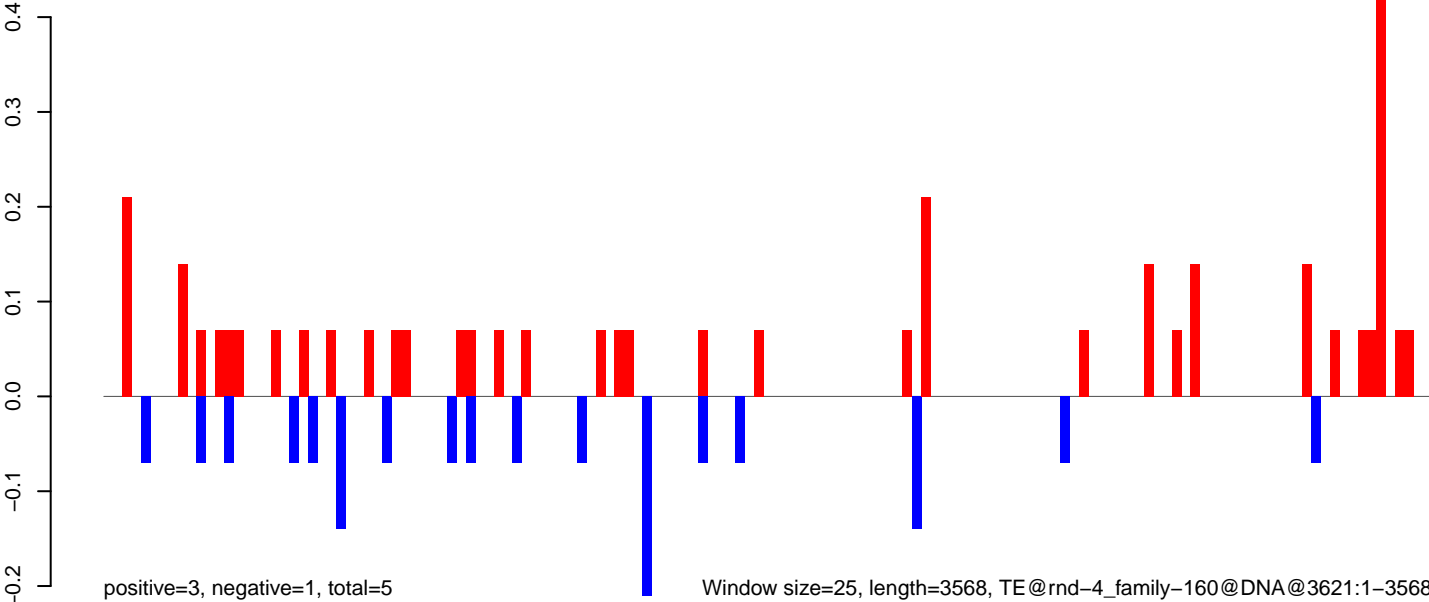
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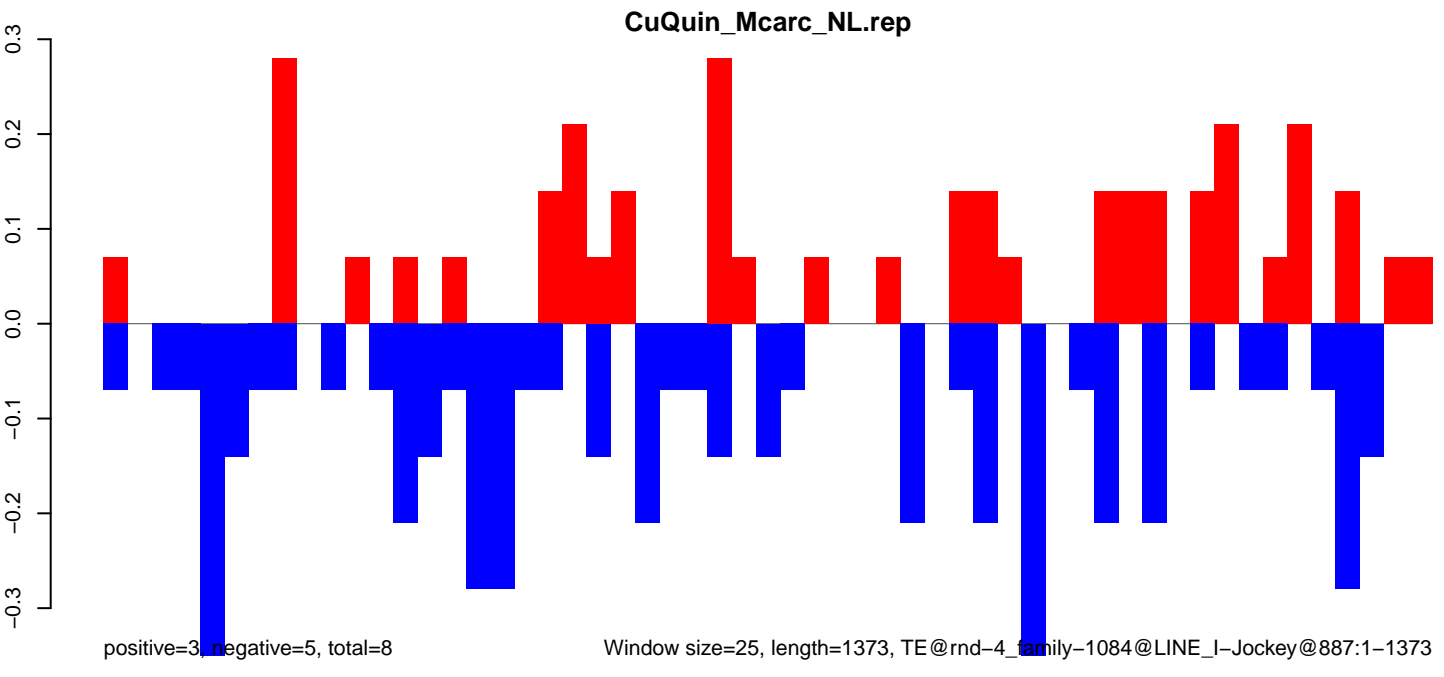
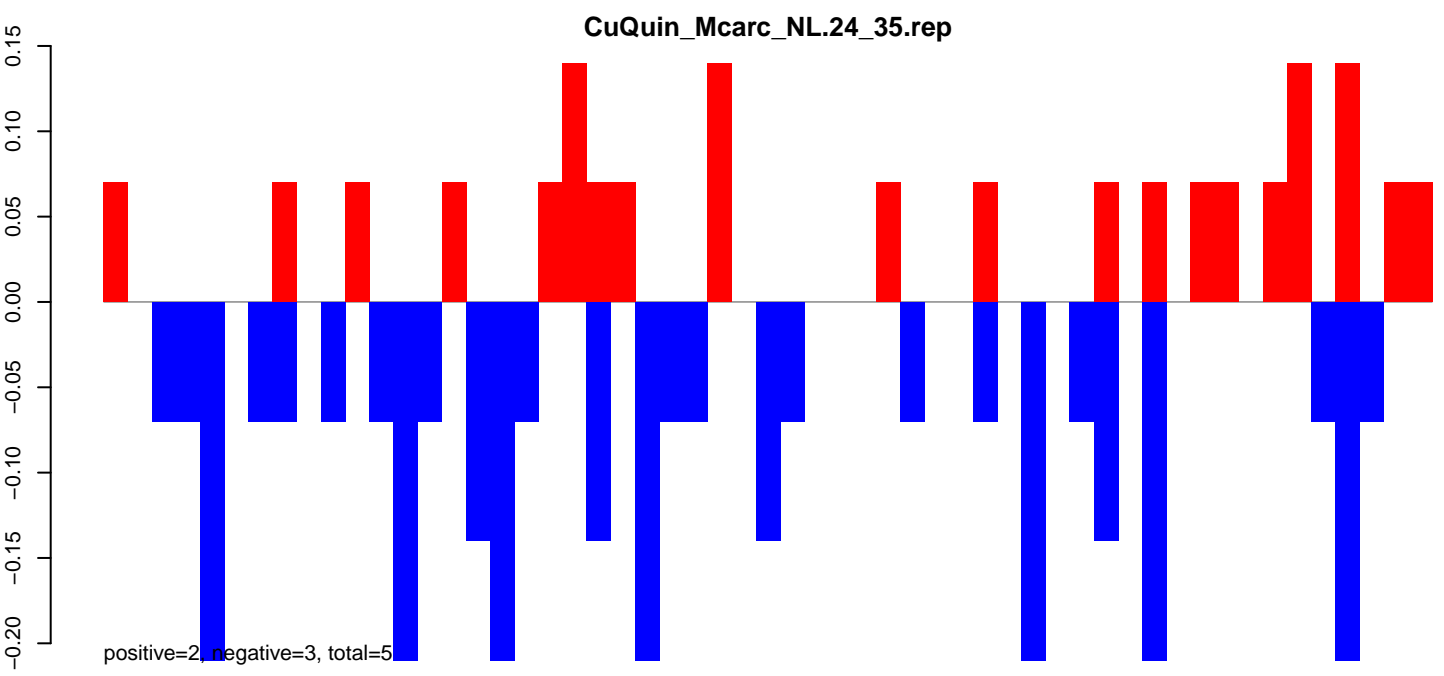
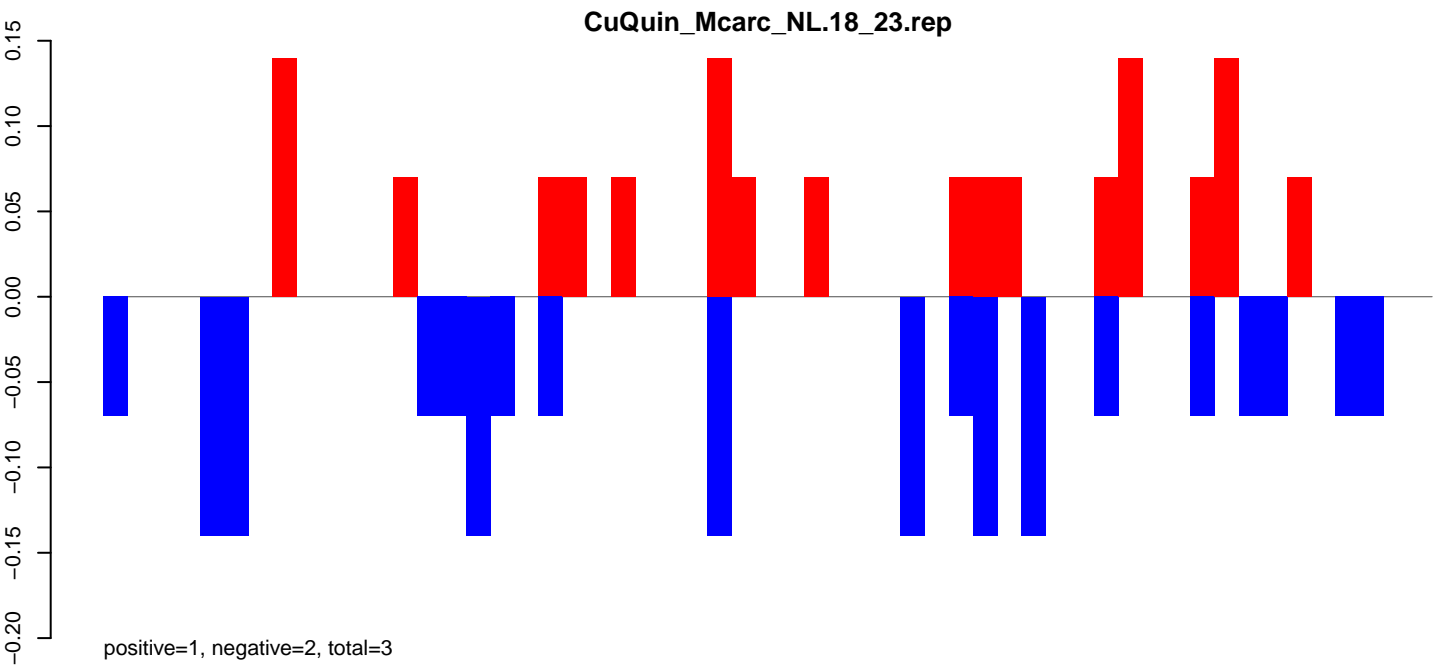


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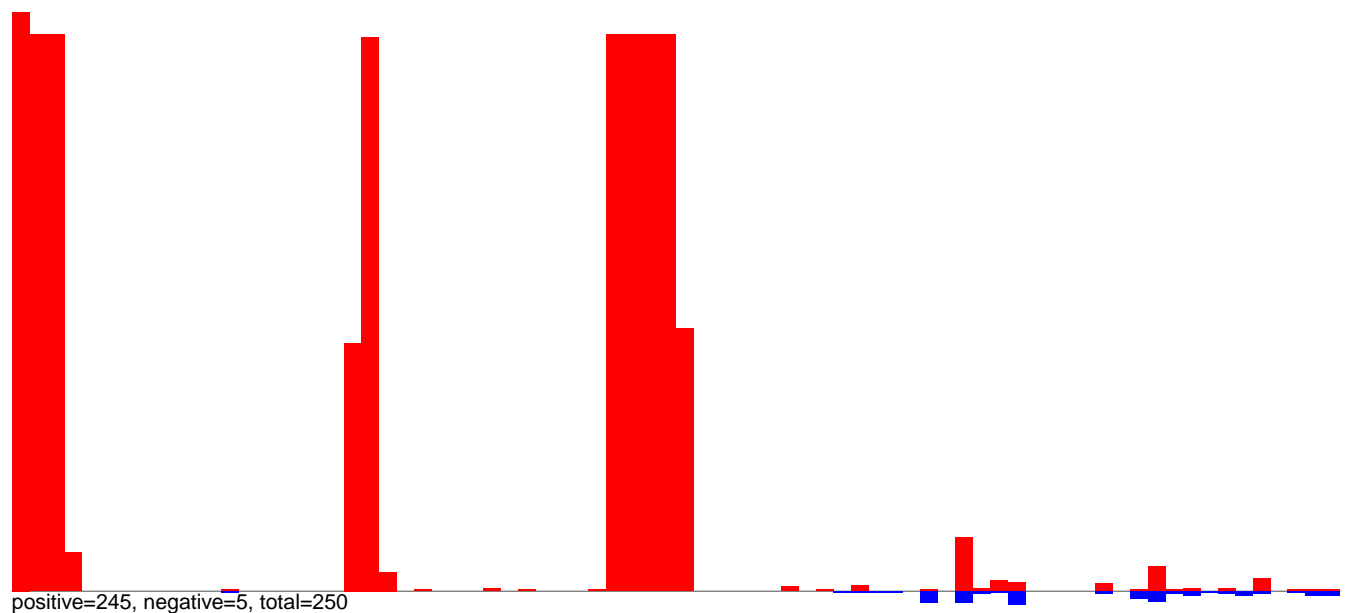
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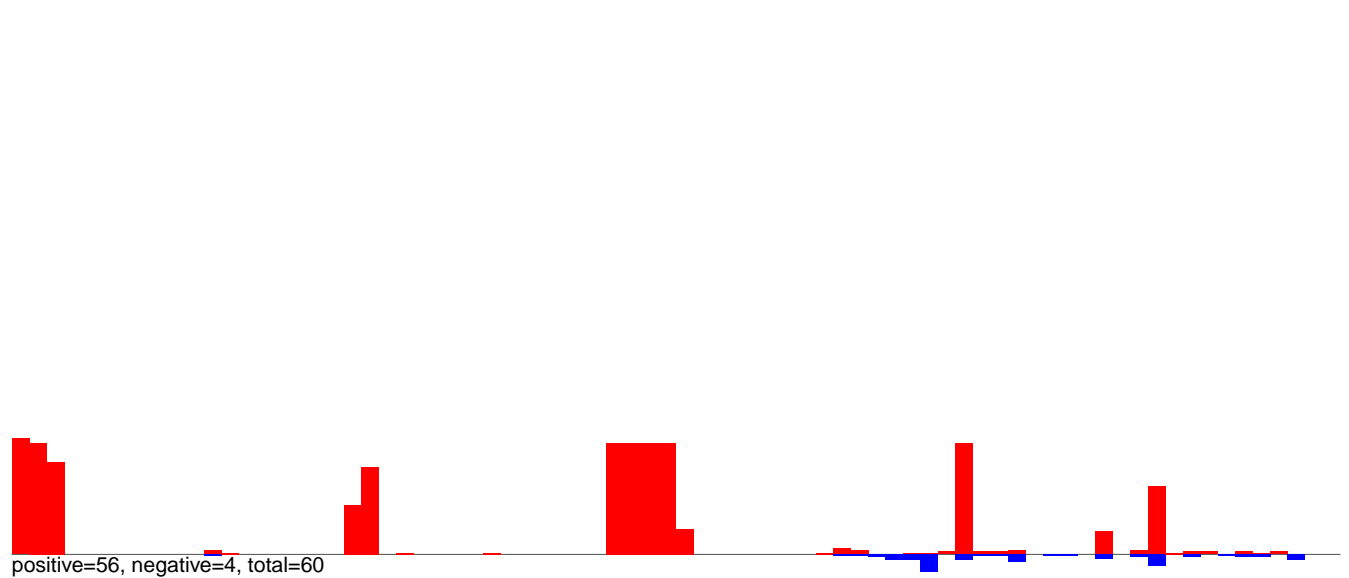


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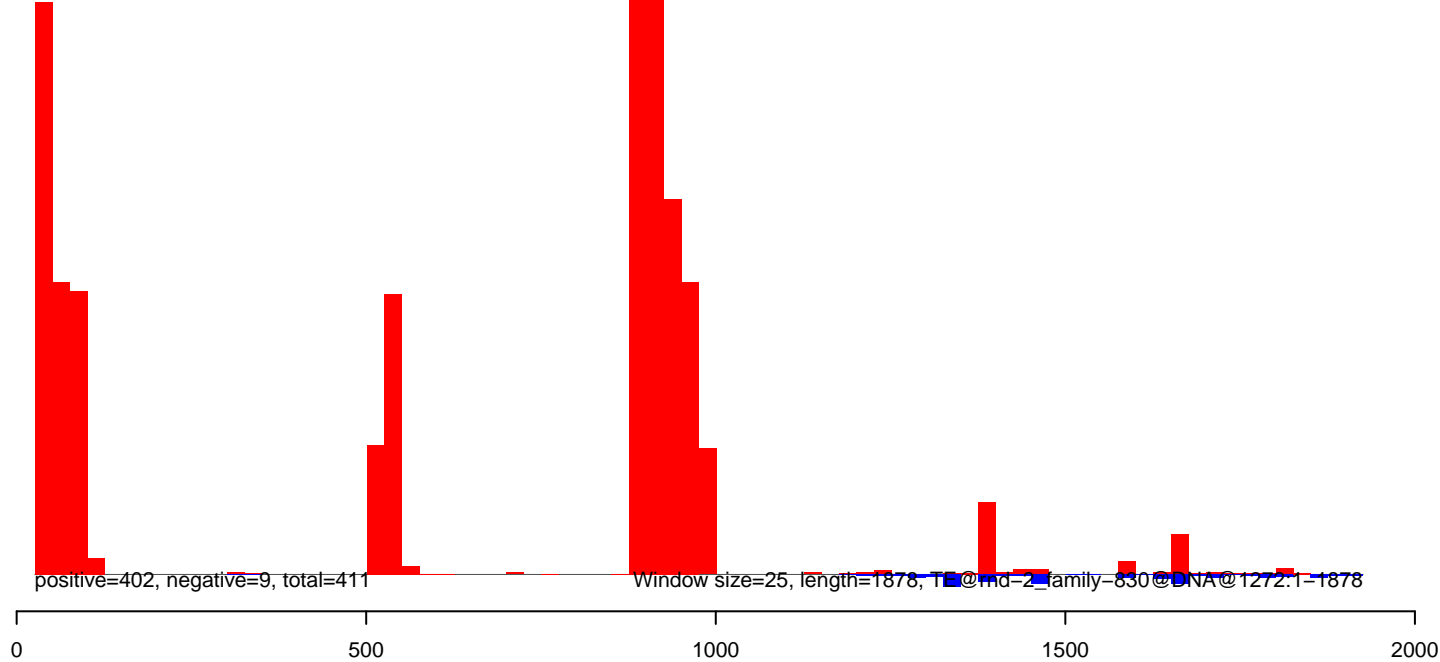
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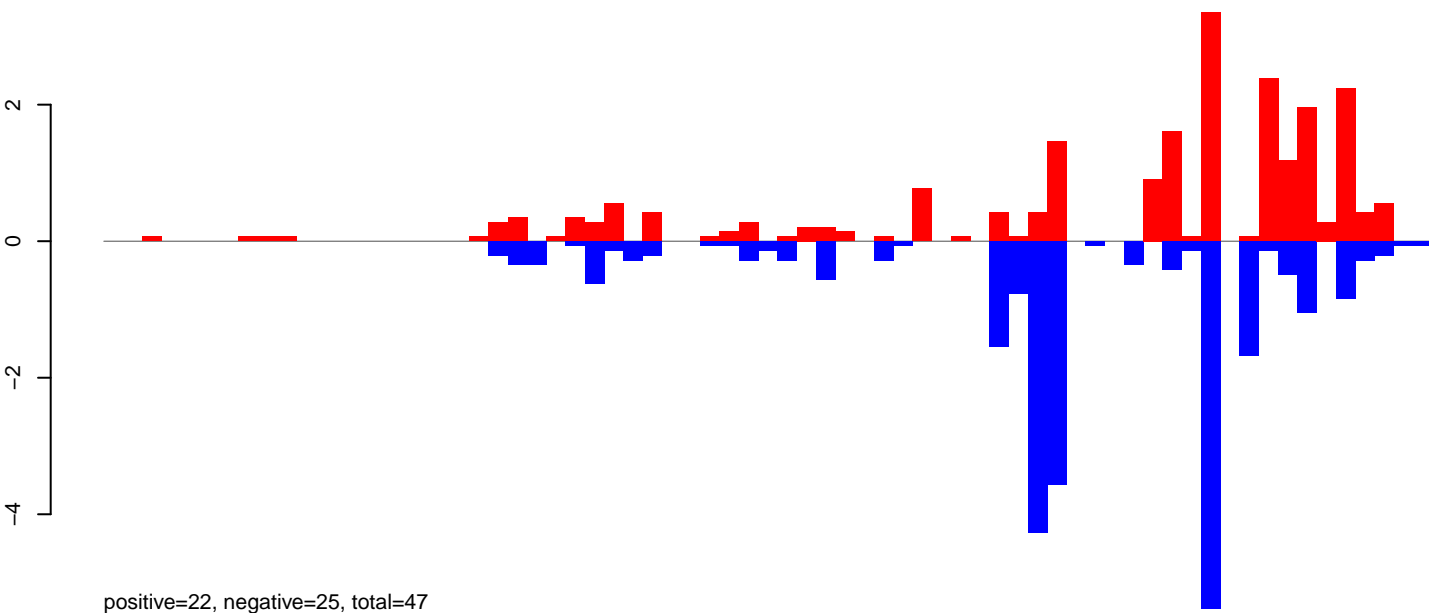
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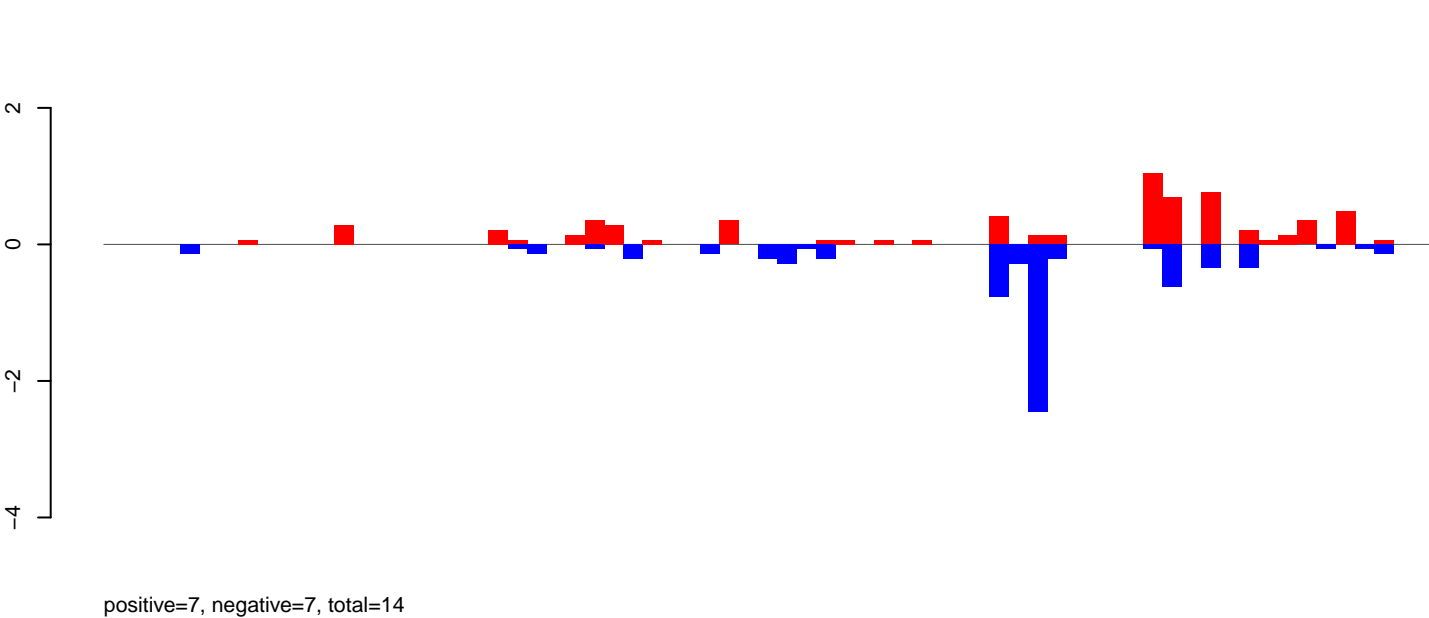
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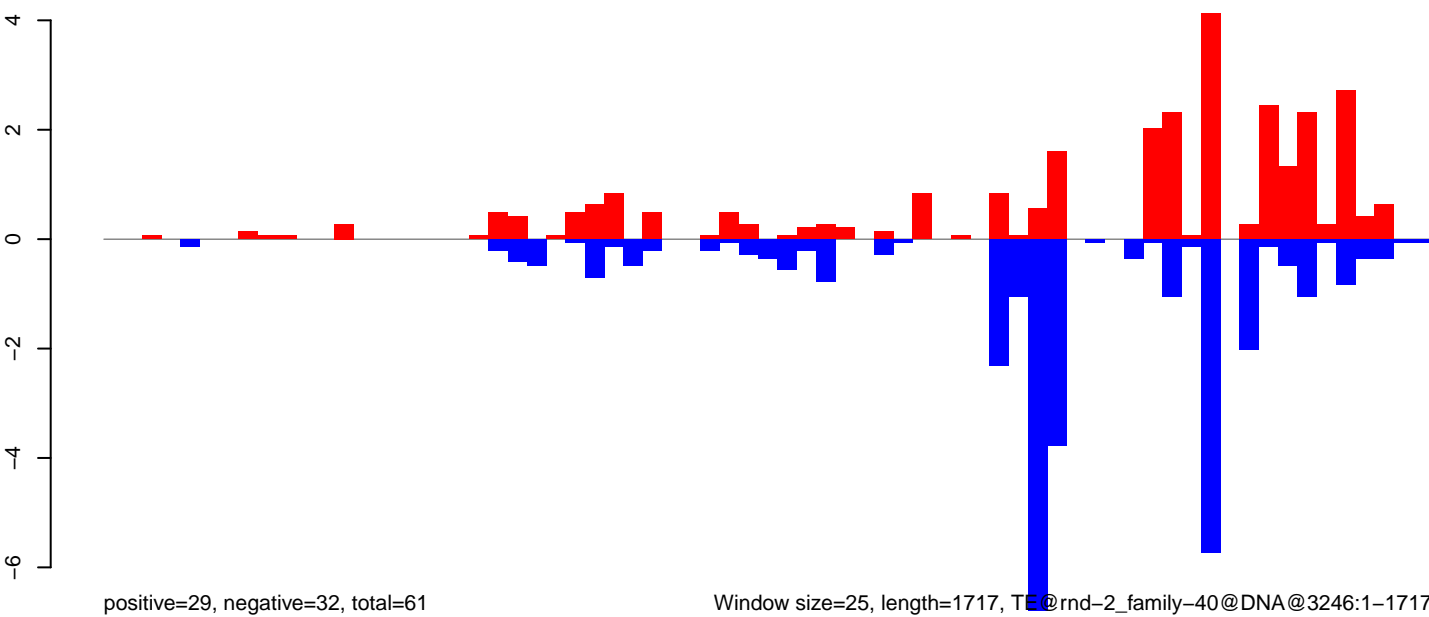
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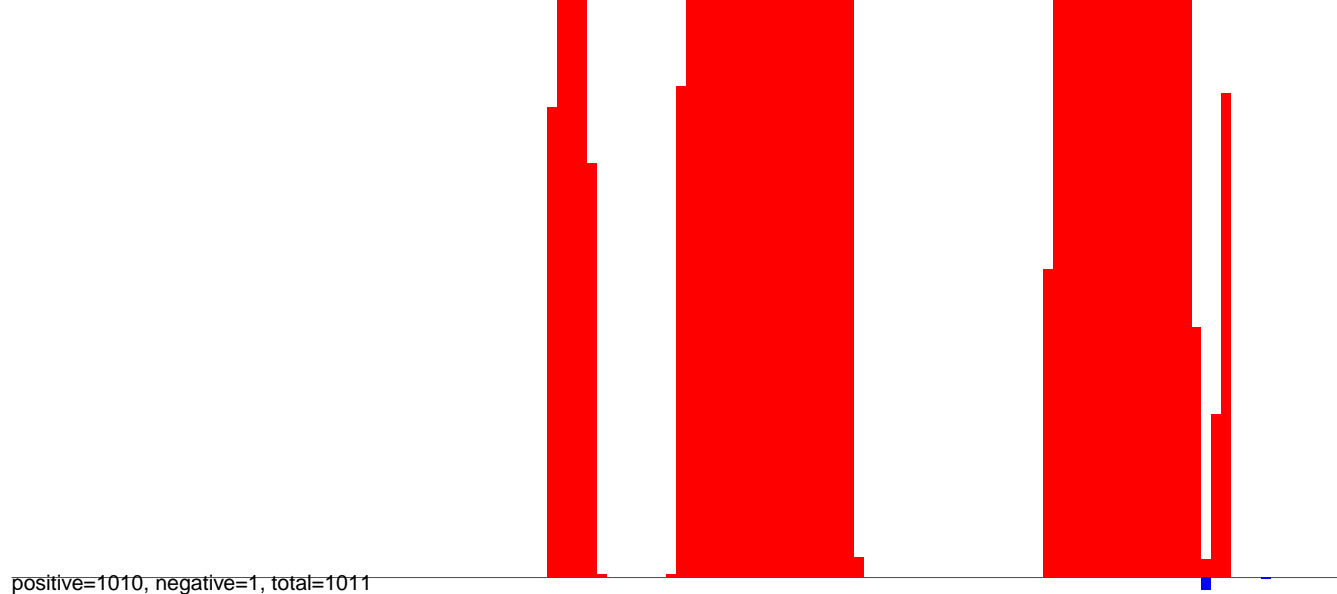
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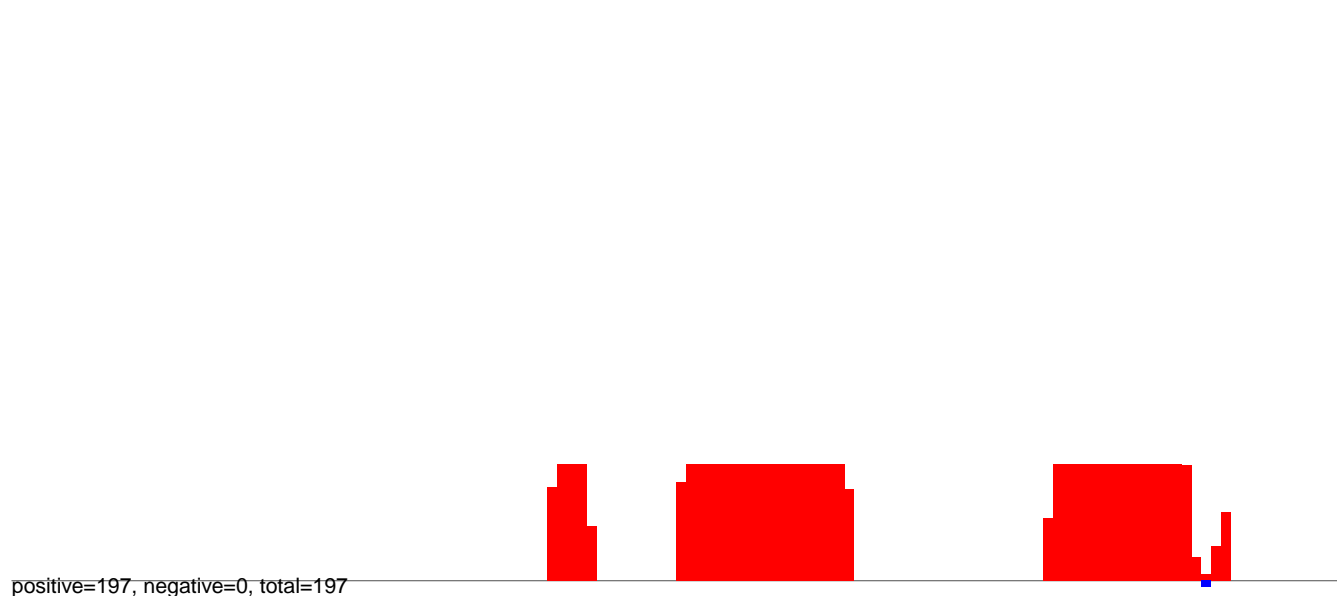
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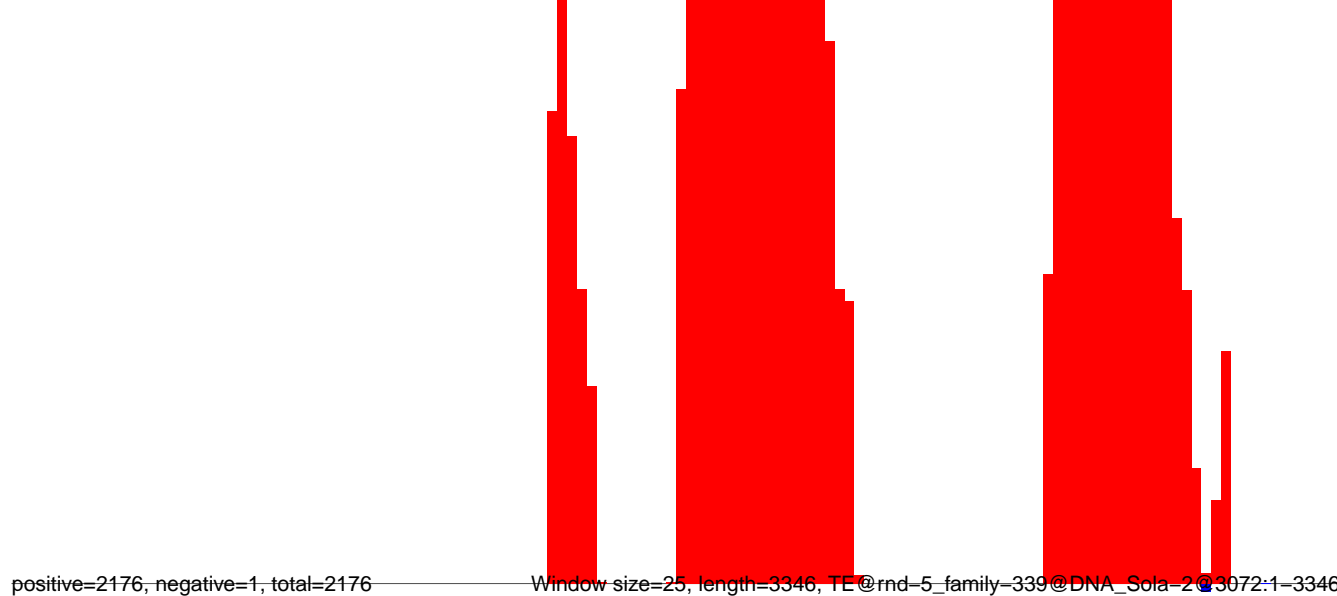
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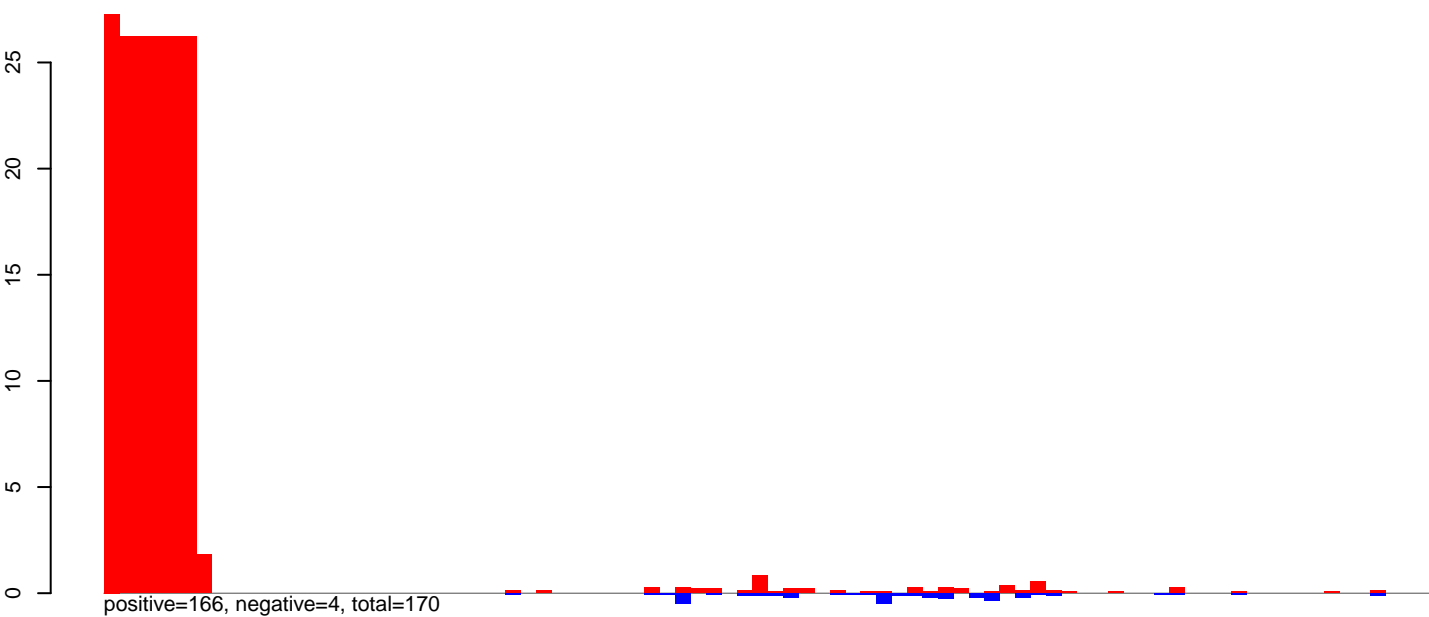
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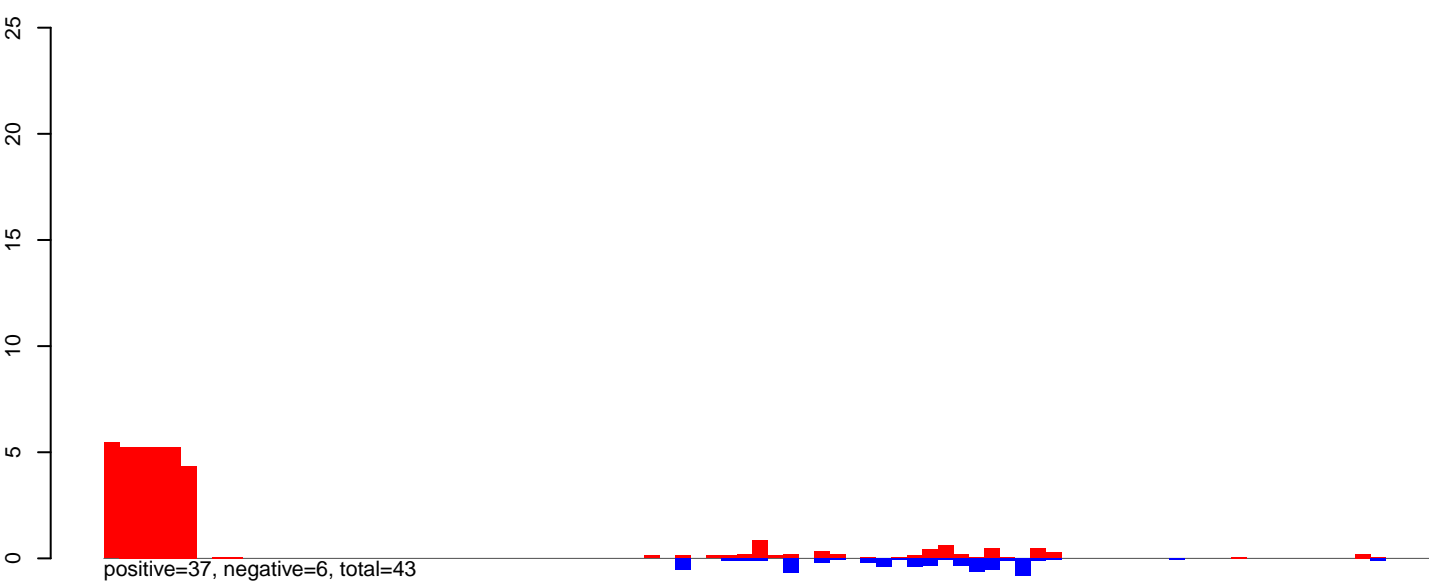
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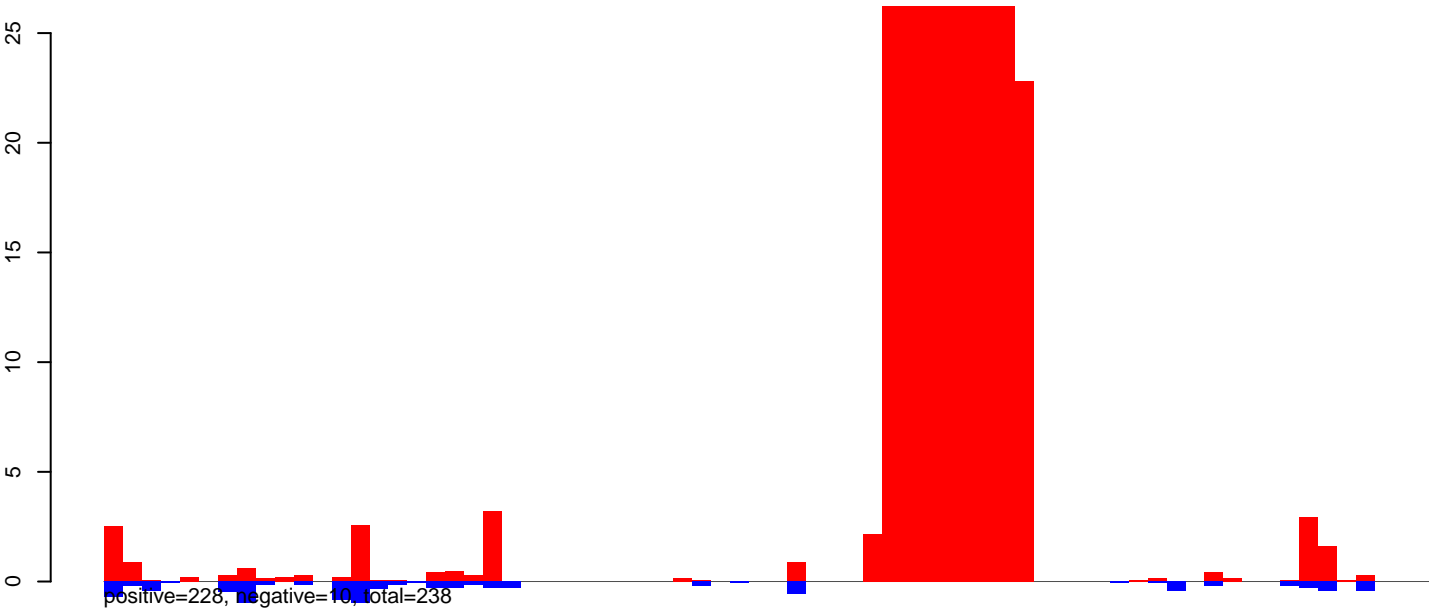
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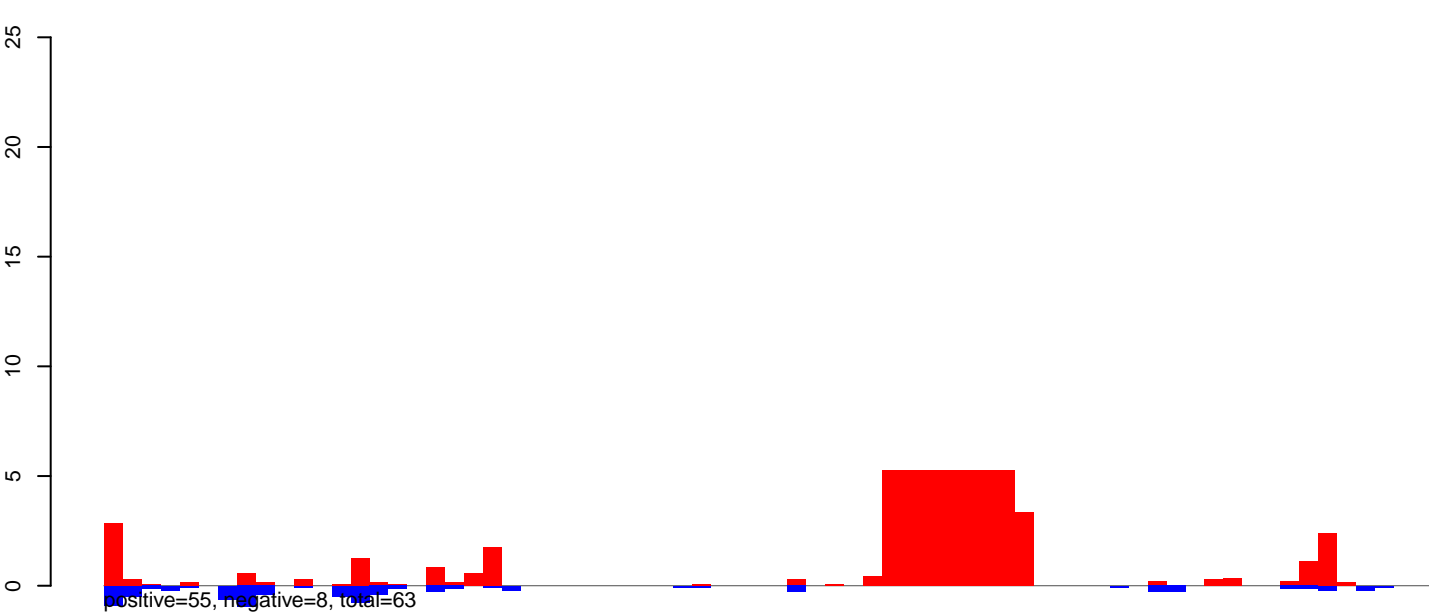
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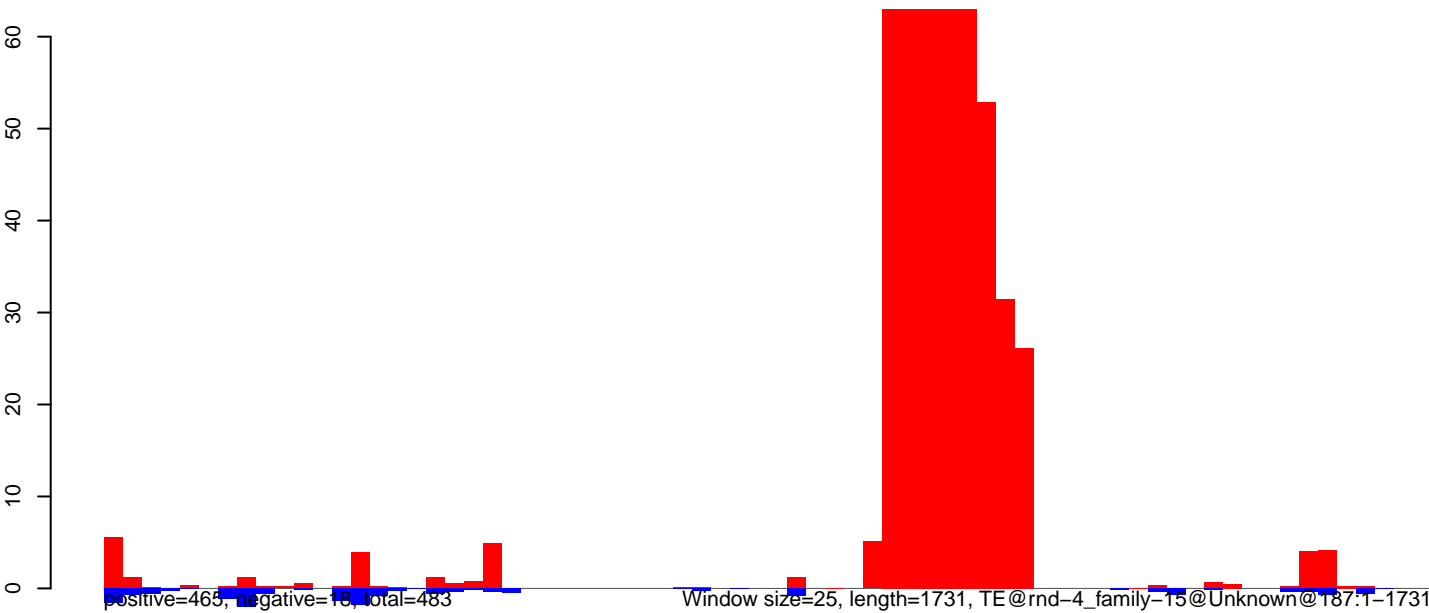
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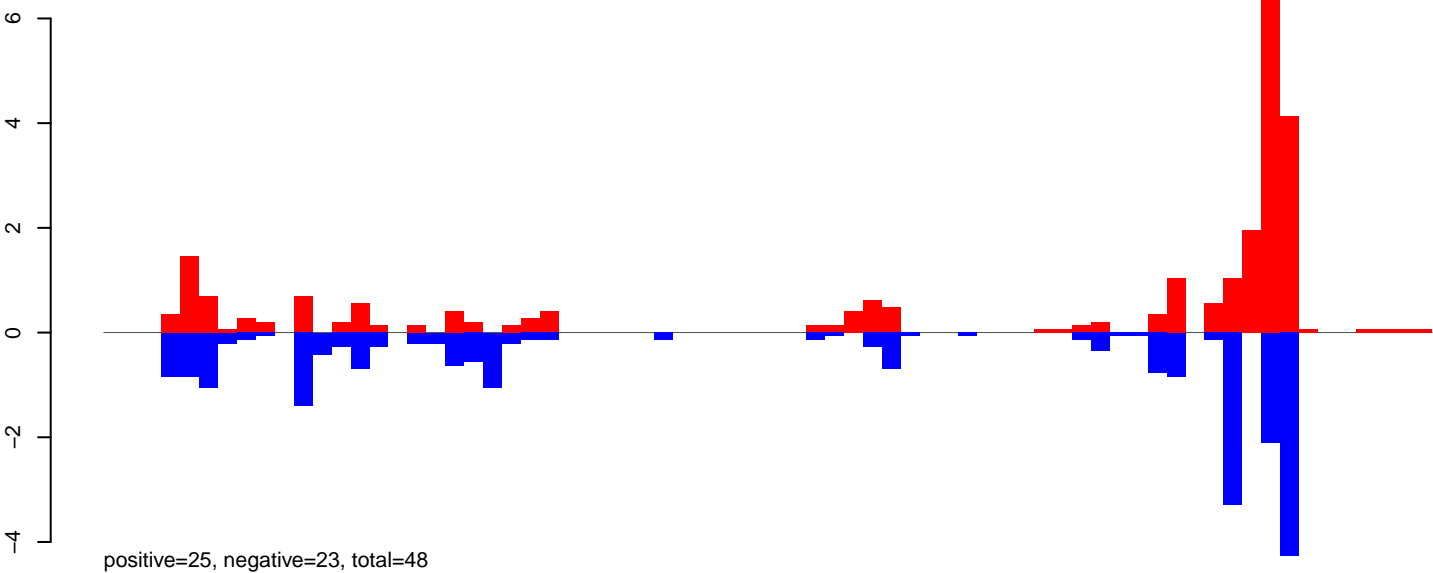
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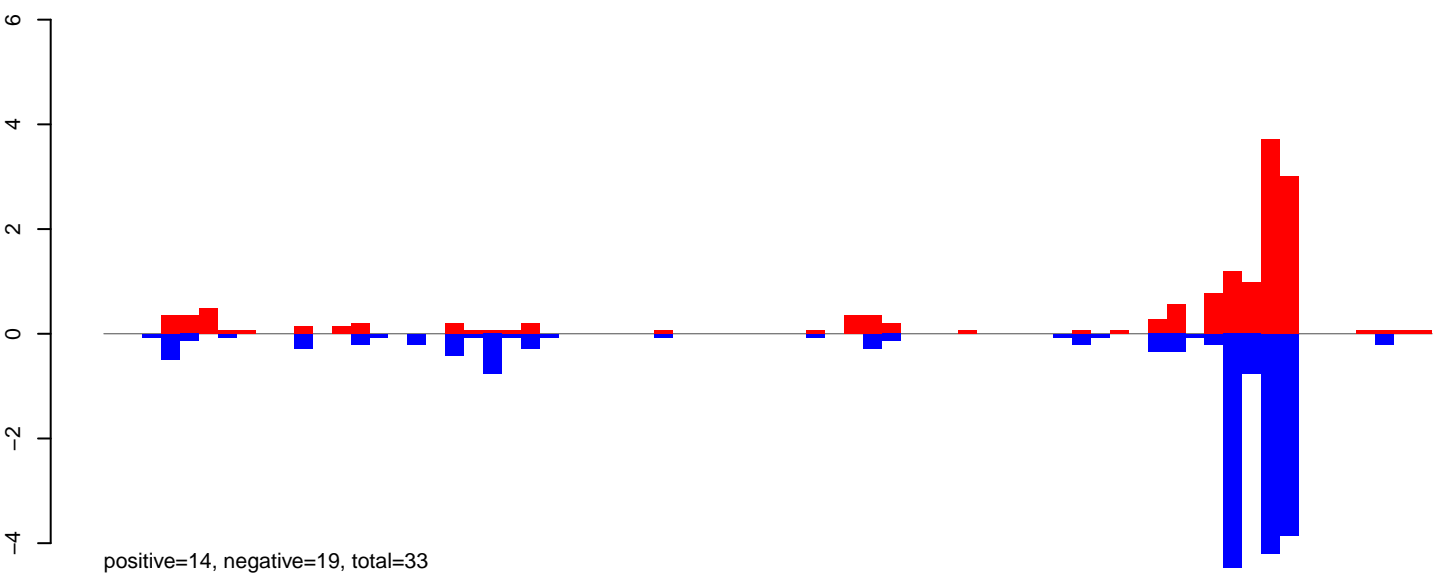
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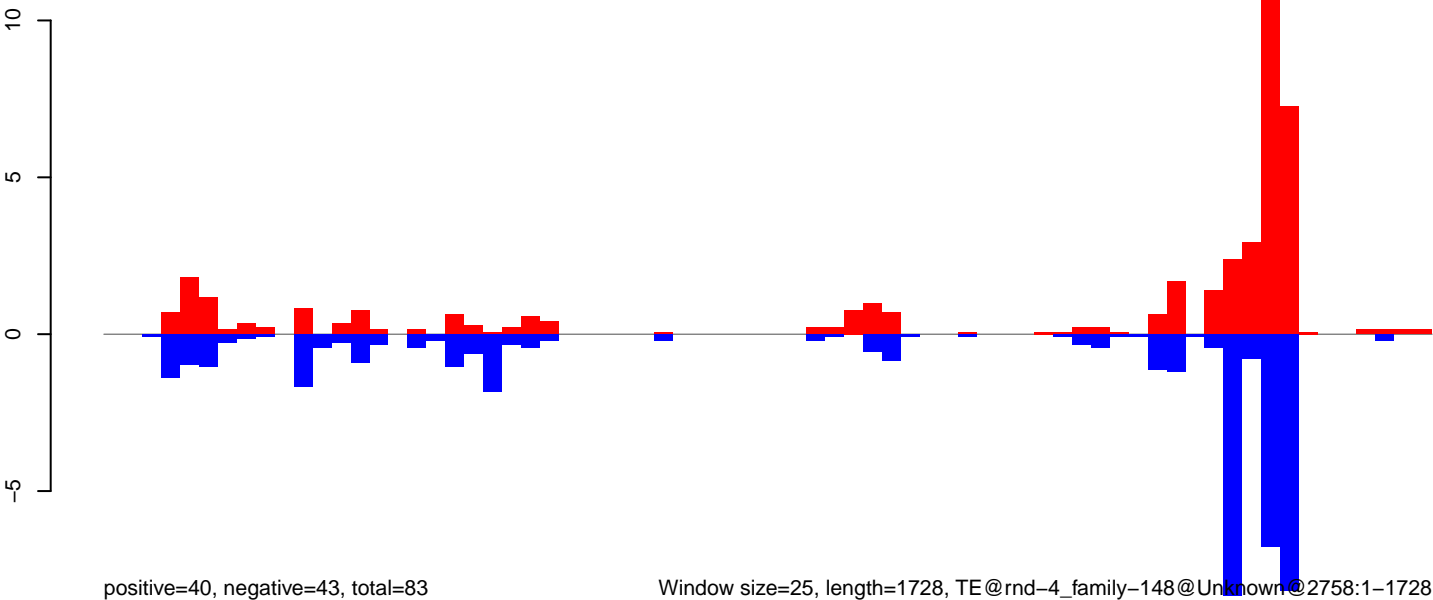
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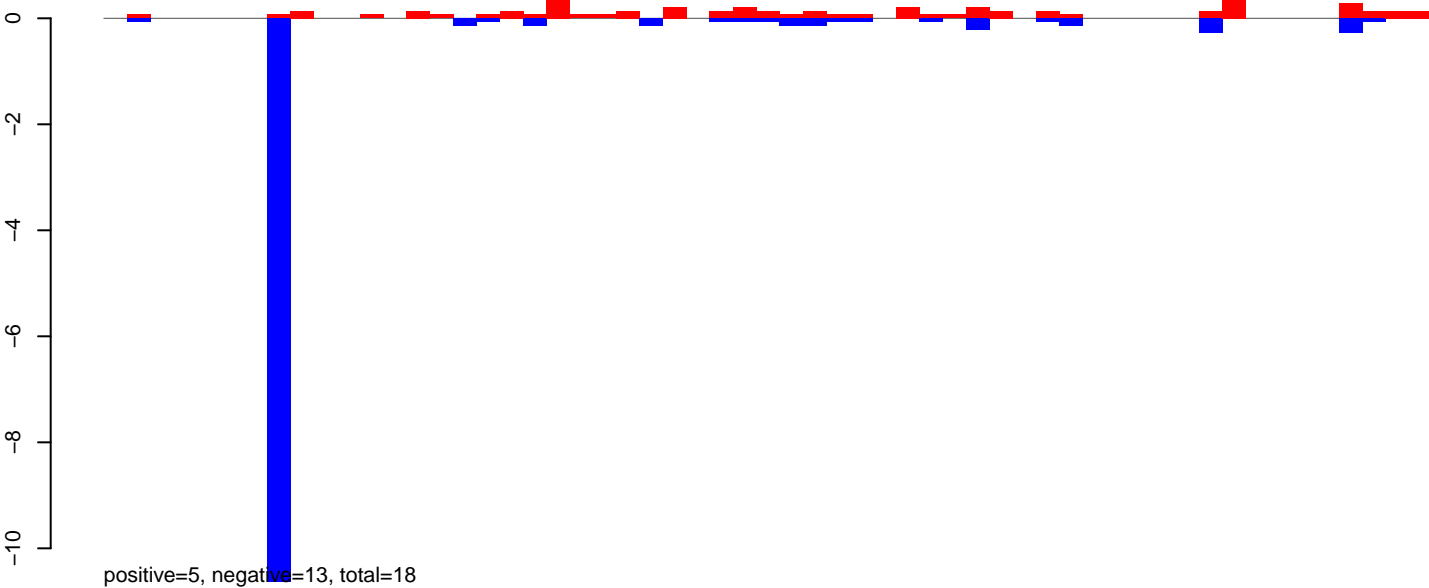


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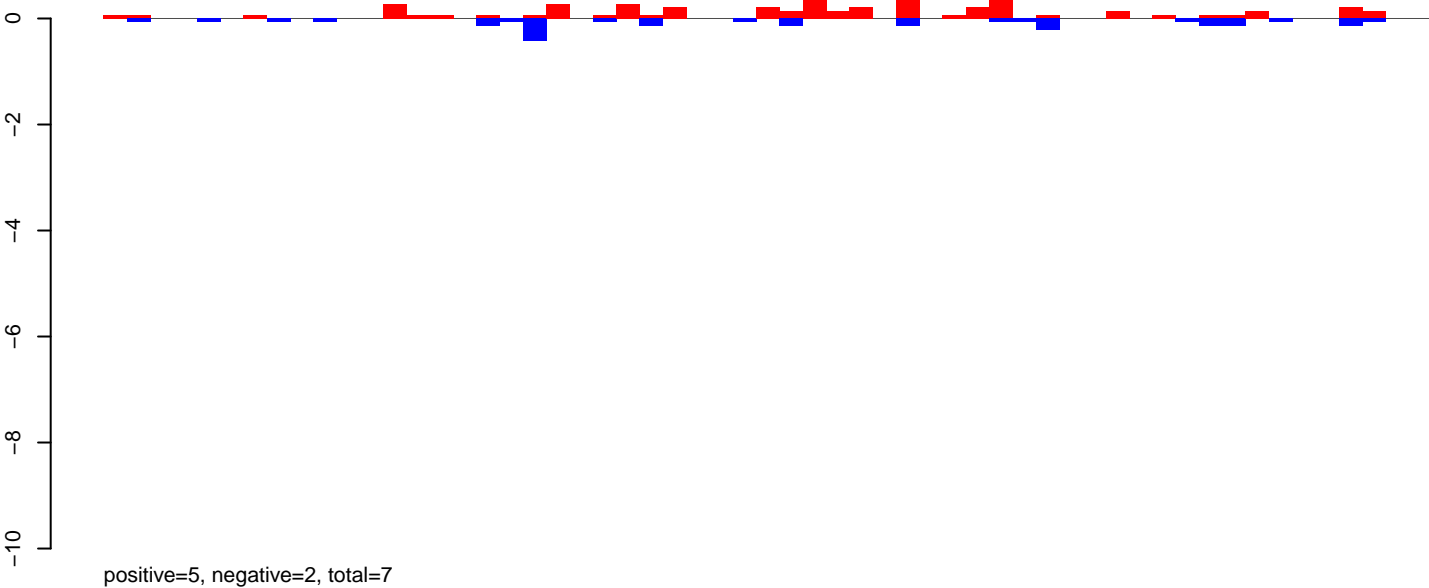


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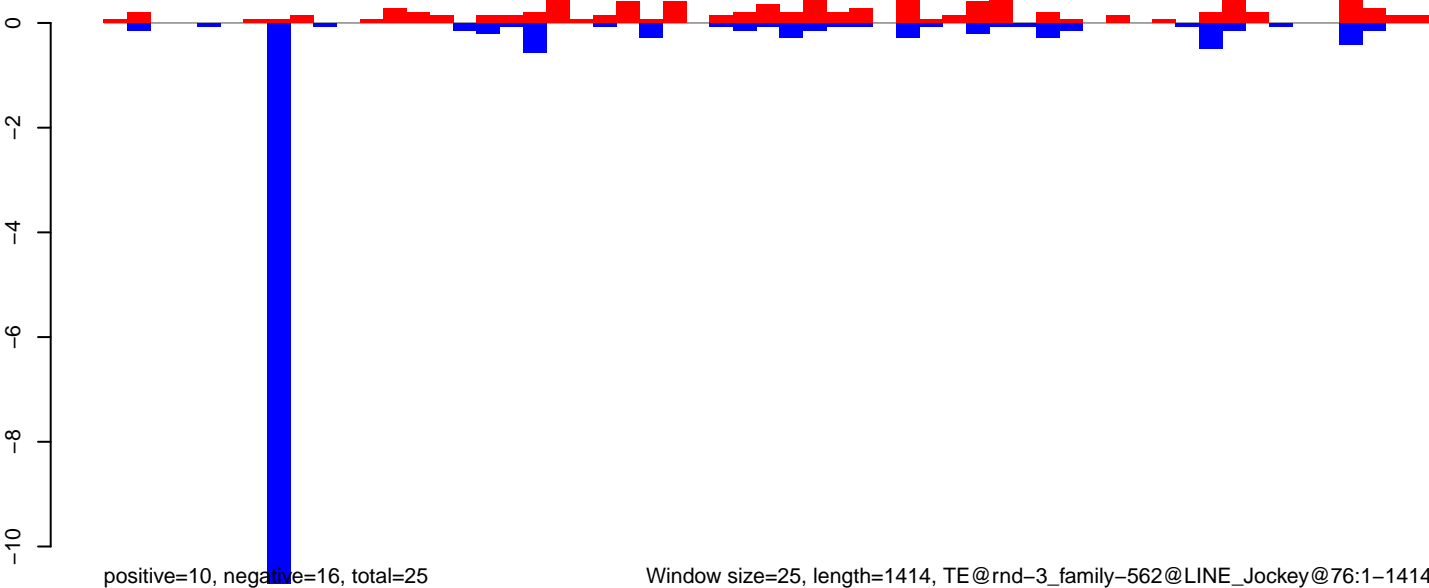
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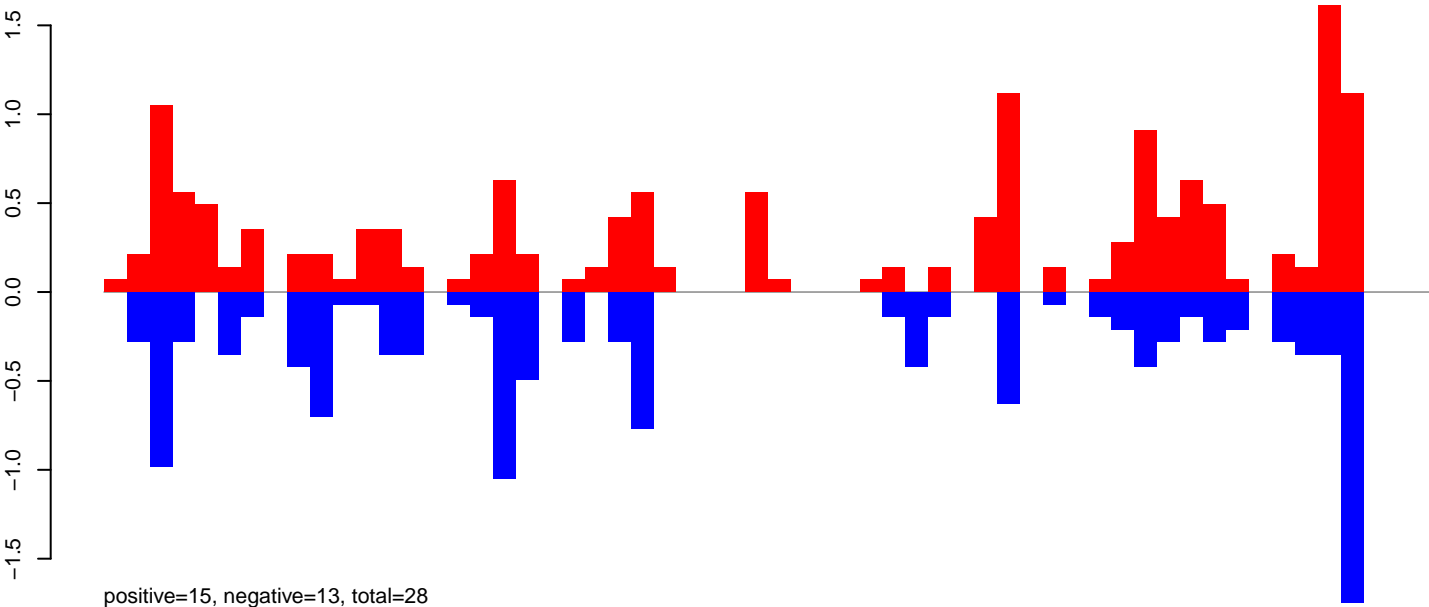
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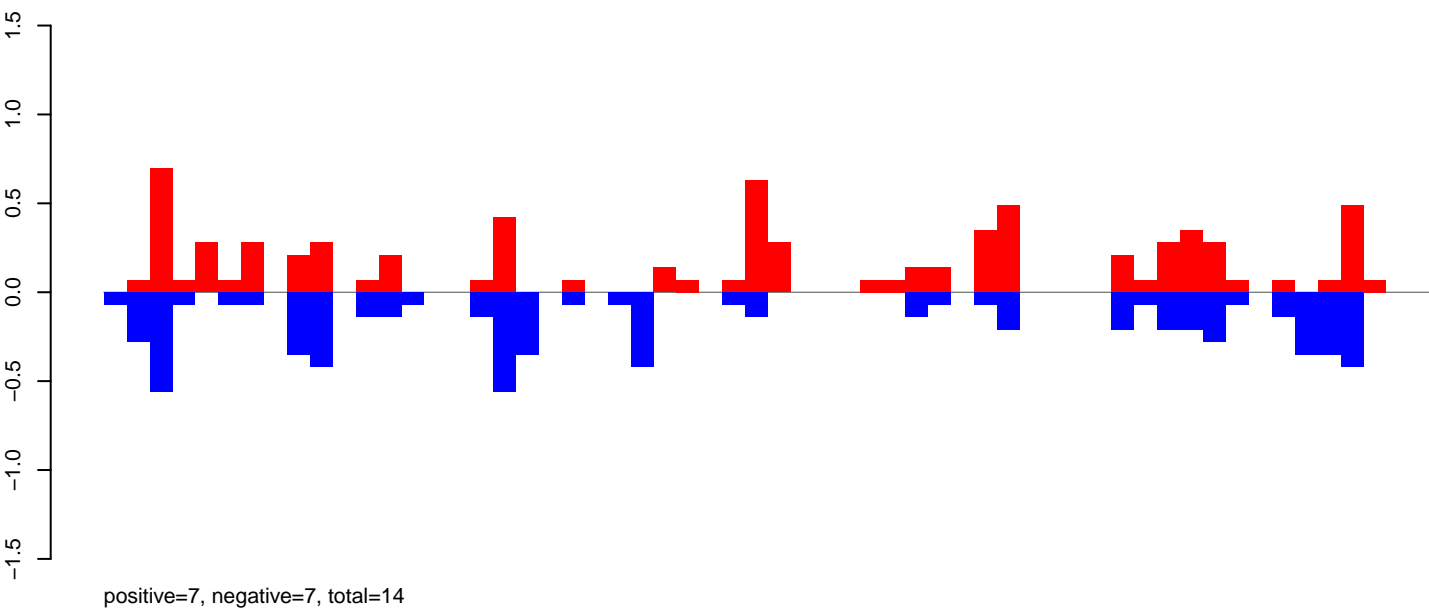
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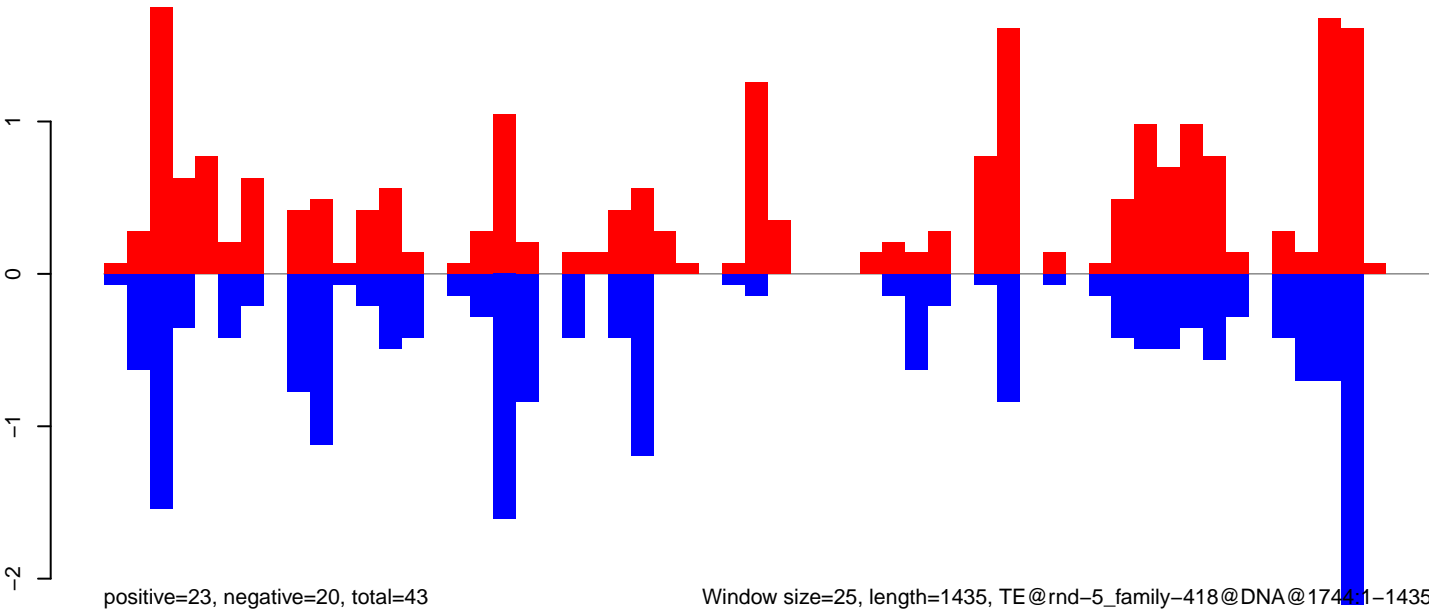
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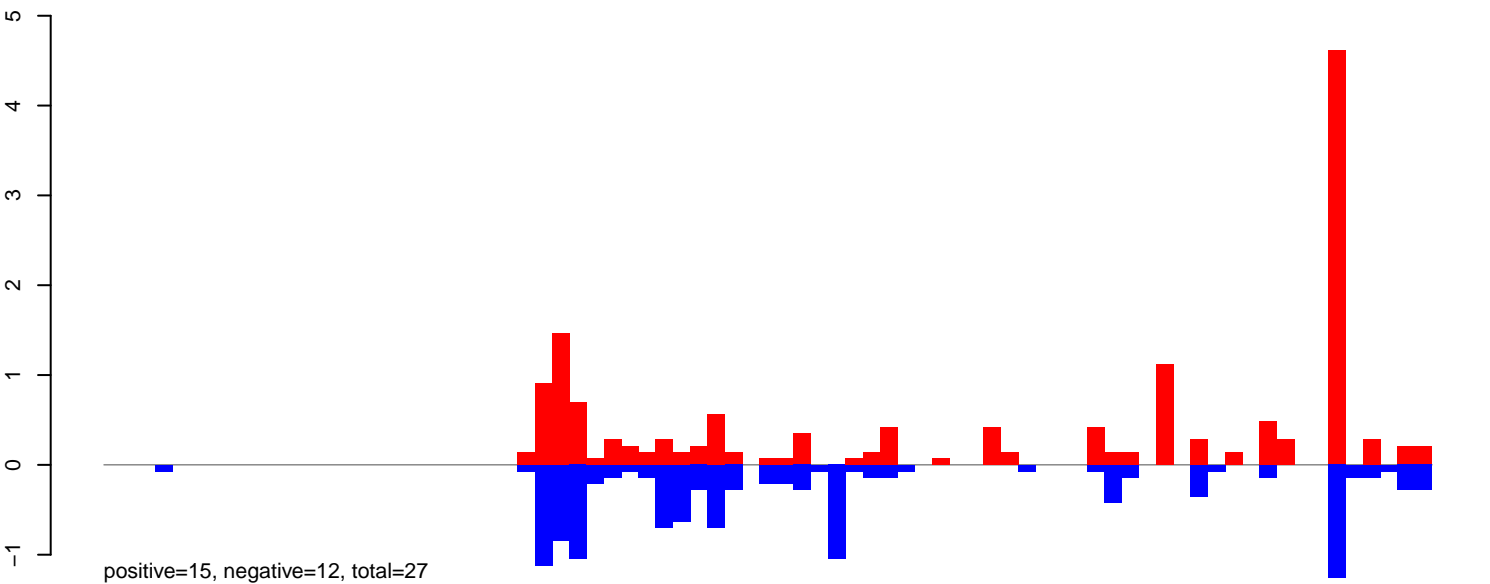


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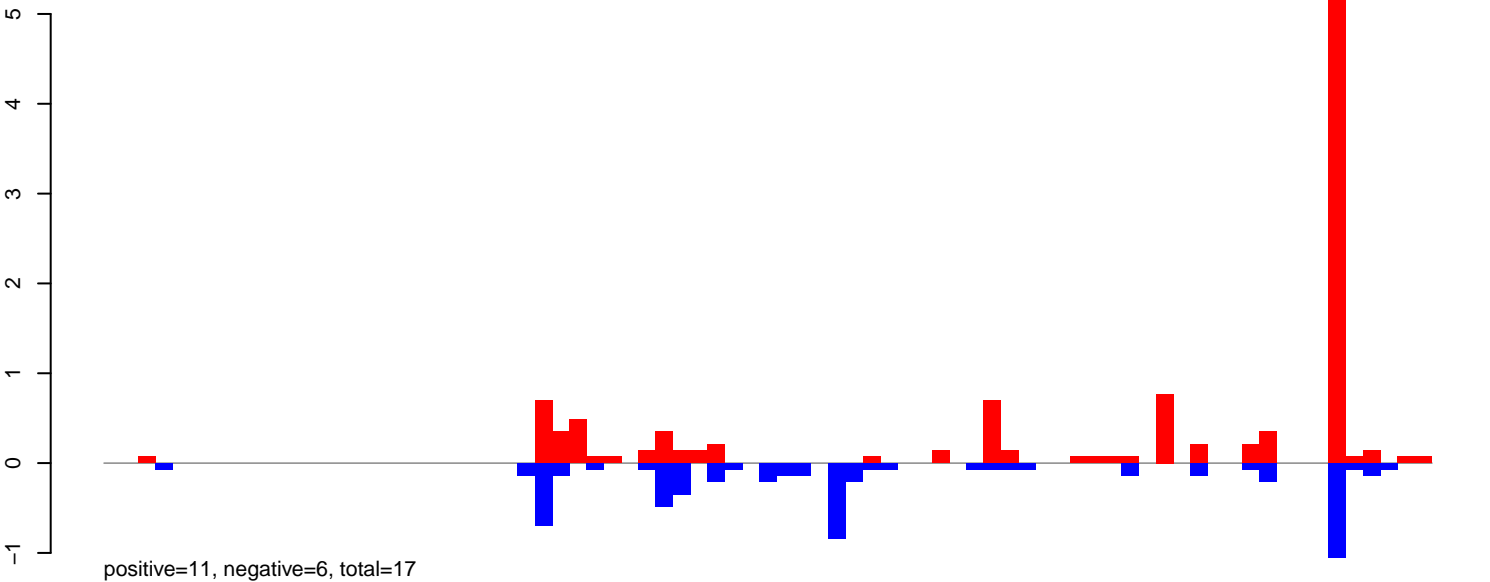


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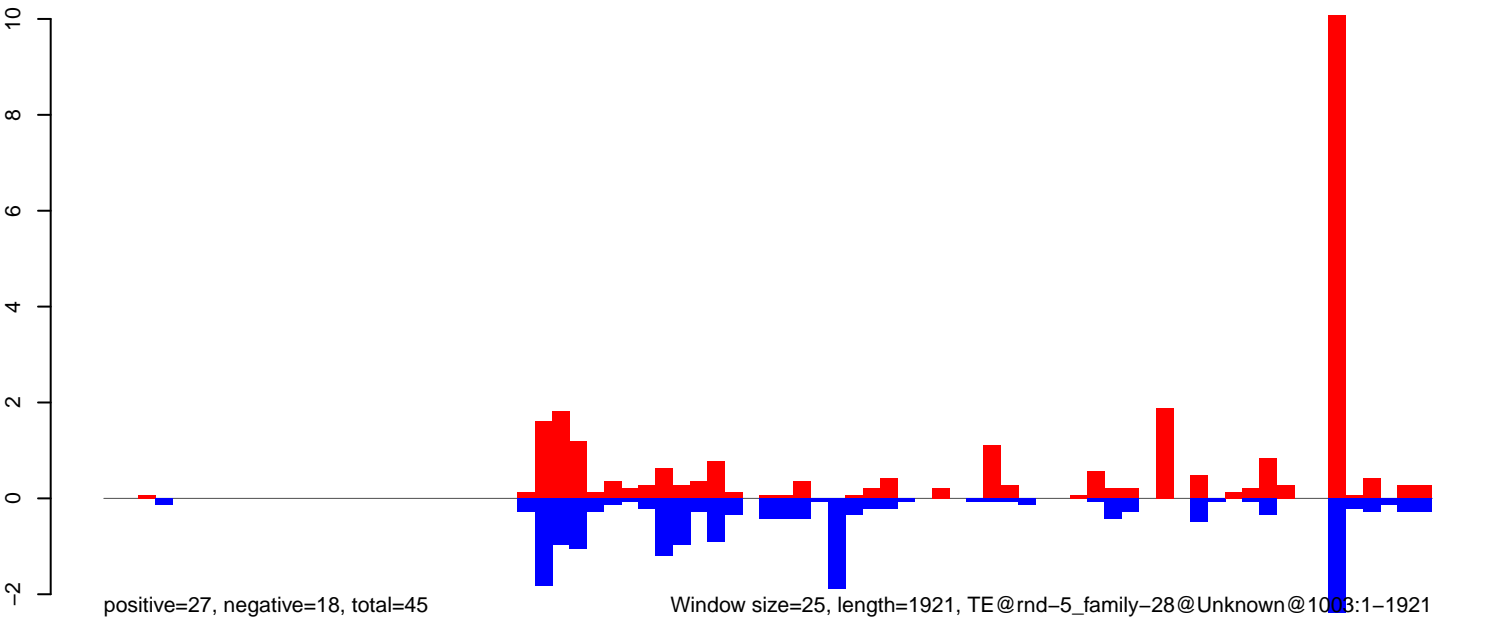
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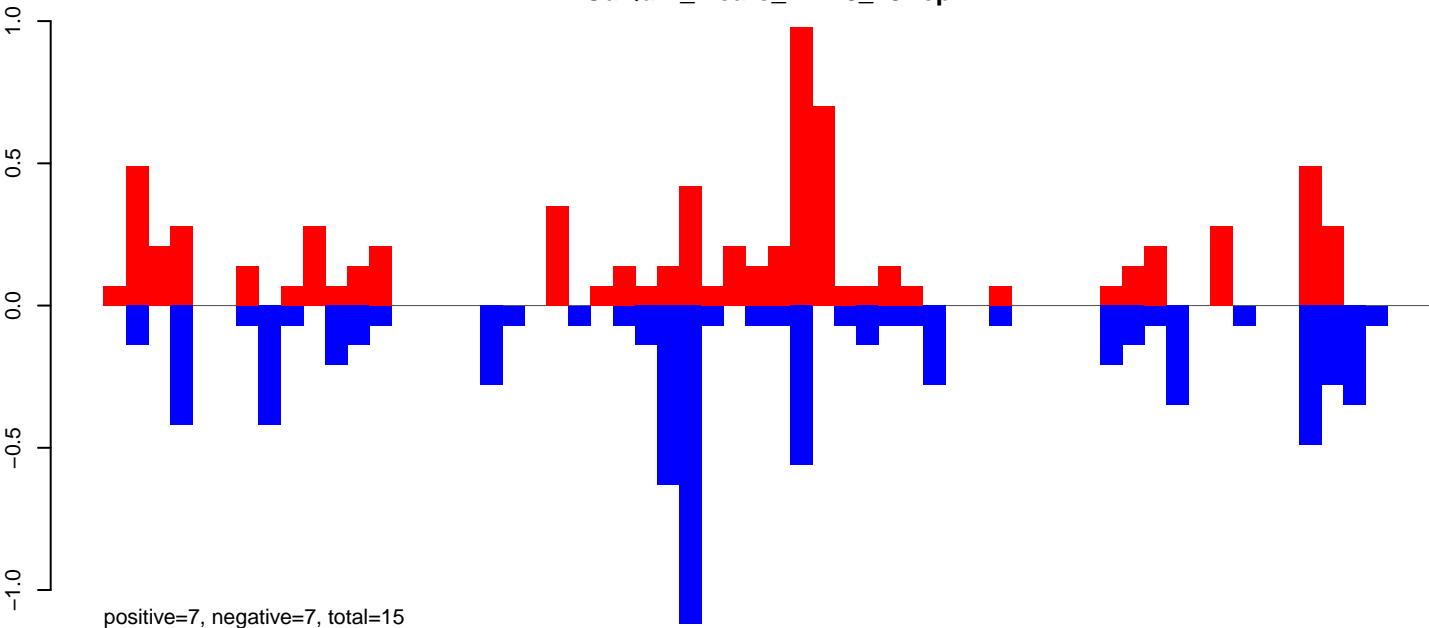
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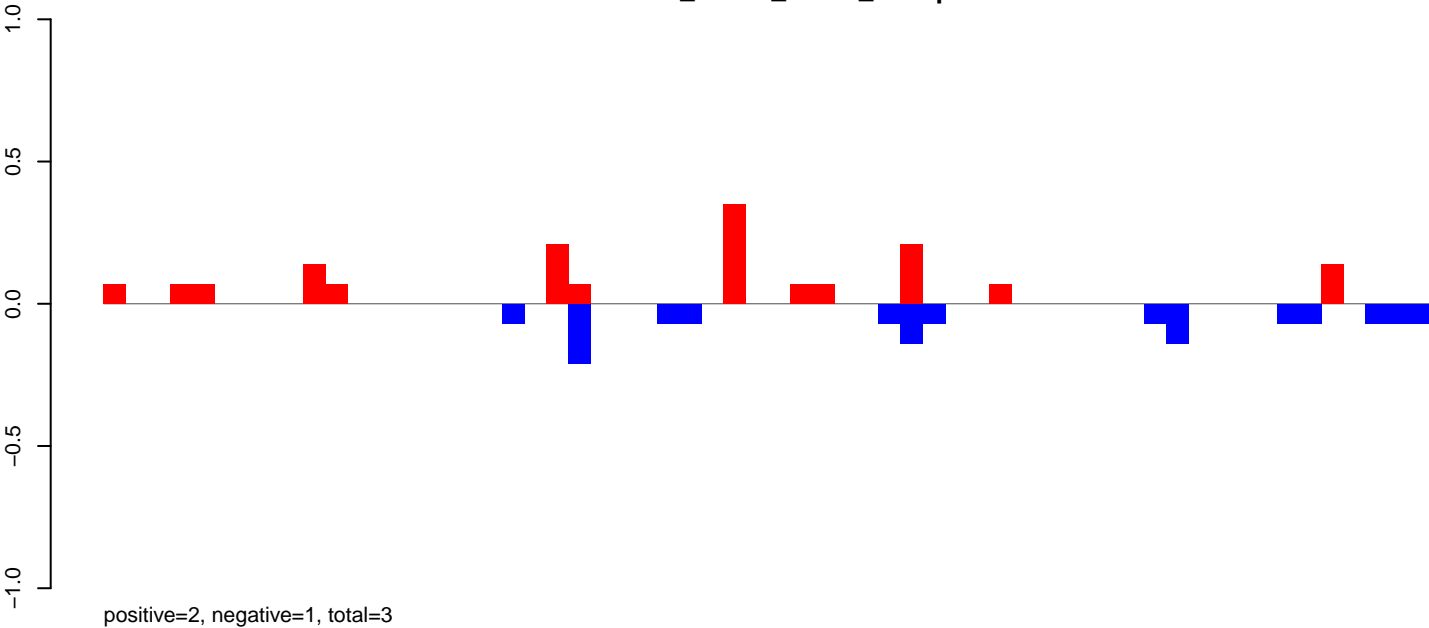
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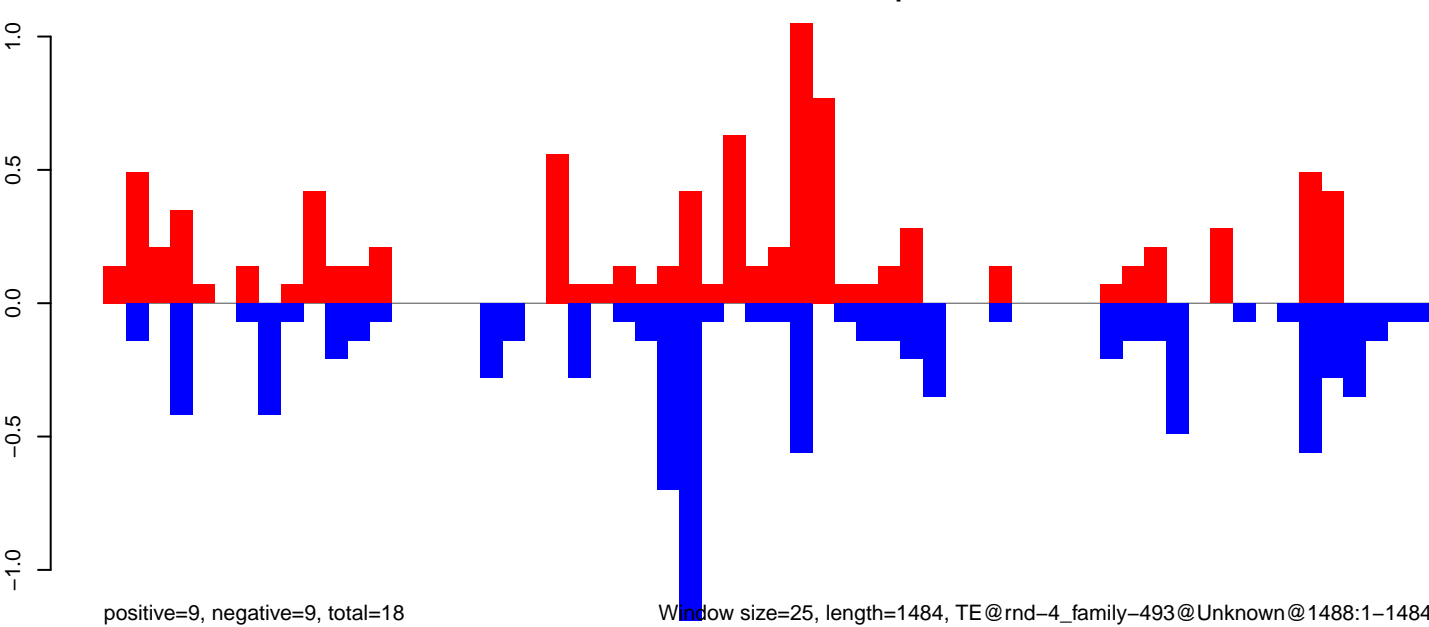
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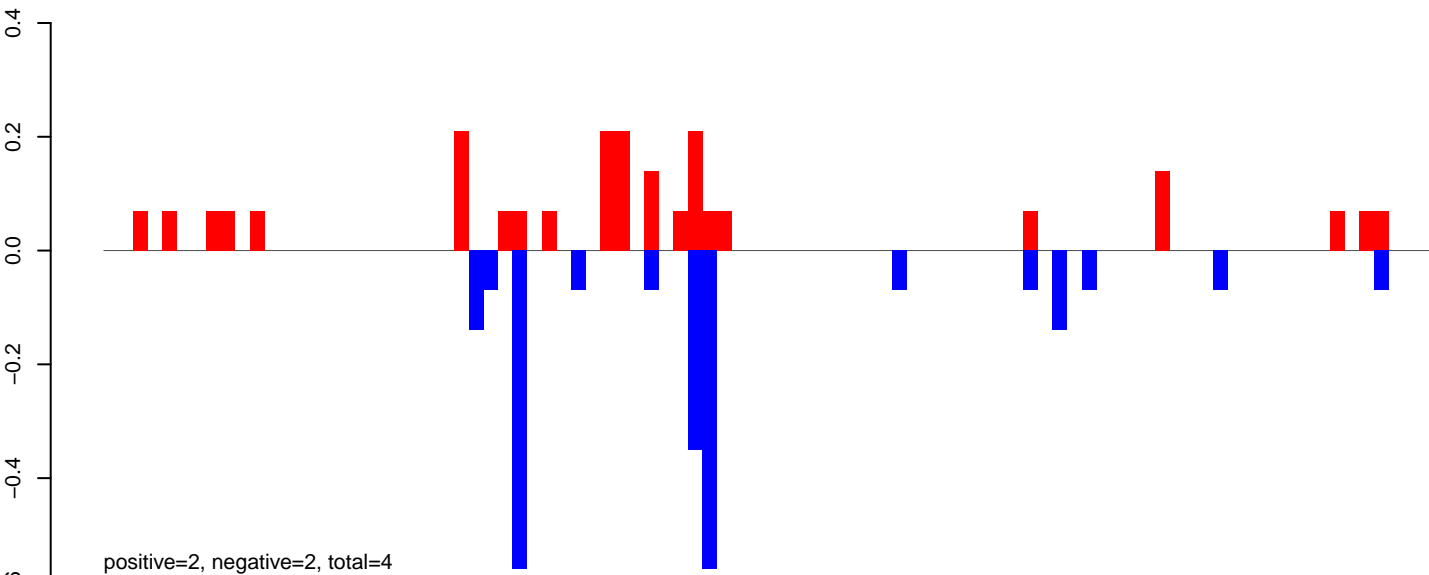


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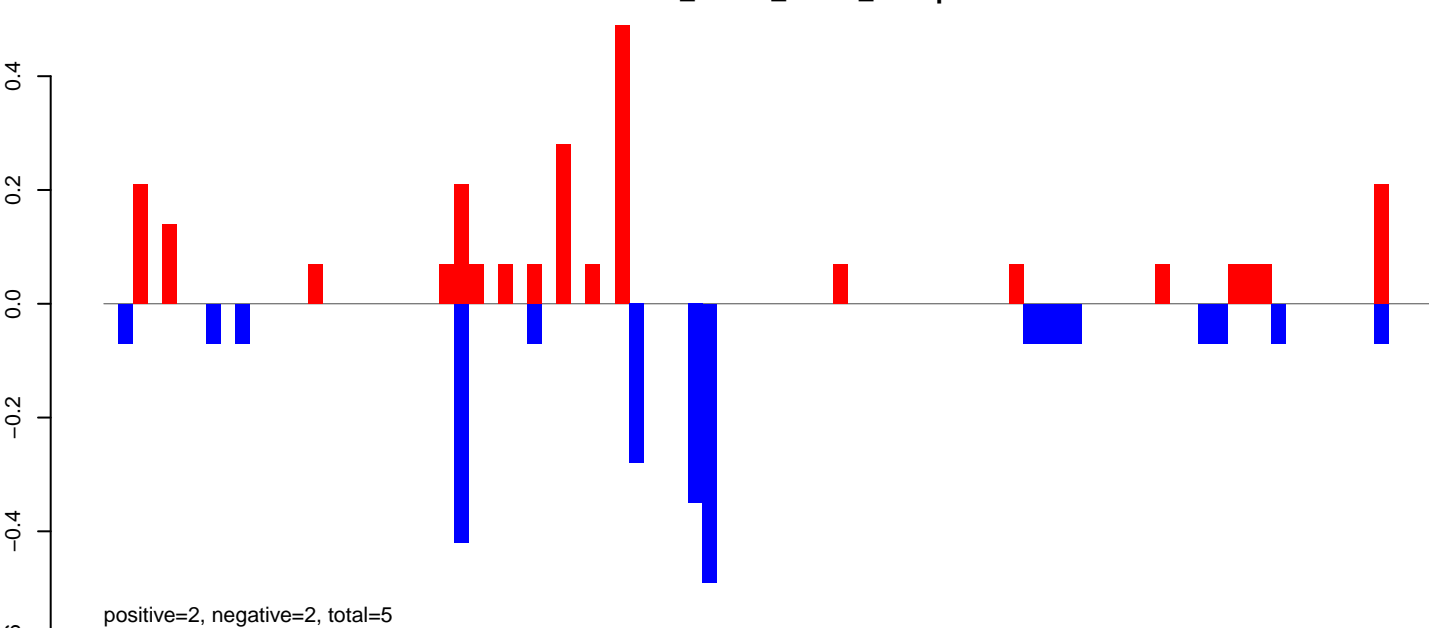


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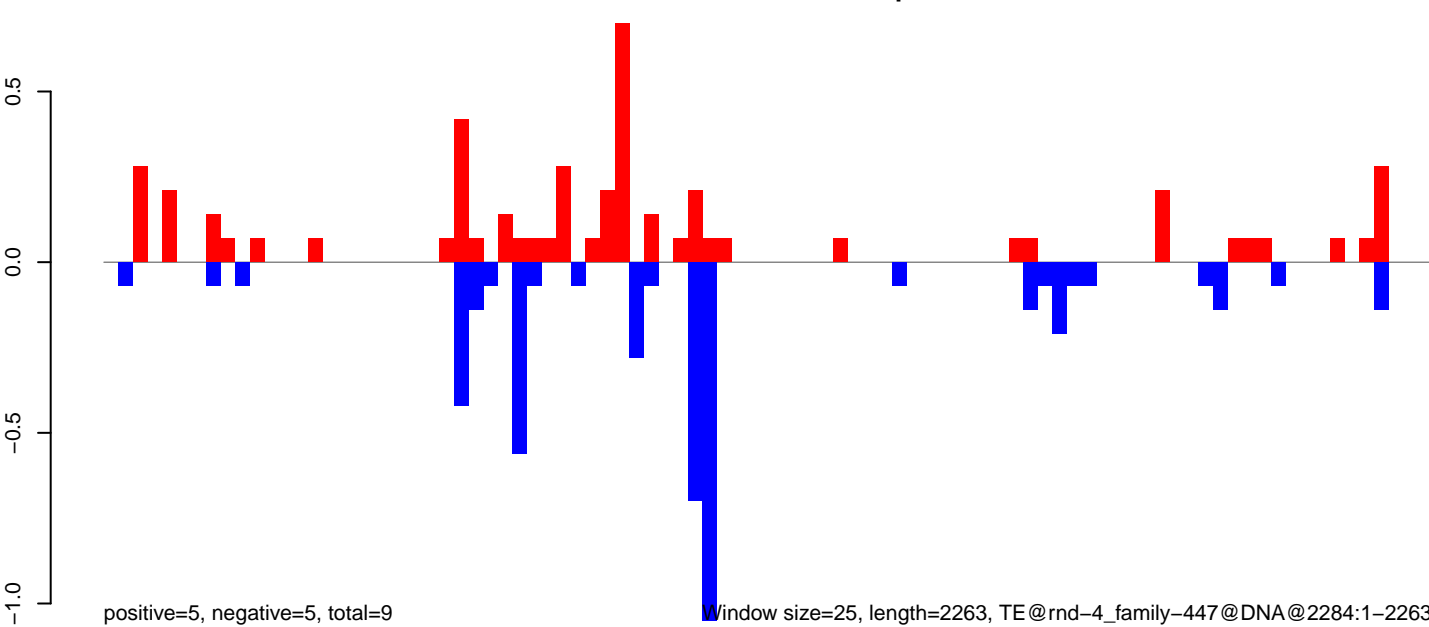
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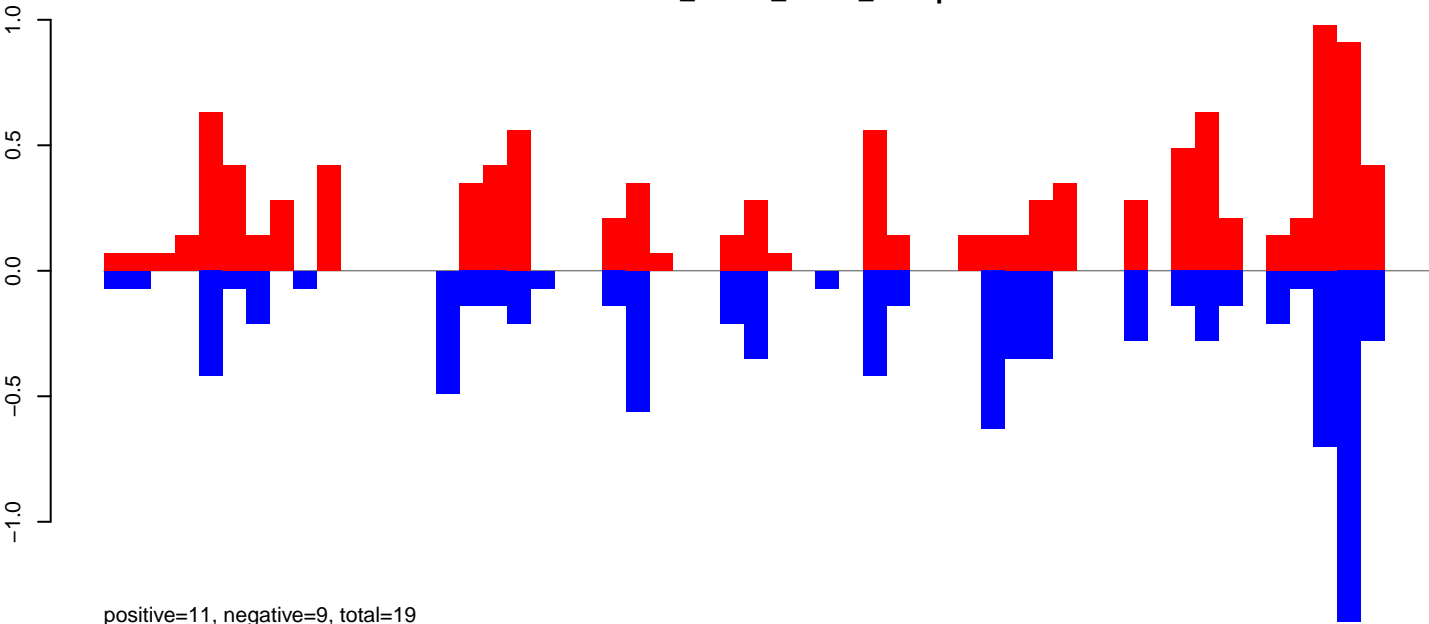
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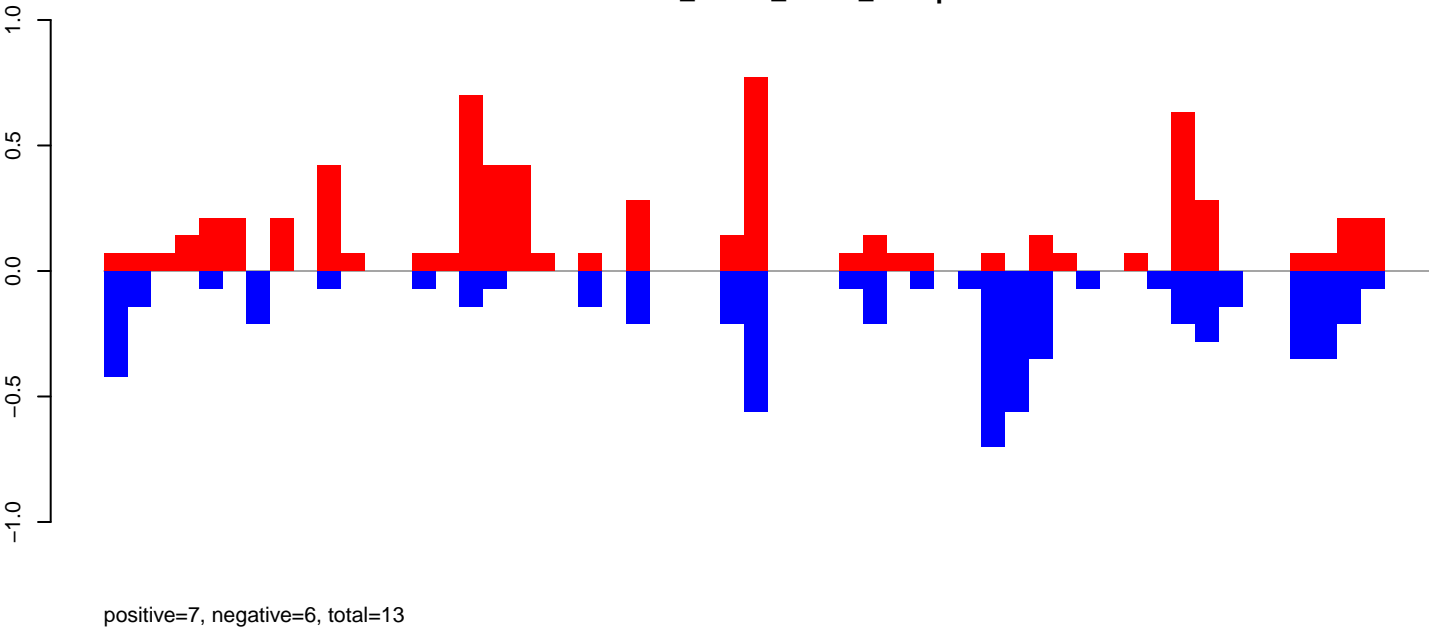
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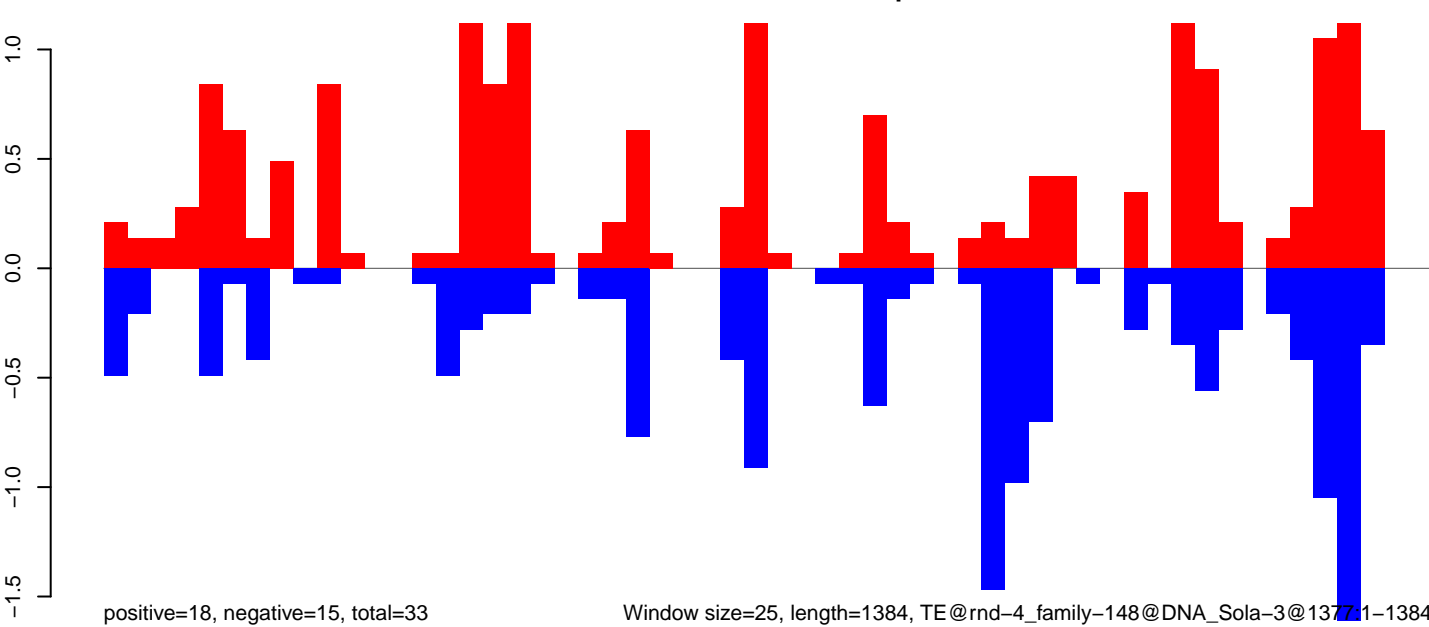
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CuQuin_Mcarc_NL.24_35.rep



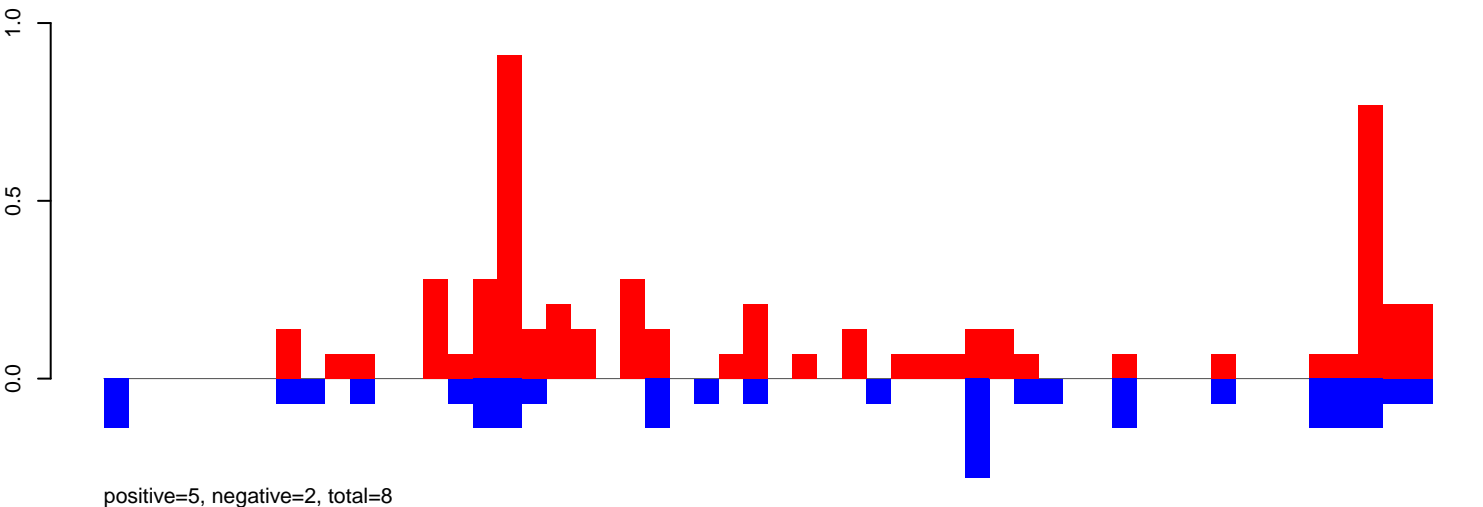
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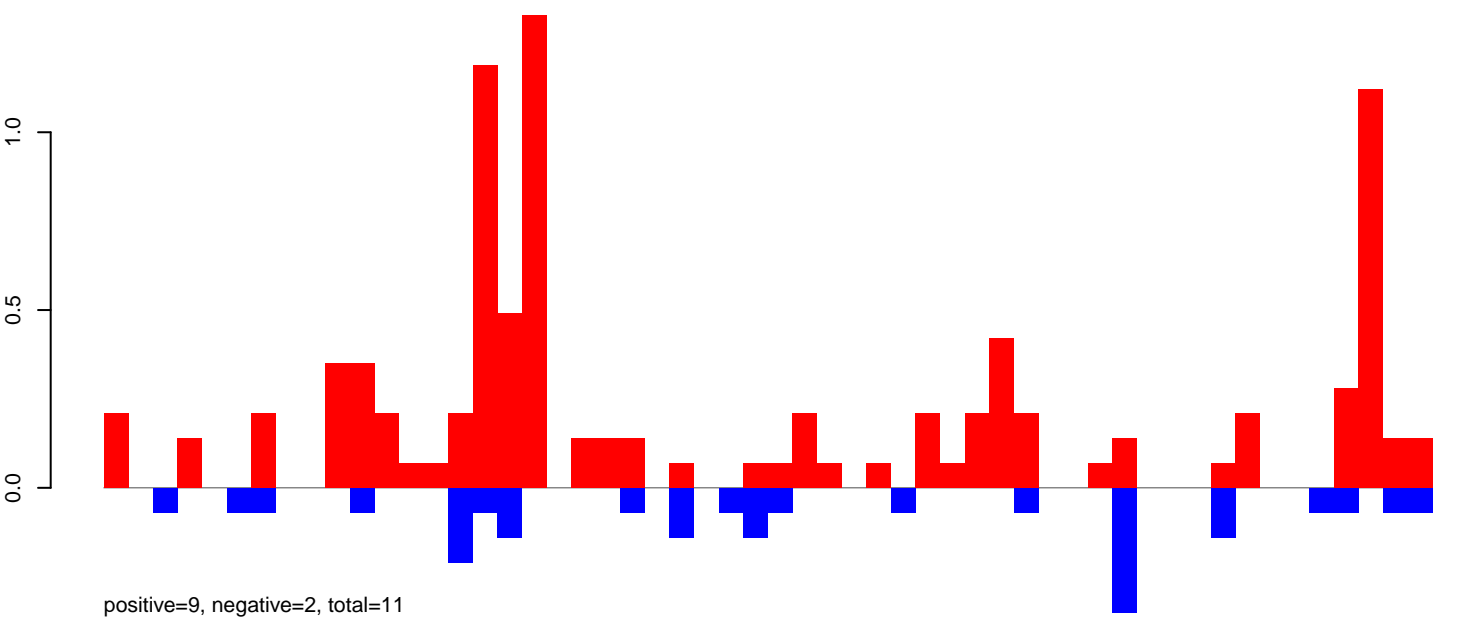
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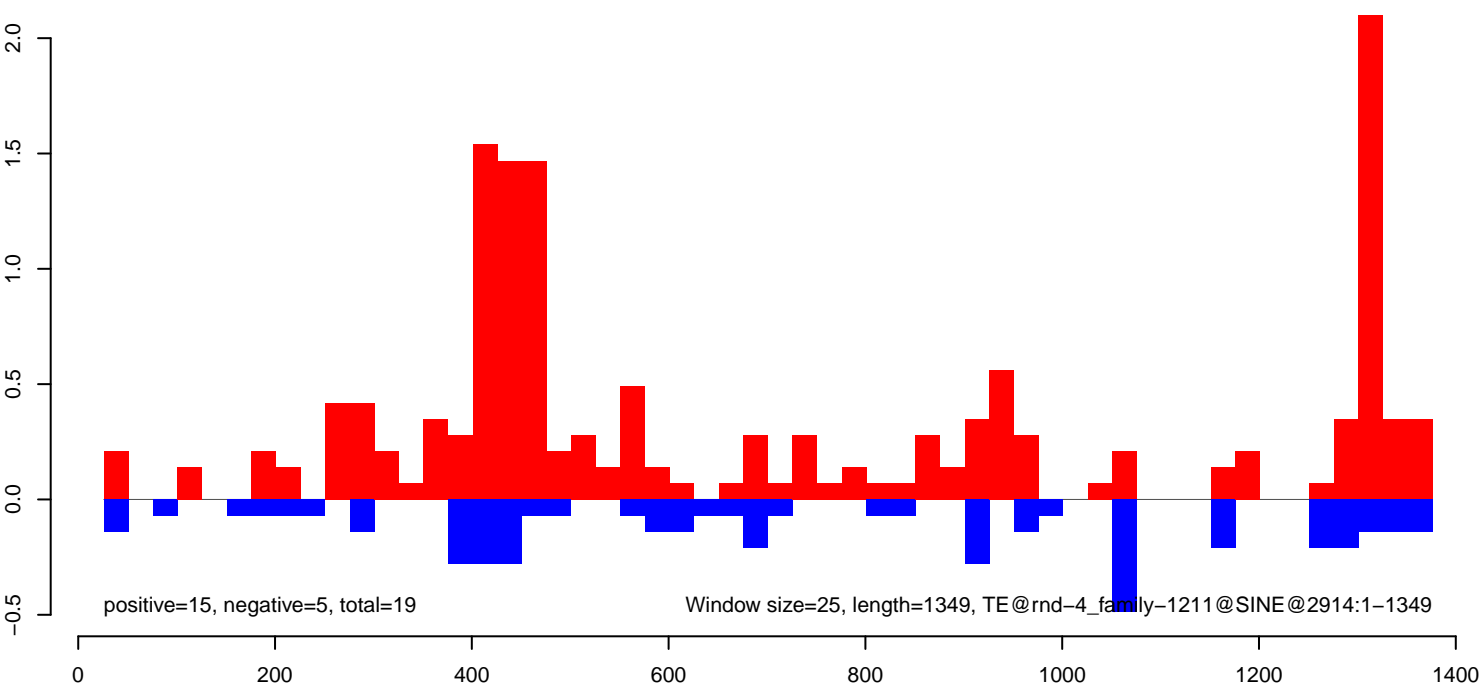
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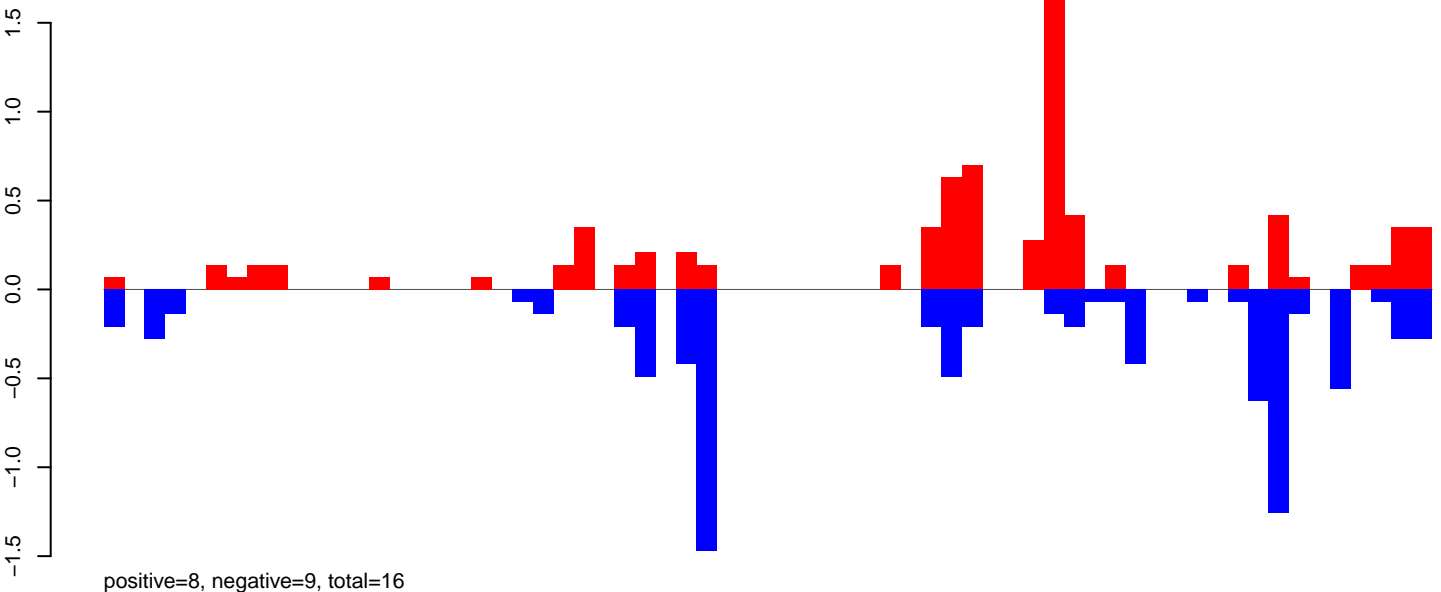
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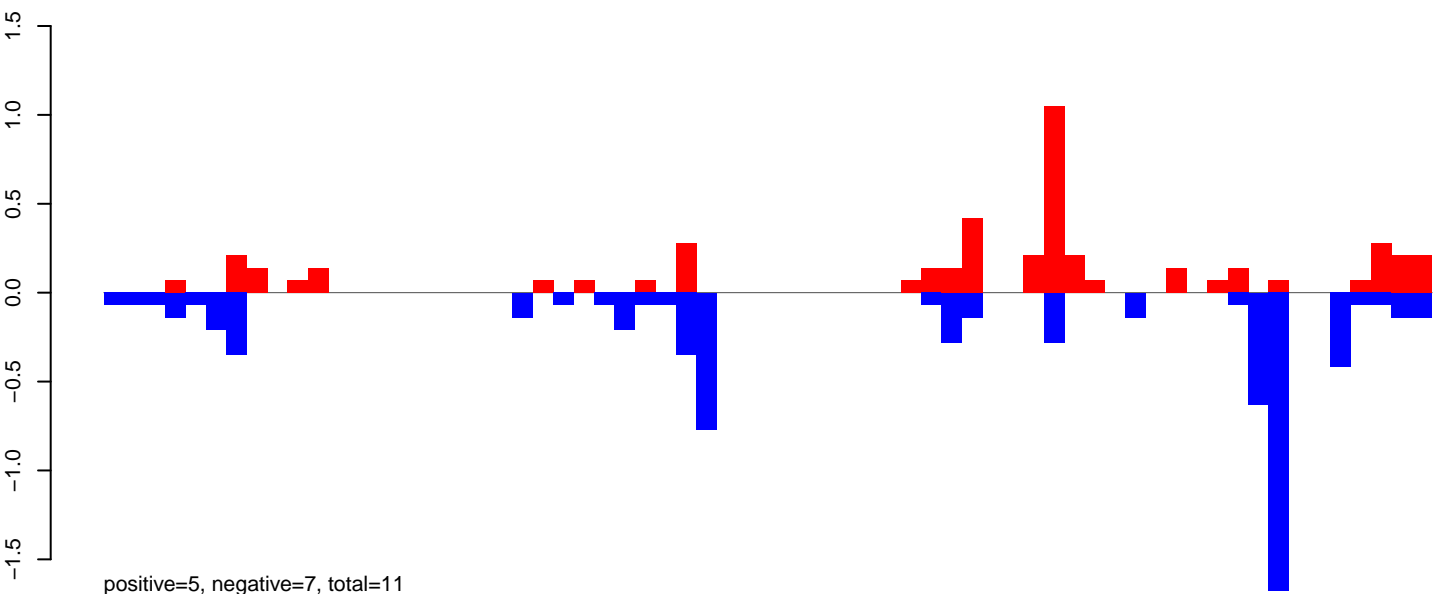
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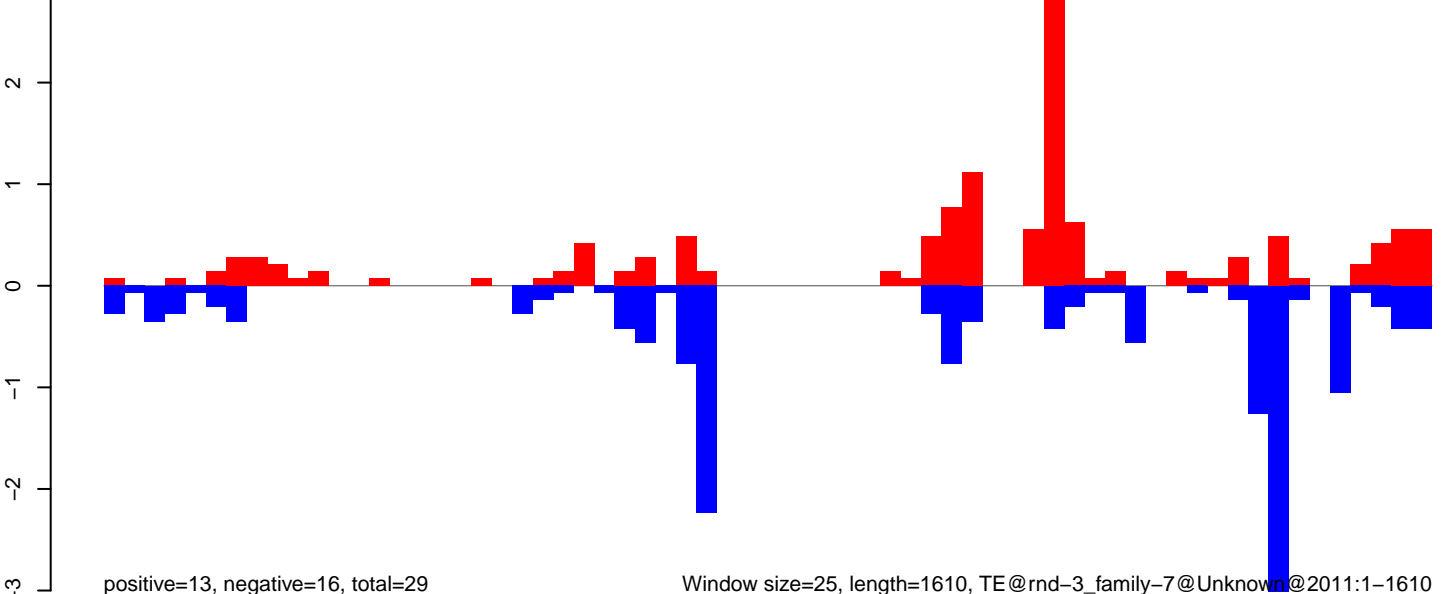
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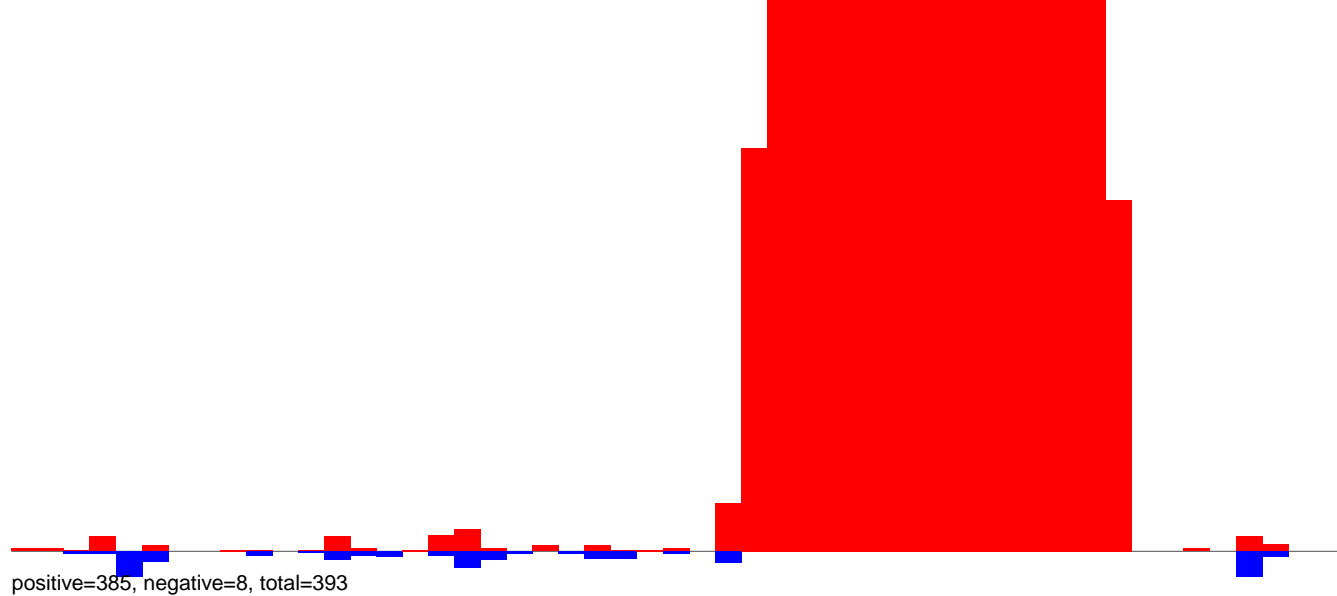
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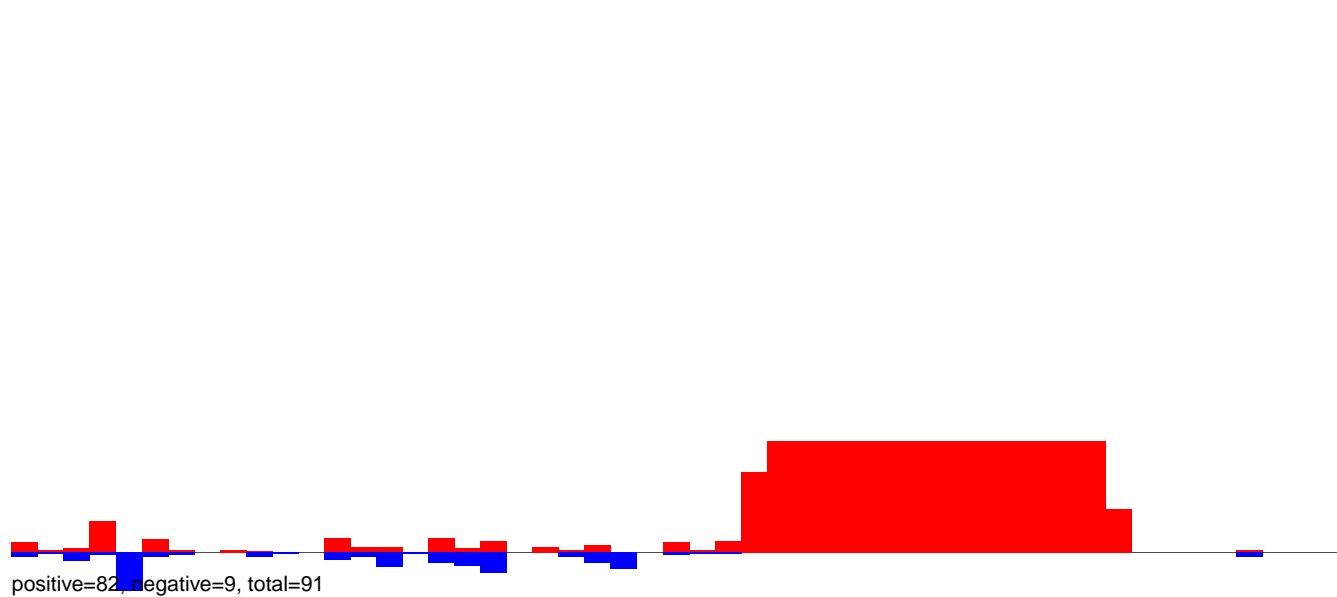
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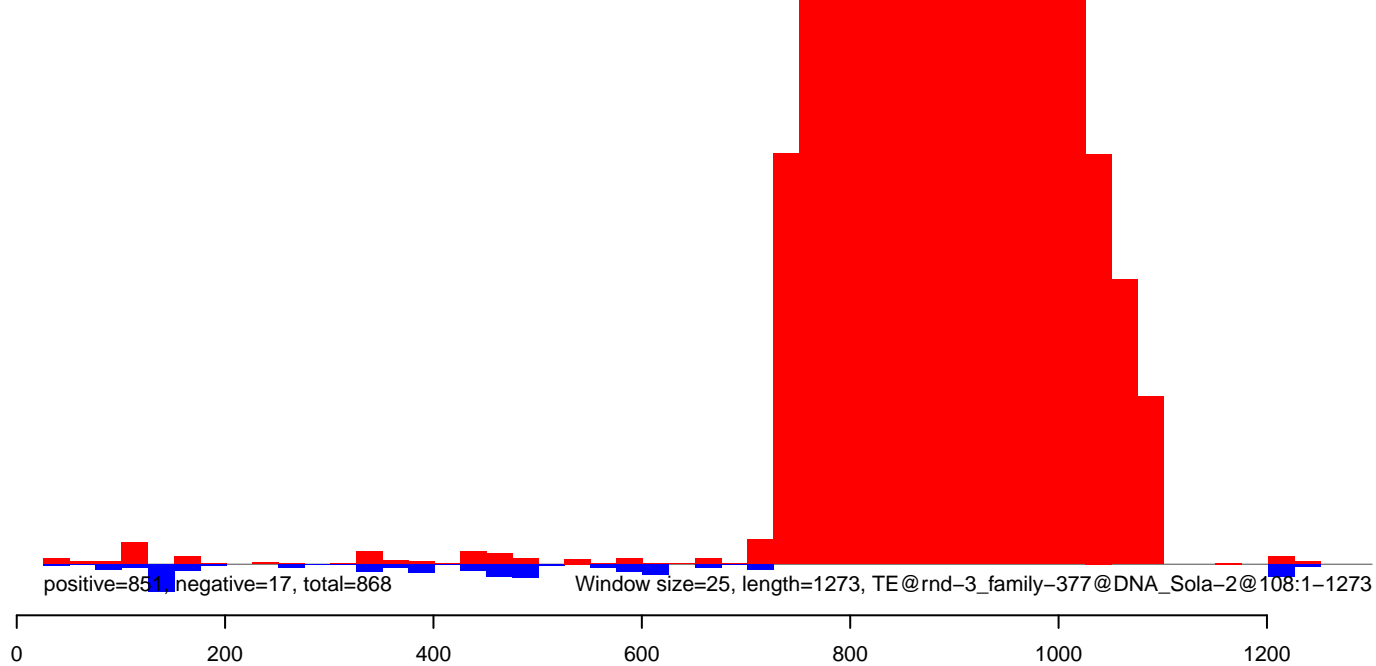
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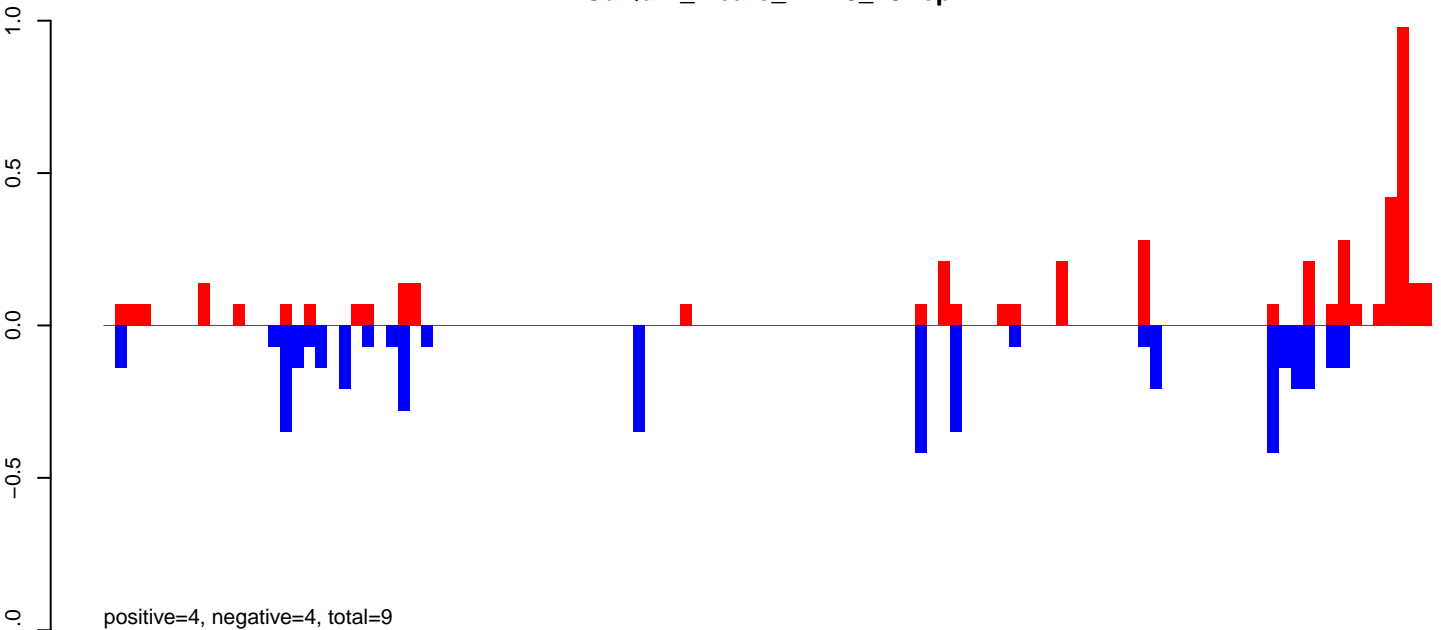
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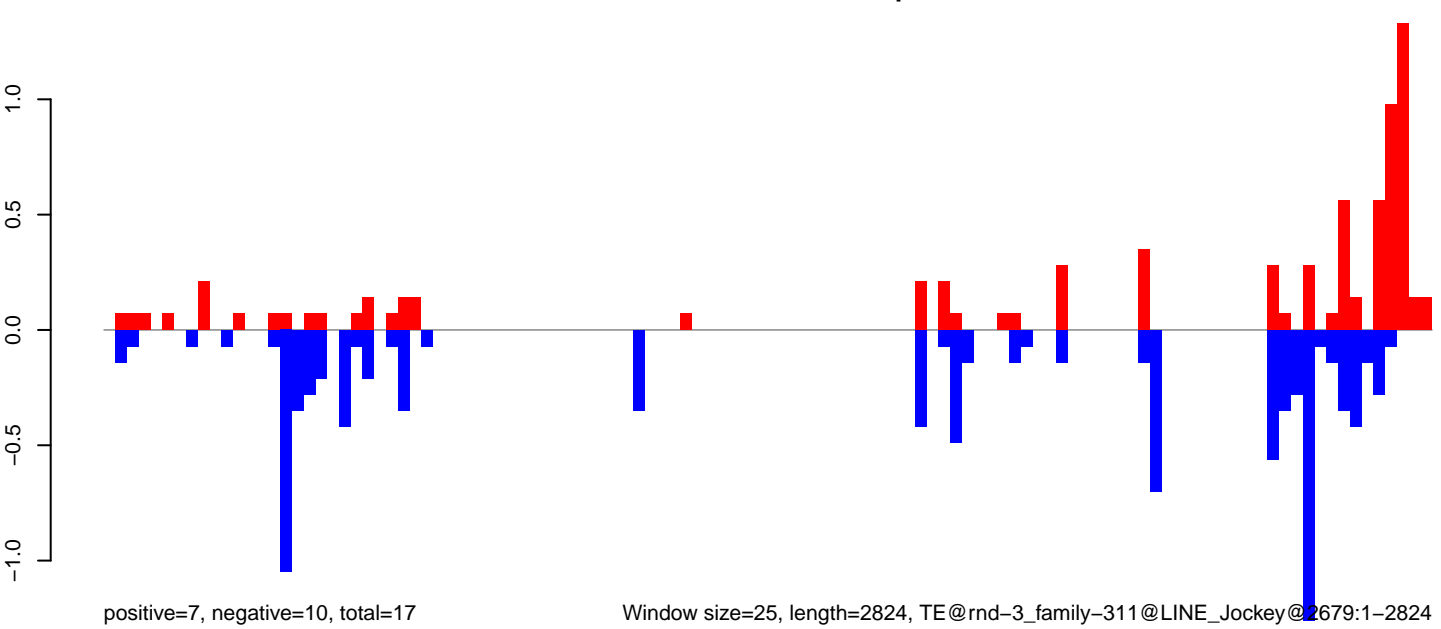
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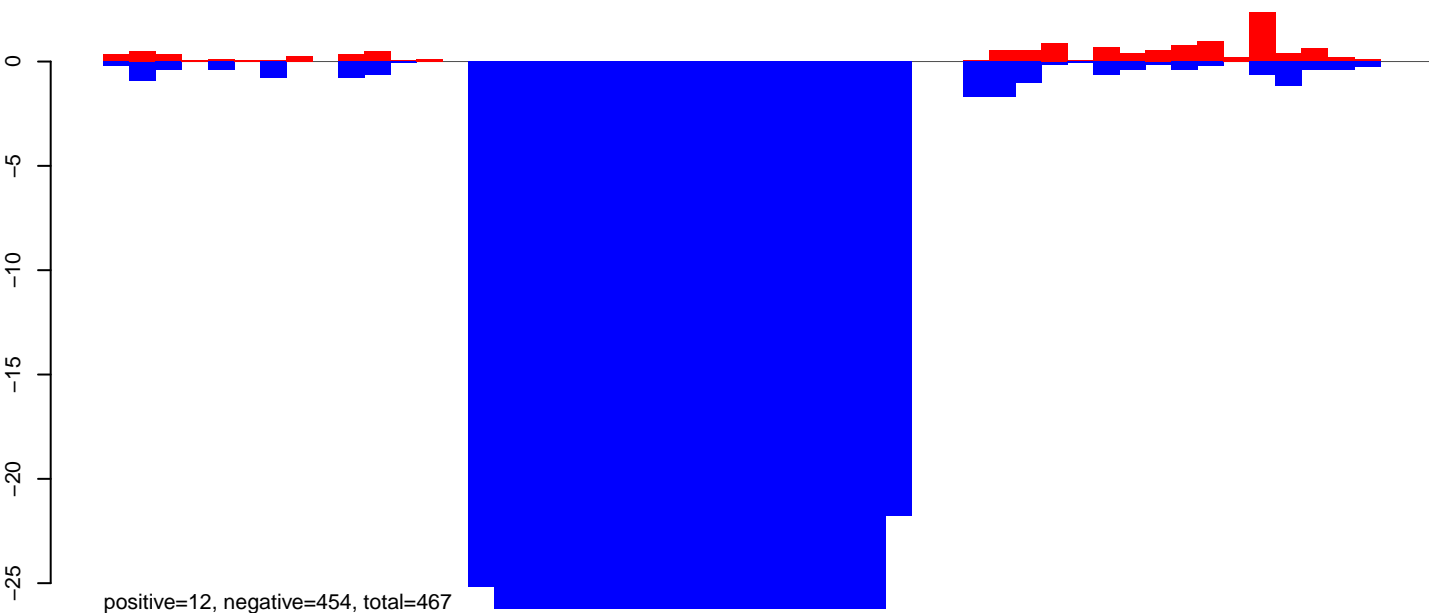


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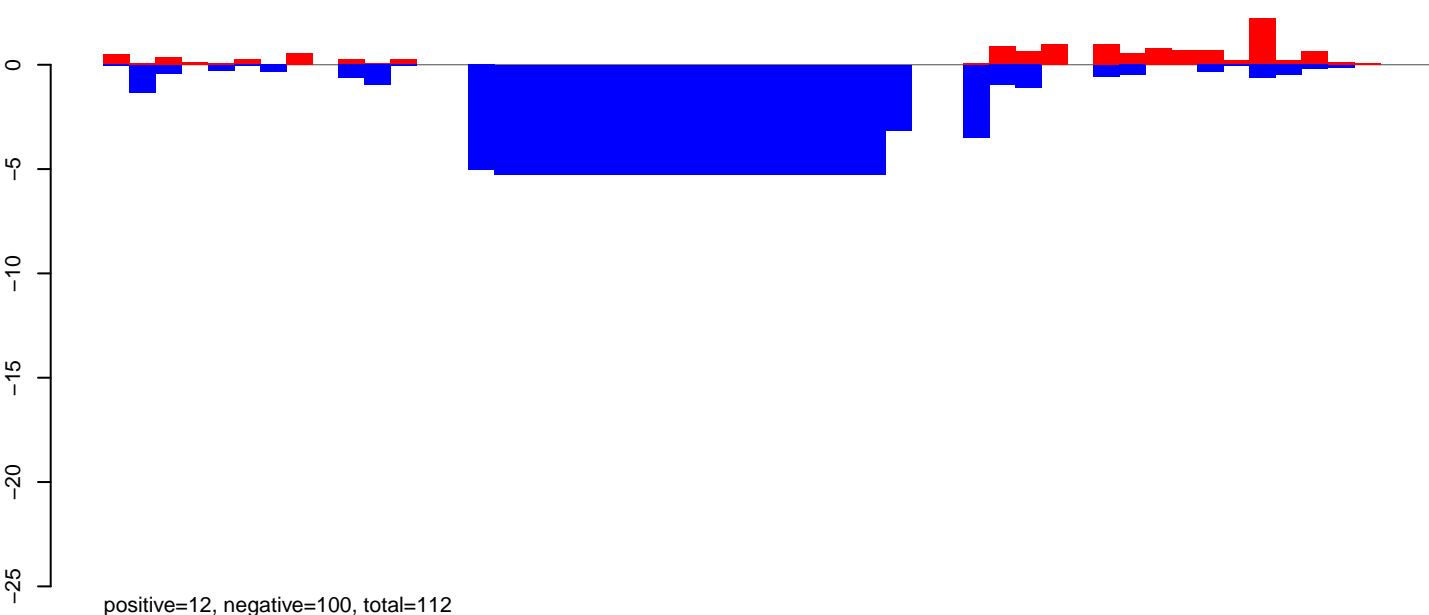


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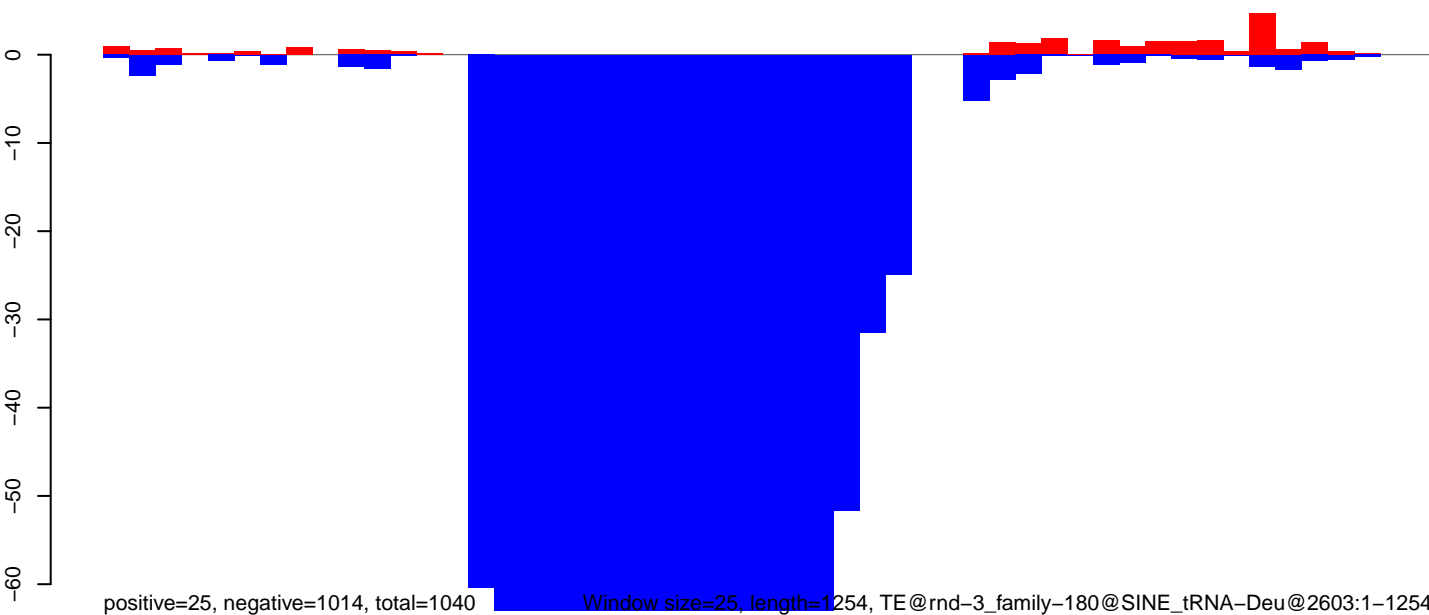
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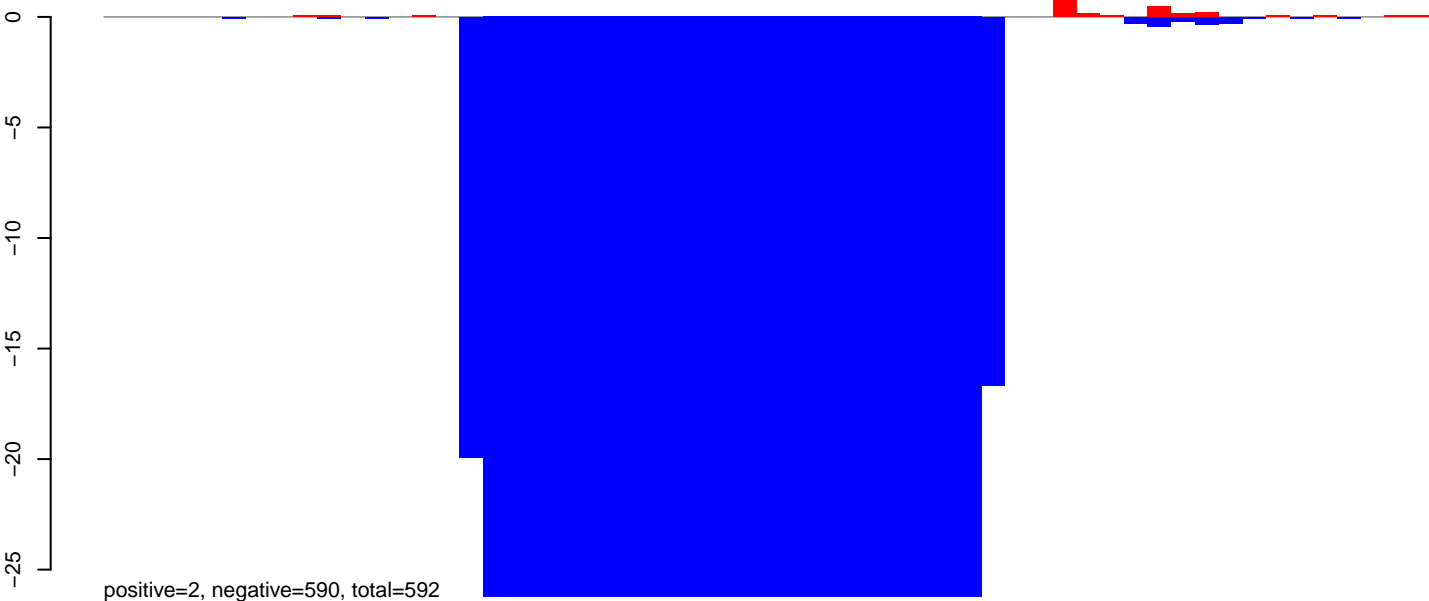


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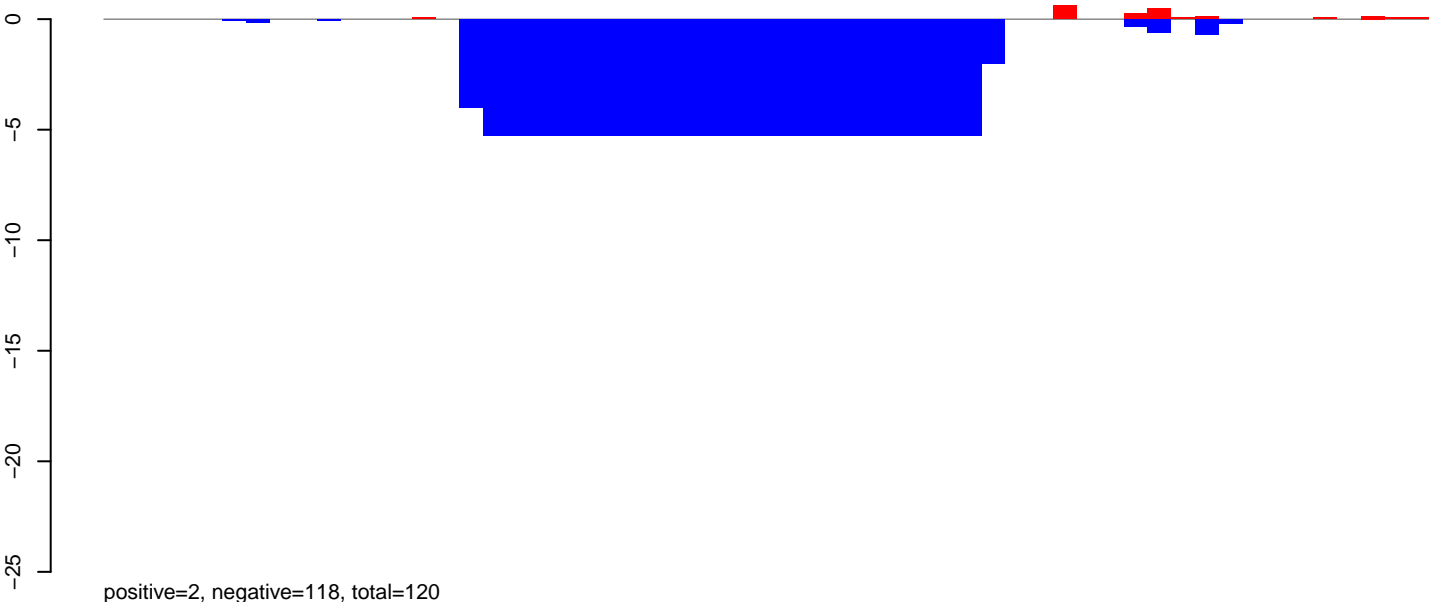


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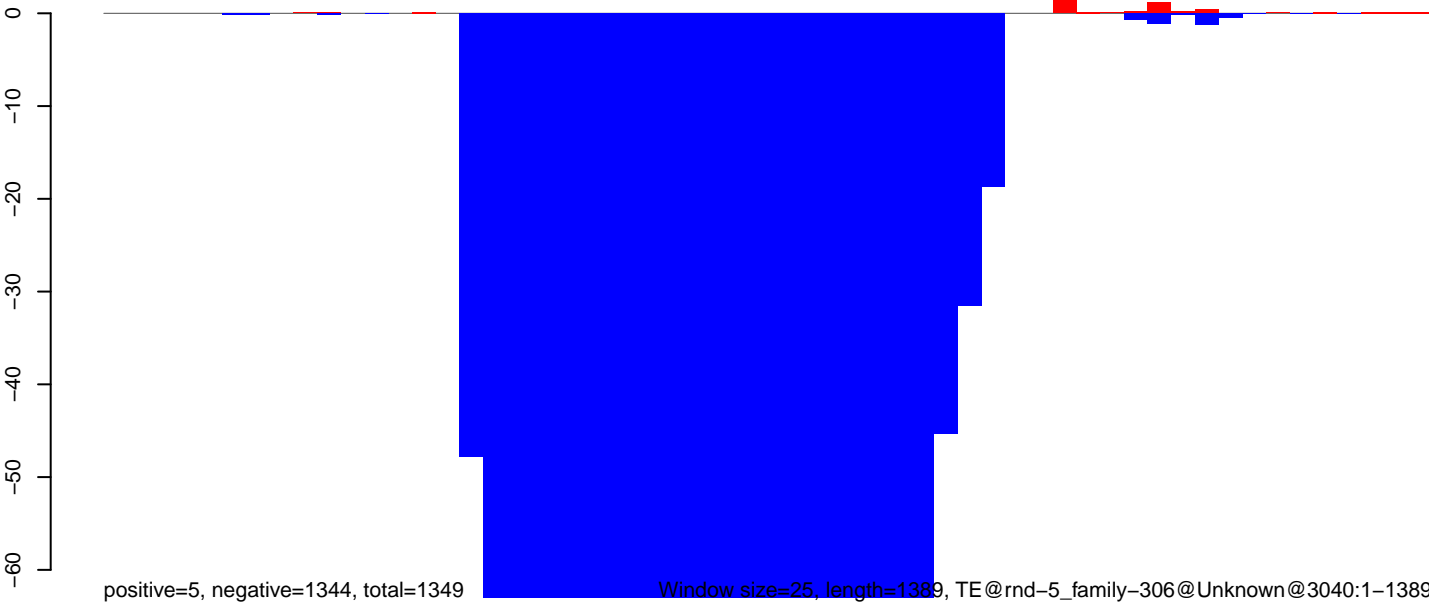
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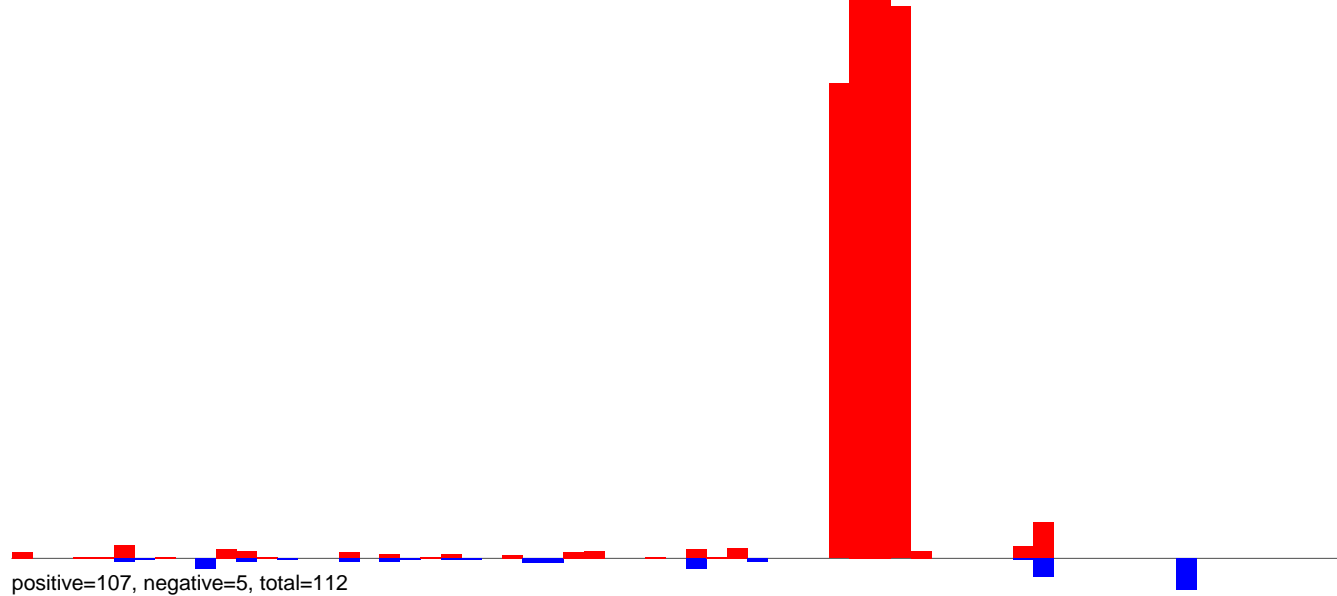


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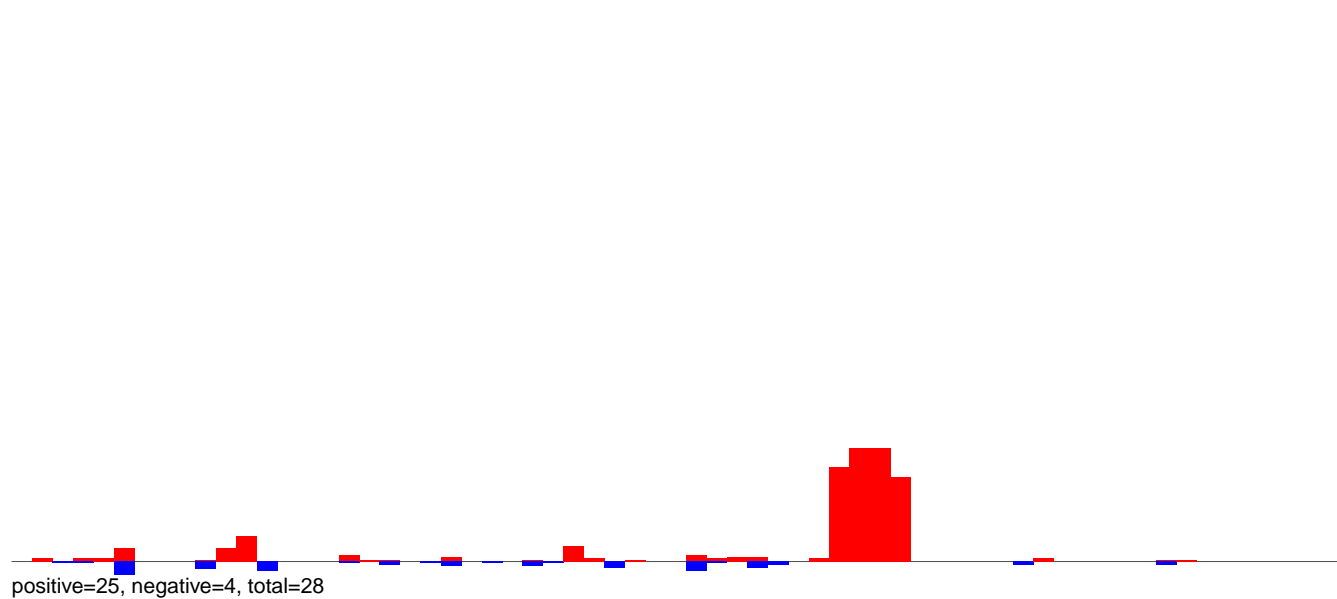


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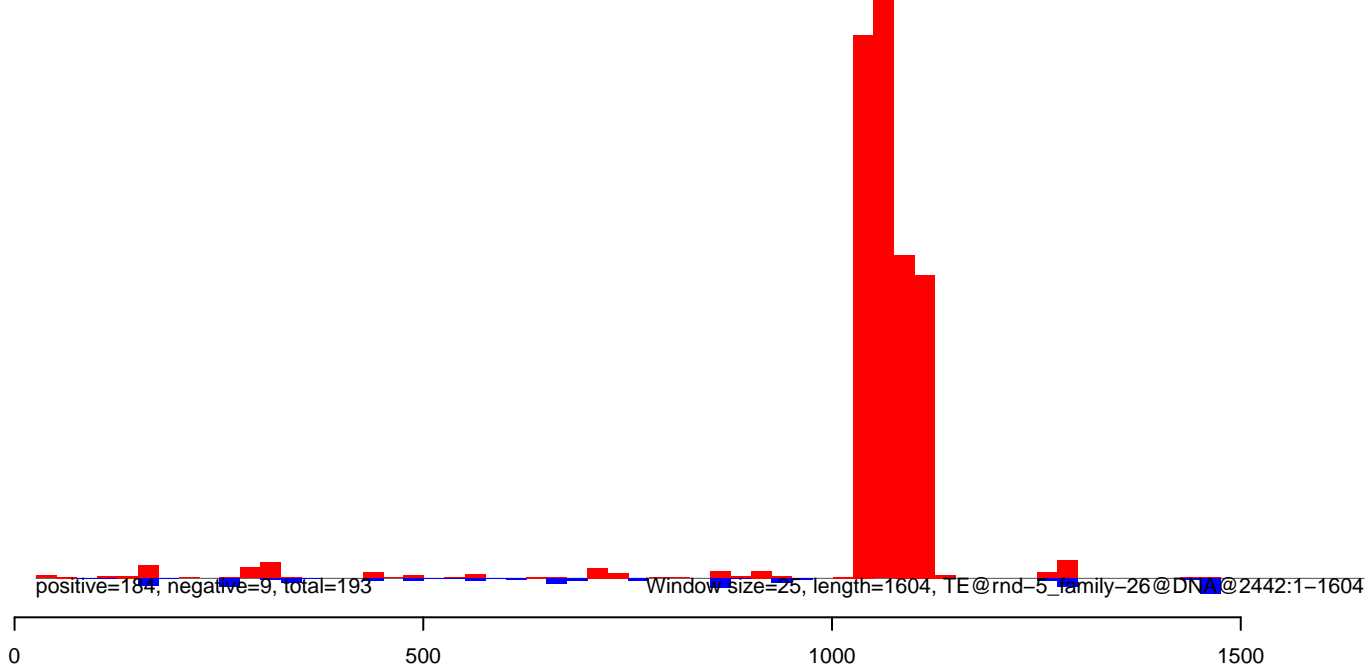
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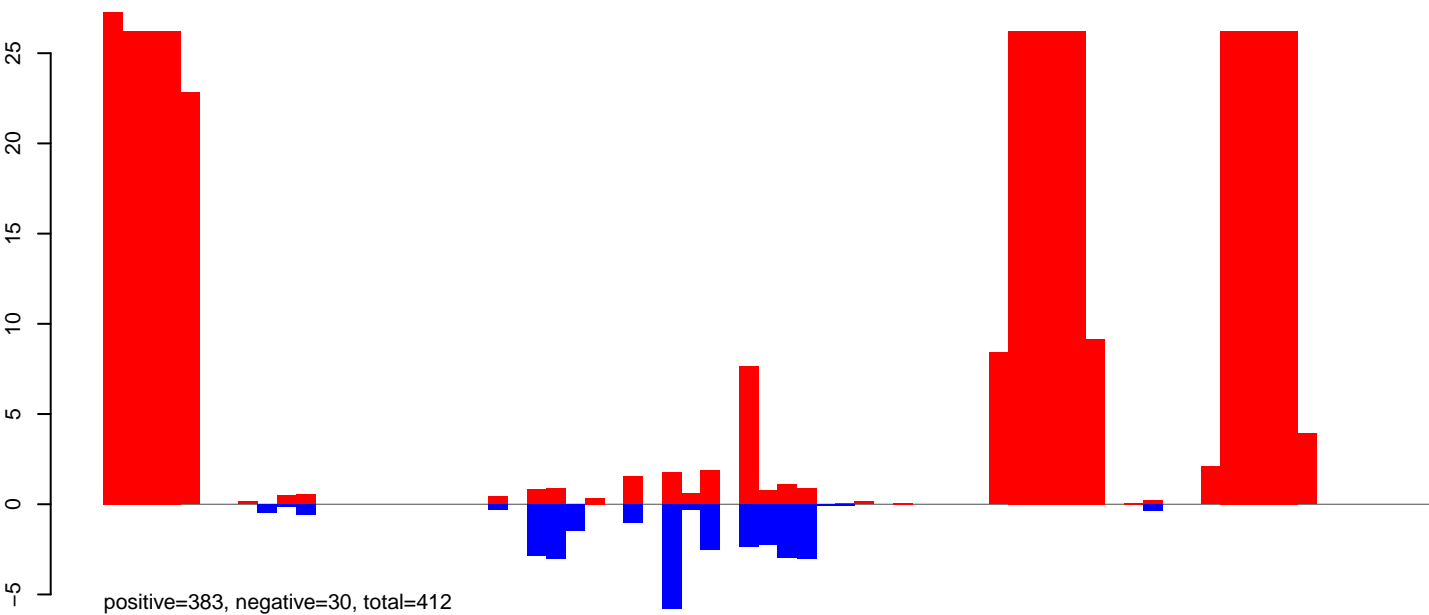
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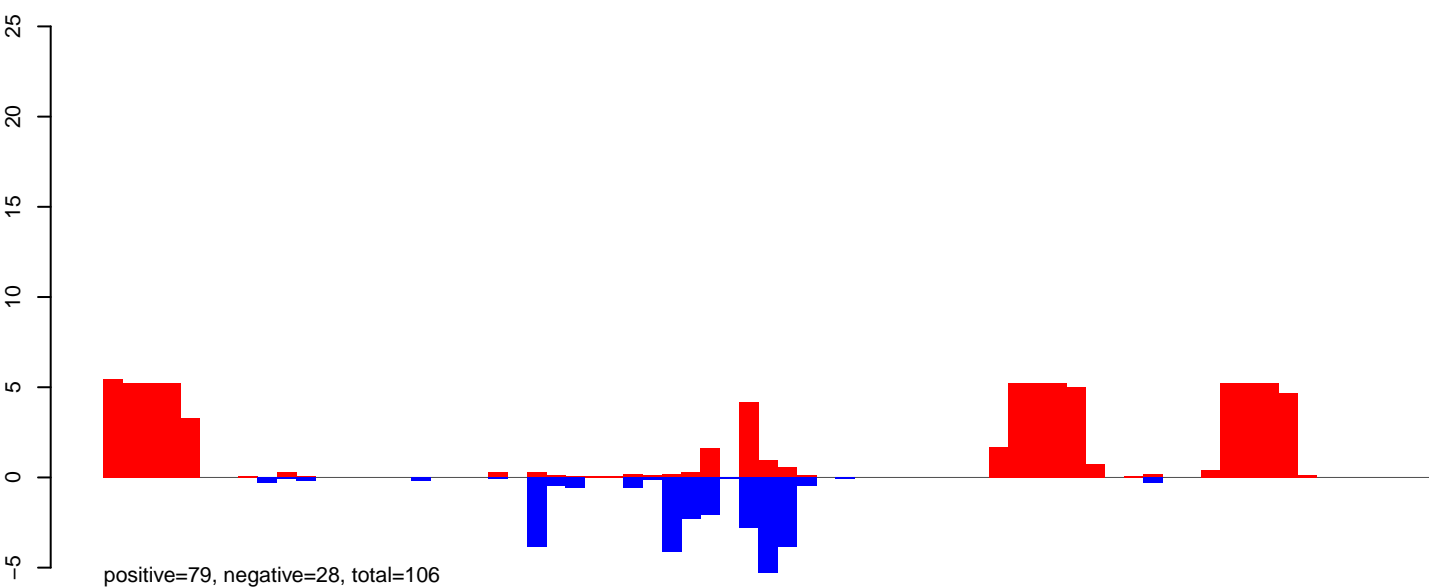
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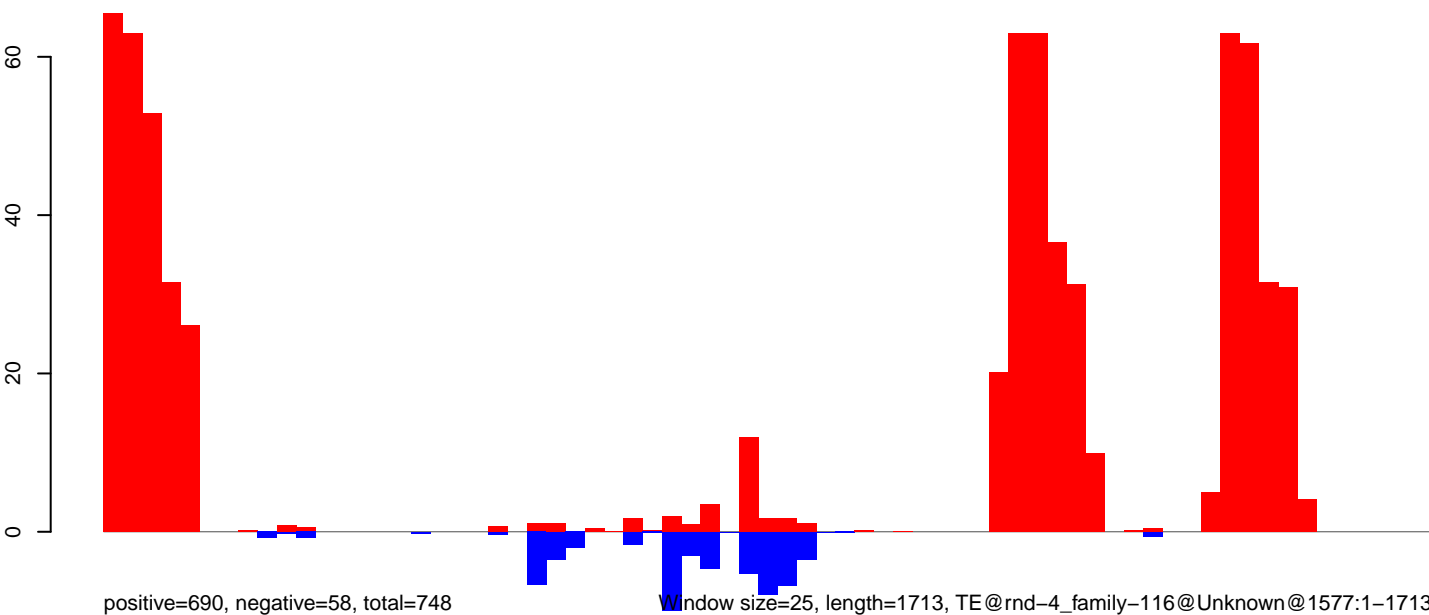
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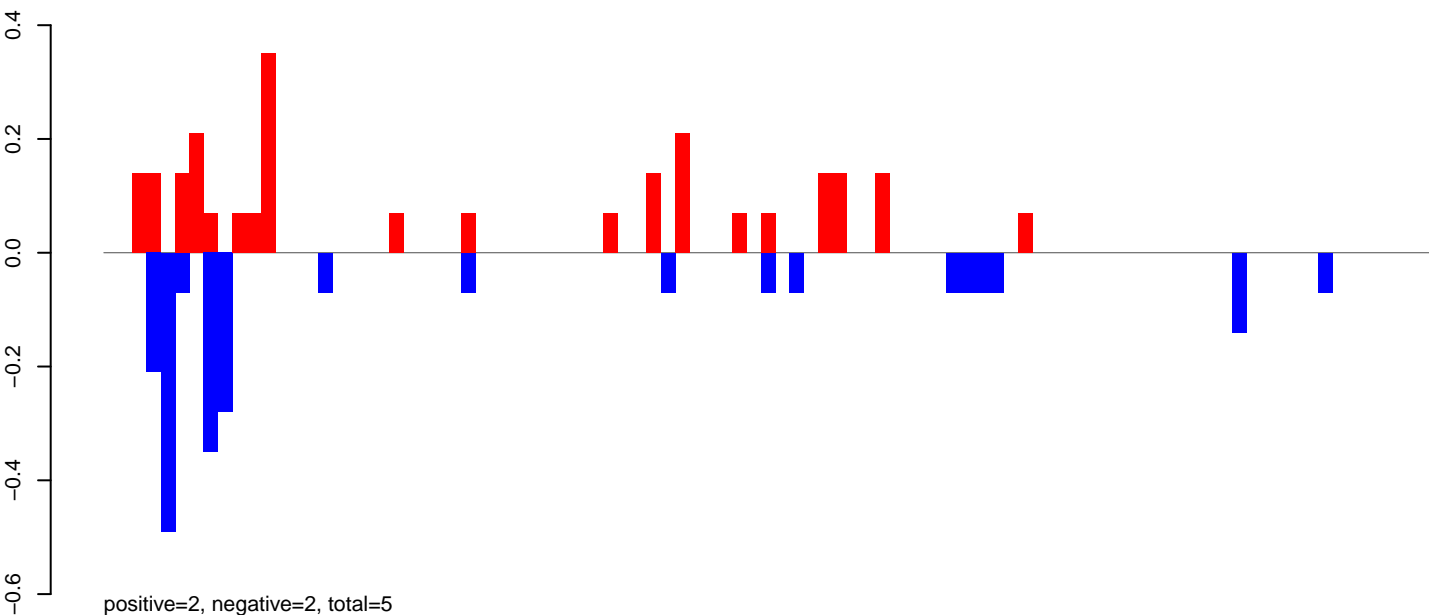


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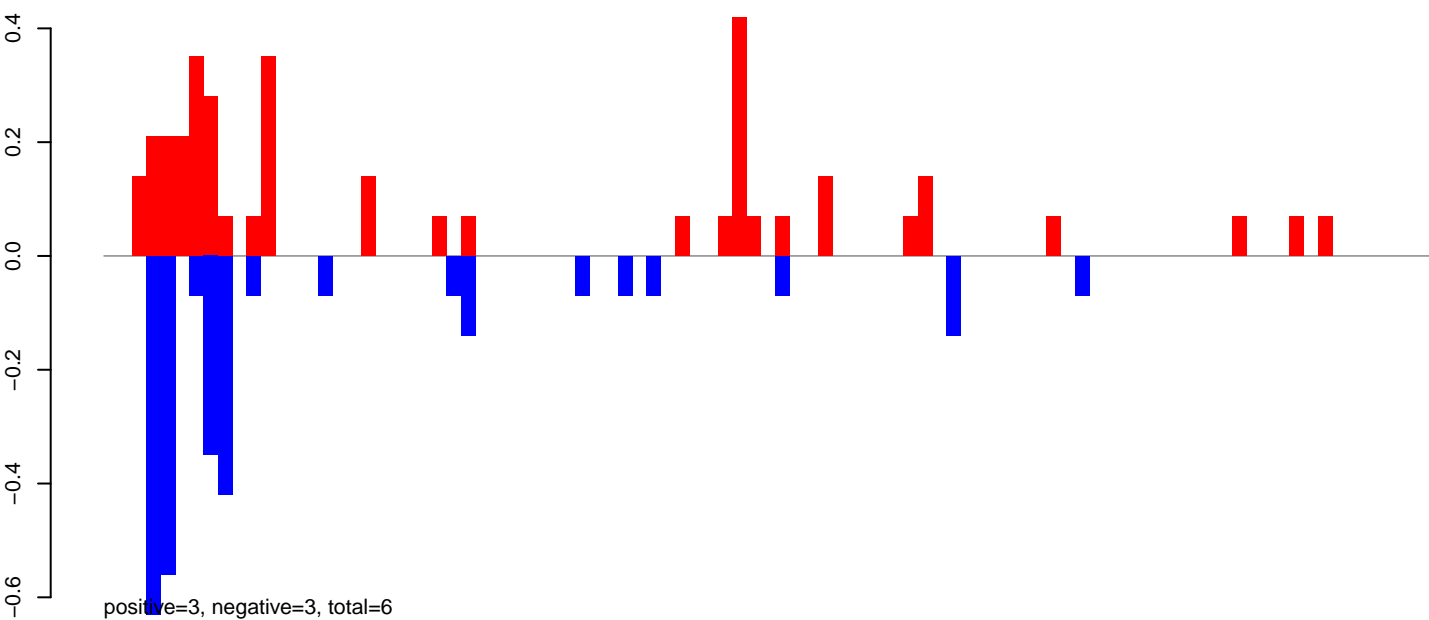


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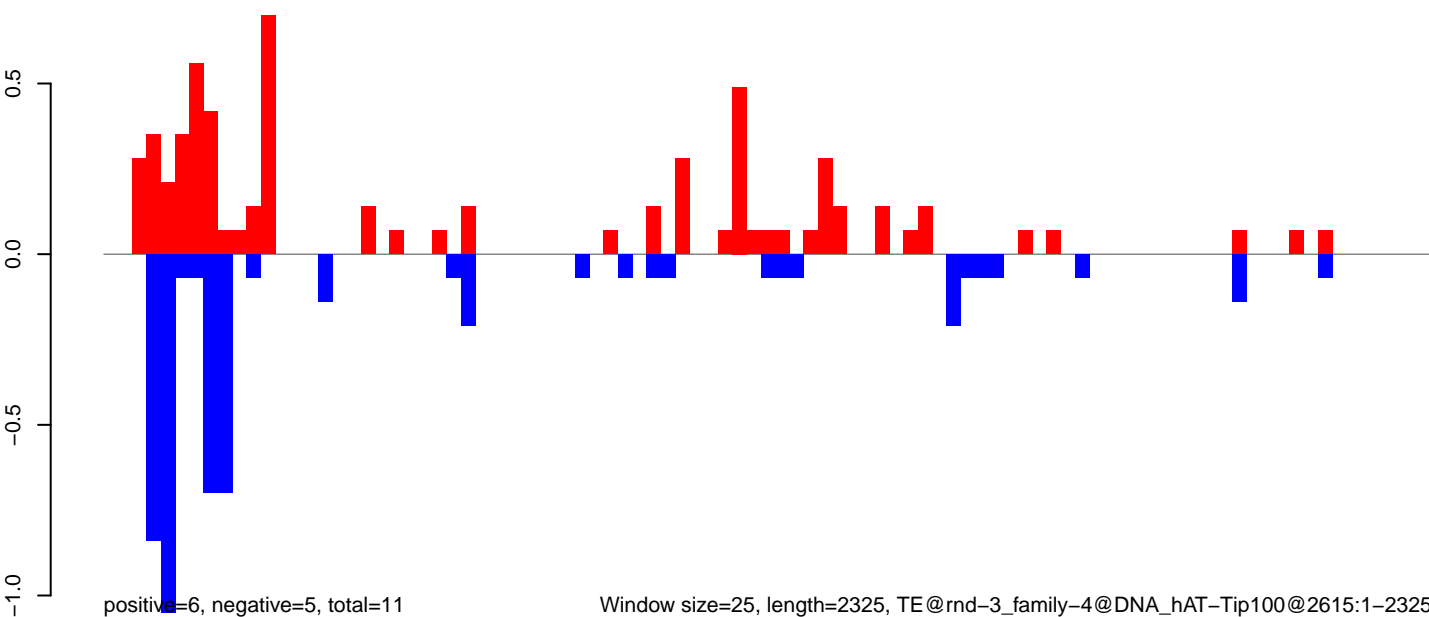
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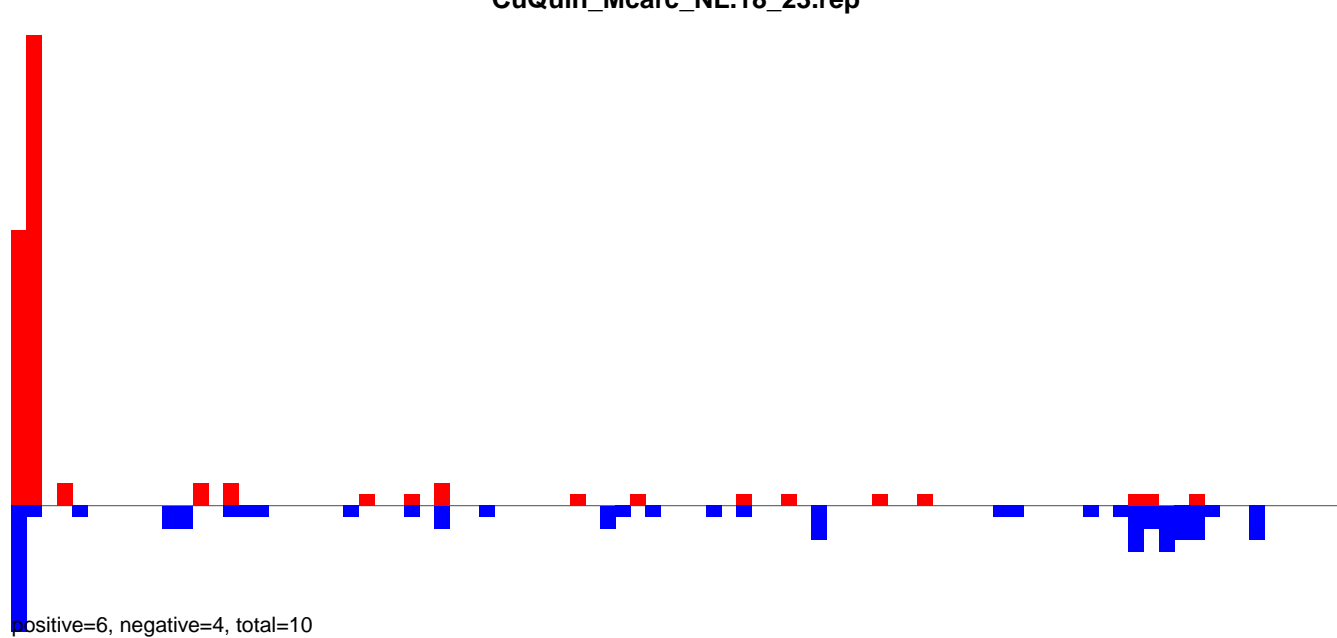
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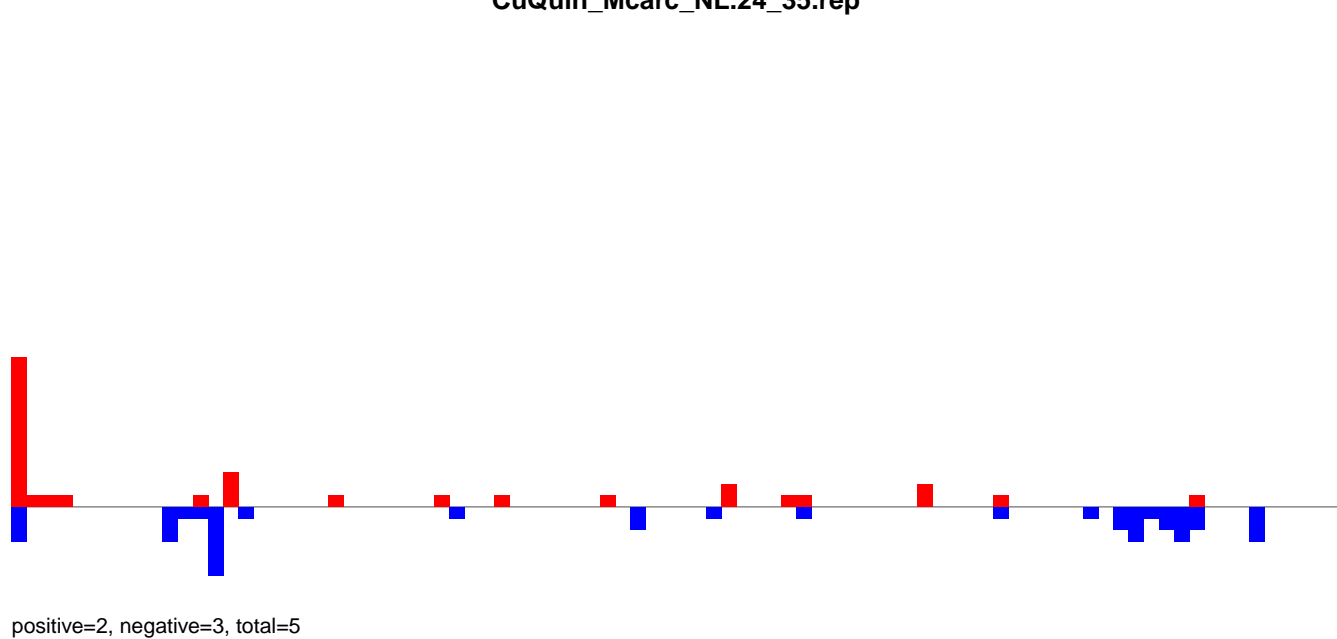
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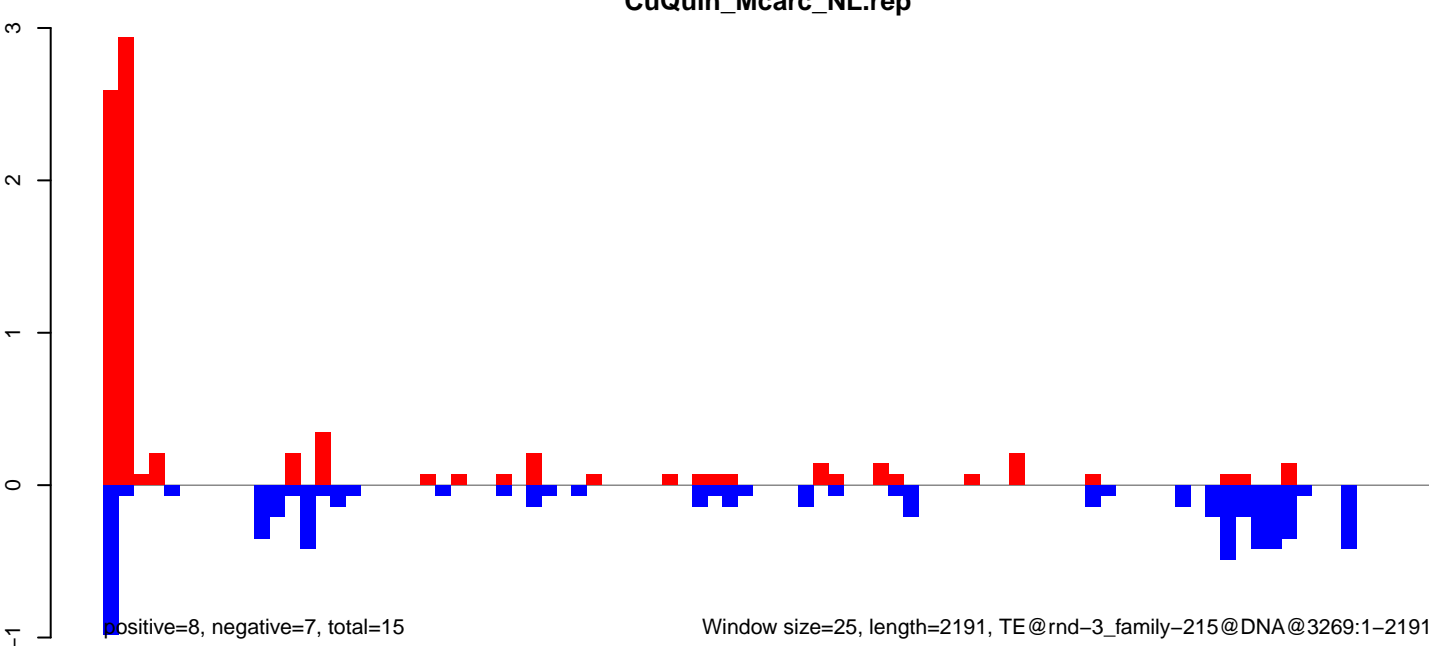
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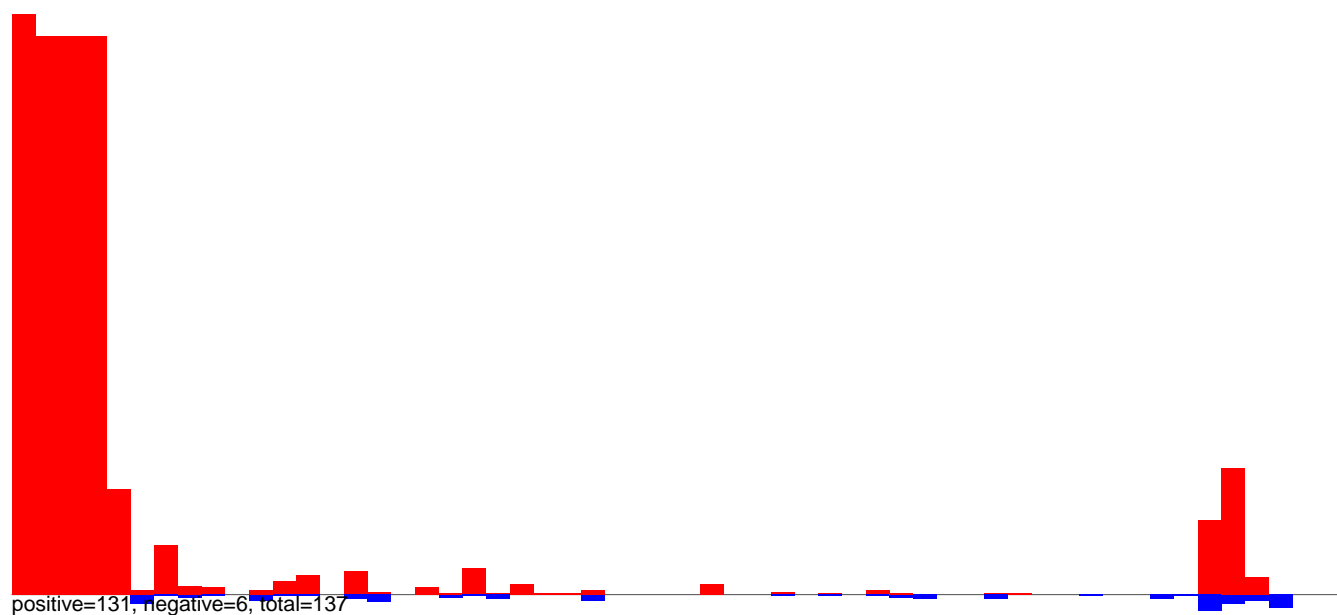
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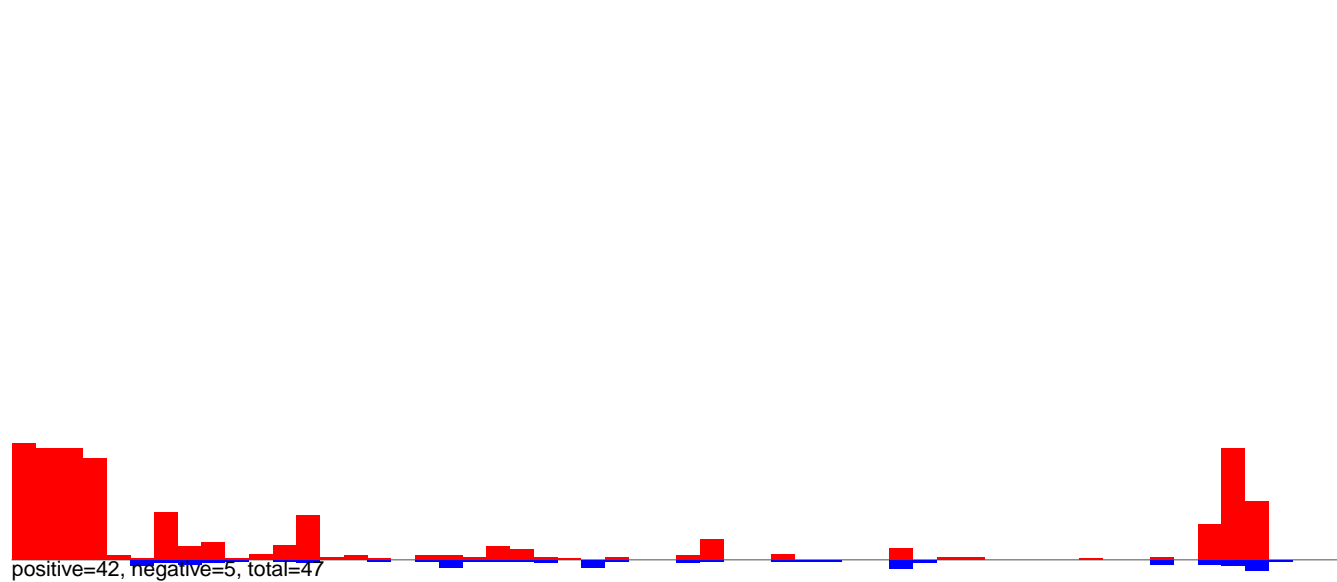
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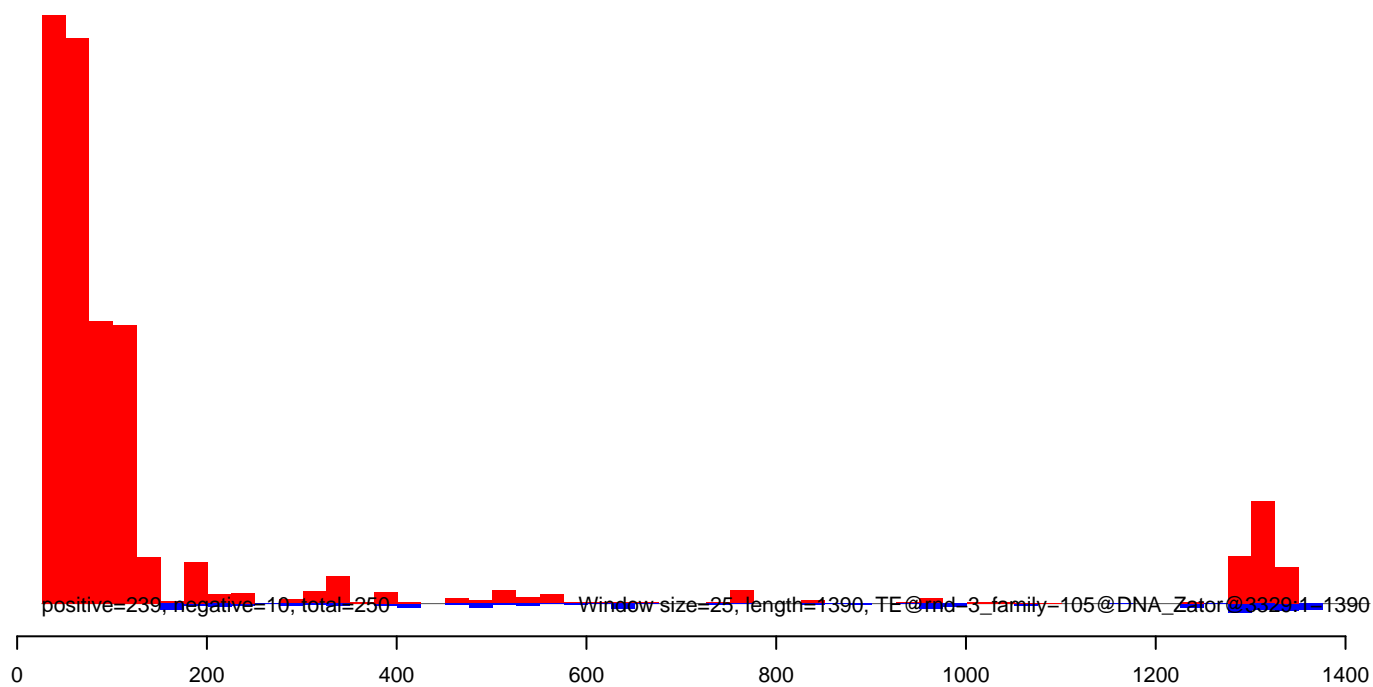
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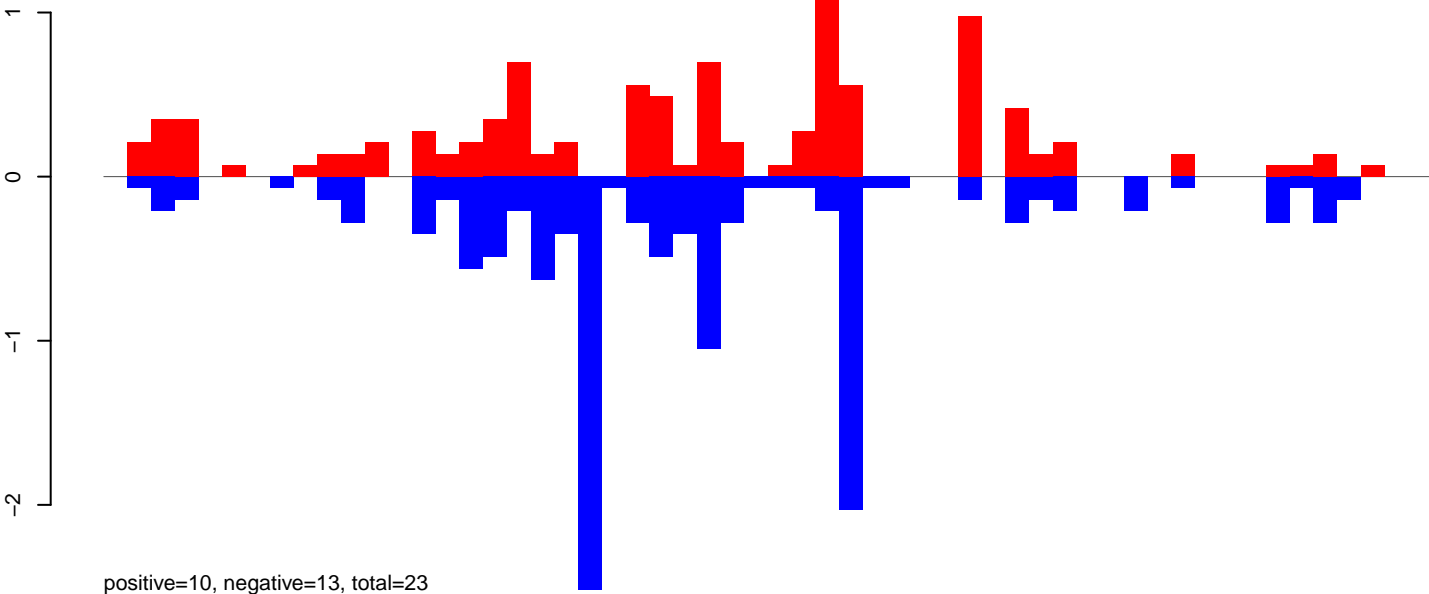
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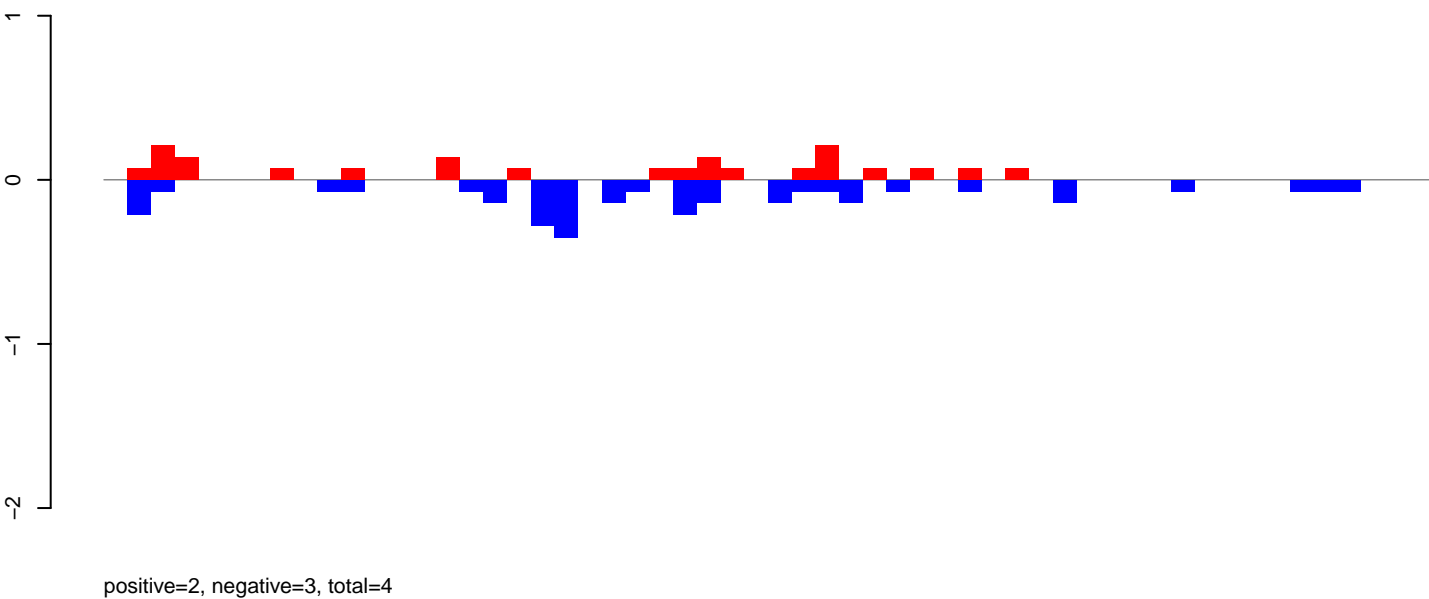
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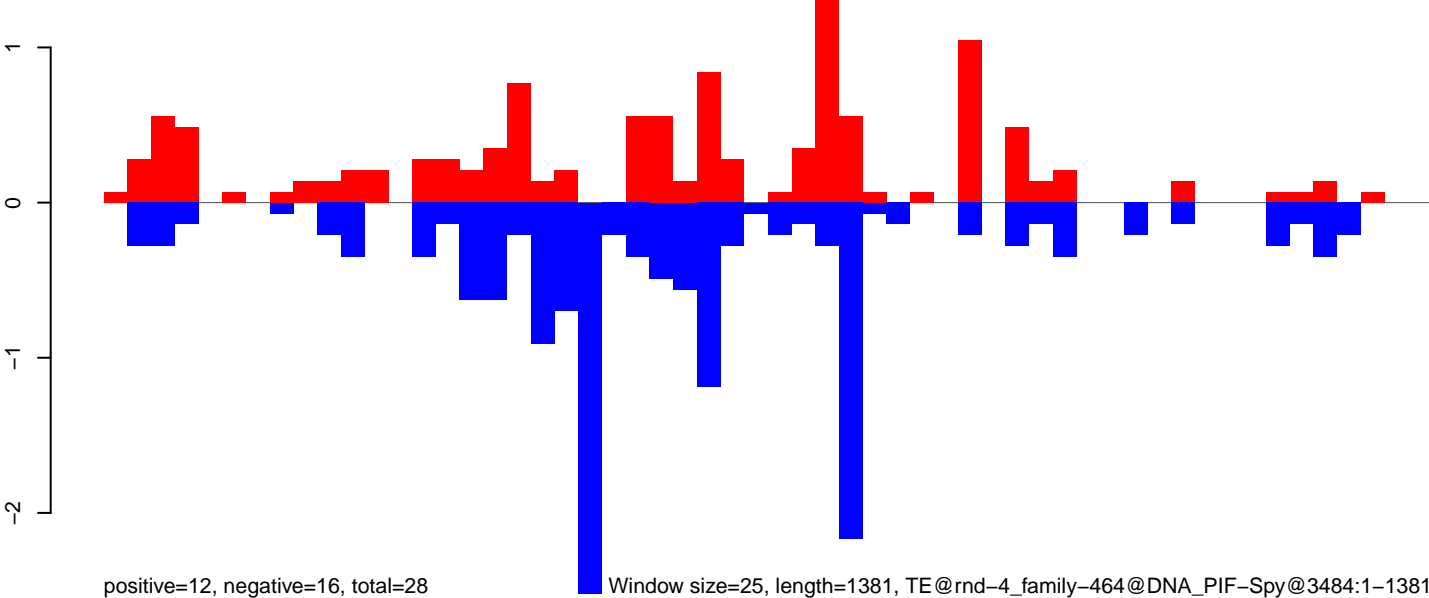
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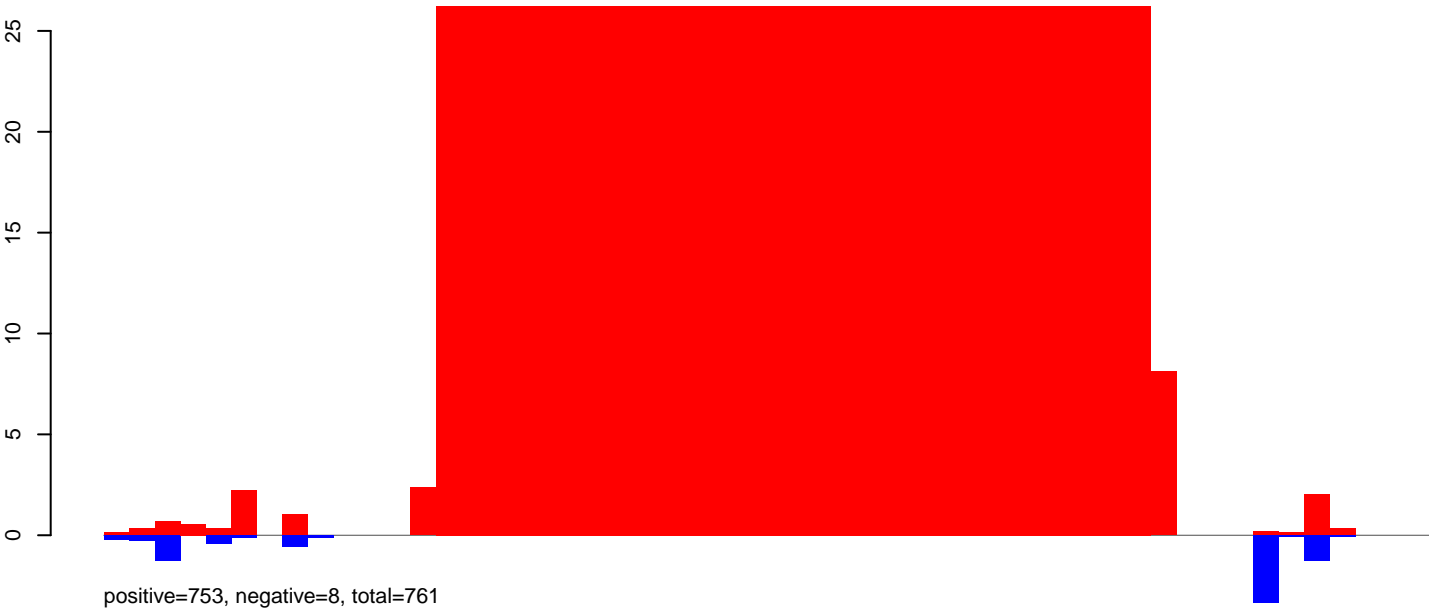
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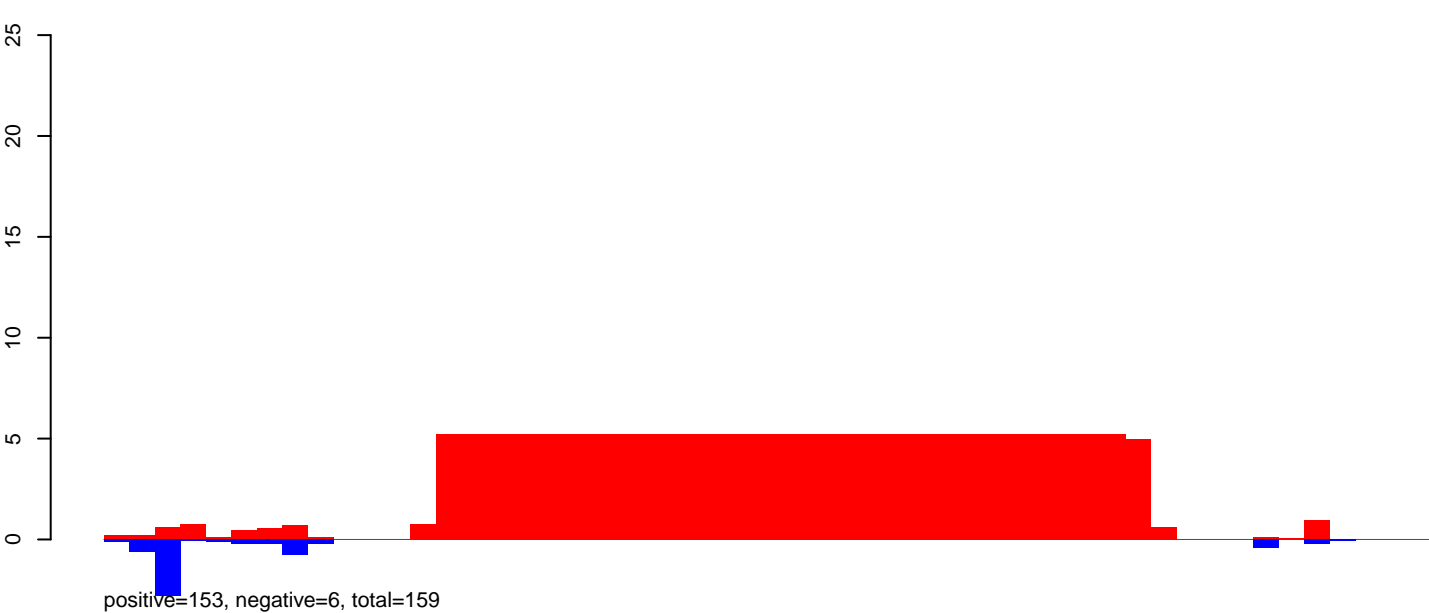
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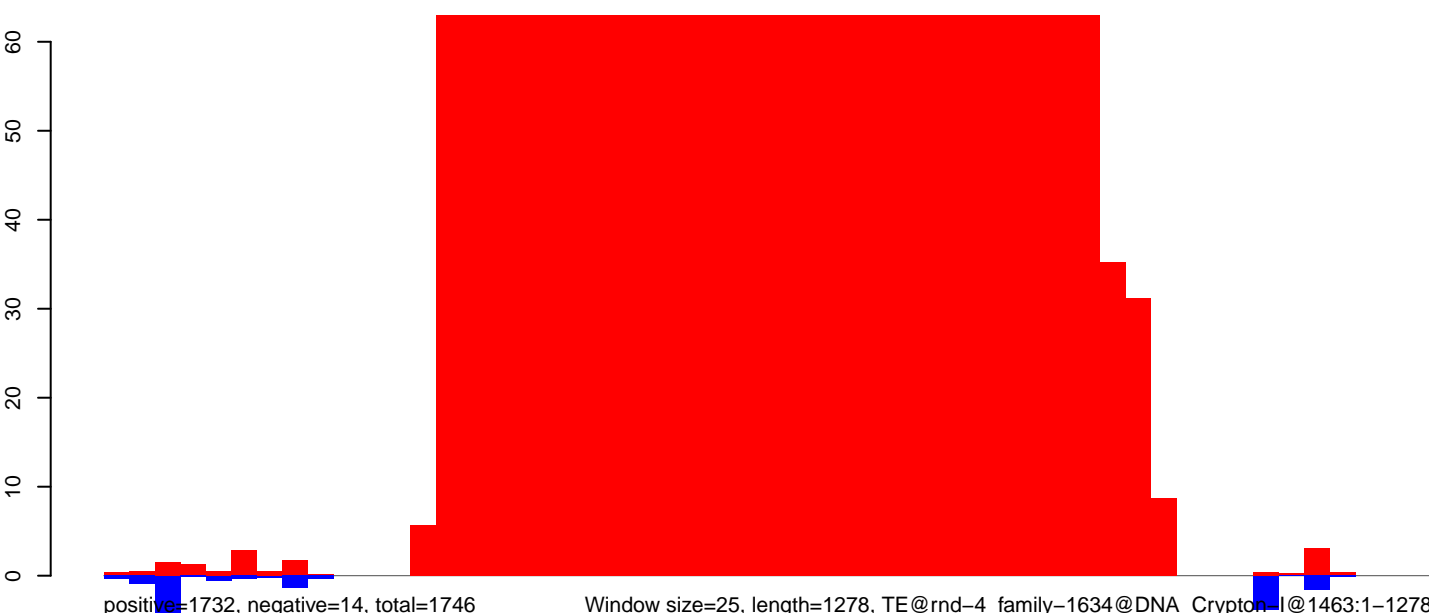
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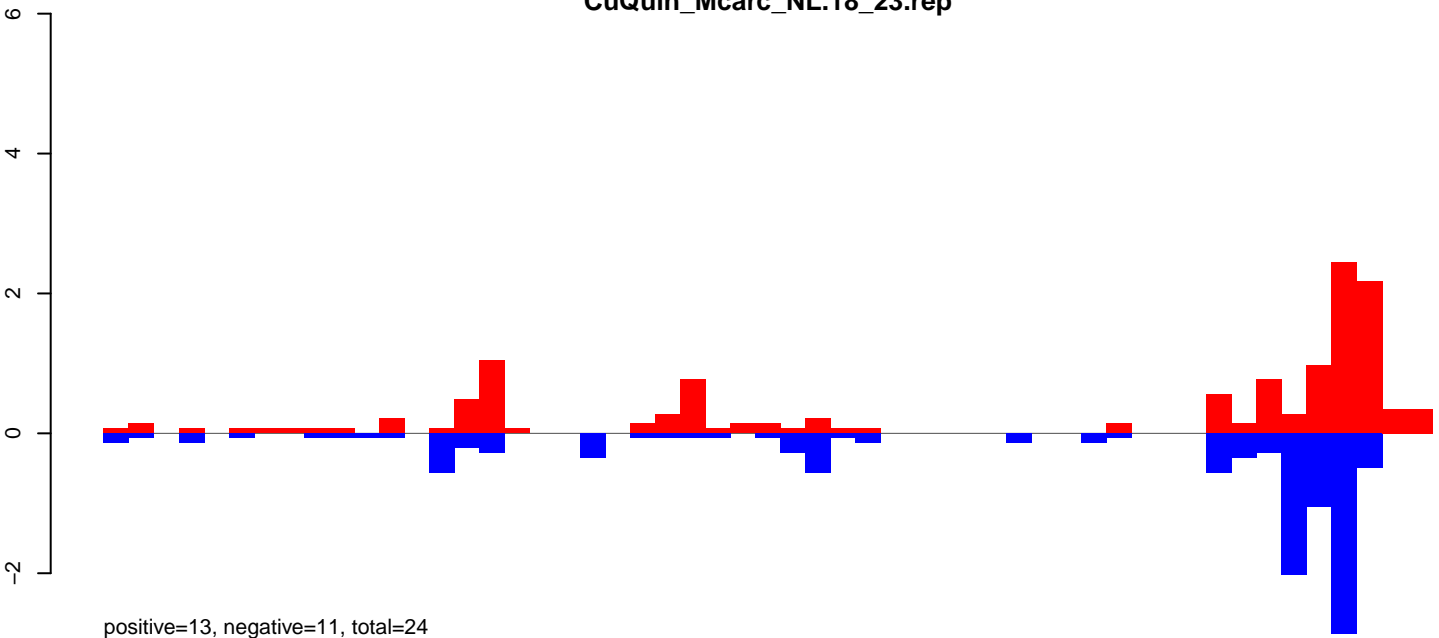
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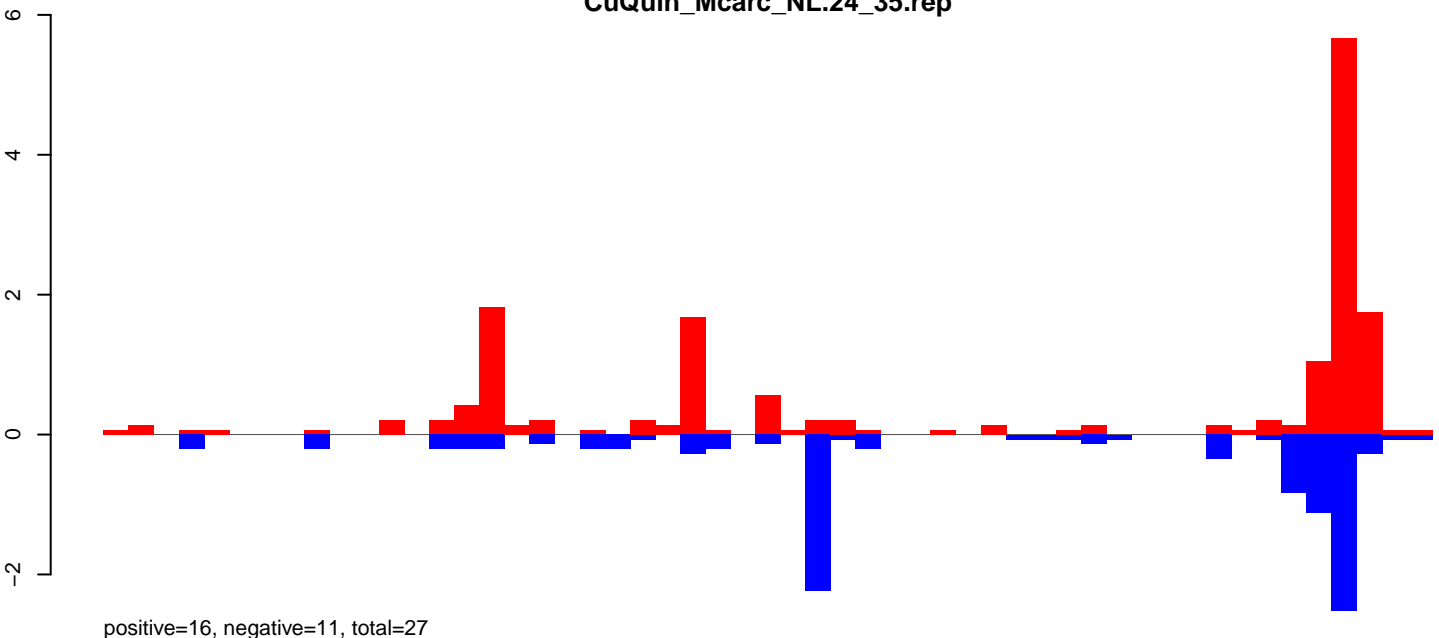
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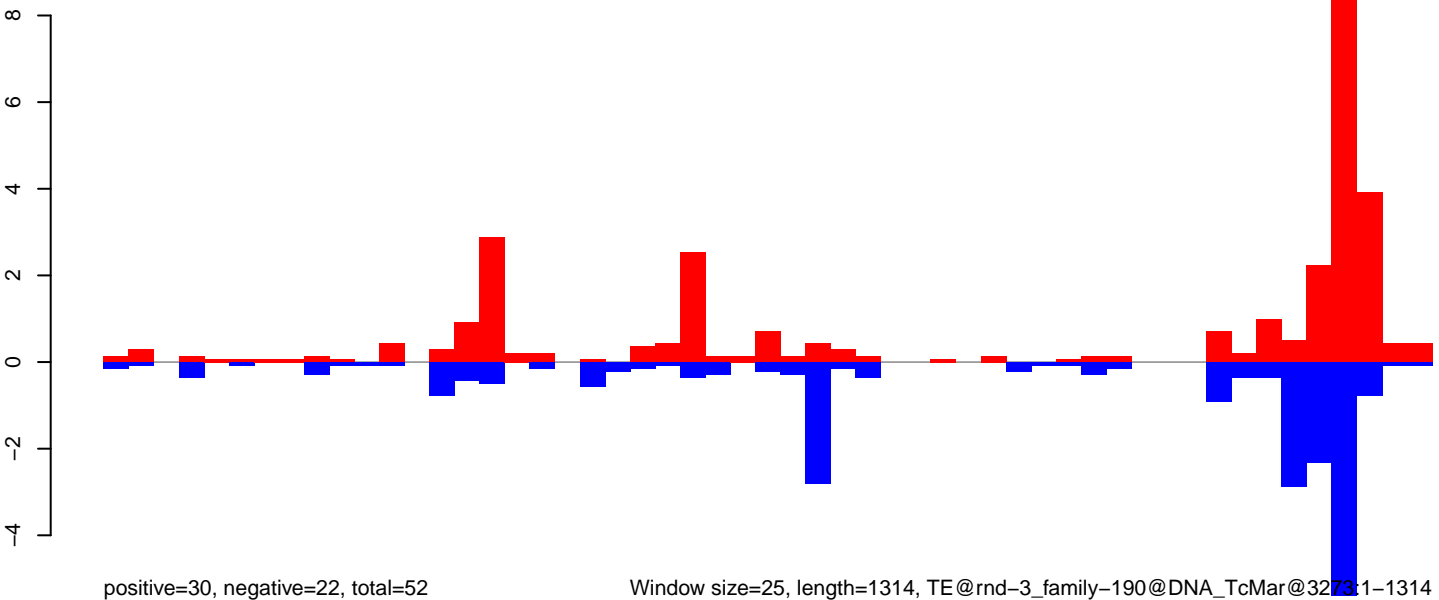
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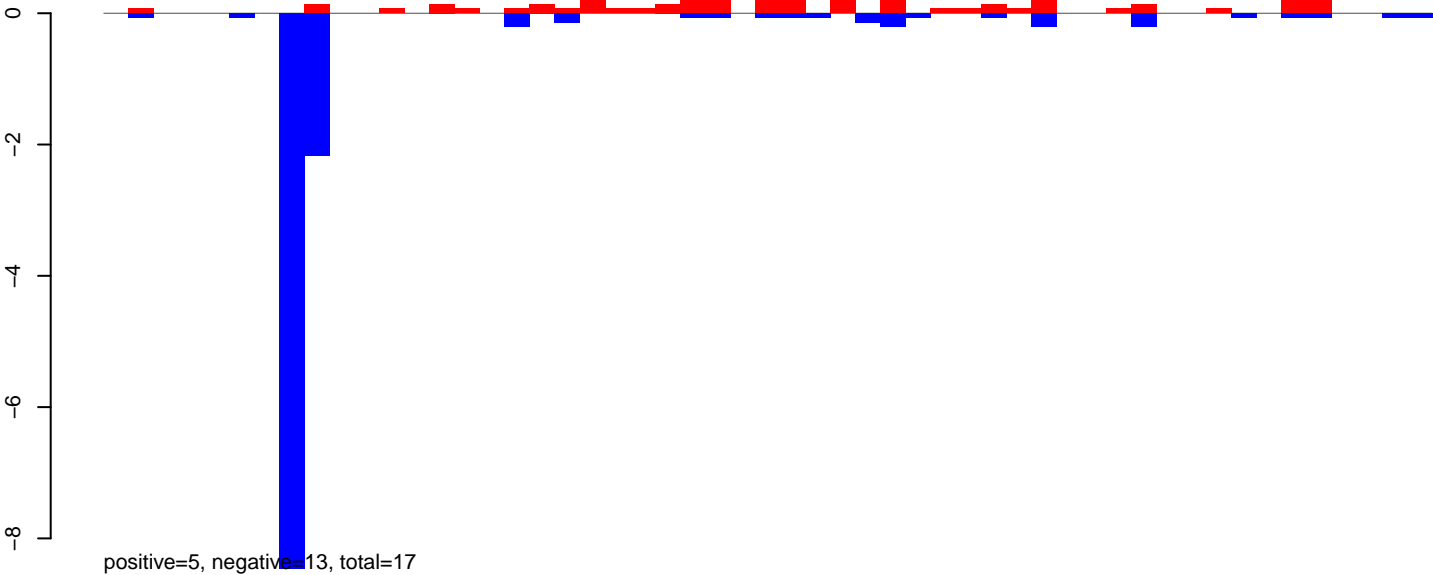


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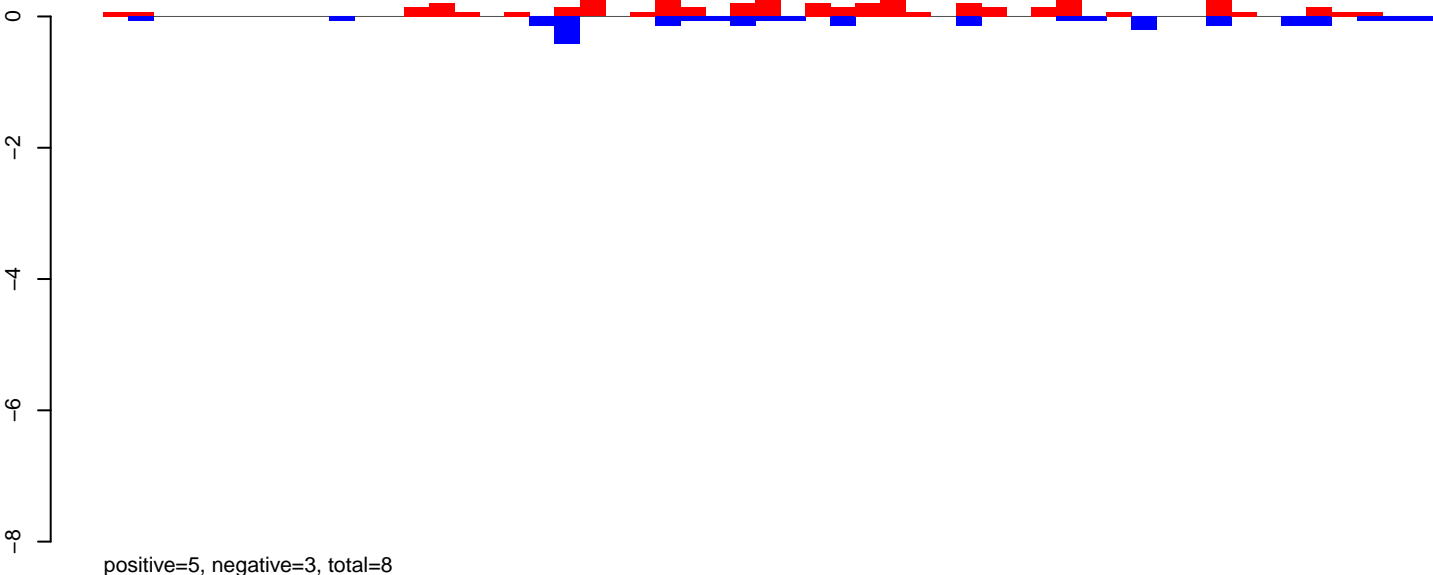


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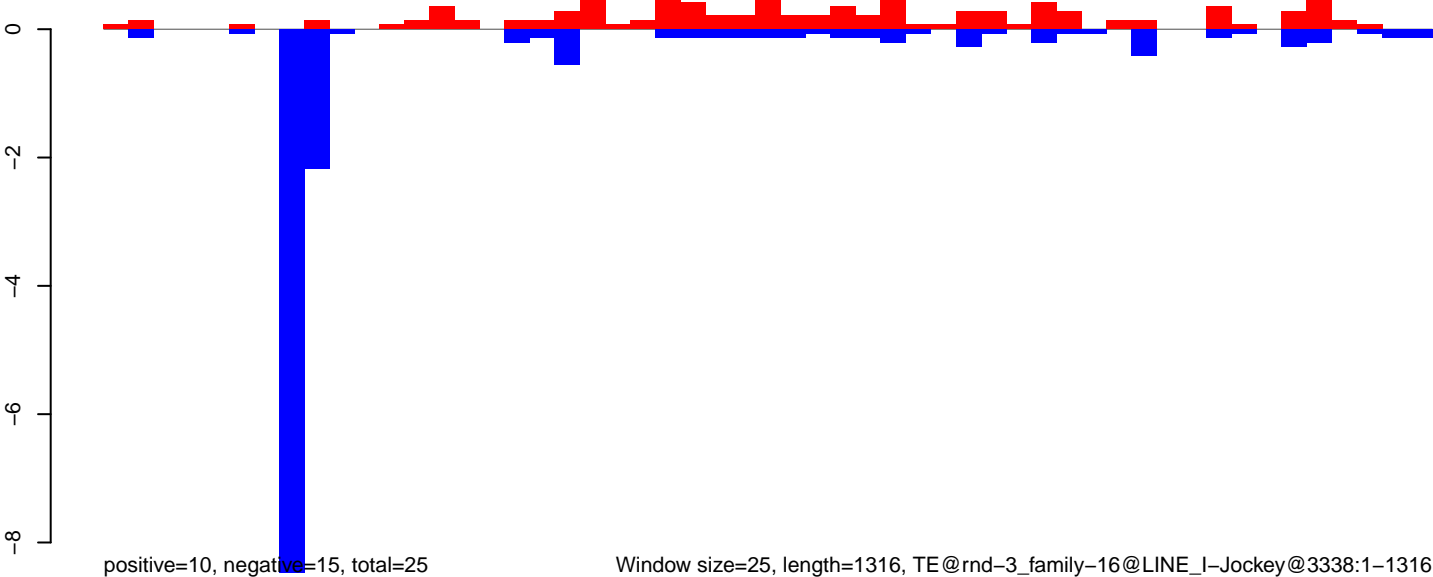
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CuQuin_Mcarc_NL.24_35.rep

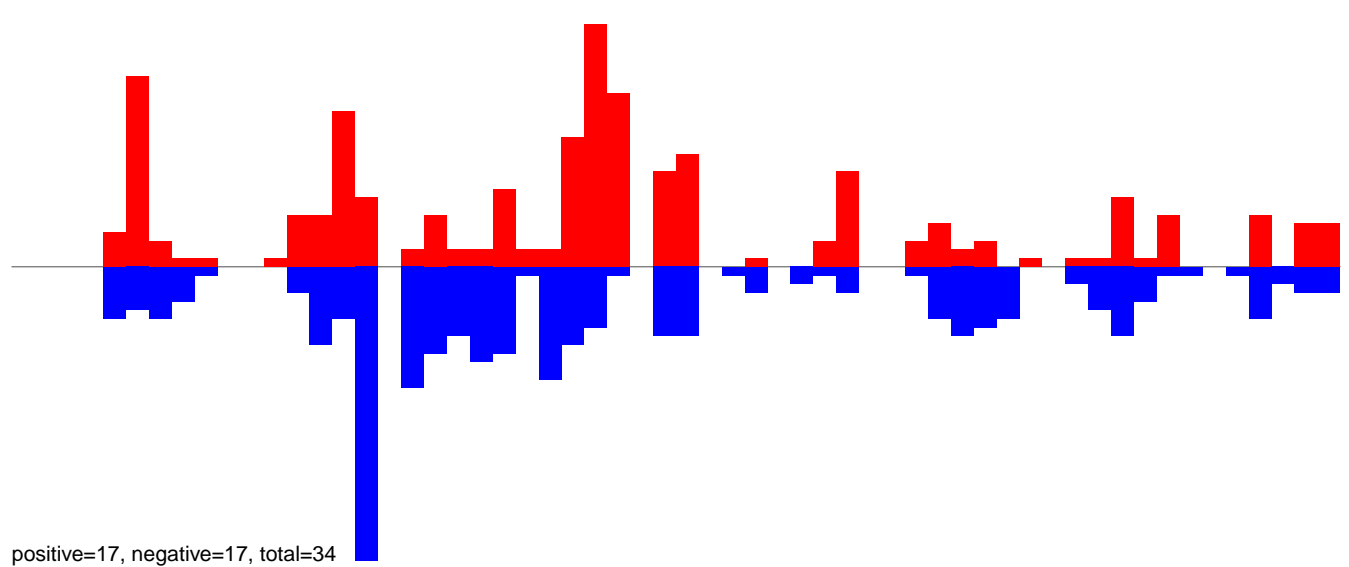


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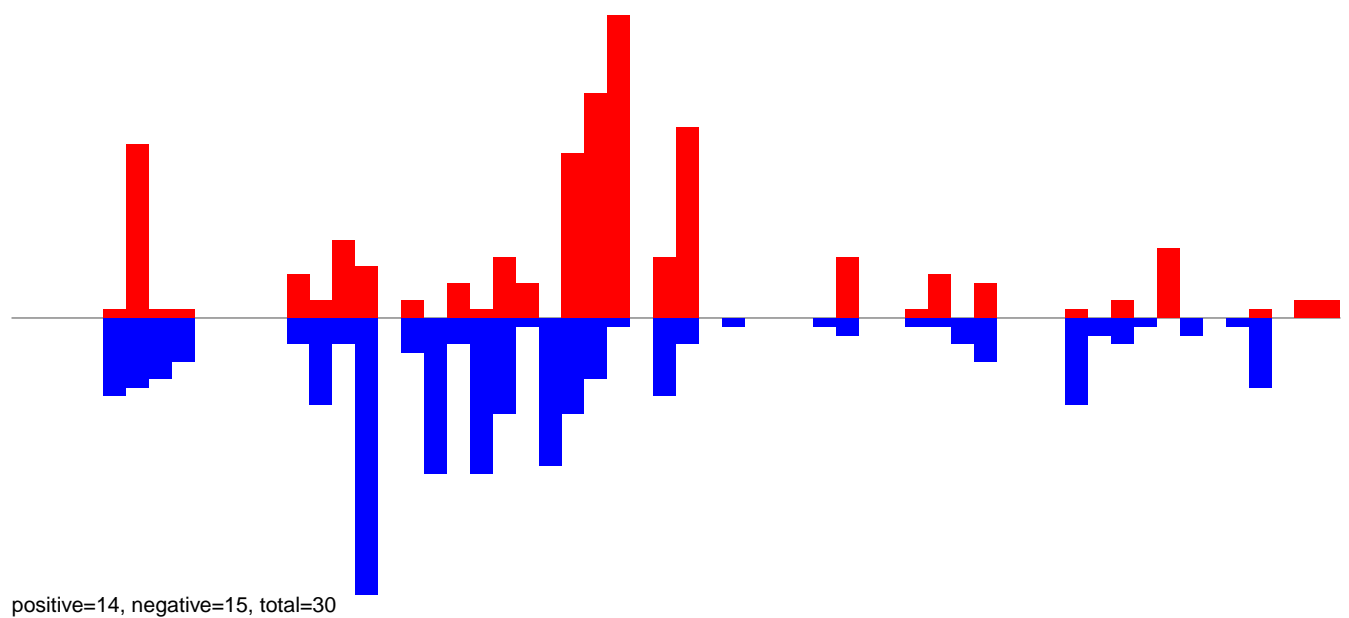


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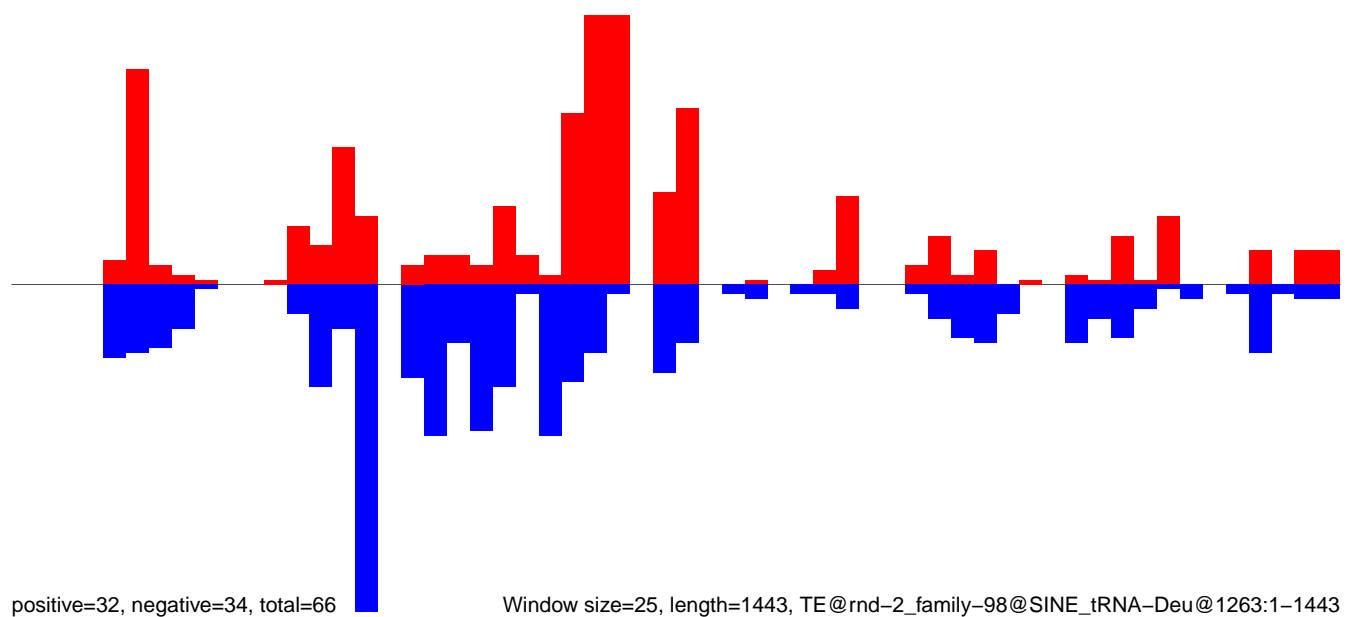
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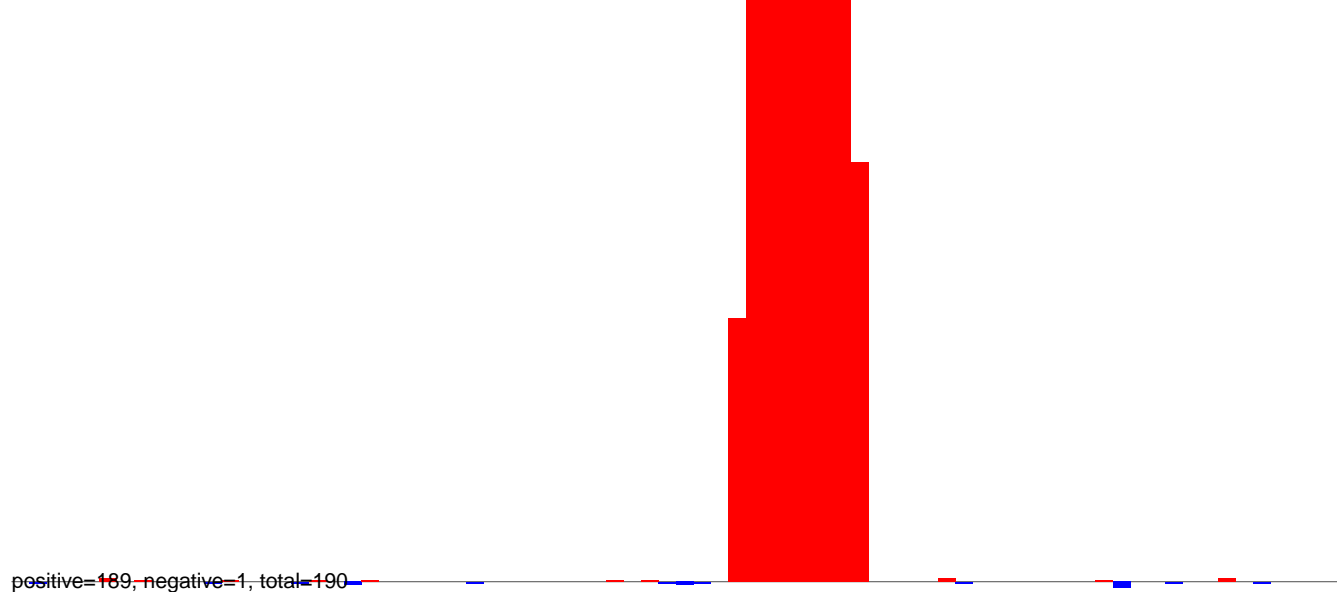
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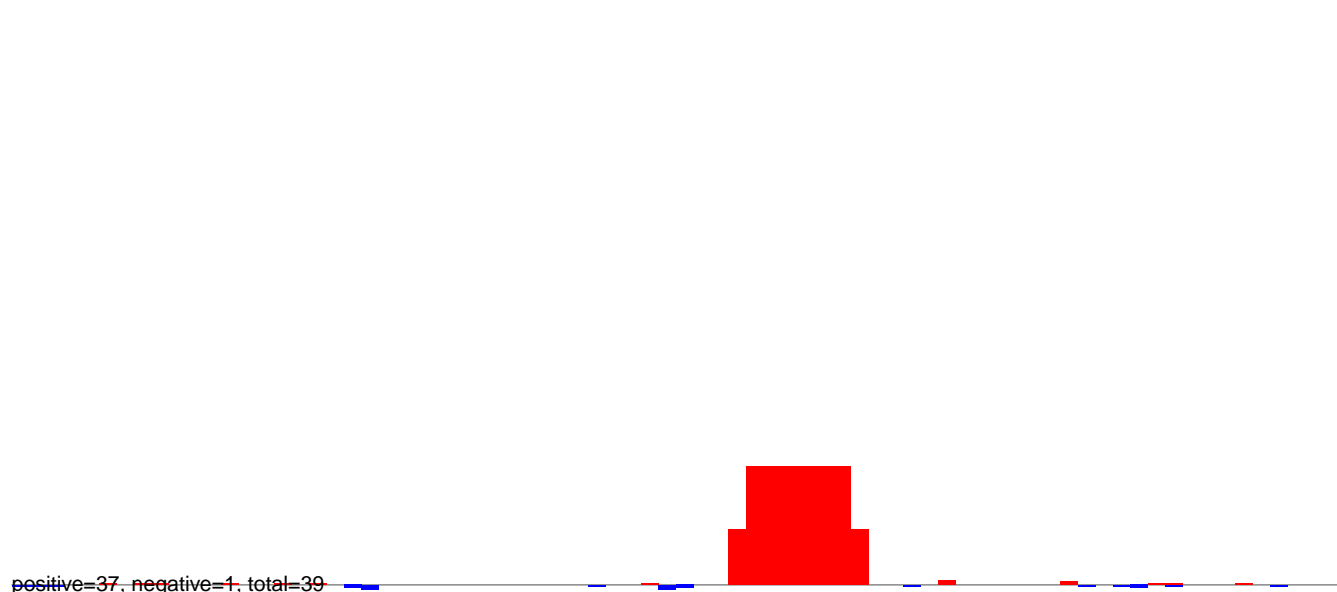
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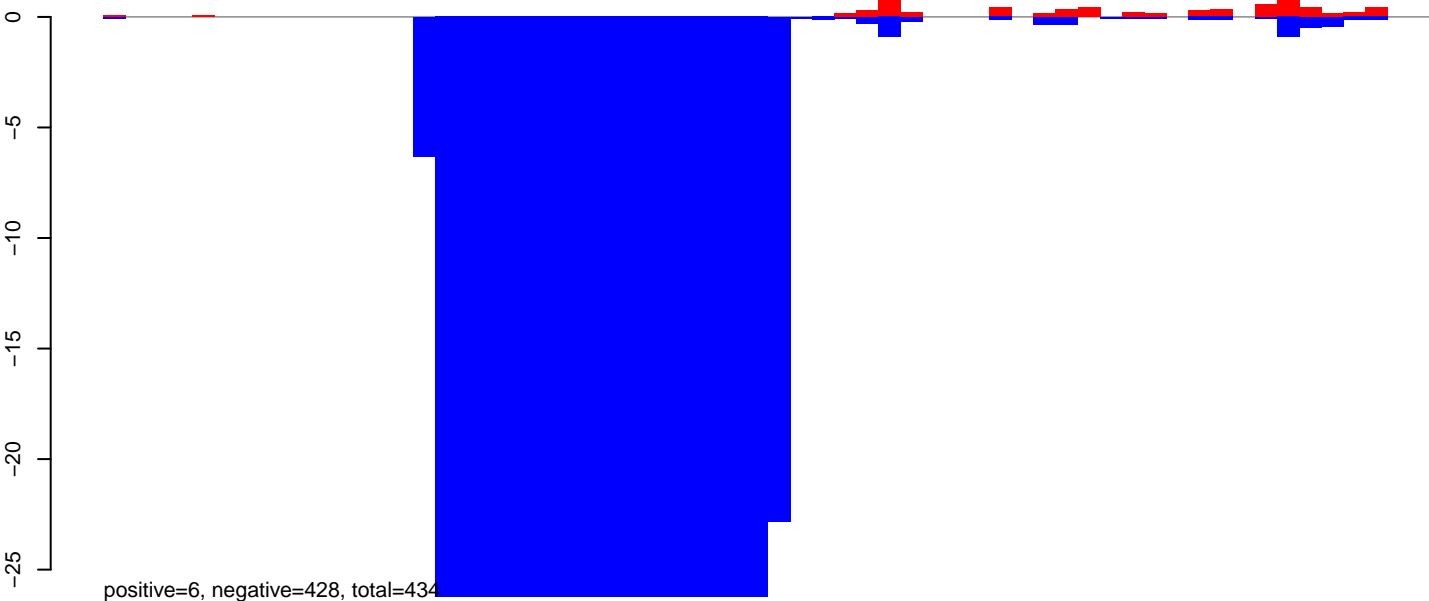
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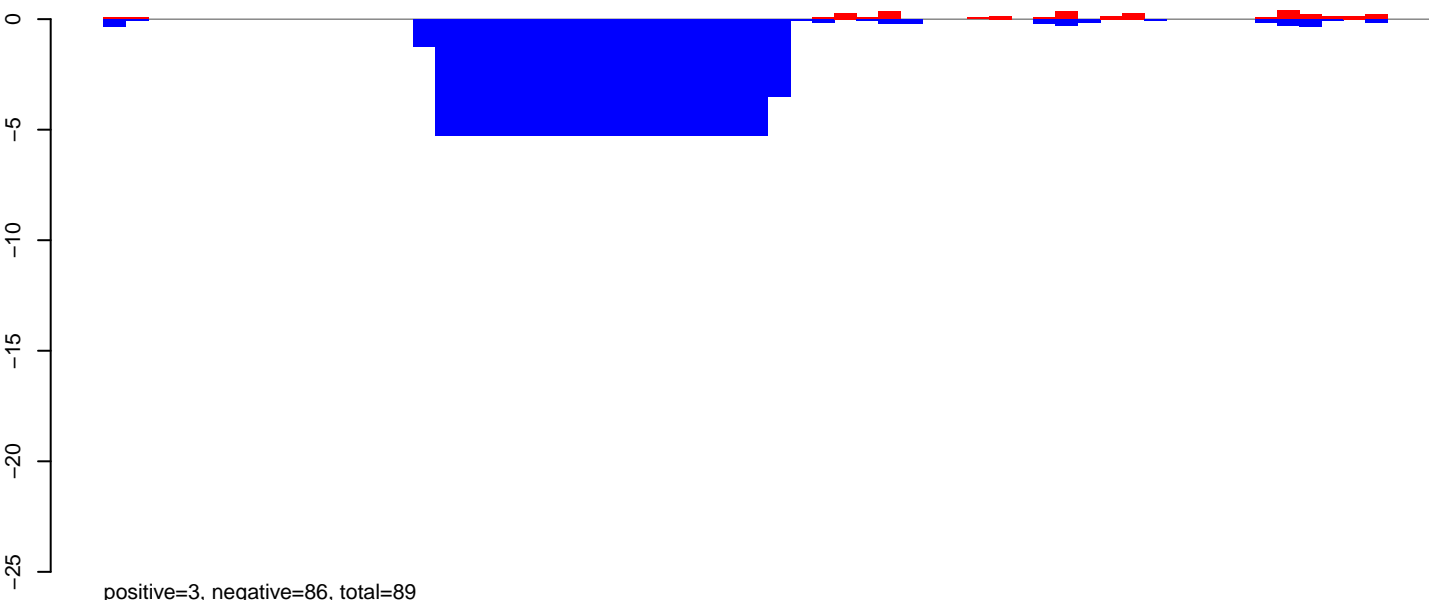
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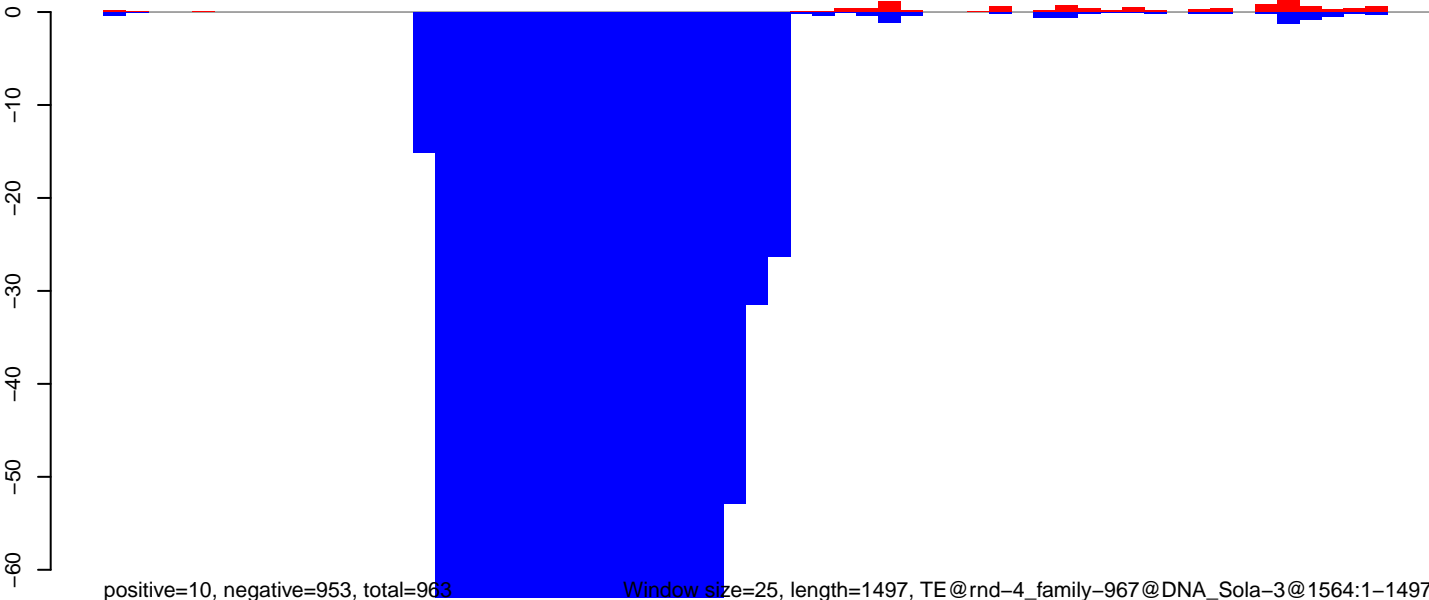
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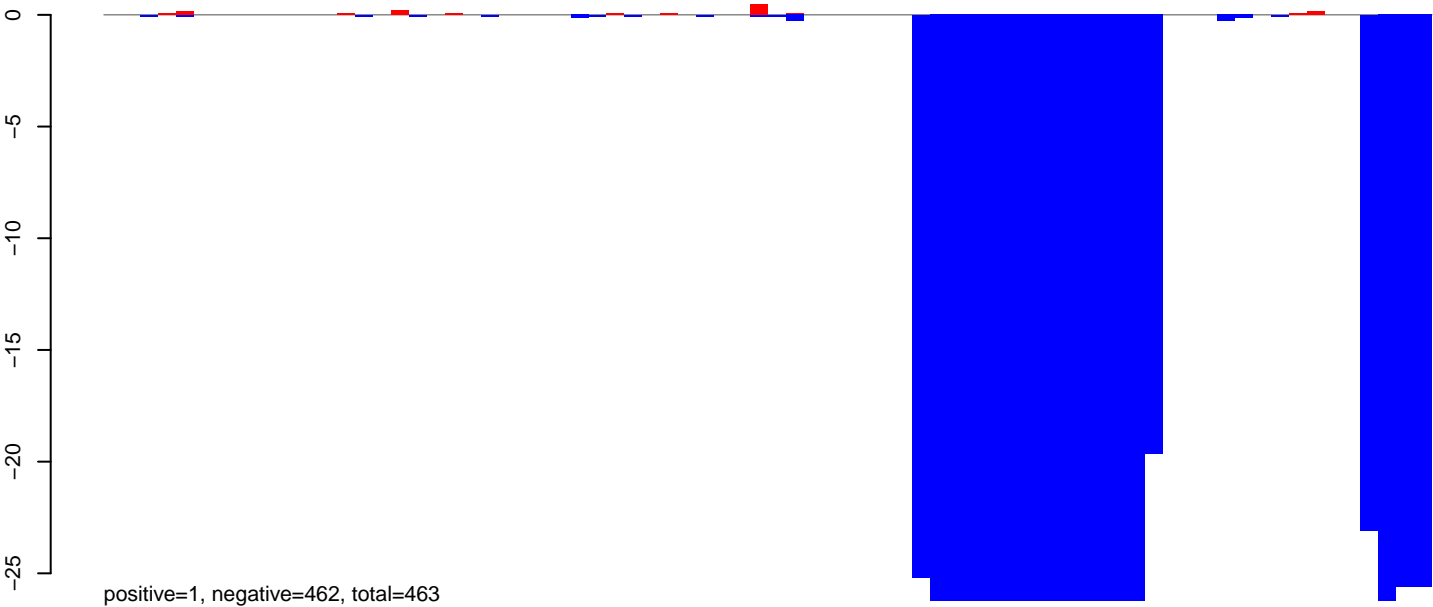


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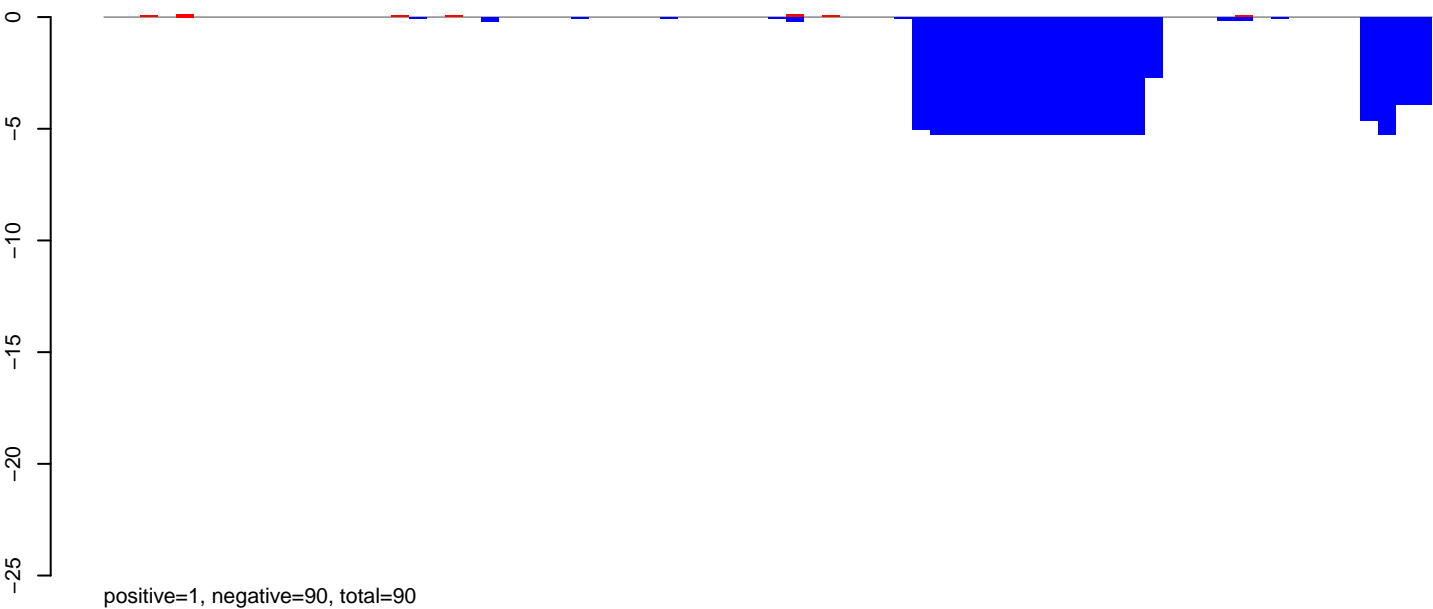


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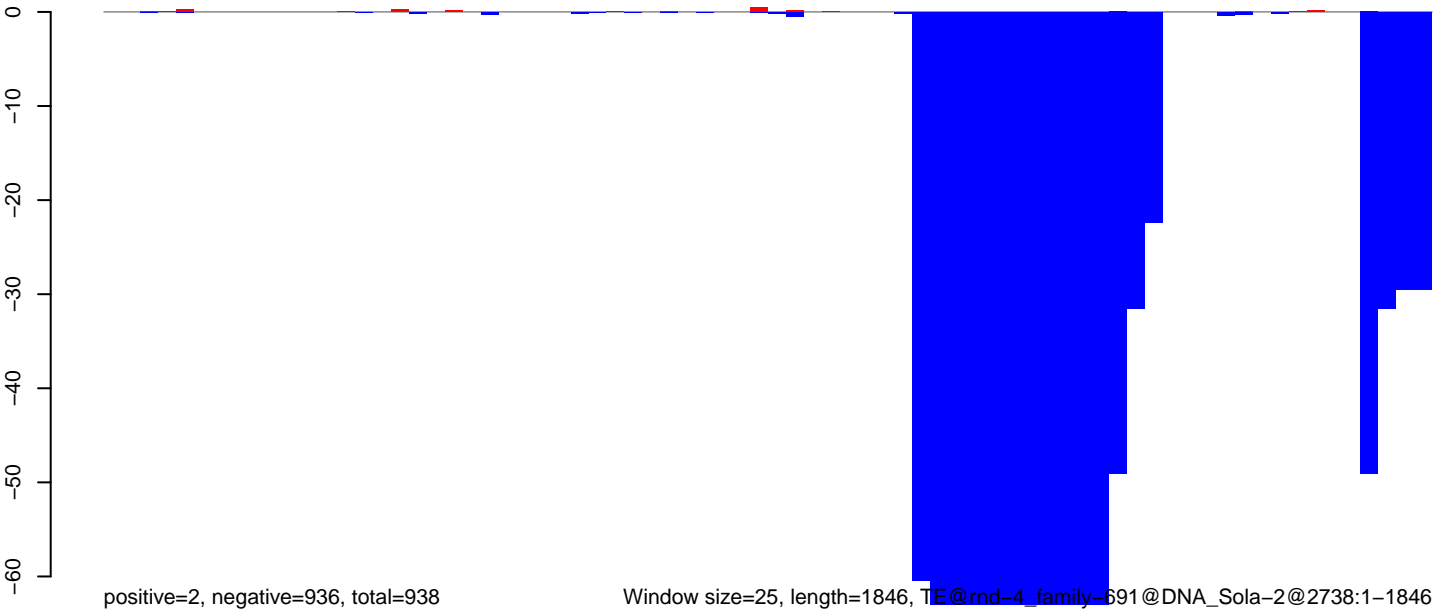
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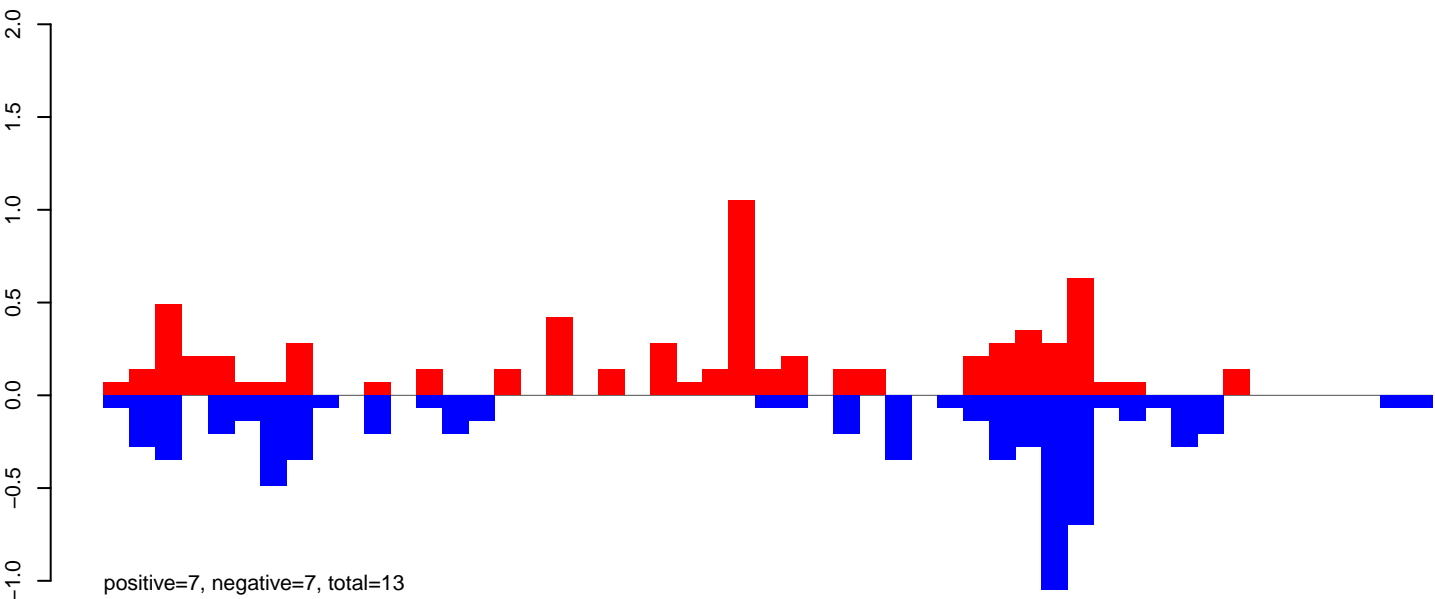
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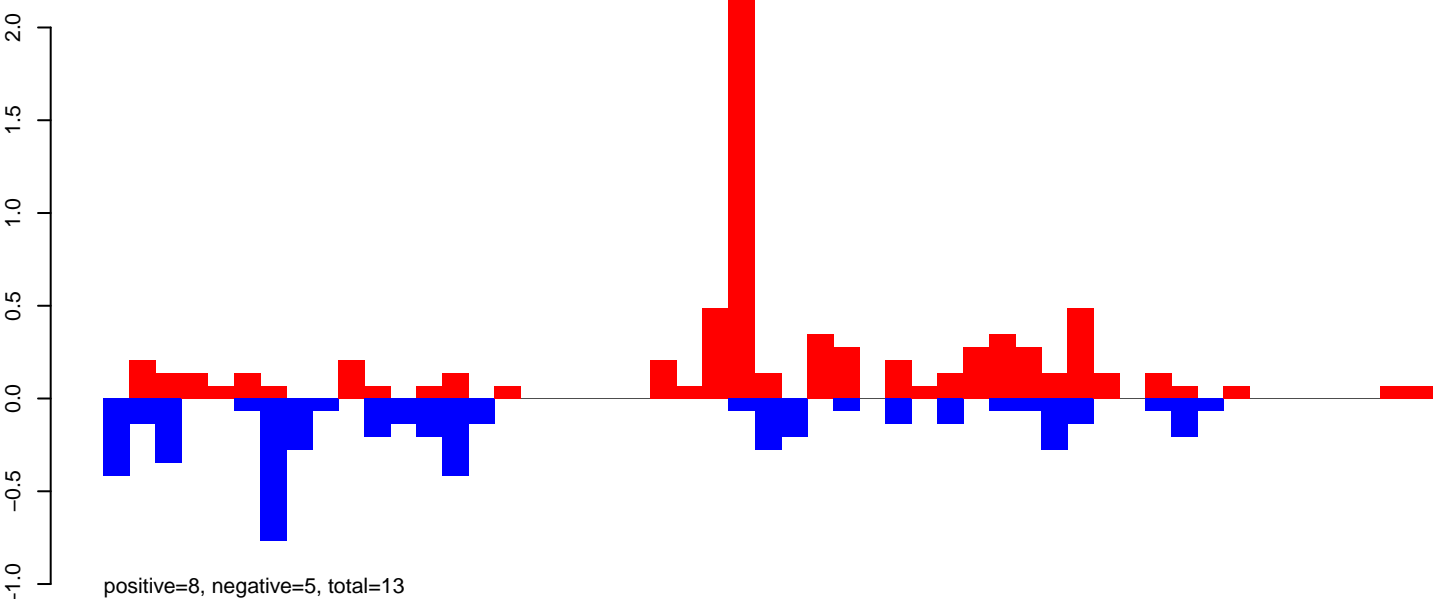
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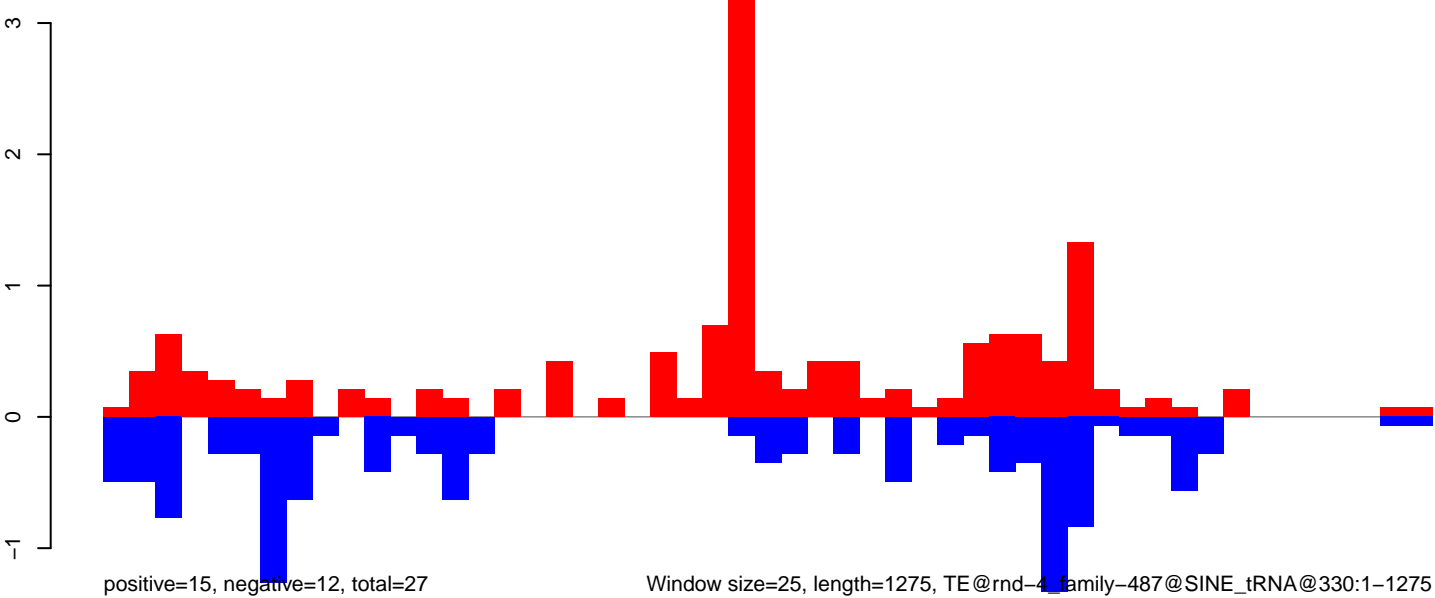
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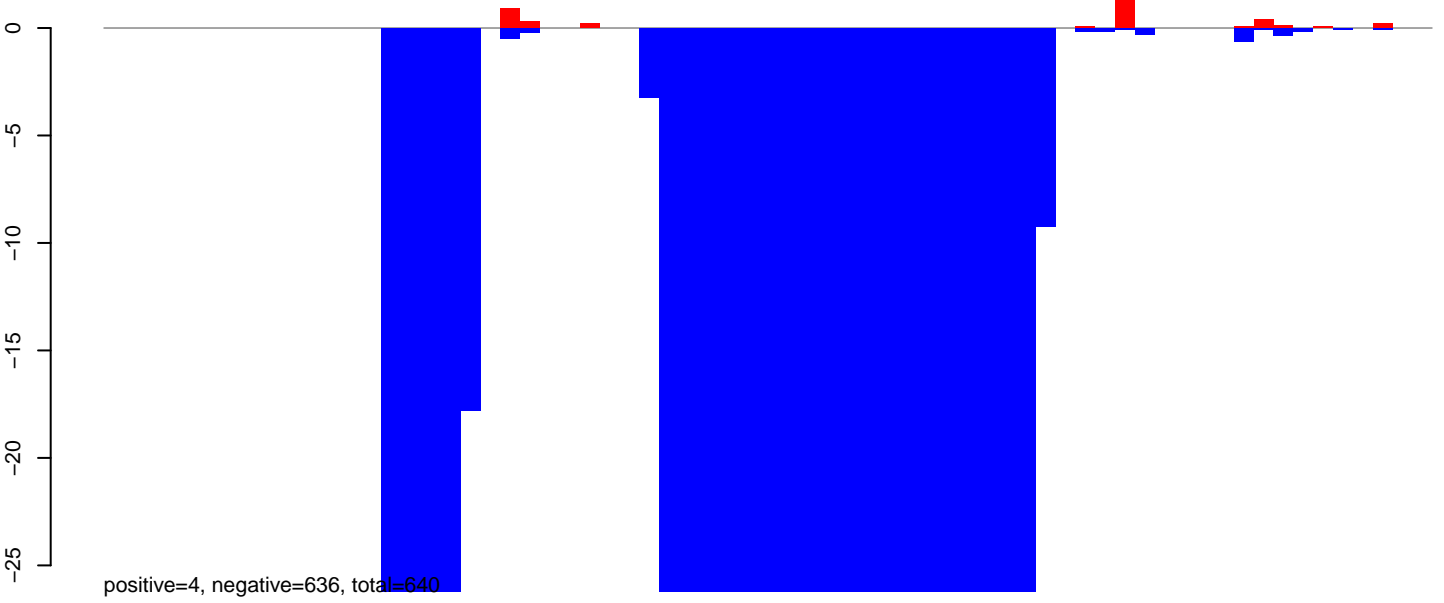


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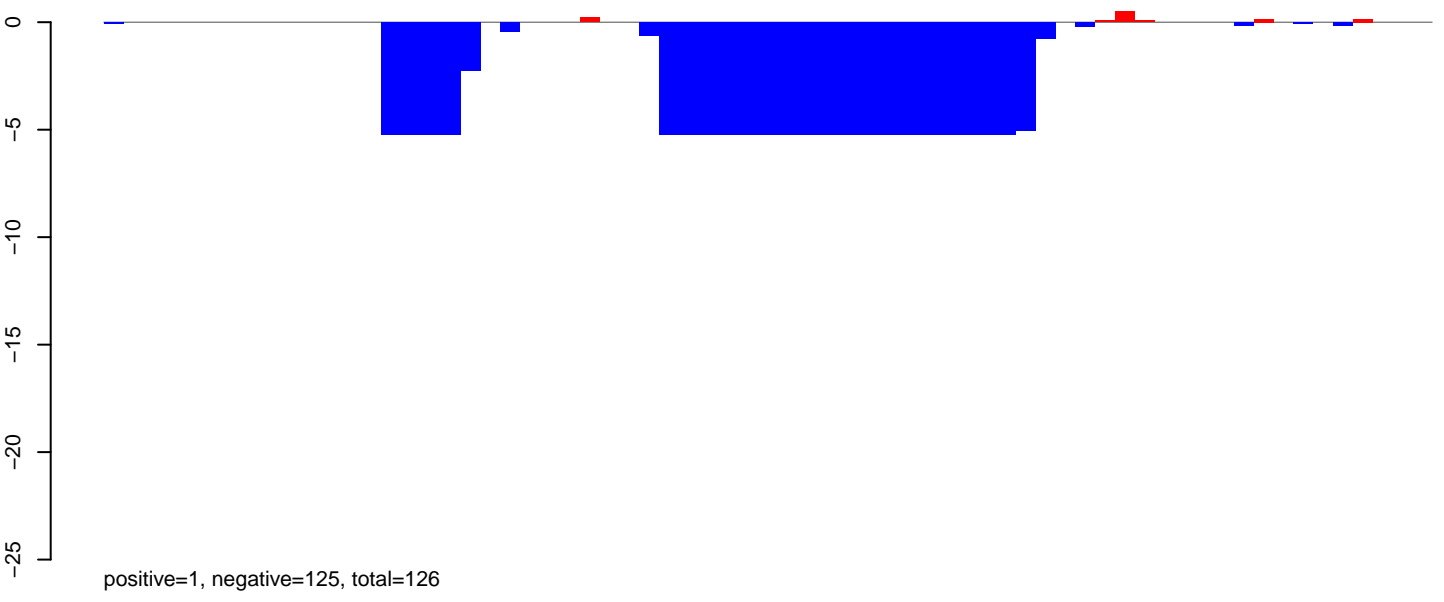


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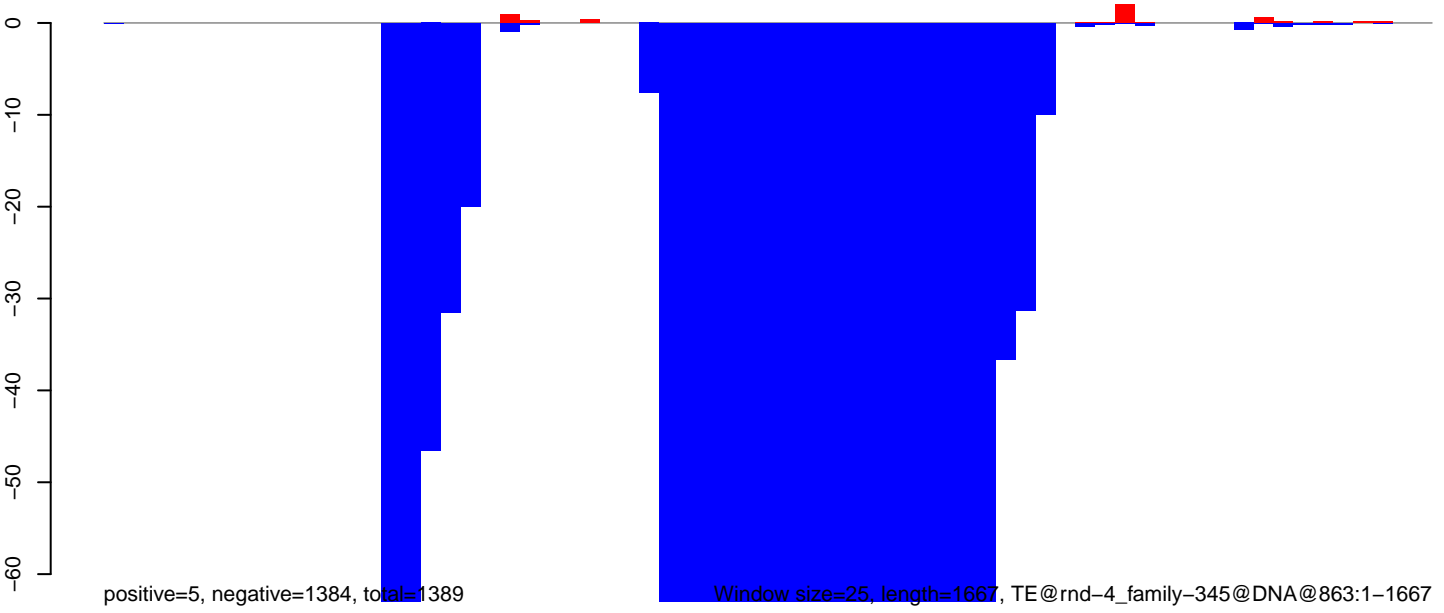
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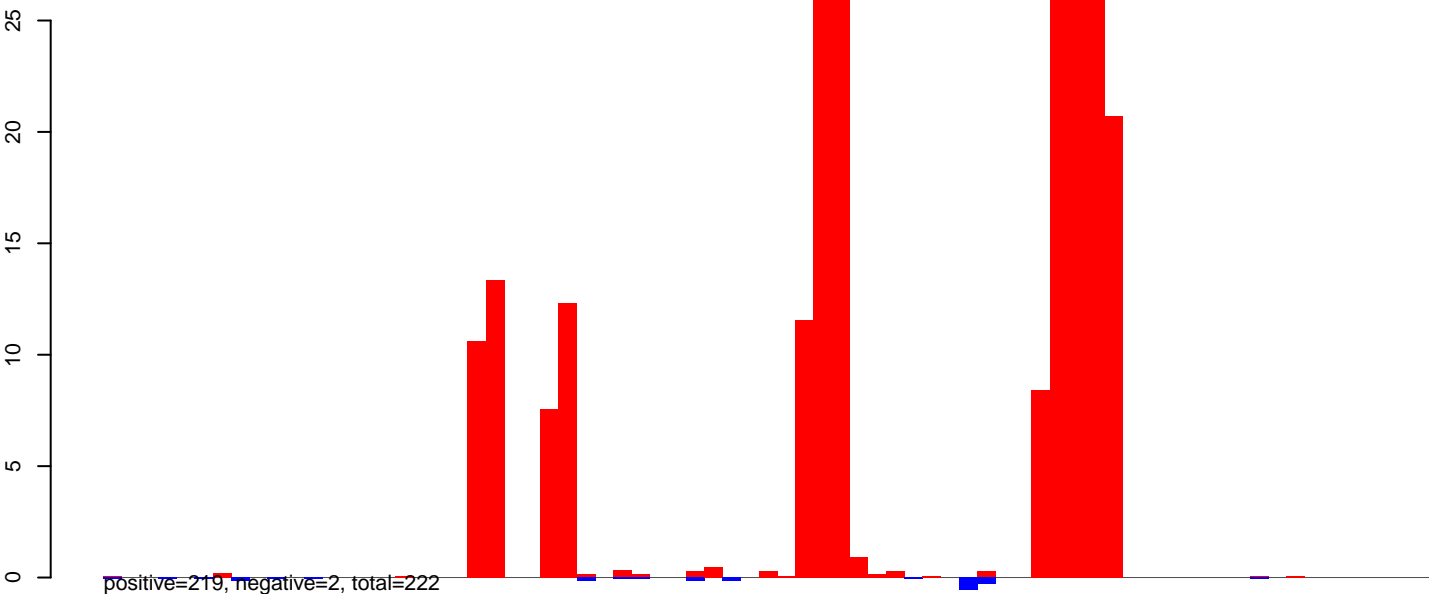
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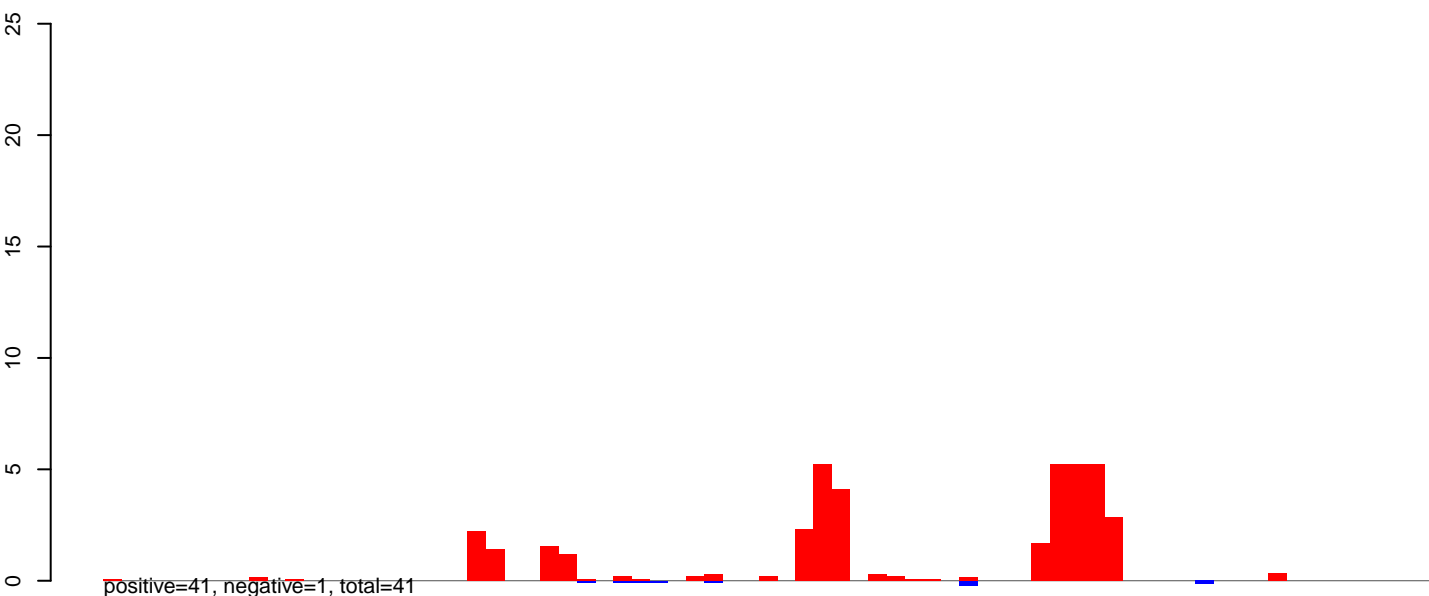
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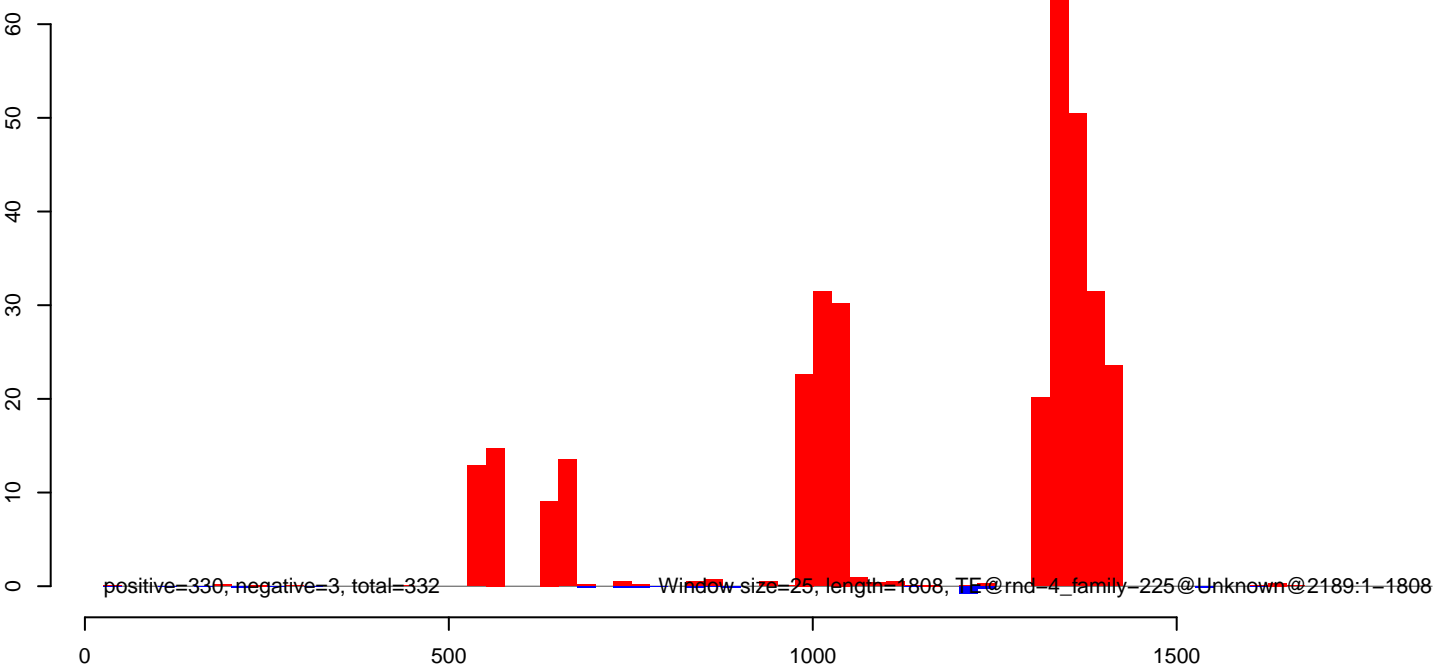
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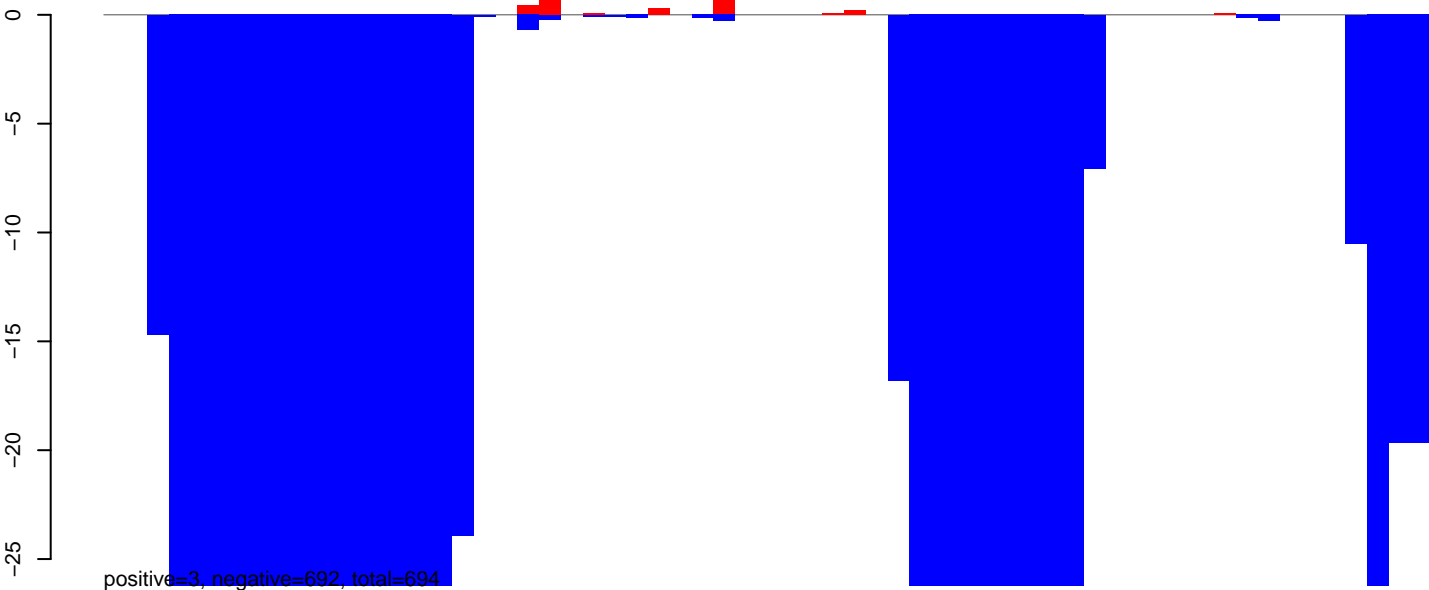
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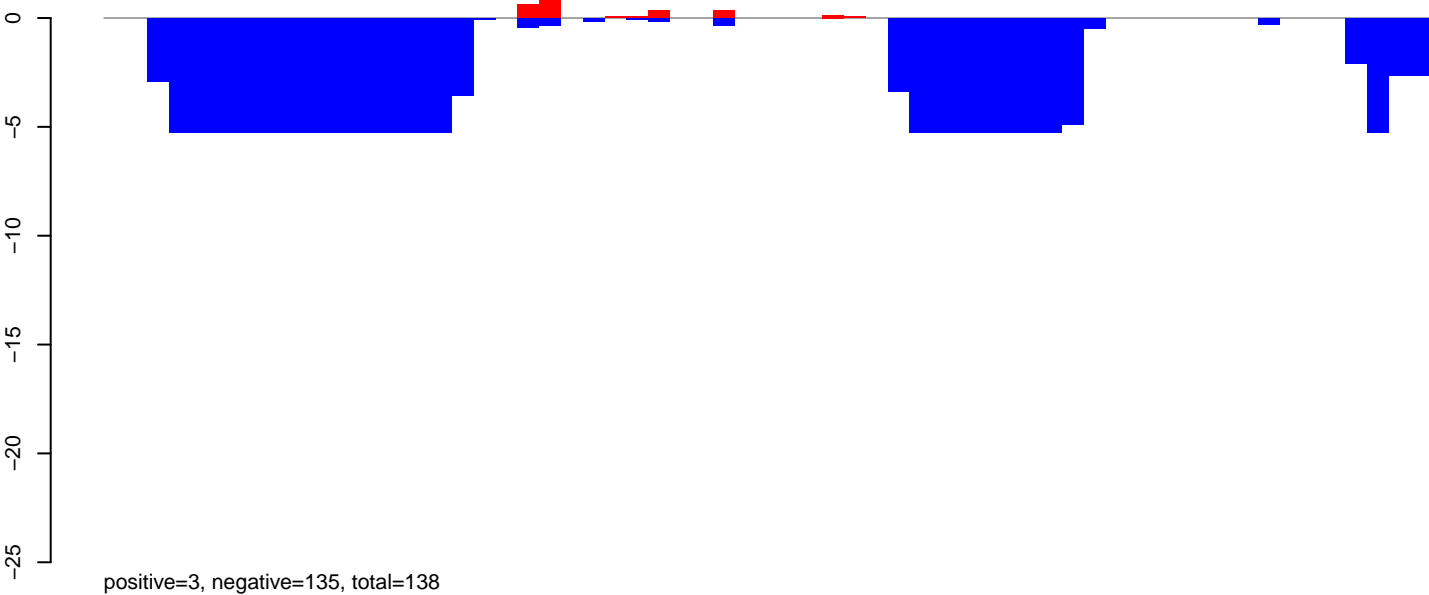
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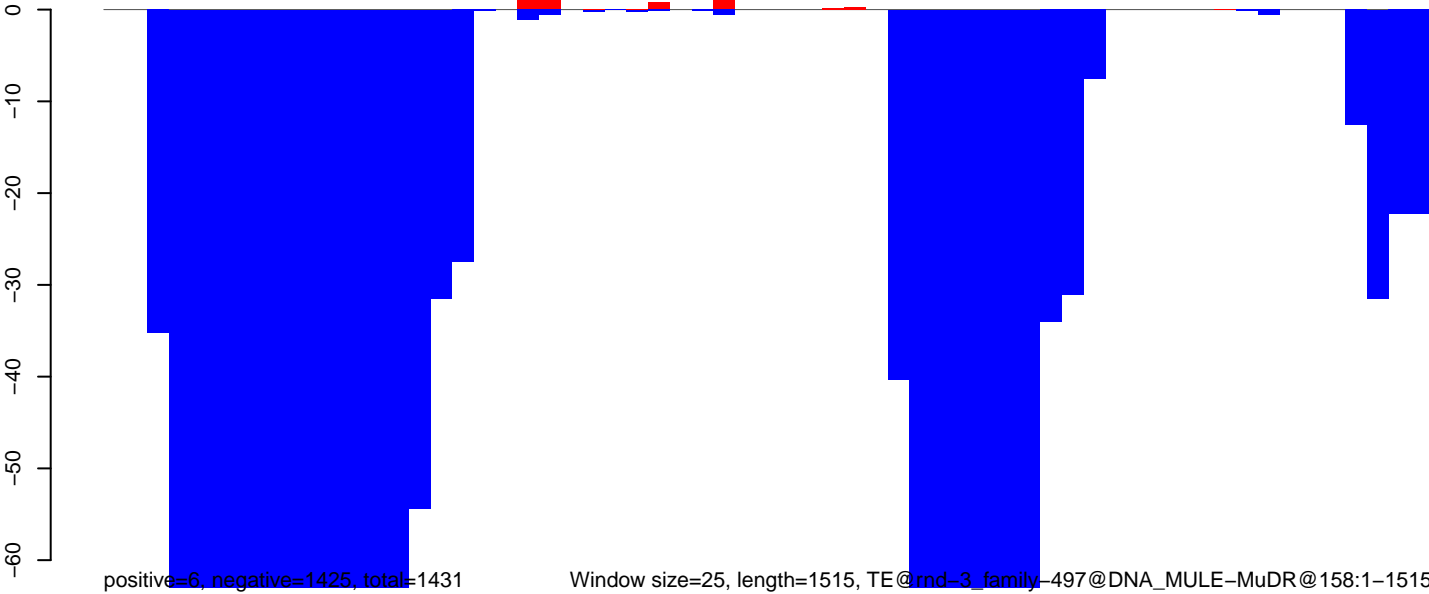
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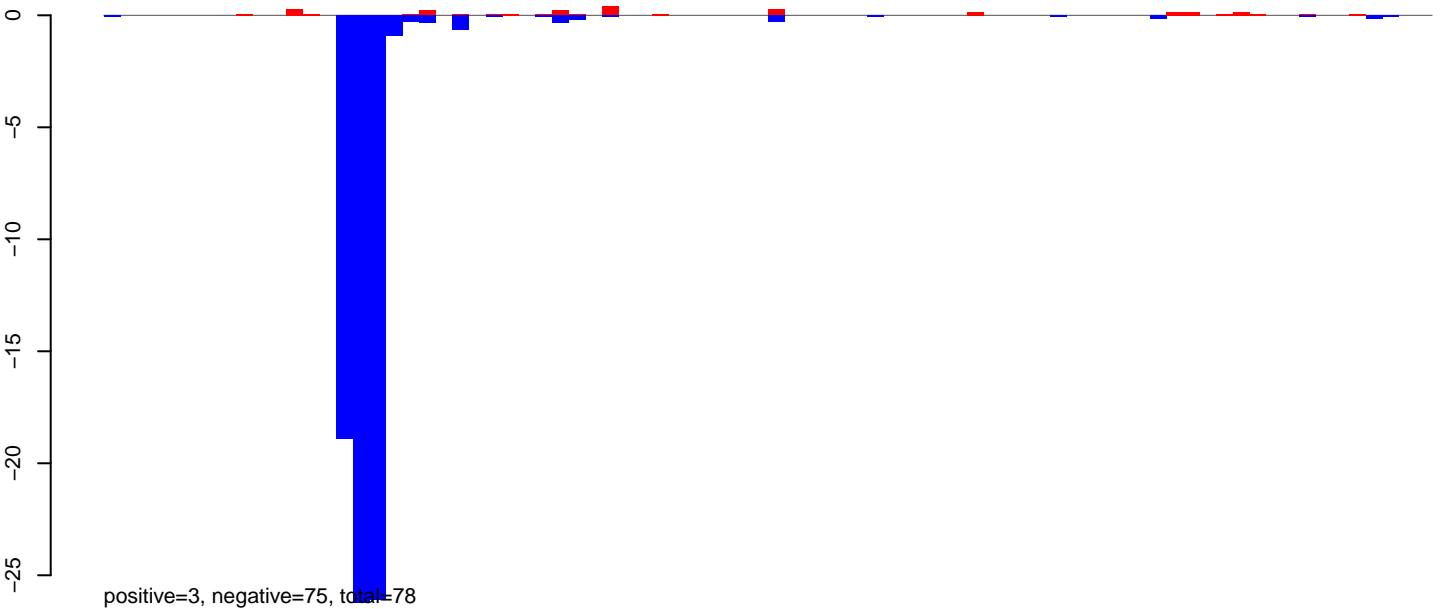
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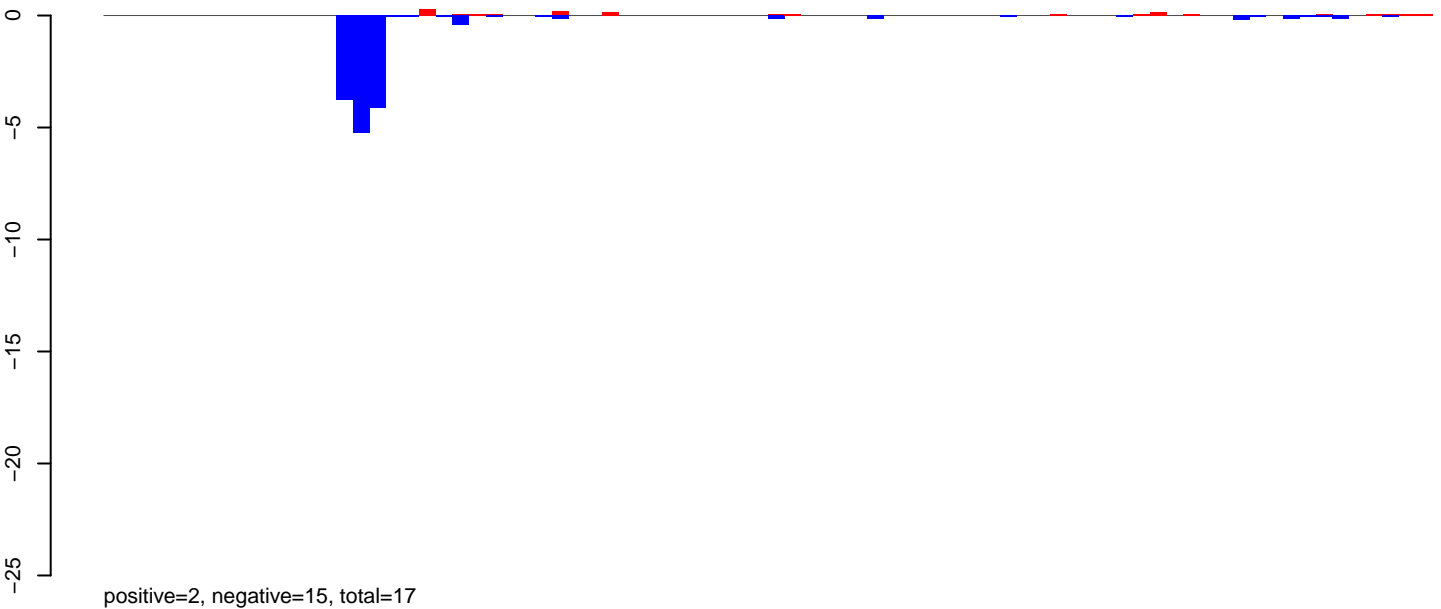
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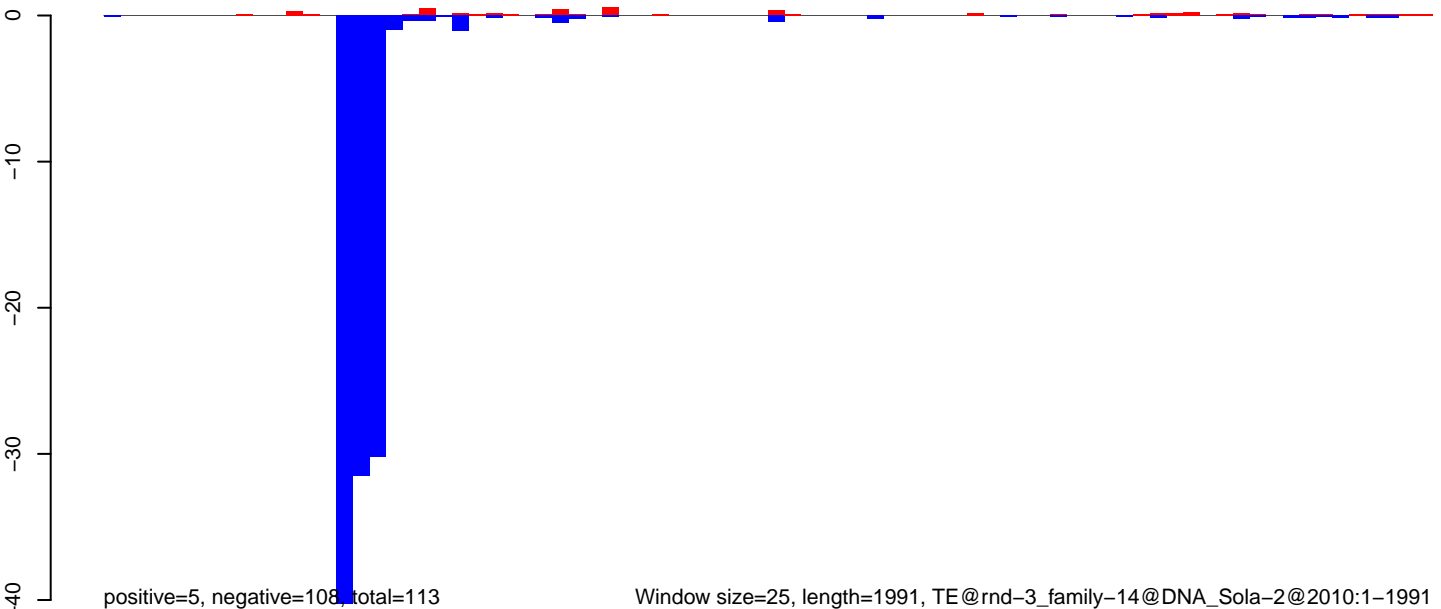
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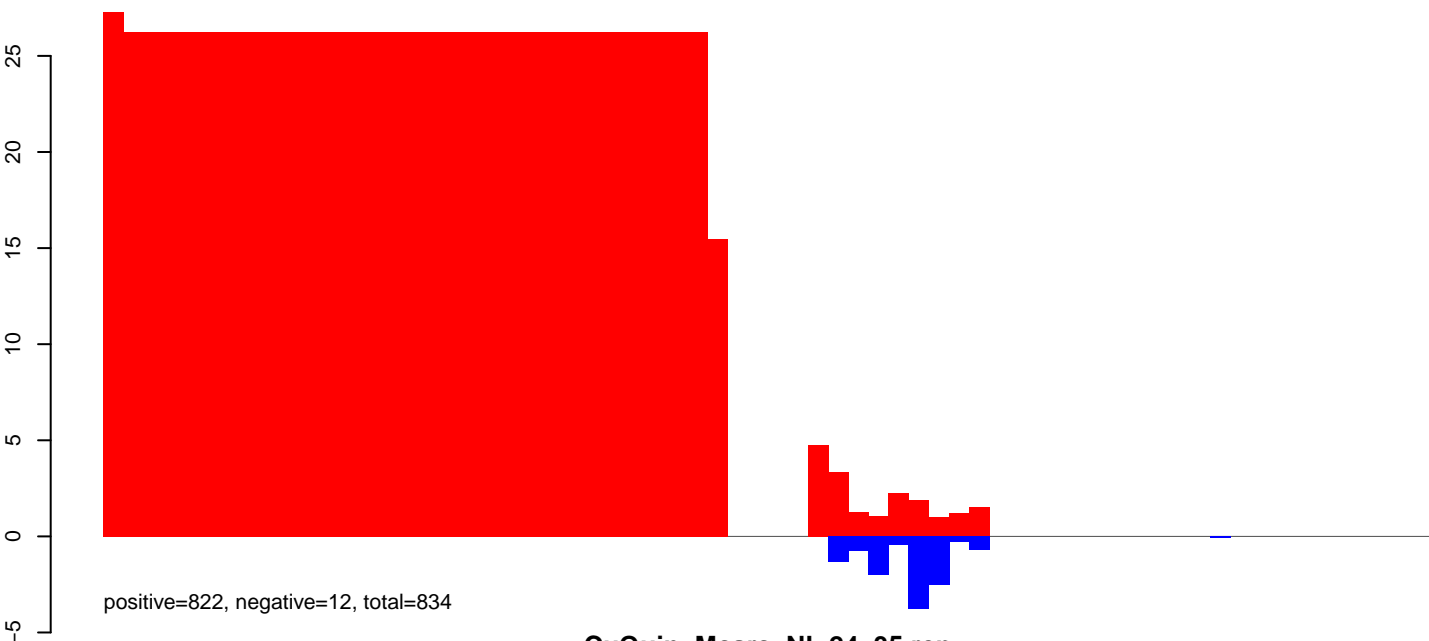
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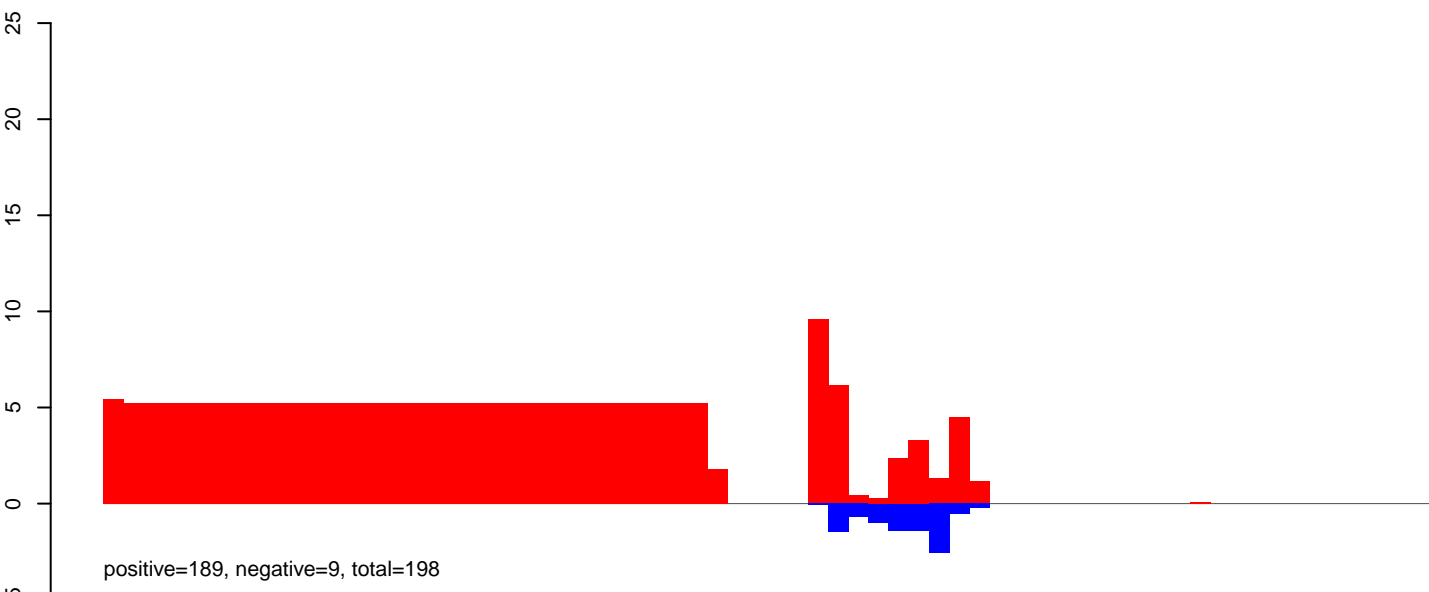
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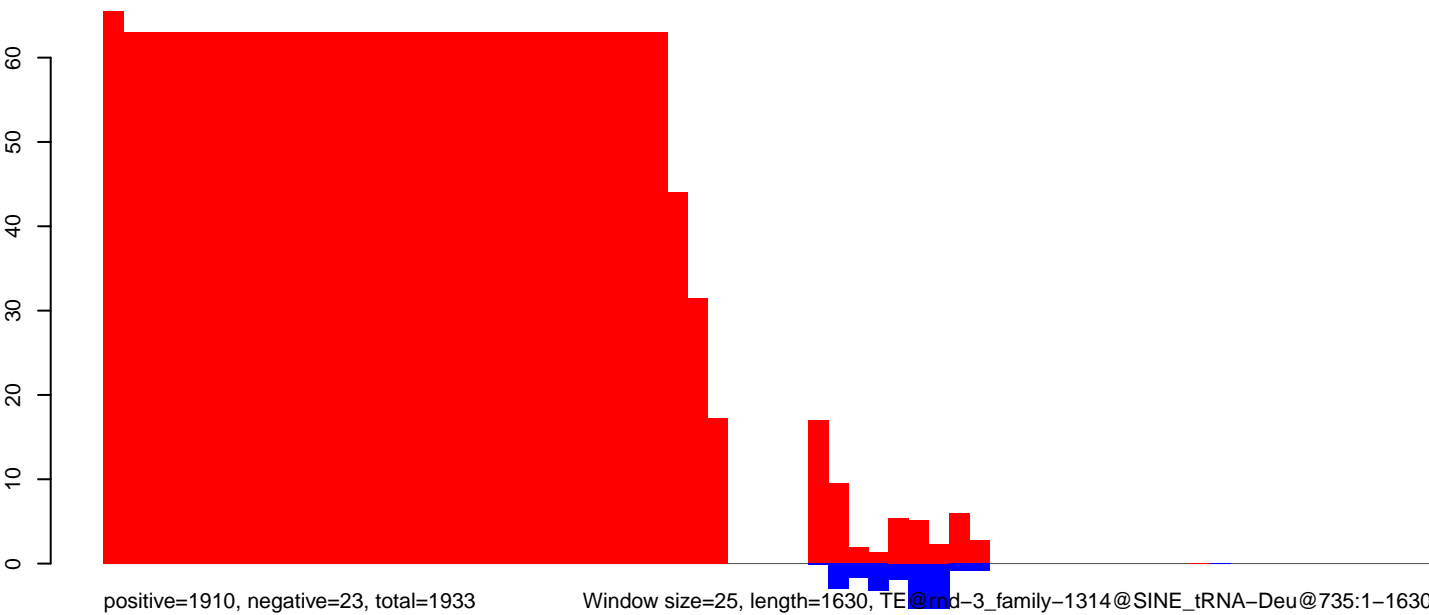
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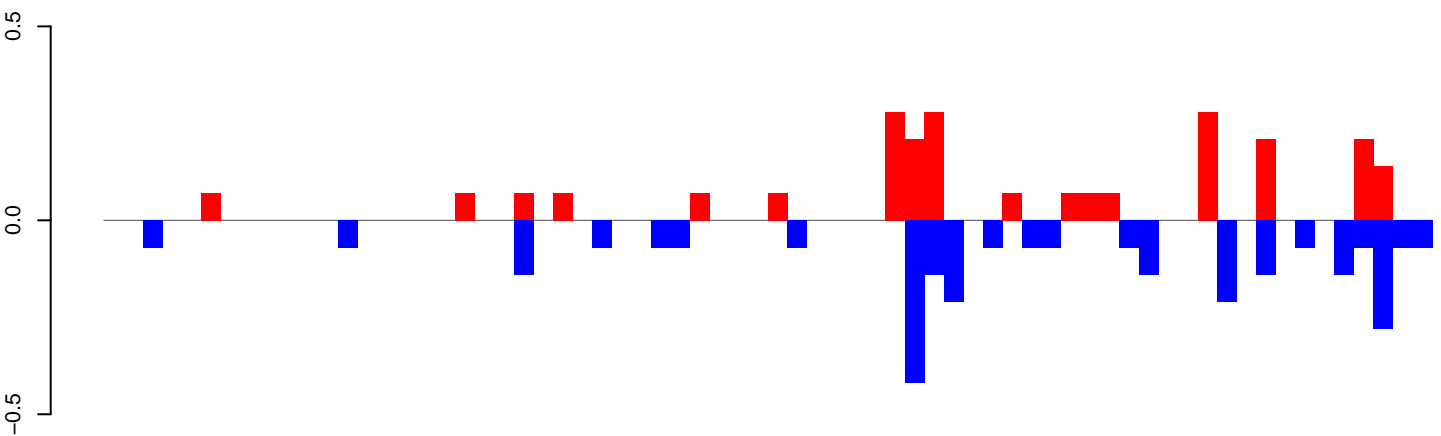


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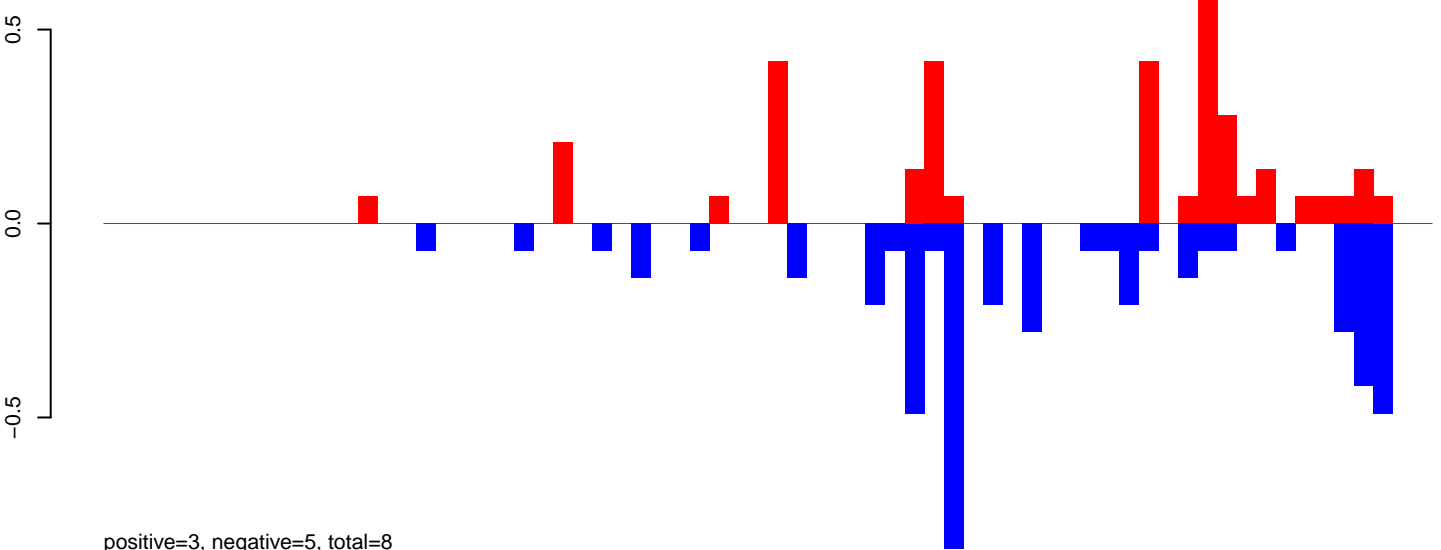
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CuQuin_Mcarc_NL.18_23.rep



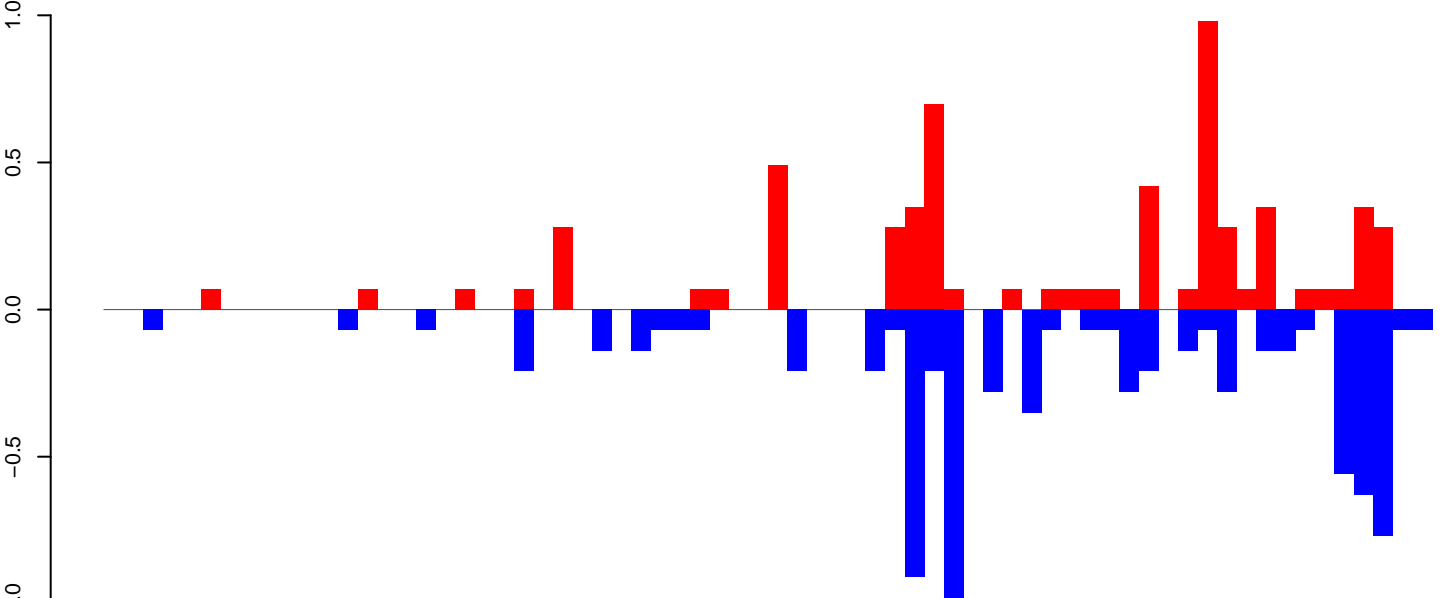
positive=2, negative=3, total=5

CuQuin_Mcarc_NL.24_35.rep



positive=3, negative=5, total=8

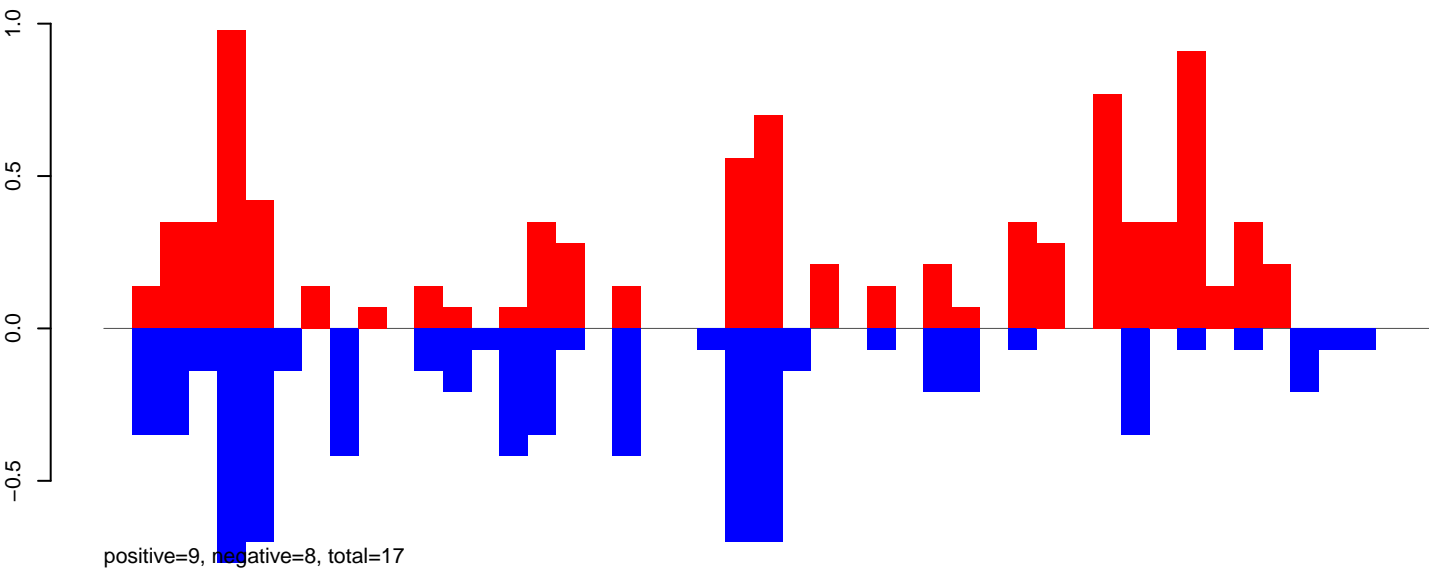
CuQuin_Mcarc_NL.rep



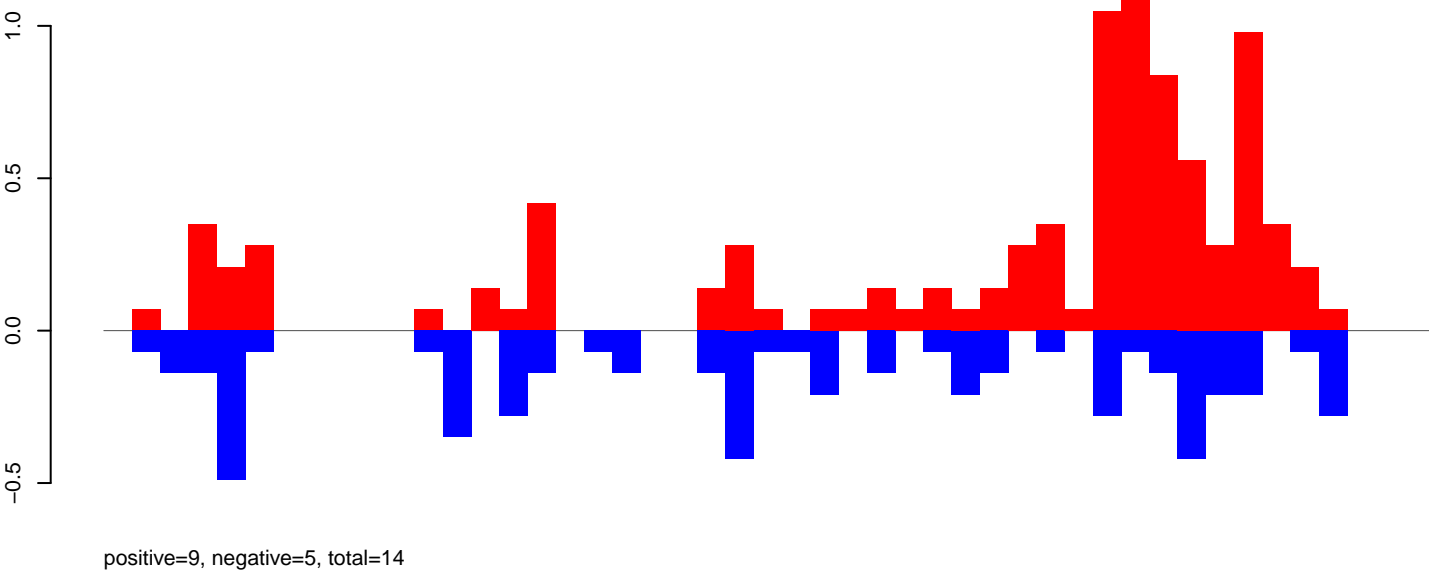
positive=6, negative=8, total=14

Window size=25, length=1687, TE@md-3_family-111@DNA_hAT-hAT19@665:1-1687

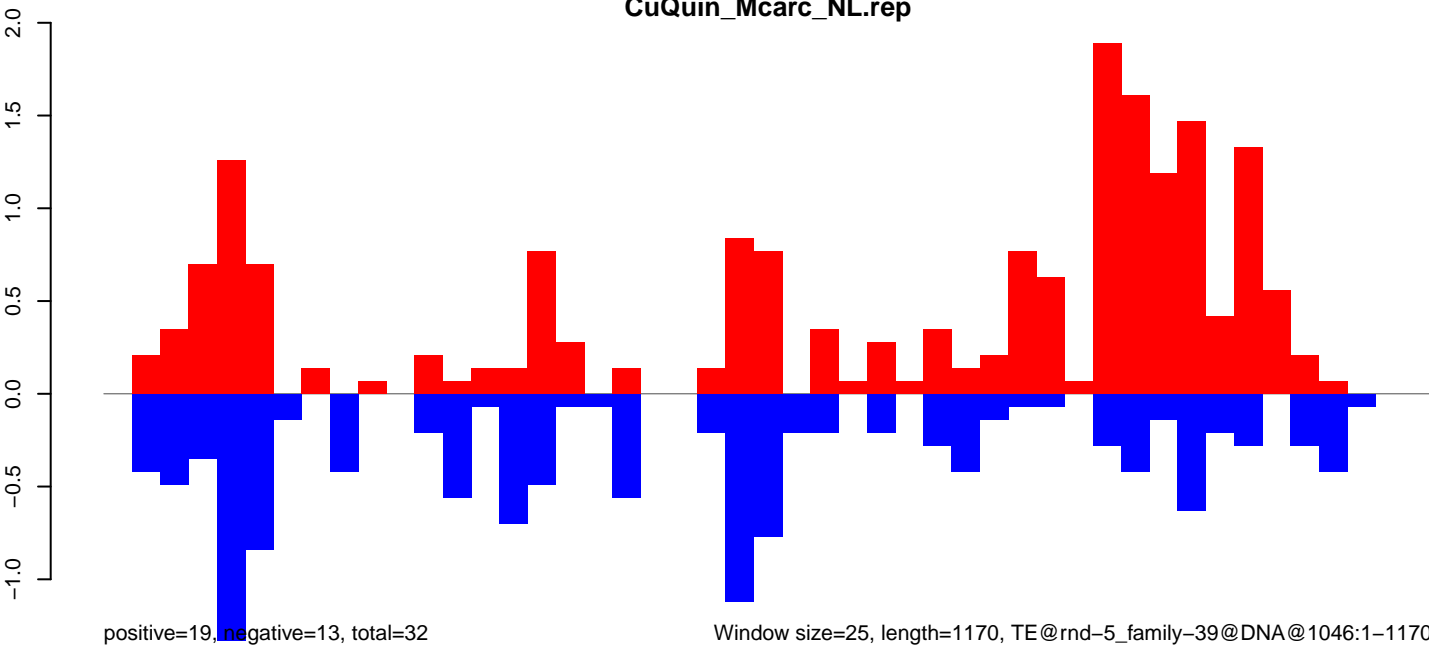
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



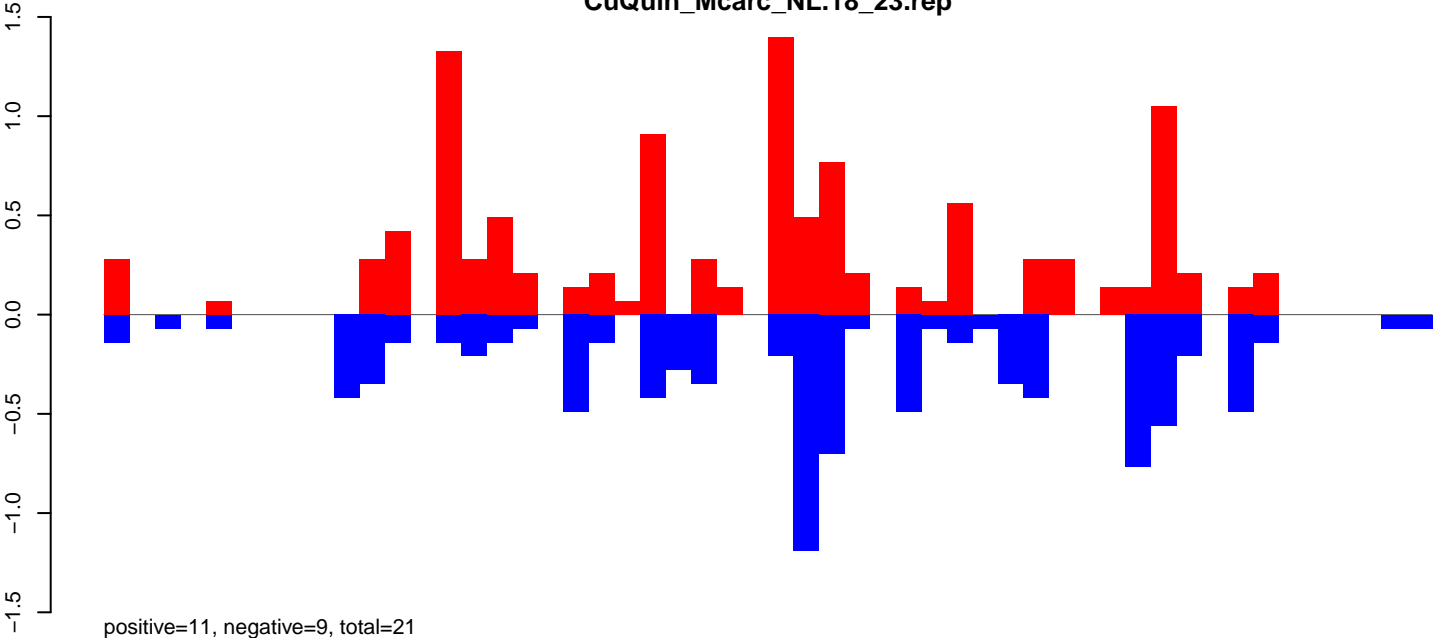
CuQuin_Mcarc_NL.rep



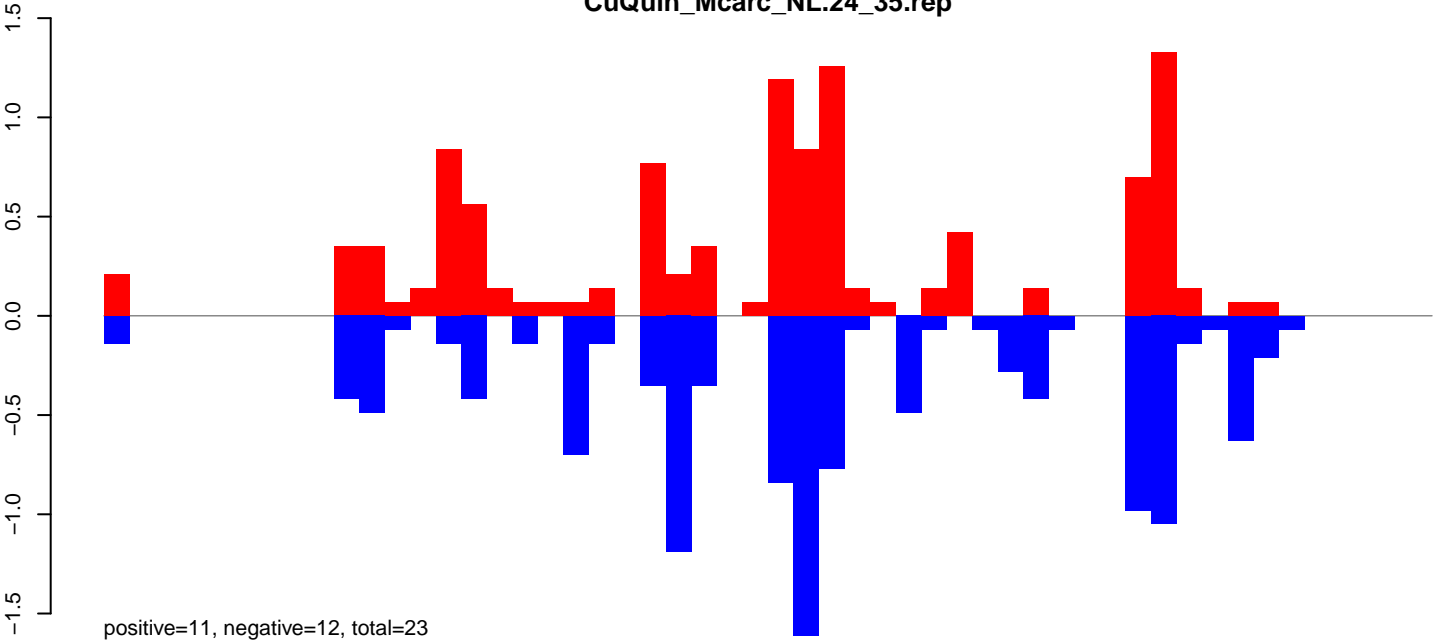
Window size=25, length=1170, TE@rnd-5_family-39@DNA@1046:1-1170

0 200 400 600 800 1000 1200

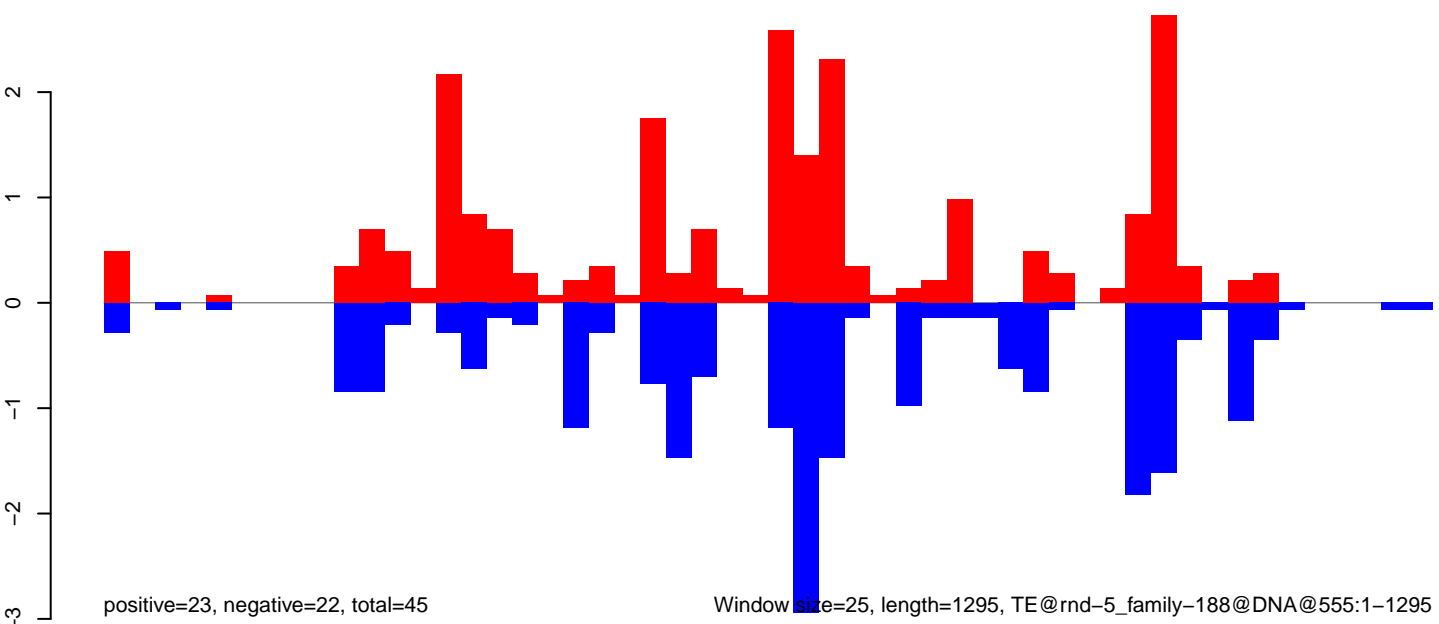
CuQuin_Mcarc_NL.18_23.rep



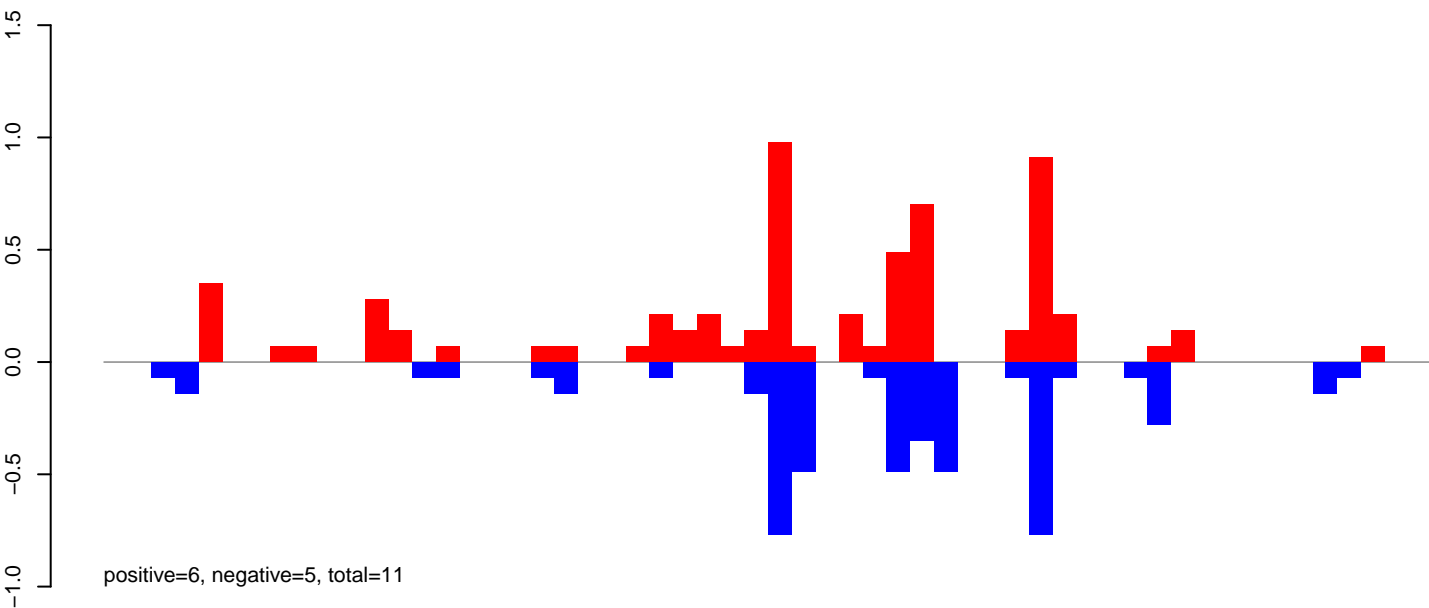
CuQuin_Mcarc_NL.24_35.rep



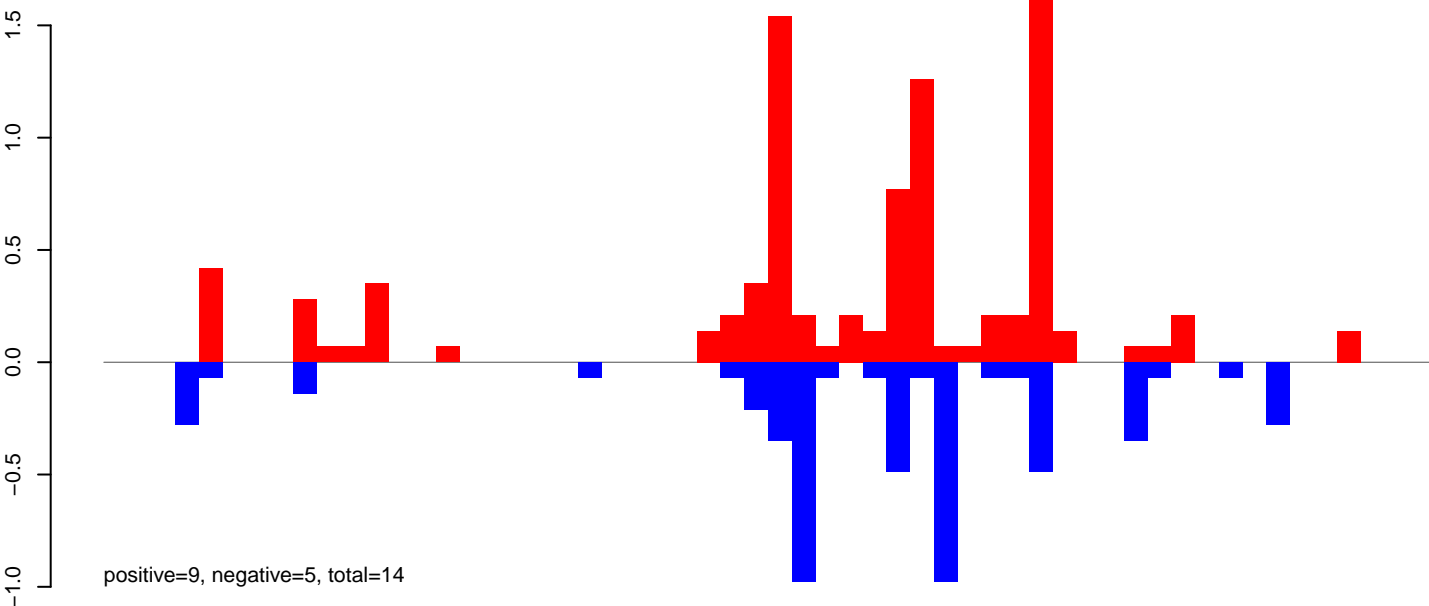
CuQuin_Mcarc_NL.rep



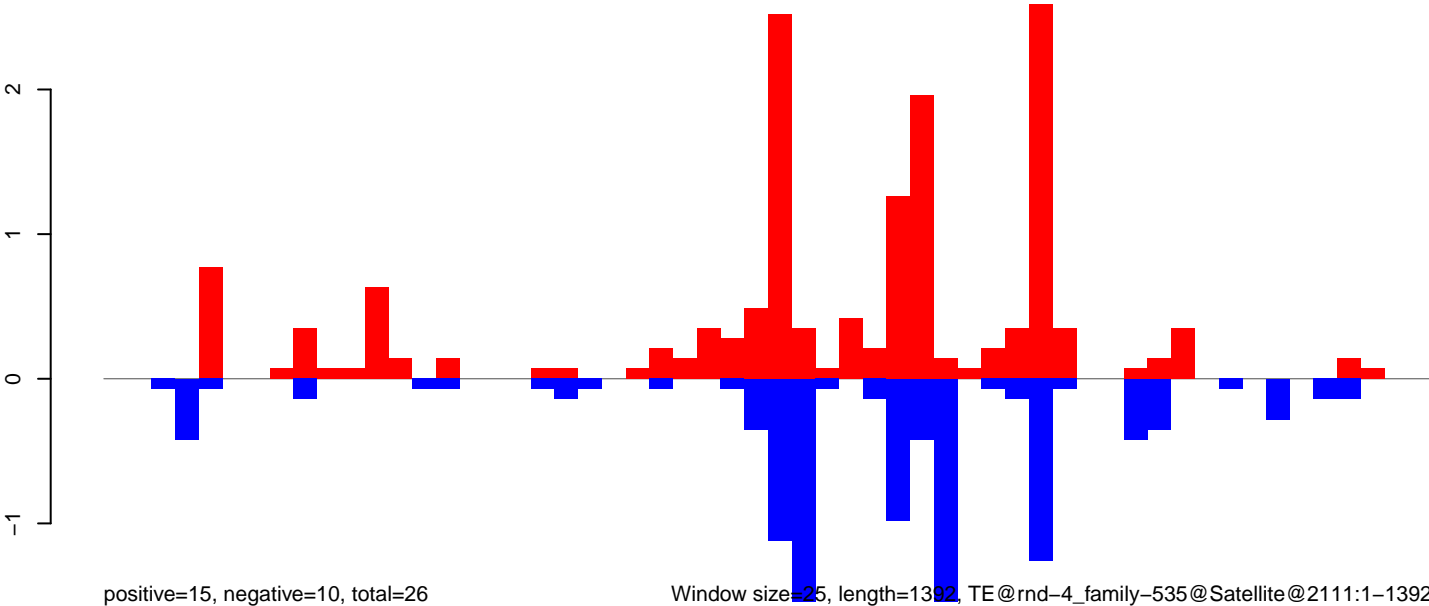
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



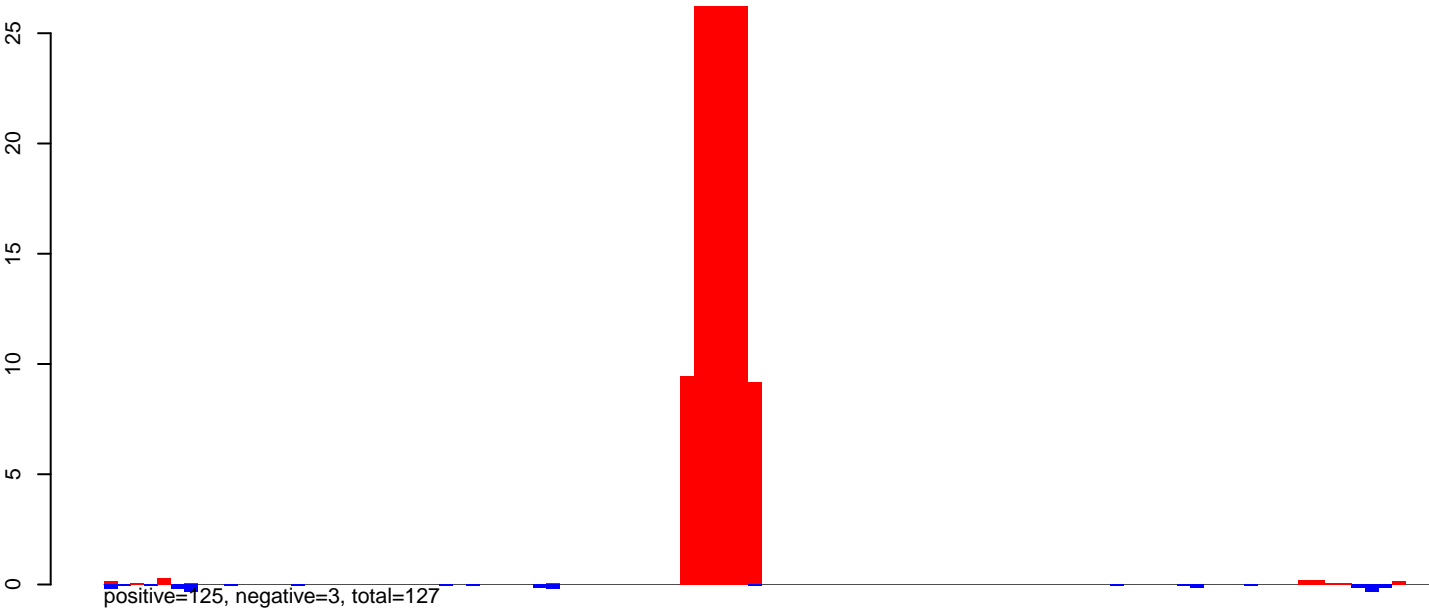
CuQuin_Mcarc_NL.rep



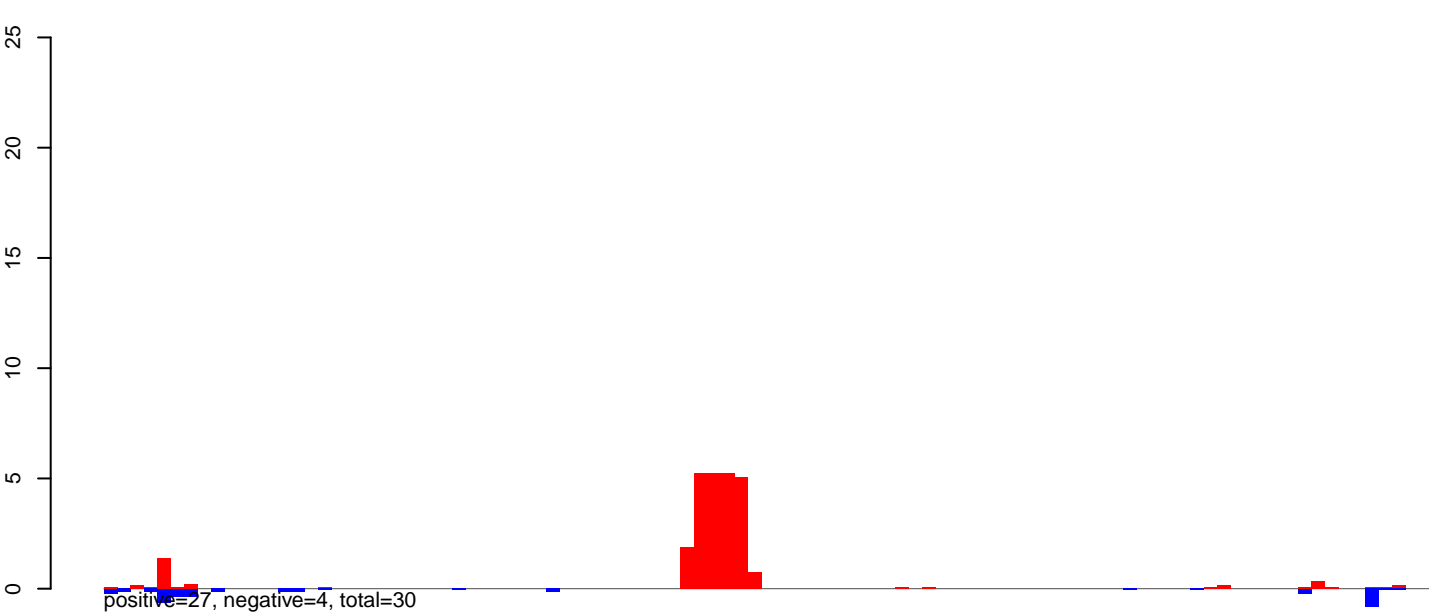
Window size=25, length=1392, TE@rnd-4_family-535@Satellite@2111:1-1392

0 200 400 600 800 1000 1200 1400

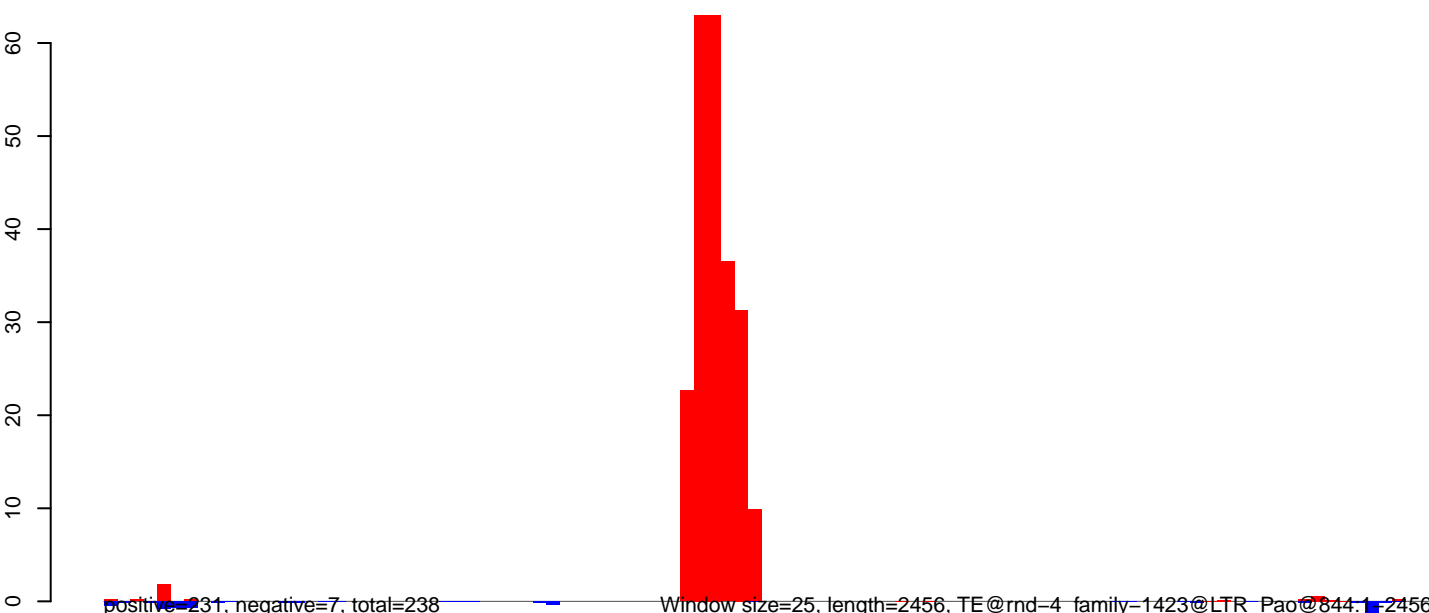
CuQuin_Mcarc_NL.18_23.rep



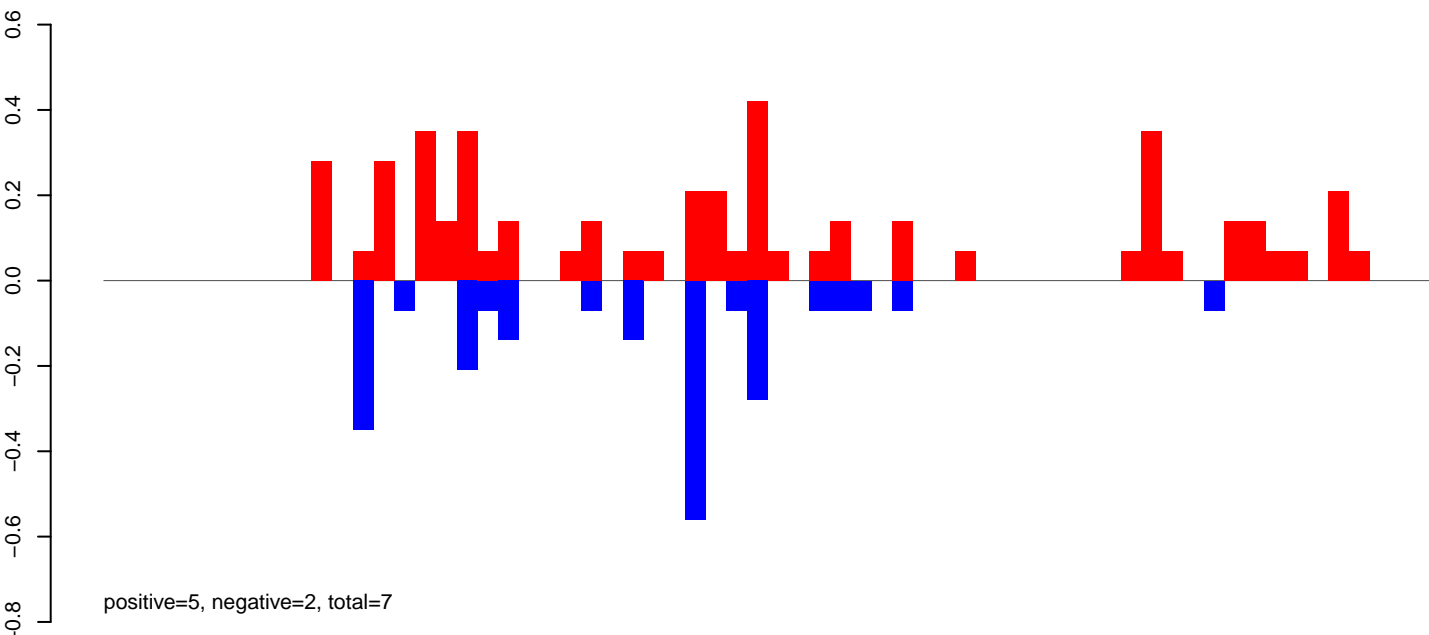
CuQuin_Mcarc_NL.24_35.rep



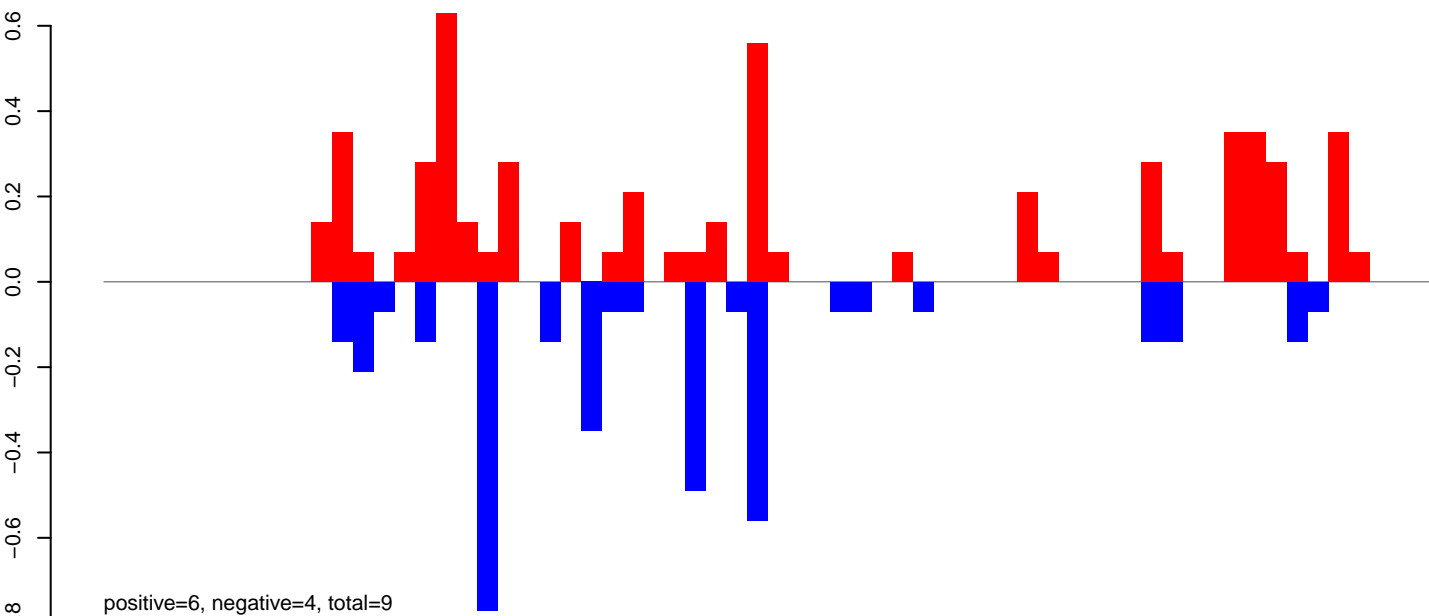
CuQuin_Mcarc_NL.rep



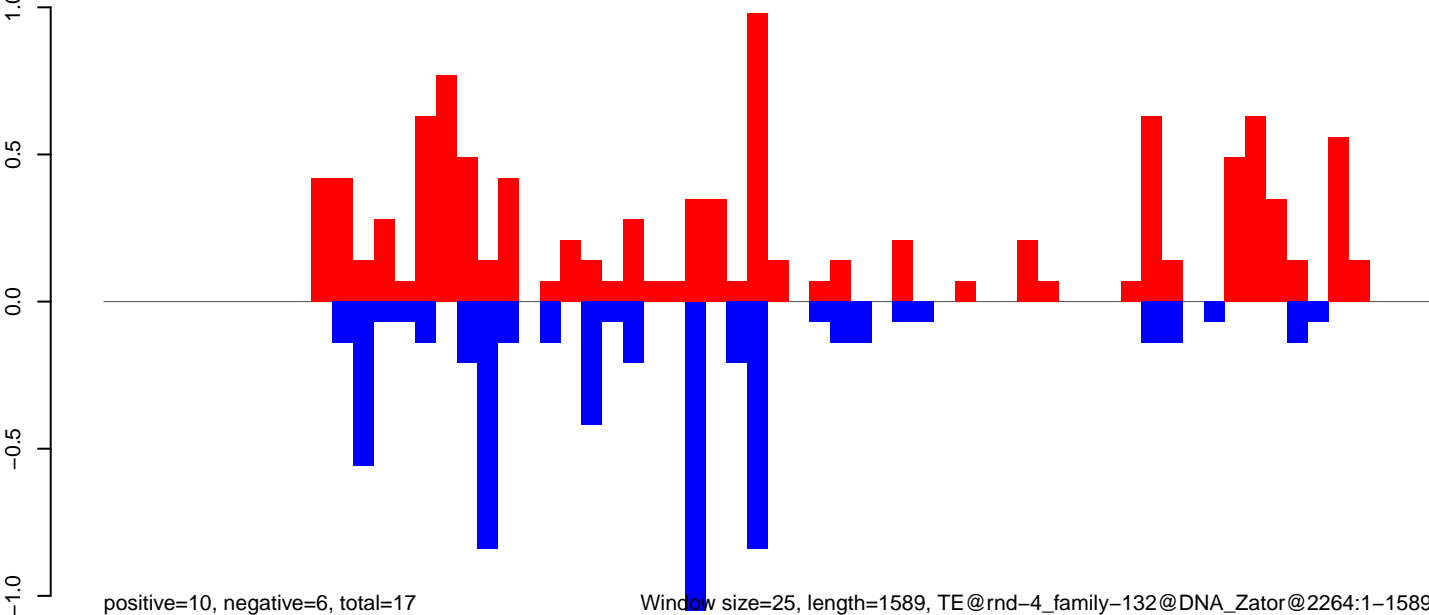
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



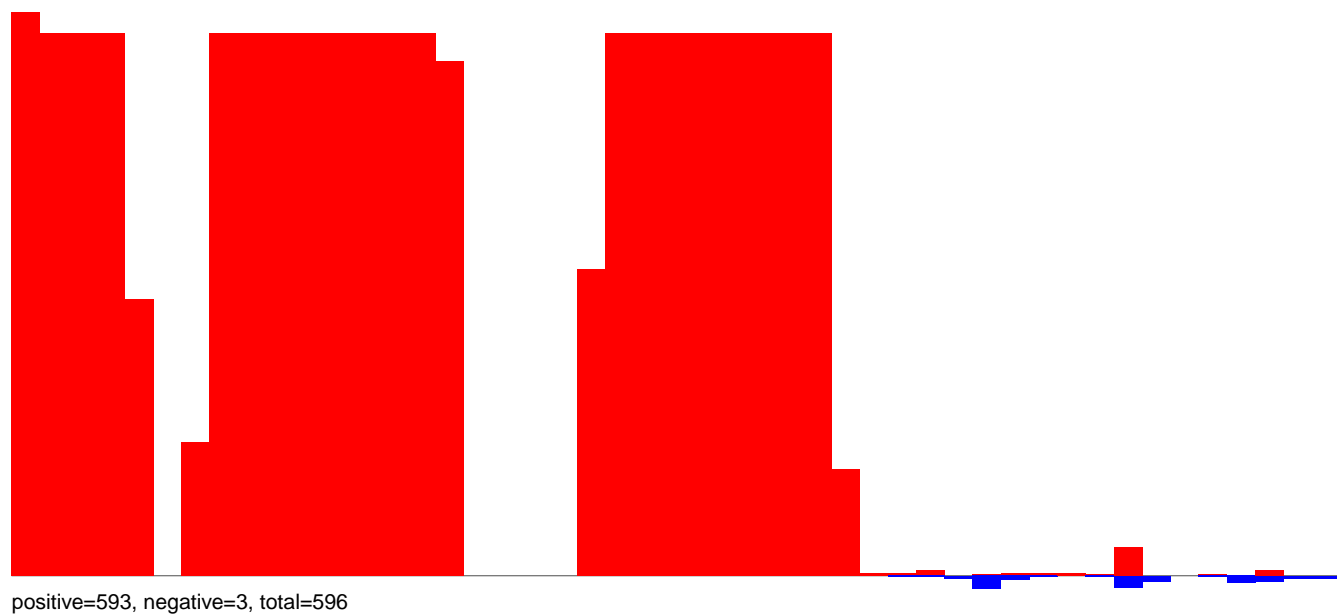
CuQuin_Mcarc_NL.rep



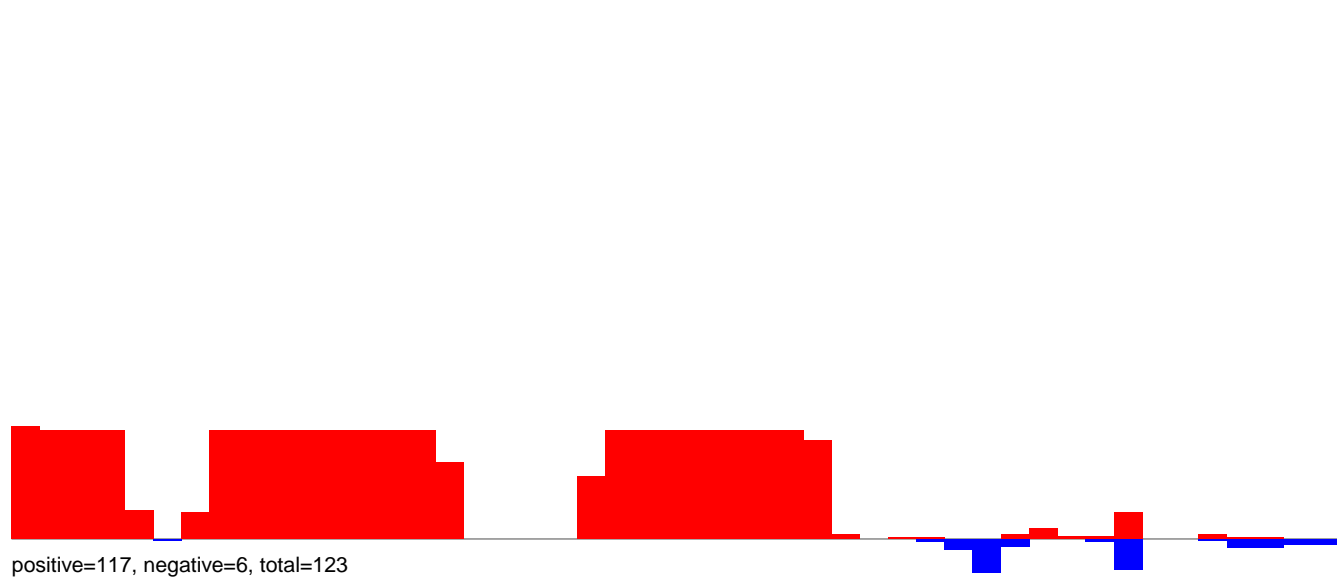
Window size=25, length=1589, TE@rnd-4_family-132@DNA_Zator@2264:1-1589

0 500 1000 1500

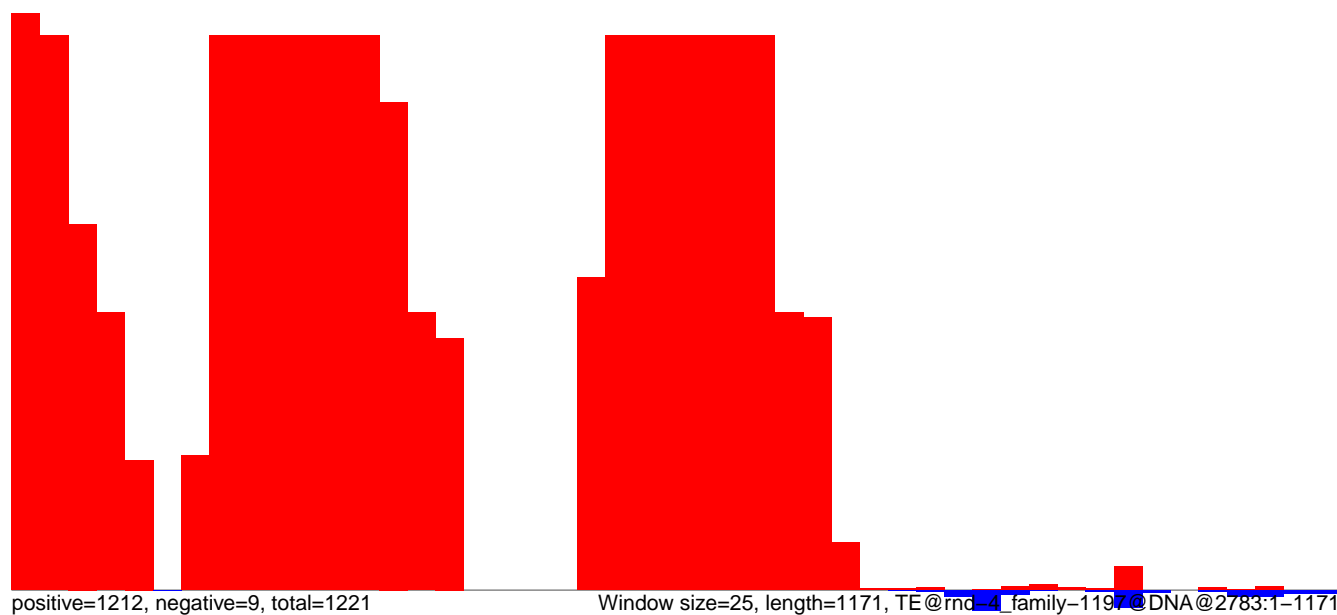
CuQuin_Mcarc_NL.18_23.rep



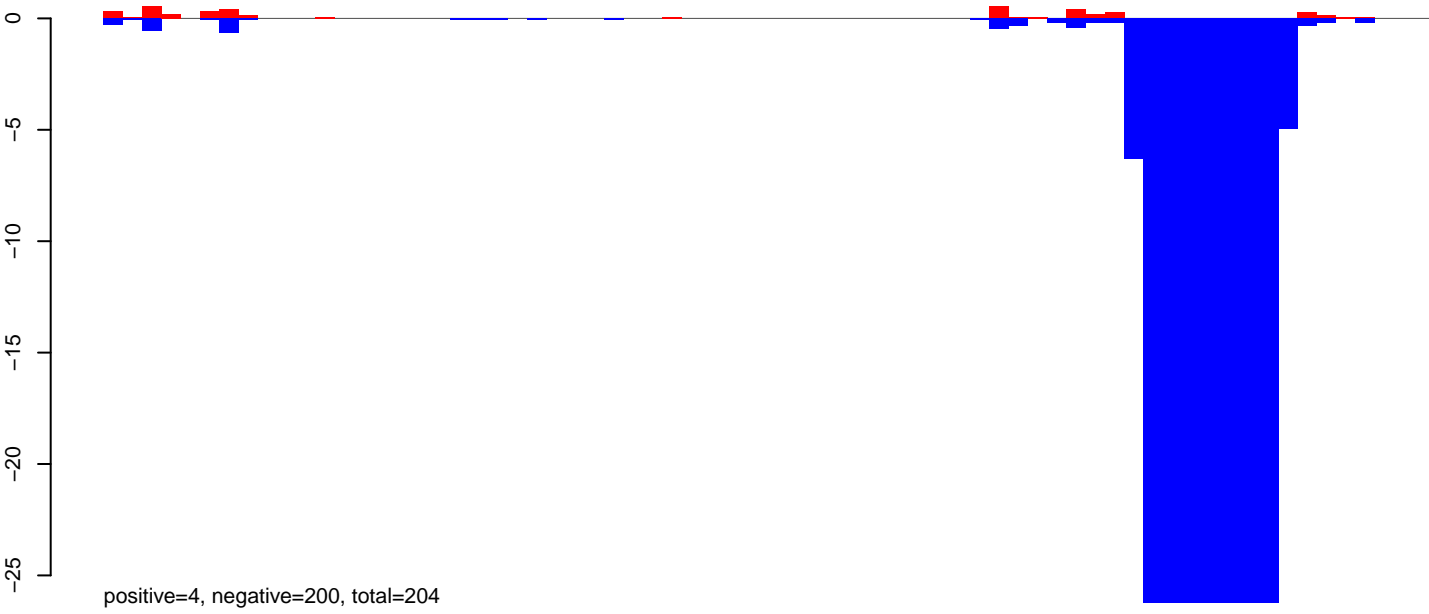
CuQuin_Mcarc_NL.24_35.rep



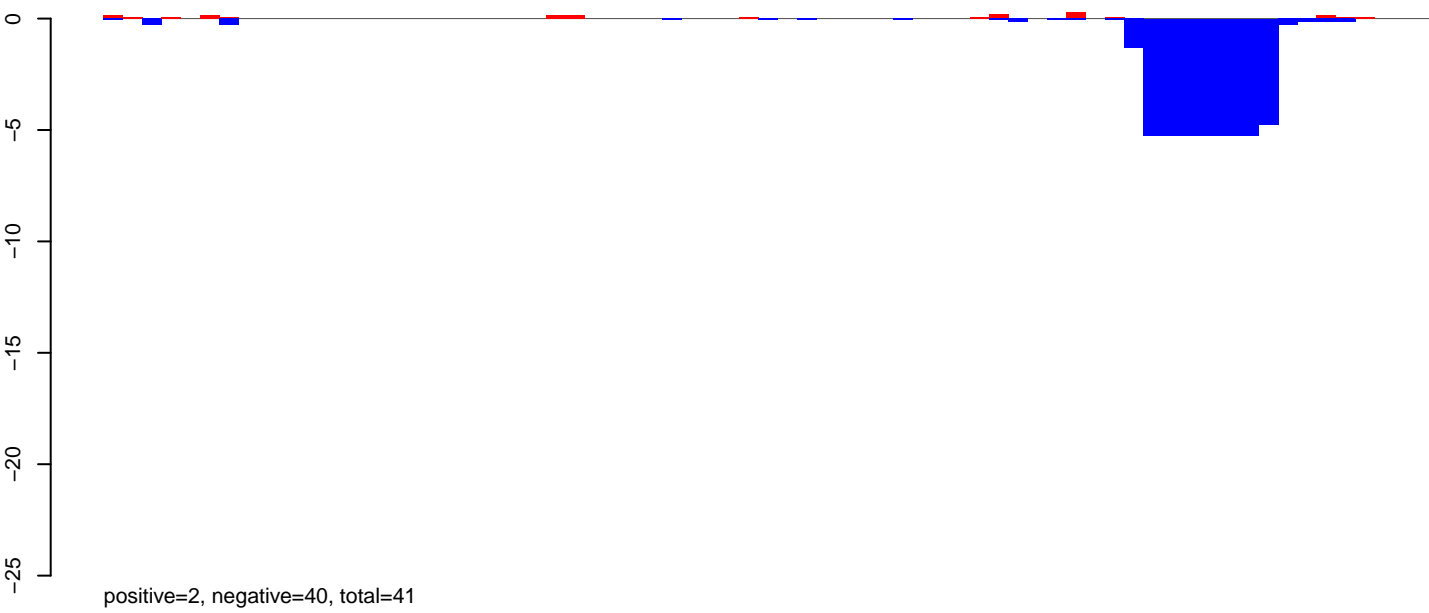
CuQuin_Mcarc_NL.rep



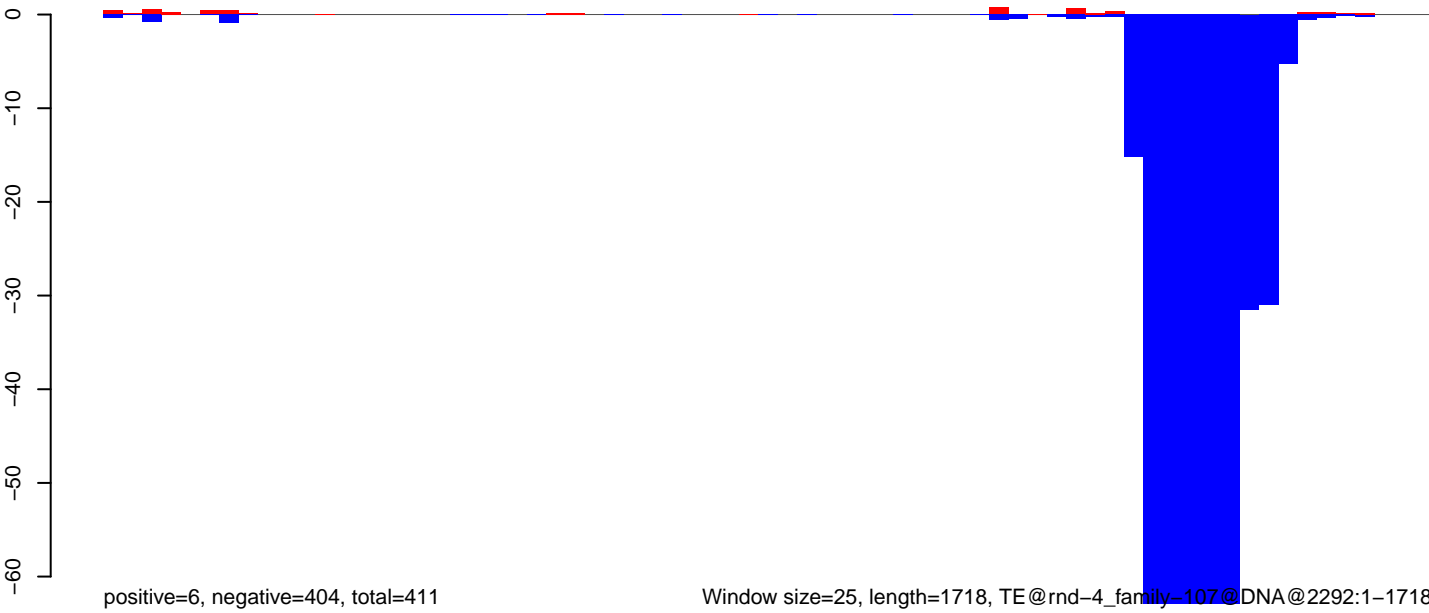
CuQuin_Mcarc_NL.18_23.rep



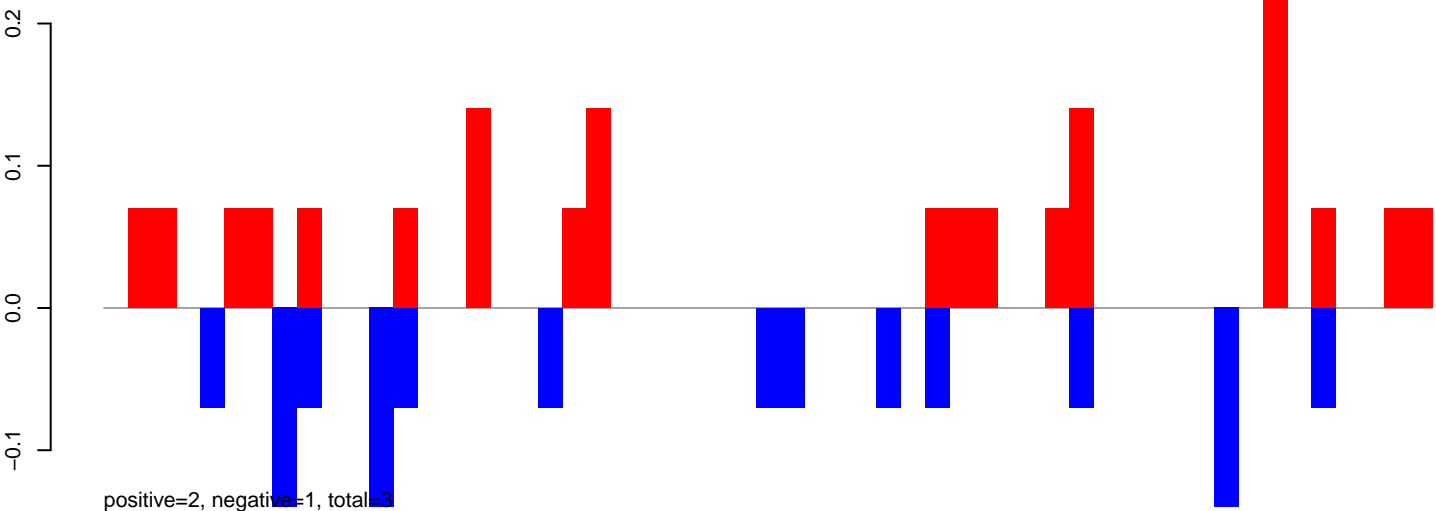
CuQuin_Mcarc_NL.24_35.rep



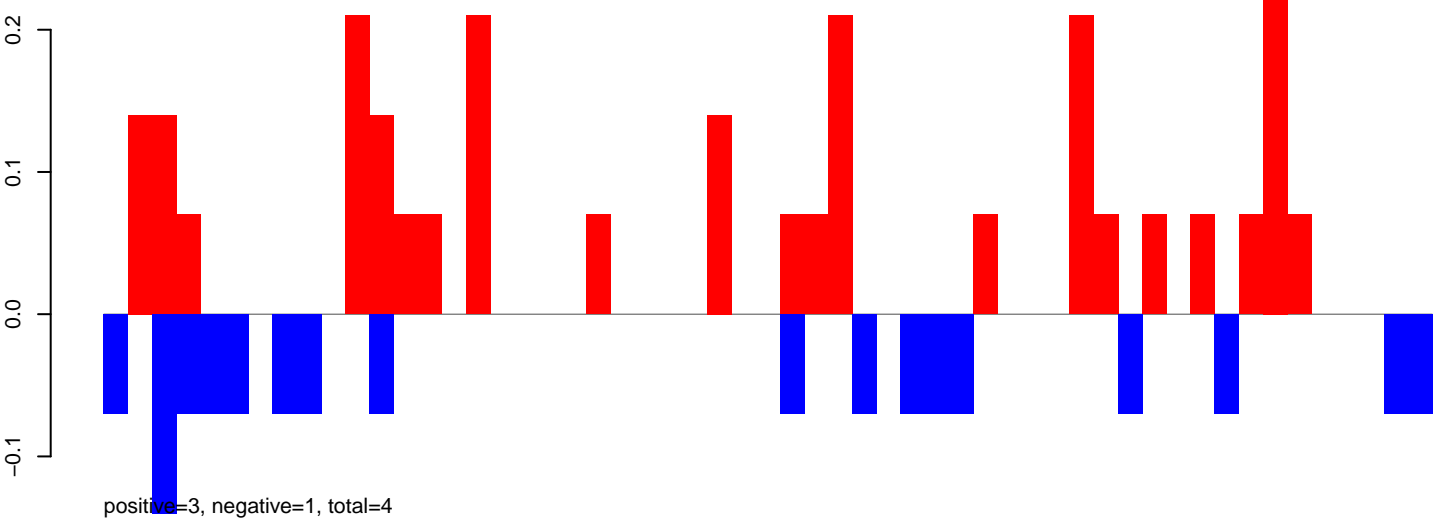
CuQuin_Mcarc_NL.rep



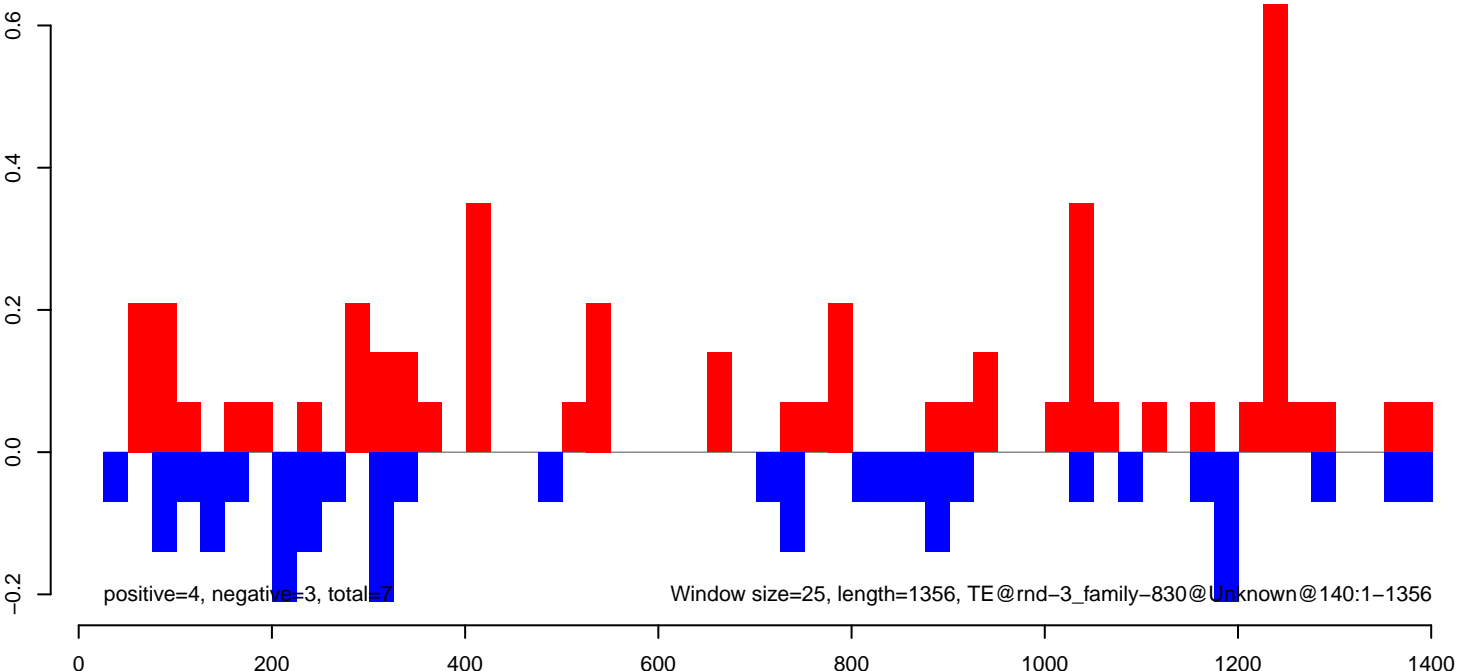
CuQuin_Mcarc_NL.18_23.rep



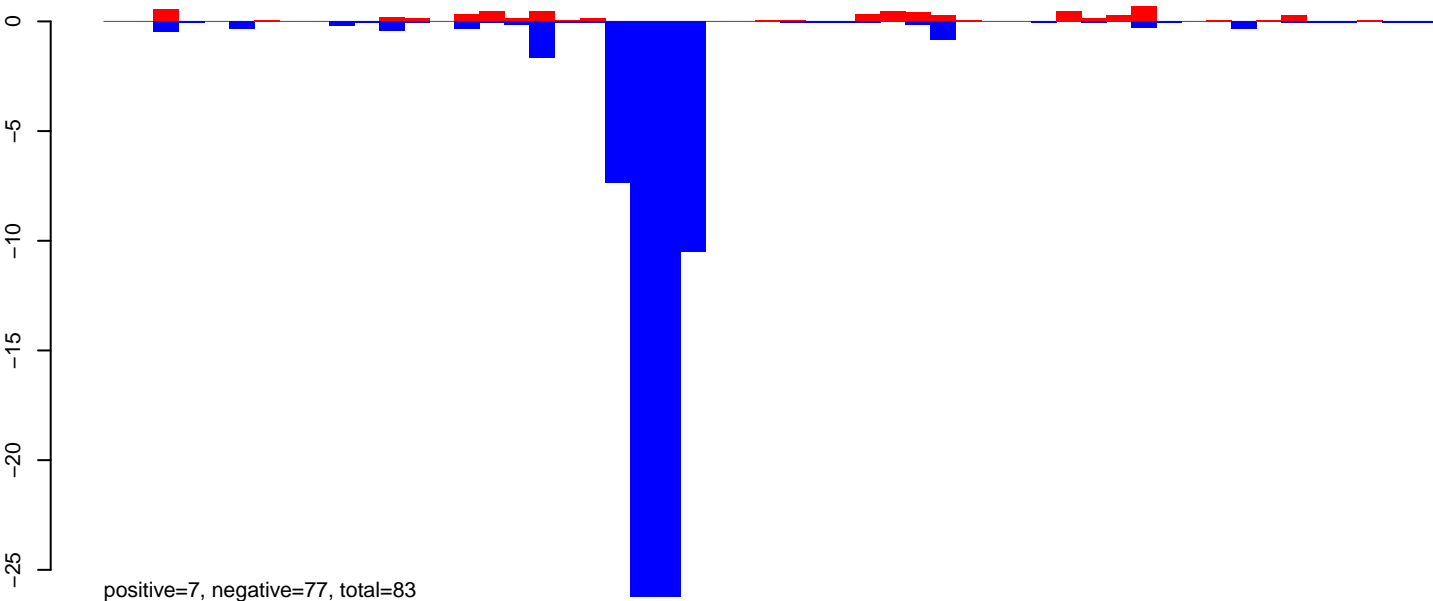
CuQuin_Mcarc_NL.24_35.rep



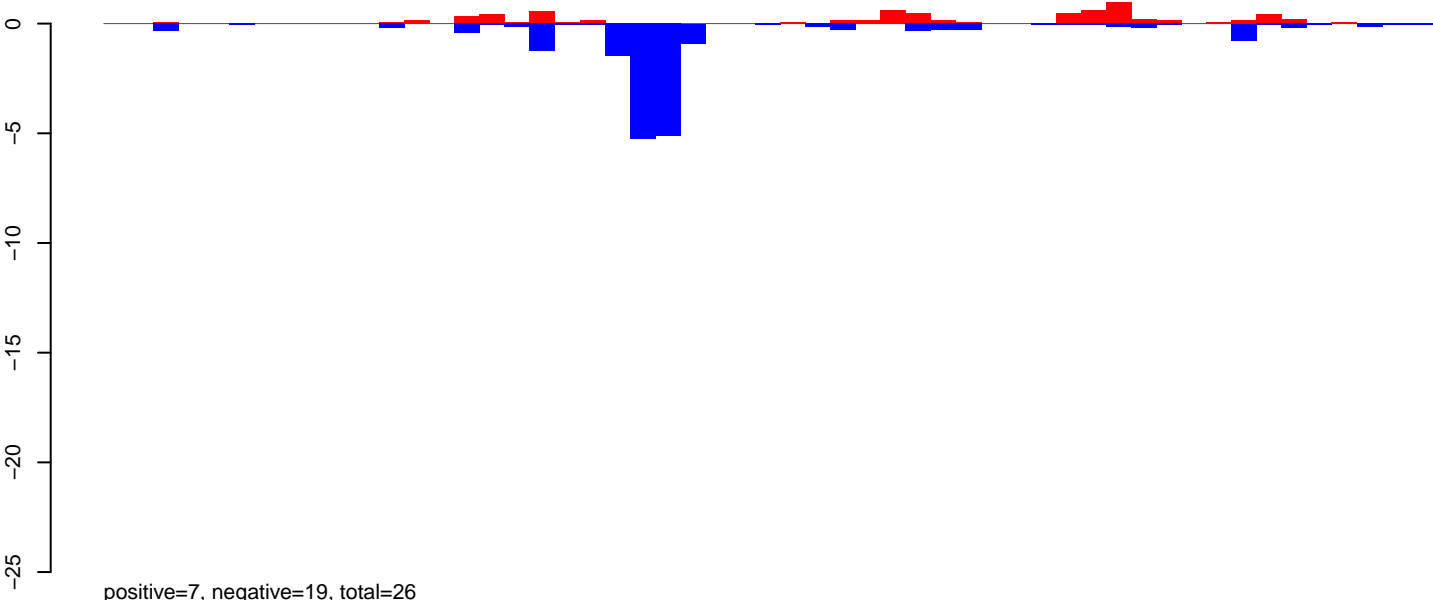
CuQuin_Mcarc_NL.rep



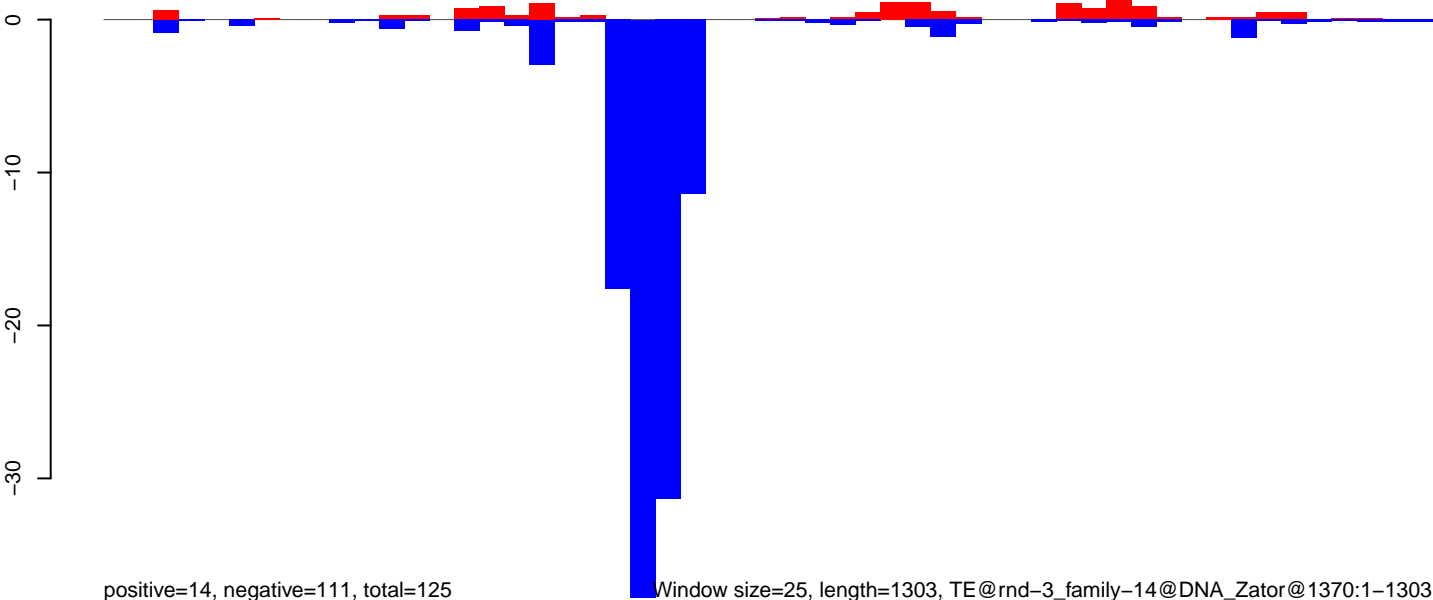
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



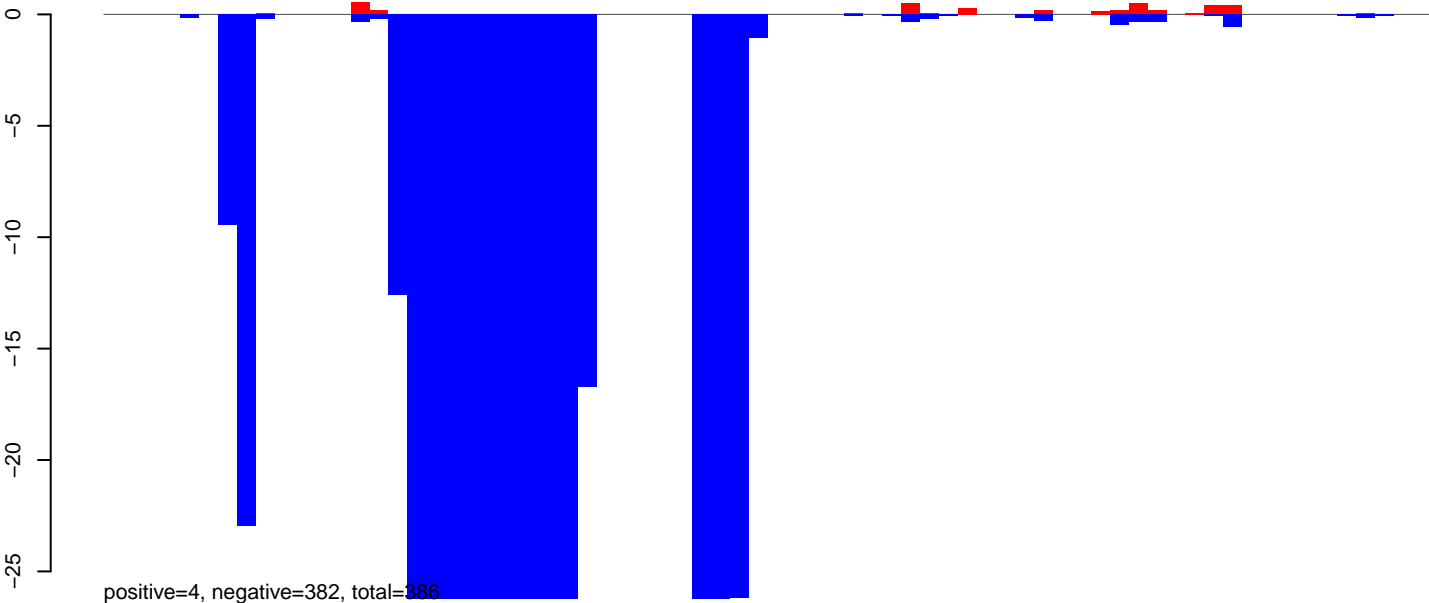
CuQuin_Mcarc_NL.rep



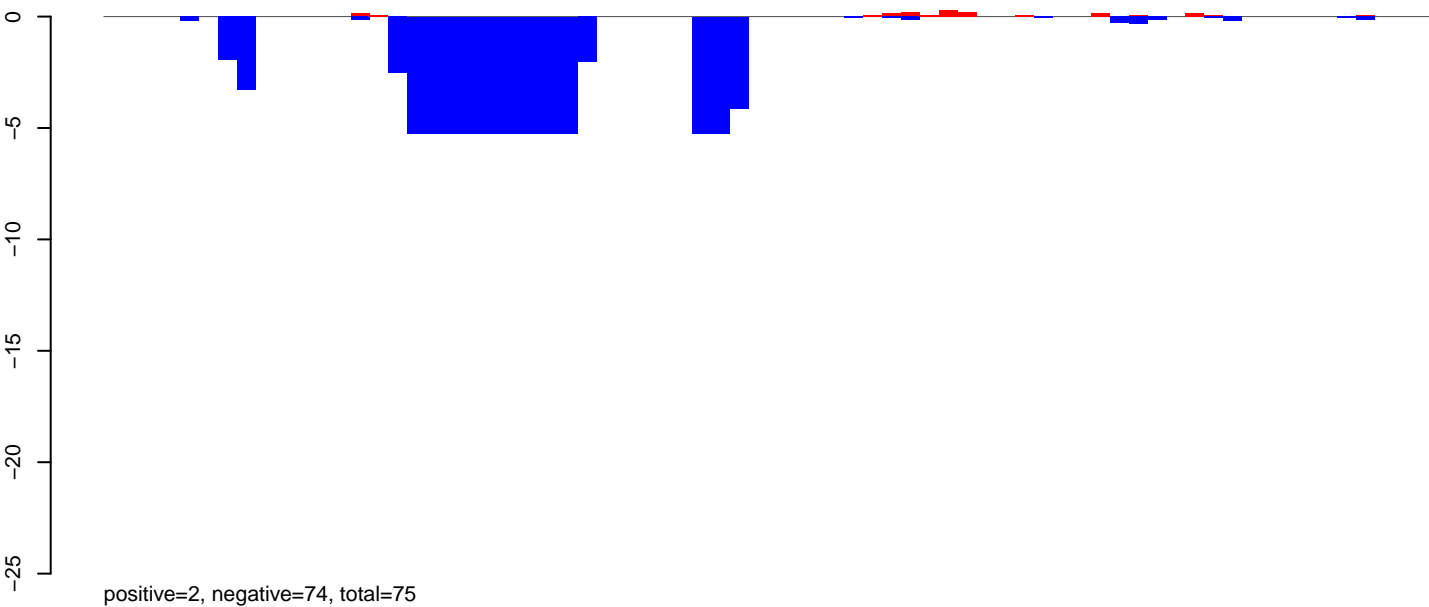
Window size=25, length=1303, TE@rnd-3_family-14@DNA_Zator@1370:1-1303

0 200 400 600 800 1000 1200 1400

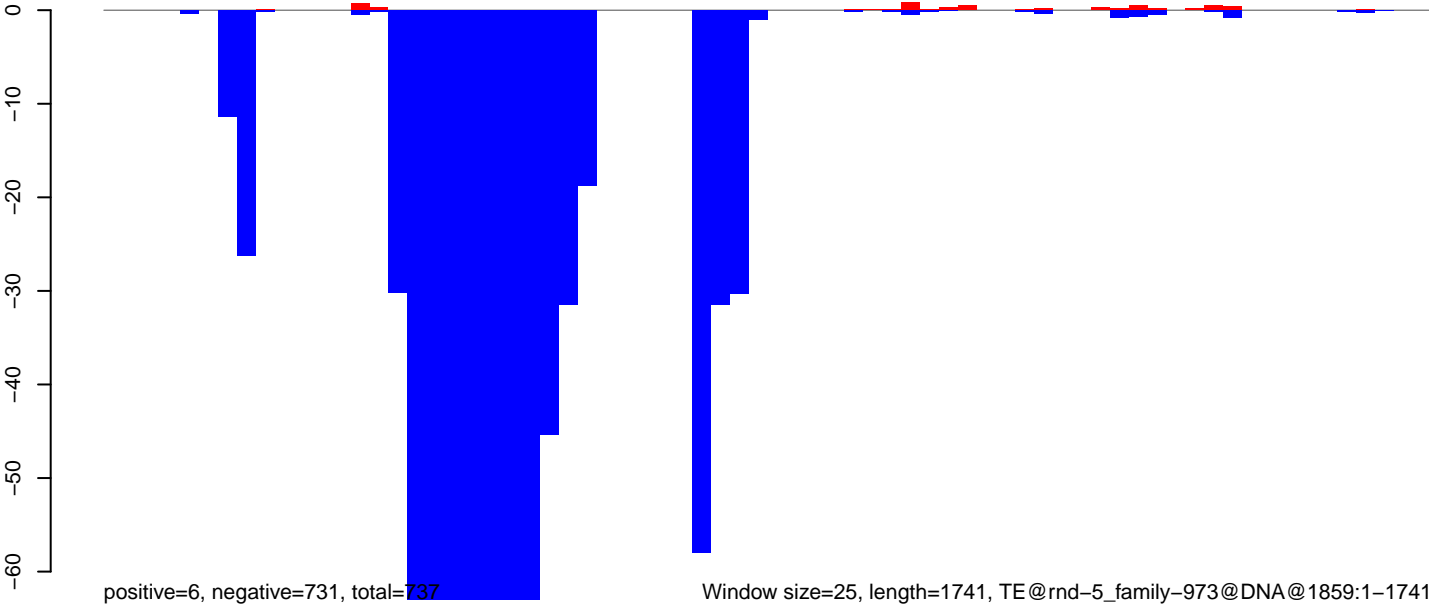
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



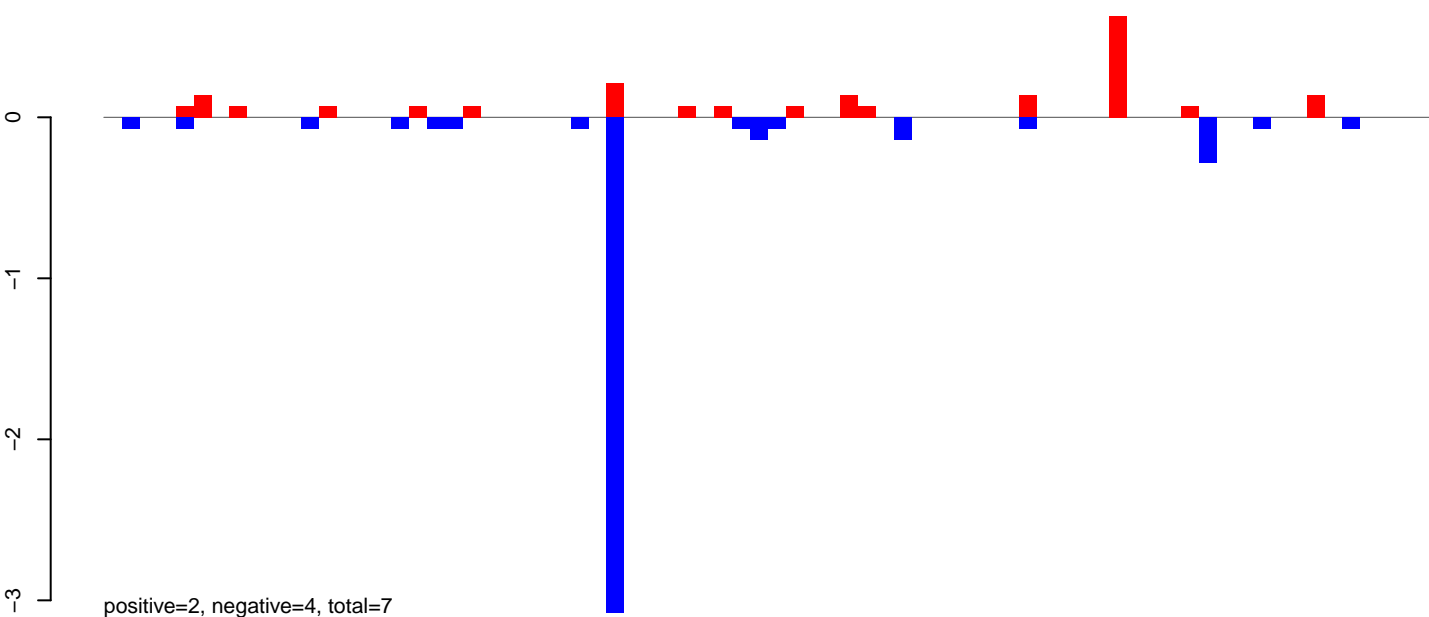
CuQuin_Mcarc_NL.rep



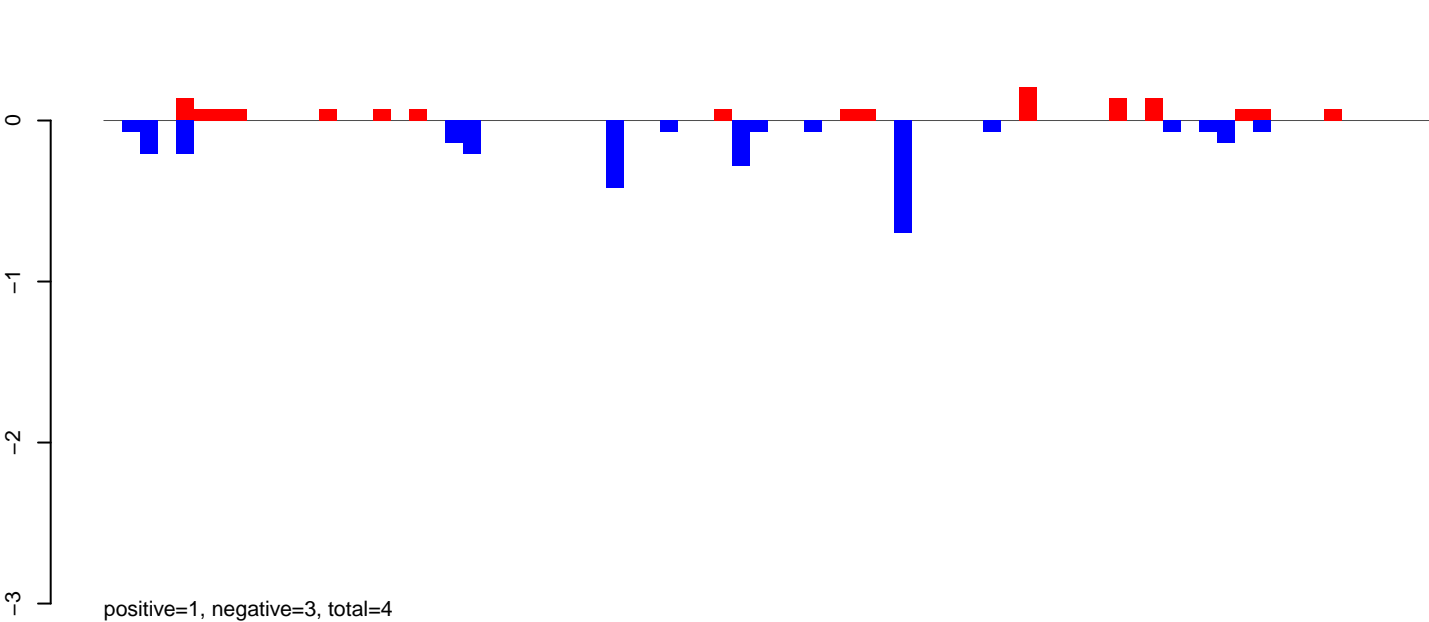
Window size=25, length=1741, TE@rnd-5_family-973@DNA@1859:1-1741

0 500 1000 1500

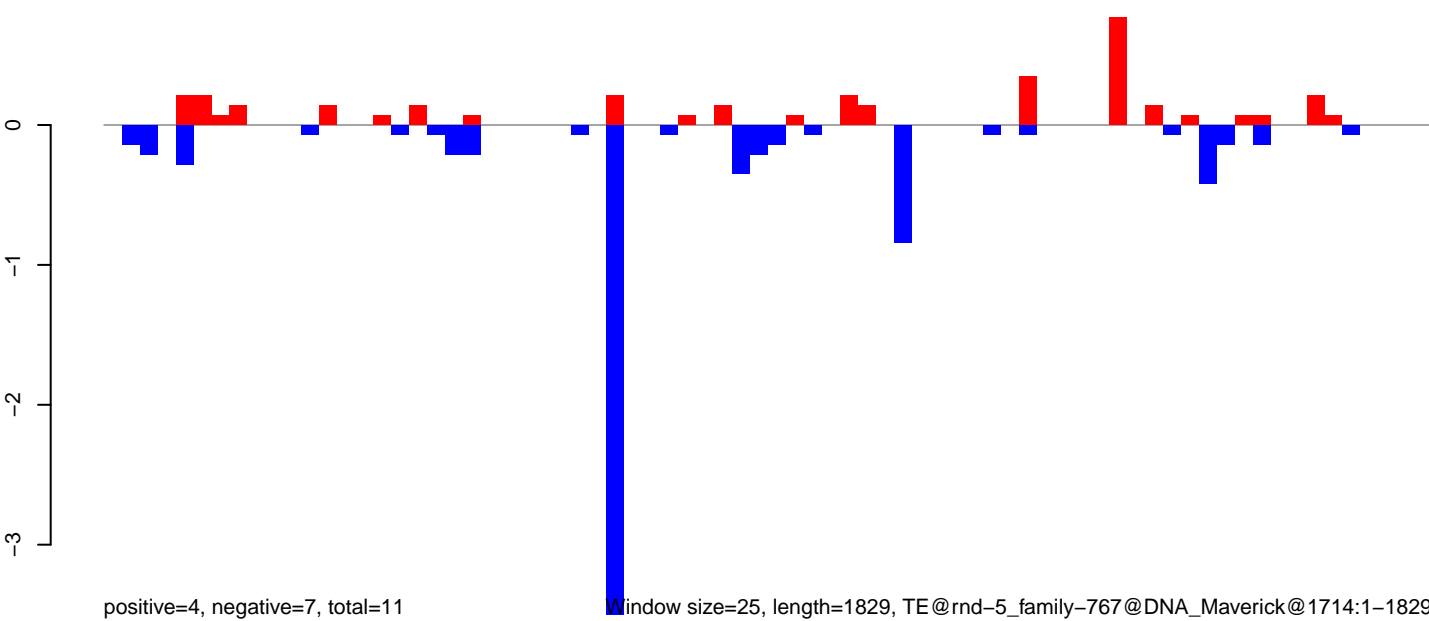
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



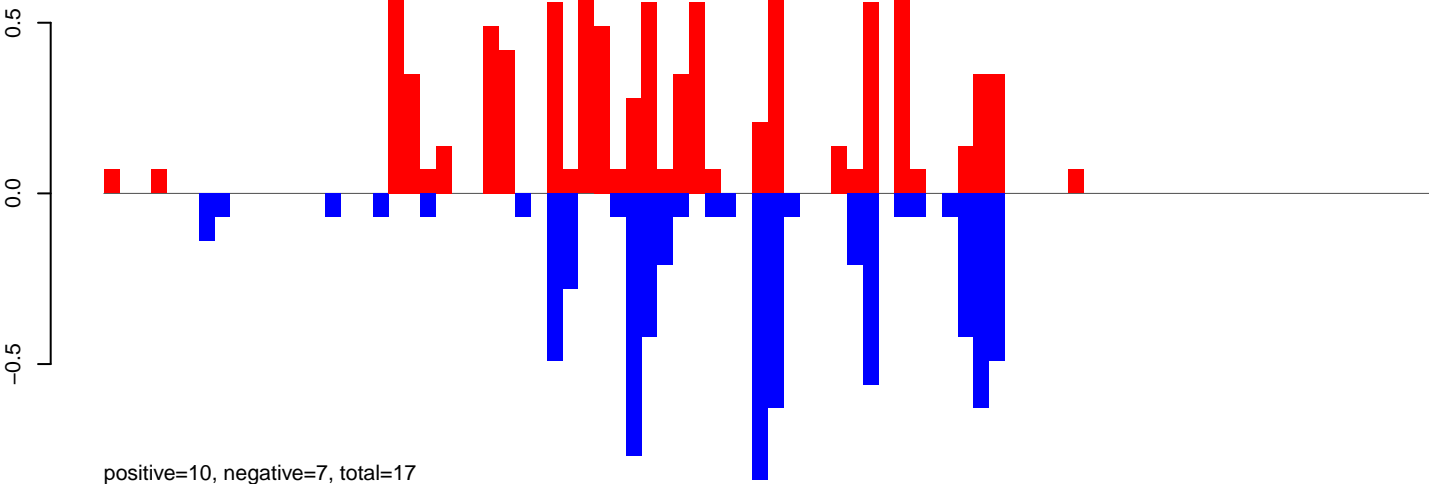
CuQuin_Mcarc_NL.rep



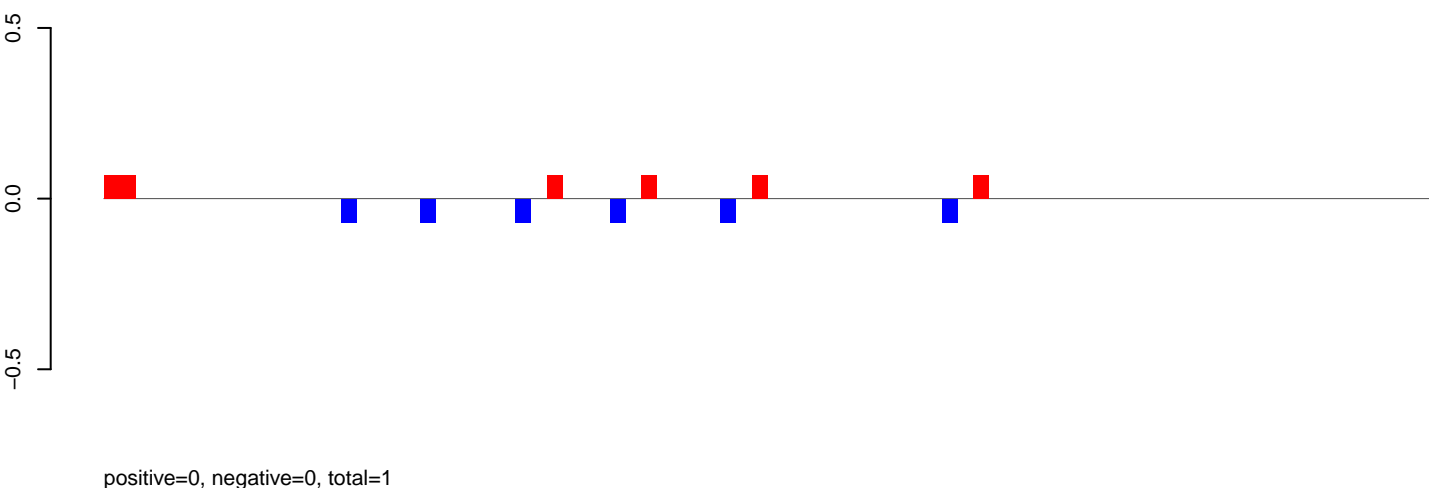
Window size=25, length=1829, TE@rnd-5_family-767@DNA_Maverick@1714:1-1829

0 500 1000 1500

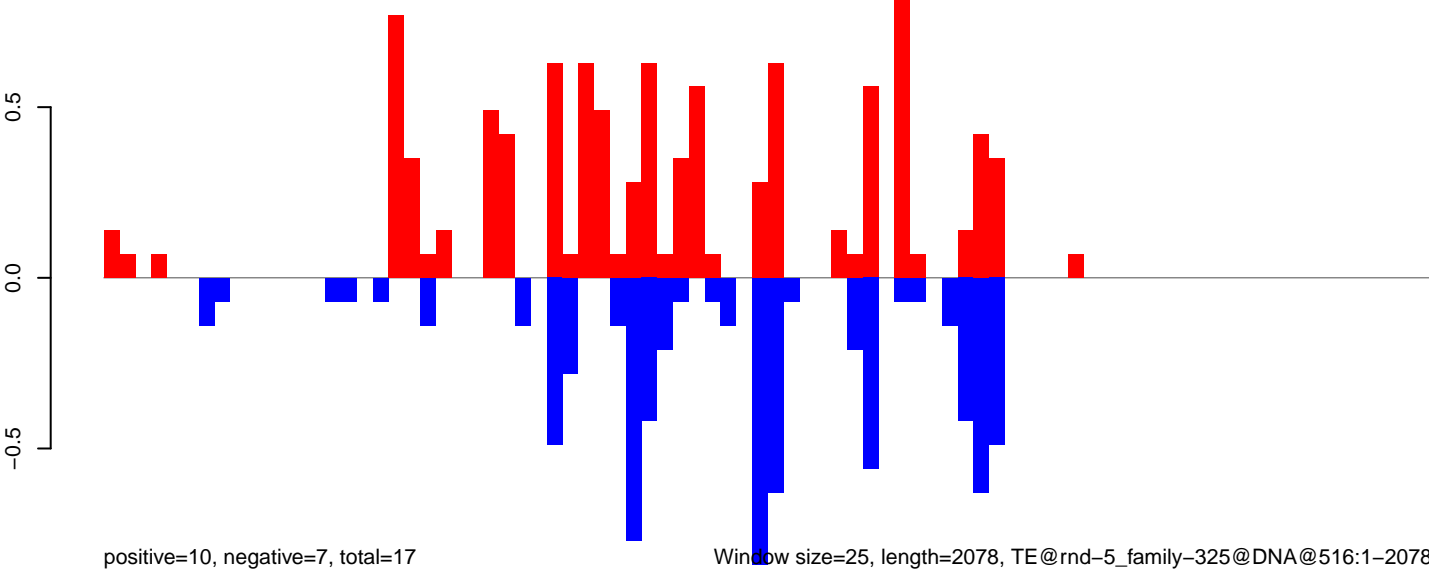
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



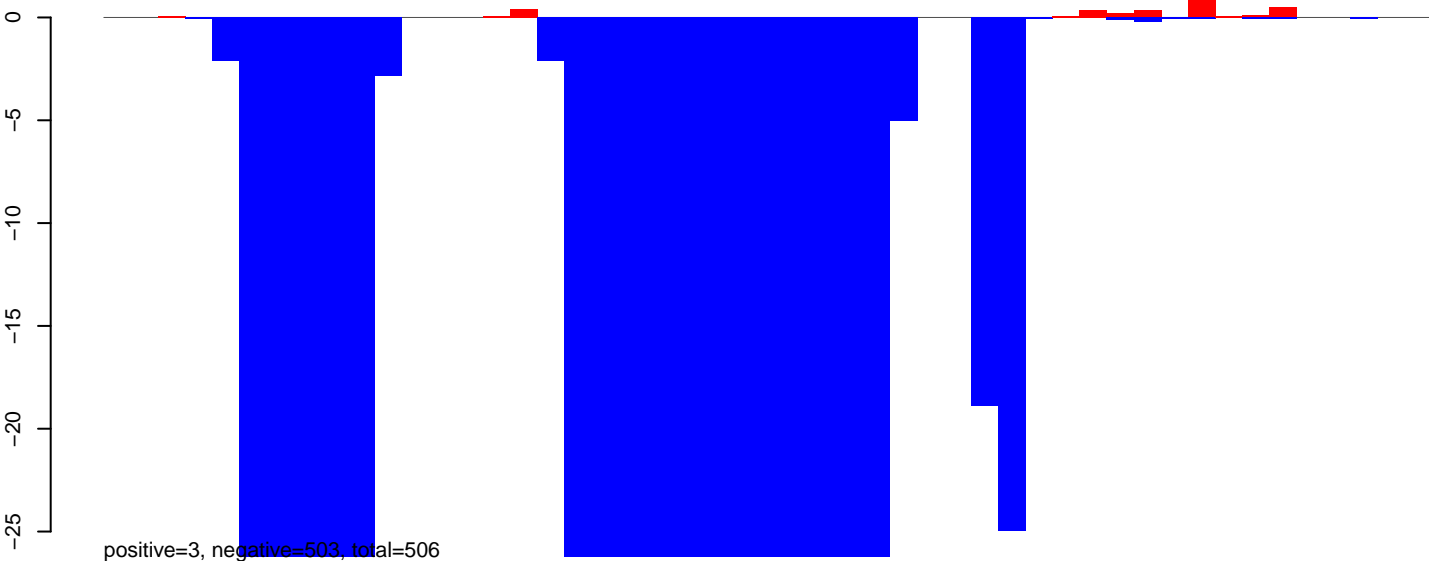
CuQuin_Mcarc_NL.rep



Window size=25, length=2078, TE@rnd-5_family-325@DNA@516:1-2078

0 500 1000 1500 2000

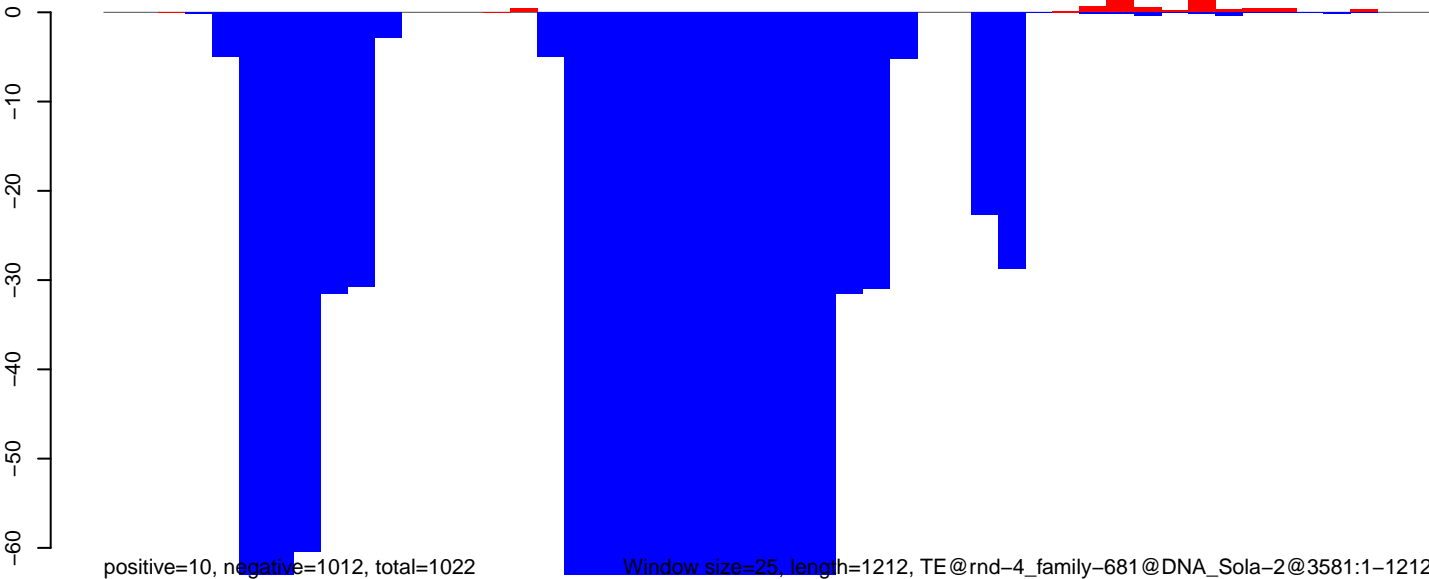
CuQuin_Mcarc_NL.18_23.rep



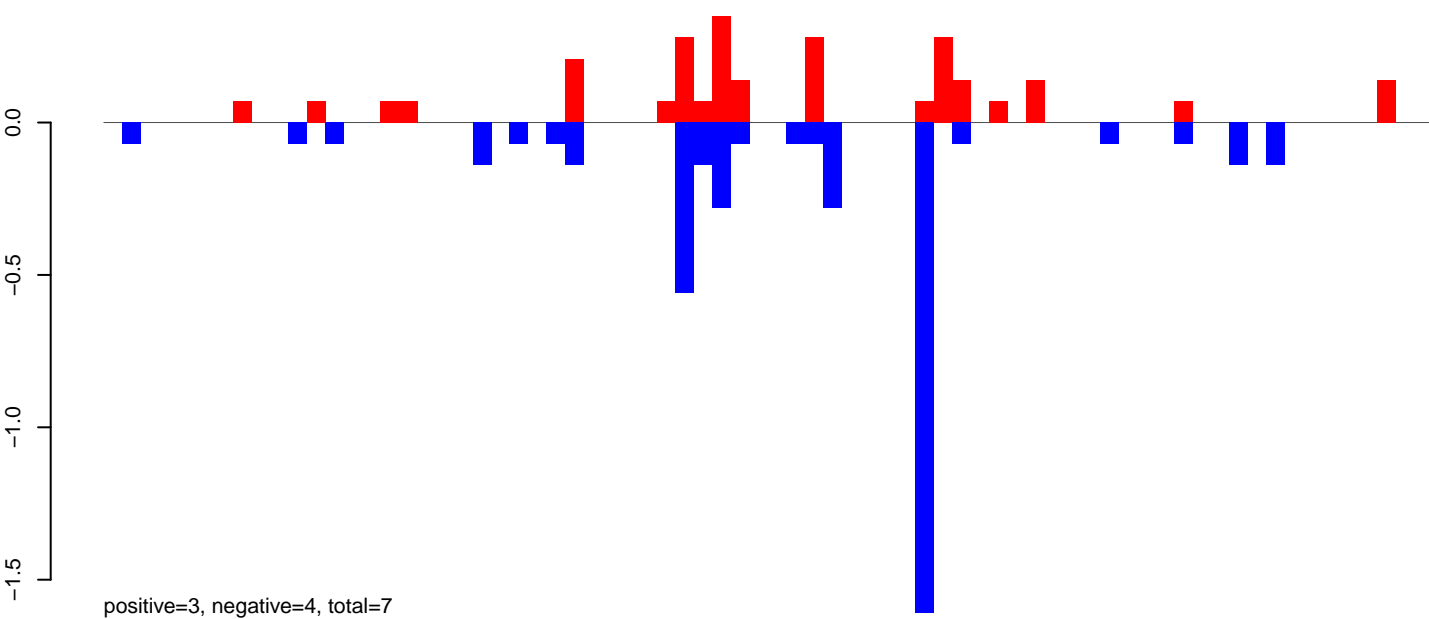
CuQuin_Mcarc_NL.24_35.rep



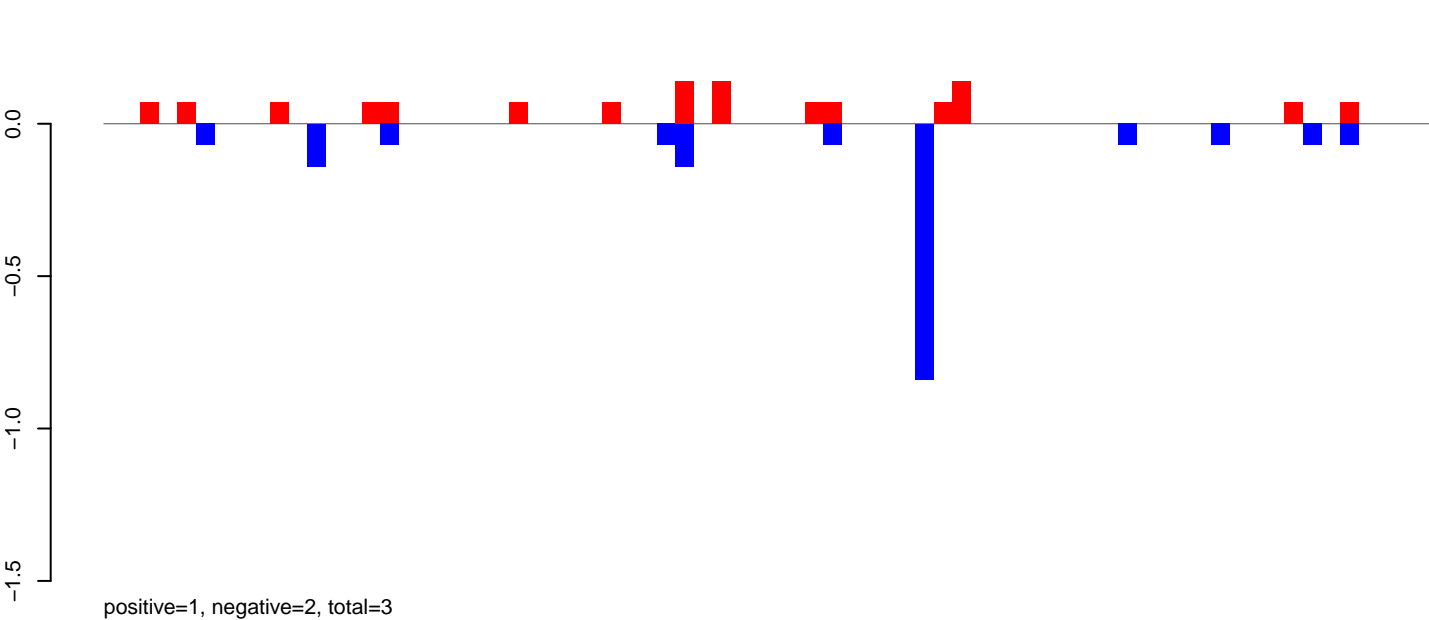
CuQuin_Mcarc_NL.rep



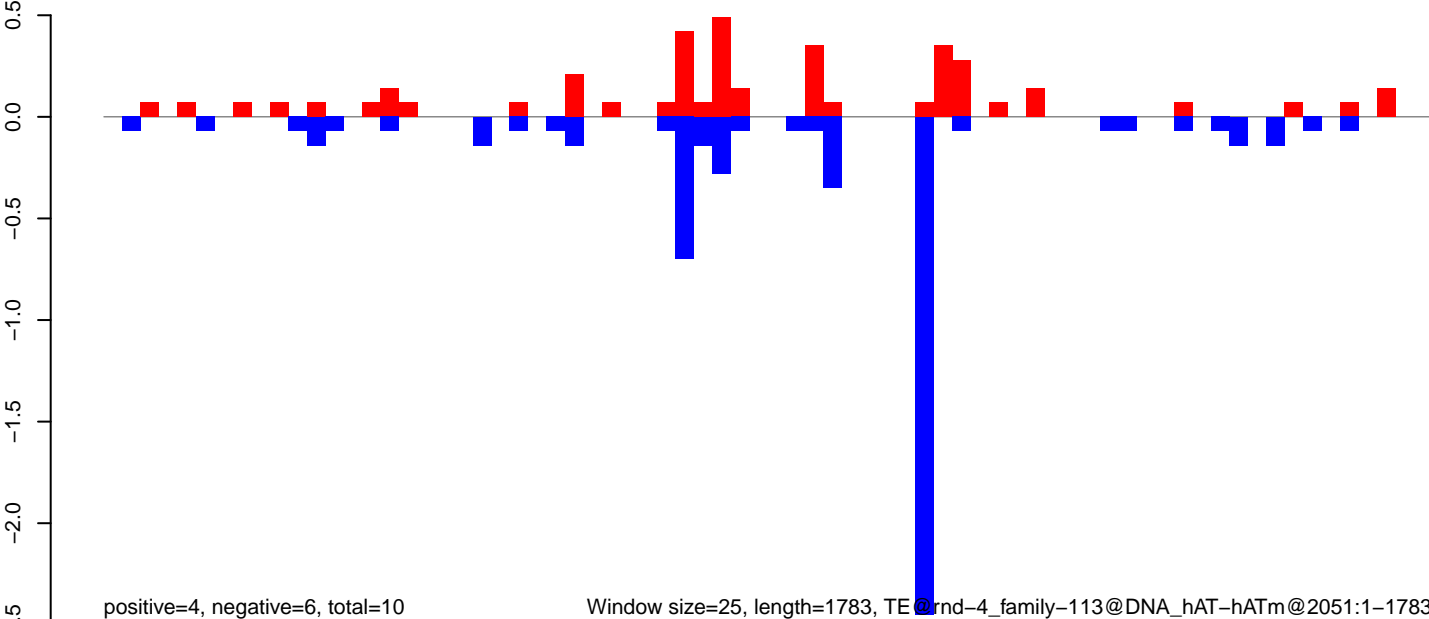
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

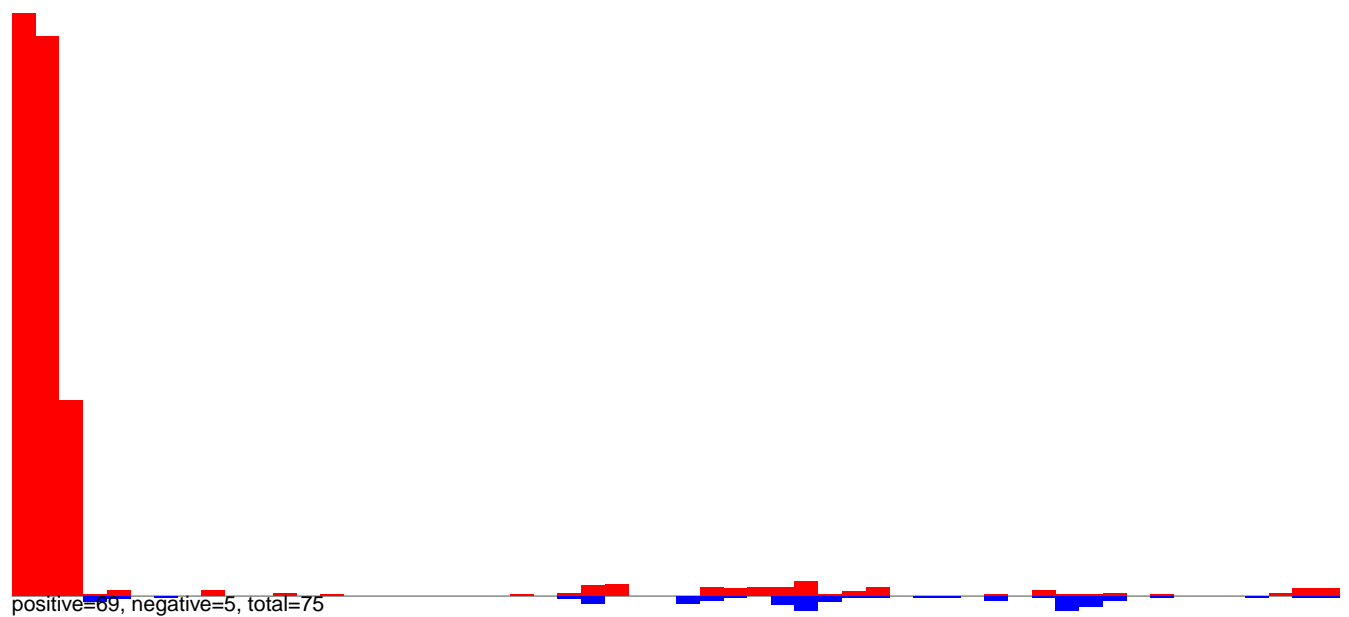


CuQuin_Mcarc_NL.rep

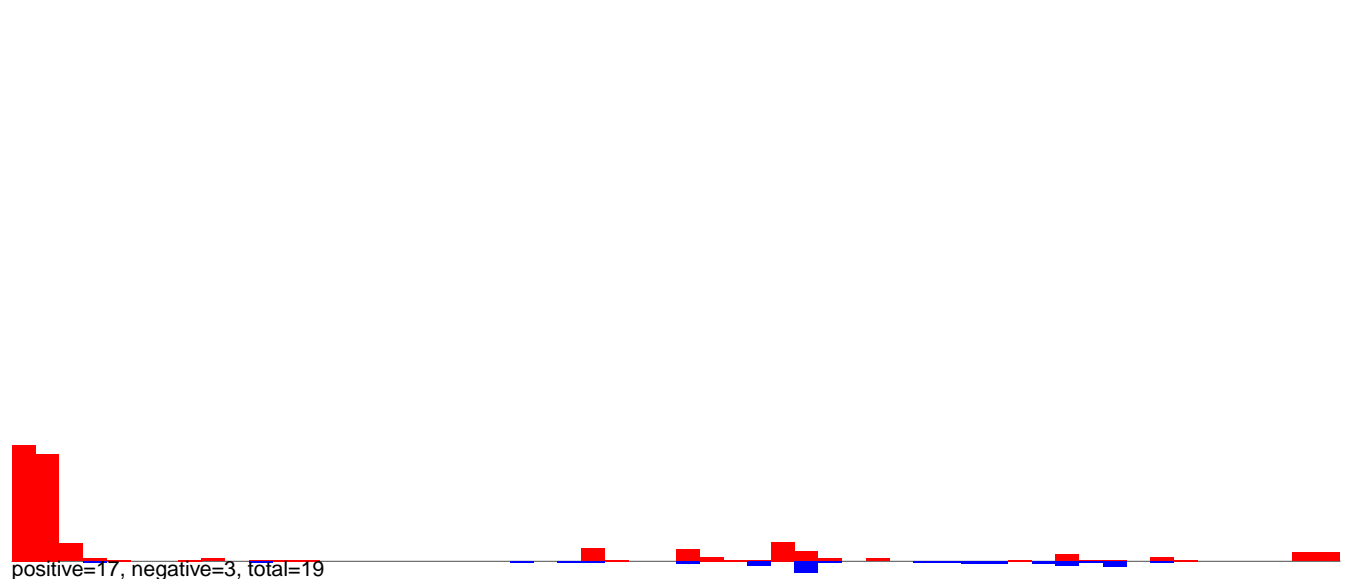


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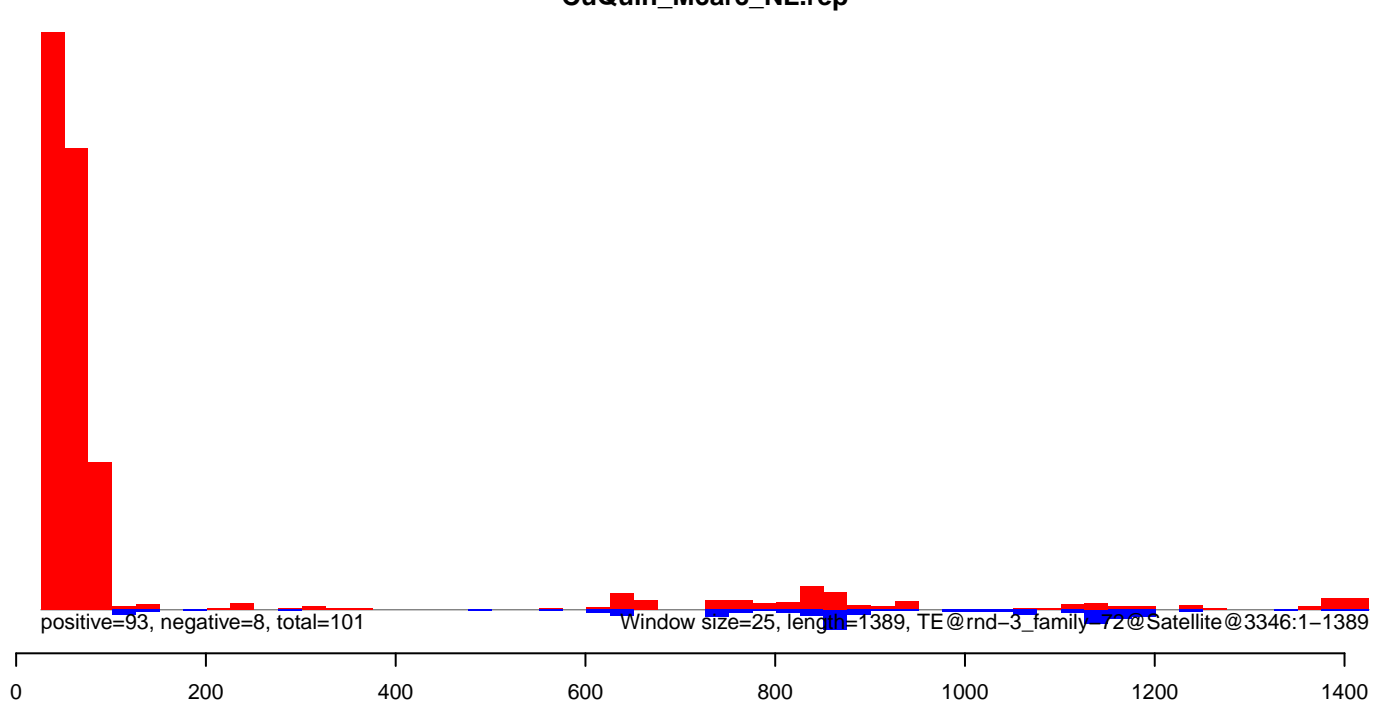
CuQuin_Mcarc_NL.18_23.rep



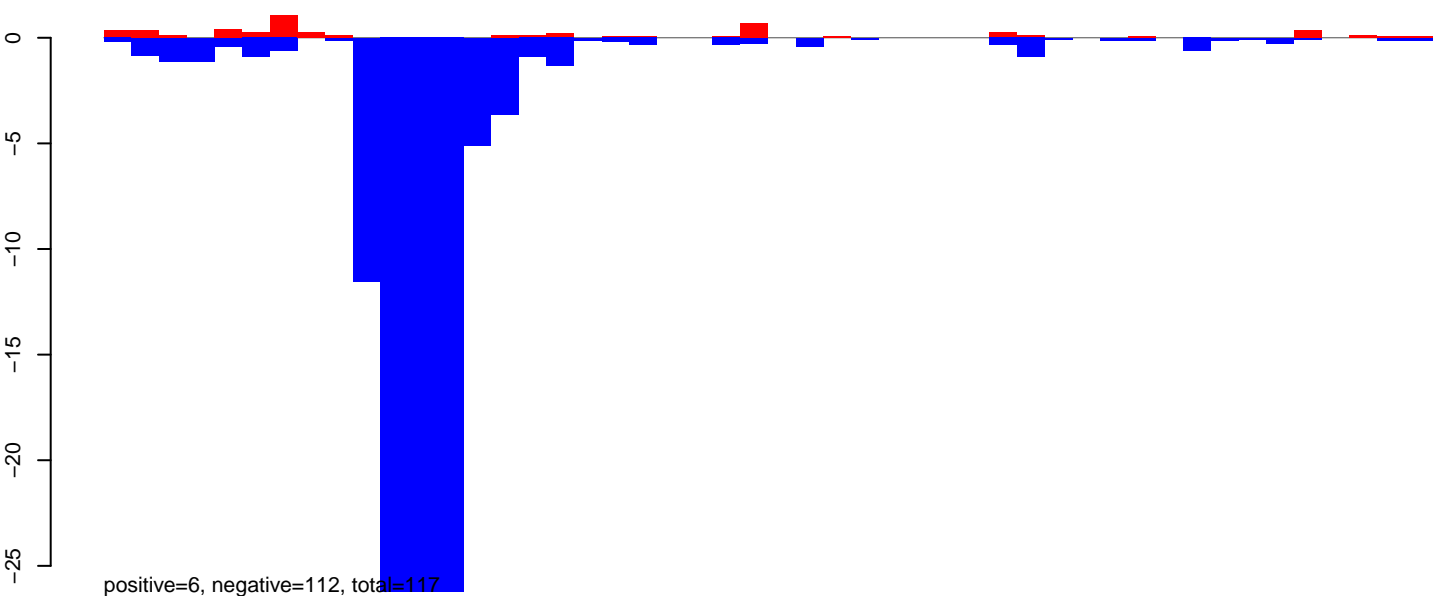
CuQuin_Mcarc_NL.24_35.rep



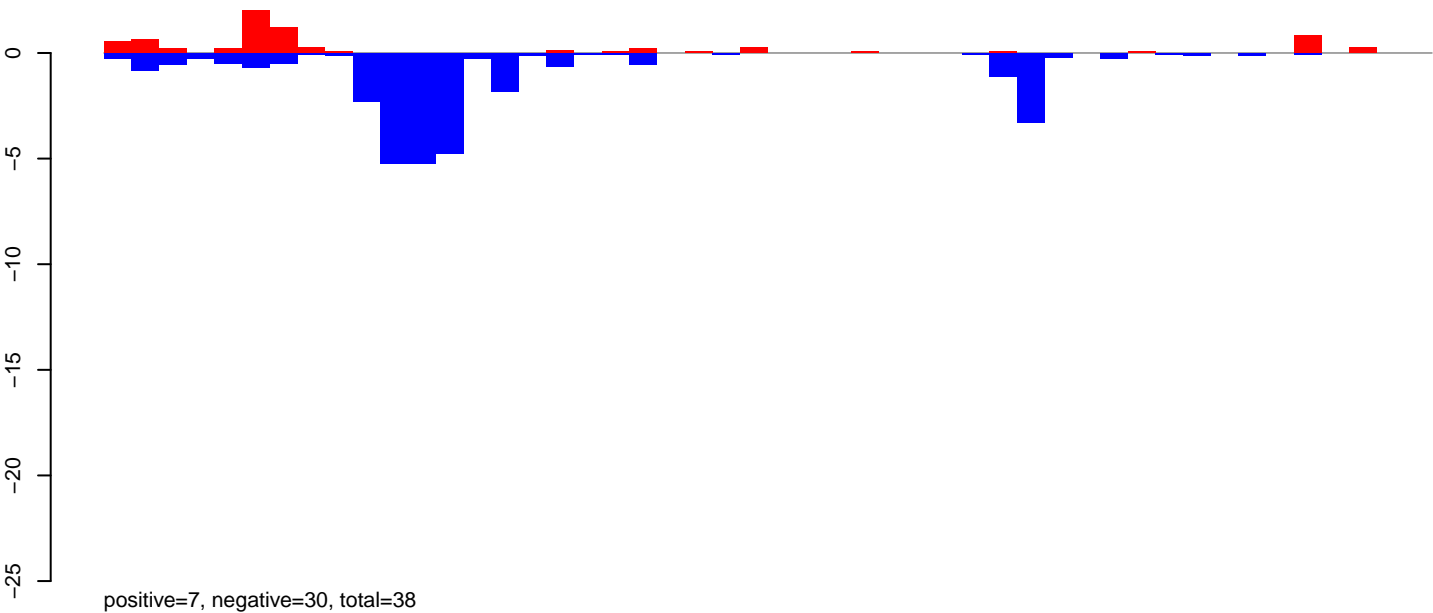
CuQuin_Mcarc_NL.rep



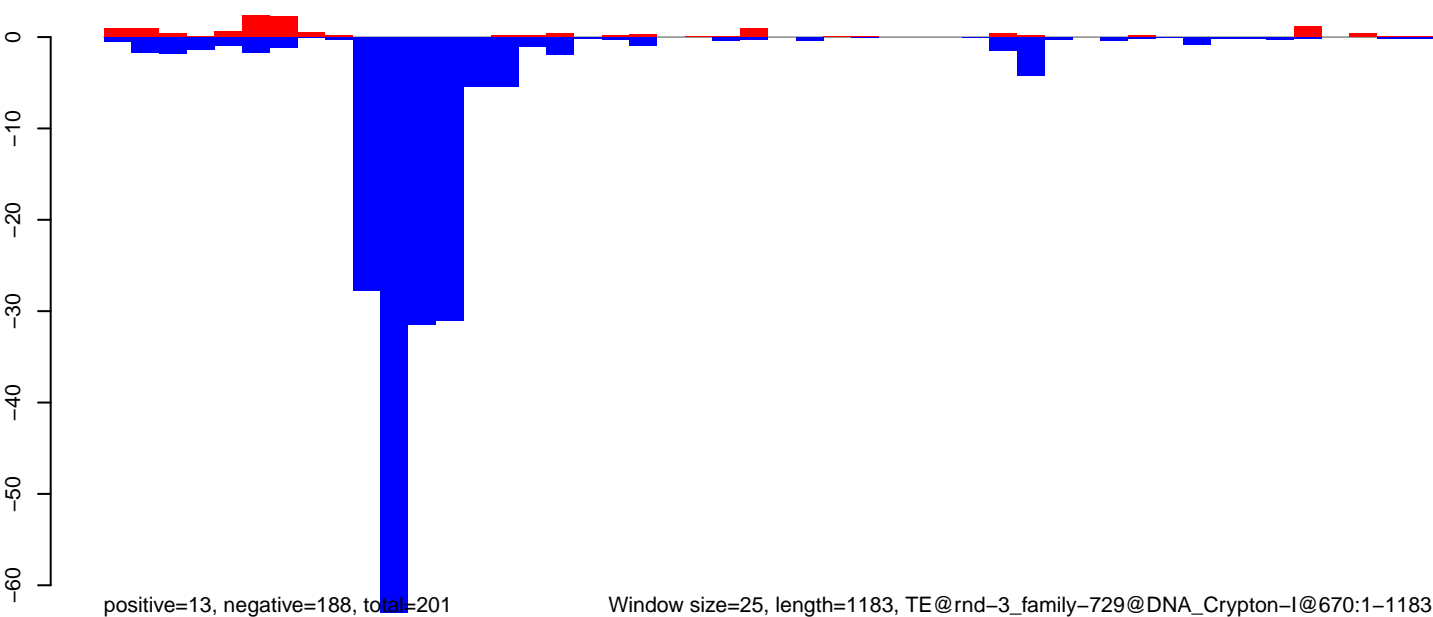
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



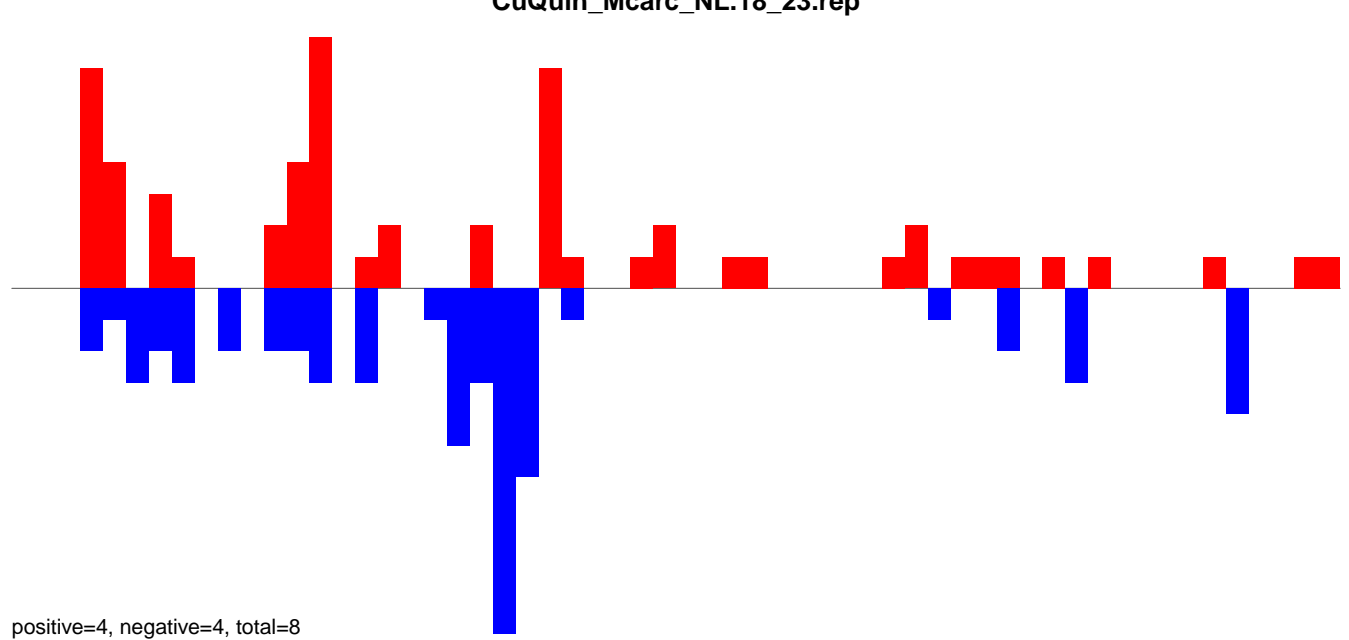
CuQuin_Mcarc_NL.rep



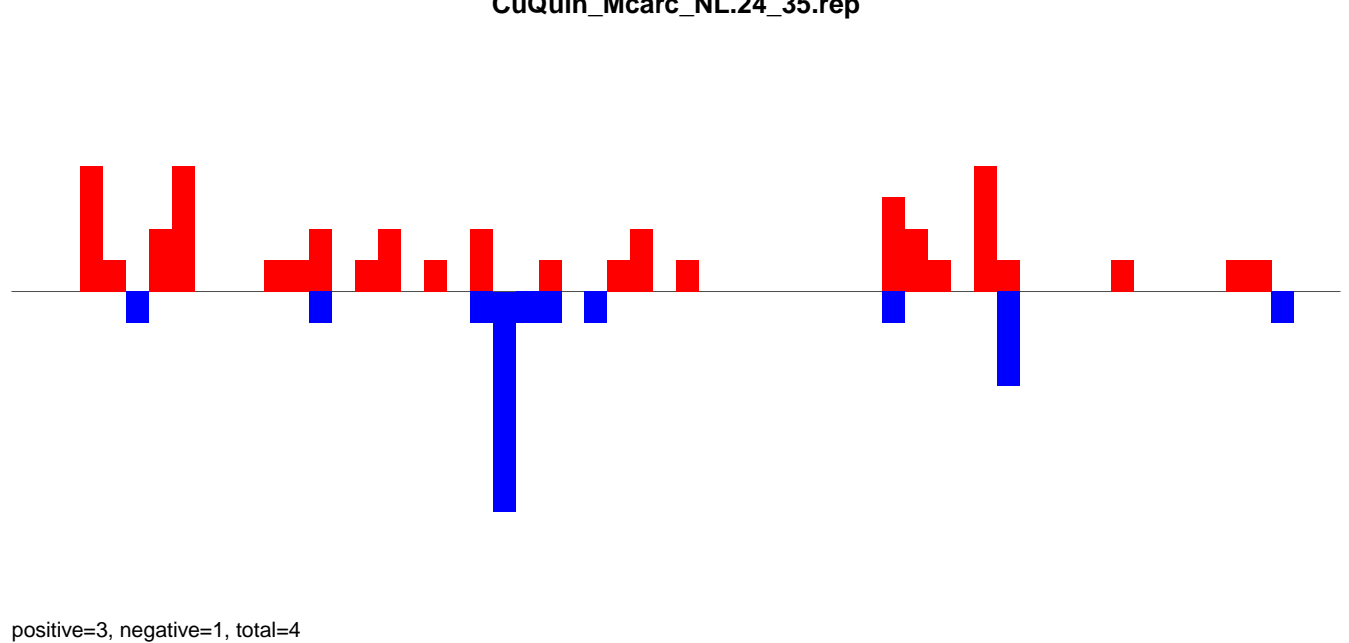
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0 200 400 600 800 1000 1200

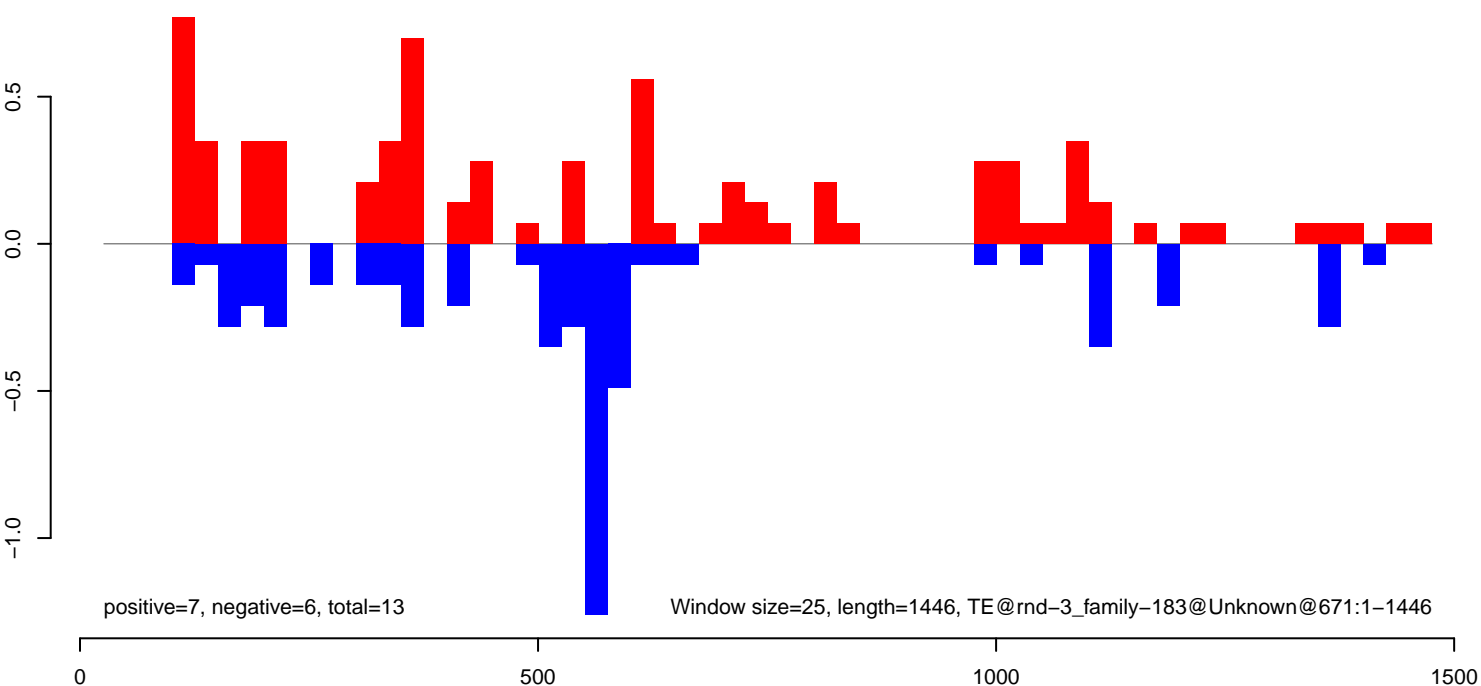
CuQuin_Mcarc_NL.18_23.rep



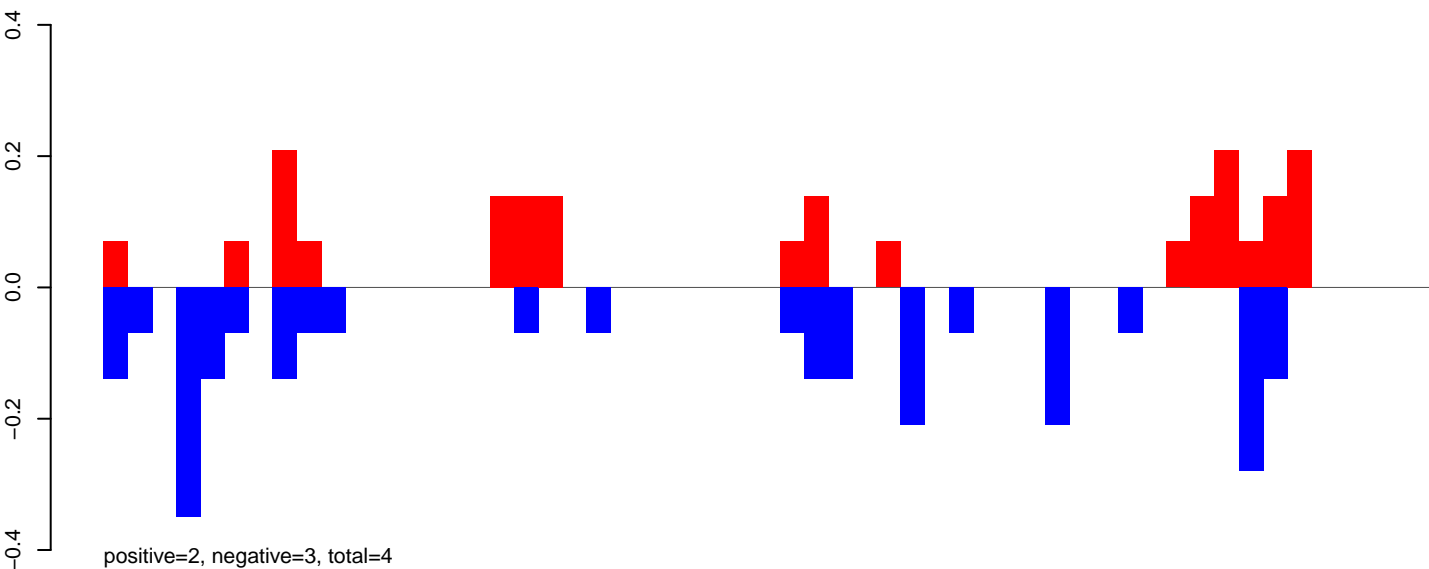
CuQuin_Mcarc_NL.24_35.rep



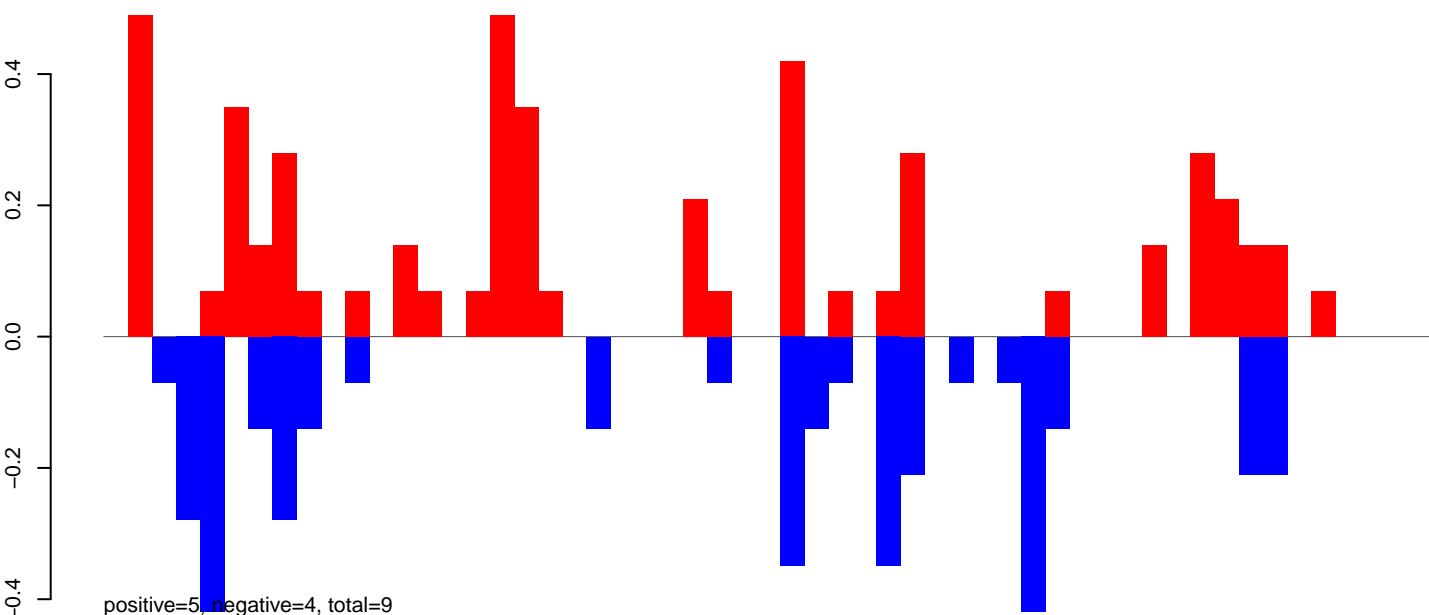
CuQuin_Mcarc_NL.rep



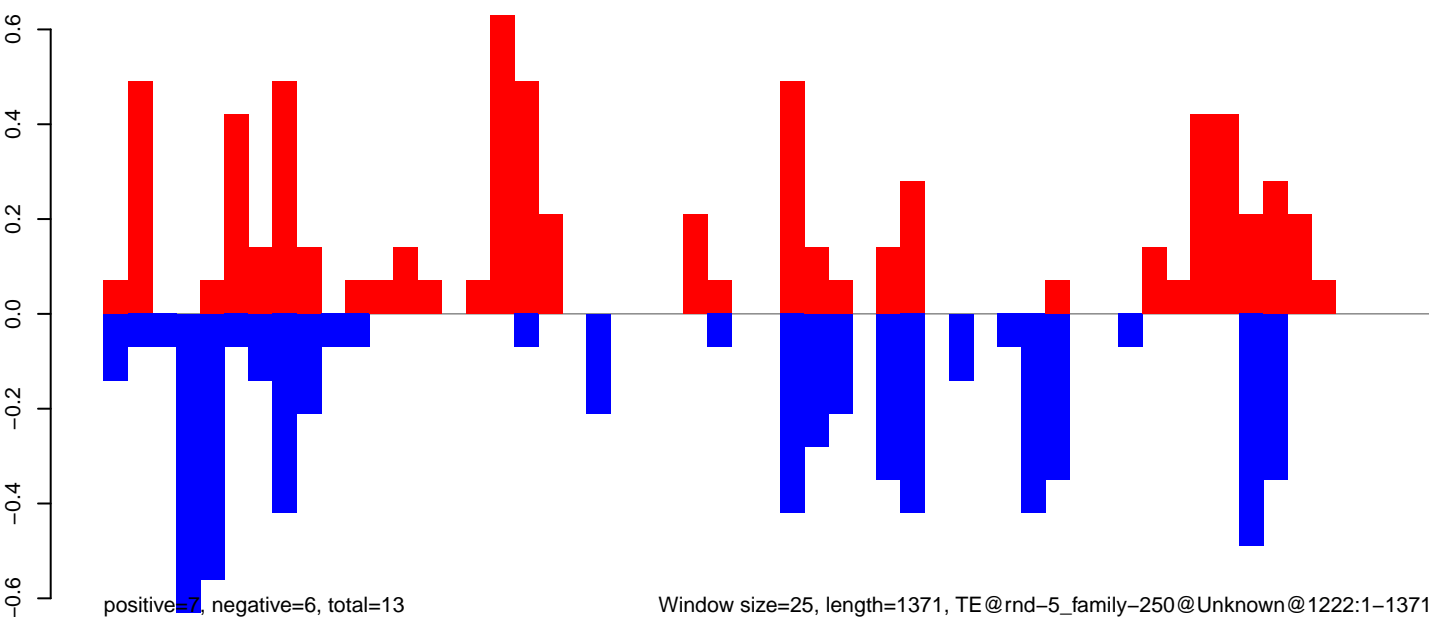
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

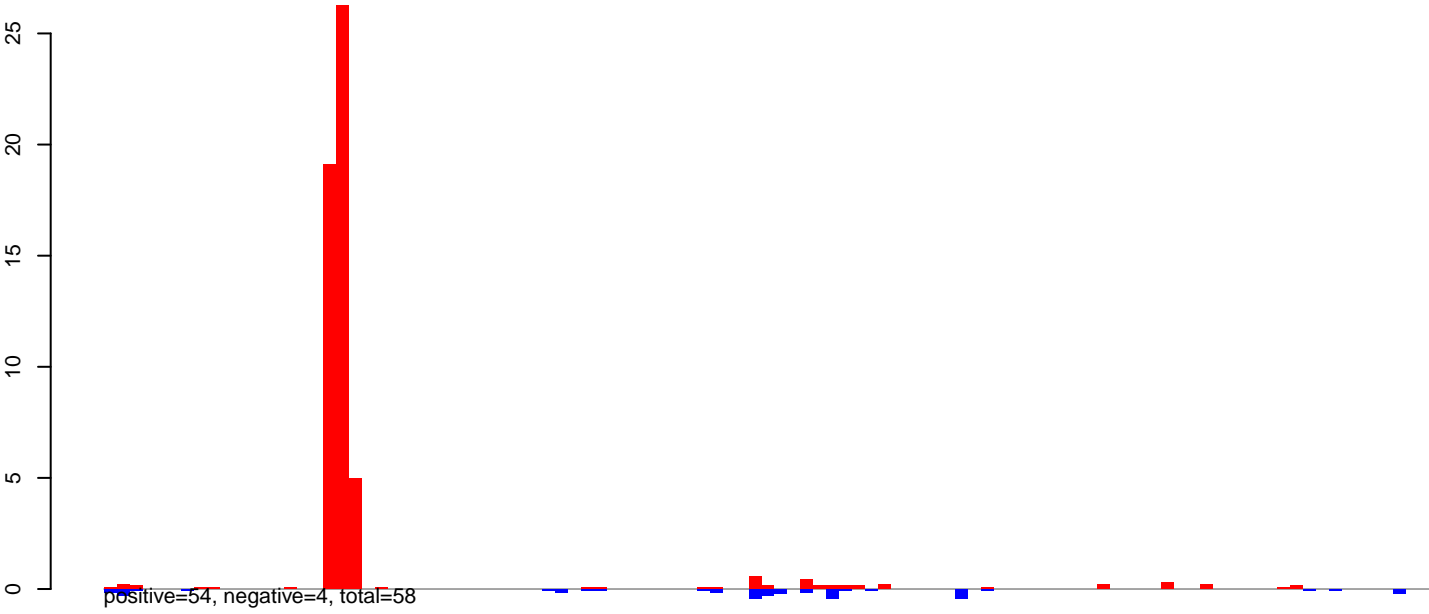


CuQuin_Mcarc_NL.rep

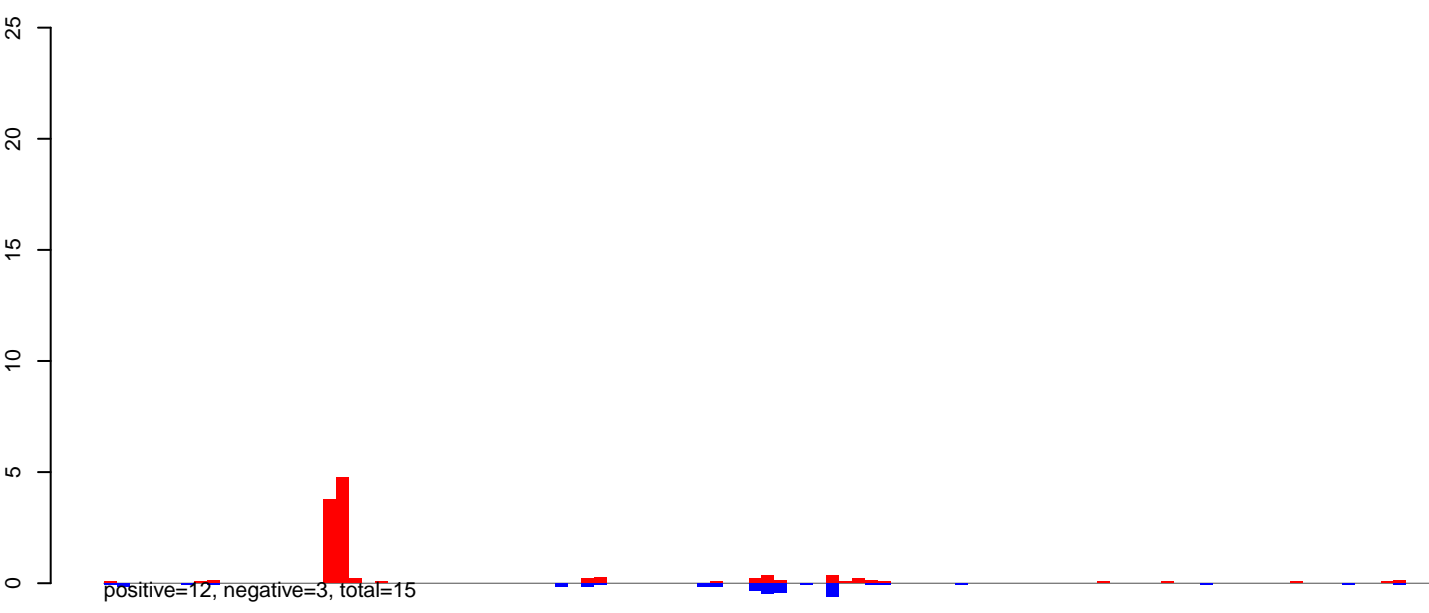


Window size=25, length=1371, TE@rnd-5_family-250@Unknown@1222:1-1371

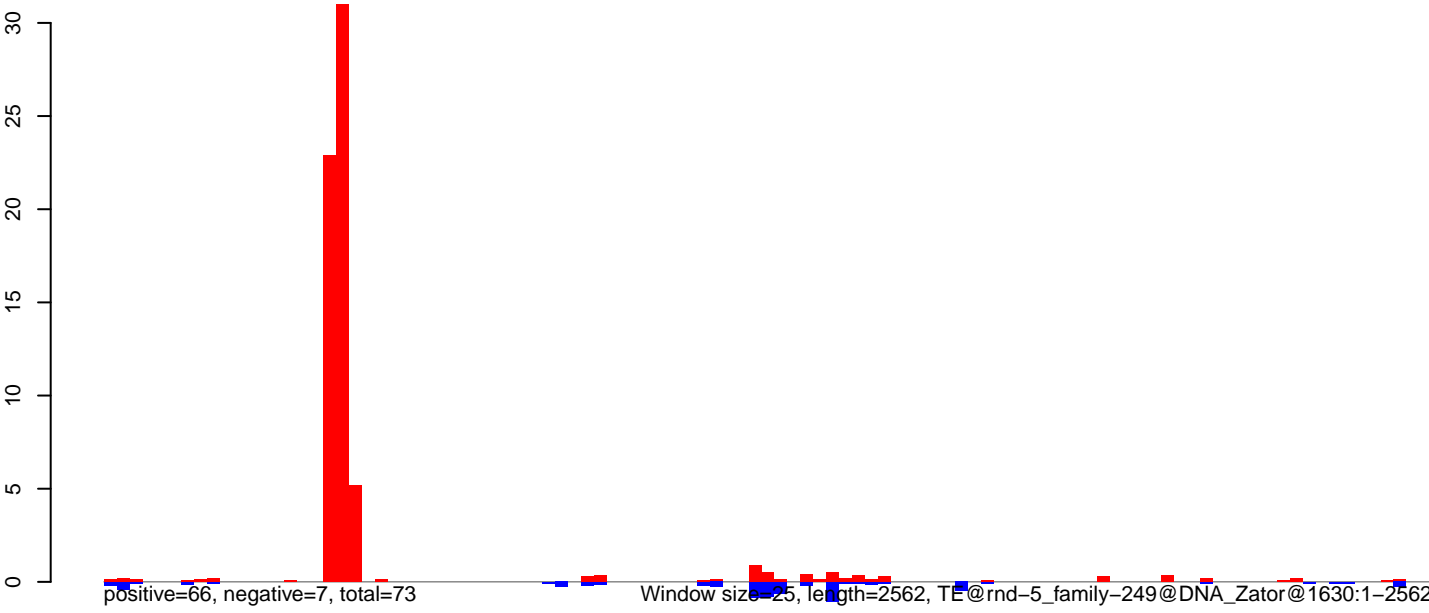
CuQuin_Mcarc_NL.18_23.rep



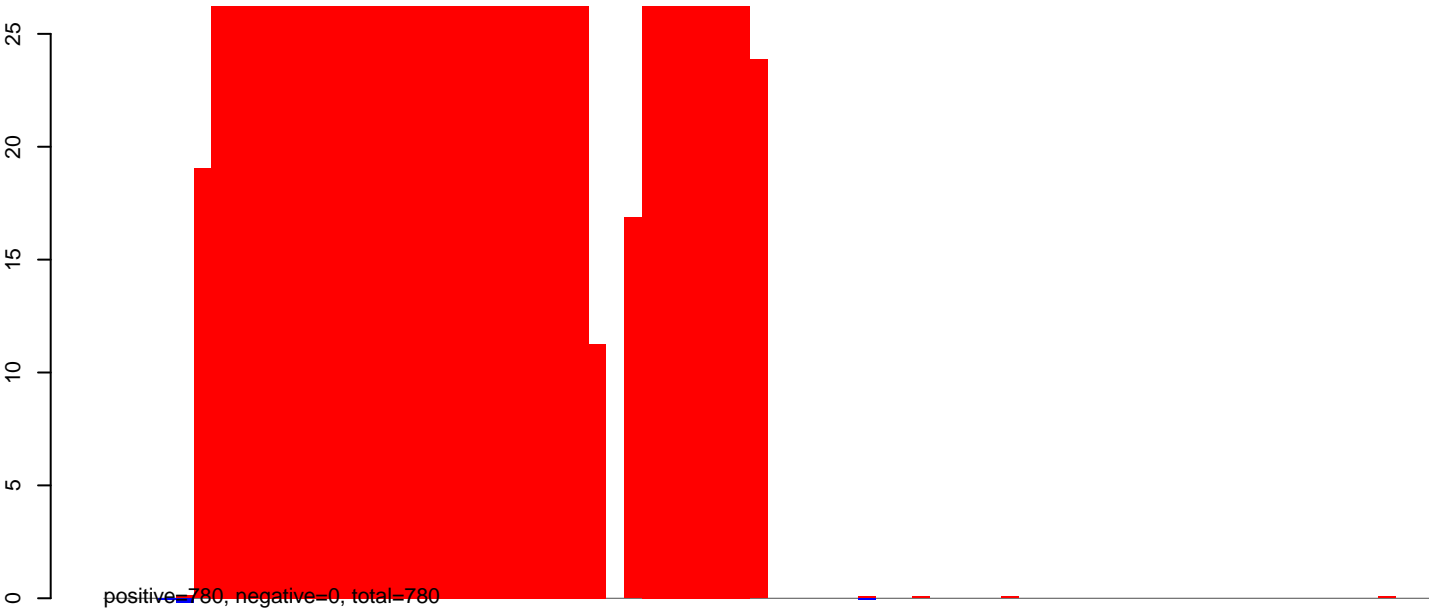
CuQuin_Mcarc_NL.24_35.rep



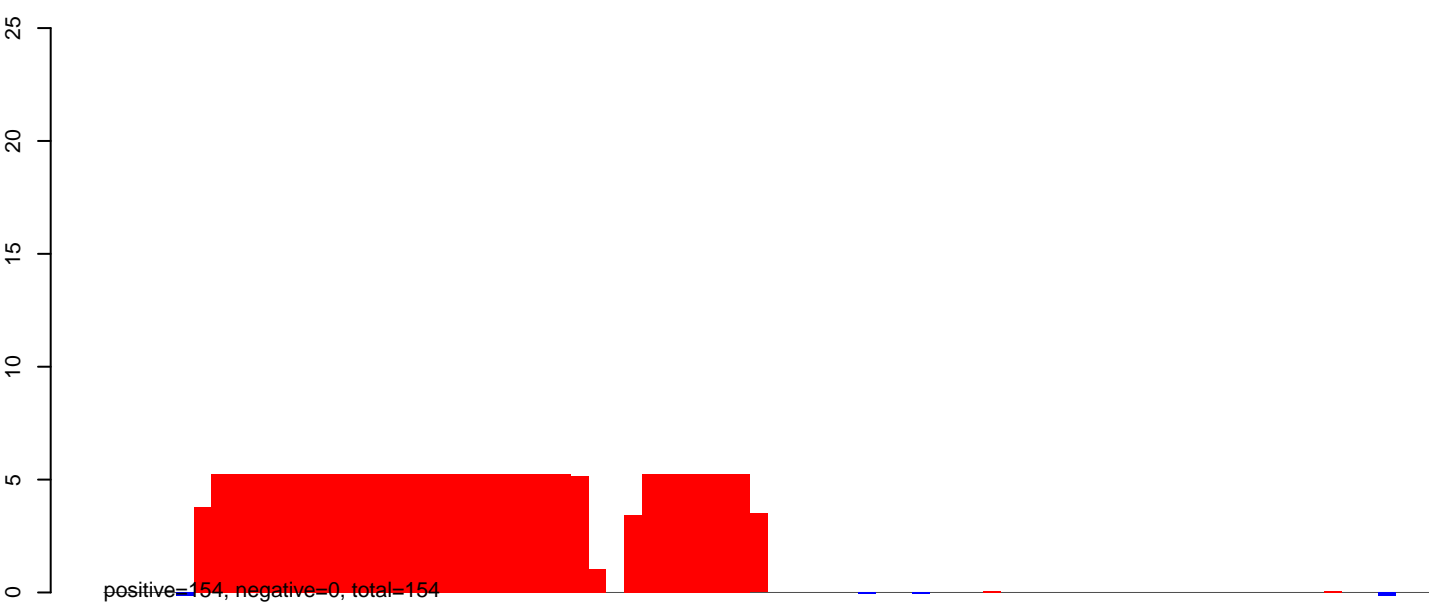
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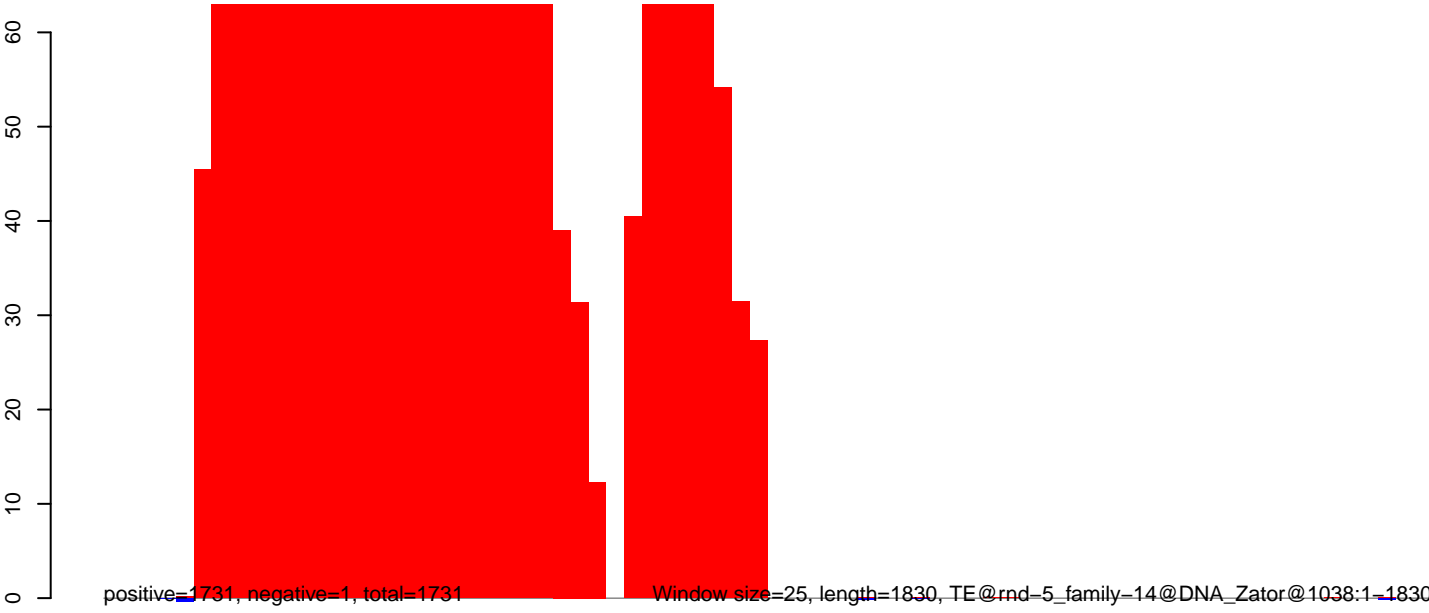
CuQuin_Mcarc_NL.18_23.rep



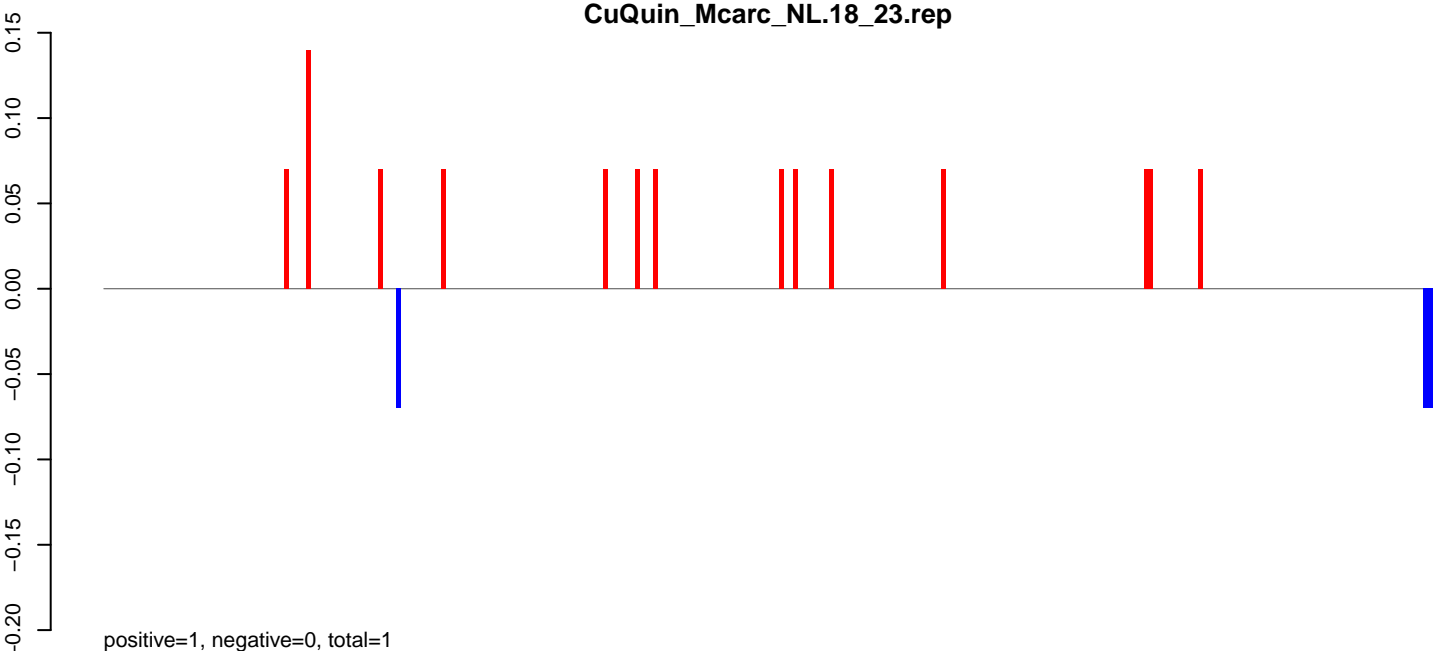
CuQuin_Mcarc_NL.24_35.rep



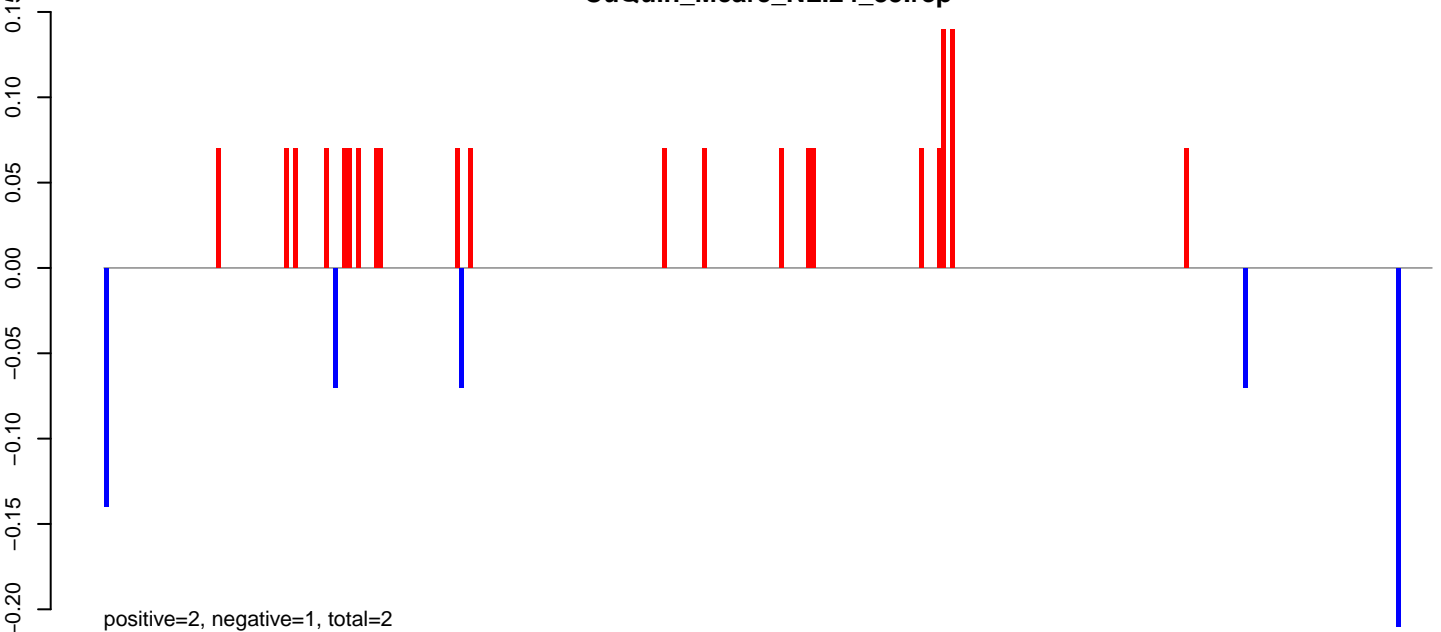
CuQuin_Mcarc_NL.rep



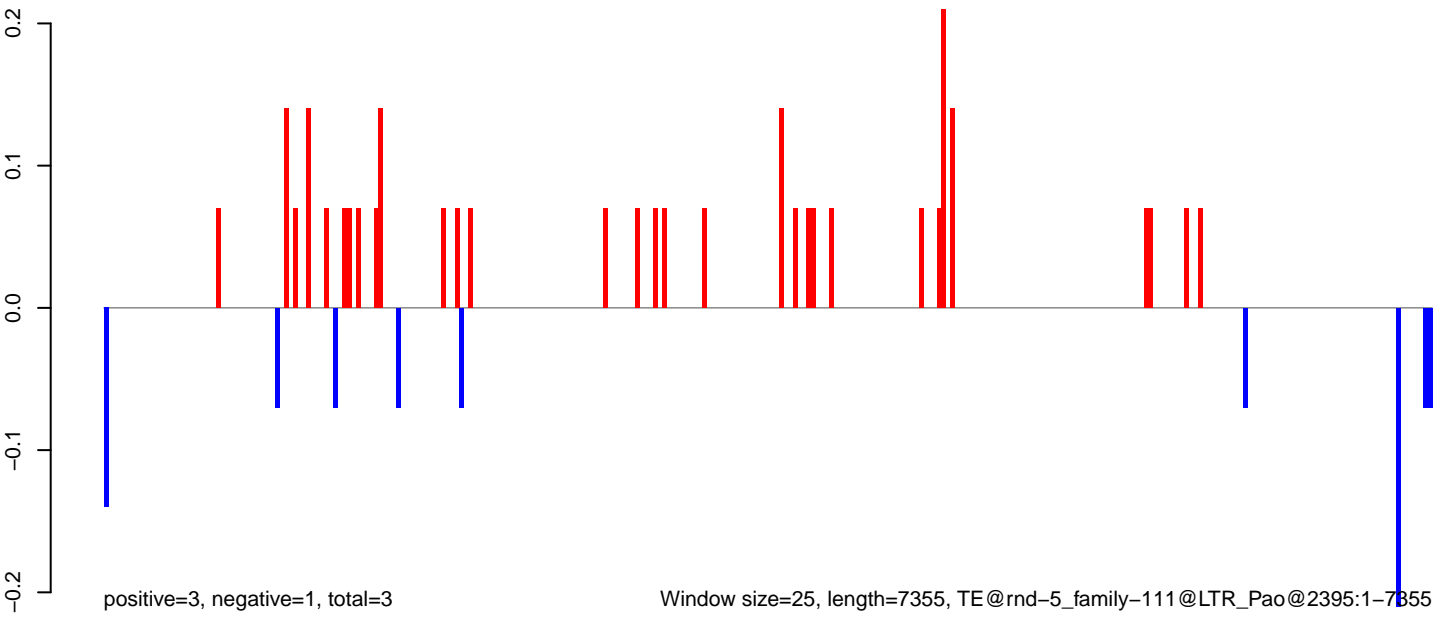
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CuQuin_Mcarc_NL.24_35.rep

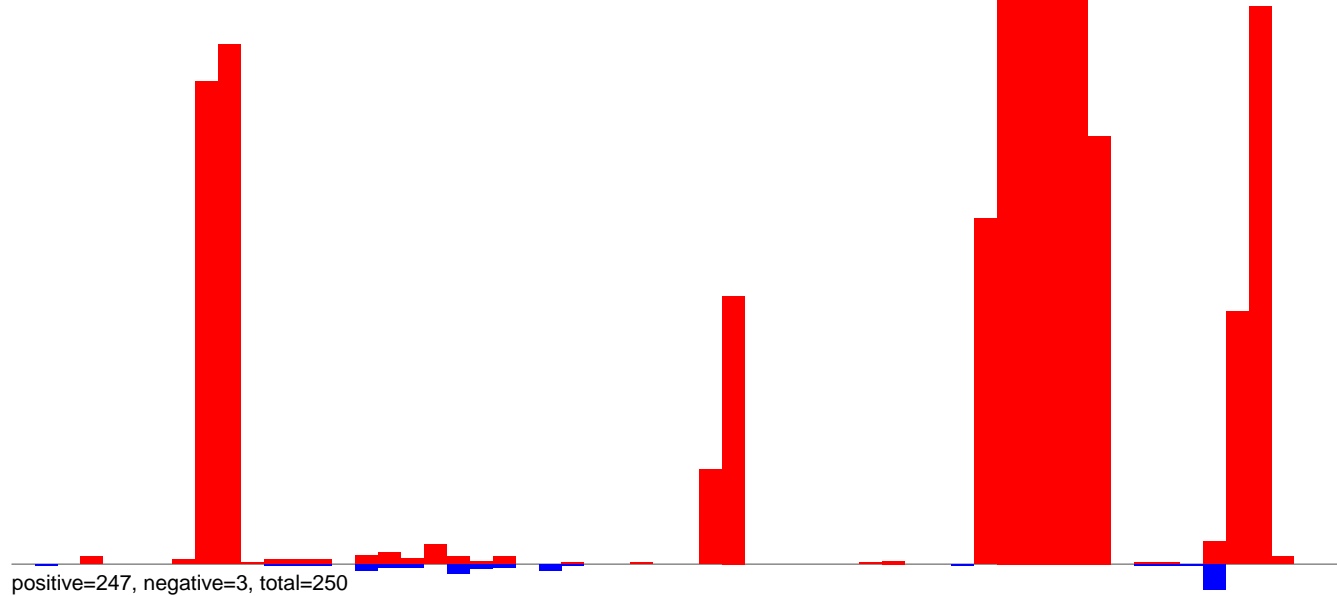


CuQuin_Mcarc_NL.rep

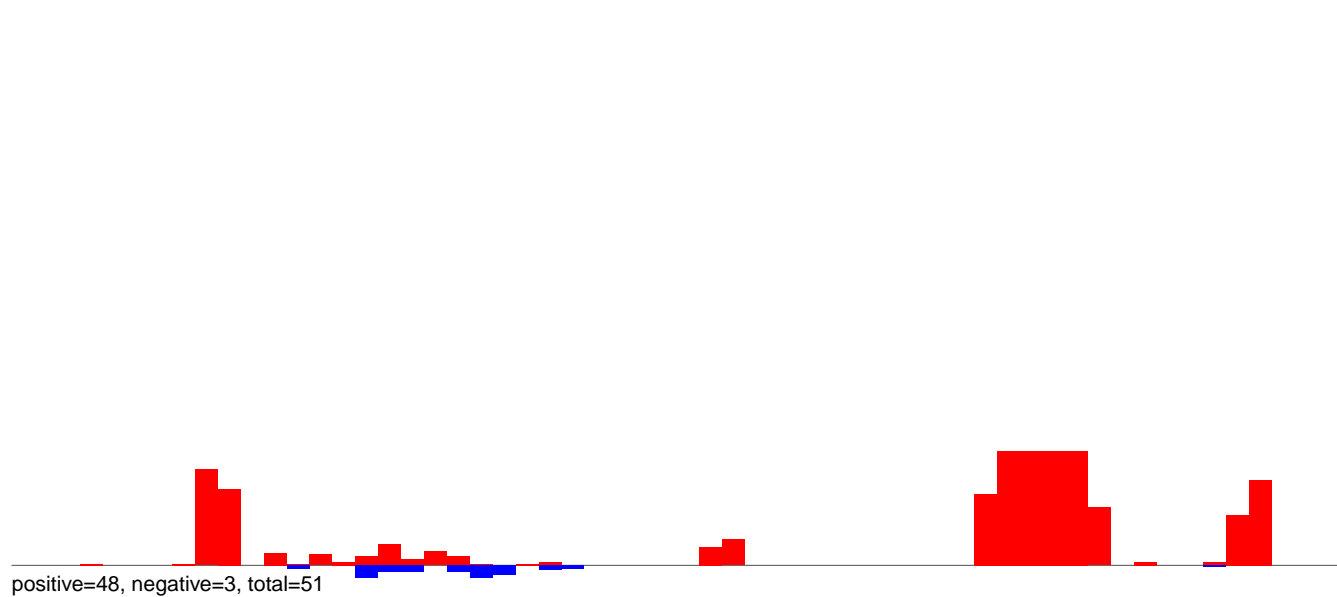


Window size=25, length=7355, TE@rnd-5_family-111@LTR_Pao@2395:1-7355

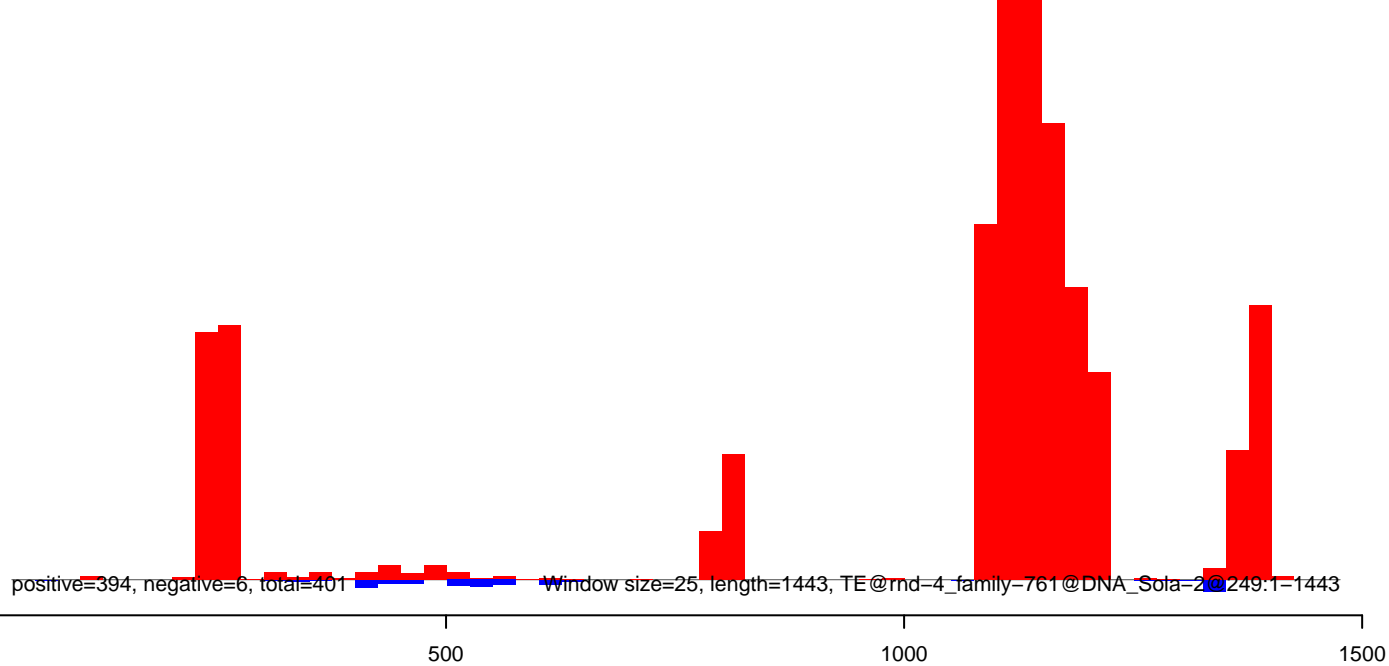
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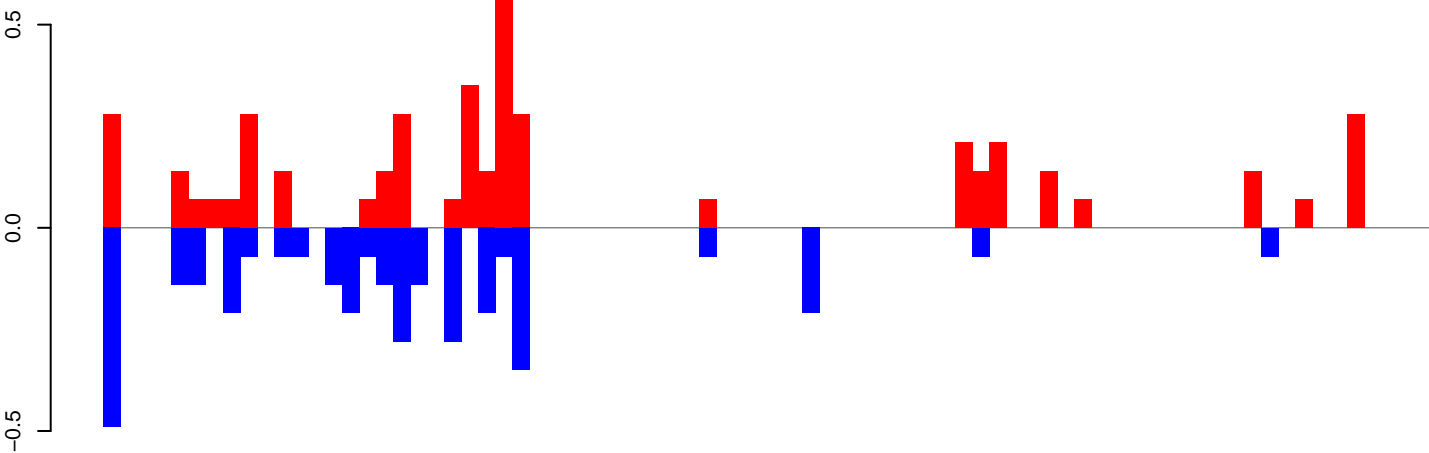
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

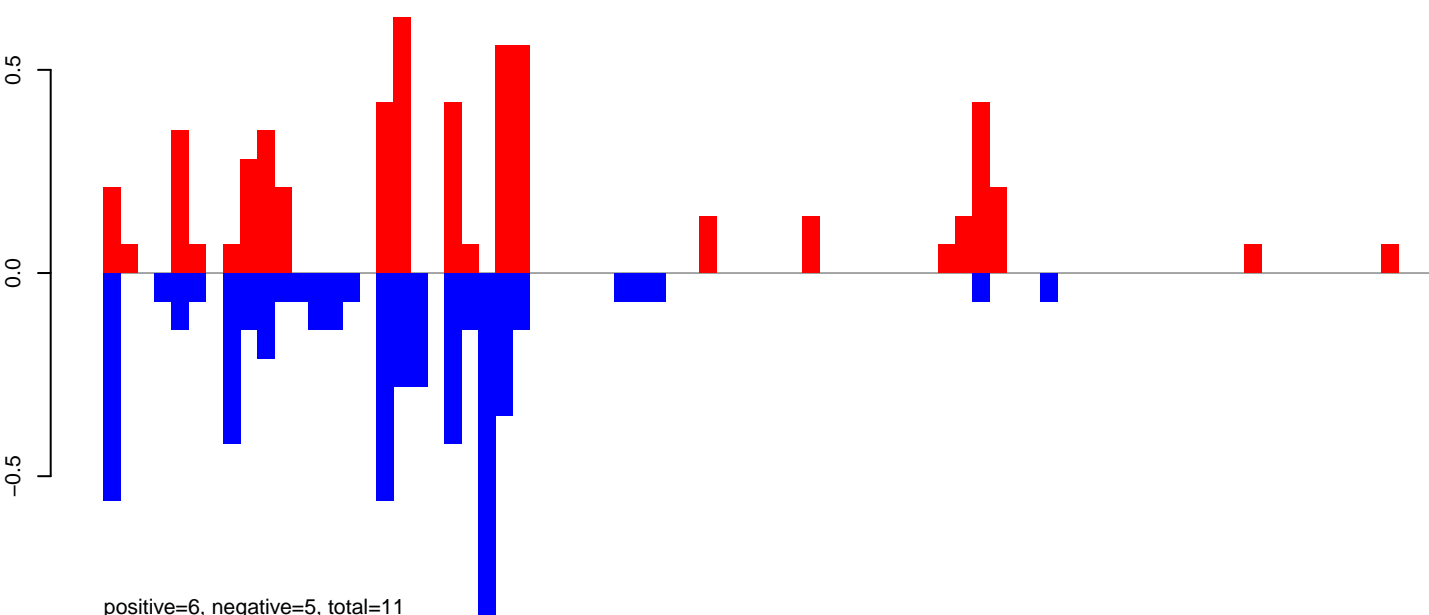


CuQuin_Mcarc_NL.18_23.rep



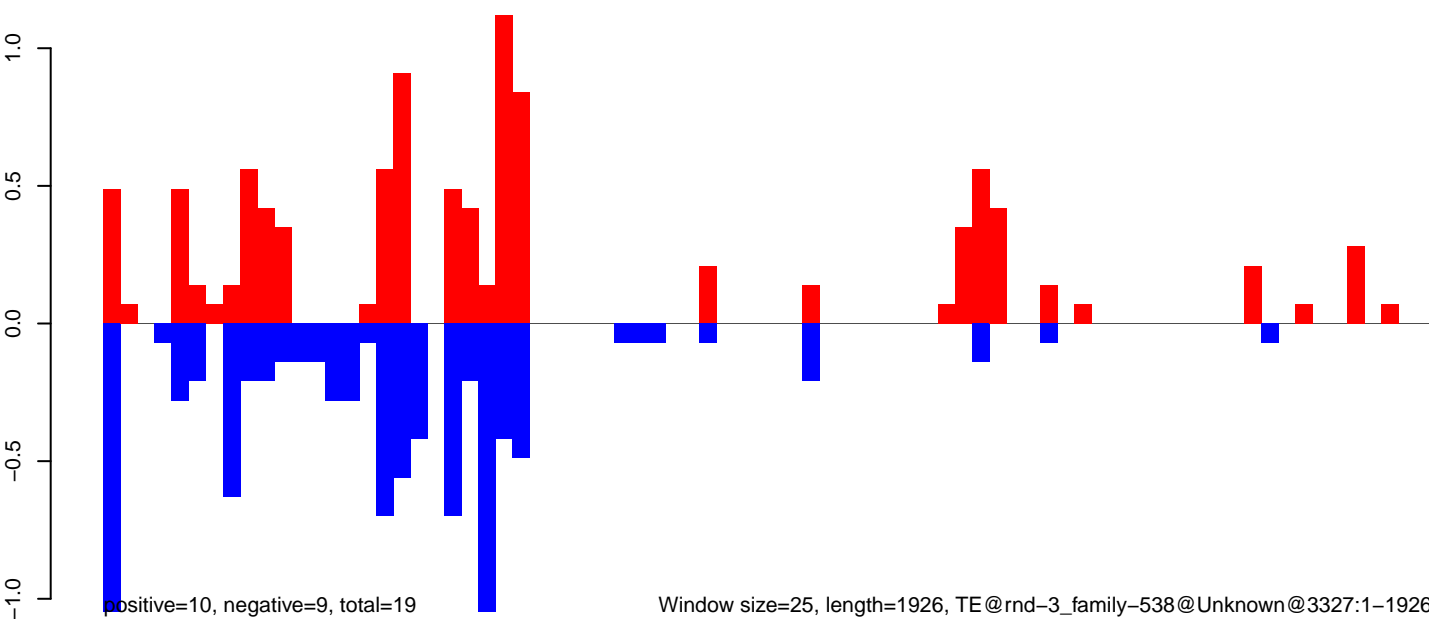
positive=4, negative=3, total=8

CuQuin_Mcarc_NL.24_35.rep



positive=6, negative=5, total=11

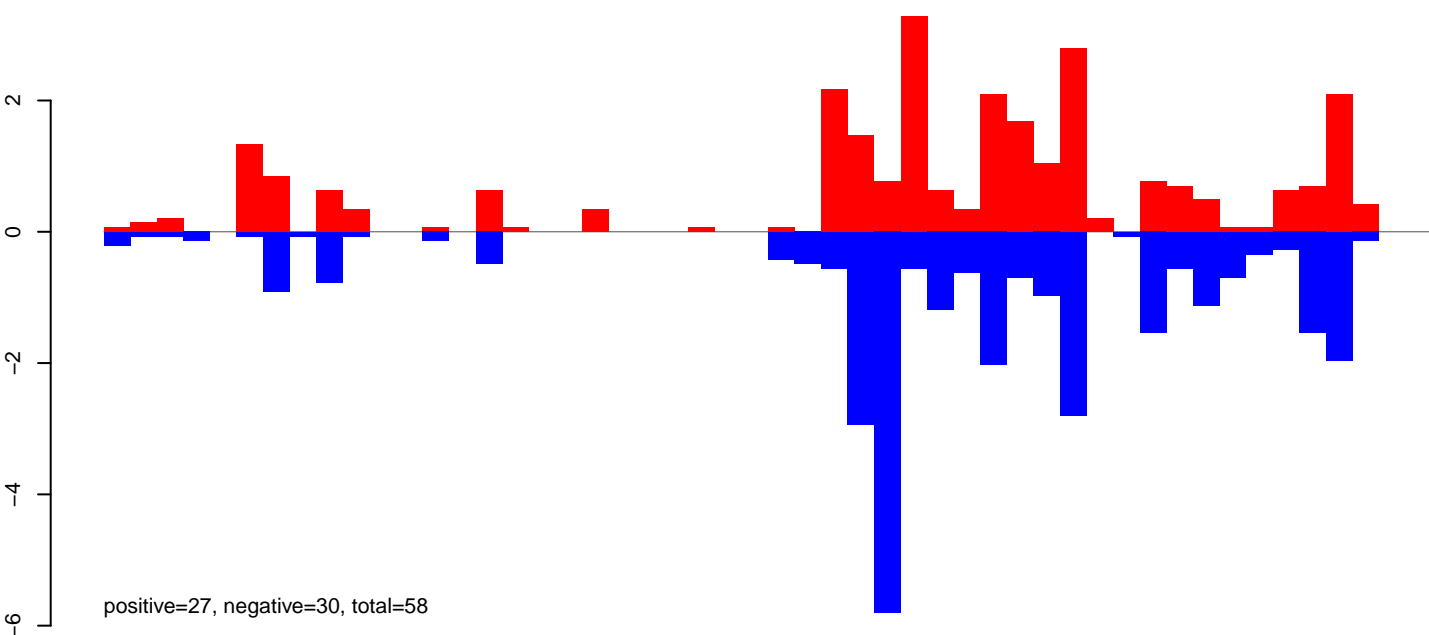
CuQuin_Mcarc_NL.rep



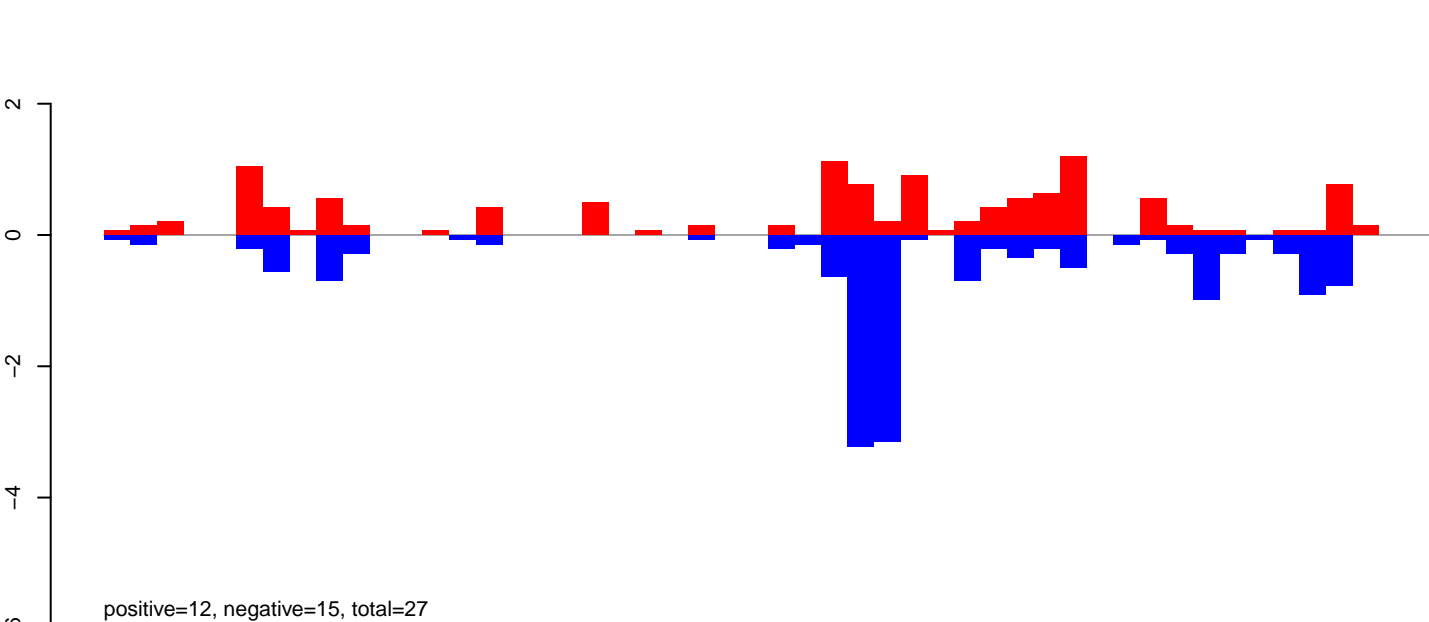
positive=10, negative=9, total=19

Window size=25, length=1926, TE@rnd-3_family-538@Unknown@3327:1-1926

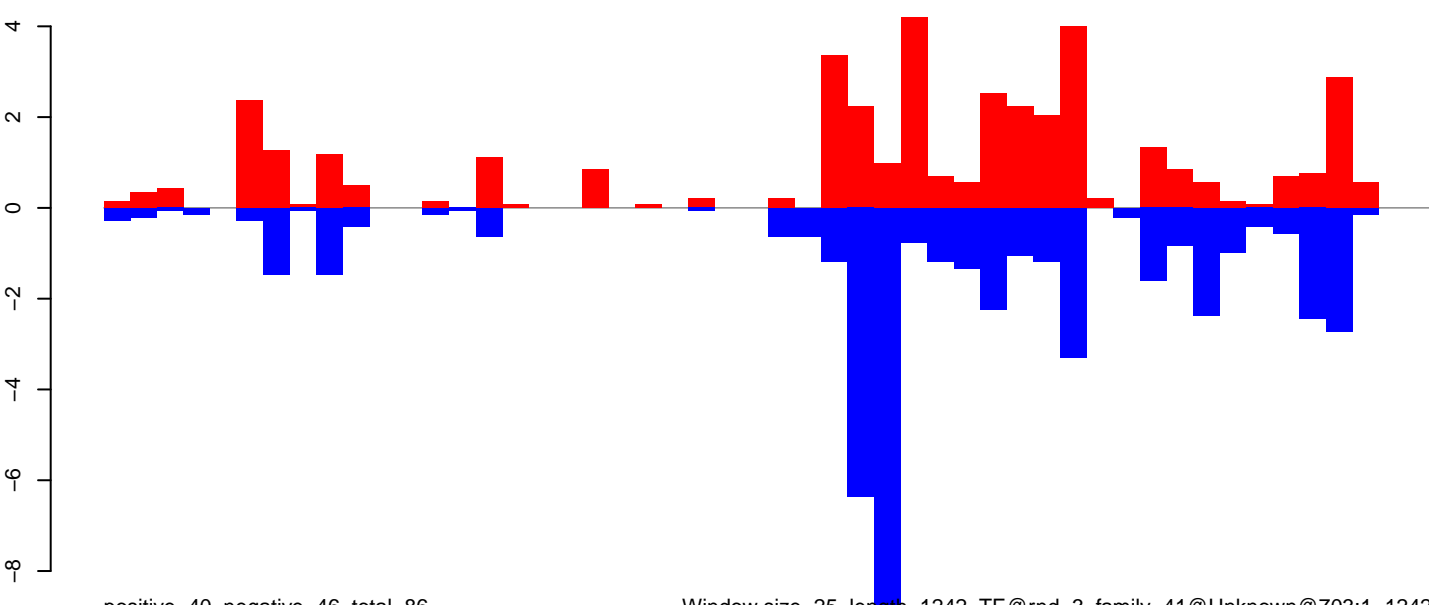
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



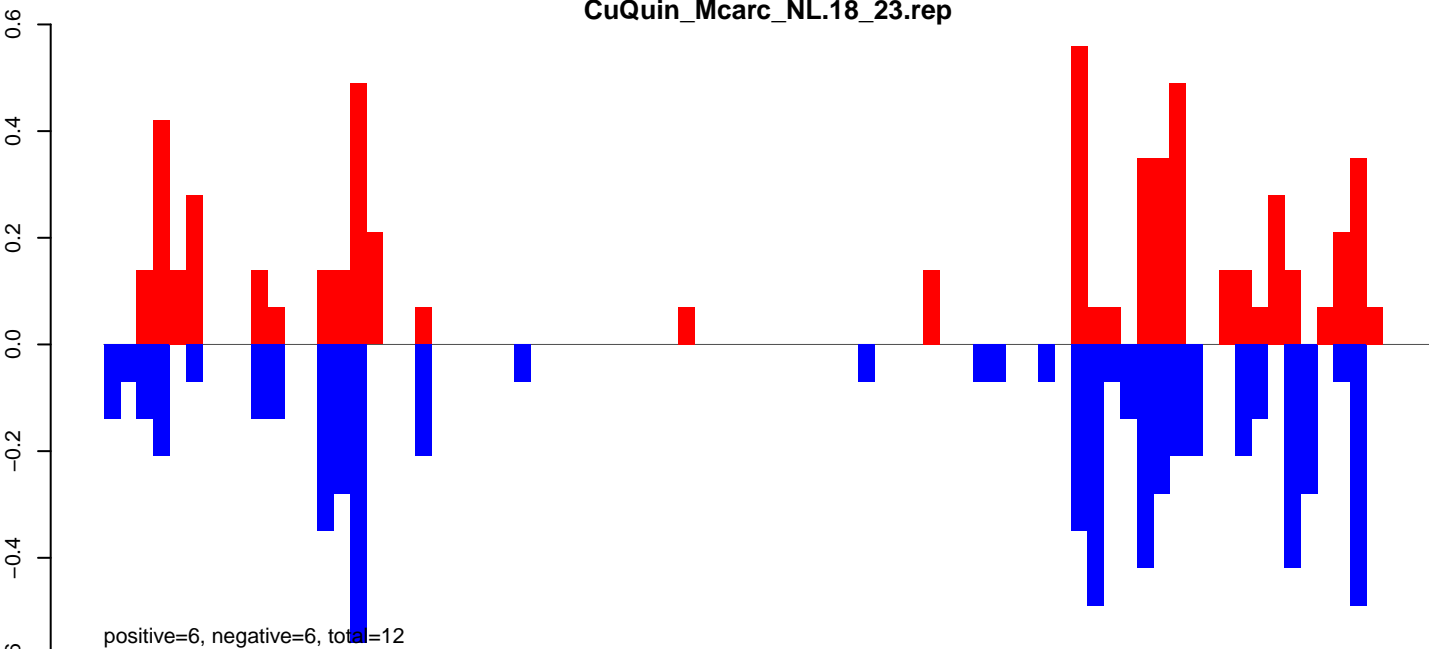
CuQuin_Mcarc_NL.rep



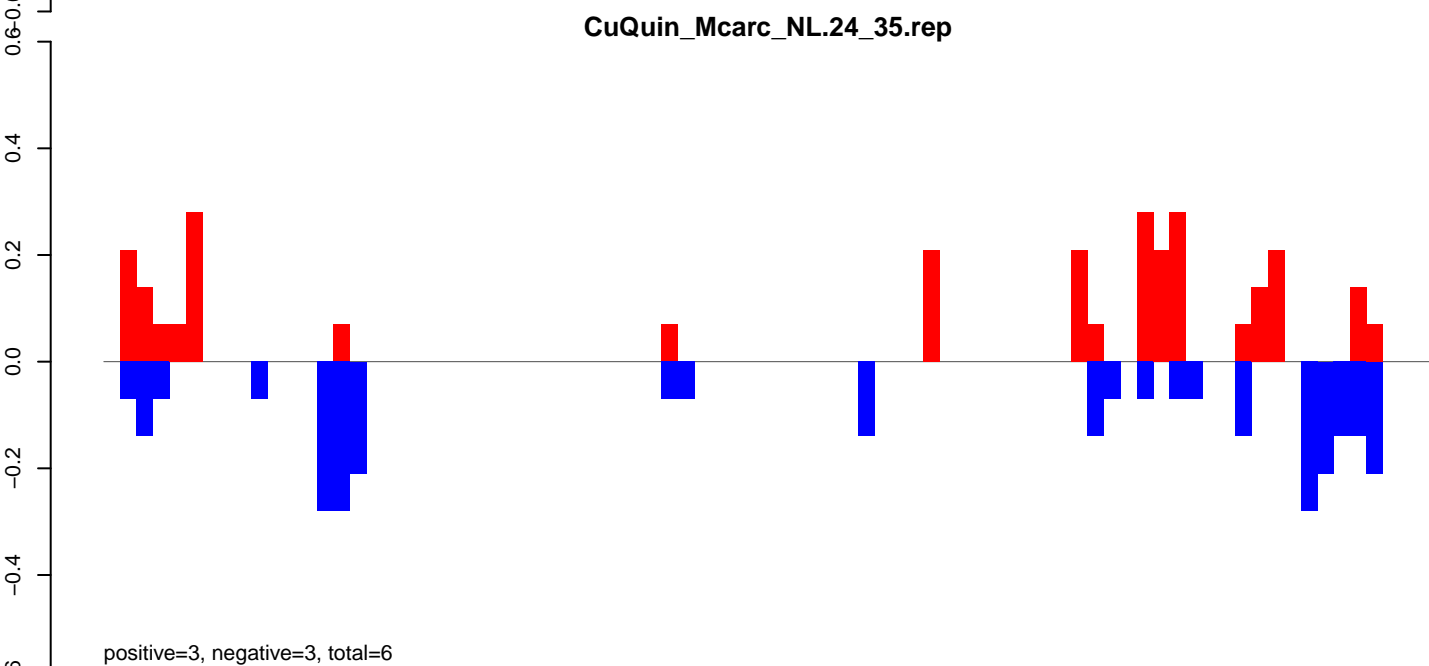
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0 200 400 600 800 1000 1200

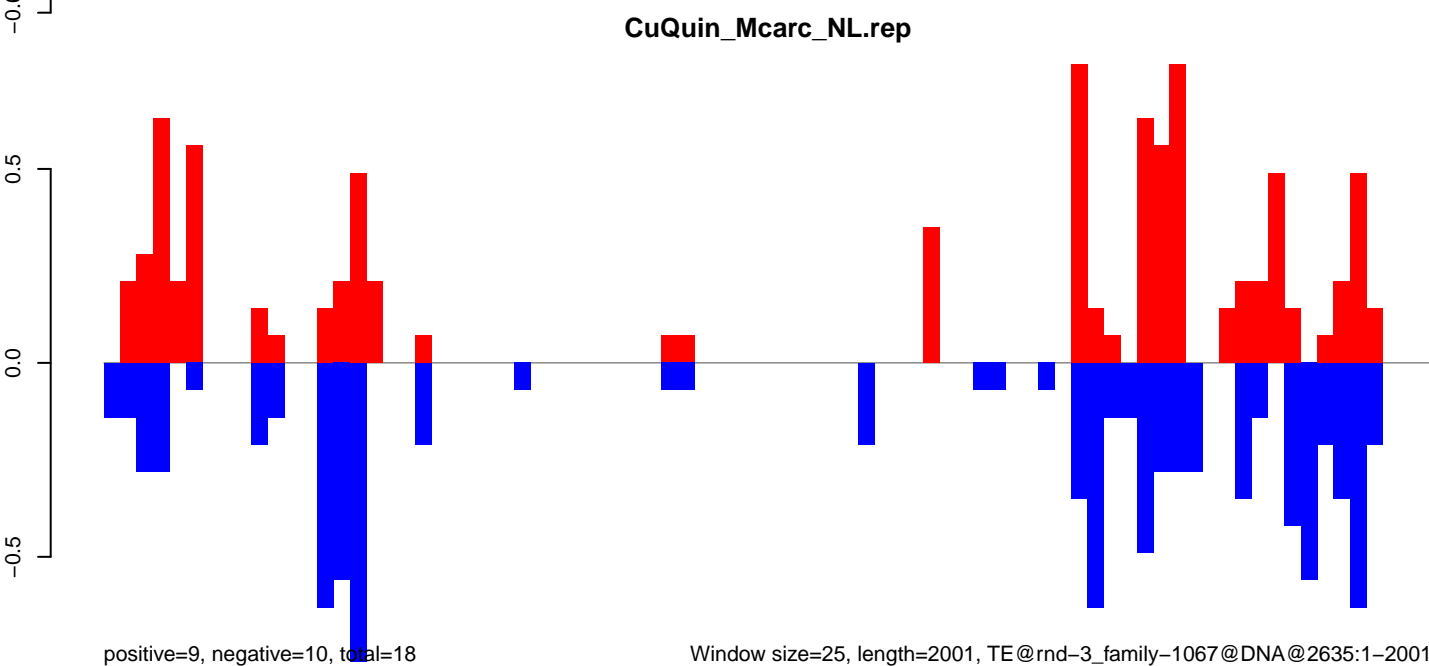
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



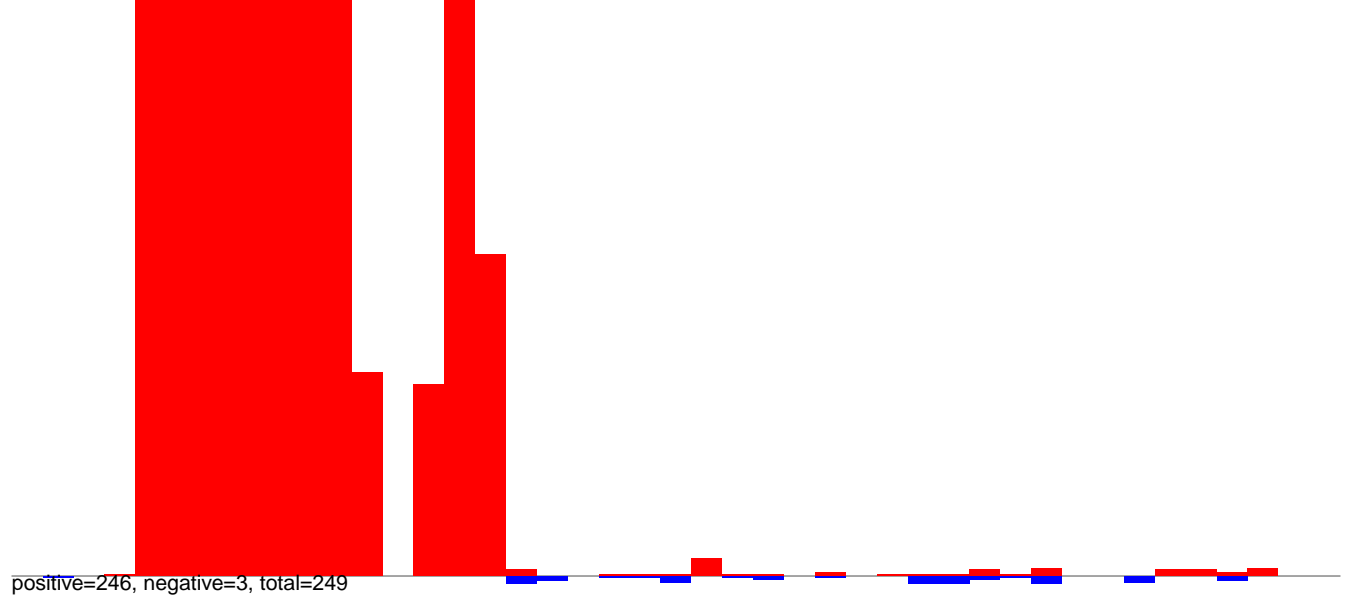
CuQuin_Mcarc_NL.rep



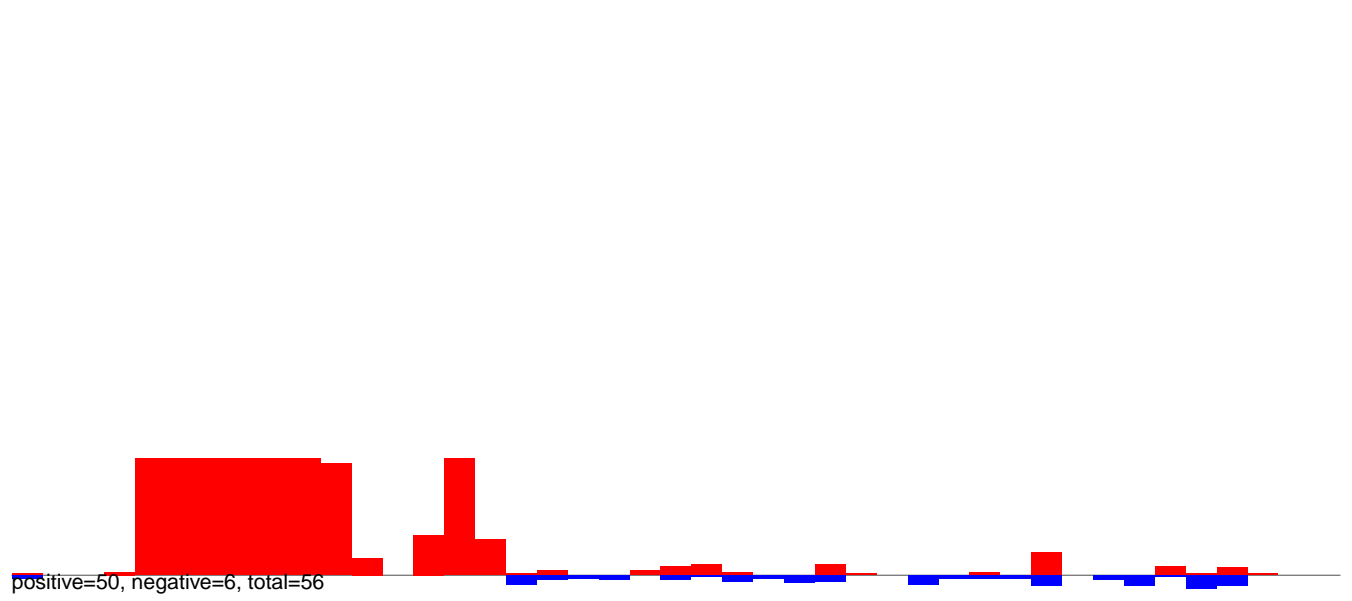
Window size=25, length=2001, TE@rnd-3_family-1067@DNA@2635:1-2001

0 500 1000 1500 2000

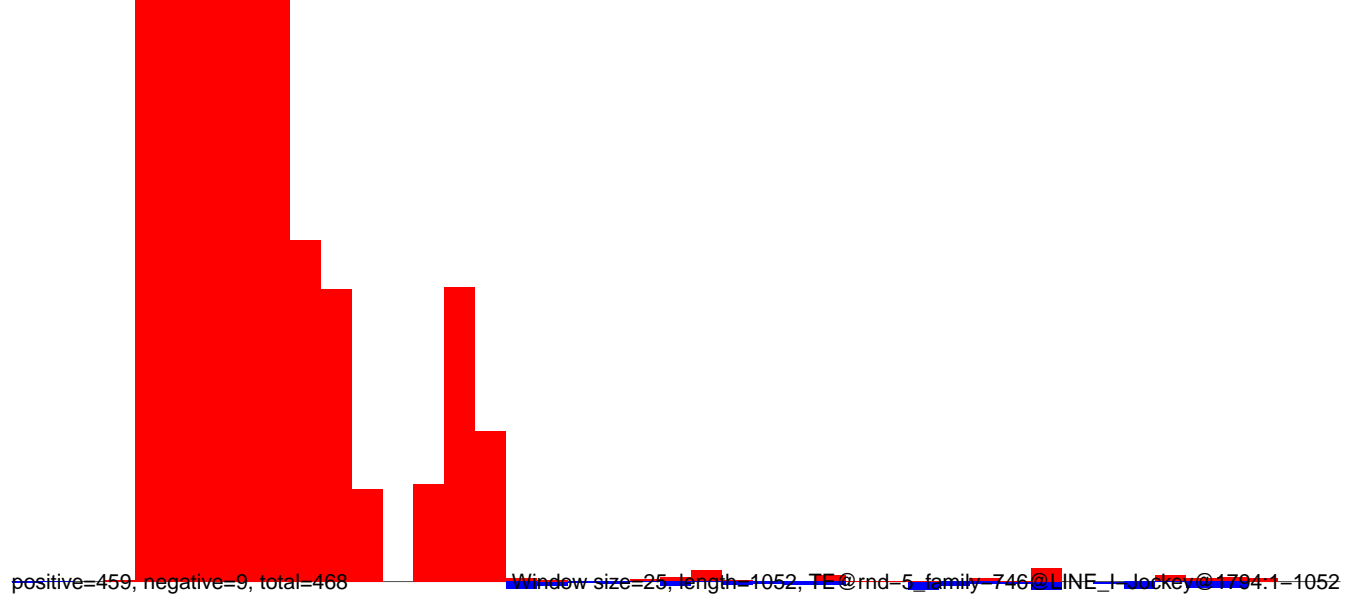
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

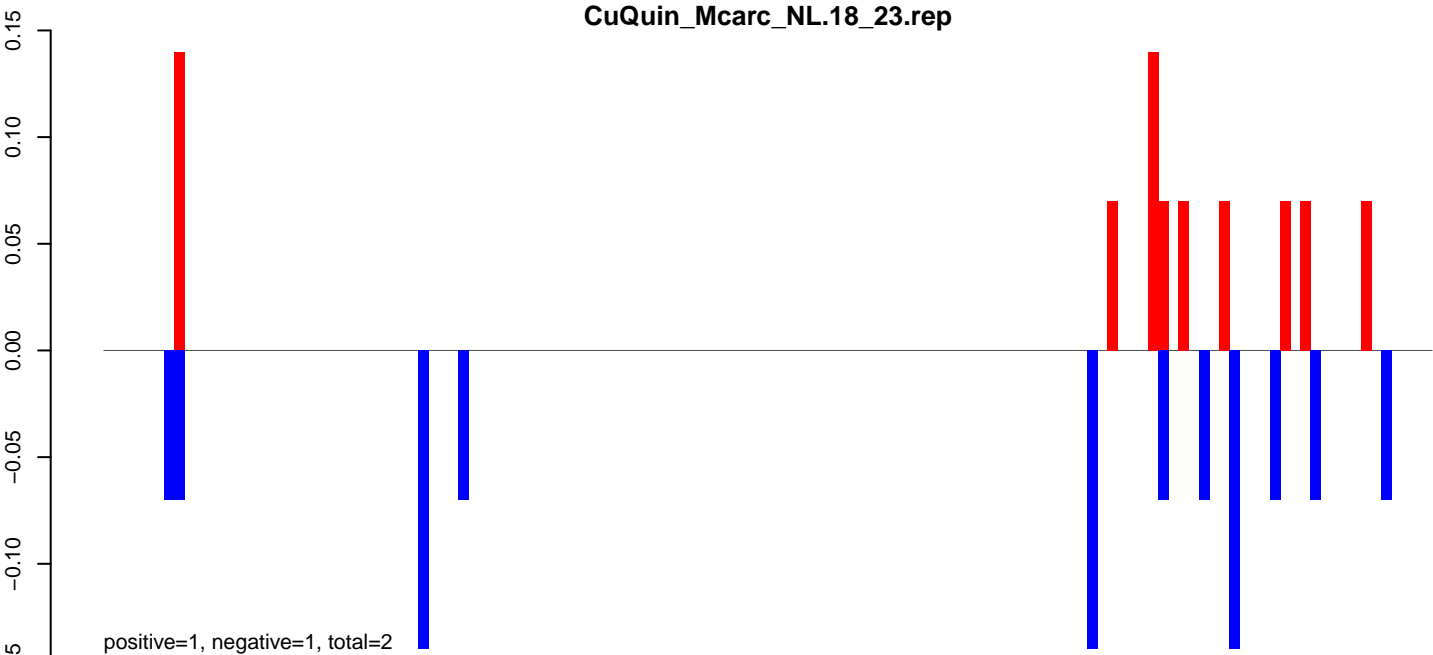


CuQuin_Mcarc_NL.rep

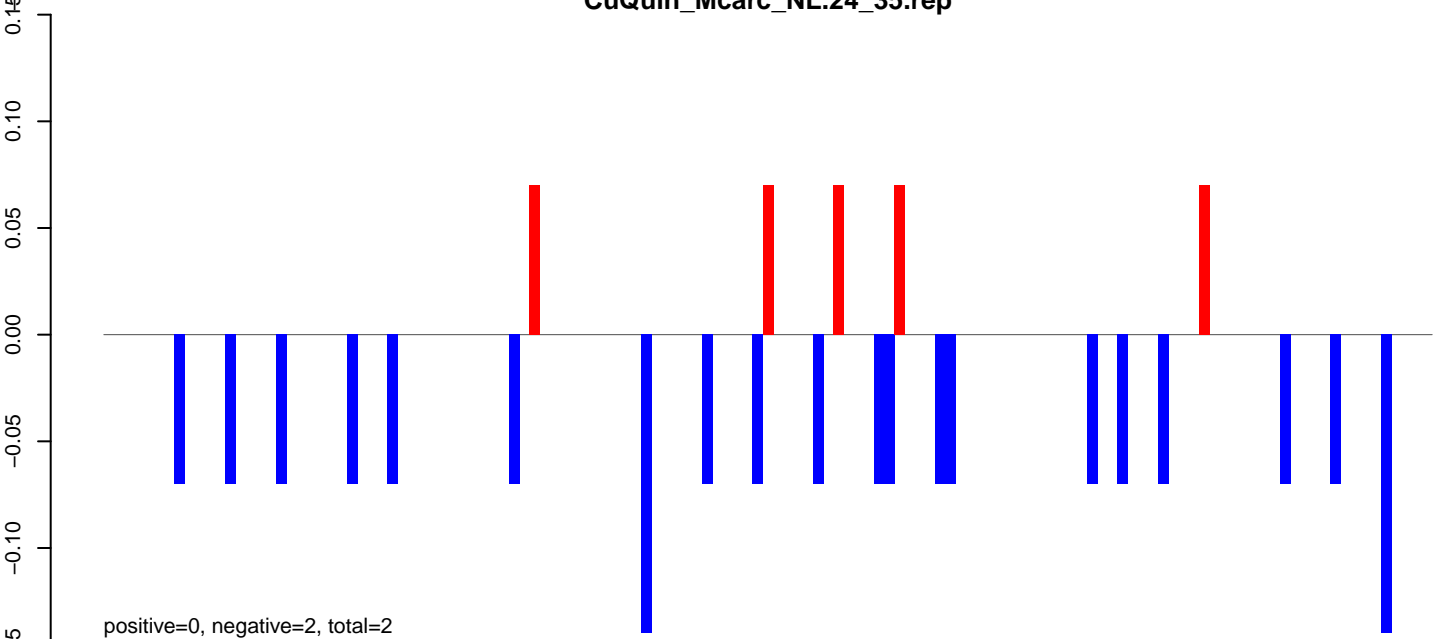


Window size=25, length=1052, TE@rnd=5, family=746@LINE_I, Jockey@1794.1-1052

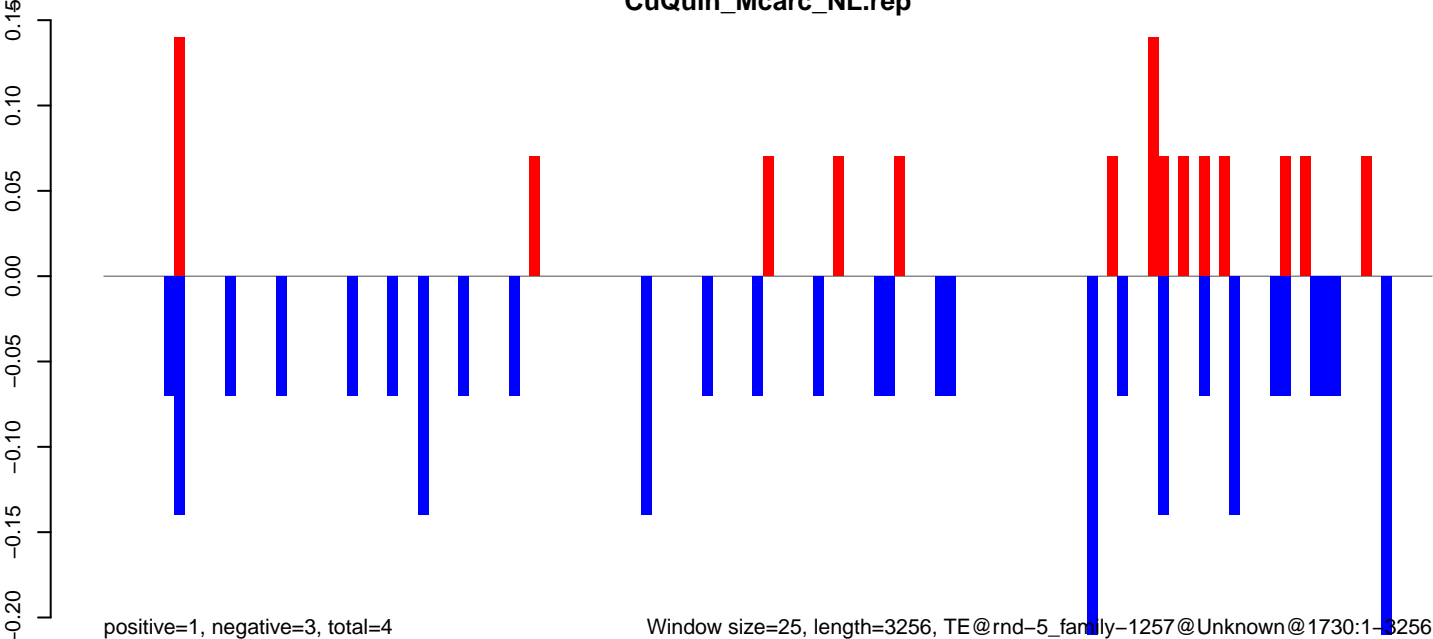
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

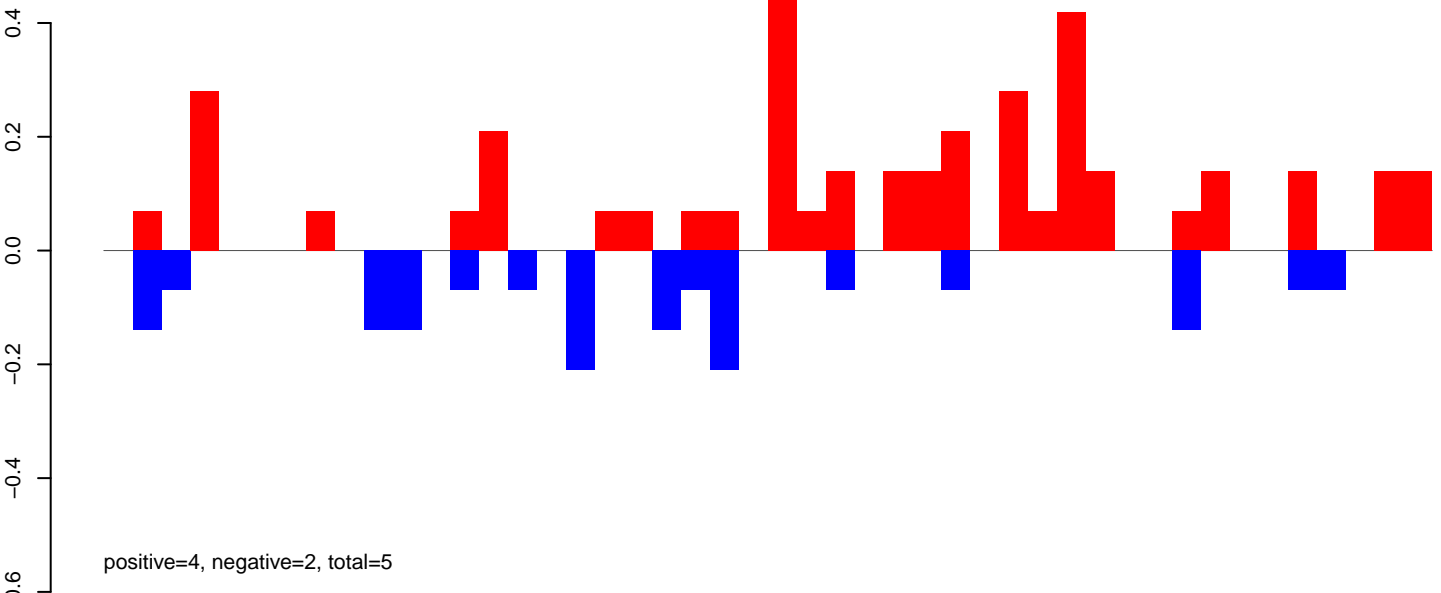


CuQuin_Mcarc_NL.rep

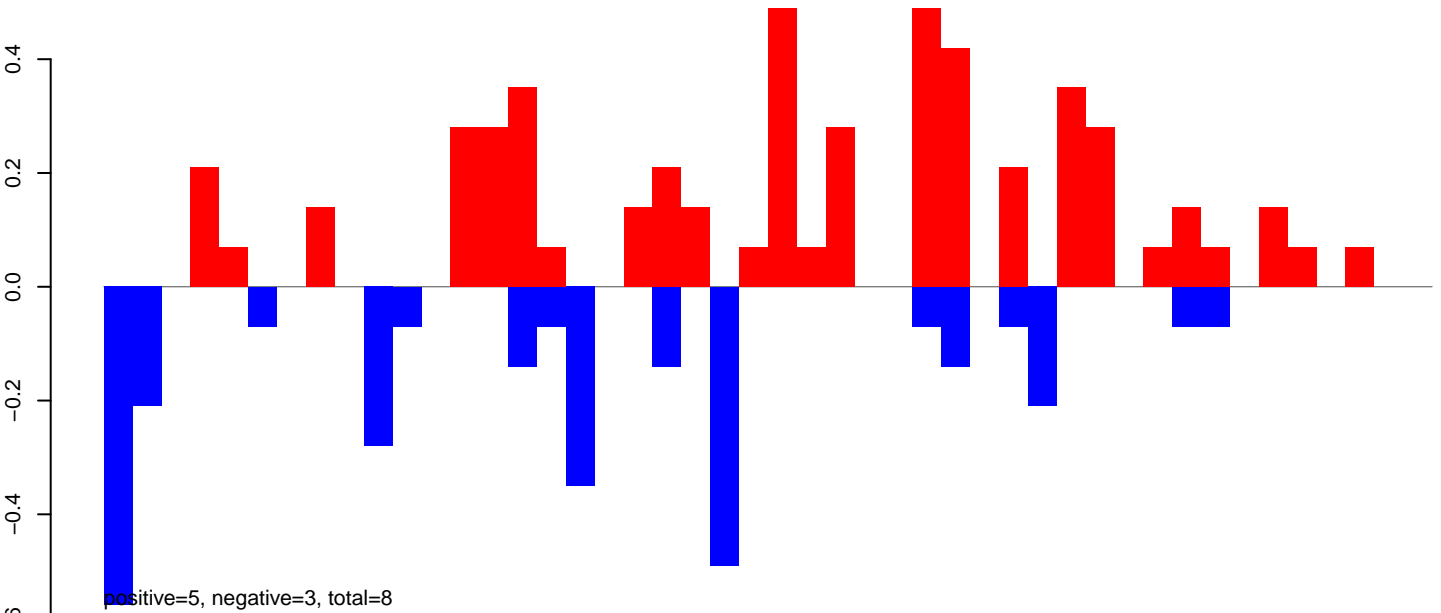


Window size=25, length=3256, TE@rnd-5_family-1257@Unknown@1730:1-3256

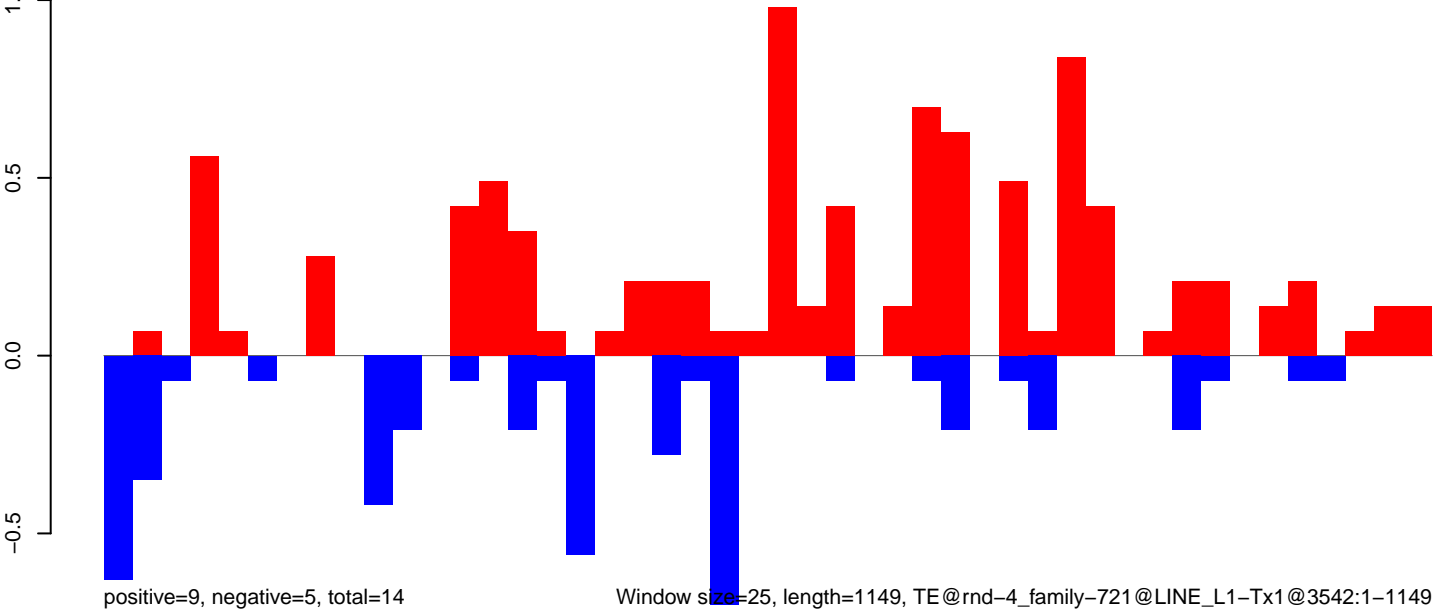
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

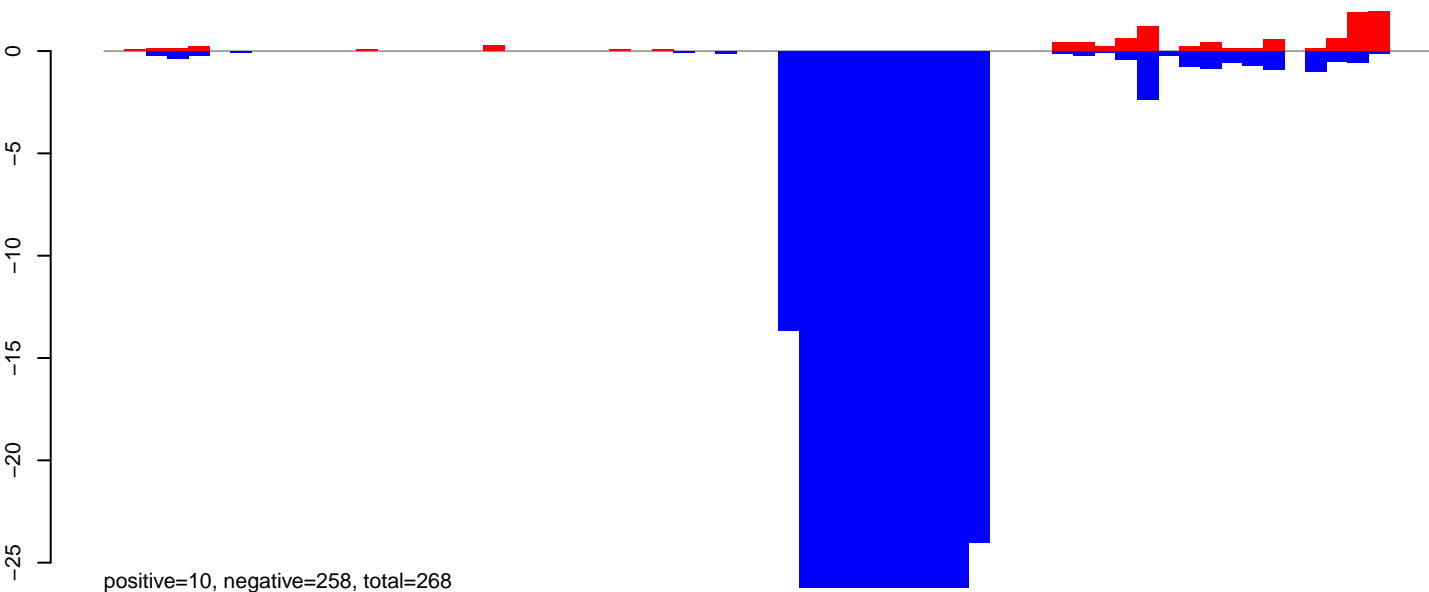


CuQuin_Mcarc_NL.rep

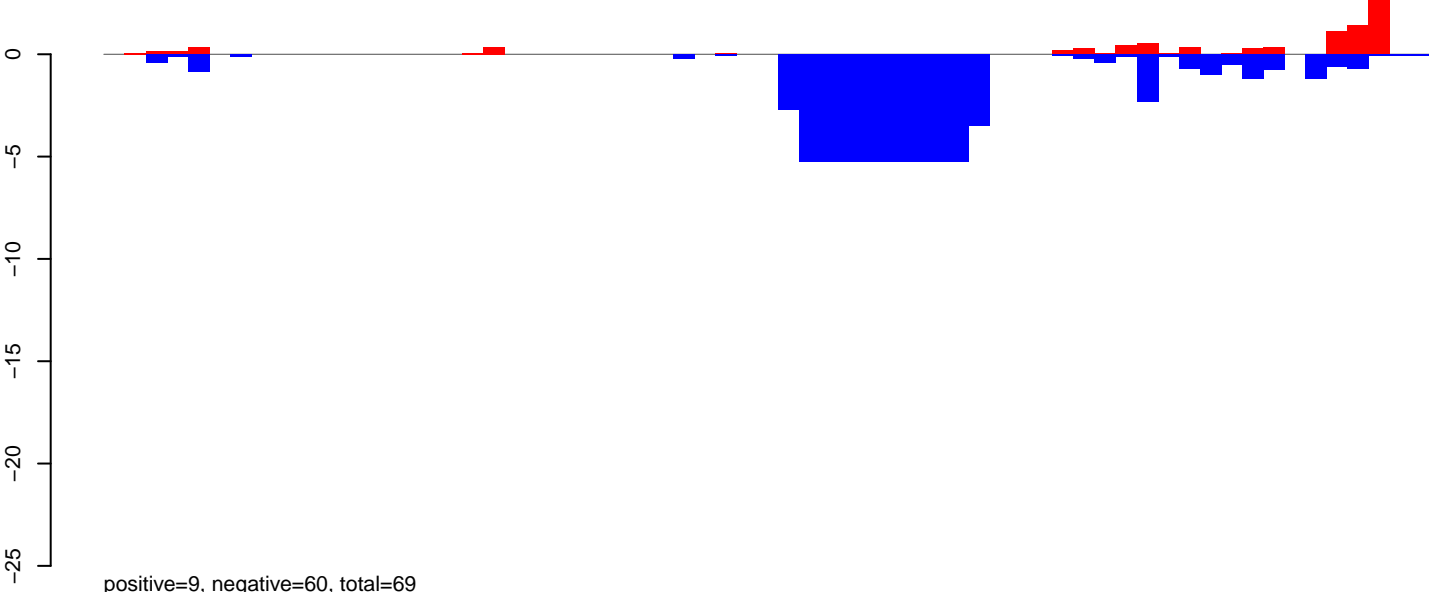


Window size=25, length=1149, TE@rnd-4_family-721@LINE_L1-Tx1@3542:1-1149

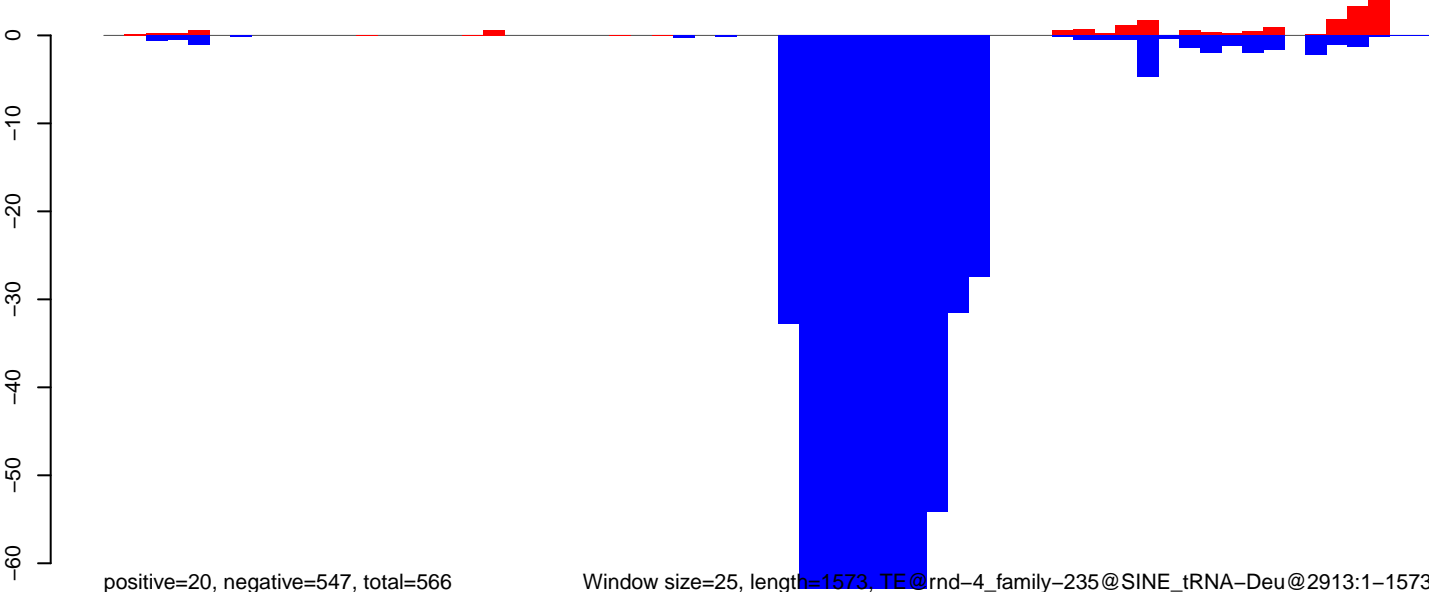
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



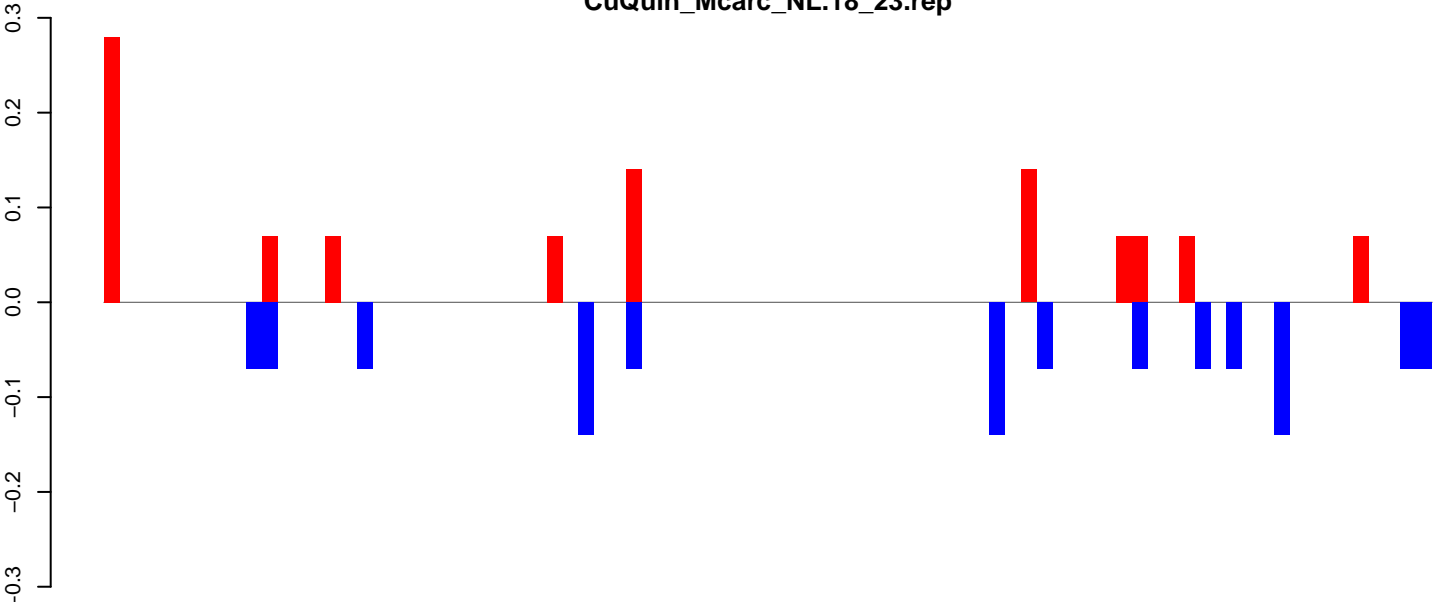
CuQuin_Mcarc_NL.rep



Window size=25, length=1573, TE@rnd-4_family-235@SINE_tRNA-Deu@2913:1-1573

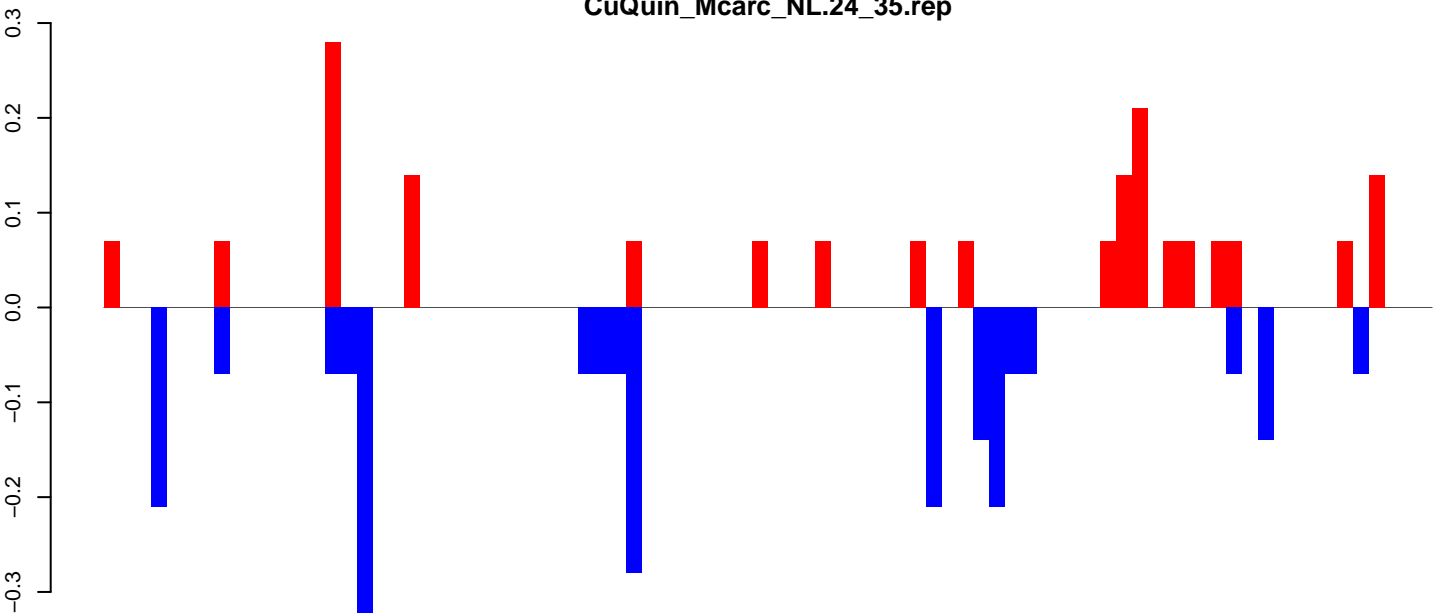
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CuQuin_Mcarc_NL.18_23.rep



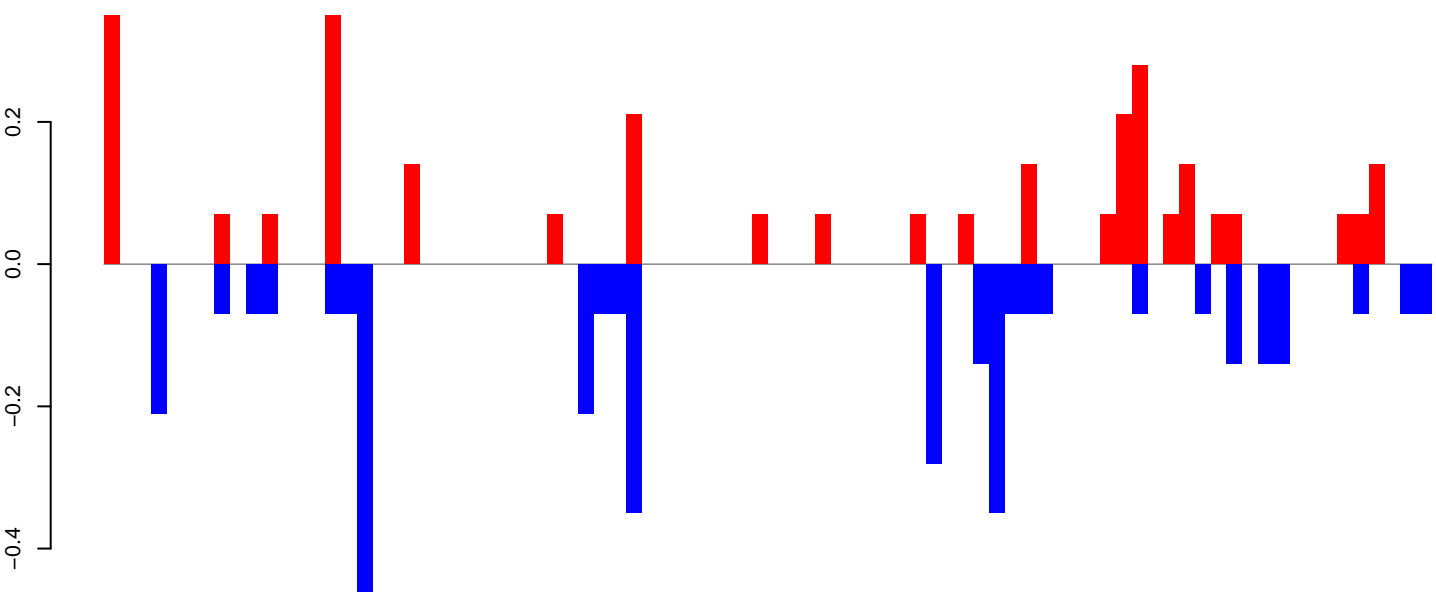
positive=1, negative=1, total=2

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=2, total=4

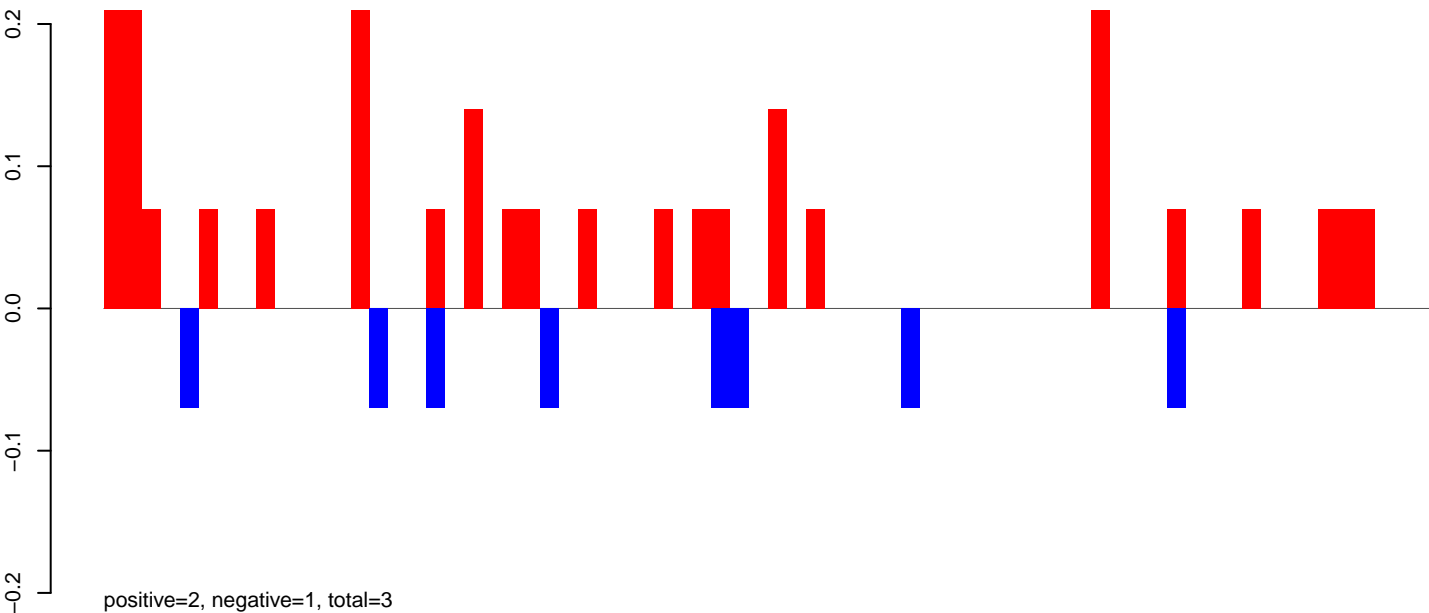
CuQuin_Mcarc_NL.rep



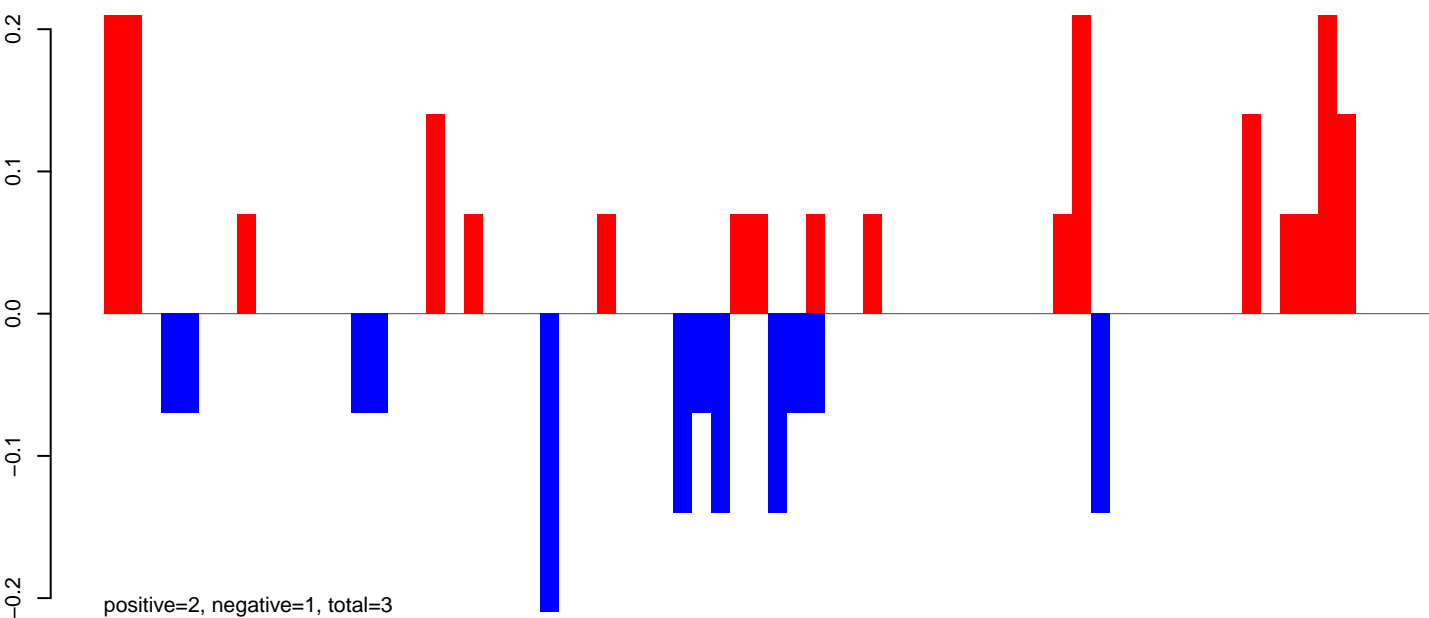
positive=3, negative=3, total=6

Window size=25, length=2088, TE@rnd-4_family-135@DNA_Zator@173:1-2088

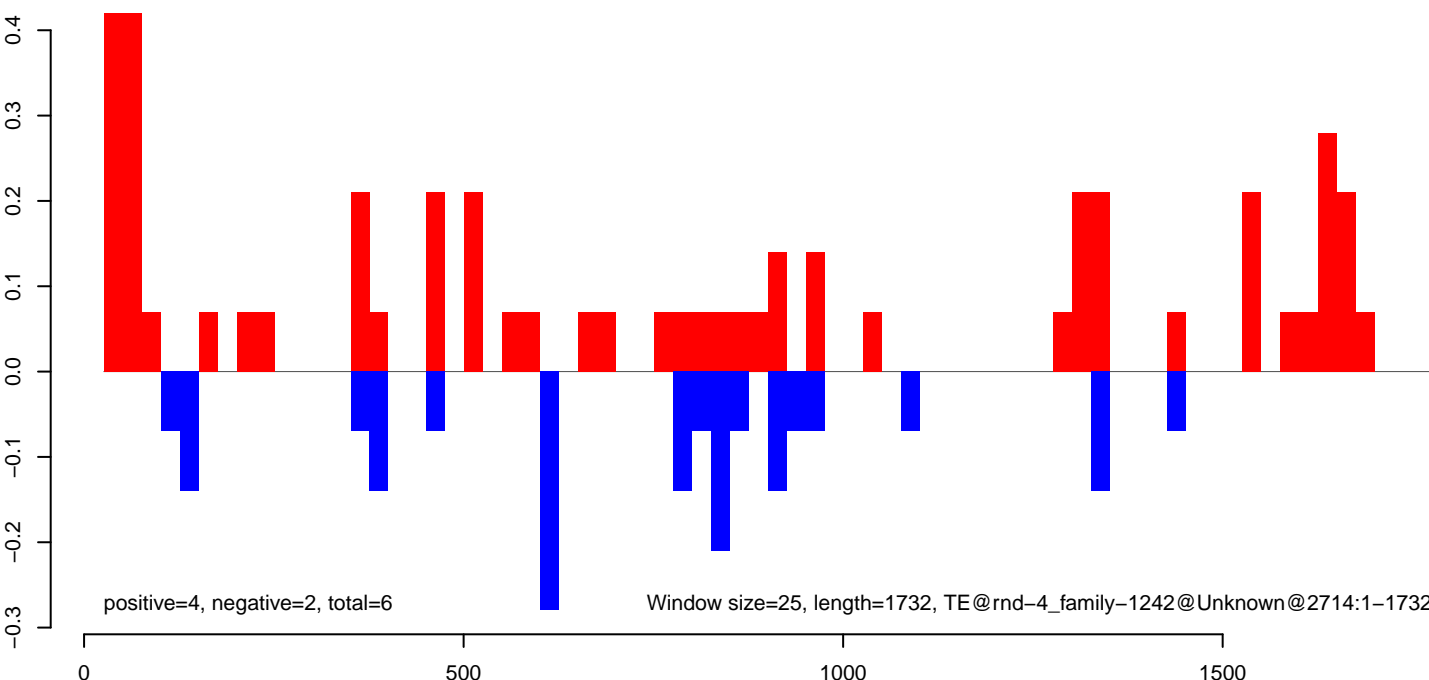
CuQuin_Mcarc_NL.18_23.rep



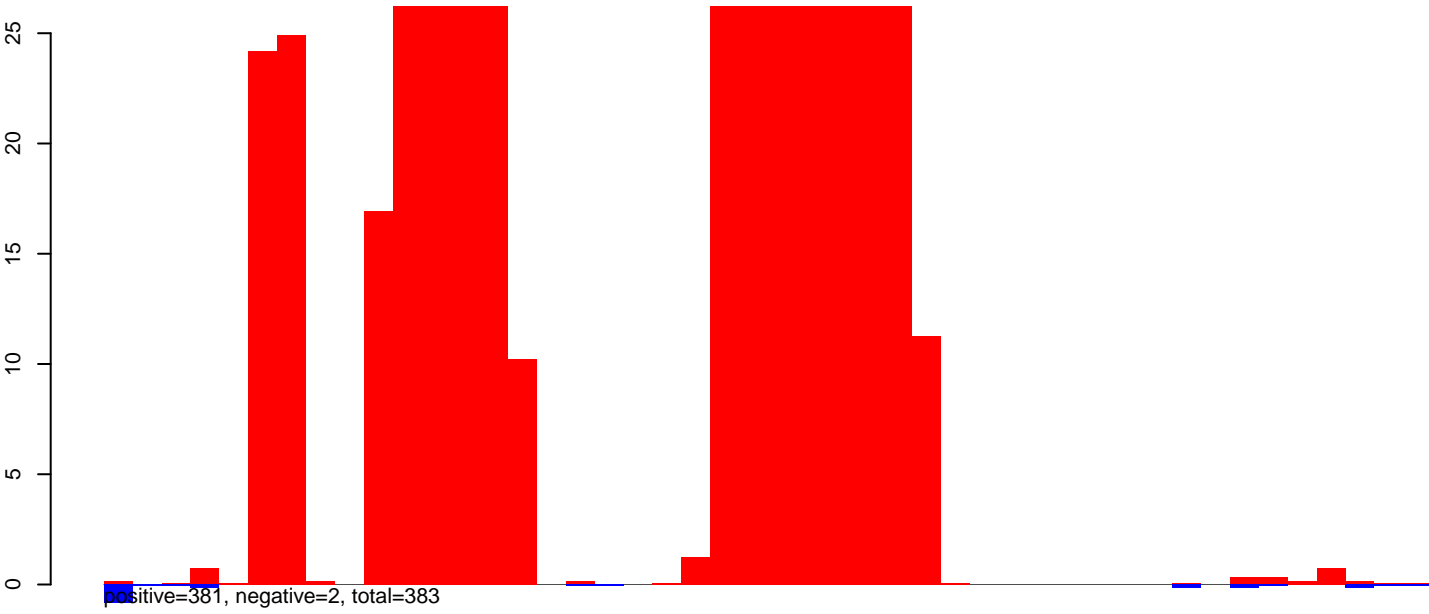
CuQuin_Mcarc_NL.24_35.rep



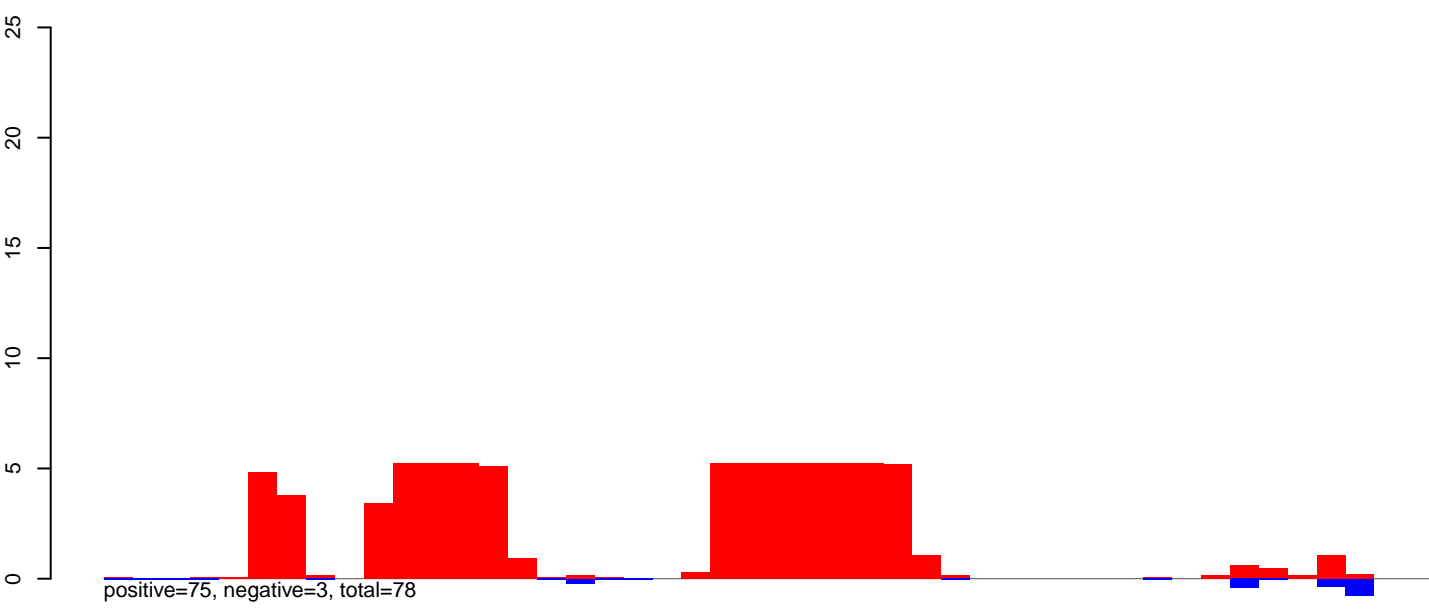
CuQuin_Mcarc_NL.rep



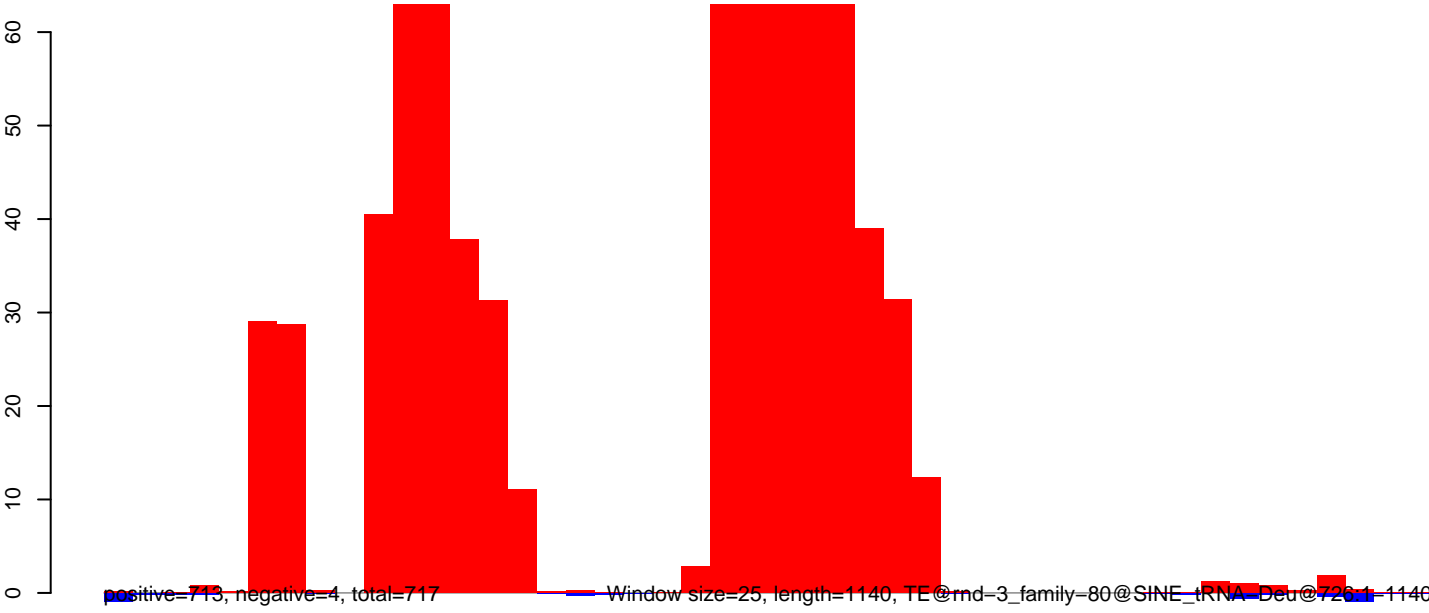
CuQuin_Mcarc_NL.18_23.rep



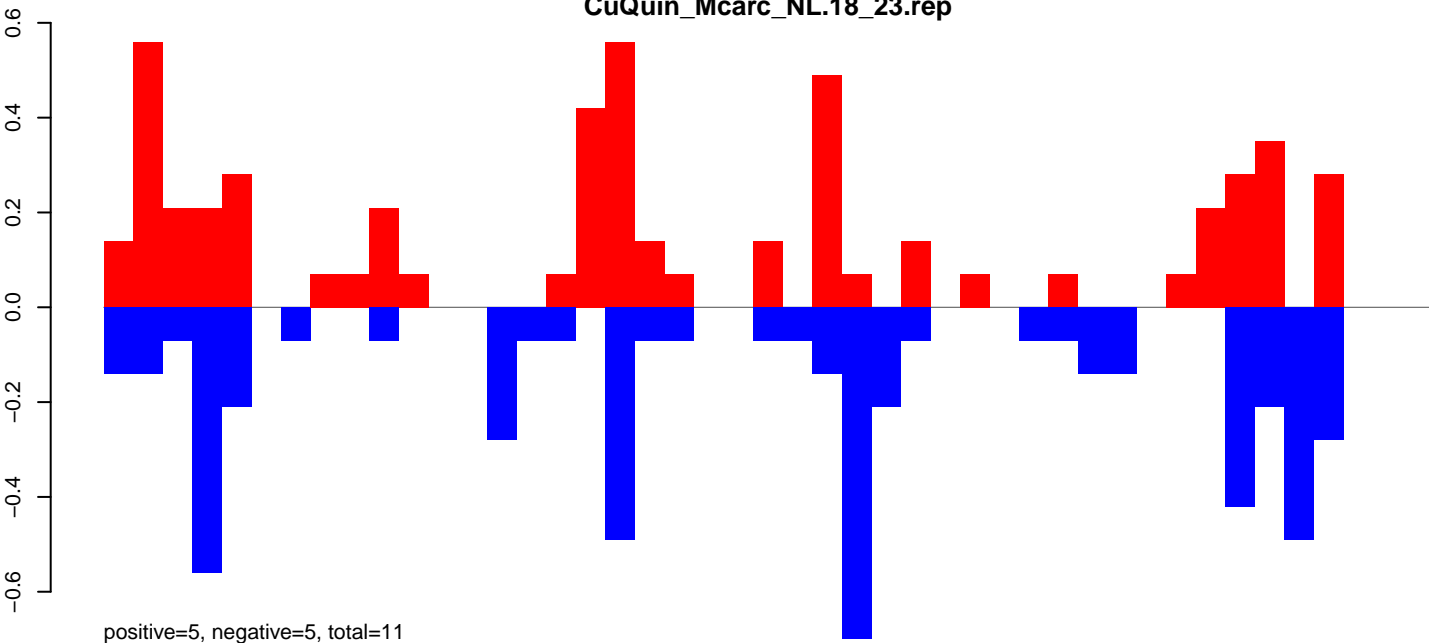
CuQuin_Mcarc_NL.24_35.rep



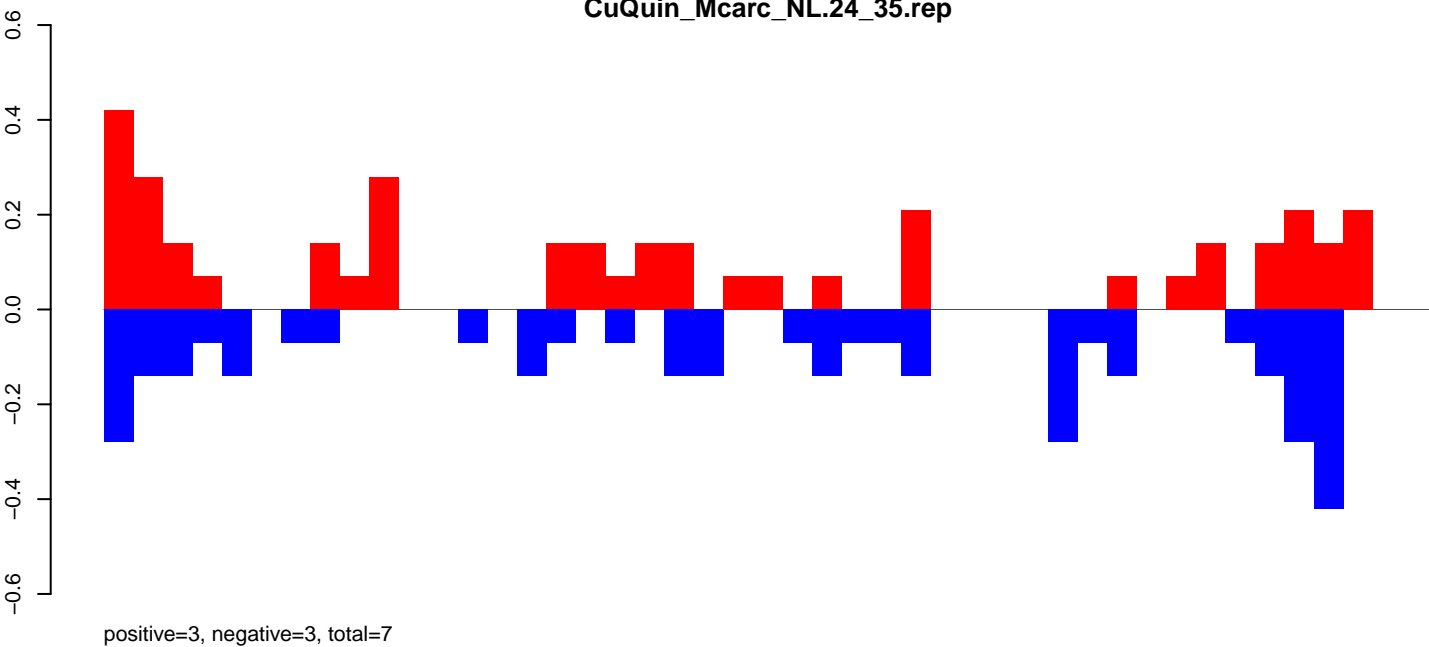
CuQuin_Mcarc_NL.rep



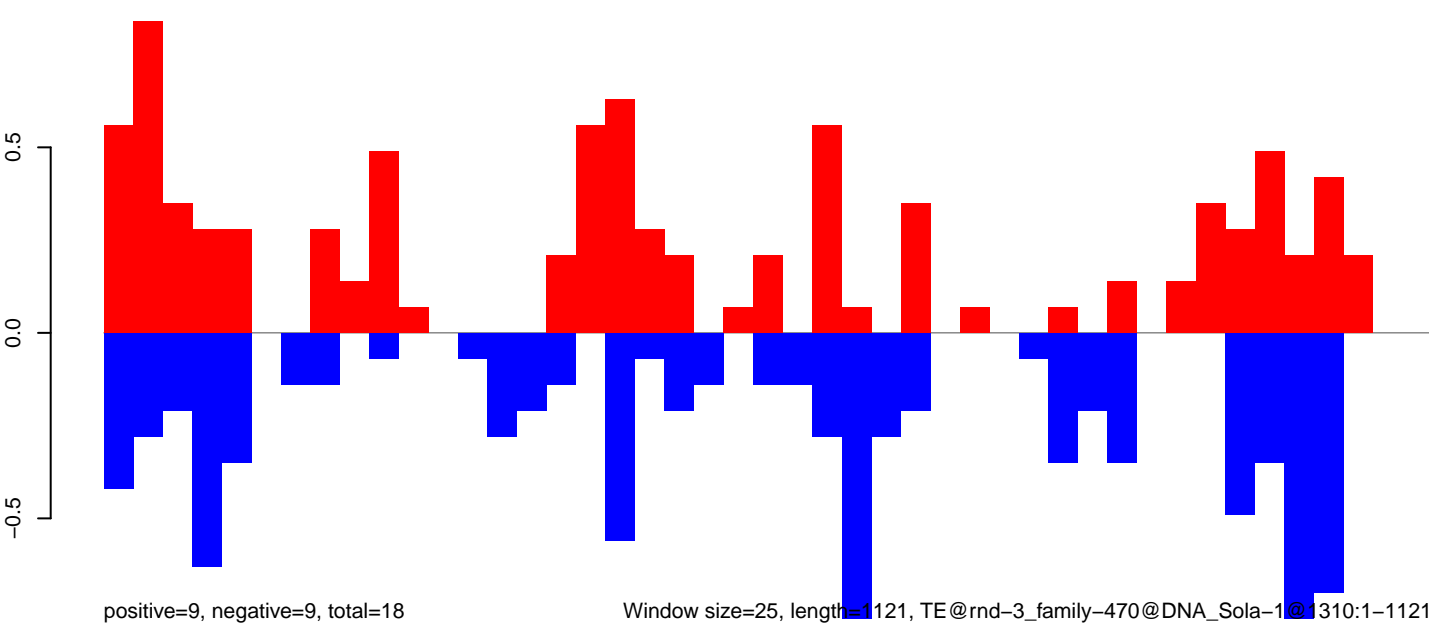
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

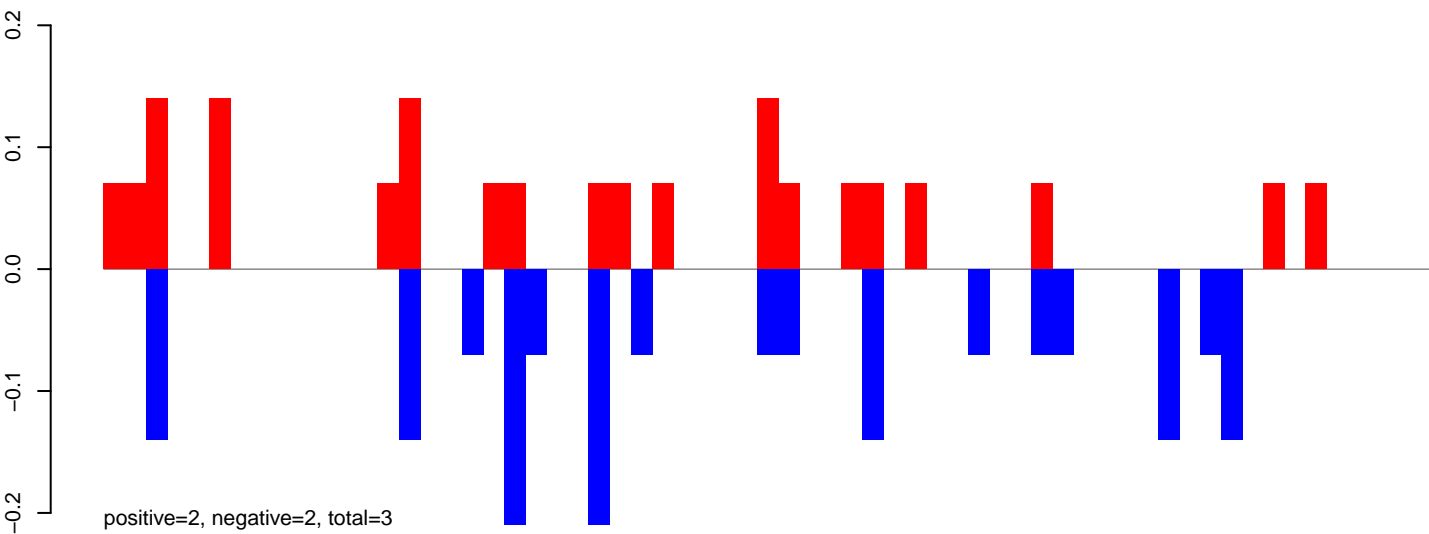


CuQuin_Mcarc_NL.rep

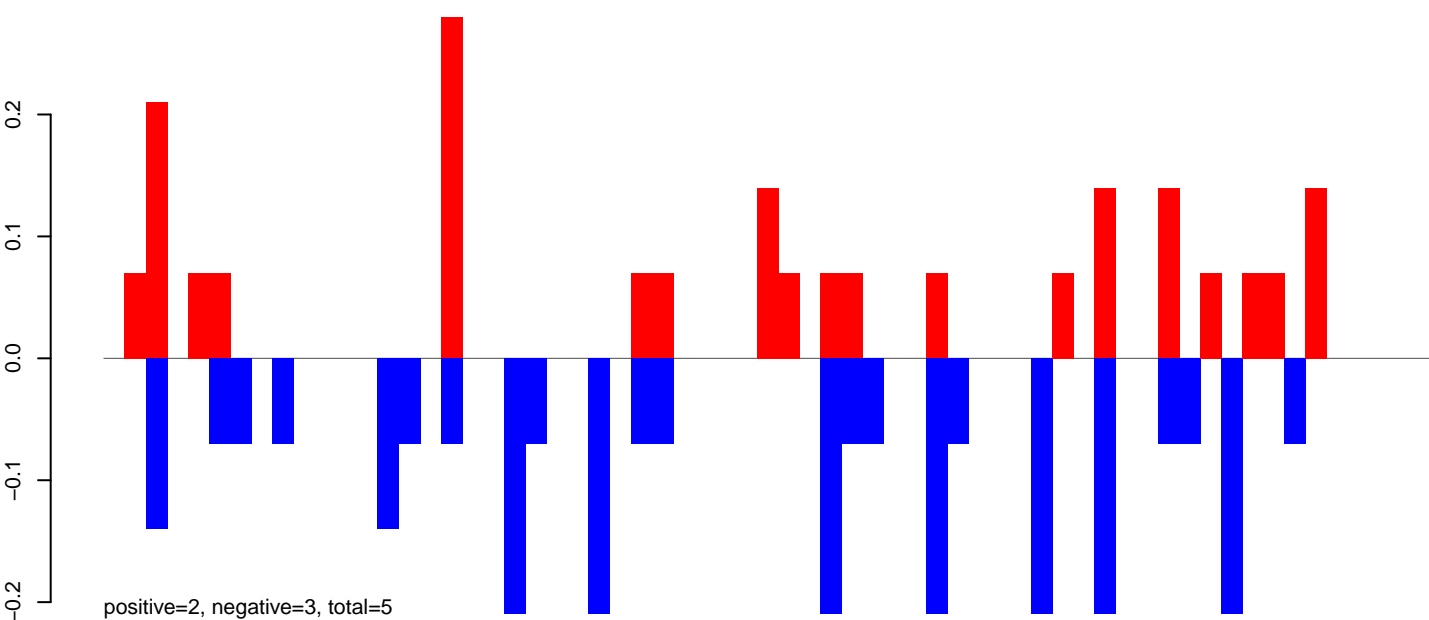


Window size=25, length=1121, TE@rnd-3_family-470@DNA_Sola-1@1310:1-1121

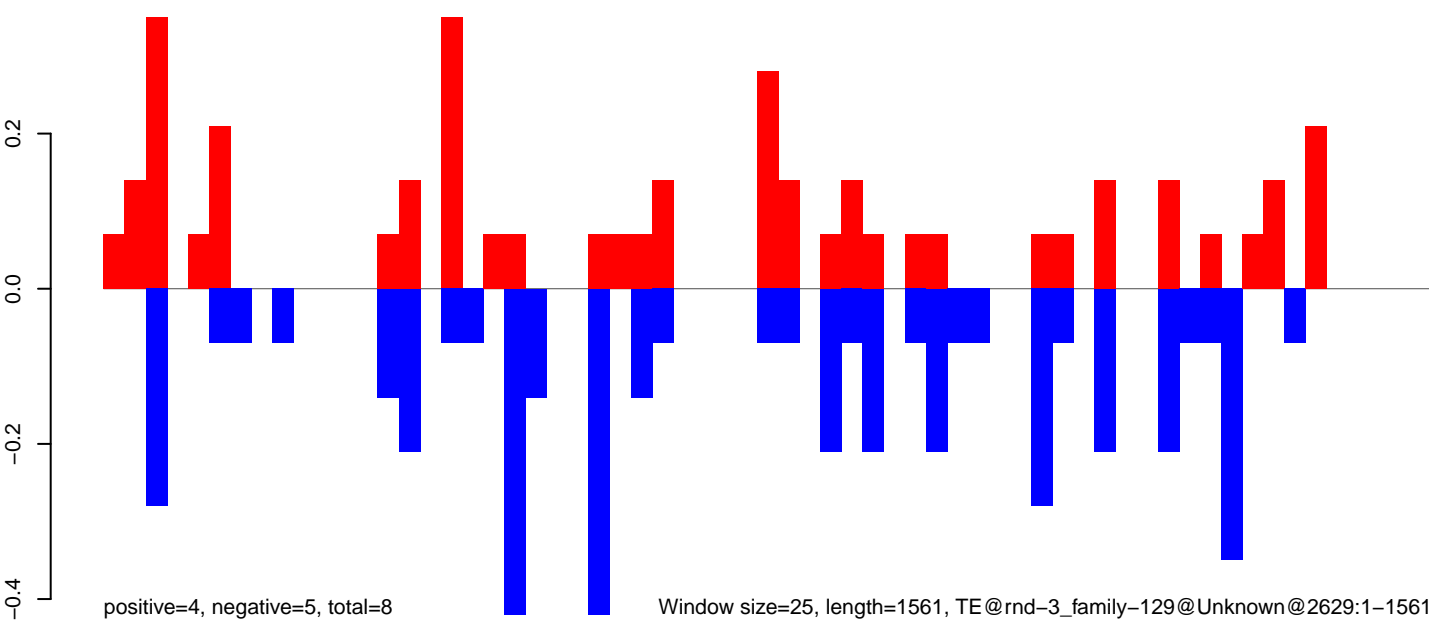
CuQuin_Mcarc_NL.18_23.rep



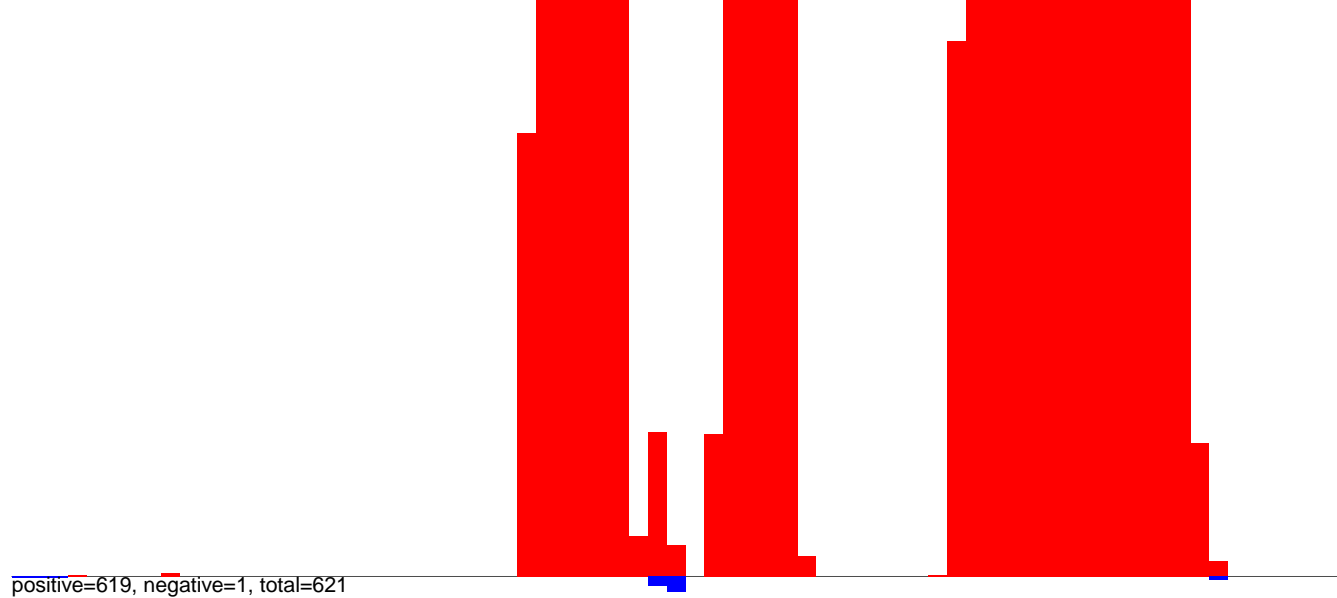
CuQuin_Mcarc_NL.24_35.rep



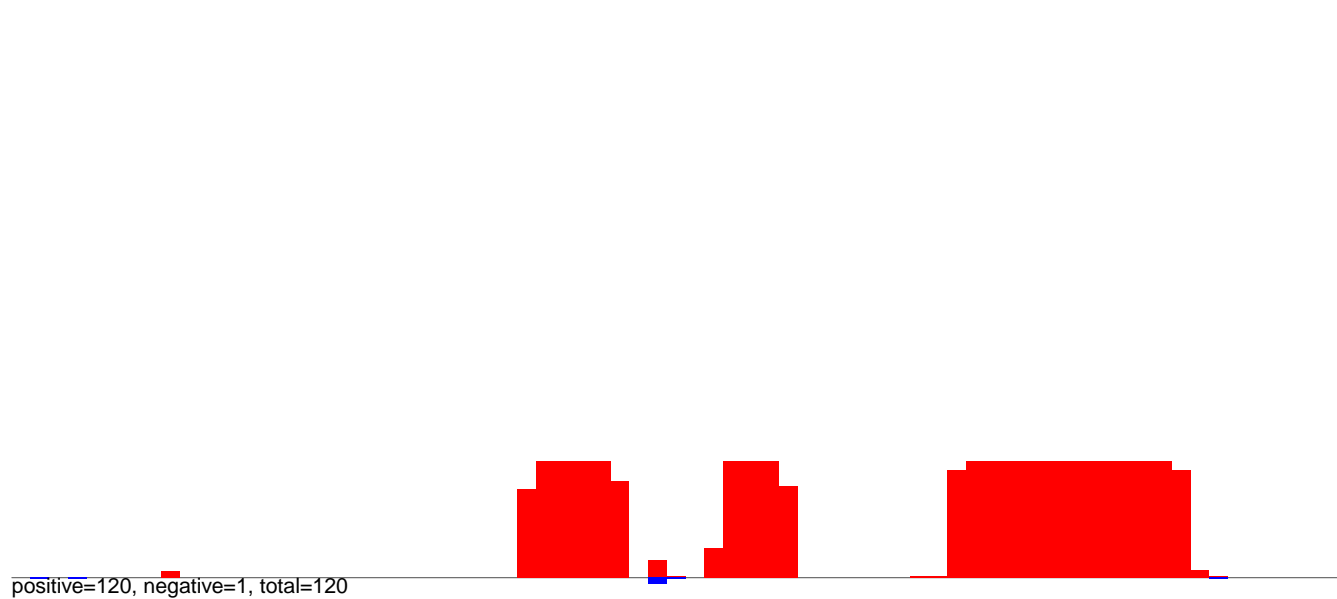
CuQuin_Mcarc_NL.rep



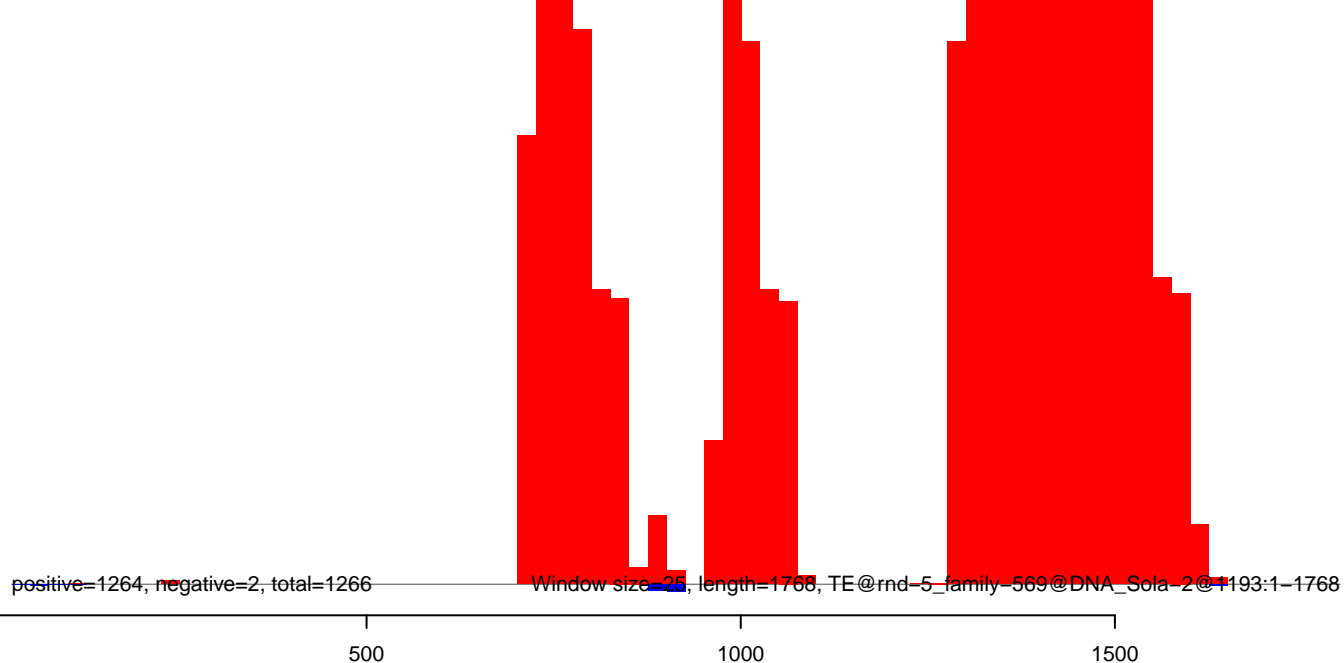
CuQuin_Mcarc_NL.18_23.rep



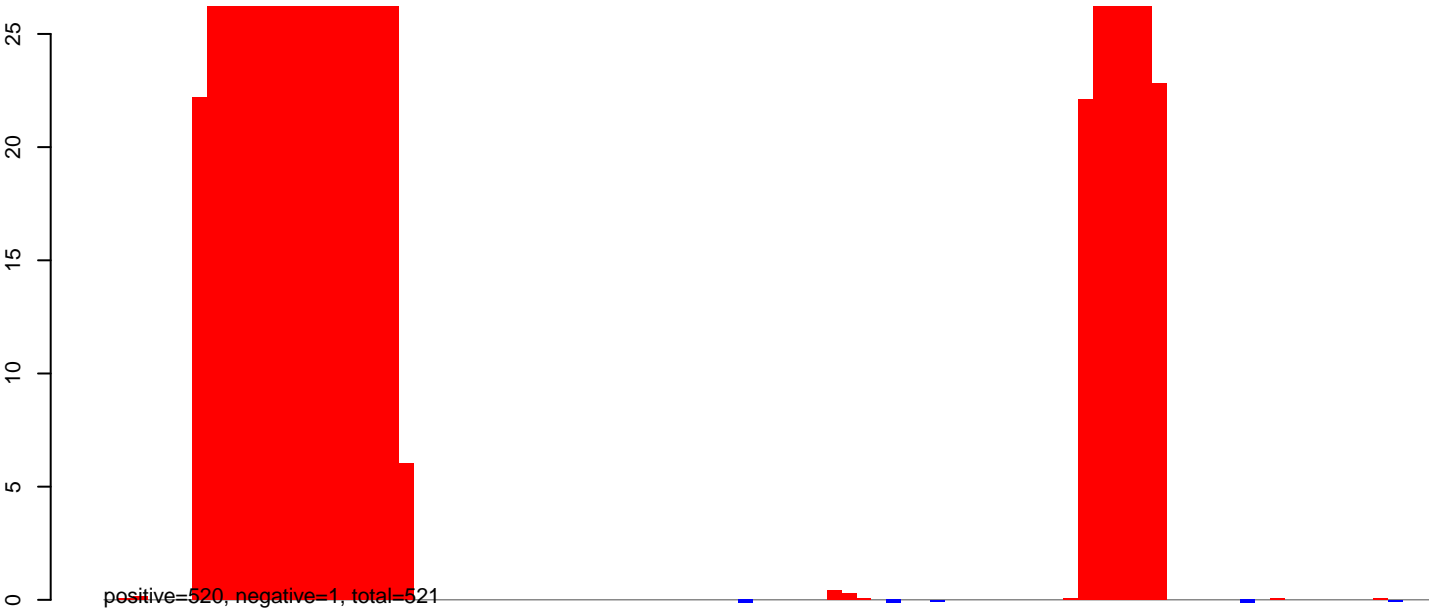
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CuQuin_Mcarc_NL.rep



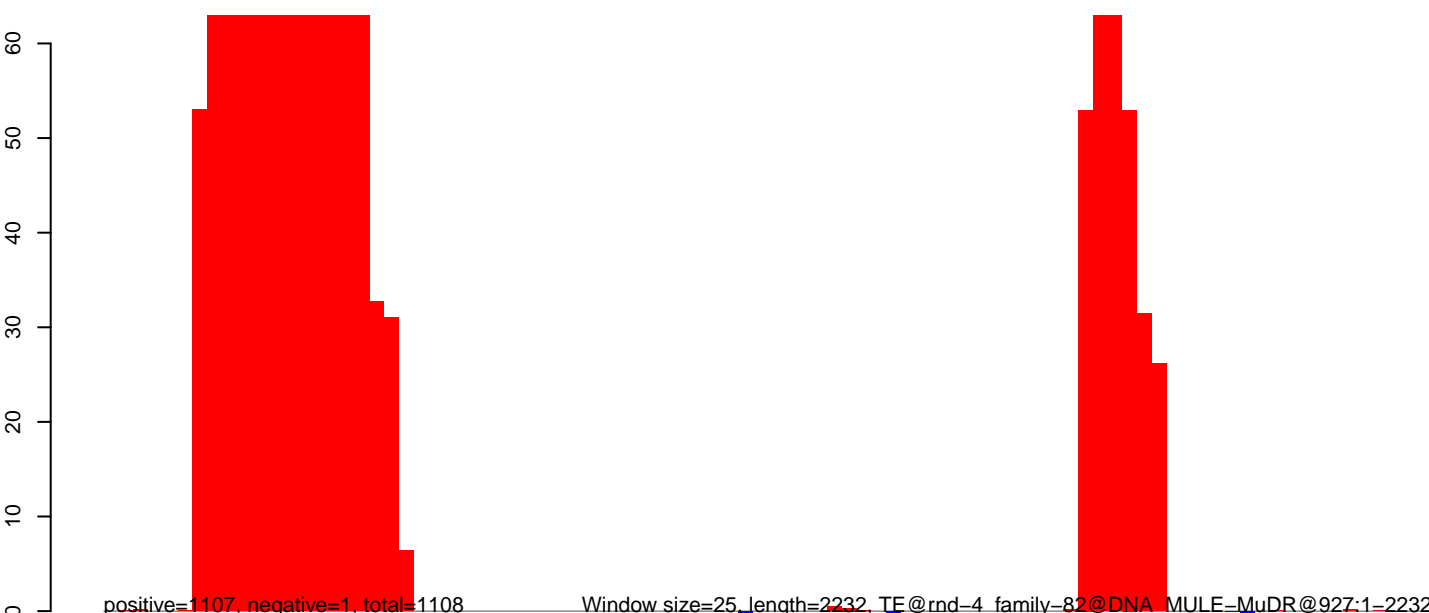
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CuQuin_Mcarc_NL.24_35.rep

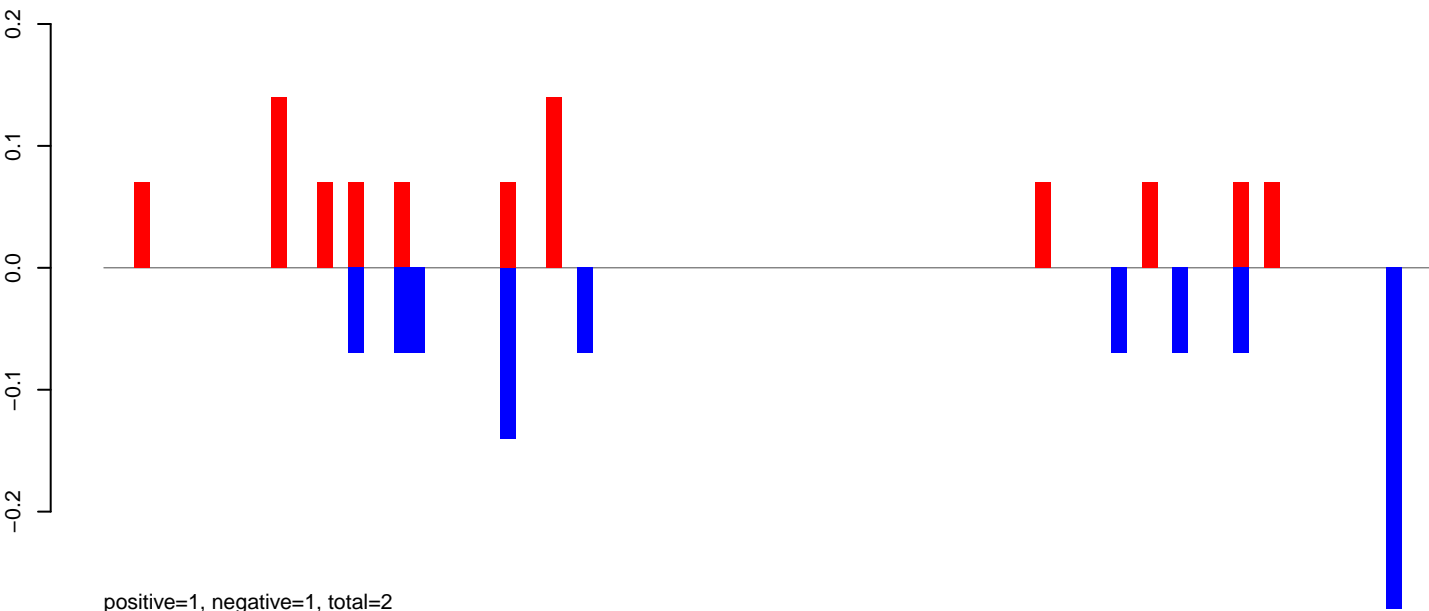


CuQuin_Mcarc_NL.rep

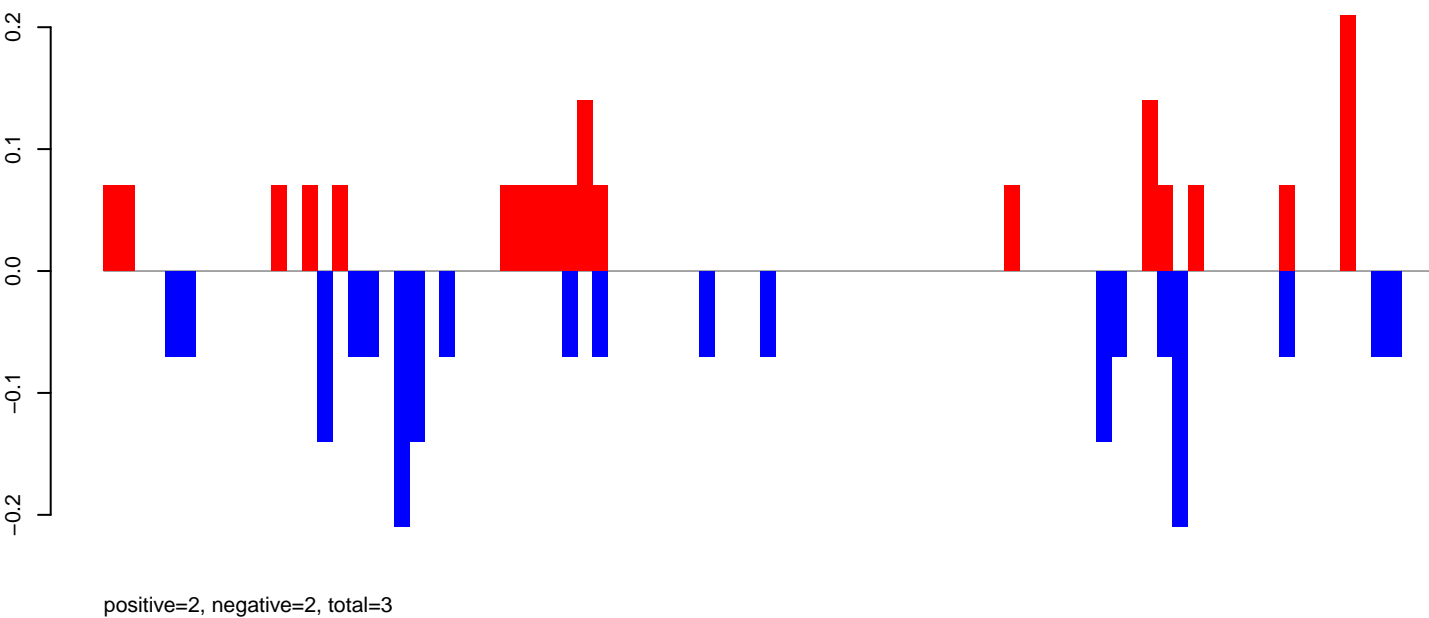


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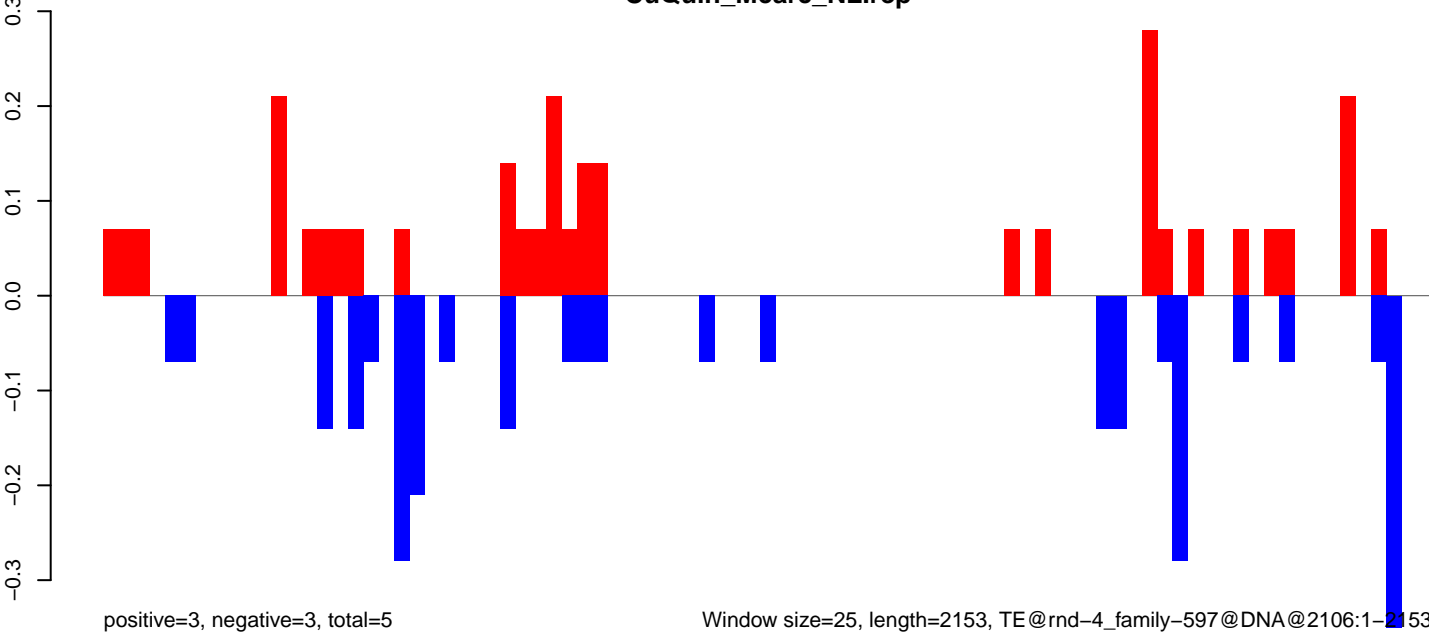
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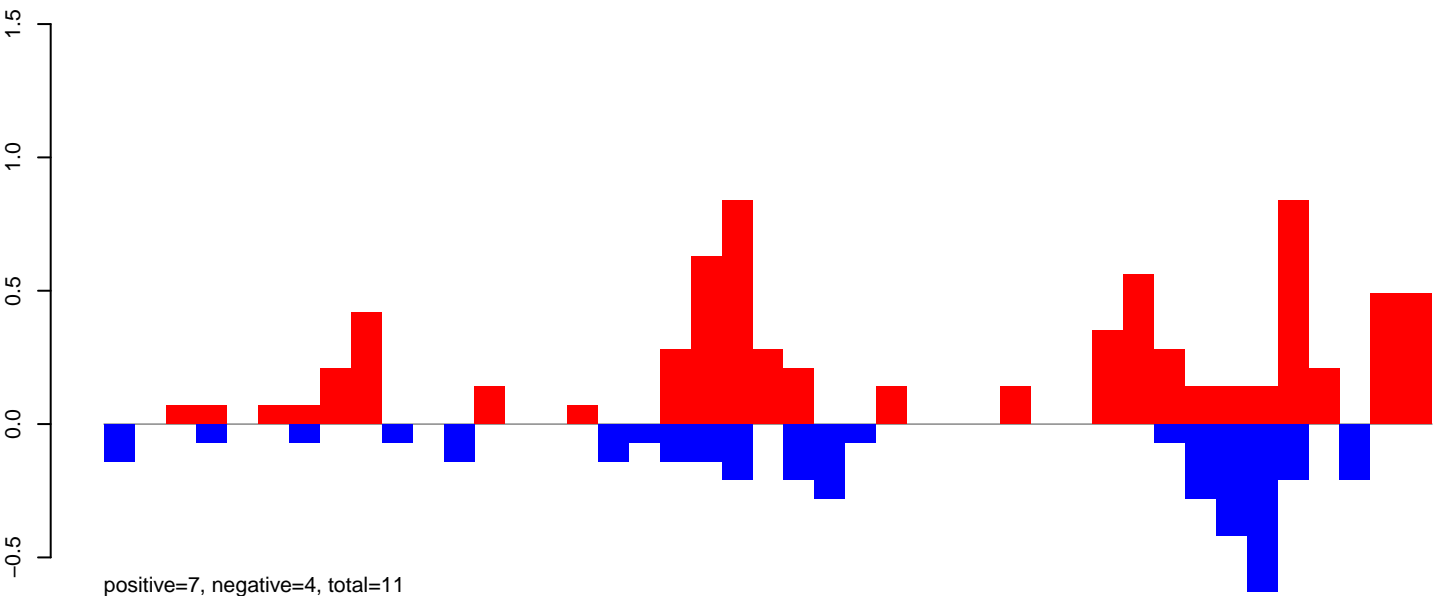
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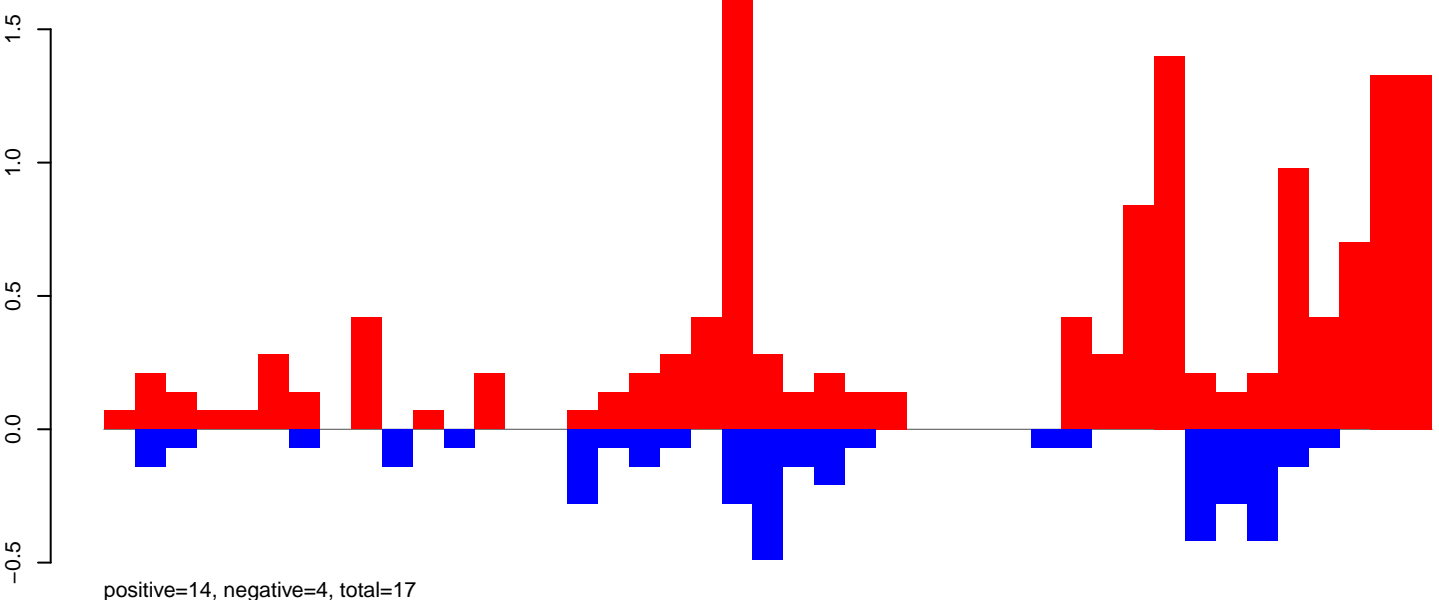
CuQuin_Mcarc_NL.rep



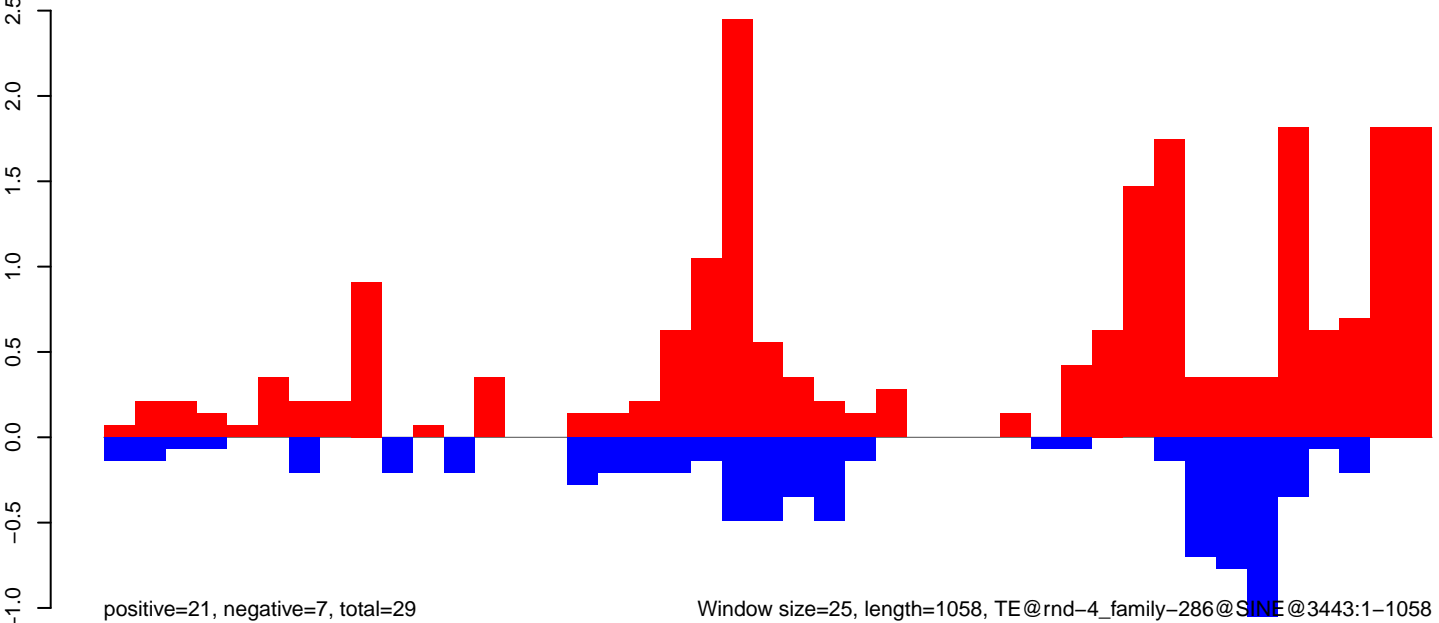
CuQuin_Mcarc_NL.18_23.rep



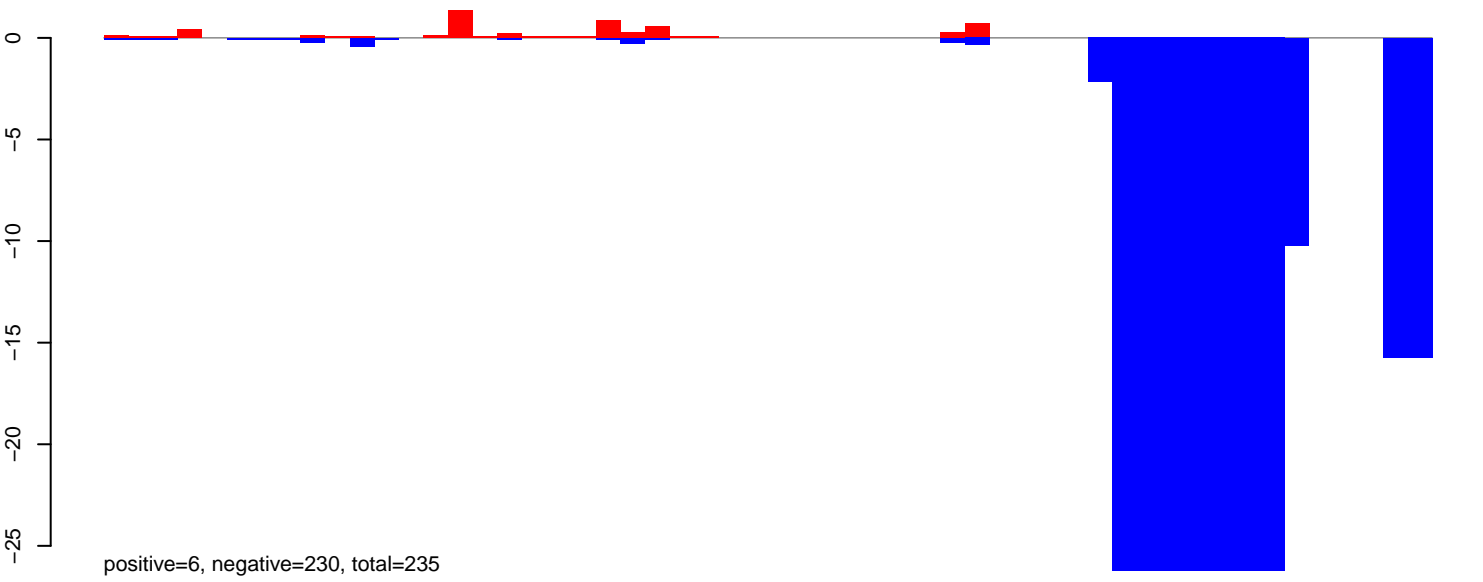
CuQuin_Mcarc_NL.24_35.rep



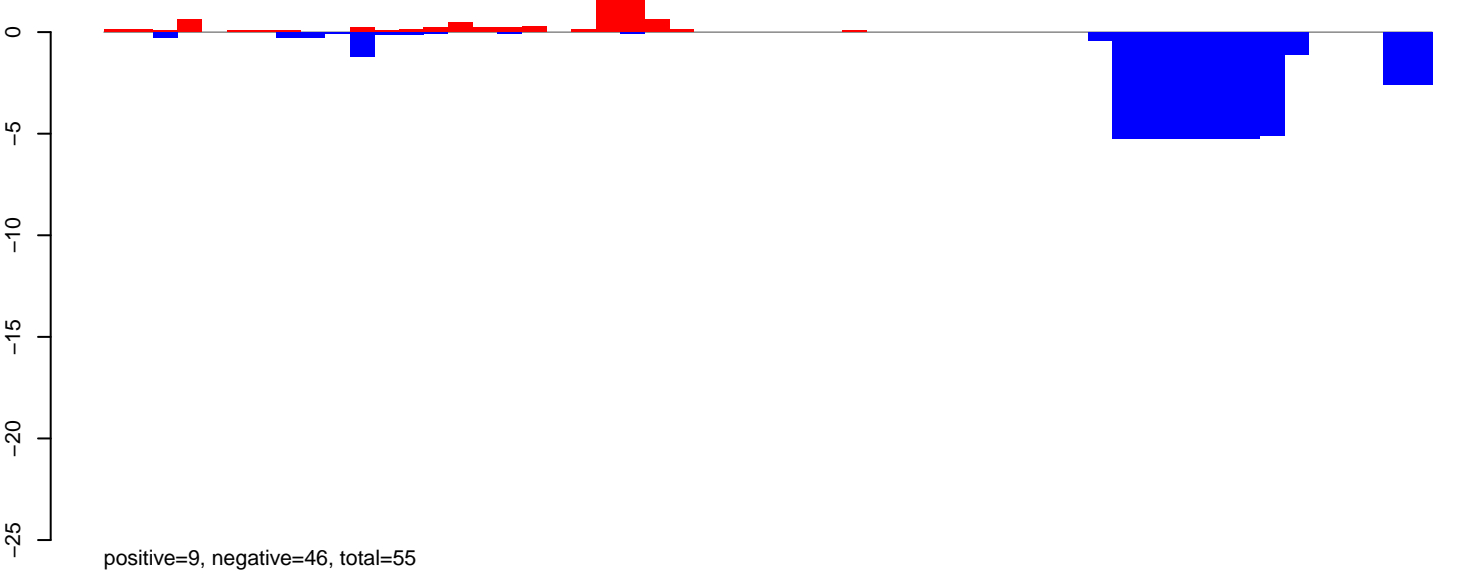
CuQuin_Mcarc_NL.rep



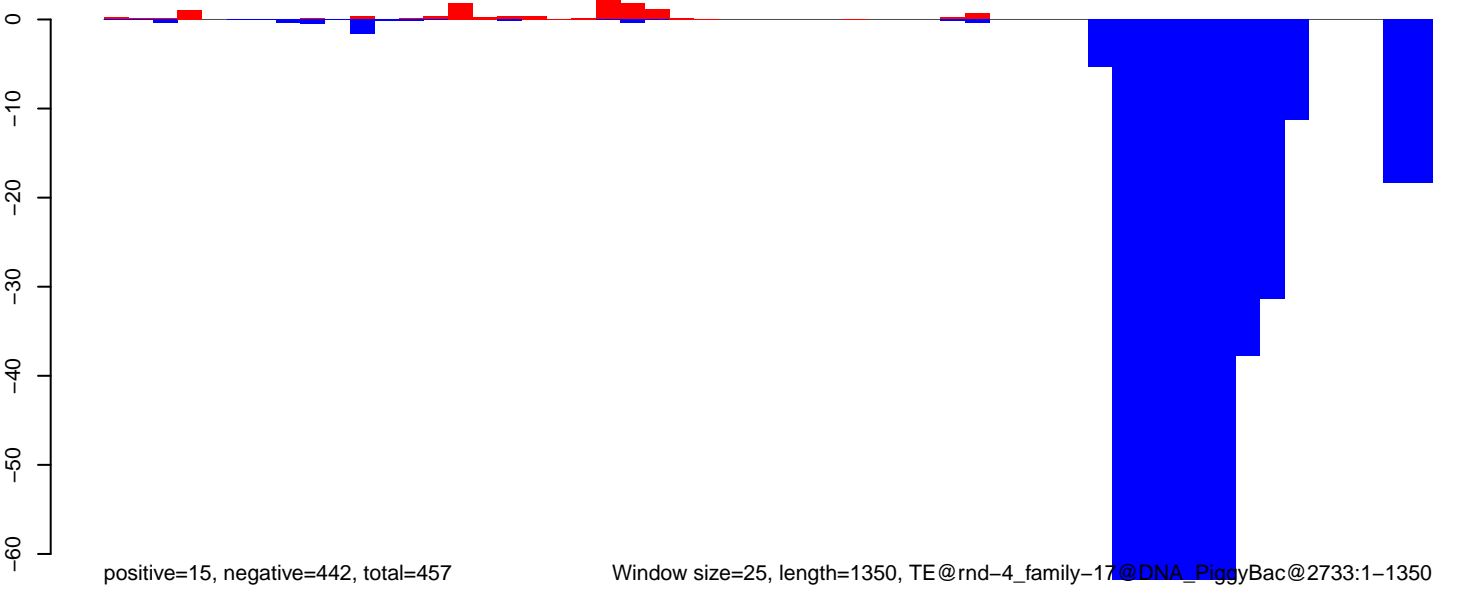
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

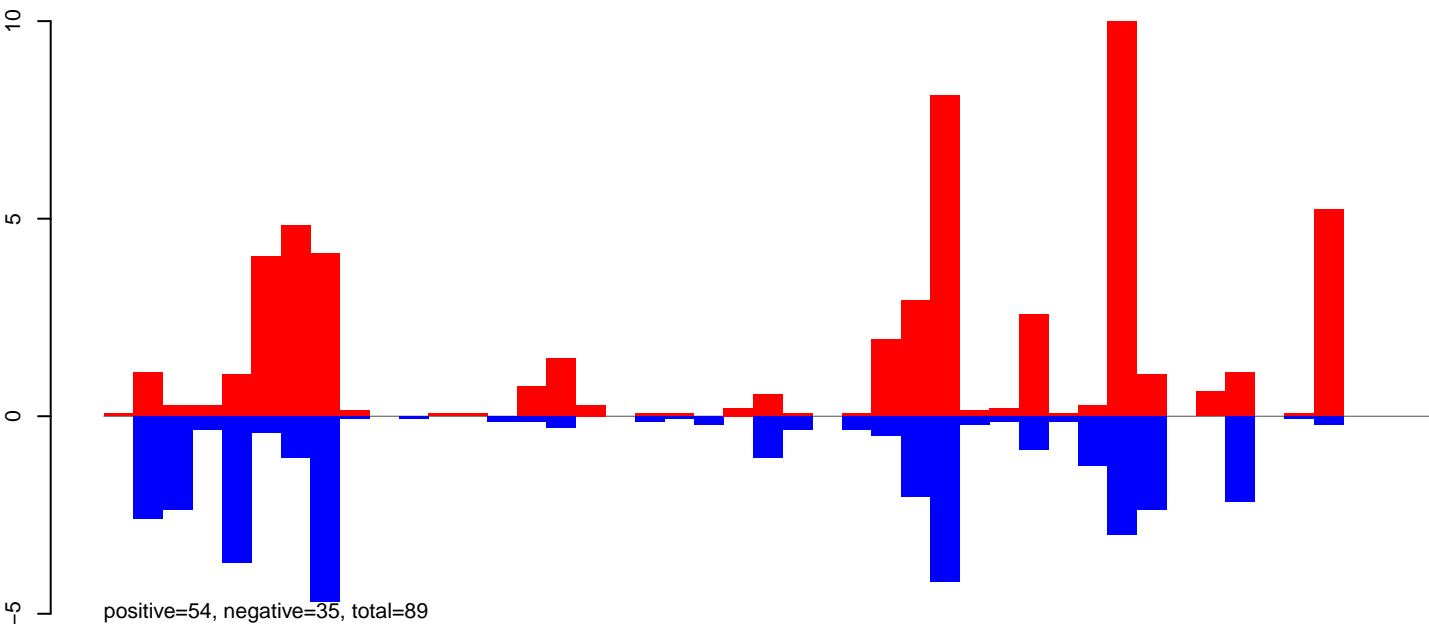


CuQuin_Mcarc_NL.rep

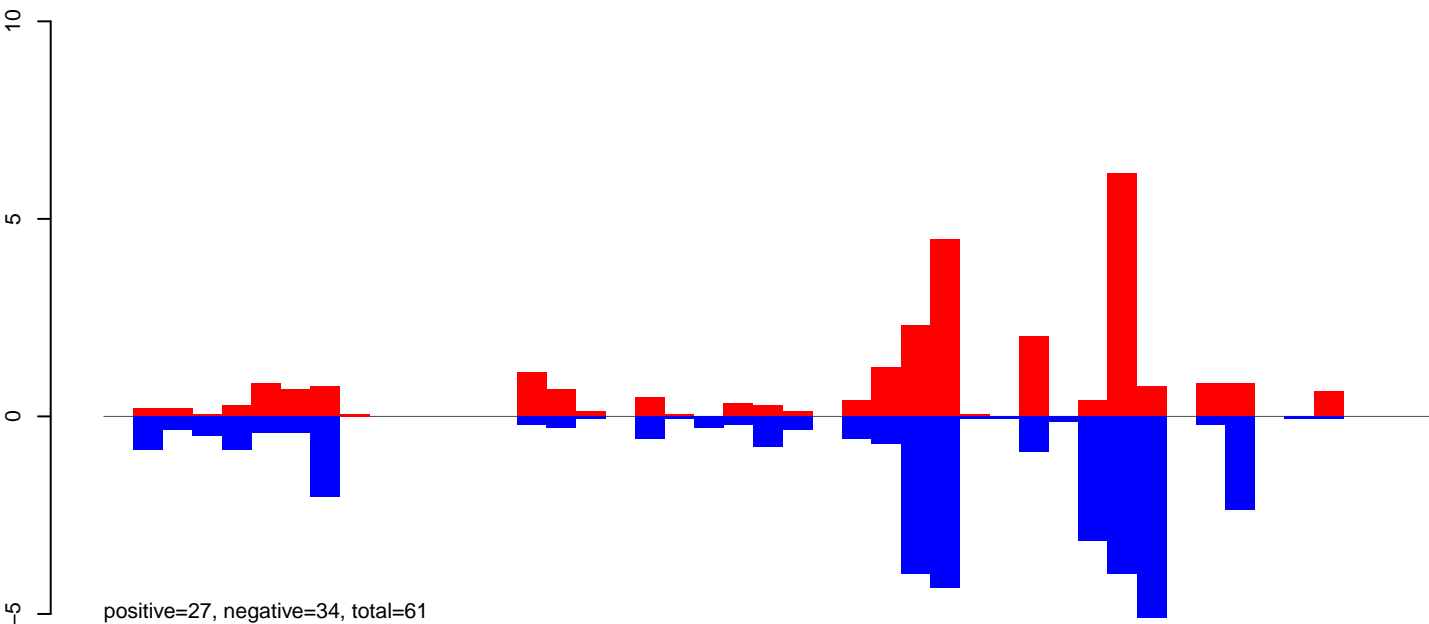


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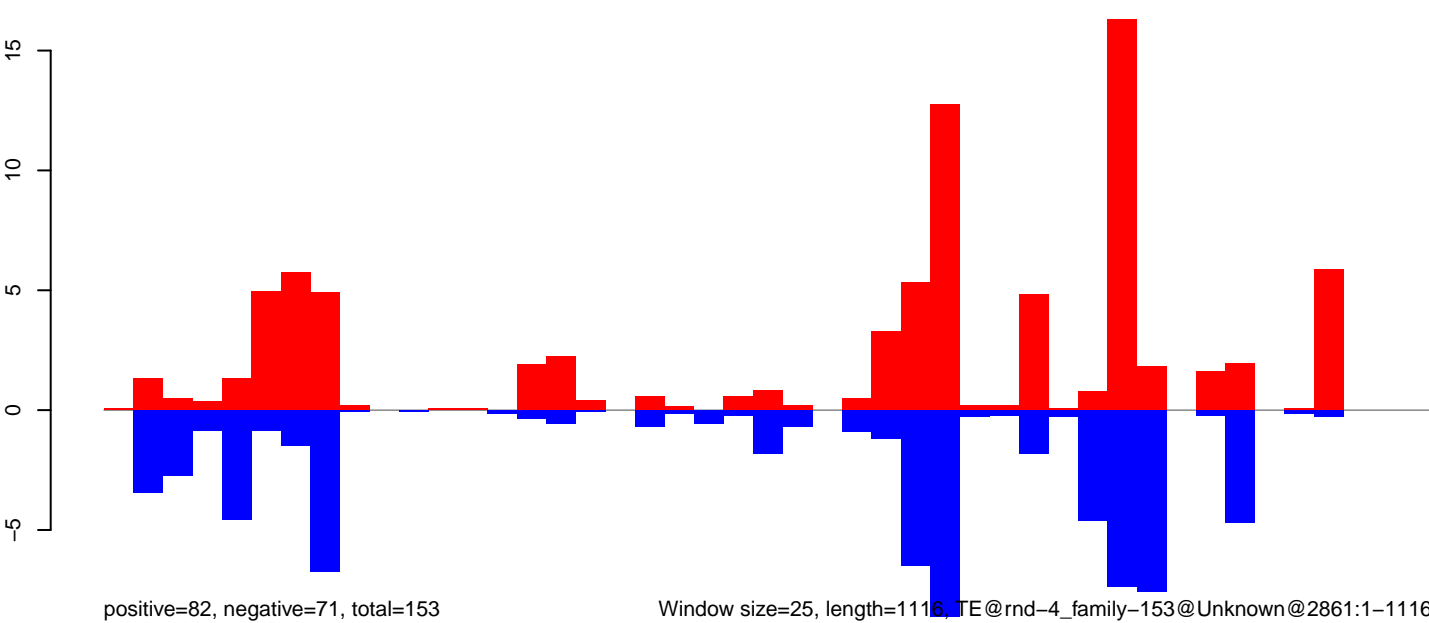
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



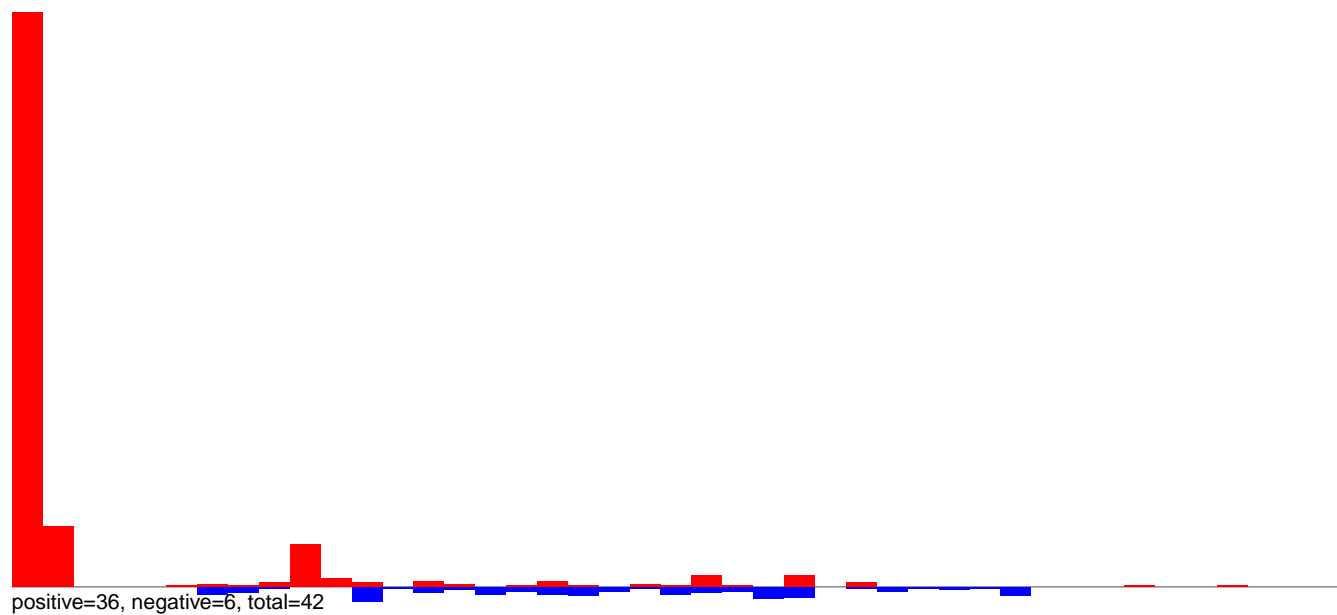
CuQuin_Mcarc_NL.rep



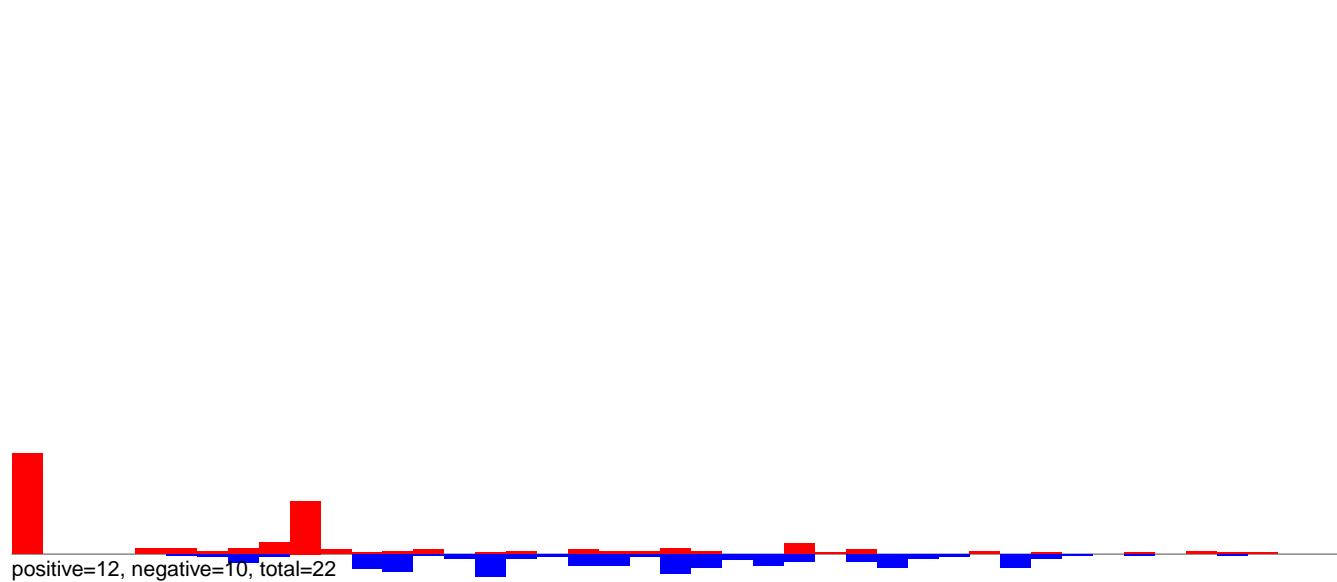
Window size=25, length=1116, TE@rnd-4_family-153@Unknown@2861:1-1116

0 200 400 600 800 1000

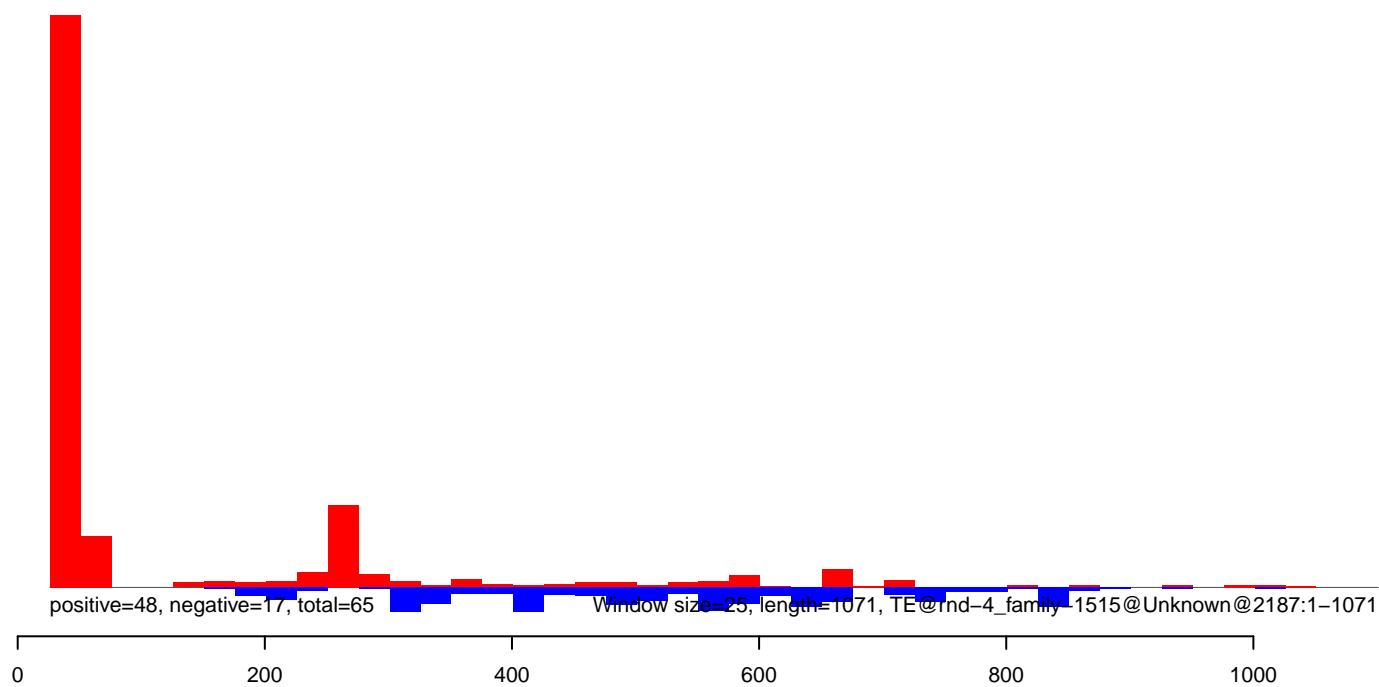
CuQuin_Mcarc_NL.18_23.rep



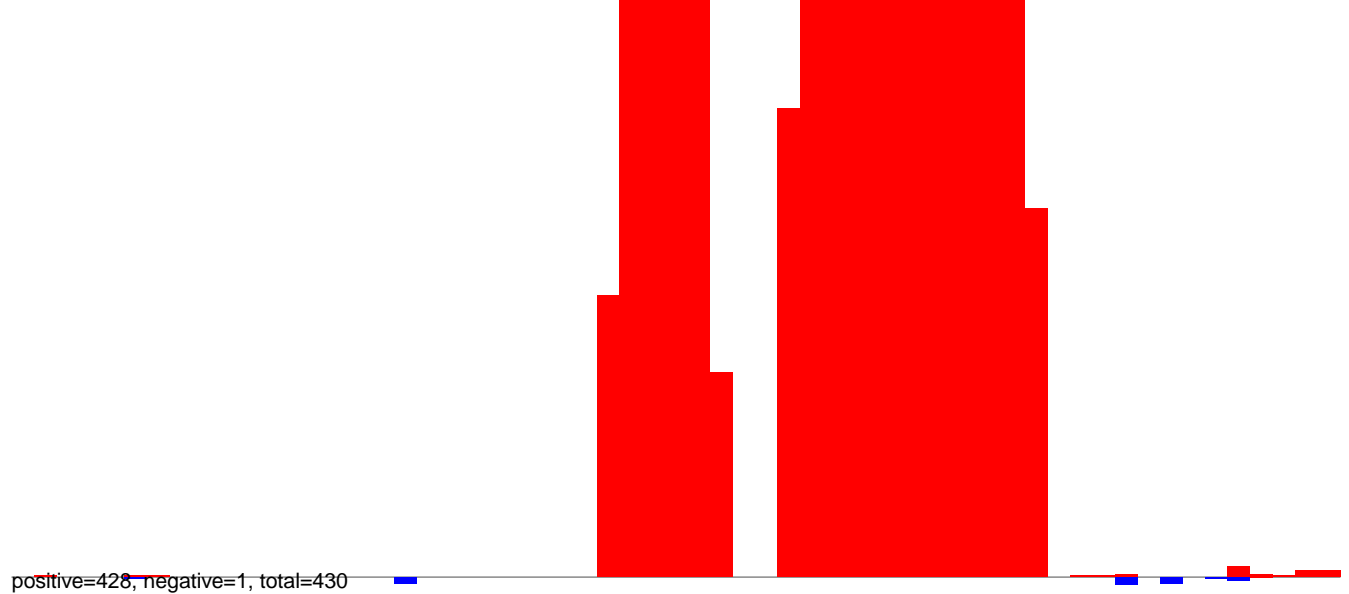
CuQuin_Mcarc_NL.24_35.rep



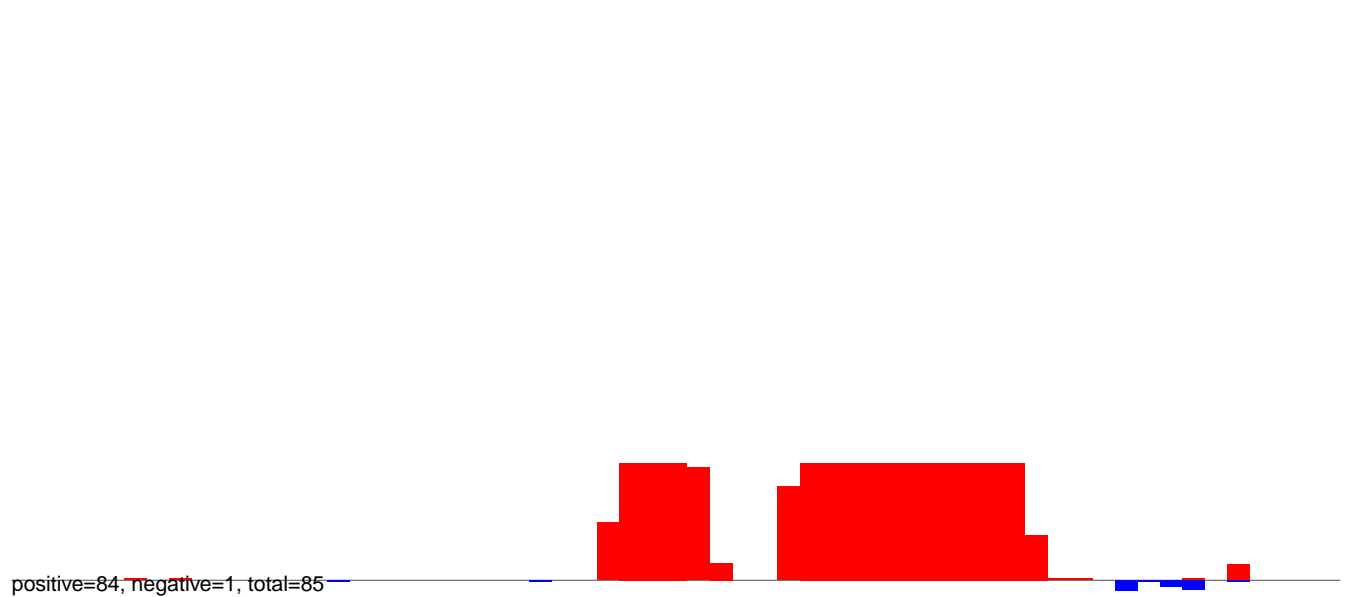
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CuQuin_Mcarc_NL.18_23.rep



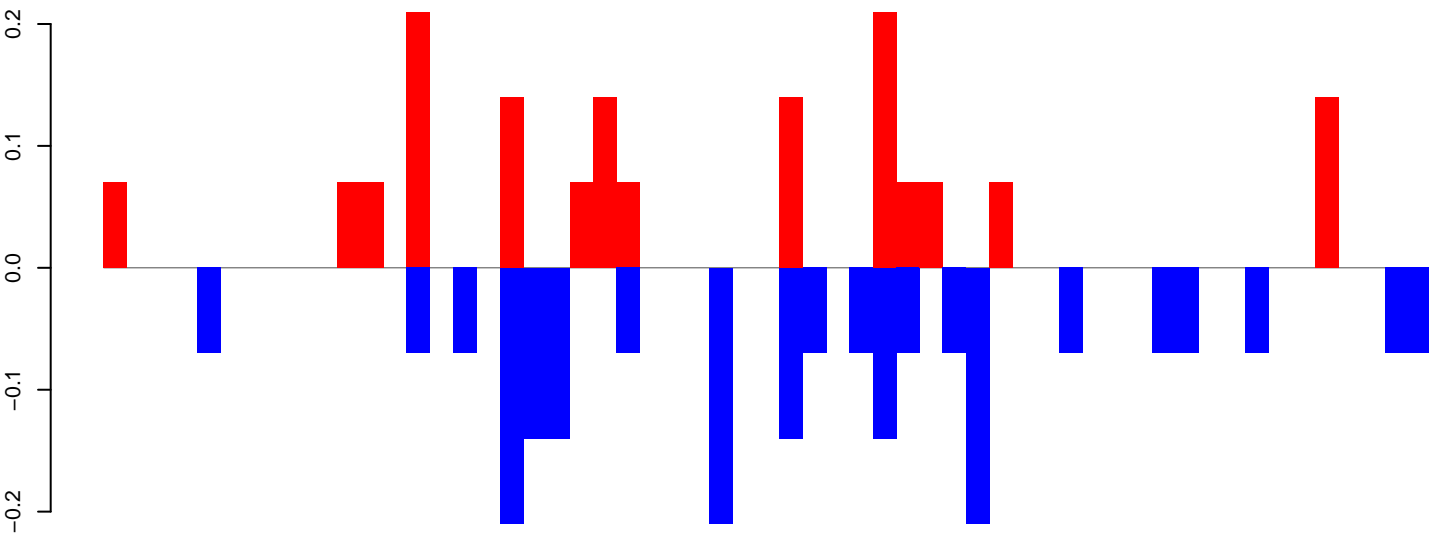
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CuQuin_Mcarc_NL.rep

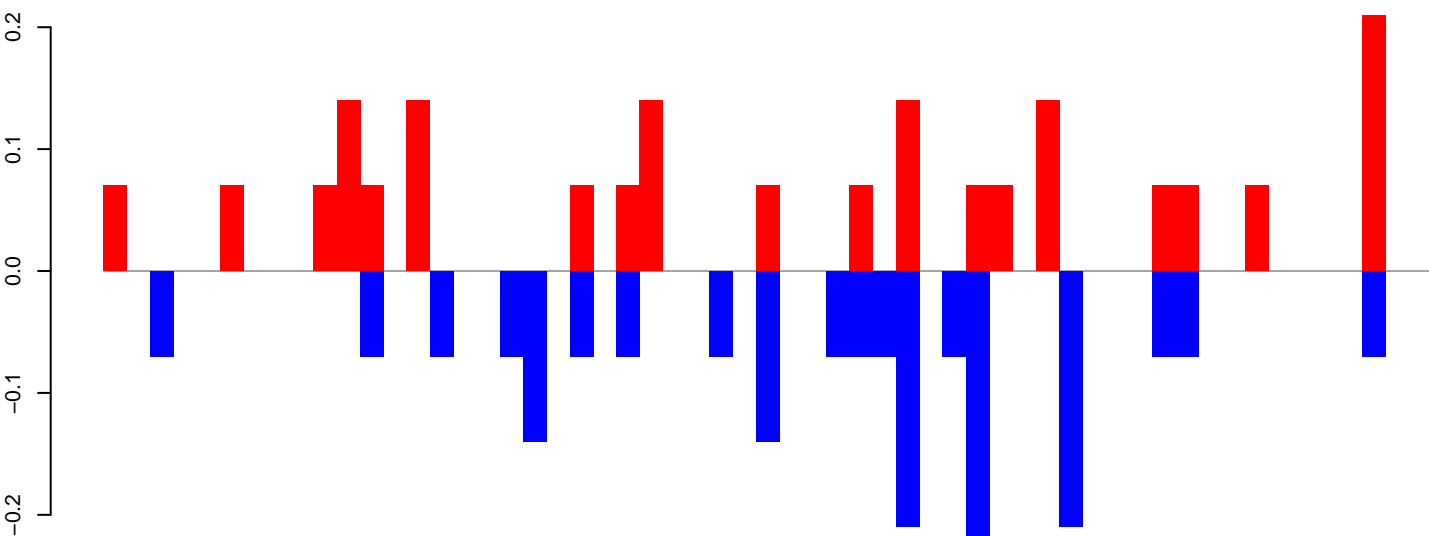


CuQuin_Mcarc_NL.18_23.rep



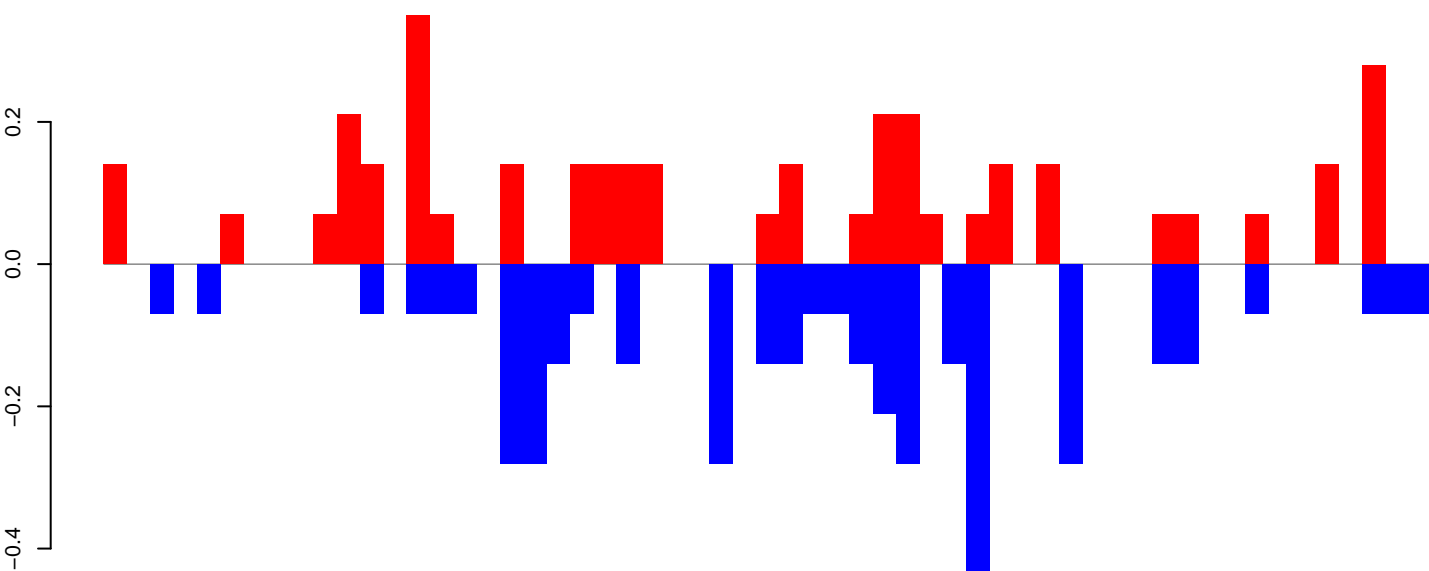
positive=2, negative=2, total=4

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=2, total=4

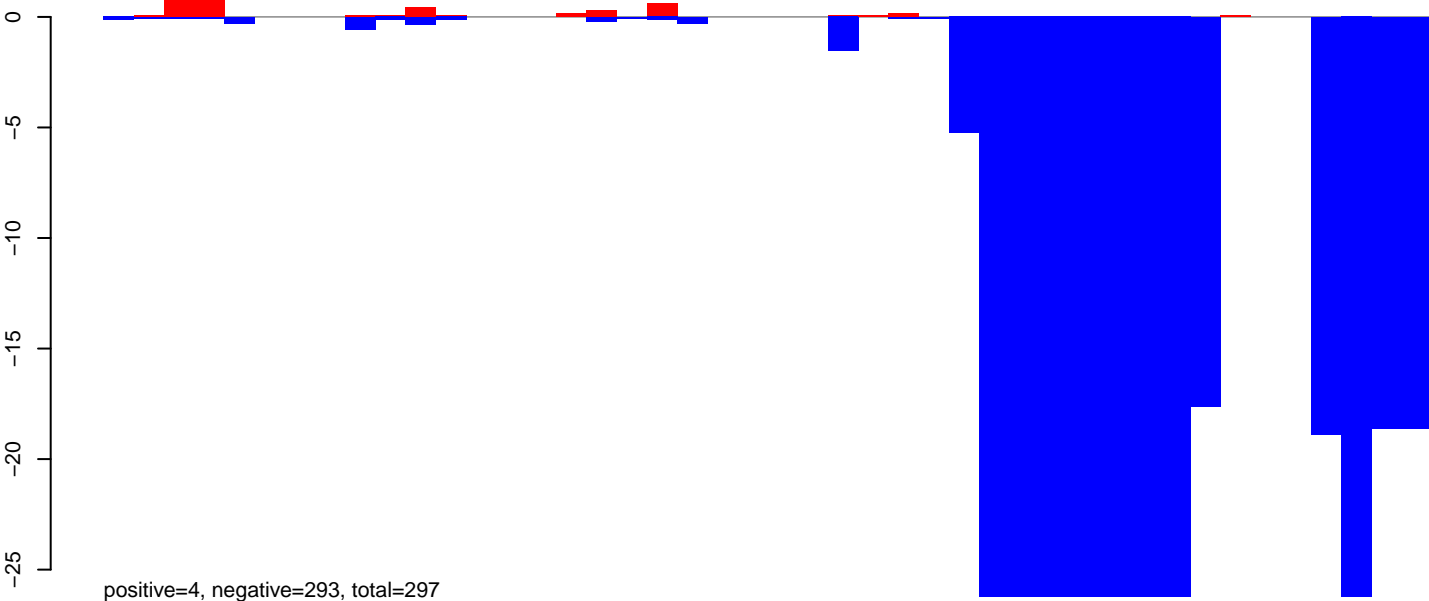
CuQuin_Mcarc_NL.rep



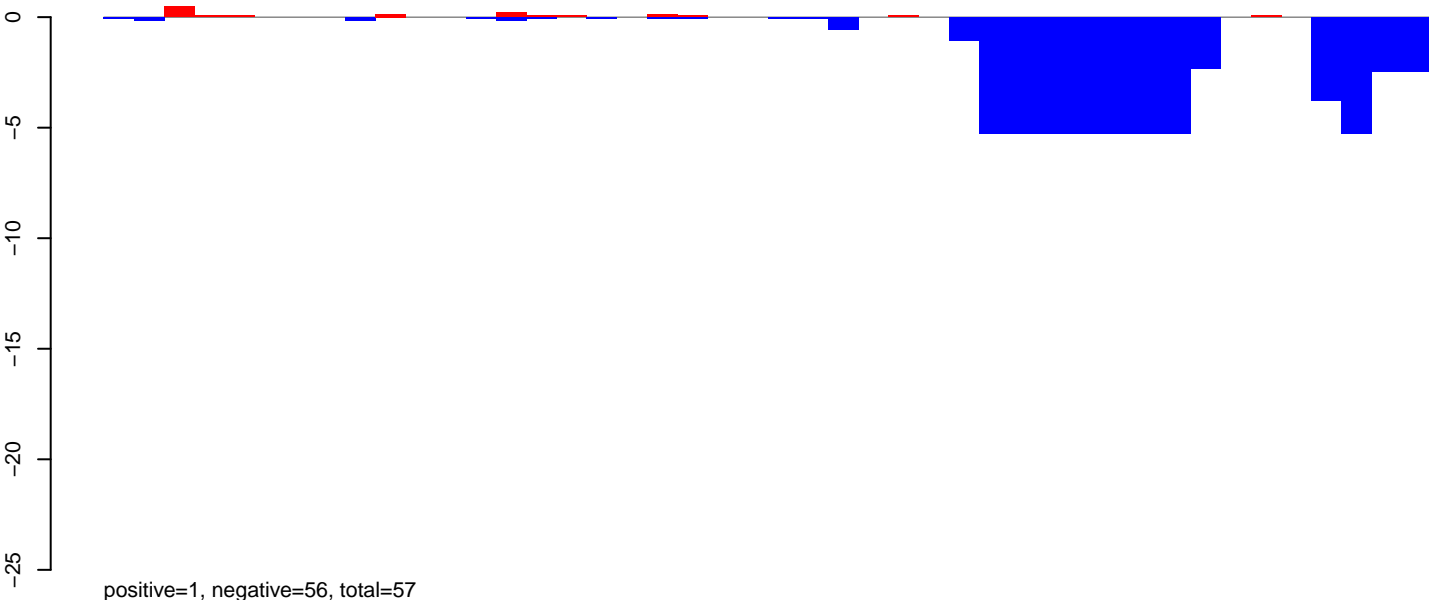
positive=3, negative=4, total=8

Window size=25, length=1414 TE@rnd-3_family-54@Unknown@75:1-1414

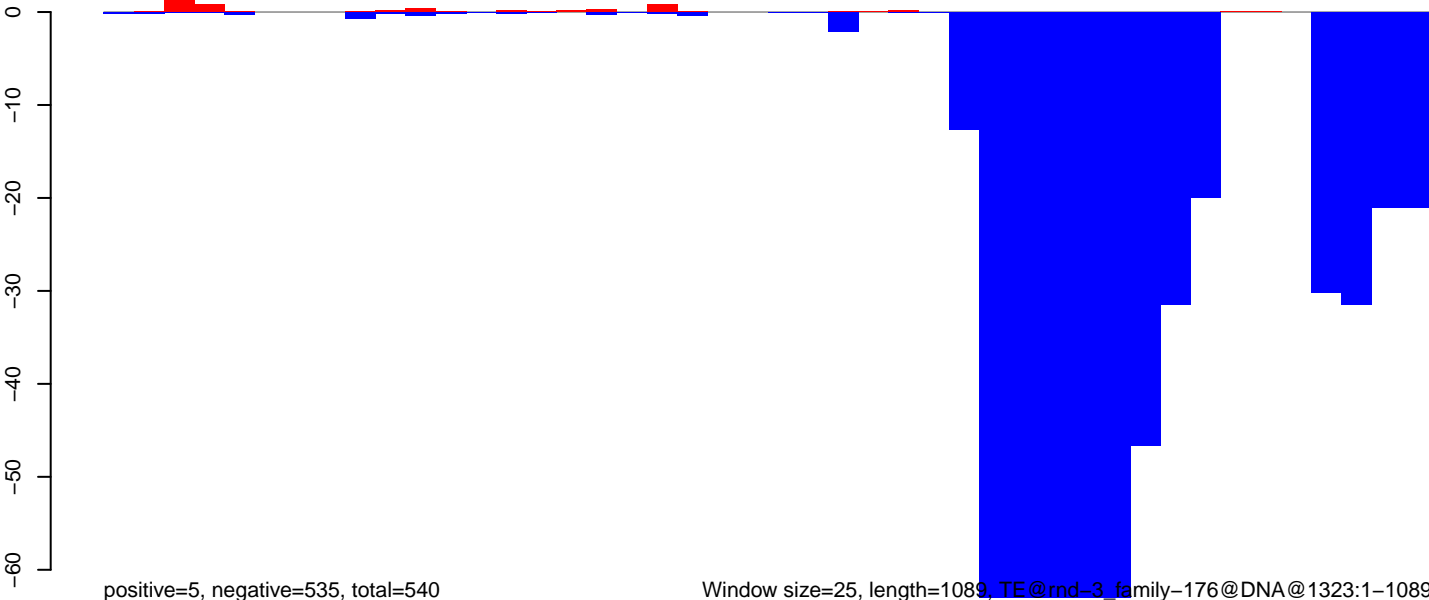
CuQuin_Mcarc_NL.18_23.rep



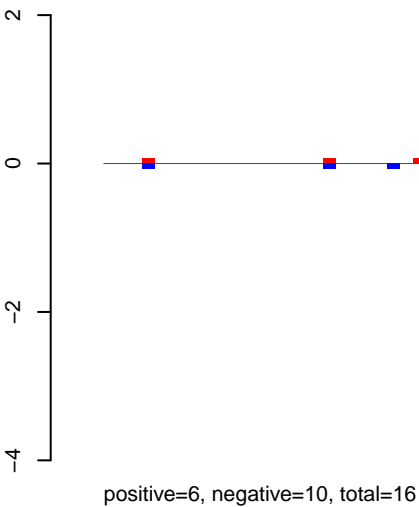
CuQuin_Mcarc_NL.24_35.rep



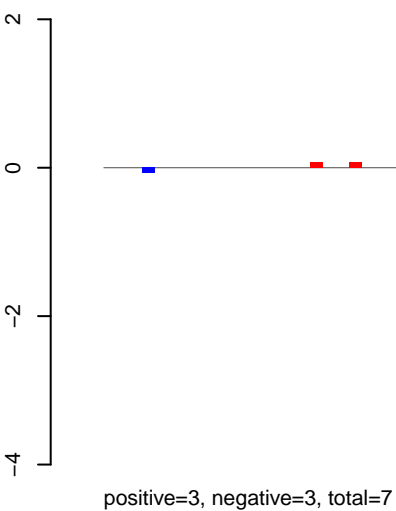
CuQuin_Mcarc_NL.rep



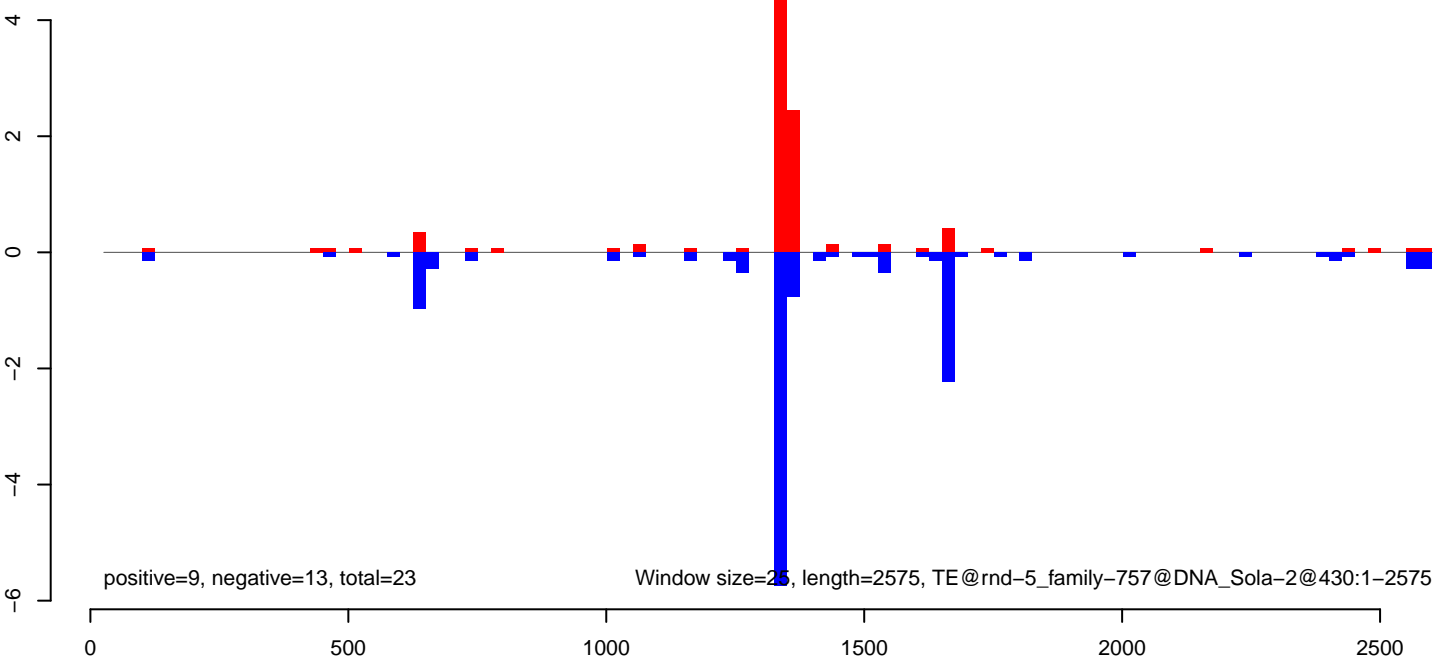
CuQuin_Mcarc_NL.18_23.rep



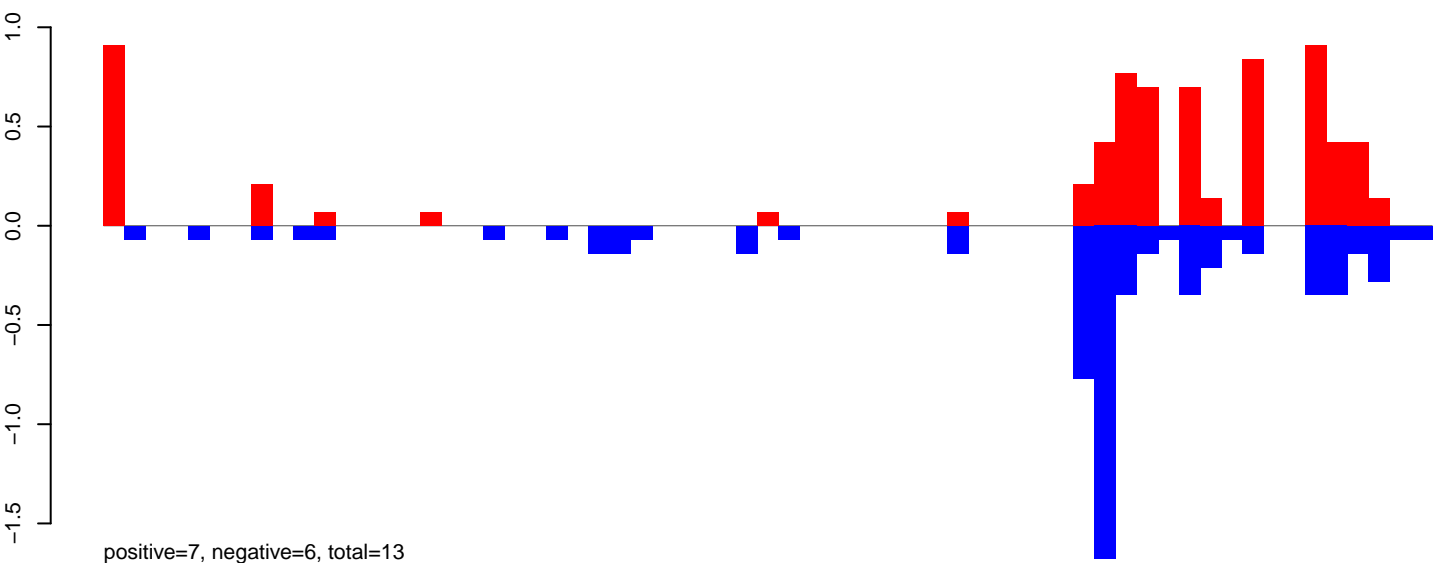
CuQuin_Mcarc_NL.24_35.rep



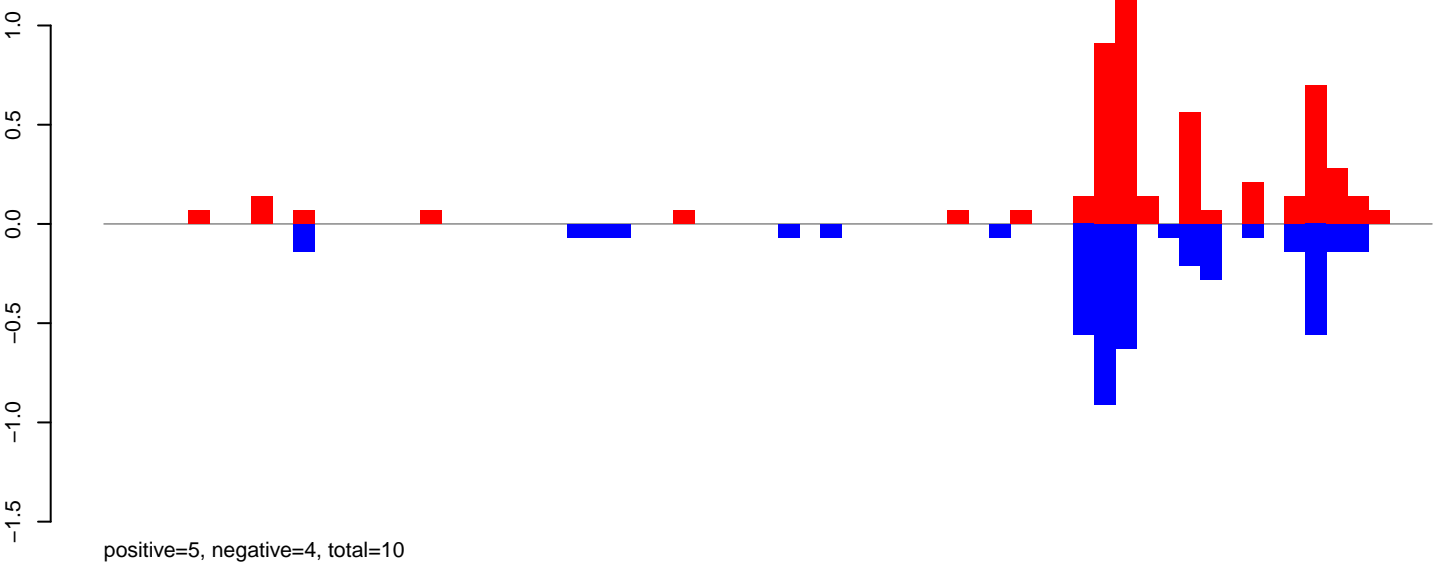
CuQuin_Mcarc_NL.rep



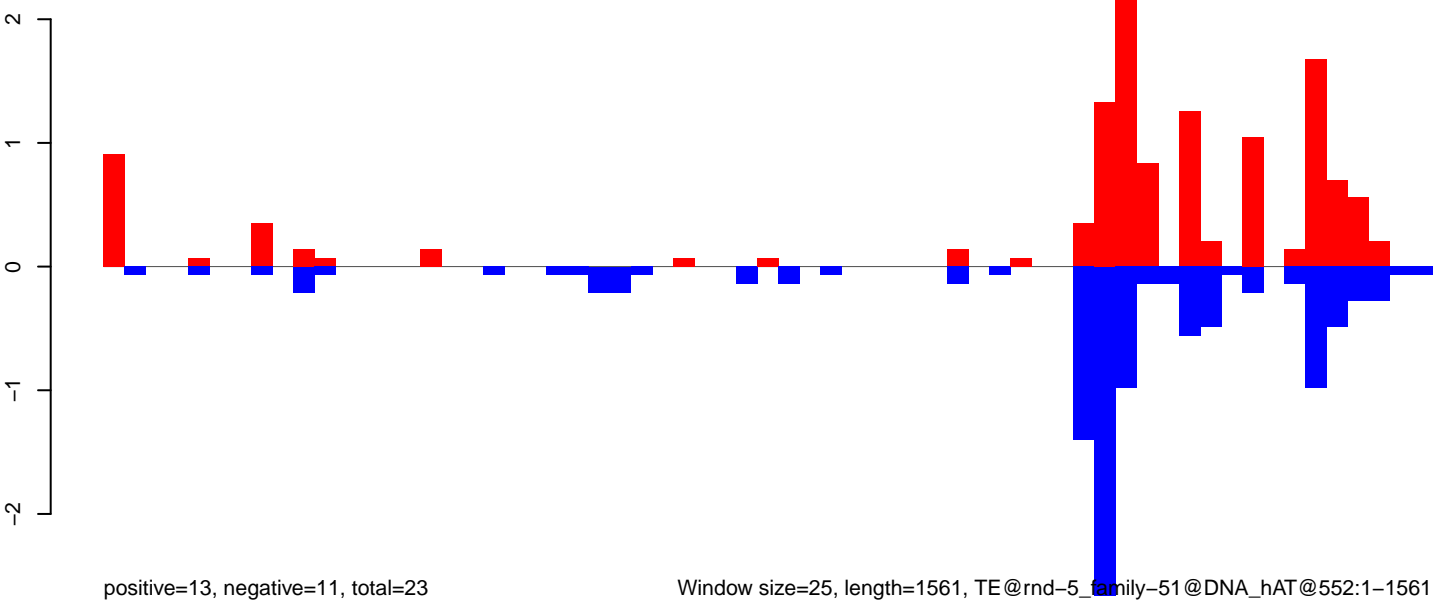
CuQuin_Mcarc_NL.18_23.rep



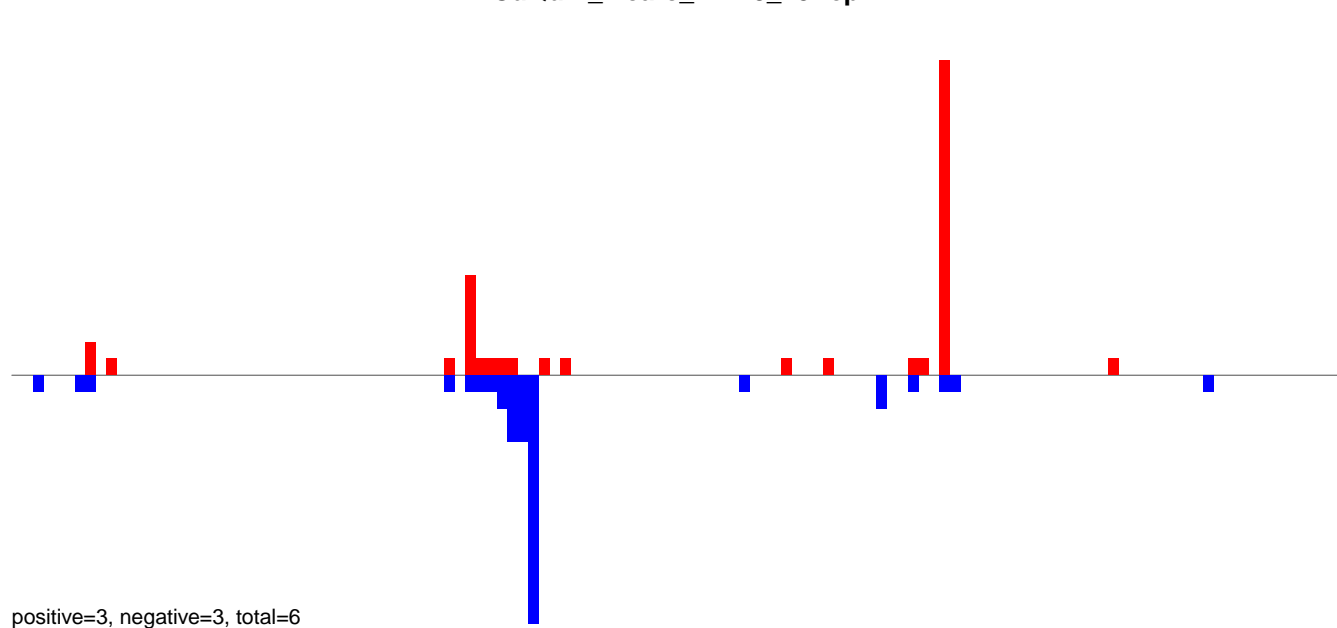
CuQuin_Mcarc_NL.24_35.rep



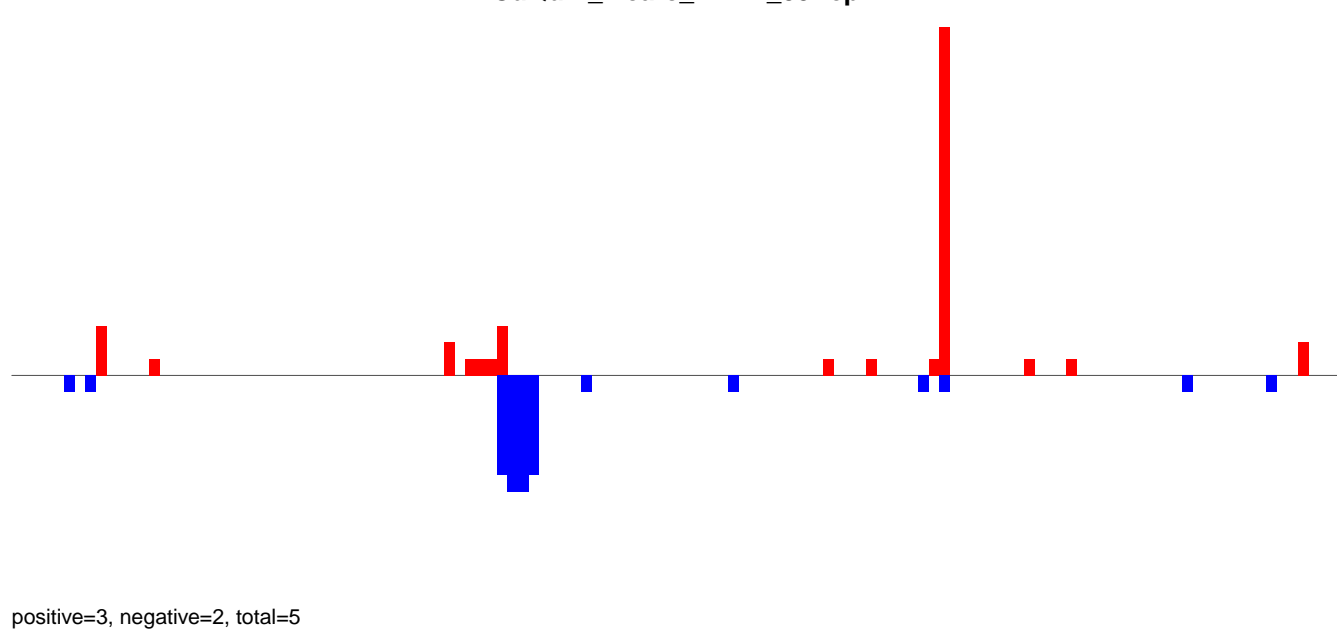
CuQuin_Mcarc_NL.rep



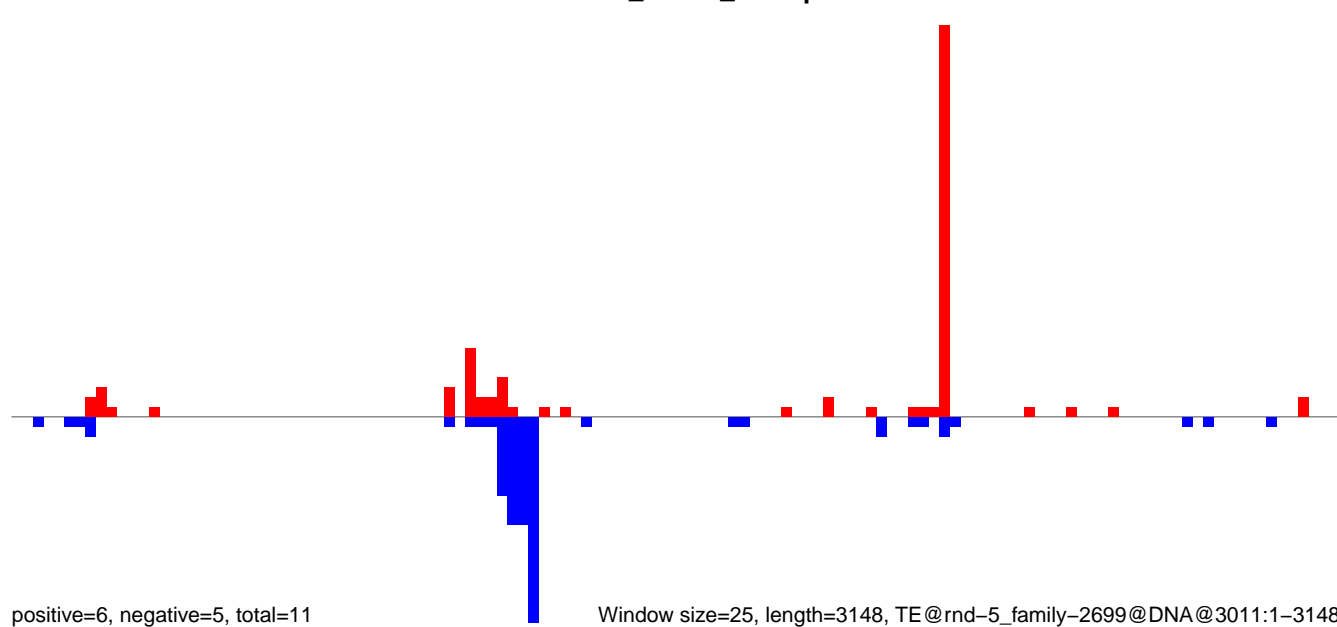
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

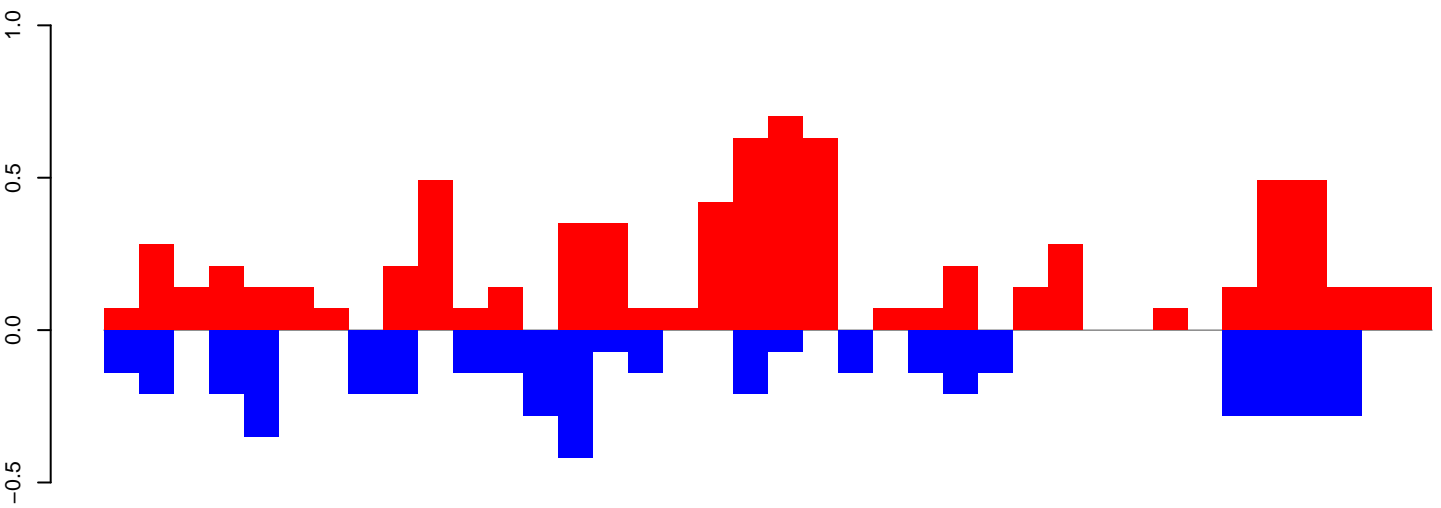


CuQuin_Mcarc_NL.rep



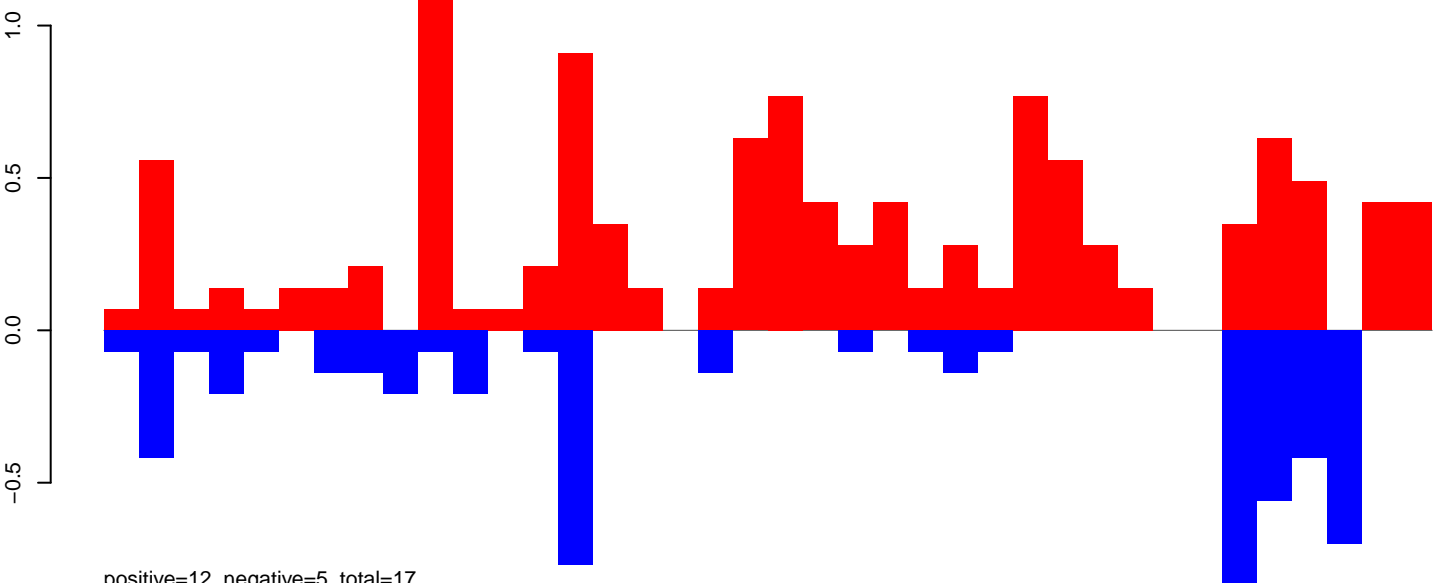
Window size=25, length=3148, TE@rnd-5_family-2699@DNA@3011:1-3148

CuQuin_Mcarc_NL.18_23.rep



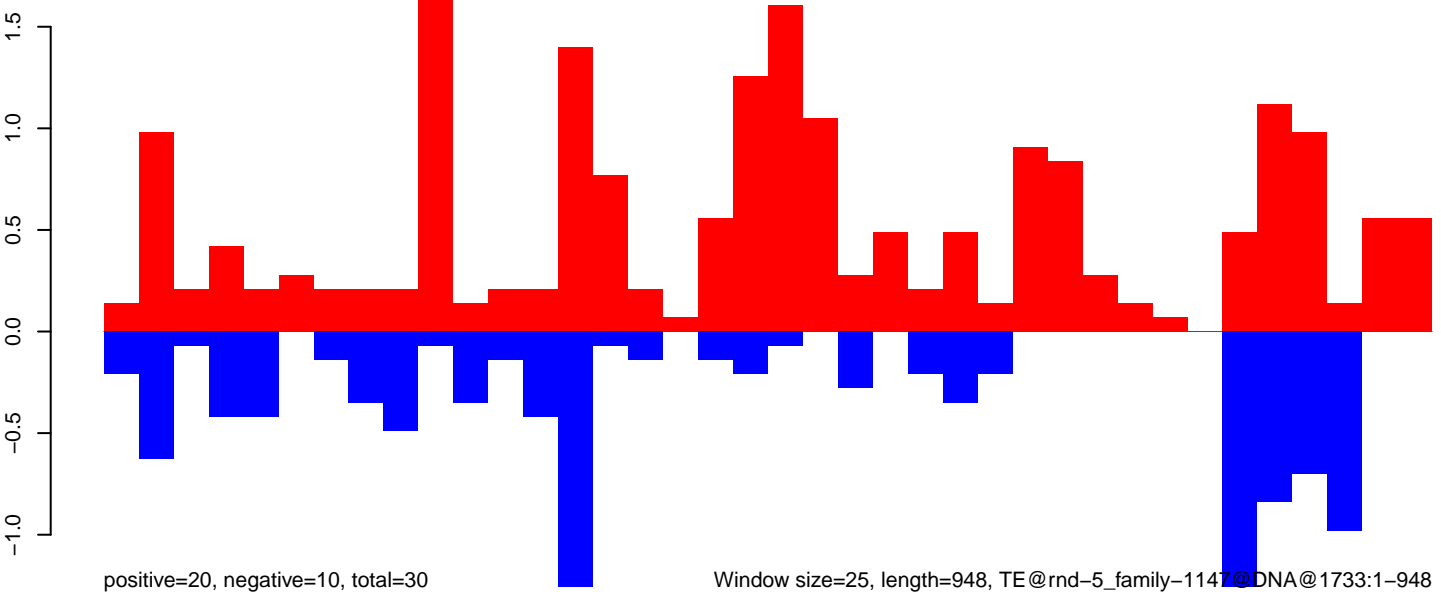
positive=8, negative=5, total=12

CuQuin_Mcarc_NL.24_35.rep



positive=12, negative=5, total=17

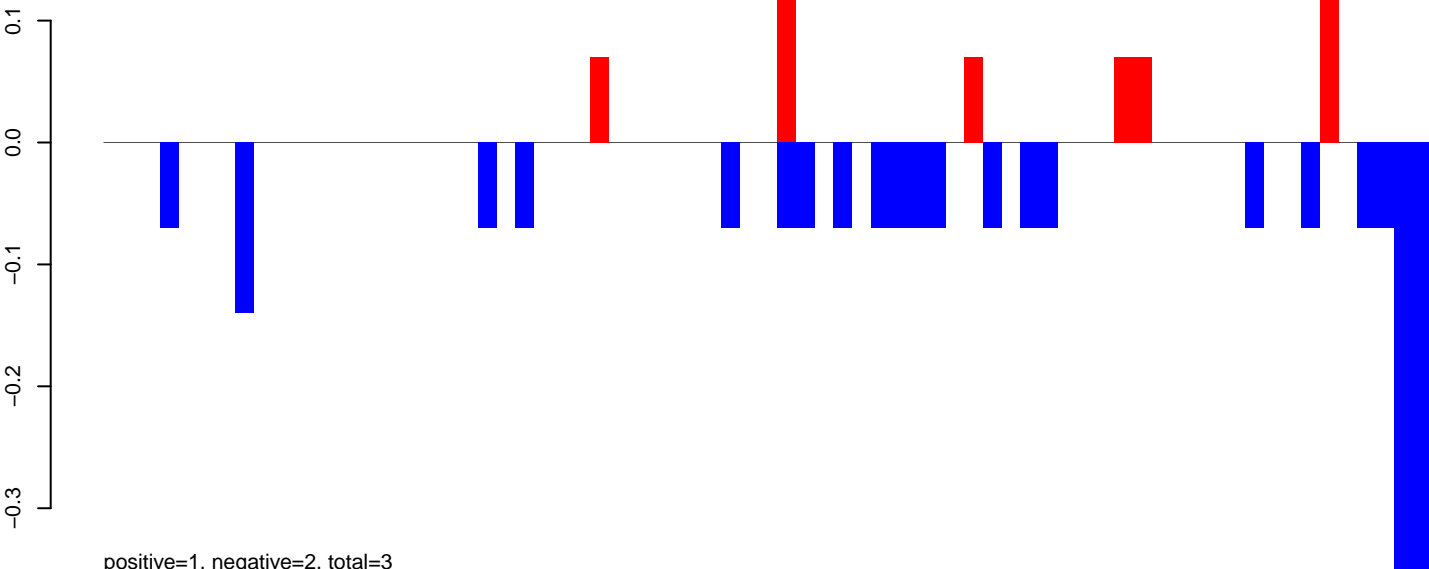
CuQuin_Mcarc_NL.rep



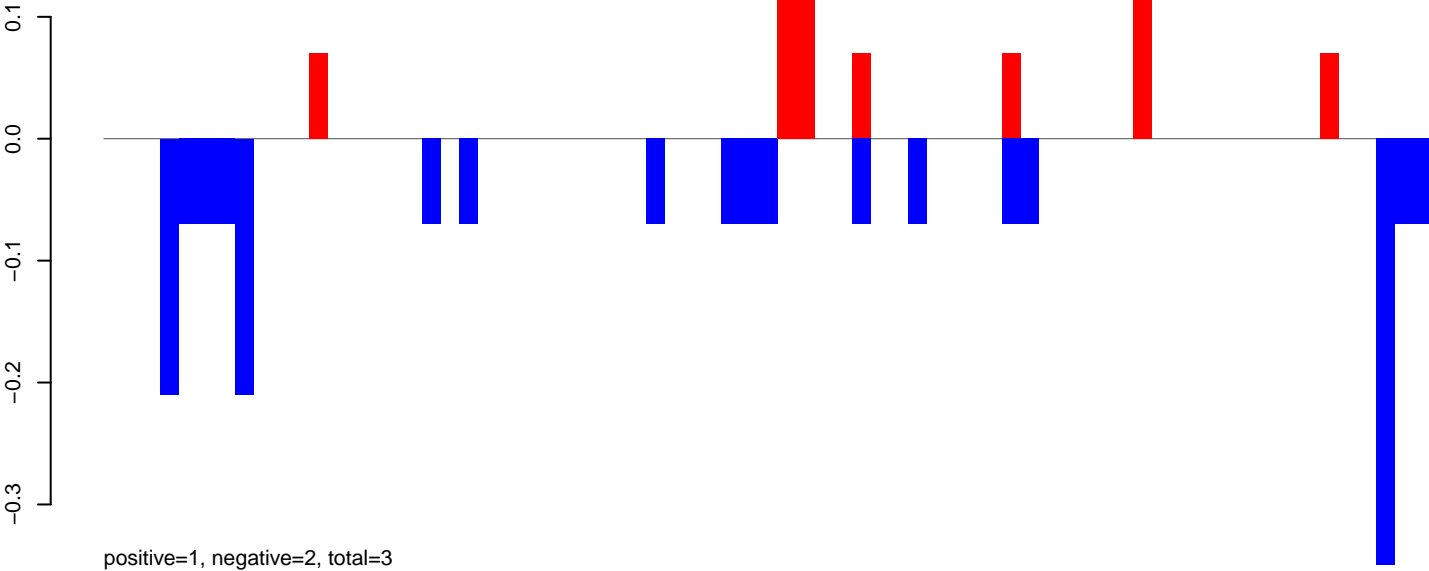
positive=20, negative=10, total=30

Window size=25, length=948, TE@rnd-5_family-1147@DNA@1733:1-948

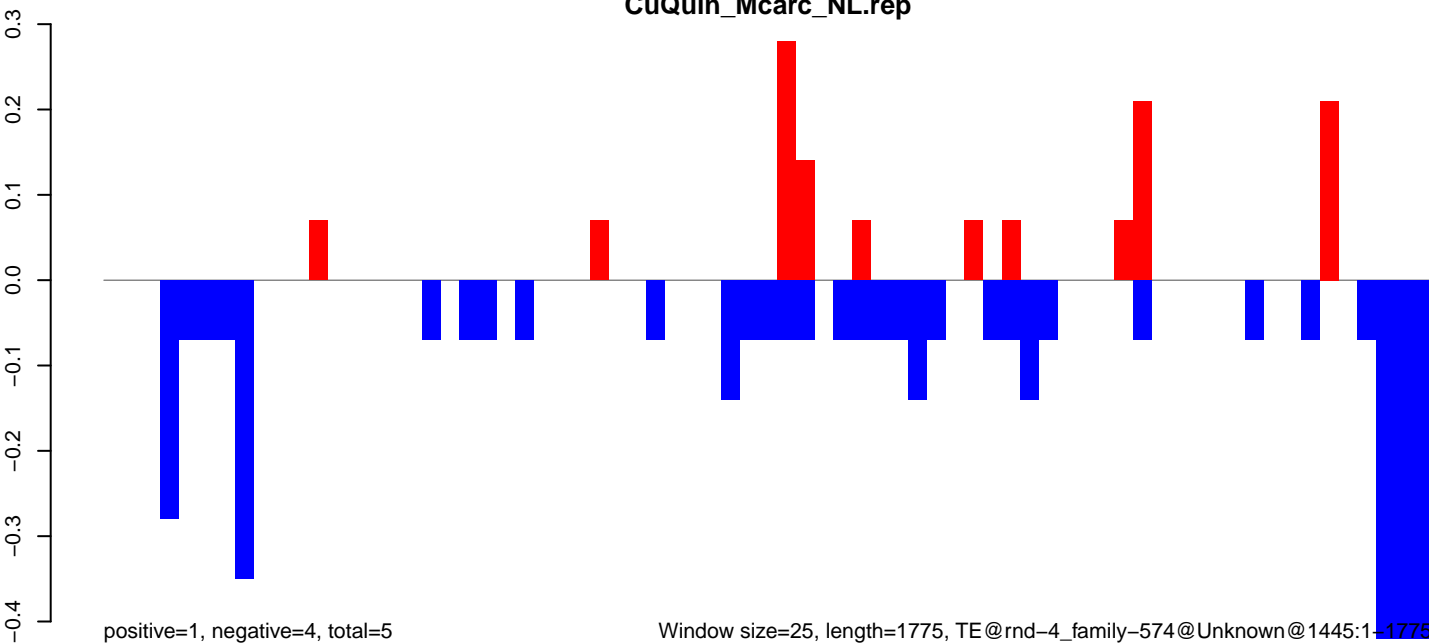
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

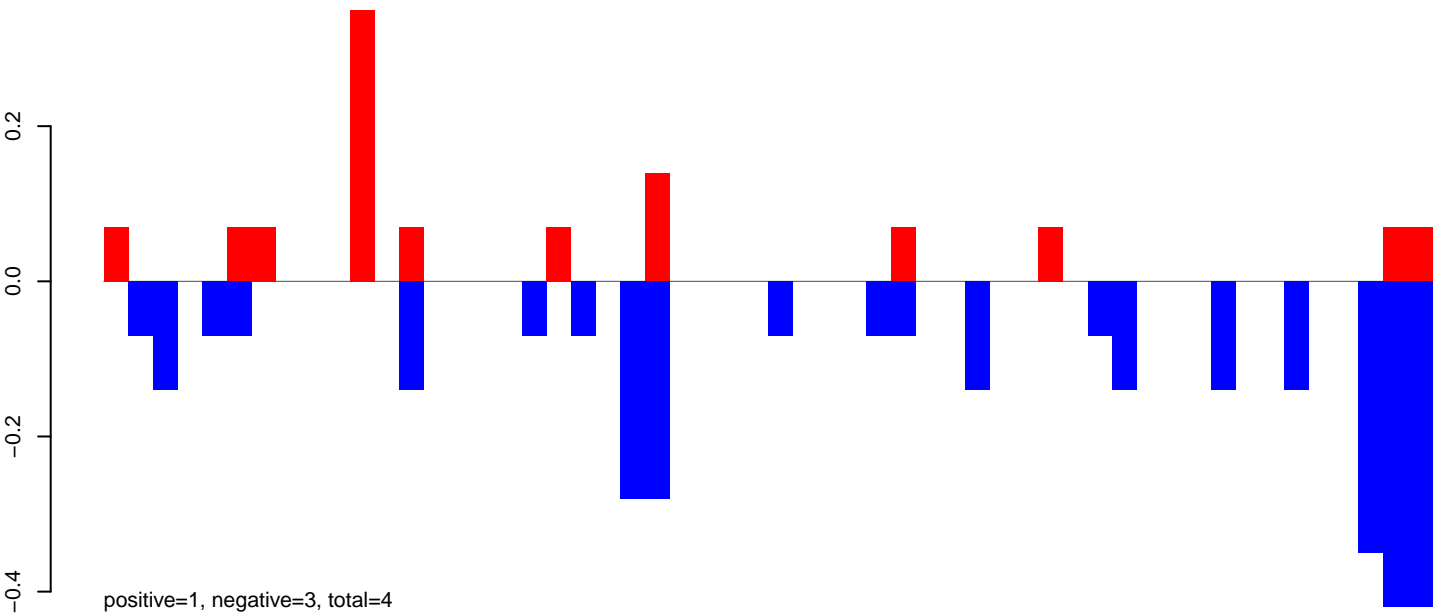


CuQuin_Mcarc_NL.rep

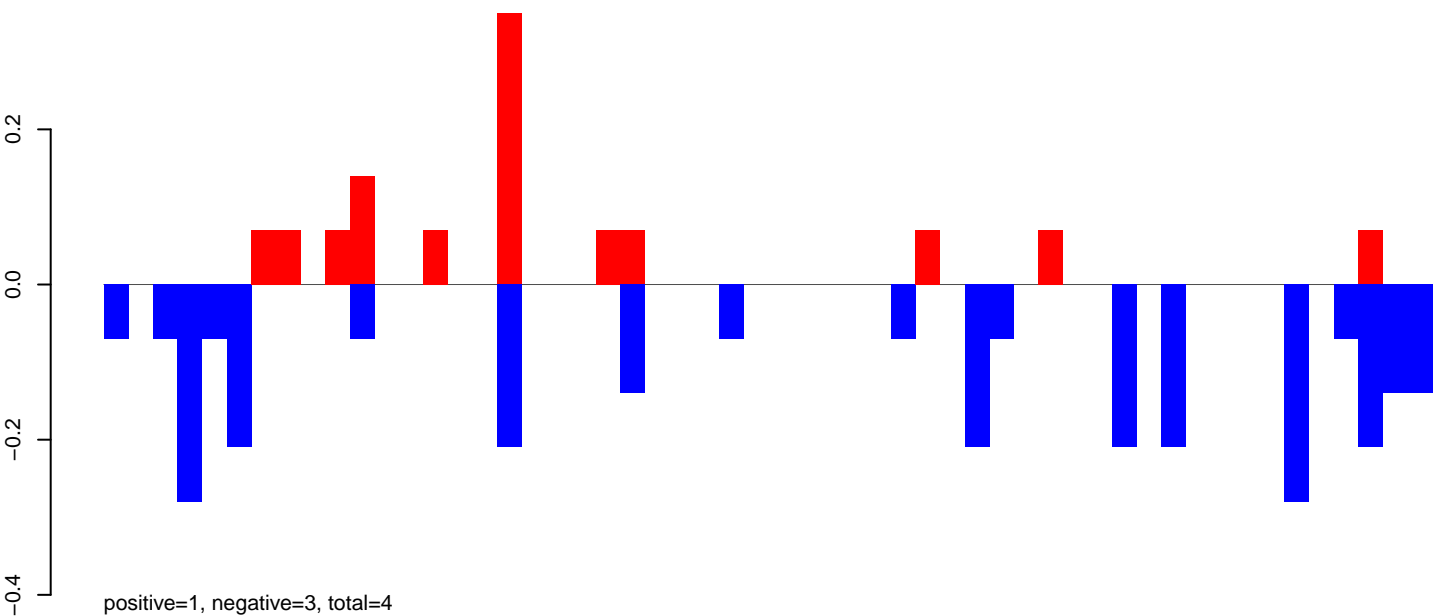


Window size=25, length=1775, TE@rnd-4_family-574@Unknown@1445:1-1775

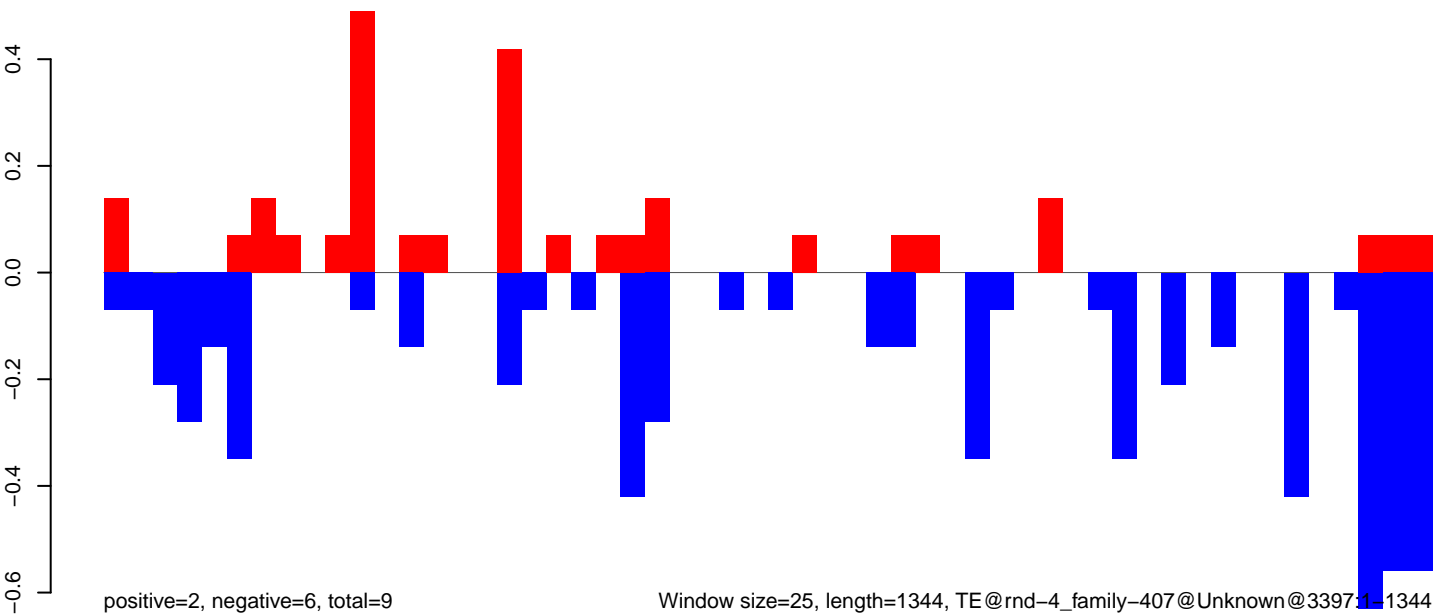
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

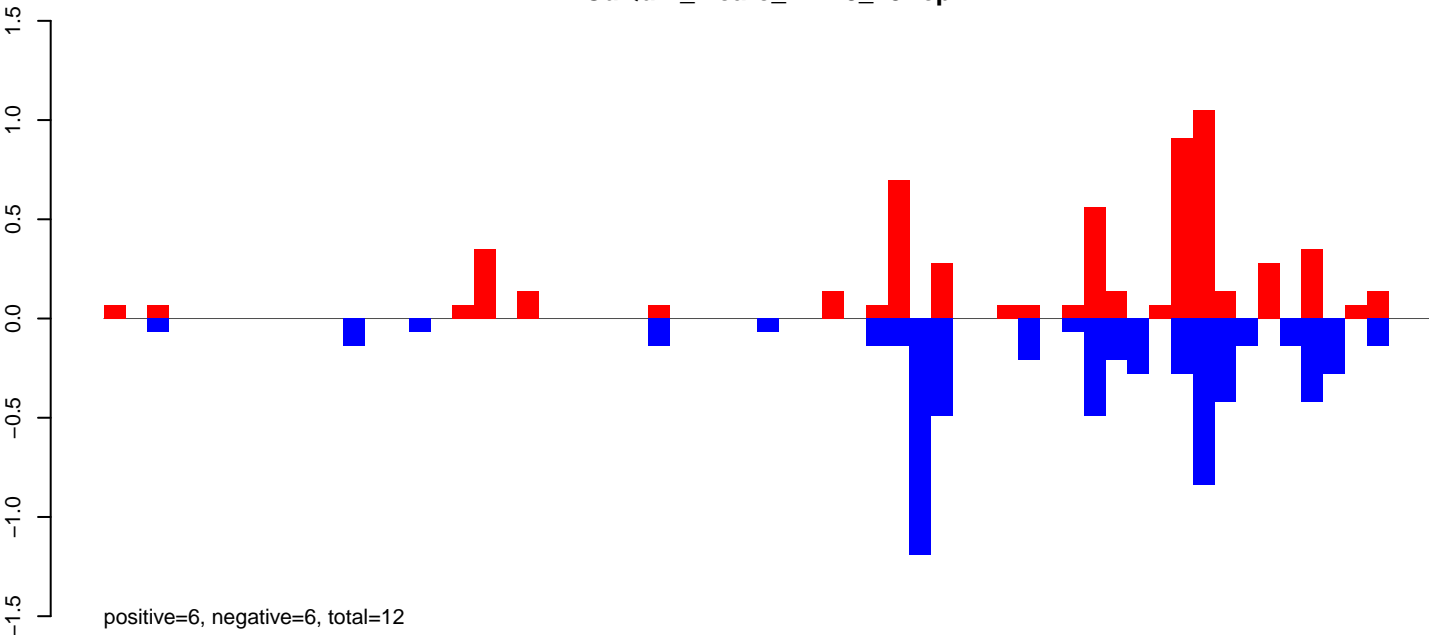


CuQuin_Mcarc_NL.rep

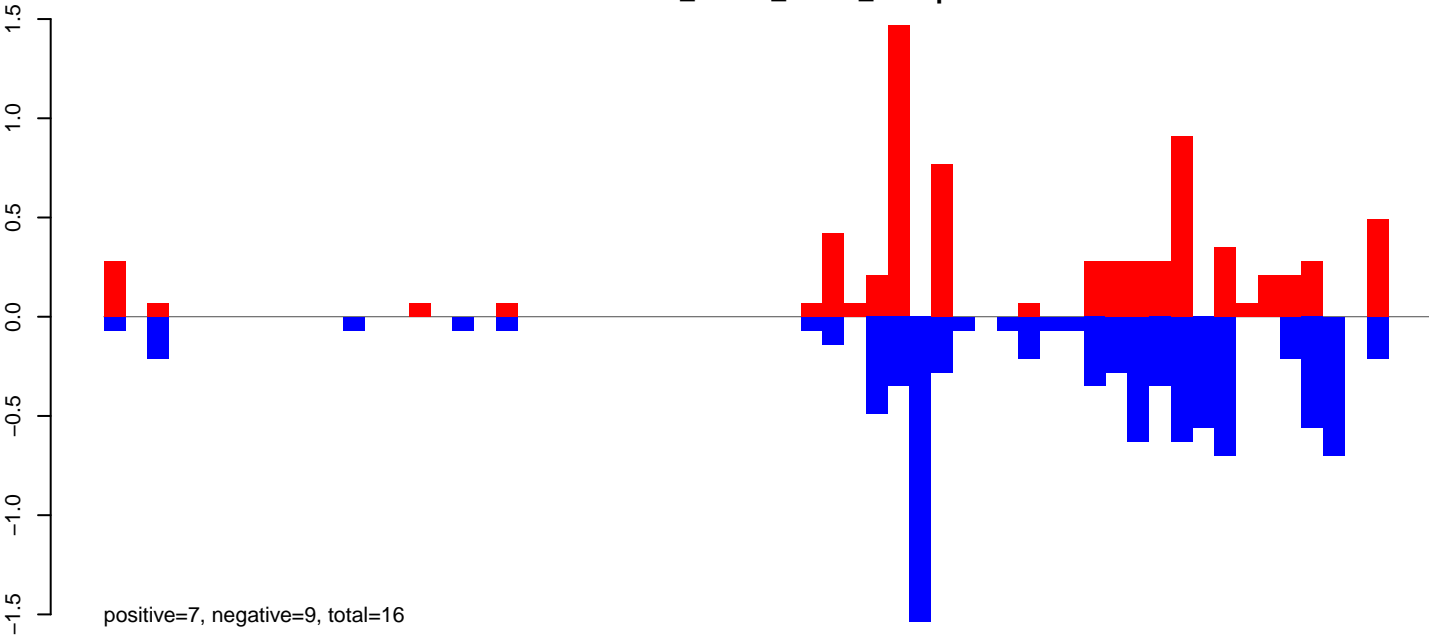


Window size=25, length=1344, TE@rnd-4_family-407@Unknown@33971-1344

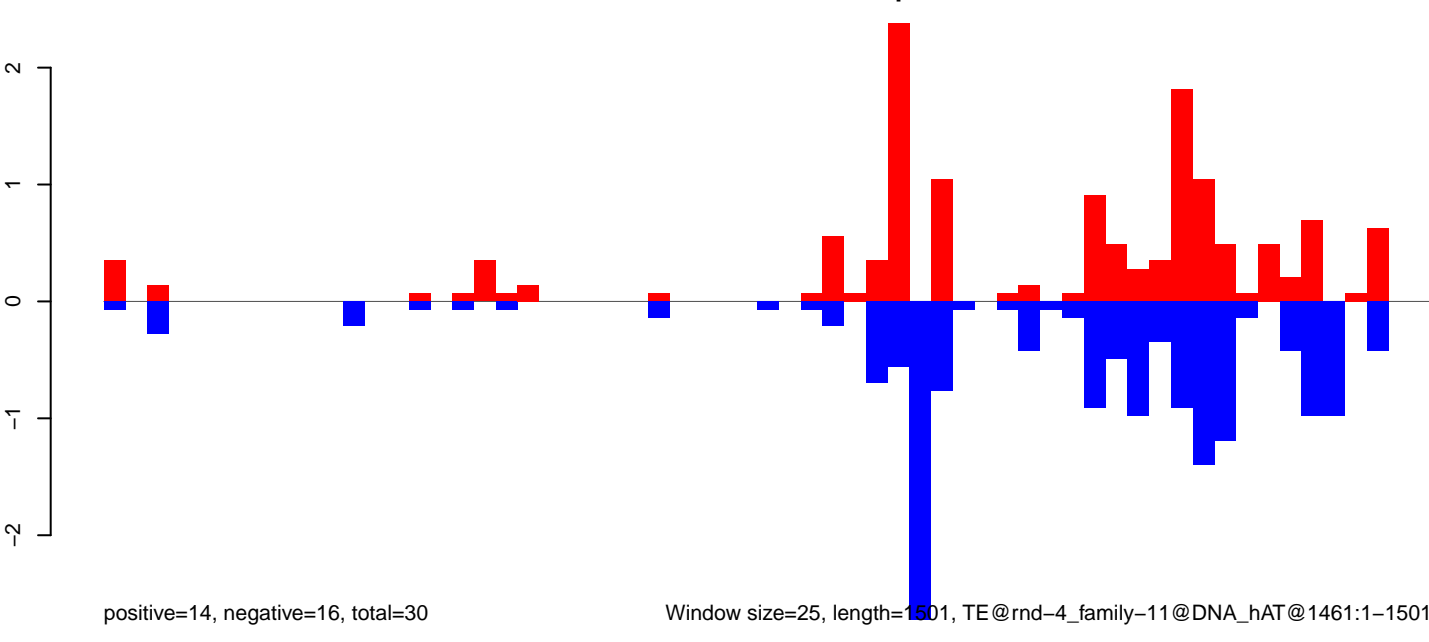
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

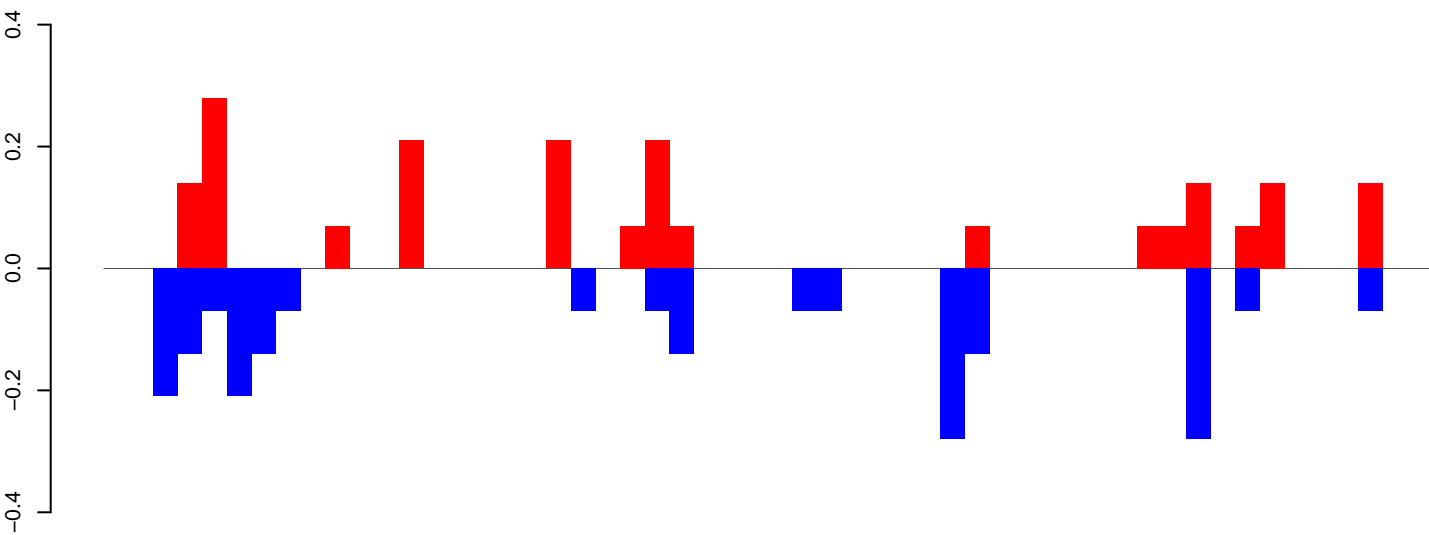


CuQuin_Mcarc_NL.rep



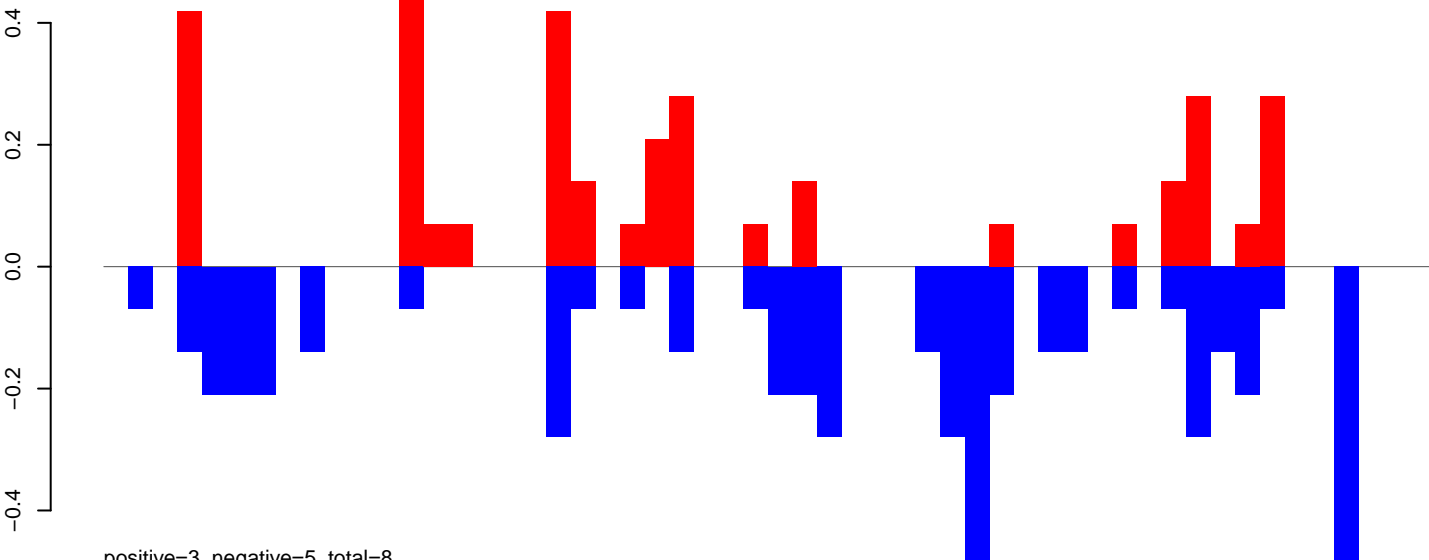
Window size=25, length=1501, TE@rnd-4_family-11@DNA_hAT@1461:1-1501

CuQuin_Mcarc_NL.18_23.rep



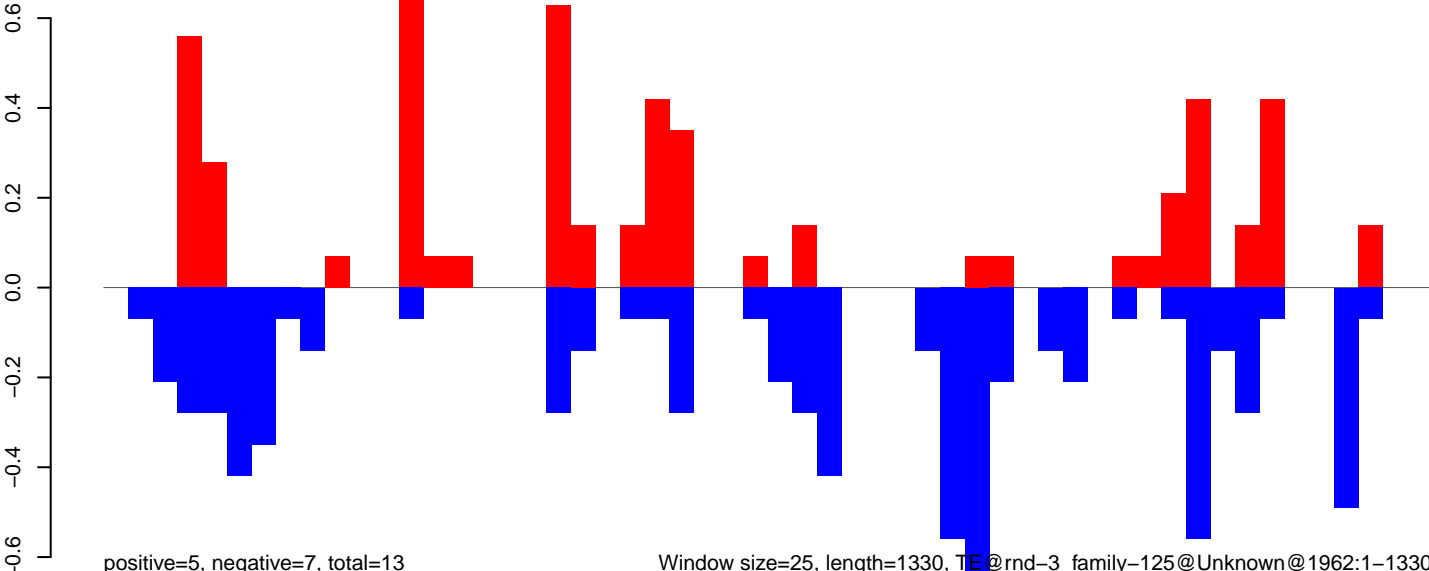
positive=2, negative=2, total=4

CuQuin_Mcarc_NL.24_35.rep



positive=3, negative=5, total=8

CuQuin_Mcarc_NL.rep

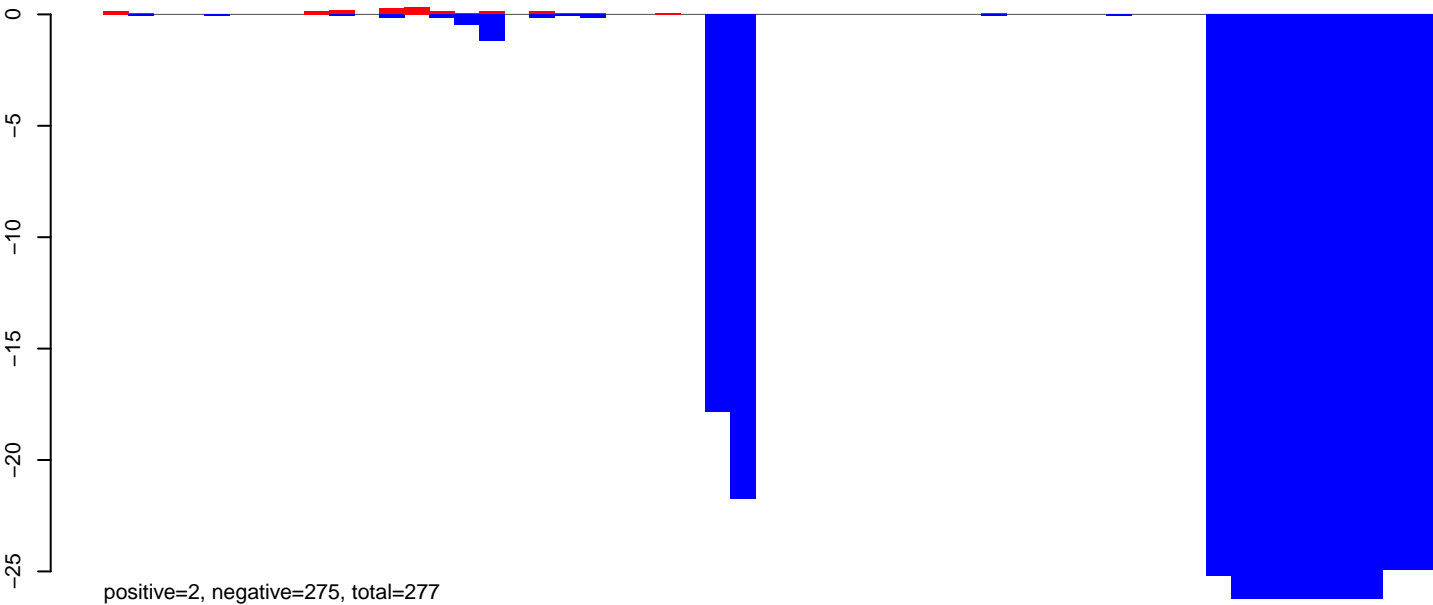


positive=5, negative=7, total=13

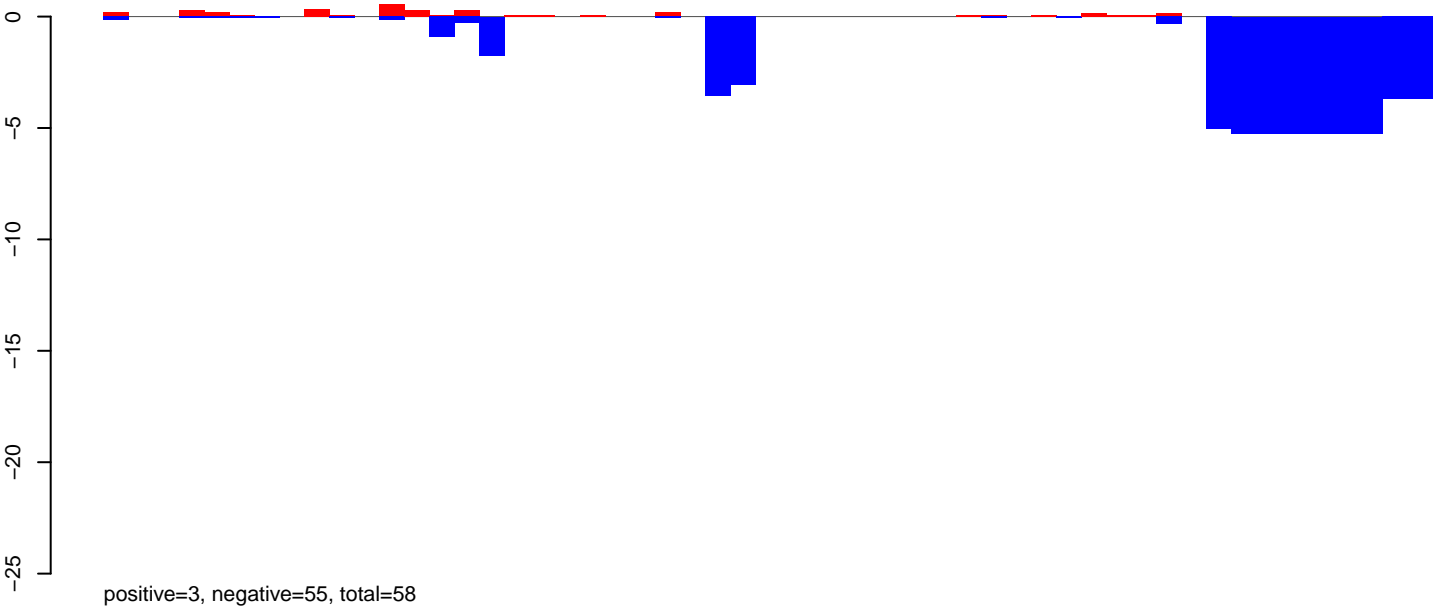
Window size=25, length=1330, TE@rnd-3_family-125@Unknown@1962:1-1330

0 200 400 600 800 1000 1200 1400

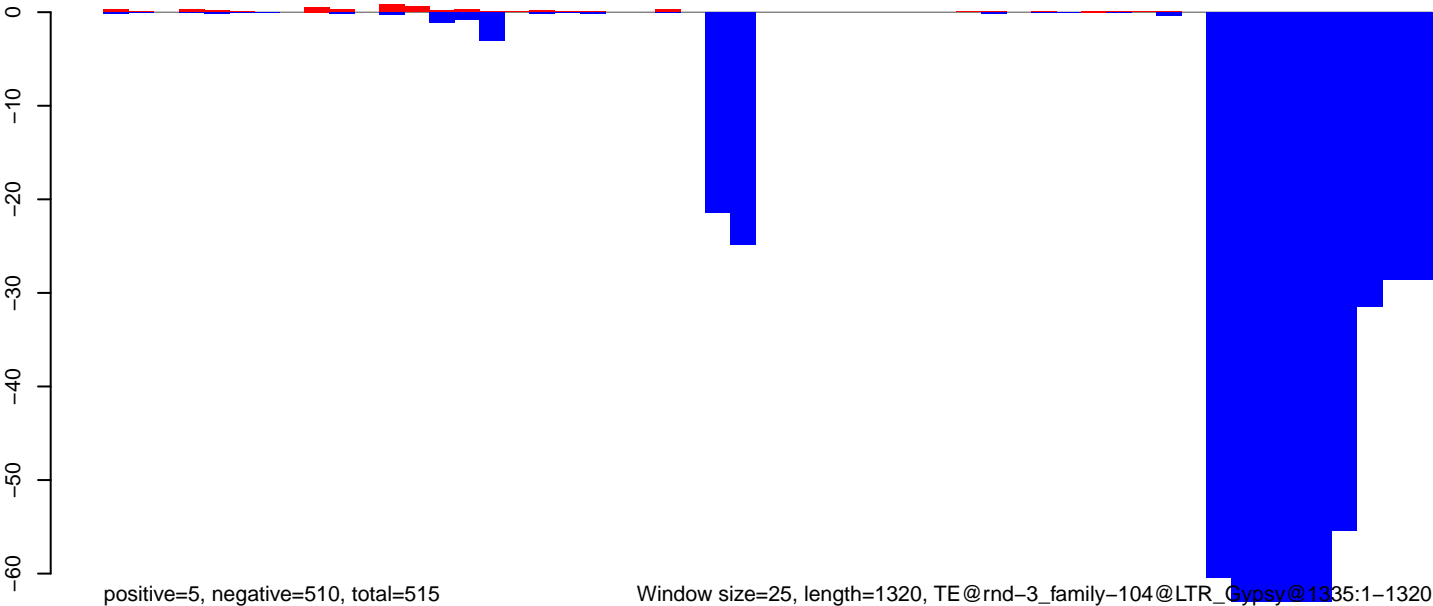
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

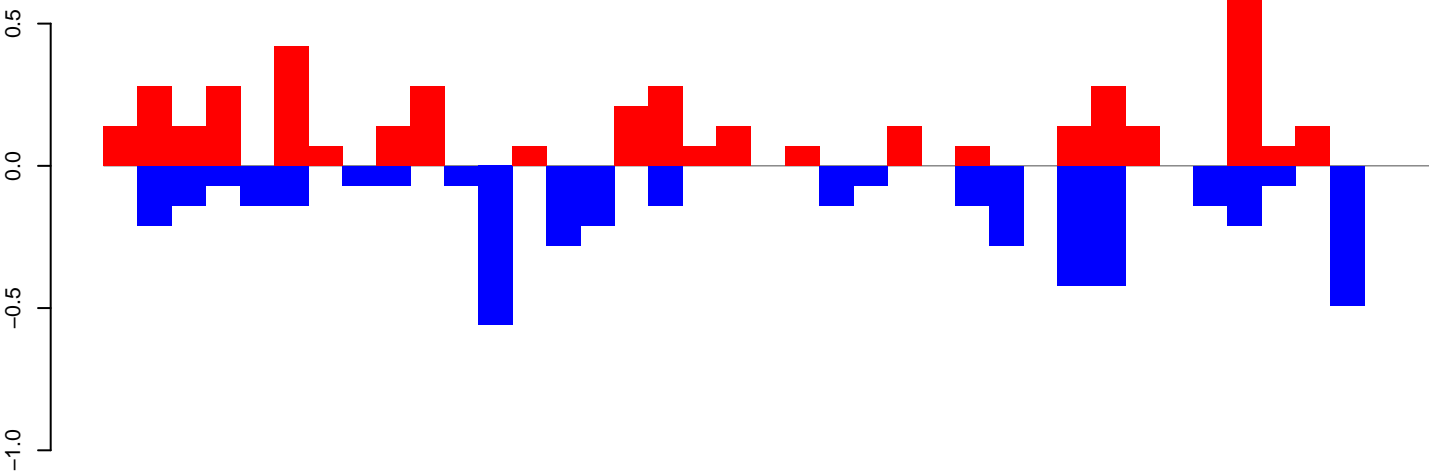


CuQuin_Mcarc_NL.rep



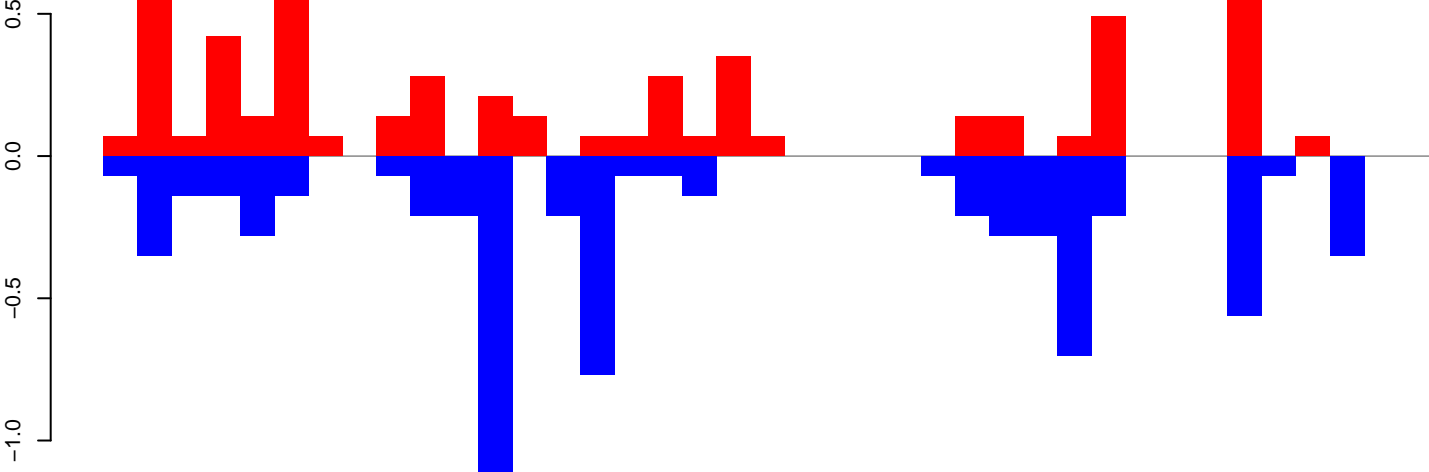
Window size=25, length=1320, TE@rnd-3_family-104@LTR_Gypsy@1335:1-1320

CuQuin_Mcarc_NL.18_23.rep



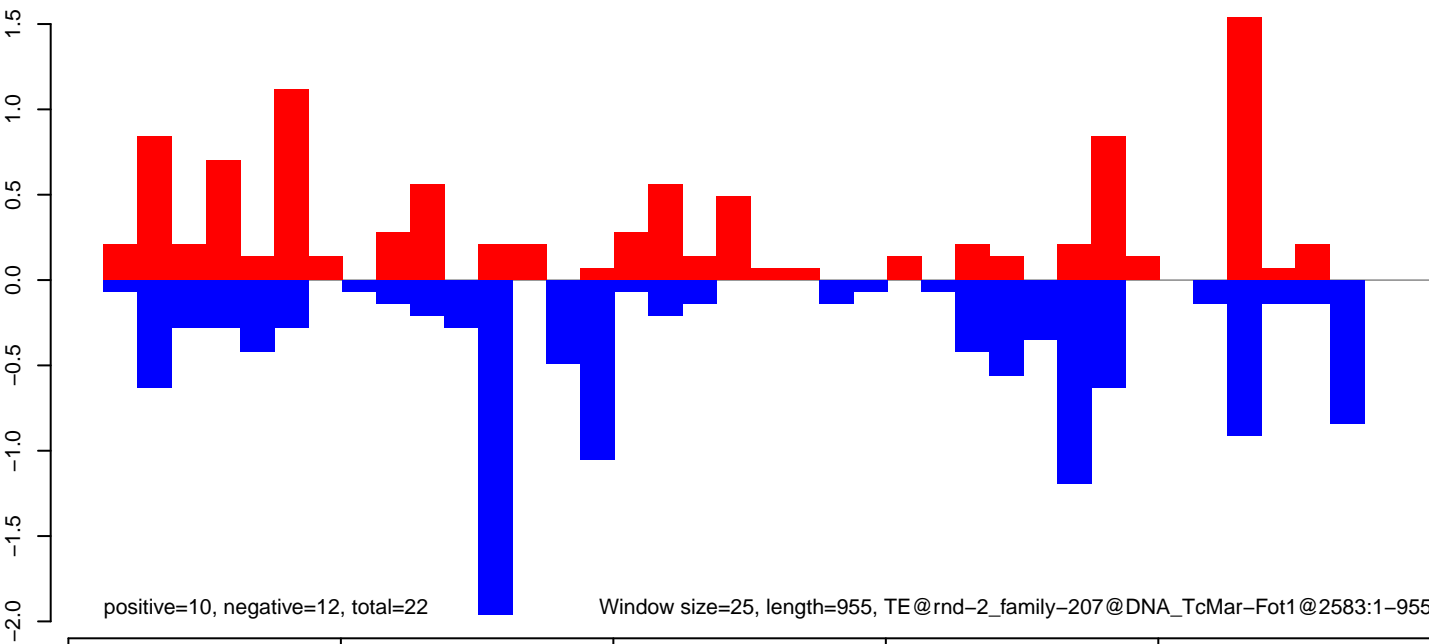
positive=4, negative=4, total=9

CuQuin_Mcarc_NL.24_35.rep



positive=5, negative=7, total=12

CuQuin_Mcarc_NL.rep

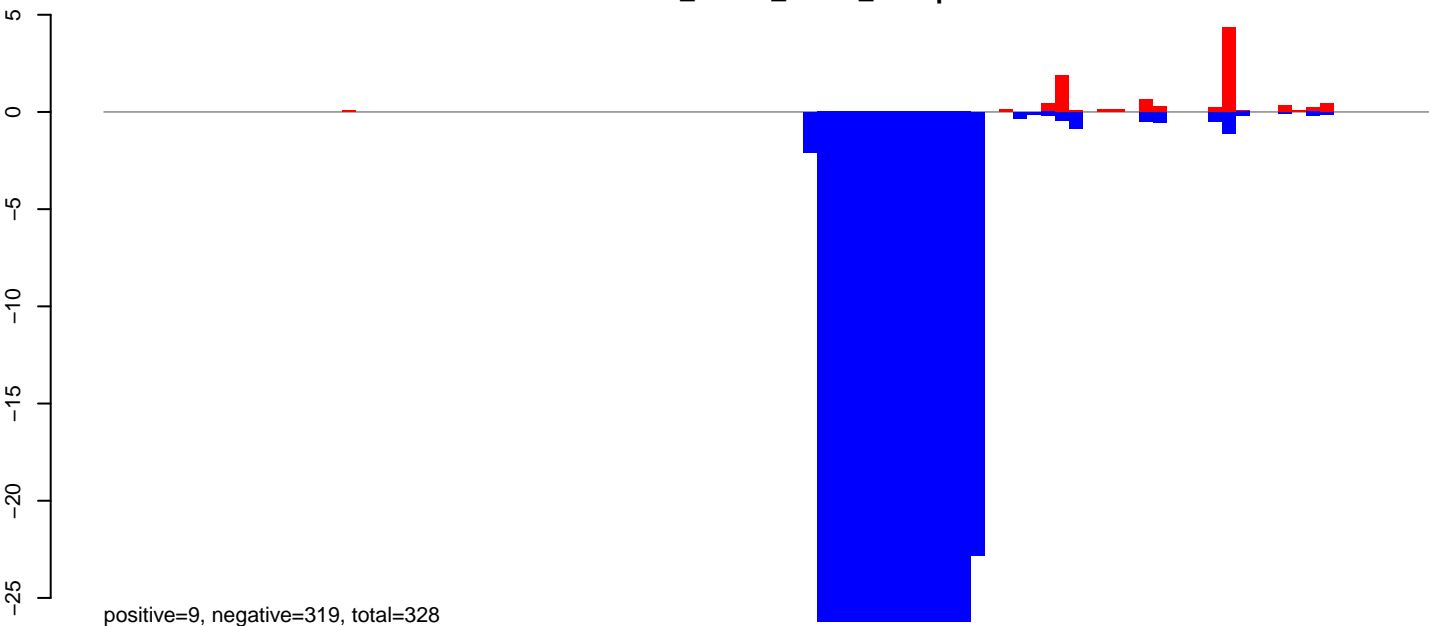


positive=10, negative=12, total=22

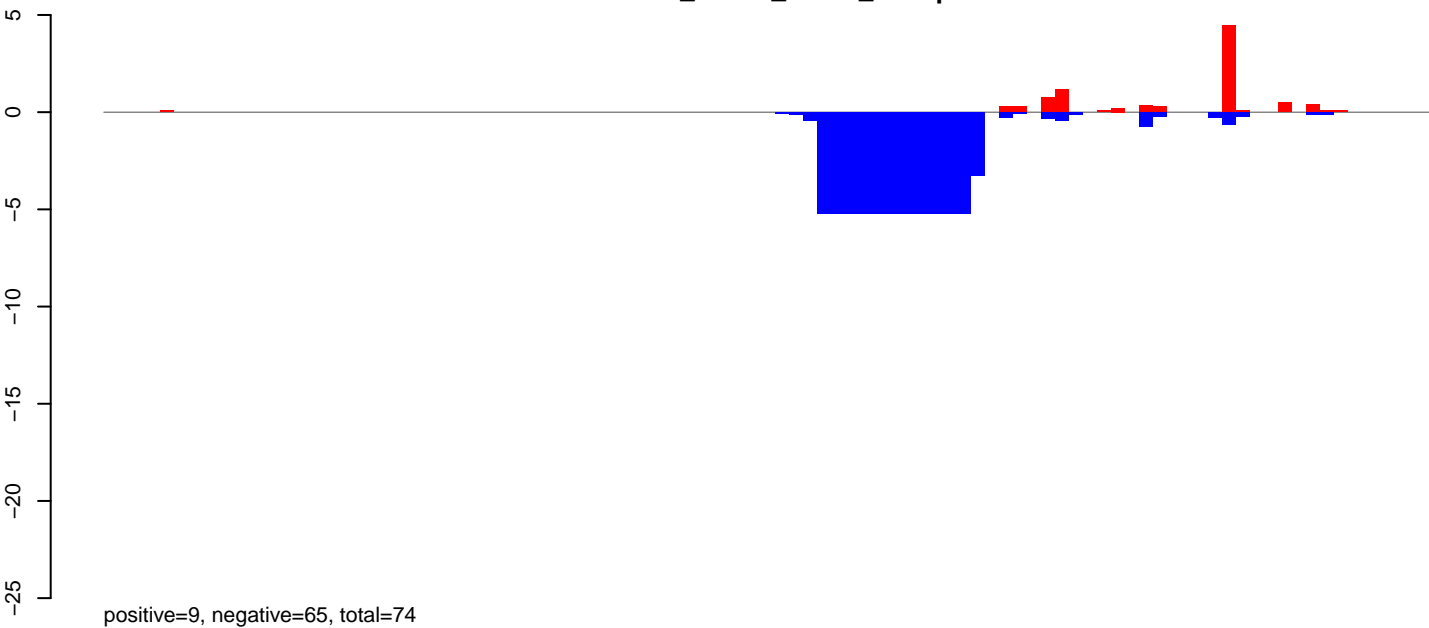
Window size=25, length=955, TE@rnd-2_family-207@DNA_TcMar-Fot1@2583:1-955

0 200 400 600 800 1000

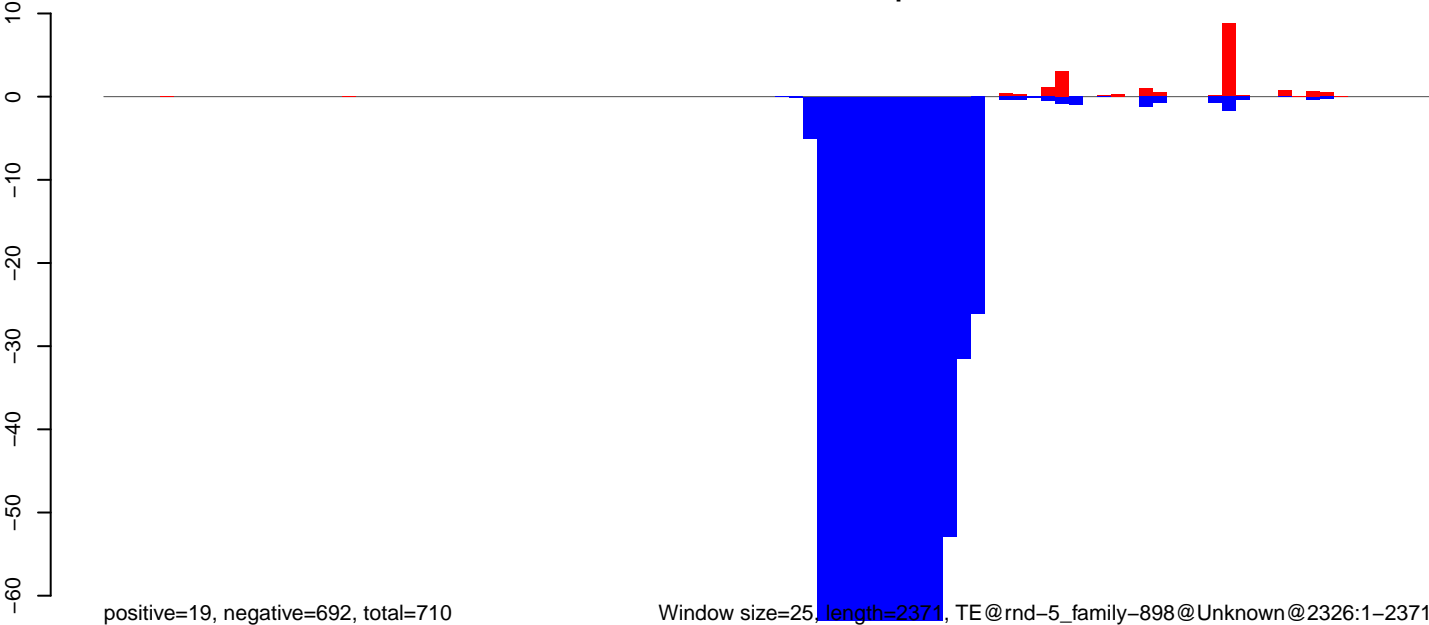
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



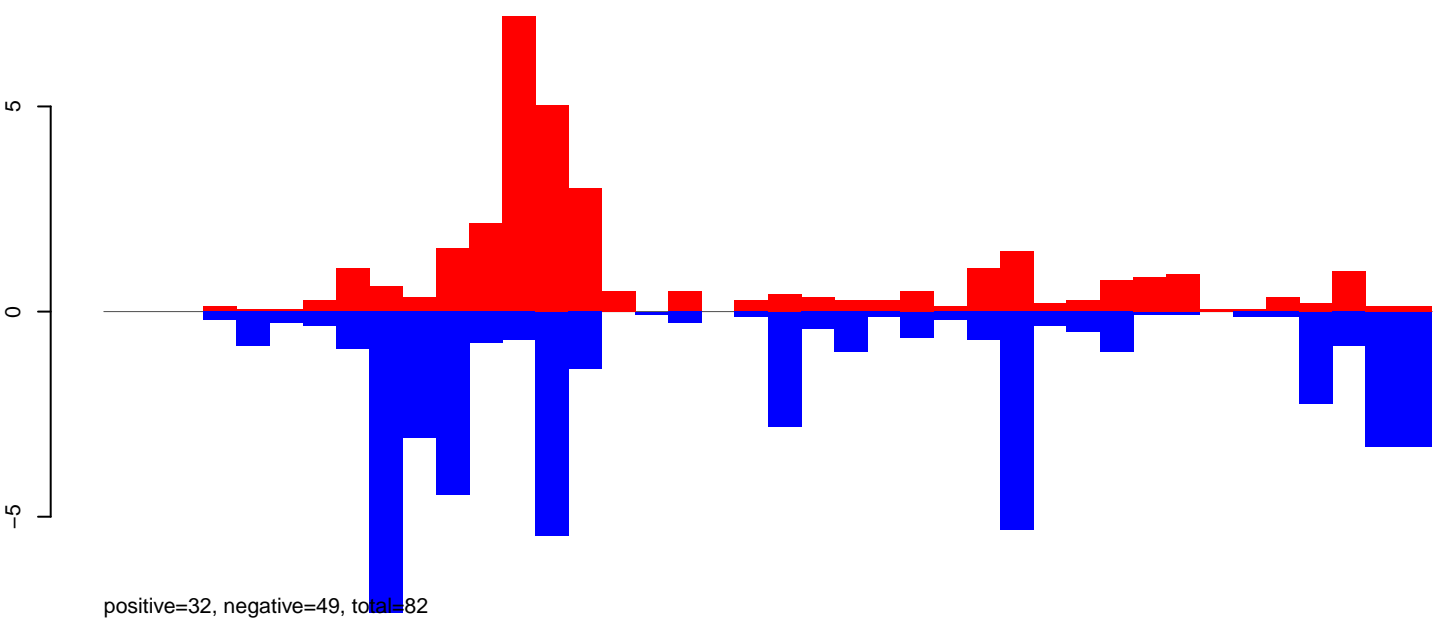
CuQuin_Mcarc_NL.rep



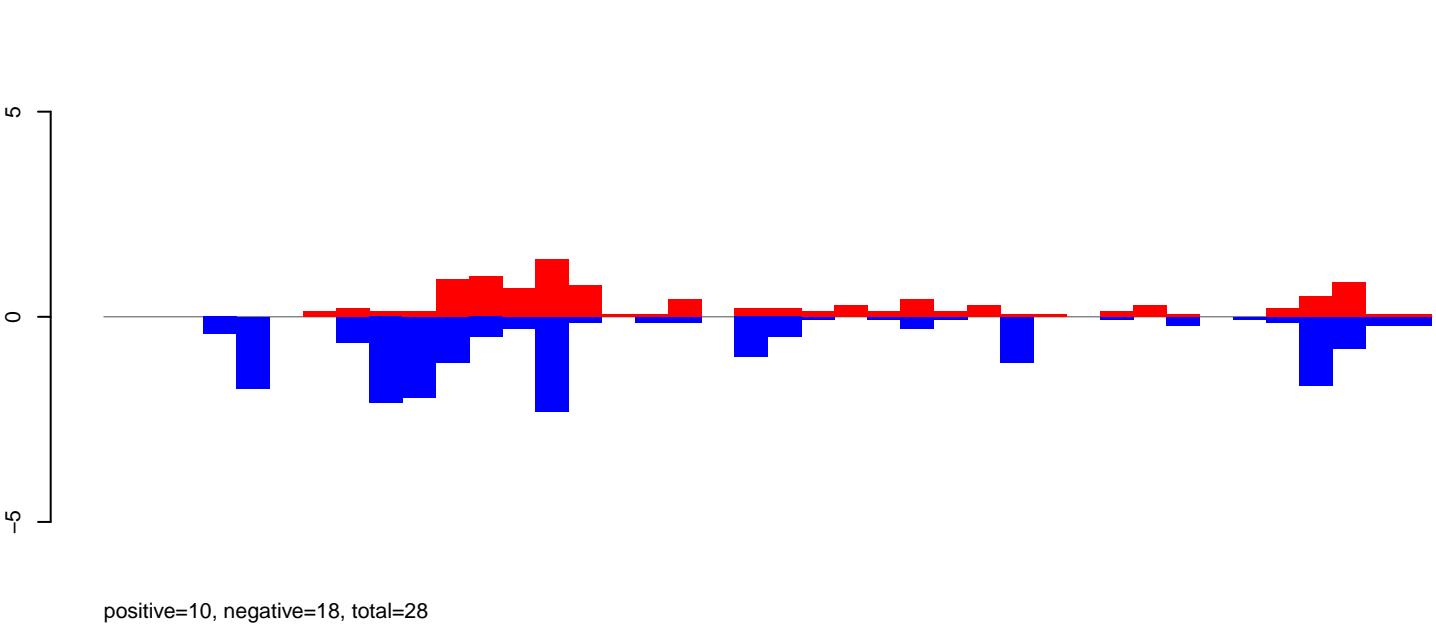
Window size=25, length=2371, TE@rnd-5_family-898@Unknown@2326:1-2371

0 500 1000 1500 2000

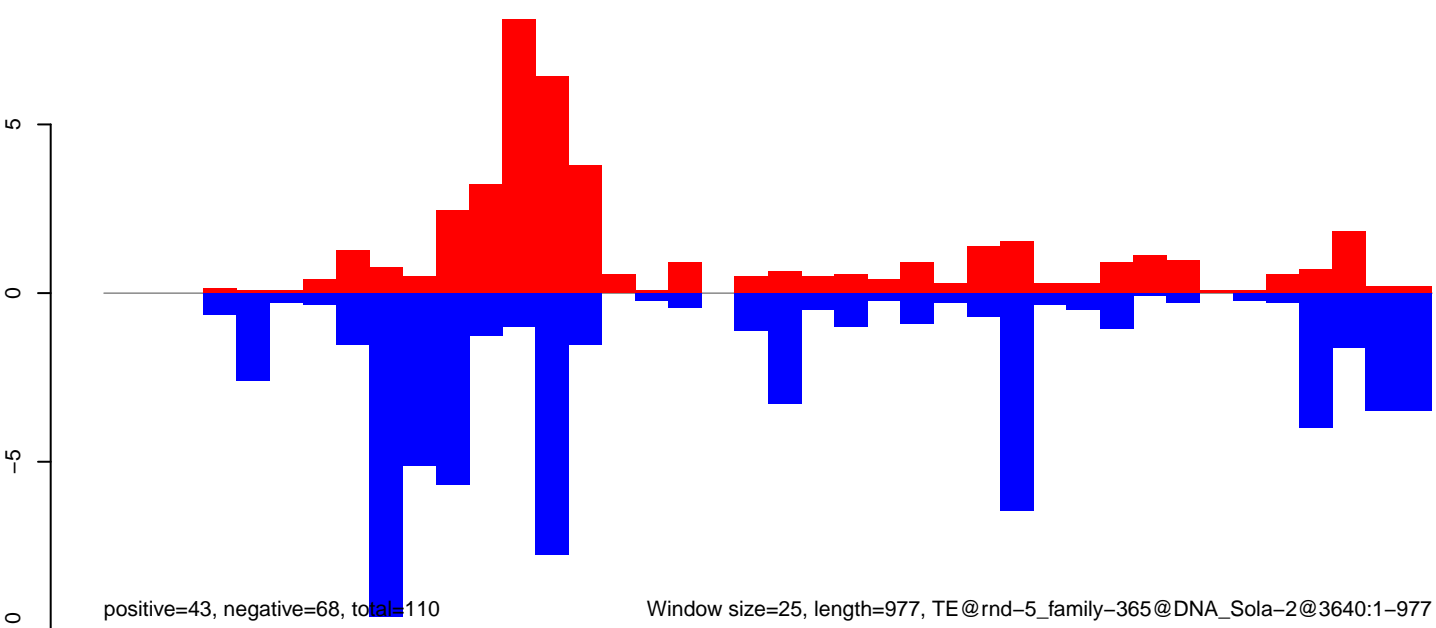
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

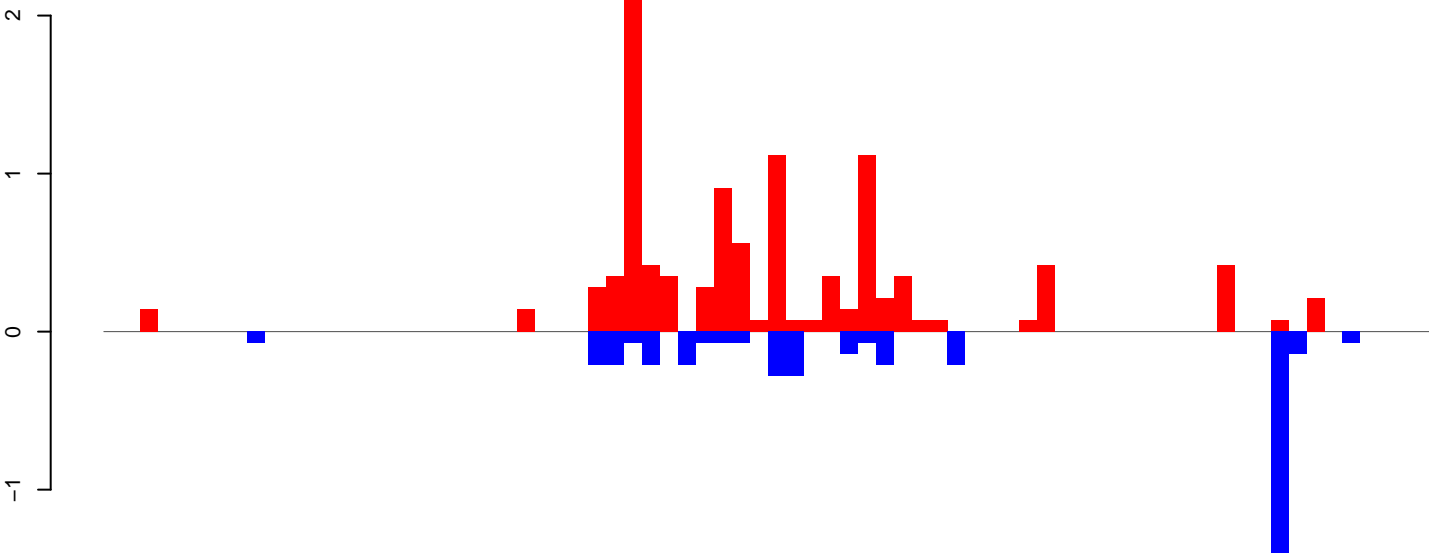


CuQuin_Mcarc_NL.rep



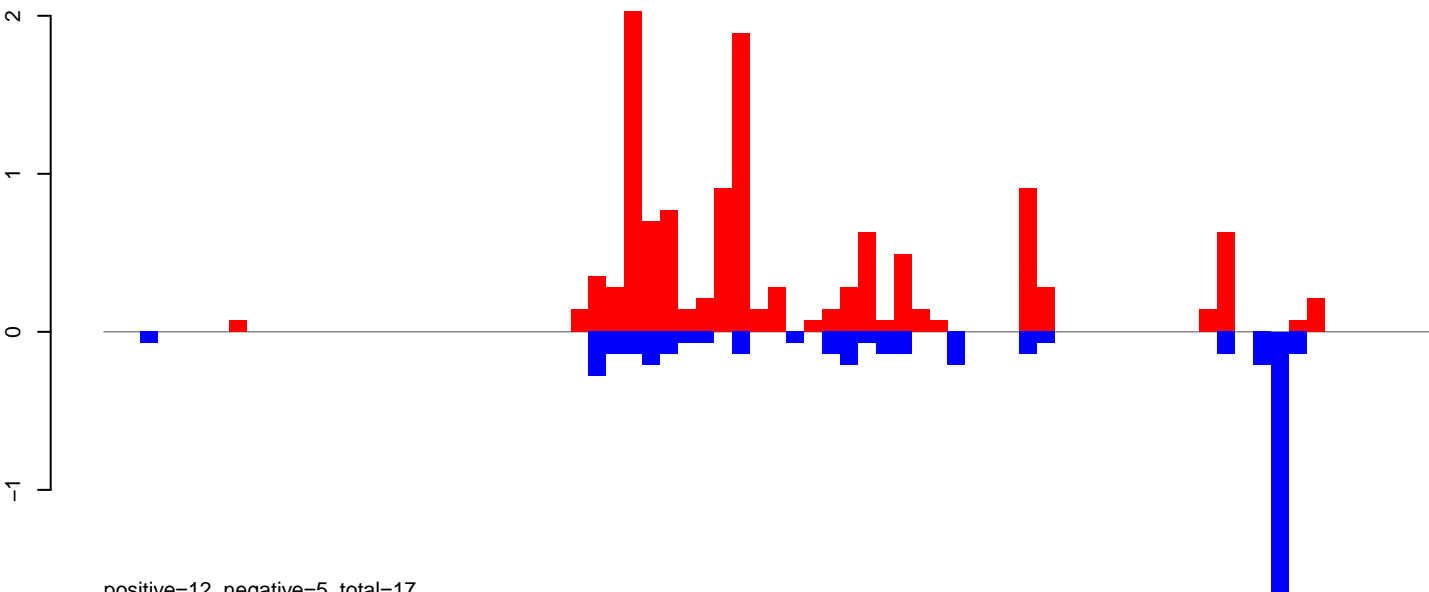
Window size=25, length=977, TE@rnd-5_family-365@DNA_Sola-2@3640:1-977

CuQuin_Mcarc_NL.18_23.rep



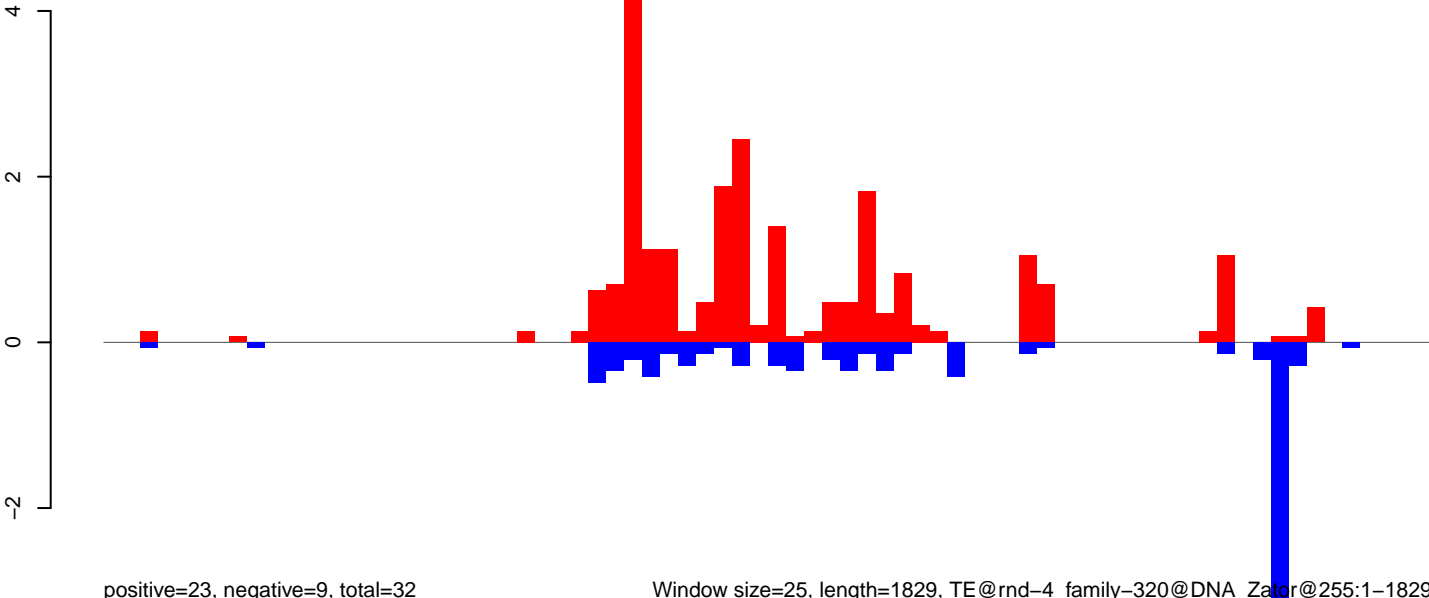
positive=10, negative=4, total=14

CuQuin_Mcarc_NL.24_35.rep



positive=12, negative=5, total=17

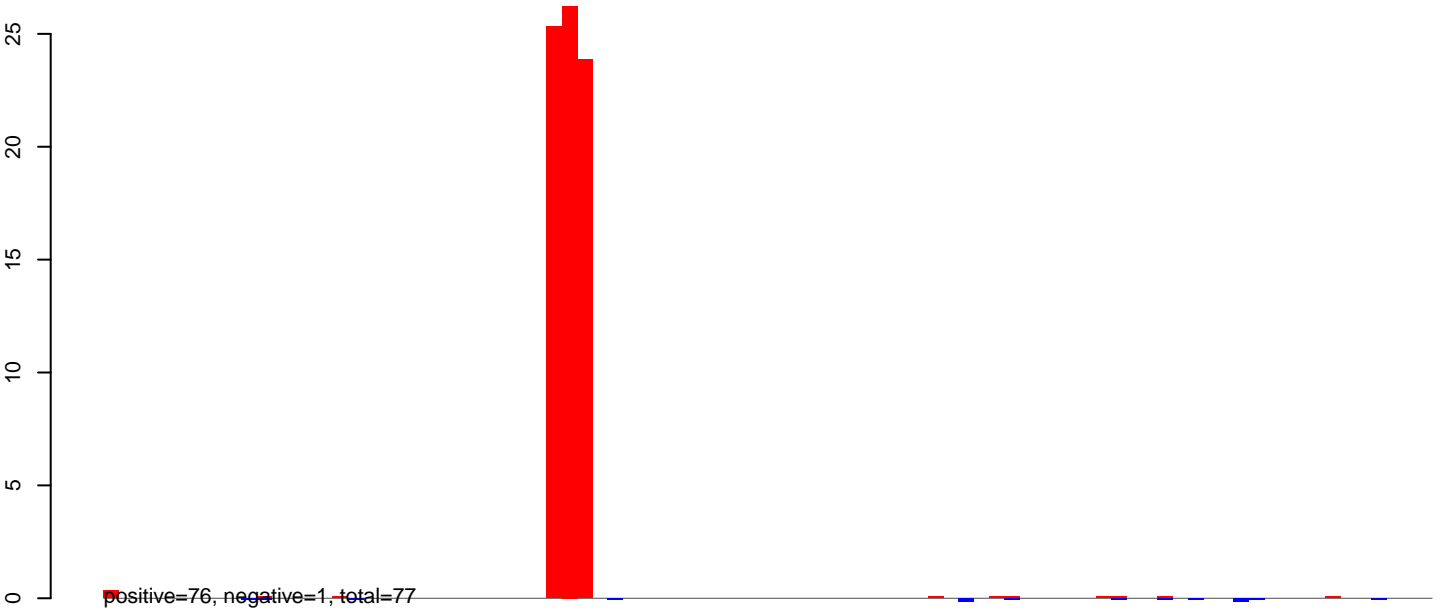
CuQuin_Mcarc_NL.rep



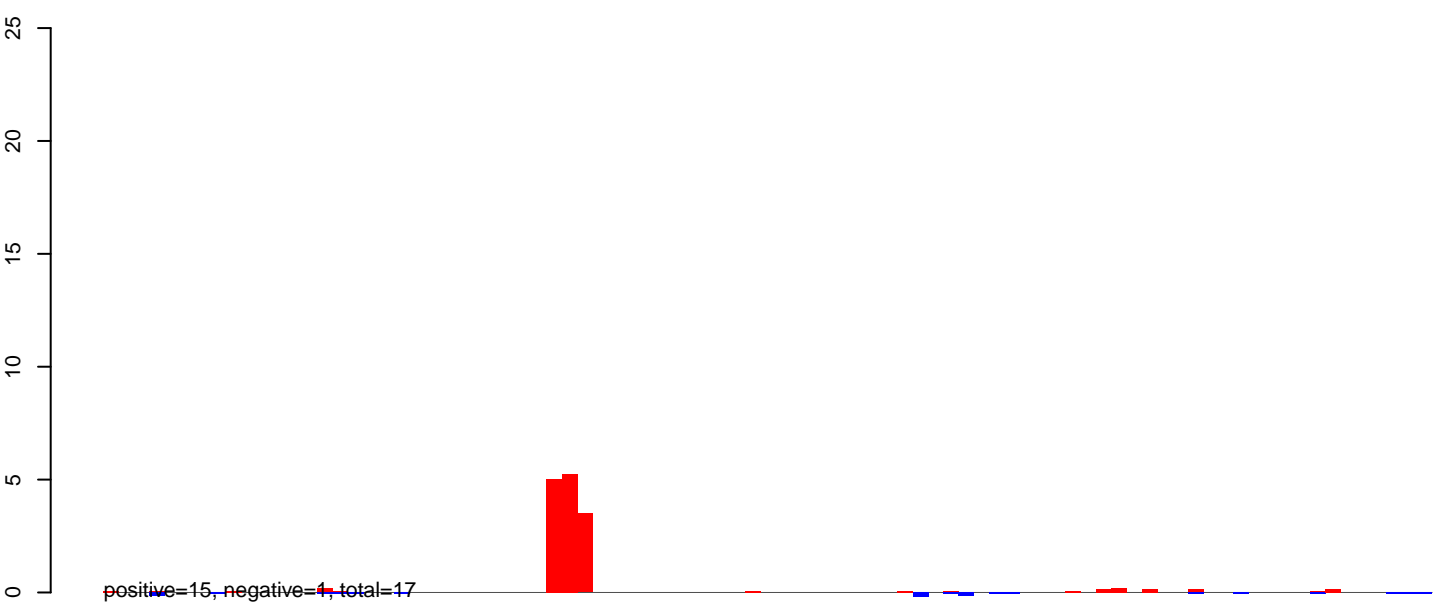
positive=23, negative=9, total=32

Window size=25, length=1829, TE@rnd-4_family-320@DNA_Zalor@255:1-1829

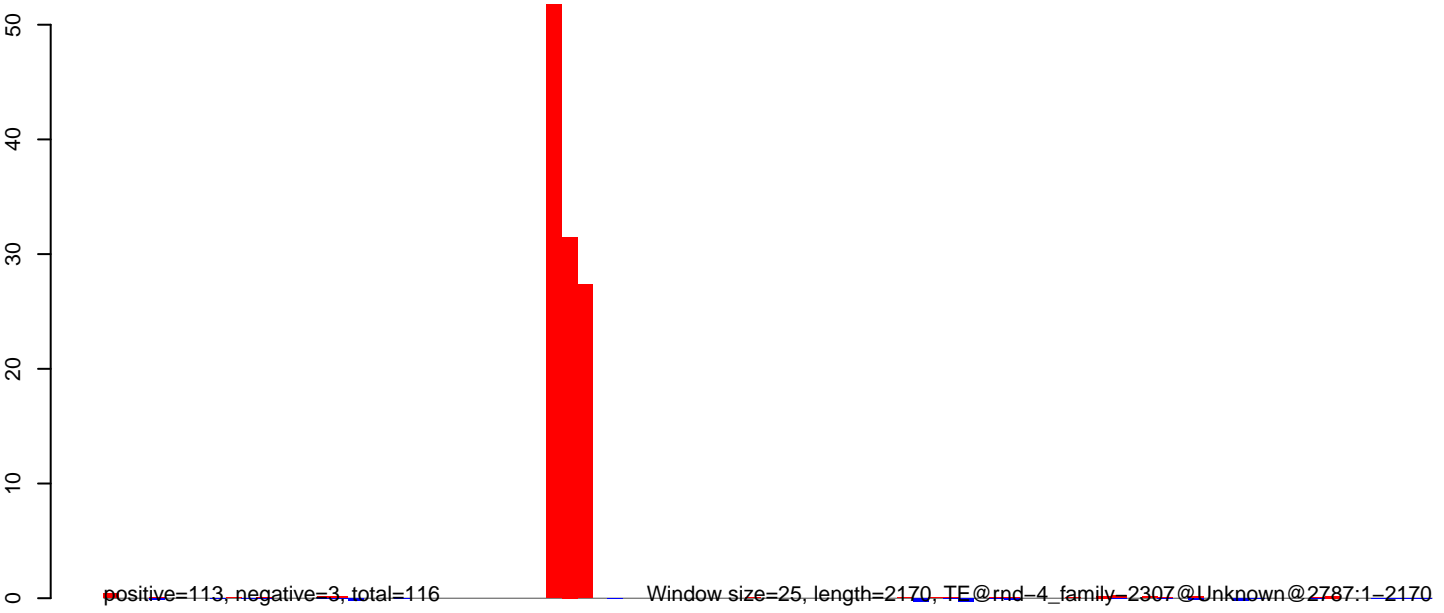
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

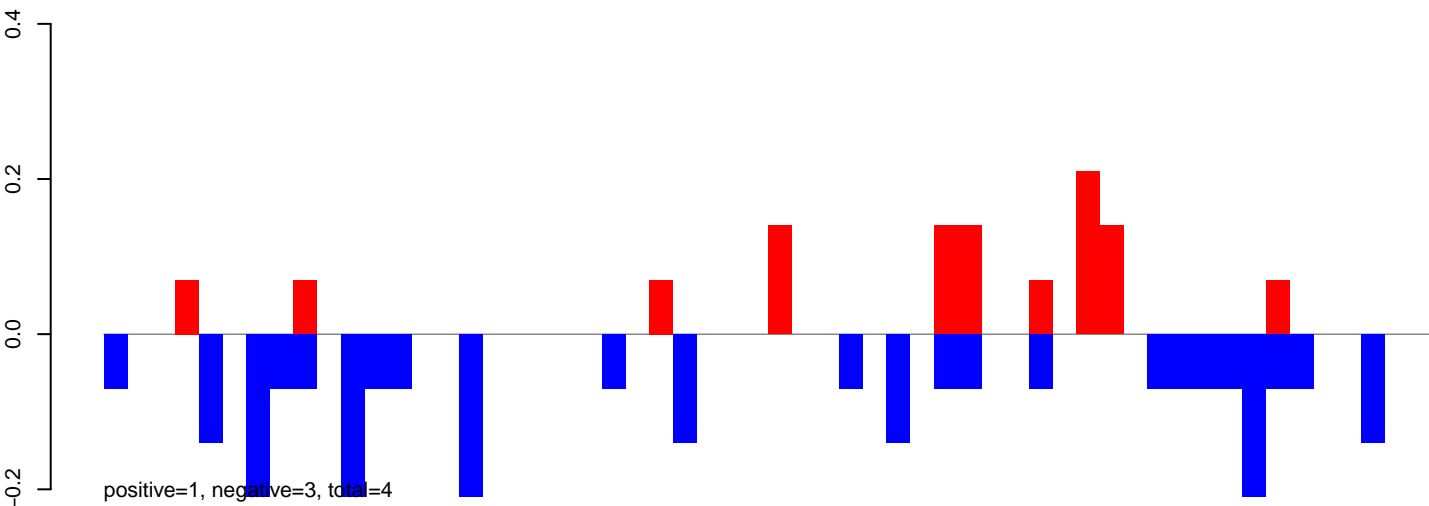


CuQuin_Mcarc_NL.rep

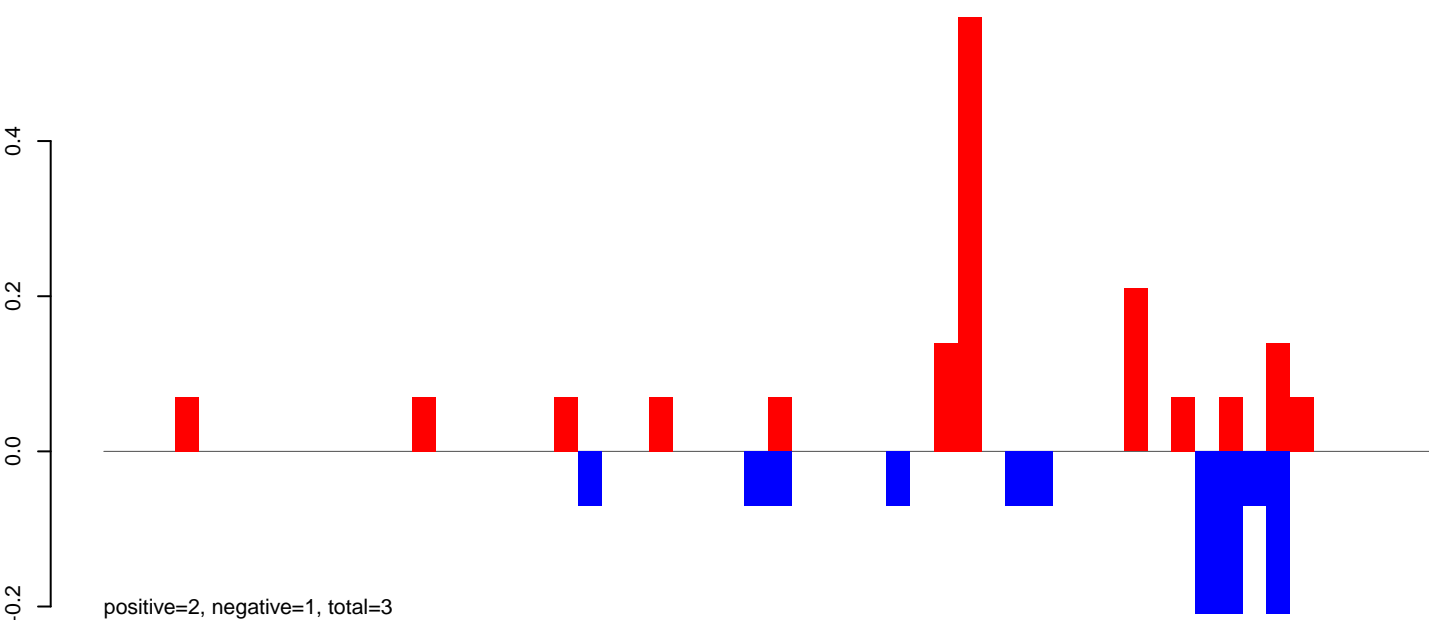


Window size=25, length=2170, TE@rnd-4_family-2307@Unknown@2787:1-2170

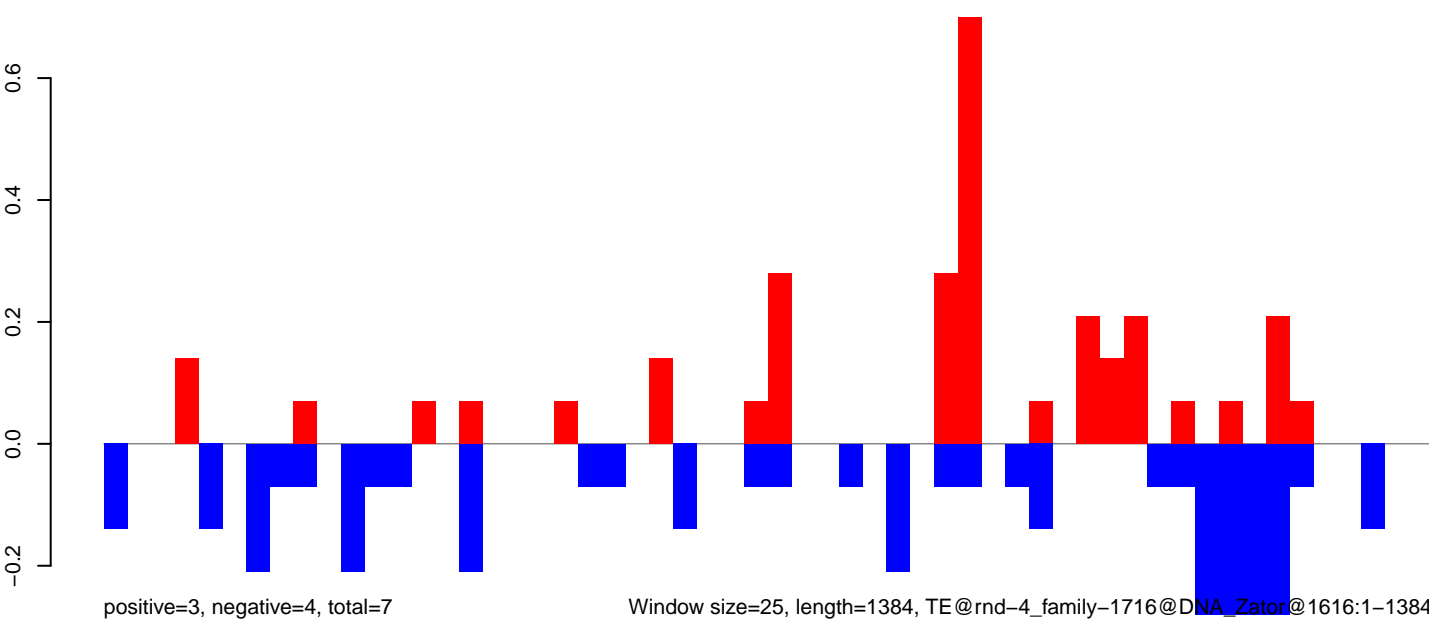
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

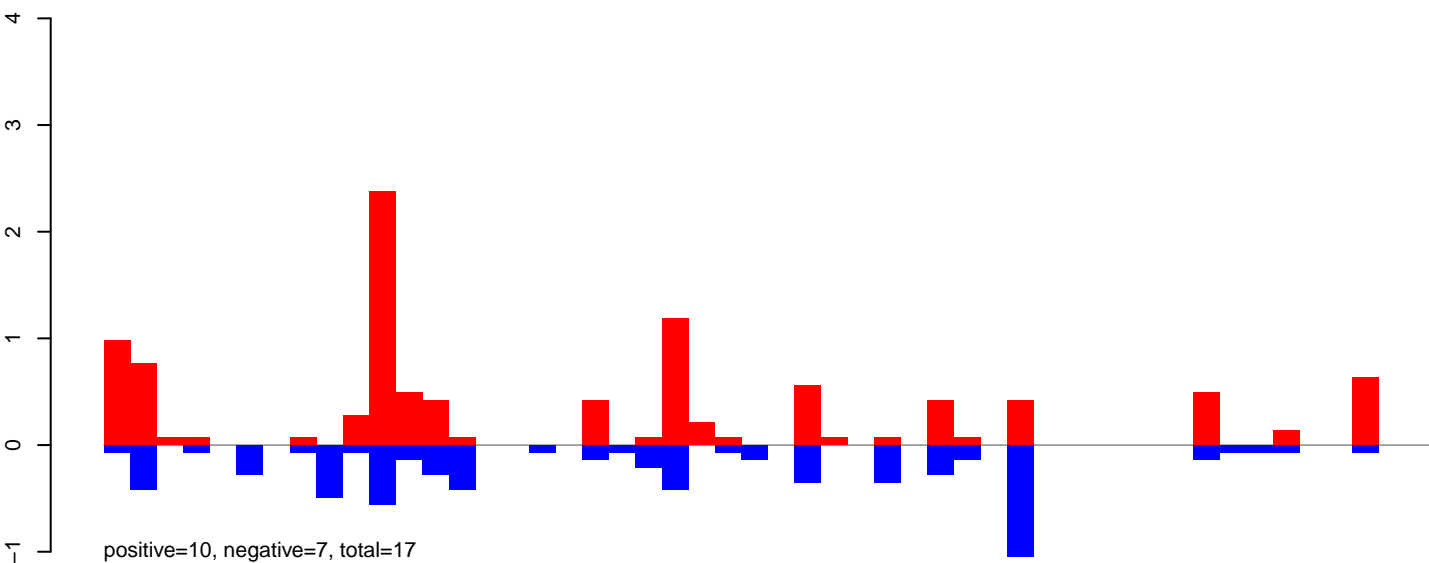


CuQuin_Mcarc_NL.rep

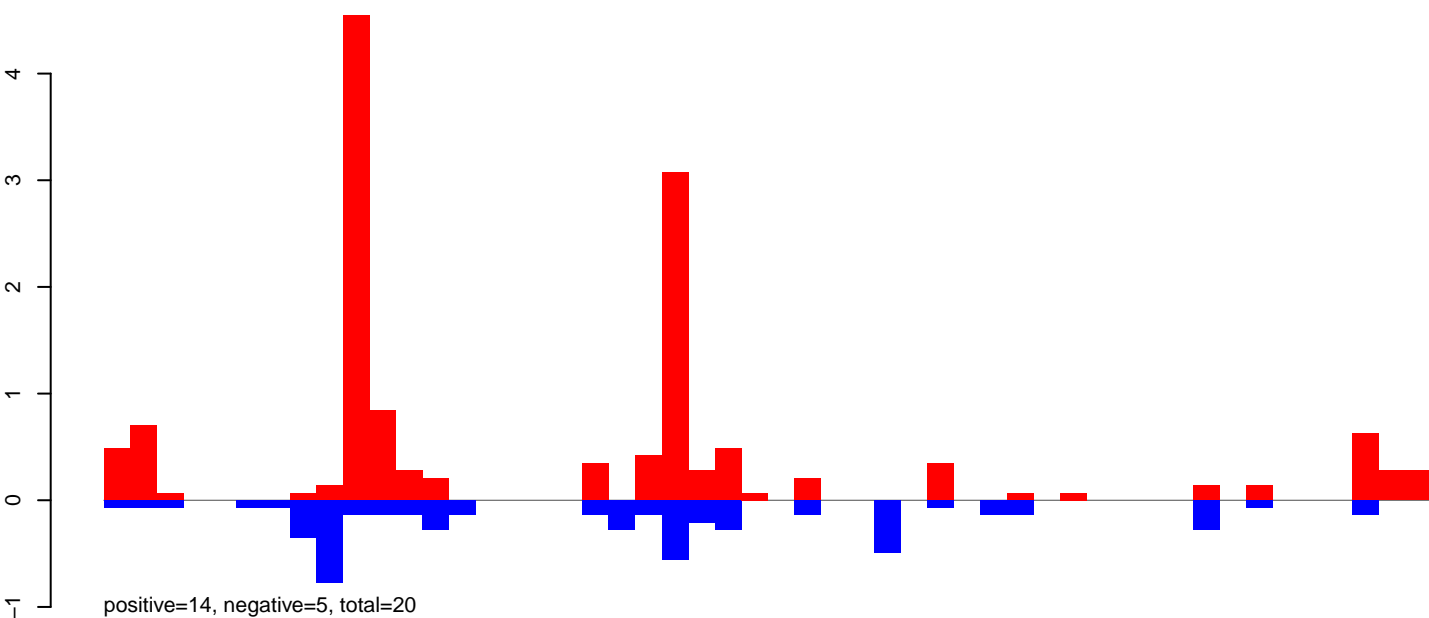


Window size=25, length=1384, TE@rnd-4_family-1716@DNA_Zator@1616:1-1384

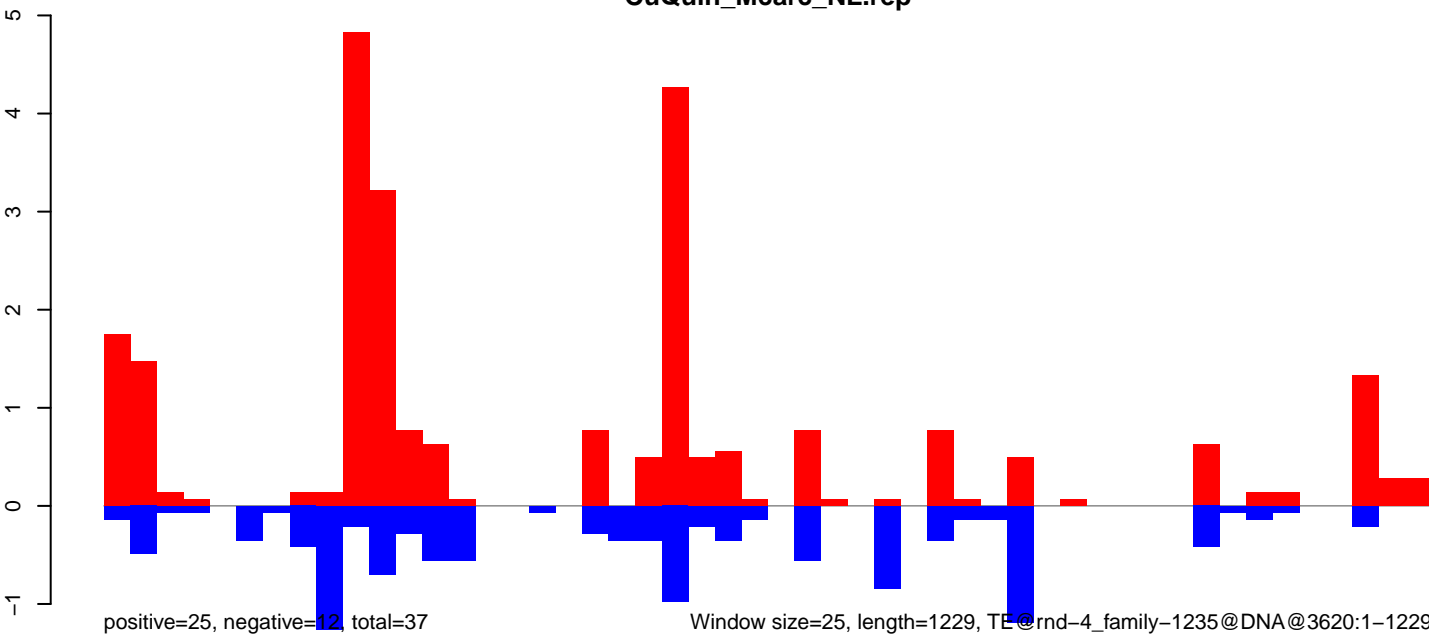
CuQuin_Mcarc_NL.18_23.rep



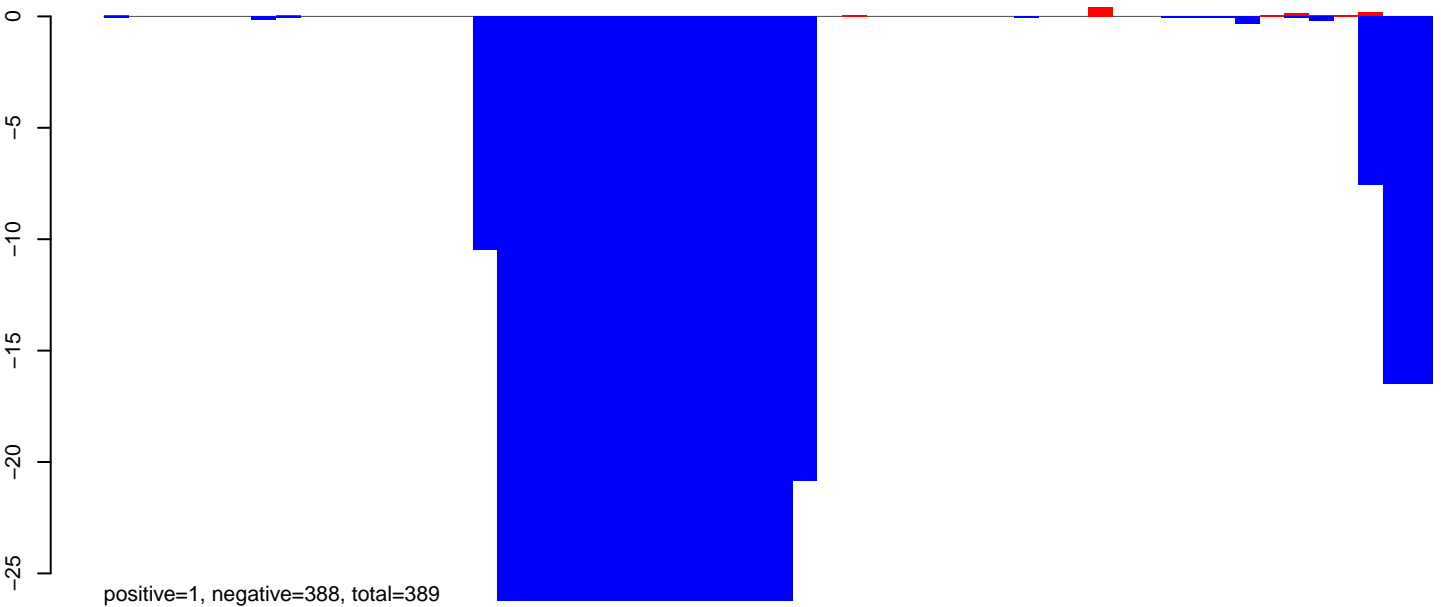
CuQuin_Mcarc_NL.24_35.rep



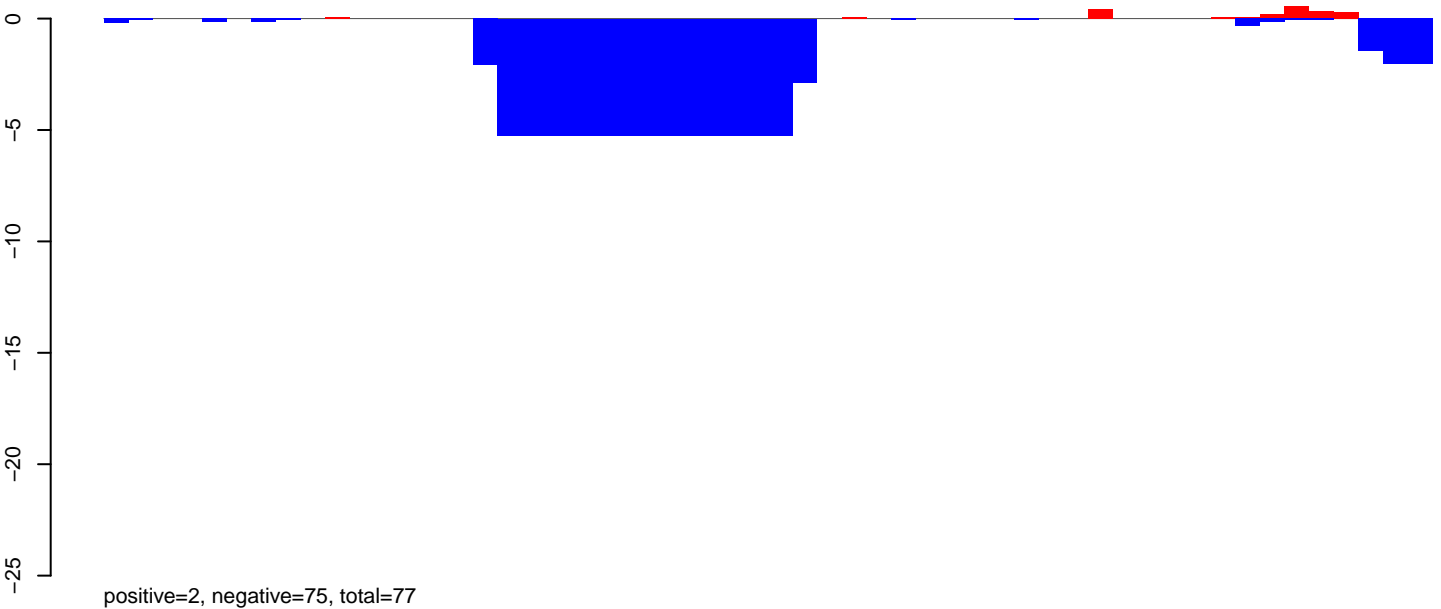
CuQuin_Mcarc_NL.rep



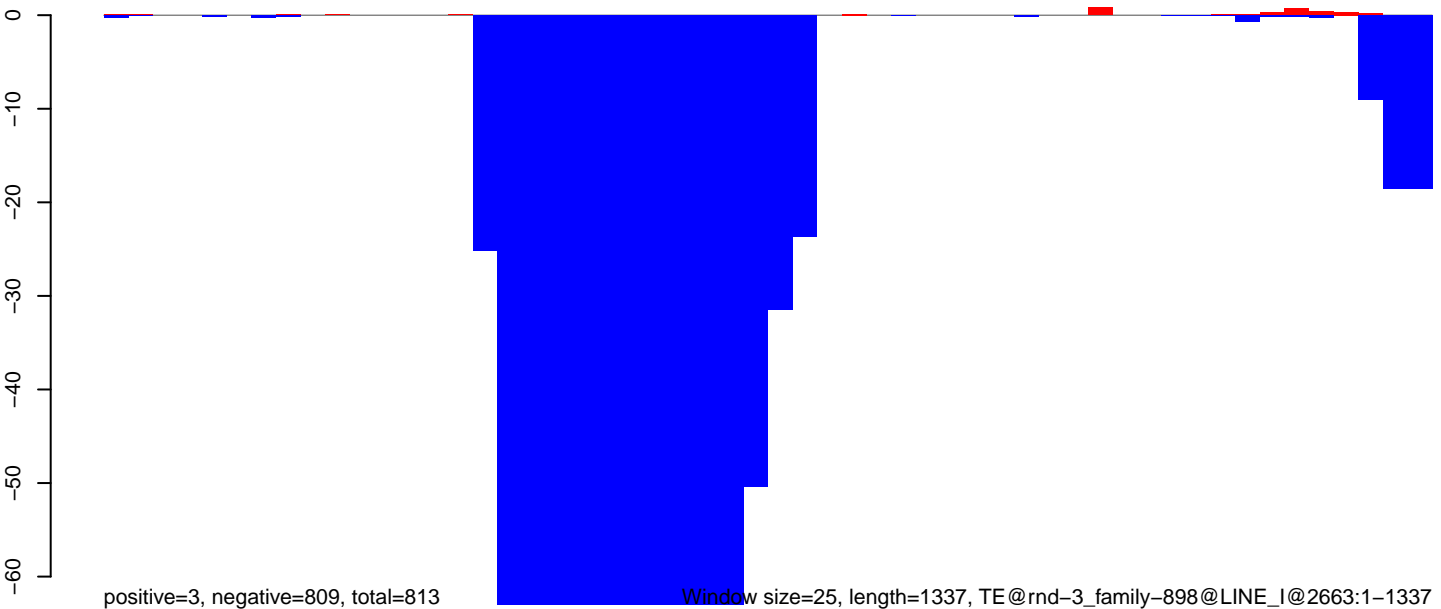
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



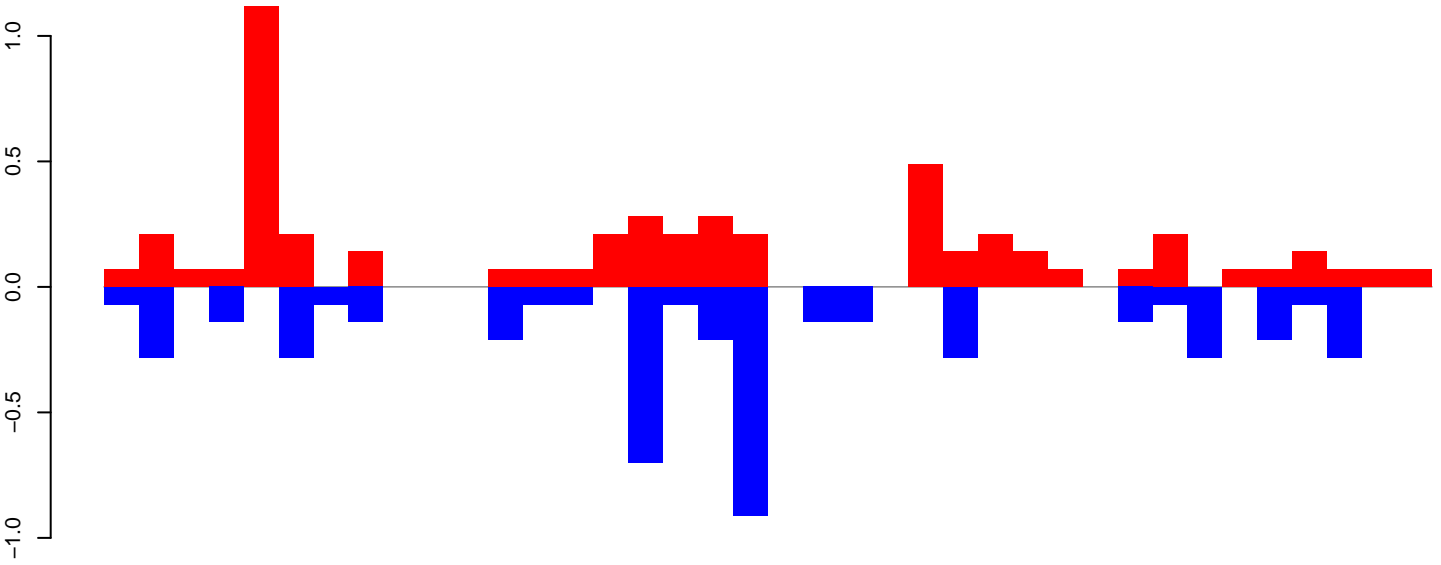
CuQuin_Mcarc_NL.rep



Window size=25, length=1337, TE@rnd-3_family-898@LINE_I@2663:1-1337

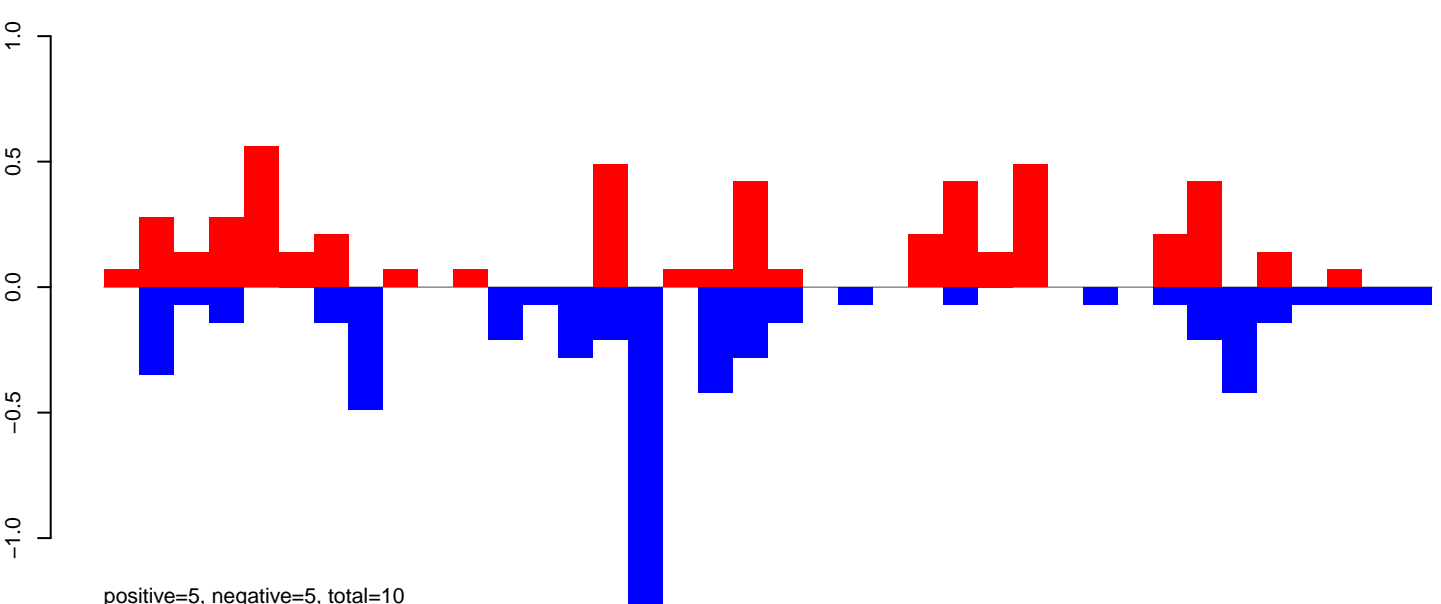
0 200 400 600 800 1000 1200 1400

CuQuin_Mcarc_NL.18_23.rep



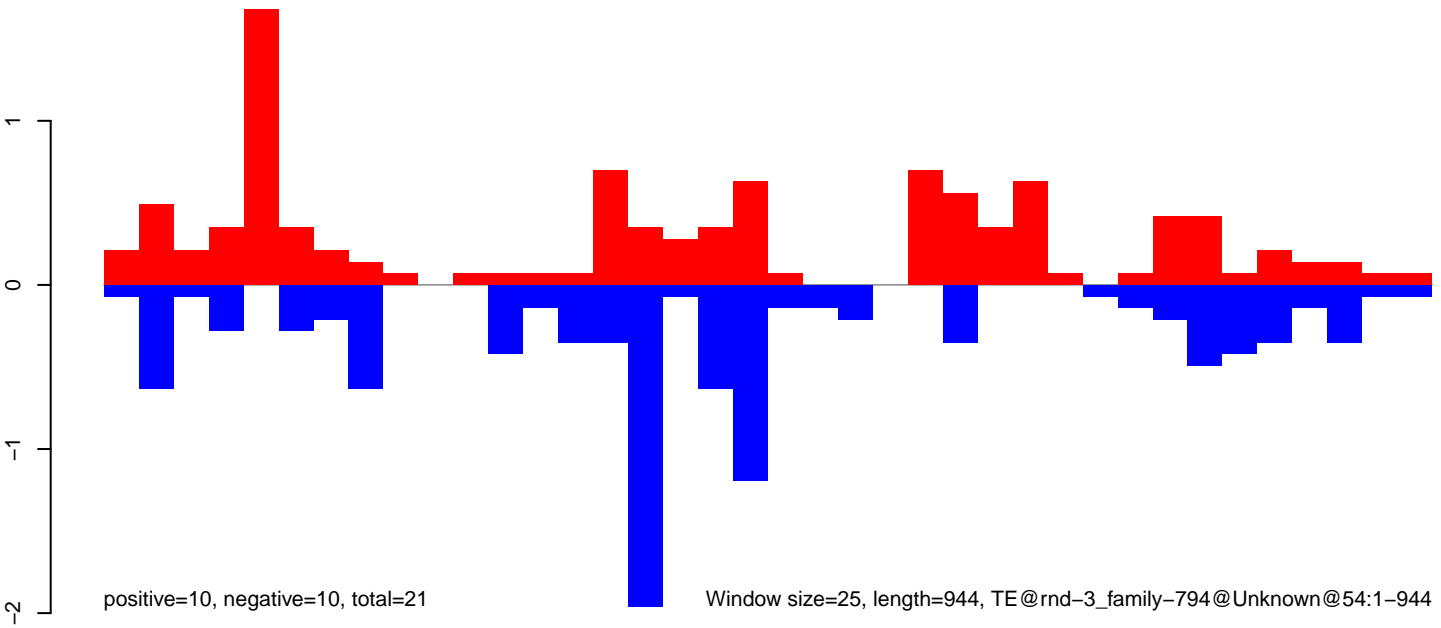
positive=5, negative=5, total=10

CuQuin_Mcarc_NL.24_35.rep



positive=5, negative=5, total=10

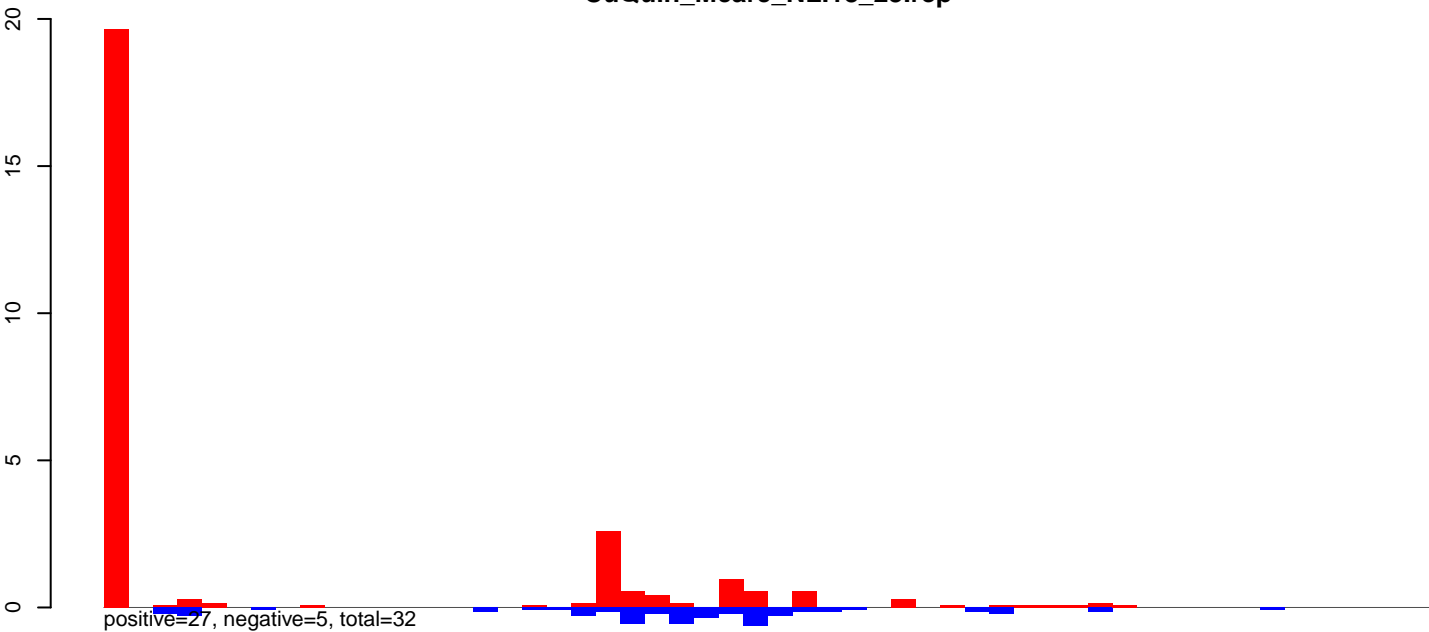
CuQuin_Mcarc_NL.rep



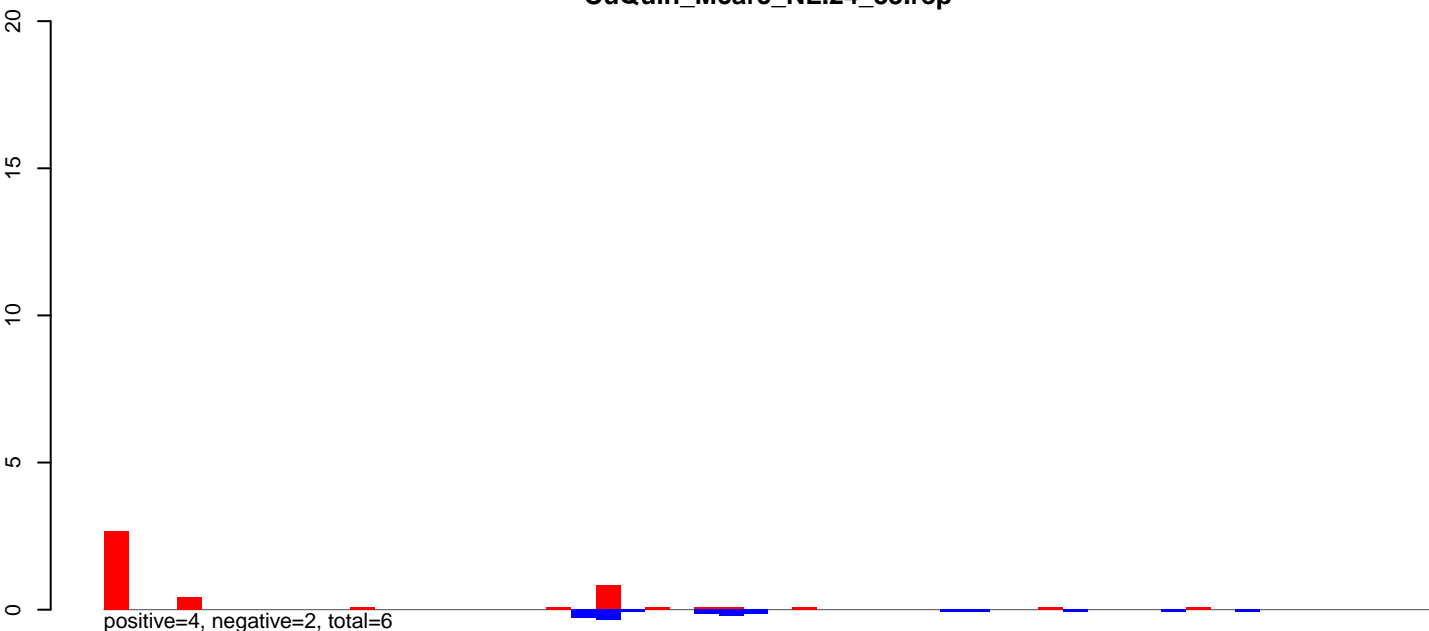
positive=10, negative=10, total=21

Window size=25, length=944, TE@rnd-3_family-794@Unknown@54:1-944

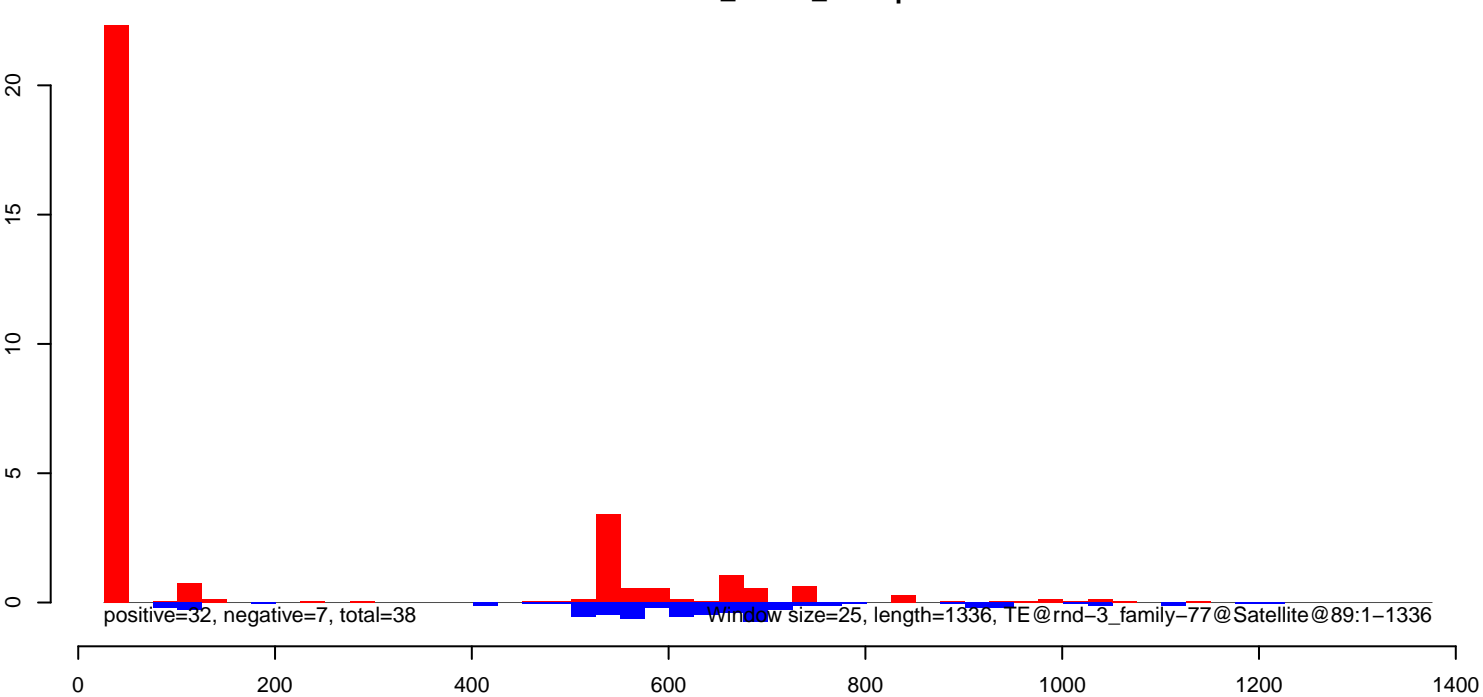
CuQuin_Mcarc_NL.18_23.rep



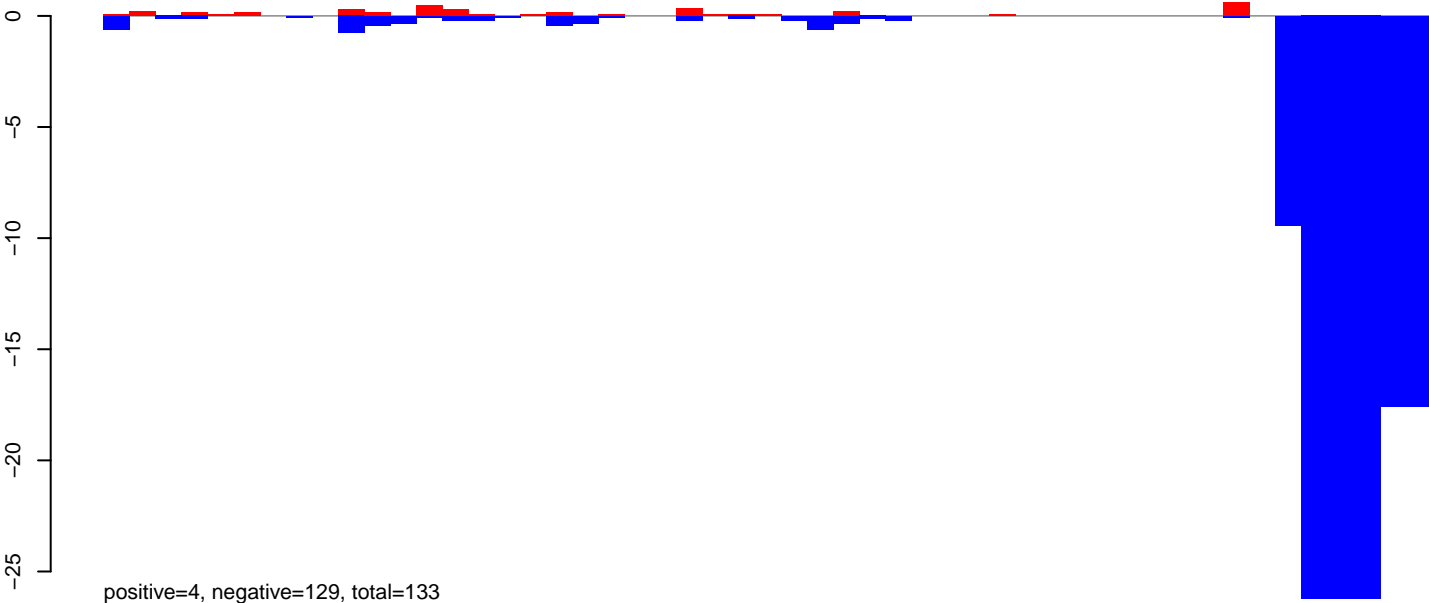
CuQuin_Mcarc_NL.24_35.rep



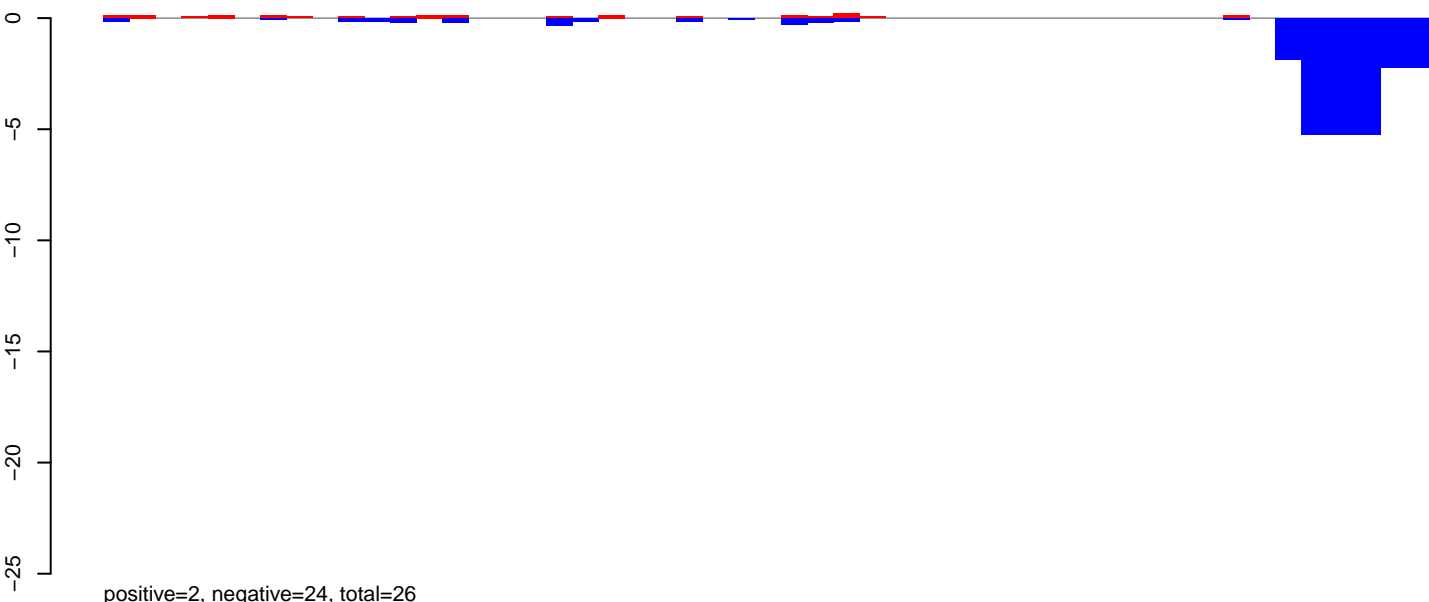
CuQuin_Mcarc_NL.rep



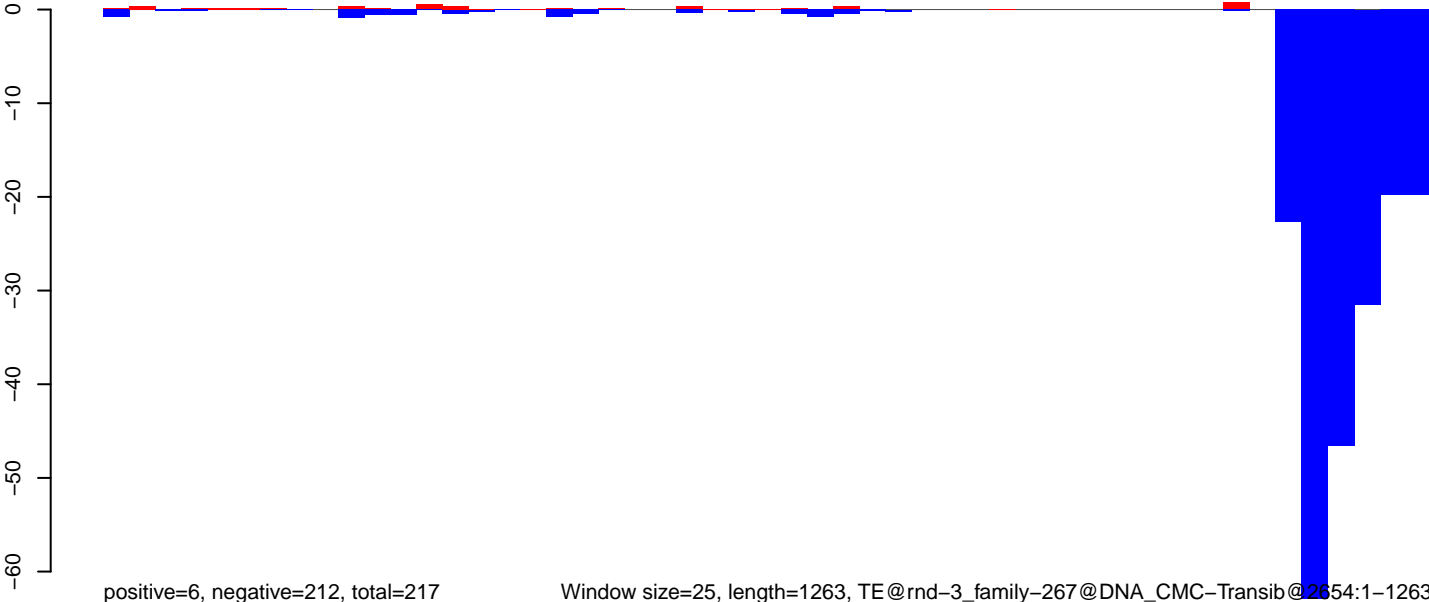
CuQuin_Mcarc_NL.18_23.rep



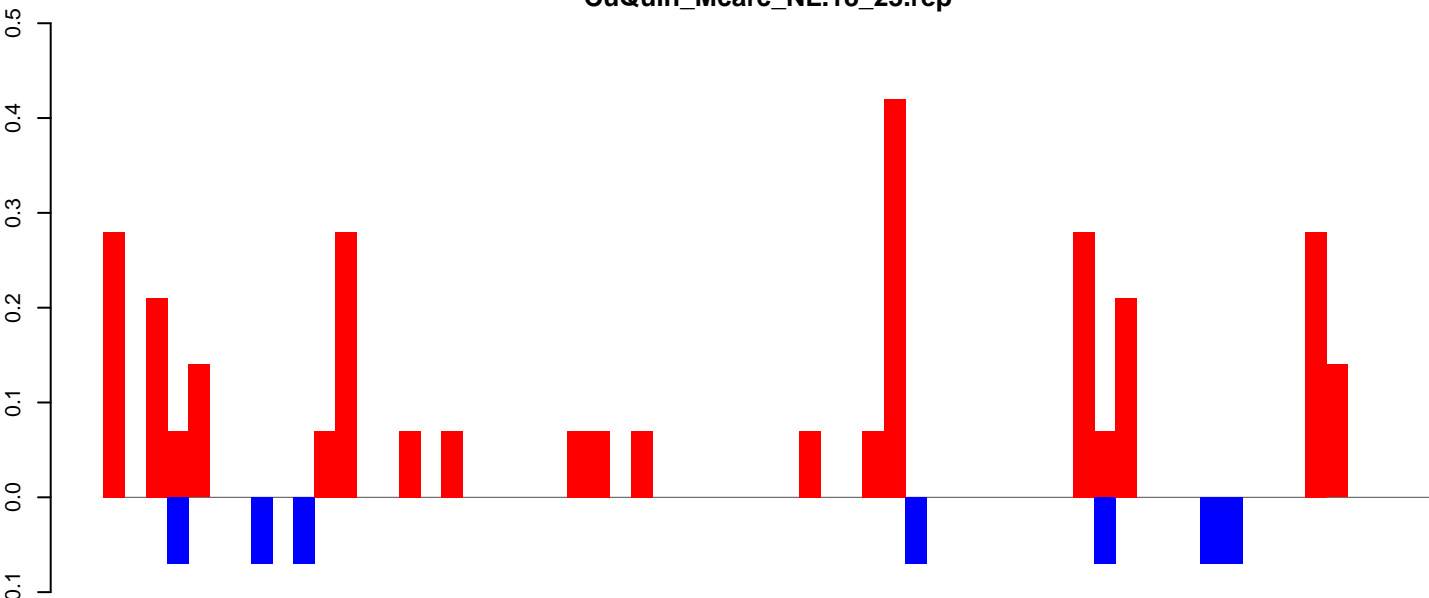
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

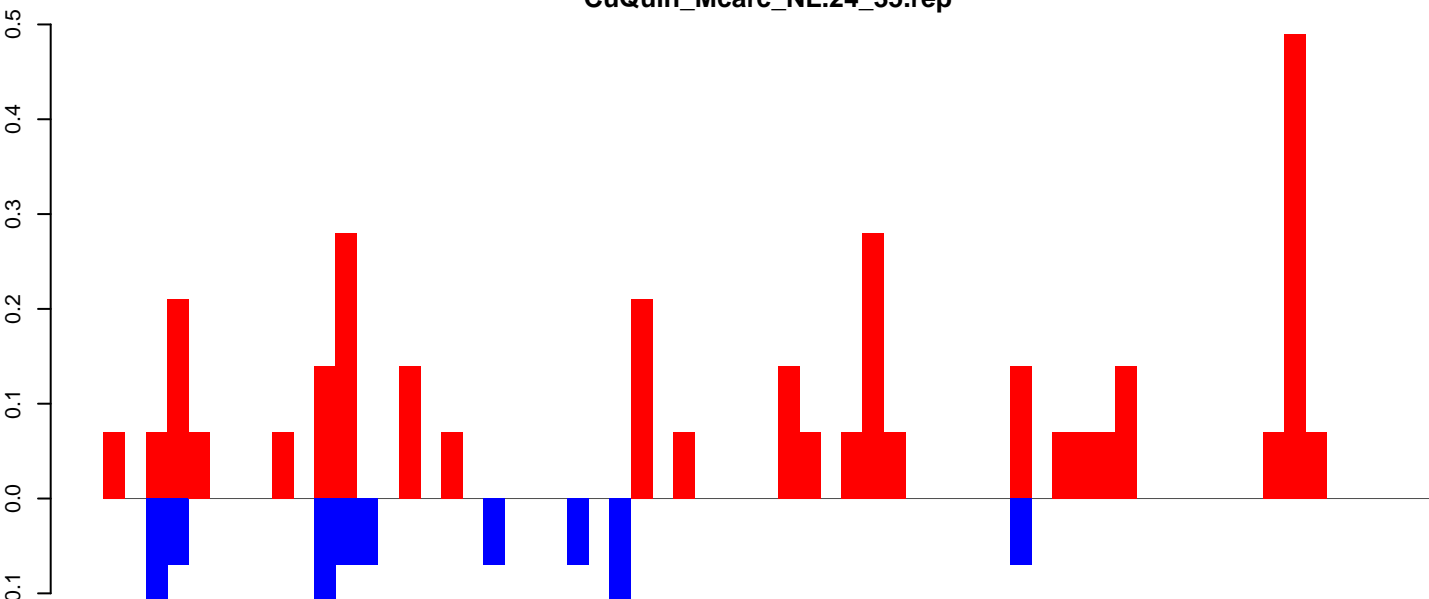


CuQuin_Mcarc_NL.18_23.rep



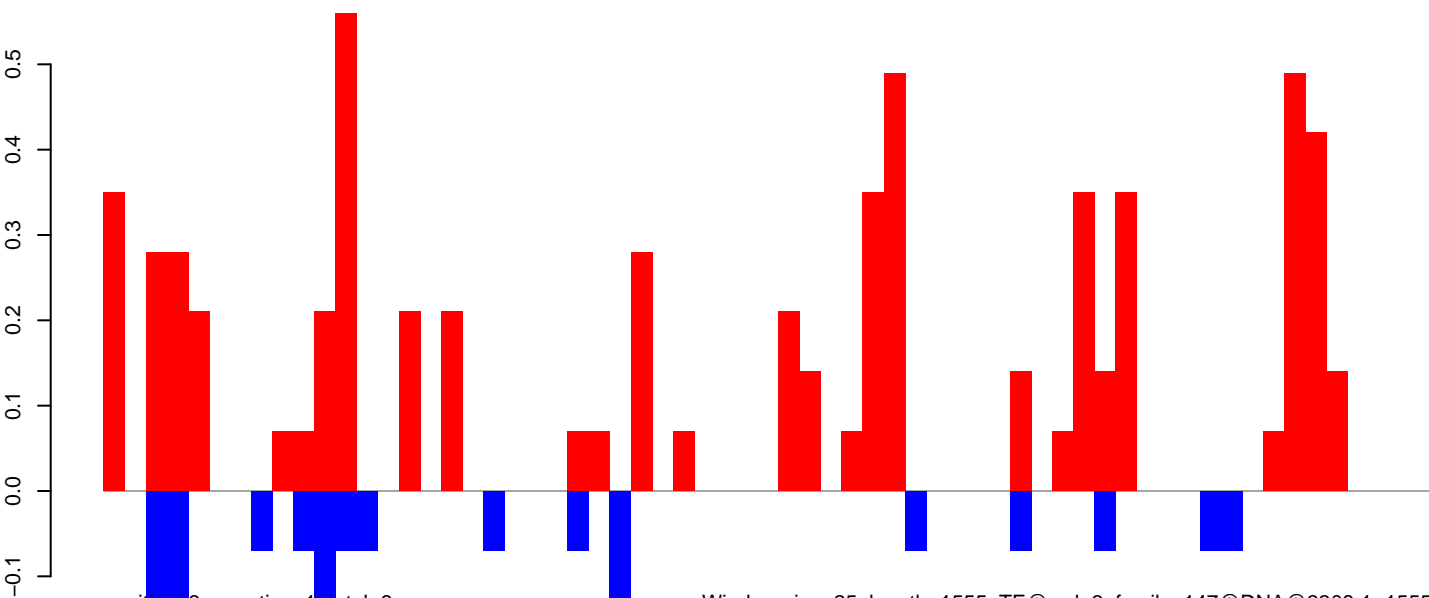
positive=3, negative=0, total=3

CuQuin_Mcarc_NL.24_35.rep



positive=3, negative=1, total=4

CuQuin_Mcarc_NL.rep

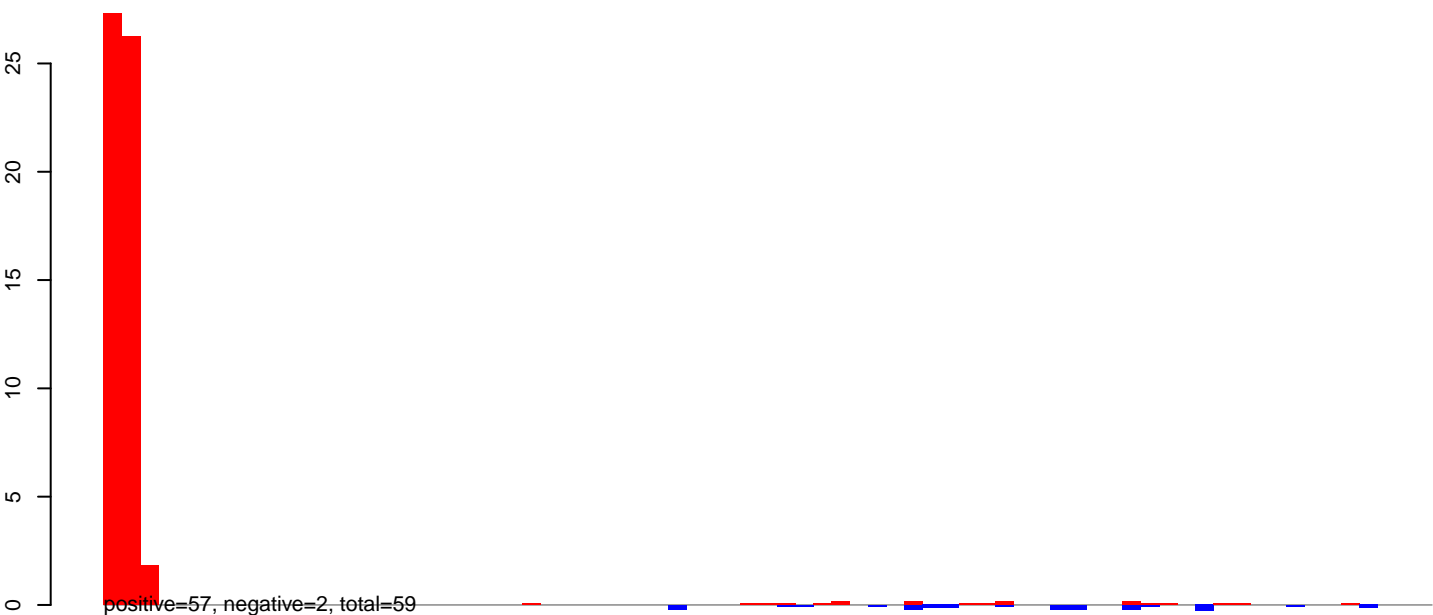


positive=6, negative=1, total=8

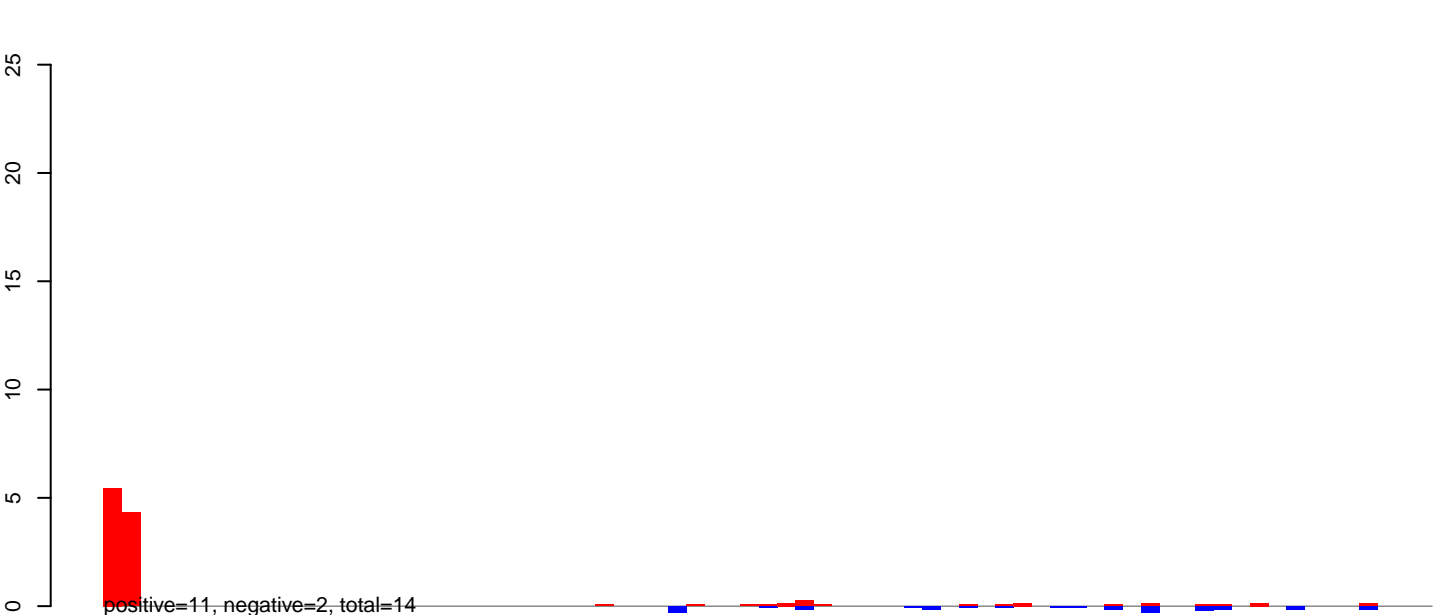
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0 500 1000 1500

CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

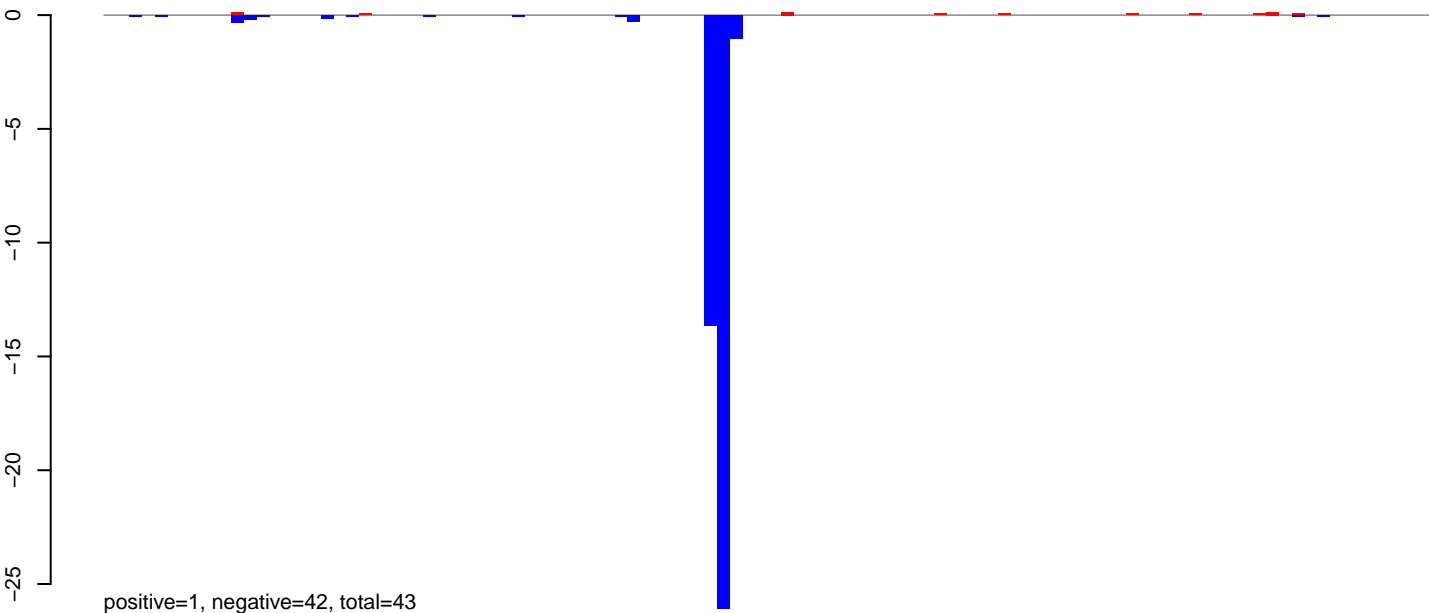


CuQuin_Mcarc_NL.rep

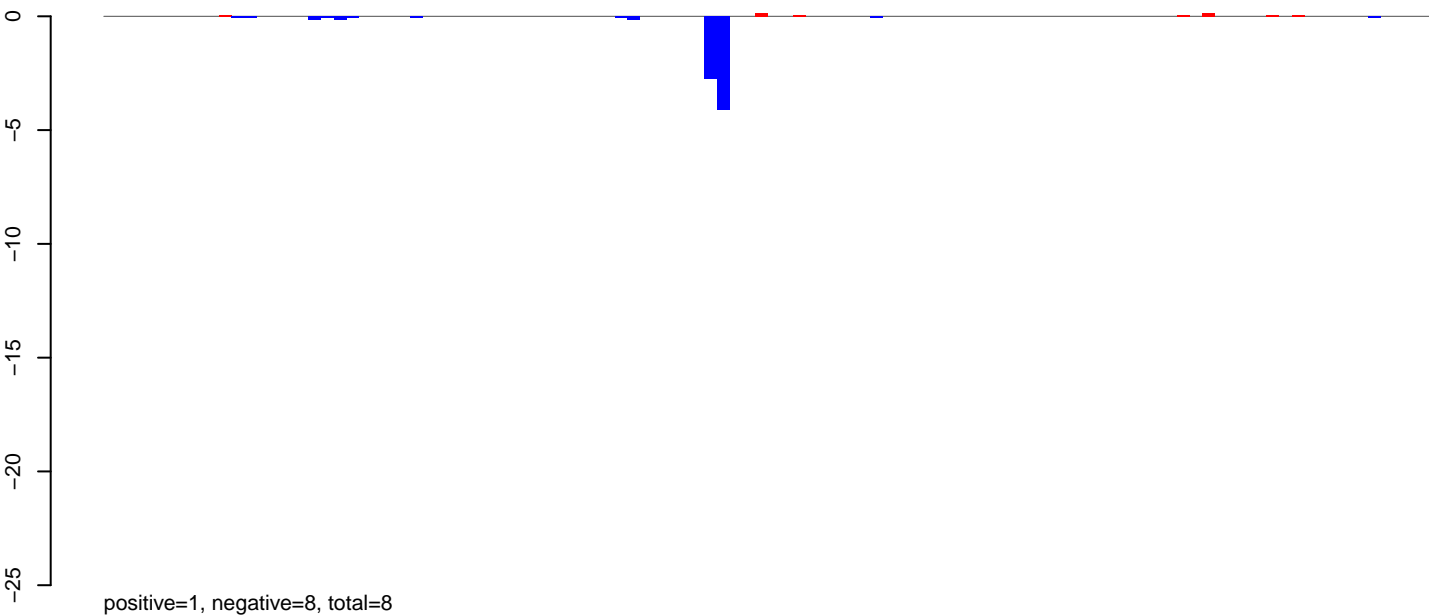


Window size=25, length=1819, TE@rnd=5, family=799@DNA_Zator@2394.4=1819

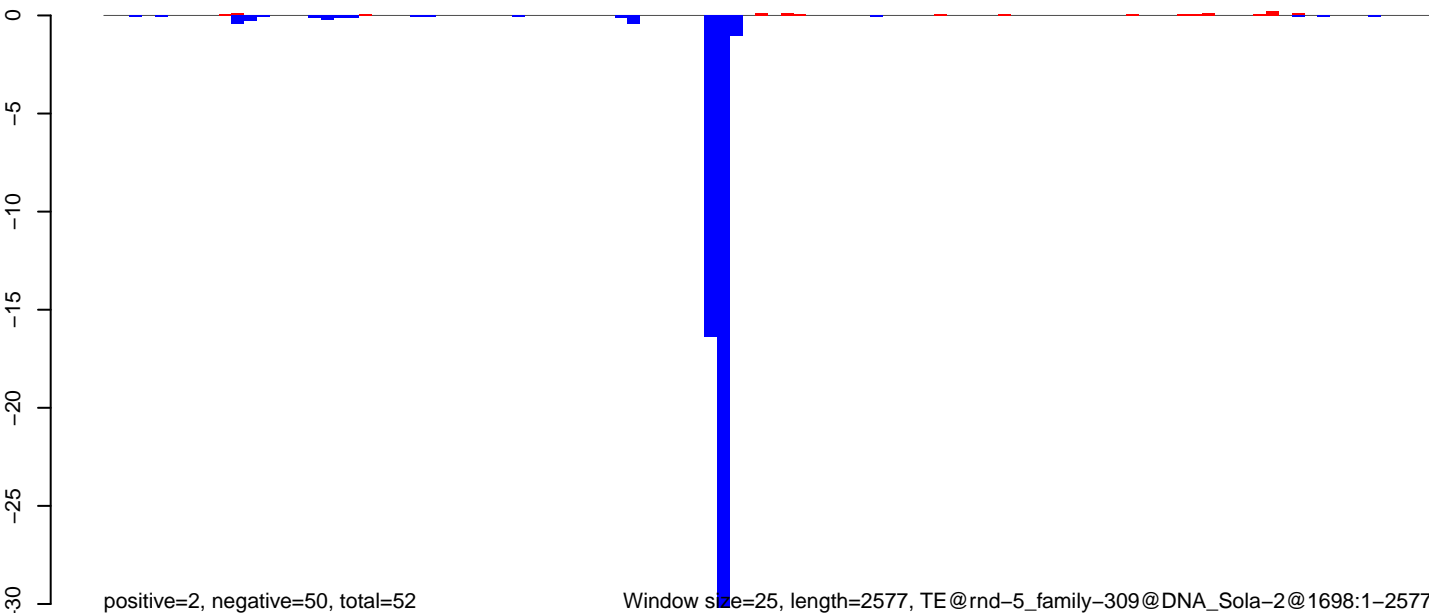
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



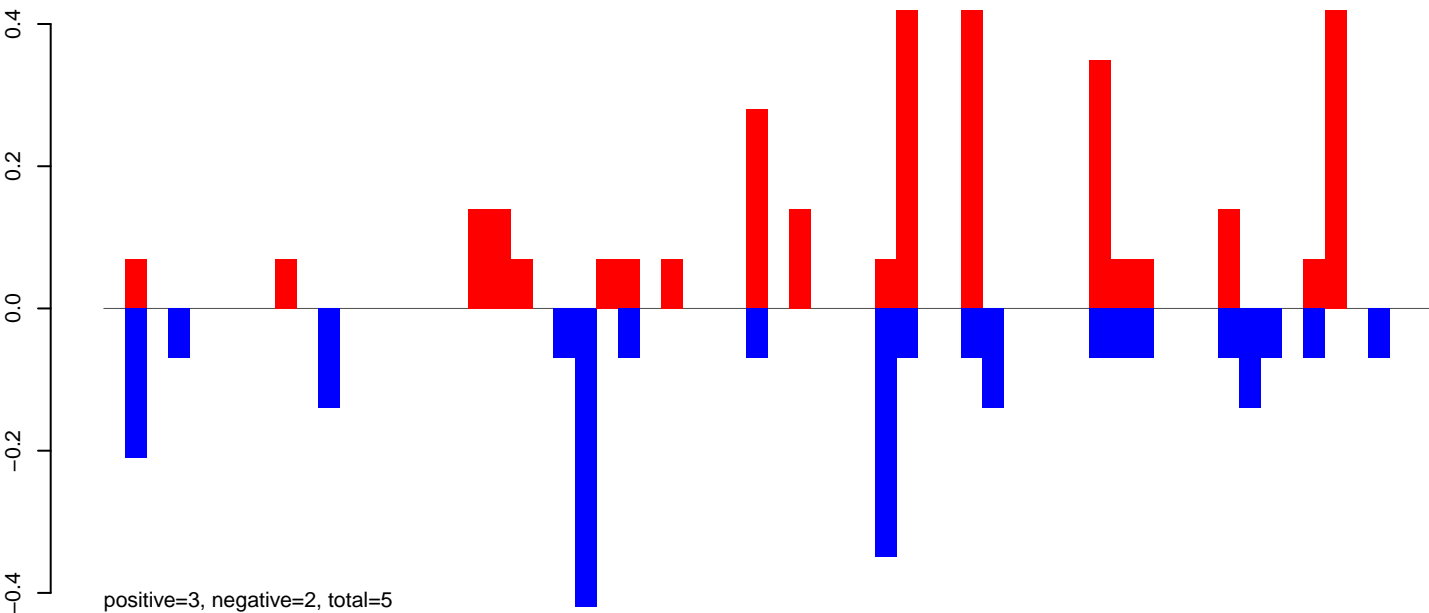
CuQuin_Mcarc_NL.rep



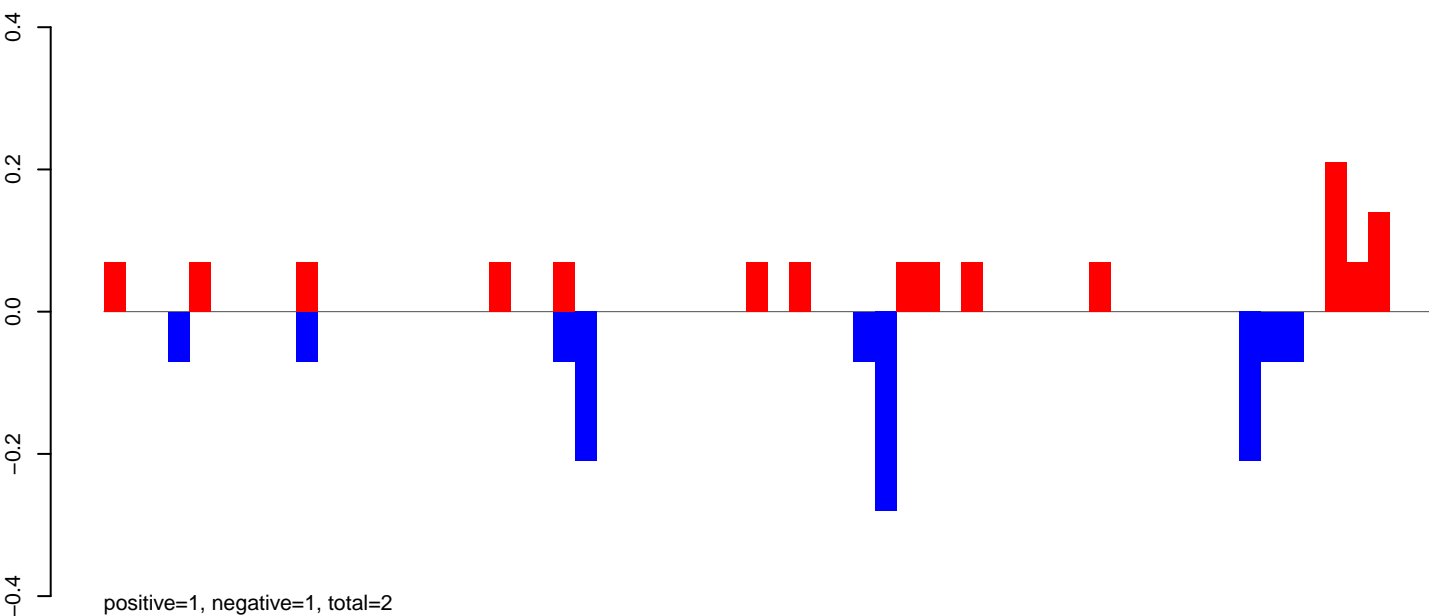
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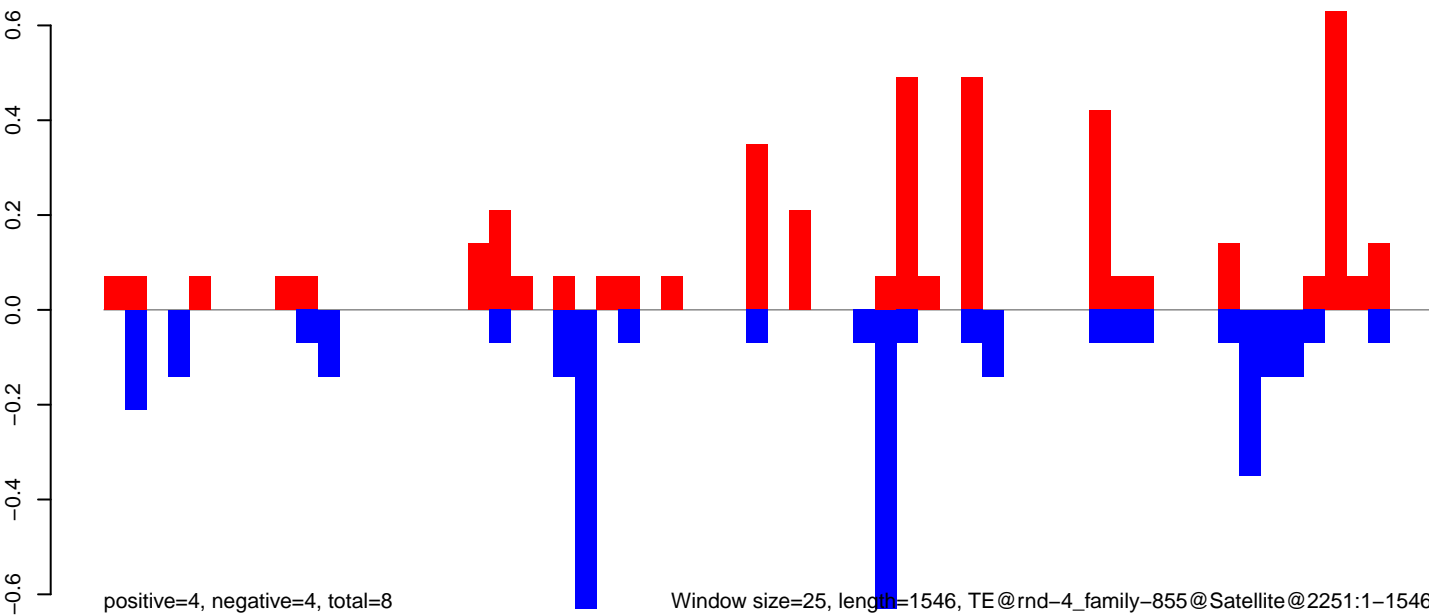
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



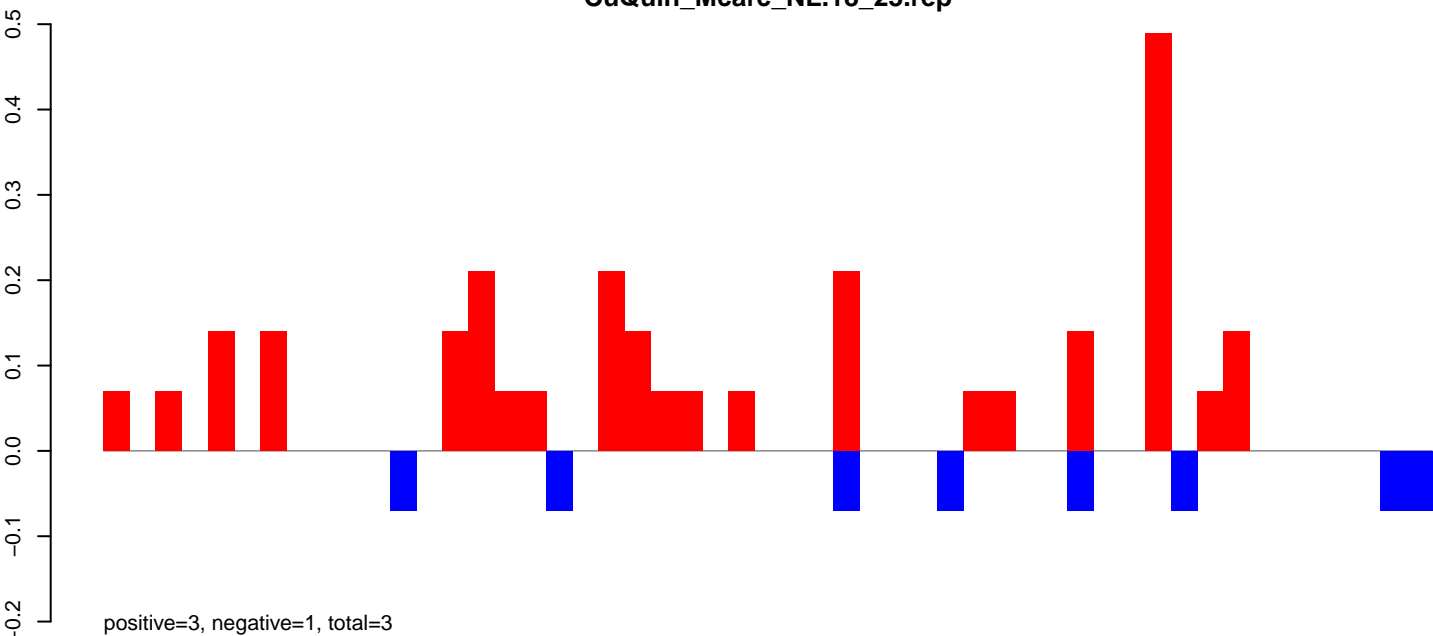
CuQuin_Mcarc_NL.rep



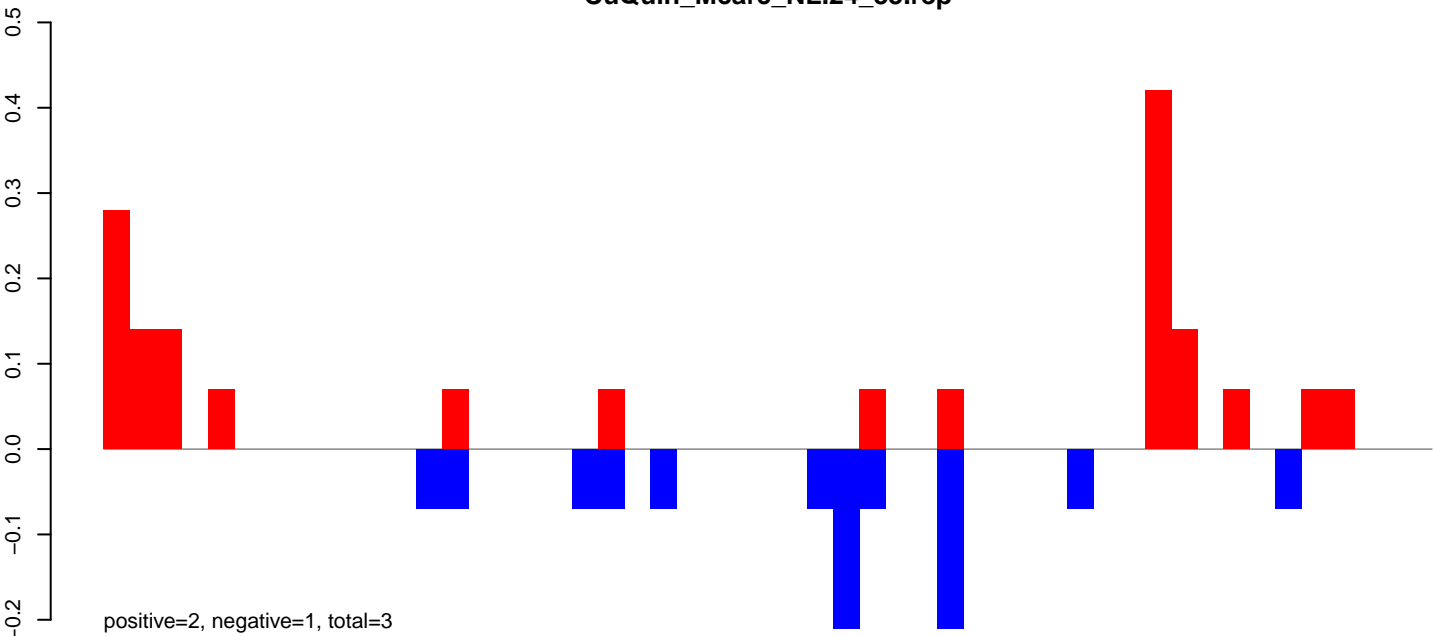
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0 500 1000 1500

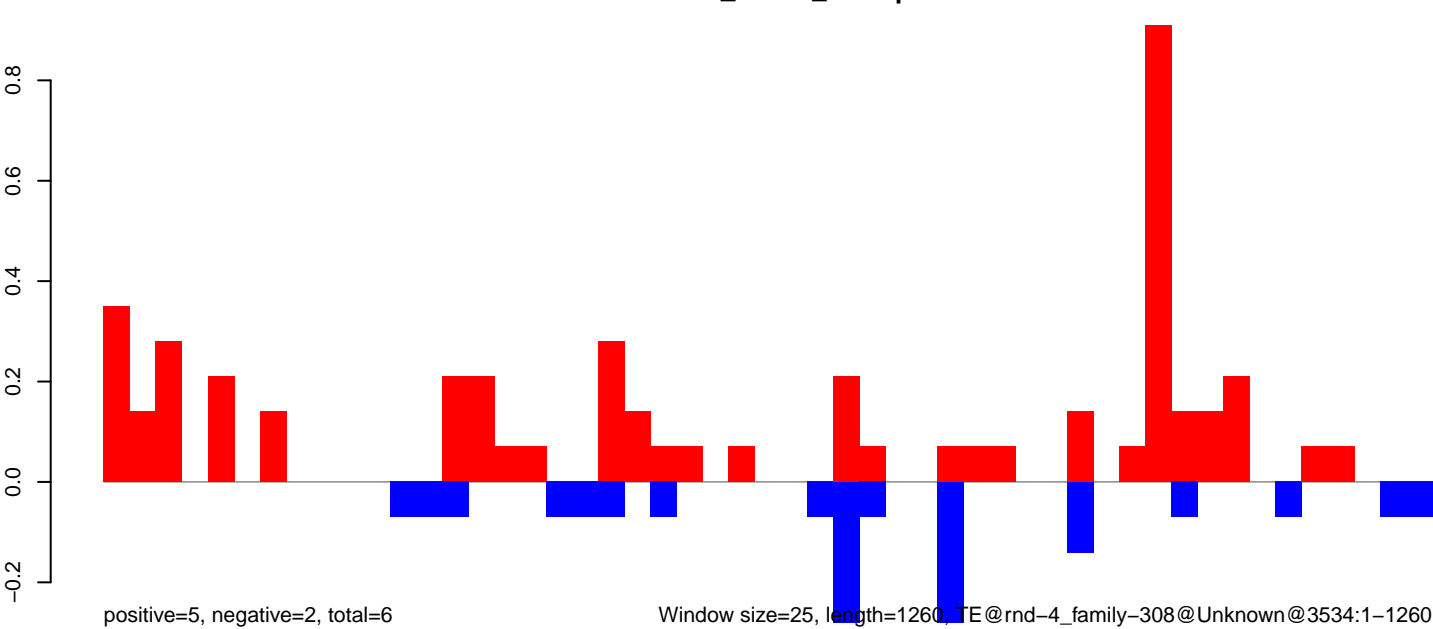
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

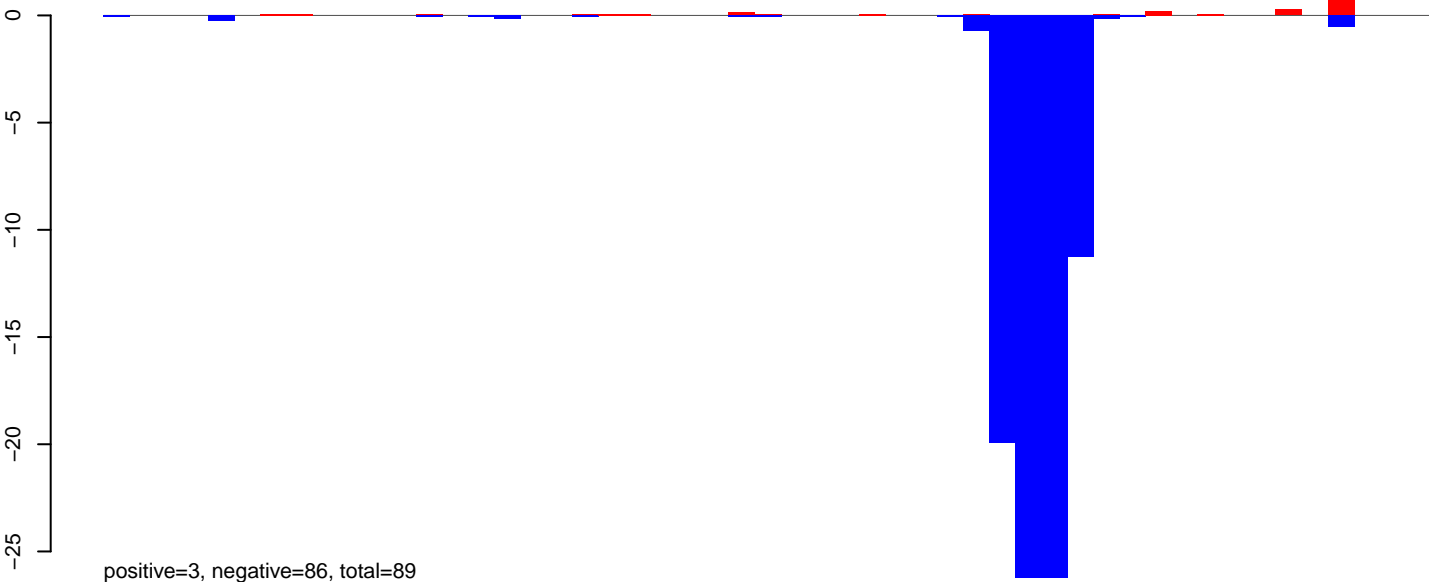


CuQuin_Mcarc_NL.rep

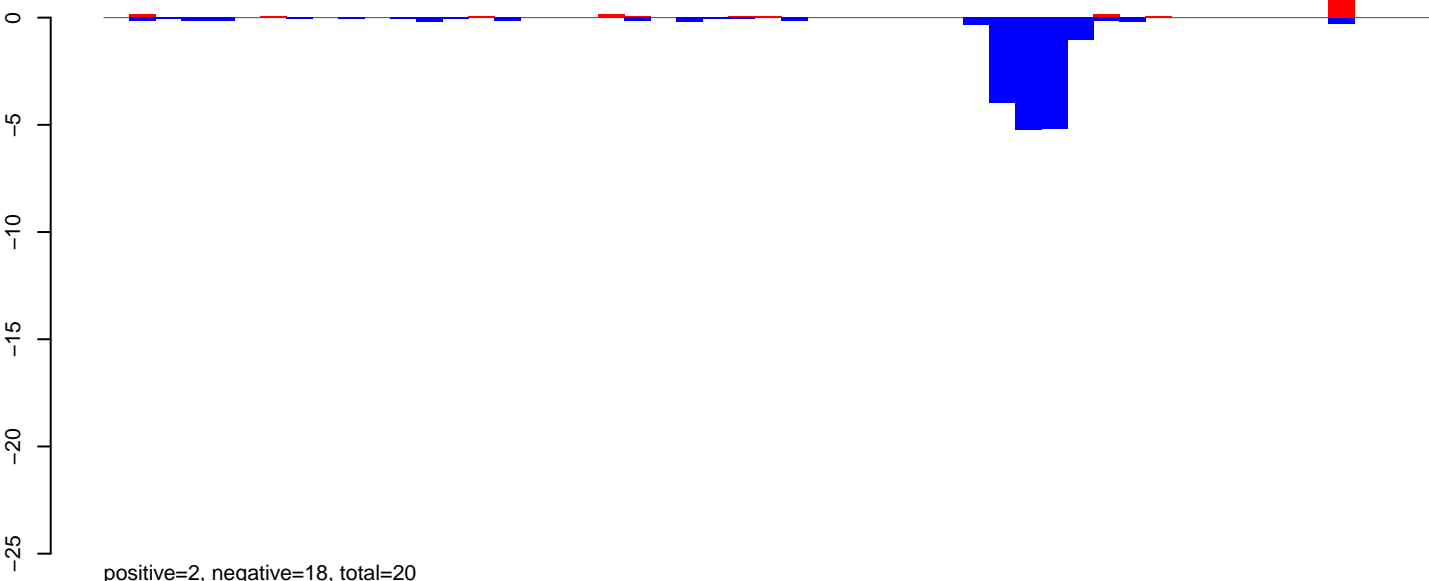


Window size=25, length=1260, TE@rnd-4_family-308@Unknown@3534:1-1260

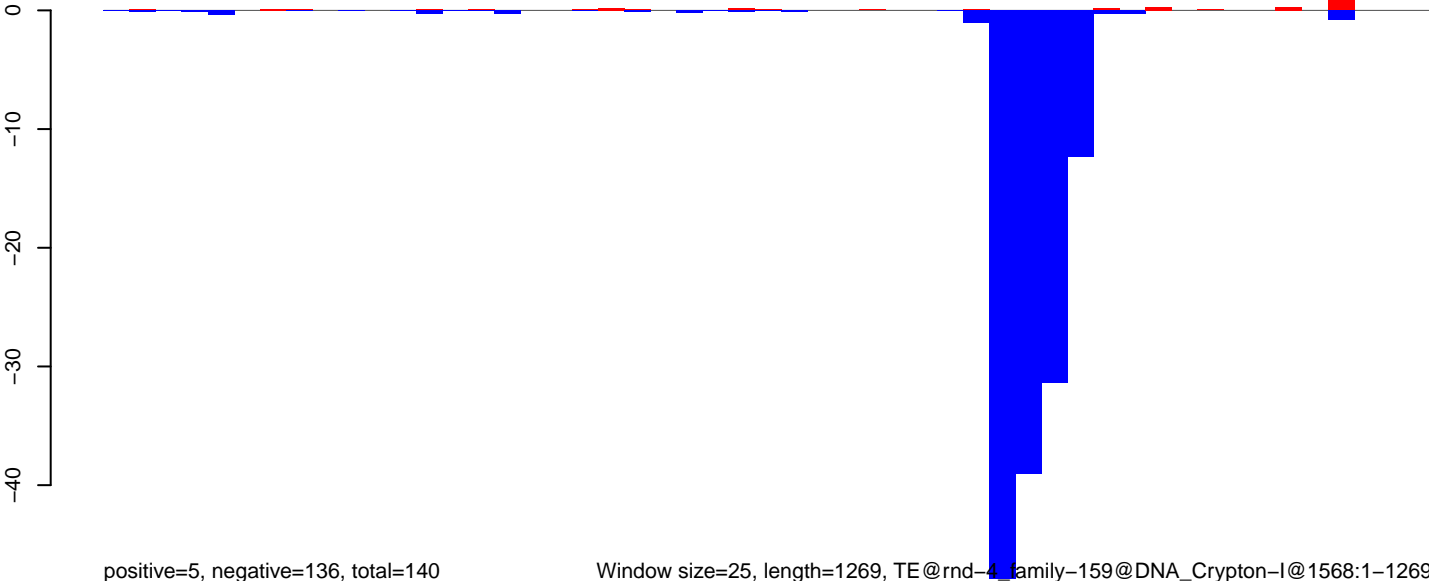
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



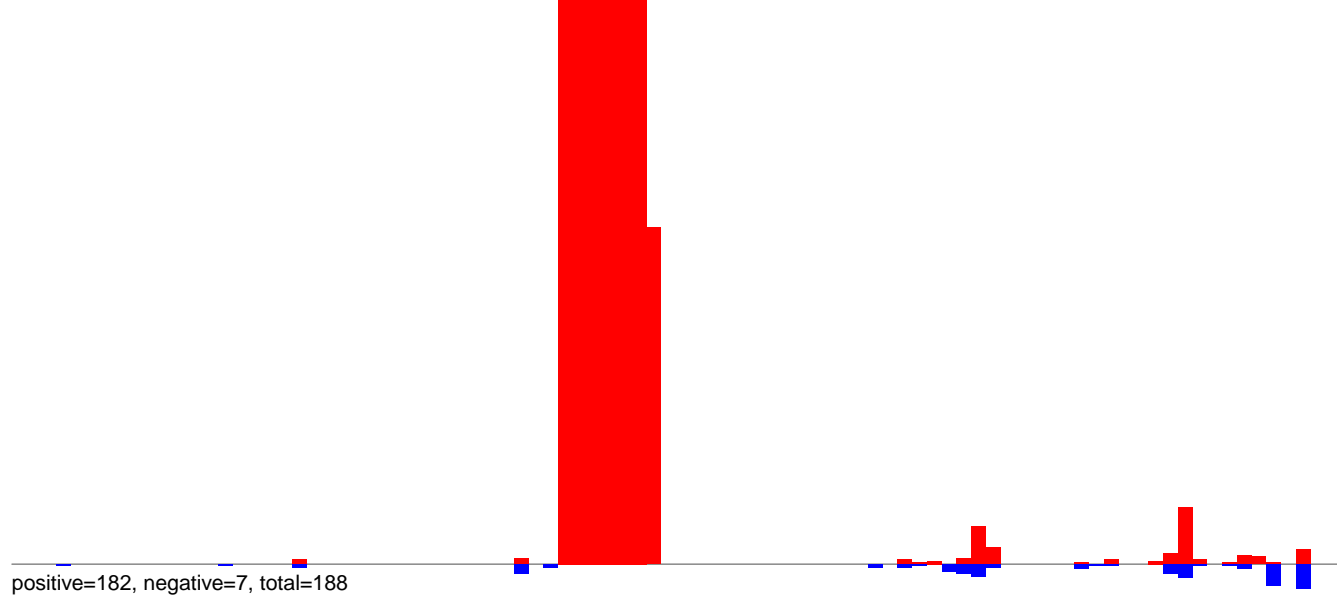
CuQuin_Mcarc_NL.rep



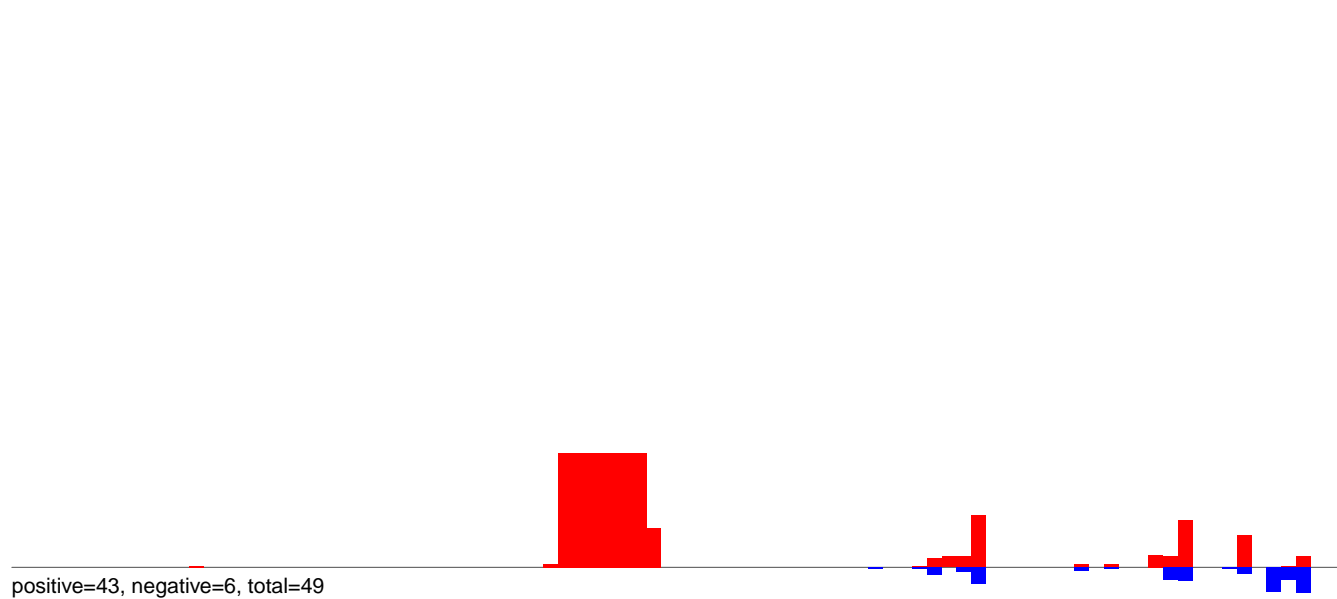
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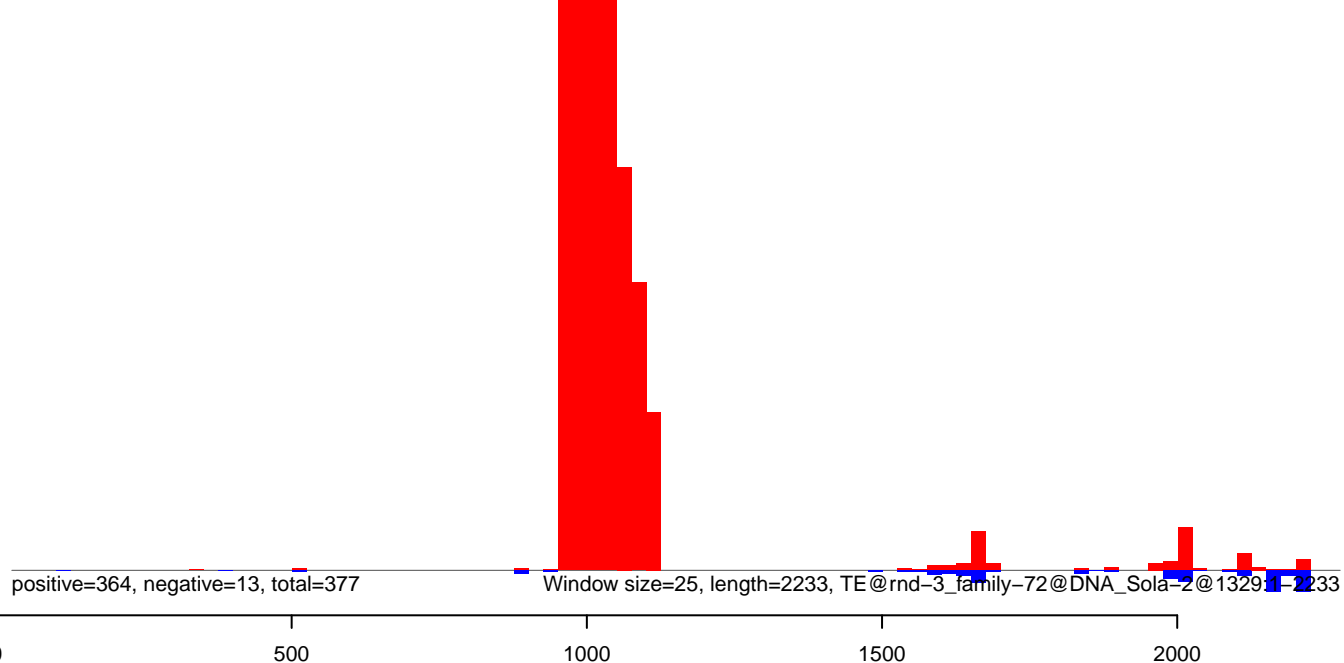
CuQuin_Mcarc_NL.18_23.rep



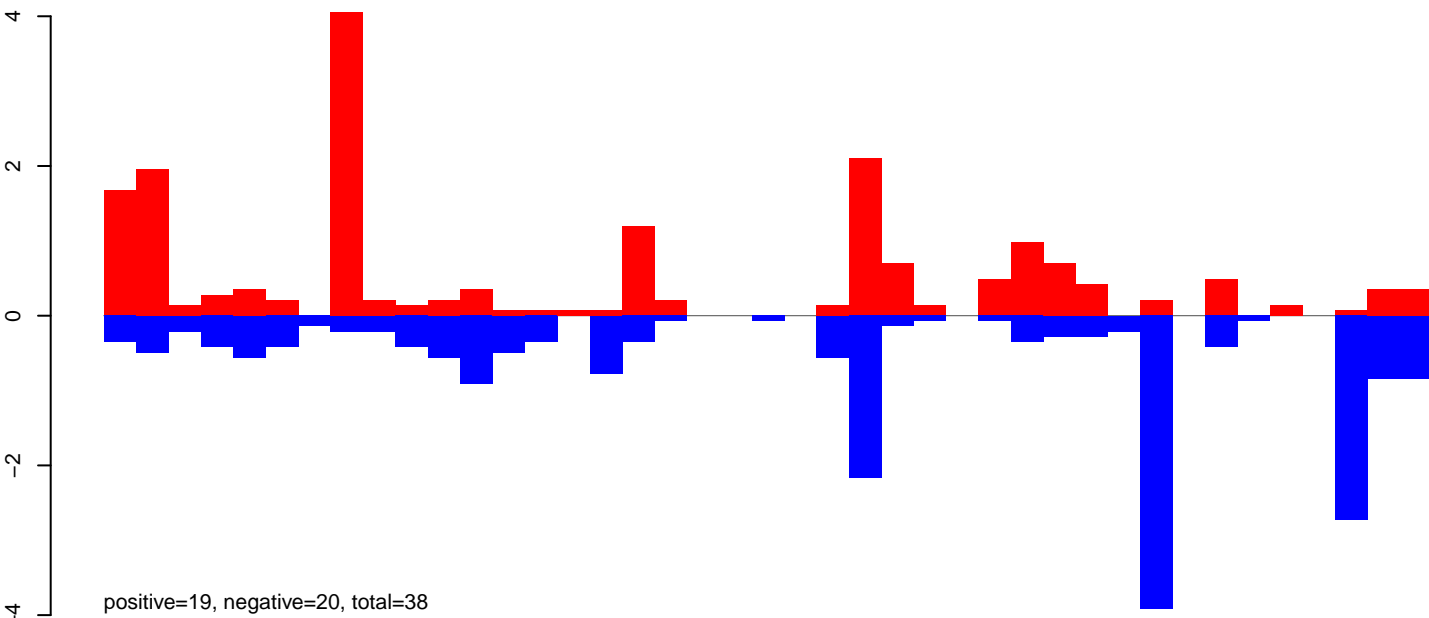
CuQuin_Mcarc_NL.24_35.rep



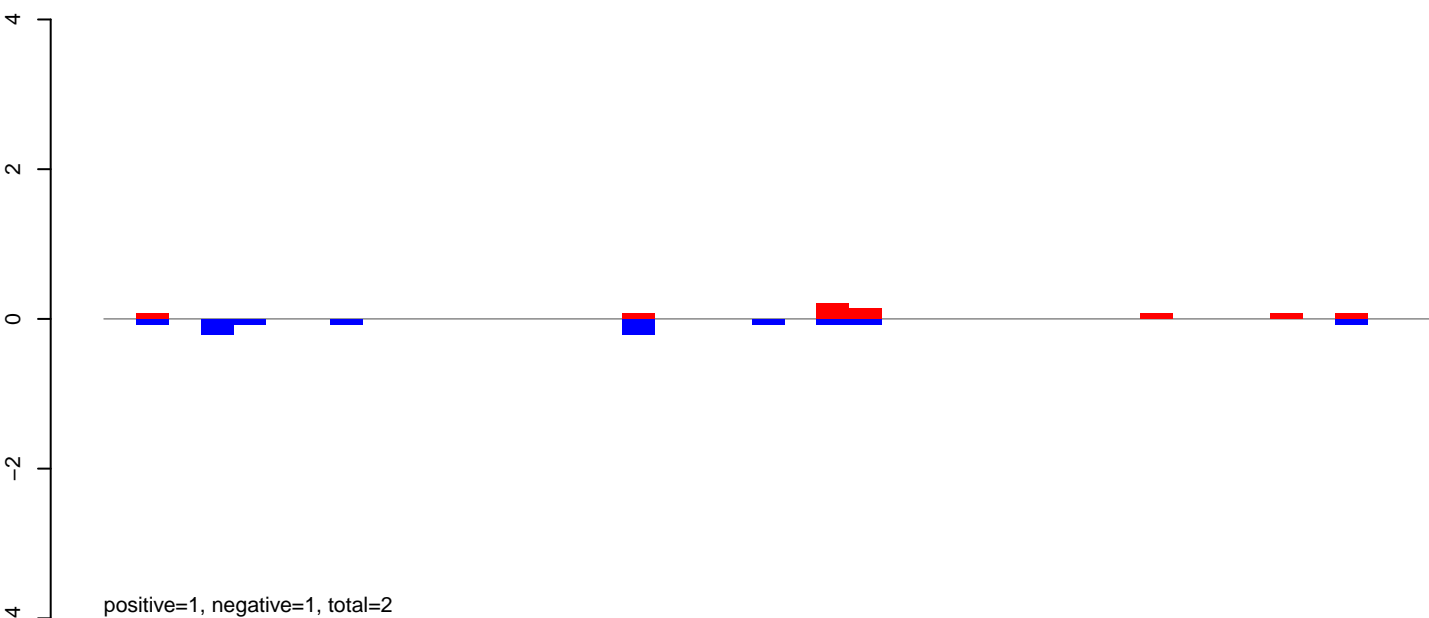
CuQuin_Mcarc_NL.rep



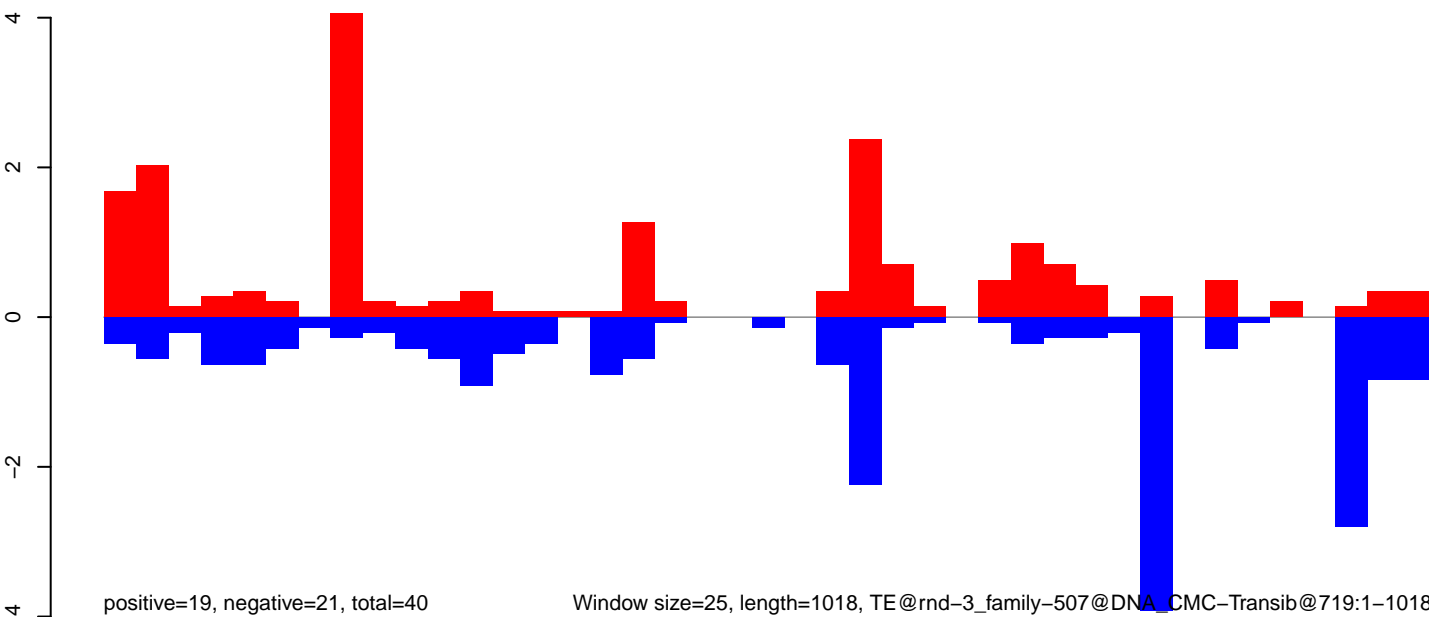
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

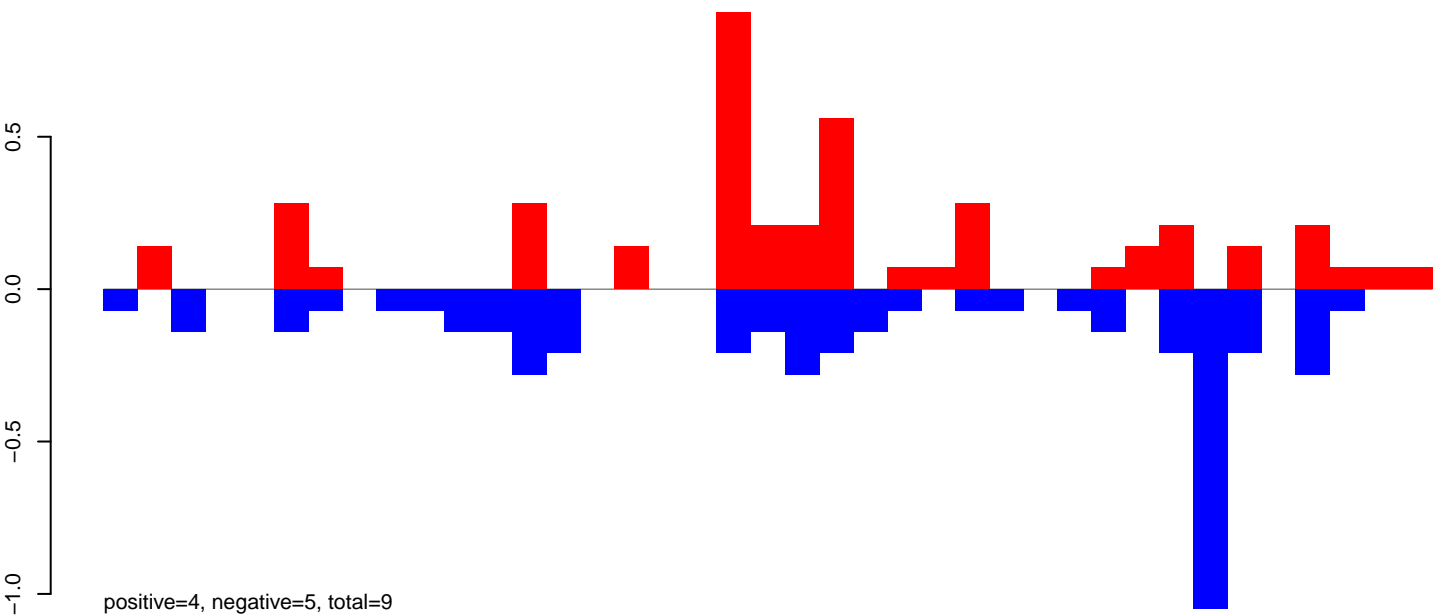


CuQuin_Mcarc_NL.rep

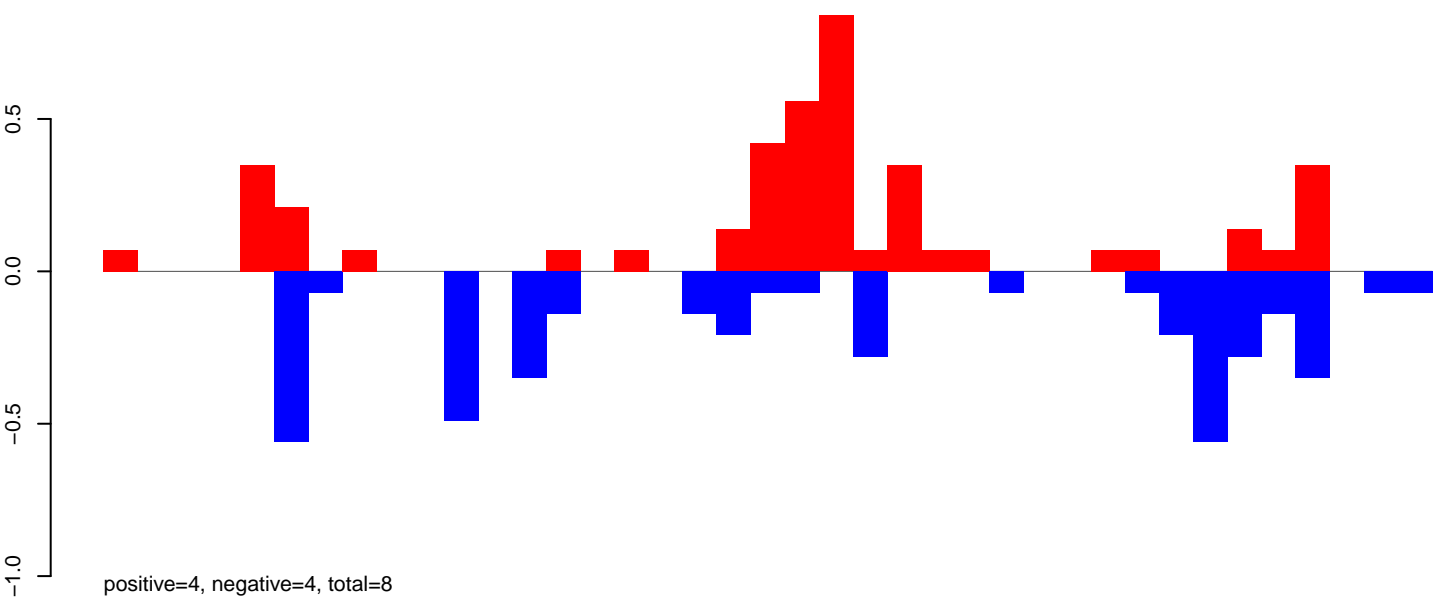


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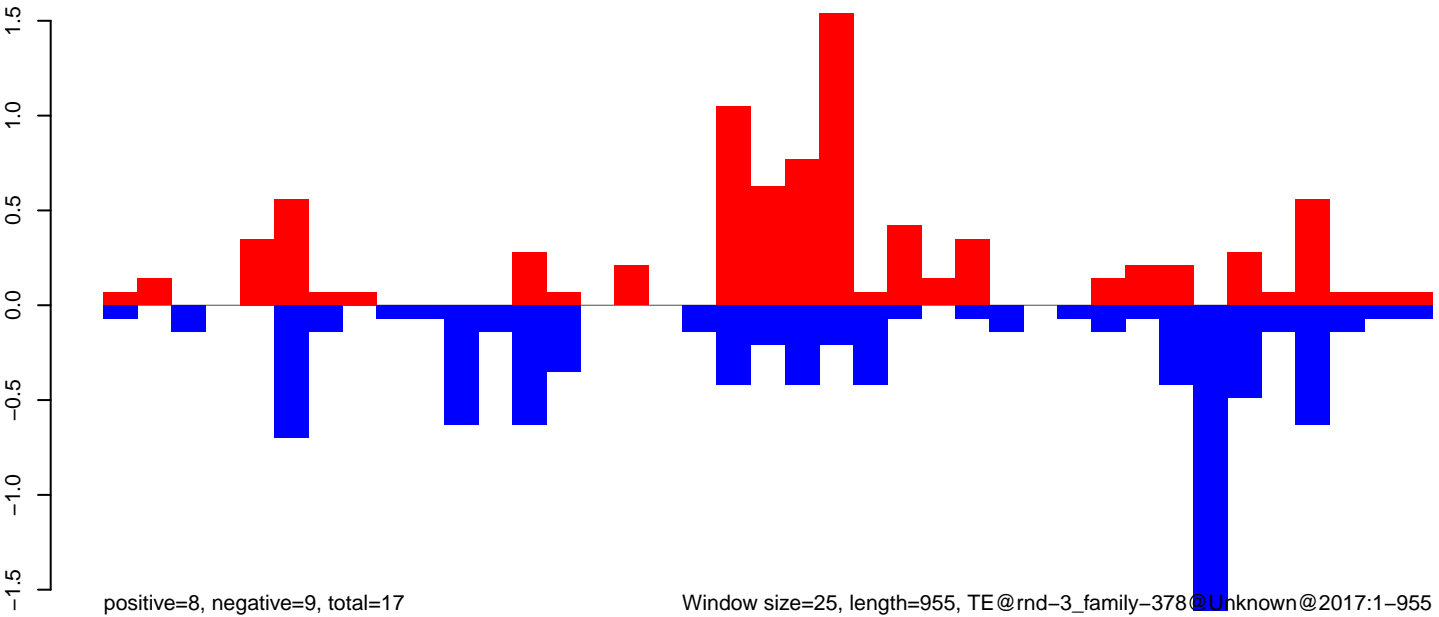
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



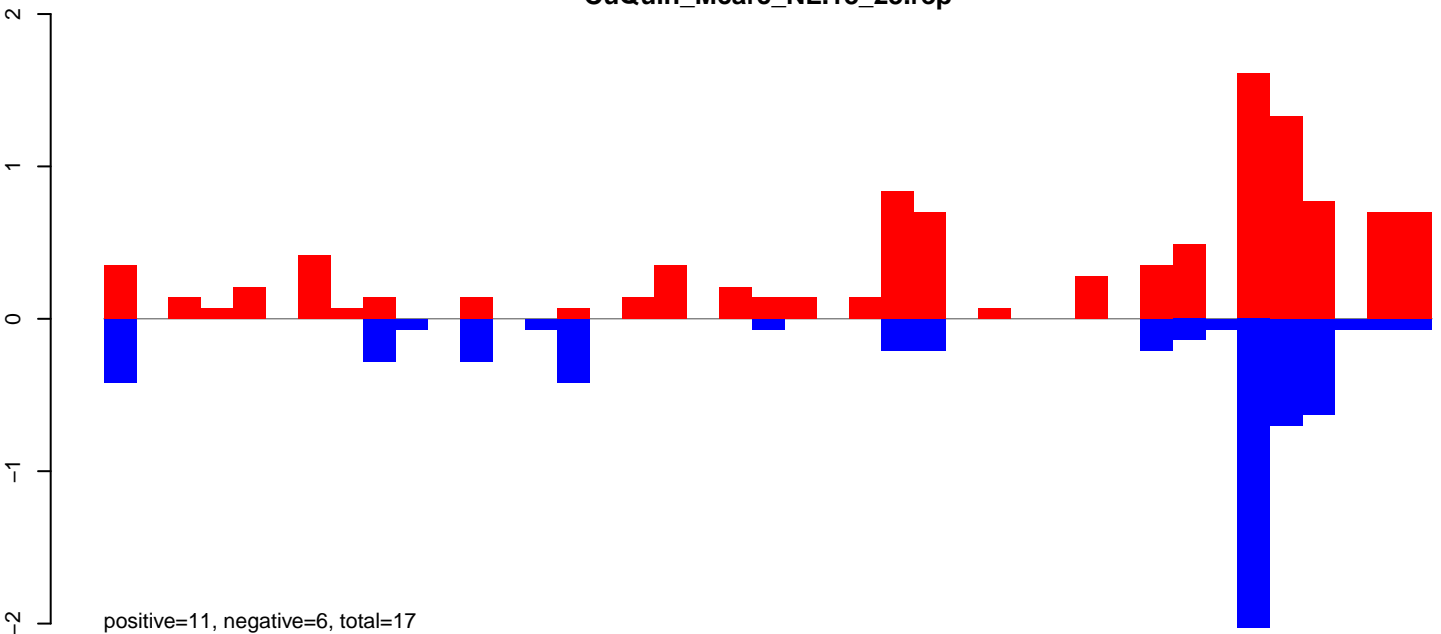
CuQuin_Mcarc_NL.rep



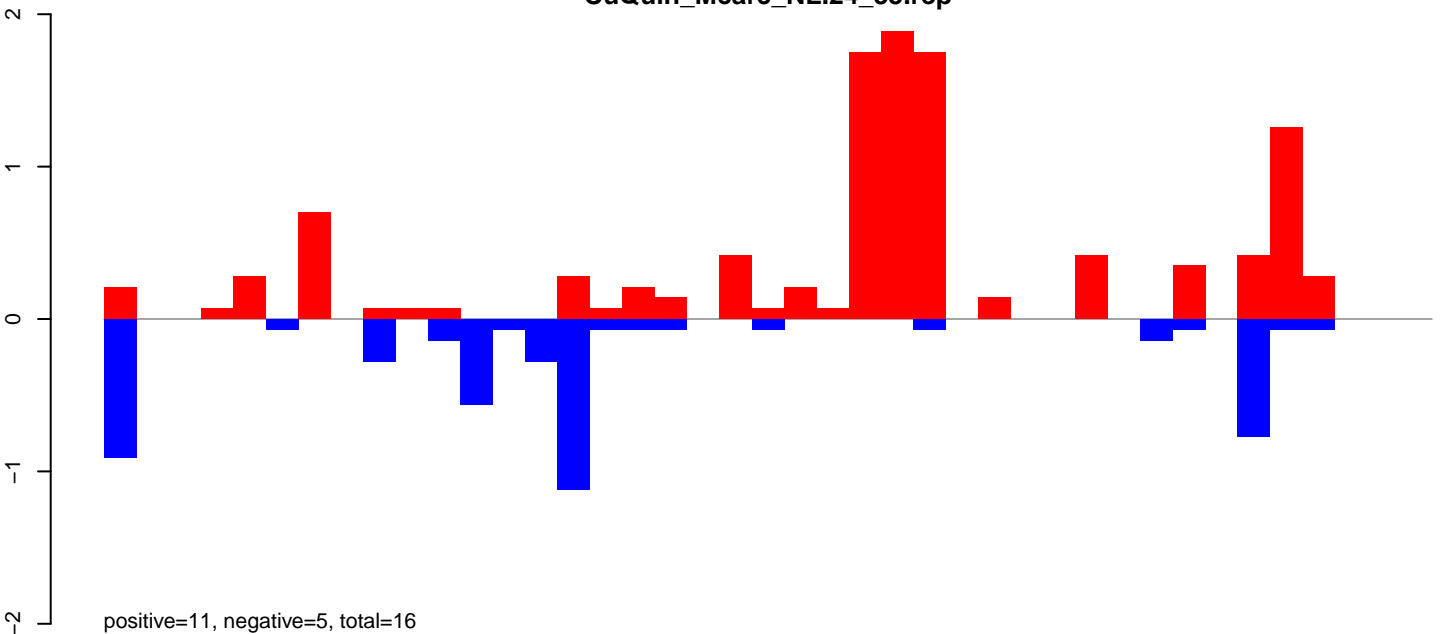
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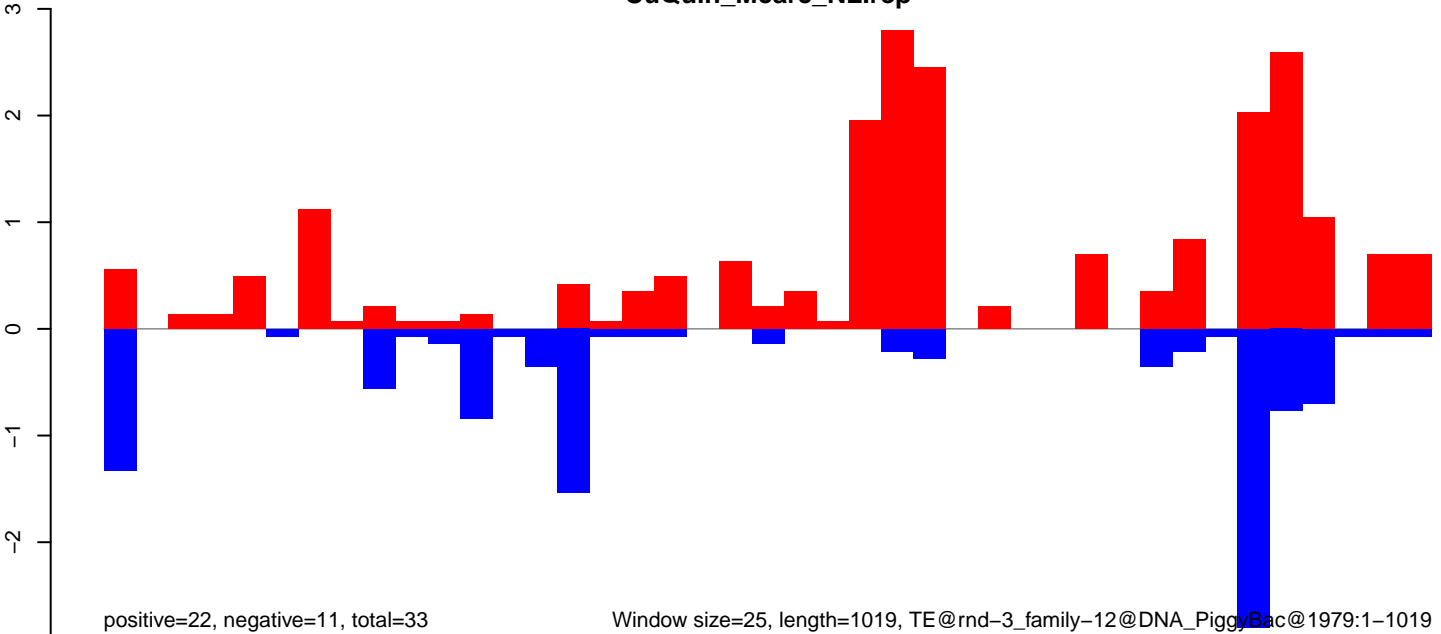
CuQuin_Mcarc_NL.18_23.rep



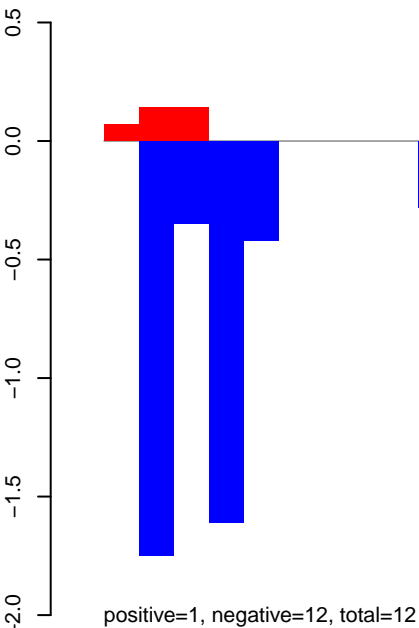
CuQuin_Mcarc_NL.24_35.rep



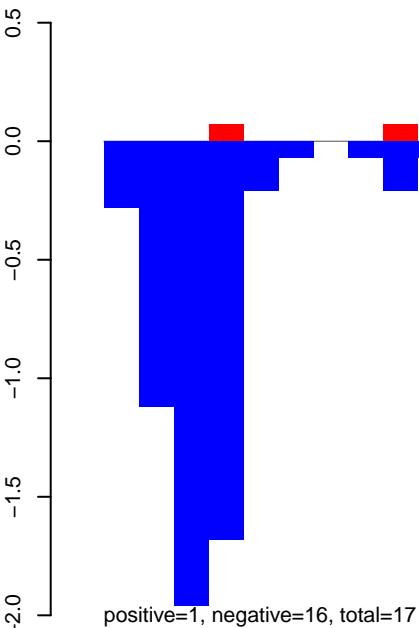
CuQuin_Mcarc_NL.rep



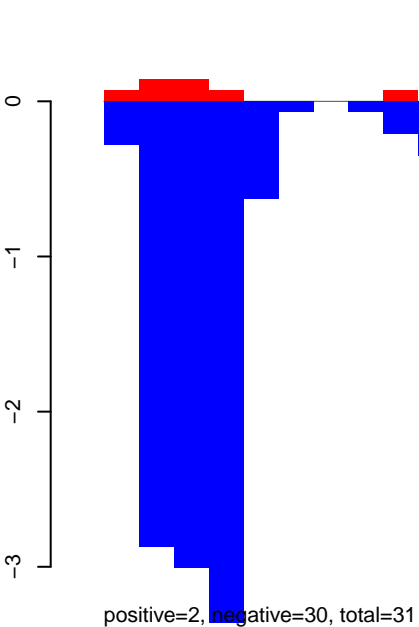
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

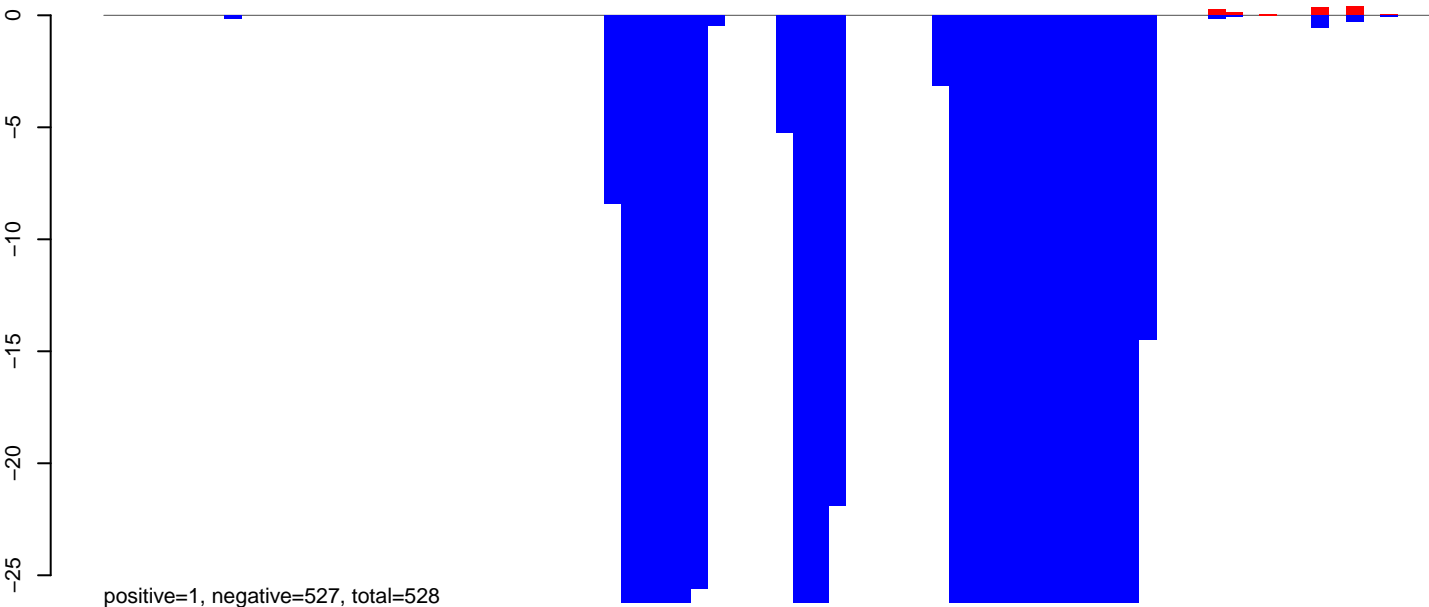


CuQuin_Mcarc_NL.rep

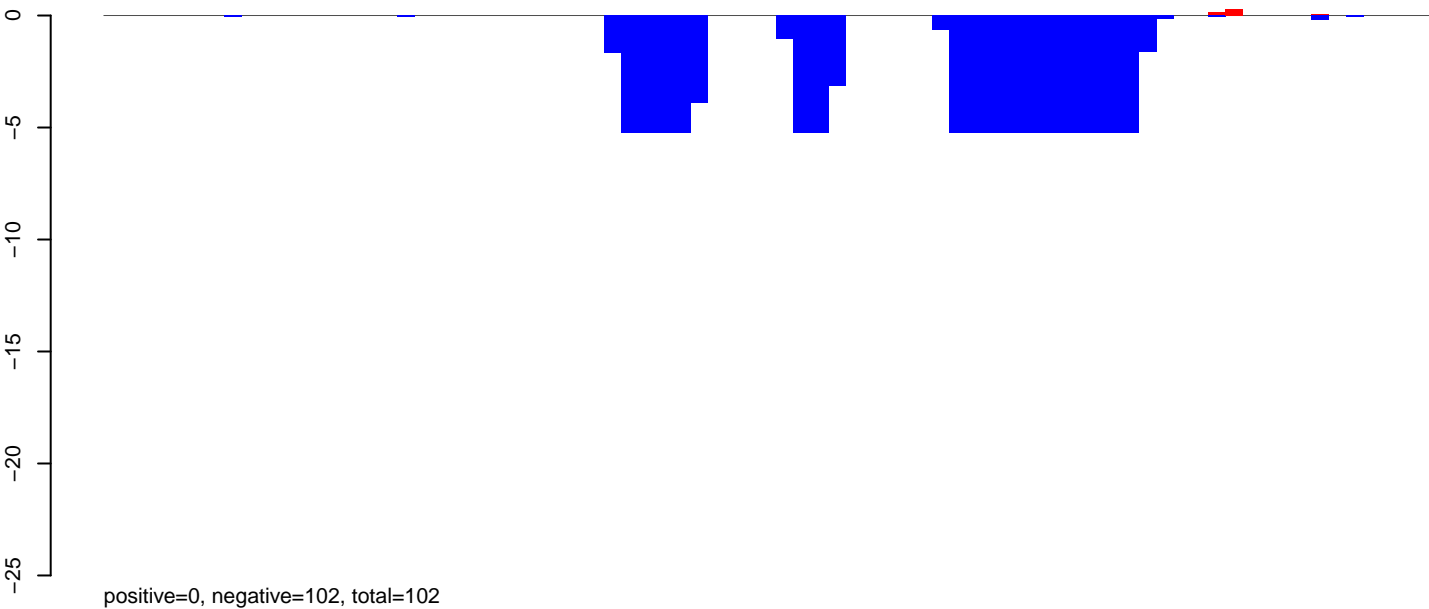


Window size=25, length=932, TE@rnd-5_family-943@Unknown@1168:1-932

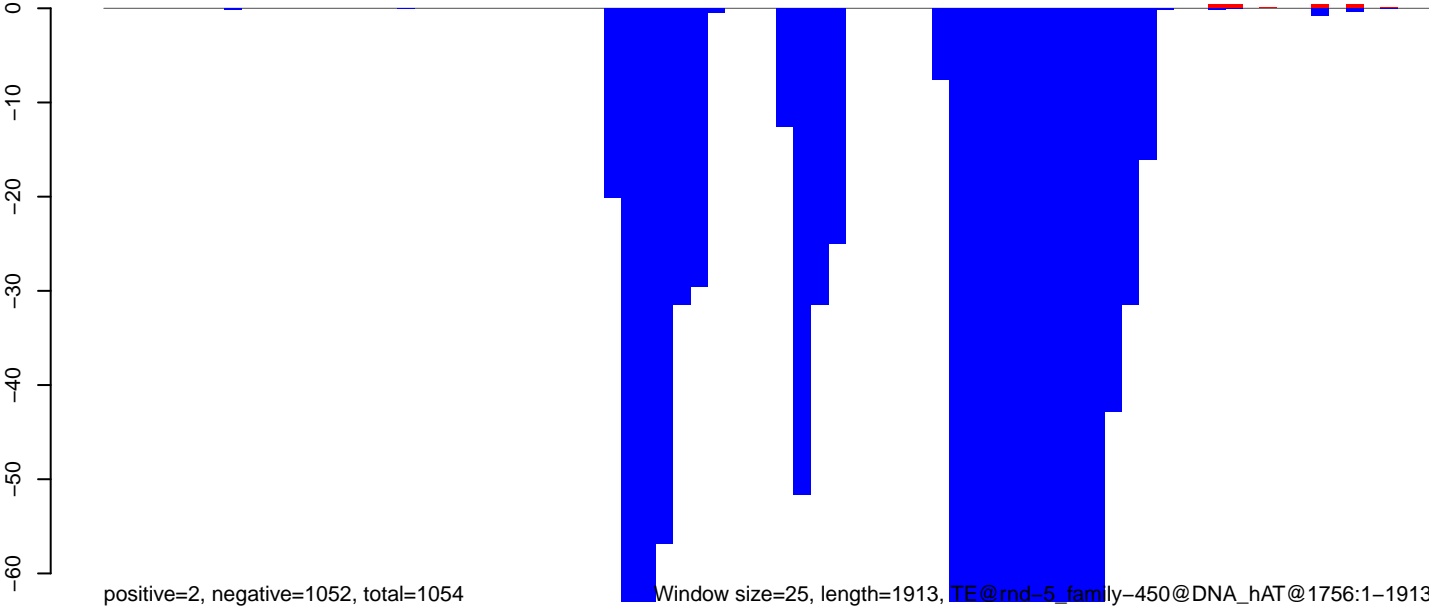
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

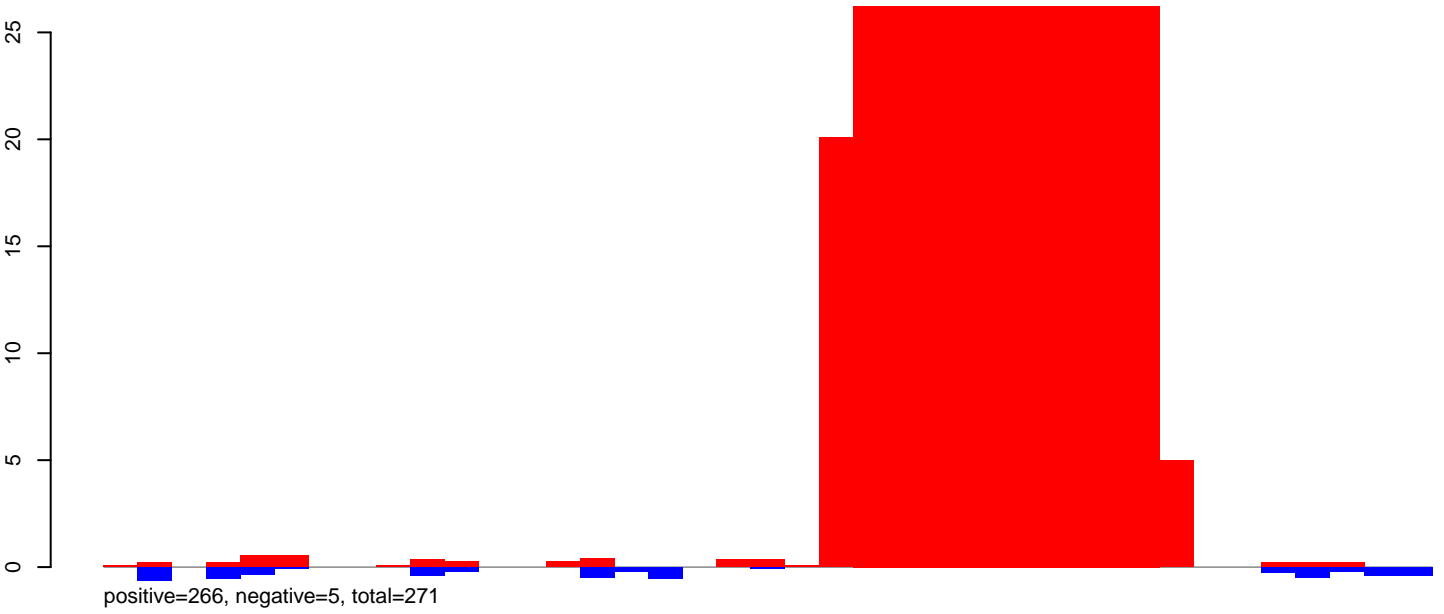


CuQuin_Mcarc_NL.rep

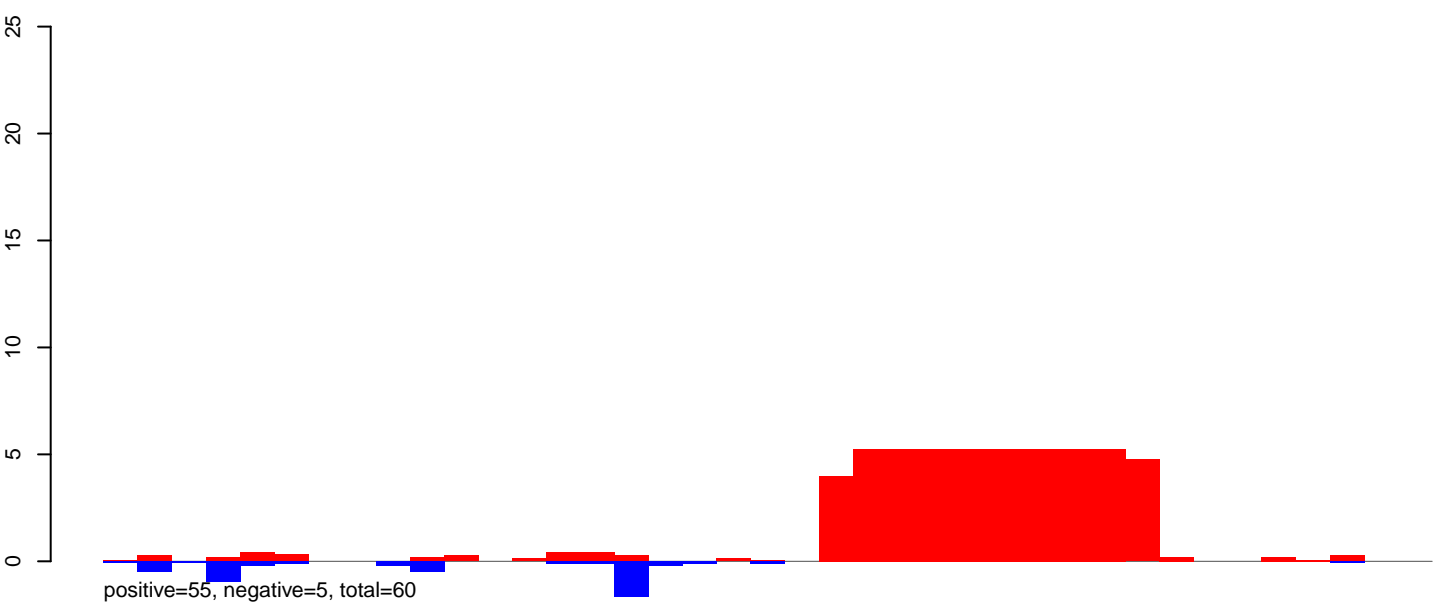


Window size=25, length=1913, TE@rnd-5_family-450@DNA_hAT@1756:1-1913

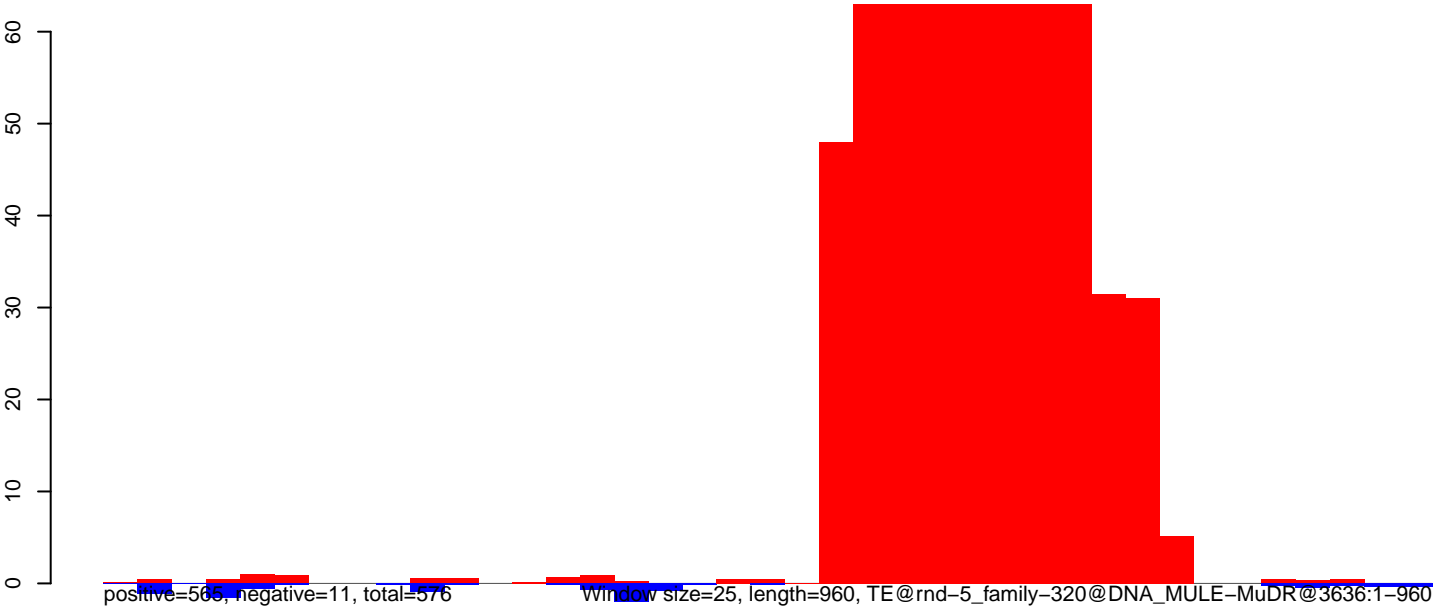
CuQuin_Mcarc_NL.18_23.rep



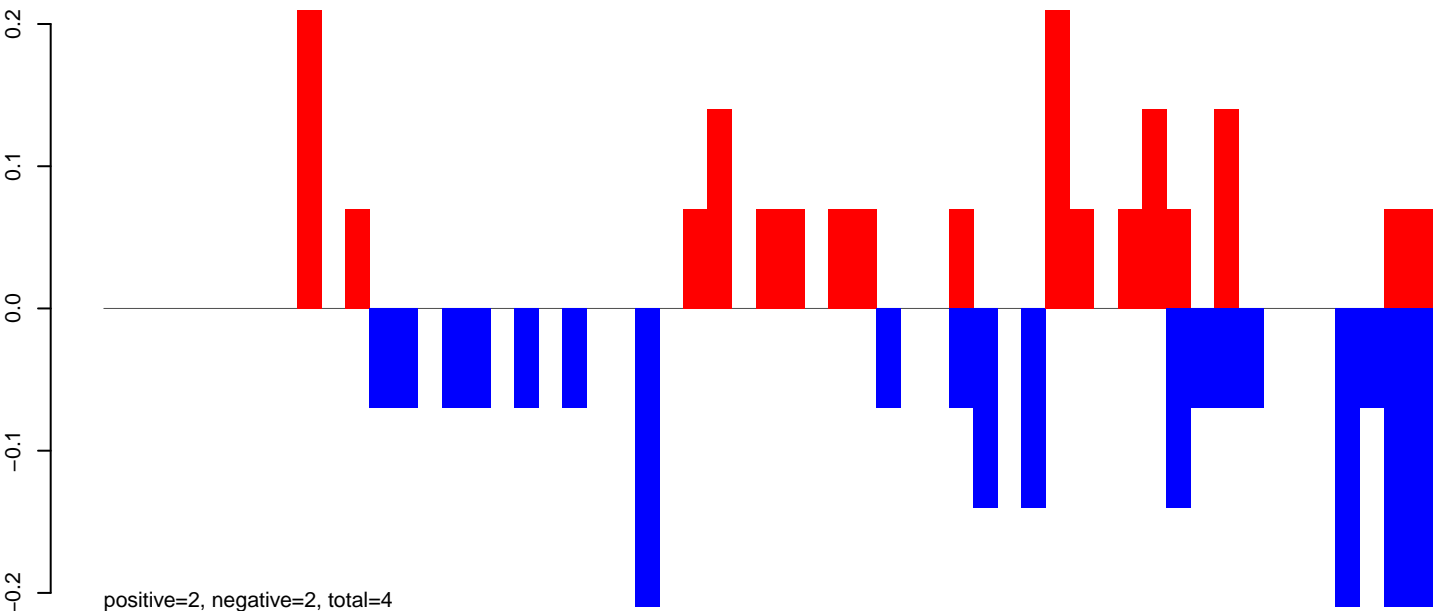
CuQuin_Mcarc_NL.24_35.rep



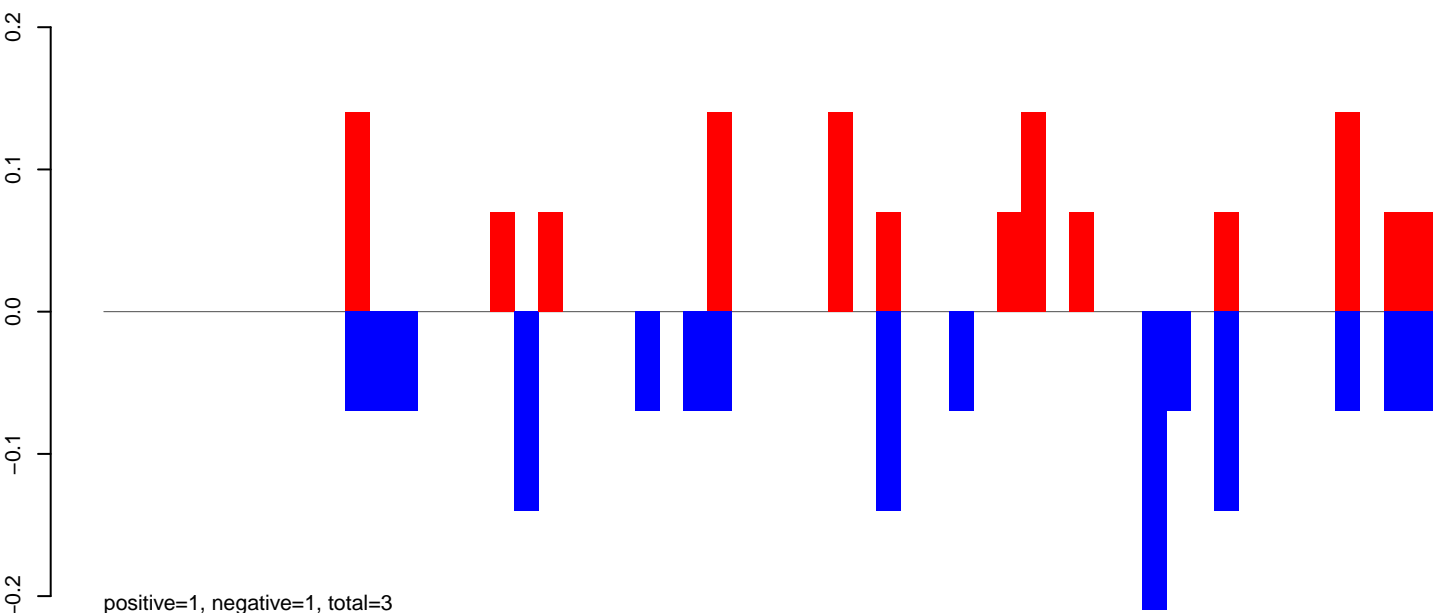
CuQuin_Mcarc_NL.rep



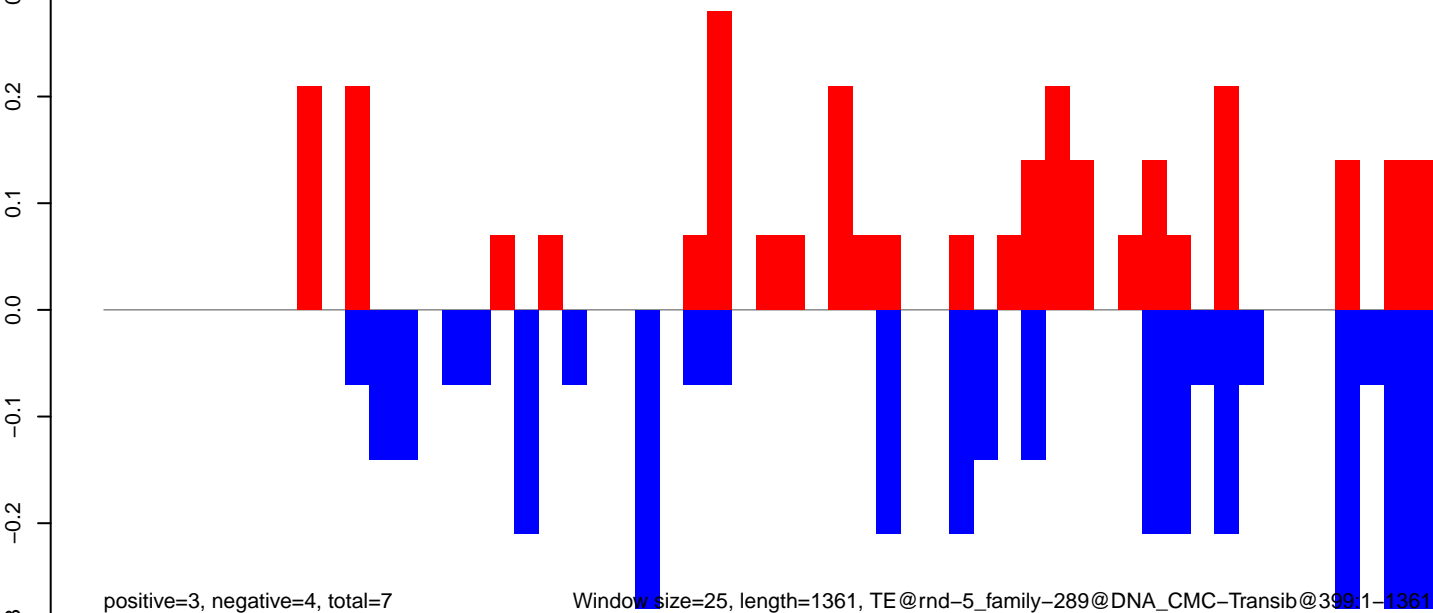
CuQuin_Mcarc_NL.18_23.rep



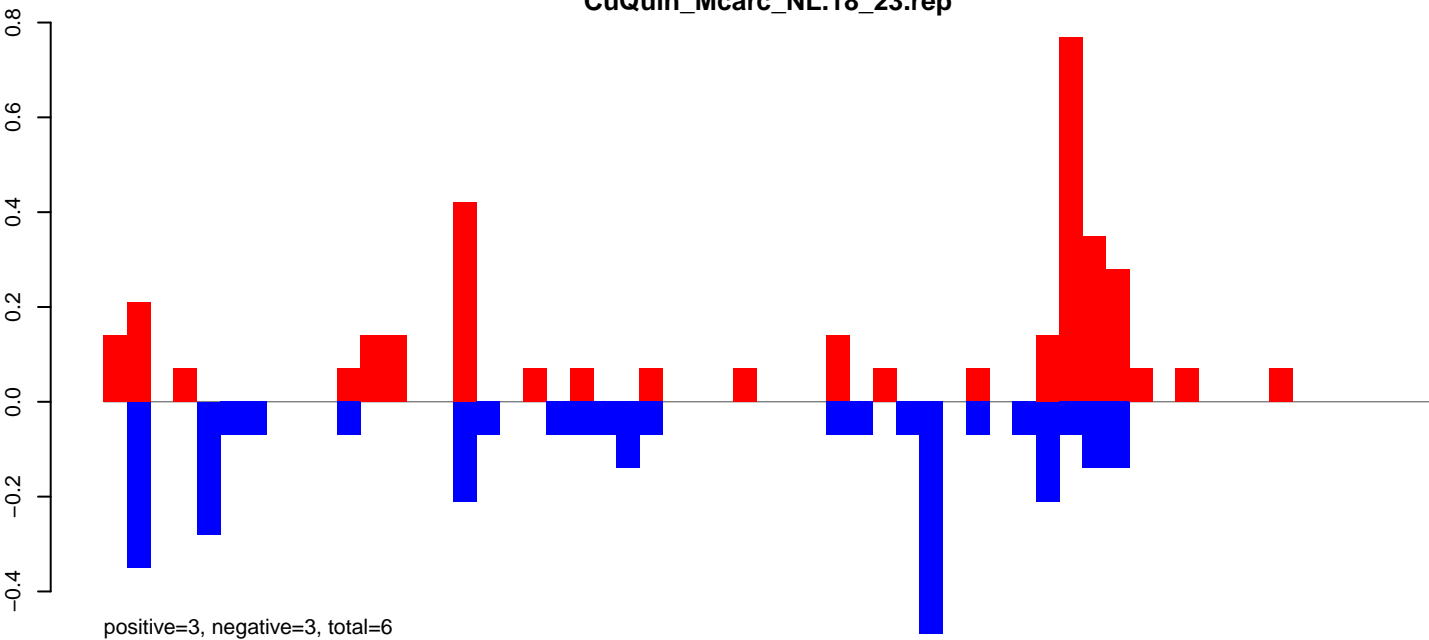
CuQuin_Mcarc_NL.24_35.rep



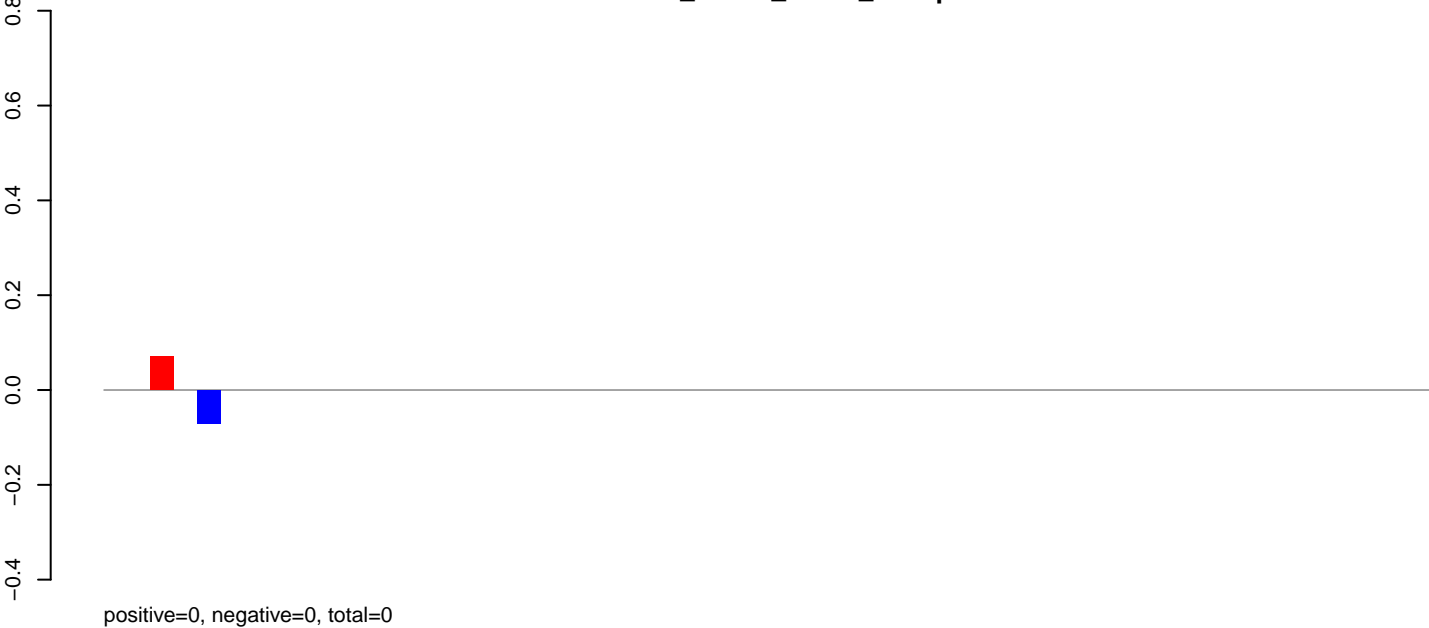
CuQuin_Mcarc_NL.rep



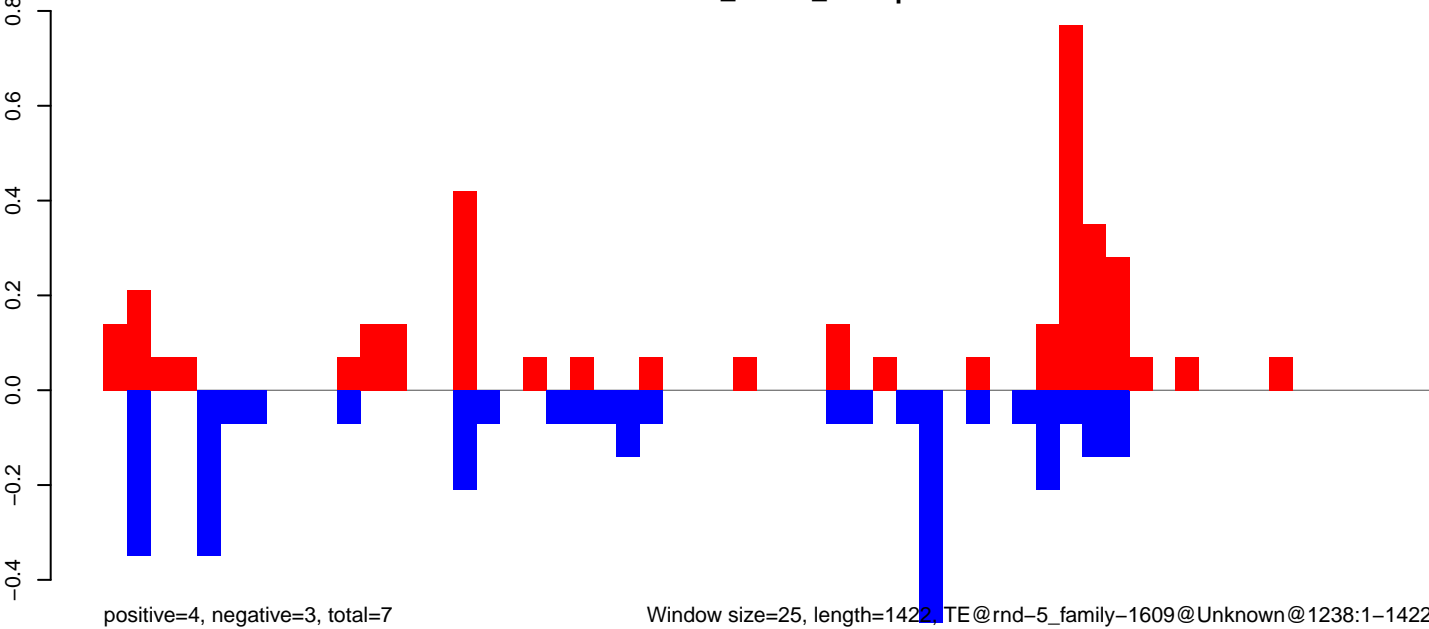
CuQuin_Mcarc_NL.18_23.rep



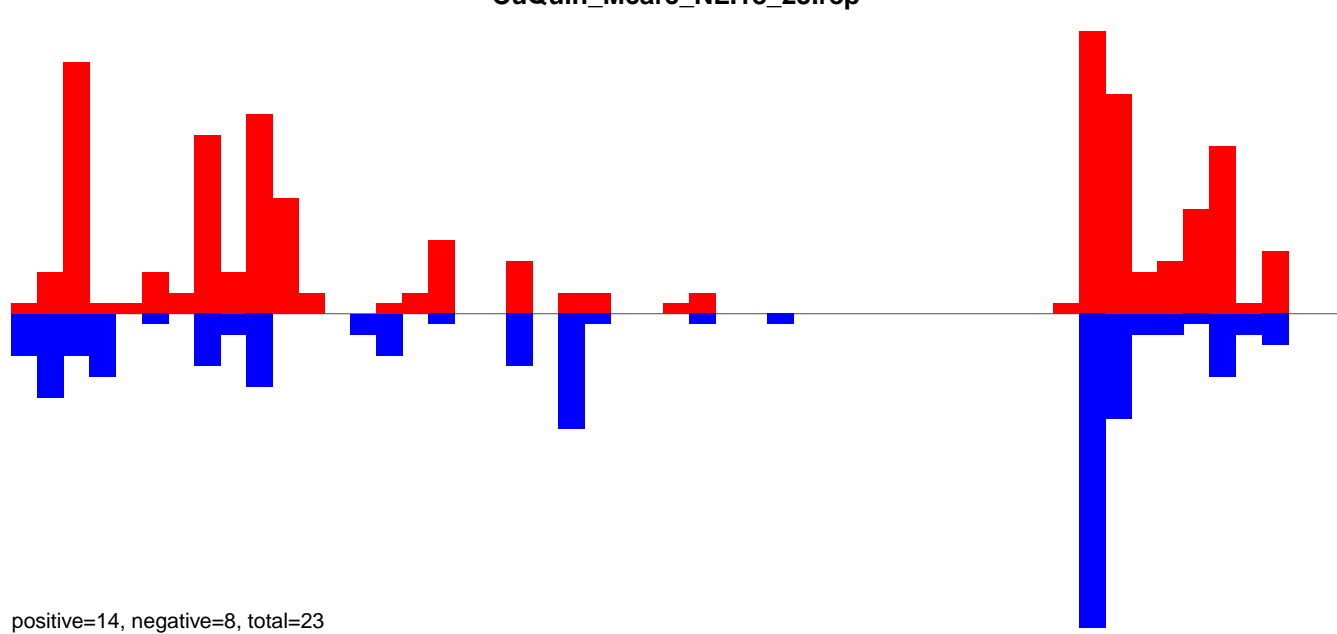
CuQuin_Mcarc_NL.24_35.rep



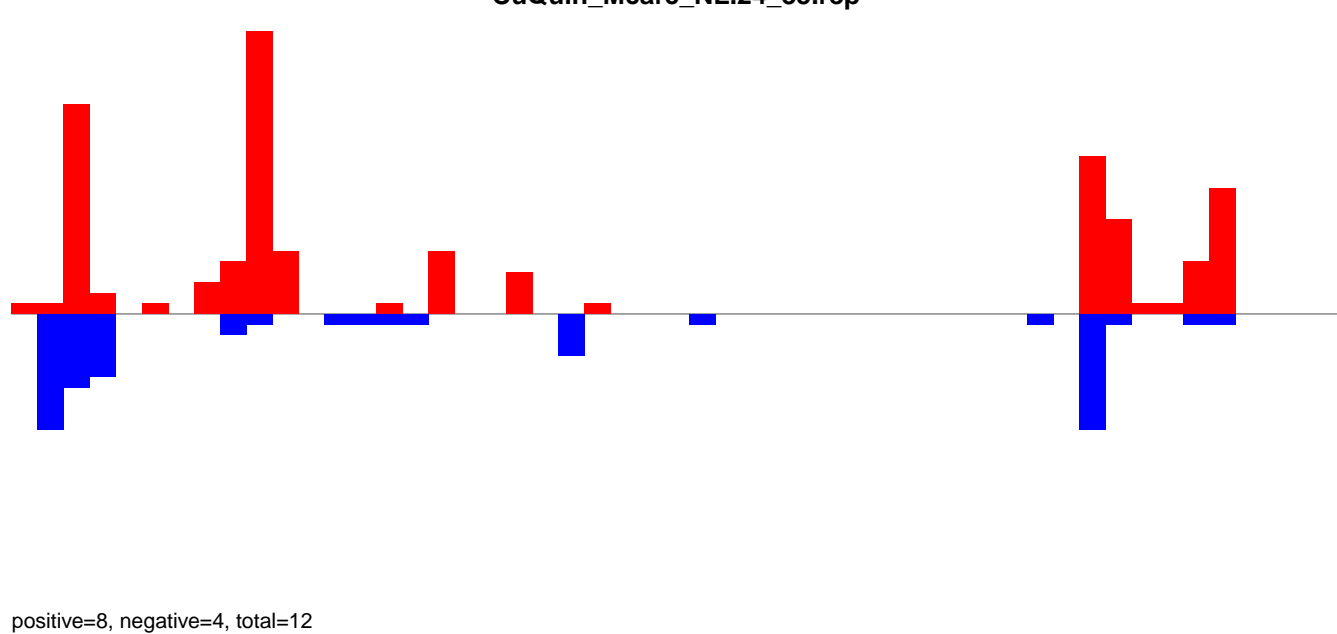
CuQuin_Mcarc_NL.rep



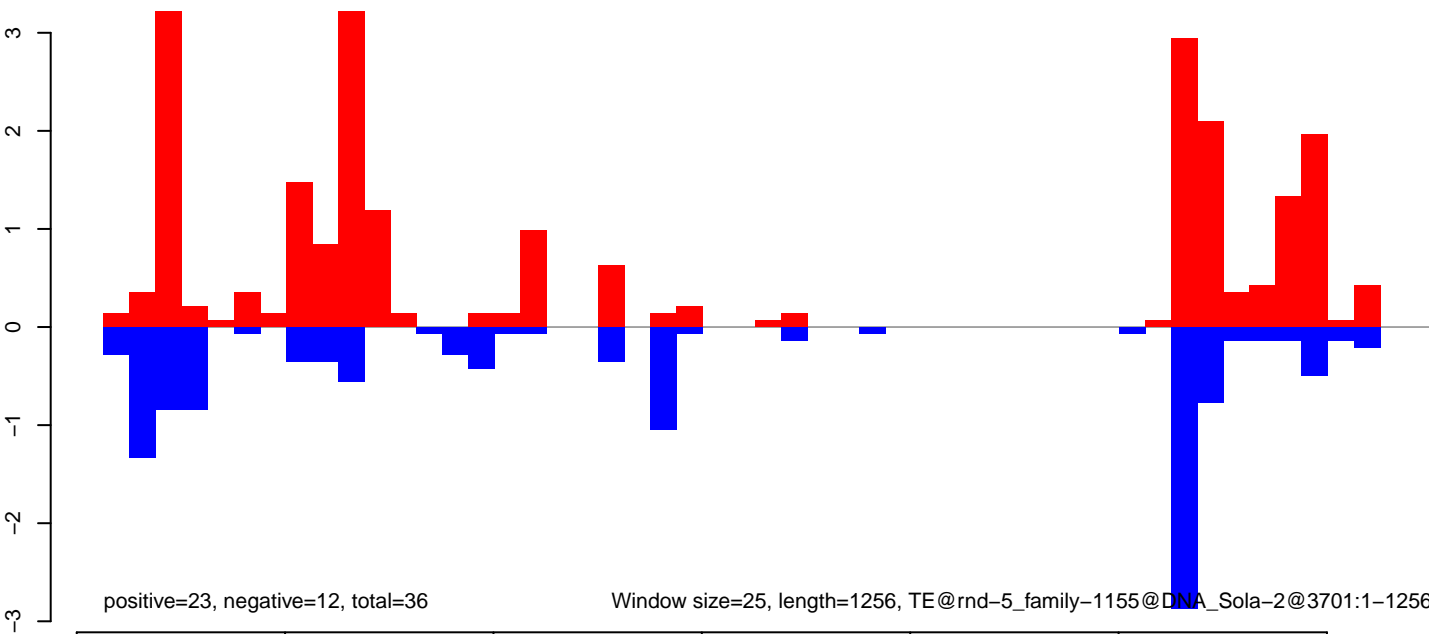
CuQuin_Mcarc_NL.18_23.rep



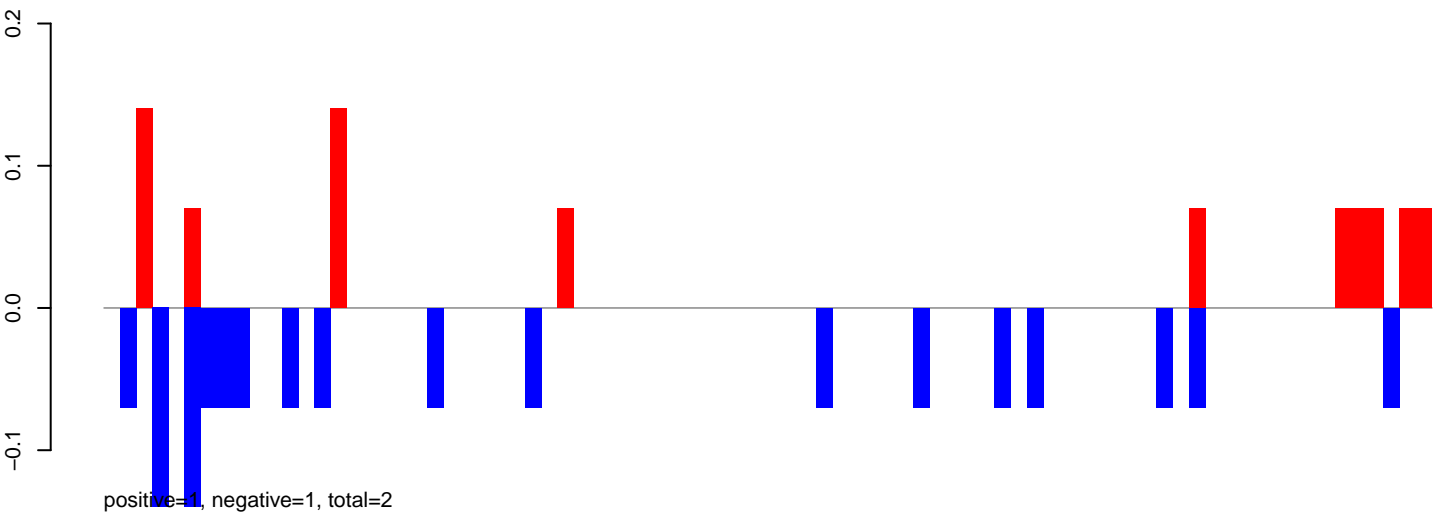
CuQuin_Mcarc_NL.24_35.rep



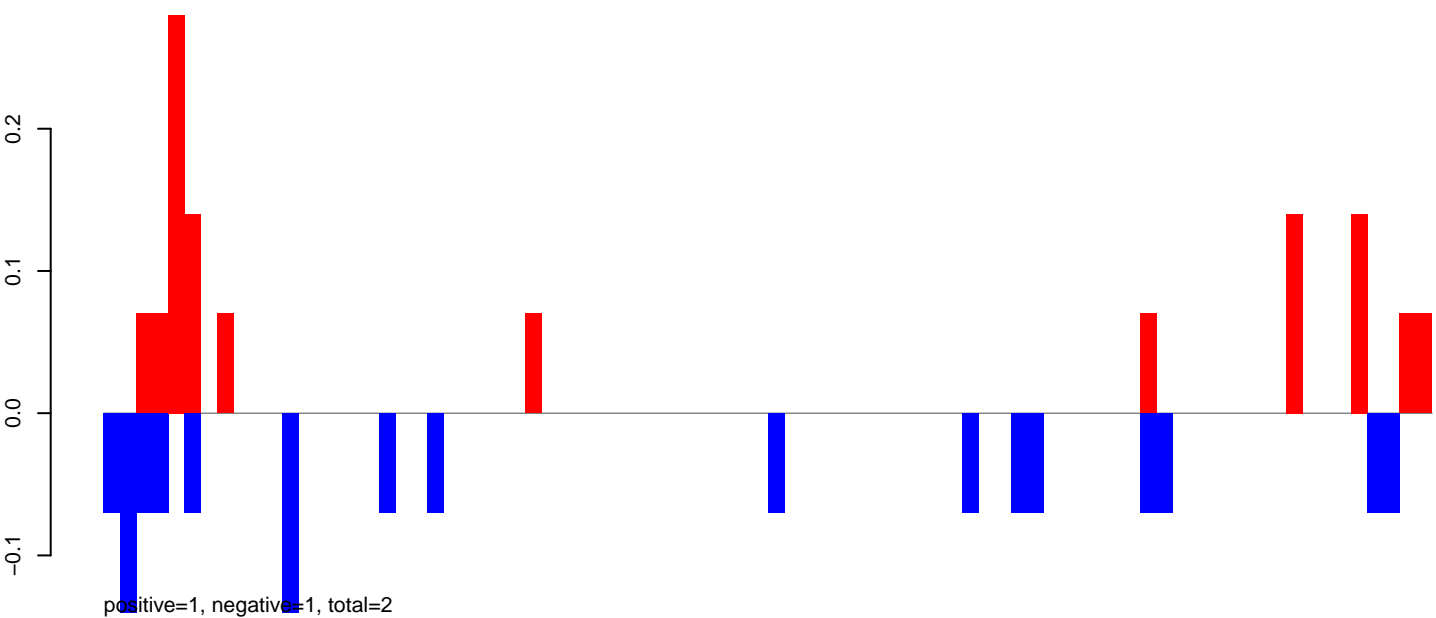
CuQuin_Mcarc_NL.rep



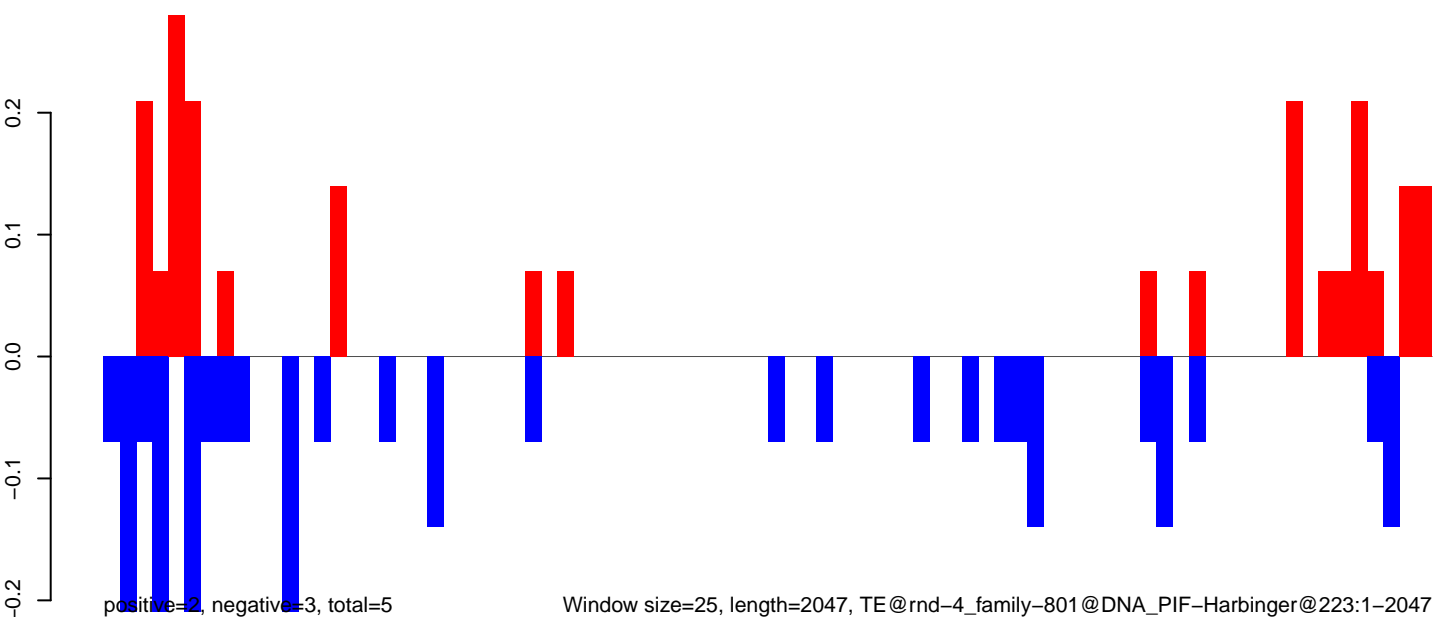
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

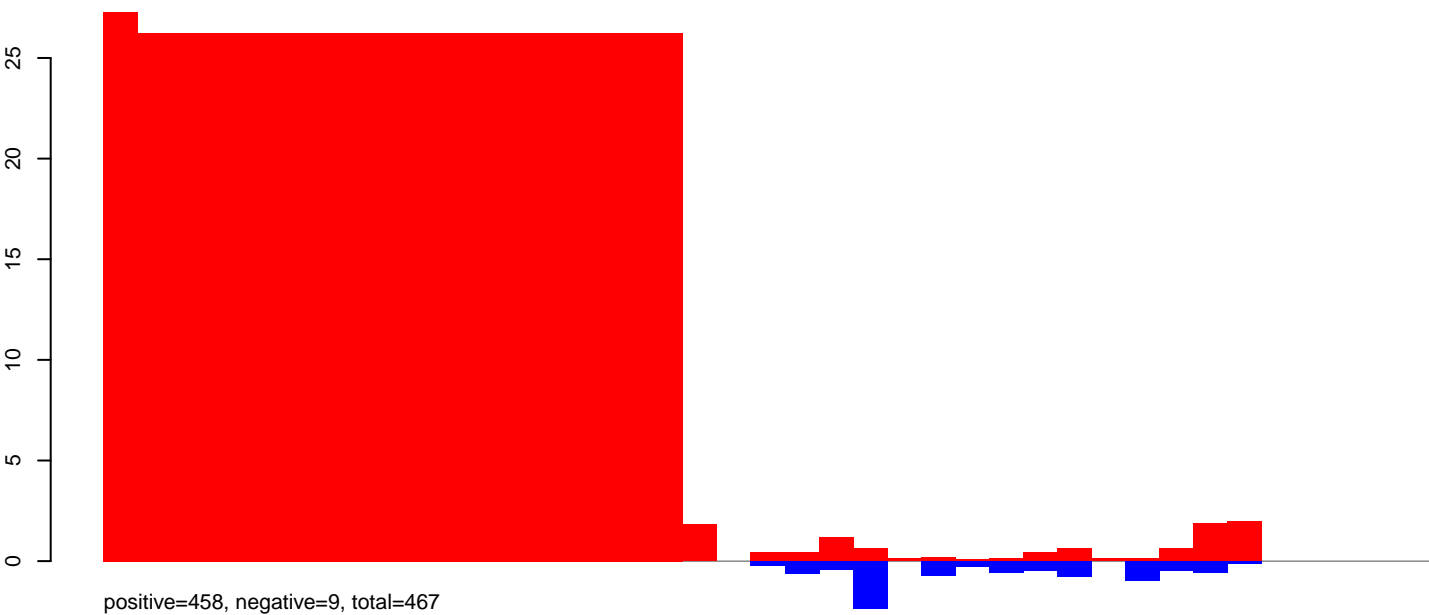


CuQuin_Mcarc_NL.rep

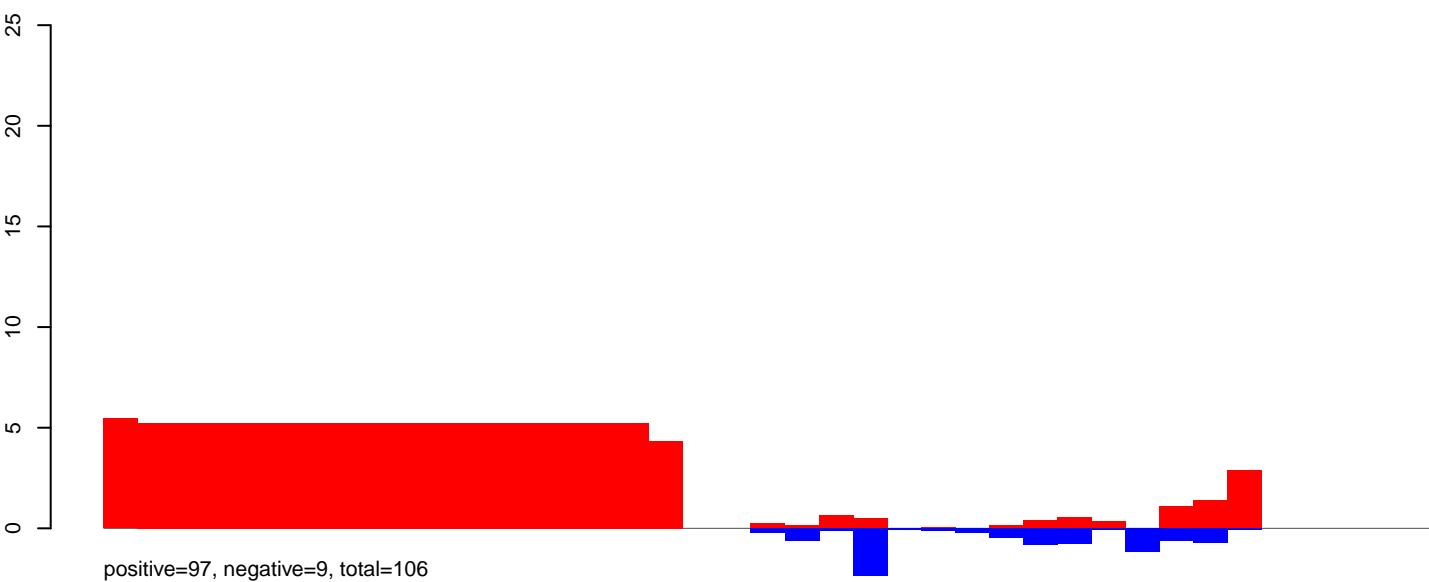


Window size=25, length=2047, TE@rnd-4_family-801@DNA_PIF-Harbinger@223:1-2047

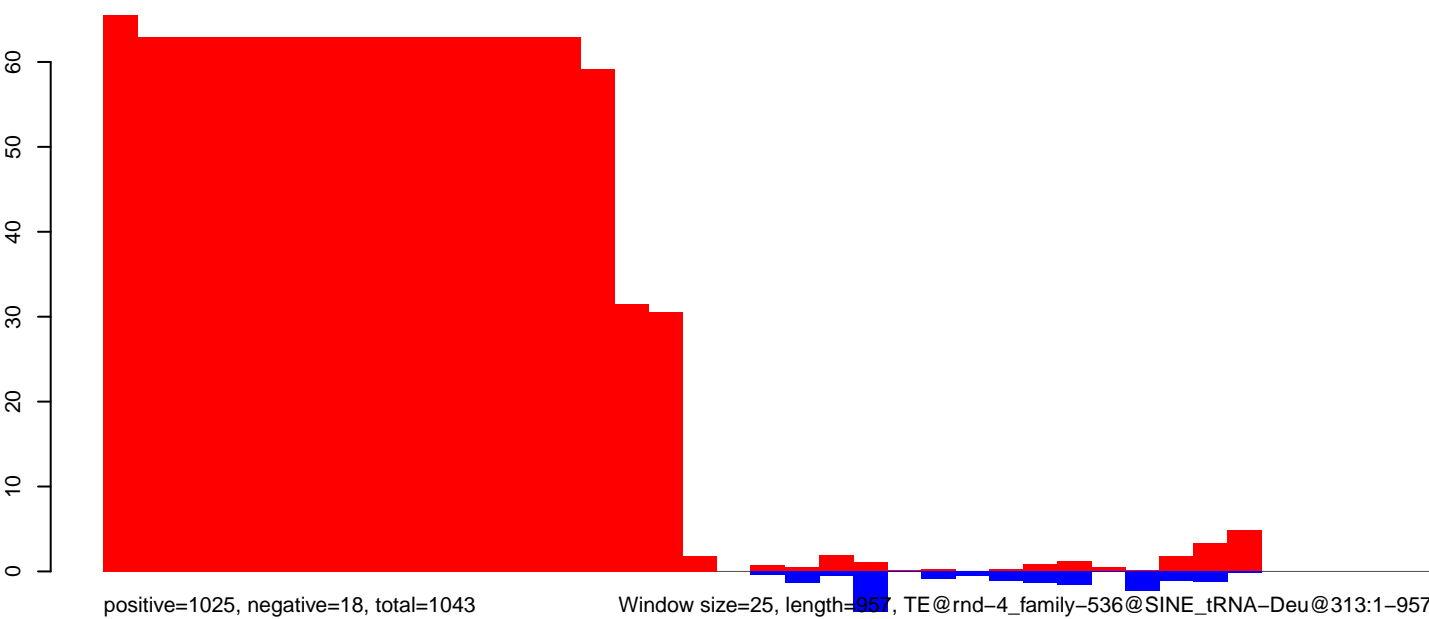
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



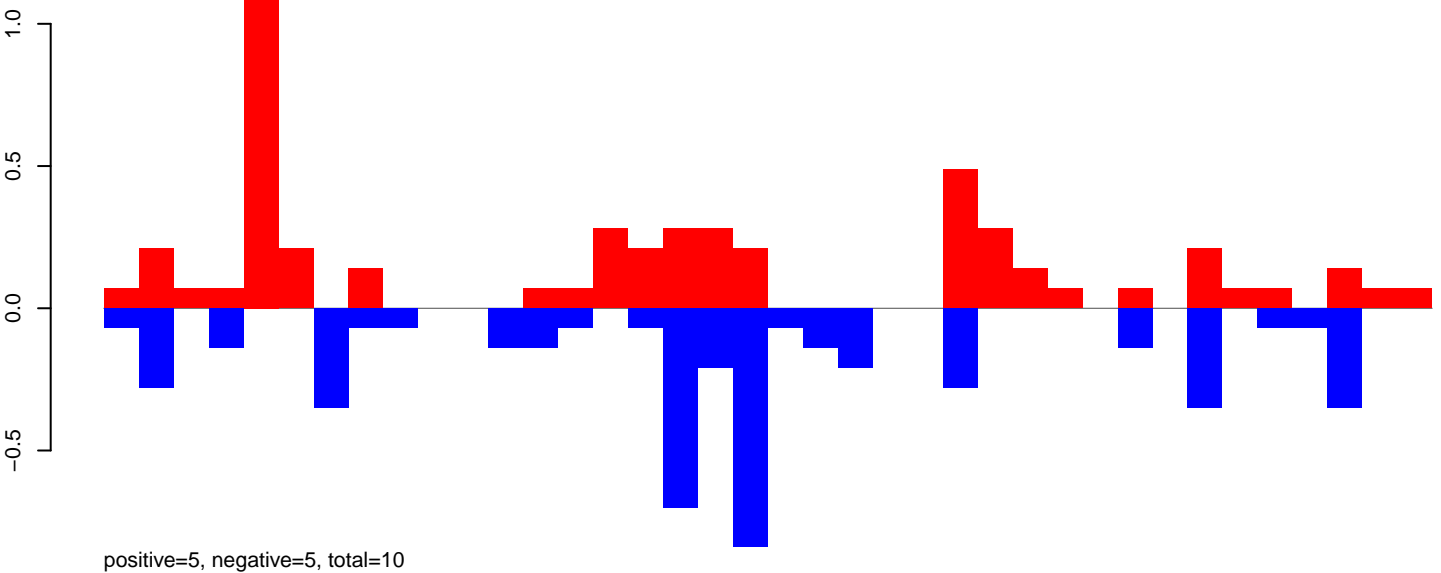
CuQuin_Mcarc_NL.rep



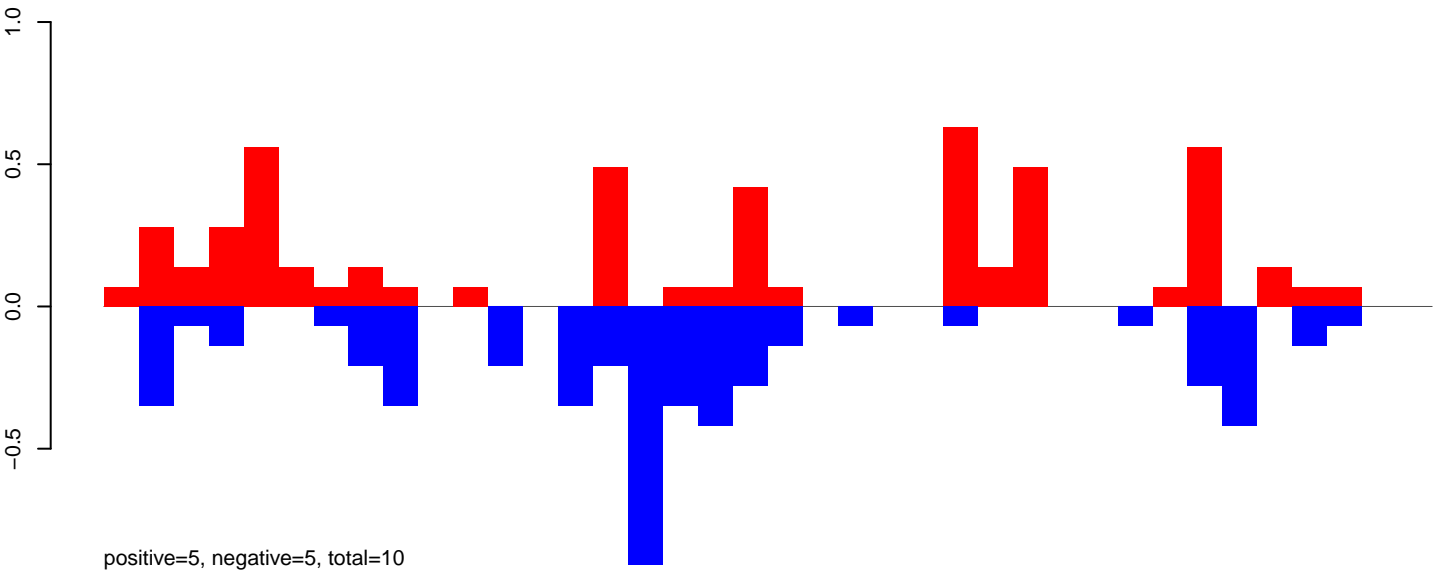
Window size=25, length=957, TE@rnd-4_family-536@SINE_tRNA-Deu@313:1-957

0 200 400 600 800 1000

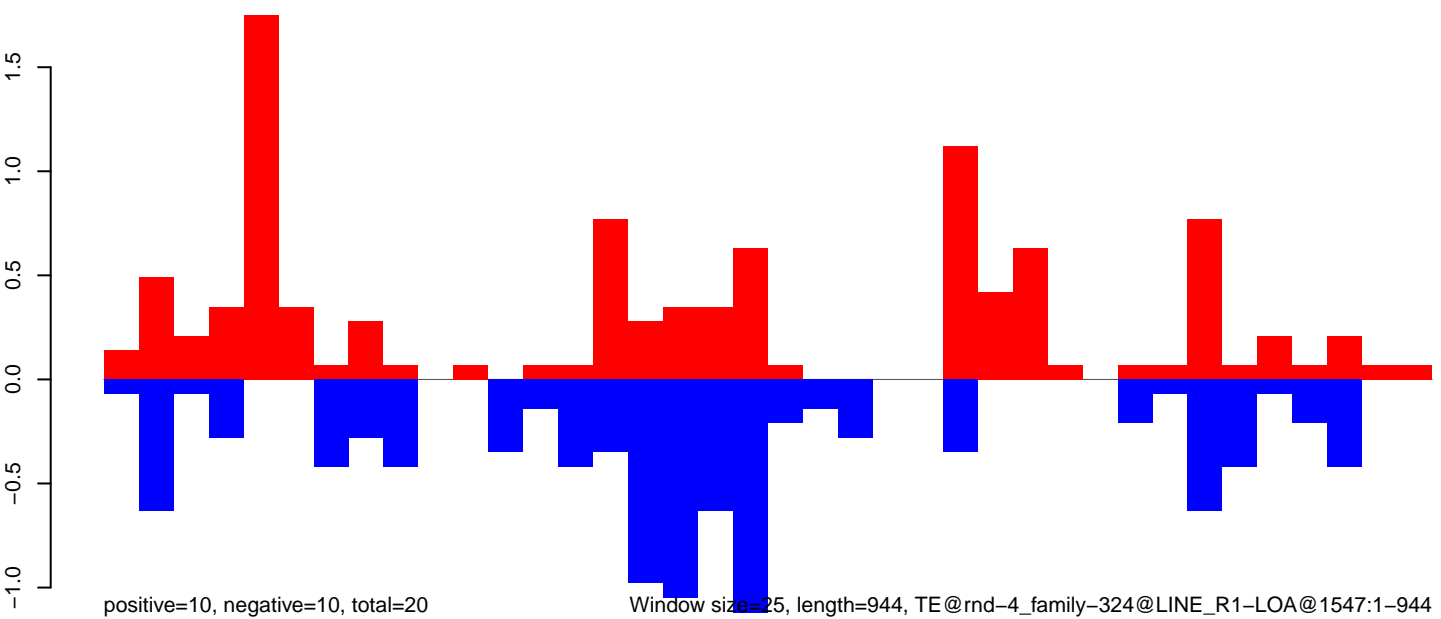
CuQuin_Mcarc_NL.18_23.rep



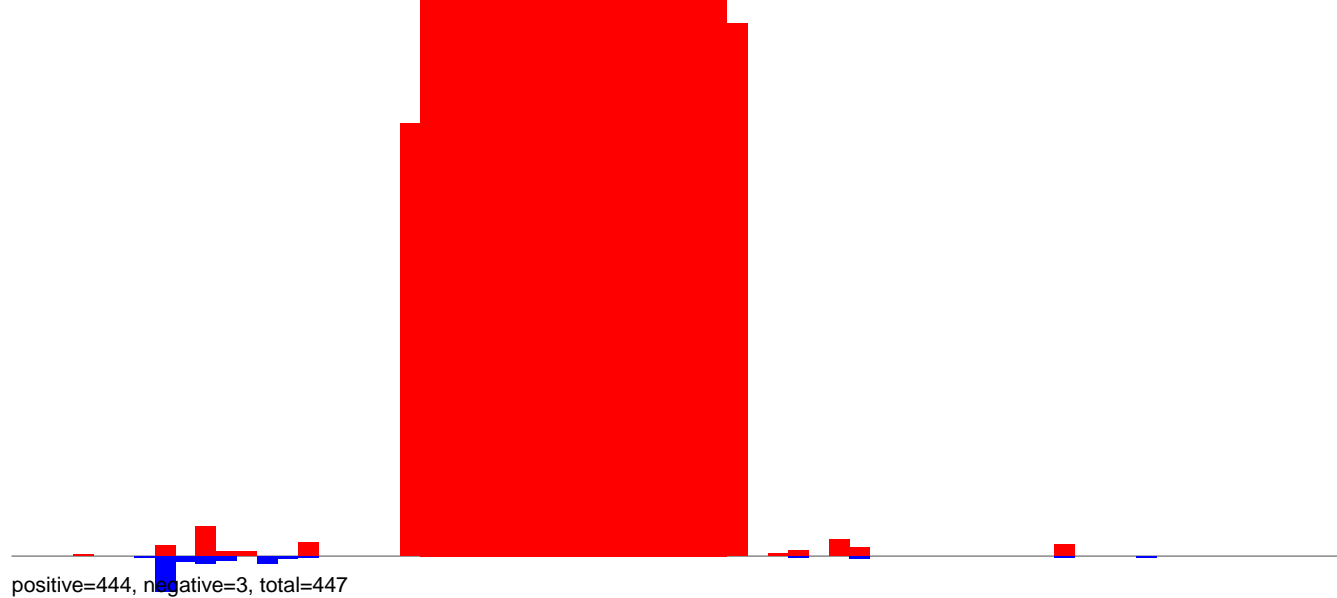
CuQuin_Mcarc_NL.24_35.rep



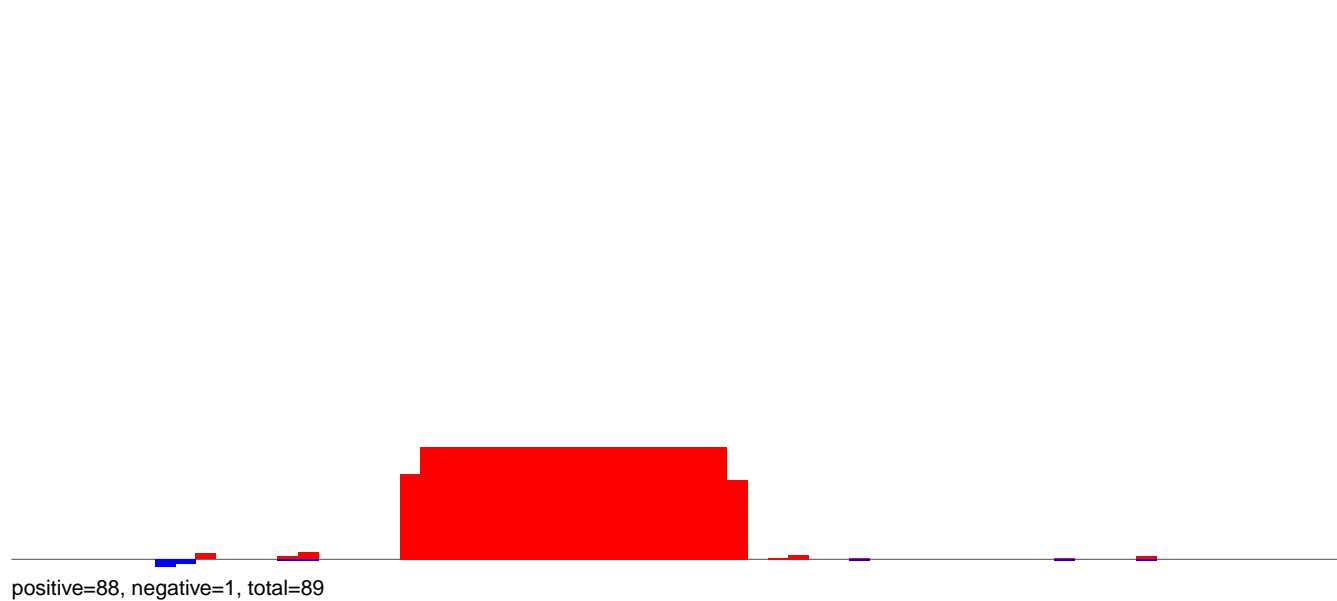
CuQuin_Mcarc_NL.rep



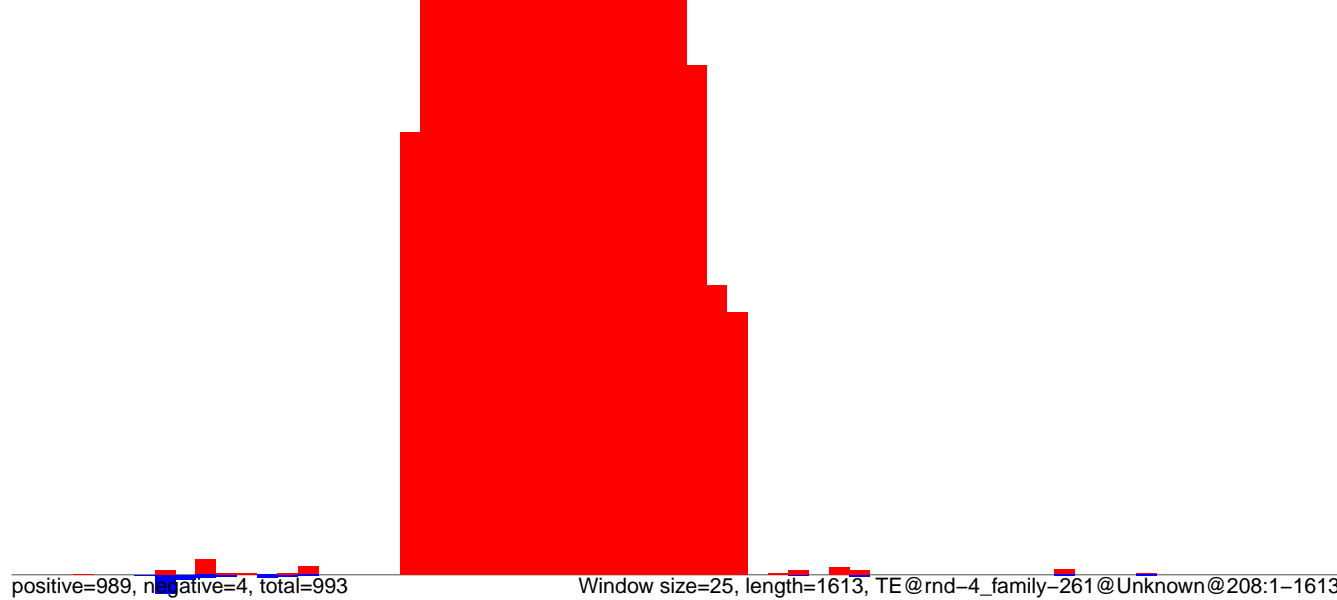
CuQuin_Mcarc_NL.18_23.rep



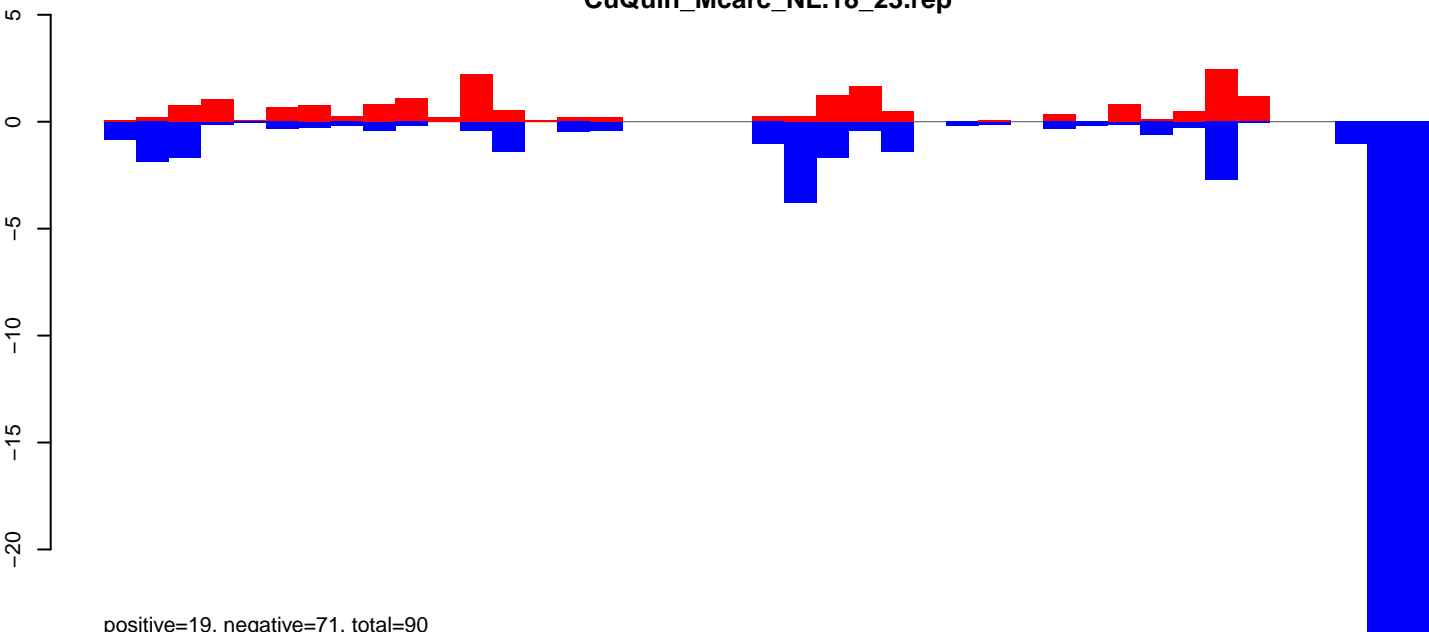
CuQuin_Mcarc_NL.24_35.rep



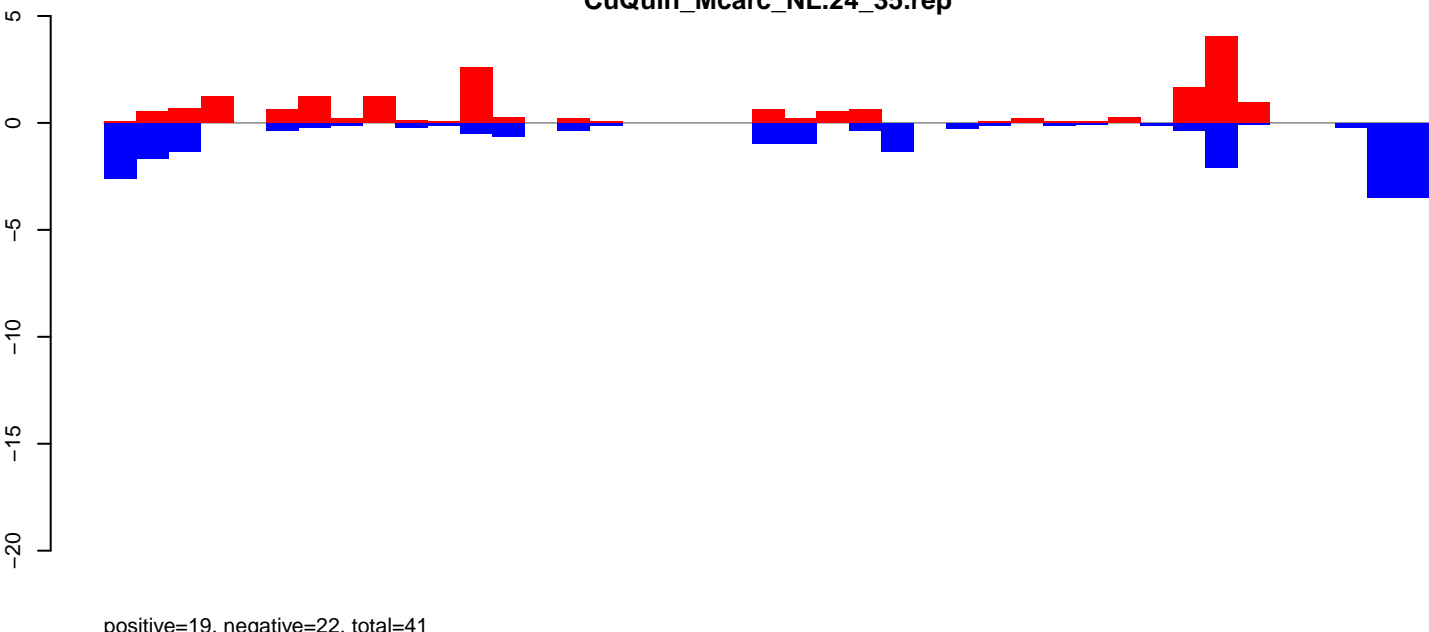
CuQuin_Mcarc_NL.rep



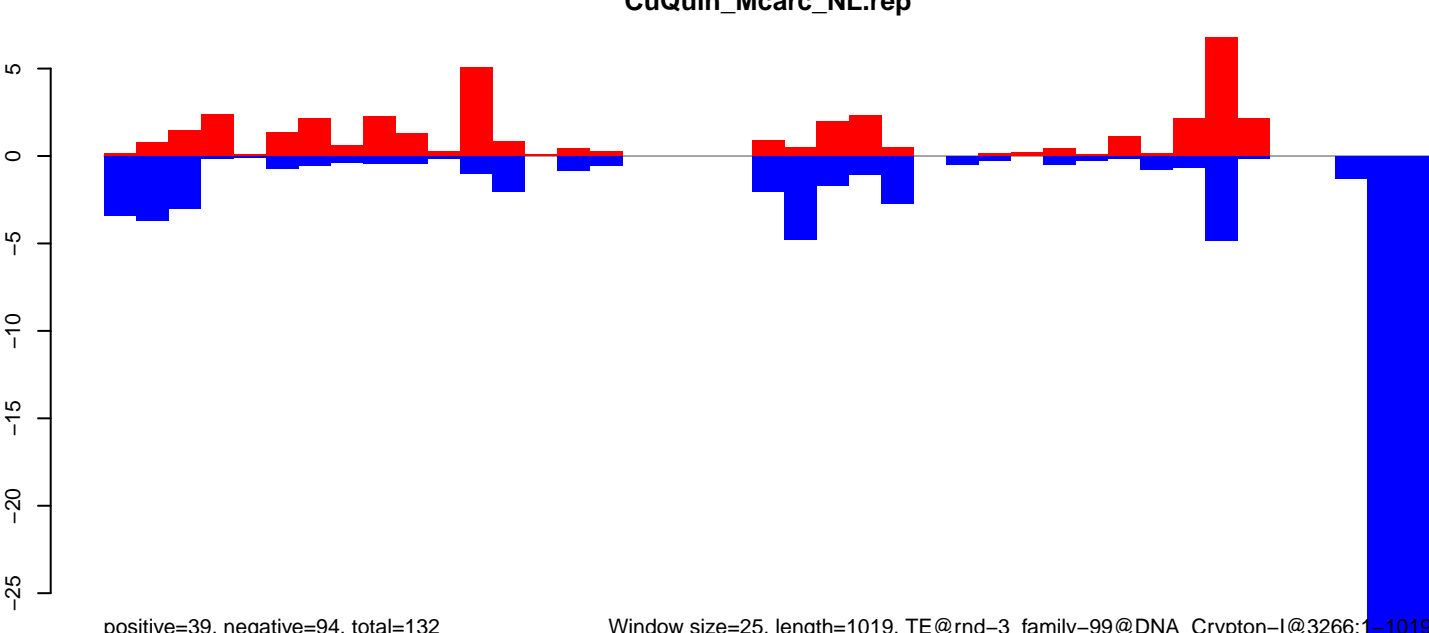
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



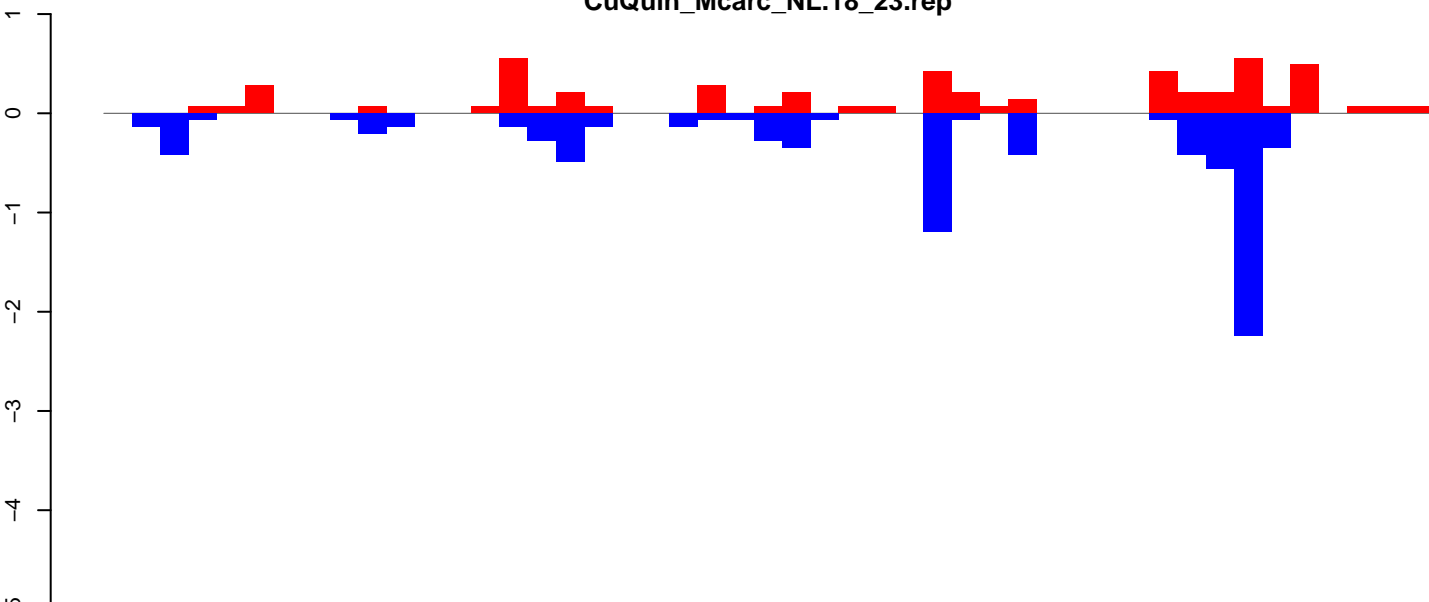
CuQuin_Mcarc_NL.rep



Window size=25, length=1019, TE@rnd-3_family-99@DNA_Crypton-I@3266:1-1019

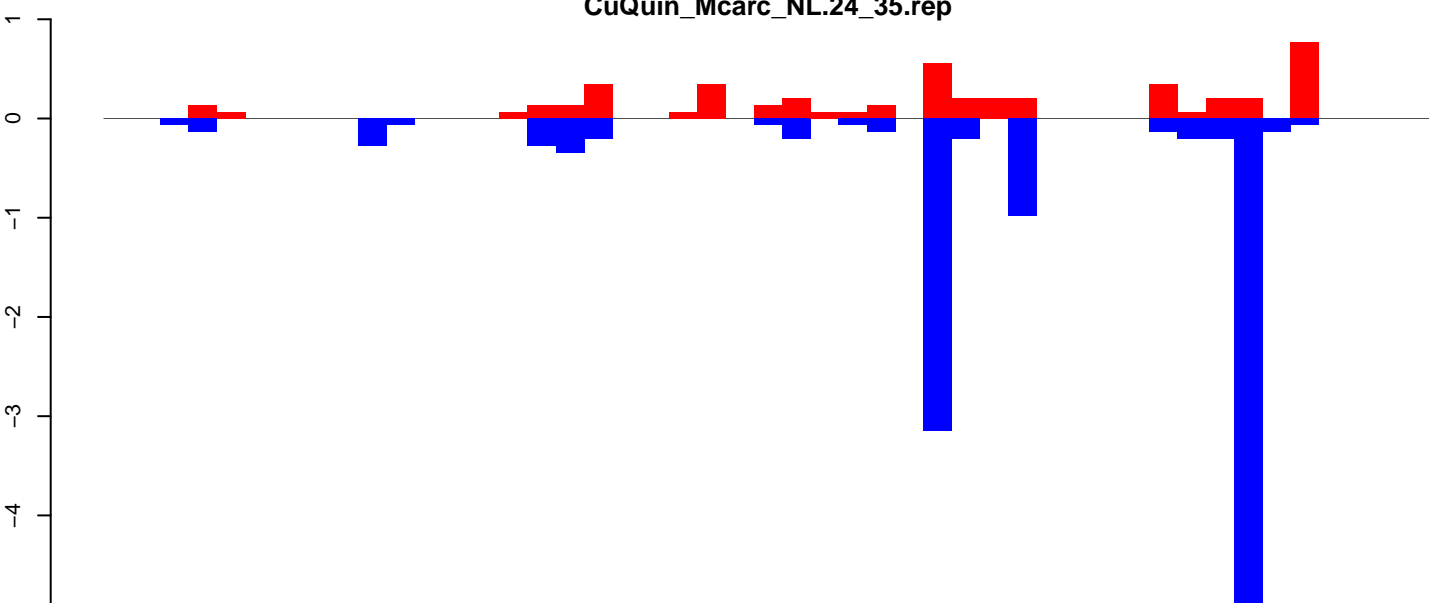
0 200 400 600 800 1000

CuQuin_Mcarc_NL.18_23.rep



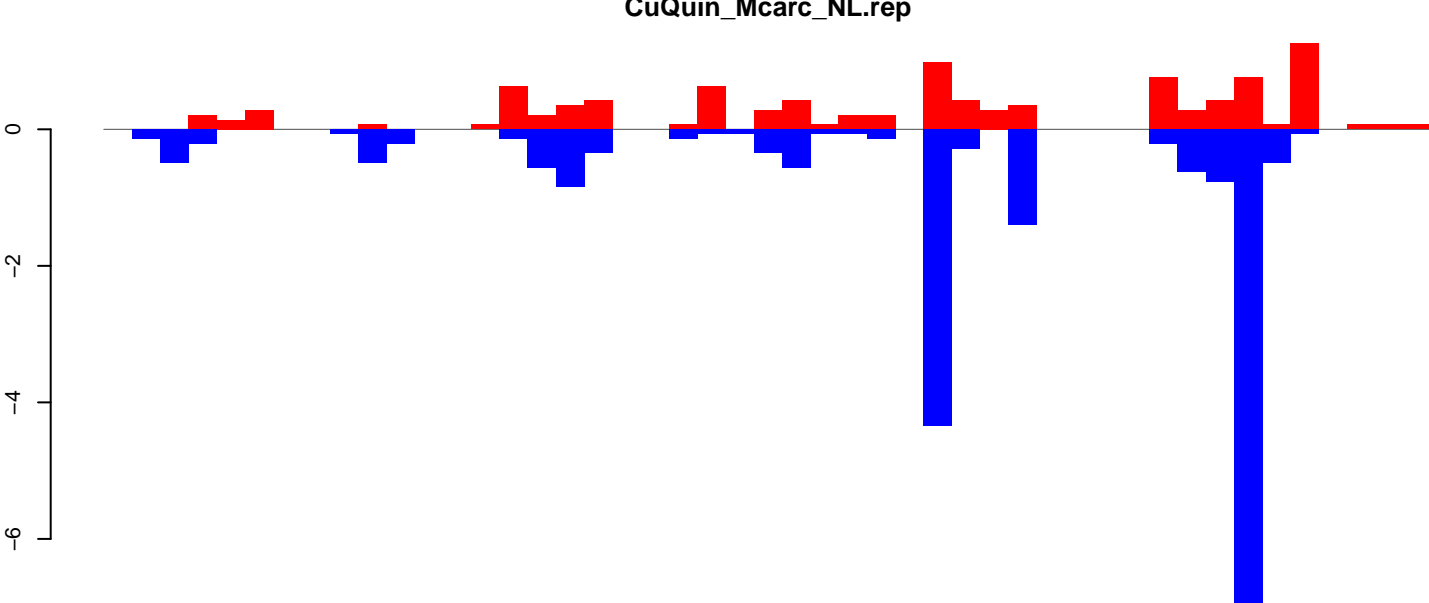
positive=5, negative=8, total=14

CuQuin_Mcarc_NL.24_35.rep



positive=5, negative=12, total=17

CuQuin_Mcarc_NL.rep

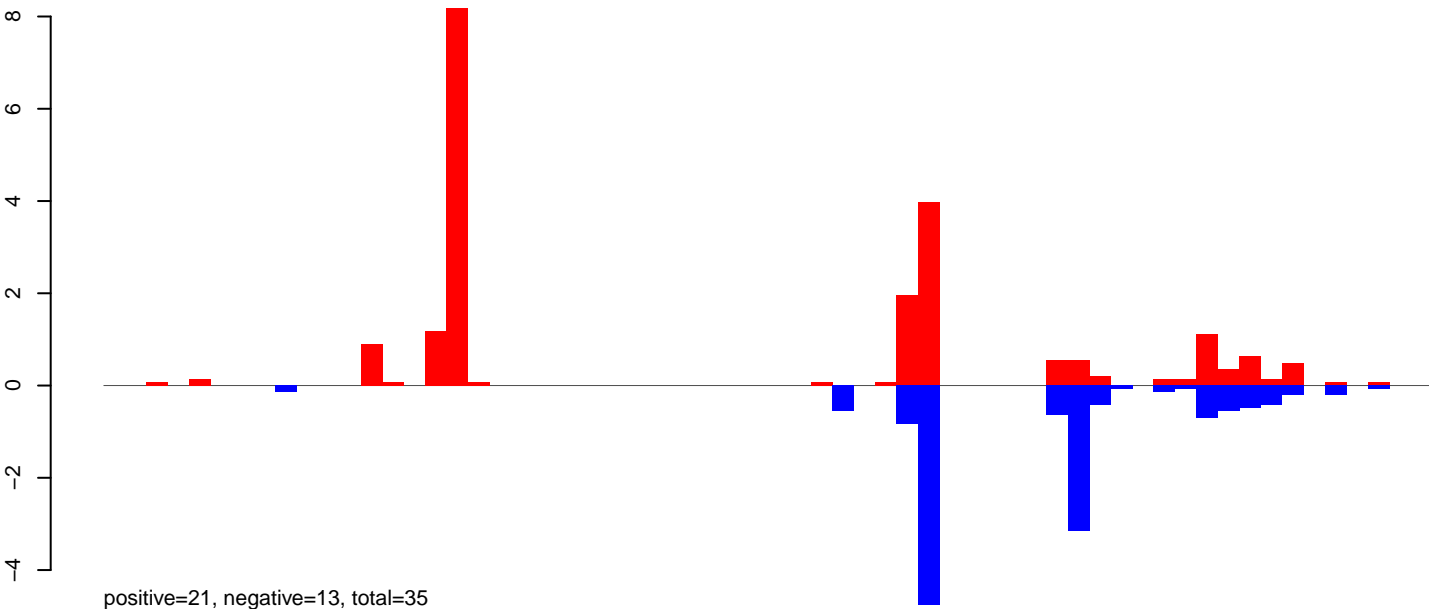


positive=10, negative=21, total=31

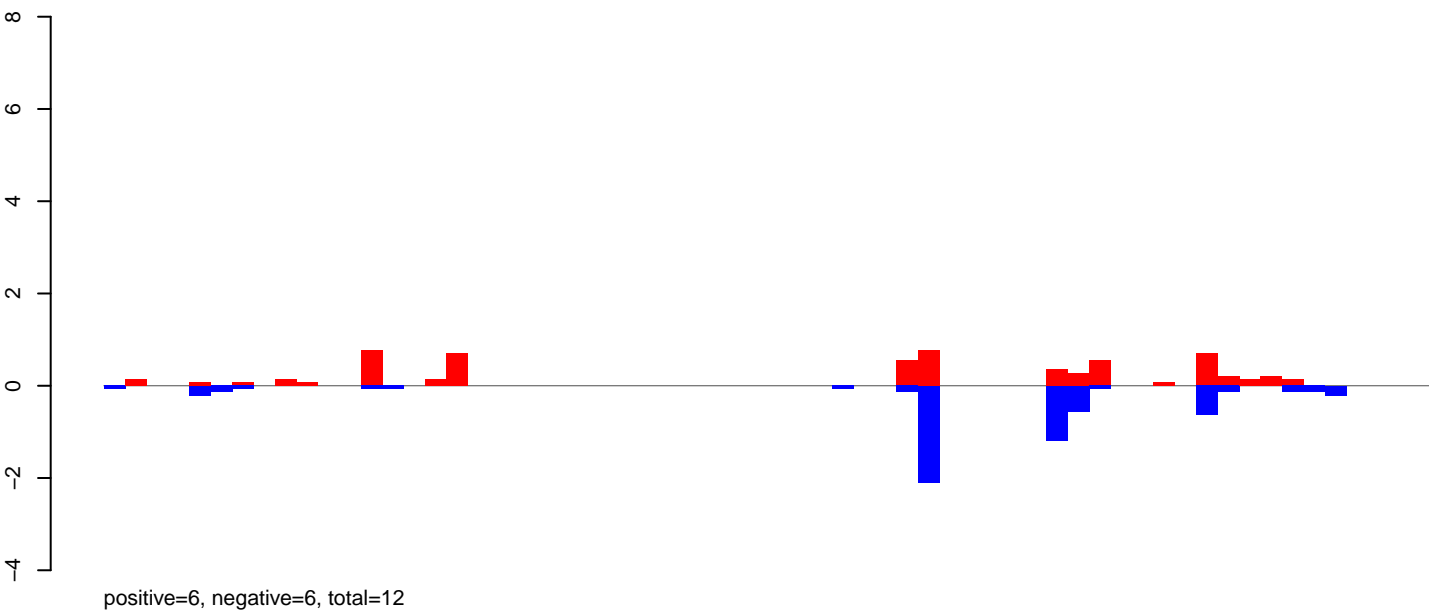
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0 200 400 600 800 1000 1200

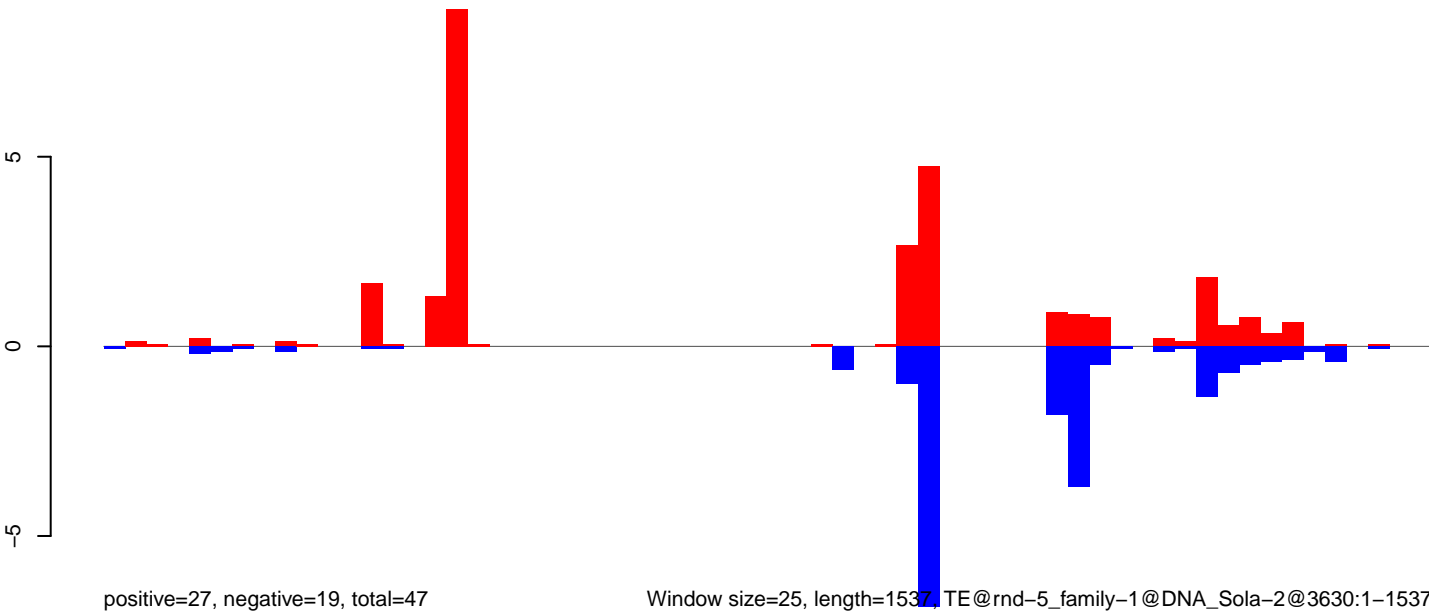
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

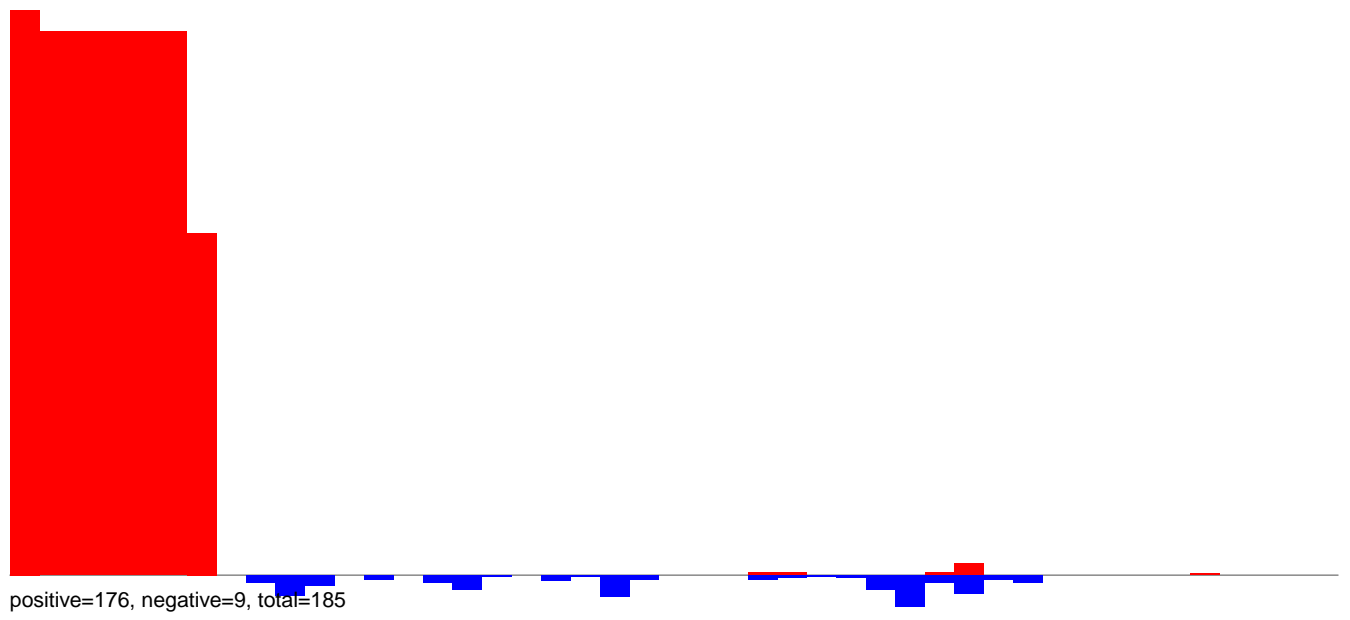


CuQuin_Mcarc_NL.rep

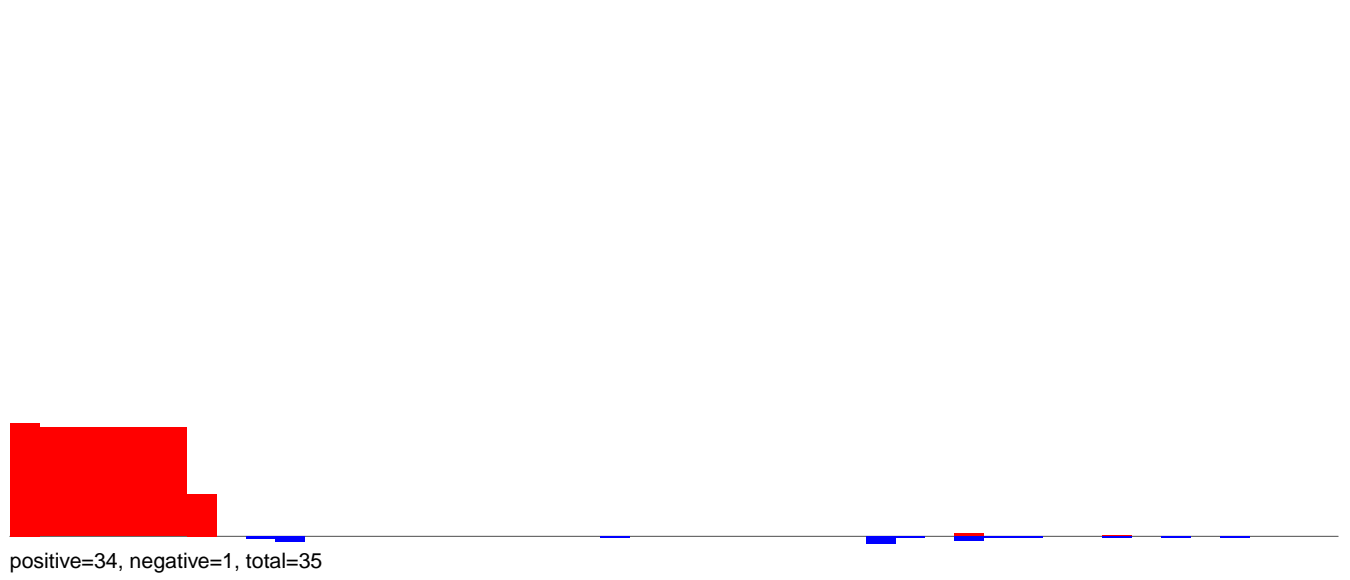


Window size=25, length=1537, TE@rnd-5_family-1@DNA_Sola-2@3630:1-1537

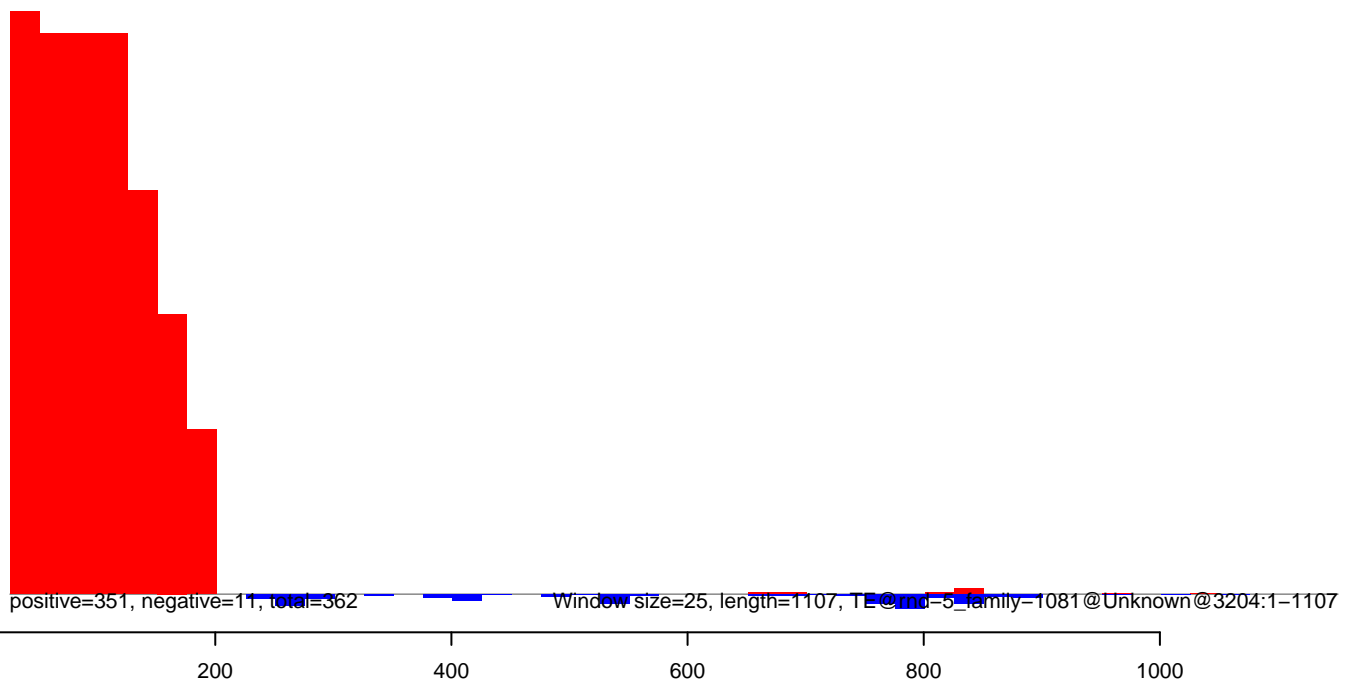
CuQuin_Mcarc_NL.18_23.rep



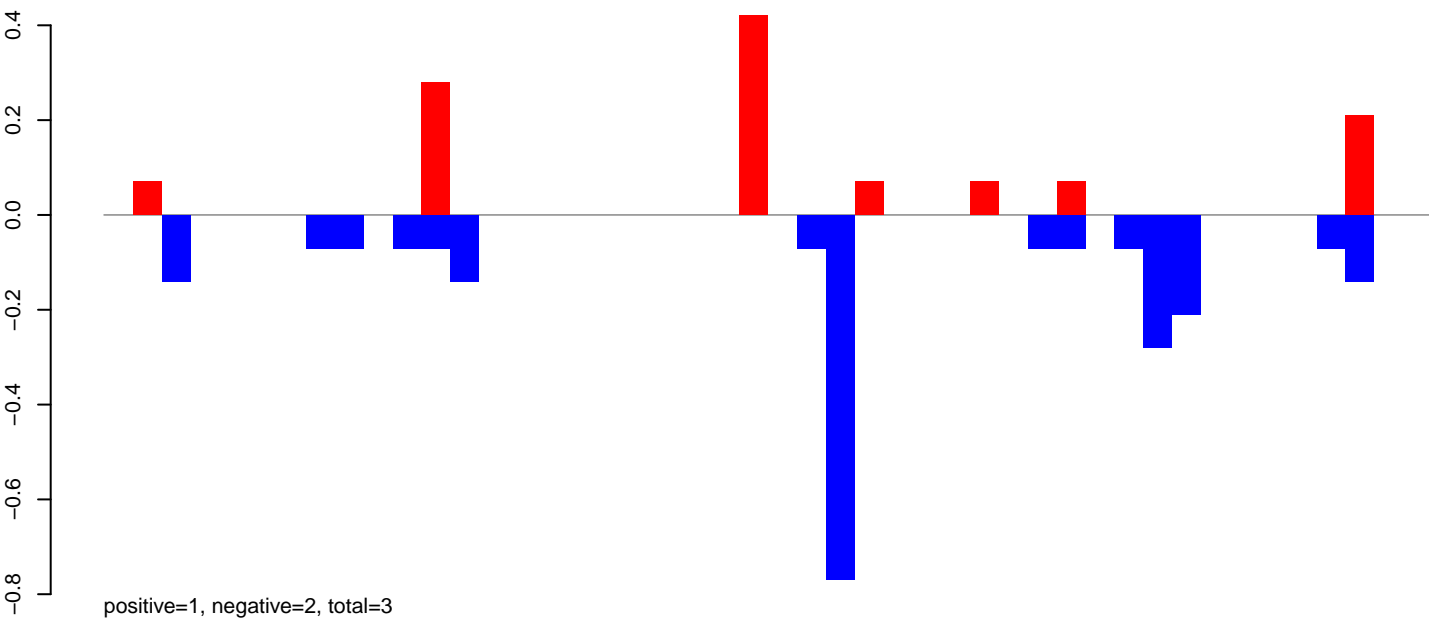
CuQuin_Mcarc_NL.24_35.rep



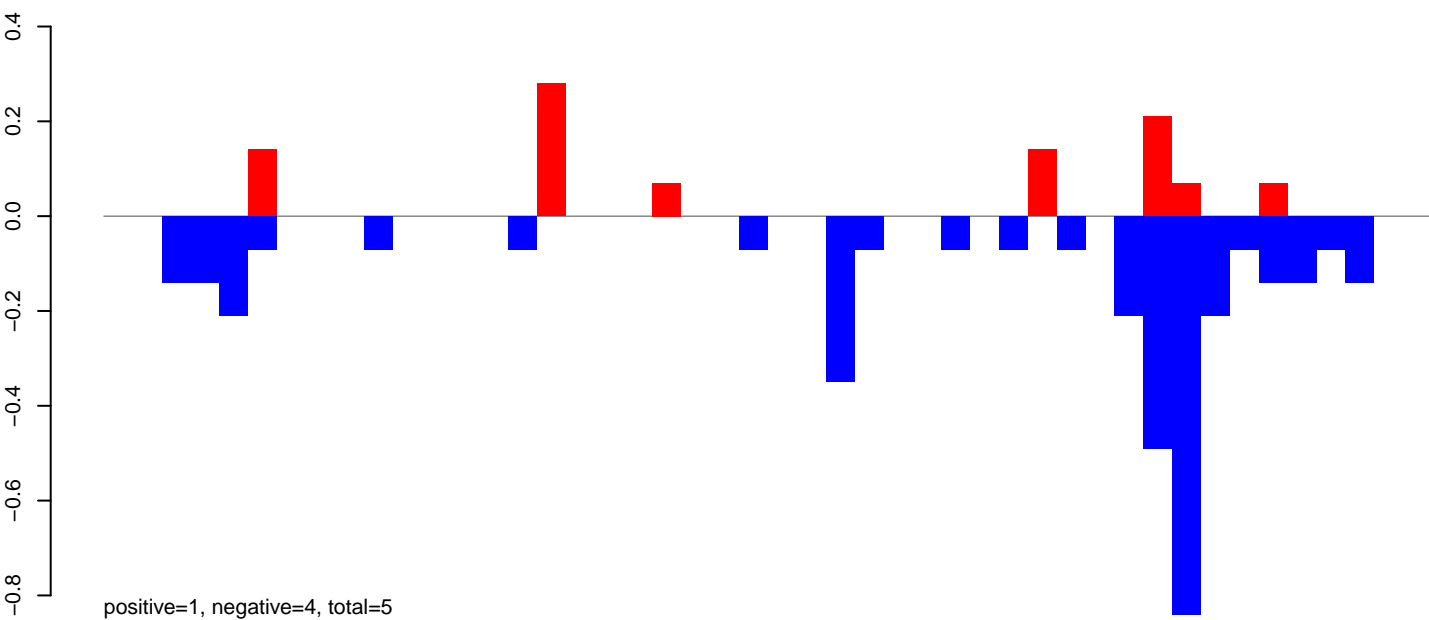
CuQuin_Mcarc_NL.rep



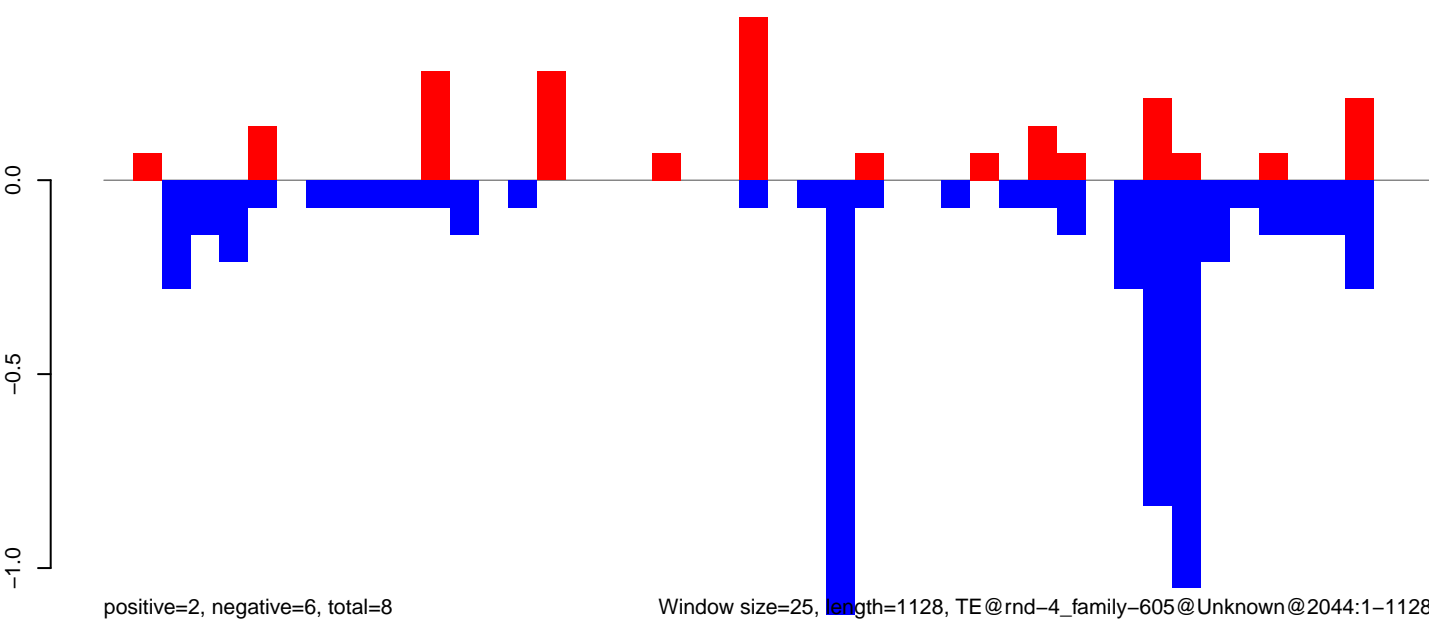
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



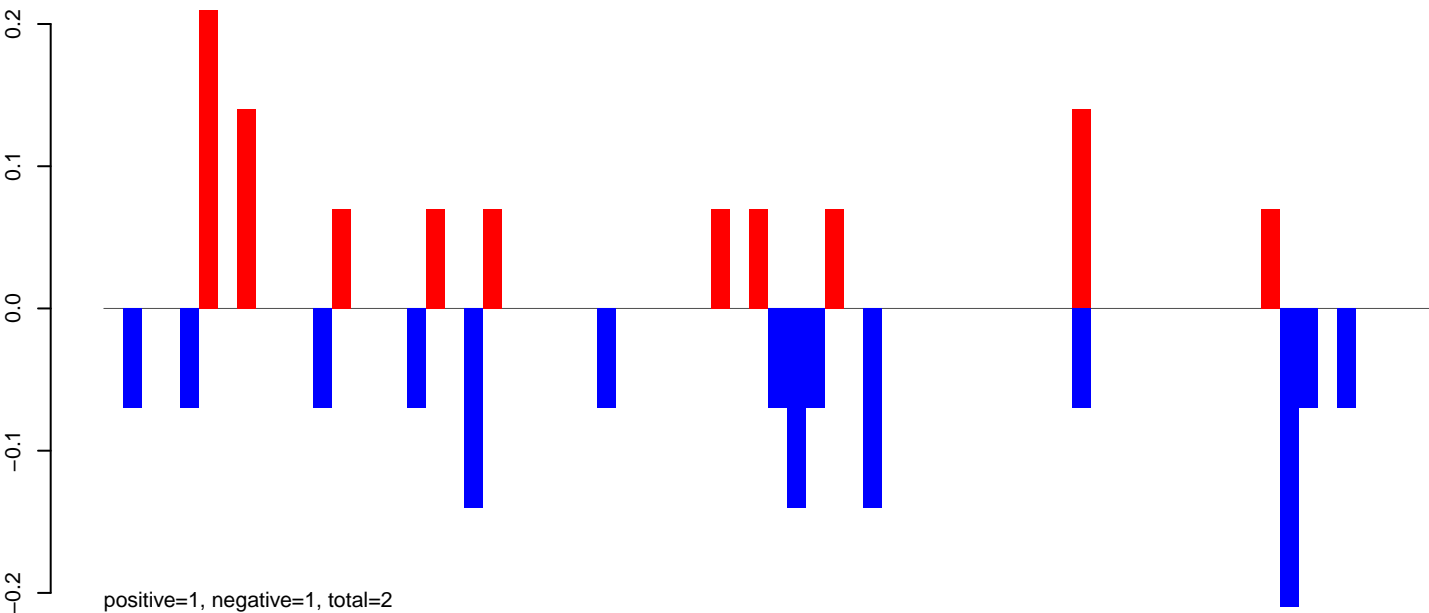
CuQuin_Mcarc_NL.rep



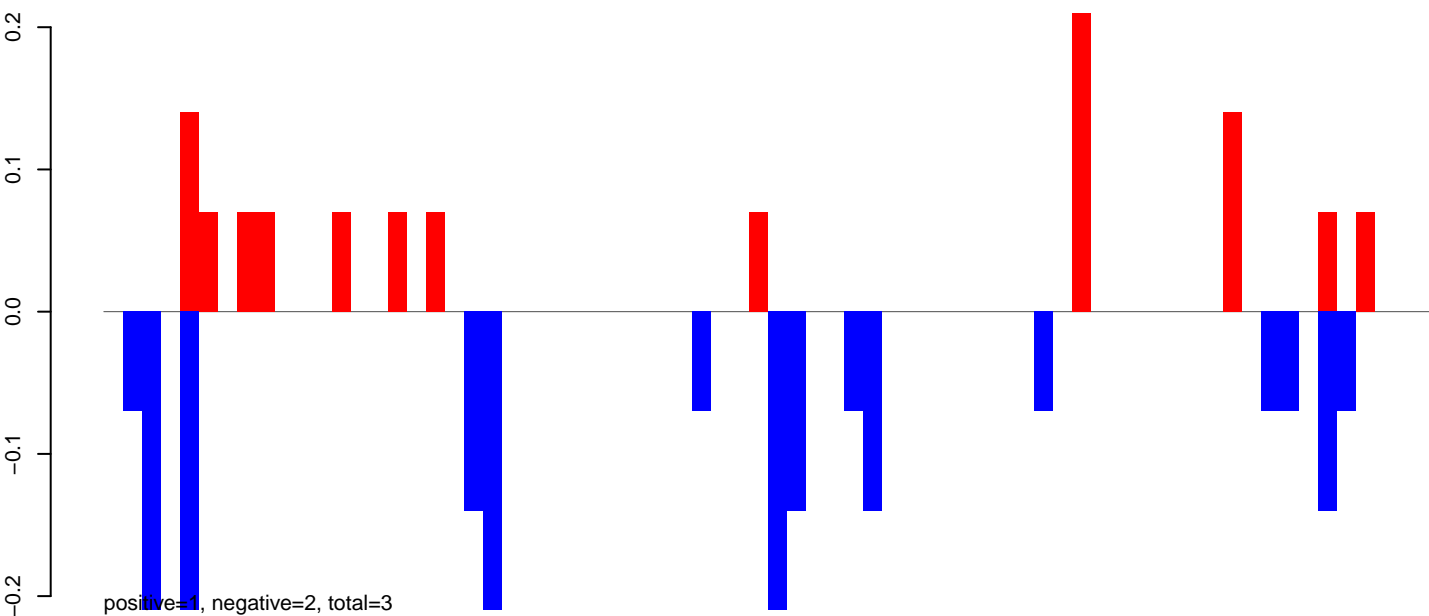
Window size=25, length=1128, TE@rnd-4_family-605@Unknown@2044:1-1128

0 200 400 600 800 1000 1200

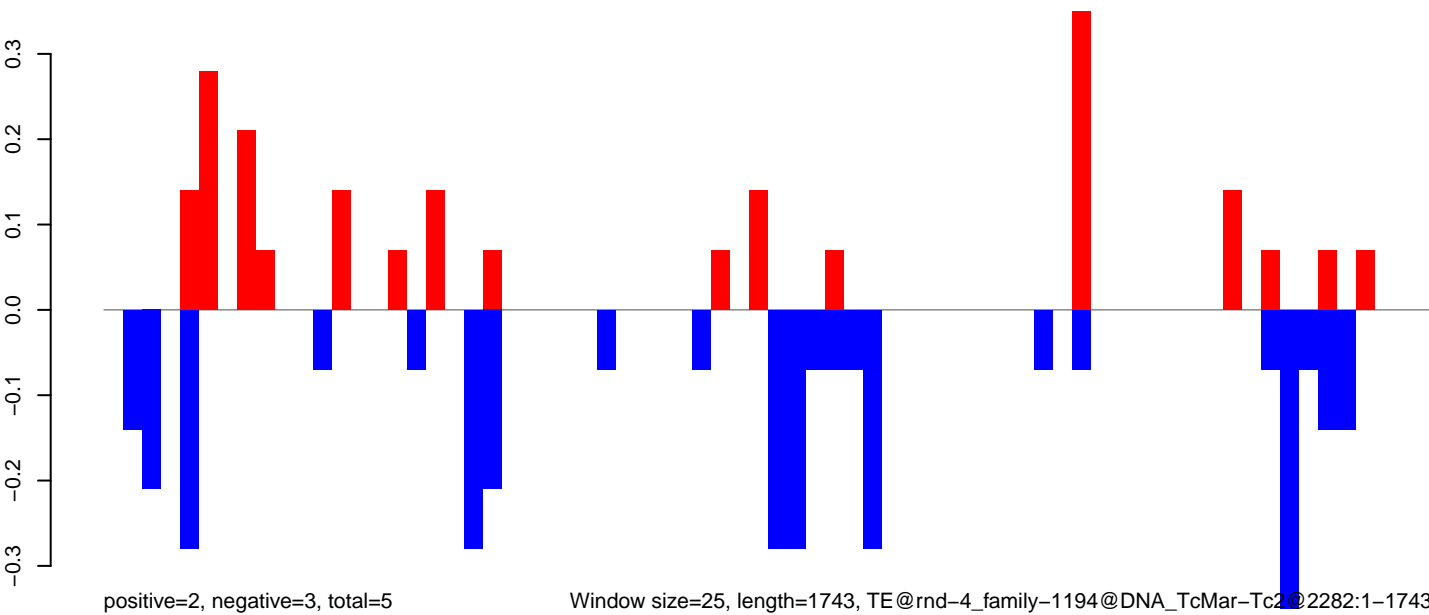
CuQuin_Mcarc_NL.18_23.rep



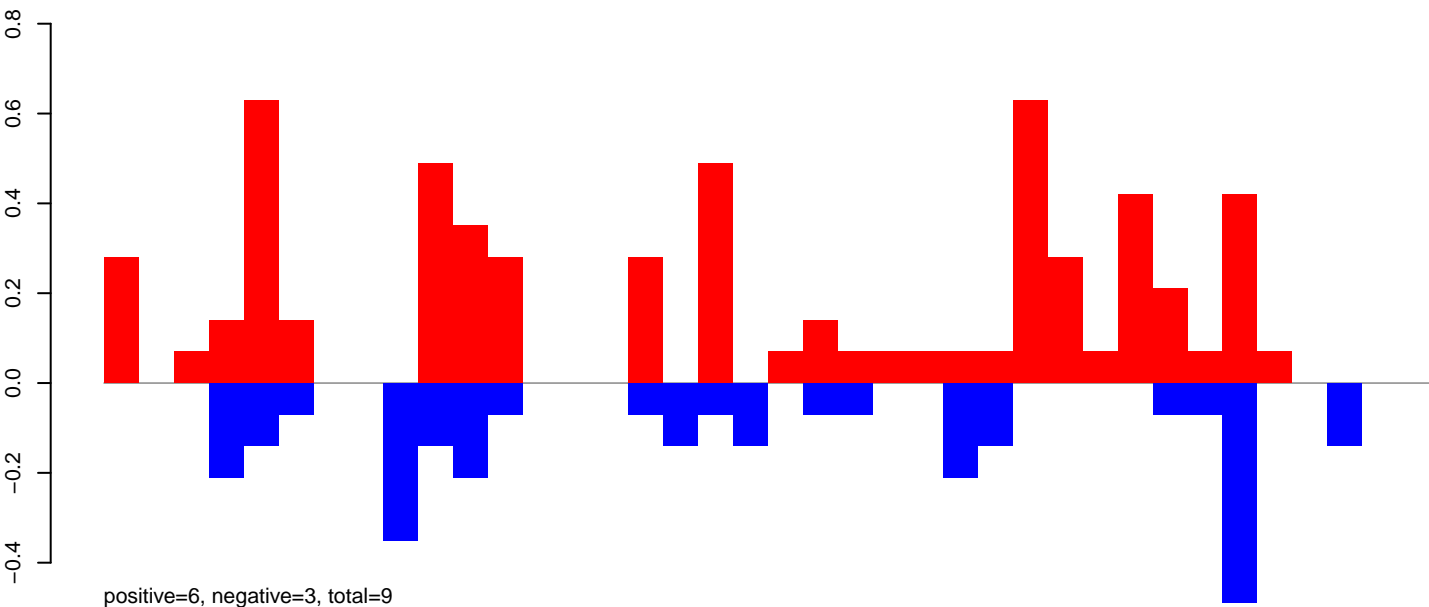
CuQuin_Mcarc_NL.24_35.rep



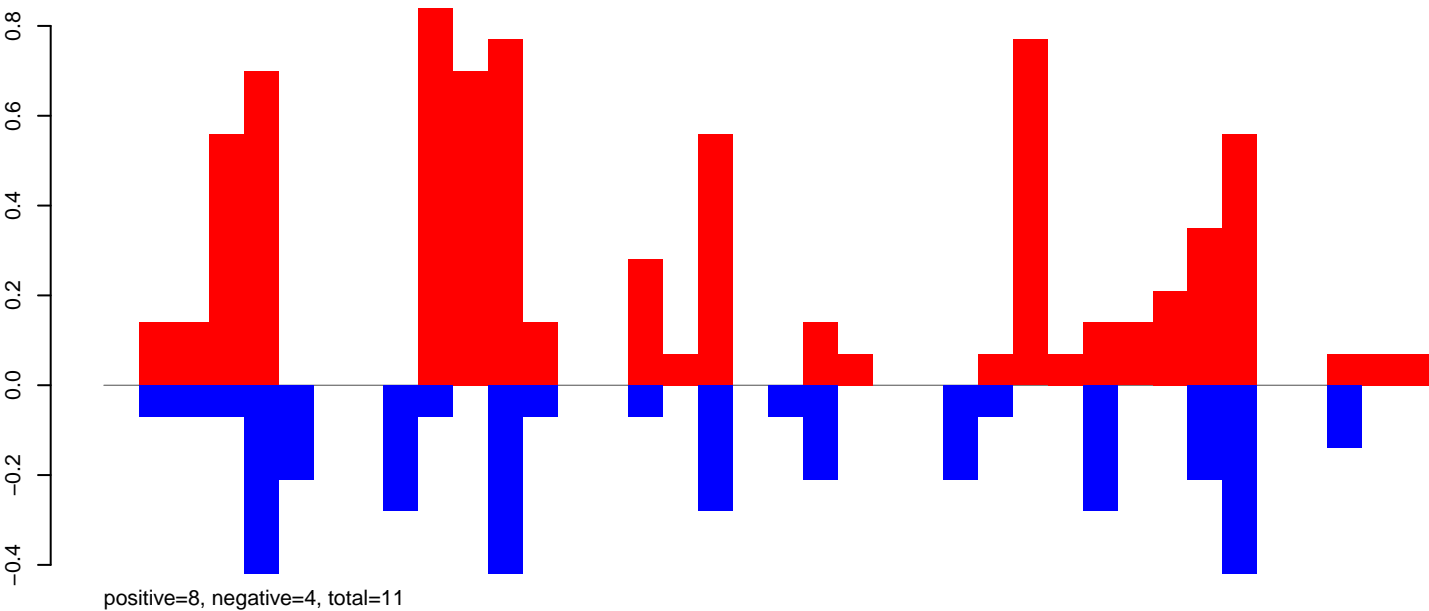
CuQuin_Mcarc_NL.rep



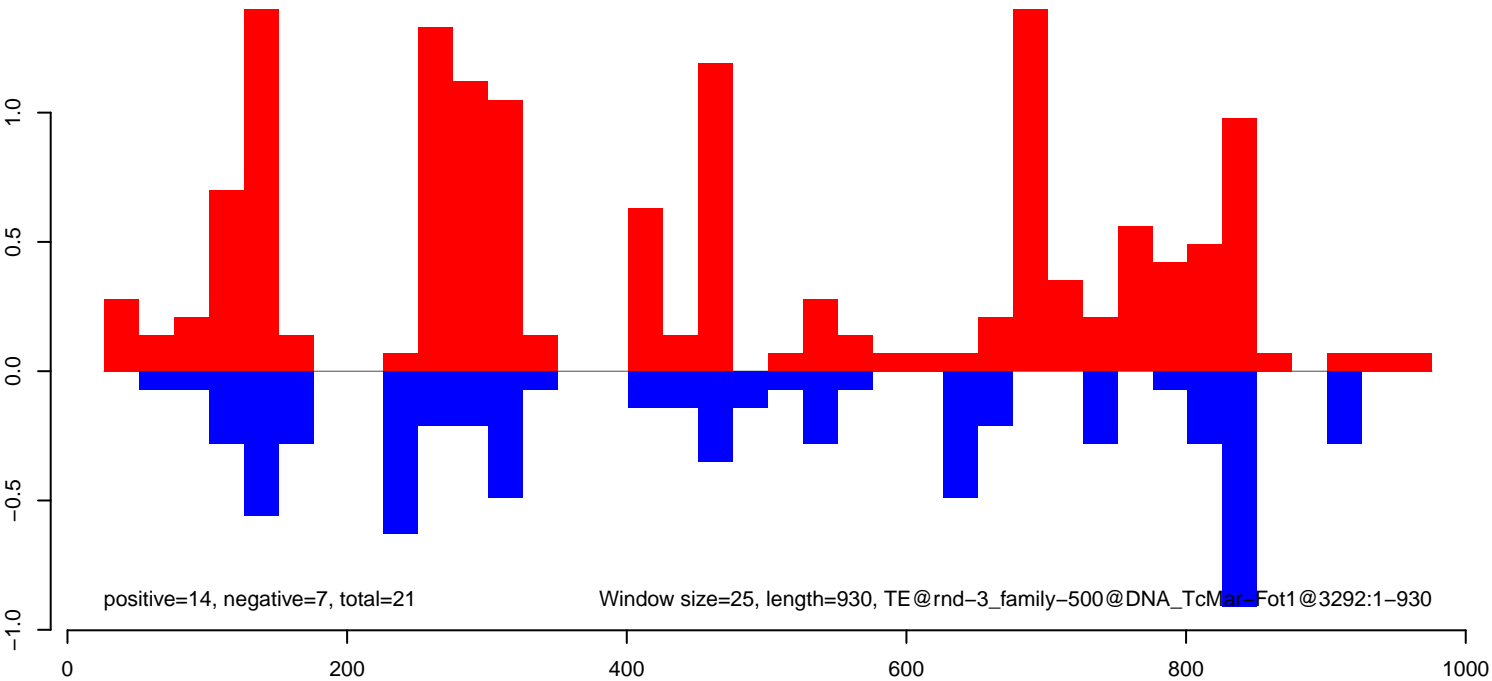
CuQuin_Mcarc_NL.18_23.rep



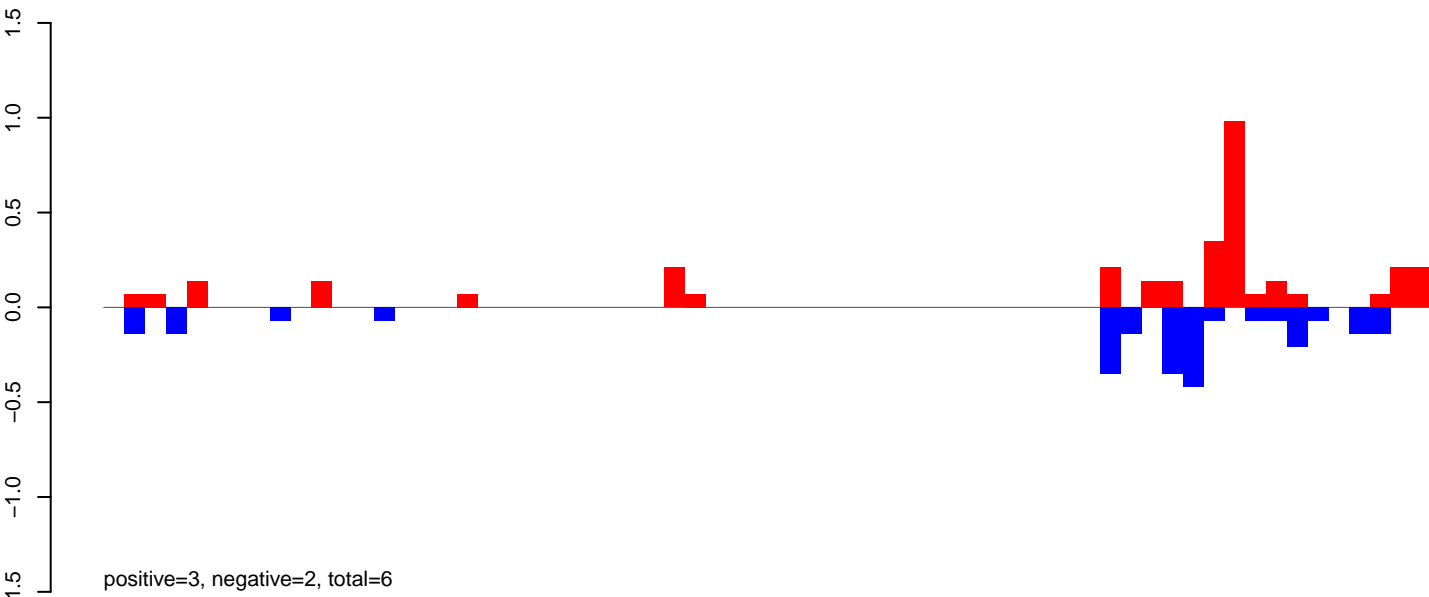
CuQuin_Mcarc_NL.24_35.rep



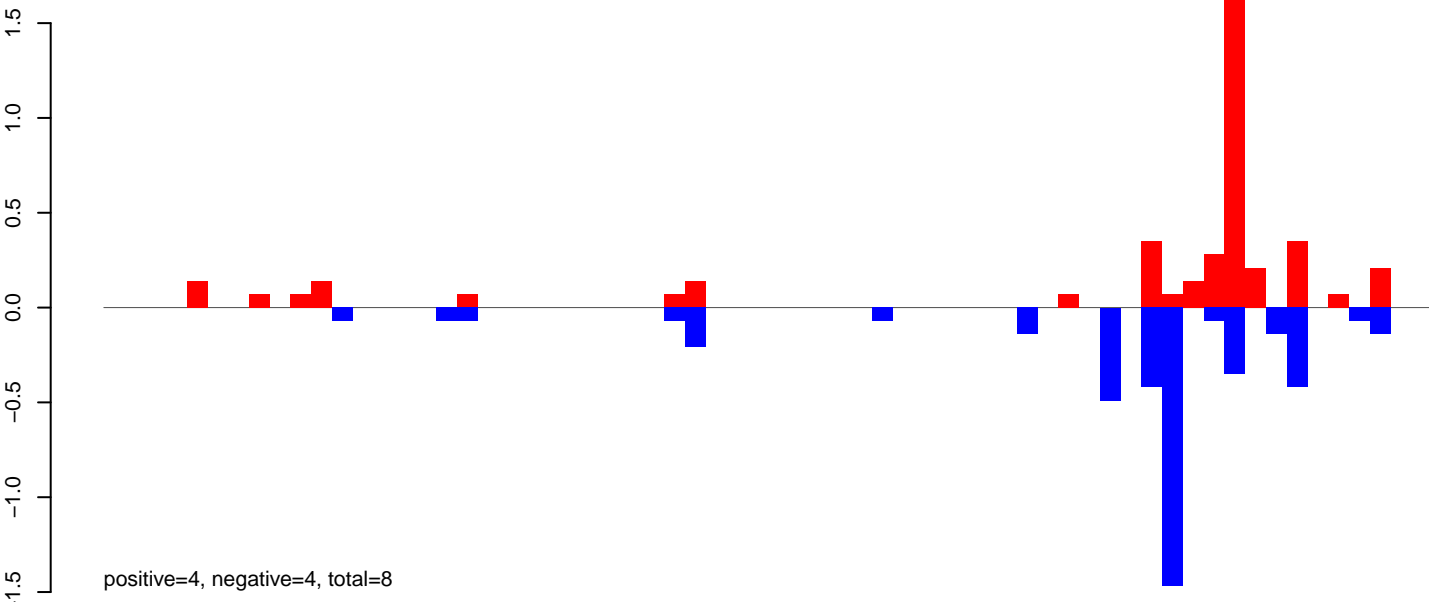
CuQuin_Mcarc_NL.rep



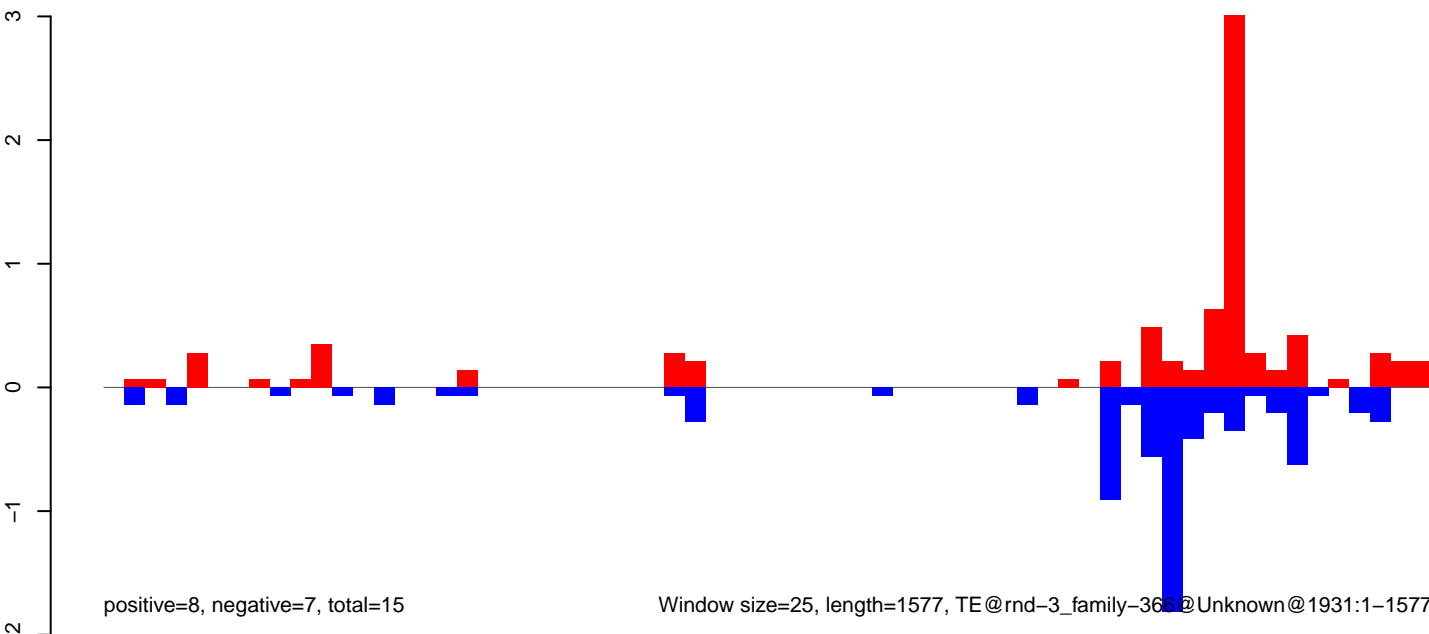
CuQuin_Mcarc_NL.18_23.rep



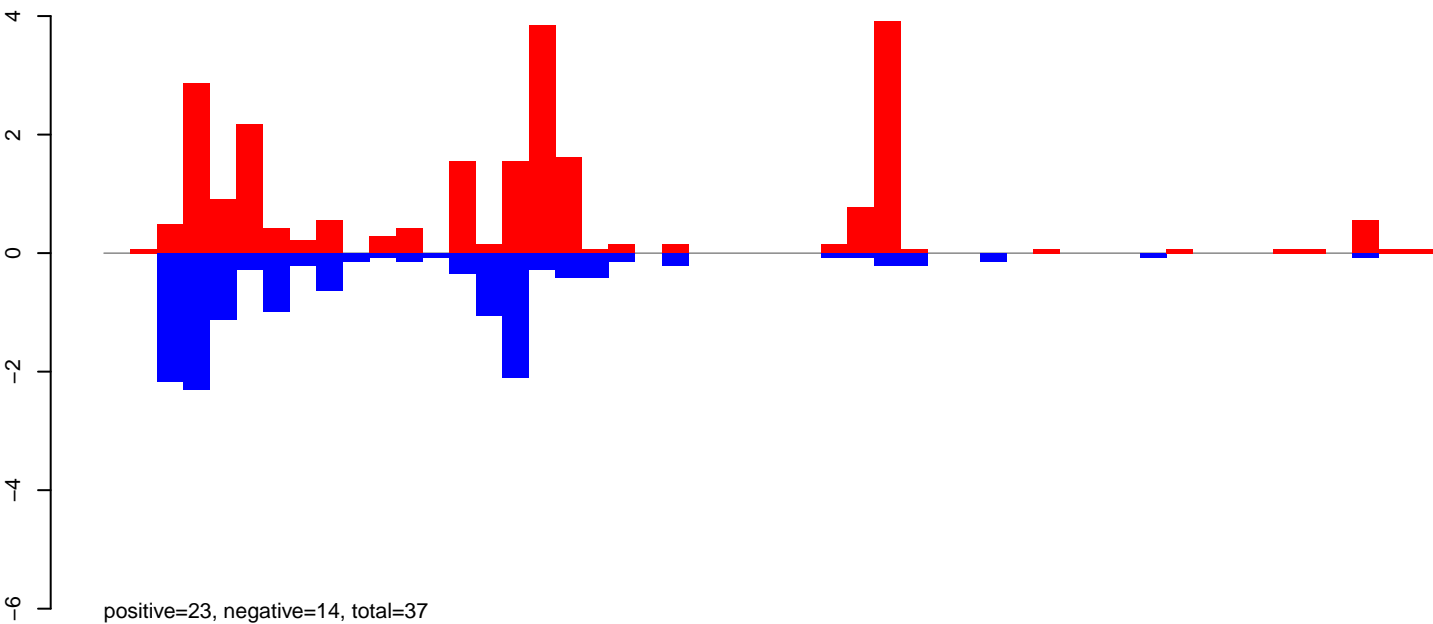
CuQuin_Mcarc_NL.24_35.rep



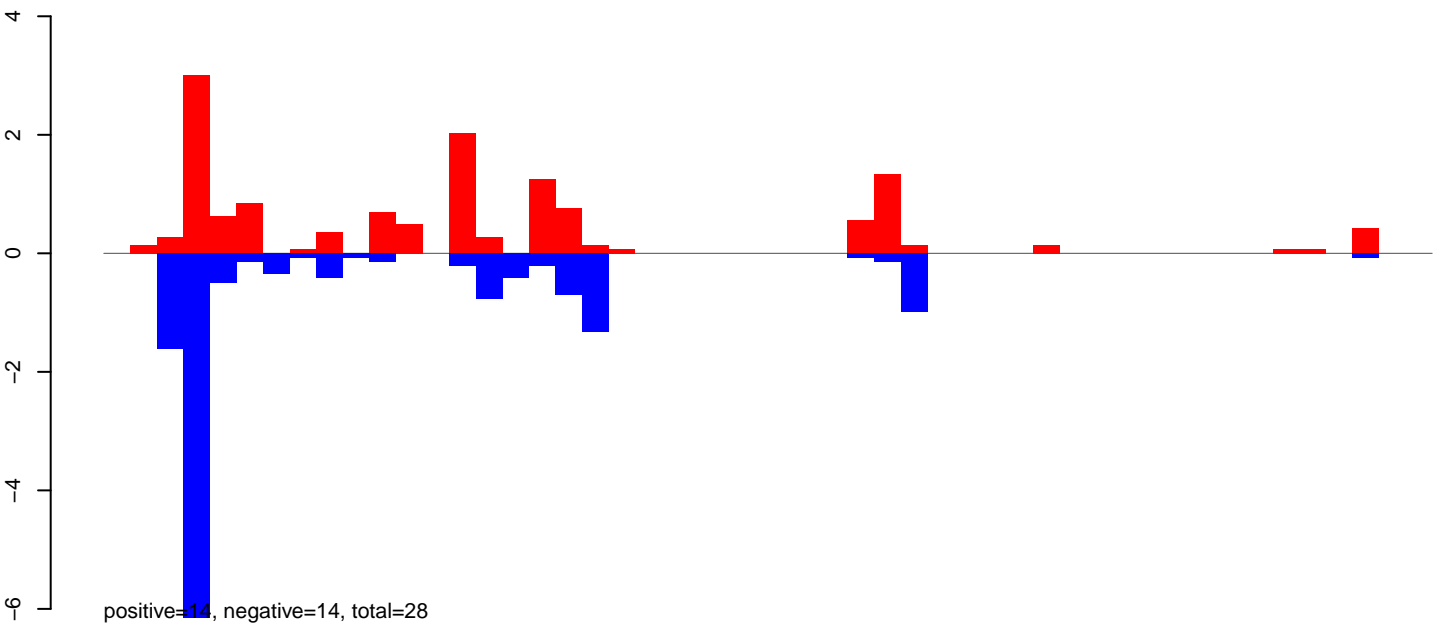
CuQuin_Mcarc_NL.rep



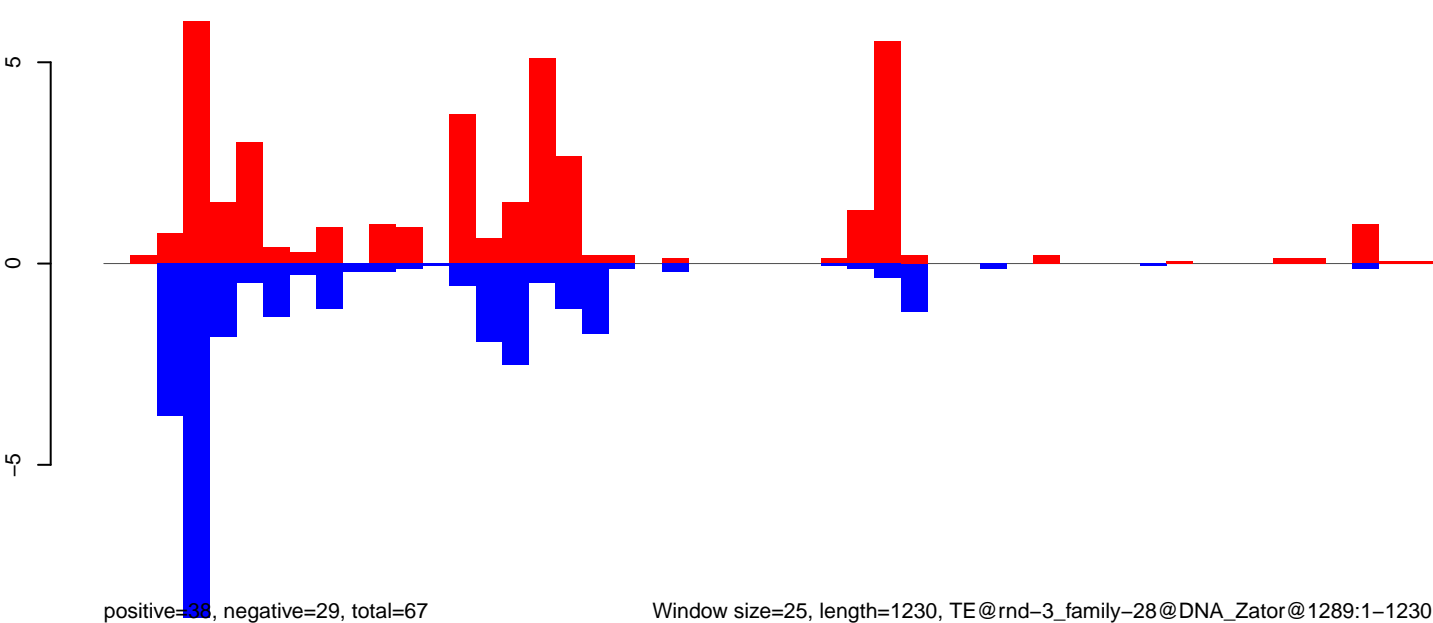
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



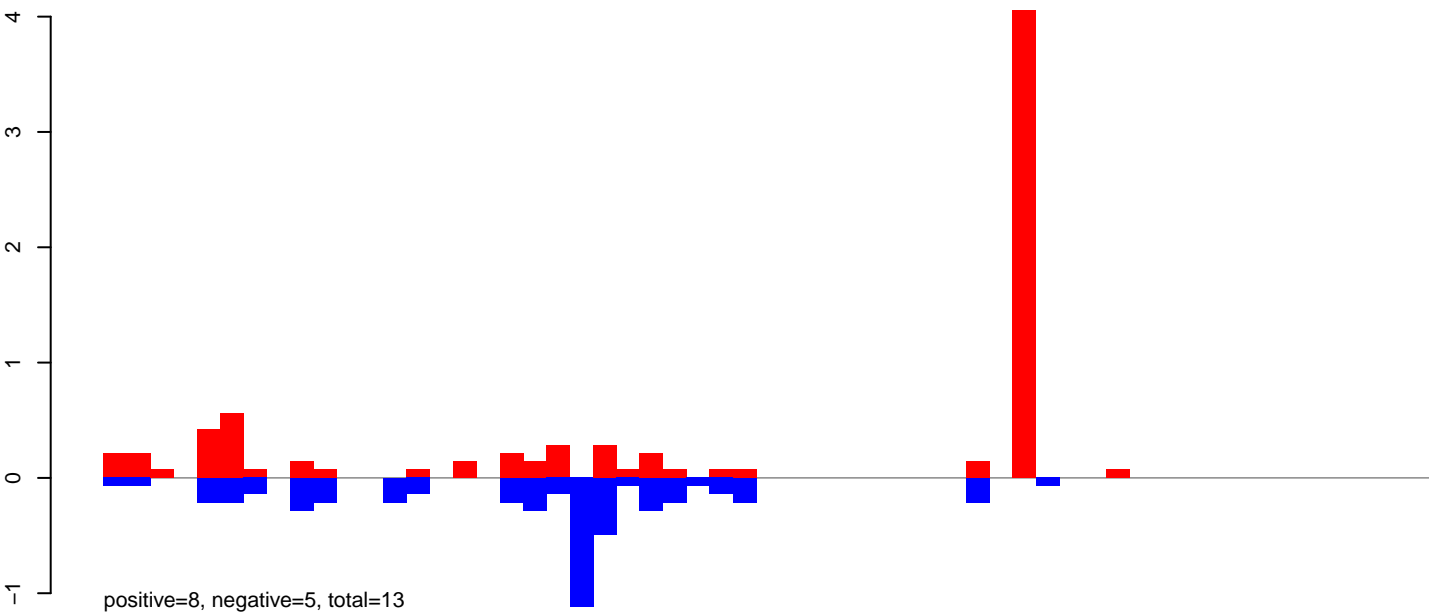
CuQuin_Mcarc_NL.rep



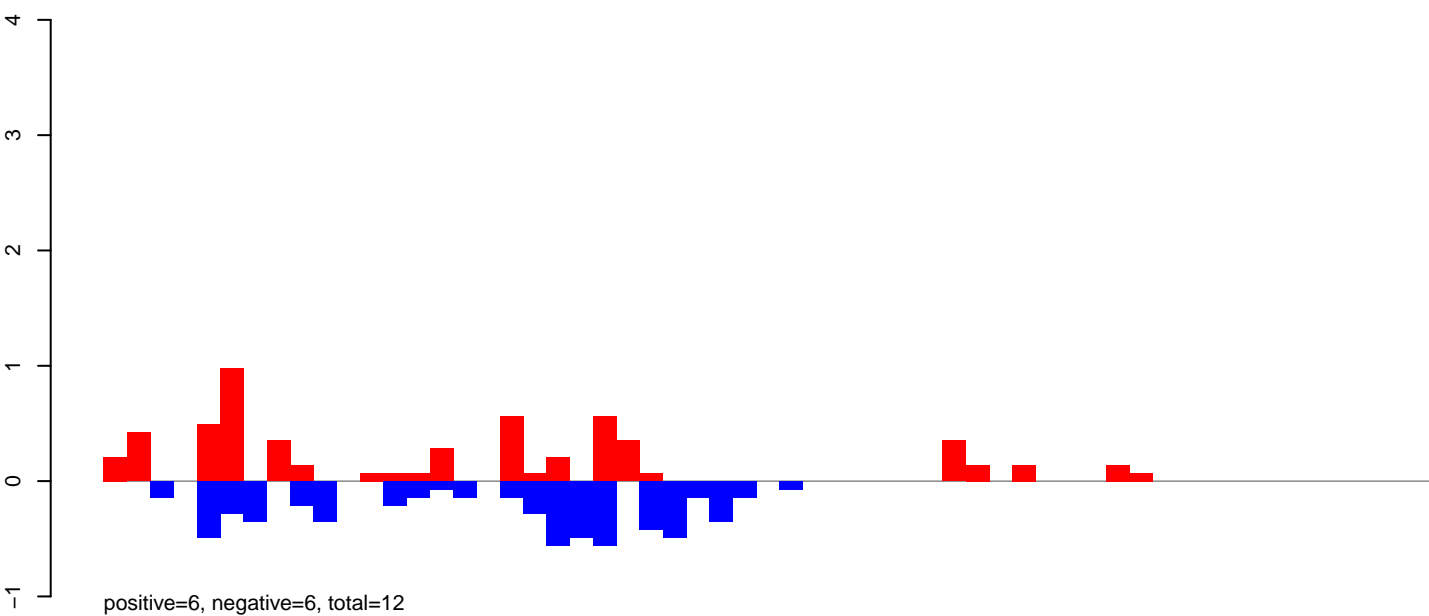
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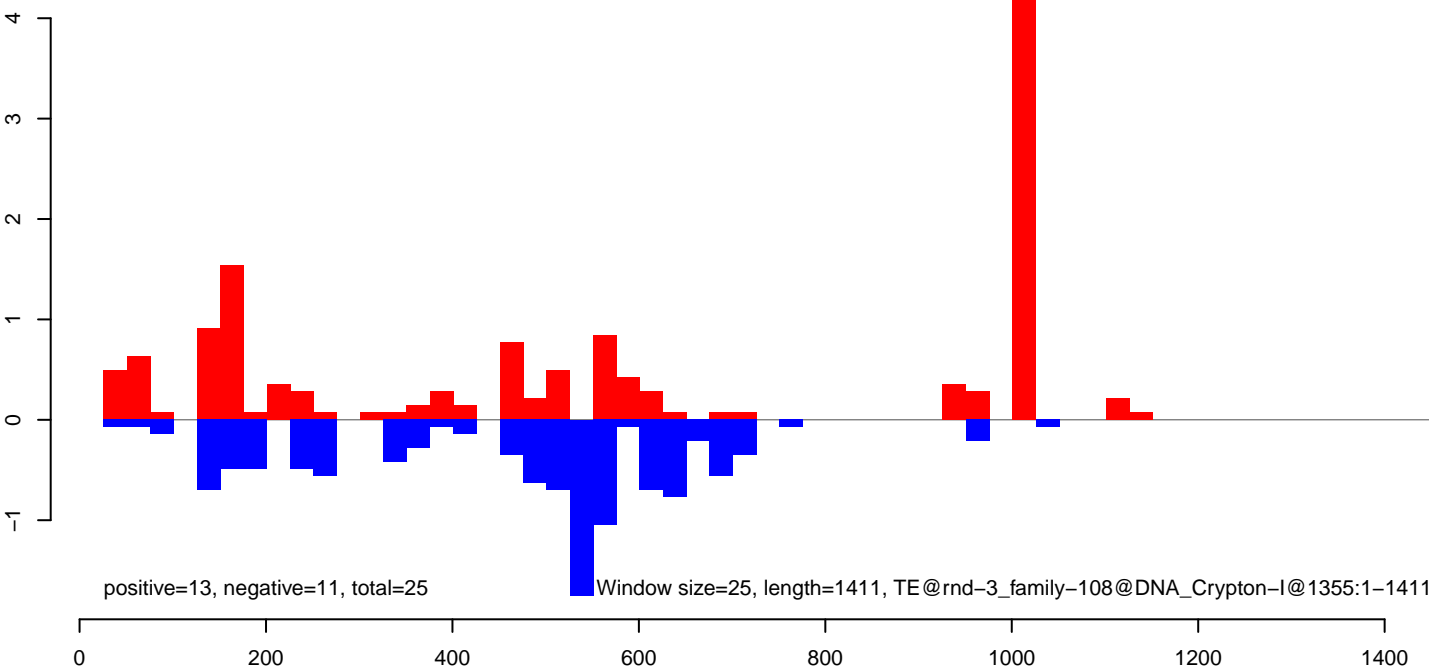
CuQuin_Mcarc_NL.18_23.rep



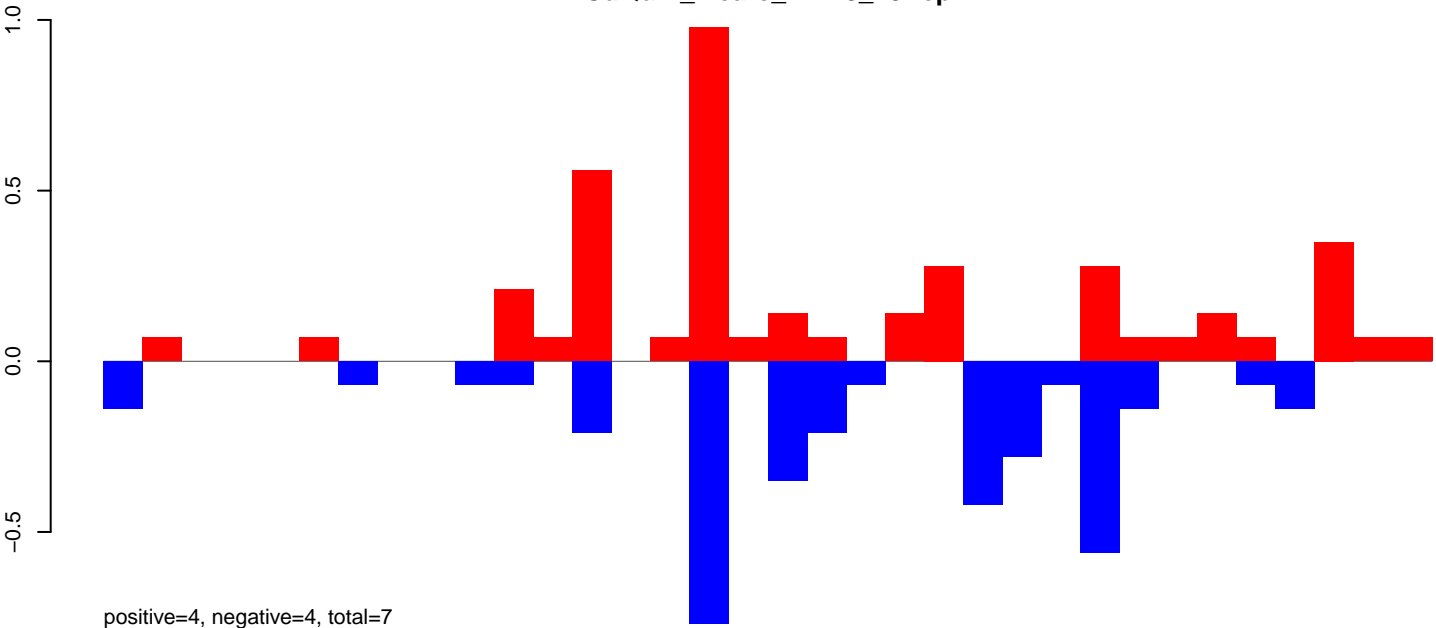
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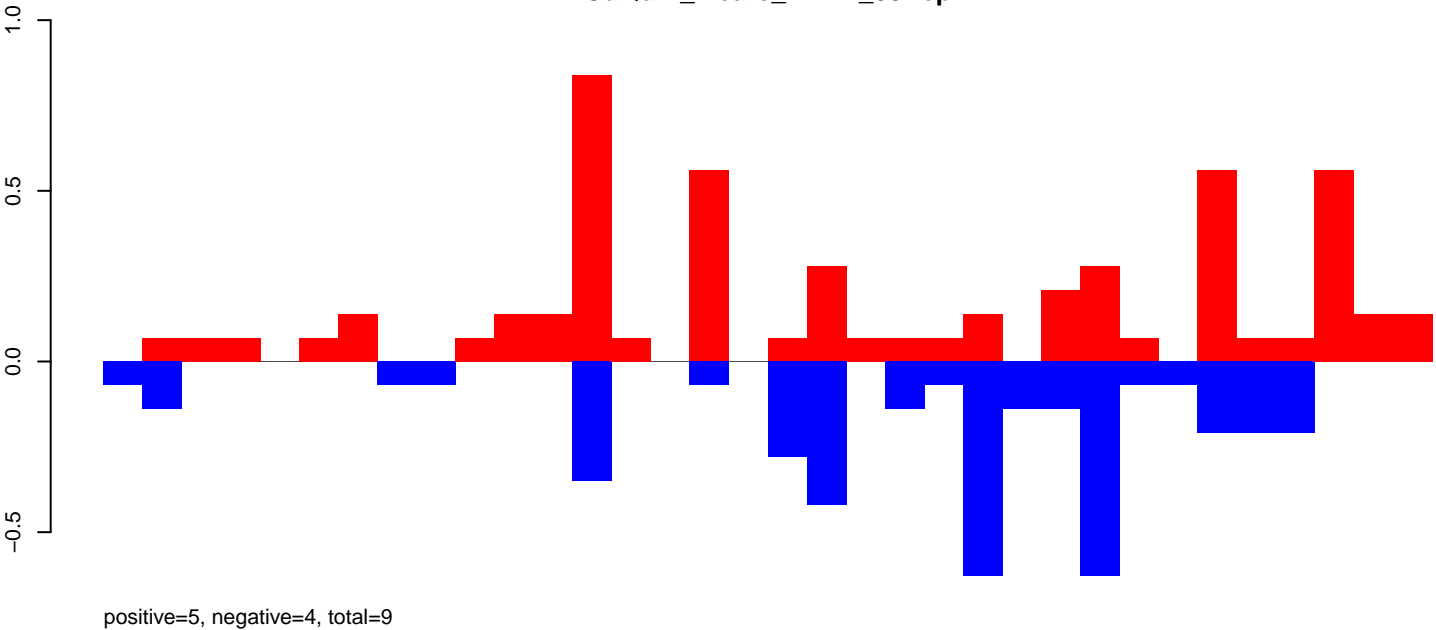
CuQuin_Mcarc_NL.rep



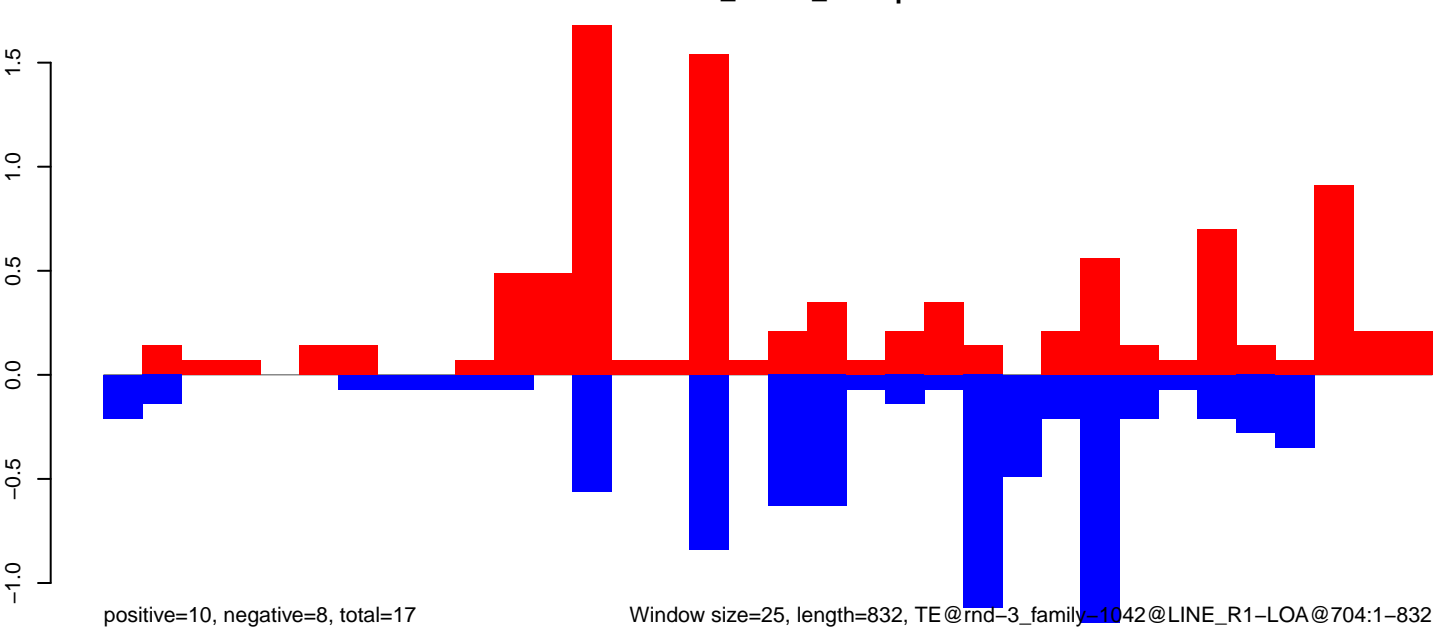
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=832, TE@rnd-3_family-1042@LINE_R1-LOA@704:1-832

CuQuin_Mcarc_NL.18_23.rep



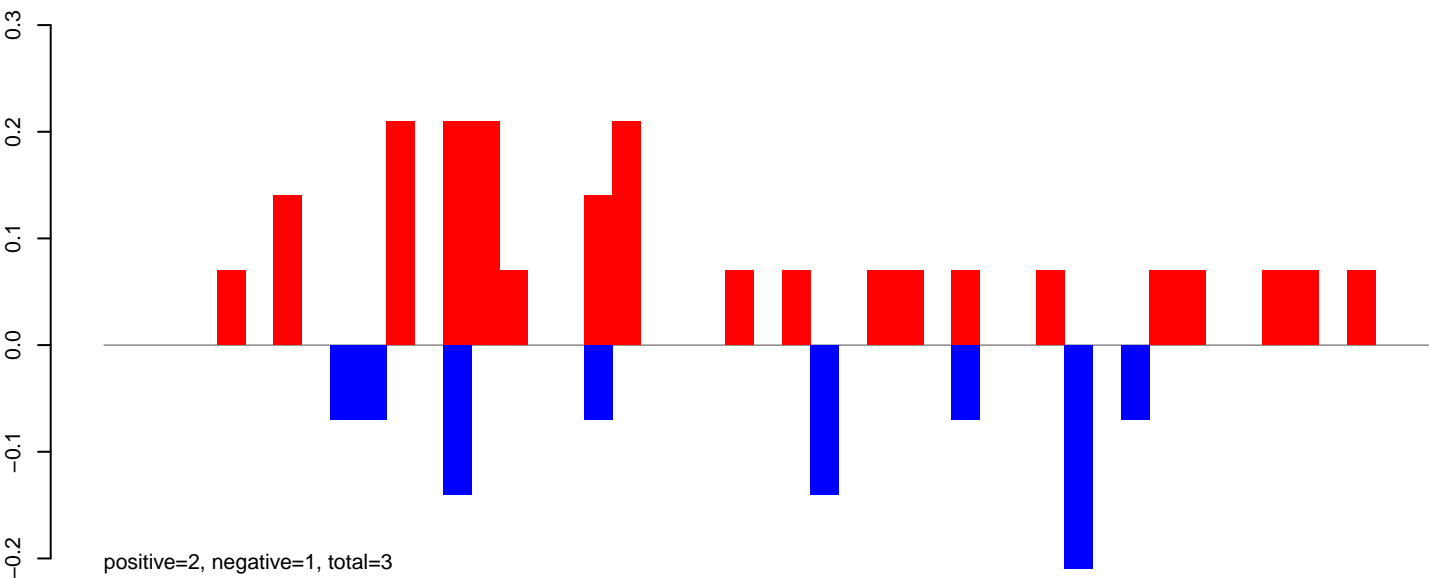
CuQuin_Mcarc_NL.24_35.rep



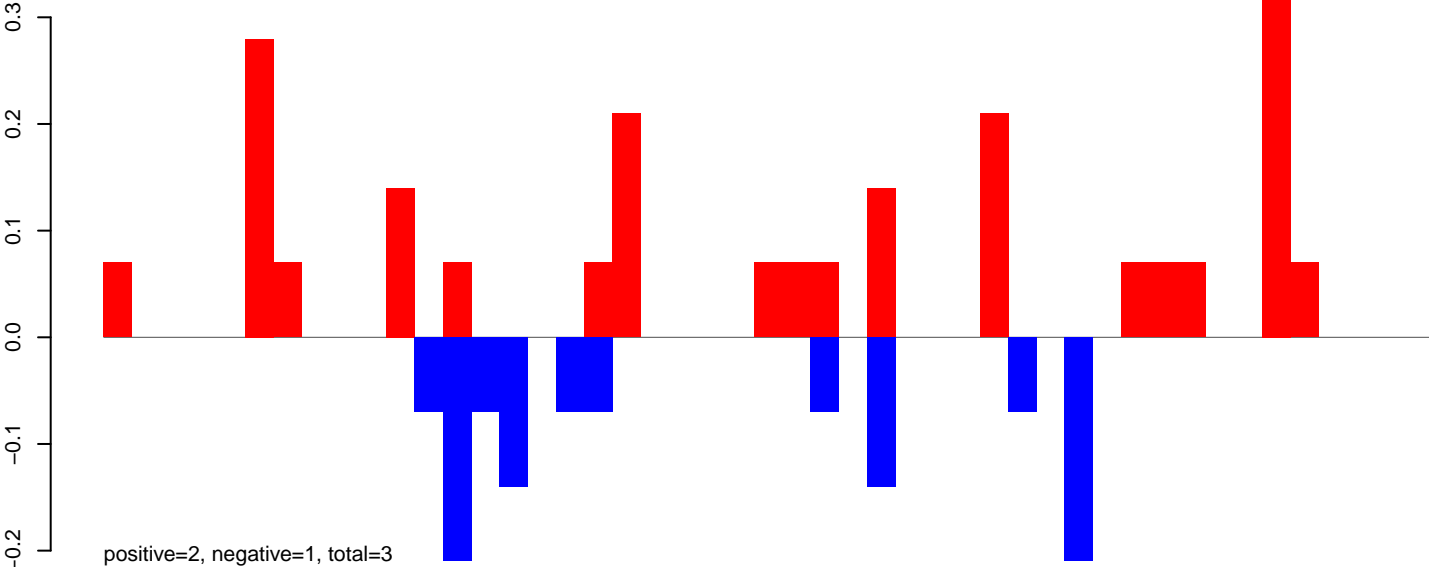
CuQuin_Mcarc_NL.rep



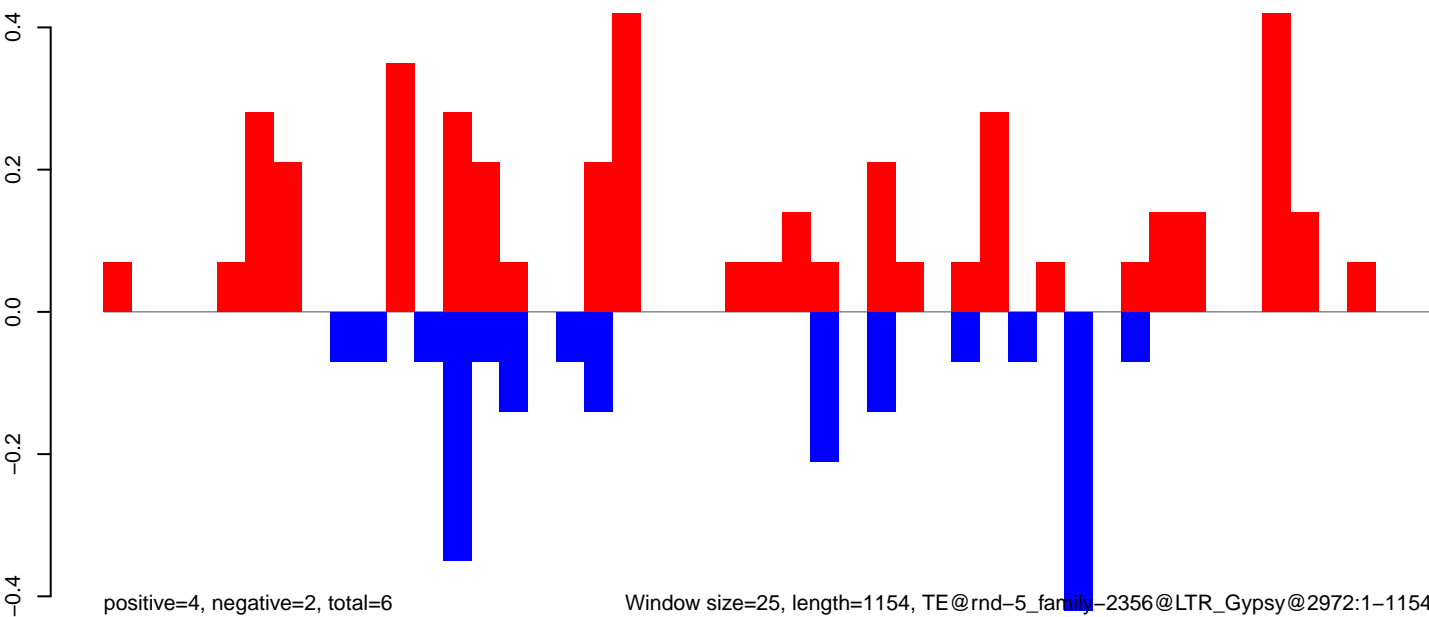
CuQuin_Mcarc_NL.18_23.rep



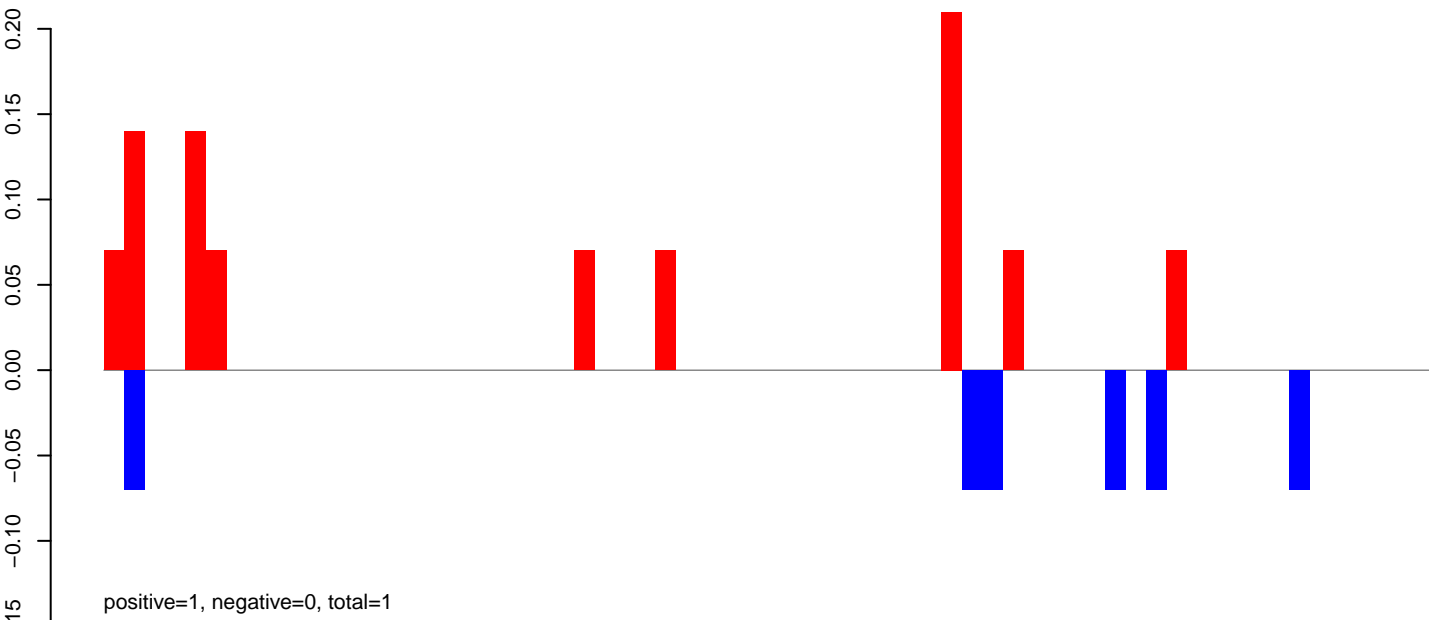
CuQuin_Mcarc_NL.24_35.rep



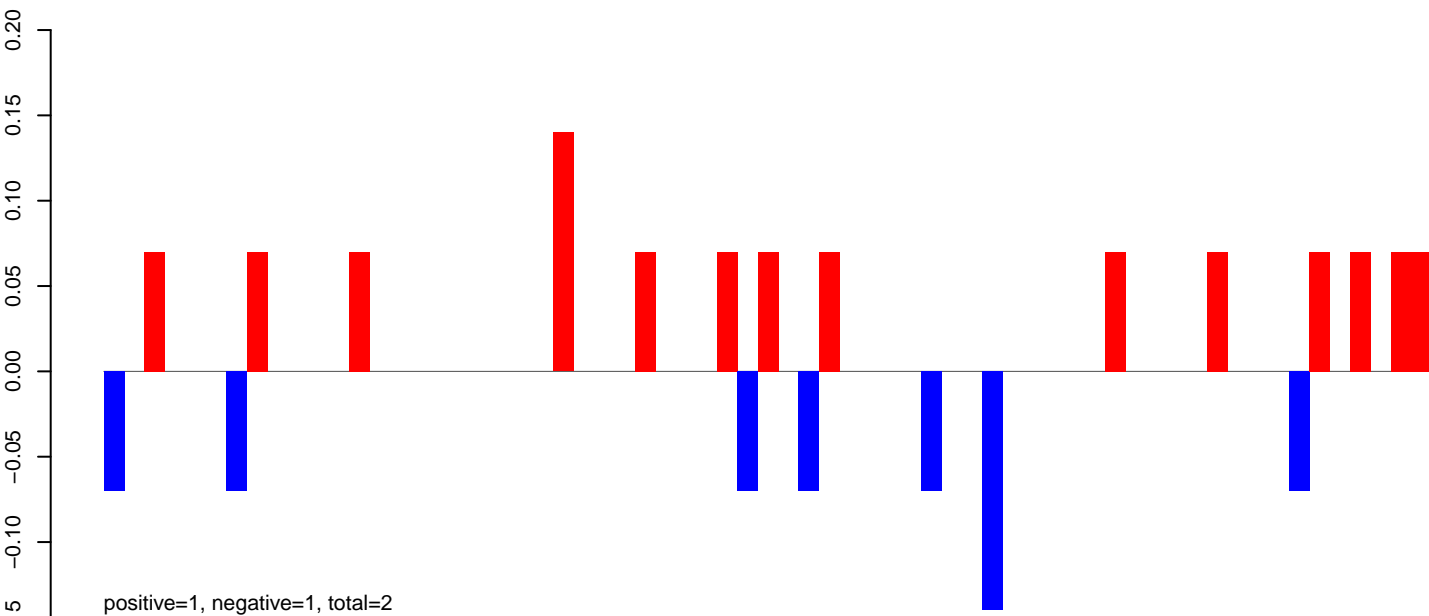
CuQuin_Mcarc_NL.rep



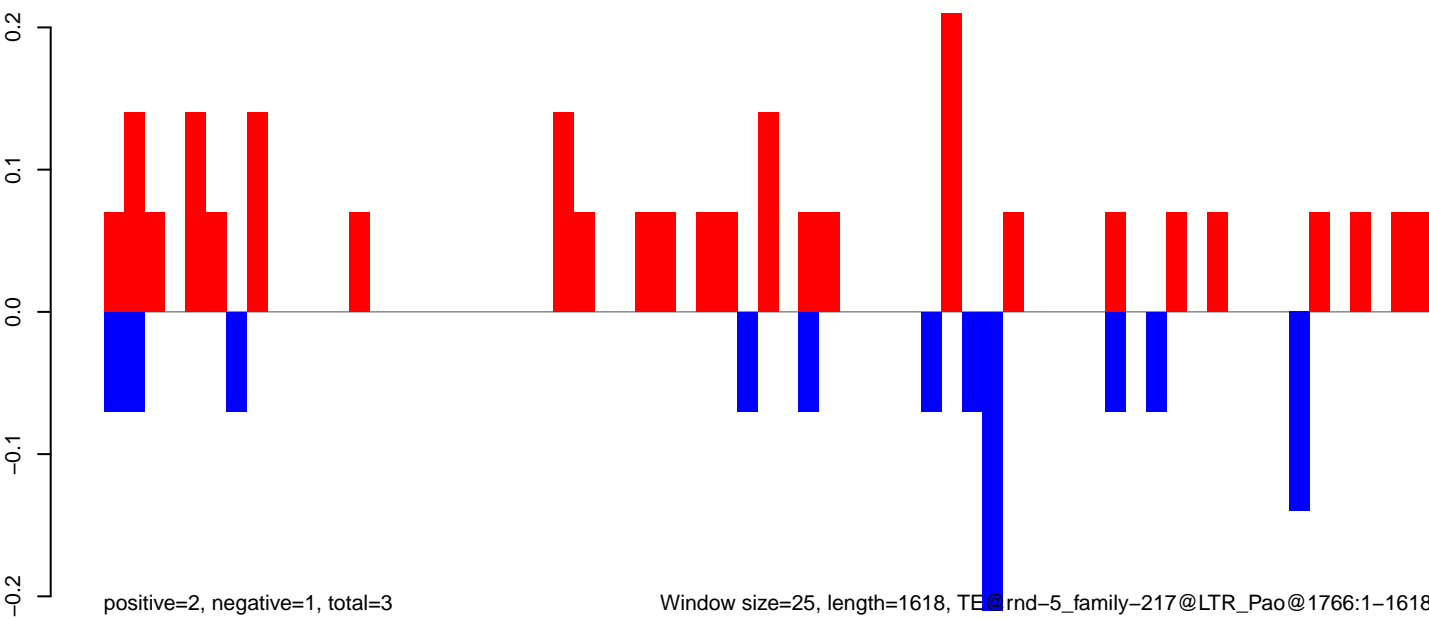
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

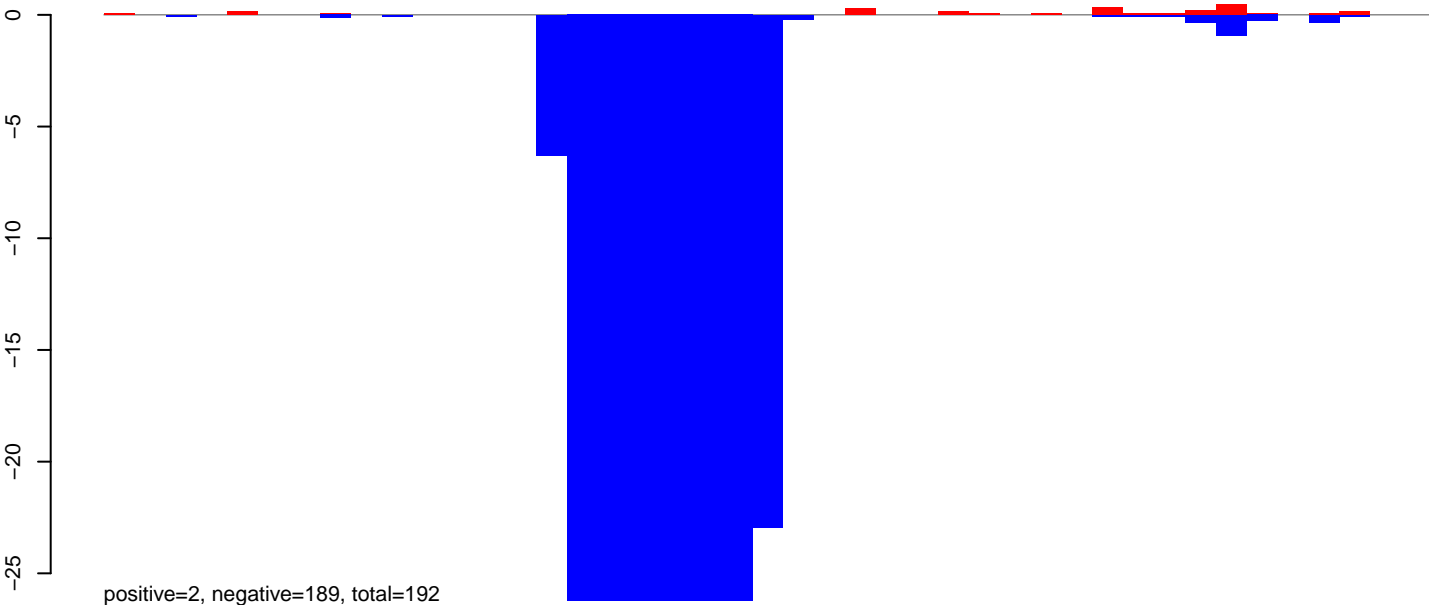


CuQuin_Mcarc_NL.rep

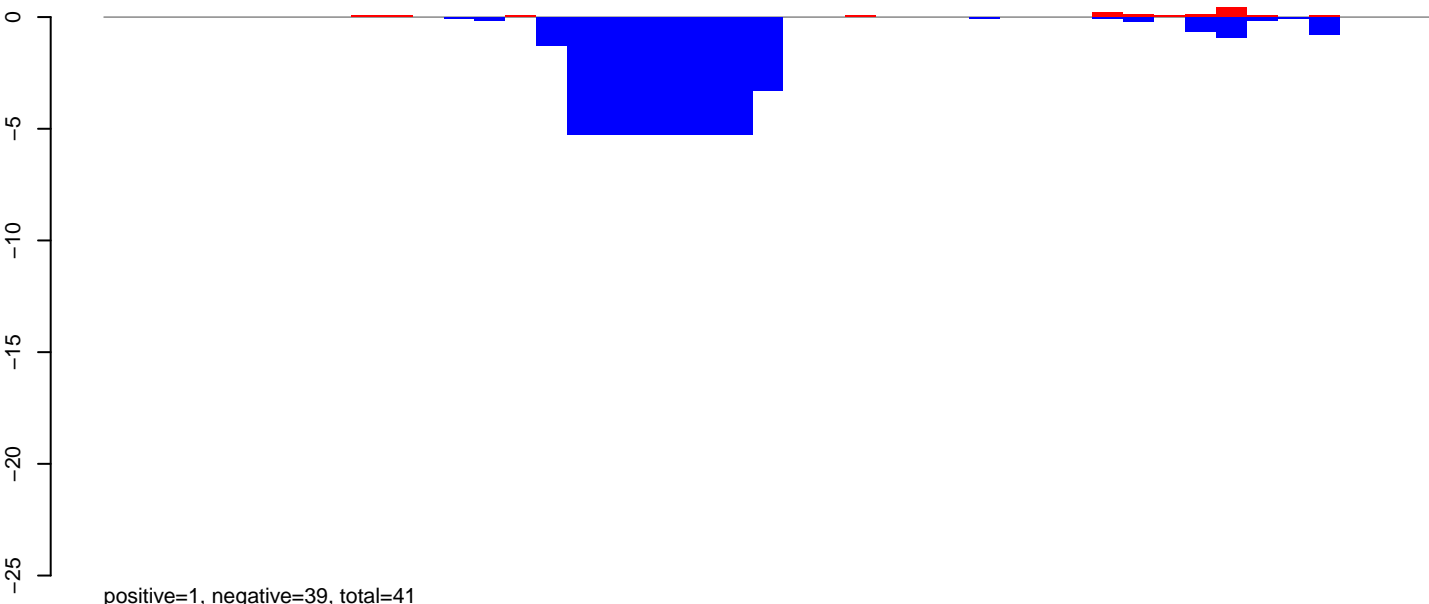


Window size=25, length=1618, TE@rnd-5_family-217@LTR_Pao@1766:1-1618

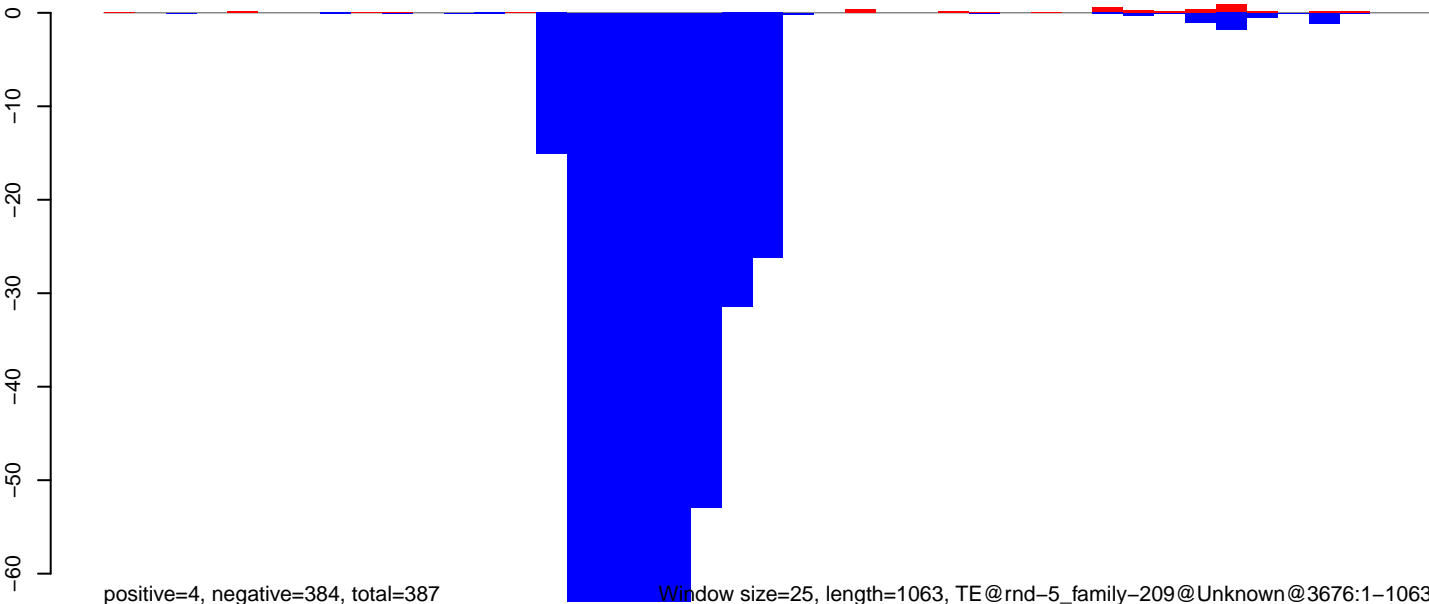
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

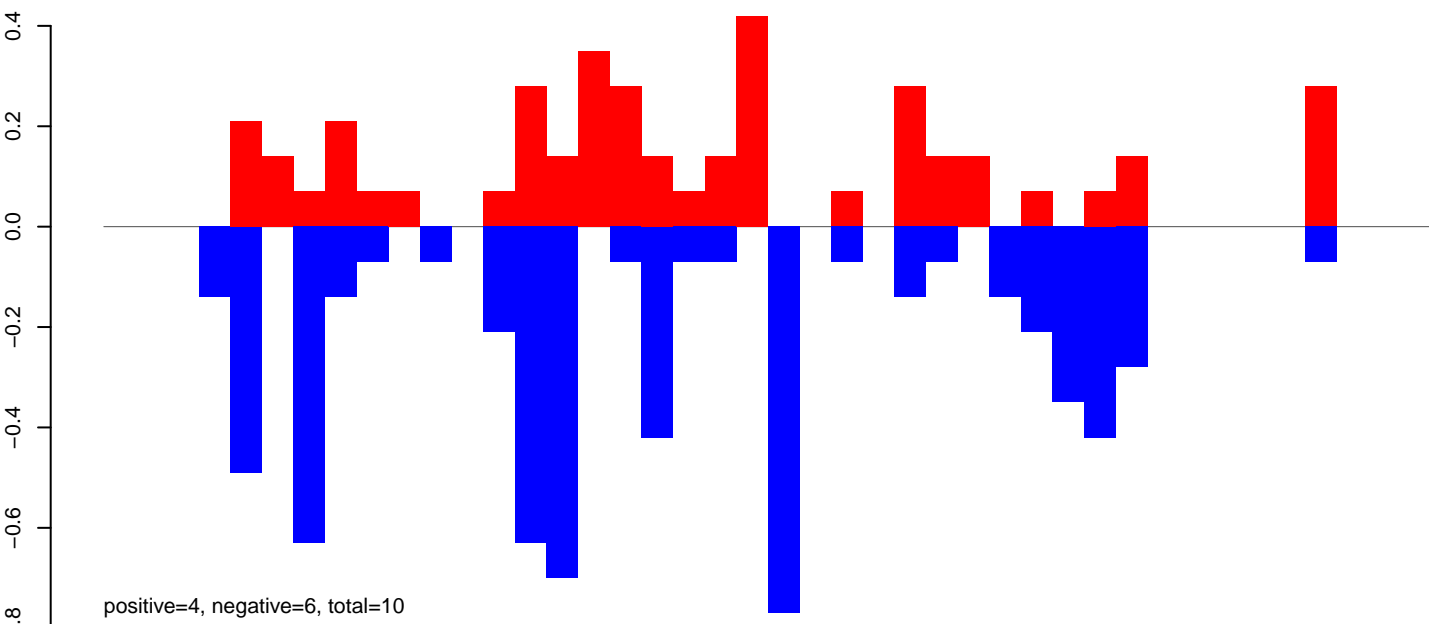


CuQuin_Mcarc_NL.rep

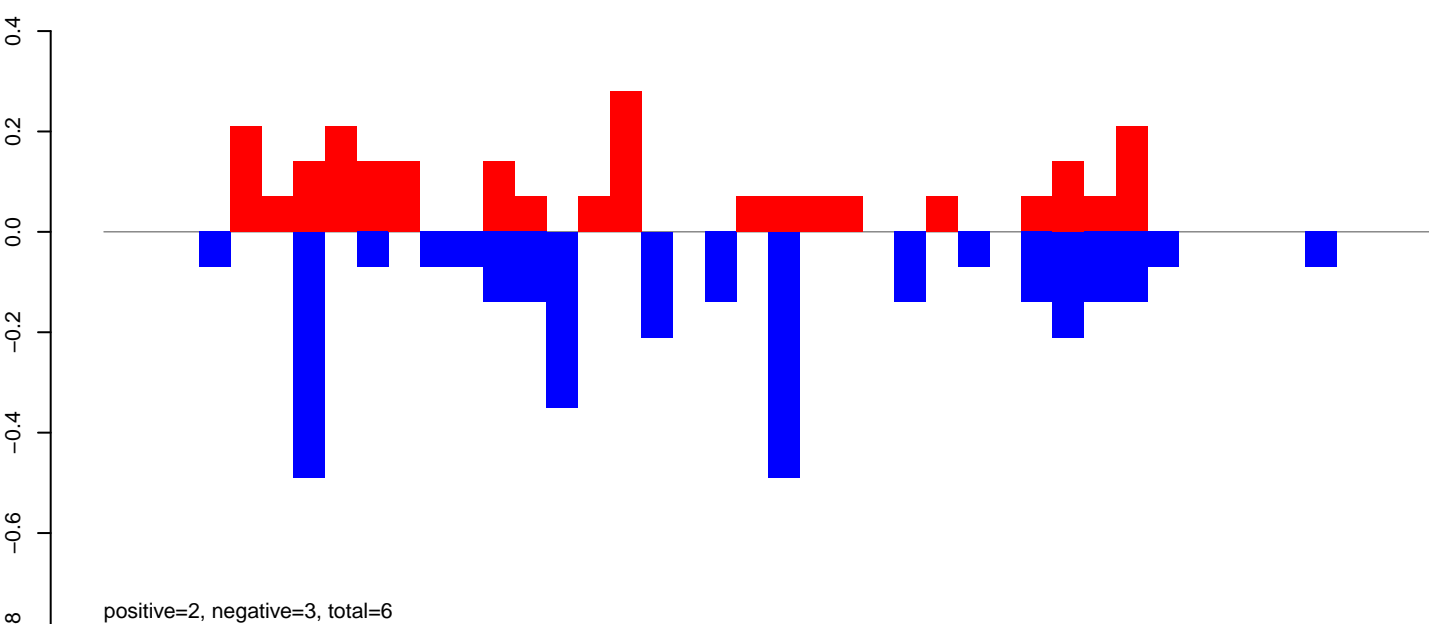


Window size=25, length=1063, TE@rnd-5_family-209@Unknown@3676:1-1063

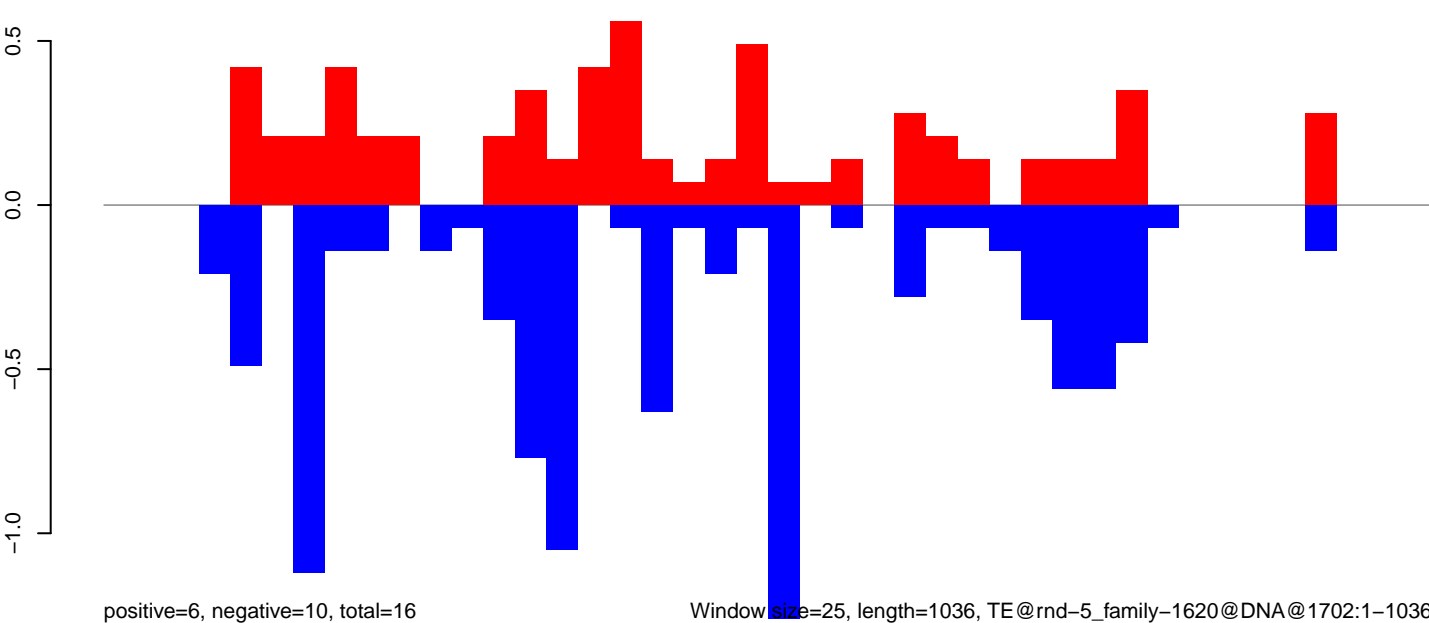
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

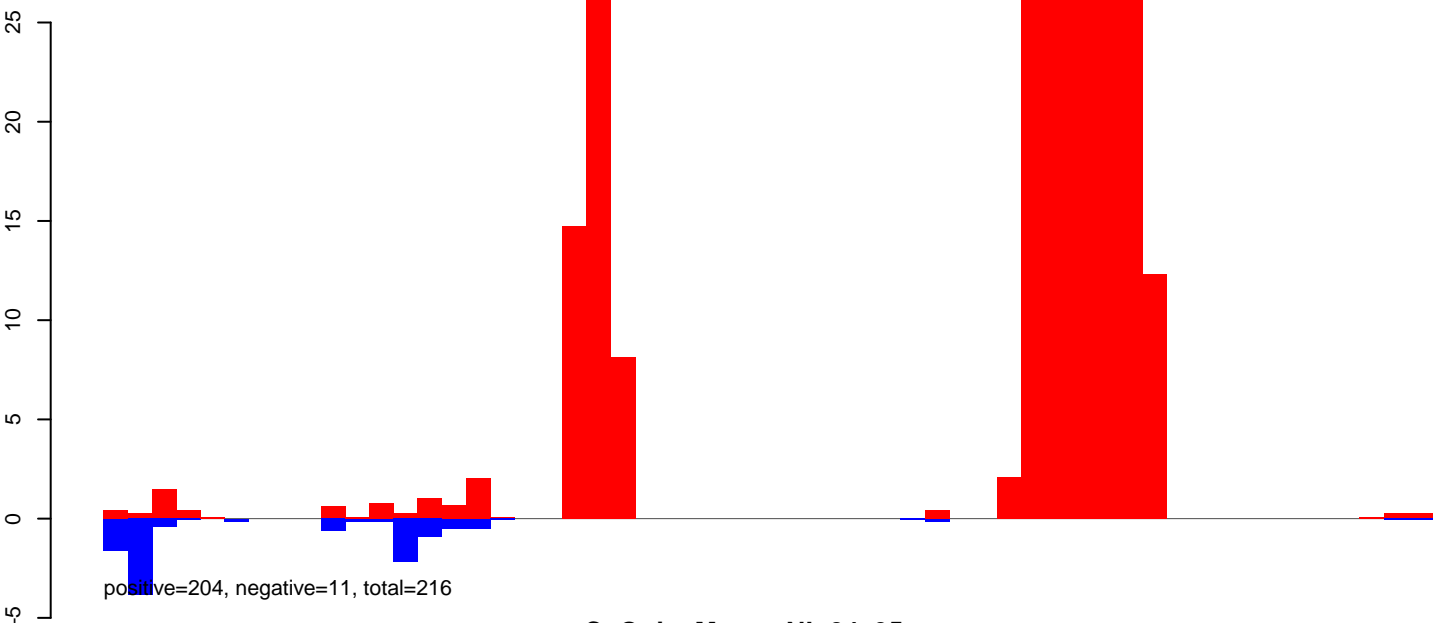


CuQuin_Mcarc_NL.rep

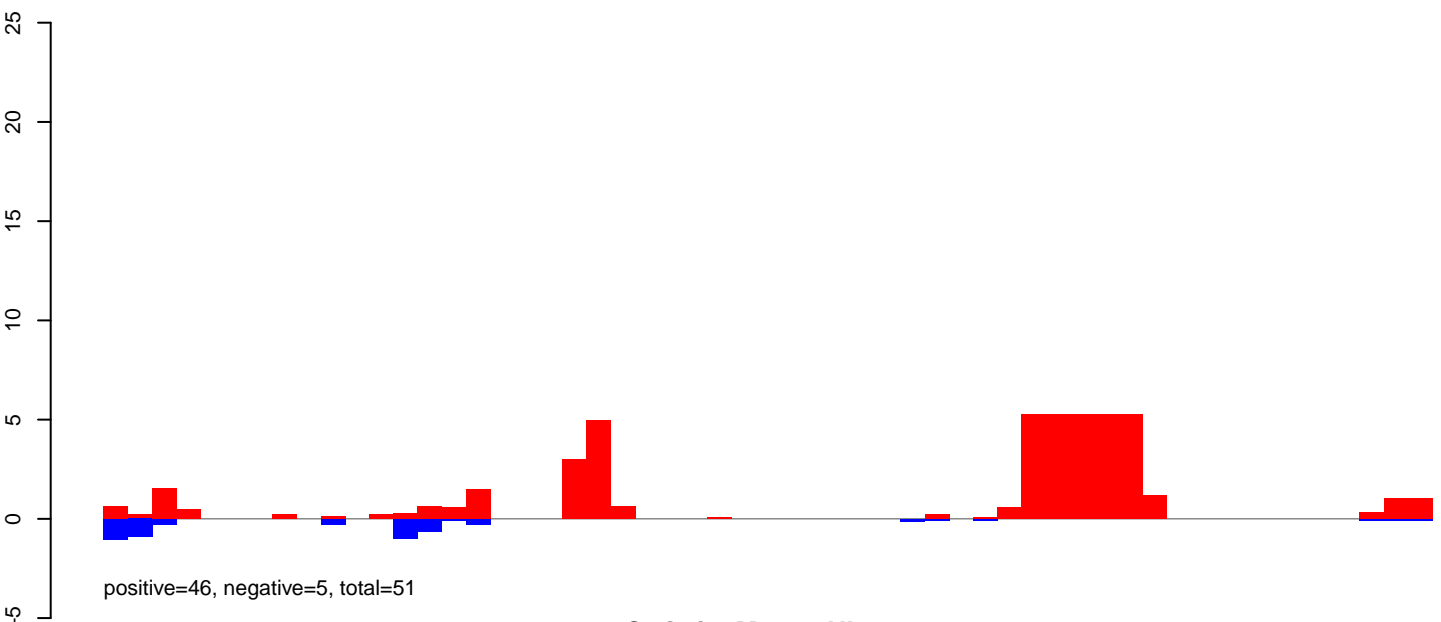


Window size=25, length=1036, TE@rnd-5_family-1620@DNA@1702:1-1036

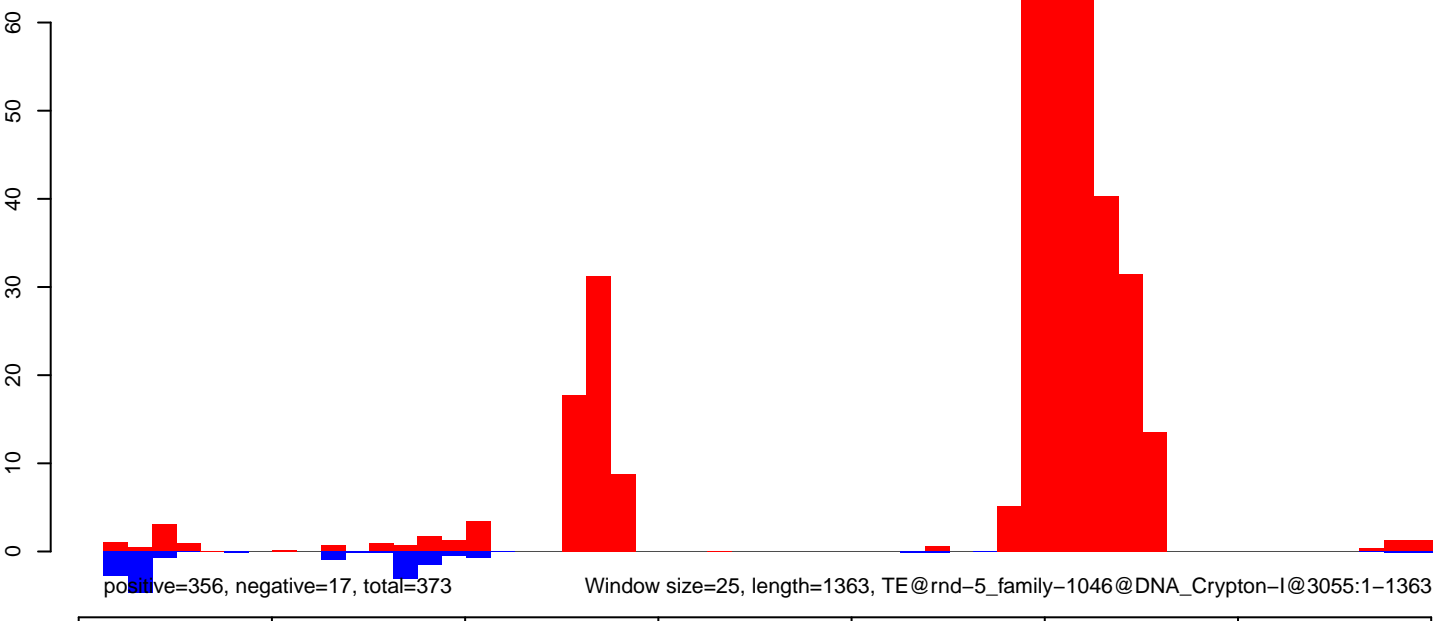
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

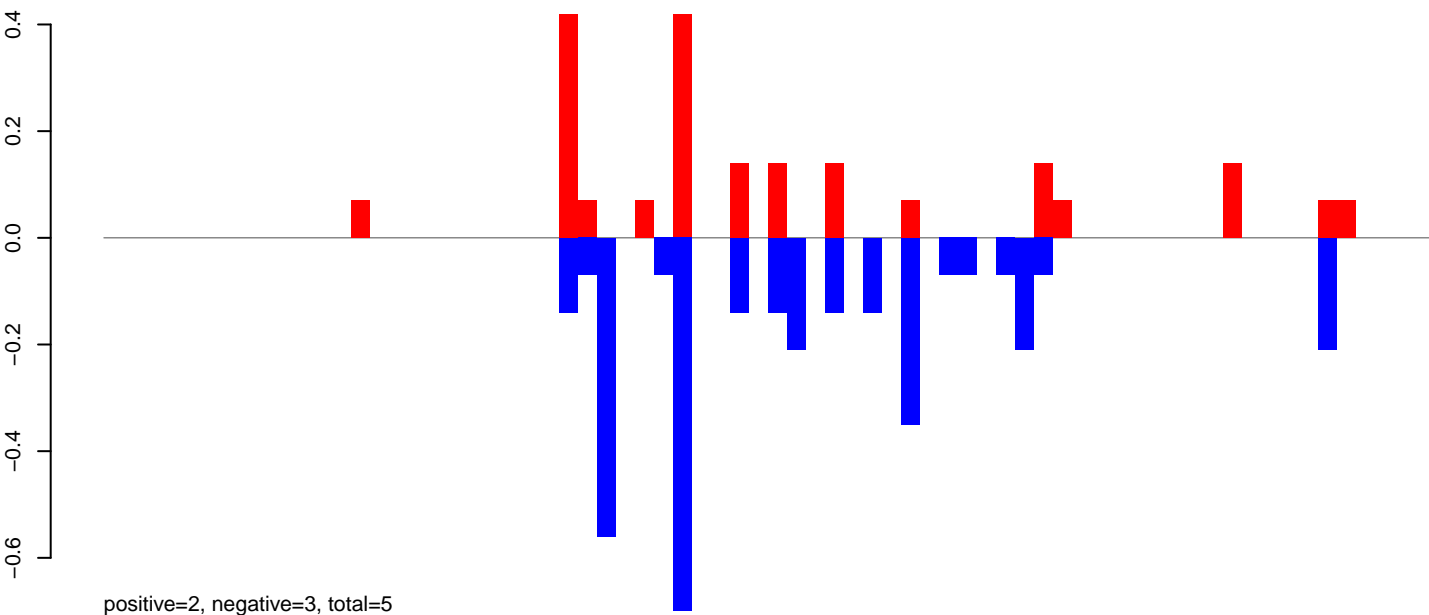


CuQuin_Mcarc_NL.rep

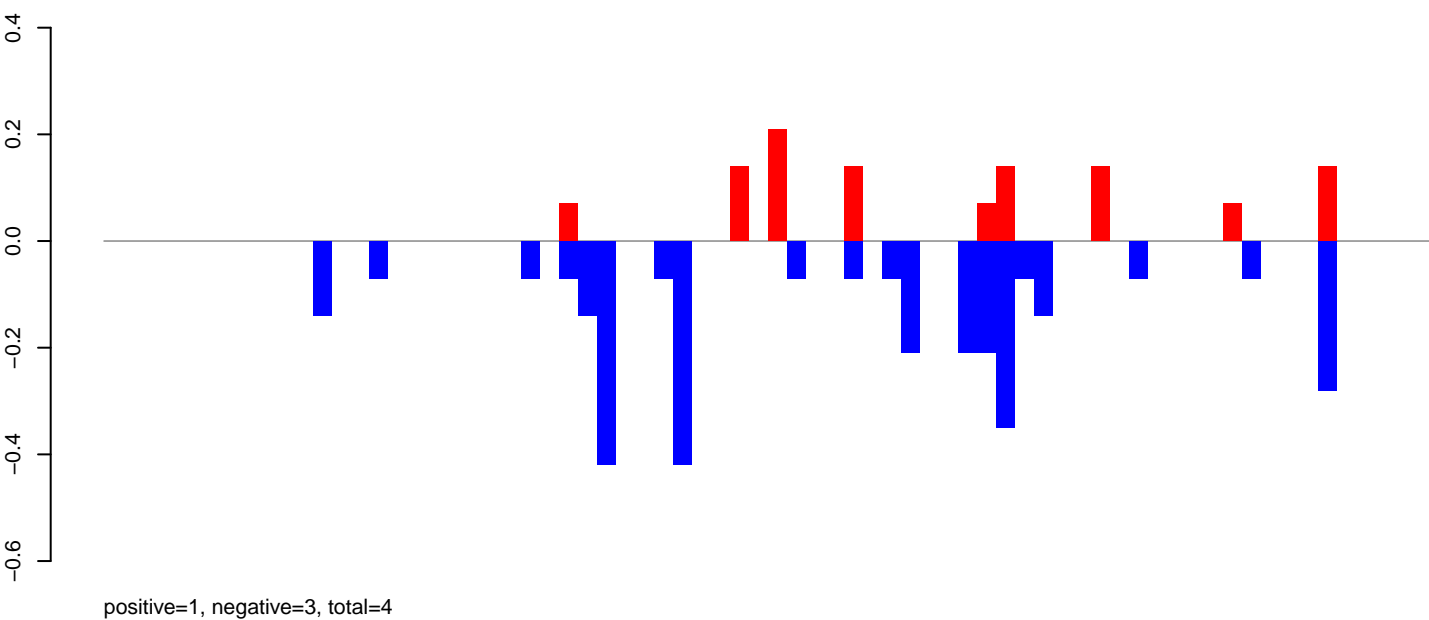


Window size=25, length=1363, TE@rnd-5_family-1046@DNA_Crypton-I@3055:1-1363

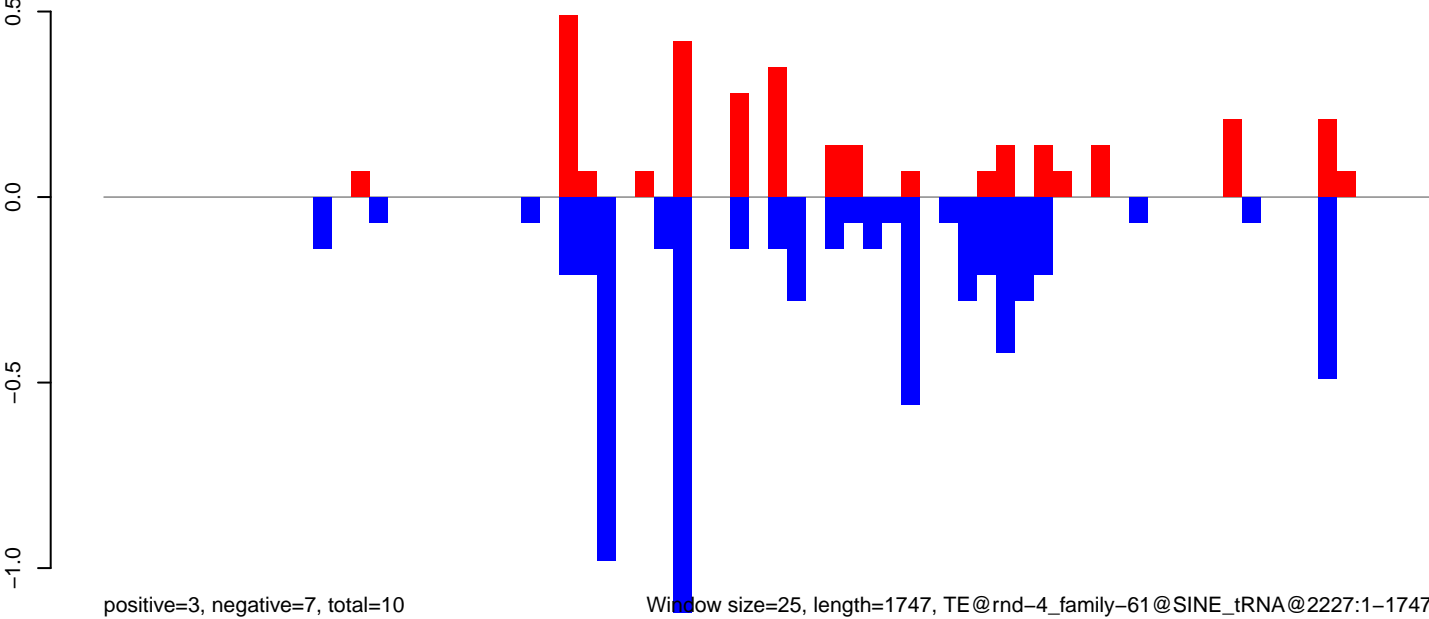
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

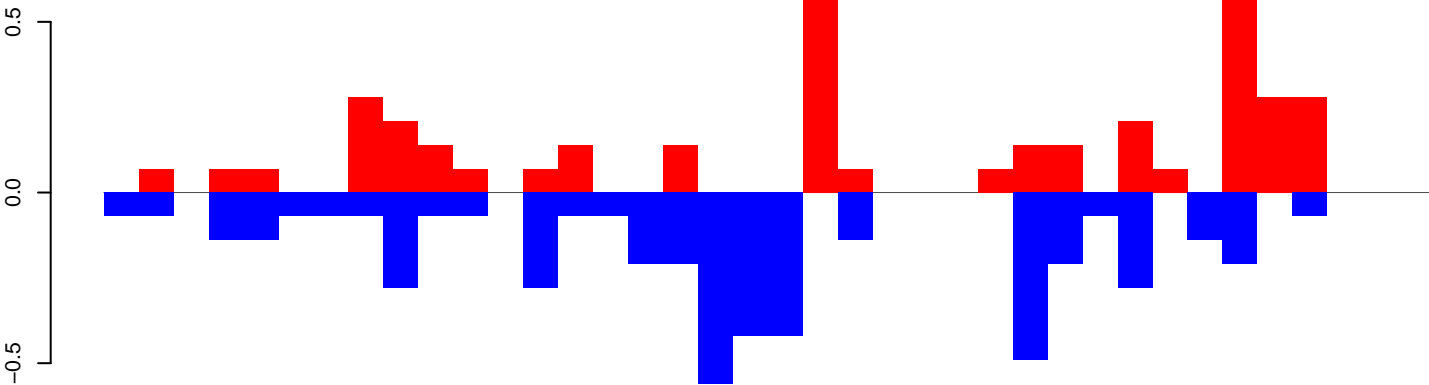


CuQuin_Mcarc_NL.rep



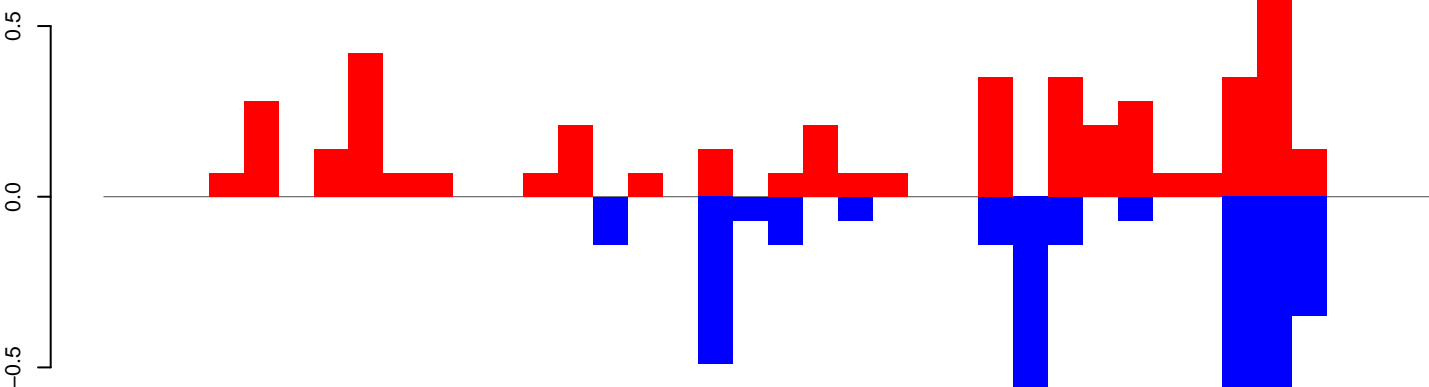
Window size=25, length=1747, TE@rnd-4_family-61@SINE_tRNA@2227:1-1747

CuQuin_Mcarc_NL.18_23.rep



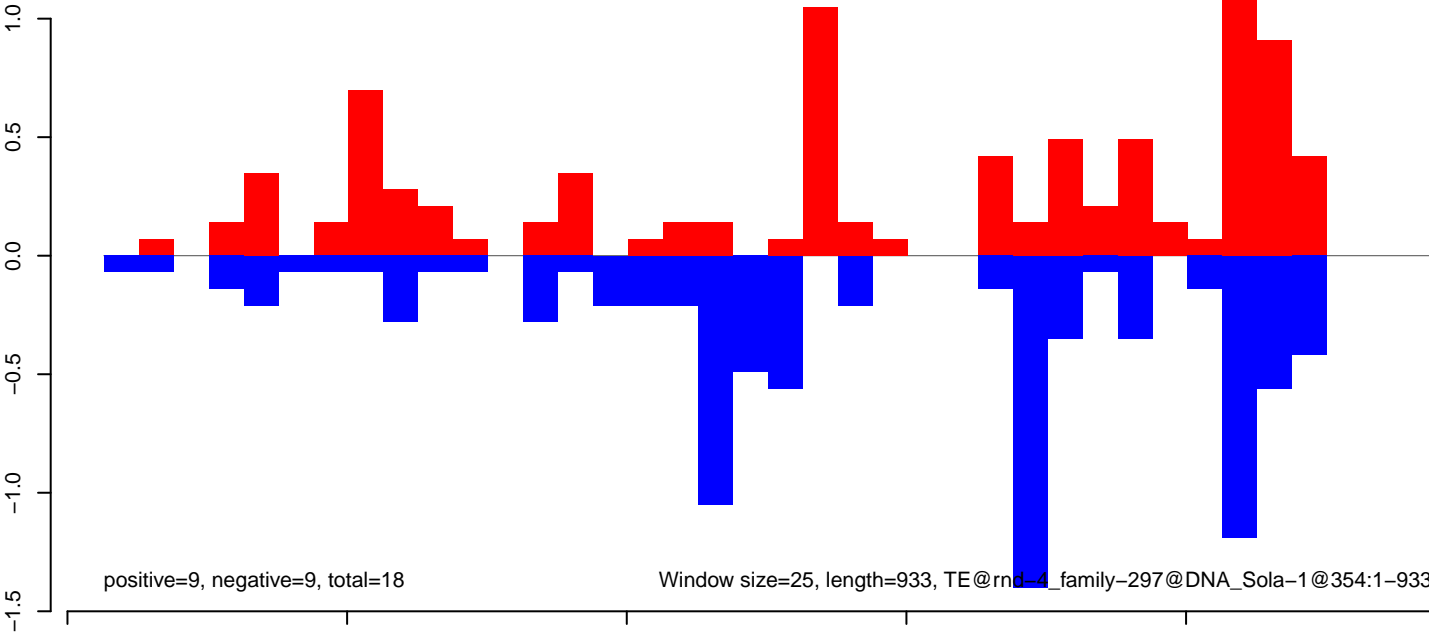
positive=4, negative=5, total=9

CuQuin_Mcarc_NL.24_35.rep



positive=4, negative=4, total=8

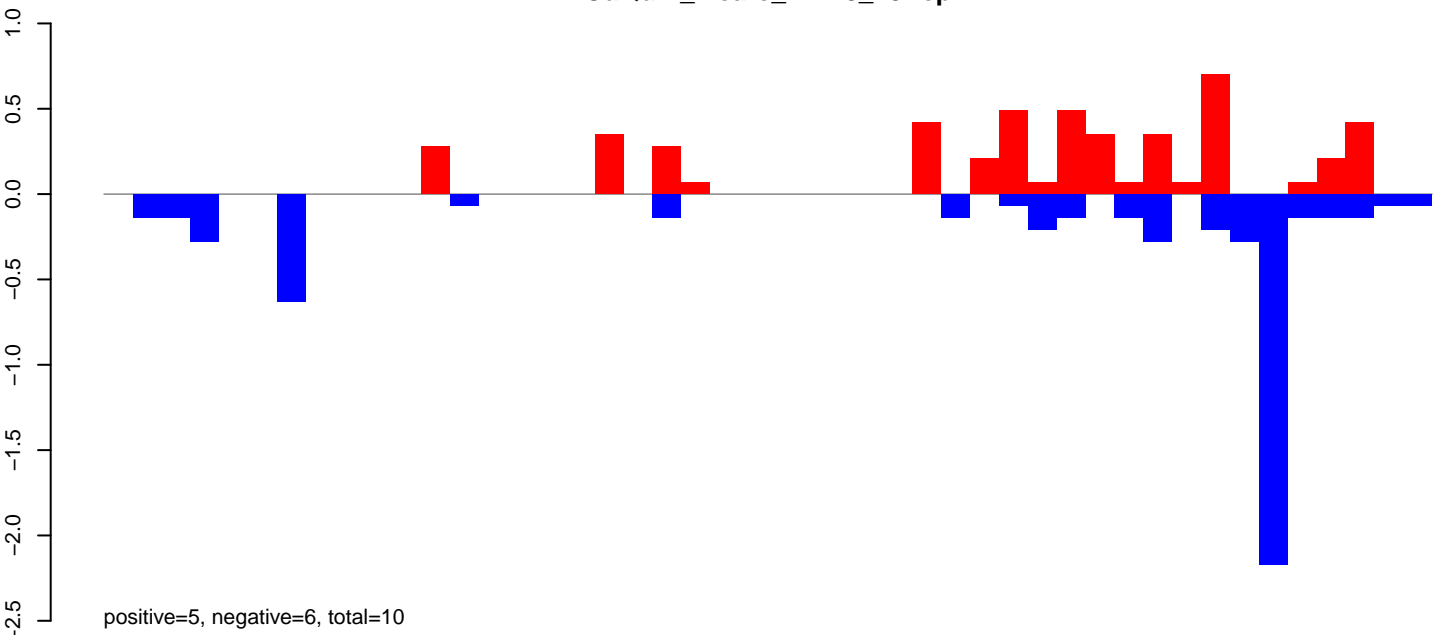
CuQuin_Mcarc_NL.rep



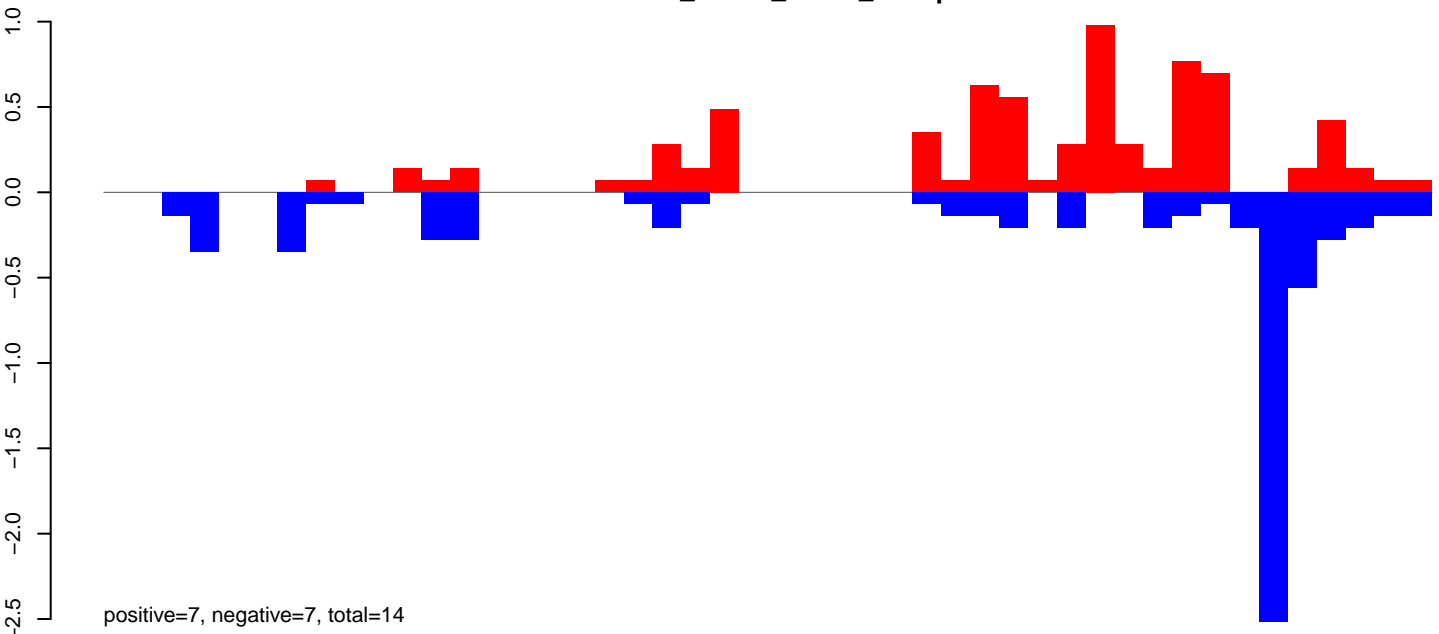
positive=9, negative=9, total=18

Window size=25, length=933, TE@rnd-4_family-297@DNA_Sola-1@354:1-933

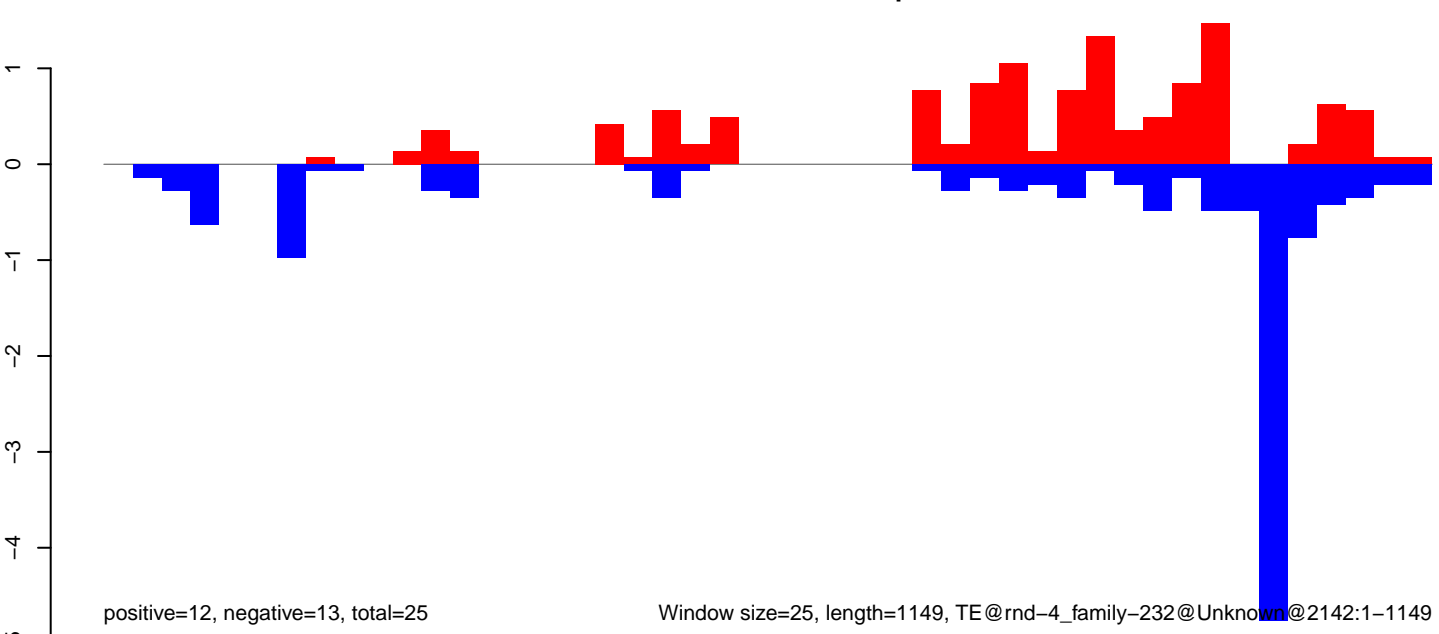
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

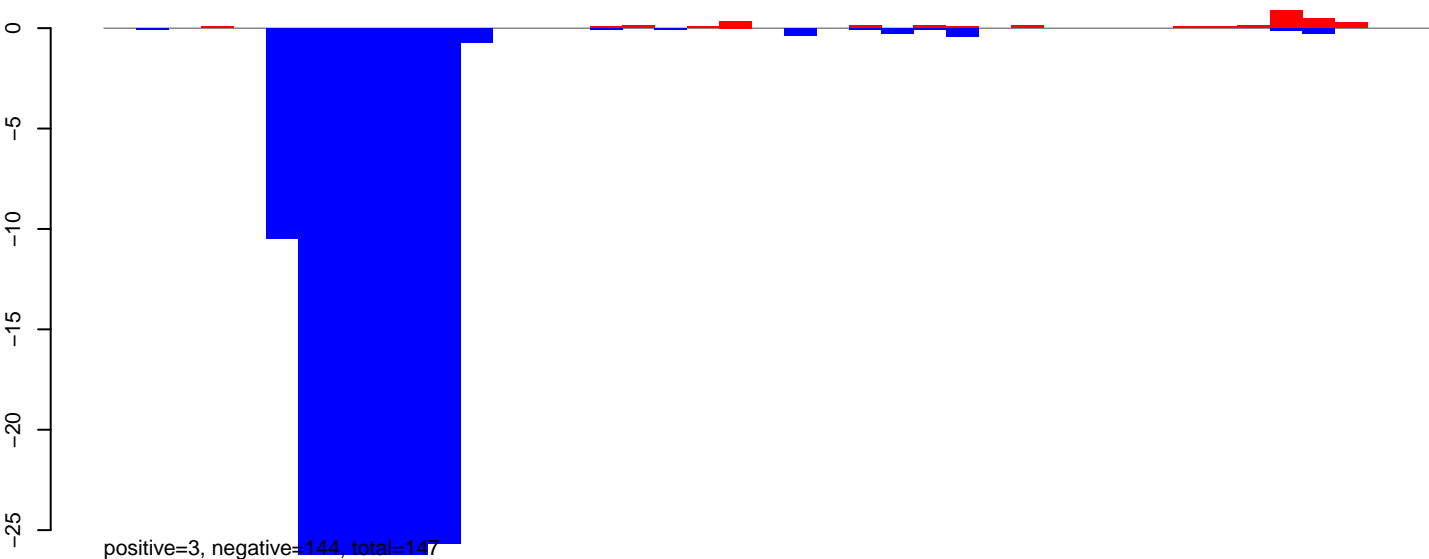


CuQuin_Mcarc_NL.rep

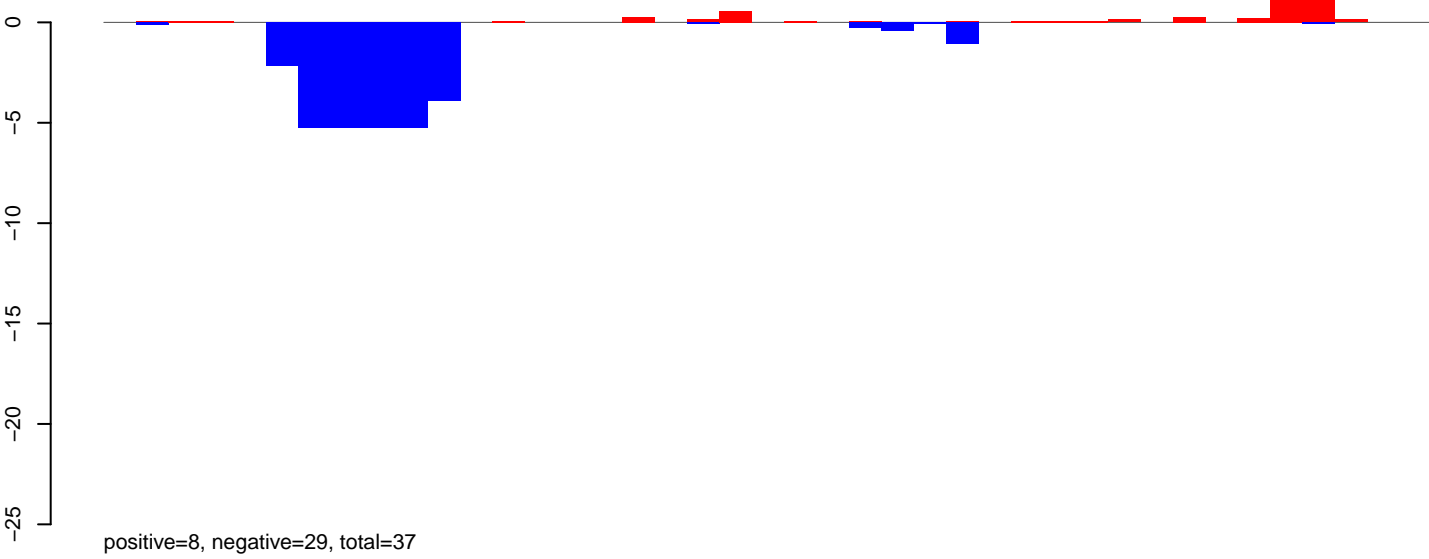


Window size=25, length=1149, TE@rnd-4_family-232@Unknown@2142:1-1149

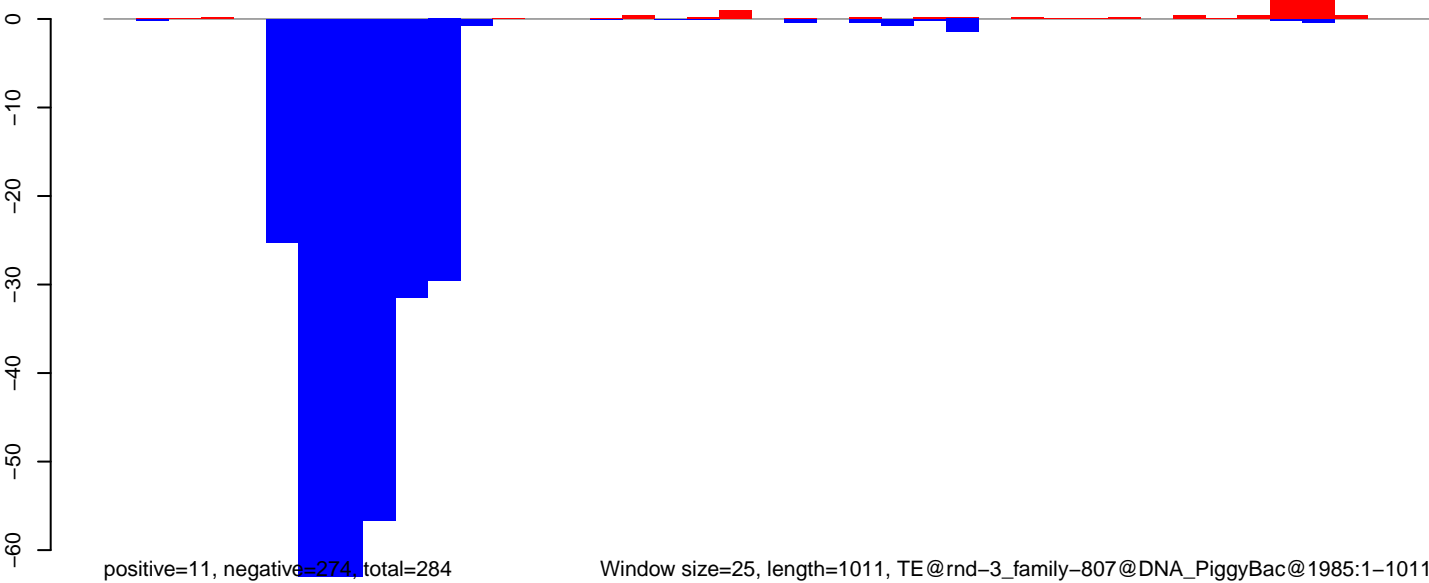
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



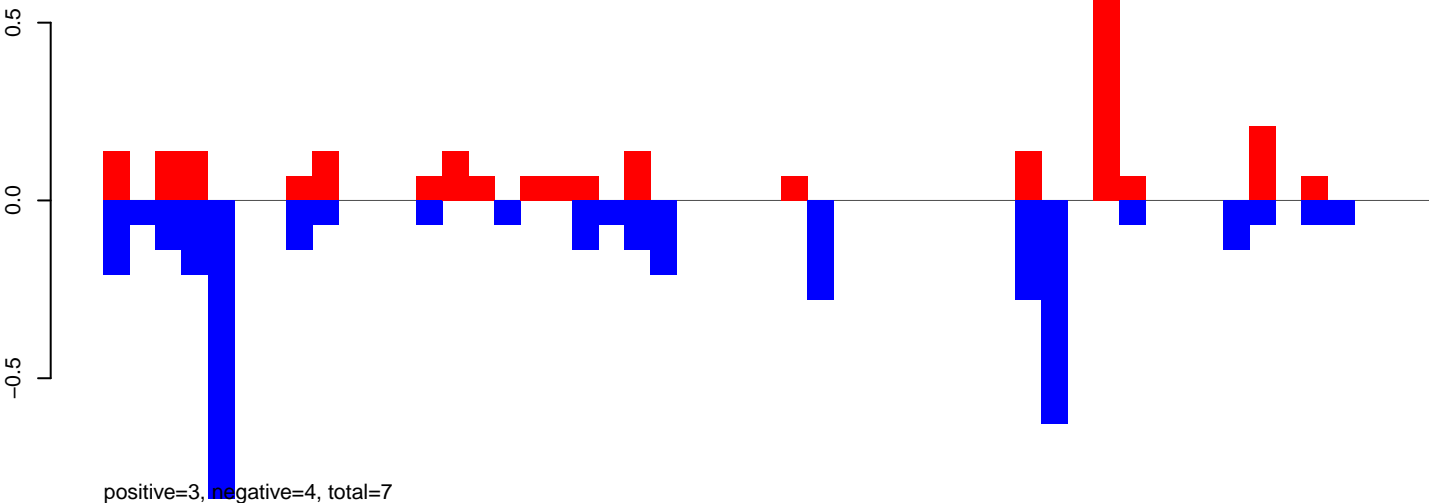
CuQuin_Mcarc_NL.rep



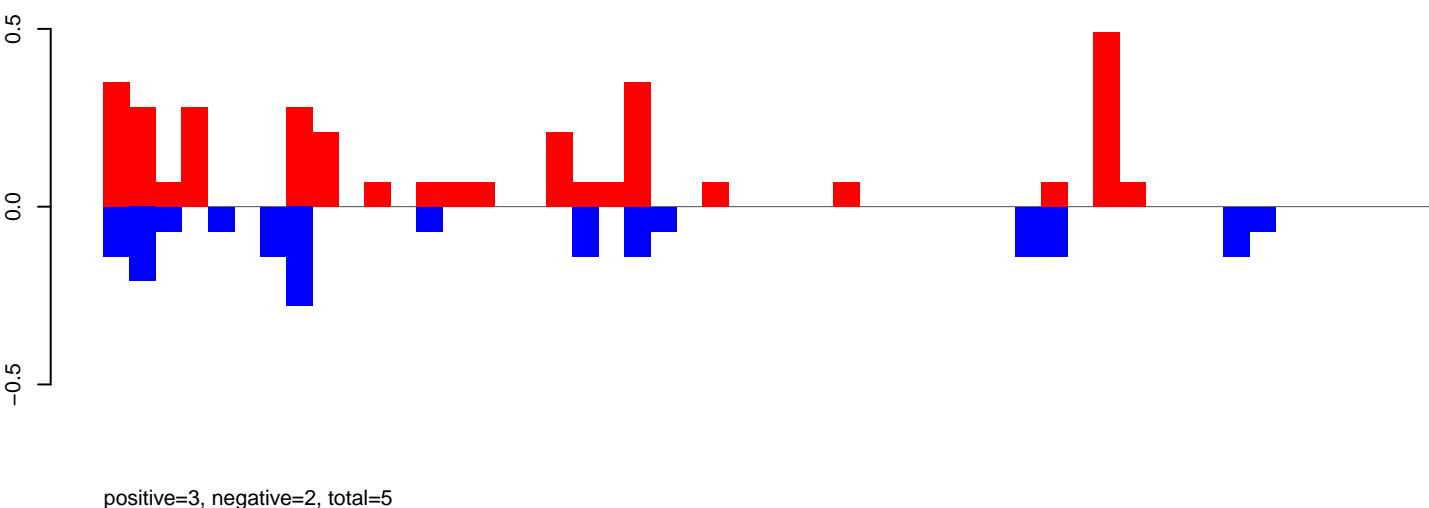
Window size=25, length=1011, TE@rnd-3_family-807@DNA_PiggyBac@1985:1-1011

0 200 400 600 800 1000

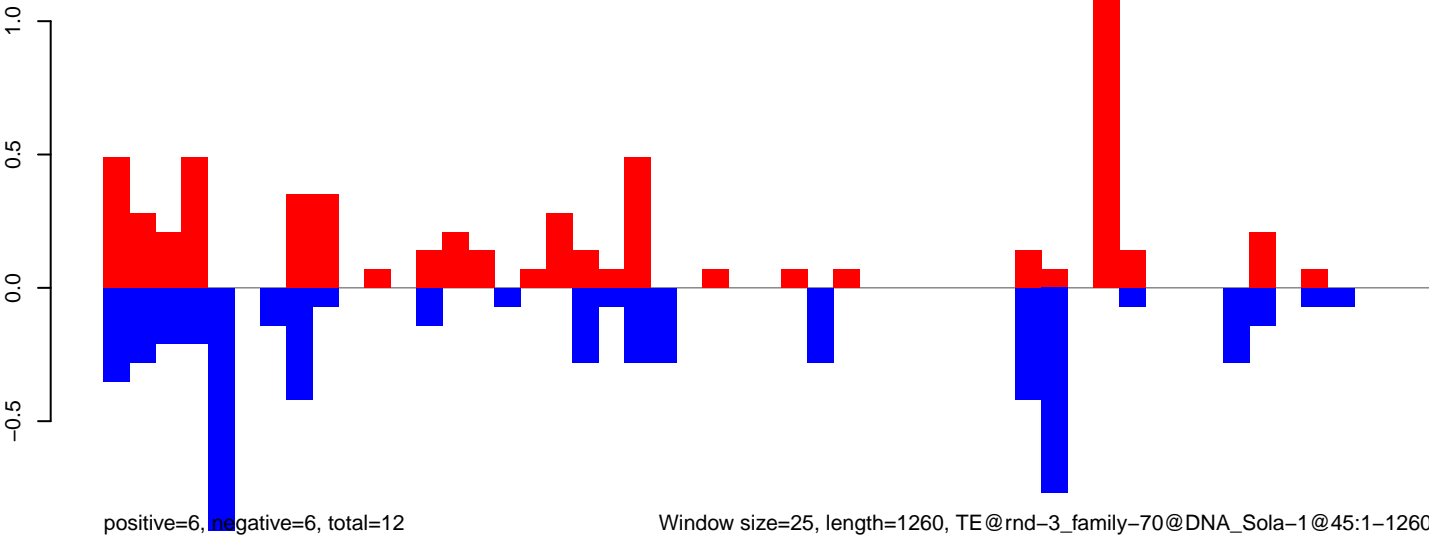
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

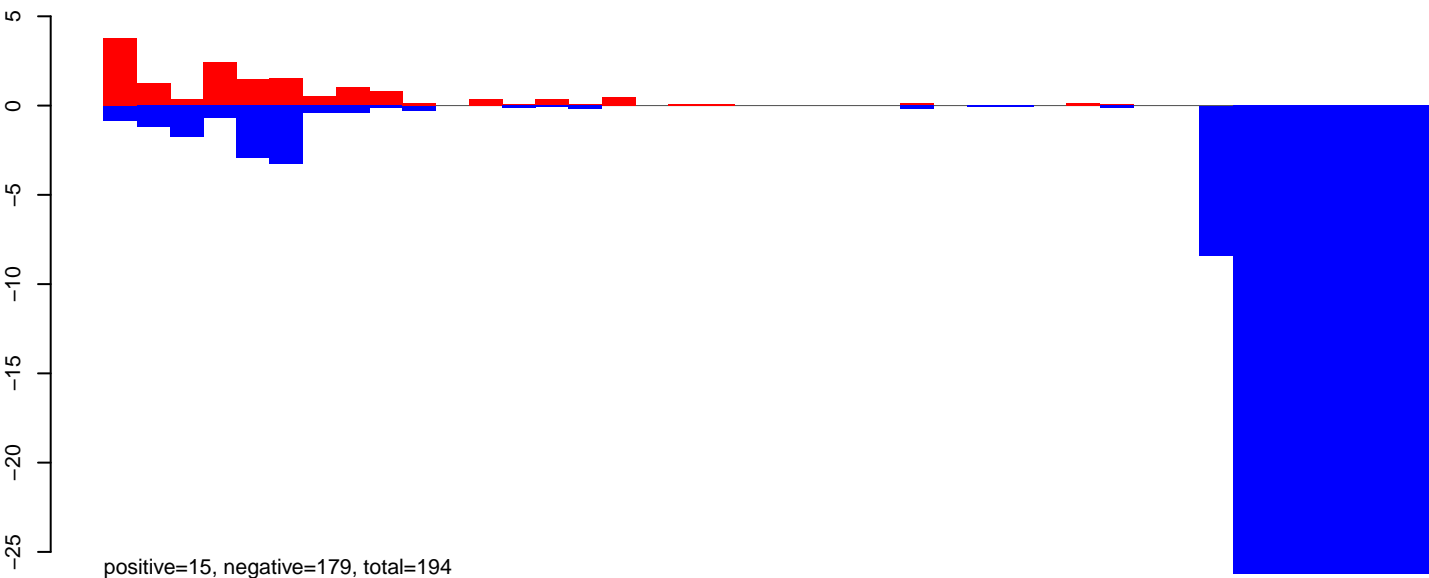


CuQuin_Mcarc_NL.rep

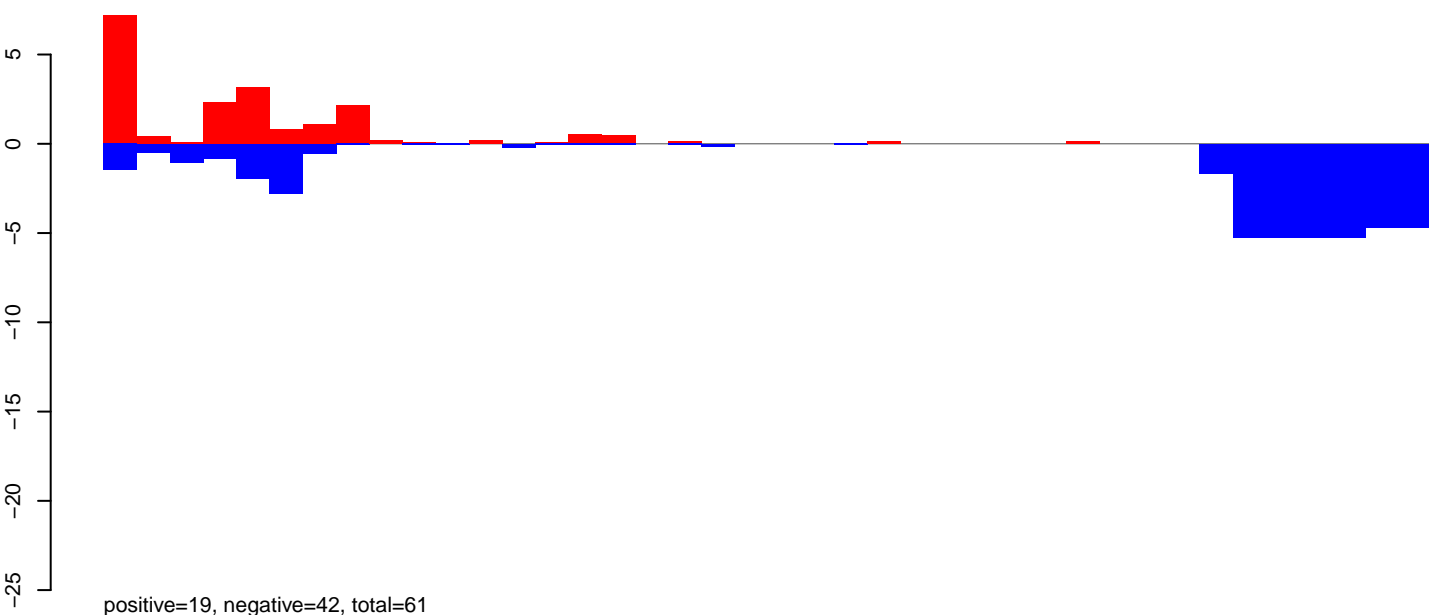


Window size=25, length=1260, TE@rnd-3_family-70@DNA_Sola-1@45:1-1260

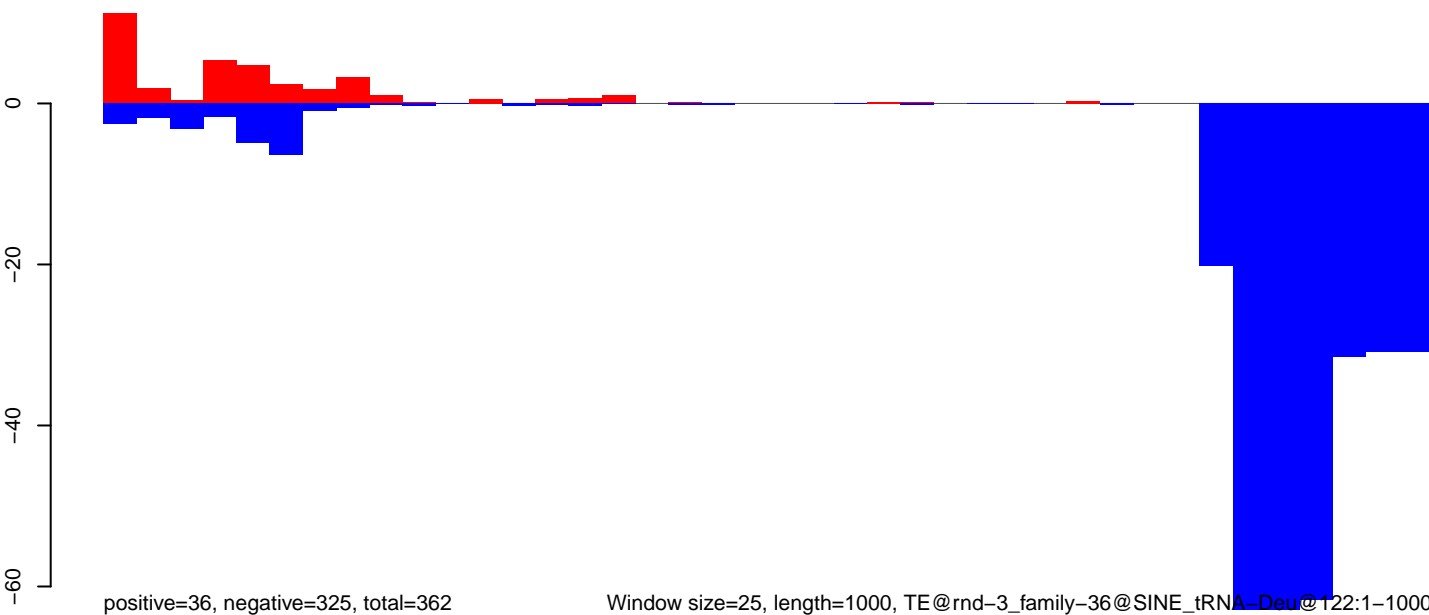
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



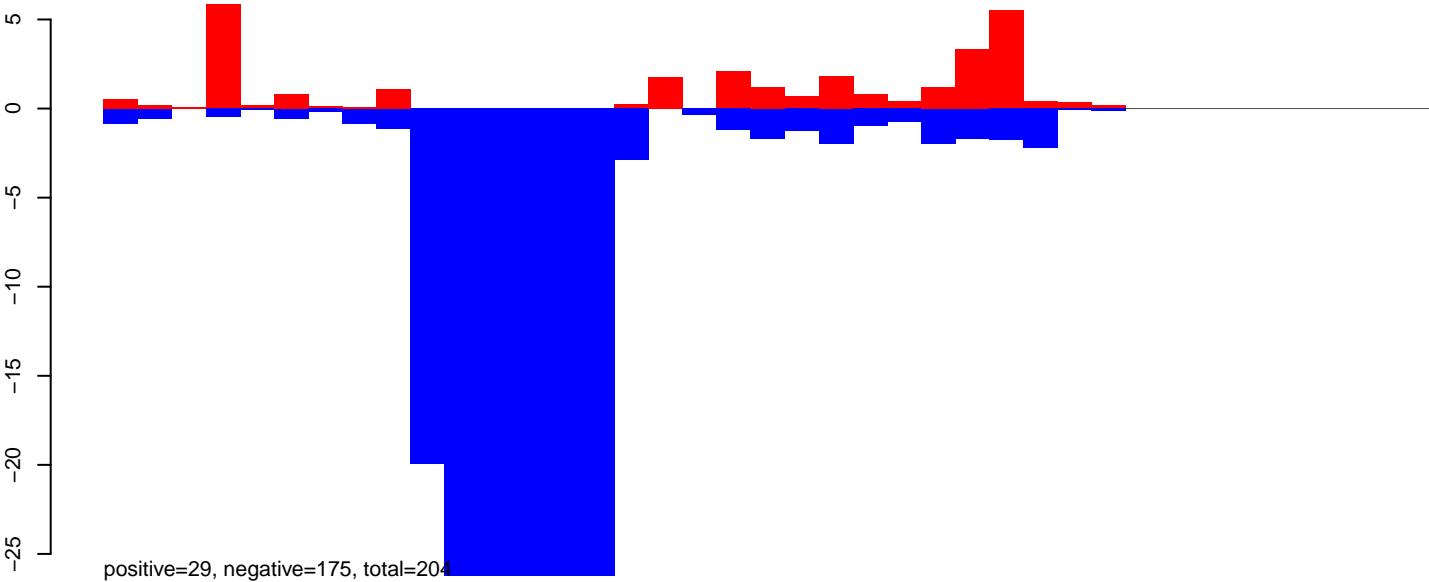
CuQuin_Mcarc_NL.rep



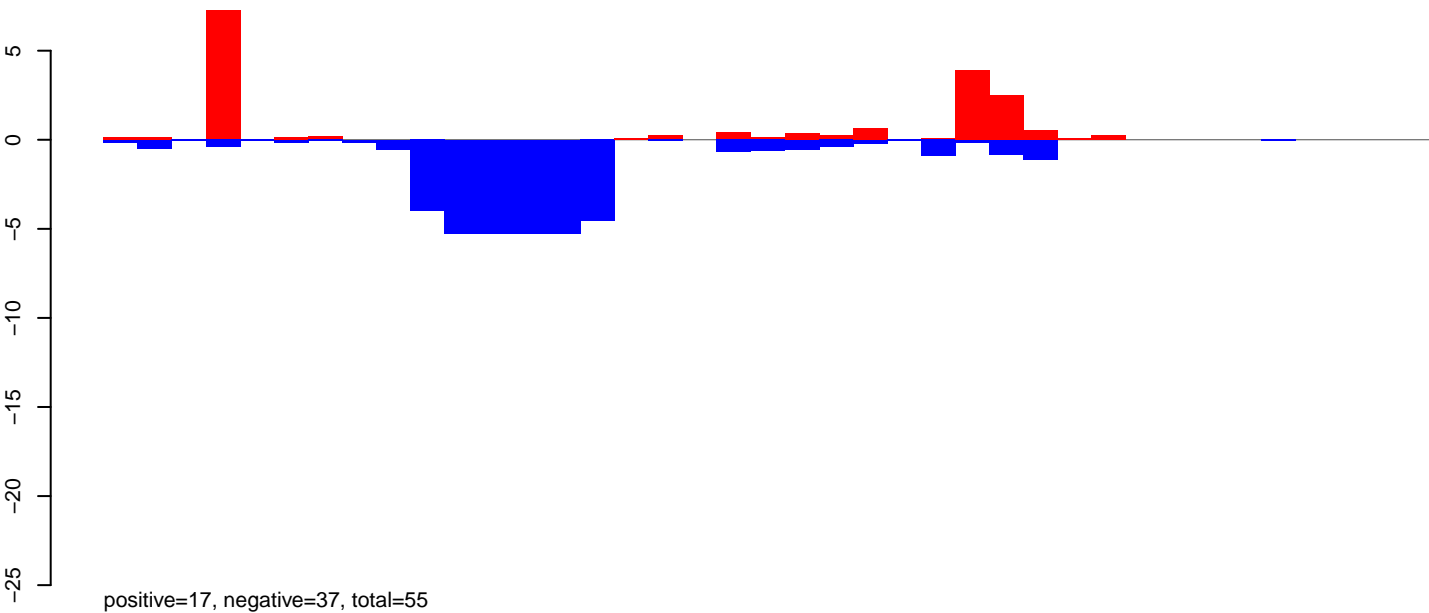
Window size=25, length=1000, TE@rnd-3_family-36@SINE_tRNA-Deu@122:1-1000

0 200 400 600 800 1000

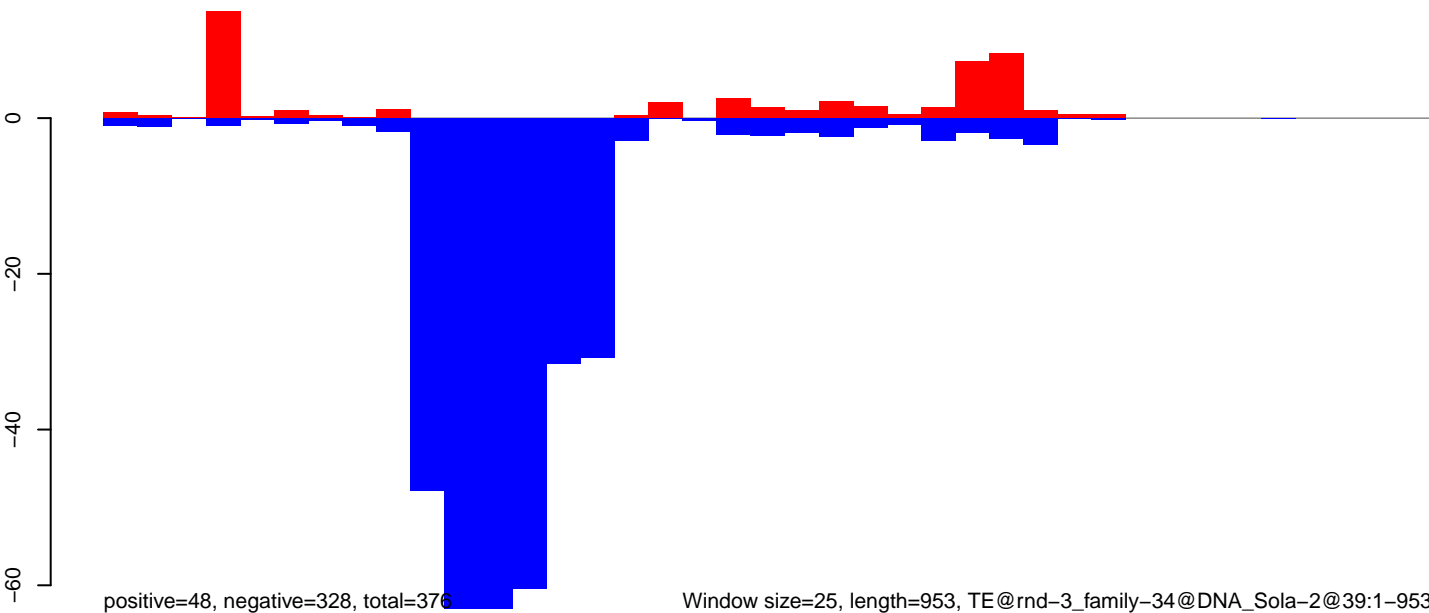
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



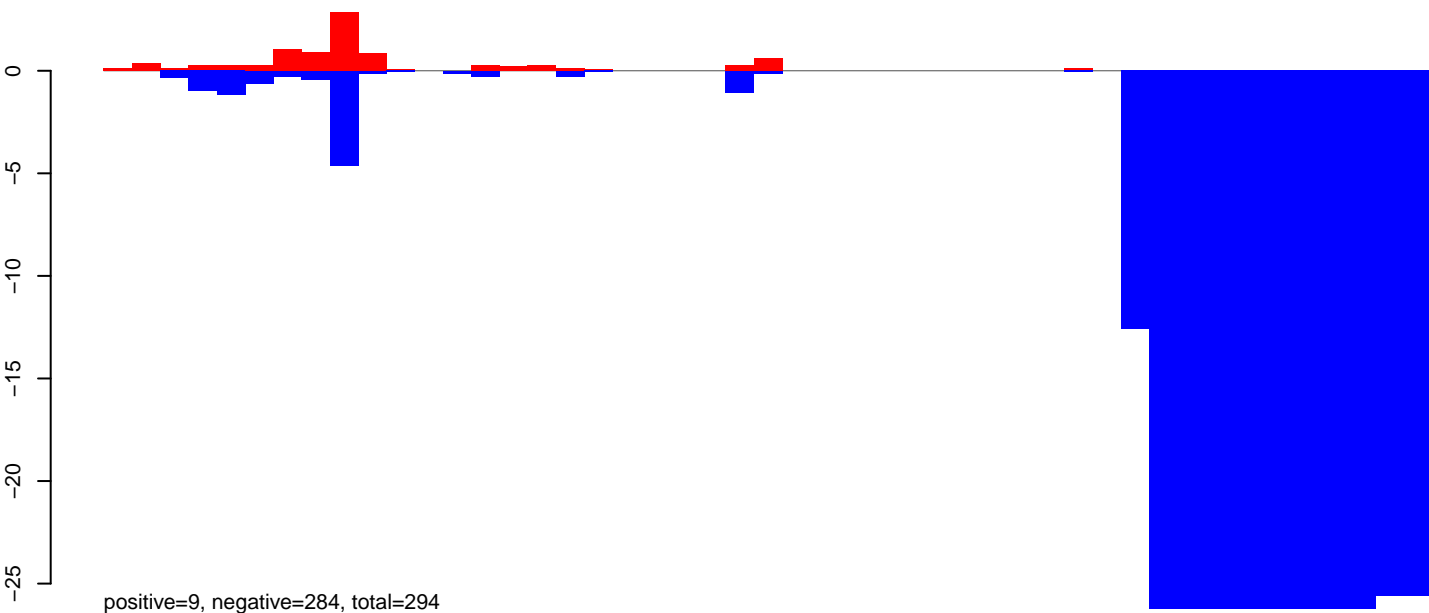
CuQuin_Mcarc_NL.rep



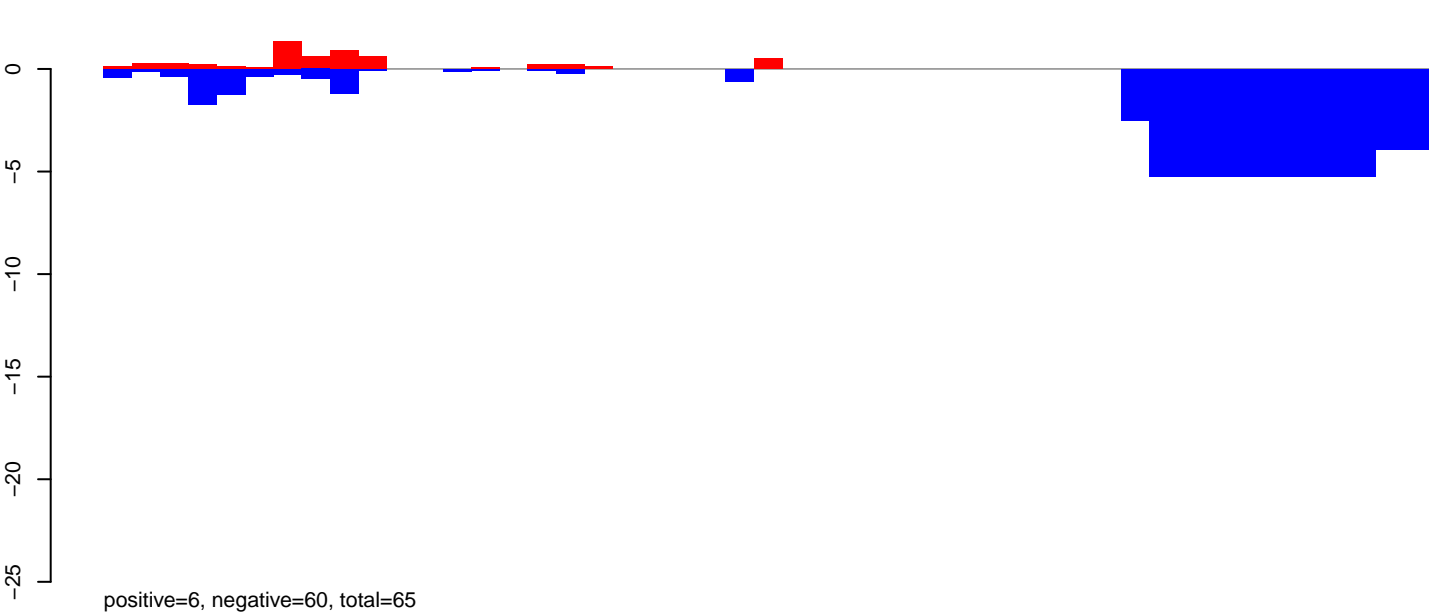
Window size=25, length=953, TE@rnd-3_family-34@DNA_Sola-2@39:1-953

0 200 400 600 800 1000

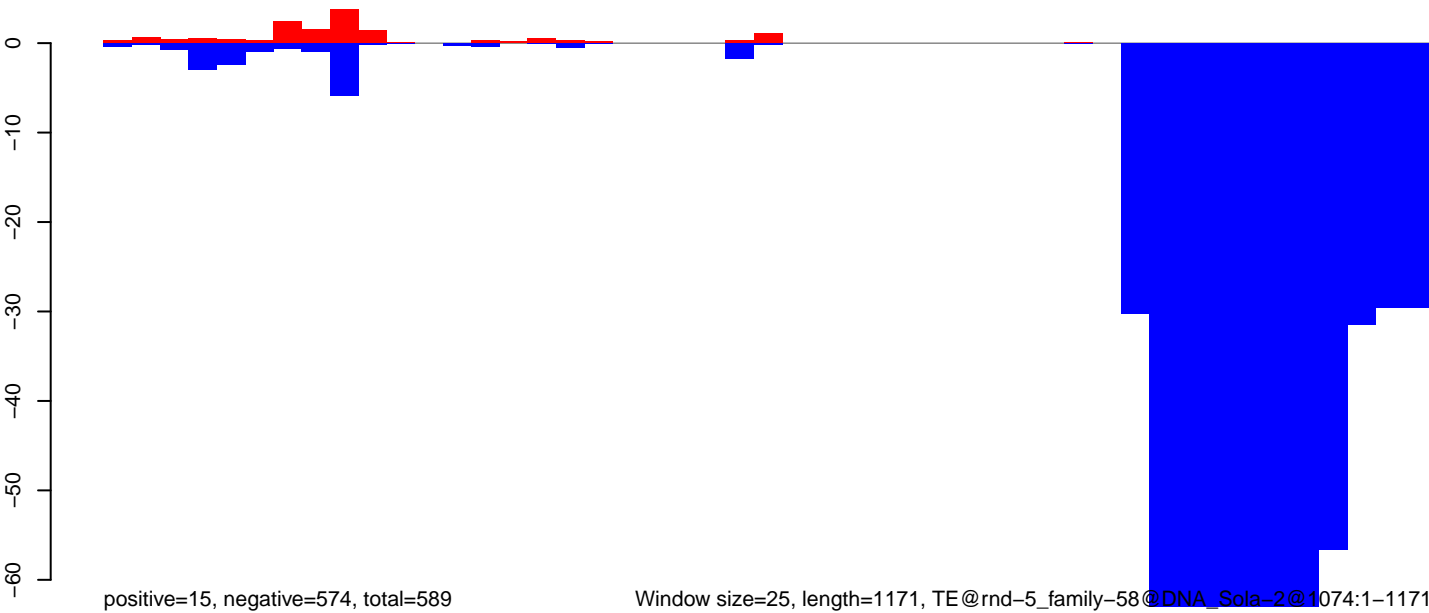
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



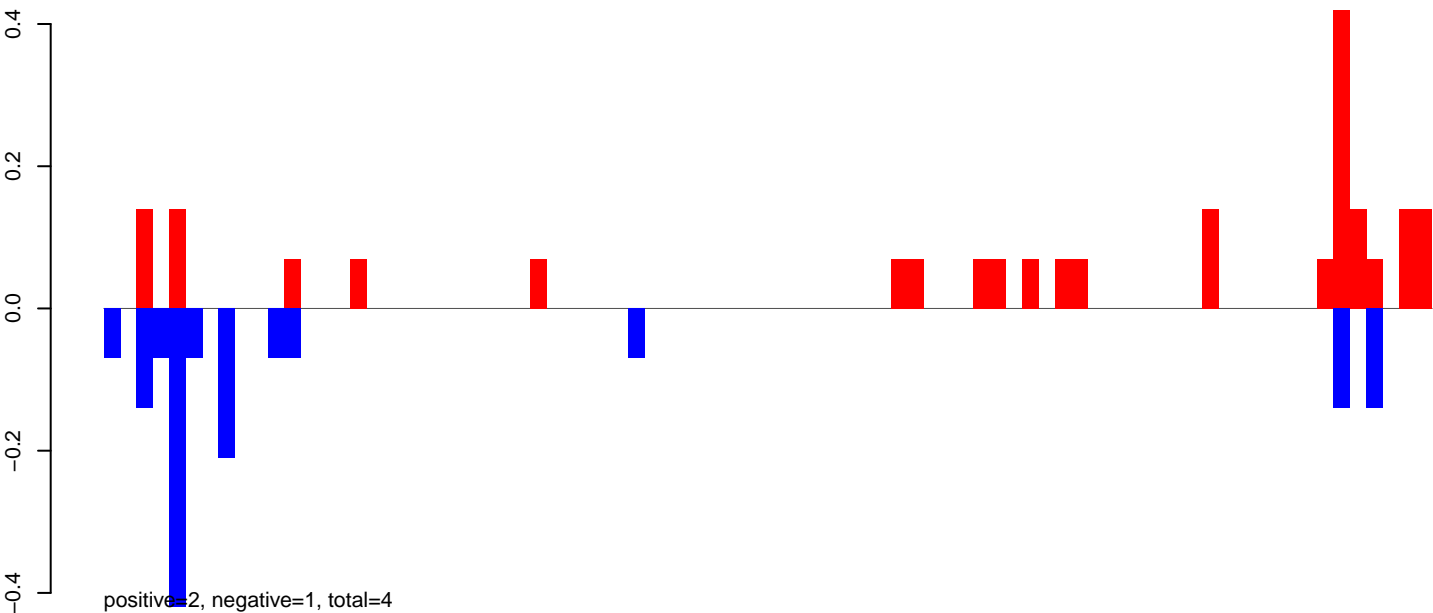
CuQuin_Mcarc_NL.rep



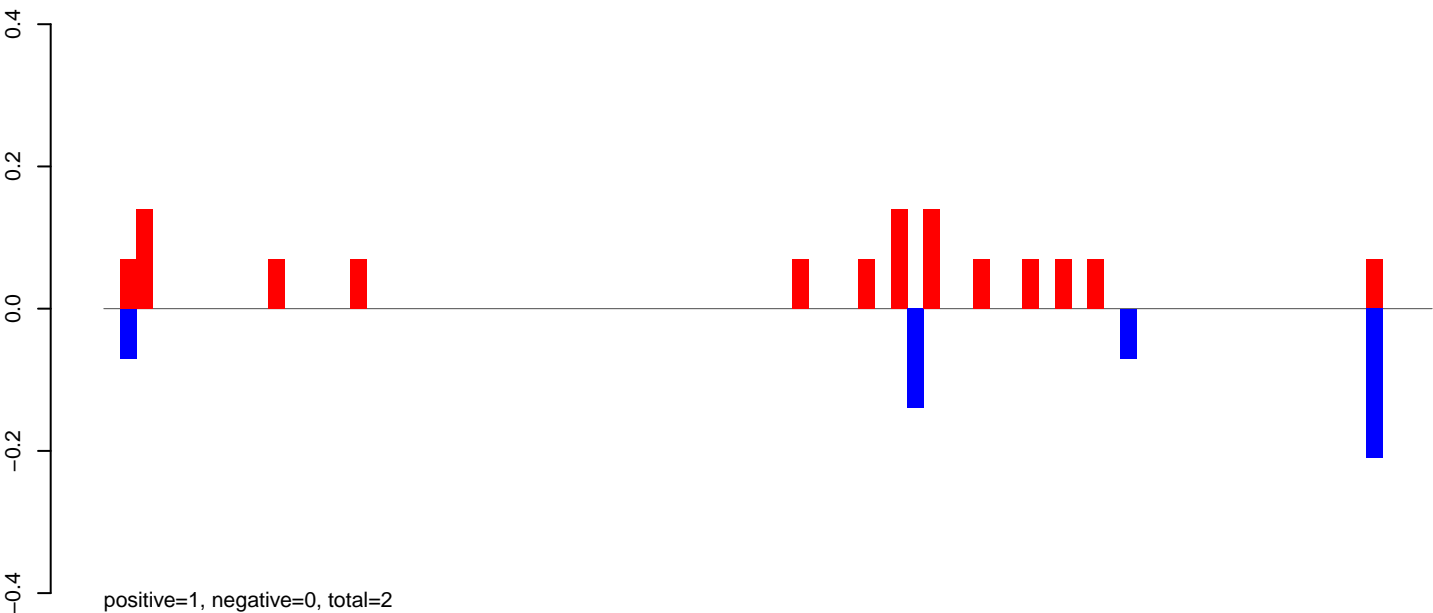
Window size=25, length=1171, TE@rnd-5_family-58@DNA_Sola-2@1074:1-1171

0 200 400 600 800 1000 1200

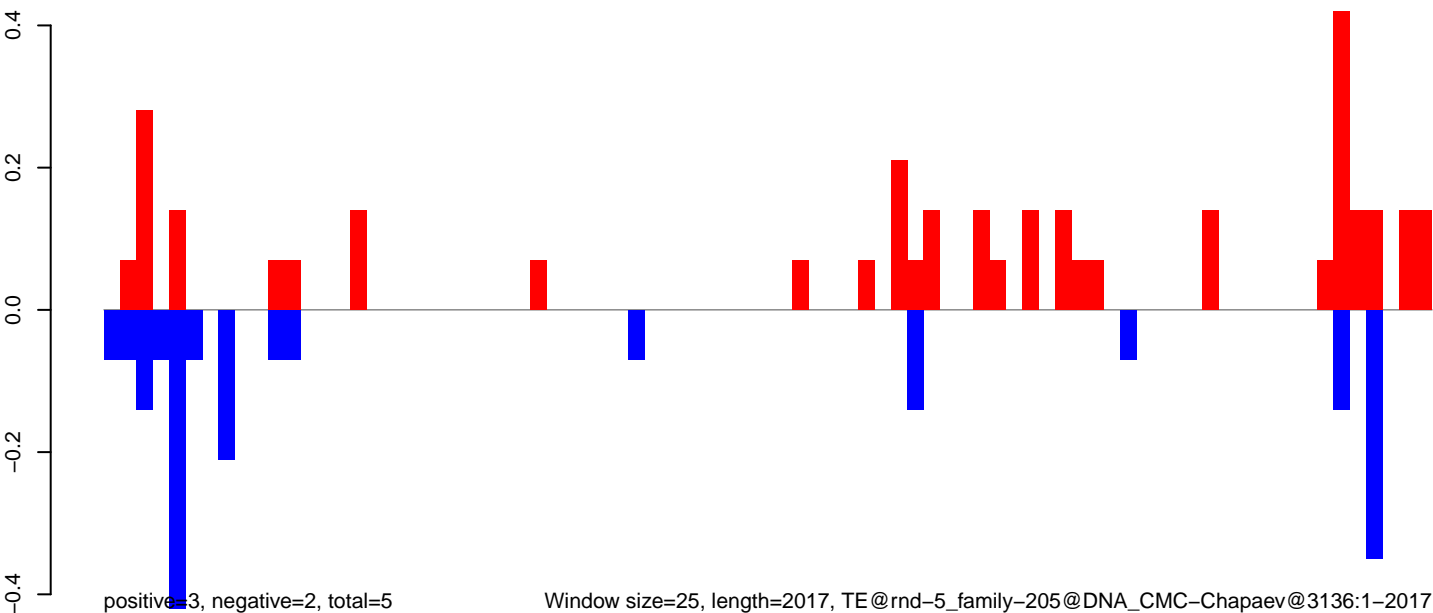
CuQuin_Mcarc_NL.18_23.rep



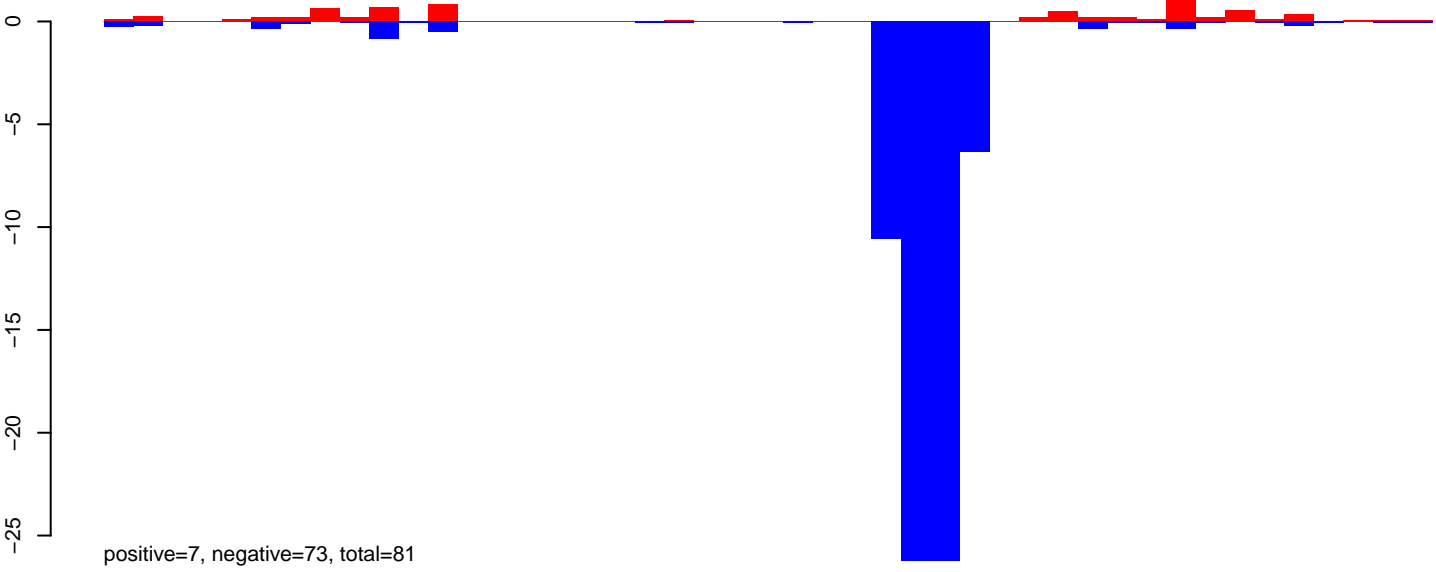
CuQuin_Mcarc_NL.24_35.rep



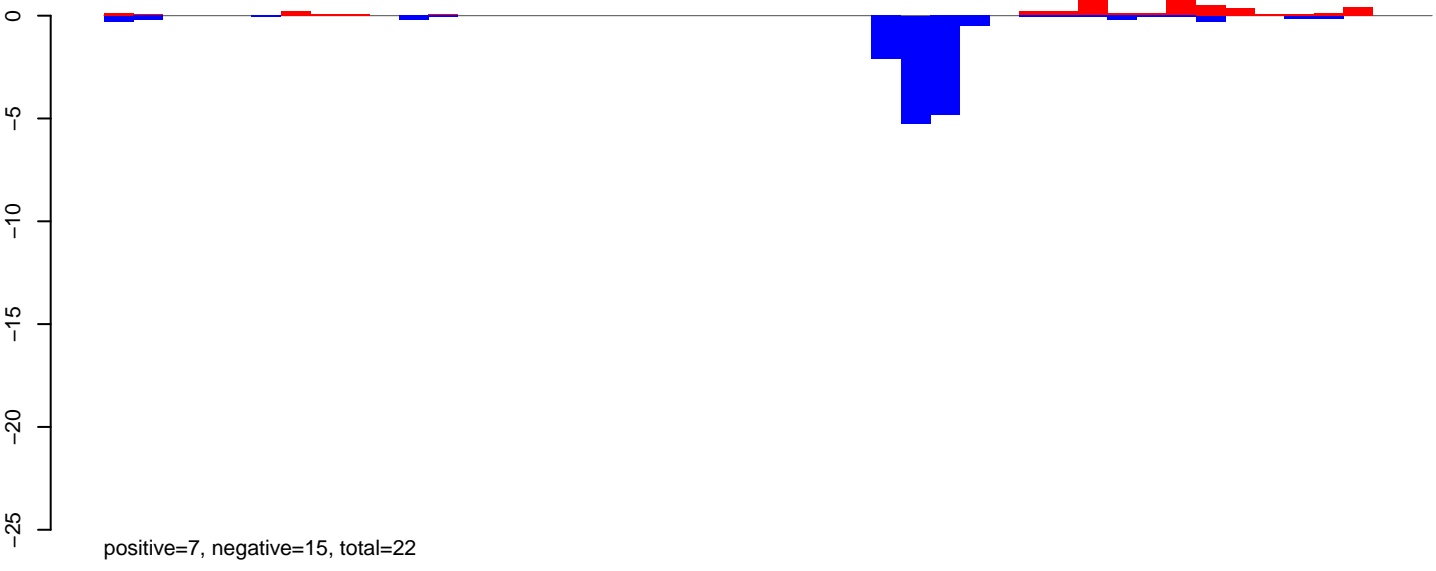
CuQuin_Mcarc_NL.rep



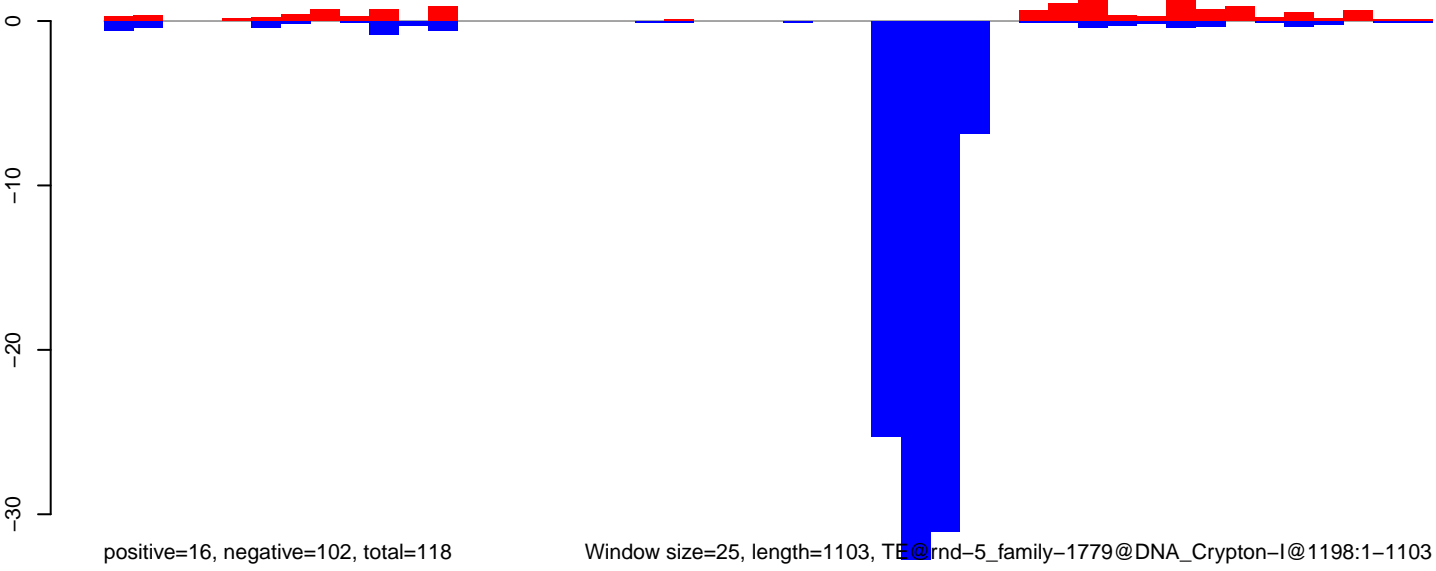
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

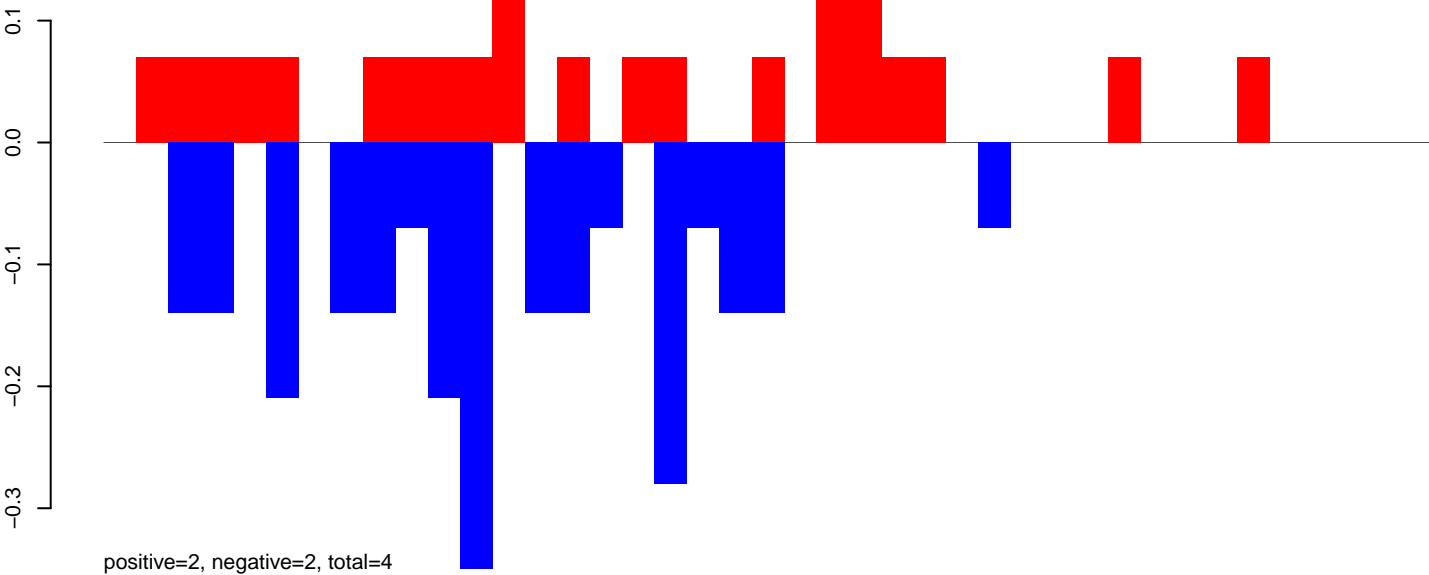


CuQuin_Mcarc_NL.rep

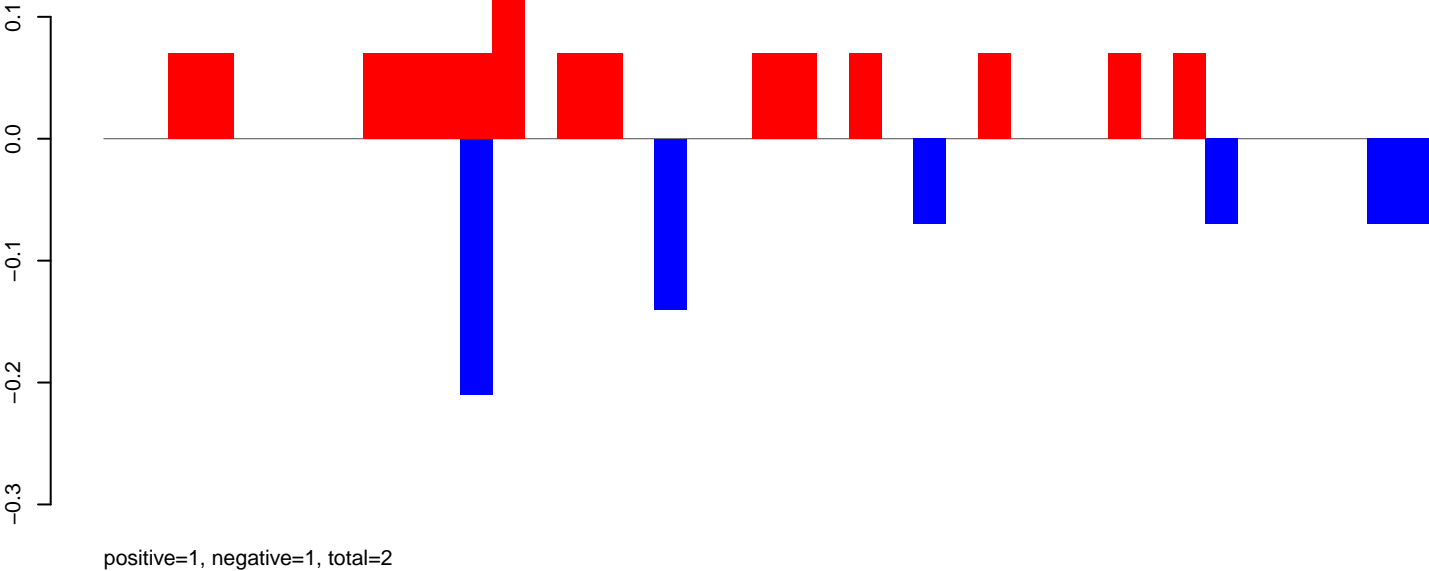


Window size=25, length=1103, TE@rnd-5_family-1779@DNA_Crypton-I@1198:1-1103

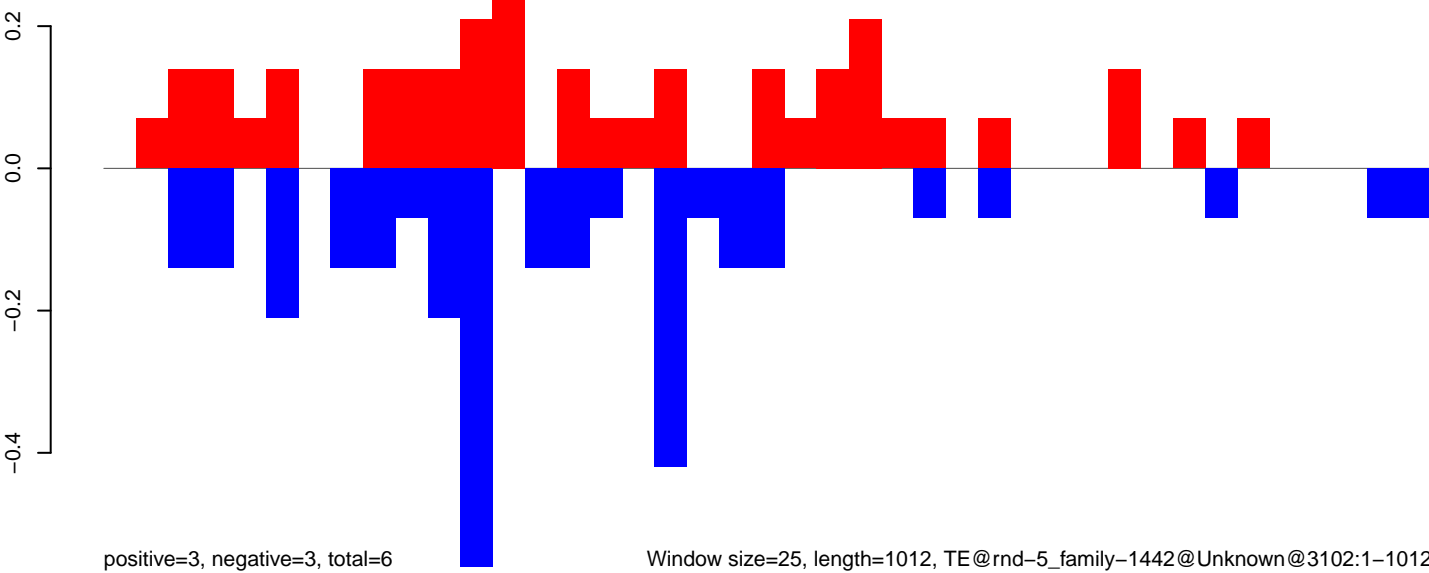
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

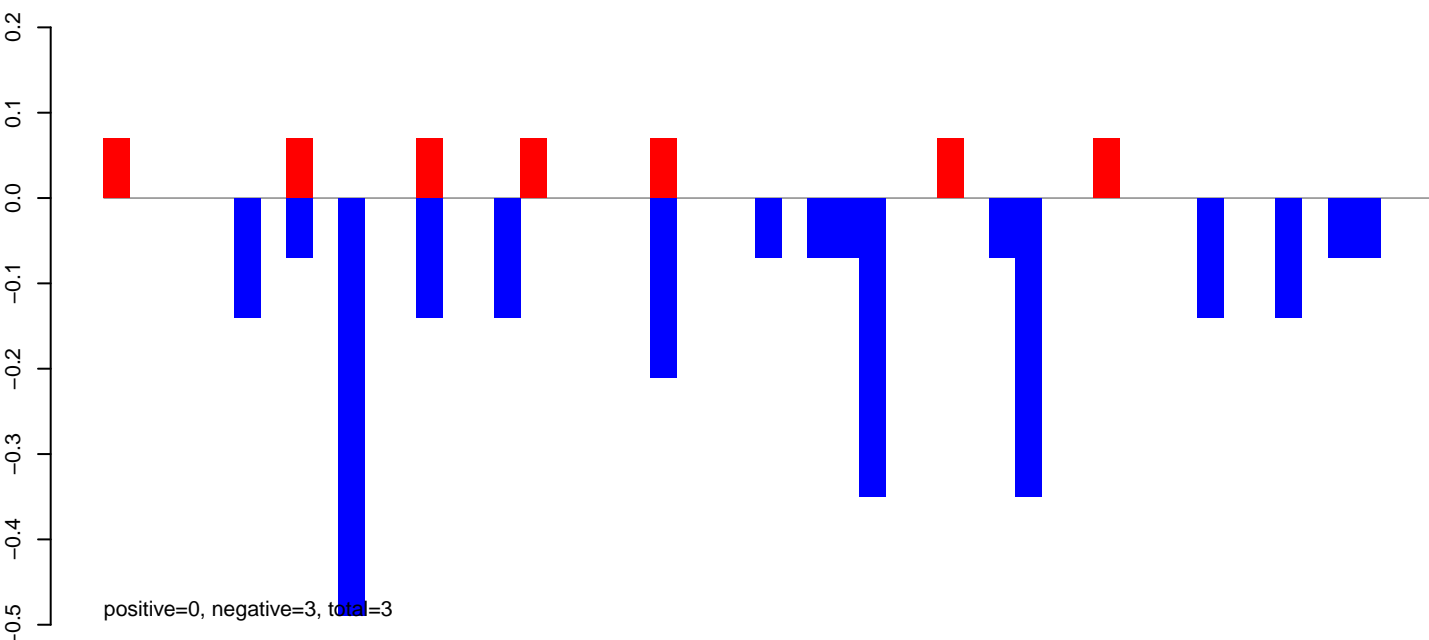


CuQuin_Mcarc_NL.rep

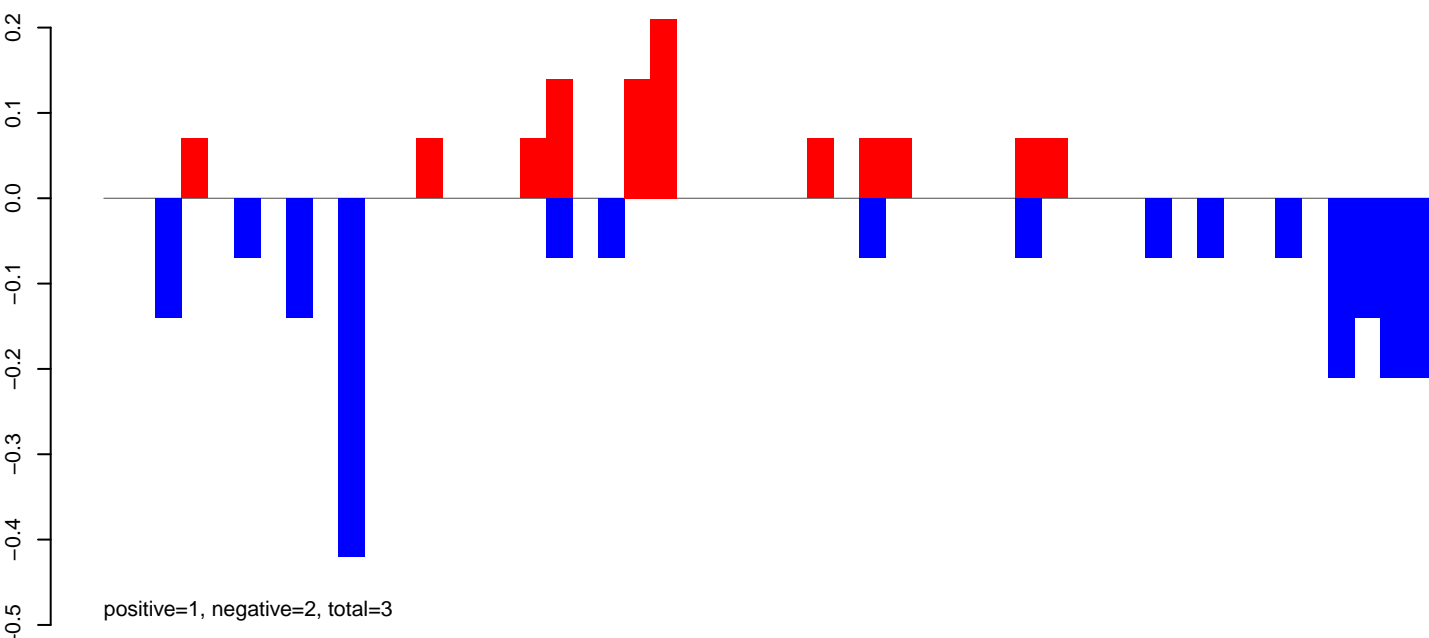


Window size=25, length=1012, TE@rnd-5_family-1442@Unknown@3102:1-1012

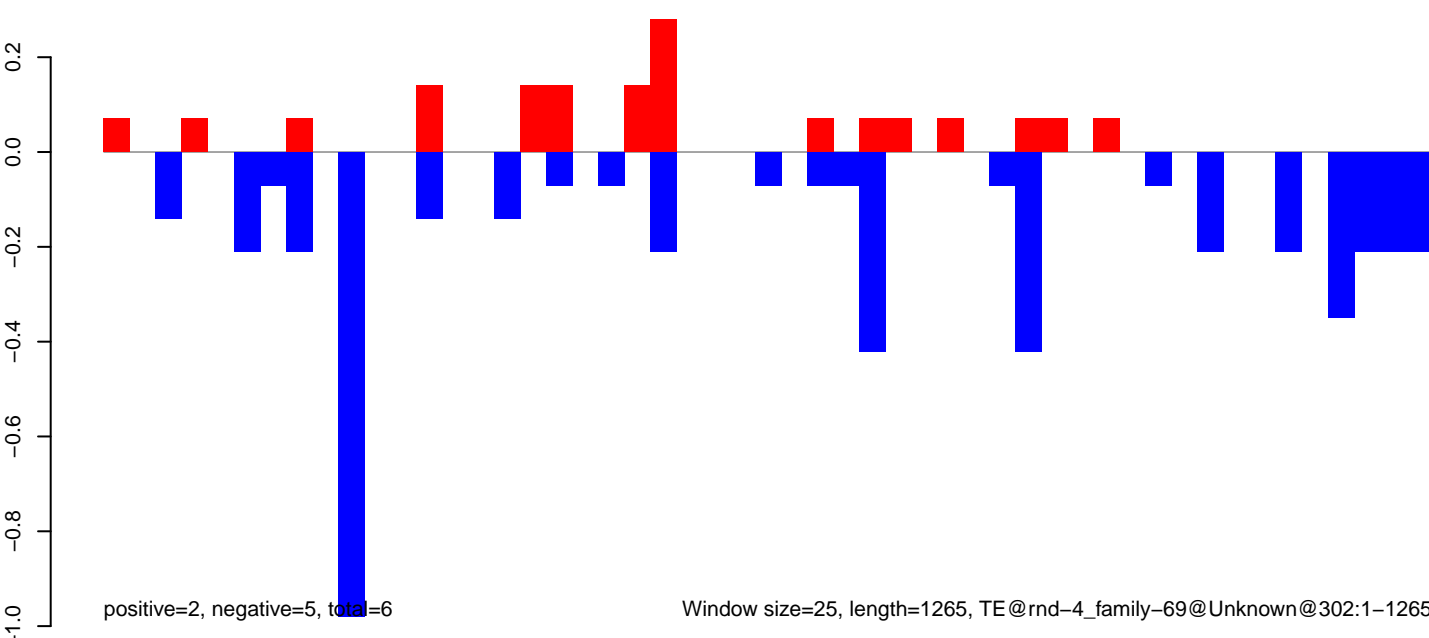
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



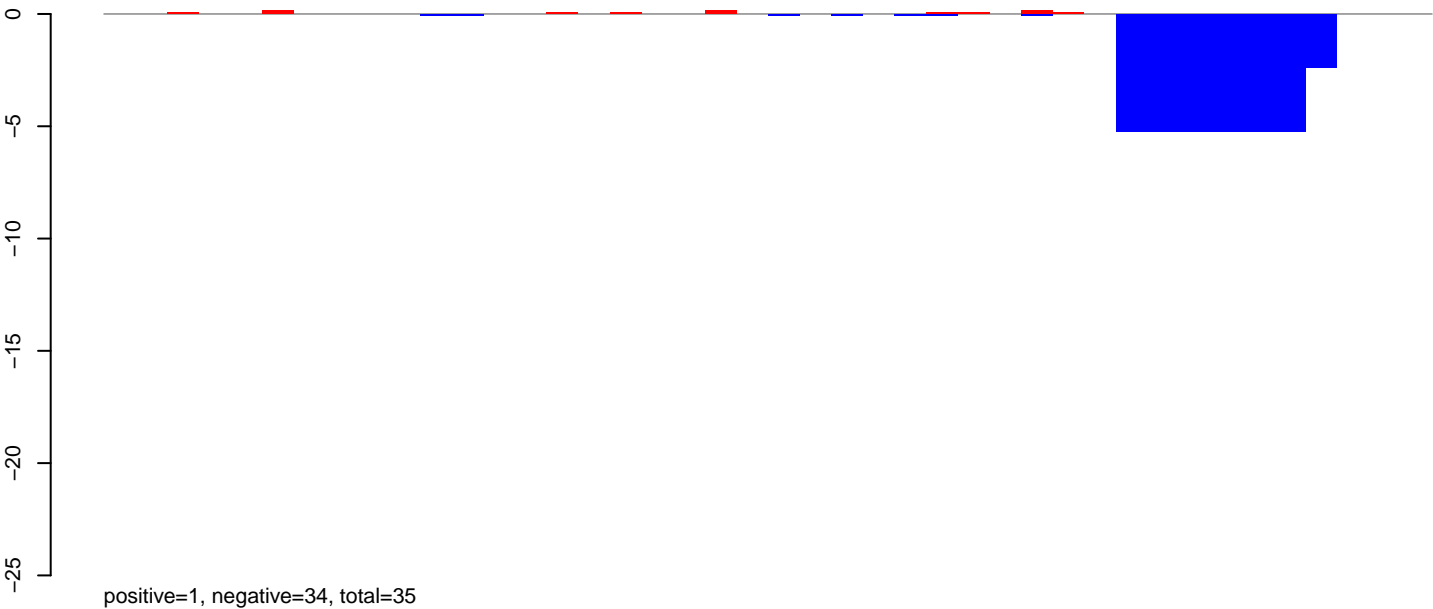
CuQuin_Mcarc_NL.rep



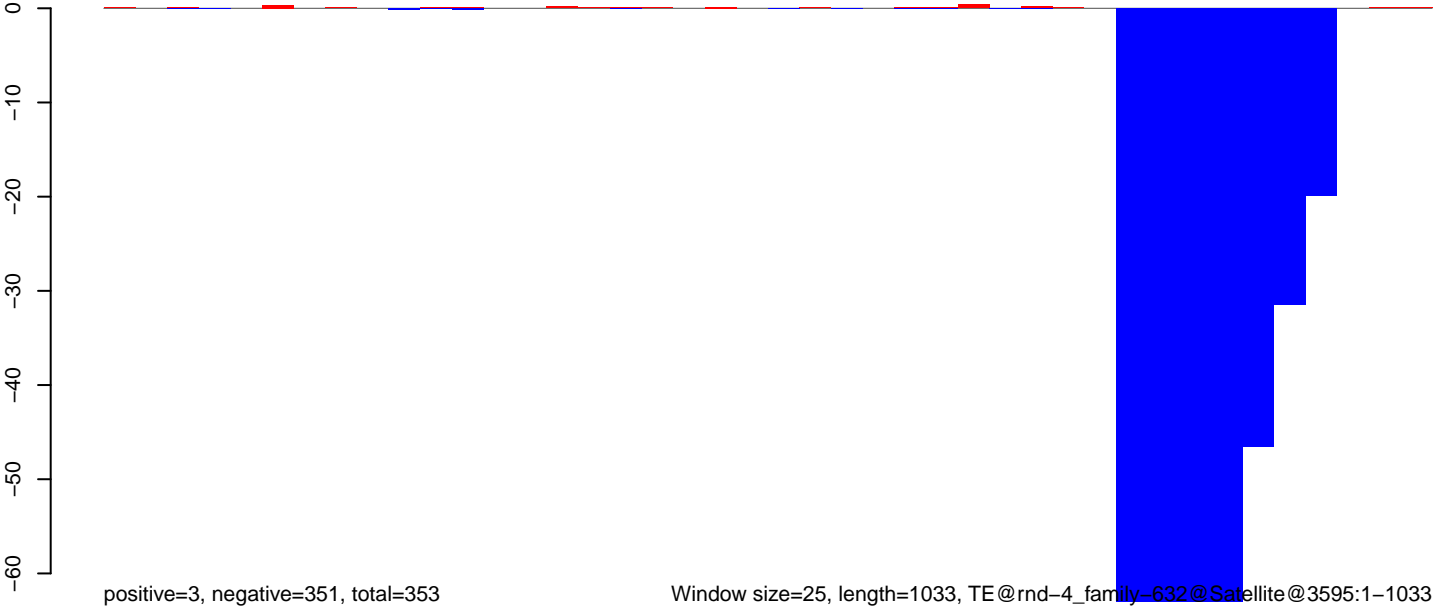
CuQuin_Mcarc_NL.18_23.rep



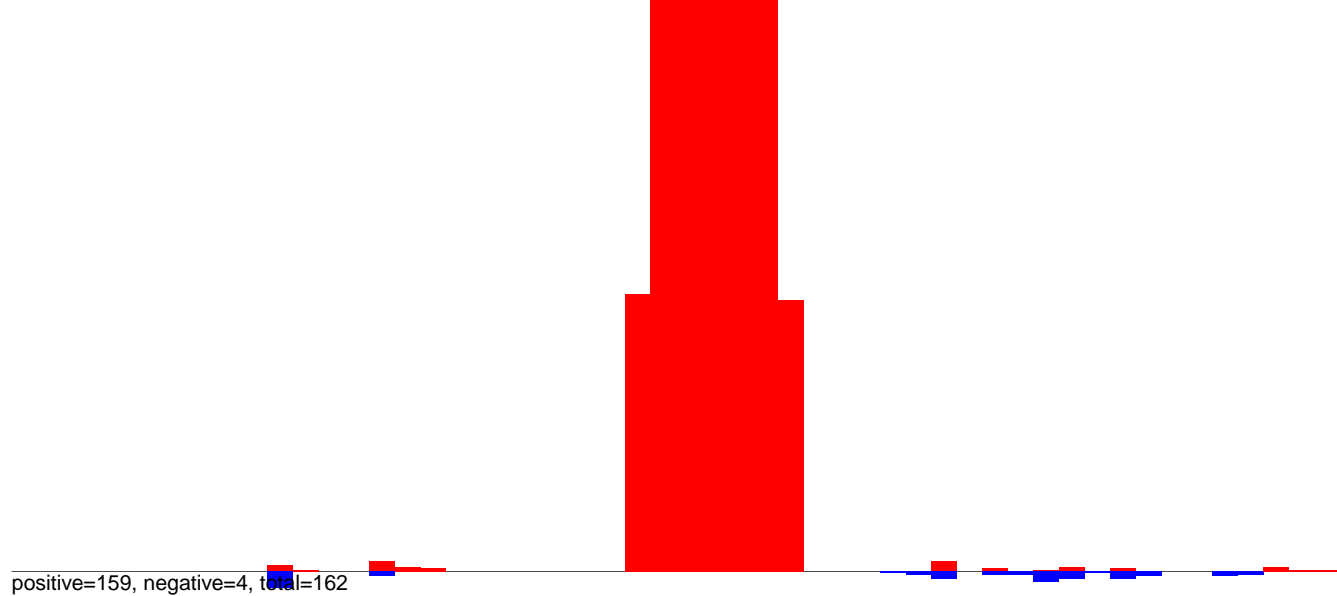
CuQuin_Mcarc_NL.24_35.rep



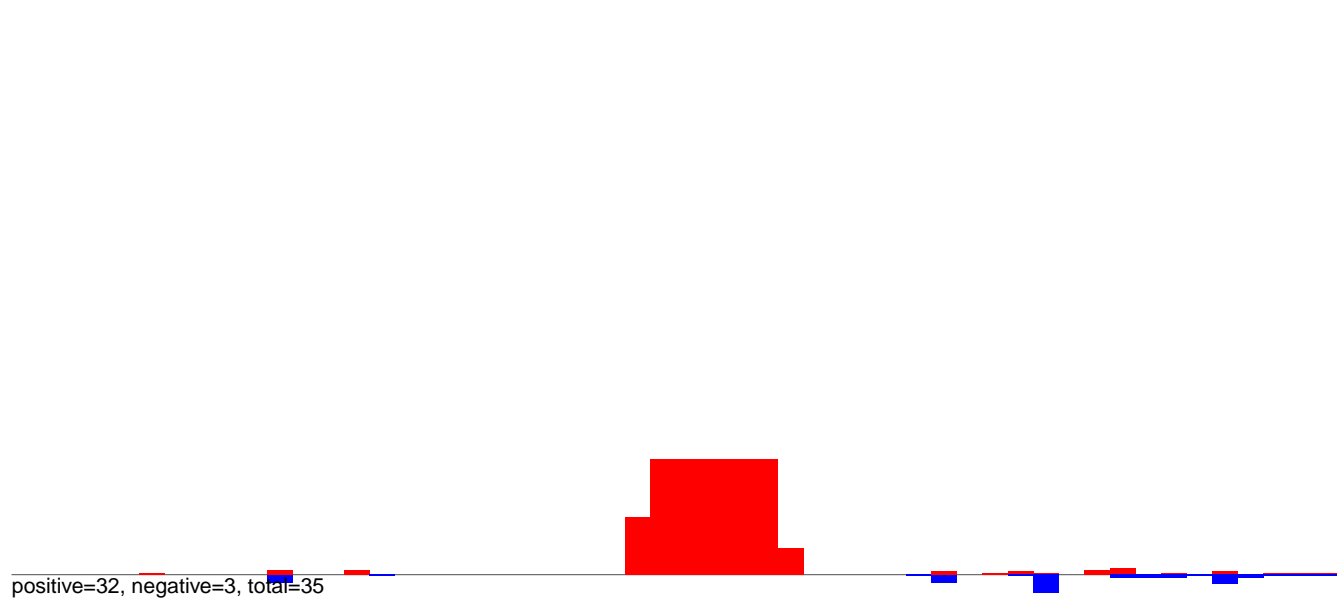
CuQuin_Mcarc_NL.rep



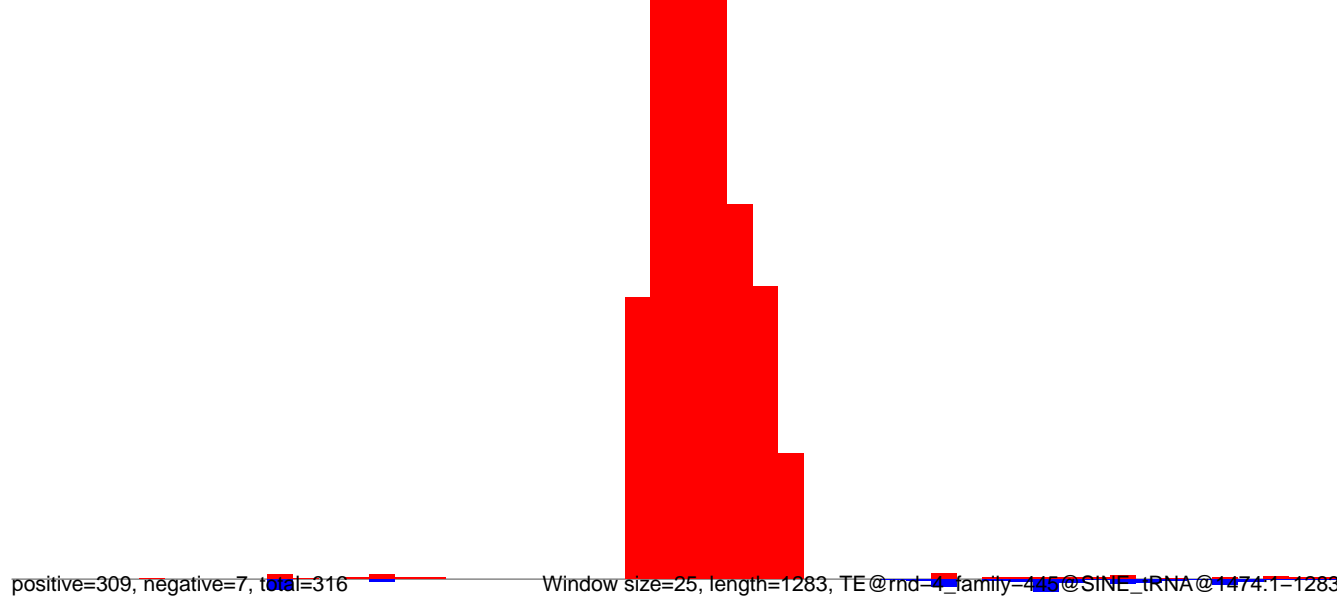
CuQuin_Mcarc_NL.18_23.rep



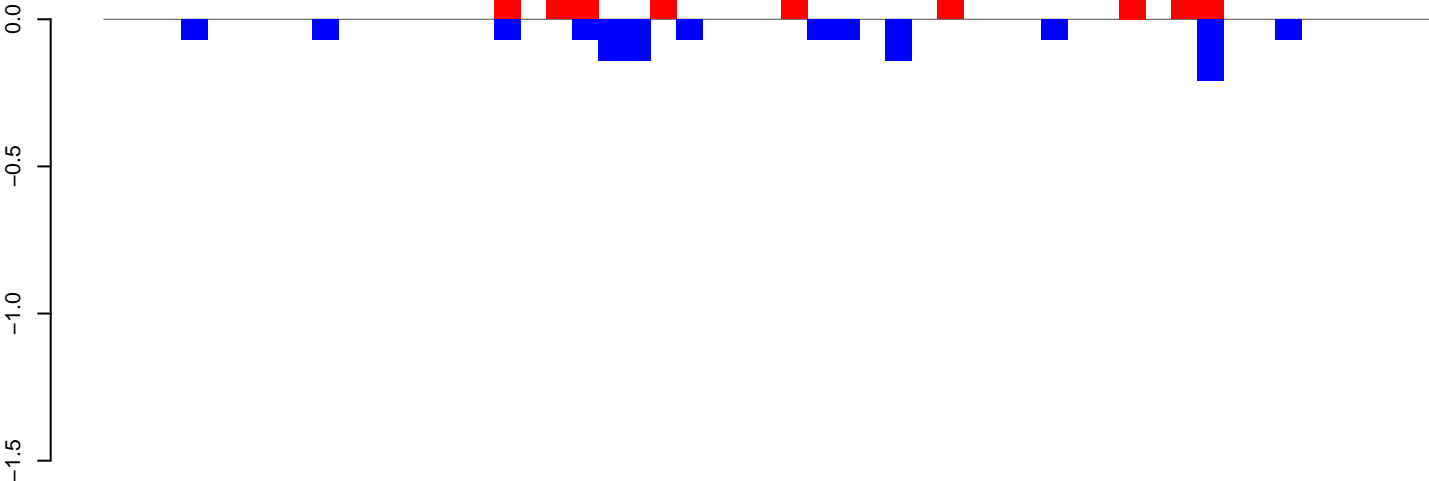
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

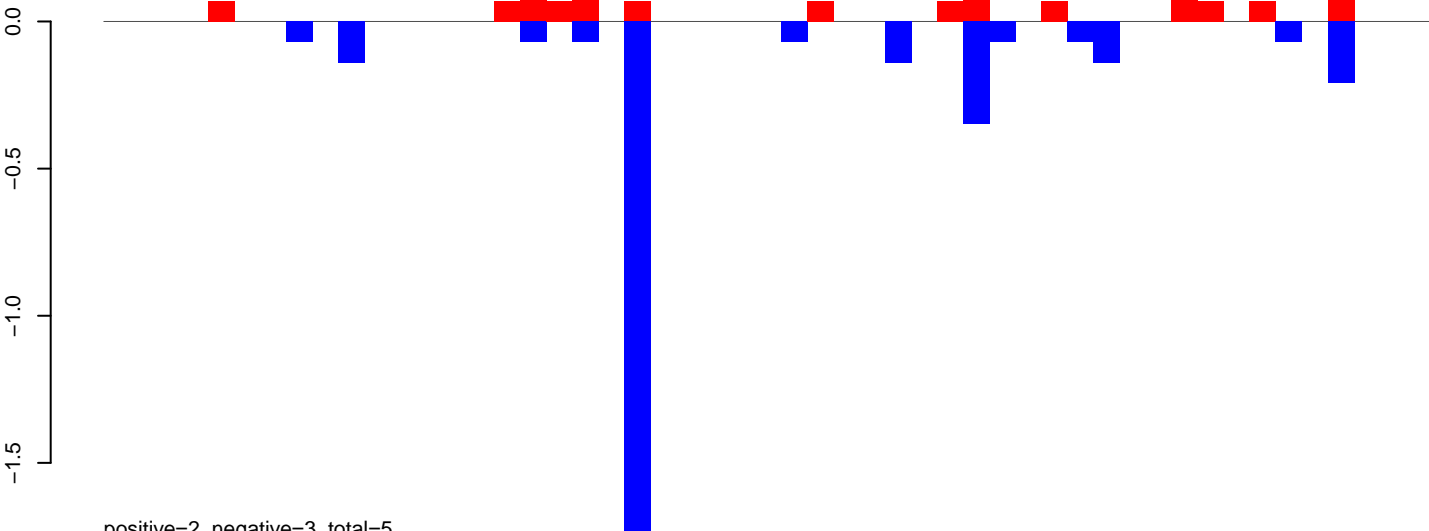


CuQuin_Mcarc_NL.18_23.rep



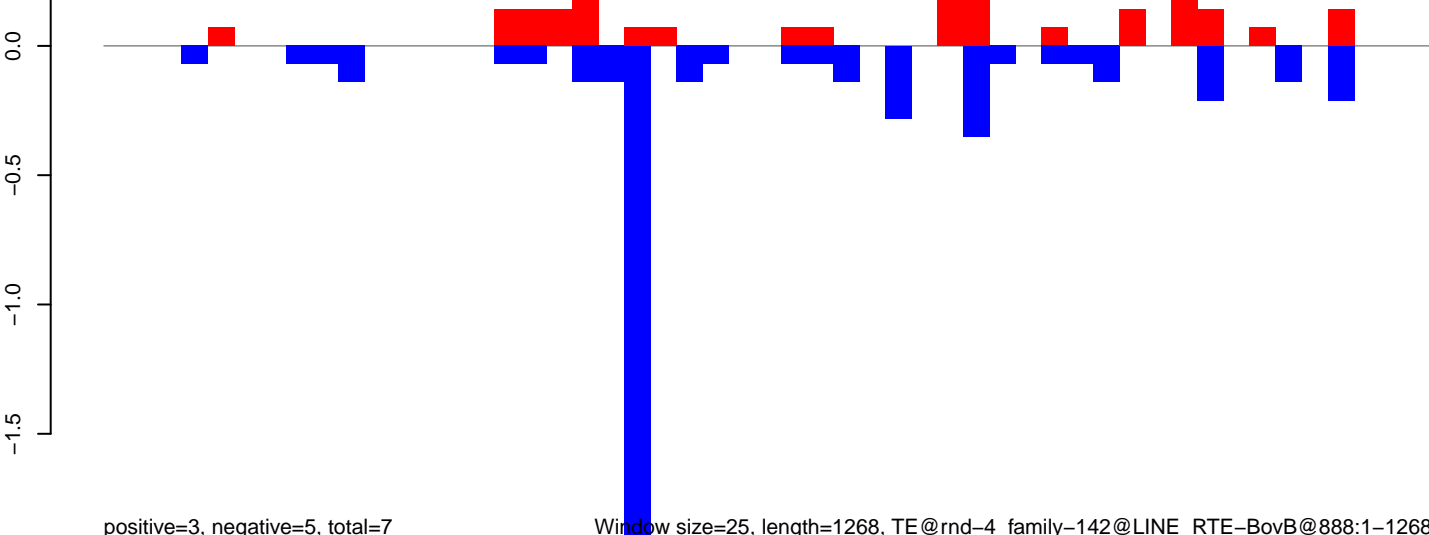
positive=1, negative=1, total=2

CuQuin_Mcarc_NL.24_35.rep



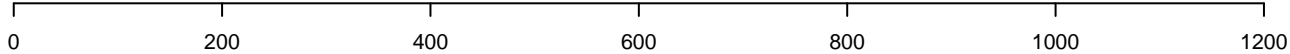
positive=2, negative=3, total=5

CuQuin_Mcarc_NL.rep

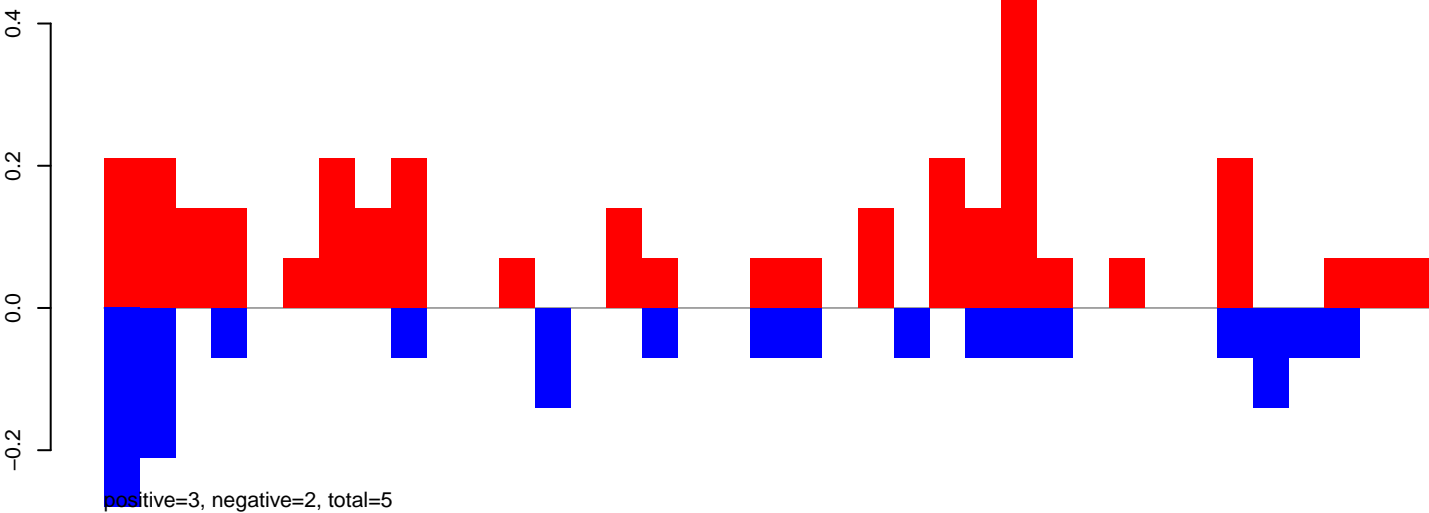


positive=3, negative=5, total=7

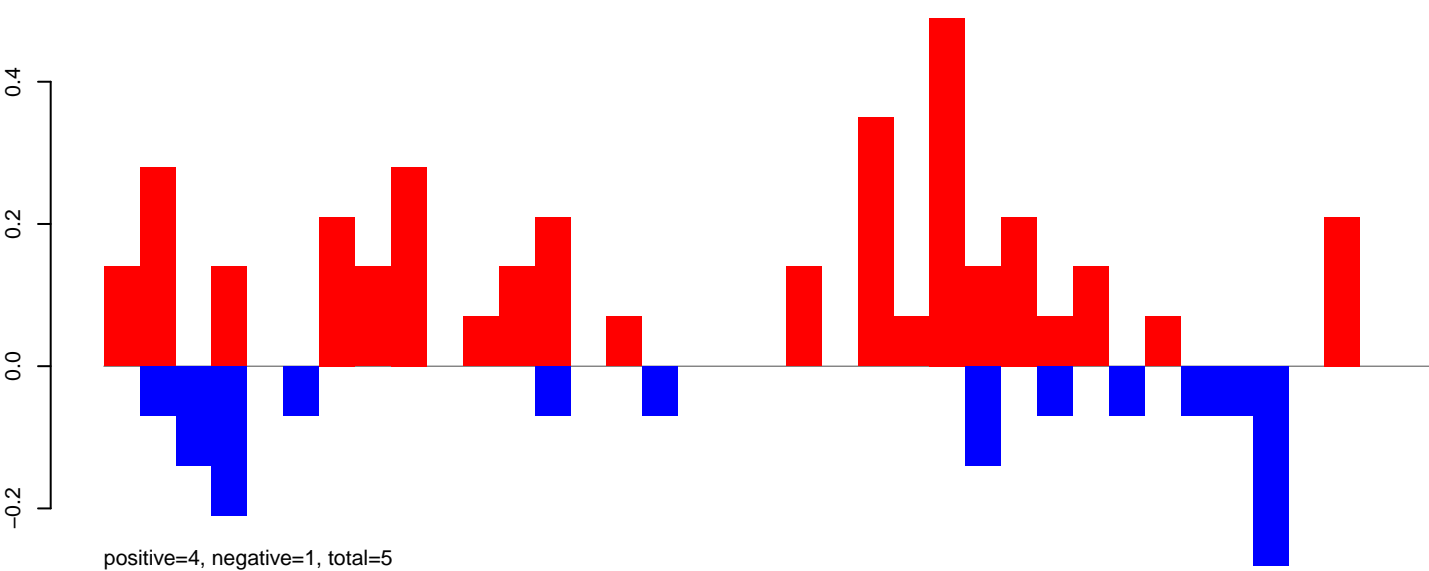
Window size=25, length=1268, TE@rnd-4_family-142@LINE_RTE-BovB@888:1-1268



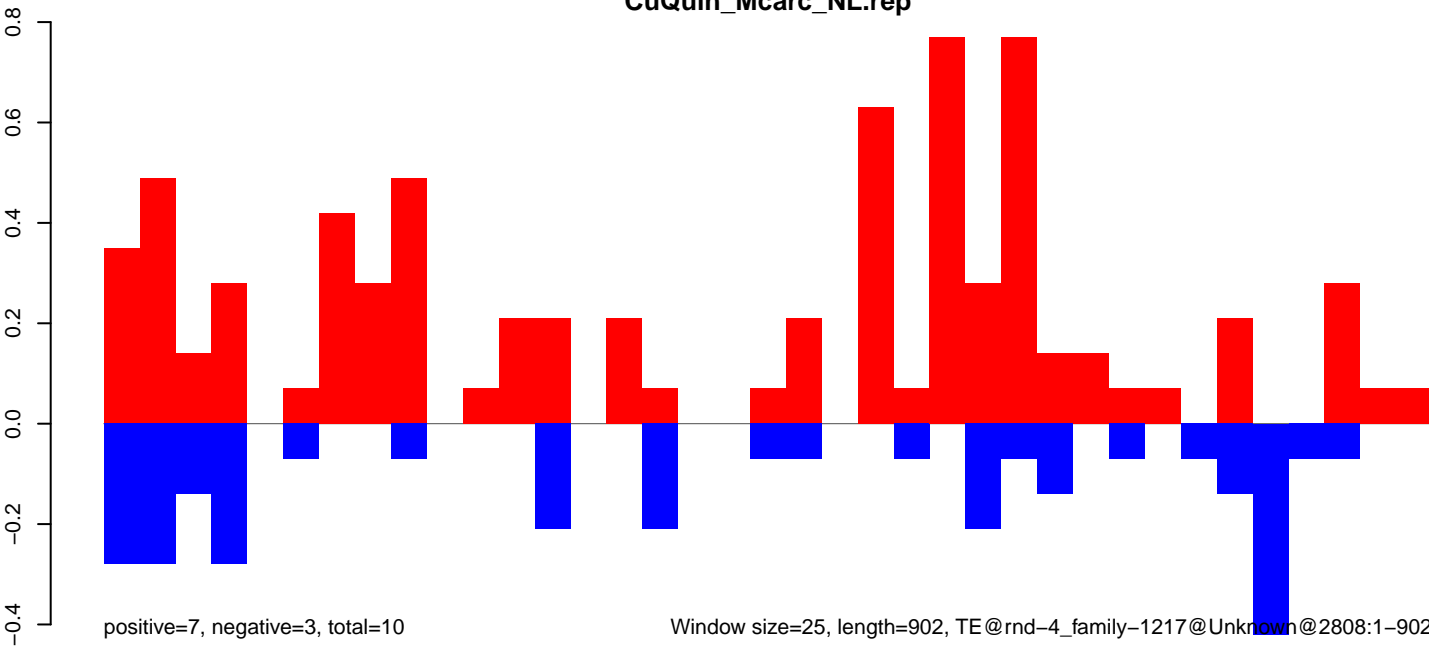
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

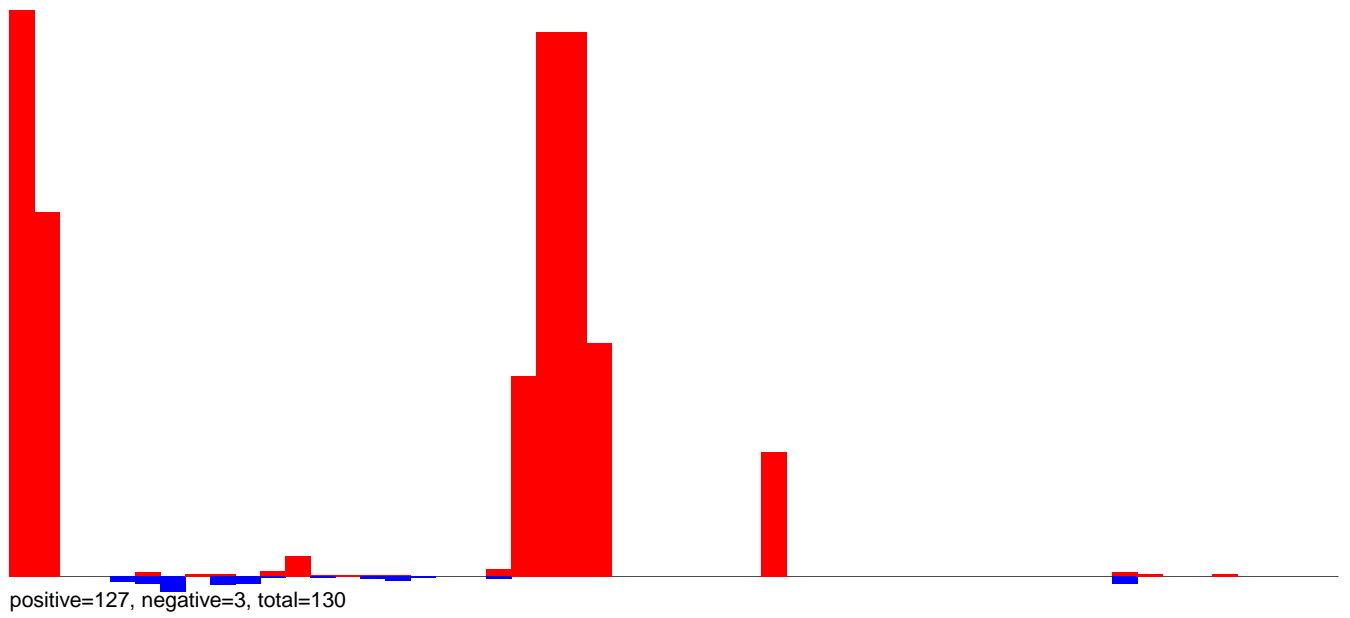


CuQuin_Mcarc_NL.rep

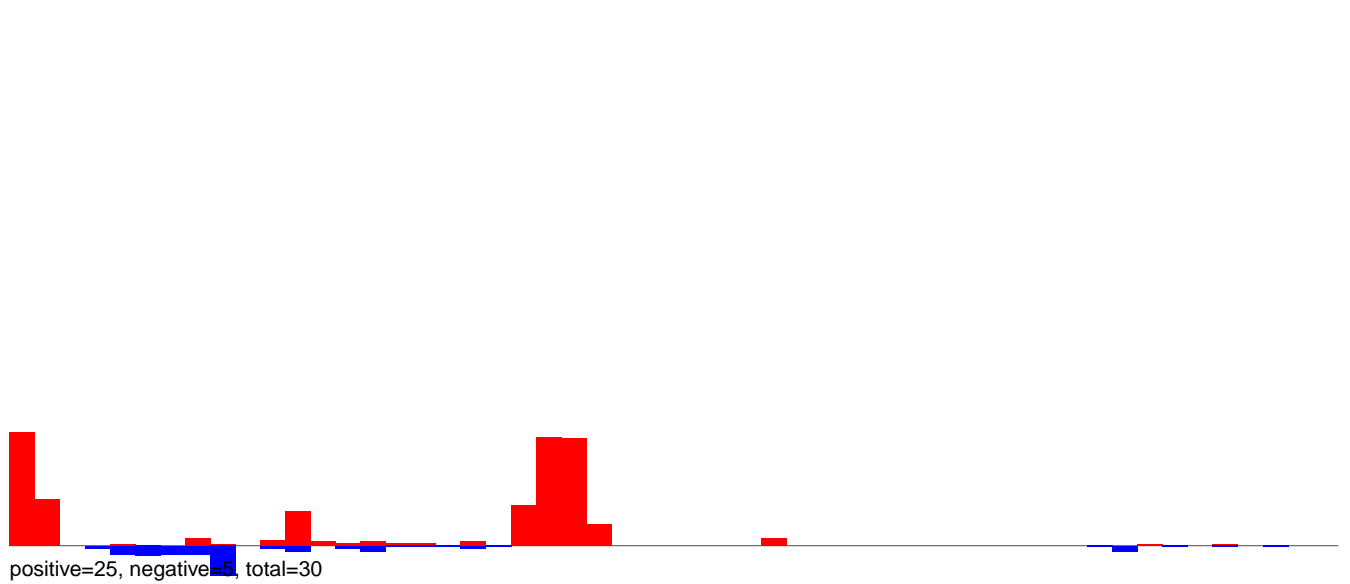


Window size=25, length=902, TE@rnd-4_family-1217@Unknown@2808:1-902

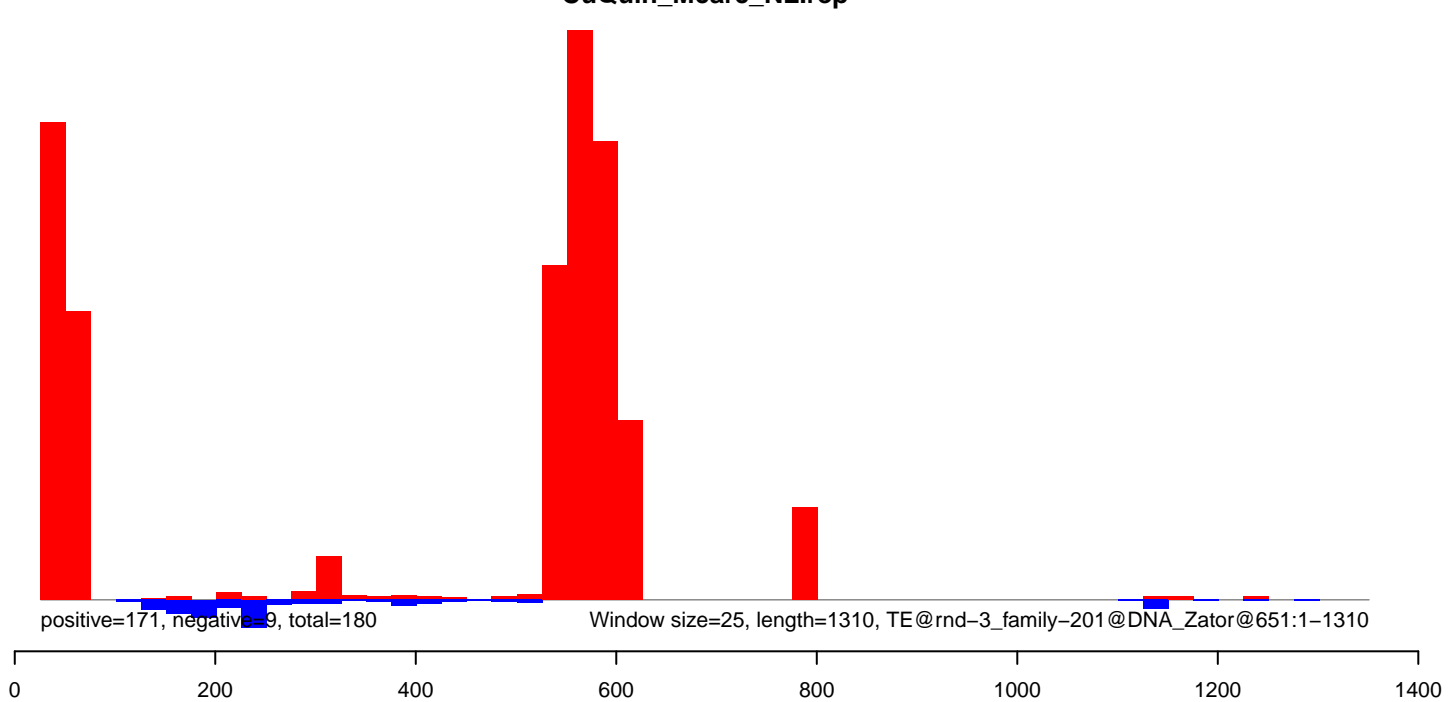
CuQuin_Mcarc_NL.18_23.rep



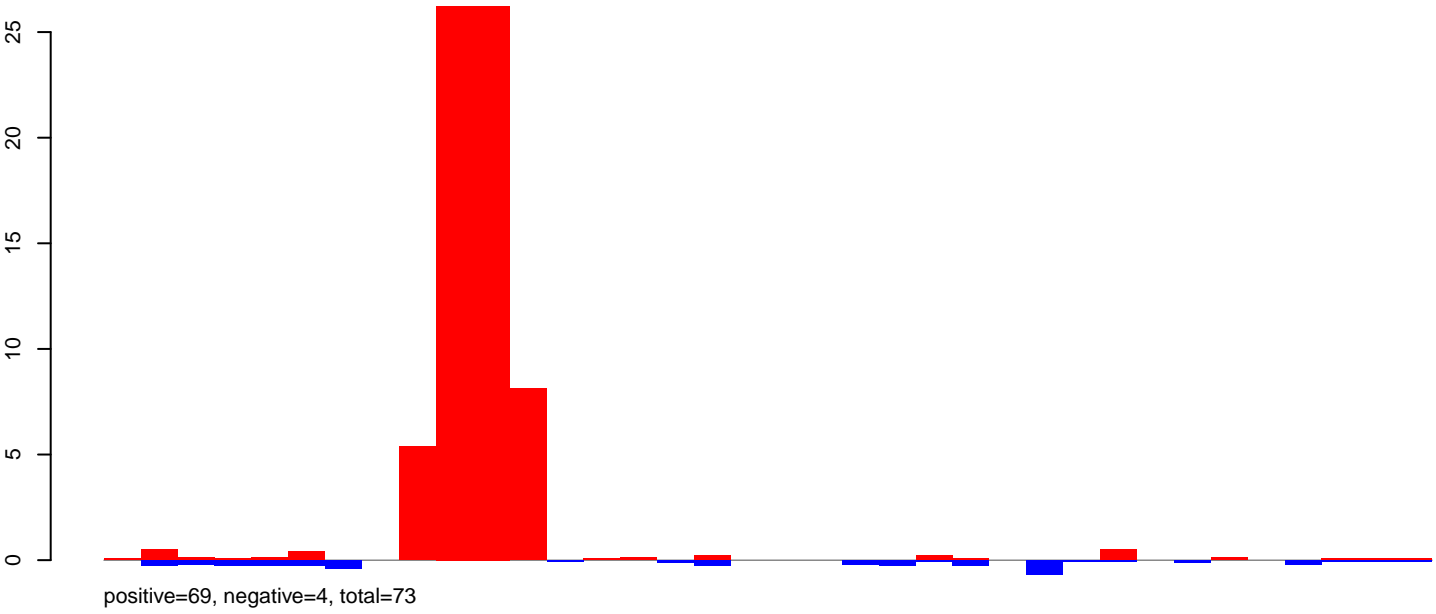
CuQuin_Mcarc_NL.24_35.rep



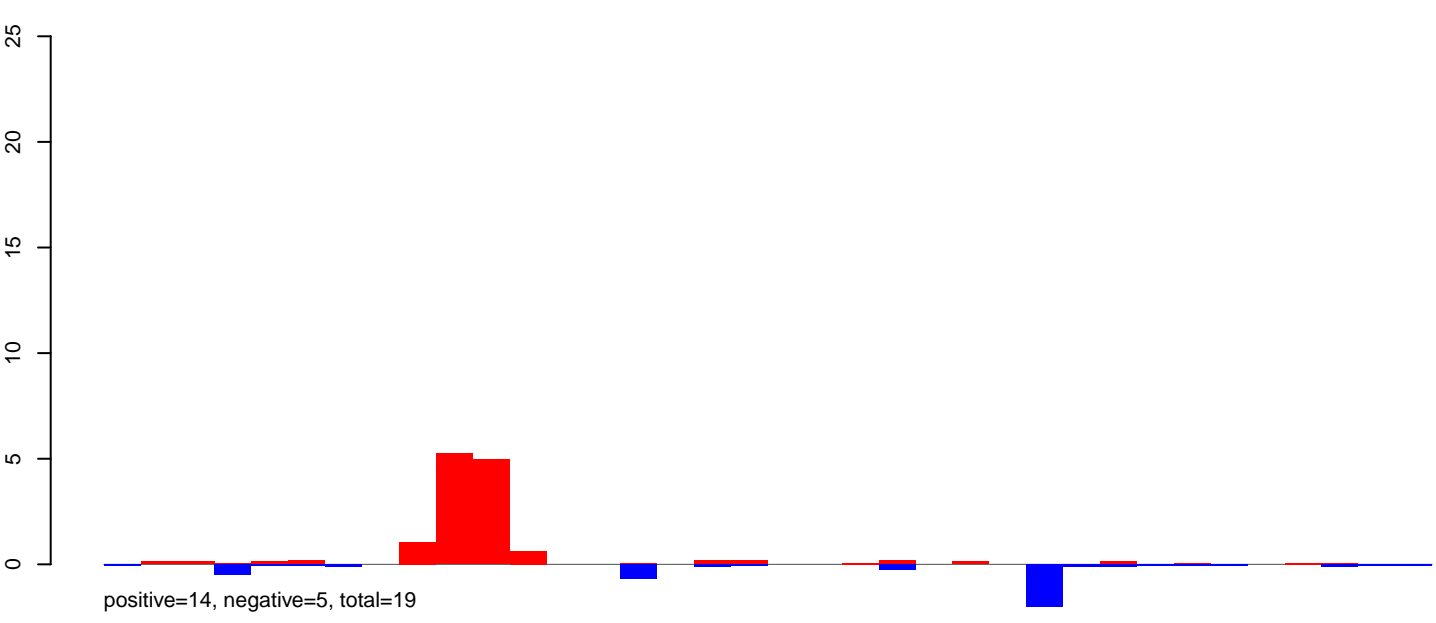
CuQuin_Mcarc_NL.rep



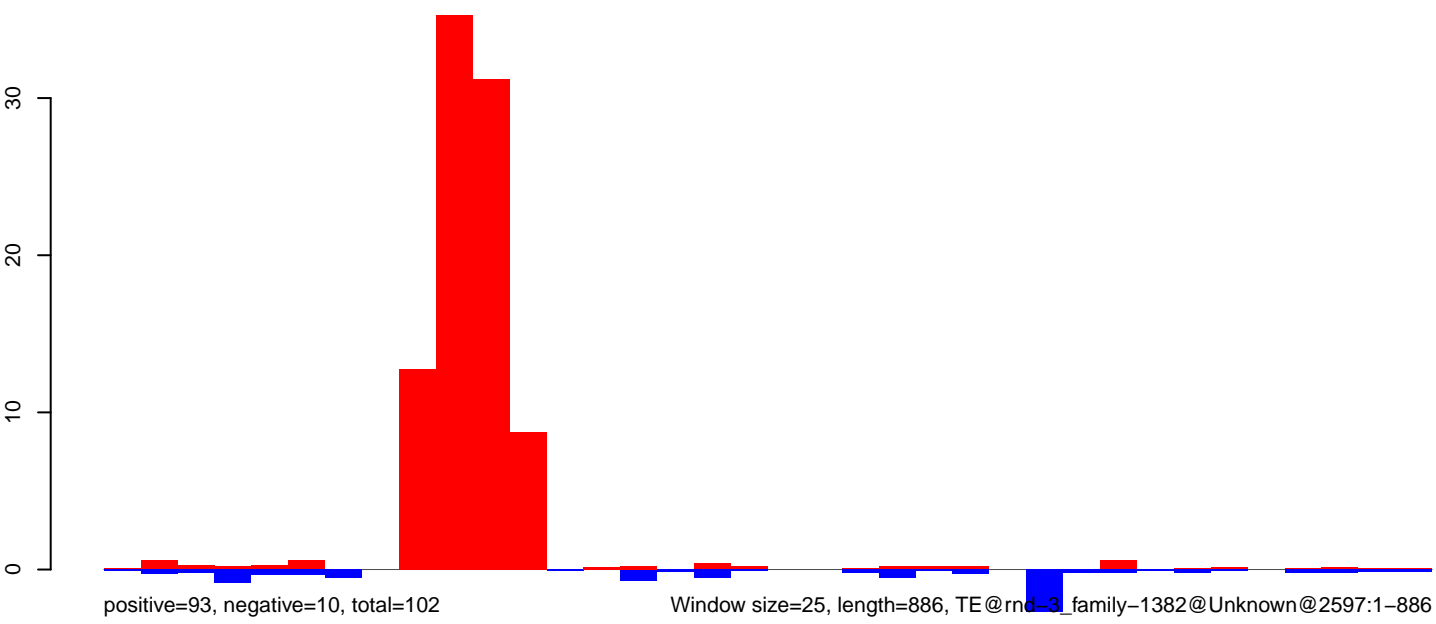
CuQuin_Mcarc_NL.18_23.rep



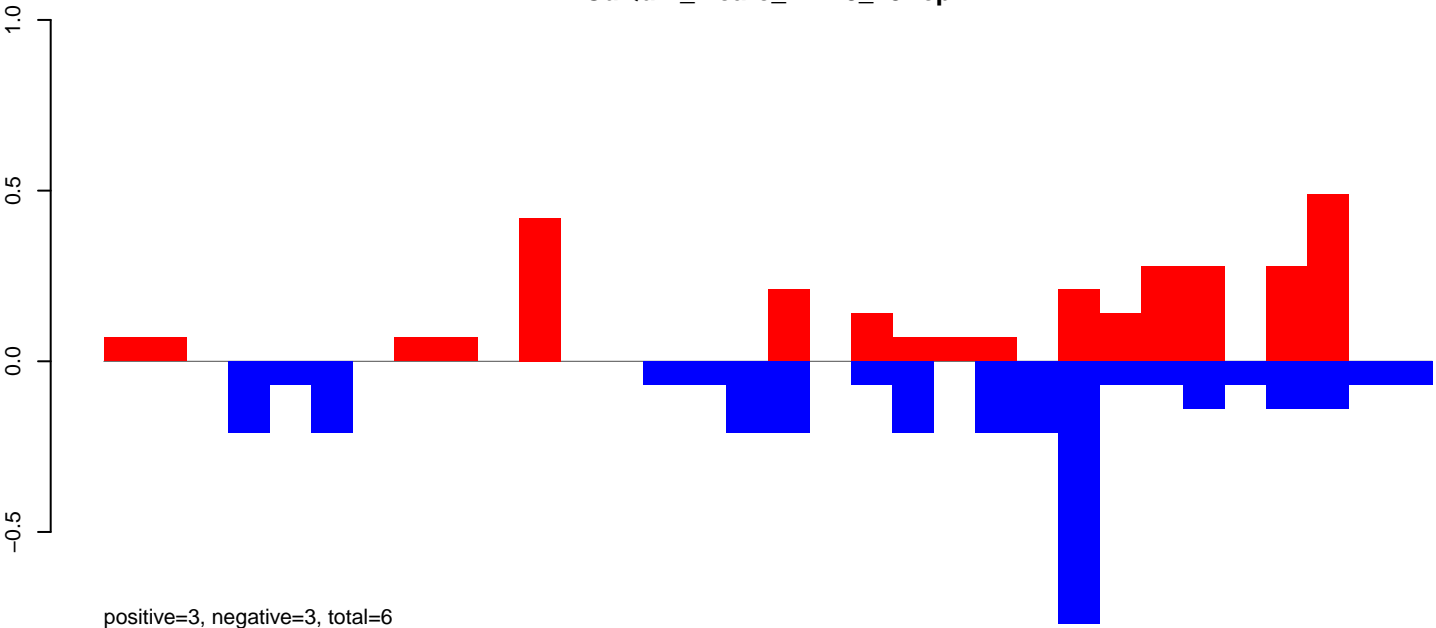
CuQuin_Mcarc_NL.24_35.rep



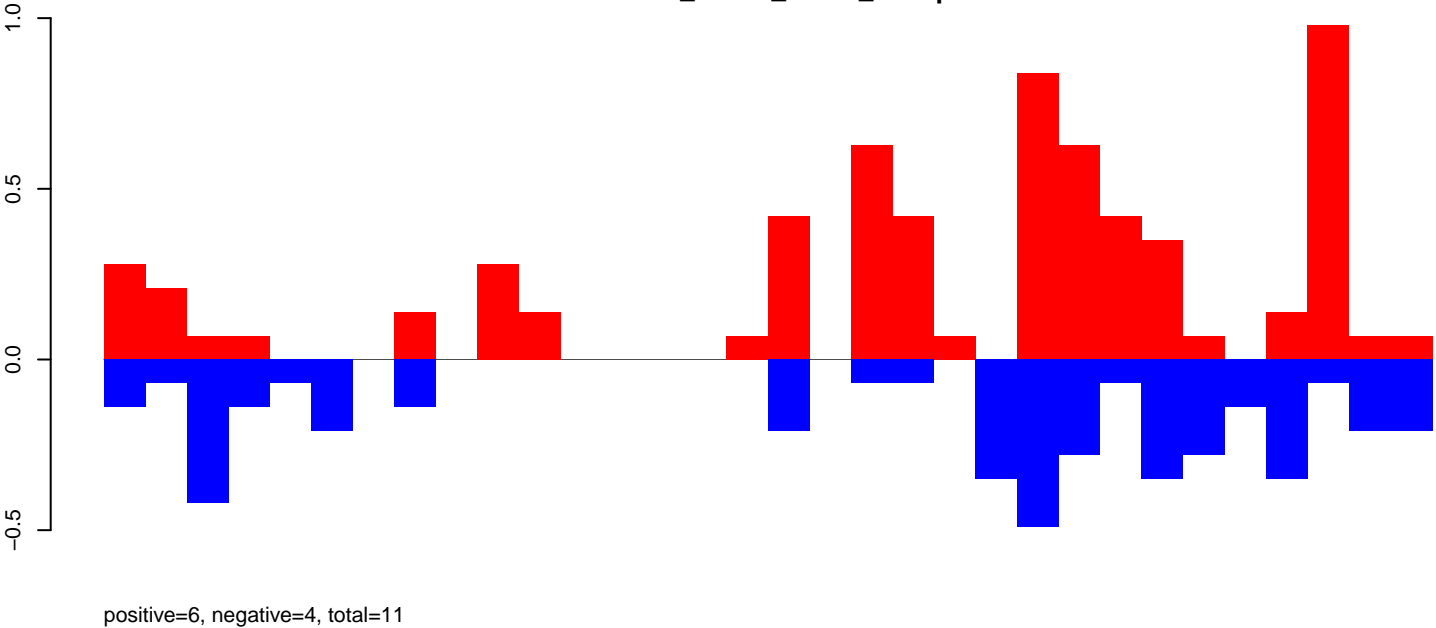
CuQuin_Mcarc_NL.rep



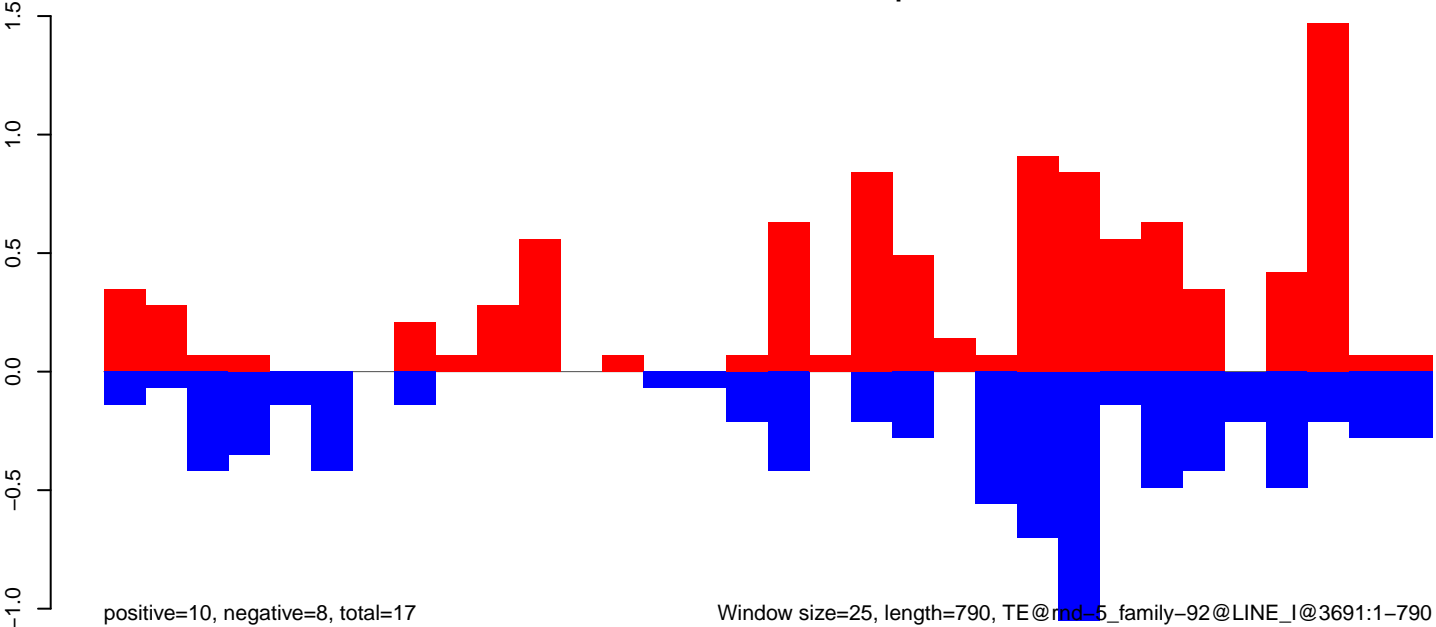
CuQuin_Mcarc_NL.18_23.rep



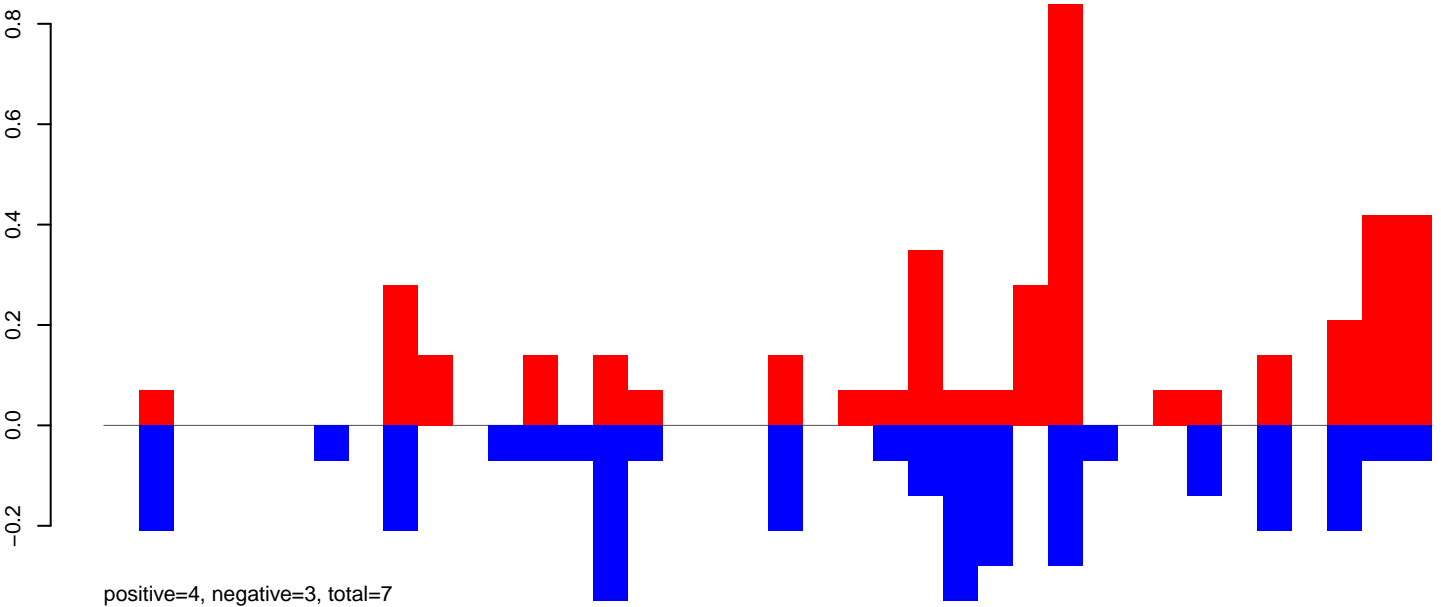
CuQuin_Mcarc_NL.24_35.rep



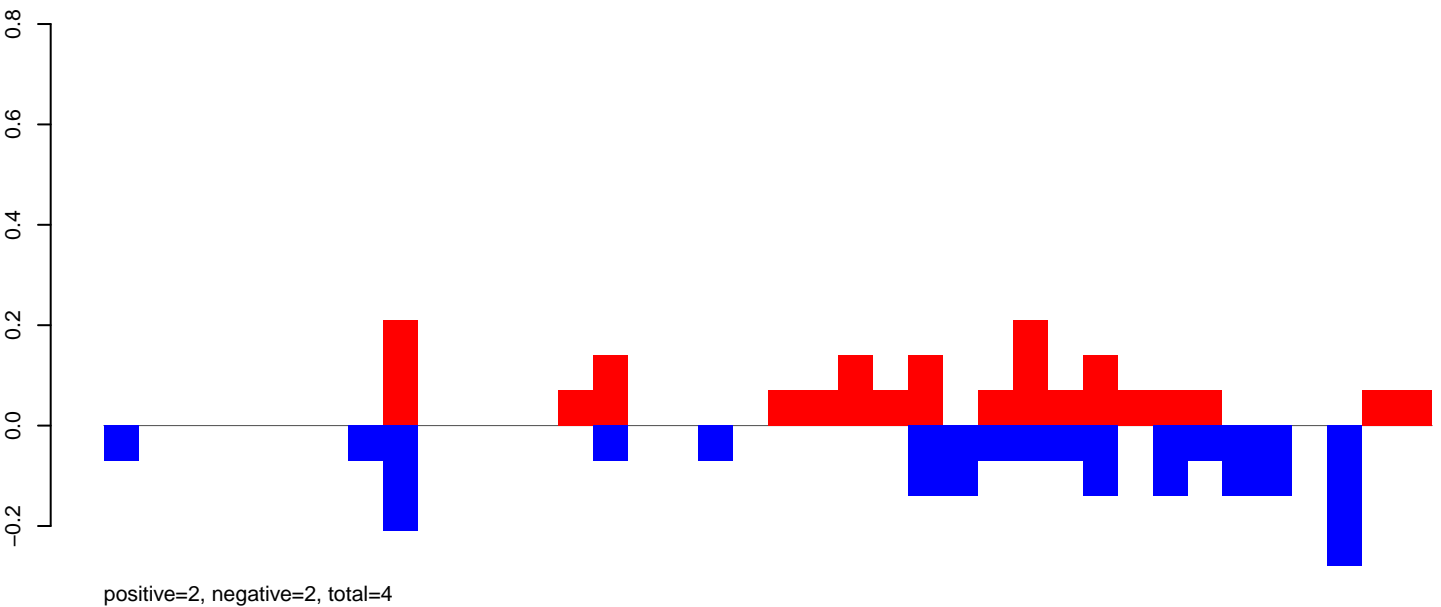
CuQuin_Mcarc_NL.rep



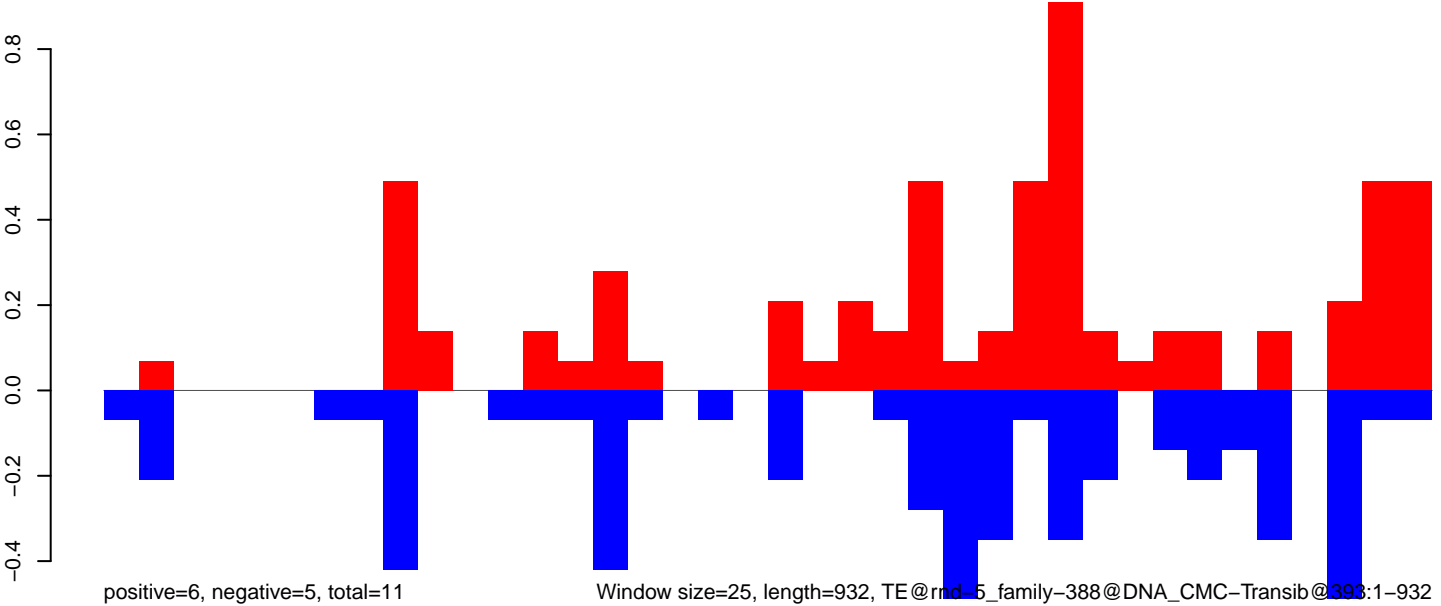
CuQuin_Mcarc_NL.18_23.rep



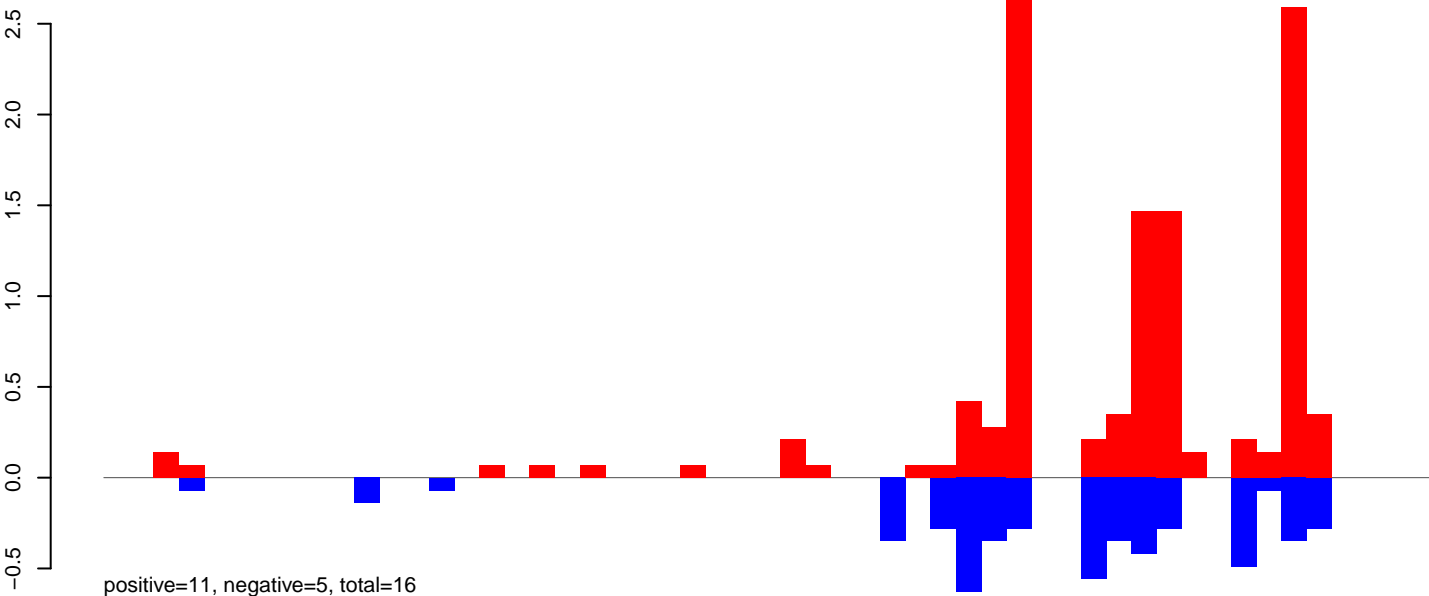
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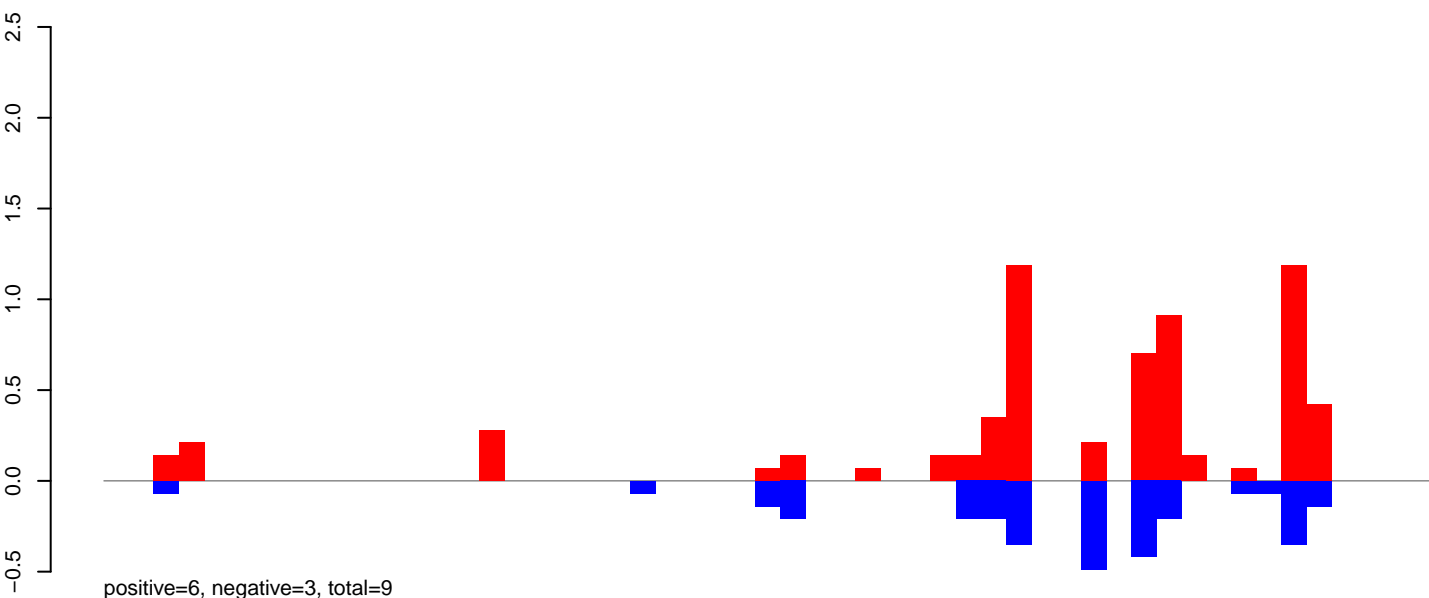
CuQuin_Mcarc_NL.rep



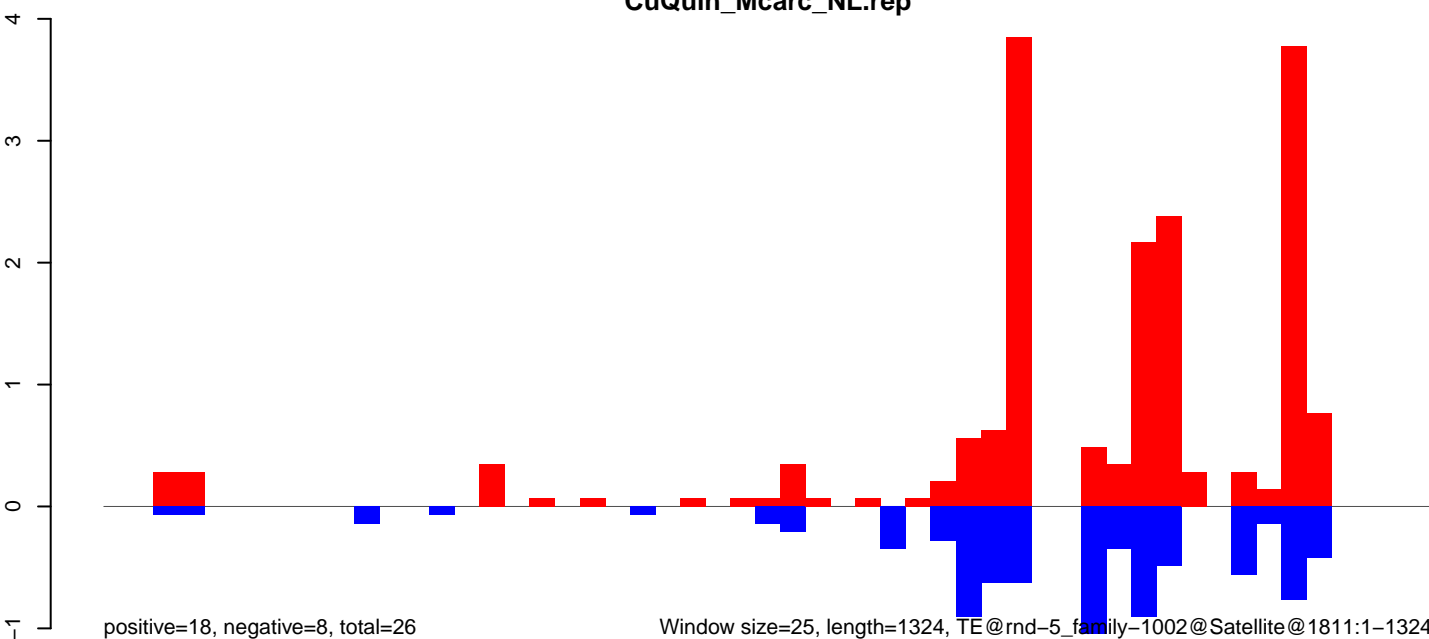
CuQuin_Mcarc_NL.18_23.rep



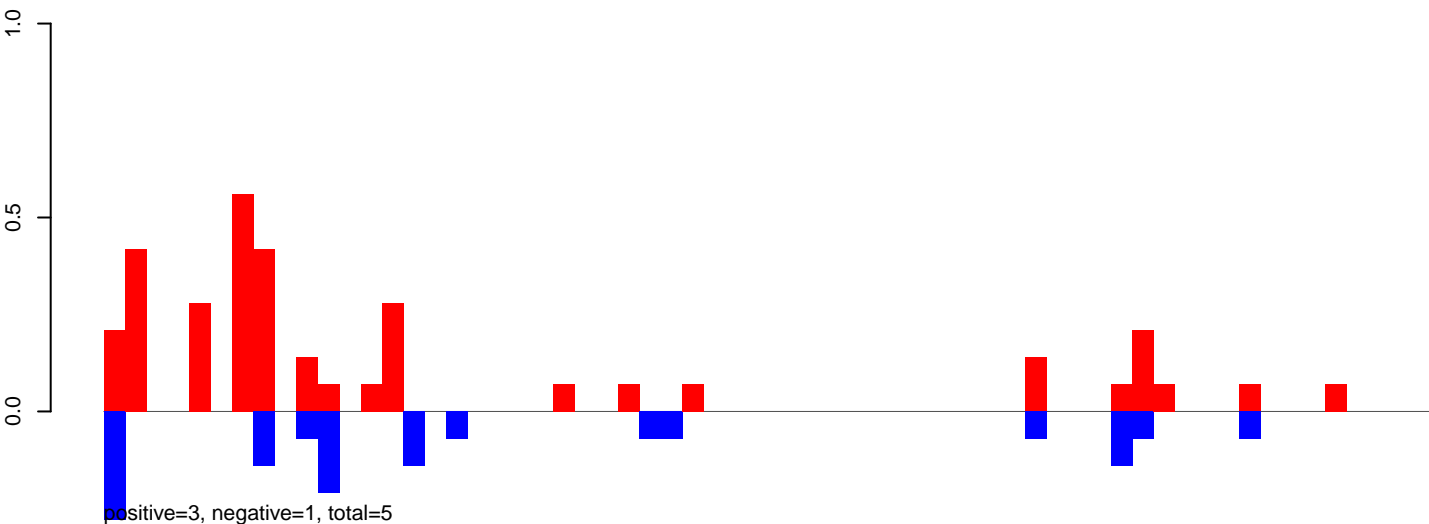
CuQuin_Mcarc_NL.24_35.rep



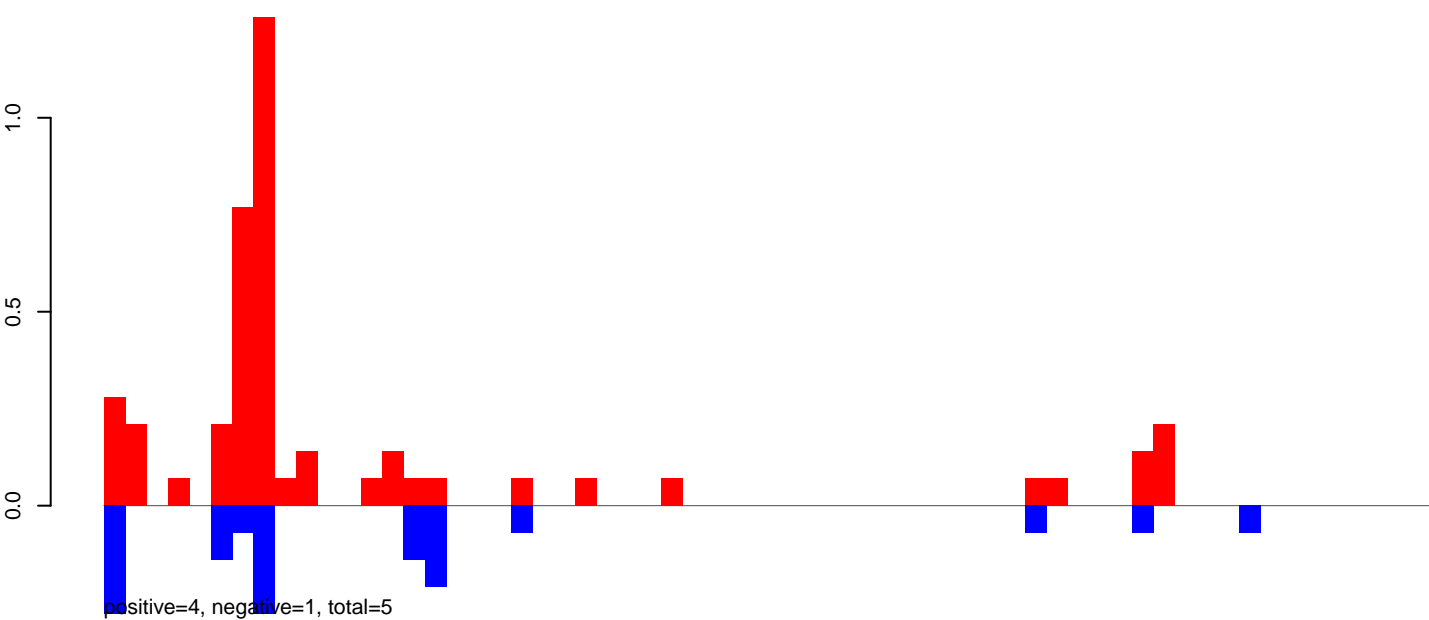
CuQuin_Mcarc_NL.rep



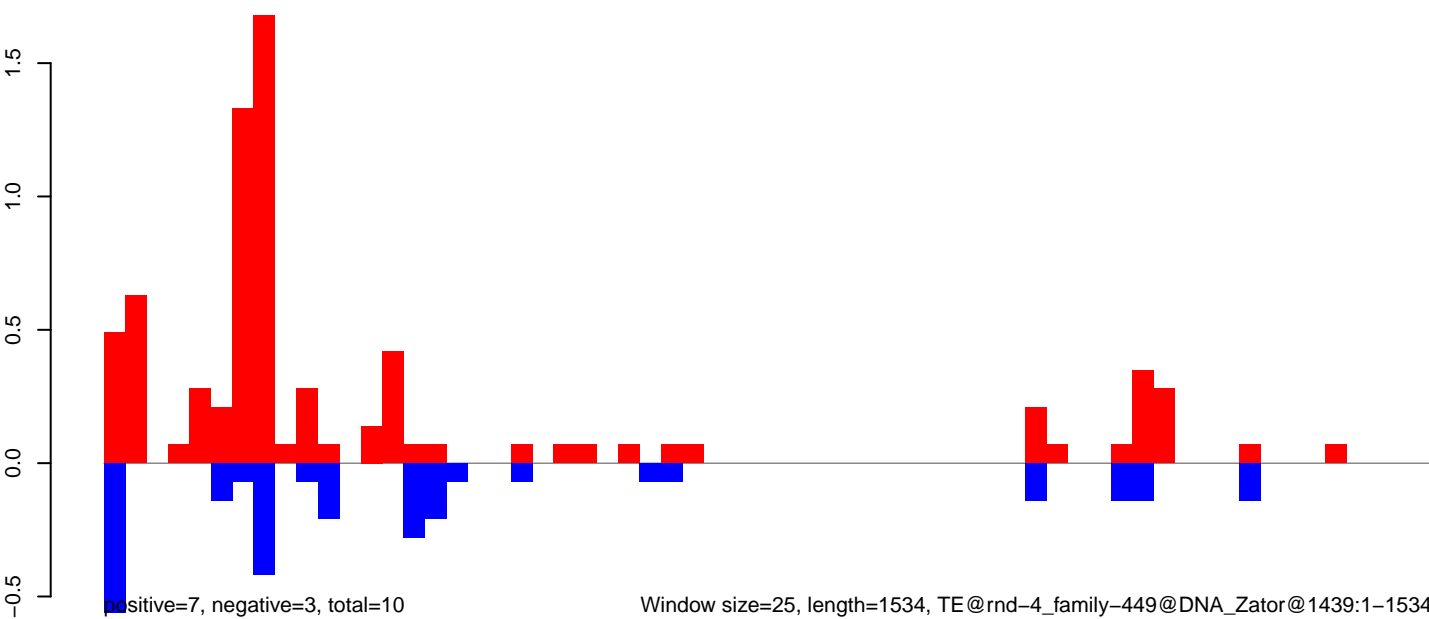
CuQuin_Mcarc_NL.18_23.rep



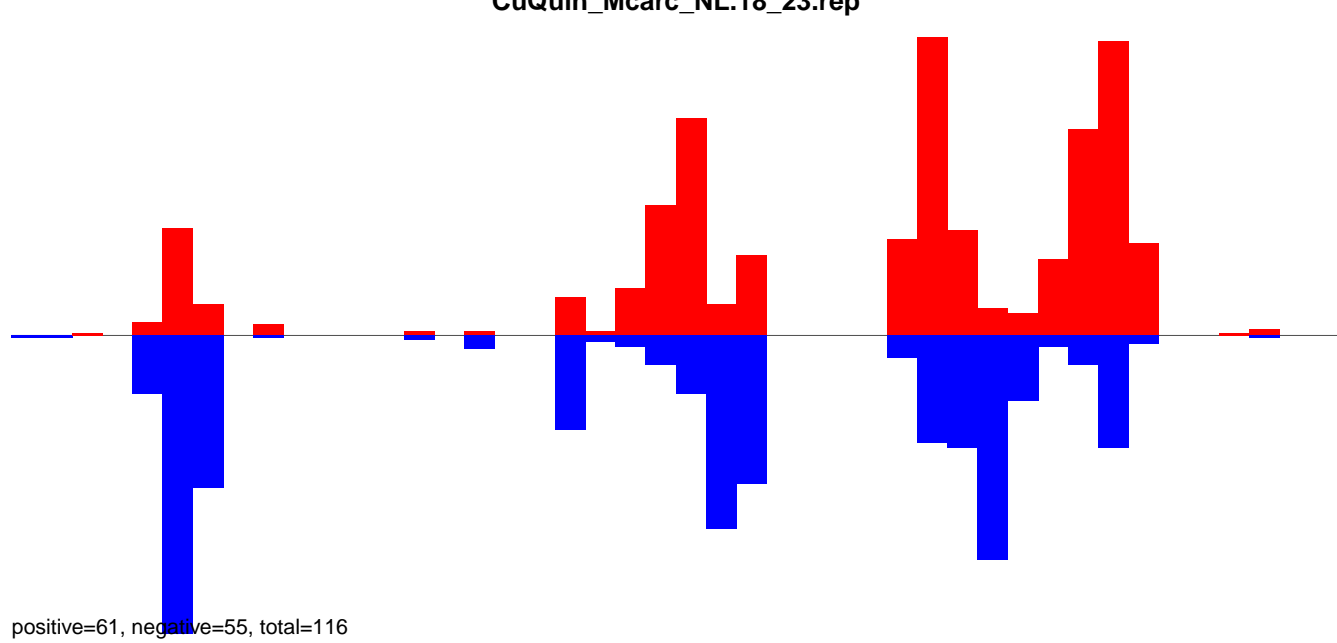
CuQuin_Mcarc_NL.24_35.rep



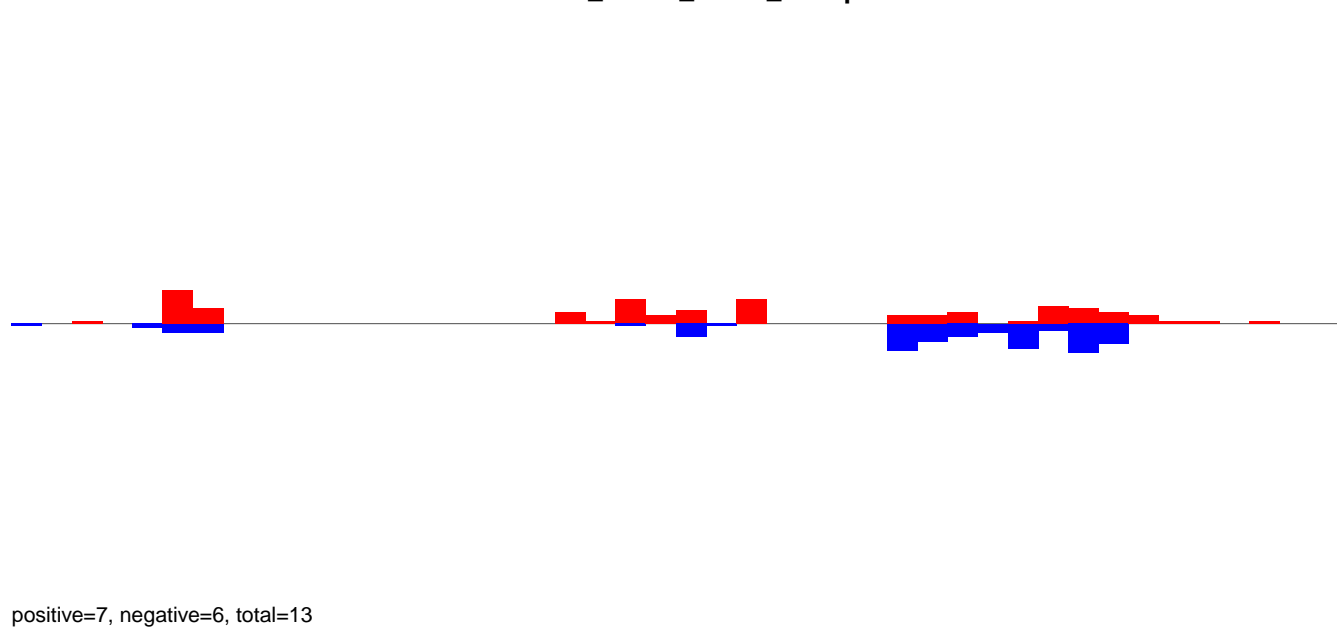
CuQuin_Mcarc_NL.rep



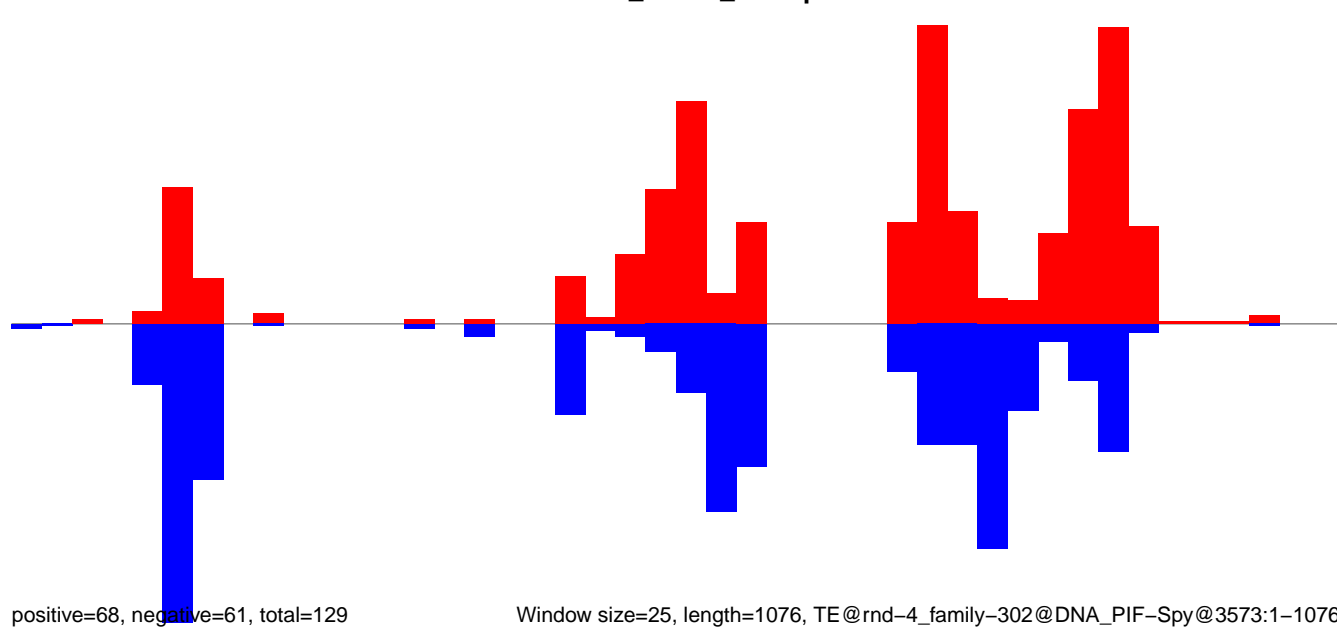
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

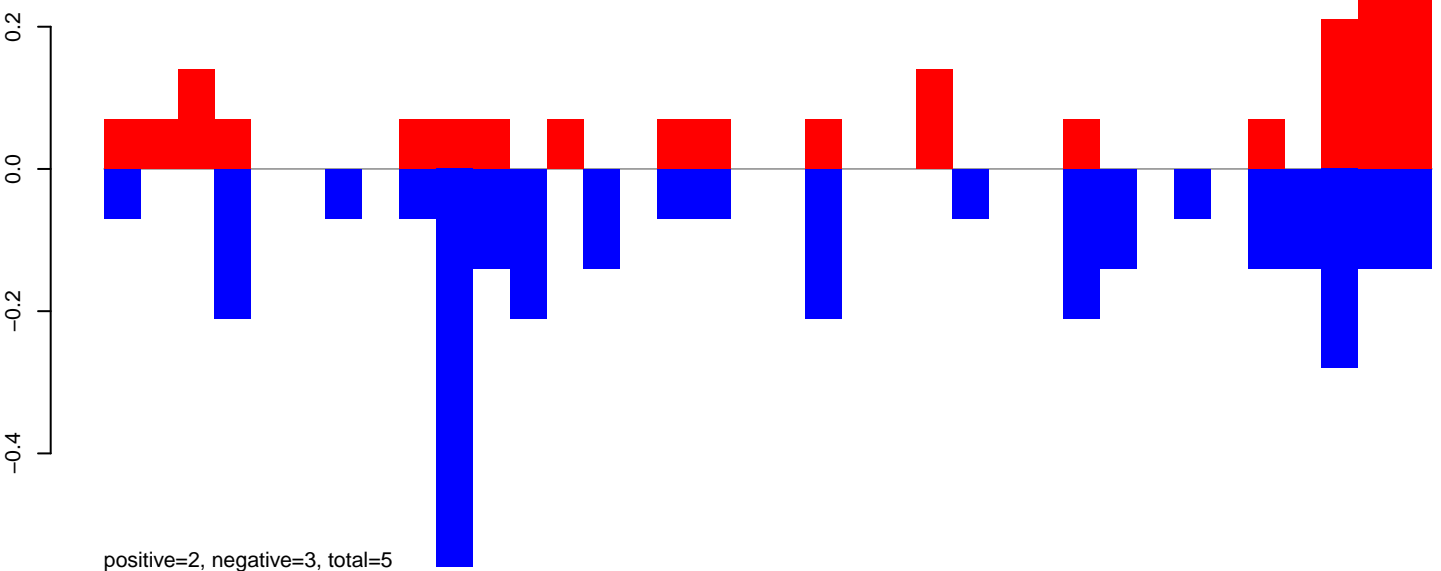


CuQuin_Mcarc_NL.rep

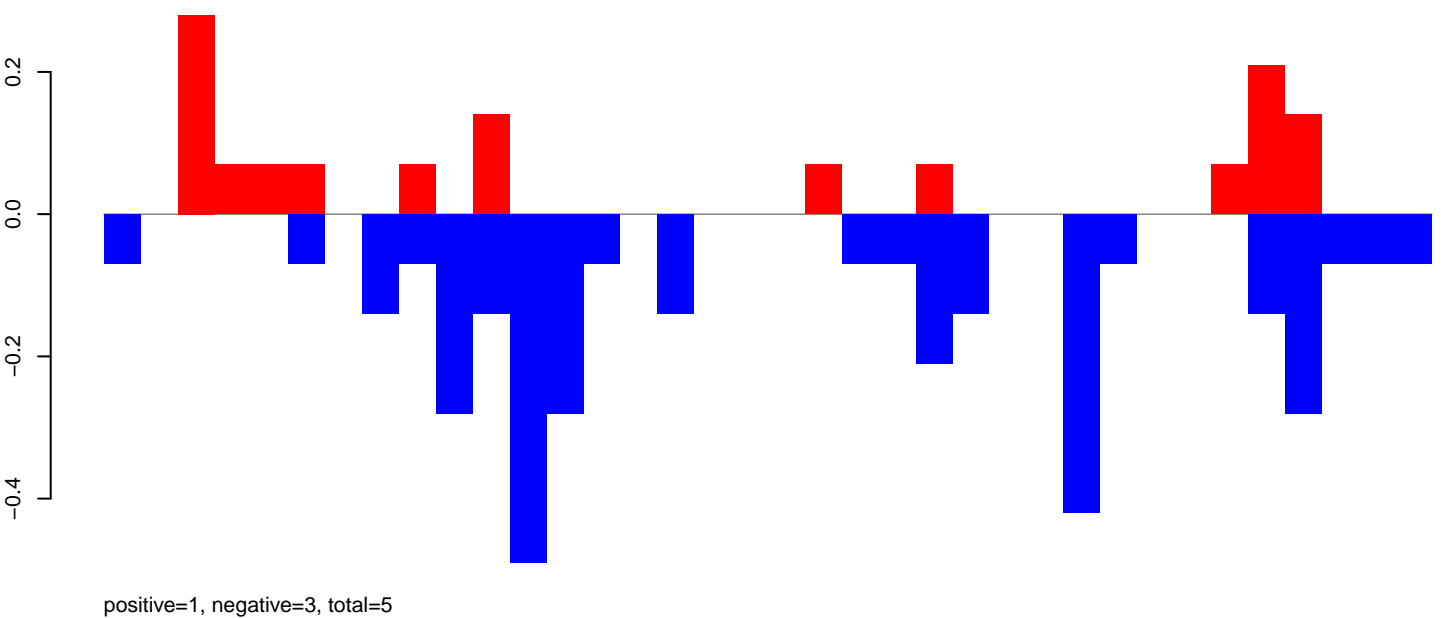


Window size=25, length=1076, TE@rnd-4_family-302@DNA_PIF-Spy@3573:1-1076

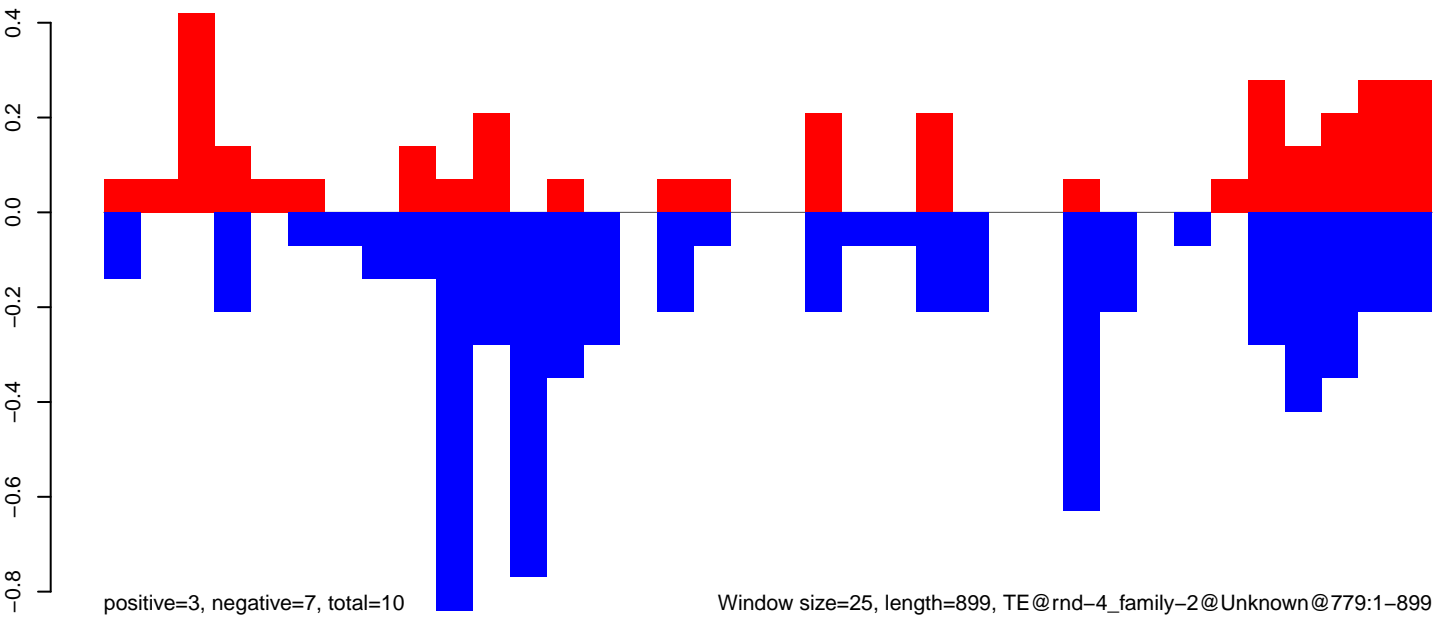
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

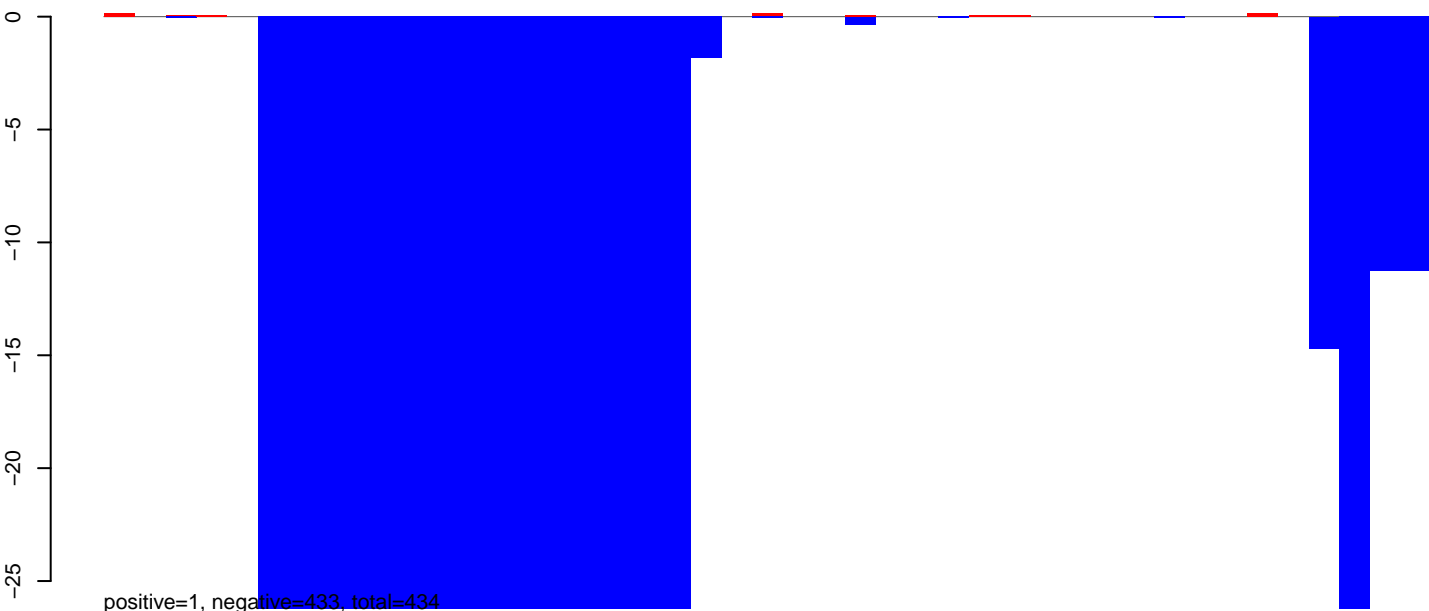


CuQuin_Mcarc_NL.rep

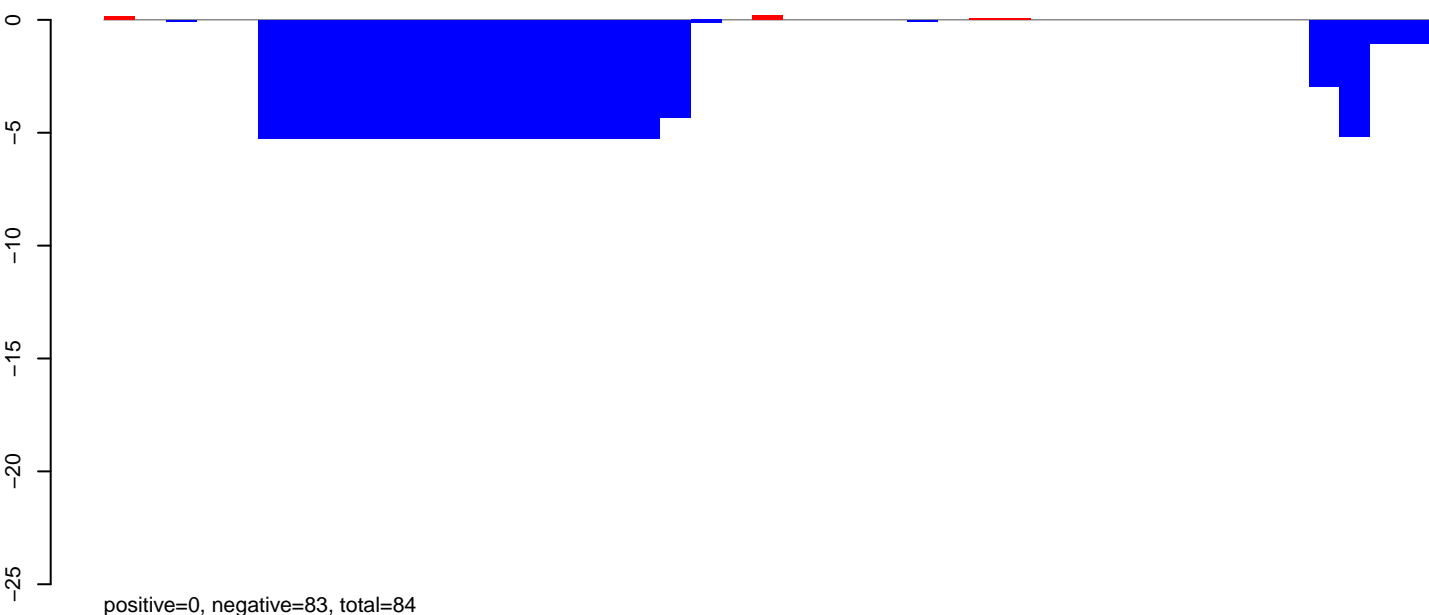


Window size=25, length=899, TE@rnd-4_family-2@Unknown@779:1-899

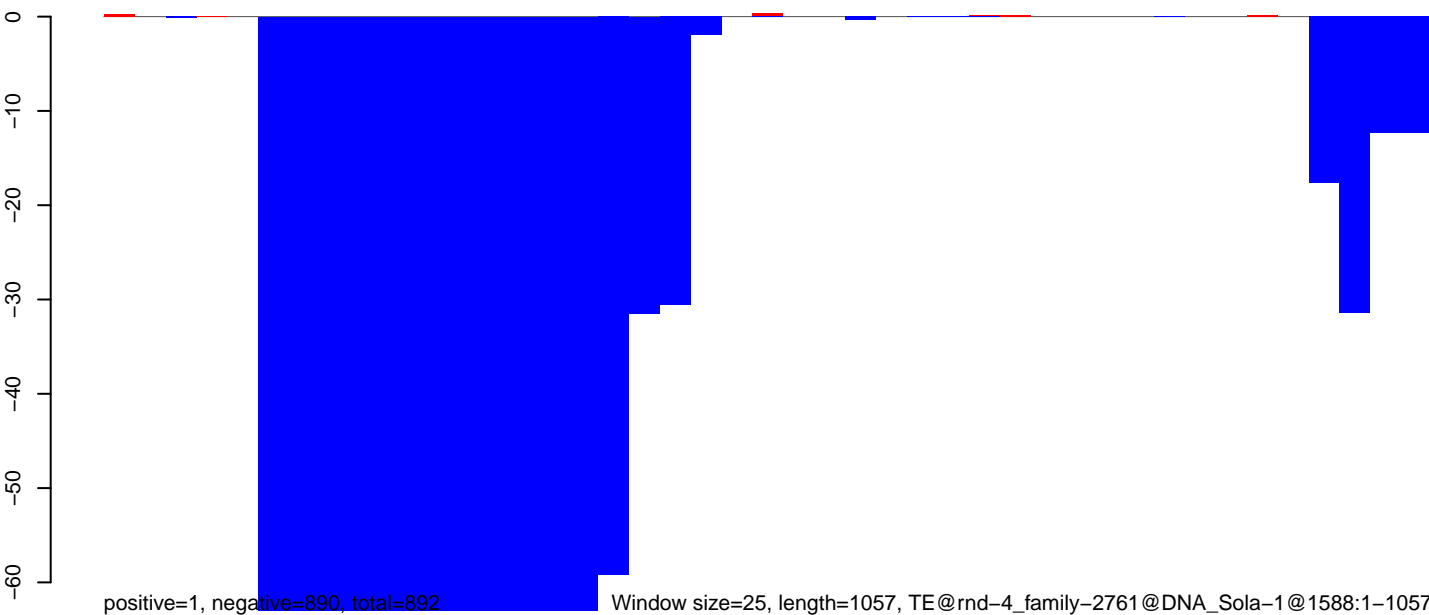
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

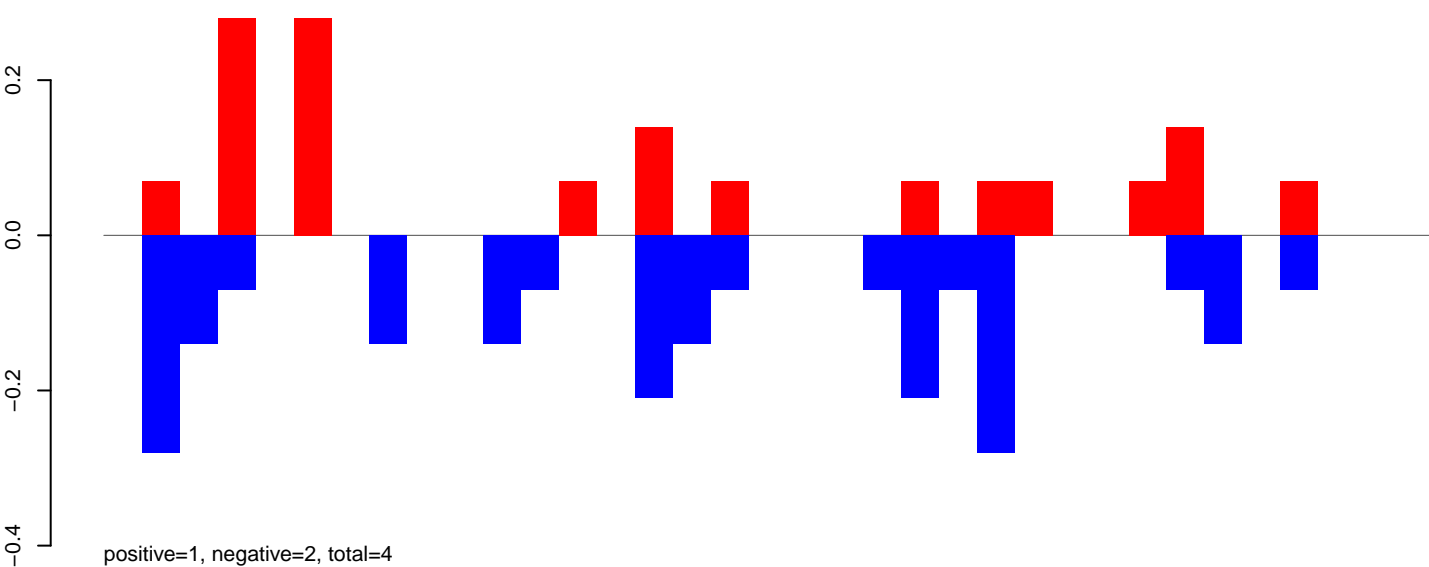


CuQuin_Mcarc_NL.rep

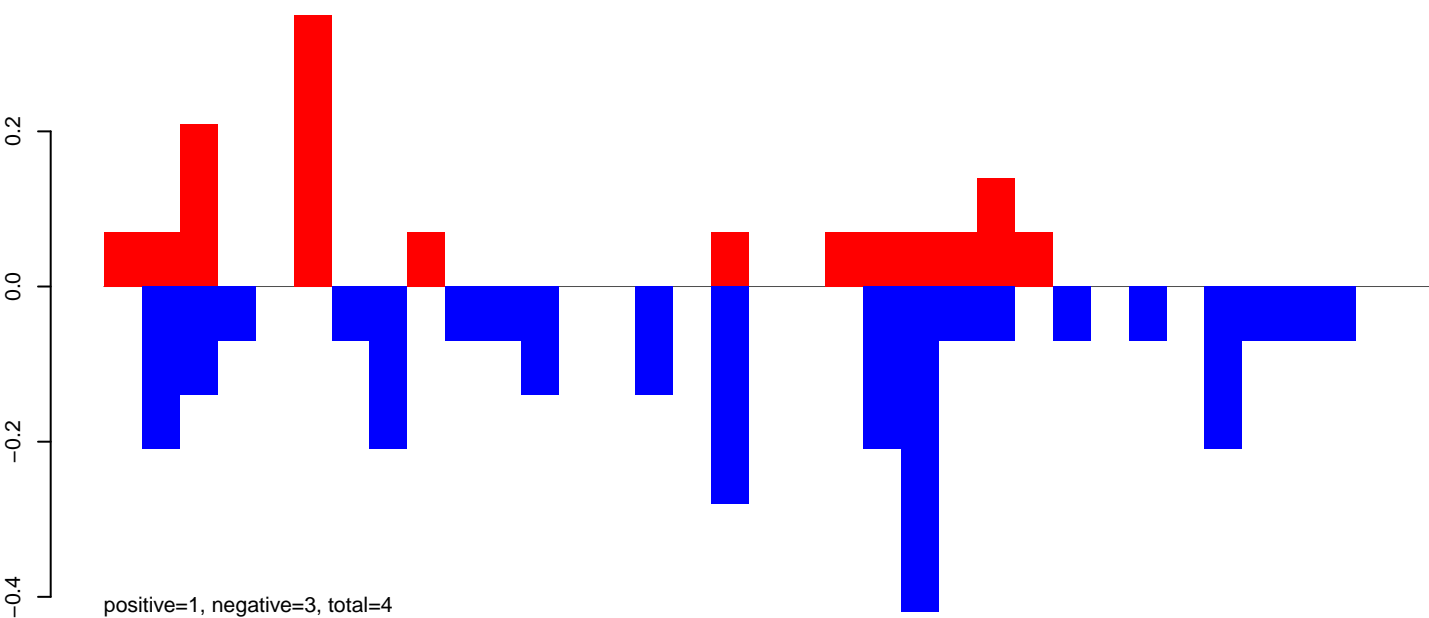


Window size=25, length=1057, TE@rnd-4_family-2761@DNA_Sola-1@1588:1-1057

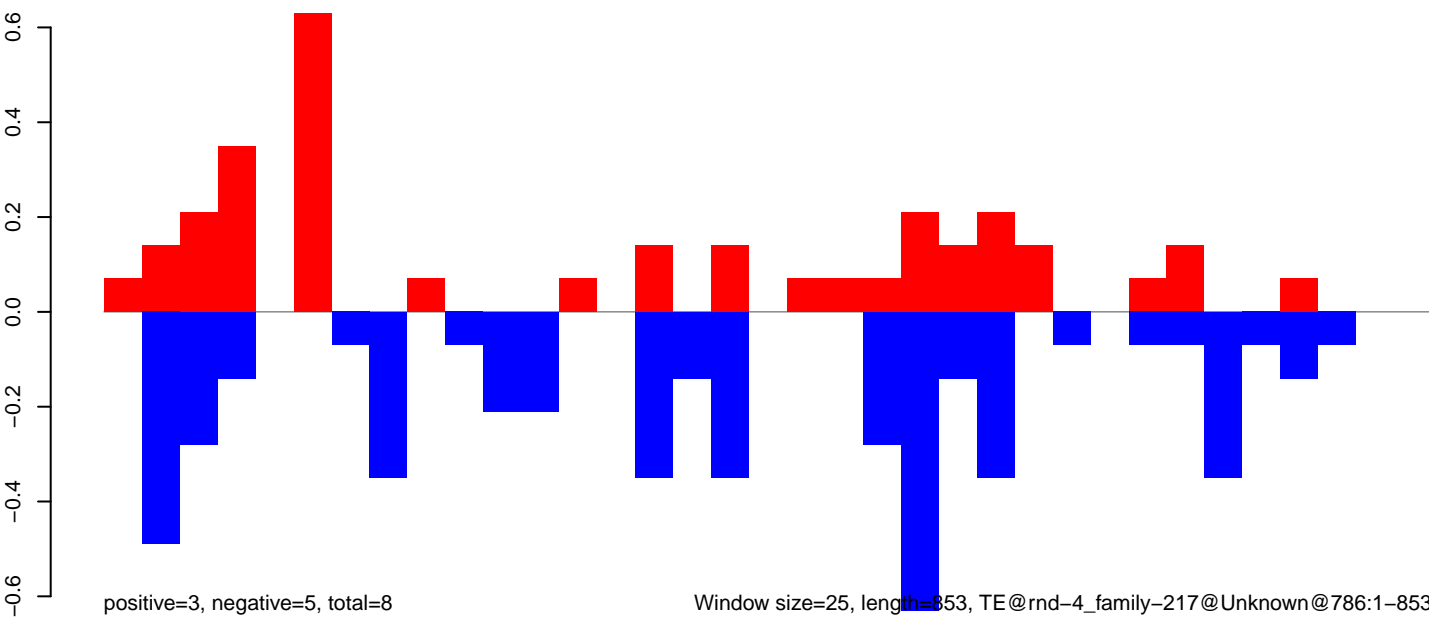
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

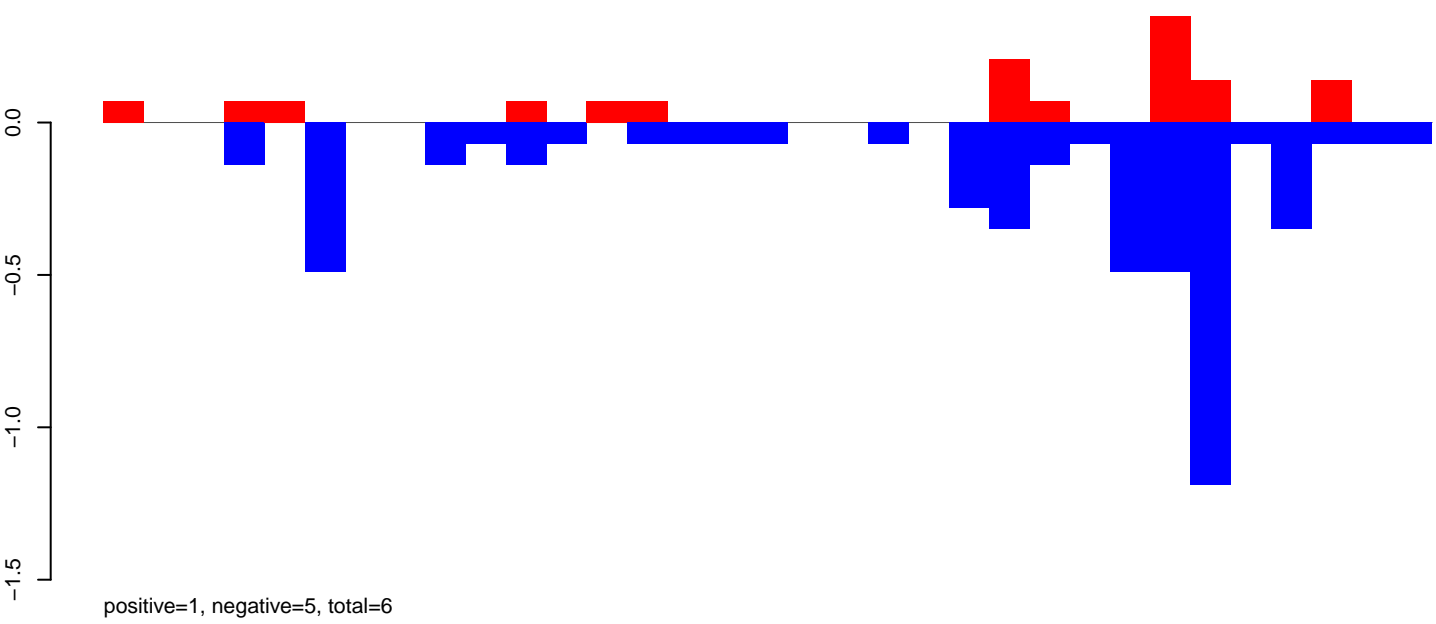


CuQuin_Mcarc_NL.rep

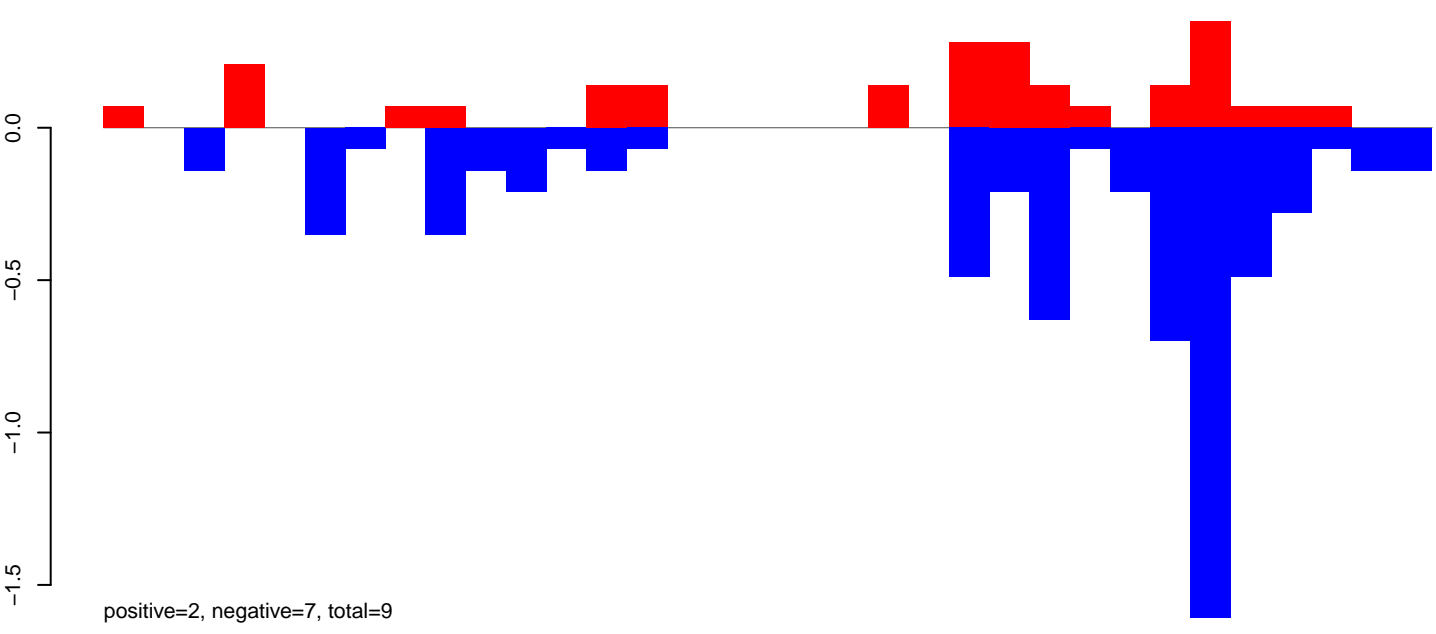


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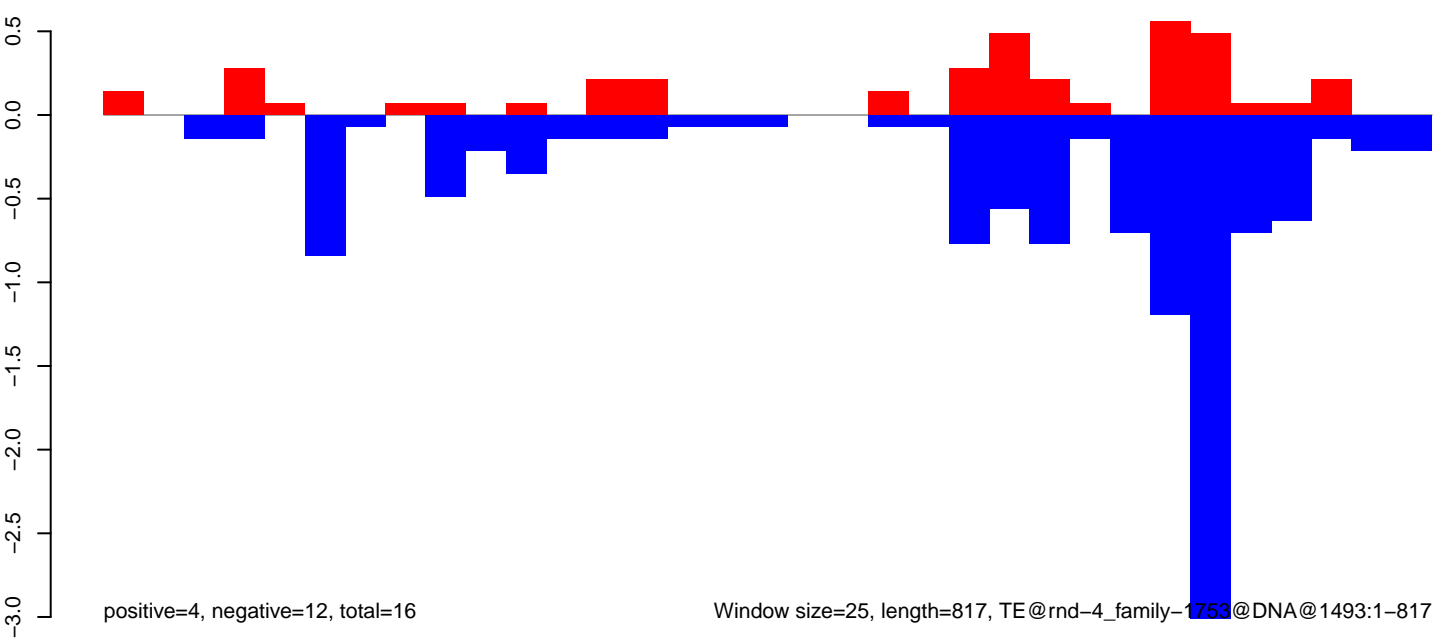
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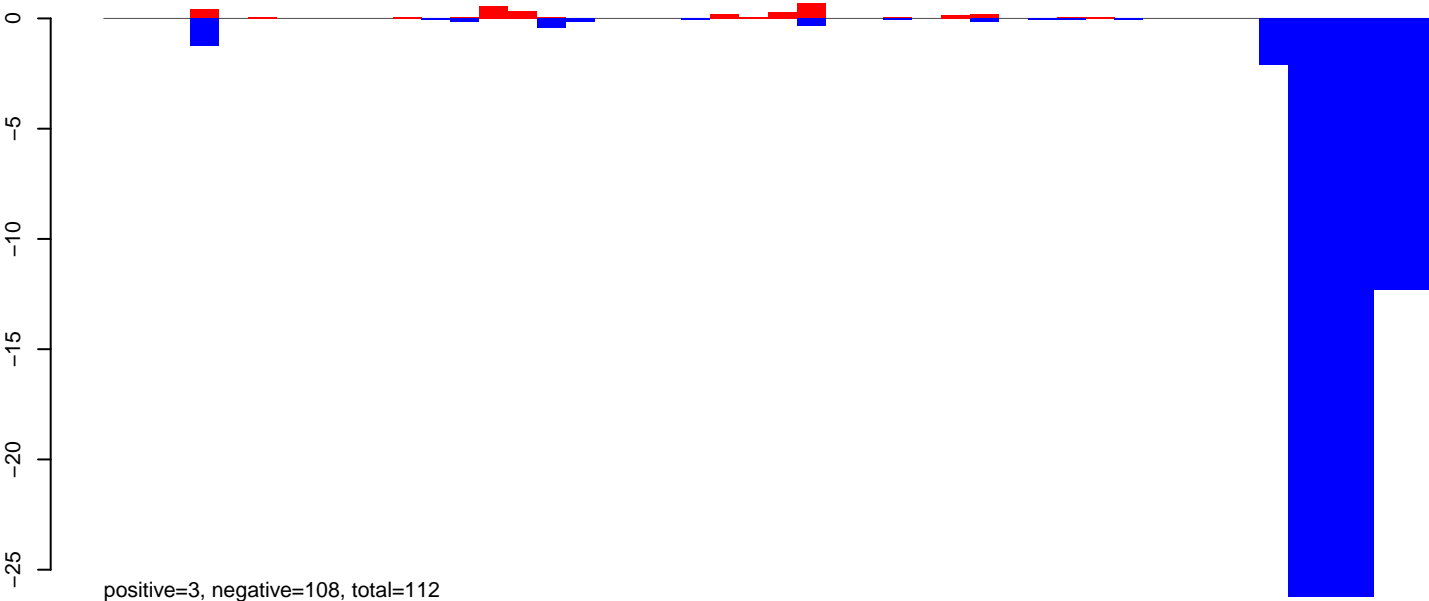
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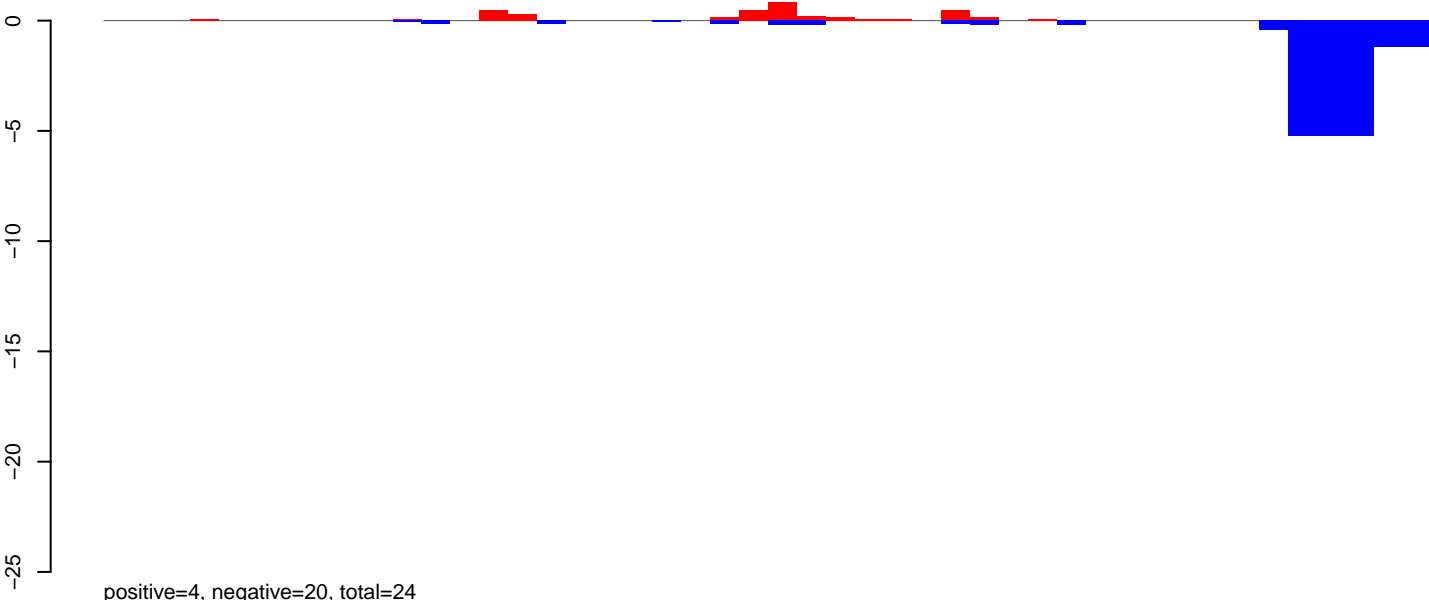
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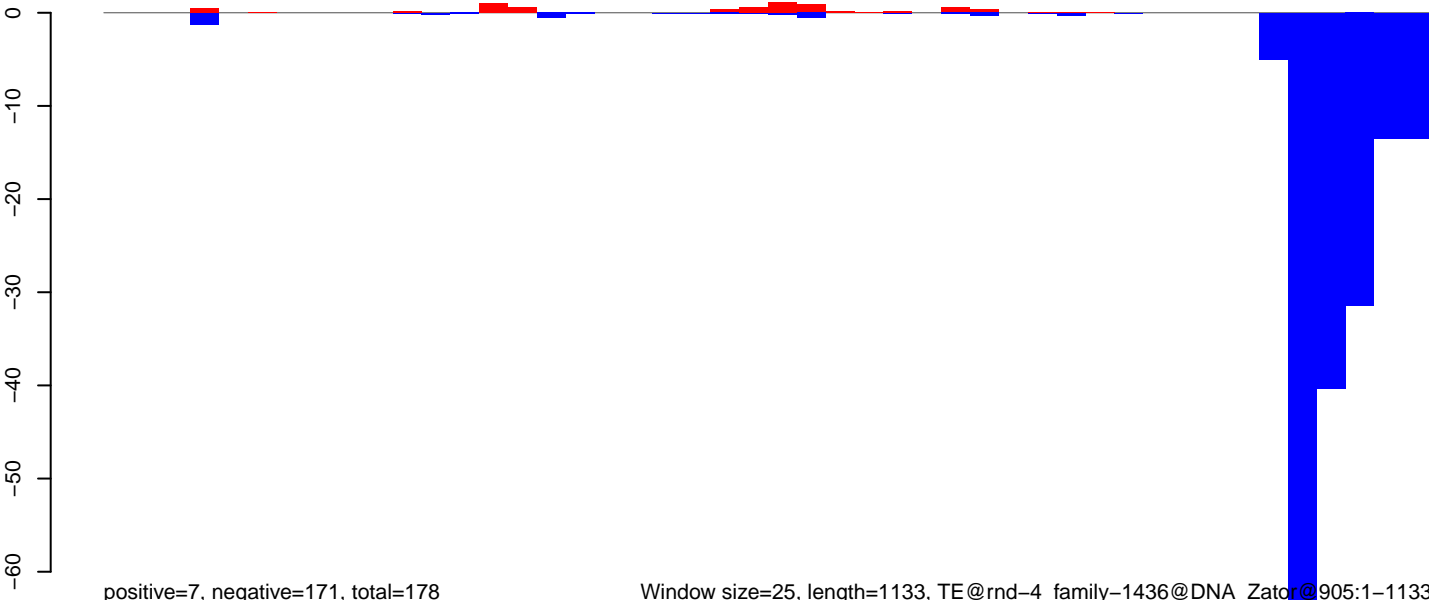
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CuQuin_Mcarc_NL.24_35.rep



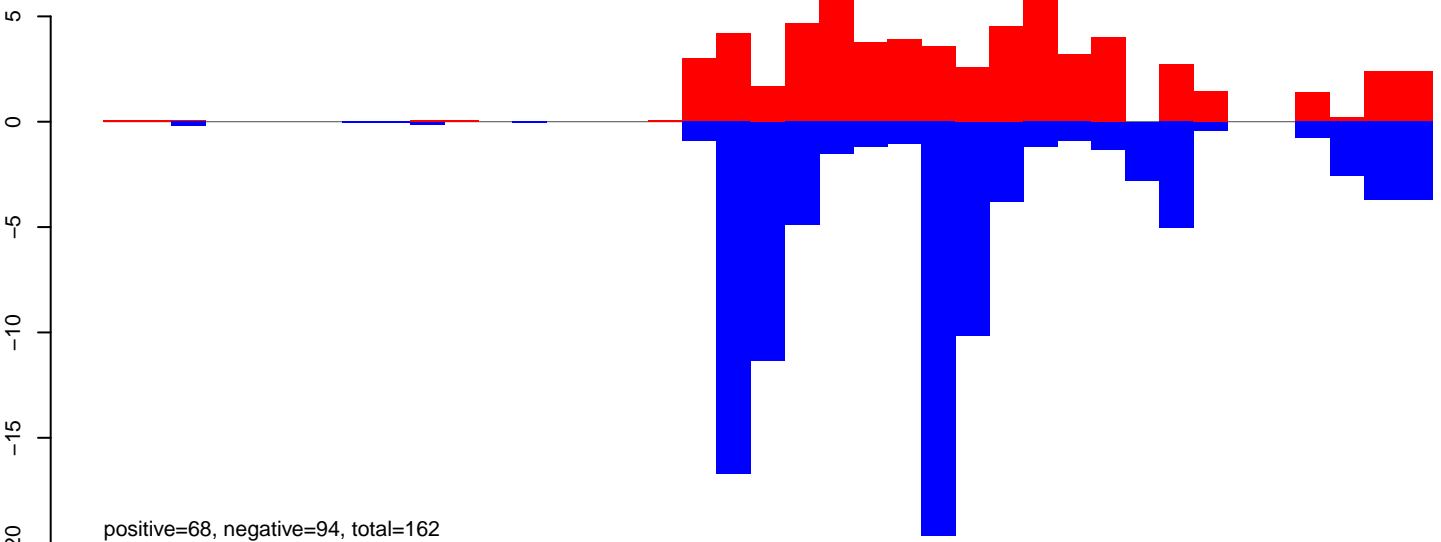
CuQuin_Mcarc_NL.rep



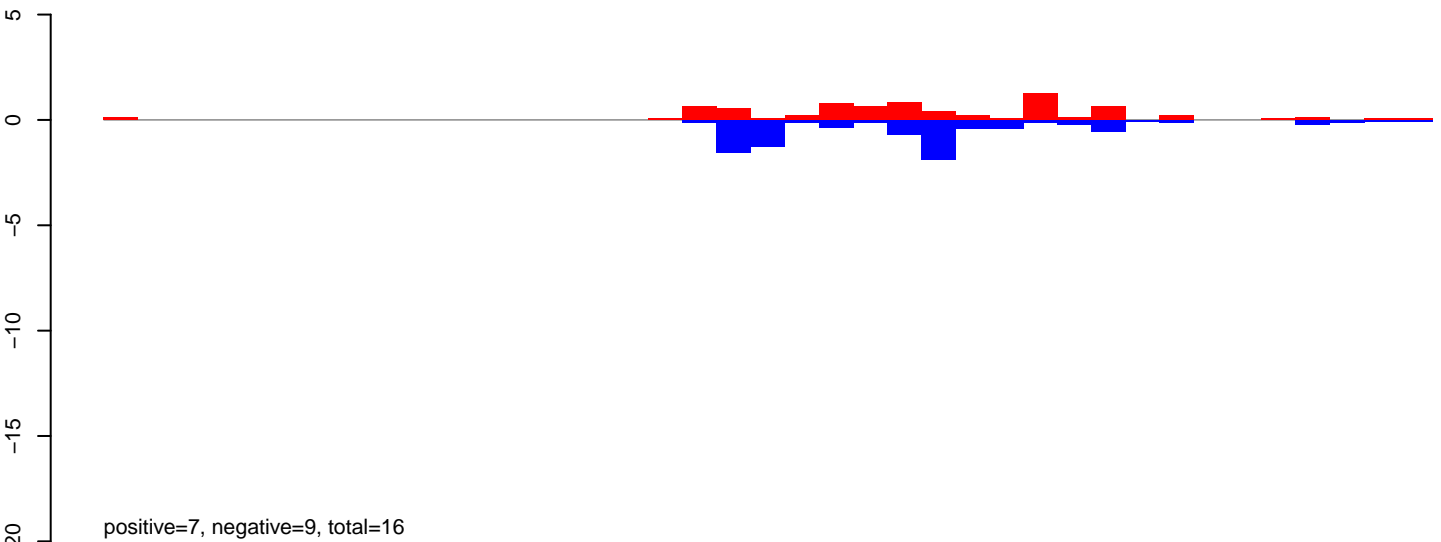
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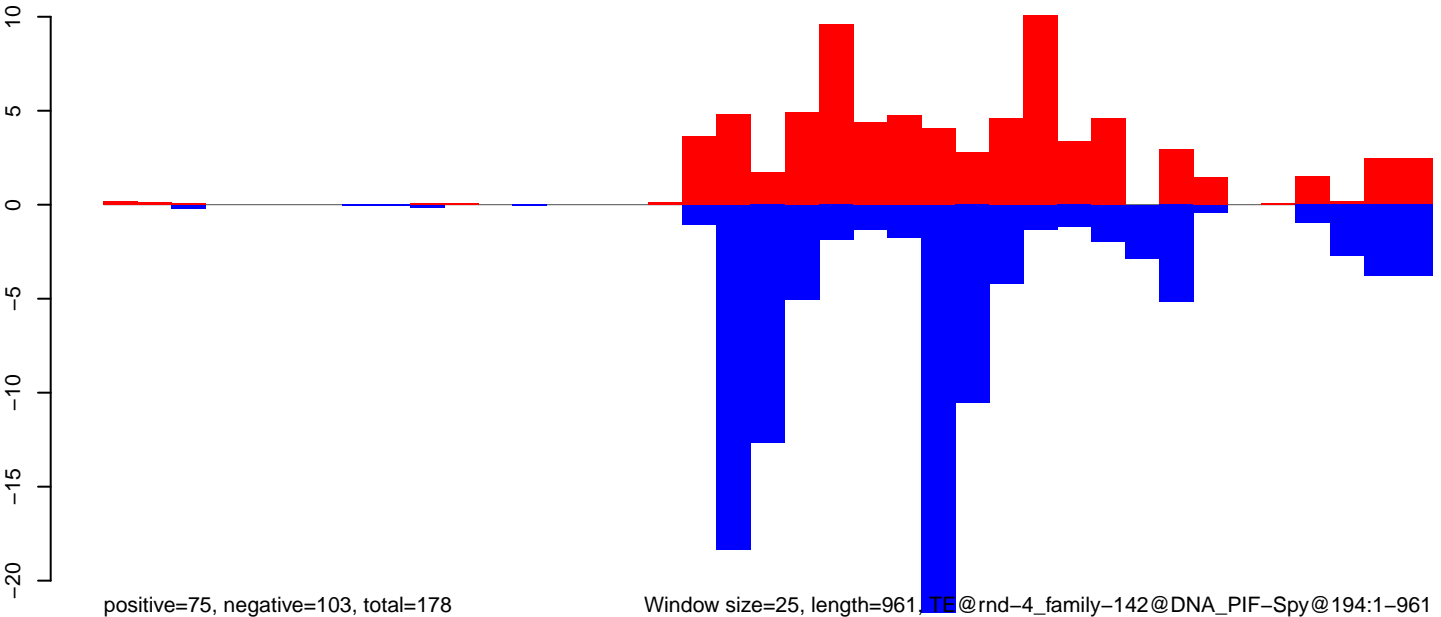
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

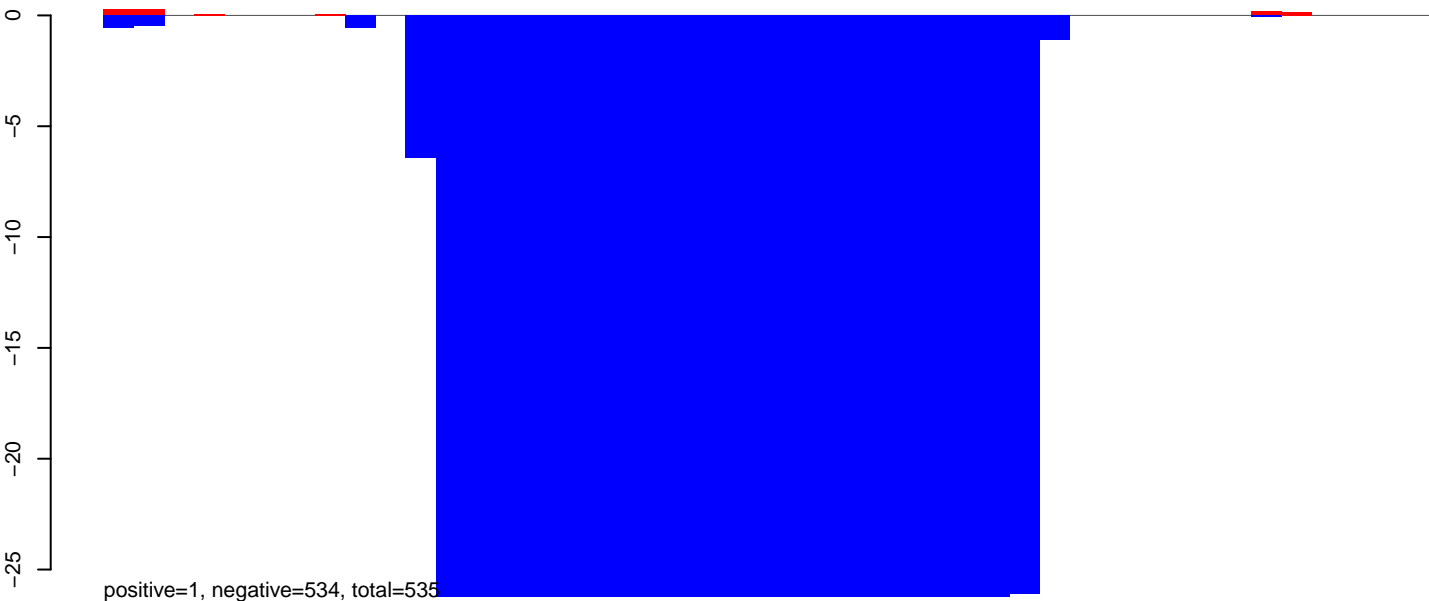


CuQuin_Mcarc_NL.rep

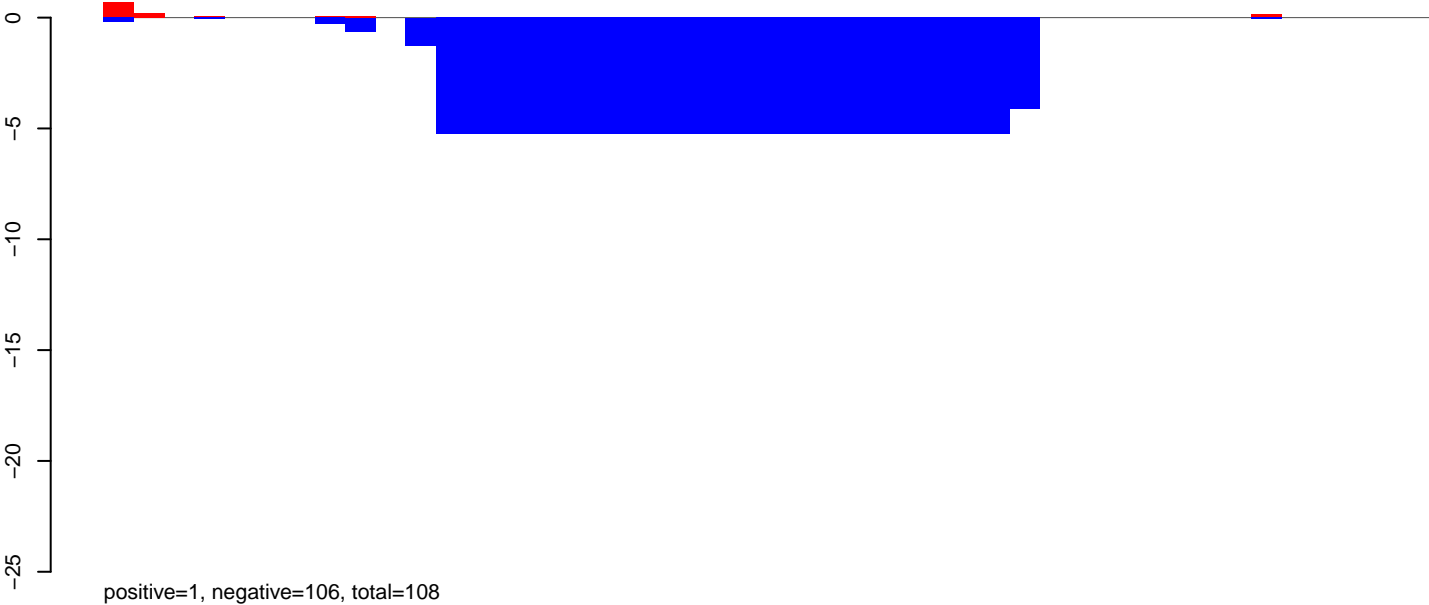


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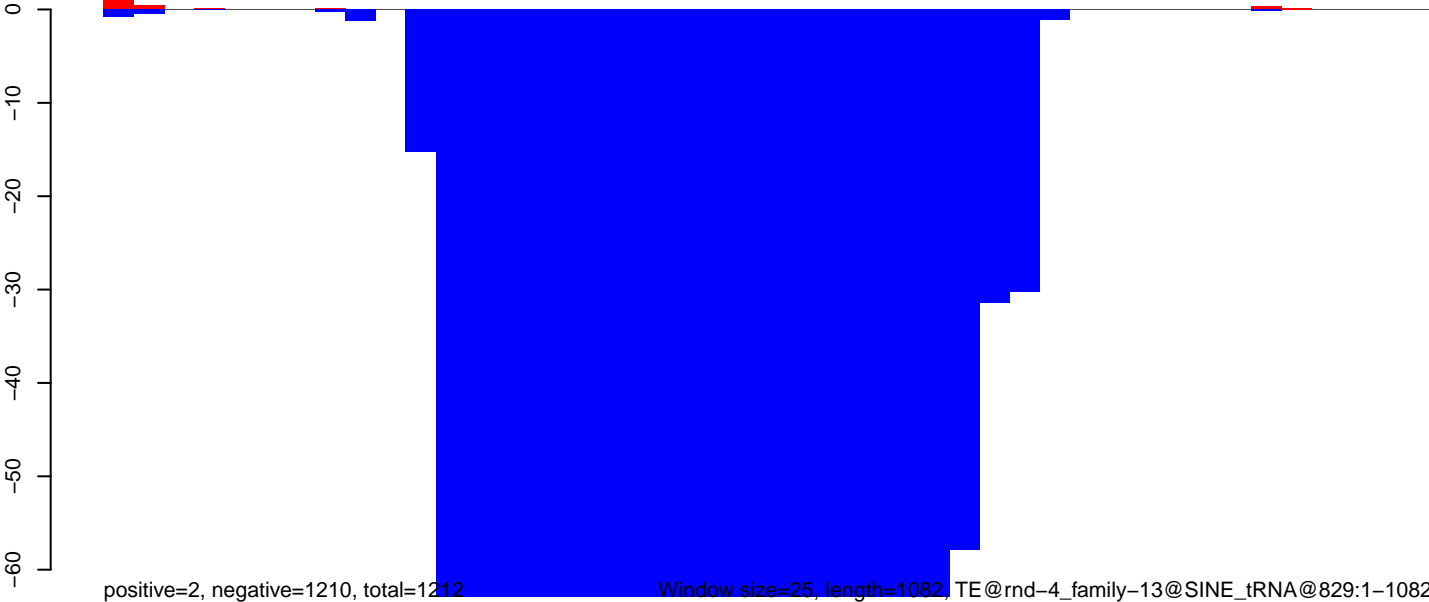
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CuQuin_Mcarc_NL.24_35.rep



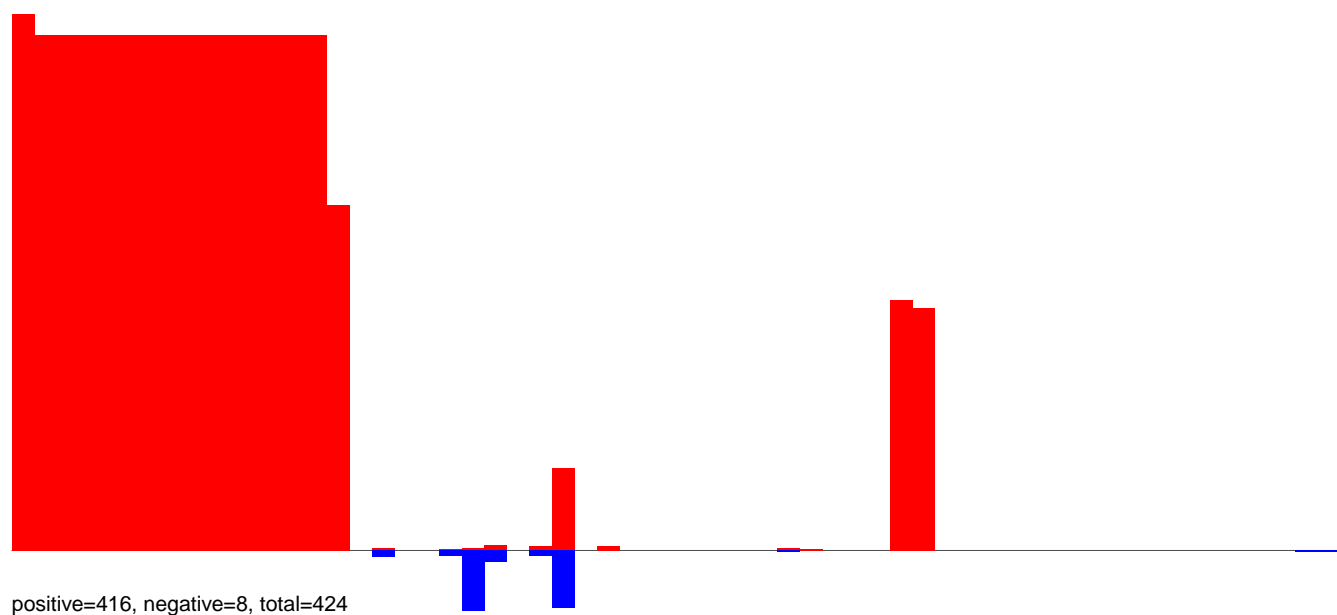
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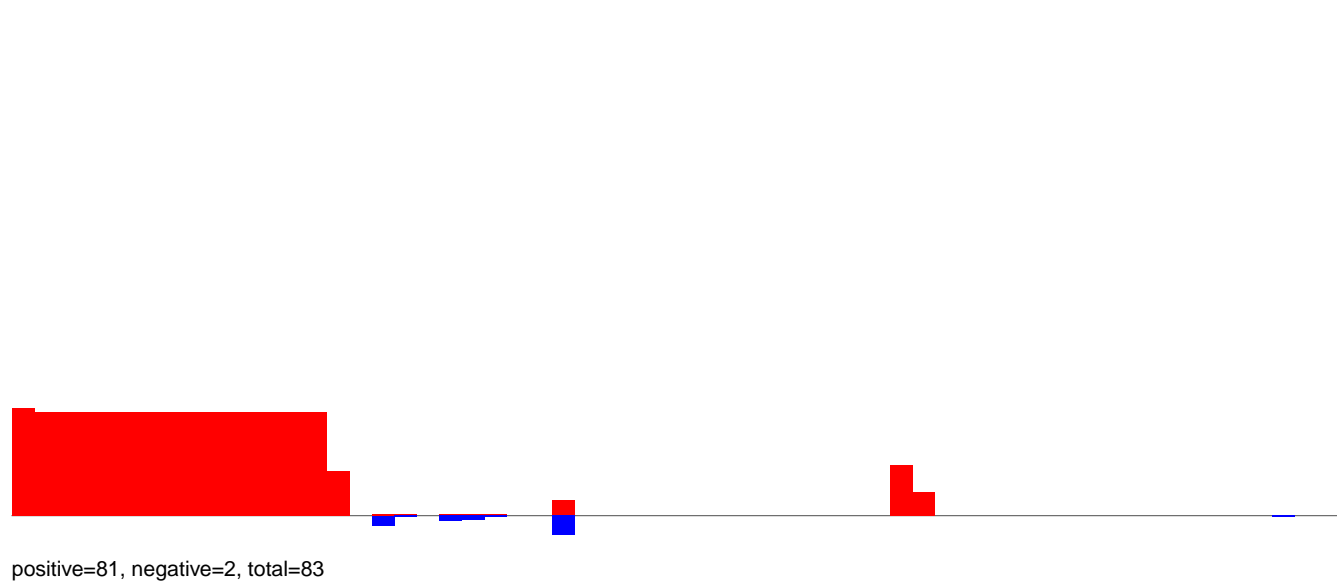
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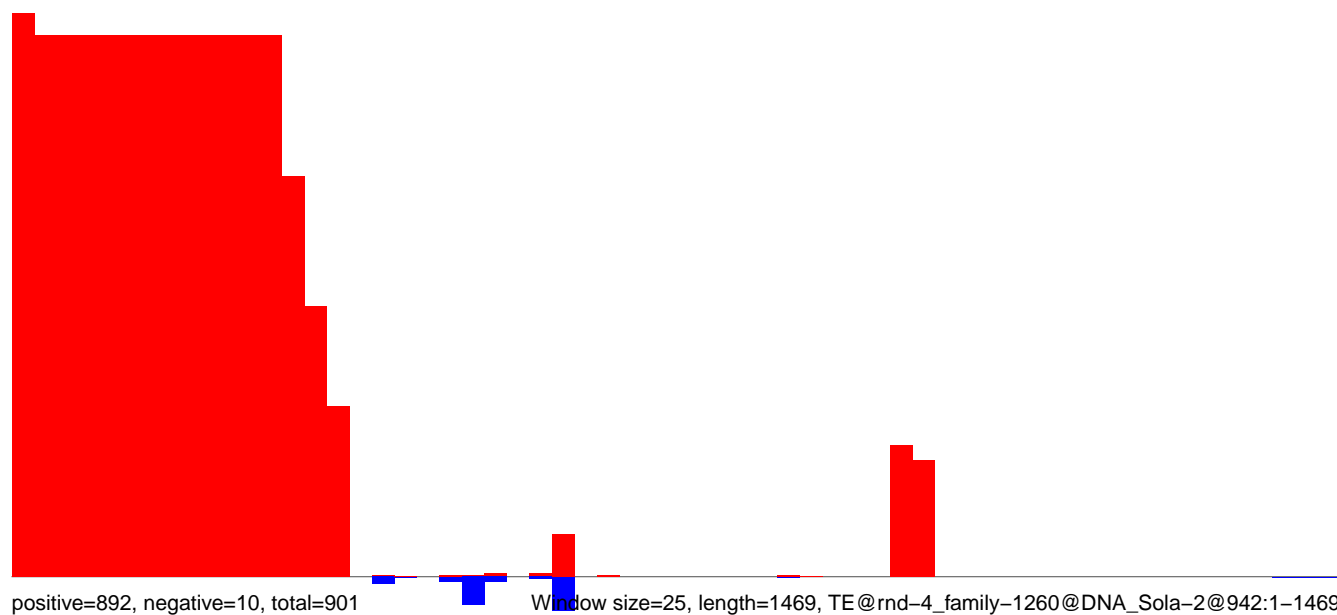
CuQuin_Mcarc_NL.18_23.rep



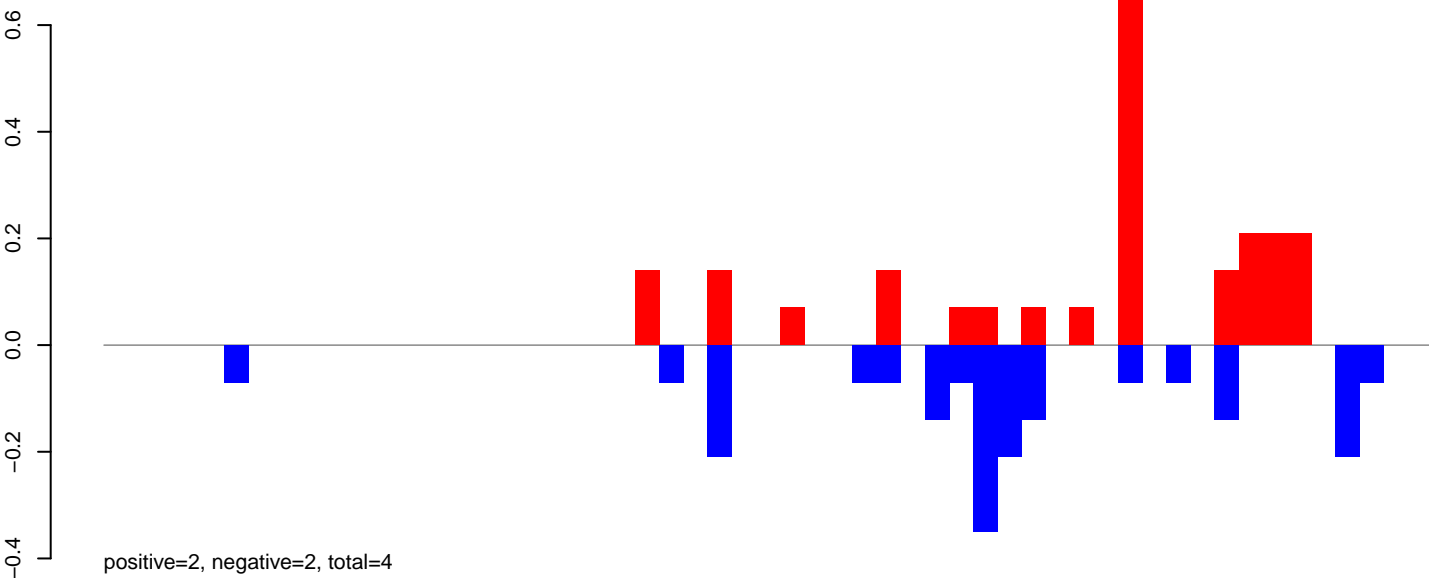
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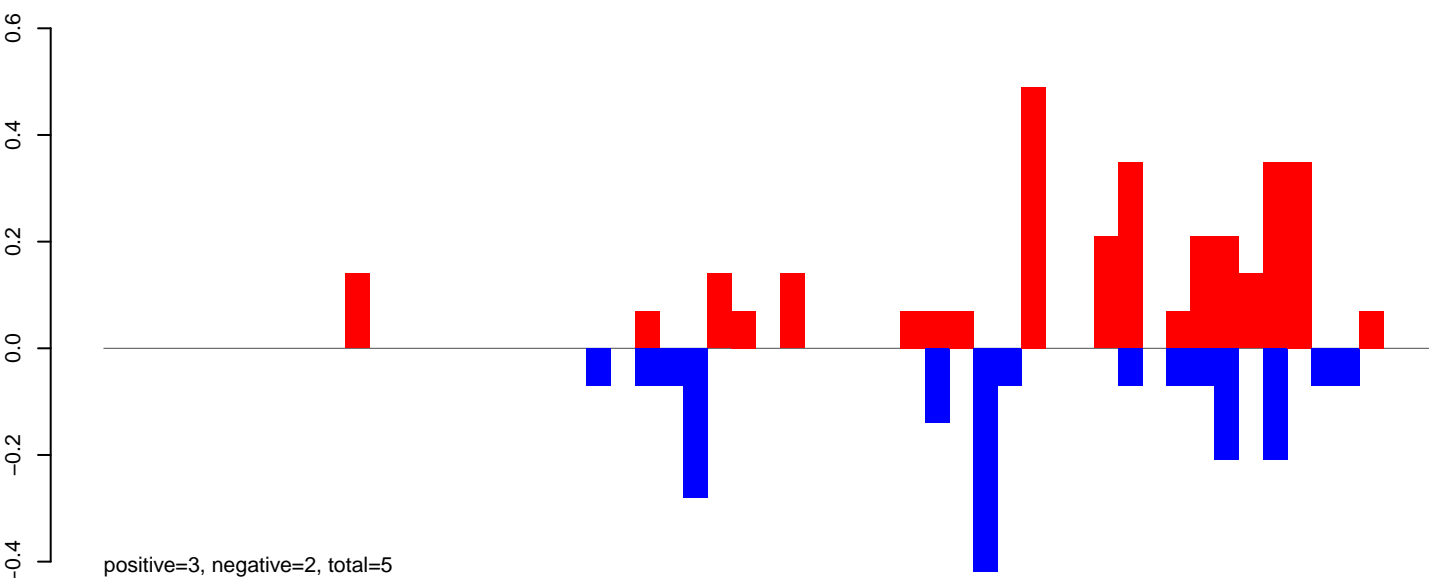
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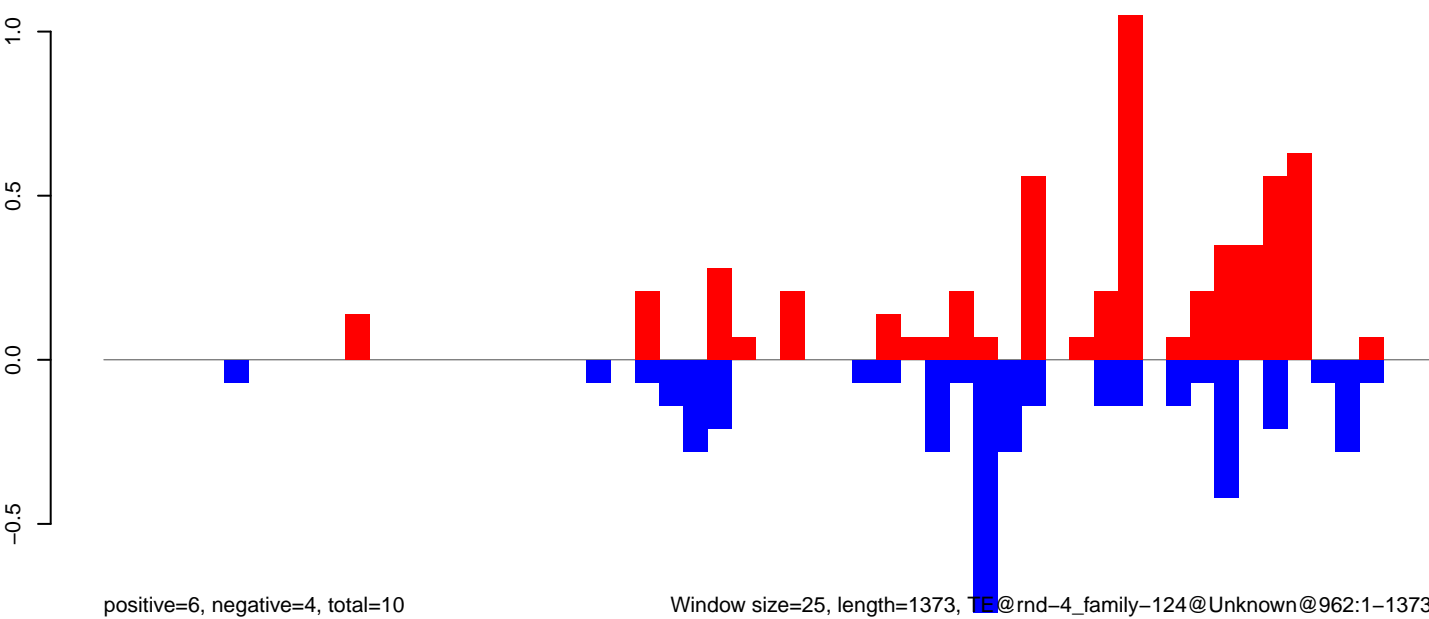
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CuQuin_Mcarc_NL.24_35.rep



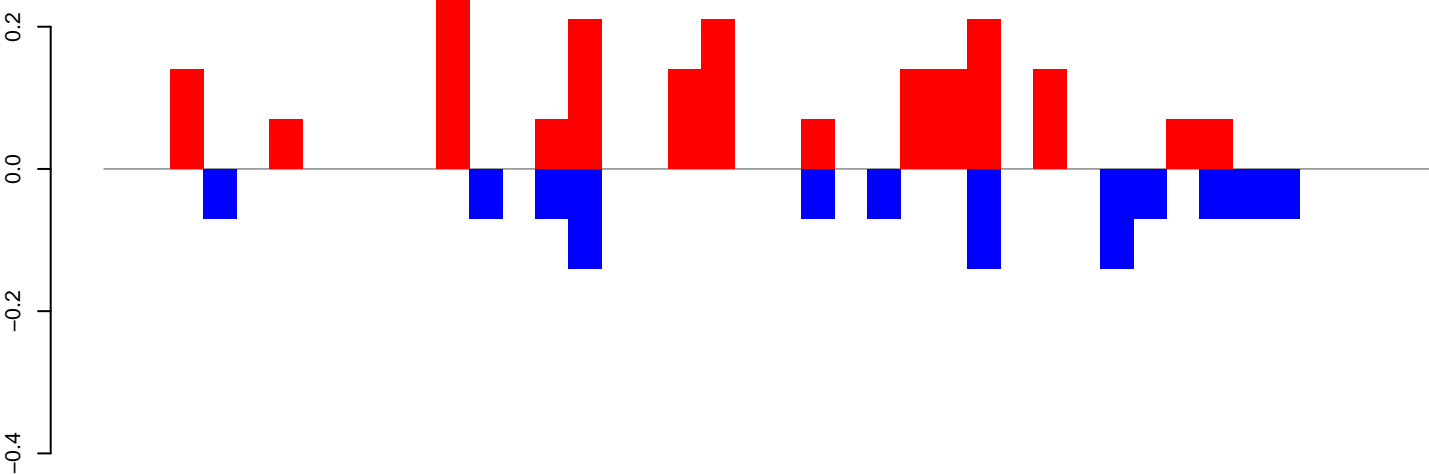
CuQuin_Mcarc_NL.rep



Window size=25, length=1373, TE@rnd-4_family-124@Unknown@962:1-1373

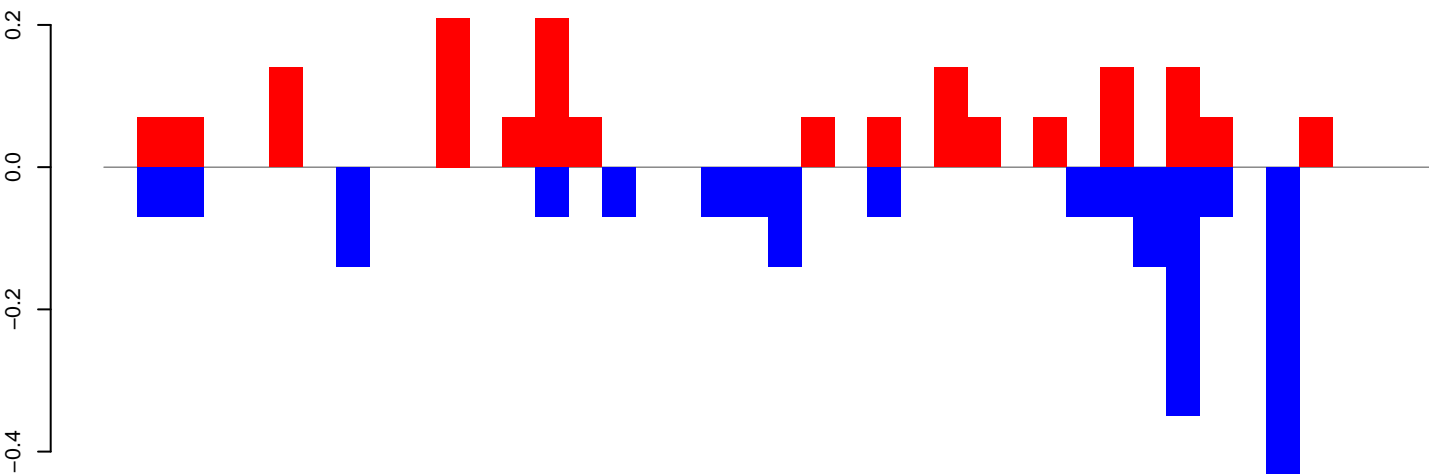
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CuQuin_Mcarc_NL.18_23.rep



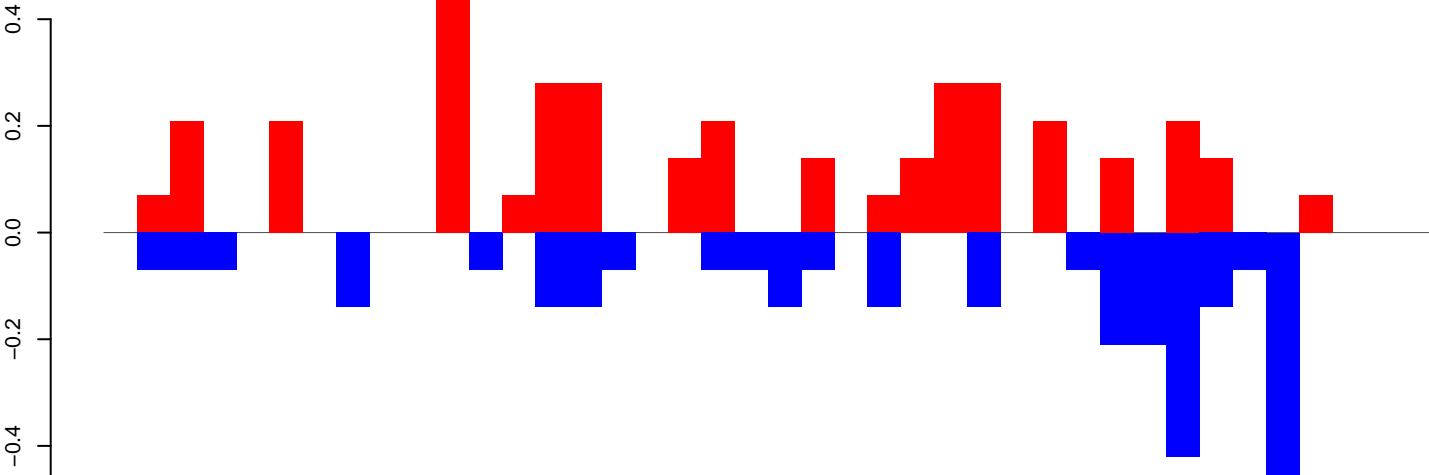
positive=2, negative=1, total=3

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=2, total=4

CuQuin_Mcarc_NL.rep

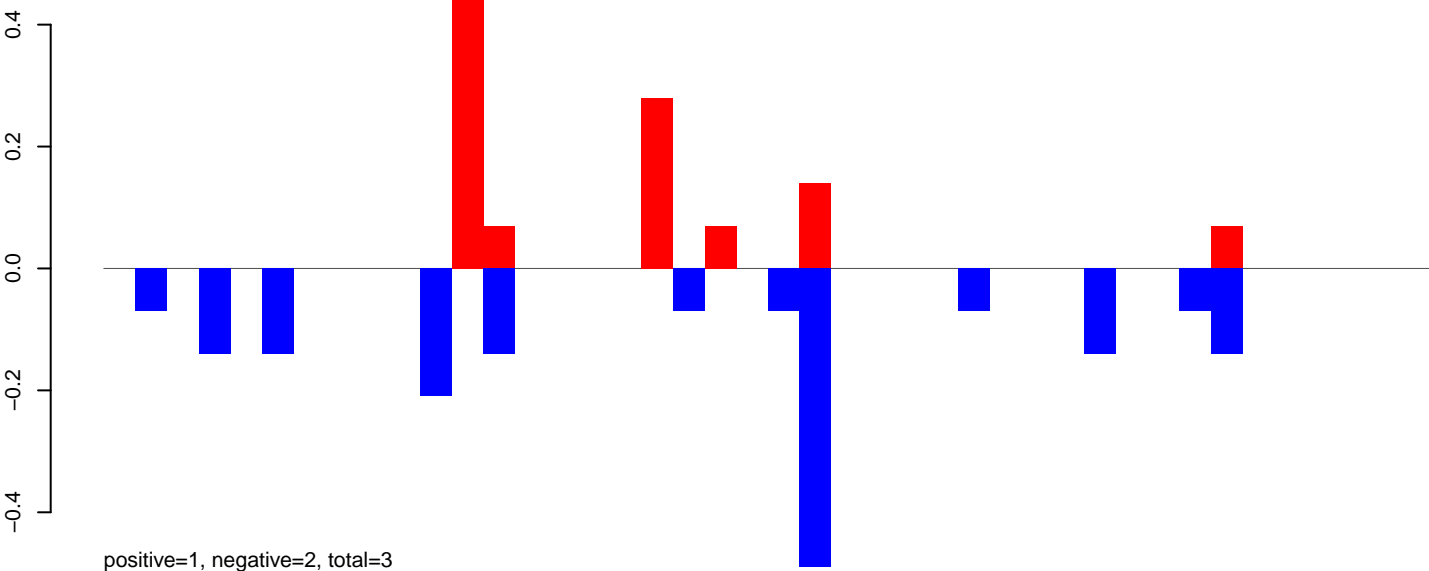


positive=4, negative=3, total=7

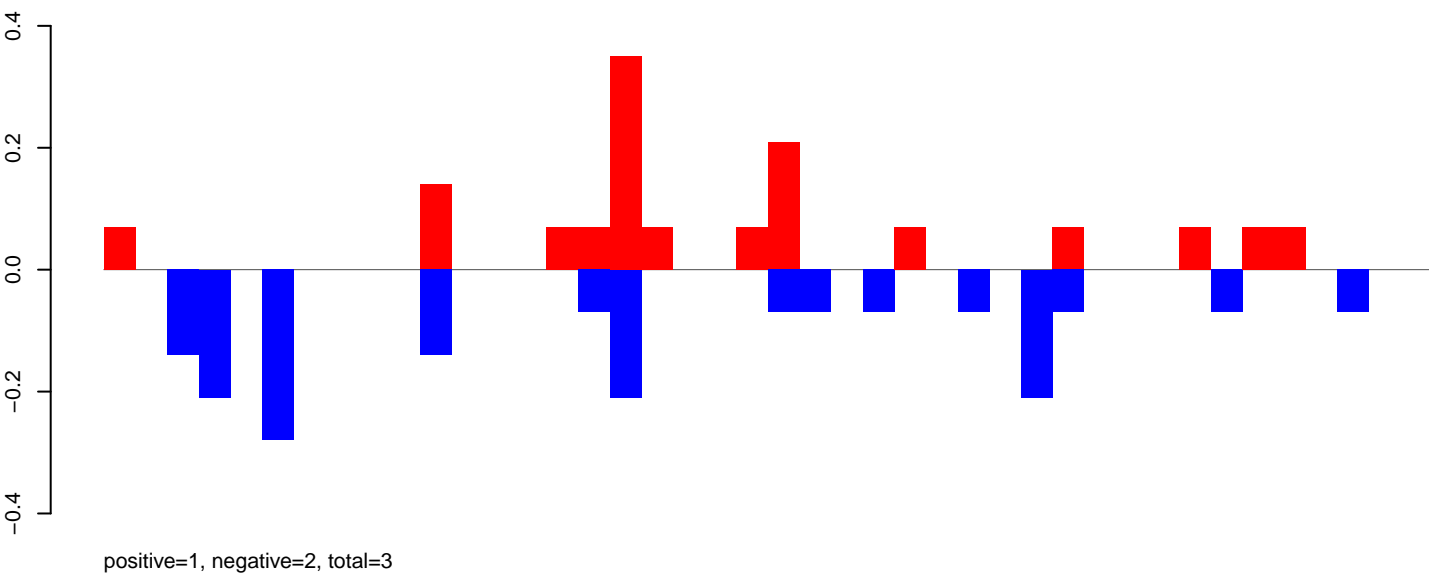
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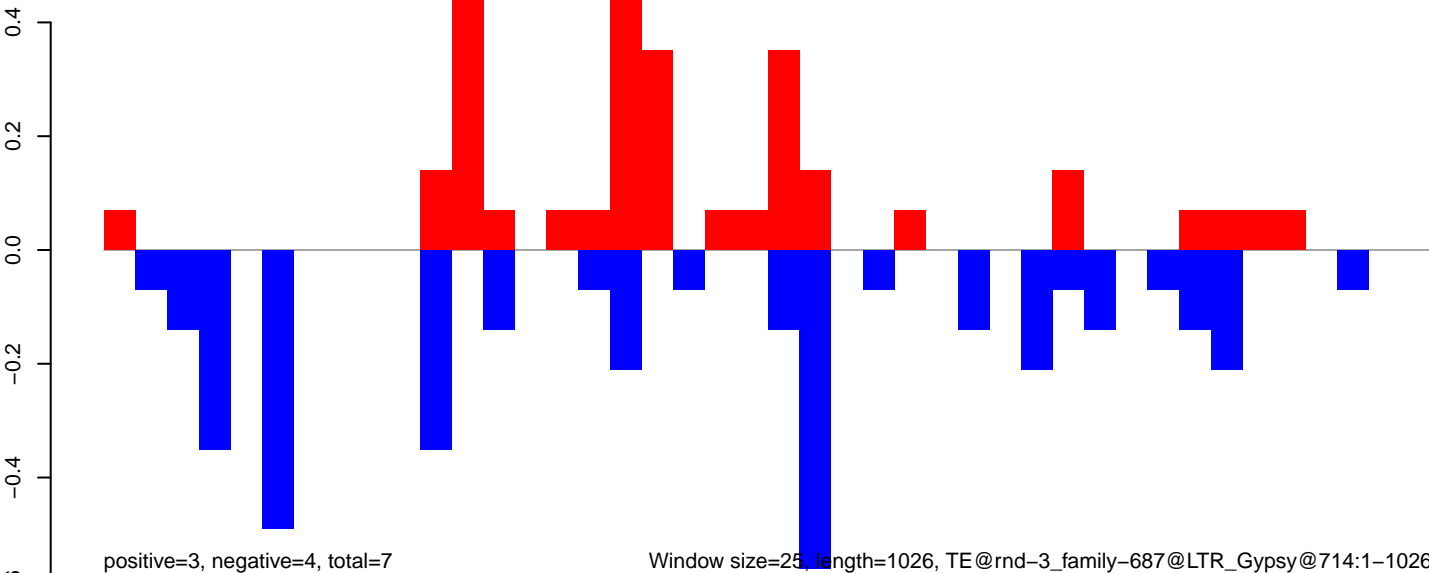
CuQuin_Mcarc_NL.18_23.rep



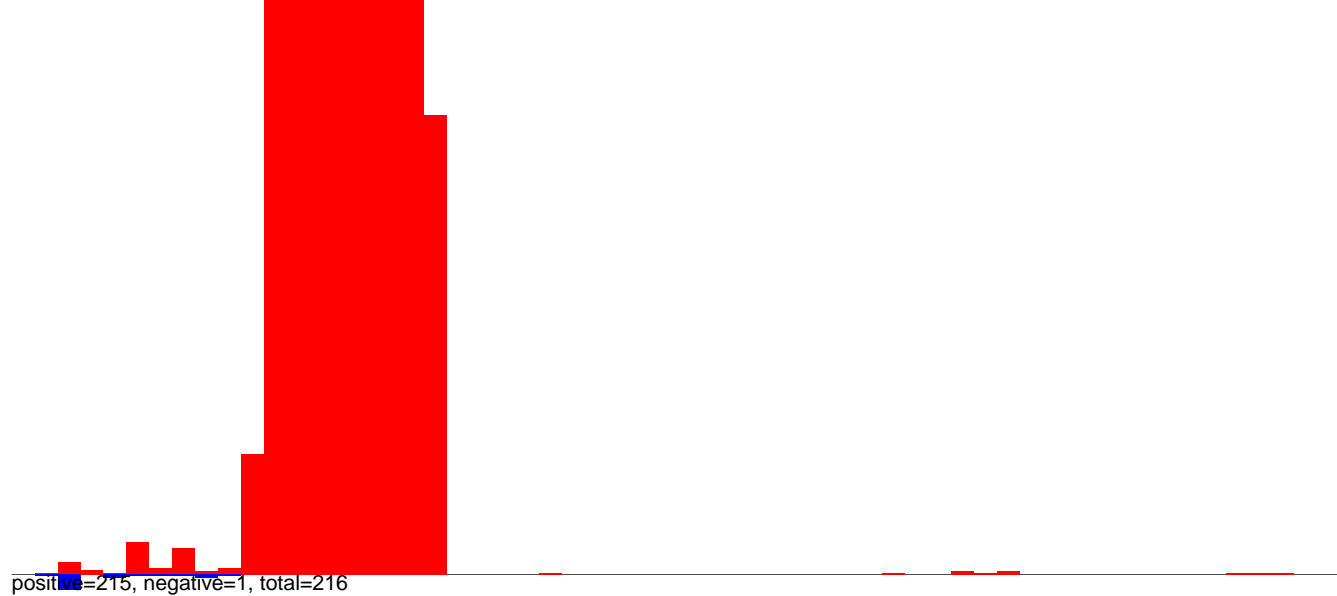
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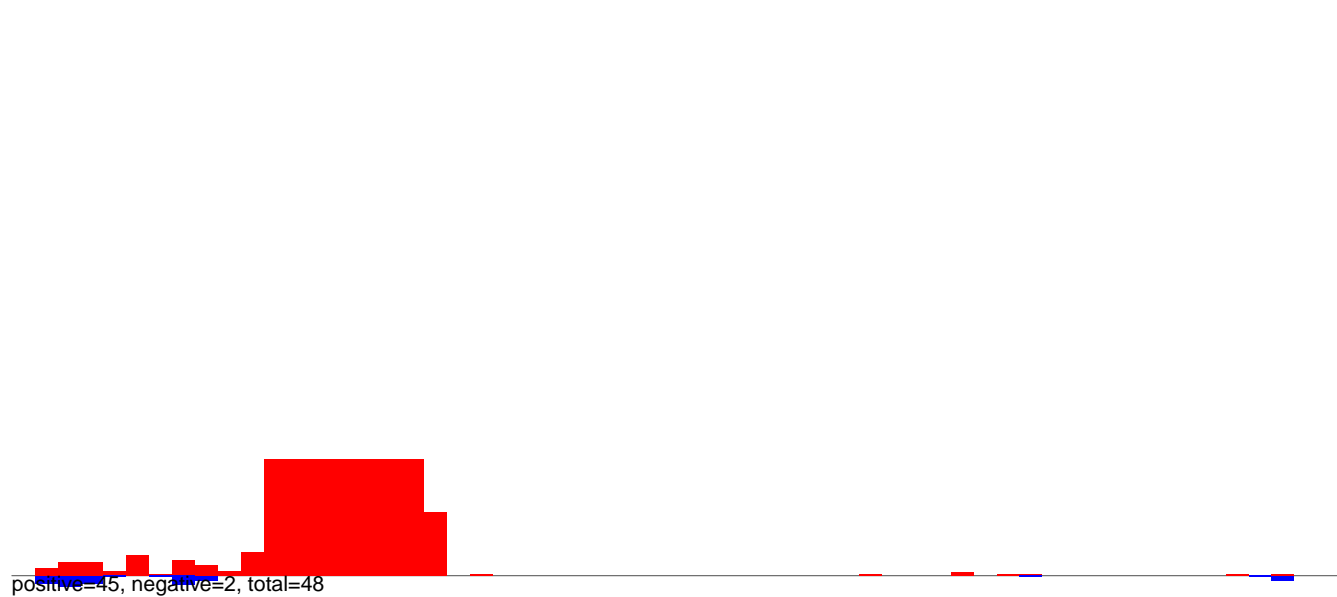
CuQuin_Mcarc_NL.rep



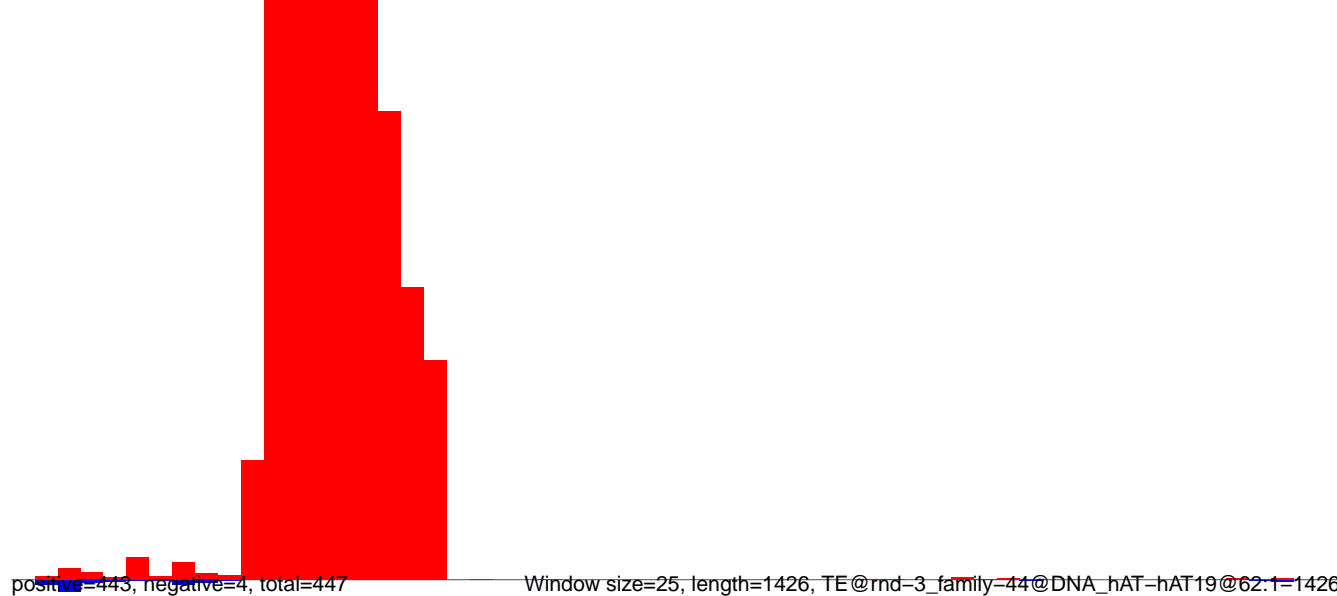
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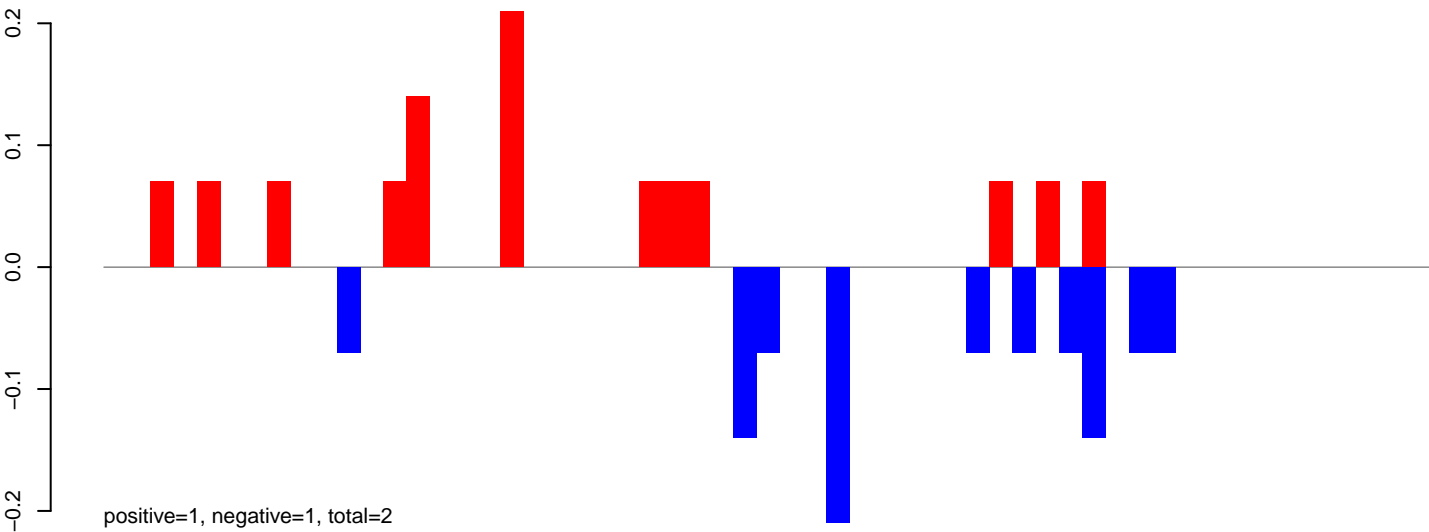
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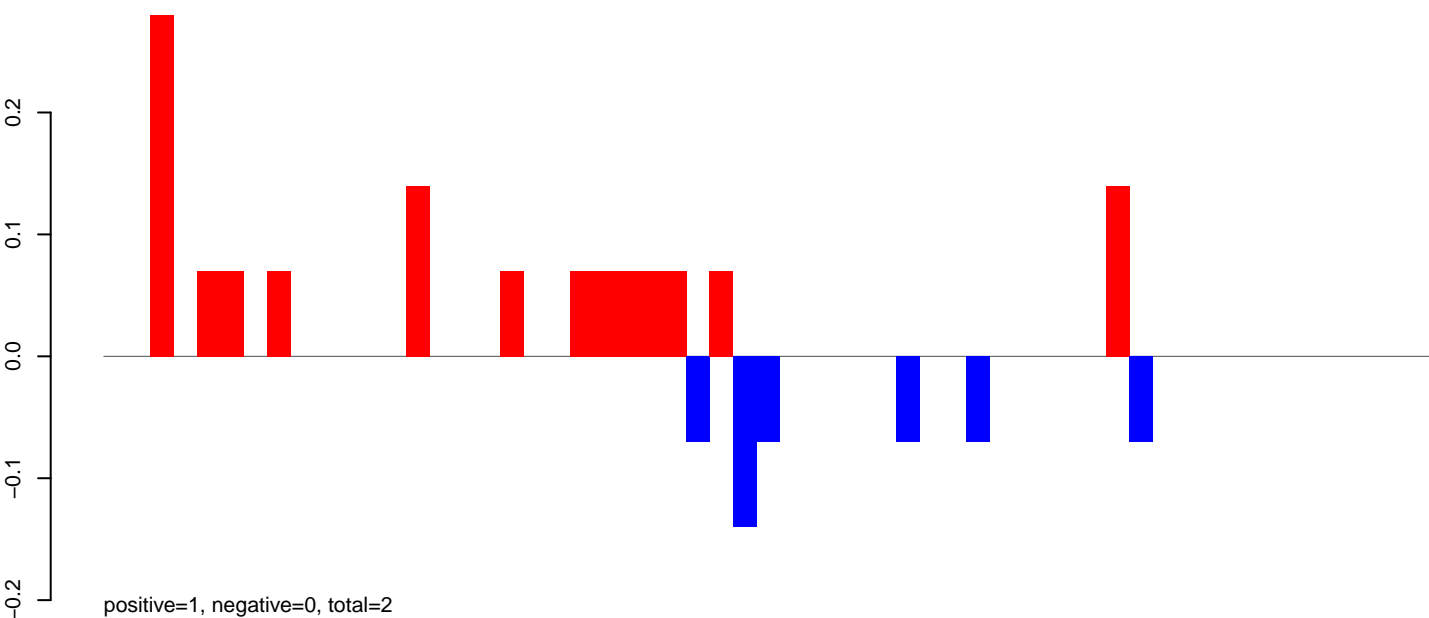
CuQuin_Mcarc_NL.rep



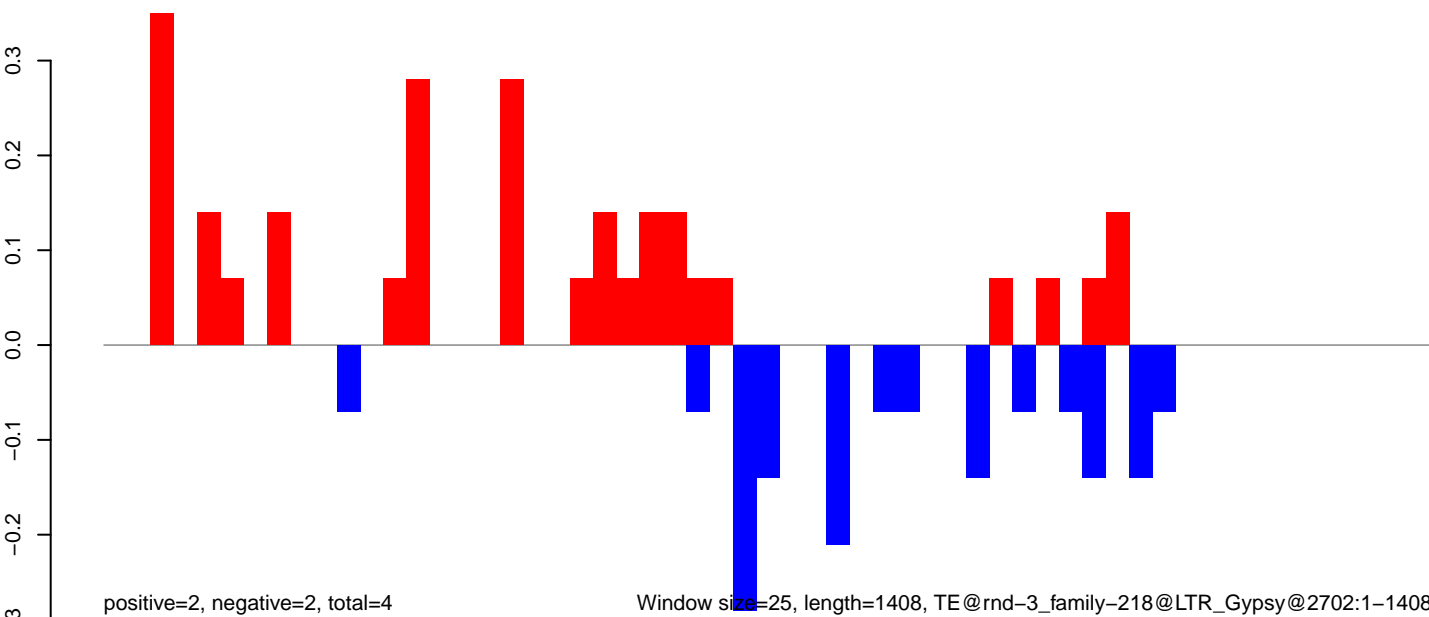
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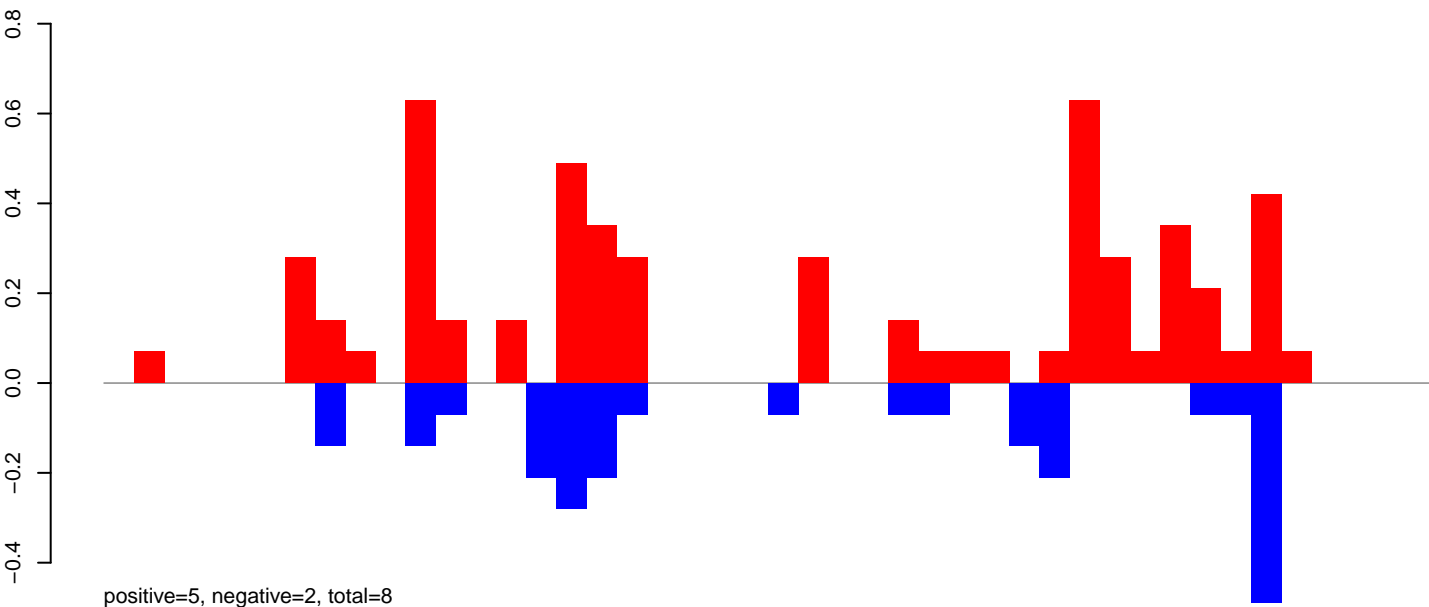
CuQuin_Mcarc_NL.24_35.rep



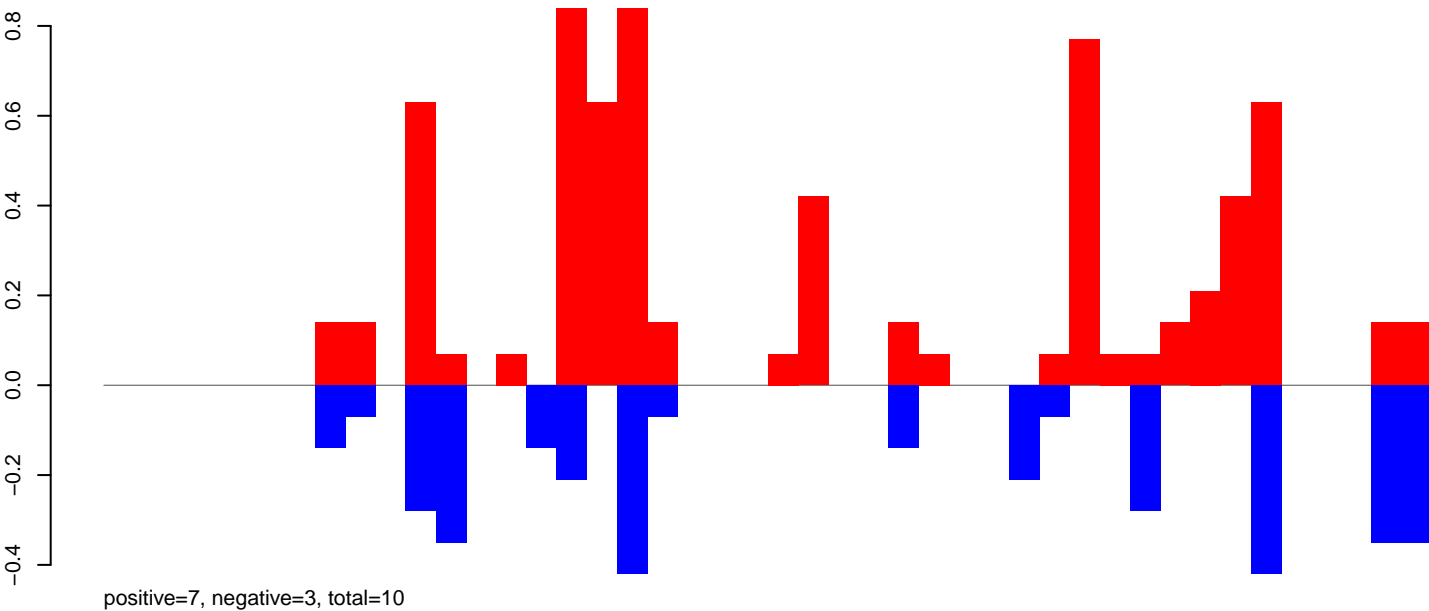
CuQuin_Mcarc_NL.rep



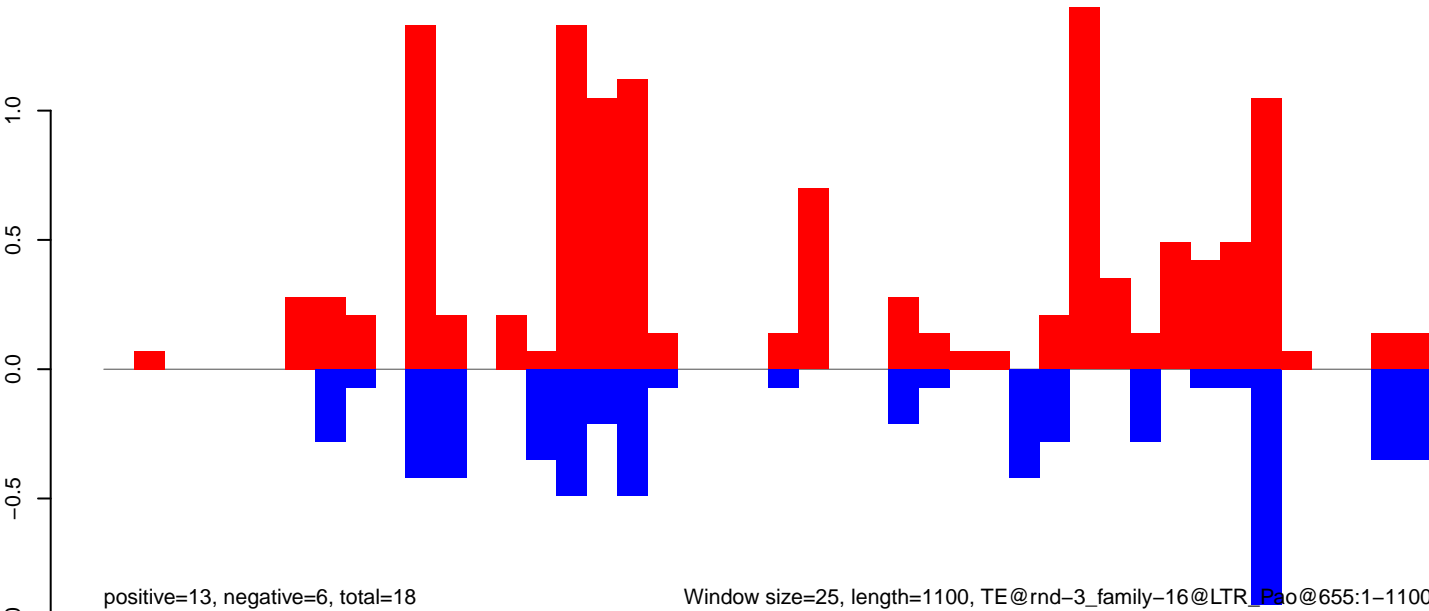
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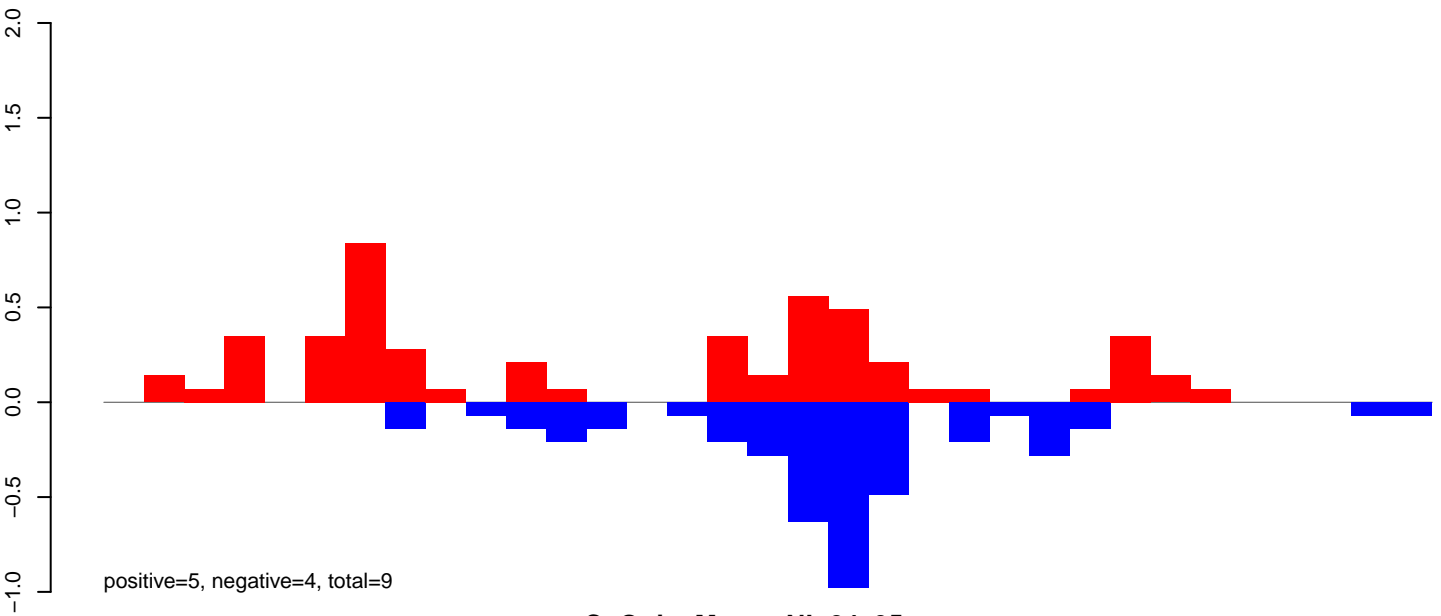
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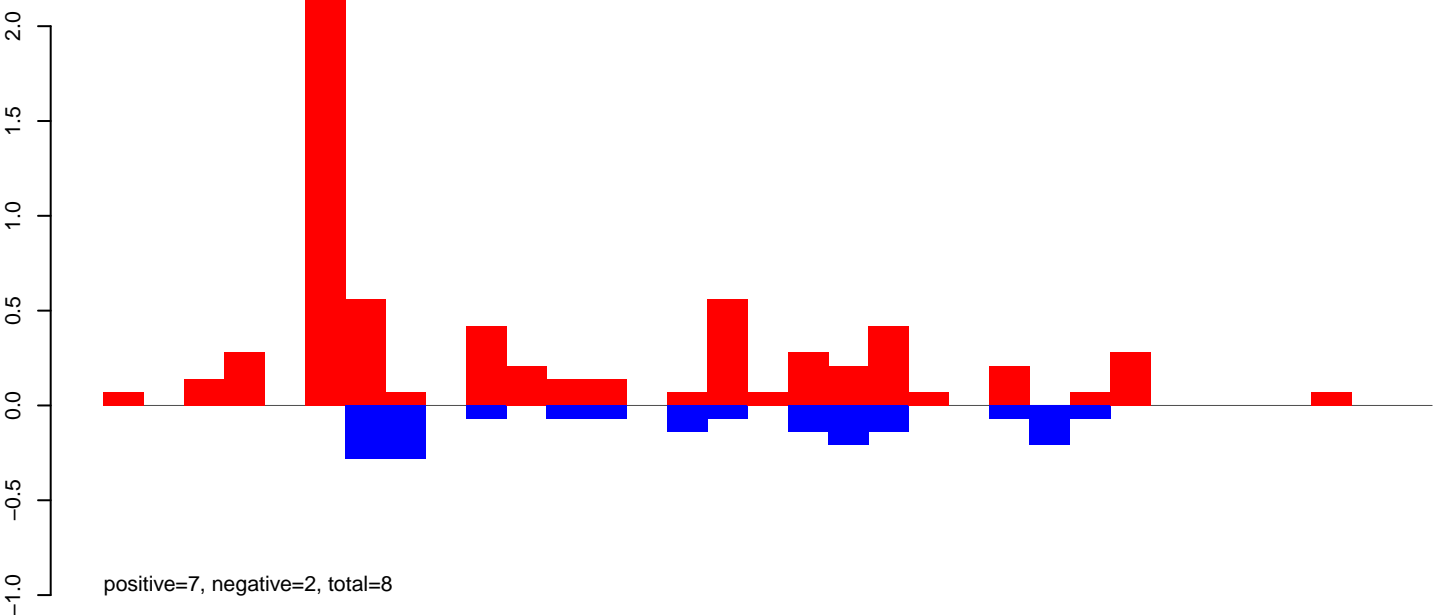
CuQuin_Mcarc_NL.rep



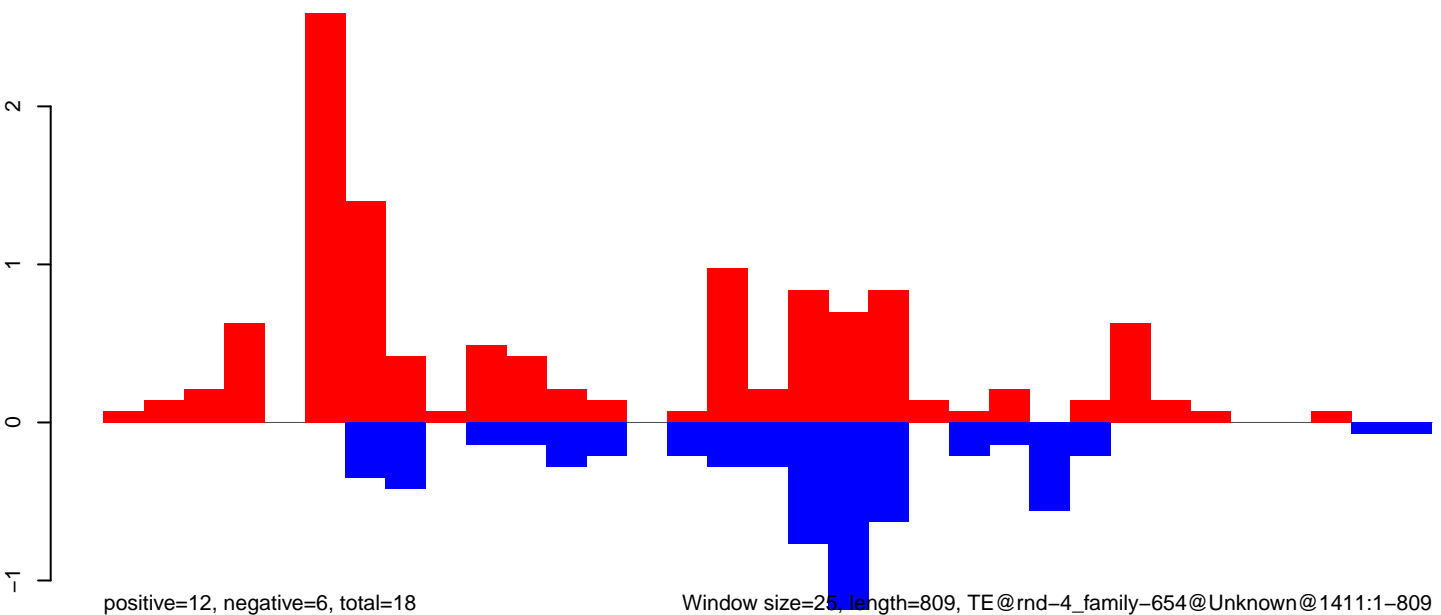
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

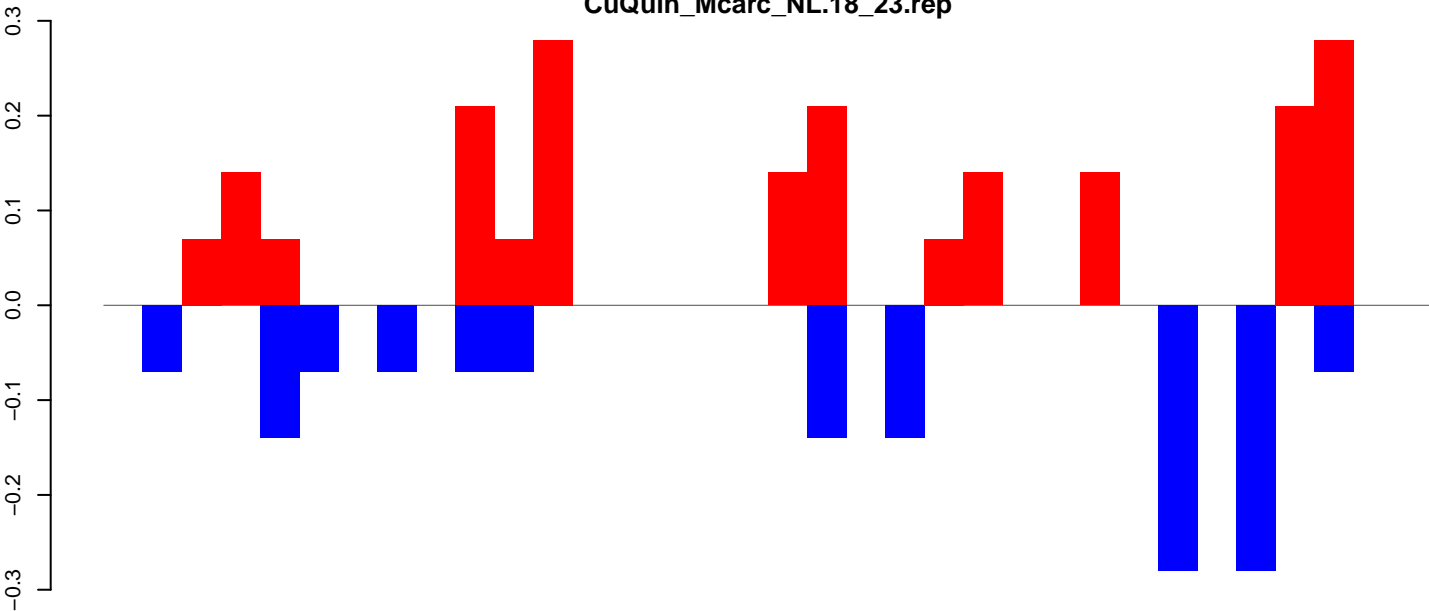


CuQuin_Mcarc_NL.rep



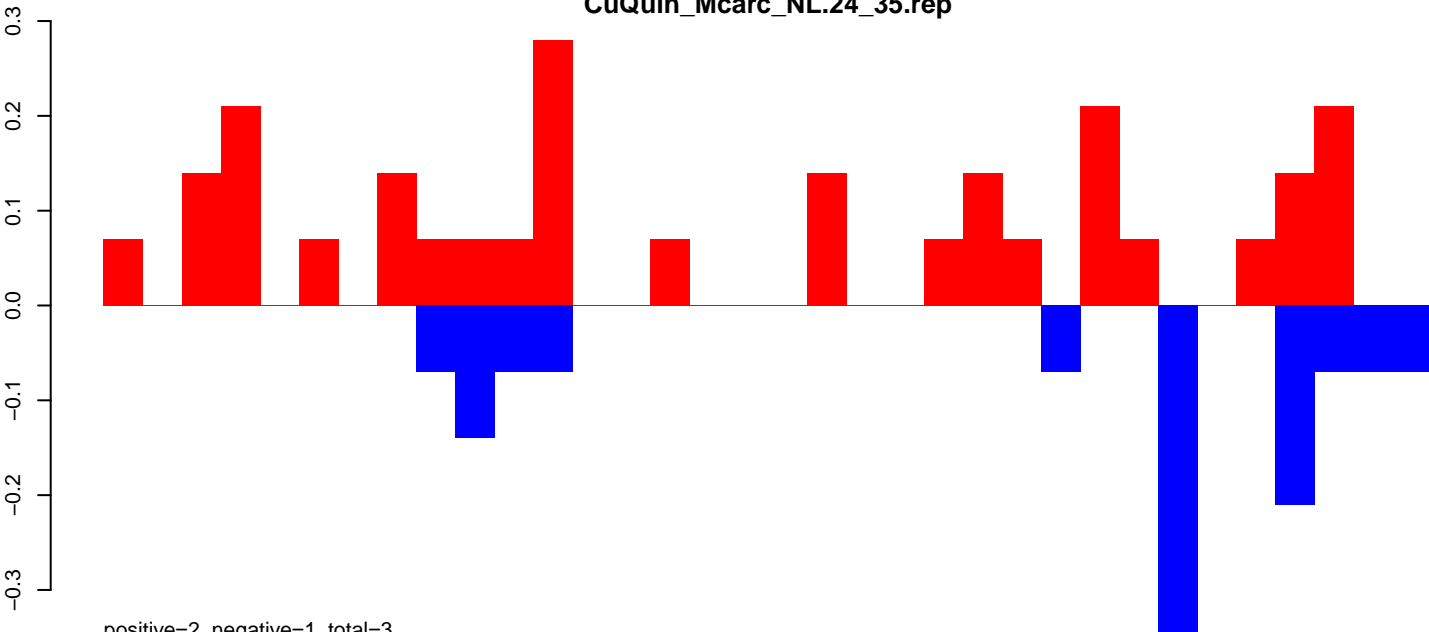
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CuQuin_Mcarc_NL.18_23.rep



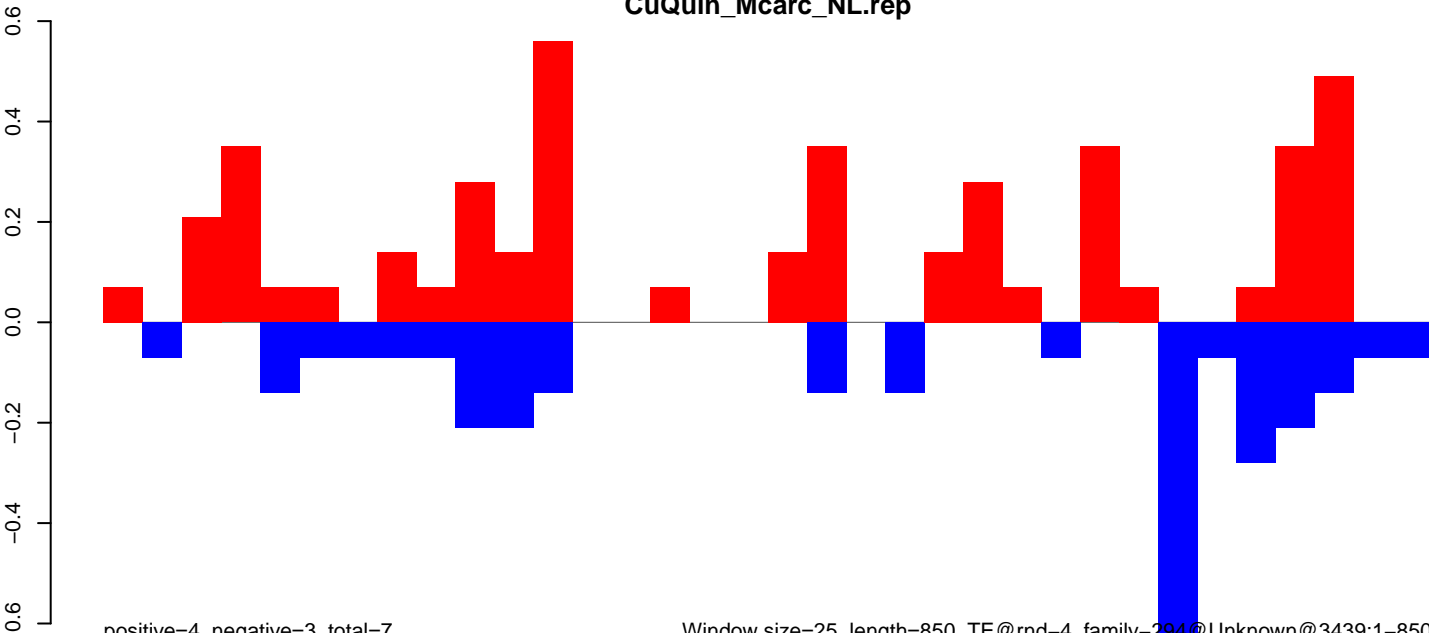
positive=2, negative=1, total=3

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=1, total=3

CuQuin_Mcarc_NL.rep

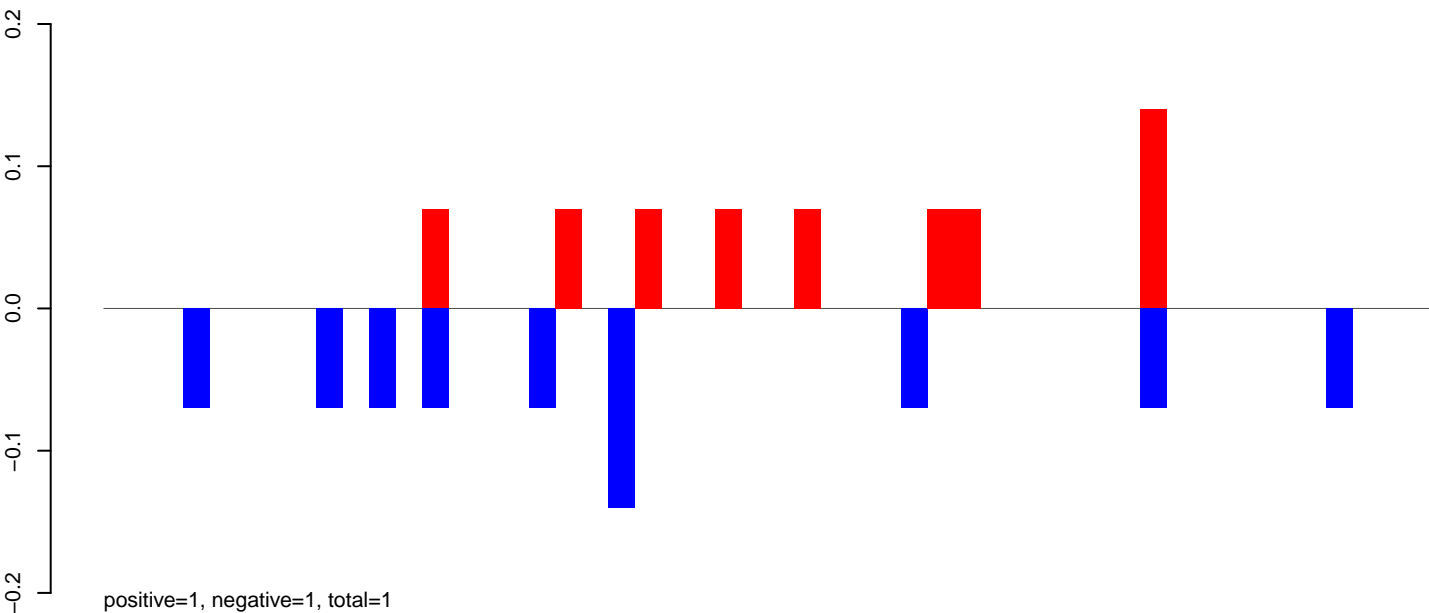


positive=4, negative=3, total=7

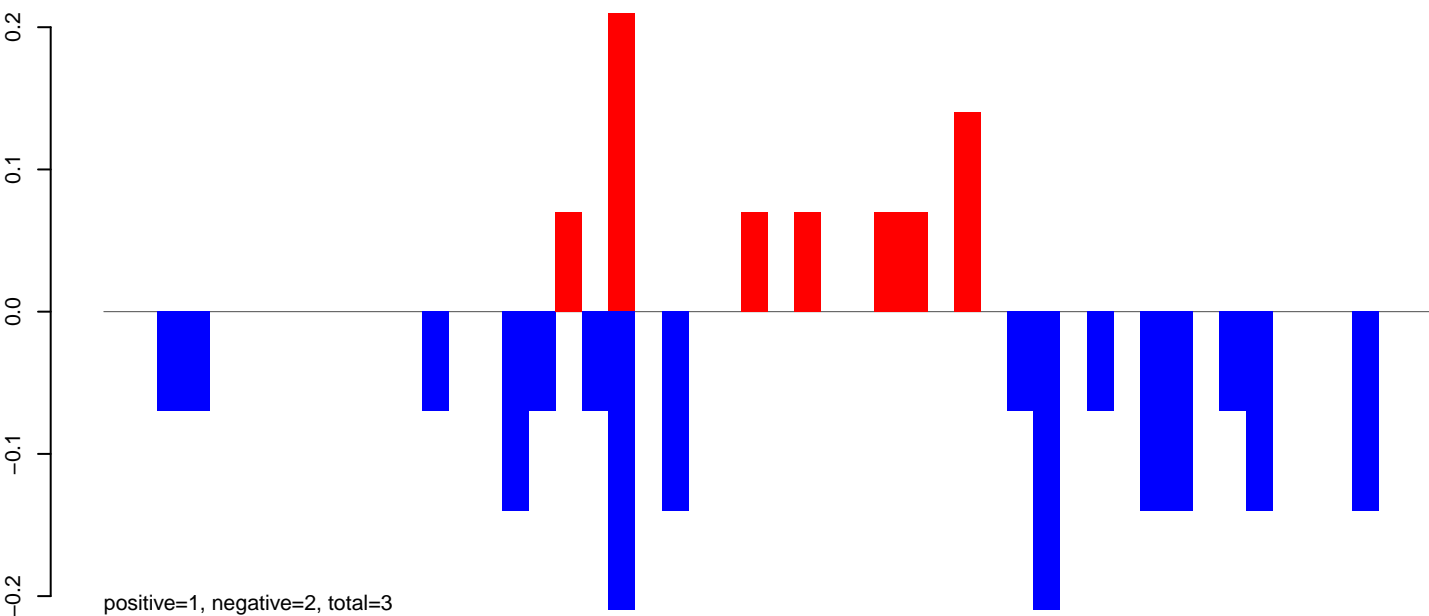
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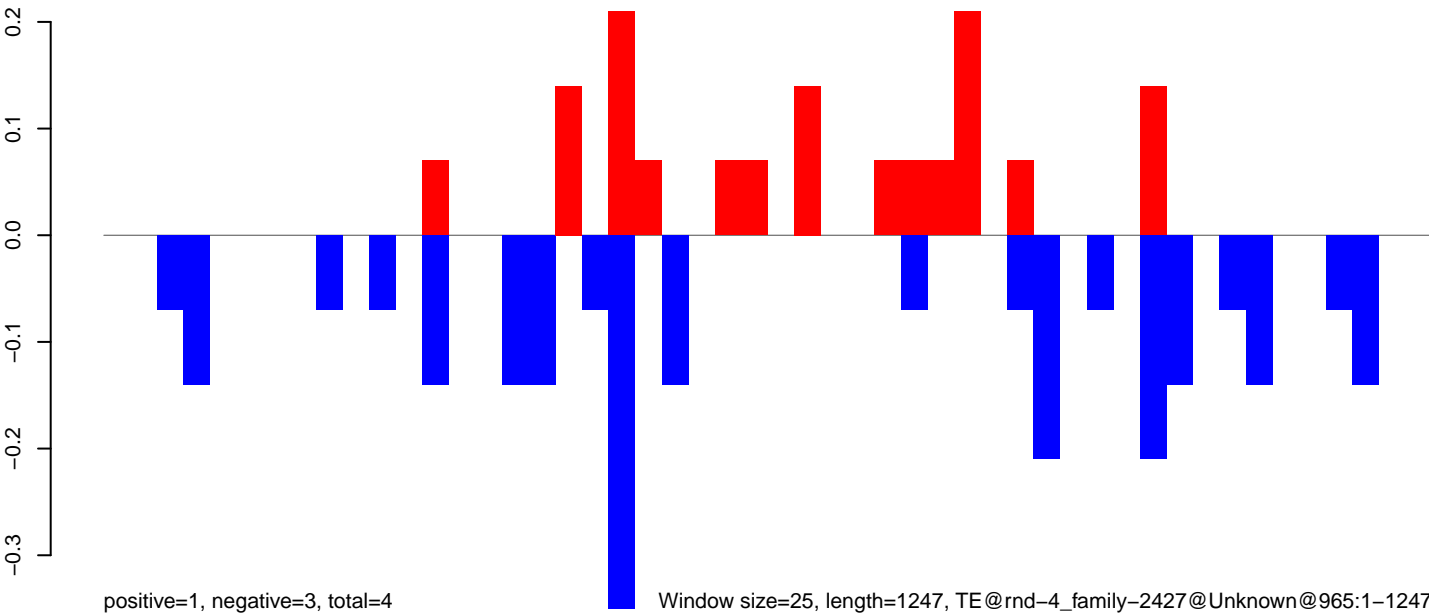
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

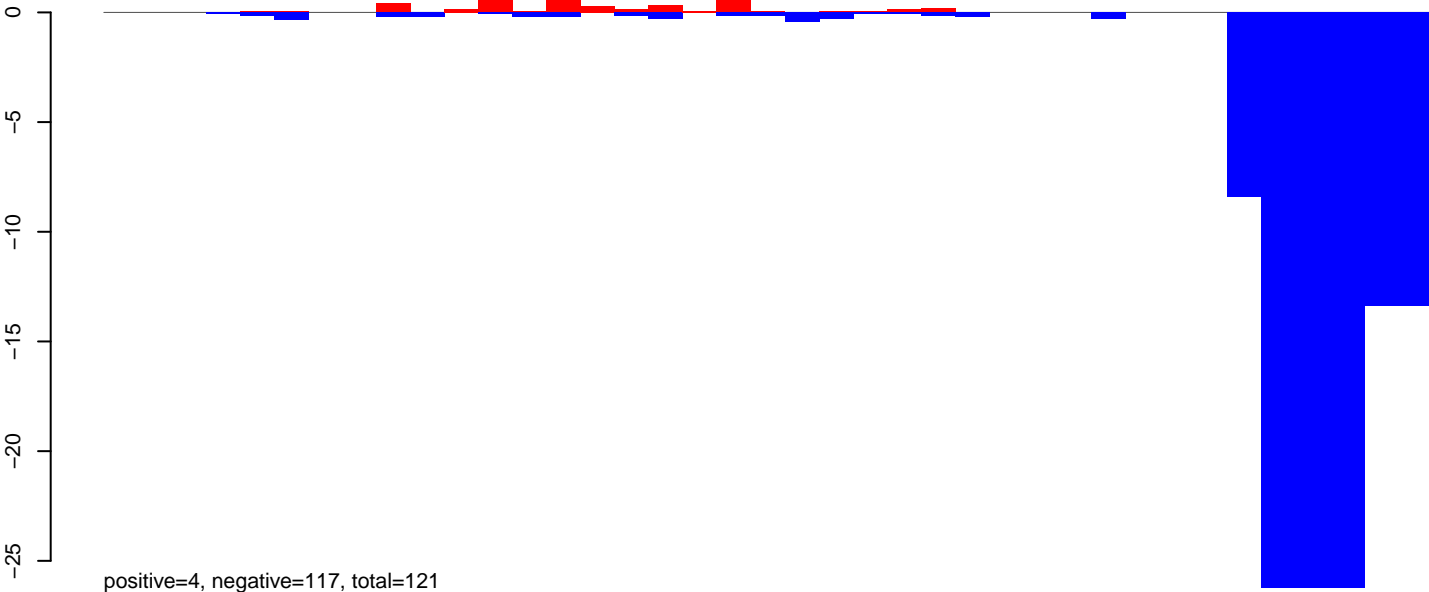


CuQuin_Mcarc_NL.rep

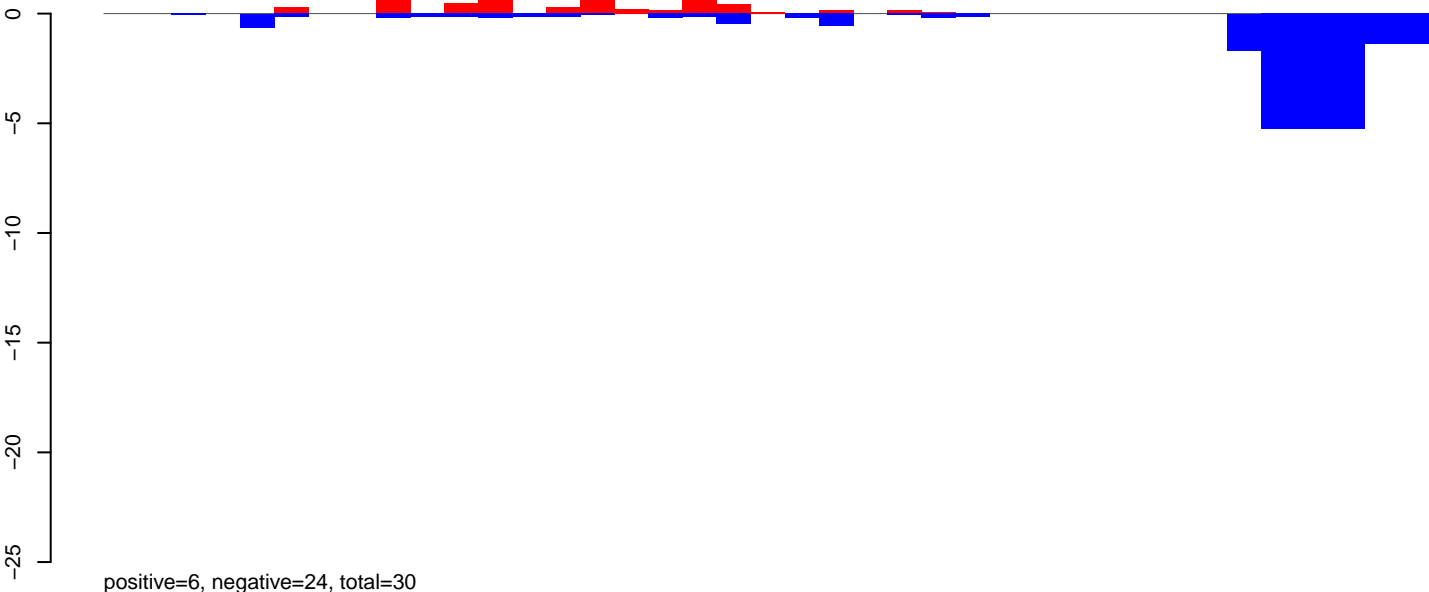


Window size=25, length=1247, TE@rnd-4_family-2427@Unknown@965:1-1247

CuQuin_Mcarc_NL.18_23.rep



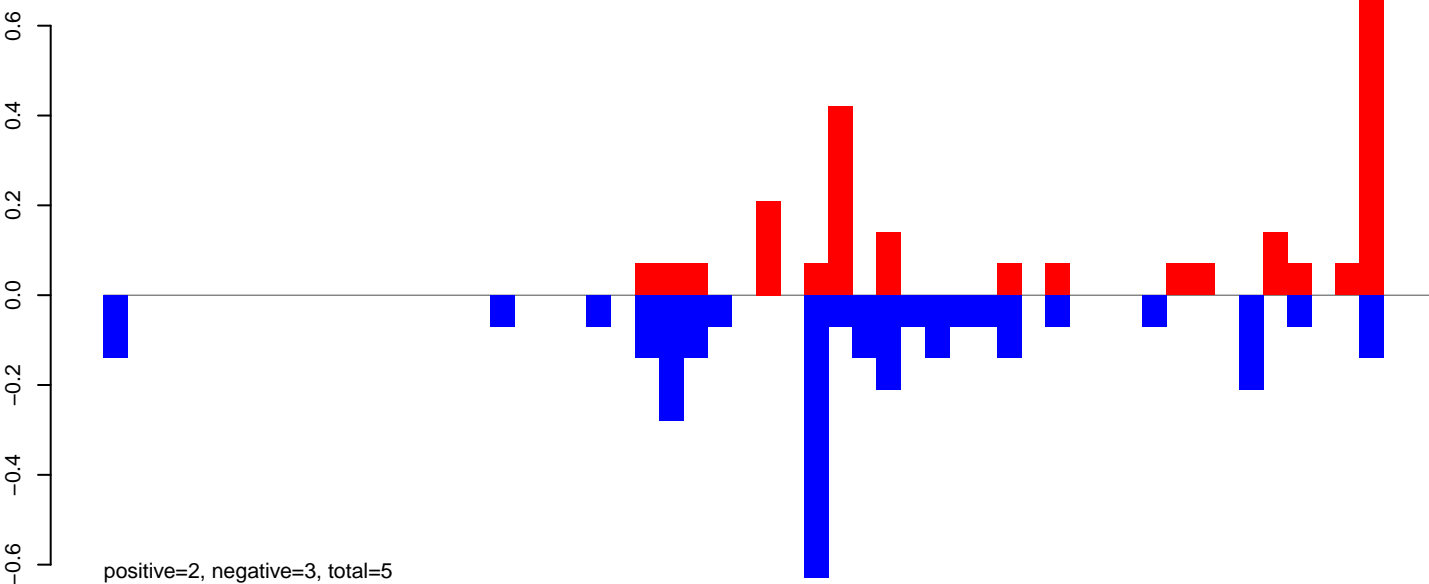
CuQuin_Mcarc_NL.24_35.rep



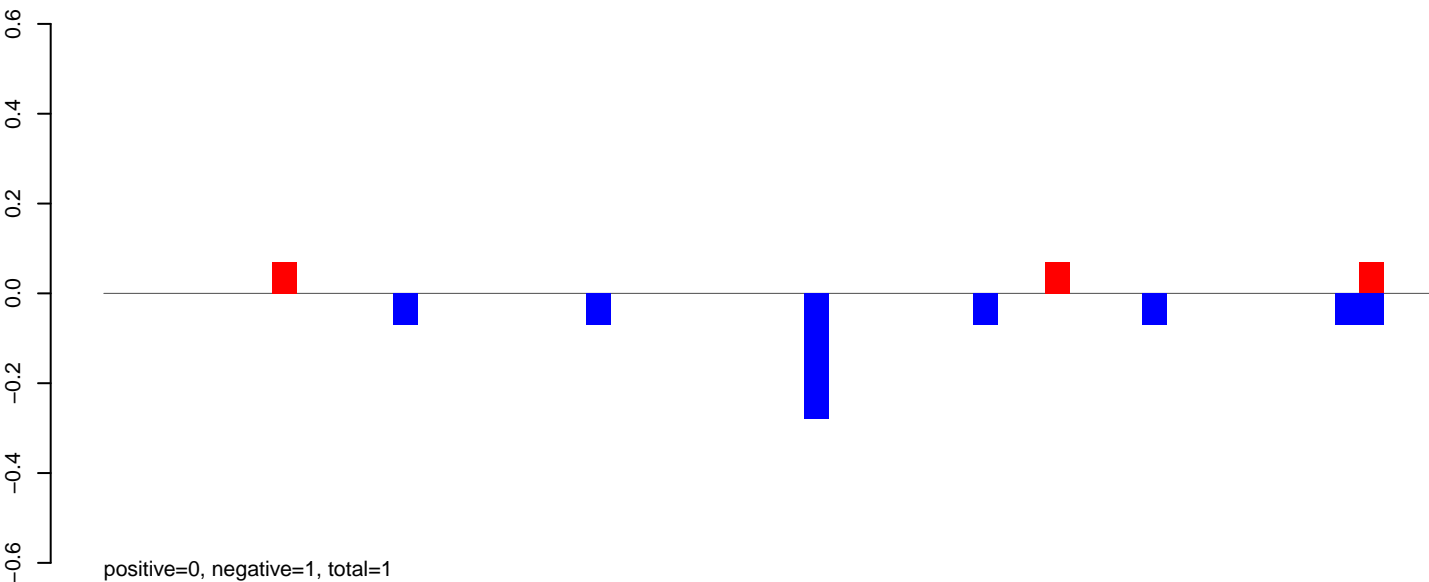
CuQuin_Mcarc_NL.rep



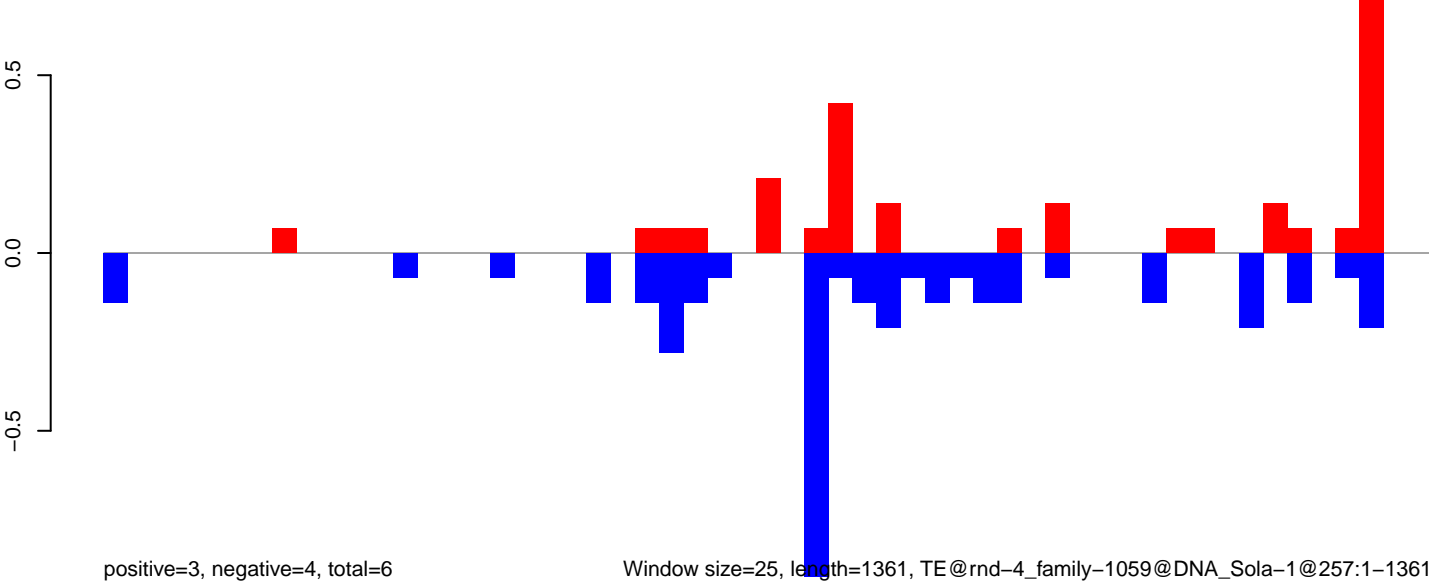
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

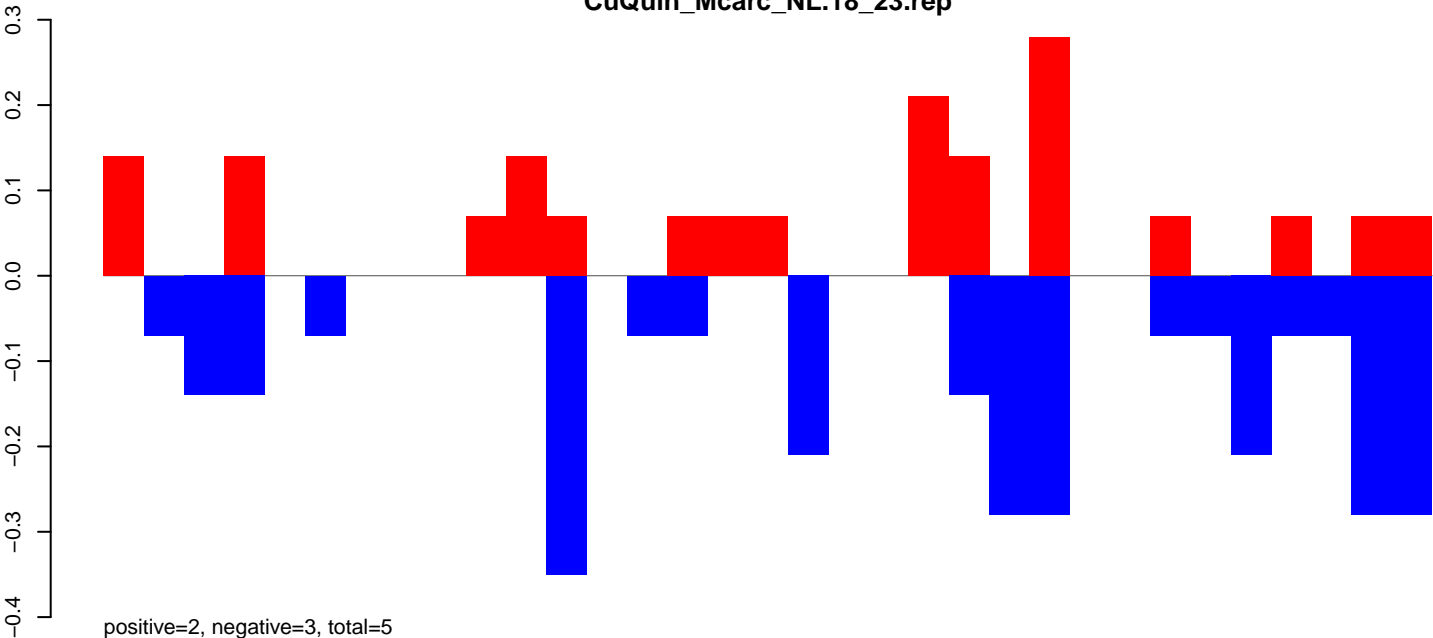


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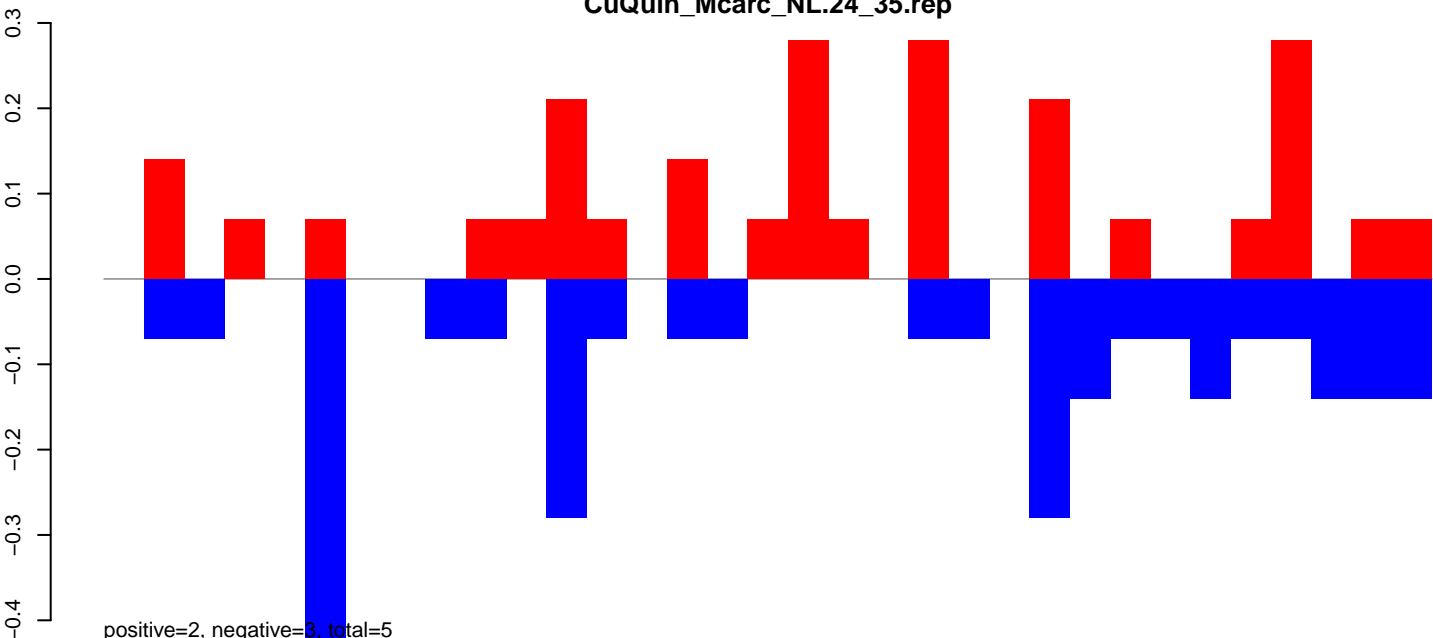
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CuQuin_Mcarc_NL.18_23.rep



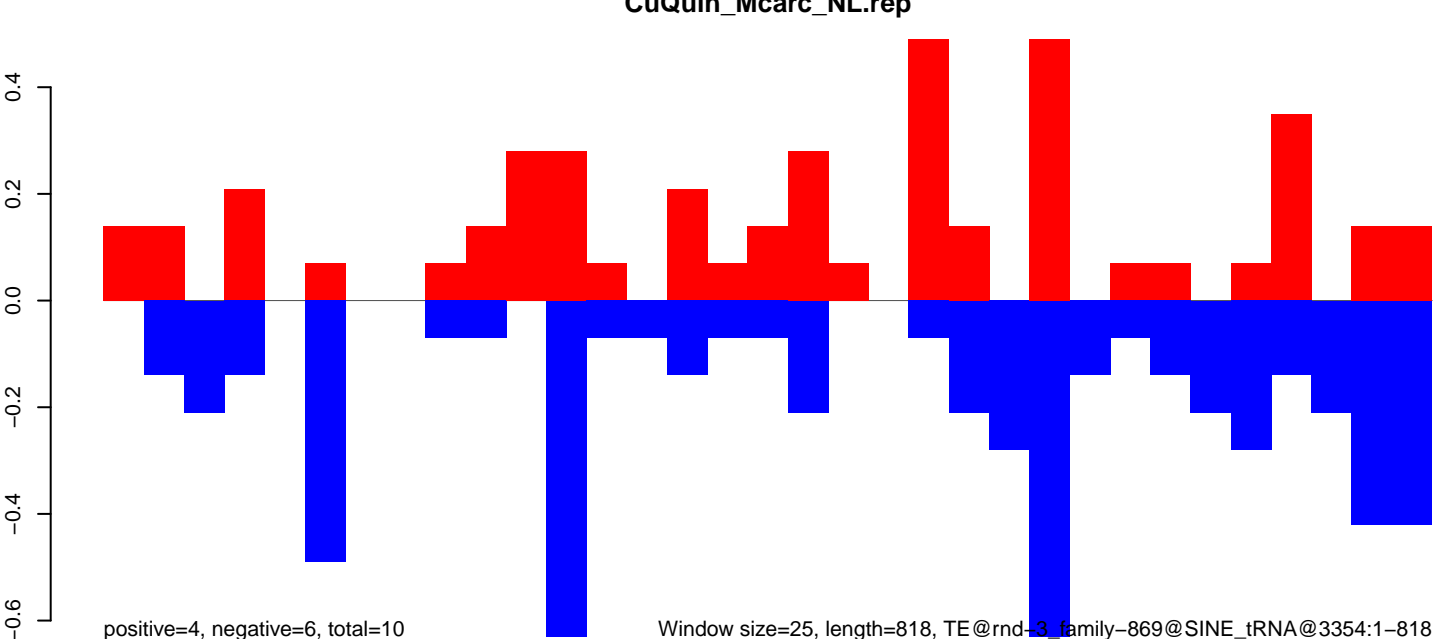
positive=2, negative=3, total=5

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=3, total=5

CuQuin_Mcarc_NL.rep



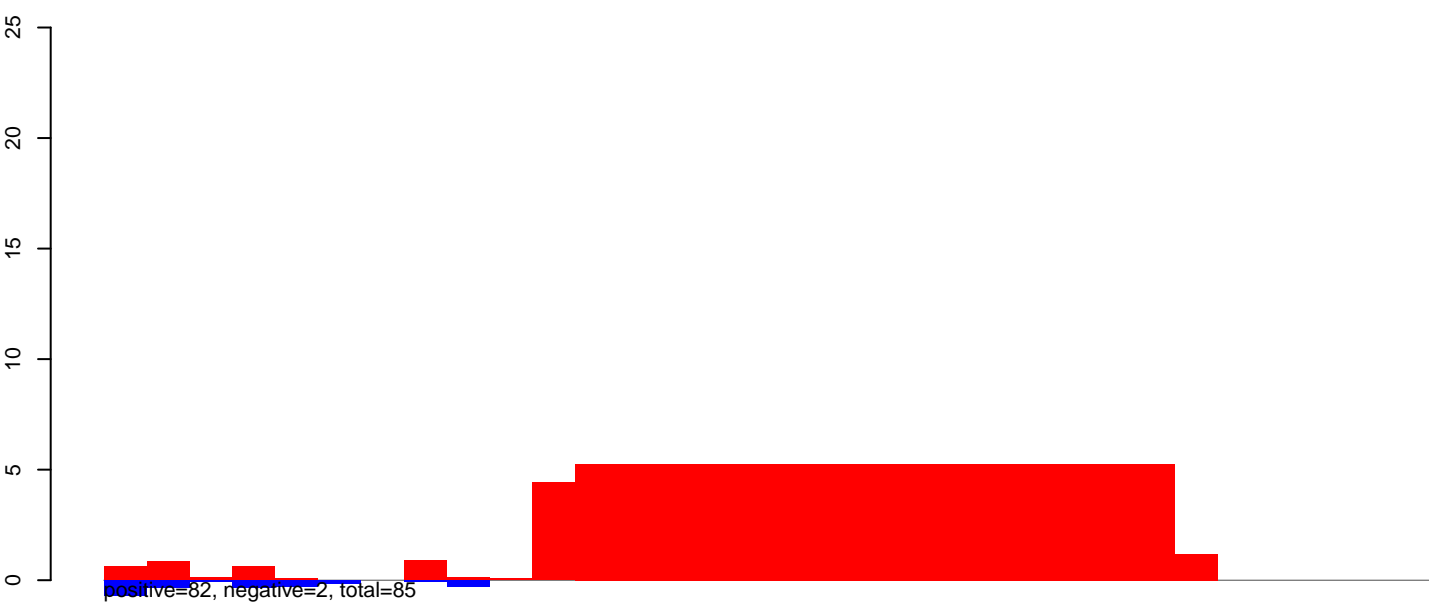
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Window size=25, length=818, TE@rnd-3, family-869@SINE_tRNA@3354:1-818

CuQuin_Mcarc_NL.18_23.rep



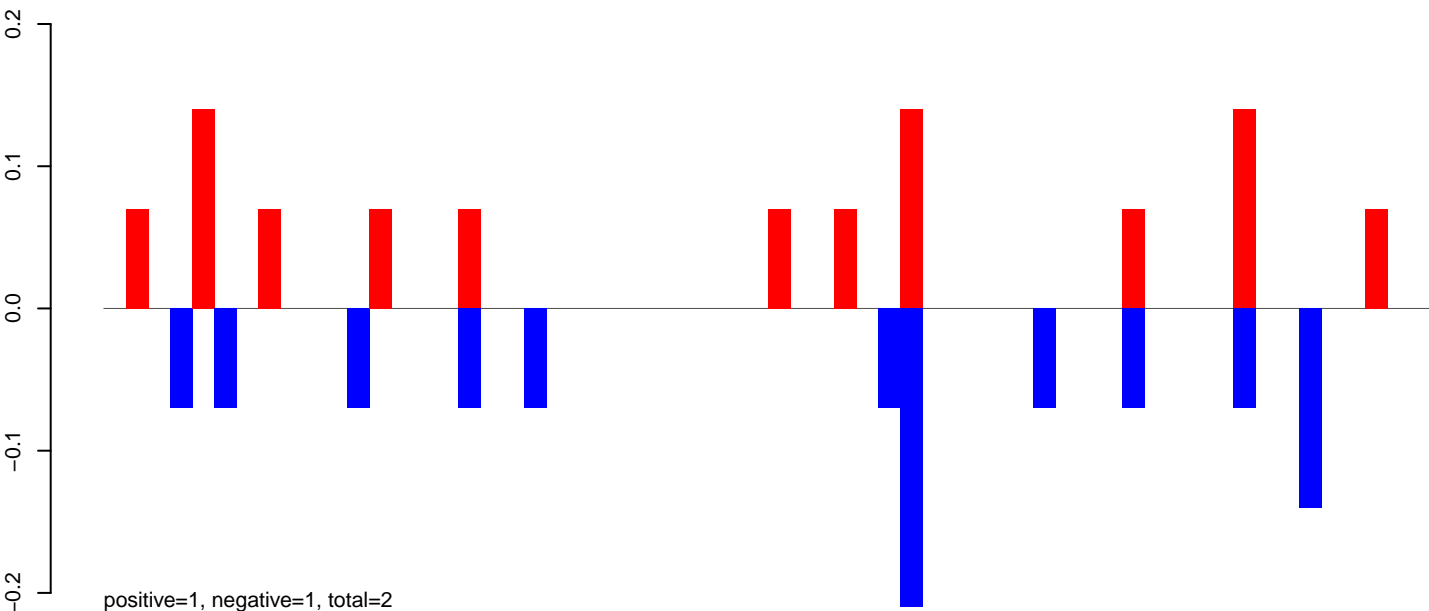
CuQuin_Mcarc_NL.24_35.rep



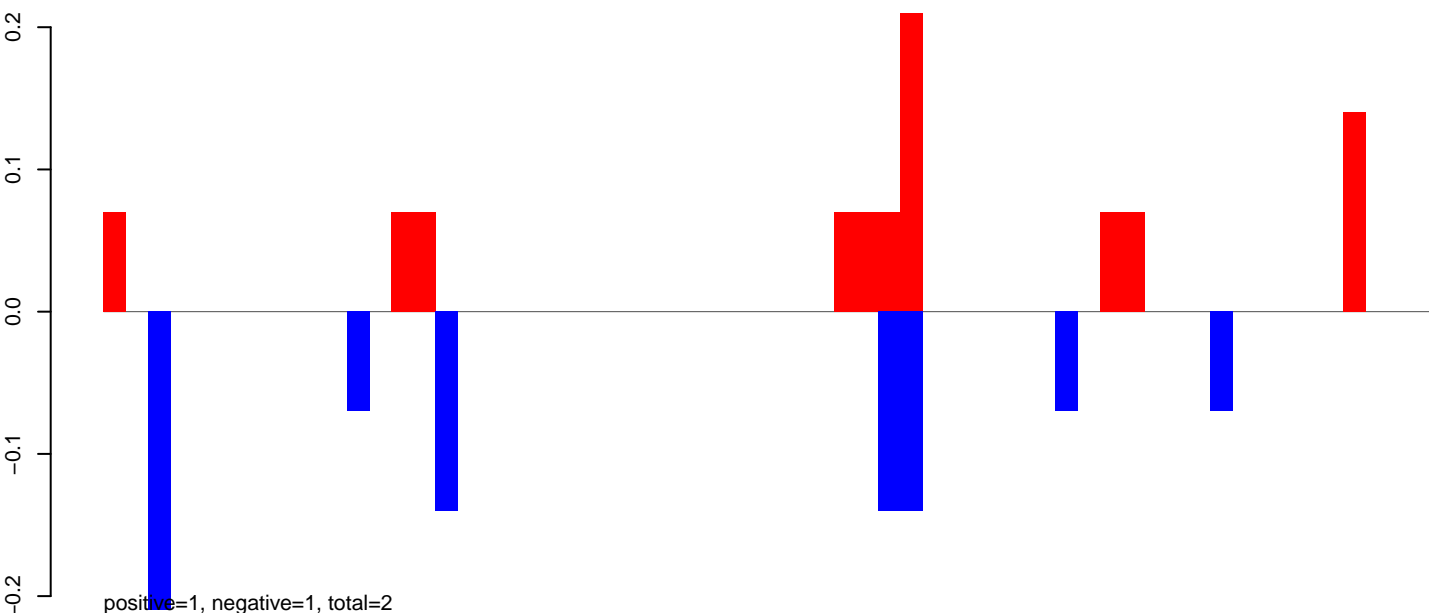
CuQuin_Mcarc_NL.rep



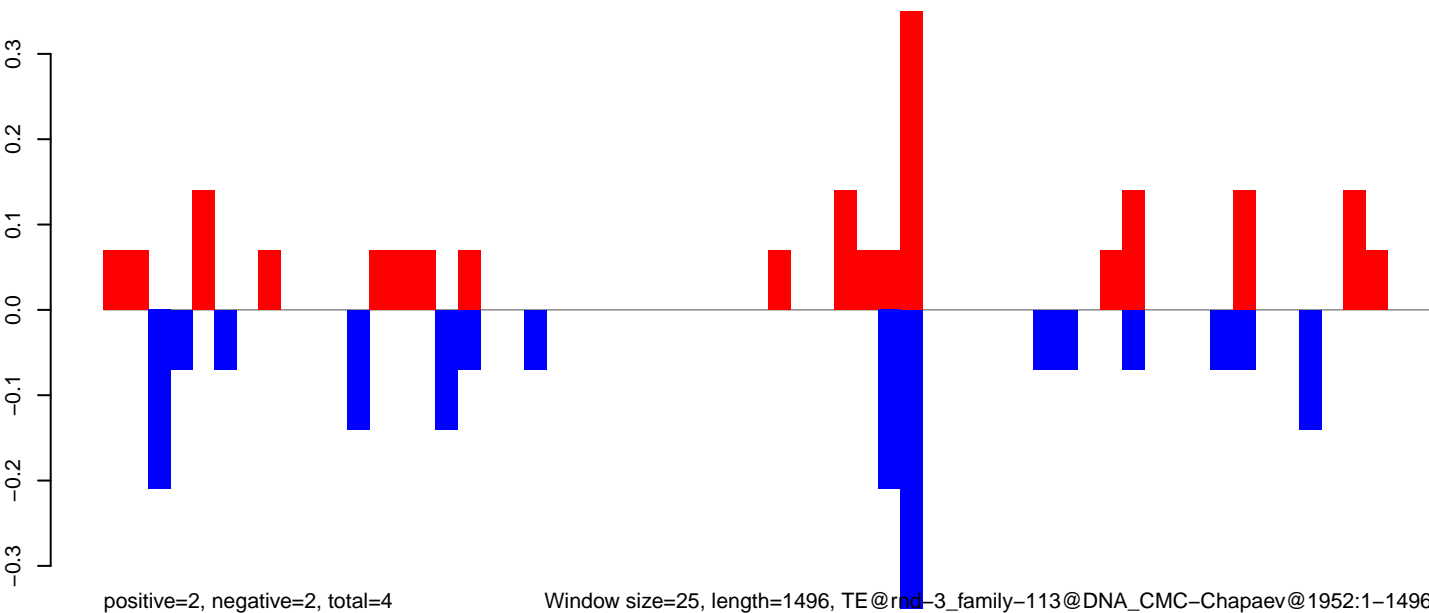
CuQuin_Mcarc_NL.18_23.rep



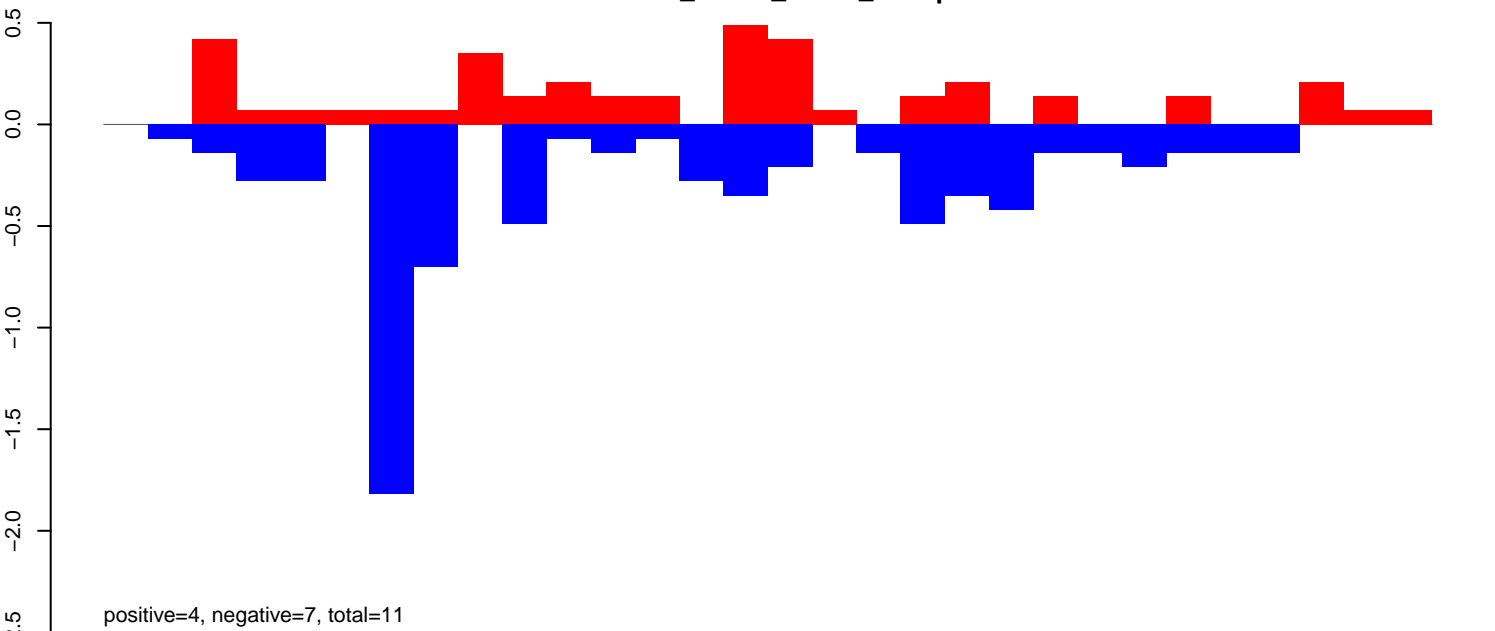
CuQuin_Mcarc_NL.24_35.rep



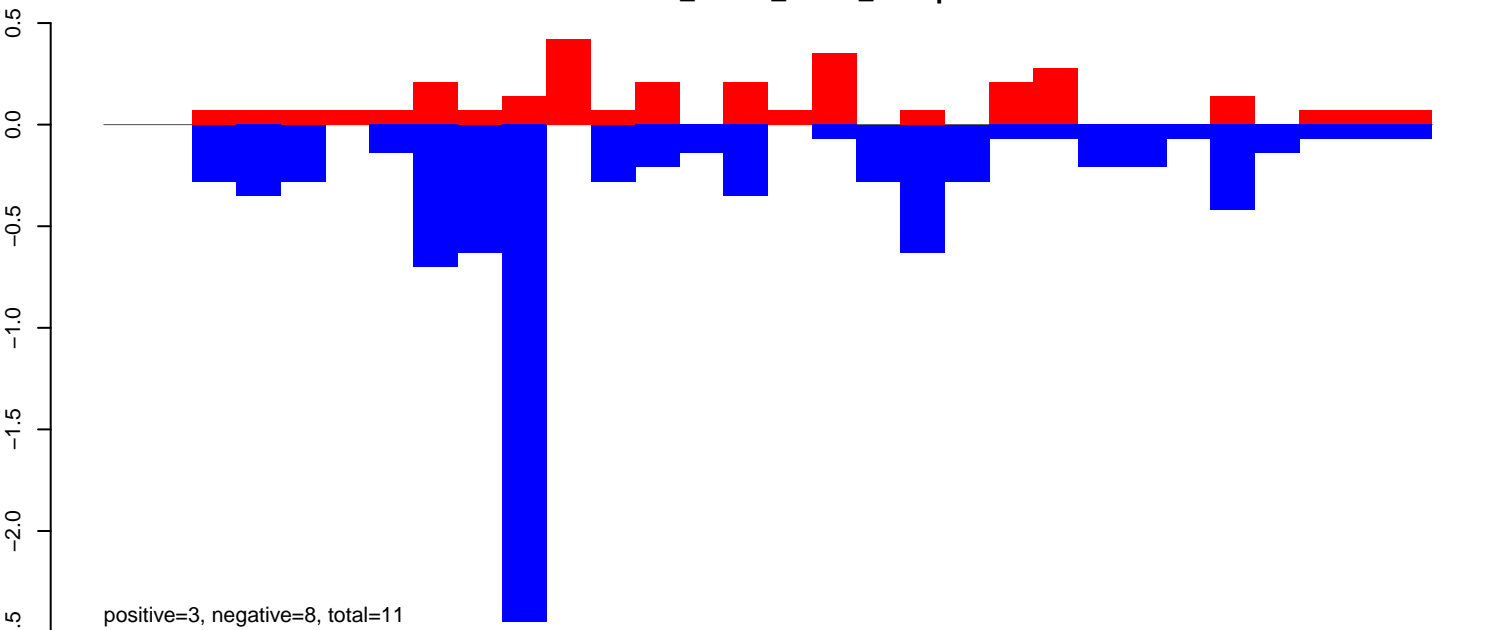
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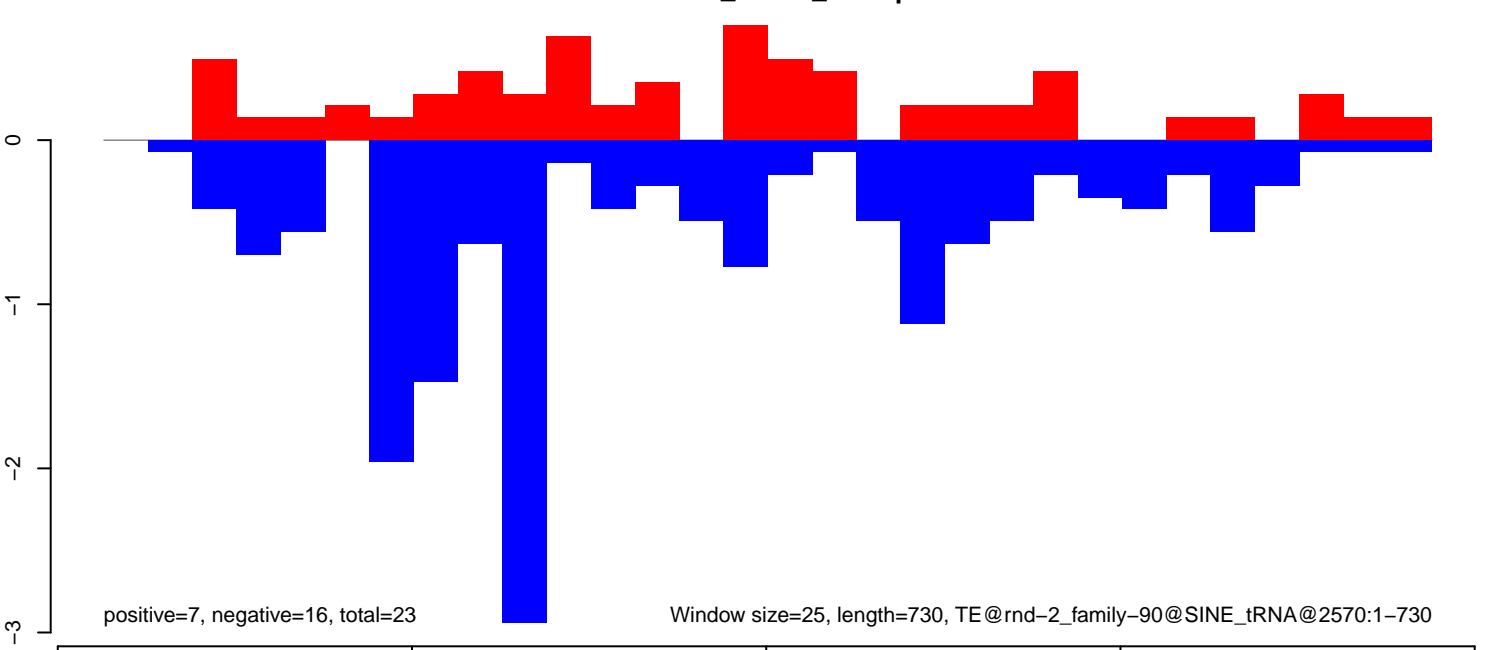
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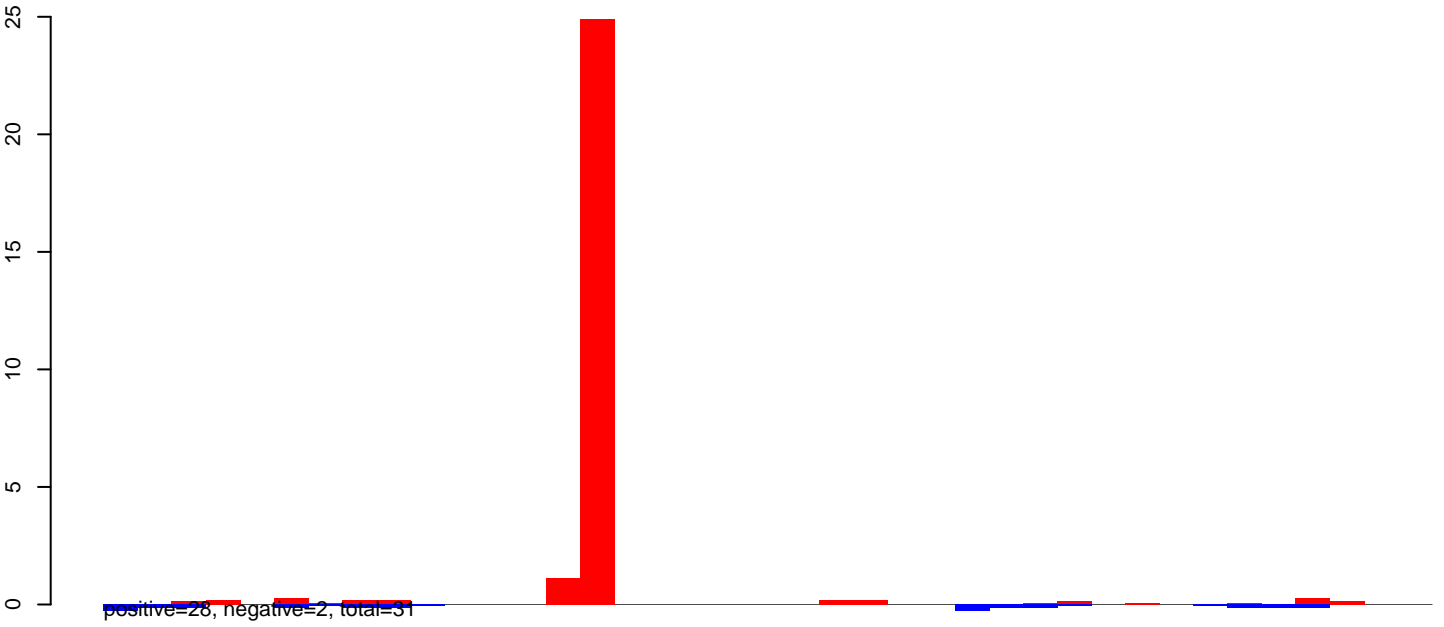
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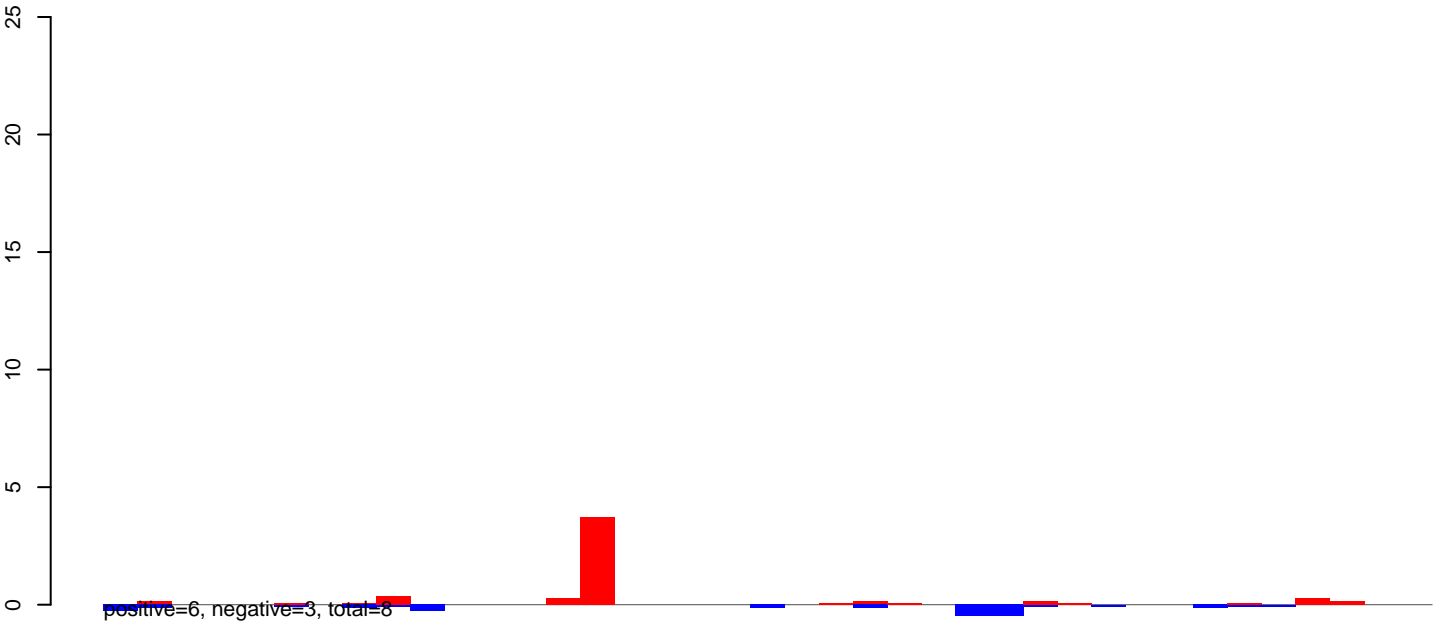
CuQuin_Mcarc_NL.rep



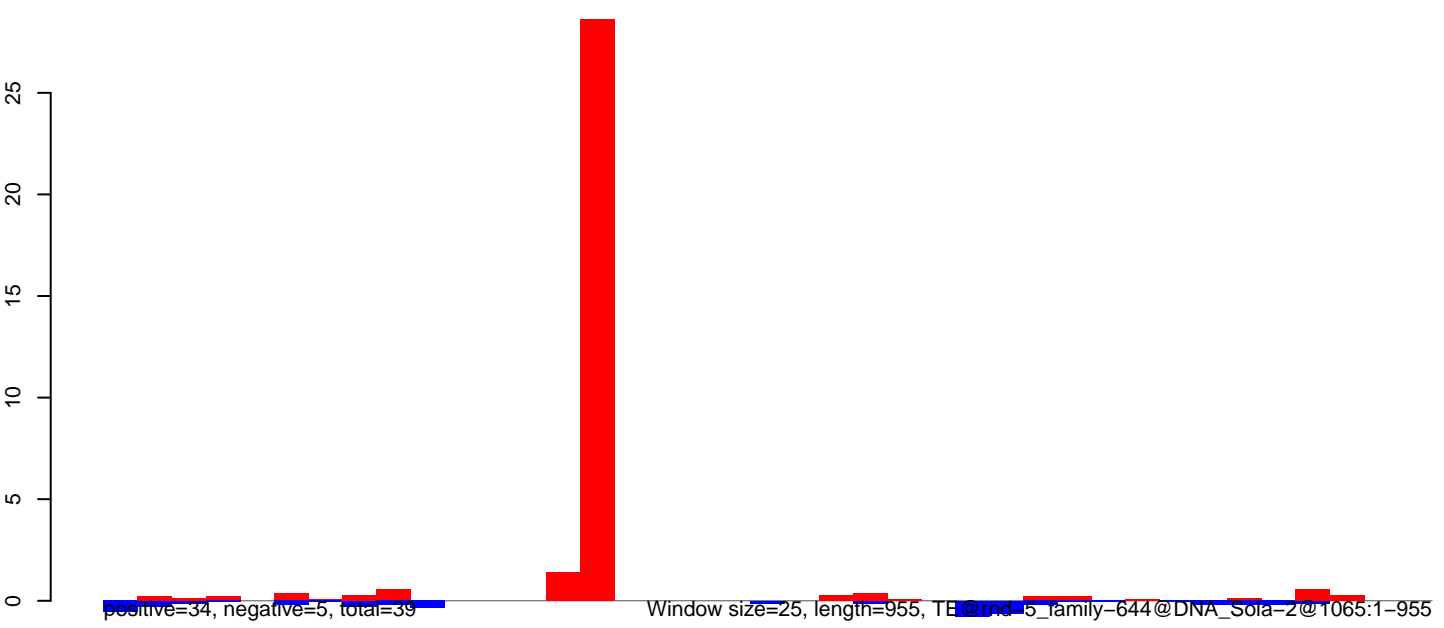
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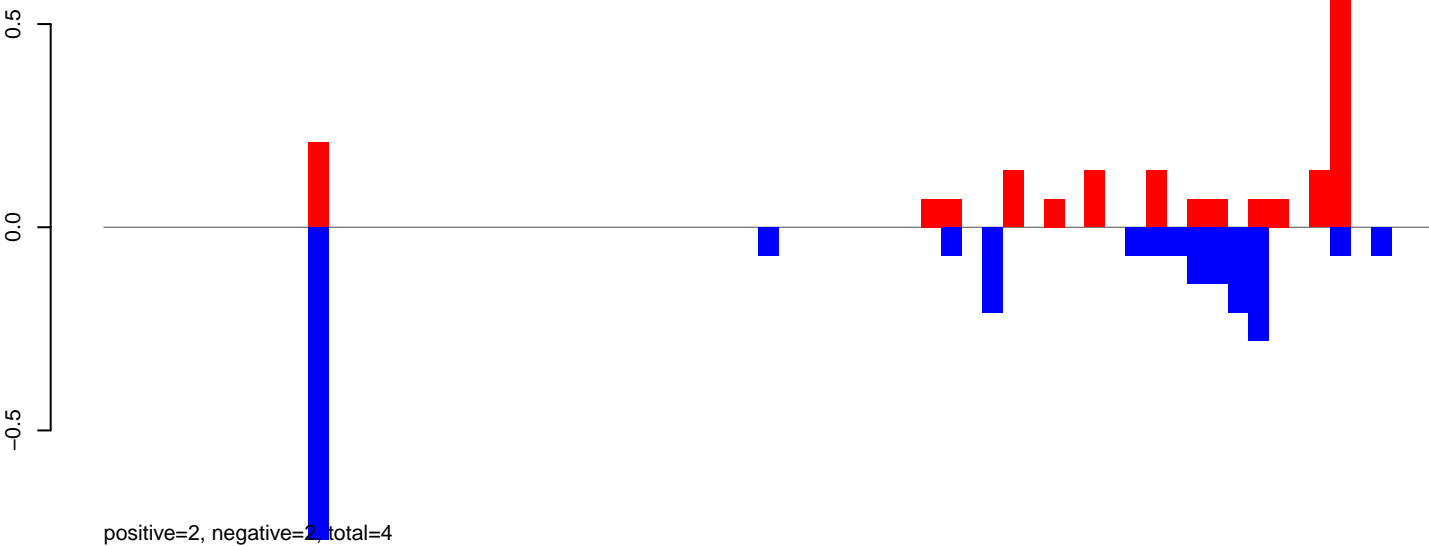
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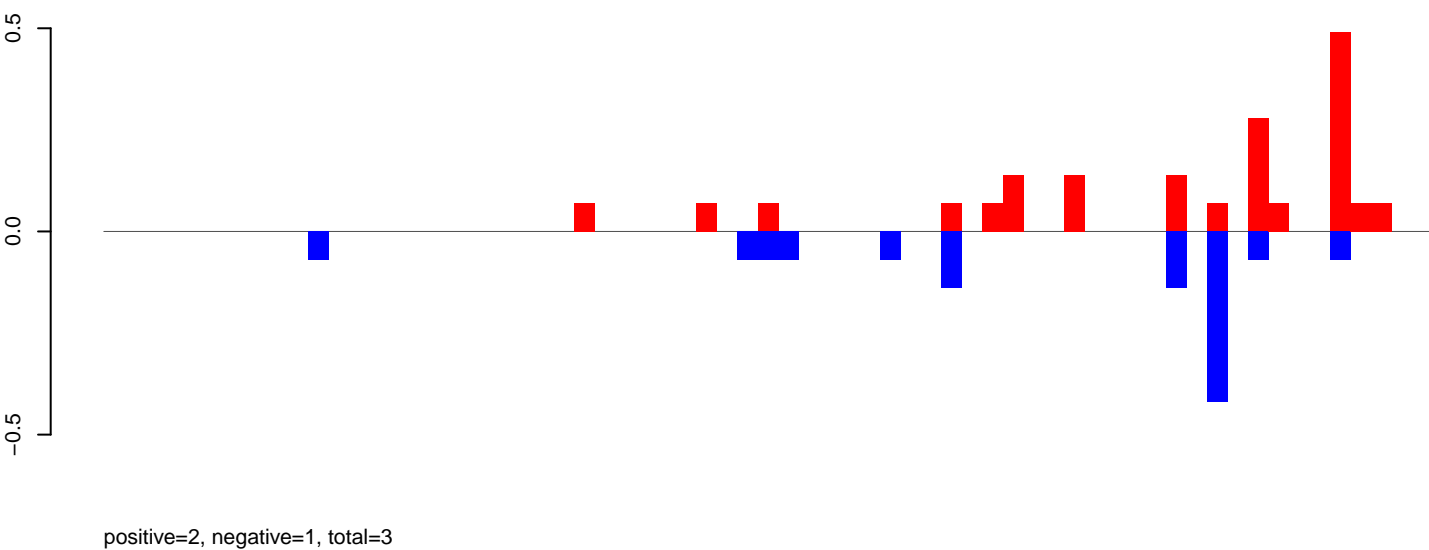
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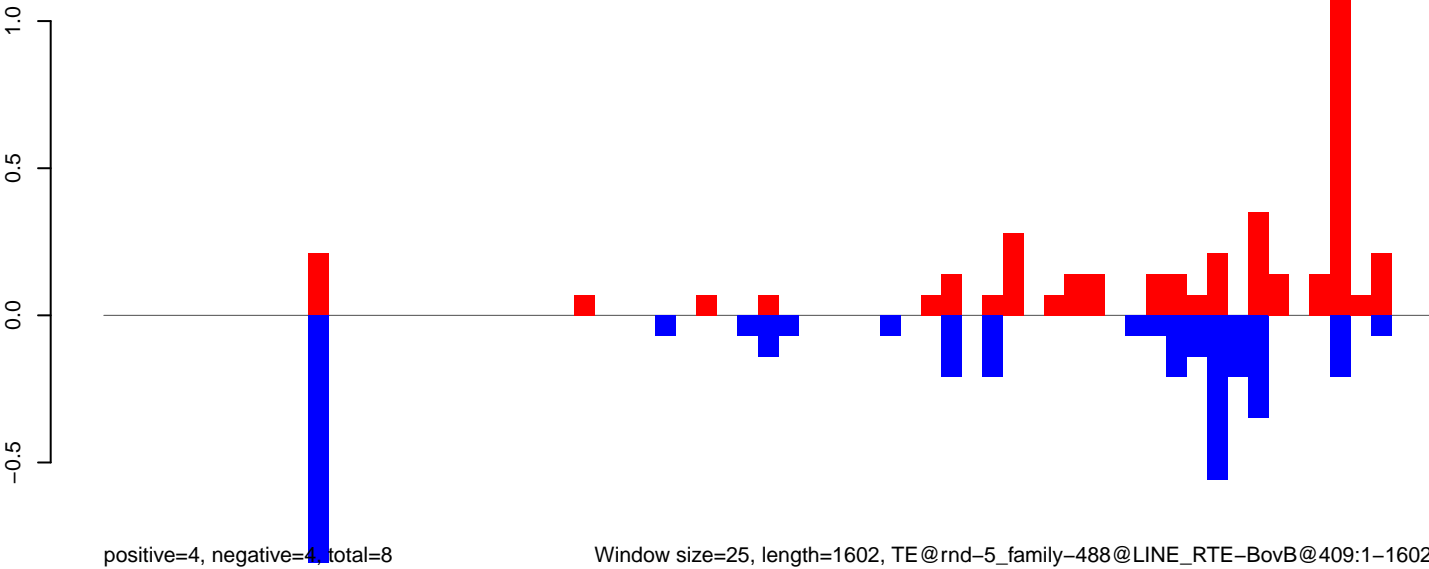
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CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep



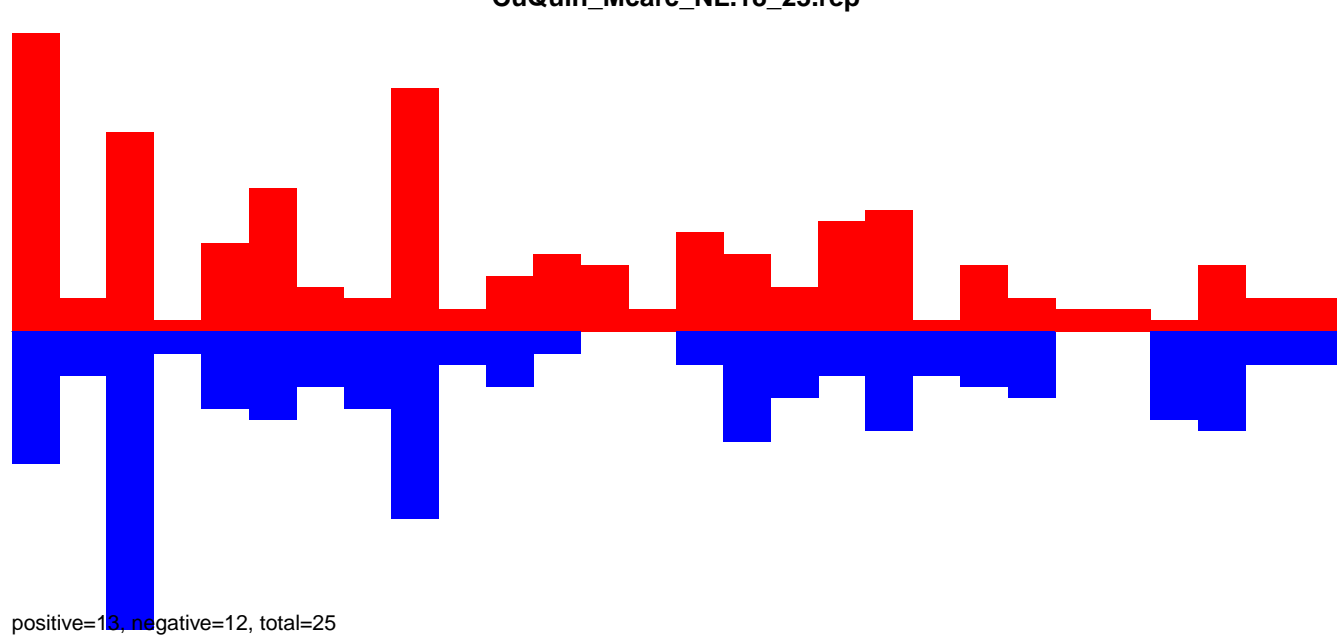
CuQuin_Mcarc_NL.24_35.rep



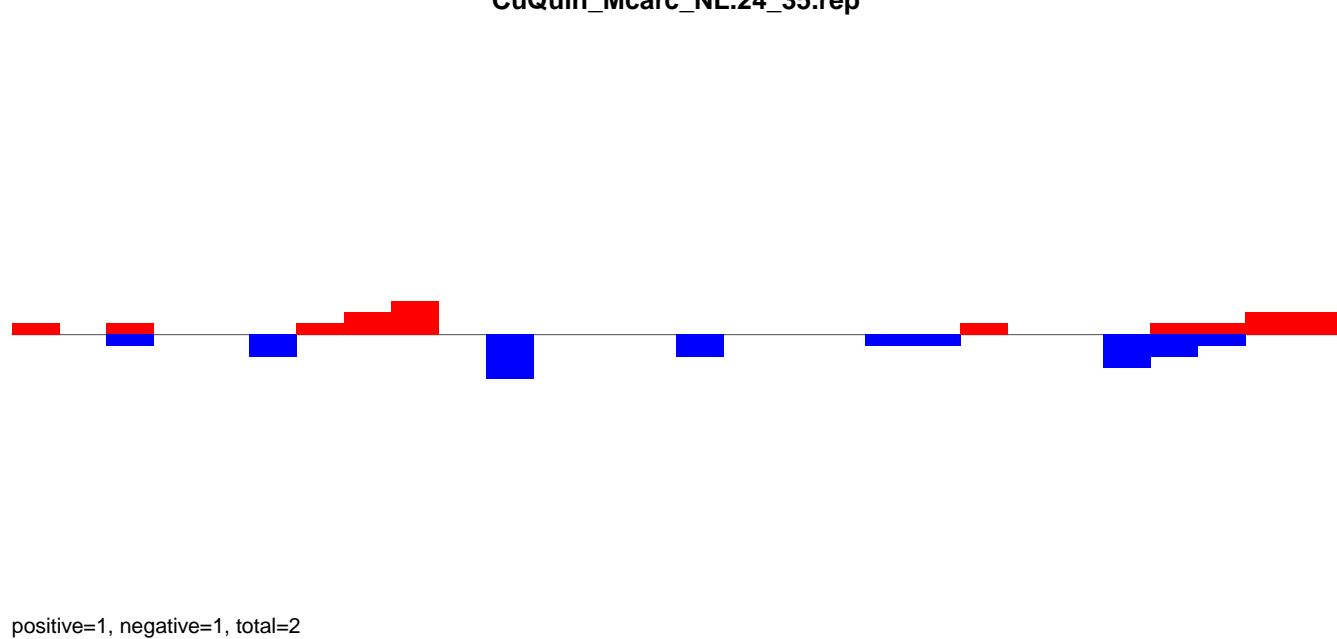
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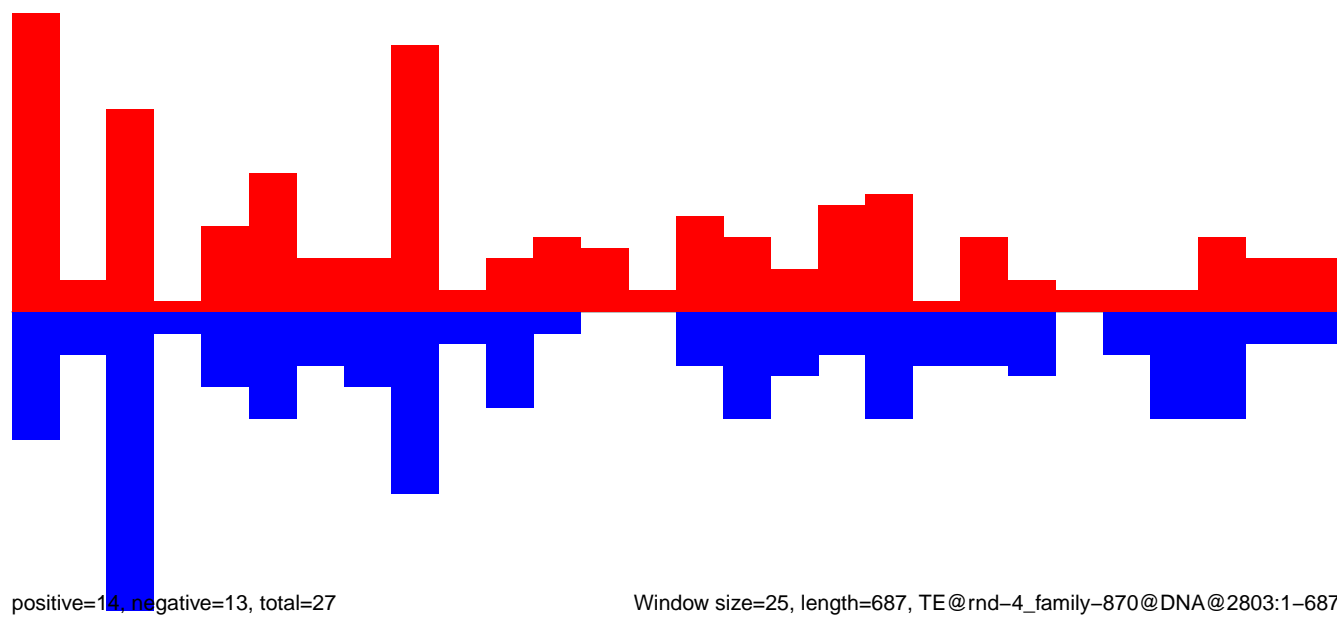
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

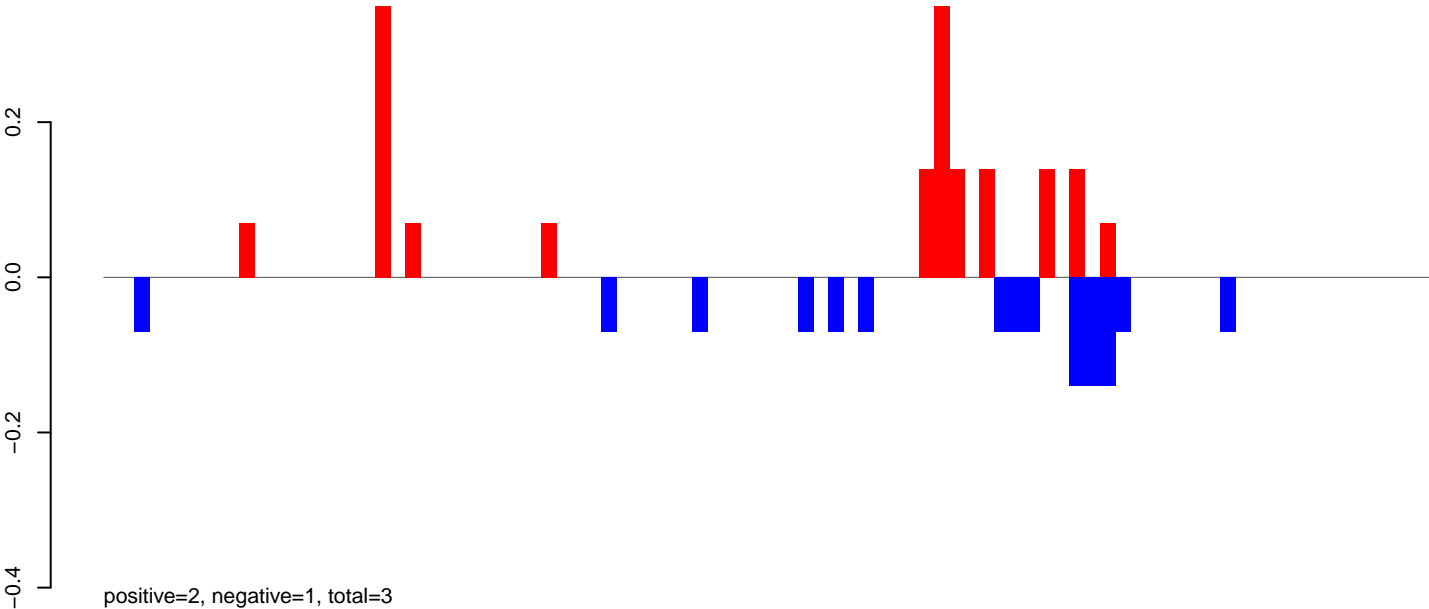


CuQuin_Mcarc_NL.rep

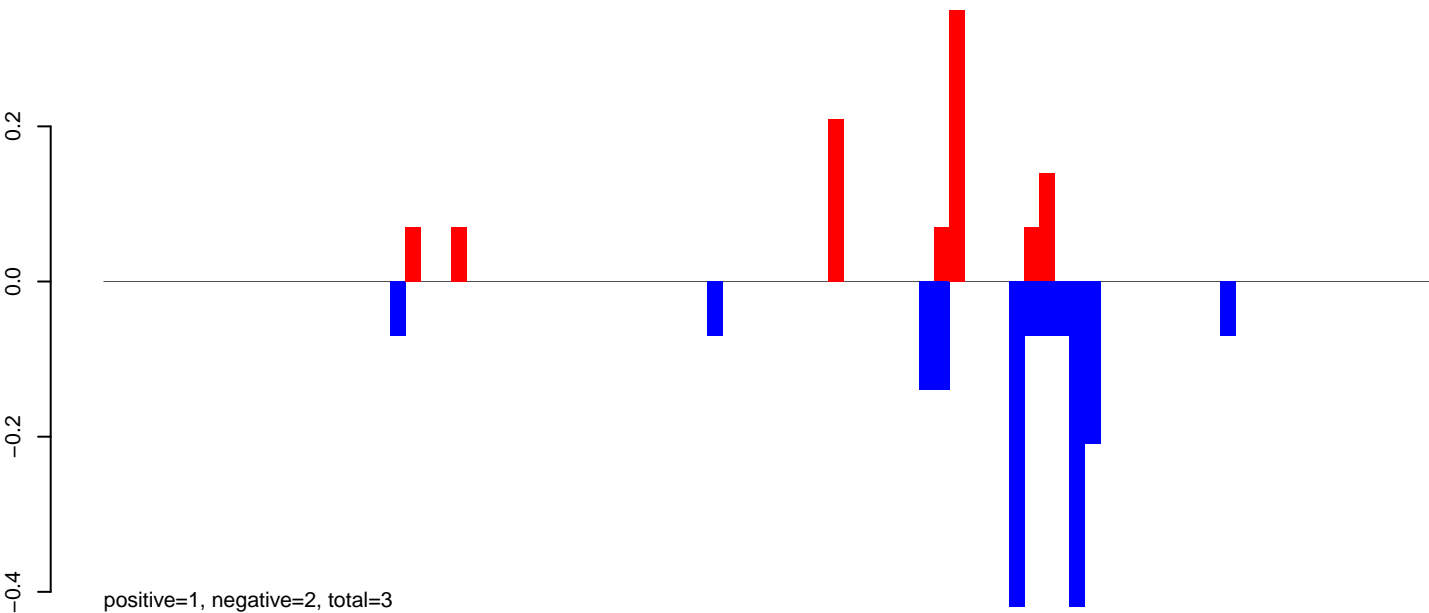


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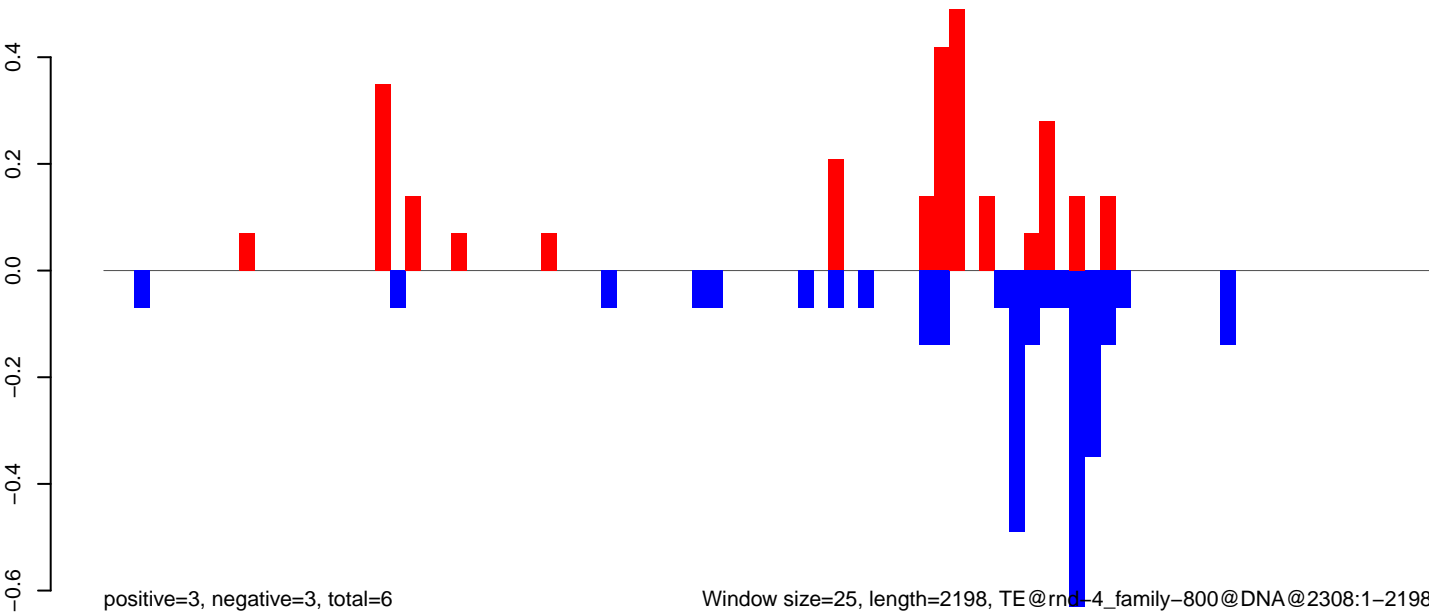
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



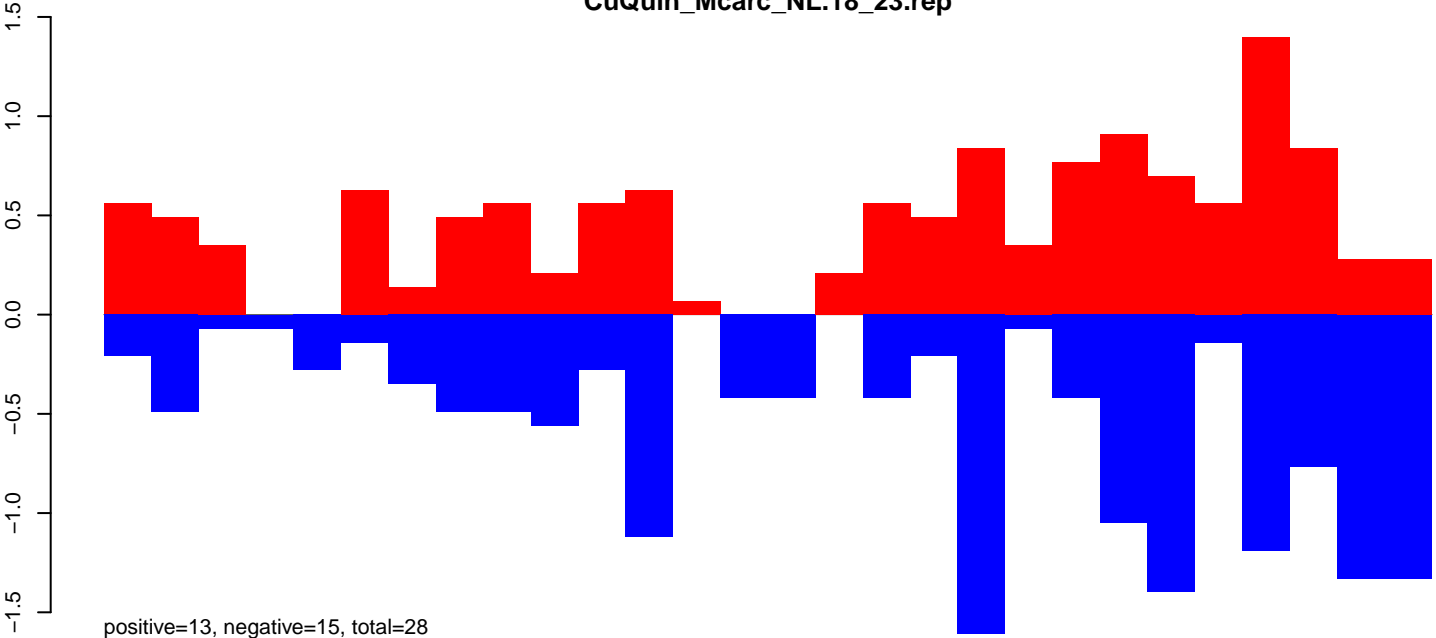
CuQuin_Mcarc_NL.rep



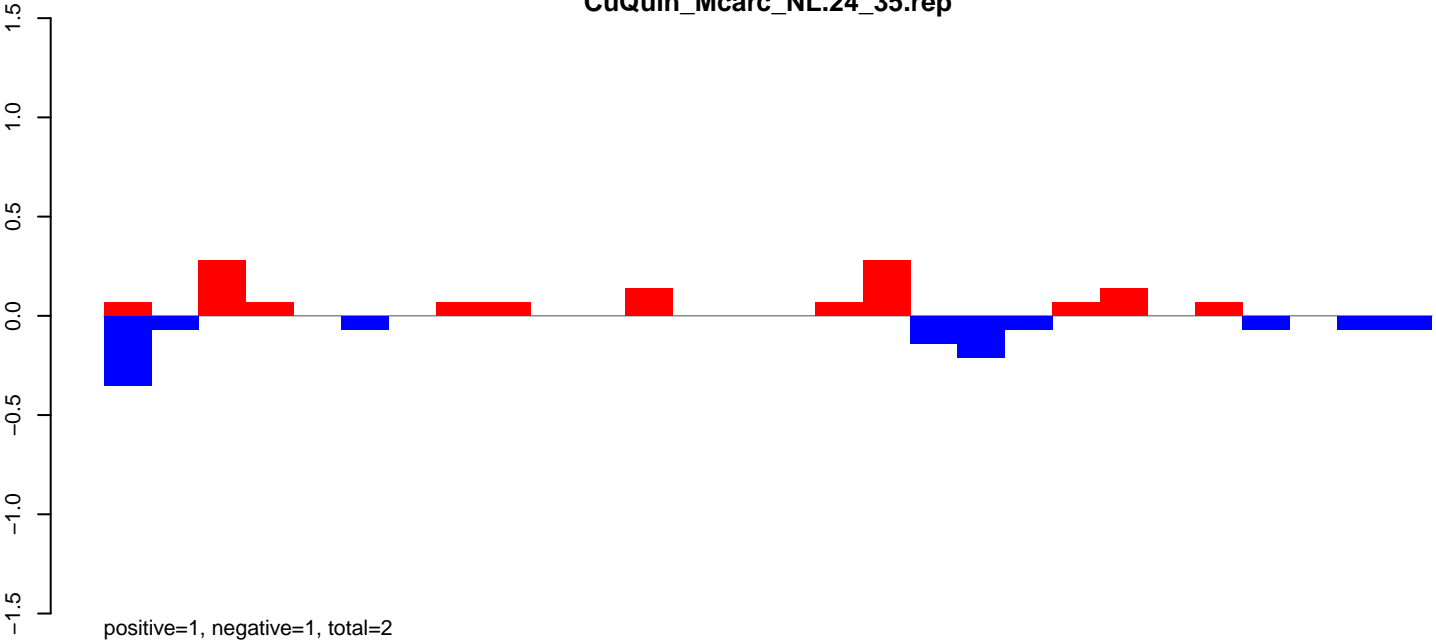
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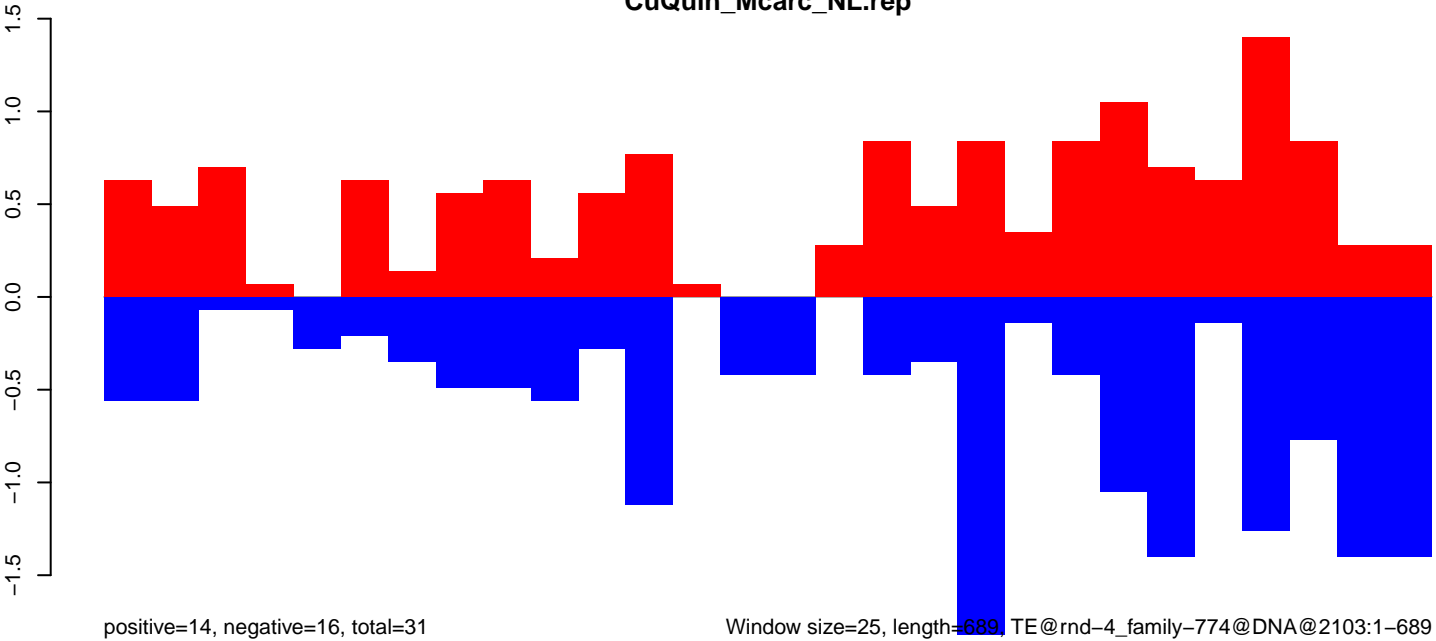
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

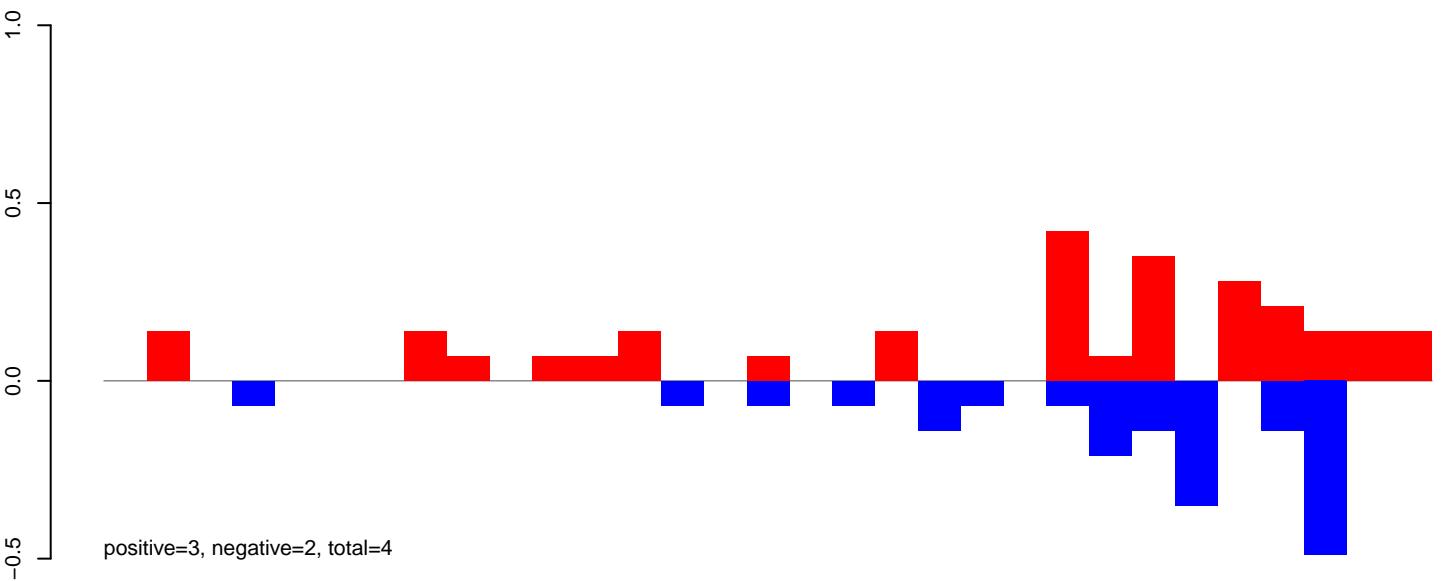


CuQuin_Mcarc_NL.rep

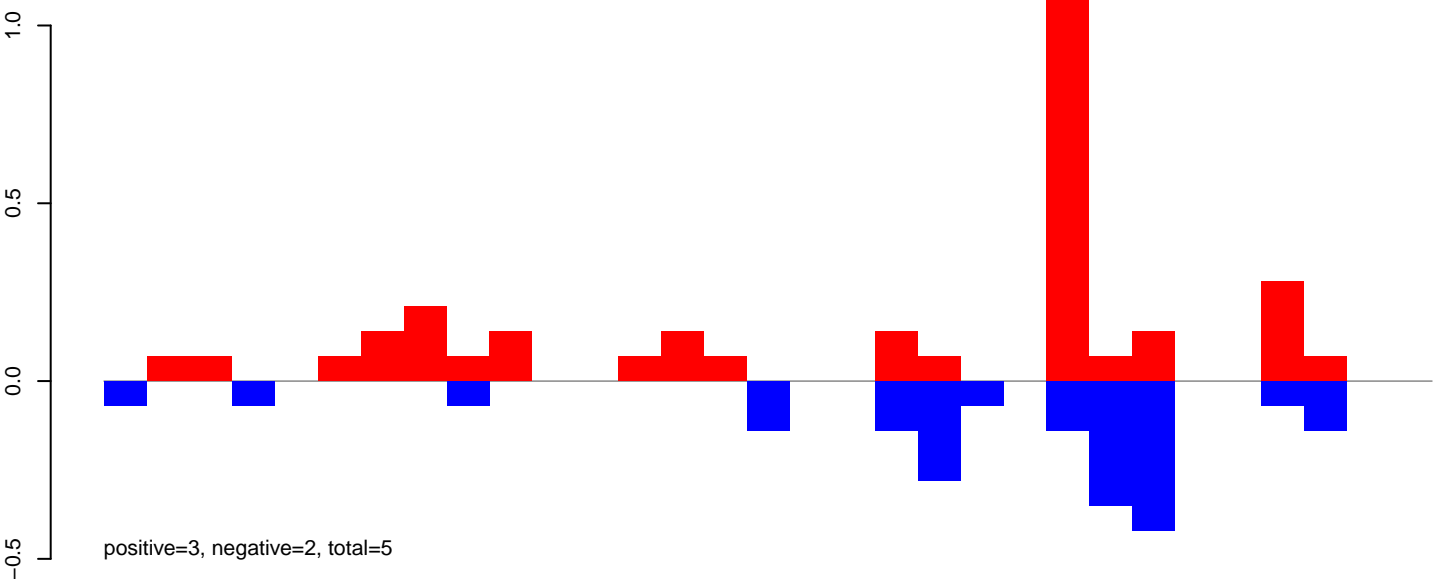


Window size=25, length=689, TE@rnd-4_family-774@DNA@2103:1-689

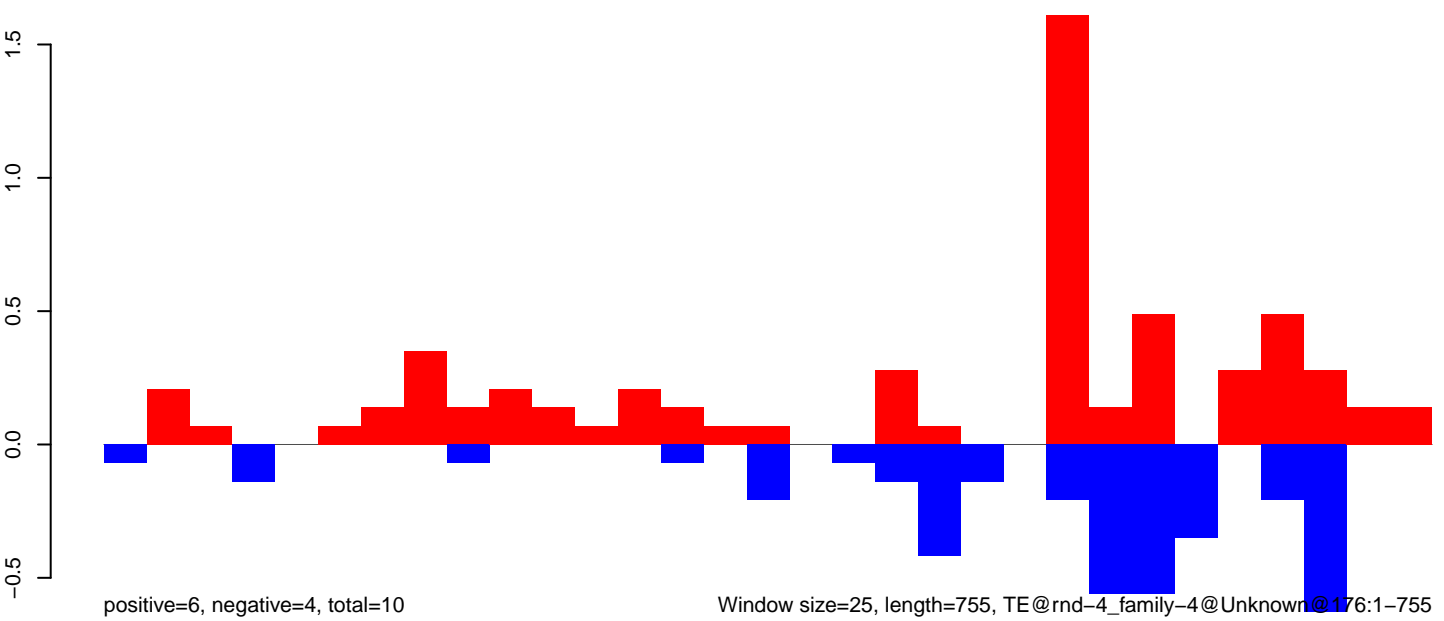
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

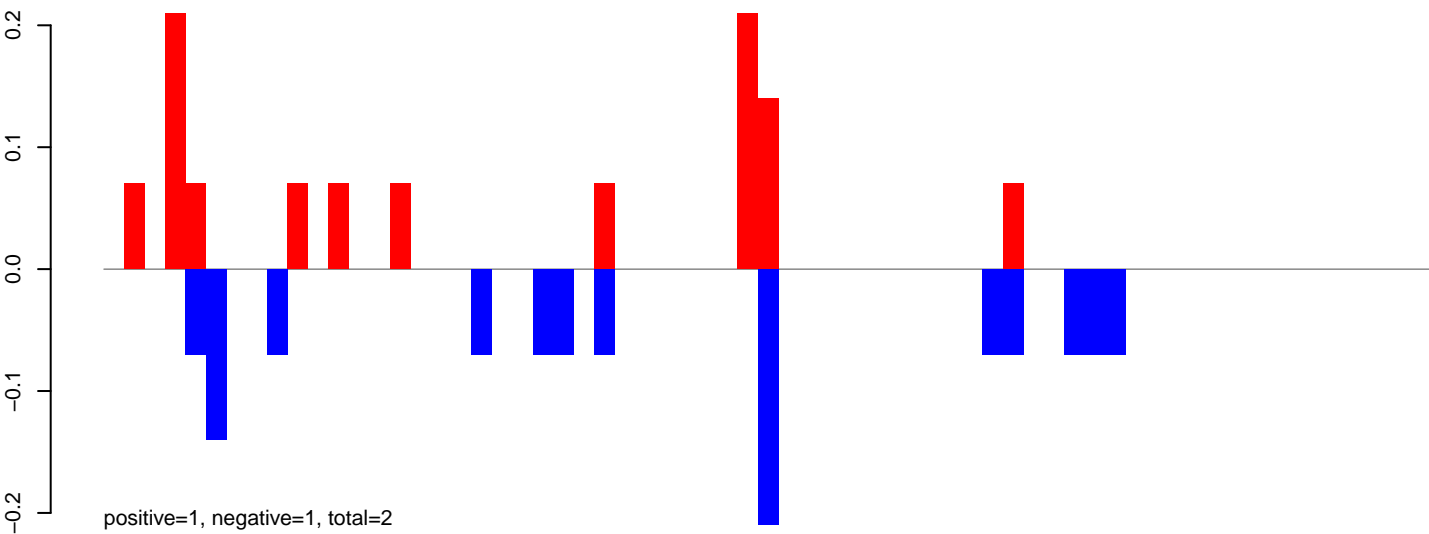


CuQuin_Mcarc_NL.rep

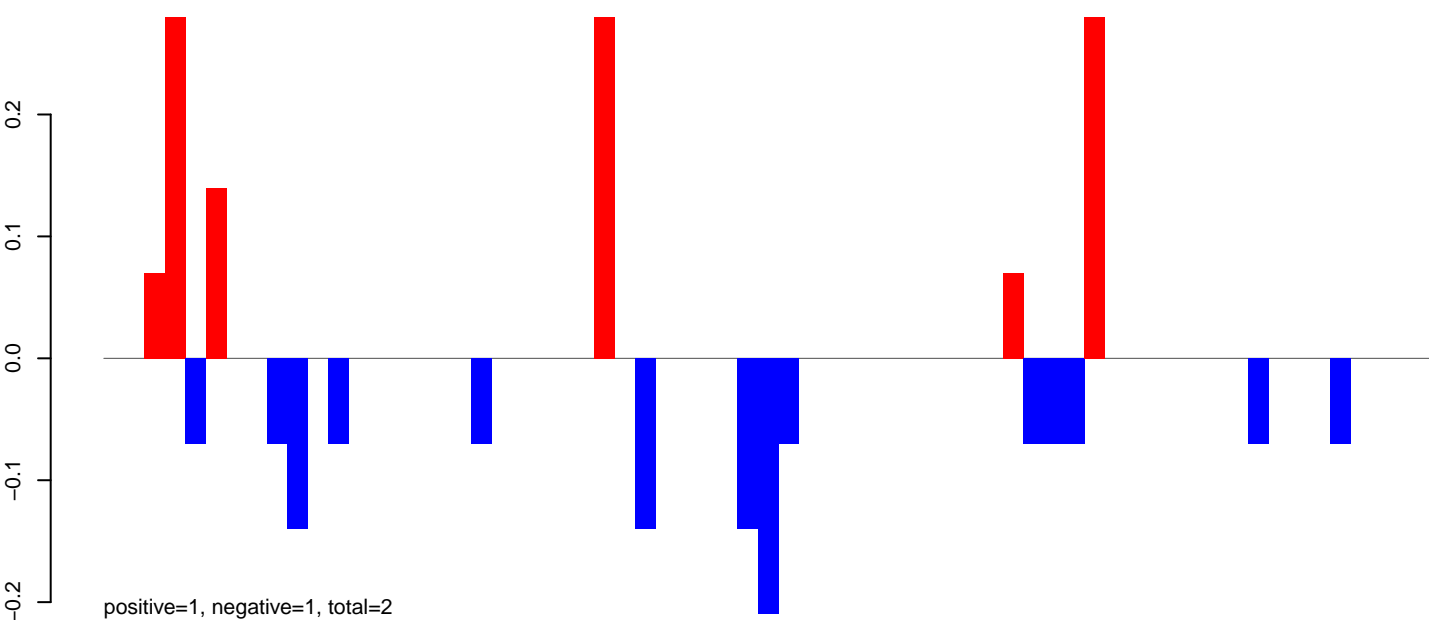


Window size=25, length=755, TE@rnd-4_family-4@Unknown@176:1-755

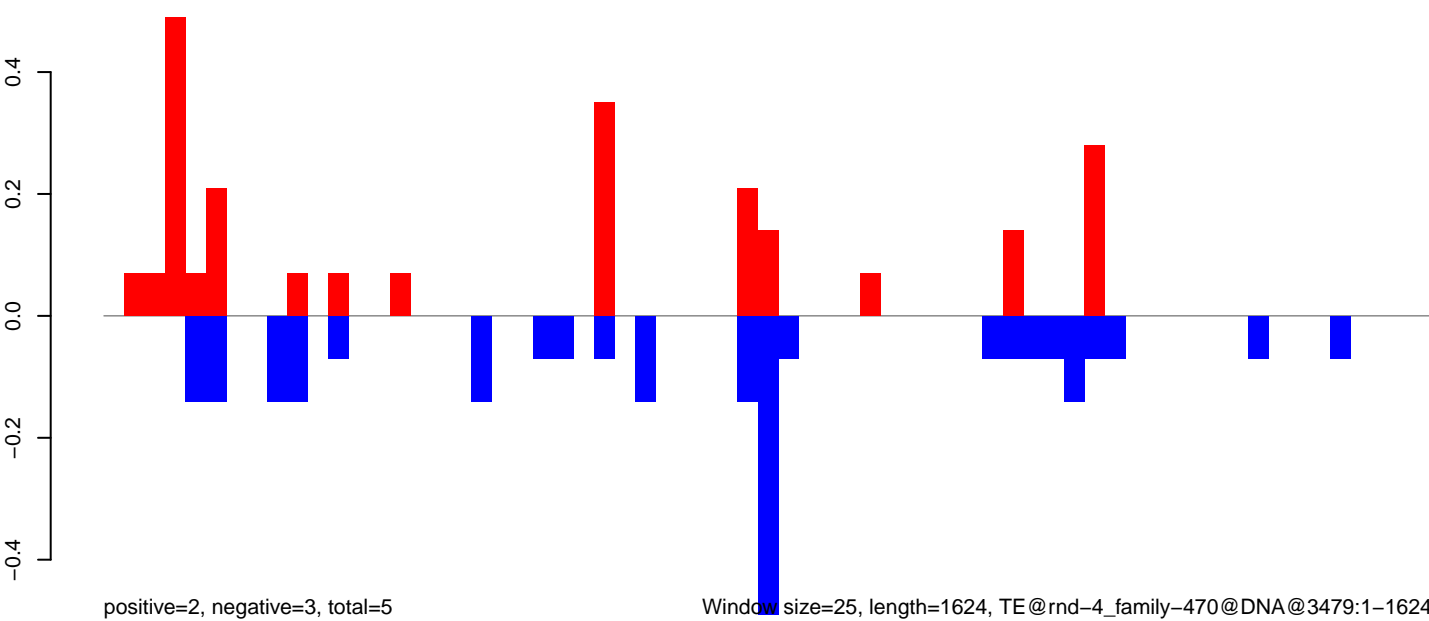
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

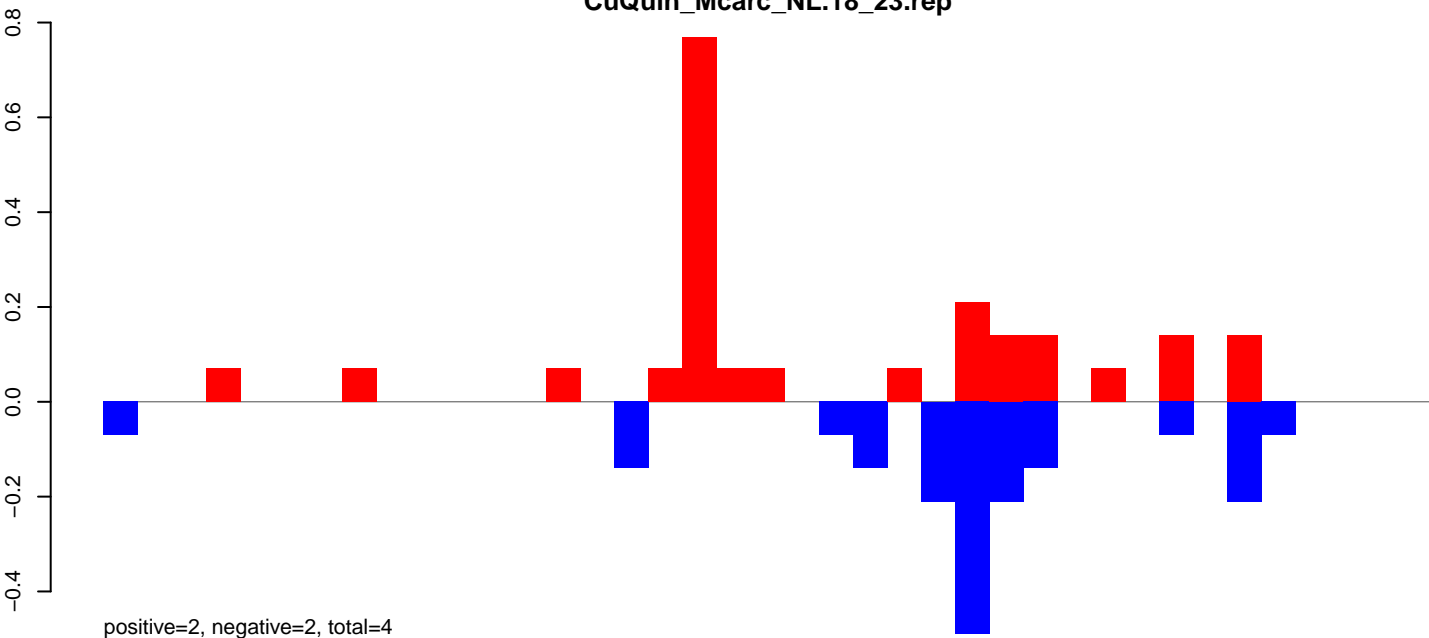


CuQuin_Mcarc_NL.rep

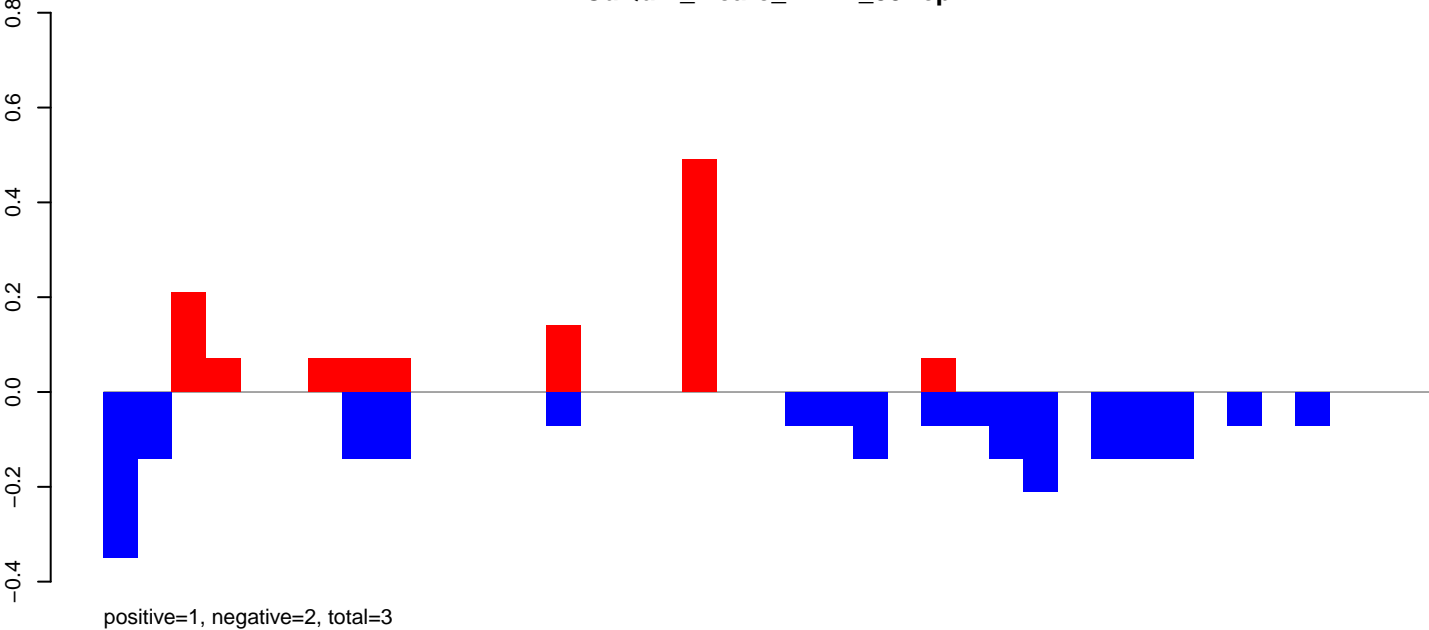


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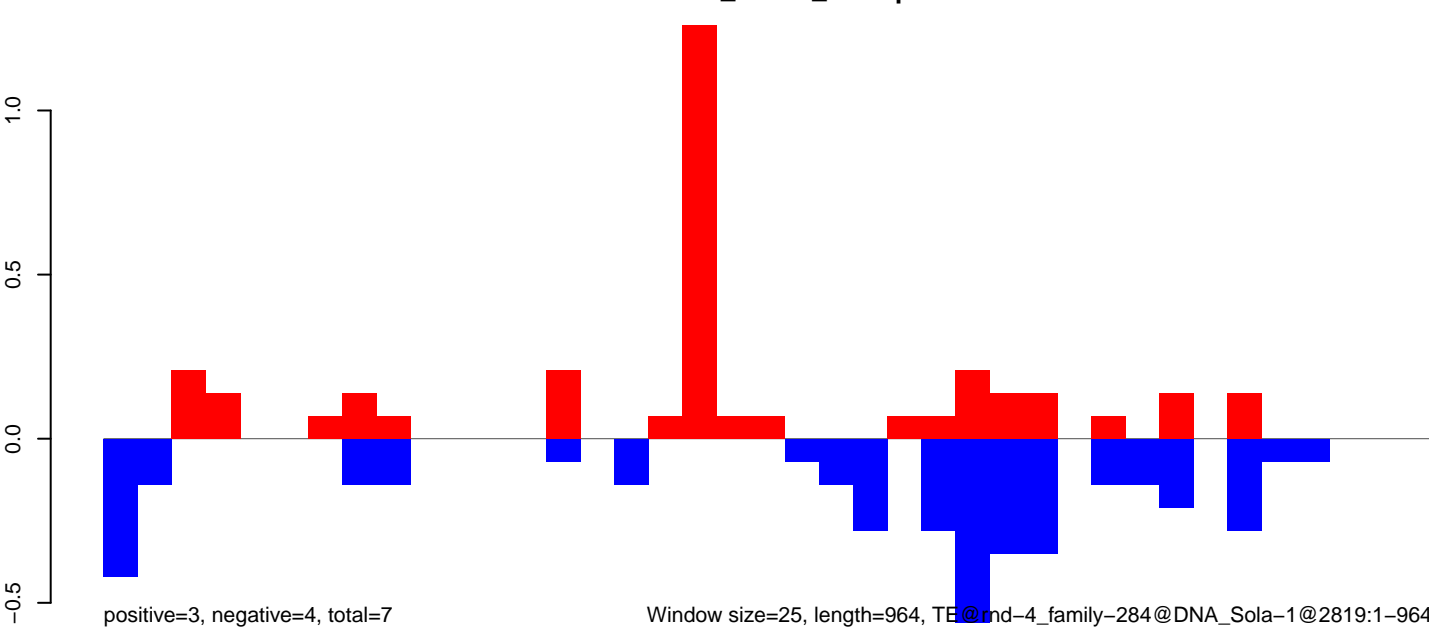
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

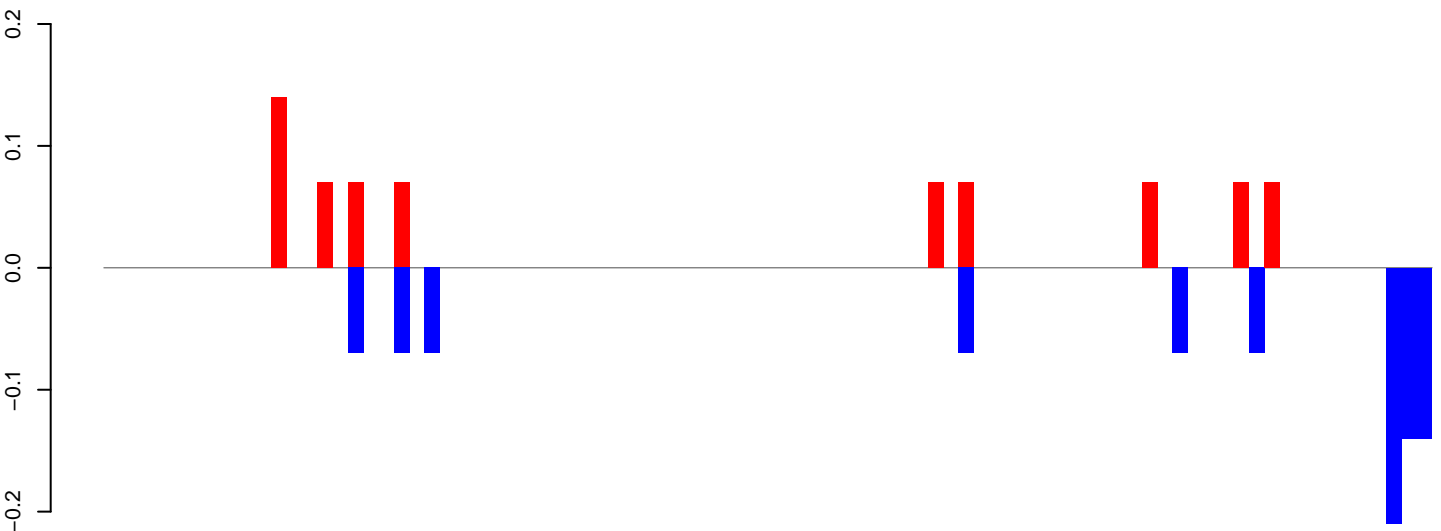


CuQuin_Mcarc_NL.rep



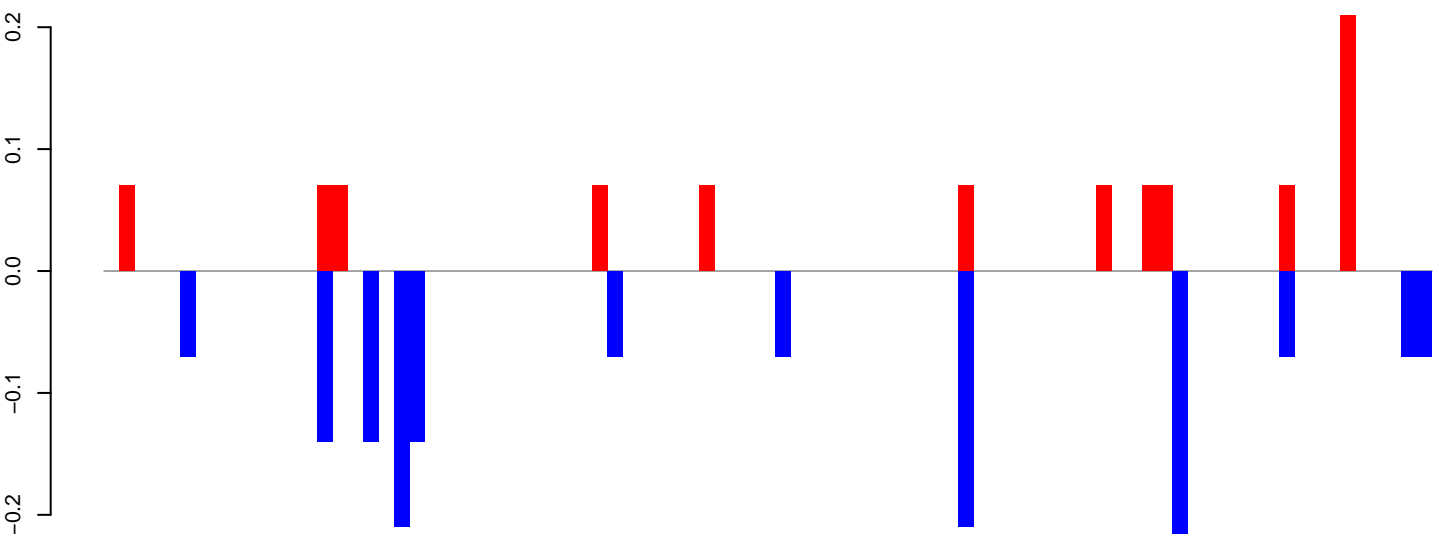
Window size=25, length=964, TE@rnd-4_family-284@DNA_Sola-1@2819:1-964

CuQuin_Mcarc_NL.18_23.rep



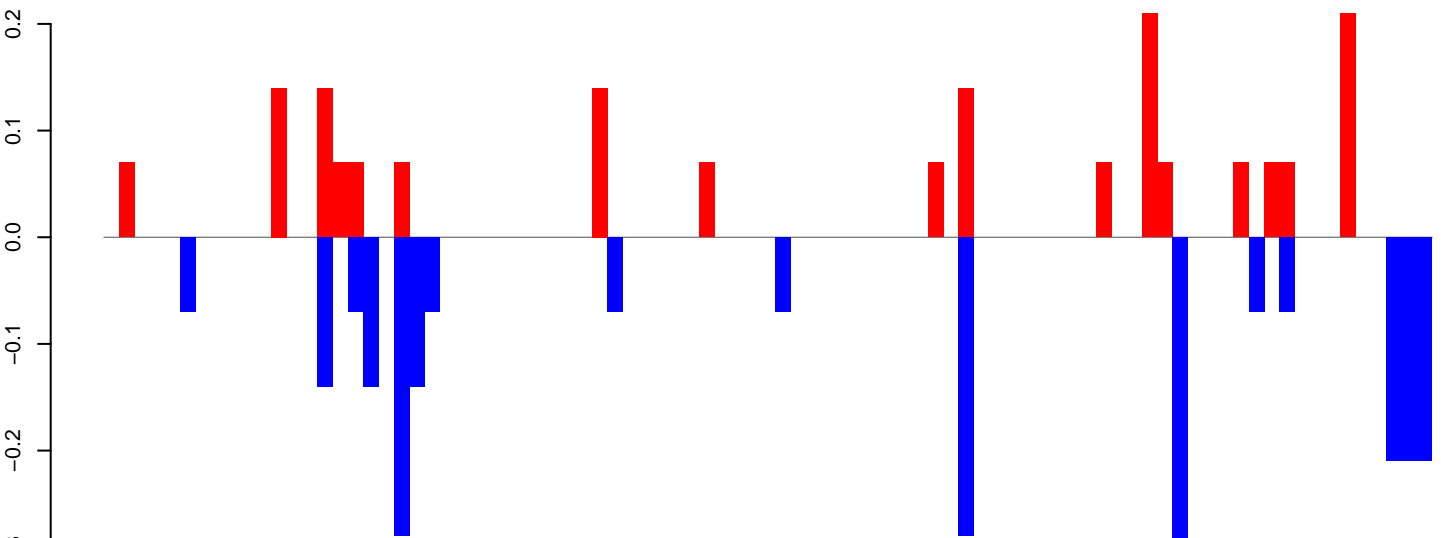
positive=1, negative=1, total=2

CuQuin_Mcarc_NL.24_35.rep



positive=1, negative=2, total=2

CuQuin_Mcarc_NL.rep

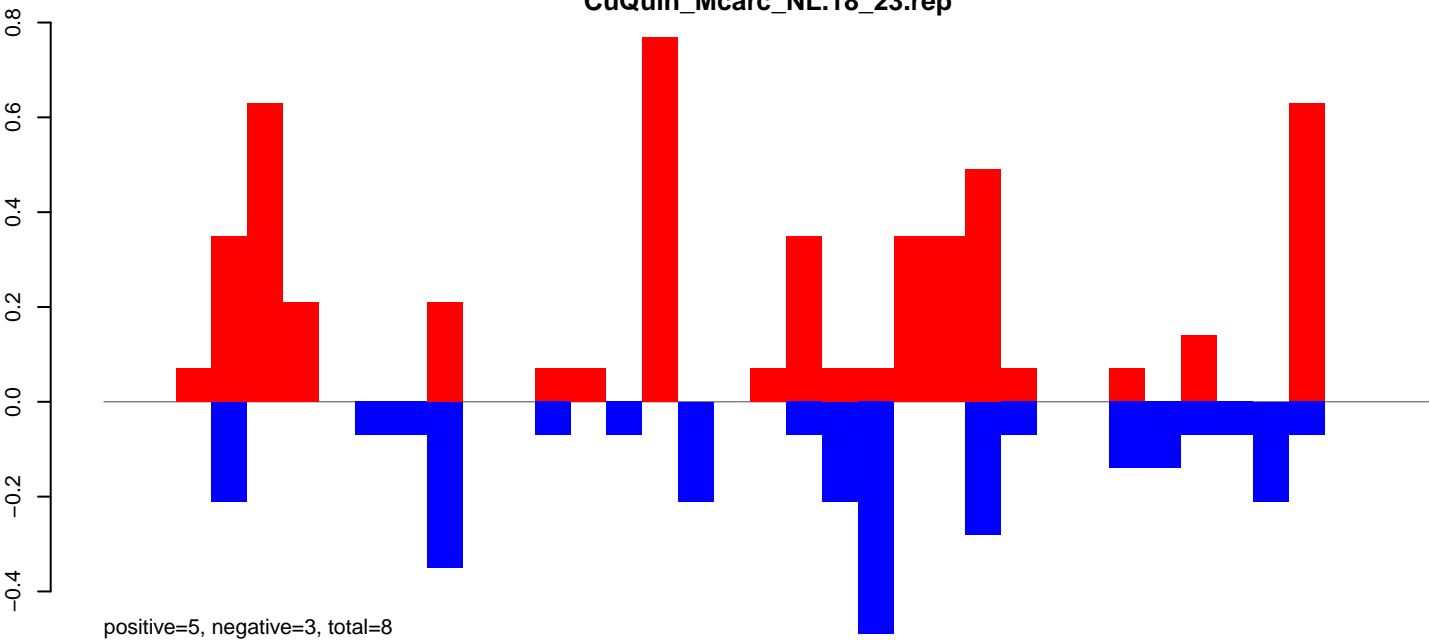


positive=2, negative=2, total=4

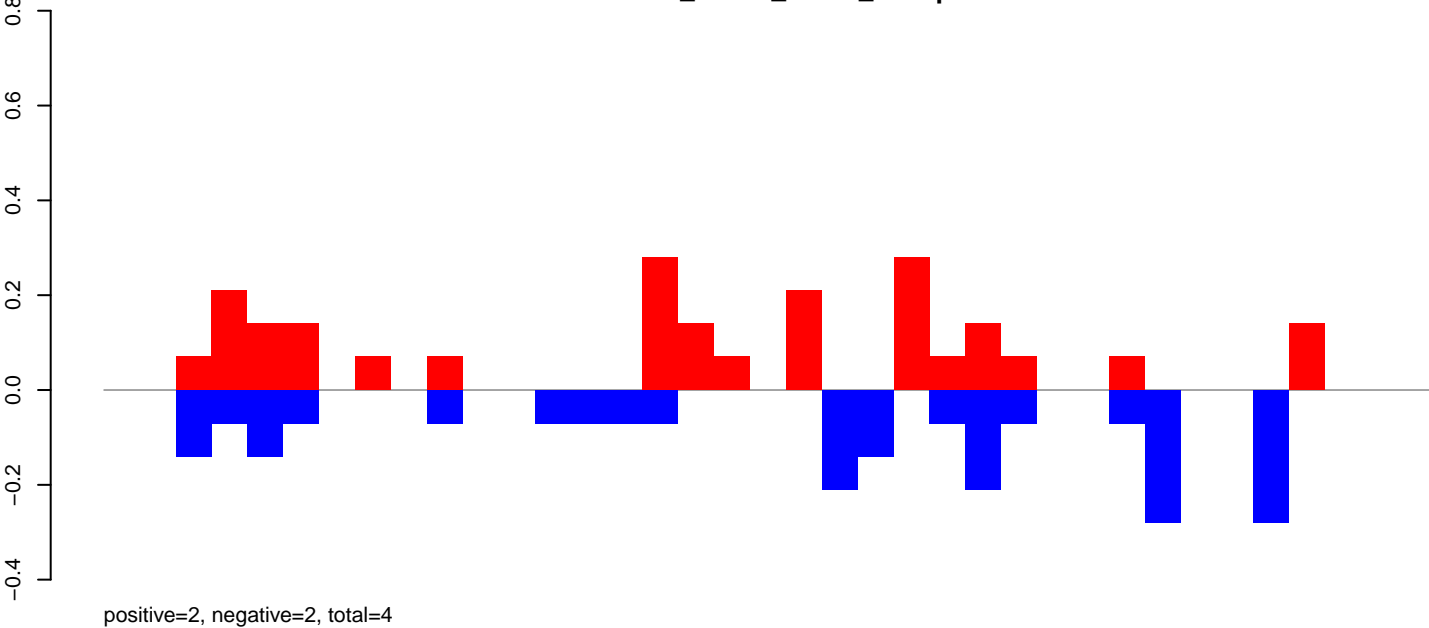
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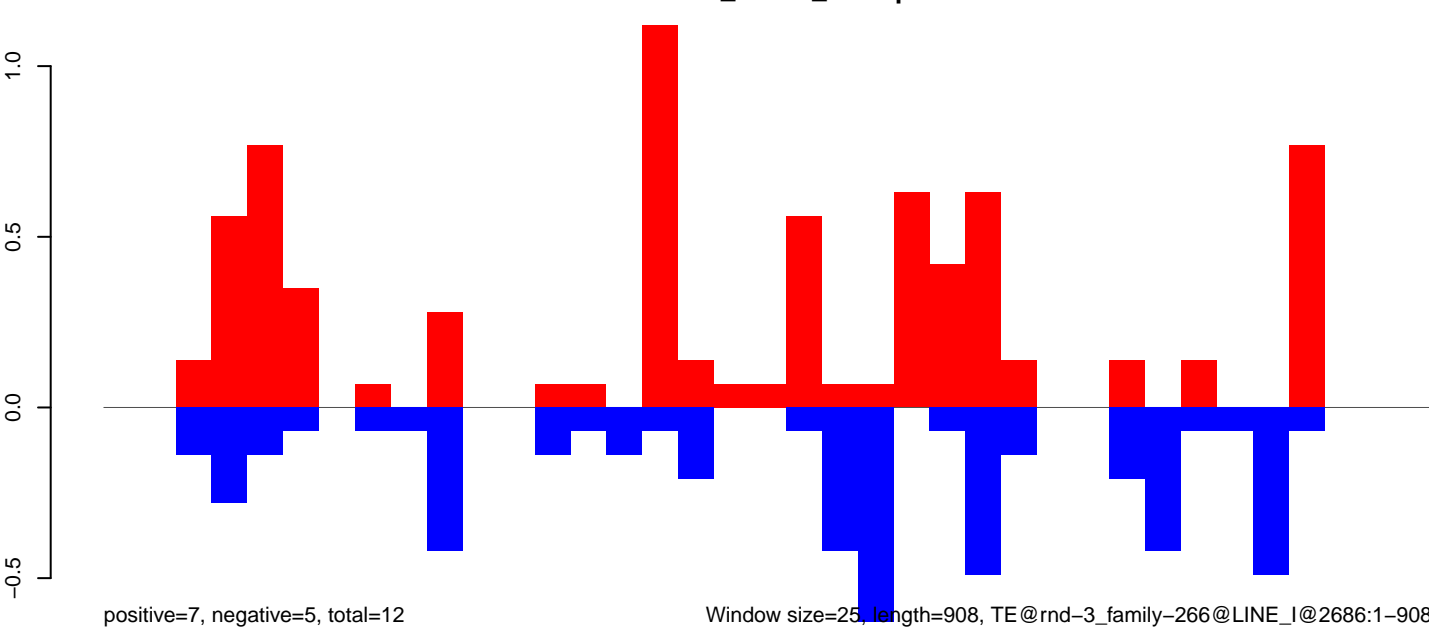
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



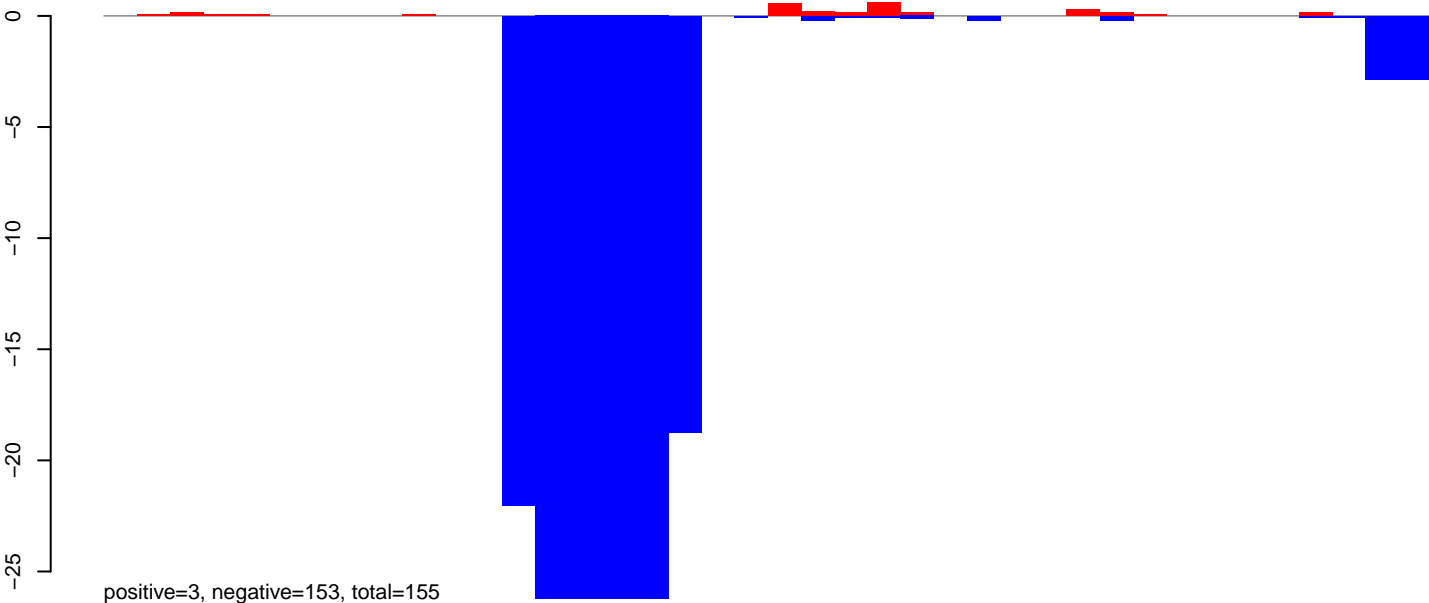
CuQuin_Mcarc_NL.rep



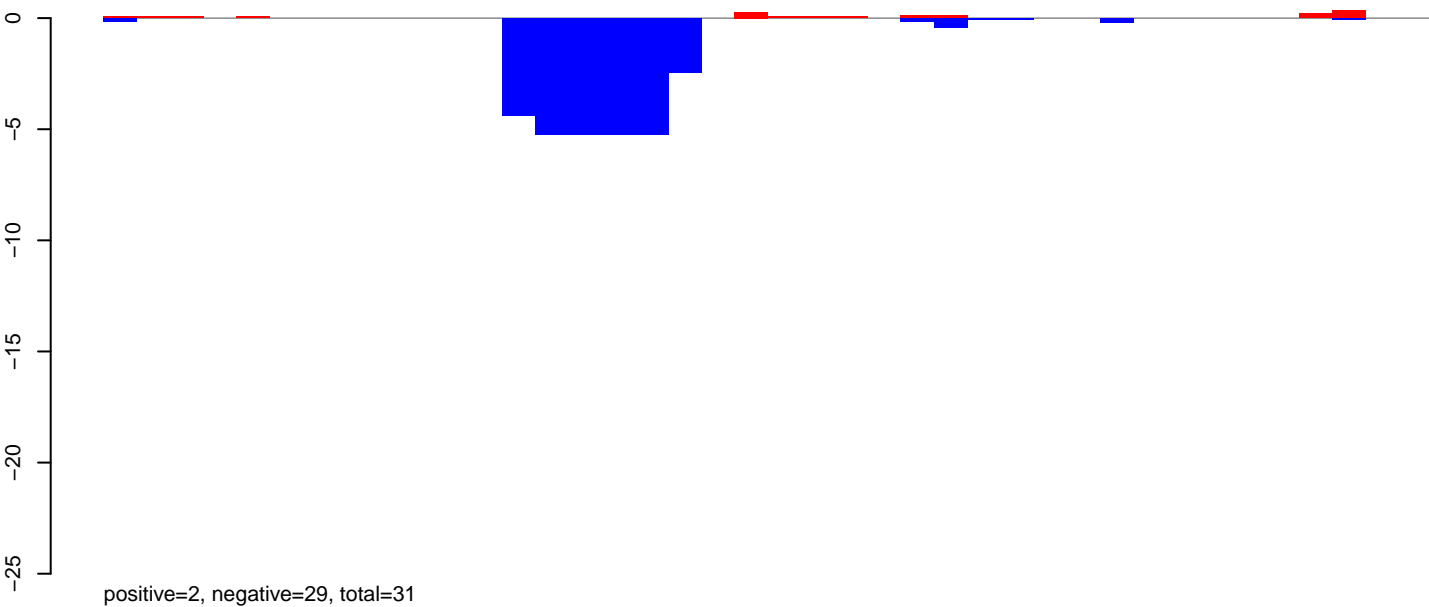
Window size=25, length=908, TE@rnd-3_family-266@LINE_I@2686:1-908

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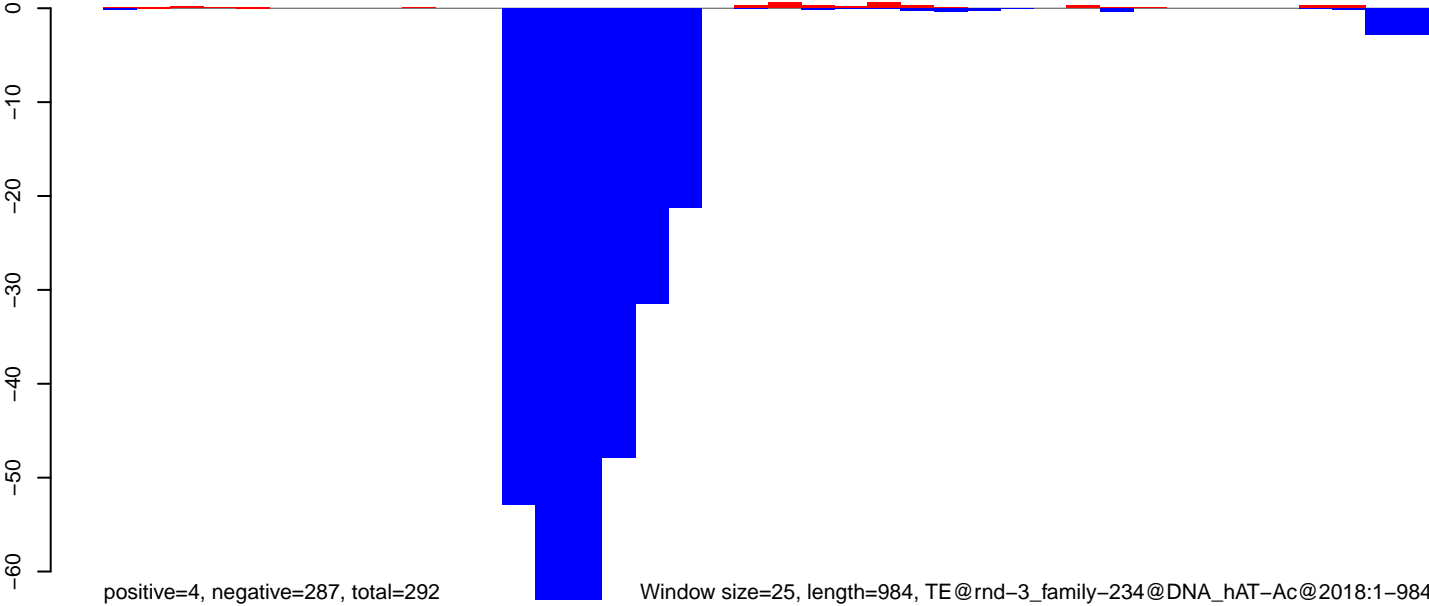
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

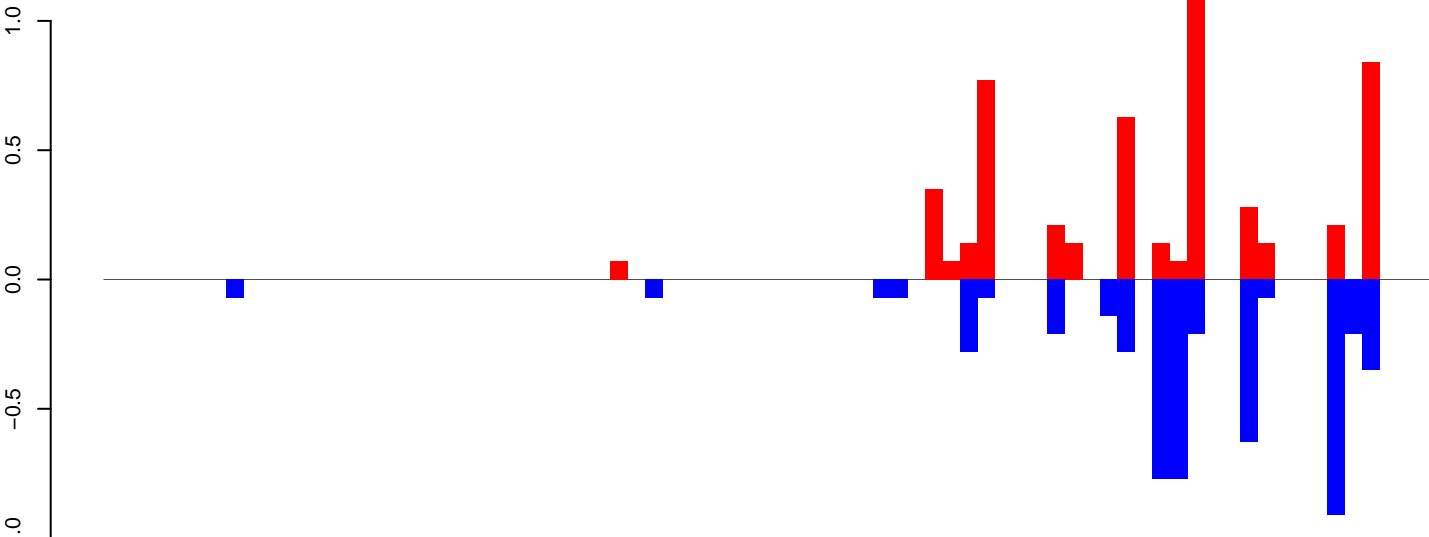


CuQuin_Mcarc_NL.rep



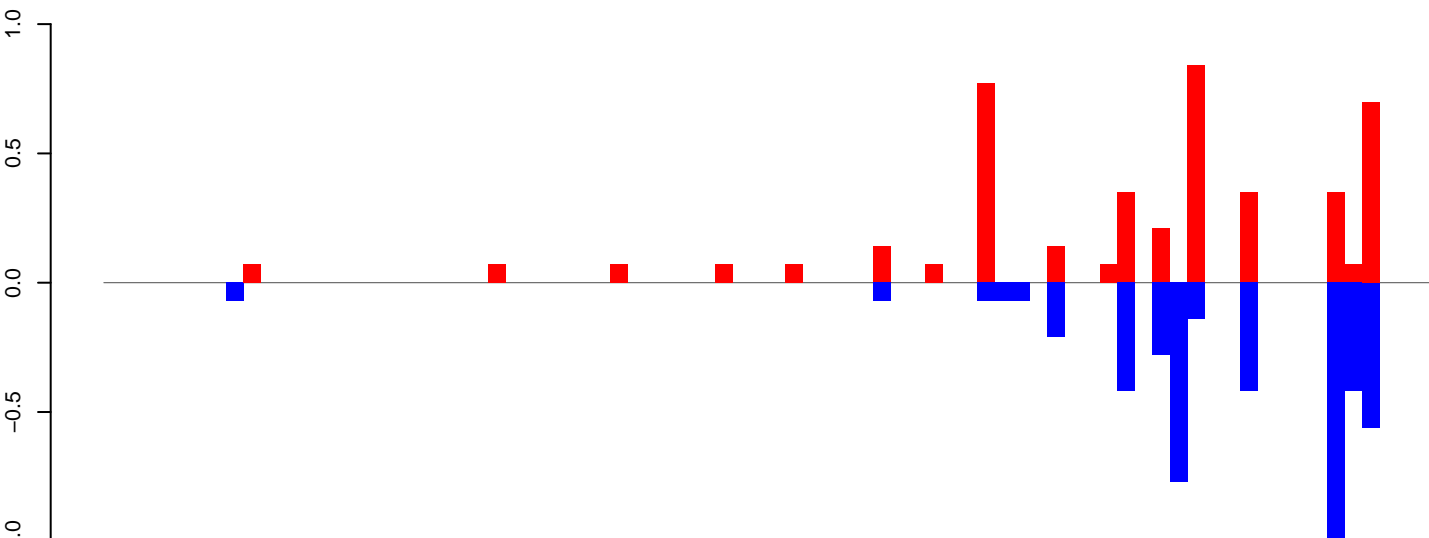
Window size=25, length=984, TE@rnd-3_family-234@DNA_hAT-Ac@2018:1-984

CuQuin_Mcarc_NL.18_23.rep



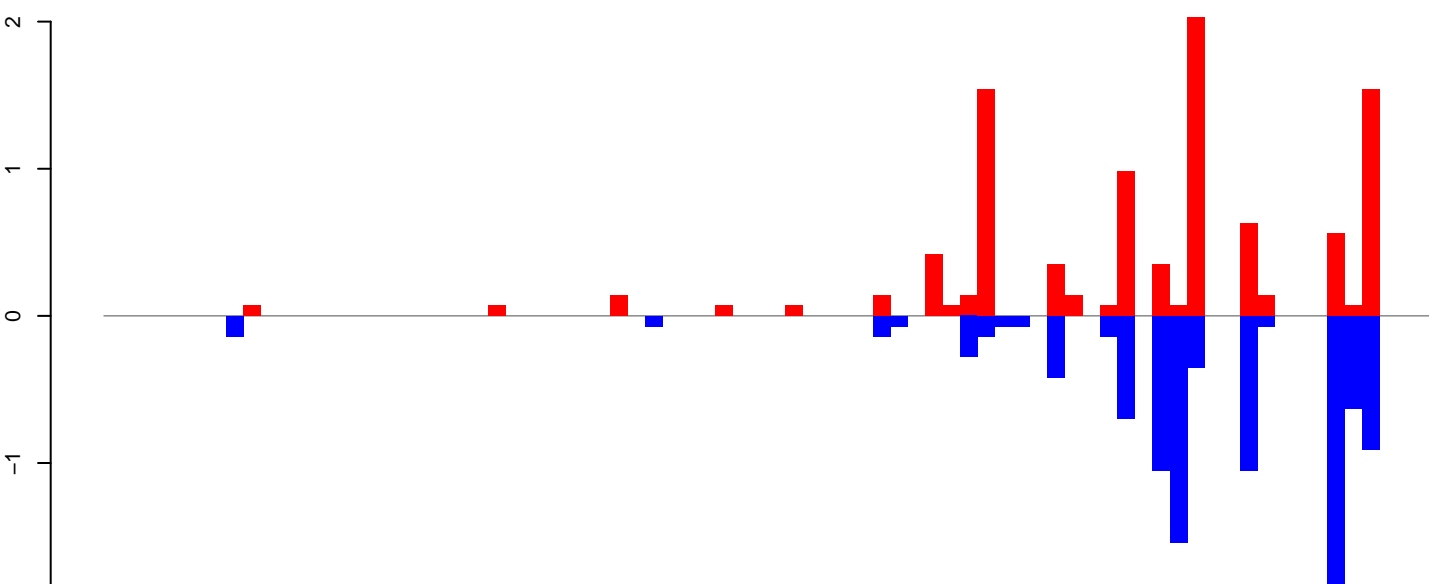
positive=5, negative=5, total=10

CuQuin_Mcarc_NL.24_35.rep



positive=4, negative=5, total=9

CuQuin_Mcarc_NL.rep

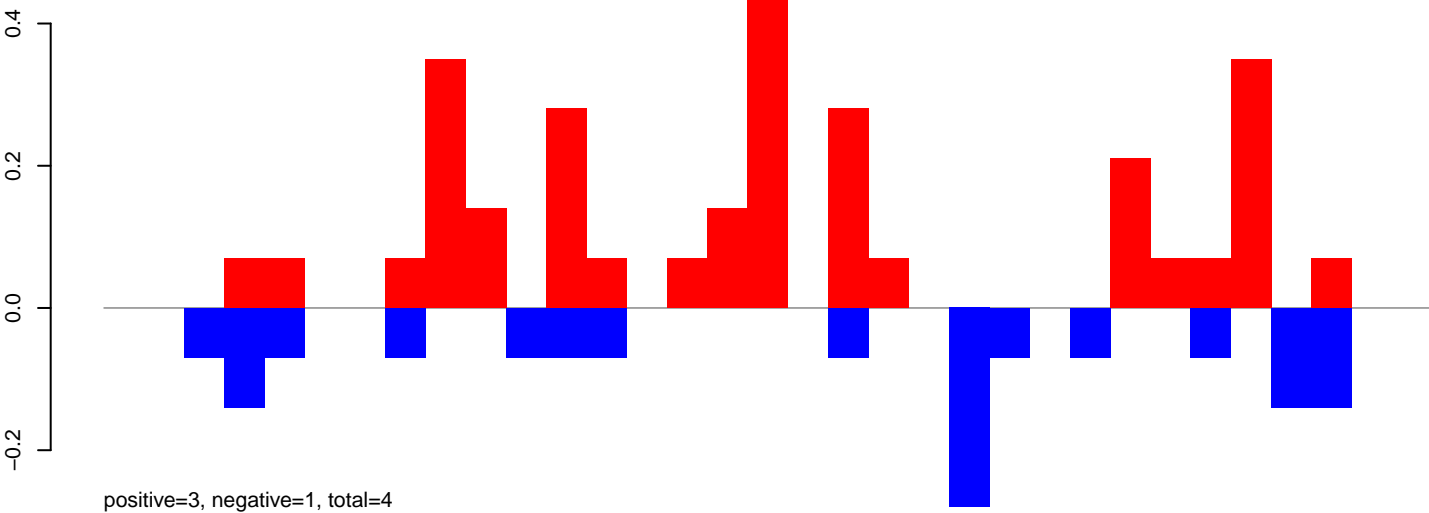


positive=10, negative=10, total=20

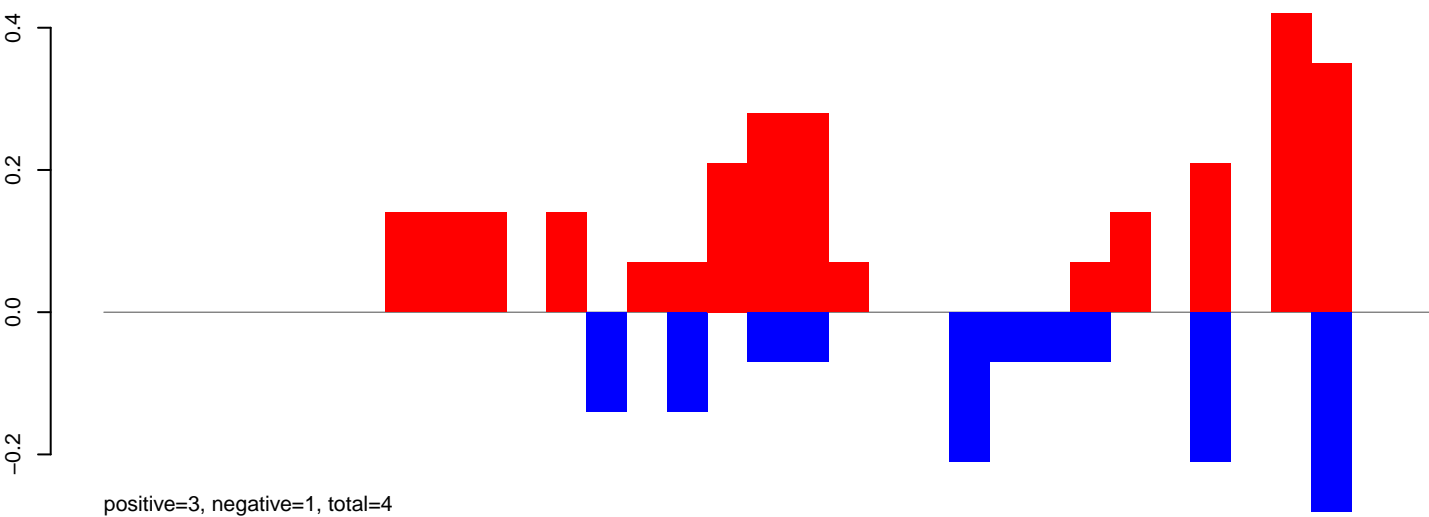
Window size=25, length=1885, TE@rnd-3_family-161@DNA_MULE-MuDR@3330:1-1885

0 500 1000 1500 2000

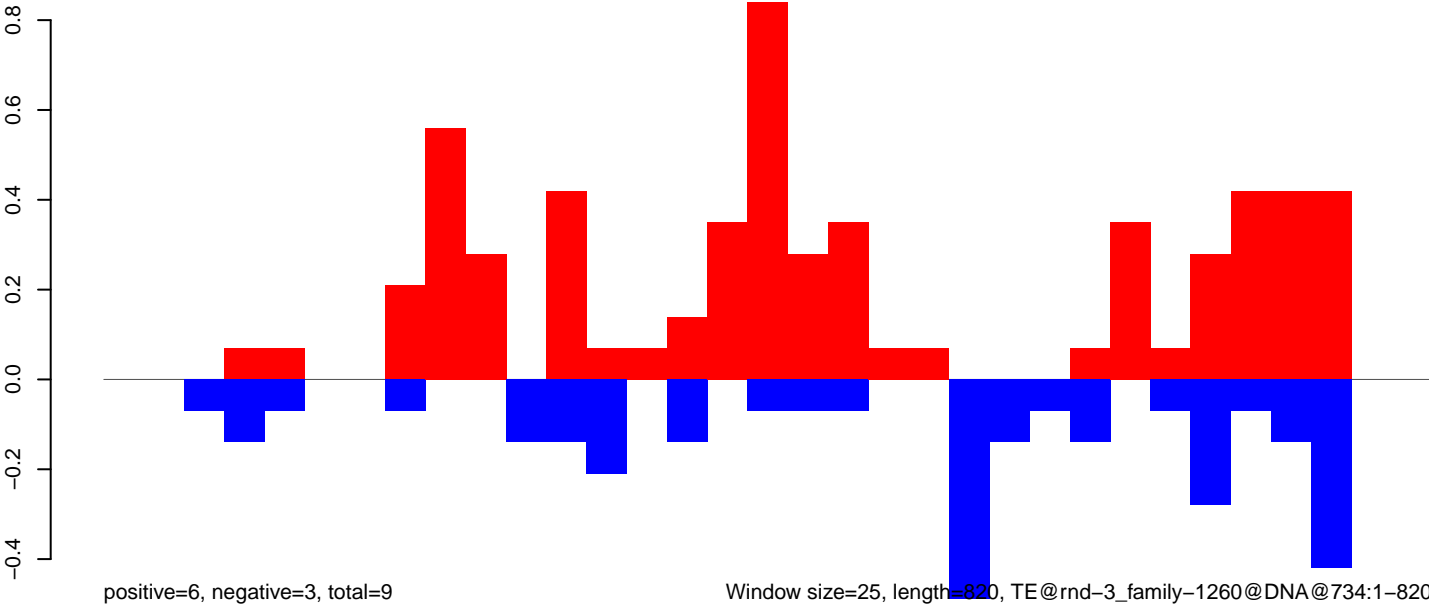
CuQuin_Mcarc_NL.18_23.rep



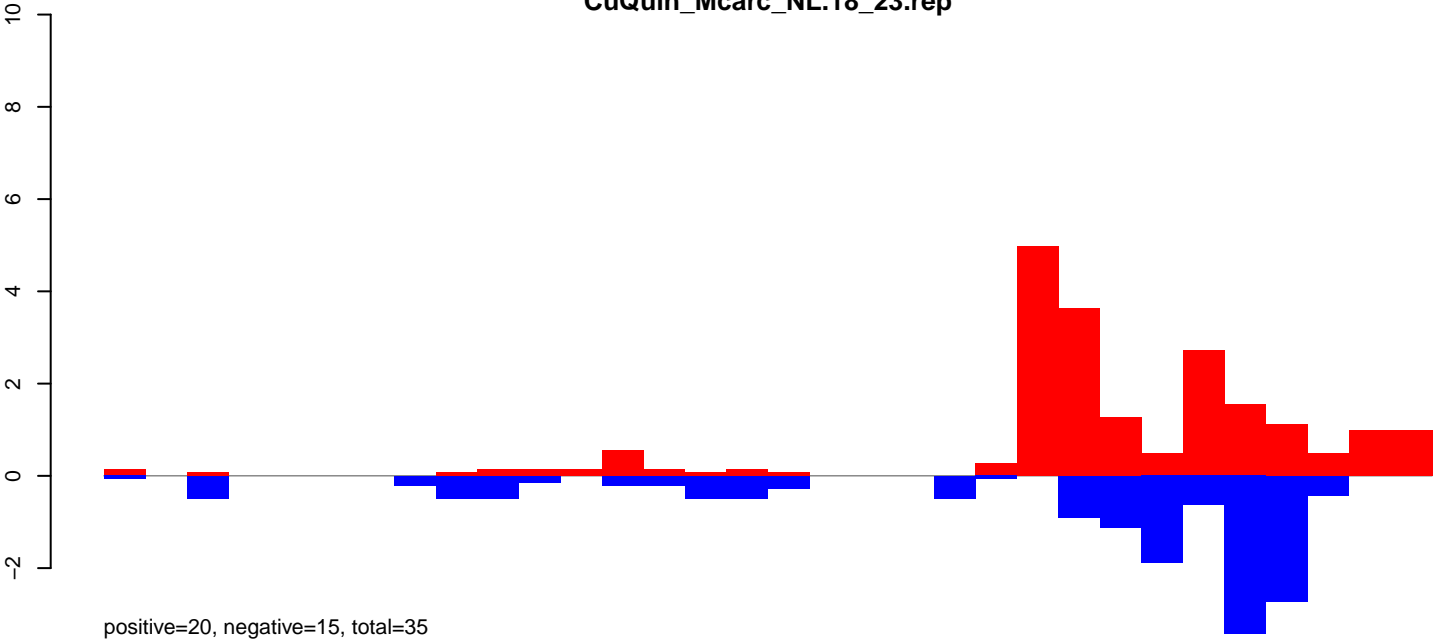
CuQuin_Mcarc_NL.24_35.rep



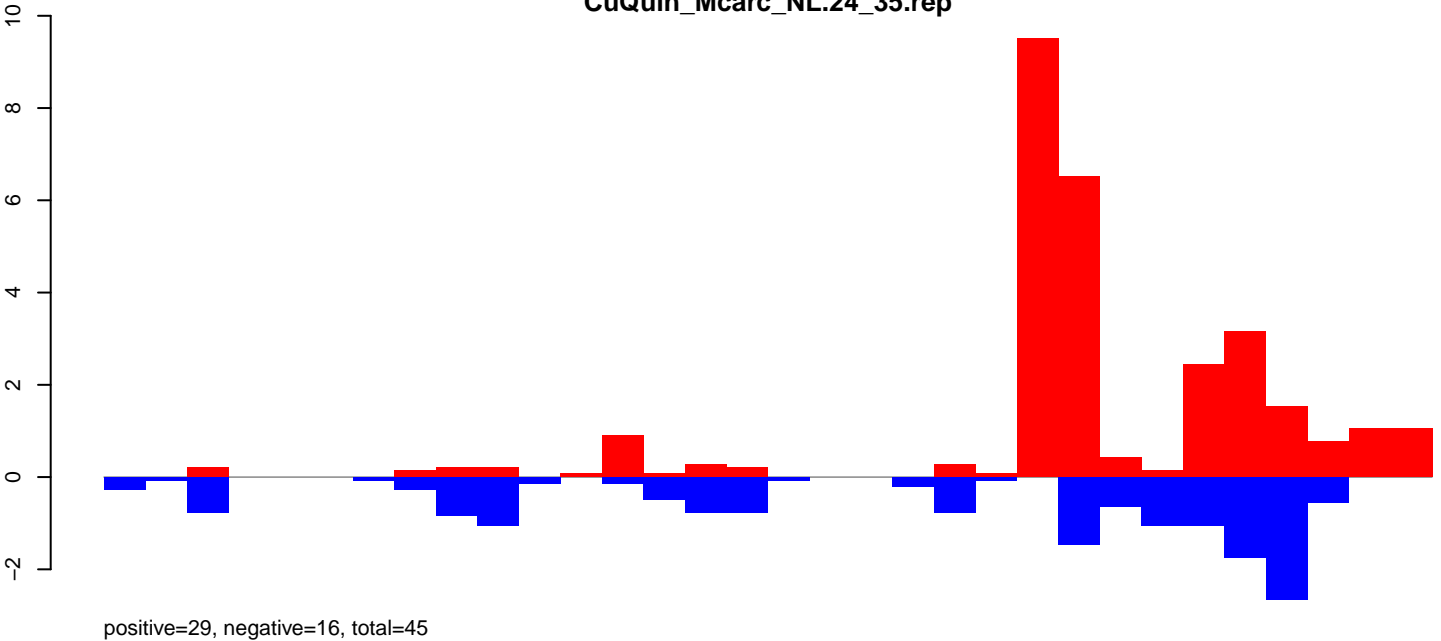
CuQuin_Mcarc_NL.rep



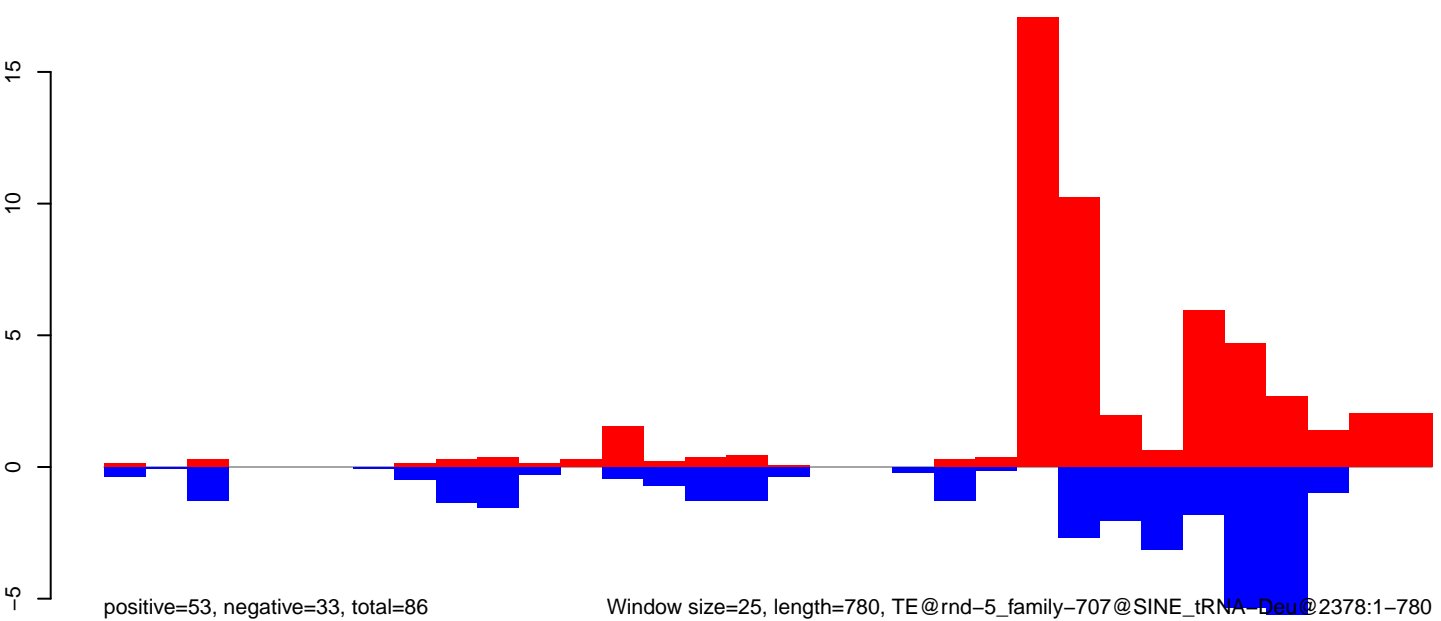
CuQuin_Mcarc_NL.18_23.rep



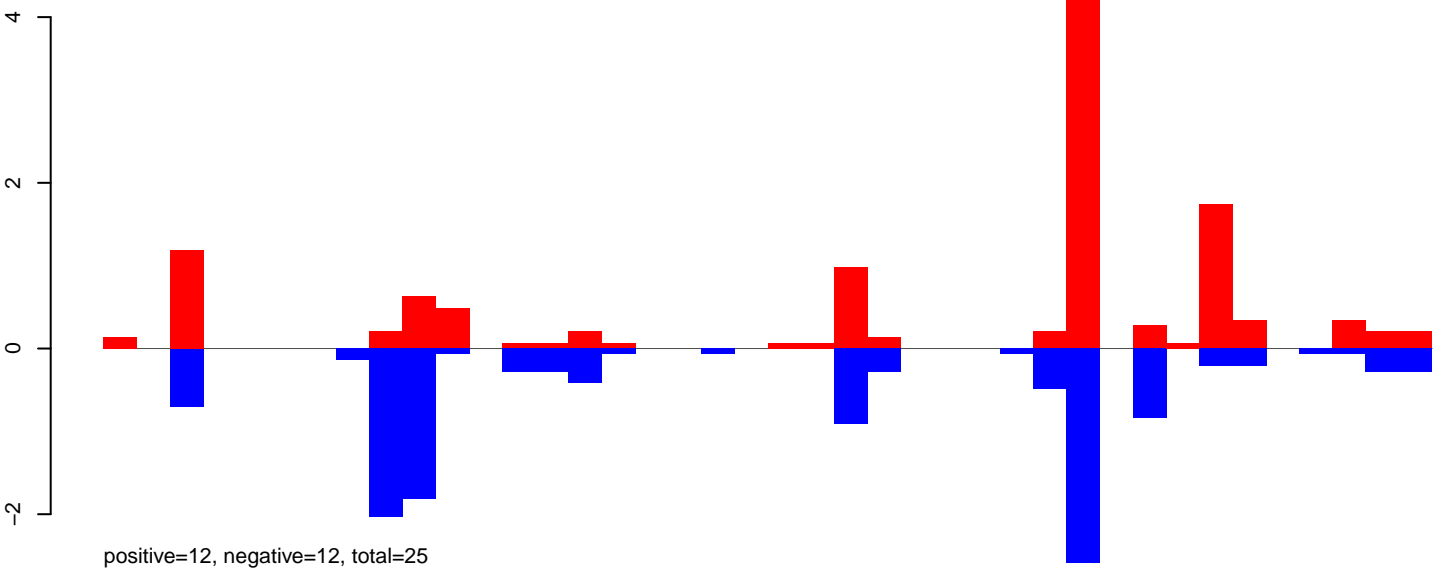
CuQuin_Mcarc_NL.24_35.rep



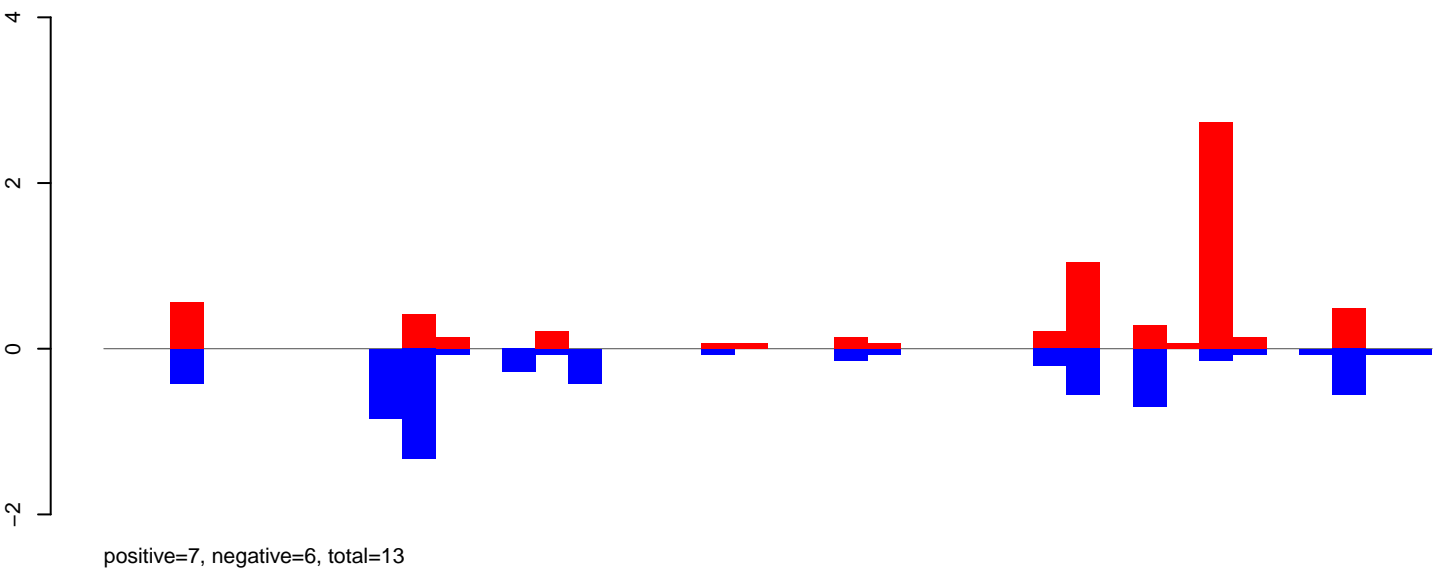
CuQuin_Mcarc_NL.rep



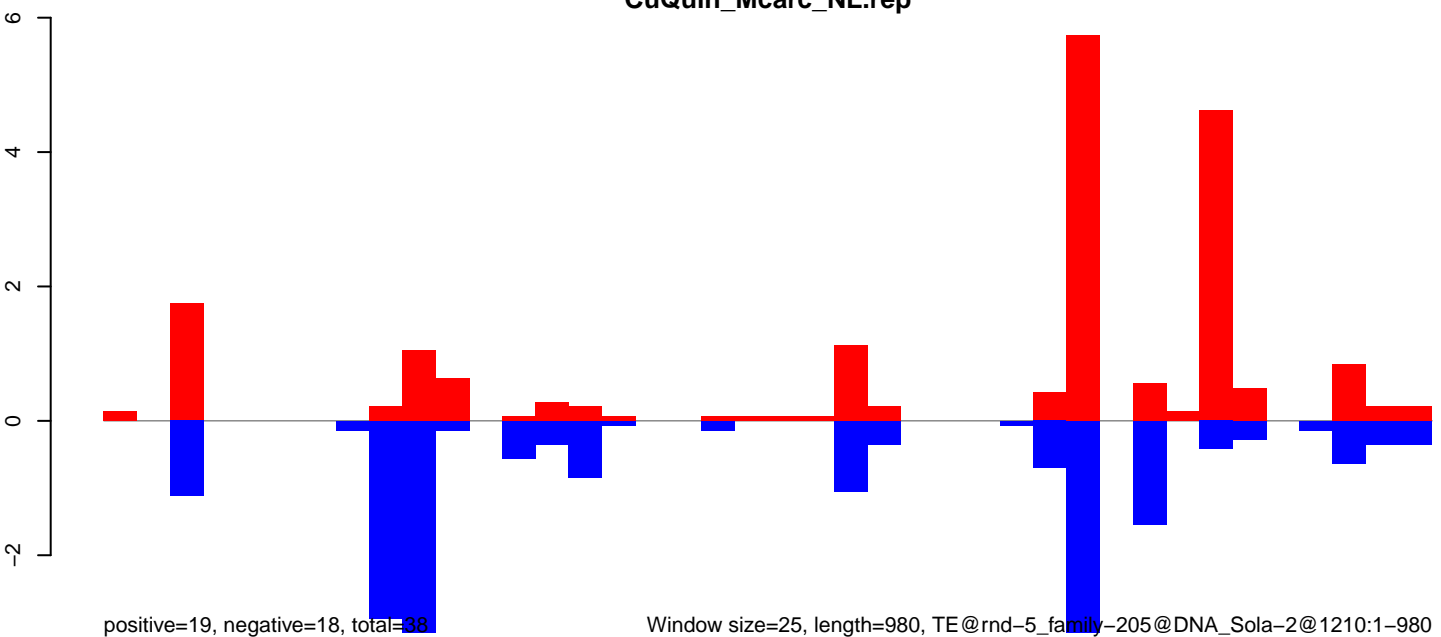
CuQuin_Mcarc_NL.18_23.rep



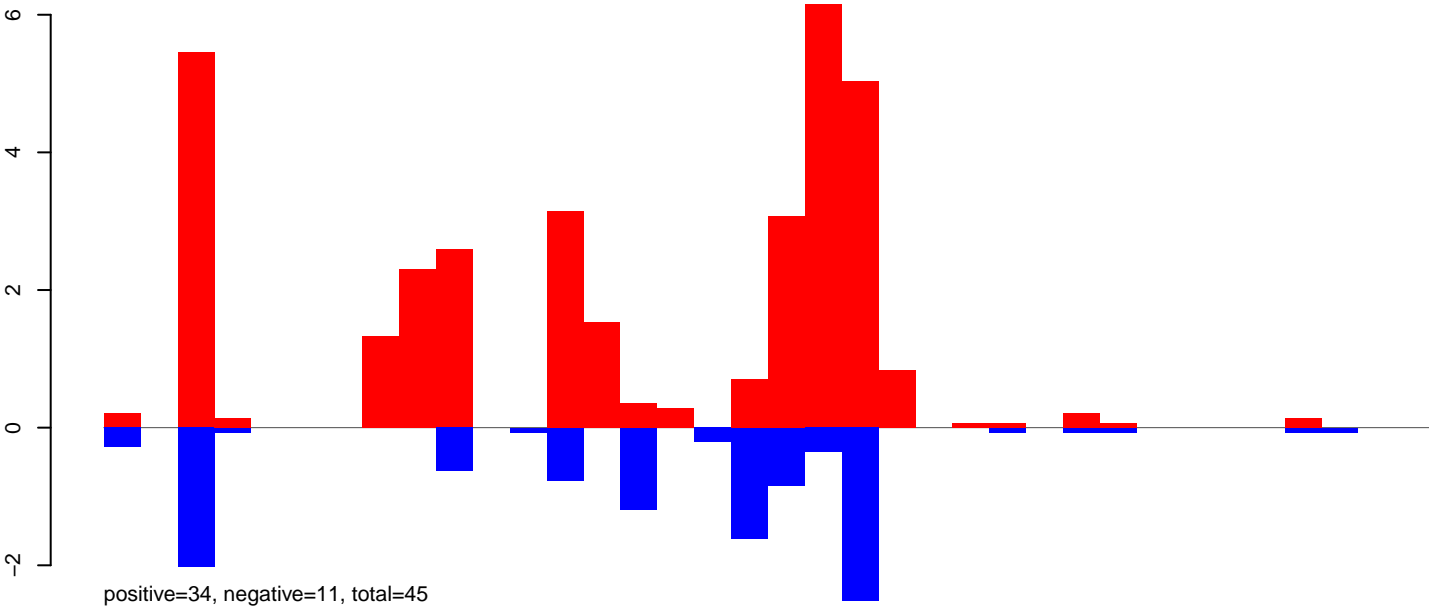
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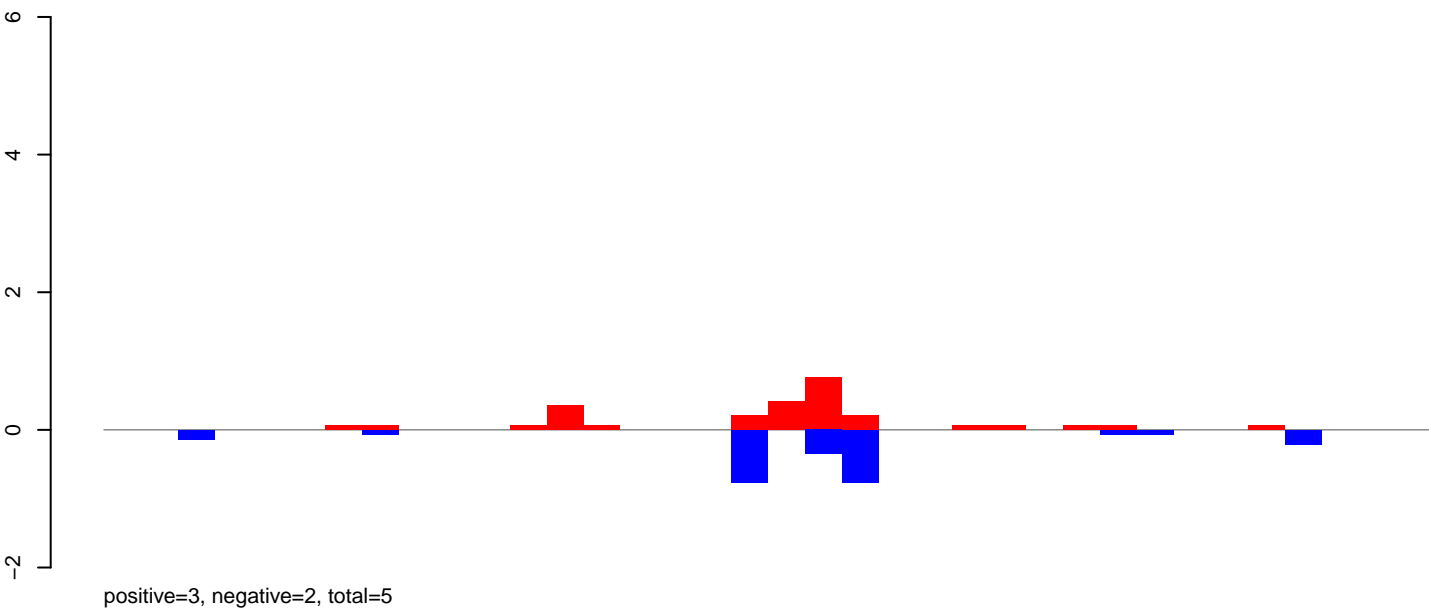
CuQuin_Mcarc_NL.rep



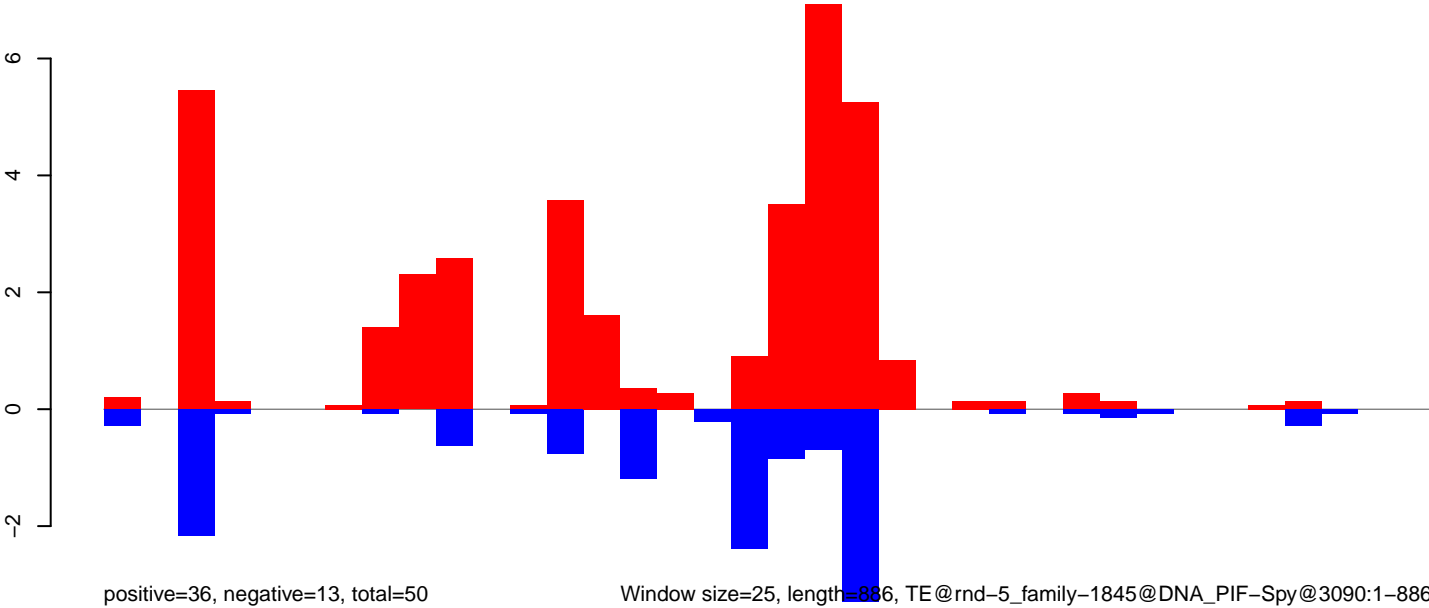
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



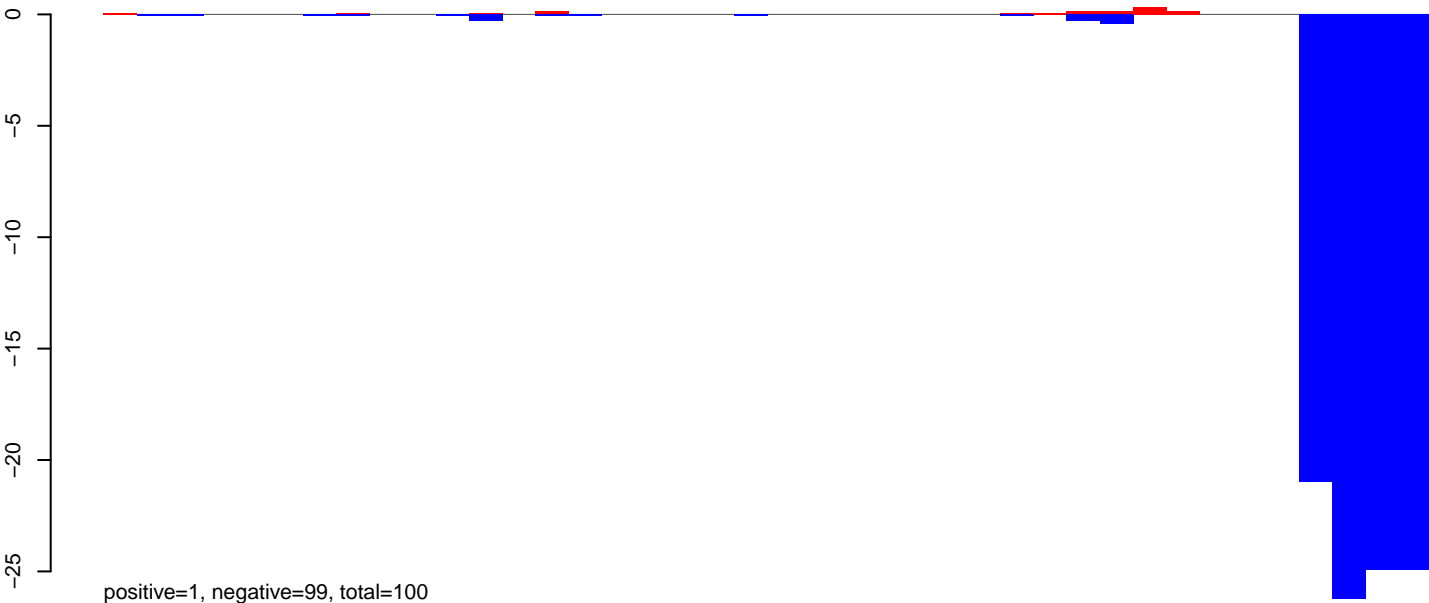
CuQuin_Mcarc_NL.rep



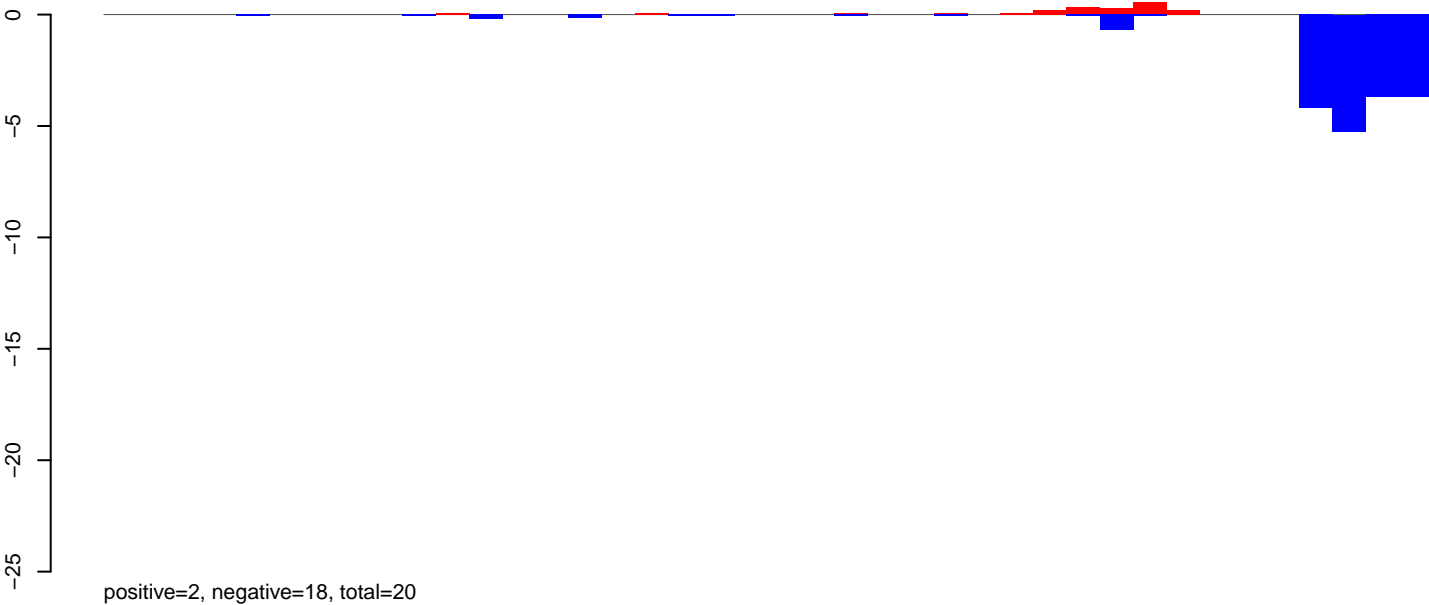
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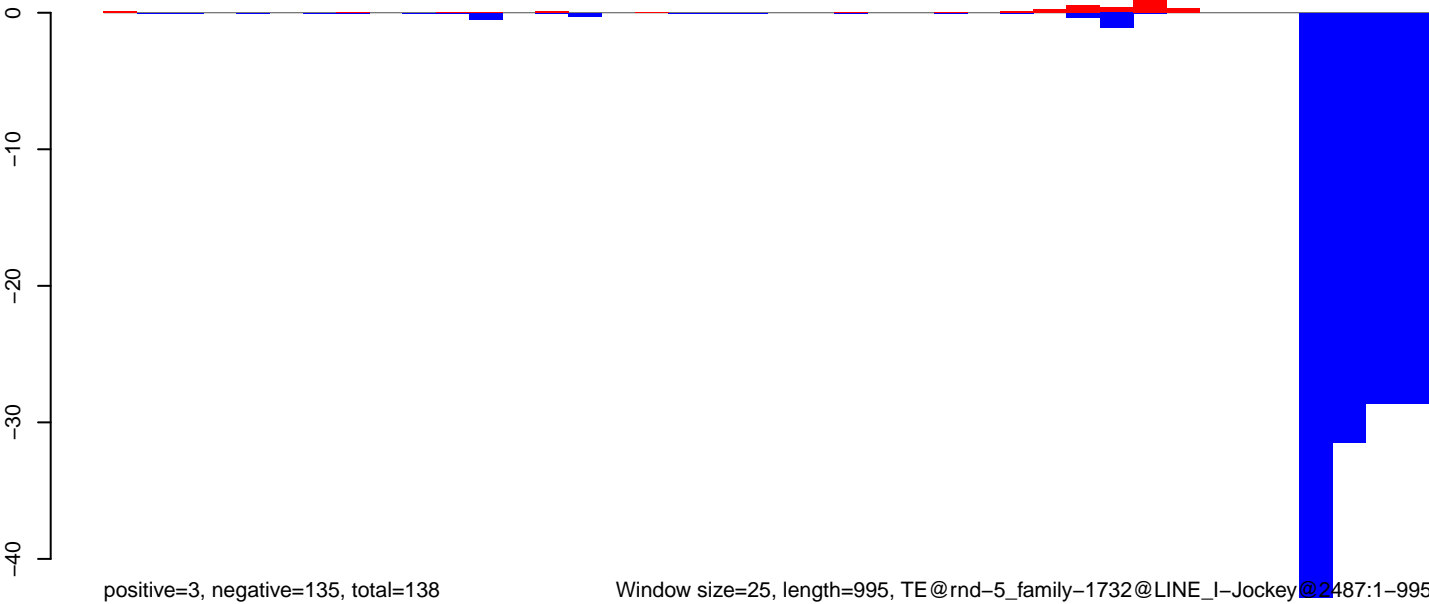
CuQuin_Mcarc_NL.18_23.rep



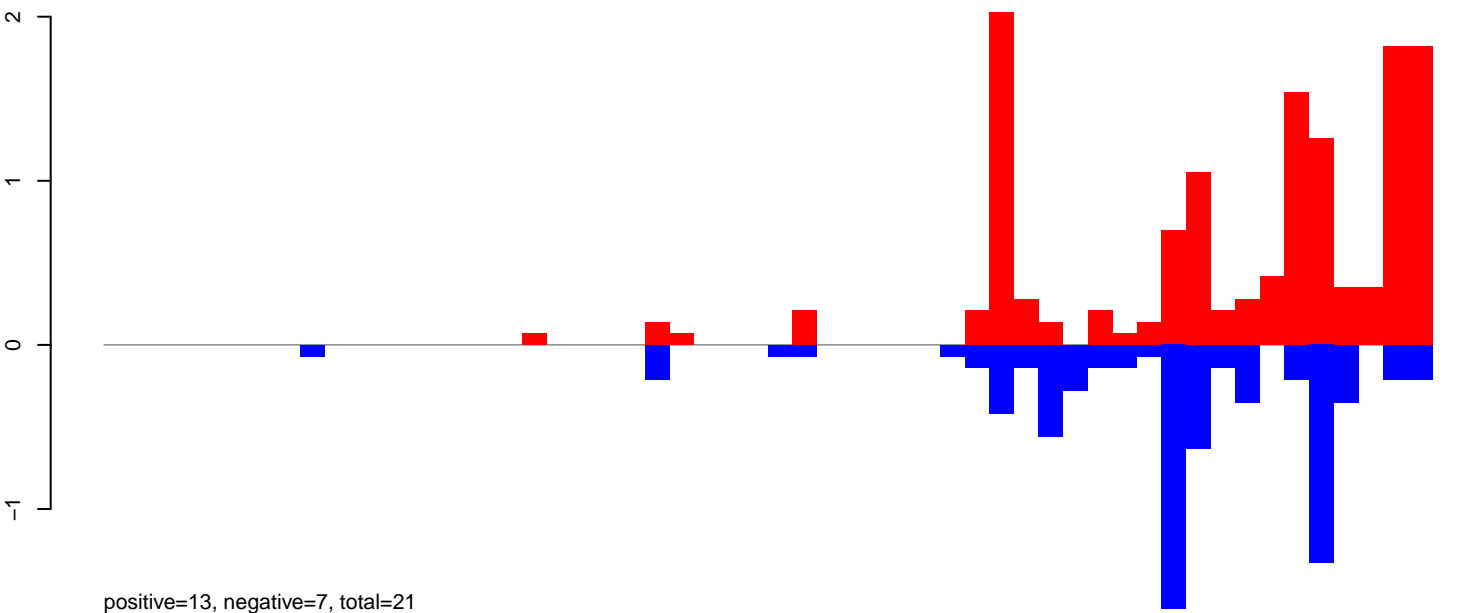
CuQuin_Mcarc_NL.24_35.rep



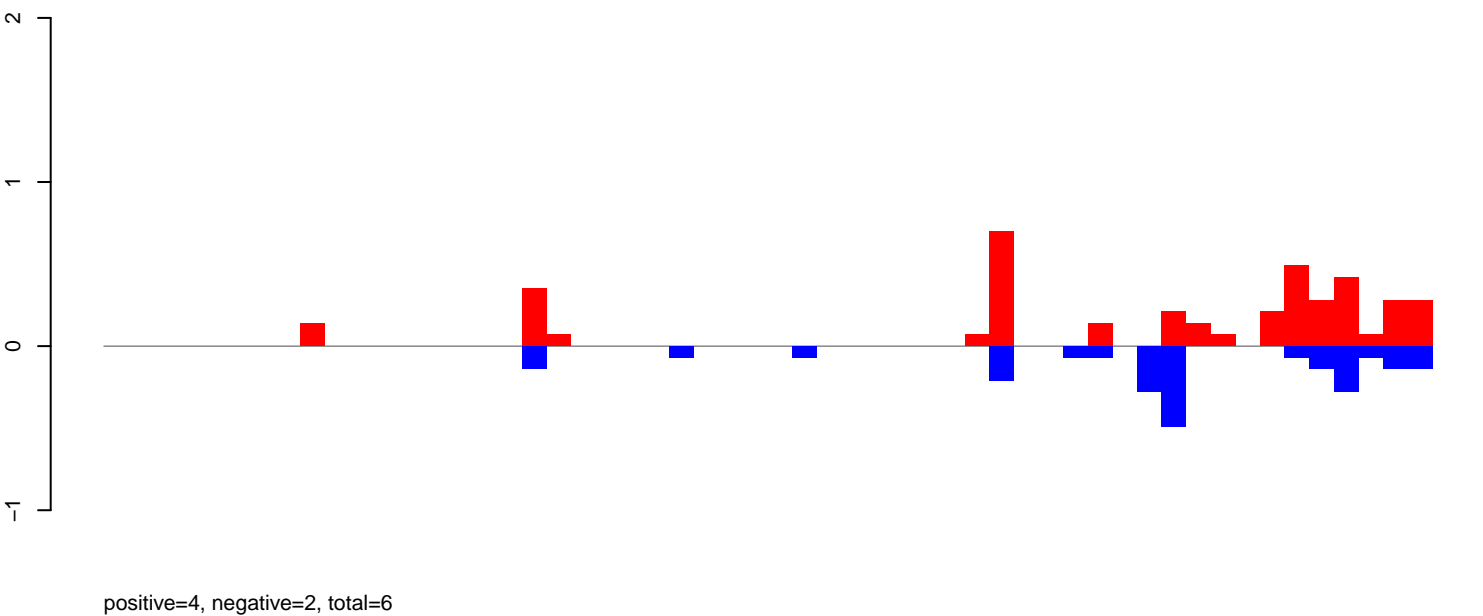
CuQuin_Mcarc_NL.rep



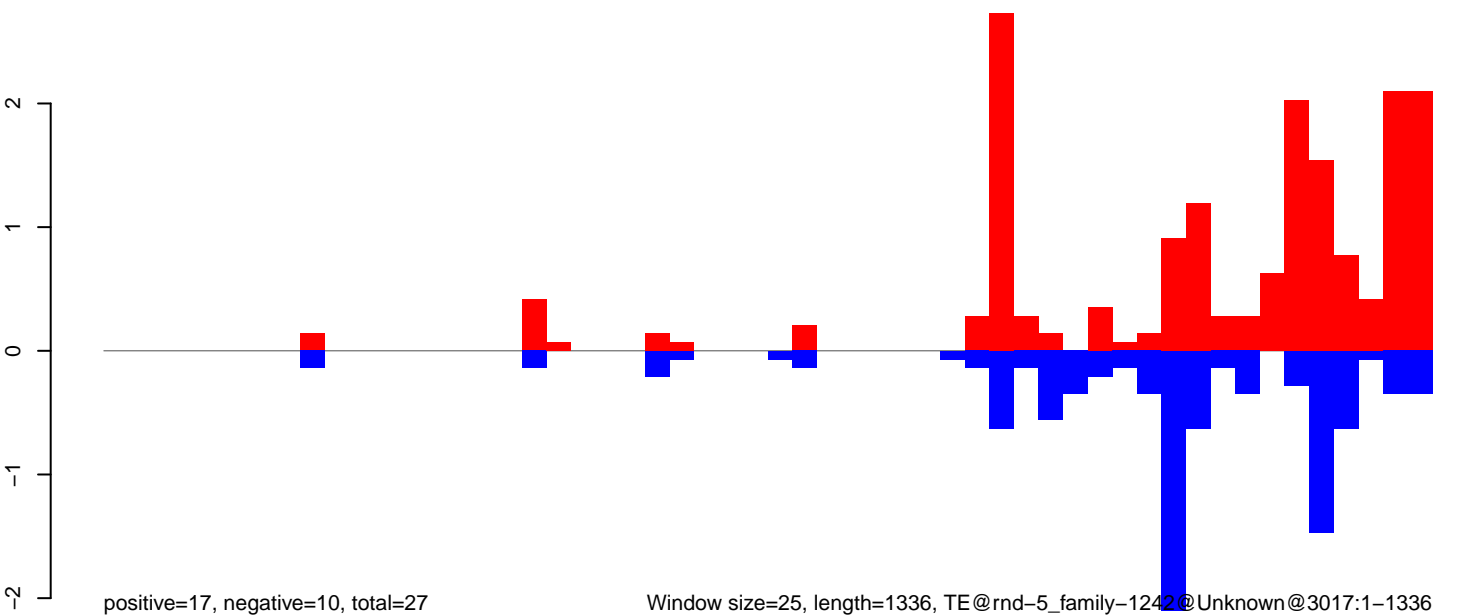
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

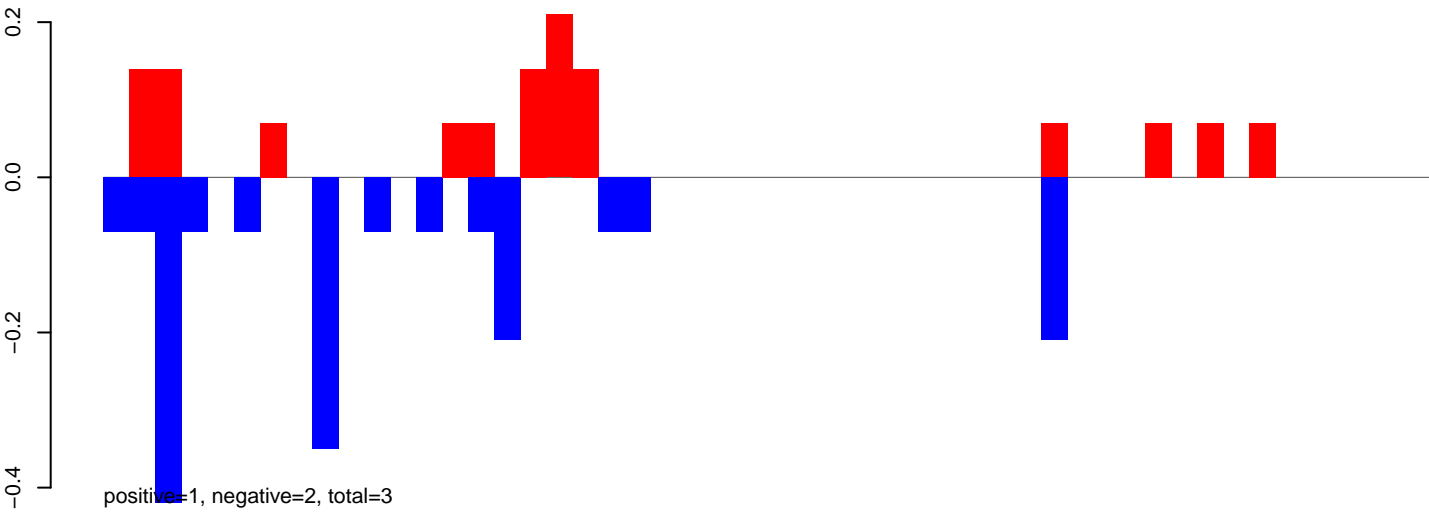


CuQuin_Mcarc_NL.rep

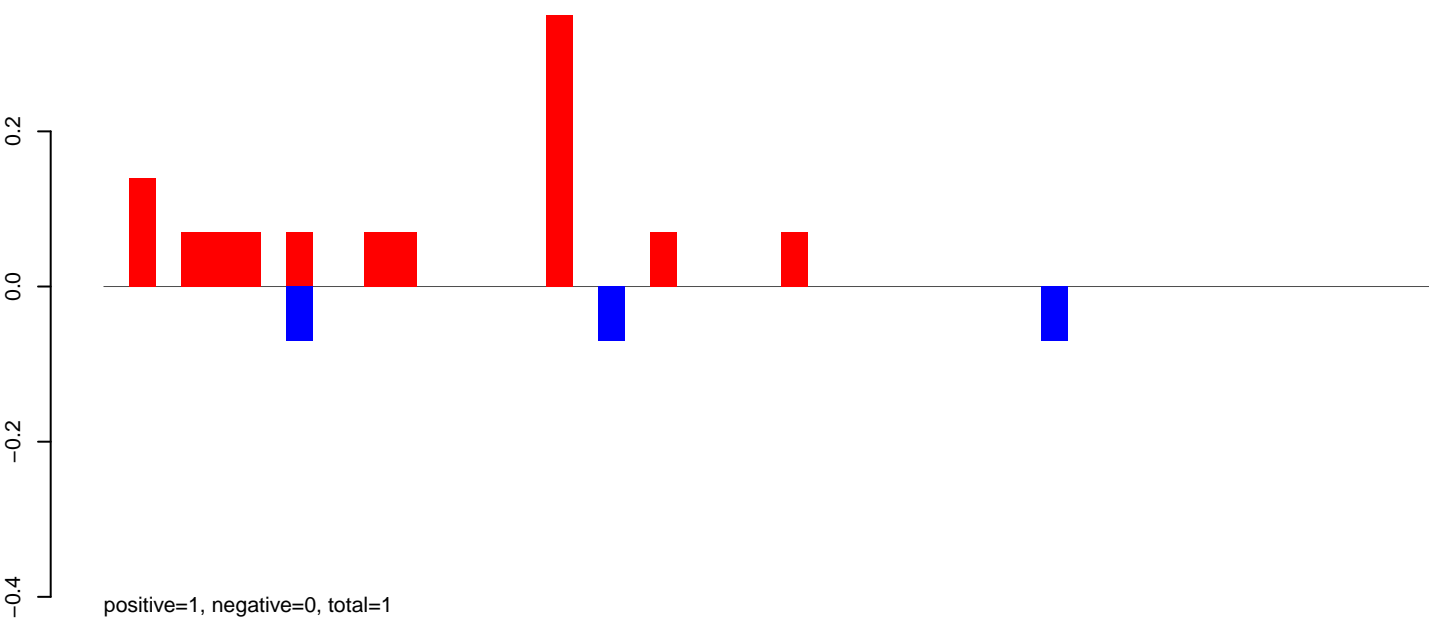


Window size=25, length=1336, TE@rnd-5_family-1242@Unknown@3017:1-1336

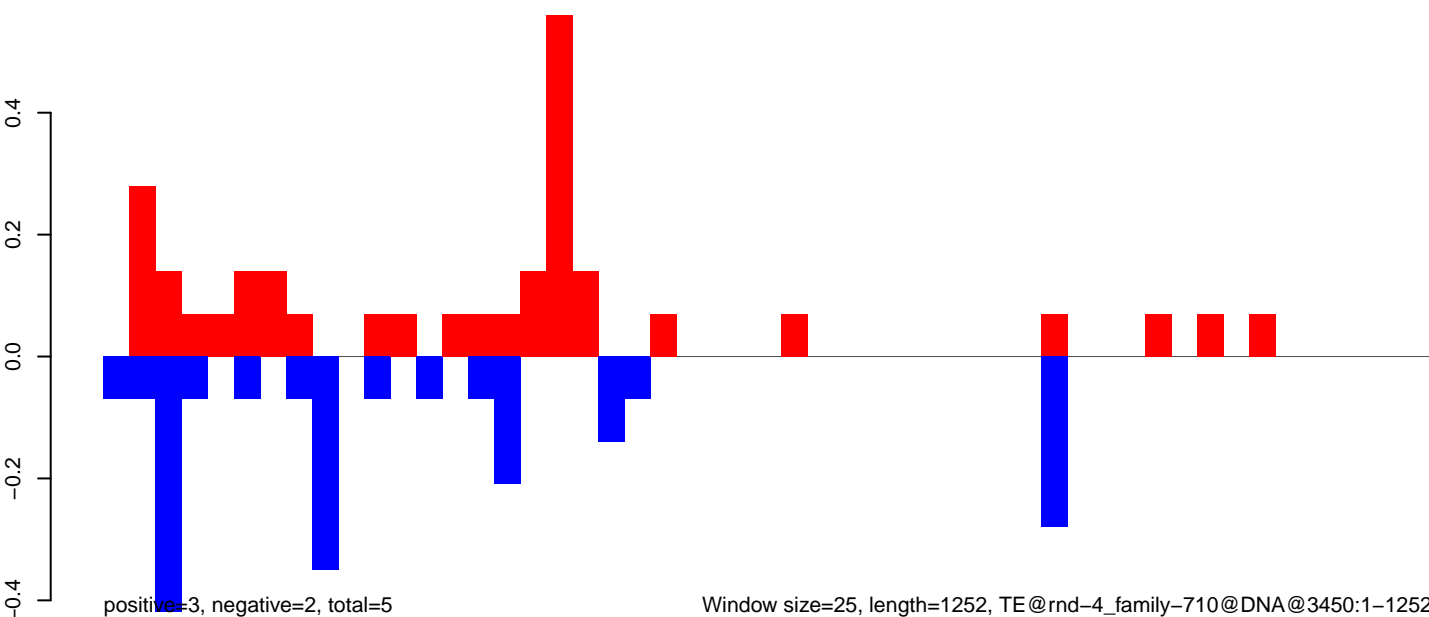
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



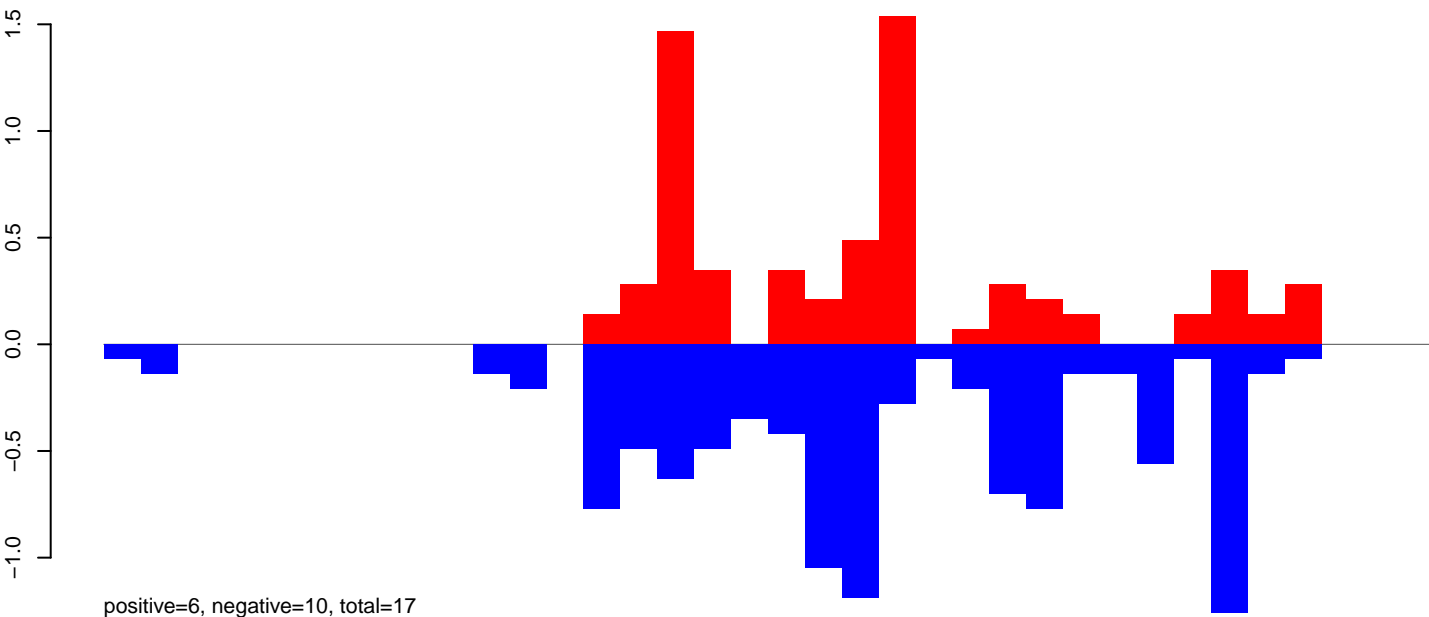
CuQuin_Mcarc_NL.rep



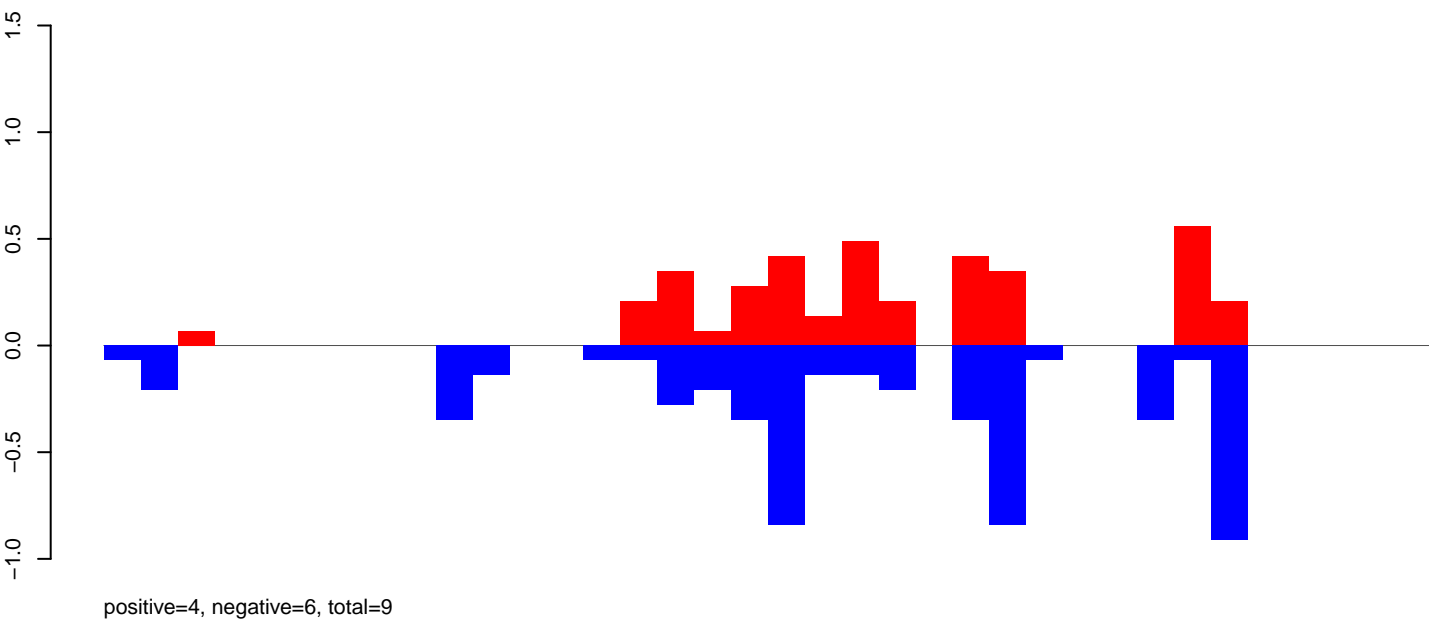
Window size=25, length=1252, TE@rnd-4_family-710@DNA@3450:1-1252

0 200 400 600 800 1000 1200

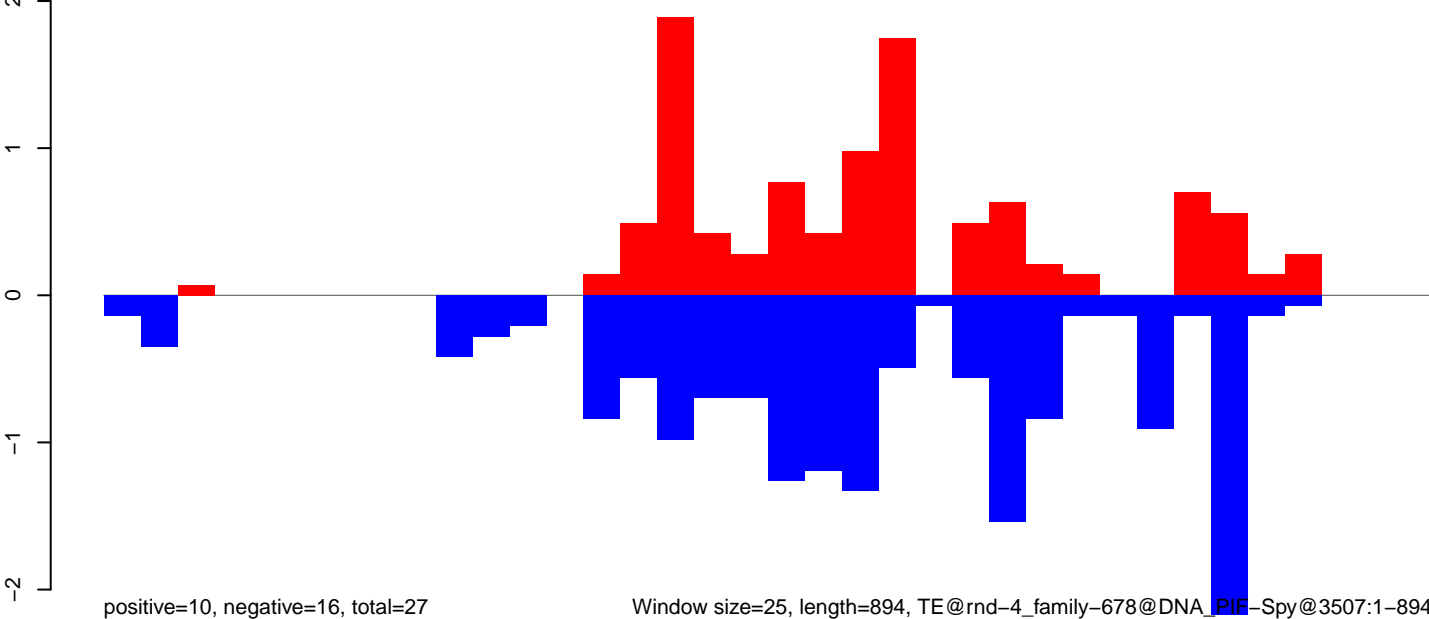
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

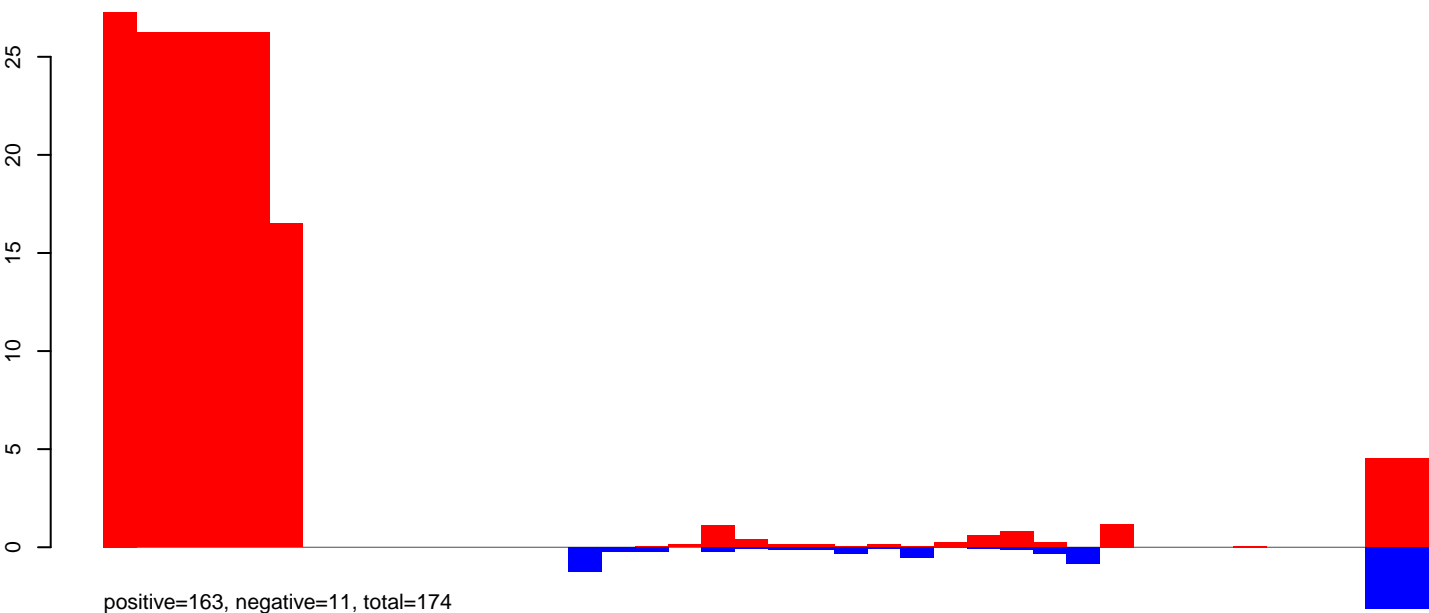


CuQuin_Mcarc_NL.rep

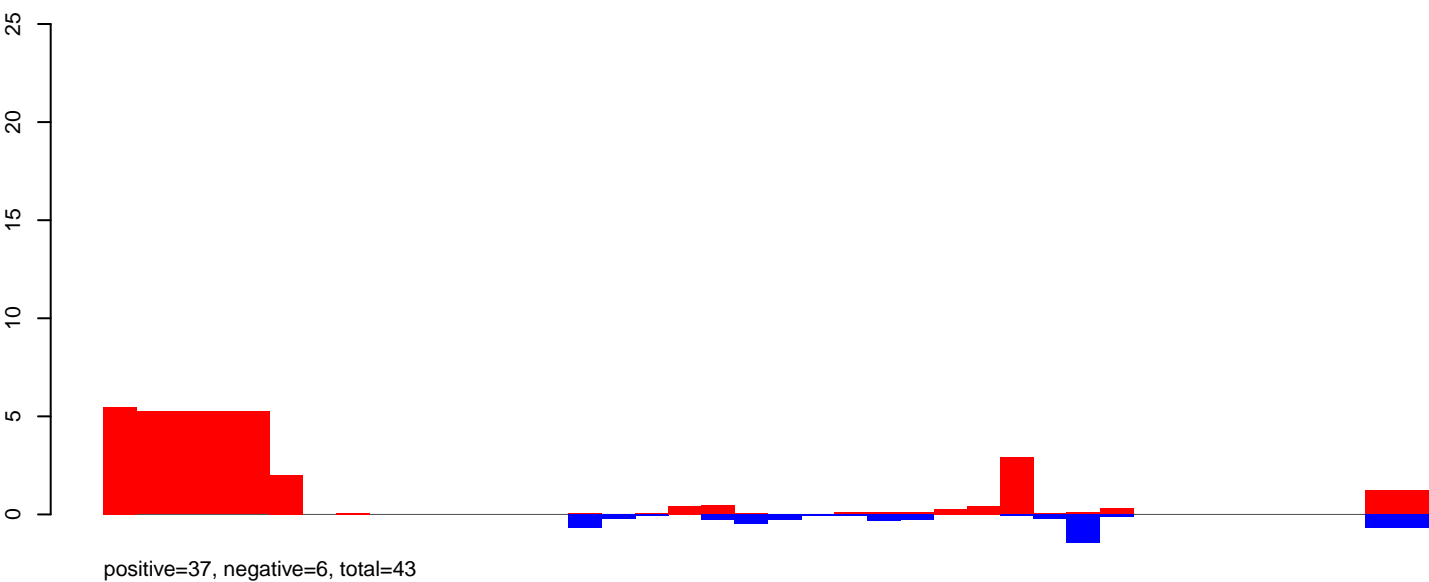


Window size=25, length=894, TE@rnd-4_family-678@DNA_PIF-Spy@3507:1-894

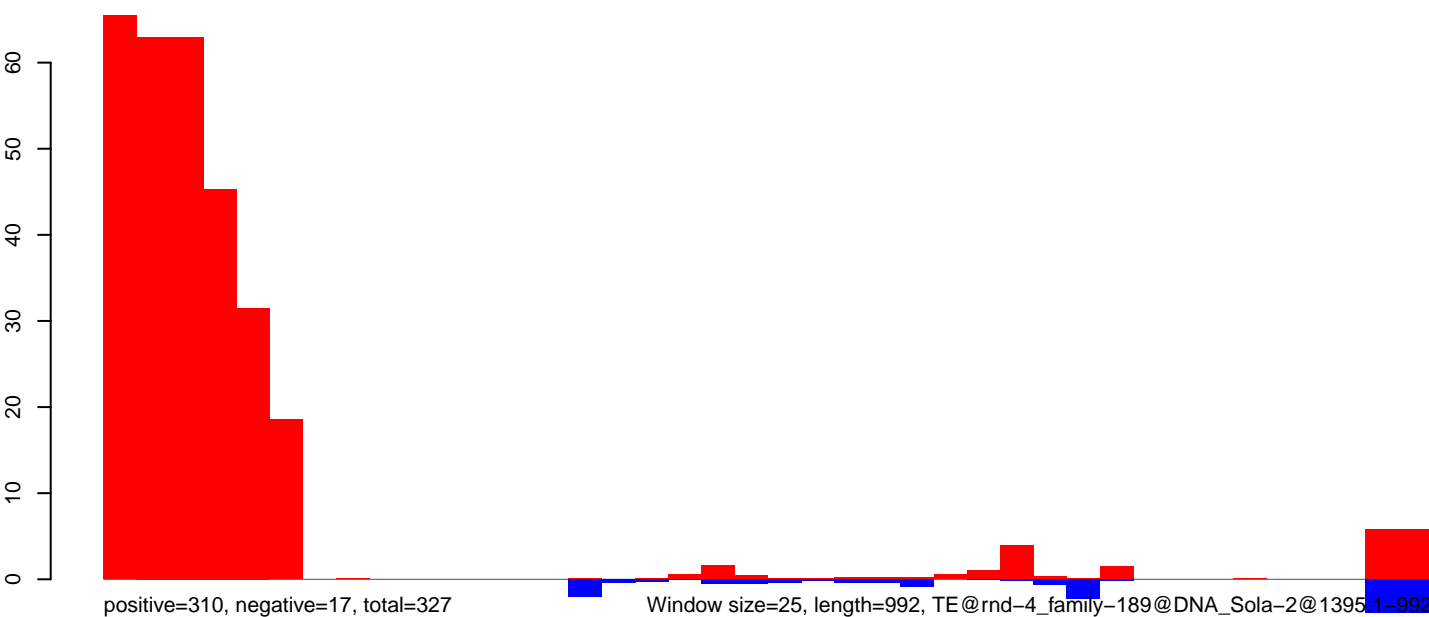
CuQuin_Mcarc_NL.18_23.rep



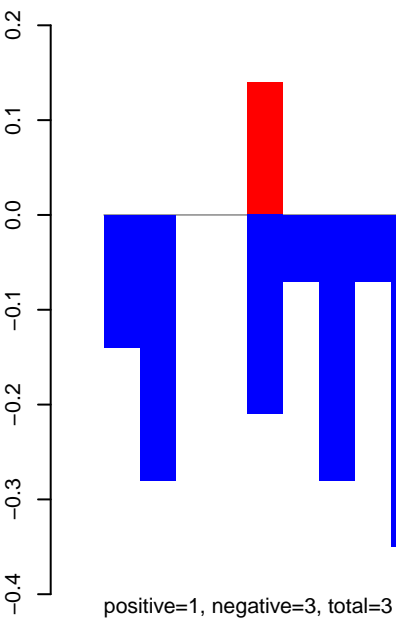
CuQuin_Mcarc_NL.24_35.rep



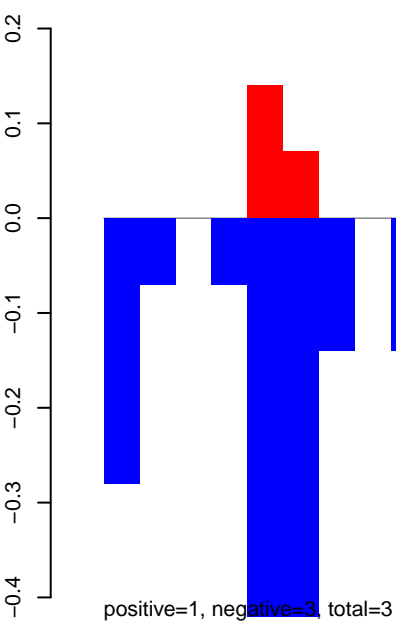
CuQuin_Mcarc_NL.rep



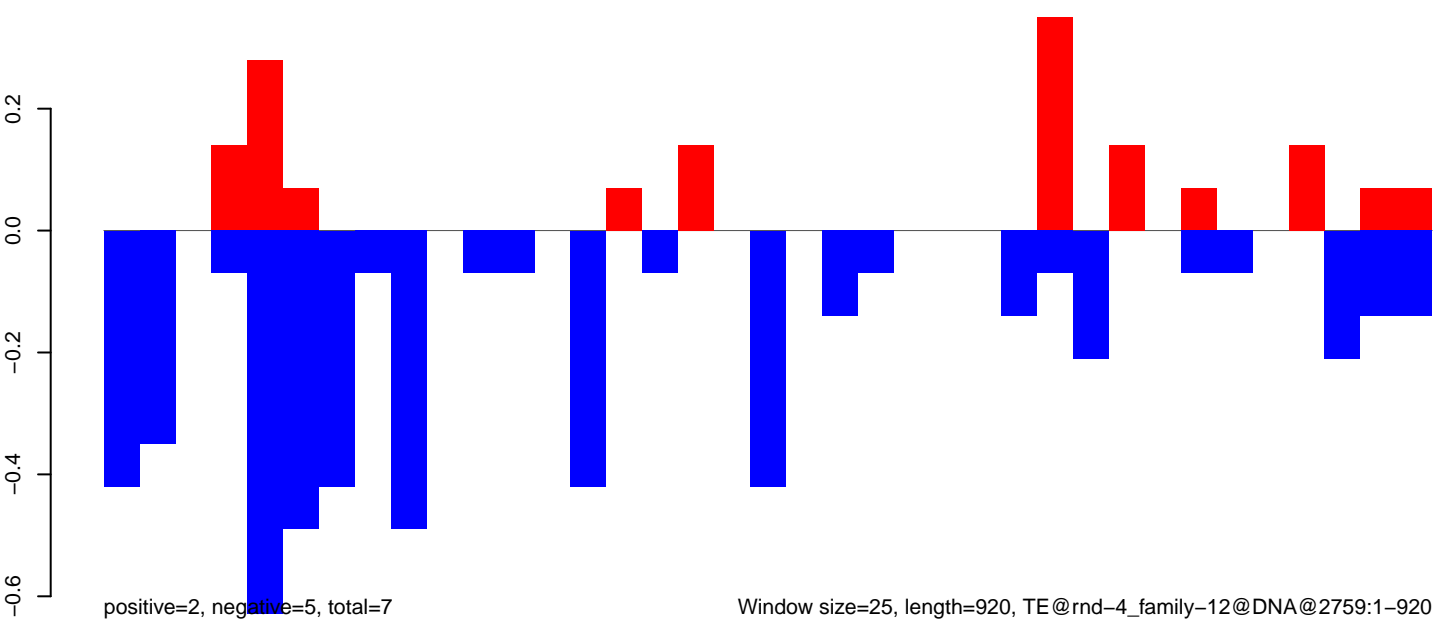
CuQuin_Mcarc_NL.18_23.rep



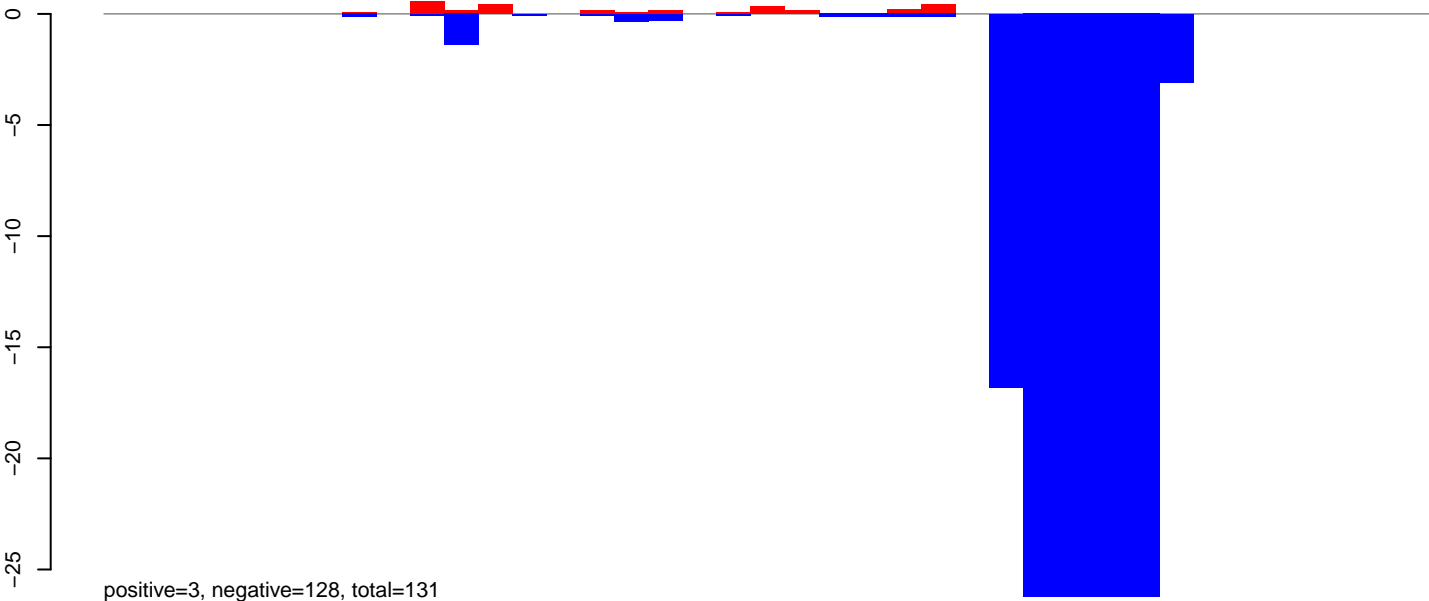
CuQuin_Mcarc_NL.24_35.rep



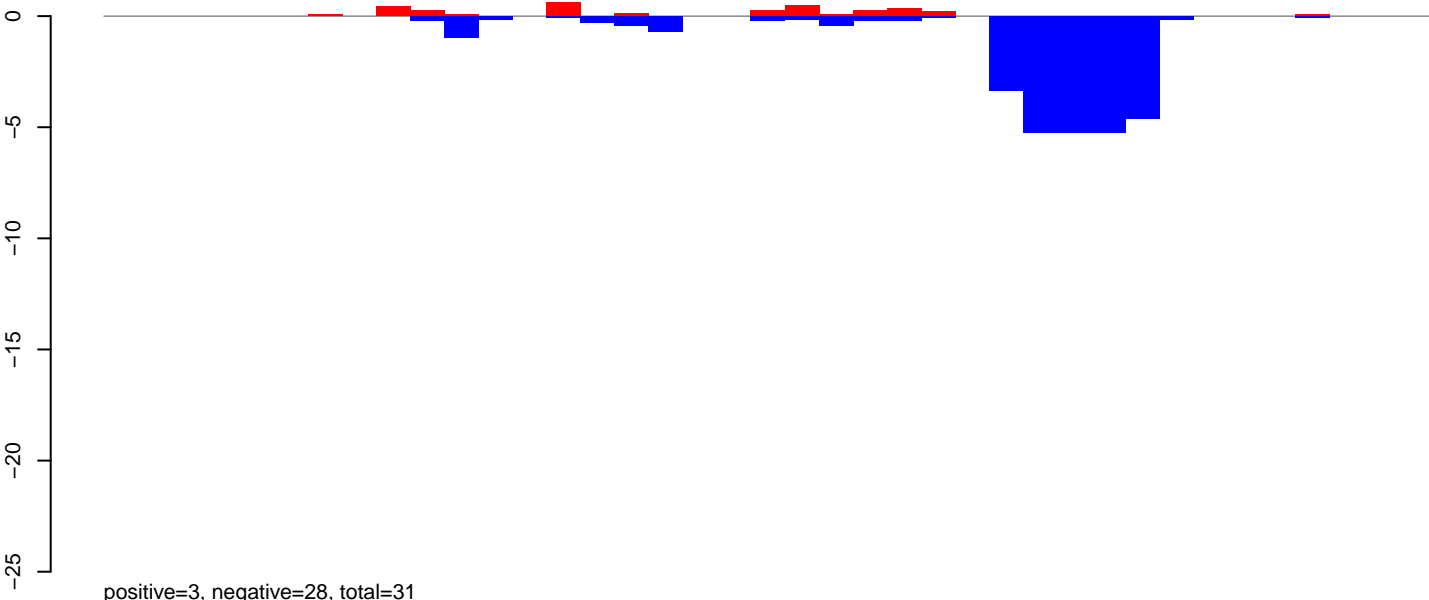
CuQuin_Mcarc_NL.rep



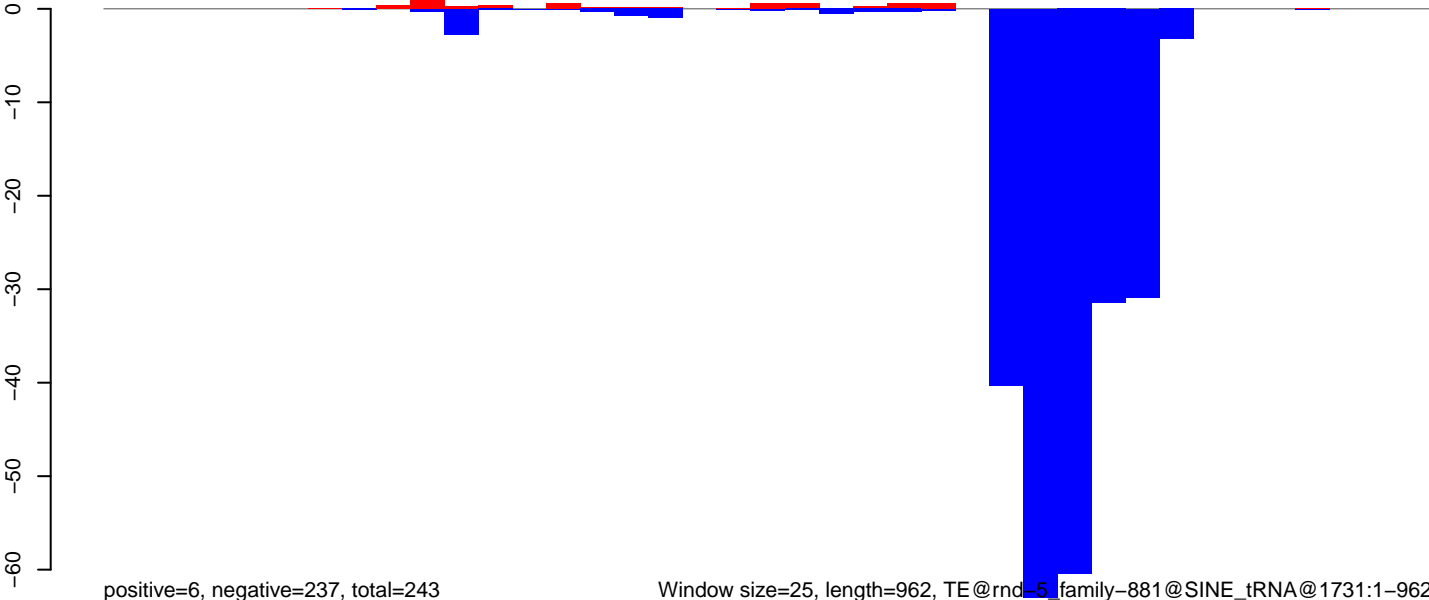
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



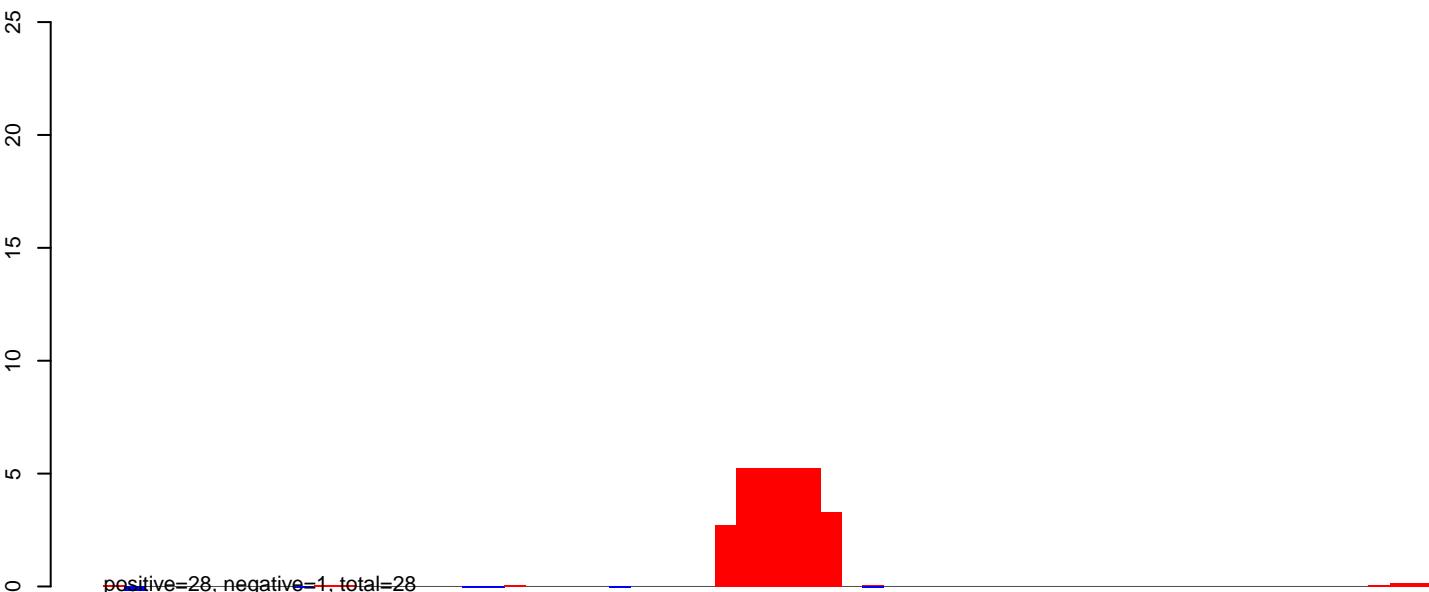
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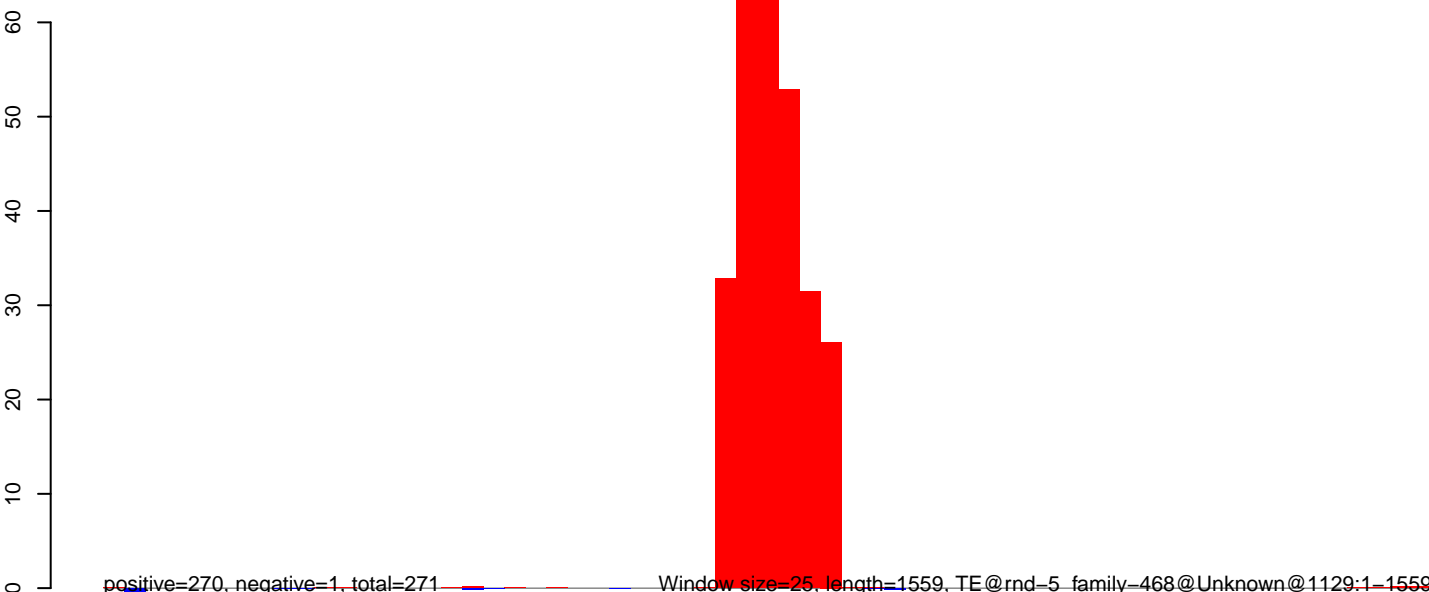
CuQuin_Mcarc_NL.18_23.rep

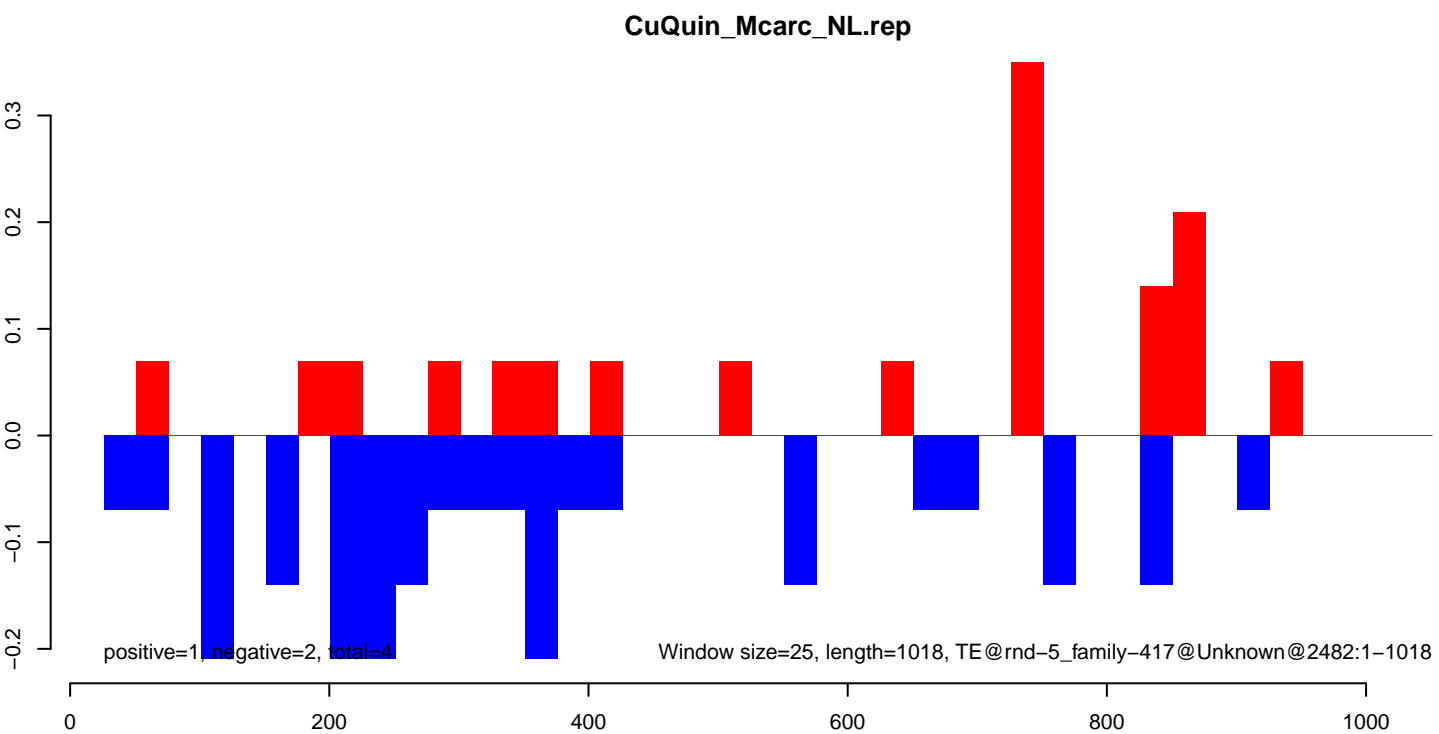
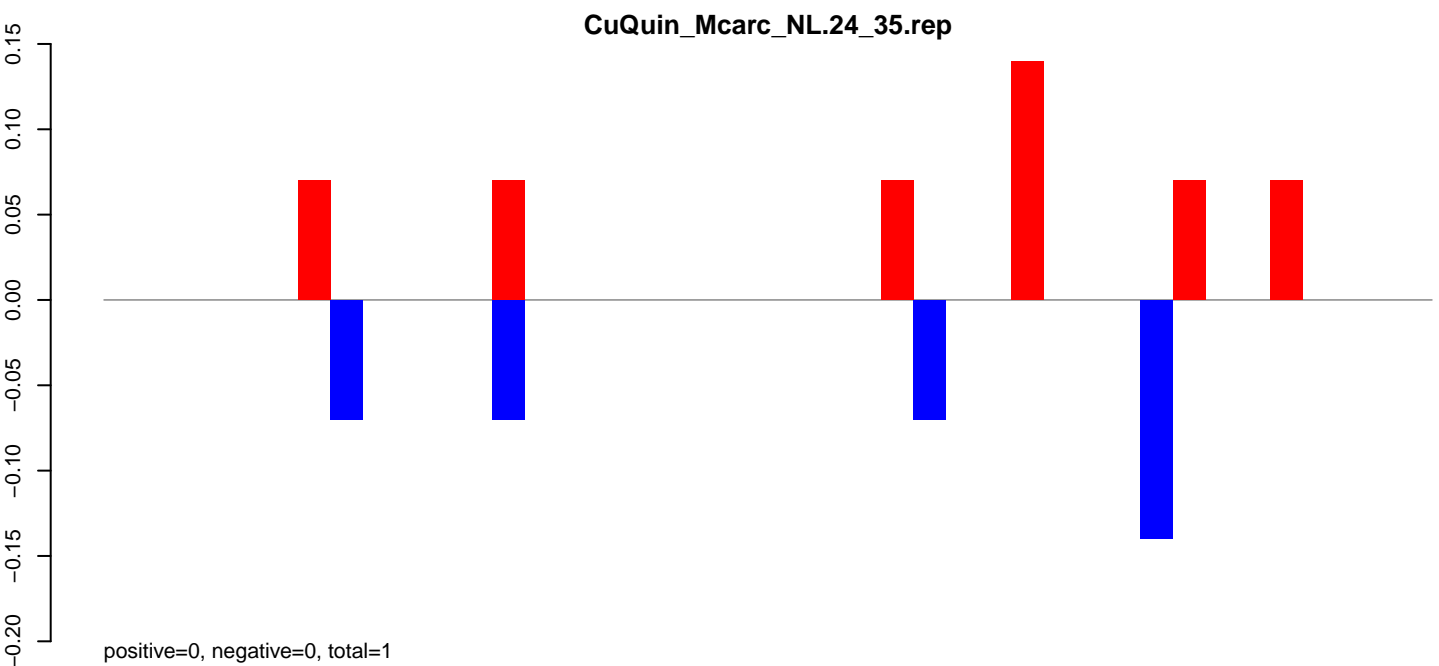
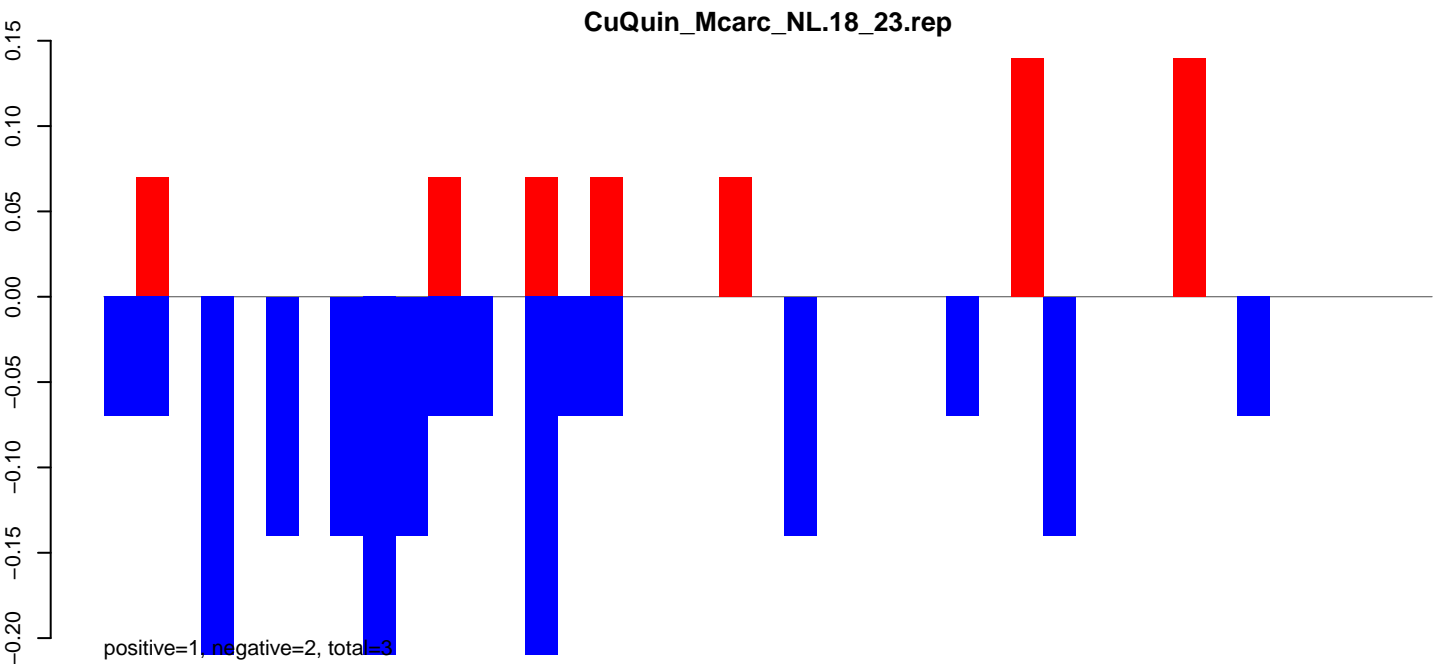


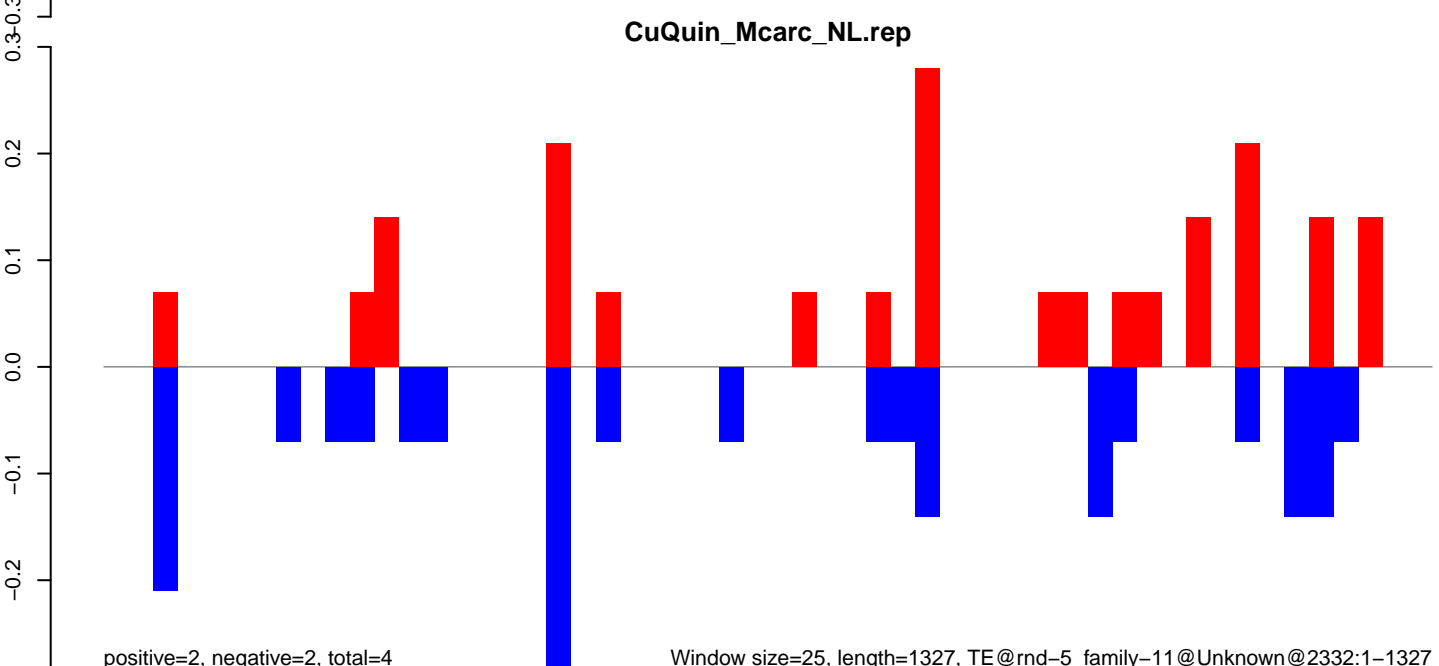
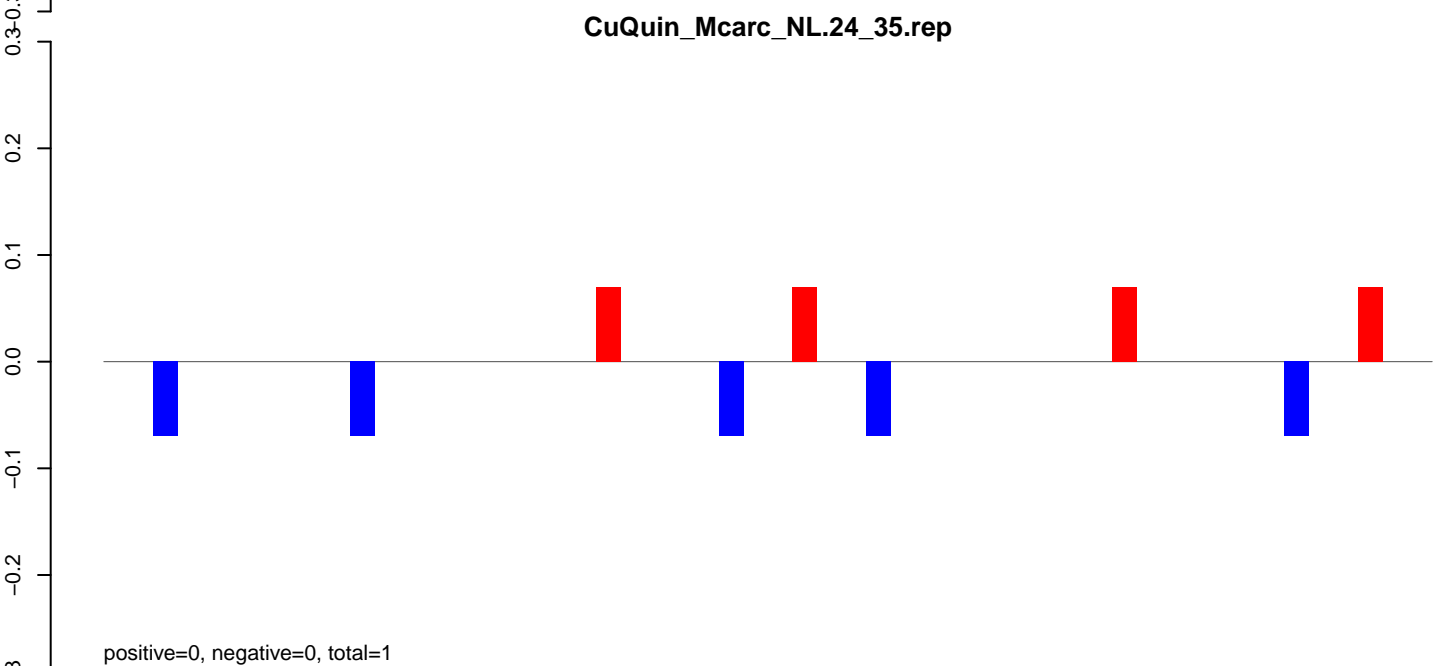
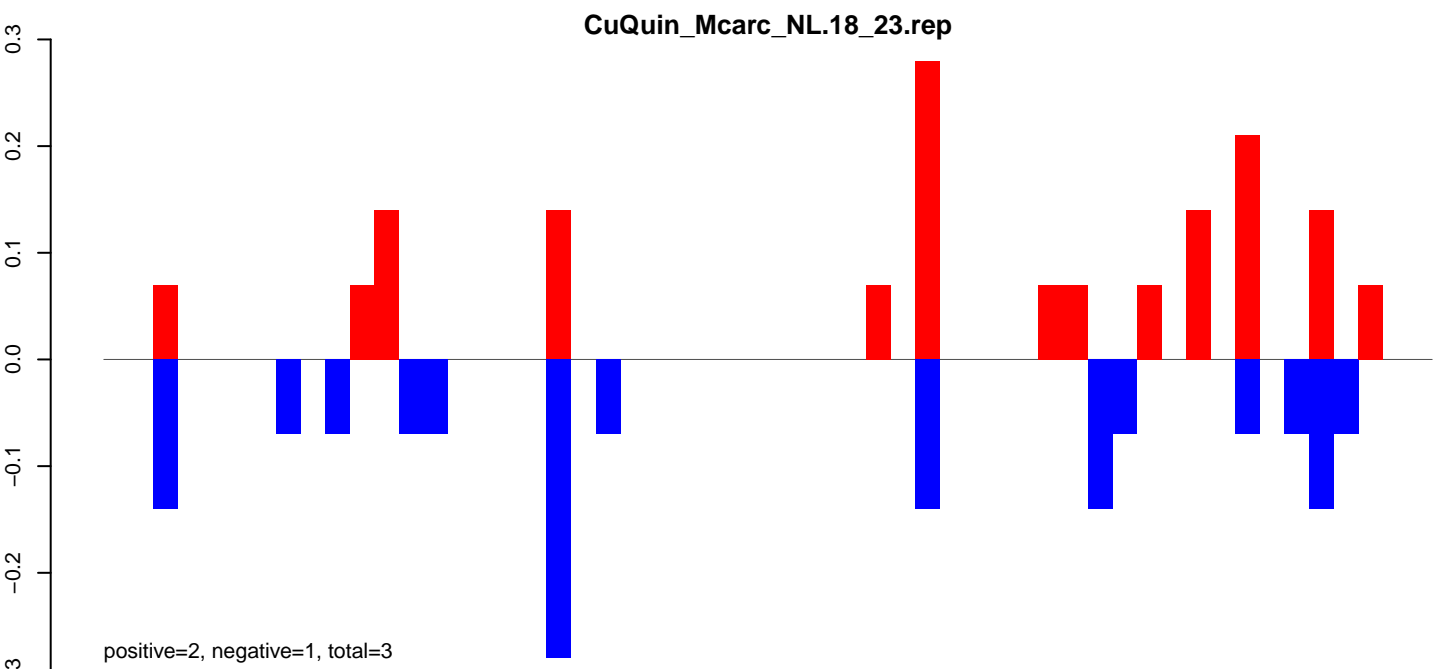
CuQuin_Mcarc_NL.24_35.rep



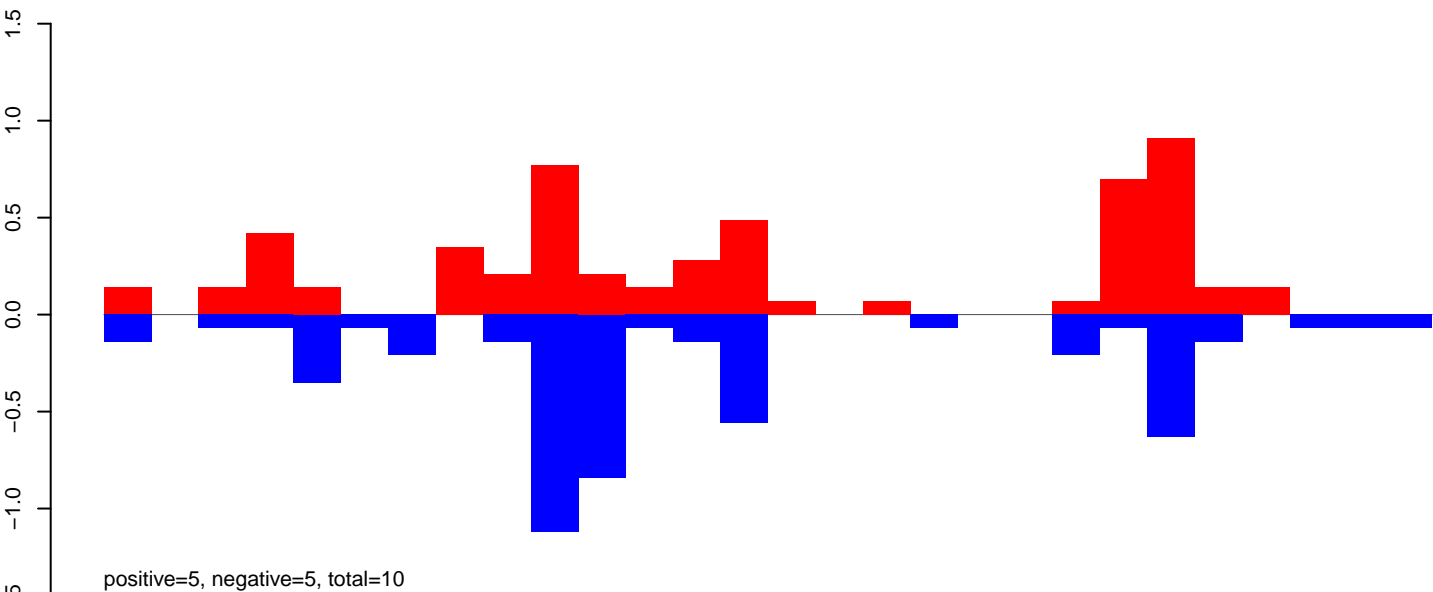
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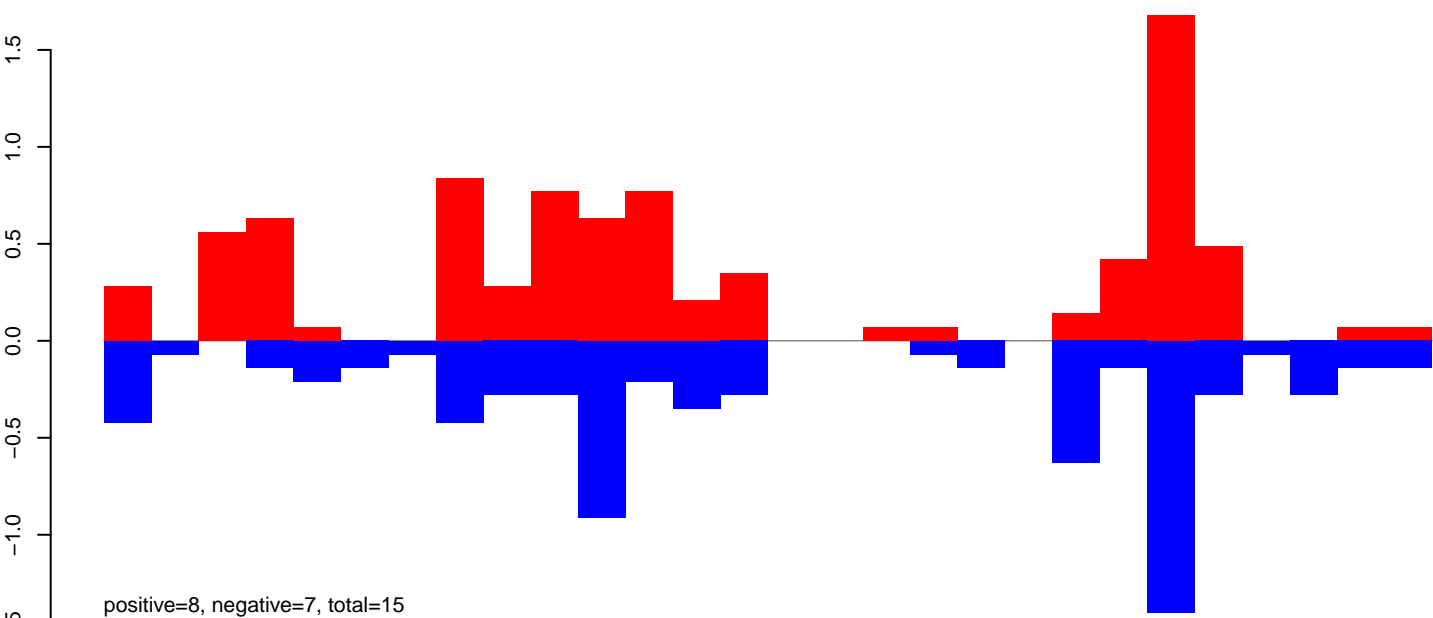




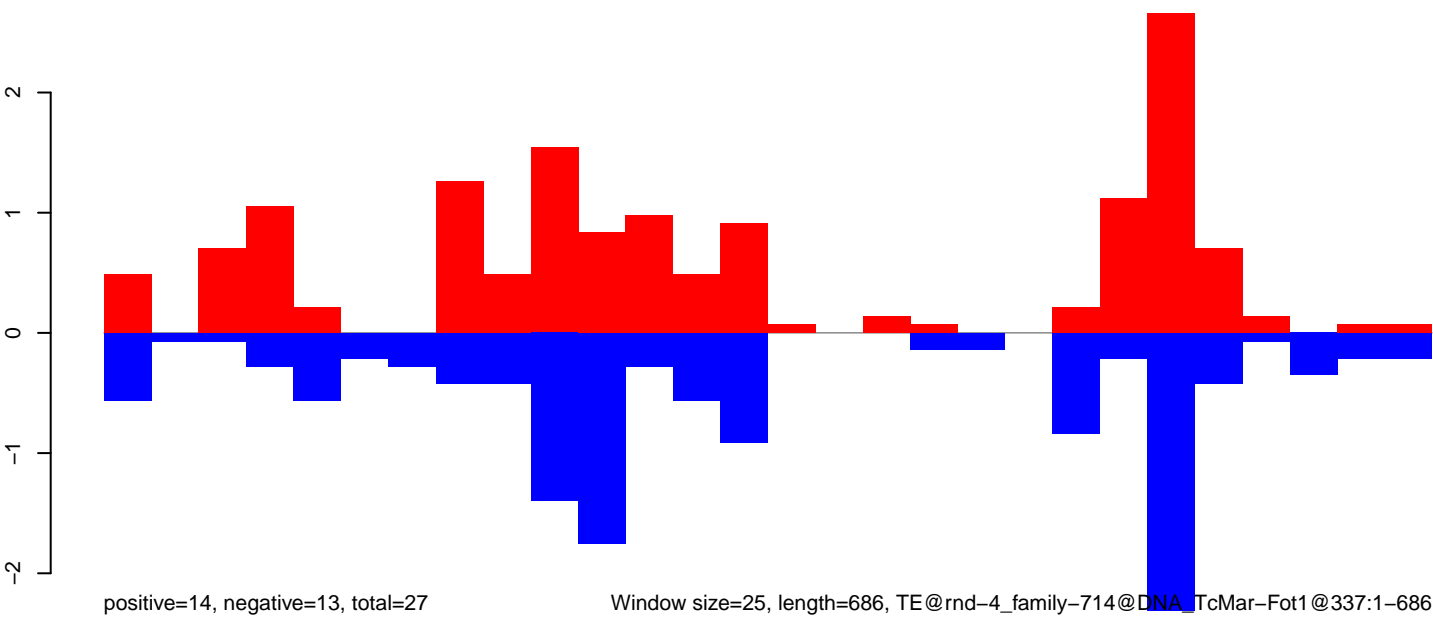
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

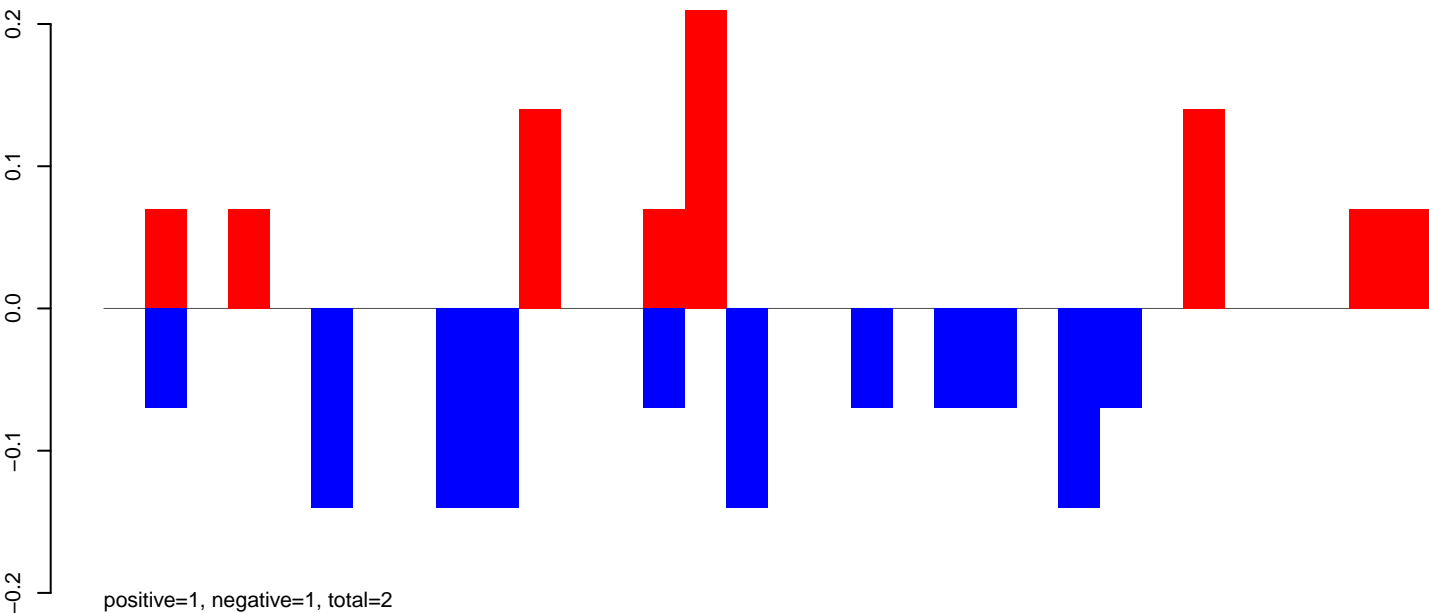


CuQuin_Mcarc_NL.rep

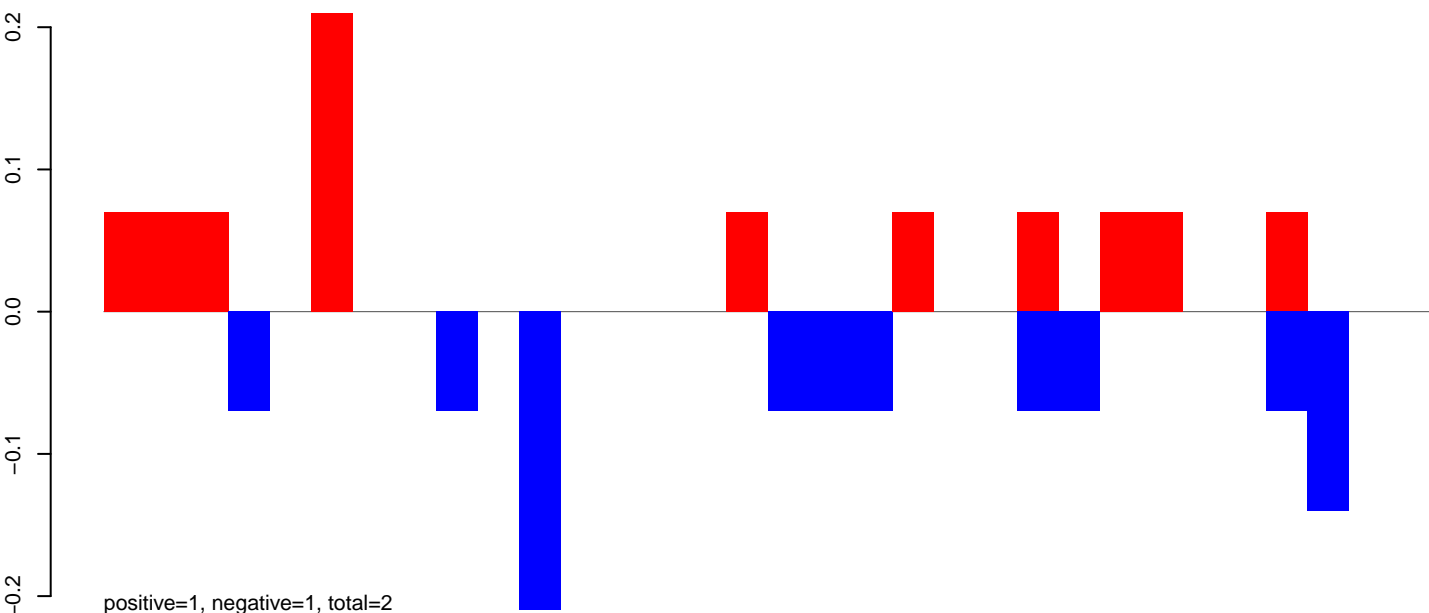


Window size=25, length=686, TE@rnd-4_family-714@DNA, TcMar-Fot1@337:1-686

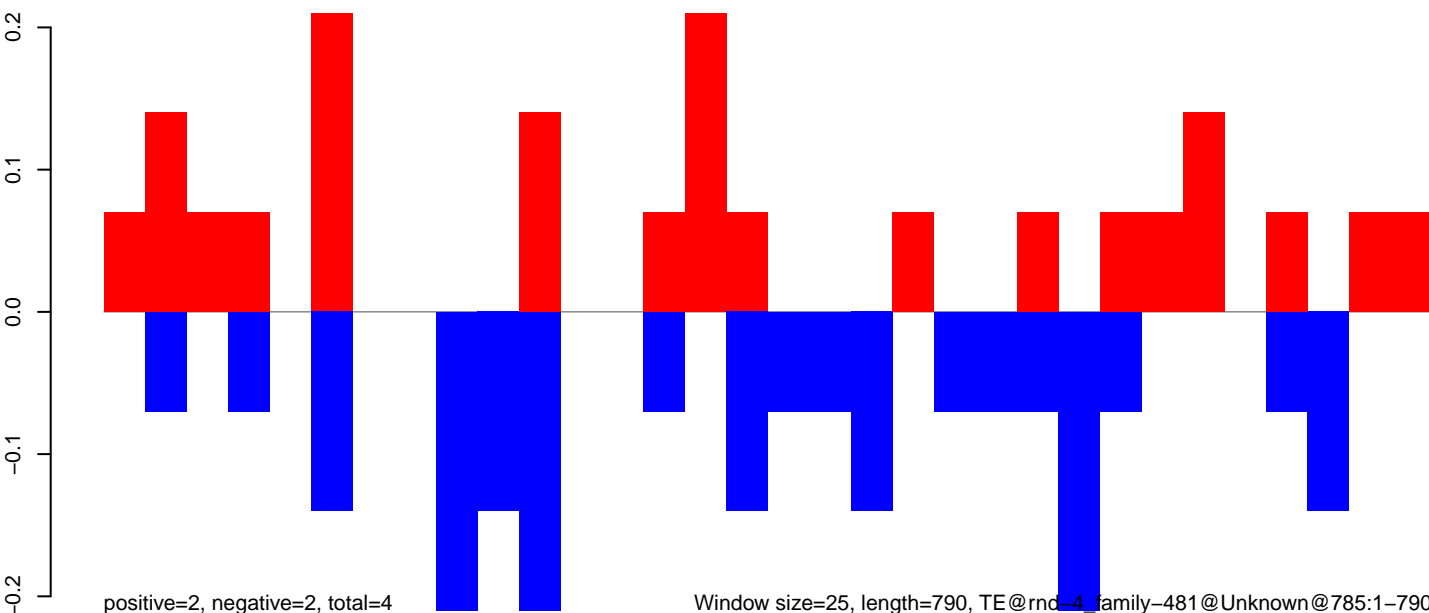
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

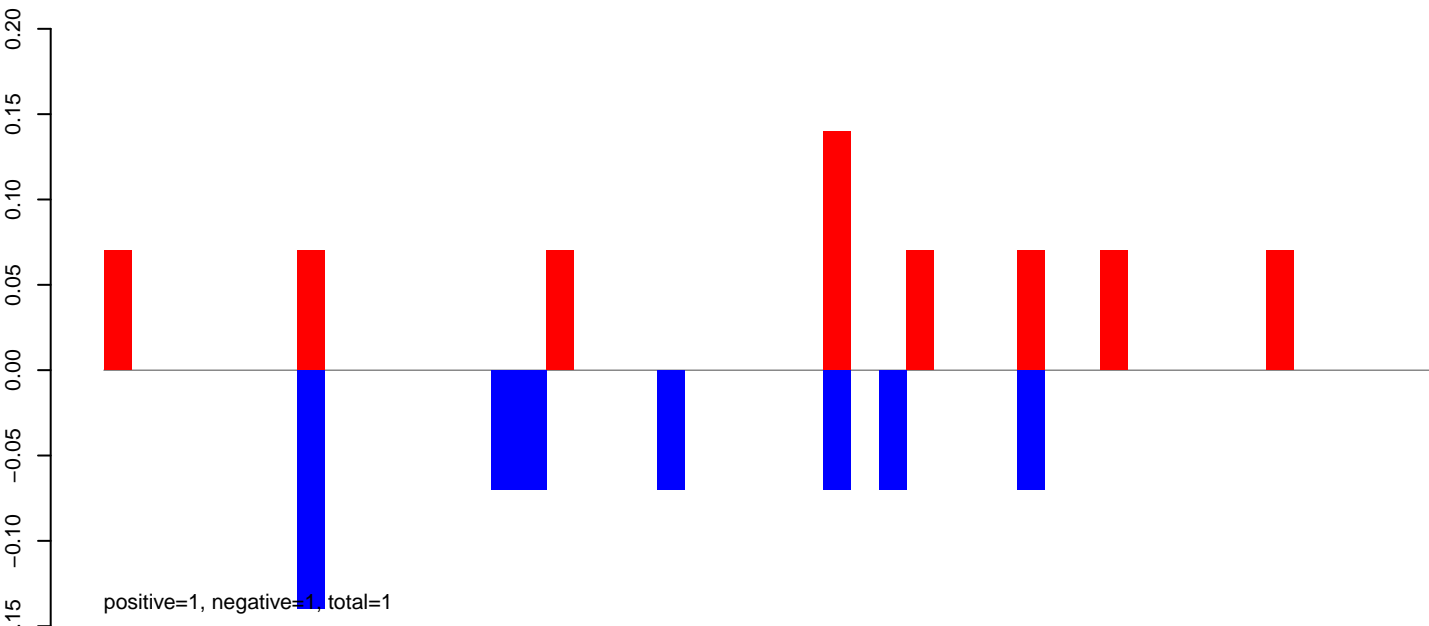


CuQuin_Mcarc_NL.rep

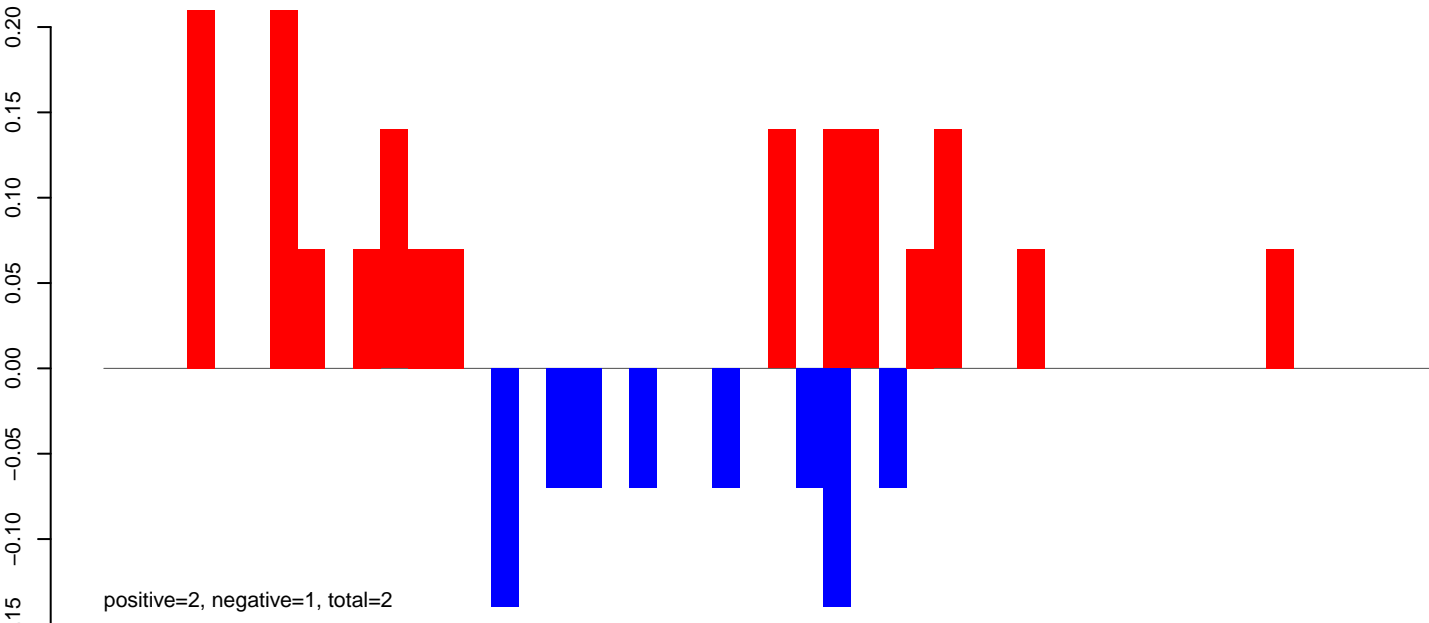


Window size=25, length=790, TE@rnd=4, family=481@Unknown@785:1-790

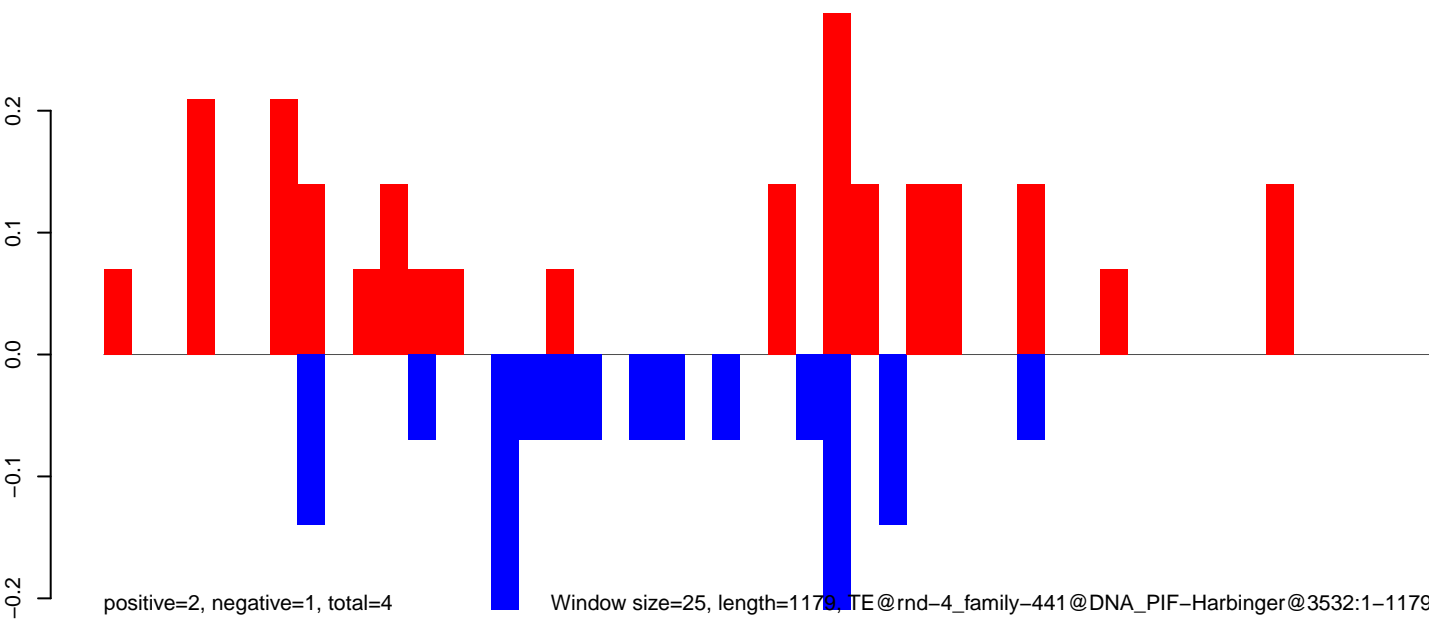
CuQuin_Mcarc_NL.18_23.rep



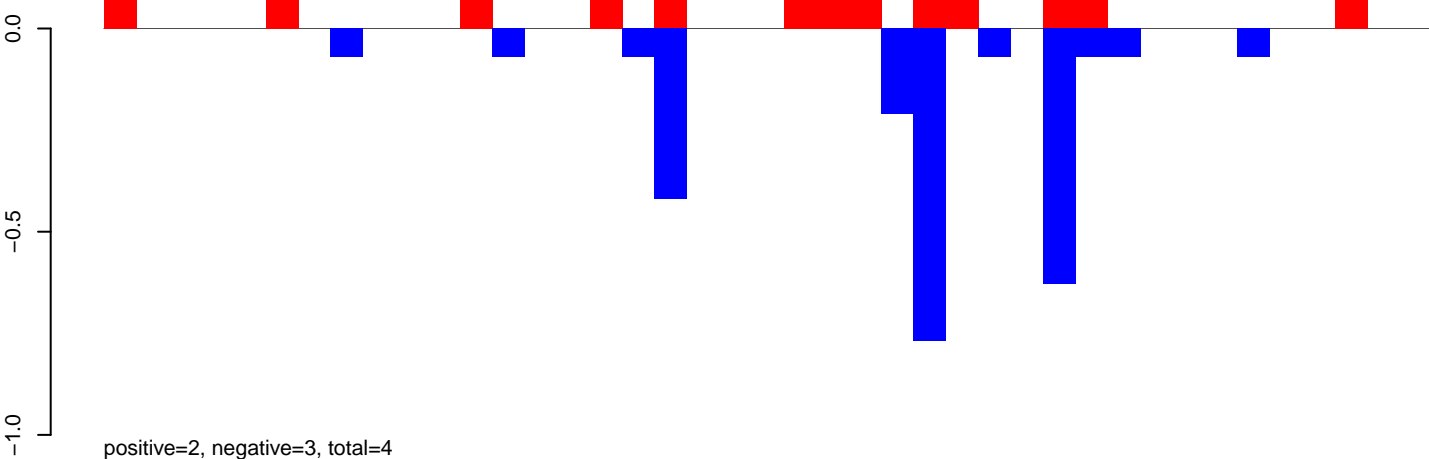
CuQuin_Mcarc_NL.24_35.rep



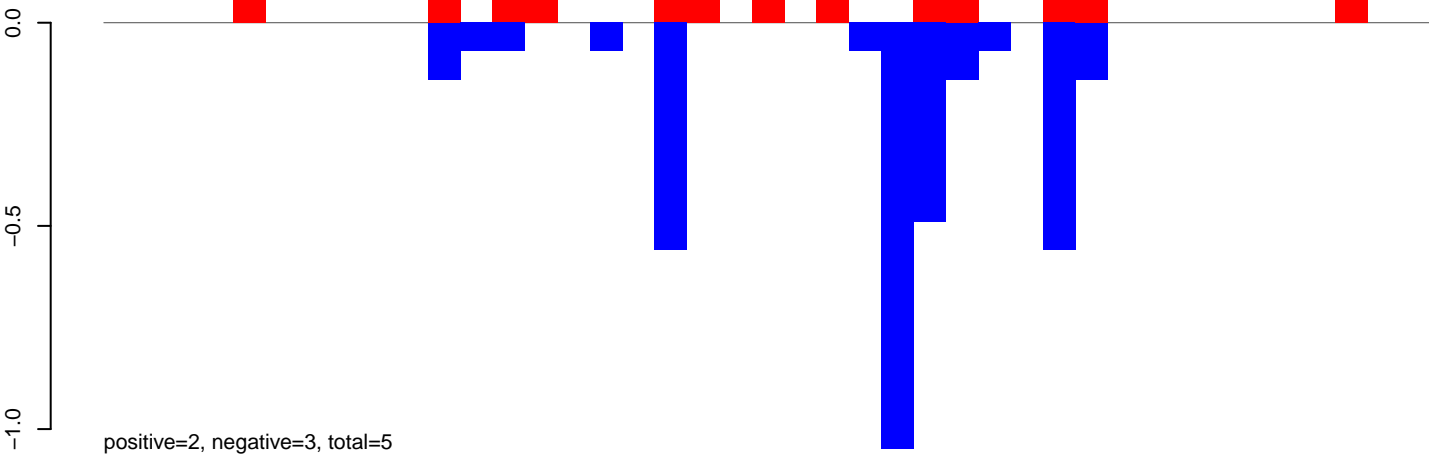
CuQuin_Mcarc_NL.rep



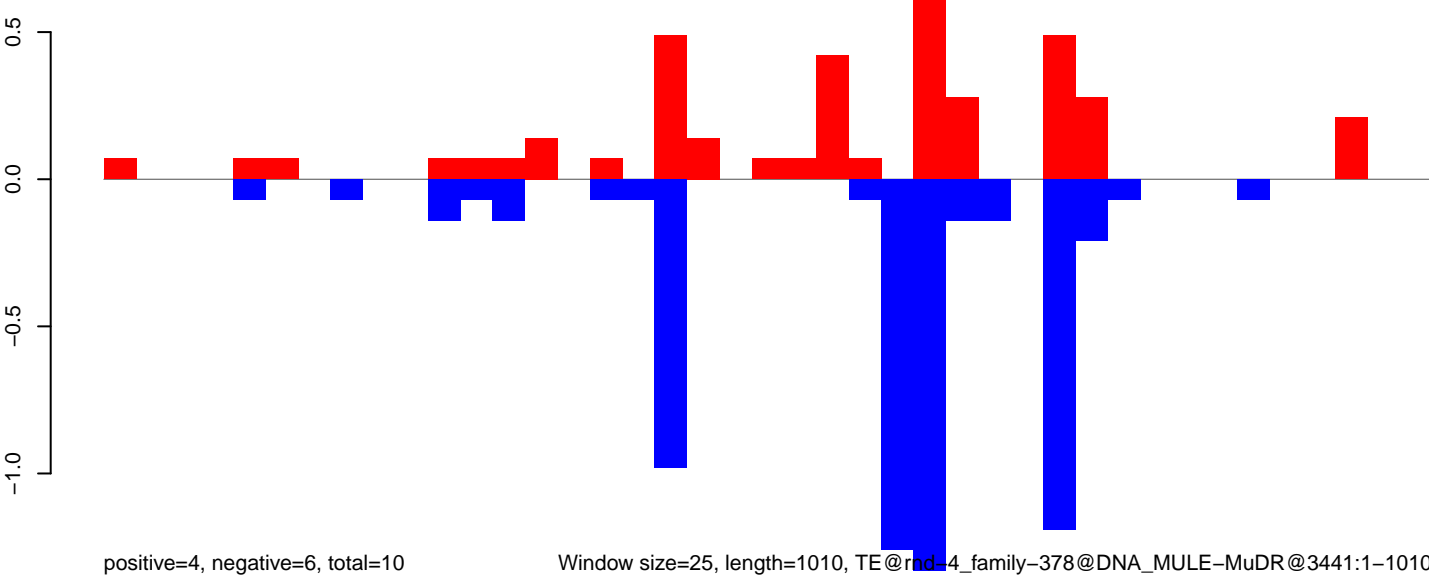
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



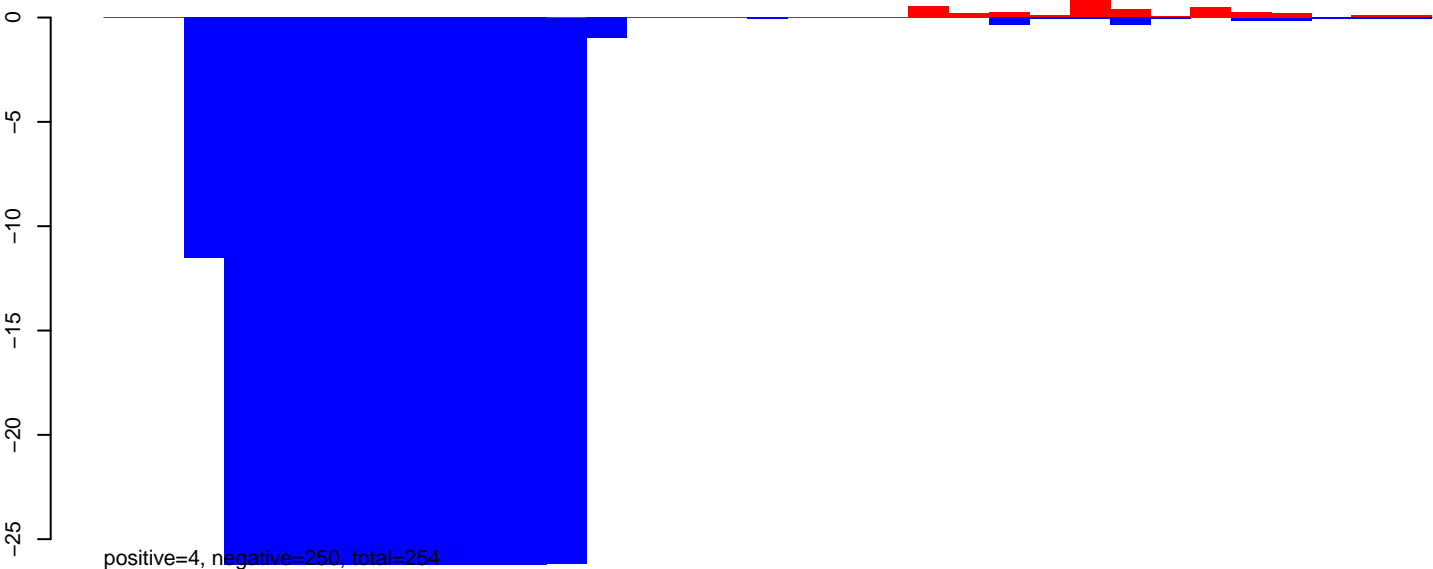
CuQuin_Mcarc_NL.rep



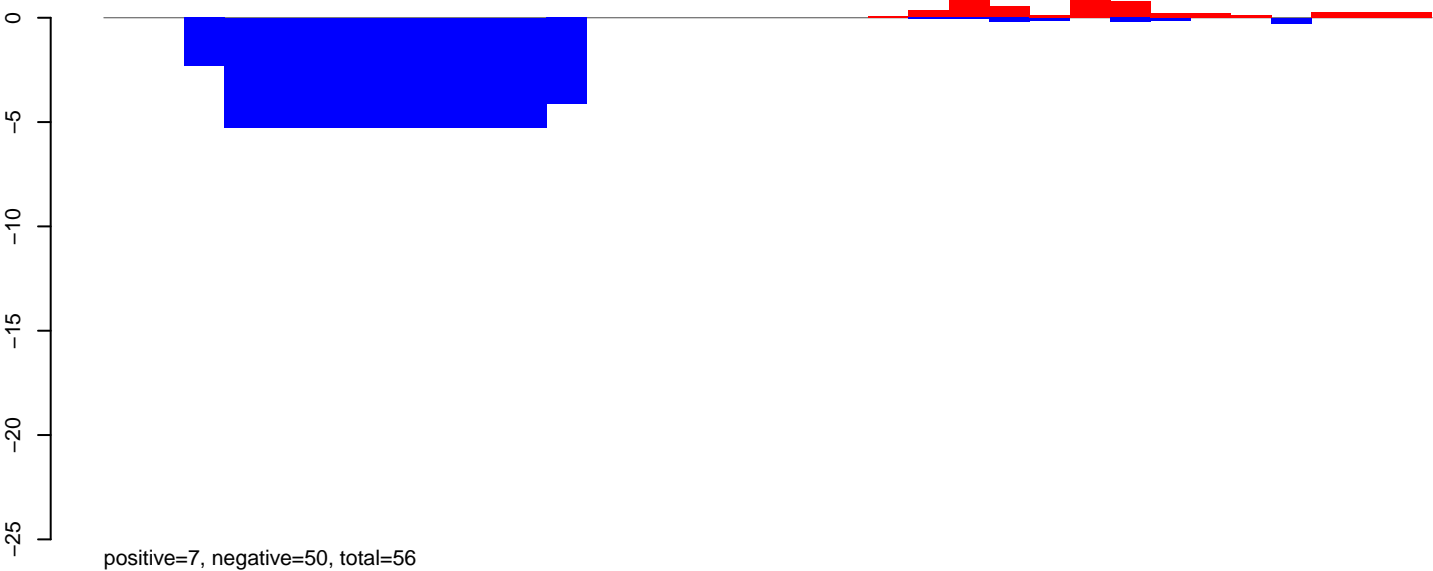
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0 200 400 600 800 1000

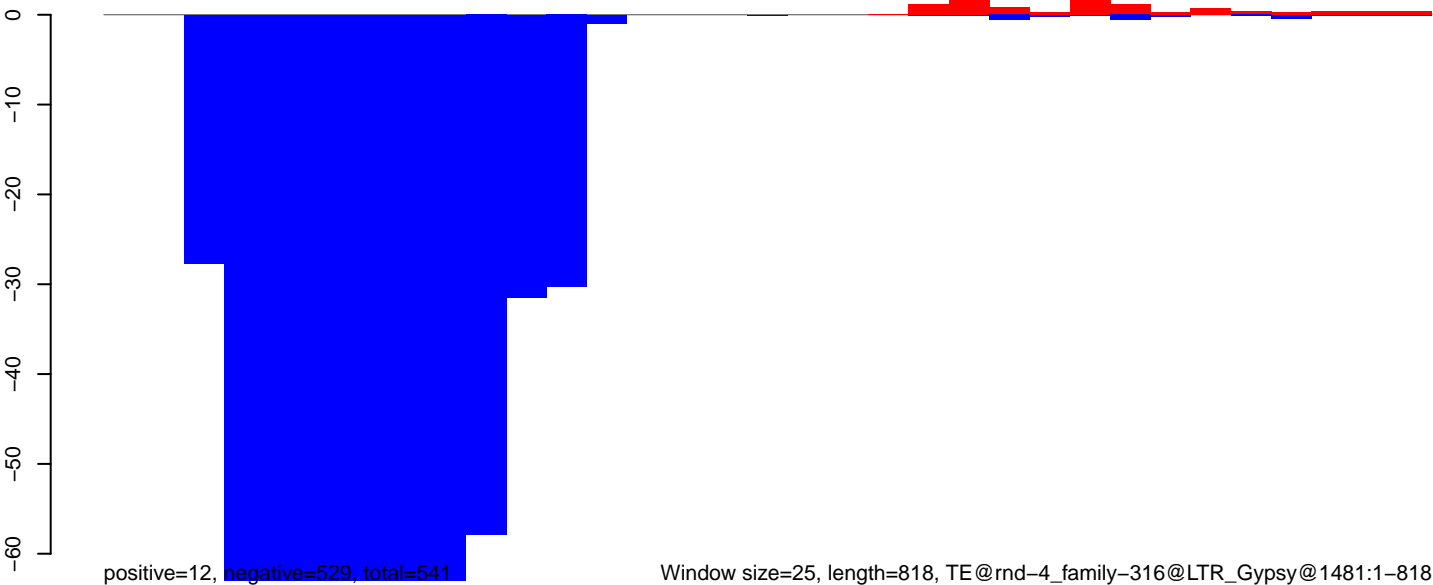
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CuQuin_Mcarc_NL.24_35.rep

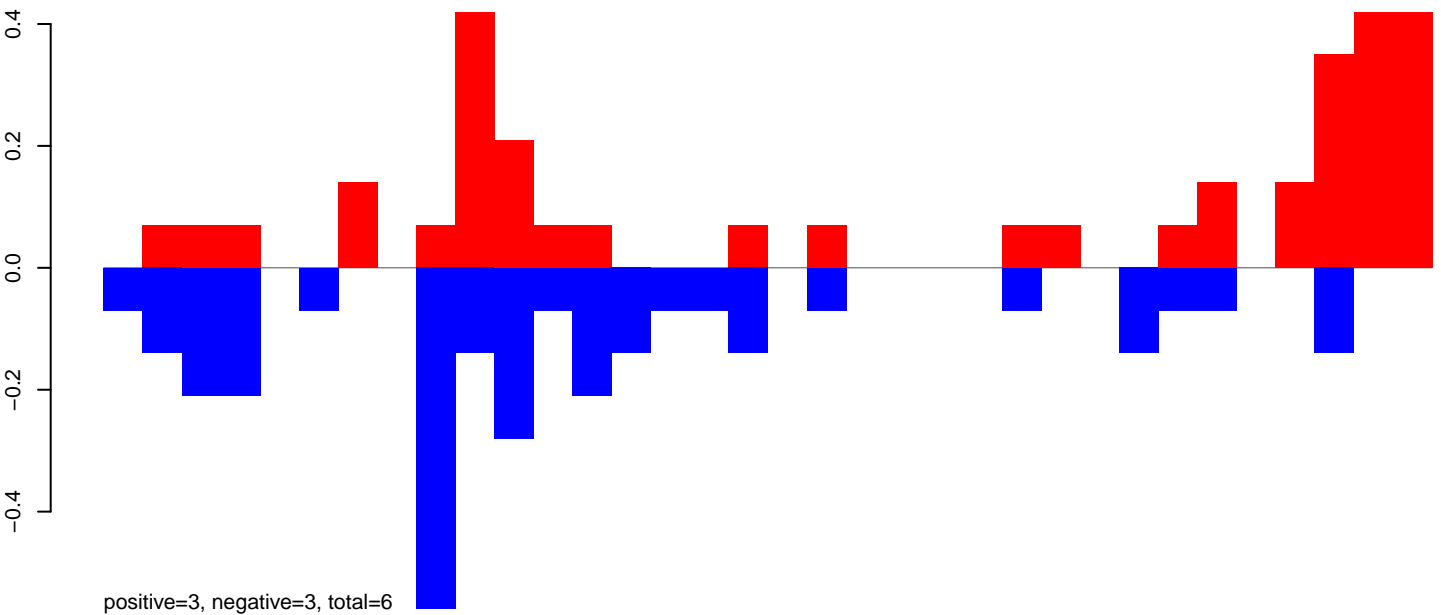


CuQuin_Mcarc_NL.rep

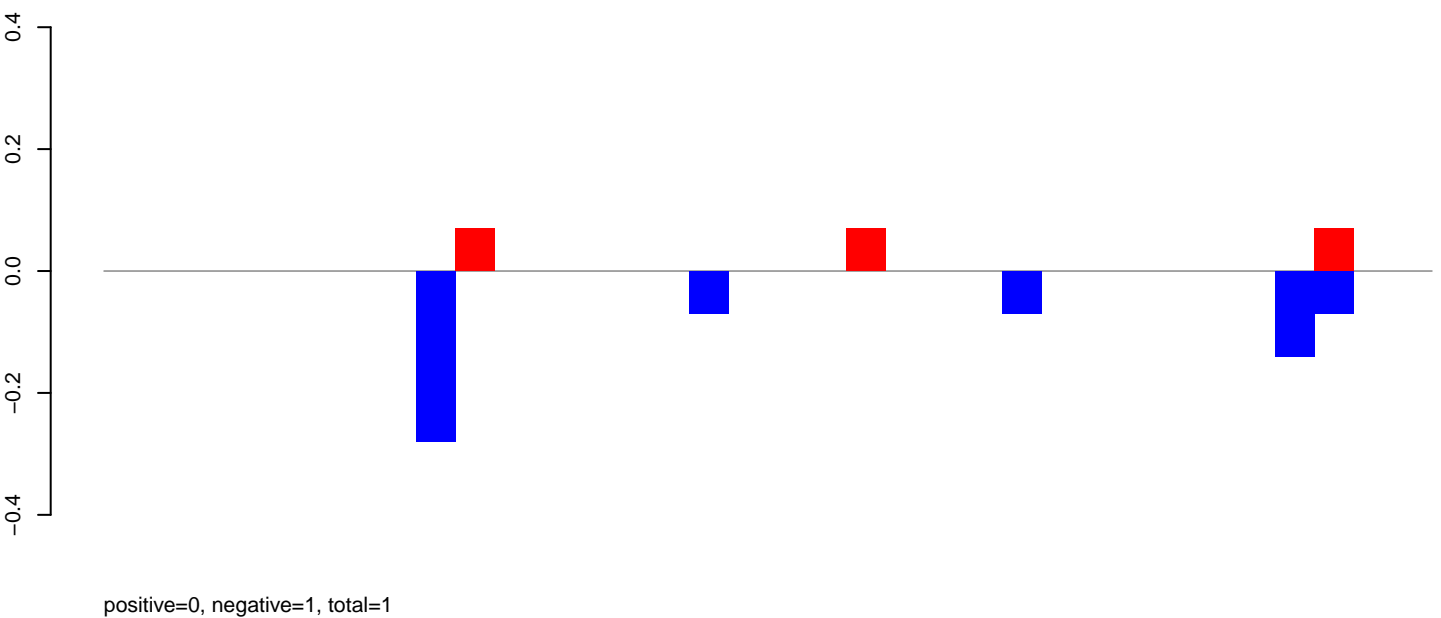


Window size=25, length=818, TE@rnd-4_family-316@LTR_Gypsy@1481:1-818

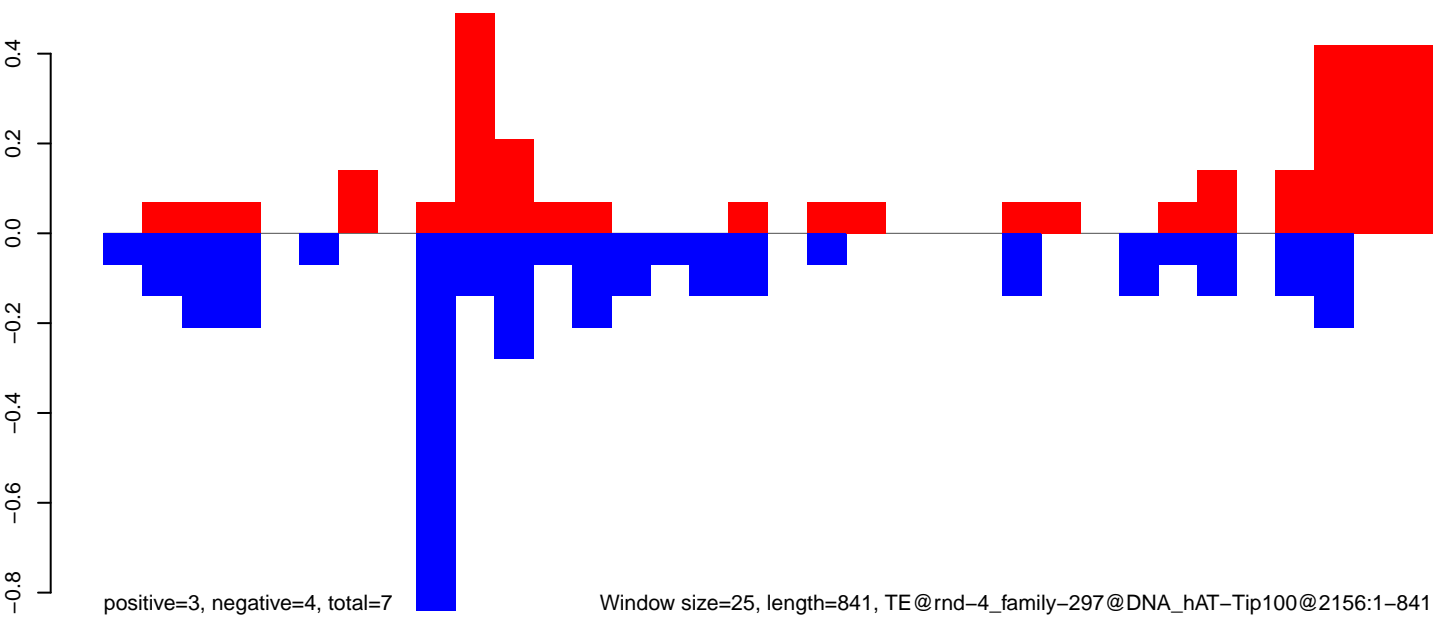
CuQuin_Mcarc_NL.18_23.rep



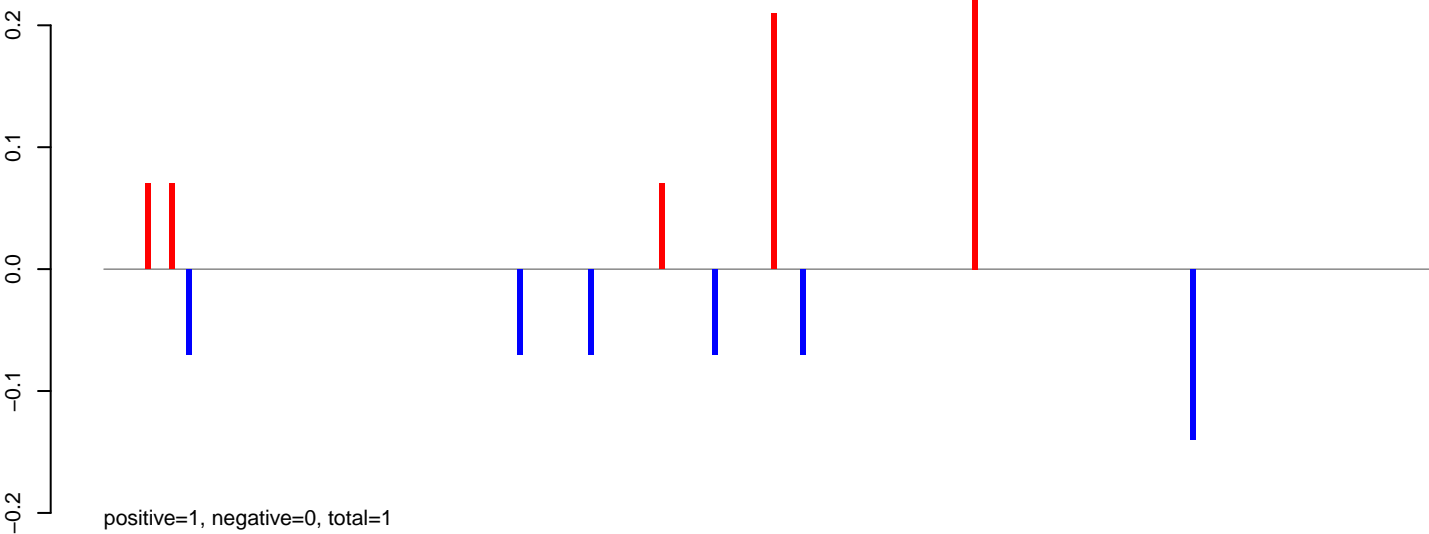
CuQuin_Mcarc_NL.24_35.rep



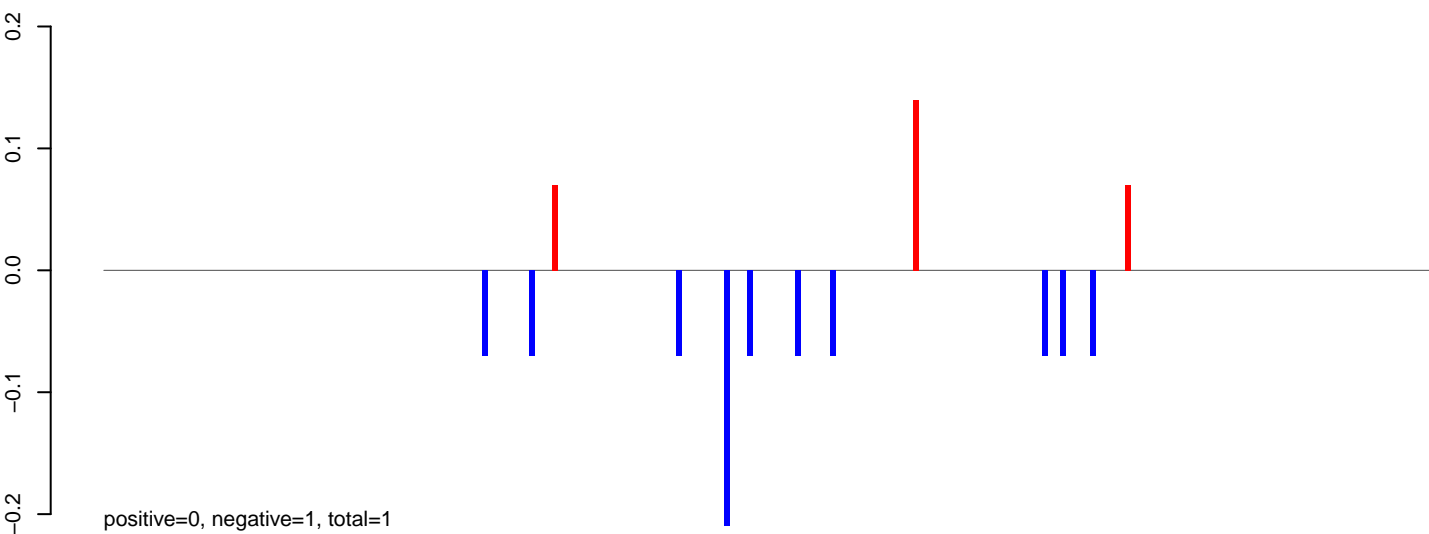
CuQuin_Mcarc_NL.rep



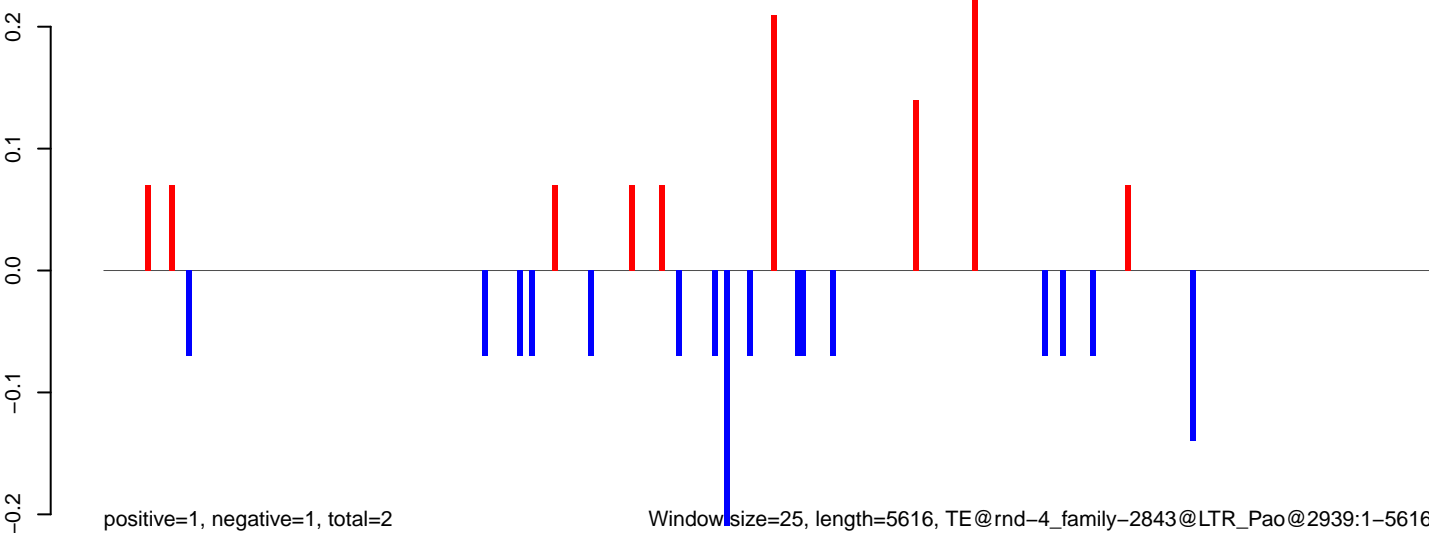
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



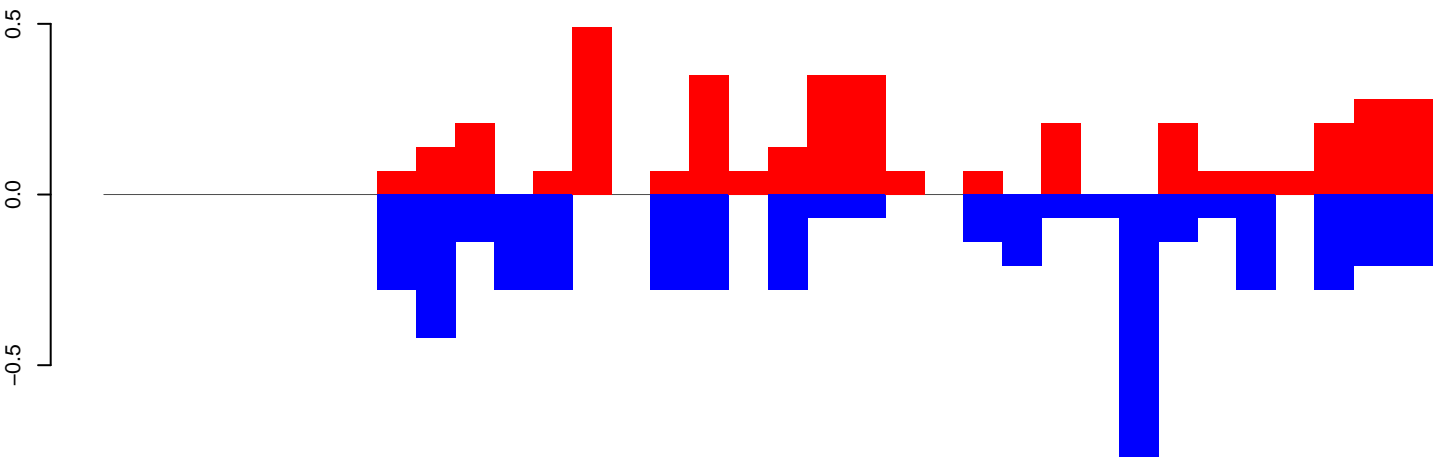
CuQuin_Mcarc_NL.rep



Window size=25, length=5616, TE@rnd-4_family-2843@LTR_Pao@2939:1-5616

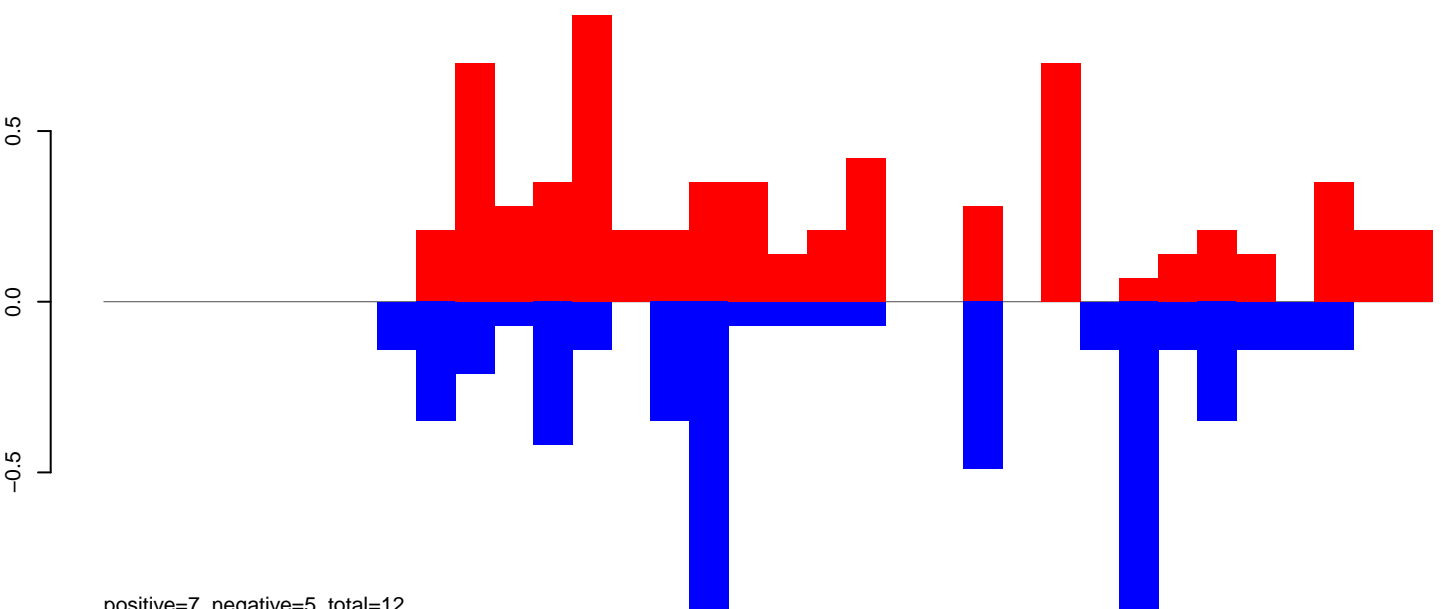
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CuQuin_Mcarc_NL.18_23.rep



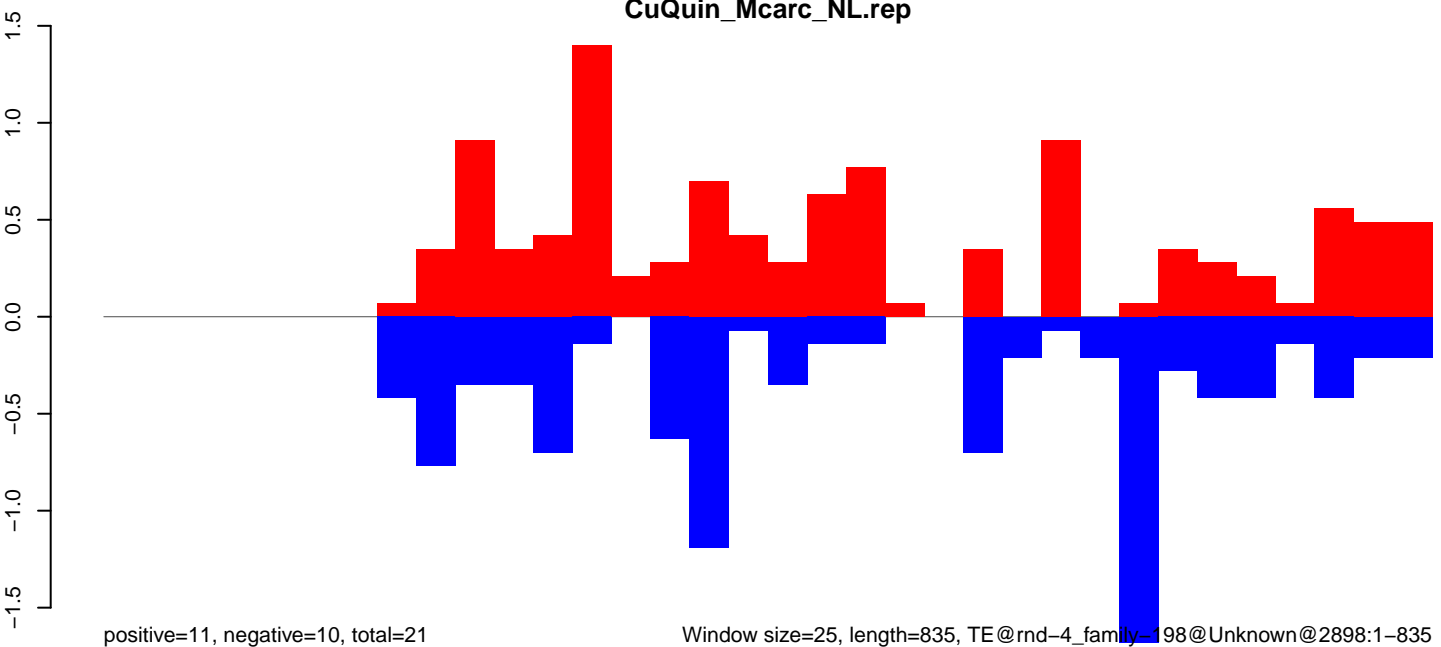
positive=4, negative=5, total=9

CuQuin_Mcarc_NL.24_35.rep



positive=7, negative=5, total=12

CuQuin_Mcarc_NL.rep

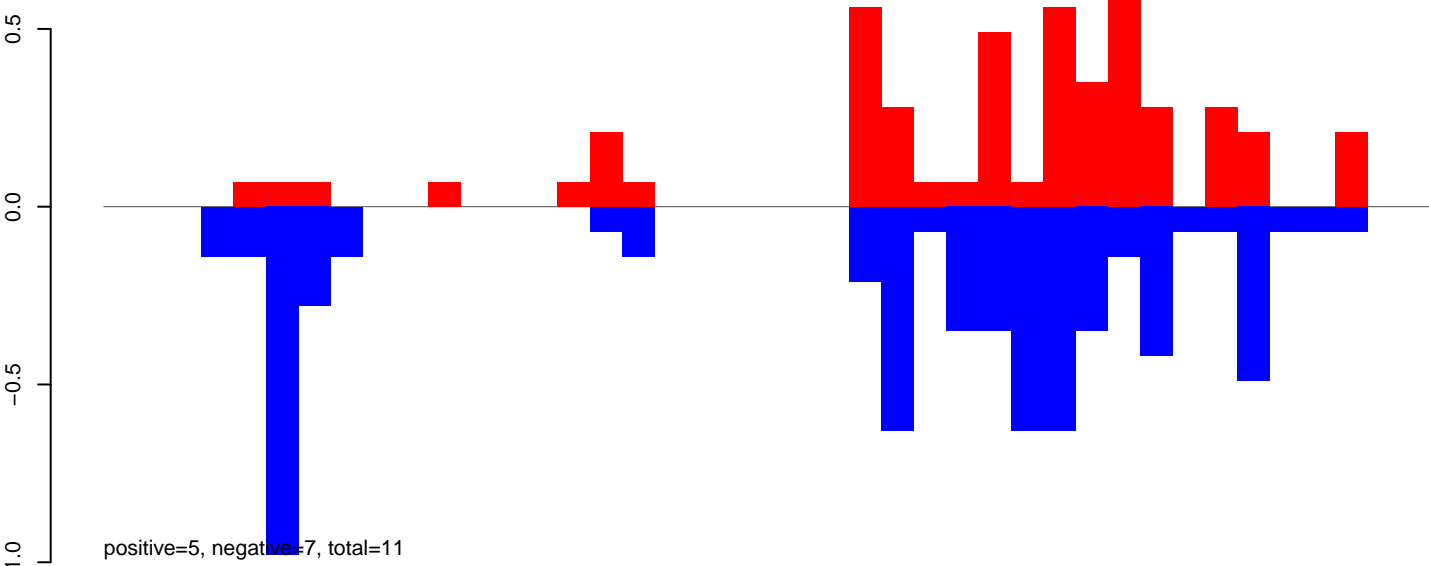


positive=11, negative=10, total=21

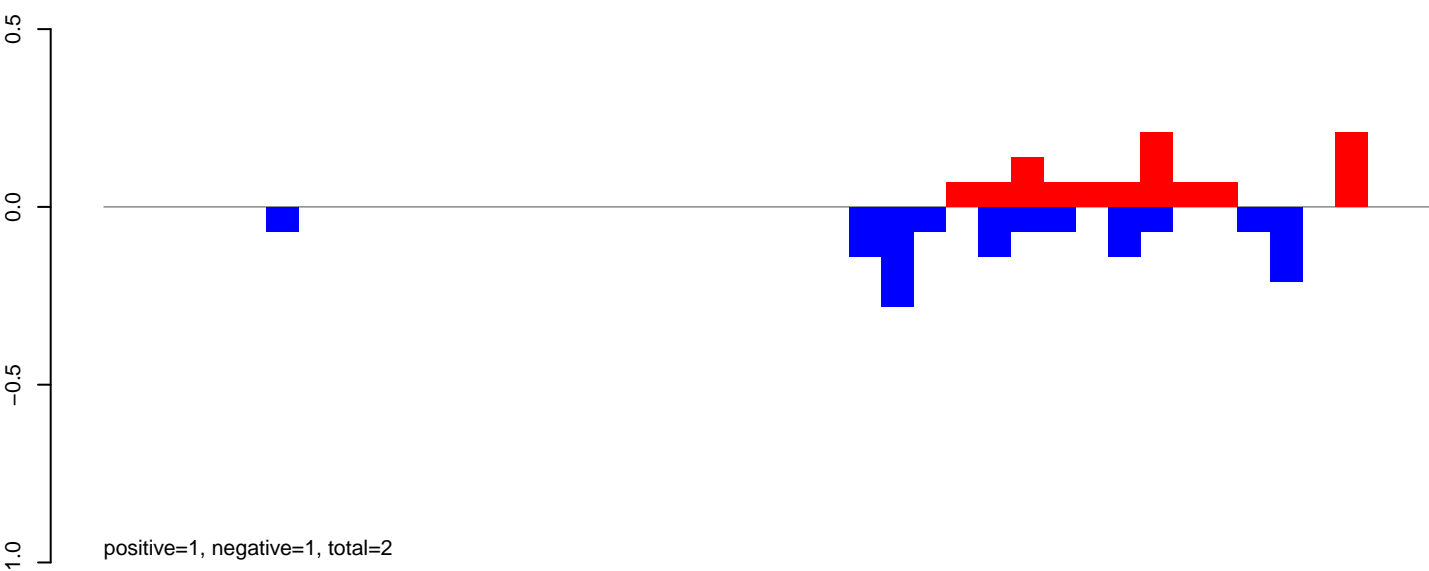
Window size=25, length=835, TE@rnd-4_family-198@Unknown@2898:1-835

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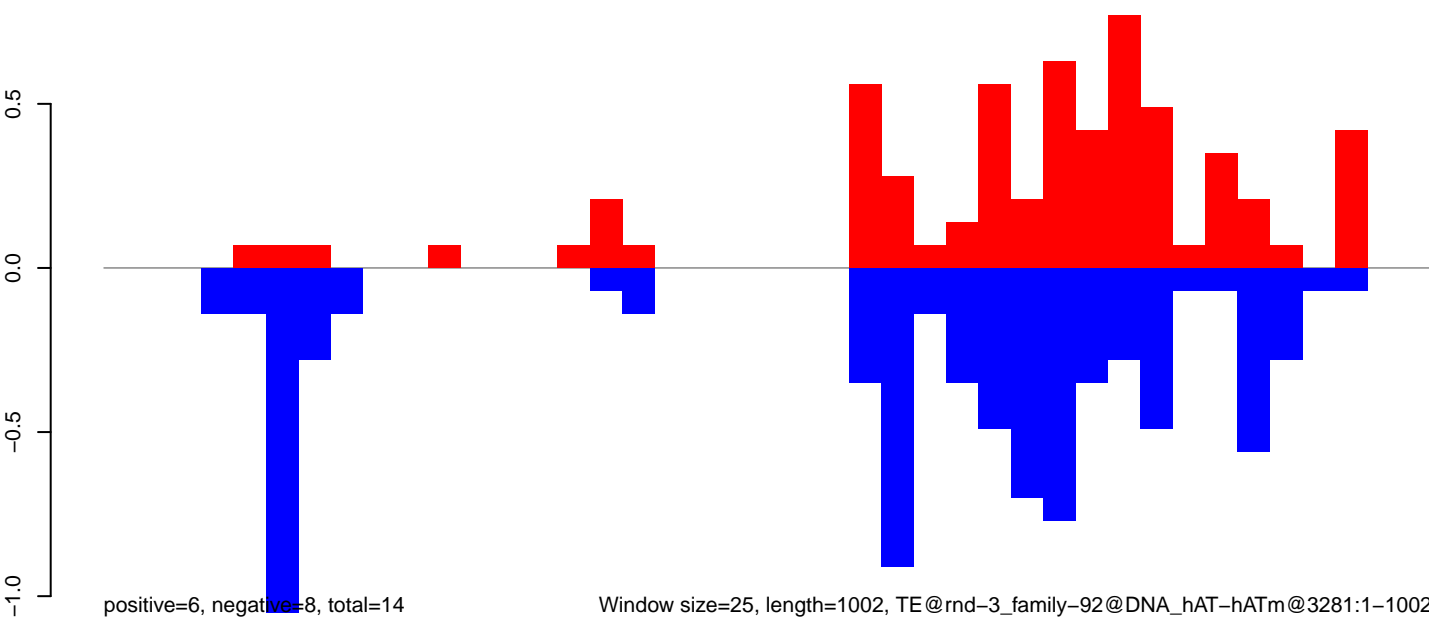
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



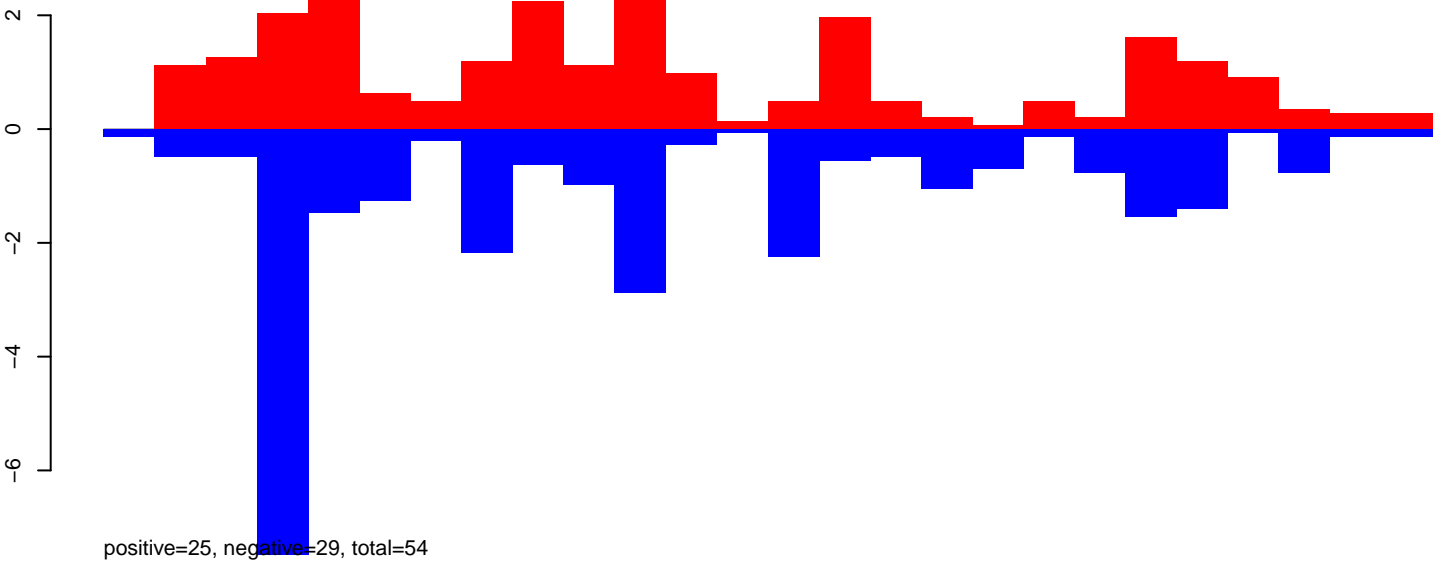
CuQuin_Mcarc_NL.rep



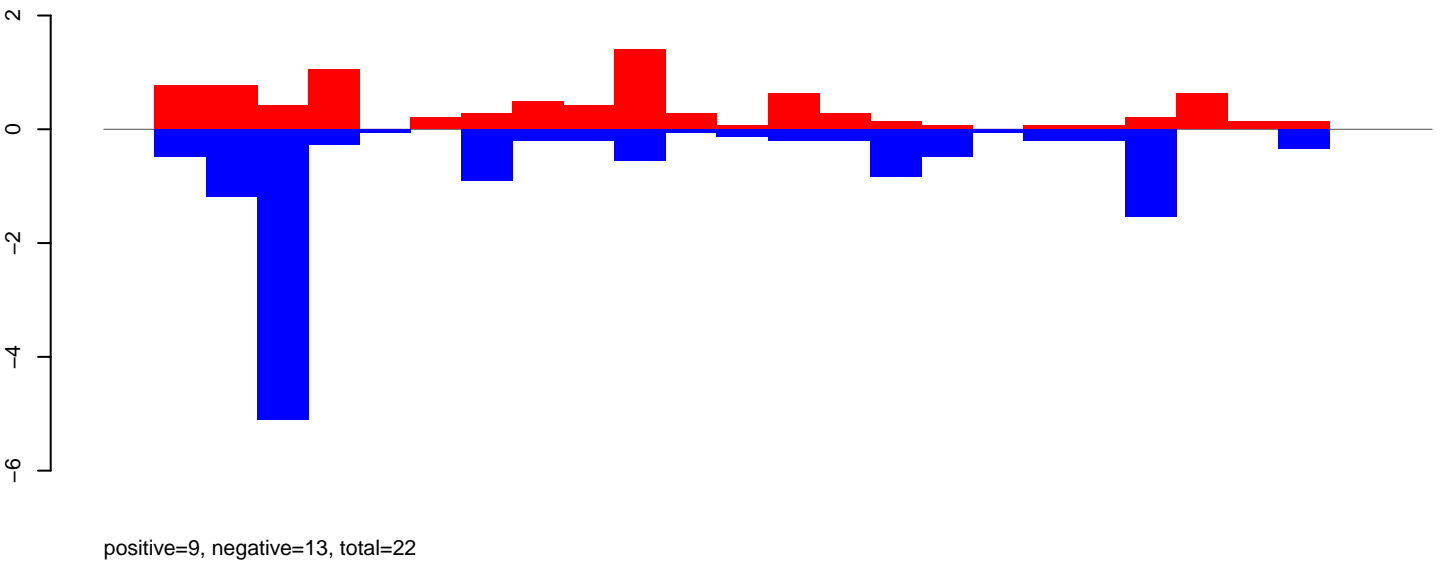
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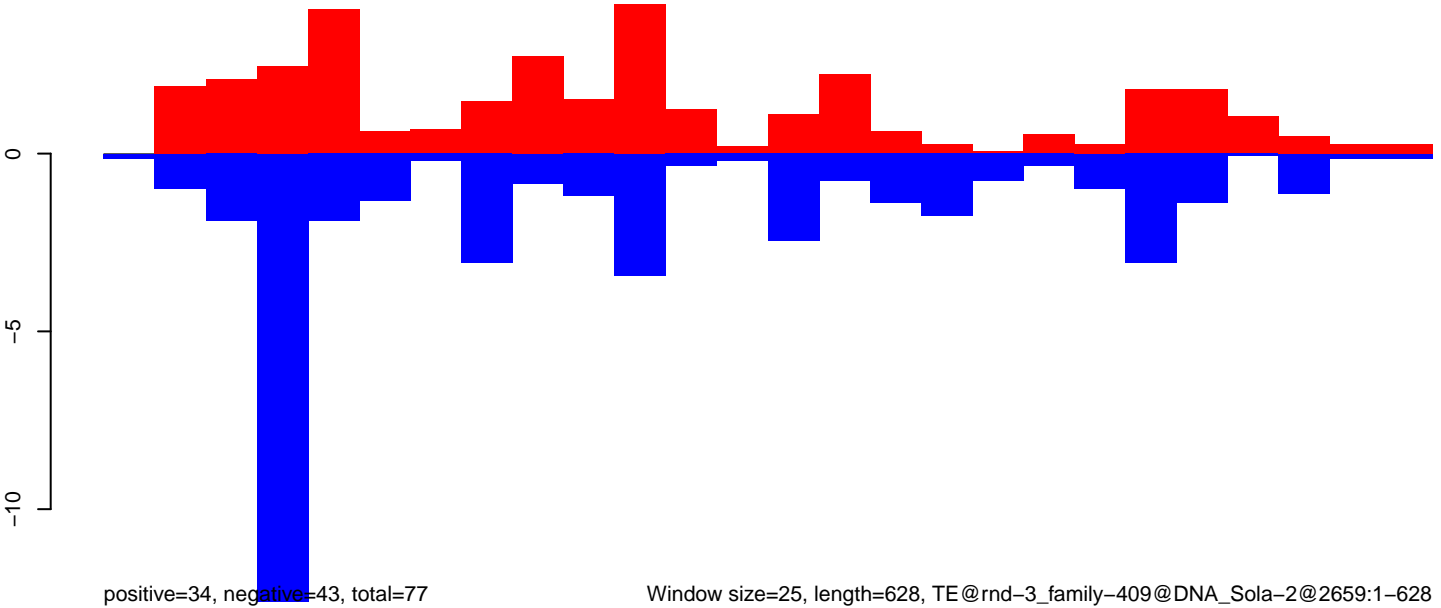
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



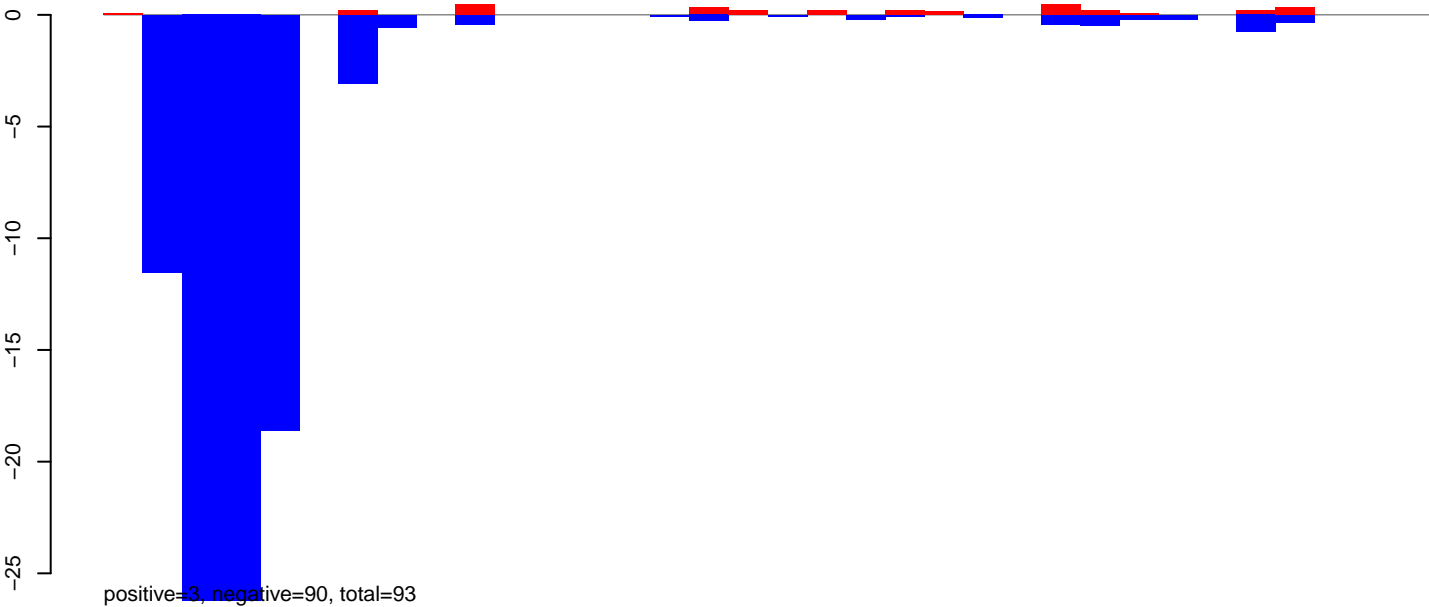
CuQuin_Mcarc_NL.rep



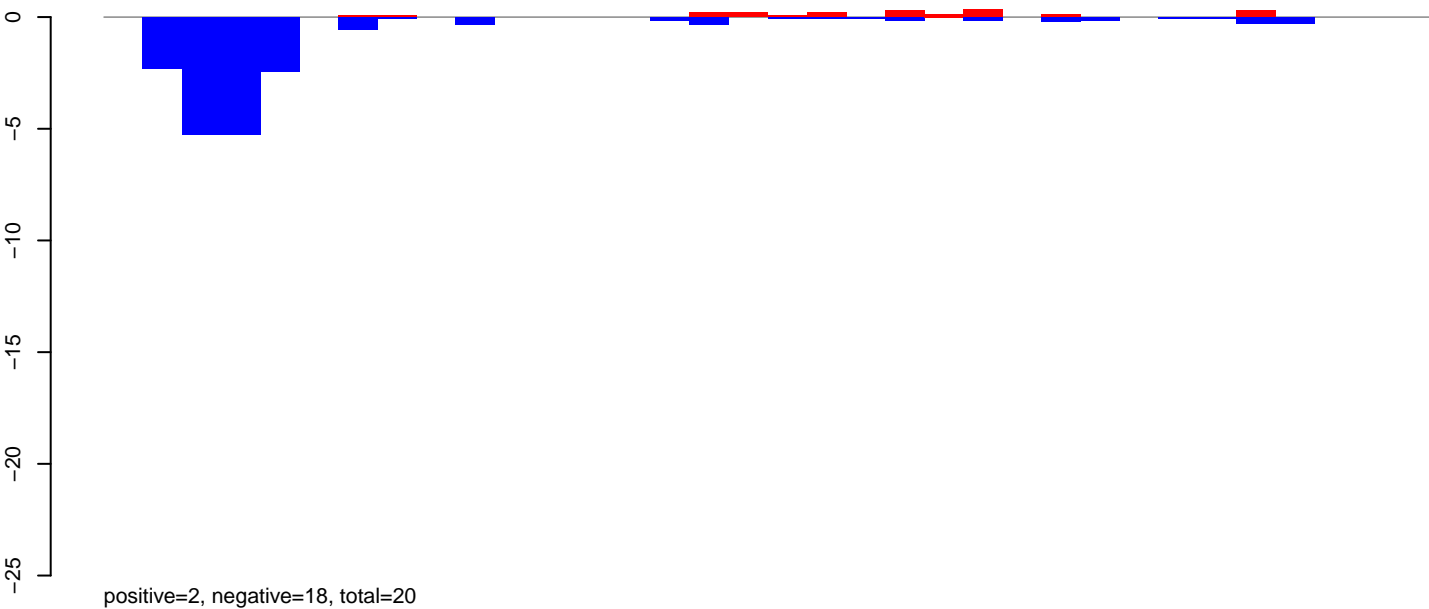
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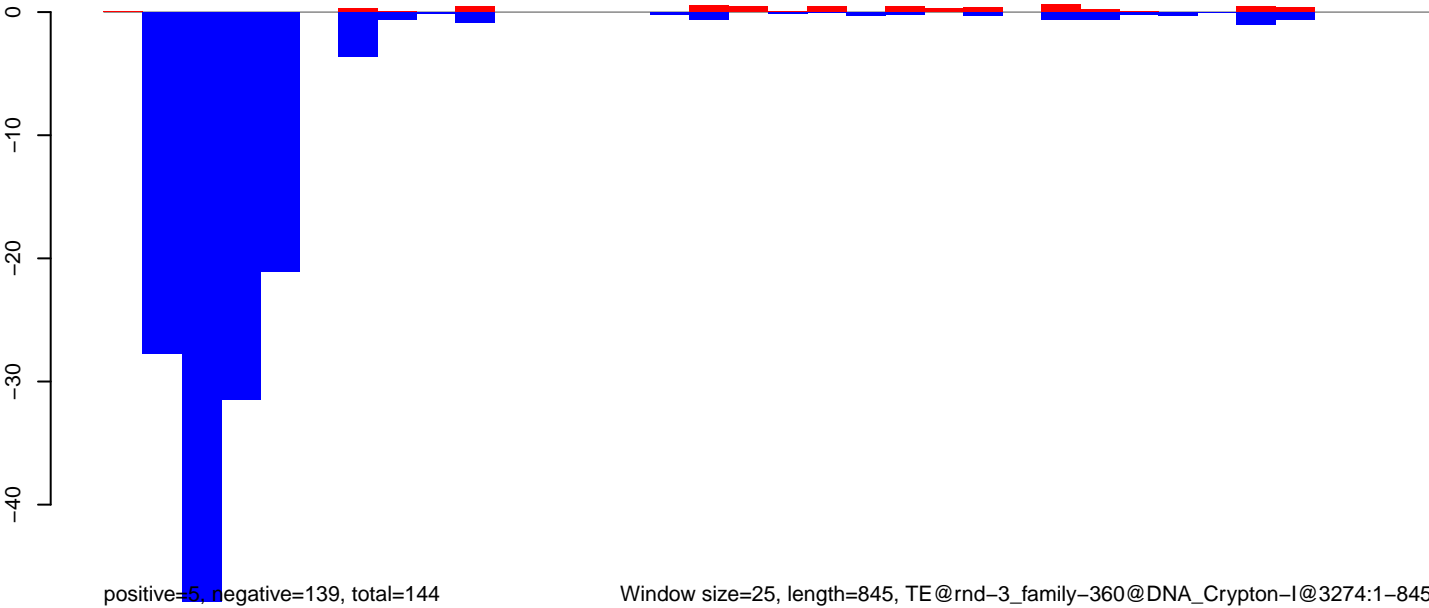
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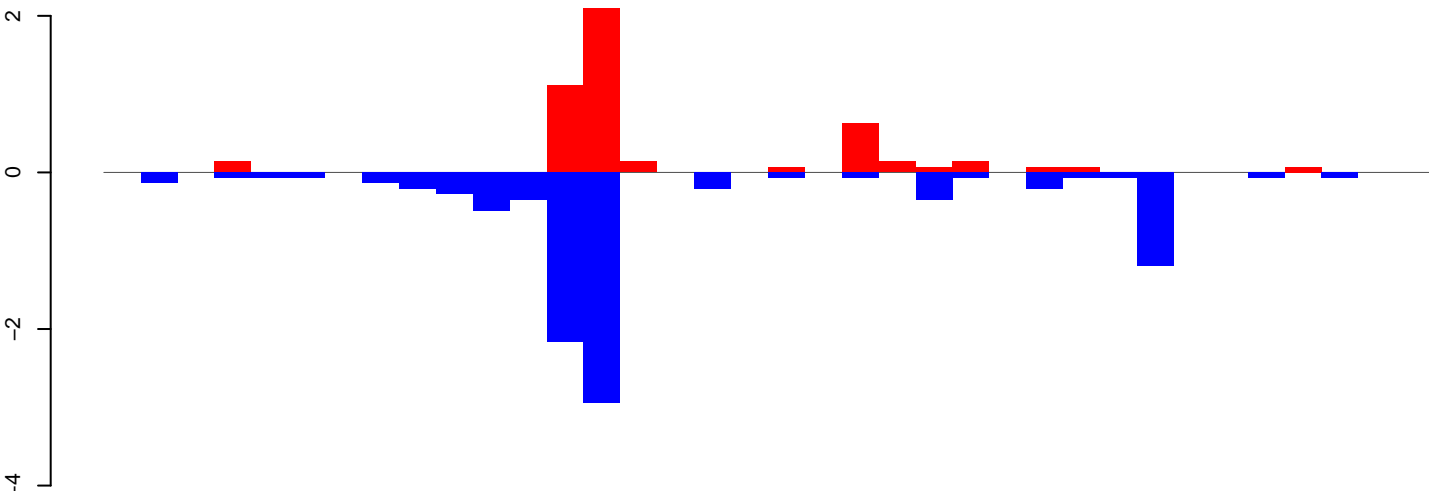
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

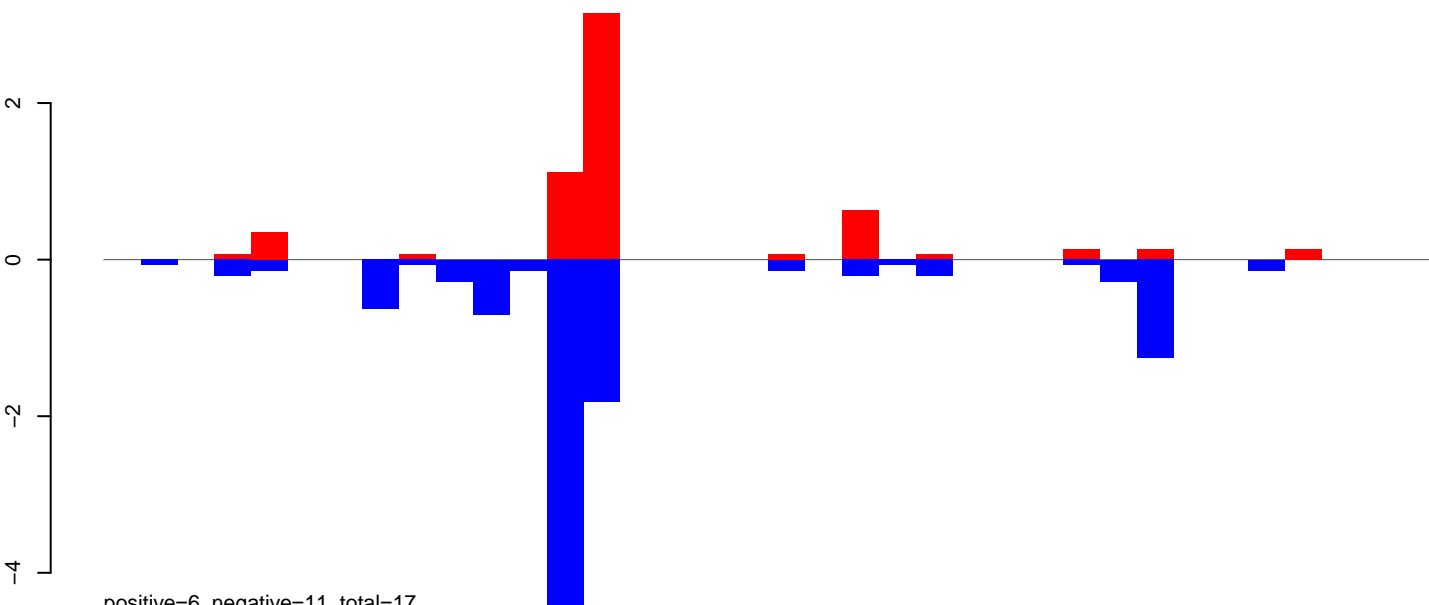


CuQuin_Mcarc_NL.18_23.rep



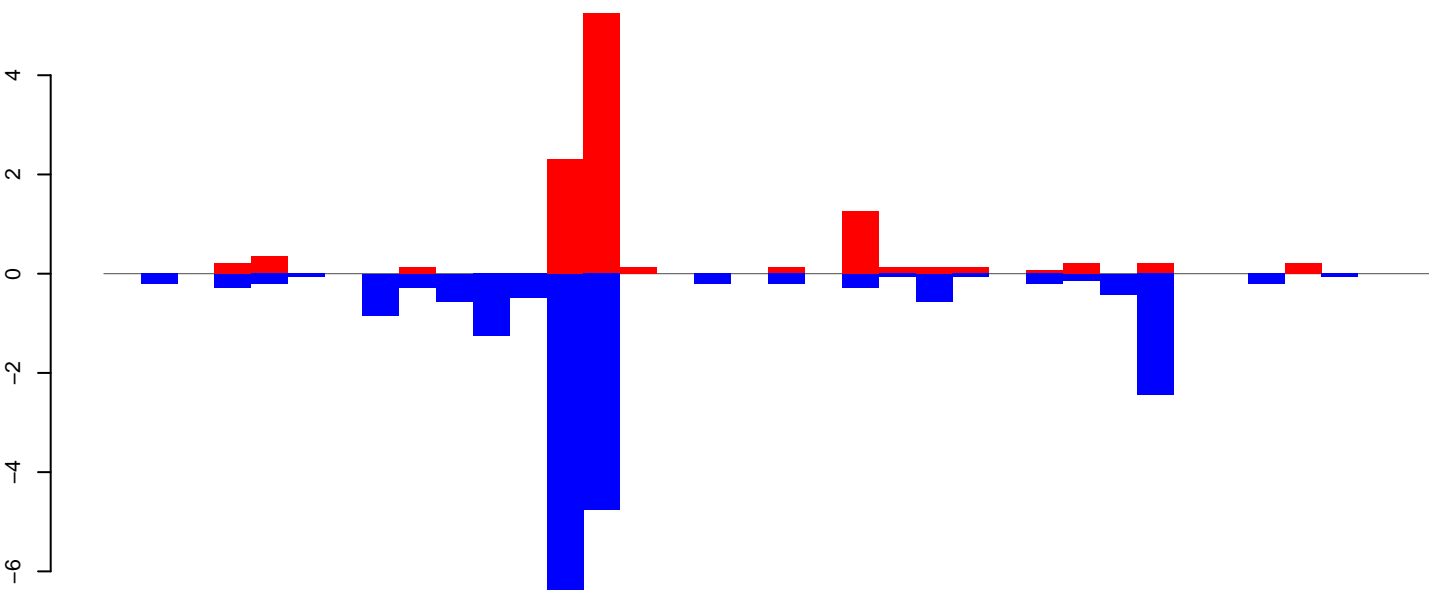
positive=5, negative=9, total=14

CuQuin_Mcarc_NL.24_35.rep



positive=6, negative=11, total=17

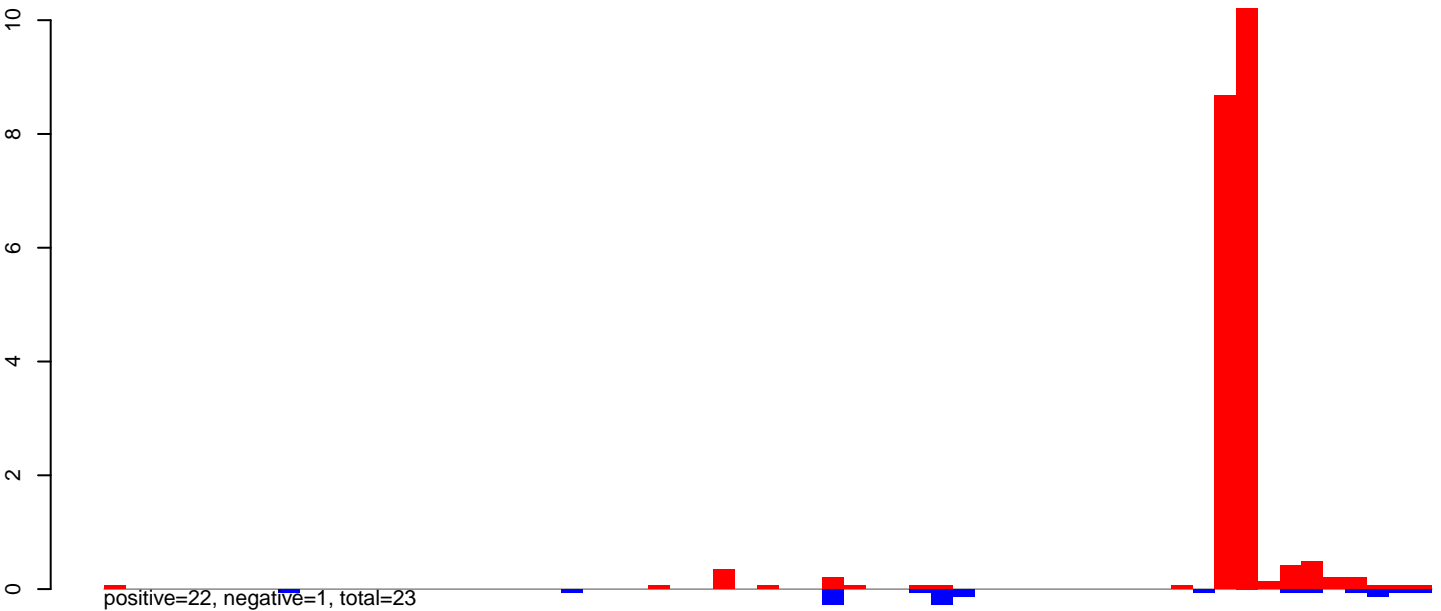
CuQuin_Mcarc_NL.rep



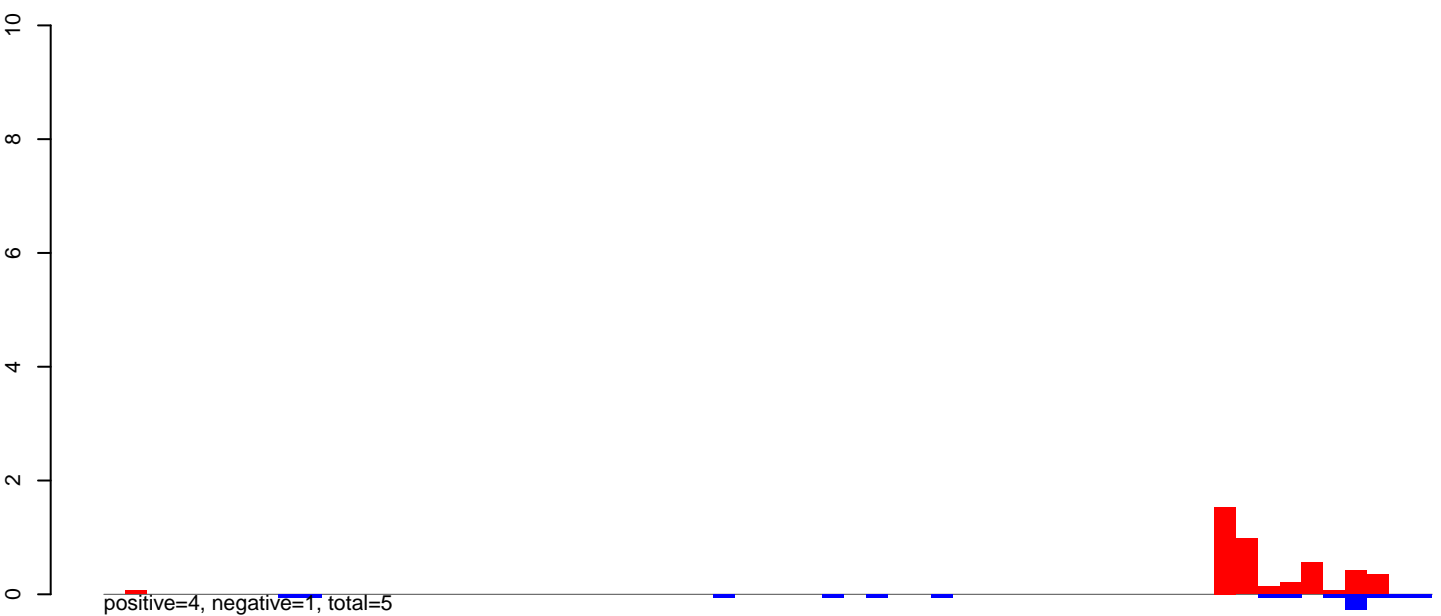
positive=11, negative=21, total=32

Window size=25, length=881, TE@rnd-3_family-287@Unknown@1284:1-881

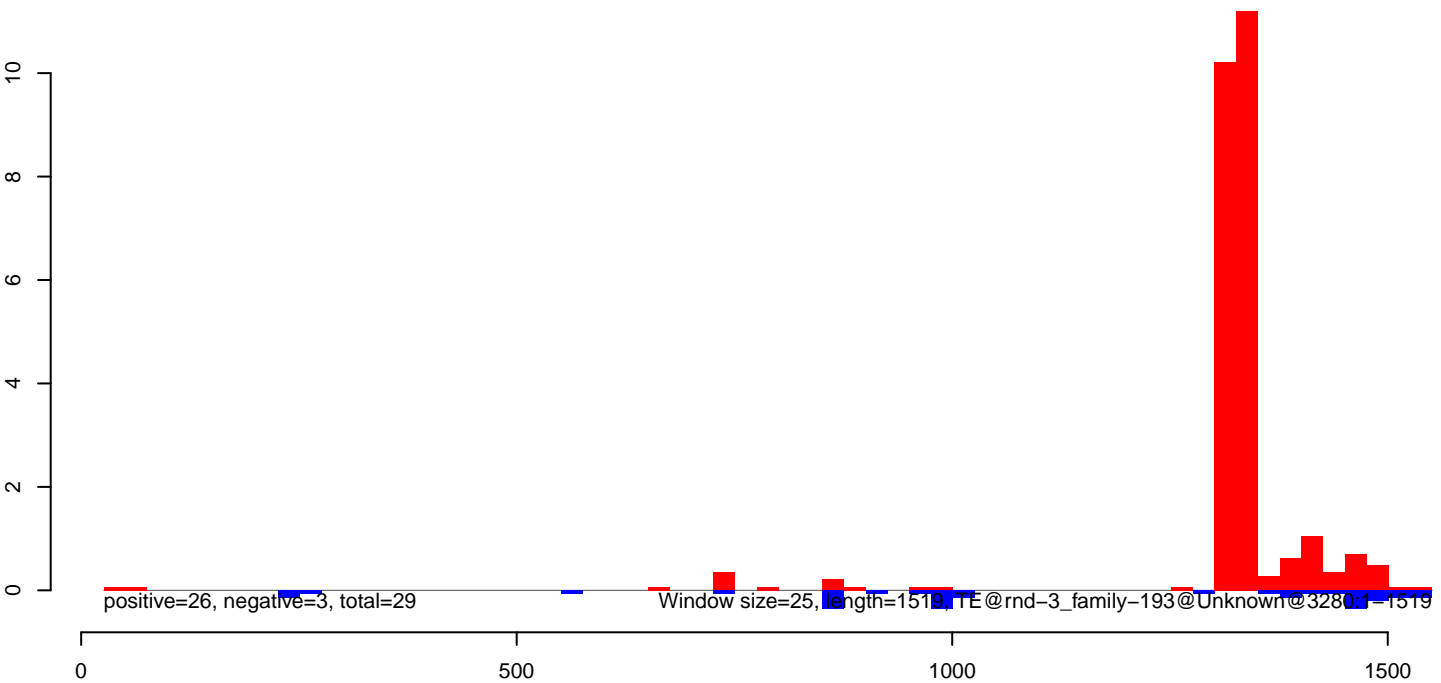
CuQuin_Mcarc_NL.18_23.rep



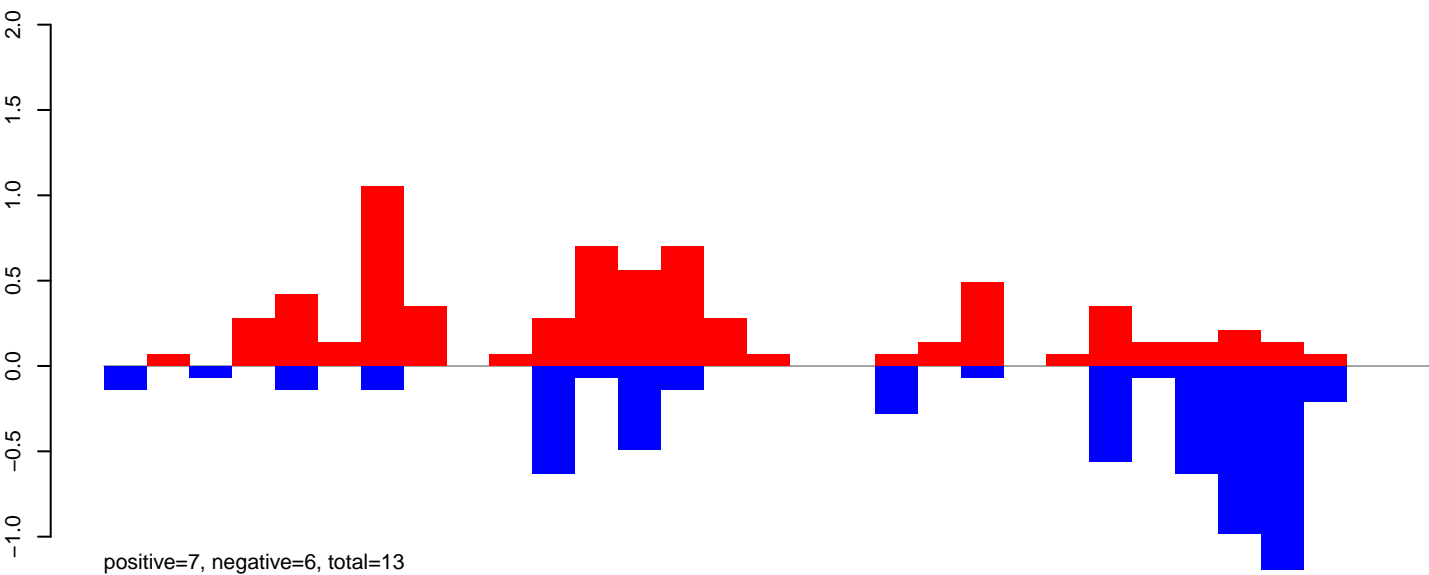
CuQuin_Mcarc_NL.24_35.rep



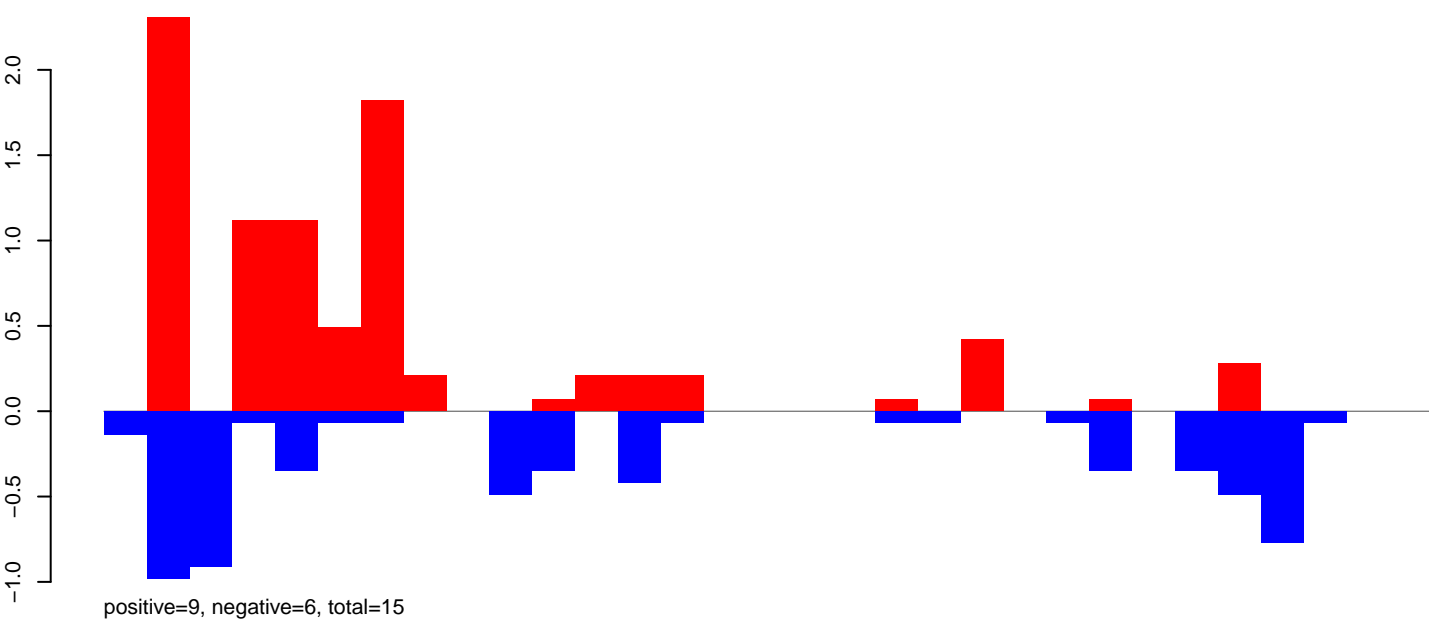
CuQuin_Mcarc_NL.rep



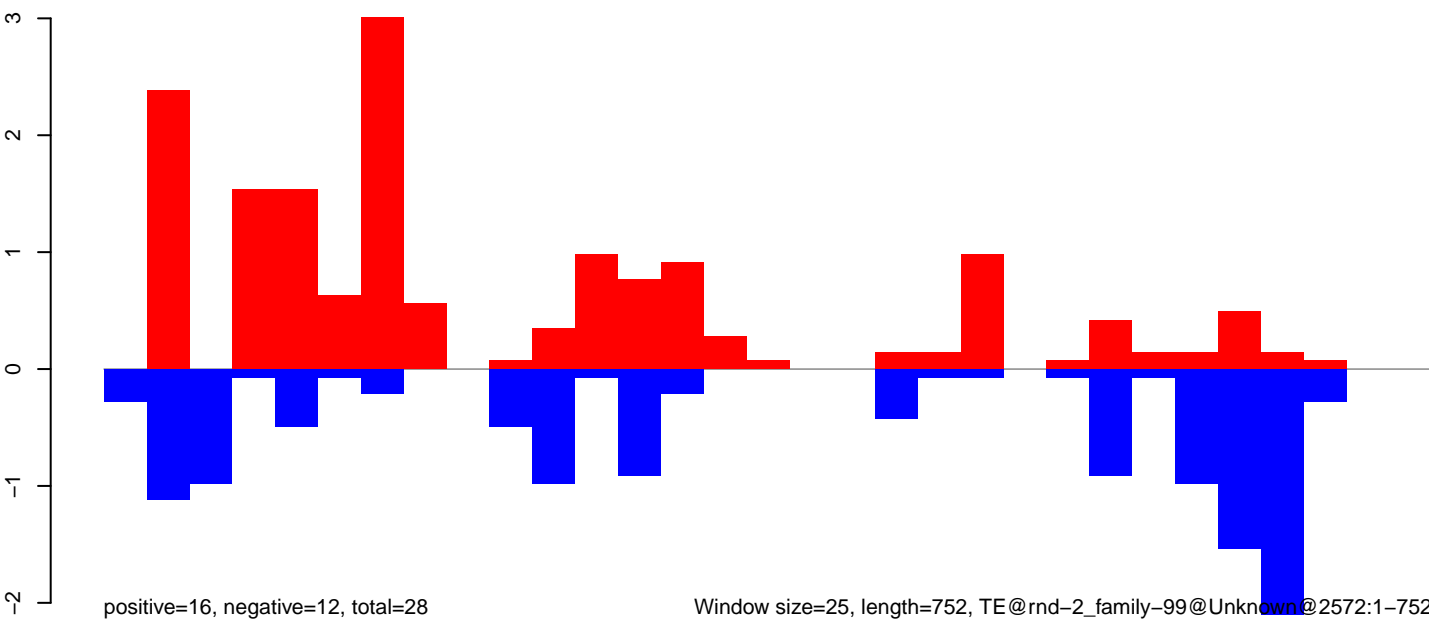
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CuQuin_Mcarc_NL.24_35.rep



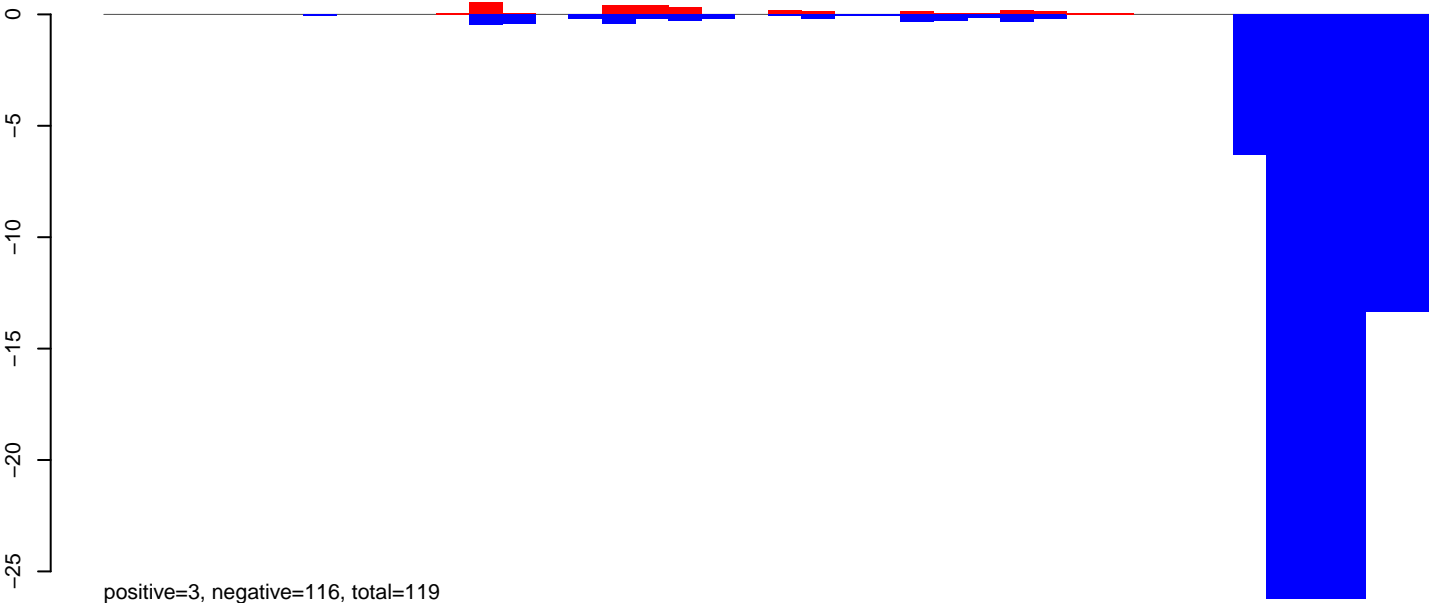
CuQuin_Mcarc_NL.rep



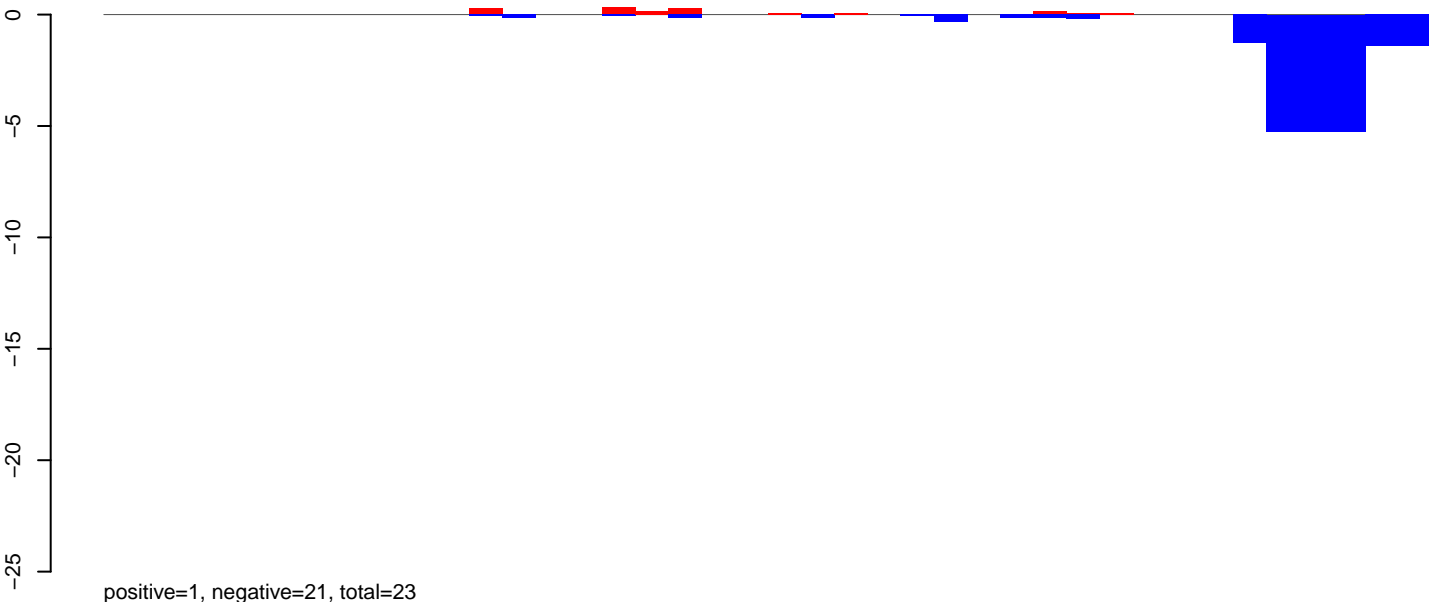
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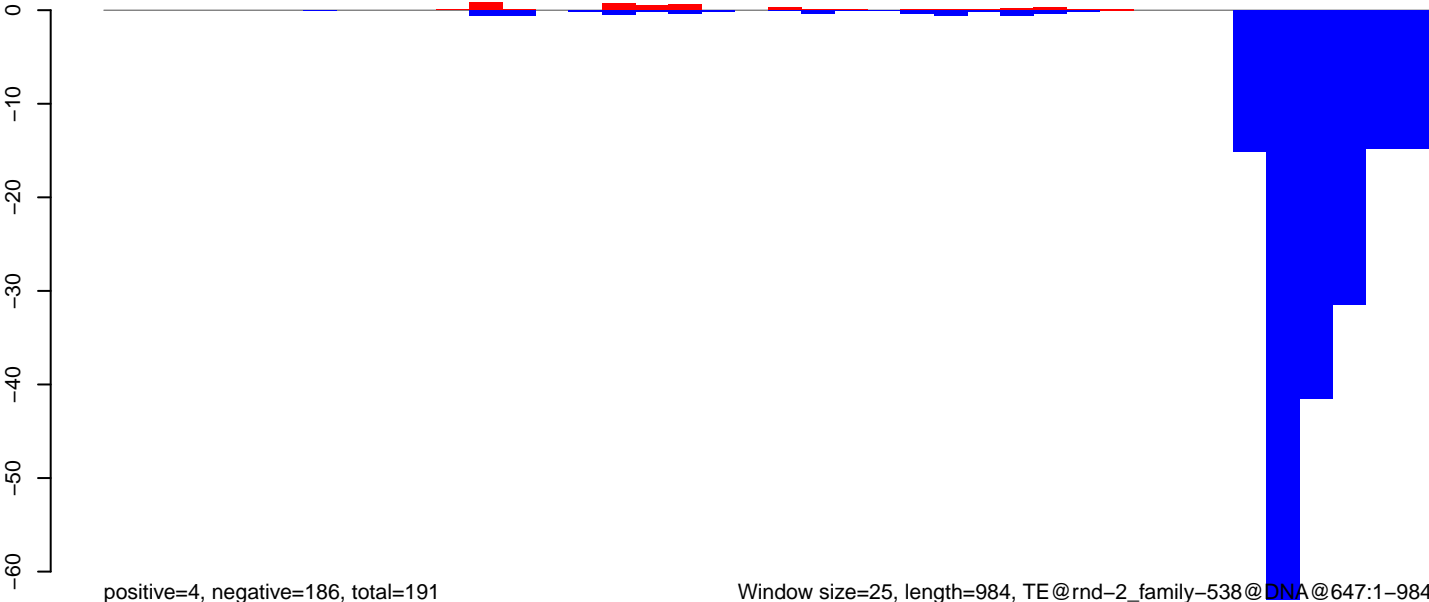
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

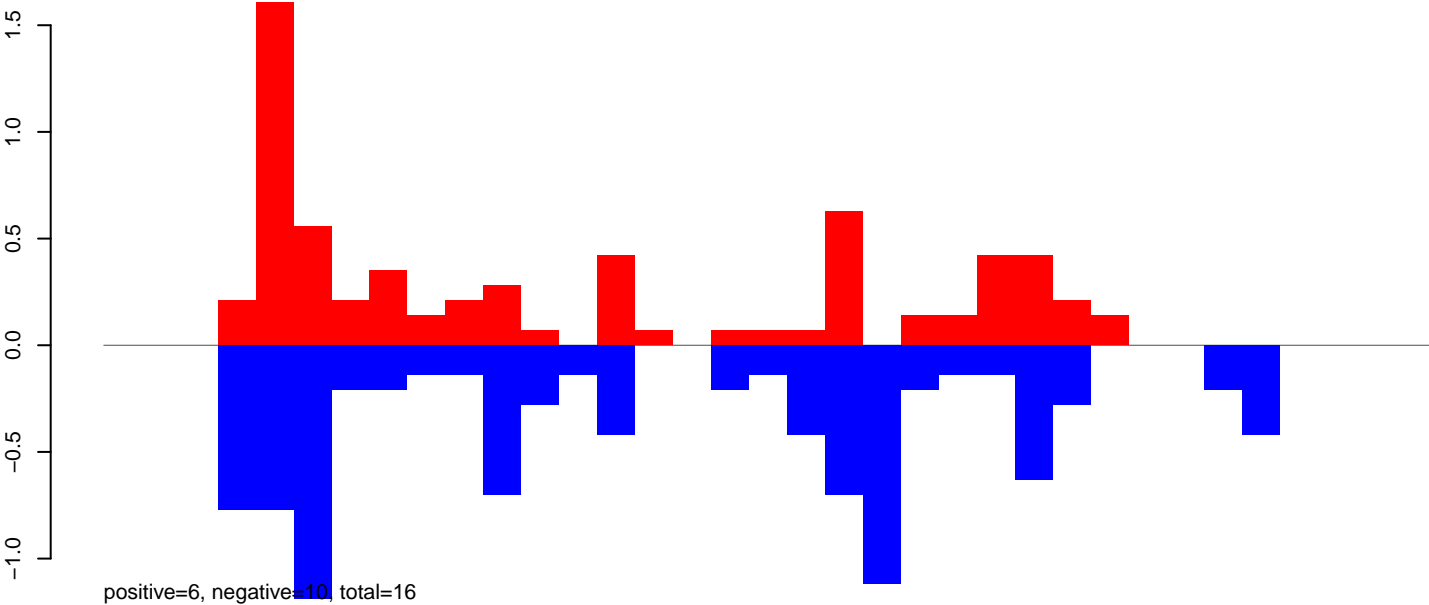


CuQuin_Mcarc_NL.rep

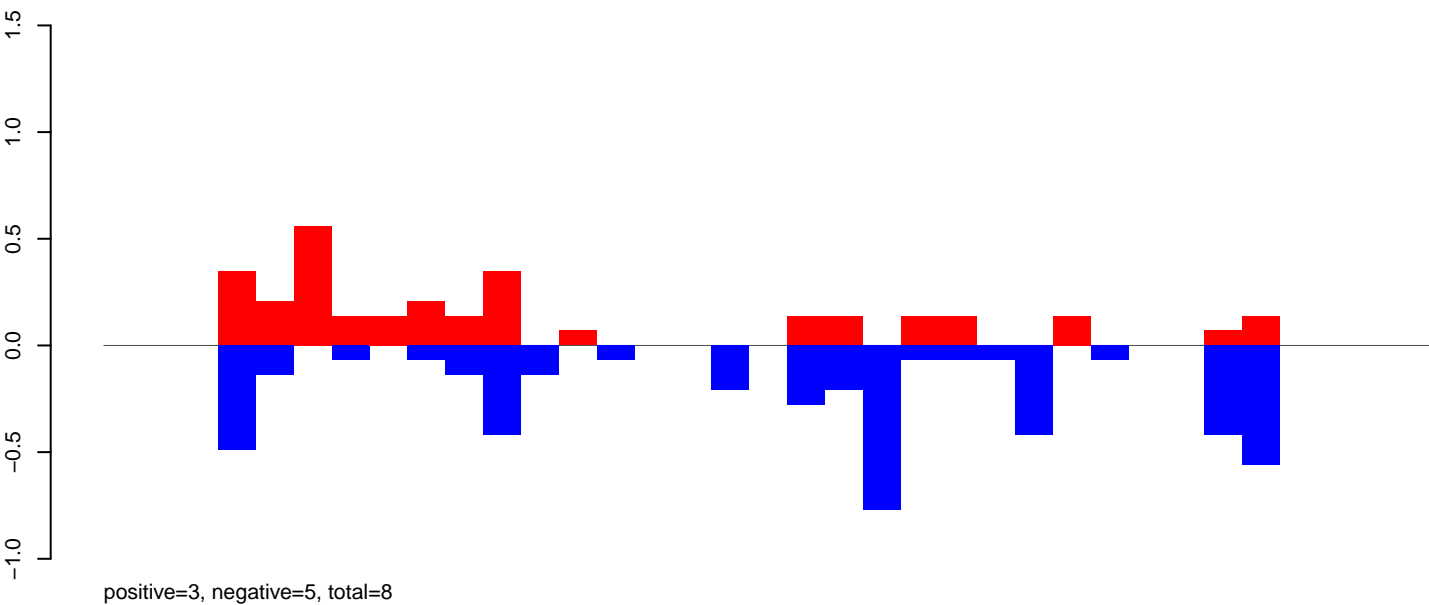


Window size=25, length=984, TE@rnd-2_family-538@DNA@647:1-984

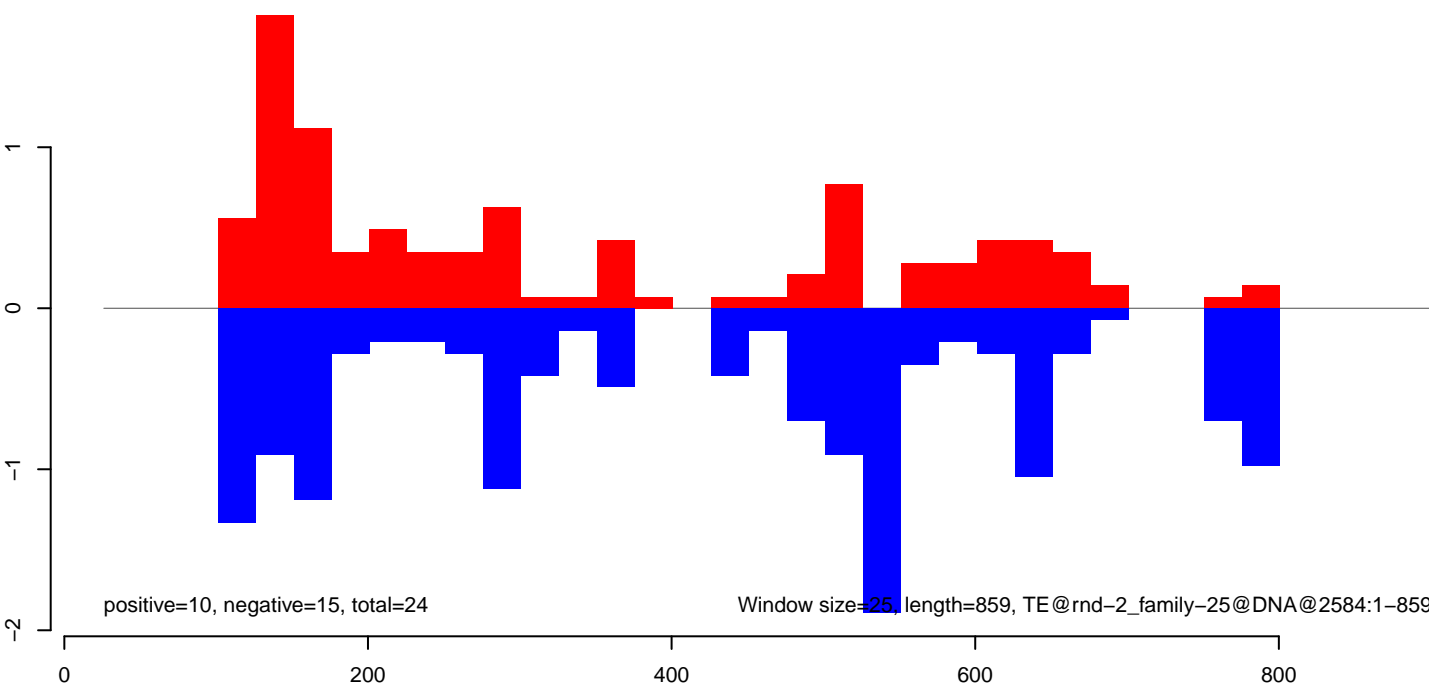
CuQuin_Mcarc_NL.18_23.rep



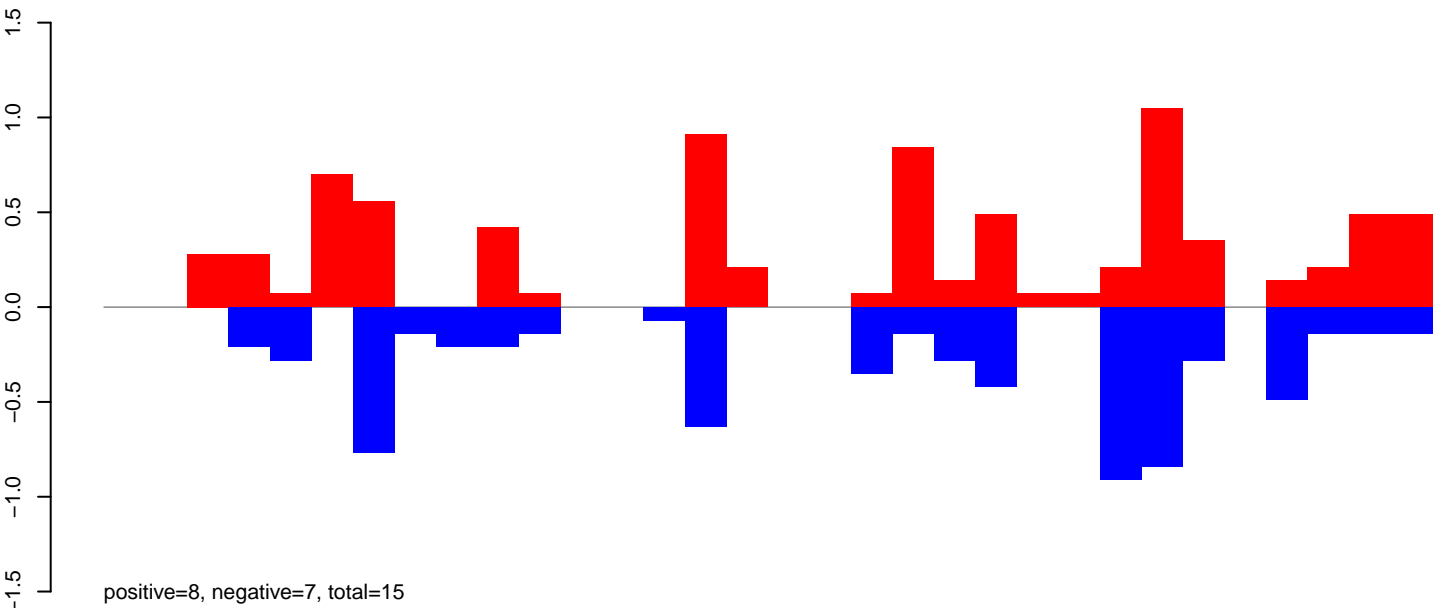
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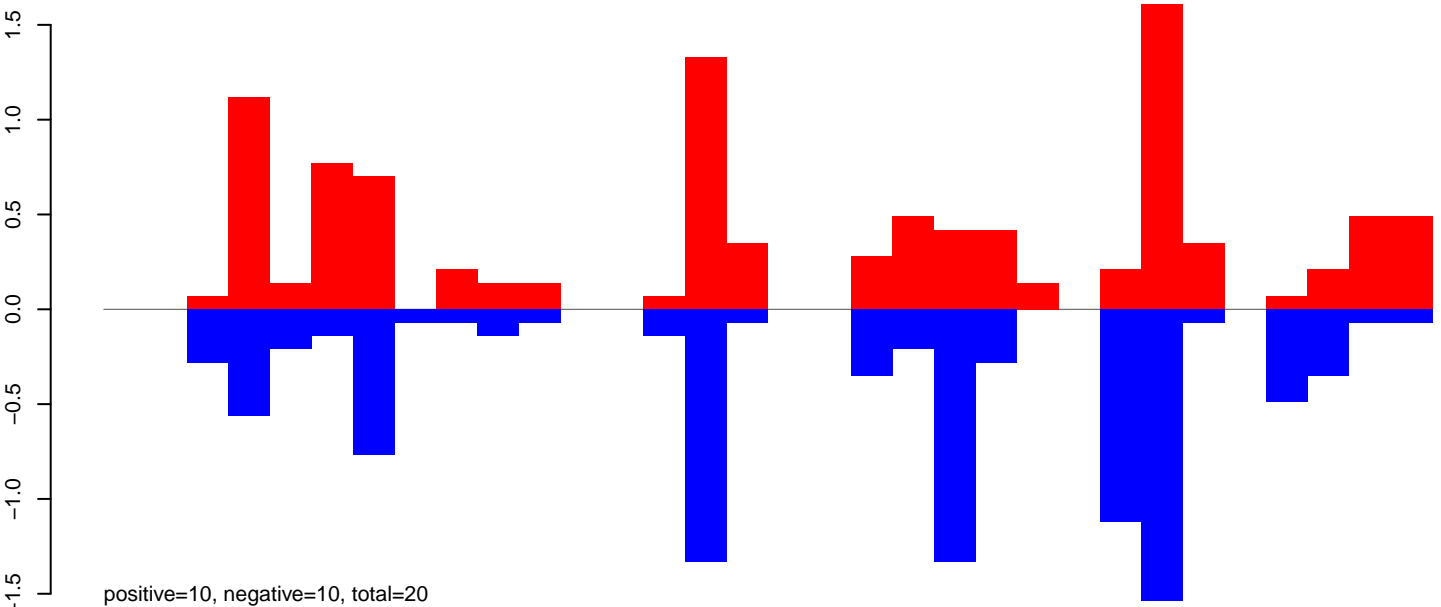
CuQuin_Mcarc_NL.rep



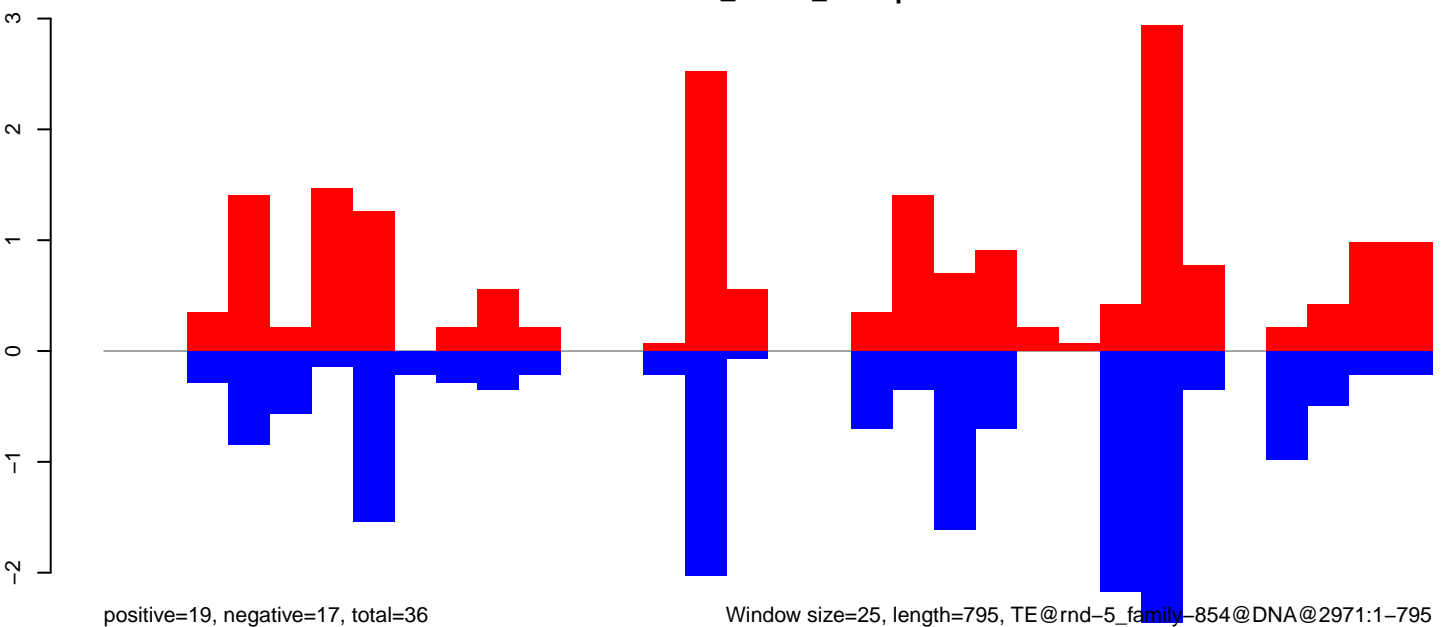
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CuQuin_Mcarc_NL.24_35.rep



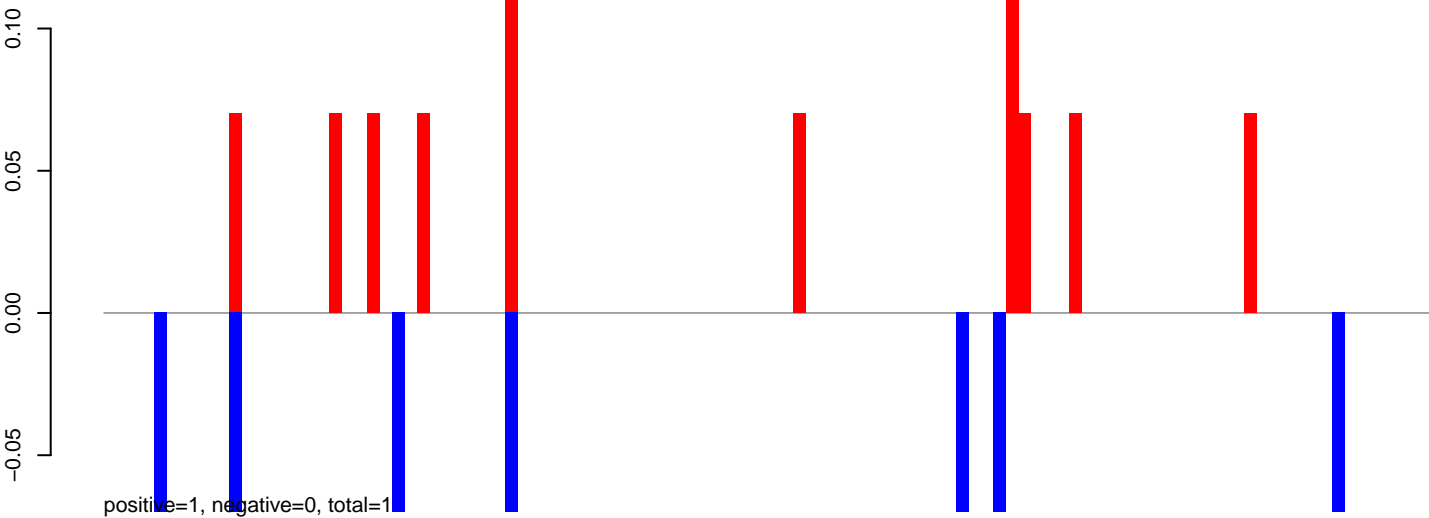
CuQuin_Mcarc_NL.rep



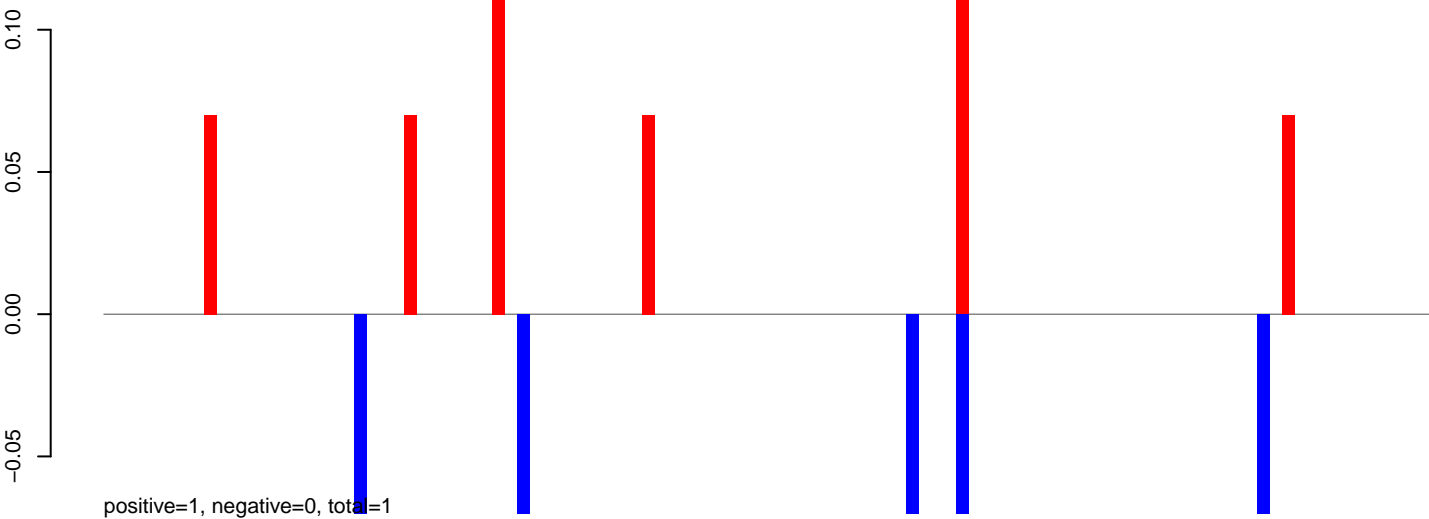
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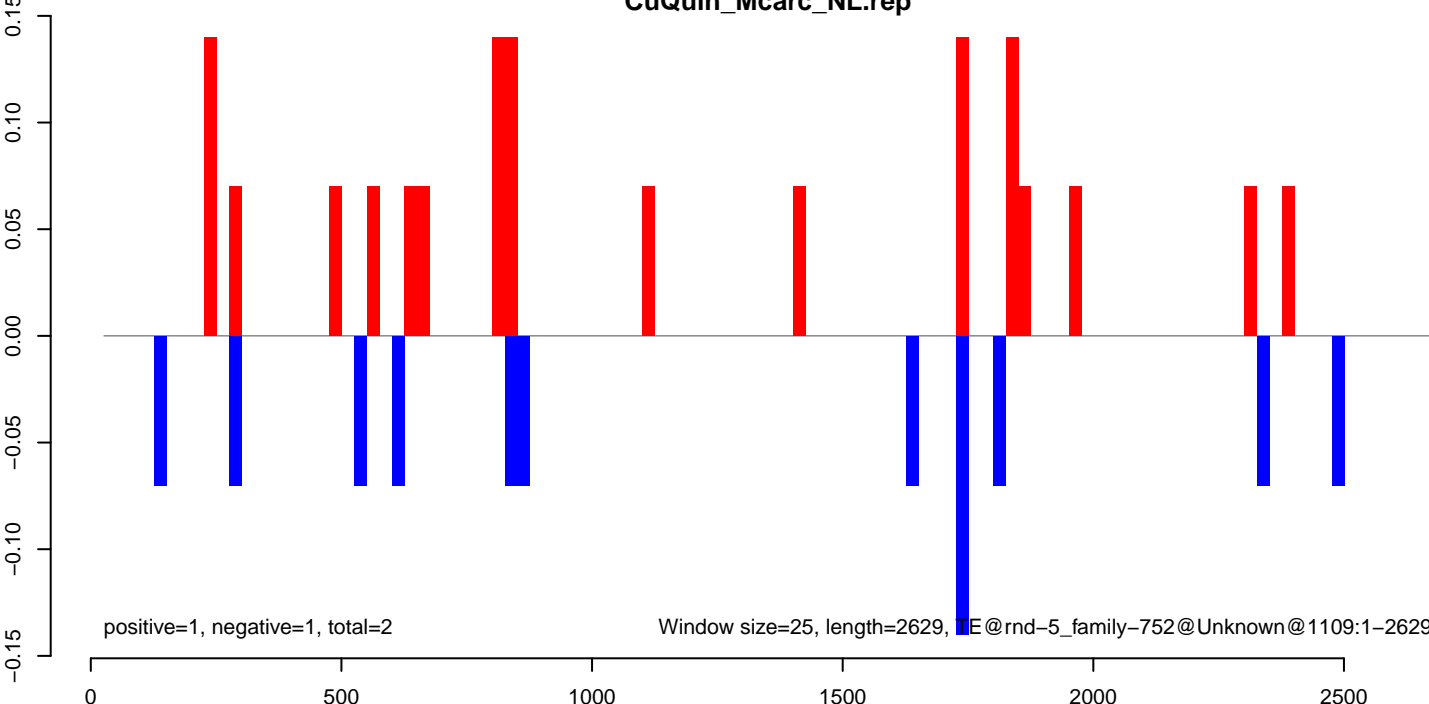
CuQuin_Mcarc_NL.18_23.rep



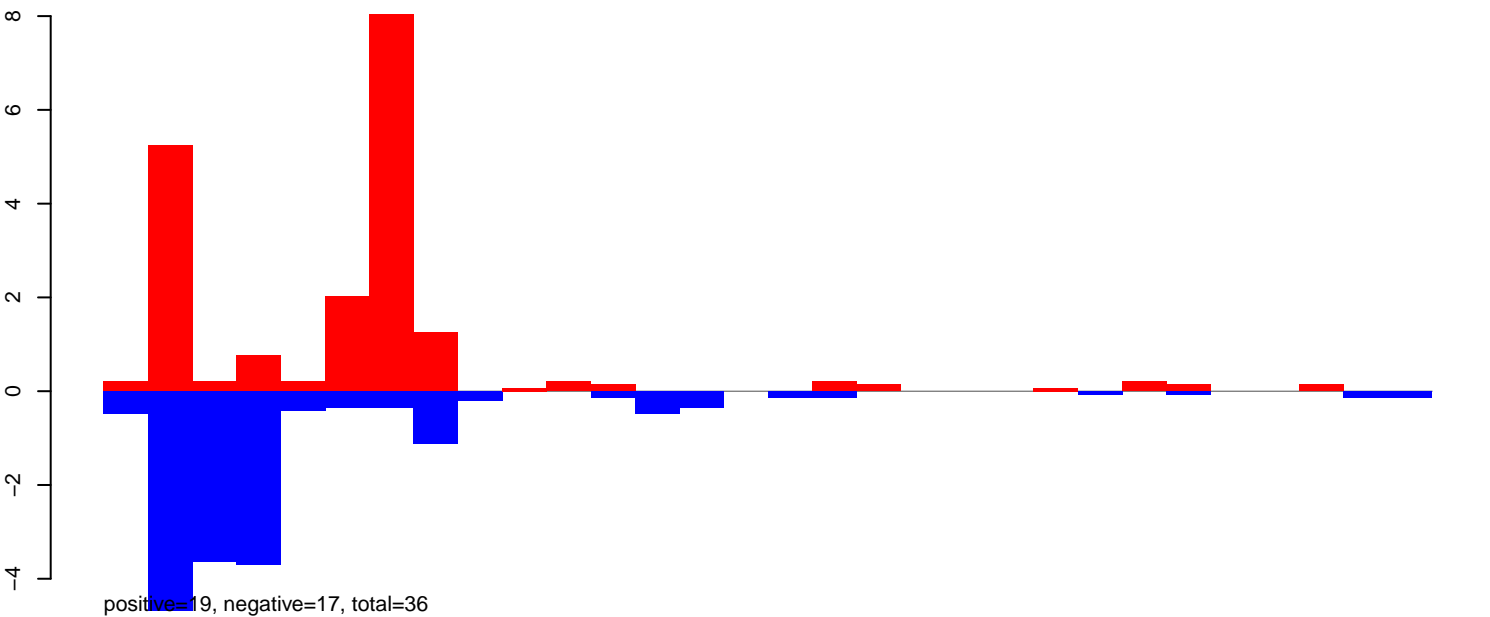
CuQuin_Mcarc_NL.24_35.rep



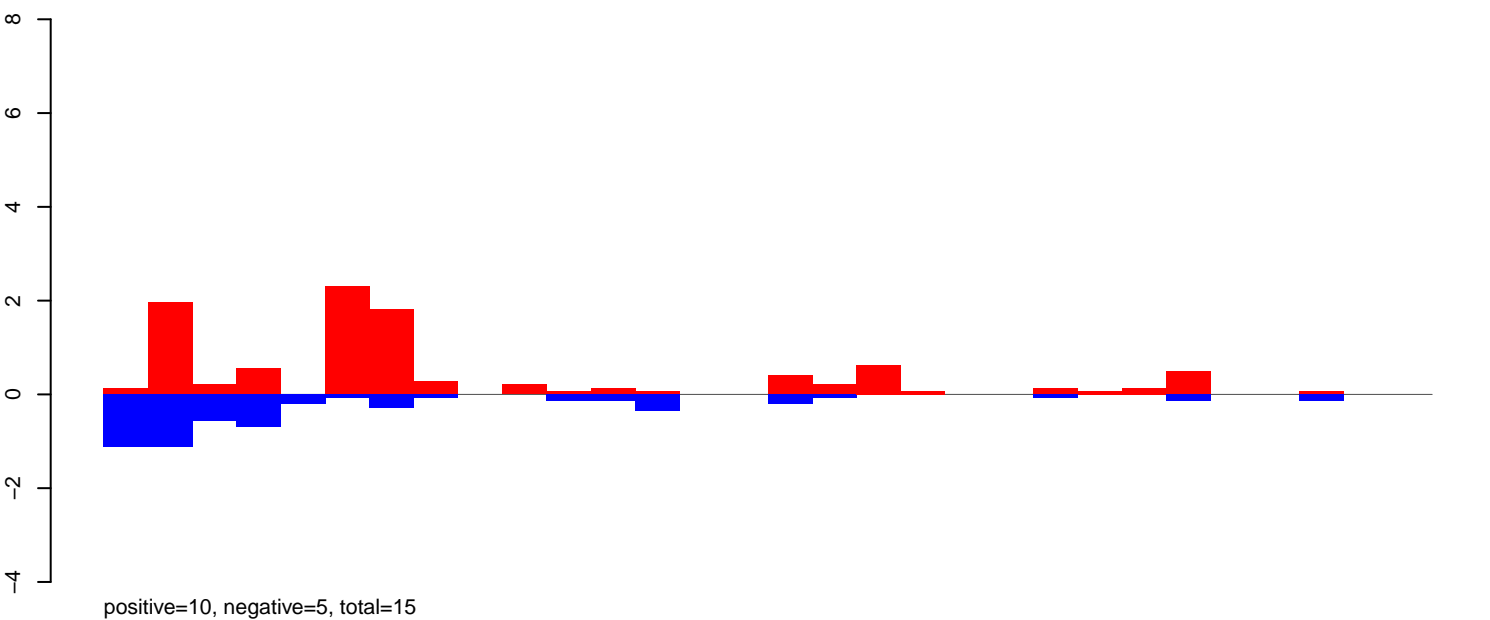
CuQuin_Mcarc_NL.rep



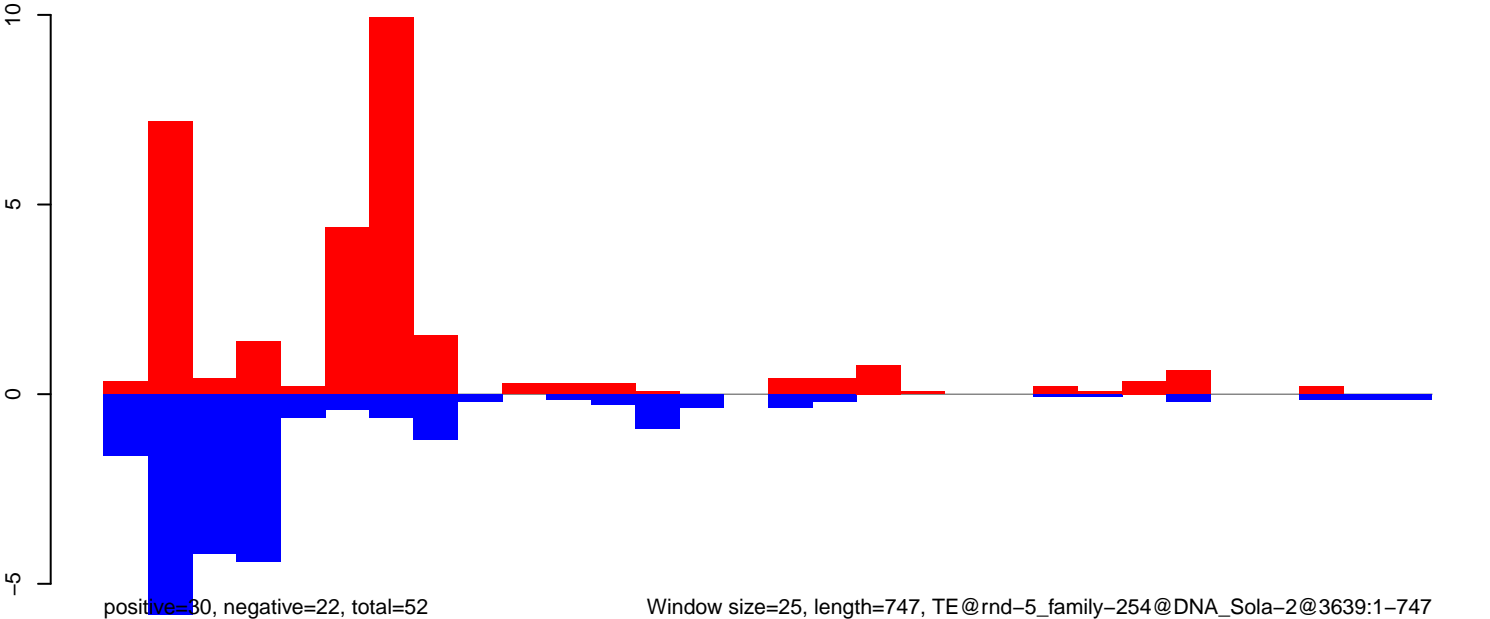
CuQuin_Mcarc_NL.18_23.rep



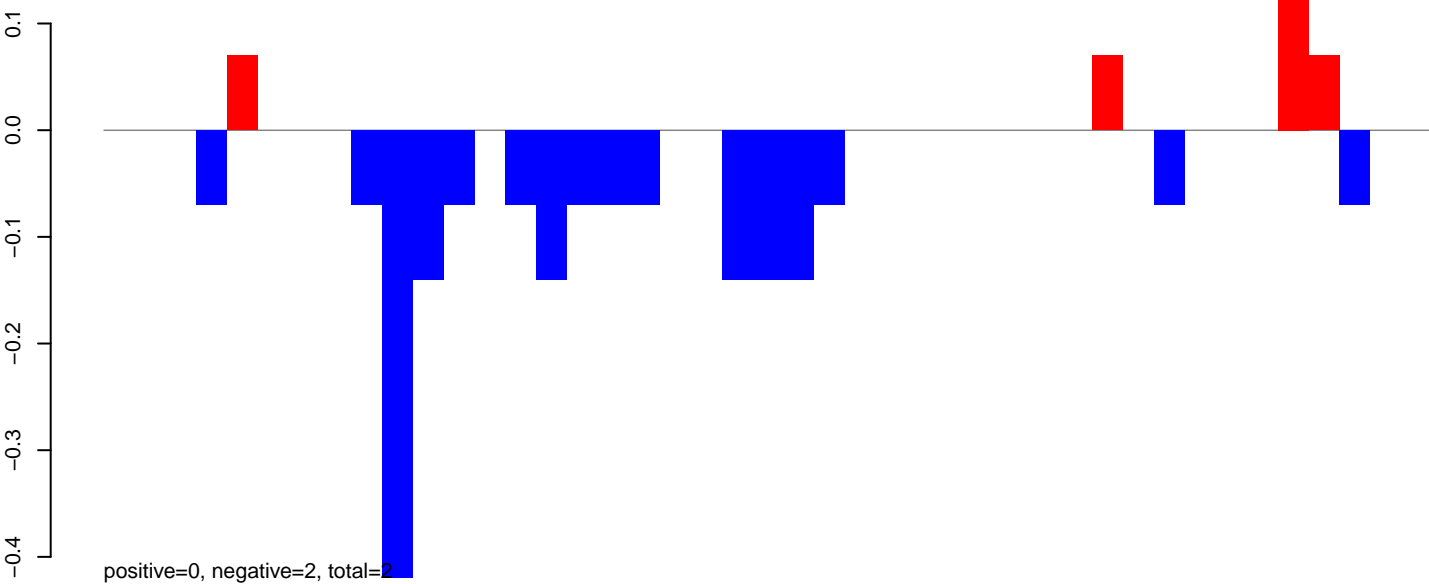
CuQuin_Mcarc_NL.24_35.rep



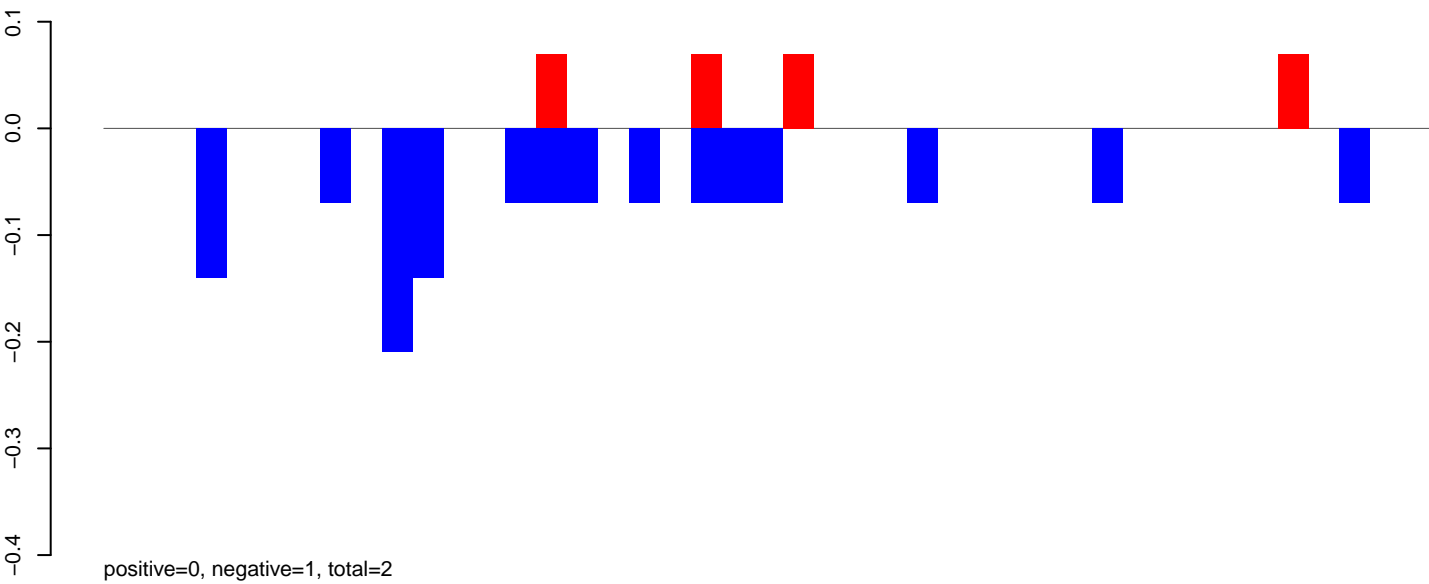
CuQuin_Mcarc_NL.rep



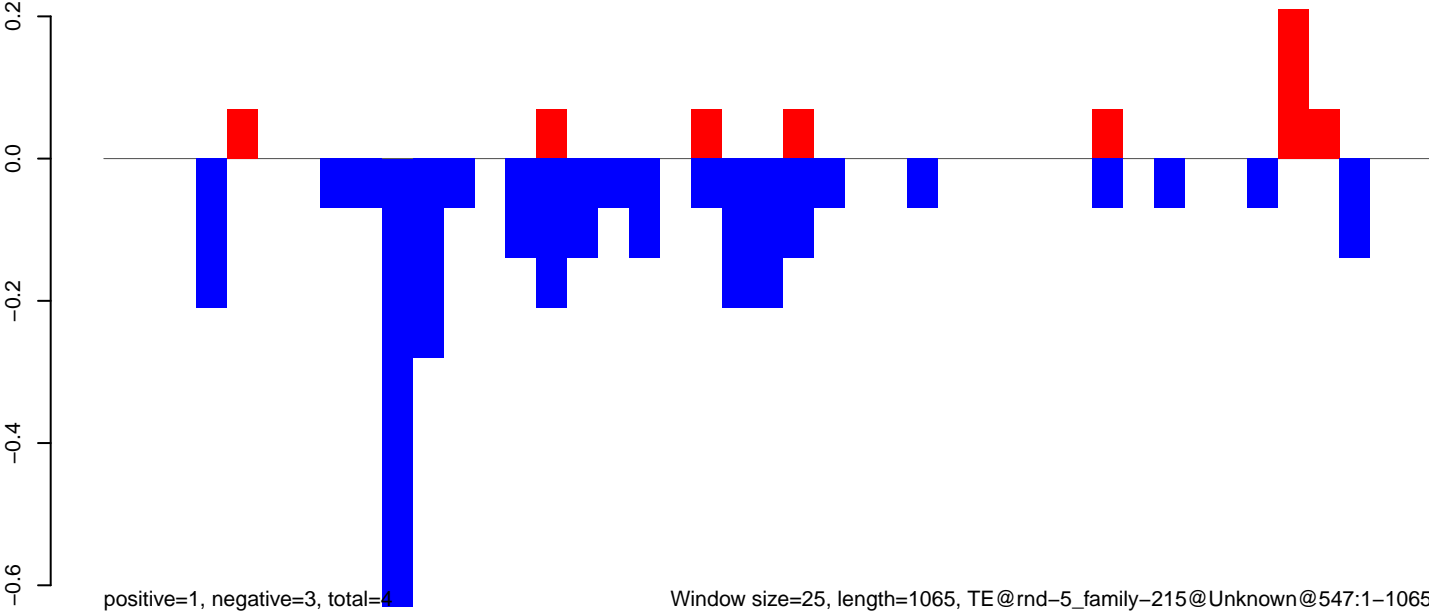
CuQuin_Mcarc_NL.18_23.rep



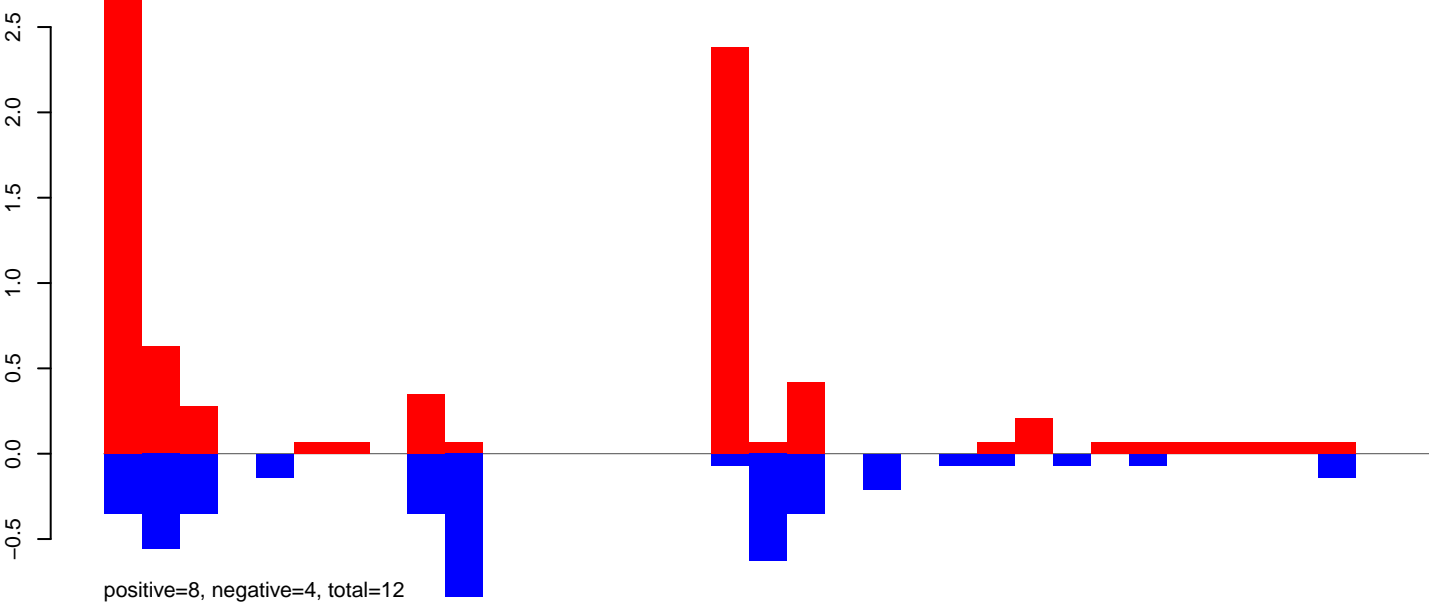
CuQuin_Mcarc_NL.24_35.rep



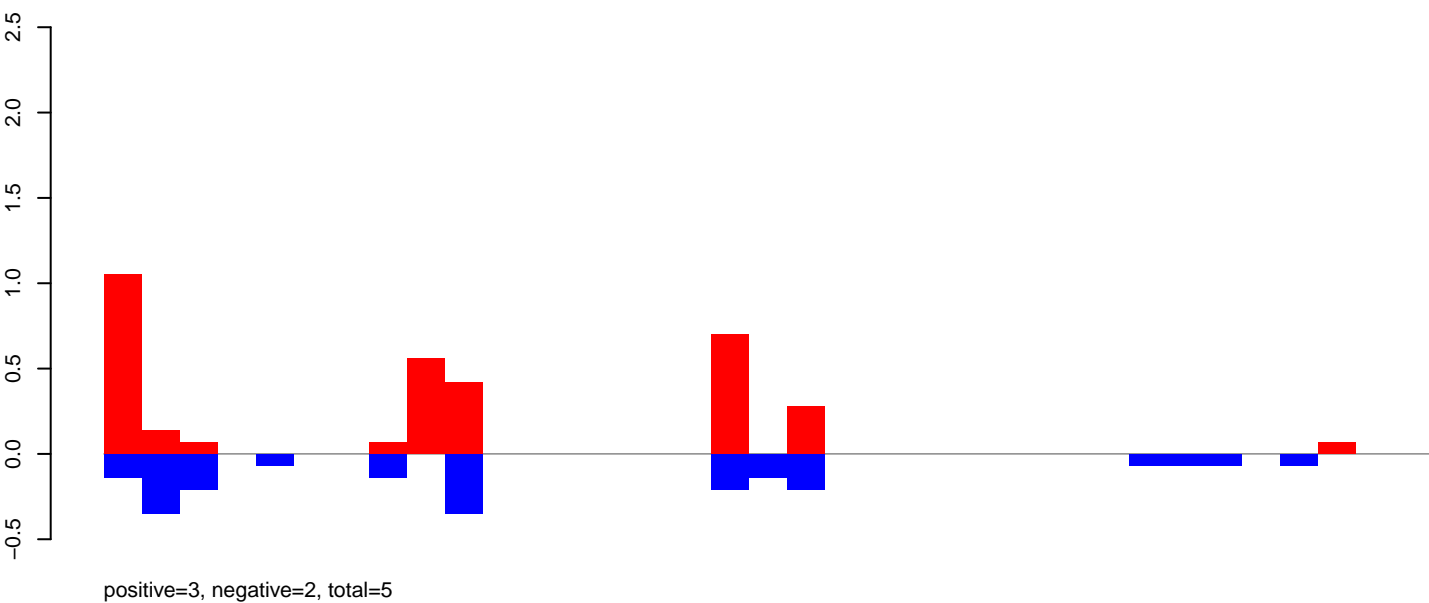
CuQuin_Mcarc_NL.rep



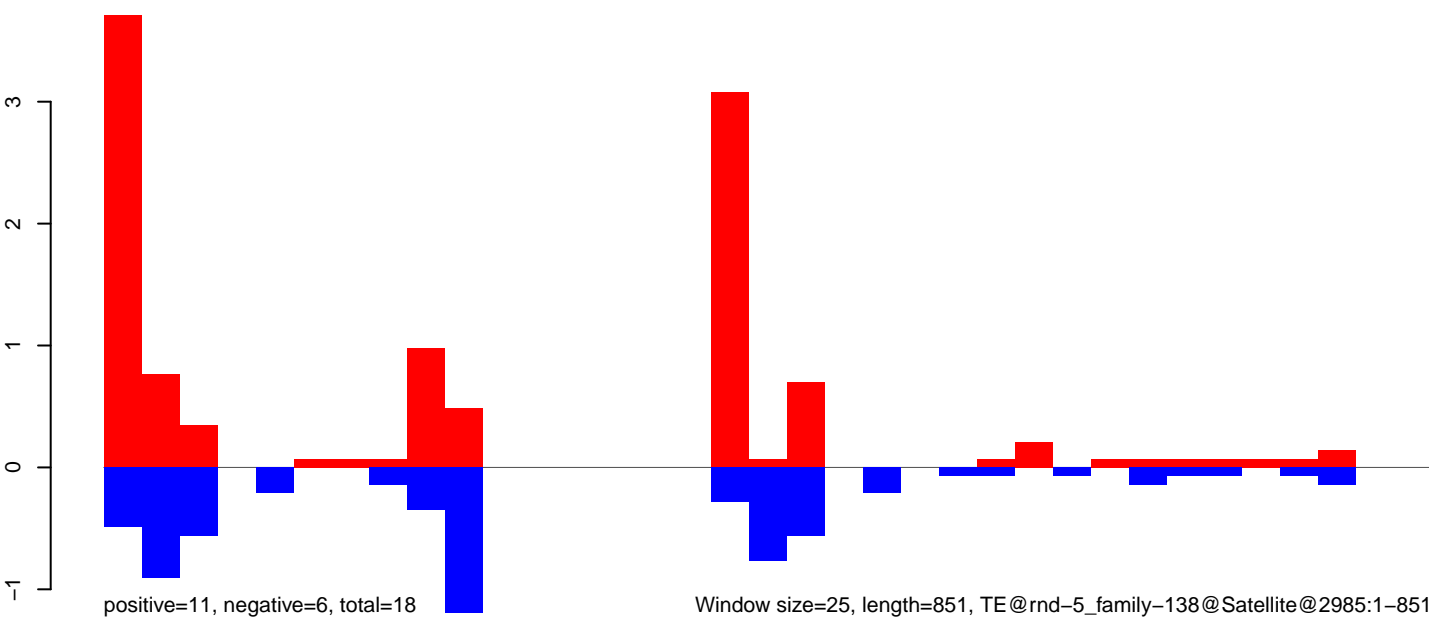
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



0

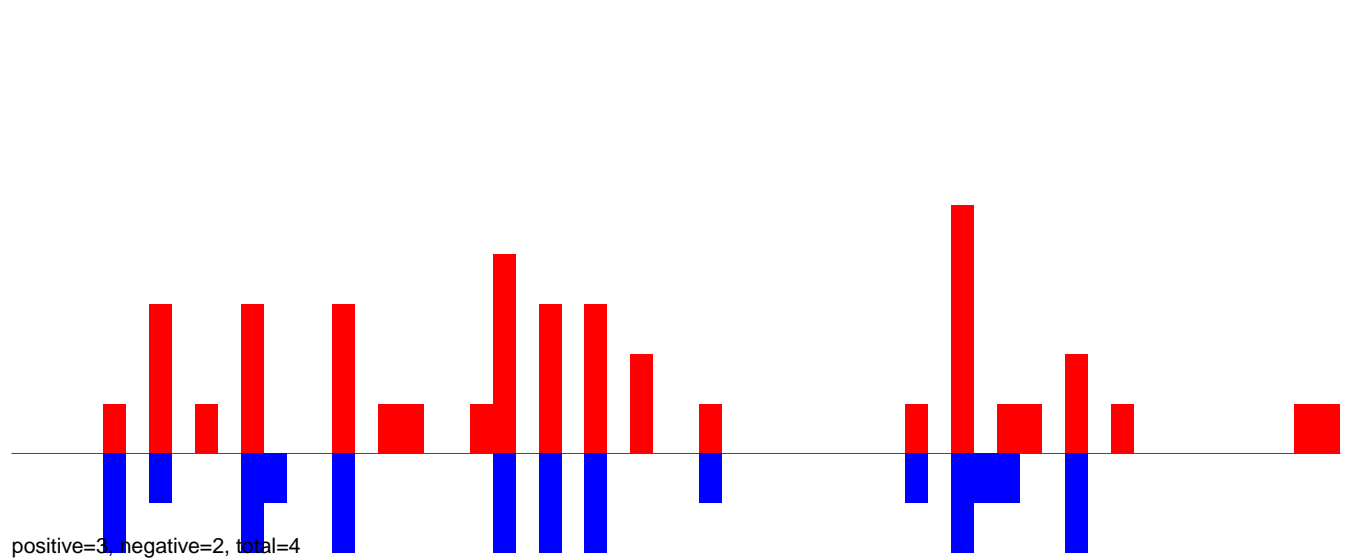
200

400

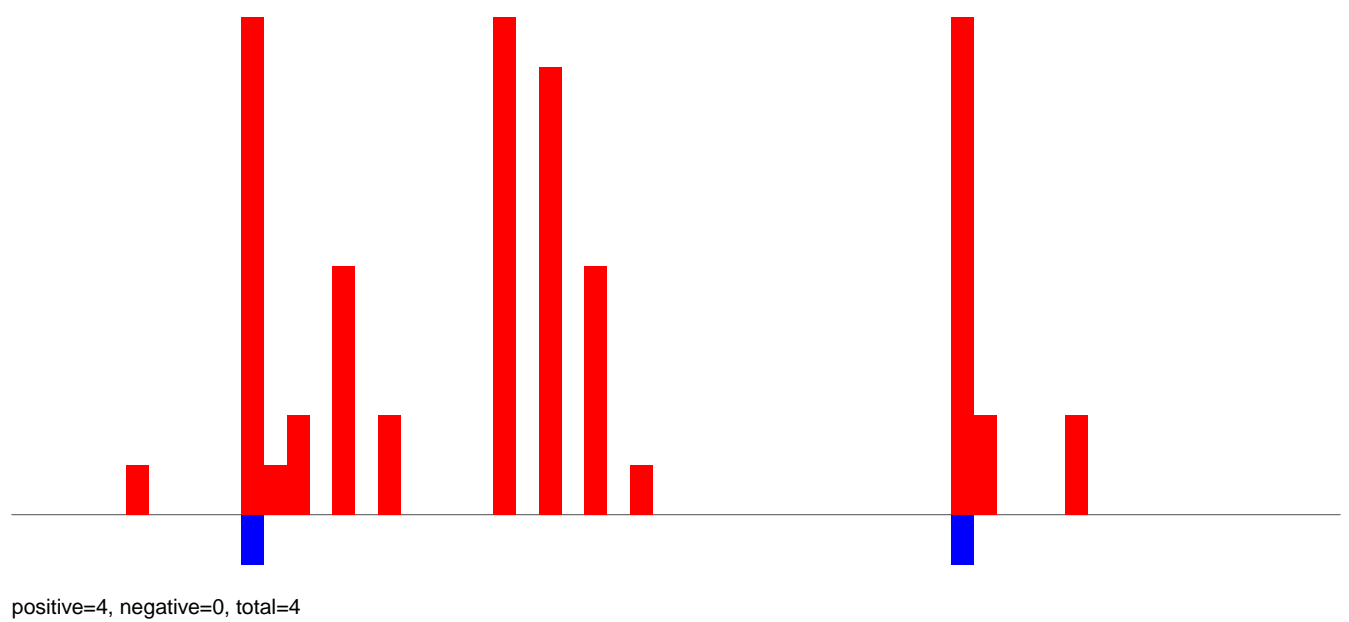
600

800

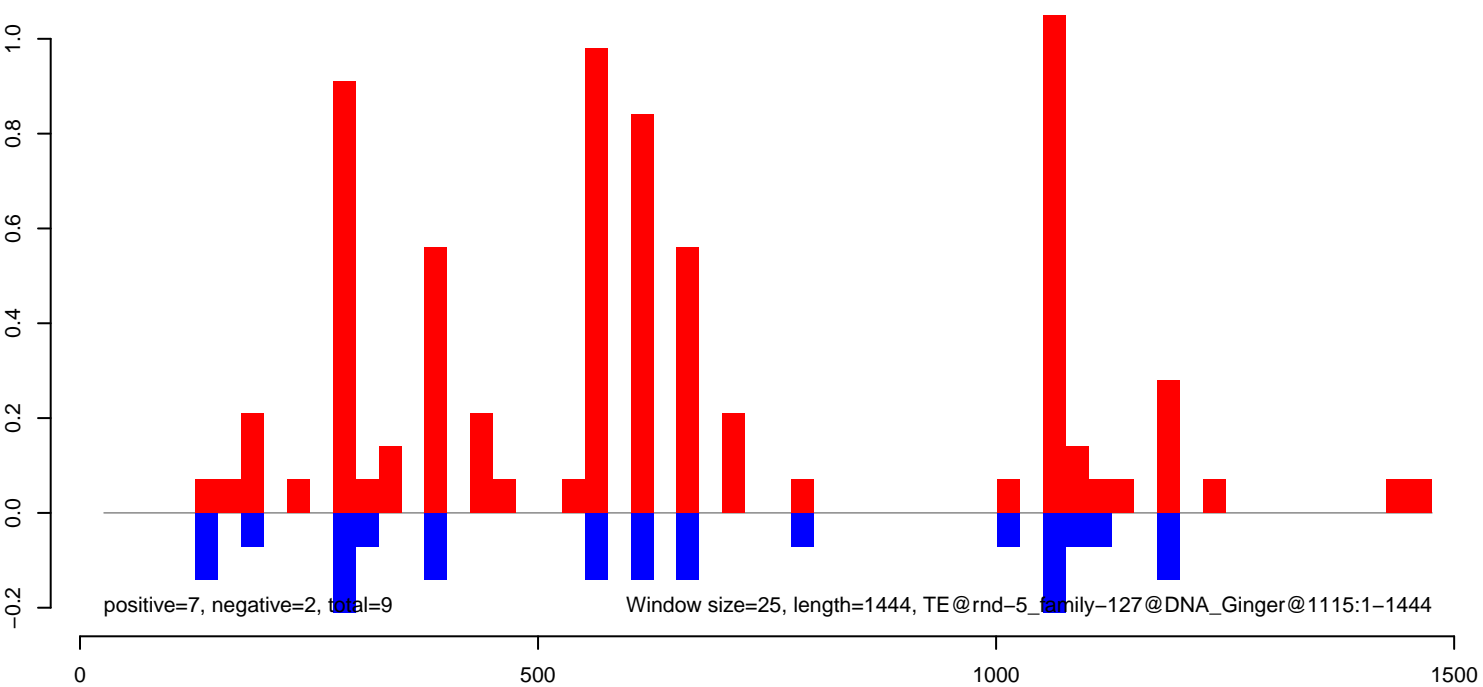
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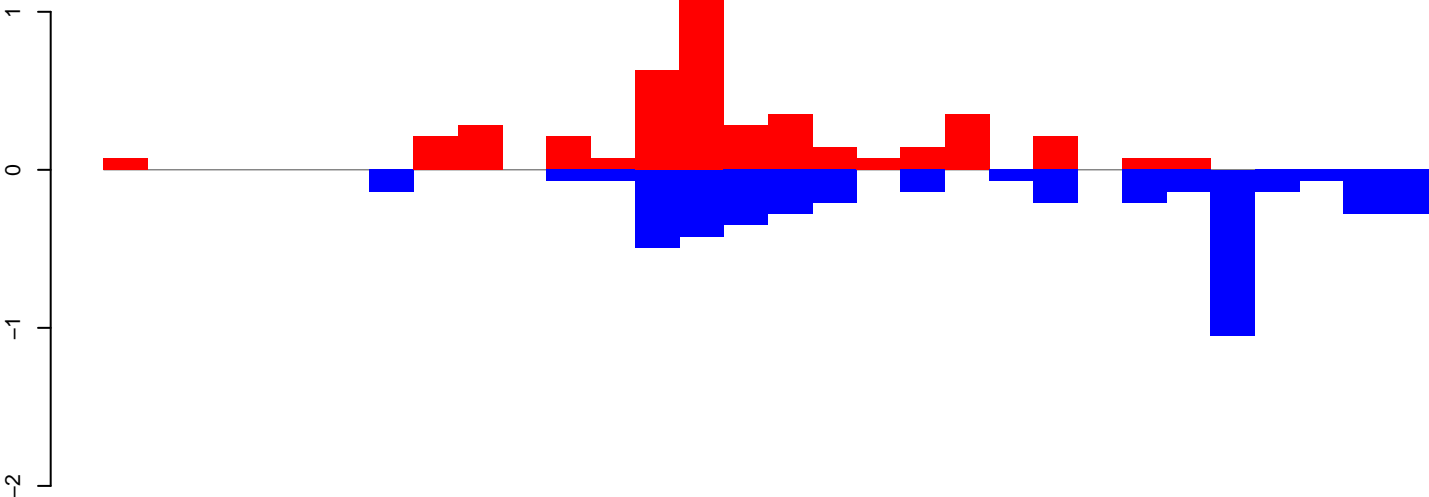
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

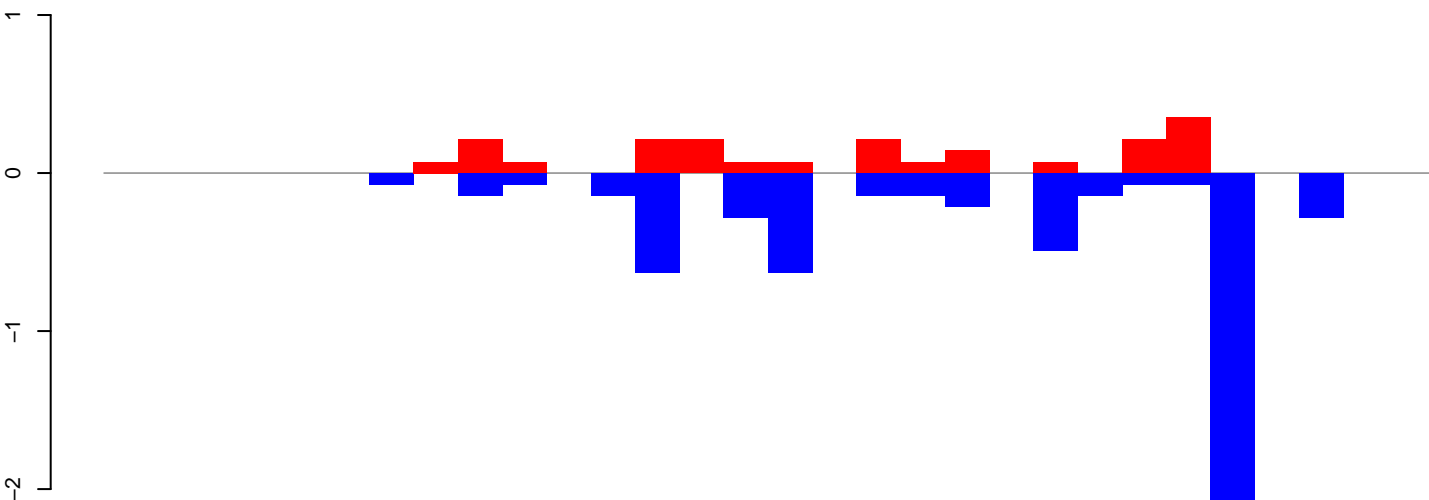


CuQuin_Mcarc_NL.18_23.rep



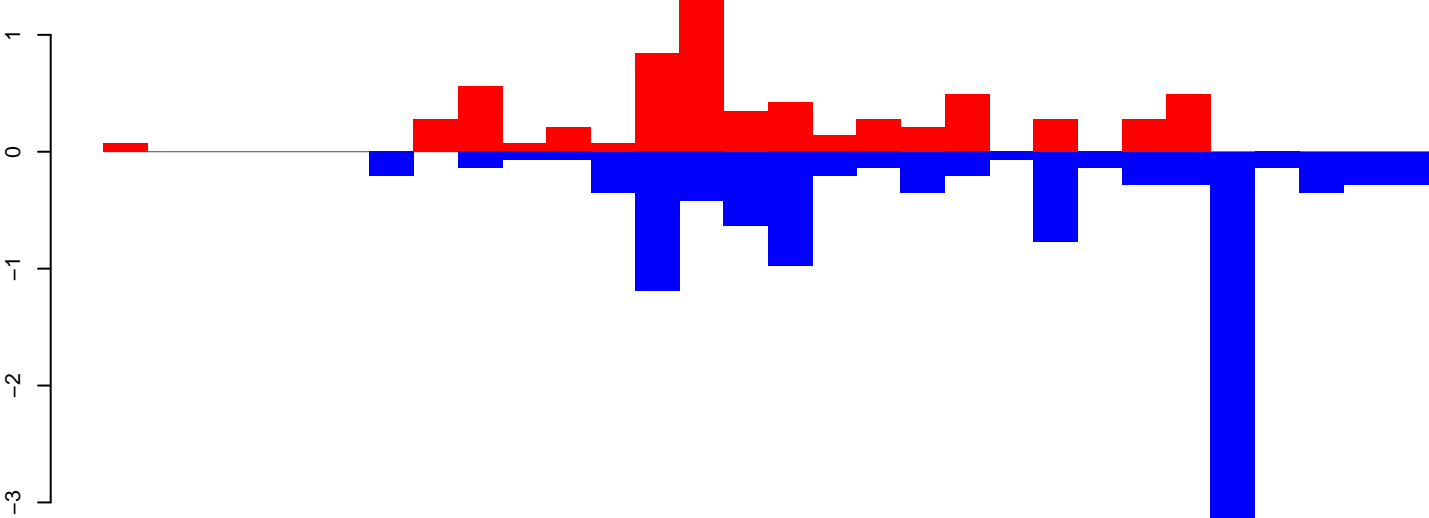
positive=4, negative=5, total=9

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=6, total=8

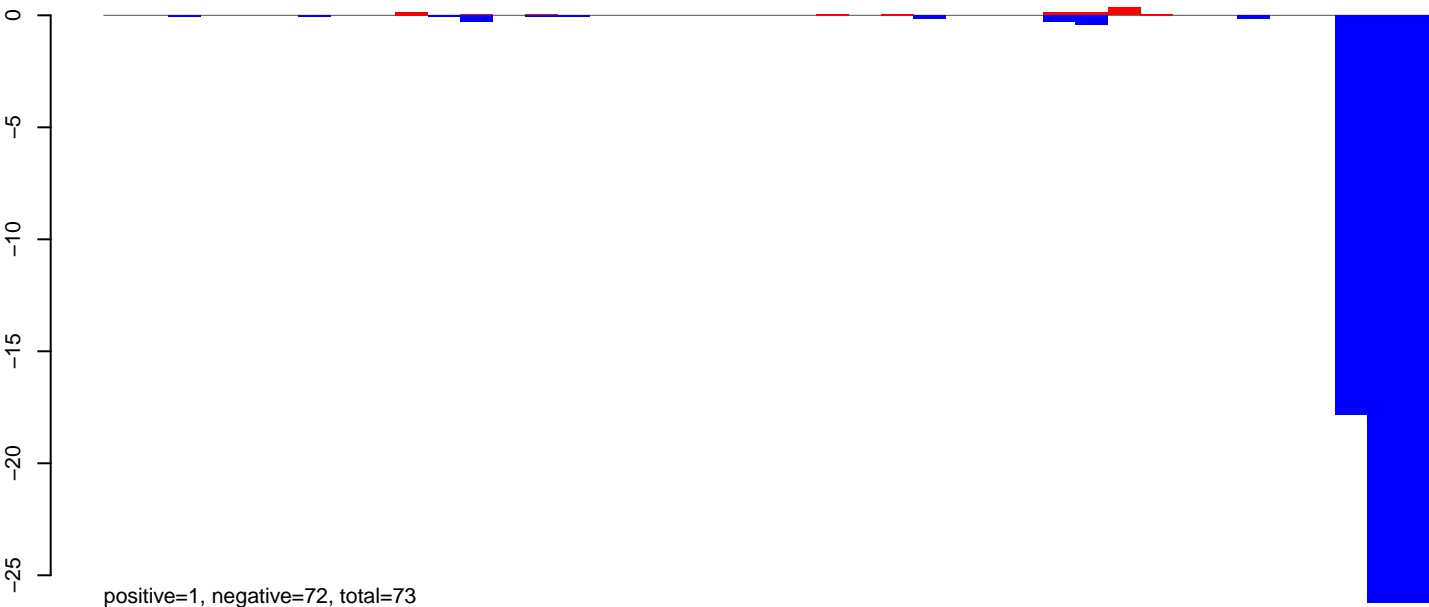
CuQuin_Mcarc_NL.rep



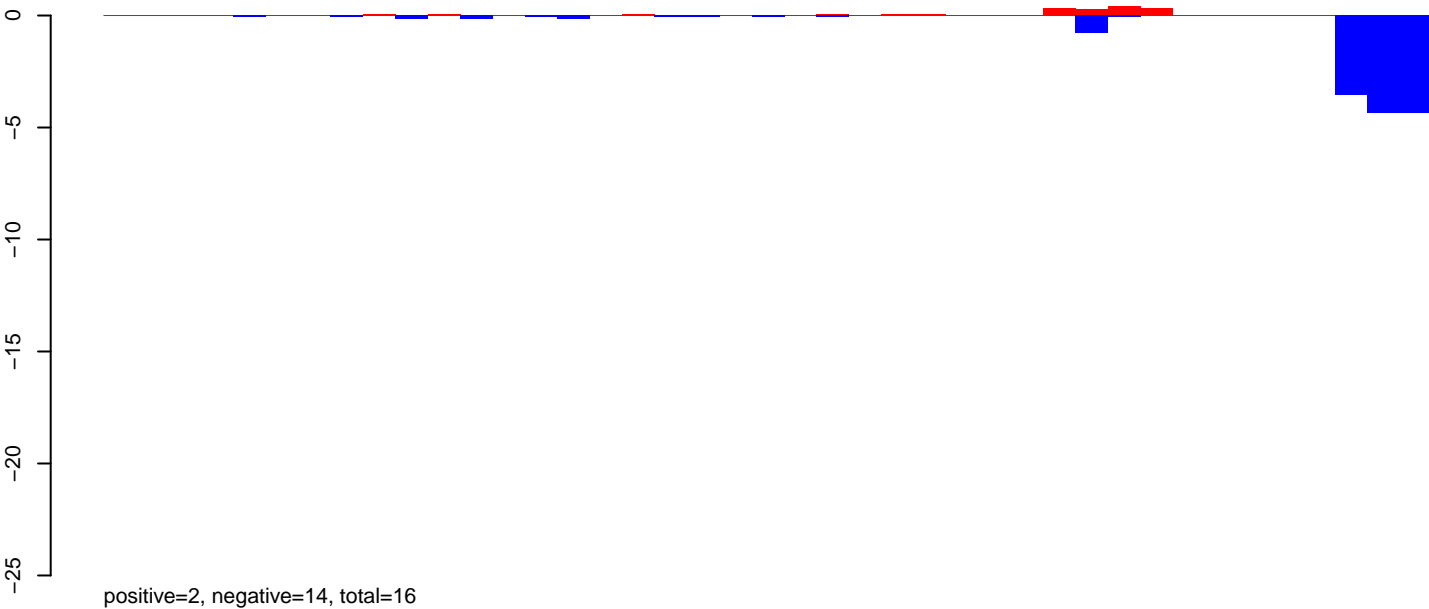
positive=6, negative=11, total=18

Window size=25, length=729, TE@rnd-4_family-6@Unknown@2060:1-729

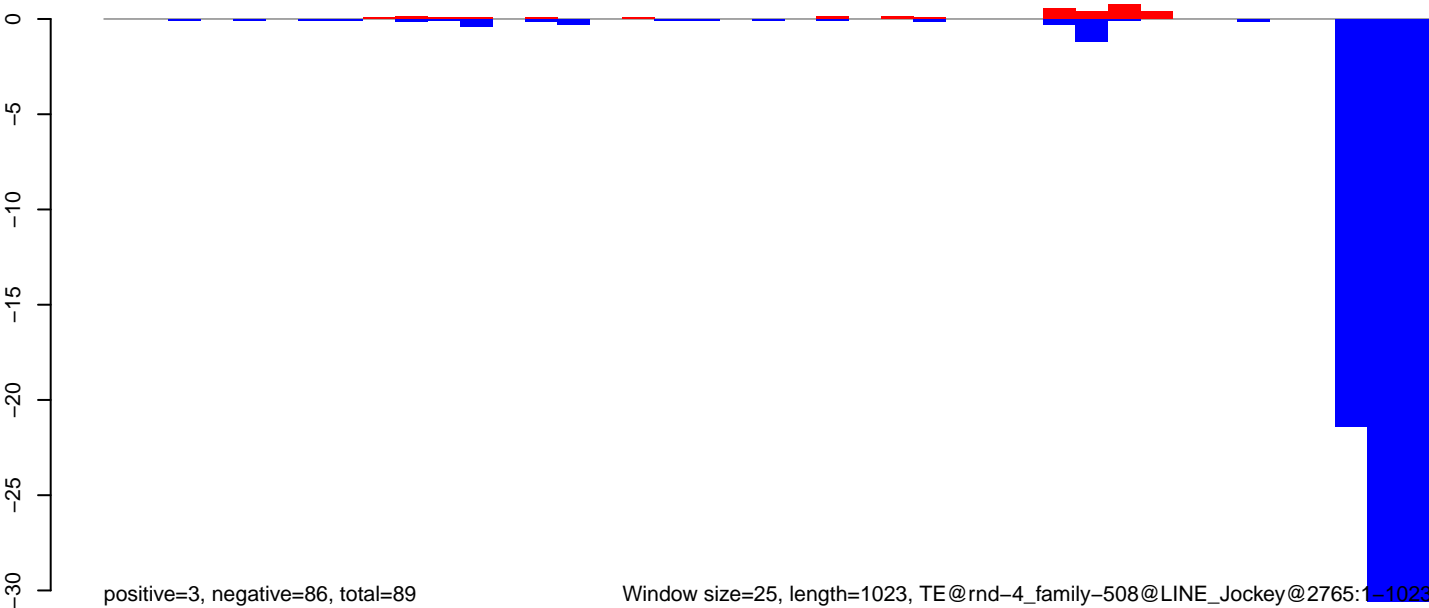
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

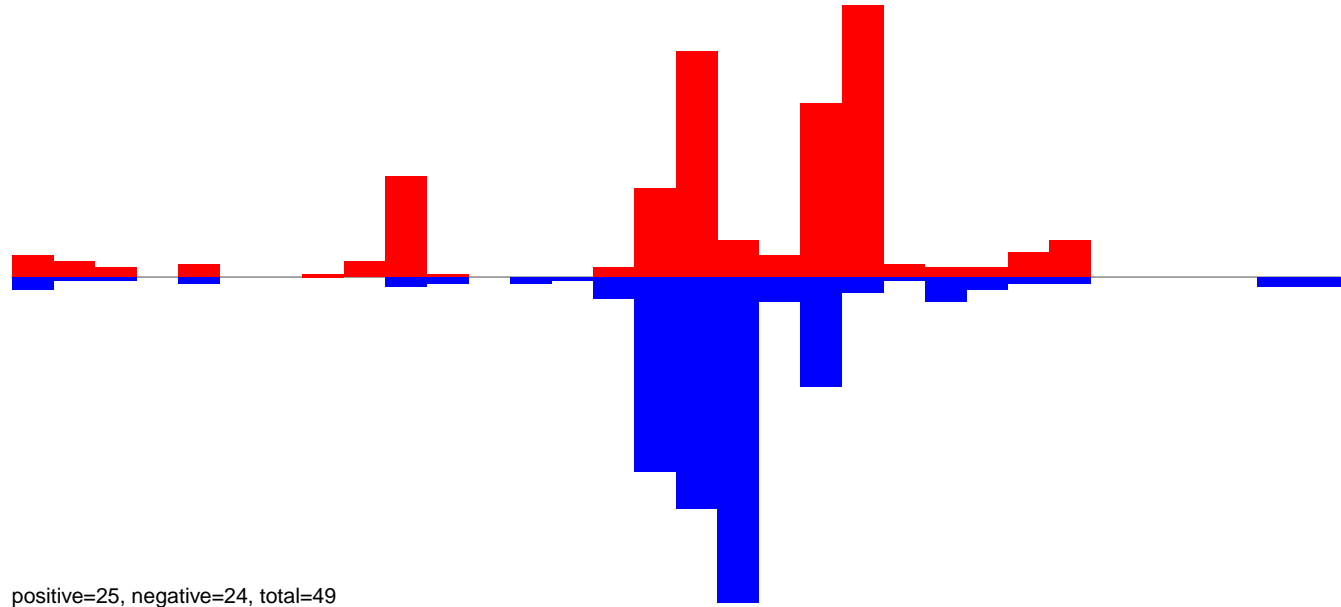


CuQuin_Mcarc_NL.rep

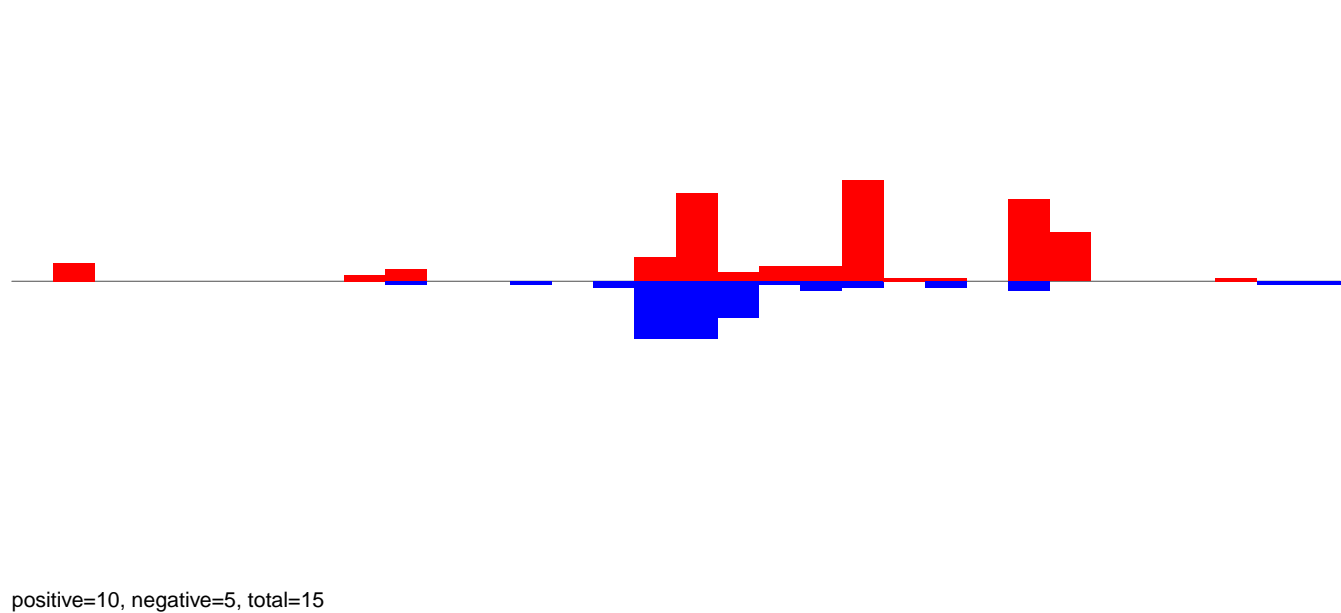


Window size=25, length=1023, TE@rnd-4_family-508@LINE_Jockey@2765:1-1023

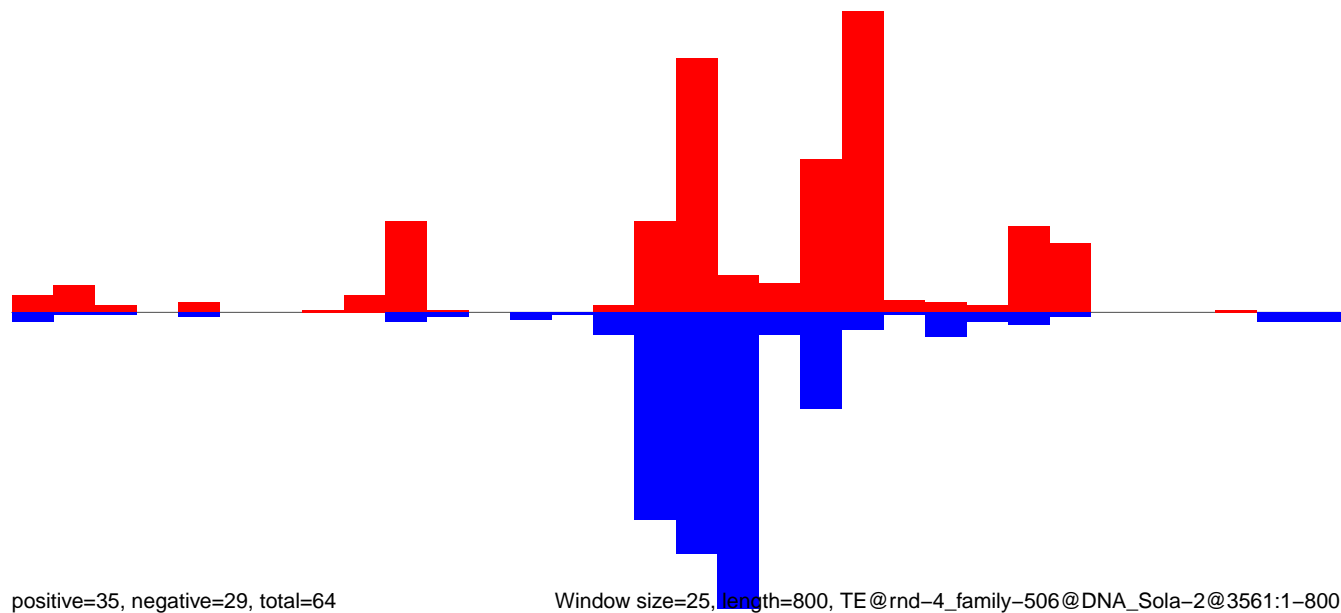
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

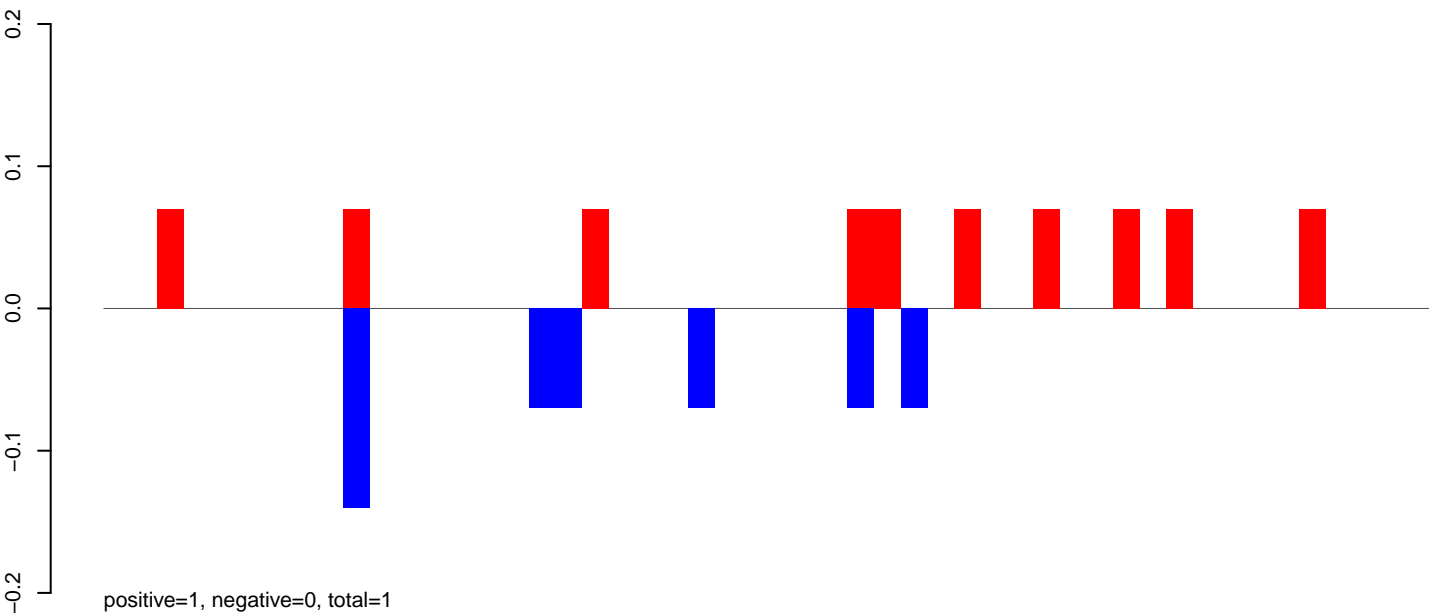


CuQuin_Mcarc_NL.rep

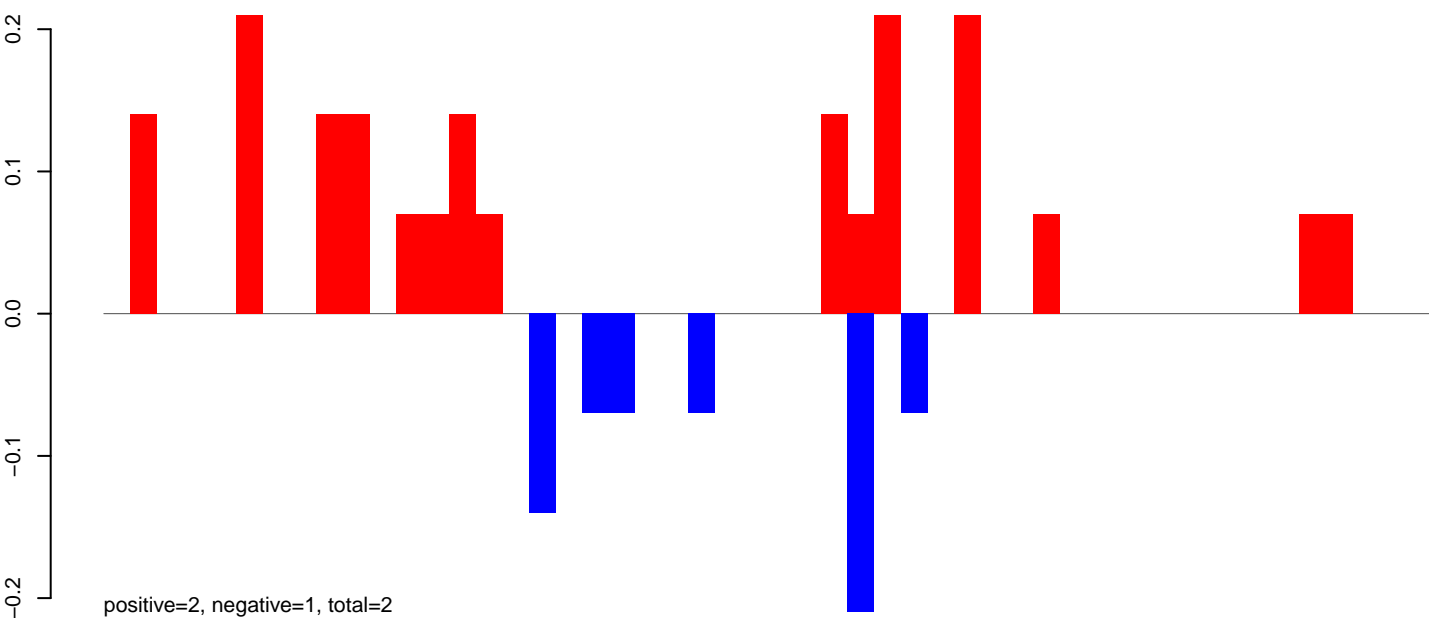


Window size=25, length=800, TE@rnd-4_family-506@DNA_Sola-2@3561:1-800

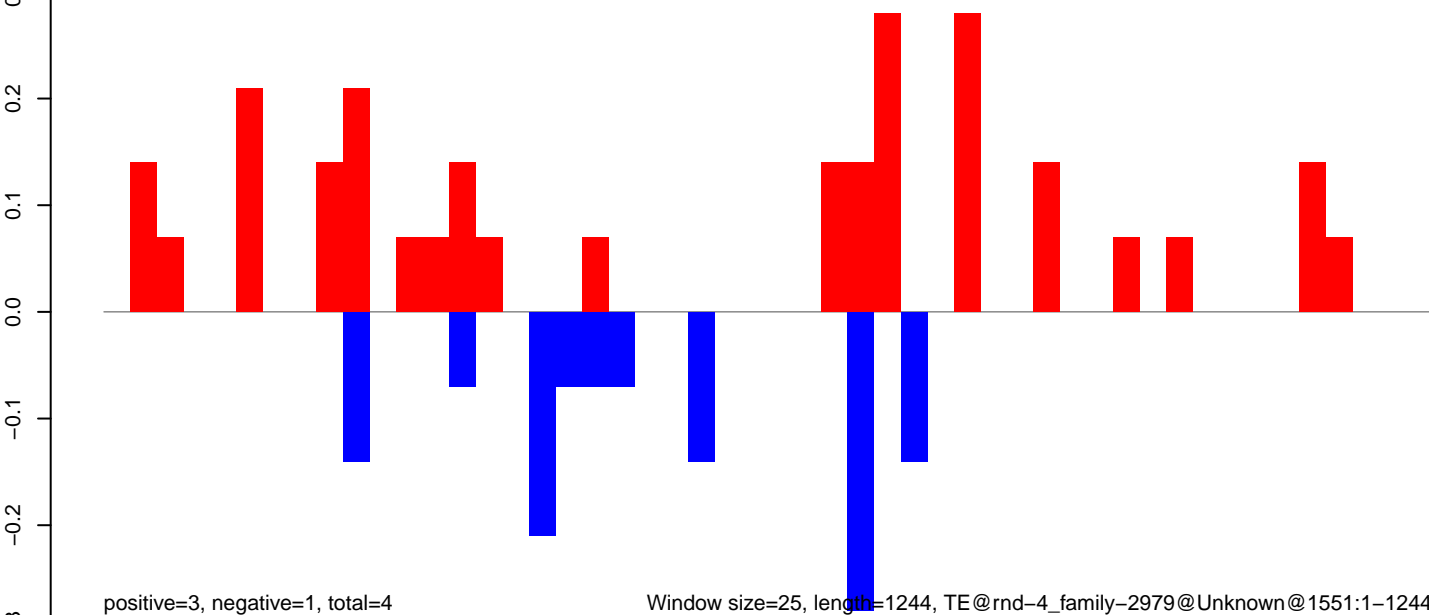
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

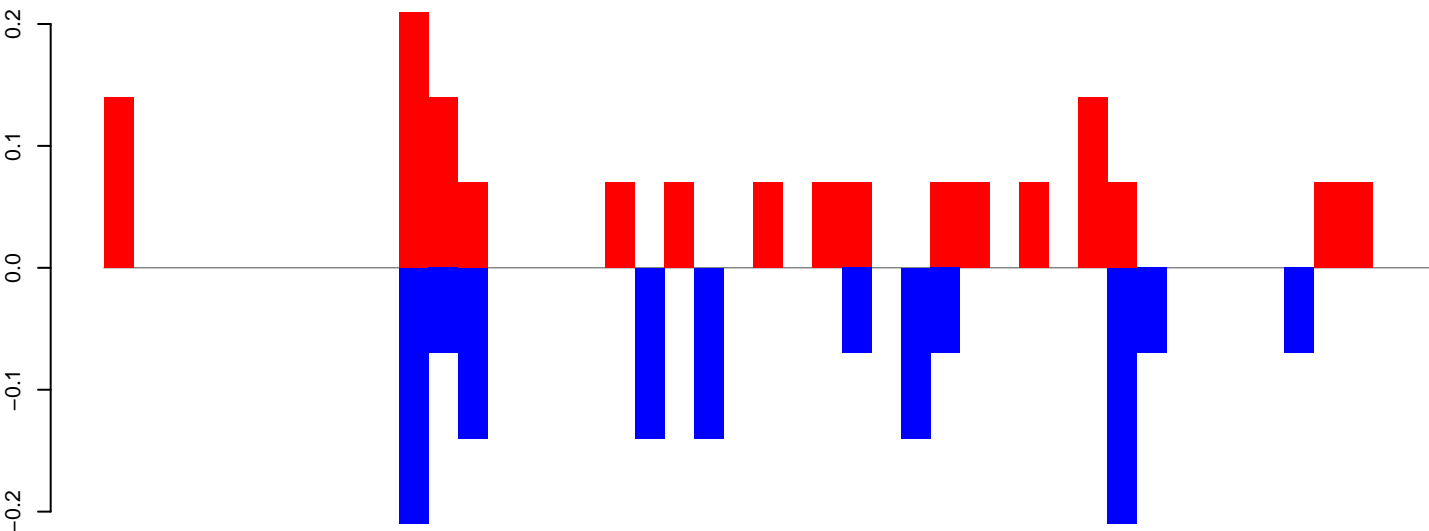


CuQuin_Mcarc_NL.rep



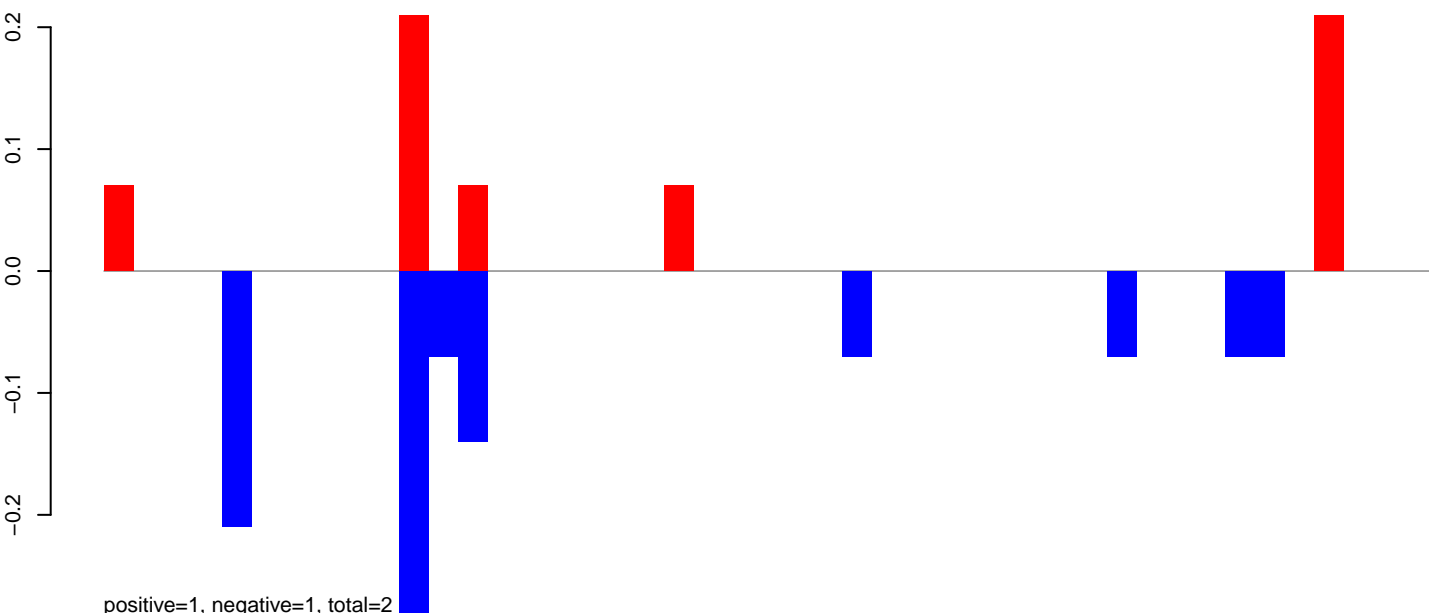
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CuQuin_Mcarc_NL.18_23.rep



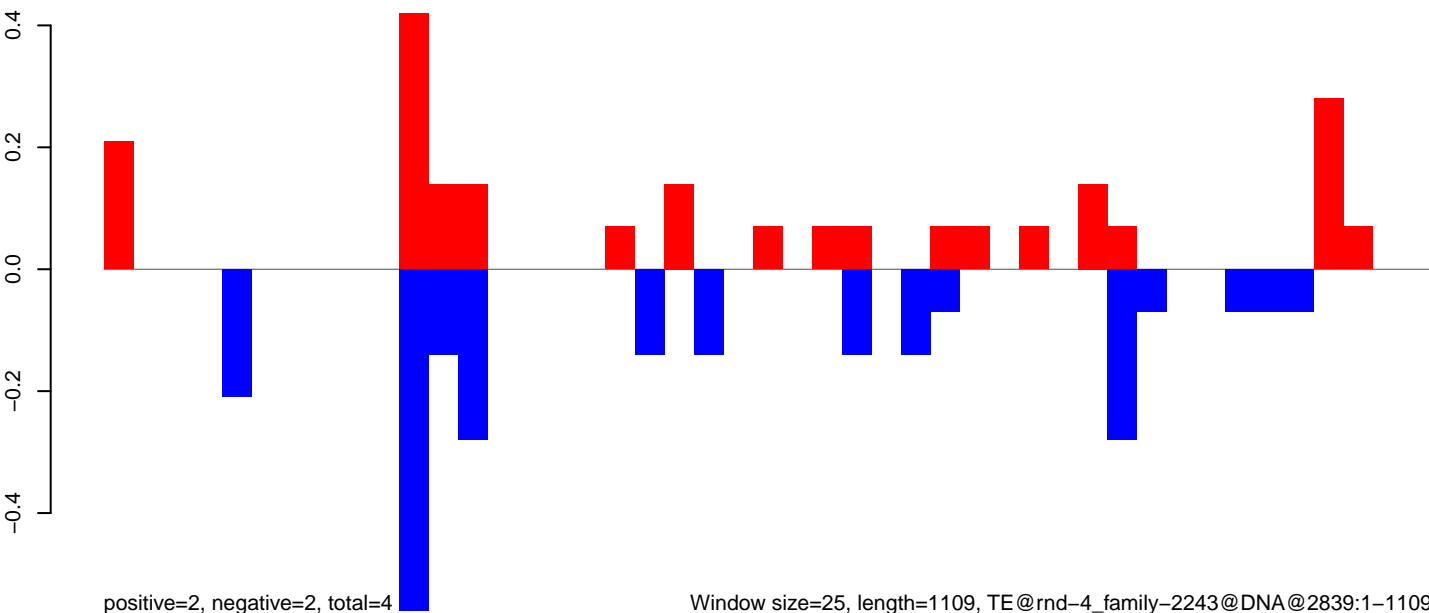
positive=1, negative=1, total=3

CuQuin_Mcarc_NL.24_35.rep



positive=1, negative=1, total=2

CuQuin_Mcarc_NL.rep

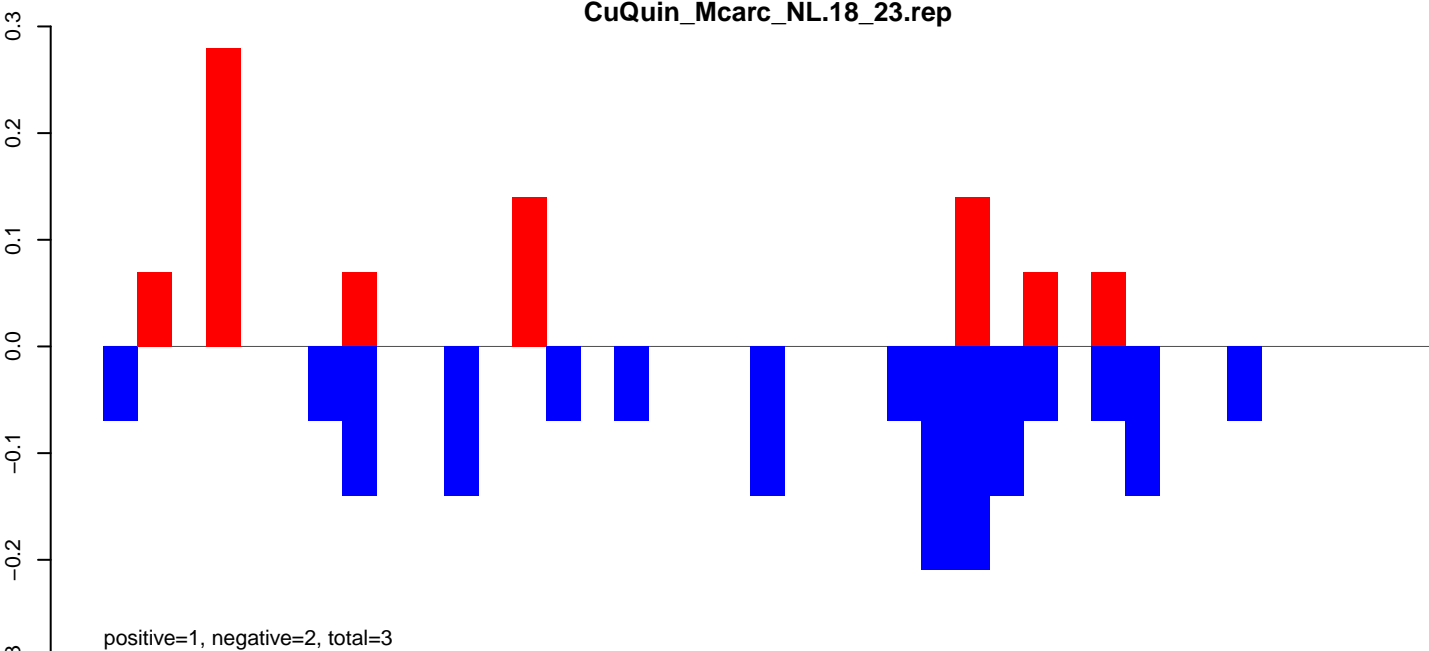


positive=2, negative=2, total=4

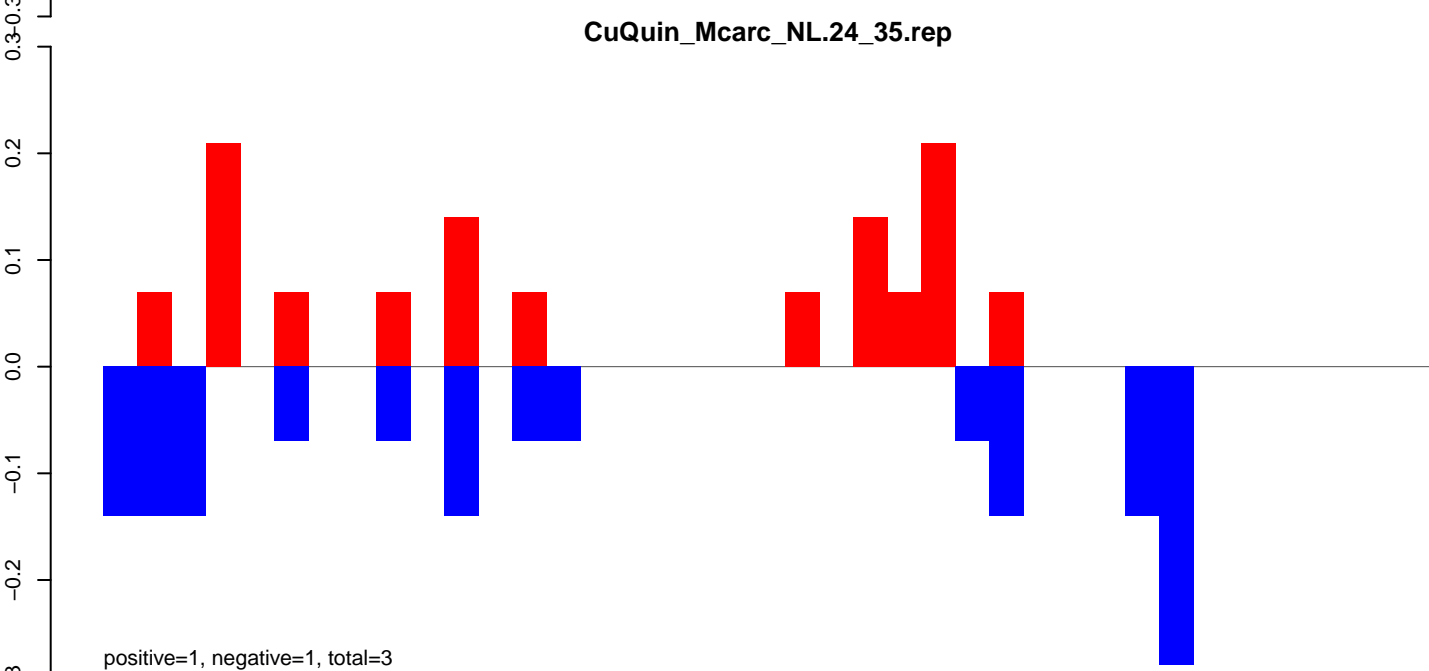
Window size=25, length=1109, TE@rnd-4_family-2243@DNA@2839:1-1109

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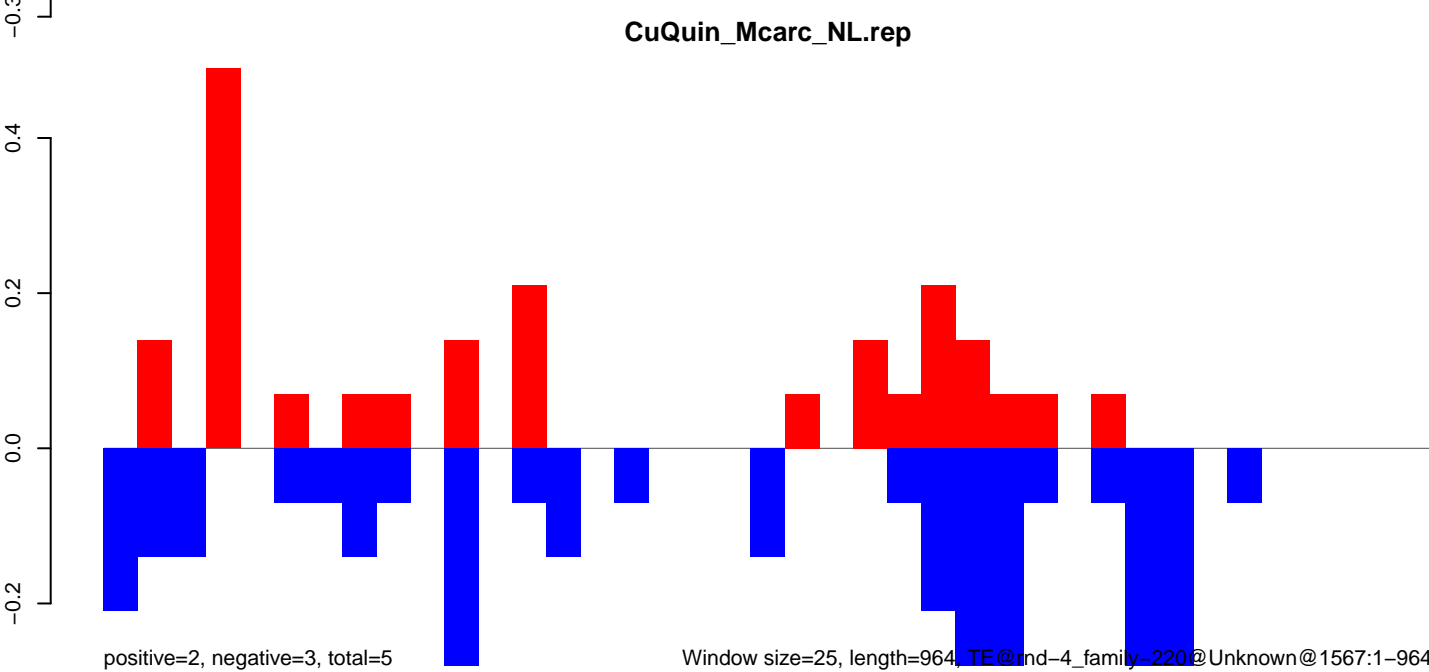
CuQuin_Mcarc_NL.18_23.rep



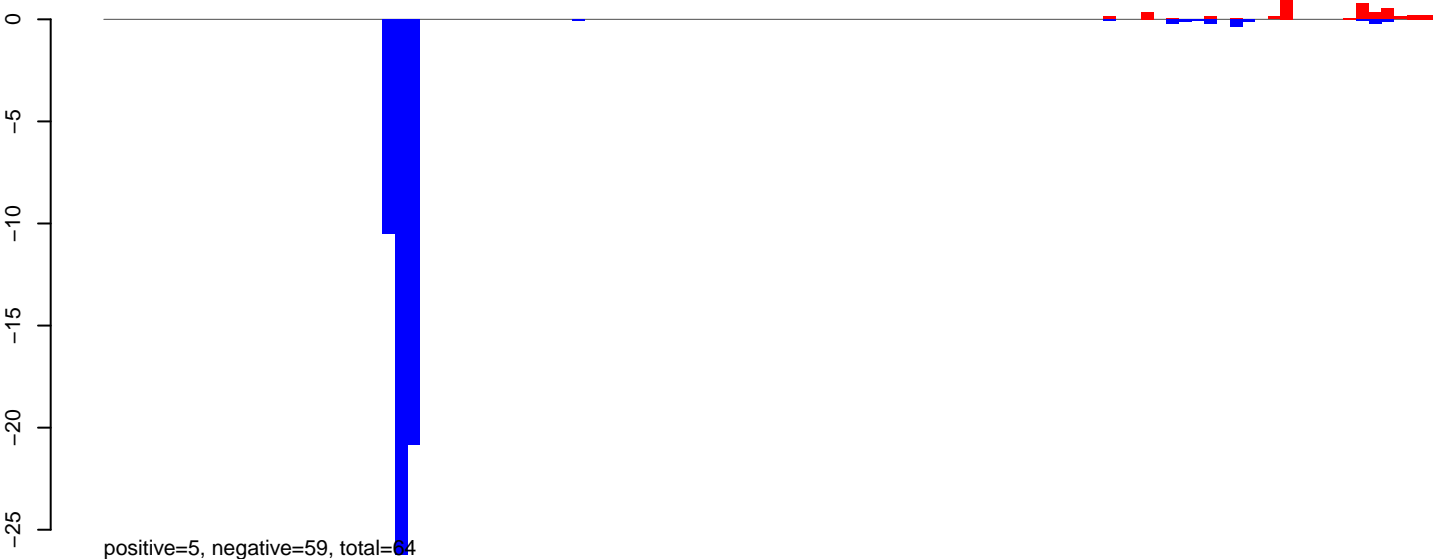
CuQuin_Mcarc_NL.24_35.rep



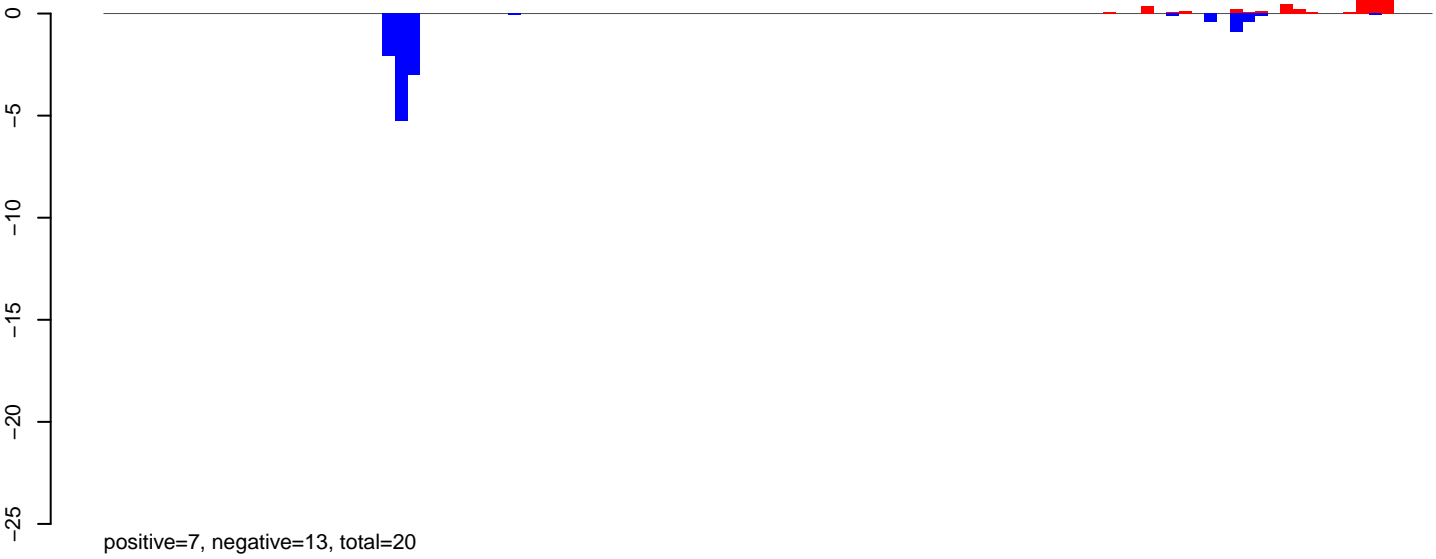
CuQuin_Mcarc_NL.rep



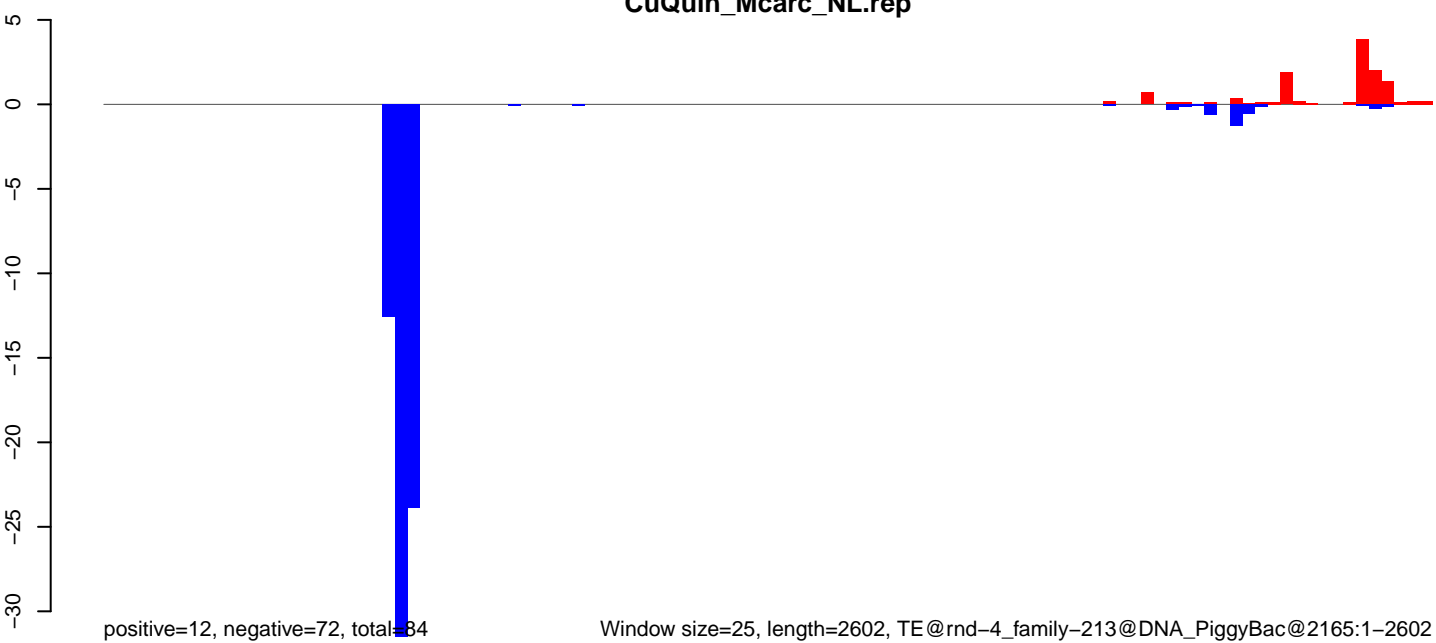
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

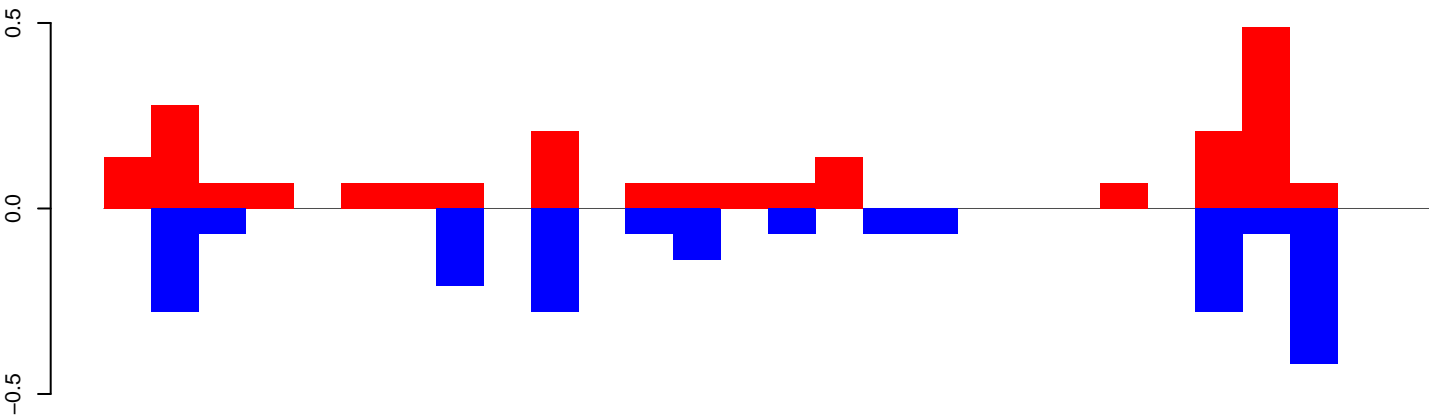


CuQuin_Mcarc_NL.rep



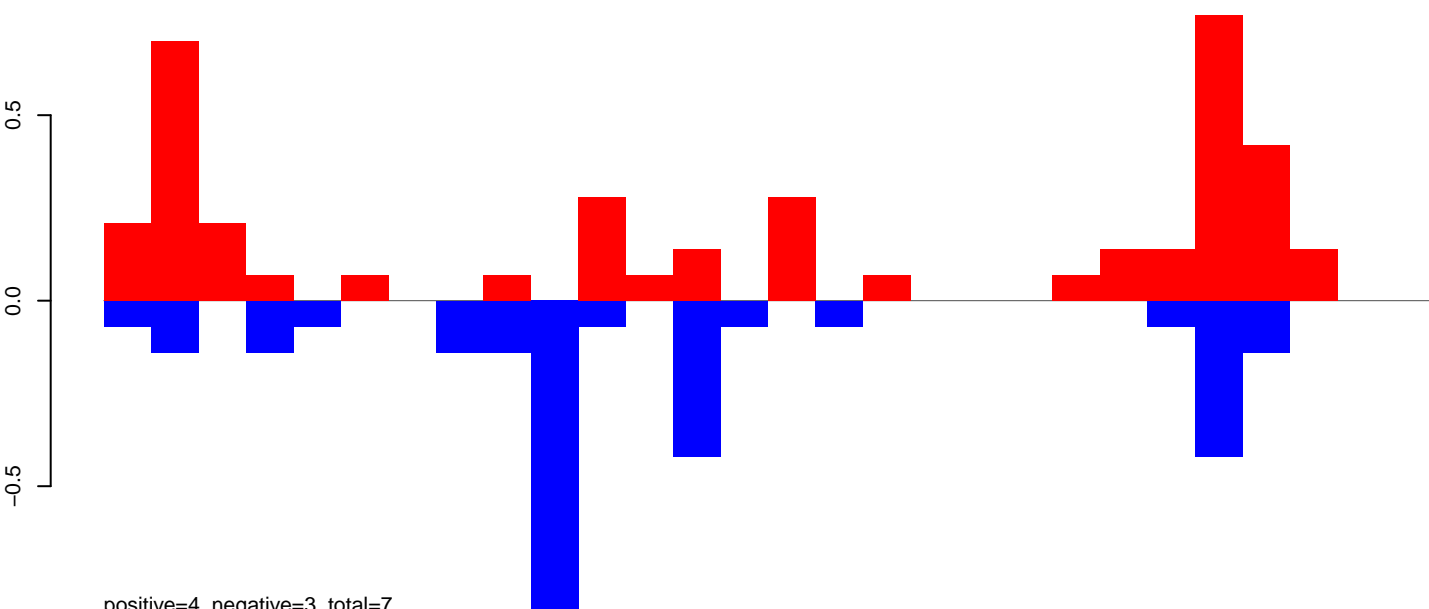
Window size=25, length=2602, TE@rnd-4_family-213@DNA_PiggyBac@2165:1-2602

CuQuin_Mcarc_NL.18_23.rep



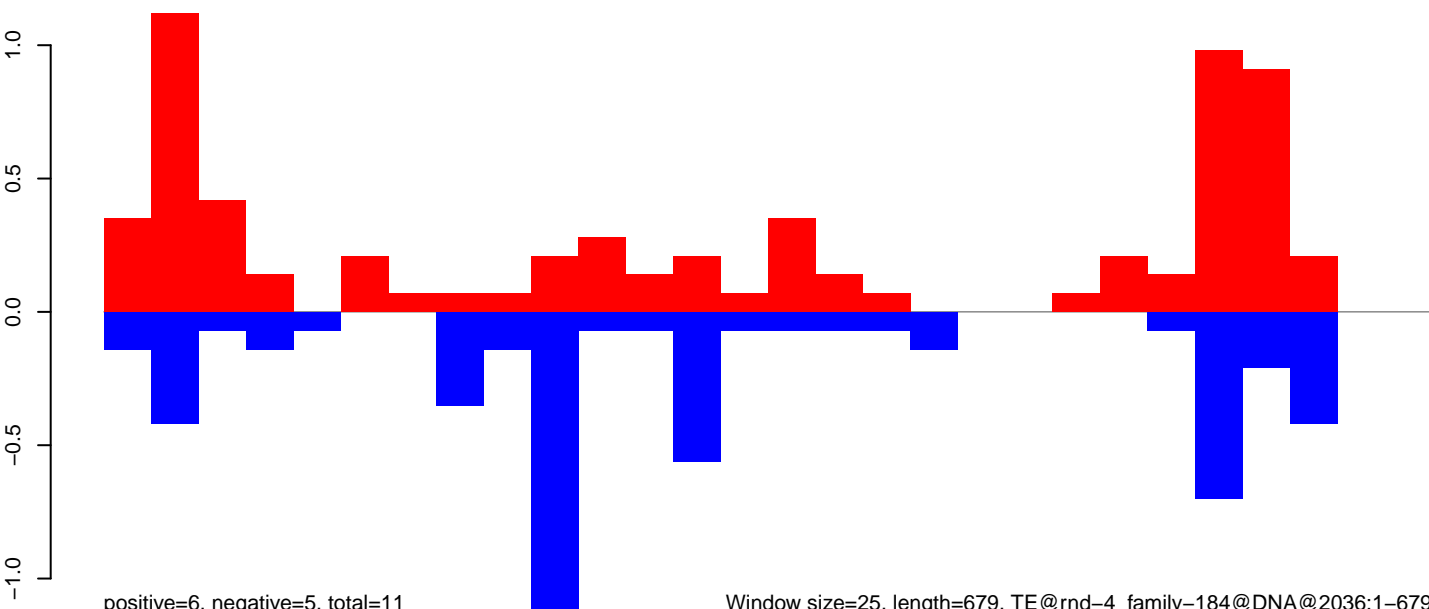
positive=2, negative=2, total=4

CuQuin_Mcarc_NL.24_35.rep



positive=4, negative=3, total=7

CuQuin_Mcarc_NL.rep

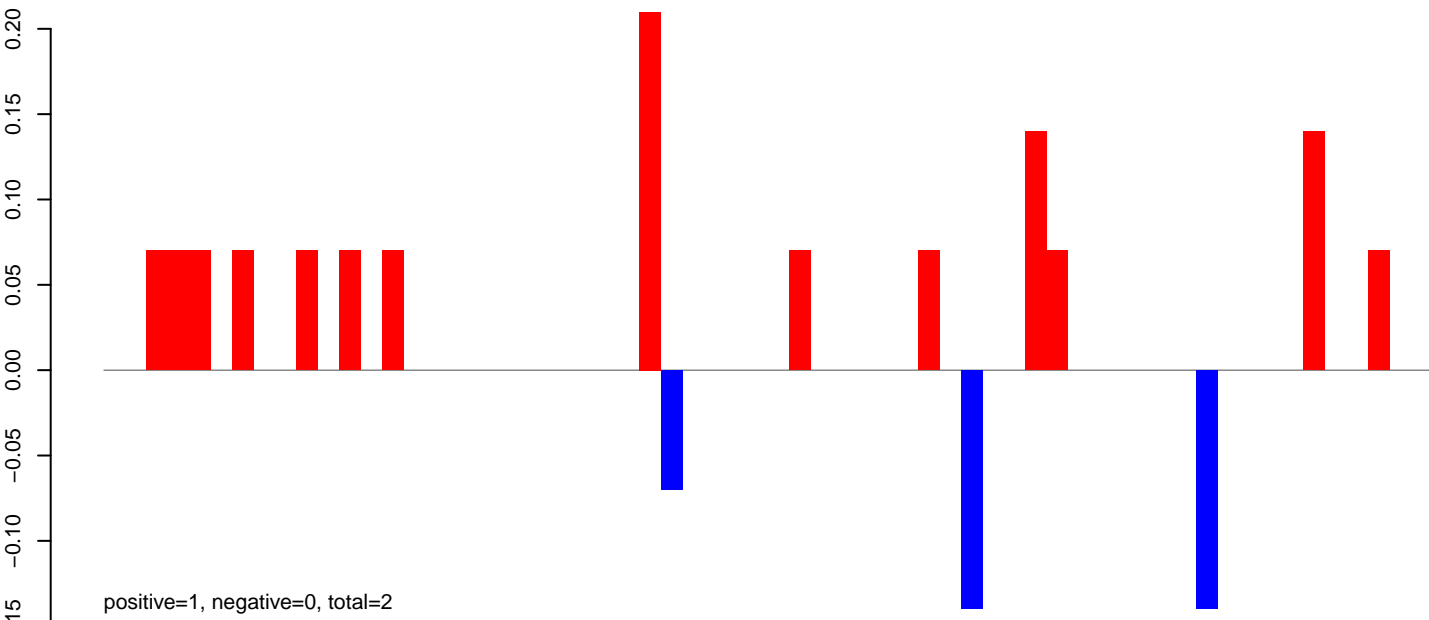


positive=6, negative=5, total=11

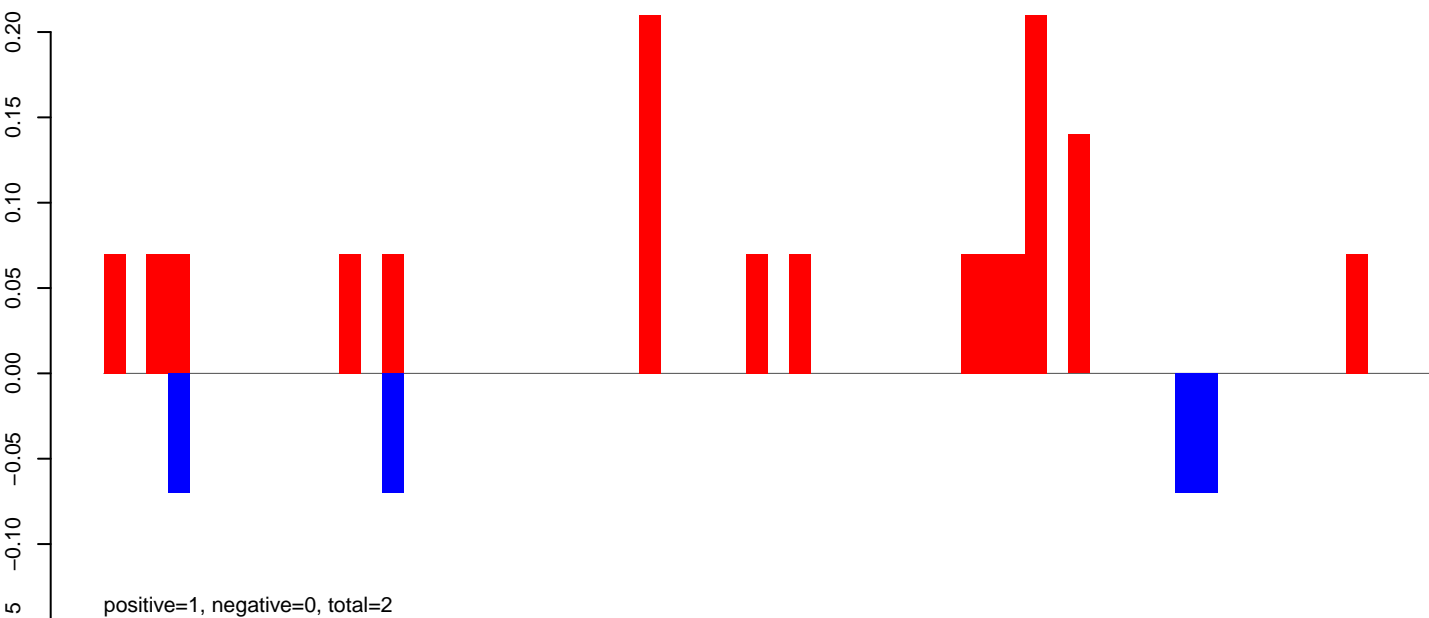
Window size=25, length=679, TE@rnd-4_family-184@DNA@2036:1-679

0 100 200 300 400 500 600 700

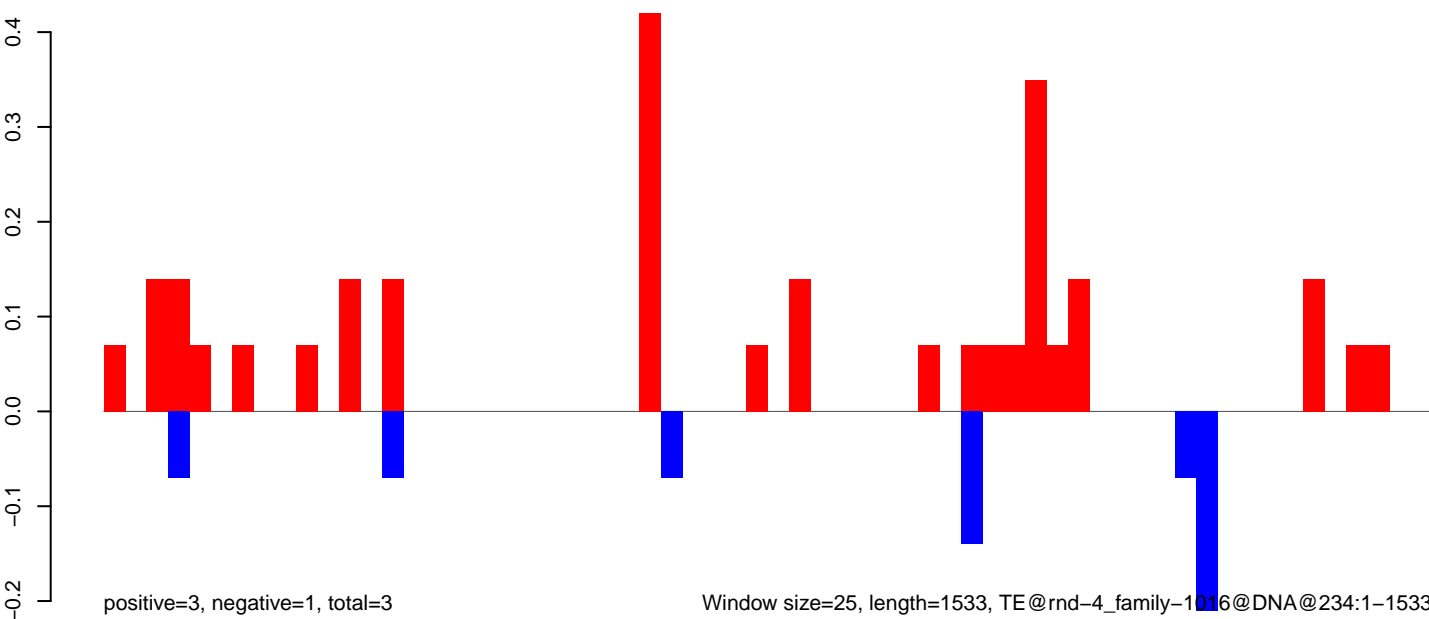
CuQuin_Mcarc_NL.18_23.rep



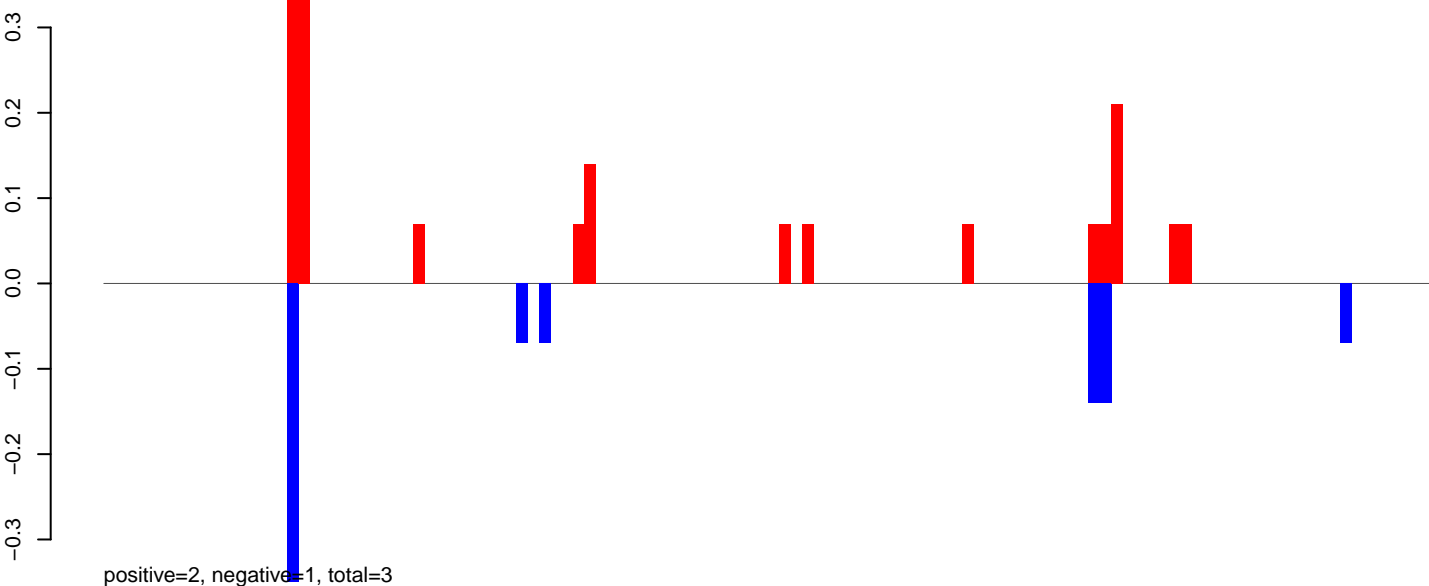
CuQuin_Mcarc_NL.24_35.rep



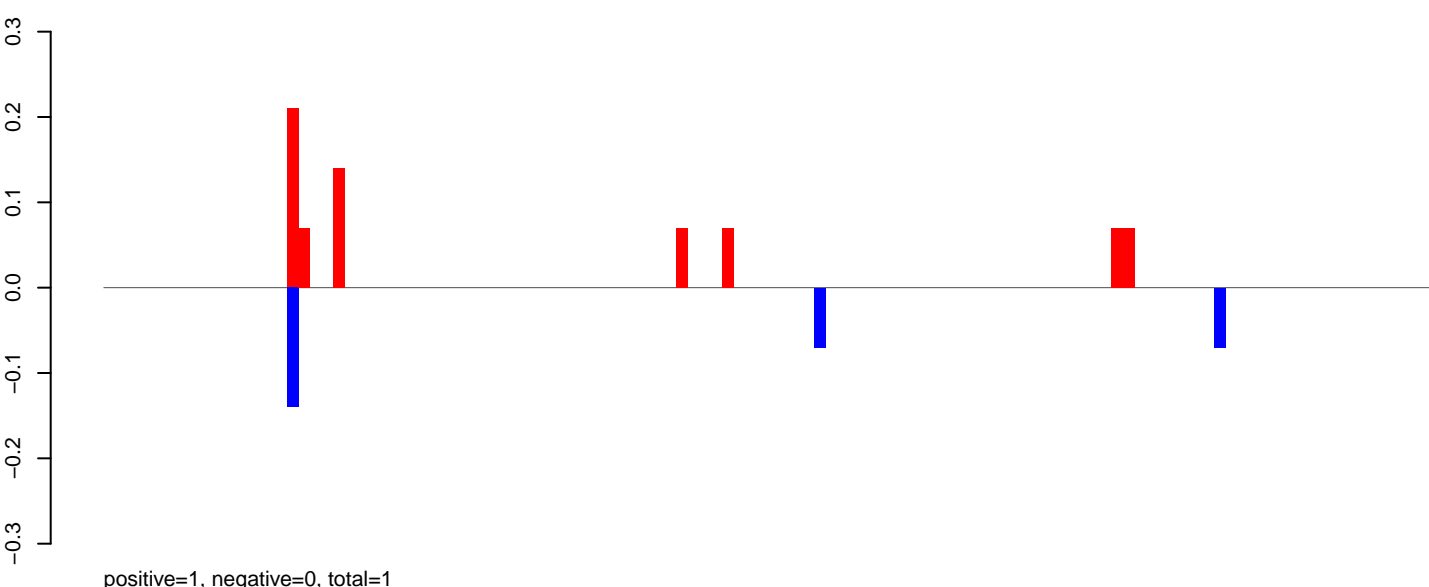
CuQuin_Mcarc_NL.rep



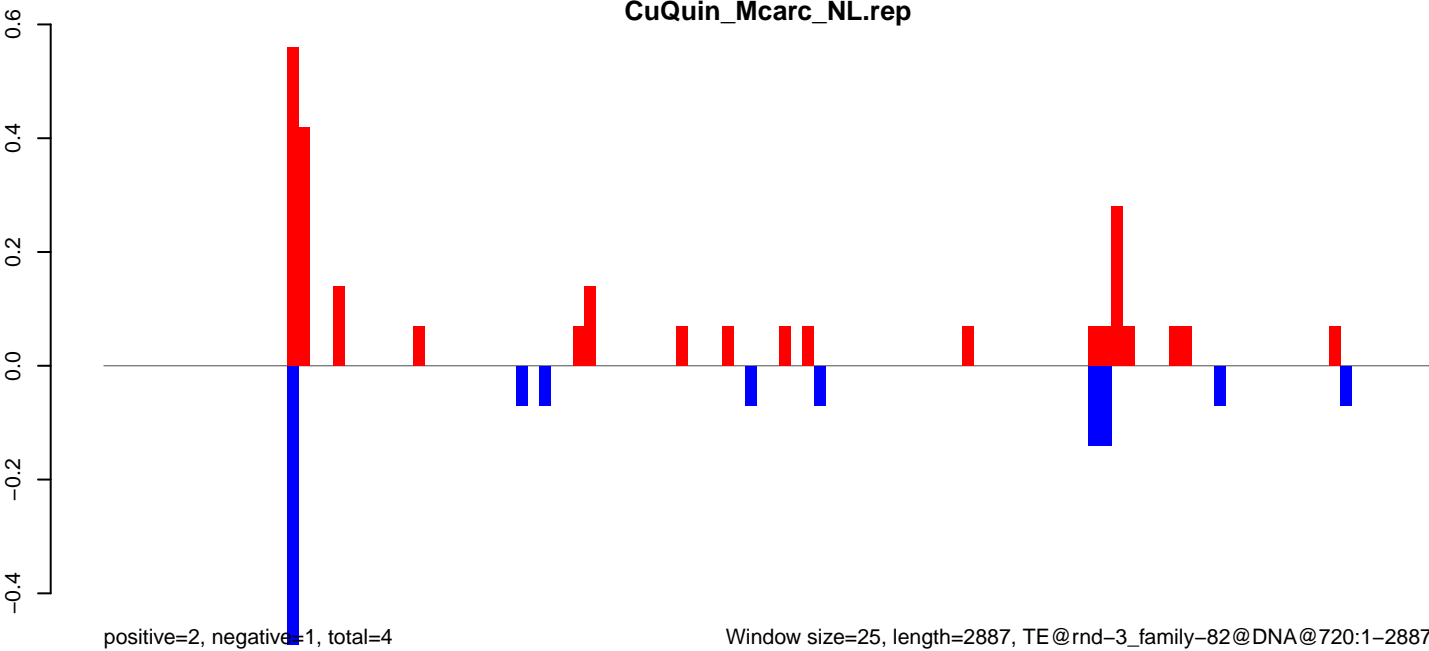
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



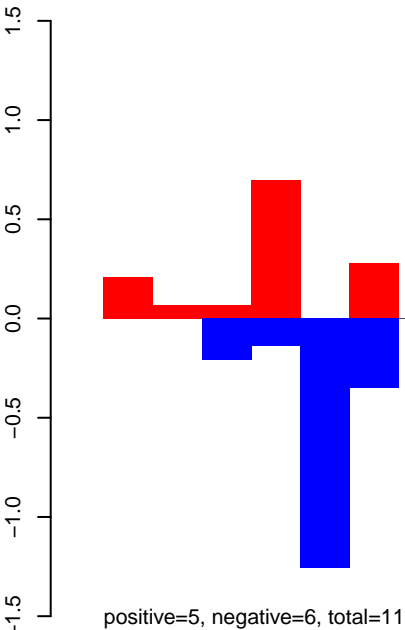
CuQuin_Mcarc_NL.rep



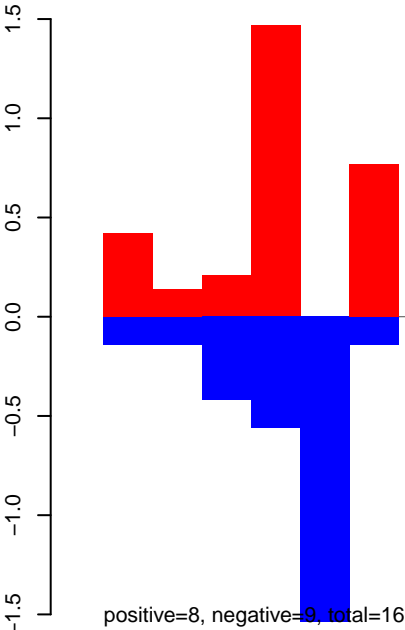
Window size=25, length=2887, TE@rnd-3_family-82@DNA@720:1-2887

0 500 1000 1500 2000 2500 3000

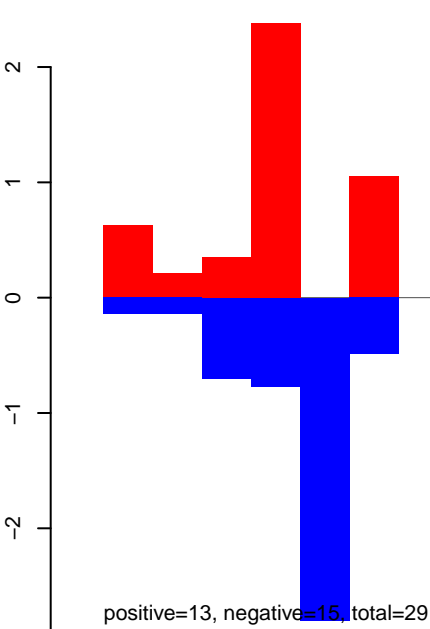
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

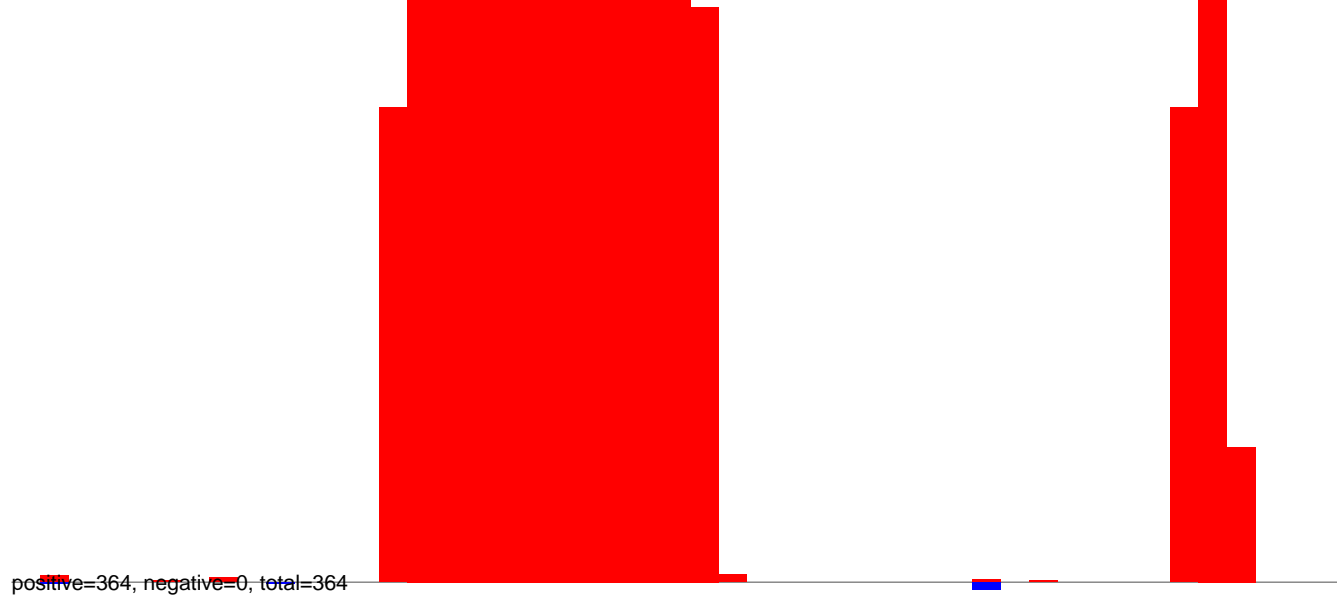


CuQuin_Mcarc_NL.rep

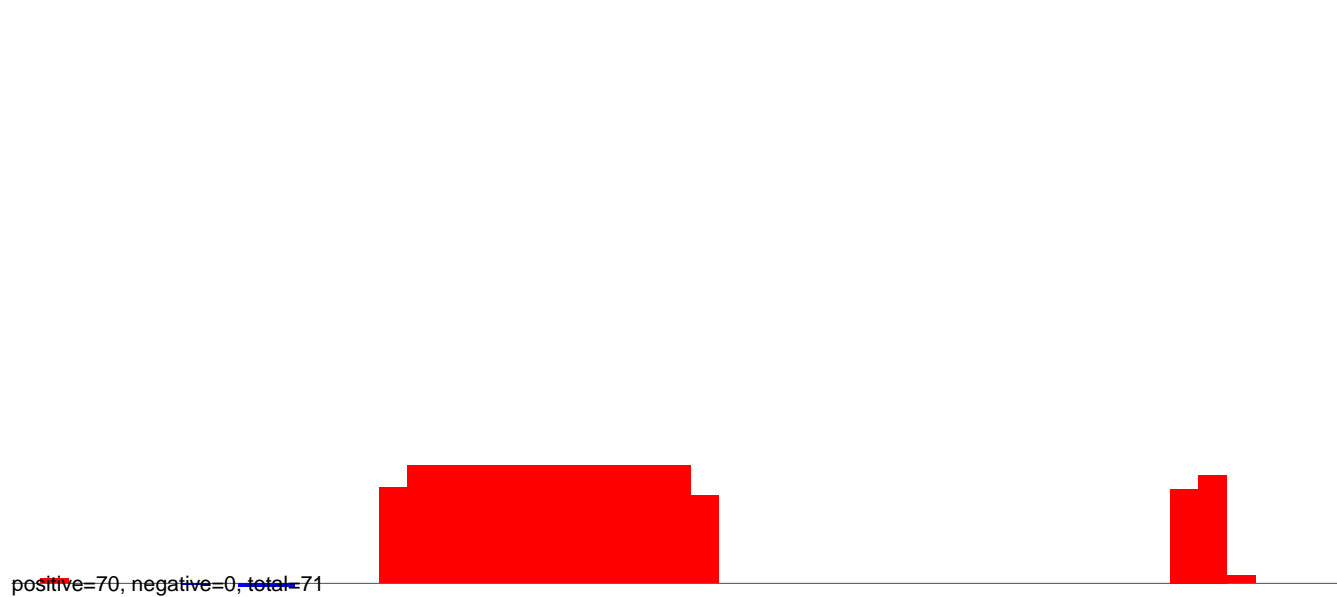


Window size=25, length=671, TE@rnd-3_family-187@DNA_TcMar-Fot1@654:1-671

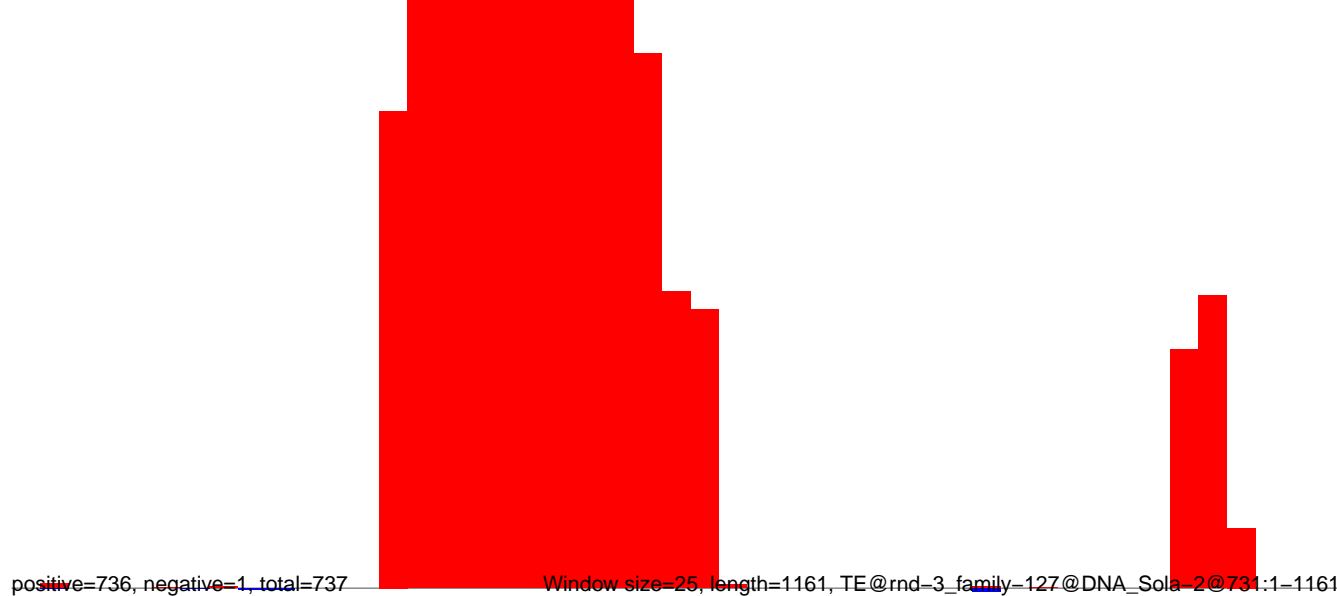
CuQuin_Mcarc_NL.18_23.rep



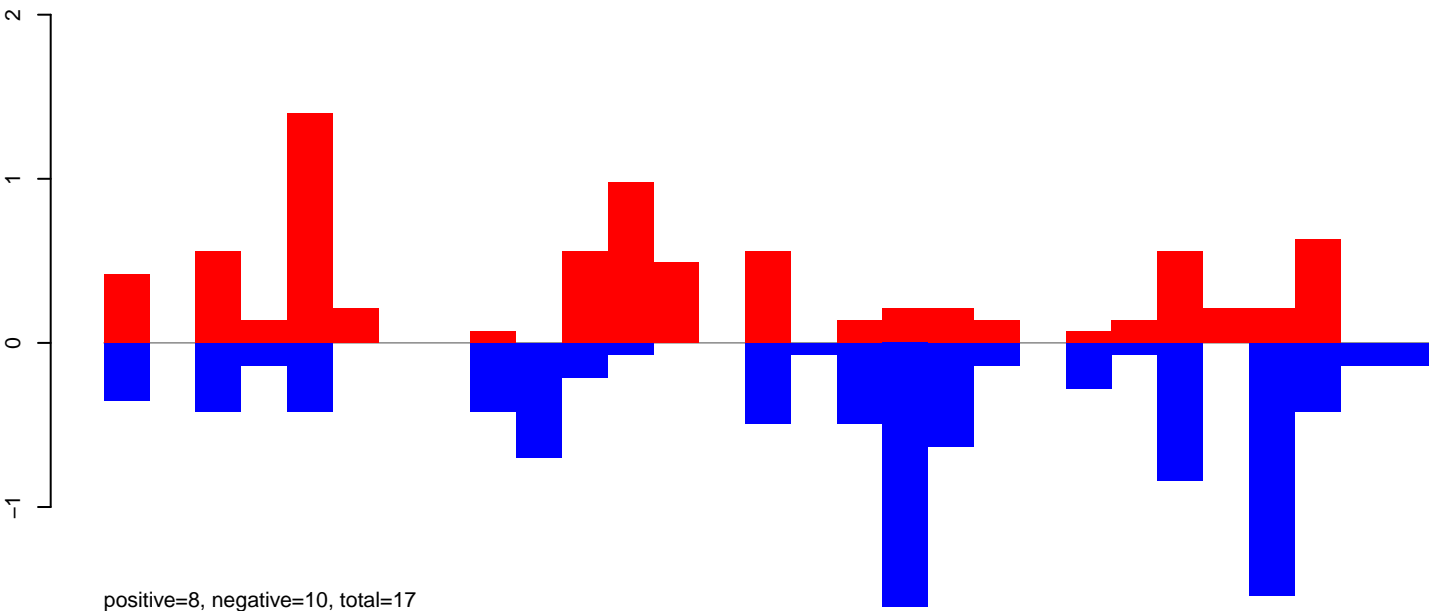
CuQuin_Mcarc_NL.24_35.rep



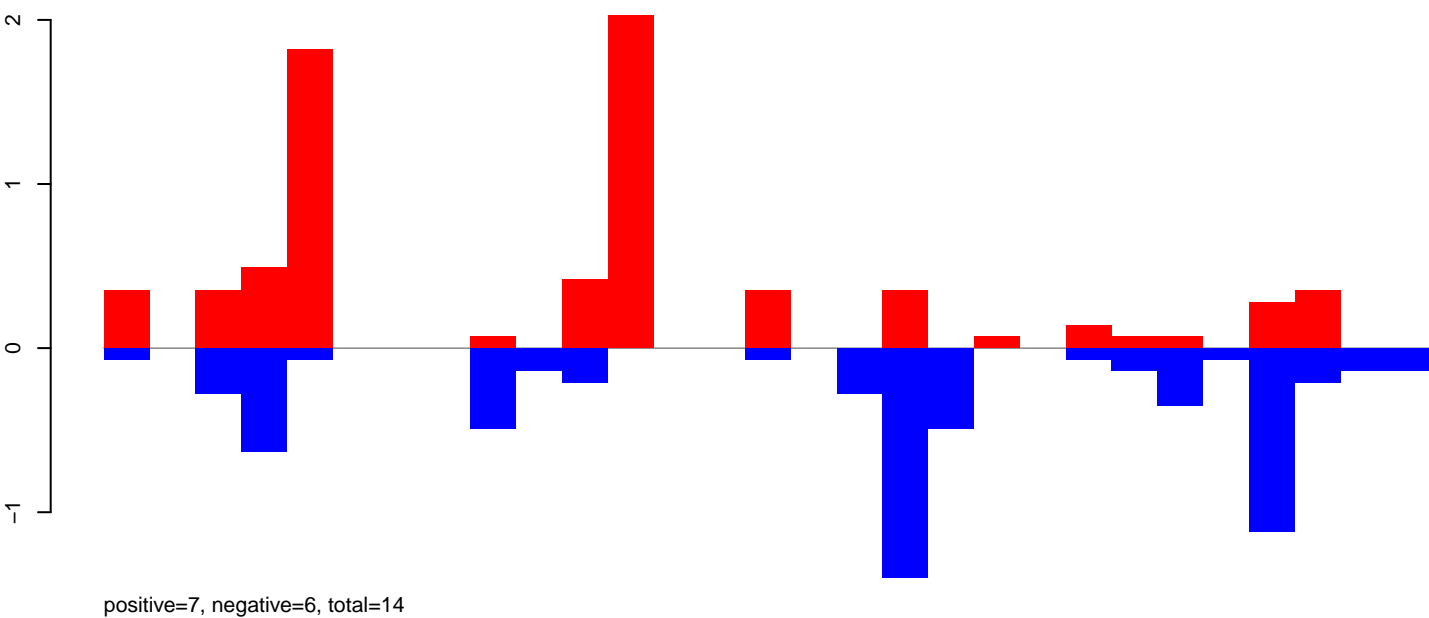
CuQuin_Mcarc_NL.rep



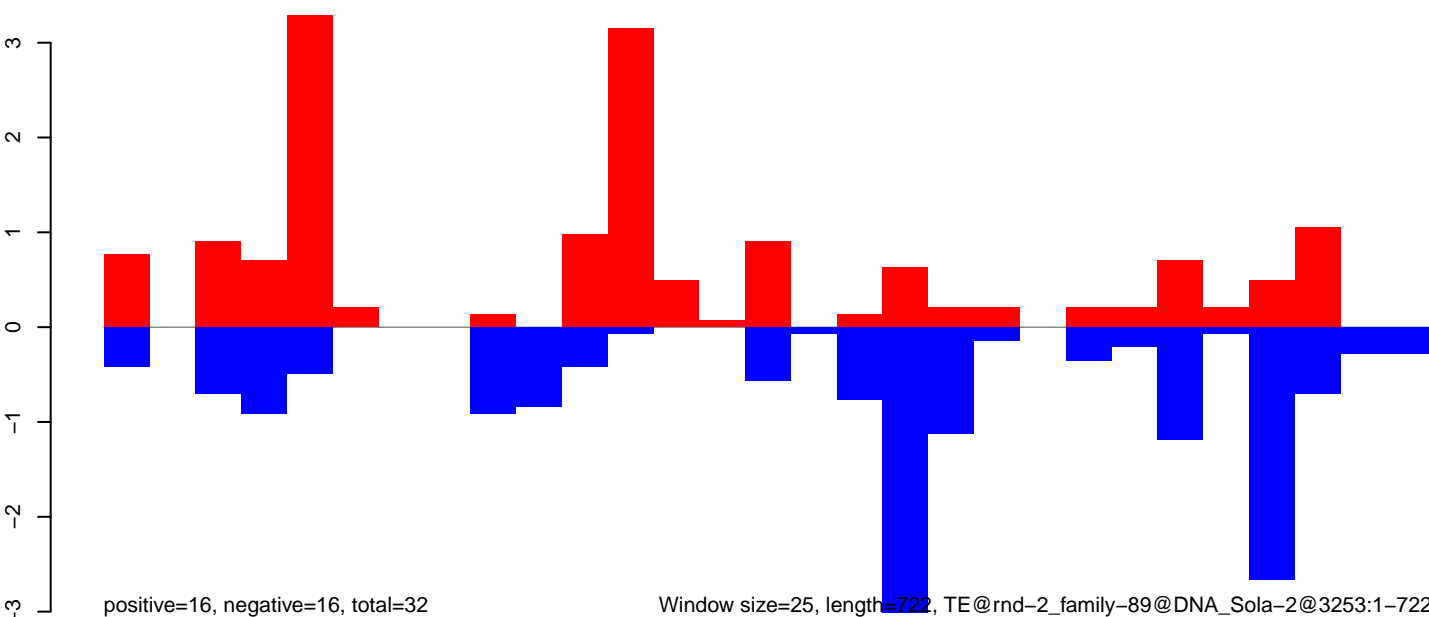
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



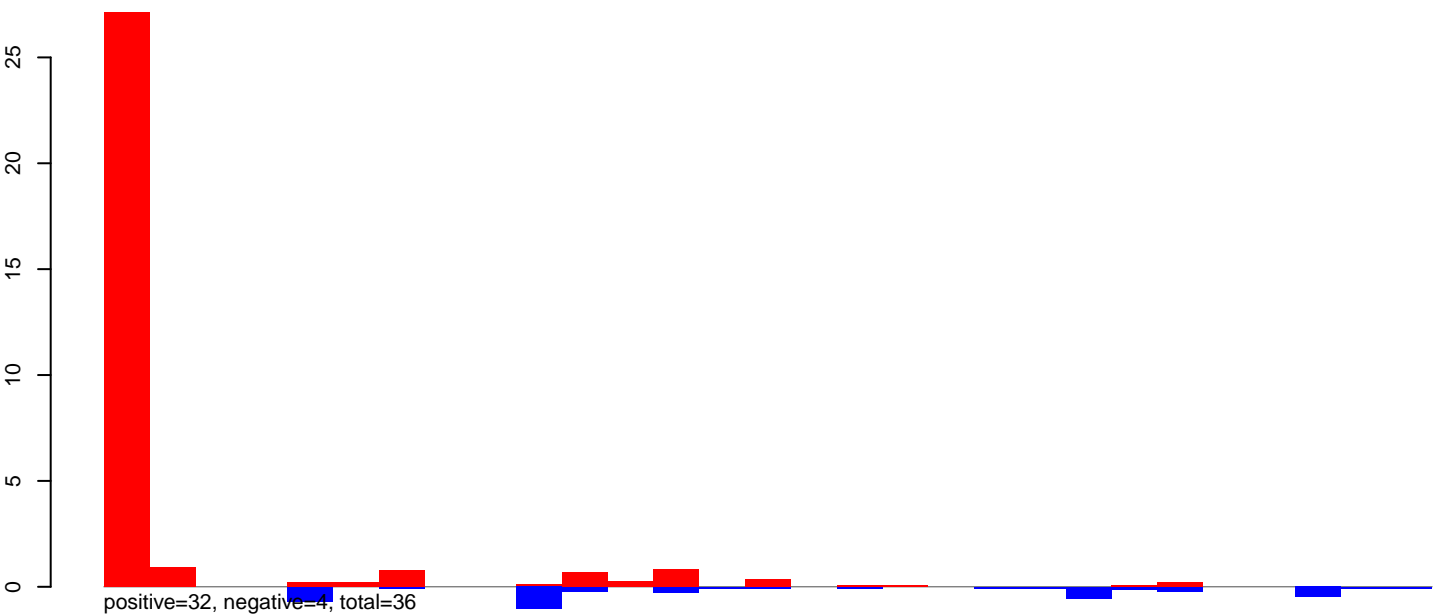
CuQuin_Mcarc_NL.rep



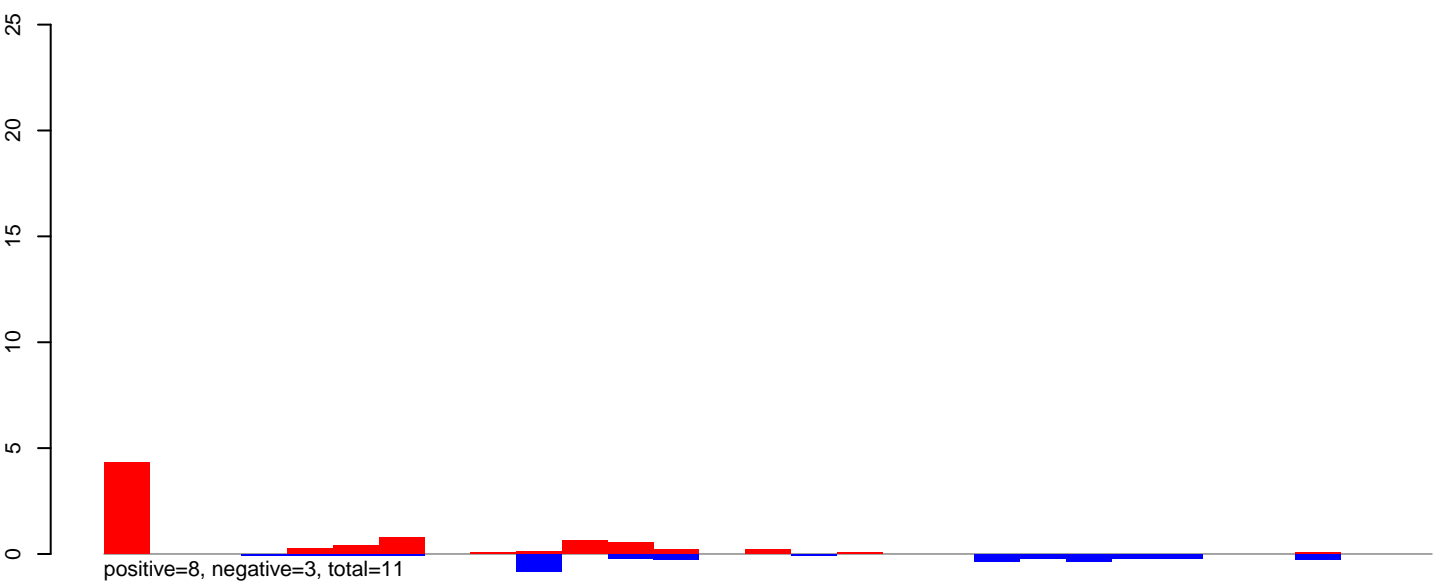
Window size=25, length=722, TE@rnd-2_family-89@DNA_Sola-2@3253:1-722

0 200 400 600

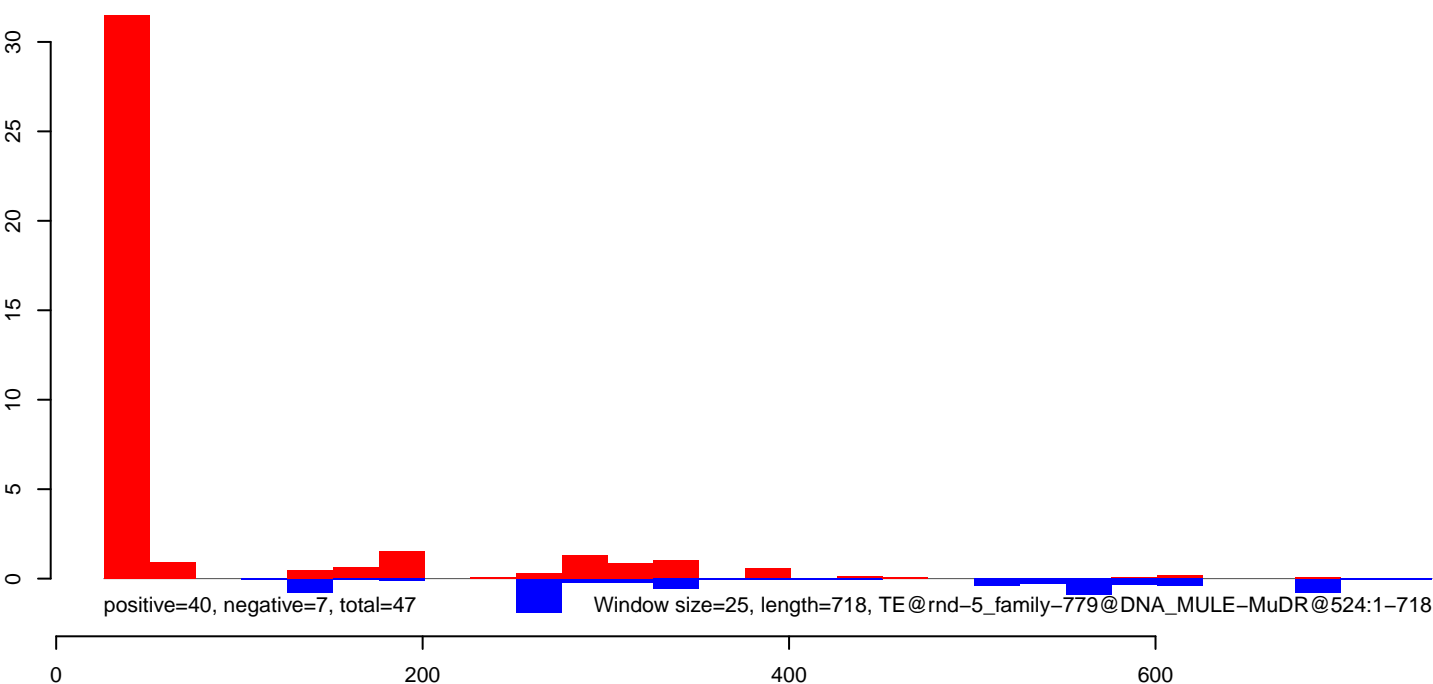
CuQuin_Mcarc_NL.18_23.rep

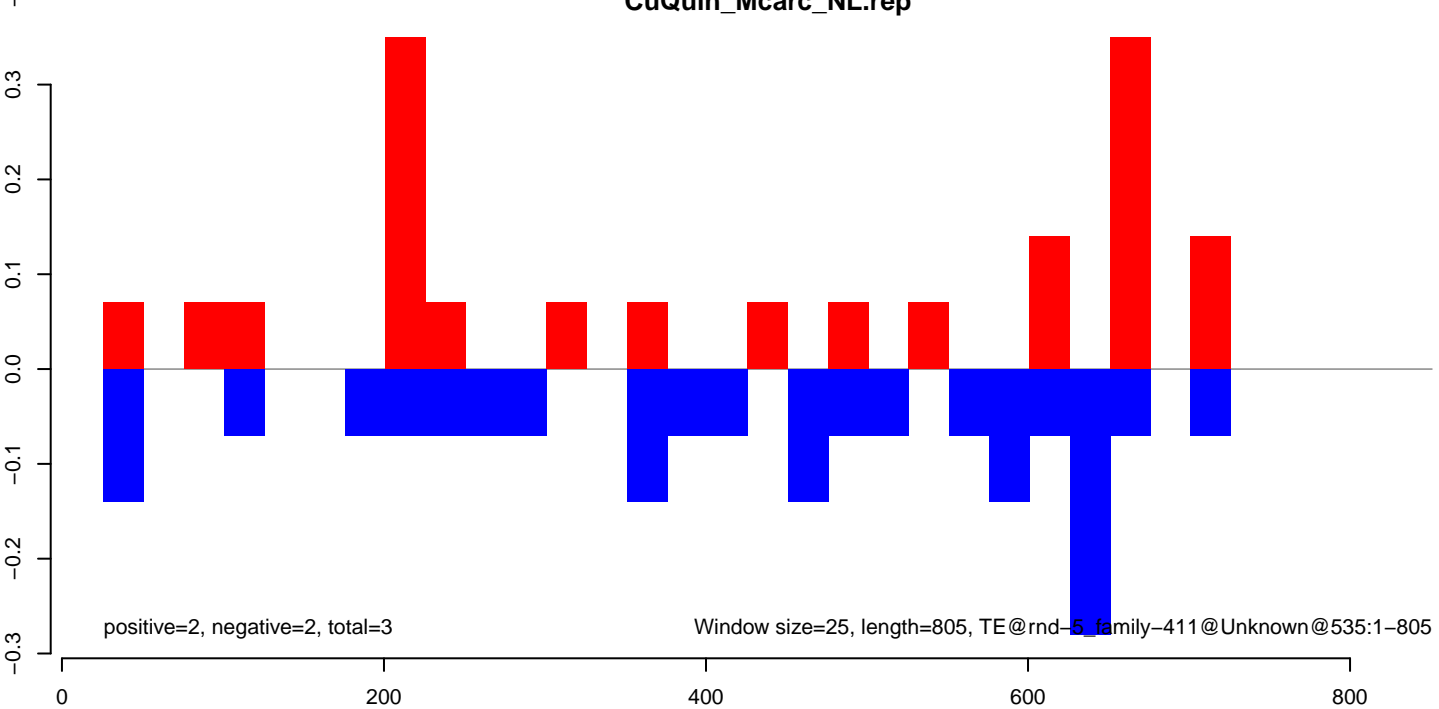
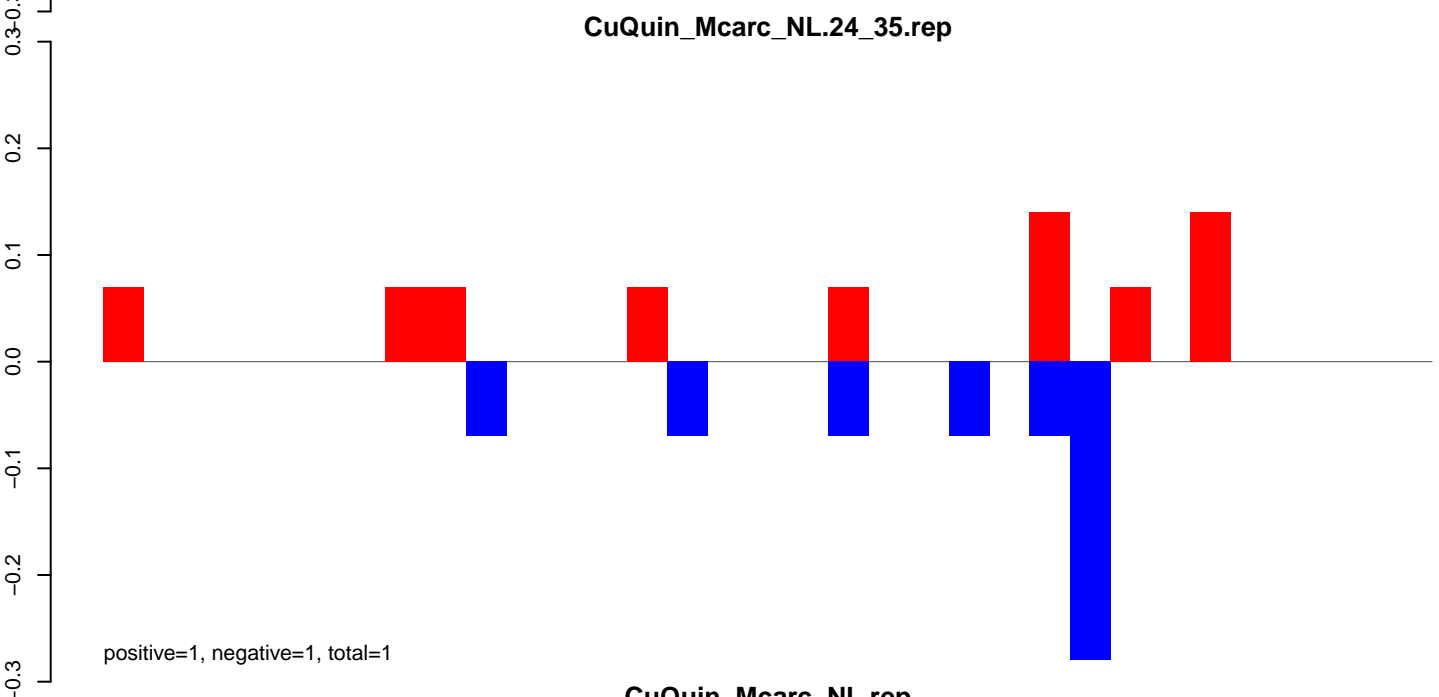
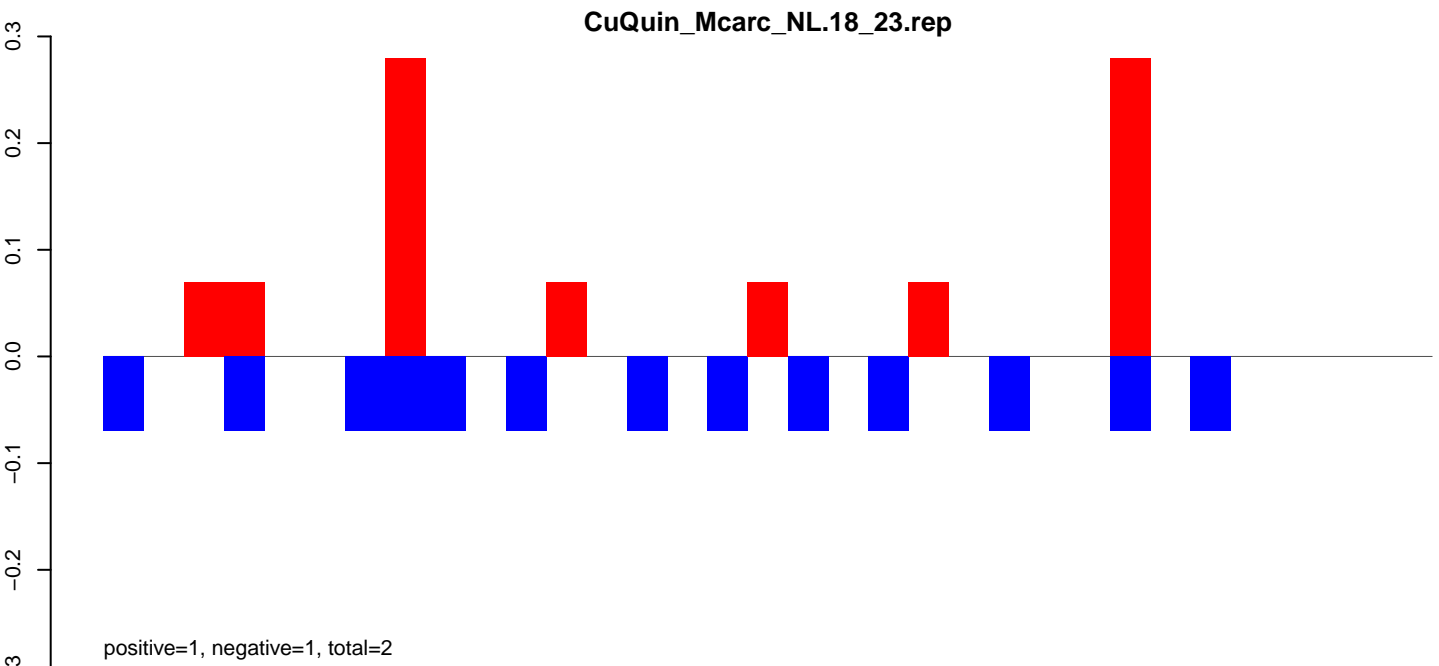


CuQuin_Mcarc_NL.24_35.rep

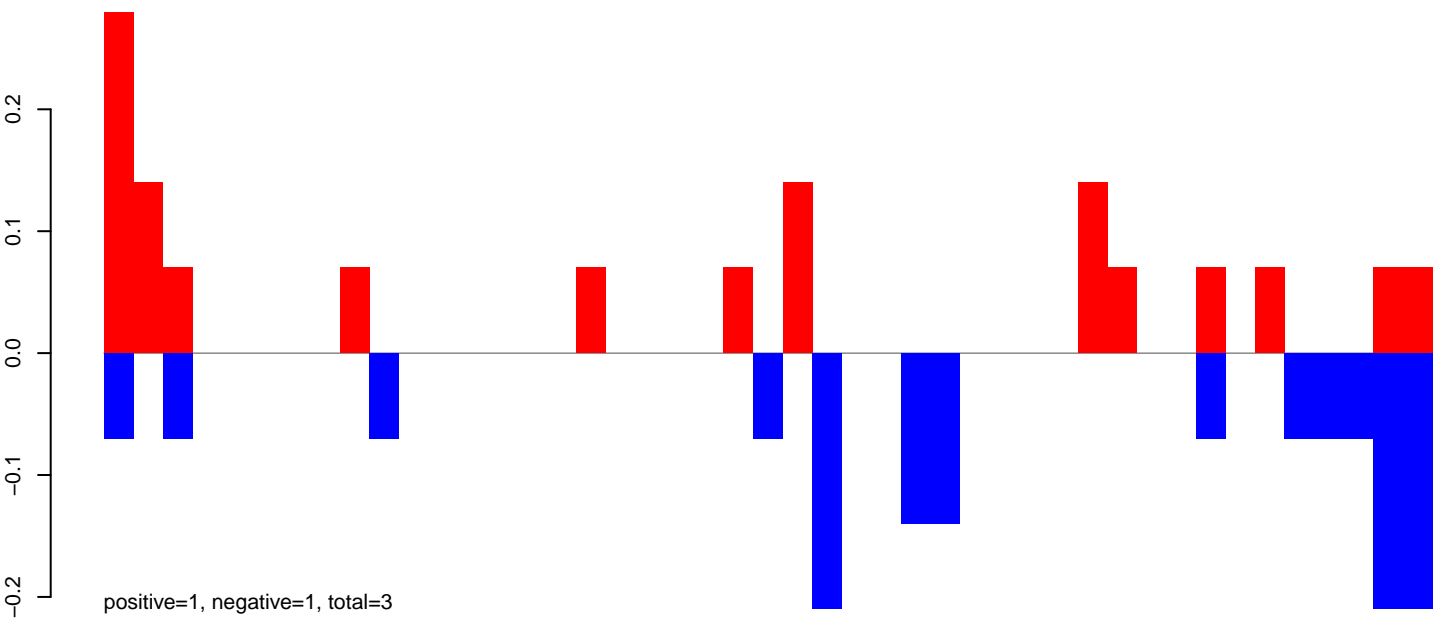


CuQuin_Mcarc_NL.rep

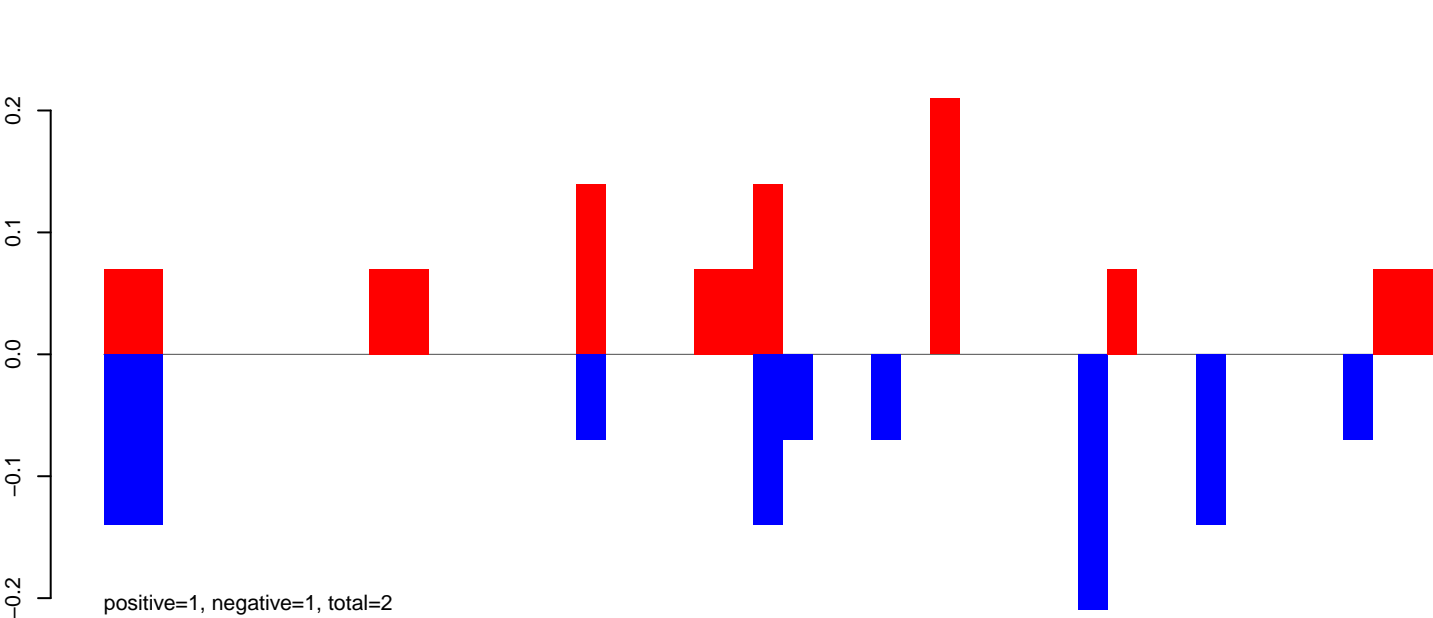




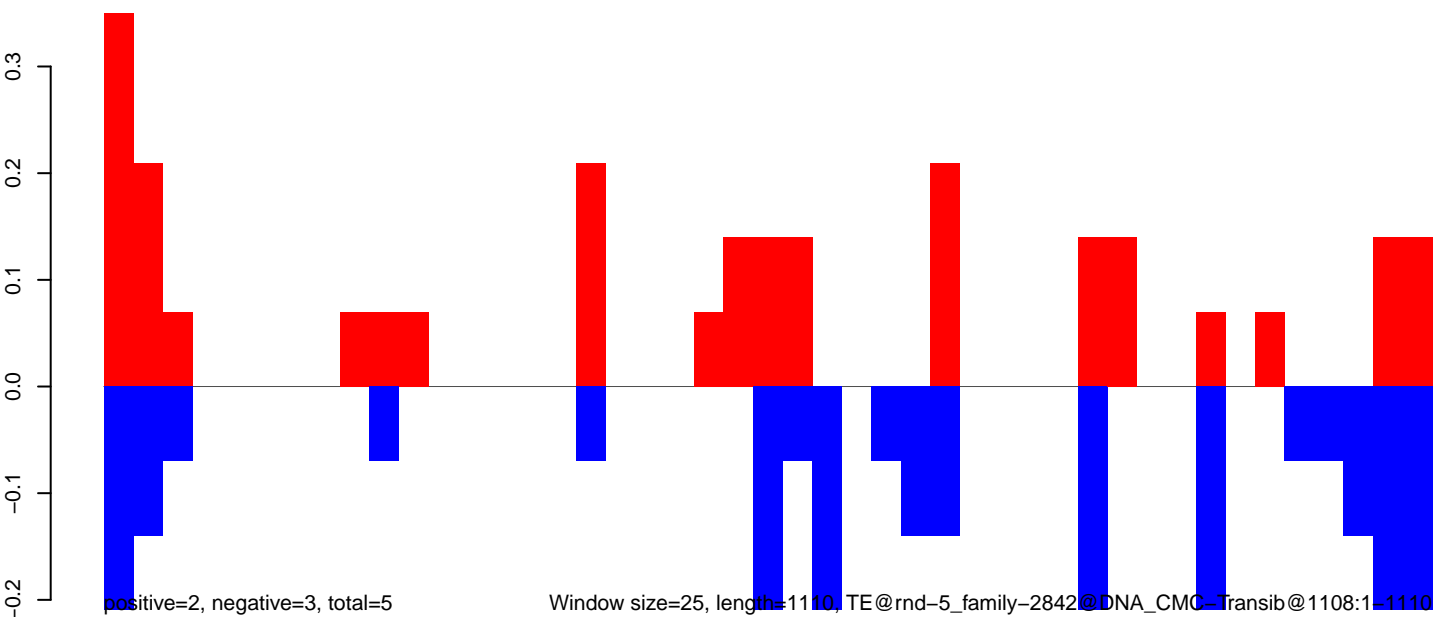
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

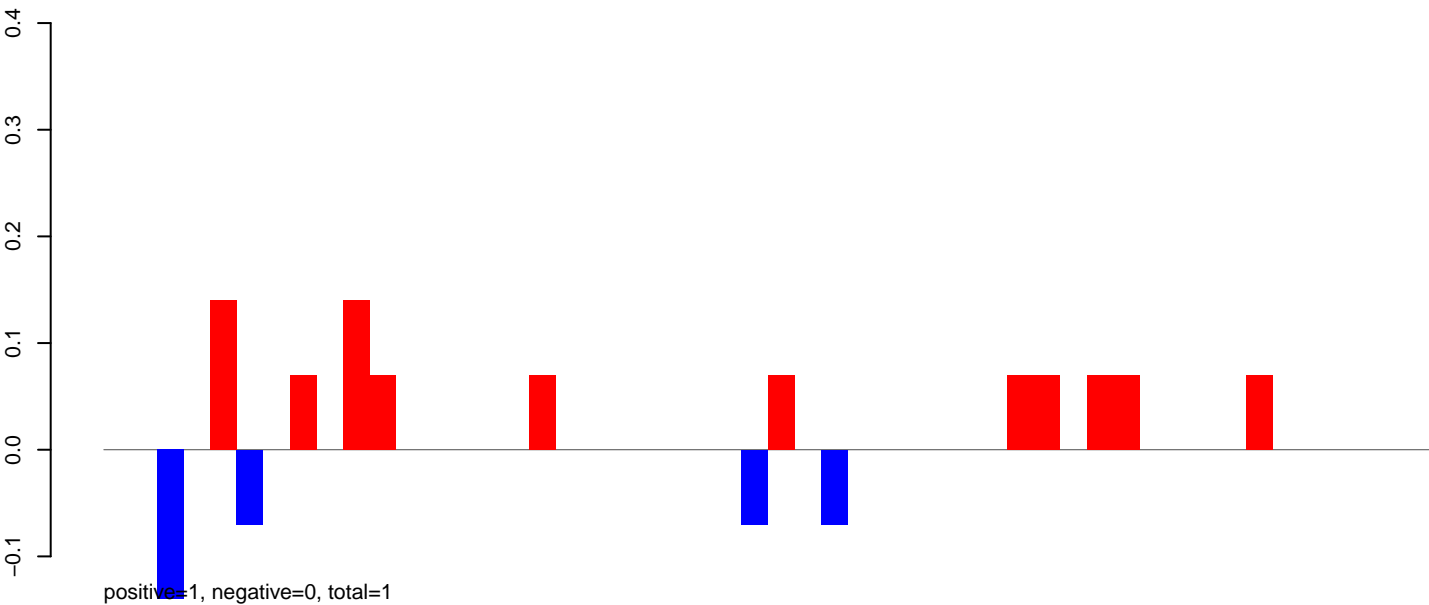


CuQuin_Mcarc_NL.rep

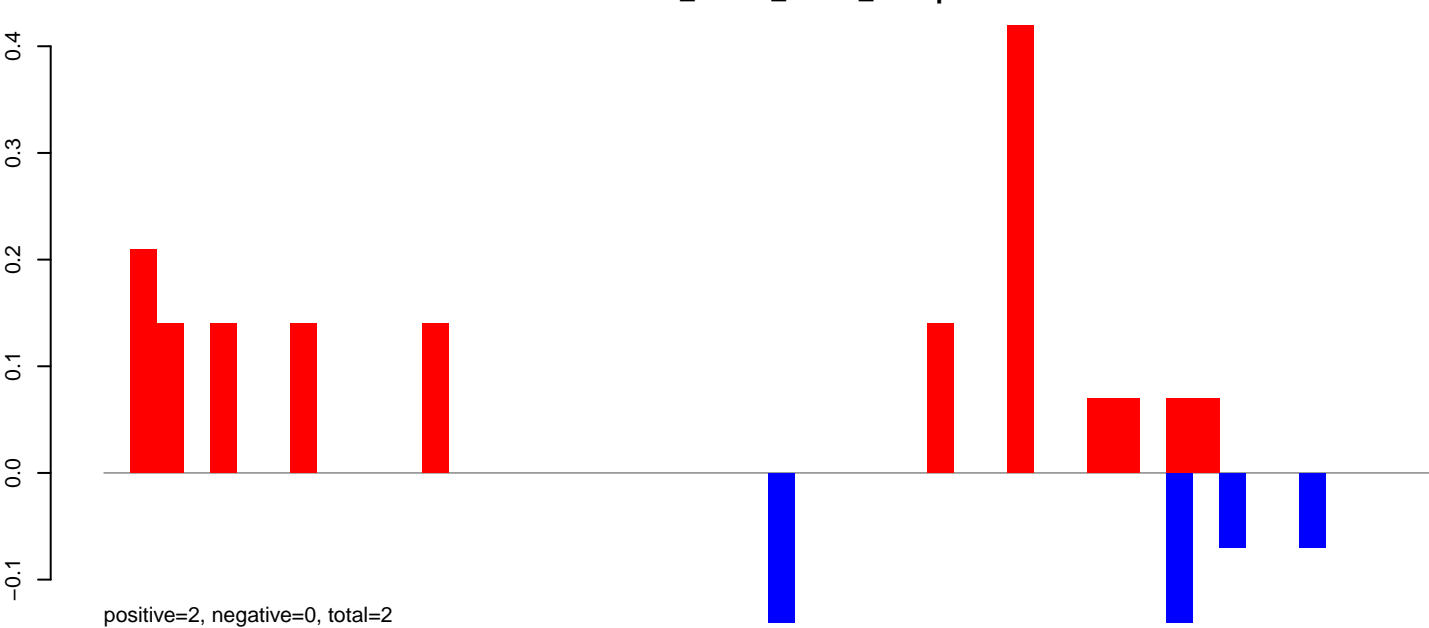


Window size=25, length=1110, TE@rnd-5_family-2842@DNA_CMC-Transib@1108:1-1110

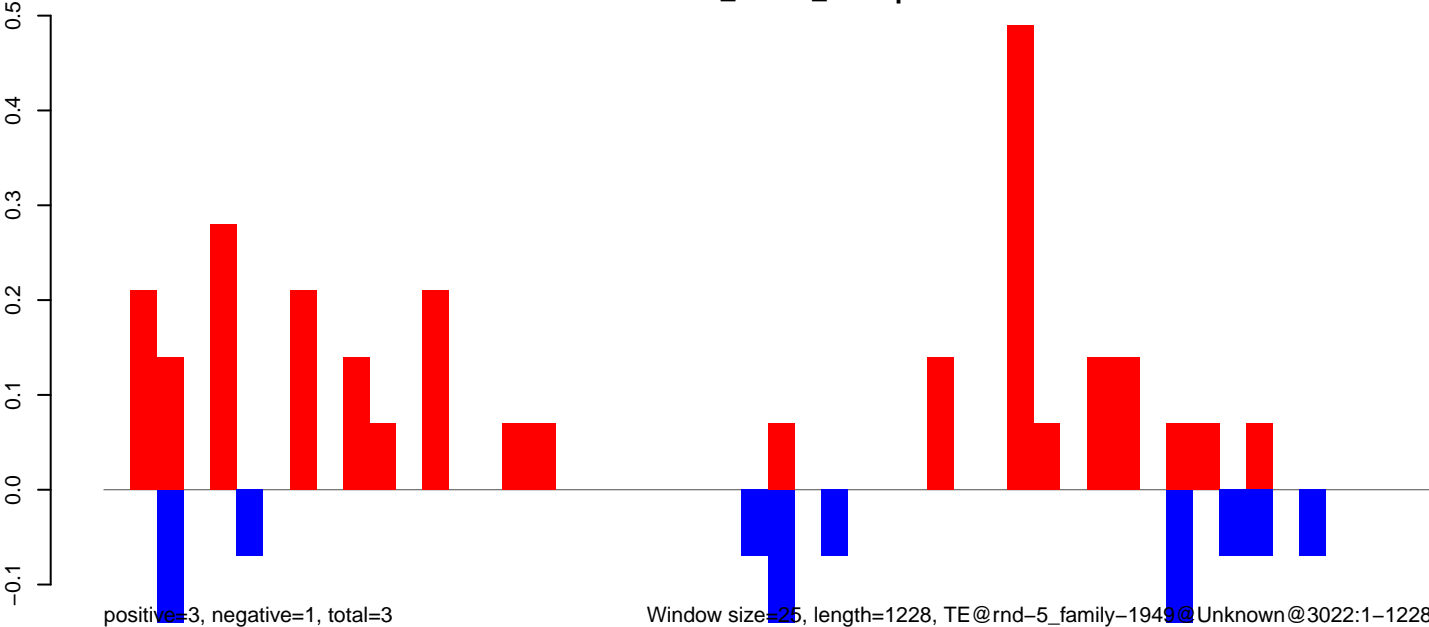
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

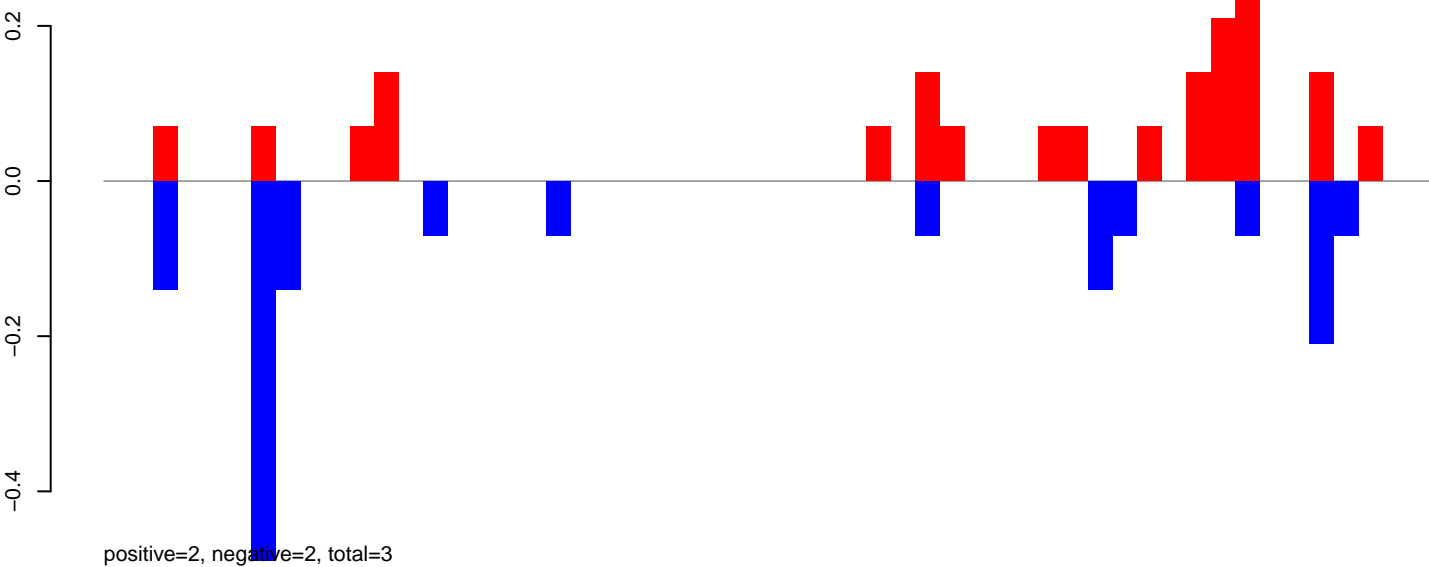


CuQuin_Mcarc_NL.rep



Window size=25, length=1228, TE@rnd-5_family-1949@Unknown@3022:1-1228

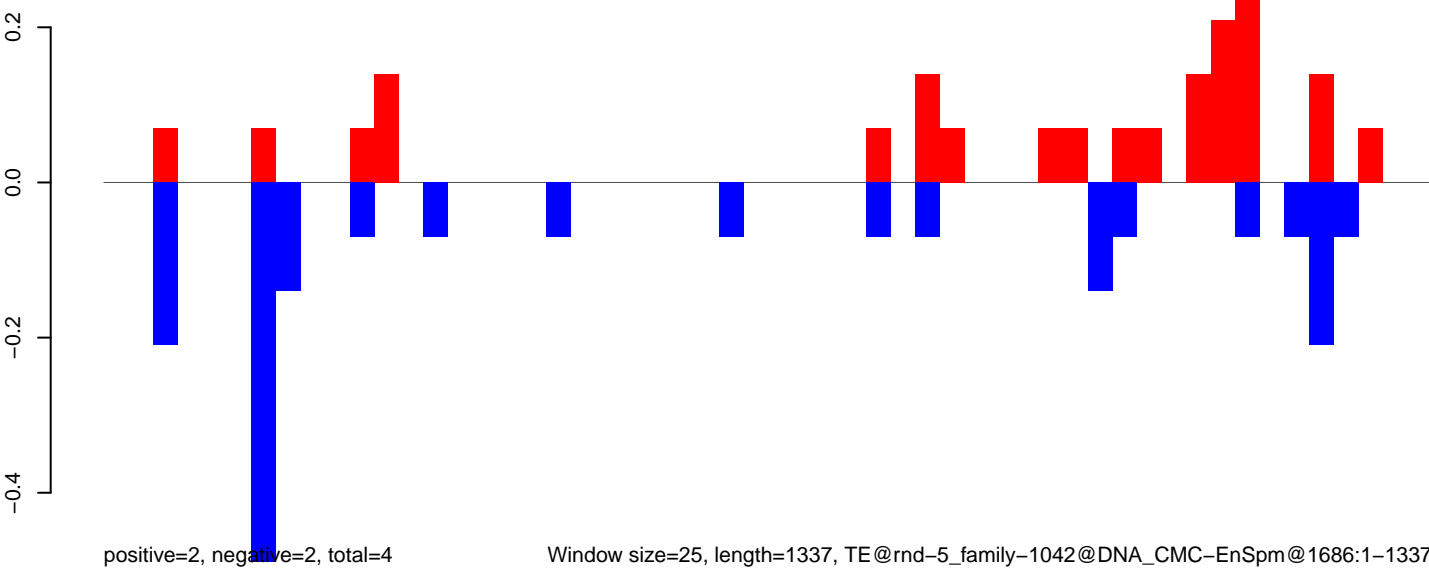
CuQuin_Mcarc_NL.18_23.rep



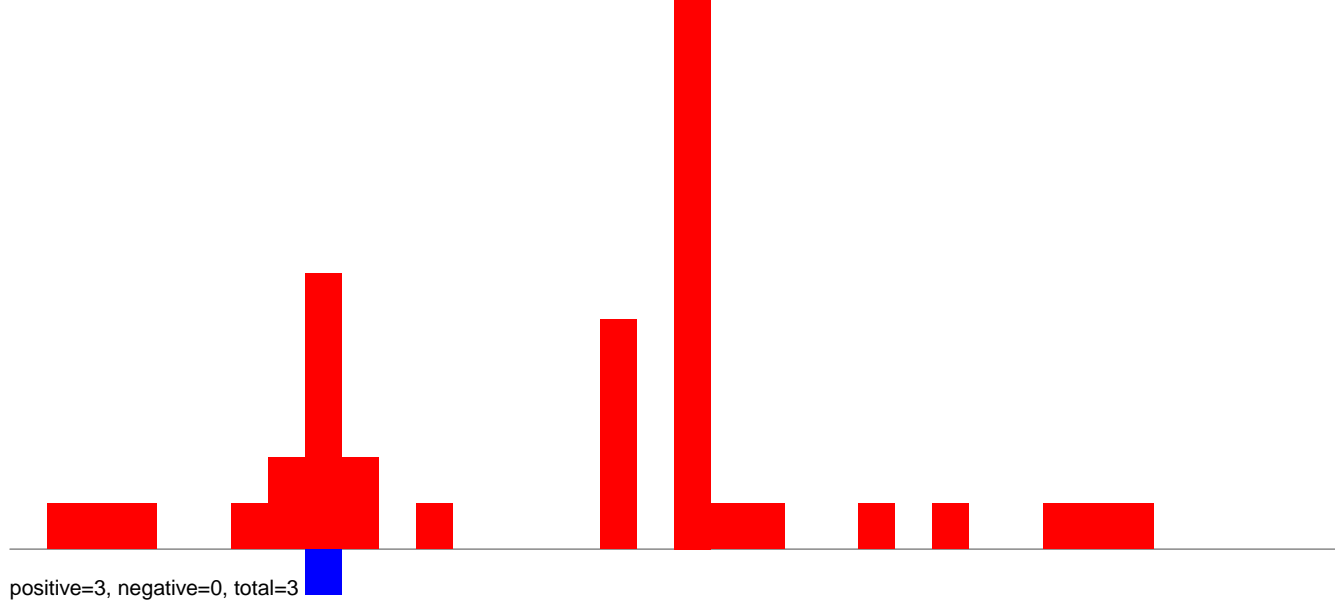
CuQuin_Mcarc_NL.24_35.rep



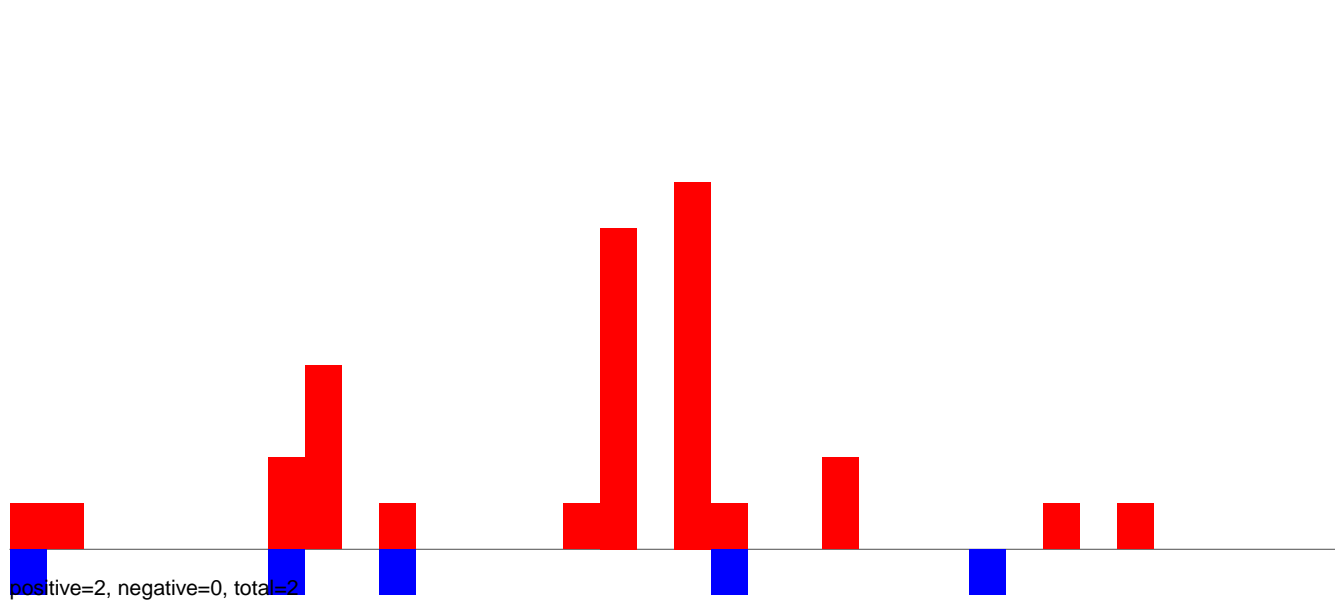
CuQuin_Mcarc_NL.rep



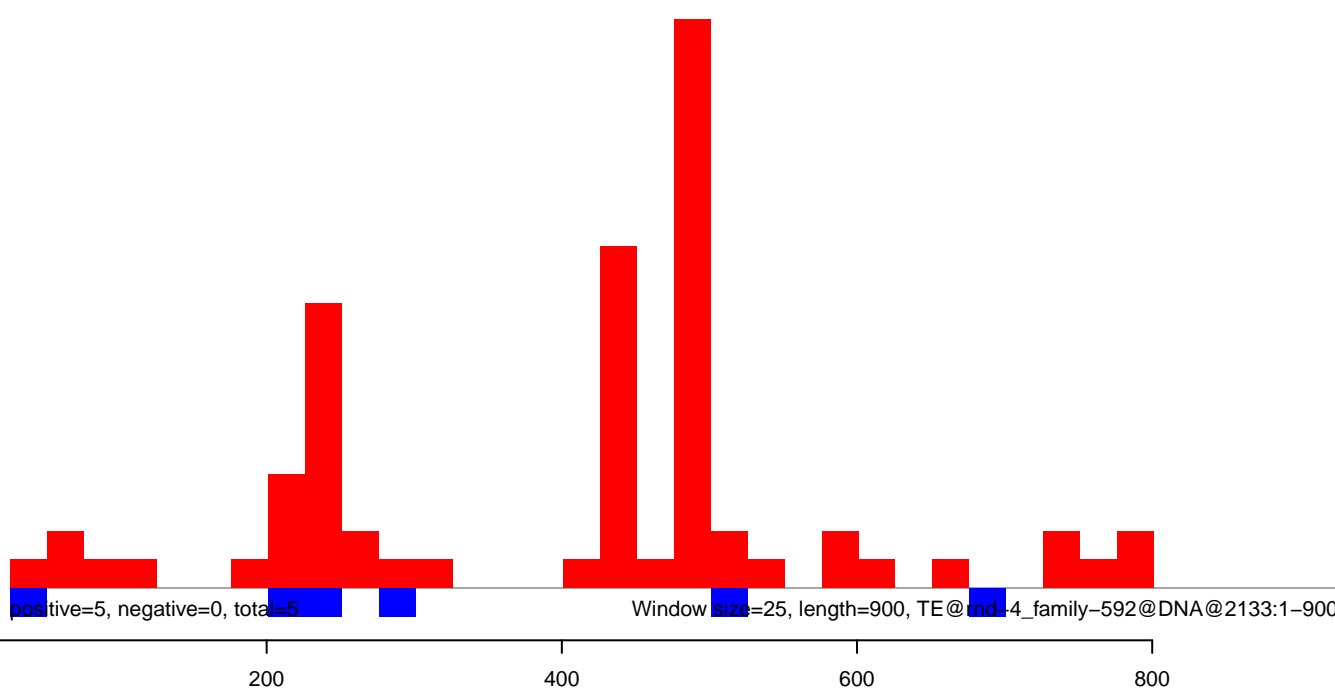
CuQuin_Mcarc_NL.18_23.rep



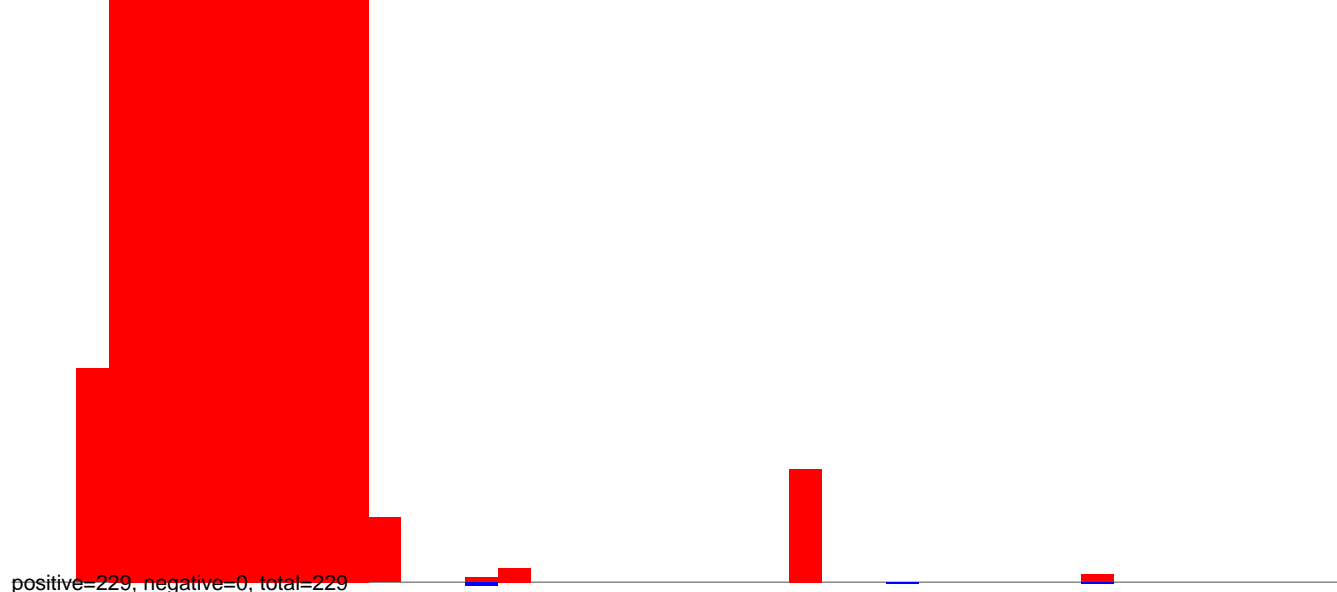
CuQuin_Mcarc_NL.24_35.rep



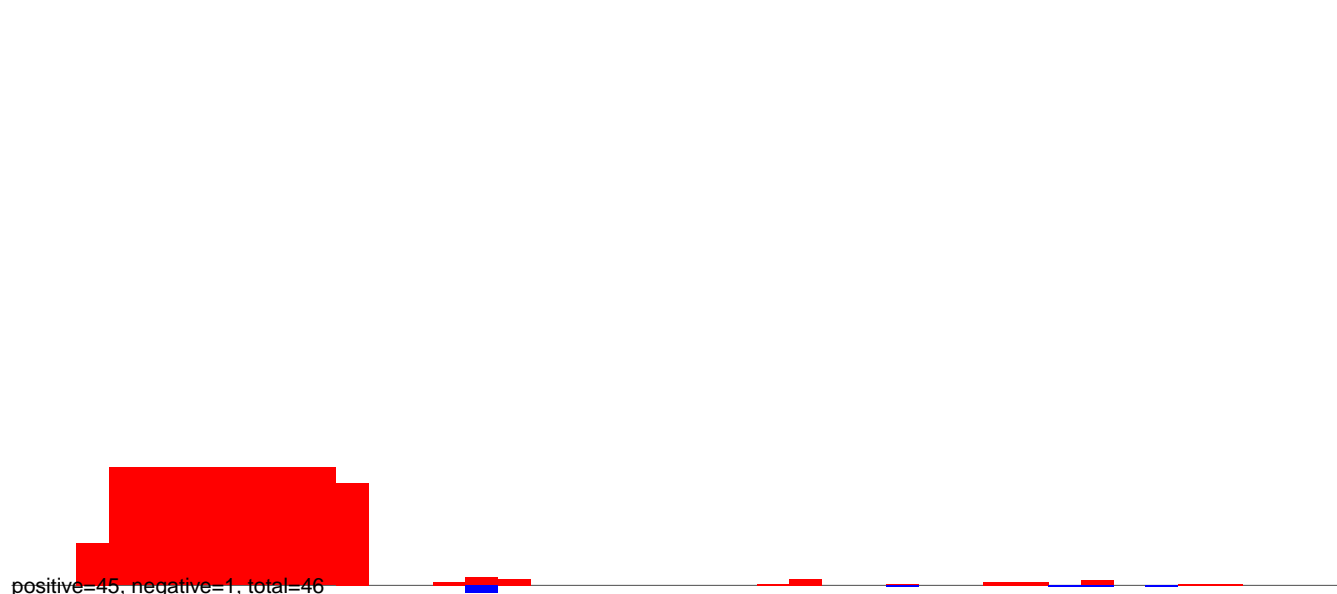
CuQuin_Mcarc_NL.rep



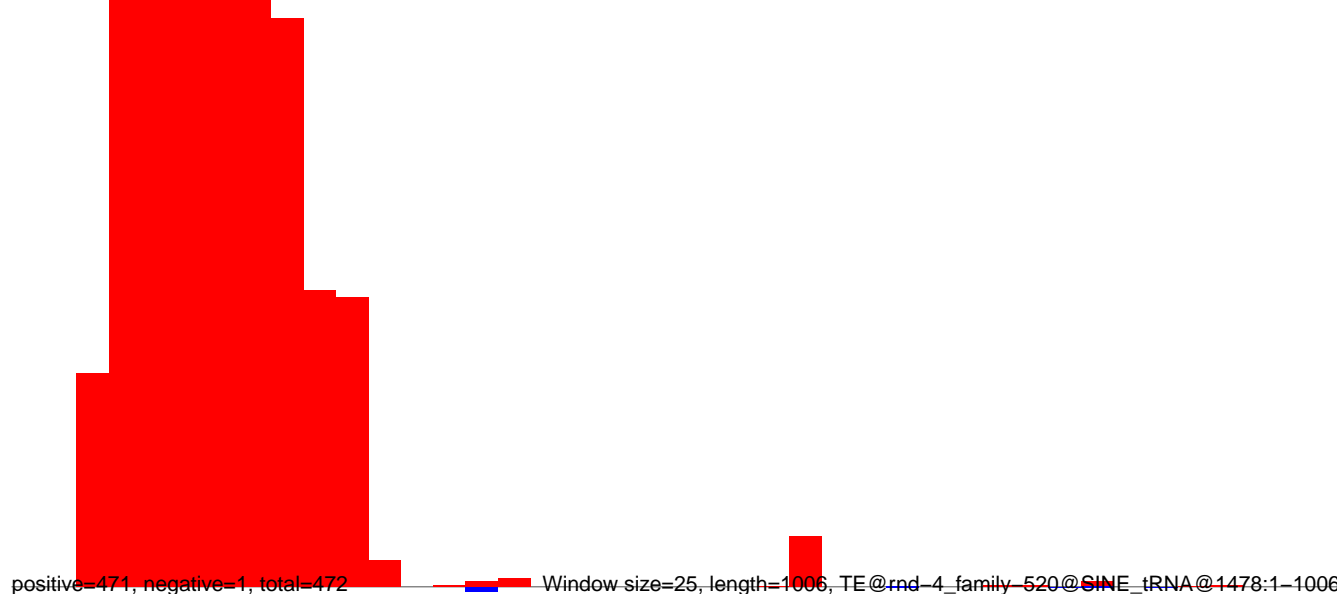
CuQuin_Mcarc_NL.18_23.rep



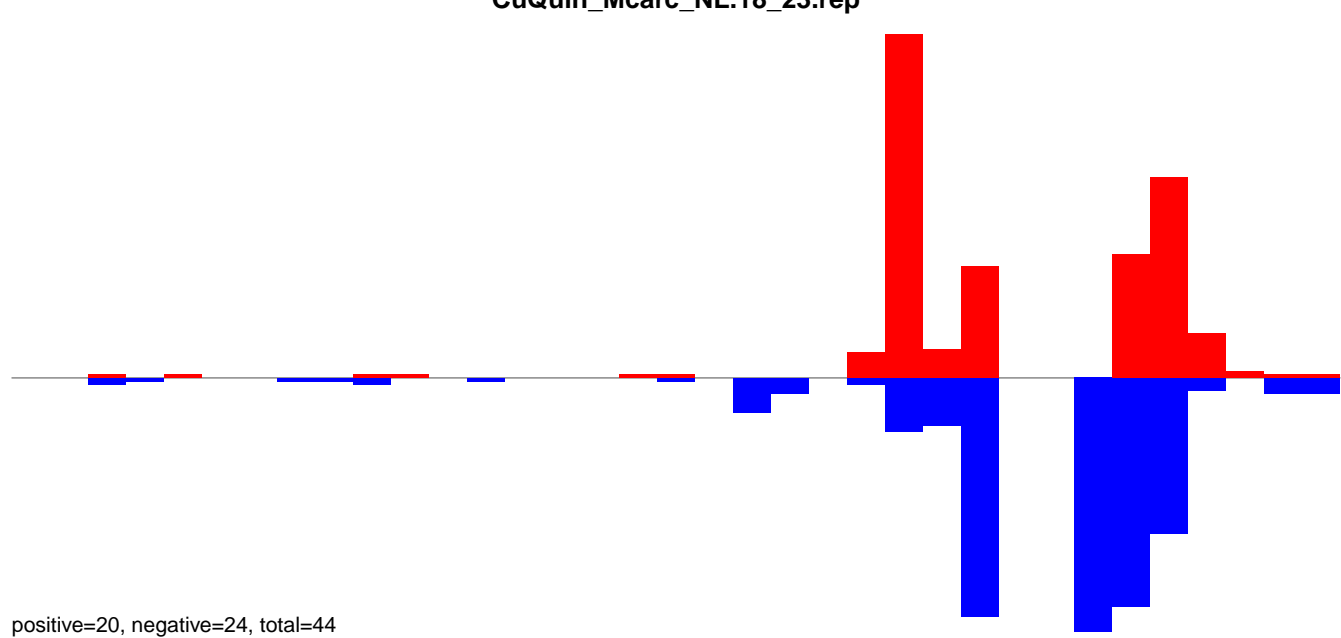
CuQuin_Mcarc_NL.24_35.rep



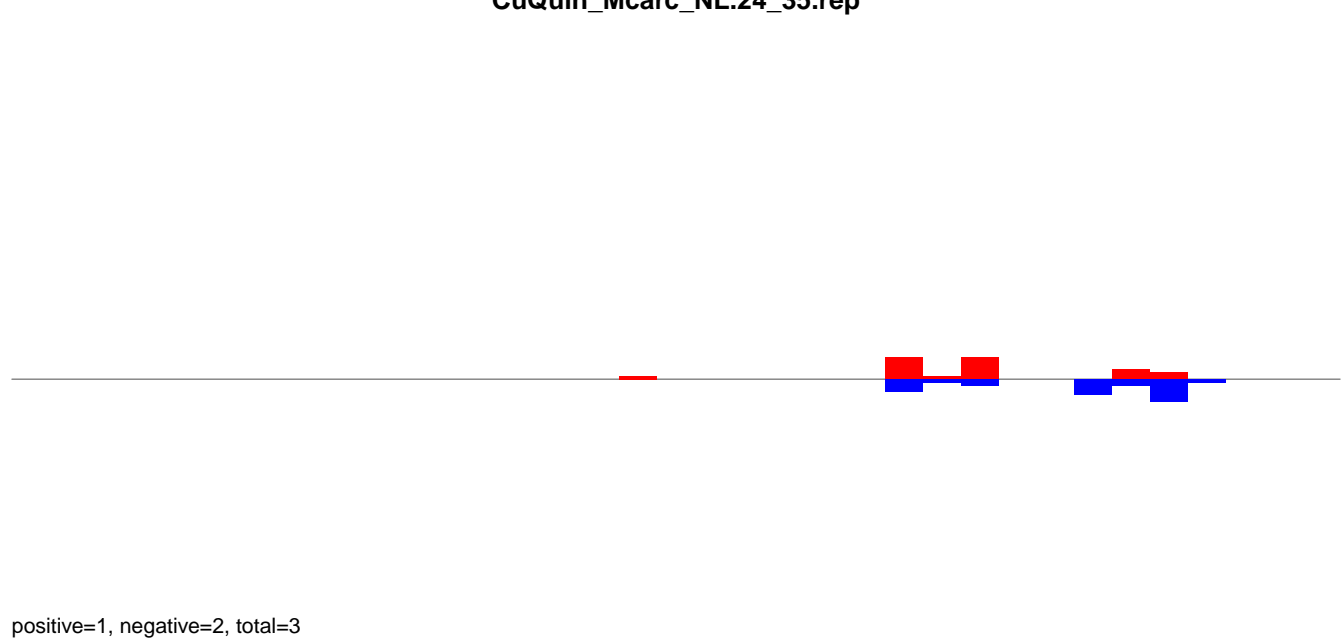
CuQuin_Mcarc_NL.rep



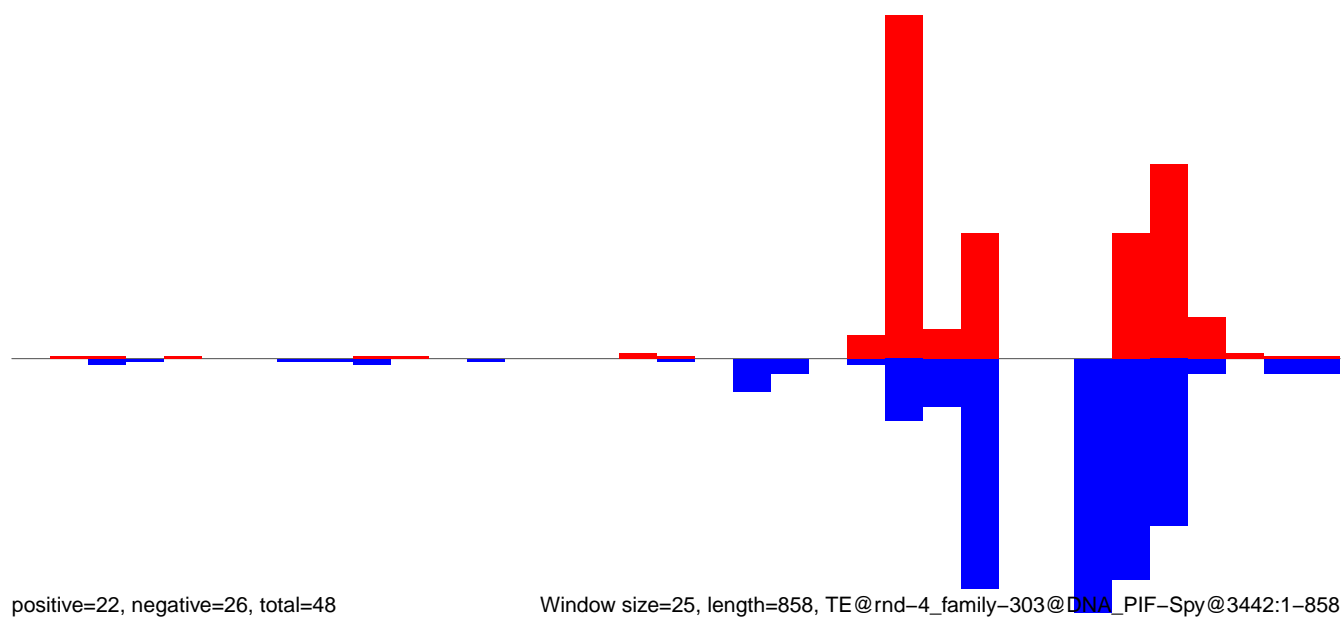
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

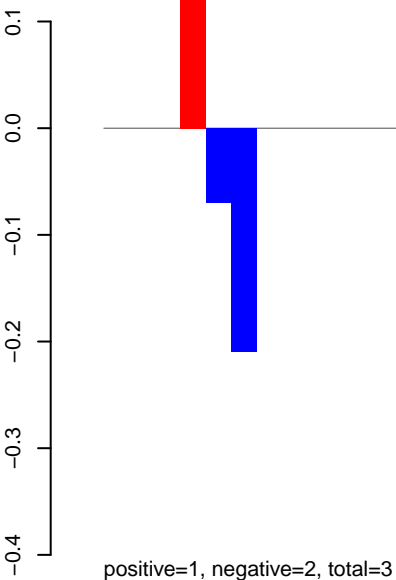


CuQuin_Mcarc_NL.rep

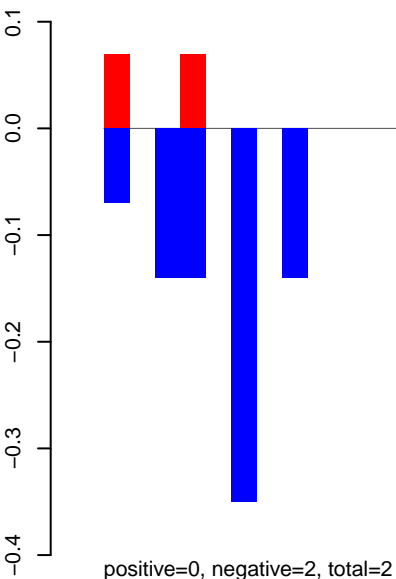


Window size=25, length=858, TE@rnd-4_family-303@DNA_PIF-Spy@3442:1-858

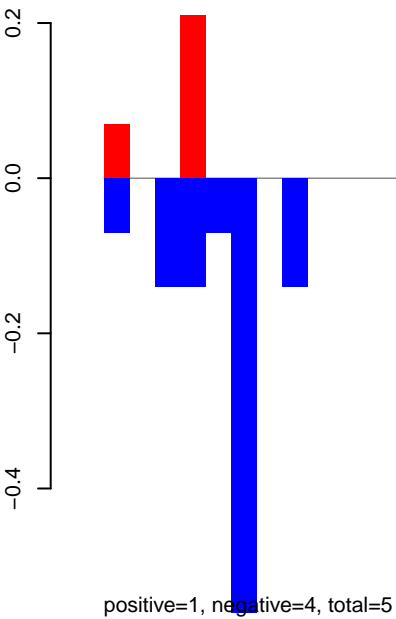
CuQuin_Mcarc_NL.18_23.rep



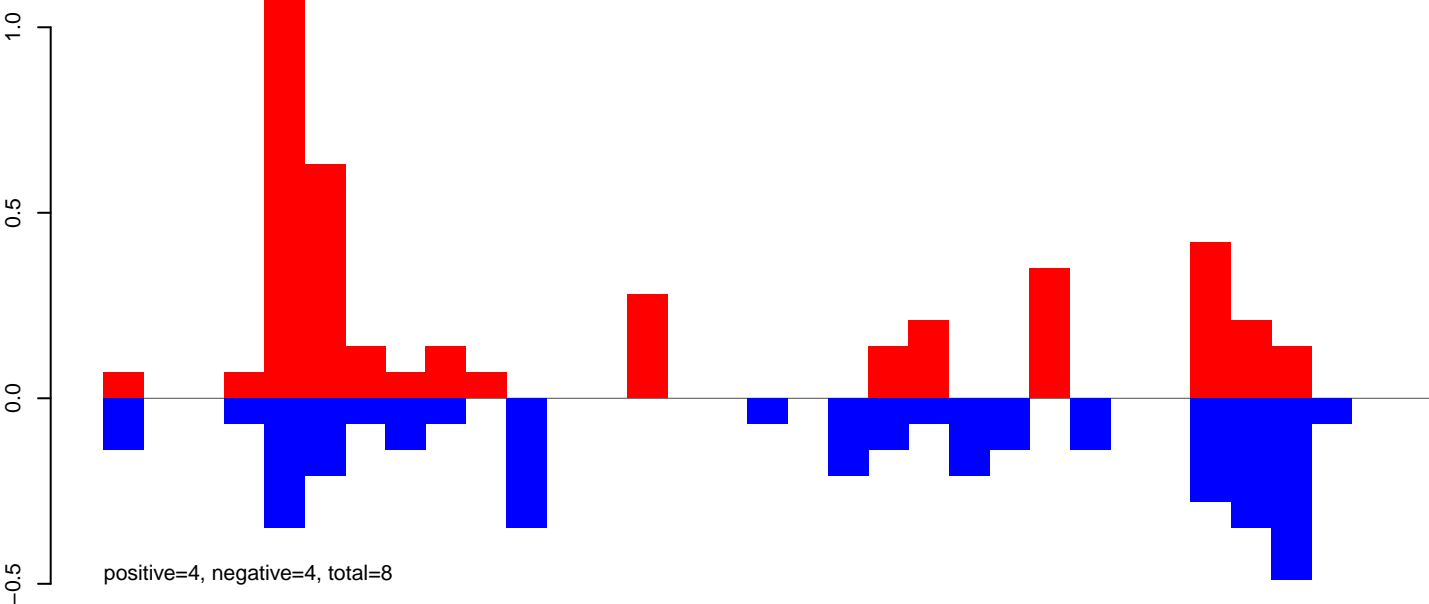
CuQuin_Mcarc_NL.24_35.rep



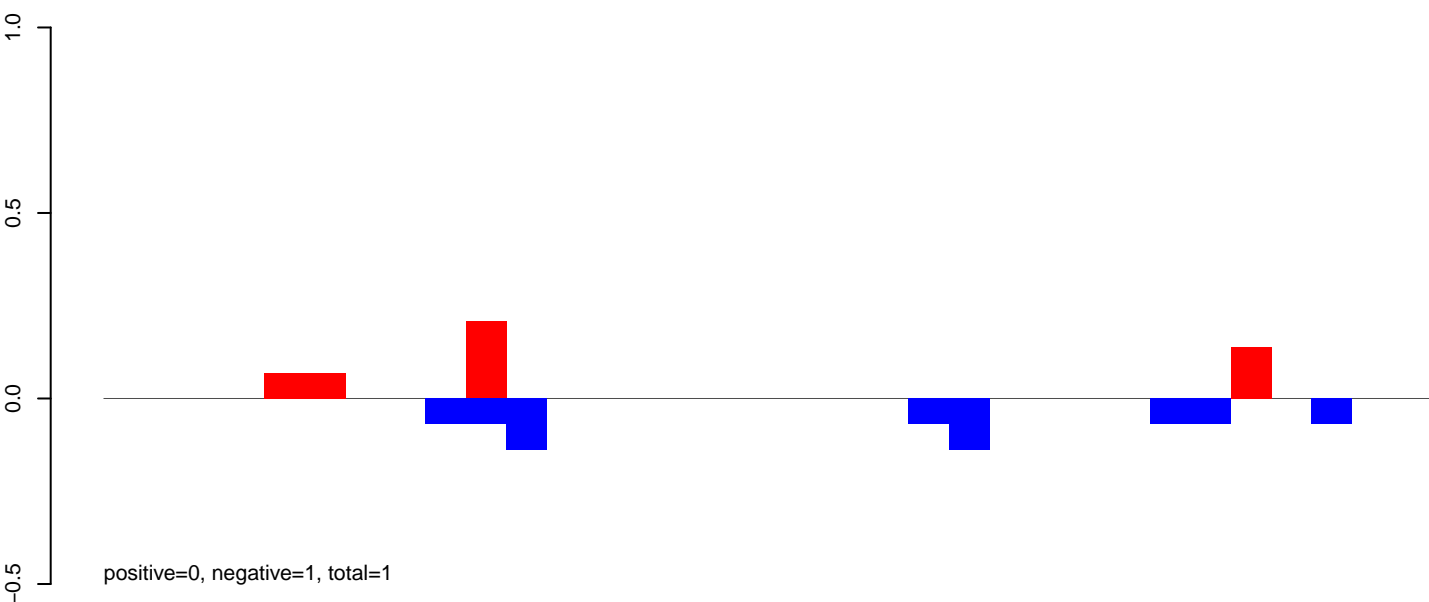
CuQuin_Mcarc_NL.rep



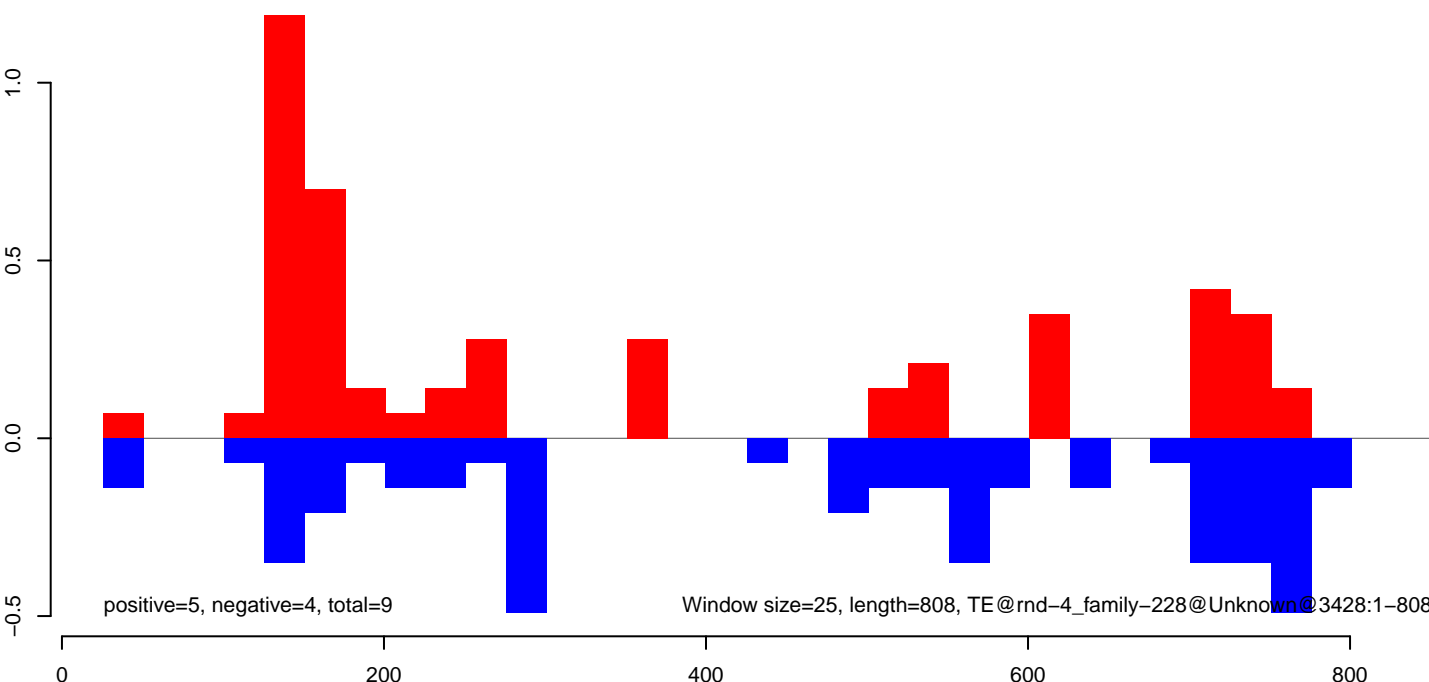
CuQuin_Mcarc_NL.18_23.rep



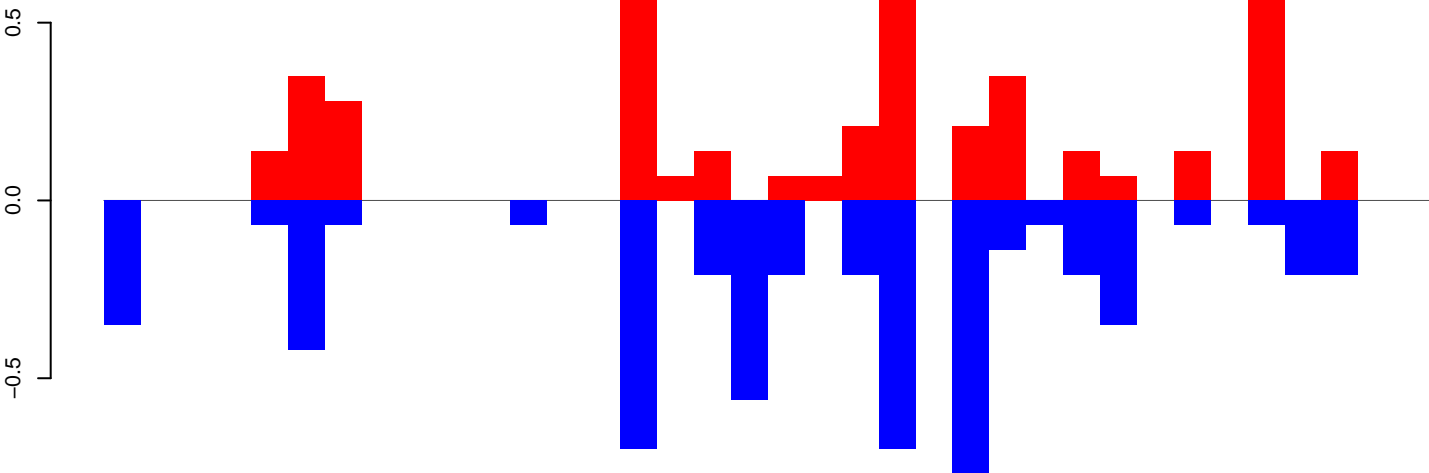
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

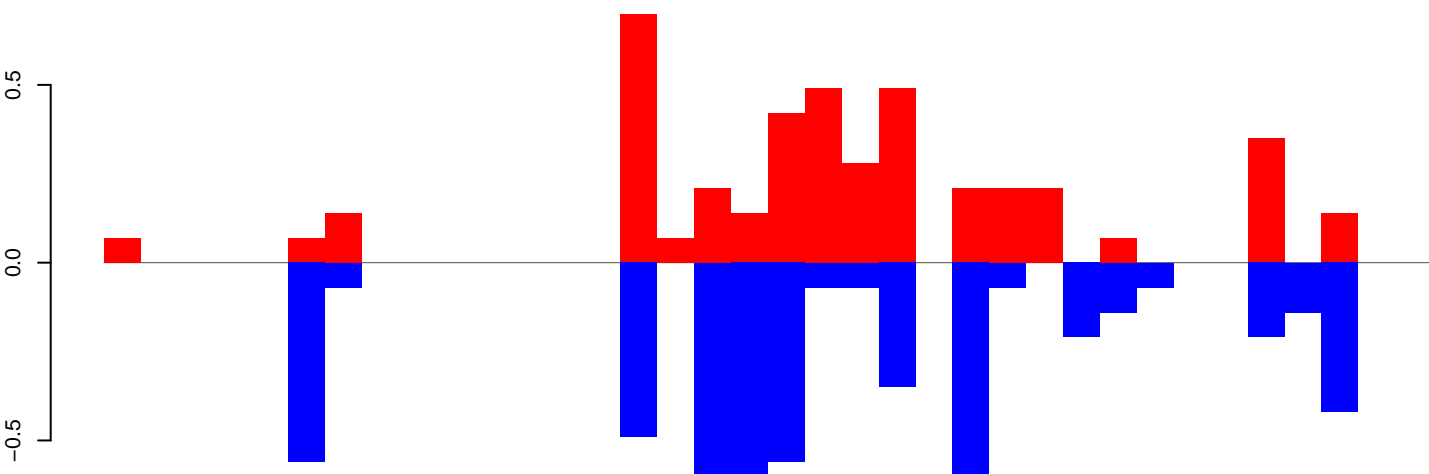


CuQuin_Mcarc_NL.18_23.rep



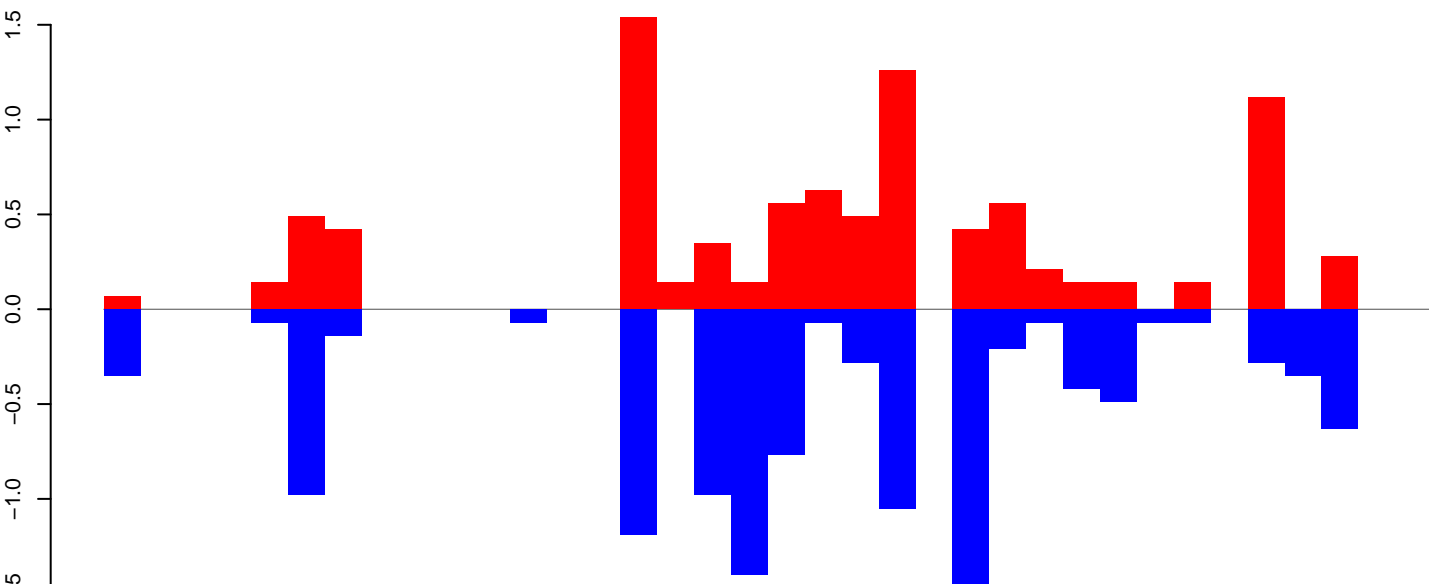
positive=5, negative=6, total=10

CuQuin_Mcarc_NL.24_35.rep



positive=4, negative=6, total=10

CuQuin_Mcarc_NL.rep



positive=9, negative=12, total=21

Window size=25, length=878, TE@rnd-4_family-151@Unknown@2786:1-878

0 200 400 600 800

CuQuin_Mcarc_NL.18_23.rep

0.3
0.2
0.1
0.0
-0.1
-0.2
-0.3

positive=2, negative=2, total=4

CuQuin_Mcarc_NL.24_35.rep

0.3
0.2
0.1
0.0
-0.1
-0.2
-0.3

positive=2, negative=2, total=3

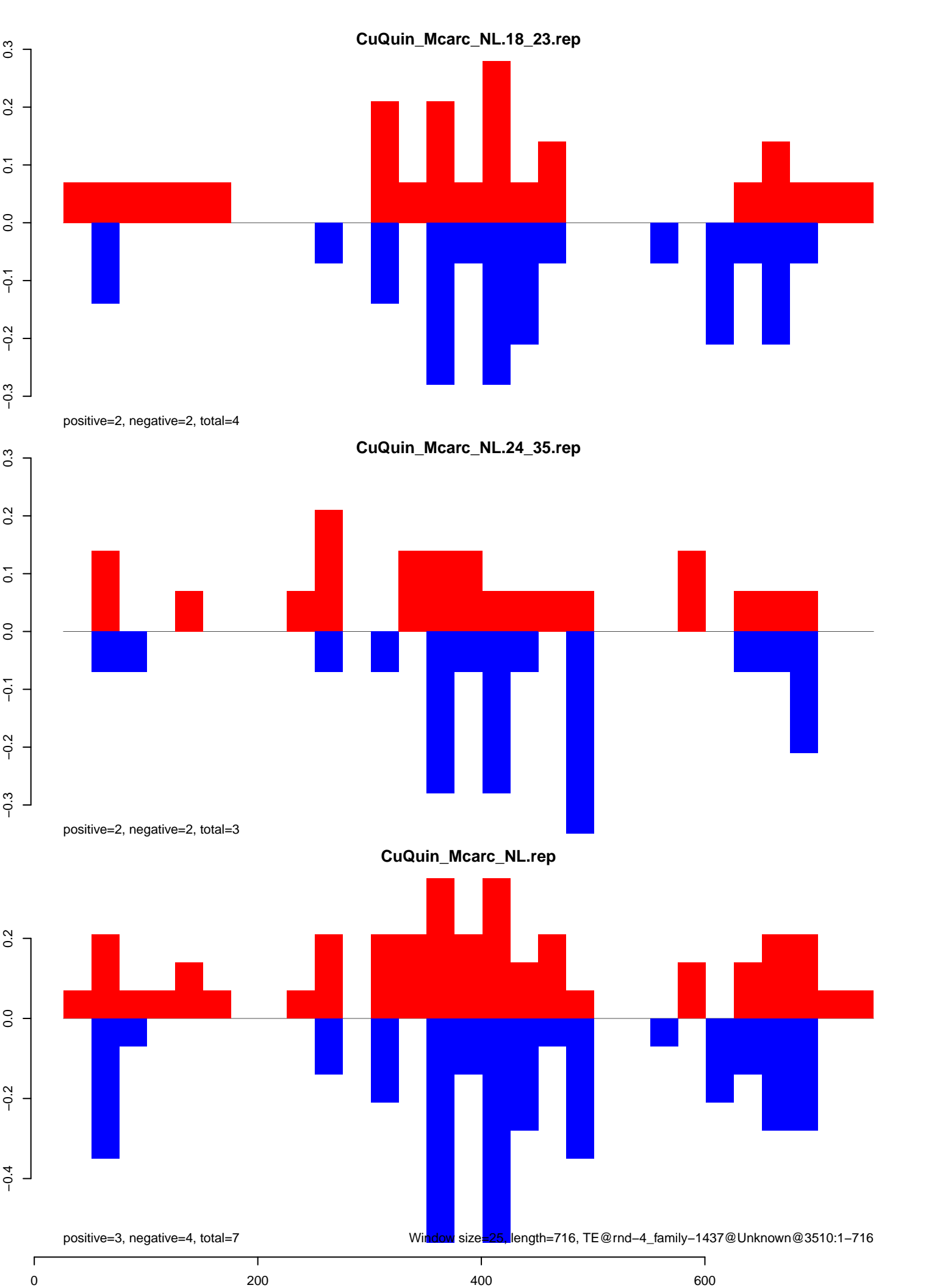
CuQuin_Mcarc_NL.rep

0.2
0.1
0.0
-0.1
-0.2
-0.3
-0.4

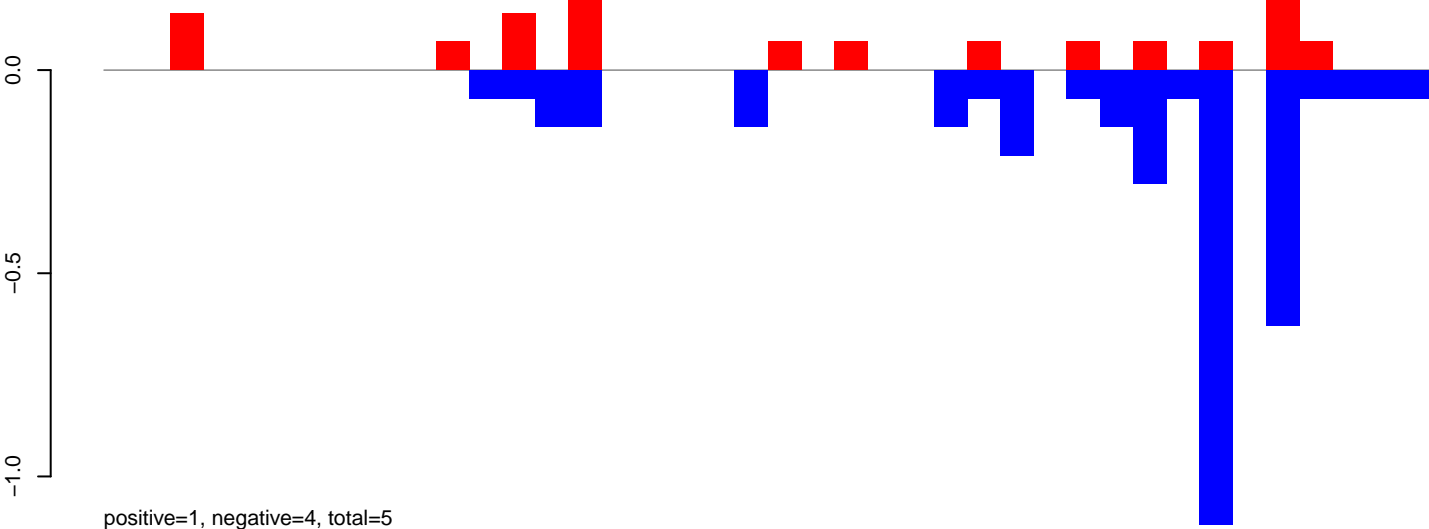
positive=3, negative=4, total=7

Window size=25, length=716, TE@rnd-4_family-1437@Unknown@3510:1-716

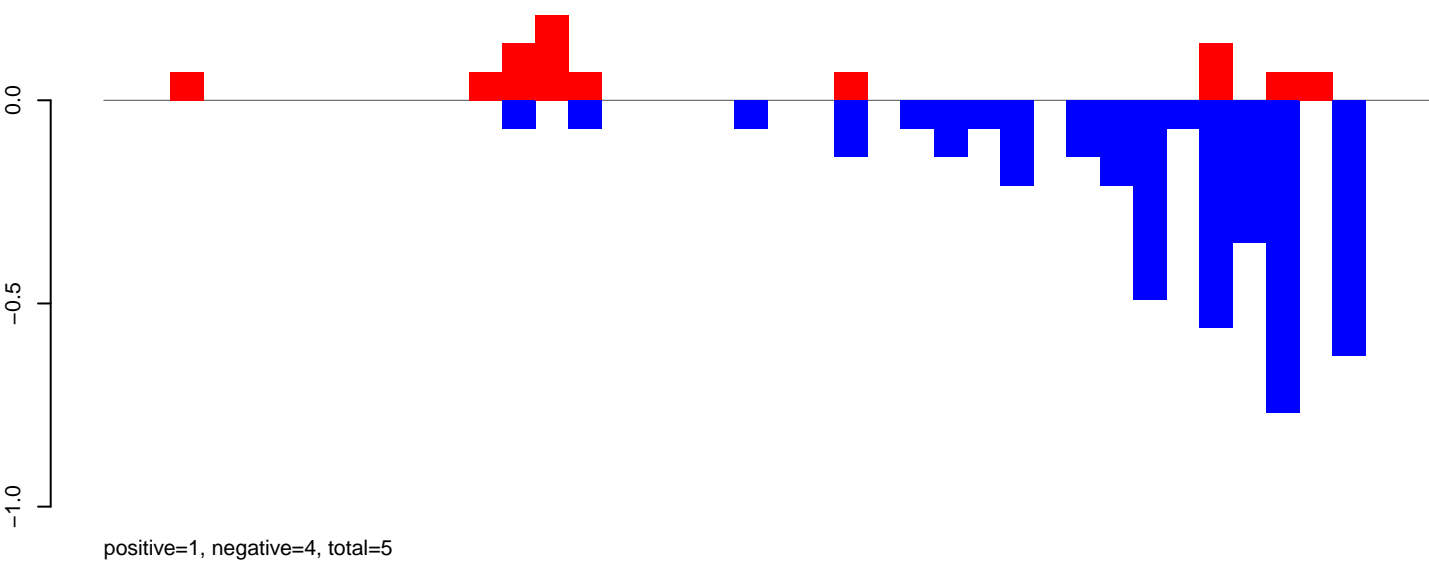
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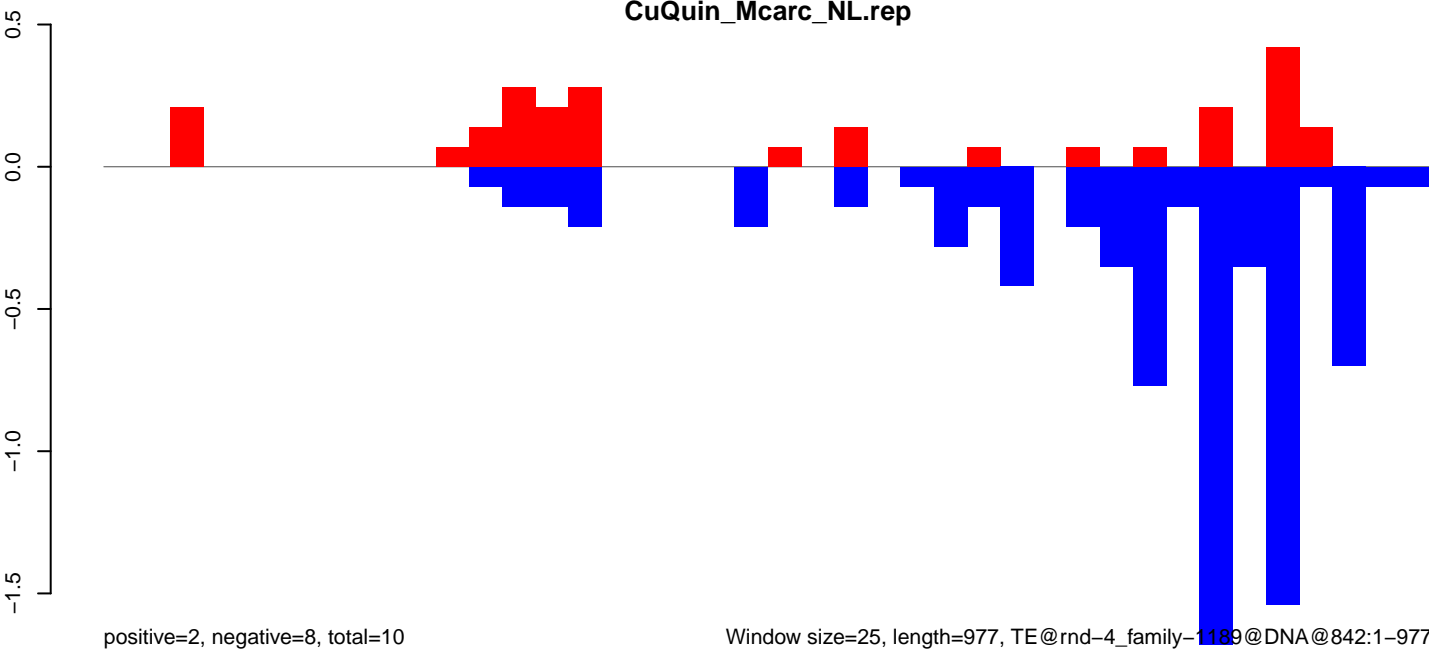
CuQuin_Mcarc_NL.18_23.rep



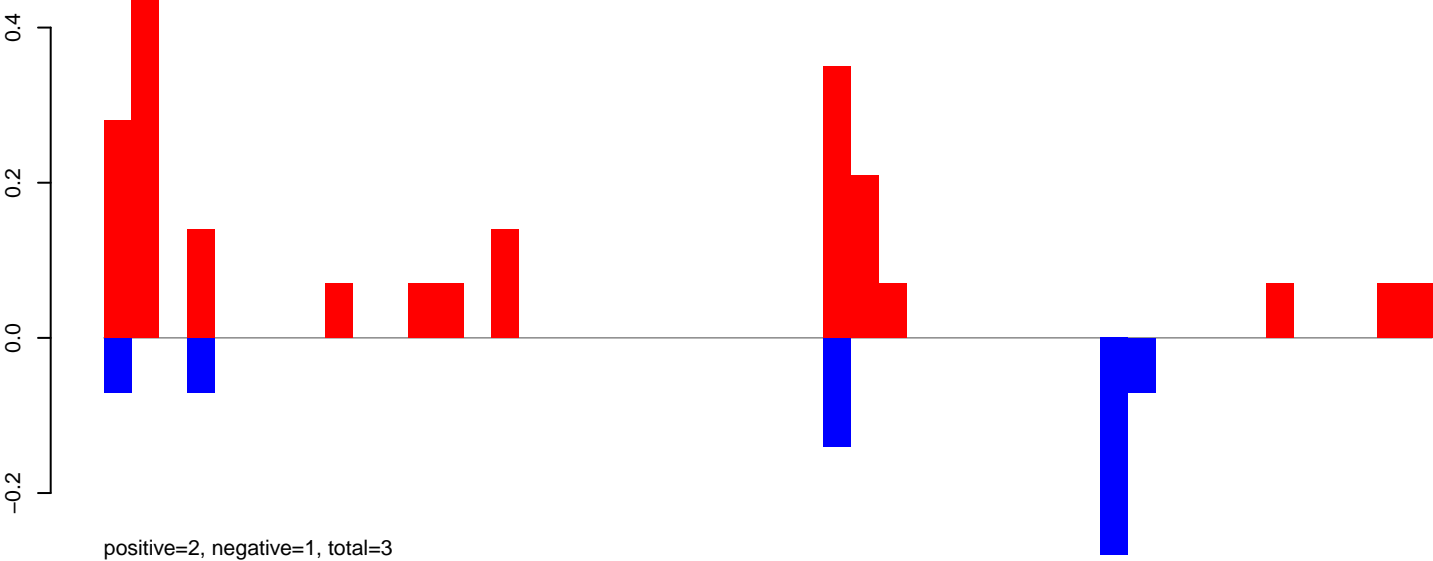
CuQuin_Mcarc_NL.24_35.rep



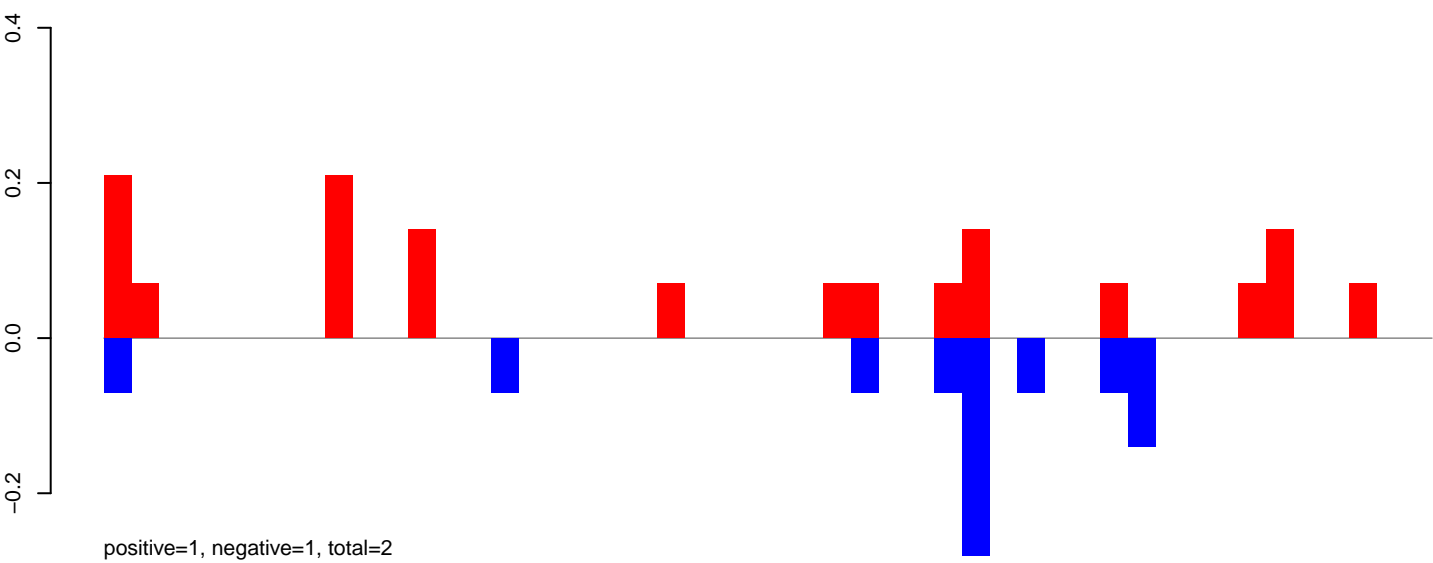
CuQuin_Mcarc_NL.rep



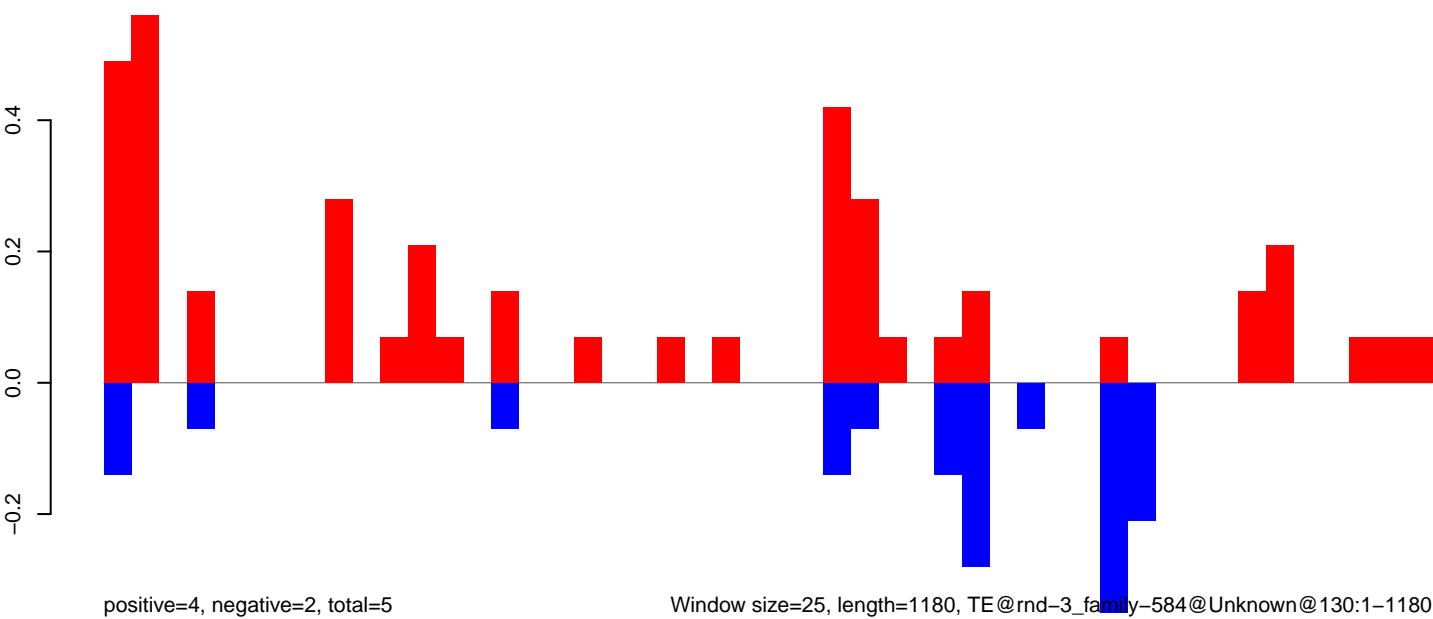
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

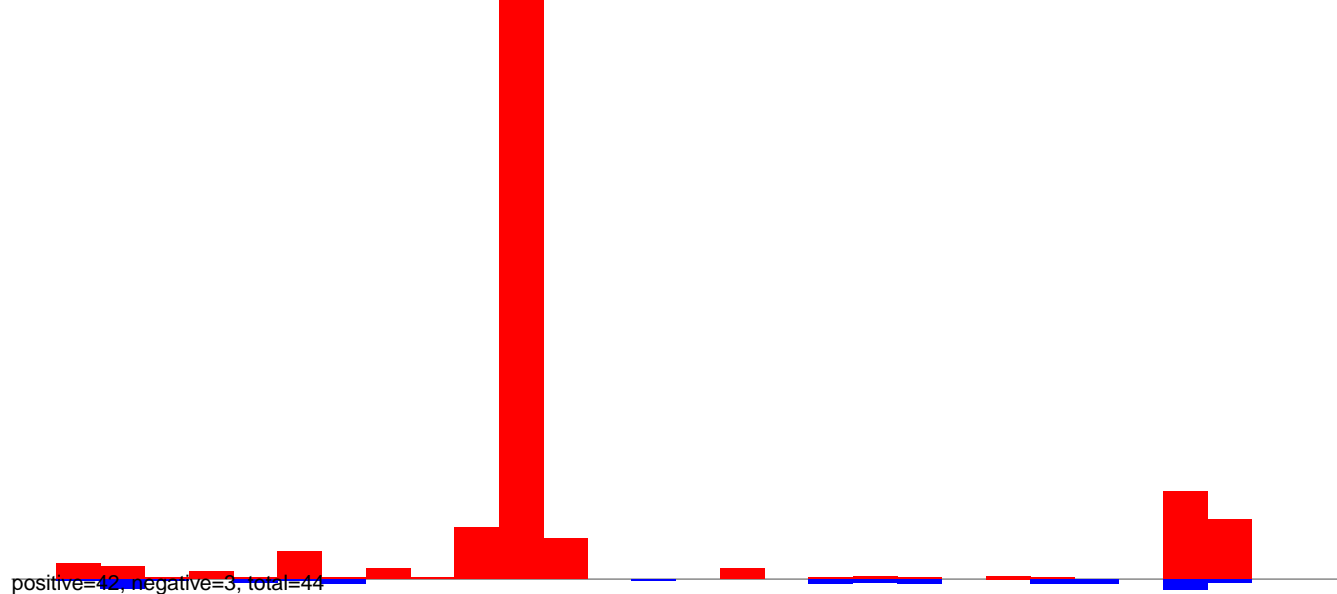


CuQuin_Mcarc_NL.rep

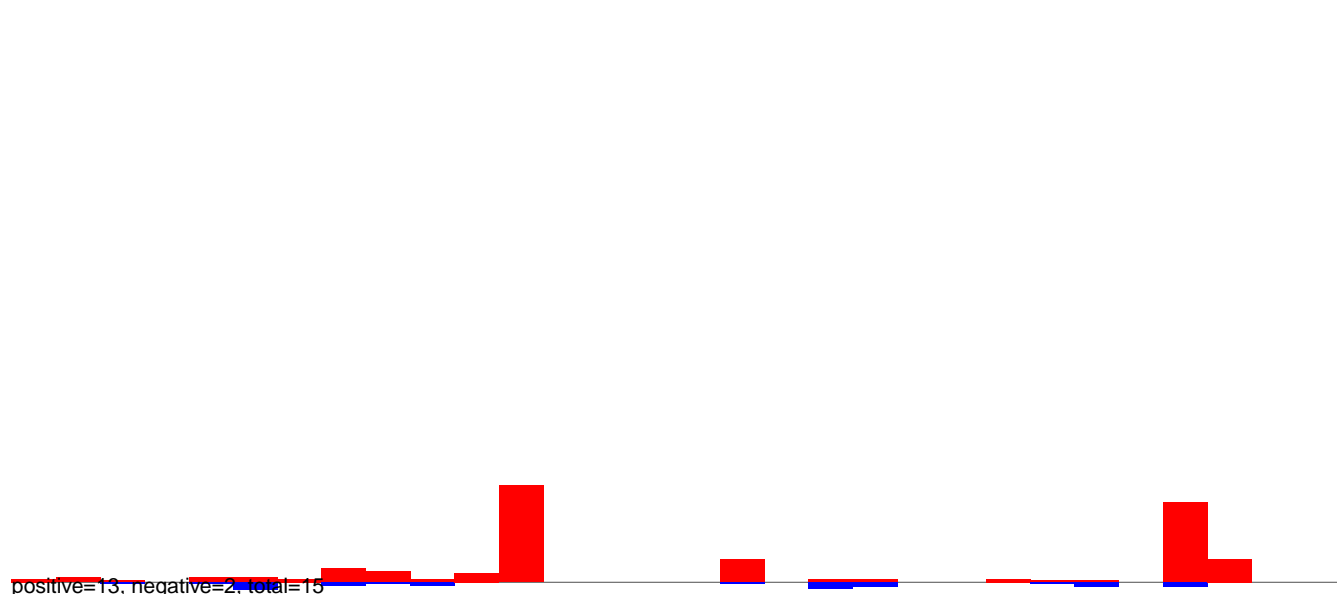


Window size=25, length=1180, TE@rnd-3_family-584@Unknown@130:1-1180

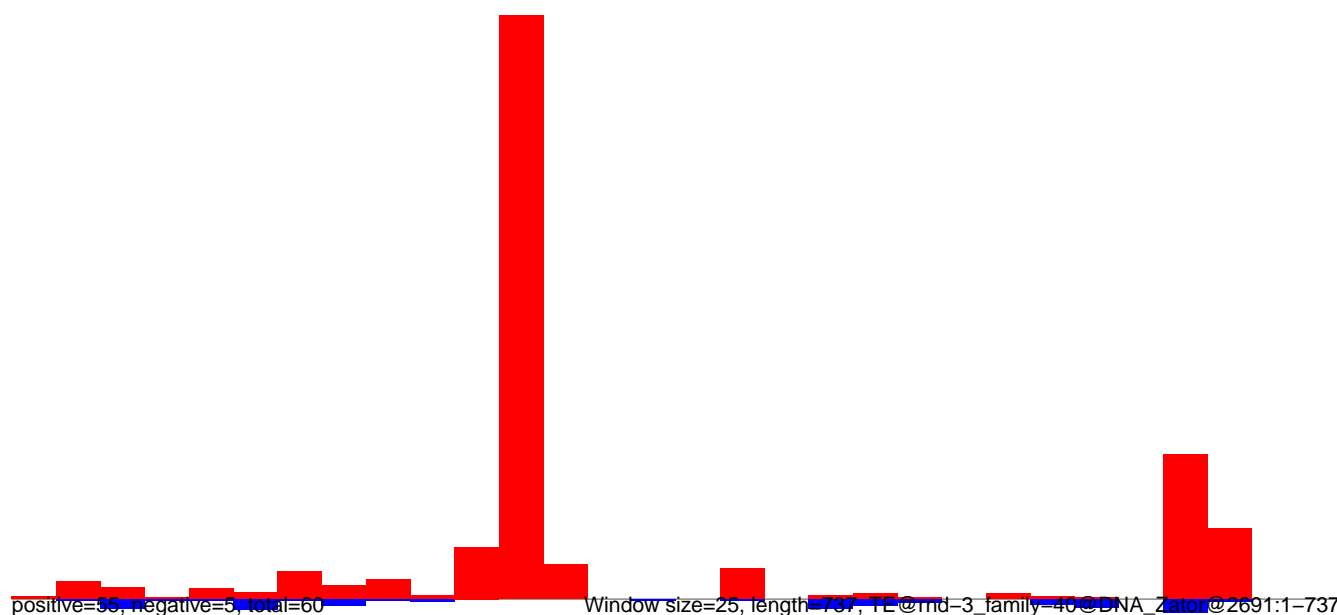
CuQuin_Mcarc_NL.18_23.rep



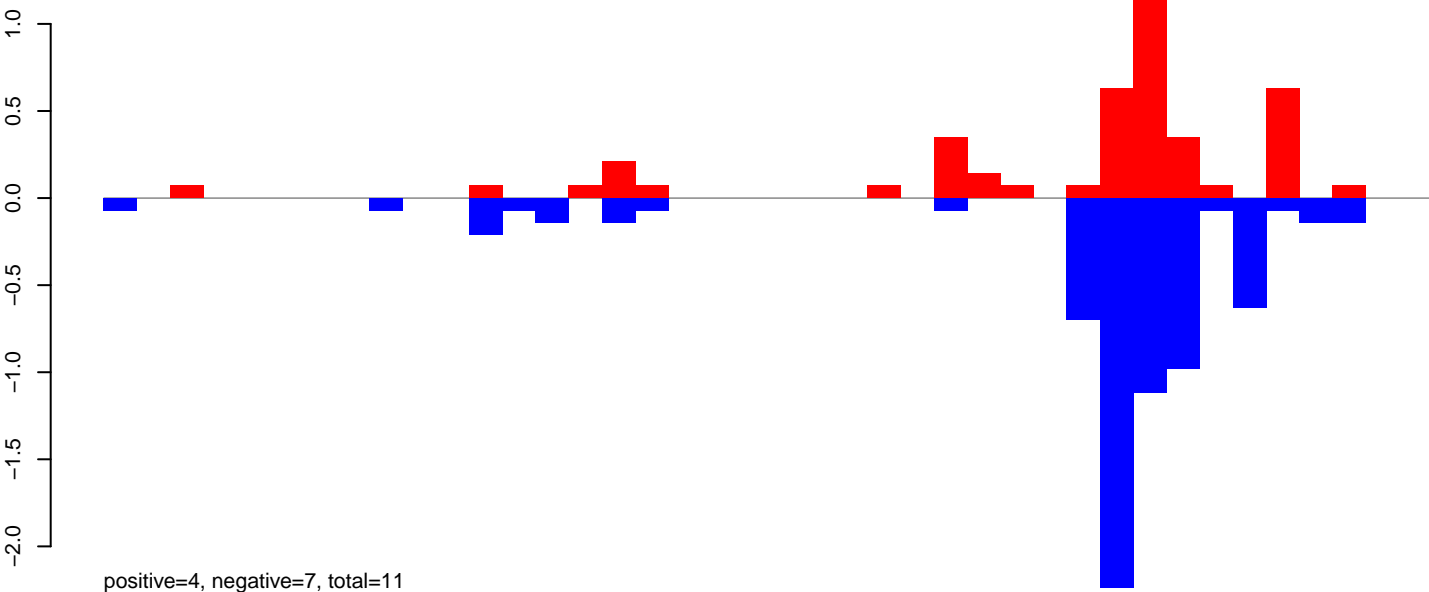
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



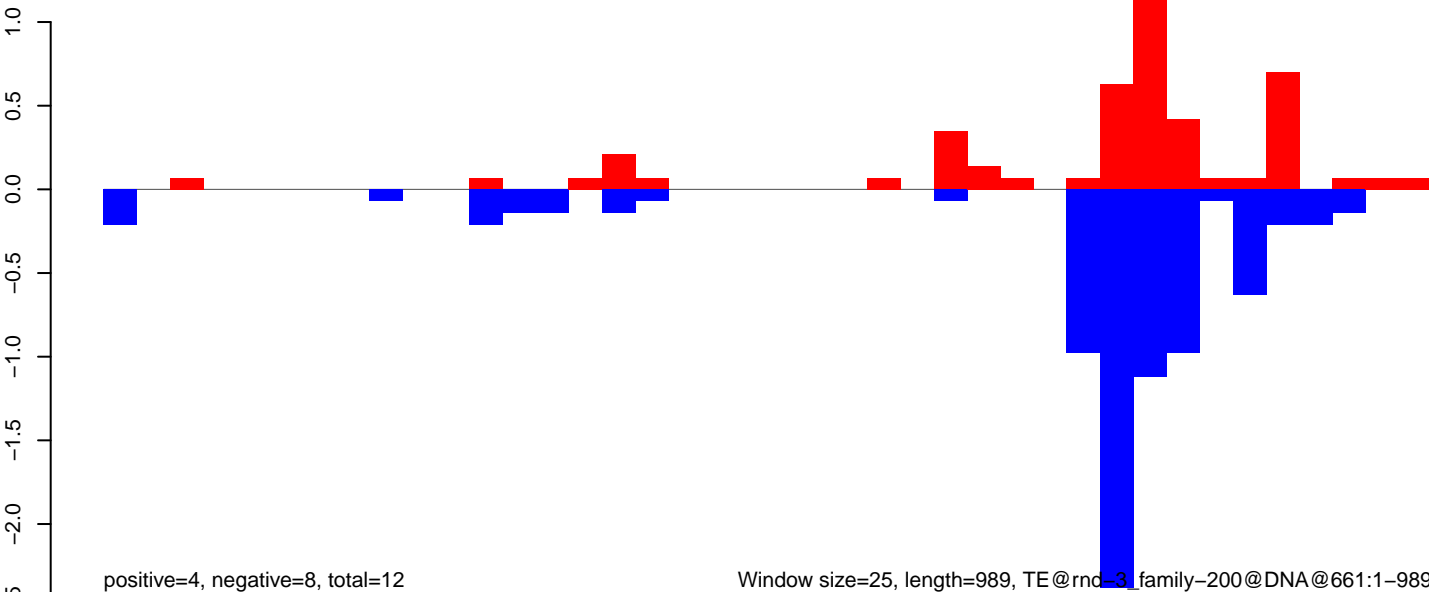
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

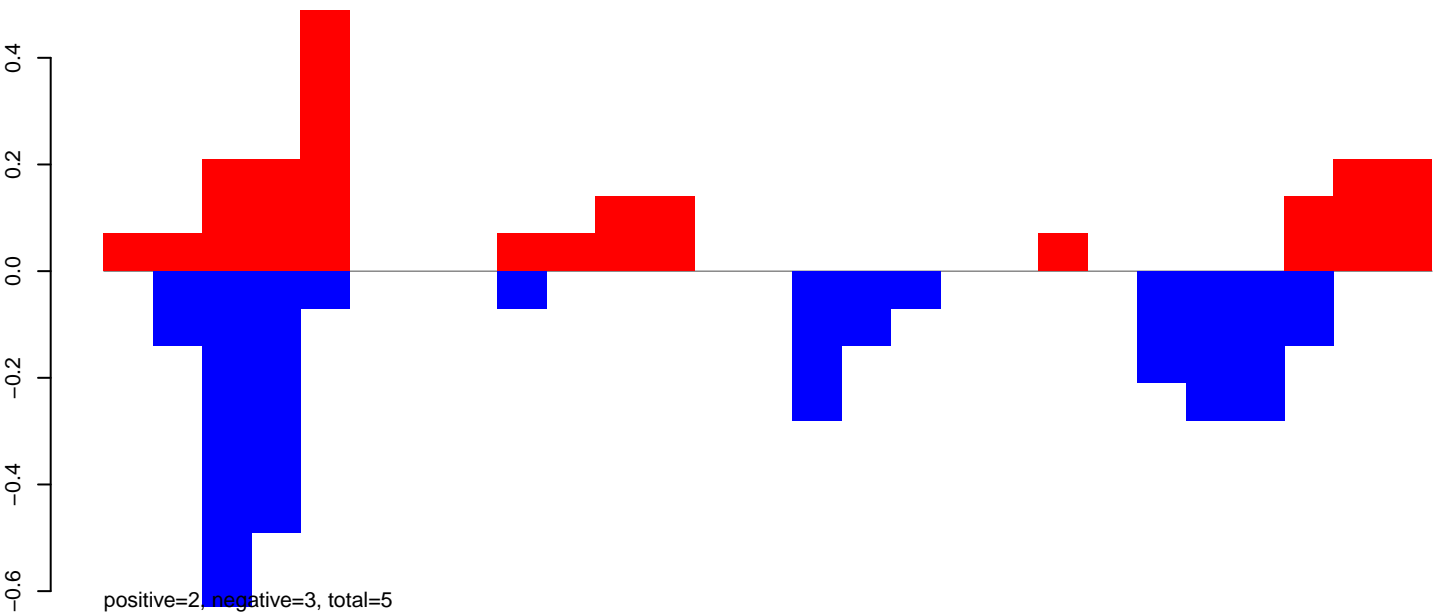


CuQuin_Mcarc_NL.rep

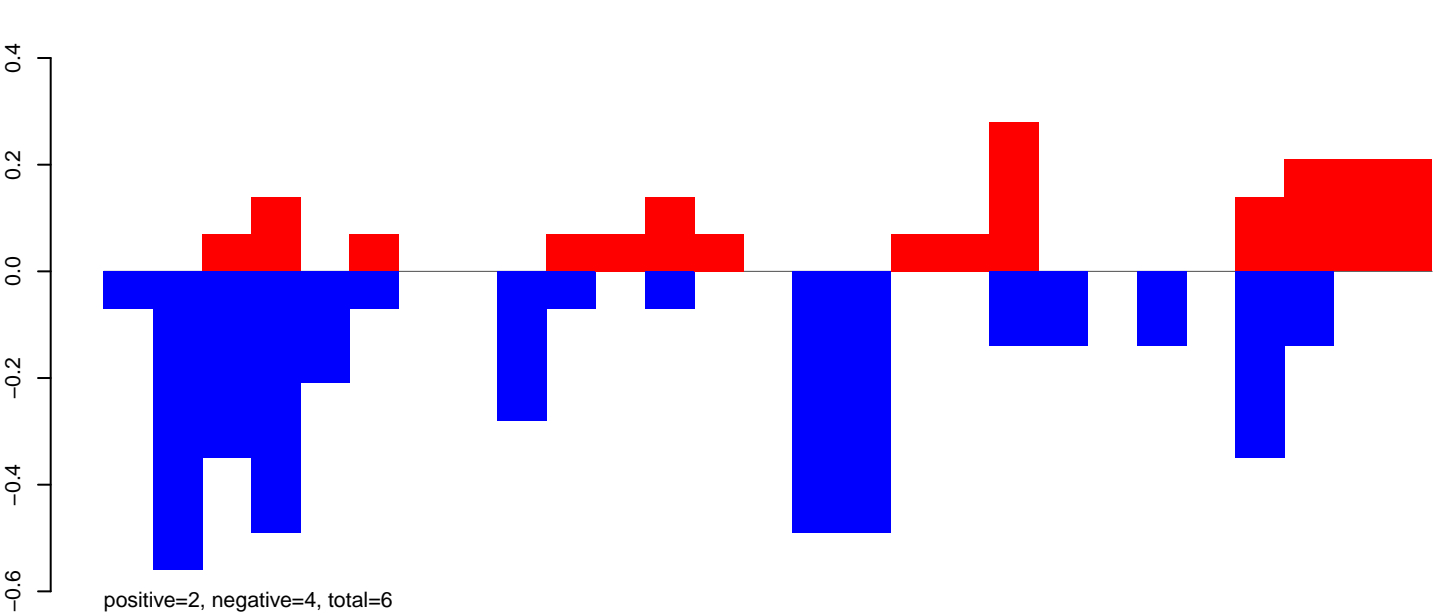


Window size=25, length=989, TE@rnd_3_family-200@DNA@661:1-989

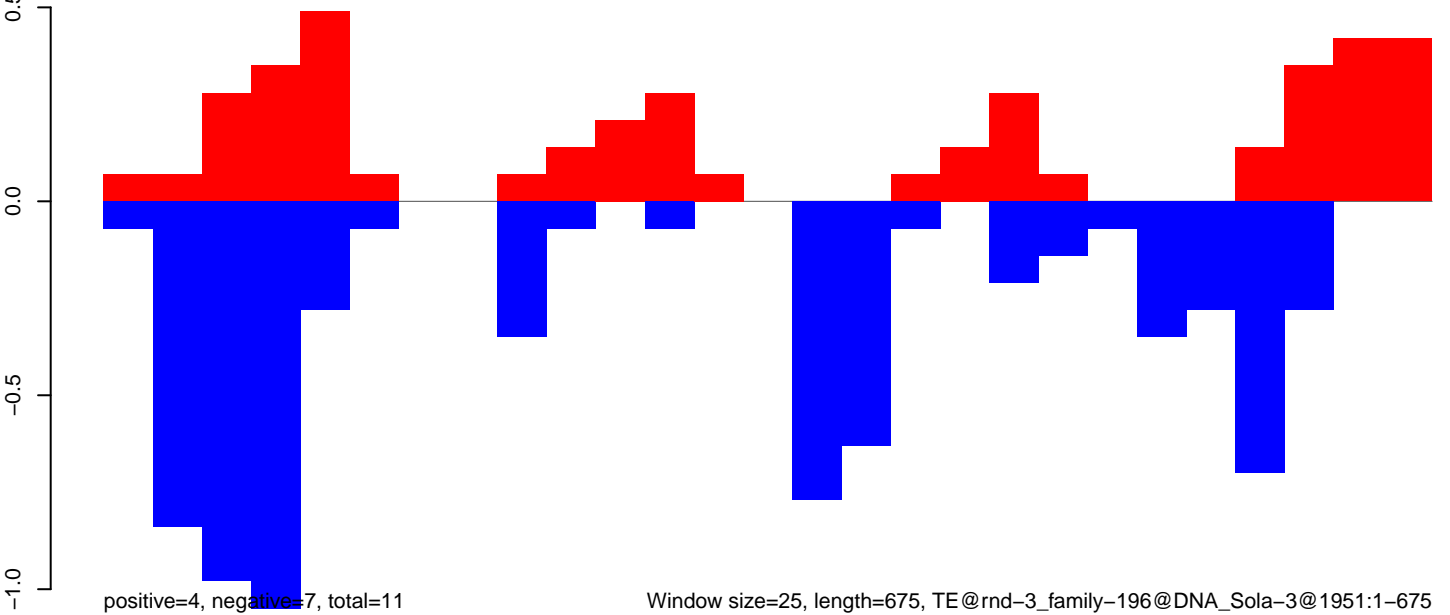
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

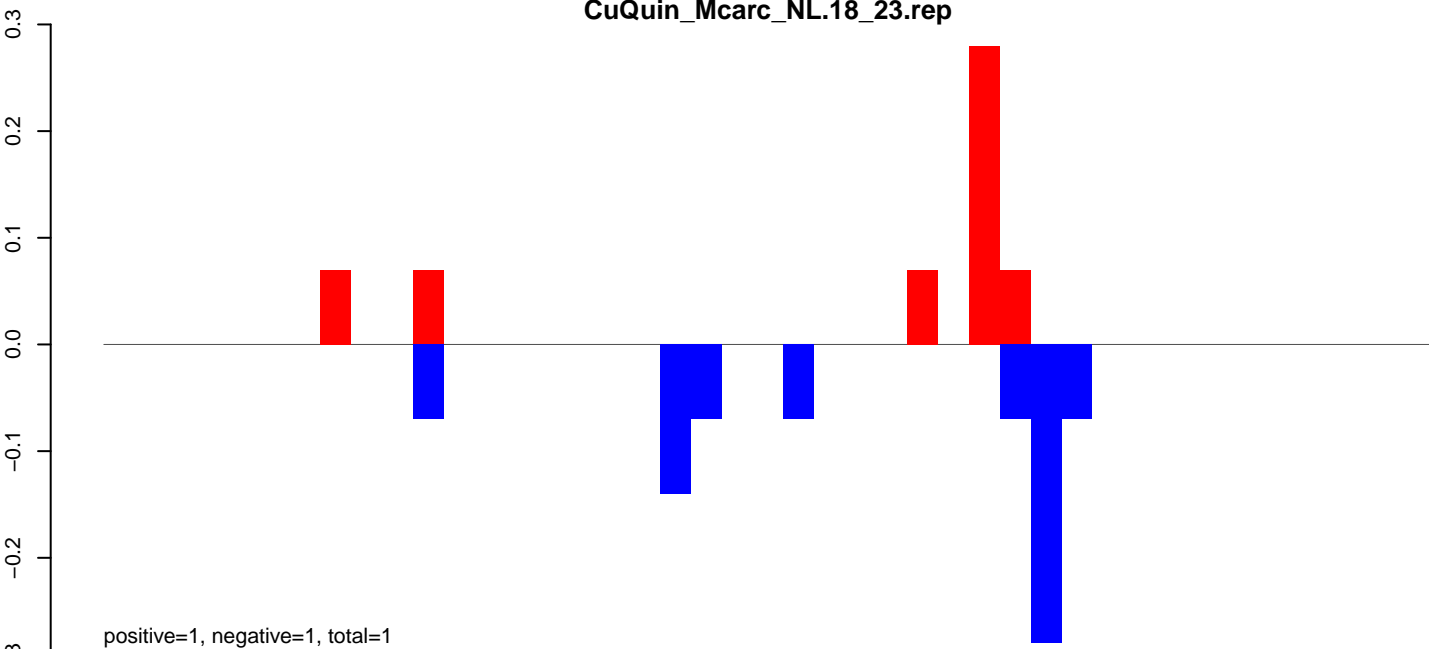


CuQuin_Mcarc_NL.rep

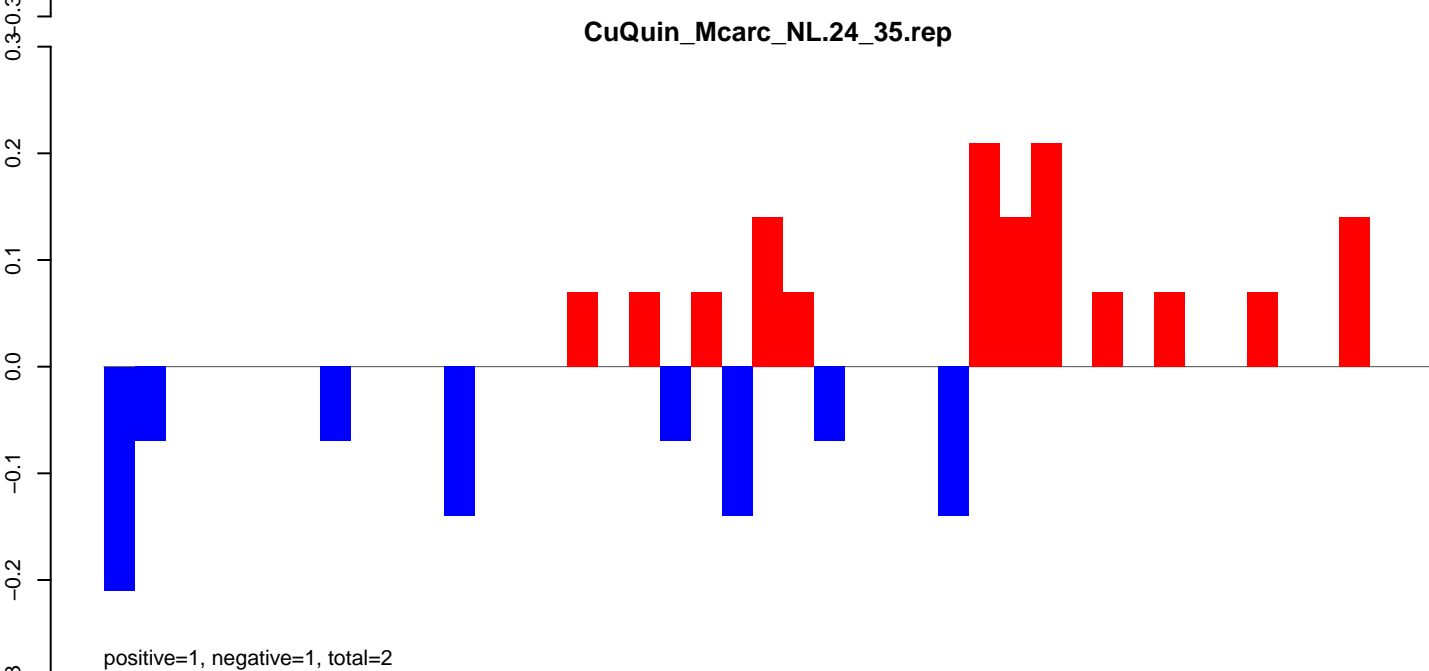


Window size=25, length=675, TE@rnd-3_family-196@DNA_Sola-3@1951:1-675

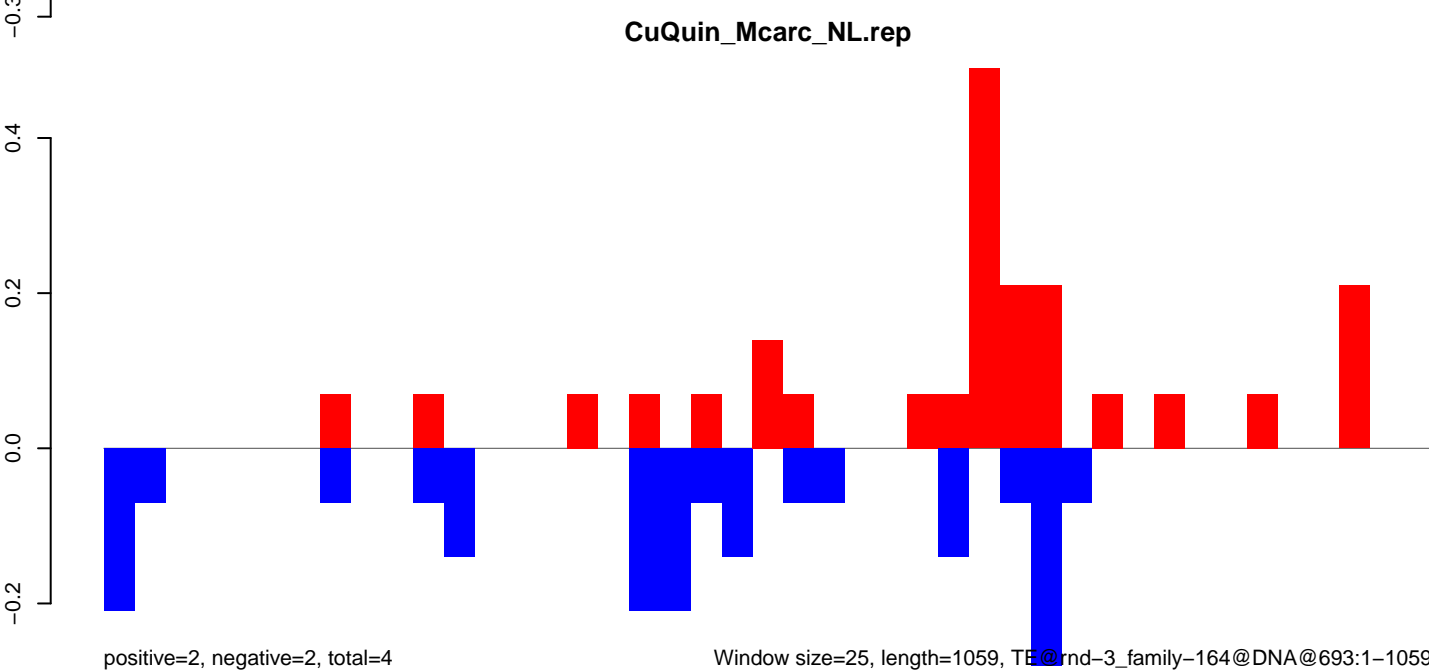
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

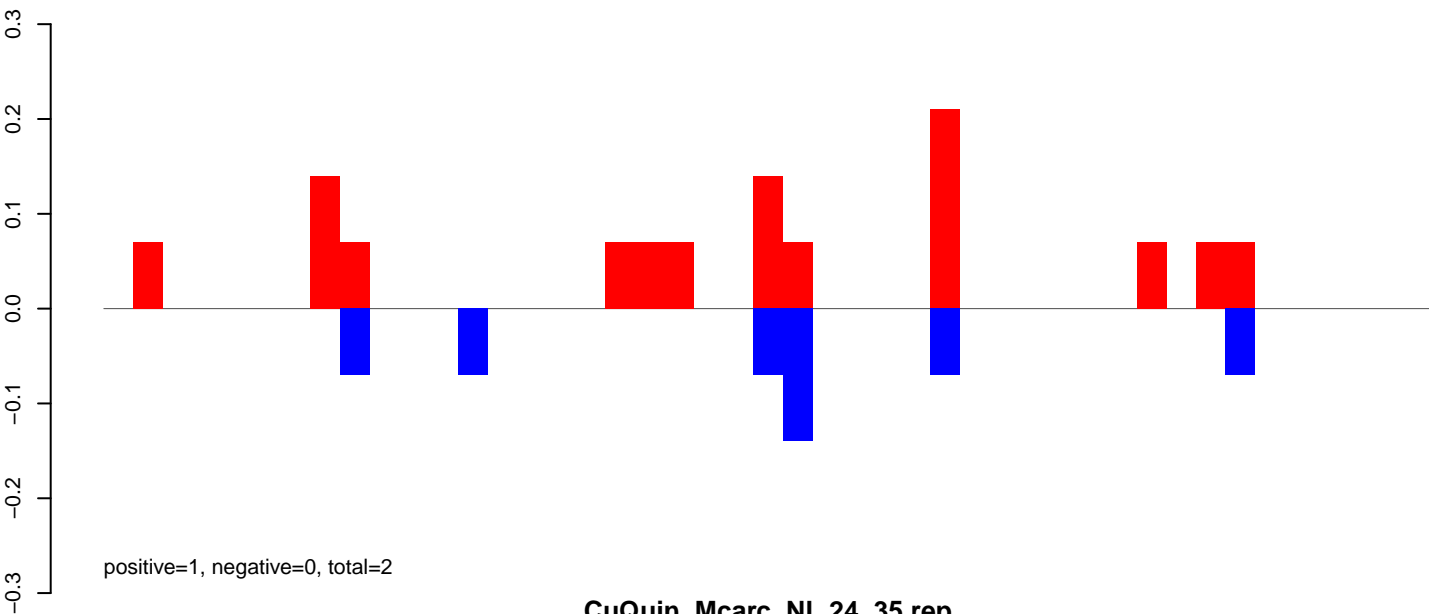


CuQuin_Mcarc_NL.rep

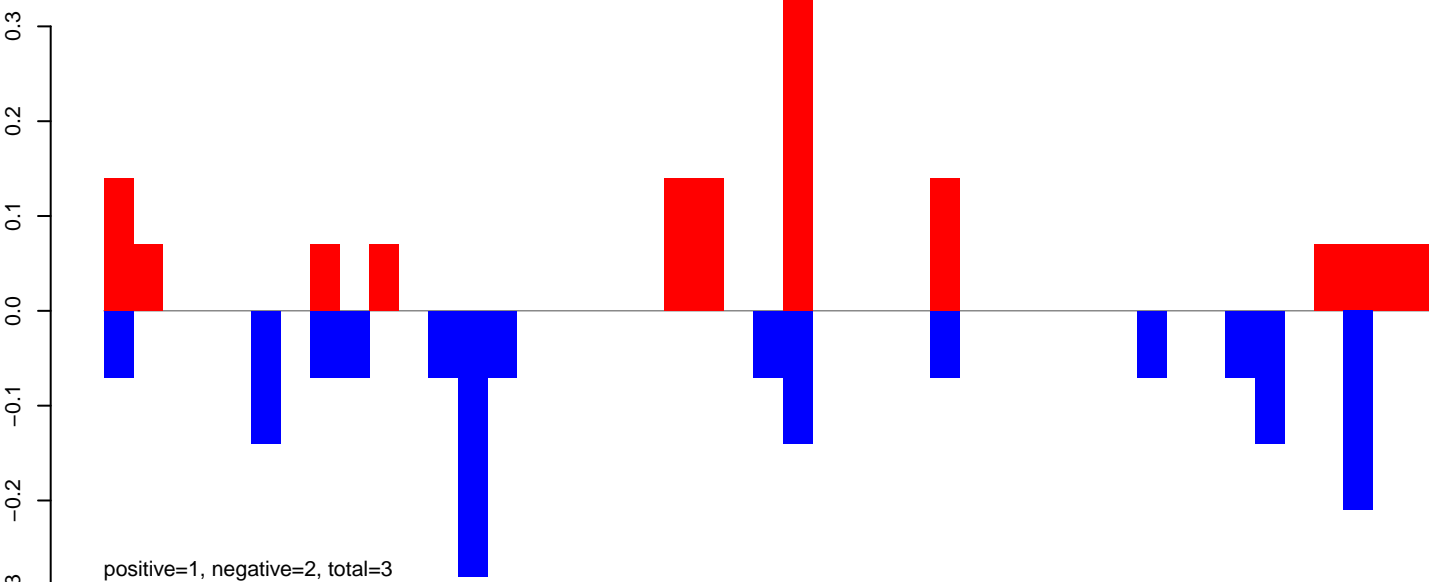


Window size=25, length=1059, TE@rnd-3_family-164@DNA@693:1-1059

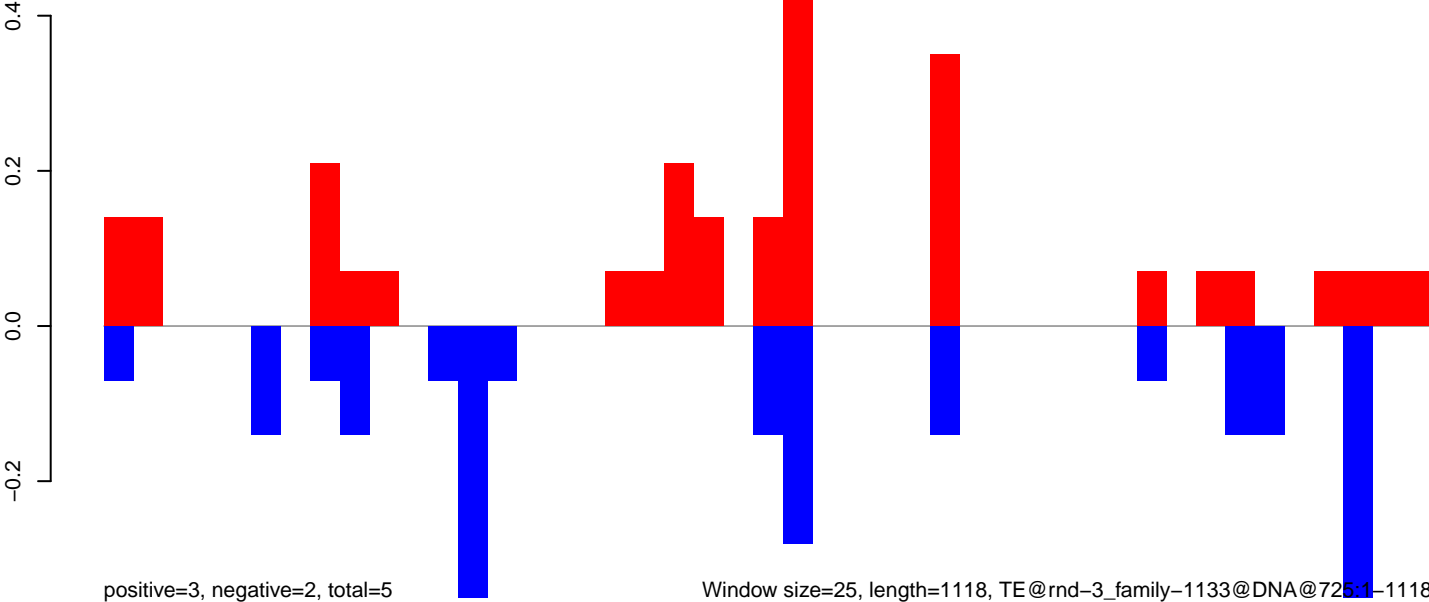
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

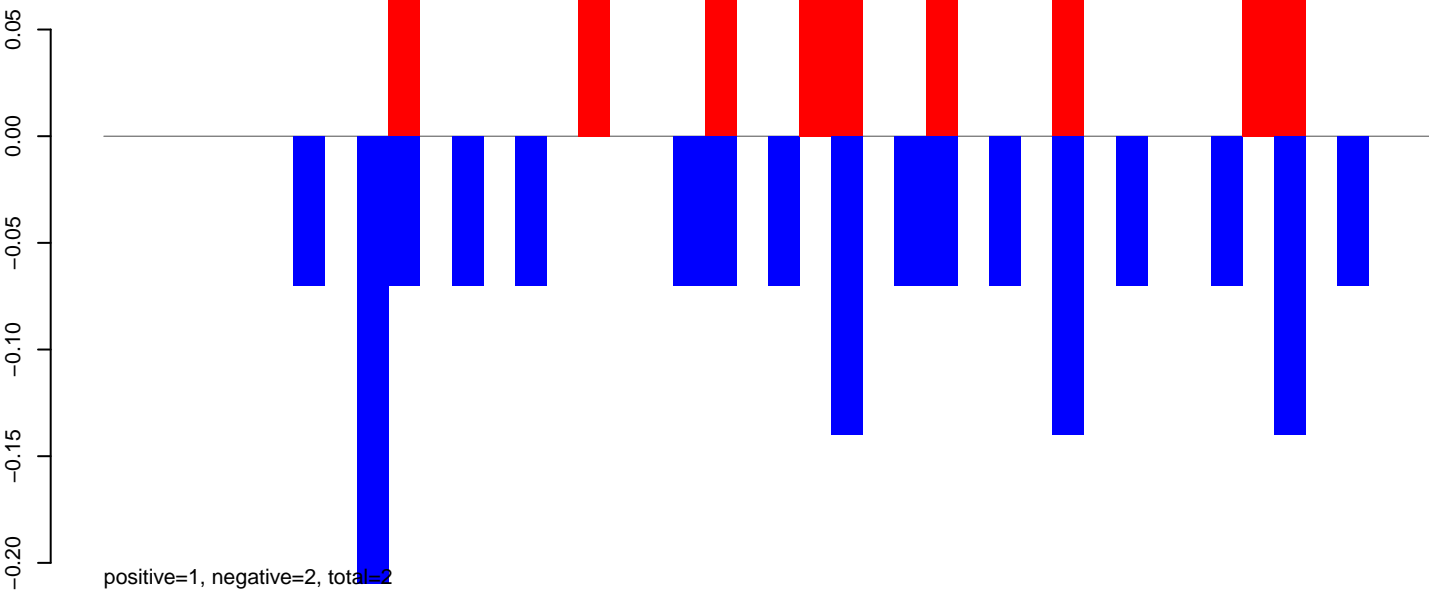


CuQuin_Mcarc_NL.rep

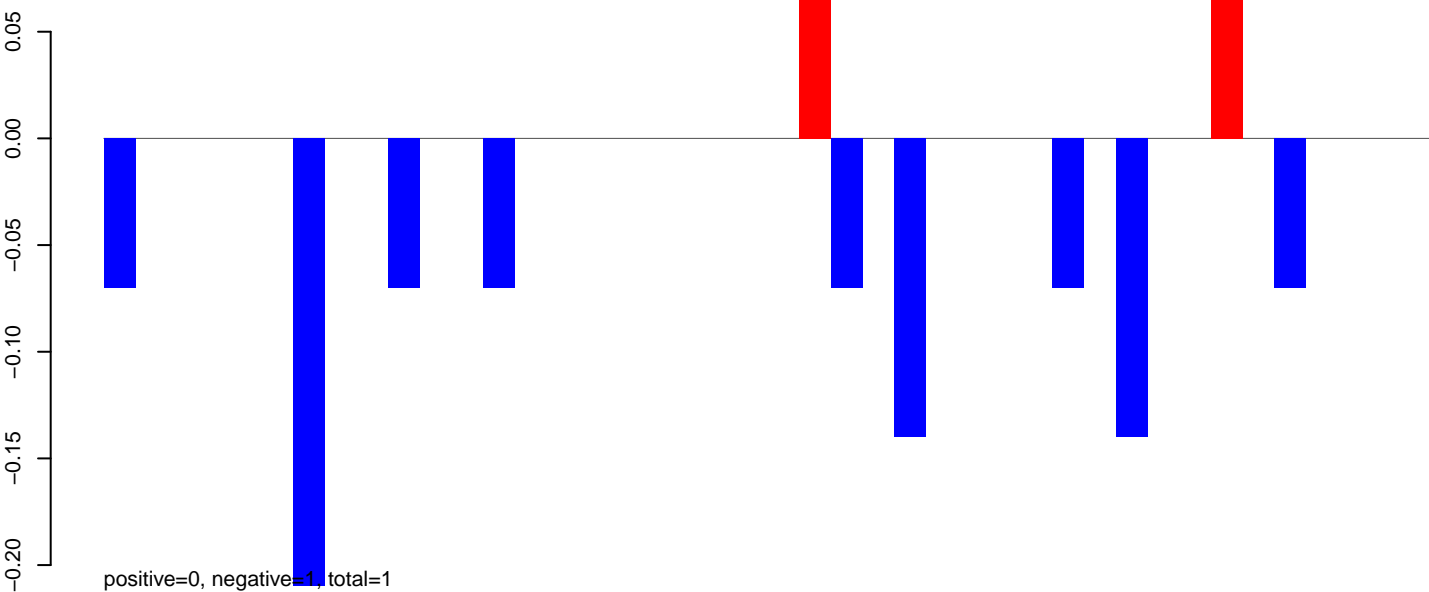


Window size=25, length=1118, TE@rnd-3_family-1133@DNA@725:1-1118

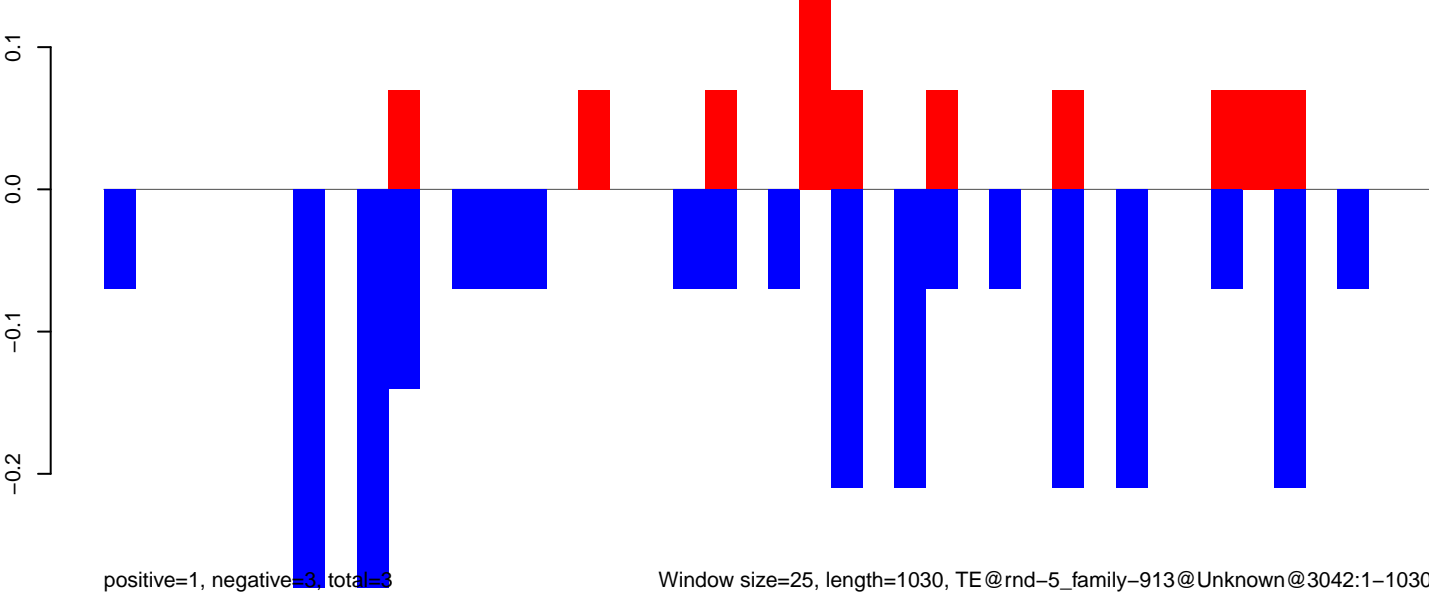
CuQuin_Mcarc_NL.18_23.rep



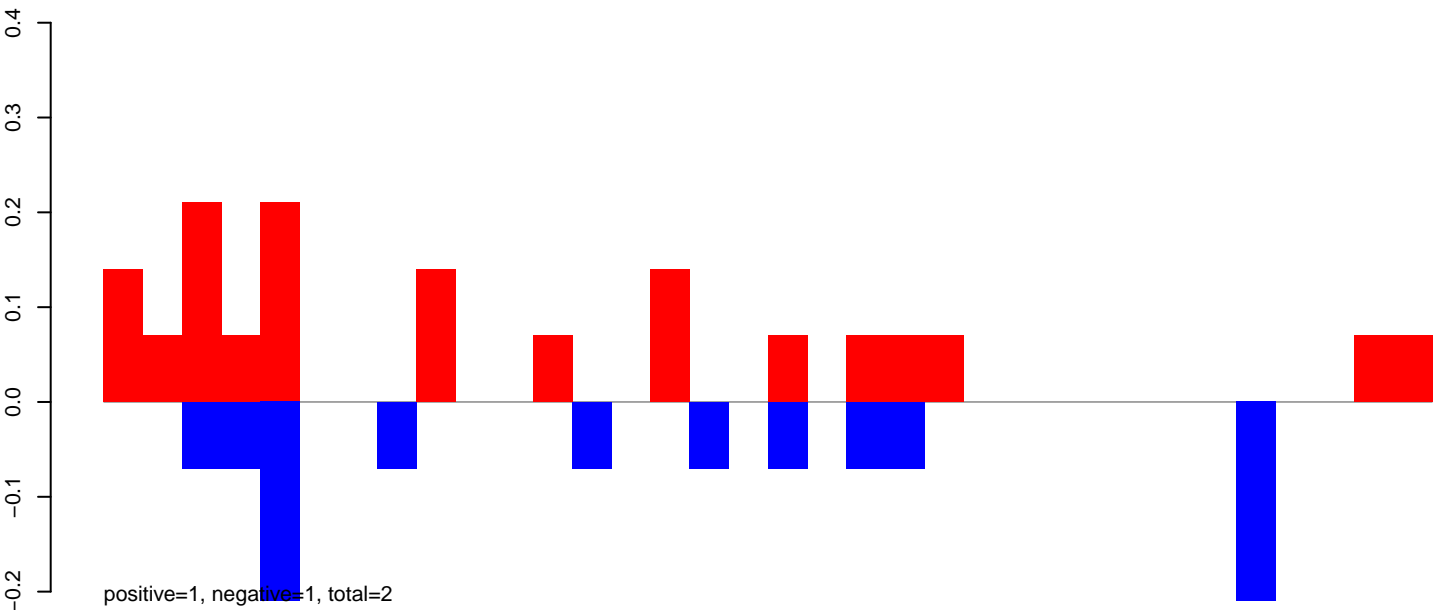
CuQuin_Mcarc_NL.24_35.rep



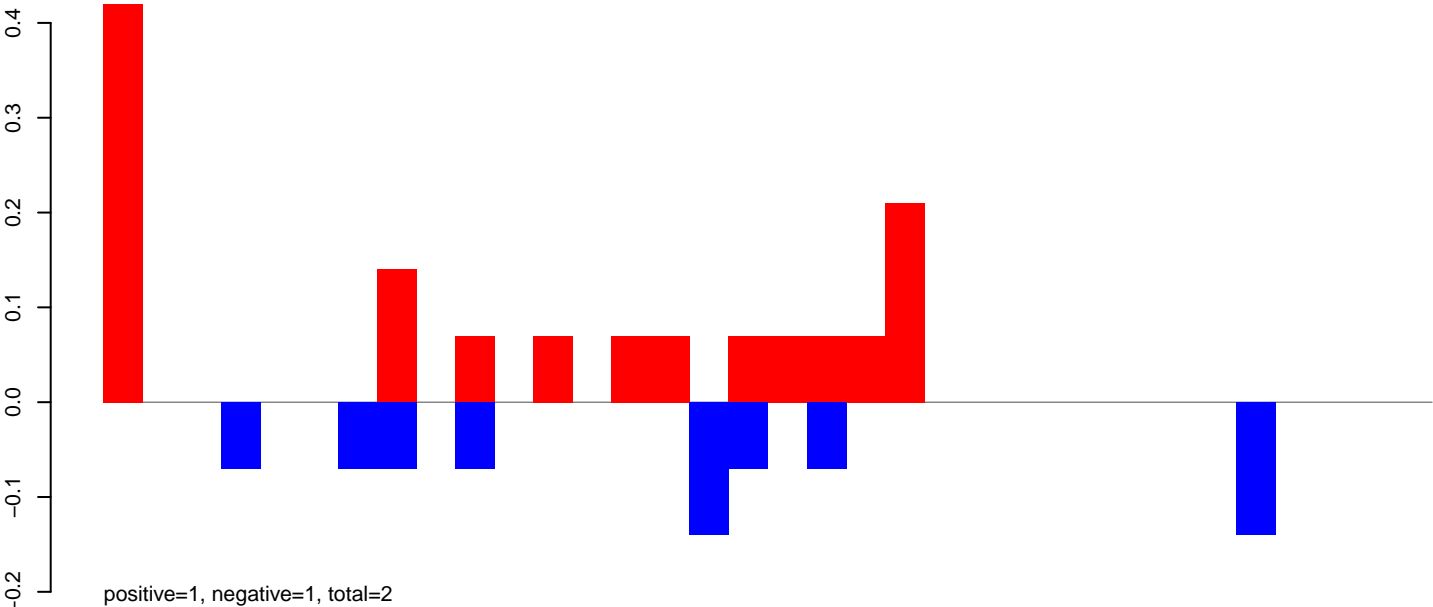
CuQuin_Mcarc_NL.rep



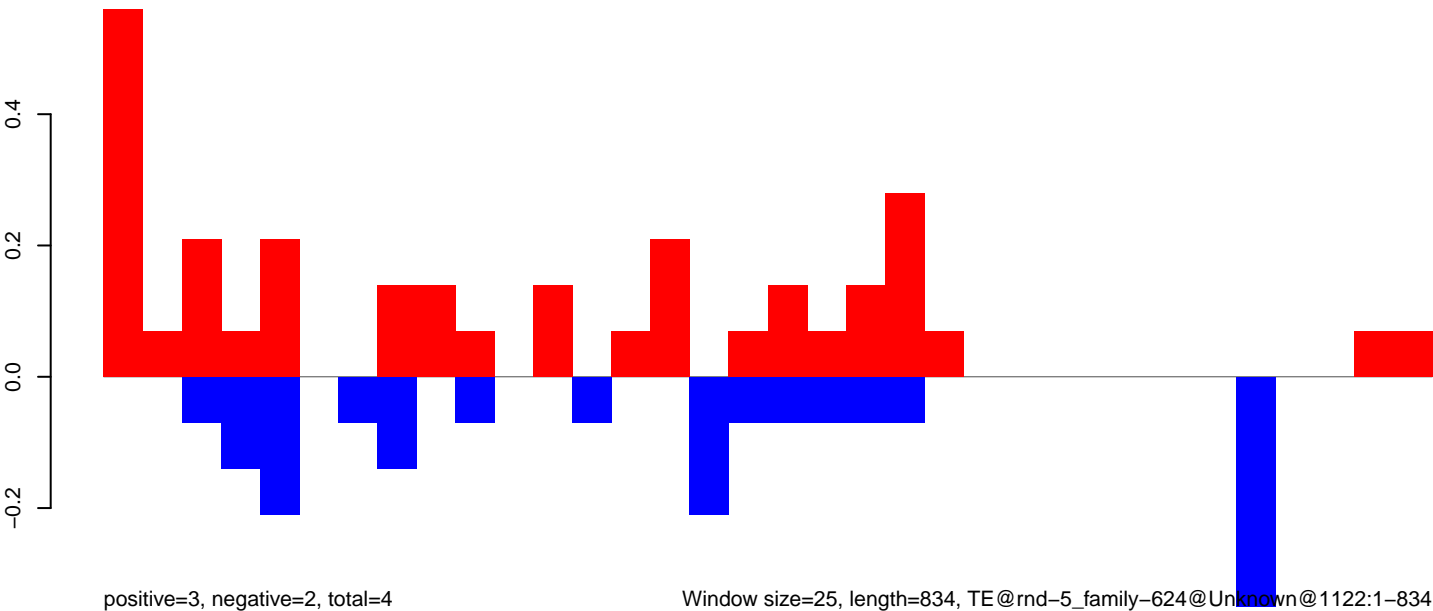
CuQuin_Mcarc_NL.18_23.rep



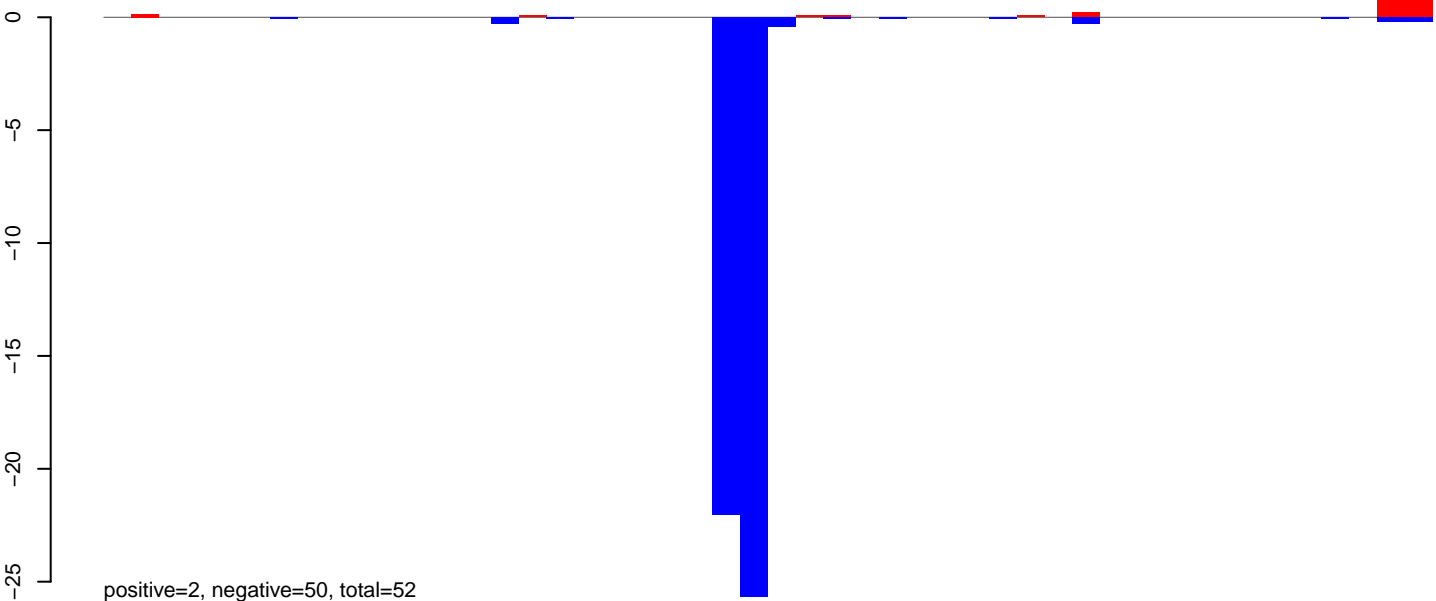
CuQuin_Mcarc_NL.24_35.rep



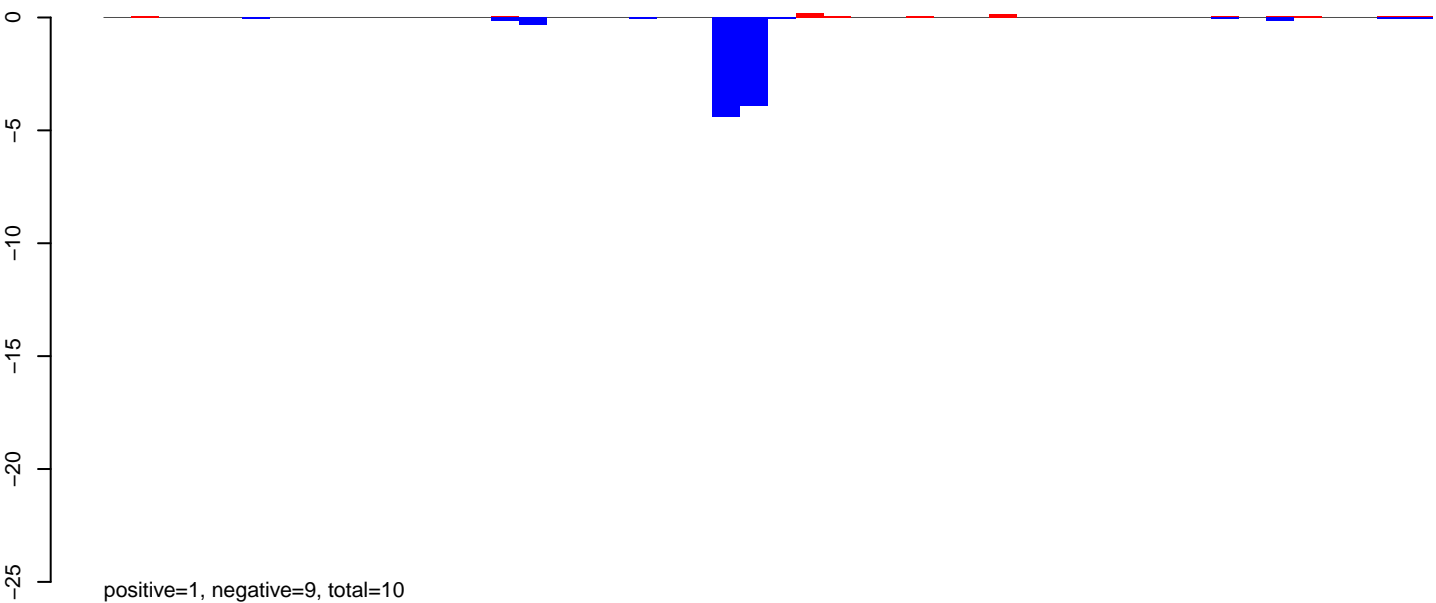
CuQuin_Mcarc_NL.rep



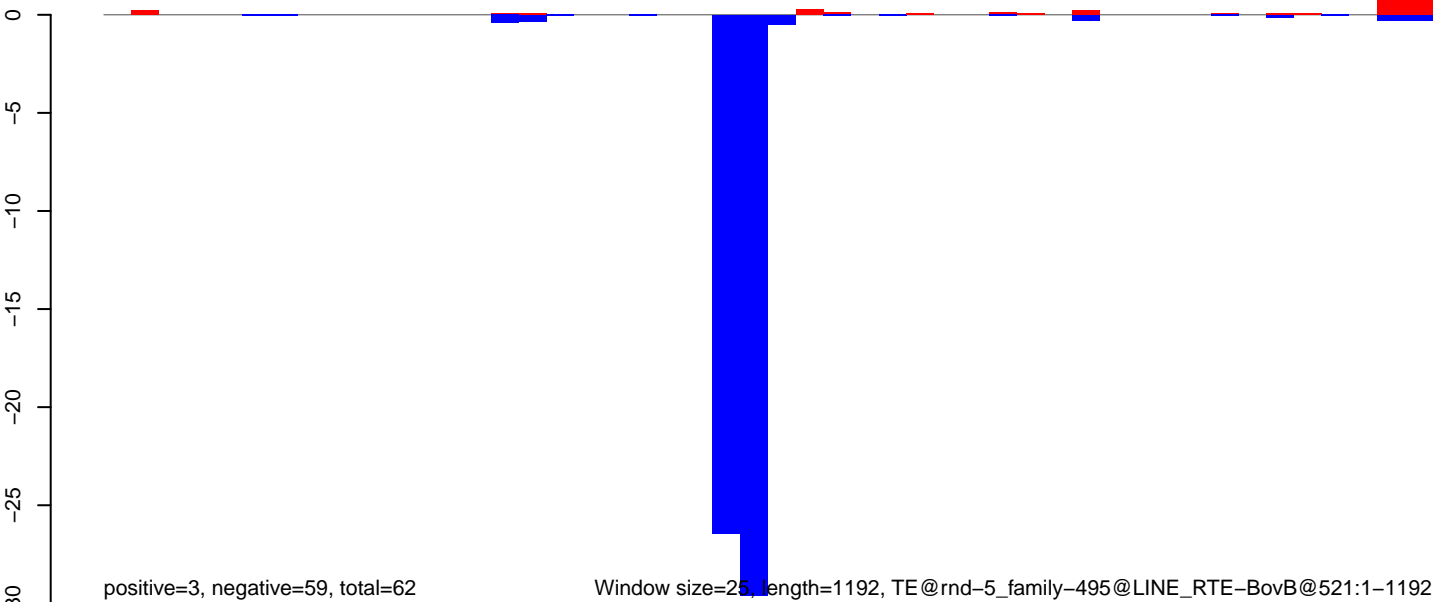
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

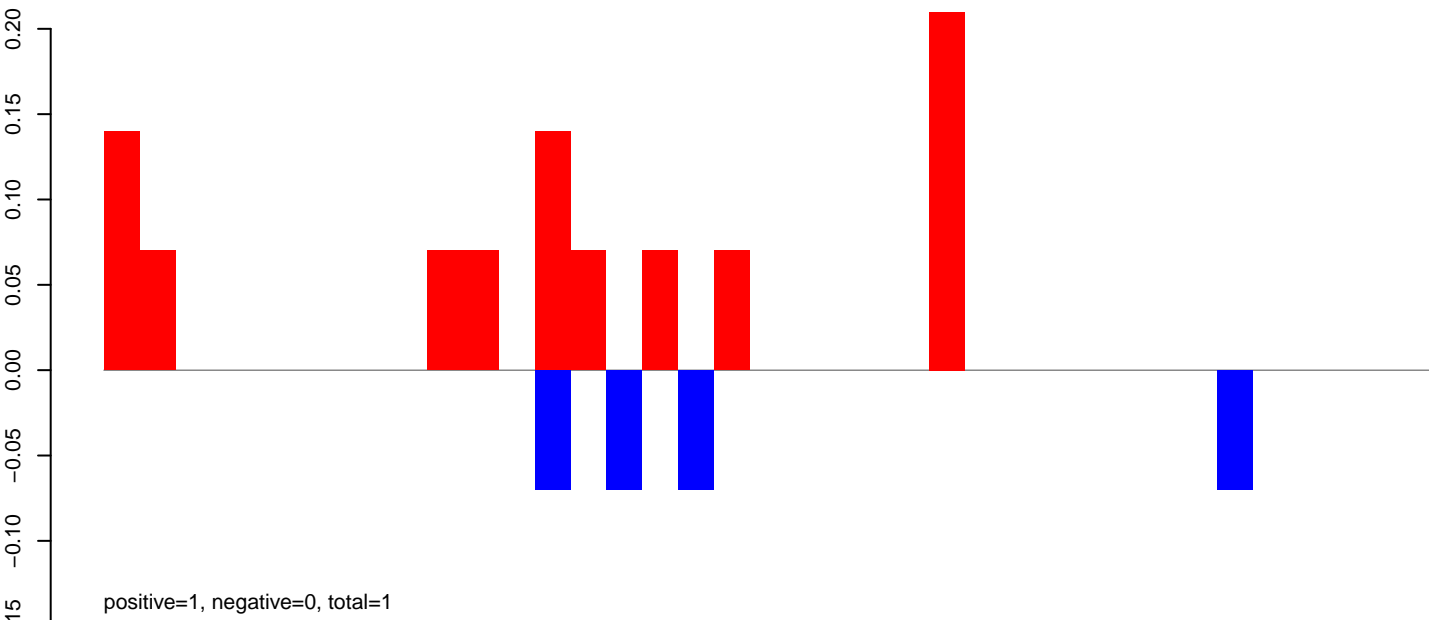


CuQuin_Mcarc_NL.rep

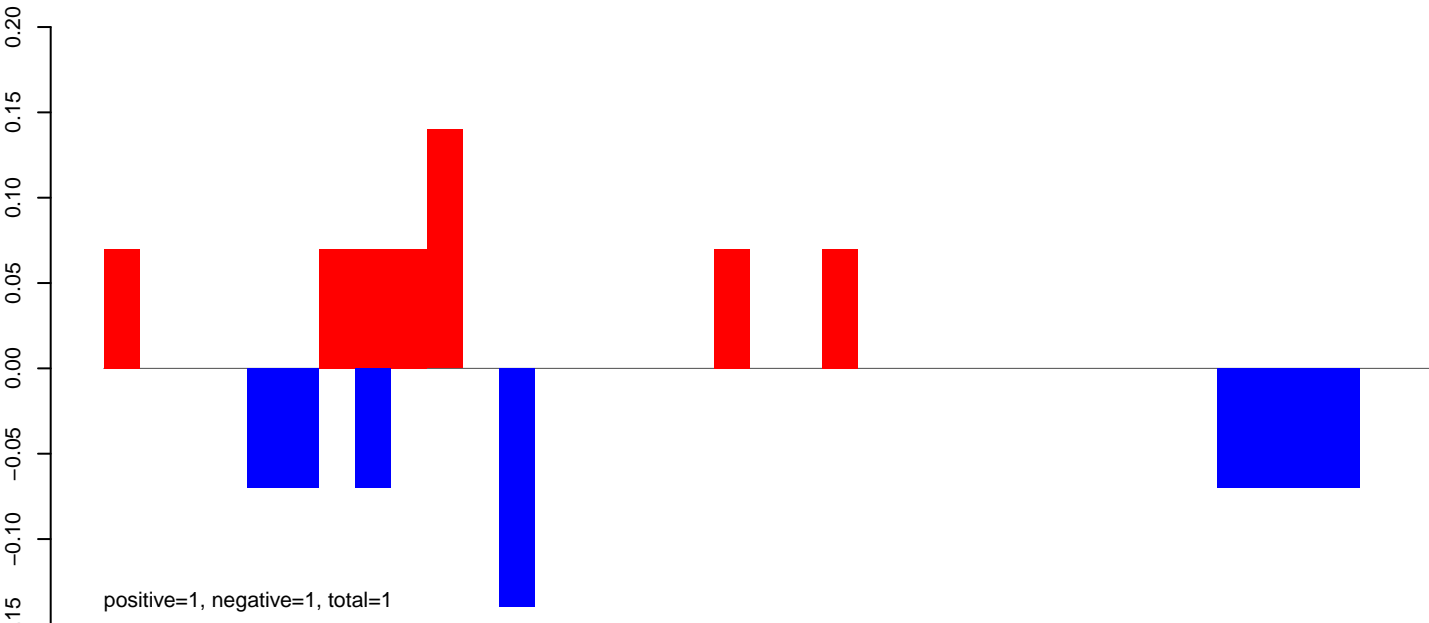


Window size=25, length=1192, TE@rnd-5_family-495@LINE_RTE-BovB@521:1-1192

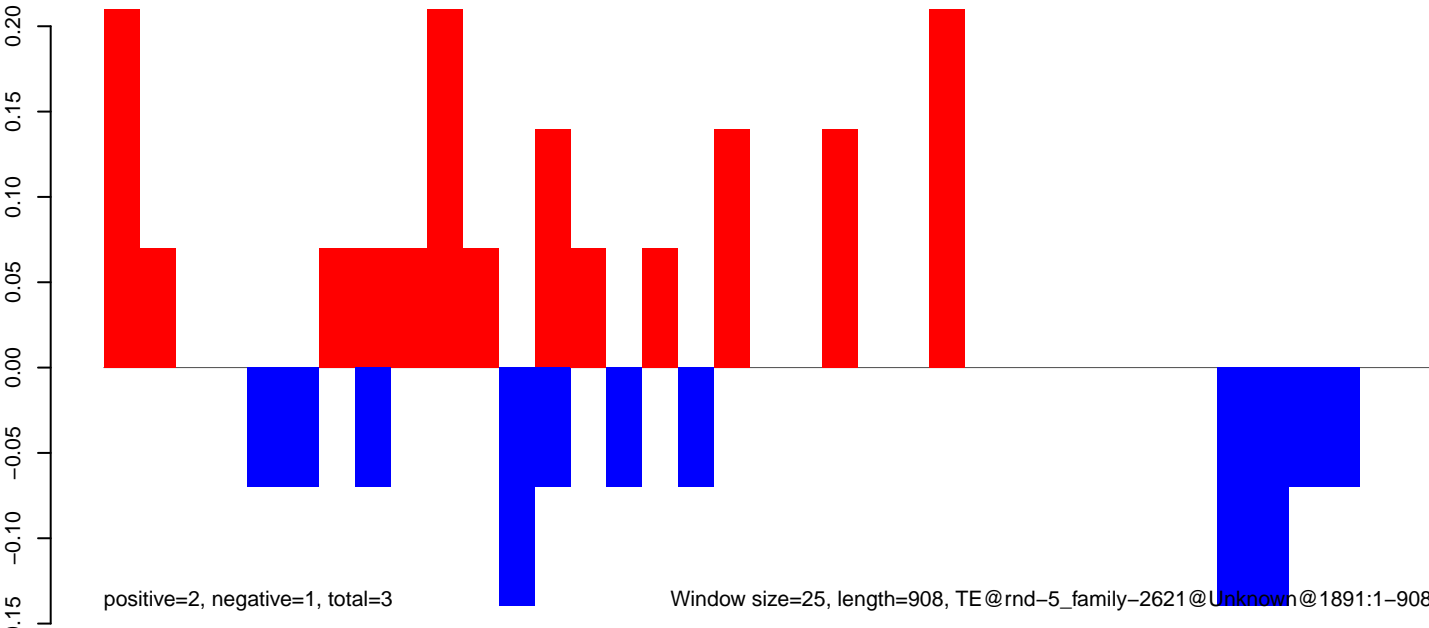
CuQuin_Mcarc_NL.18_23.rep



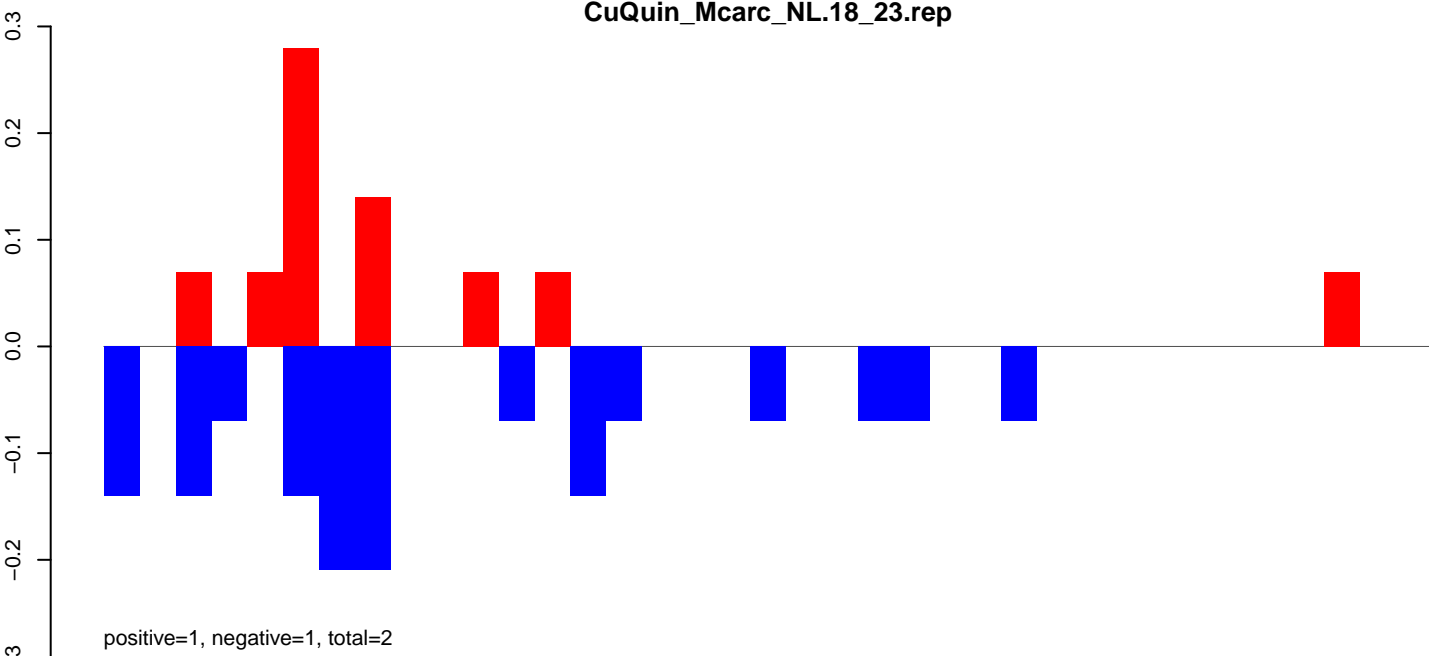
CuQuin_Mcarc_NL.24_35.rep



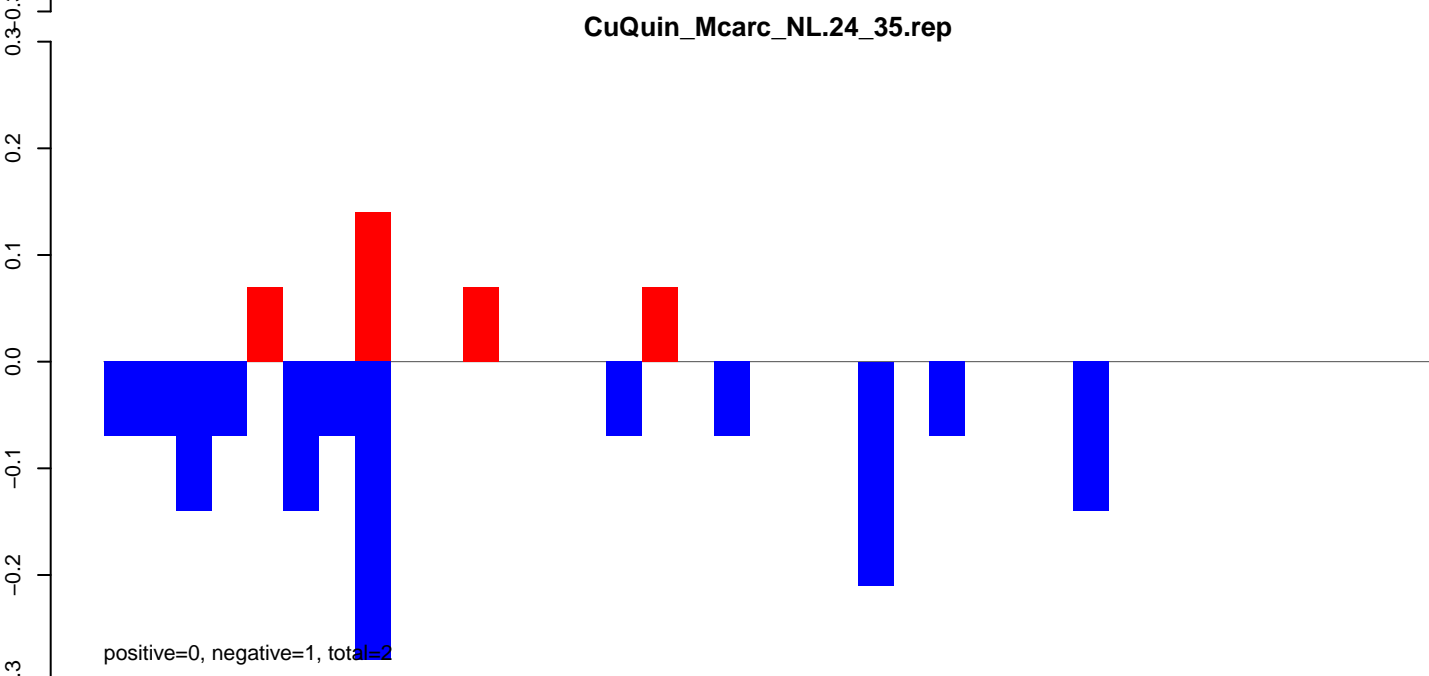
CuQuin_Mcarc_NL.rep



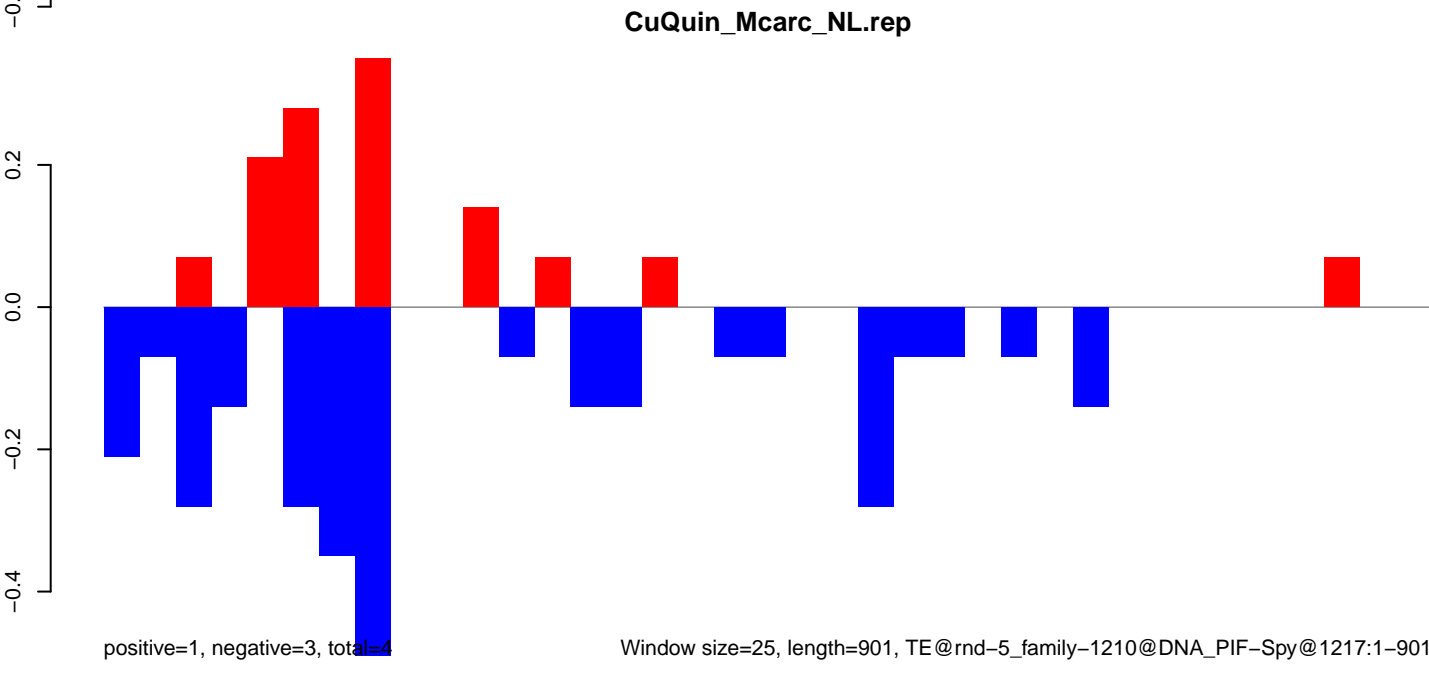
CuQuin_Mcarc_NL.18_23.rep



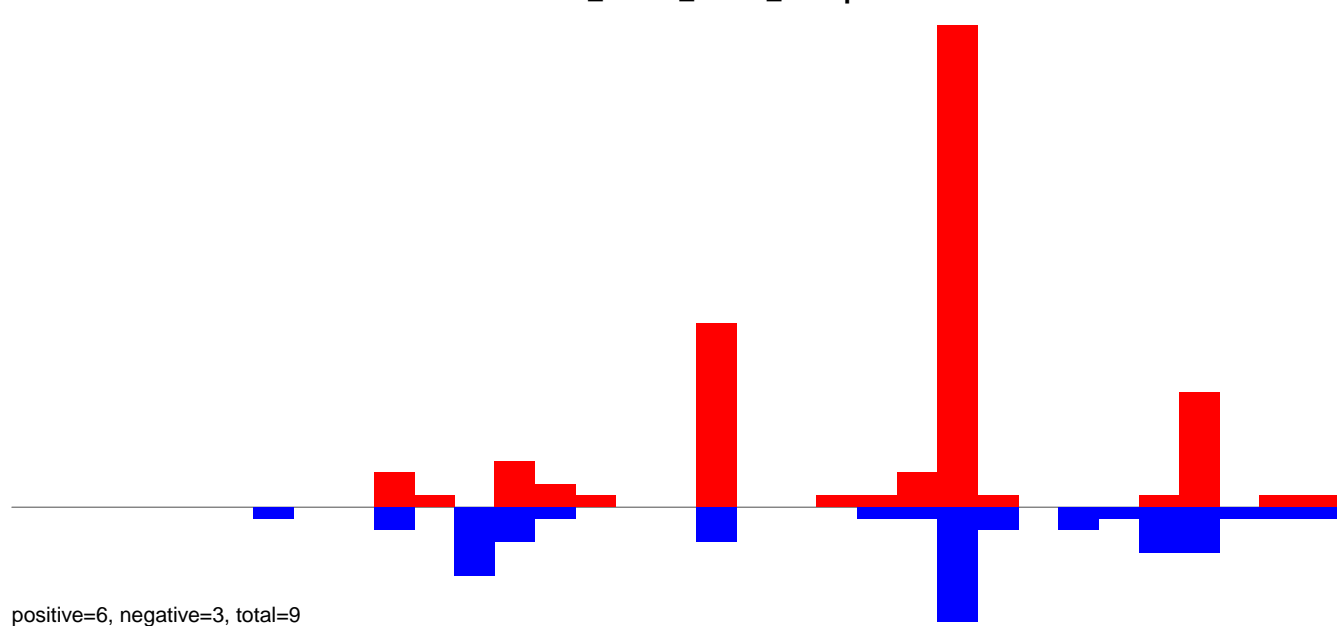
CuQuin_Mcarc_NL.24_35.rep



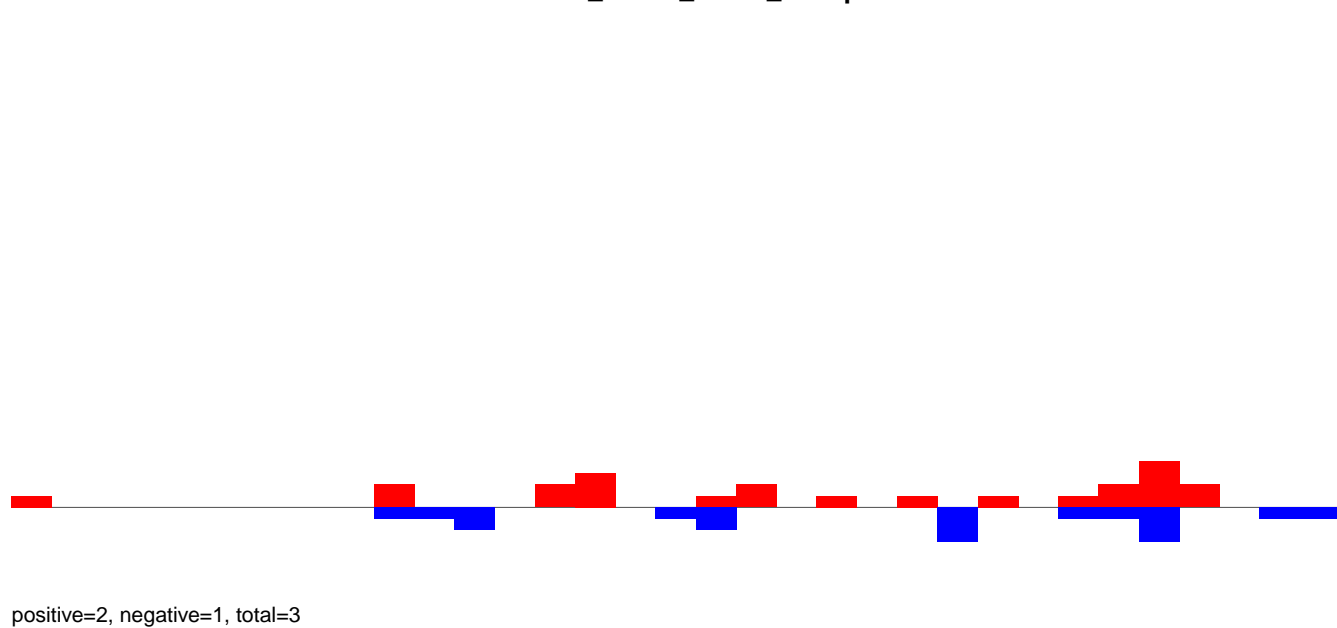
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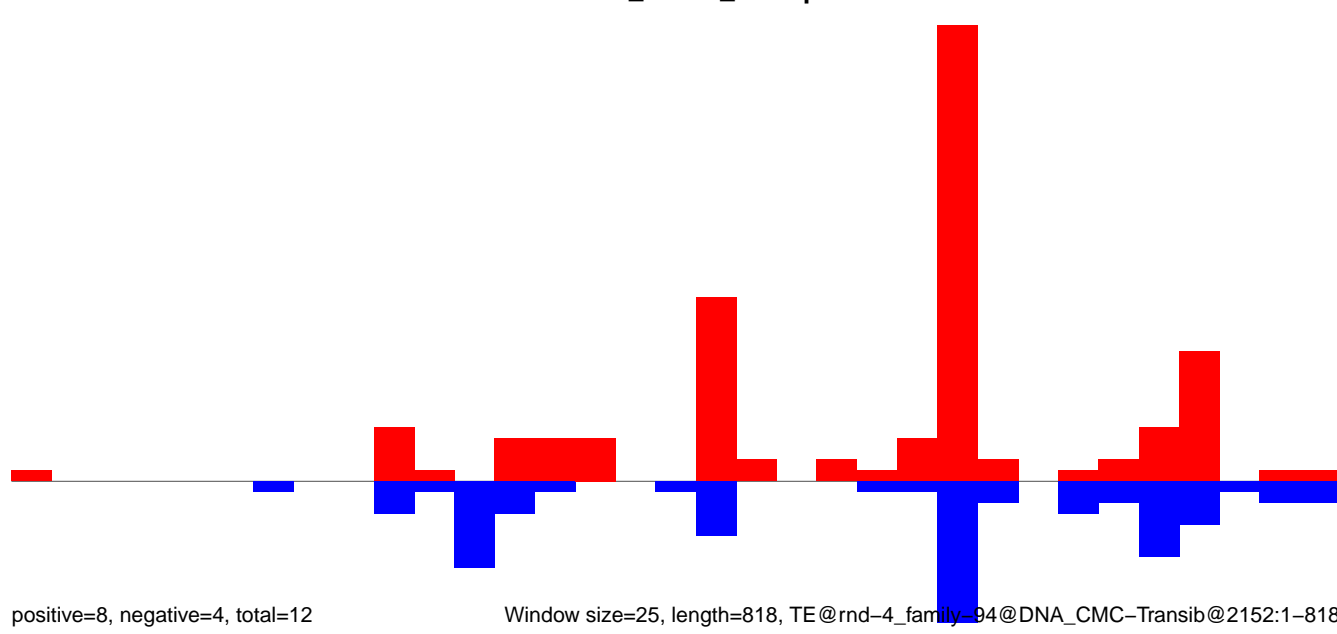
CuQuin_Mcarc_NL.18_23.rep



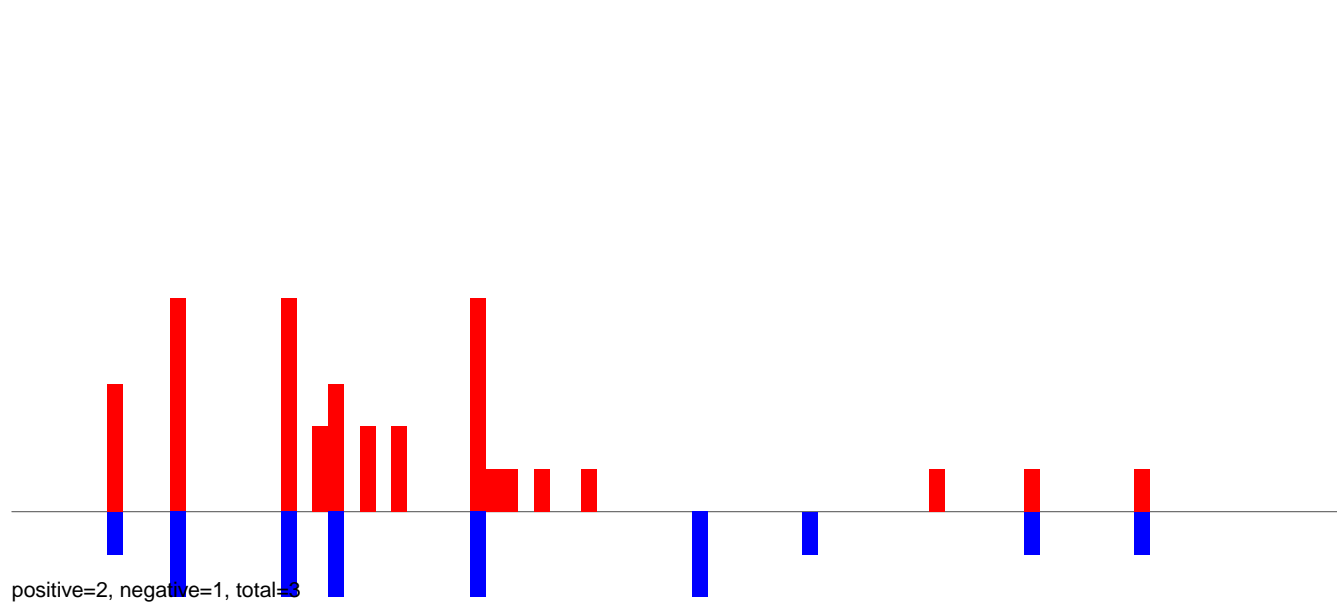
CuQuin_Mcarc_NL.24_35.rep



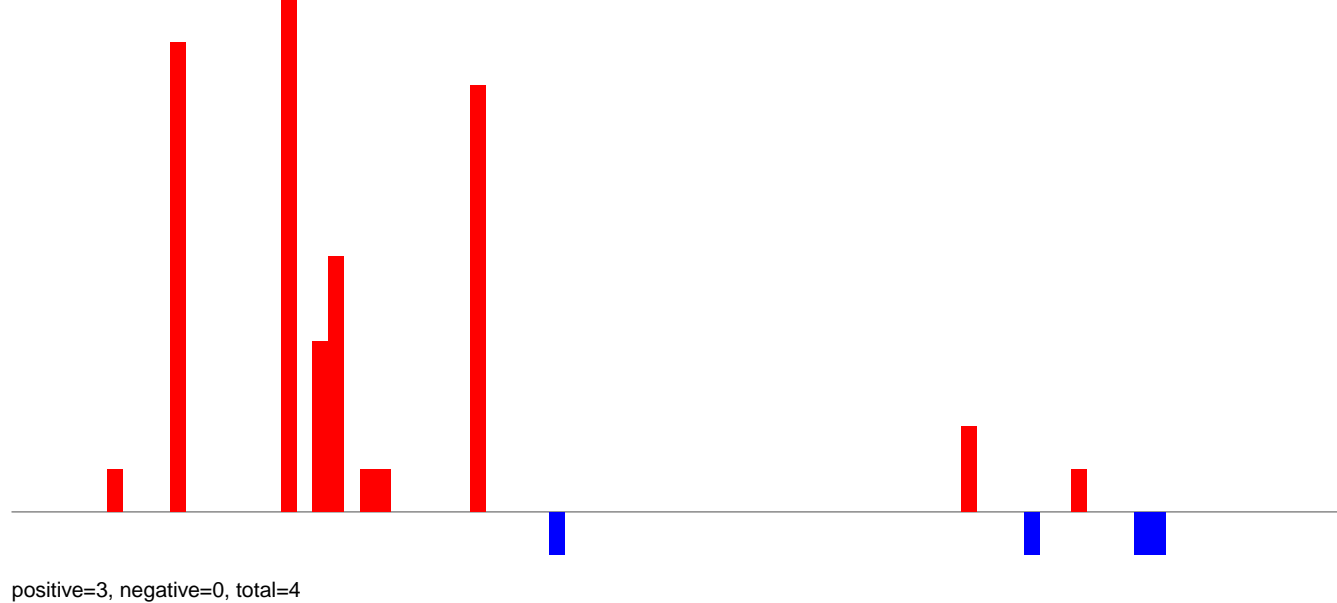
CuQuin_Mcarc_NL.rep



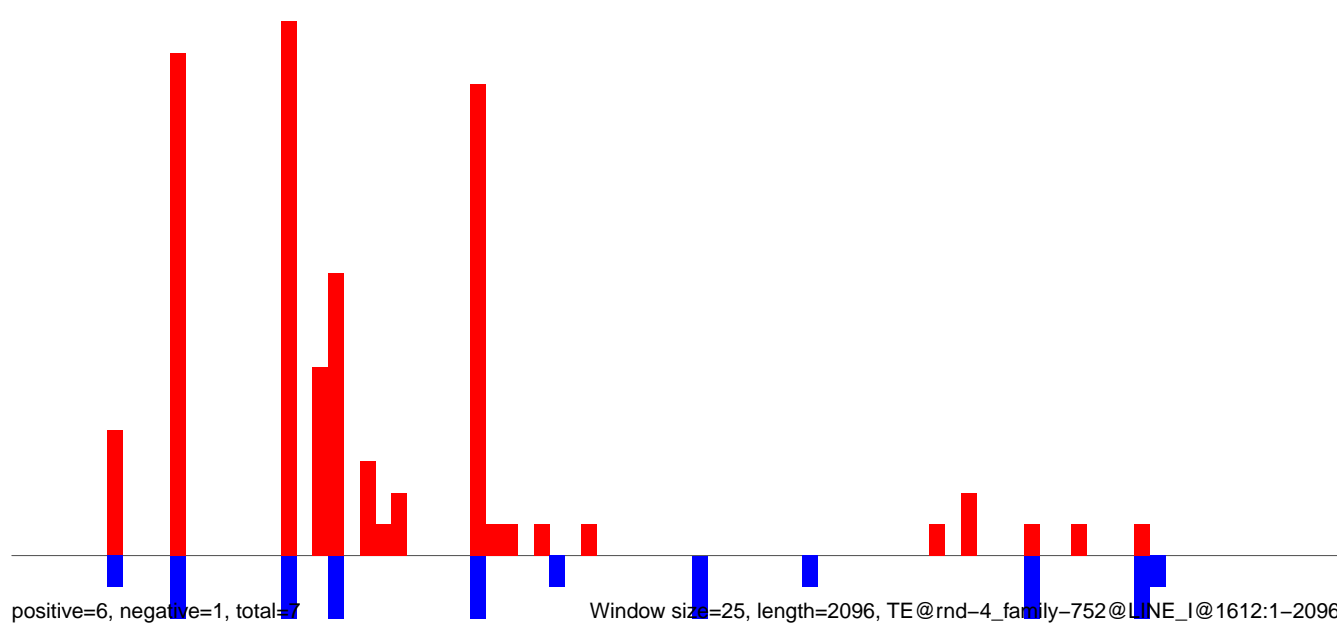
CuQuin_Mcarc_NL.18_23.rep



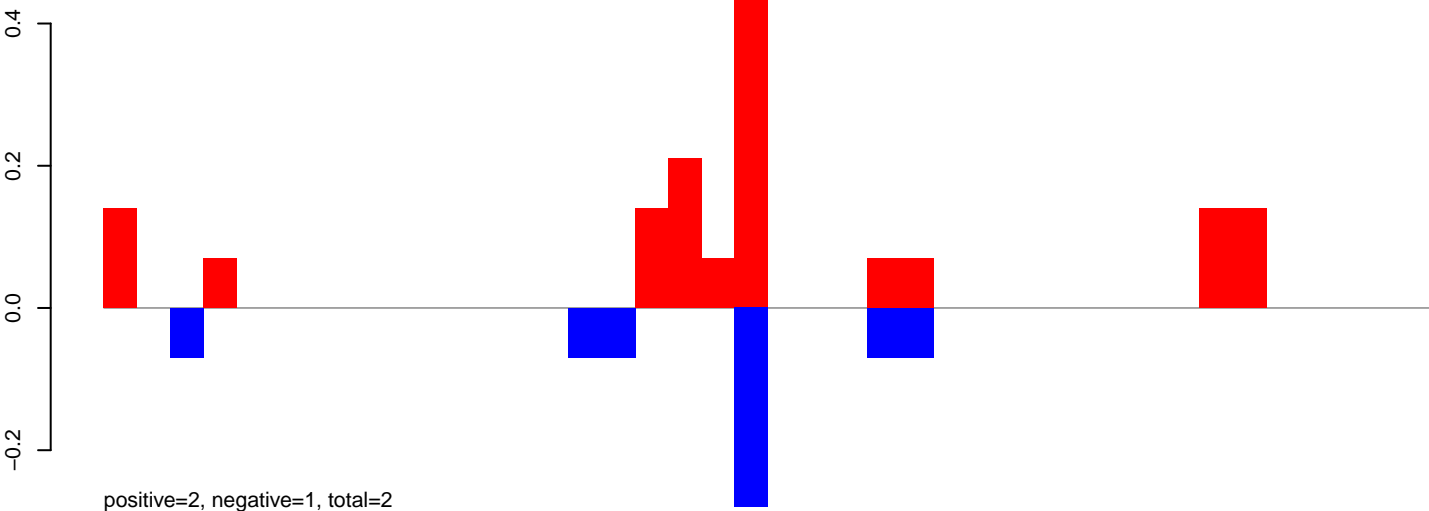
CuQuin_Mcarc_NL.24_35.rep



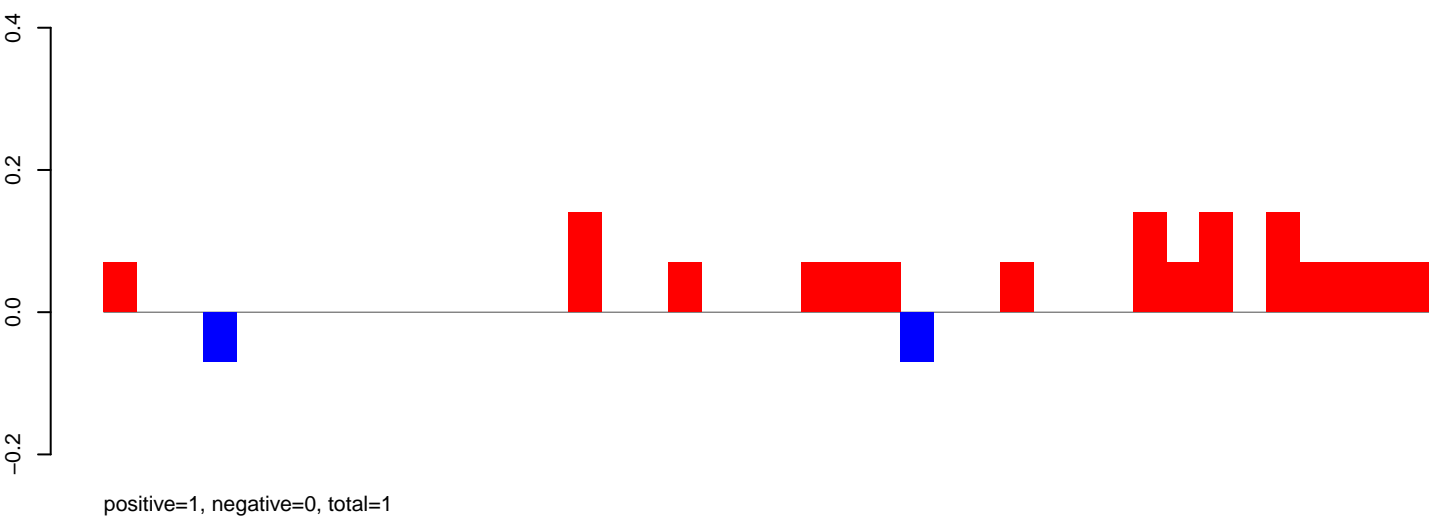
CuQuin_Mcarc_NL.rep



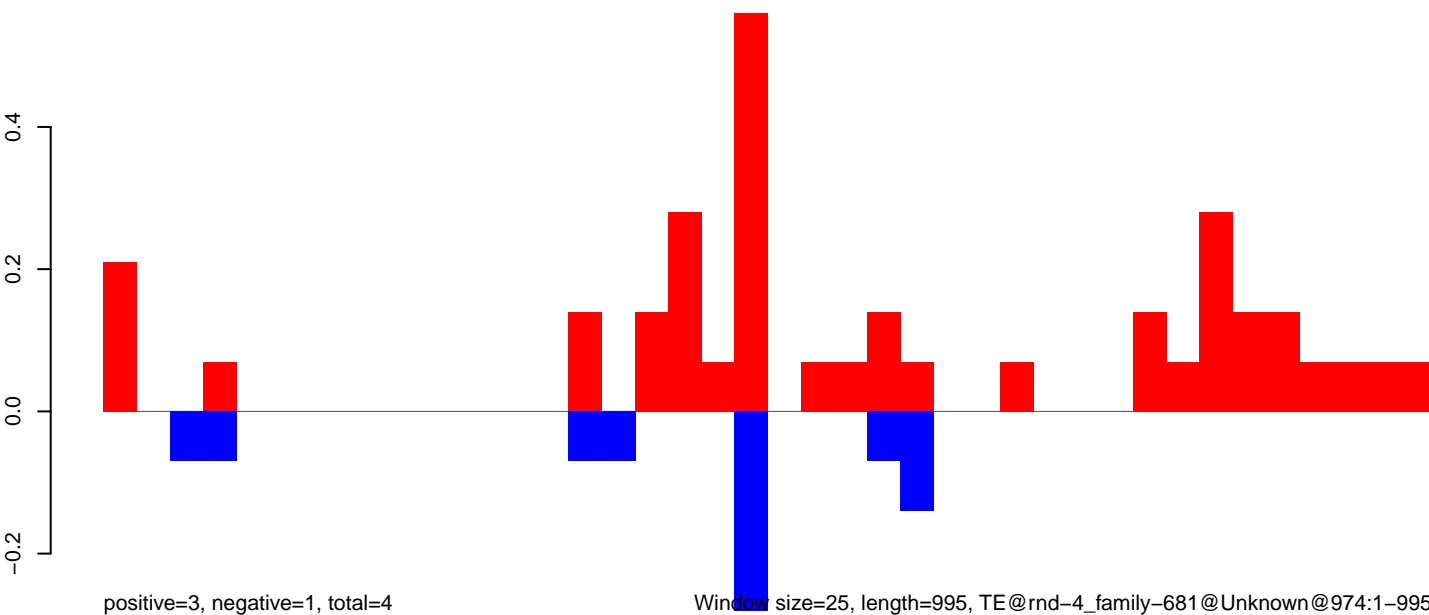
CuQuin_Mcarc_NL.18_23.rep



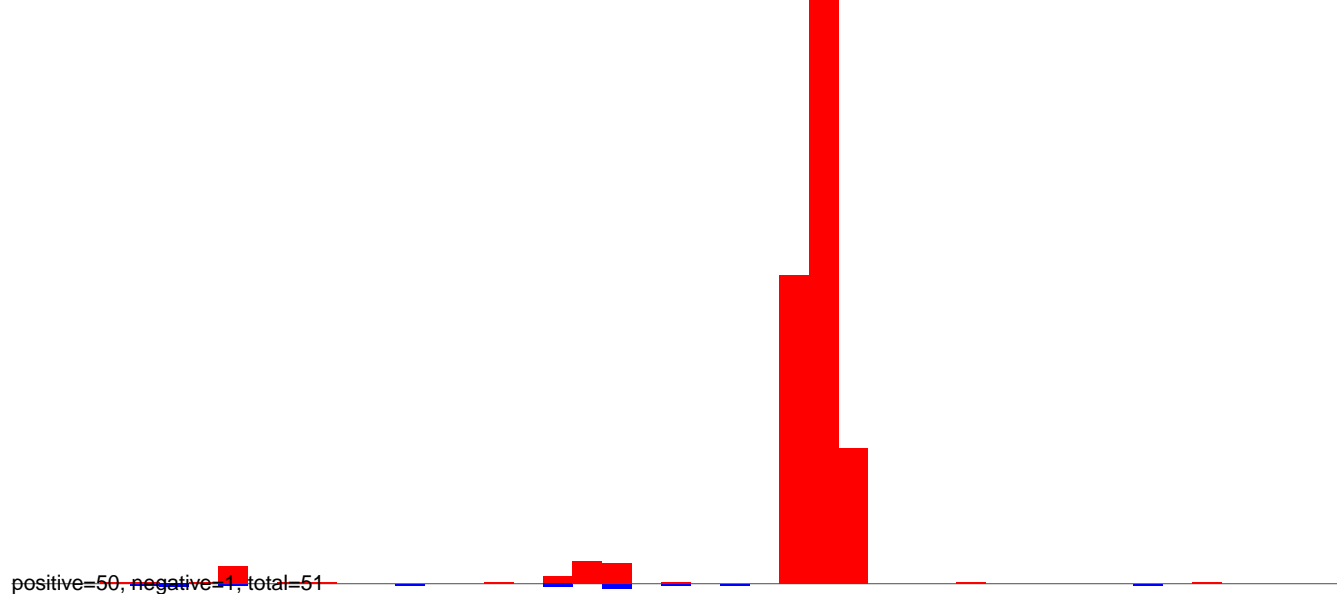
CuQuin_Mcarc_NL.24_35.rep



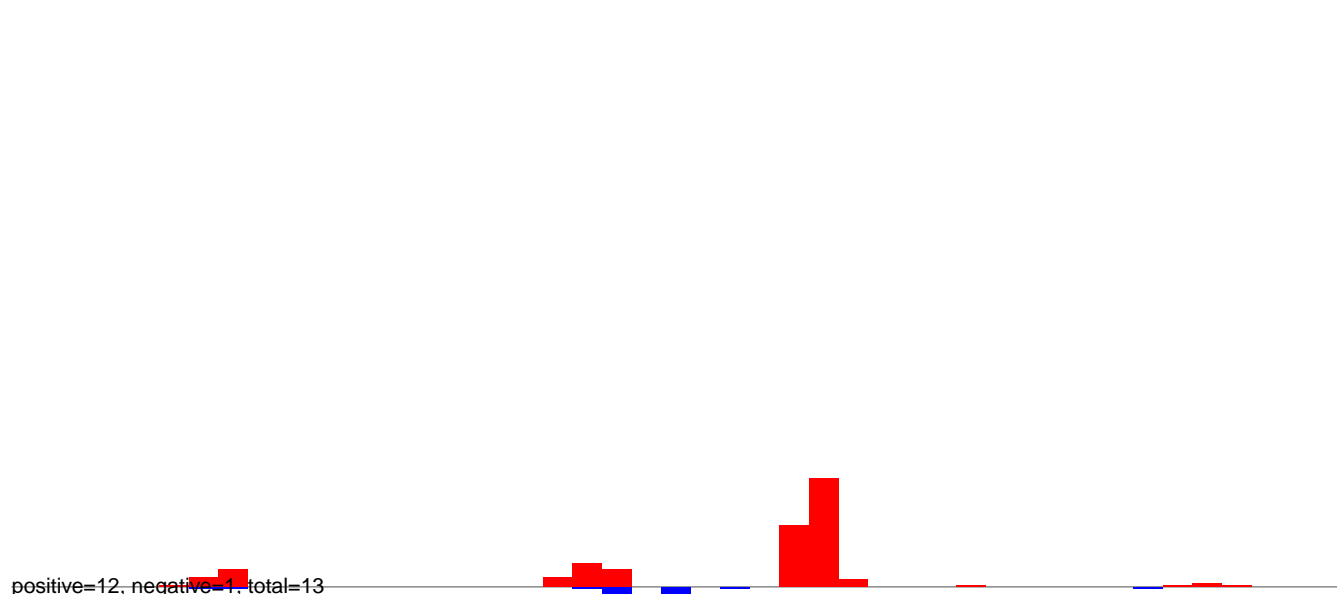
CuQuin_Mcarc_NL.rep



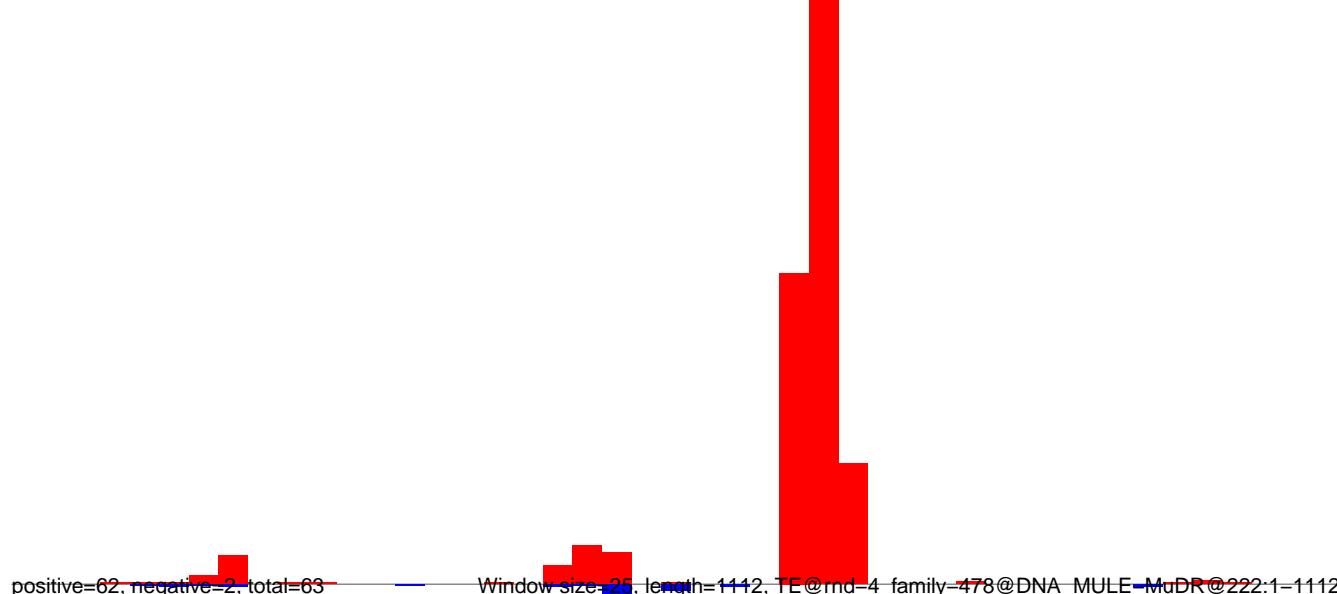
CuQuin_Mcarc_NL.18_23.rep



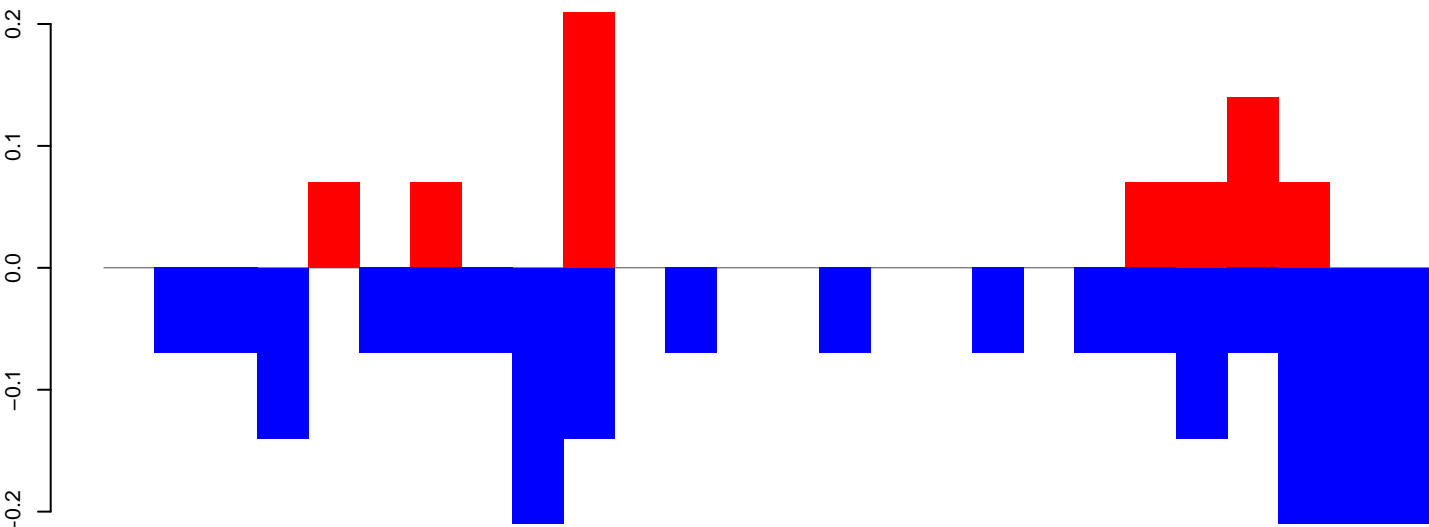
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

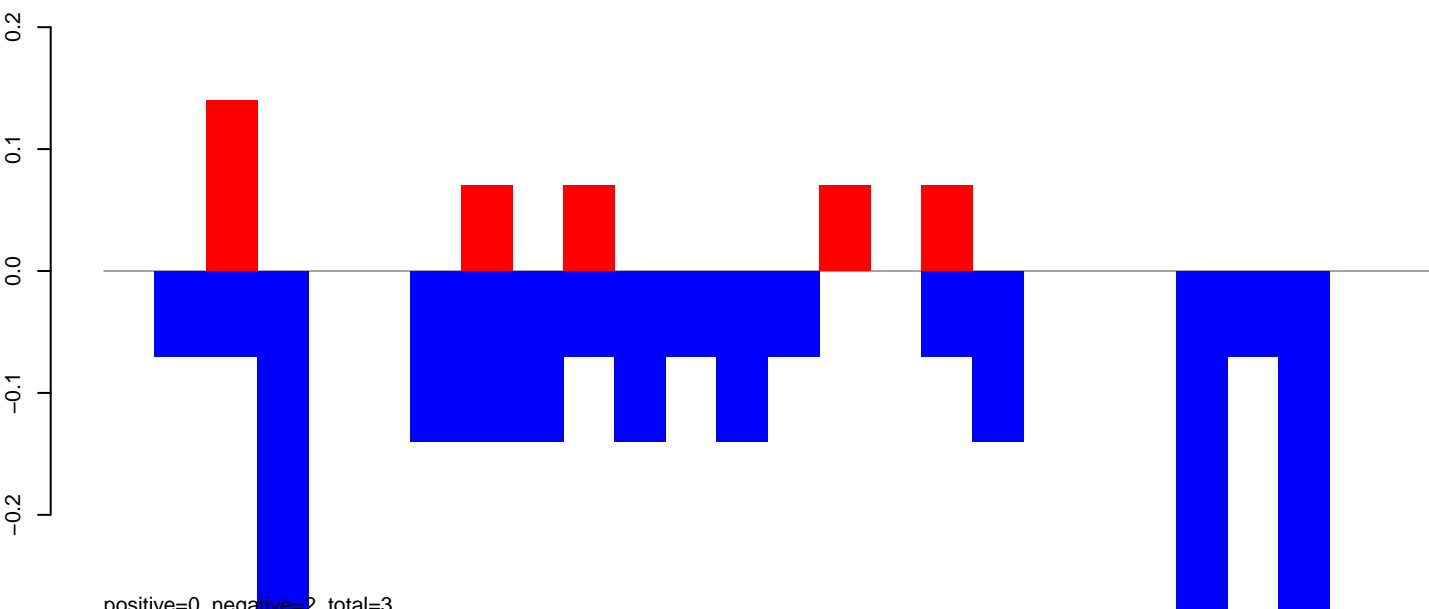


CuQuin_Mcarc_NL.18_23.rep



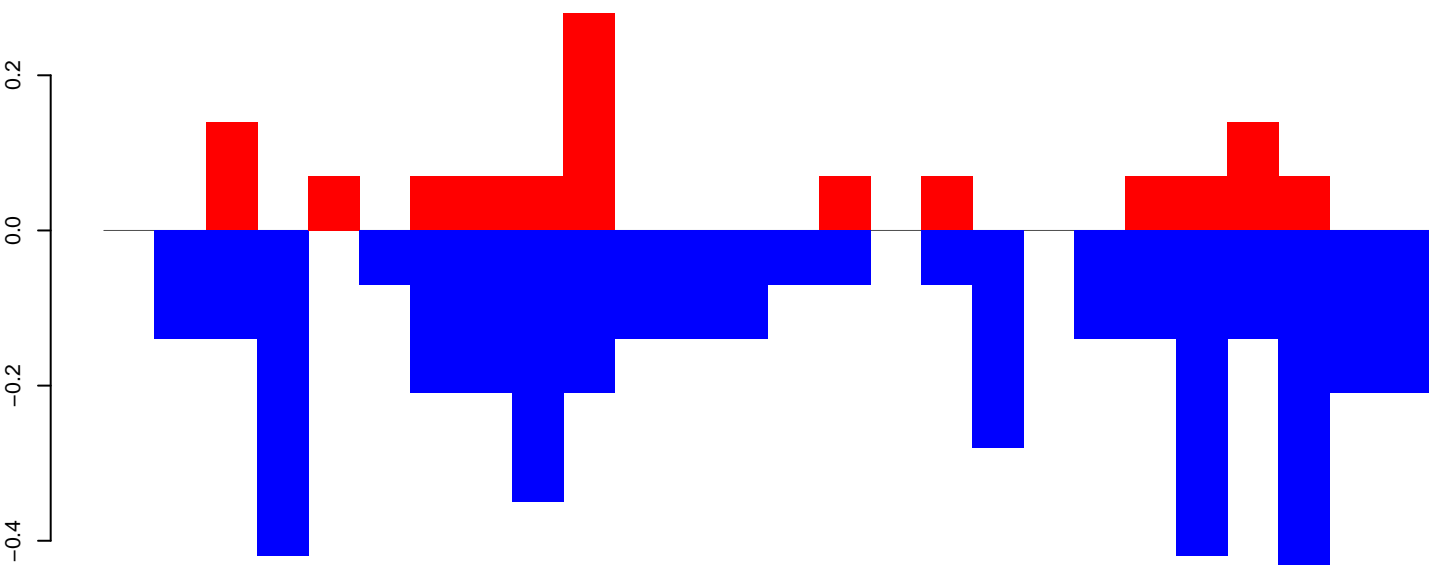
positive=1, negative=2, total=3

CuQuin_Mcarc_NL.24_35.rep



positive=0, negative=2, total=3

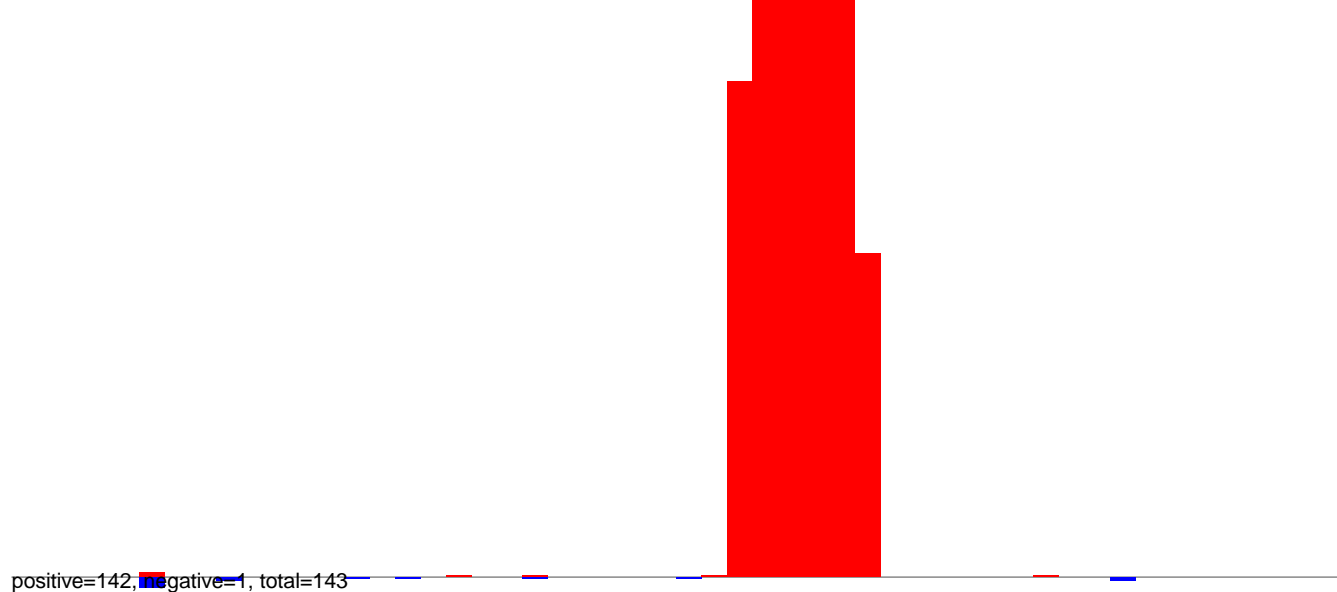
CuQuin_Mcarc_NL.rep



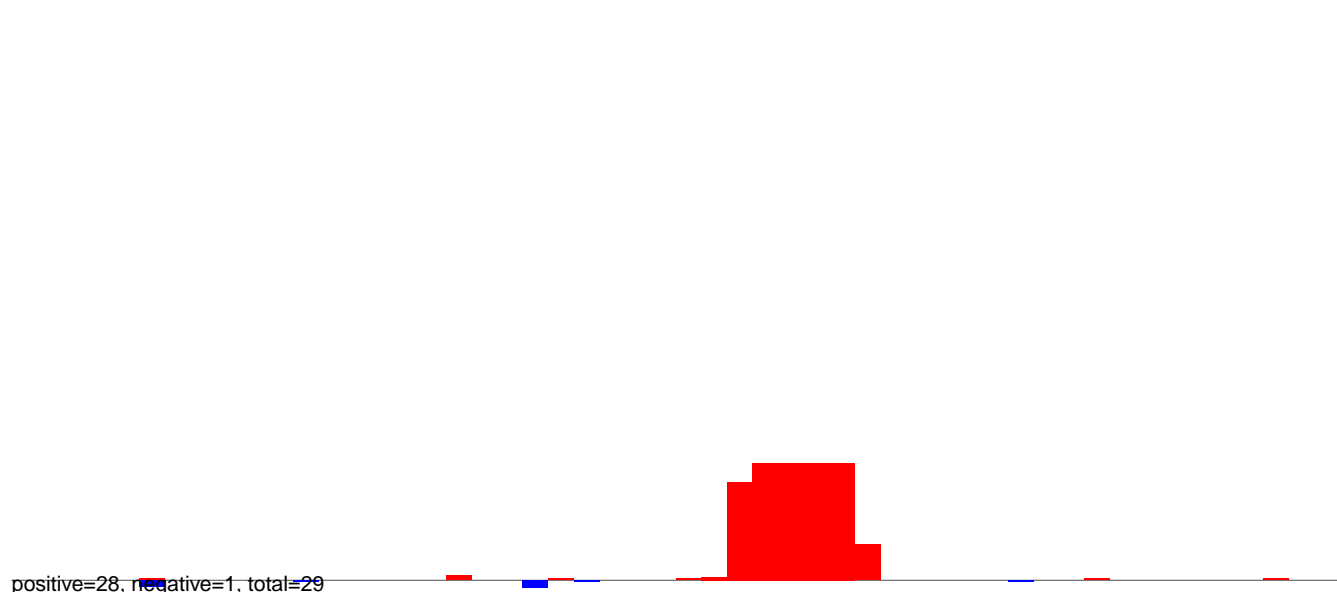
positive=1, negative=4, total=6

Window size=25, length=647, TE@rnd-4_family-403@Unknown@279:1-647

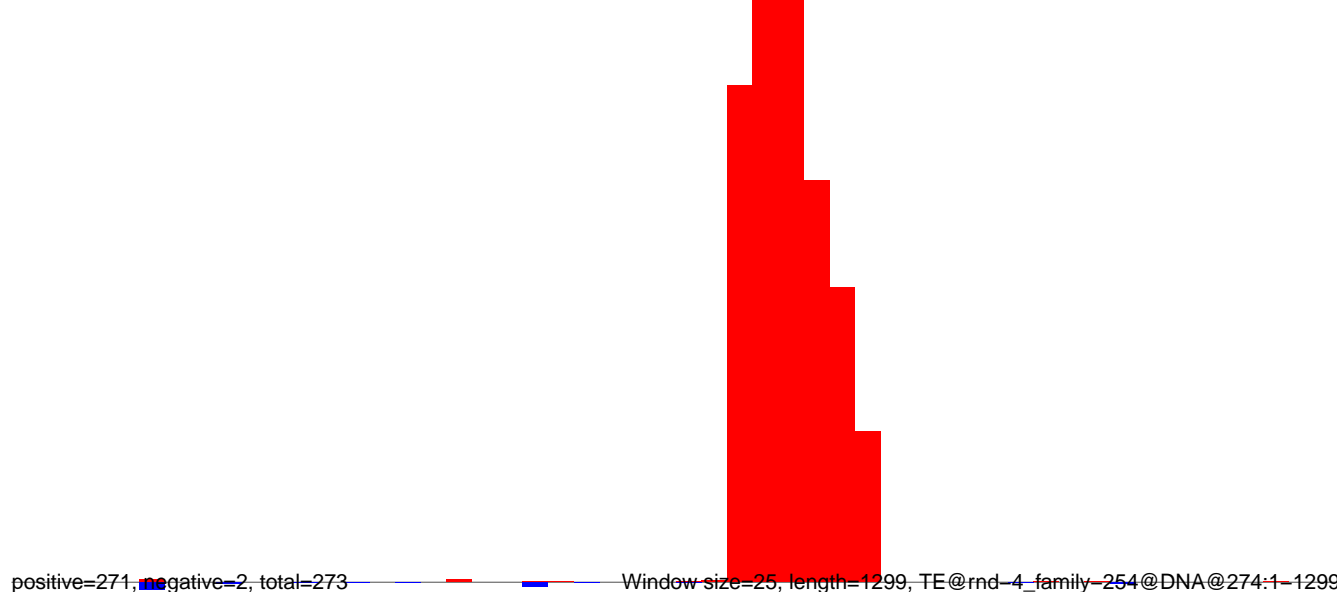
CuQuin_Mcarc_NL.18_23.rep



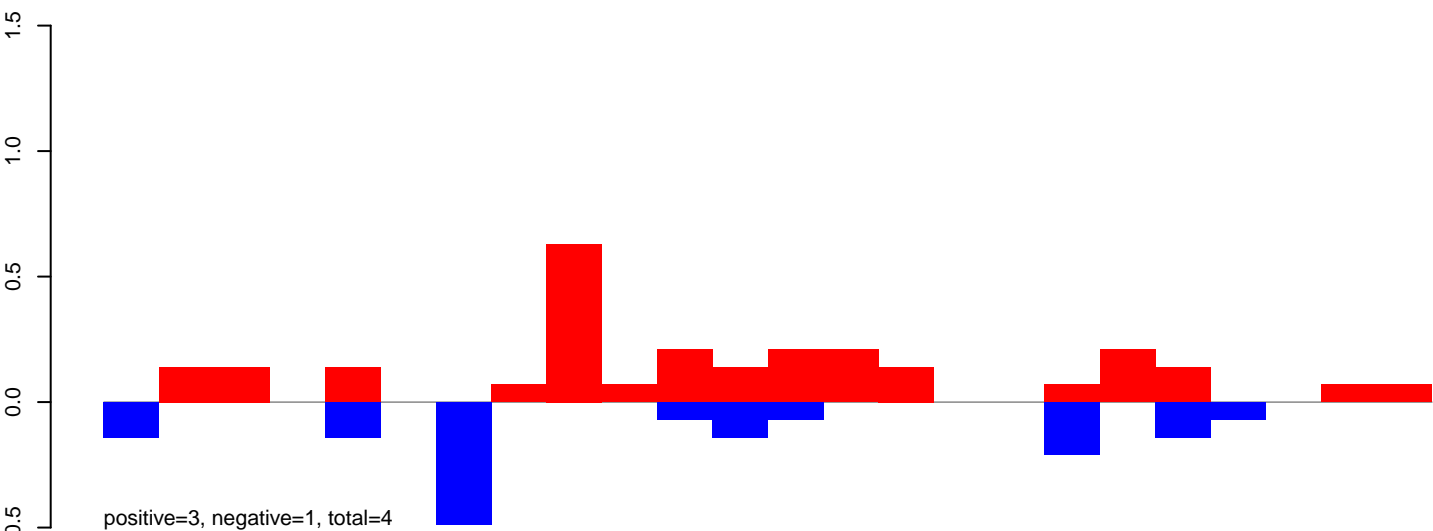
CuQuin_Mcarc_NL.24_35.rep



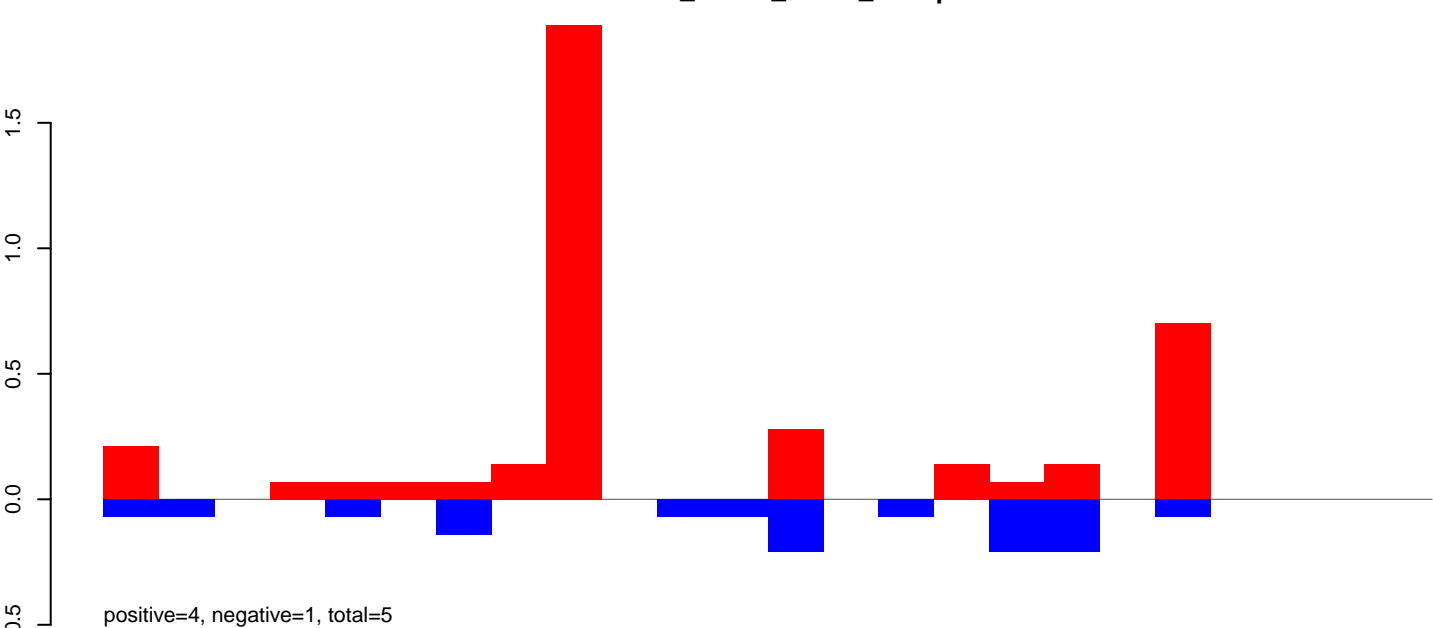
CuQuin_Mcarc_NL.rep



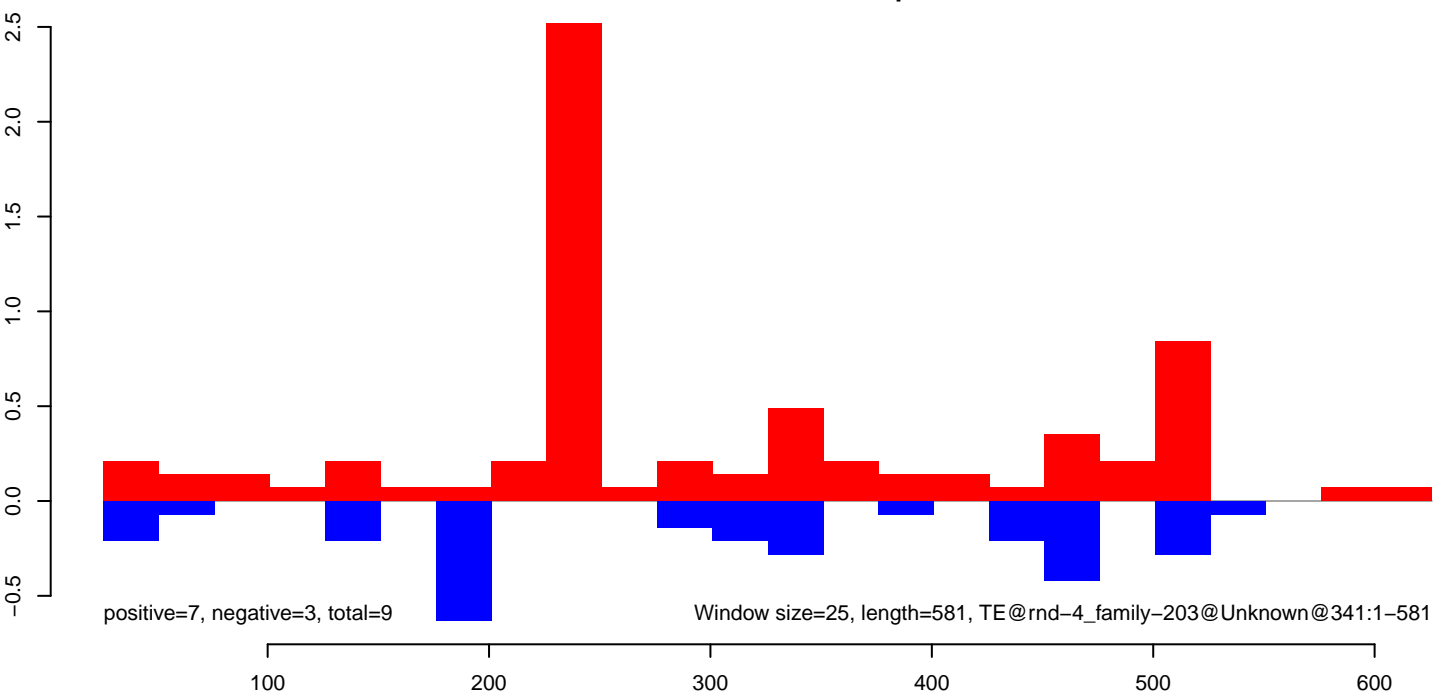
CuQuin_Mcarc_NL.18_23.rep



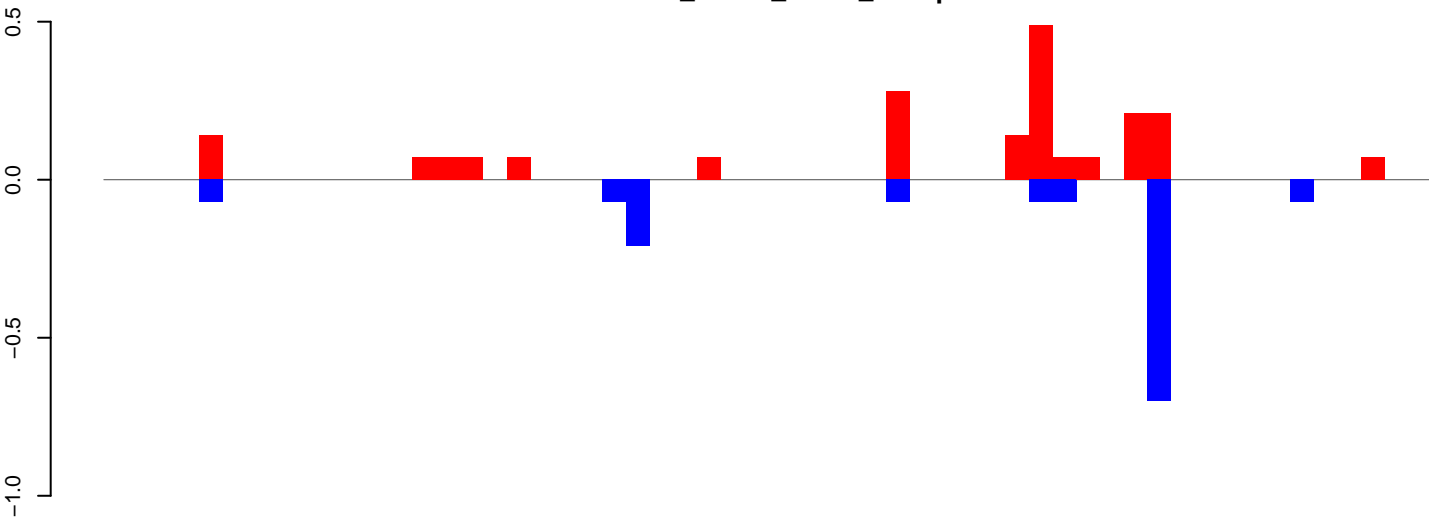
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

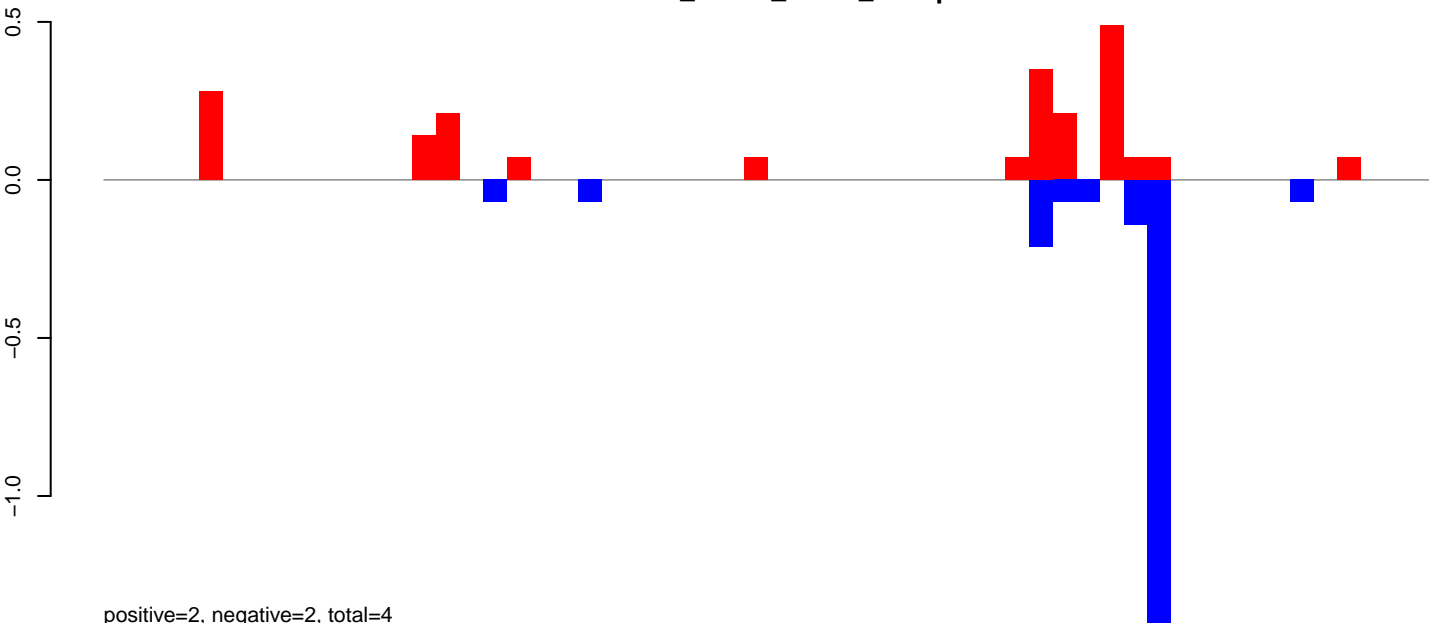


CuQuin_Mcarc_NL.18_23.rep



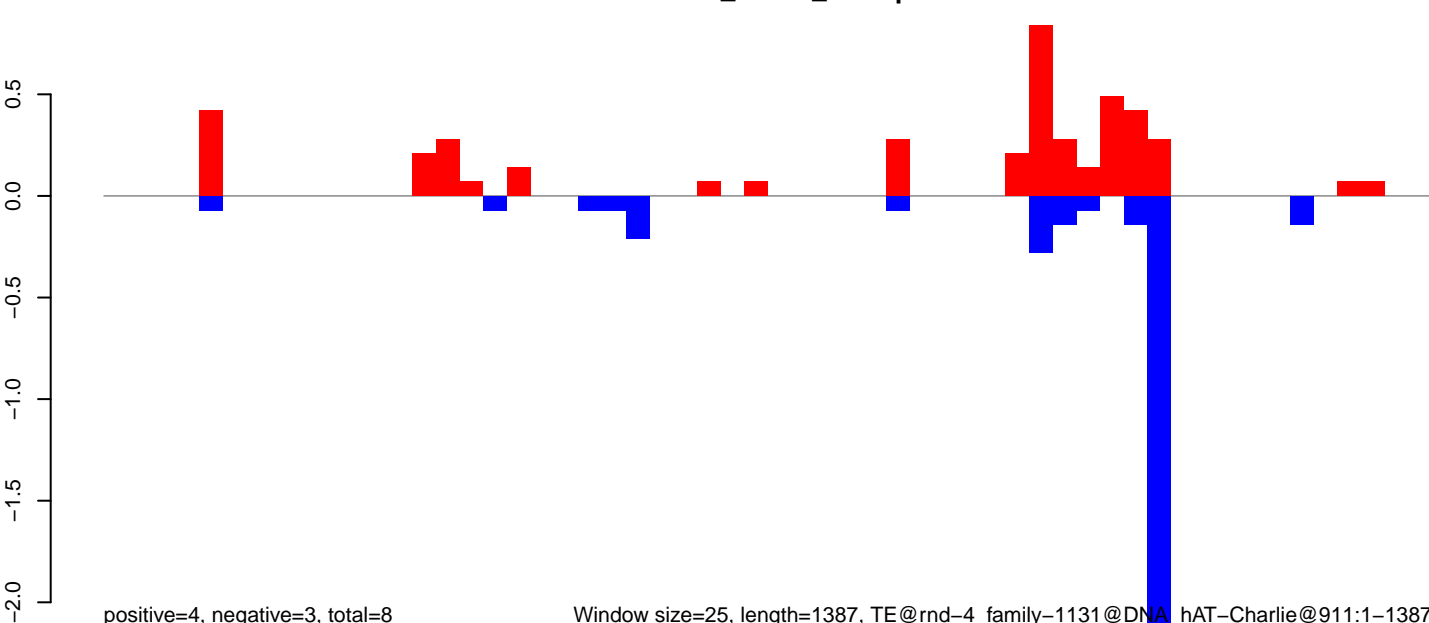
positive=2, negative=1, total=3

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=2, total=4

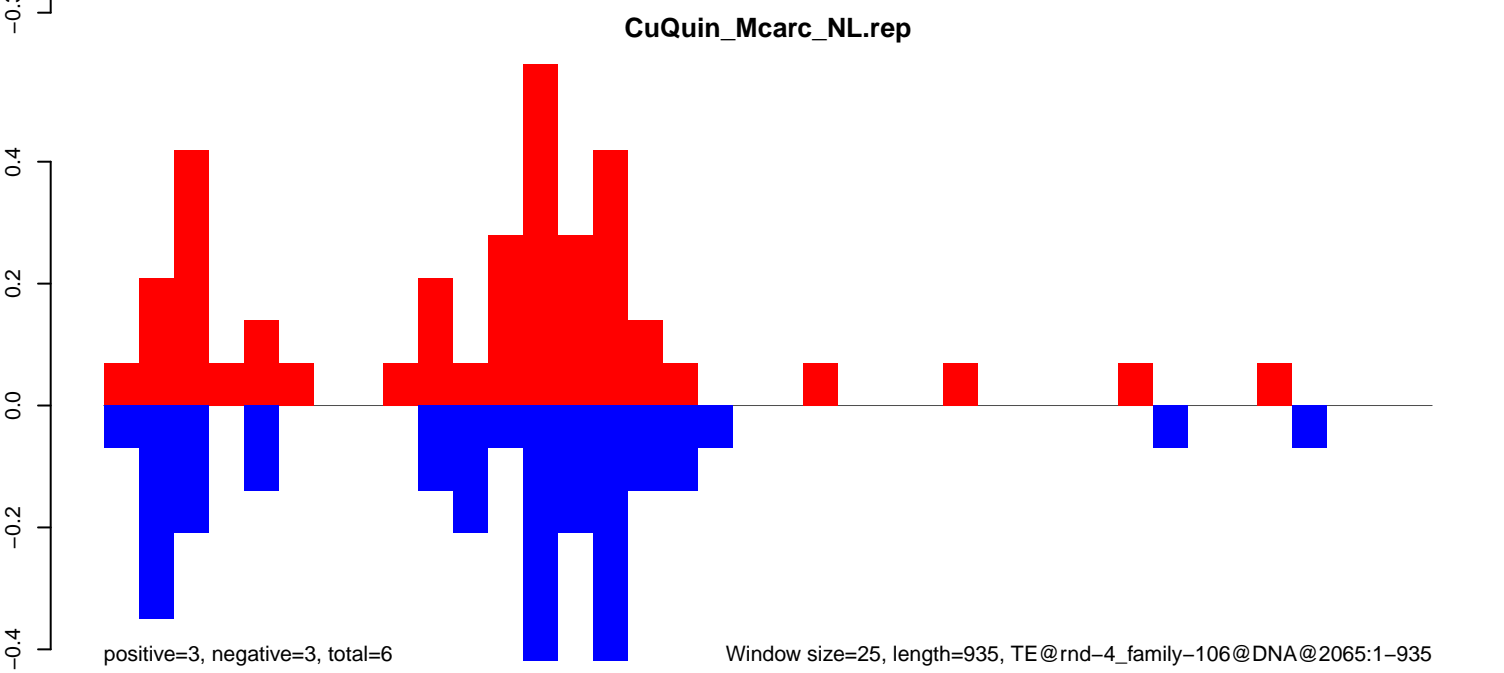
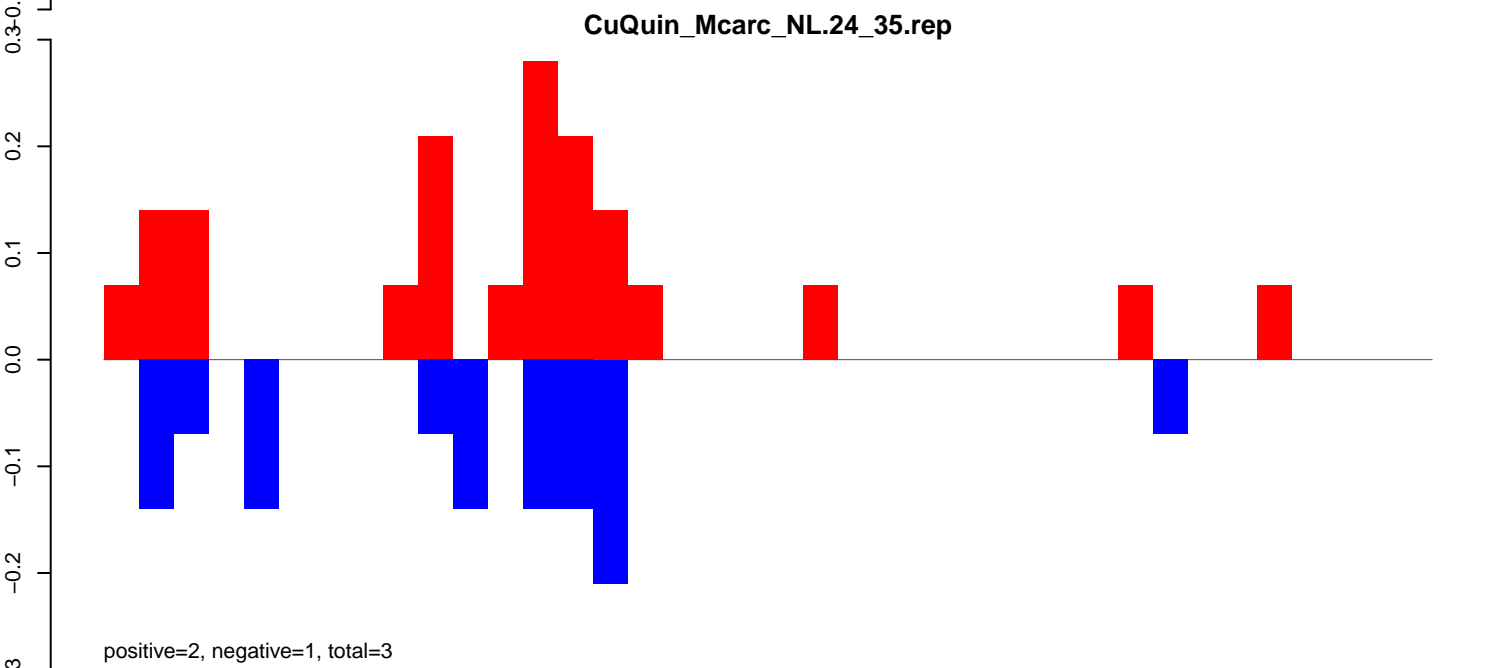
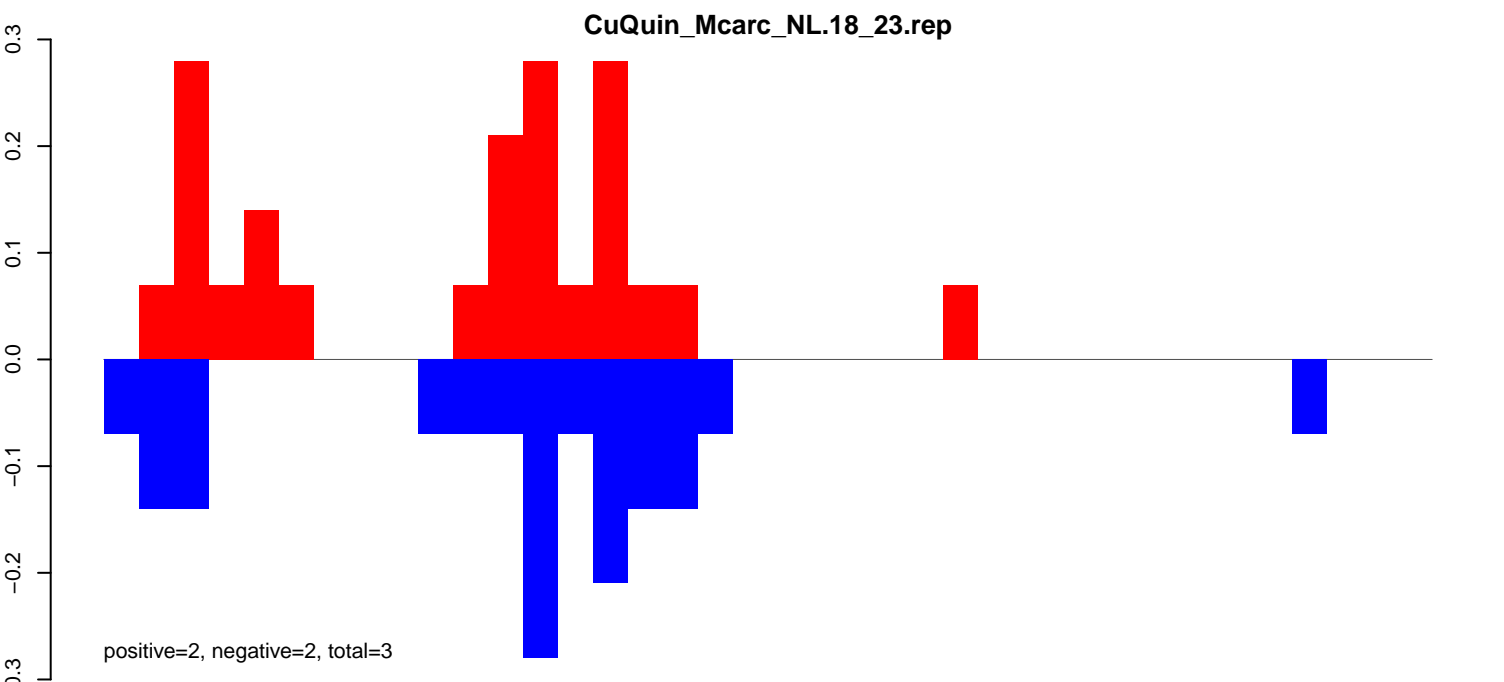
CuQuin_Mcarc_NL.rep



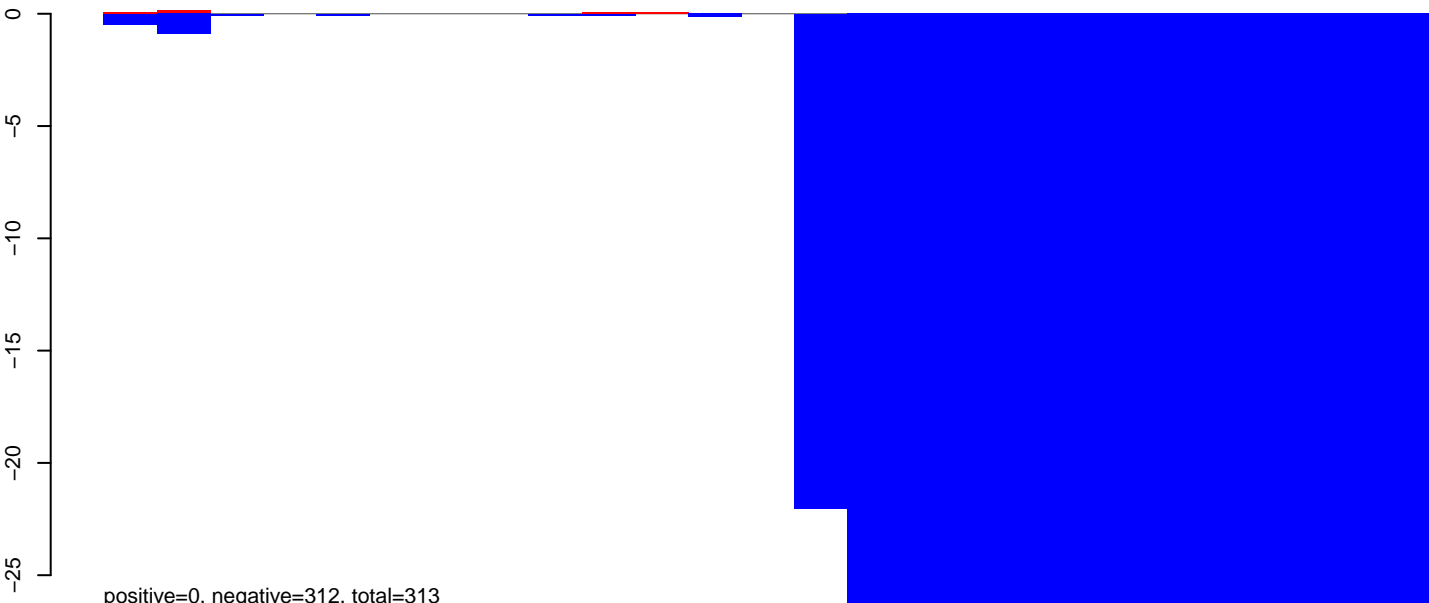
positive=4, negative=3, total=8

Window size=25, length=1387, TE@rnd-4_family-1131@DNA_hAT-Charlie@911:1-1387

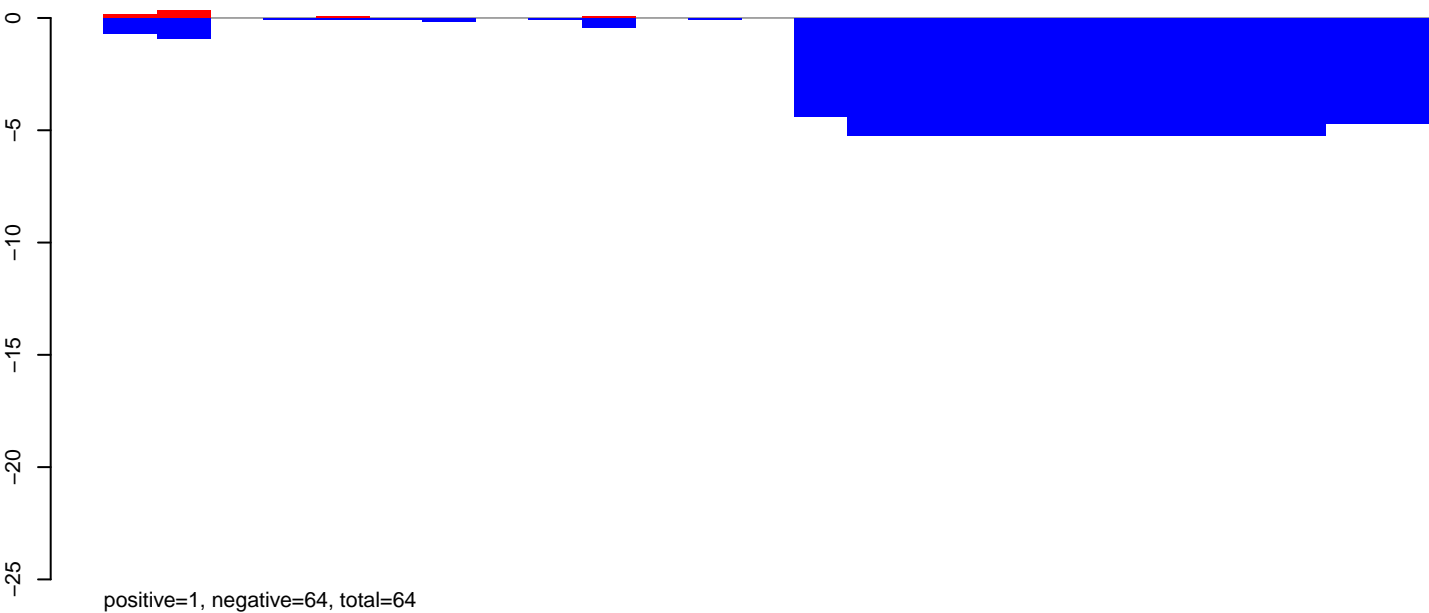
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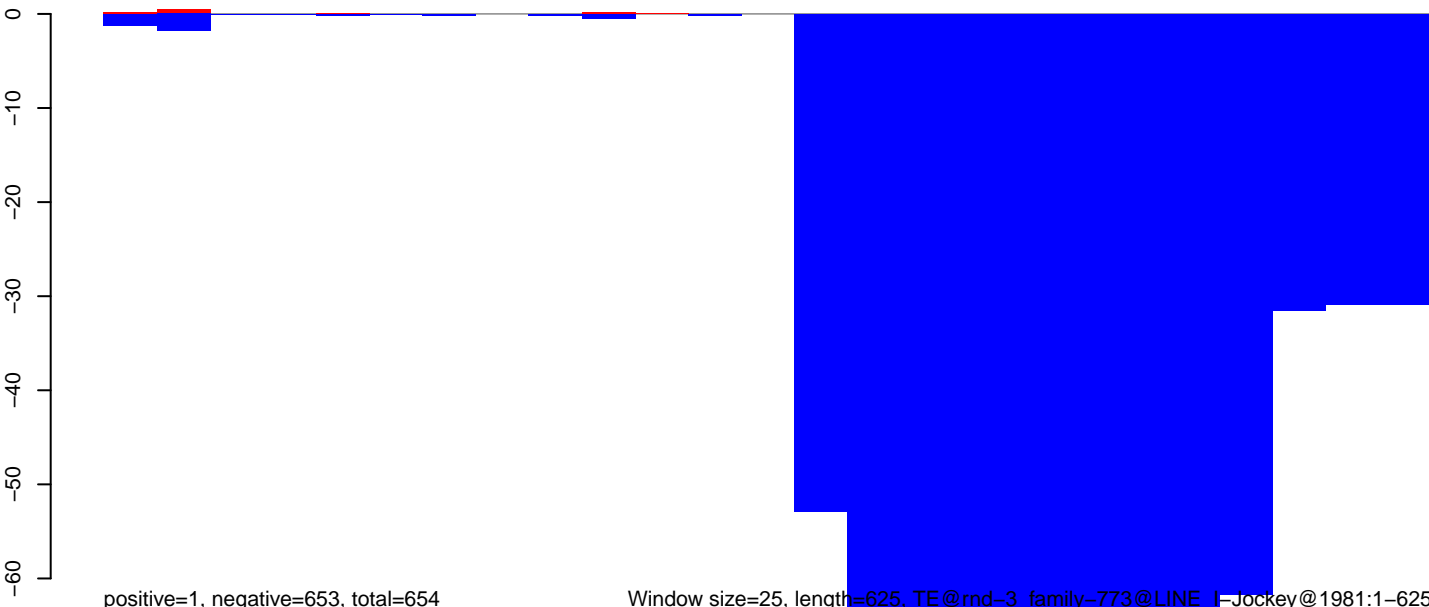
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

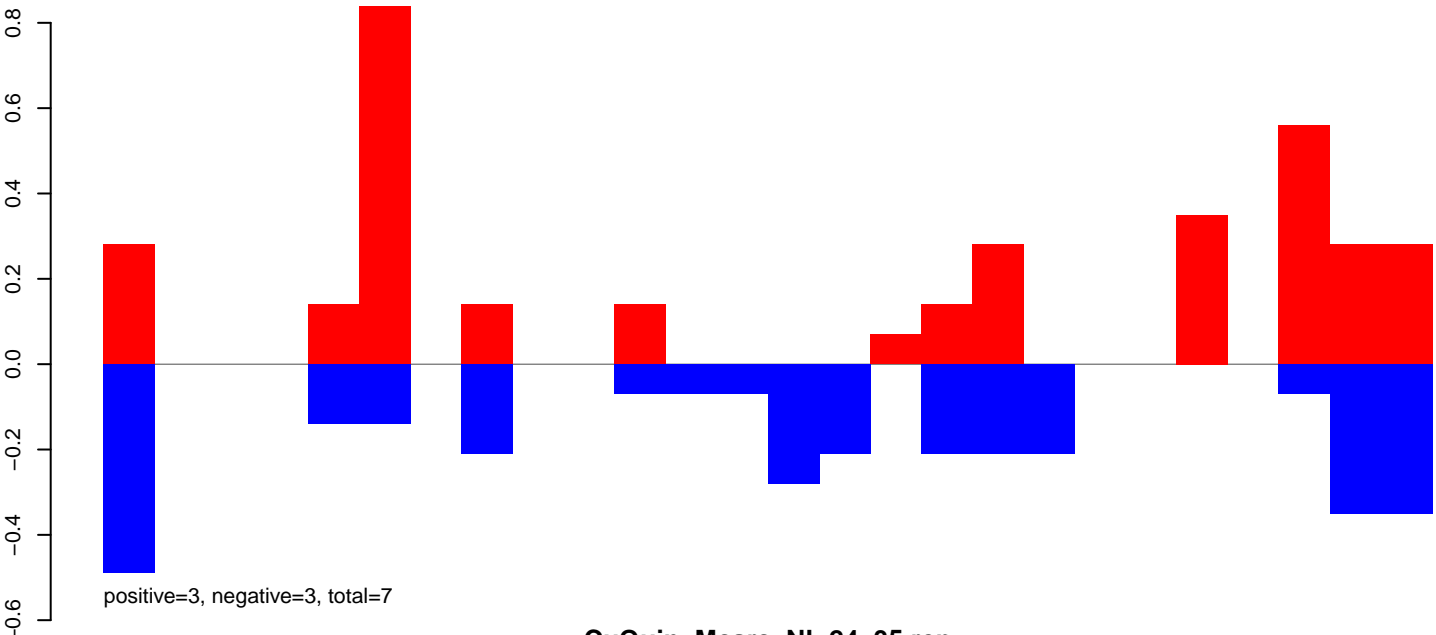


CuQuin_Mcarc_NL.rep

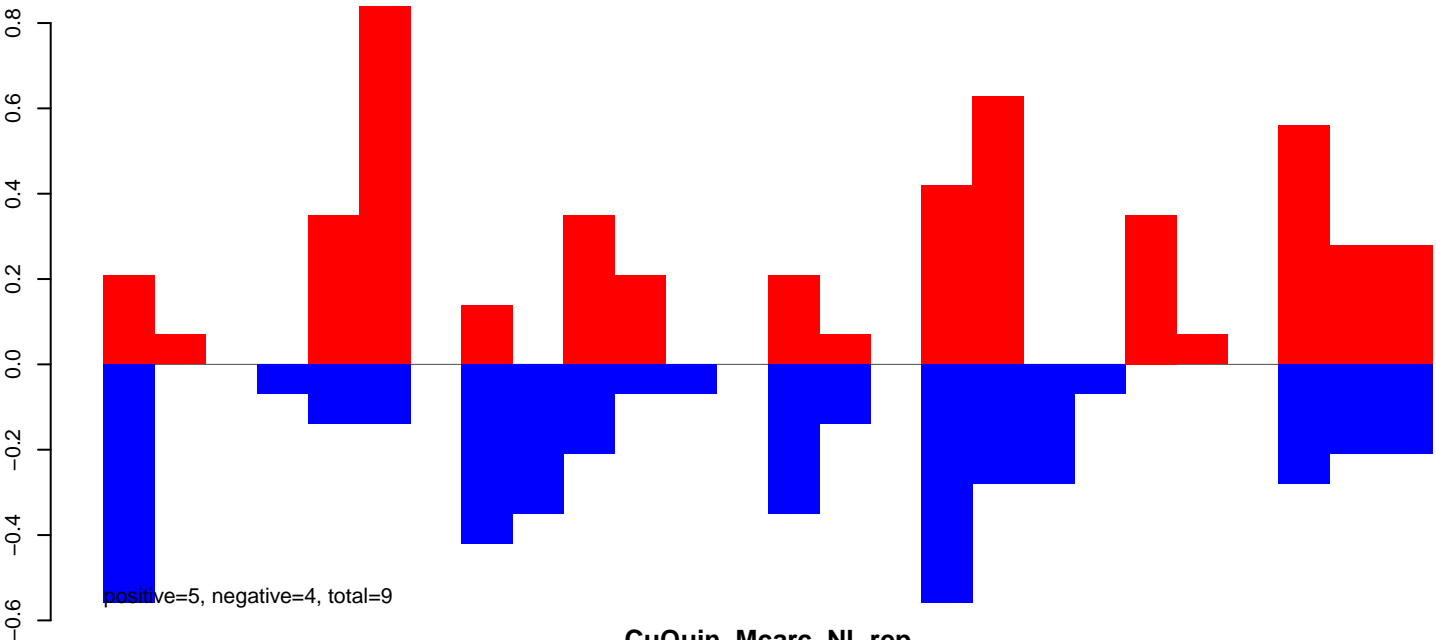


100 200 300 400 500 600

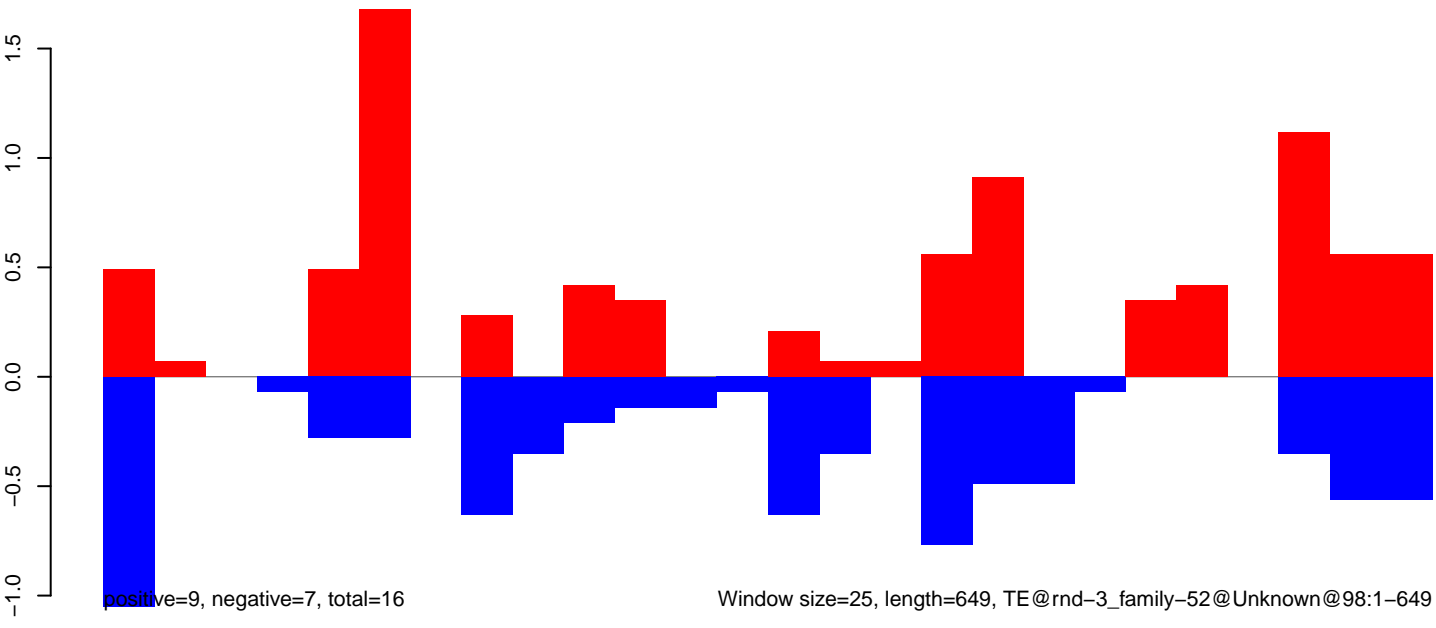
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

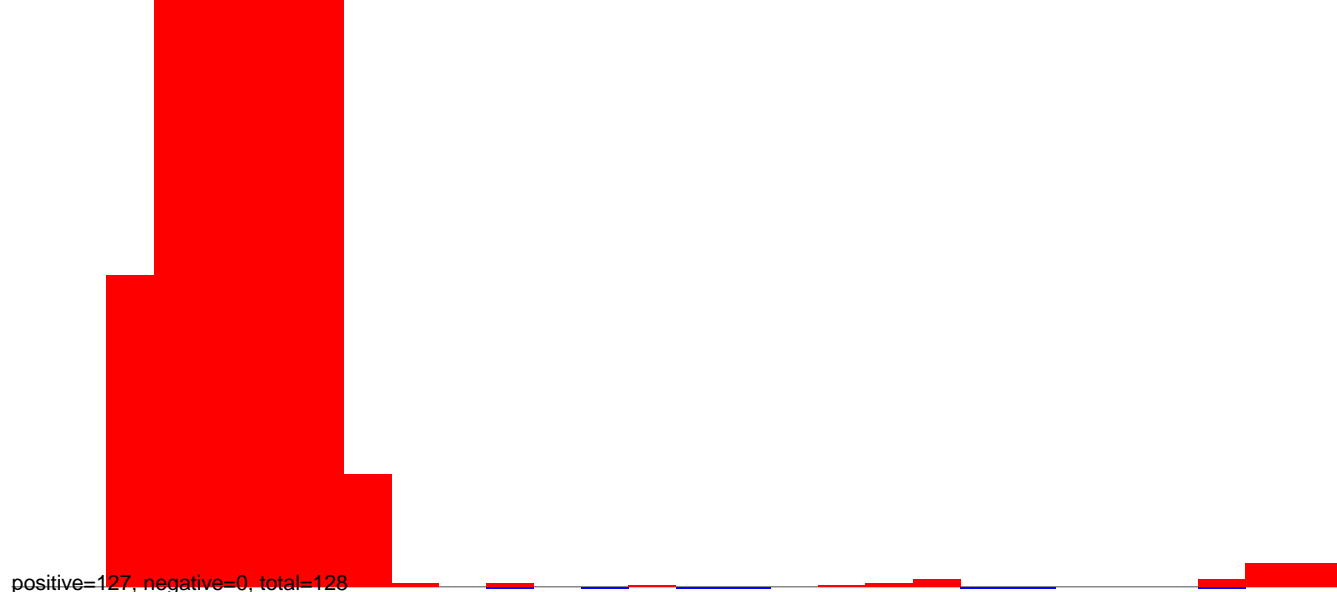


CuQuin_Mcarc_NL.rep

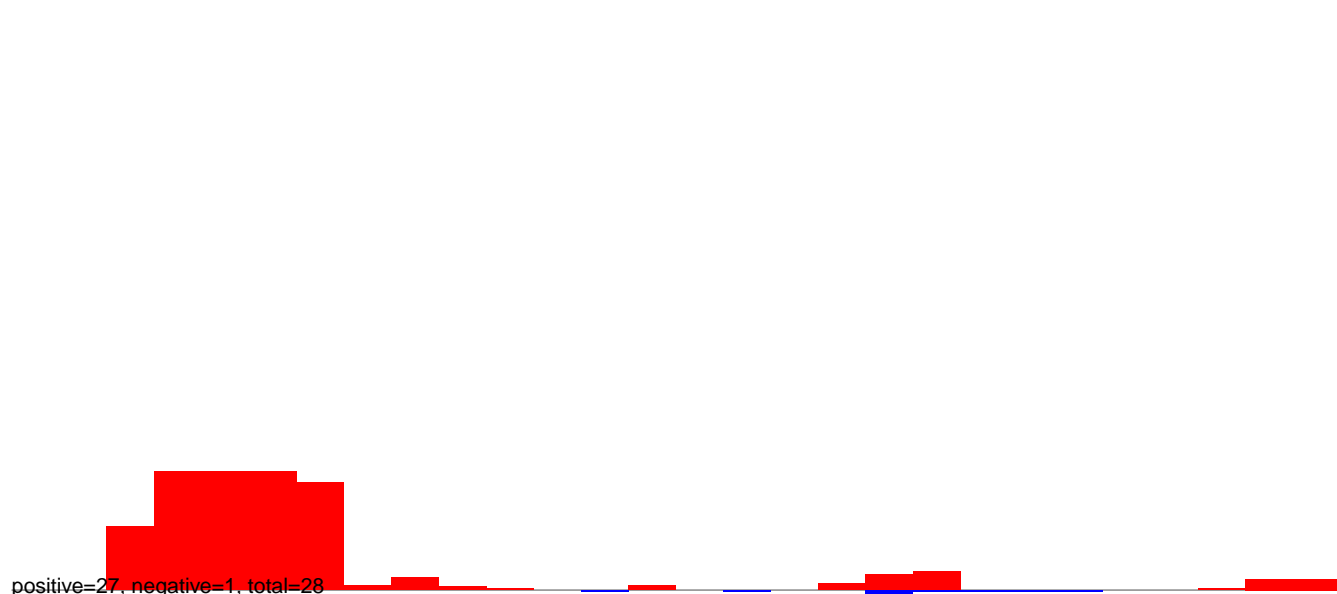


Window size=25, length=649, TE@rnd-3_family-52@Unknown@98:1-649

CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep

0.4
0.2
0.0
-0.2
-0.4

positive=1, negative=1, total=2

CuQuin_Mcarc_NL.24_35.rep

0.4
0.2
0.0
-0.2
-0.4

positive=1, negative=1, total=2

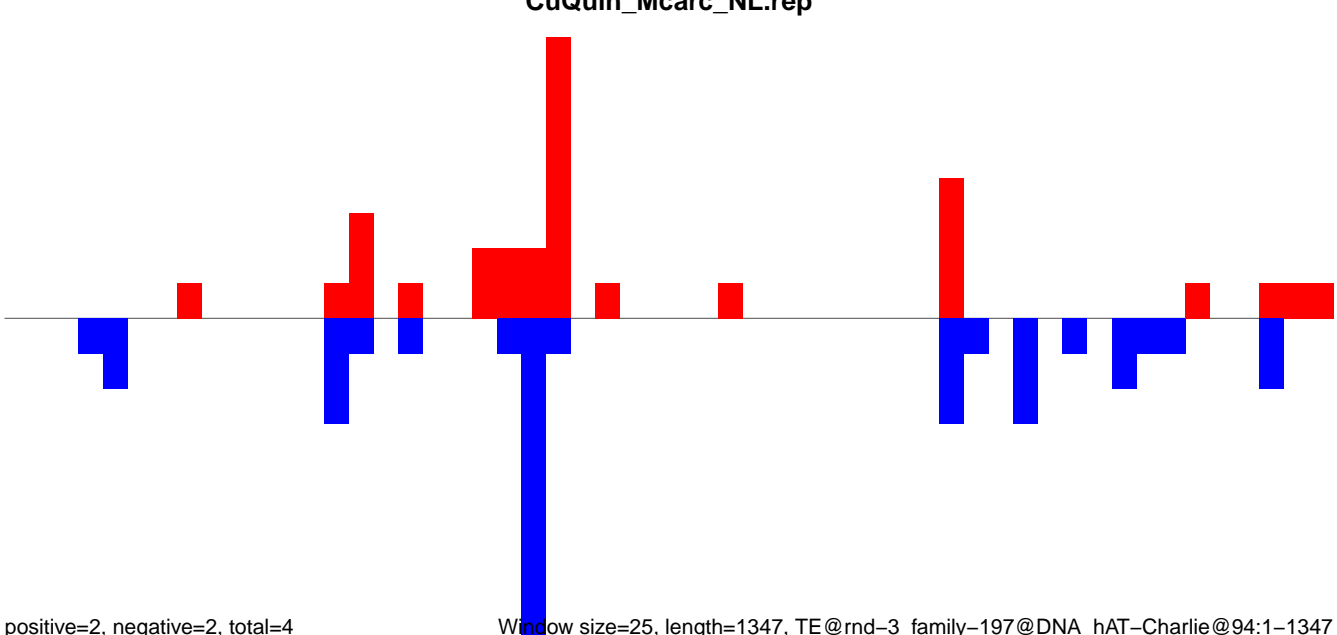
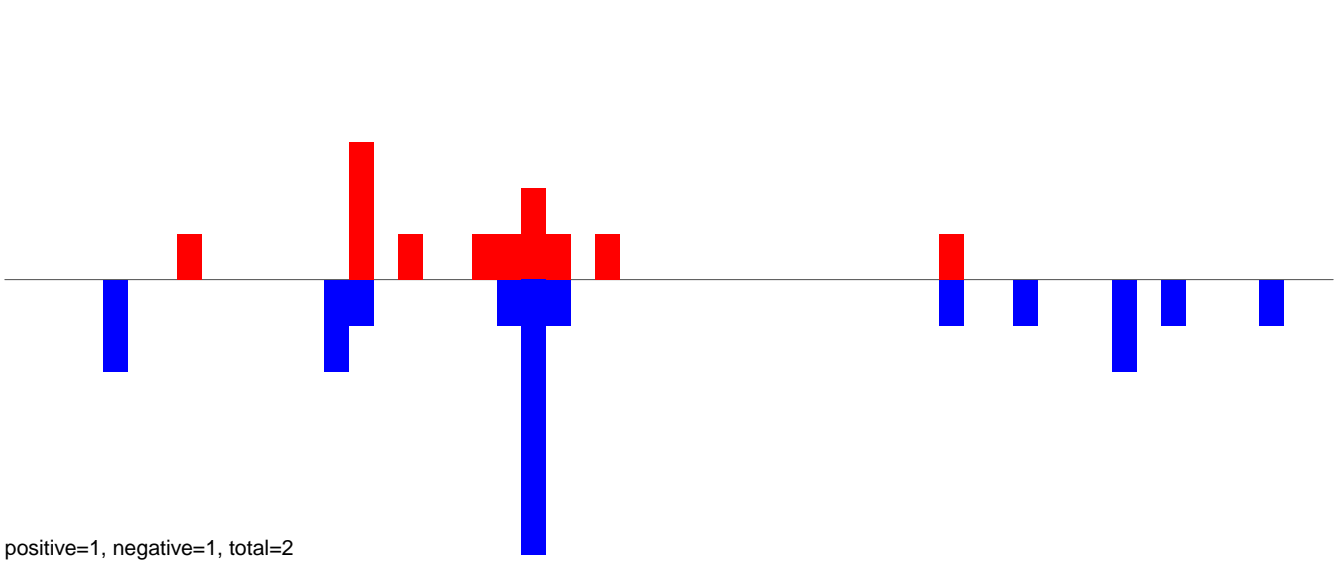
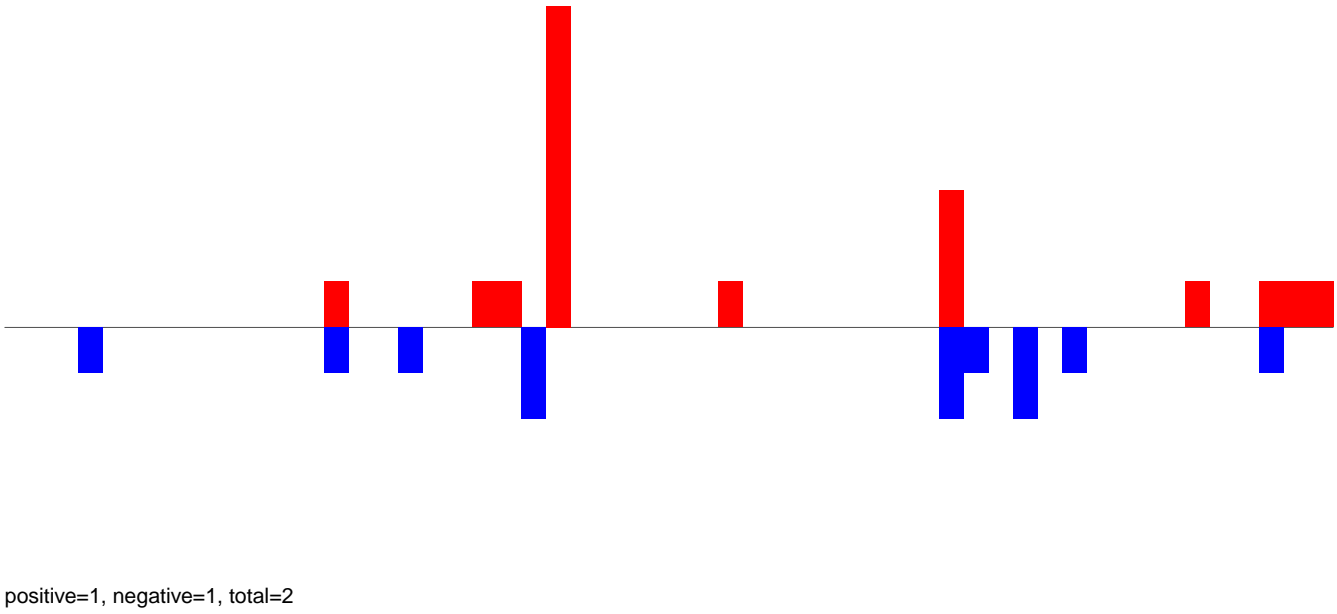
CuQuin_Mcarc_NL.rep

0.6
0.4
0.2
0.0
-0.2
-0.4
-0.6

positive=2, negative=2, total=4

Window size=25, length=1347, TE@rnd-3_family-197@DNA_hAT-Charlie@94:1-1347

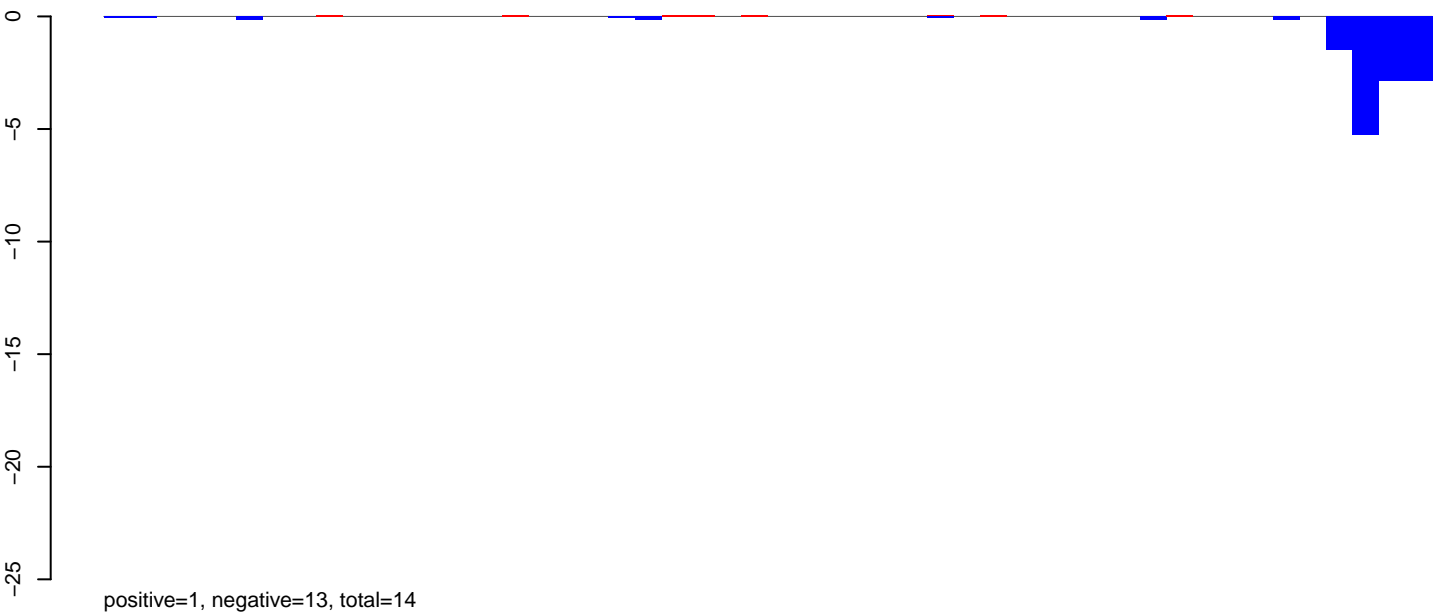
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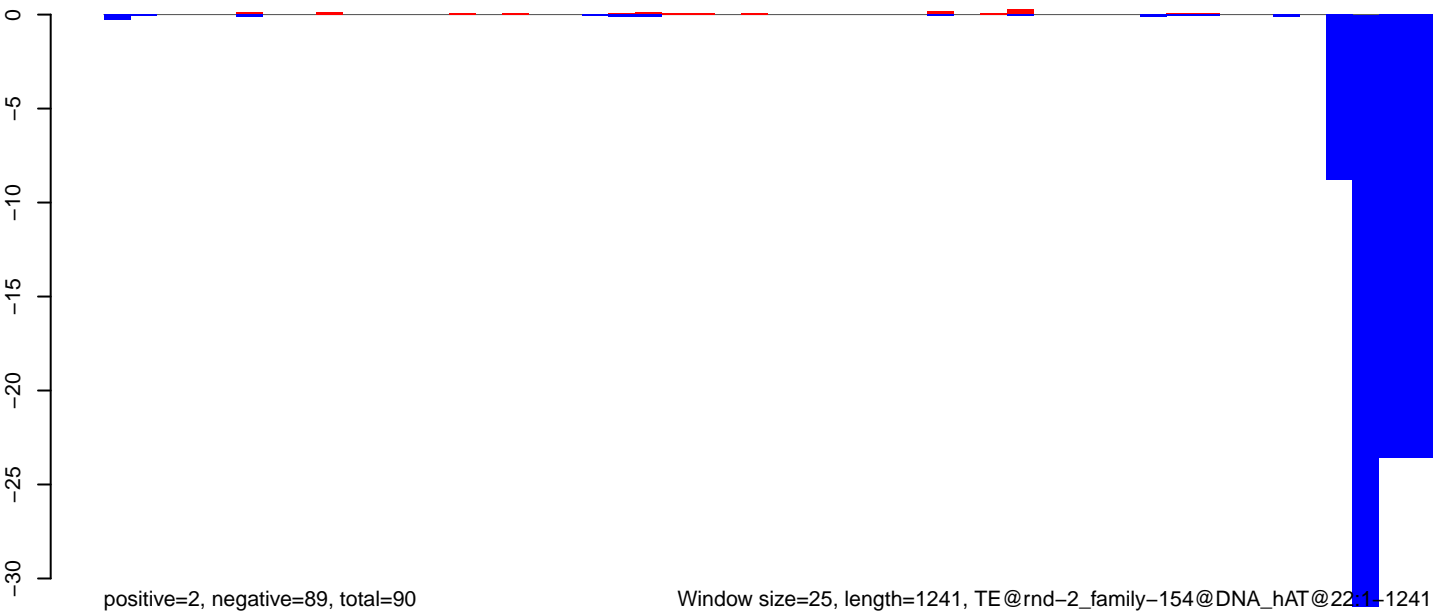
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

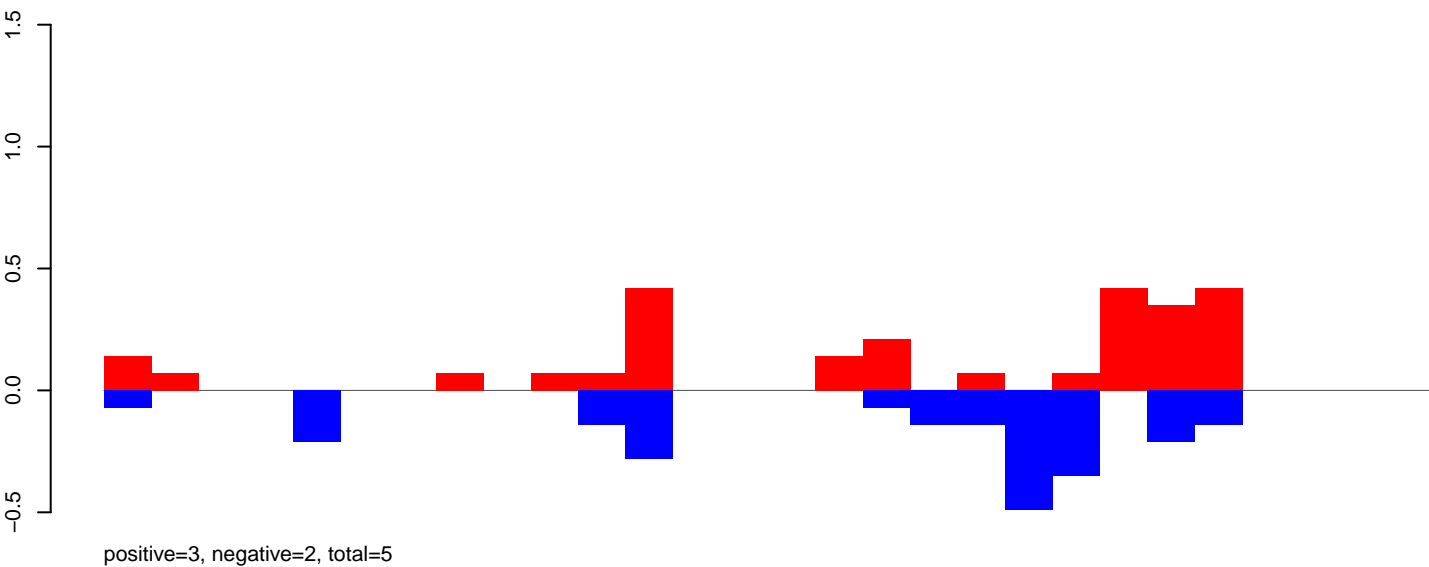


CuQuin_Mcarc_NL.rep

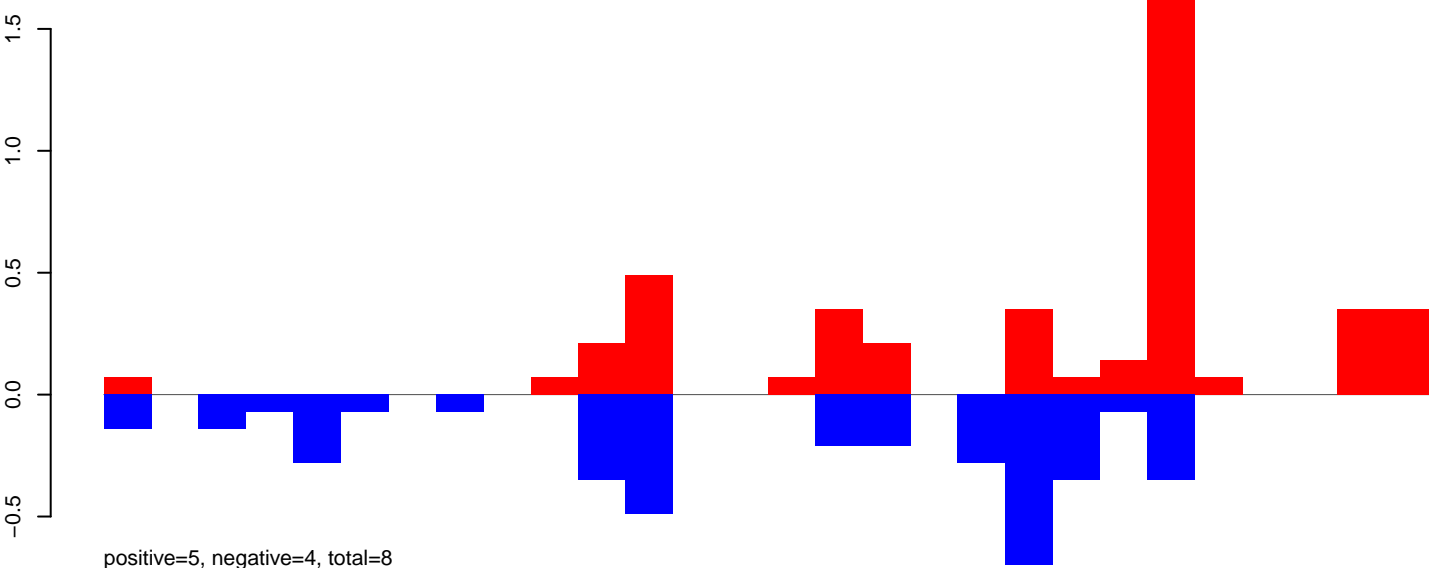


Window size=25, length=1241, TE@rnd-2_family-154@DNA_hAT@221-1241

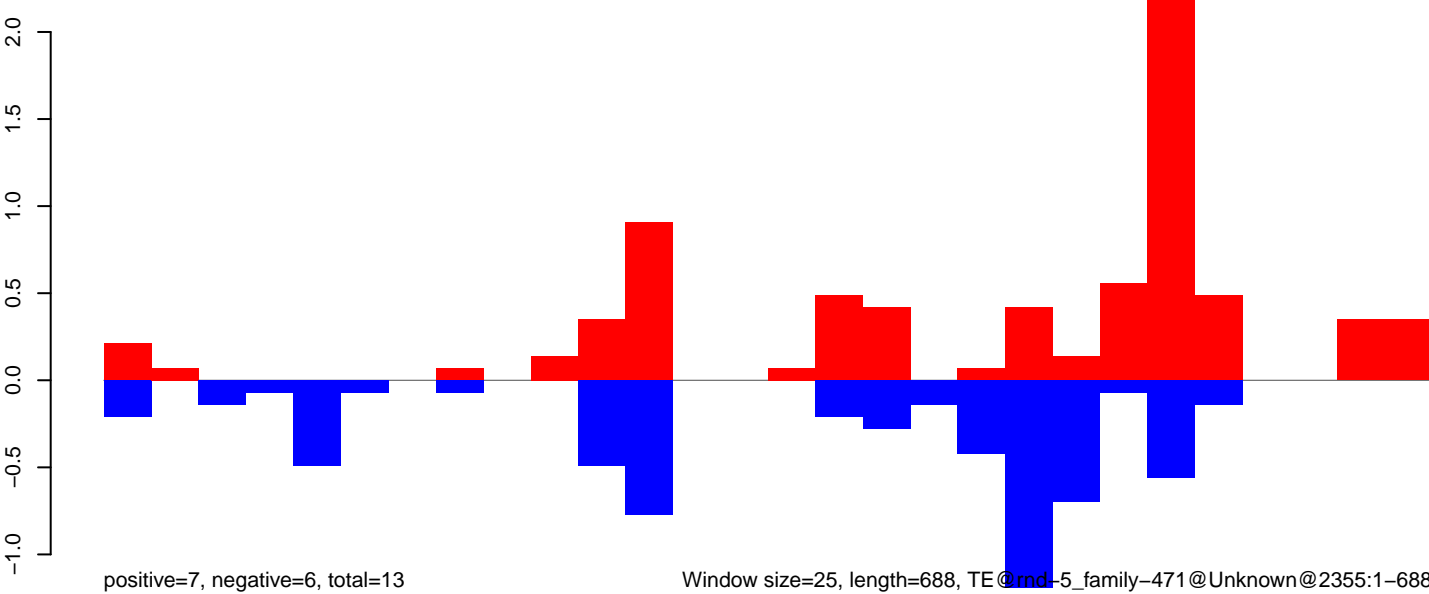
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

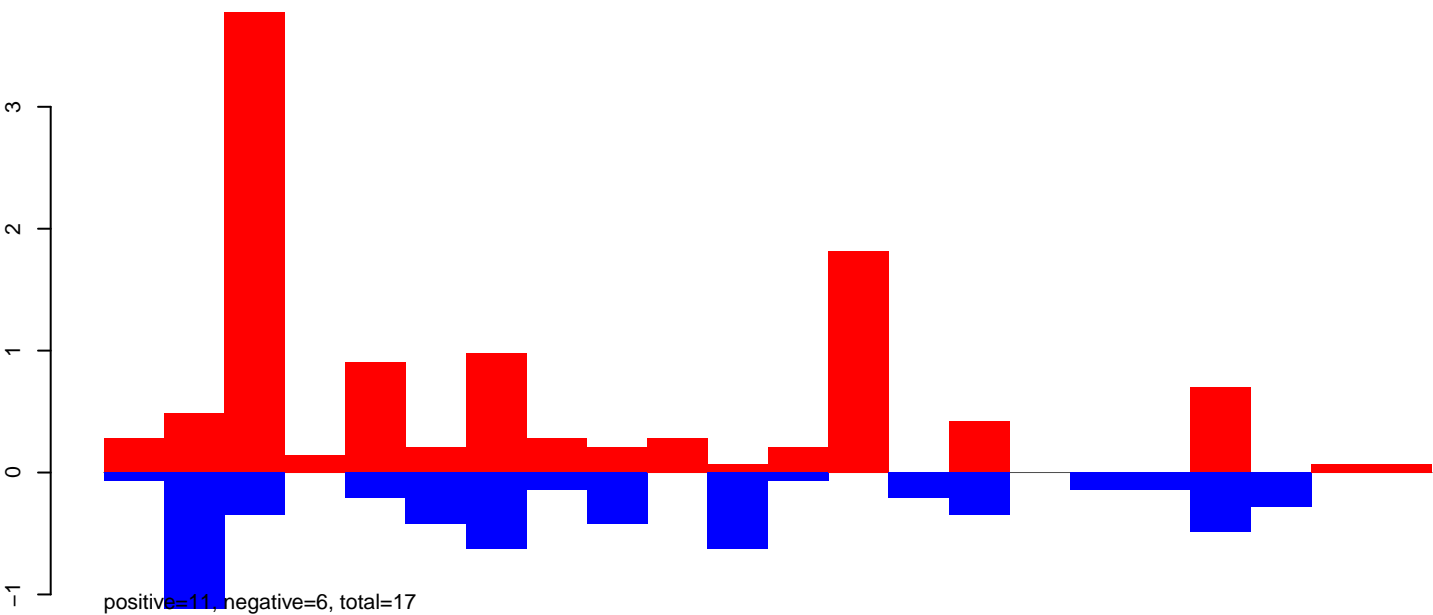


CuQuin_Mcarc_NL.rep

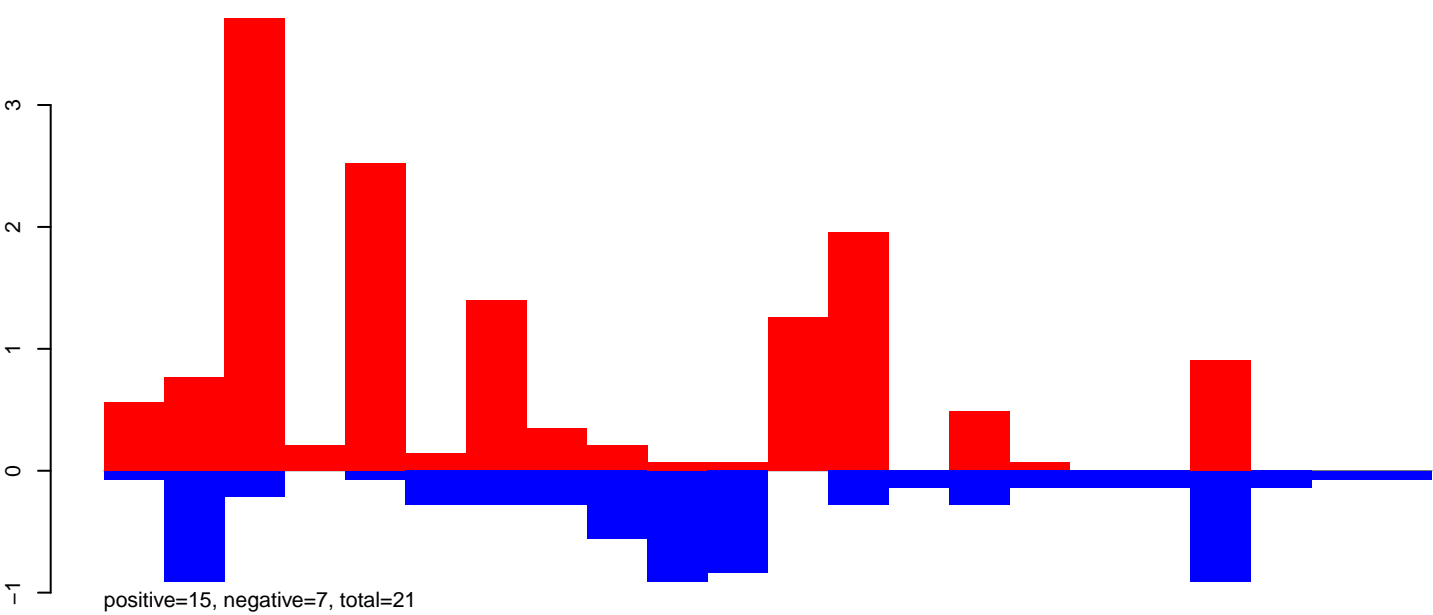


Window size=25, length=688, TE@rmd-5_family-471@Unknown@2355:1-688

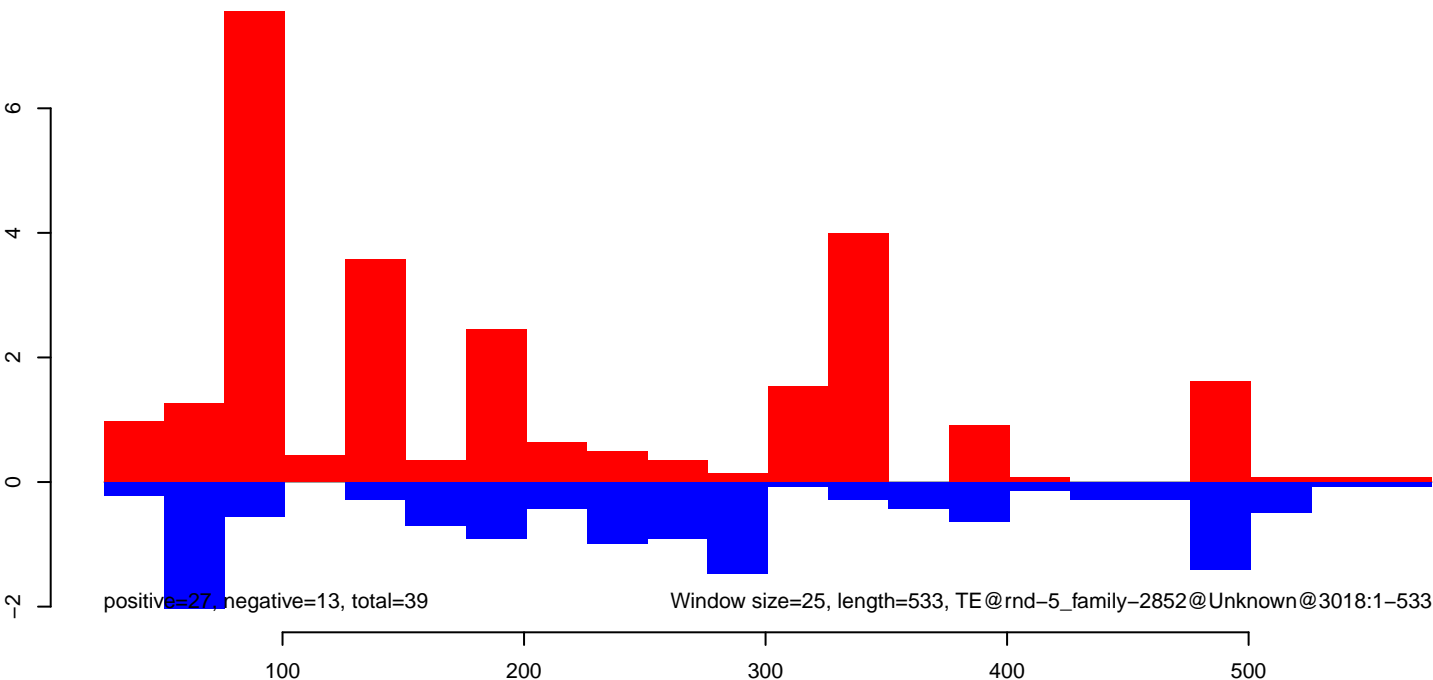
CuQuin_Mcarc_NL.18_23.rep



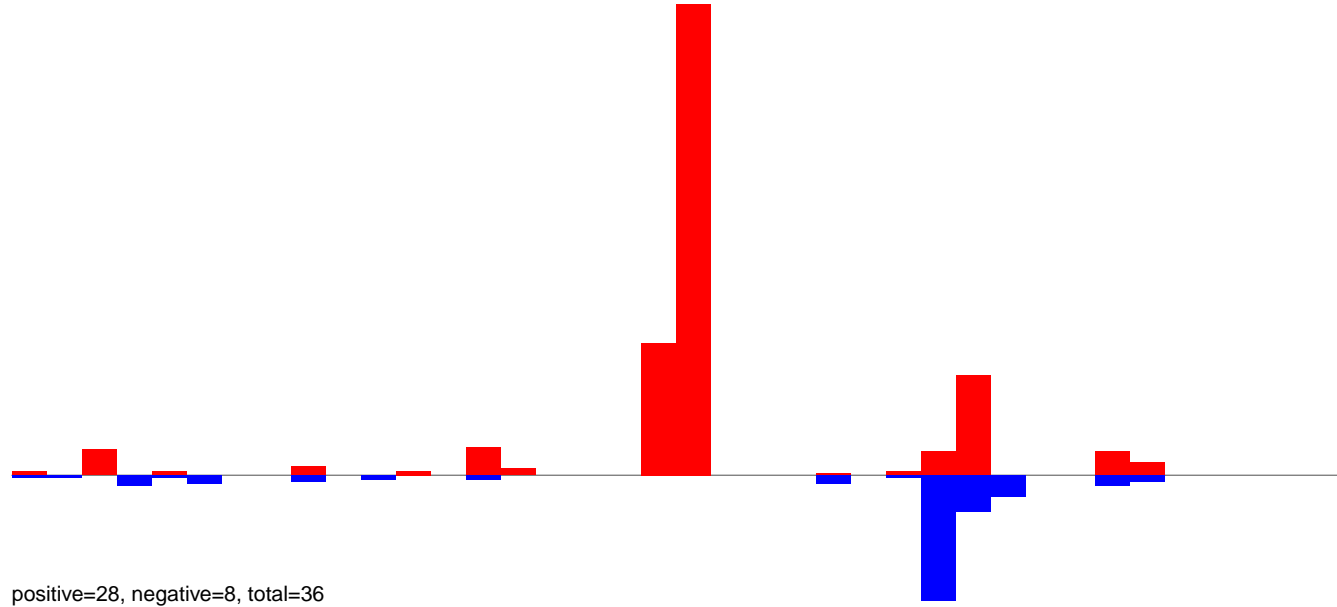
CuQuin_Mcarc_NL.24_35.rep



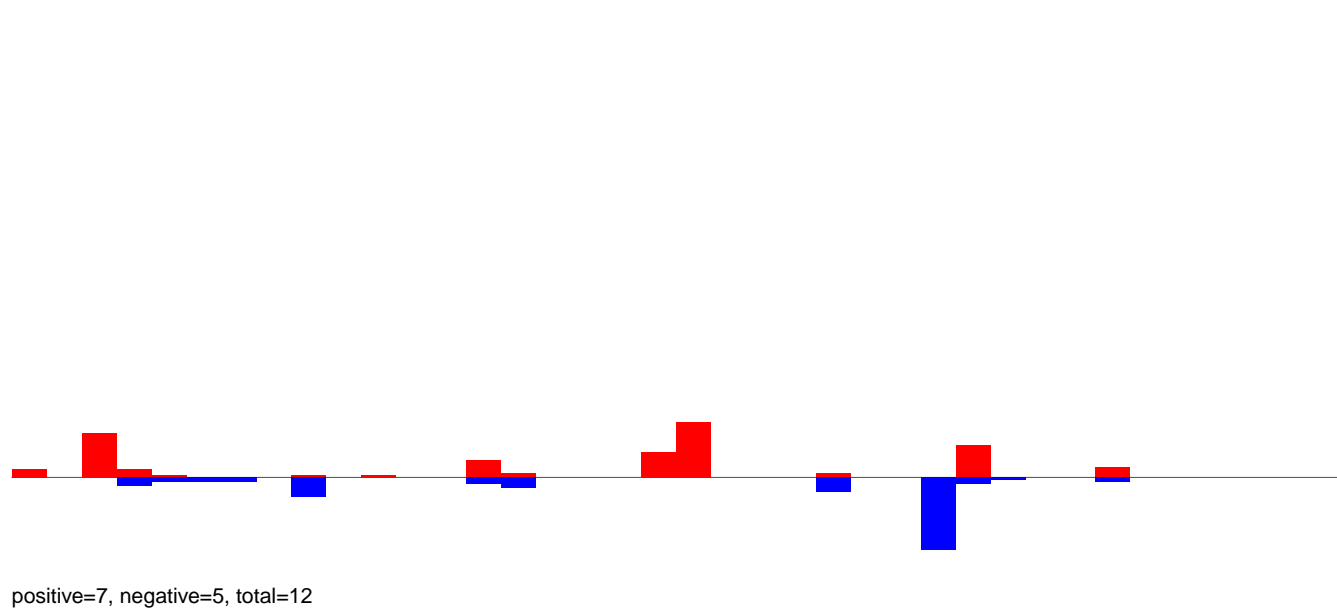
CuQuin_Mcarc_NL.rep



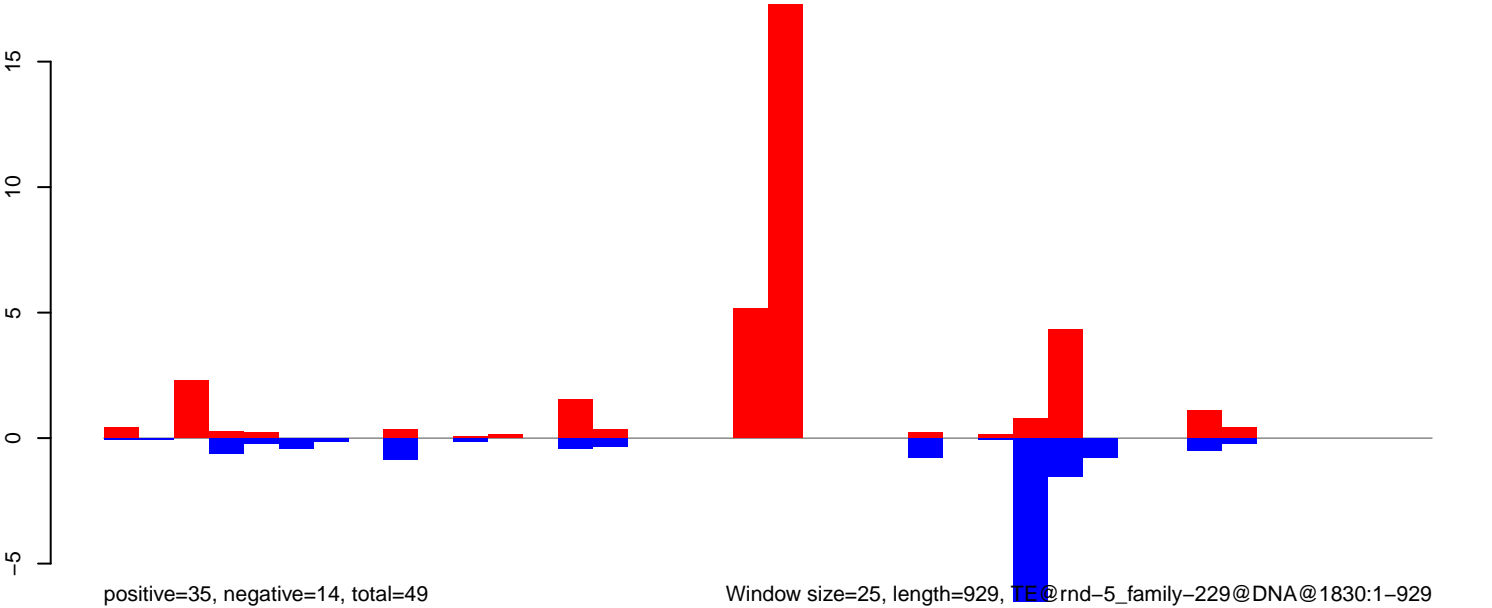
CuQuin_Mcarc_NL.18_23.rep



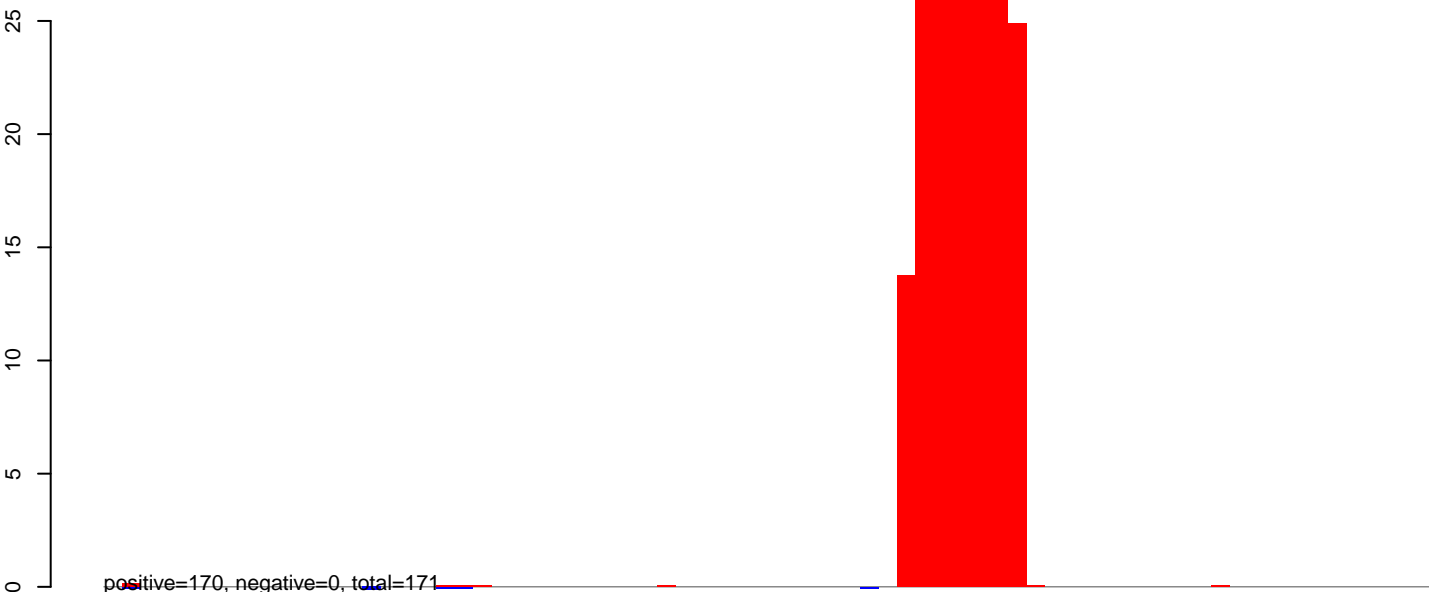
CuQuin_Mcarc_NL.24_35.rep



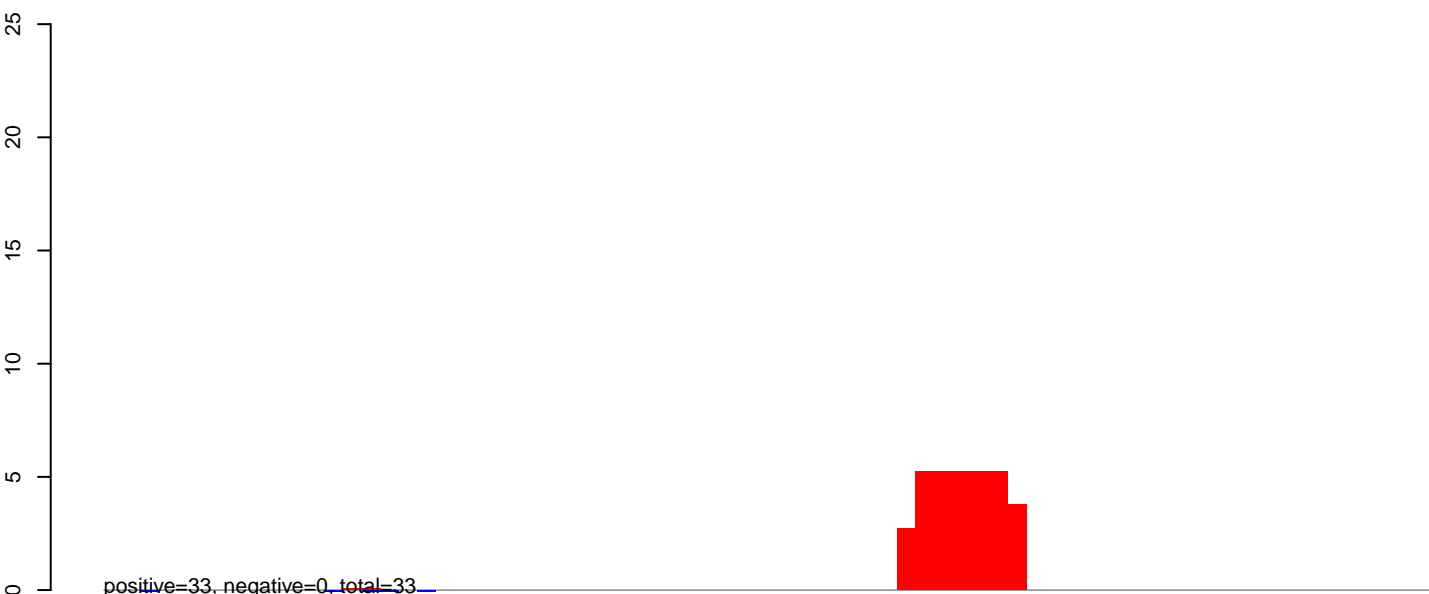
CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep



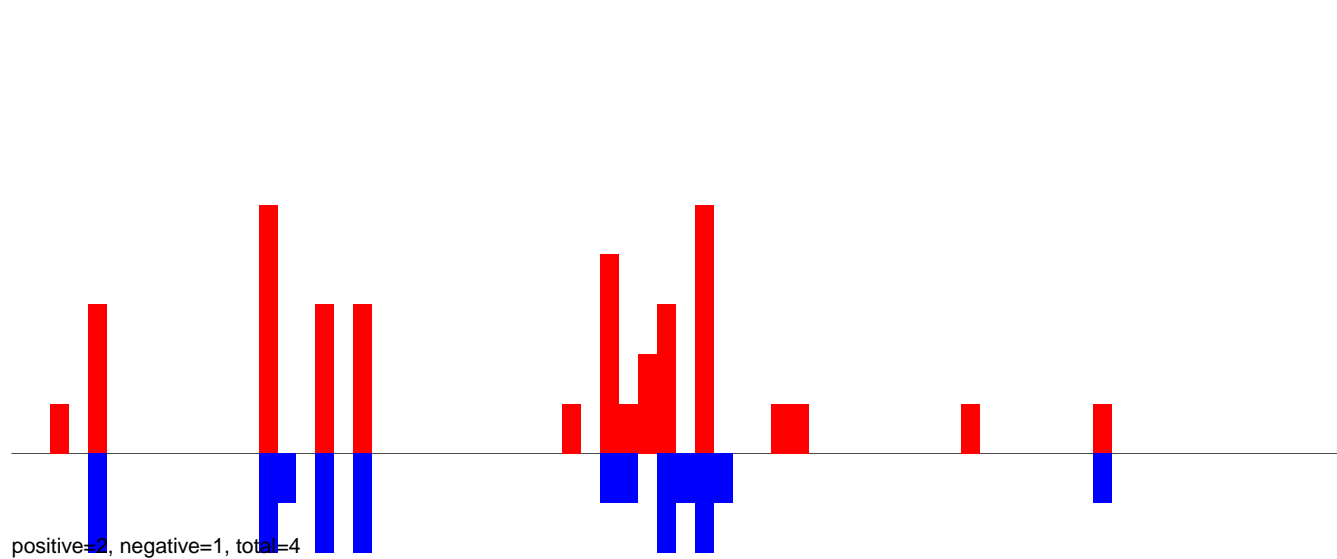
CuQuin_Mcarc_NL.24_35.rep



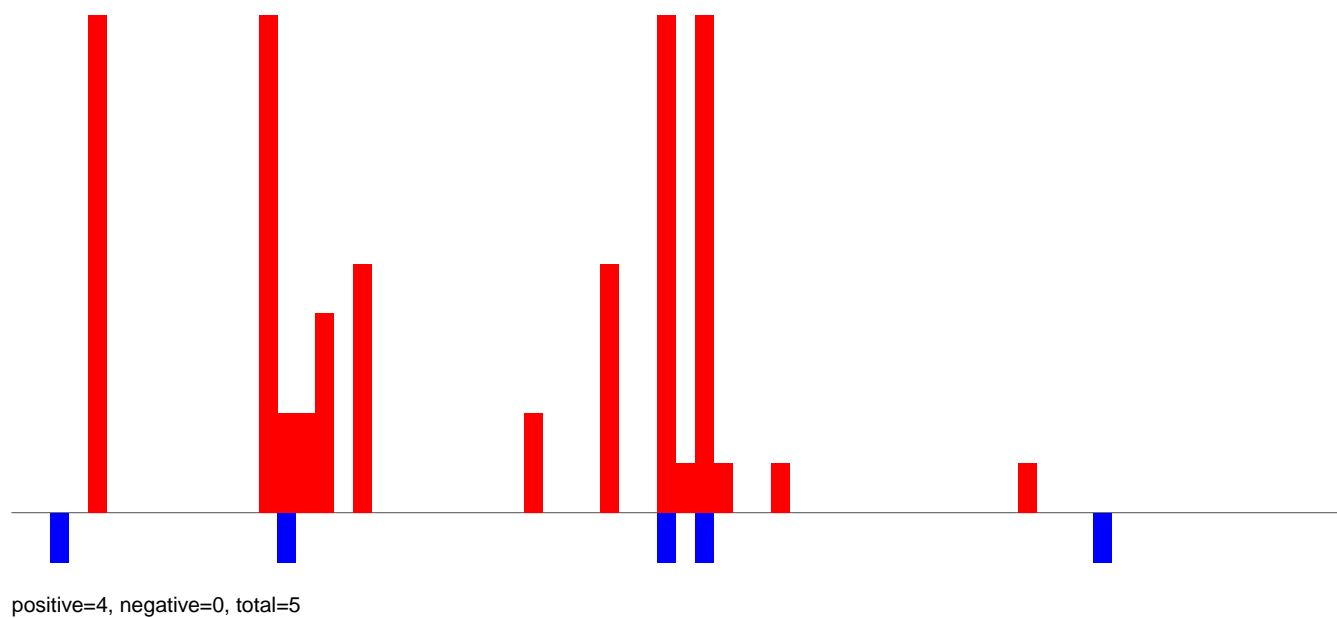
CuQuin_Mcarc_NL.rep



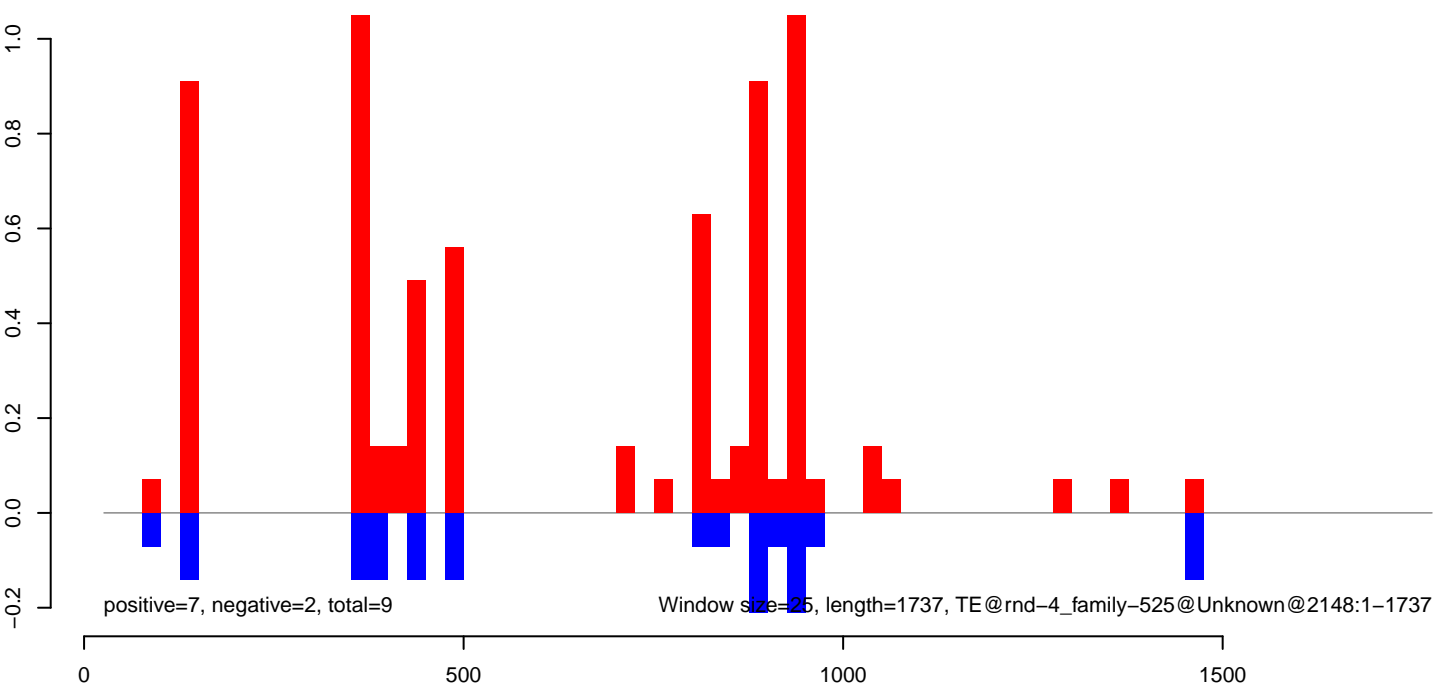
CuQuin_Mcarc_NL.18_23.rep



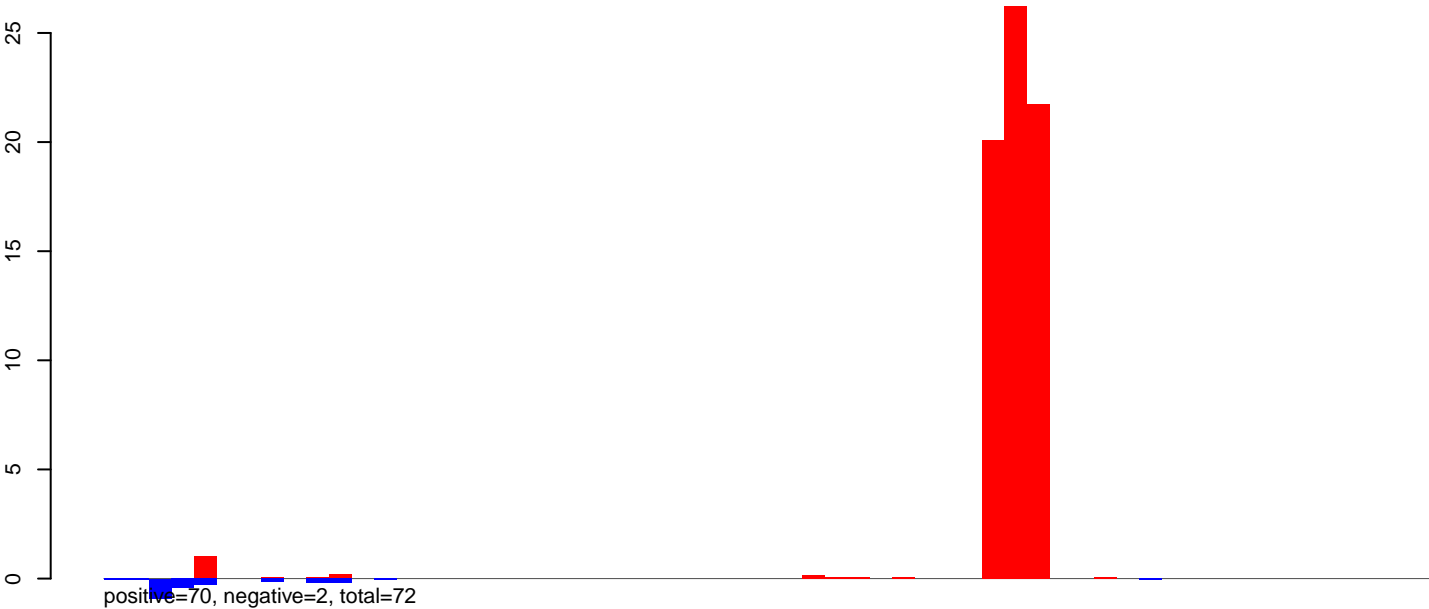
CuQuin_Mcarc_NL.24_35.rep



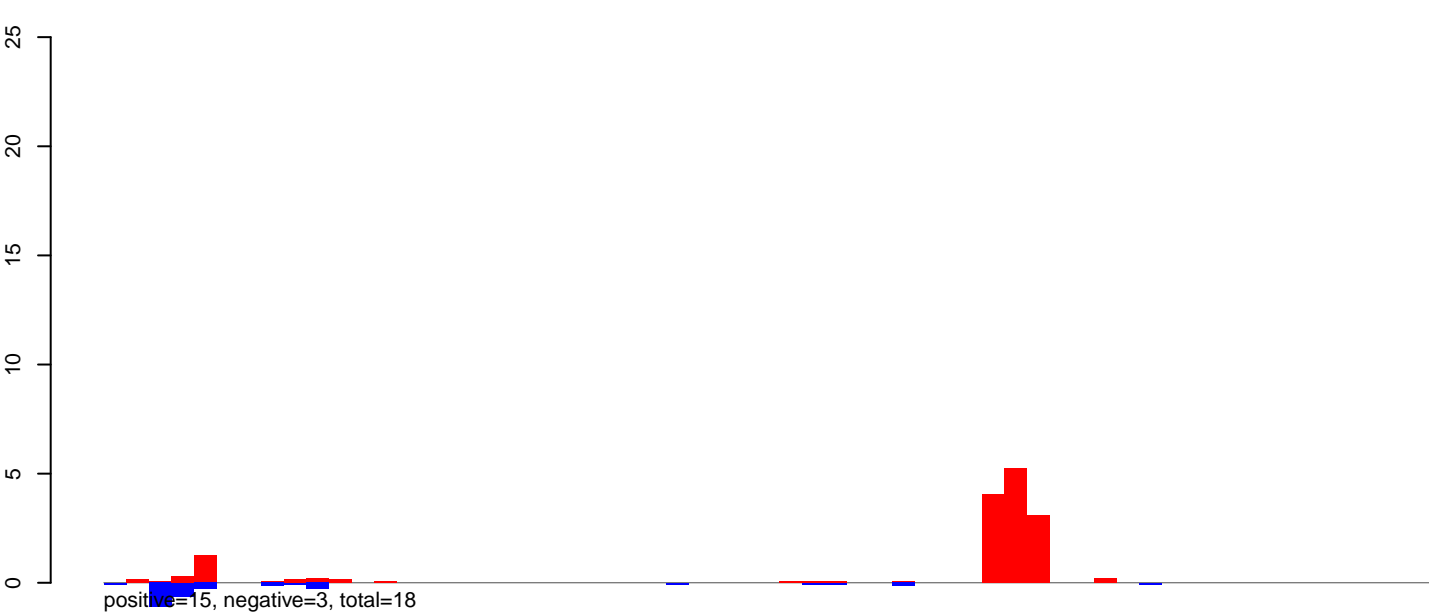
CuQuin_Mcarc_NL.rep



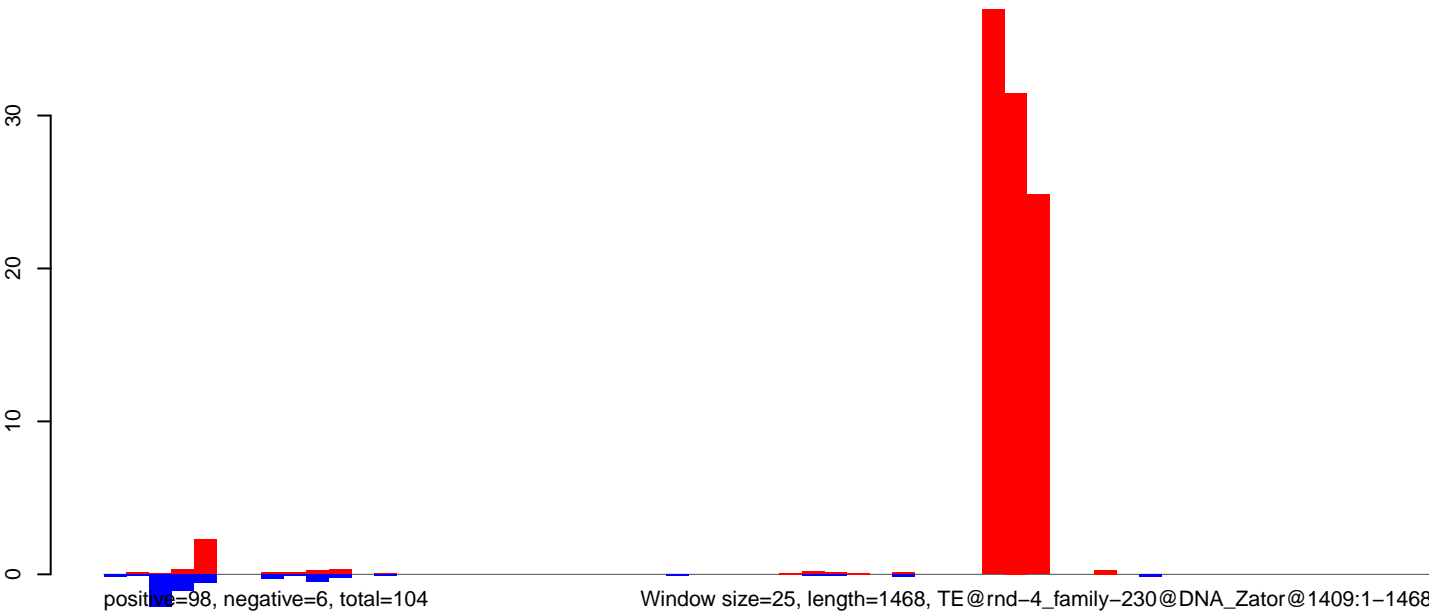
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

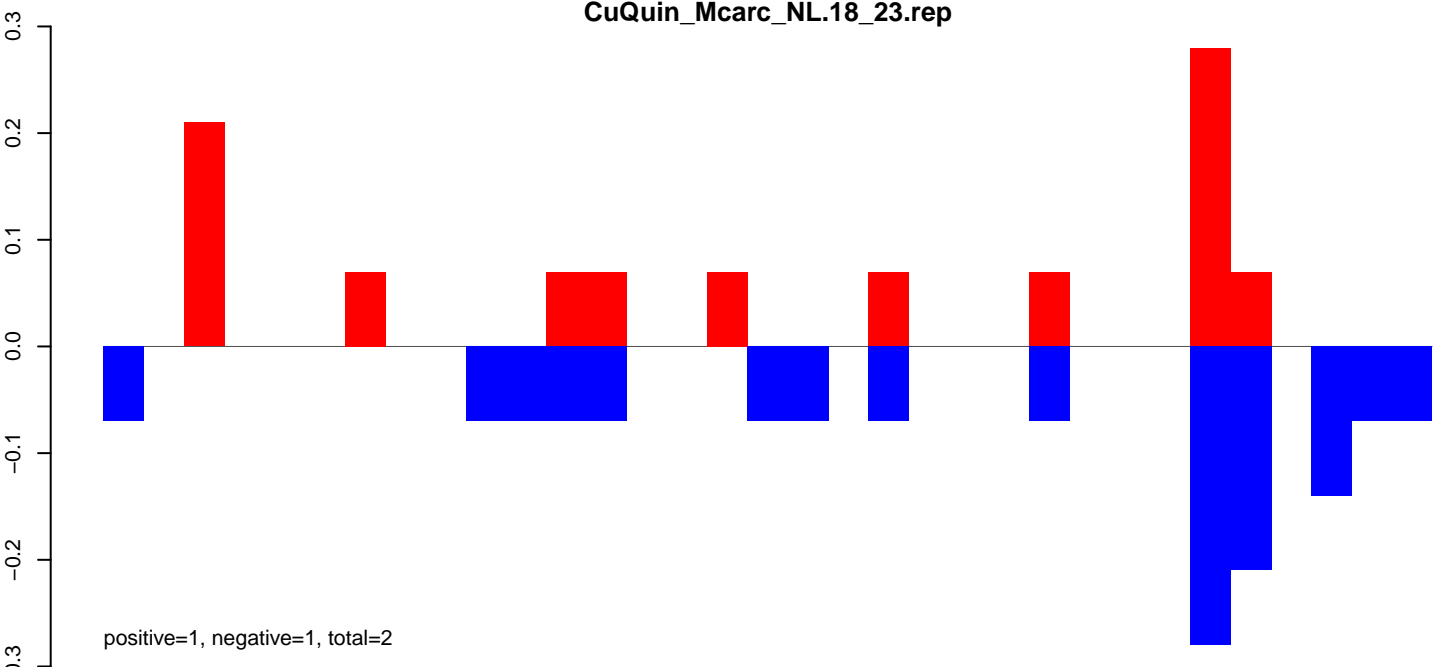


CuQuin_Mcarc_NL.rep

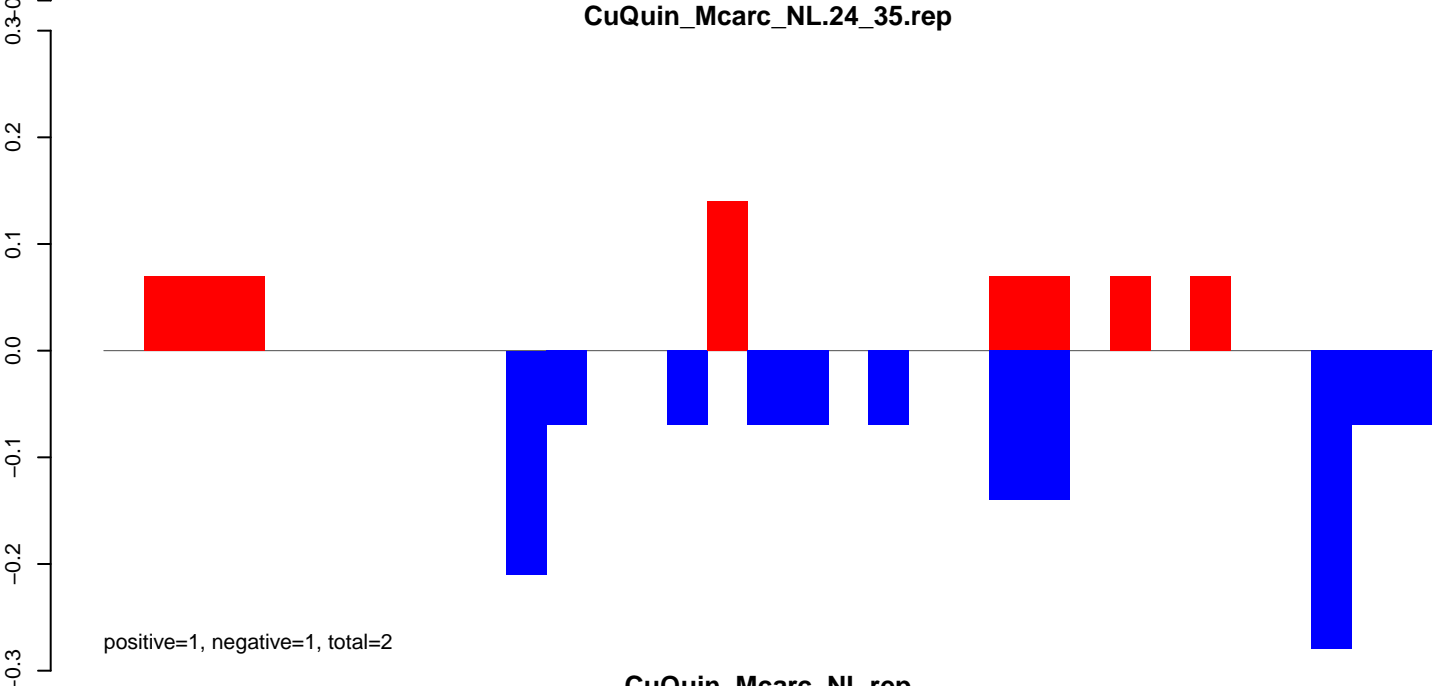


Window size=25, length=1468, TE@rnd-4_family-230@DNA_Zator@1409:1-1468

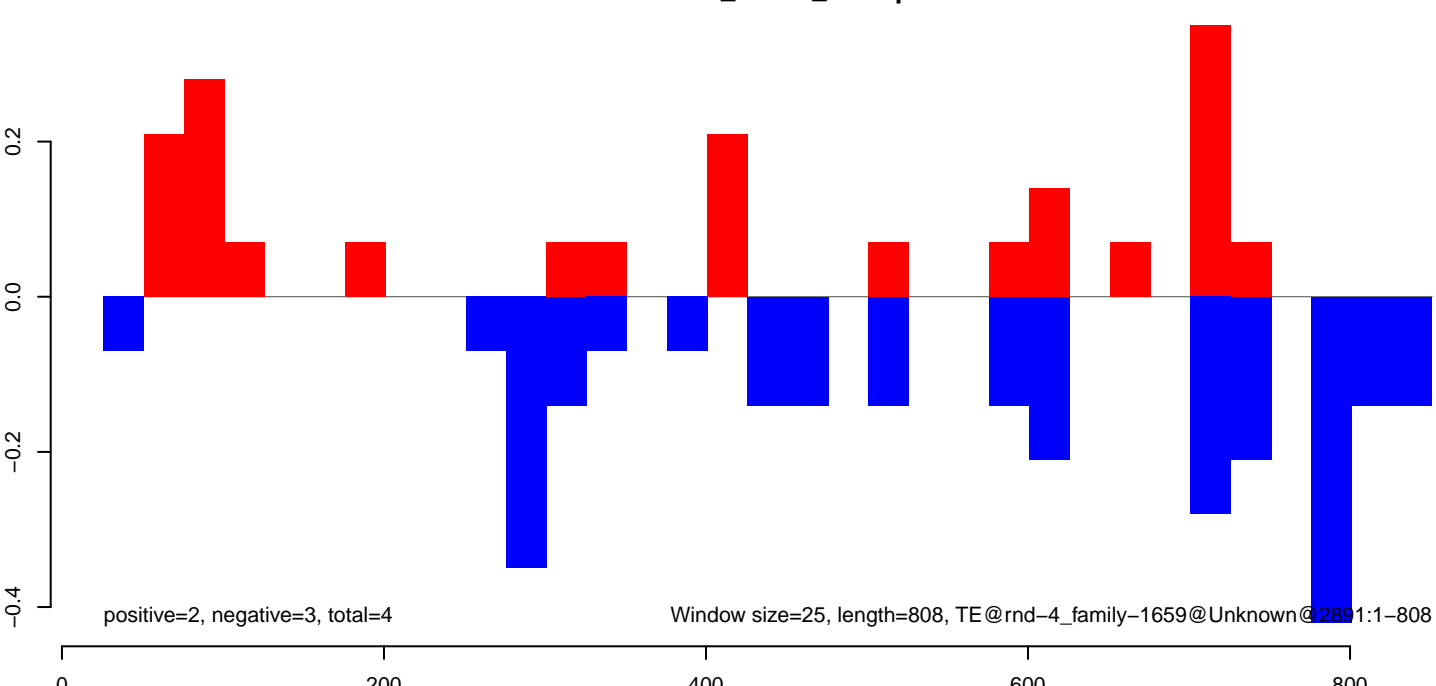
CuQuin_Mcarc_NL.18_23.rep

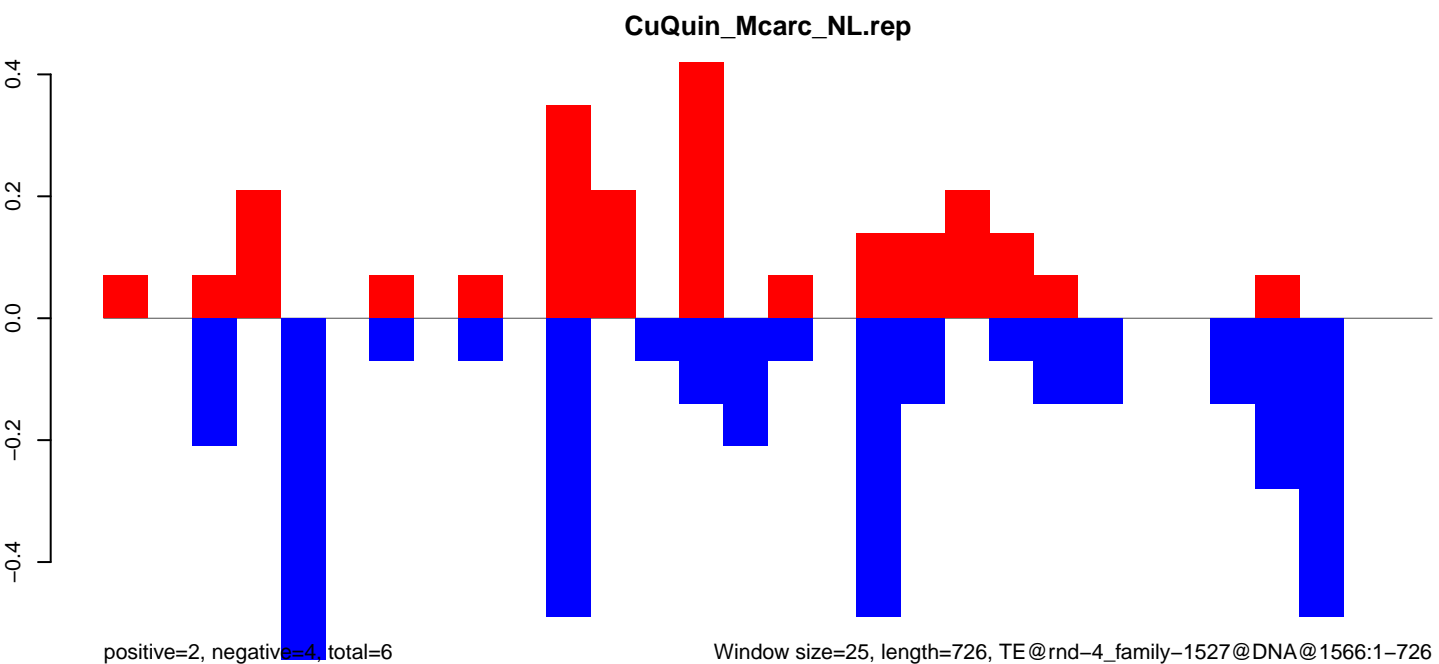
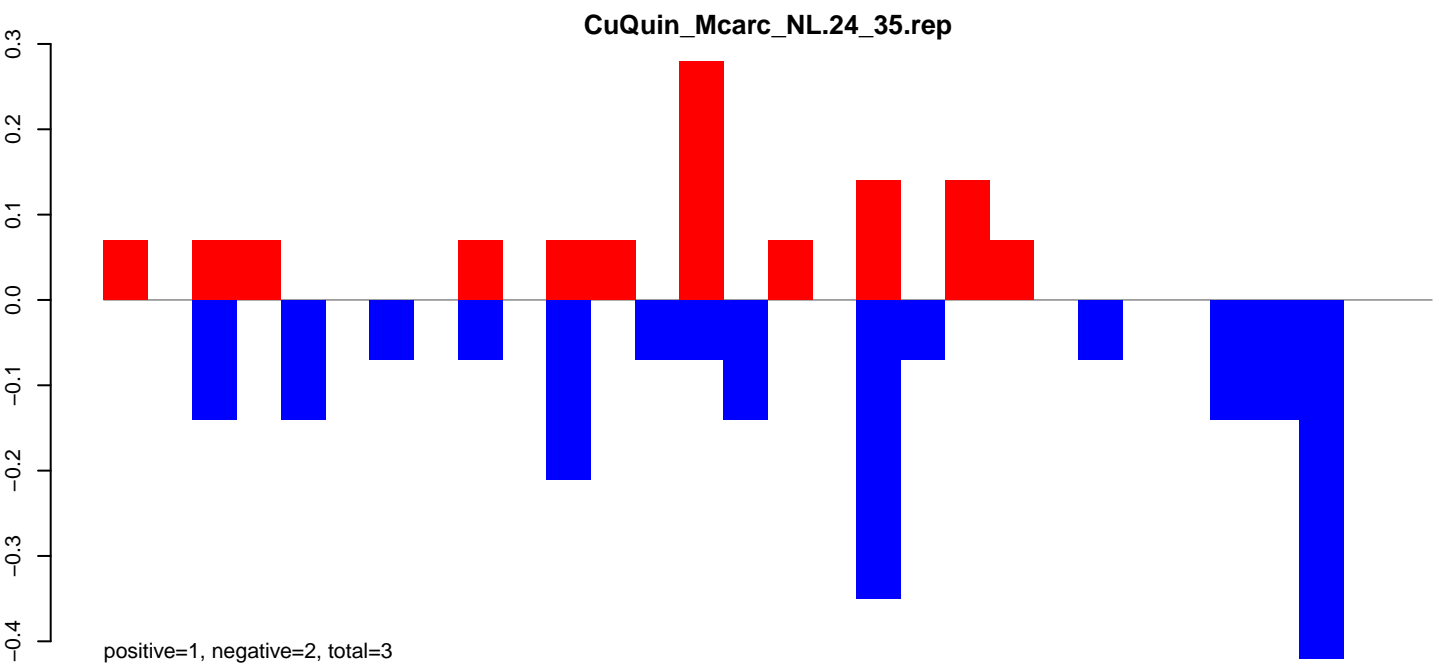
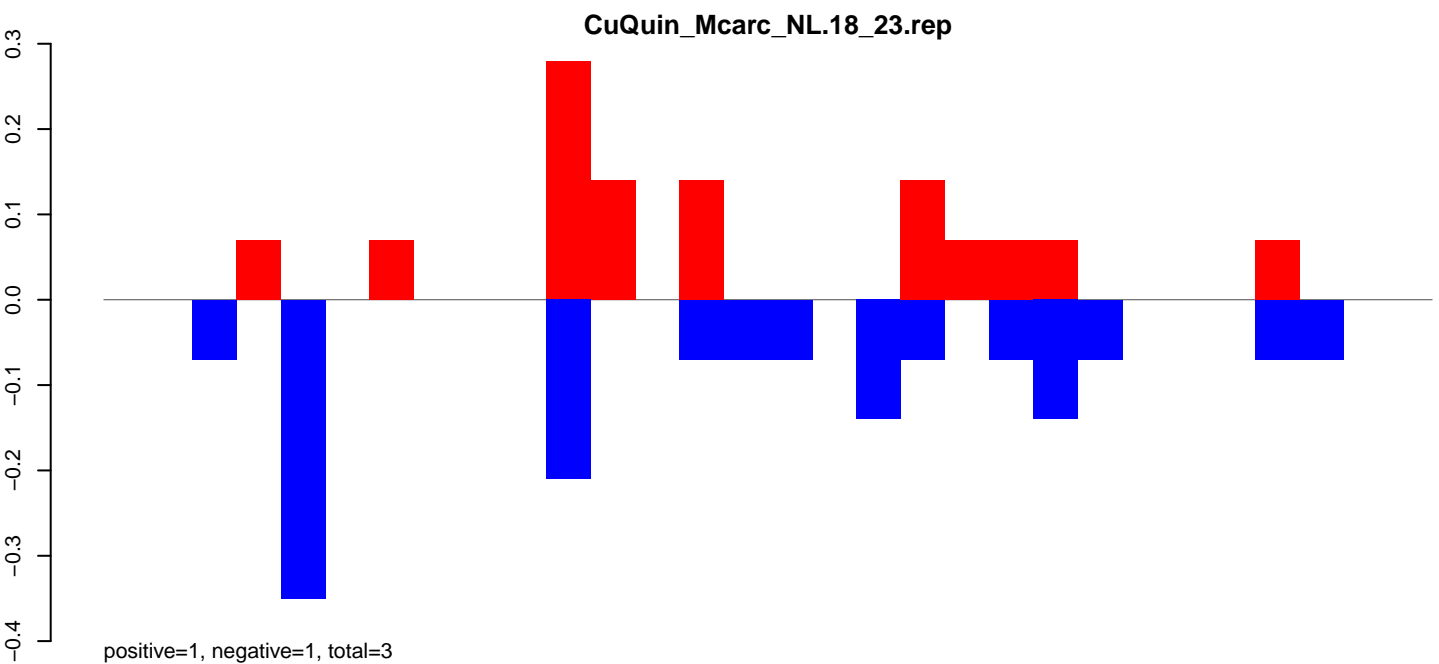


CuQuin_Mcarc_NL.24_35.rep



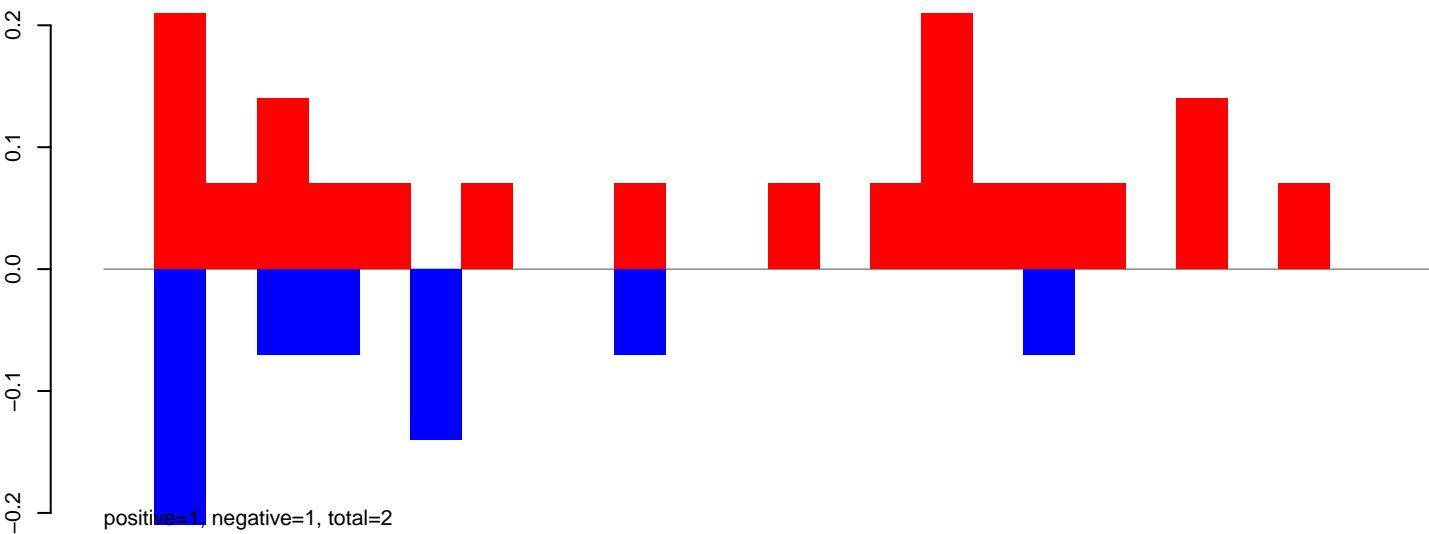
CuQuin_Mcarc_NL.rep



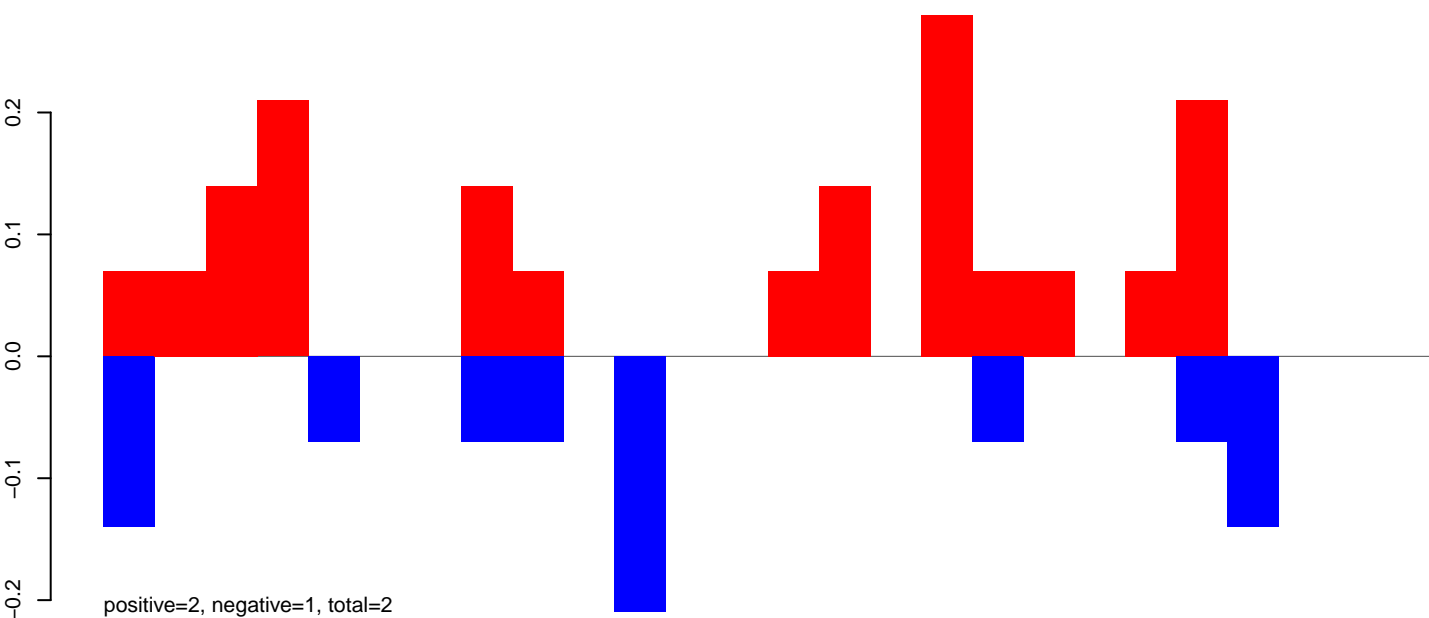


Window size=25, length=726, TE@rnd-4_family-1527@DNA@1566:1-726

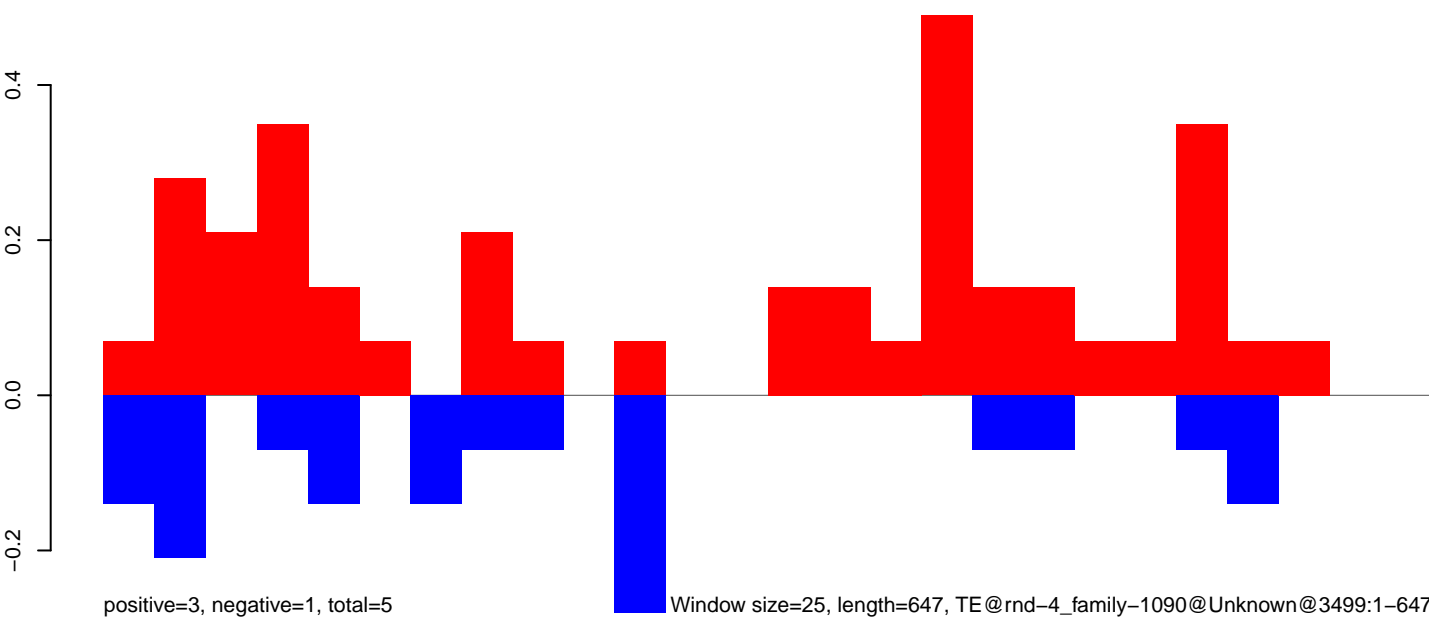
CuQuin_Mcarc_NL.18_23.rep



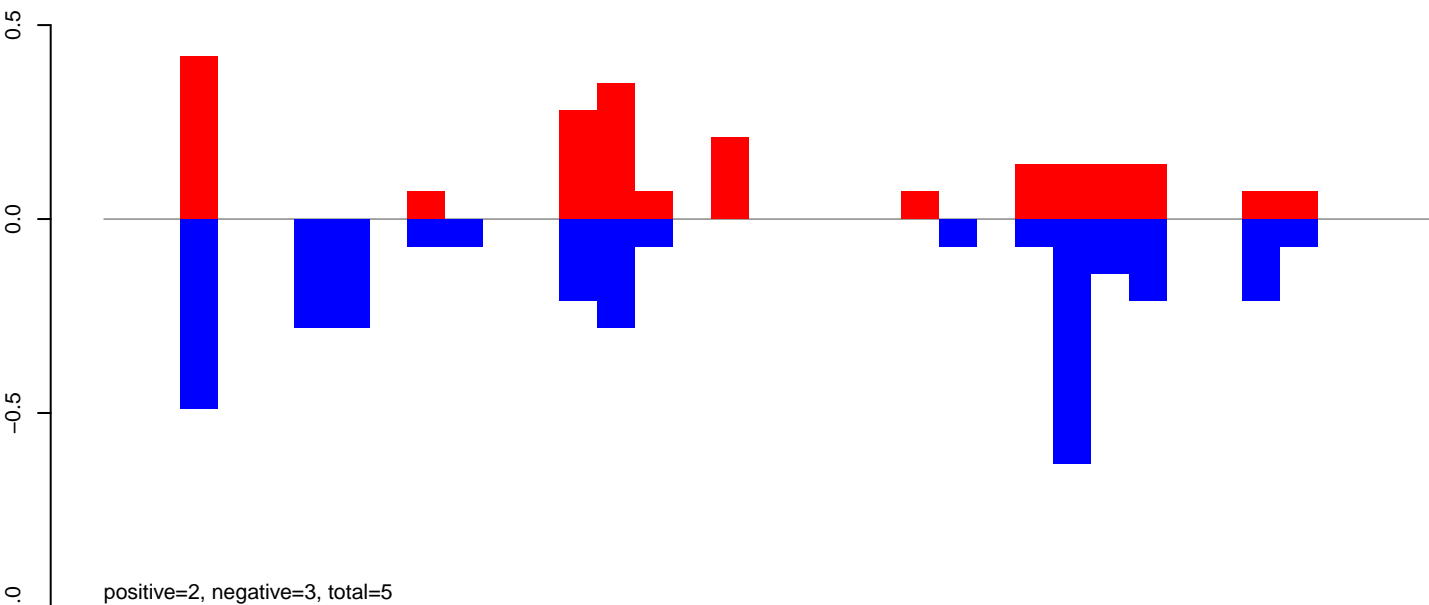
CuQuin_Mcarc_NL.24_35.rep



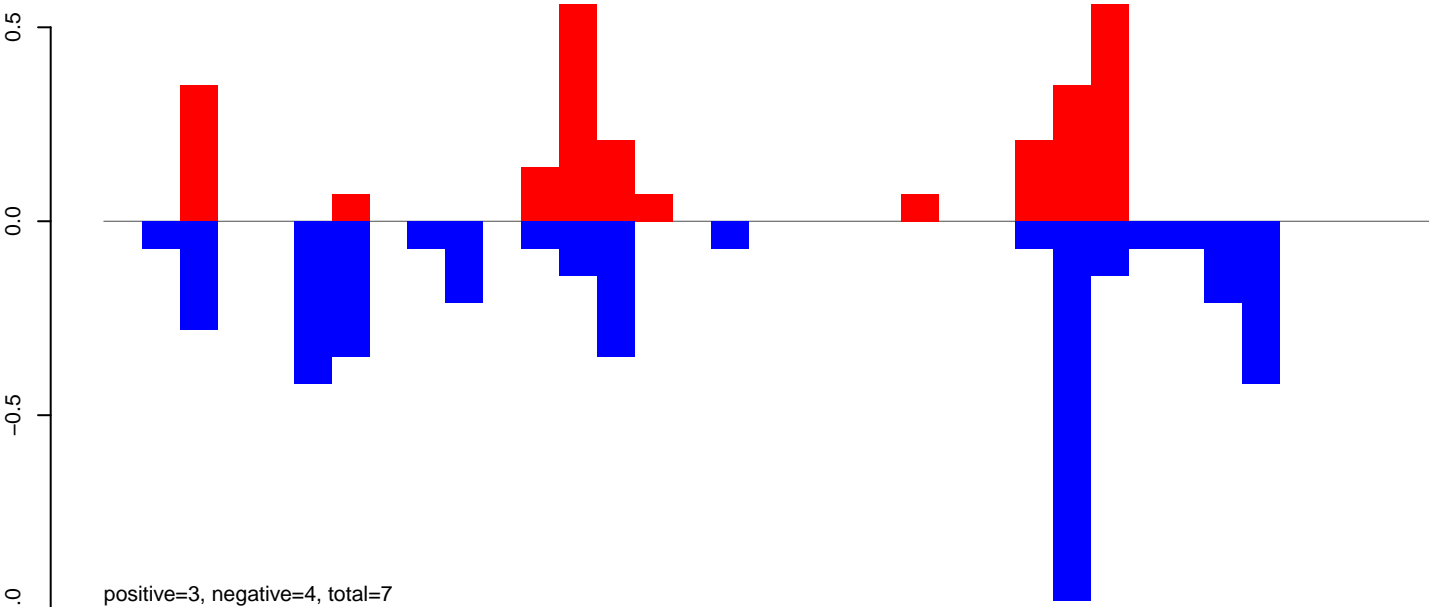
CuQuin_Mcarc_NL.rep



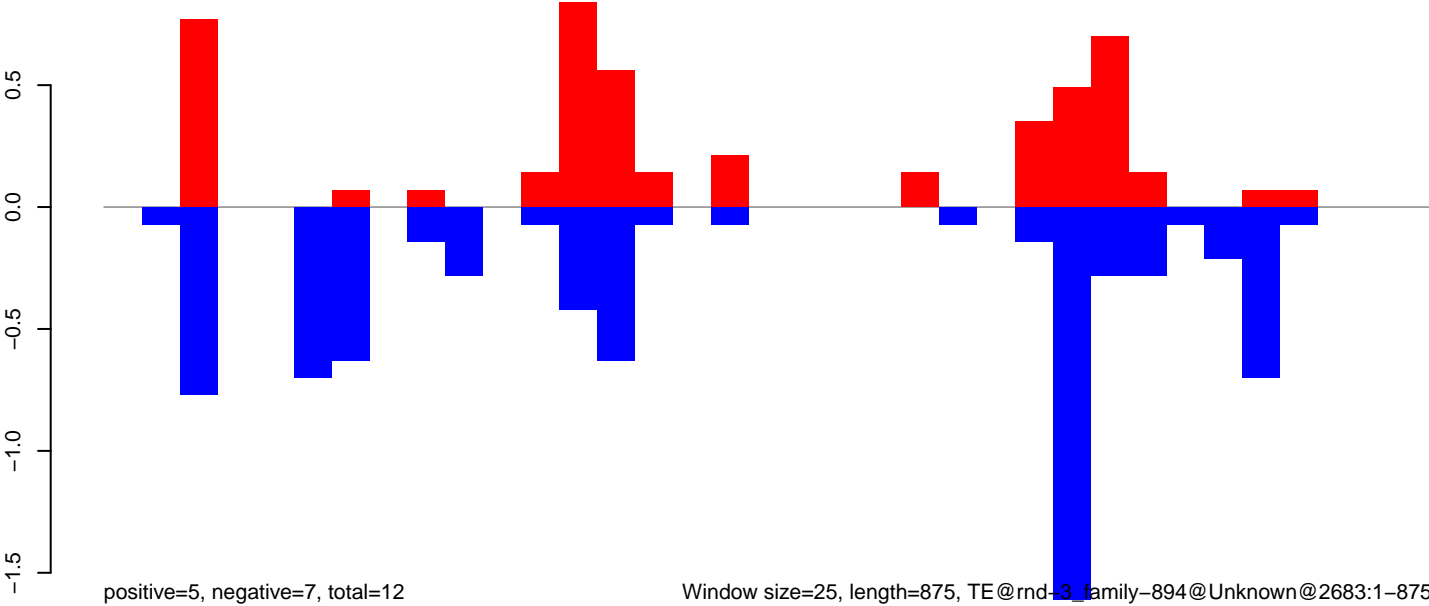
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

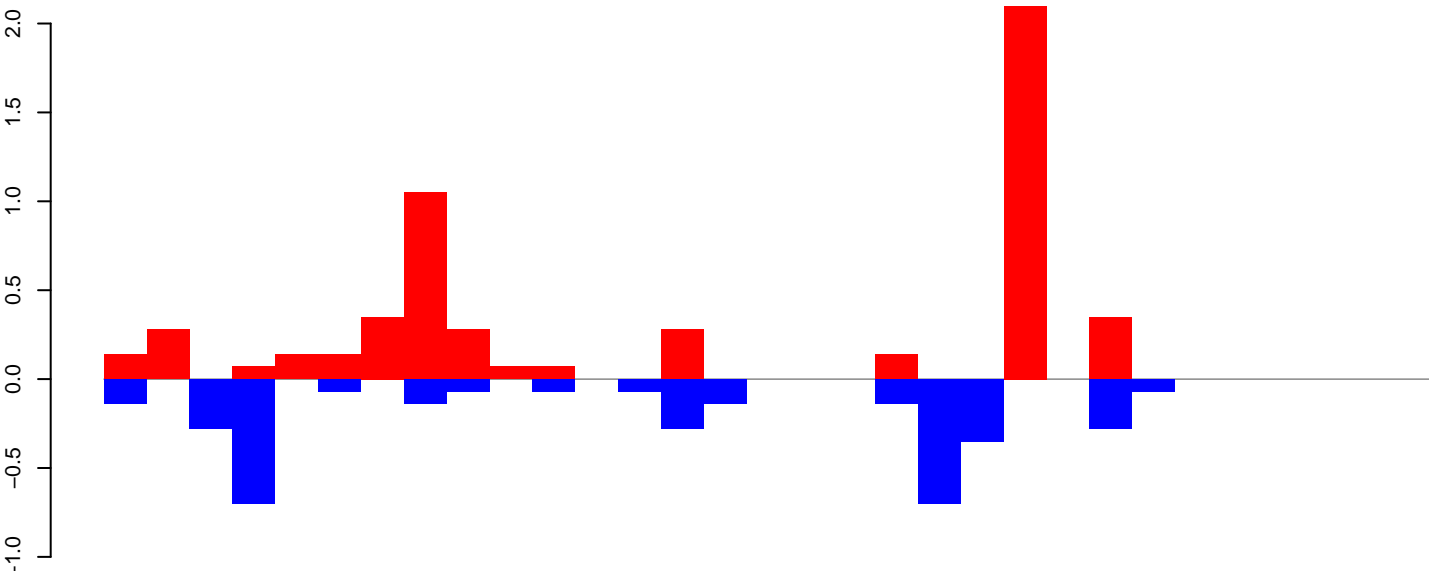


CuQuin_Mcarc_NL.rep



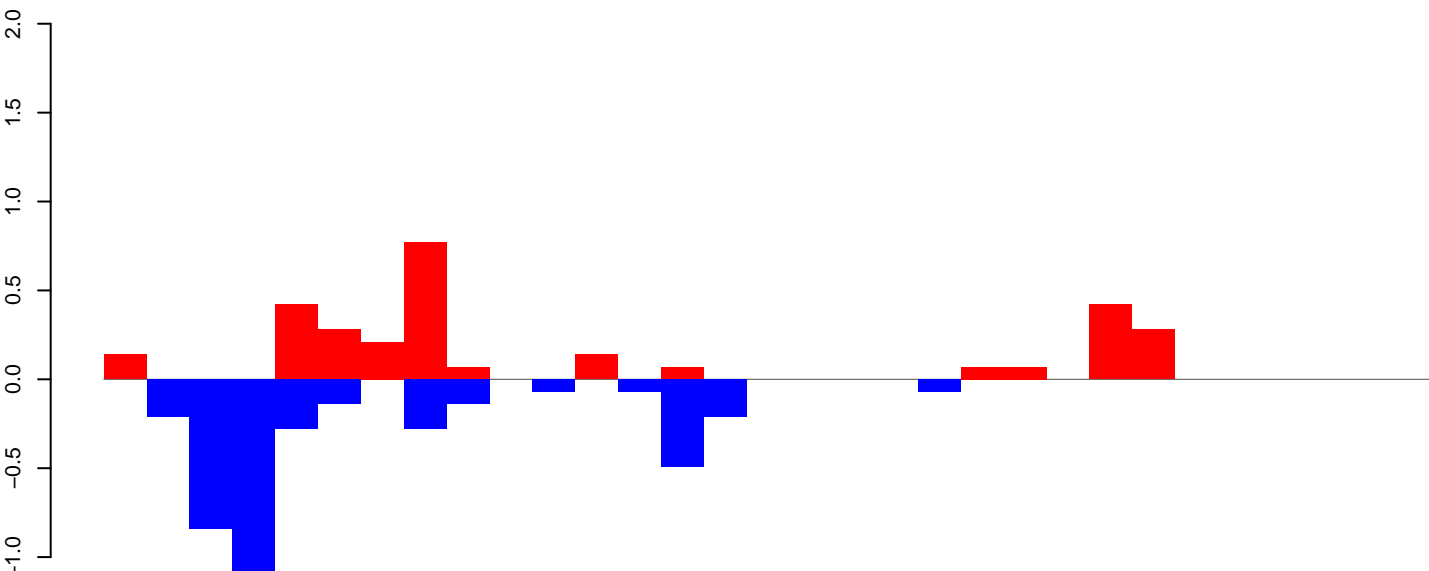
Window size=25, length=875, TE@rnd-3, family-894@Unknown@2683:1-875

CuQuin_Mcarc_NL.18_23.rep



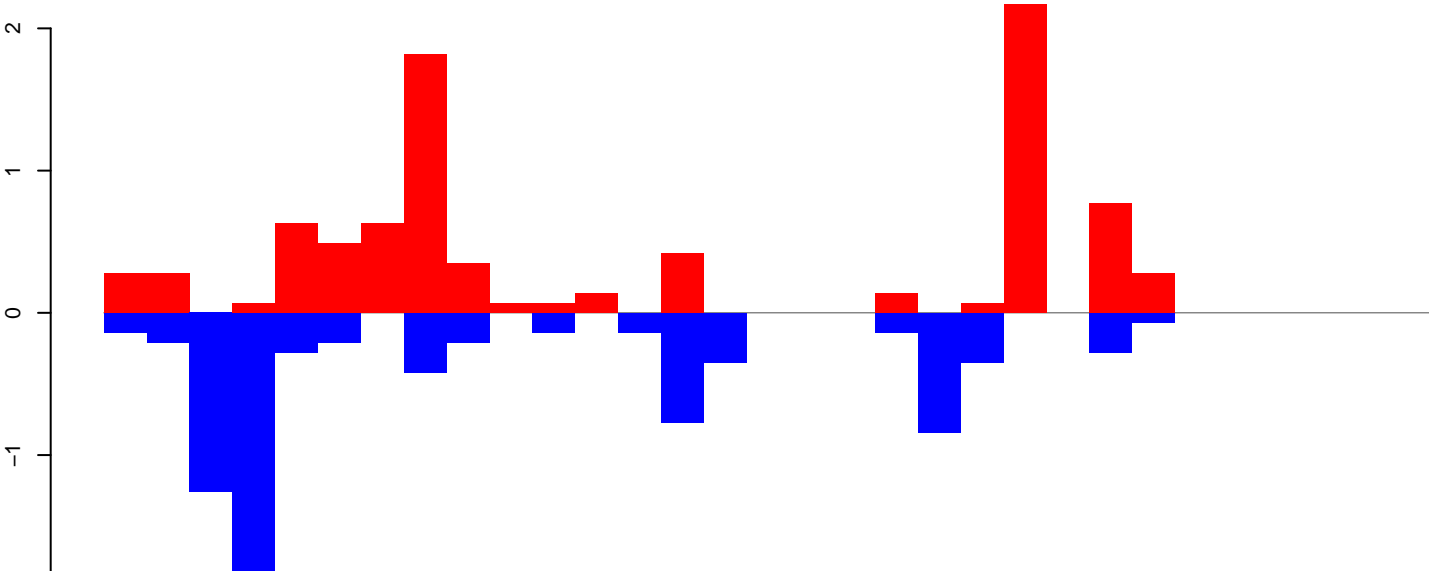
positive=5, negative=3, total=9

CuQuin_Mcarc_NL.24_35.rep



positive=3, negative=4, total=7

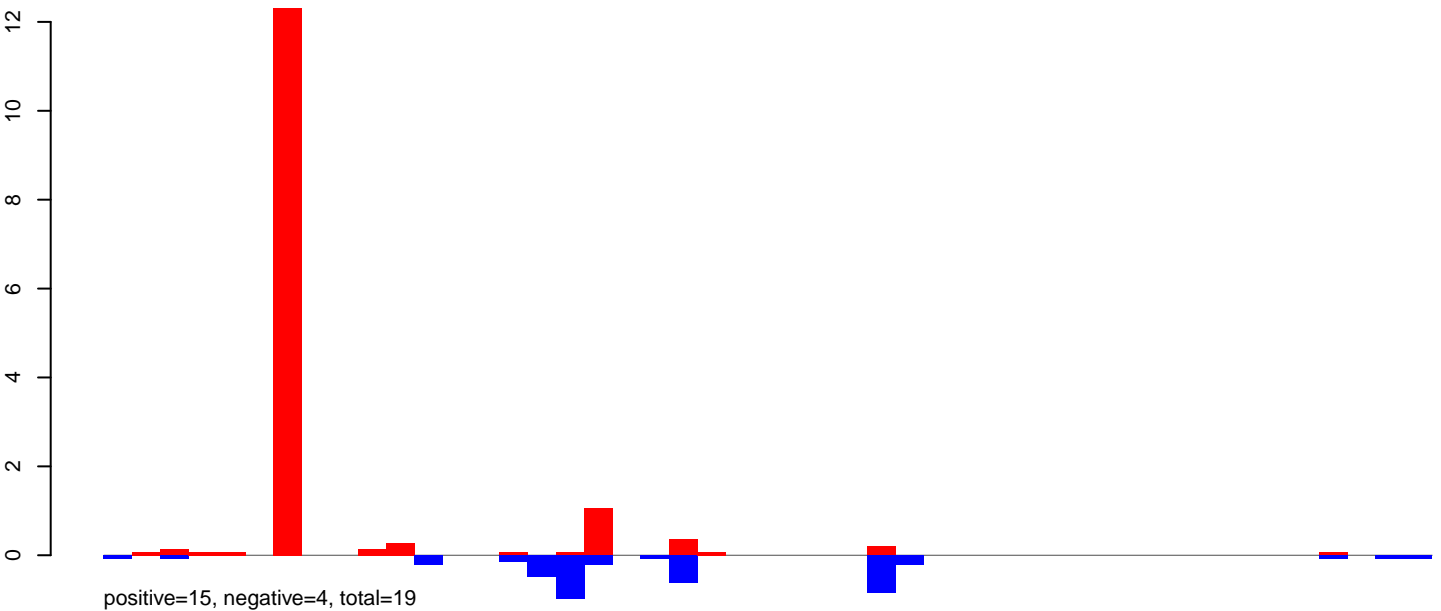
CuQuin_Mcarc_NL.rep



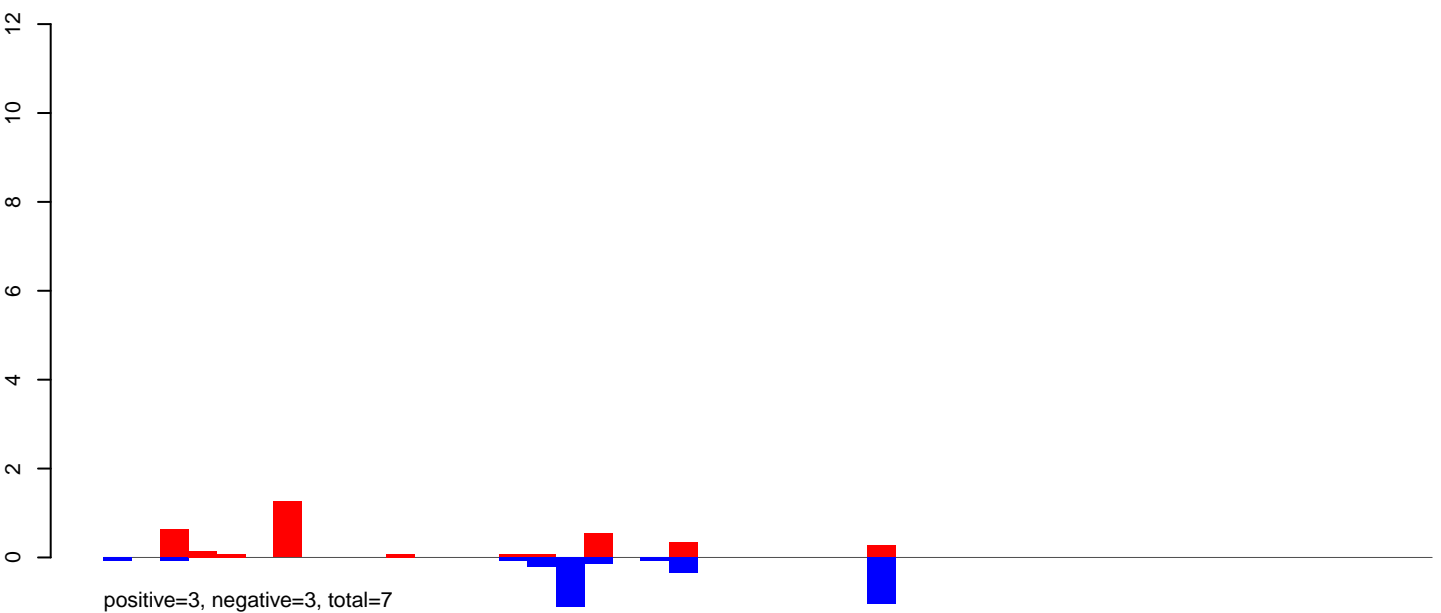
positive=9, negative=8, total=17

Window size=25, length=752, TE@rnd-3_family-75@DNA@3323:1-752

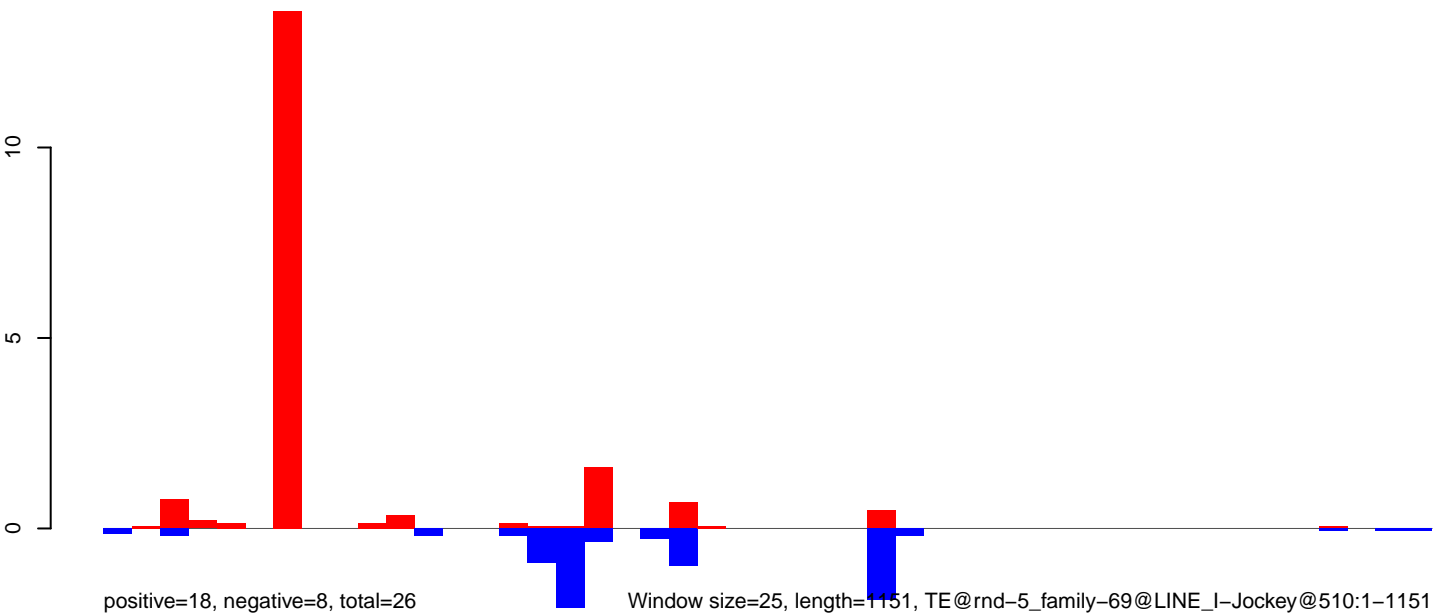
CuQuin_Mcarc_NL.18_23.rep



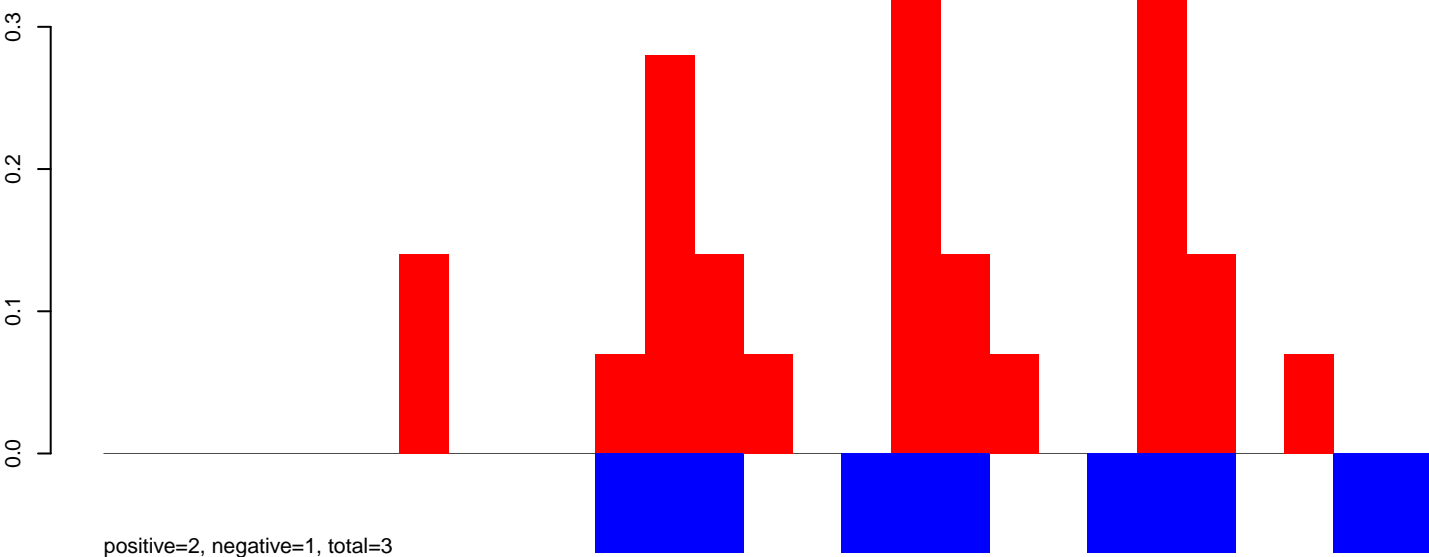
CuQuin_Mcarc_NL.24_35.rep



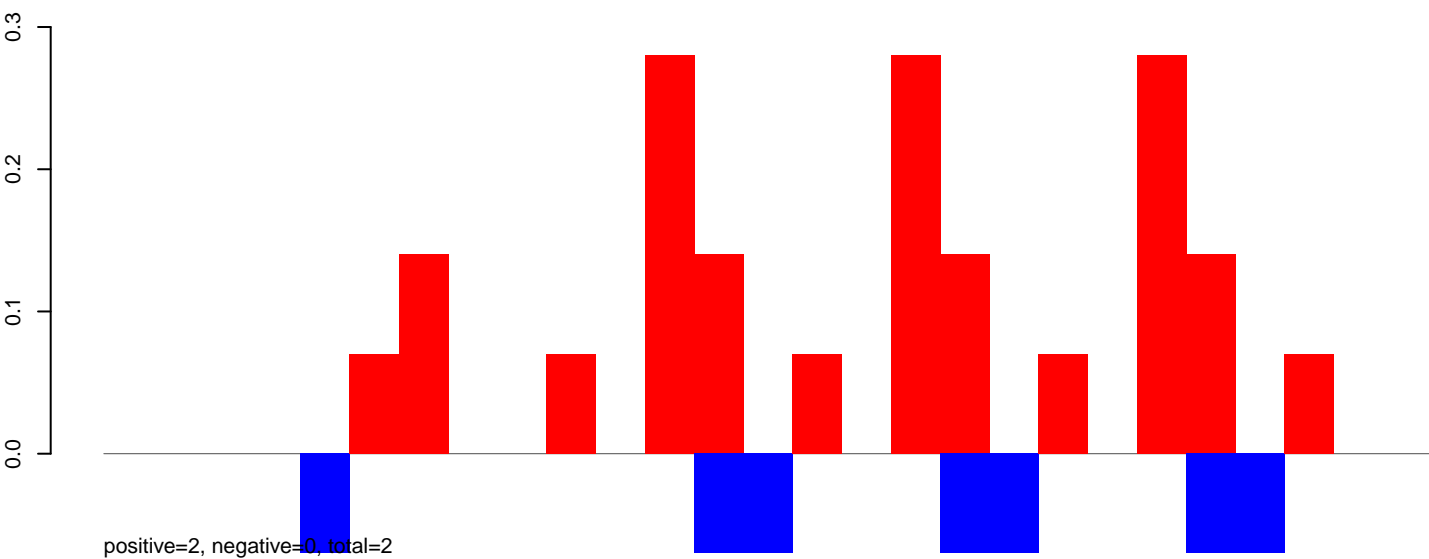
CuQuin_Mcarc_NL.rep



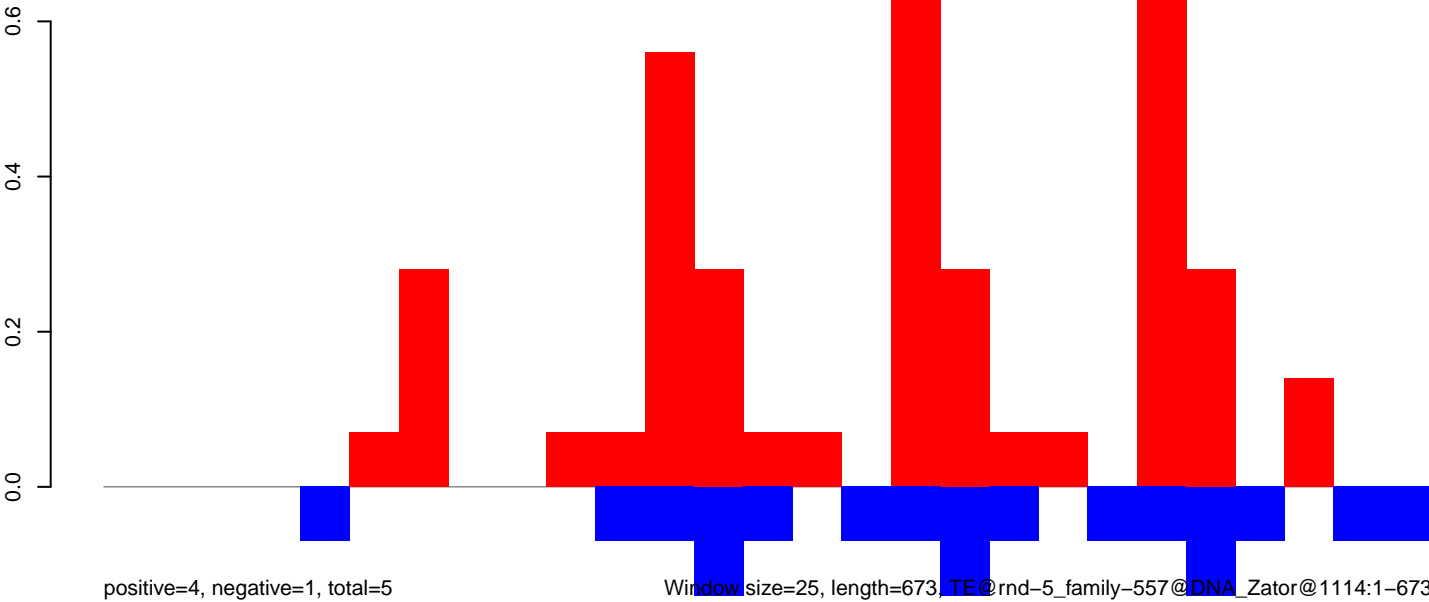
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

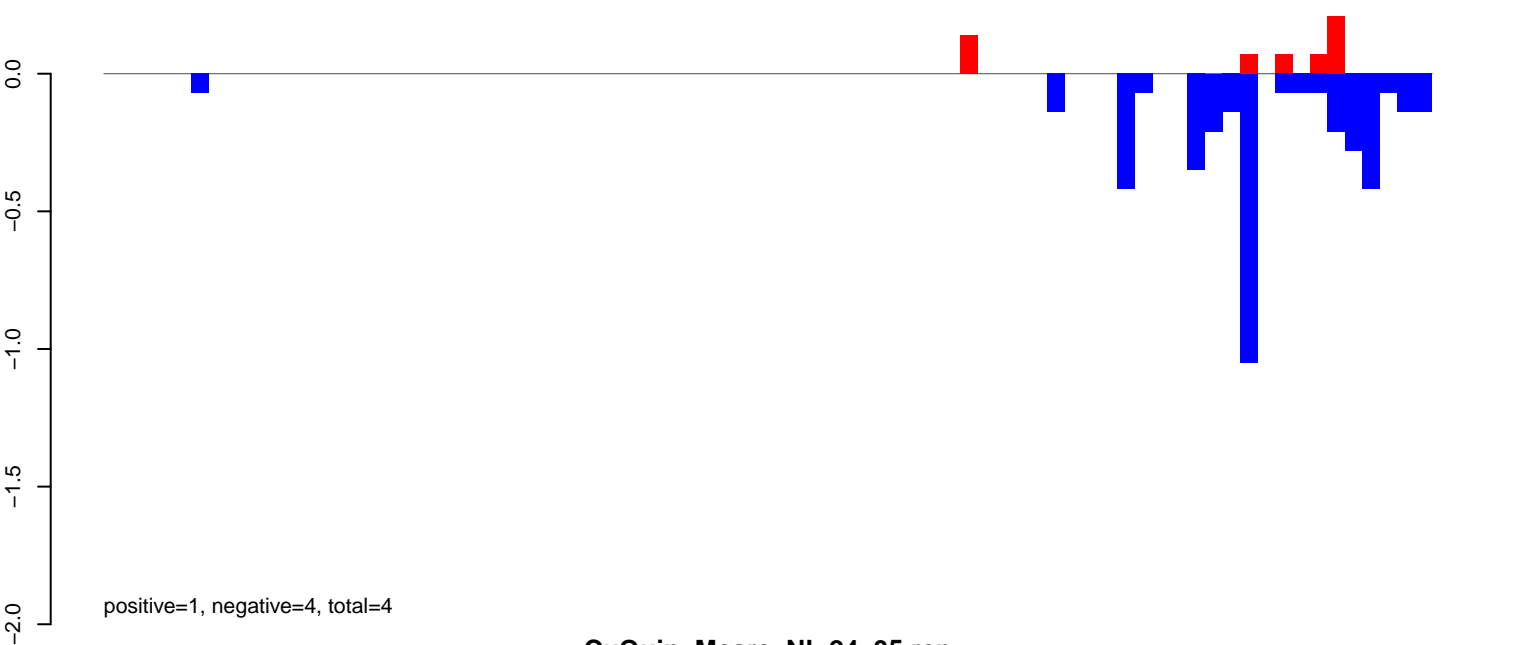


CuQuin_Mcarc_NL.rep



Window size=25, length=673, TE@rnd-5_family-557@DNA_Zator@1114:1-673

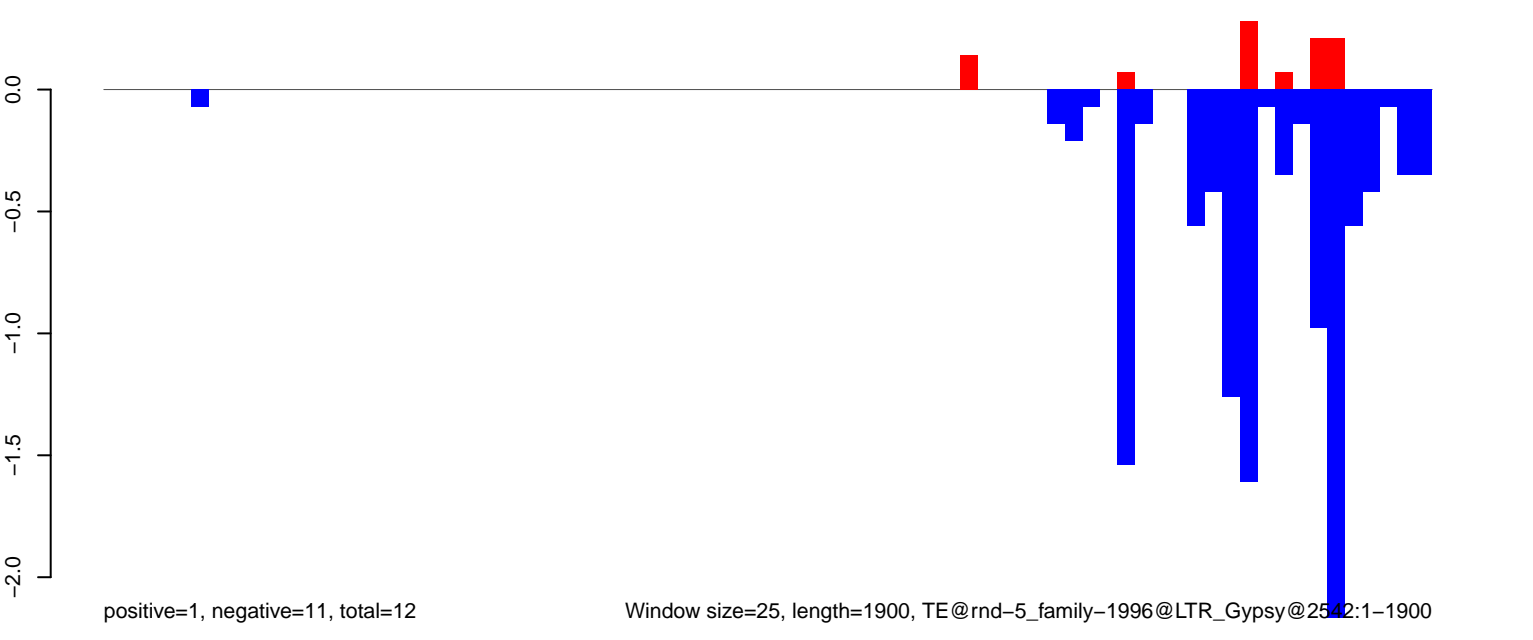
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

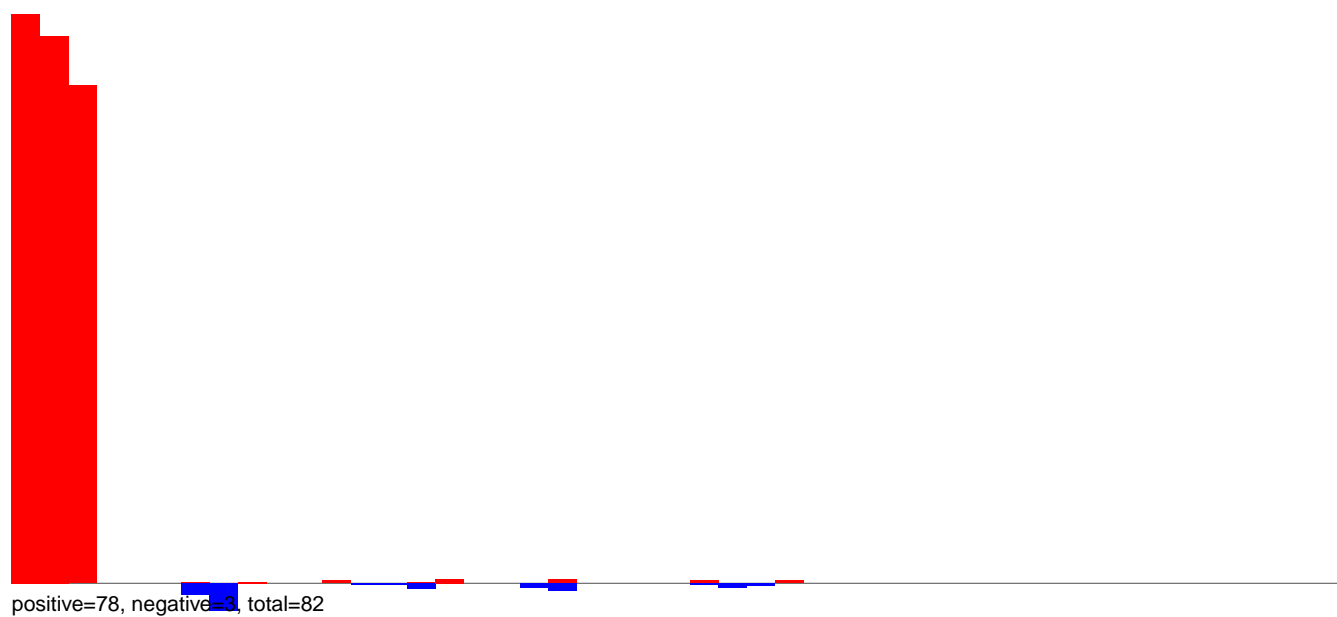


CuQuin_Mcarc_NL.rep

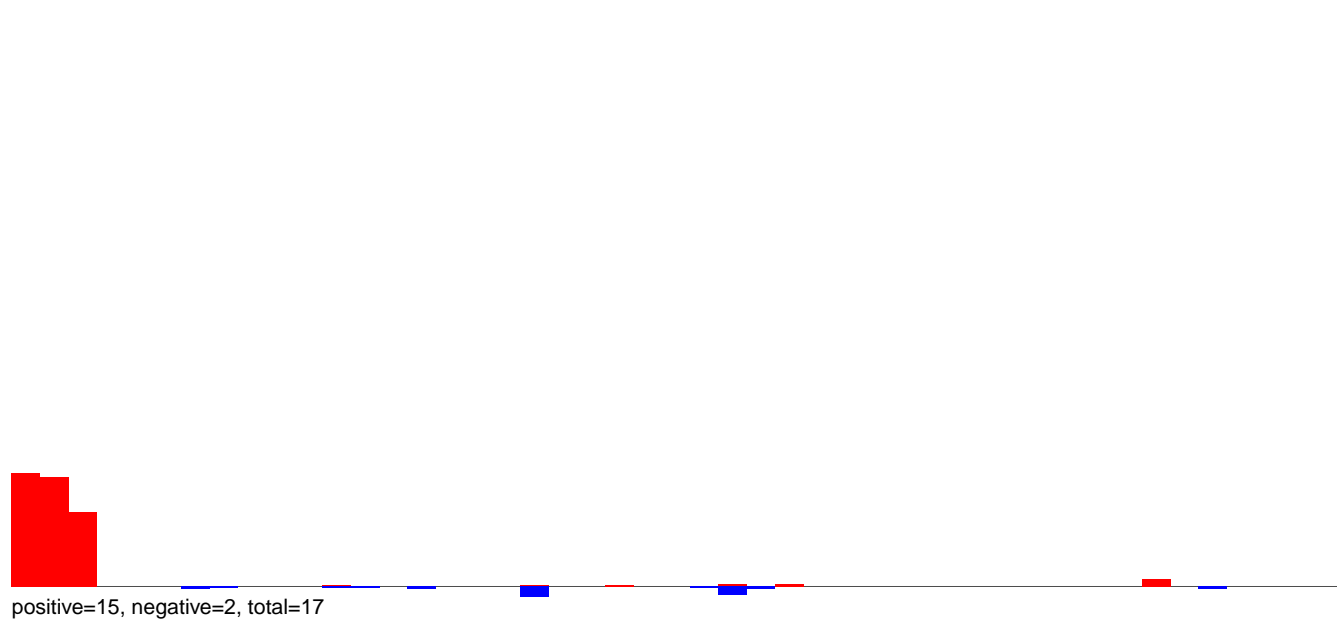


Window size=25, length=1900, TE@rnd-5_family-1996@LTR_Gypsy@2542:1-1900

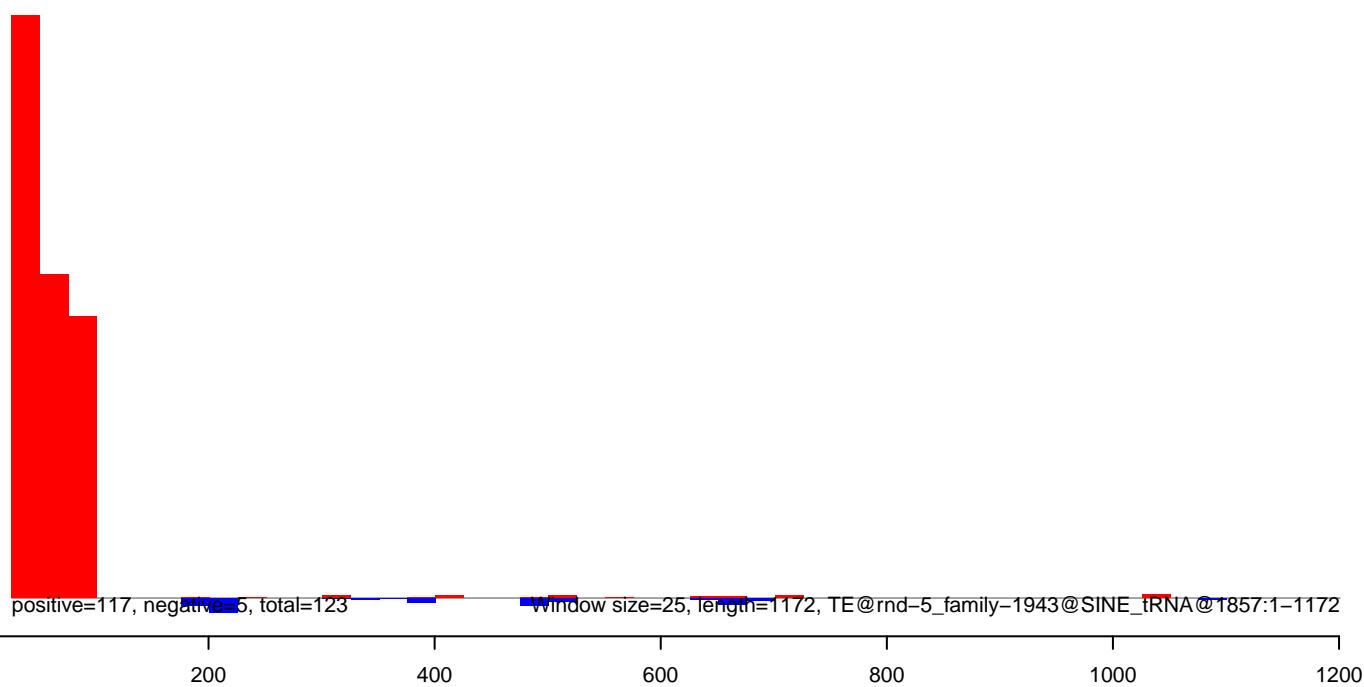
CuQuin_Mcarc_NL.18_23.rep



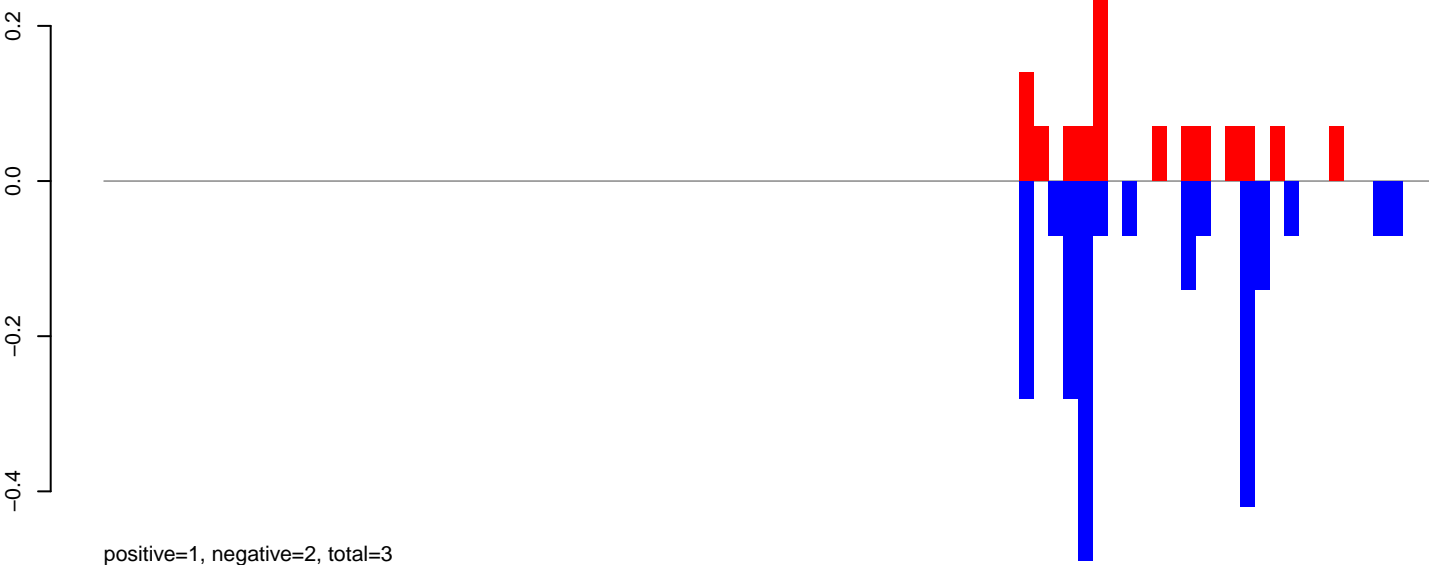
CuQuin_Mcarc_NL.24_35.rep



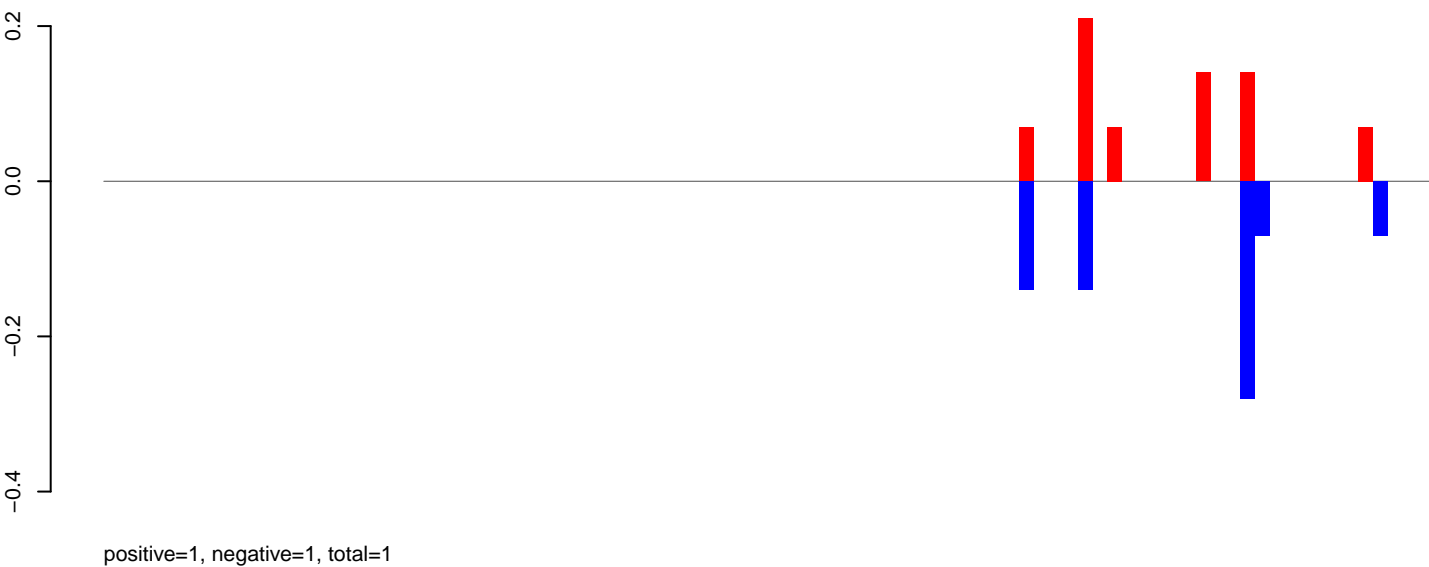
CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep



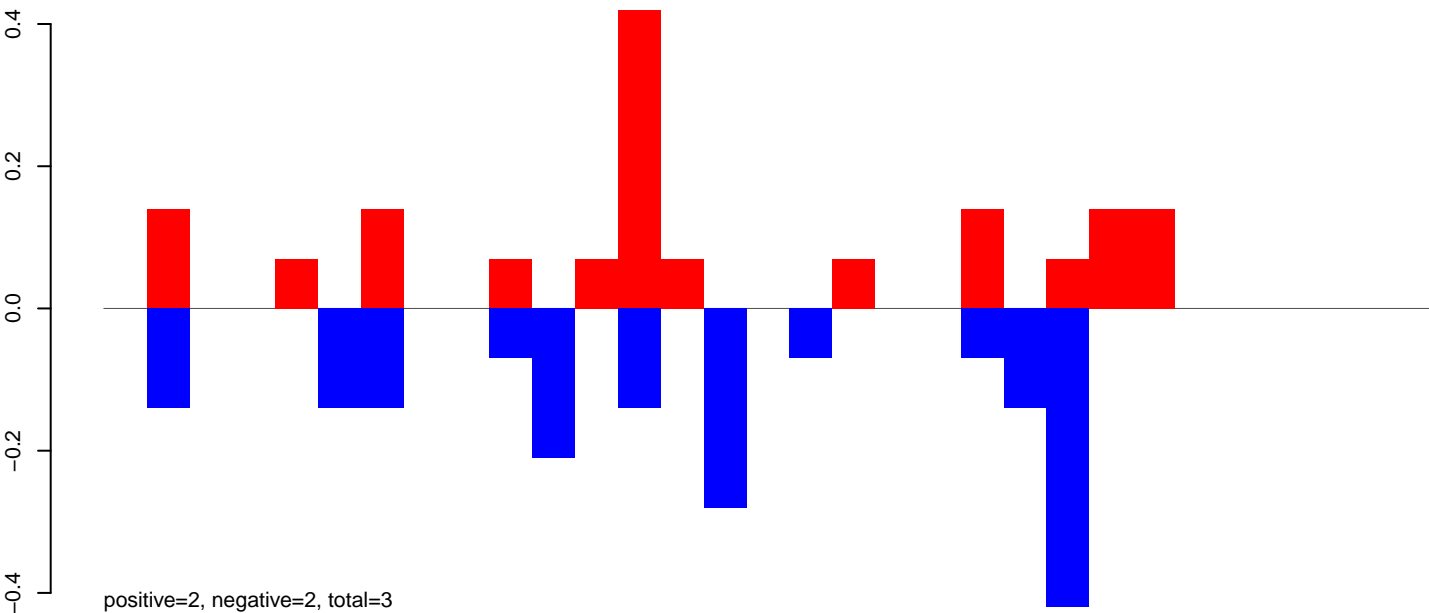
CuQuin_Mcarc_NL.24_35.rep



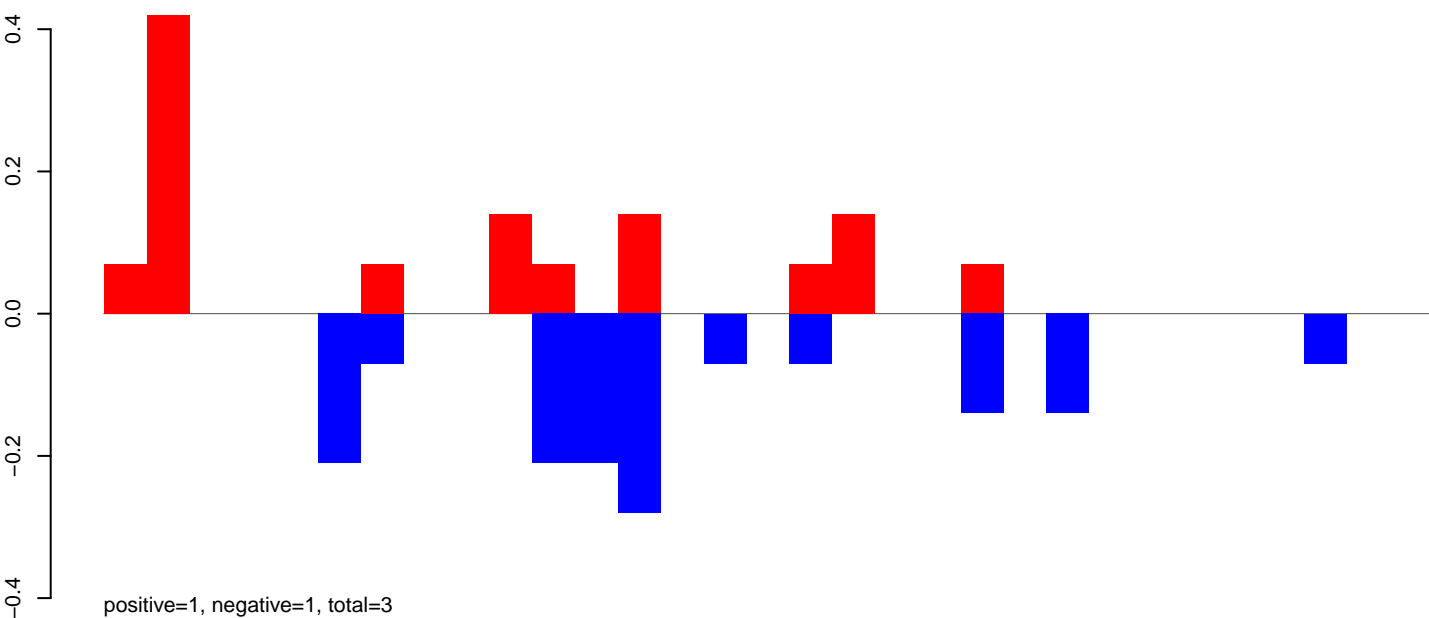
CuQuin_Mcarc_NL.rep



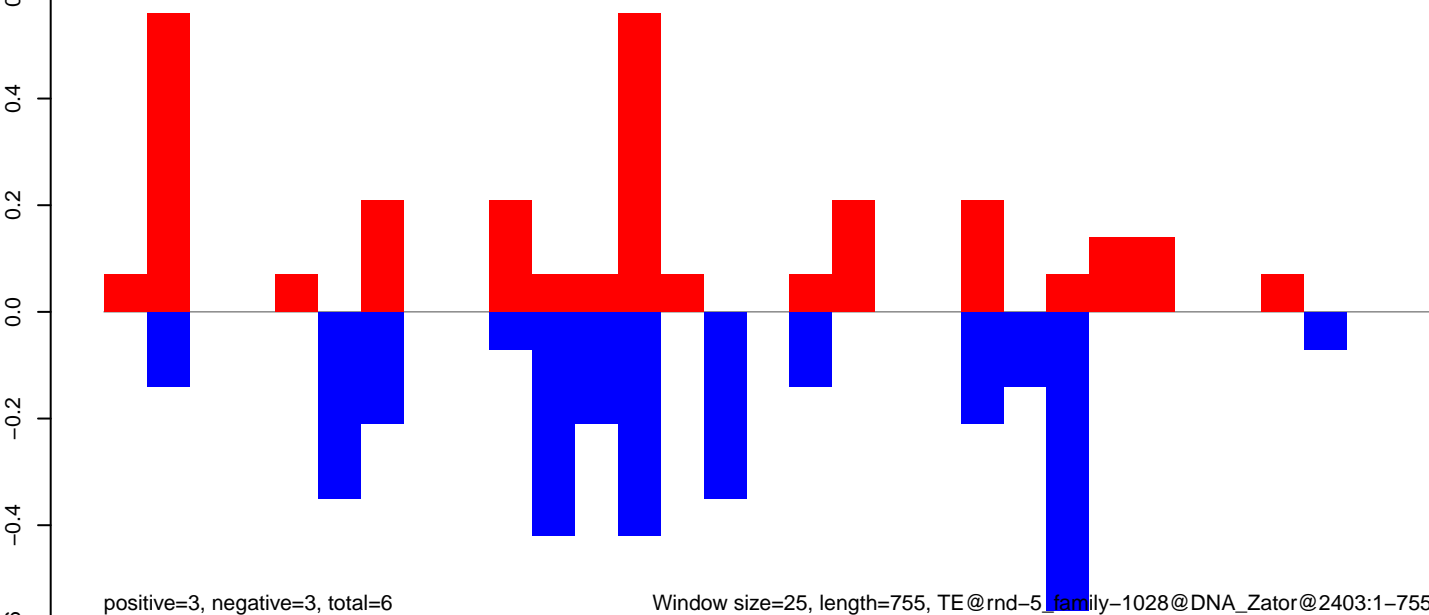
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

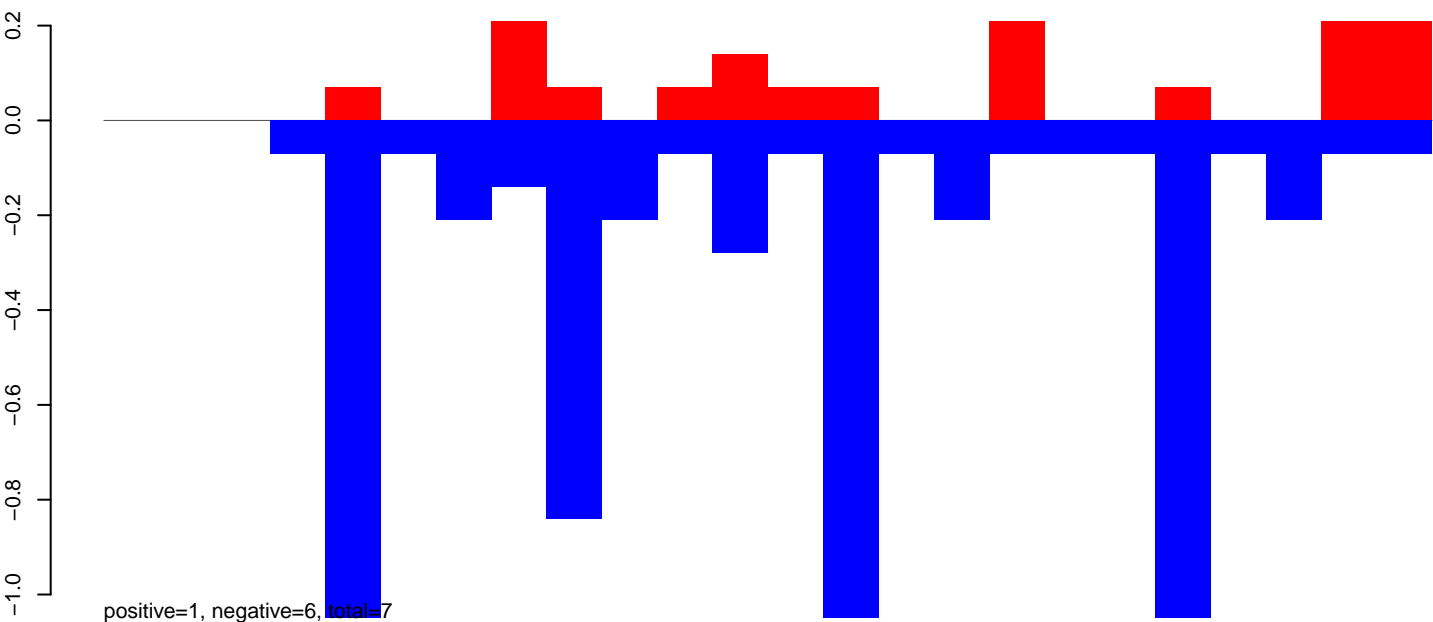


CuQuin_Mcarc_NL.rep

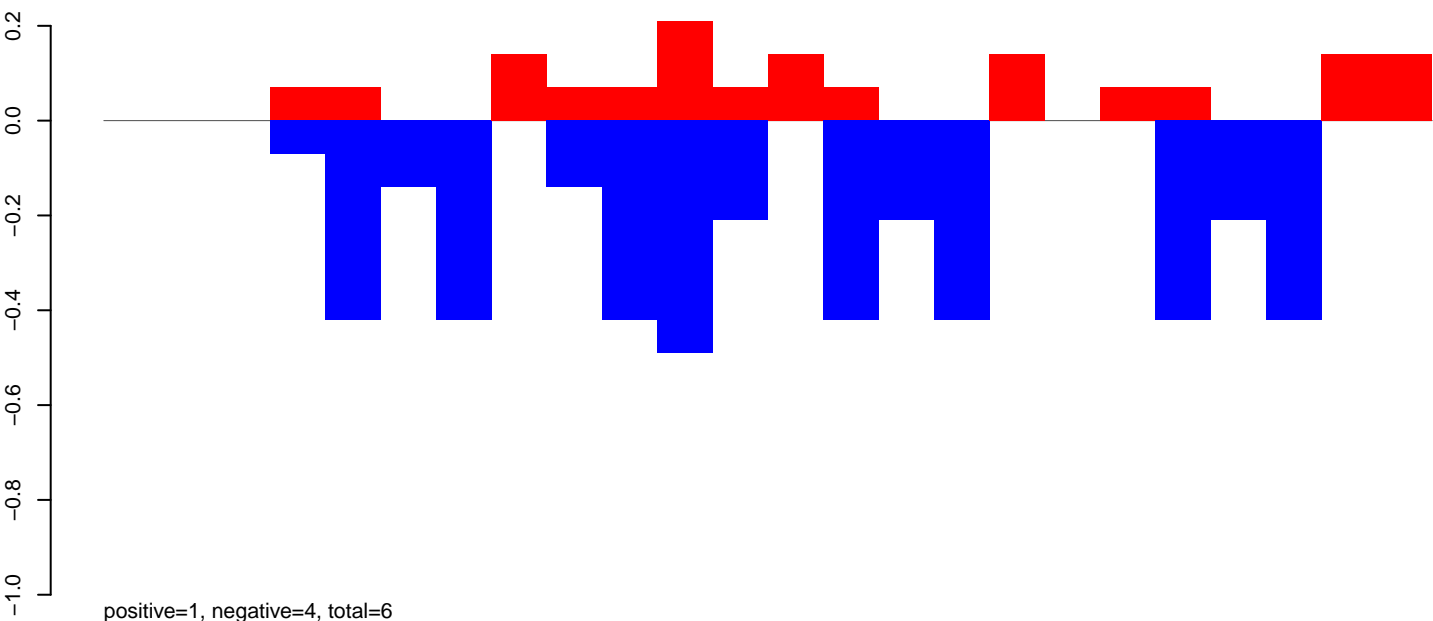


Window size=25, length=755, TE@rnd-5_family-1028@DNA_Zator@2403:1-755

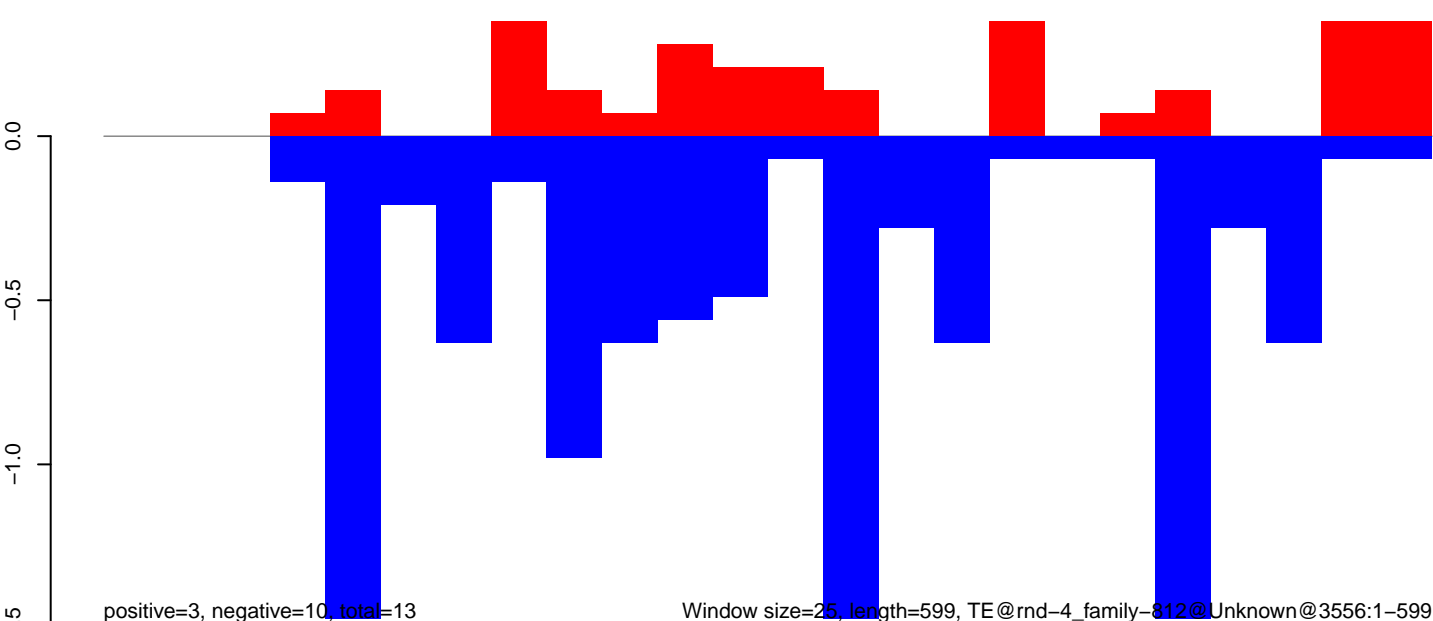
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

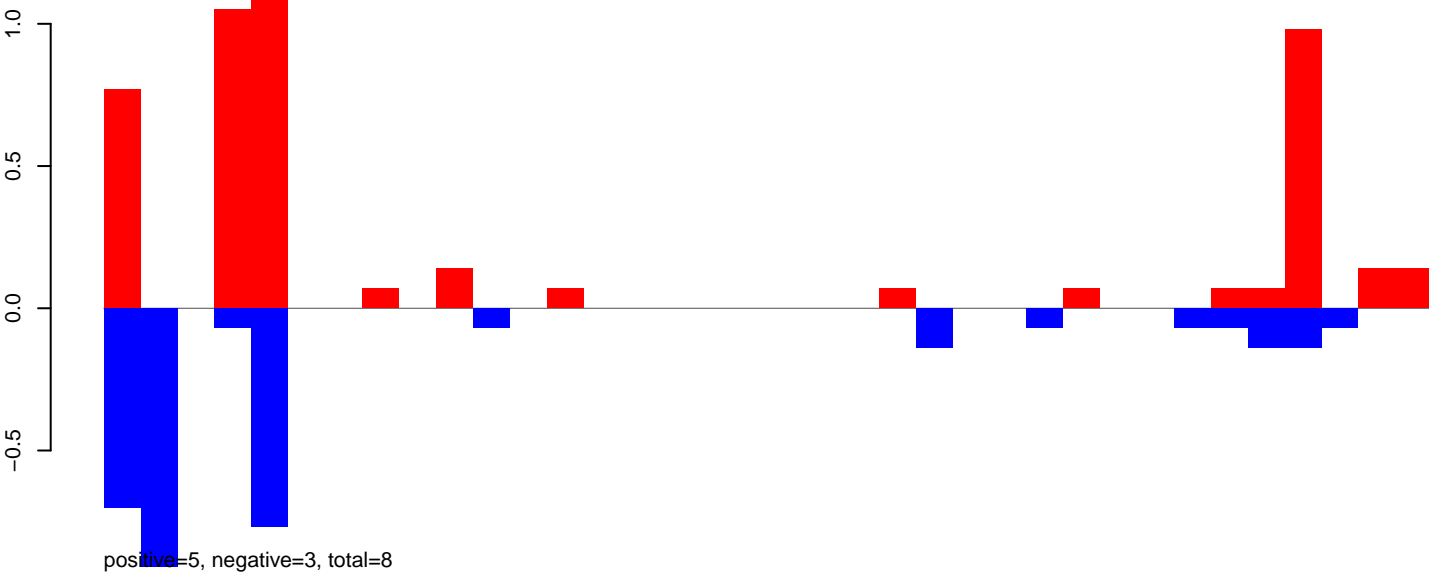


CuQuin_Mcarc_NL.rep

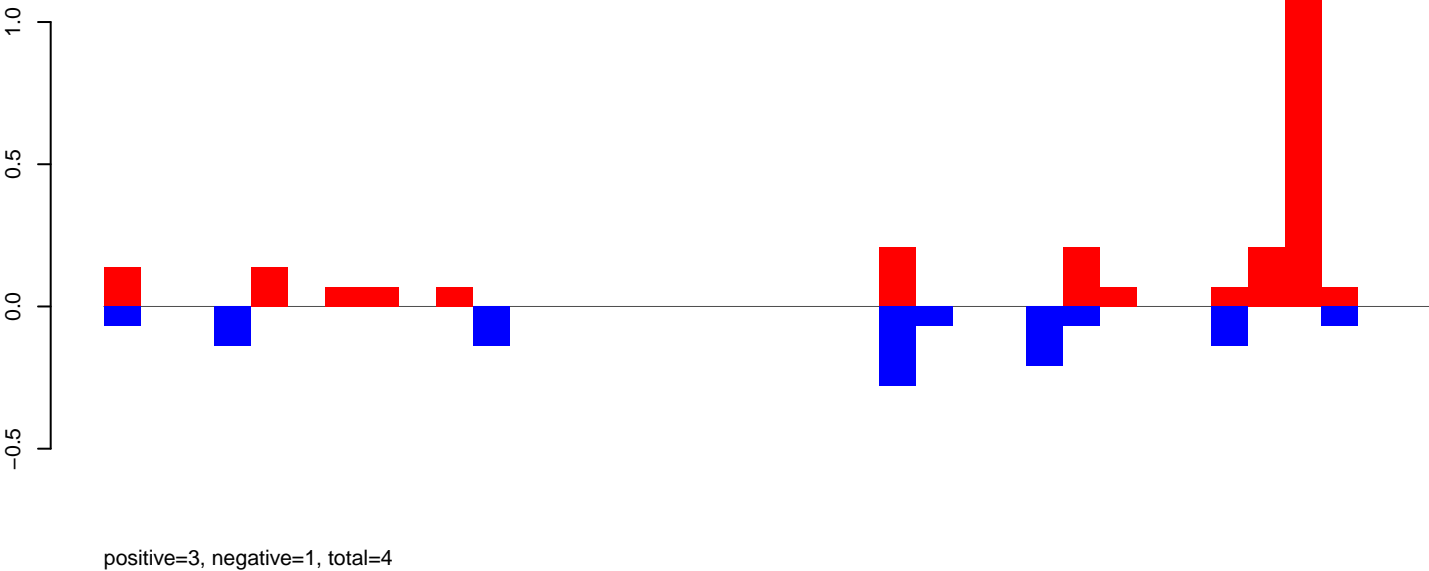


Window size=25, length=599, TE@rnd-4_family-312@Unknown@3556:1-599

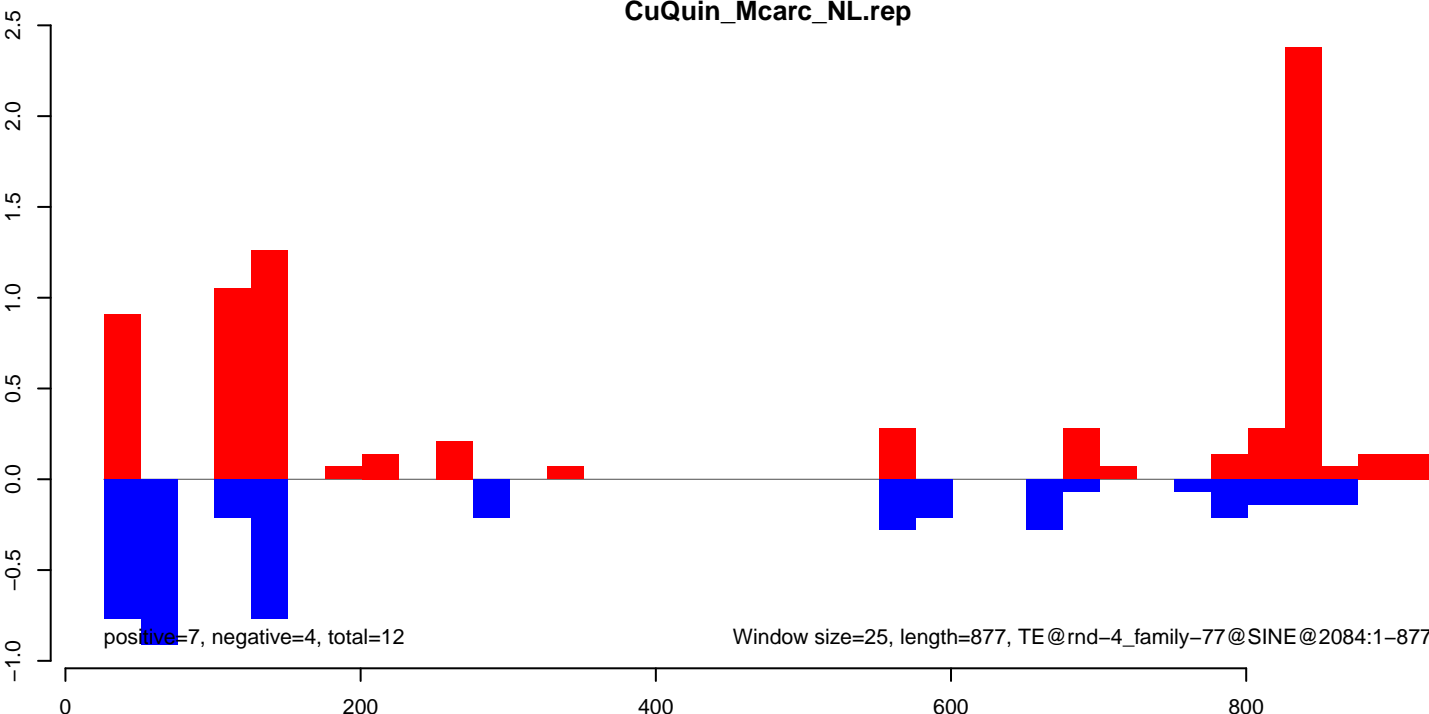
CuQuin_Mcarc_NL.18_23.rep



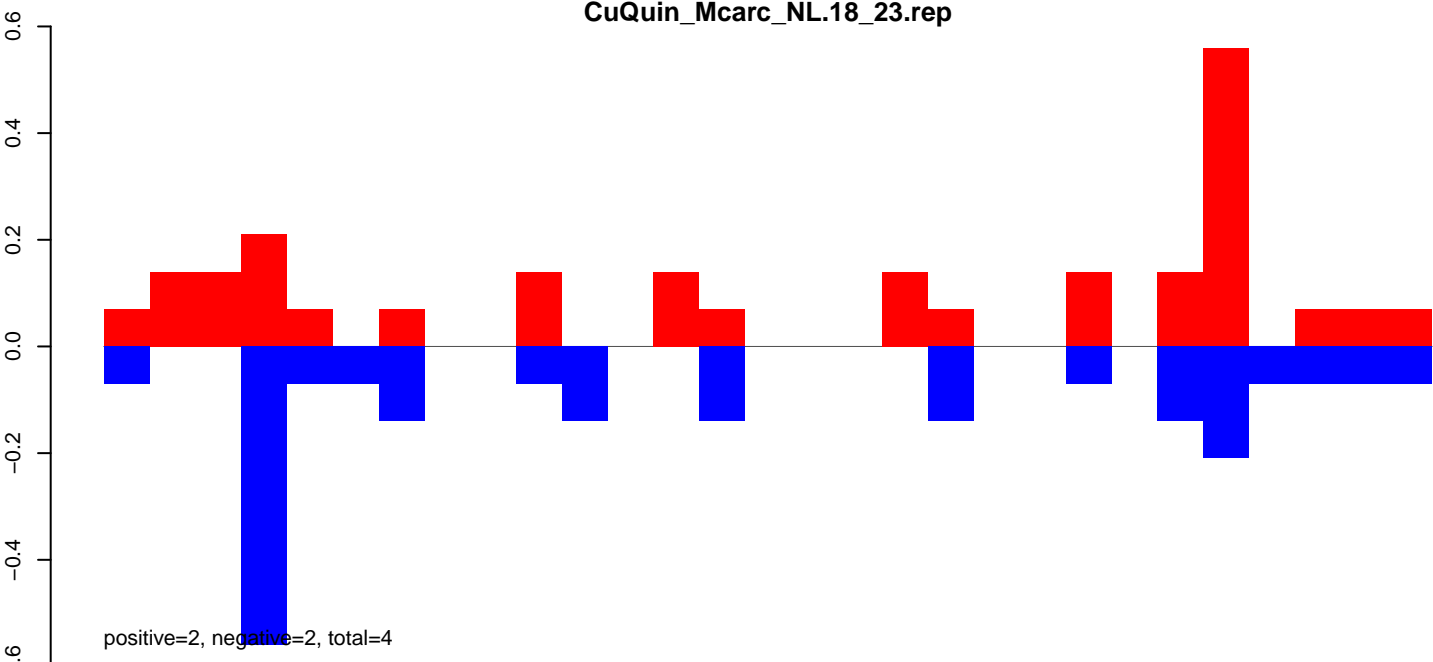
CuQuin_Mcarc_NL.24_35.rep



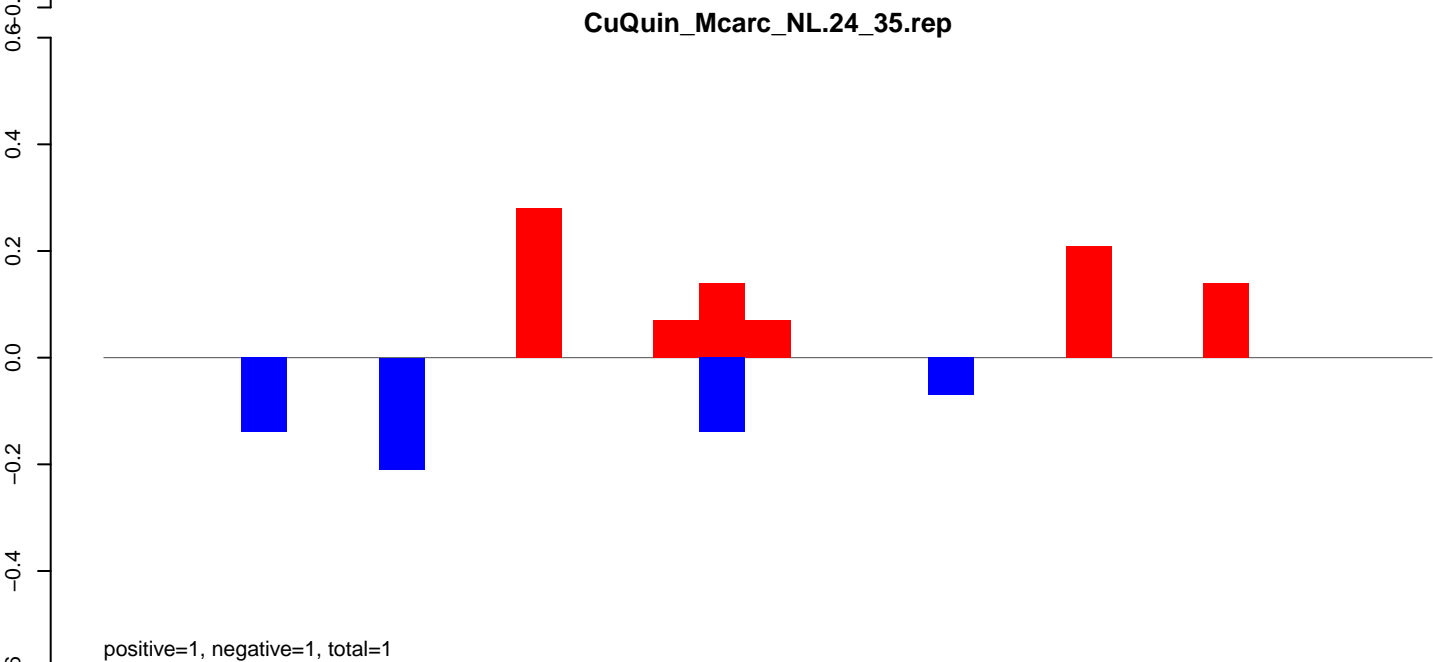
CuQuin_Mcarc_NL.rep



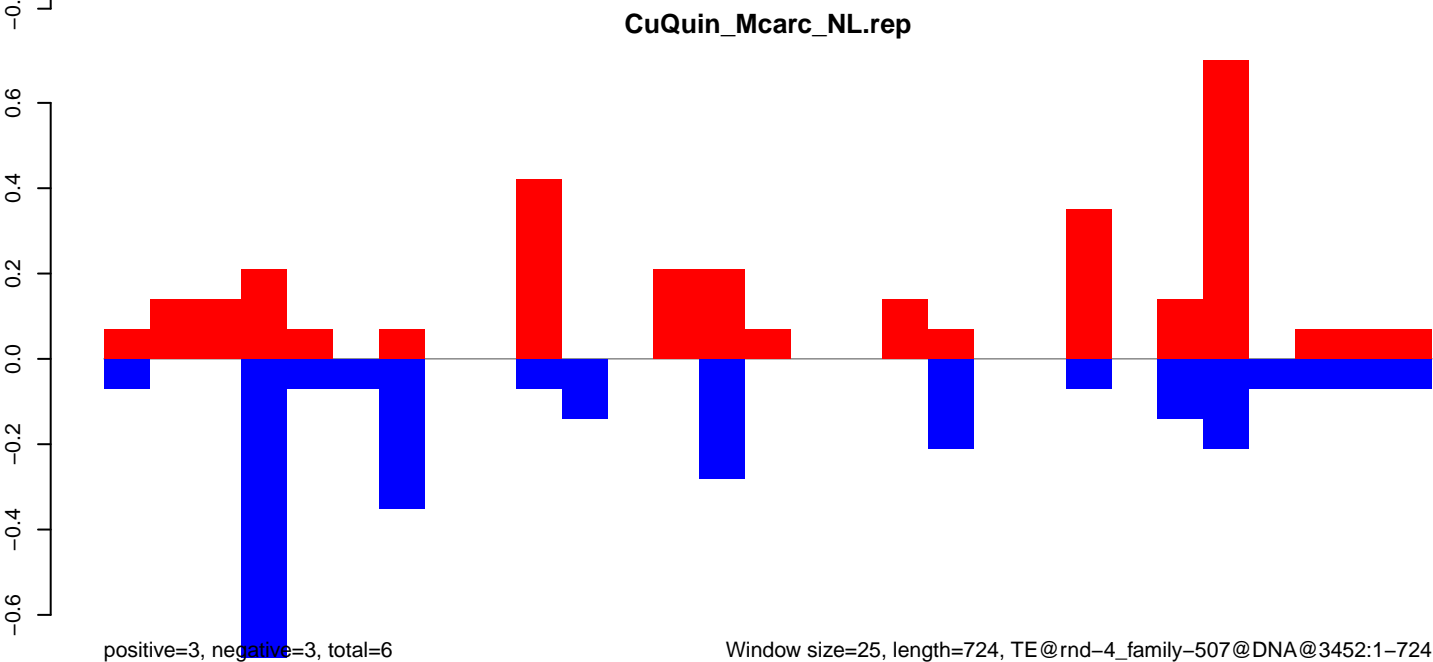
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

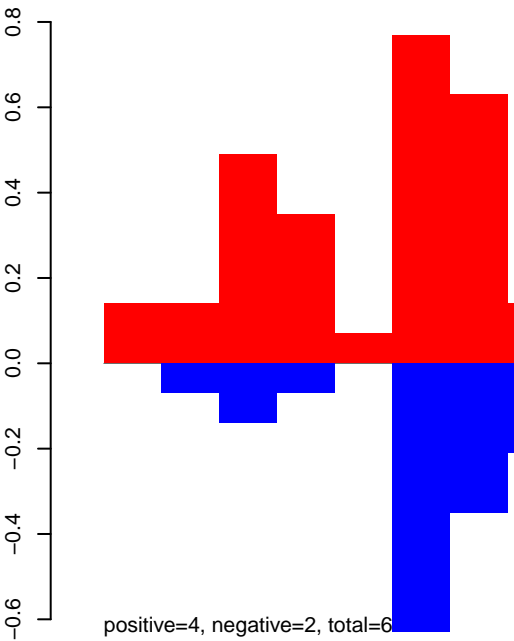


CuQuin_Mcarc_NL.rep

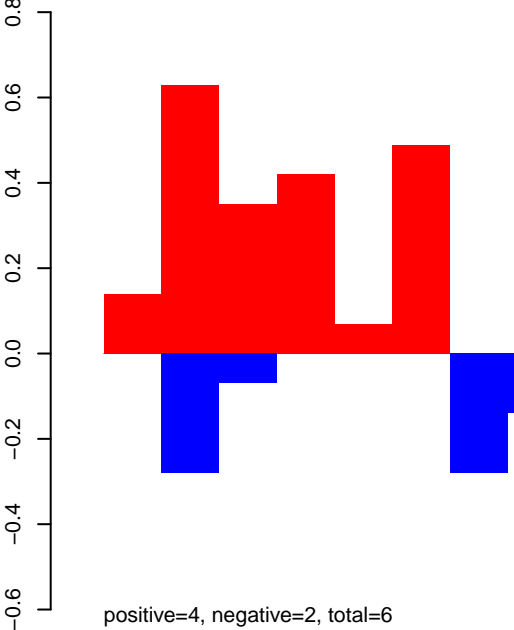


Window size=25, length=724, TE@rnd-4_family-507@DNA@3452:1-724

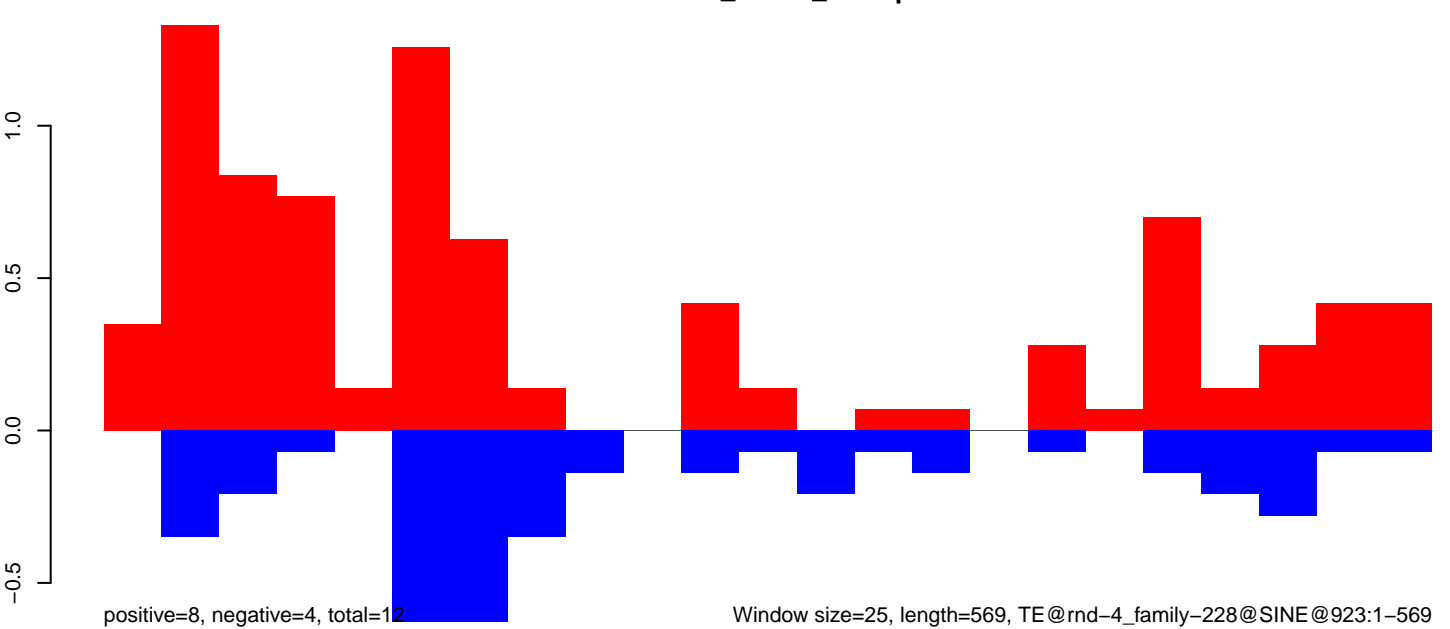
CuQuin_Mcarc_NL.18_23.rep



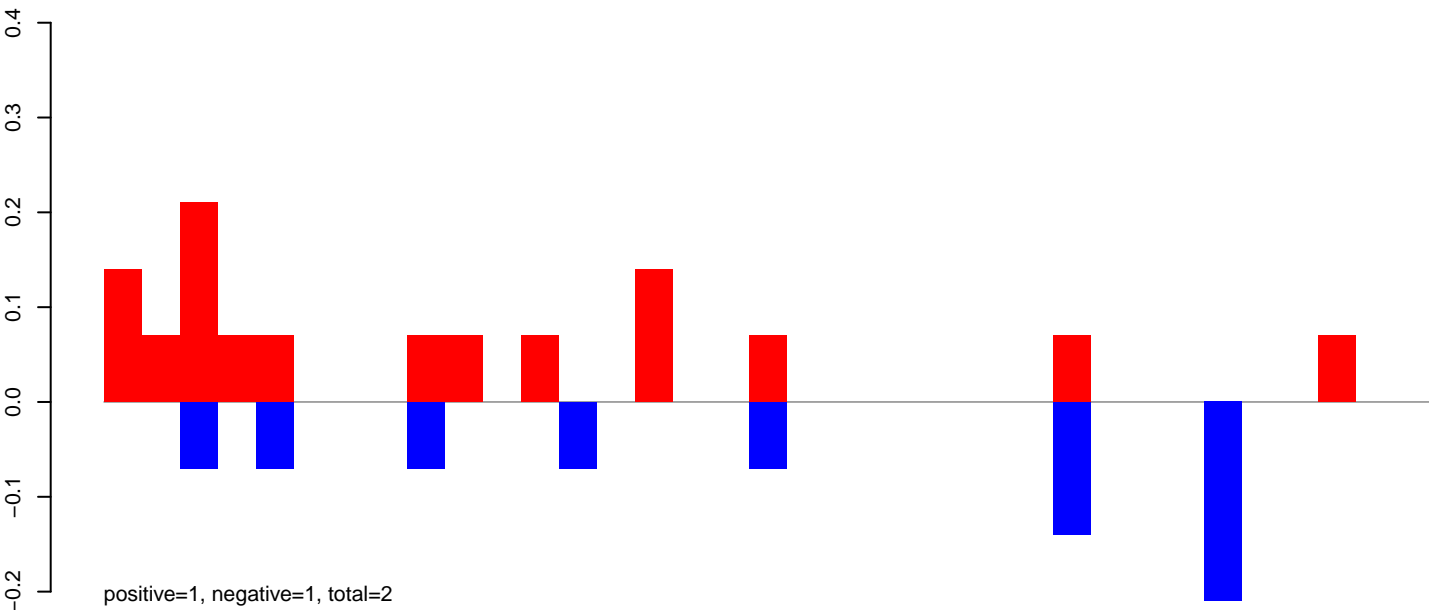
CuQuin_Mcarc_NL.24_35.rep



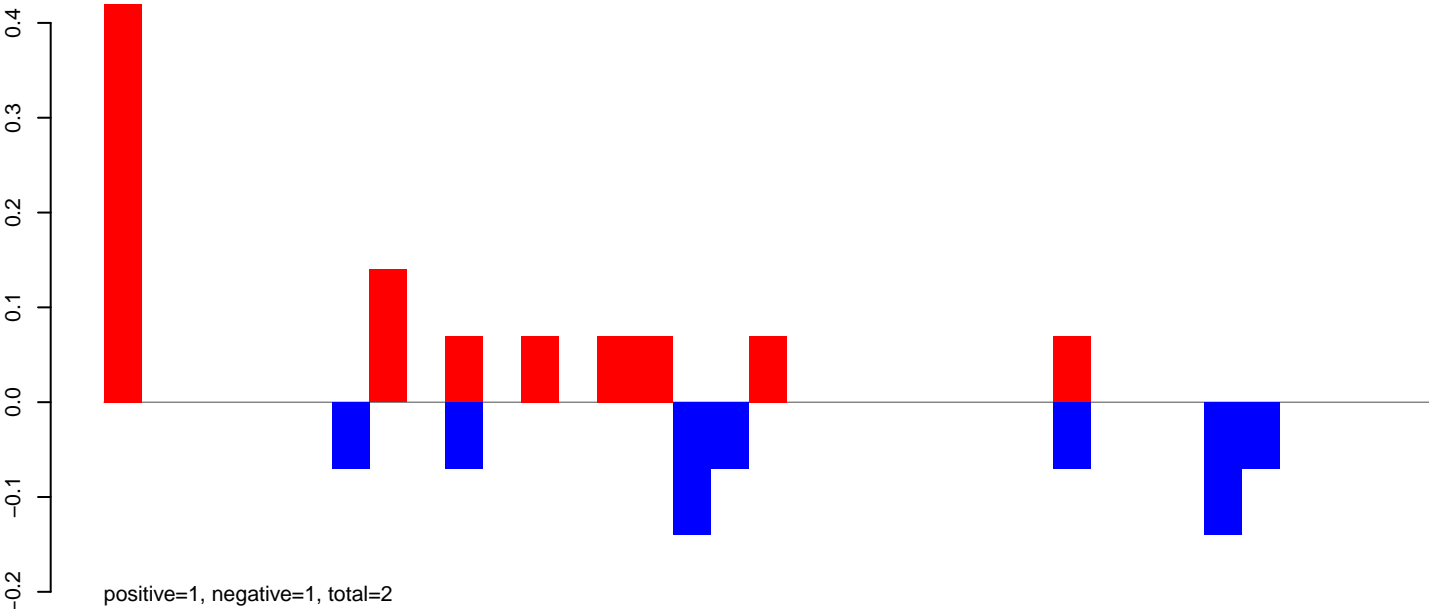
CuQuin_Mcarc_NL.rep



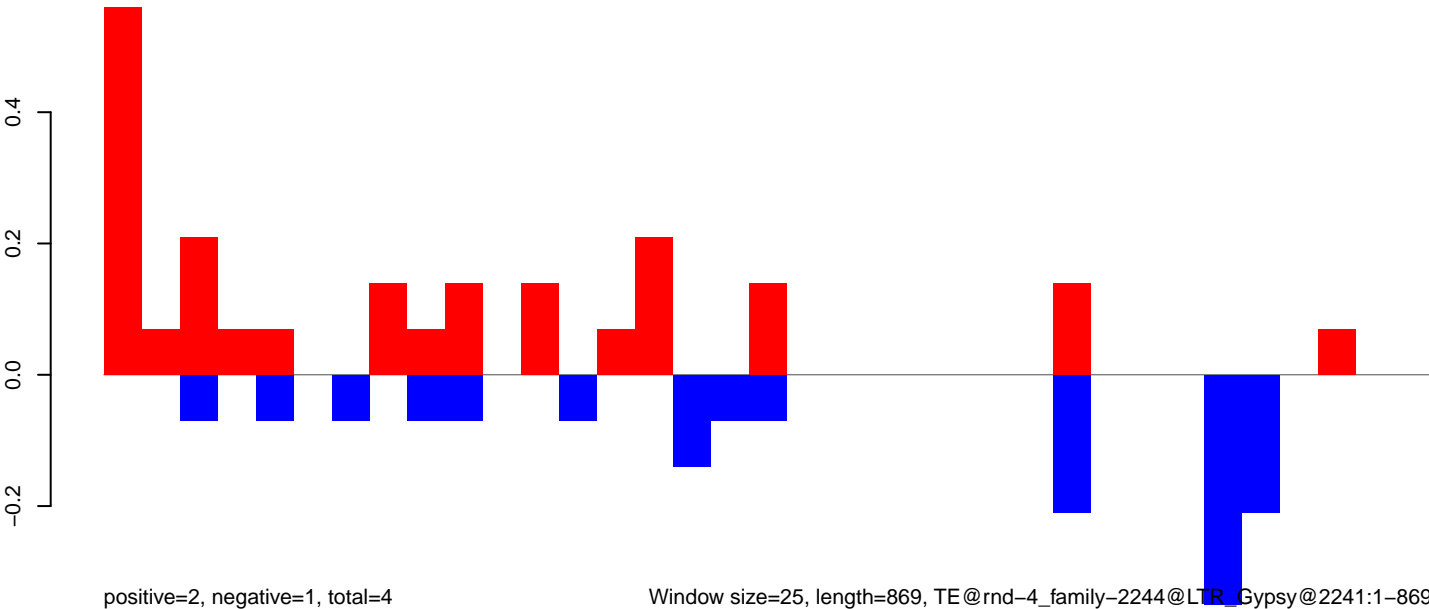
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

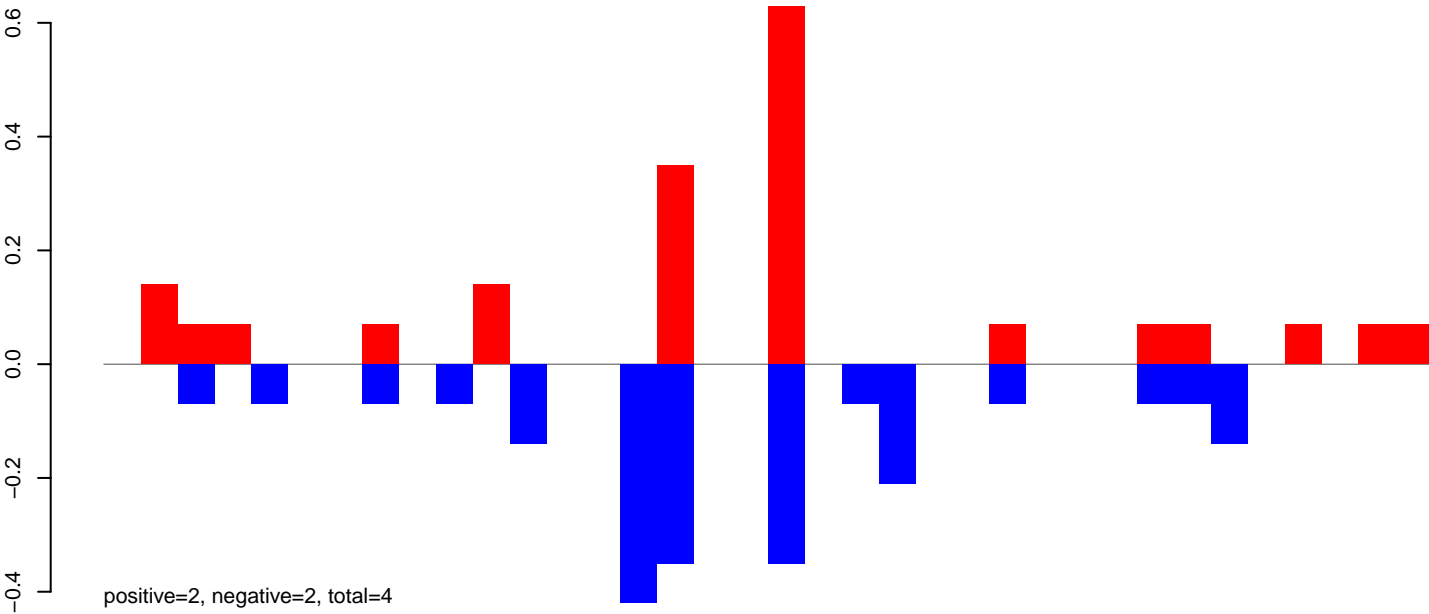


CuQuin_Mcarc_NL.rep

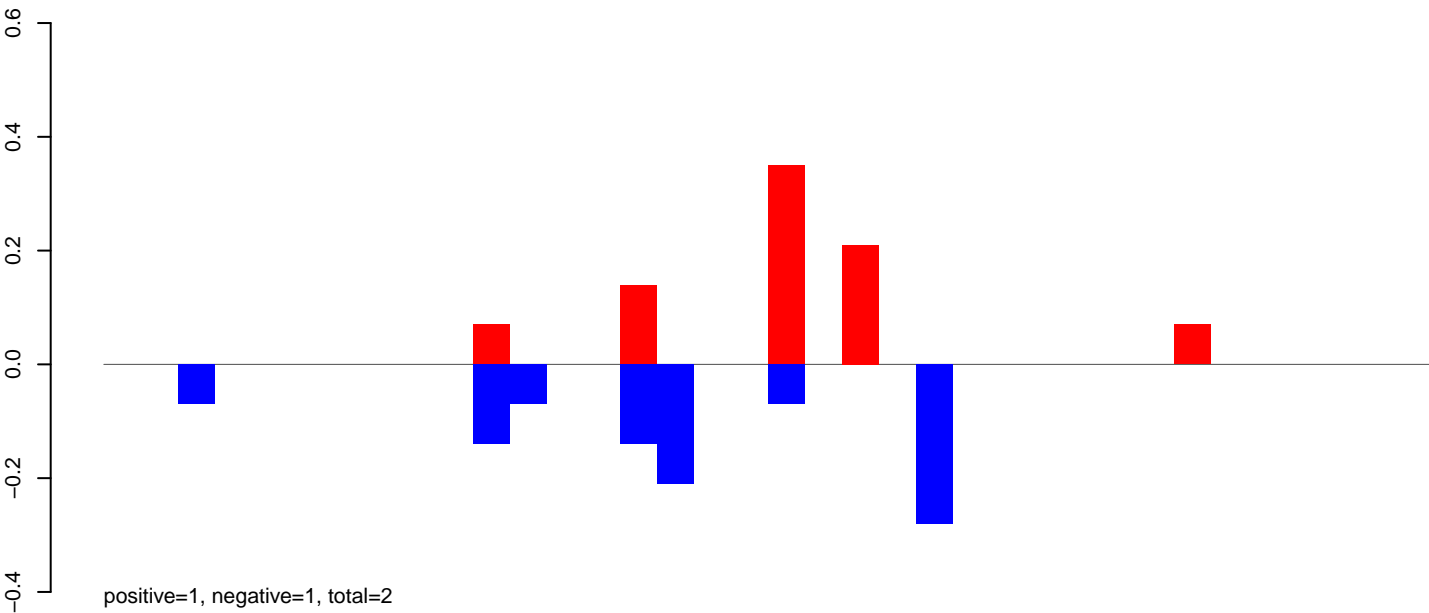


Window size=25, length=869, TE@rnd-4_family-2244@LTR_Gypsy@2241:1-869

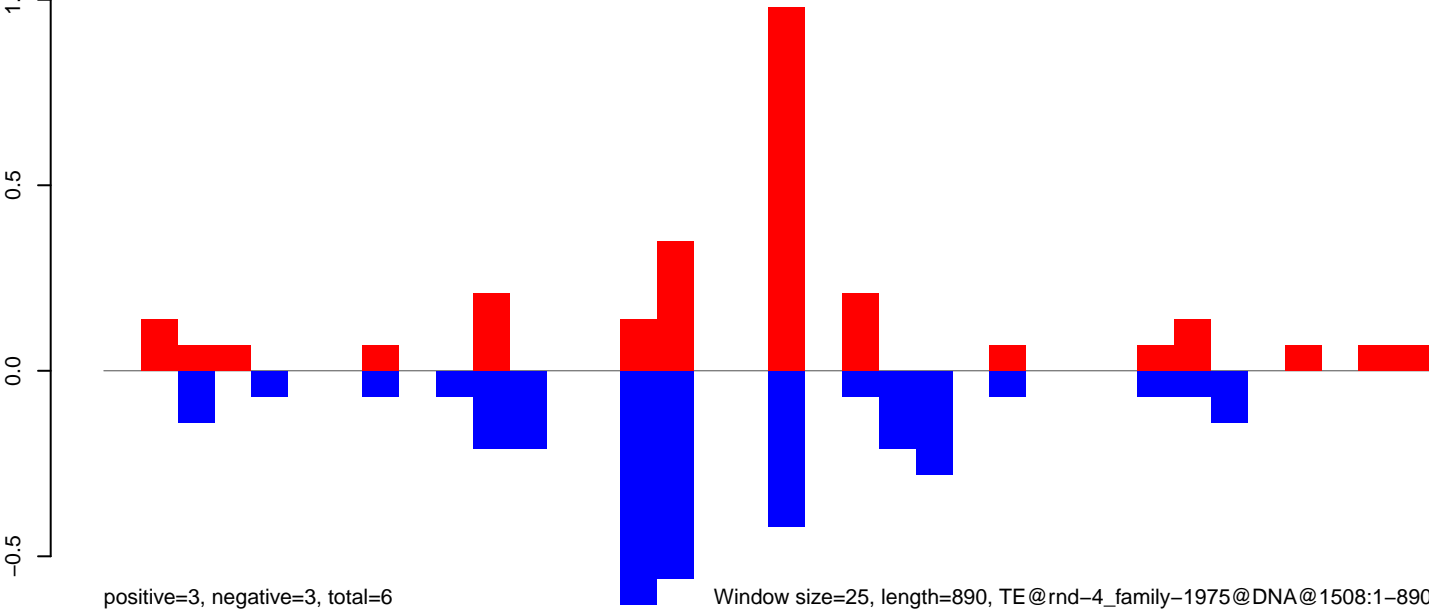
CuQuin_Mcarc_NL.18_23.rep



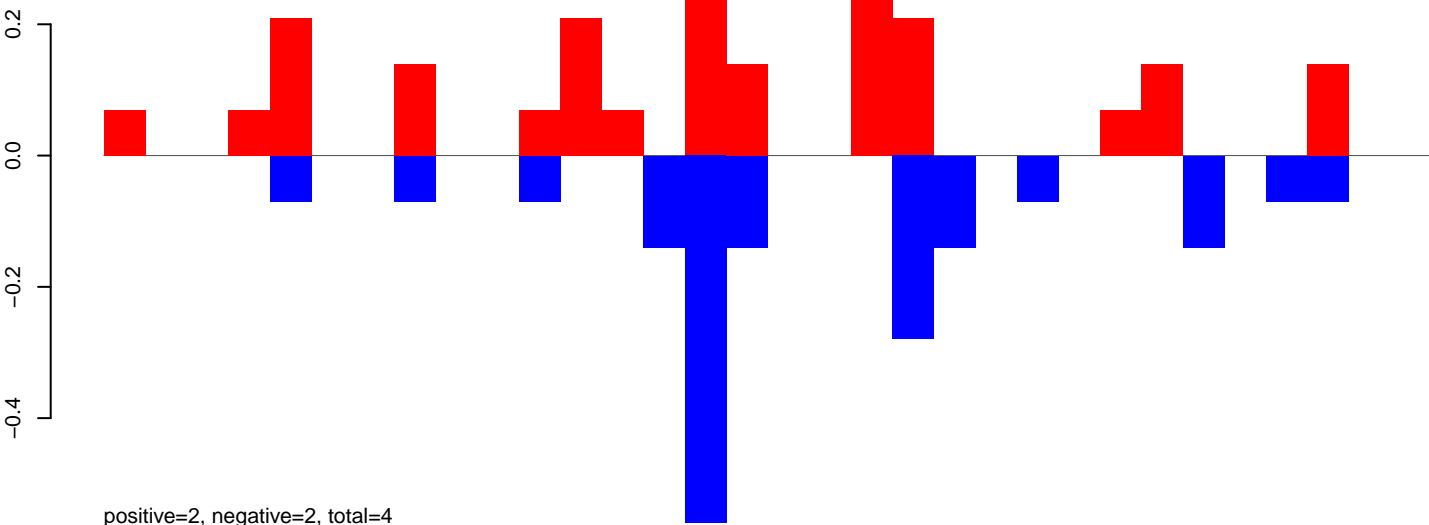
CuQuin_Mcarc_NL.24_35.rep



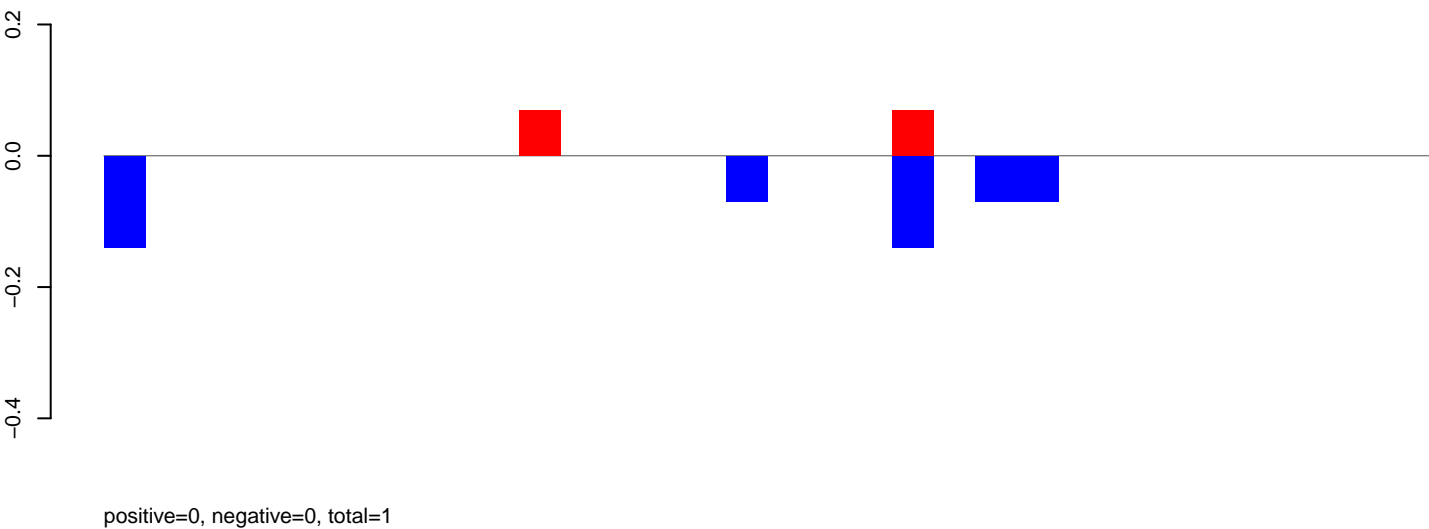
CuQuin_Mcarc_NL.rep



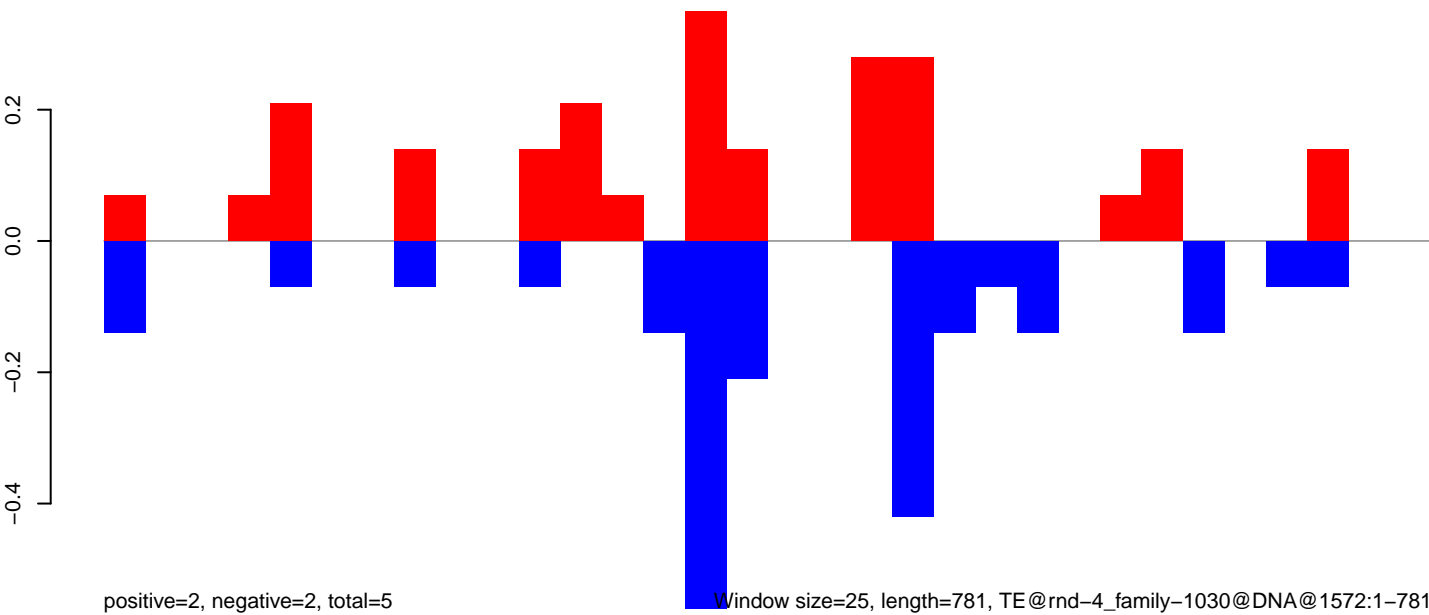
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

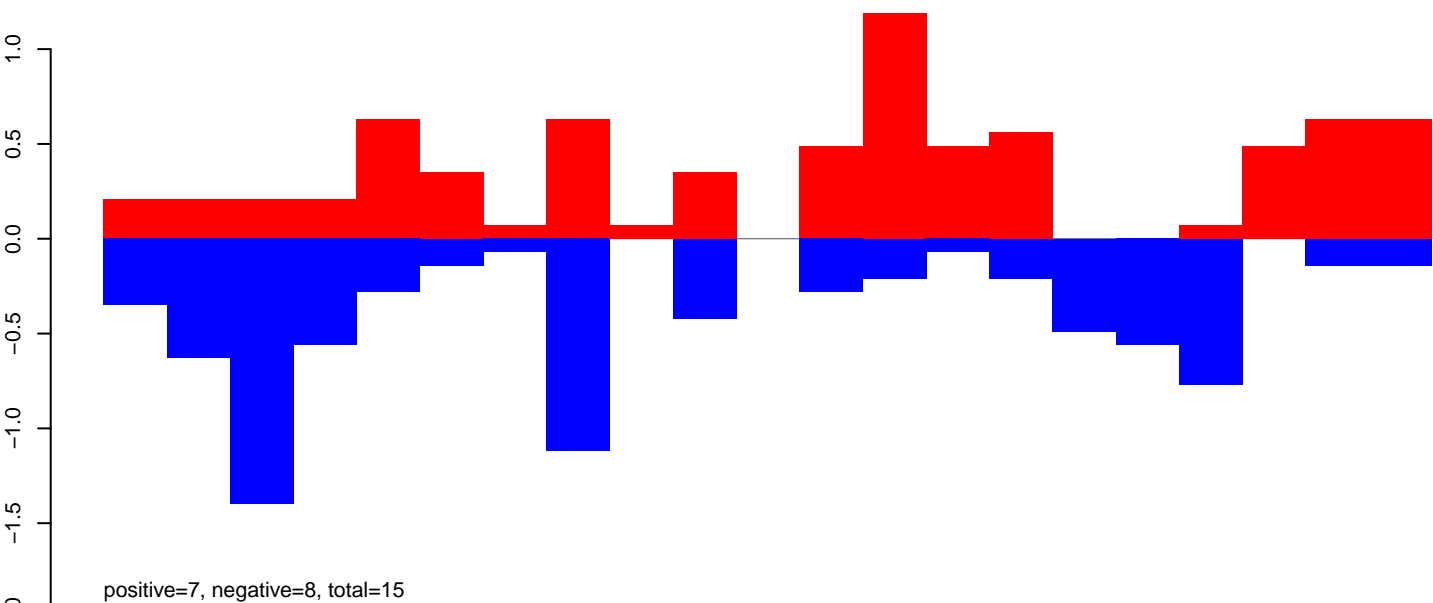


CuQuin_Mcarc_NL.rep

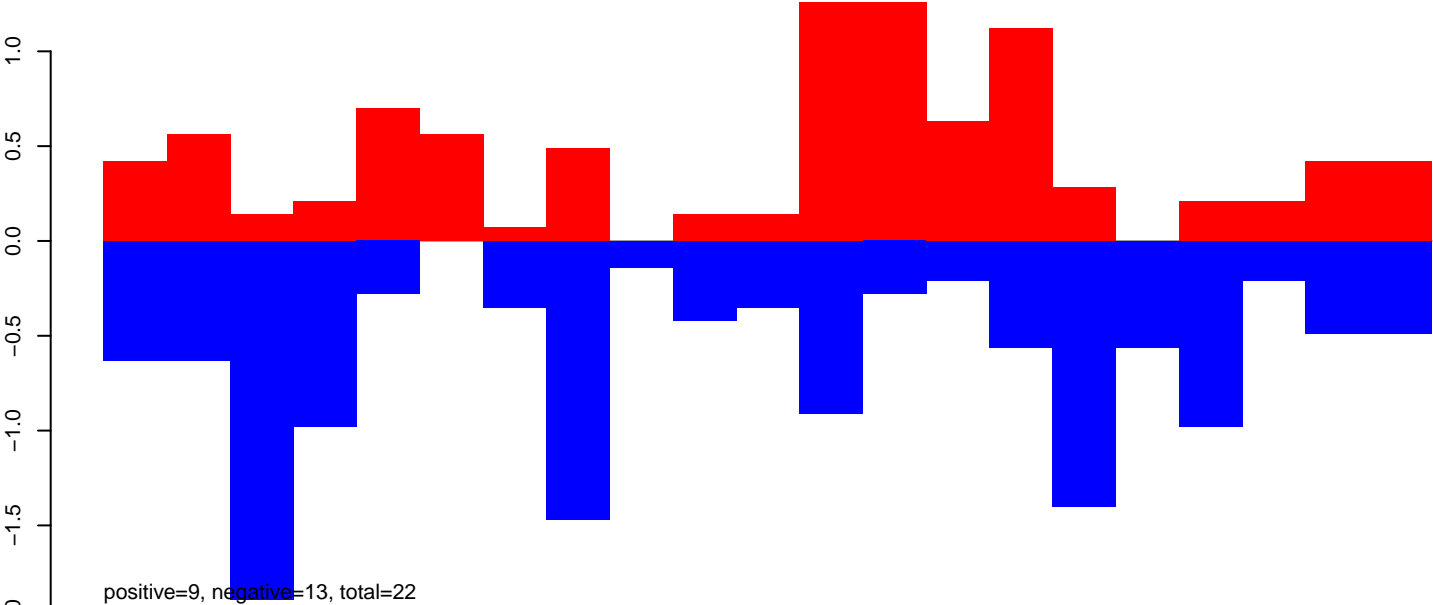


Window size=25, length=781, TE@rnd-4_family-1030@DNA@1572:1-781

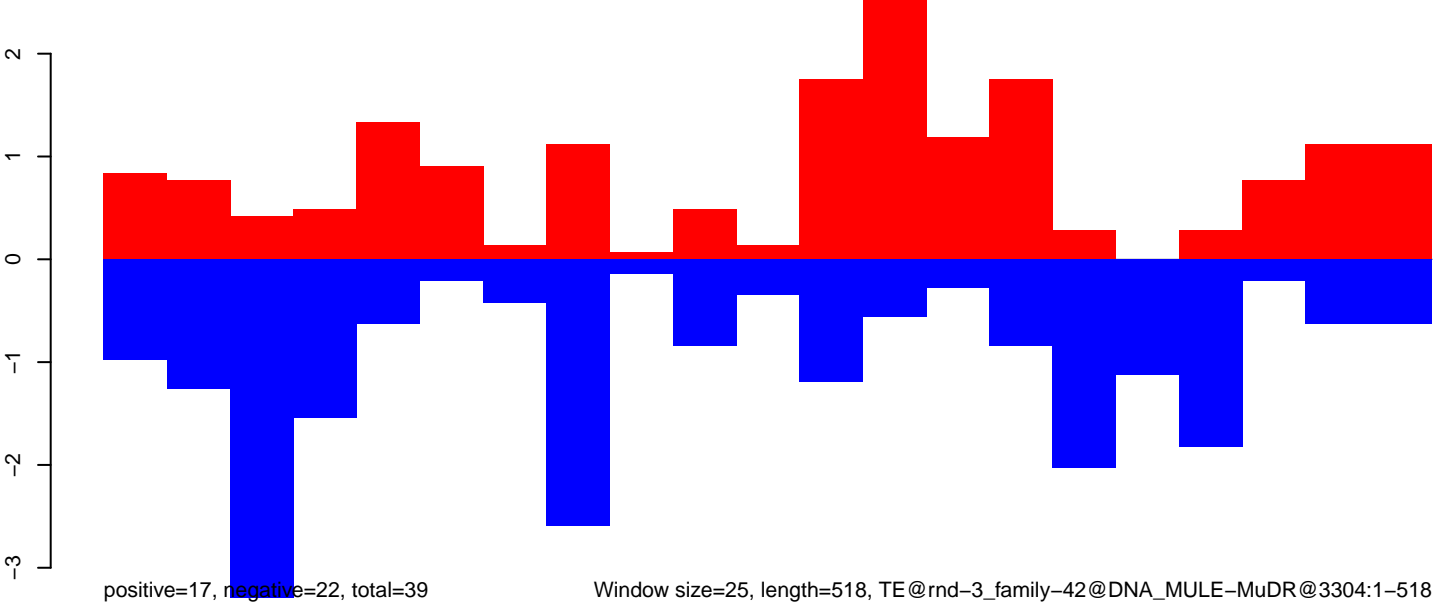
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

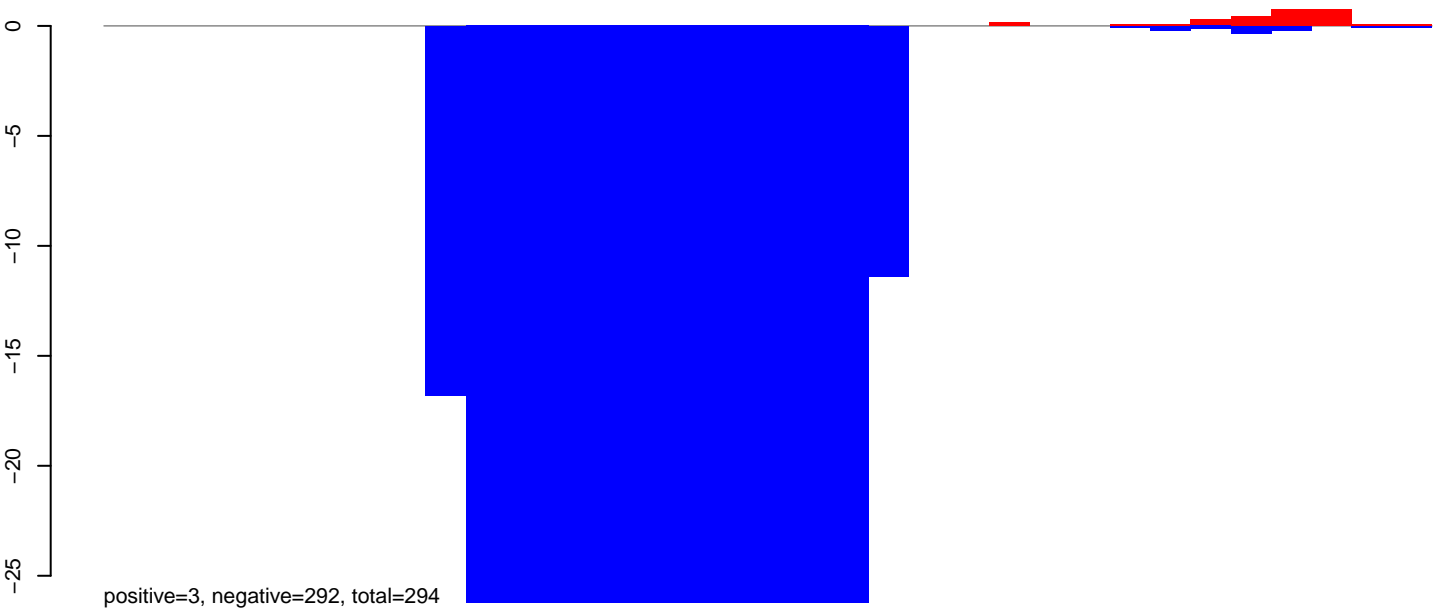


CuQuin_Mcarc_NL.rep



Window size=25, length=518, TE@rnd-3_family-42@DNA_MULE-MuDR@3304:1-518

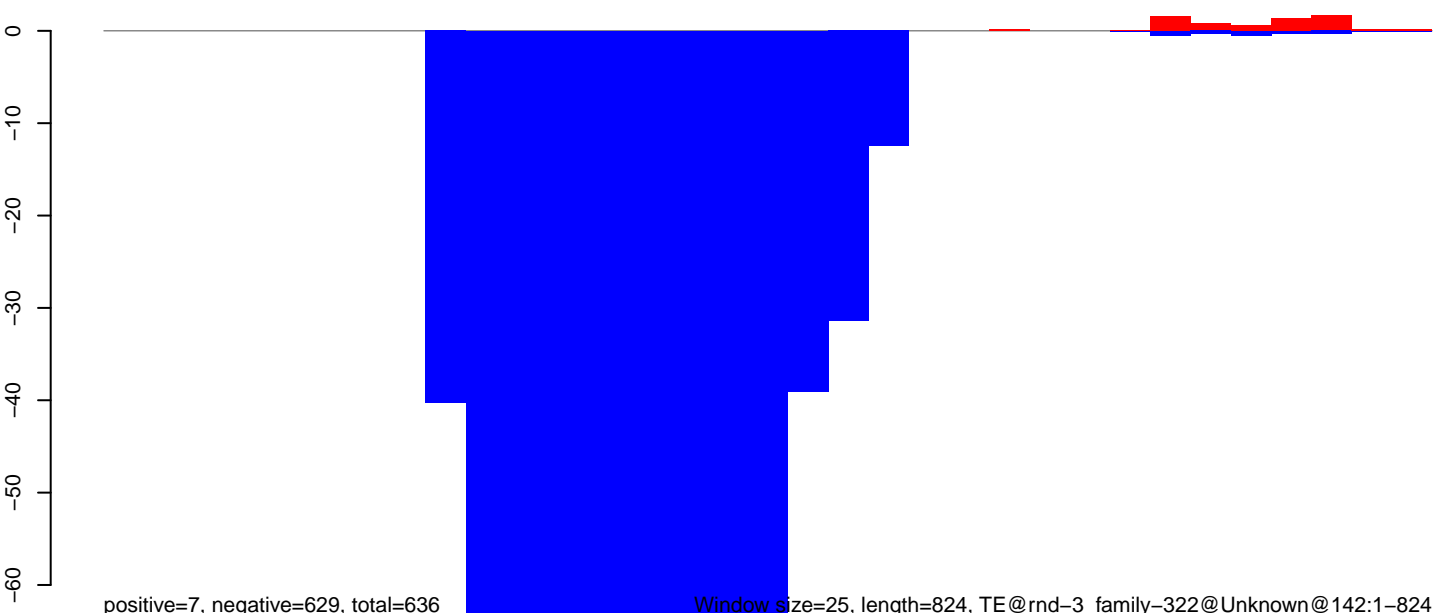
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

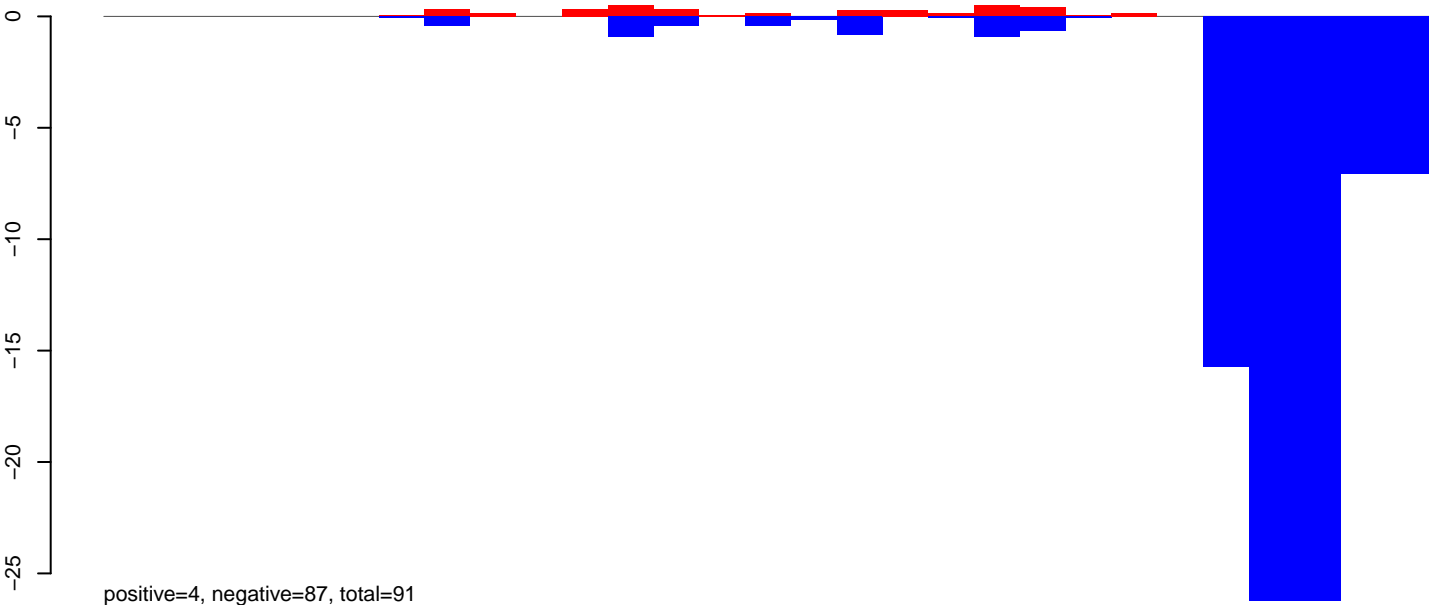


CuQuin_Mcarc_NL.rep

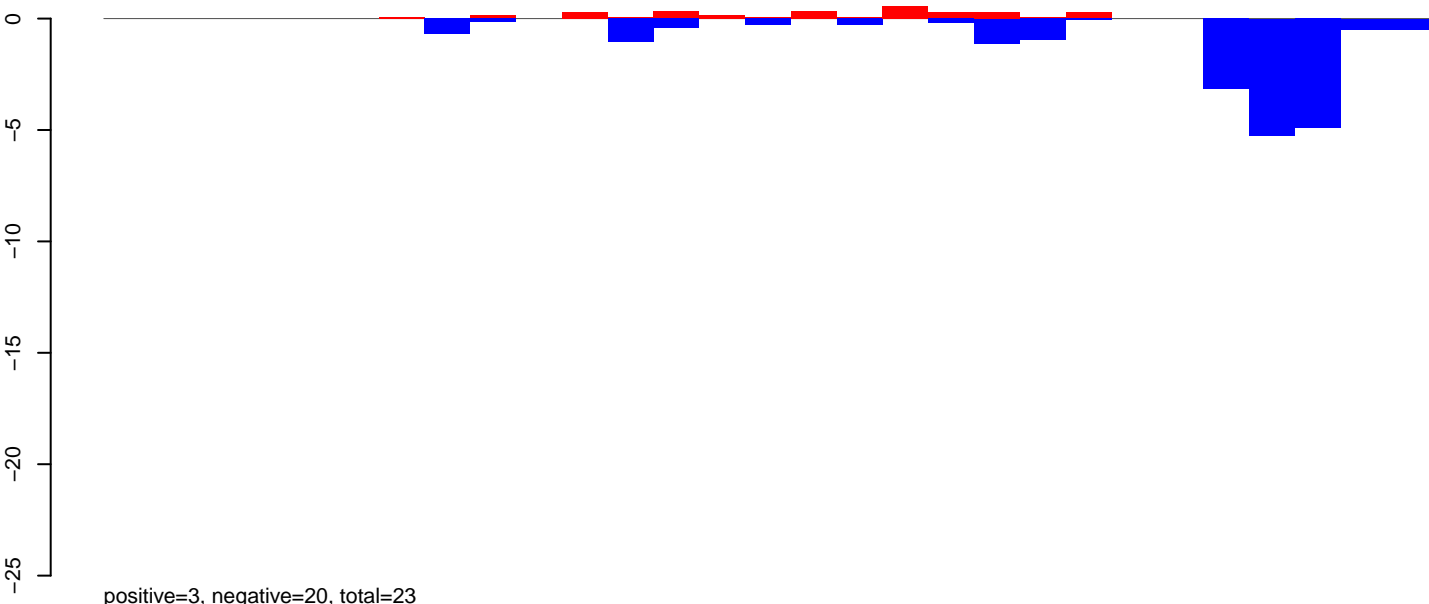


Window size=25, length=824, TE@rnd-3_family-322@Unknown@142:1-824

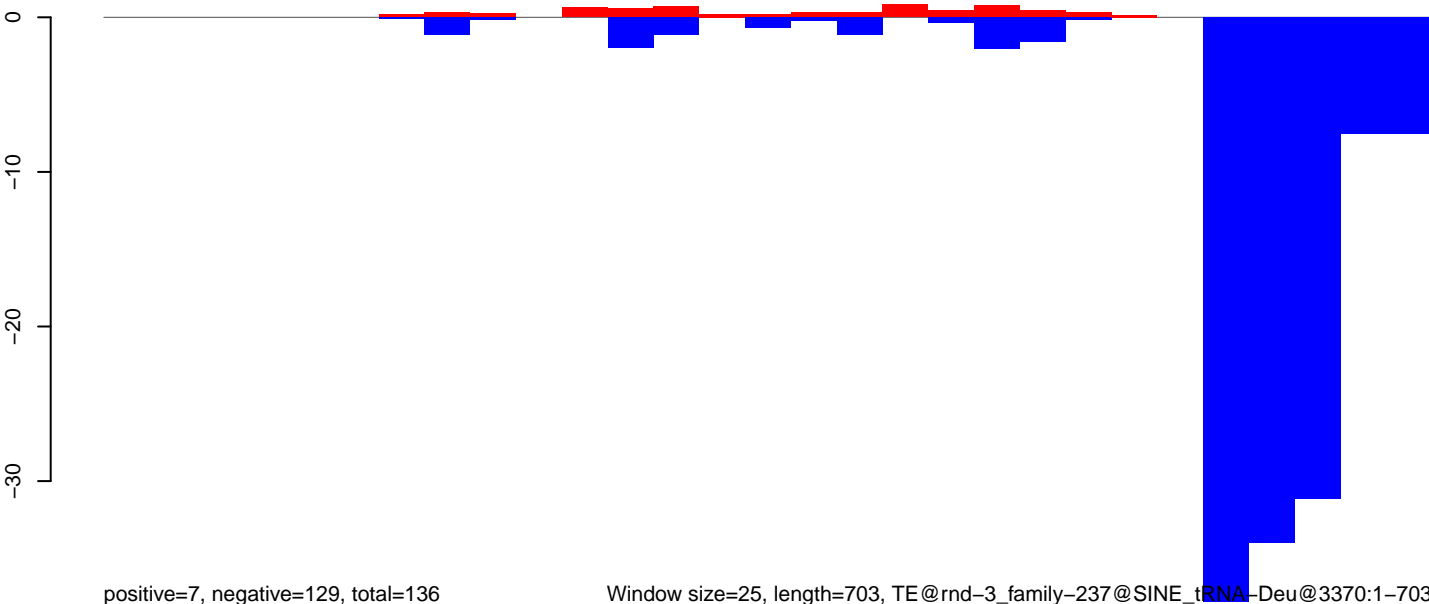
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



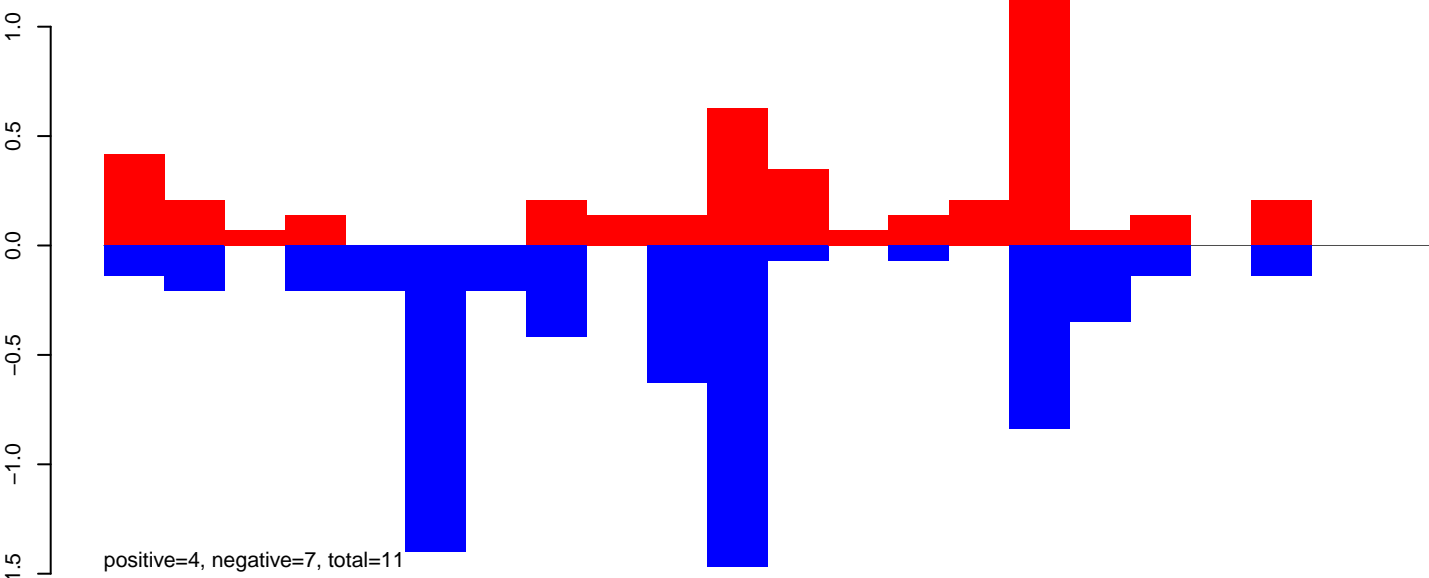
CuQuin_Mcarc_NL.rep



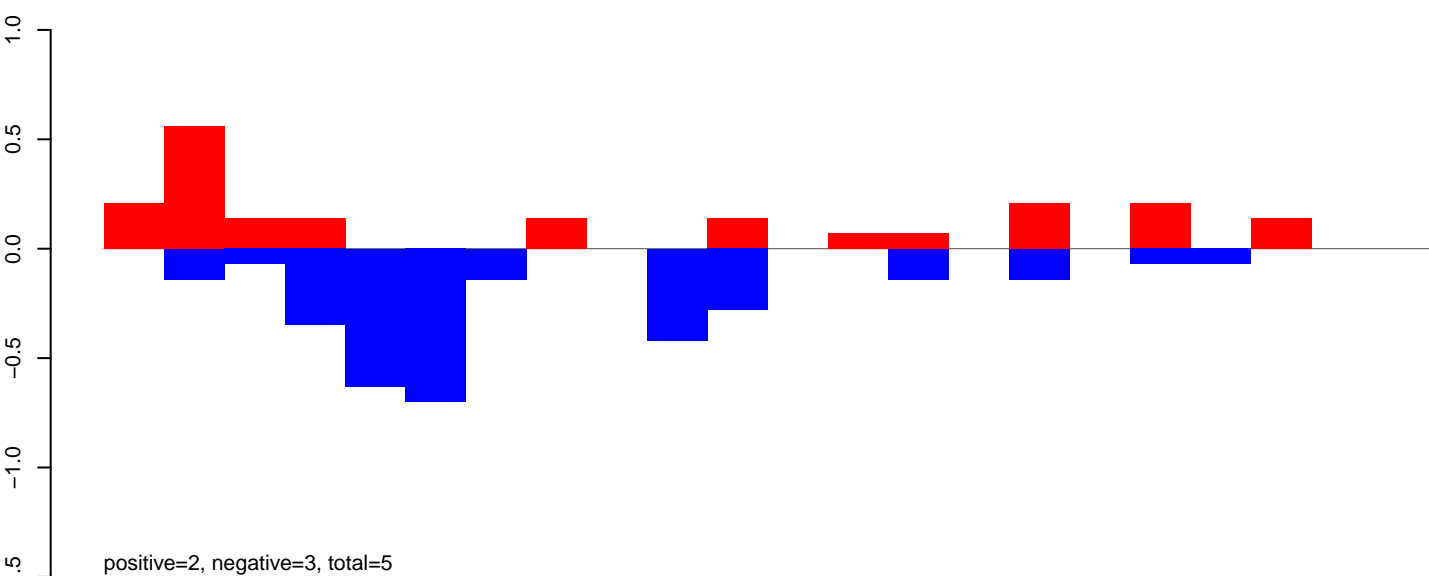
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0 200 400 600

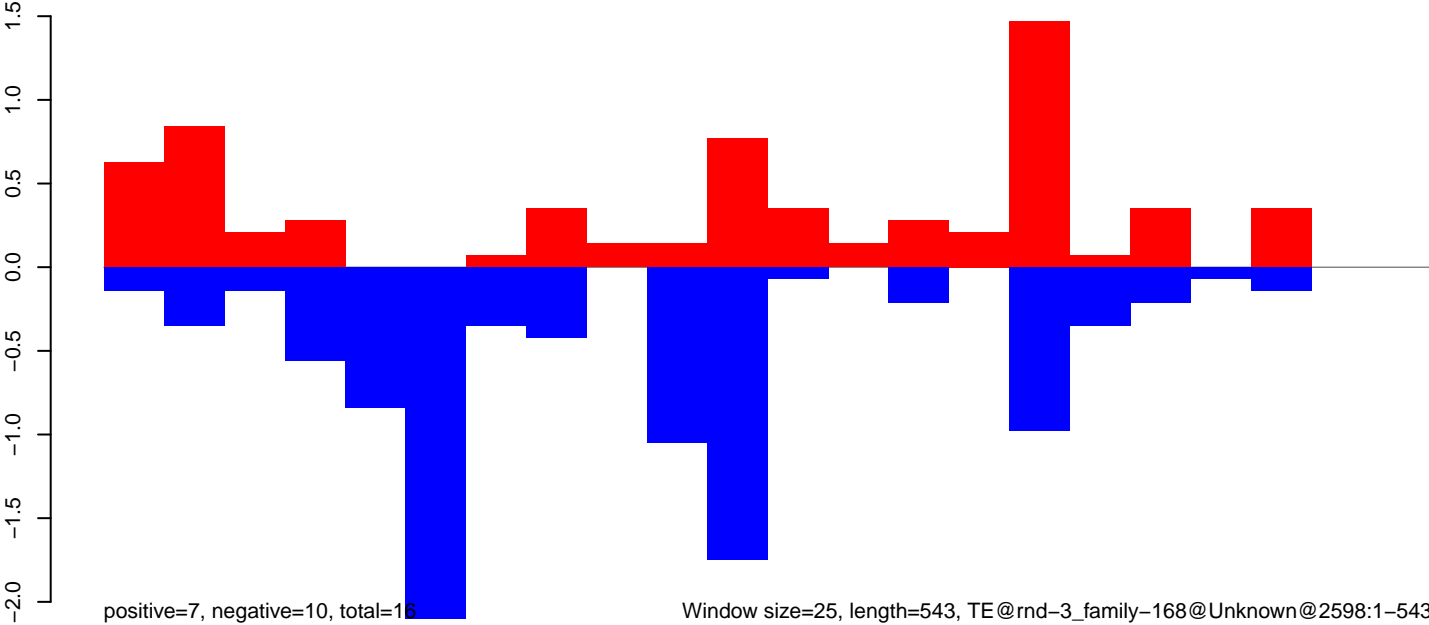
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

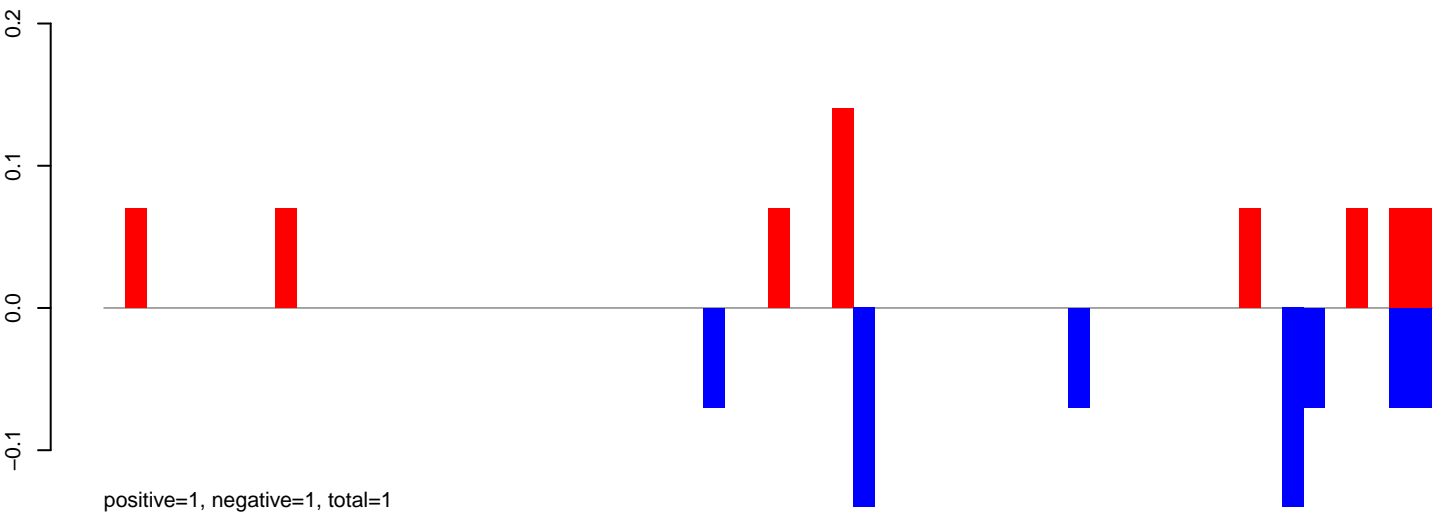


CuQuin_Mcarc_NL.rep

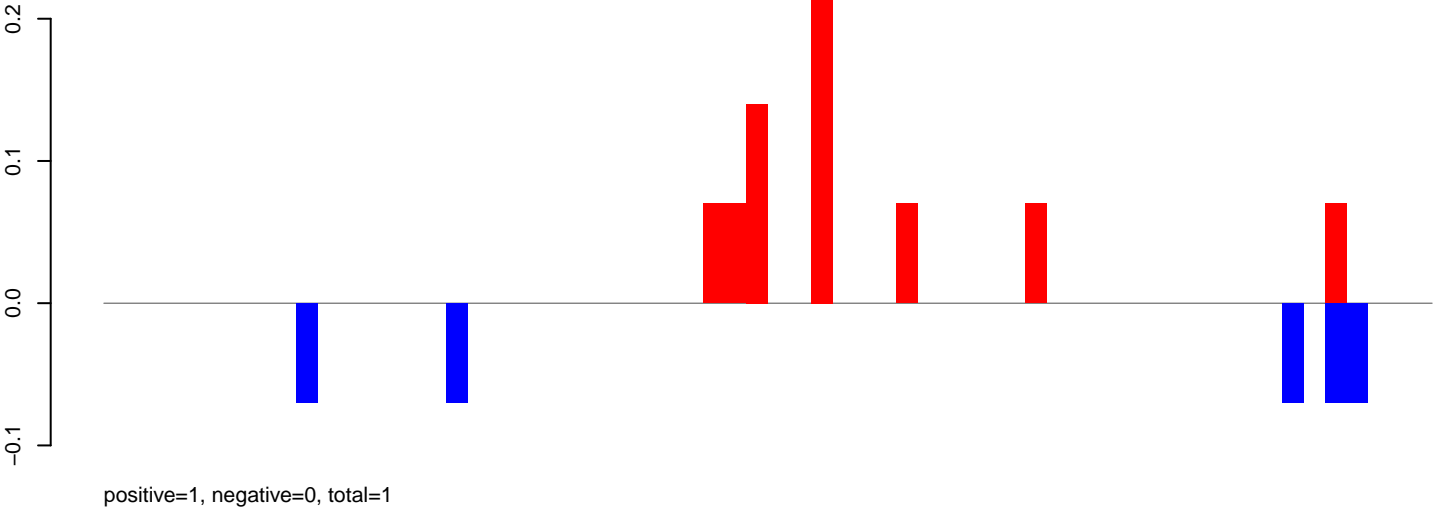


Window size=25, length=543, TE@rnd-3_family-168@Unknown@2598:1-543

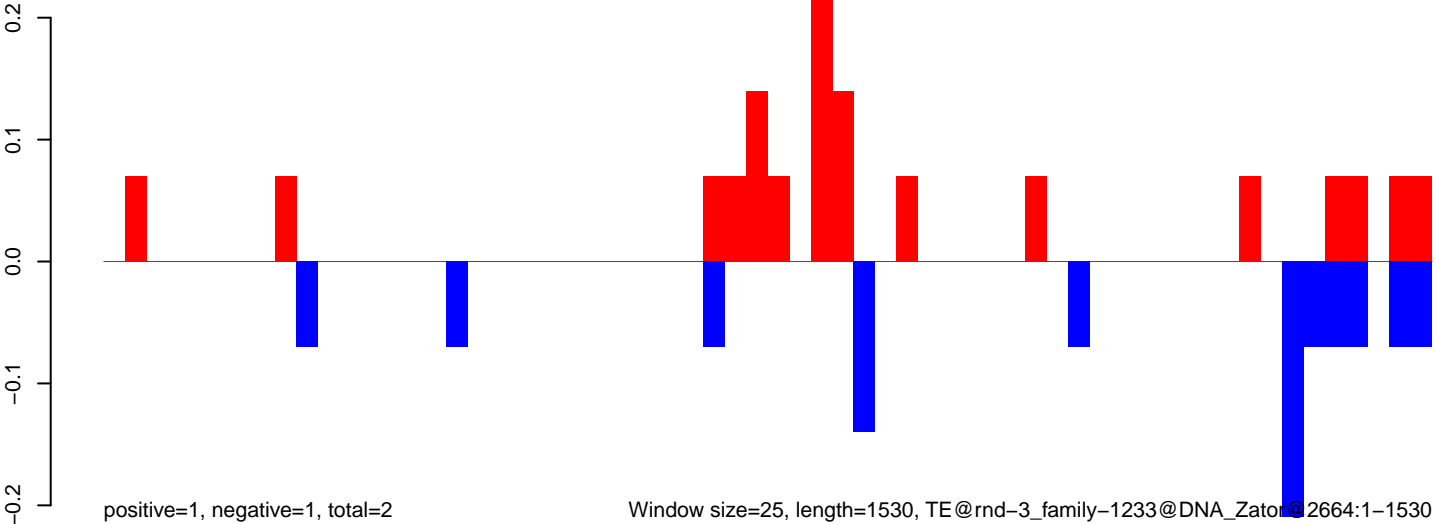
CuQuin_Mcarc_NL.18_23.rep



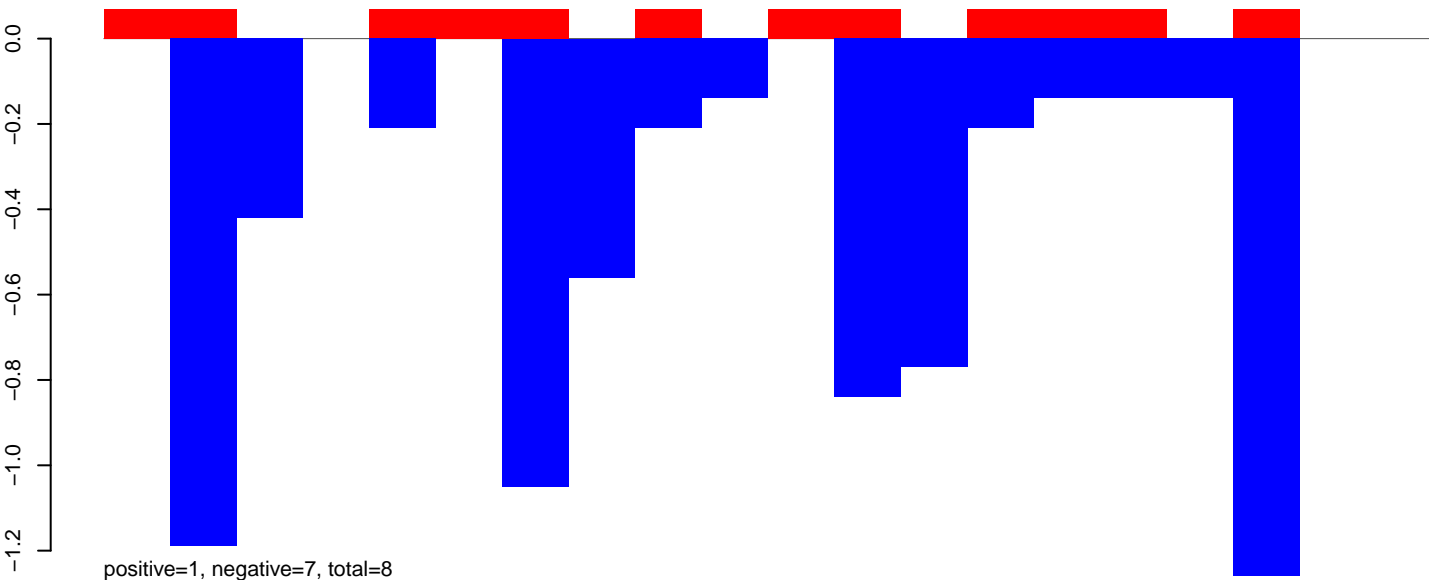
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

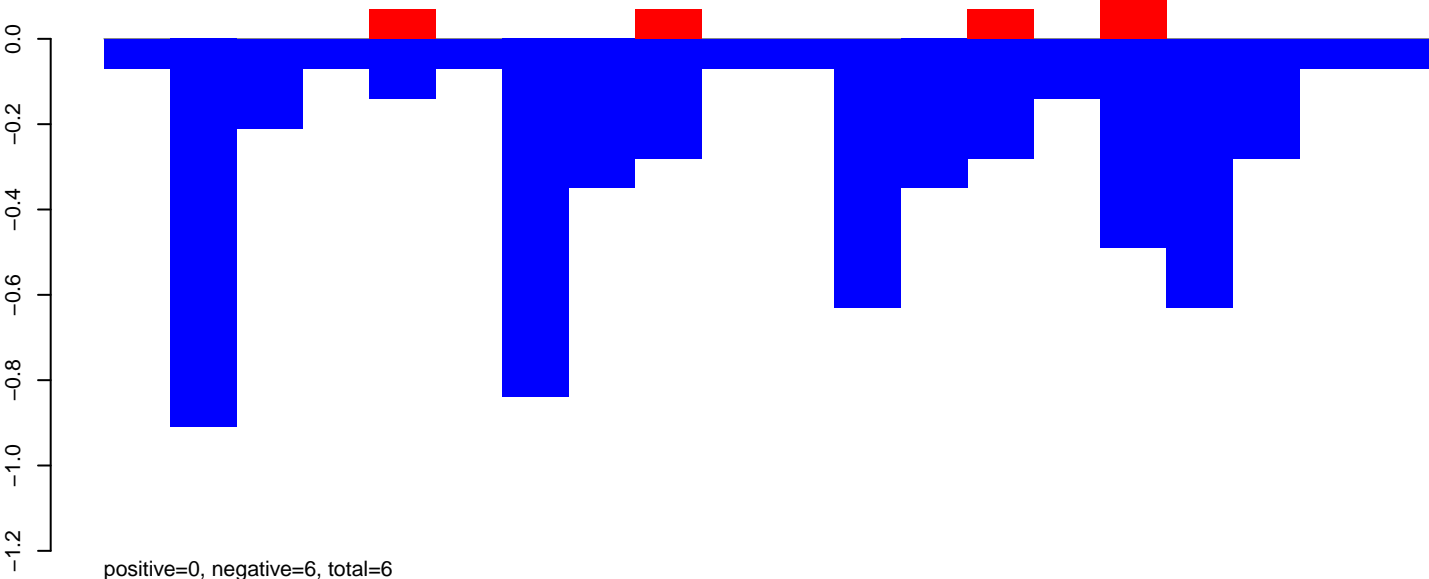


CuQuin_Mcarc_NL.18_23.rep



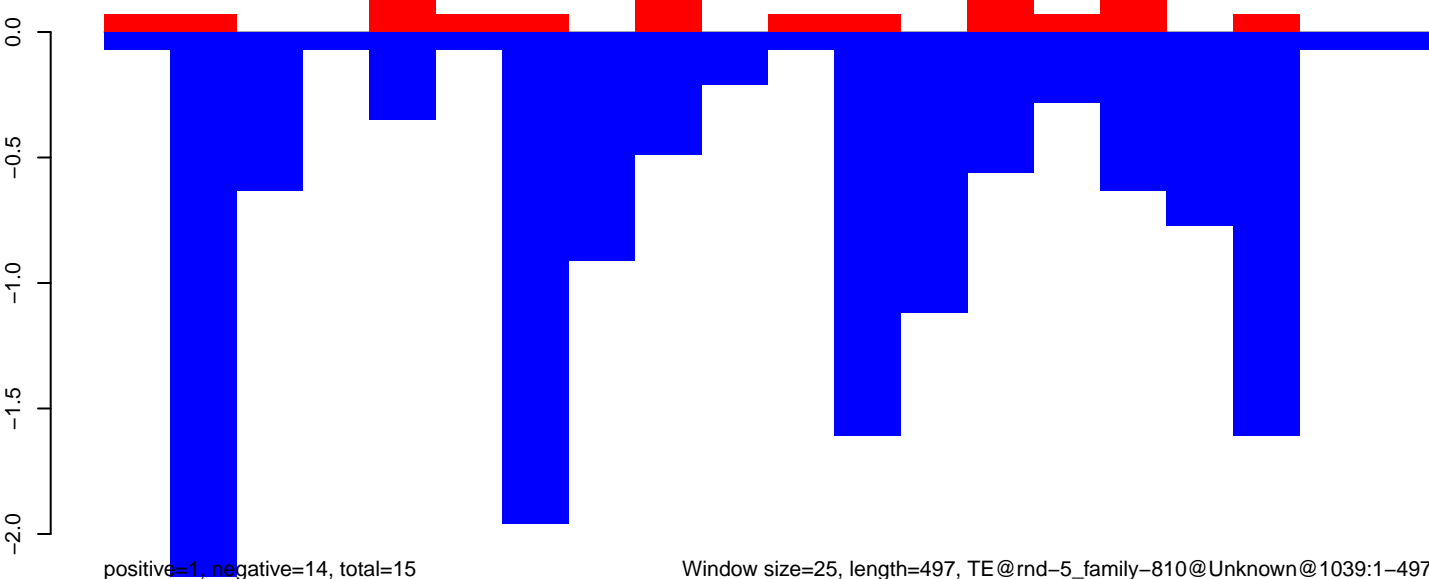
positive=1, negative=7, total=8

CuQuin_Mcarc_NL.24_35.rep



positive=0, negative=6, total=6

CuQuin_Mcarc_NL.rep

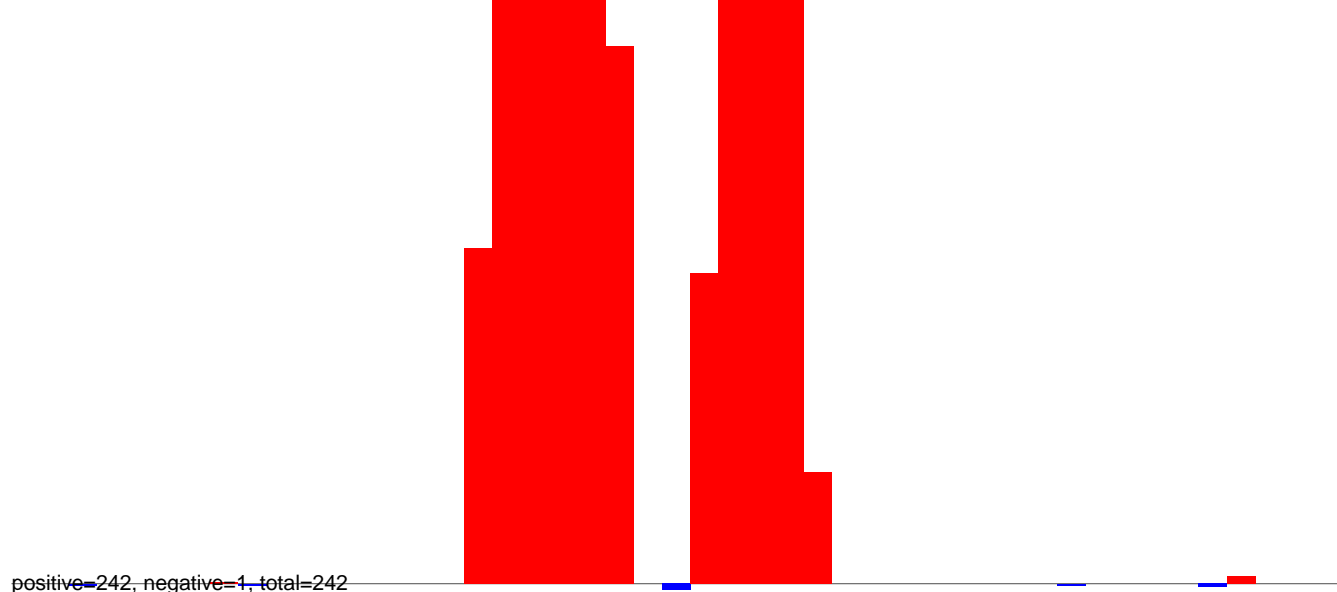


positive=1, negative=14, total=15

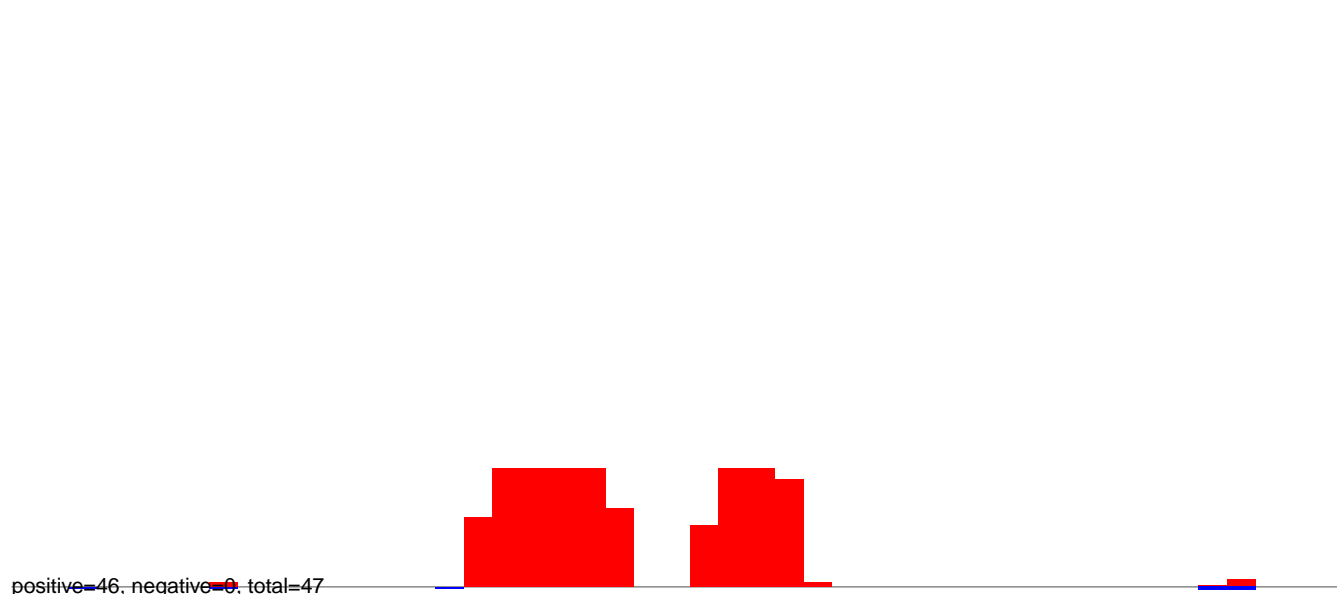
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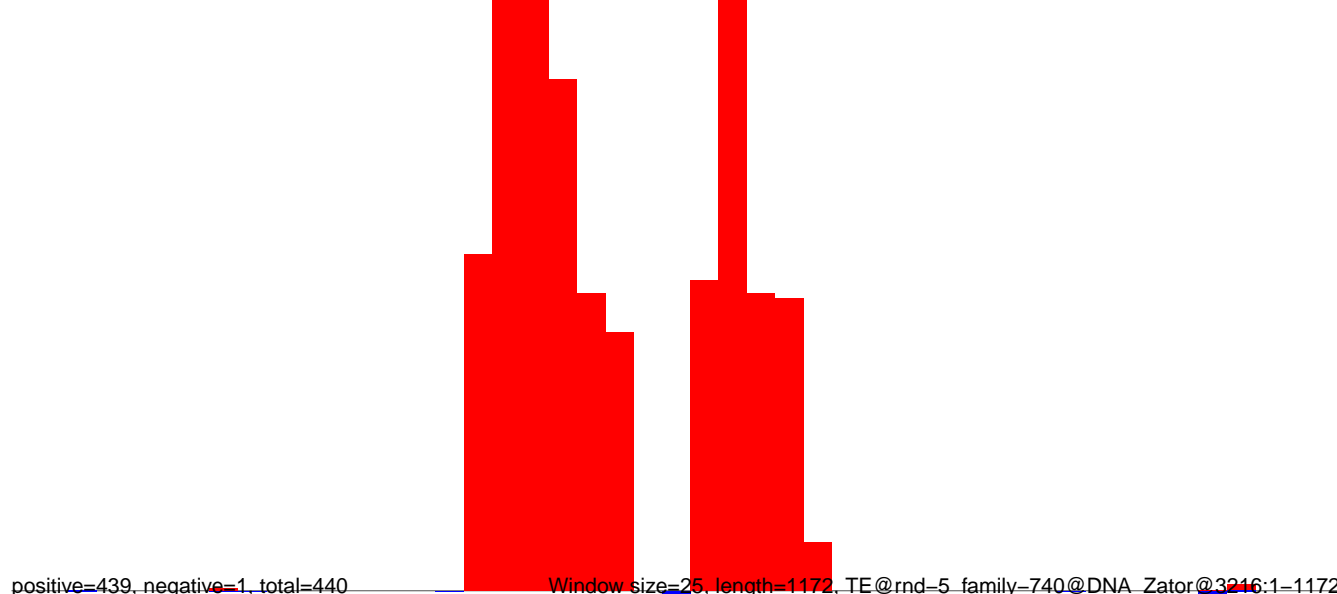
CuQuin_Mcarc_NL.18_23.rep



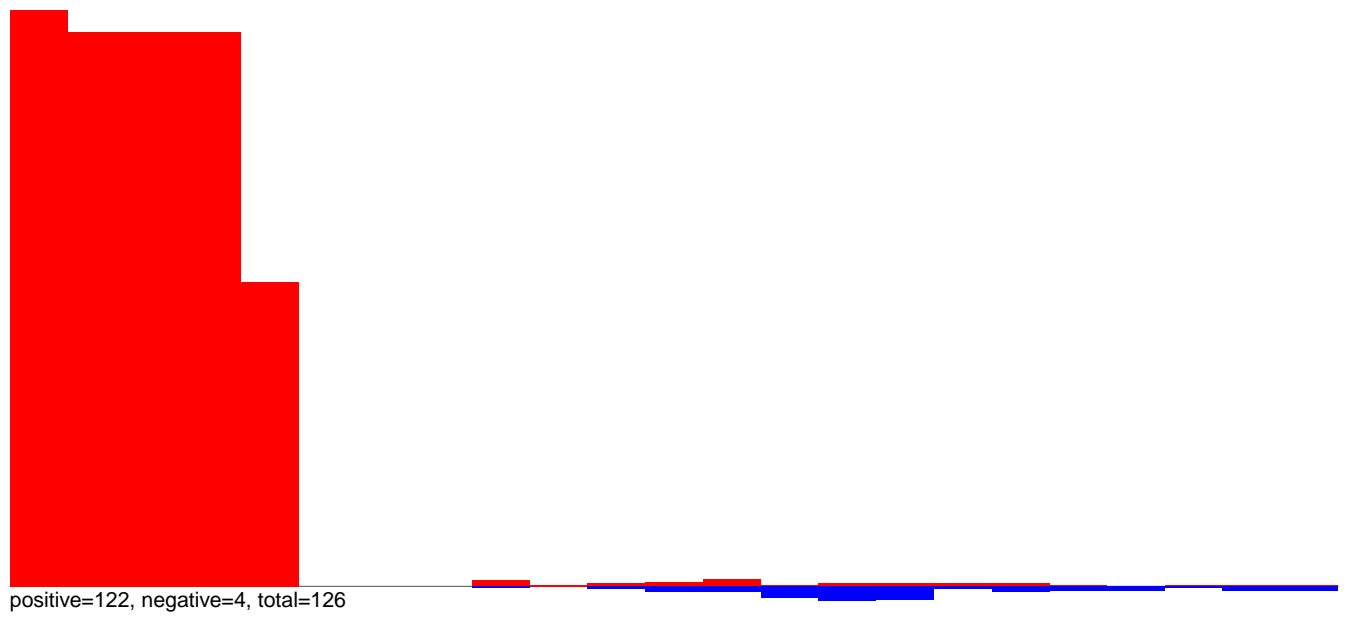
CuQuin_Mcarc_NL.24_35.rep



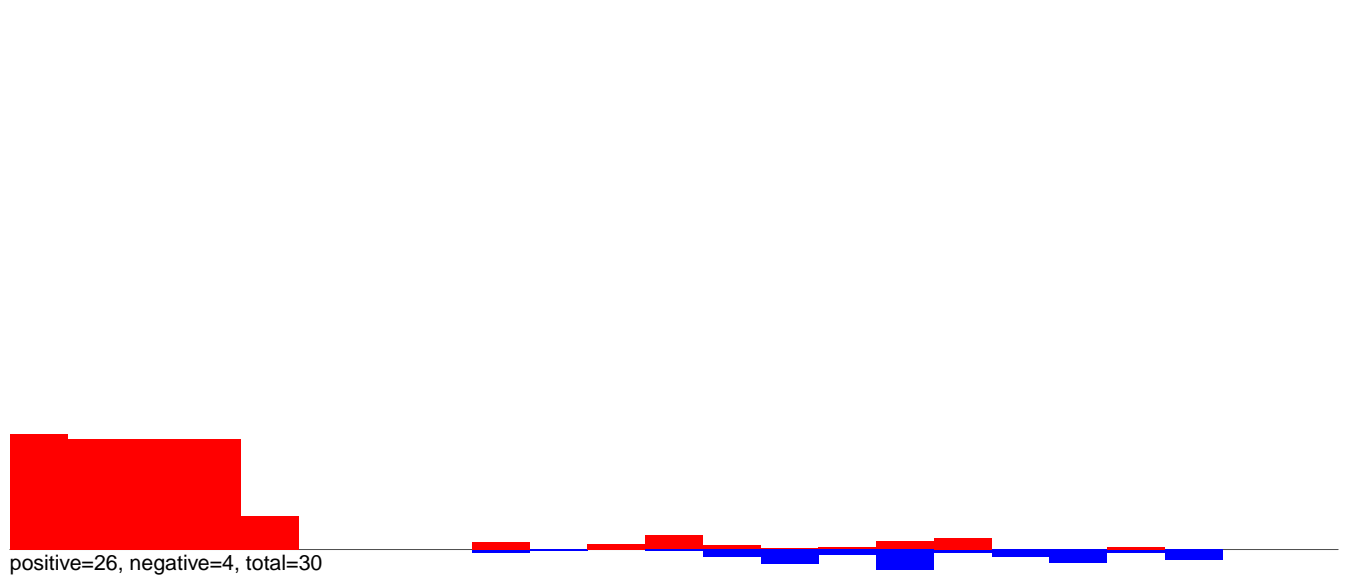
CuQuin_Mcarc_NL.rep



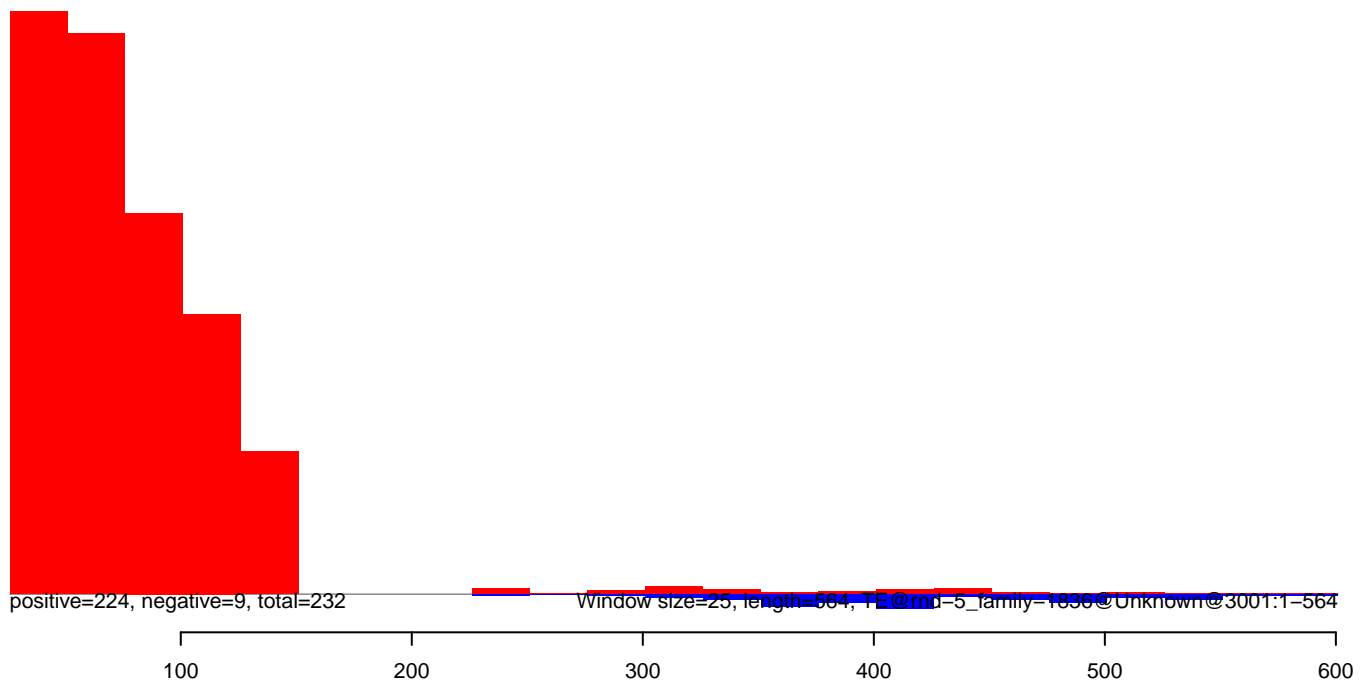
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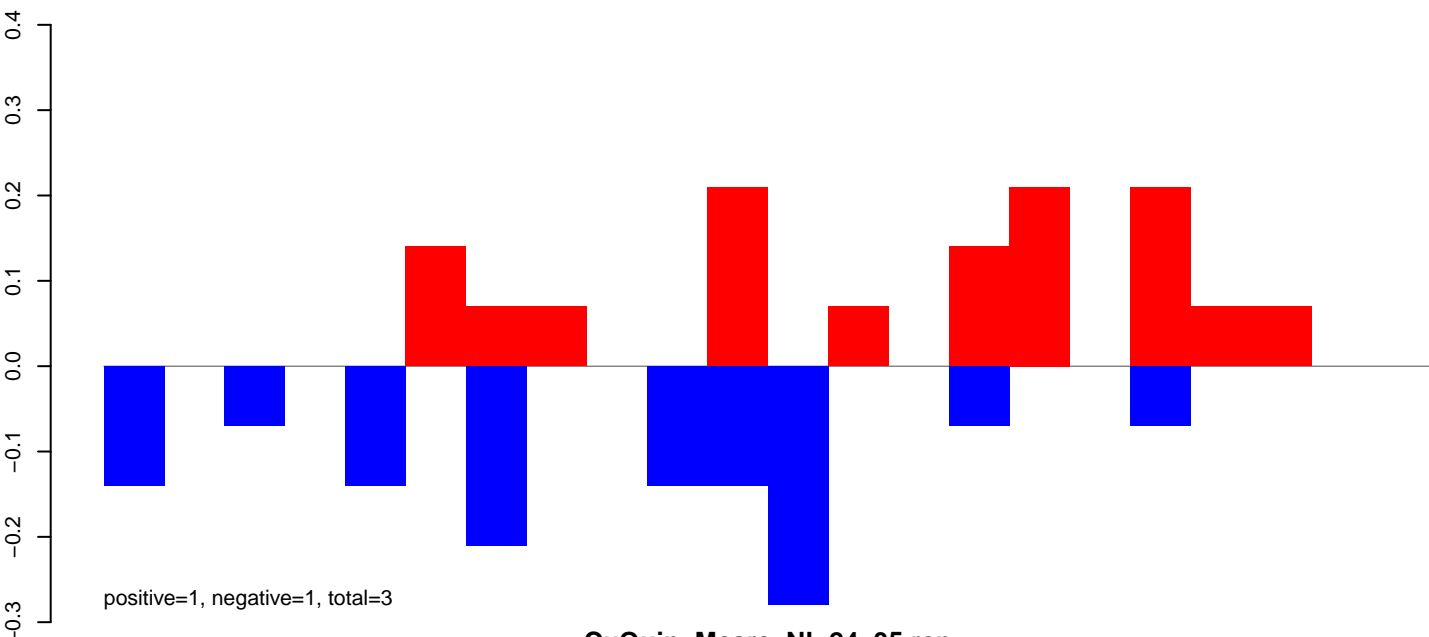
CuQuin_Mcarc_NL.24_35.rep



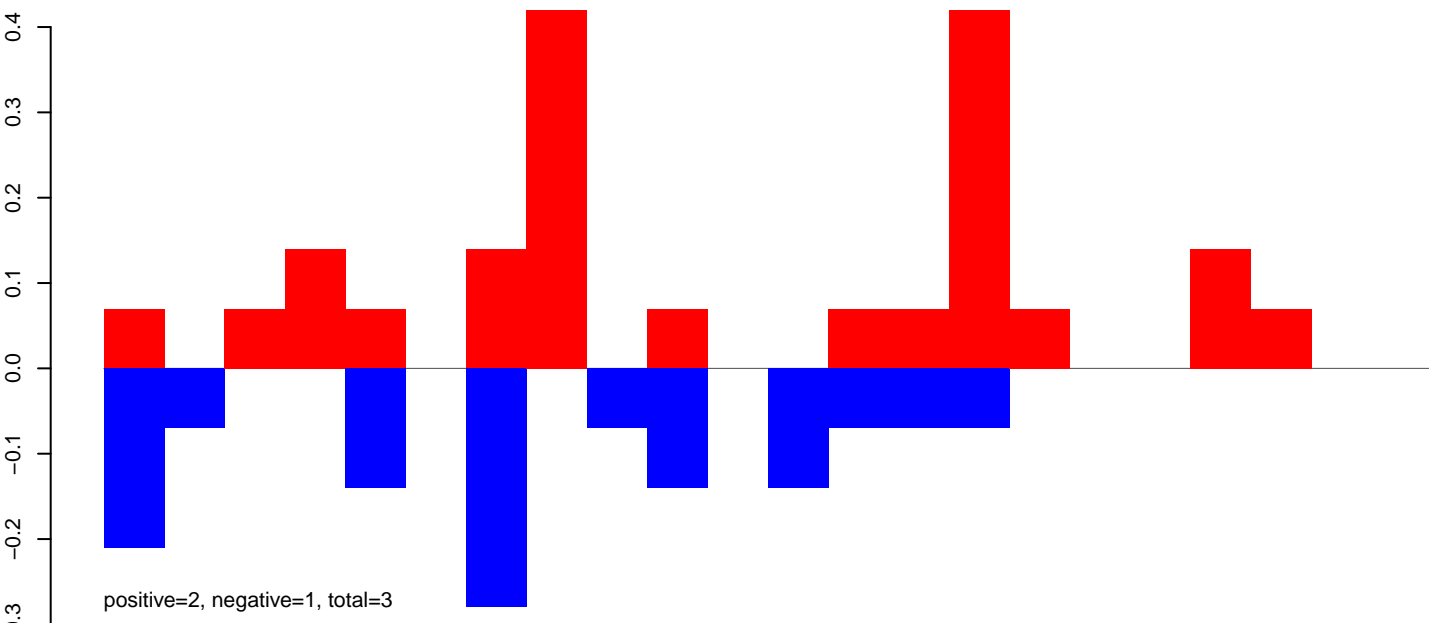
CuQuin_Mcarc_NL.rep



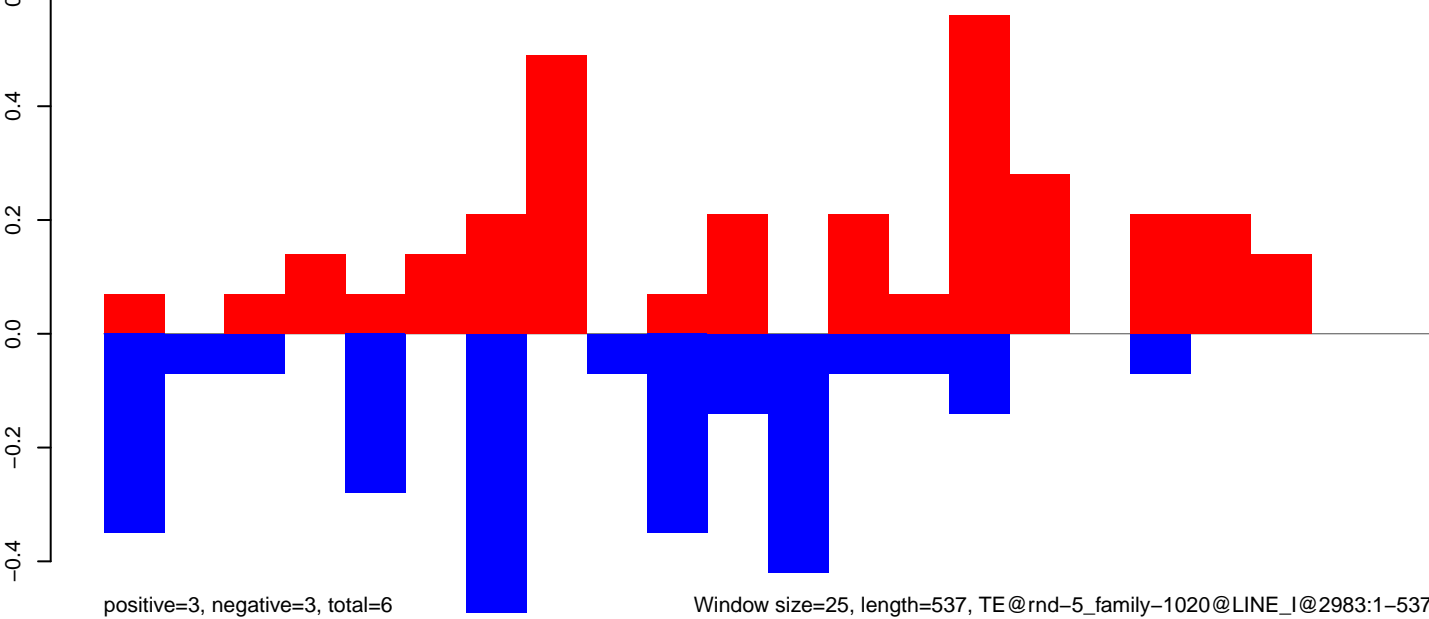
CuQuin_Mcarc_NL.18_23.rep



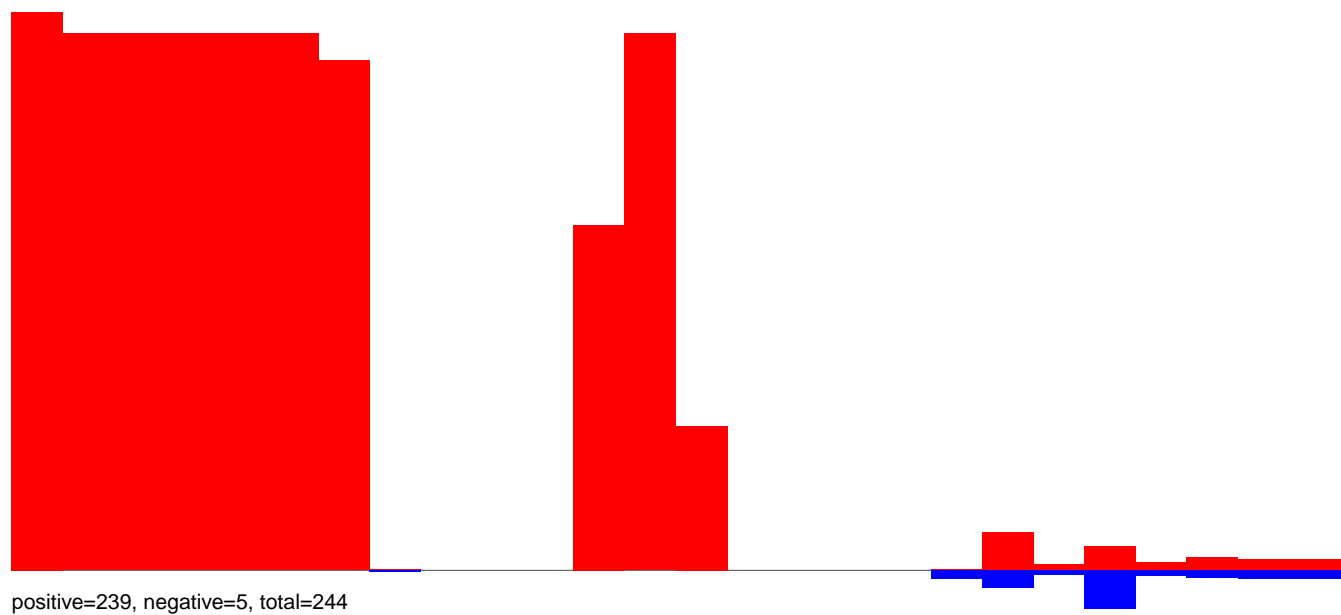
CuQuin_Mcarc_NL.24_35.rep



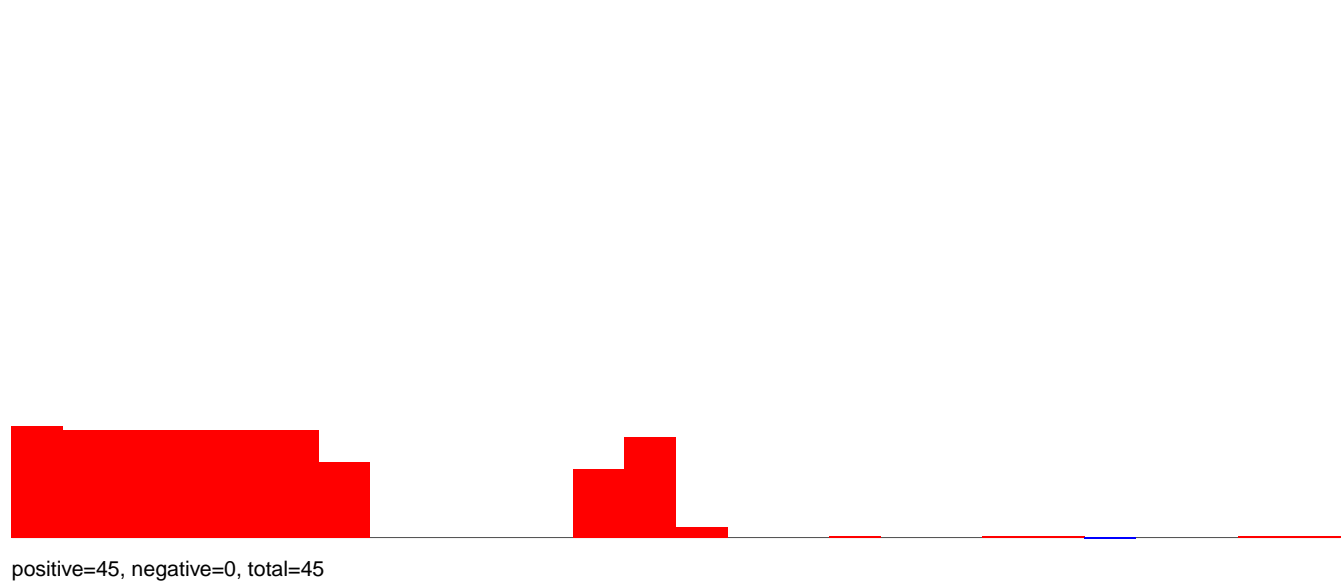
CuQuin_Mcarc_NL.rep



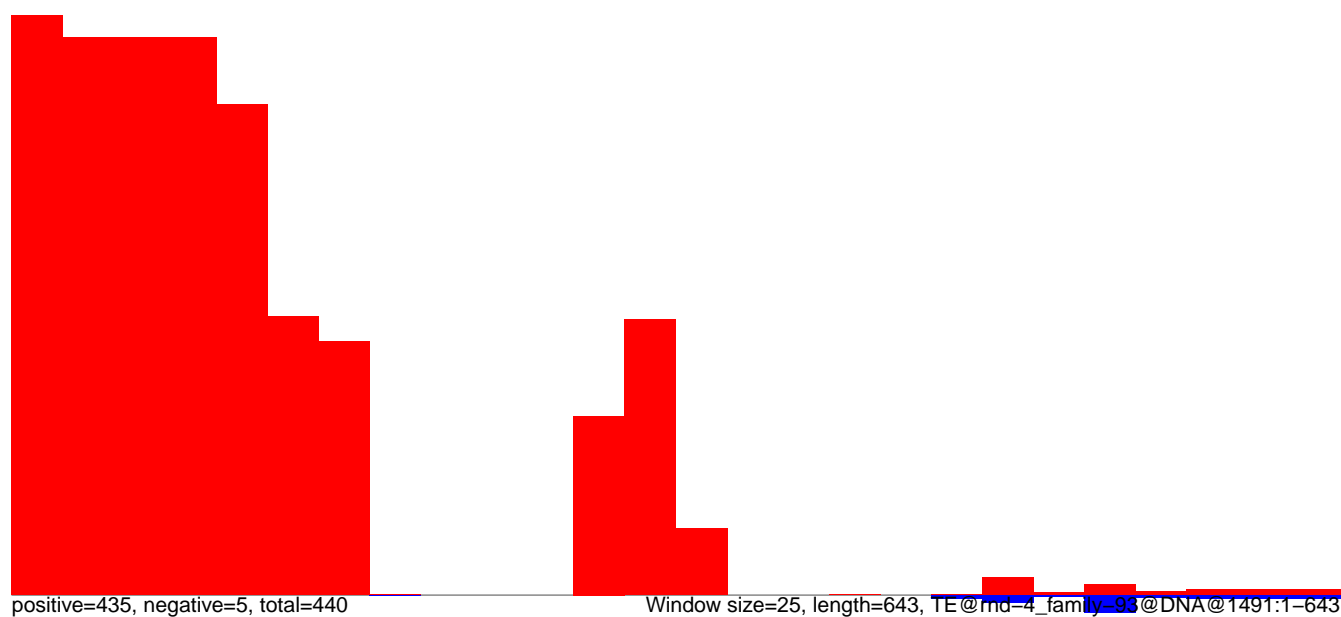
CuQuin_Mcarc_NL.18_23.rep



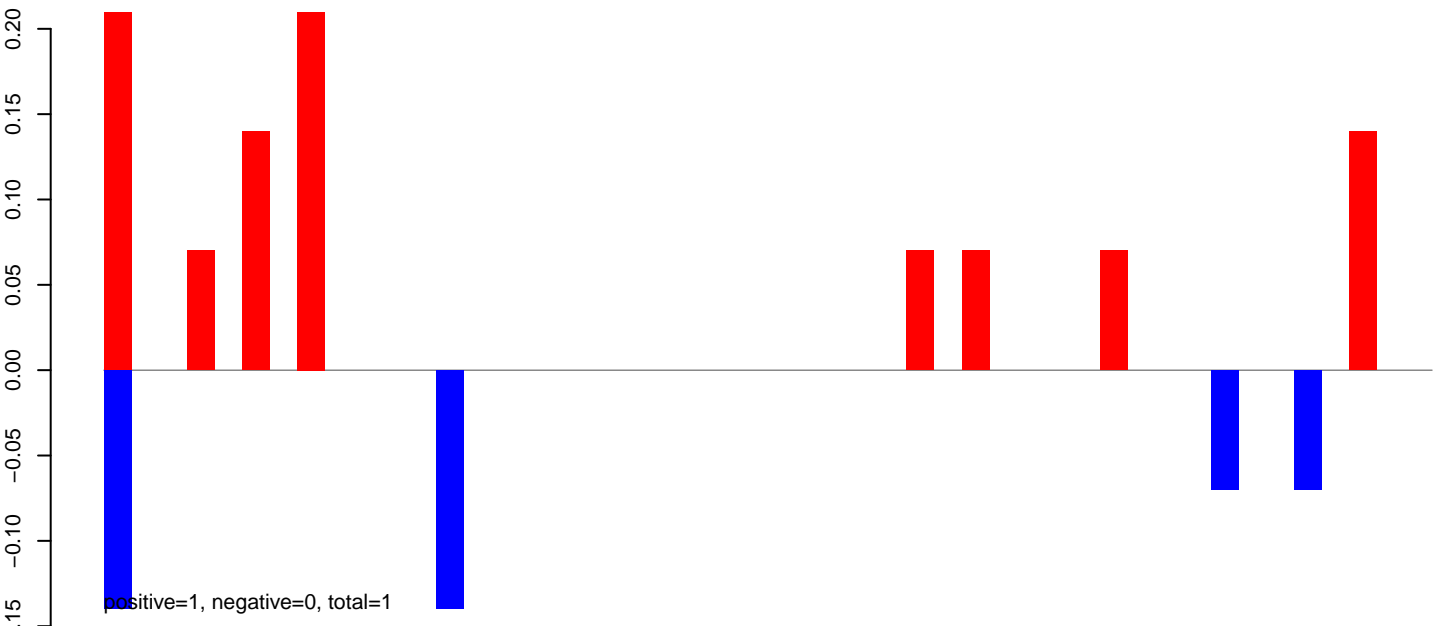
CuQuin_Mcarc_NL.24_35.rep



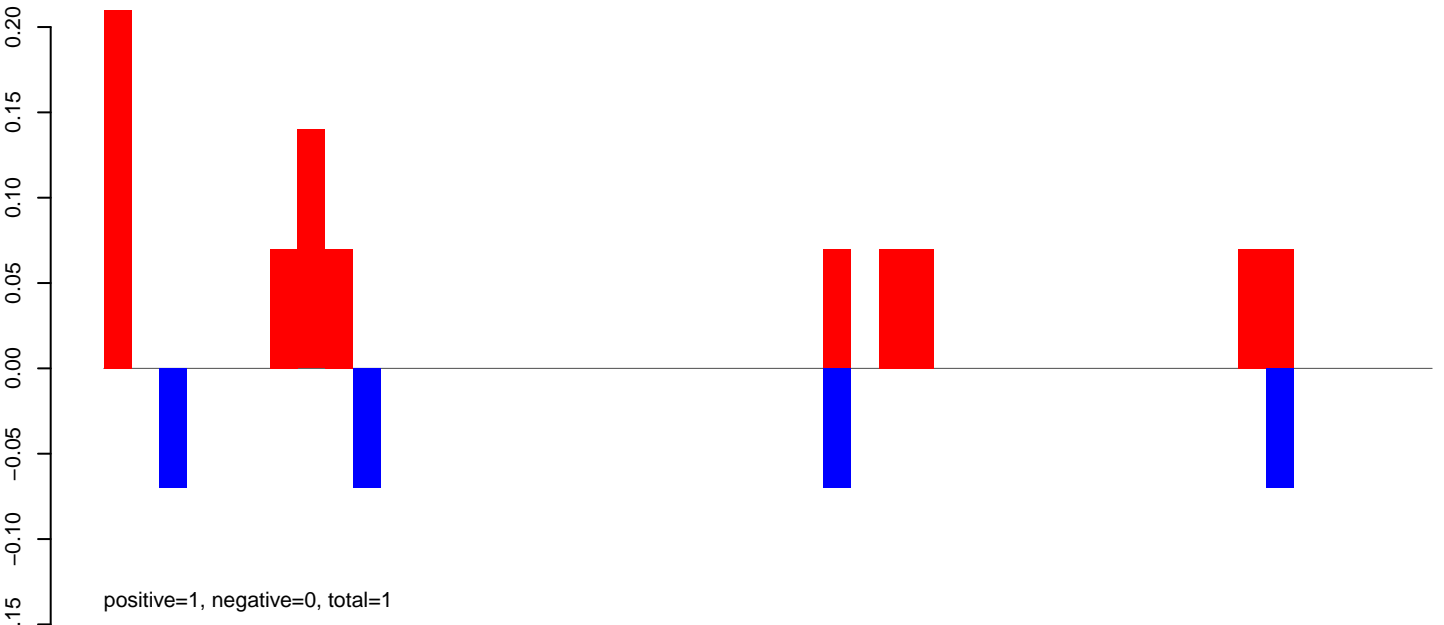
CuQuin_Mcarc_NL.rep



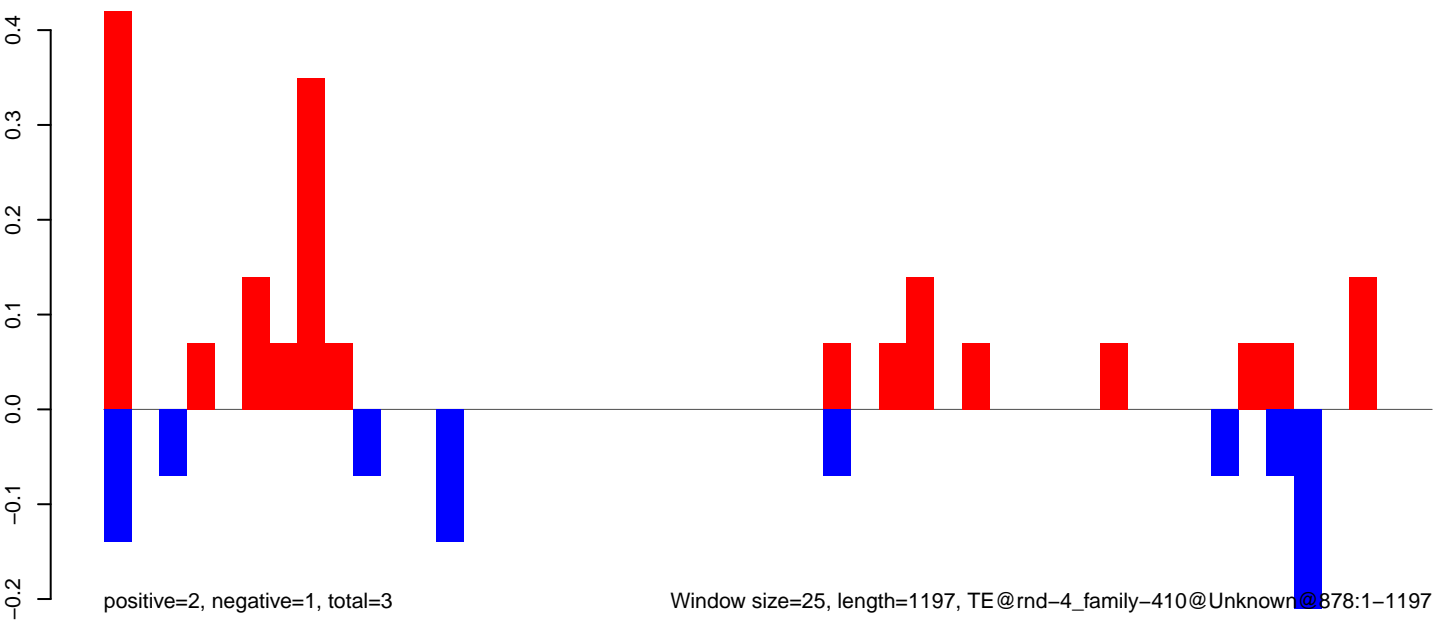
CuQuin_Mcarc_NL.18_23.rep



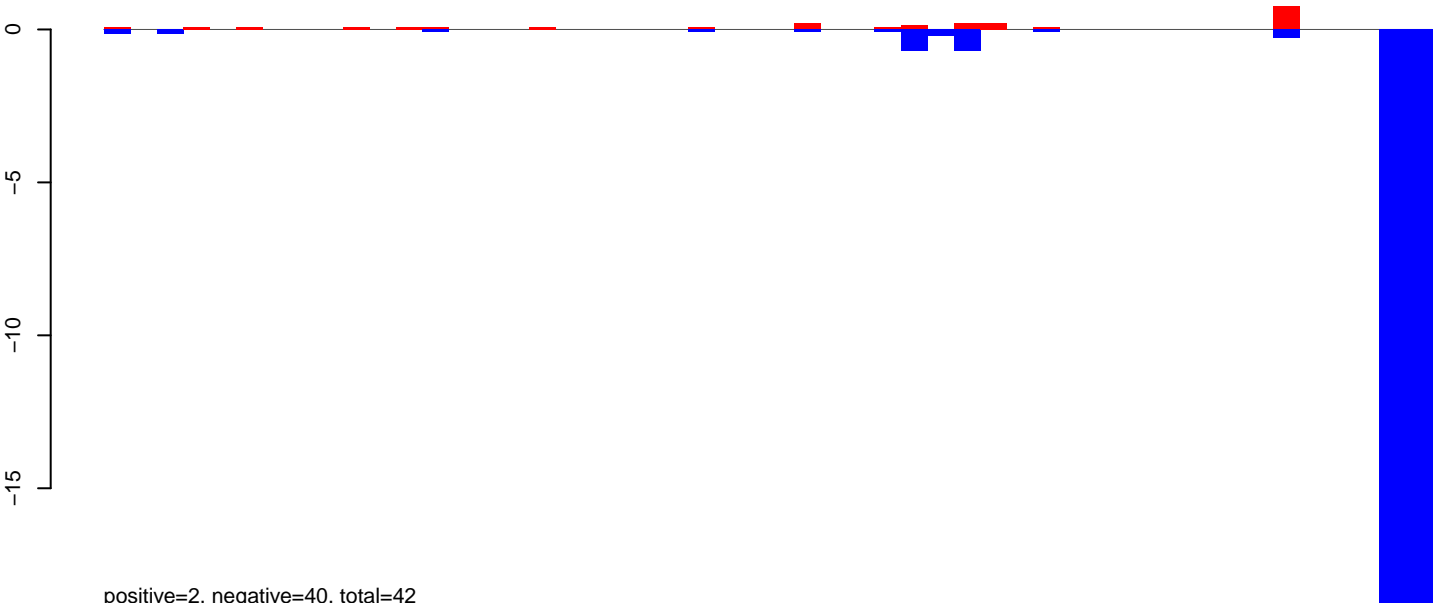
CuQuin_Mcarc_NL.24_35.rep



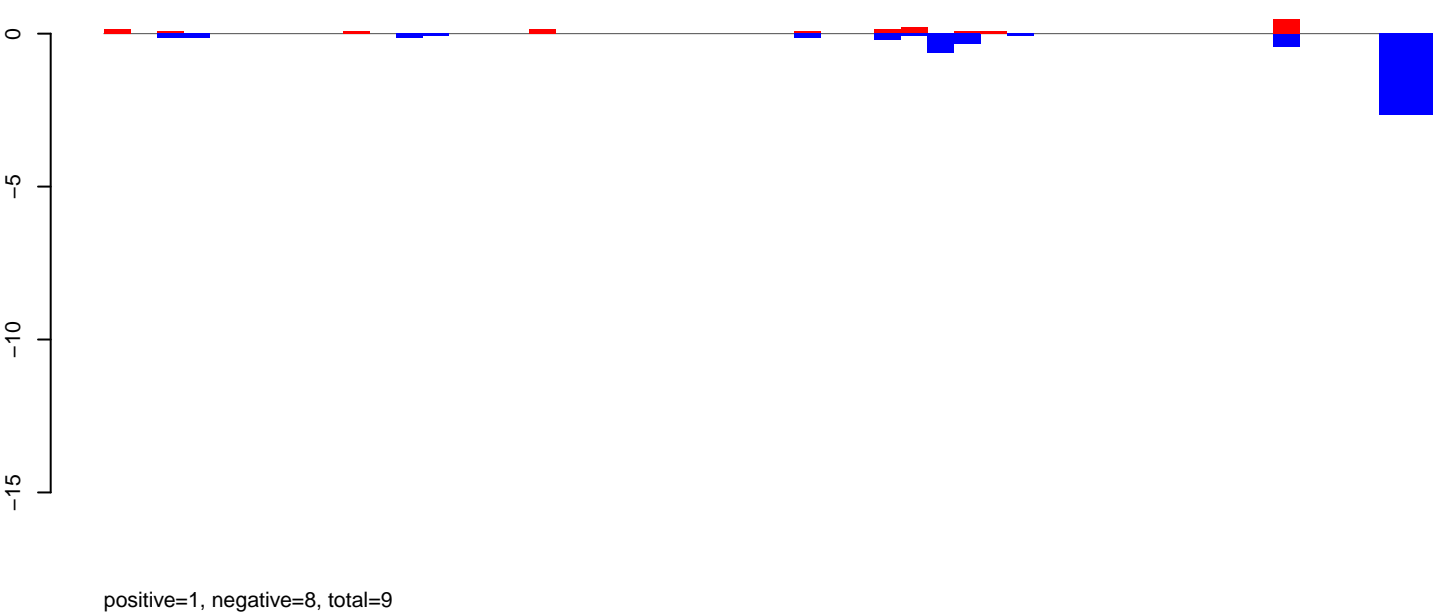
CuQuin_Mcarc_NL.rep



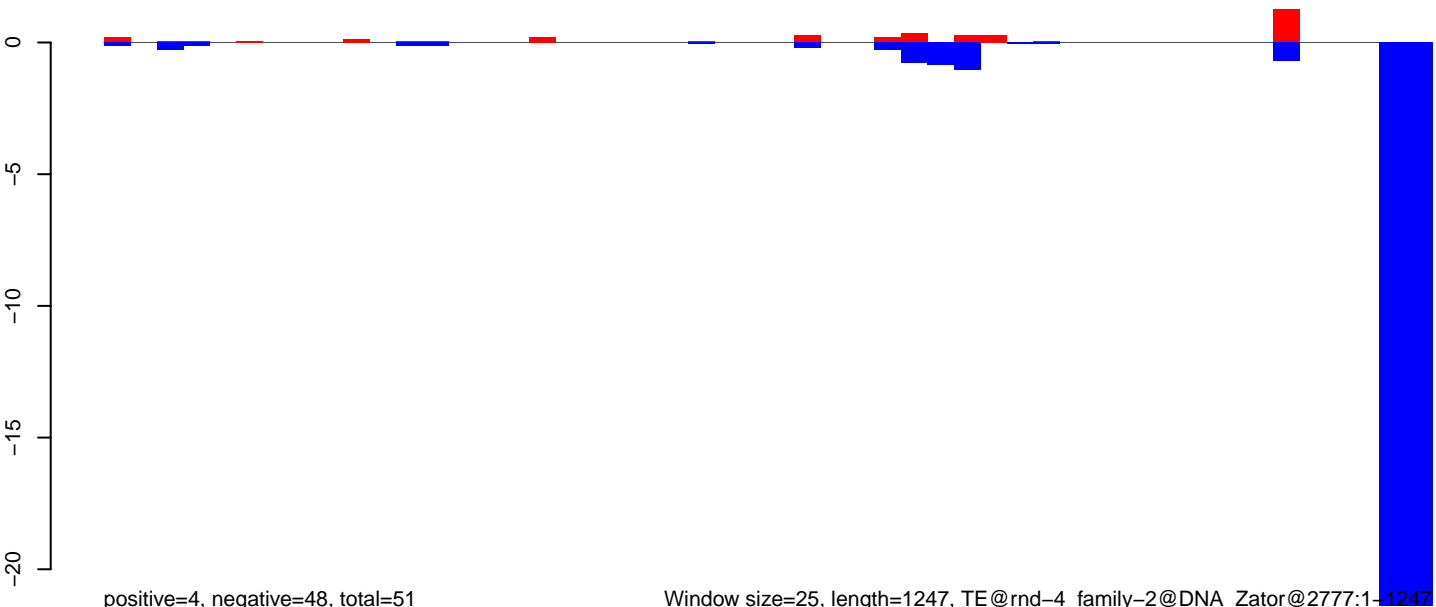
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

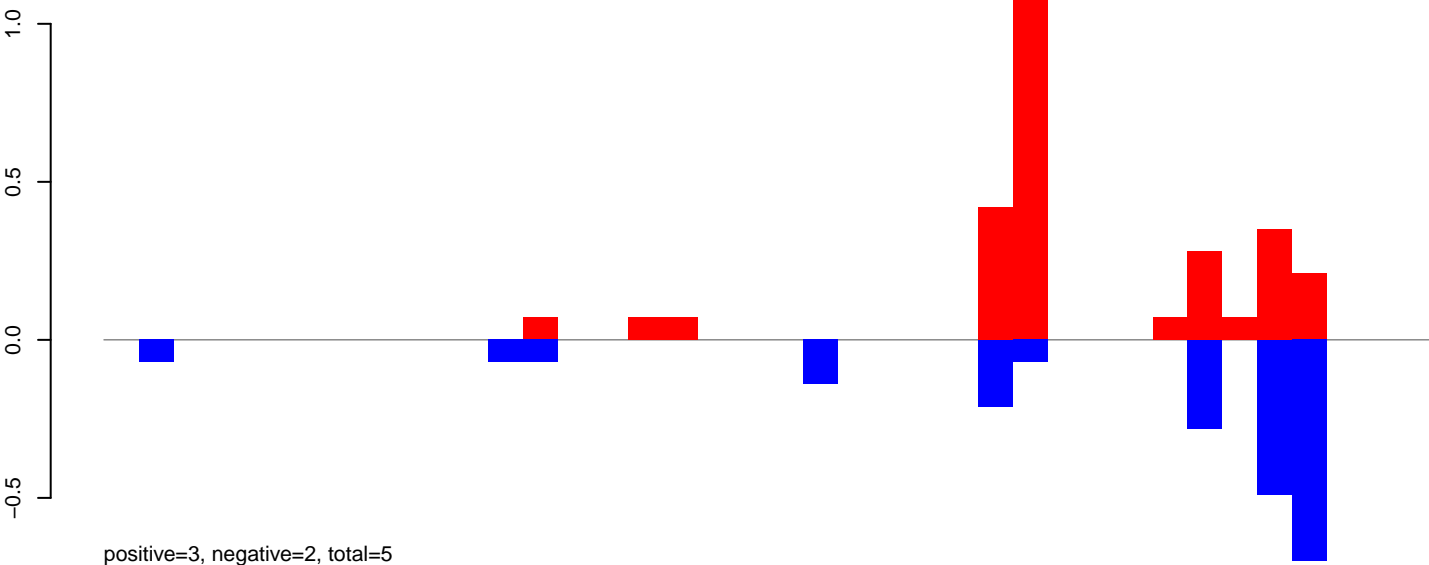


CuQuin_Mcarc_NL.rep

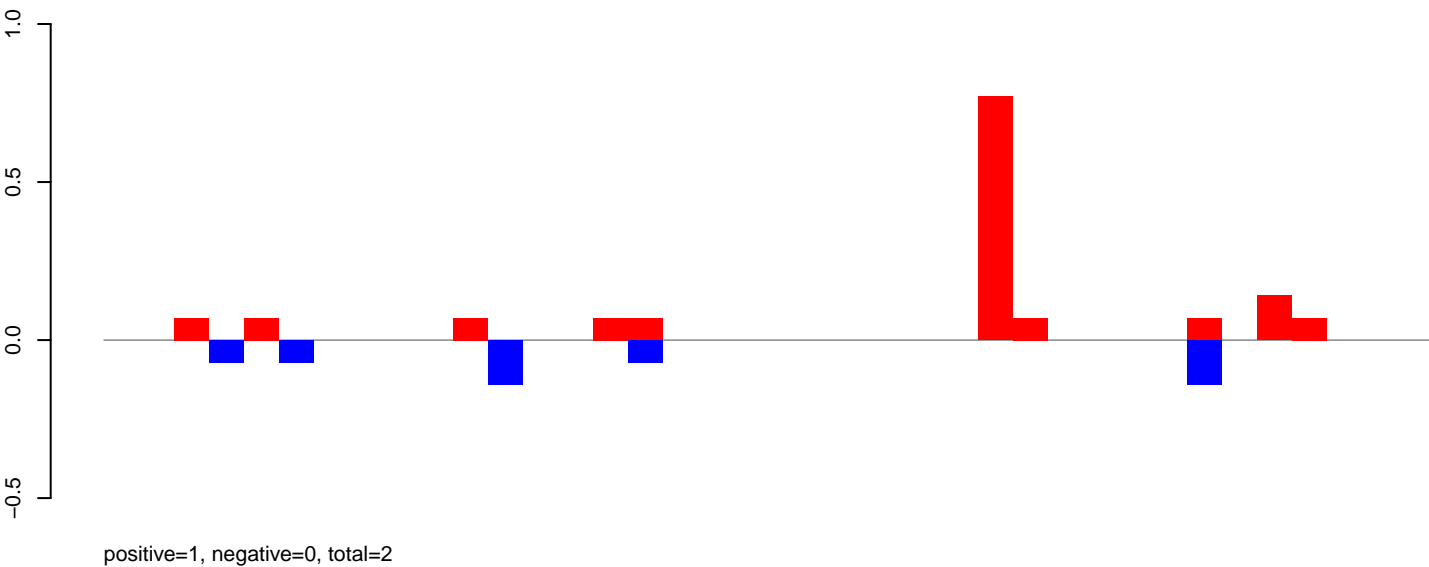


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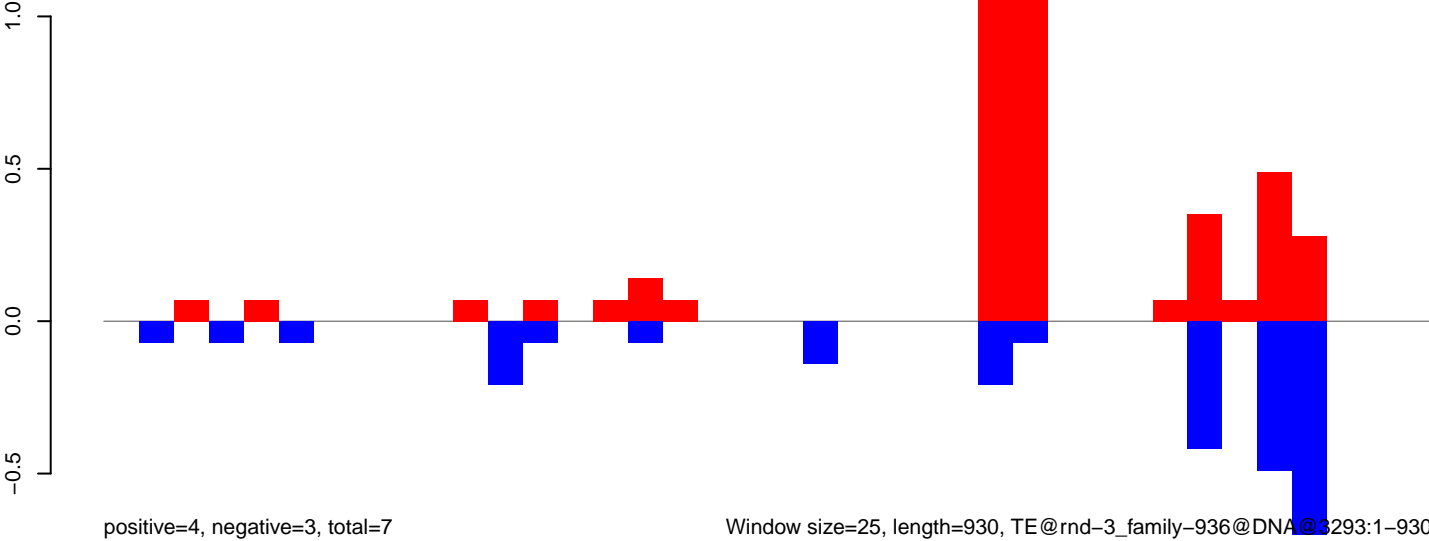
CuQuin_Mcarc_NL.18_23.rep



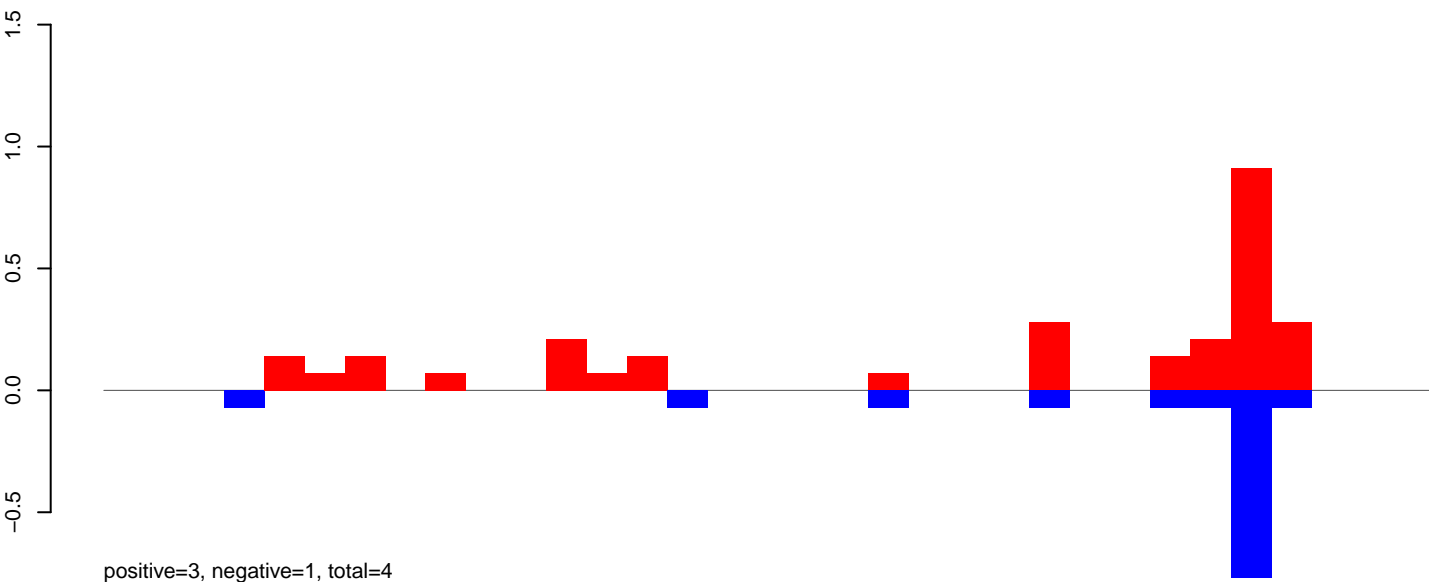
CuQuin_Mcarc_NL.24_35.rep



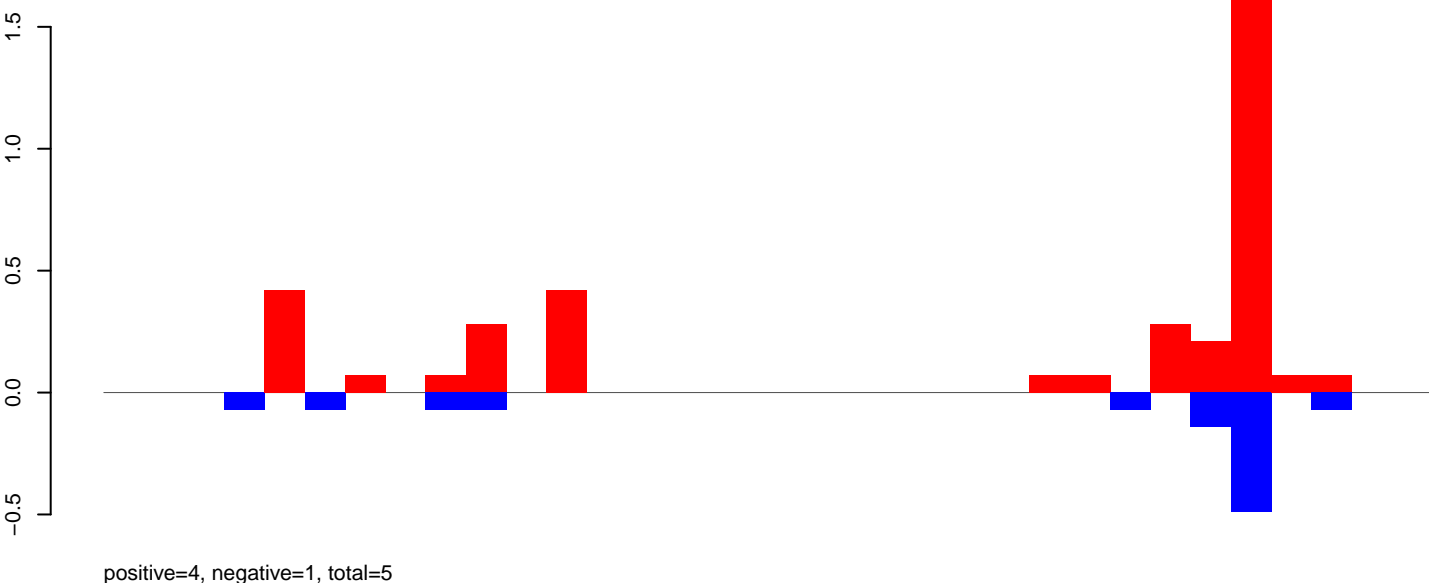
CuQuin_Mcarc_NL.rep



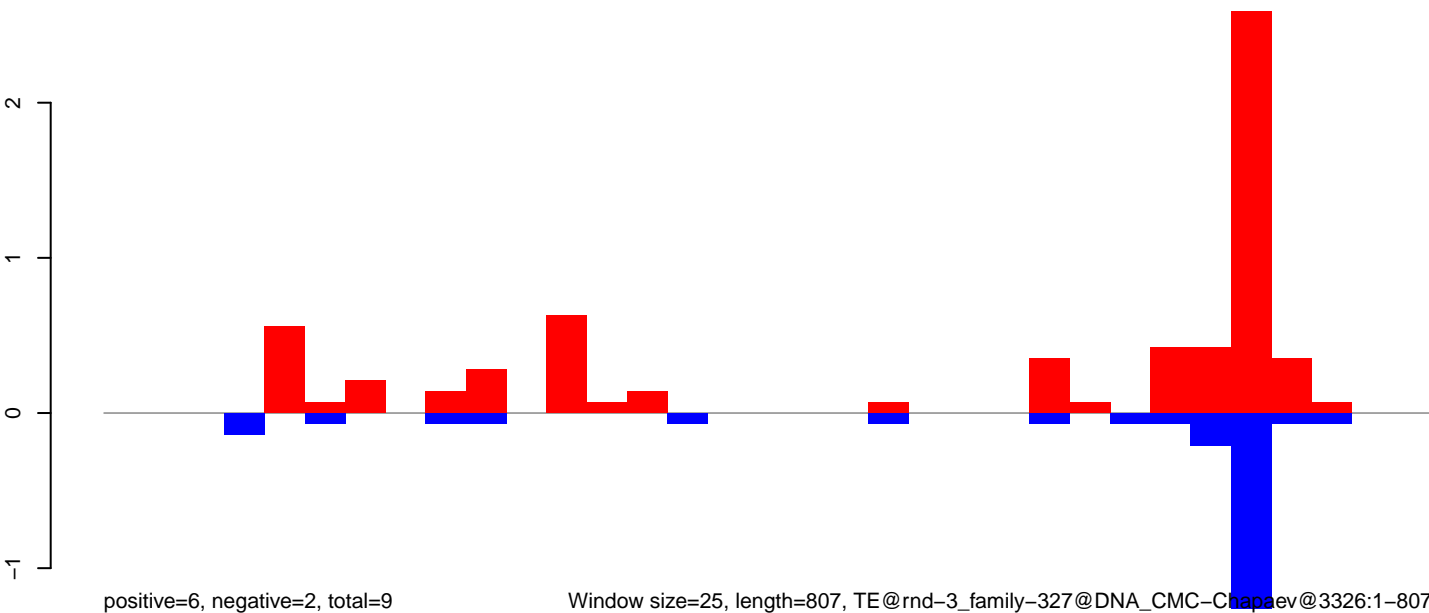
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CuQuin_Mcarc_NL.24_35.rep

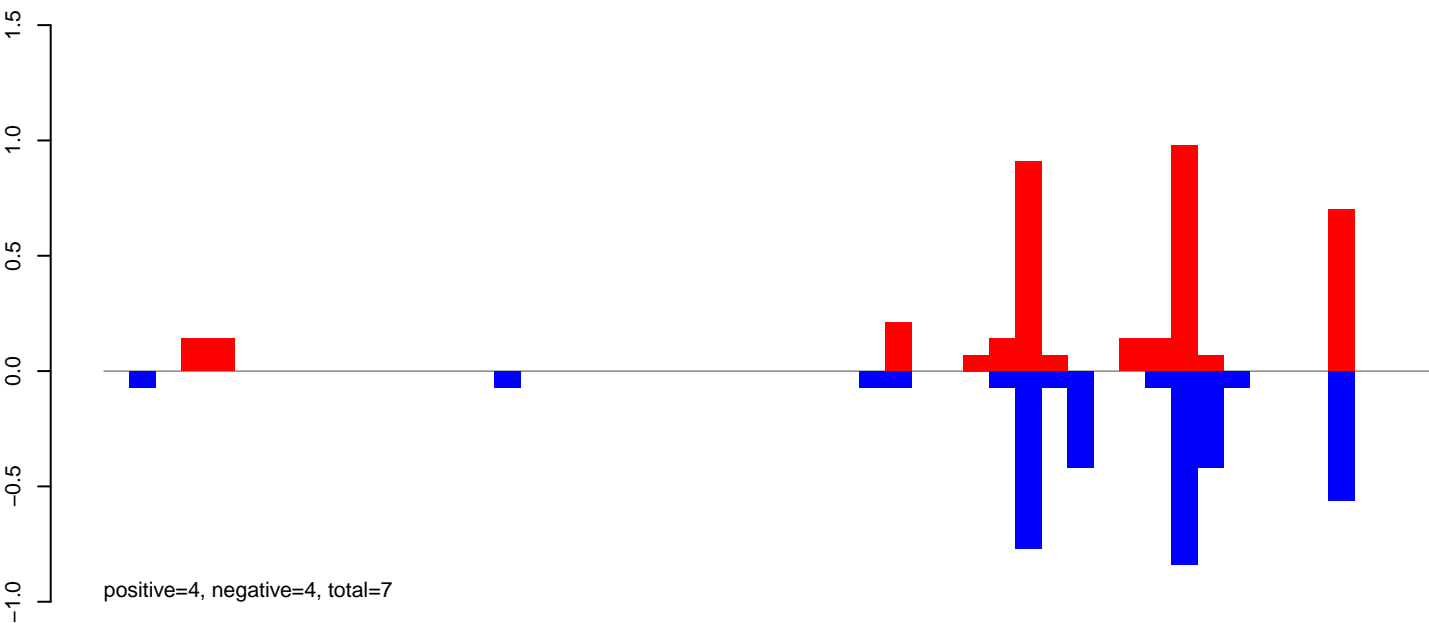


CuQuin_Mcarc_NL.rep

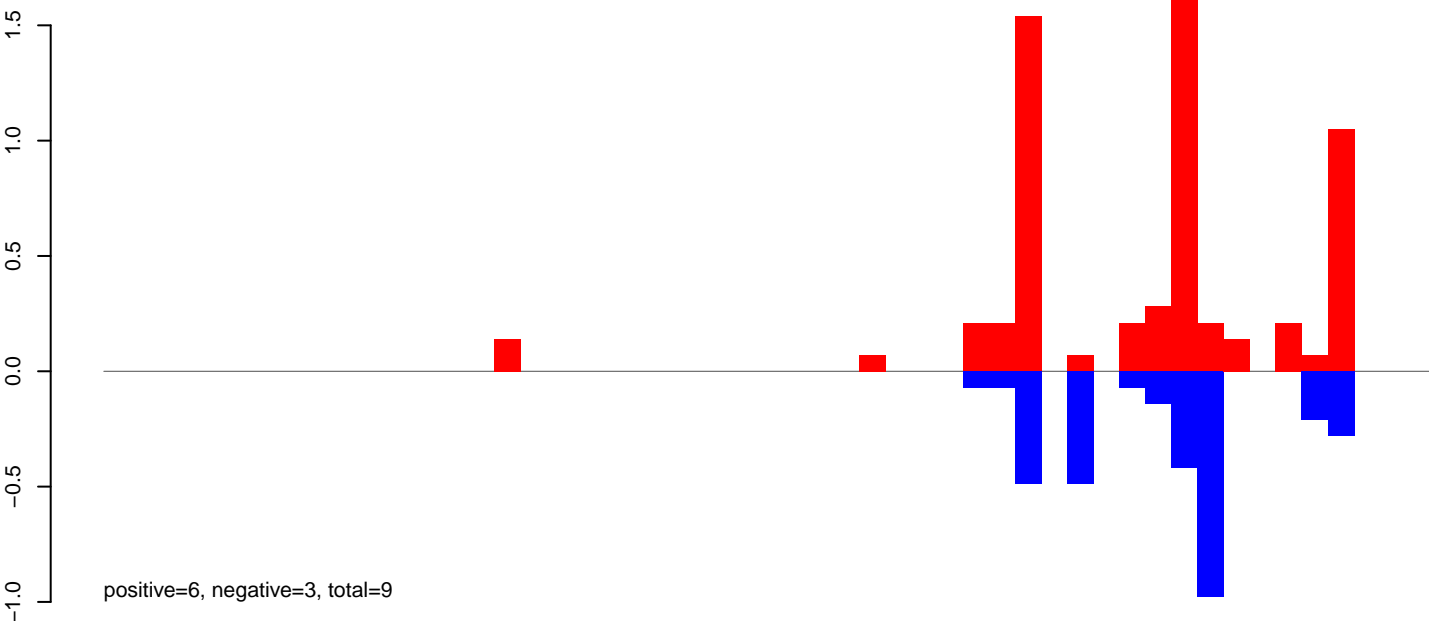


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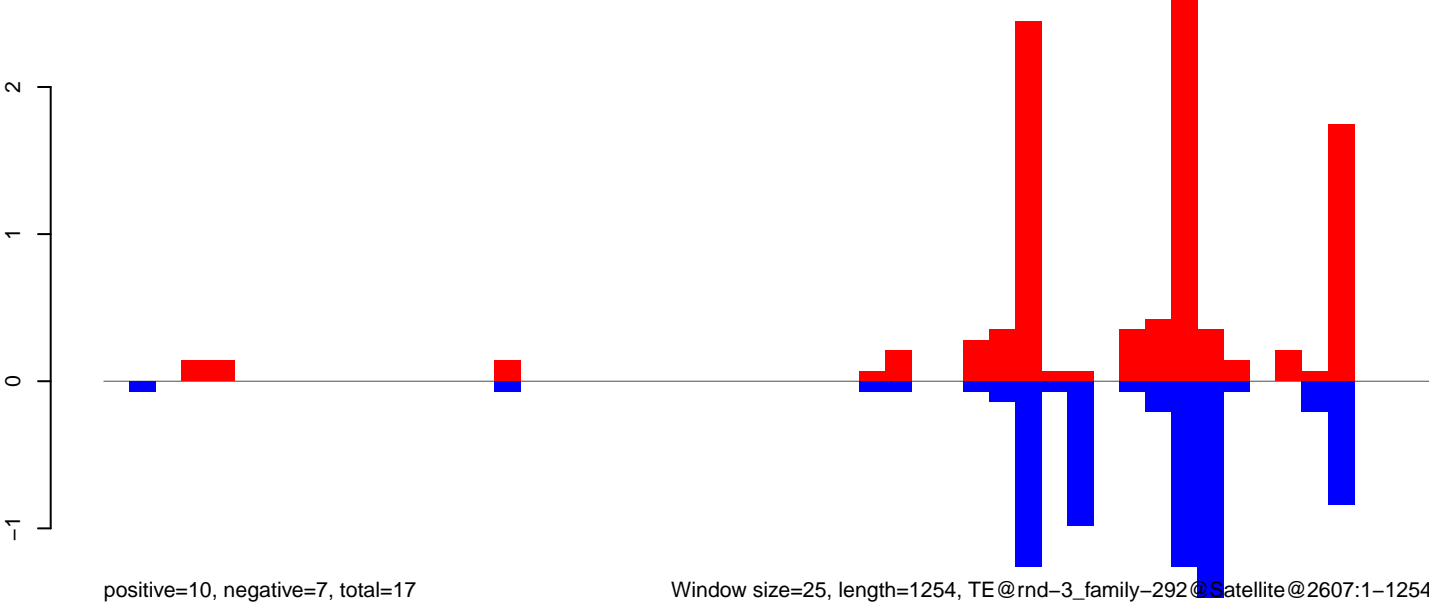
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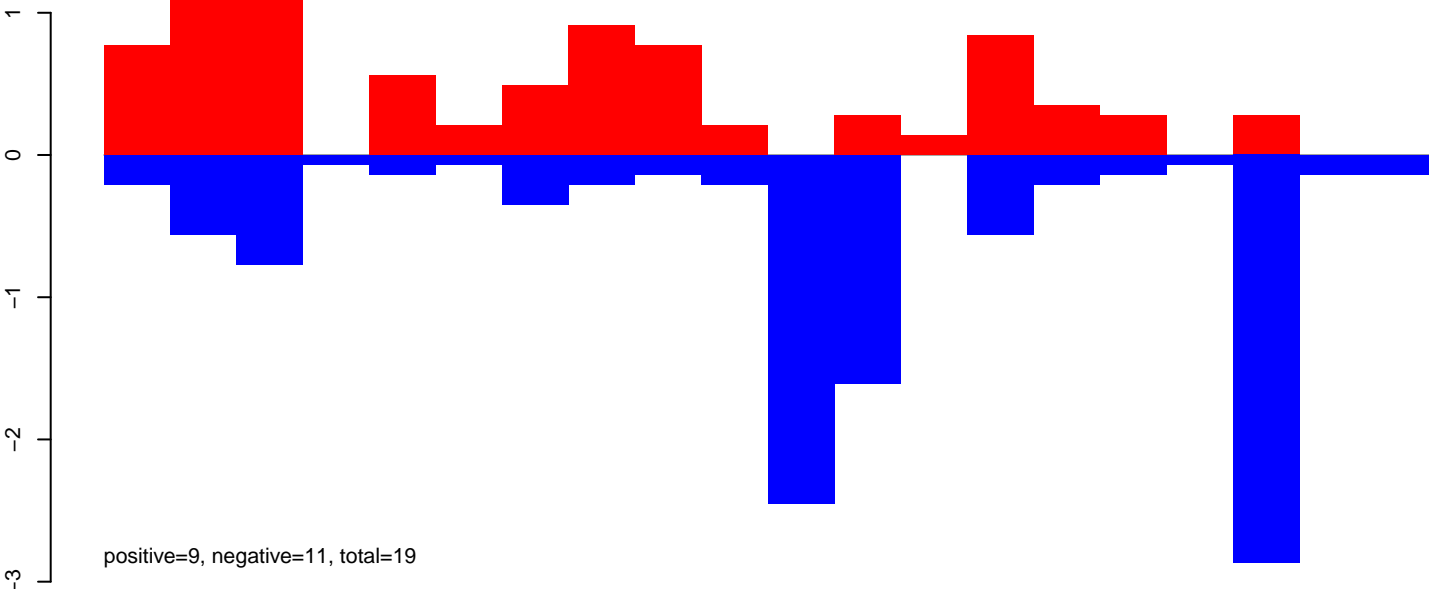
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



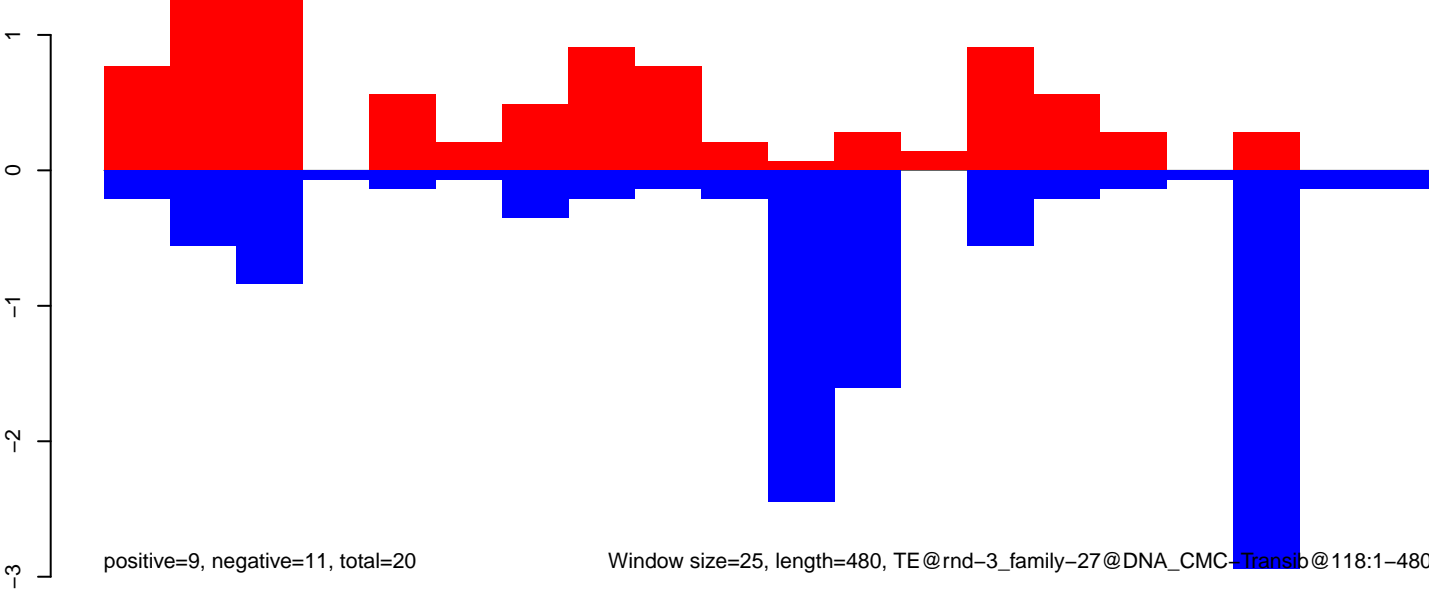
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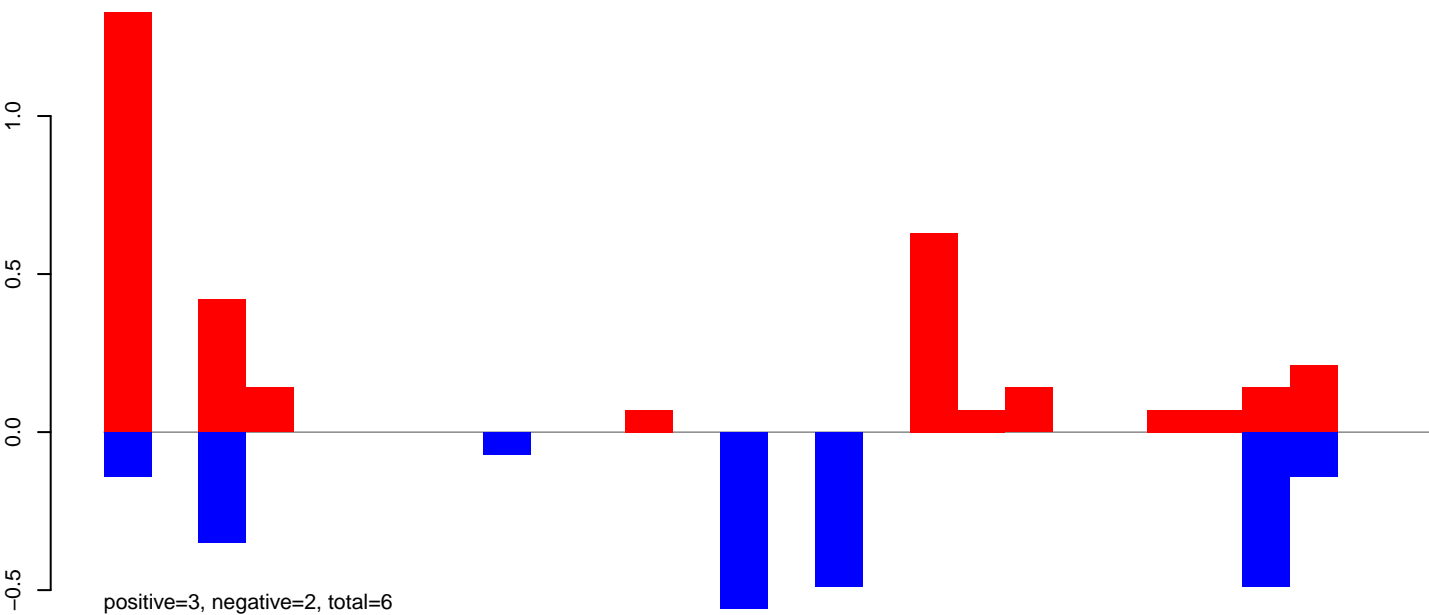
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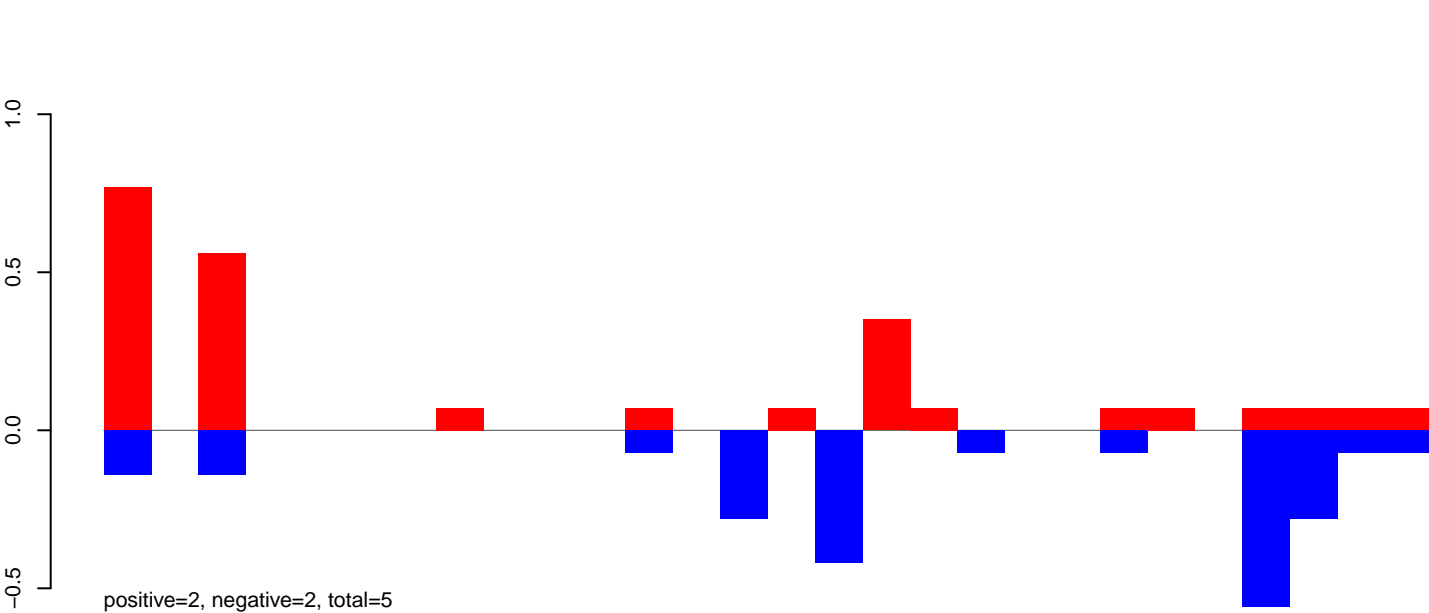
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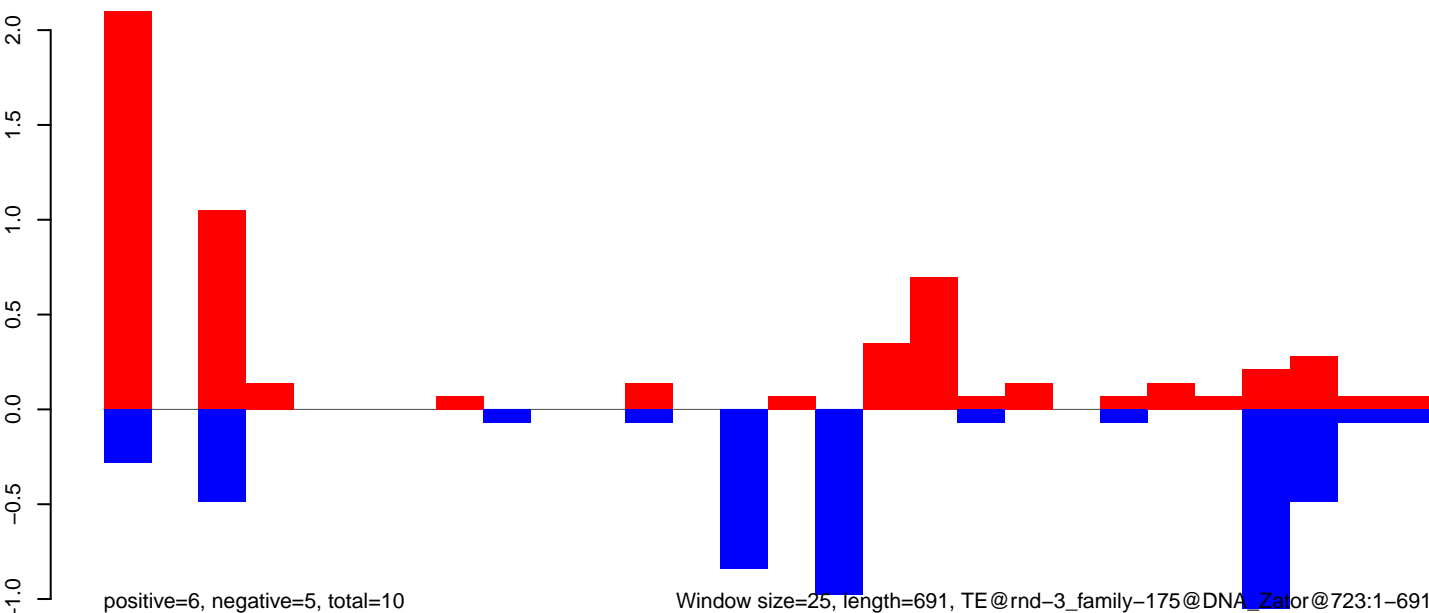
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CuQuin_Mcarc_NL.24_35.rep

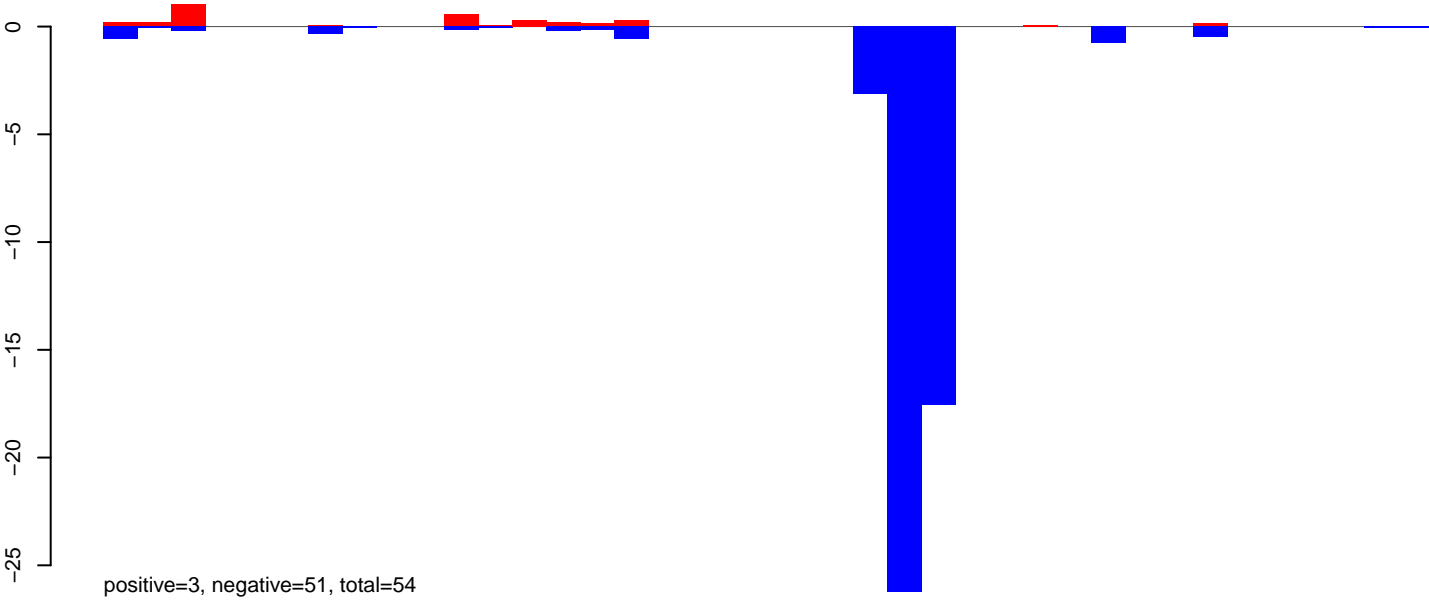


CuQuin_Mcarc_NL.rep

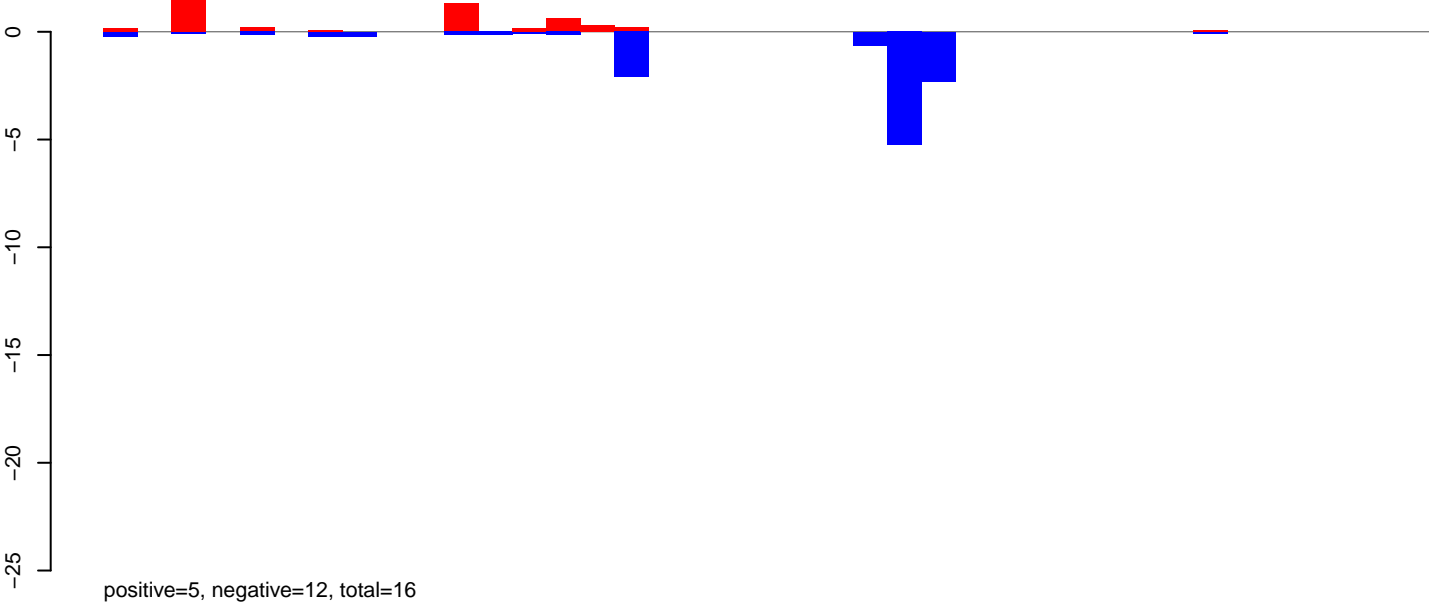


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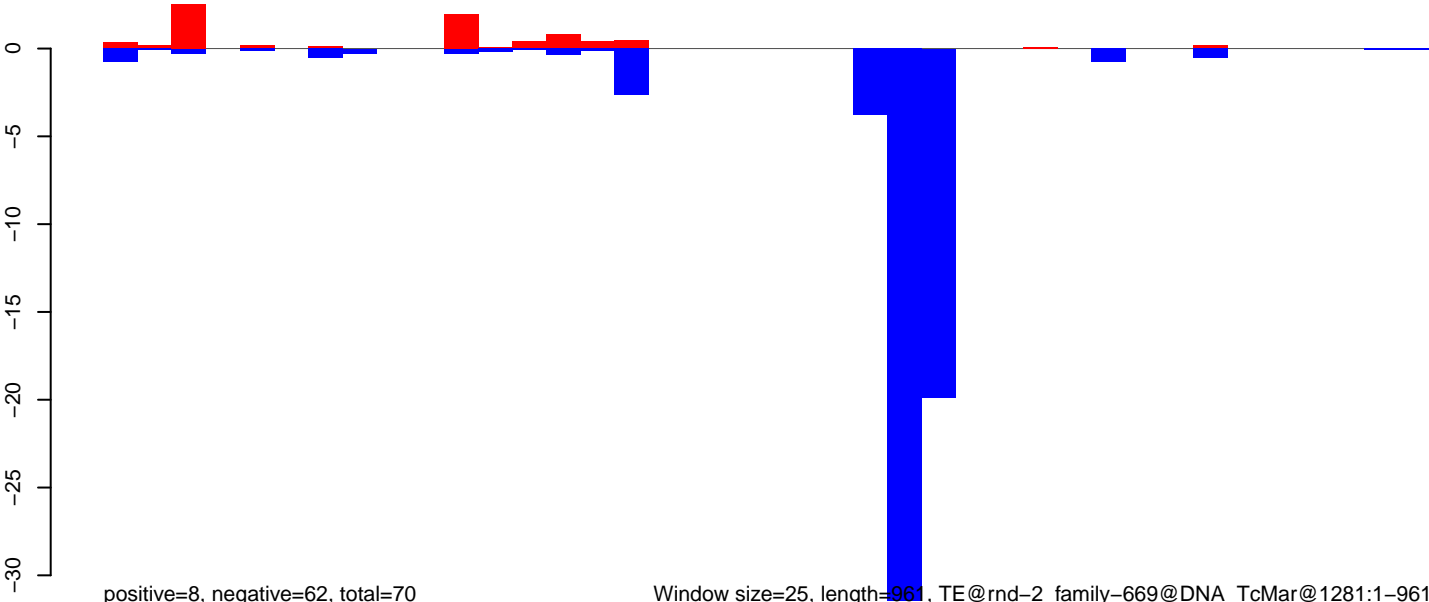
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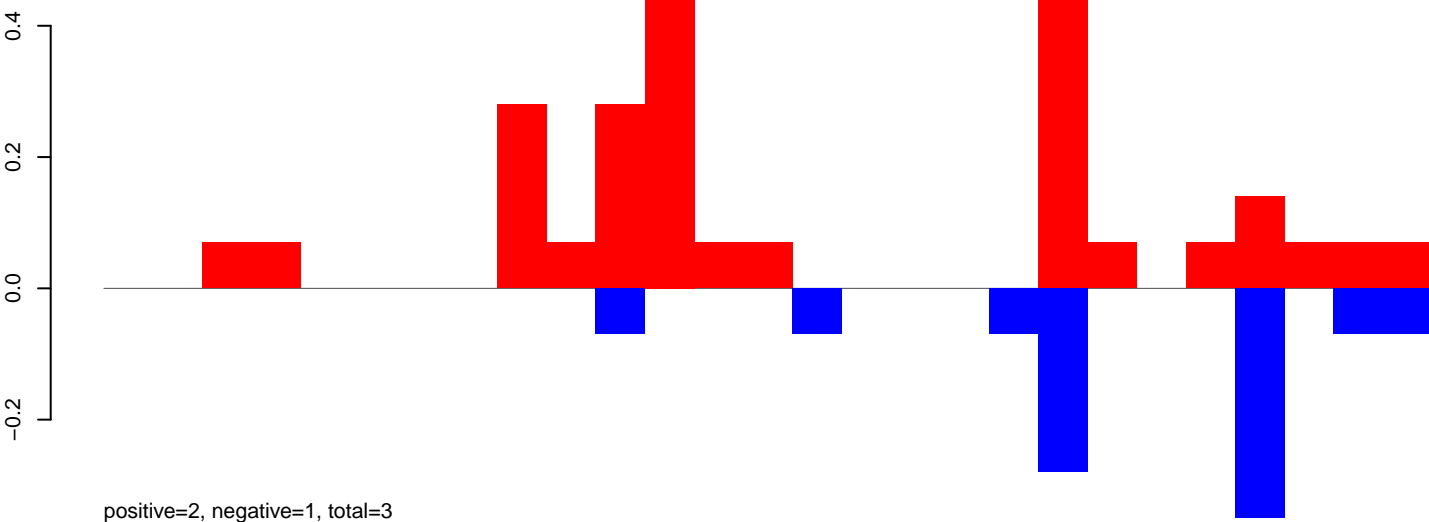
CuQuin_Mcarc_NL.24_35.rep



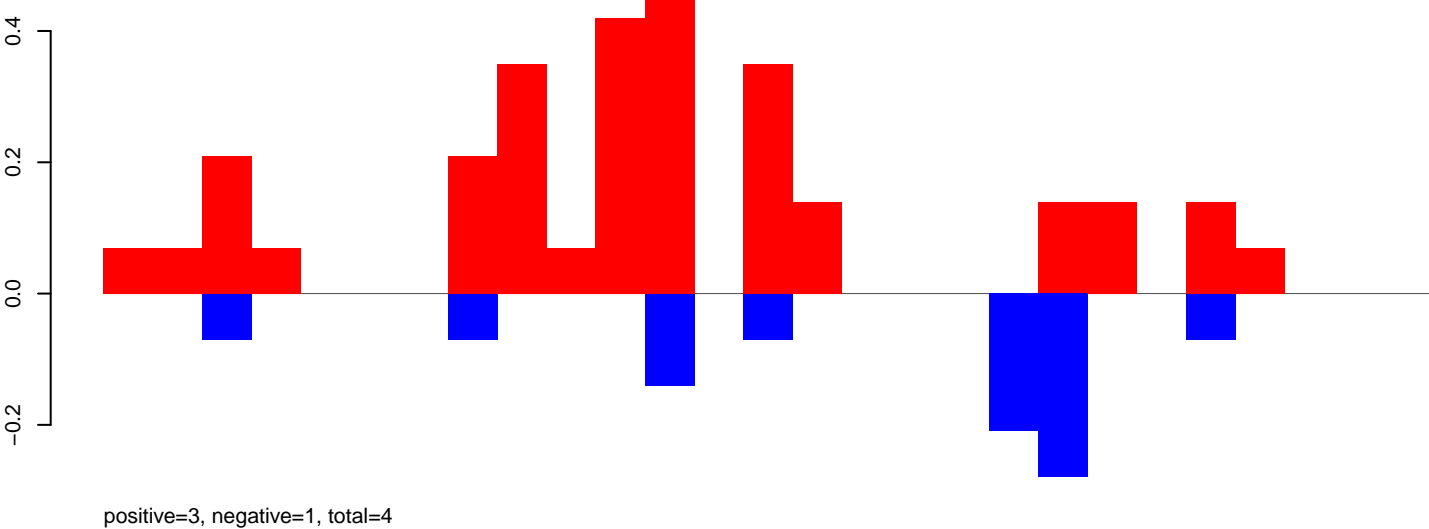
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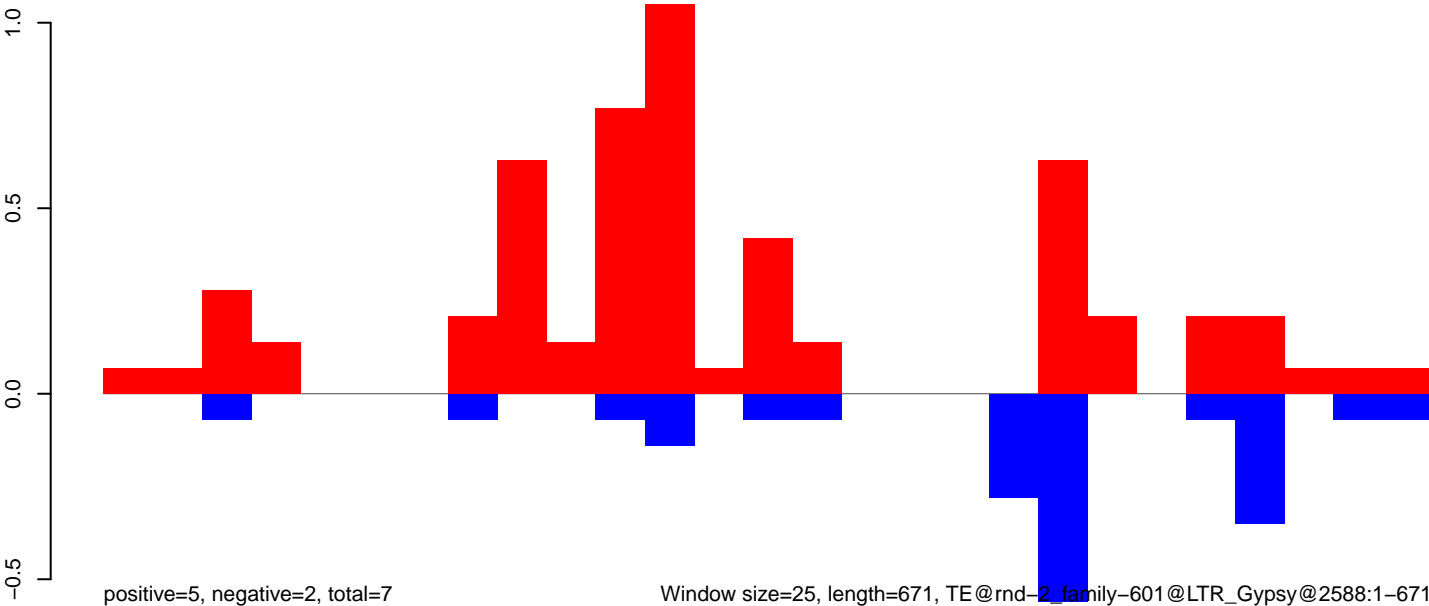
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

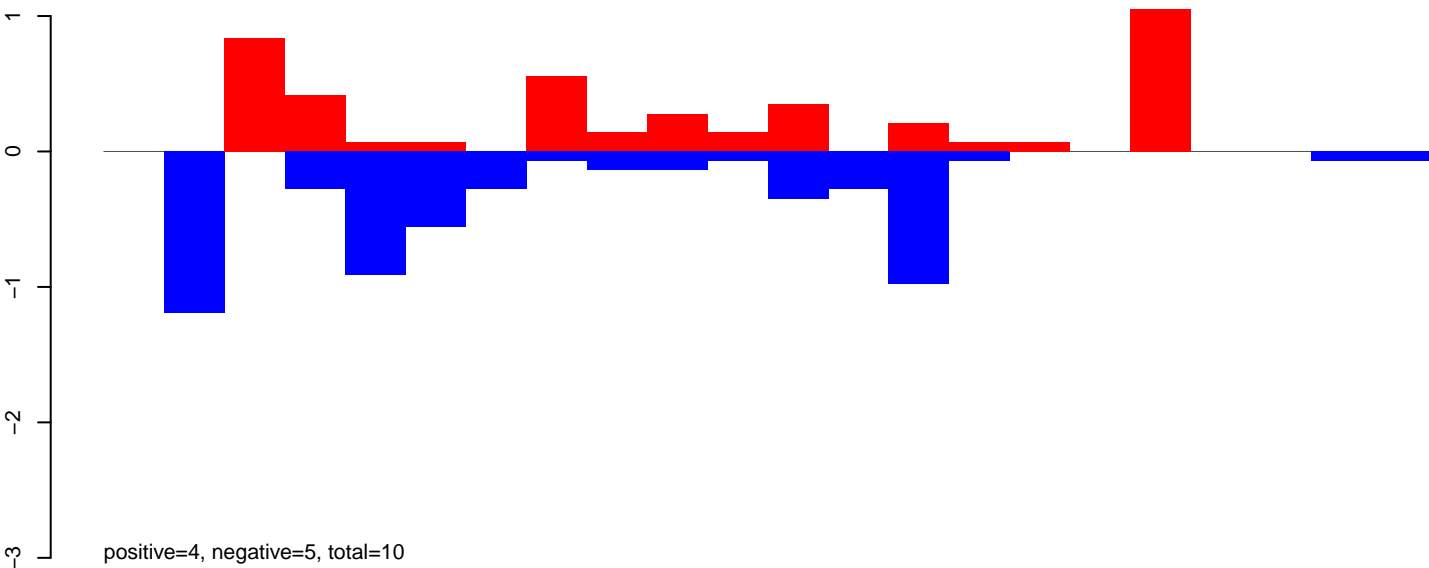


CuQuin_Mcarc_NL.rep

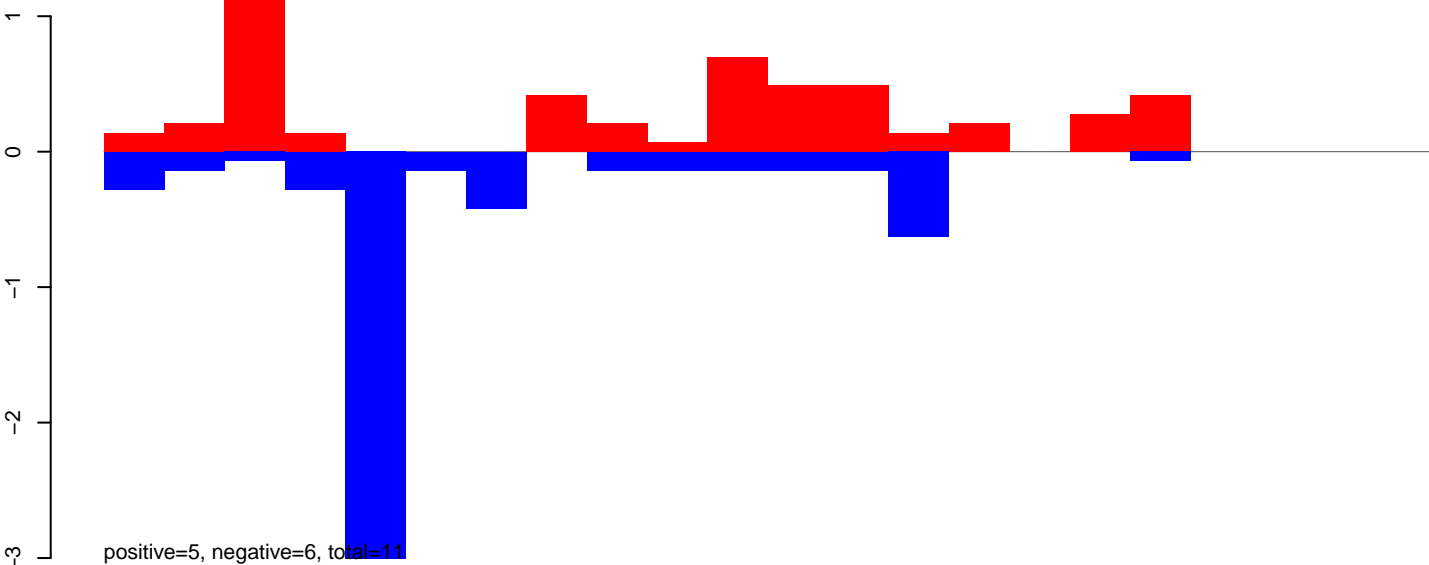


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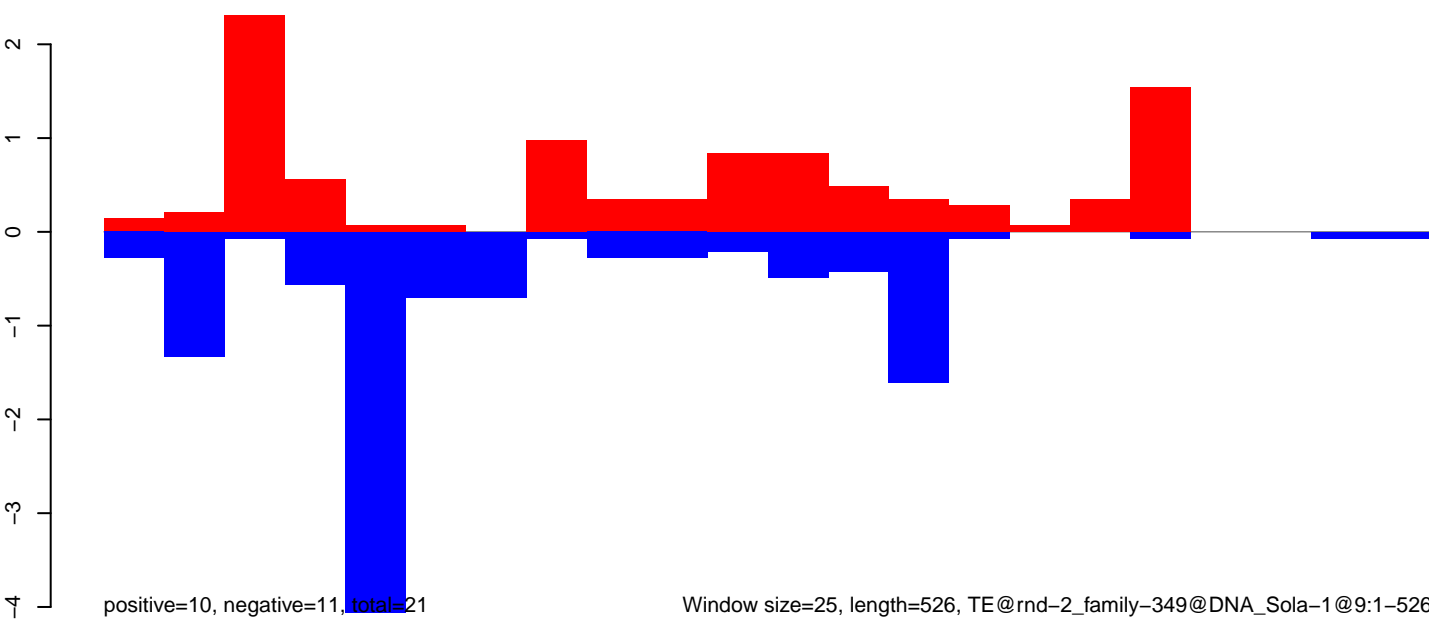
CuQuin_Mcarc_NL.18_23.rep



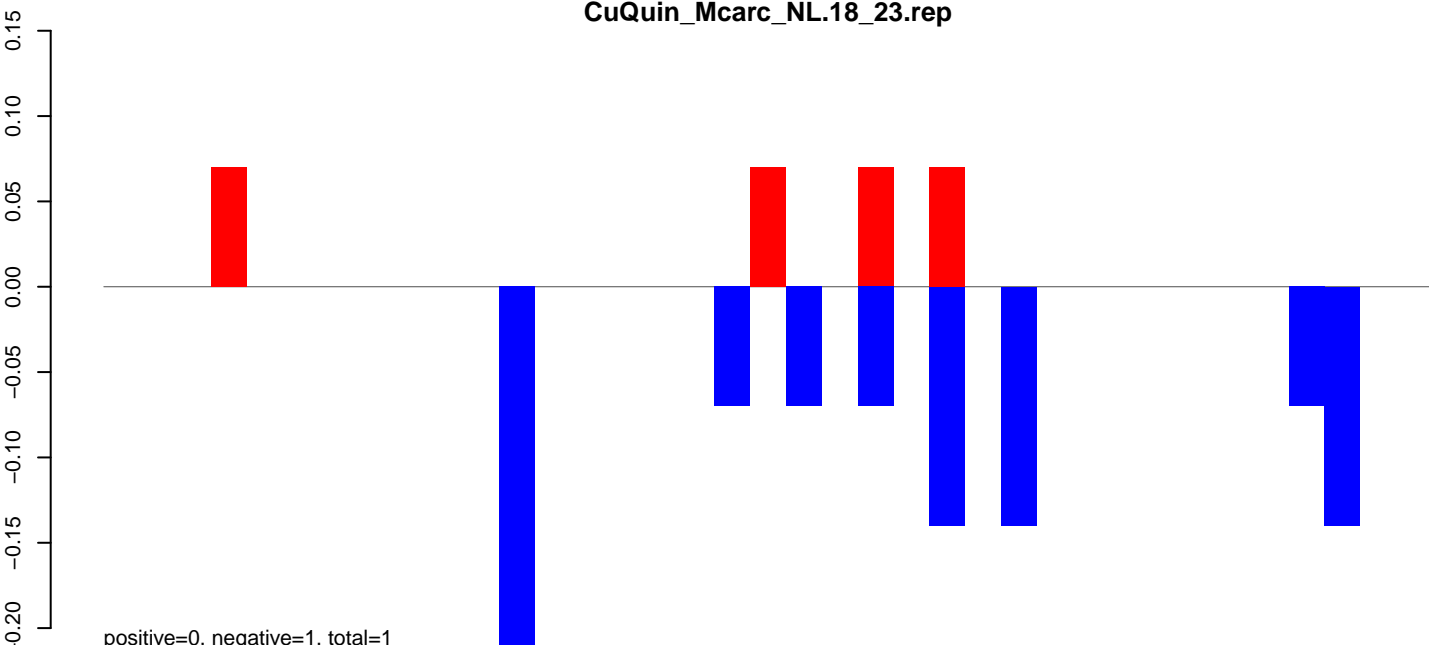
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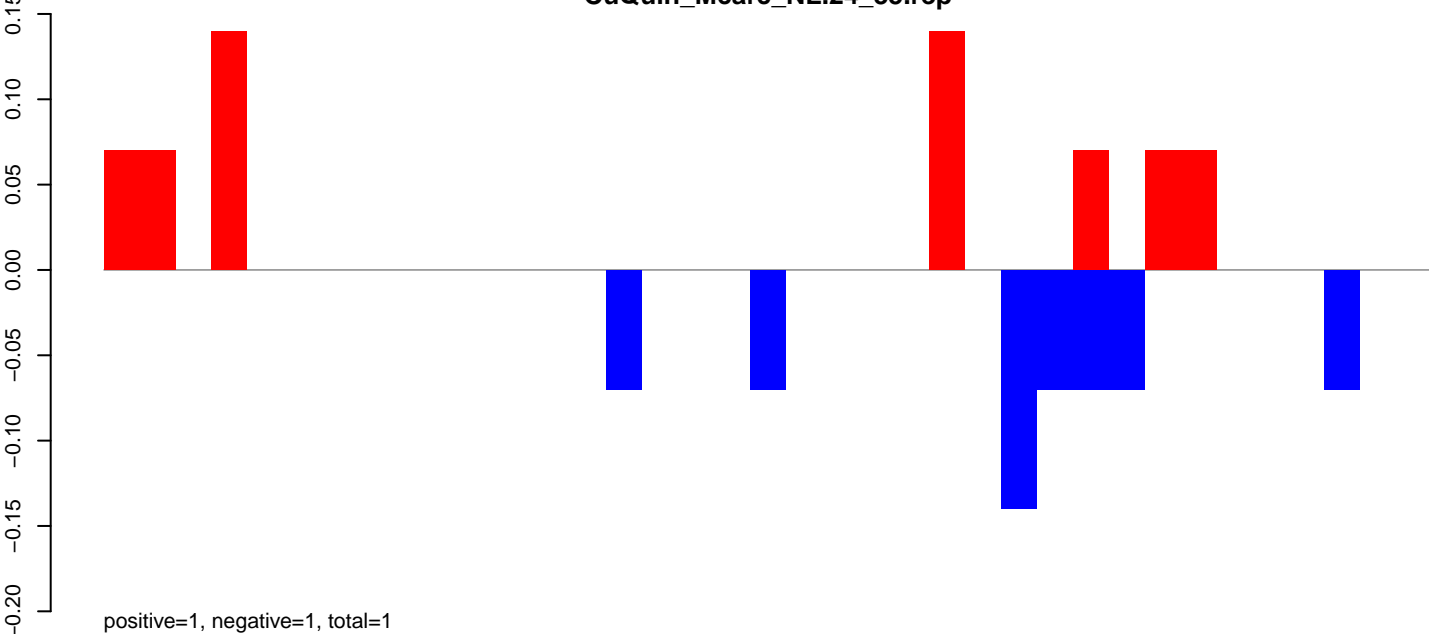
CuQuin_Mcarc_NL.rep



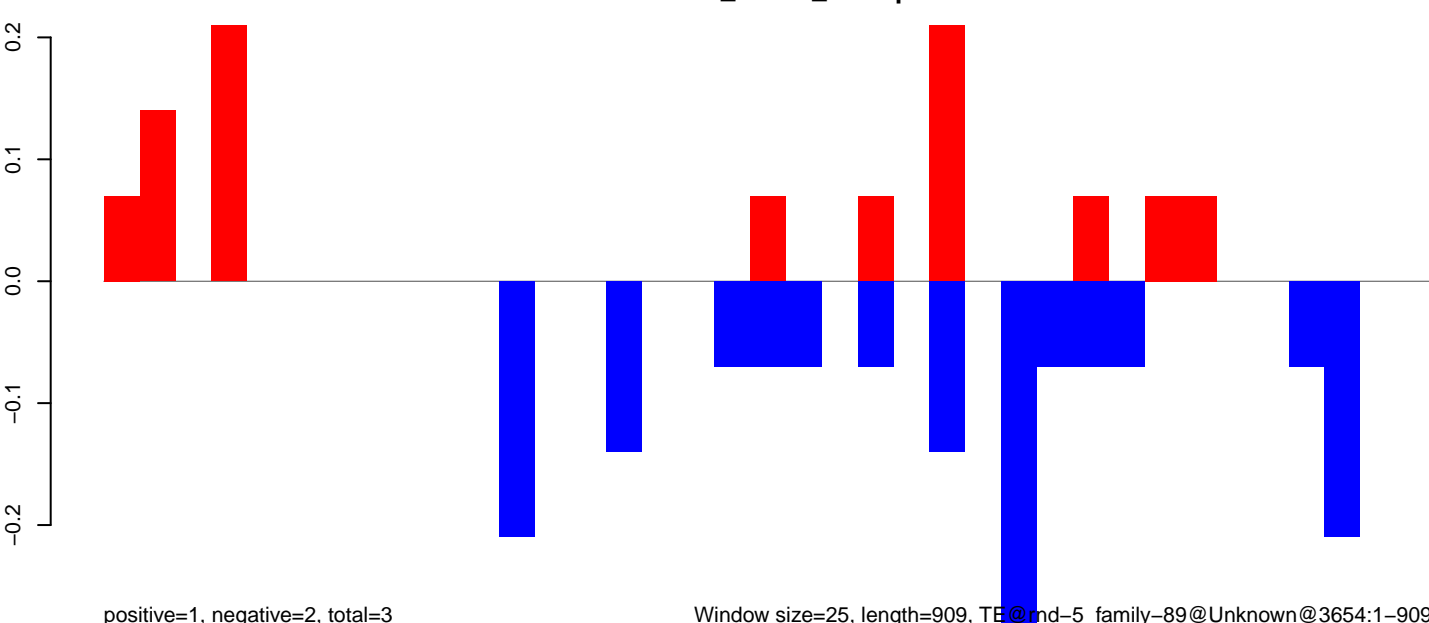
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CuQuin_Mcarc_NL.24_35.rep



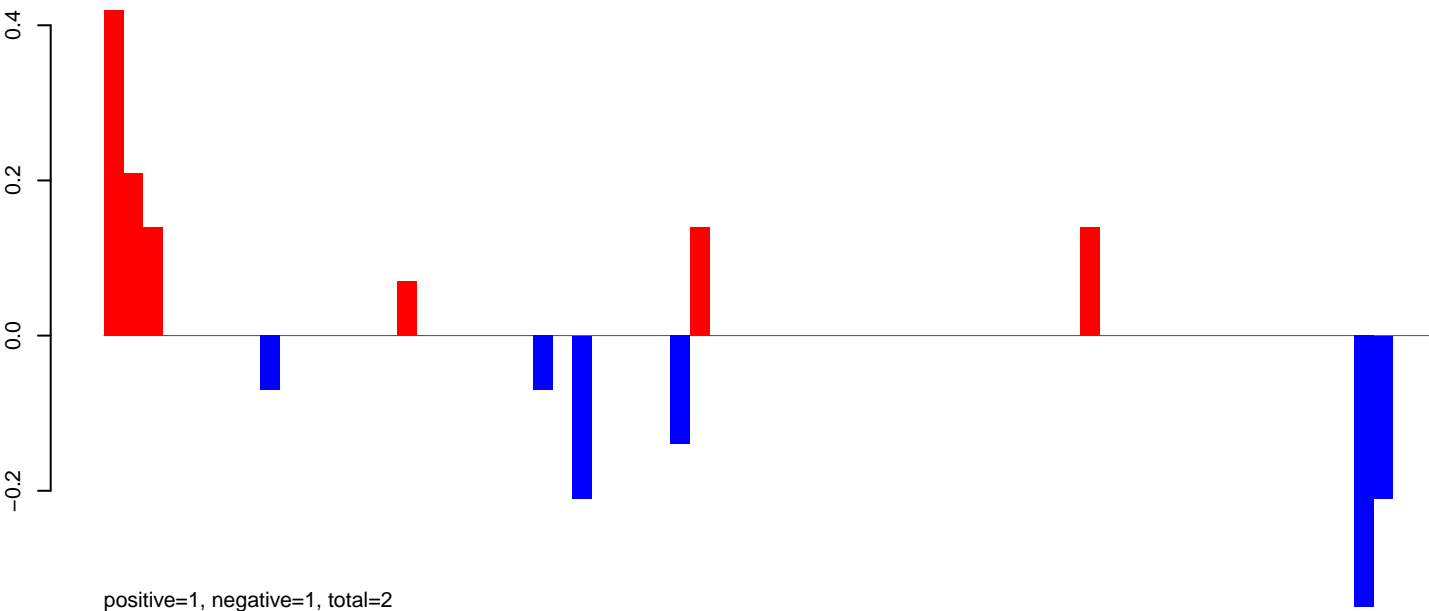
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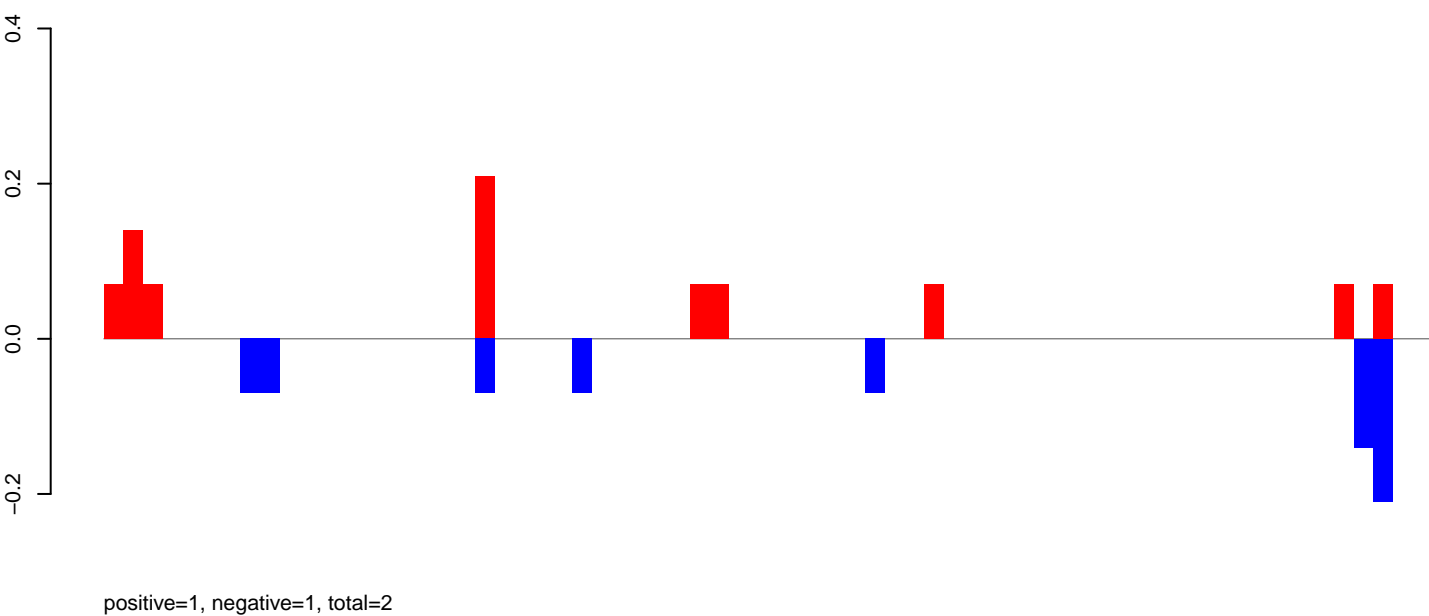
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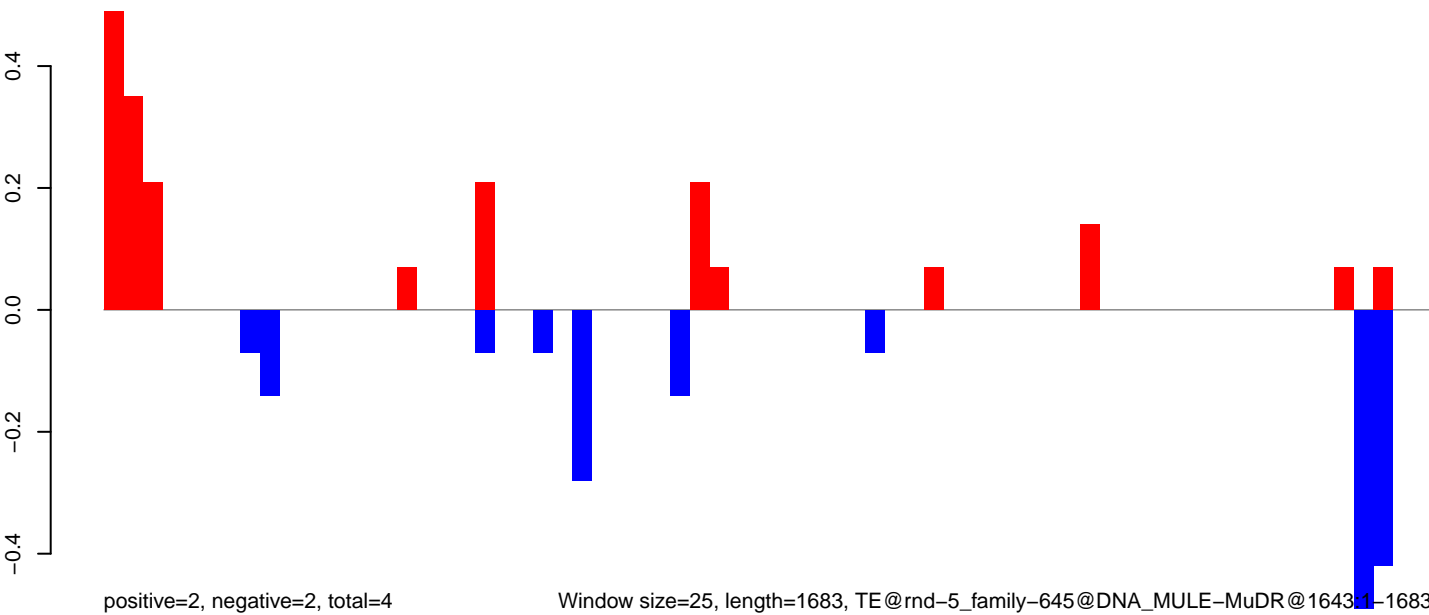
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CuQuin_Mcarc_NL.24_35.rep



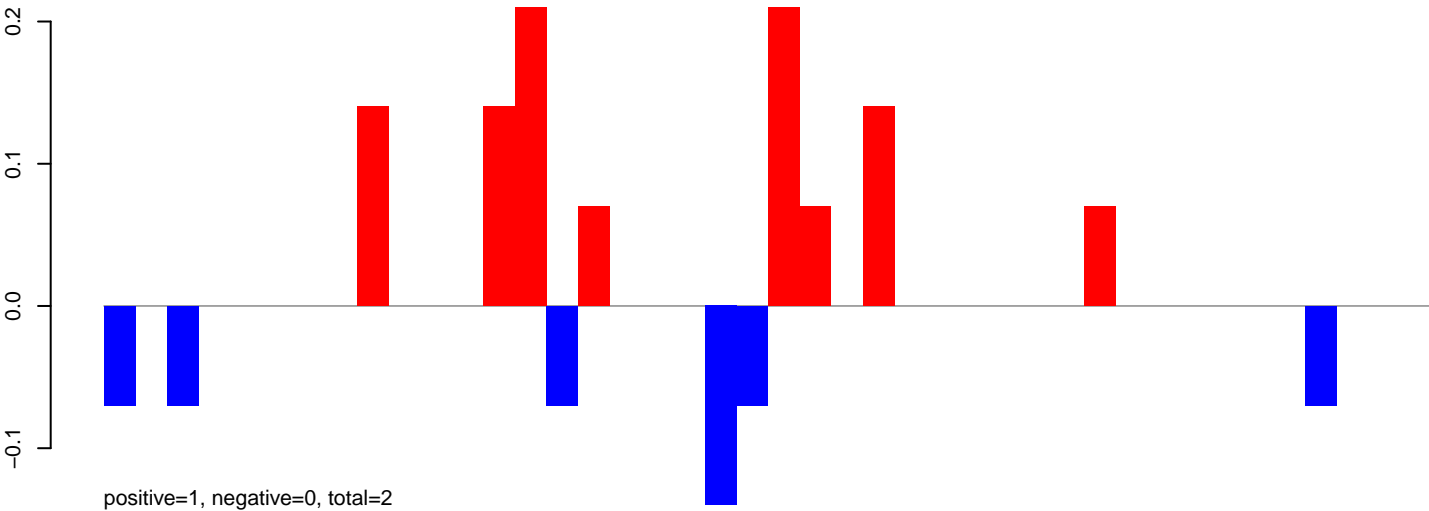
CuQuin_Mcarc_NL.rep



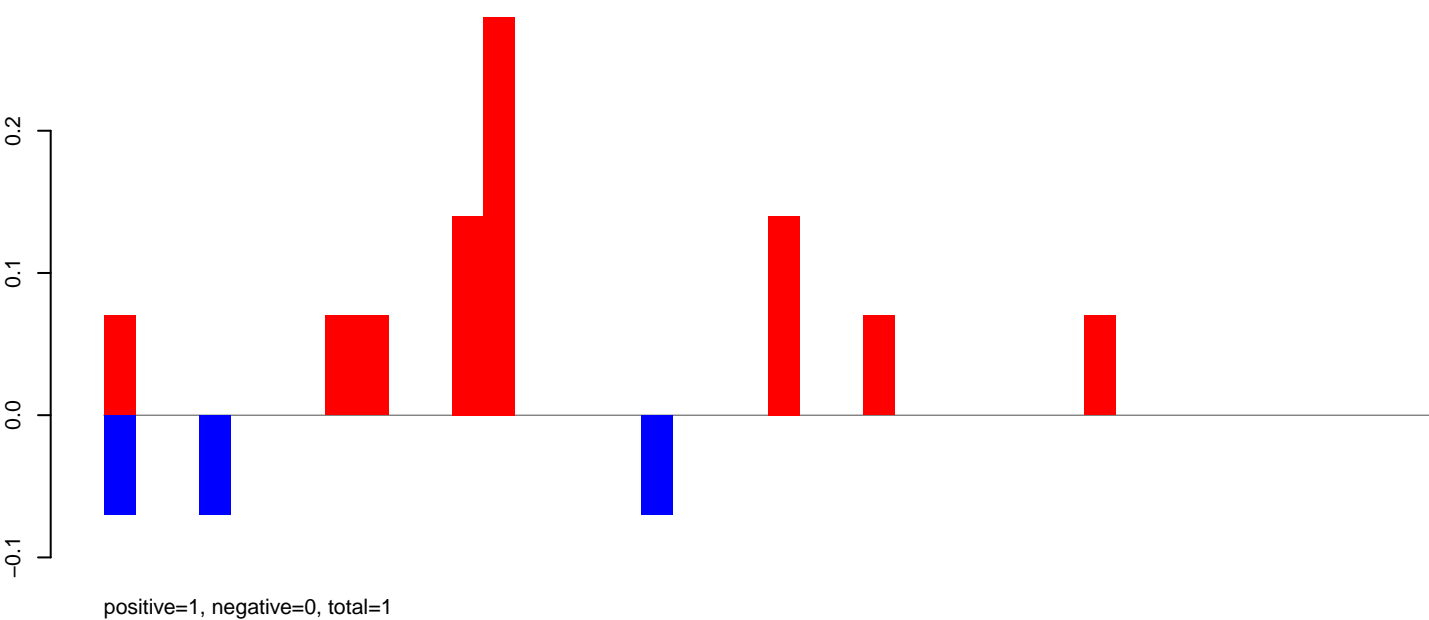
Window size=25, length=1683, TE@rnd-5_family-645@DNA_MULE-MuDR@16431-1683

0 500 1000 1500

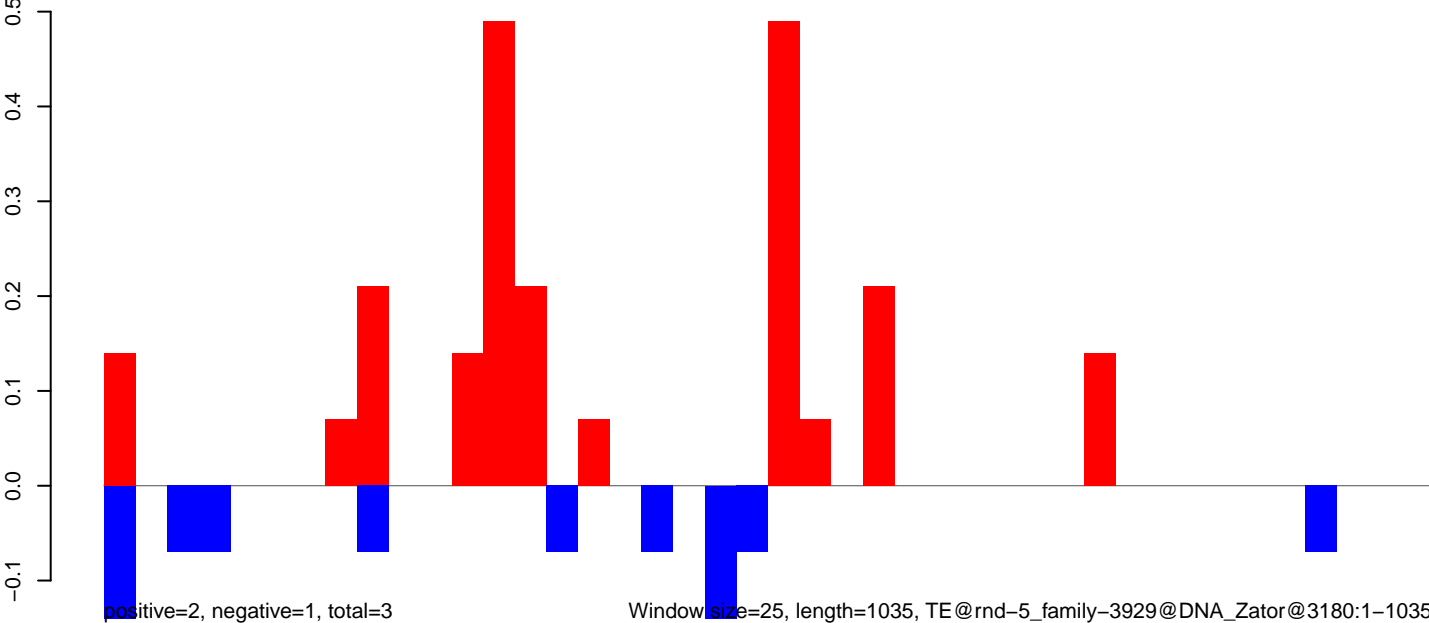
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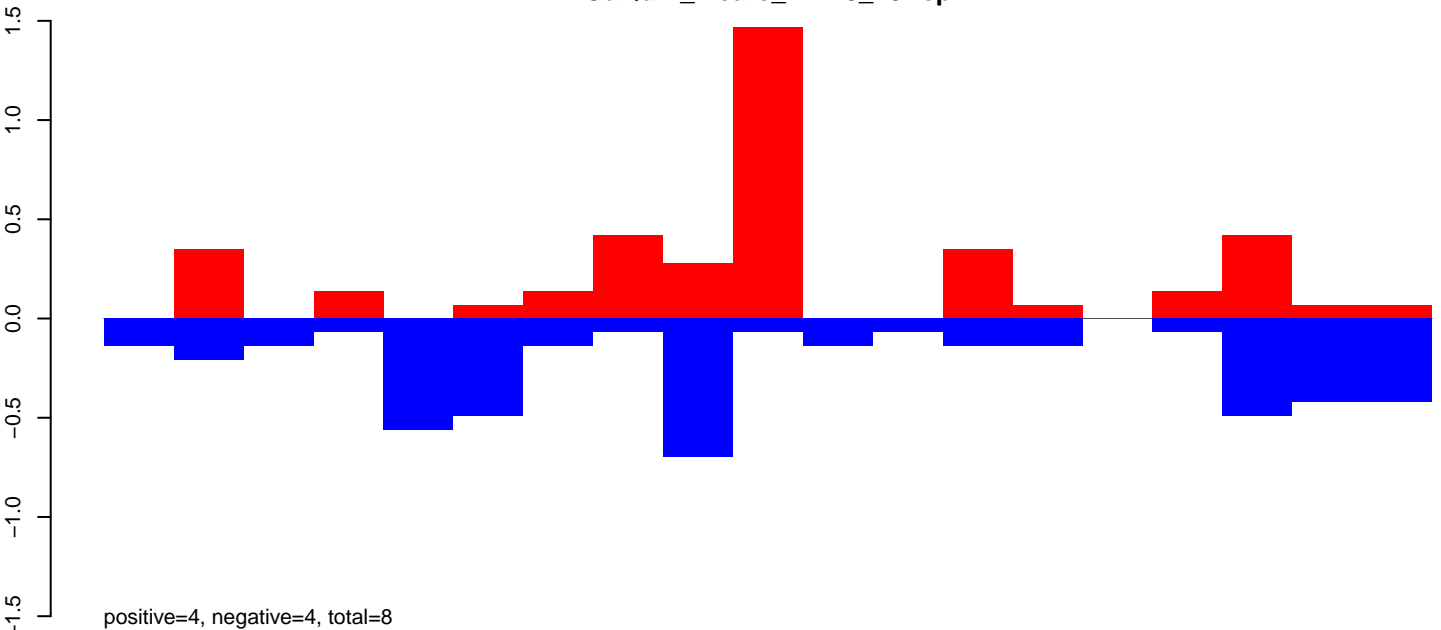
CuQuin_Mcarc_NL.24_35.rep



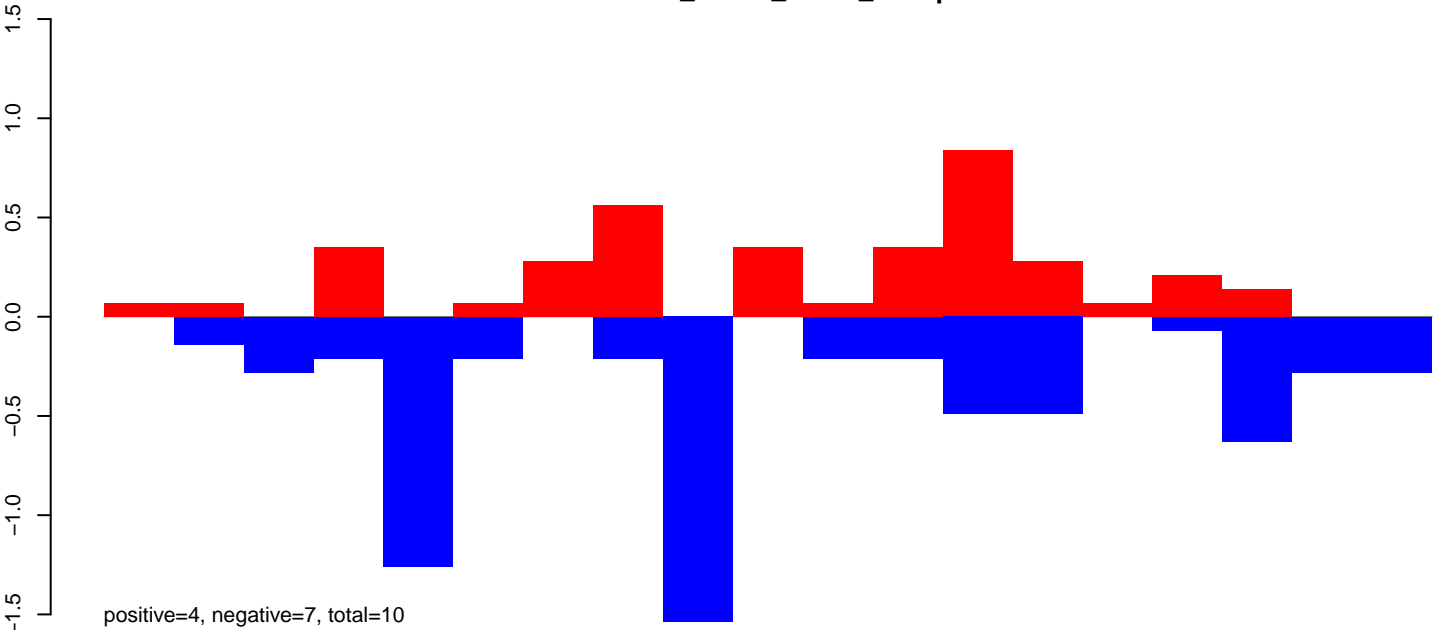
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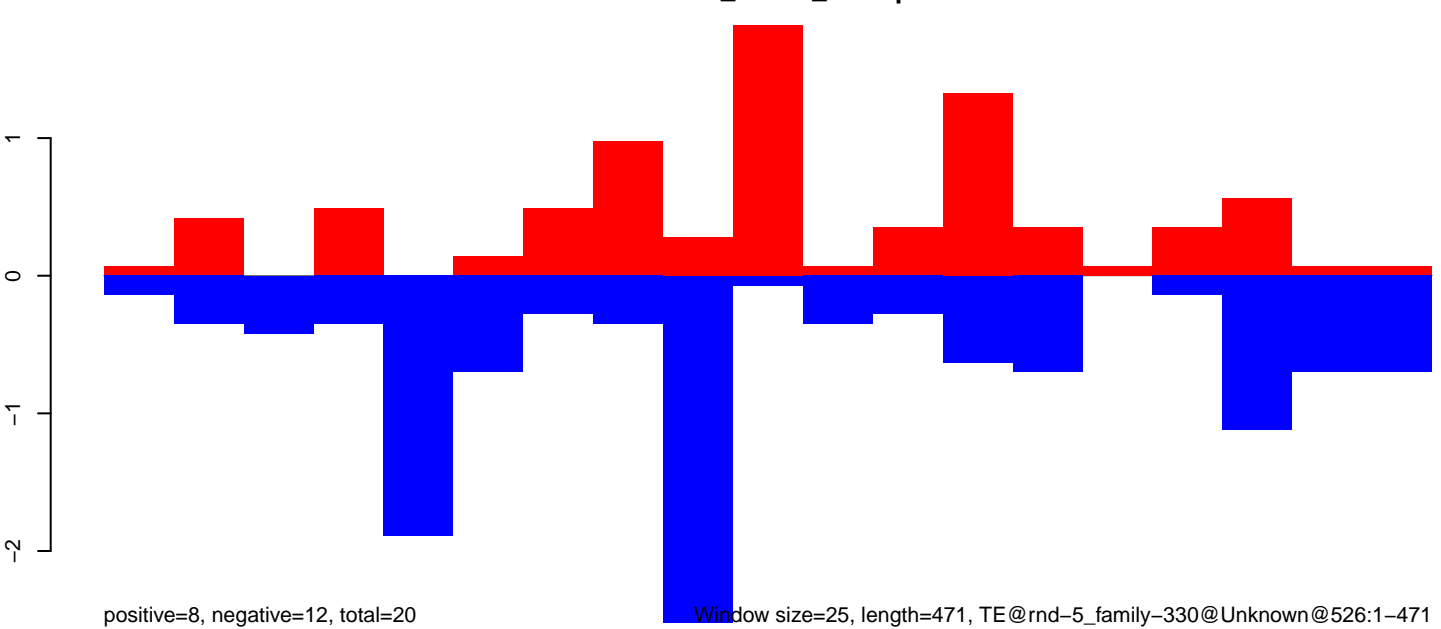
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CuQuin_Mcarc_NL.24_35.rep



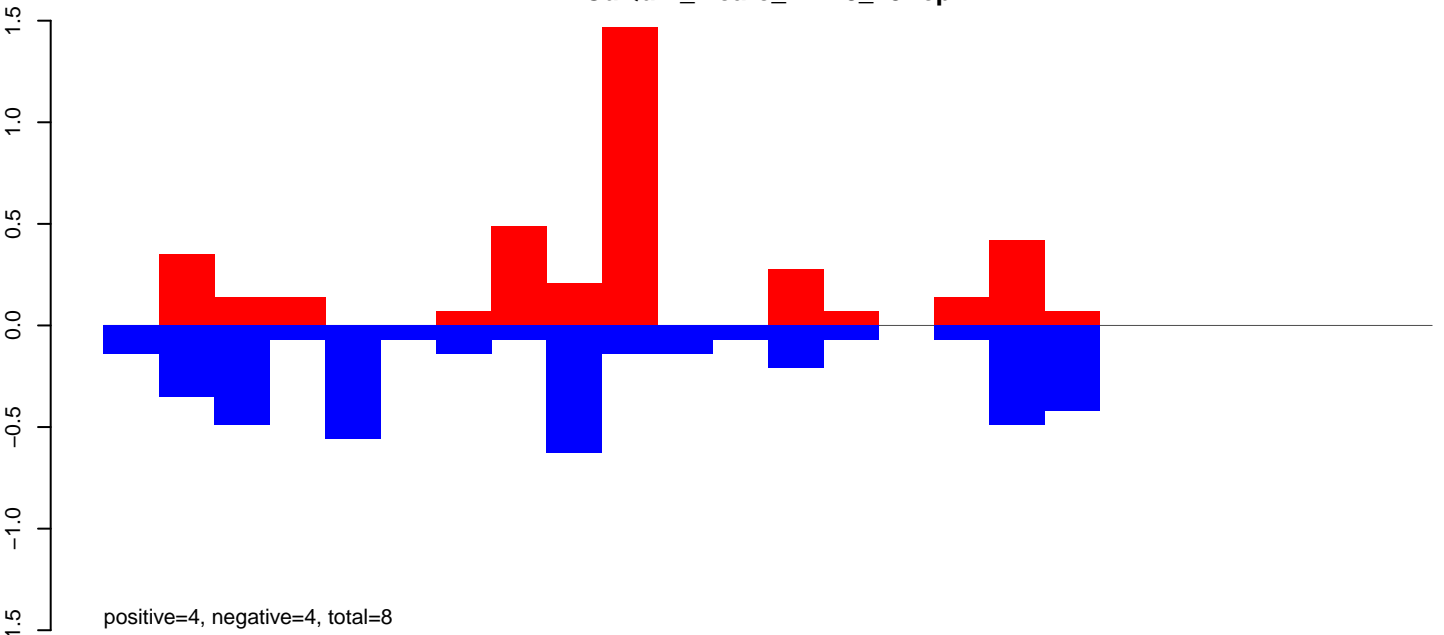
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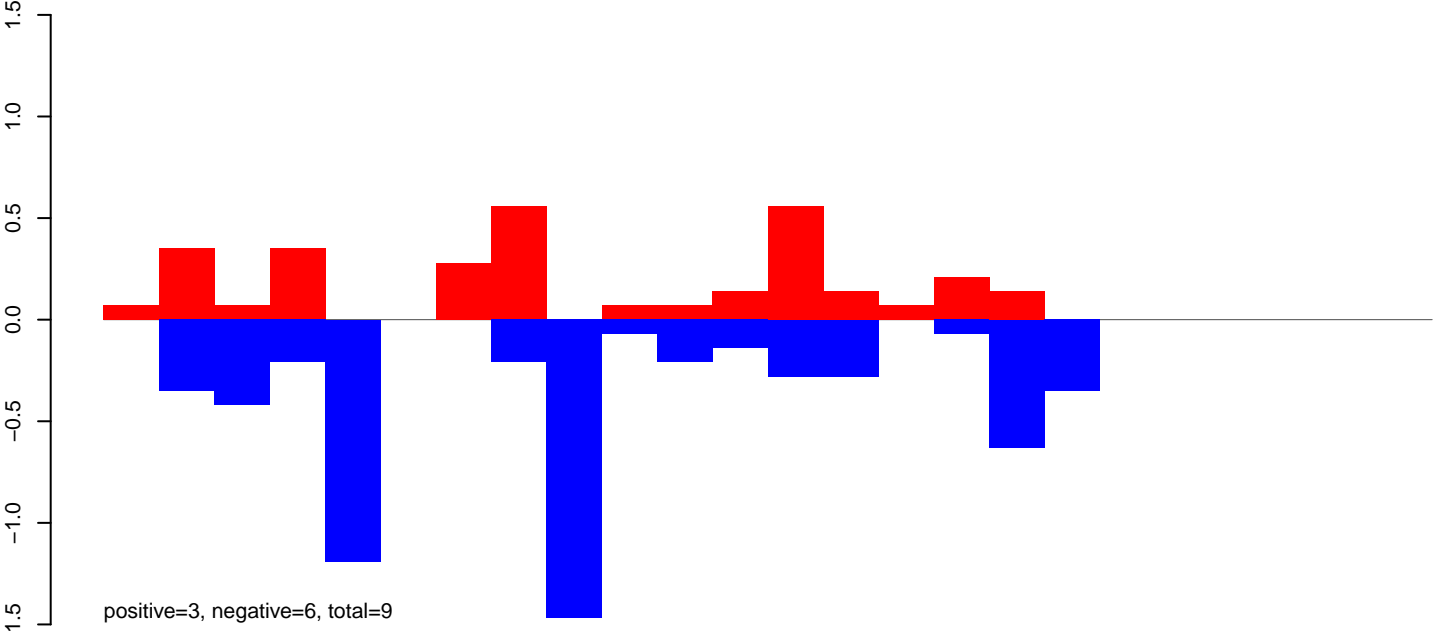
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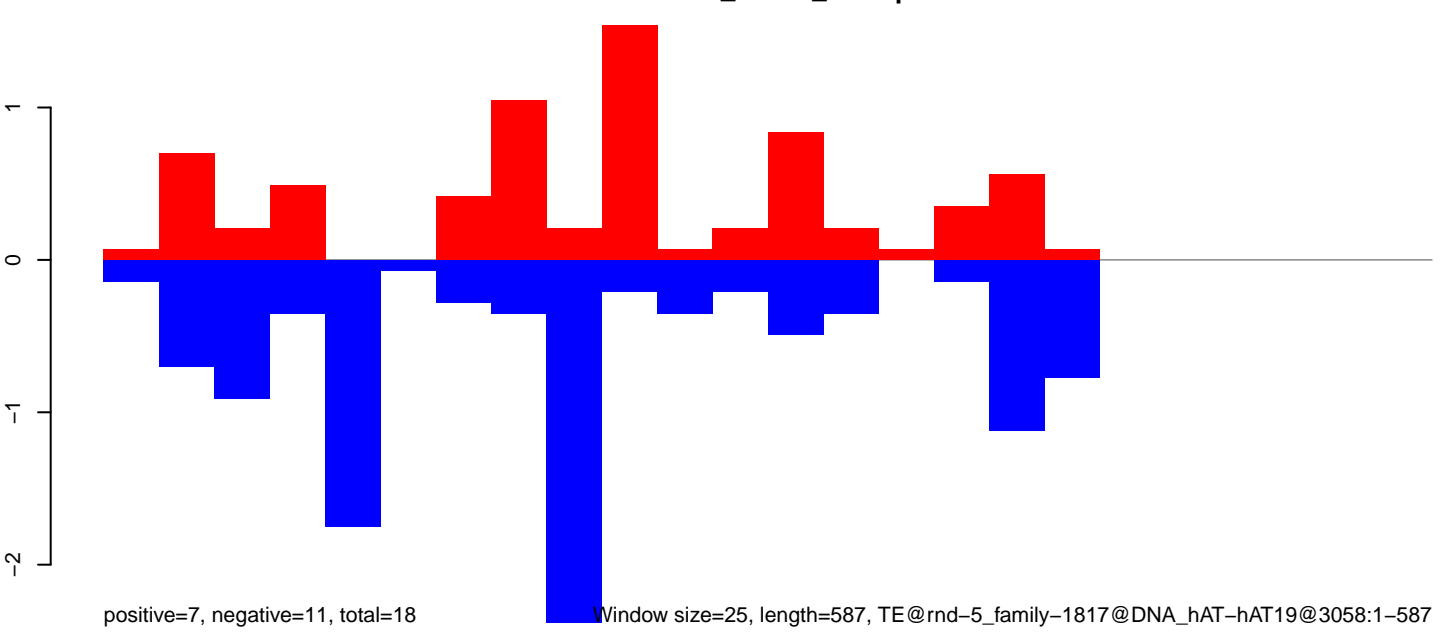
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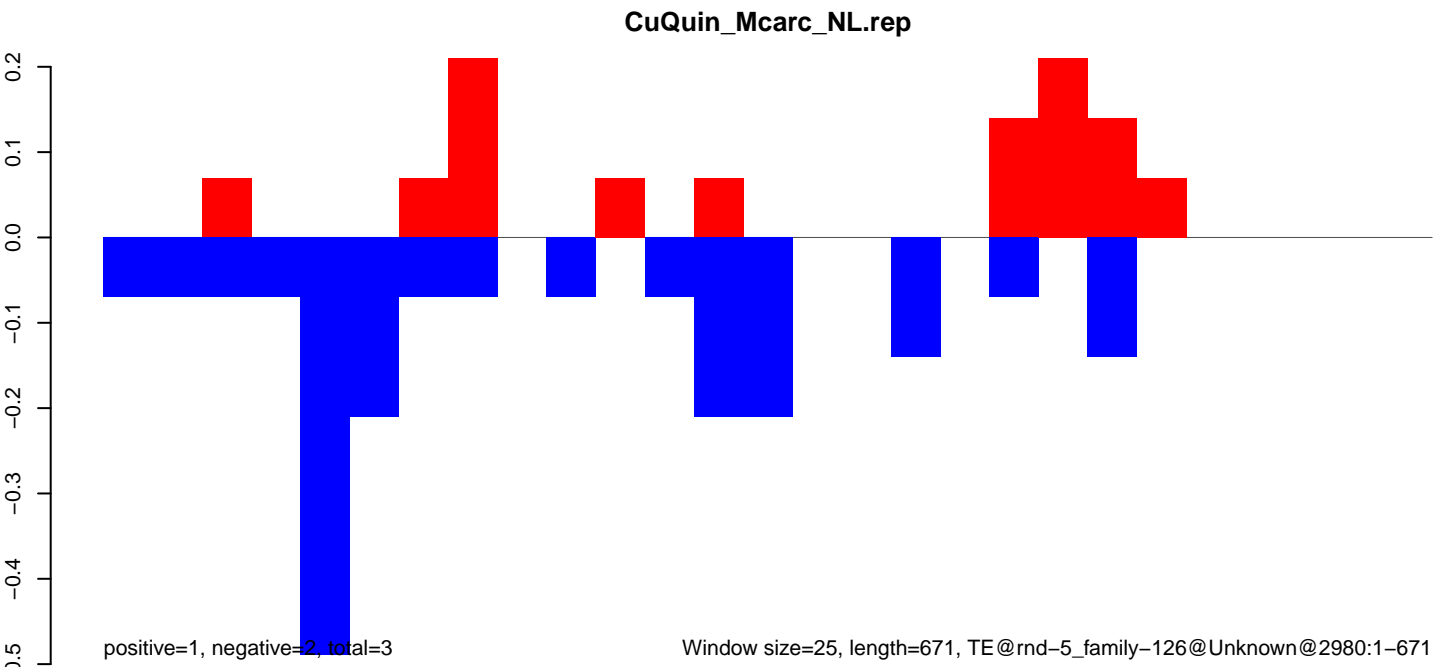
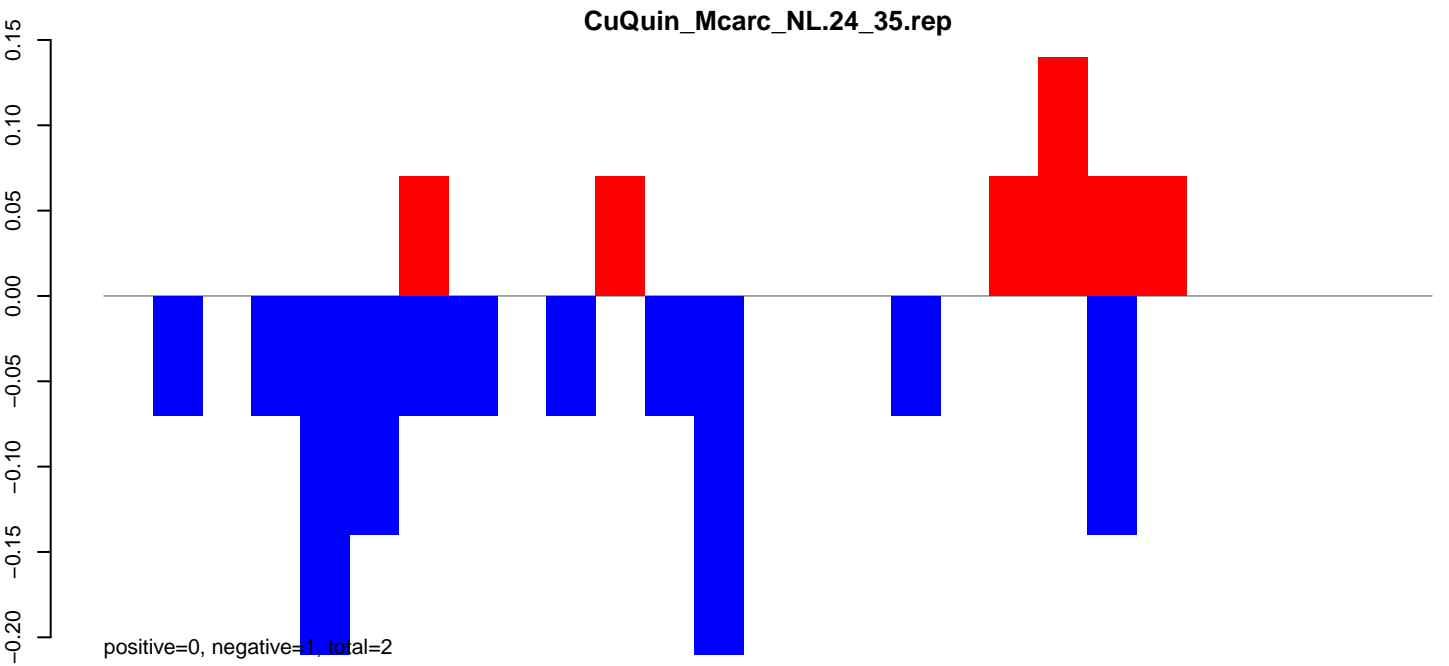
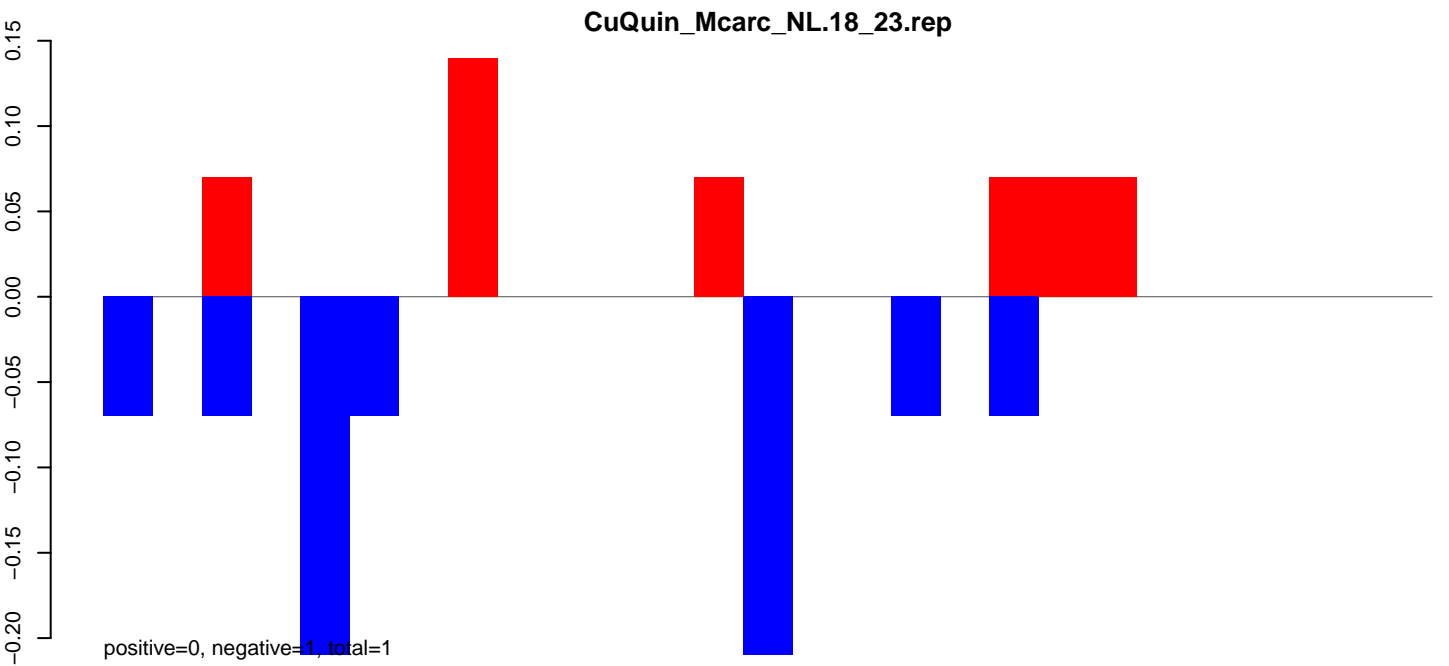
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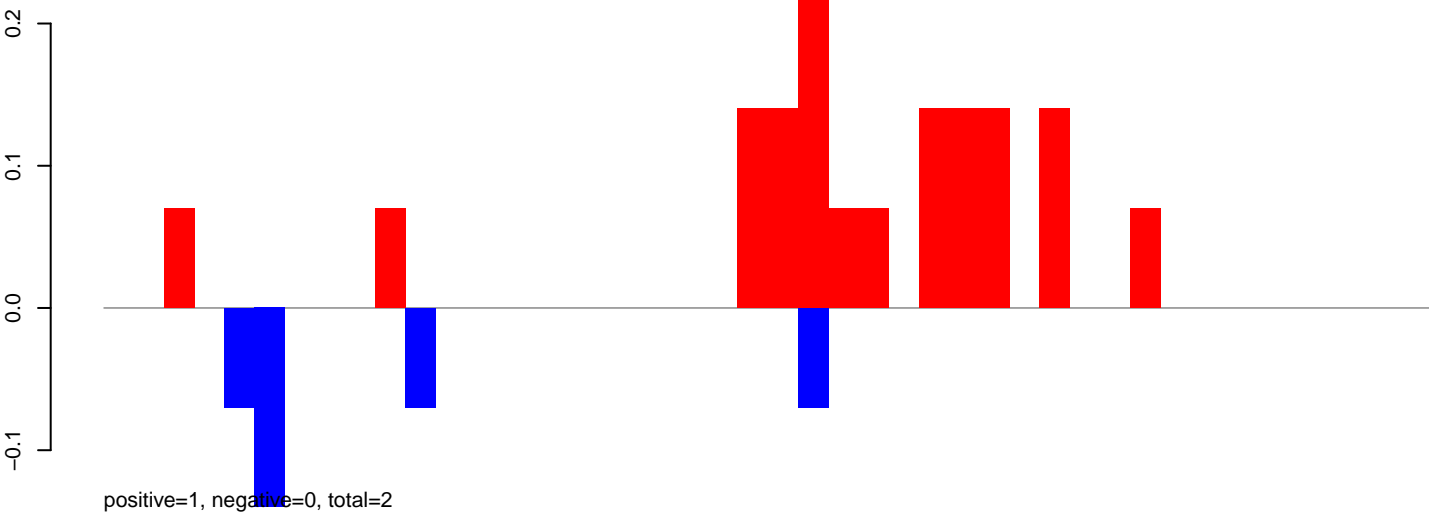
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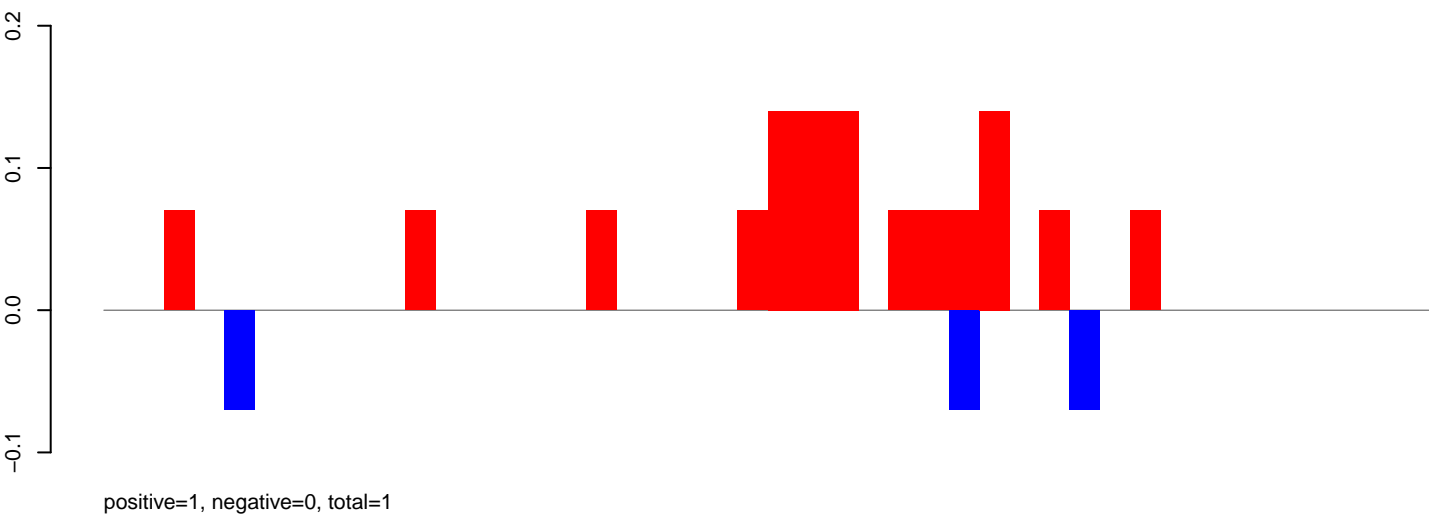
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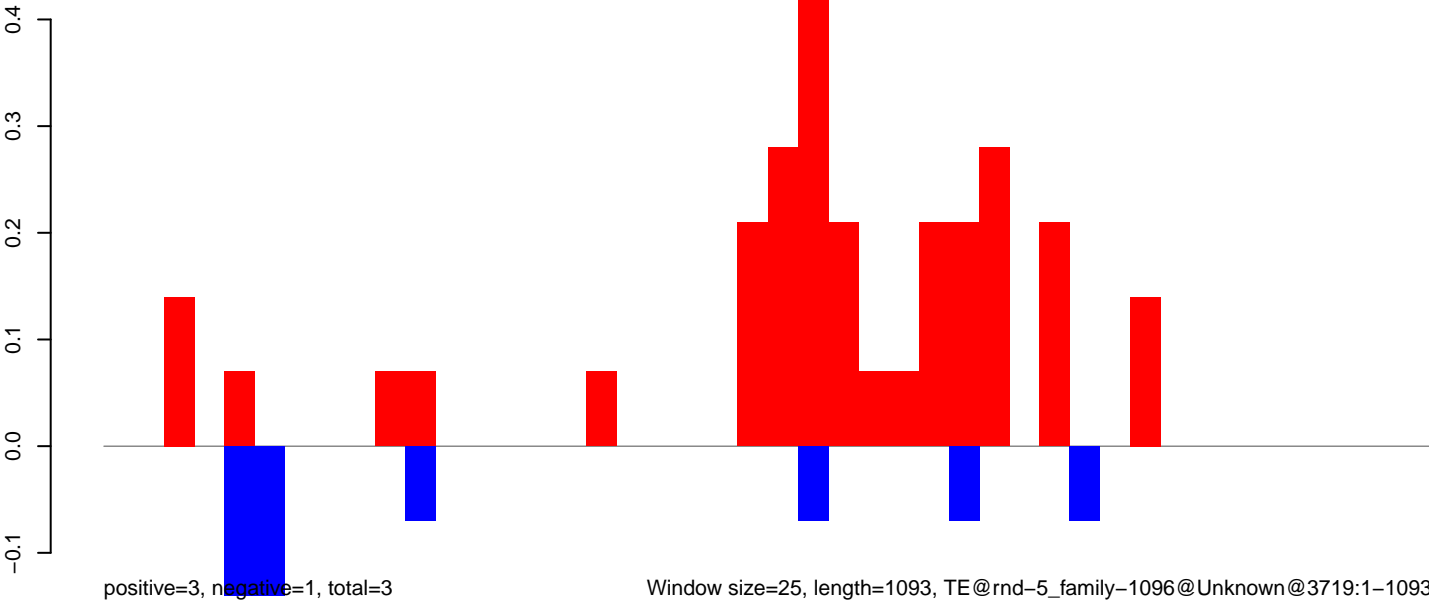
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CuQuin_Mcarc_NL.24_35.rep

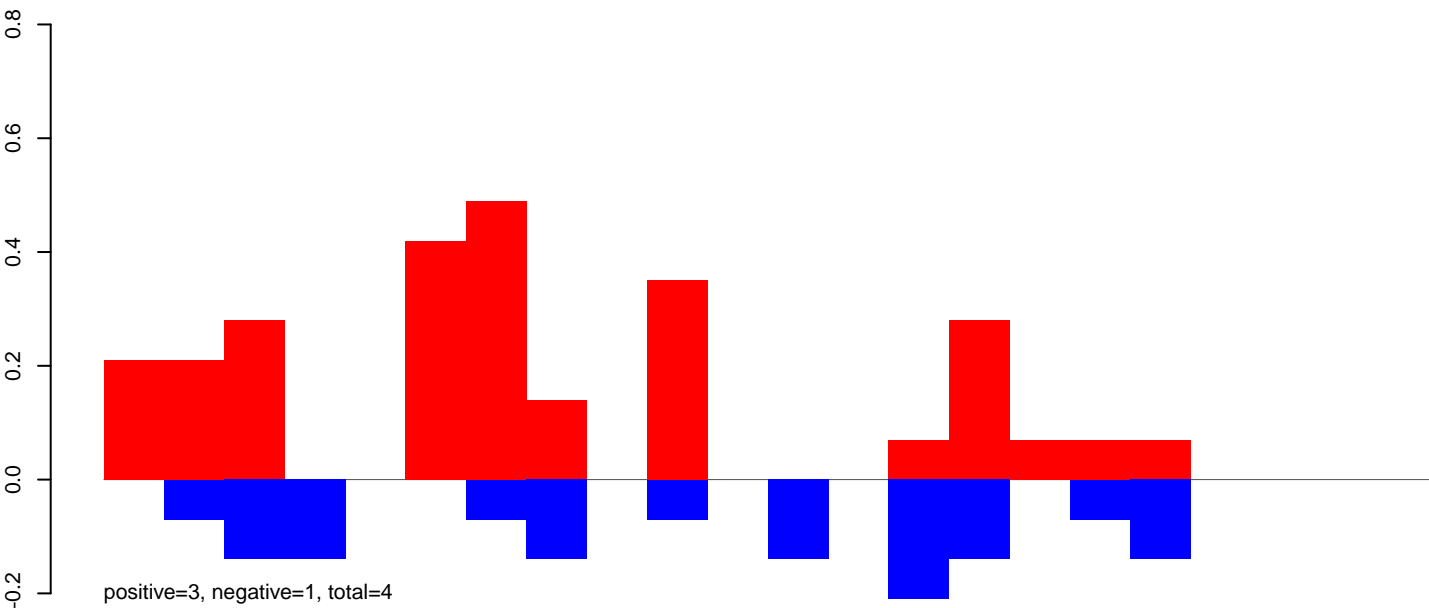


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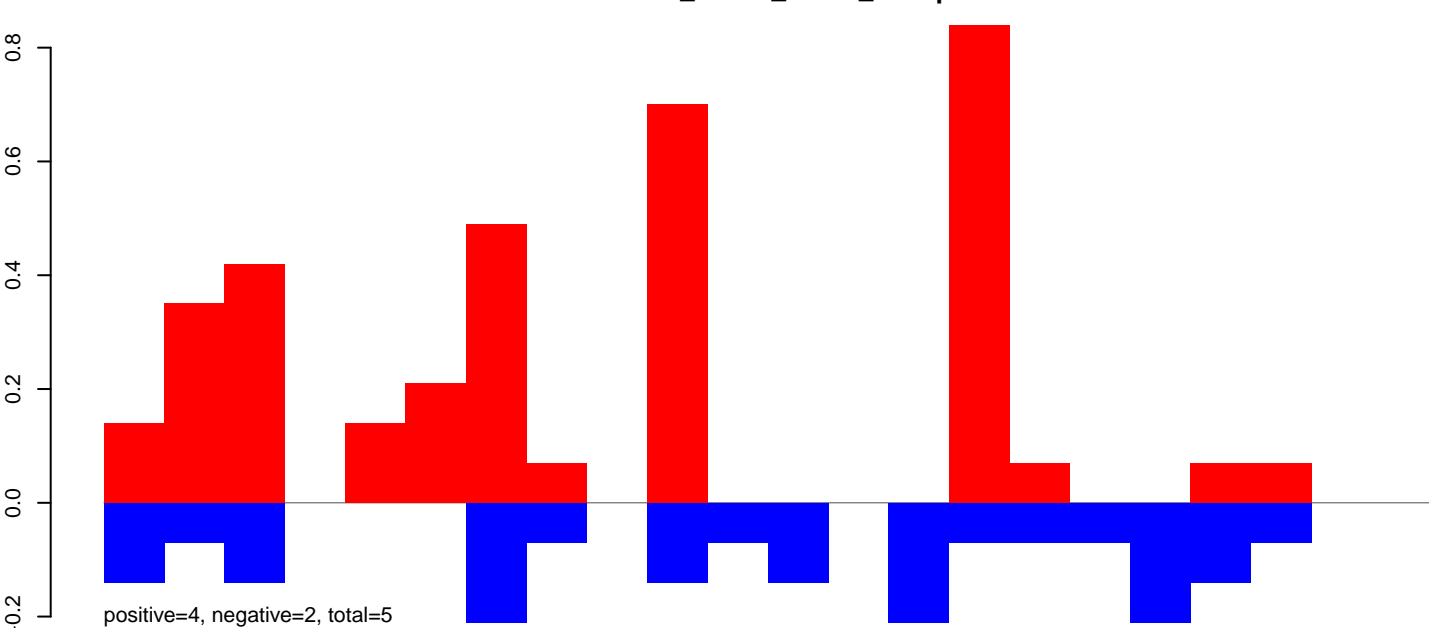


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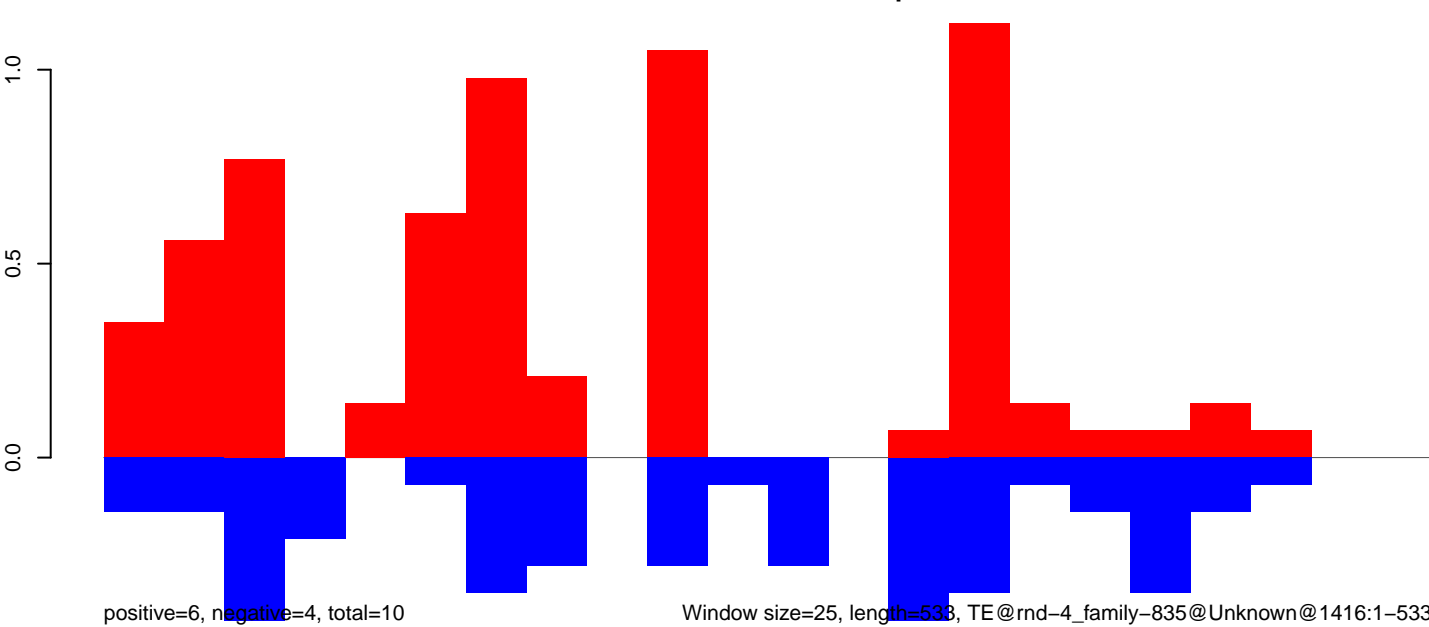
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

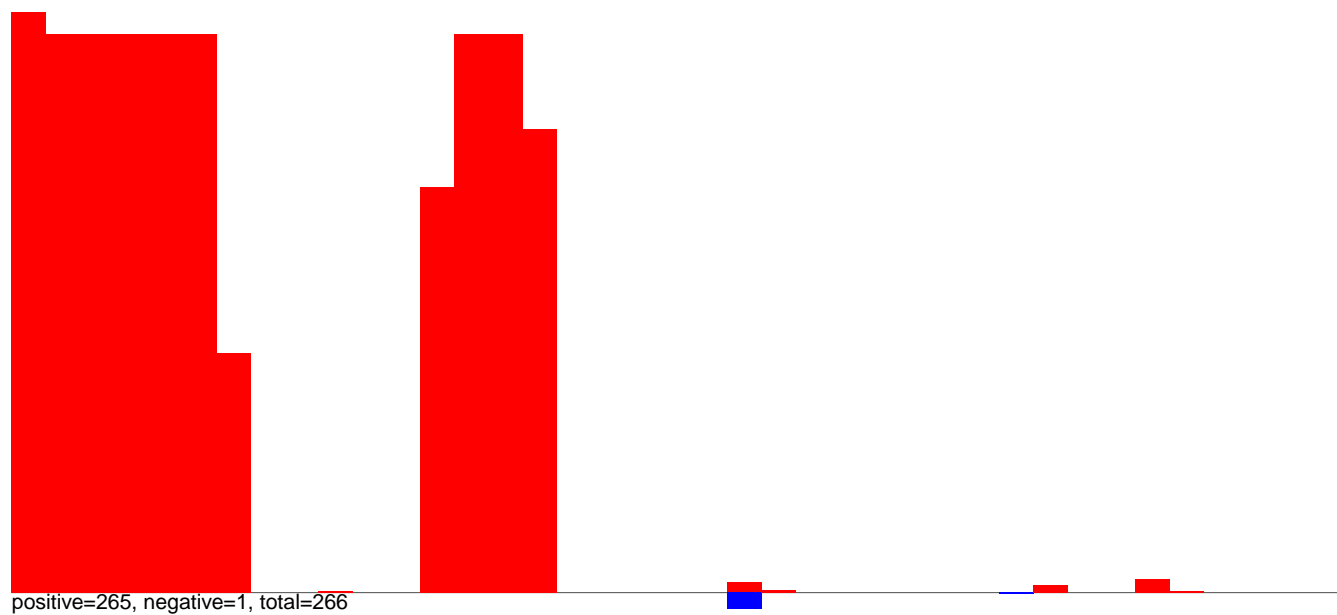


CuQuin_Mcarc_NL.rep

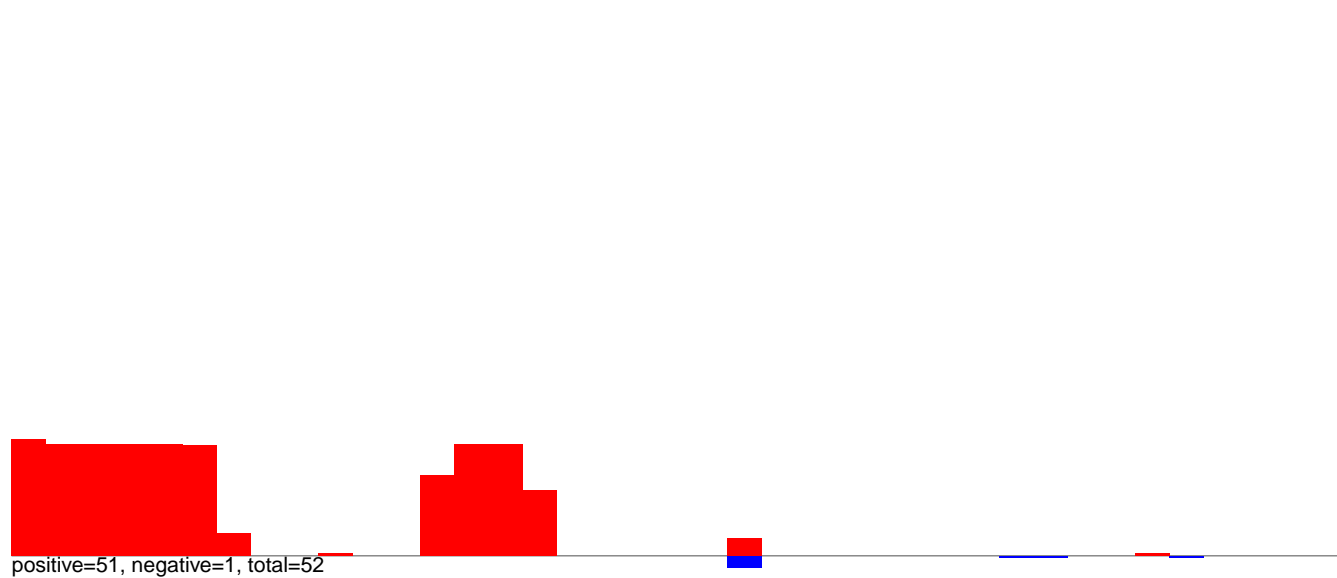


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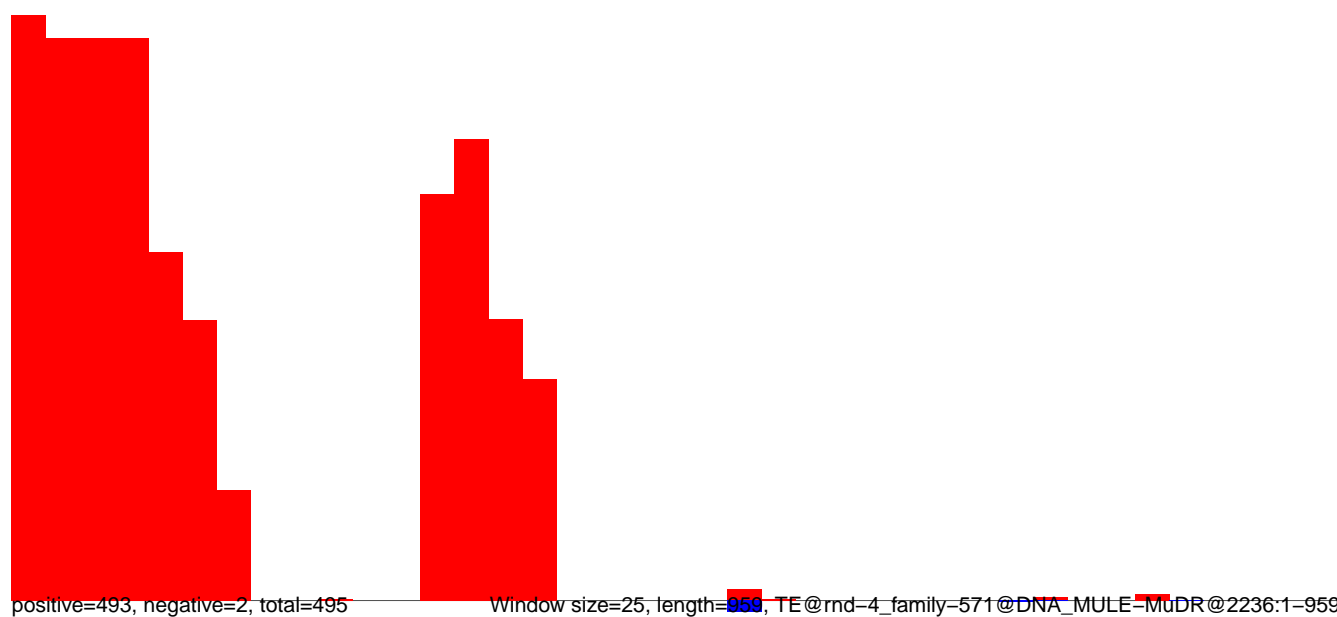
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

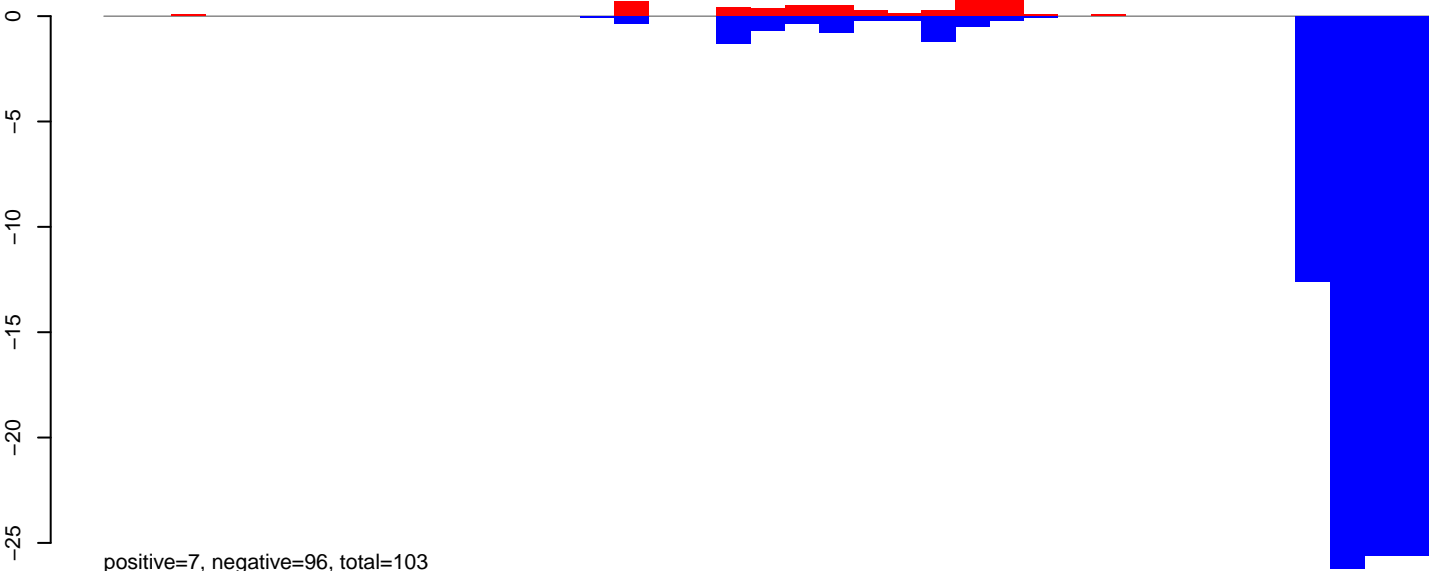


CuQuin_Mcarc_NL.rep

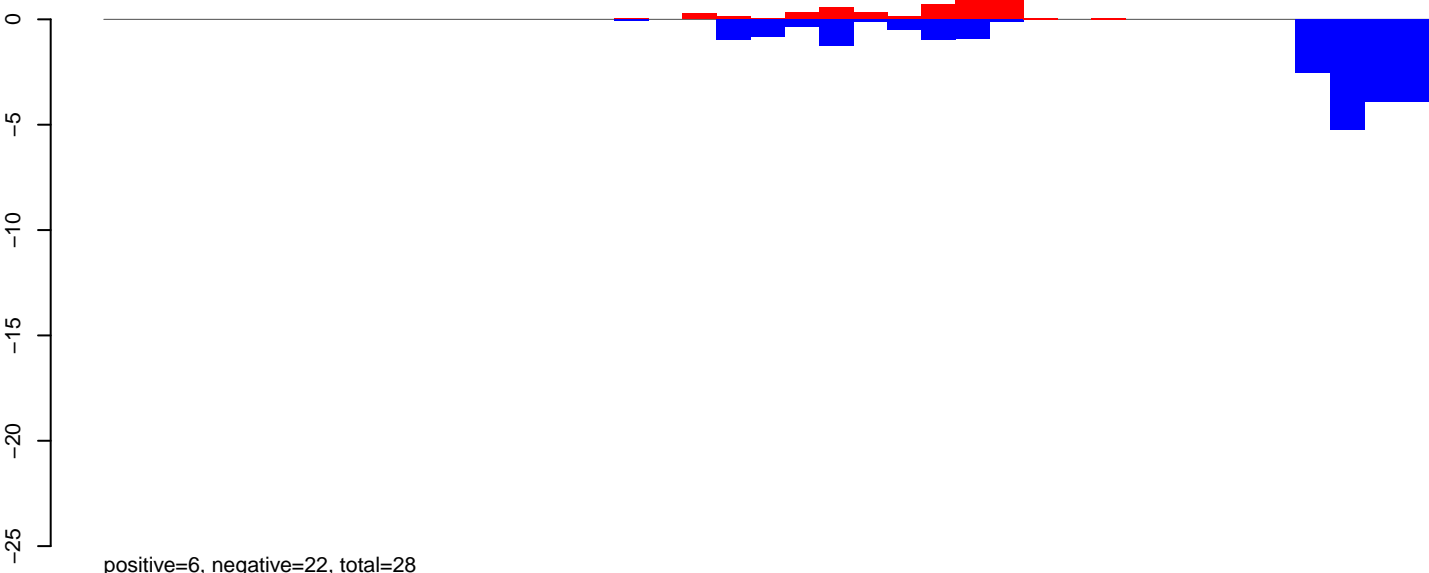


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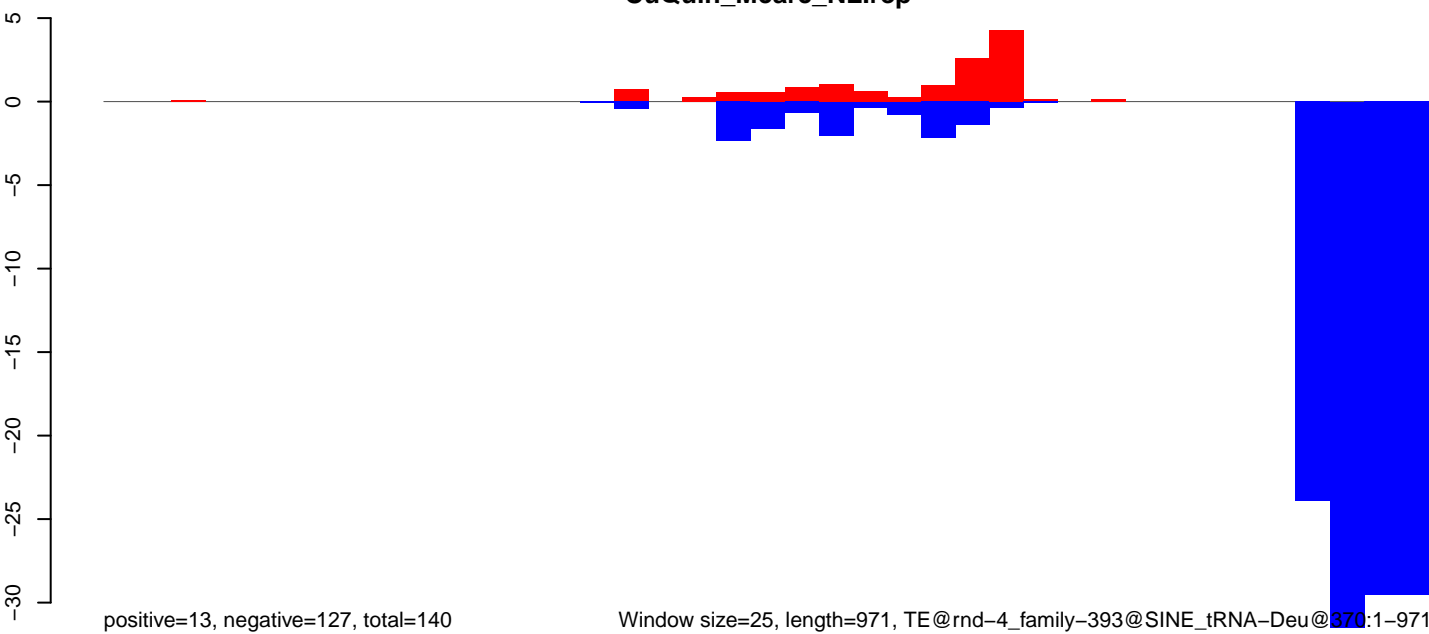
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



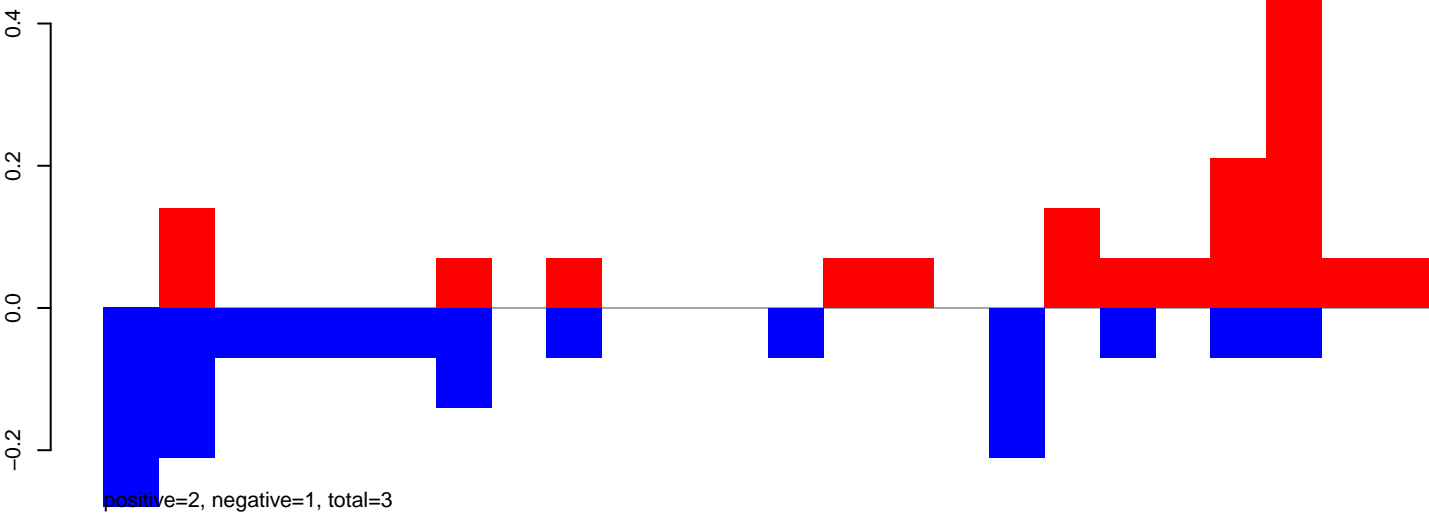
CuQuin_Mcarc_NL.rep



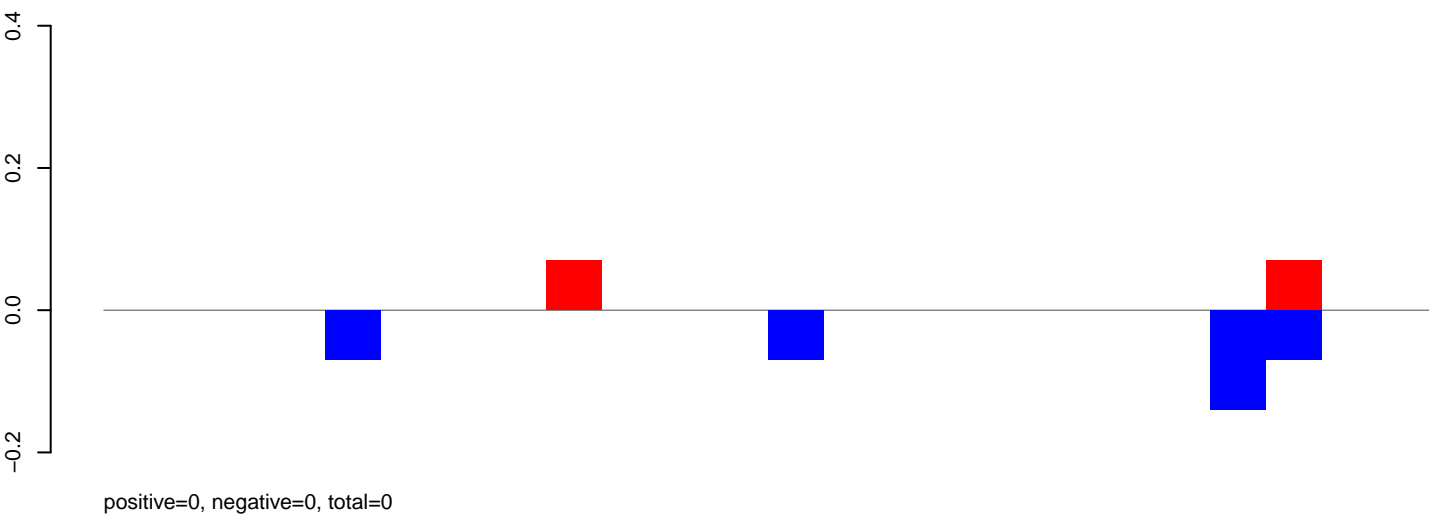
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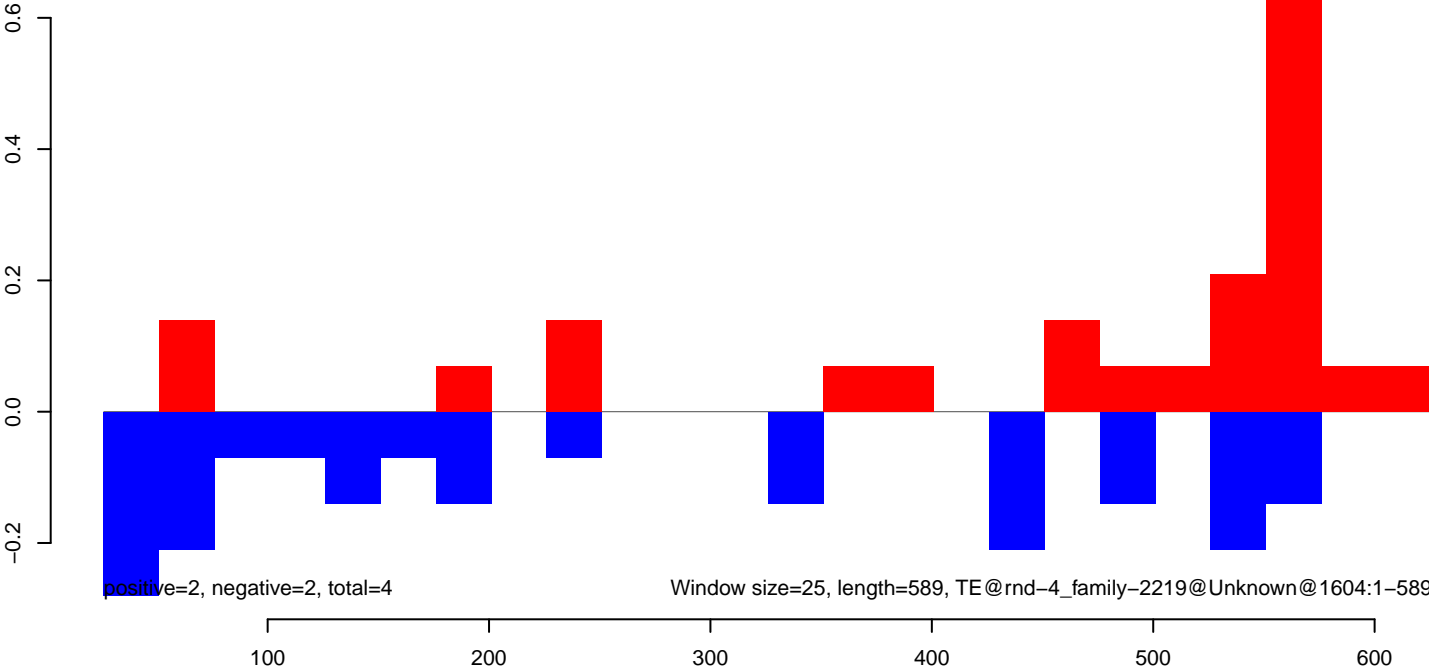
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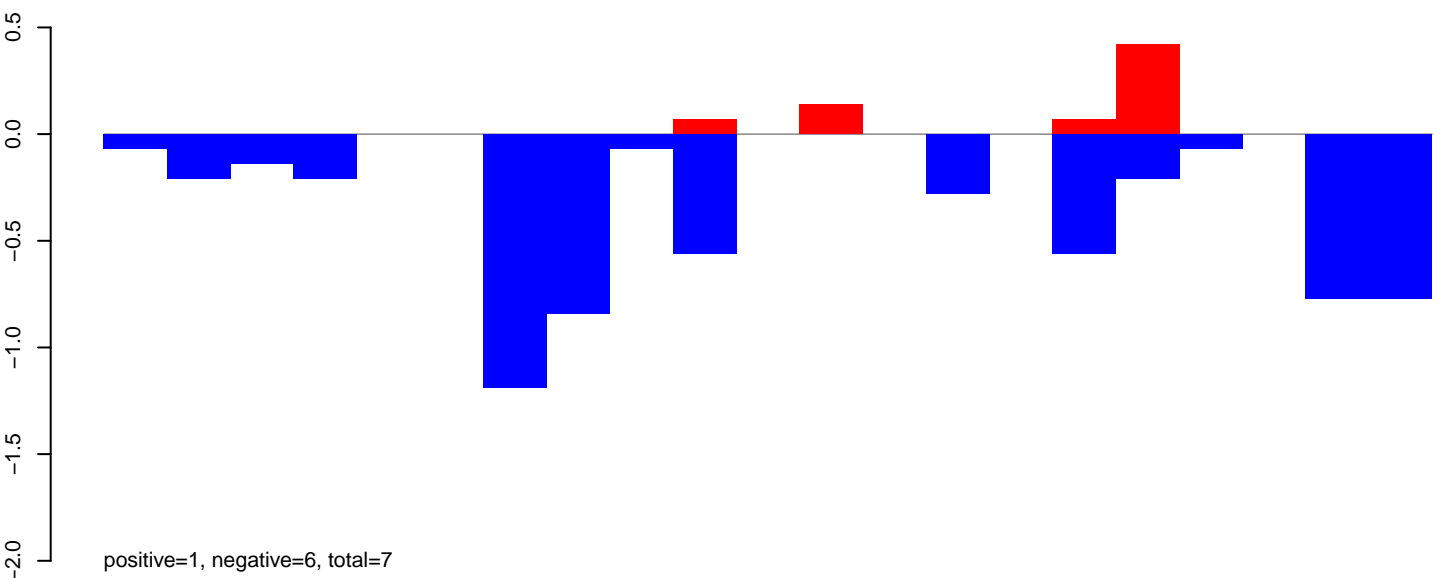
CuQuin_Mcarc_NL.24_35.rep



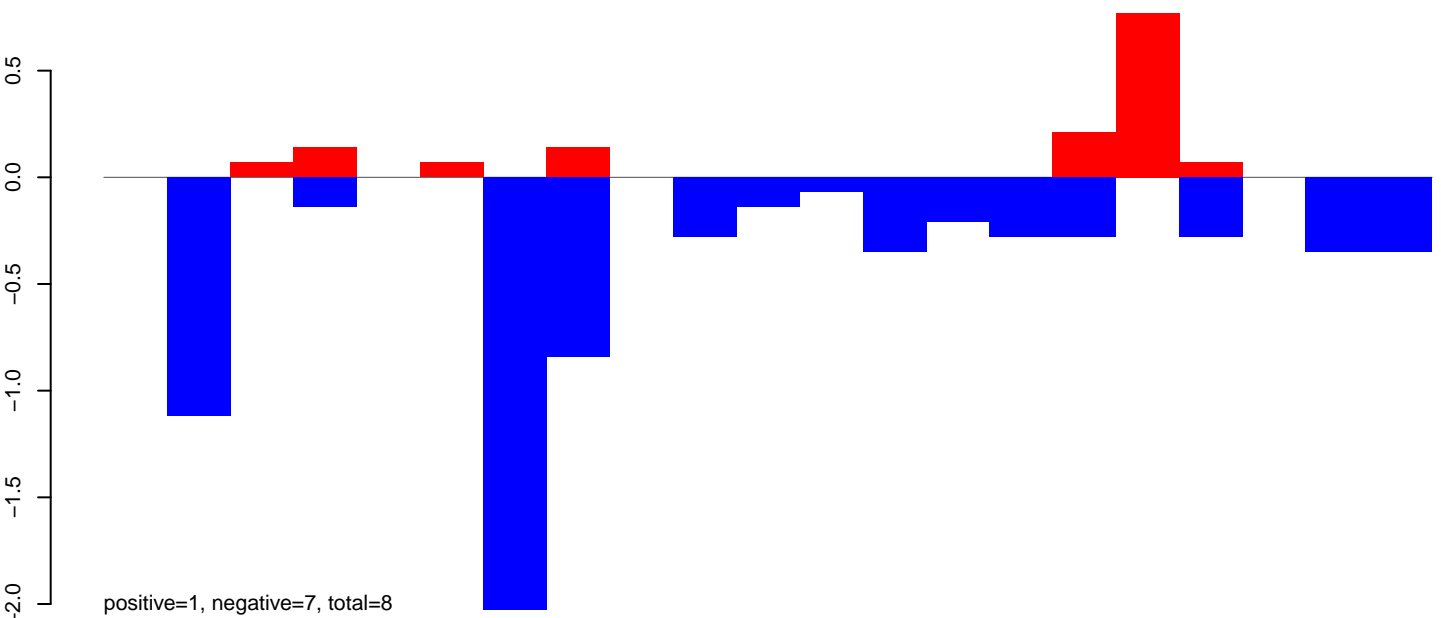
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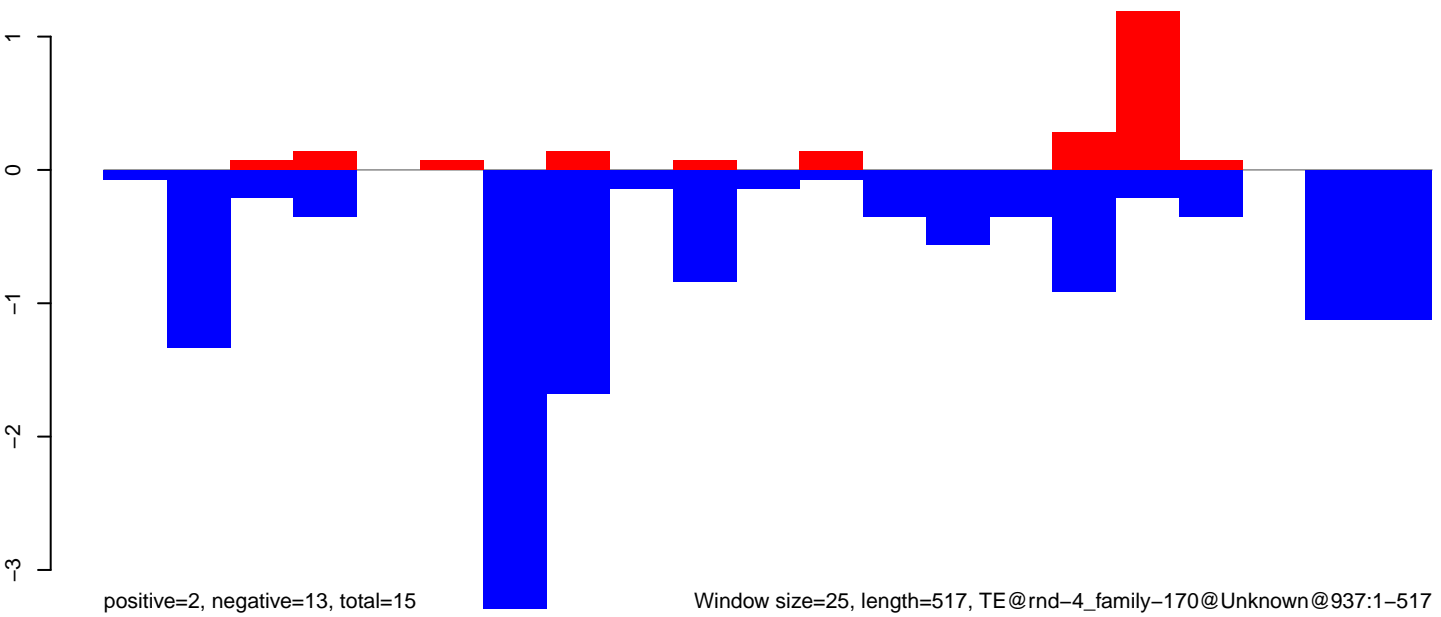
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CuQuin_Mcarc_NL.24_35.rep

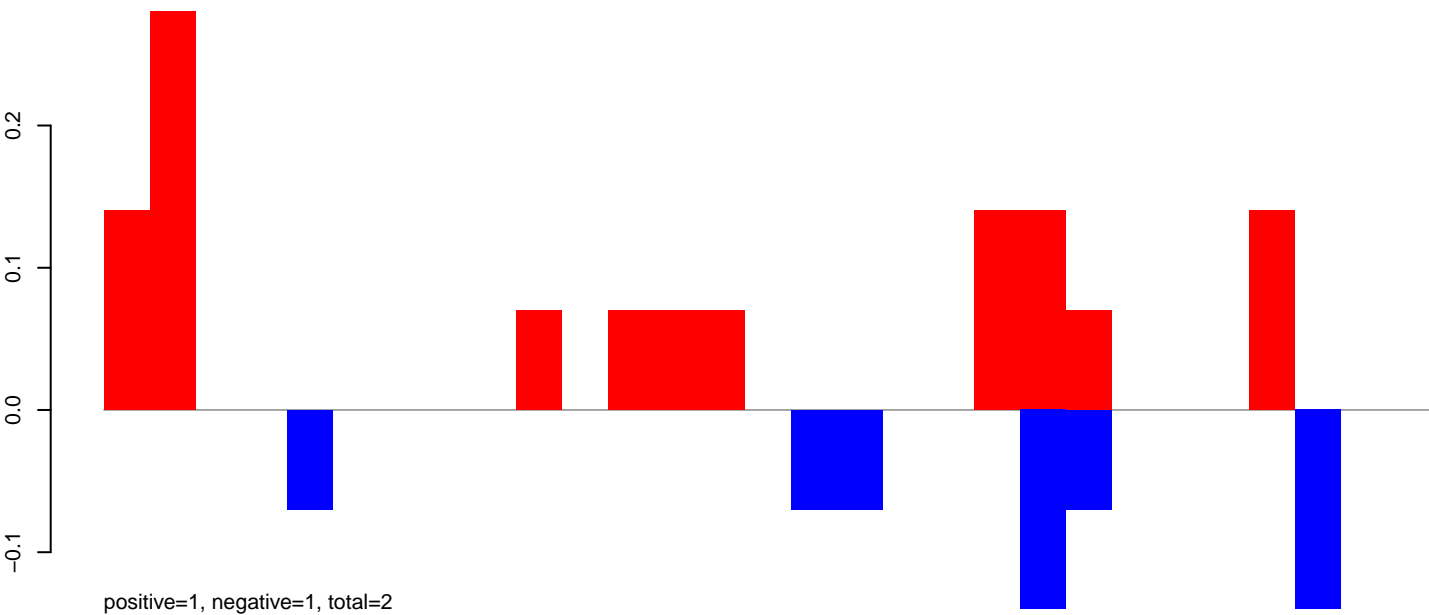


CuQuin_Mcarc_NL.rep

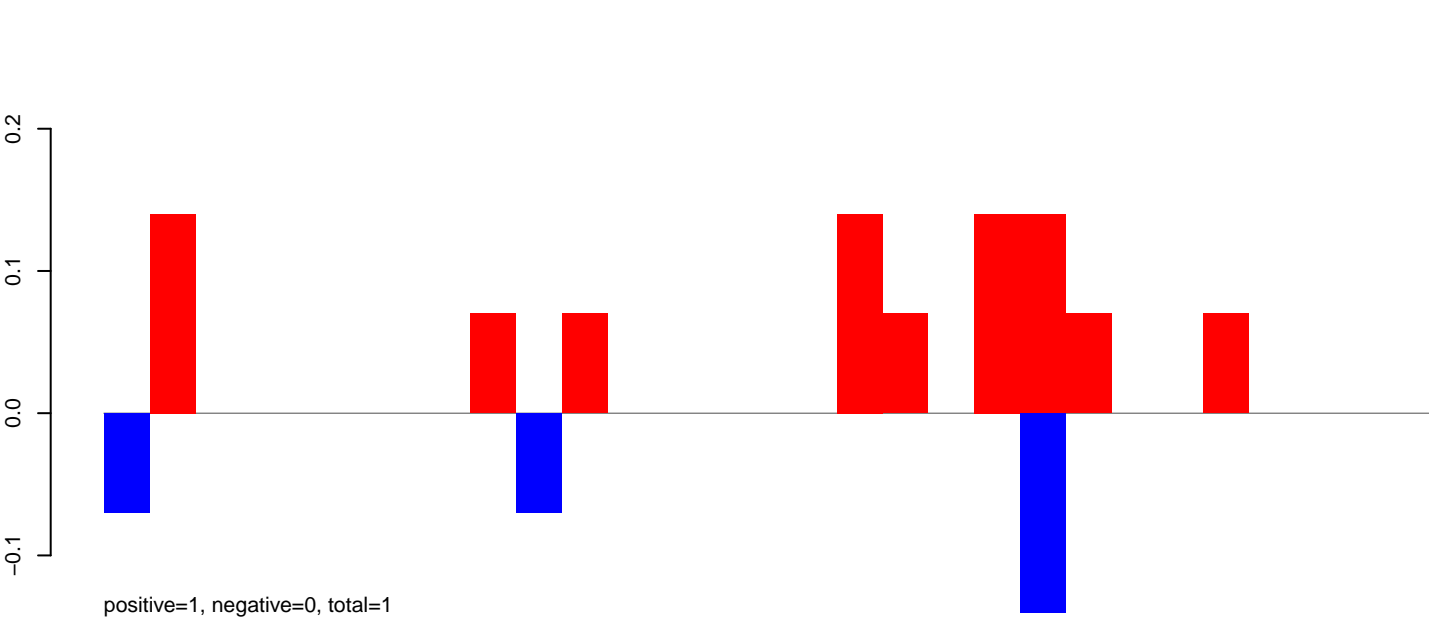


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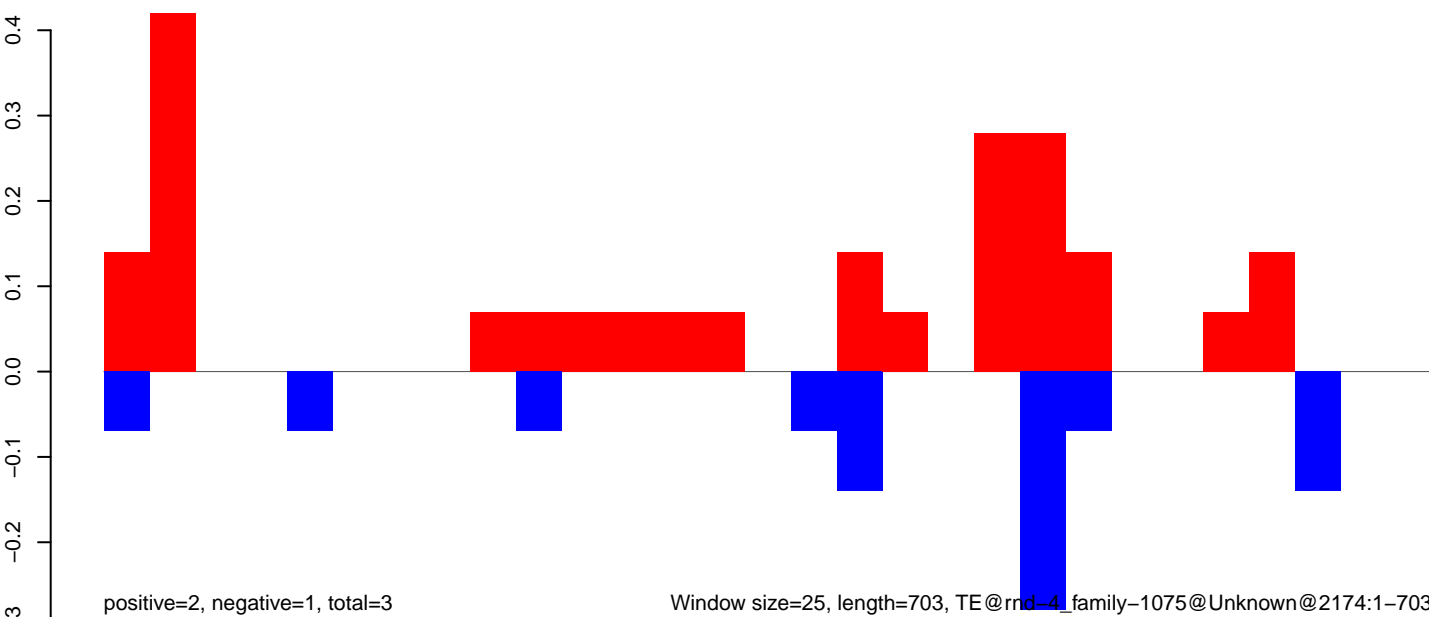
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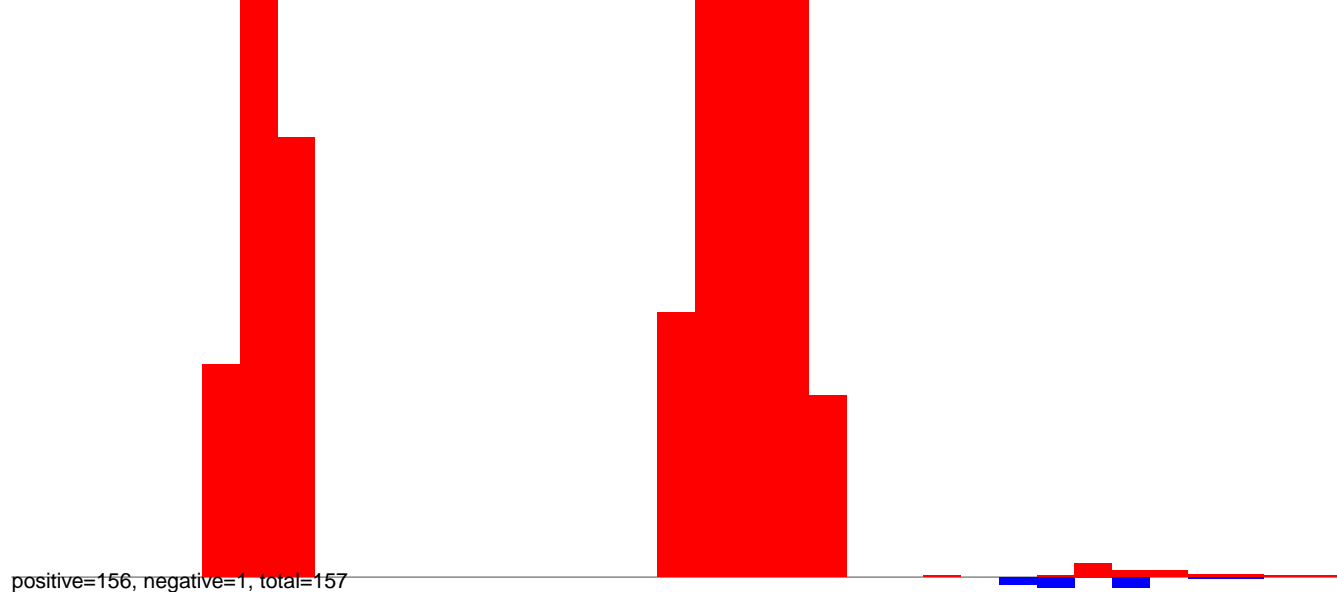
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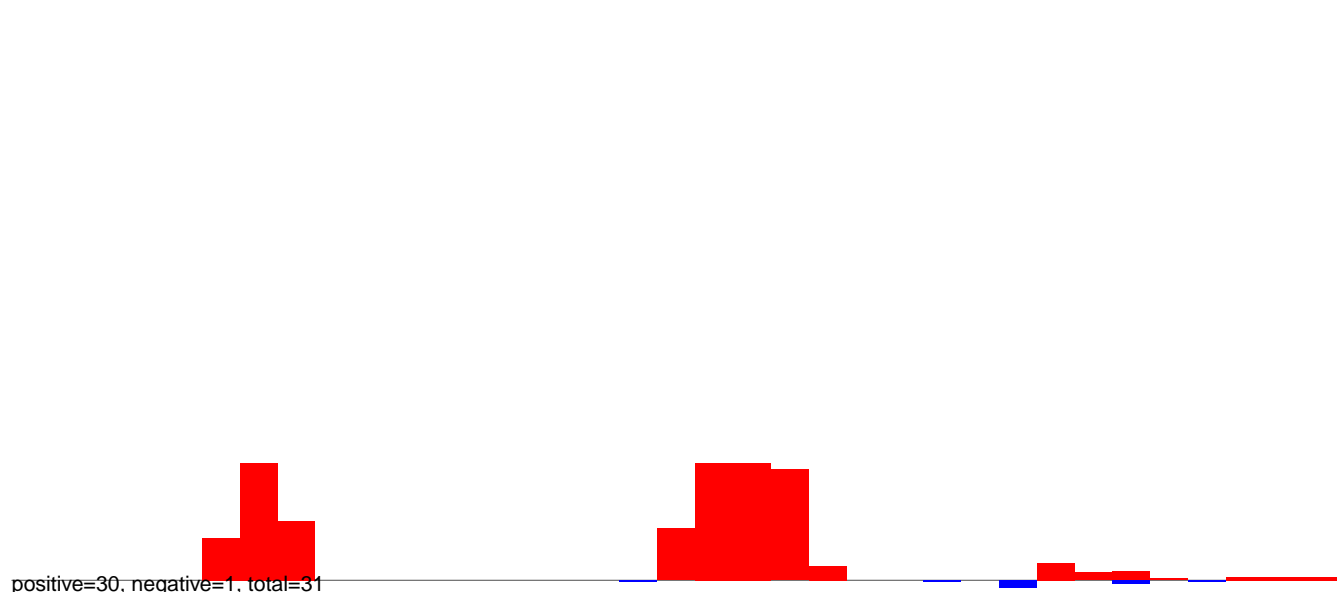
CuQuin_Mcarc_NL.rep



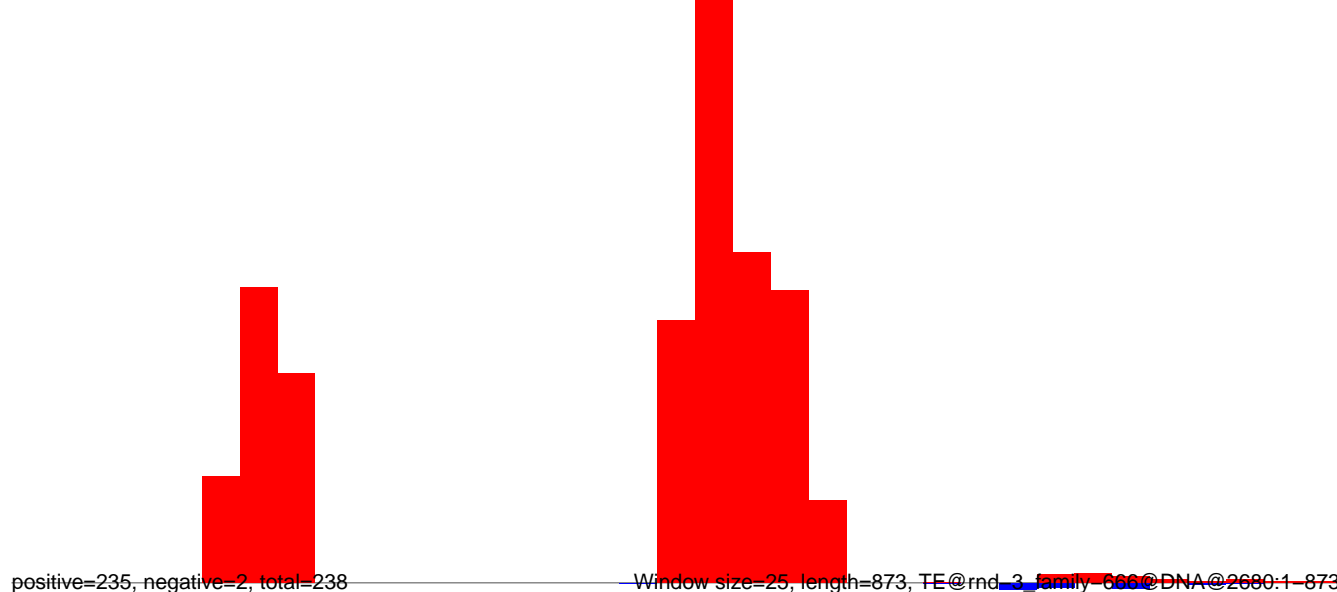
CuQuin_Mcarc_NL.18_23.rep



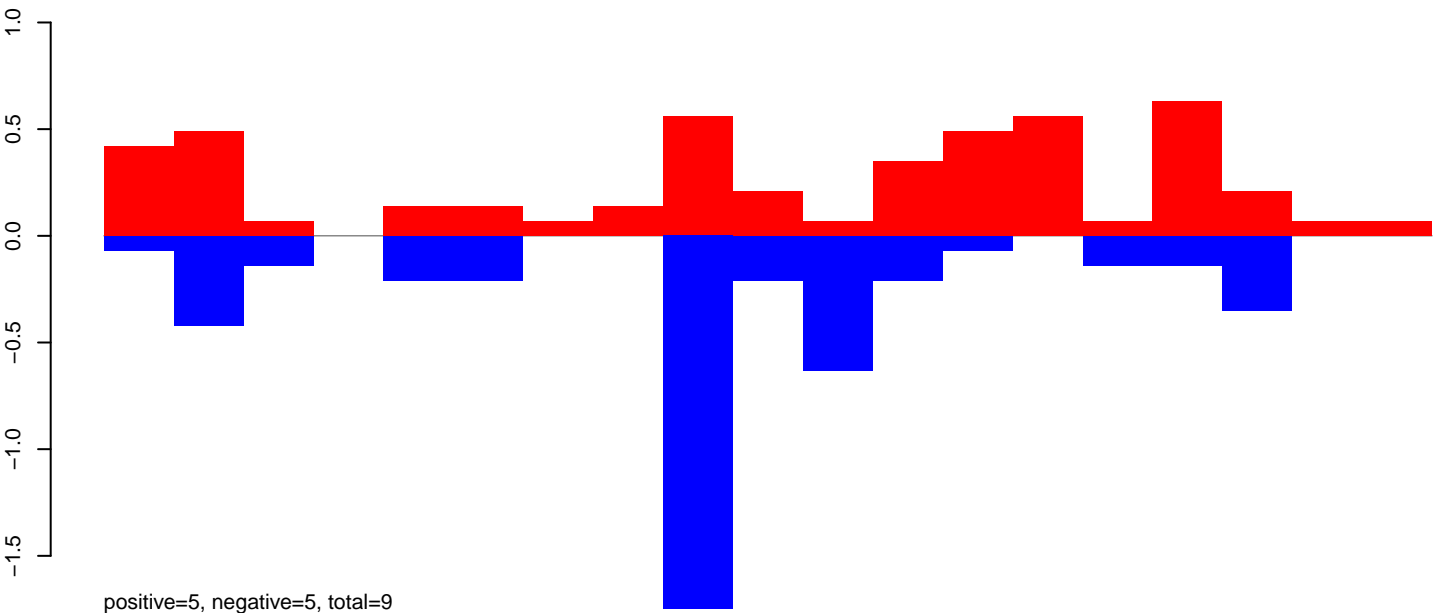
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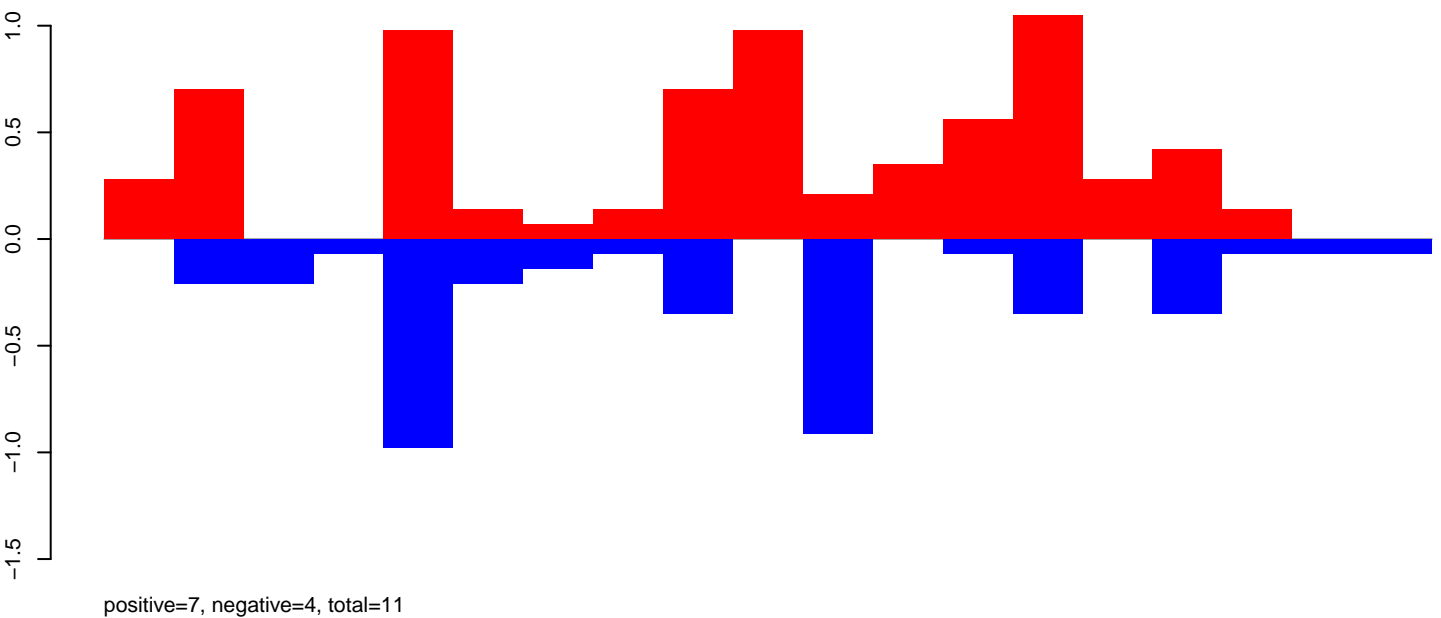
CuQuin_Mcarc_NL.rep



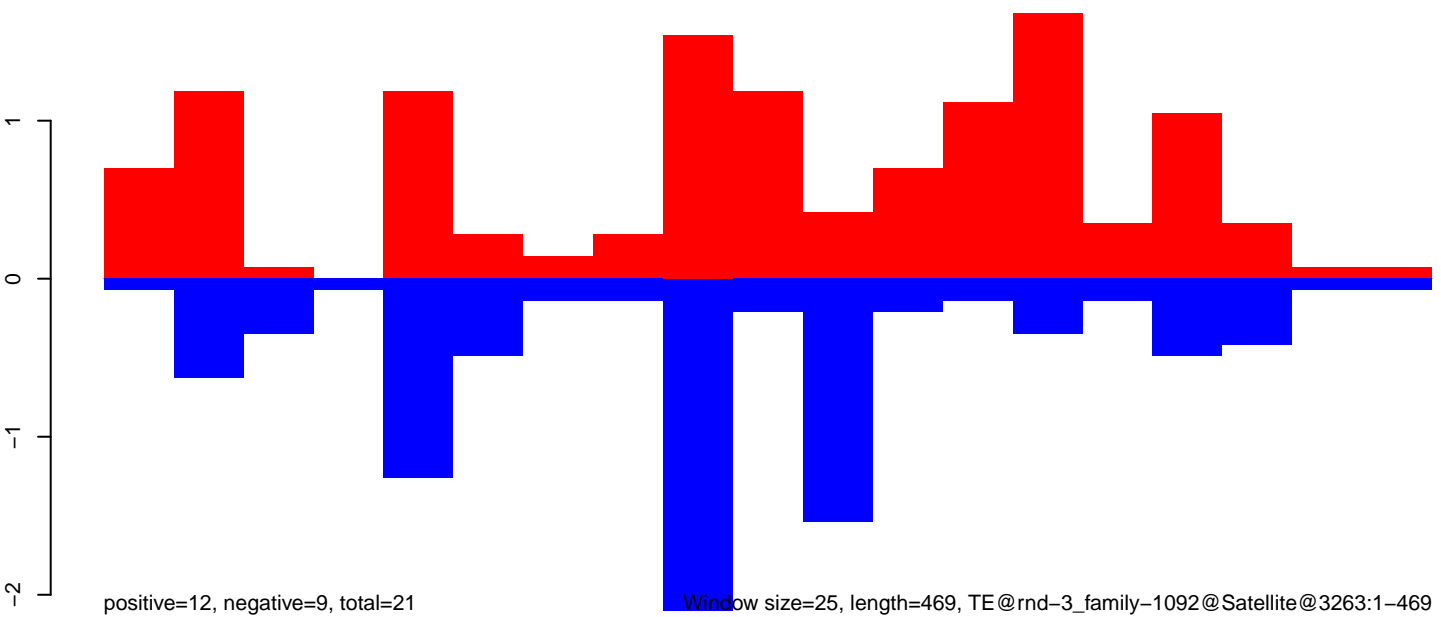
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

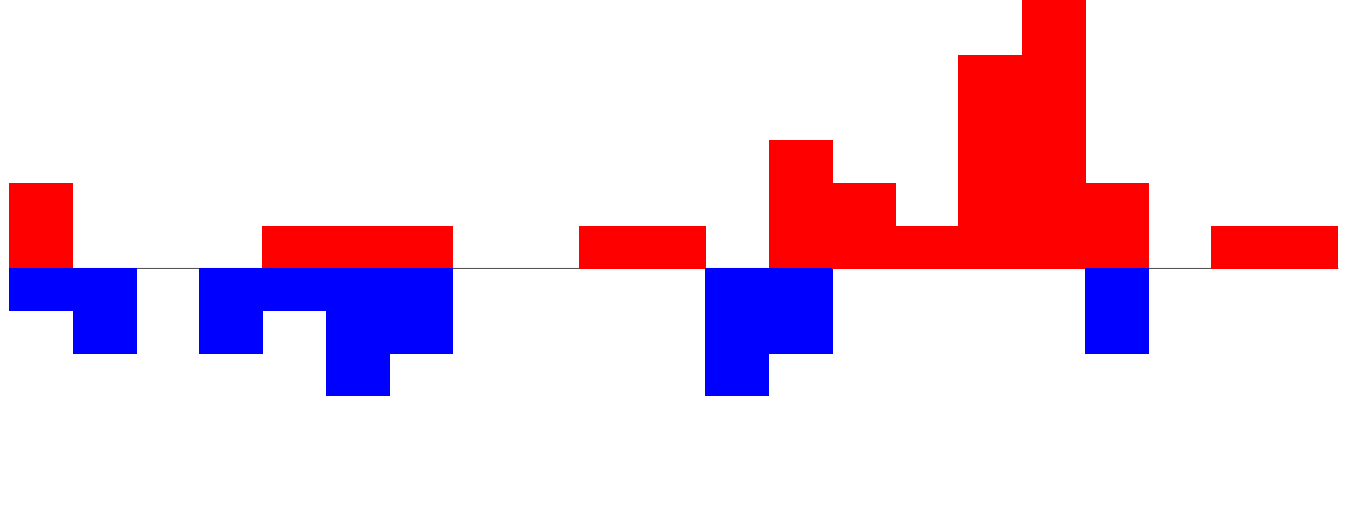


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CuQuin_Mcarc_NL.18_23.rep

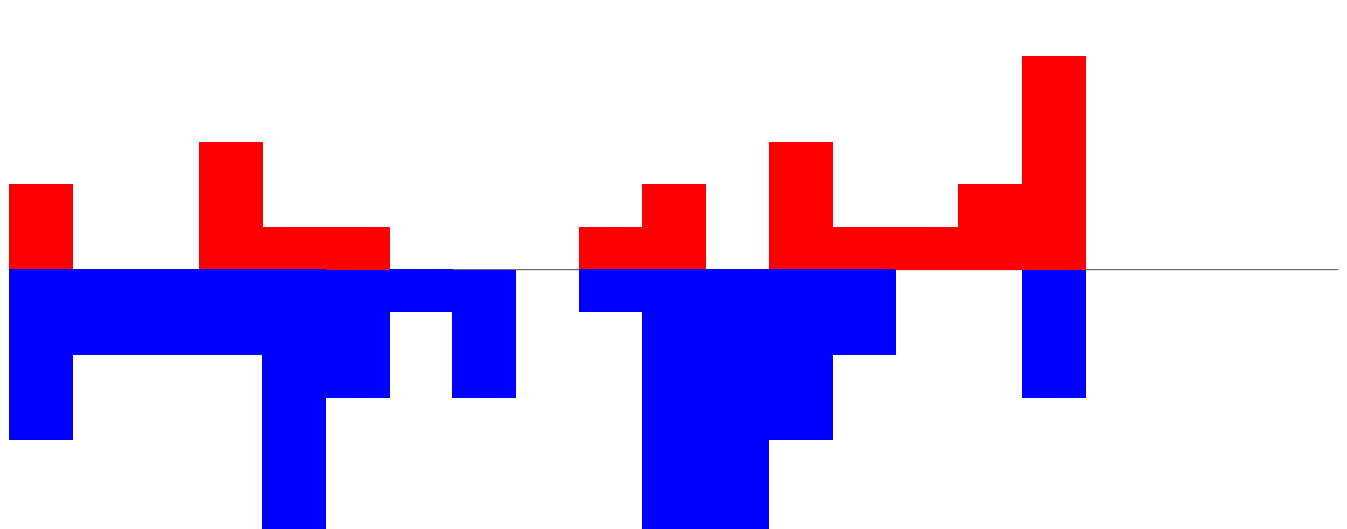
0.4
0.2
0.0
-0.2
-0.4



positive=2, negative=1, total=3

CuQuin_Mcarc_NL.24_35.rep

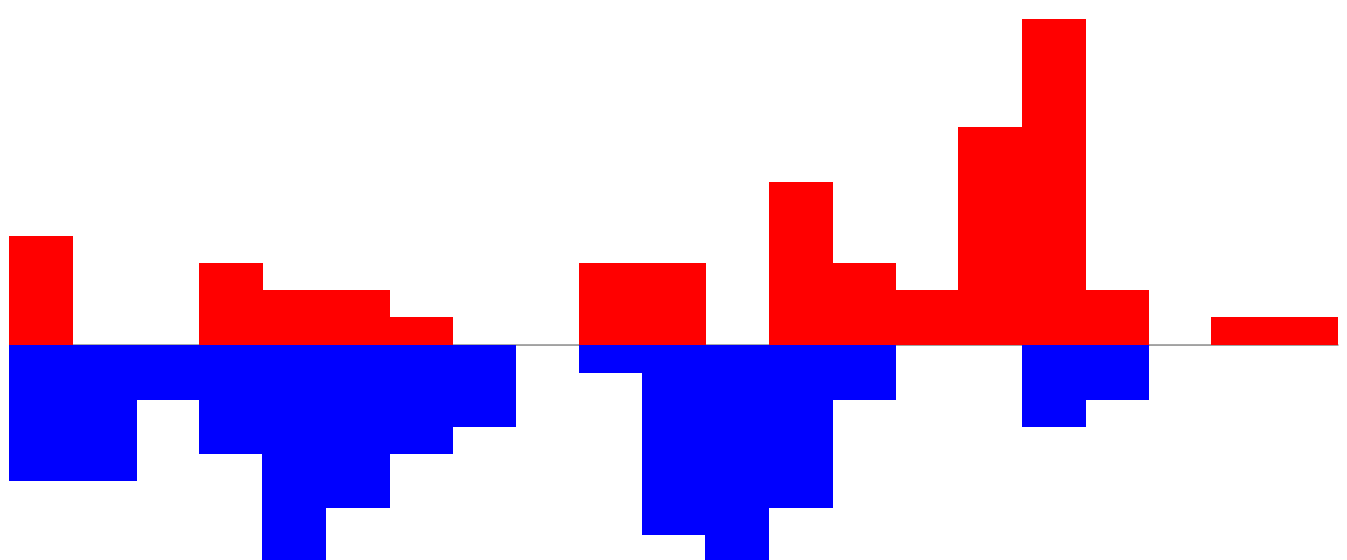
0.4
0.2
0.0
-0.2
-0.4



positive=2, negative=3, total=5

CuQuin_Mcarc_NL.rep

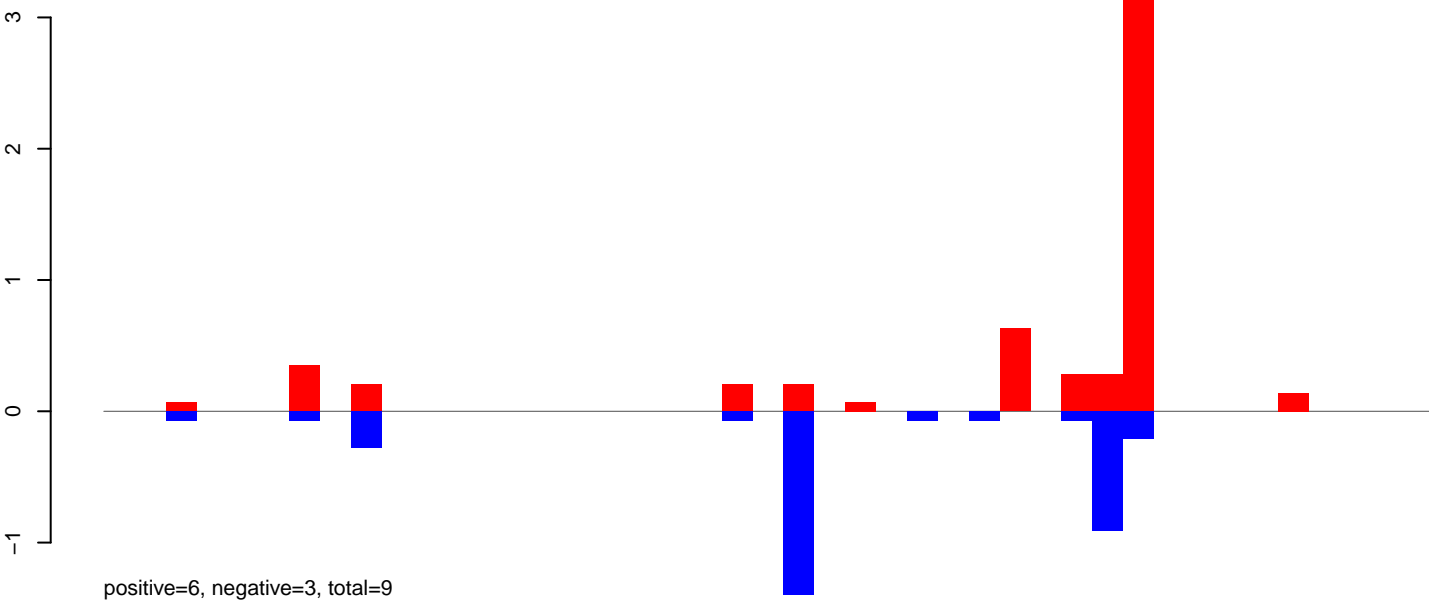
0.5
0.0
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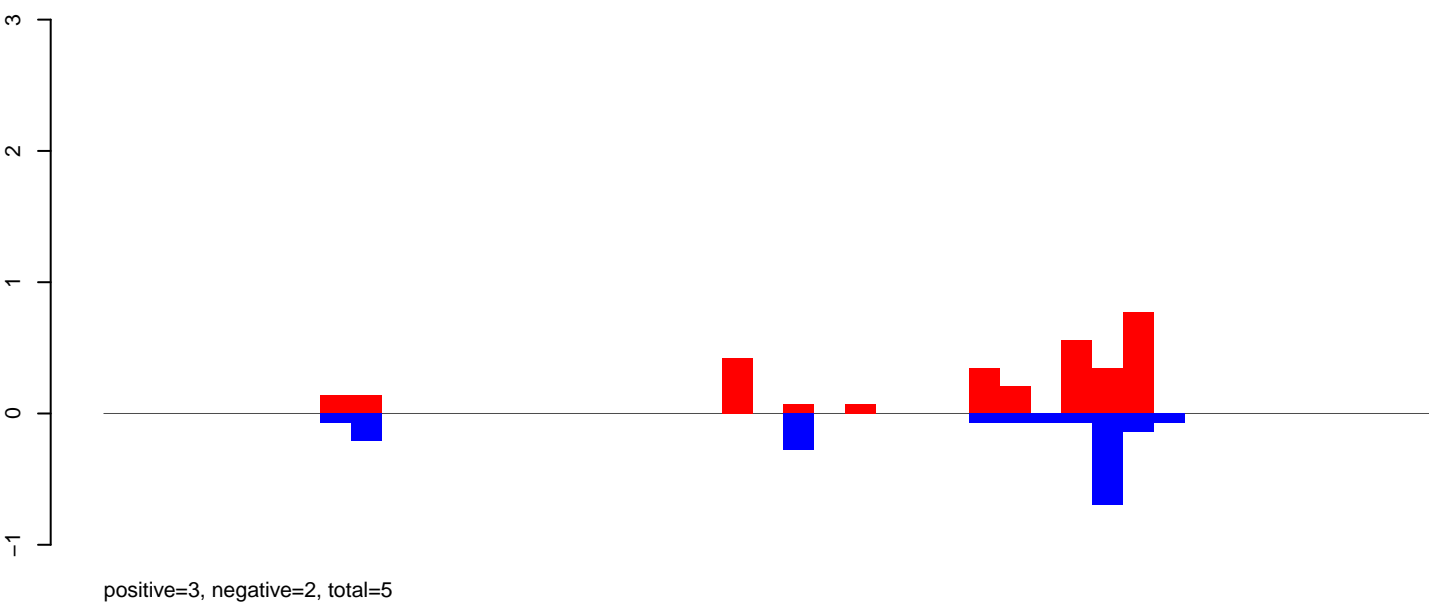
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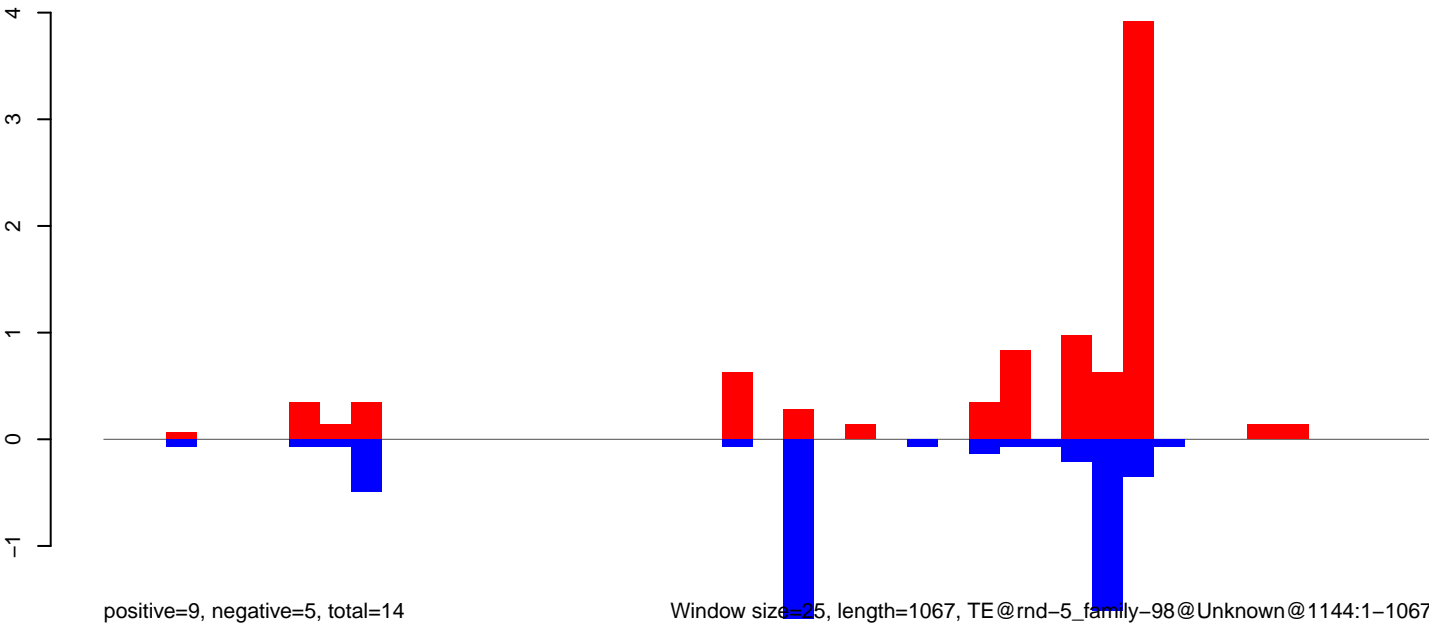
CuQuin_Mcarc_NL.18_23.rep



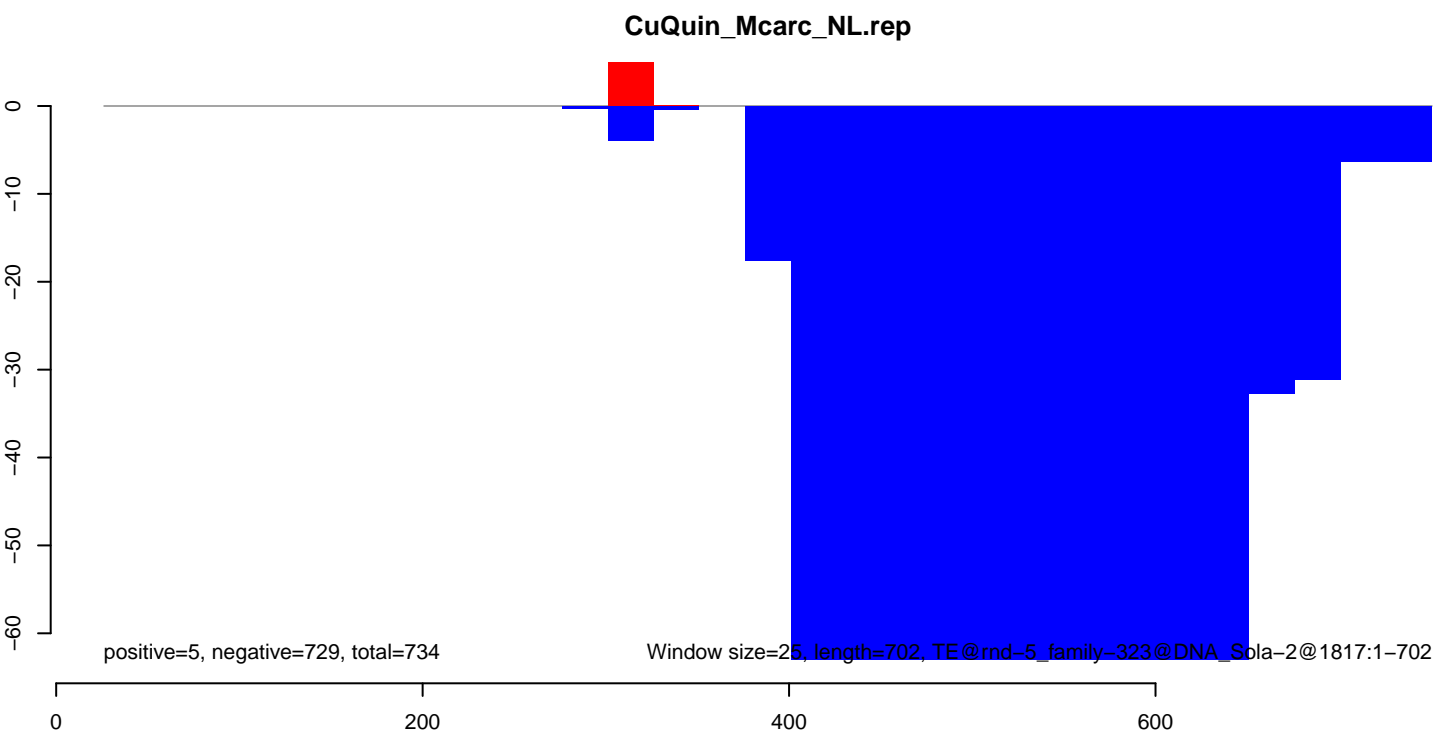
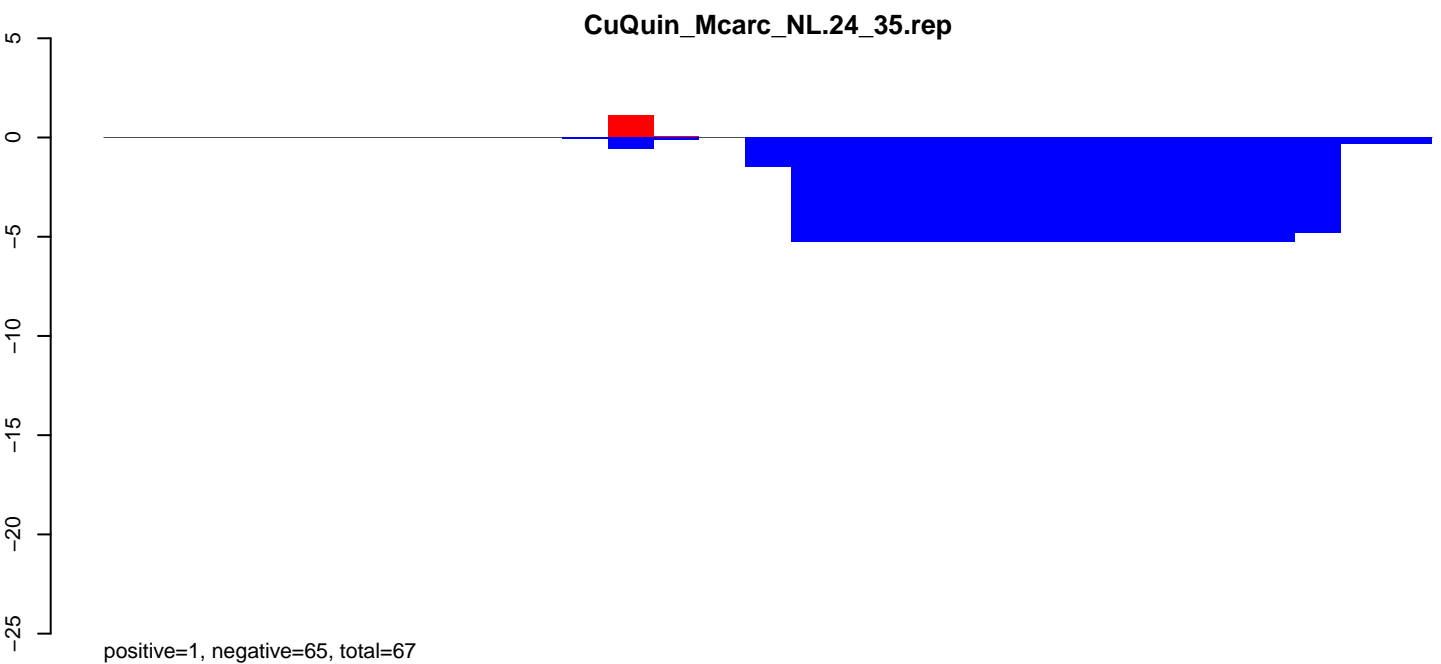
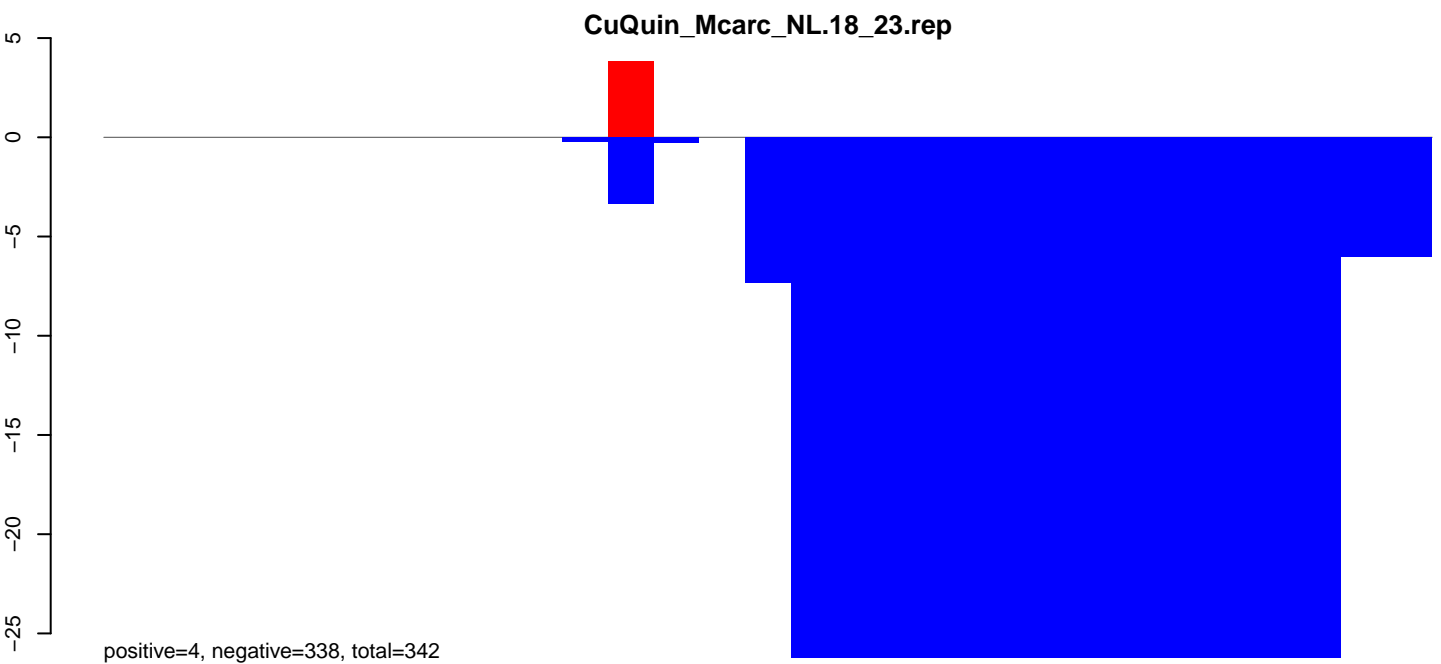
CuQuin_Mcarc_NL.24_35.rep



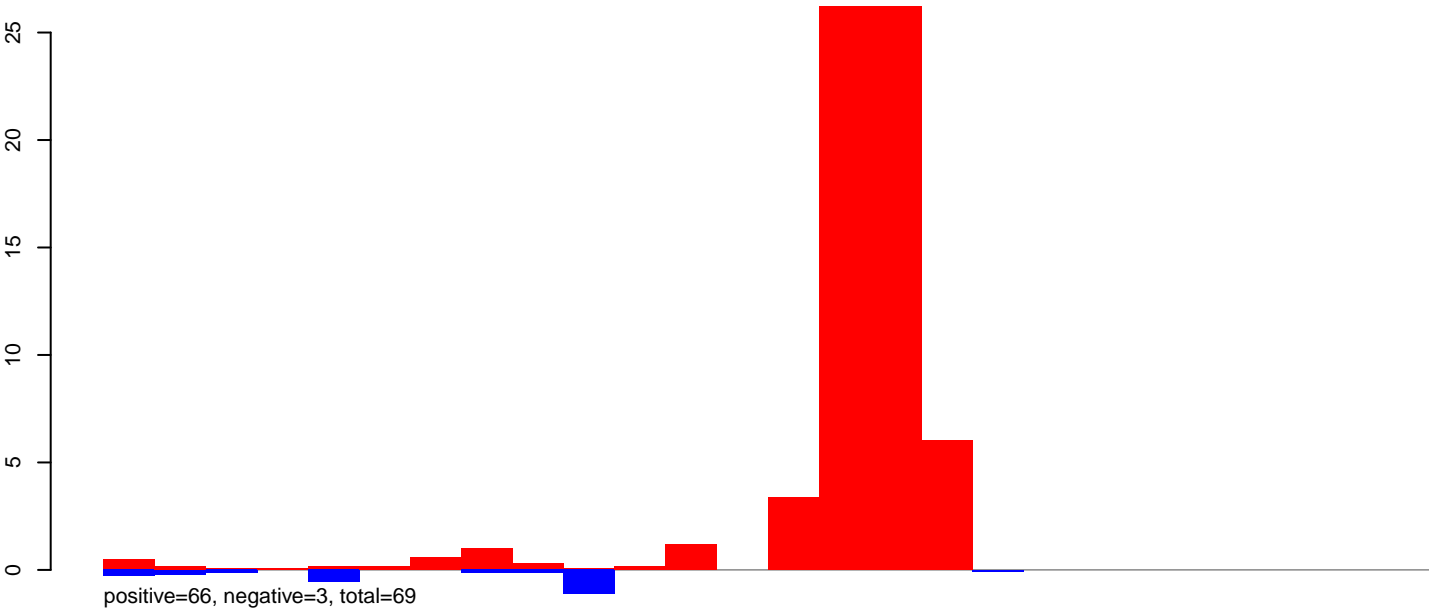
CuQuin_Mcarc_NL.rep



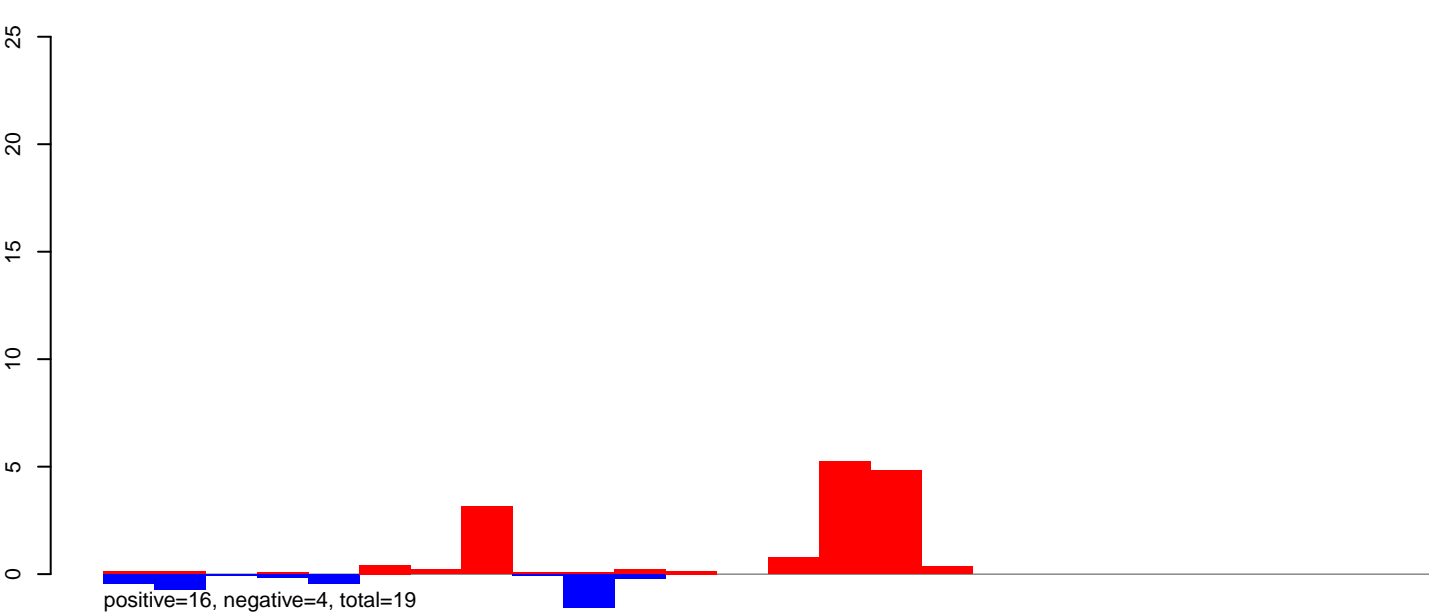
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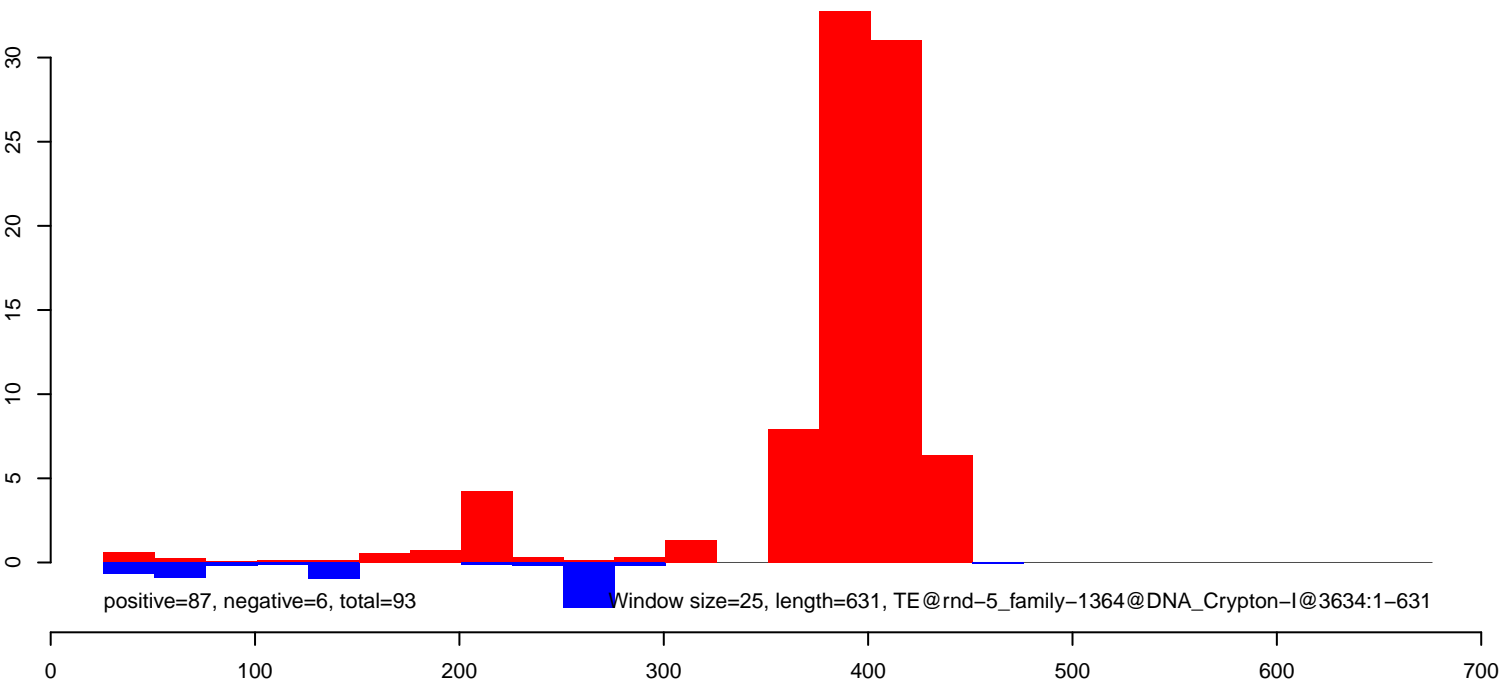
CuQuin_Mcarc_NL.18_23.rep



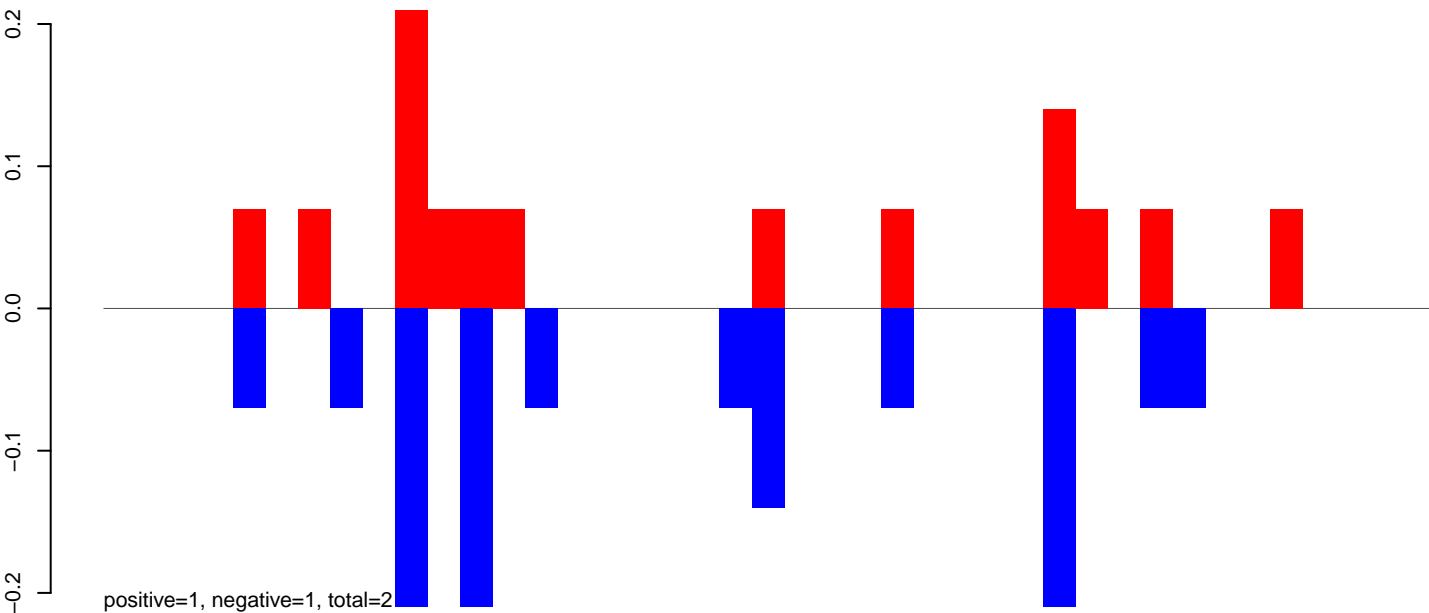
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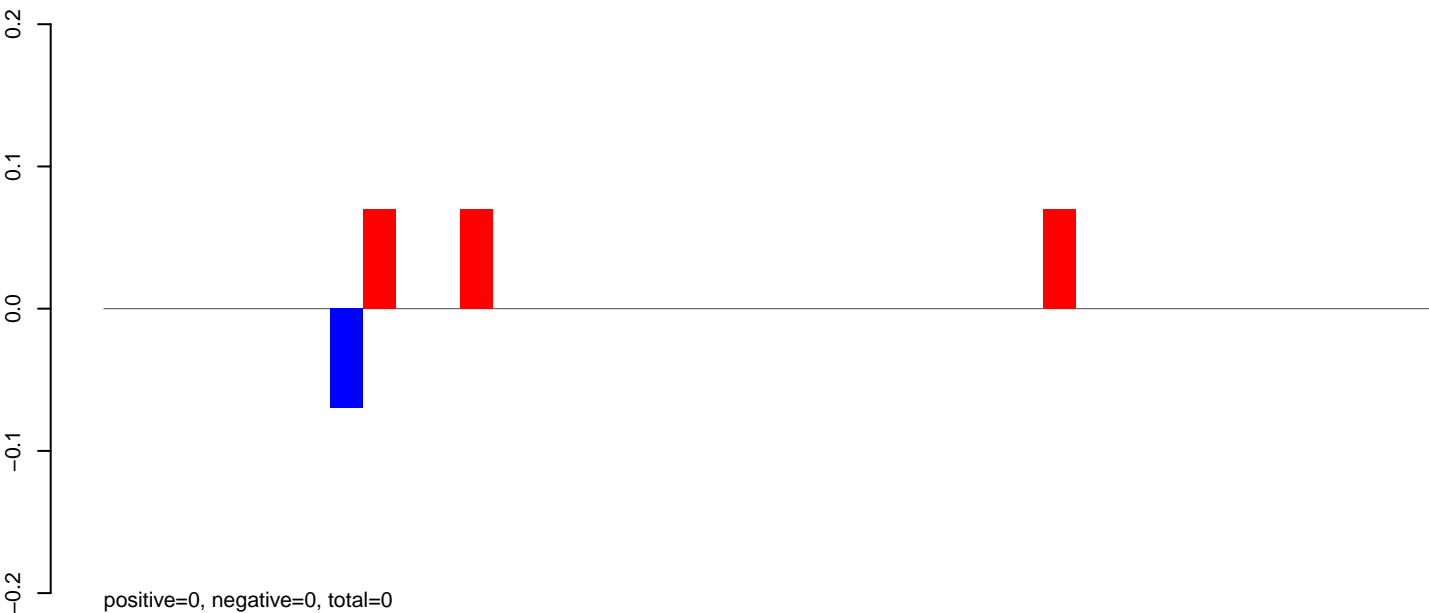
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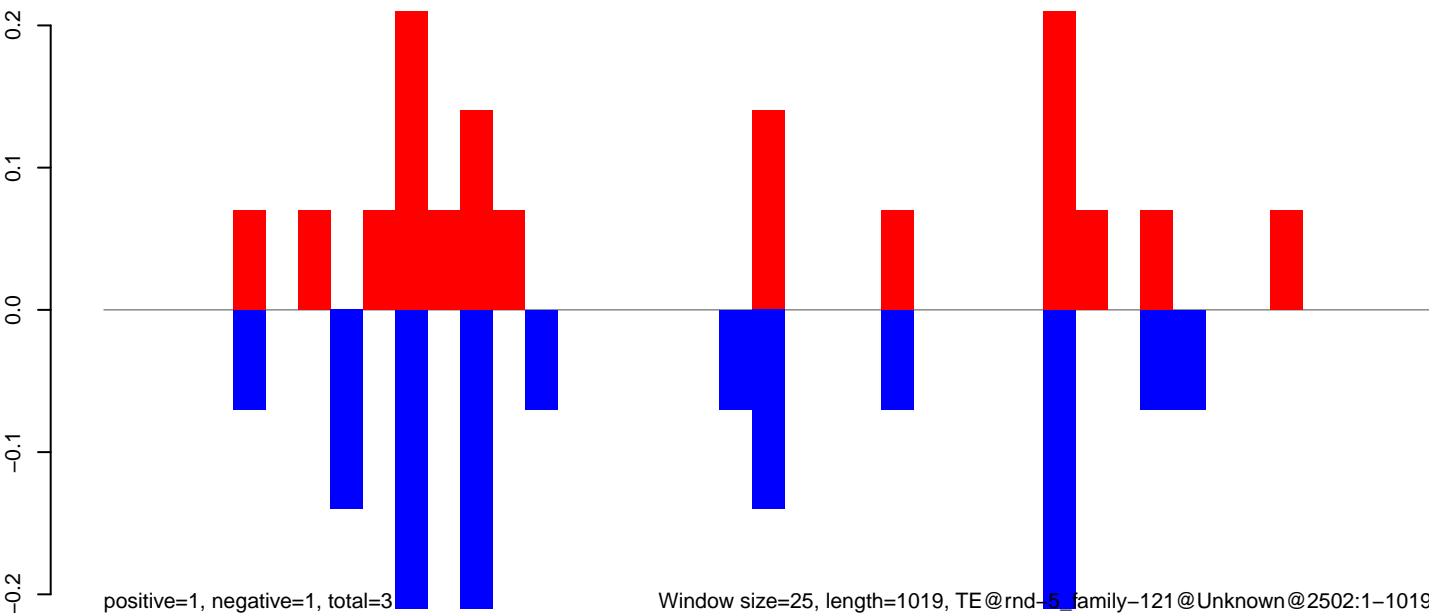
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CuQuin_Mcarc_NL.24_35.rep

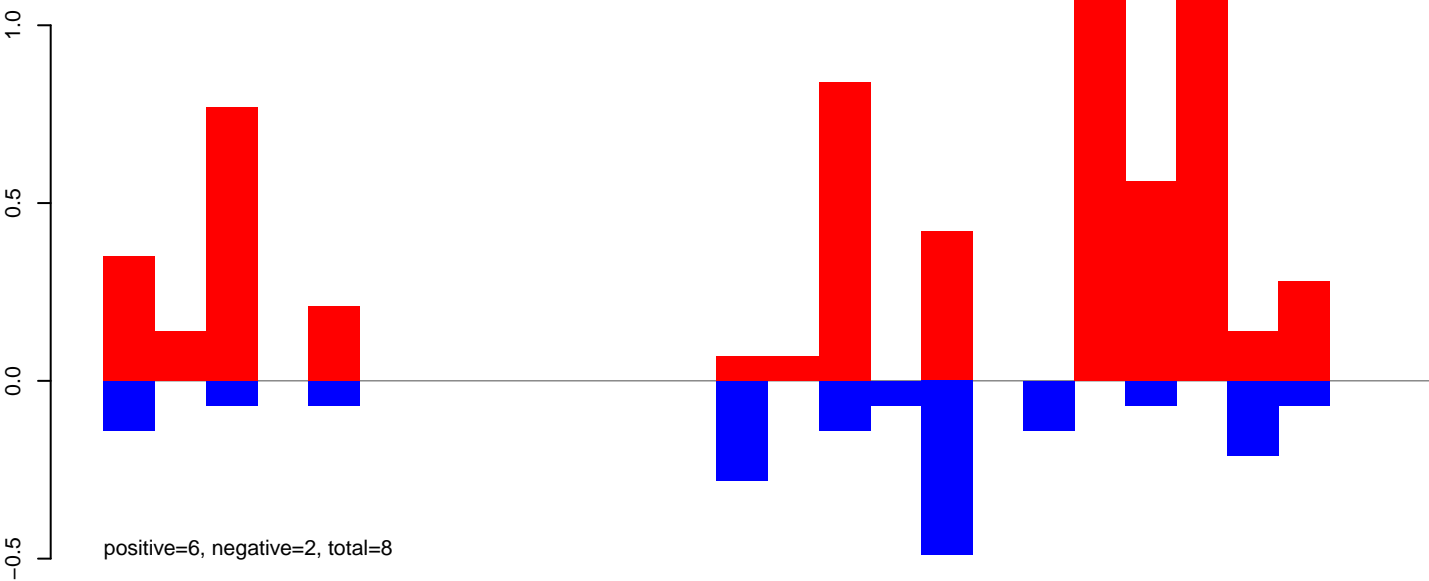


CuQuin_Mcarc_NL.rep

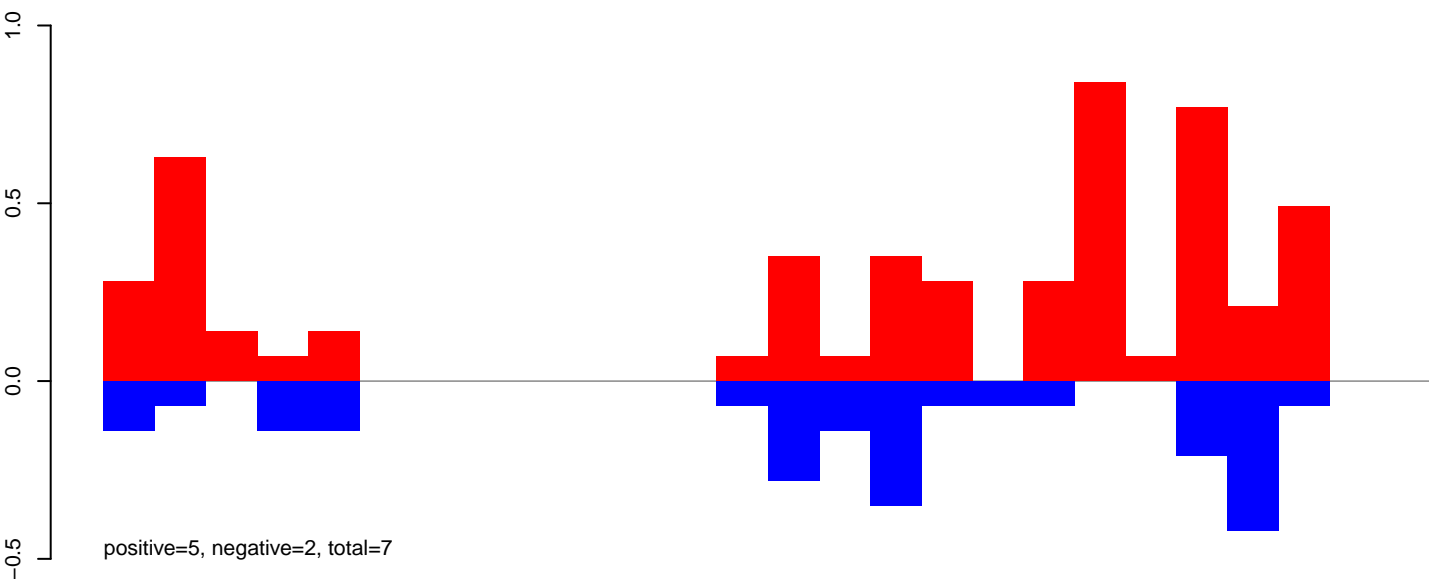


Window size=25, length=1019, TE@rnd-5_family-121@Unknown@2502:1-1019

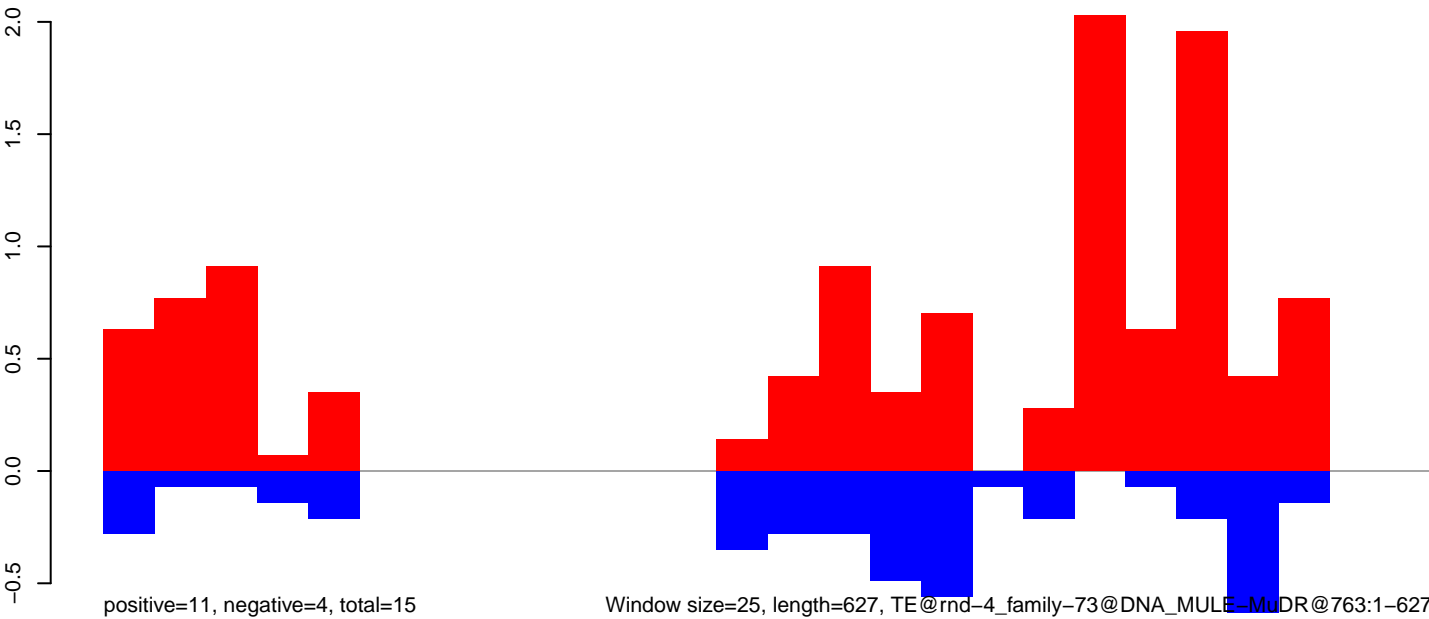
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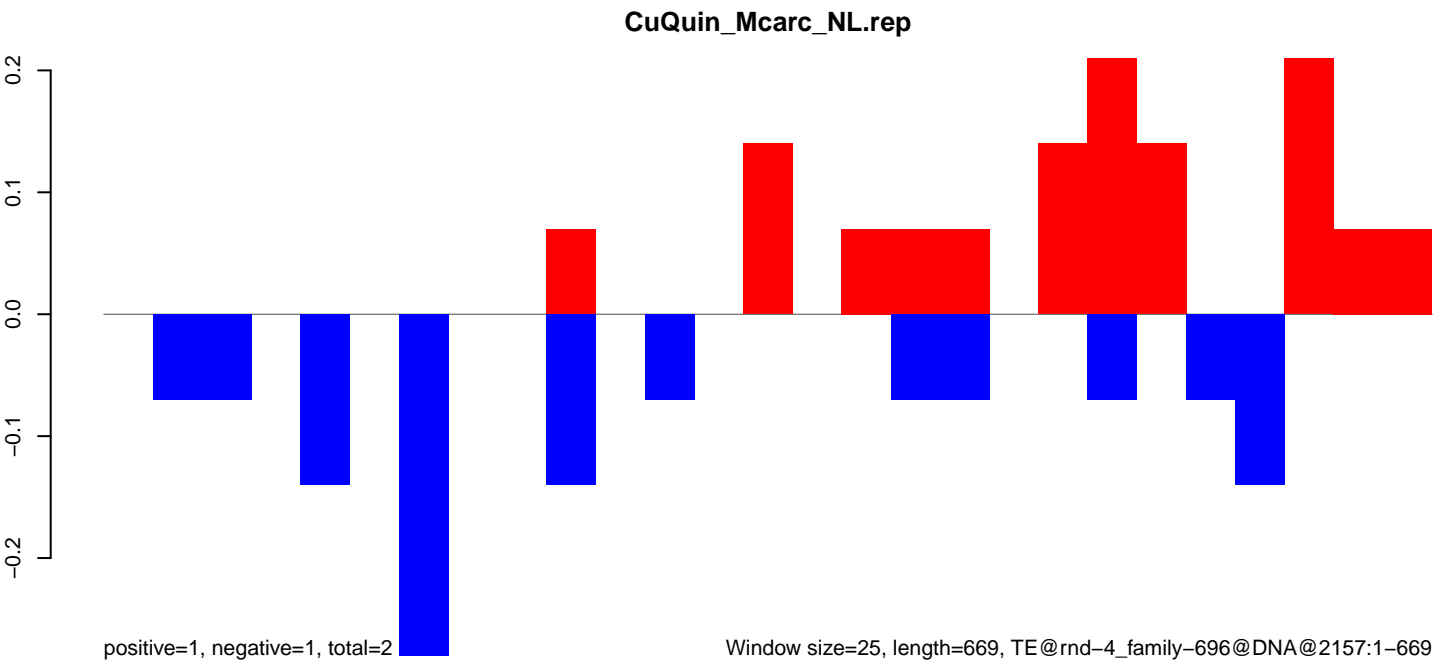
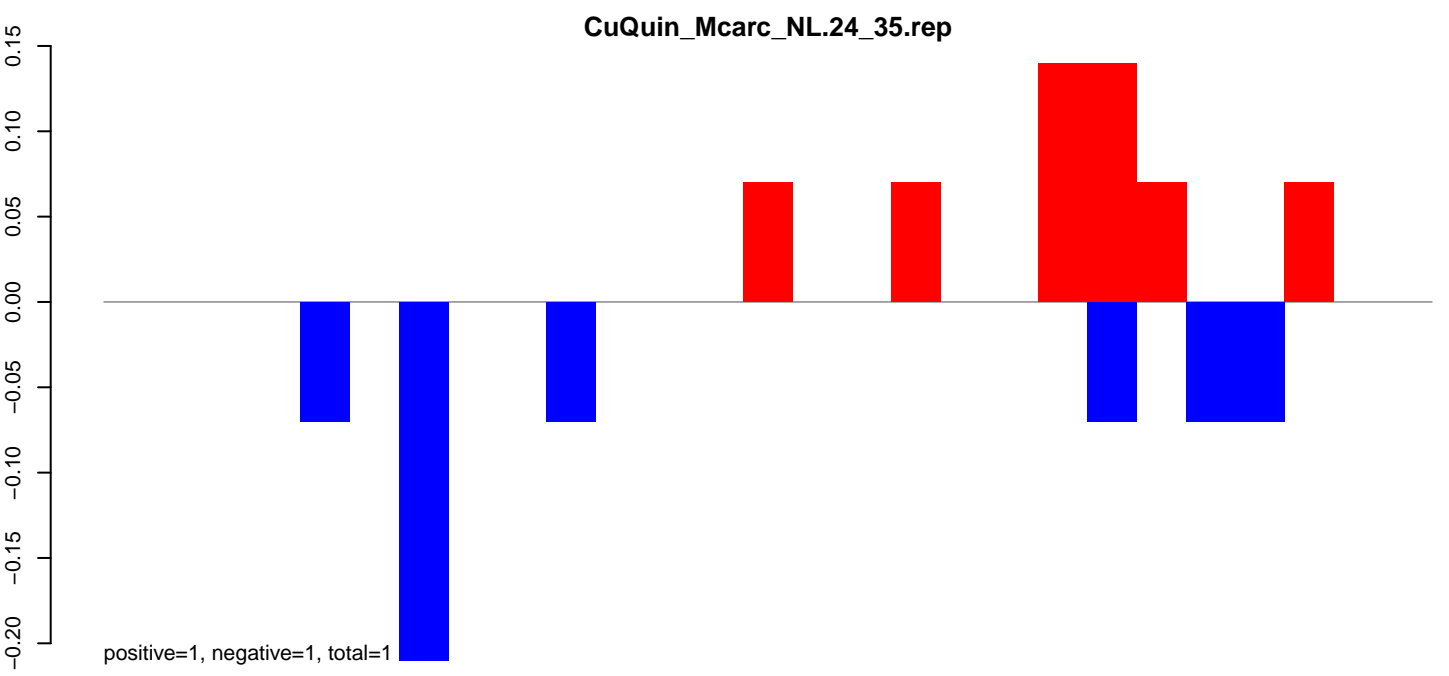
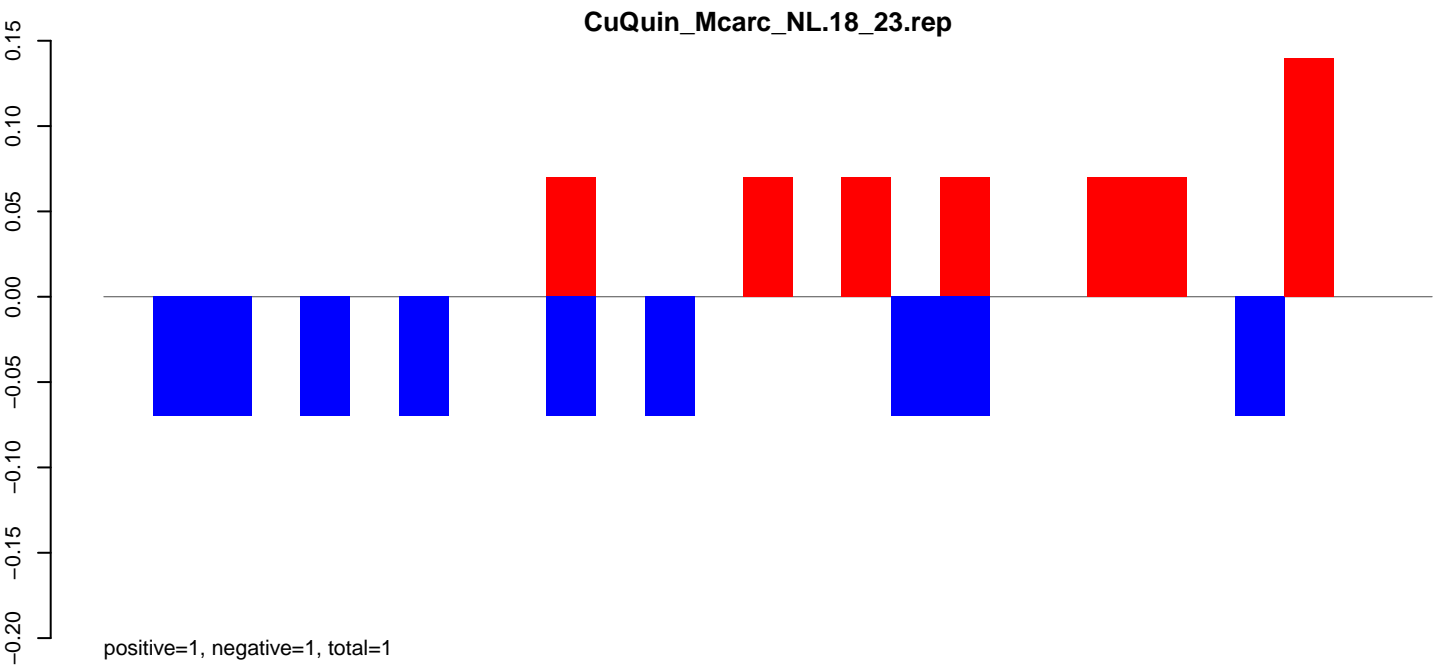


CuQuin_Mcarc_NL.24_35.rep

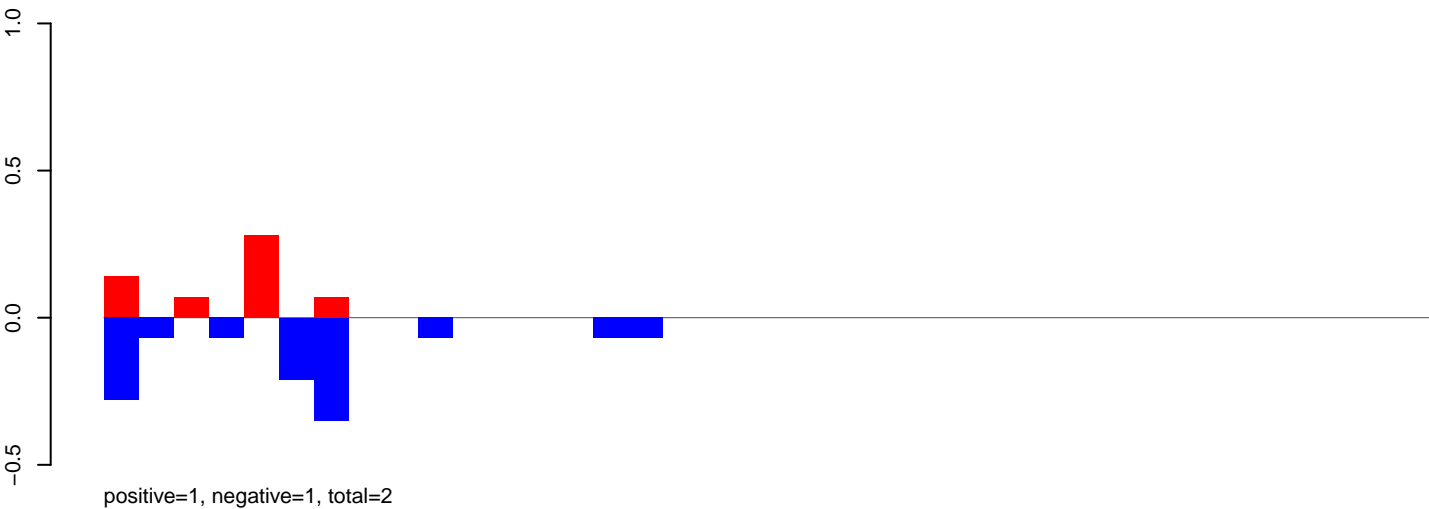


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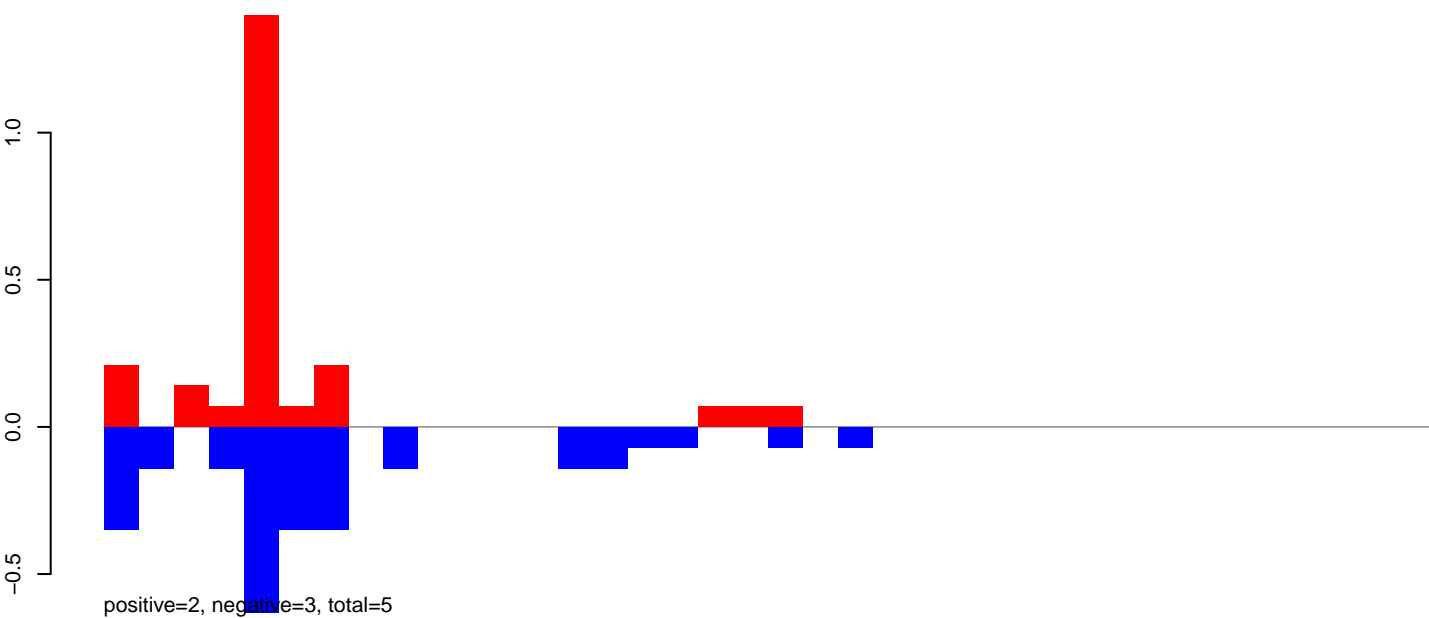




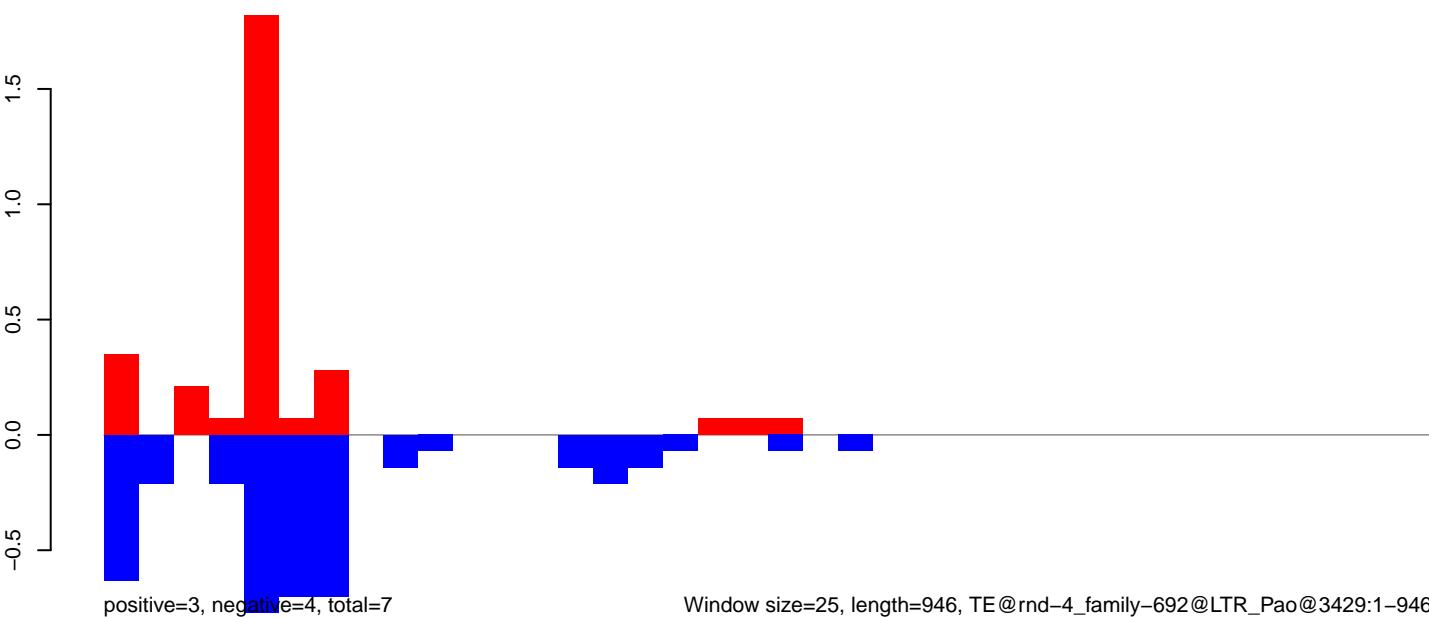
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CuQuin_Mcarc_NL.24_35.rep

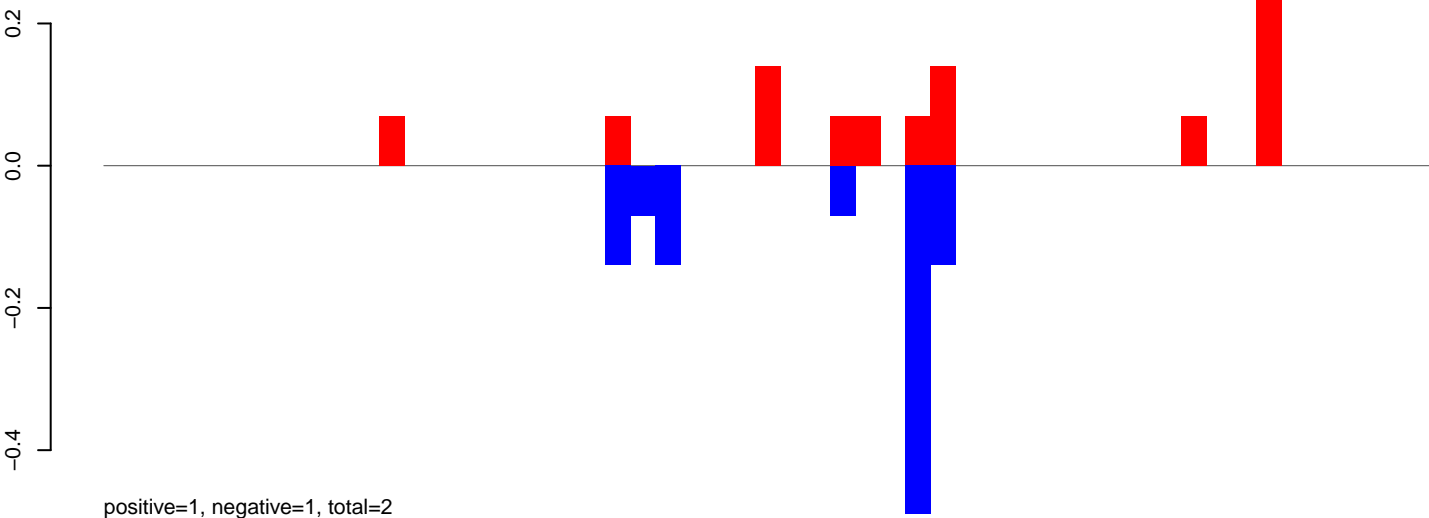


CuQuin_Mcarc_NL.rep

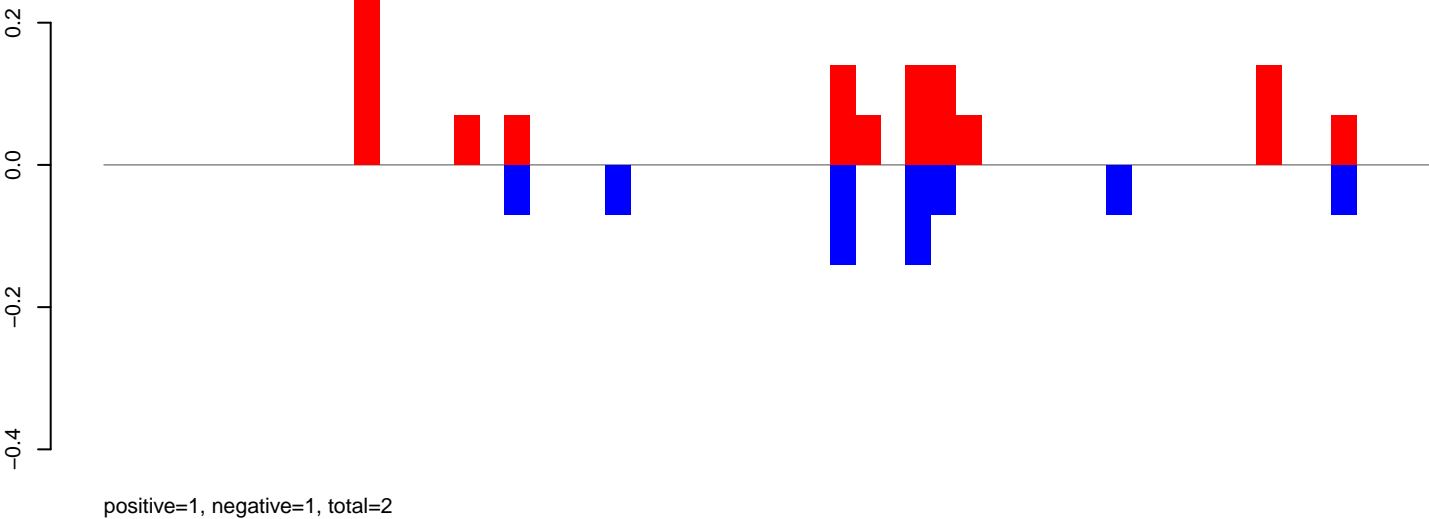


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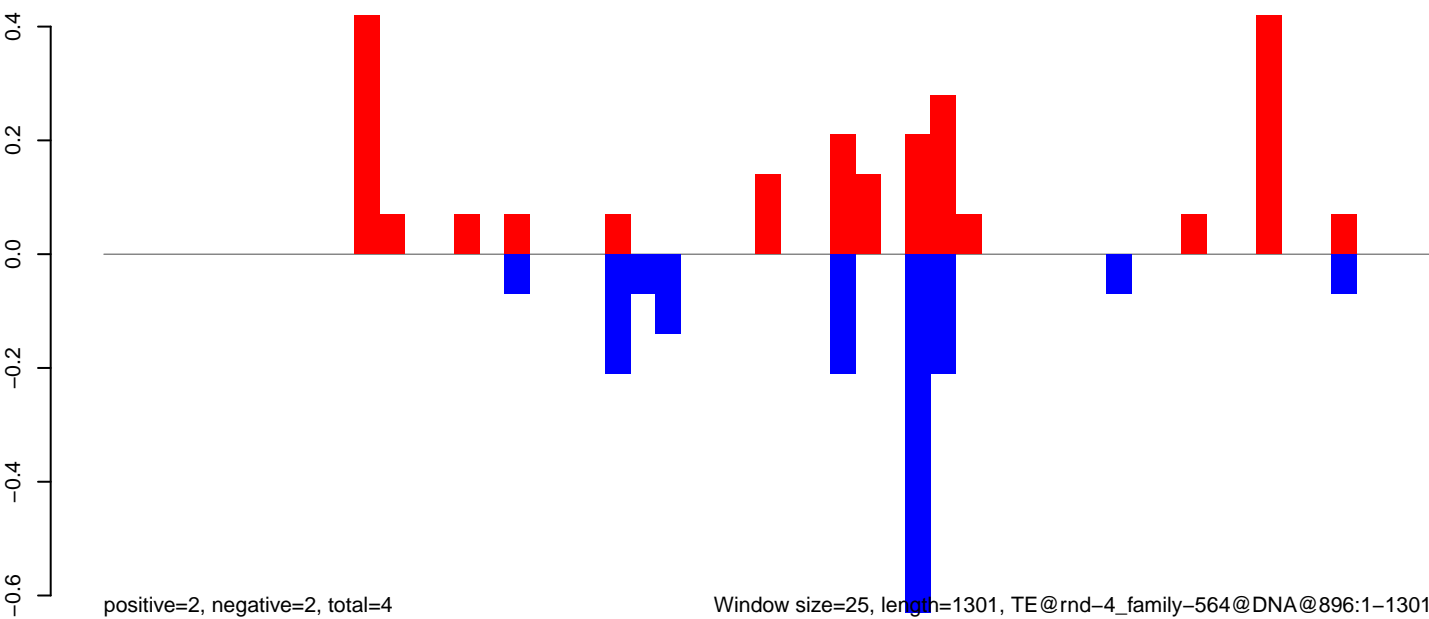
CuQuin_Mcarc_NL.18_23.rep



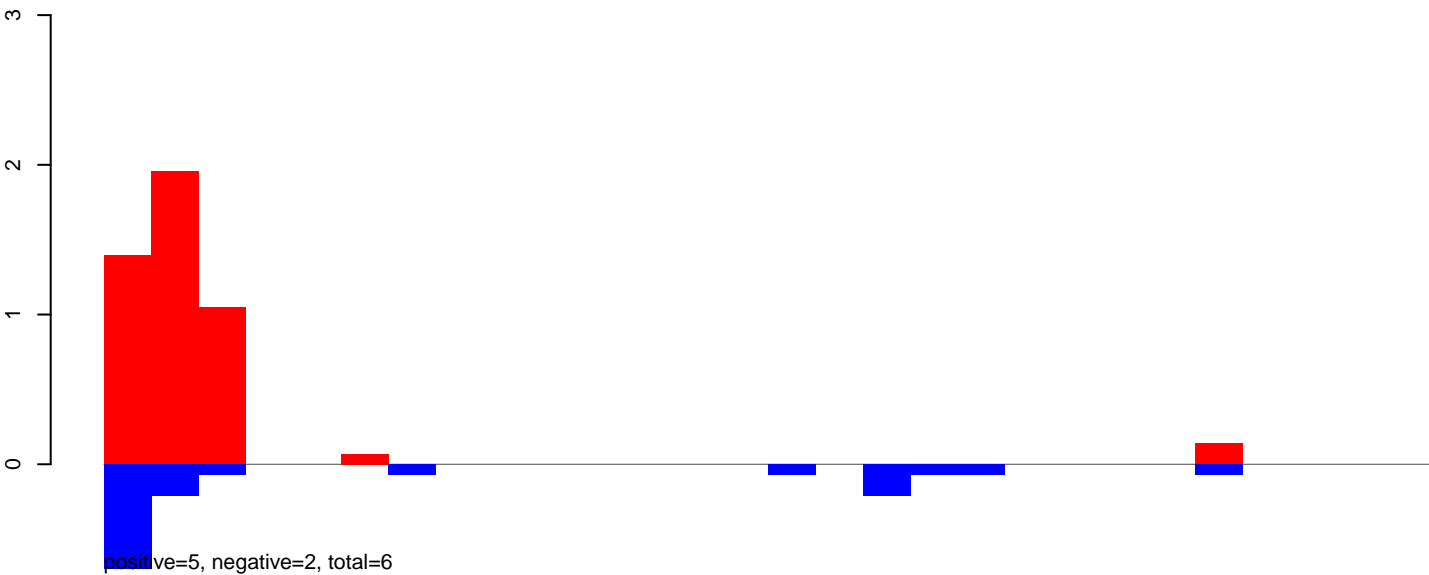
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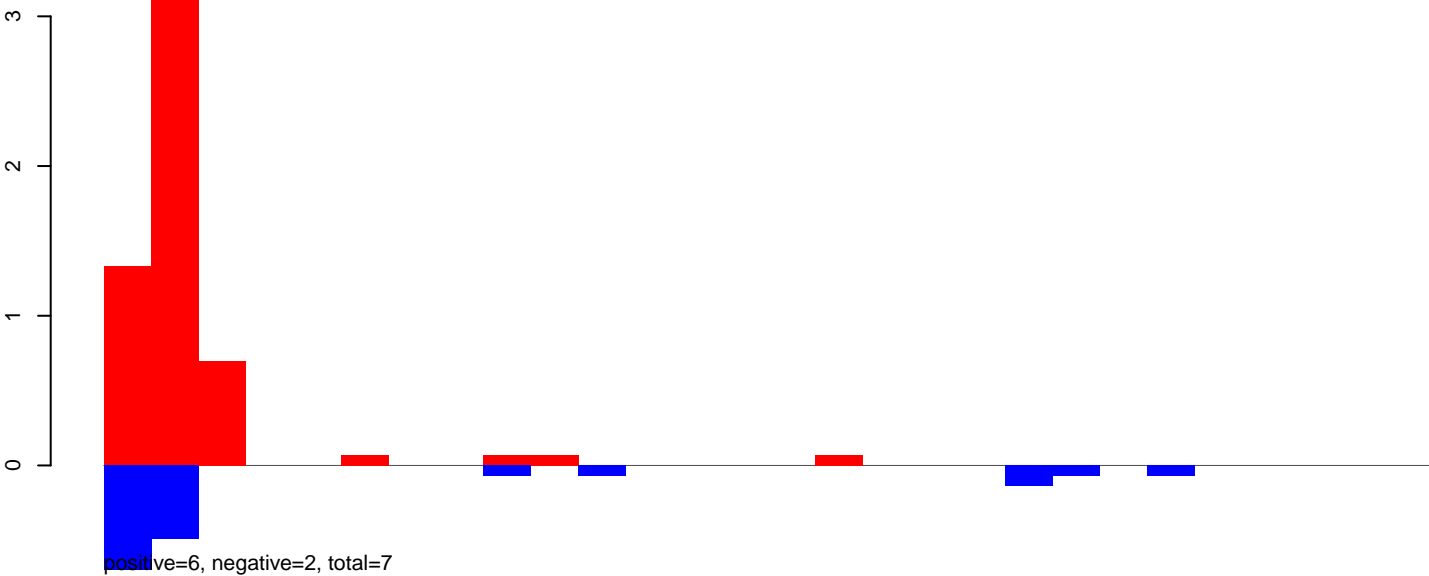
CuQuin_Mcarc_NL.rep



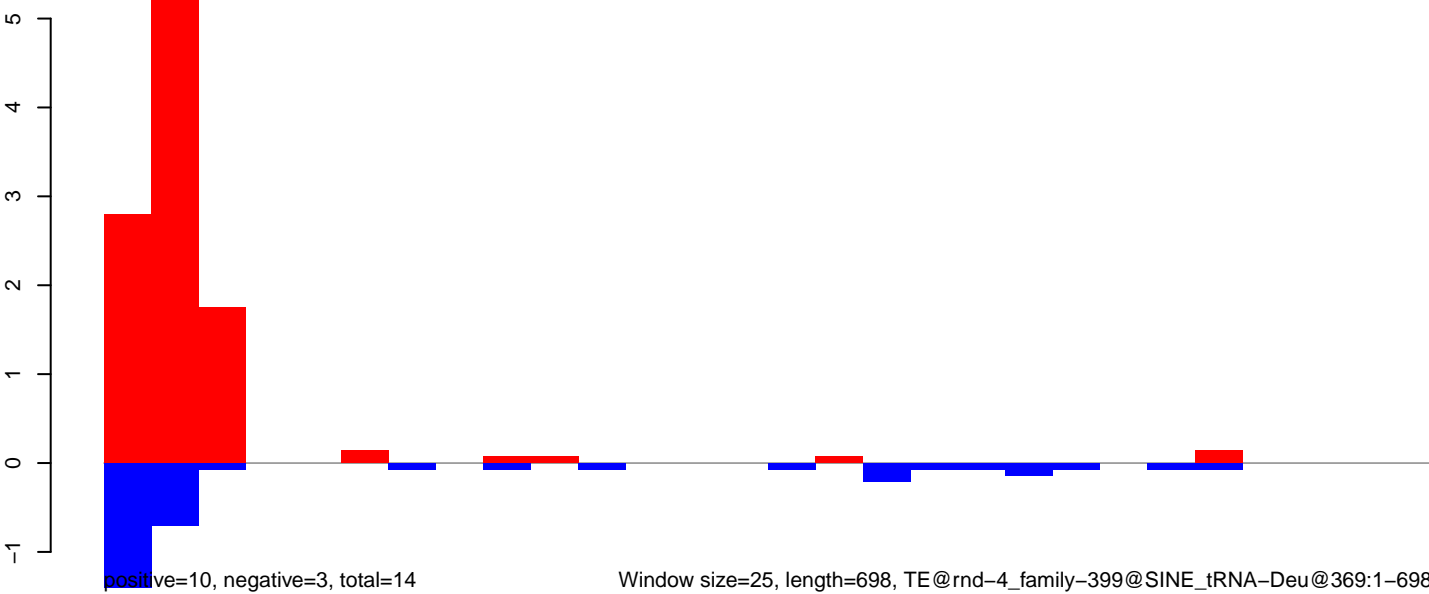
CuQuin_Mcarc_NL.18_23.rep



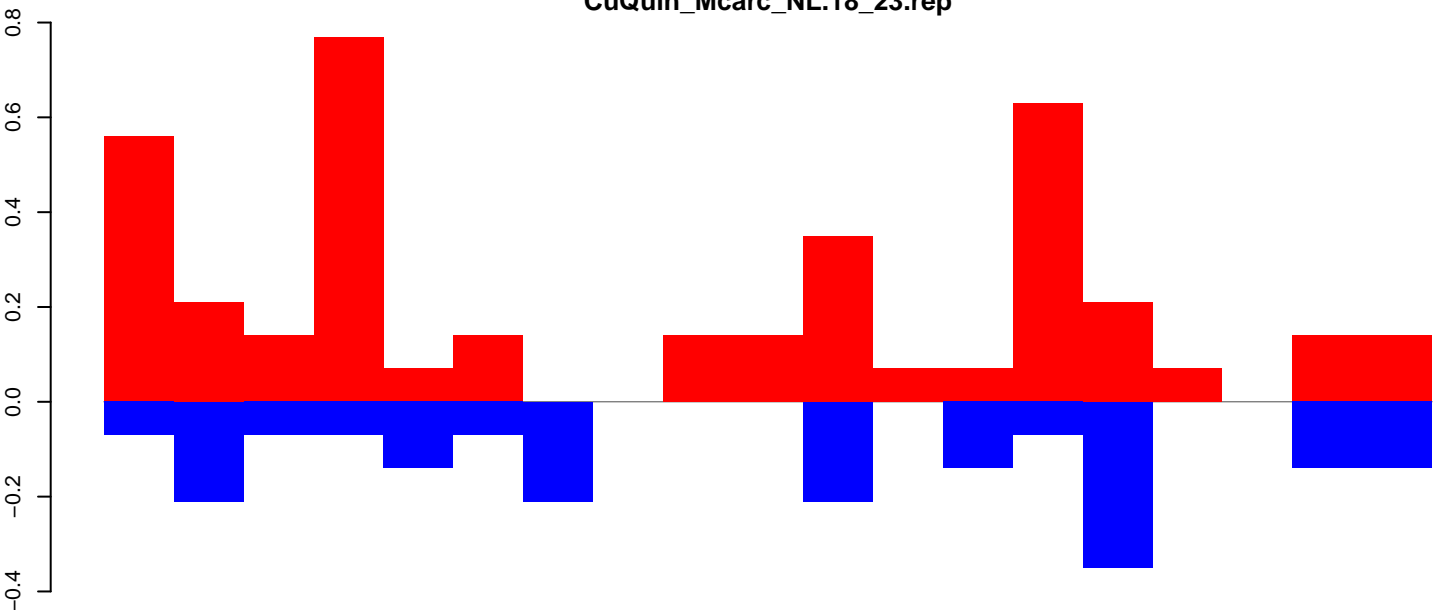
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CuQuin_Mcarc_NL.rep

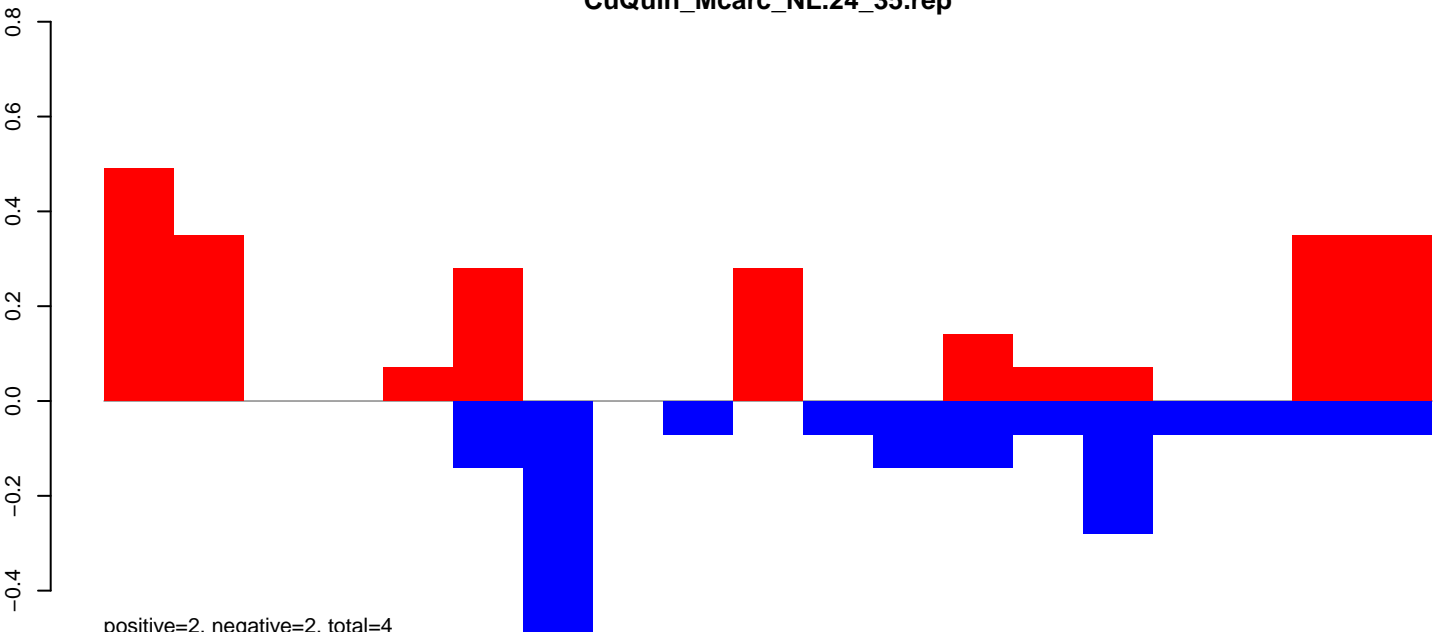


CuQuin_Mcarc_NL.18_23.rep



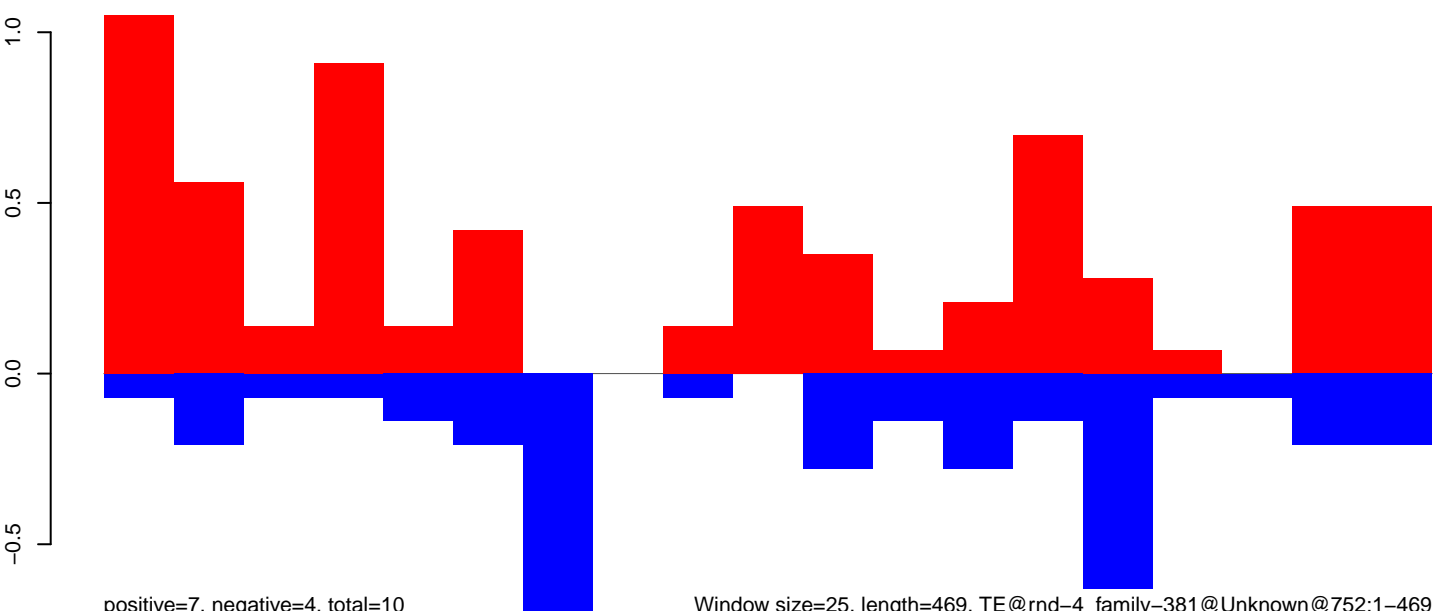
positive=4, negative=2, total=6

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=2, total=4

CuQuin_Mcarc_NL.rep

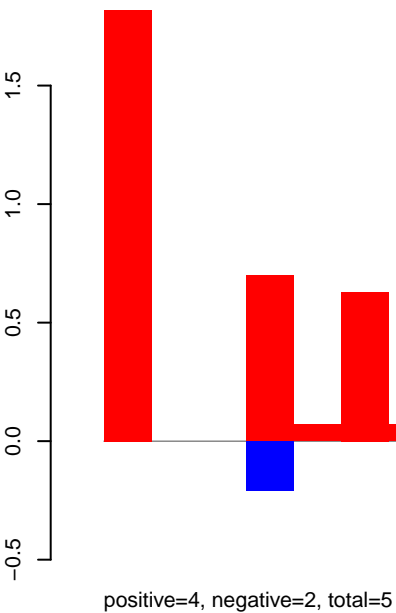


positive=7, negative=4, total=10

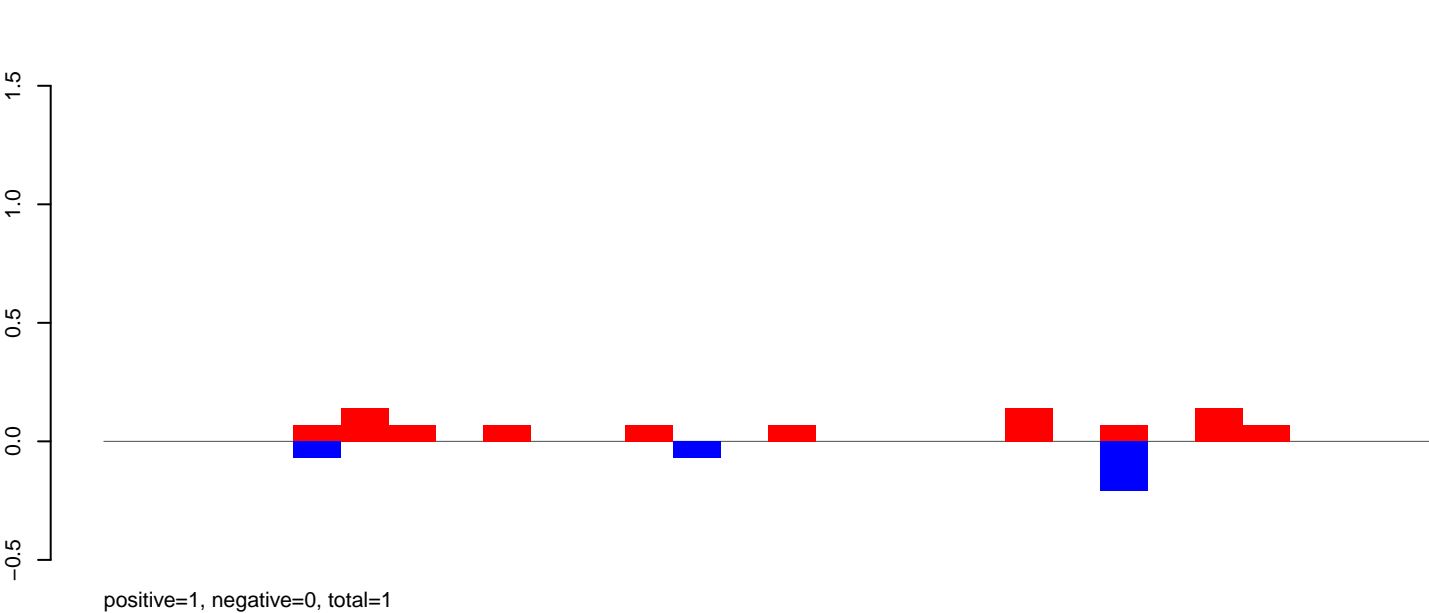
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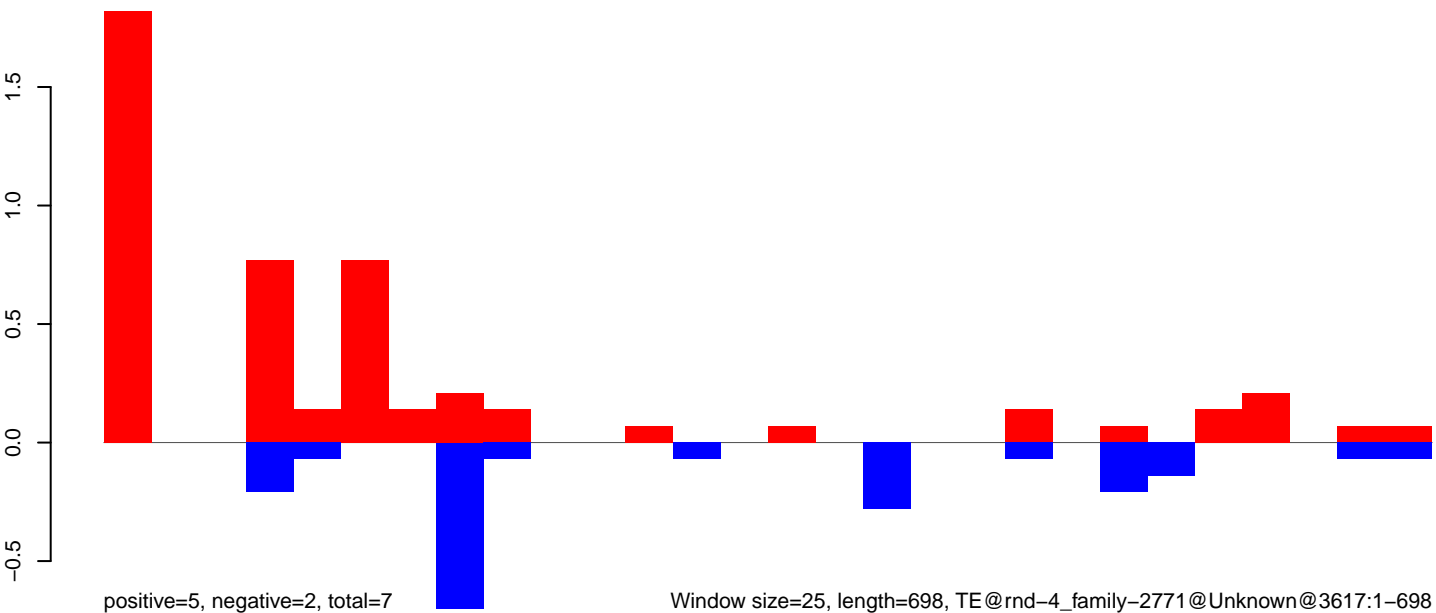
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CuQuin_Mcarc_NL.24_35.rep

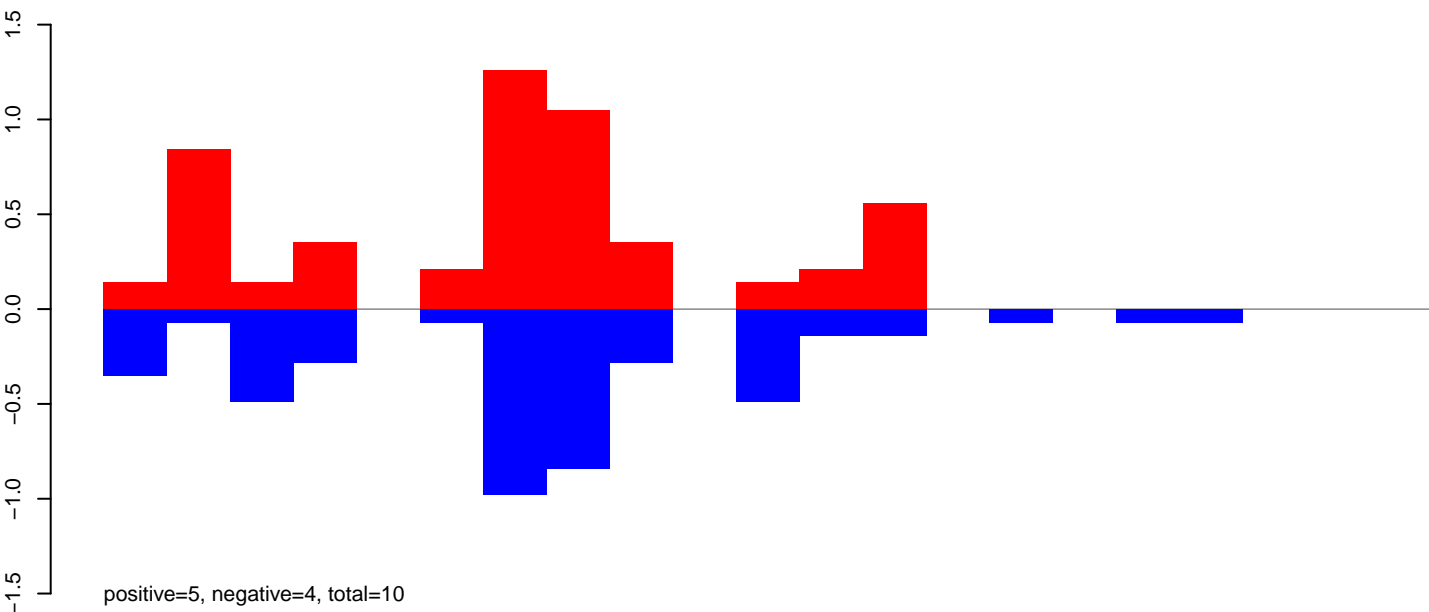


CuQuin_Mcarc_NL.rep

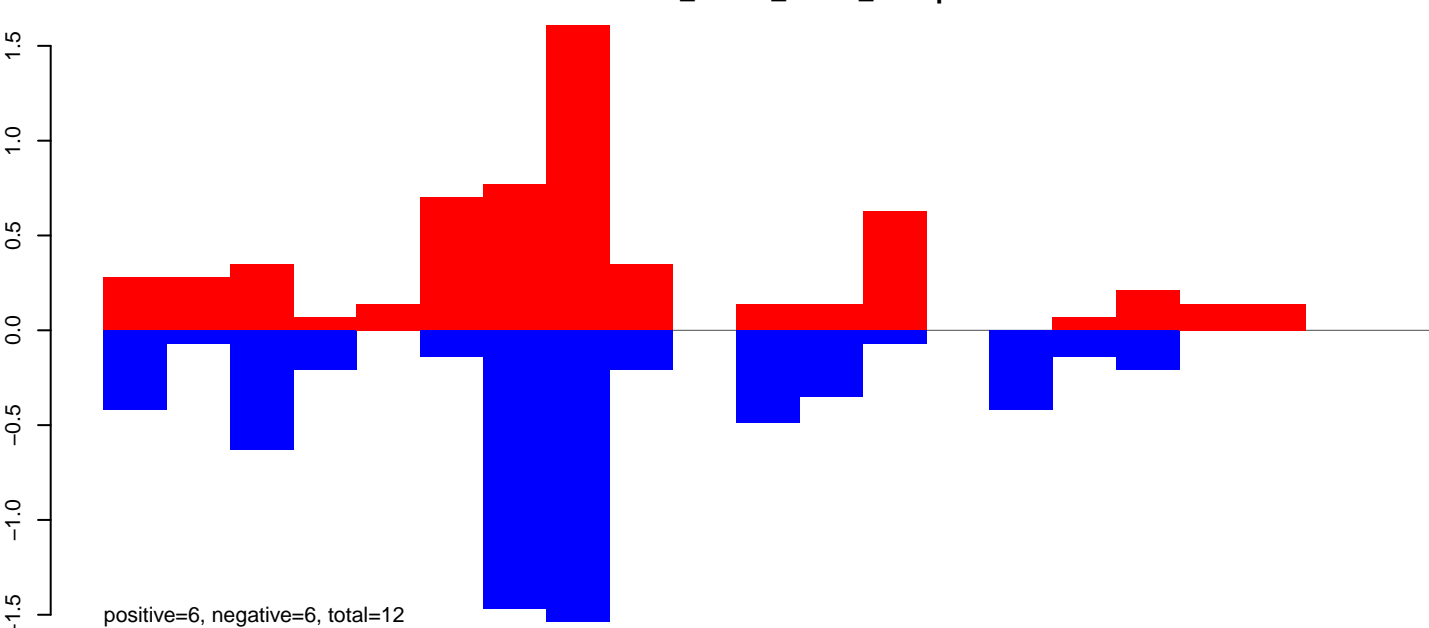


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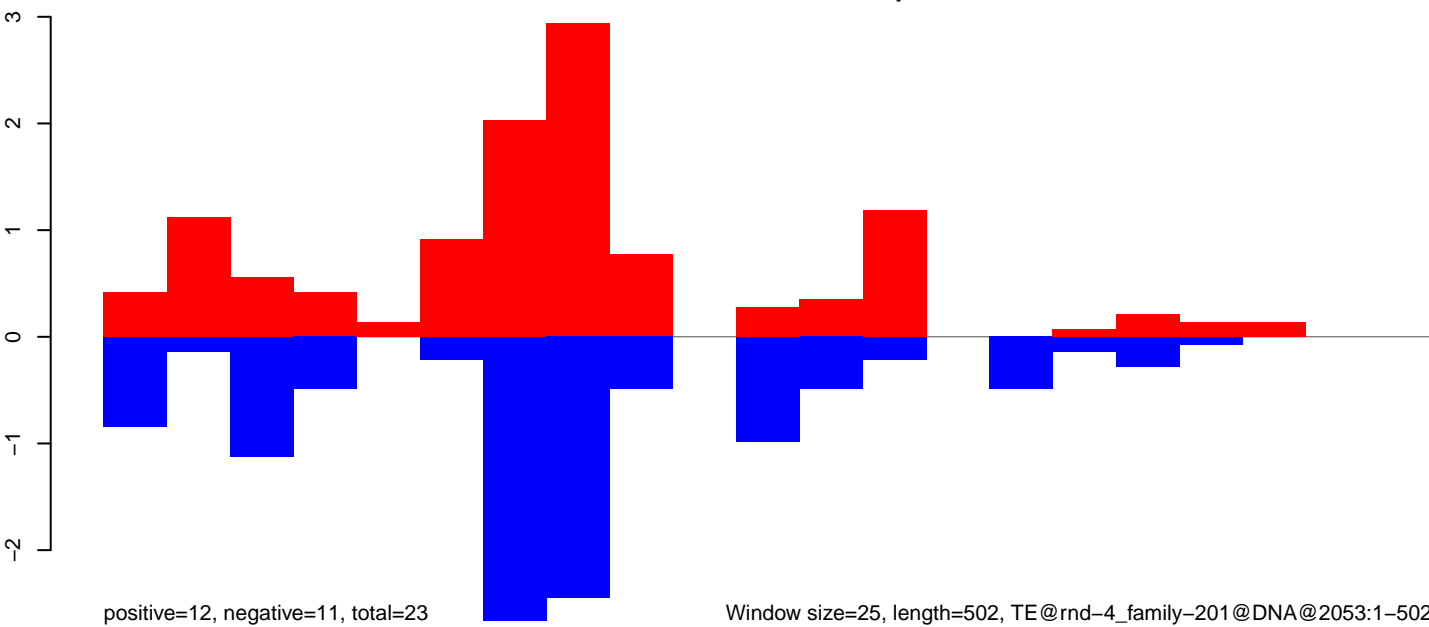
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

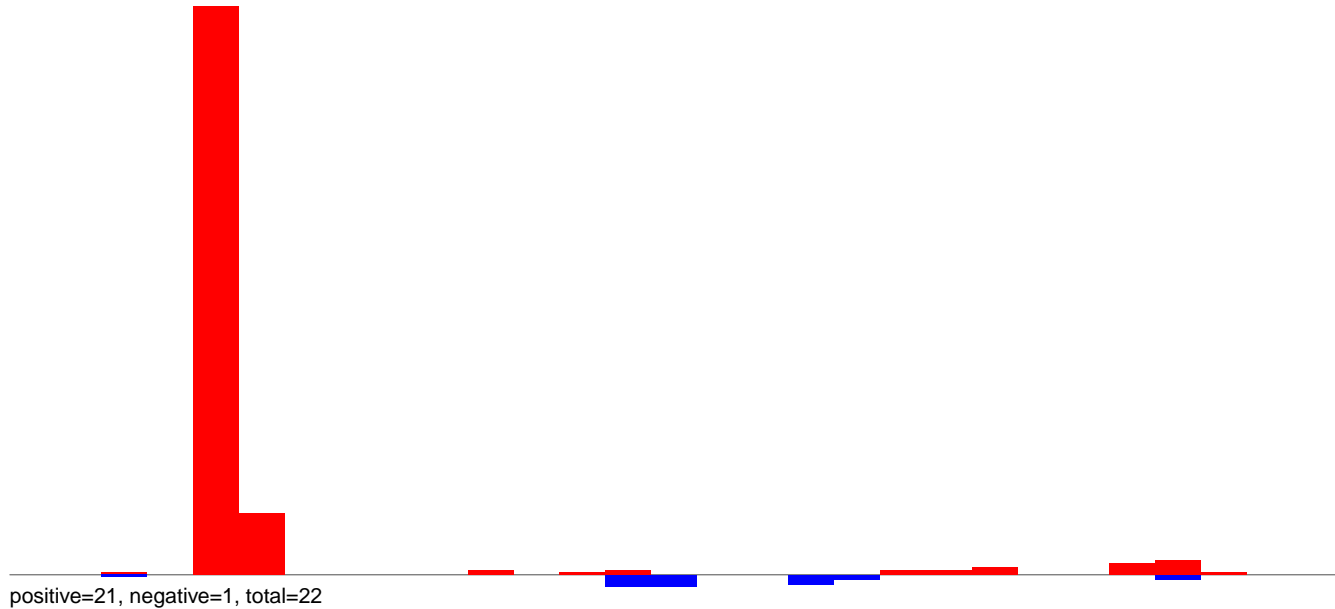


CuQuin_Mcarc_NL.rep

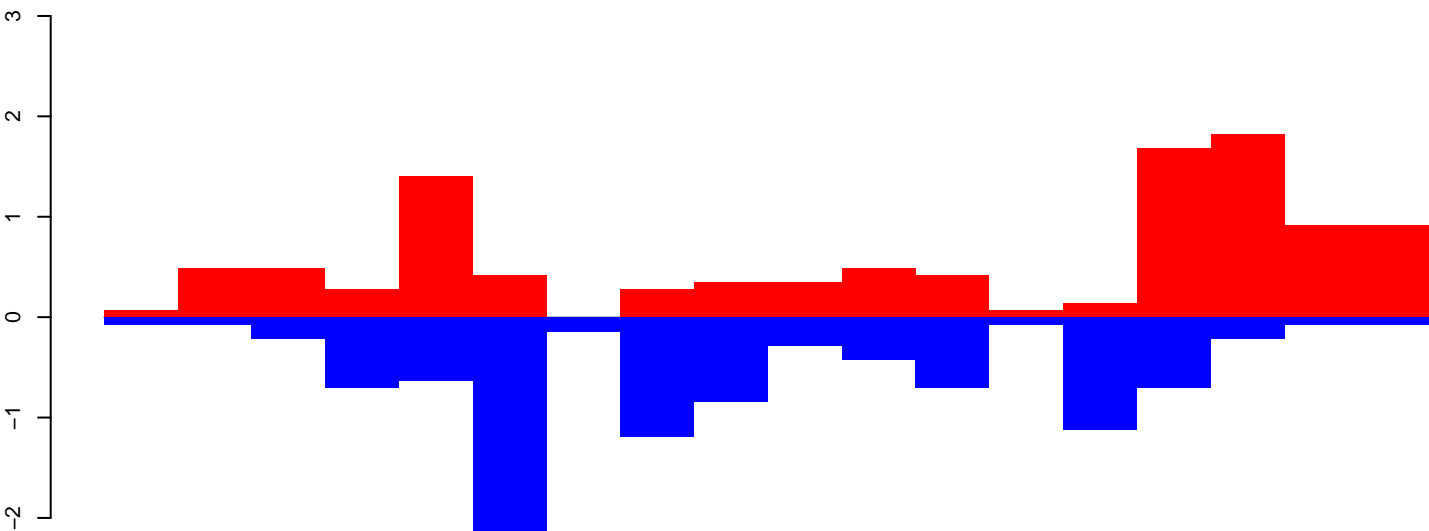


Window size=25, length=502, TE@rnd-4_family-201@DNA@2053:1-502

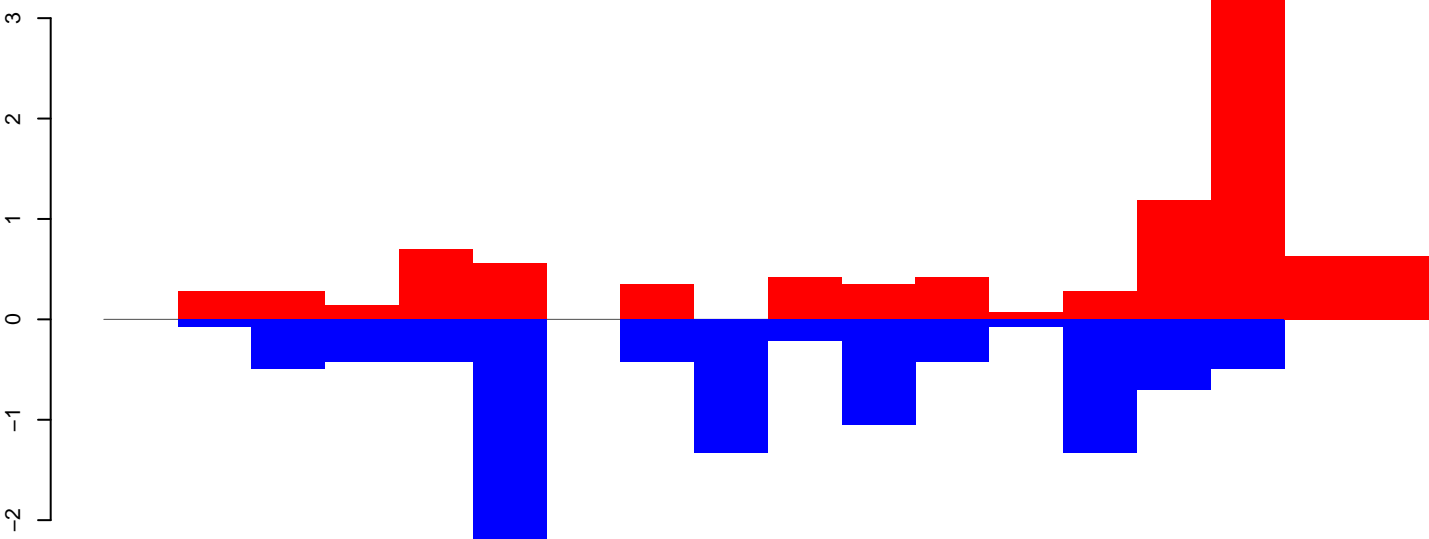
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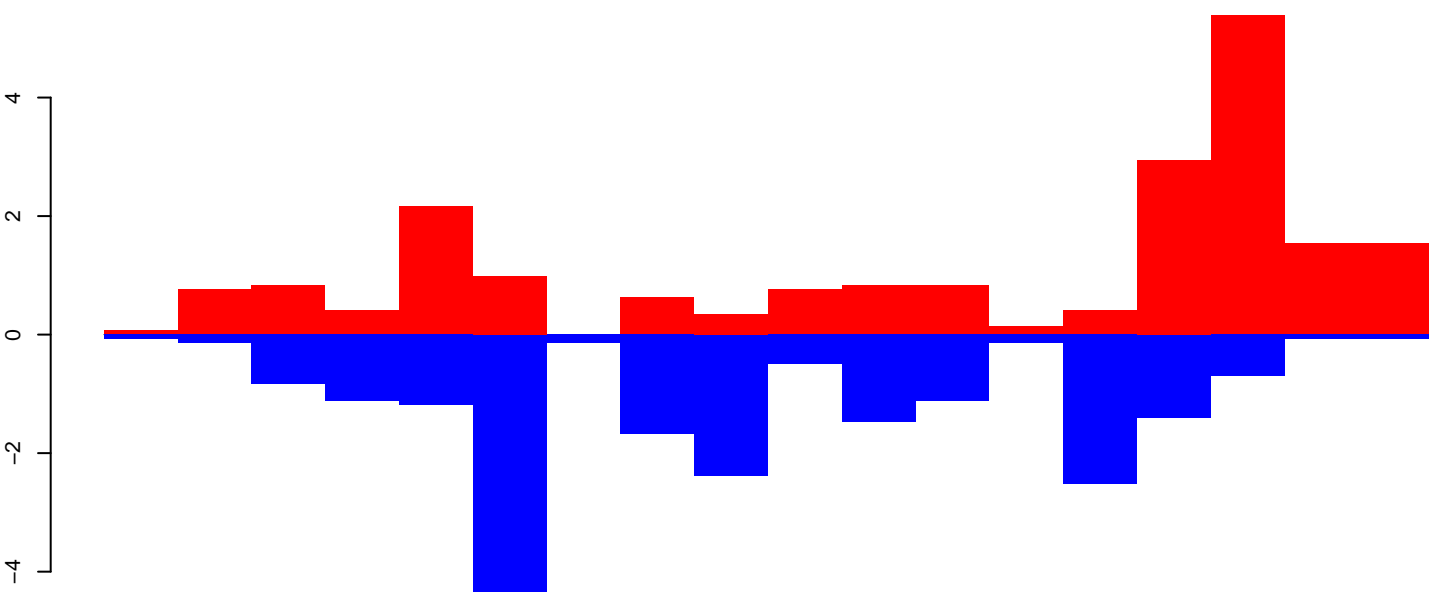
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

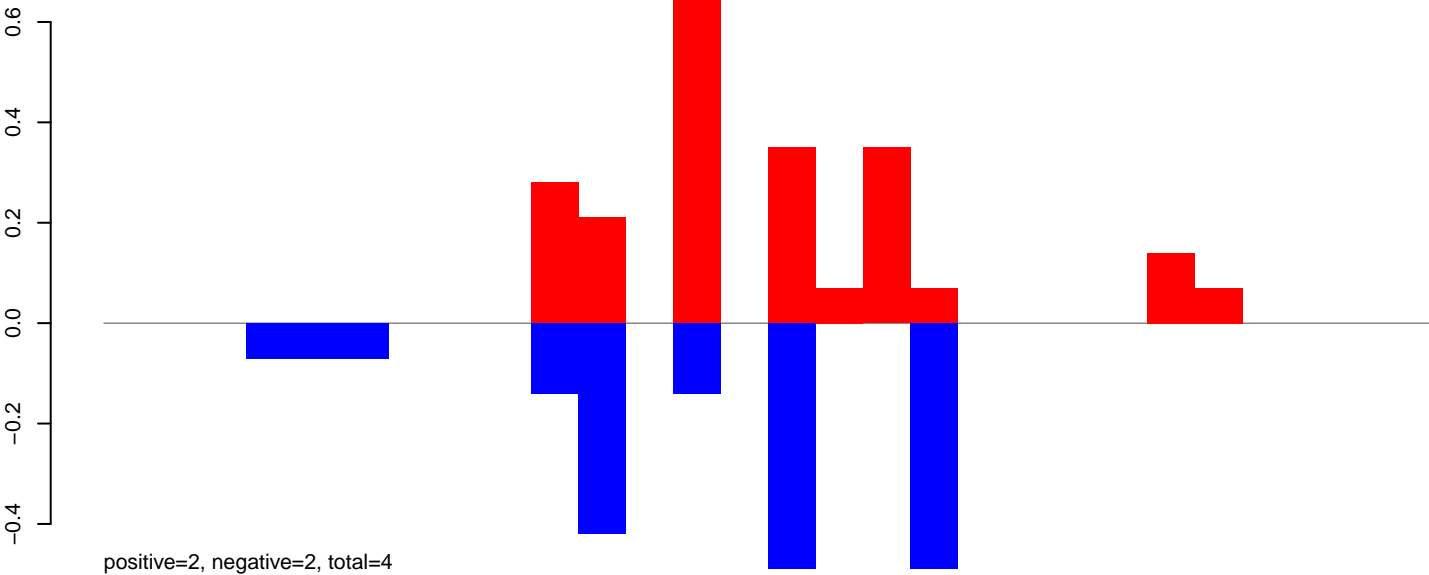


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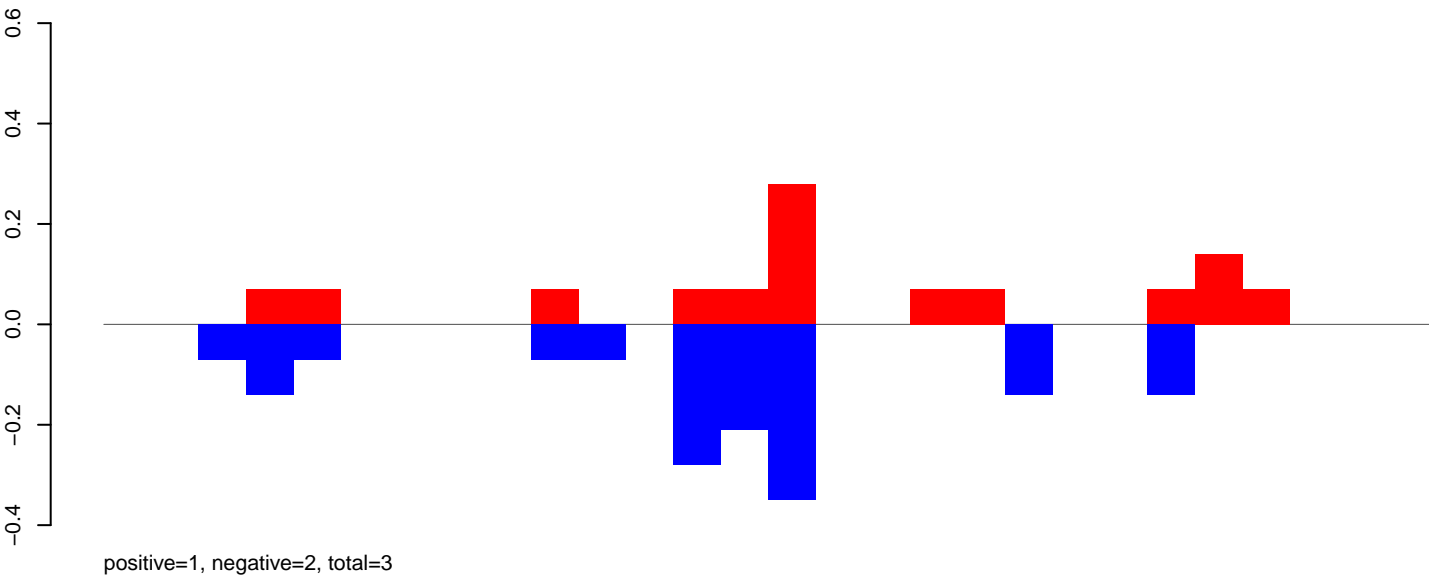


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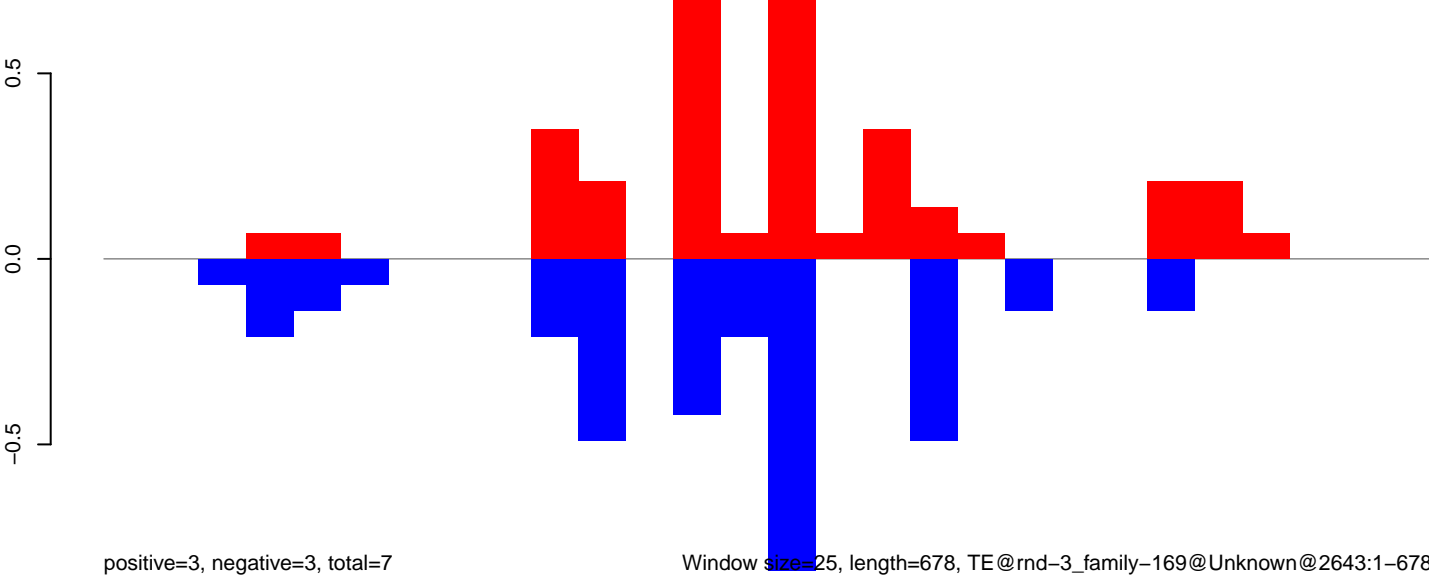
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

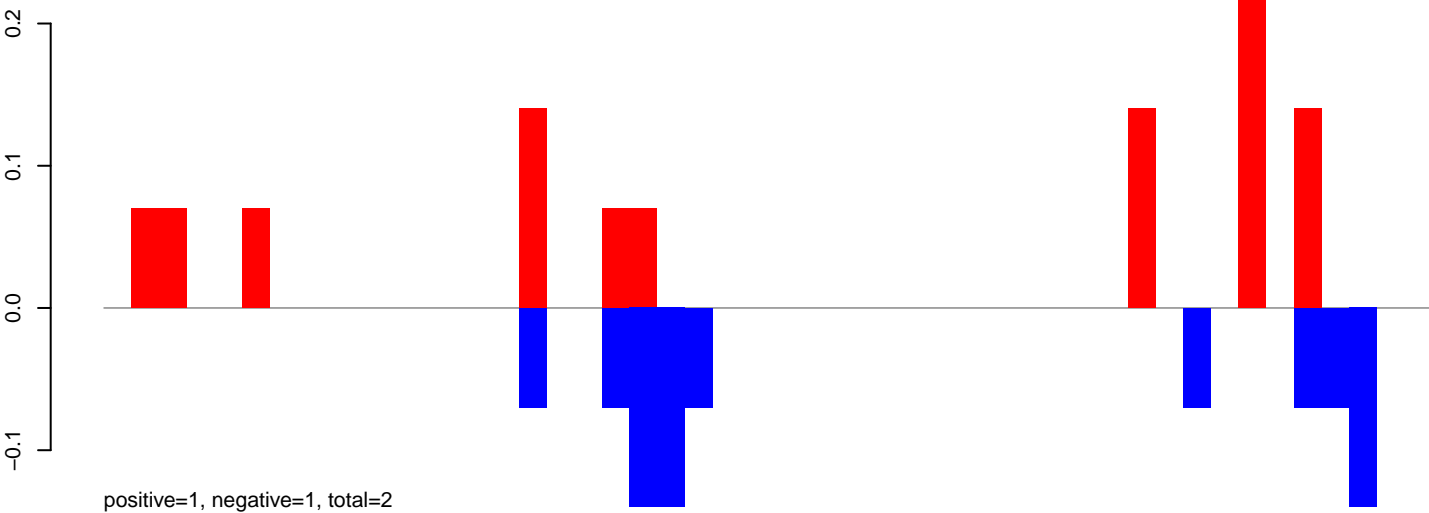


CuQuin_Mcarc_NL.rep

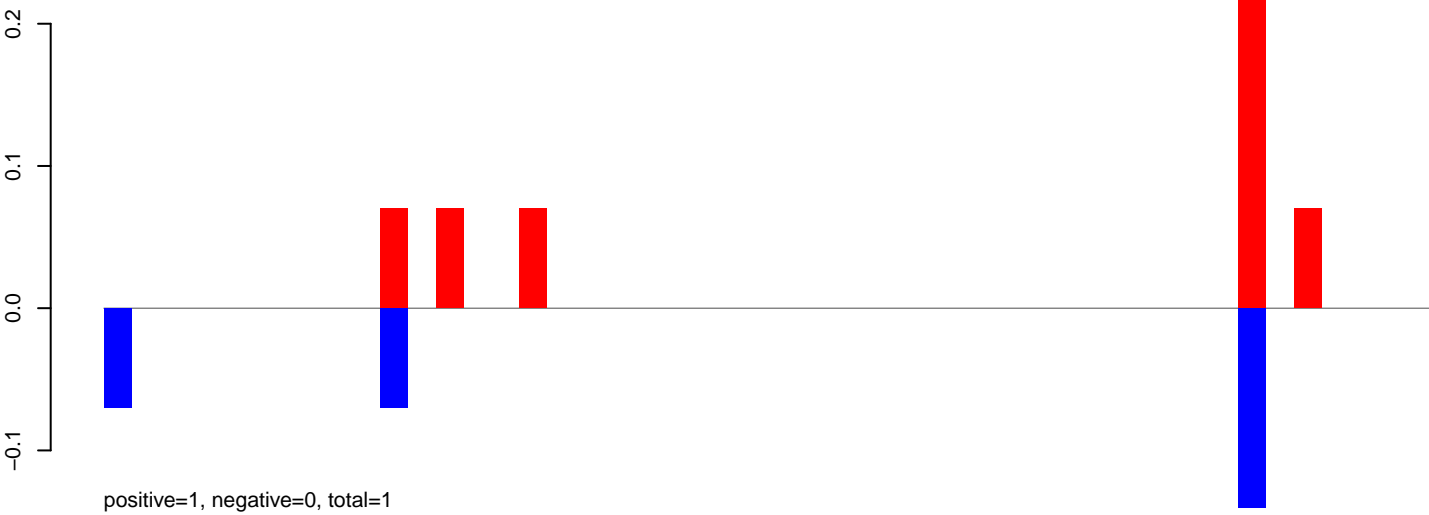


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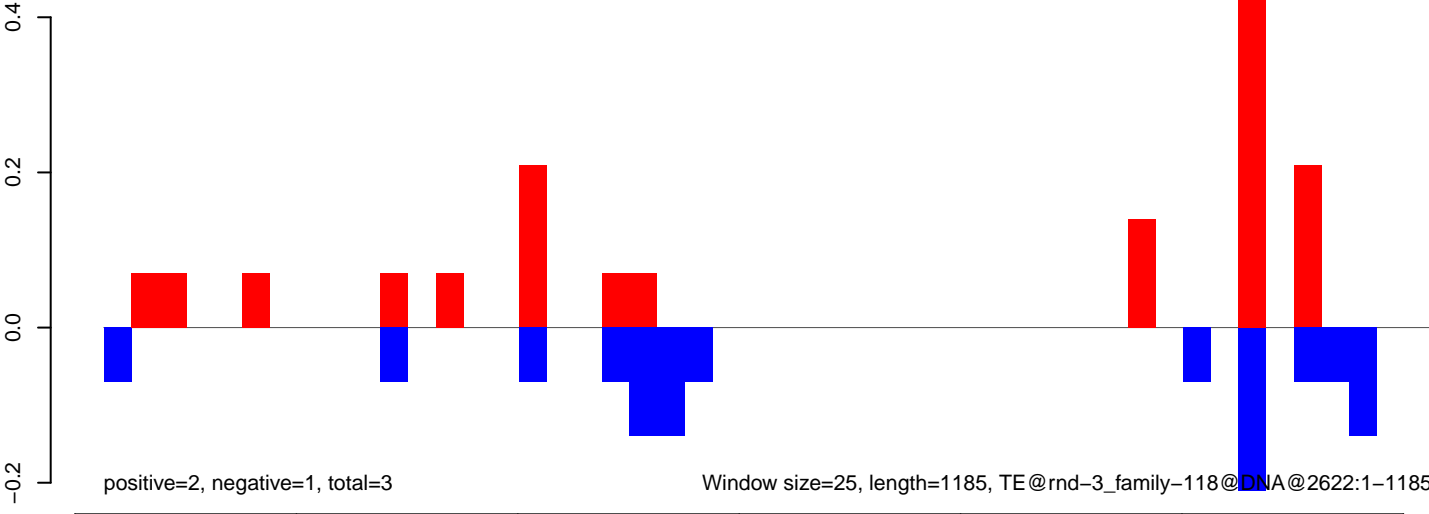
CuQuin_Mcarc_NL.18_23.rep



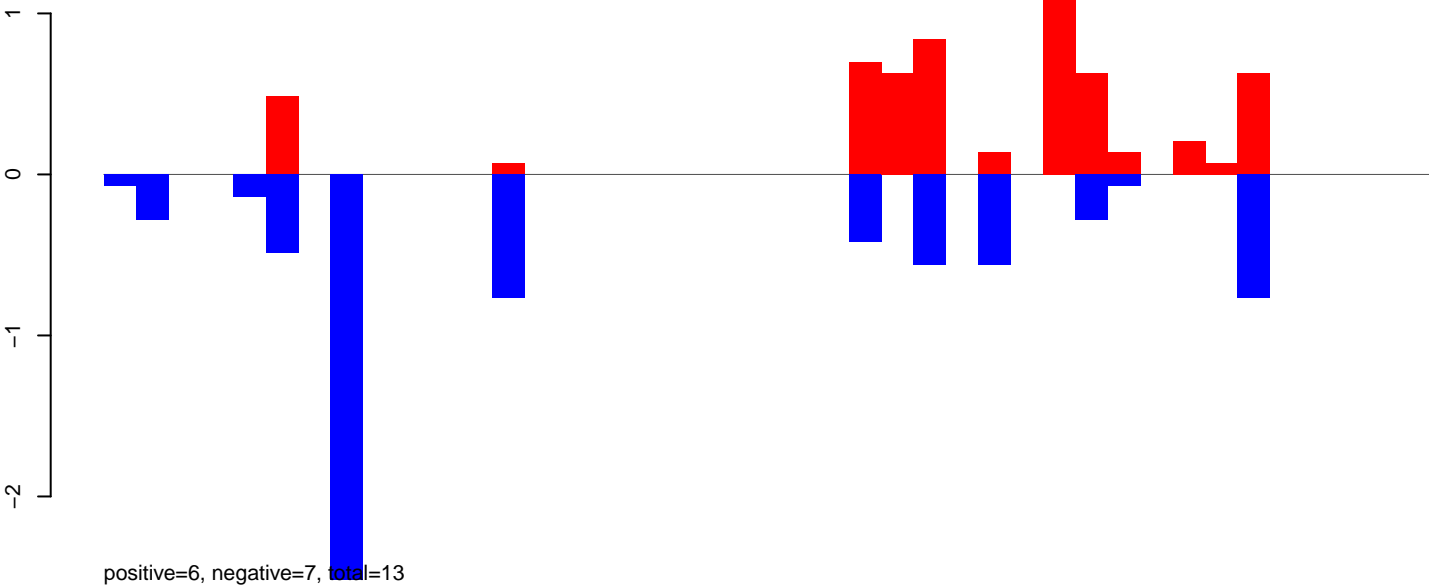
CuQuin_Mcarc_NL.24_35.rep



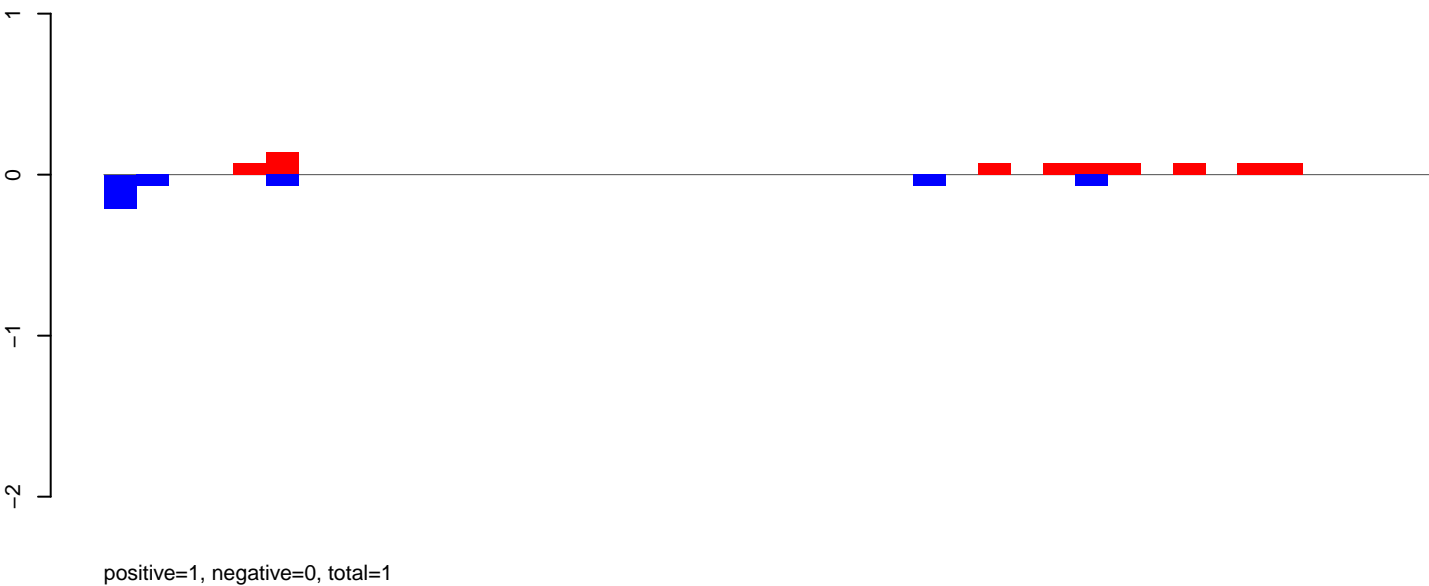
CuQuin_Mcarc_NL.rep



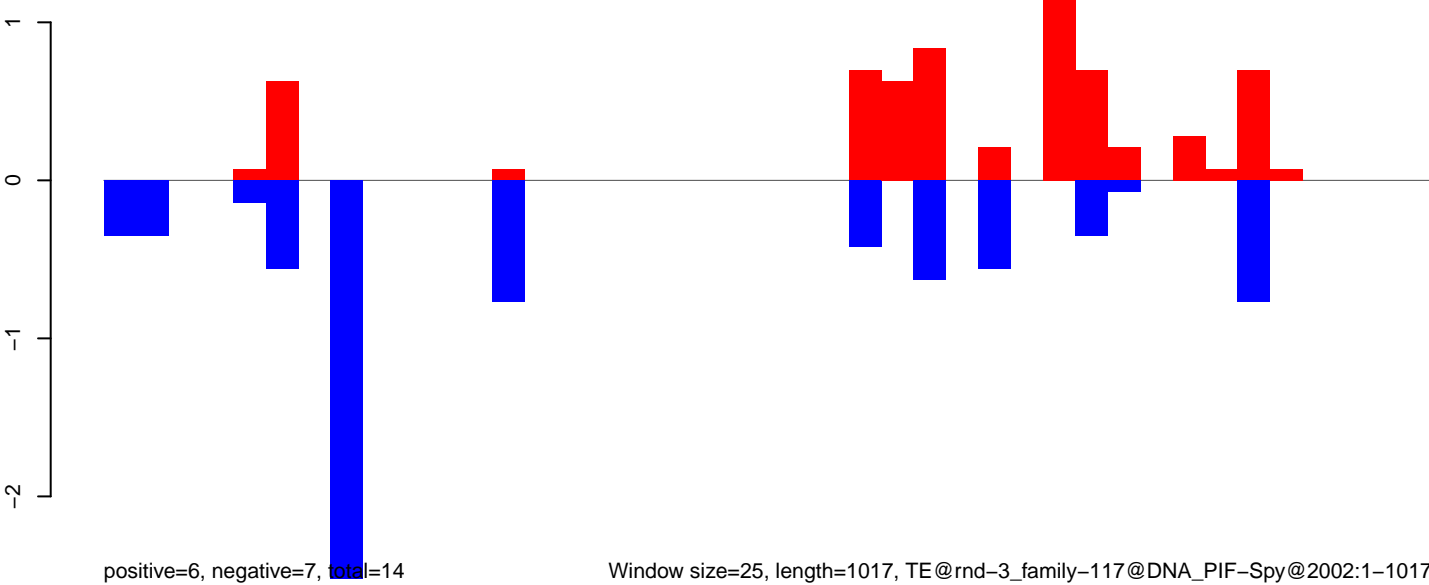
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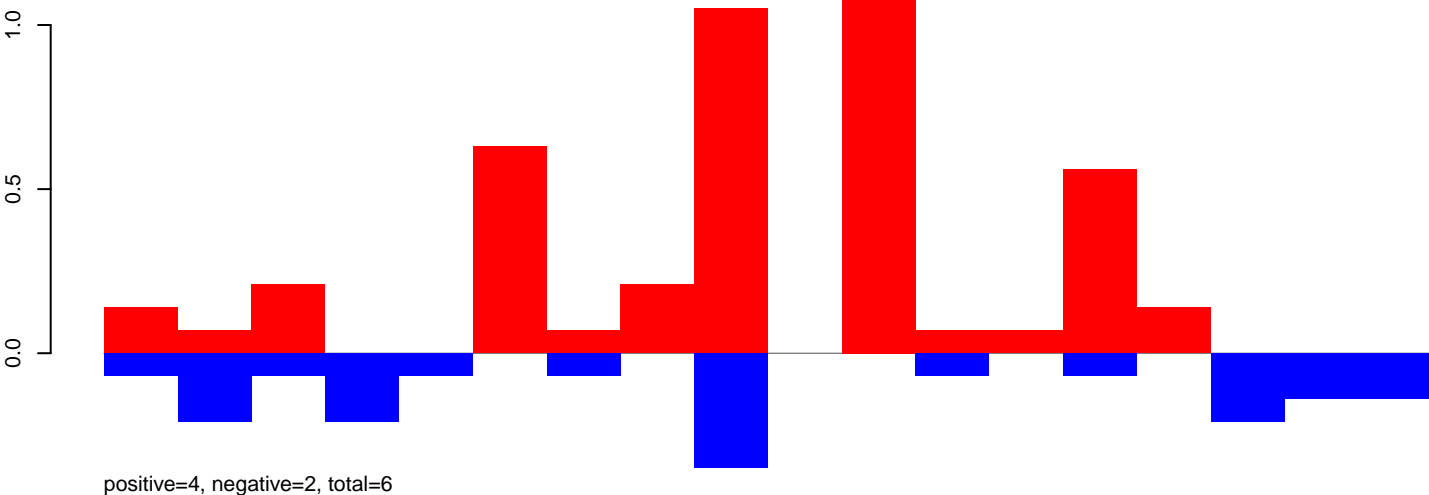
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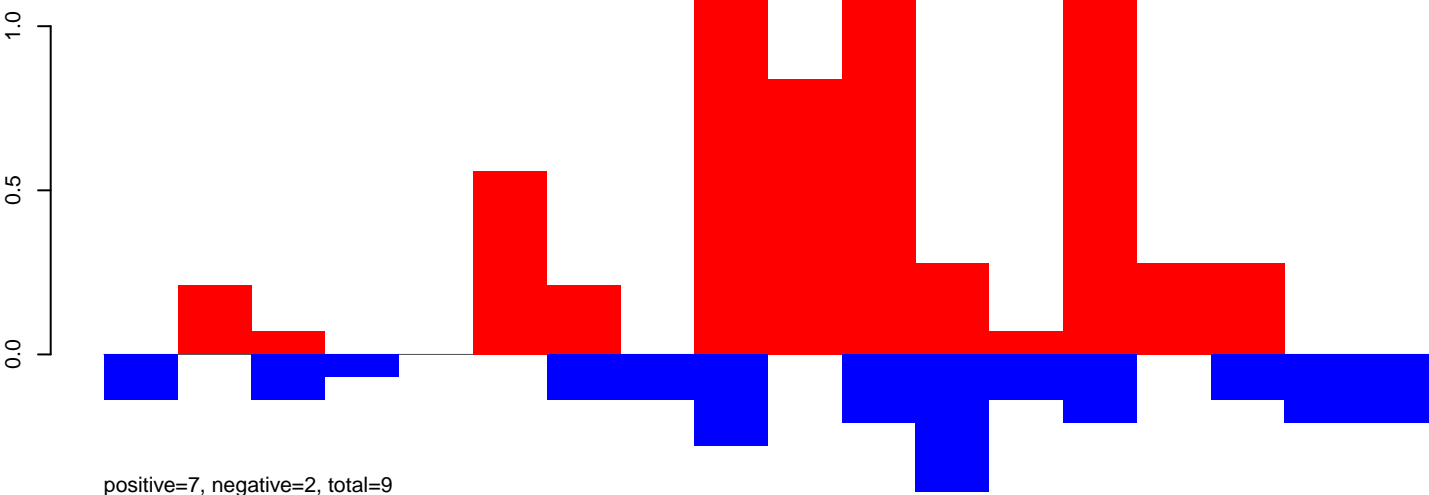
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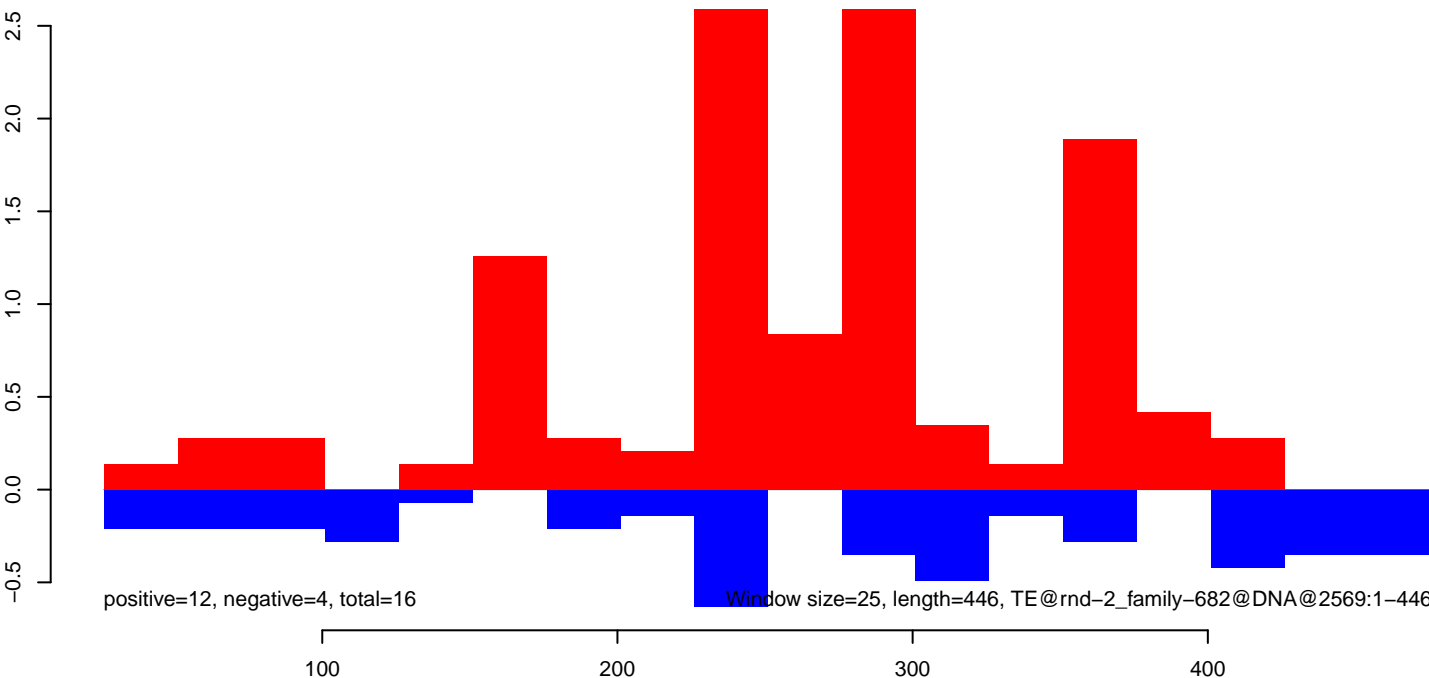
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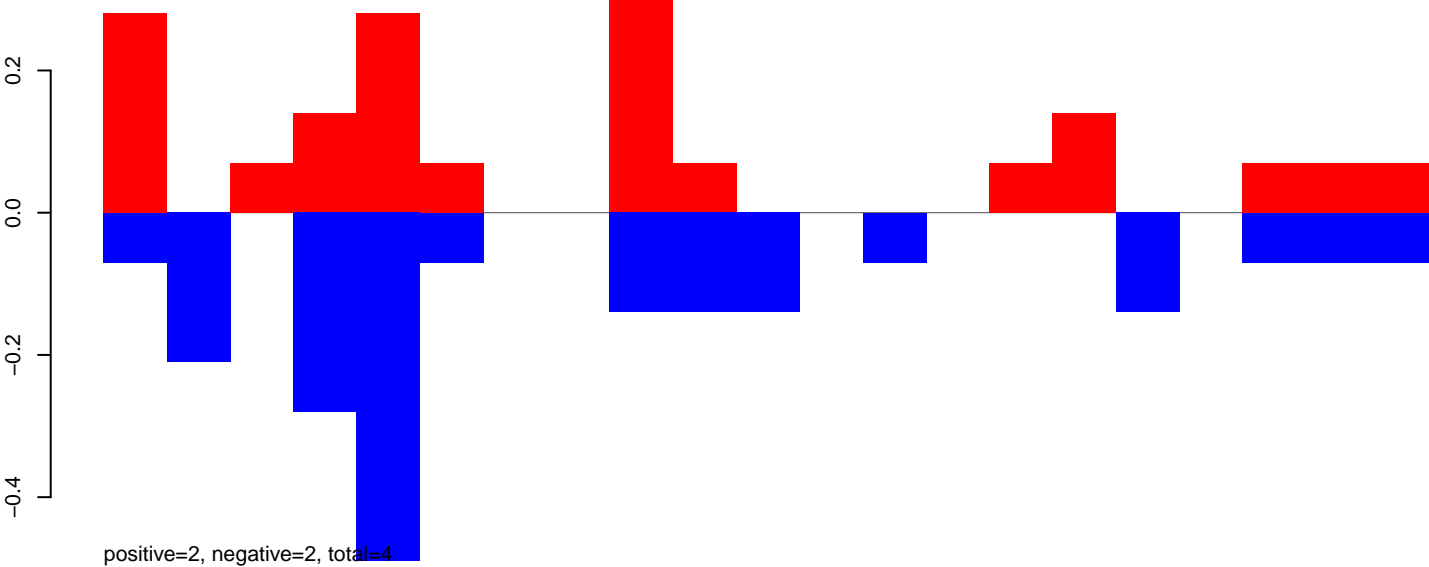
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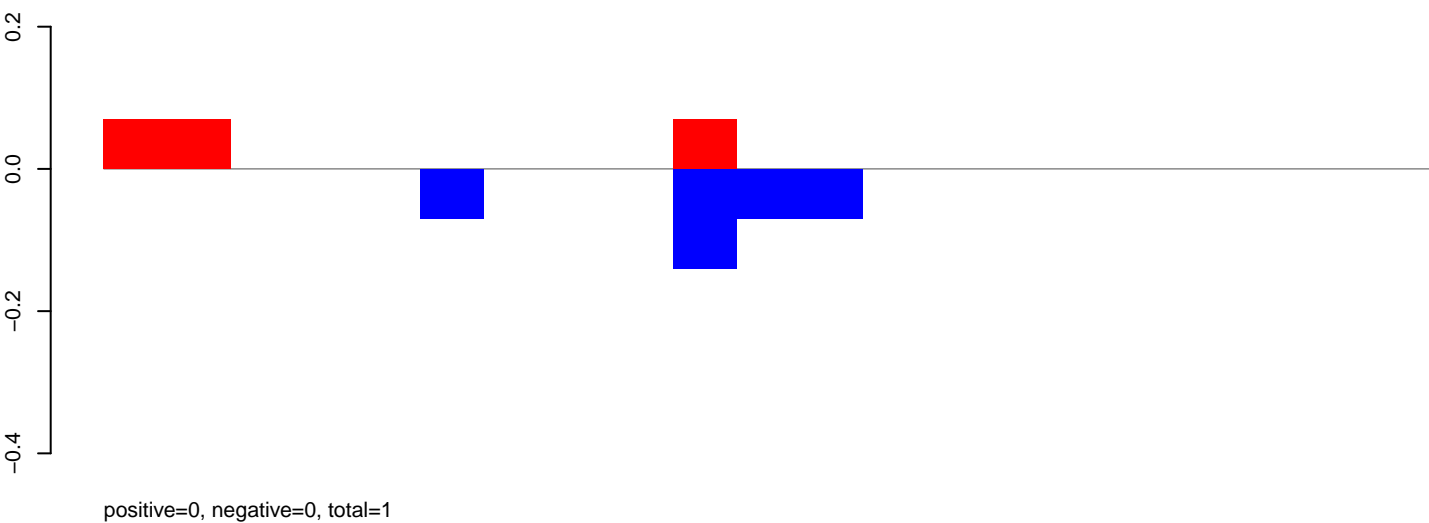
CuQuin_Mcarc_NL.rep



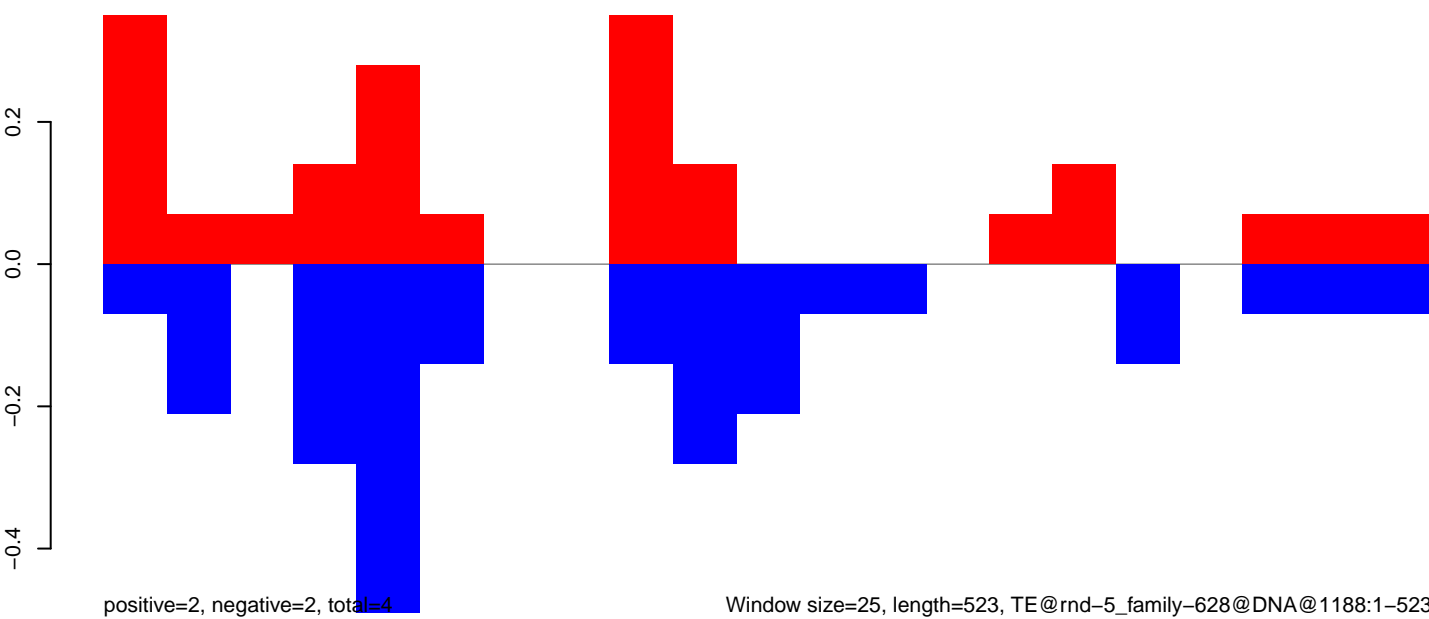
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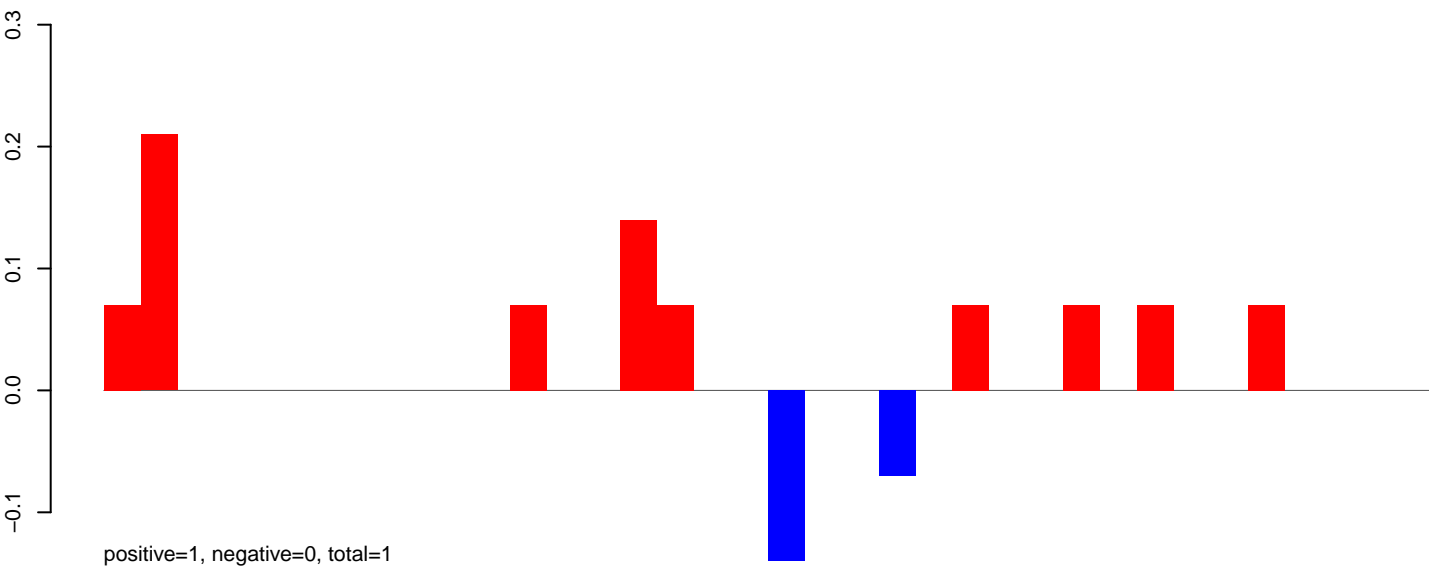
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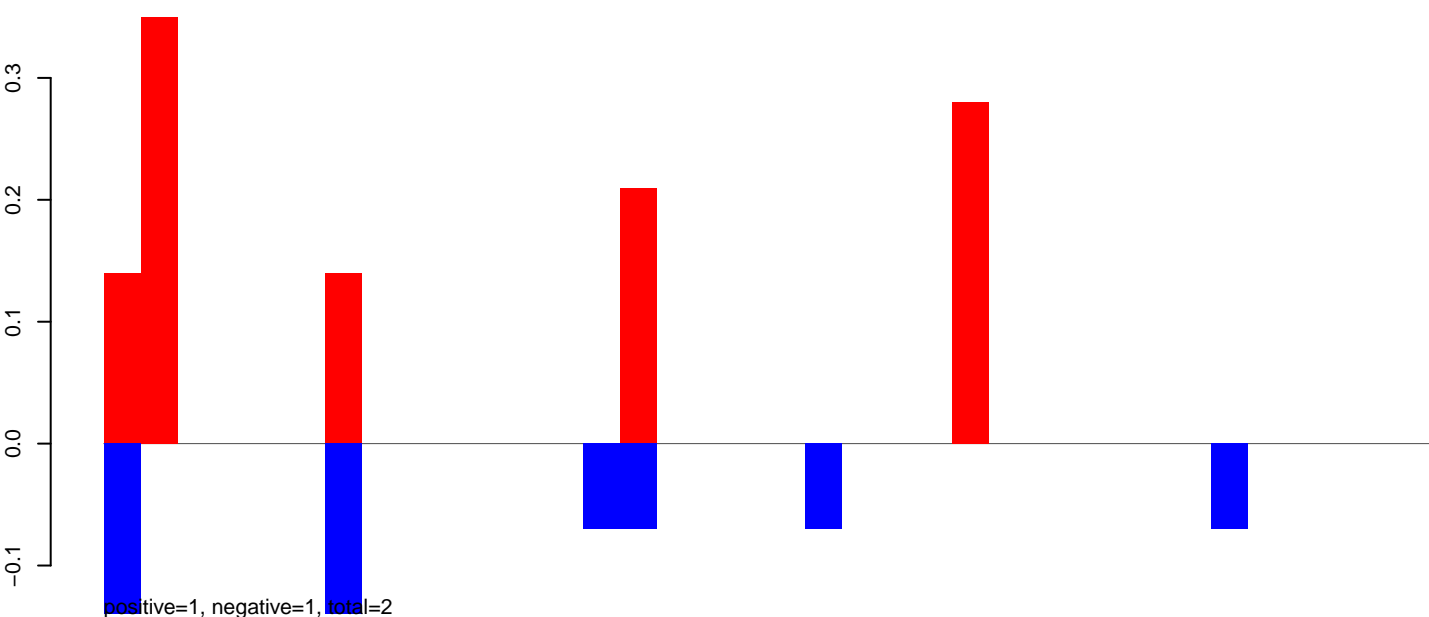
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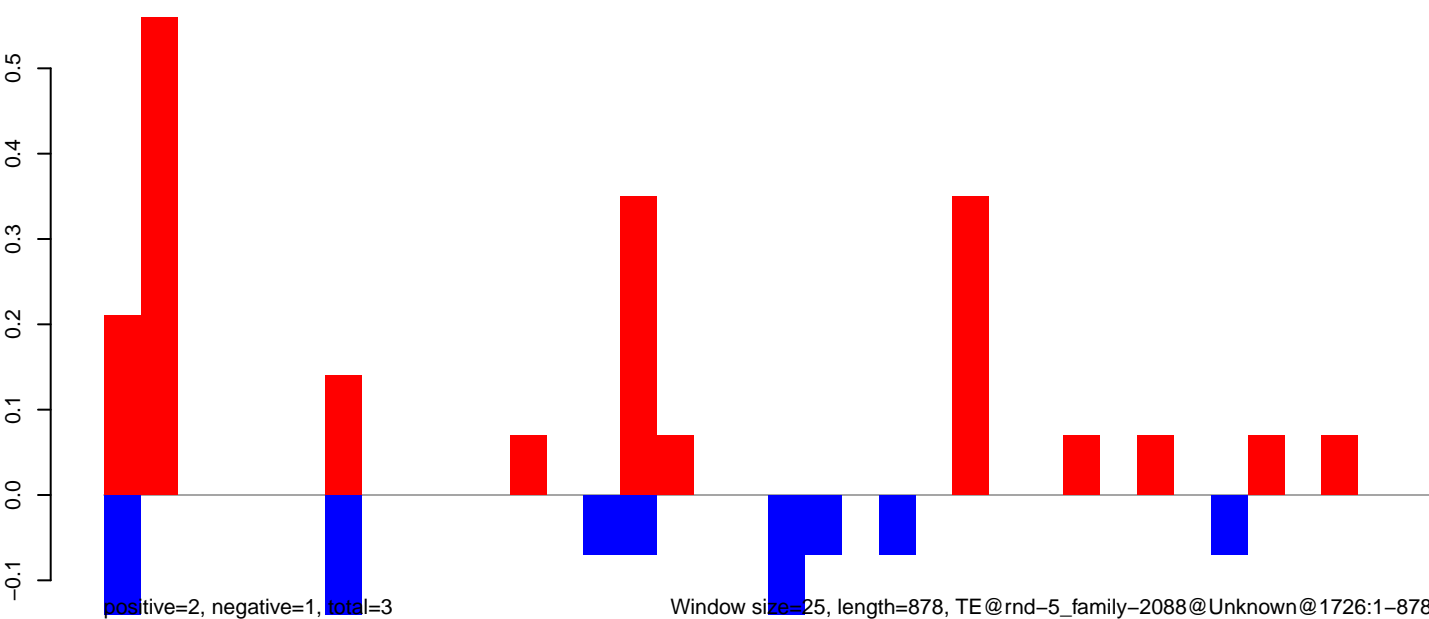
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CuQuin_Mcarc_NL.24_35.rep

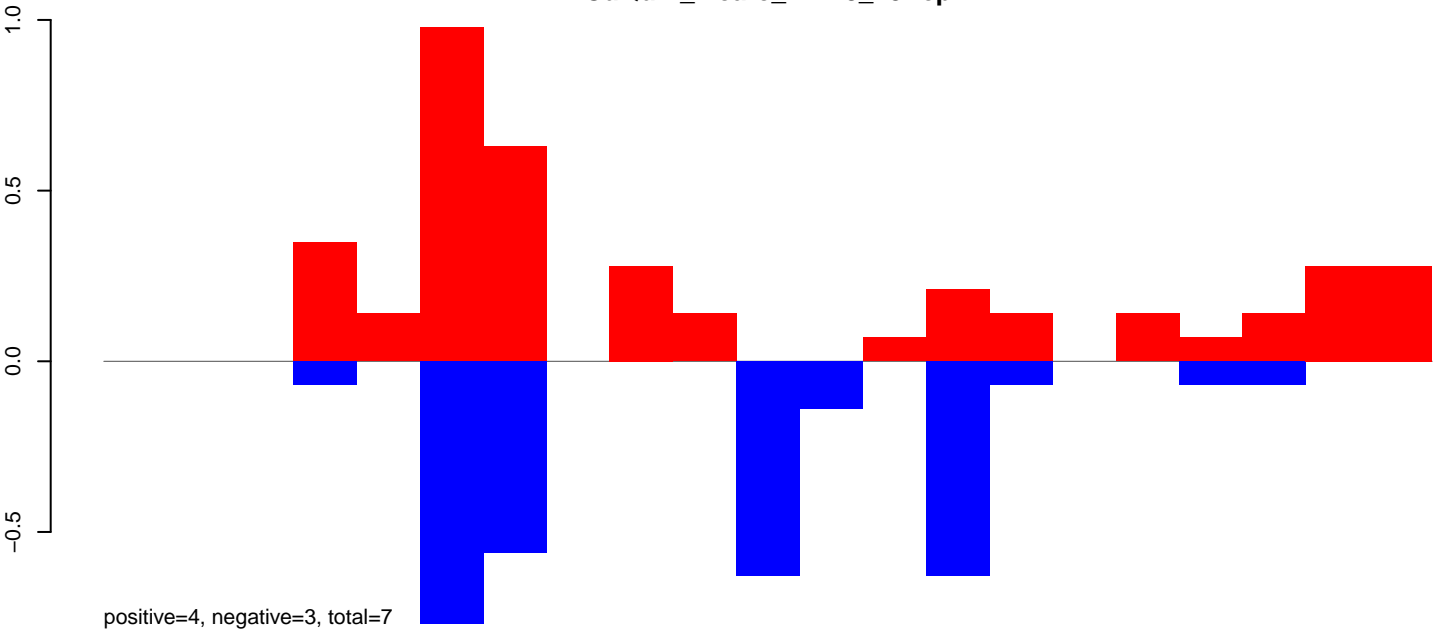


CuQuin_Mcarc_NL.rep

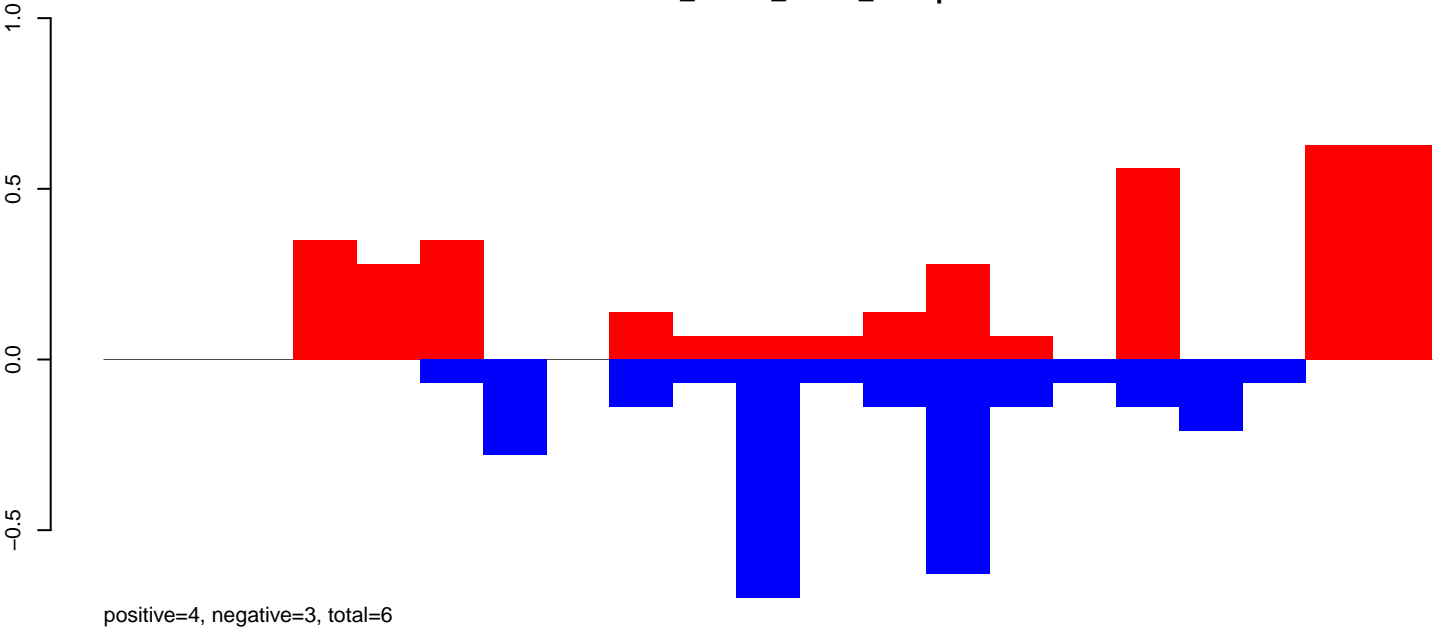


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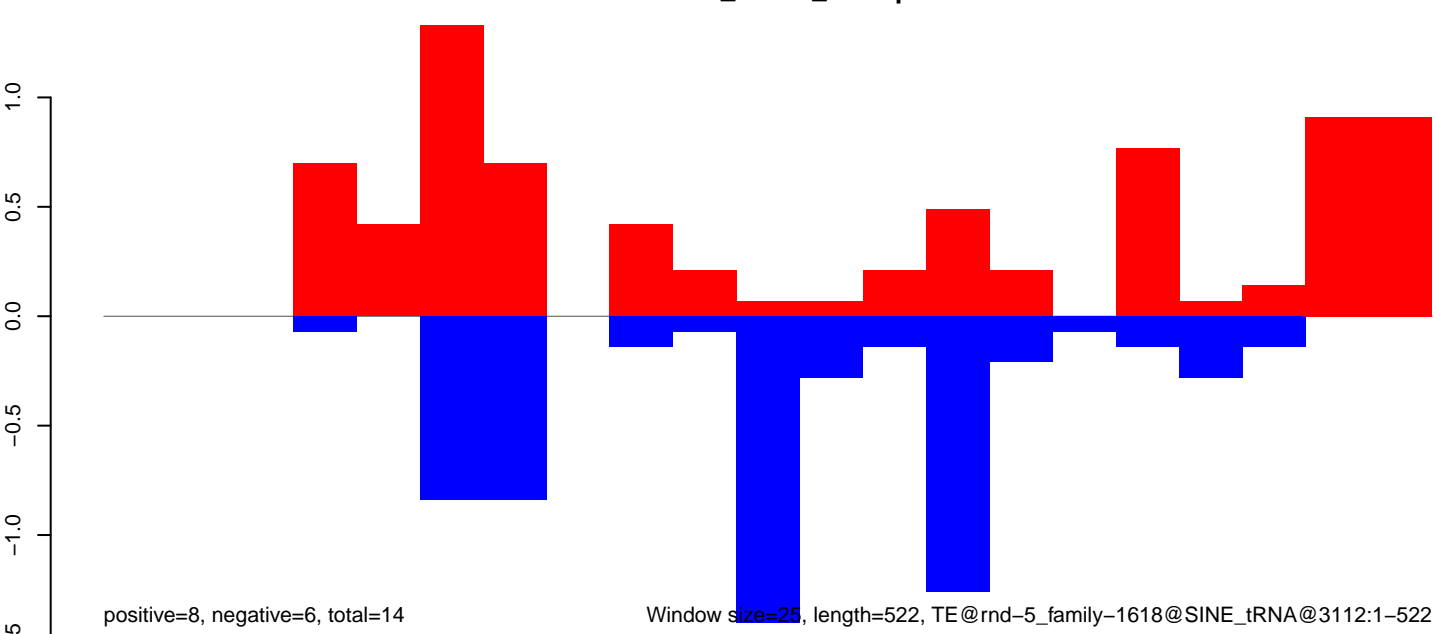
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

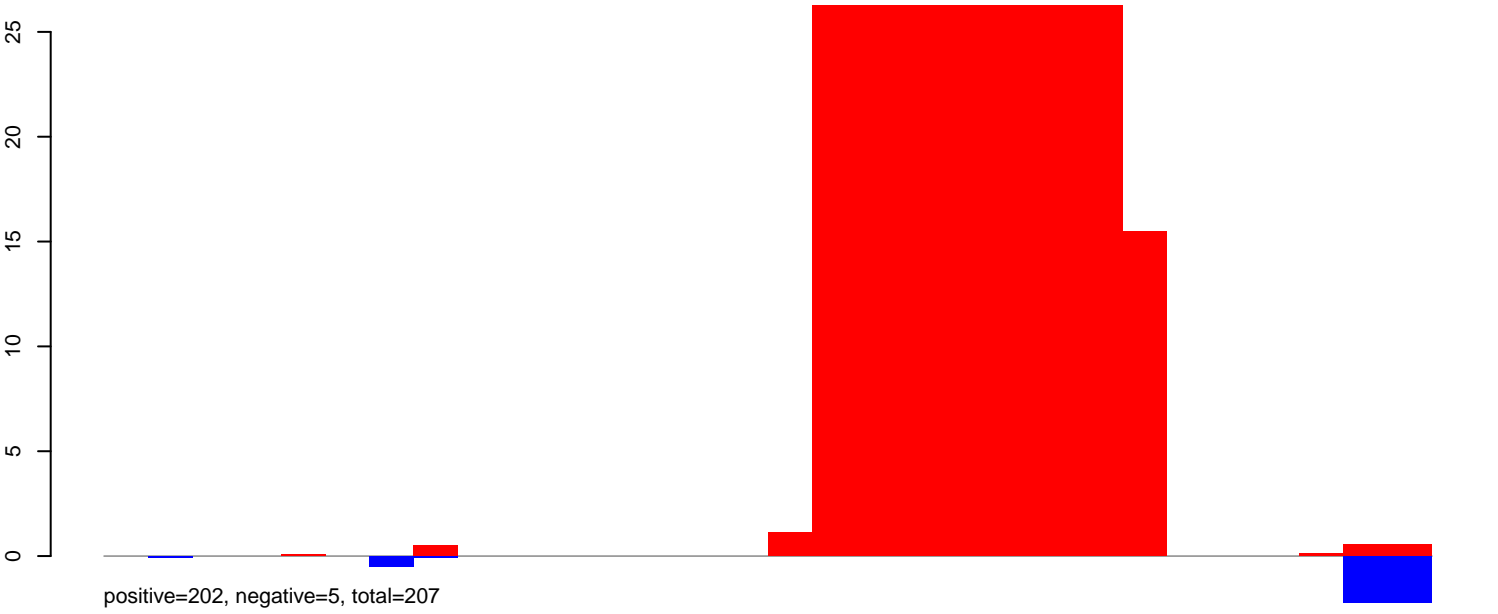


CuQuin_Mcarc_NL.rep

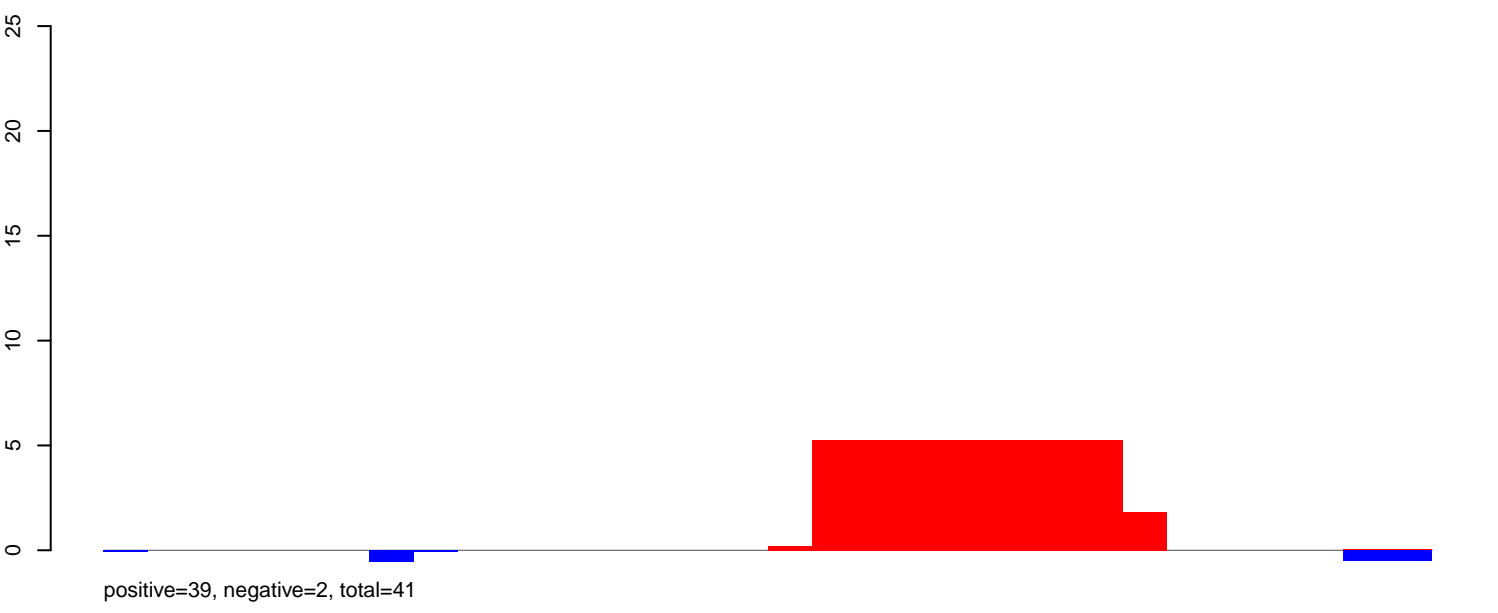


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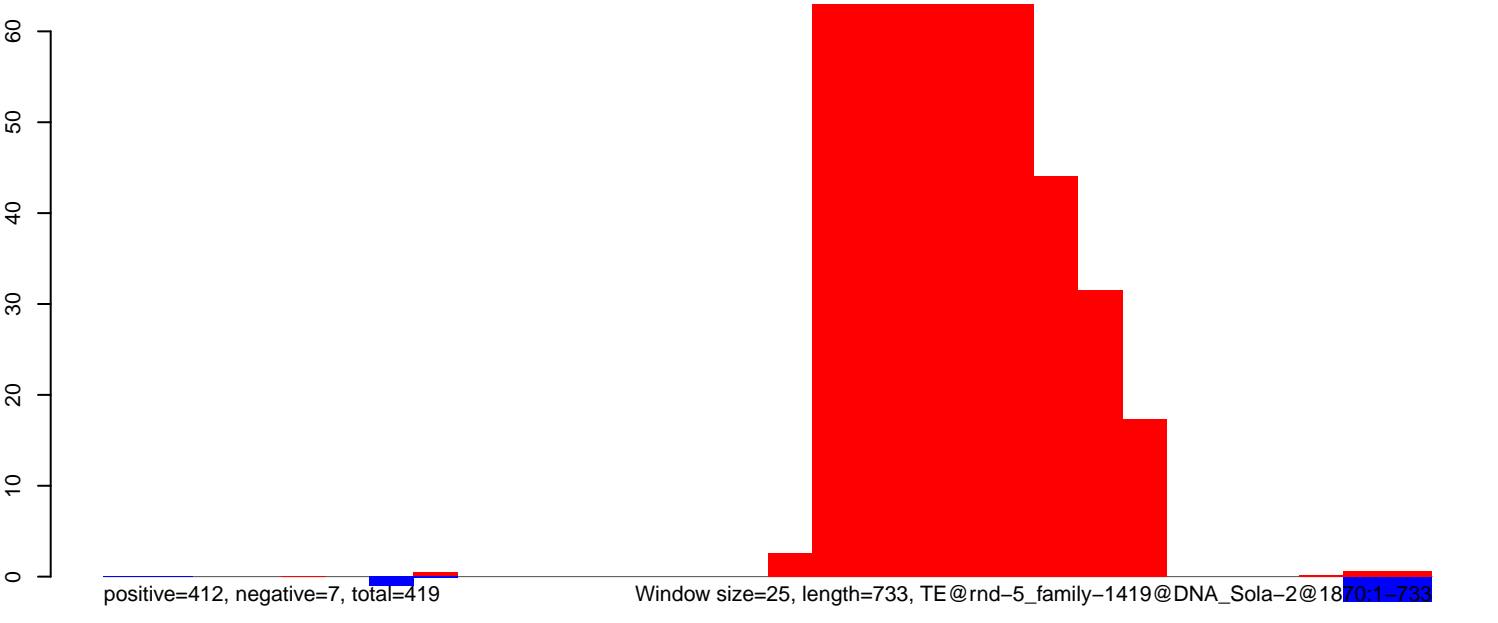
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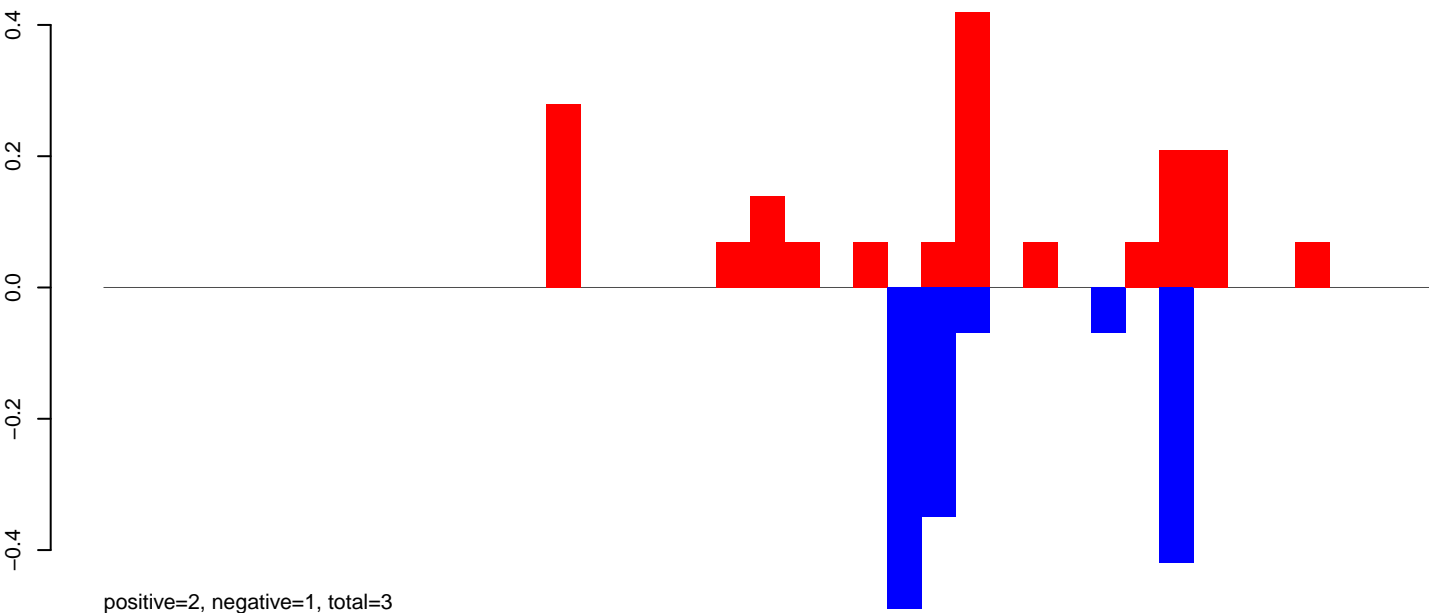
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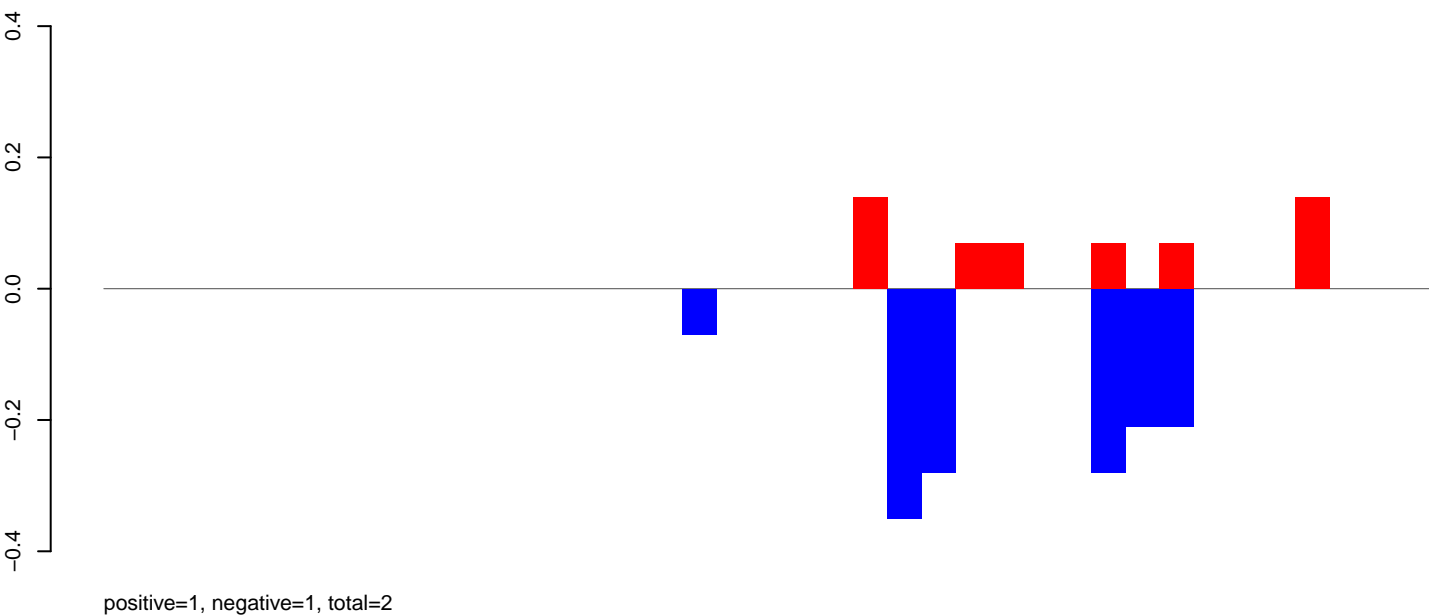
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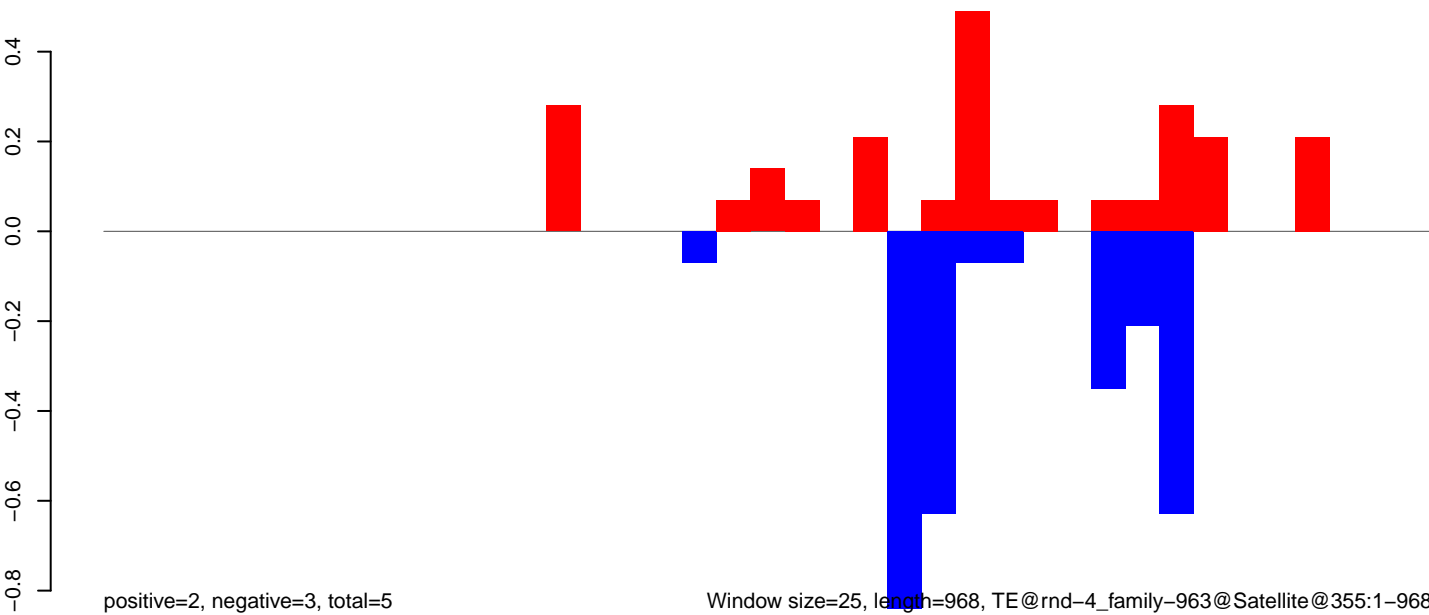
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CuQuin_Mcarc_NL.24_35.rep

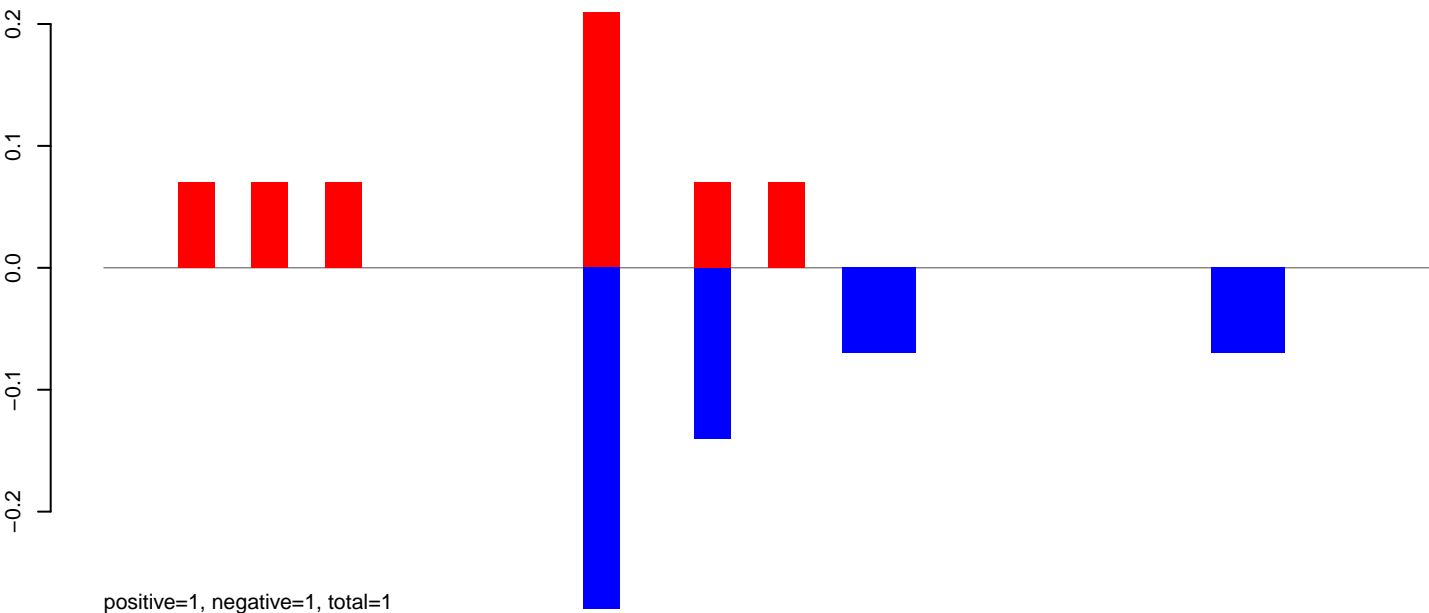


CuQuin_Mcarc_NL.rep

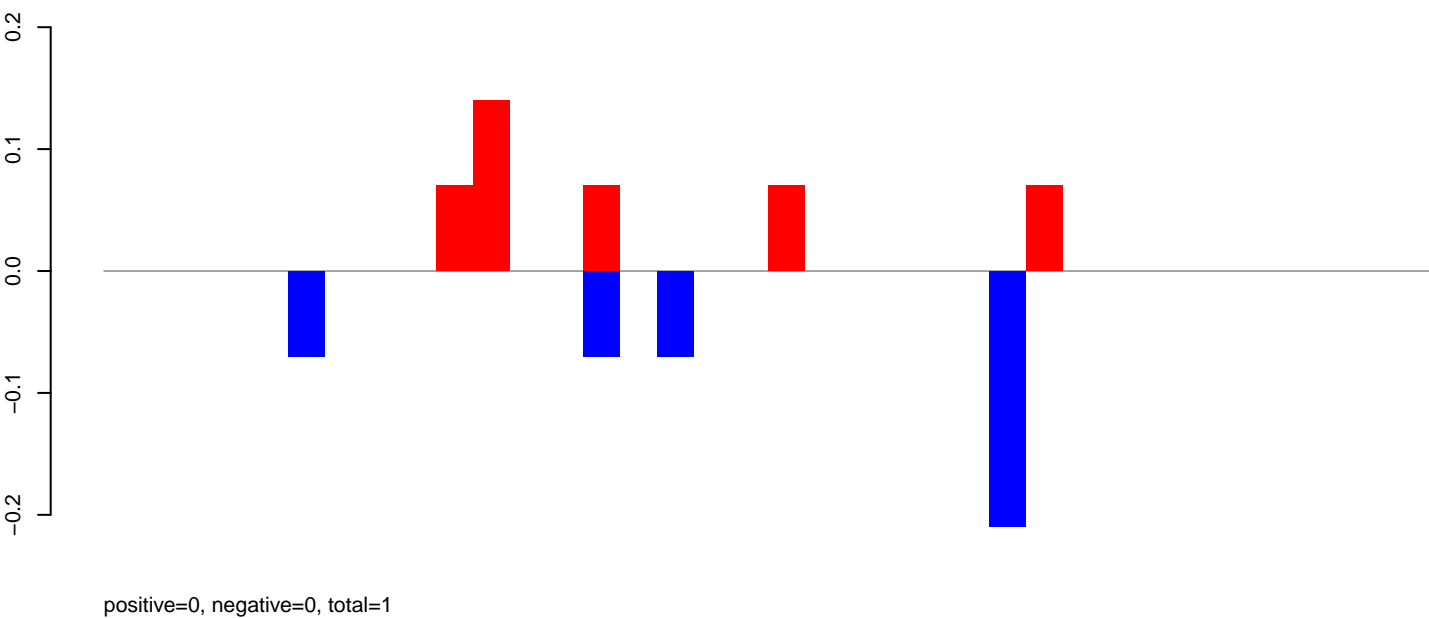


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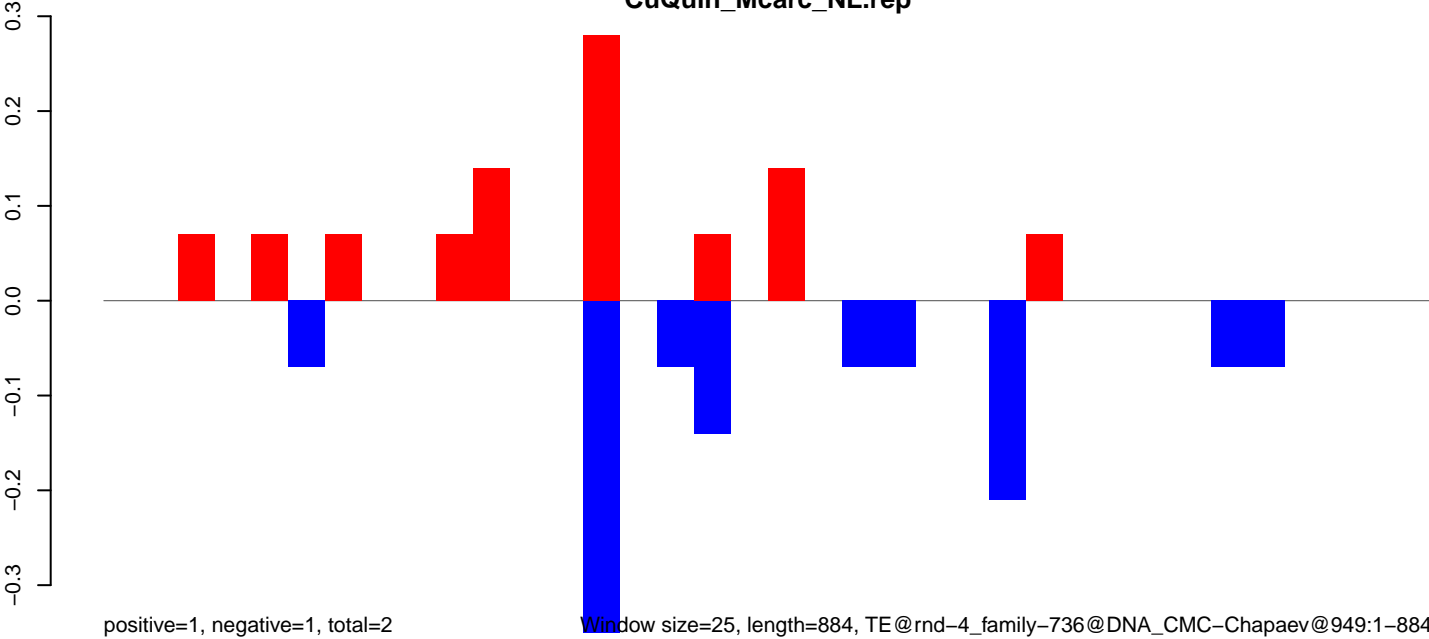
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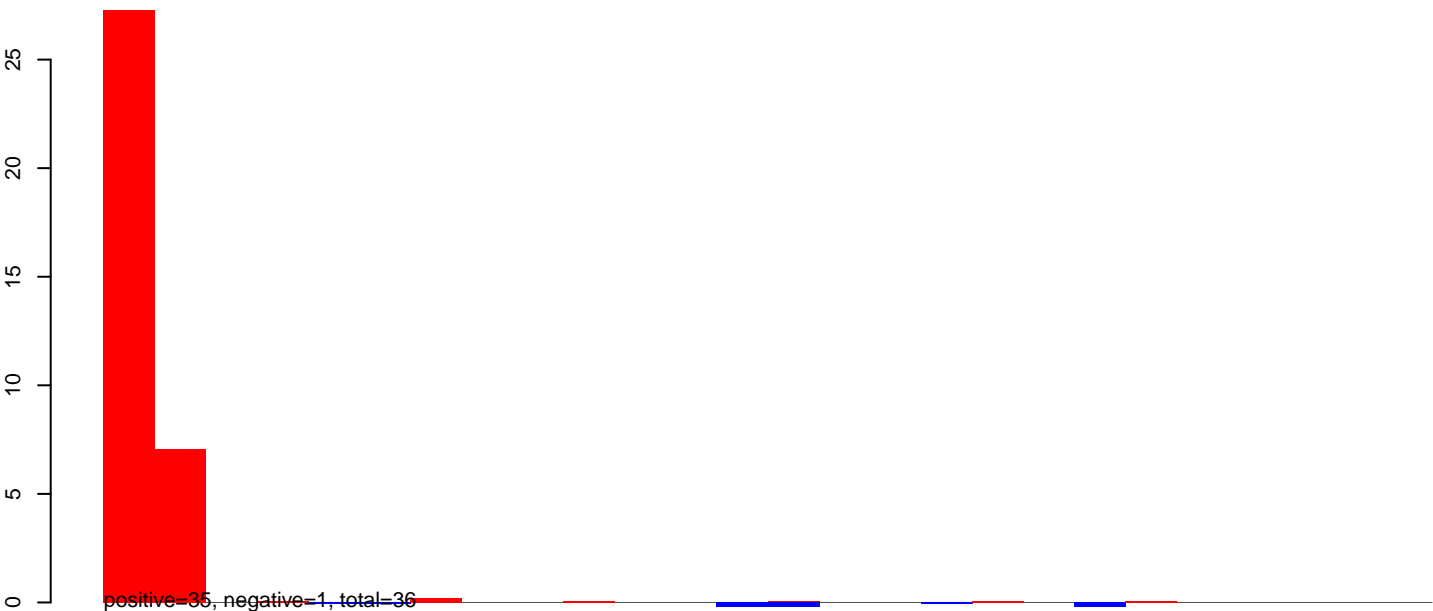
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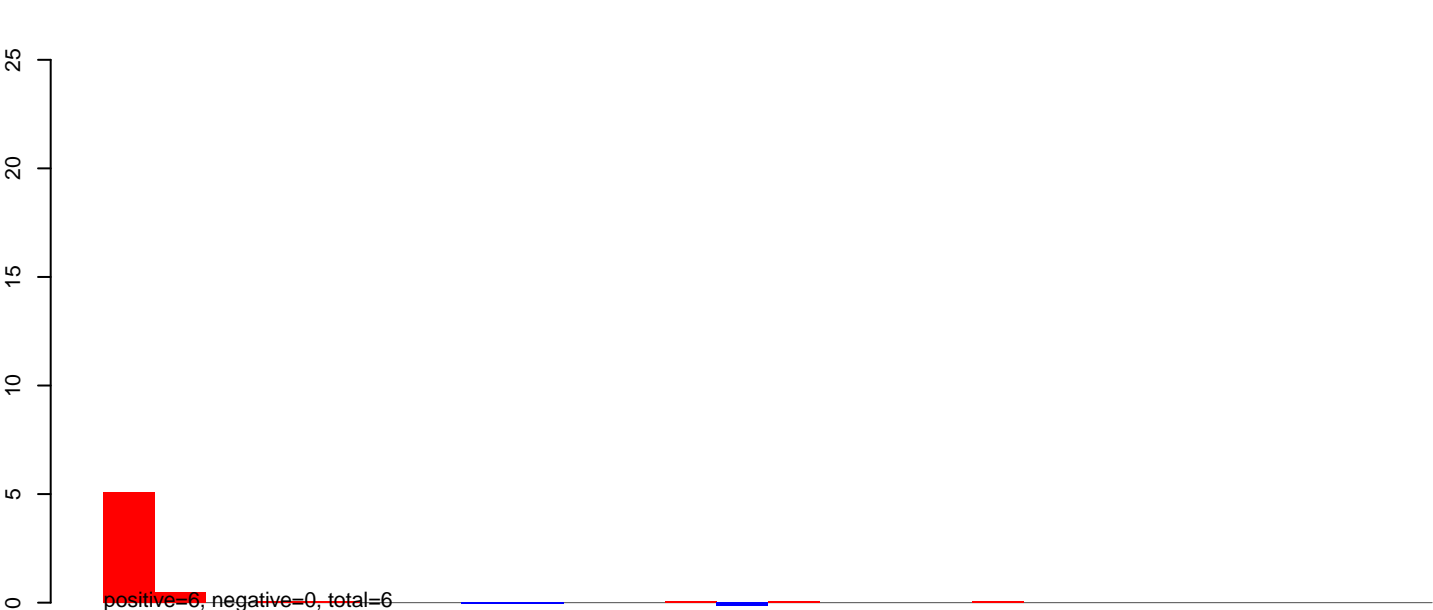
CuQuin_Mcarc_NL.rep



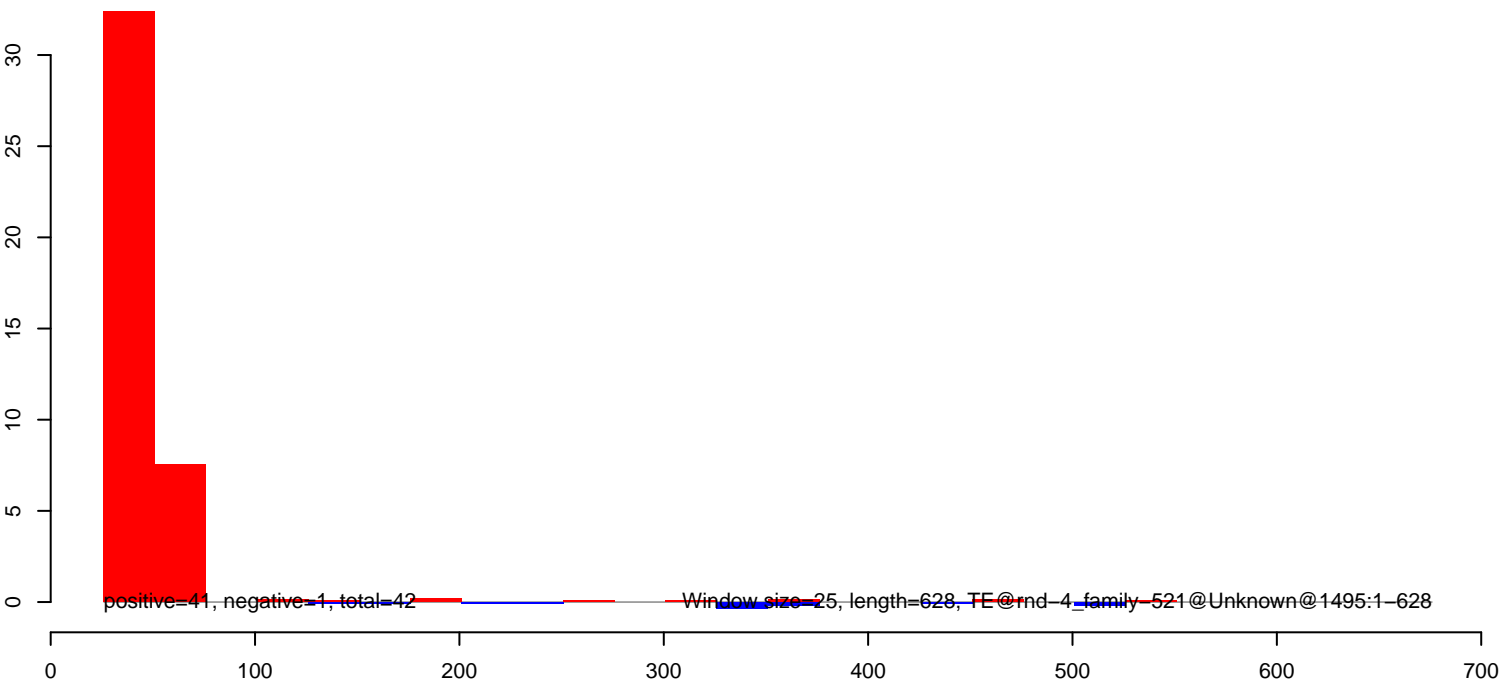
CuQuin_Mcarc_NL.18_23.rep



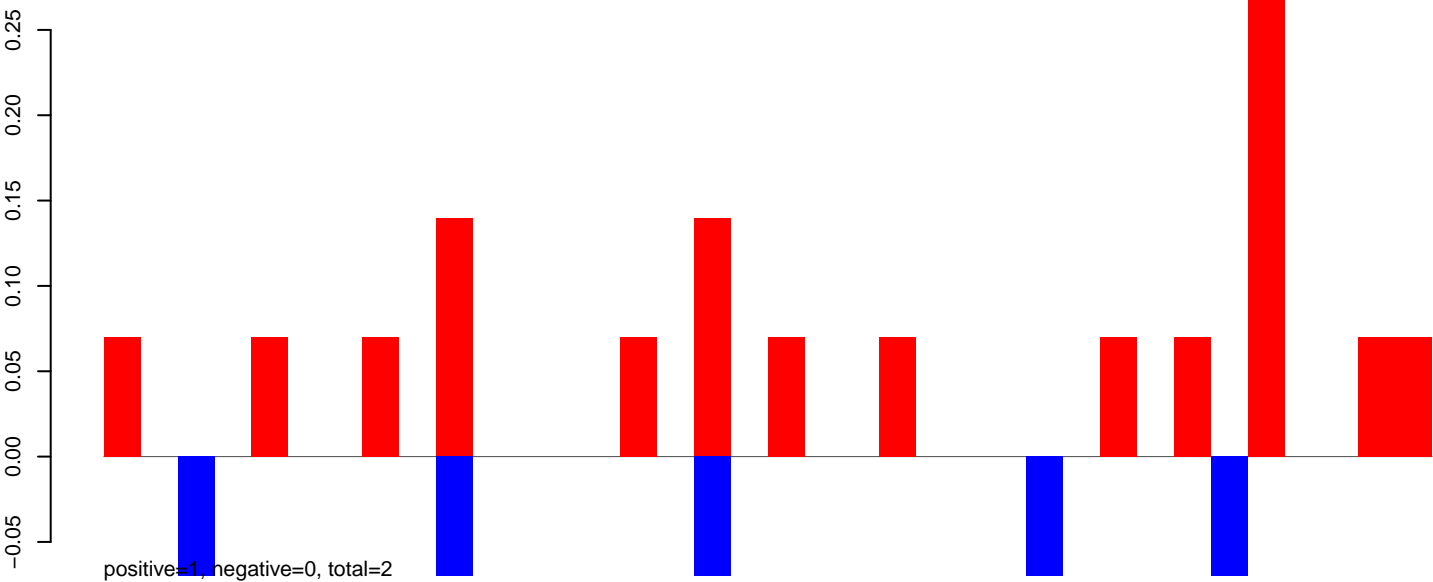
CuQuin_Mcarc_NL.24_35.rep



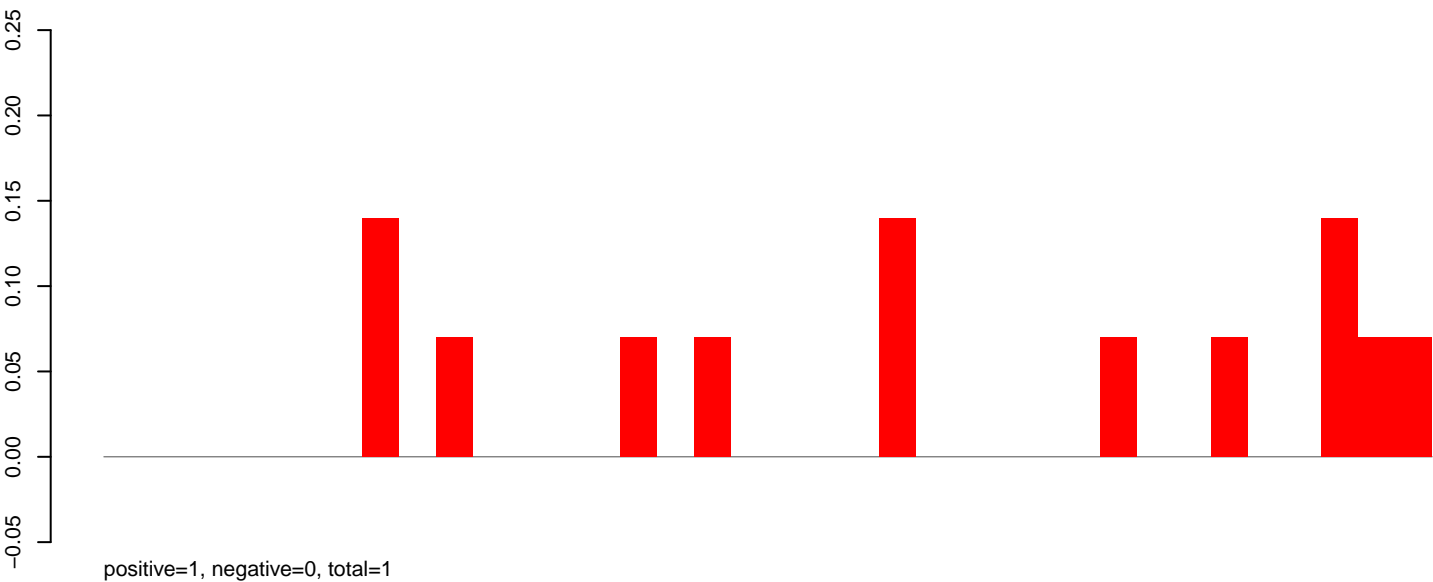
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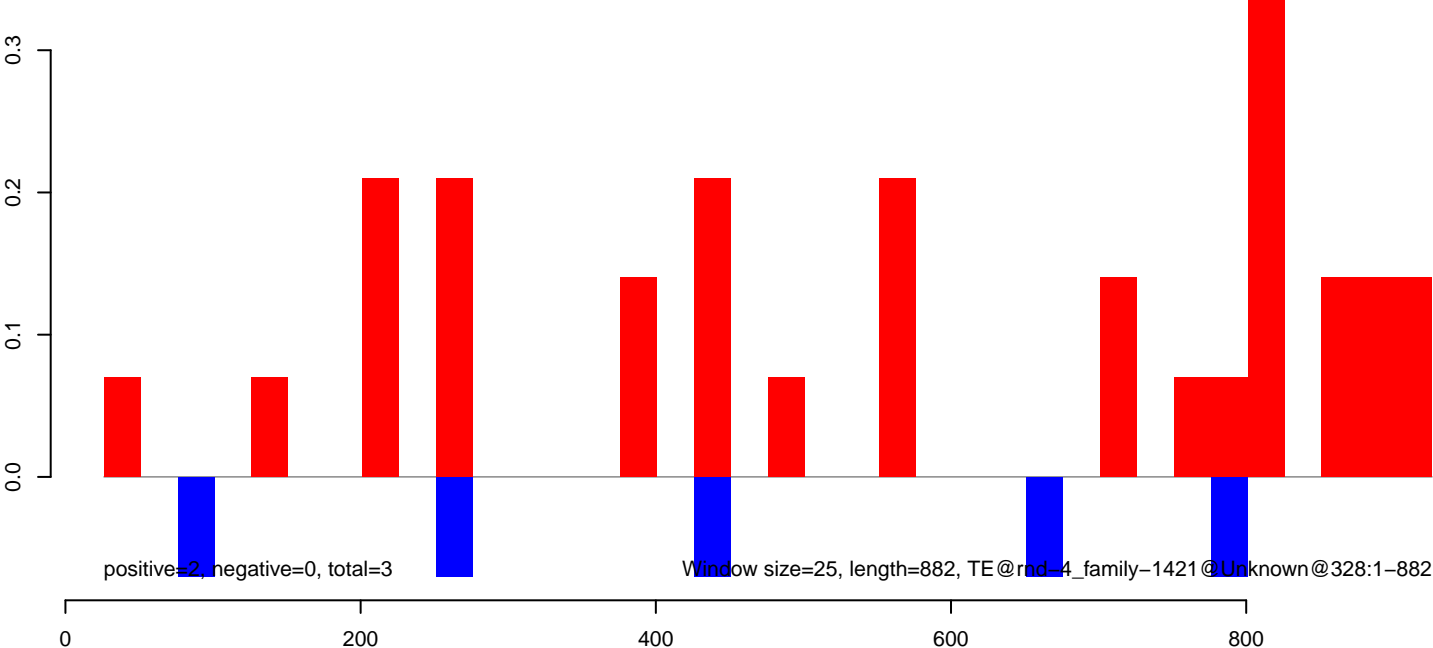
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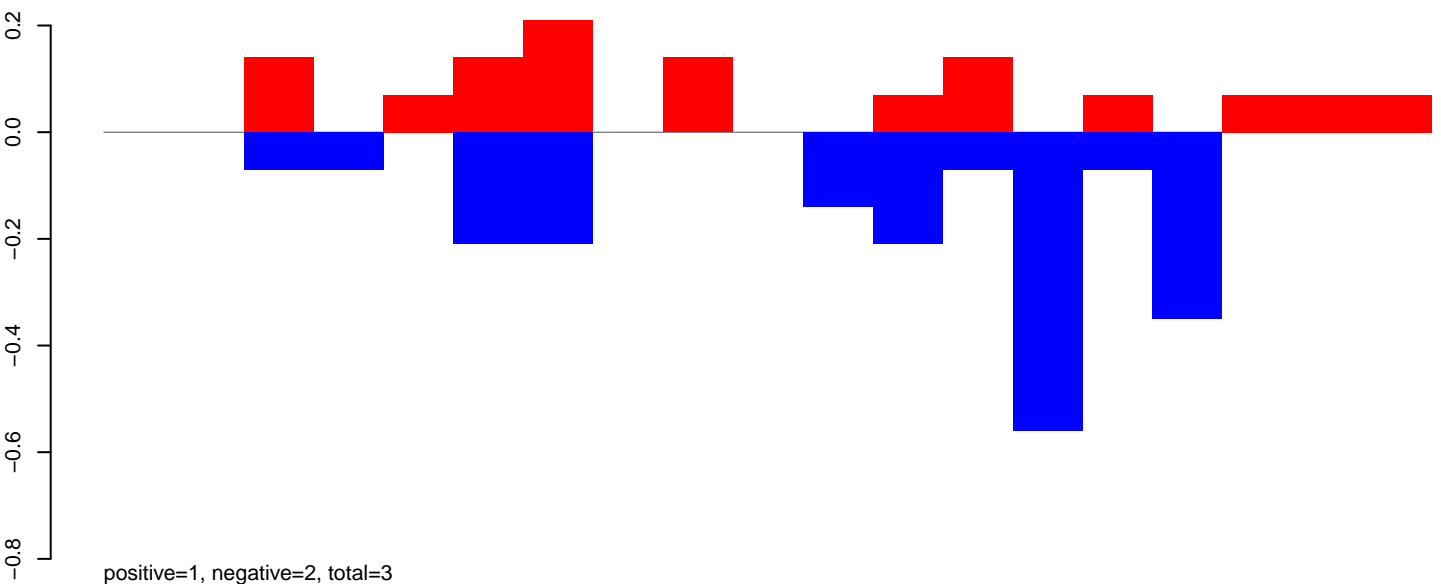
CuQuin_Mcarc_NL.24_35.rep



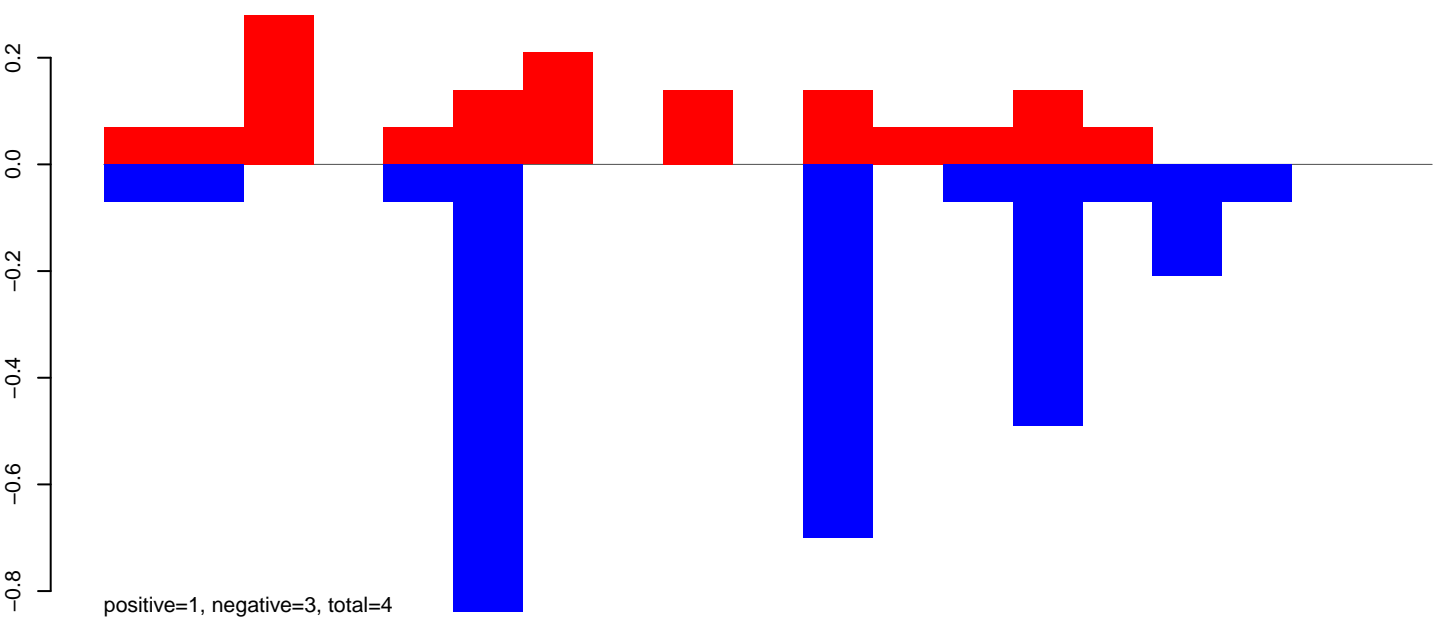
CuQuin_Mcarc_NL.rep



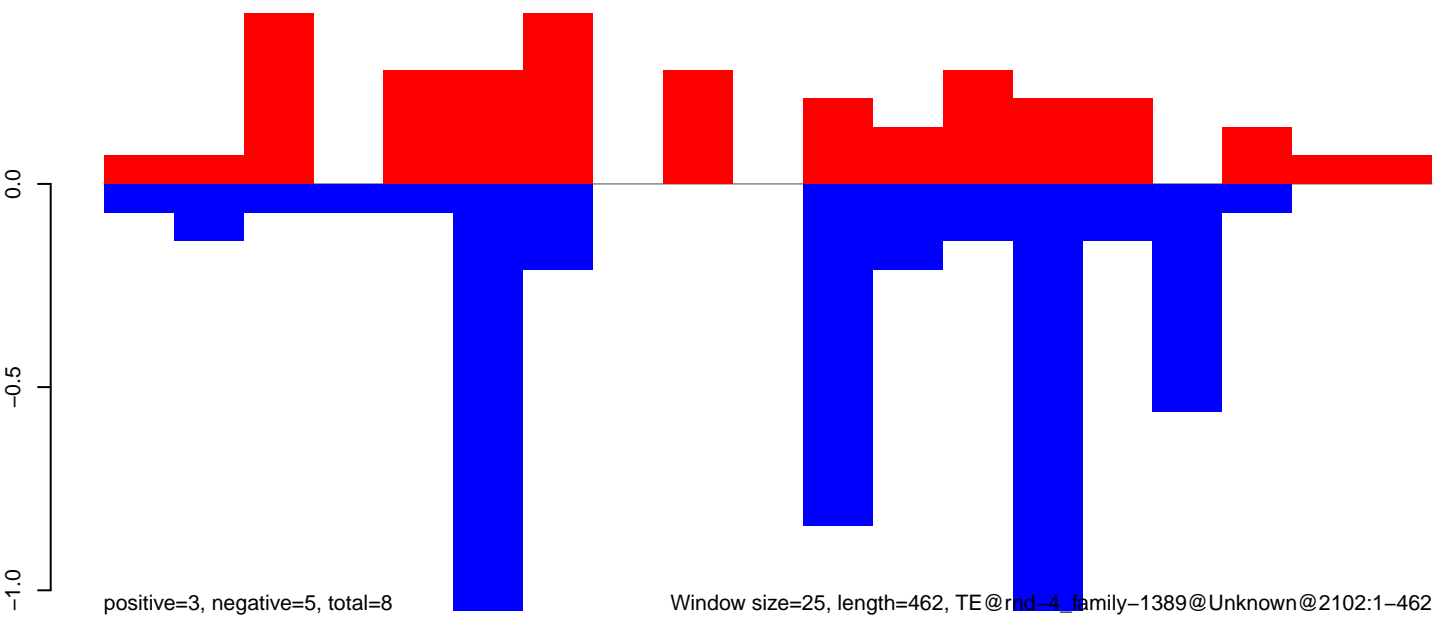
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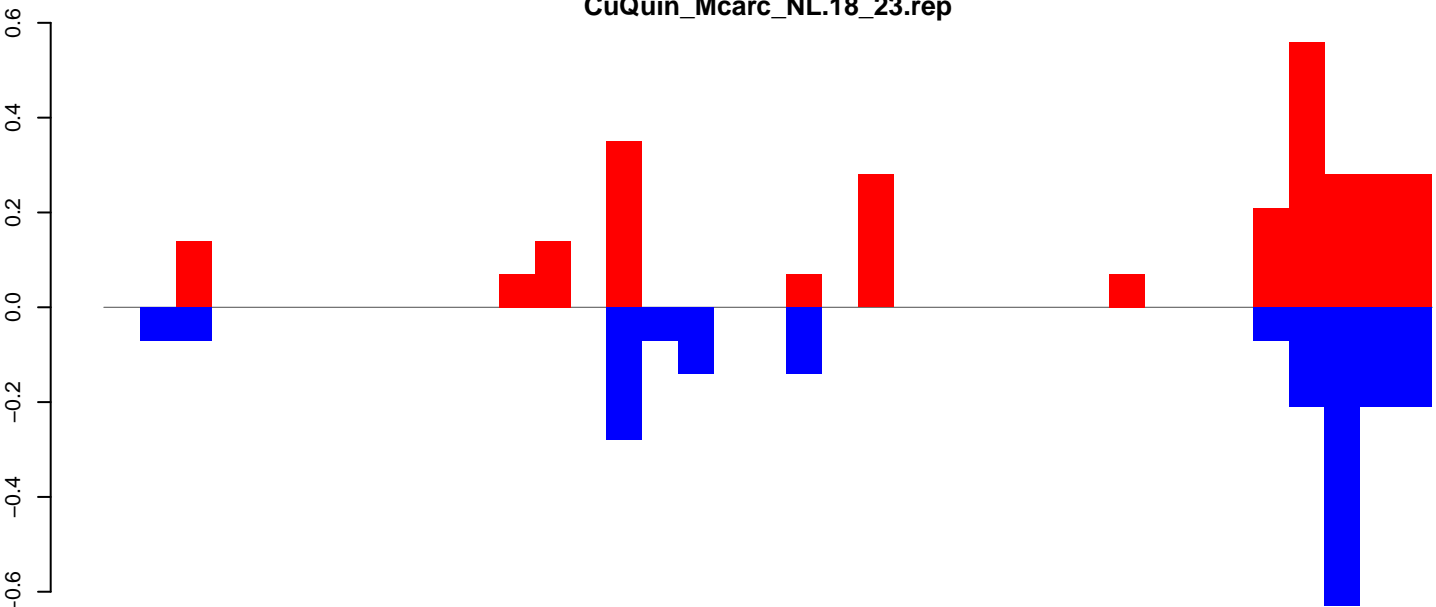
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CuQuin_Mcarc_NL.rep

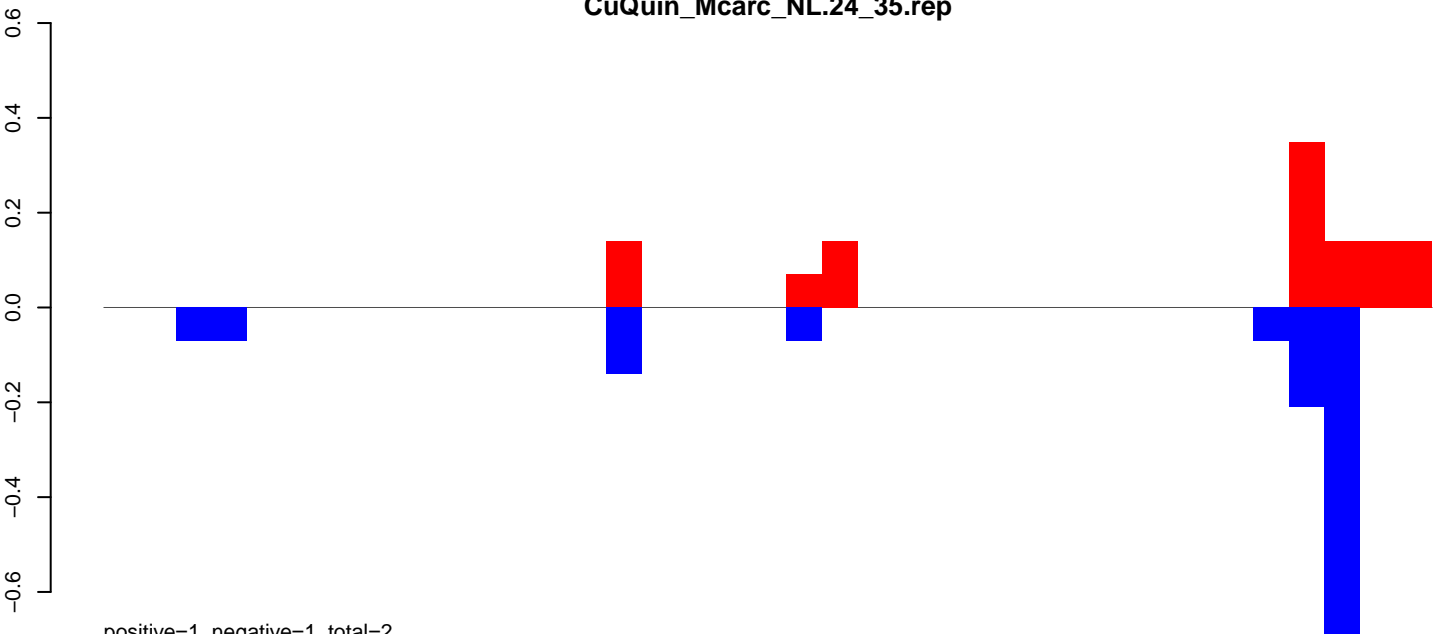


CuQuin_Mcarc_NL.18_23.rep



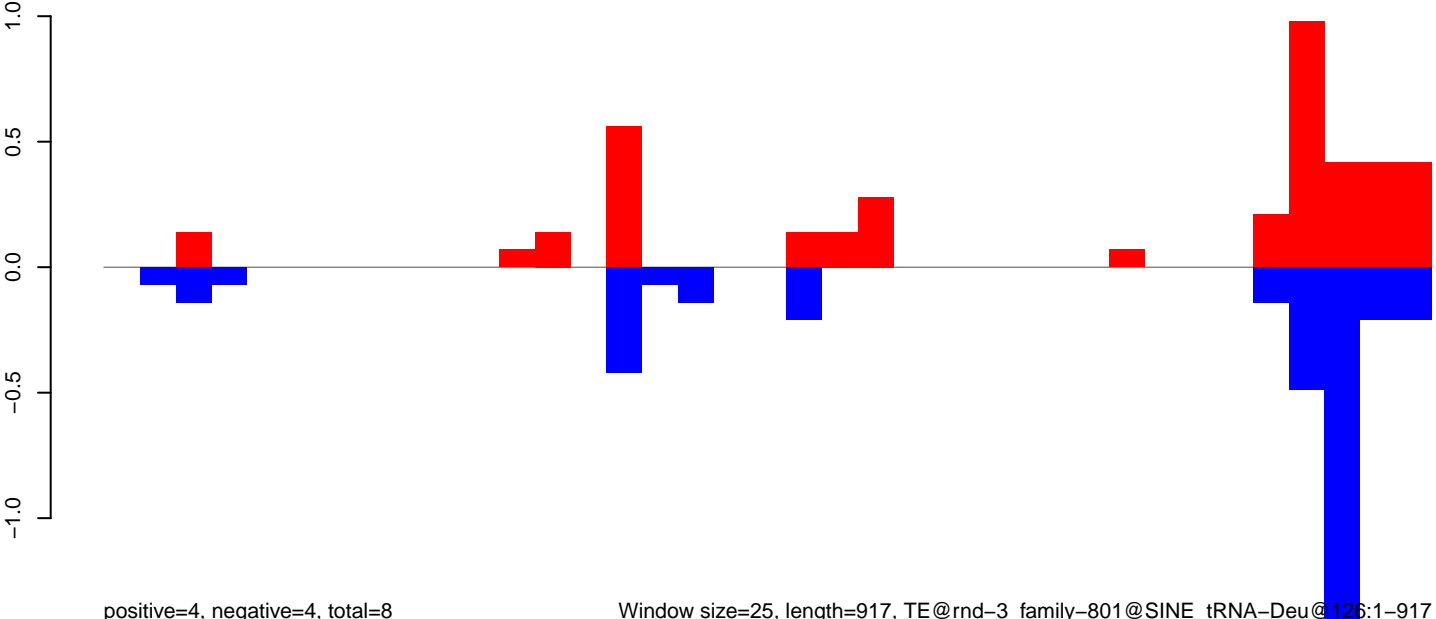
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CuQuin_Mcarc_NL.24_35.rep



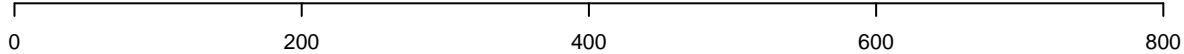
positive=1, negative=1, total=2

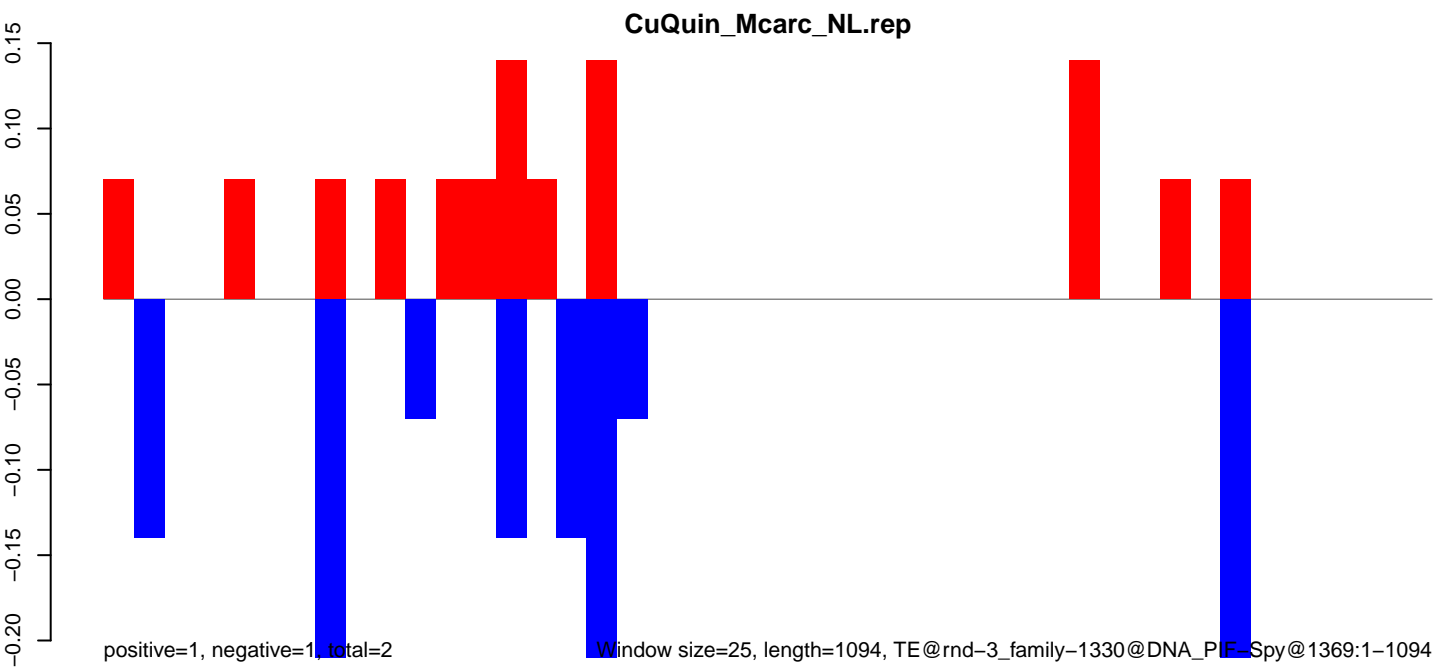
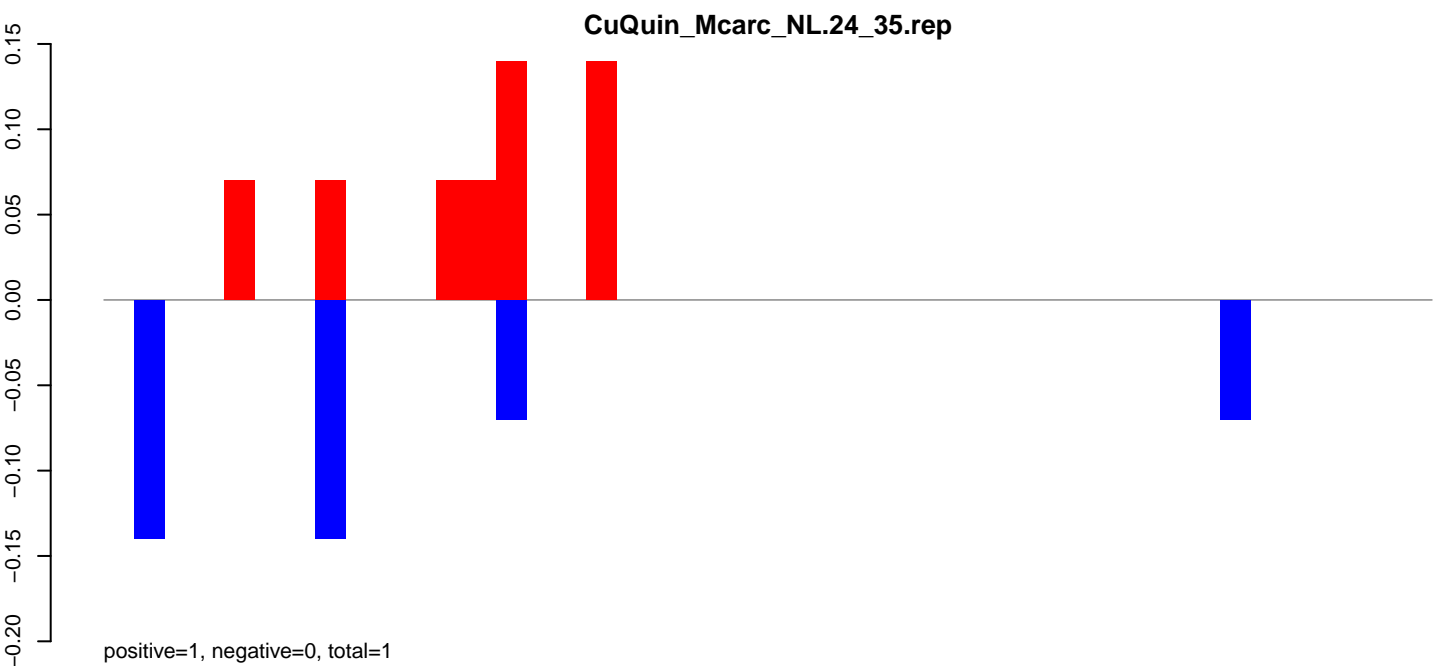
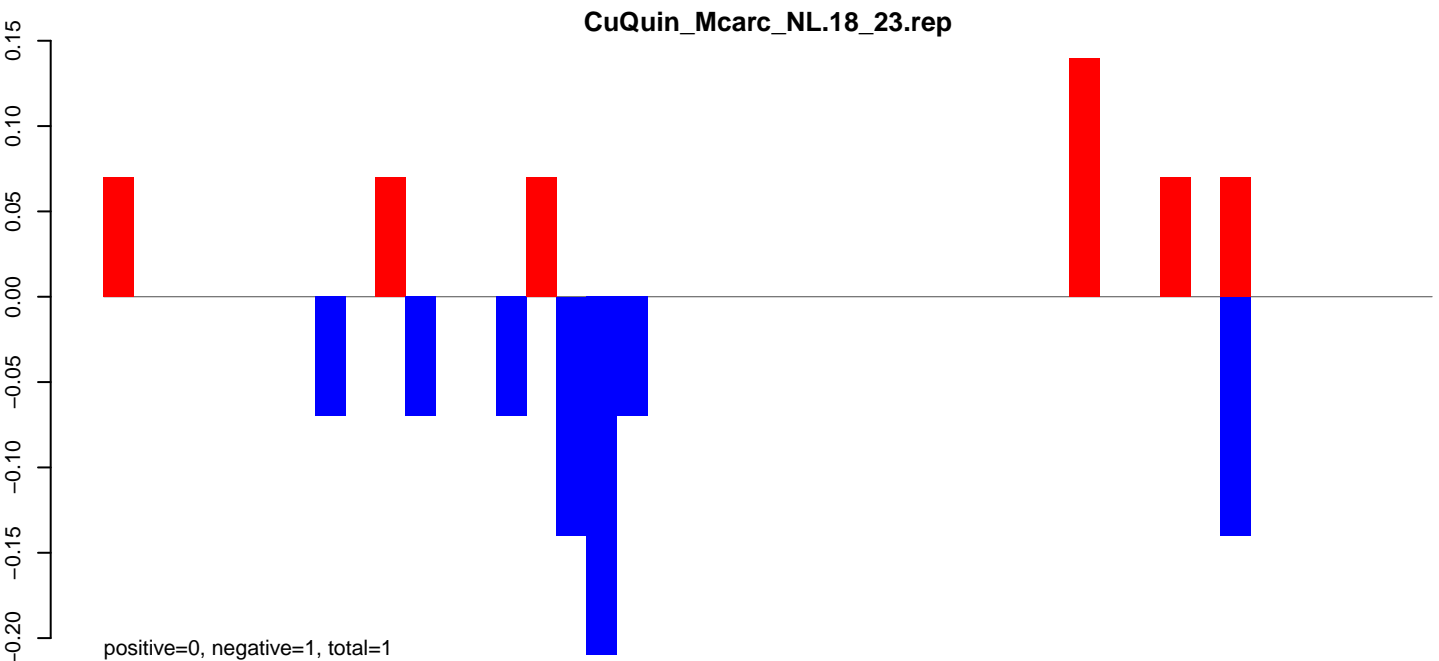
CuQuin_Mcarc_NL.rep



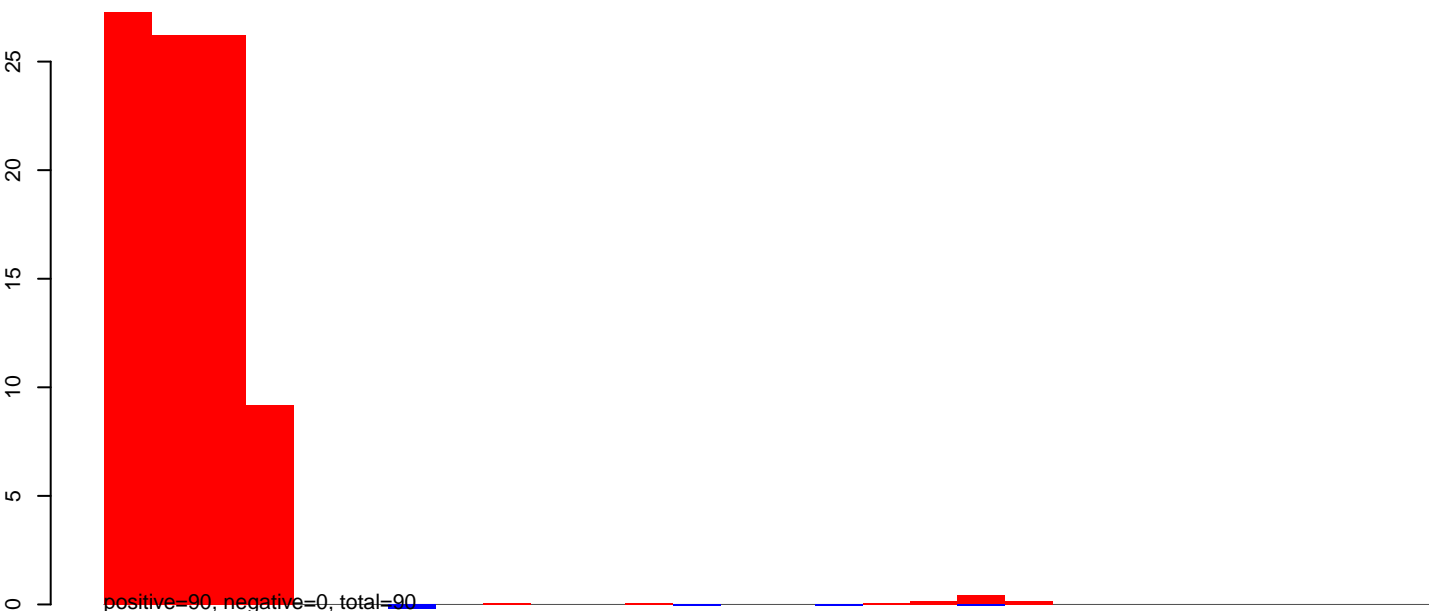
positive=4, negative=4, total=8

Window size=25, length=917, TE@rnd-3_family-801@SINE_tRNA-Deu@126:1-917

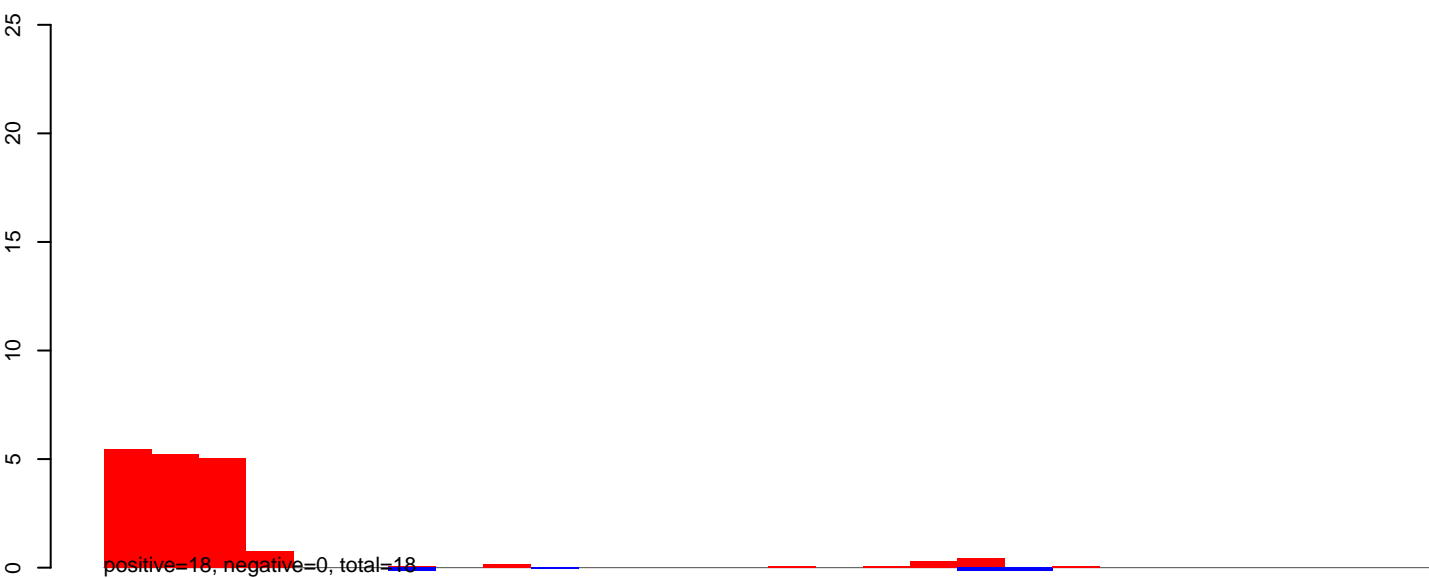




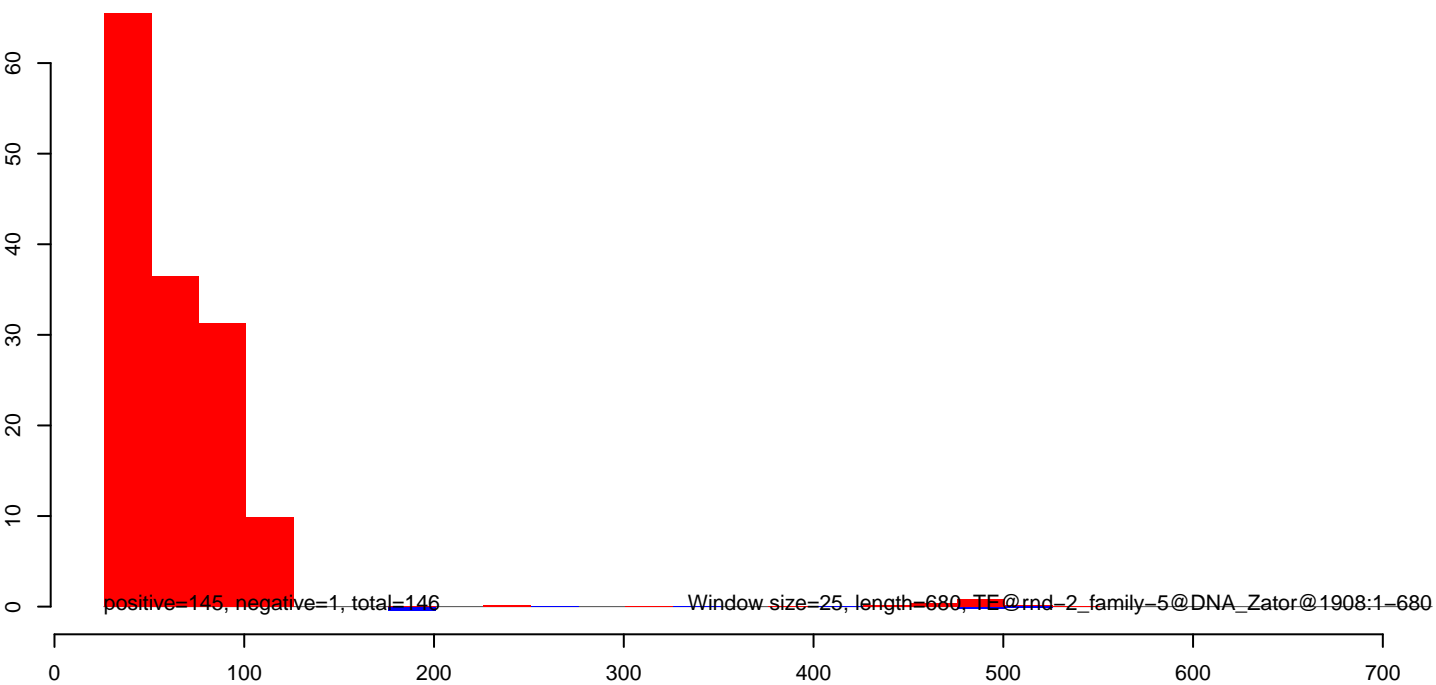
CuQuin_Mcarc_NL.18_23.rep



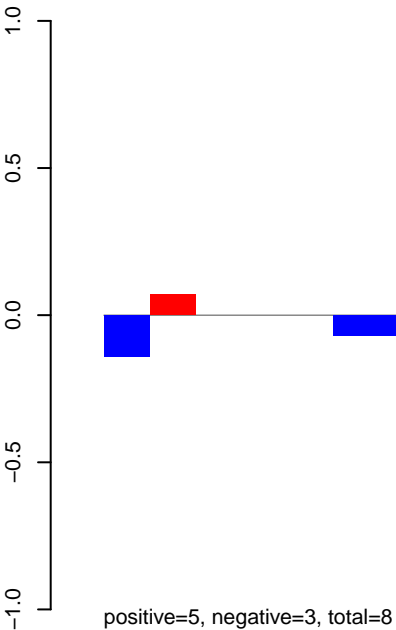
CuQuin_Mcarc_NL.24_35.rep



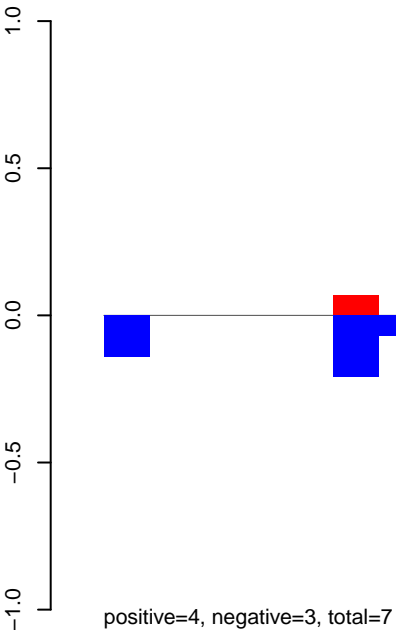
CuQuin_Mcarc_NL.rep



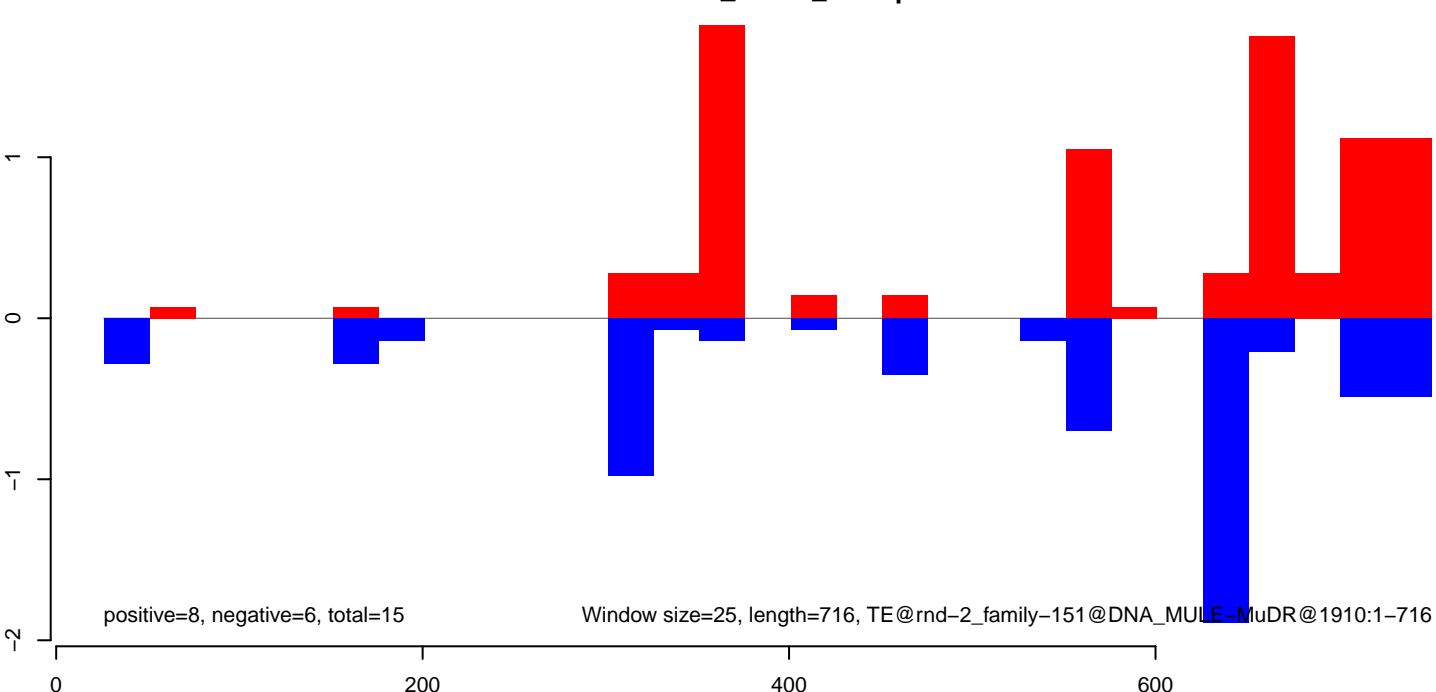
CuQuin_Mcarc_NL.18_23.rep



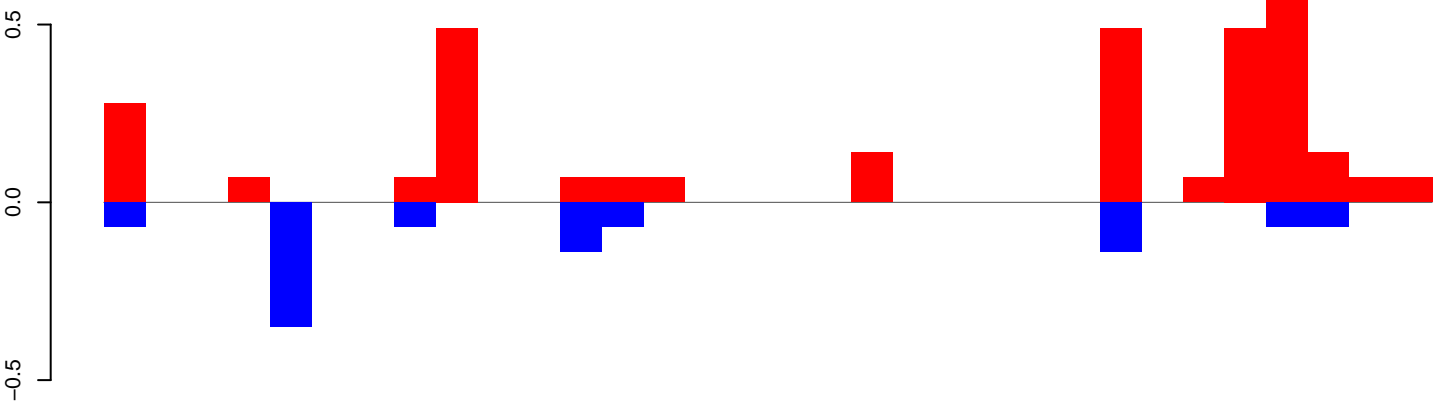
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

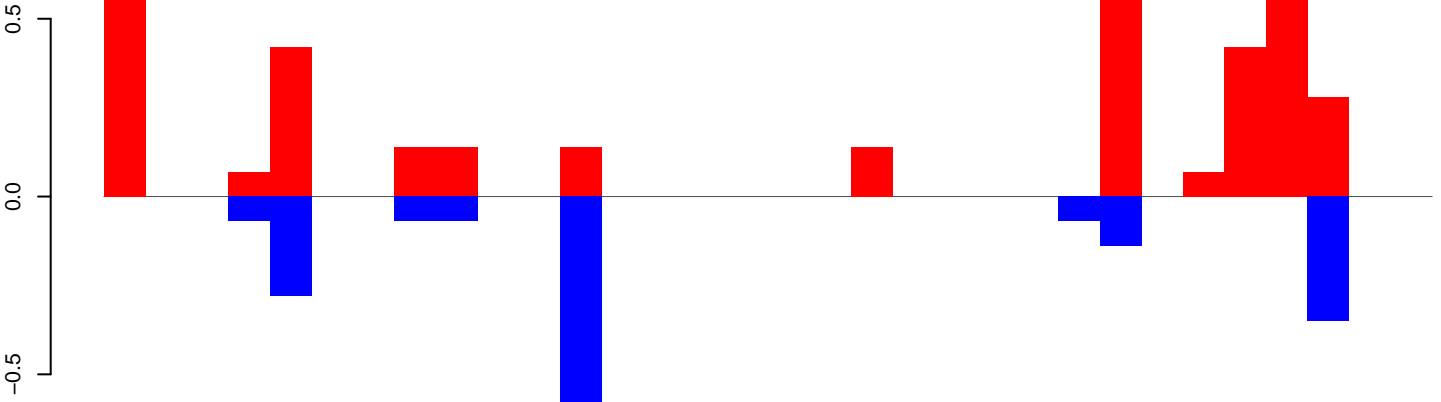


CuQuin_Mcarc_NL.18_23.rep



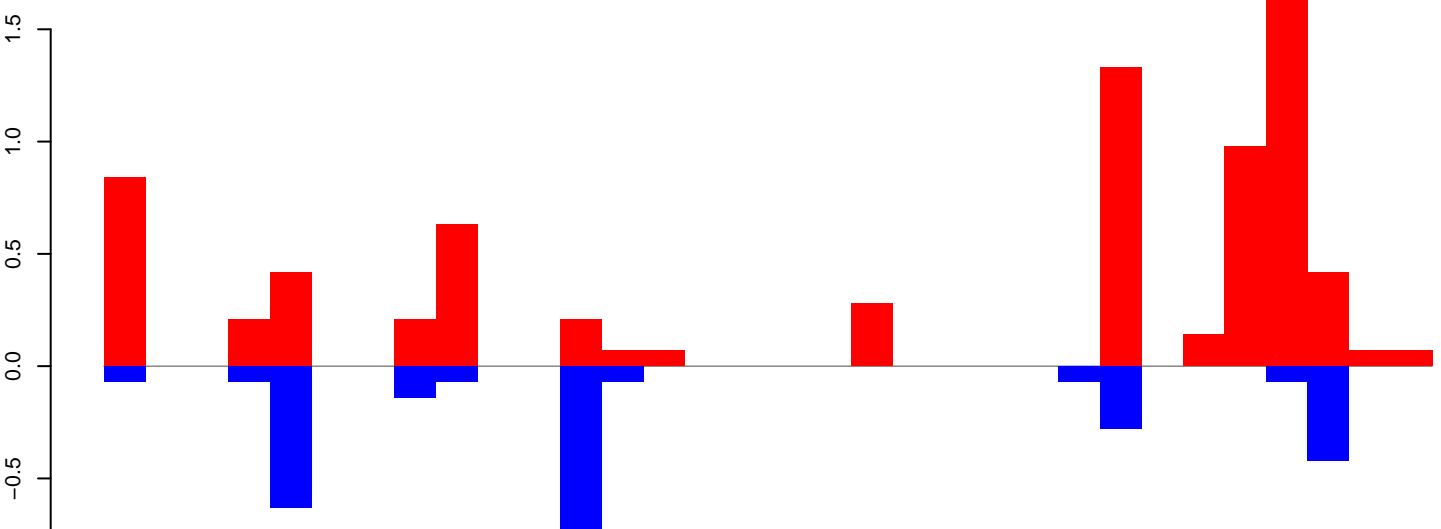
positive=3, negative=1, total=4

CuQuin_Mcarc_NL.24_35.rep



positive=4, negative=2, total=6

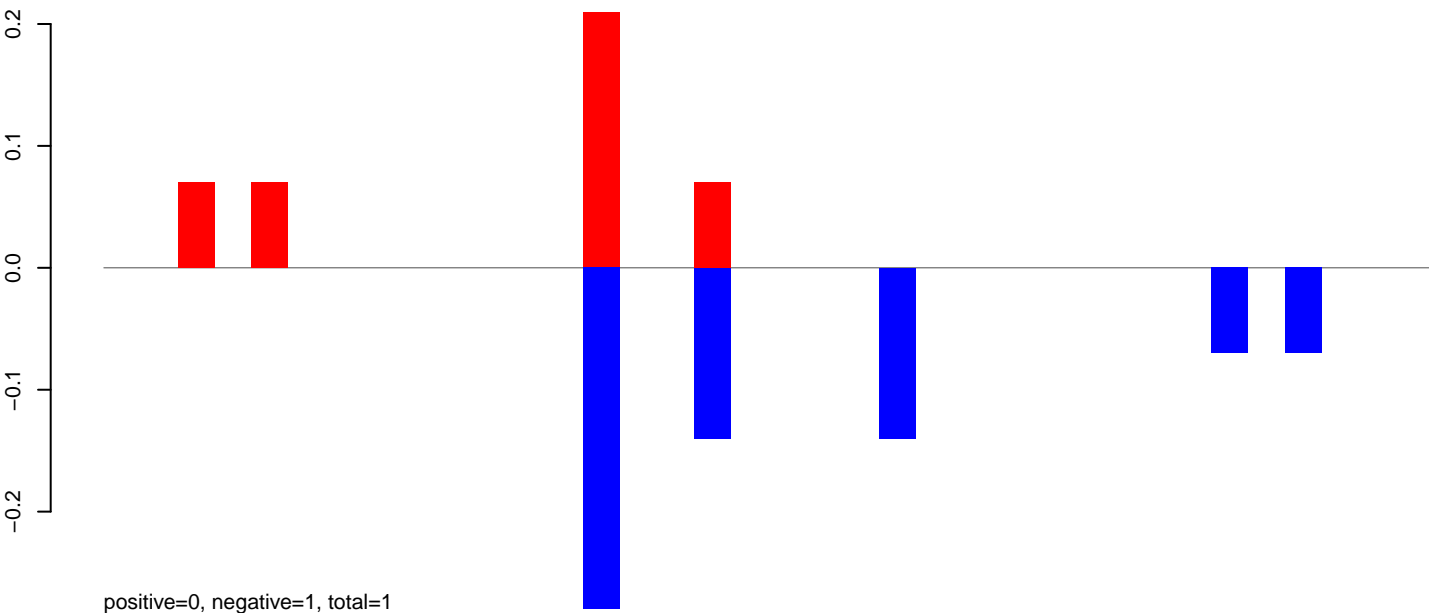
CuQuin_Mcarc_NL.rep



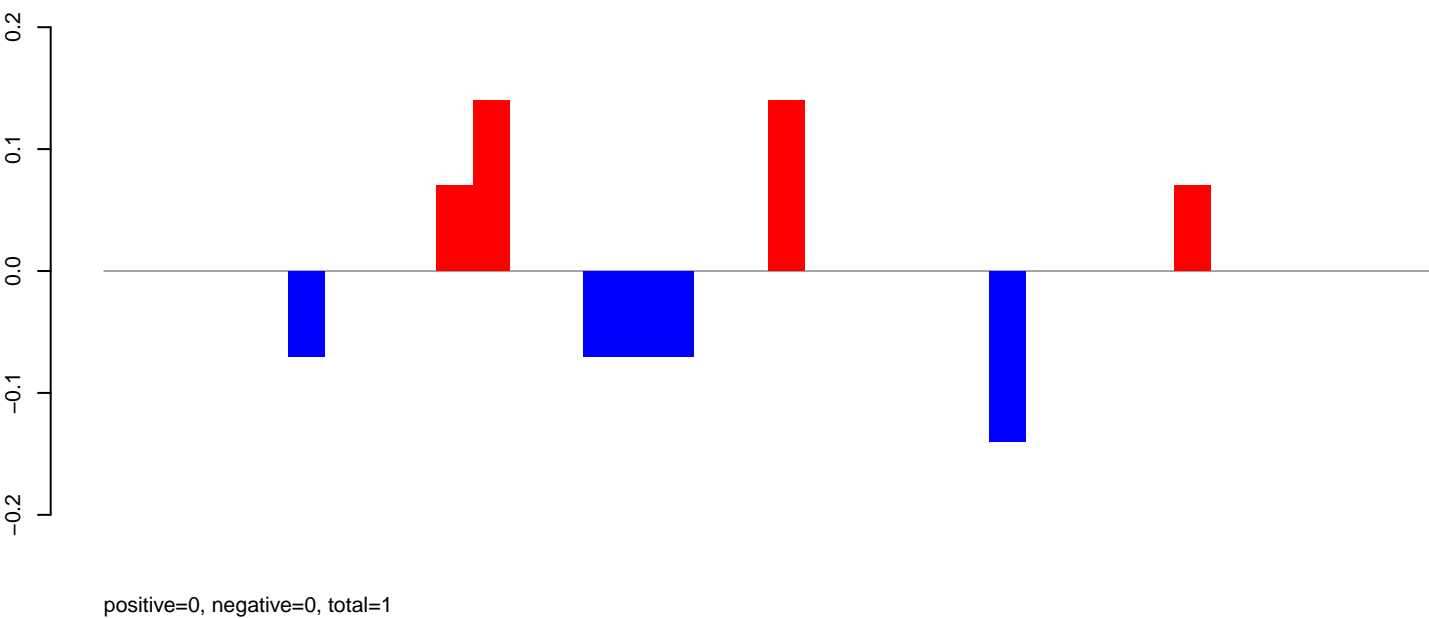
positive=8, negative=3, total=10

Window size=25, length=778, TE@rnd-2_family-143@DNA_Sola-3@27:1-778

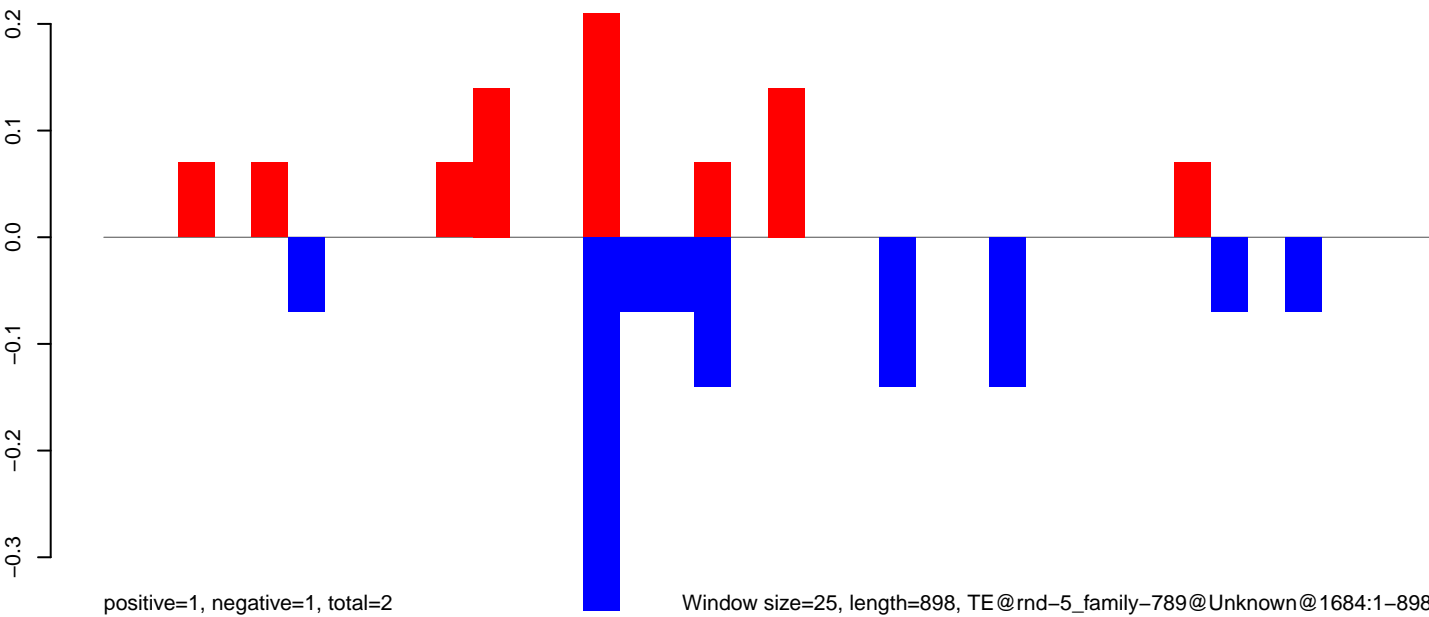
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

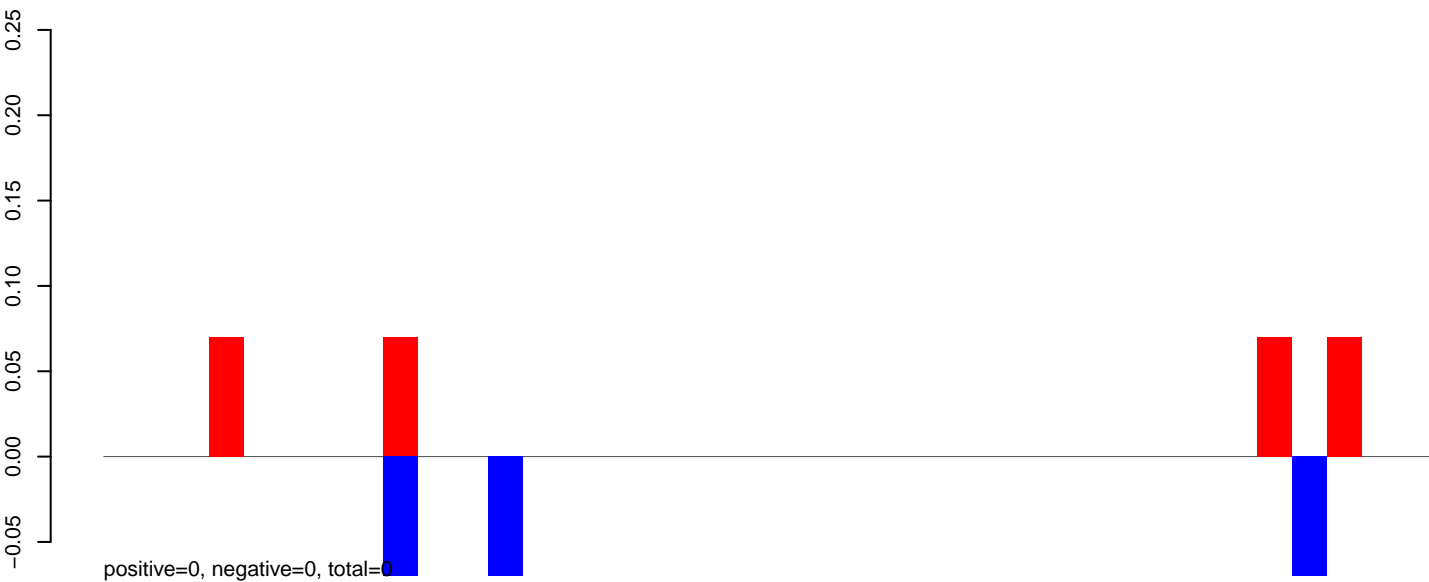


CuQuin_Mcarc_NL.rep

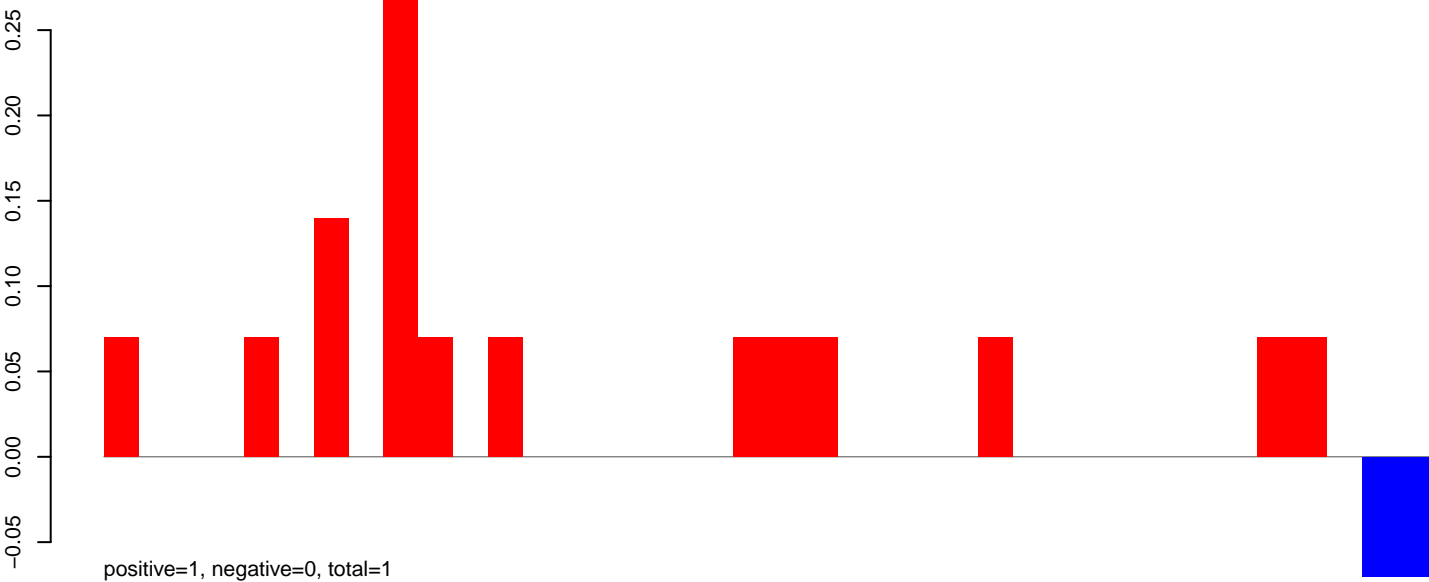


Window size=25, length=898, TE@rnd-5_family-789@Unknown@1684:1-898

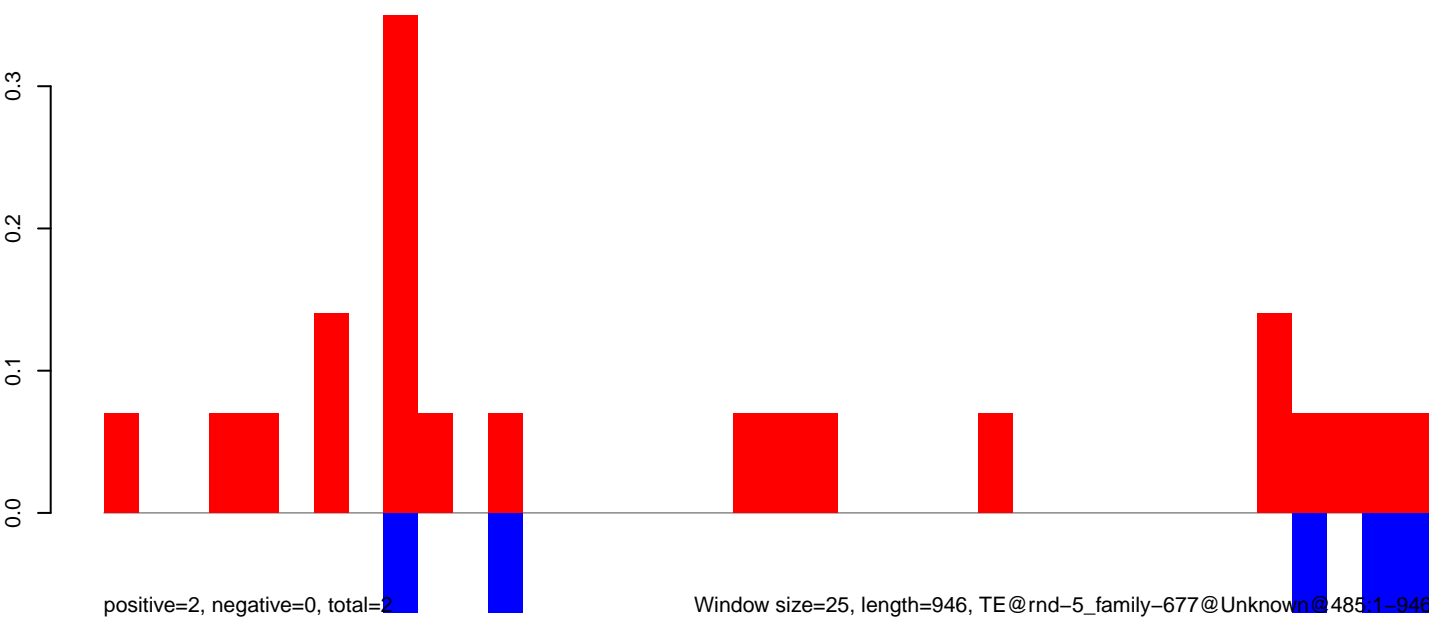
CuQuin_Mcarc_NL.18_23.rep



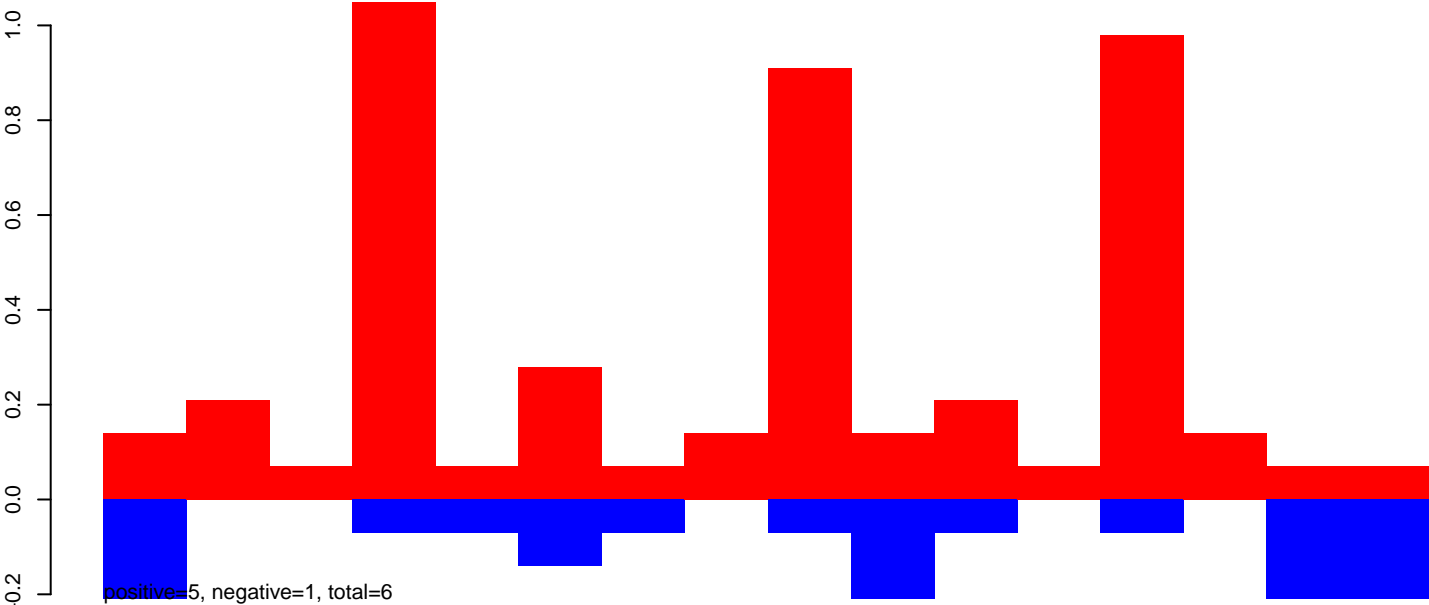
CuQuin_Mcarc_NL.24_35.rep



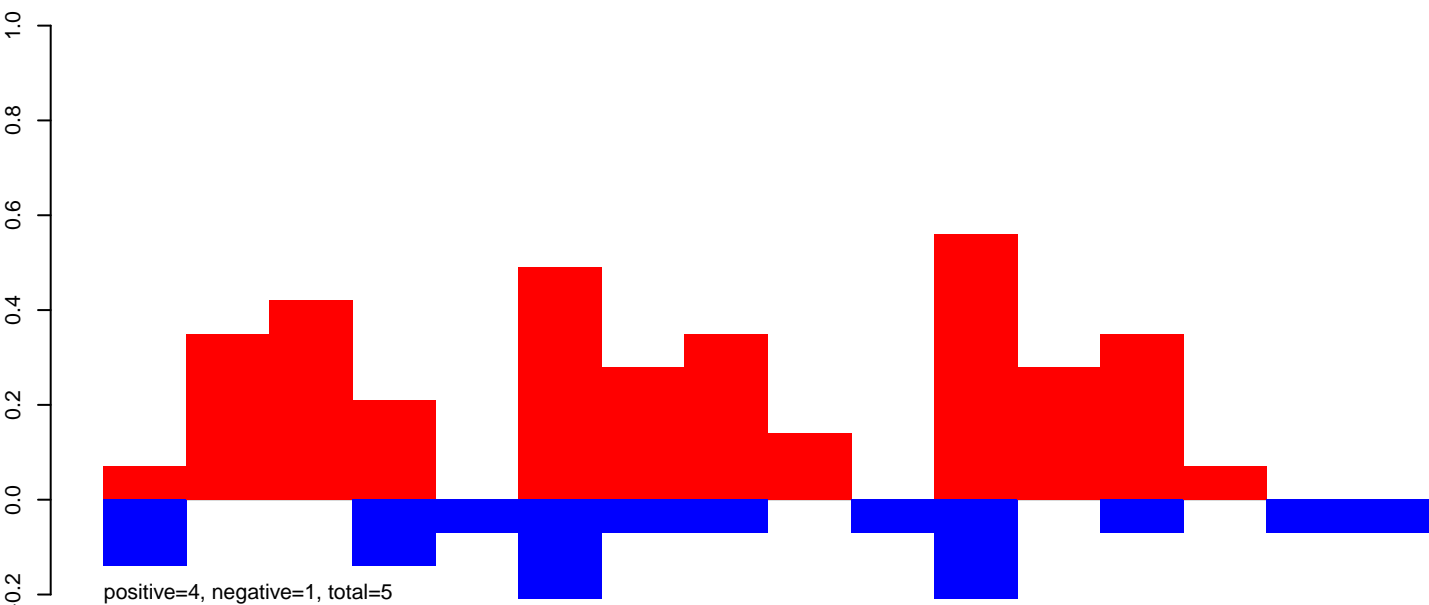
CuQuin_Mcarc_NL.rep



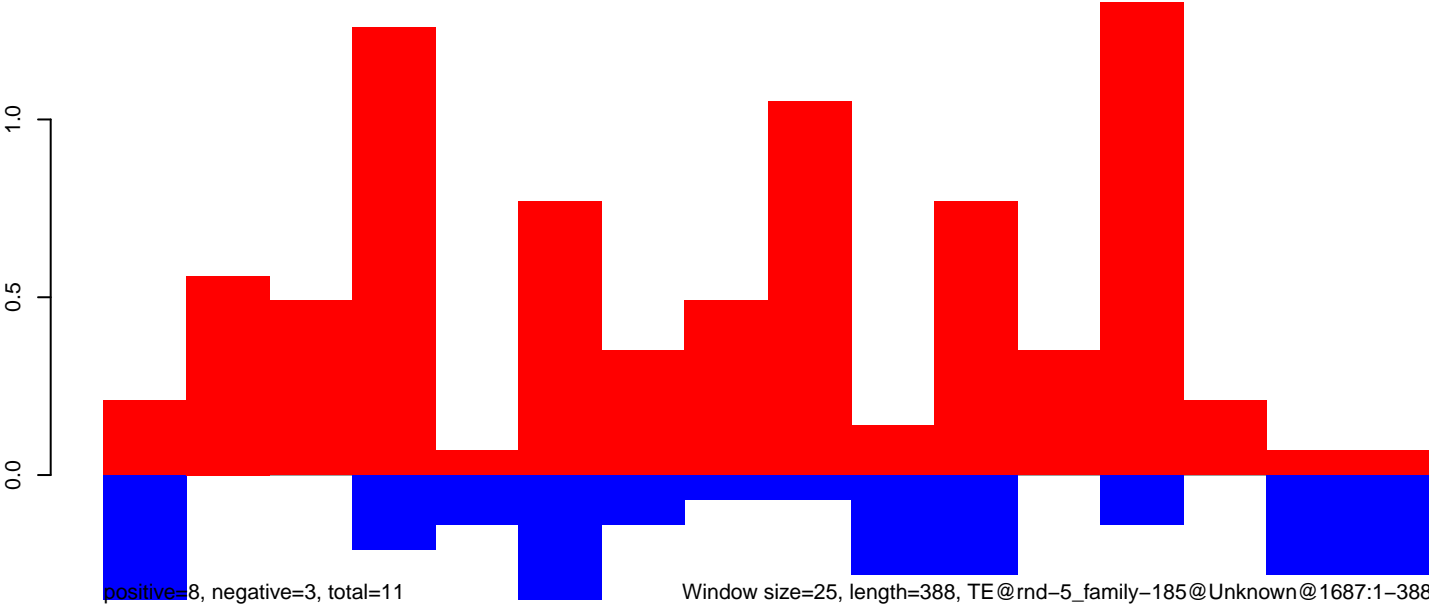
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

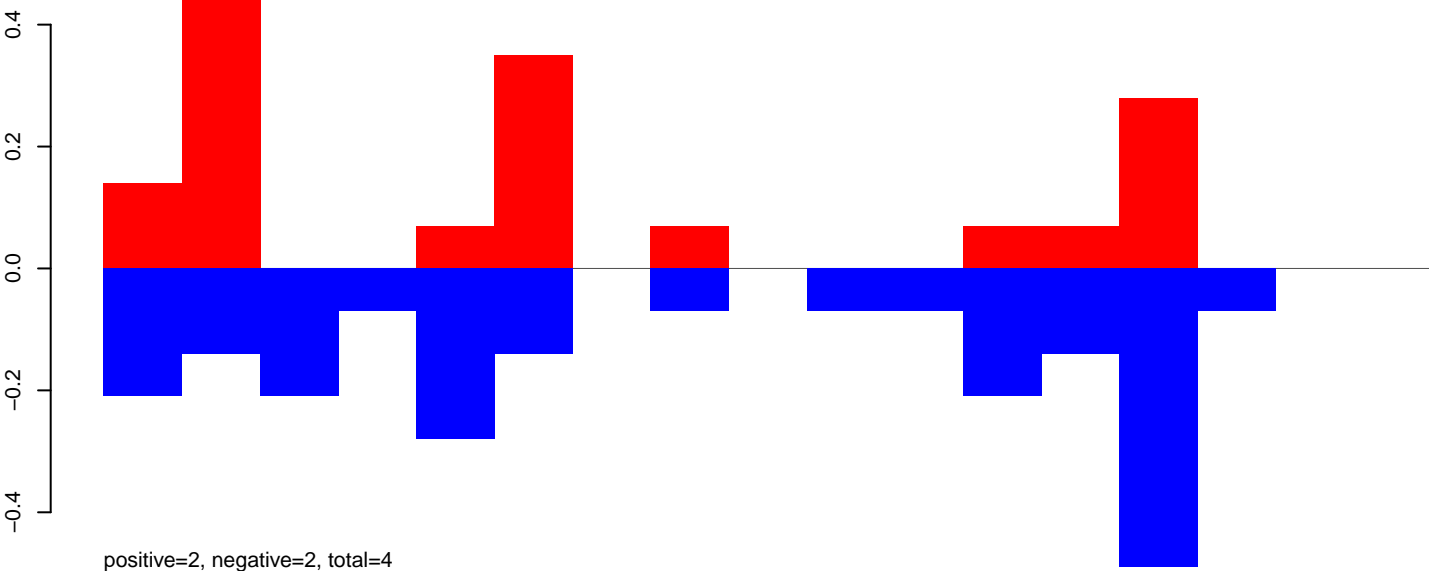


CuQuin_Mcarc_NL.rep

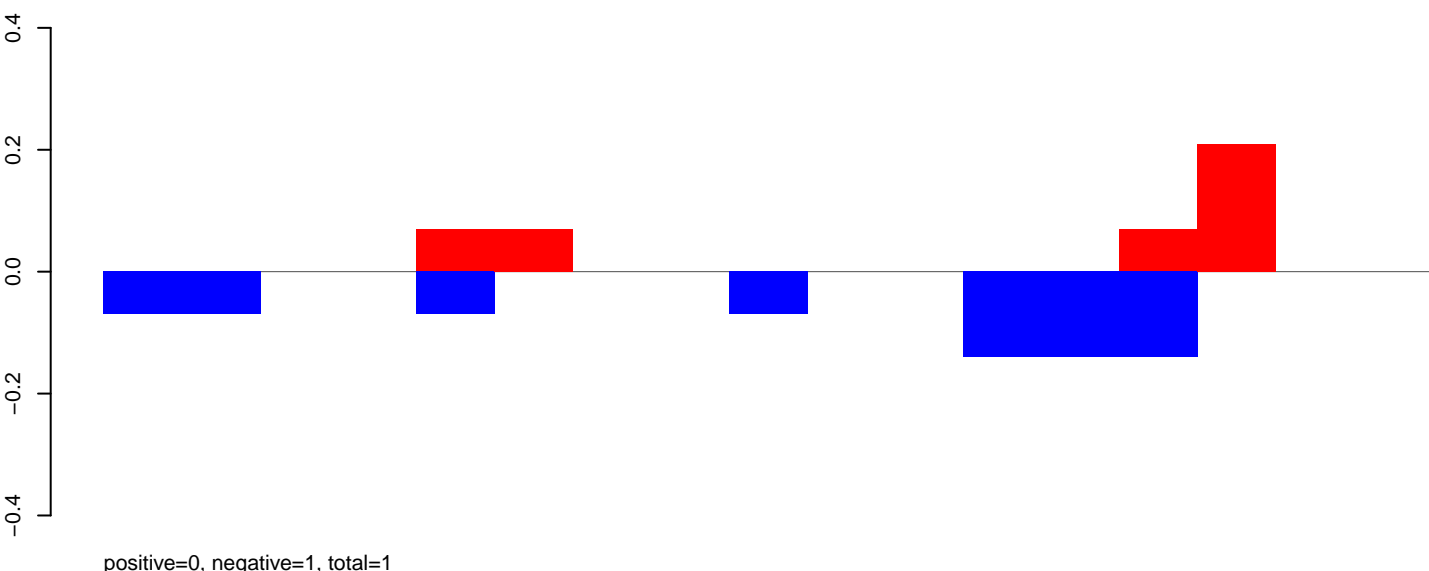


Window size=25, length=388, TE@rnd-5_family-185@Unknown@1687:1-388

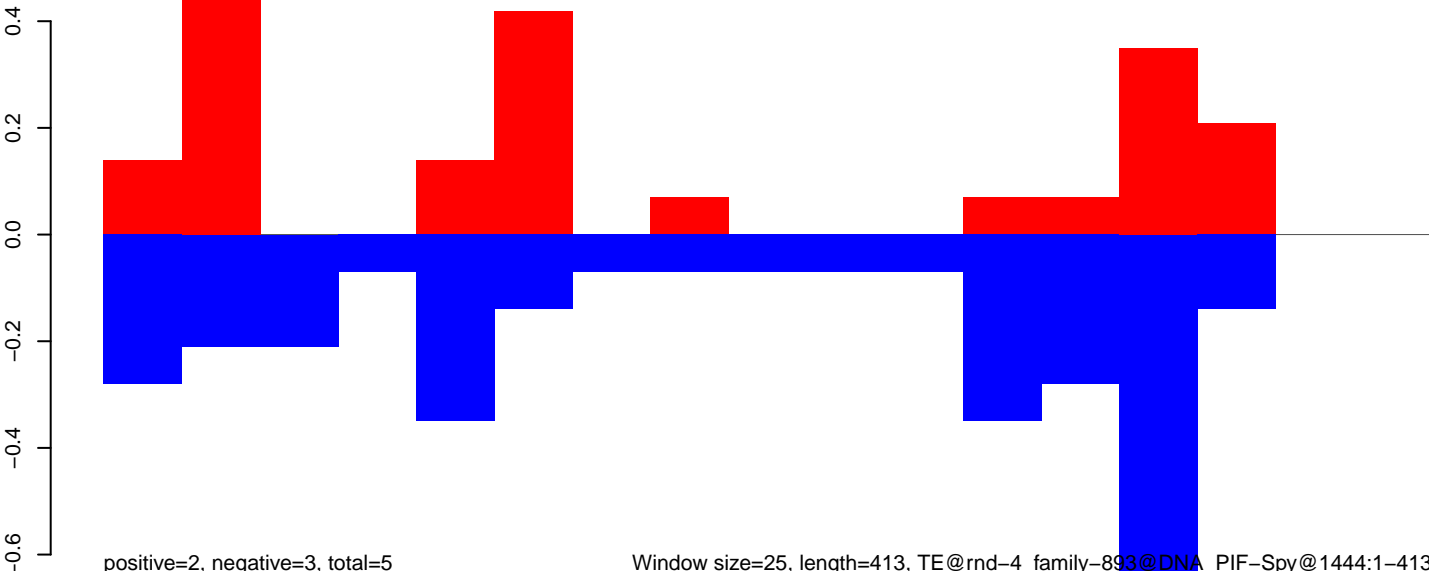
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=413, TE@rnd-4_family-893@DNA_PIF-Spy@1444:1-413

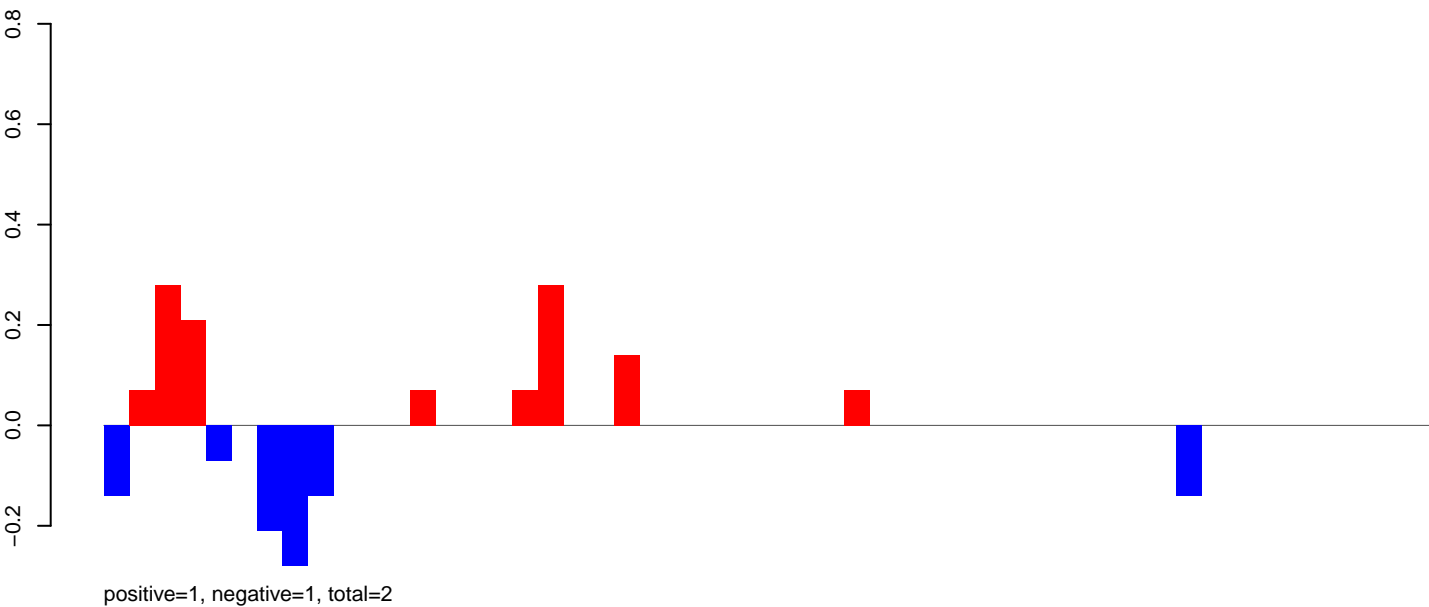
100

200

300

400

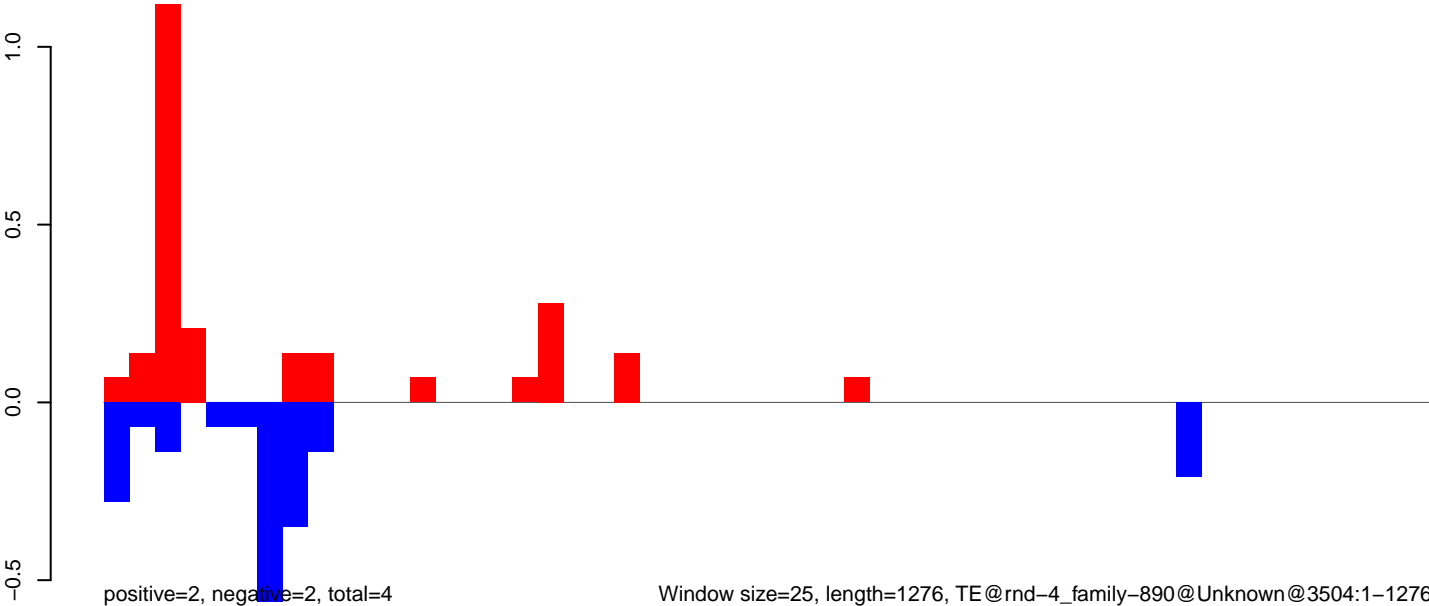
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

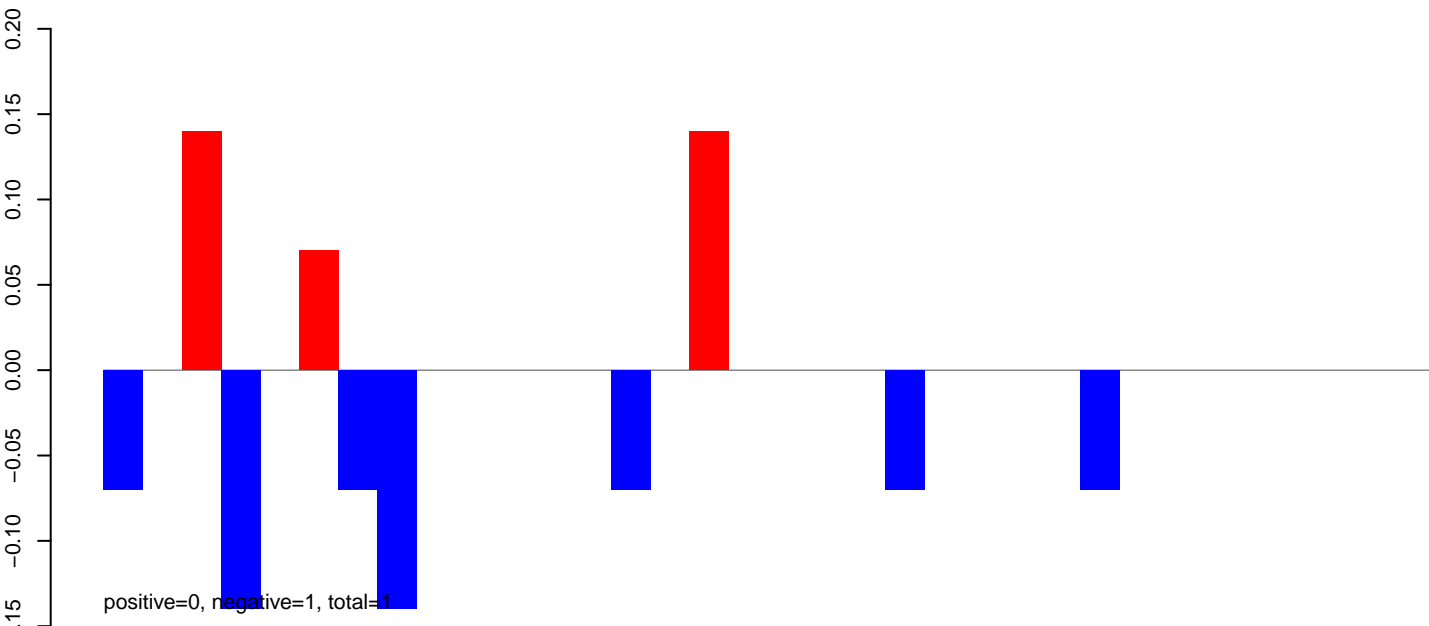


CuQuin_Mcarc_NL.rep

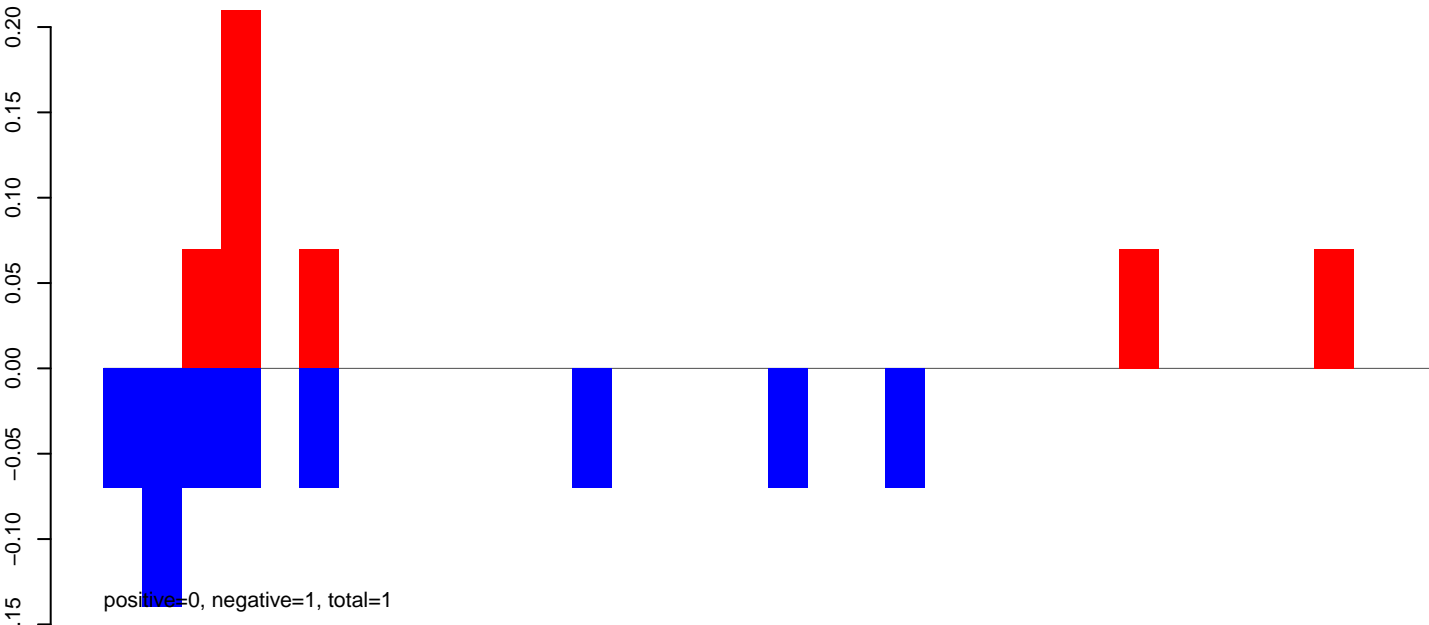


Window size=25, length=1276, TE@rnd-4_family-890@Unknown@3504:1-1276

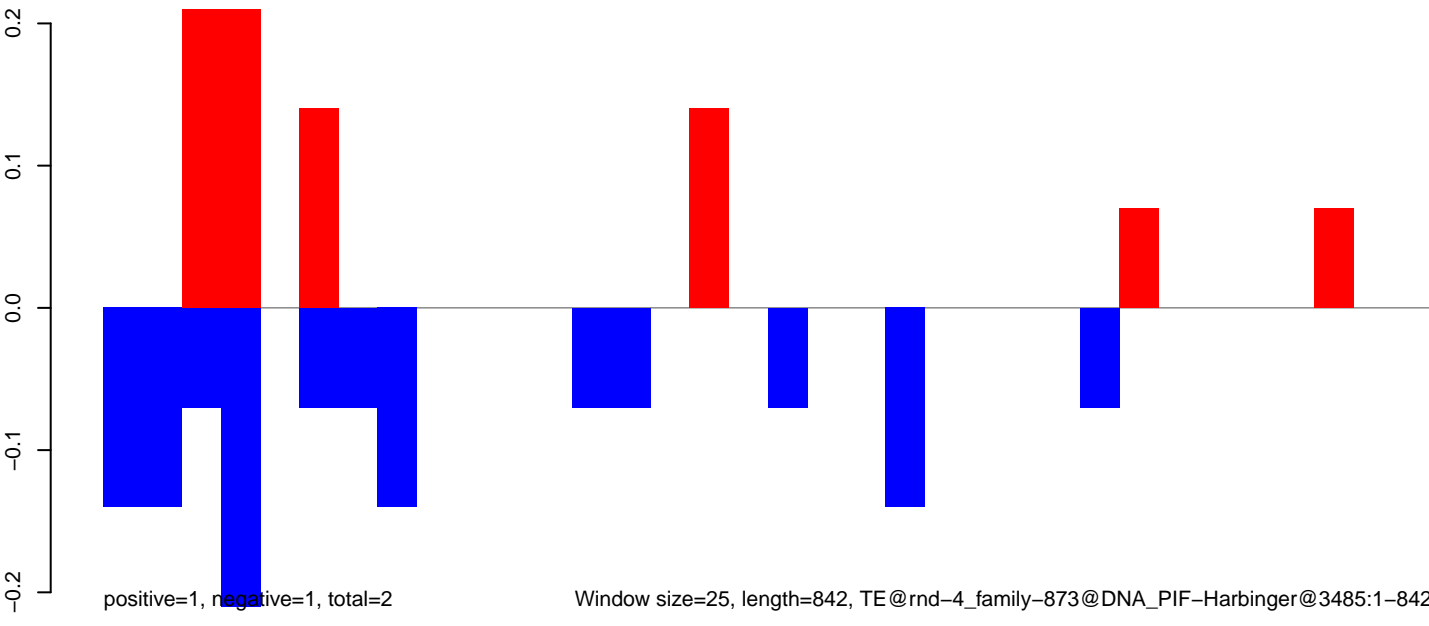
CuQuin_Mcarc_NL.18_23.rep



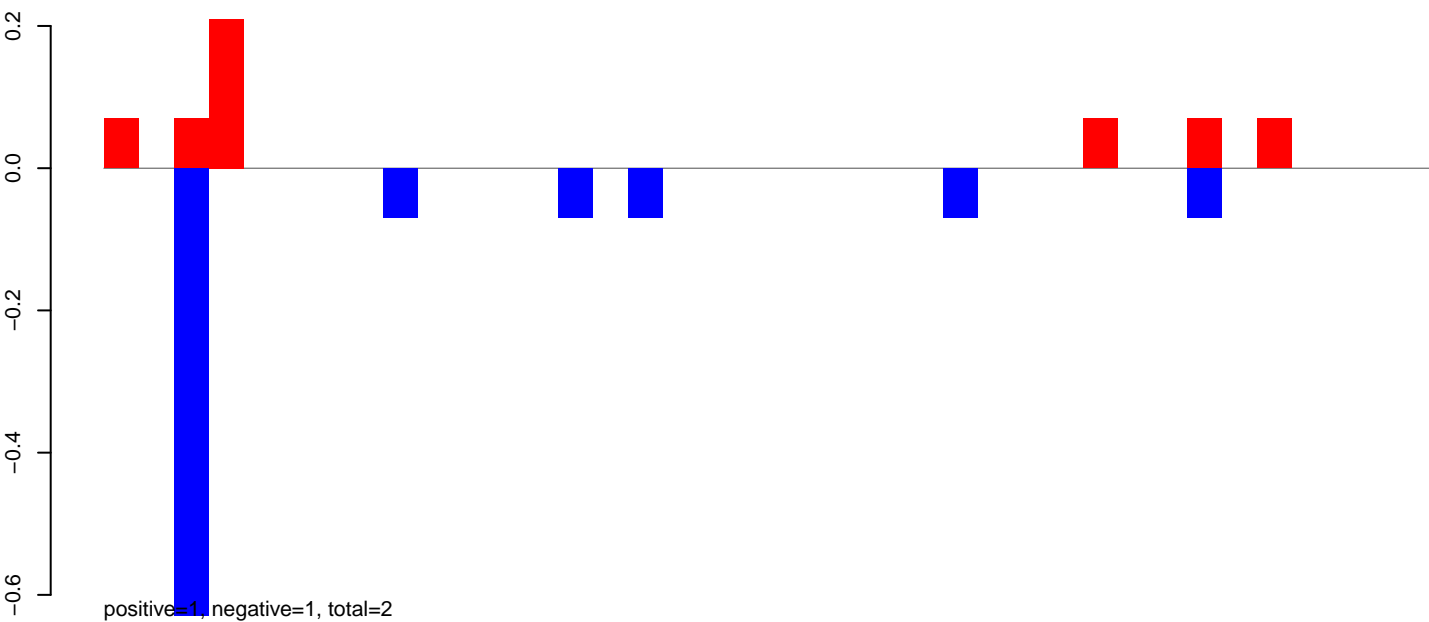
CuQuin_Mcarc_NL.24_35.rep



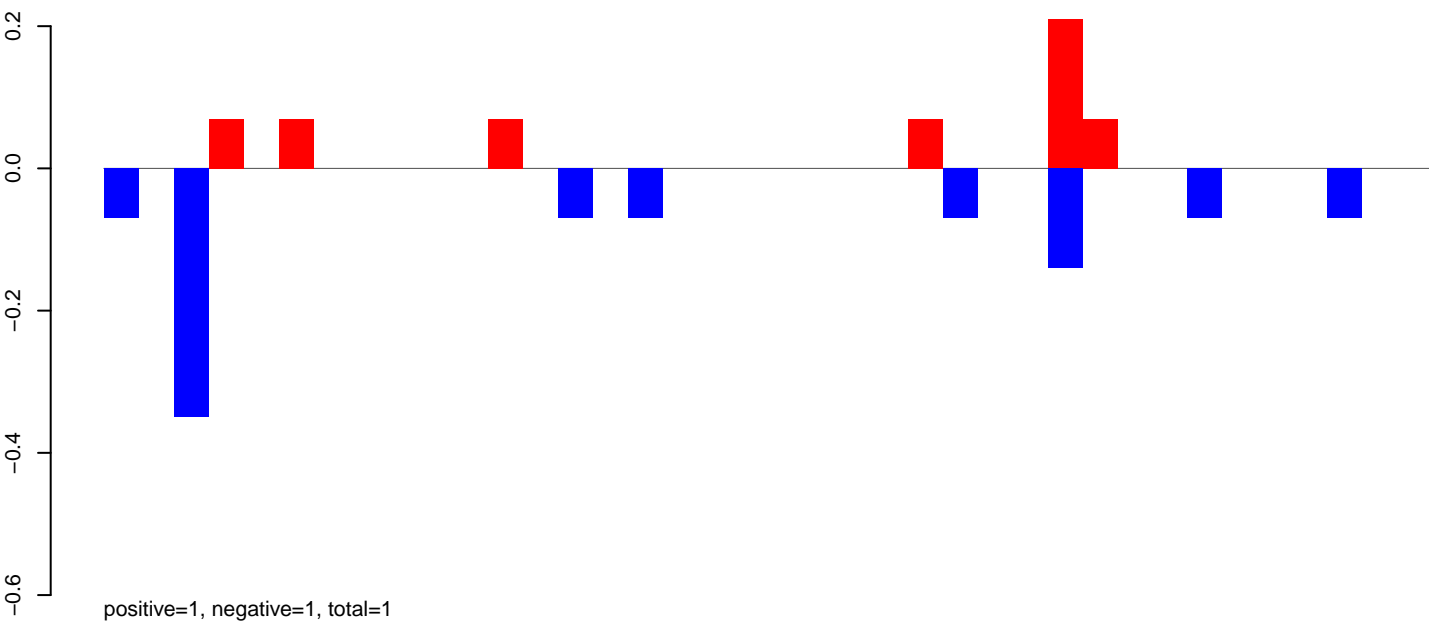
CuQuin_Mcarc_NL.rep



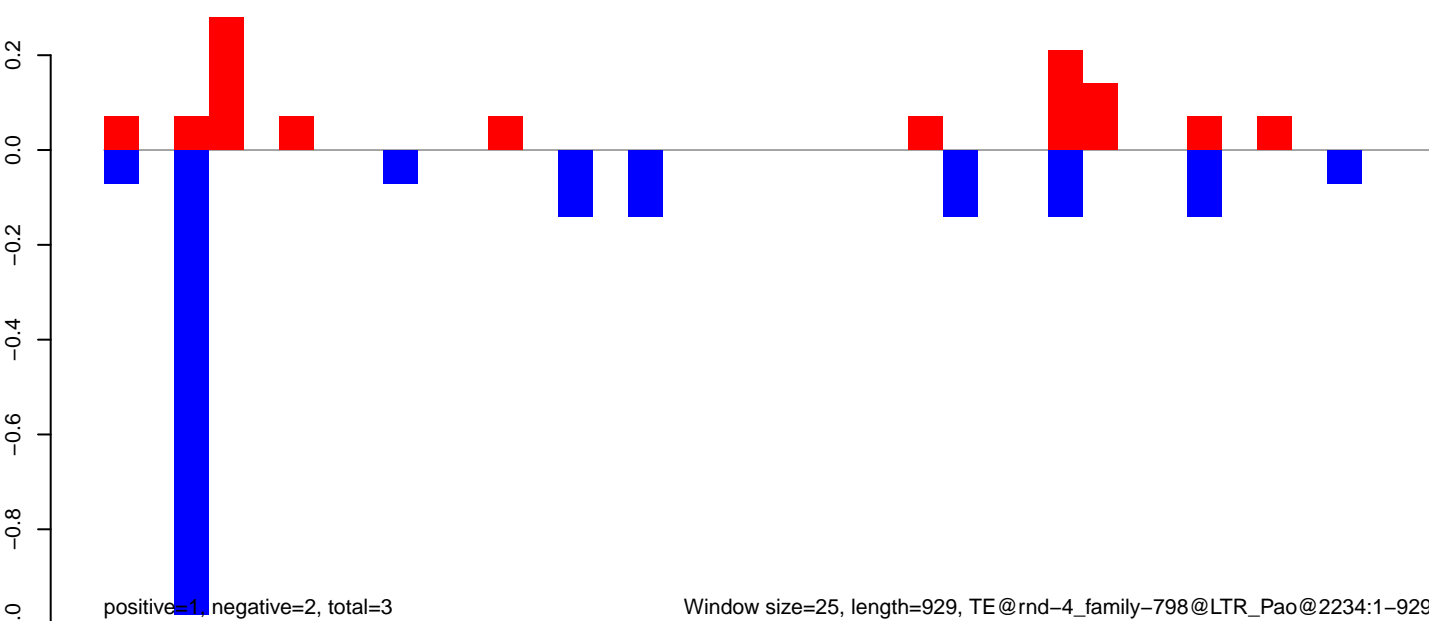
CuQuin_Mcarc_NL.18_23.rep



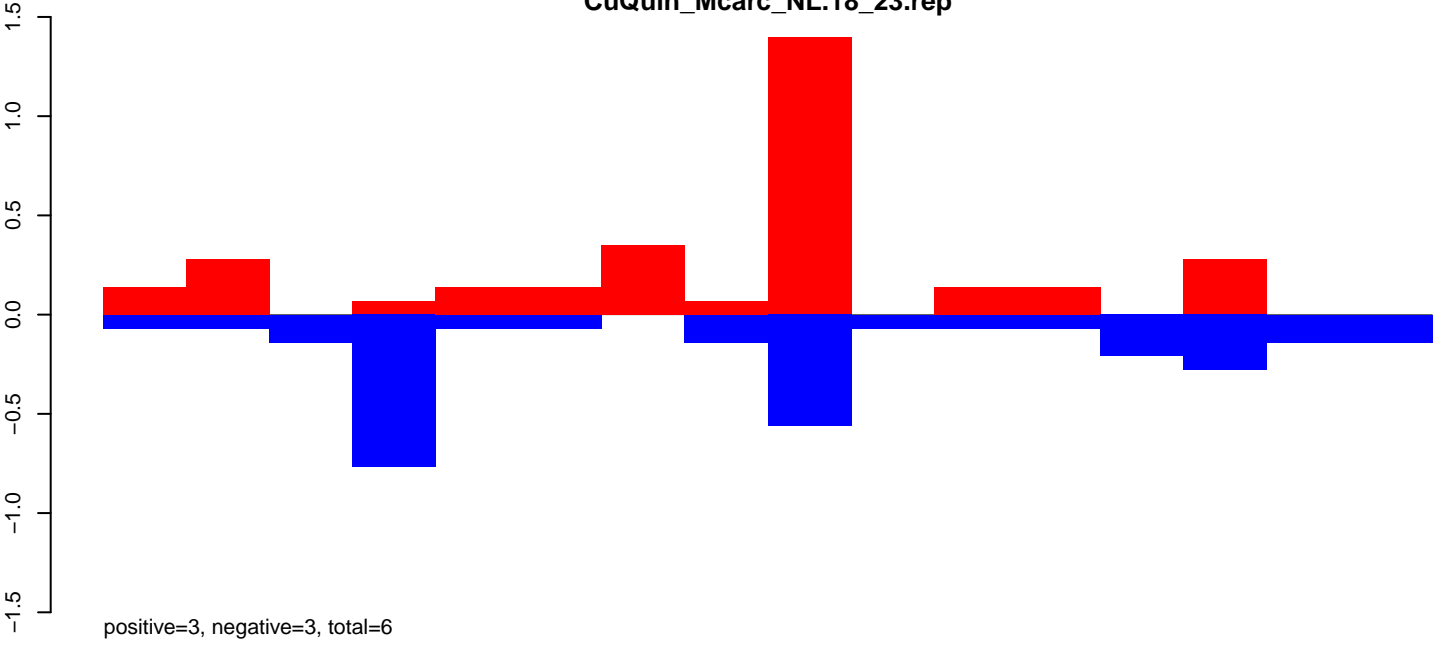
CuQuin_Mcarc_NL.24_35.rep



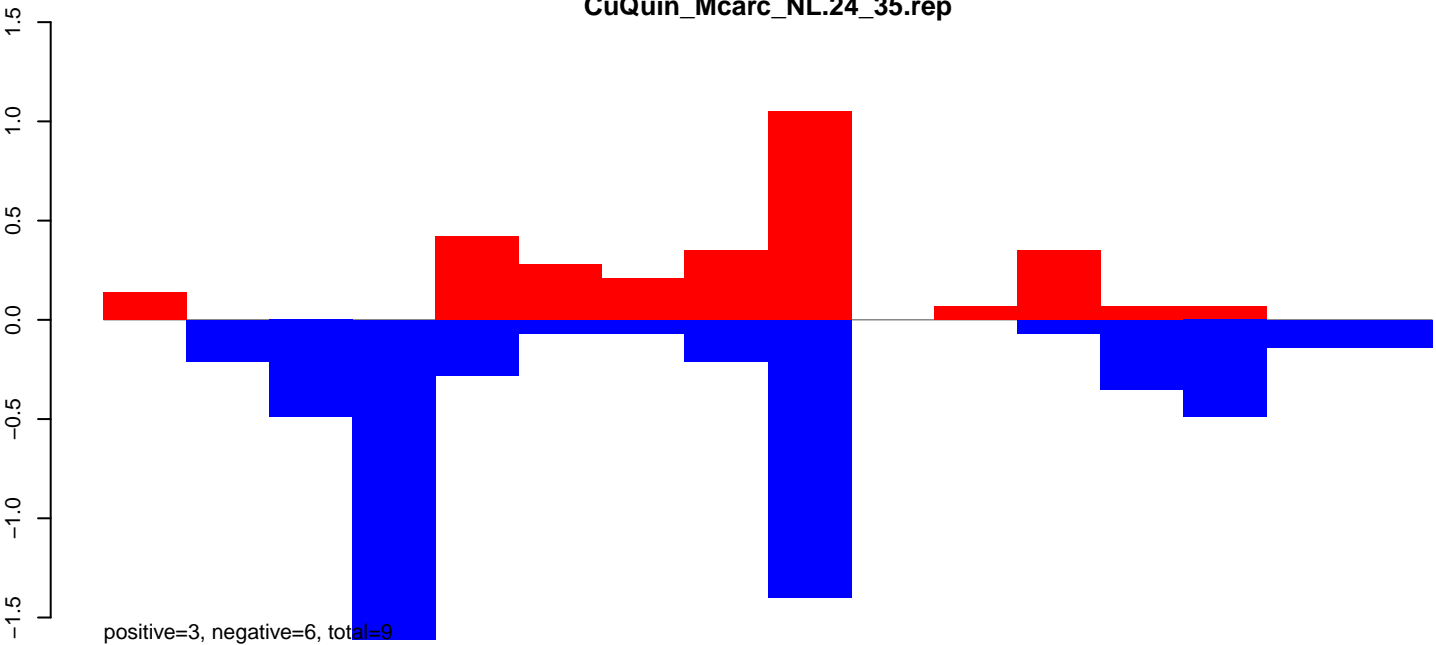
CuQuin_Mcarc_NL.rep



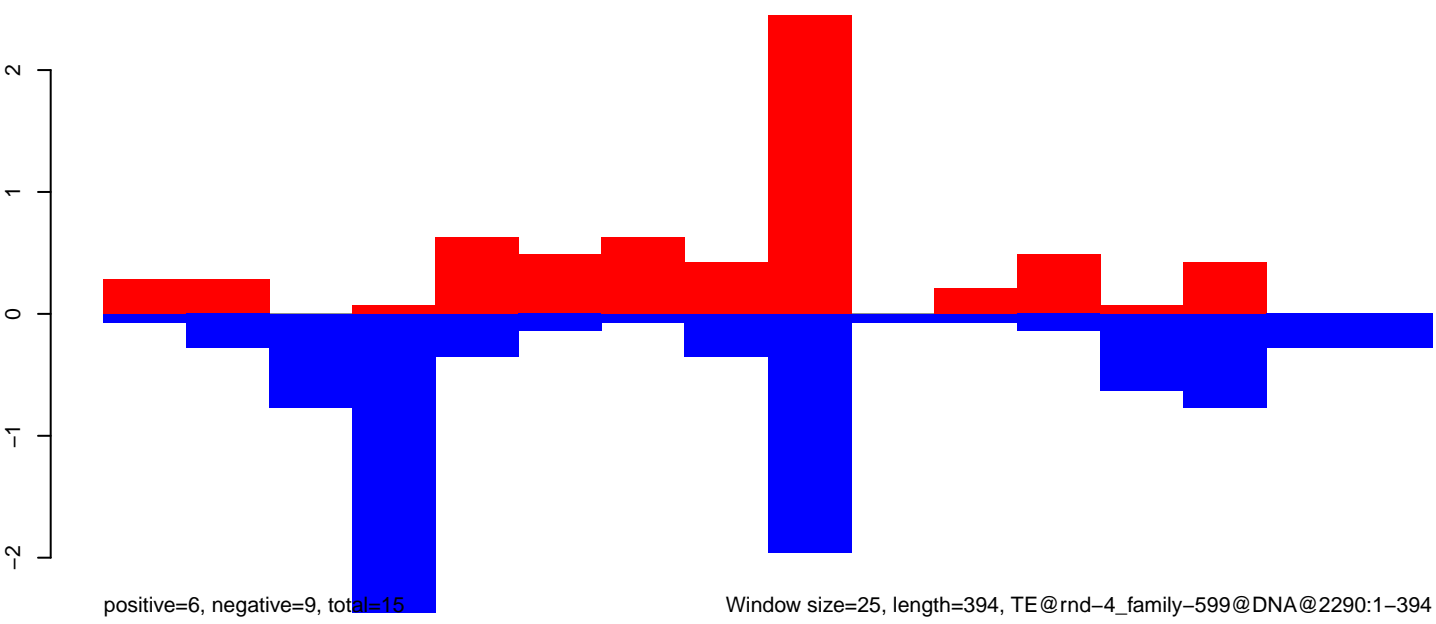
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=394, TE@rnd-4_family-599@DNA@2290:1-394

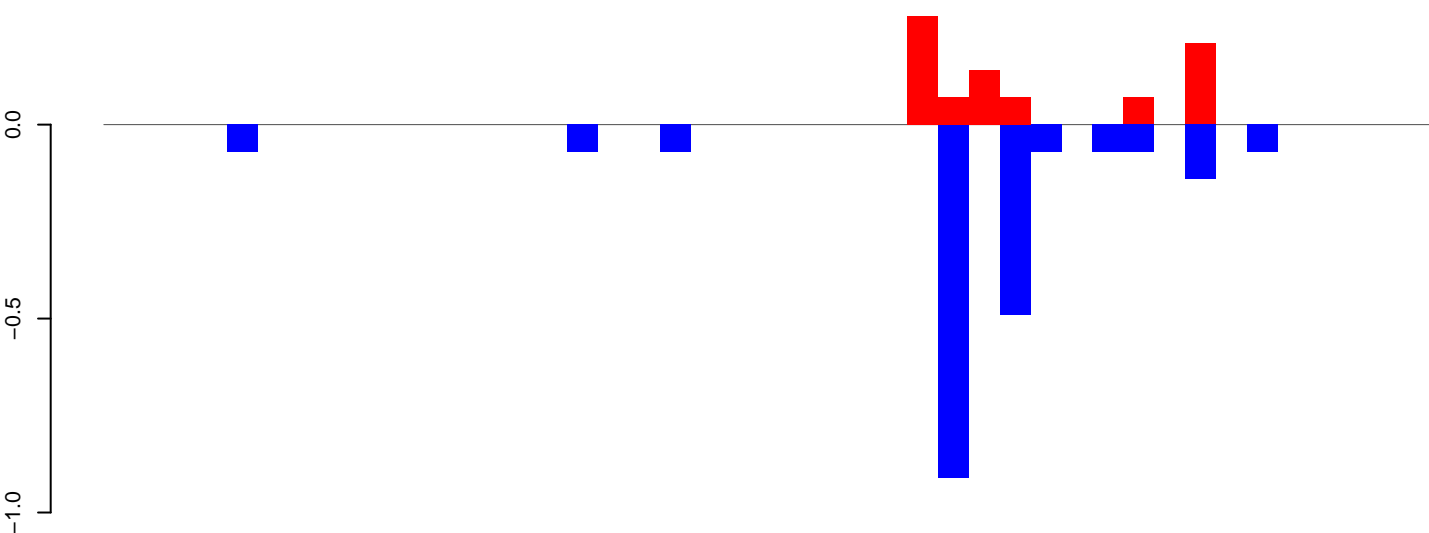
100

200

300

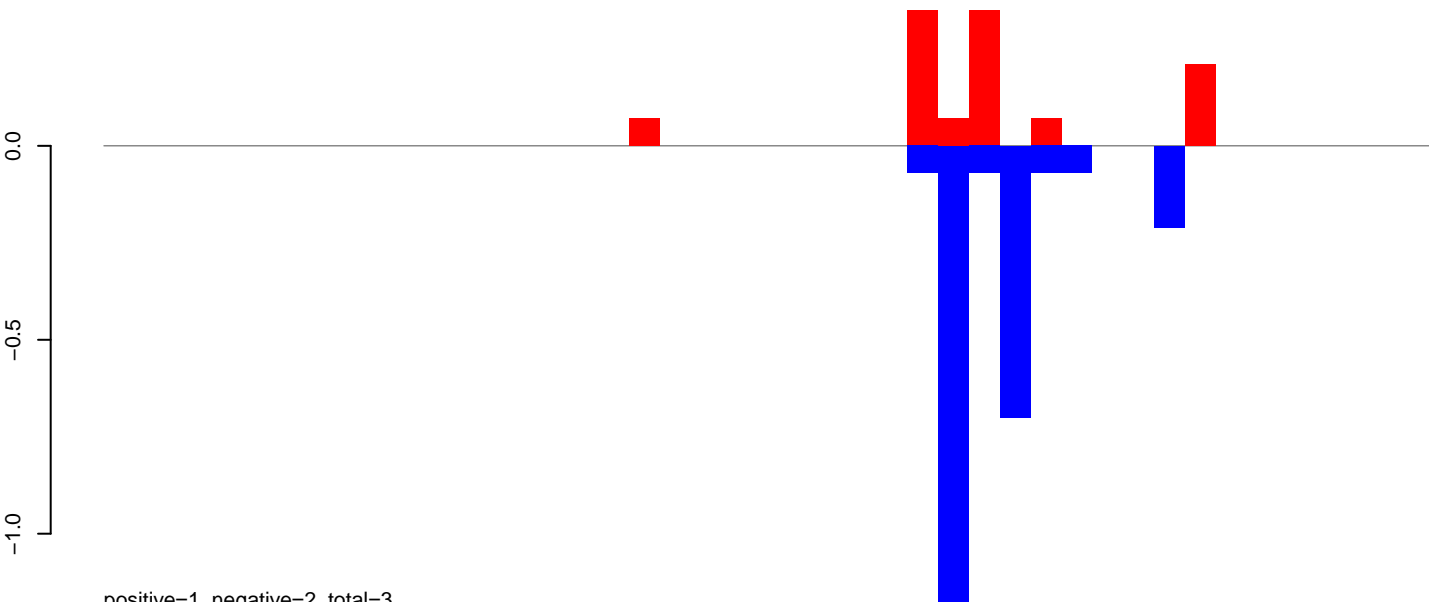
400

CuQuin_Mcarc_NL.18_23.rep



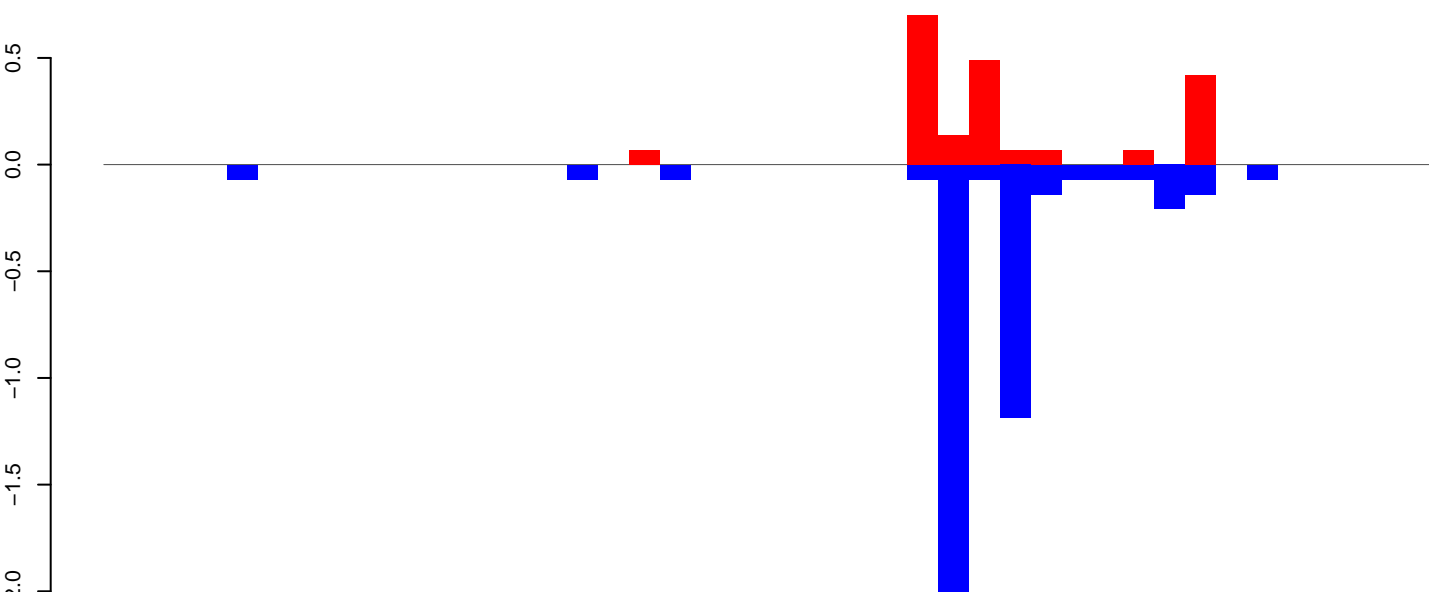
positive=1, negative=2, total=3

CuQuin_Mcarc_NL.24_35.rep



positive=1, negative=2, total=3

CuQuin_Mcarc_NL.rep

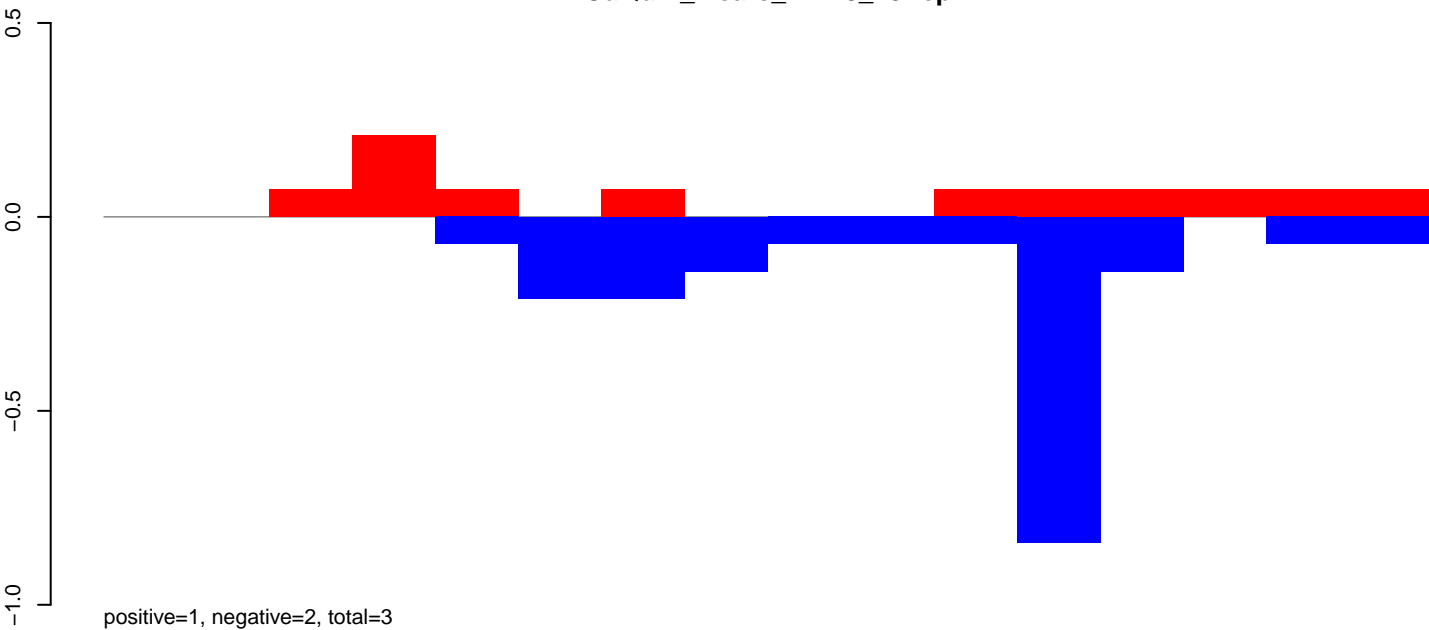


positive=2, negative=4, total=6

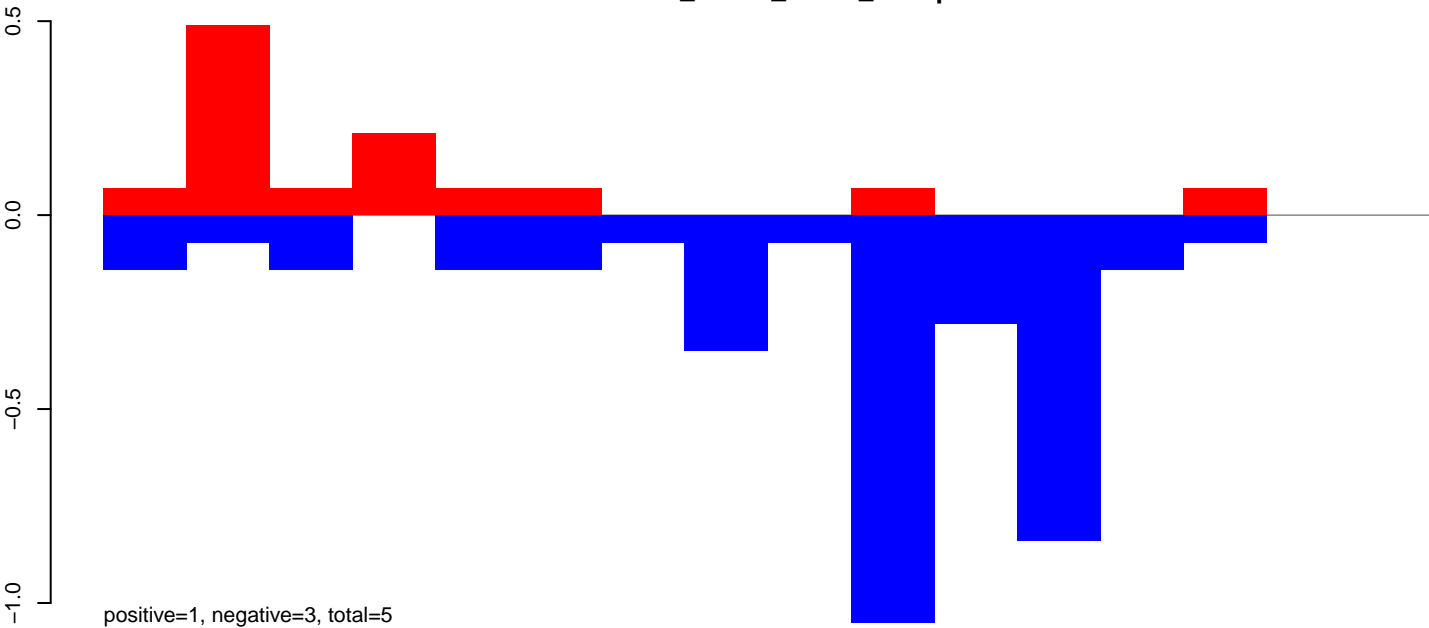
Window size=25, length=1062, TF @ rnd-4_family-2287@SINE_tRNA@1454:1-1062

0 200 400 600 800 1000

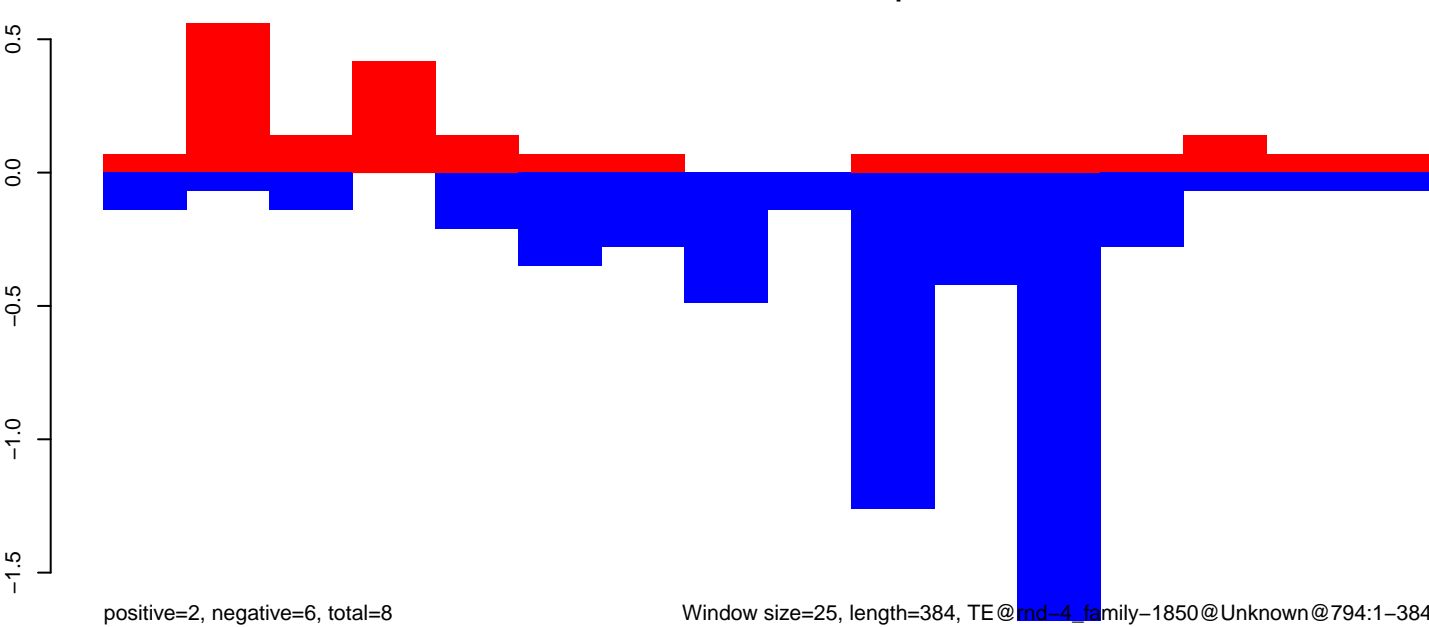
CuQuin_Mcarc_NL.18_23.rep



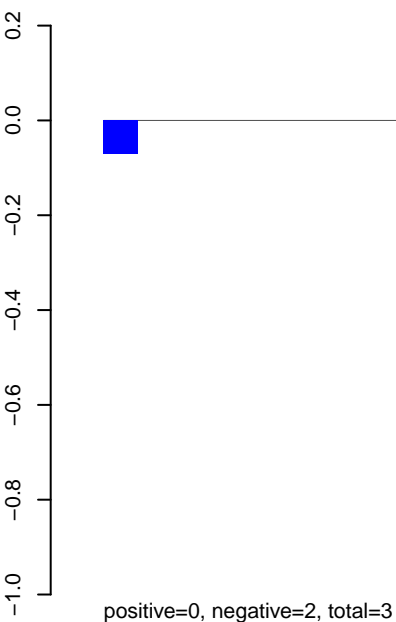
CuQuin_Mcarc_NL.24_35.rep



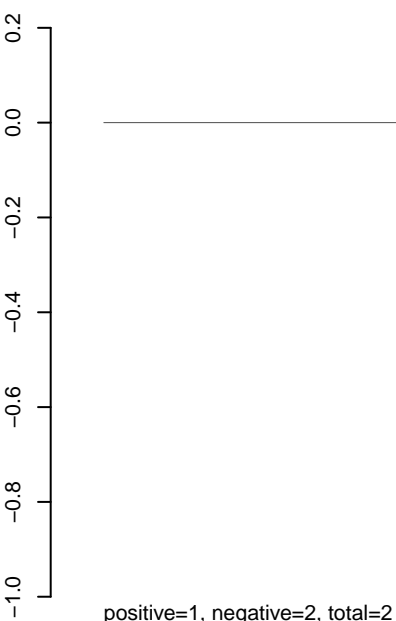
CuQuin_Mcarc_NL.rep



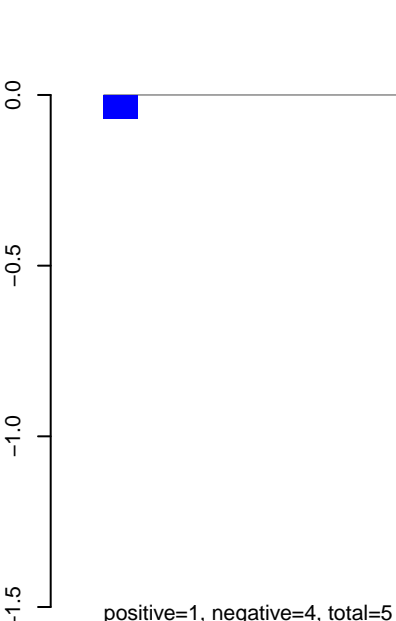
CuQuin_Mcarc_NL.18_23.rep



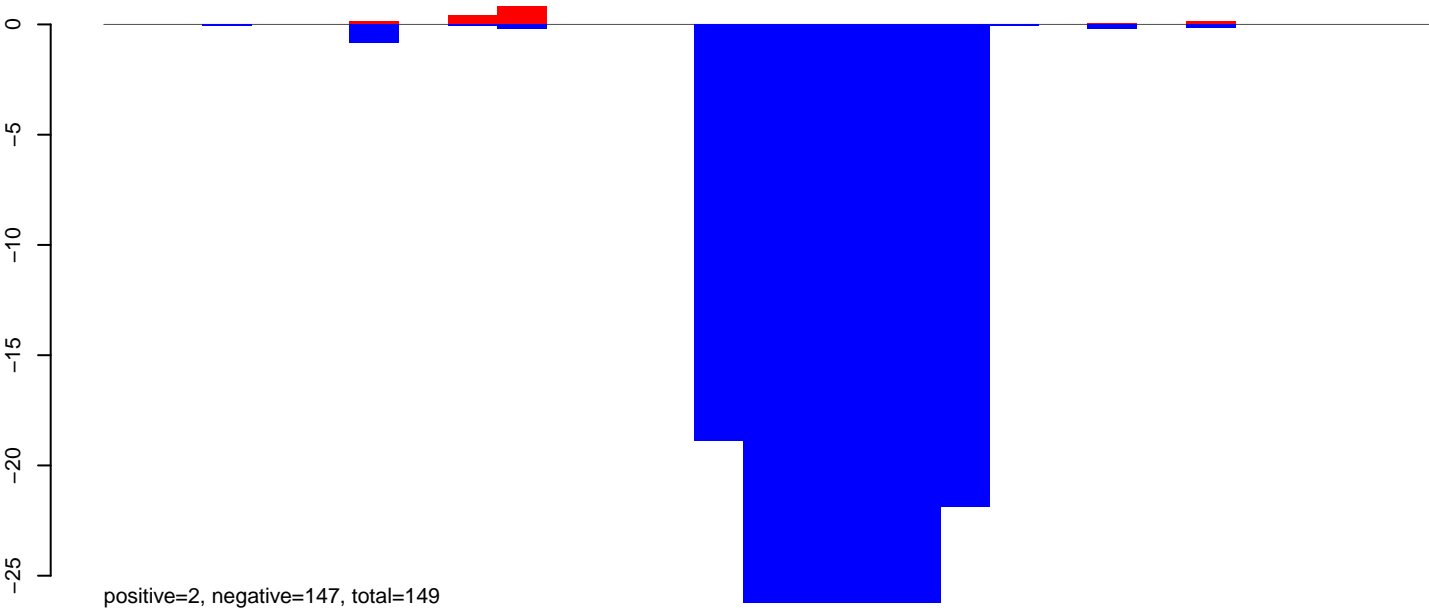
CuQuin_Mcarc_NL.24_35.rep



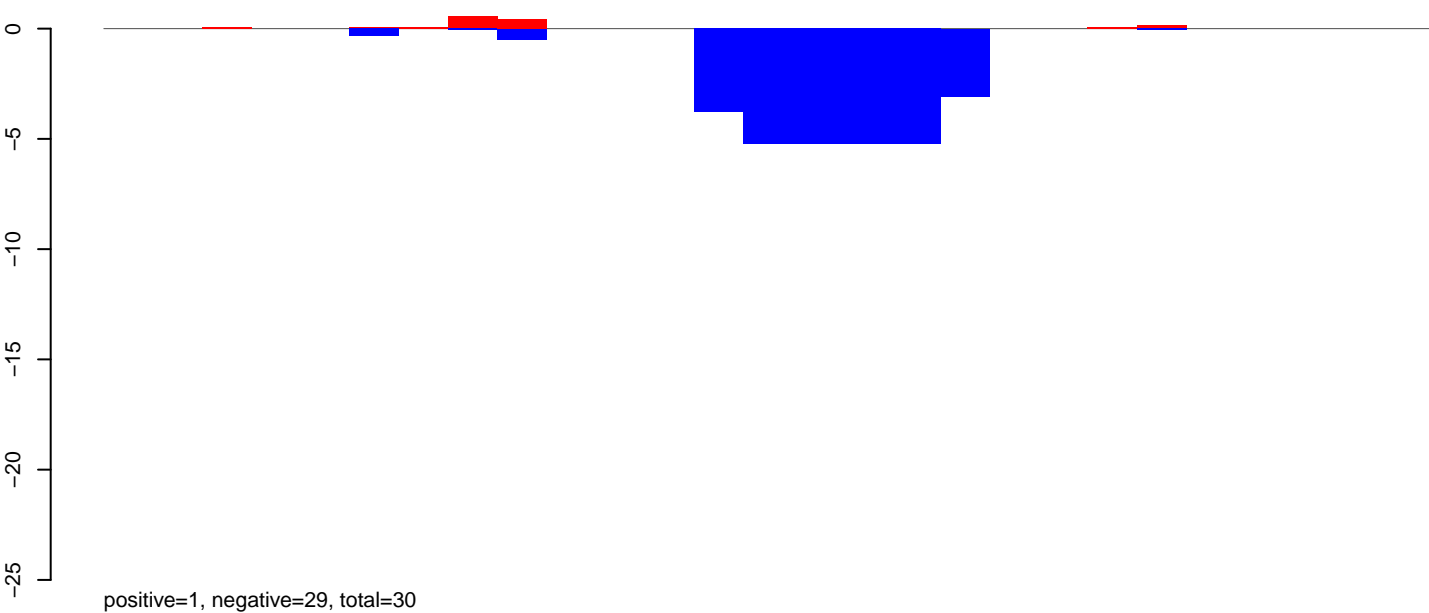
CuQuin_Mcarc_NL.rep



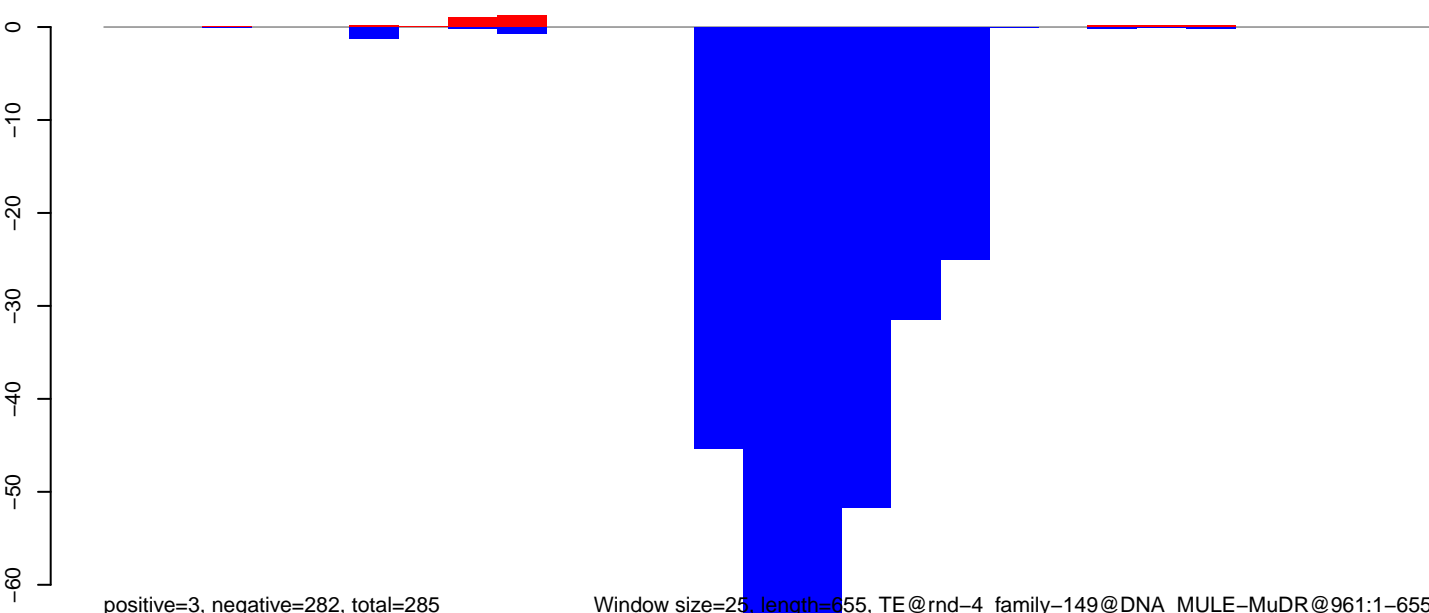
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

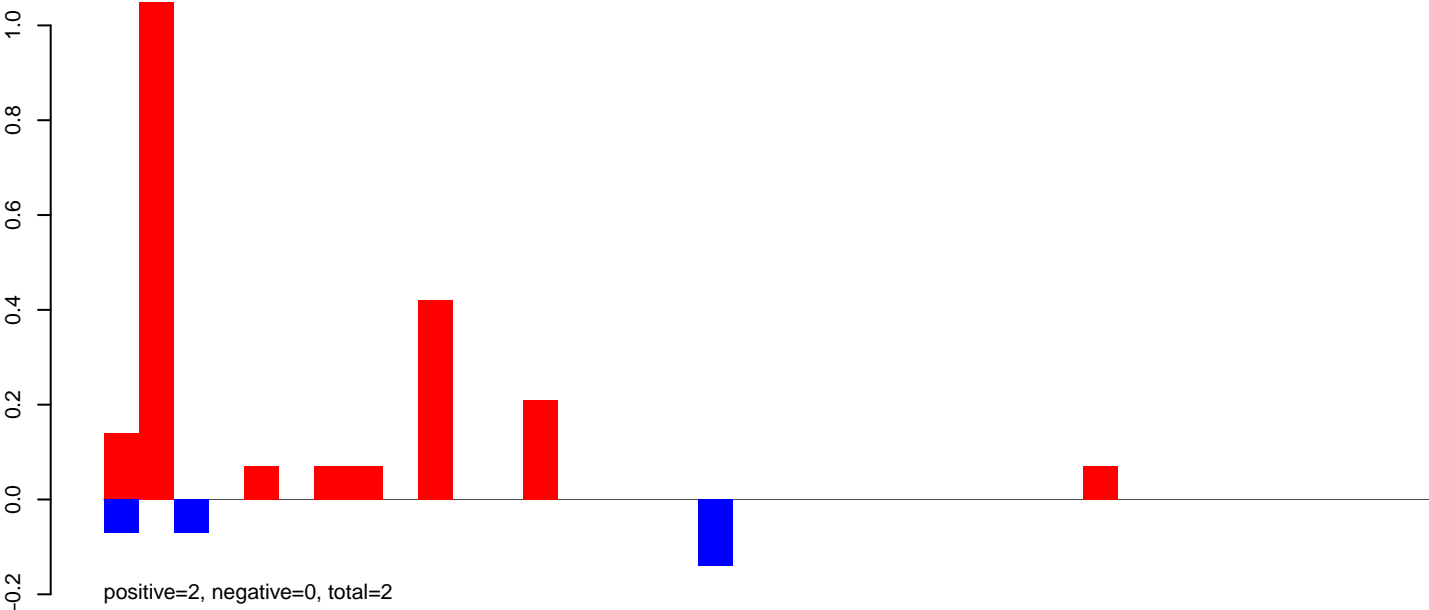


CuQuin_Mcarc_NL.rep

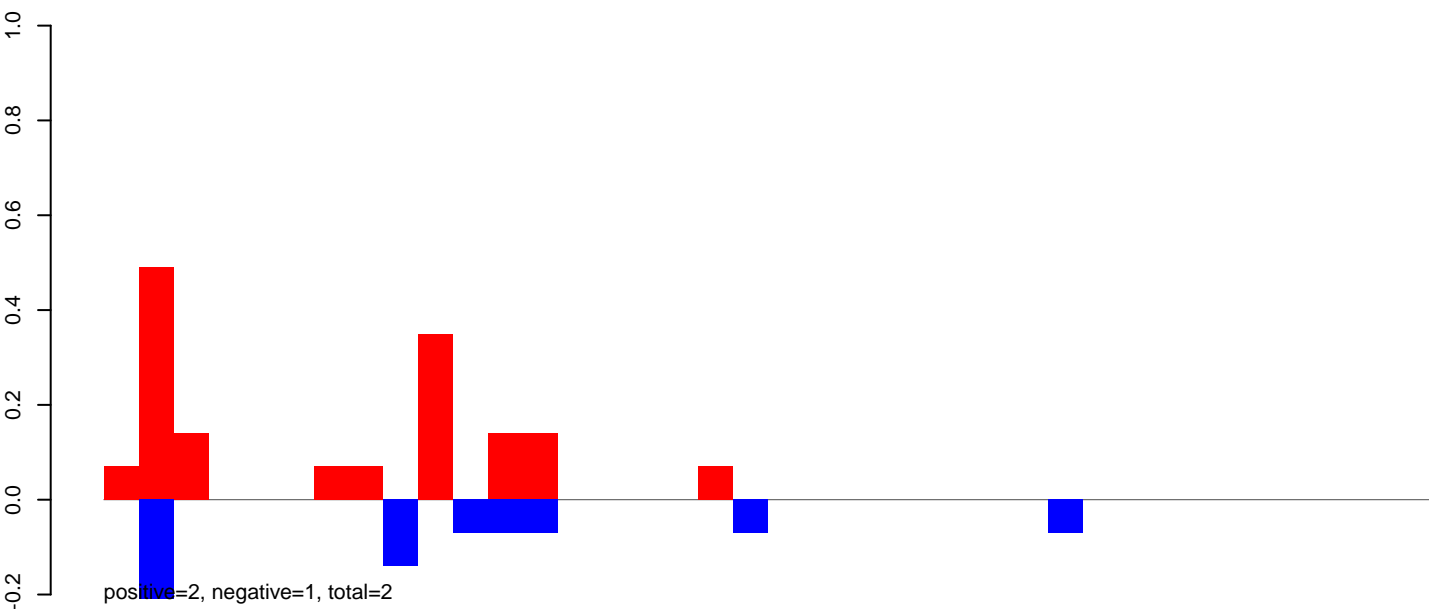


Window size=25, length=655, TE@rnd-4_family-149@DNA_MULE-MuDR@961:1-655

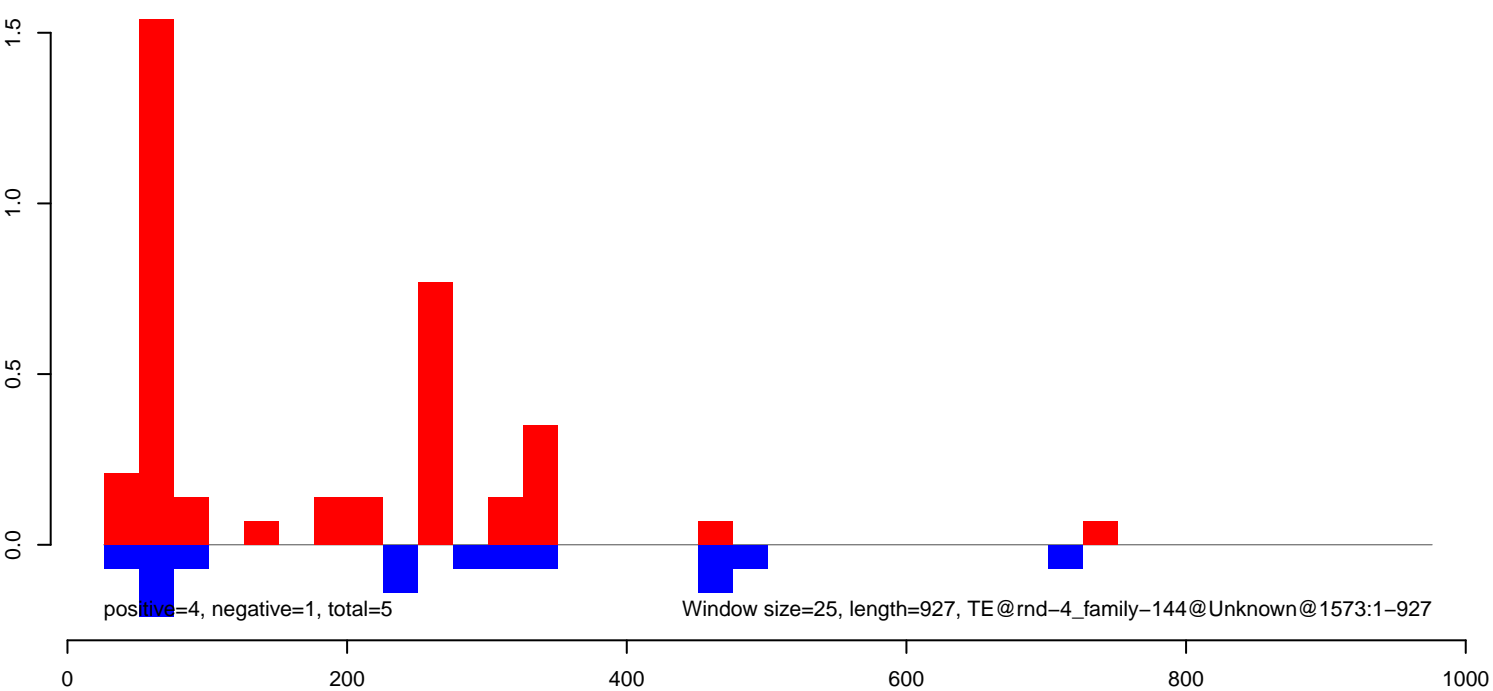
CuQuin_Mcarc_NL.18_23.rep



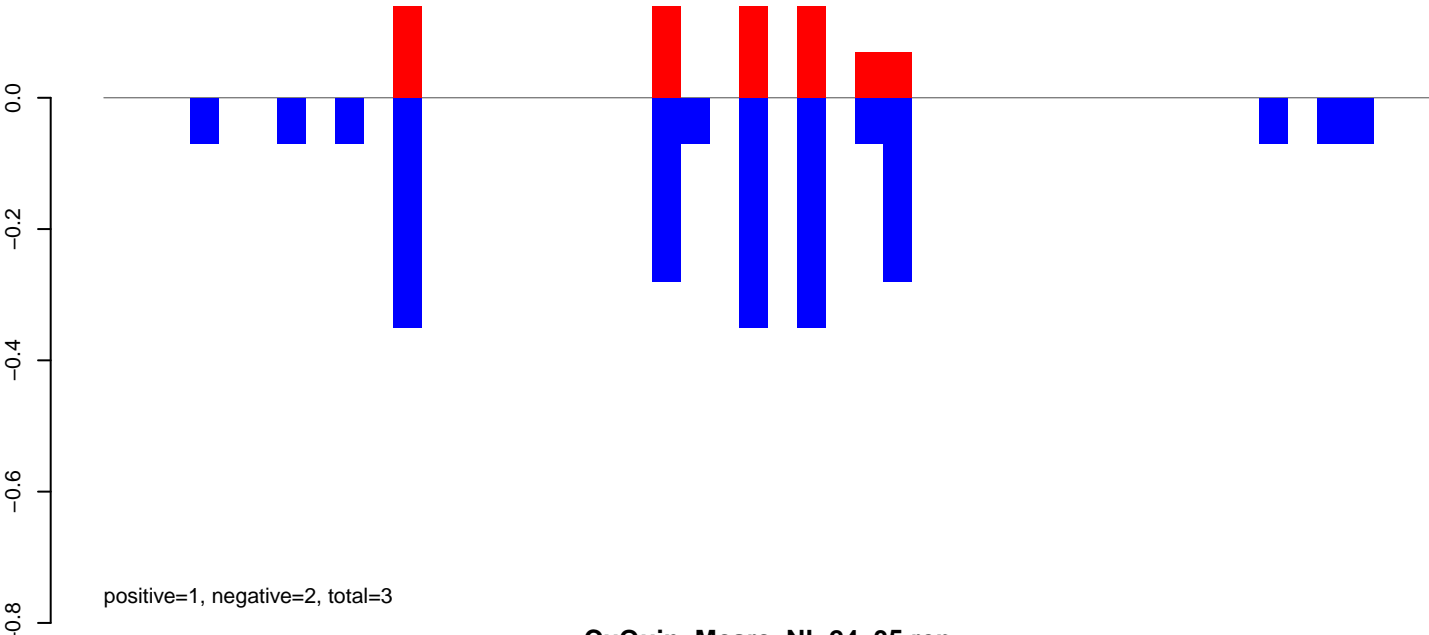
CuQuin_Mcarc_NL.24_35.rep



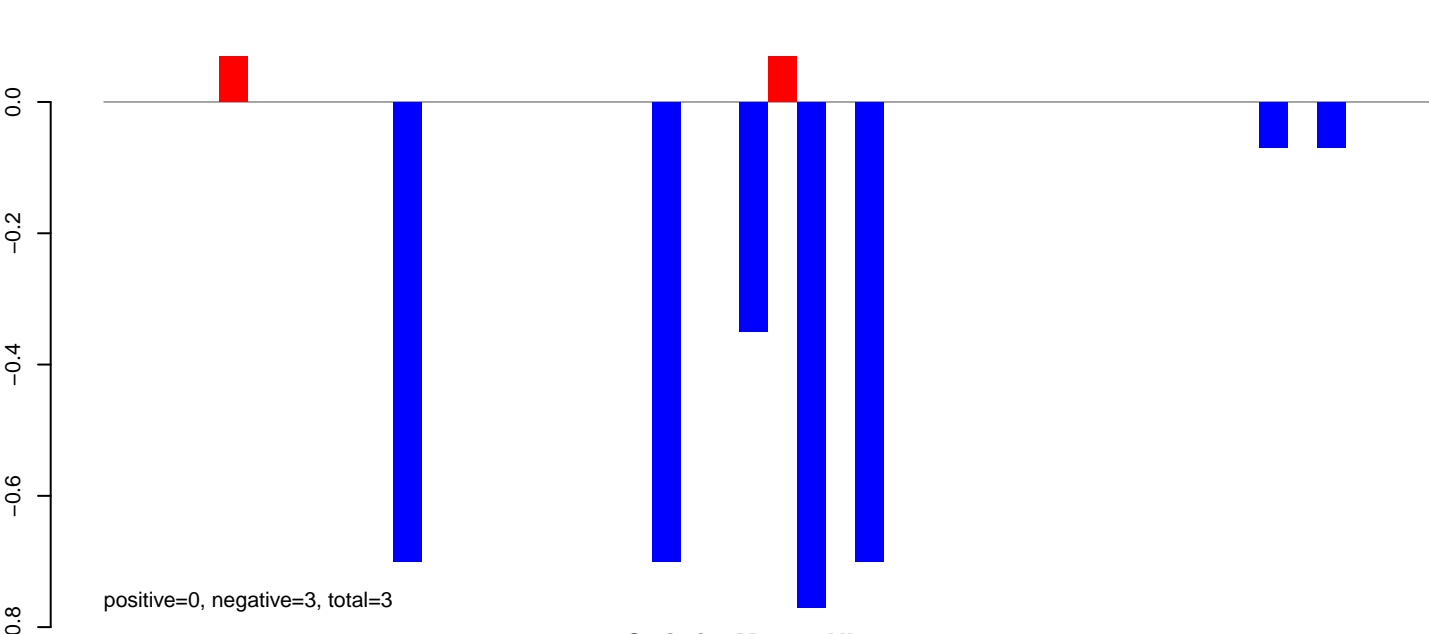
CuQuin_Mcarc_NL.rep



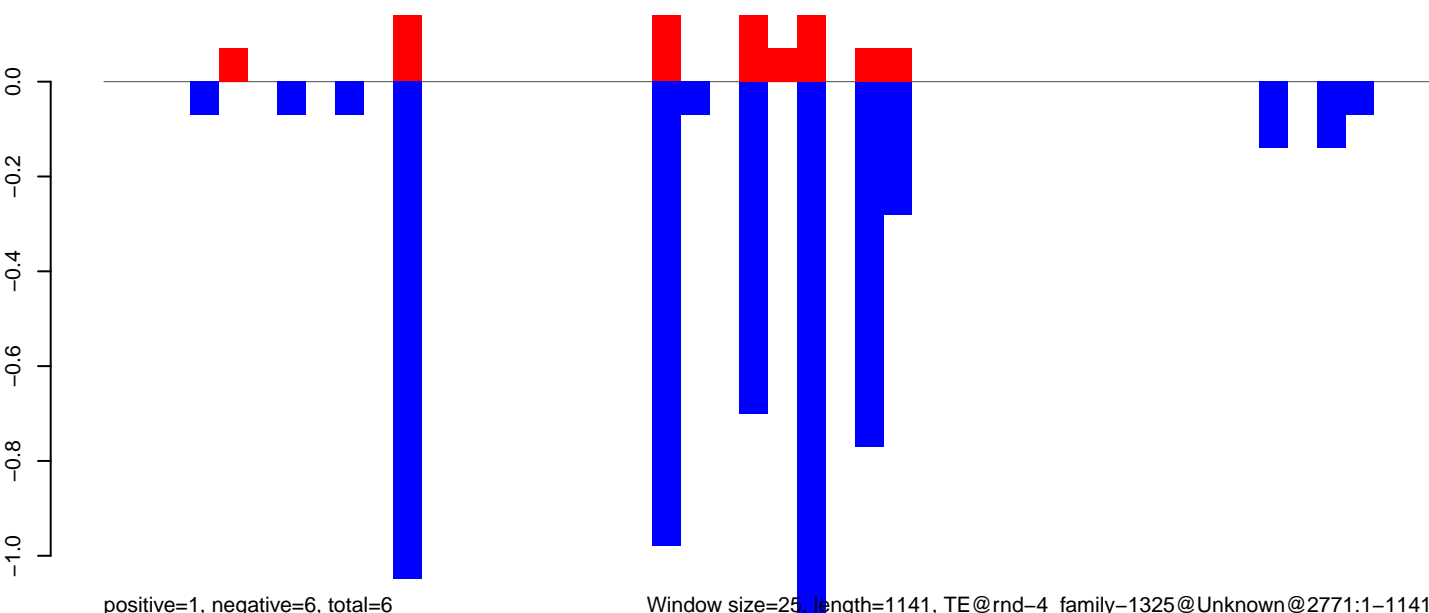
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

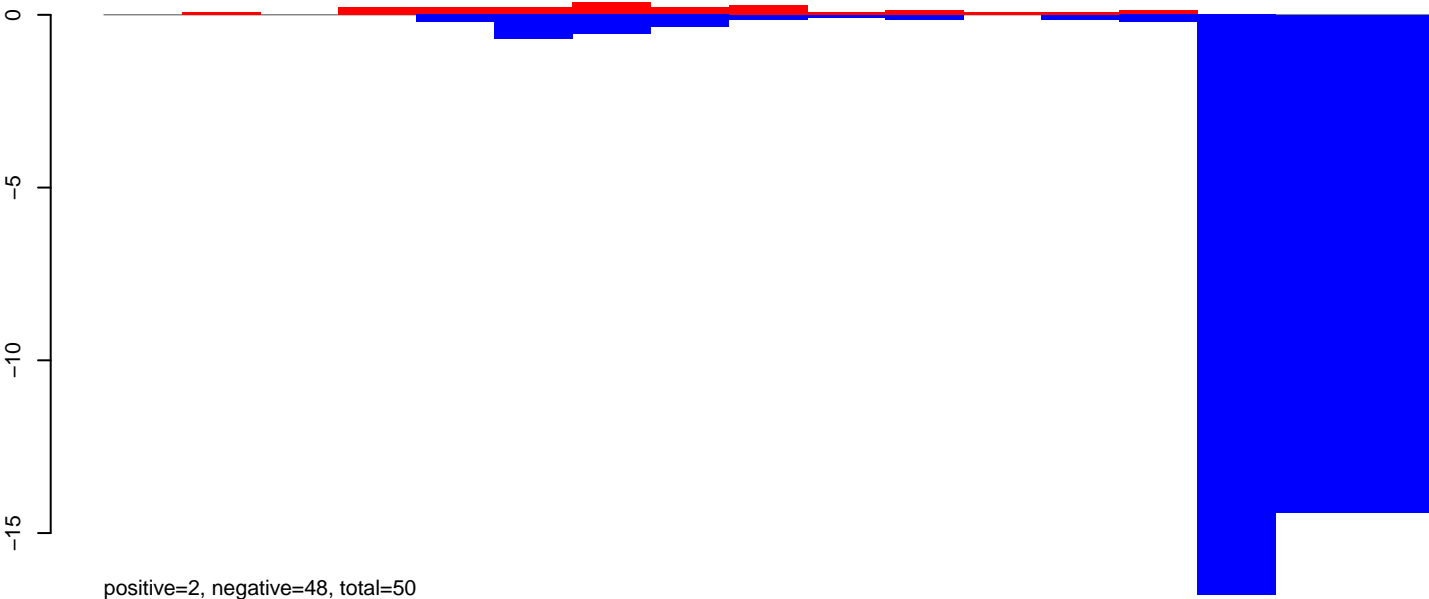


CuQuin_Mcarc_NL.rep

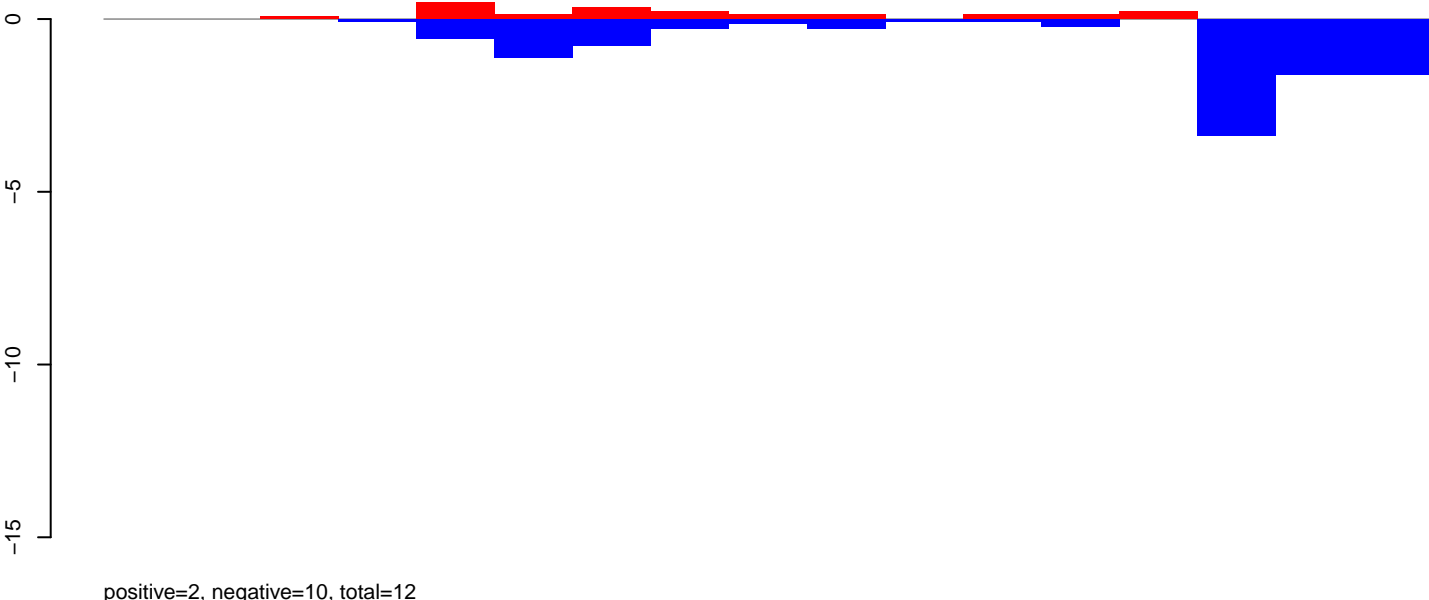


Window size=25, length=1141, TE@rnd-4_family-1325@Unknown@2771:1-1141

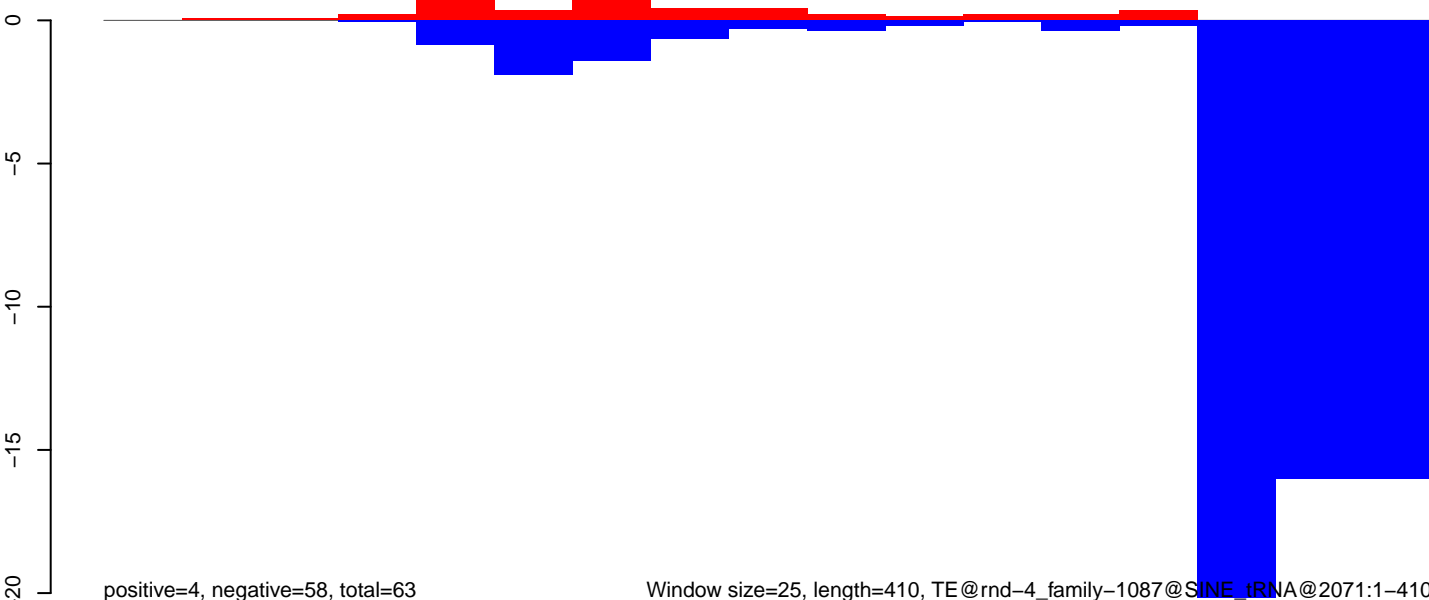
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep



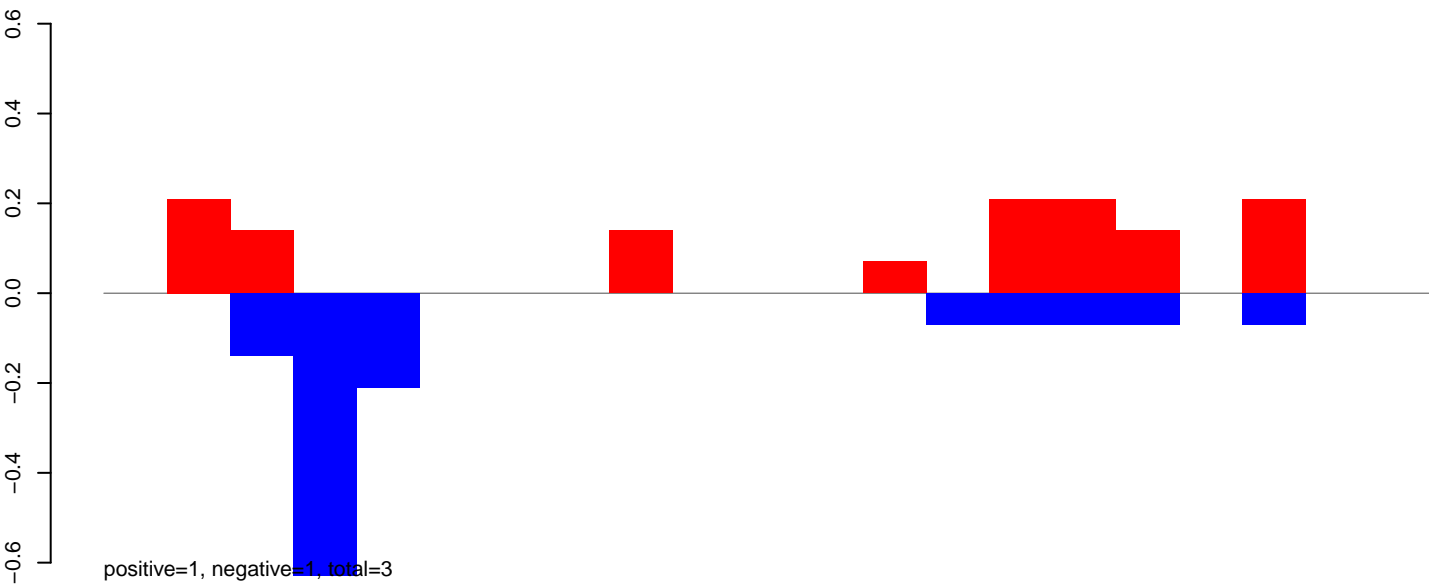
CuQuin_Mcarc_NL.24_35.rep



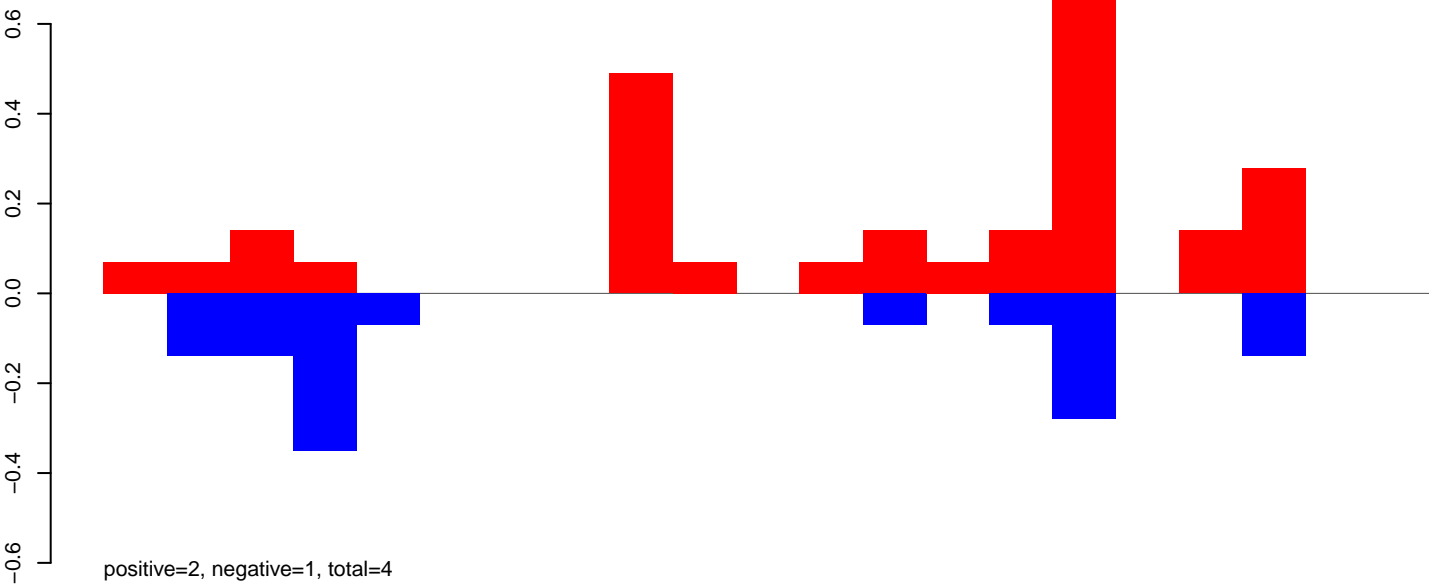
CuQuin_Mcarc_NL.rep



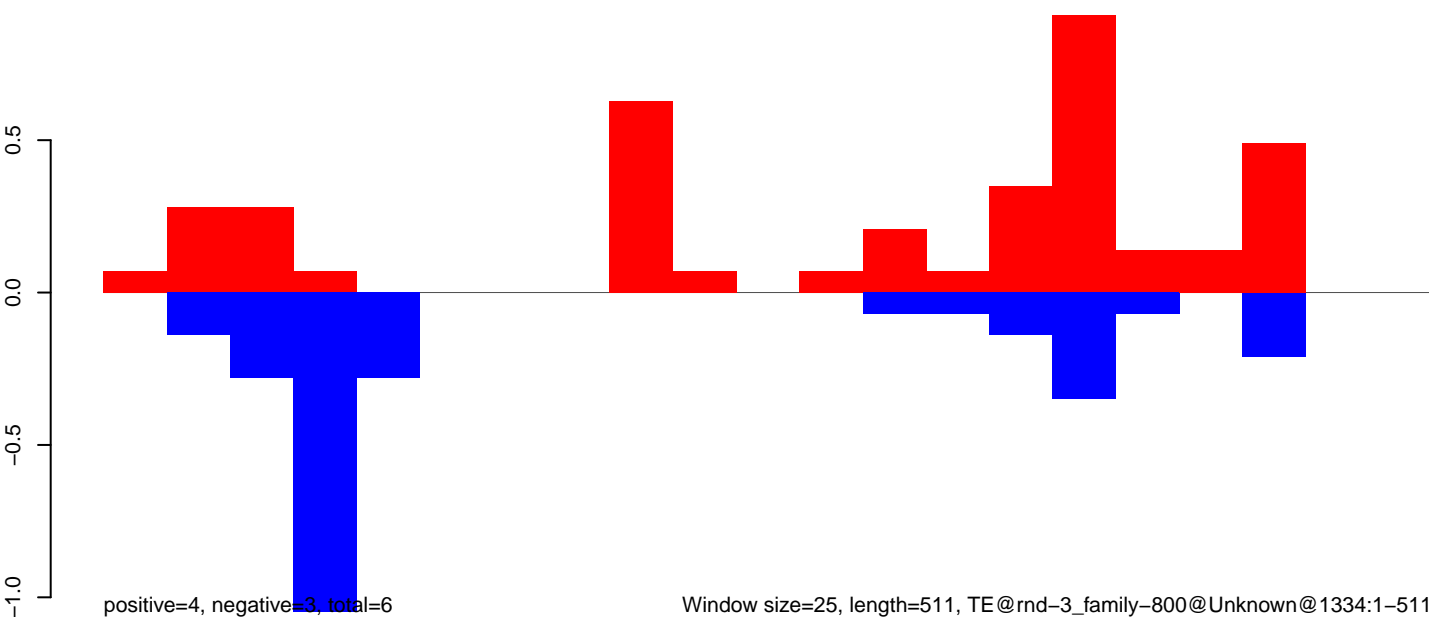
CuQuin_Mcarc_NL.18_23.rep



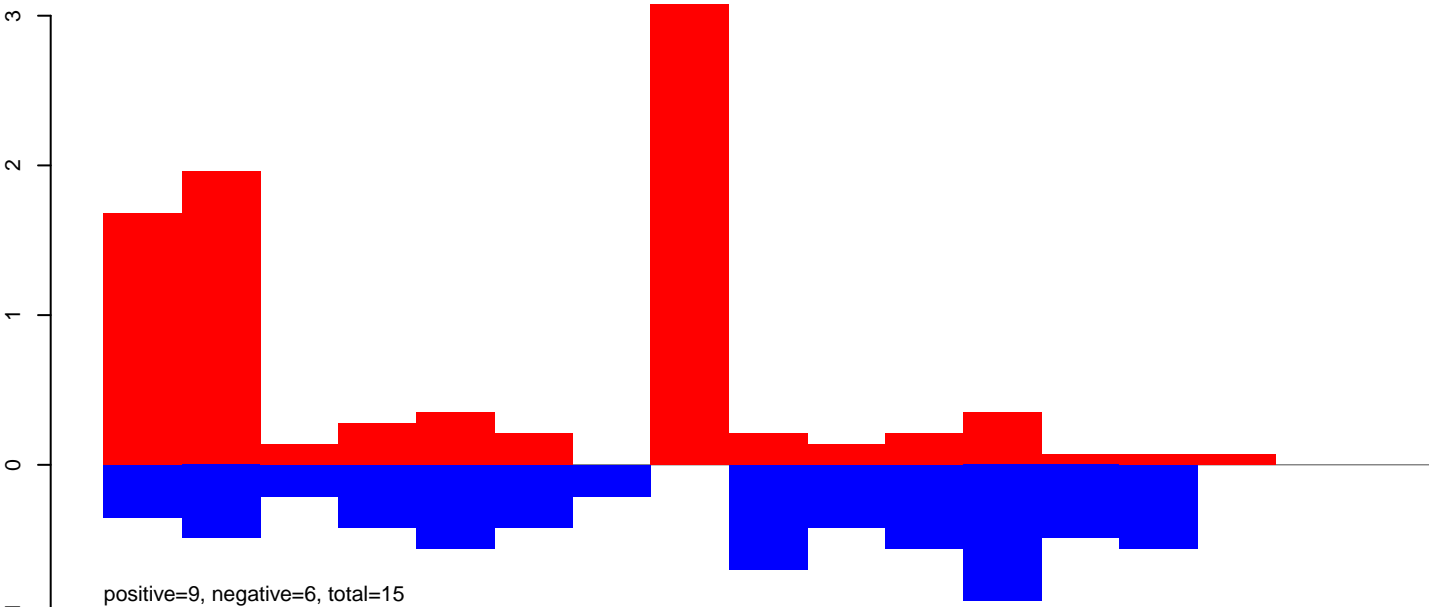
CuQuin_Mcarc_NL.24_35.rep



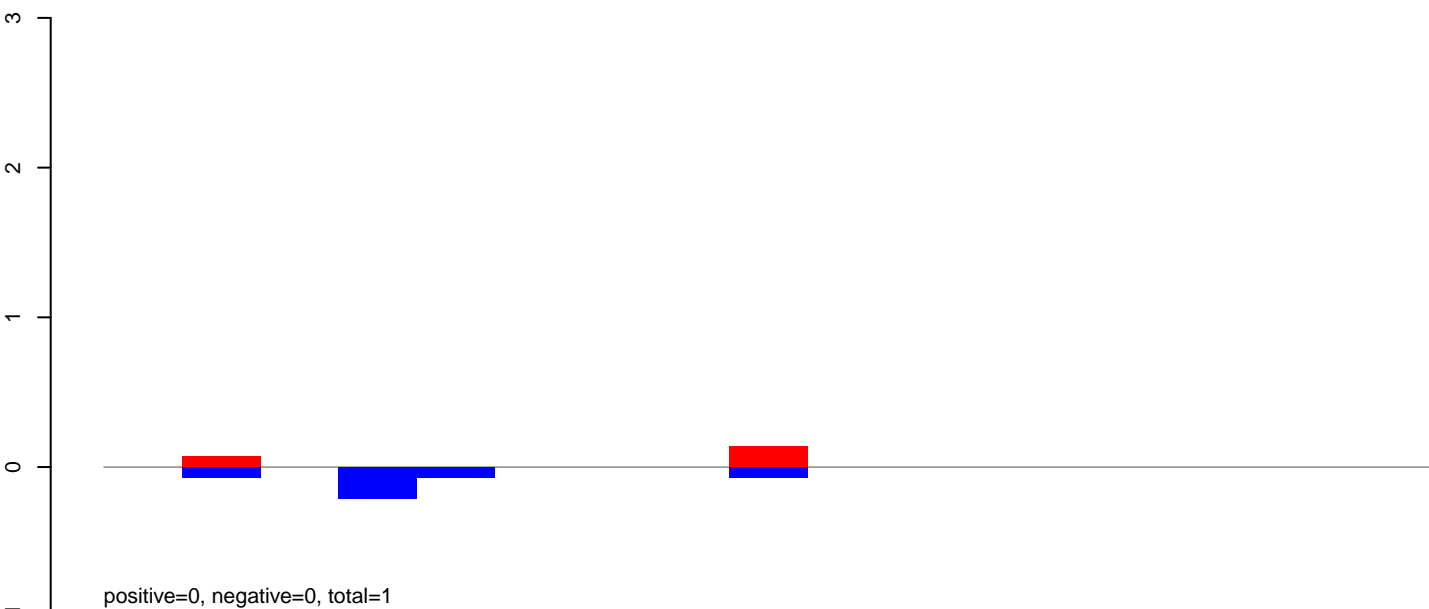
CuQuin_Mcarc_NL.rep



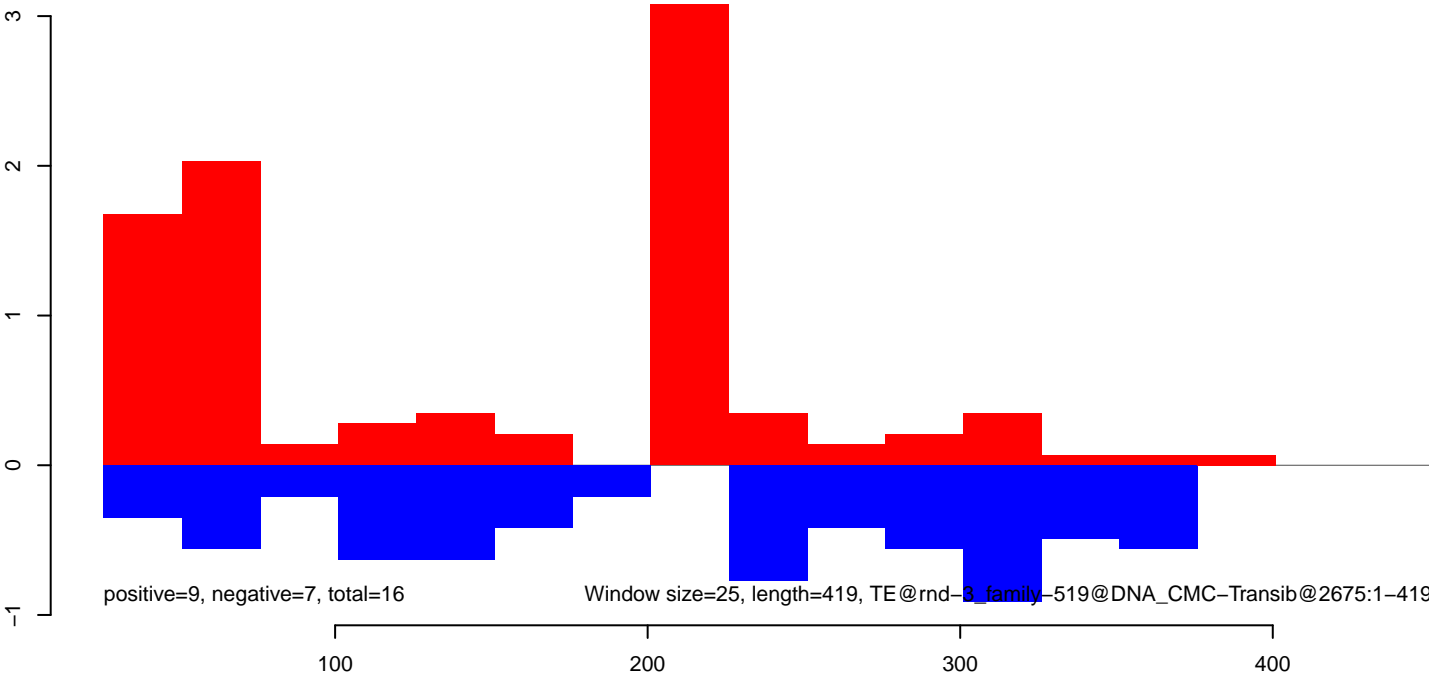
CuQuin_Mcarc_NL.18_23.rep



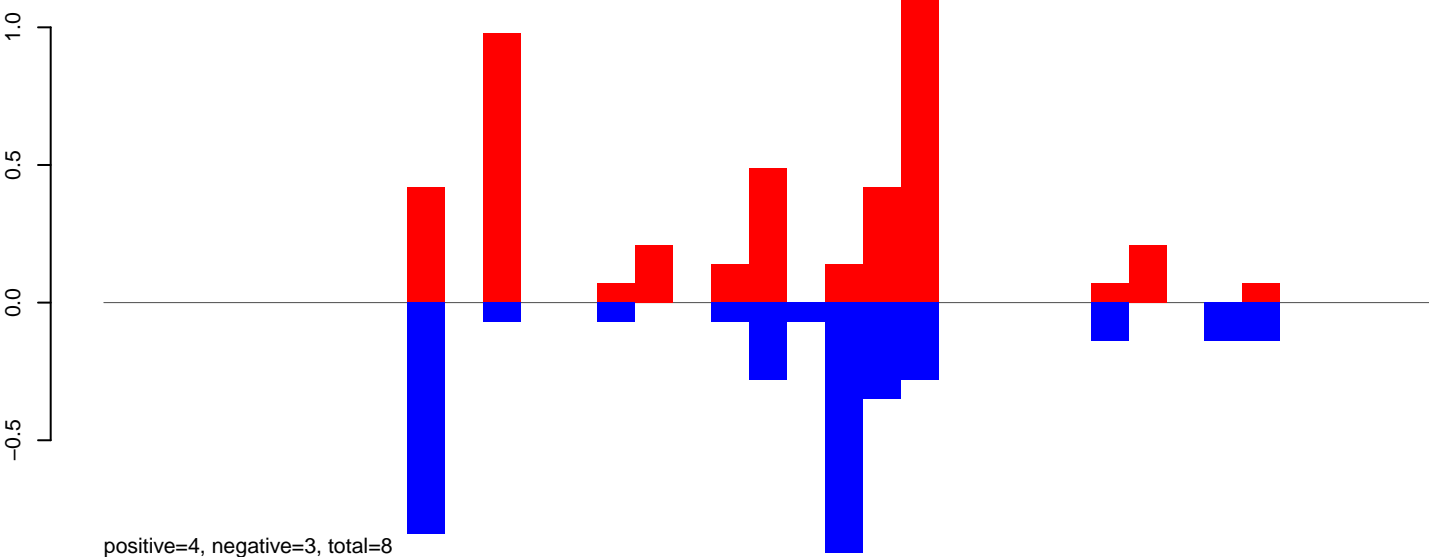
CuQuin_Mcarc_NL.24_35.rep



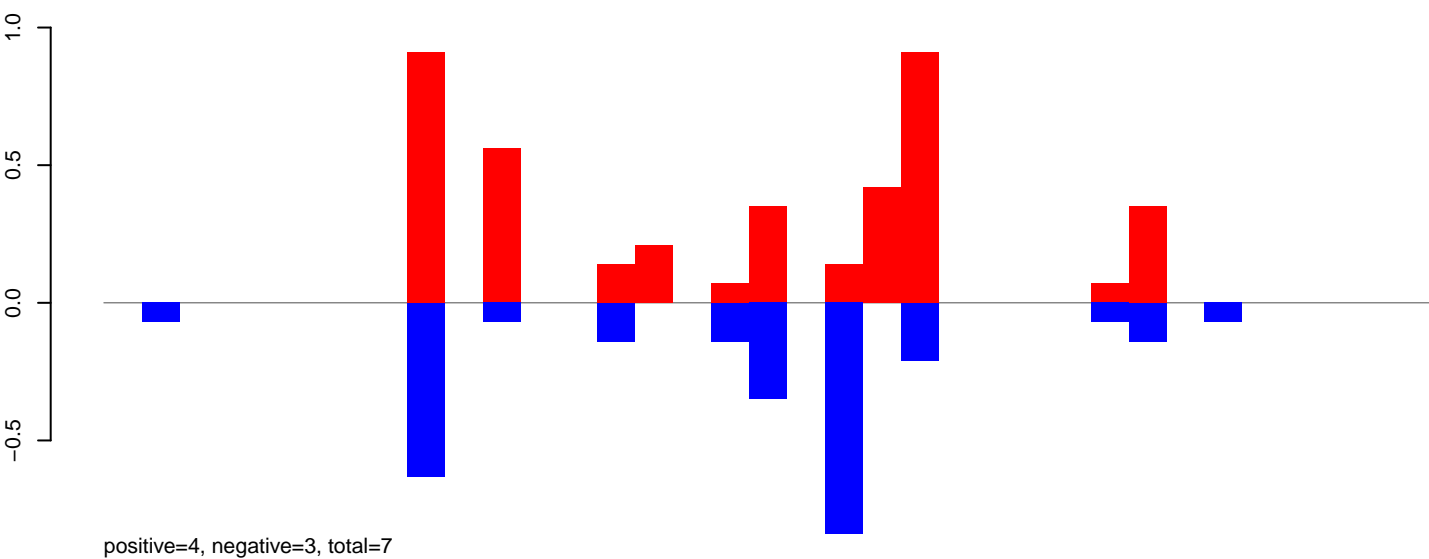
CuQuin_Mcarc_NL.rep



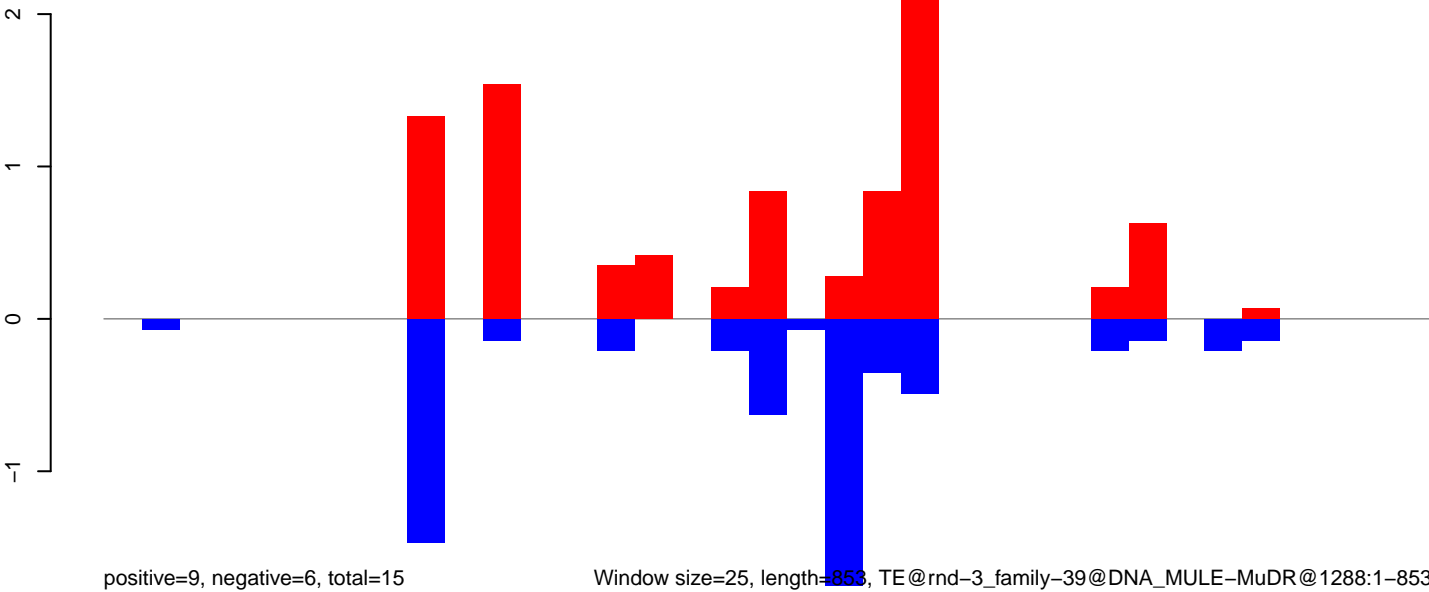
CuQuin_Mcarc_NL.18_23.rep



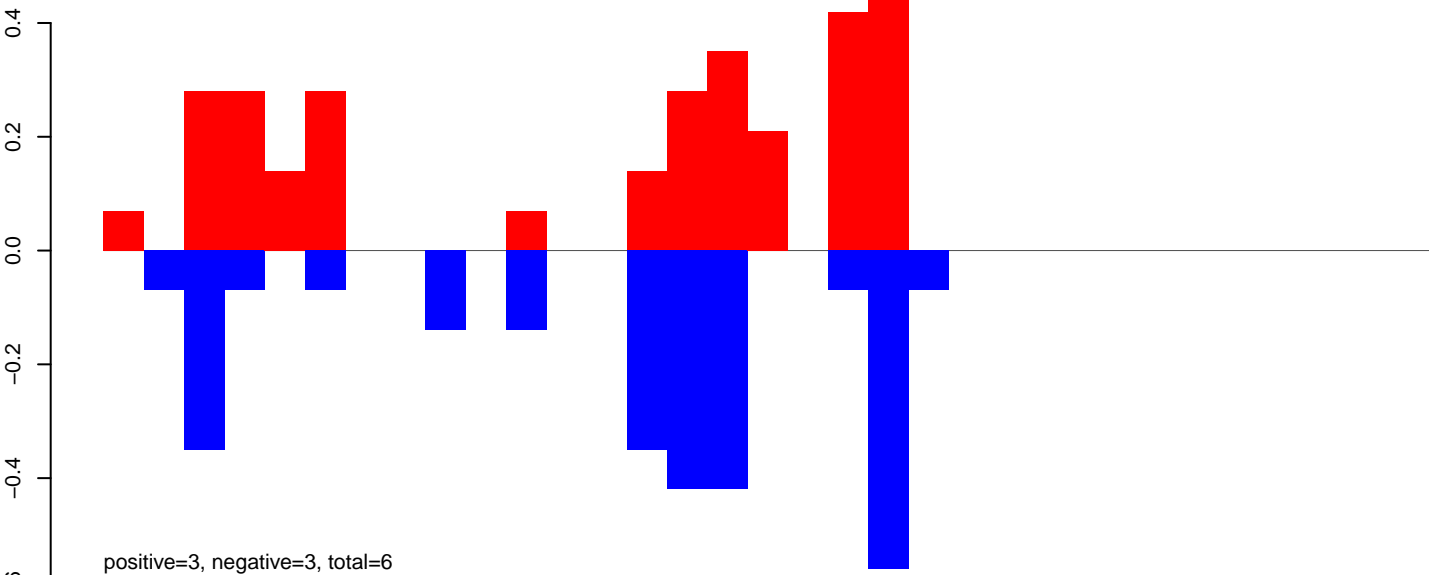
CuQuin_Mcarc_NL.24_35.rep



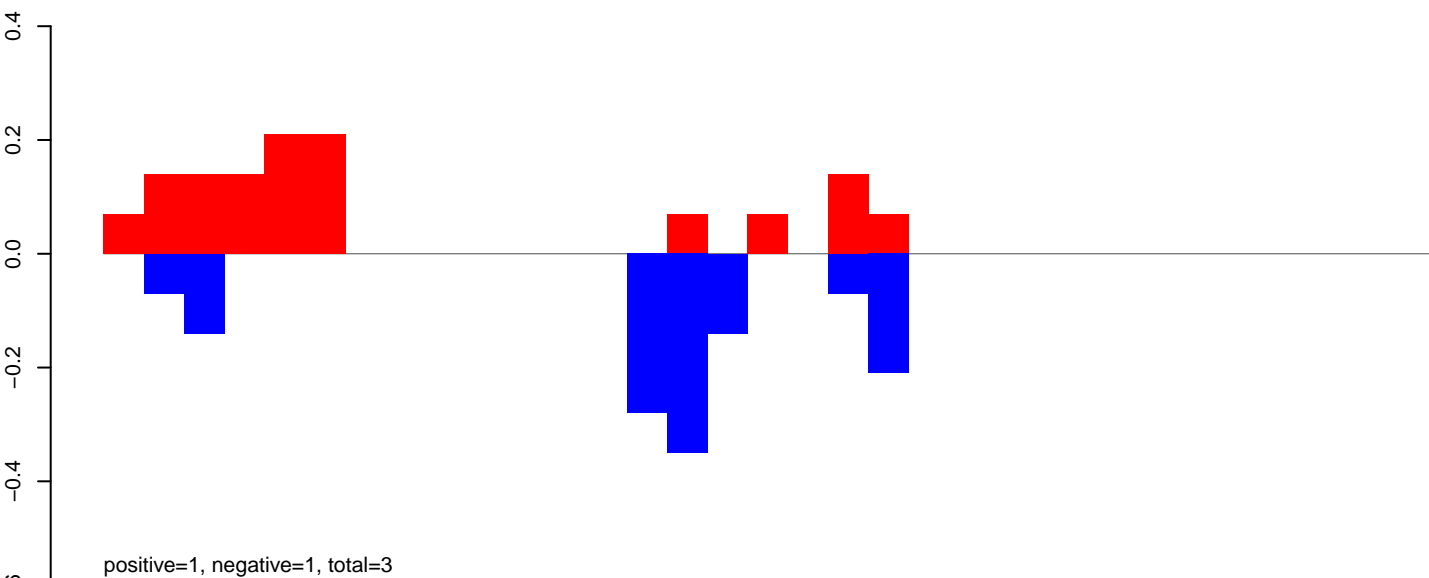
CuQuin_Mcarc_NL.rep



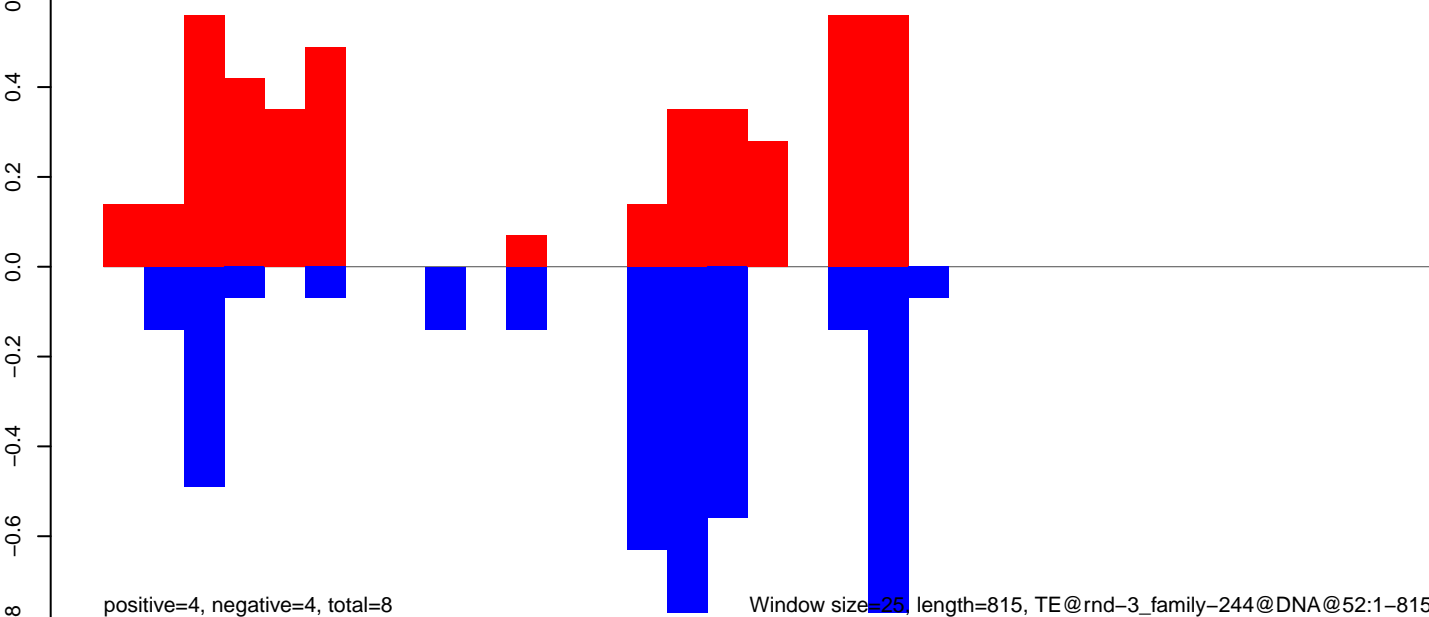
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

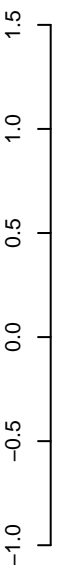


CuQuin_Mcarc_NL.18_23.rep



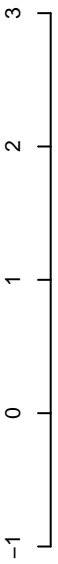
positive=3, negative=3, total=6

CuQuin_Mcarc_NL.24_35.rep



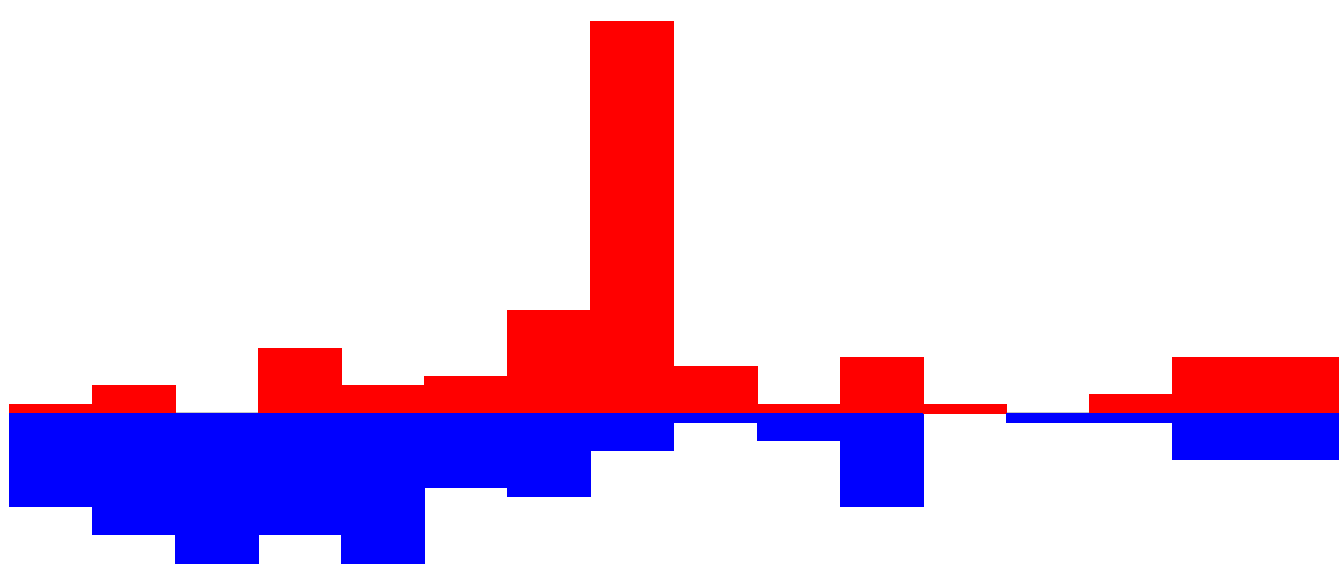
positive=4, negative=5, total=9

CuQuin_Mcarc_NL.rep

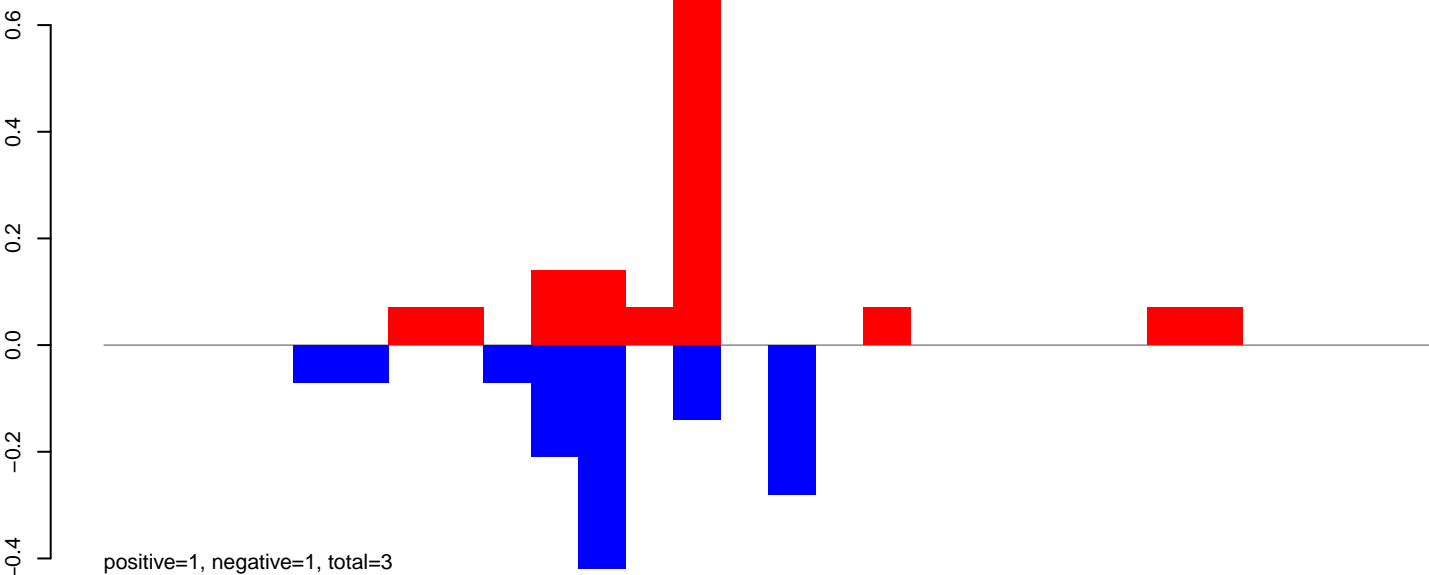


positive=7, negative=9, total=15

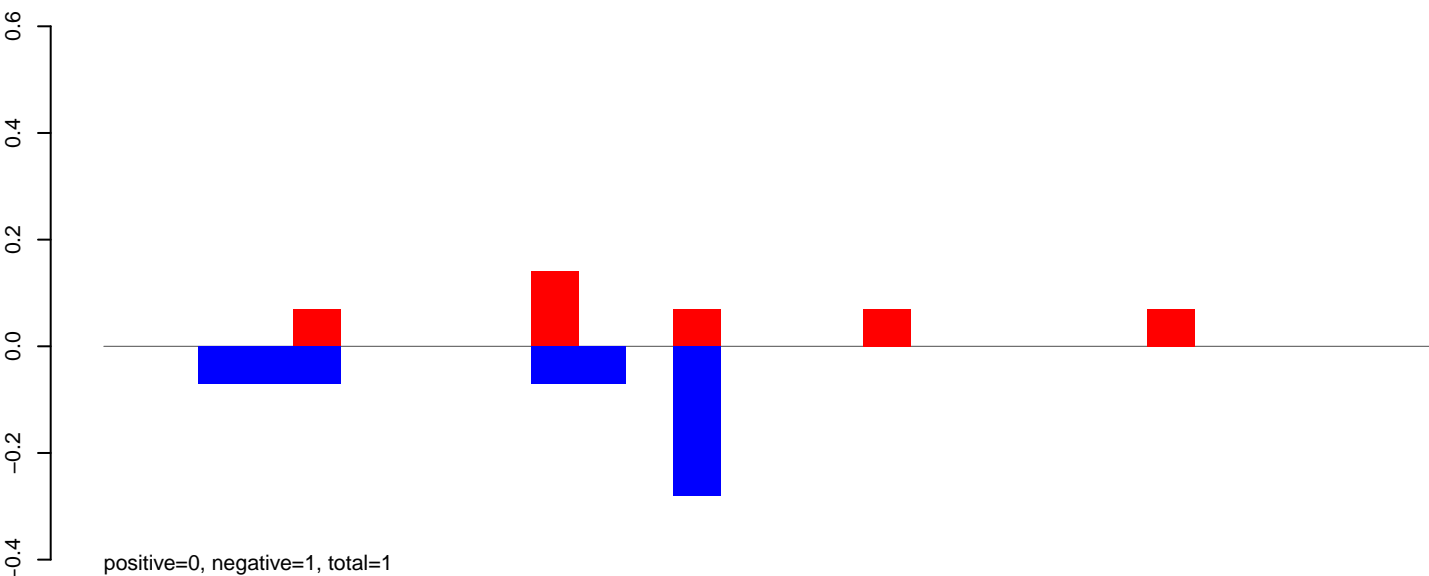
Window size=25, length=400, TE@rnd-2_family-54@Unknown@12:1-400



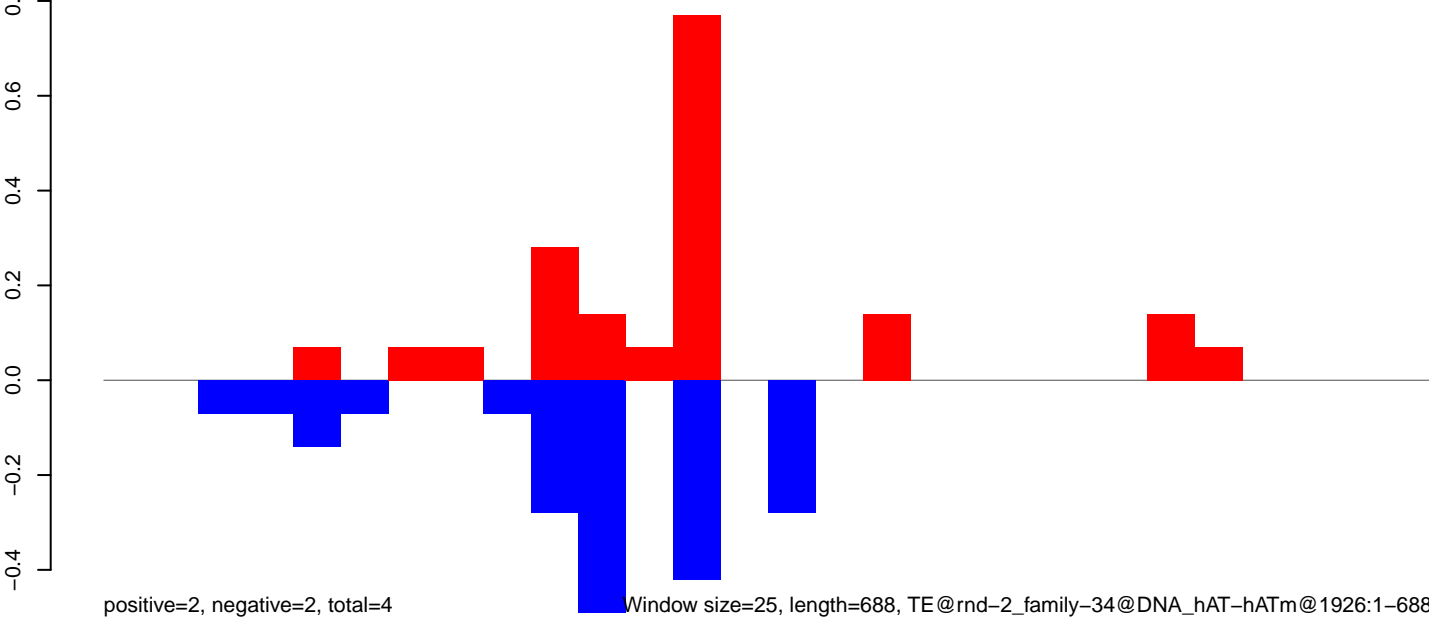
CuQuin_Mcarc_NL.18_23.rep



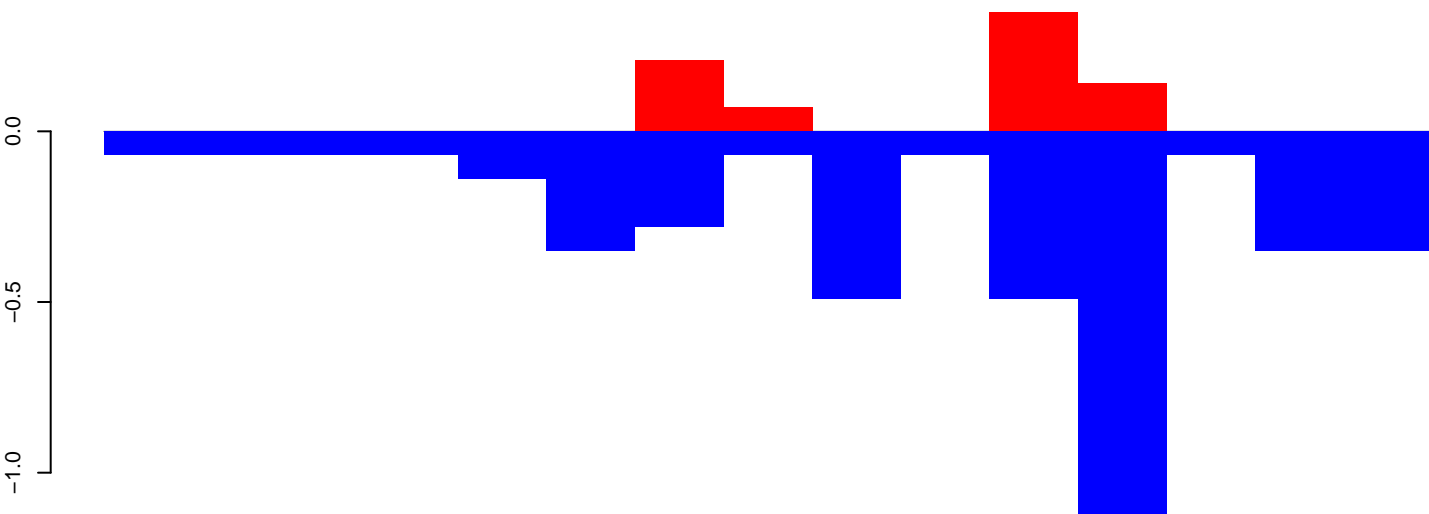
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

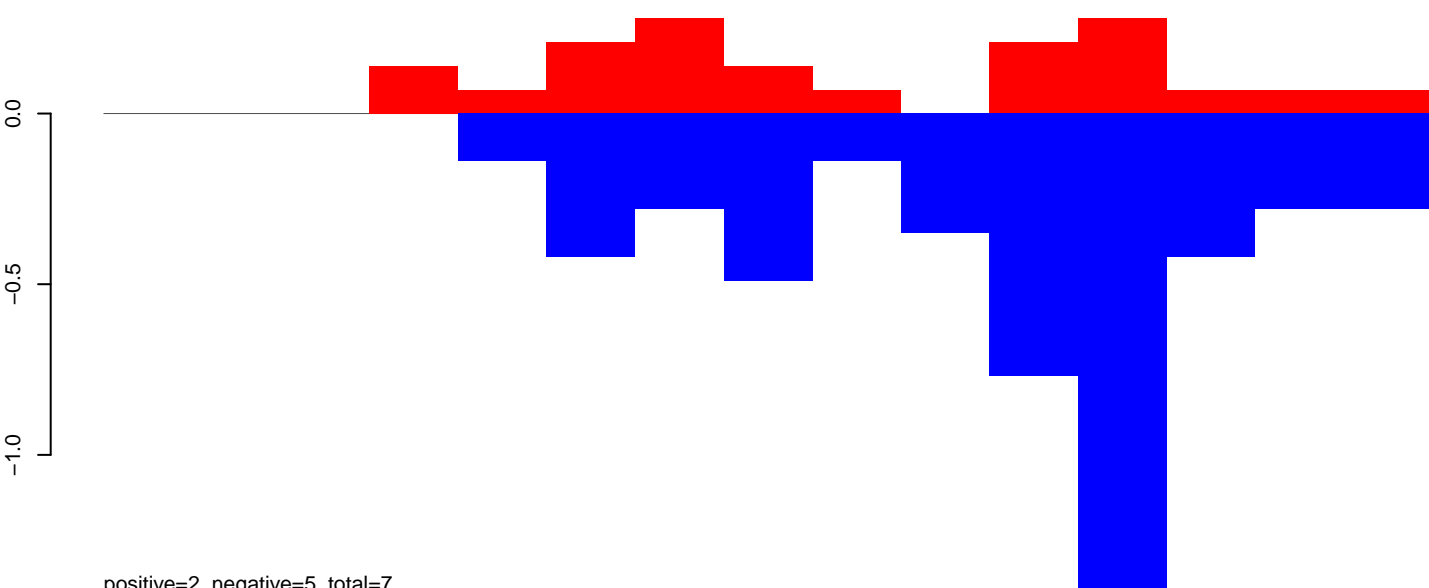


CuQuin_Mcarc_NL.18_23.rep



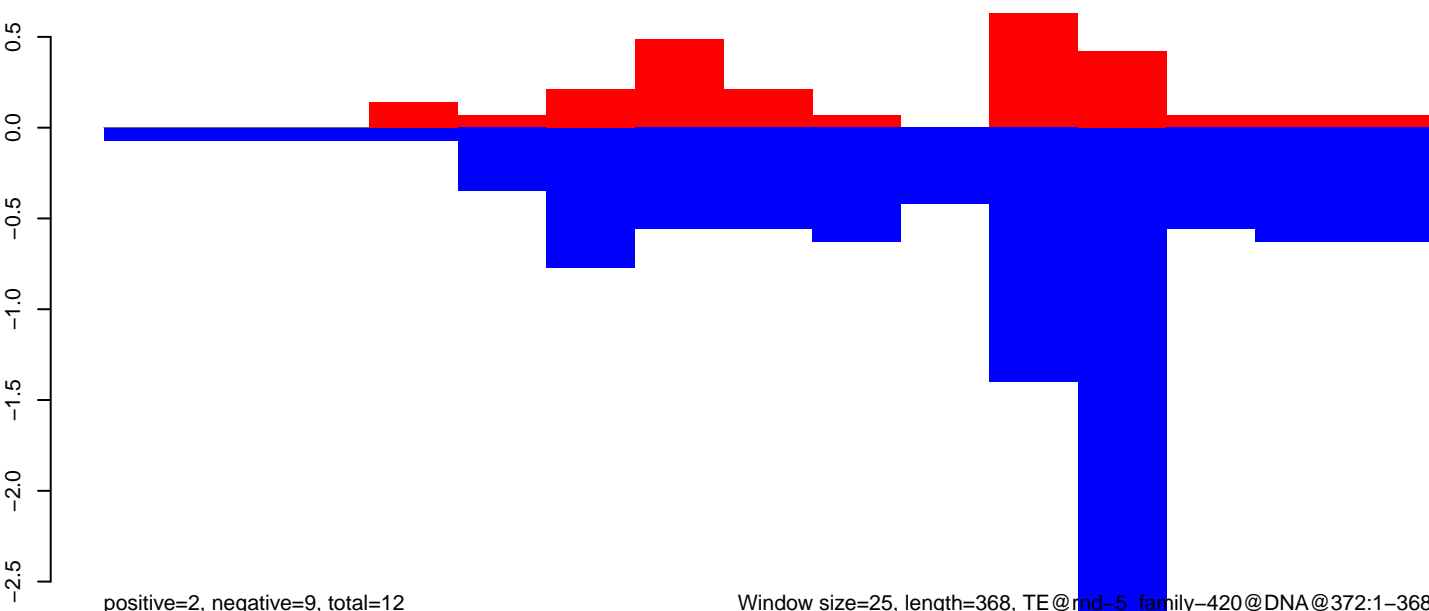
positive=1, negative=4, total=5

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=5, total=7

CuQuin_Mcarc_NL.rep



positive=2, negative=9, total=12

Window size=25, length=368, TE@md-5, family-420@DNA@372:1-368

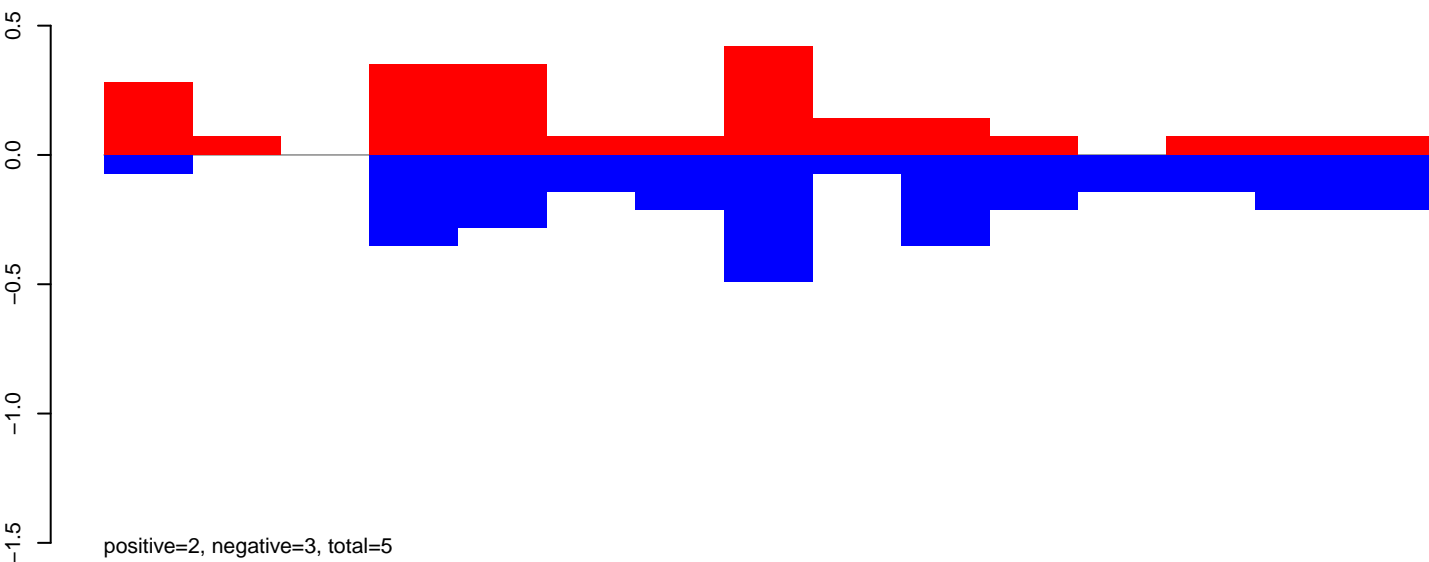
100

200

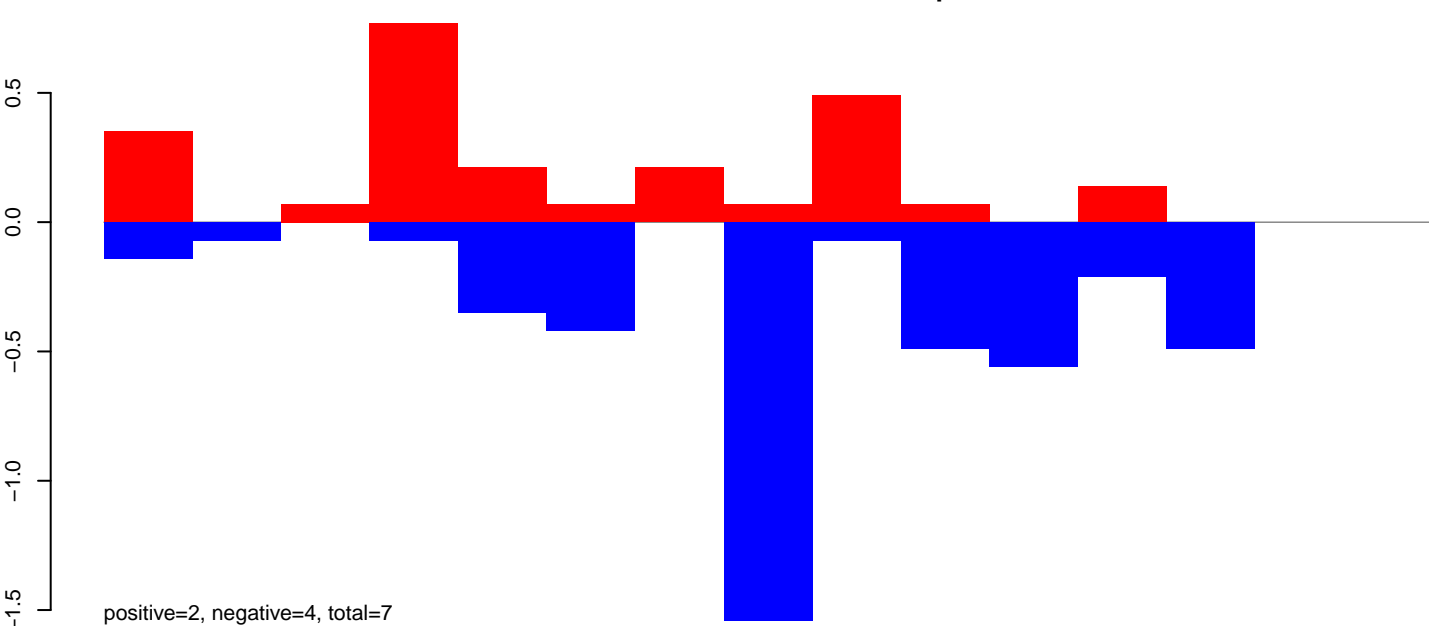
300

400

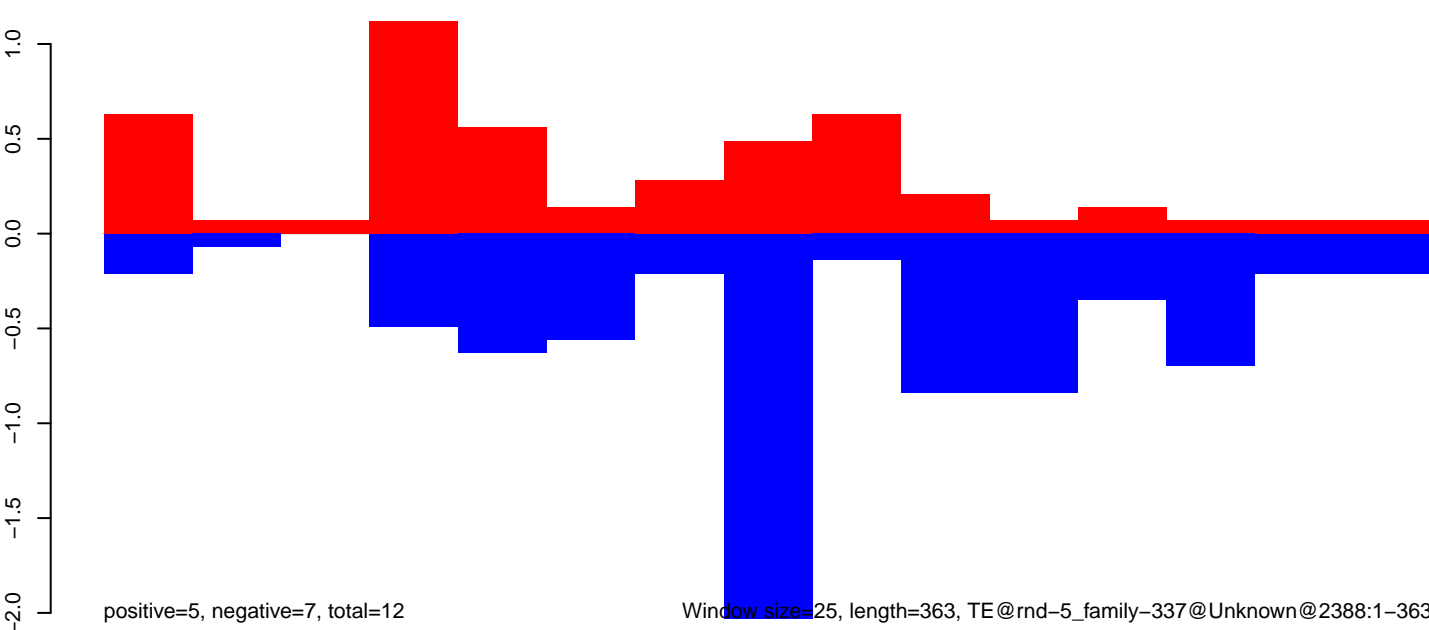
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=363, TE@rnd-5_family-337@Unknown@2388:1-363

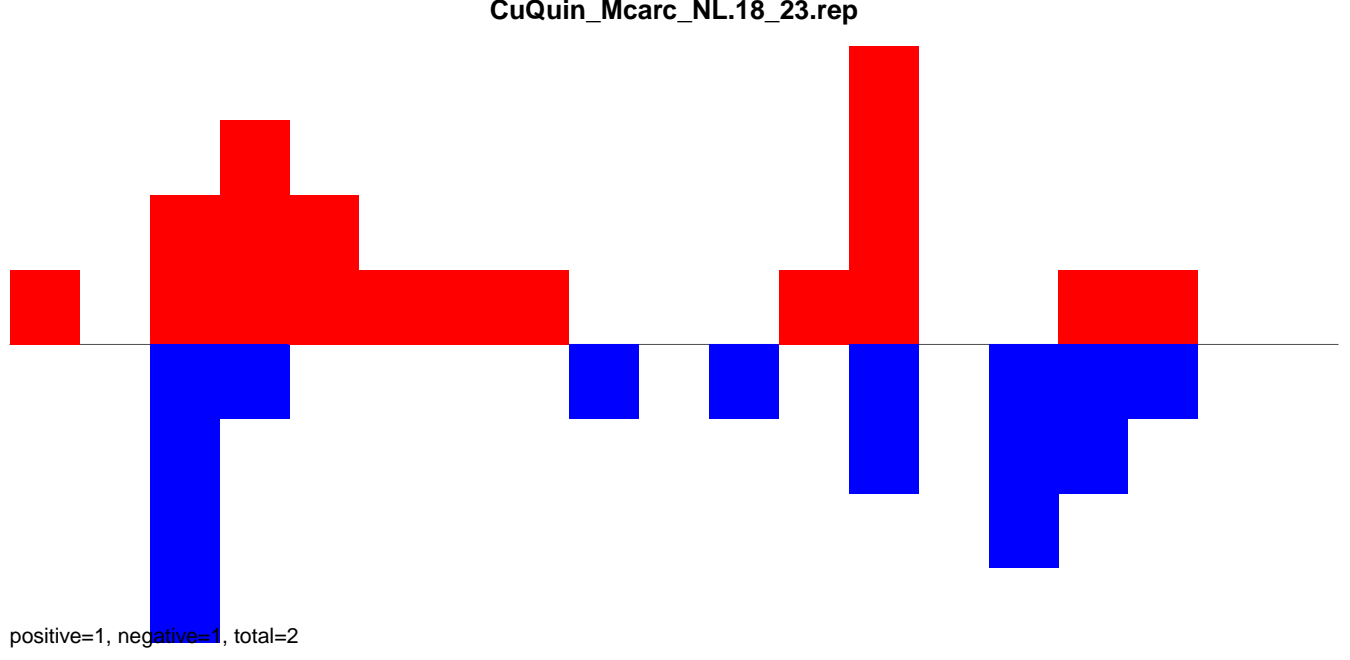
100

200

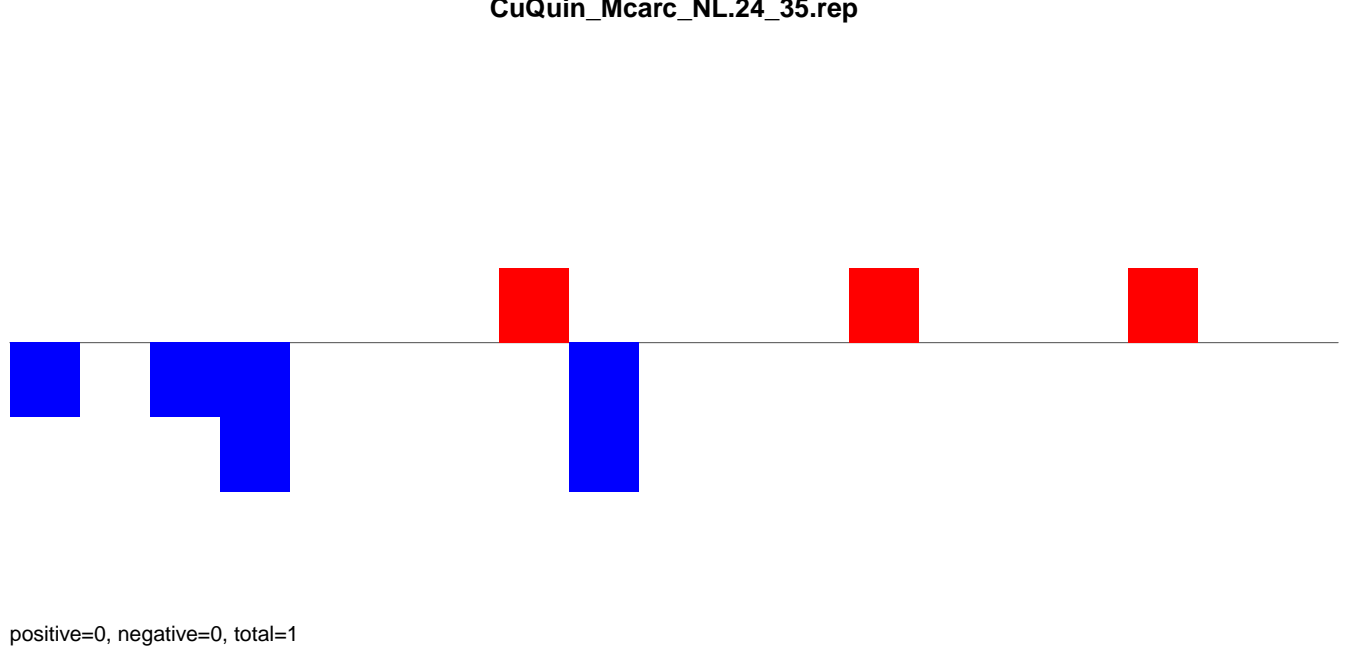
300

400

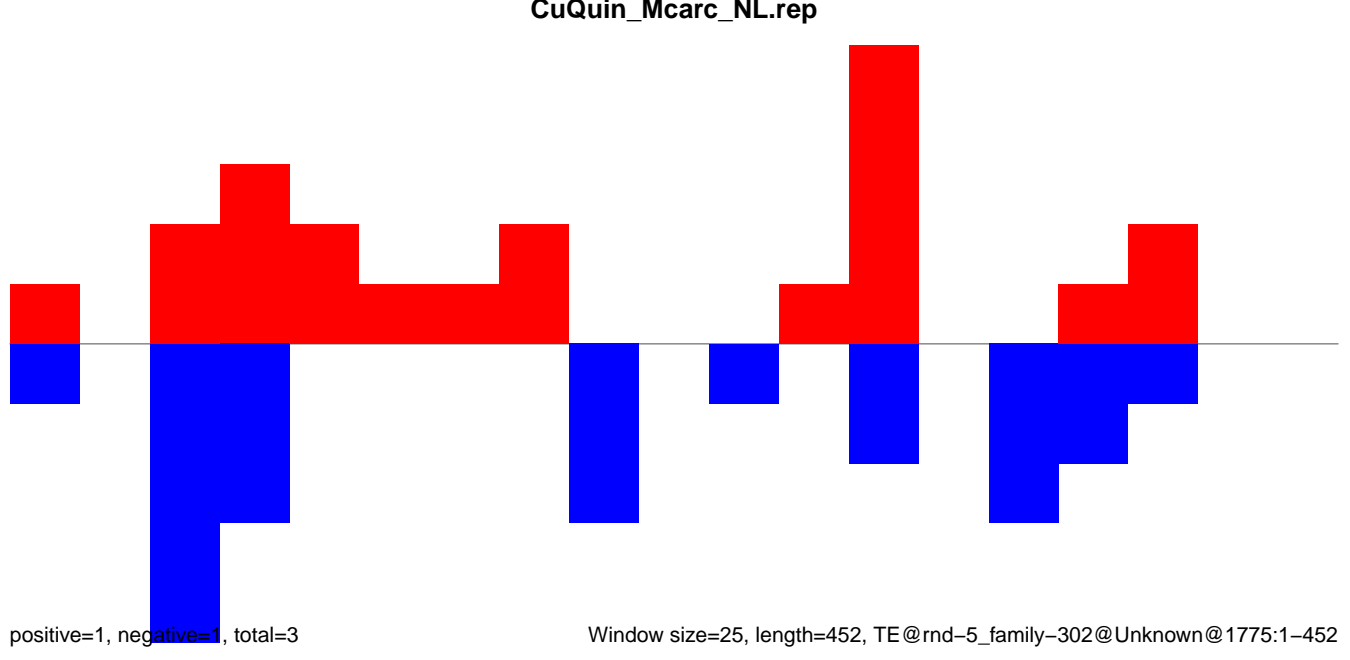
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



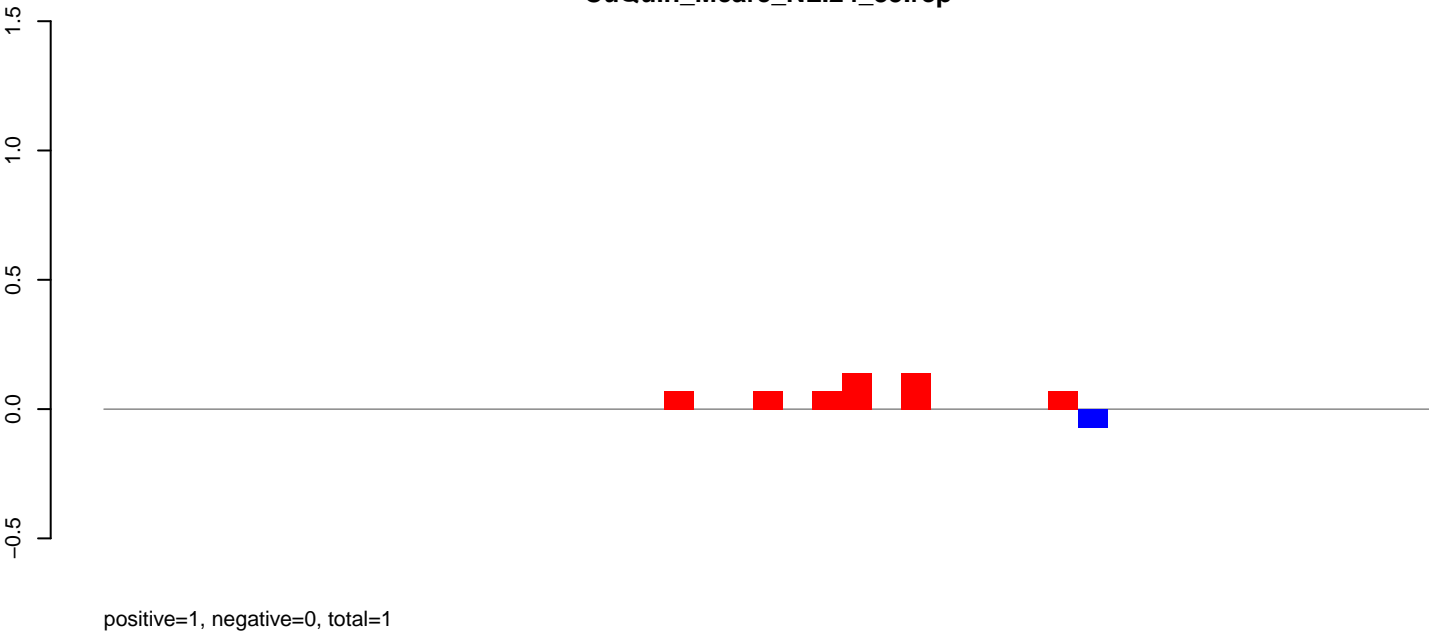
CuQuin_Mcarc_NL.rep



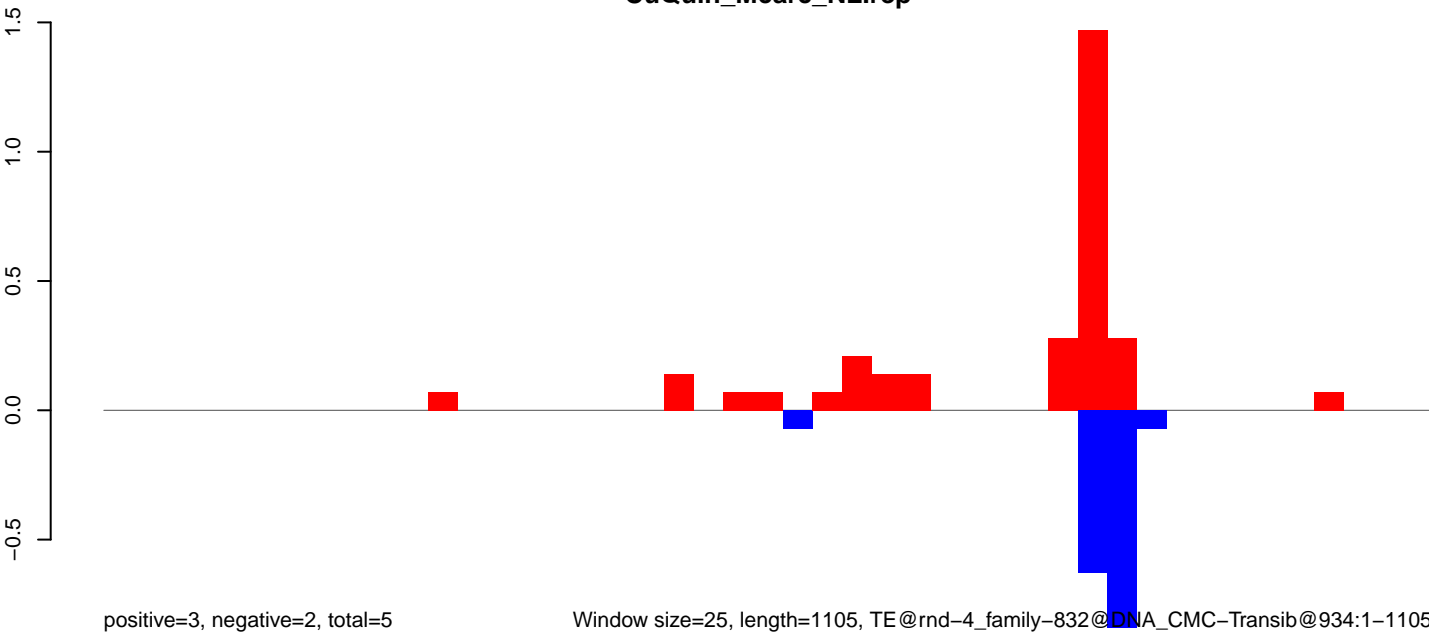
CuQuin_Mcarc_NL.18_23.rep



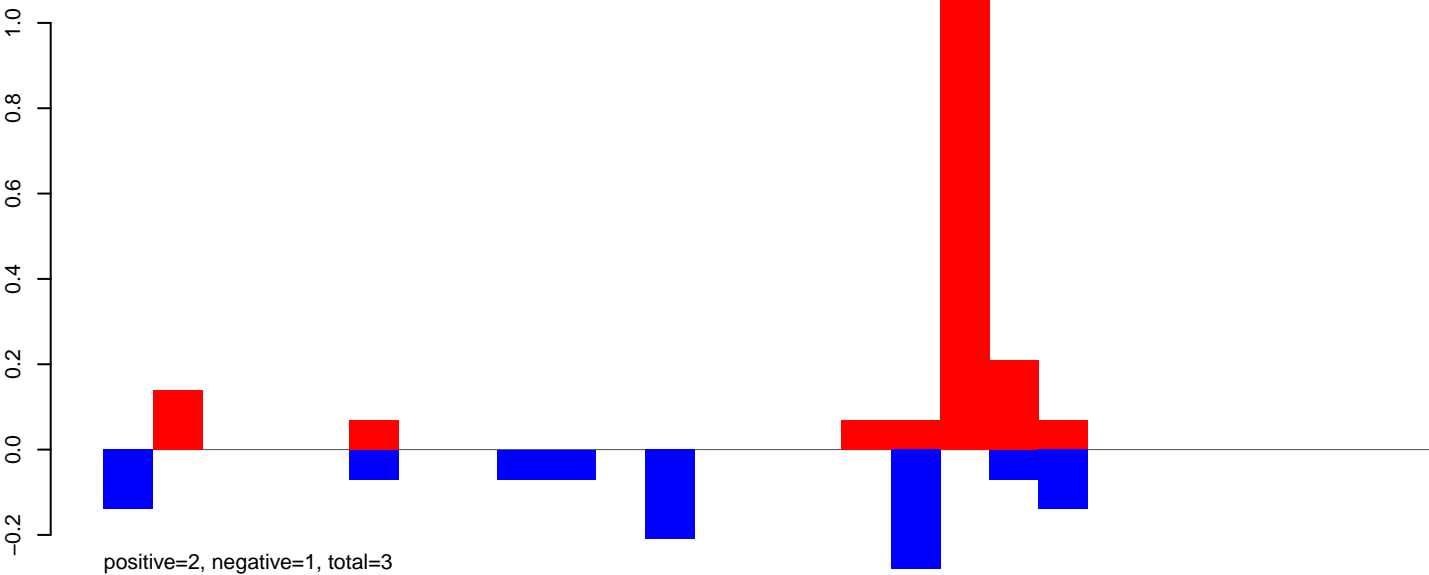
CuQuin_Mcarc_NL.24_35.rep



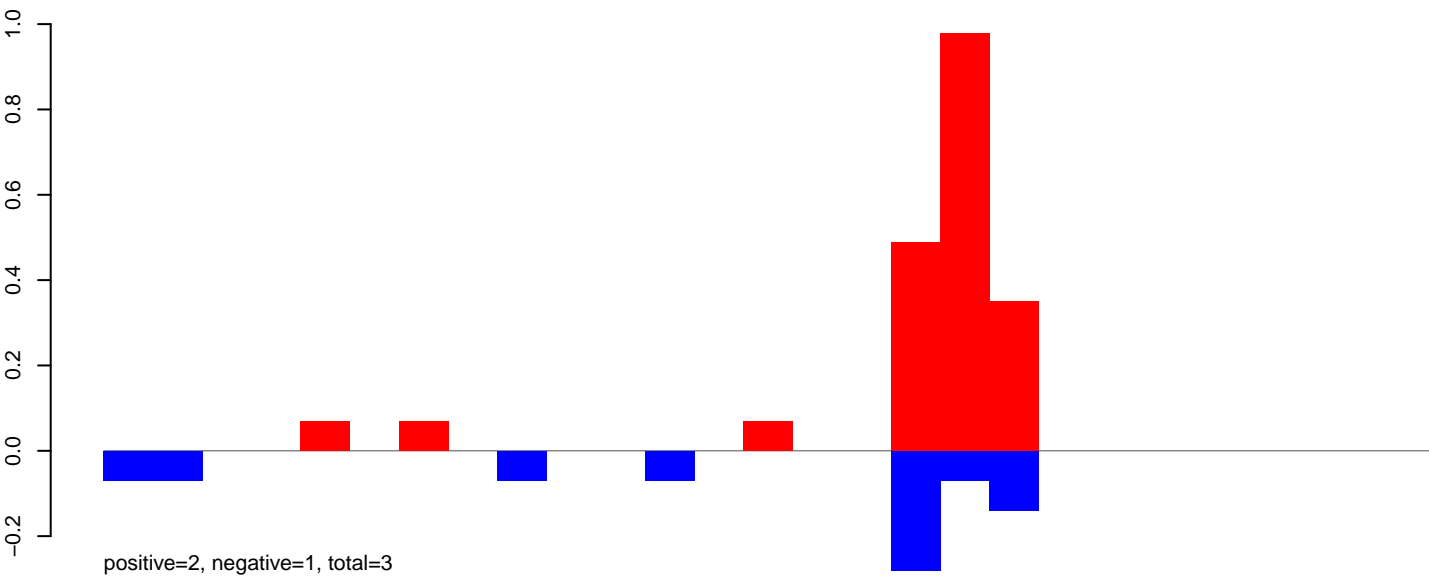
CuQuin_Mcarc_NL.rep



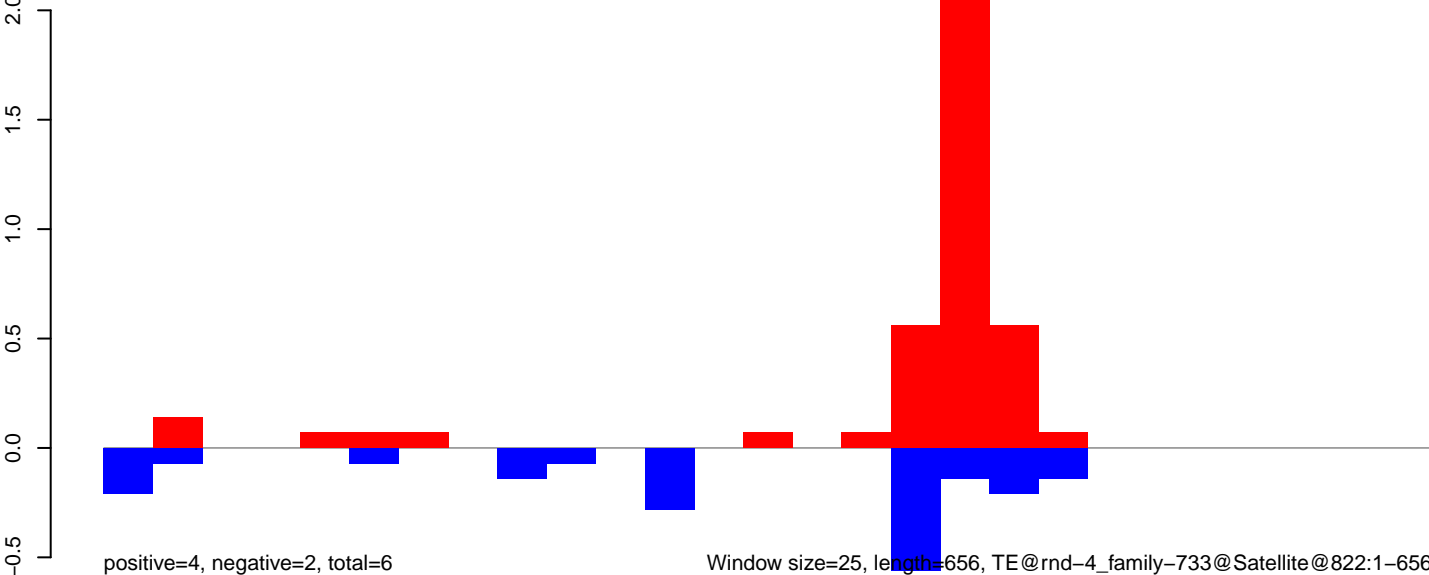
CuQuin_Mcarc_NL.18_23.rep



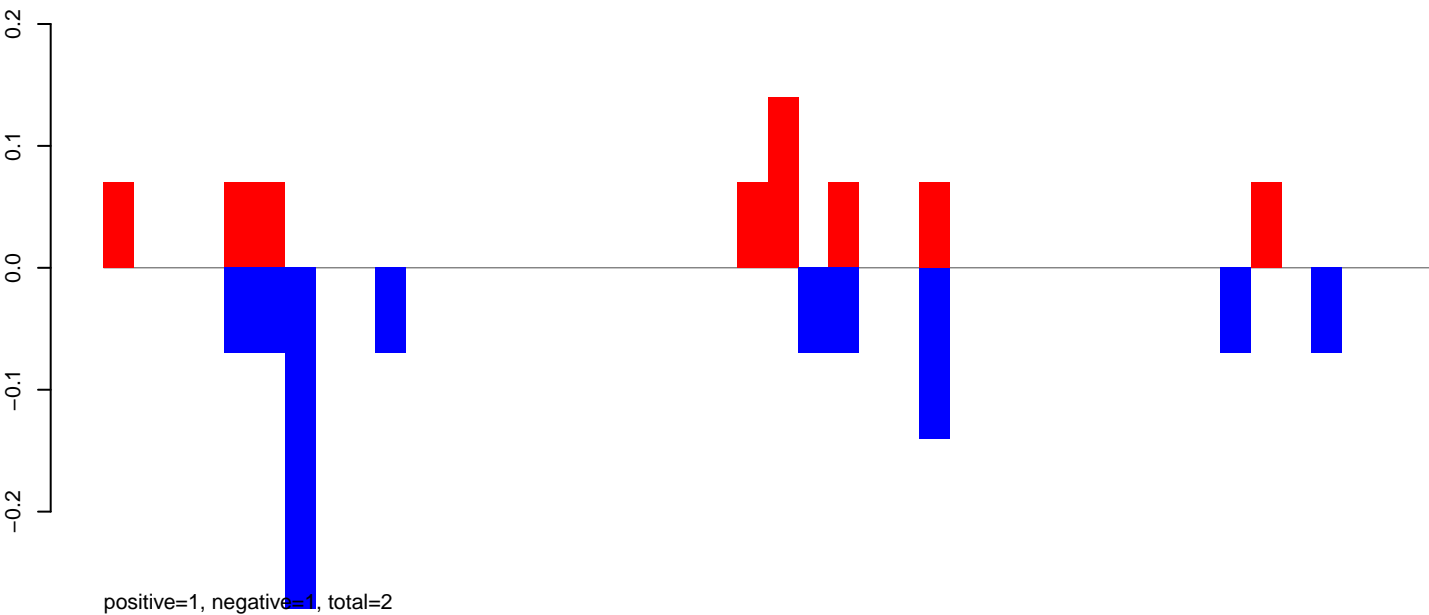
CuQuin_Mcarc_NL.24_35.rep



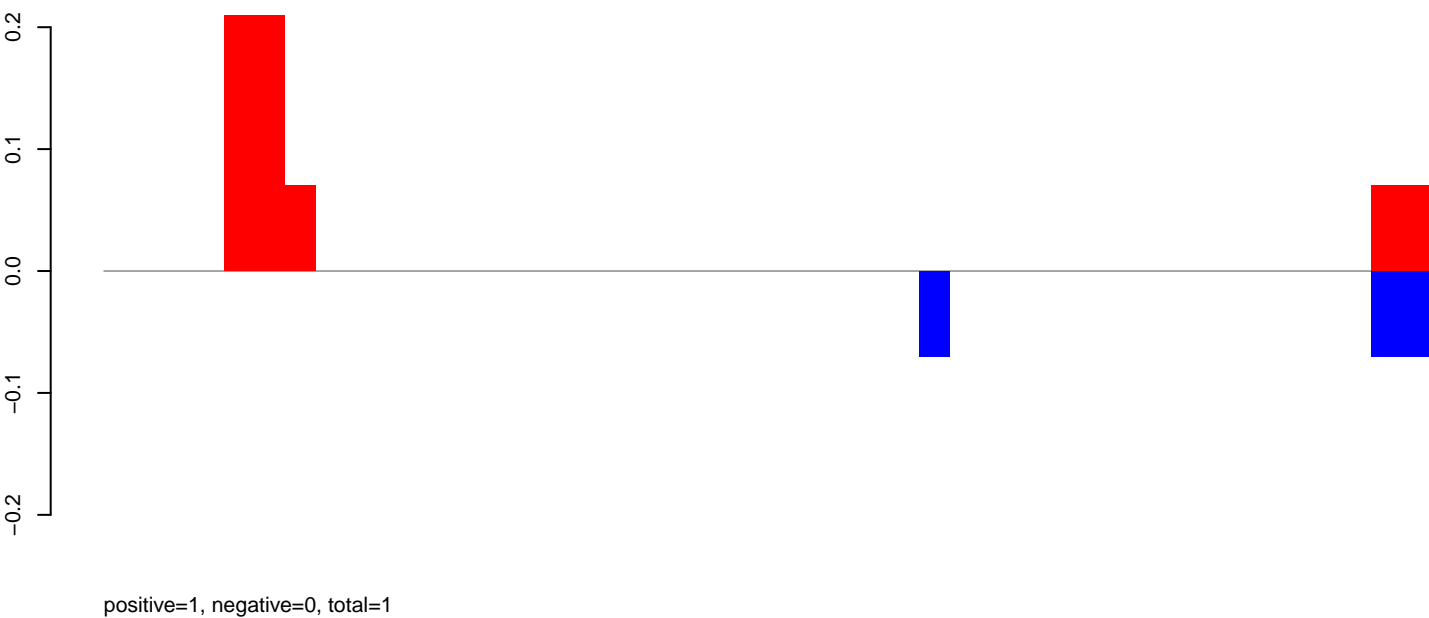
CuQuin_Mcarc_NL.rep



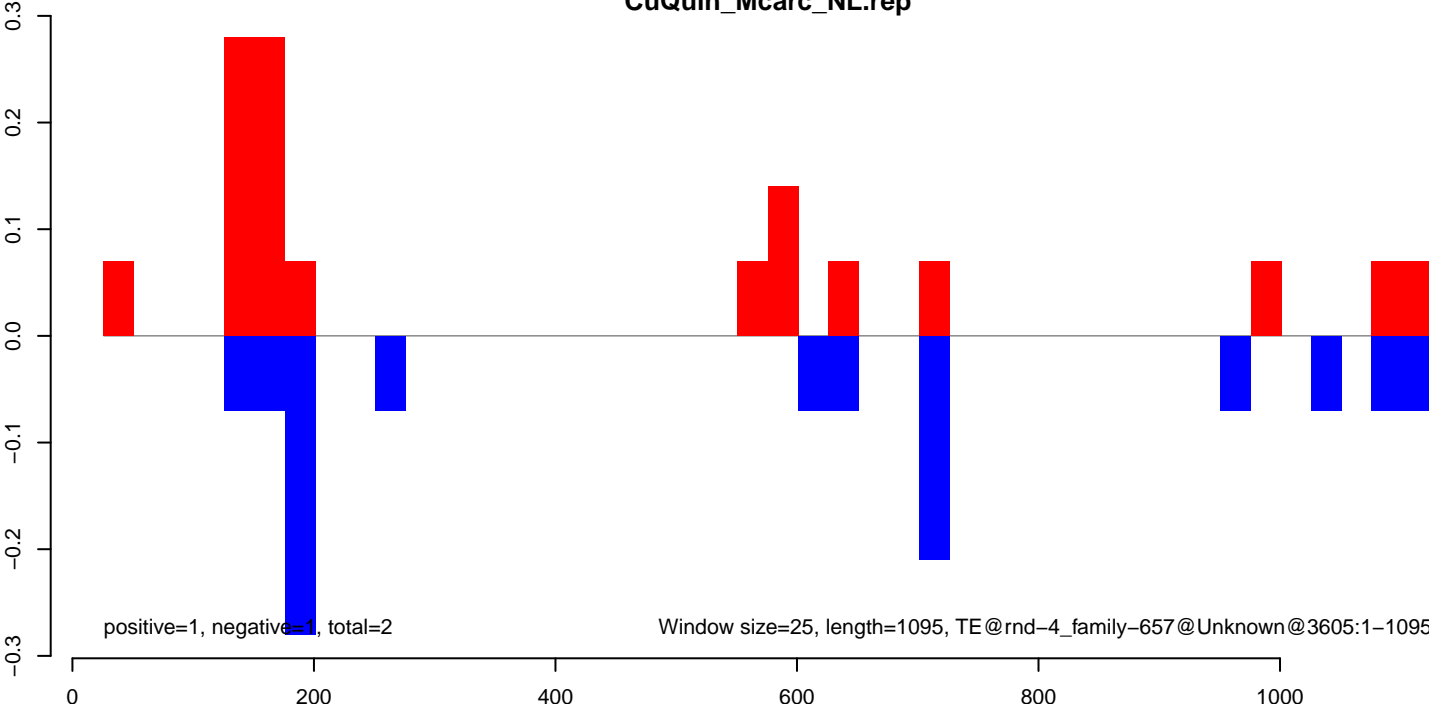
CuQuin_Mcarc_NL.18_23.rep



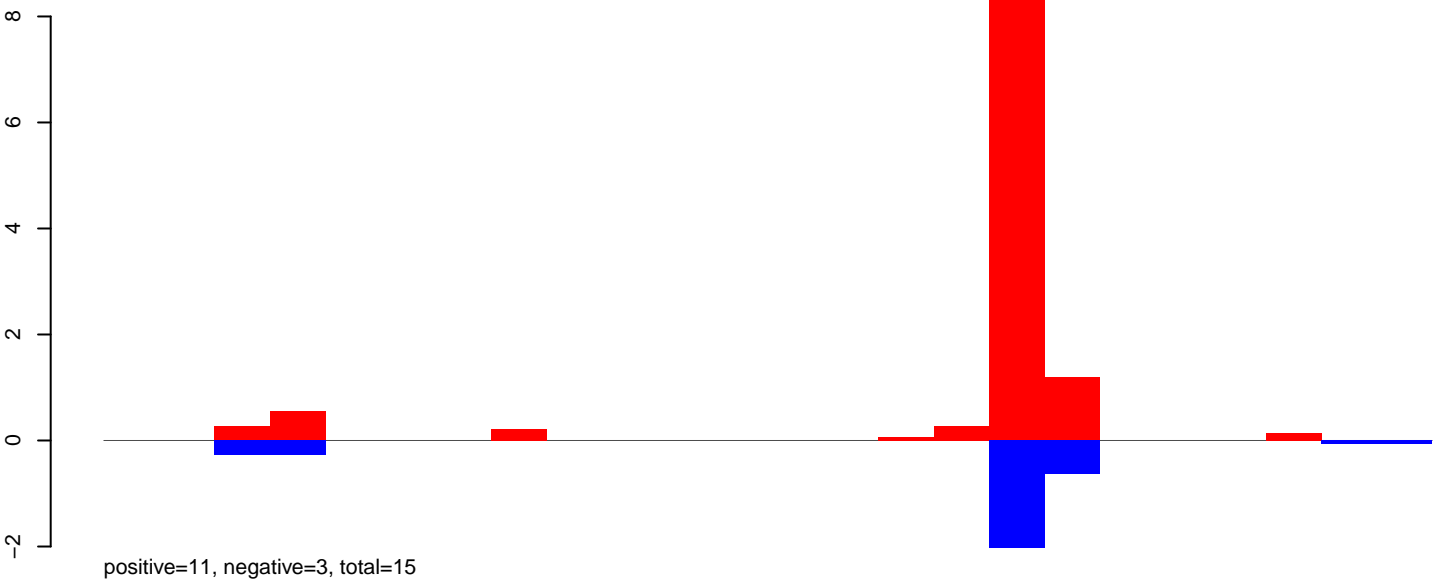
CuQuin_Mcarc_NL.24_35.rep



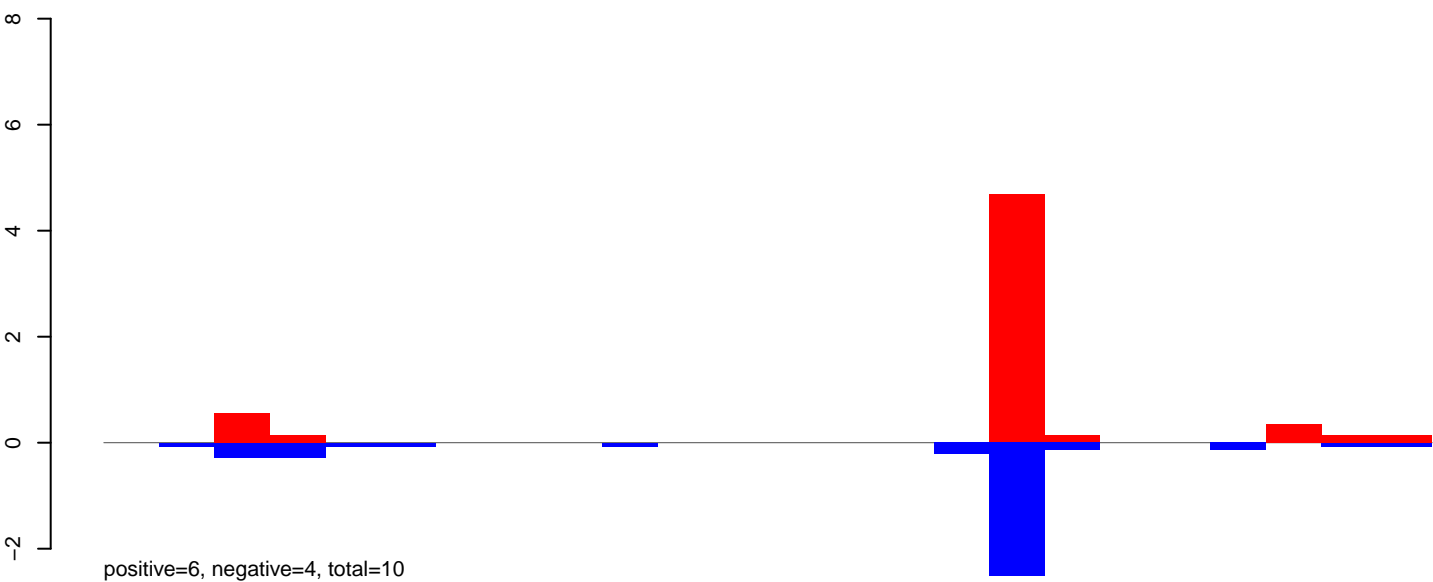
CuQuin_Mcarc_NL.rep



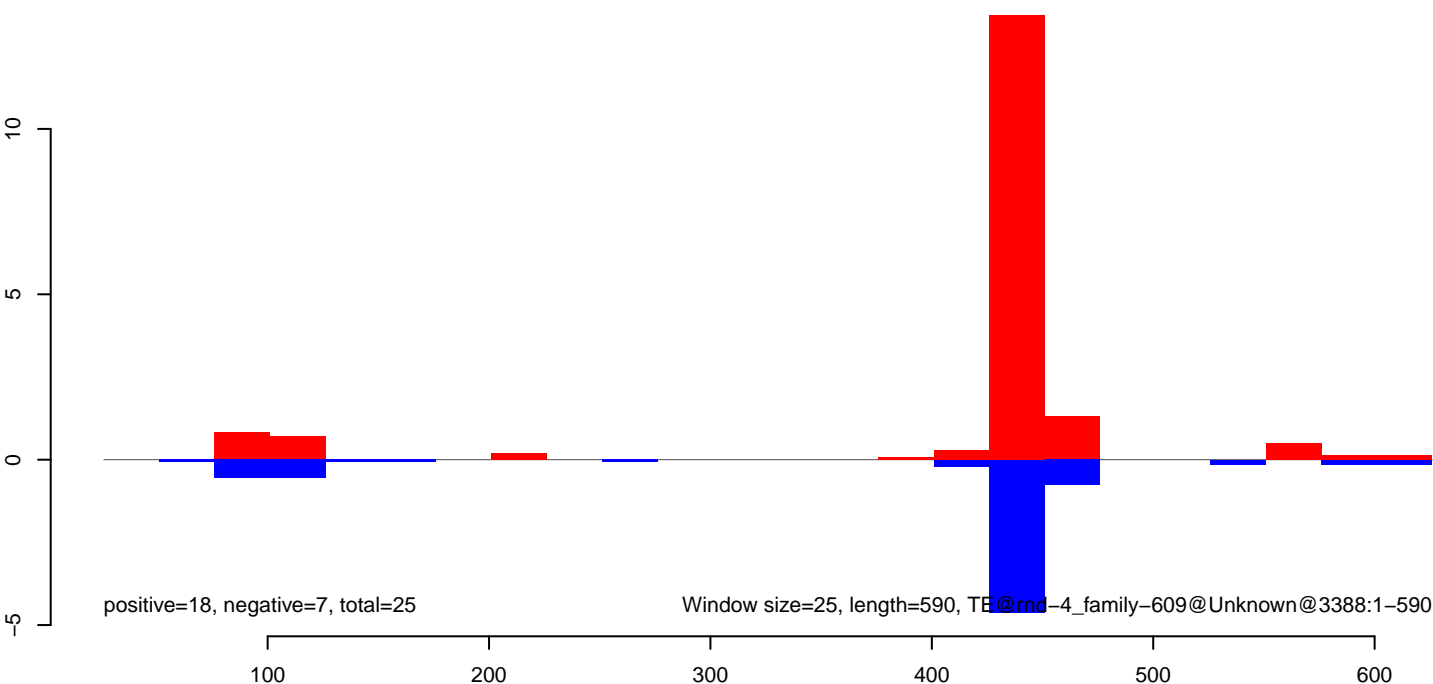
CuQuin_Mcarc_NL.18_23.rep



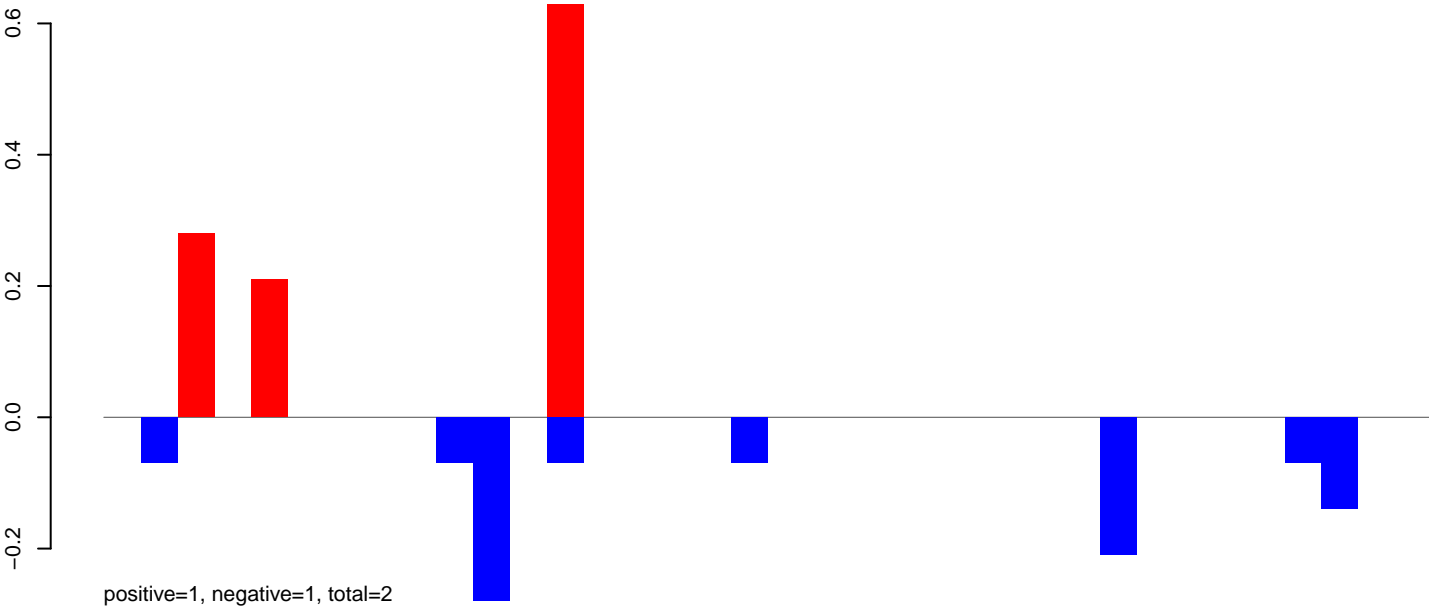
CuQuin_Mcarc_NL.24_35.rep



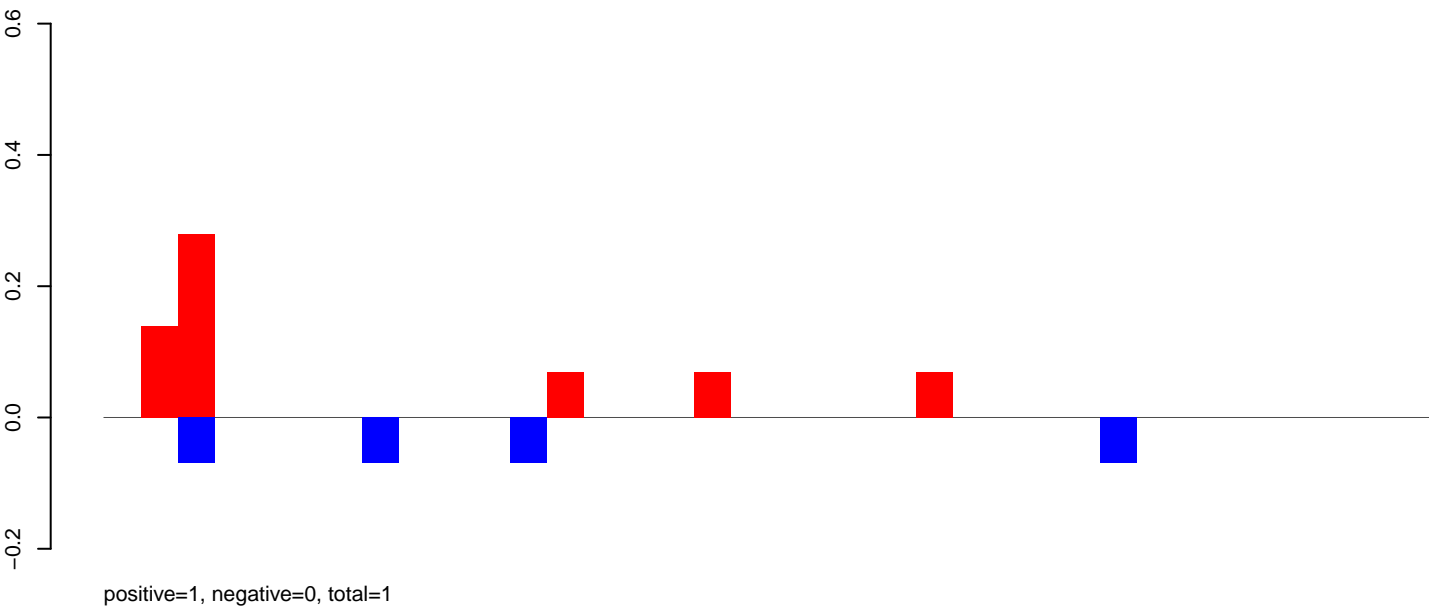
CuQuin_Mcarc_NL.rep



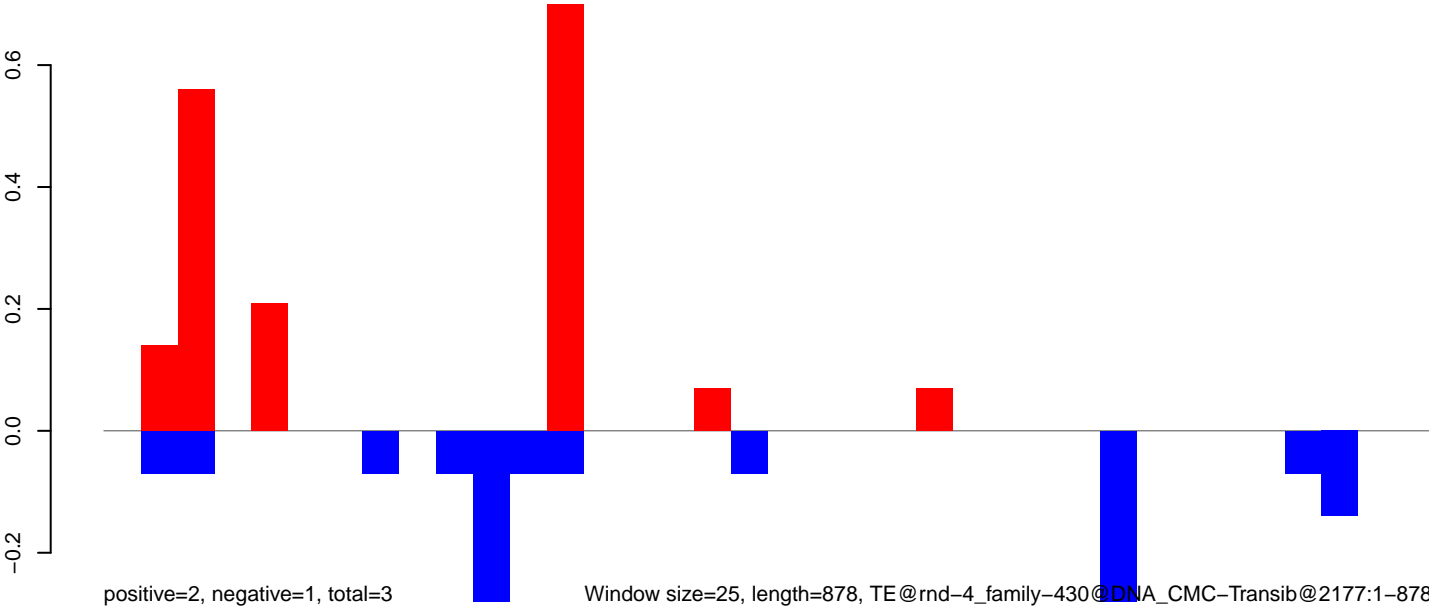
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

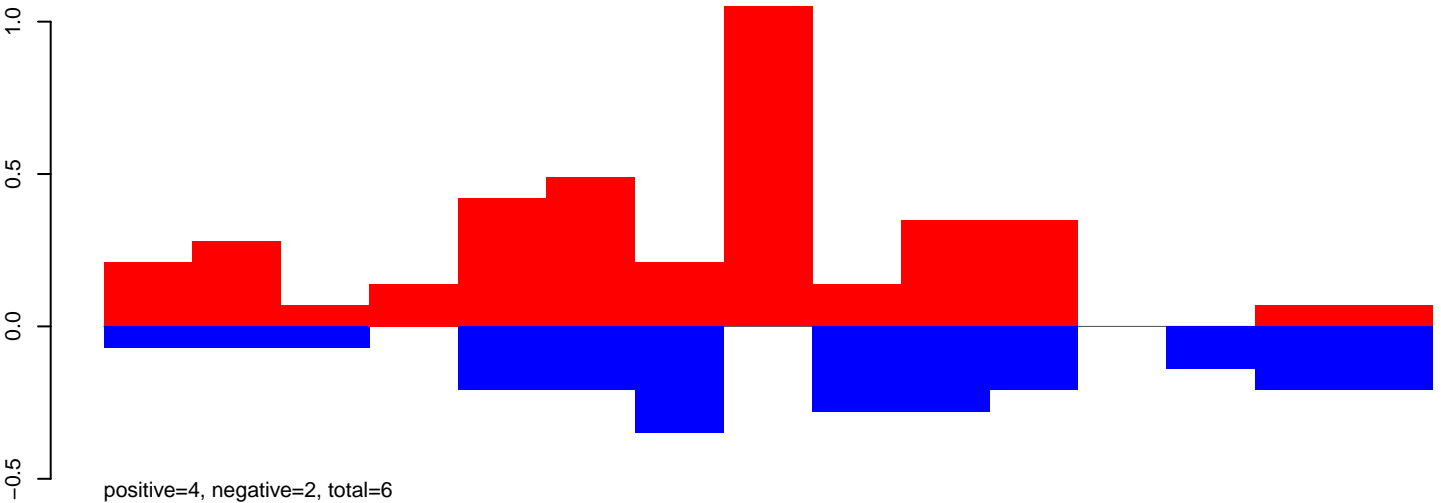


CuQuin_Mcarc_NL.rep

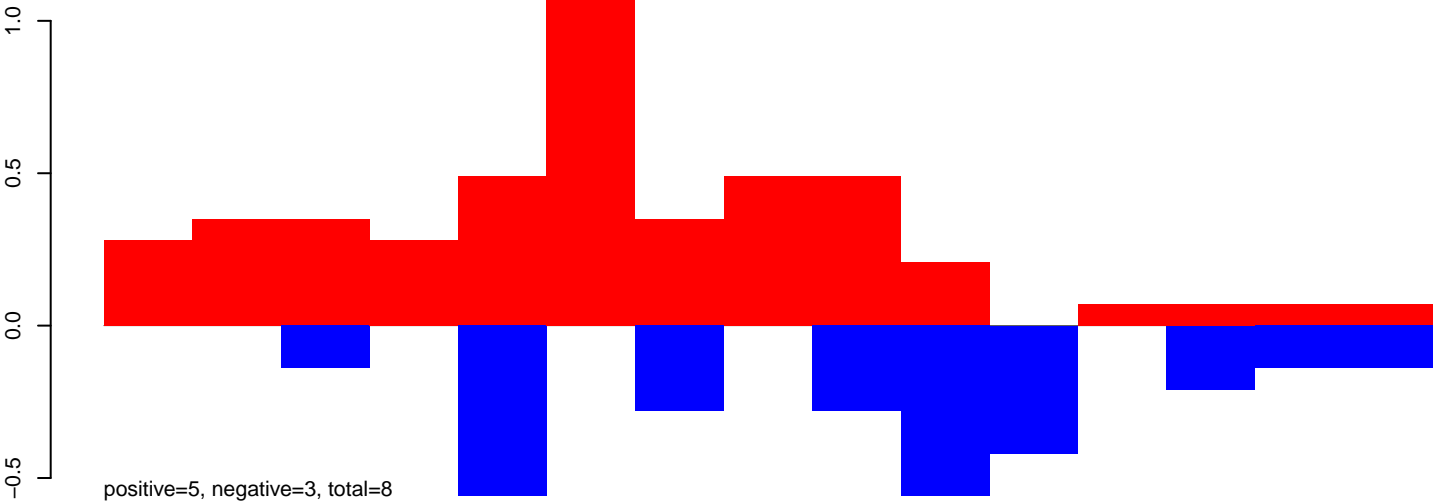


Window size=25, length=878, TE@rnd-4_family-430@DNA_CMC-Transib@2177:1-878

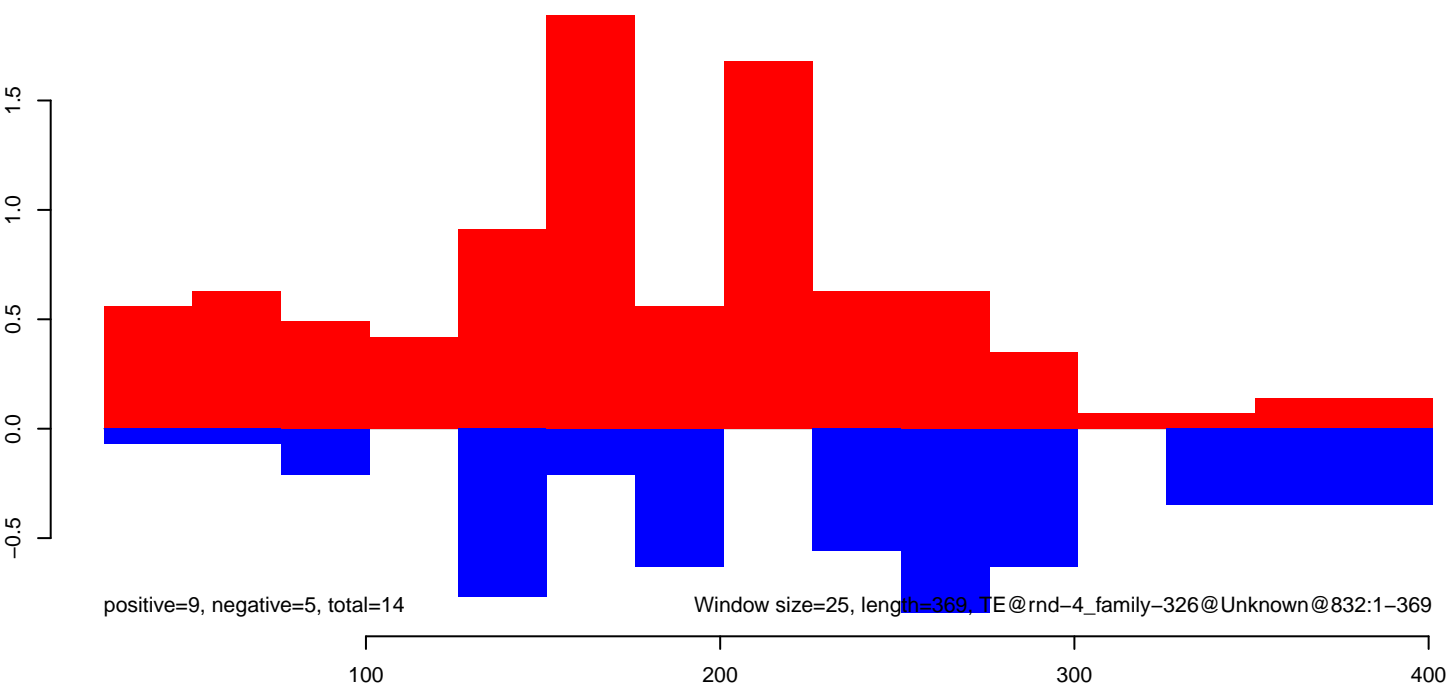
CuQuin_Mcarc_NL.18_23.rep



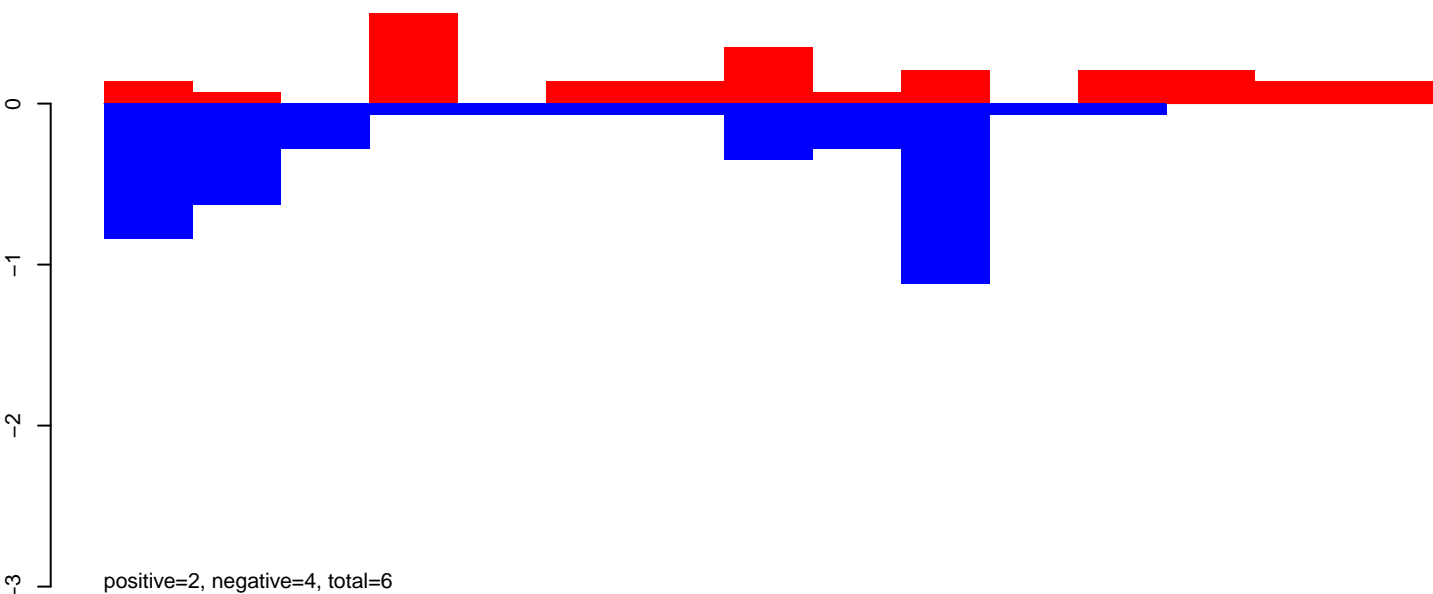
CuQuin_Mcarc_NL.24_35.rep



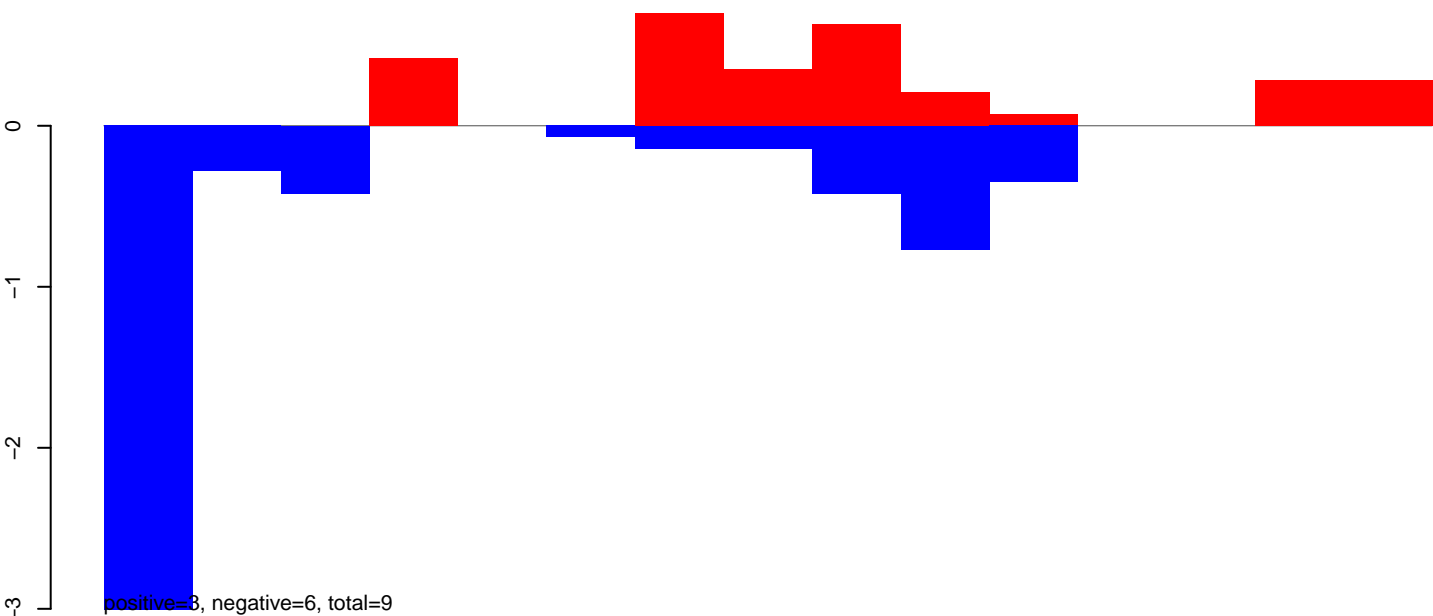
CuQuin_Mcarc_NL.rep



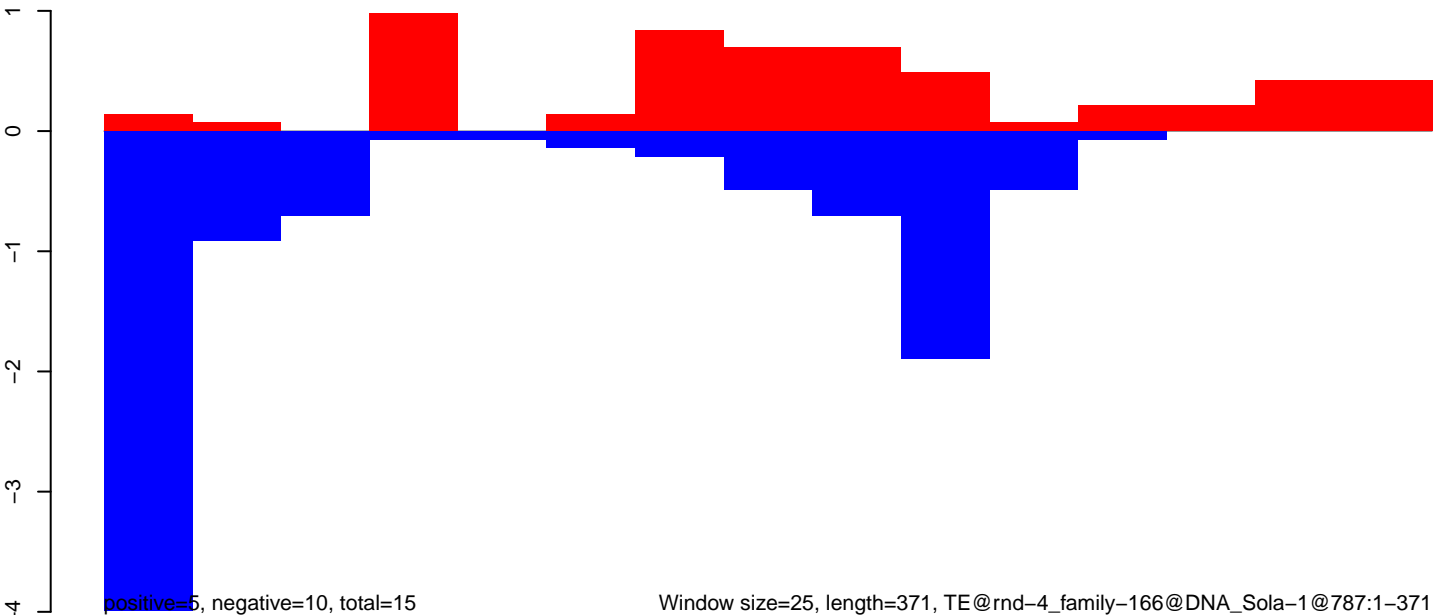
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



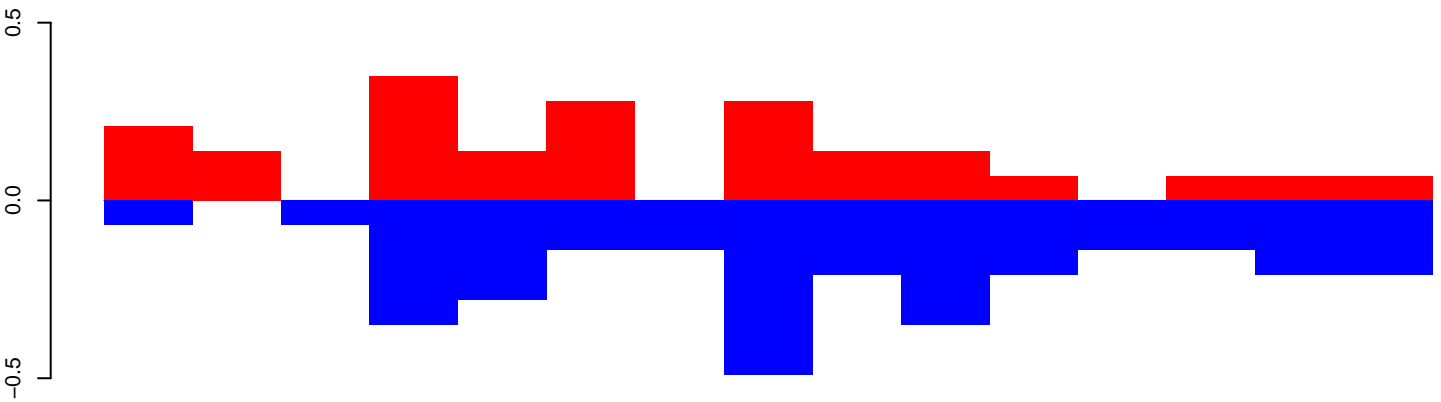
CuQuin_Mcarc_NL.rep



Window size=25, length=371, TE@rnd-4_family-166@DNA_Sola-1@787:1-371

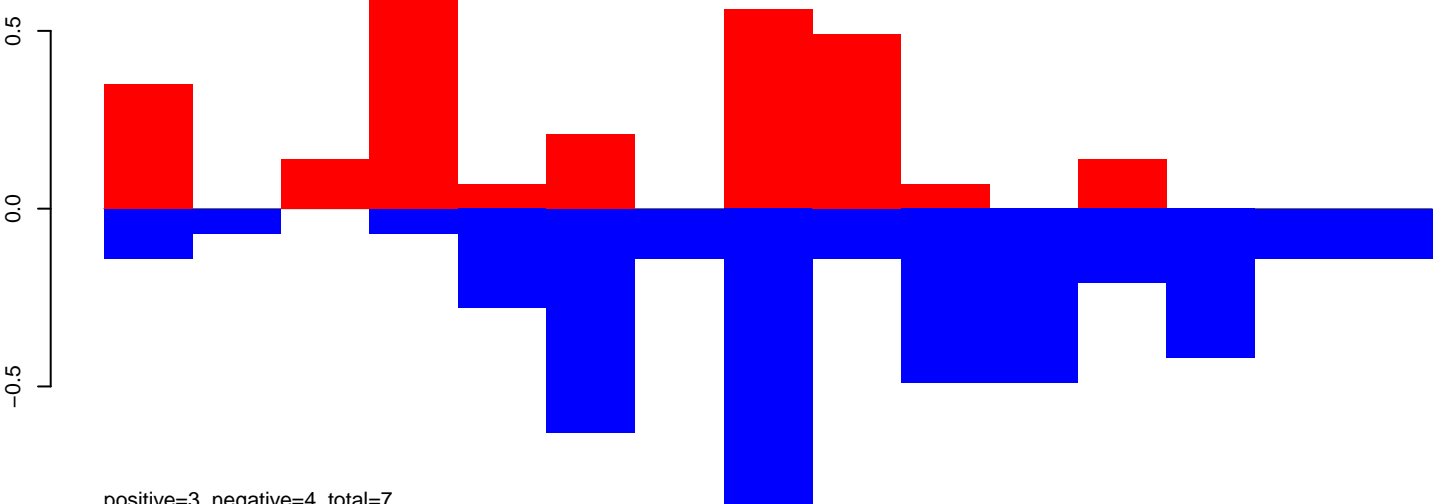
100 200 300 400

CuQuin_Mcarc_NL.18_23.rep



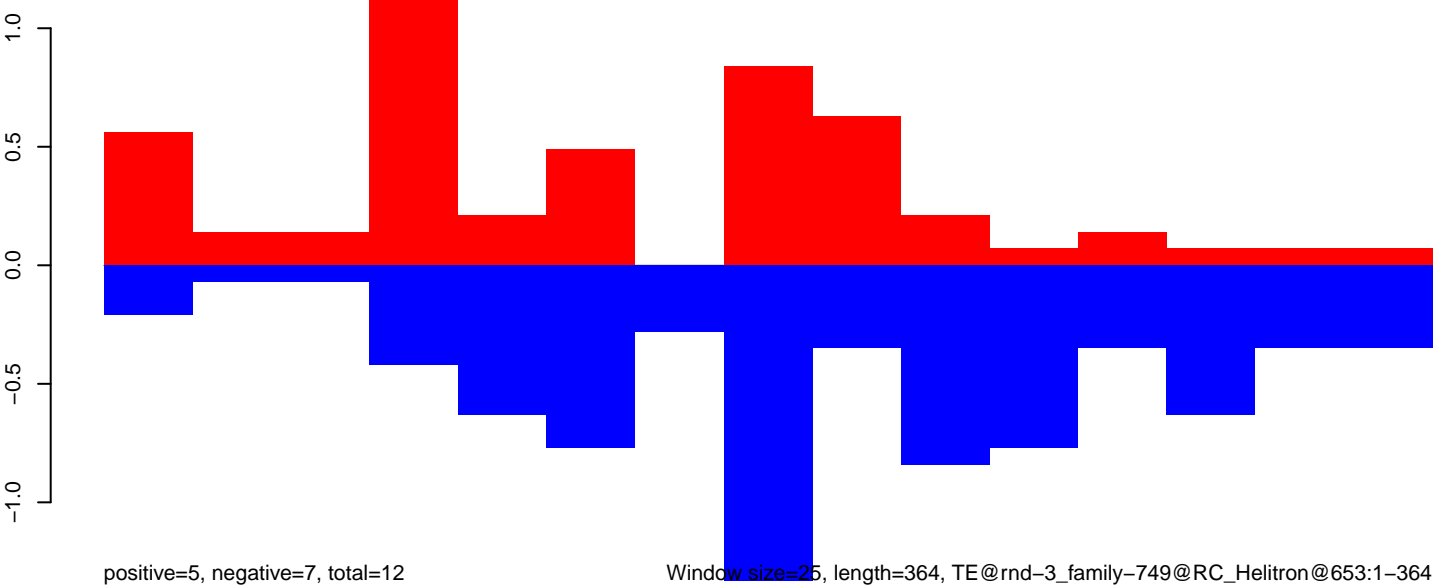
positive=2, negative=3, total=5

CuQuin_Mcarc_NL.24_35.rep



positive=3, negative=4, total=7

CuQuin_Mcarc_NL.rep



positive=5, negative=7, total=12

Window size=25, length=364, TE@rnd-3_family-749@RC_Helitron@653:1-364

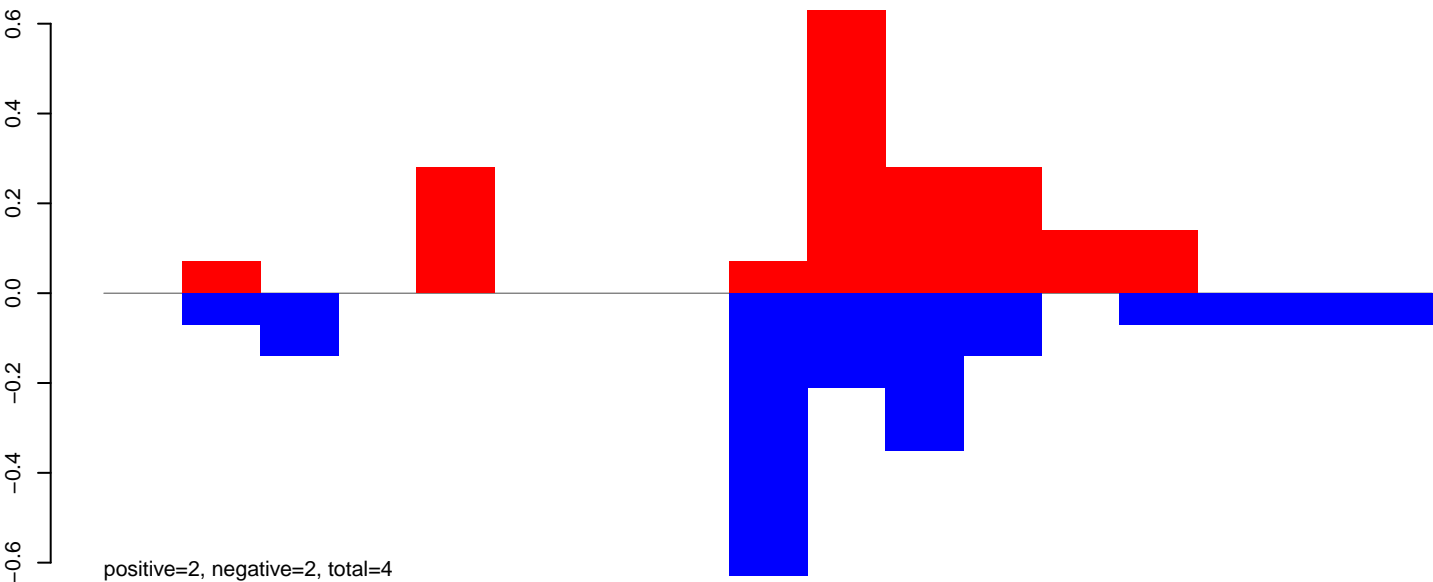
100

200

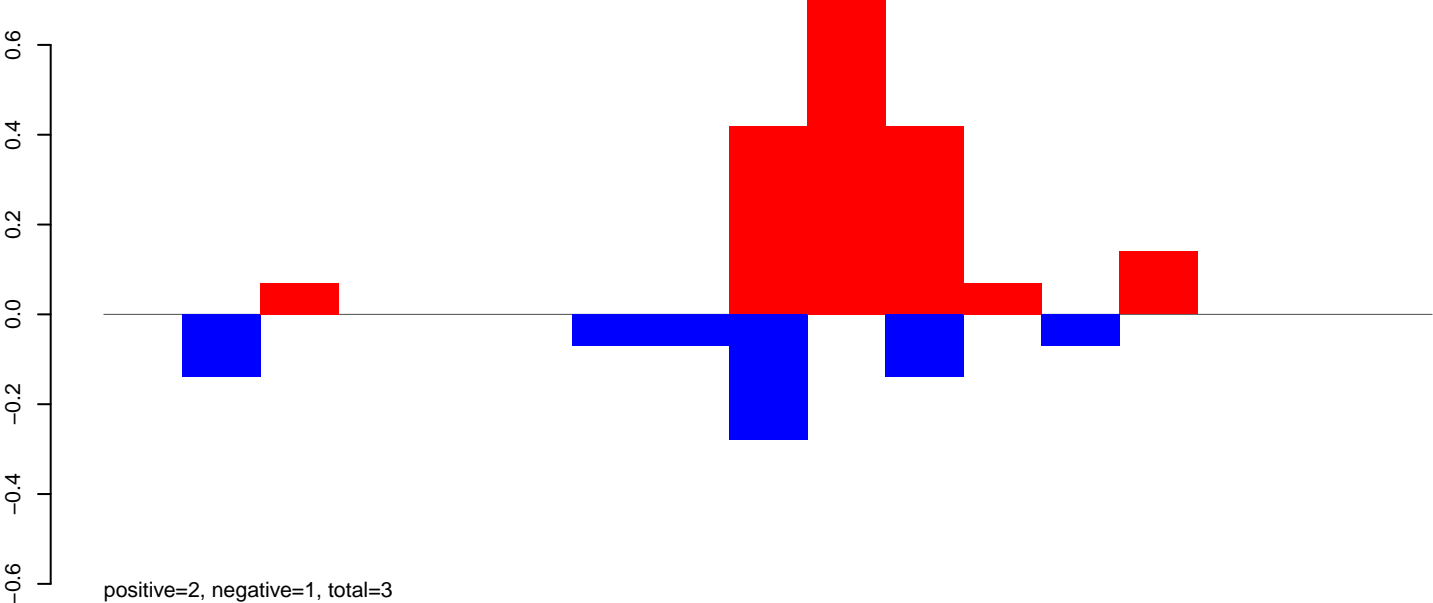
300

400

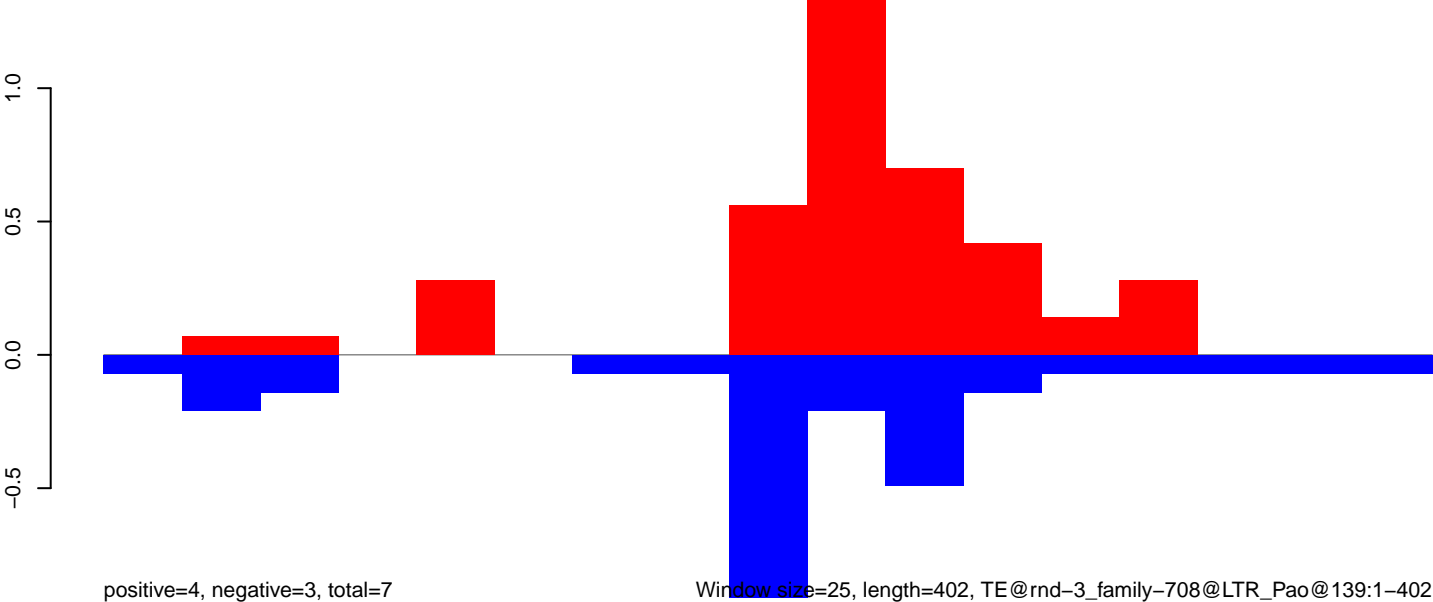
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=402, TE@rnd-3_family-708@LTR_Pao@139:1-402

100

200

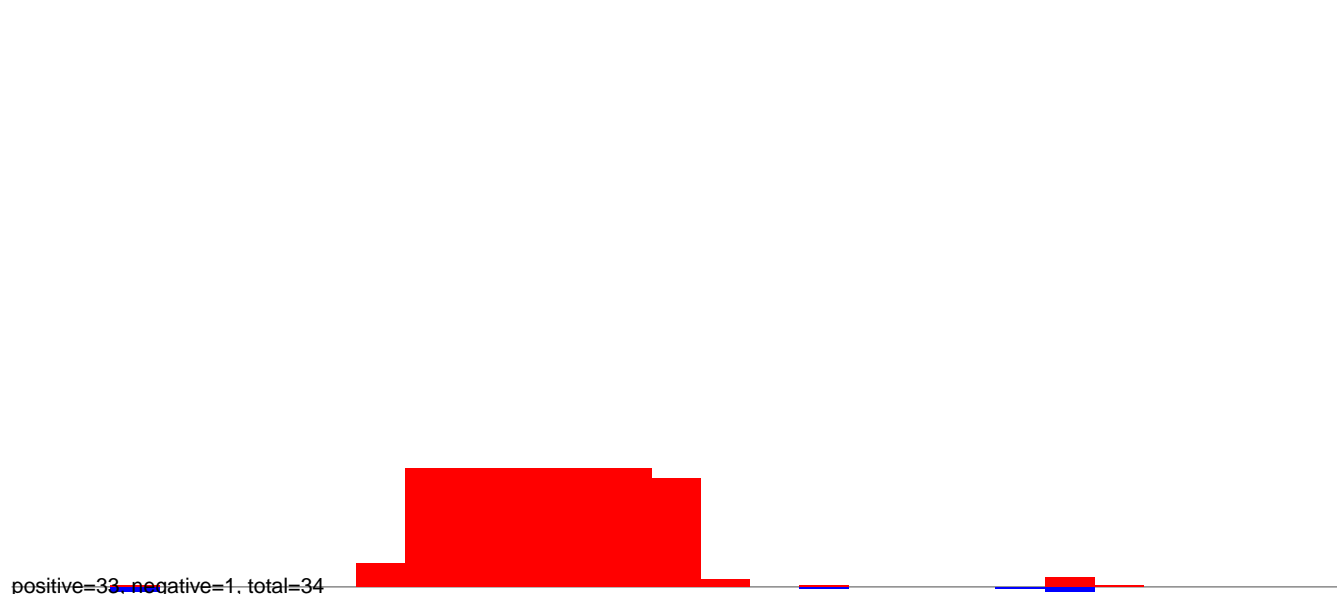
300

400

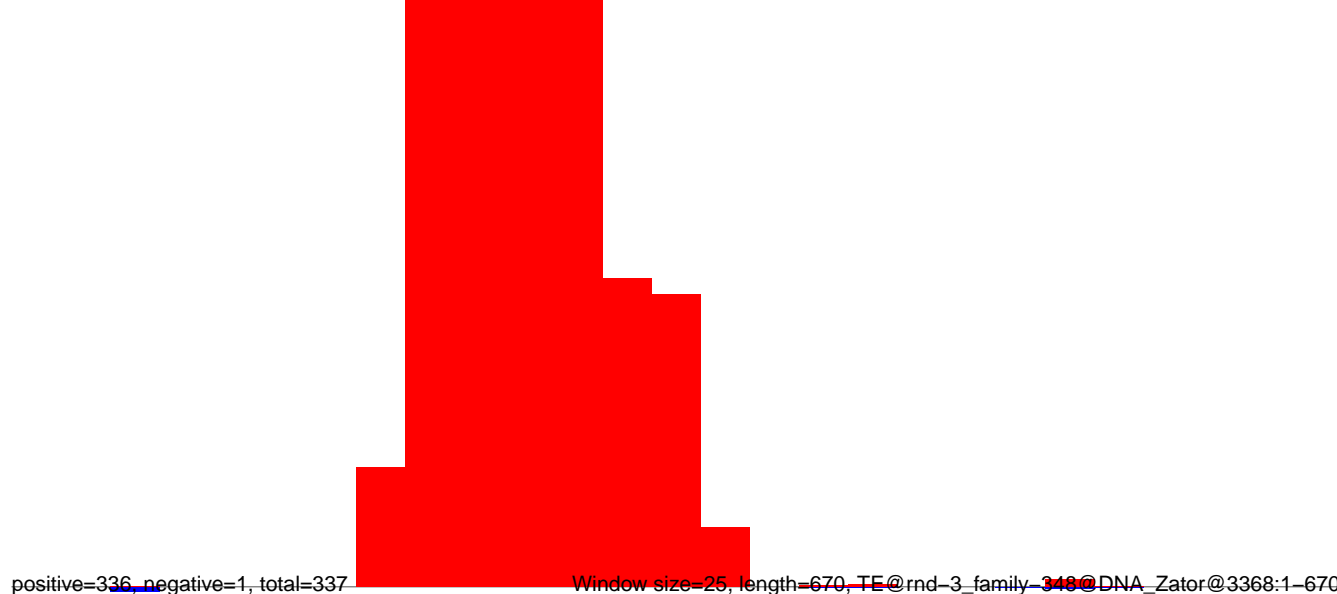
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

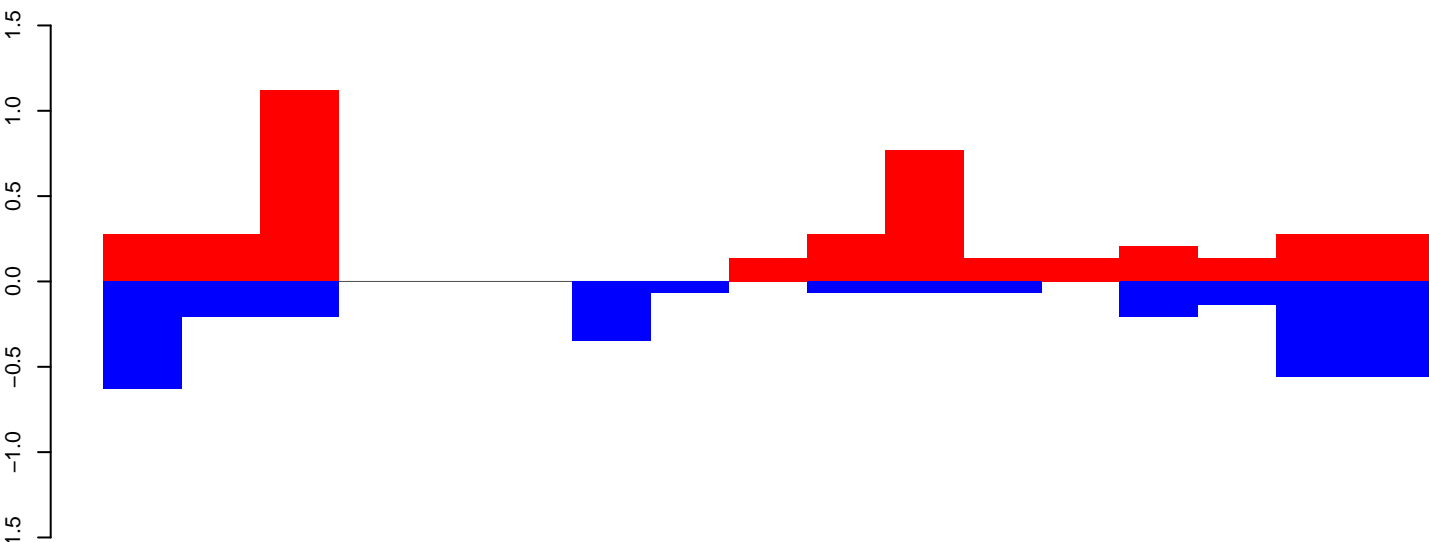


CuQuin_Mcarc_NL.rep



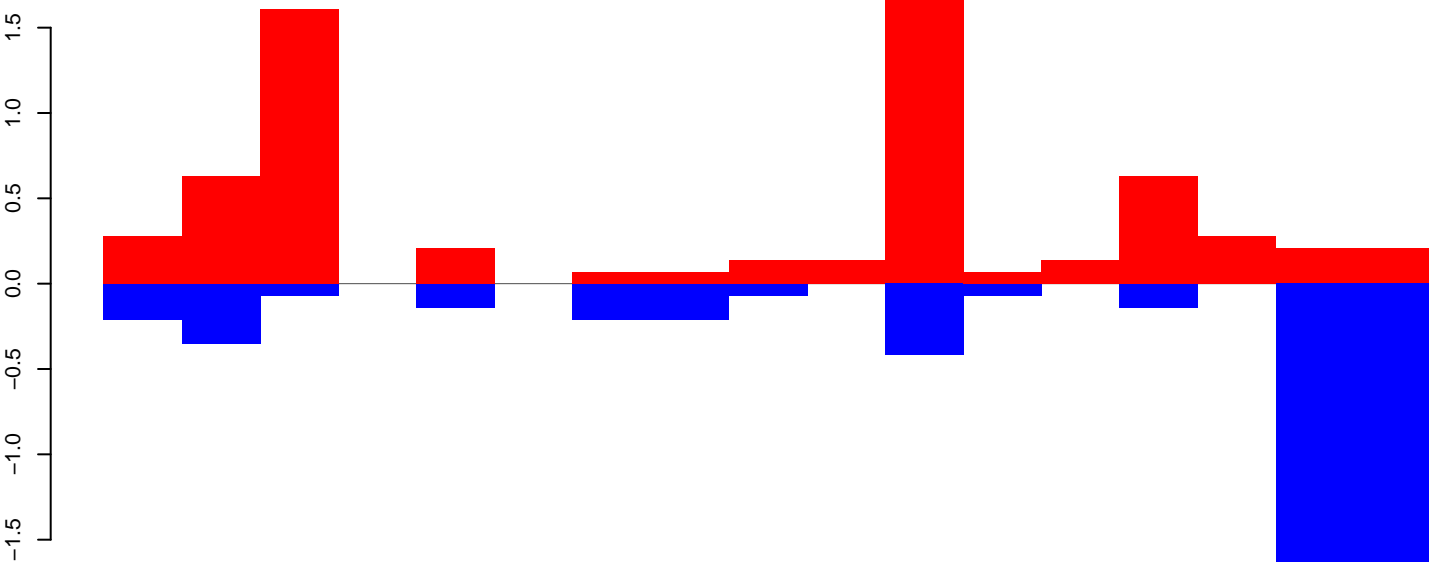
Window size=25, length=670, TE@rnd-3_family-348@DNA_Zator@3368:1-670

CuQuin_Mcarc_NL.18_23.rep



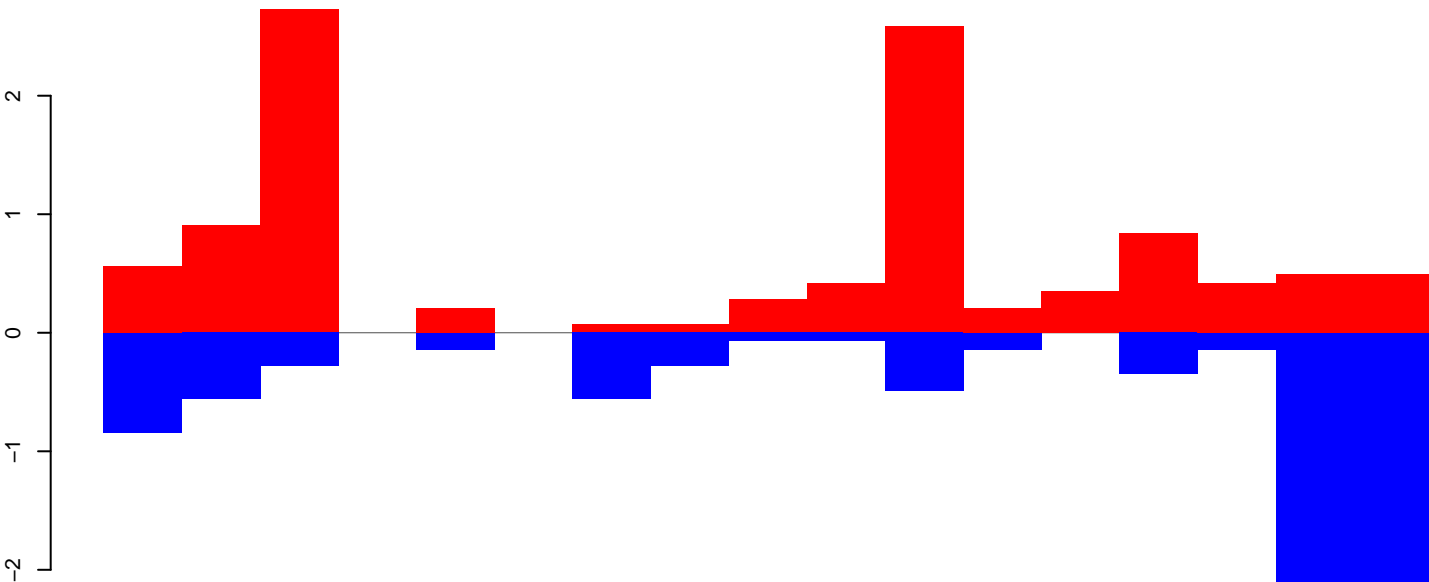
positive=4, negative=3, total=7

CuQuin_Mcarc_NL.24_35.rep



positive=6, negative=5, total=12

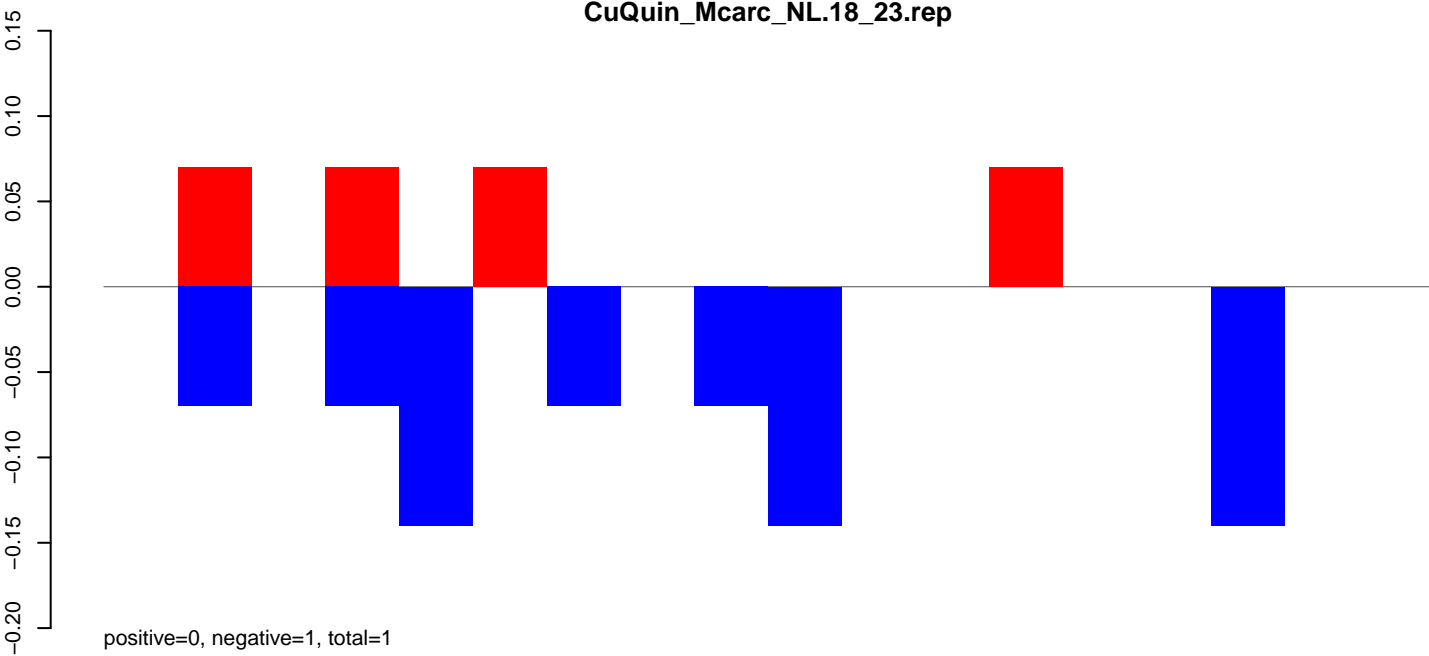
CuQuin_Mcarc_NL.rep



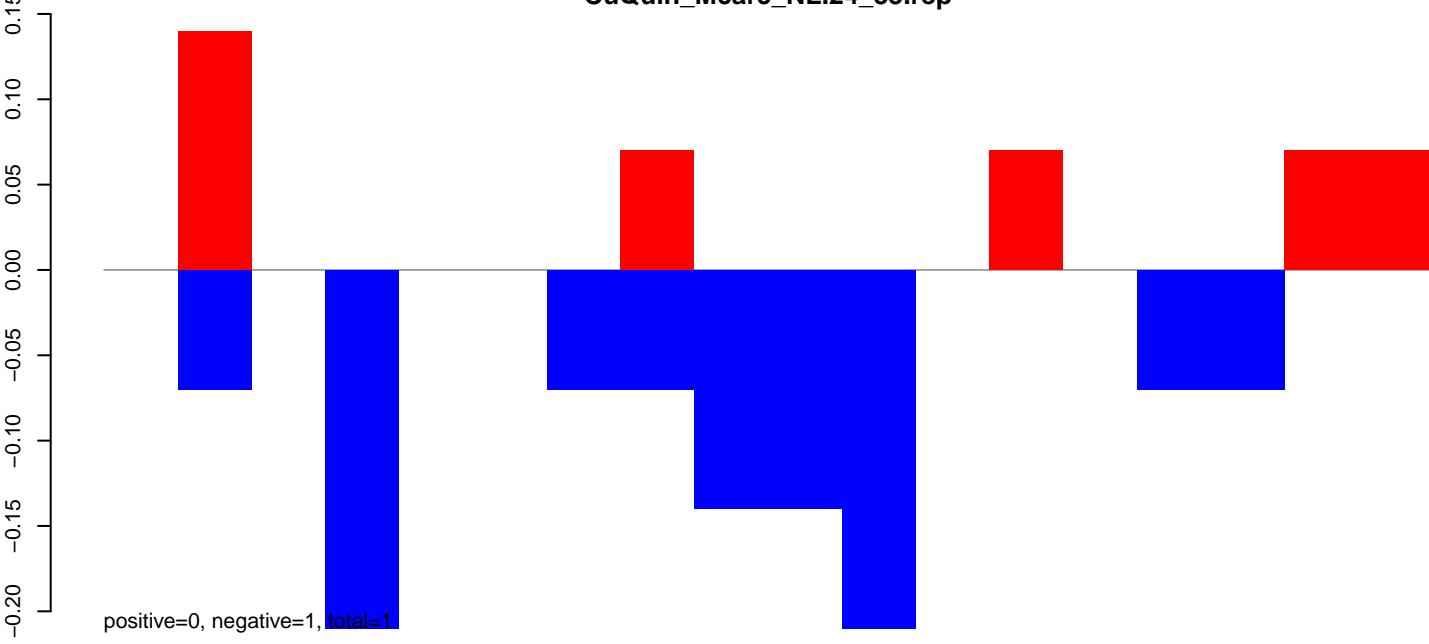
positive=11, negative=9, total=19

Window size=25, length=421, TE@rnd-3_family-326@DNA_TcMar@48:1-421

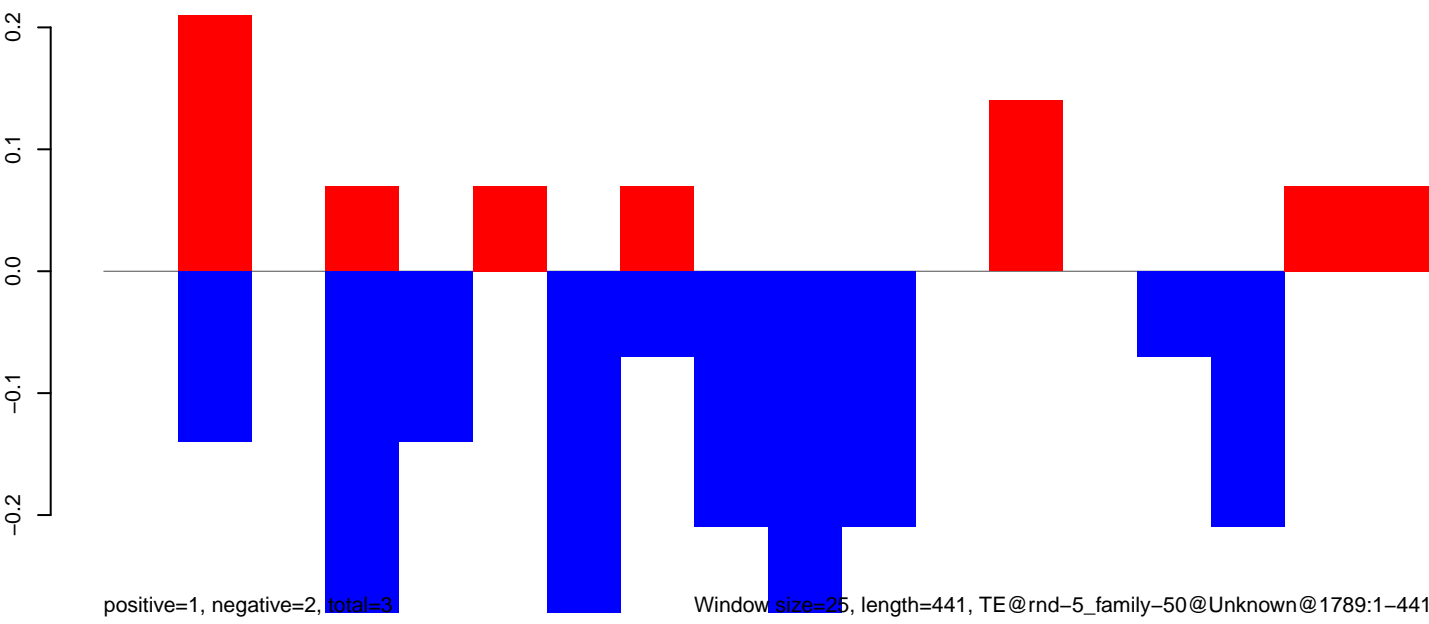
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

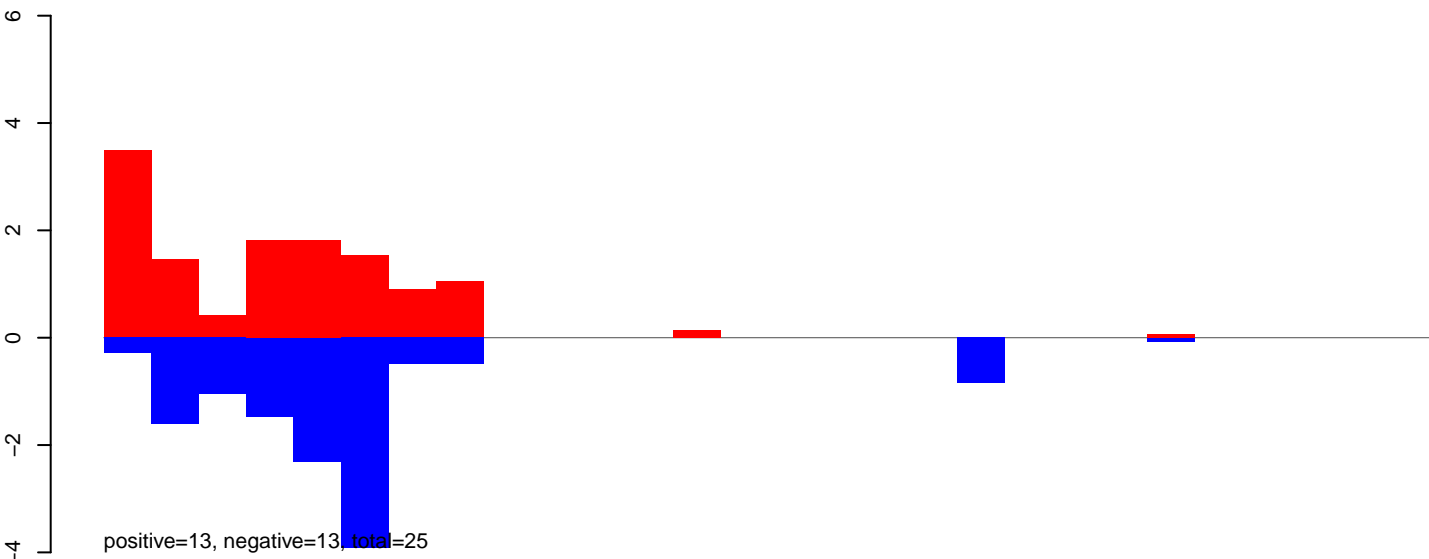


CuQuin_Mcarc_NL.rep

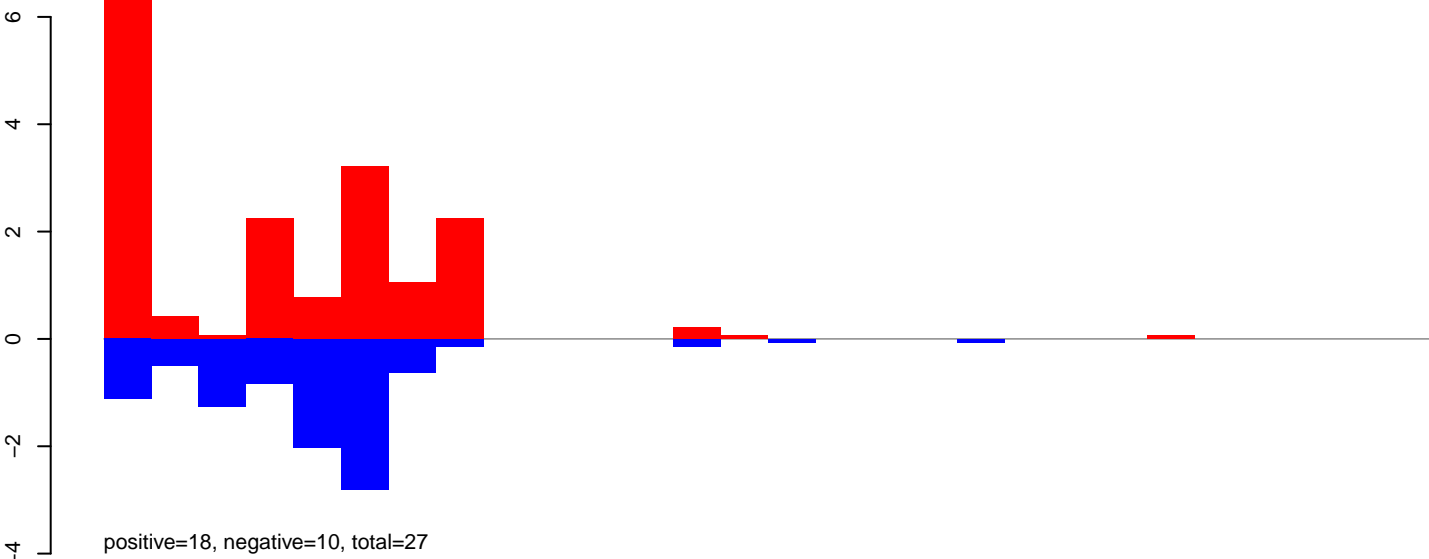


Window size=25, length=441, TE@rnd-5_family-50@Unknown@1789:1-441

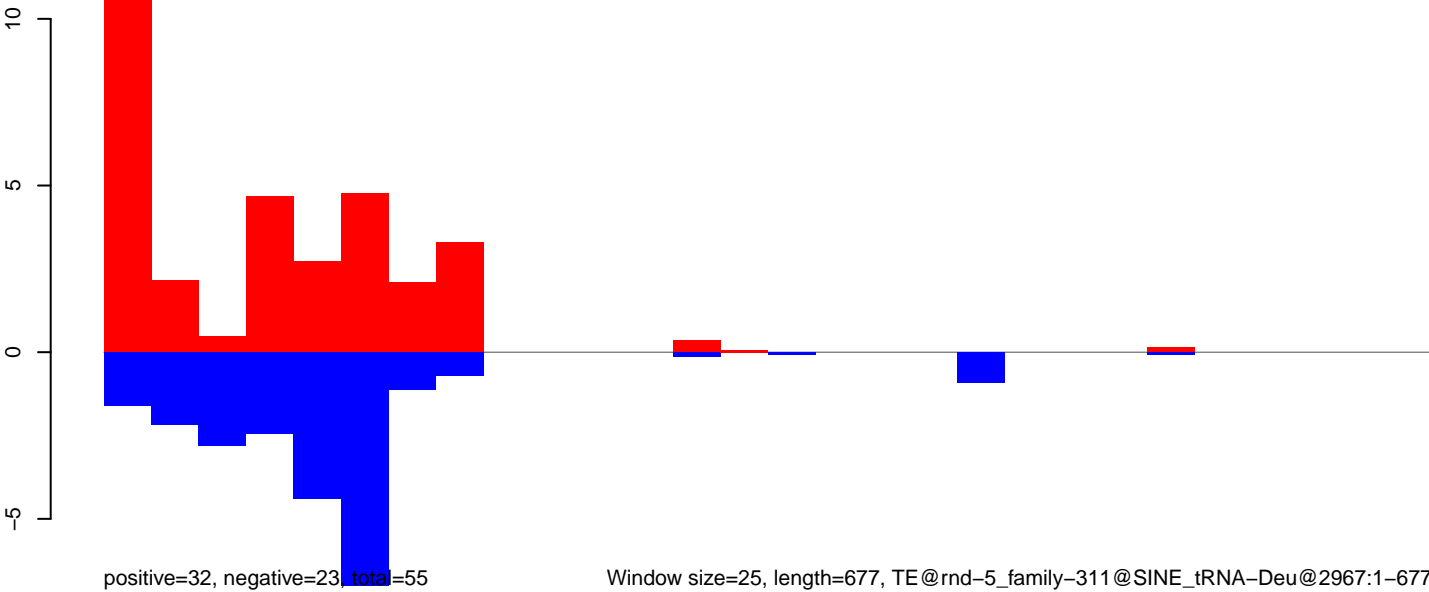
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

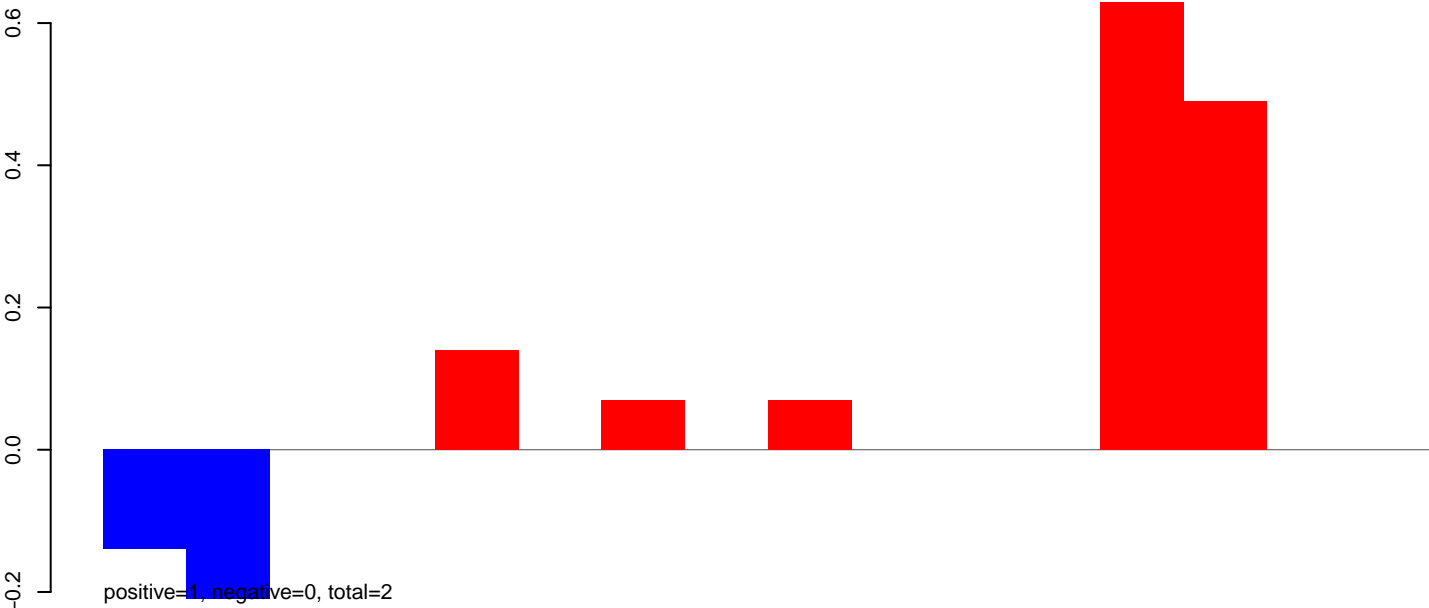


CuQuin_Mcarc_NL.rep

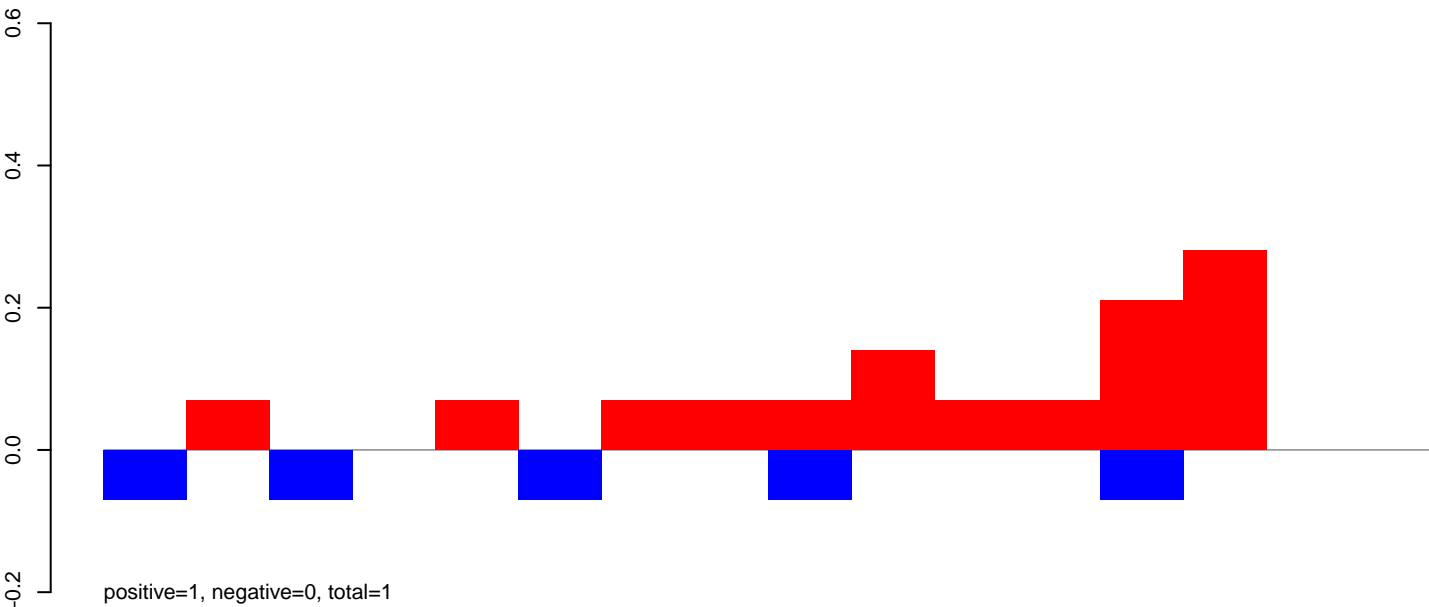


Window size=25, length=677, TE@rnd-5_family-311@SINE_tRNA-Deu@2967:1-677

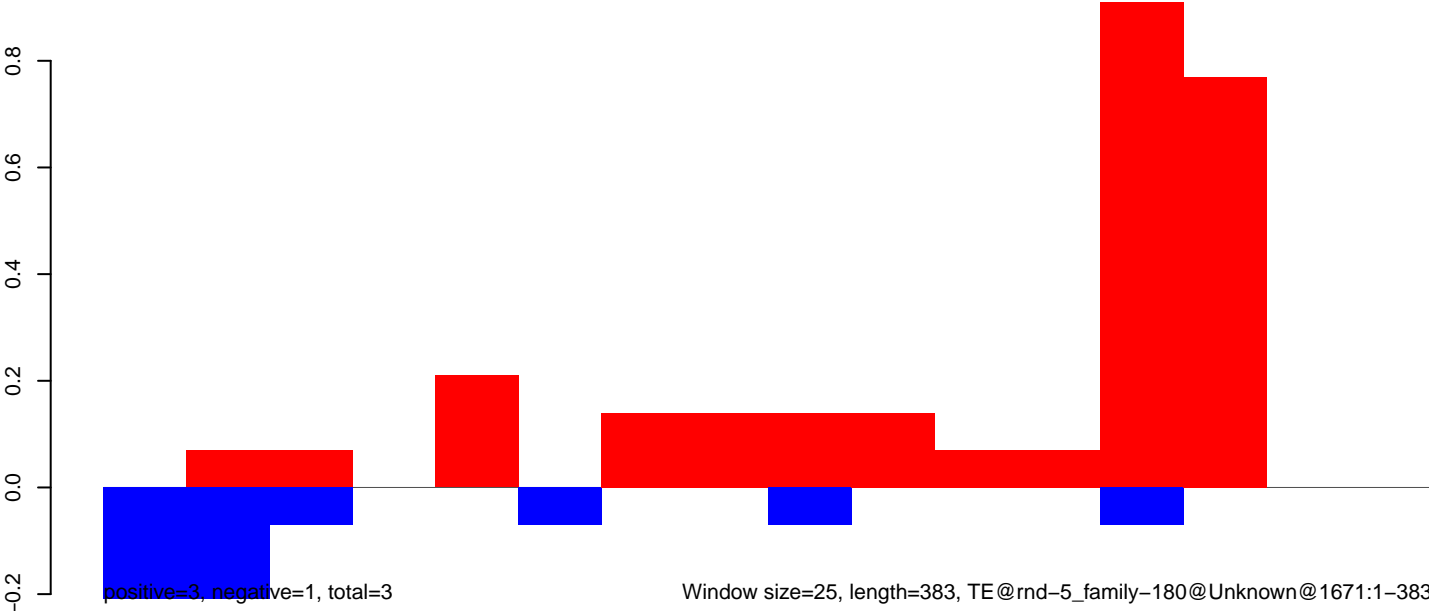
CuQuin_Mcarc_NL.18_23.rep



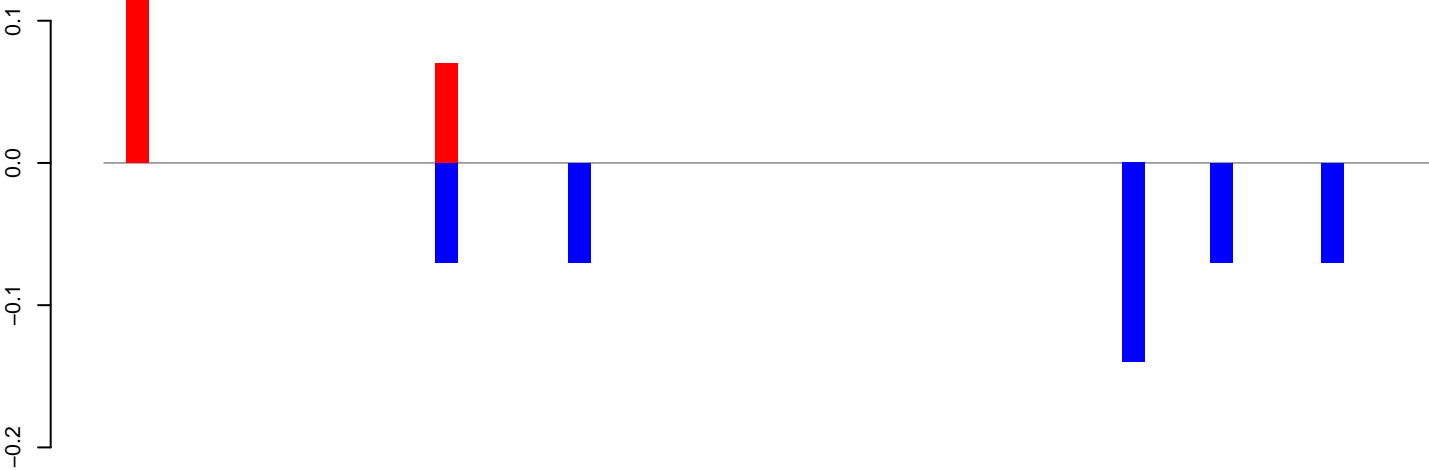
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

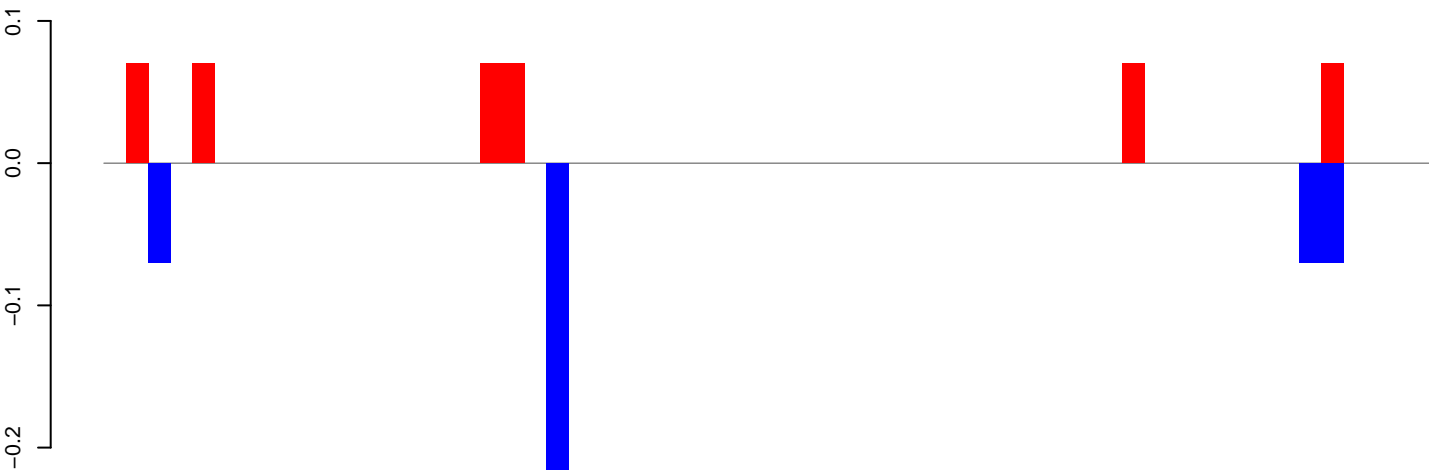


CuQuin_Mcarc_NL.18_23.rep



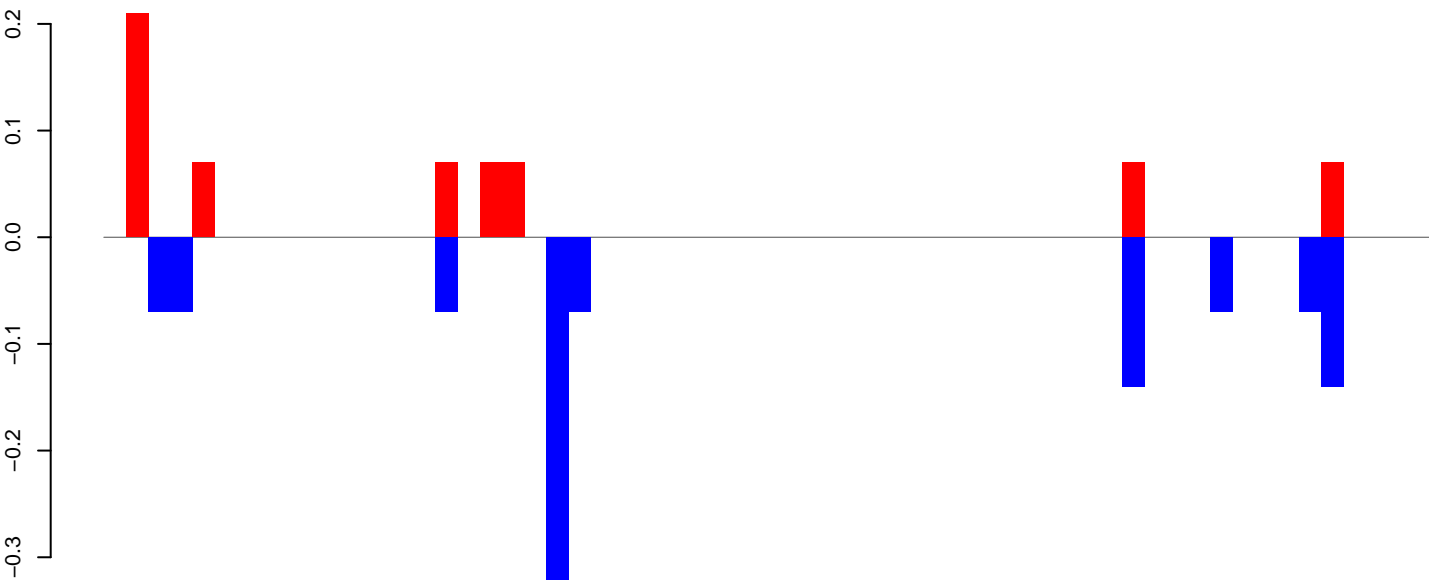
positive=0, negative=0, total=1

CuQuin_Mcarc_NL.24_35.rep



positive=0, negative=0, total=1

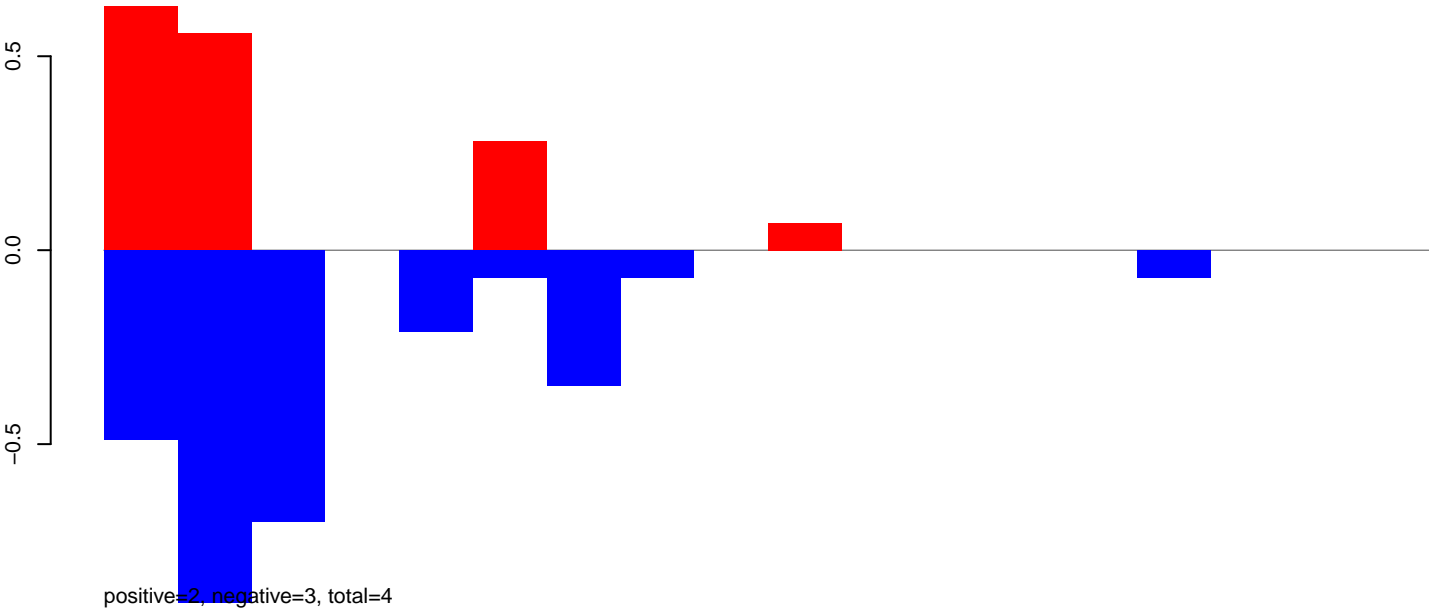
CuQuin_Mcarc_NL.rep



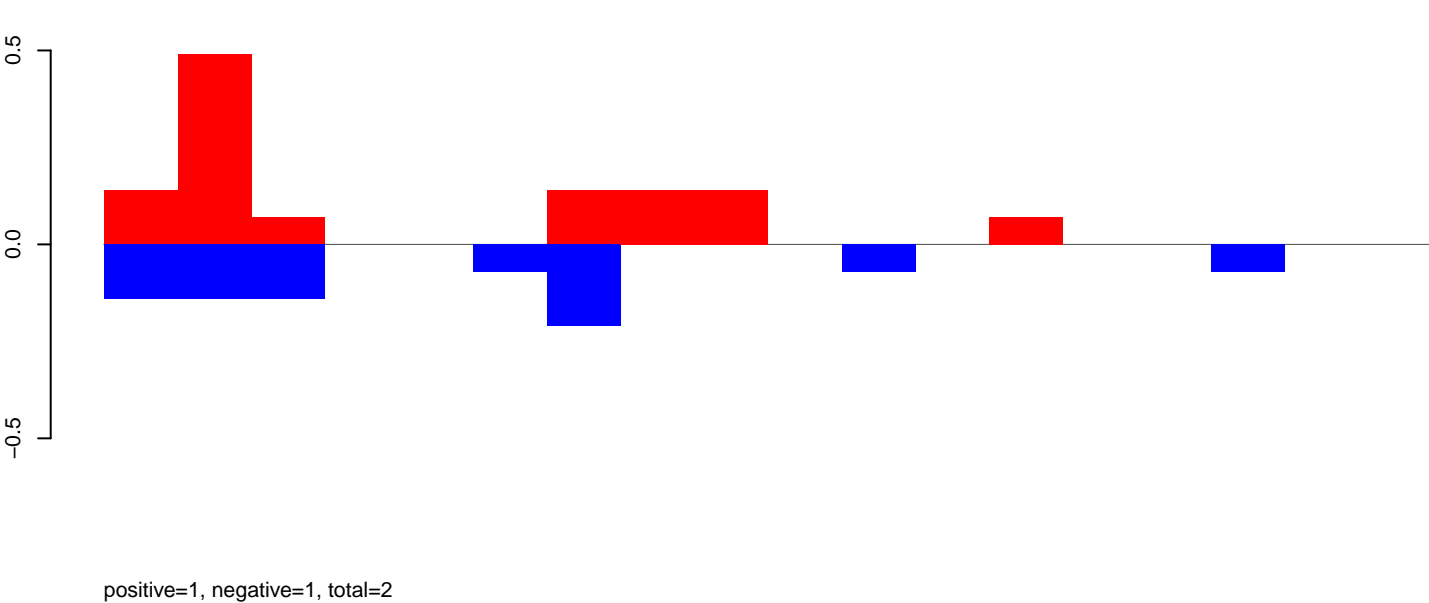
positive=1, negative=1, total=2

Window size=25, length=1479, TE@rnd-5_family-1224@Unknown@3210:1-1479

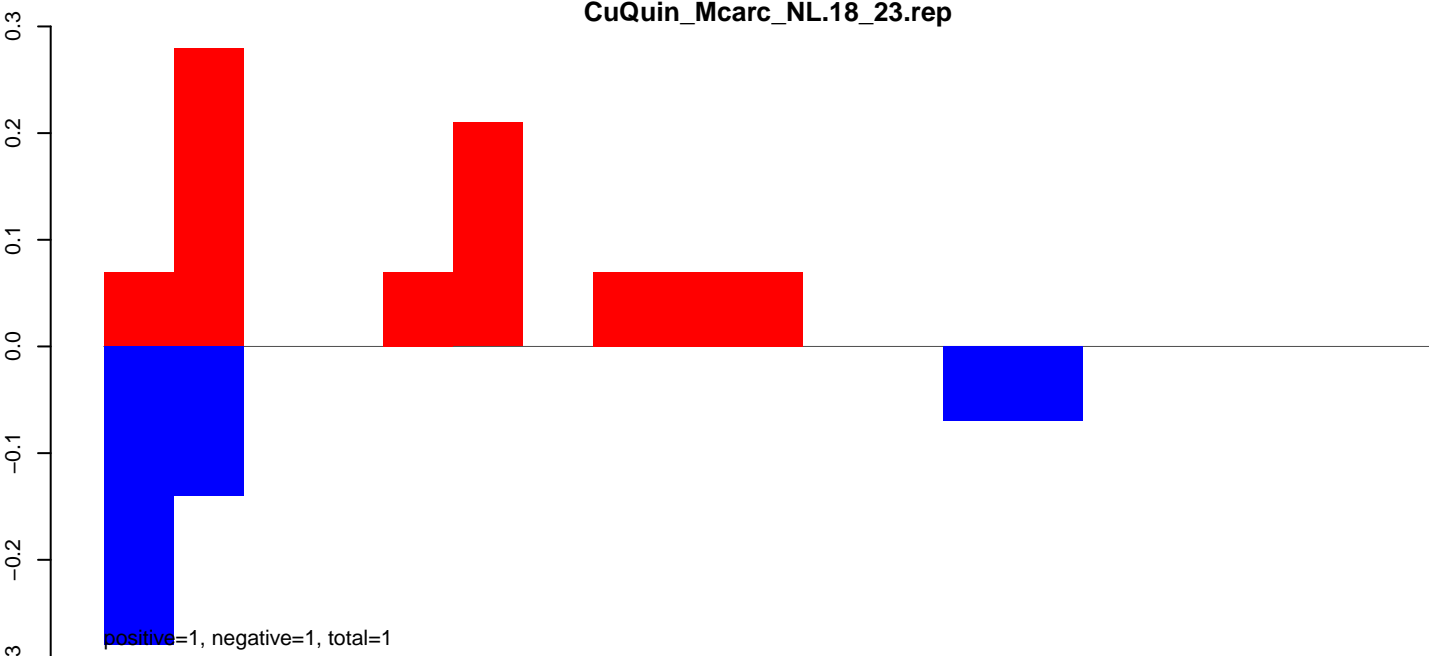
CuQuin_Mcarc_NL.18_23.rep



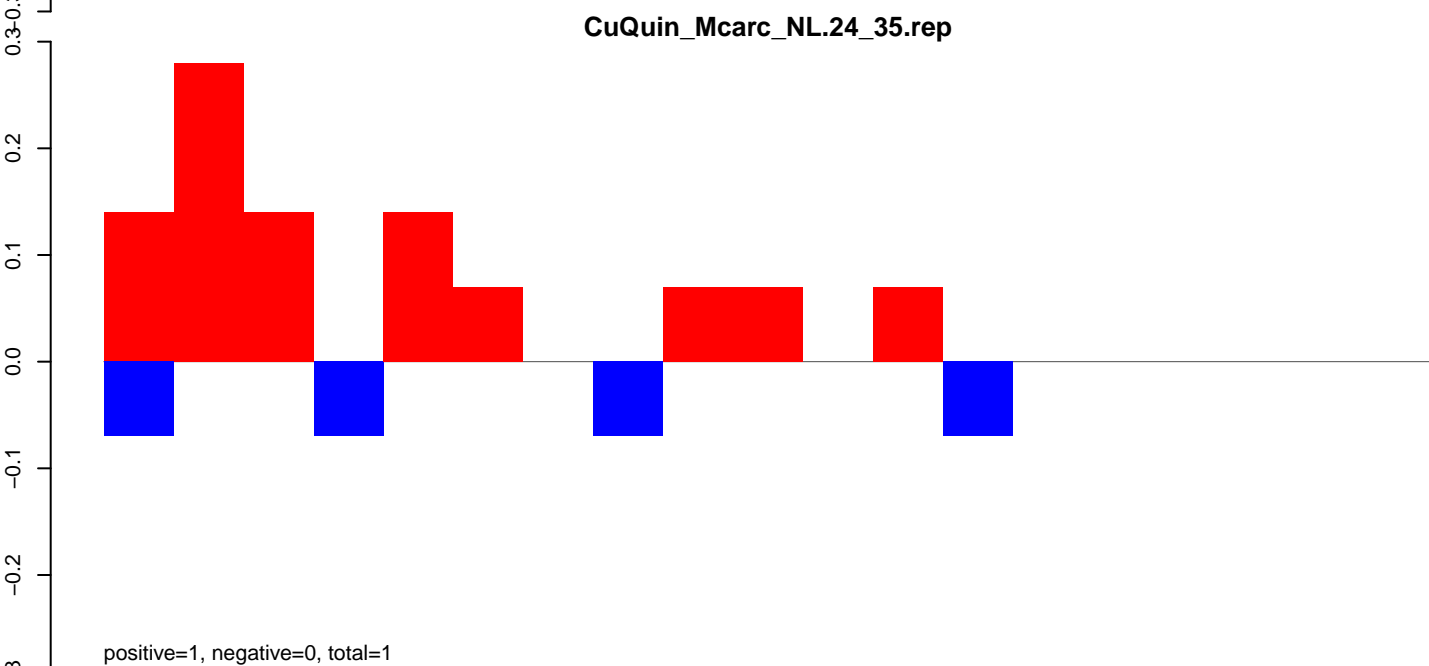
CuQuin_Mcarc_NL.24_35.rep



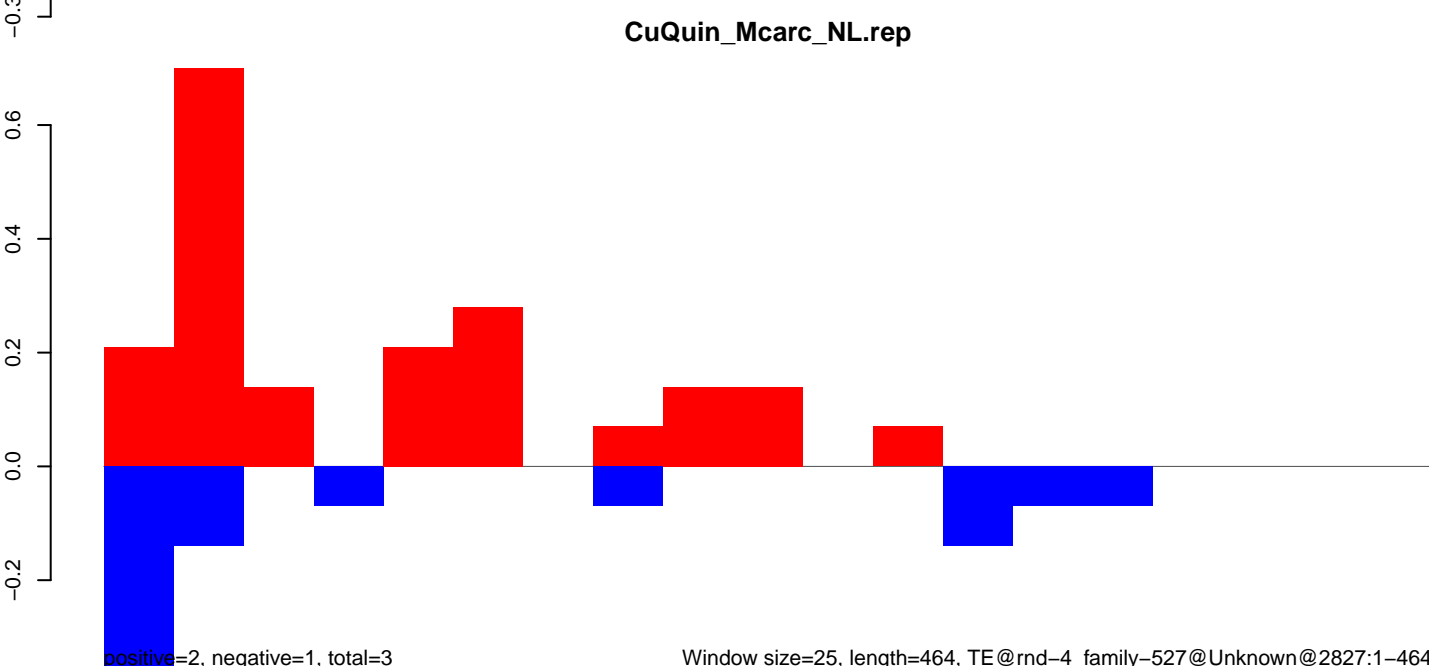
CuQuin_Mcarc_NL.18_23.rep



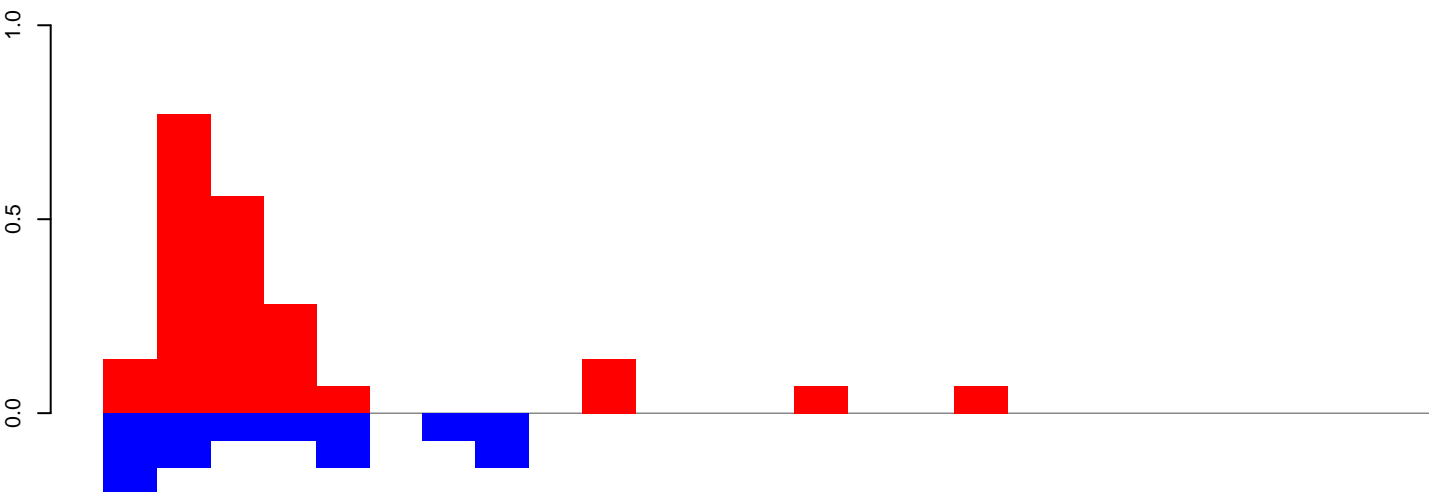
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

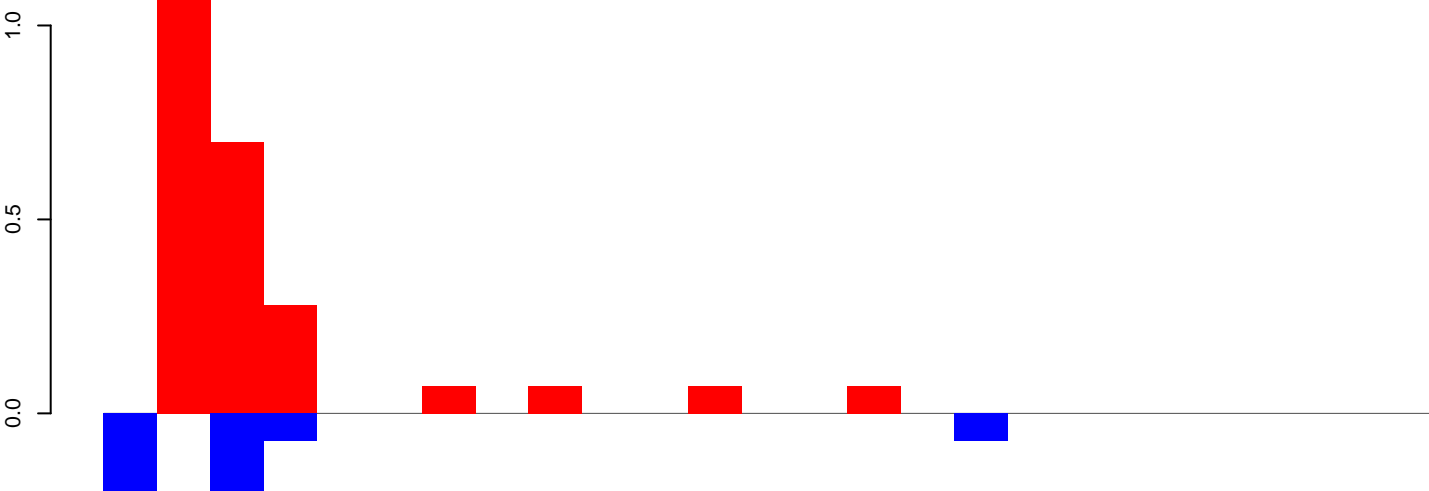


CuQuin_Mcarc_NL.18_23.rep



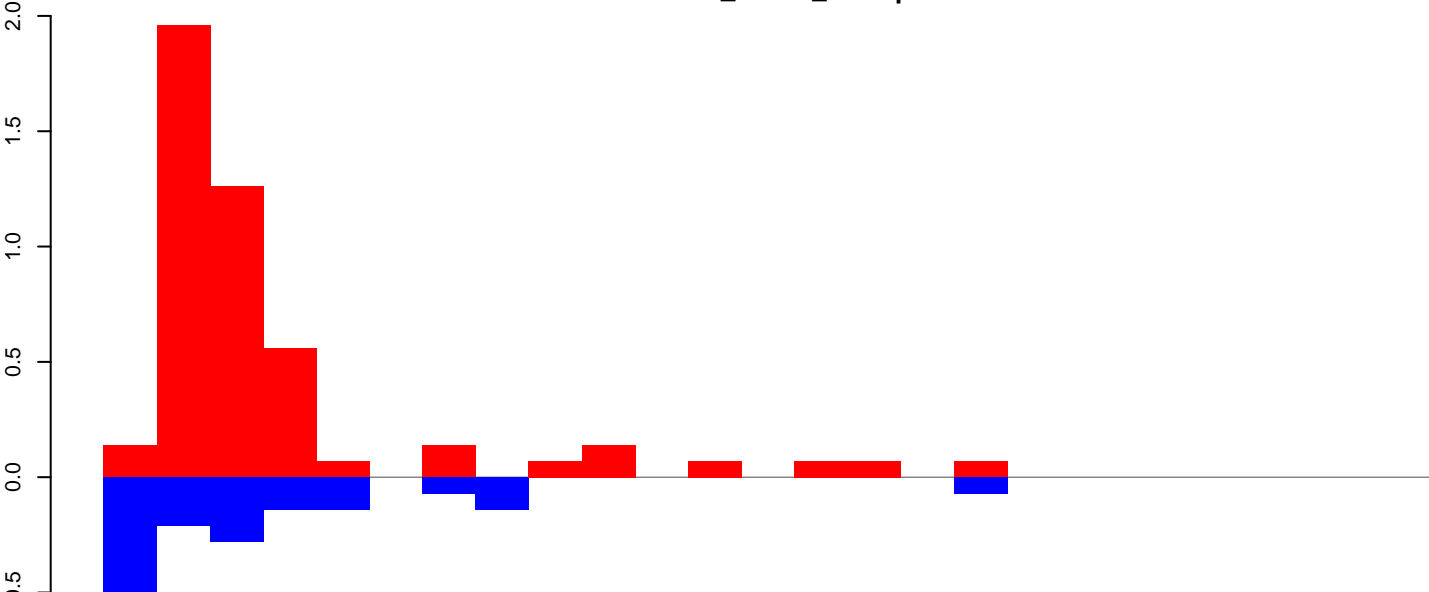
positive=2, negative=1, total=3

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=1, total=3

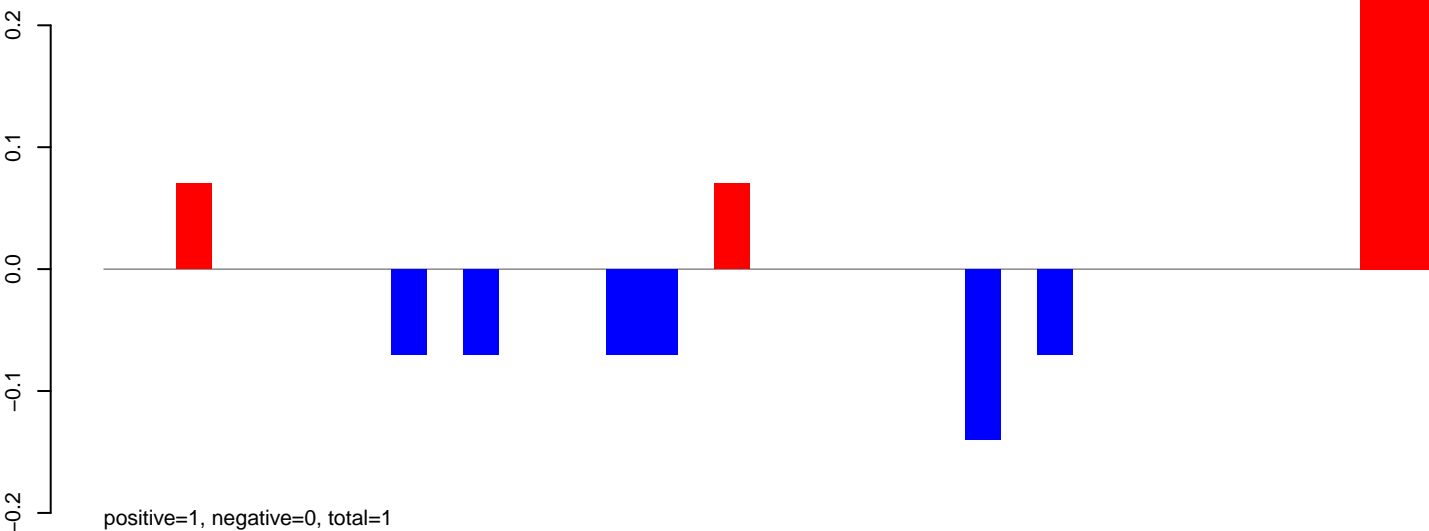
CuQuin_Mcarc_NL.rep



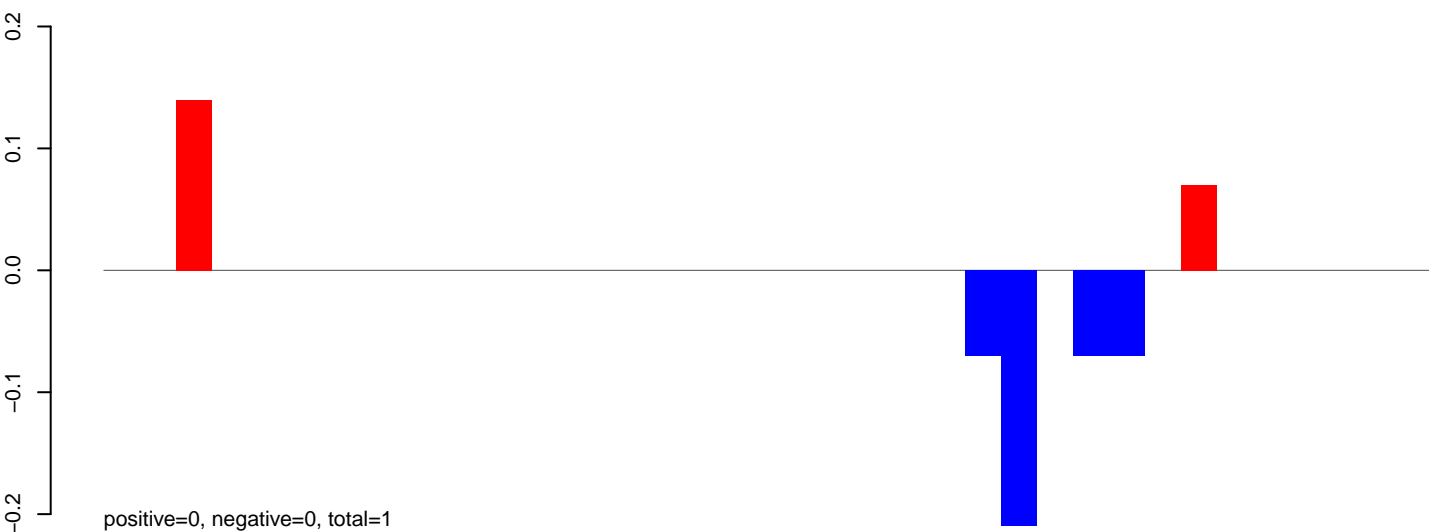
positive=5, negative=2, total=6

Window size=25, length=608, TE@rnd-4_family-203@Satellite@3385:1-608

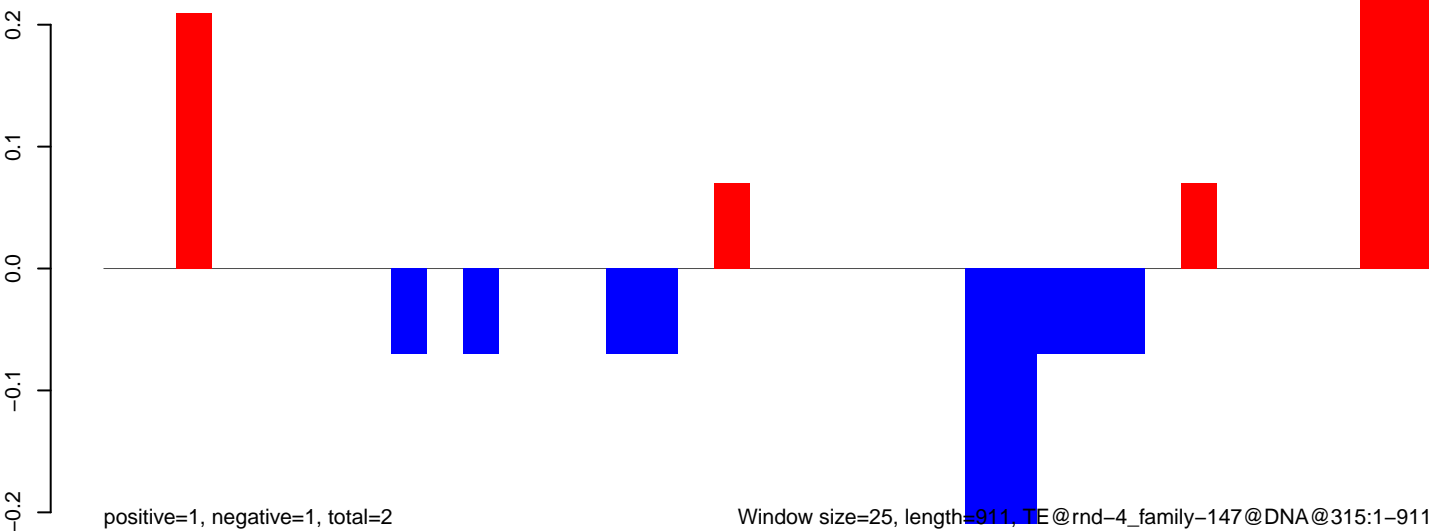
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

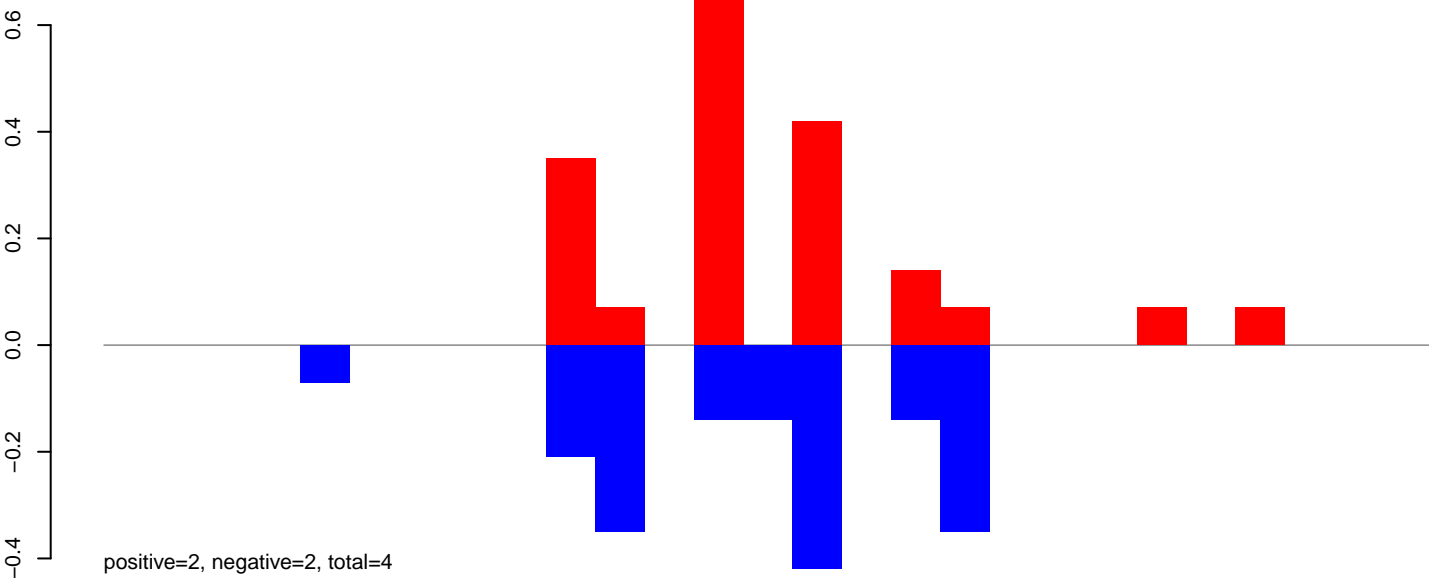


CuQuin_Mcarc_NL.rep

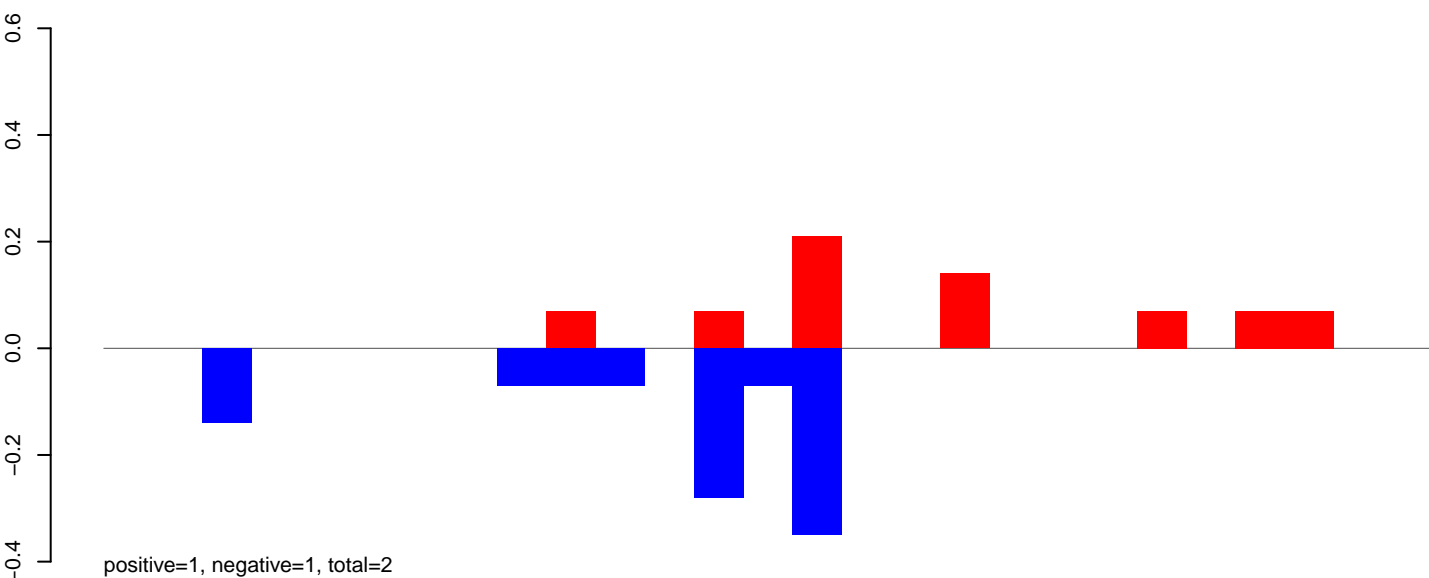


Window size=25, length=911, TE@rnd-4_family-147@DNA@315:1-911

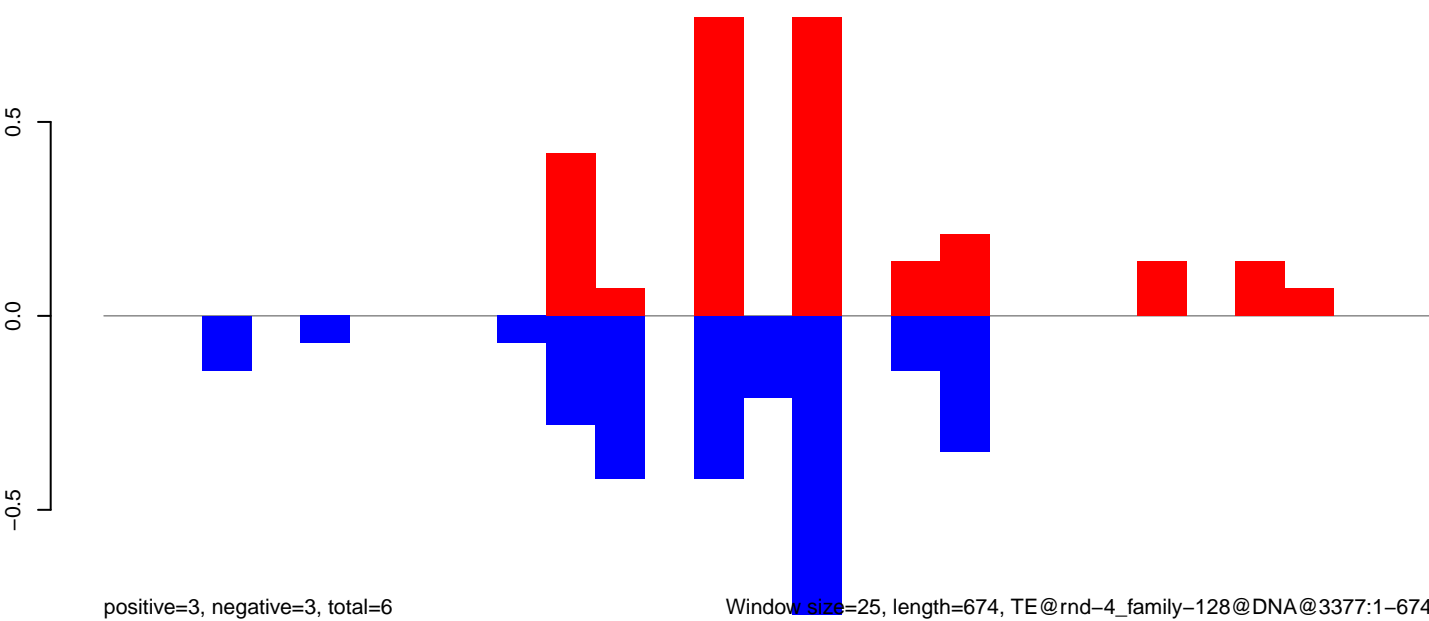
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



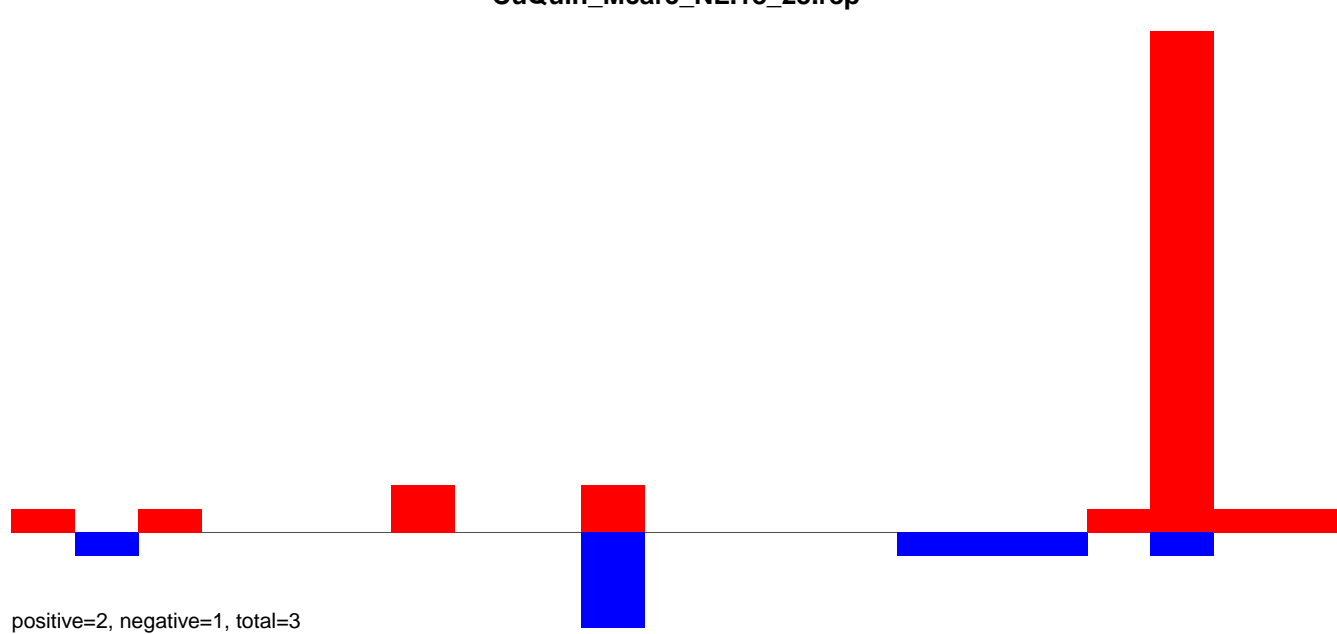
CuQuin_Mcarc_NL.rep



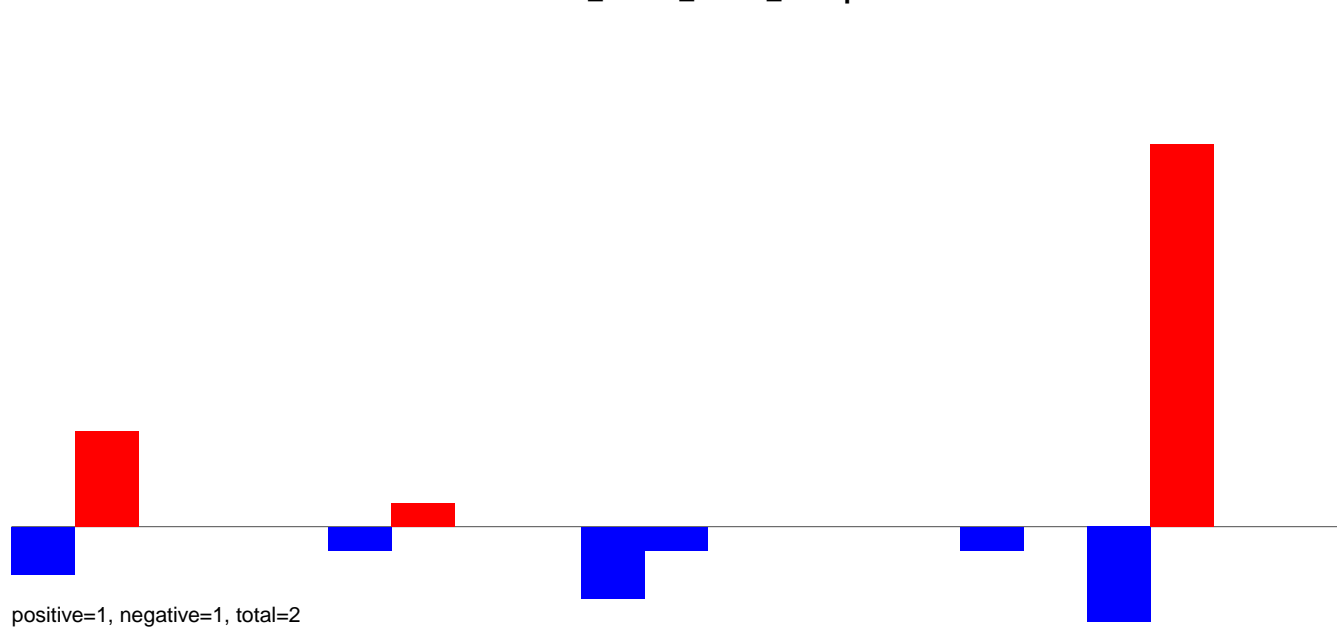
Window size=25, length=674, TE@rnd-4_family-128@DNA@3377:1-674

0 100 200 300 400 500 600 700

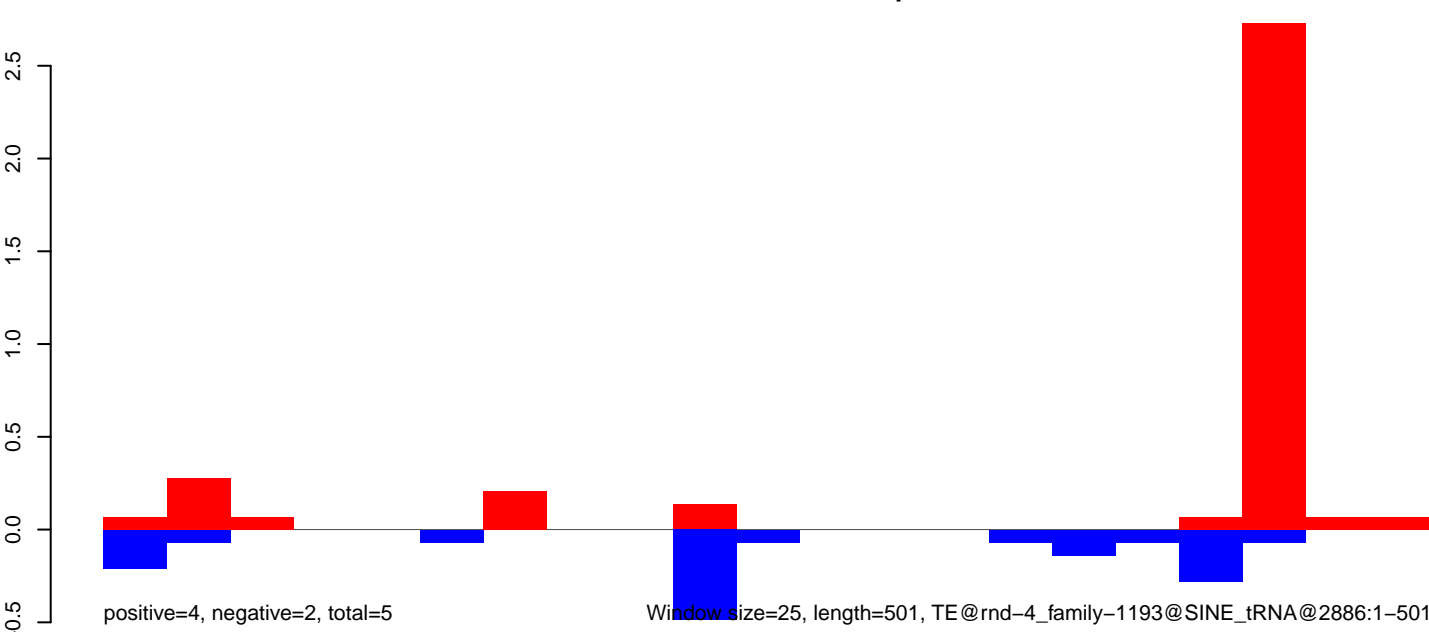
CuQuin_Mcarc_NL.18_23.rep



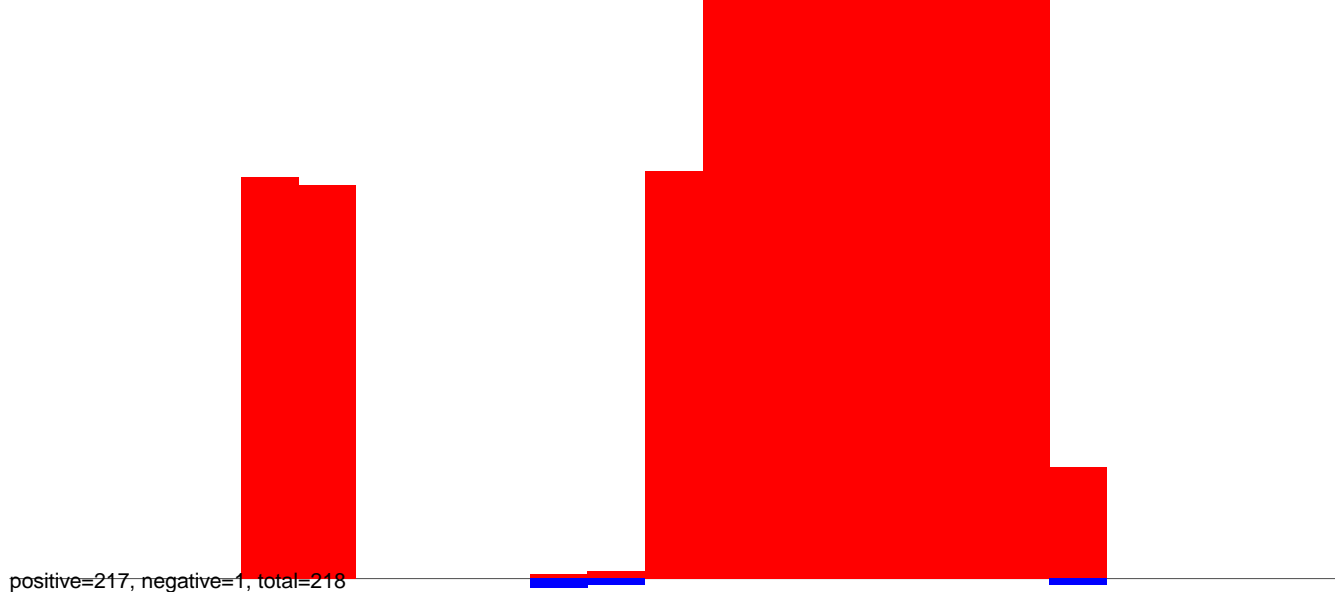
CuQuin_Mcarc_NL.24_35.rep



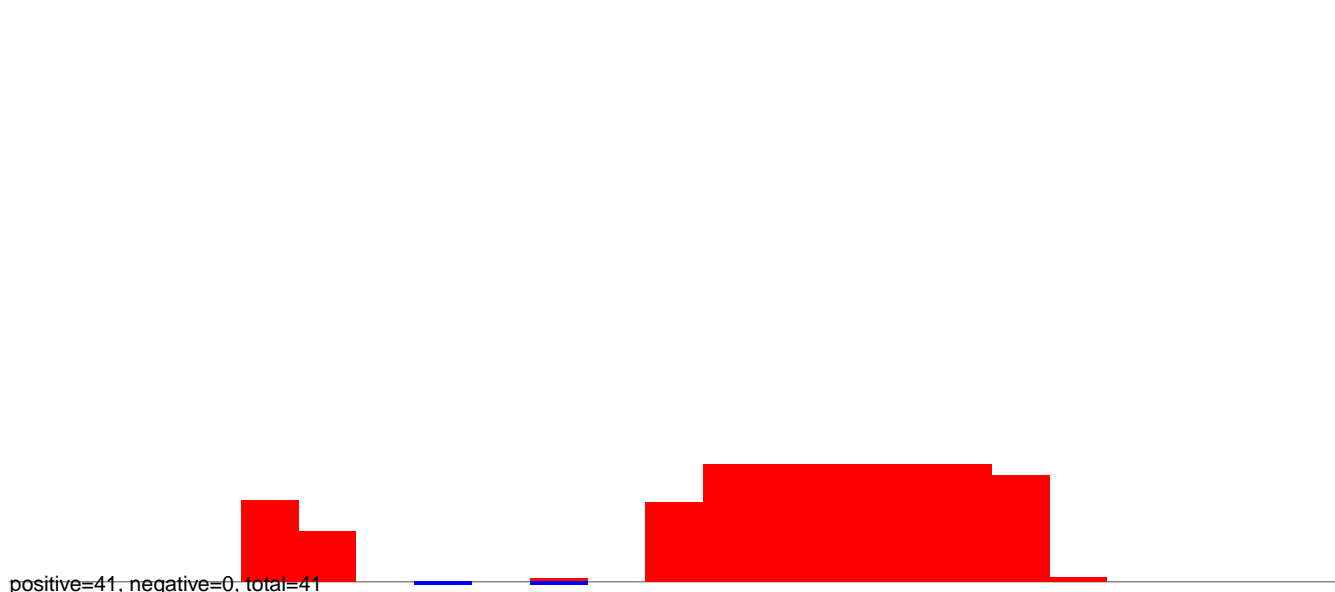
CuQuin_Mcarc_NL.rep



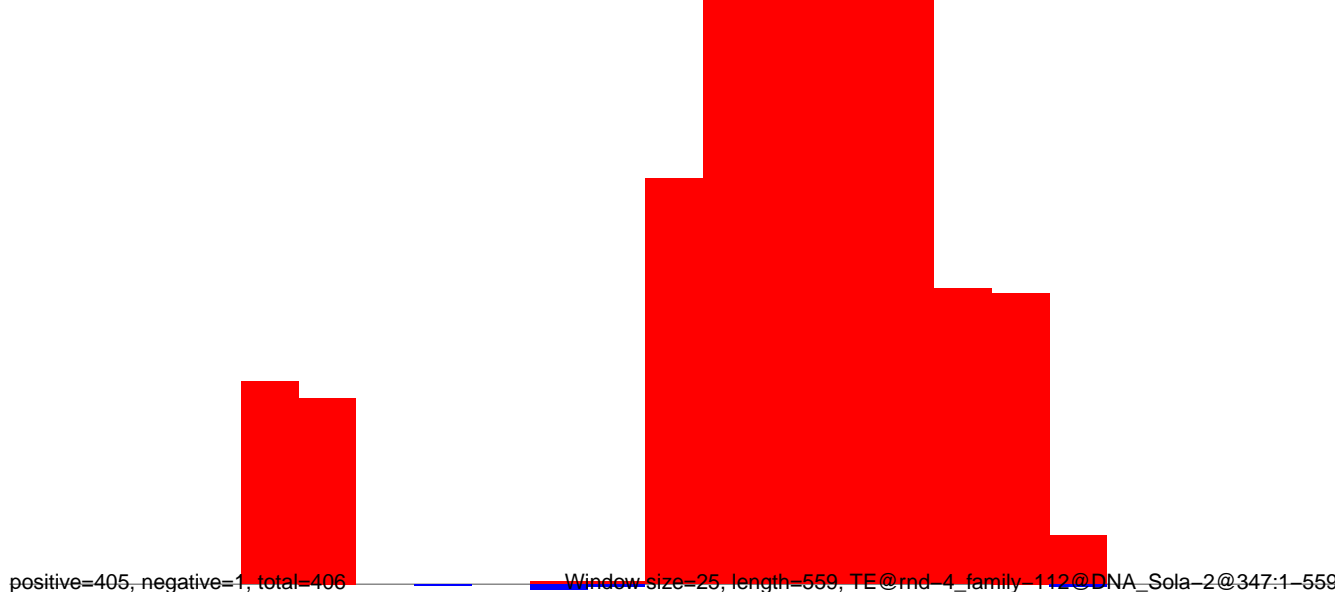
CuQuin_Mcarc_NL.18_23.rep



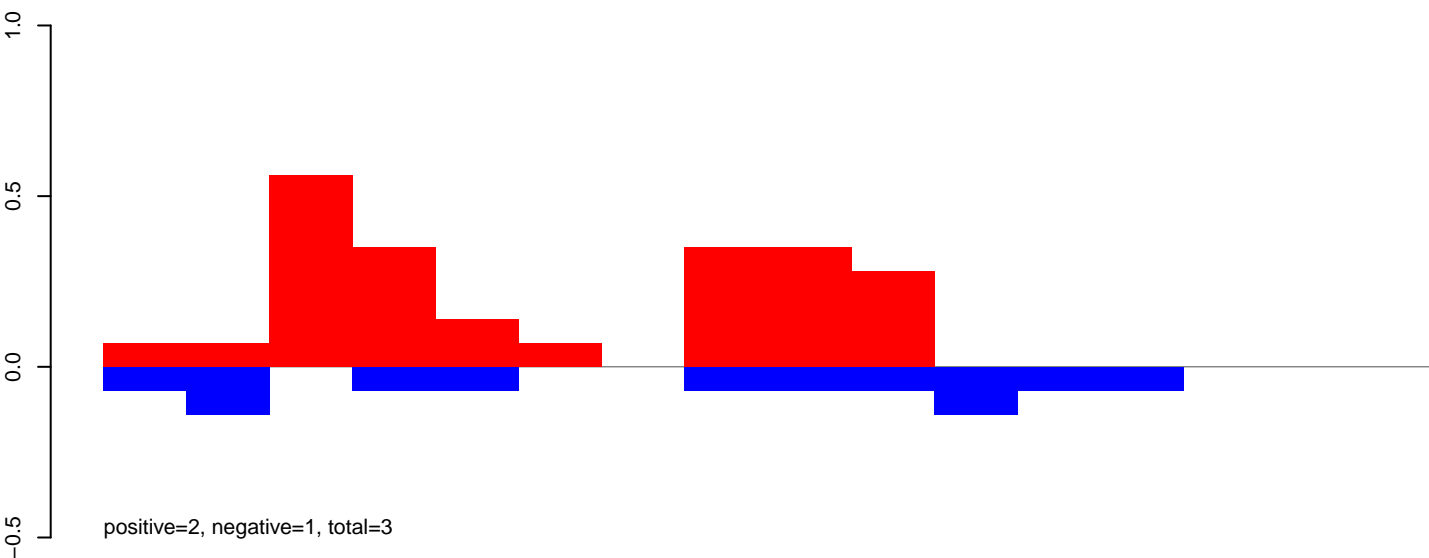
CuQuin_Mcarc_NL.24_35.rep



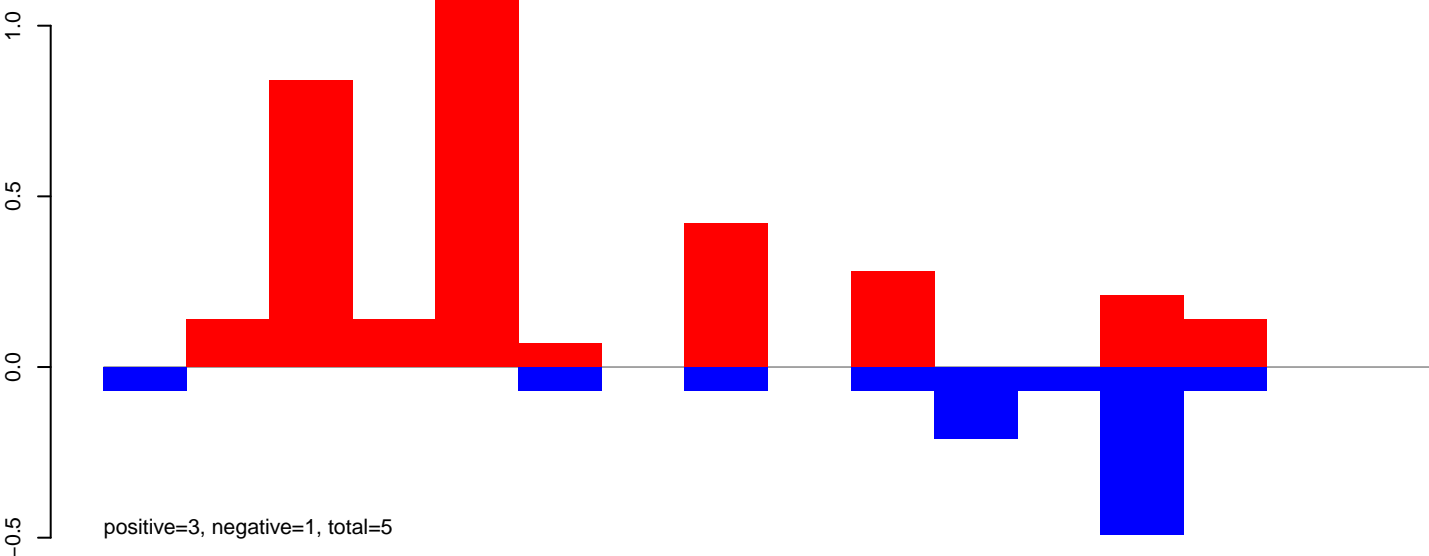
CuQuin_Mcarc_NL.rep



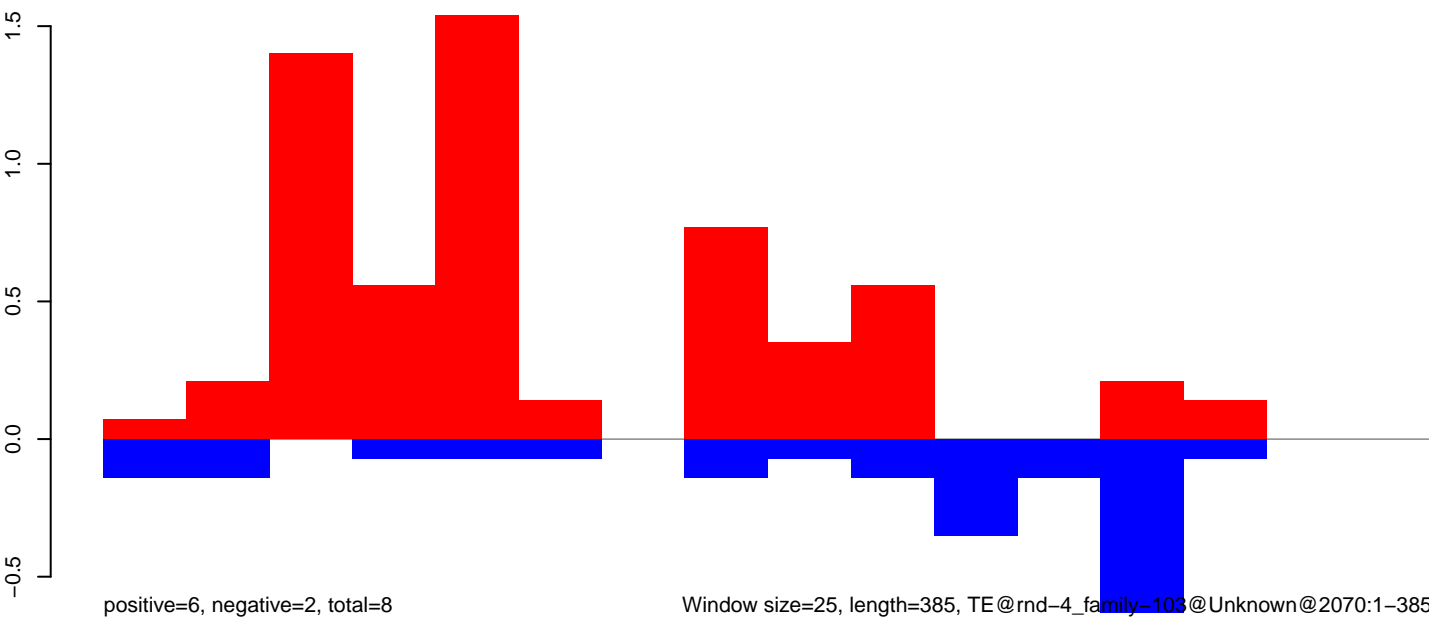
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

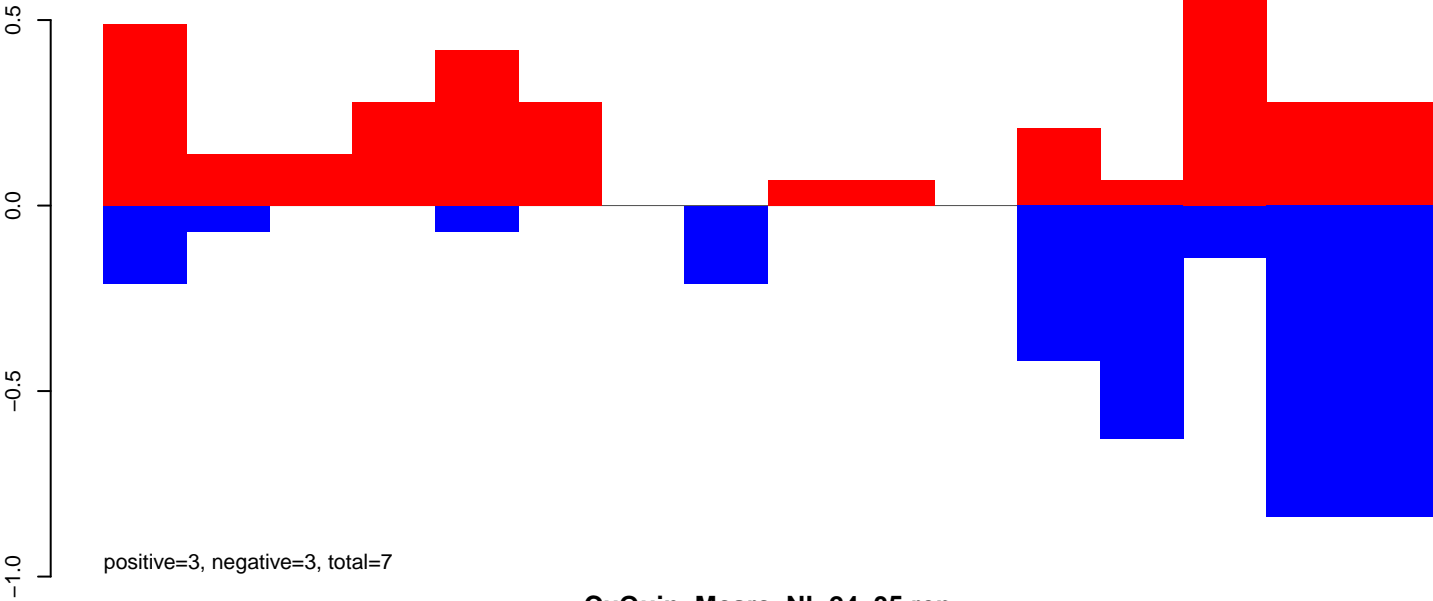


CuQuin_Mcarc_NL.rep

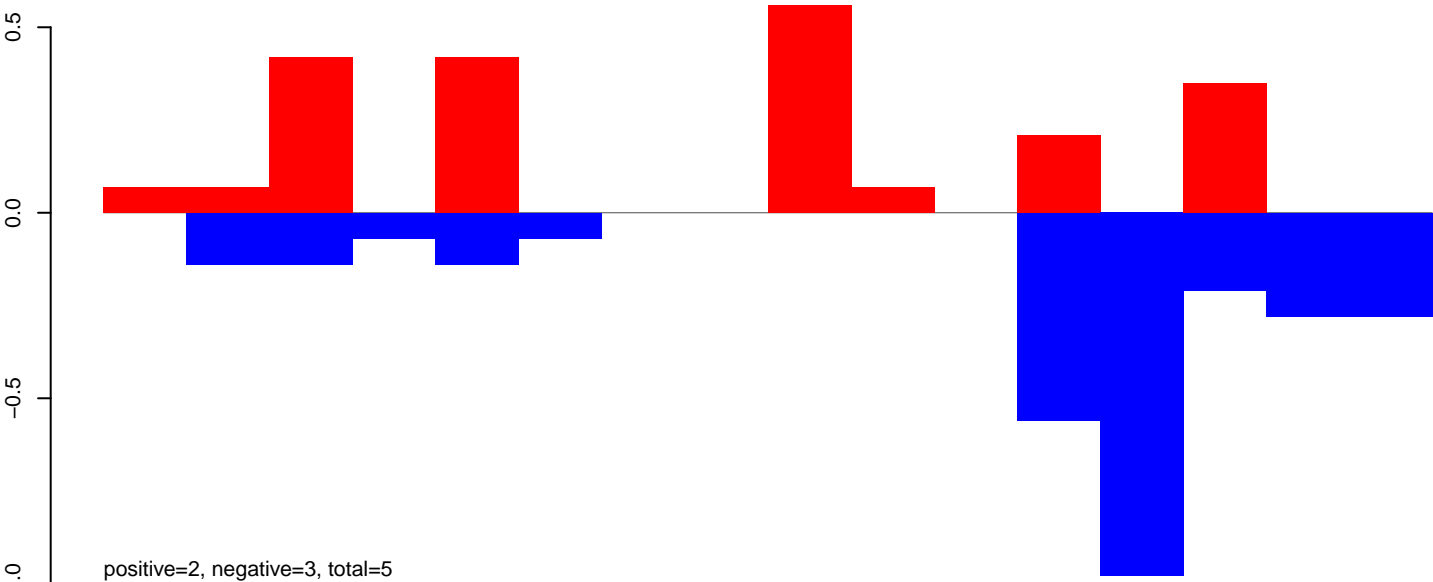


Window size=25, length=385, TE@rnd-4_family-103@Unknown@2070:1-385

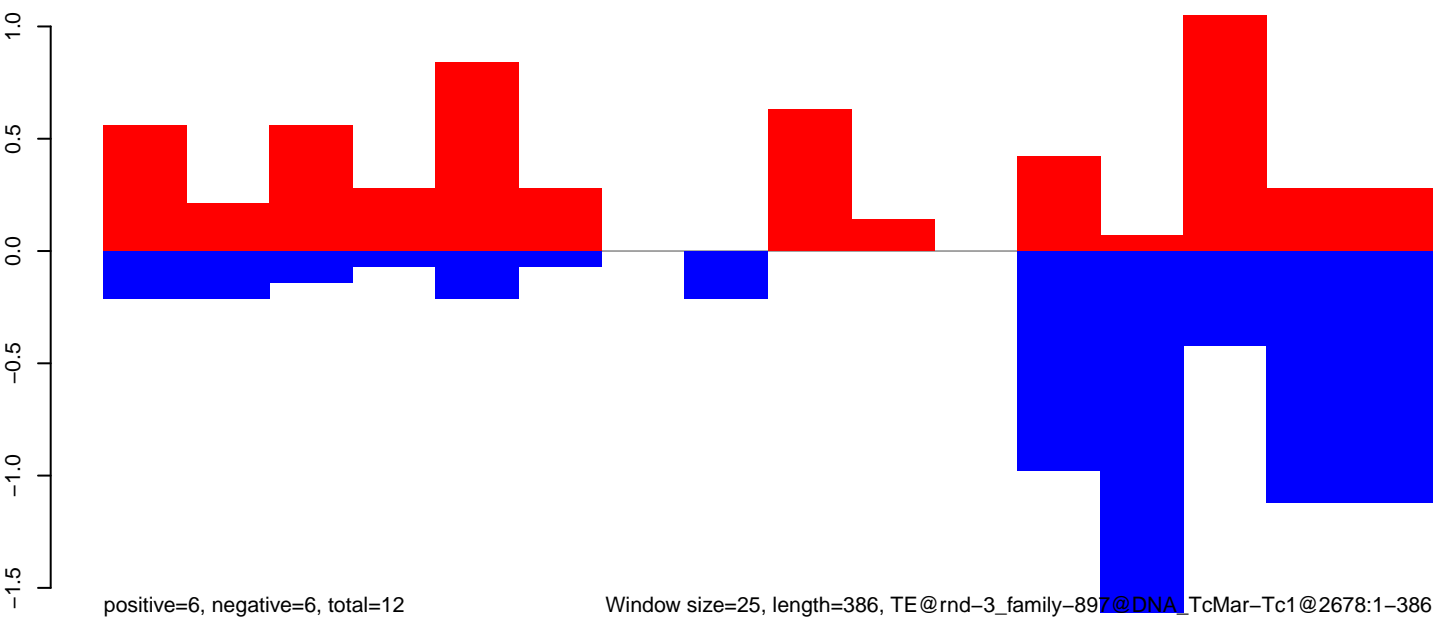
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

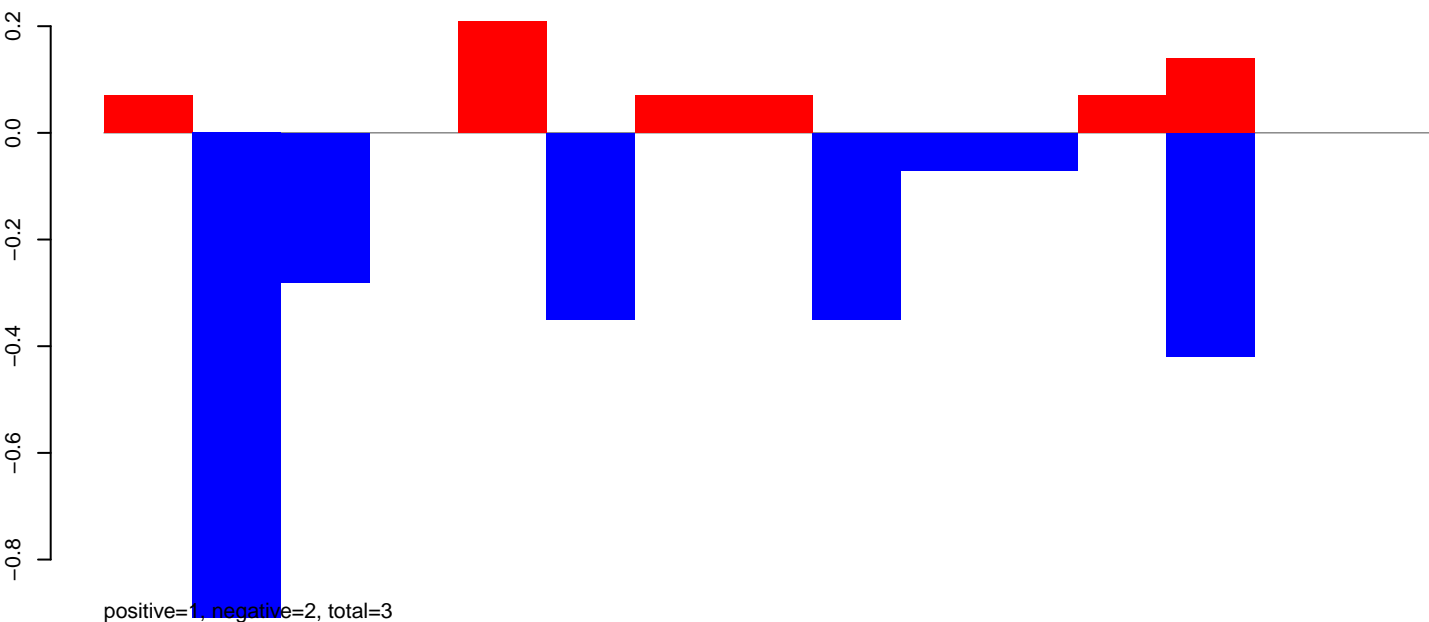


CuQuin_Mcarc_NL.rep

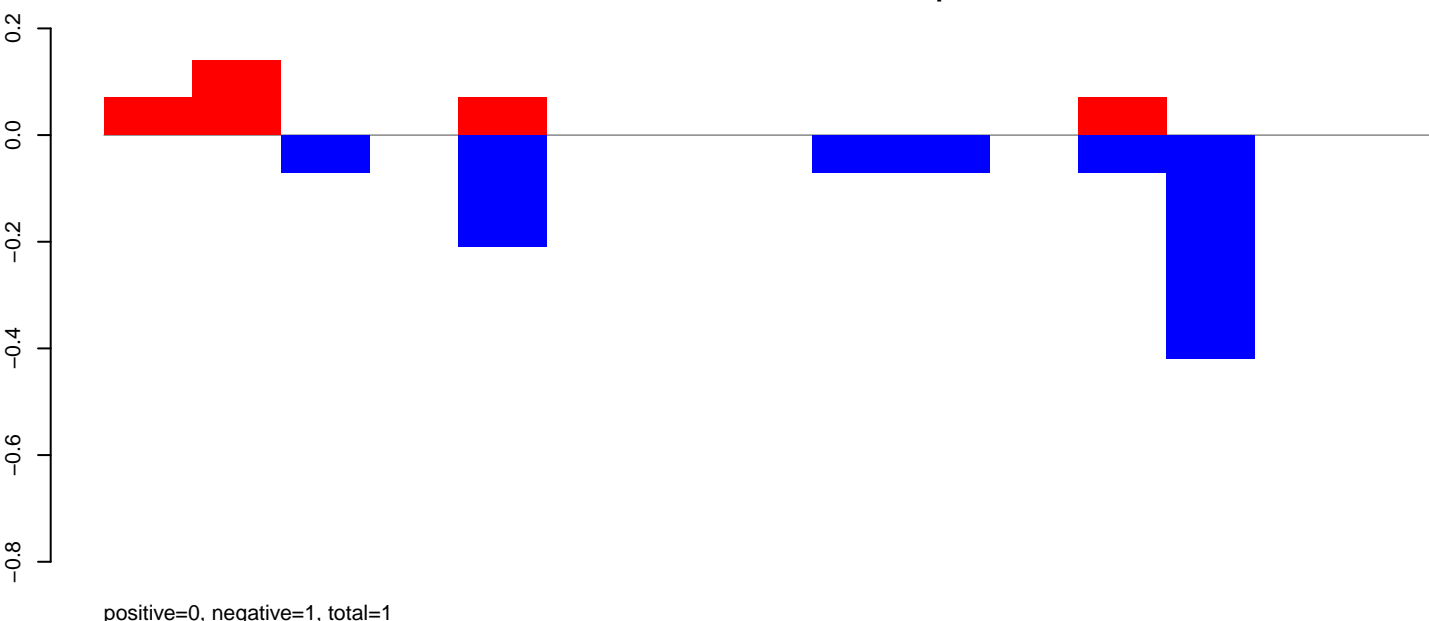


Window size=25, length=386, TE@rnd-3_family-897@DNA_TcMar-Tc1@2678:1-386

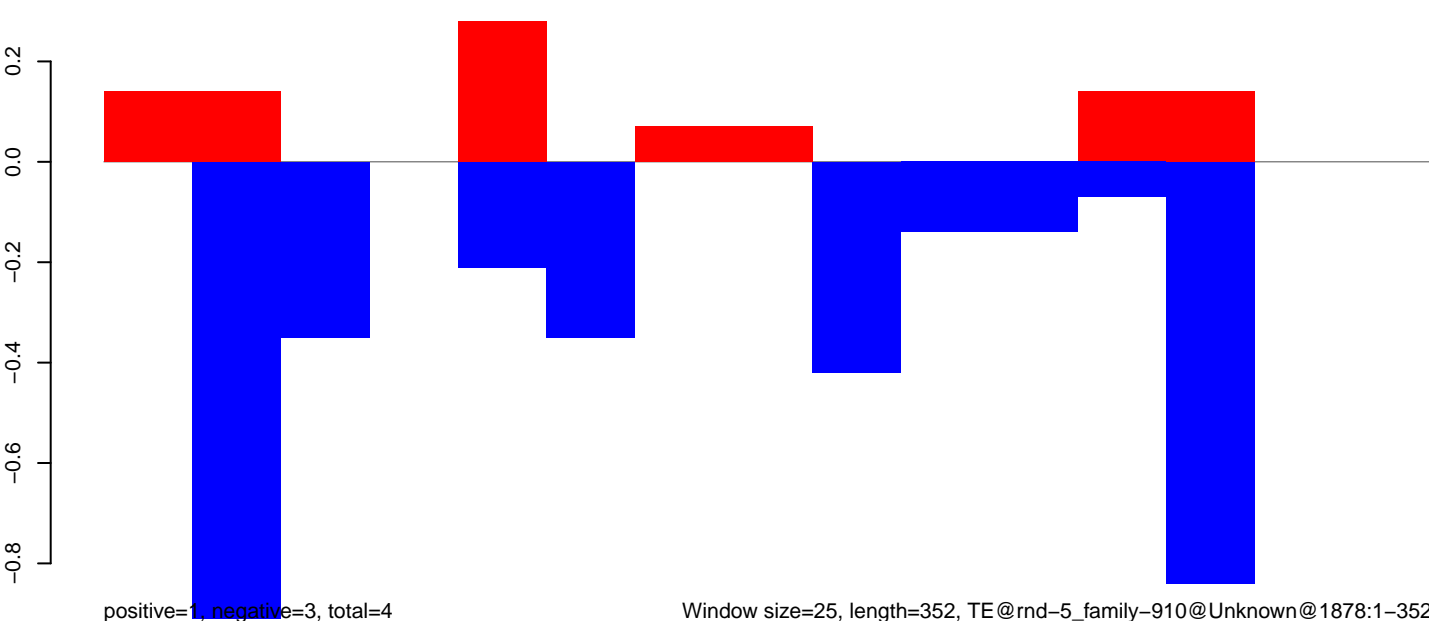
CuQuin_Mcarc_NL.18_23.rep



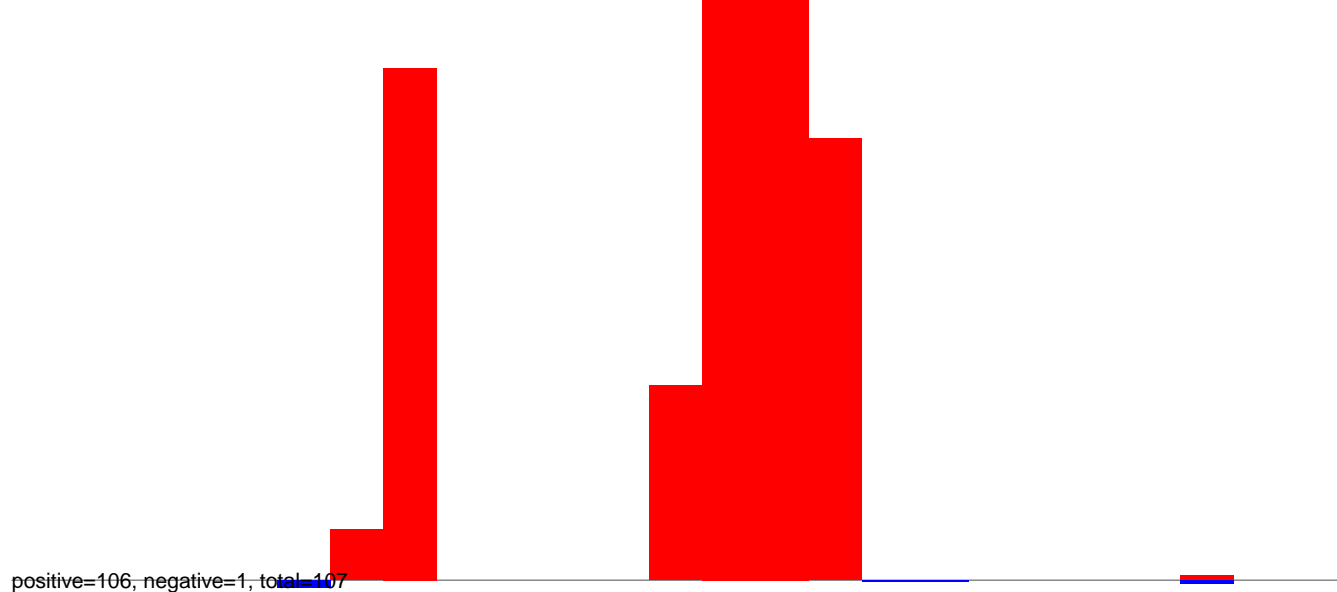
CuQuin_Mcarc_NL.24_35.rep



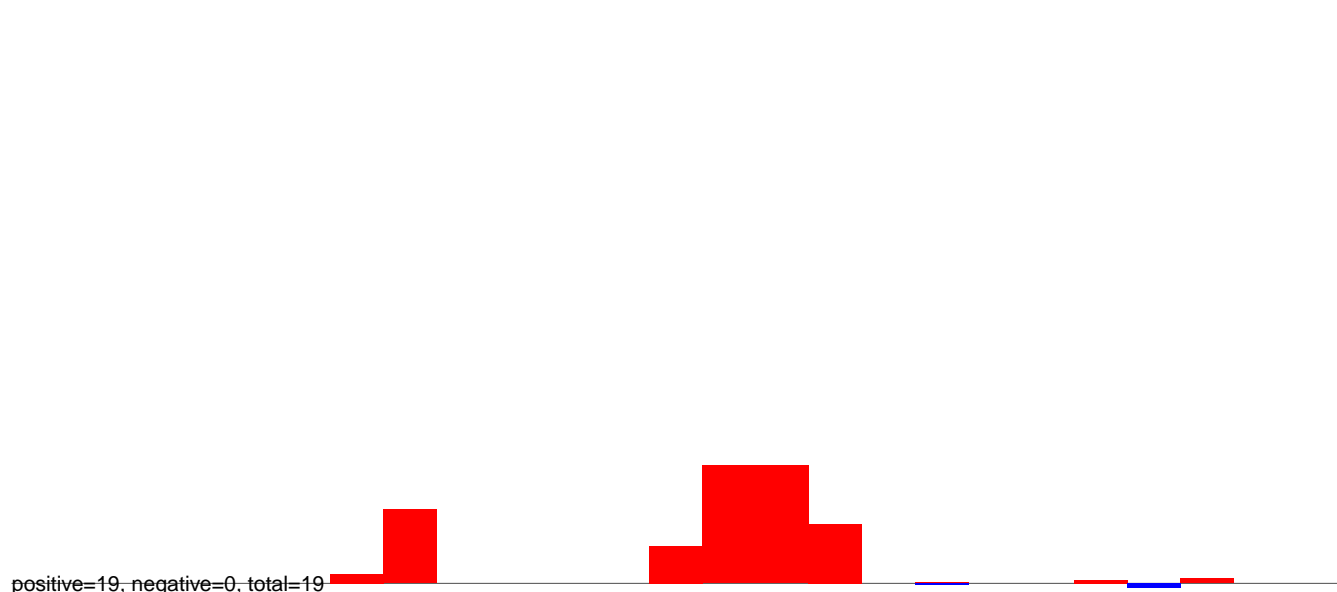
CuQuin_Mcarc_NL.rep



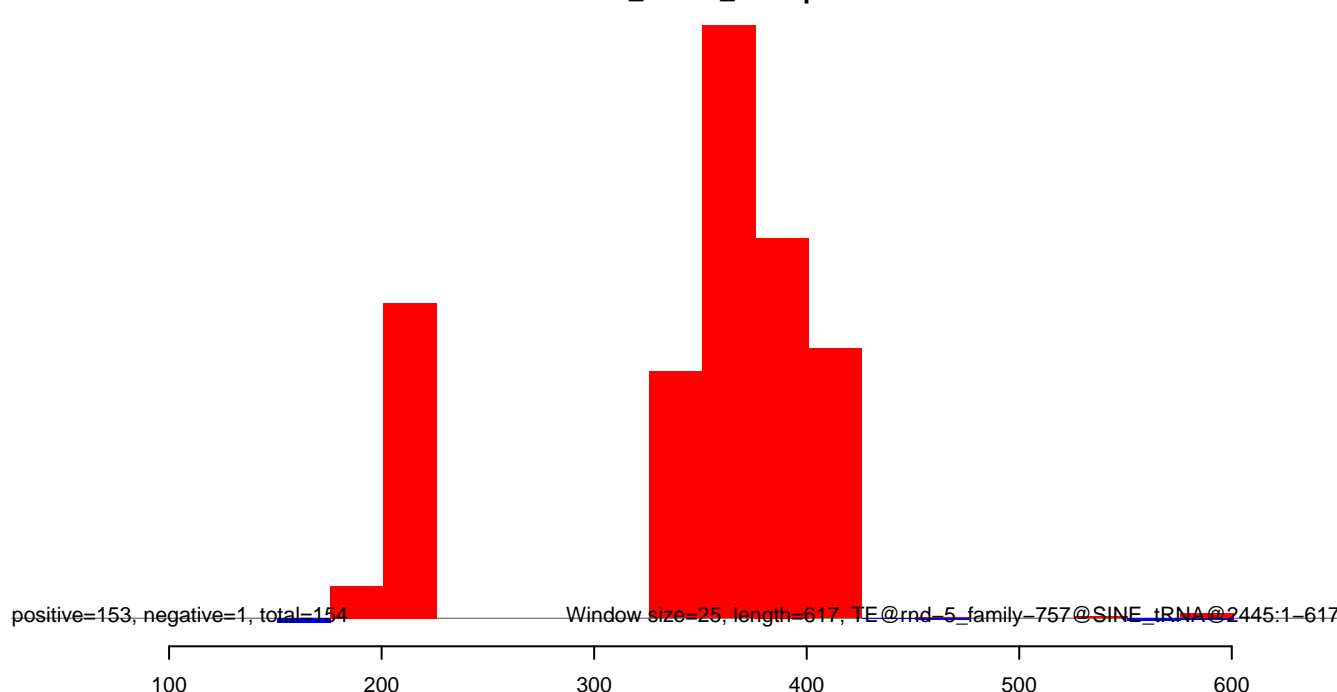
CuQuin_Mcarc_NL.18_23.rep



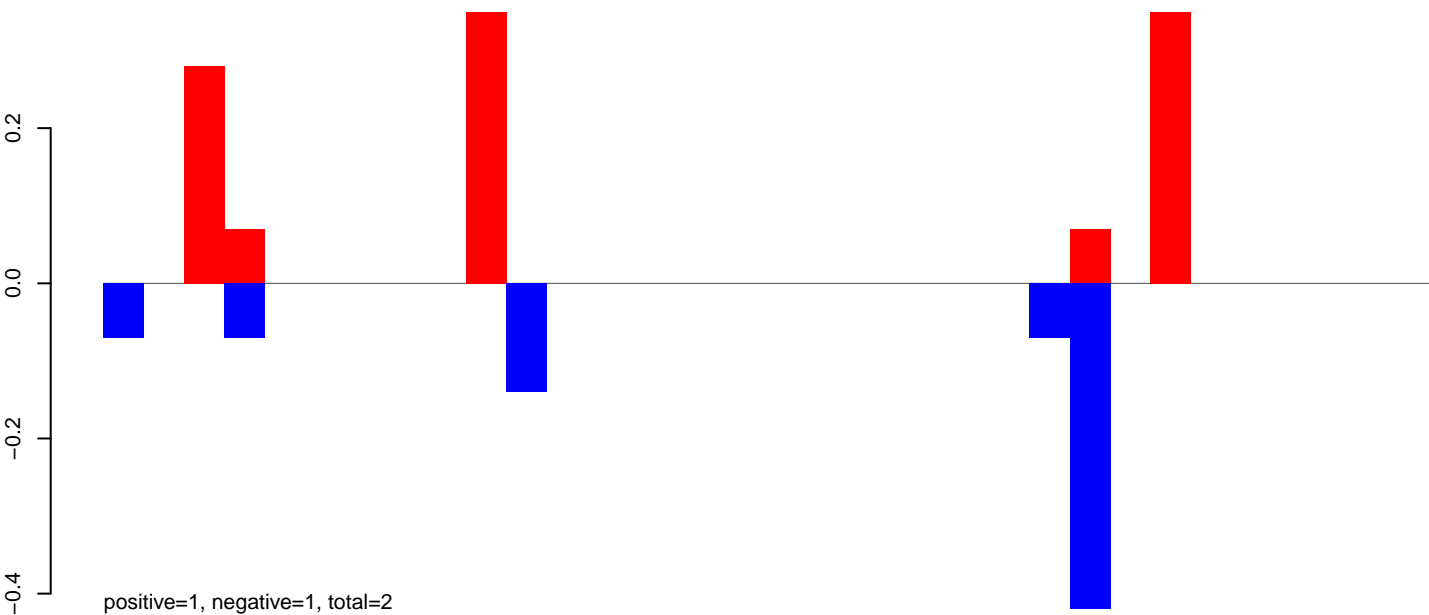
CuQuin_Mcarc_NL.24_35.rep



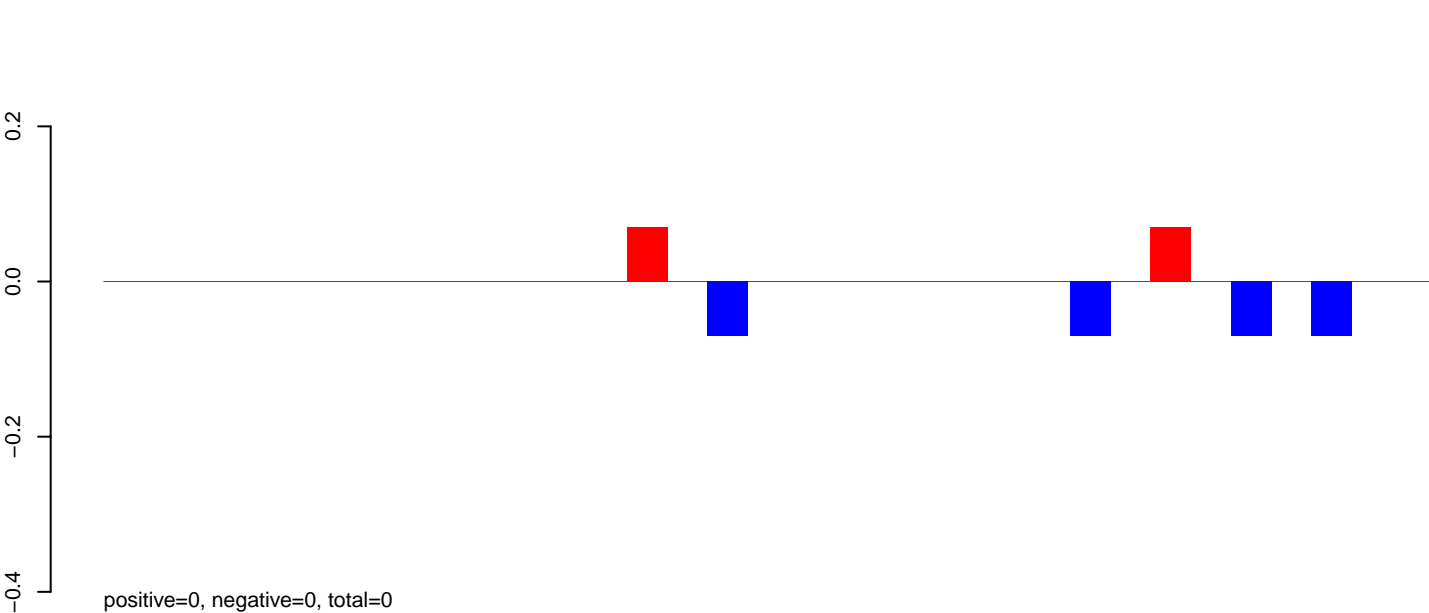
CuQuin_Mcarc_NL.rep



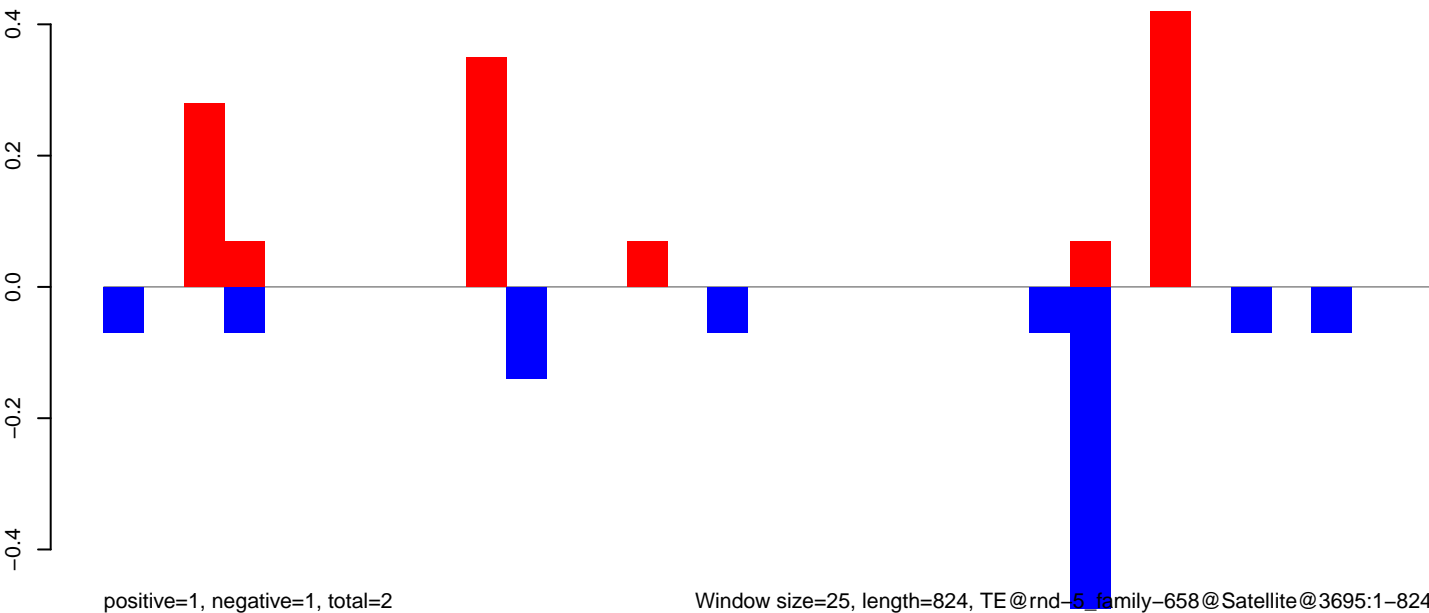
CuQuin_Mcarc_NL.18_23.rep



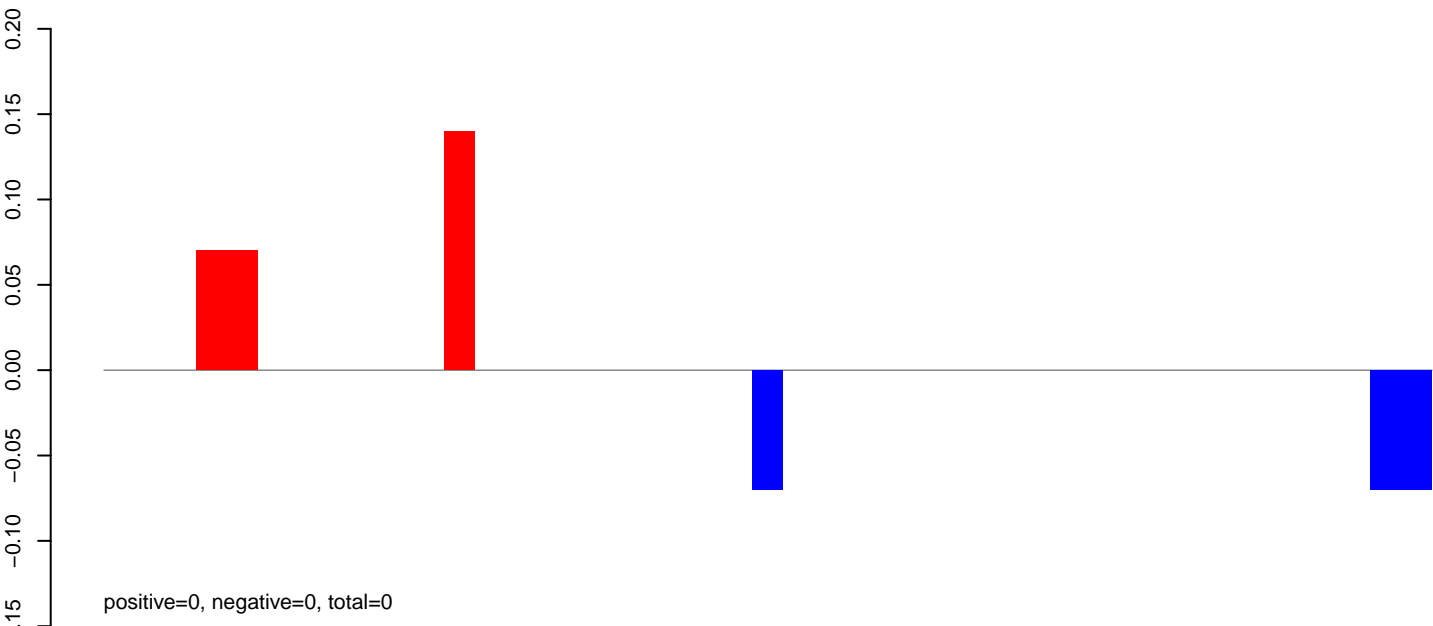
CuQuin_Mcarc_NL.24_35.rep



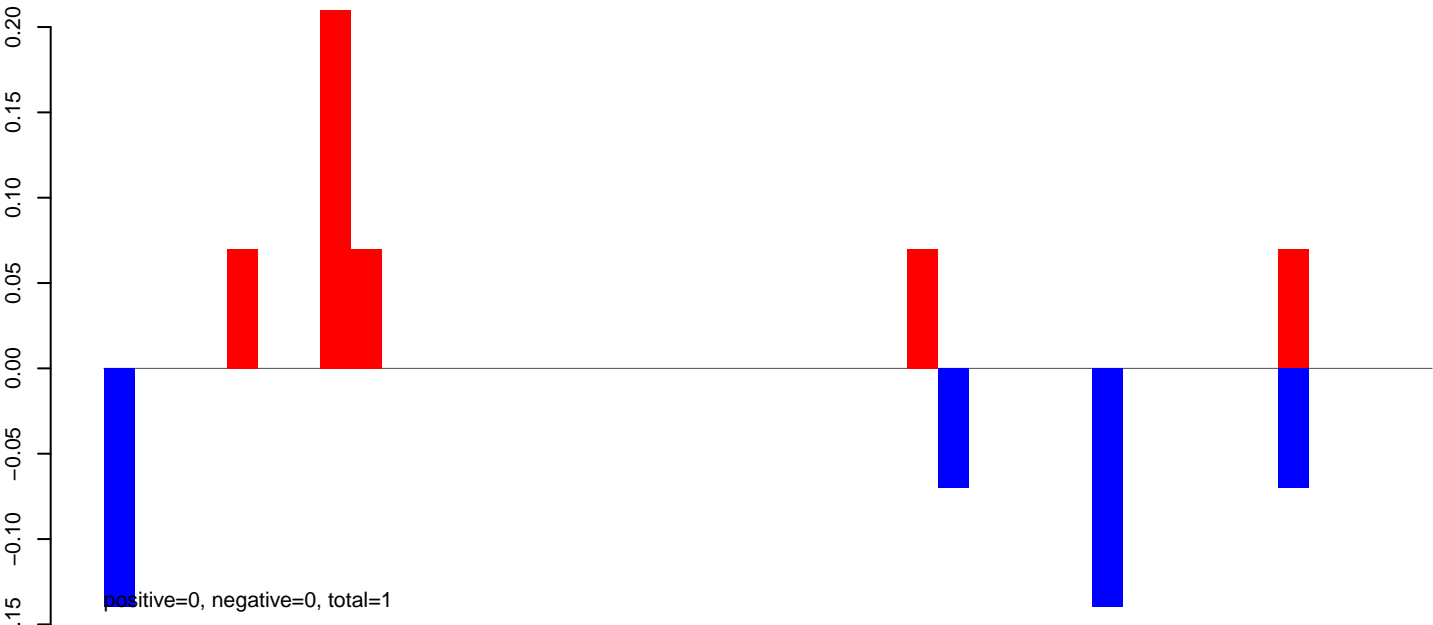
CuQuin_Mcarc_NL.rep



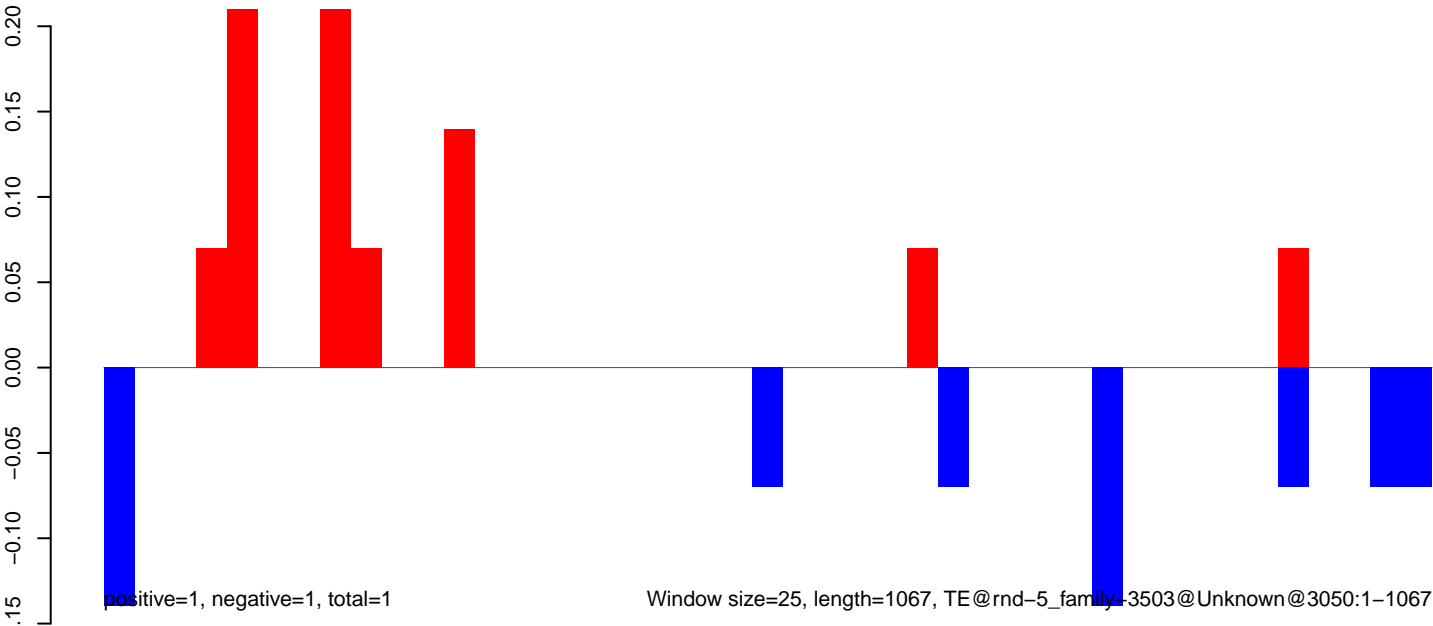
CuQuin_Mcarc_NL.18_23.rep



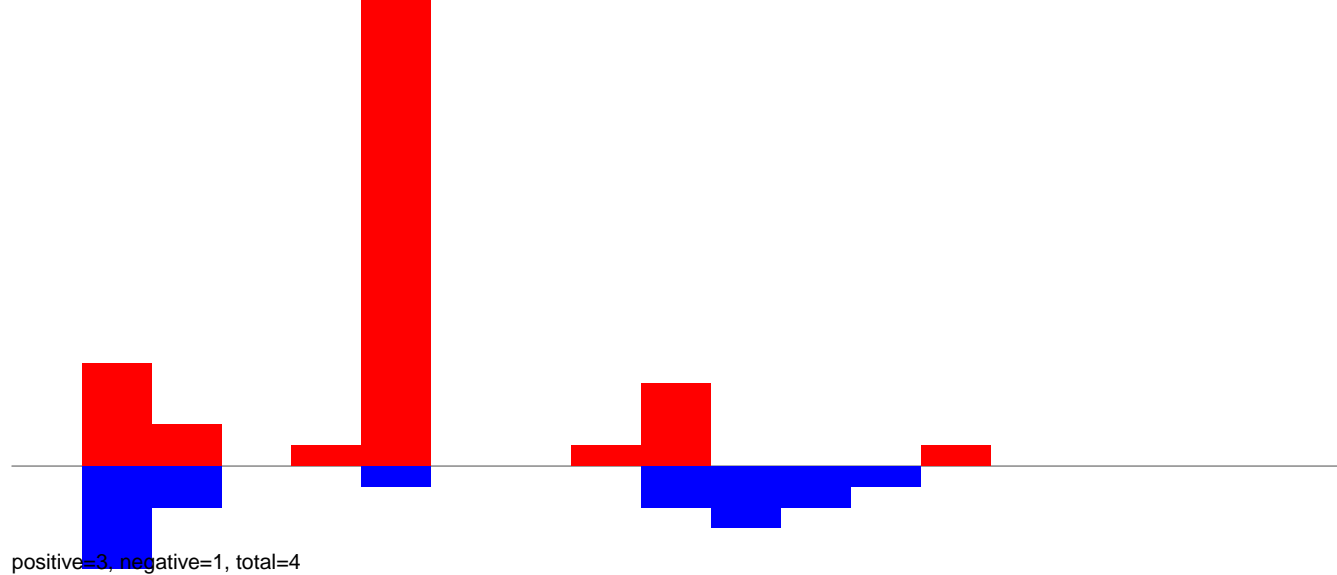
CuQuin_Mcarc_NL.24_35.rep



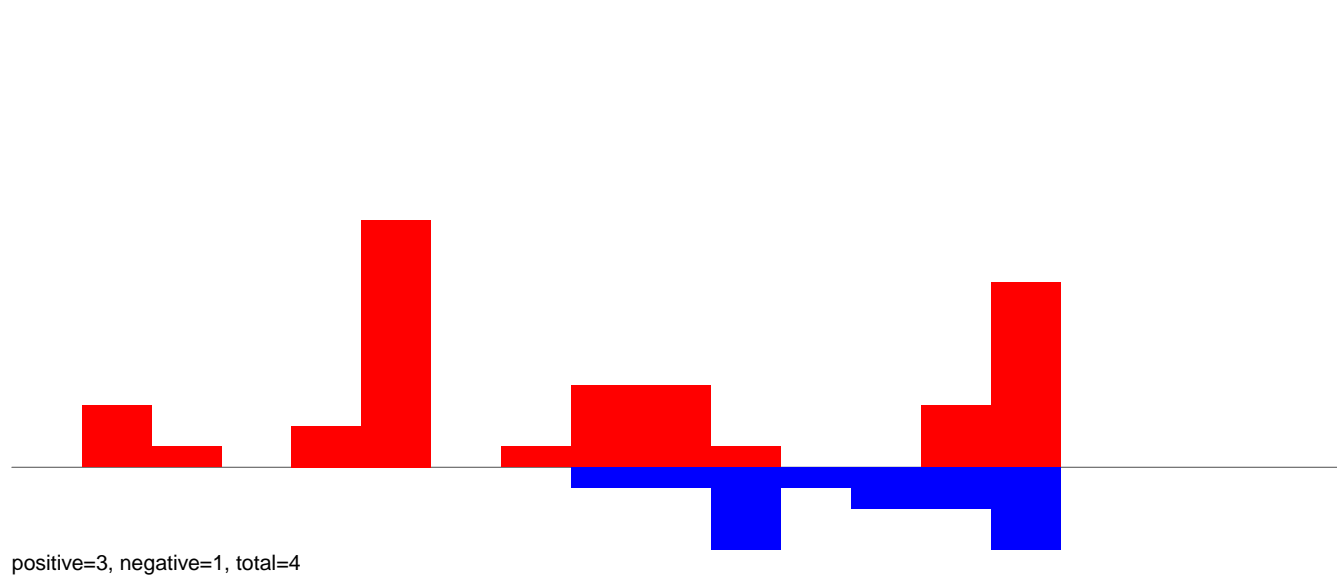
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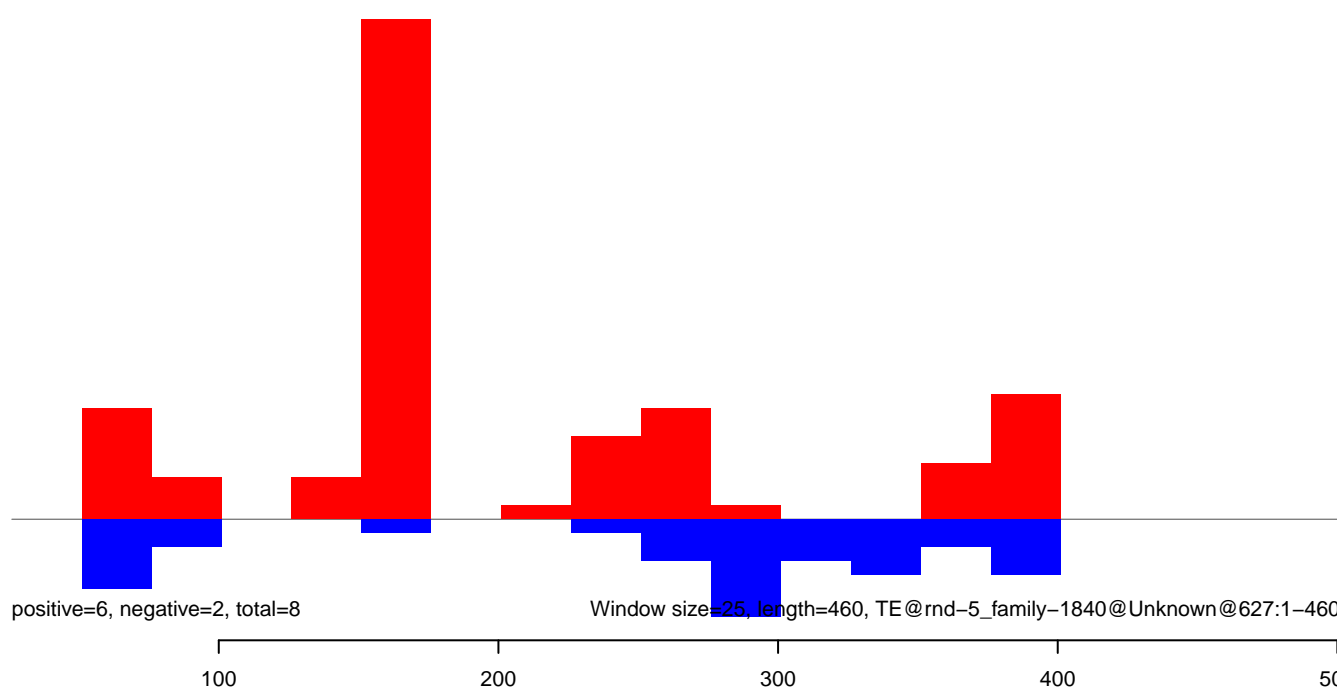
CuQuin_Mcarc_NL.18_23.rep



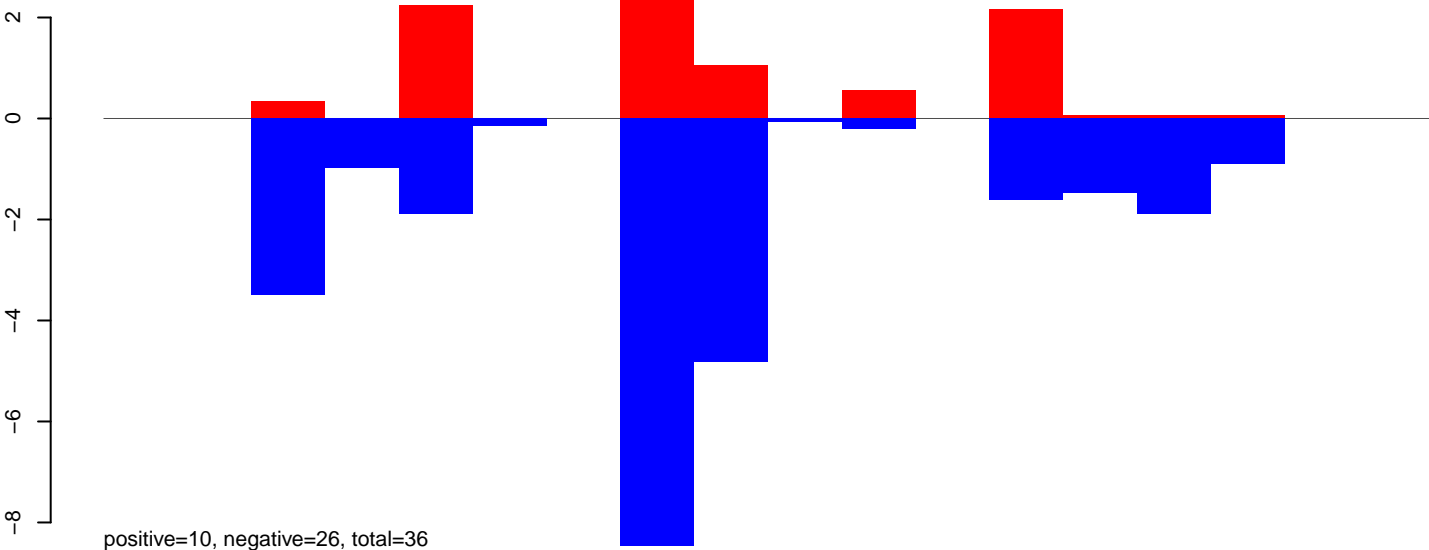
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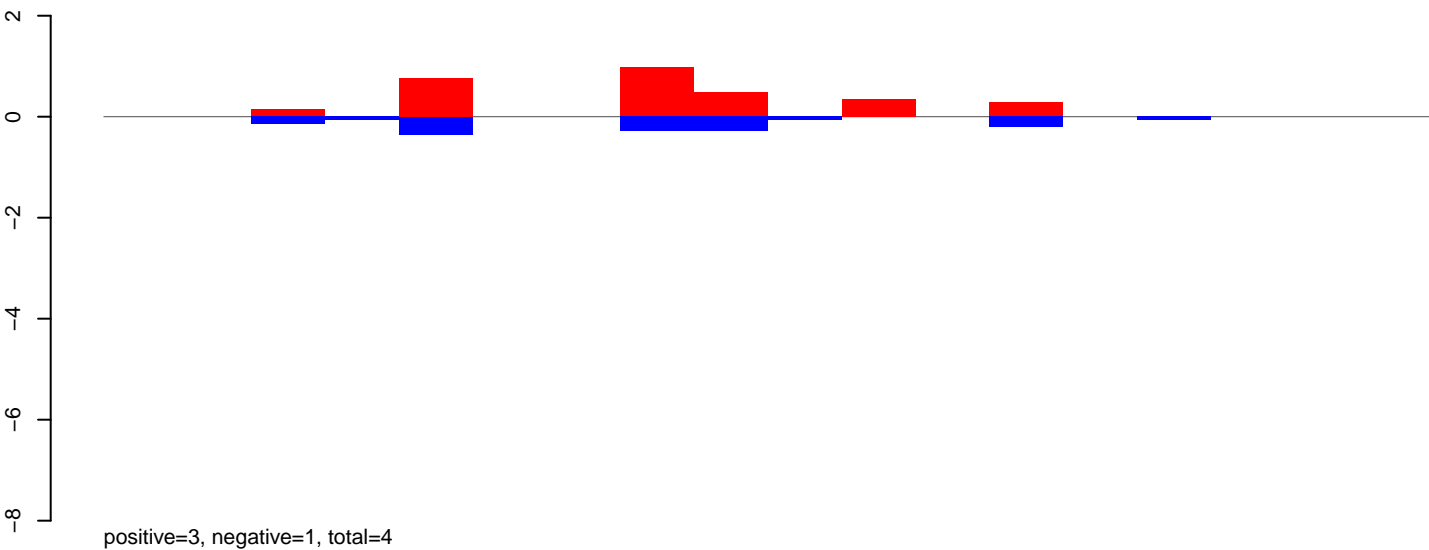
CuQuin_Mcarc_NL.rep



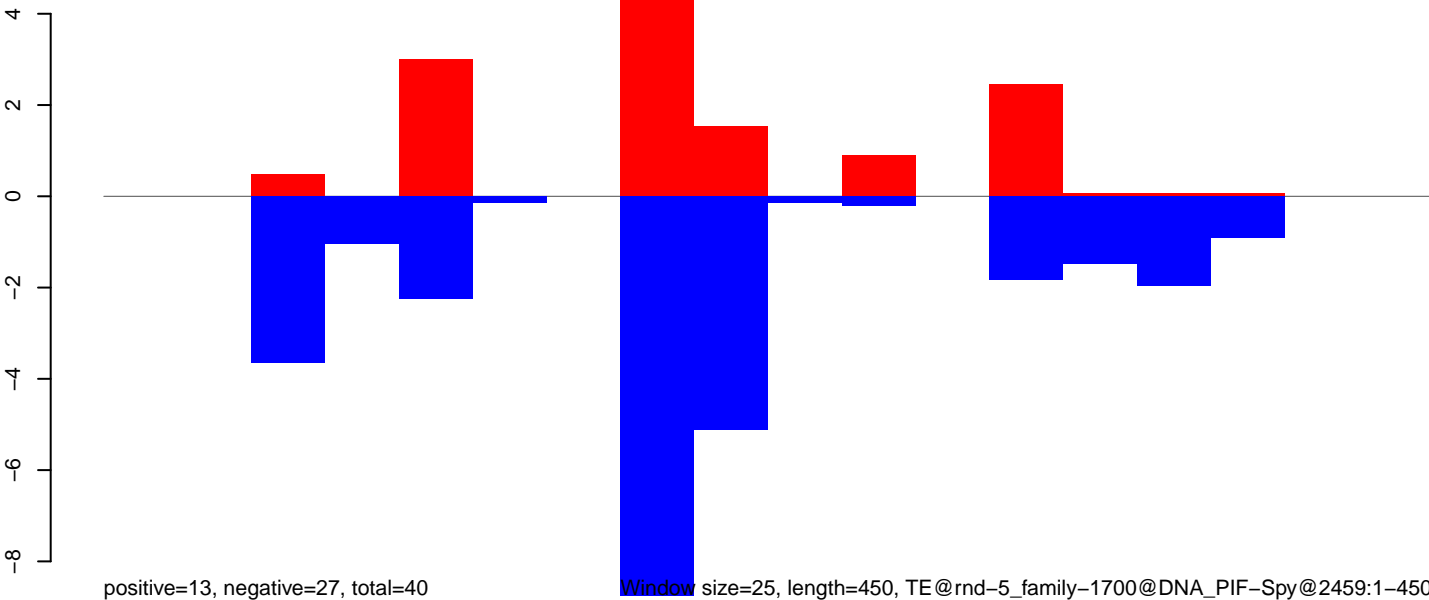
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

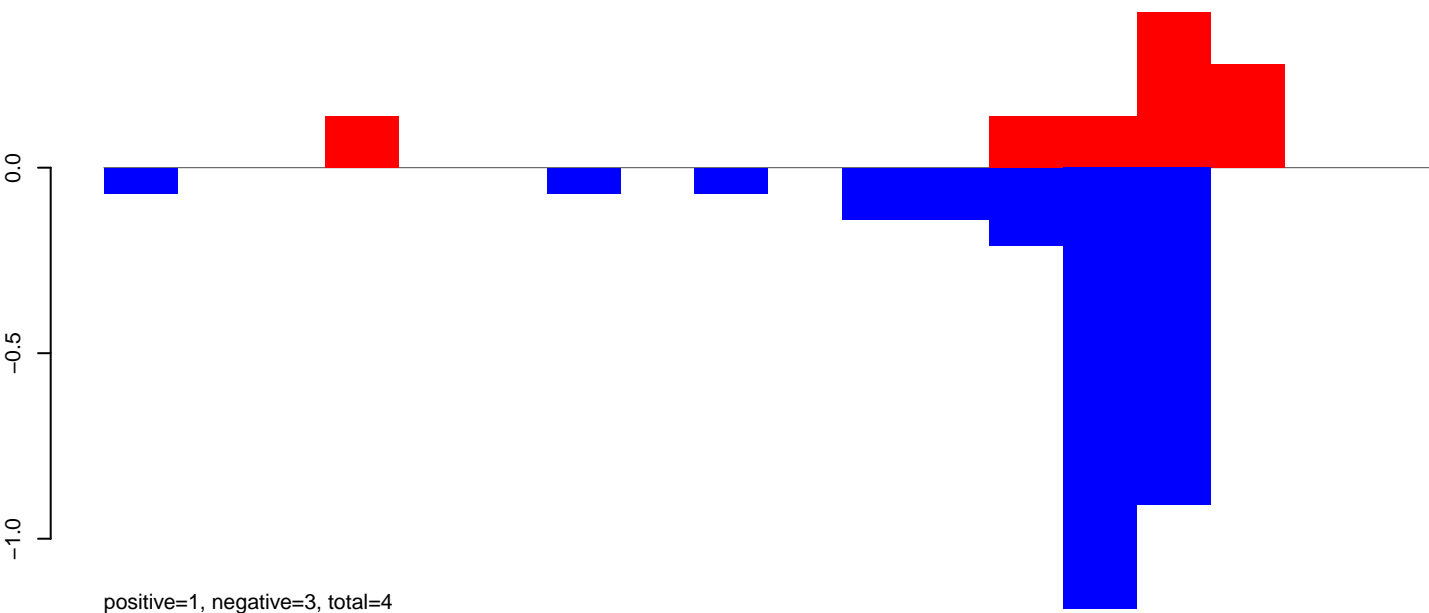


CuQuin_Mcarc_NL.rep

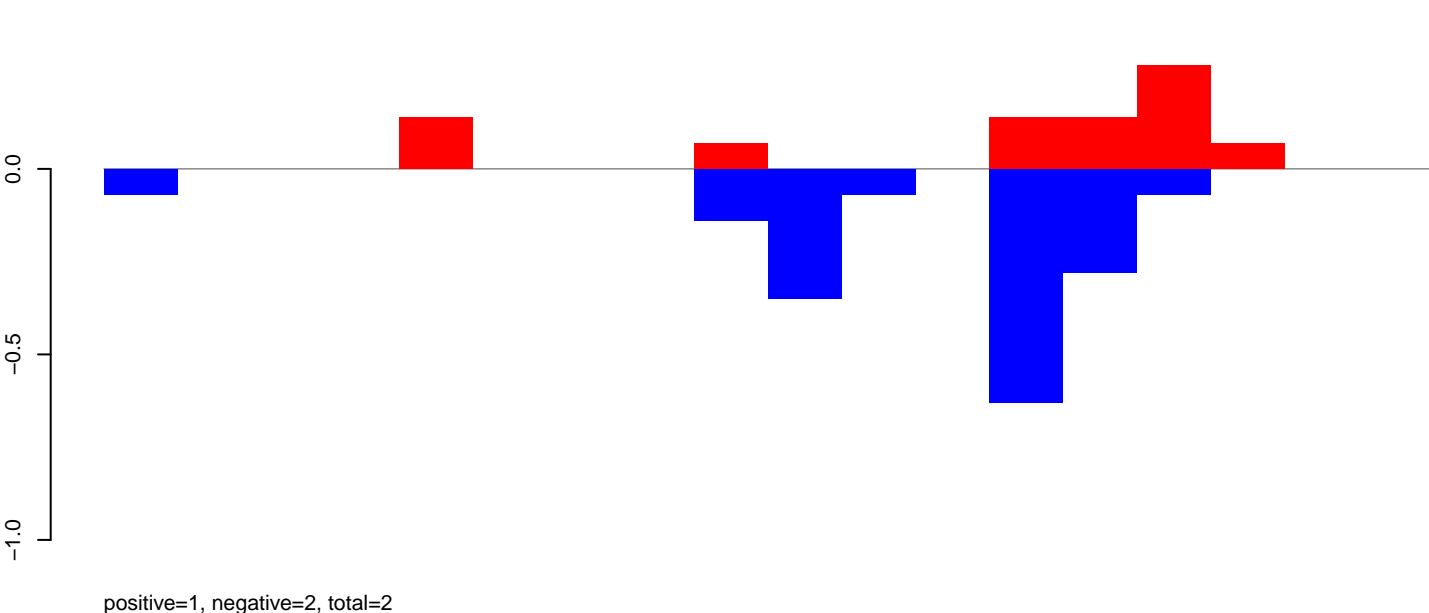


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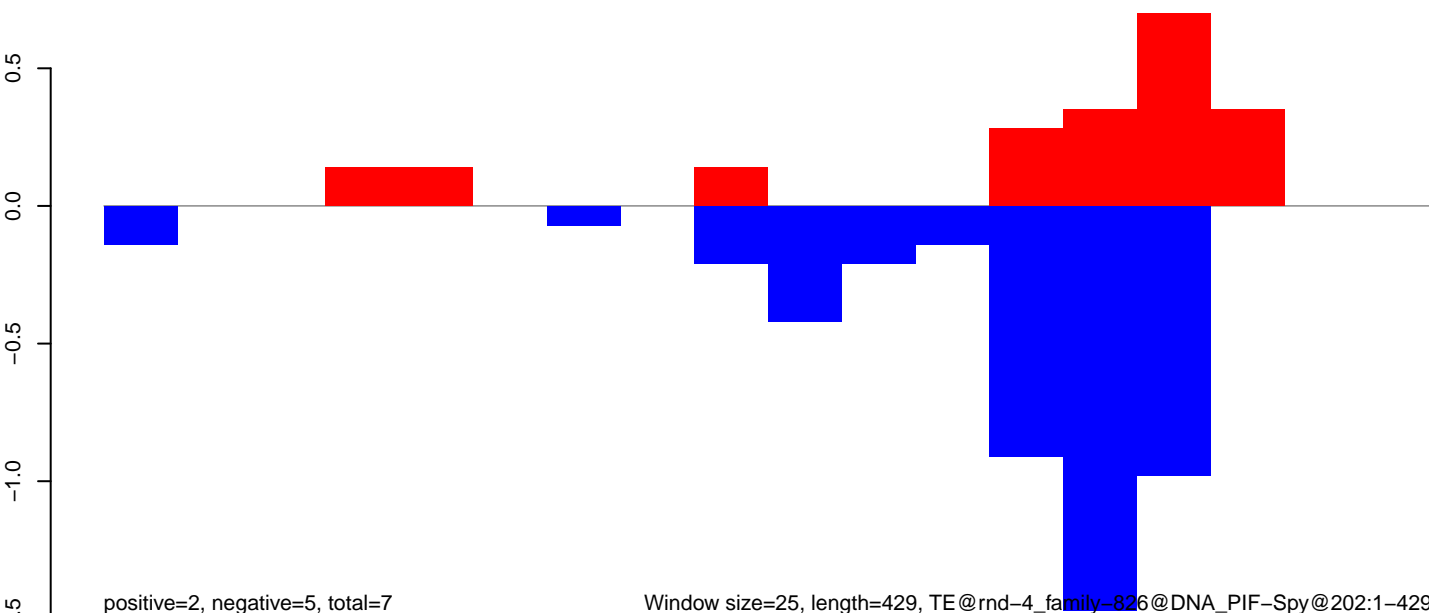
CuQuin_Mcarc_NL.18_23.rep



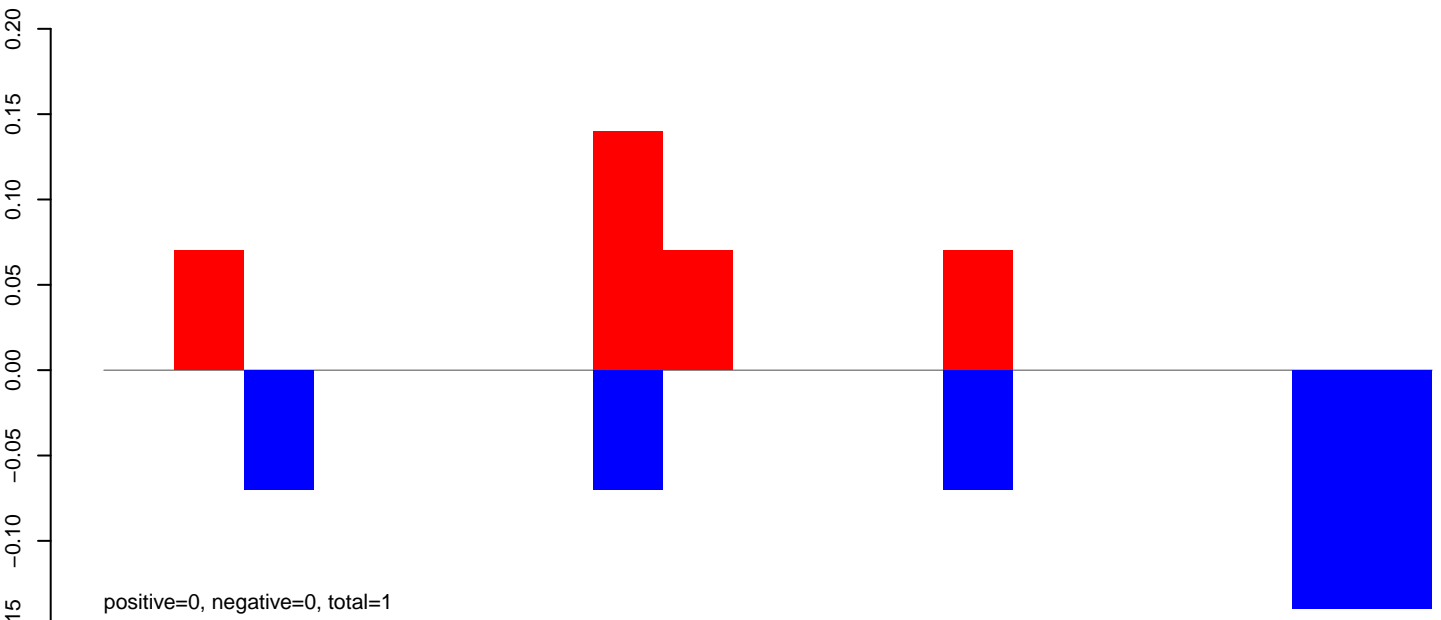
CuQuin_Mcarc_NL.24_35.rep



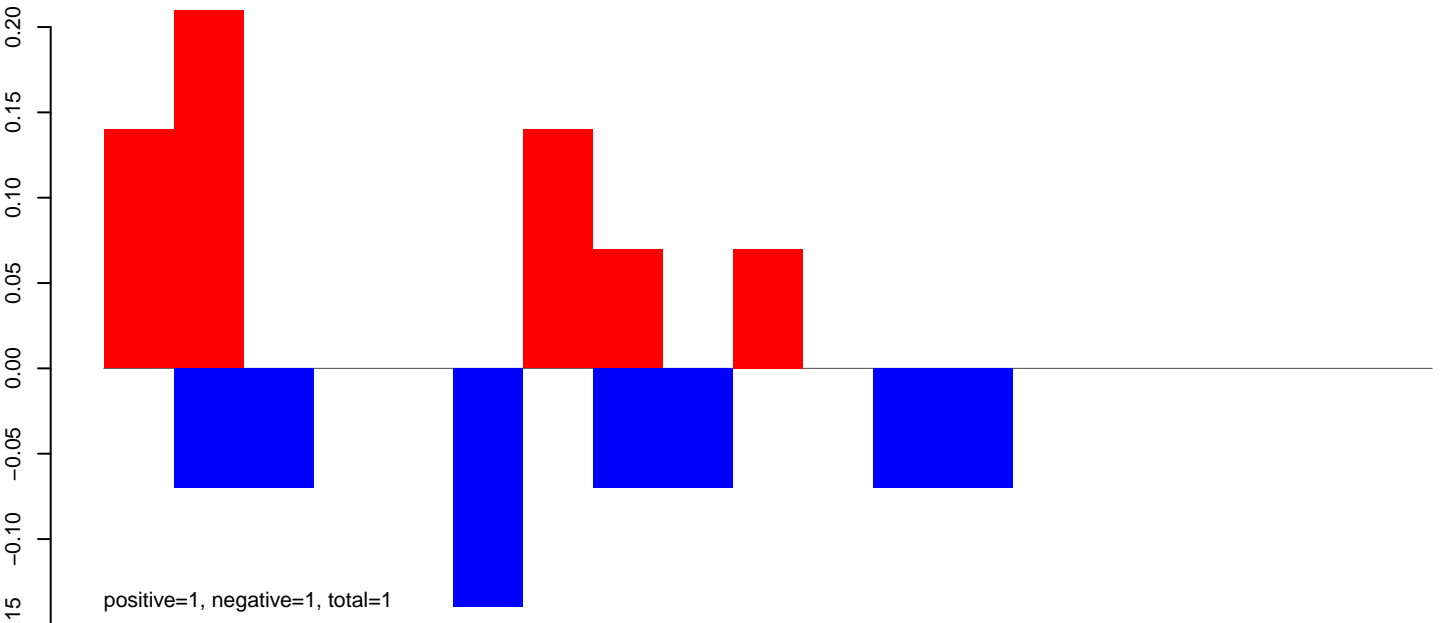
CuQuin_Mcarc_NL.rep



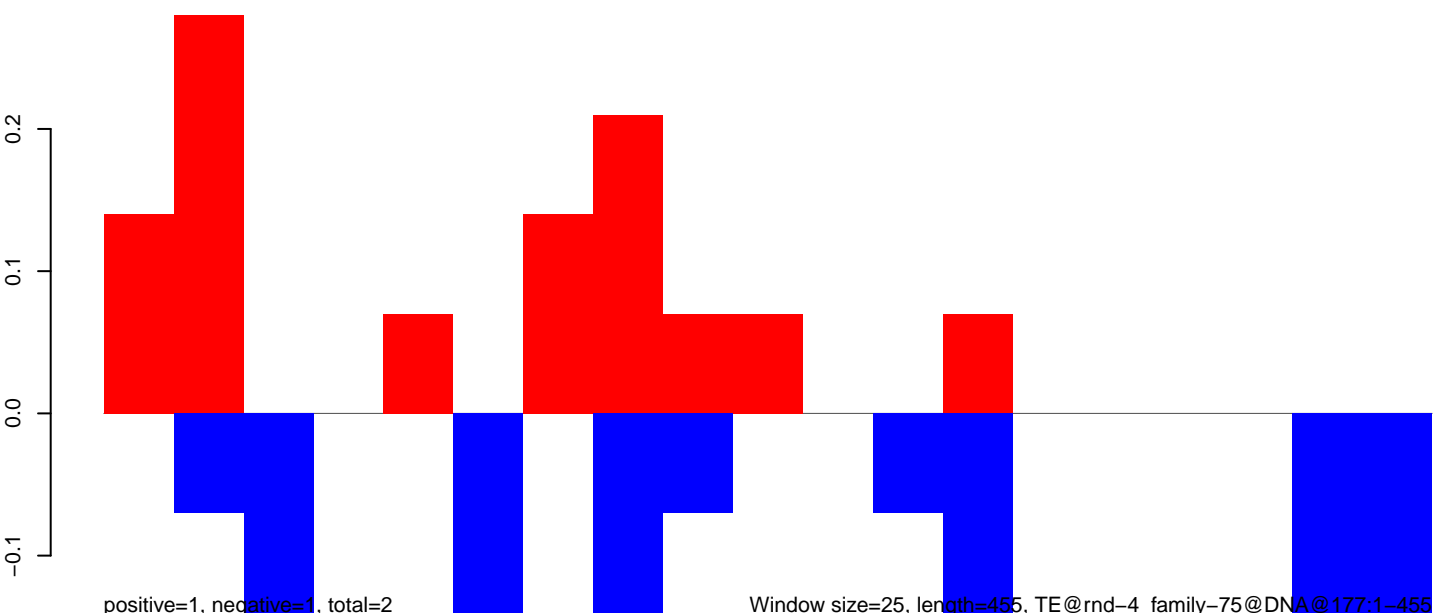
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



100

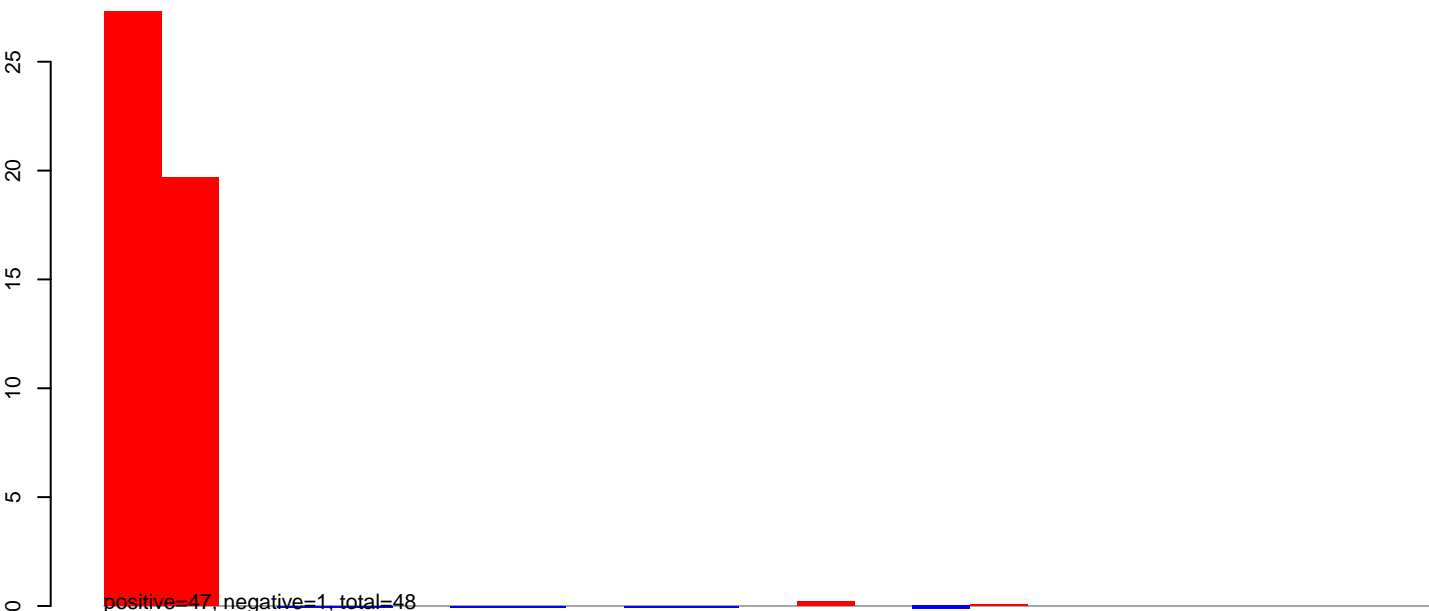
200

300

400

500

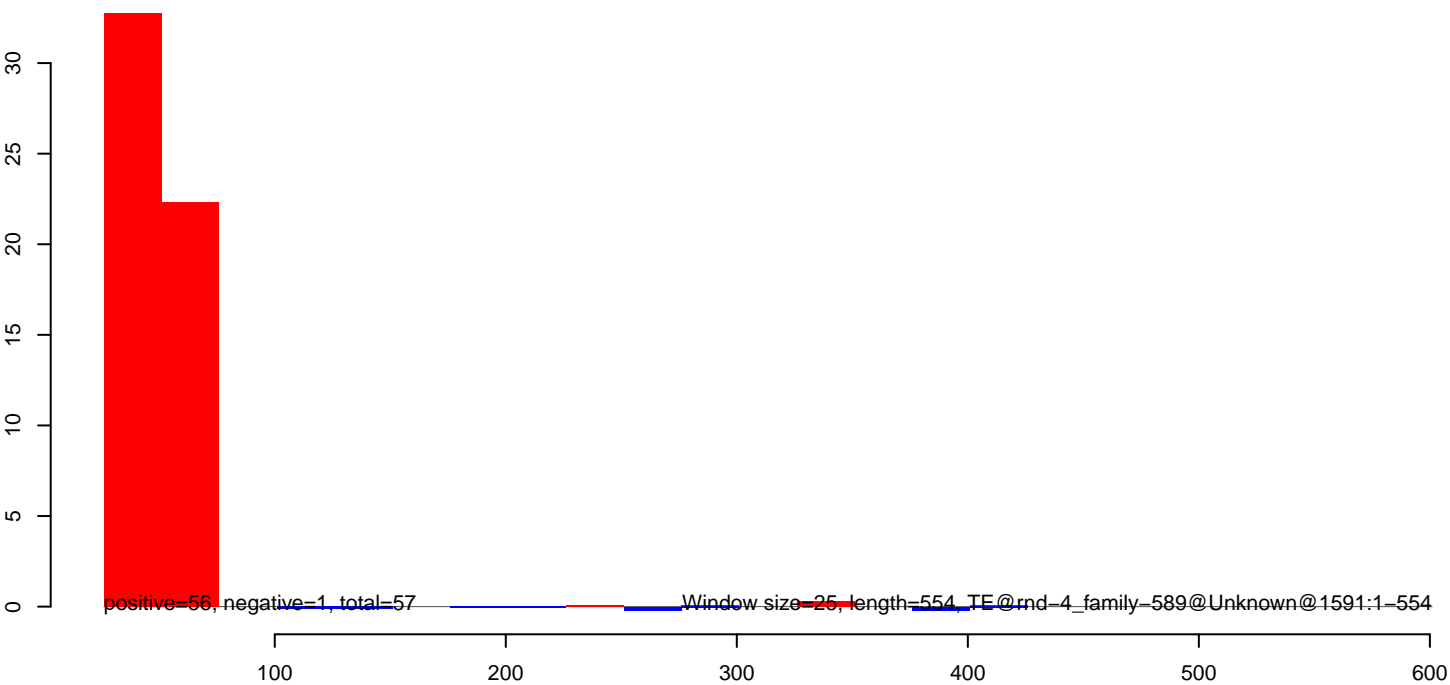
CuQuin_Mcarc_NL.18_23.rep



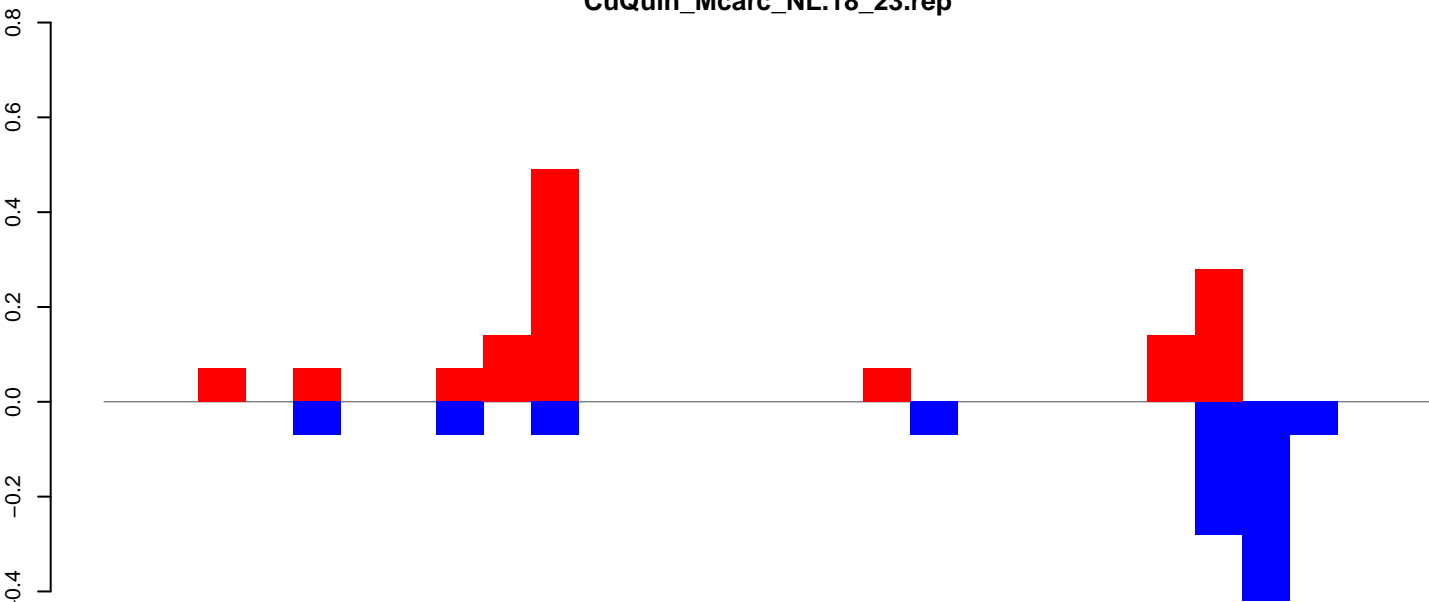
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

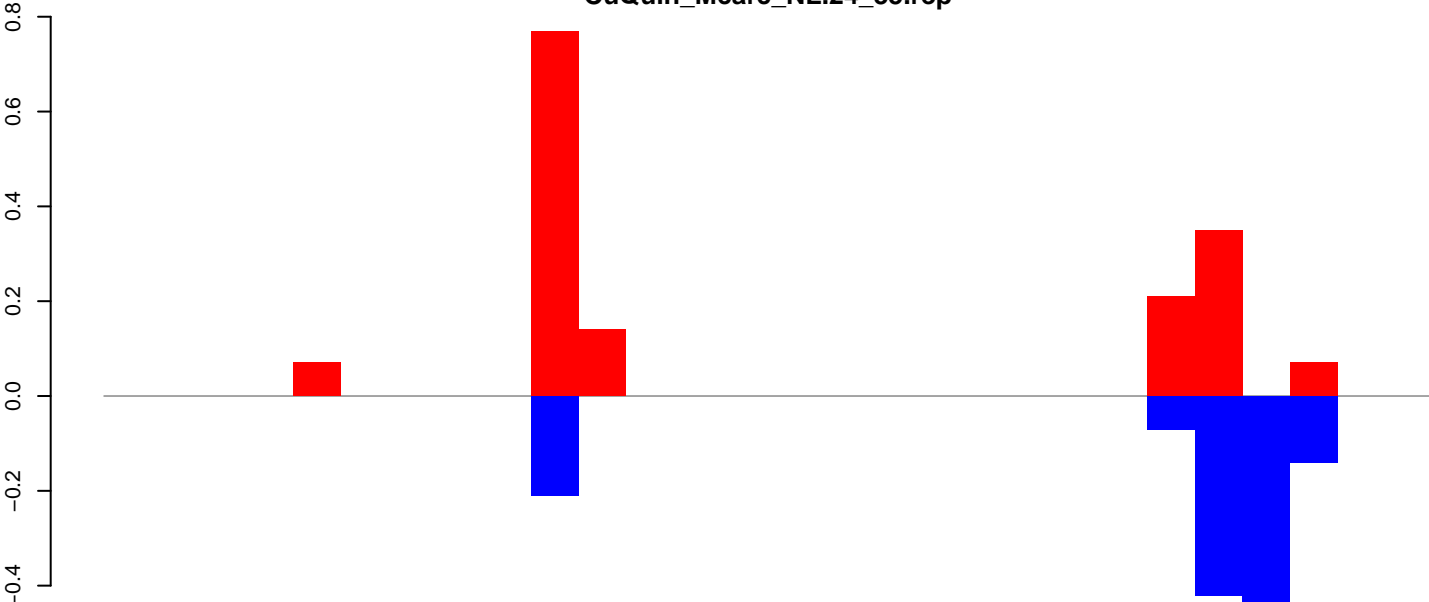


CuQuin_Mcarc_NL.18_23.rep



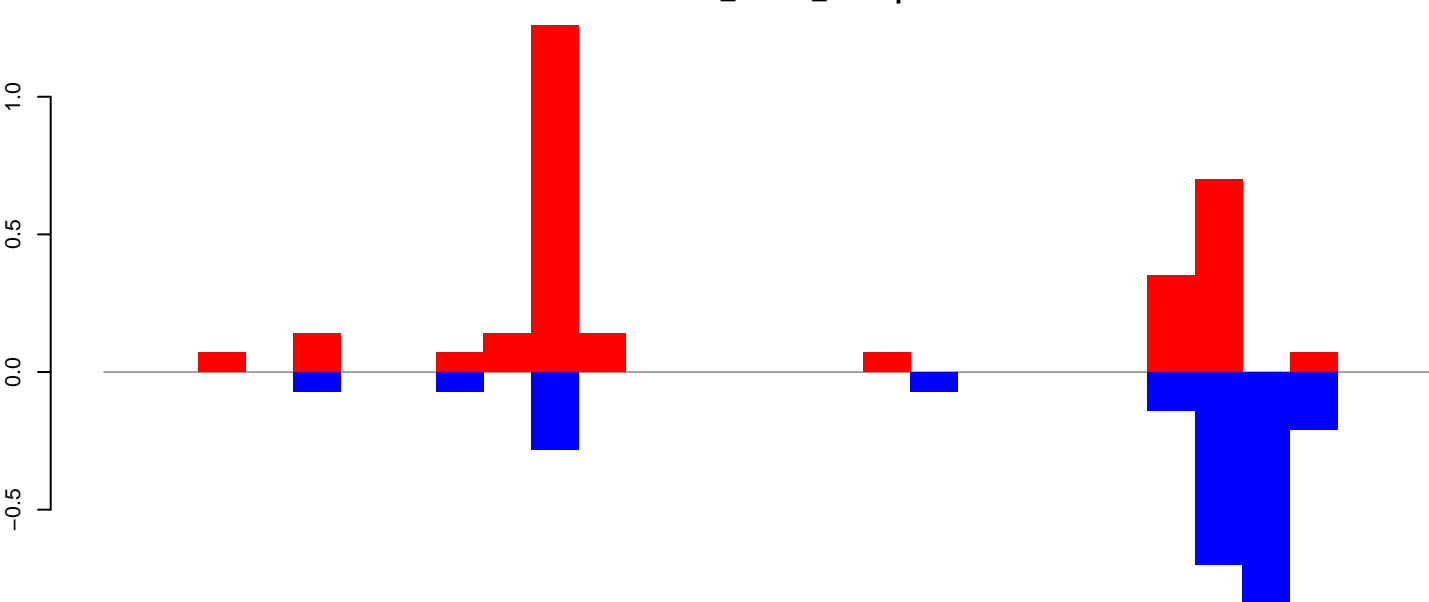
positive=1, negative=1, total=2

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=1, total=3

CuQuin_Mcarc_NL.rep

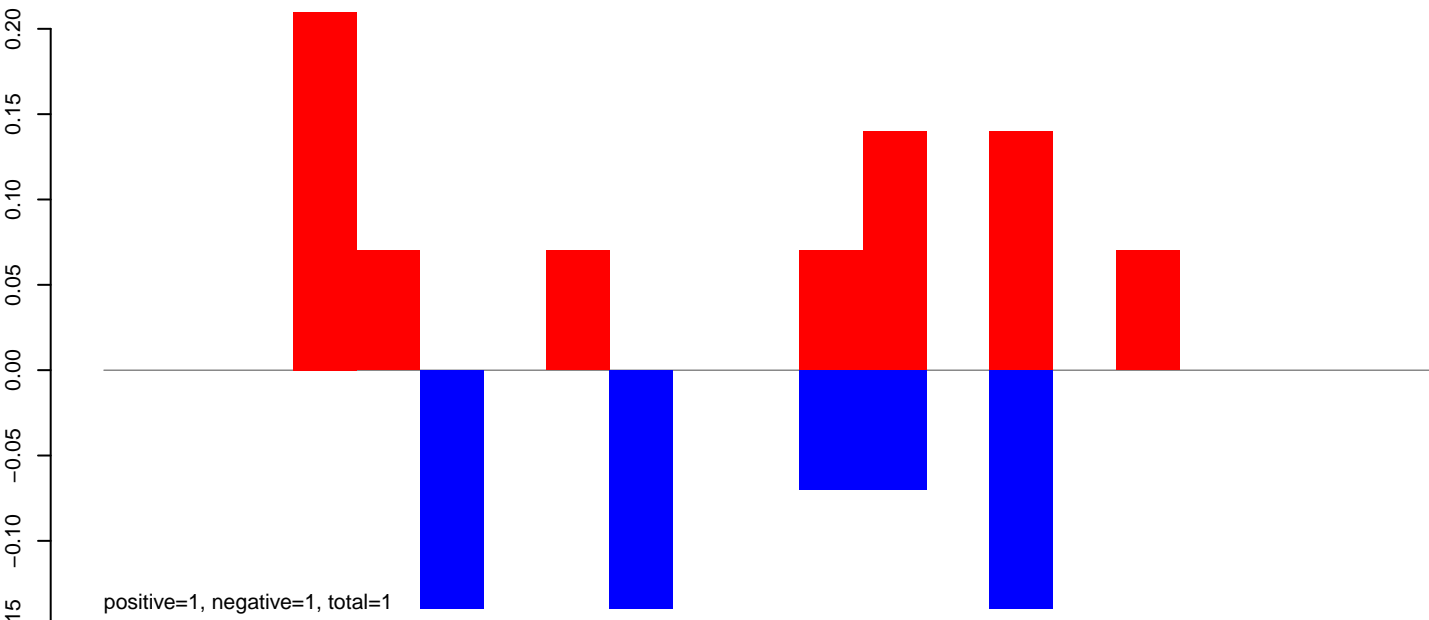


positive=3, negative=2, total=5

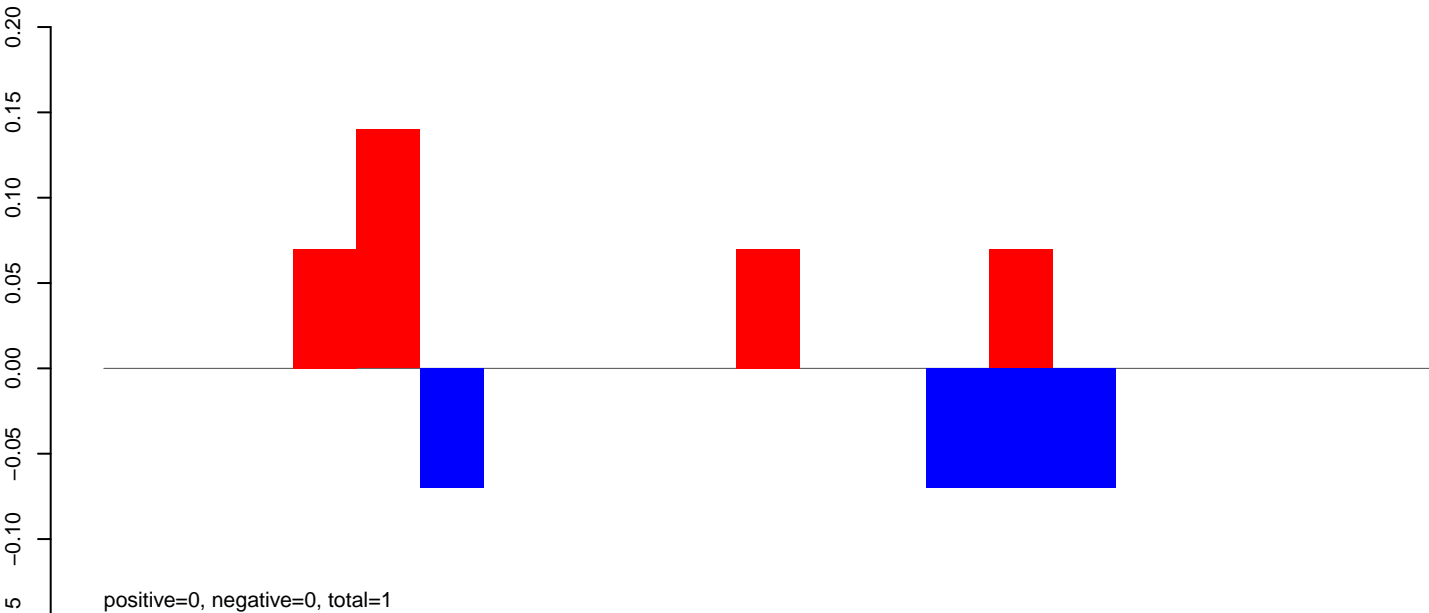
Window size=25, length=697, TE@rnd-4_family-474@DNA_MULE-MuDR@2761:1-697

0 100 200 300 400 500 600 700

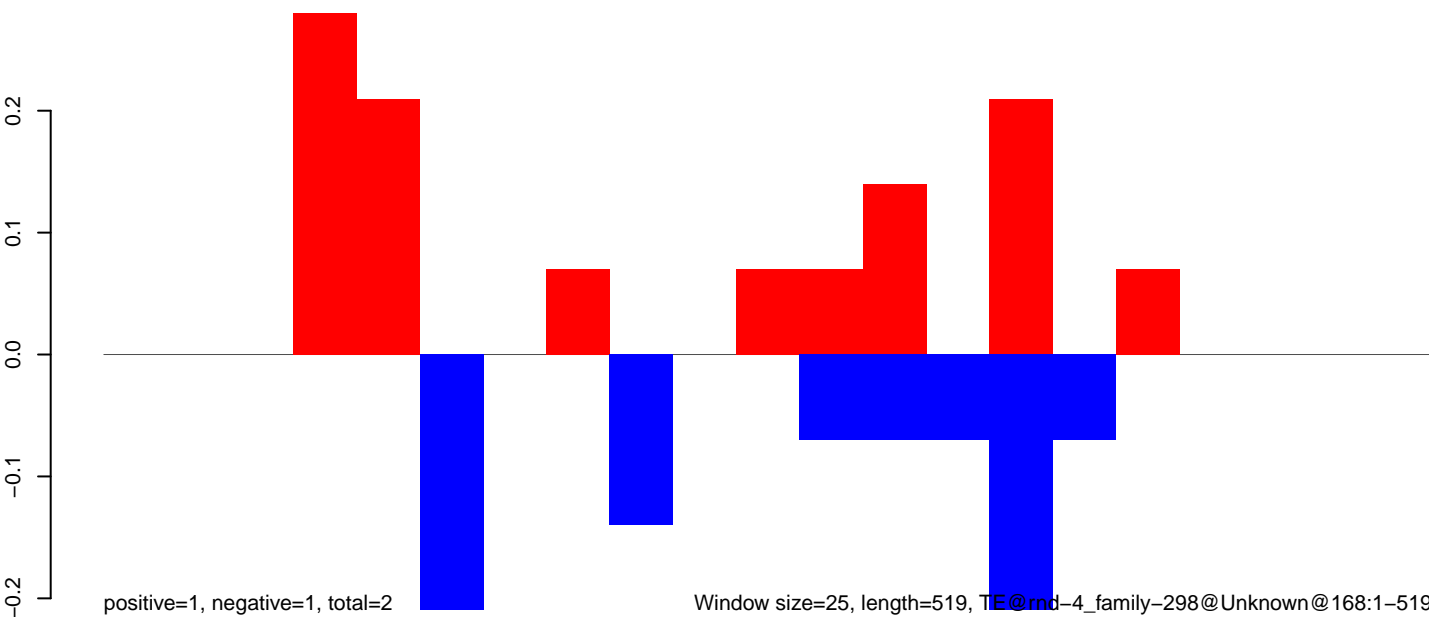
CuQuin_Mcarc_NL.18_23.rep



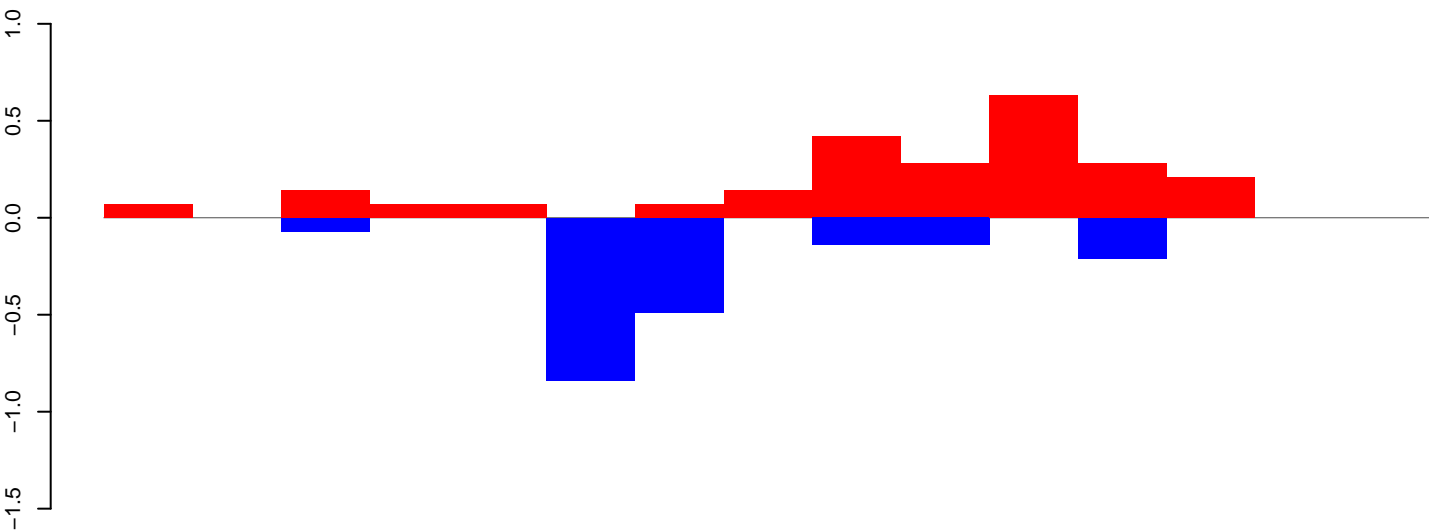
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

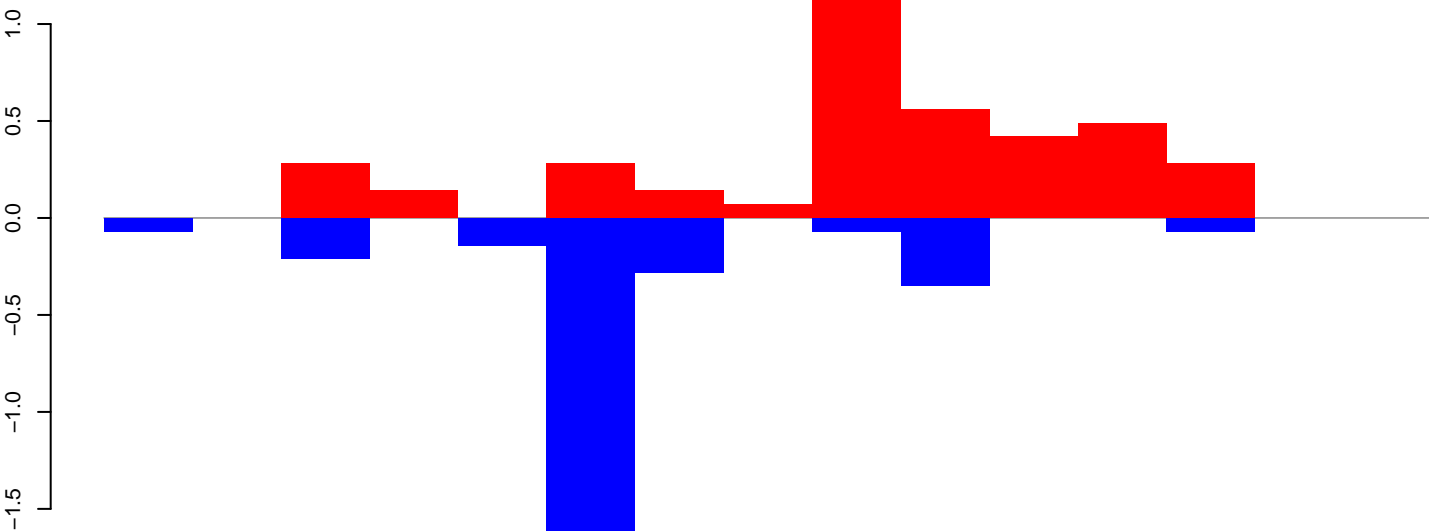


CuQuin_Mcarc_NL.18_23.rep



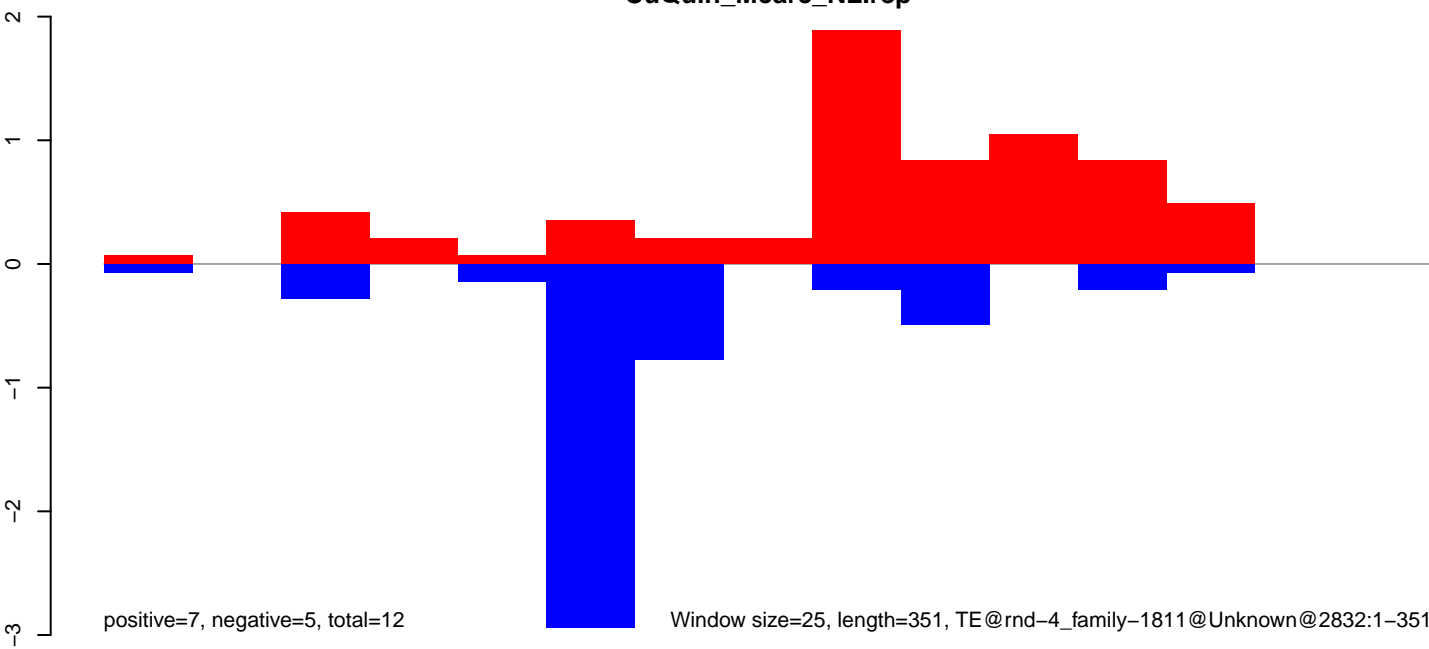
positive=2, negative=2, total=4

CuQuin_Mcarc_NL.24_35.rep



positive=4, negative=3, total=7

CuQuin_Mcarc_NL.rep



positive=7, negative=5, total=12

Window size=25, length=351, TE@rnd-4_family-1811@Unknown@2832:1-351

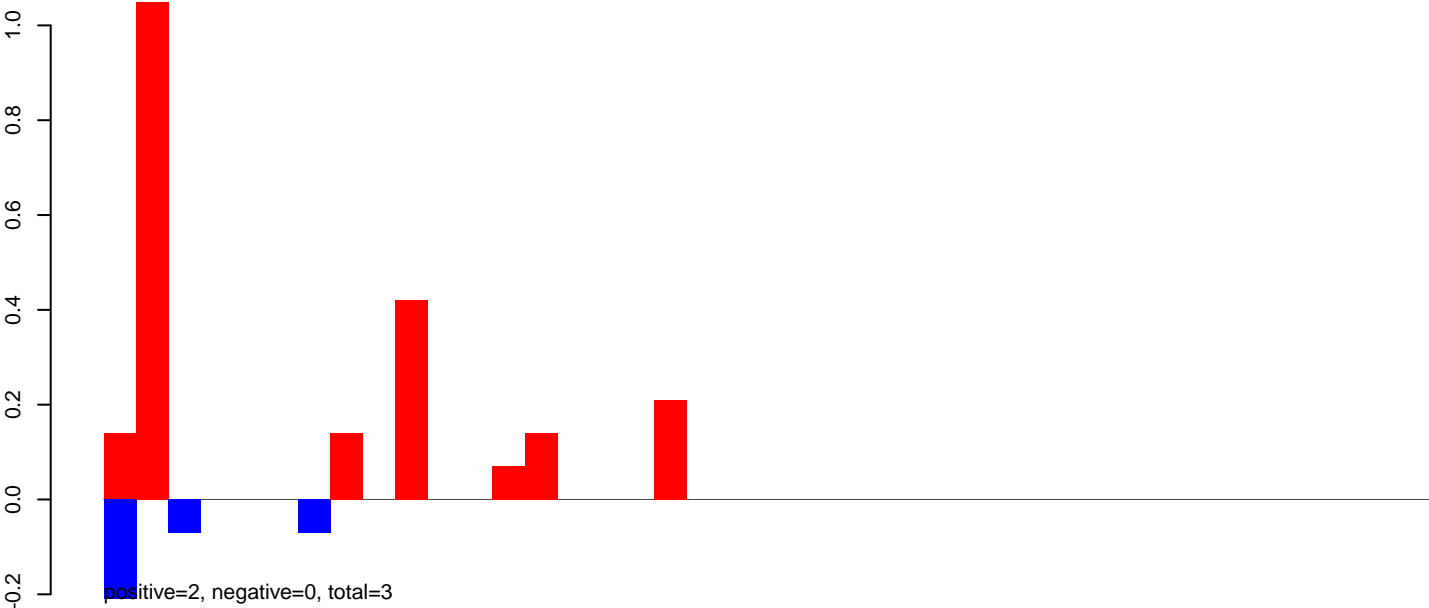
100

200

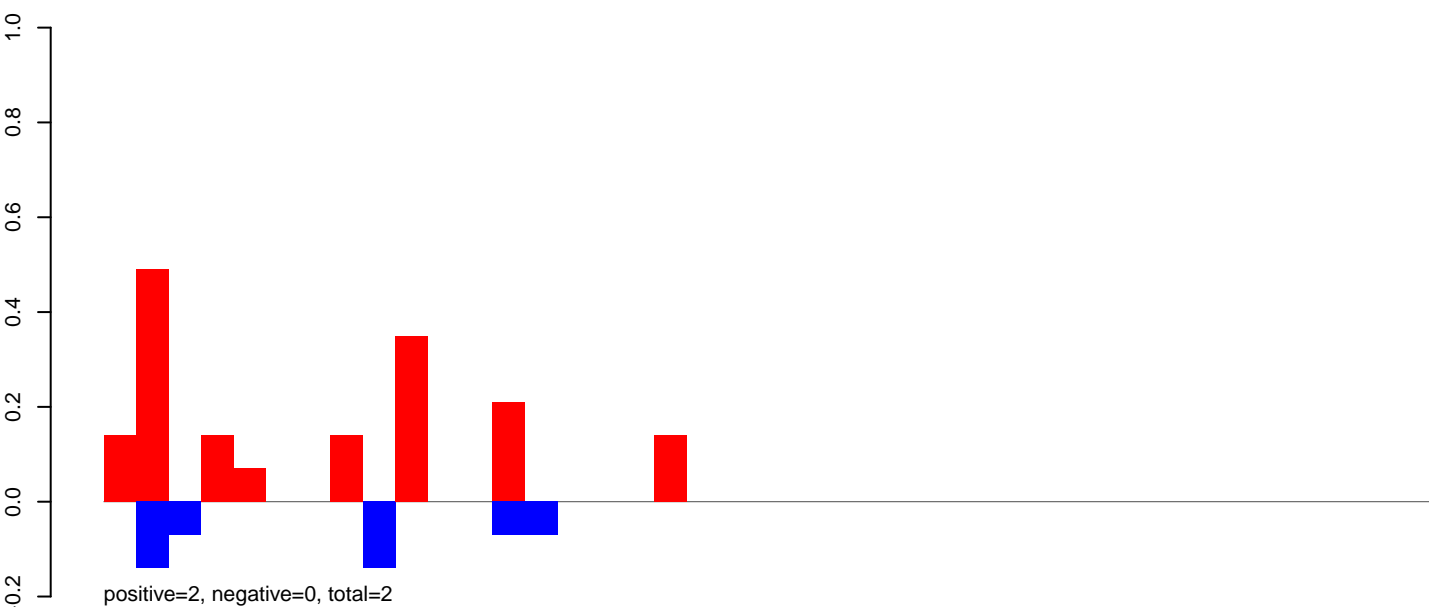
300

400

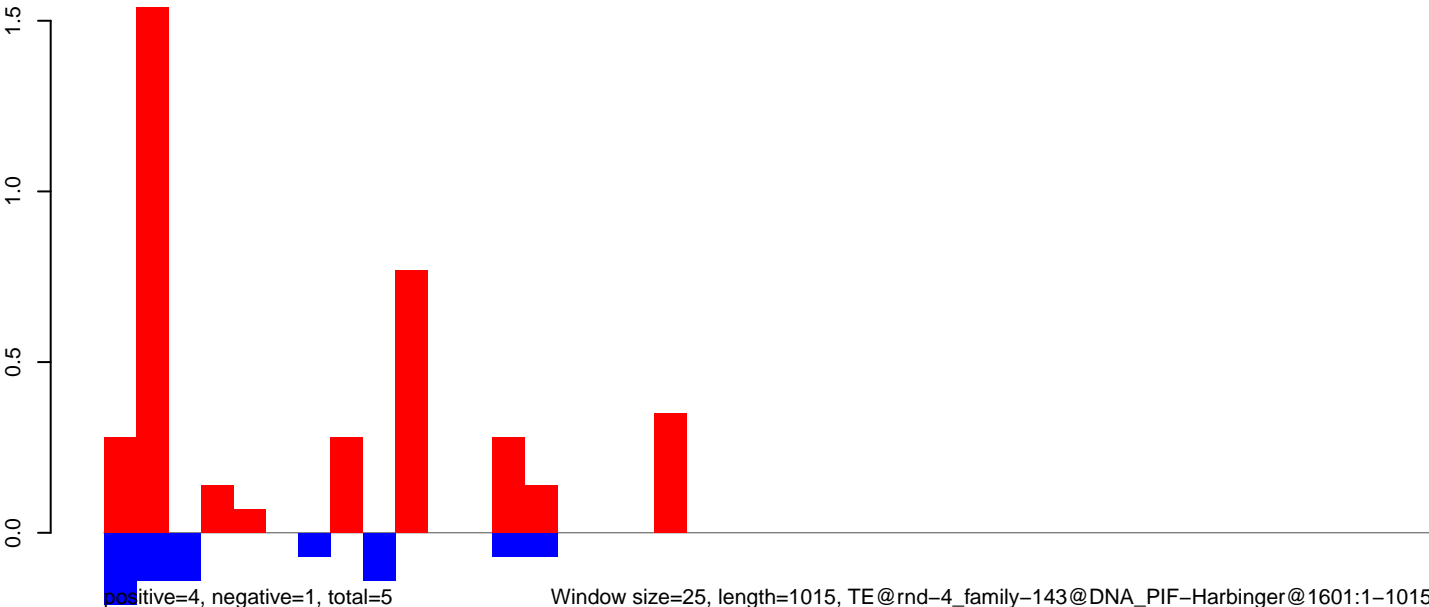
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



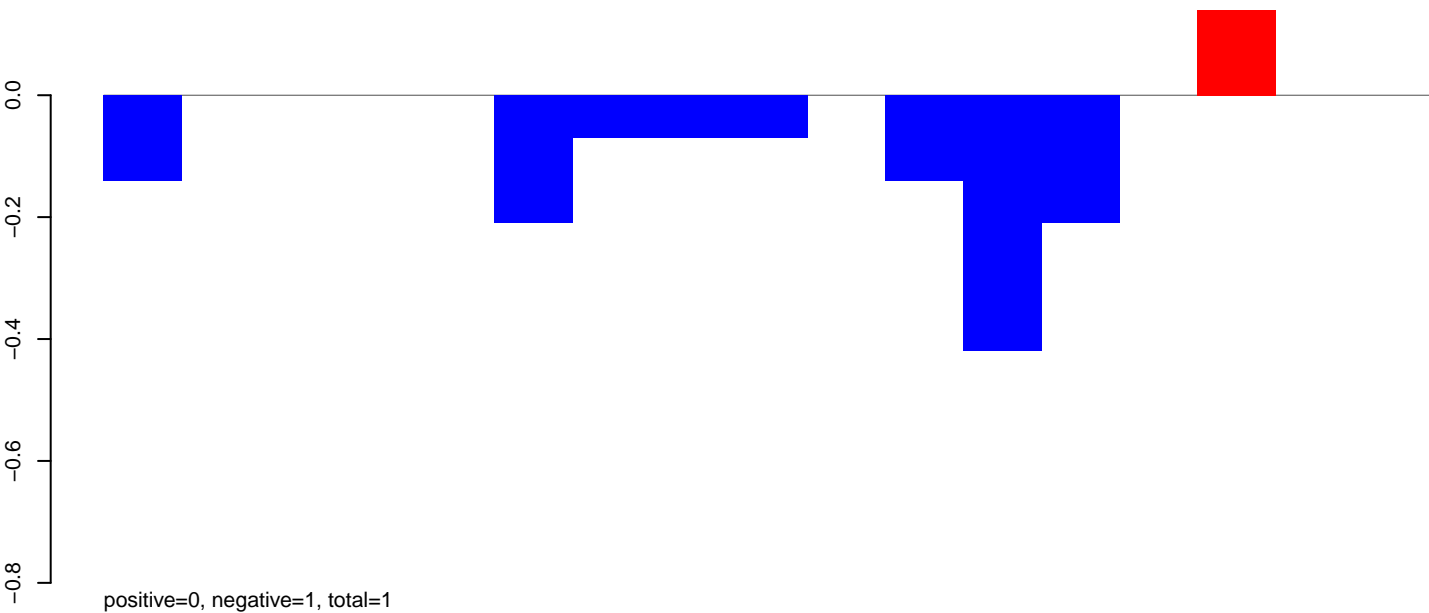
CuQuin_Mcarc_NL.rep



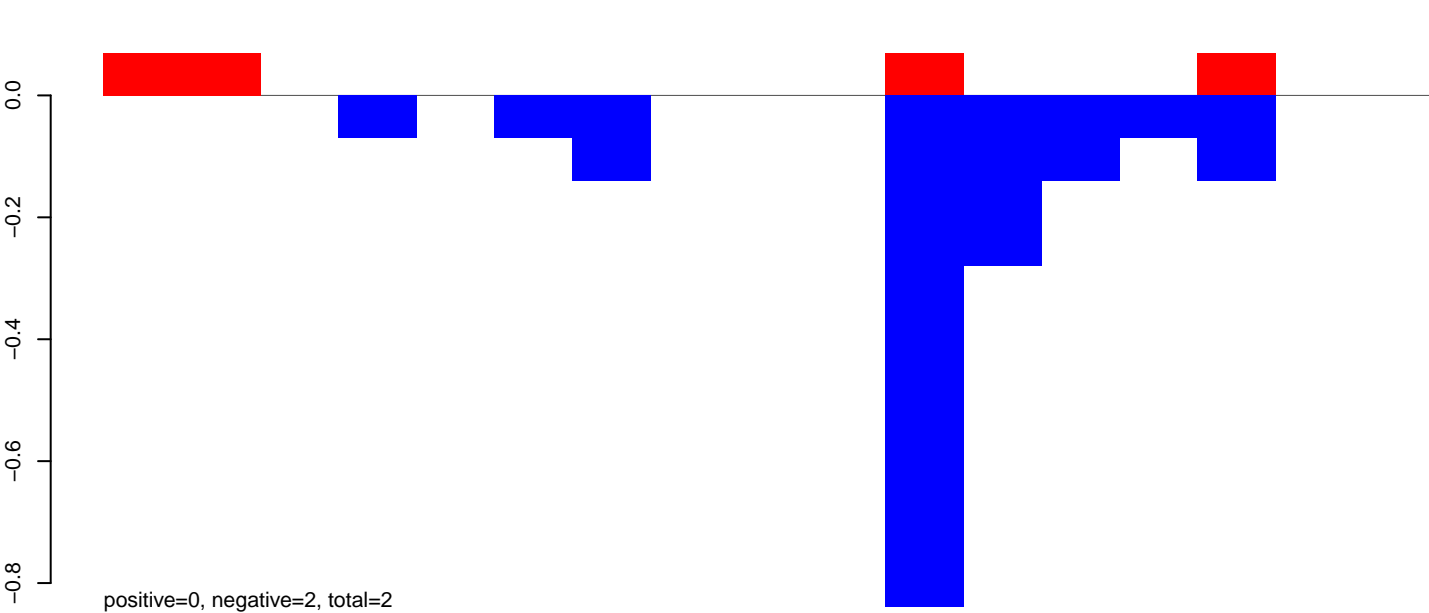
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0 200 400 600 800 1000

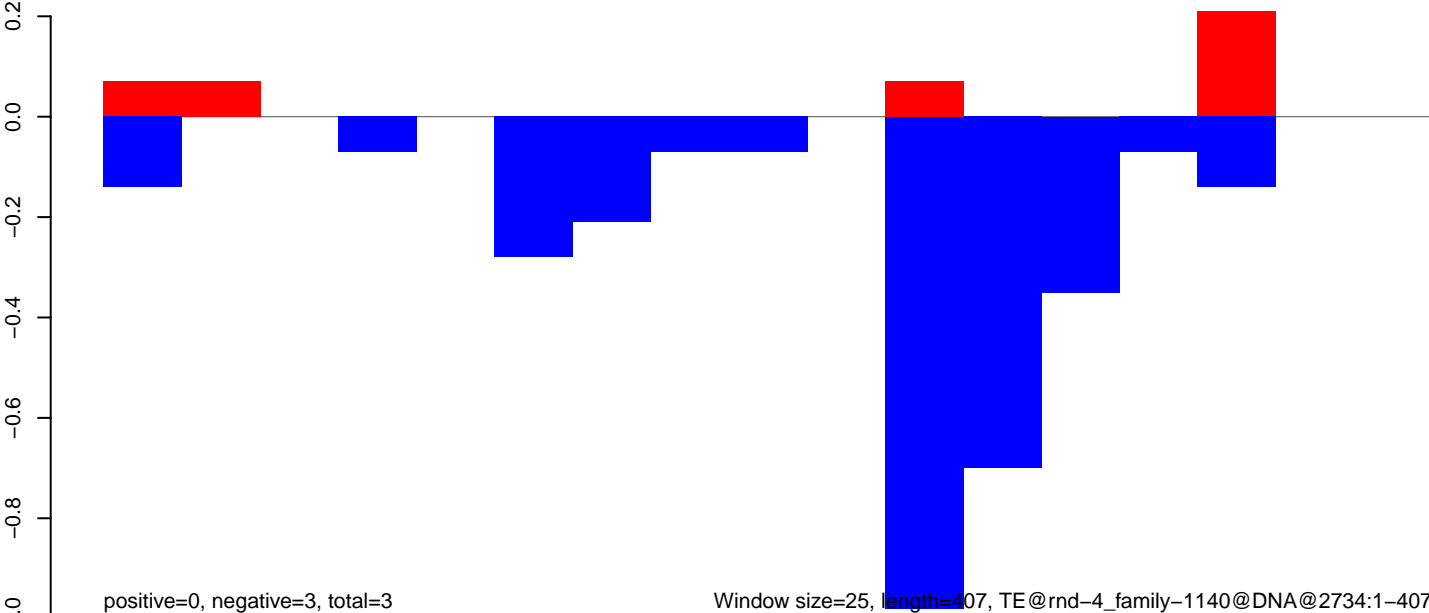
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

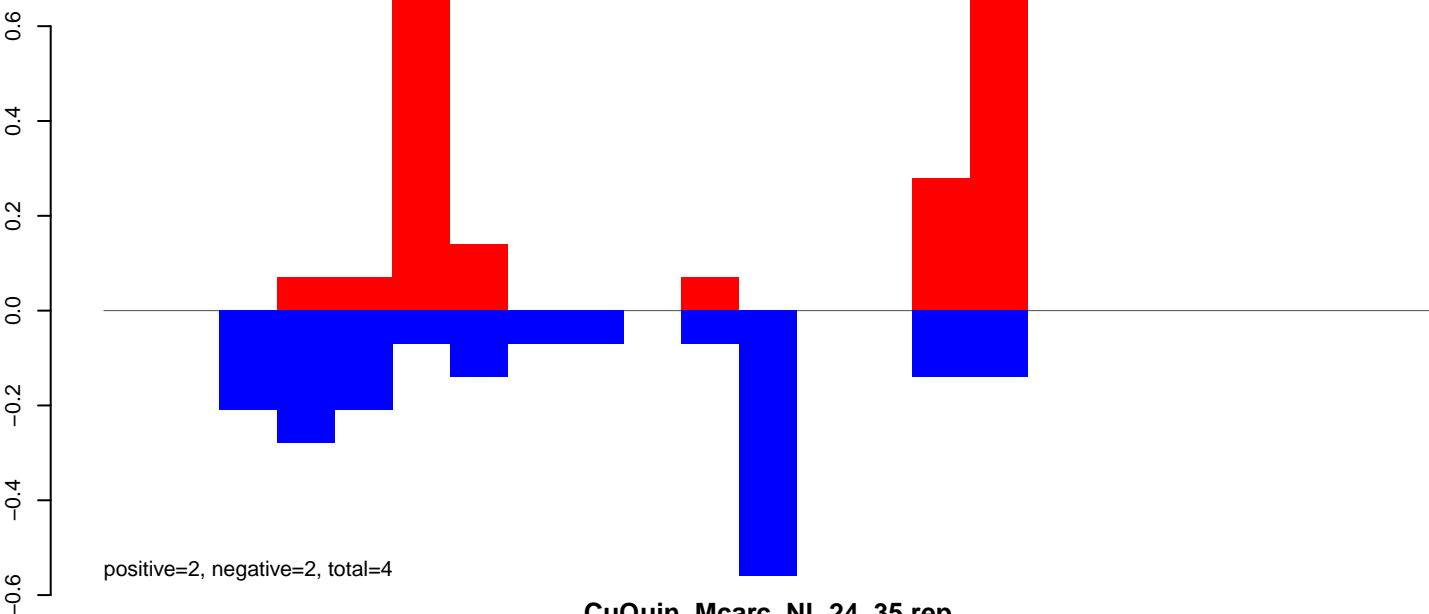


CuQuin_Mcarc_NL.rep

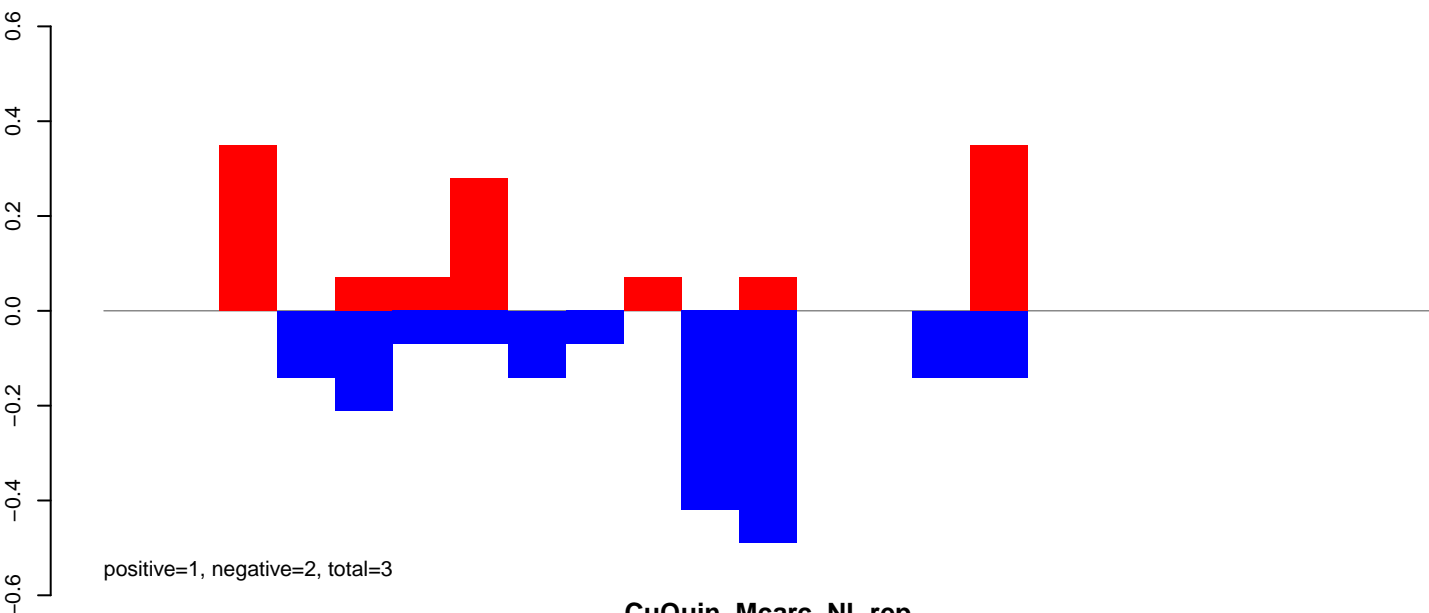


Window size=25, length=407, TE@rnd-4_family-1140@DNA@2734:1-407

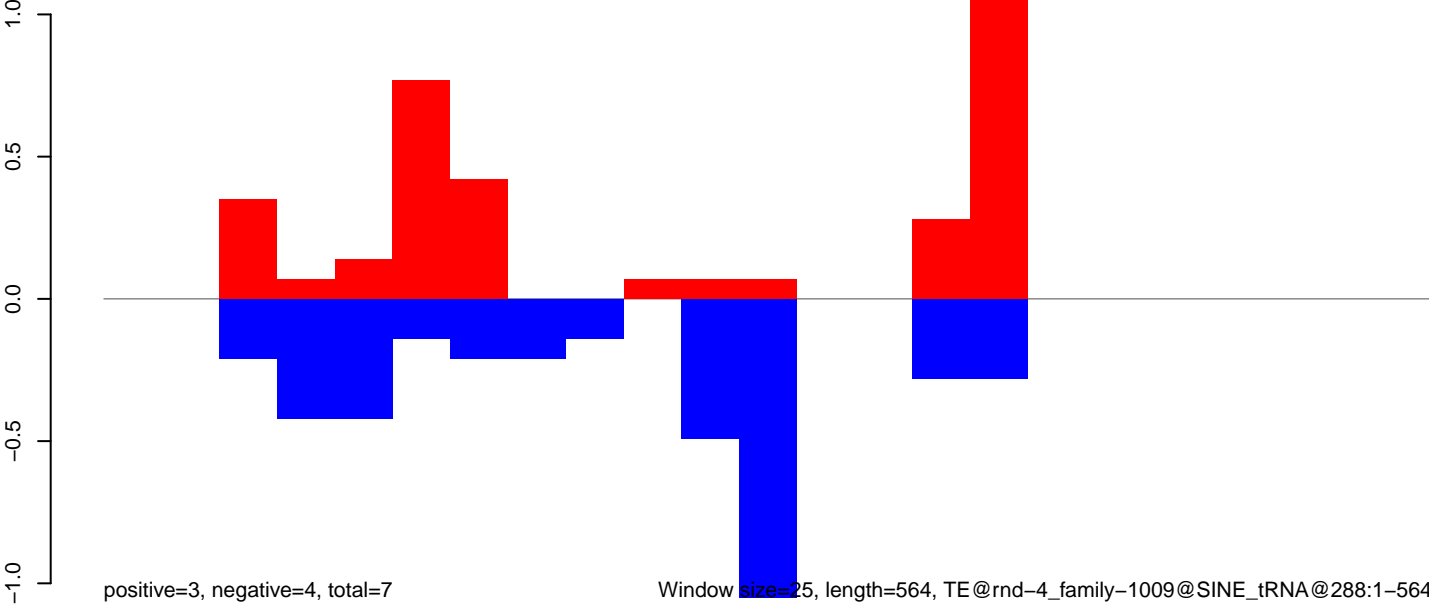
CuQuin_Mcarc_NL.18_23.rep



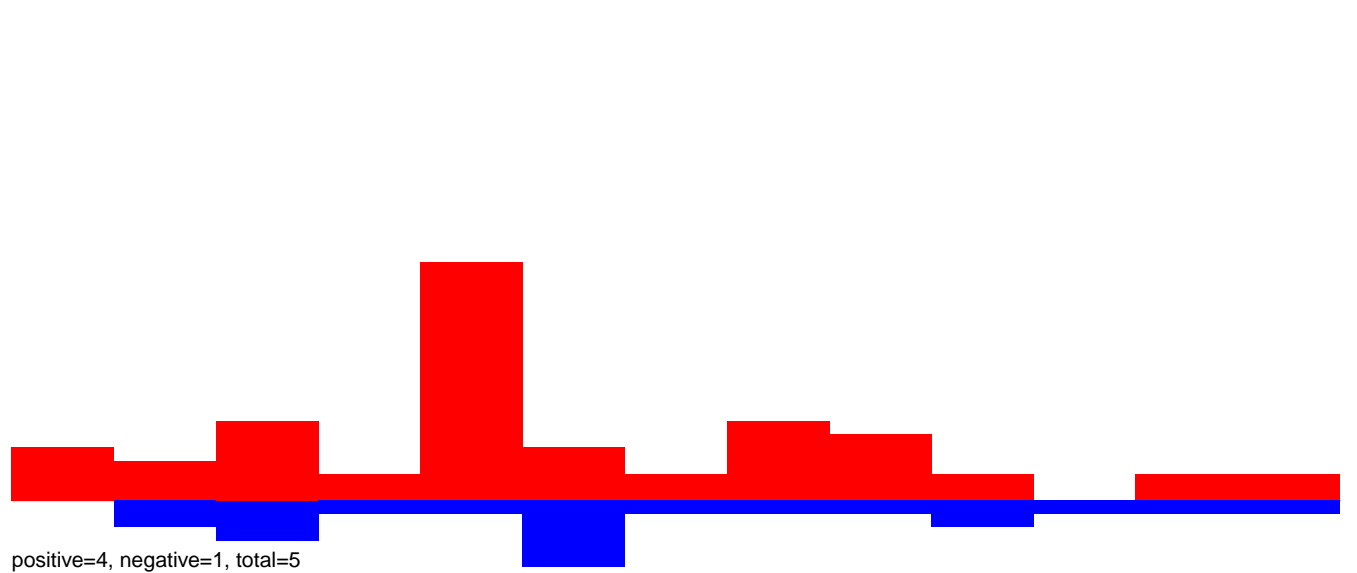
CuQuin_Mcarc_NL.24_35.rep



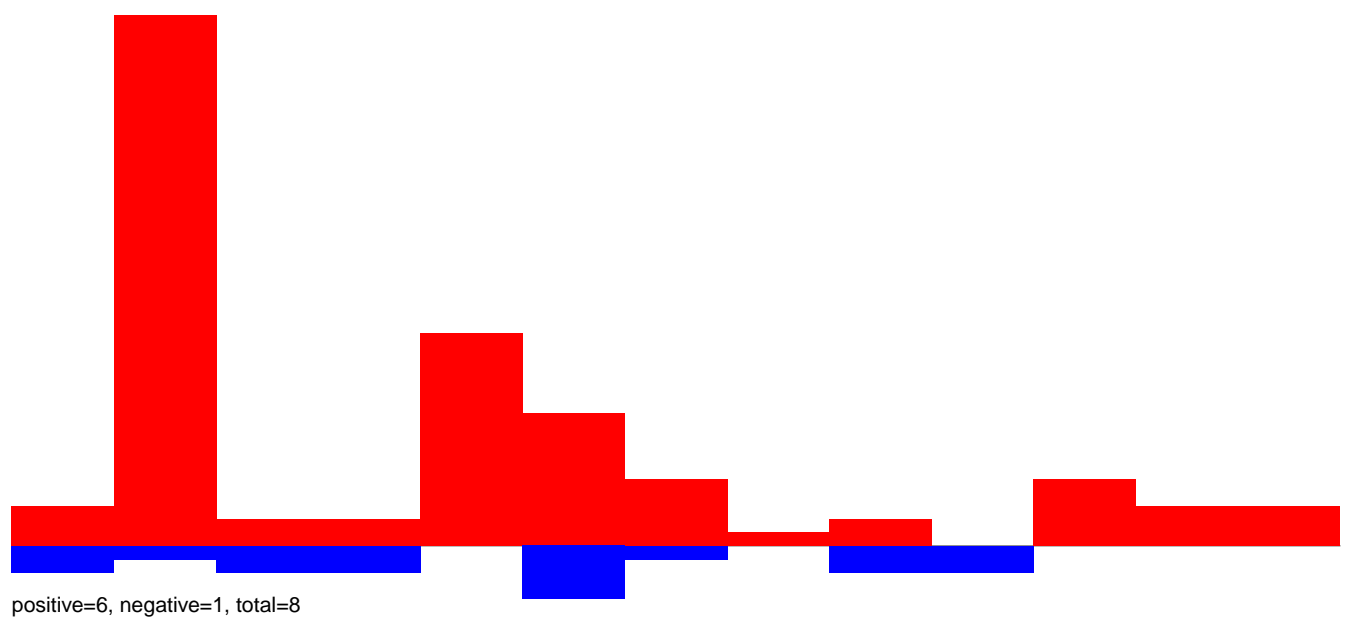
CuQuin_Mcarc_NL.rep



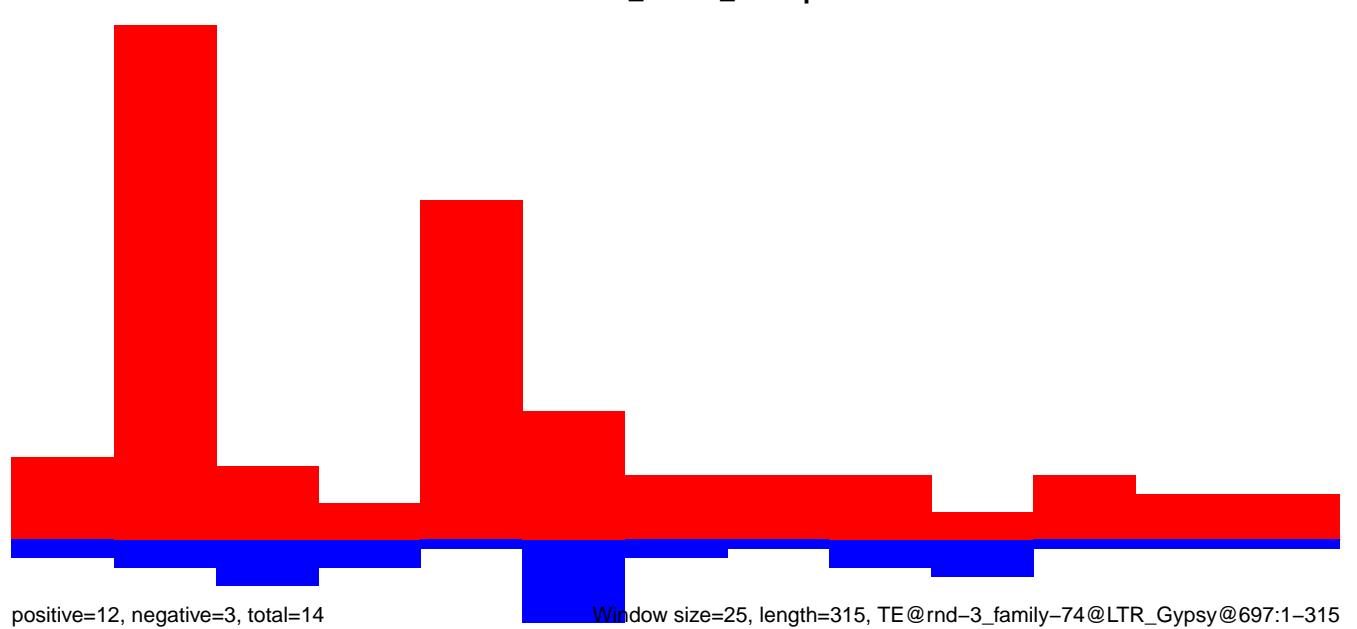
CuQuin_Mcarc_NL.18_23.rep



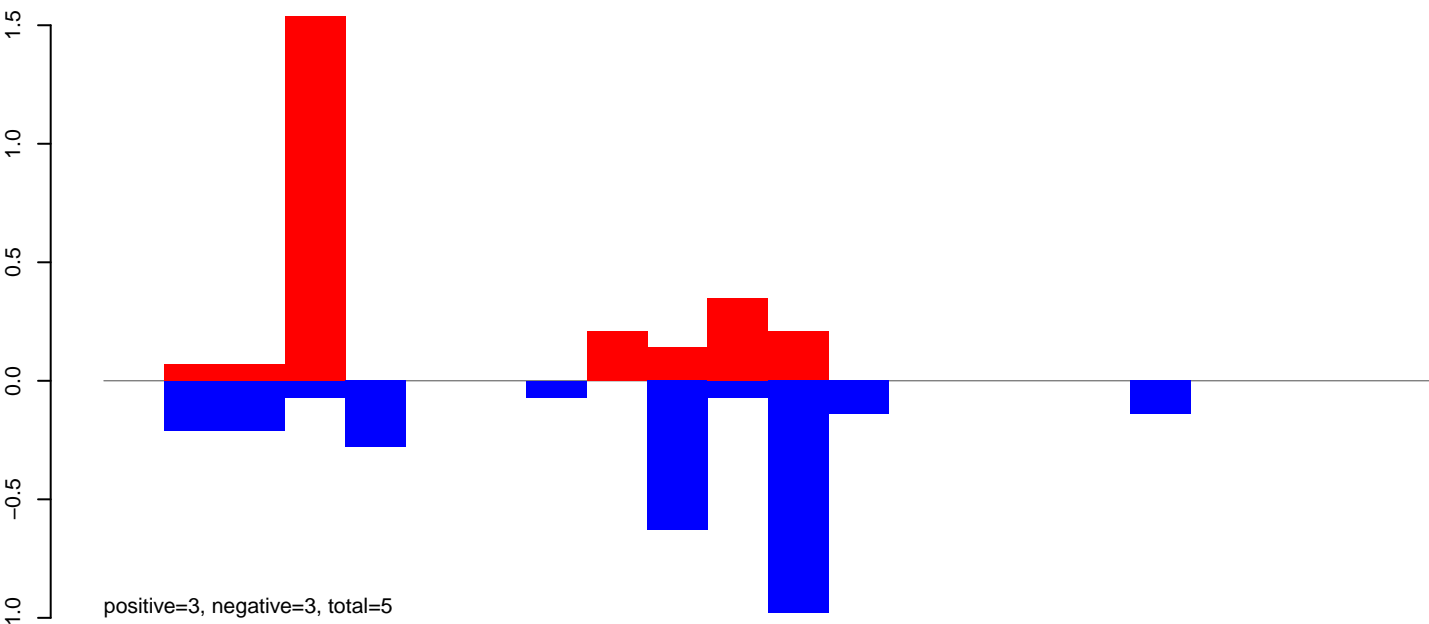
CuQuin_Mcarc_NL.24_35.rep



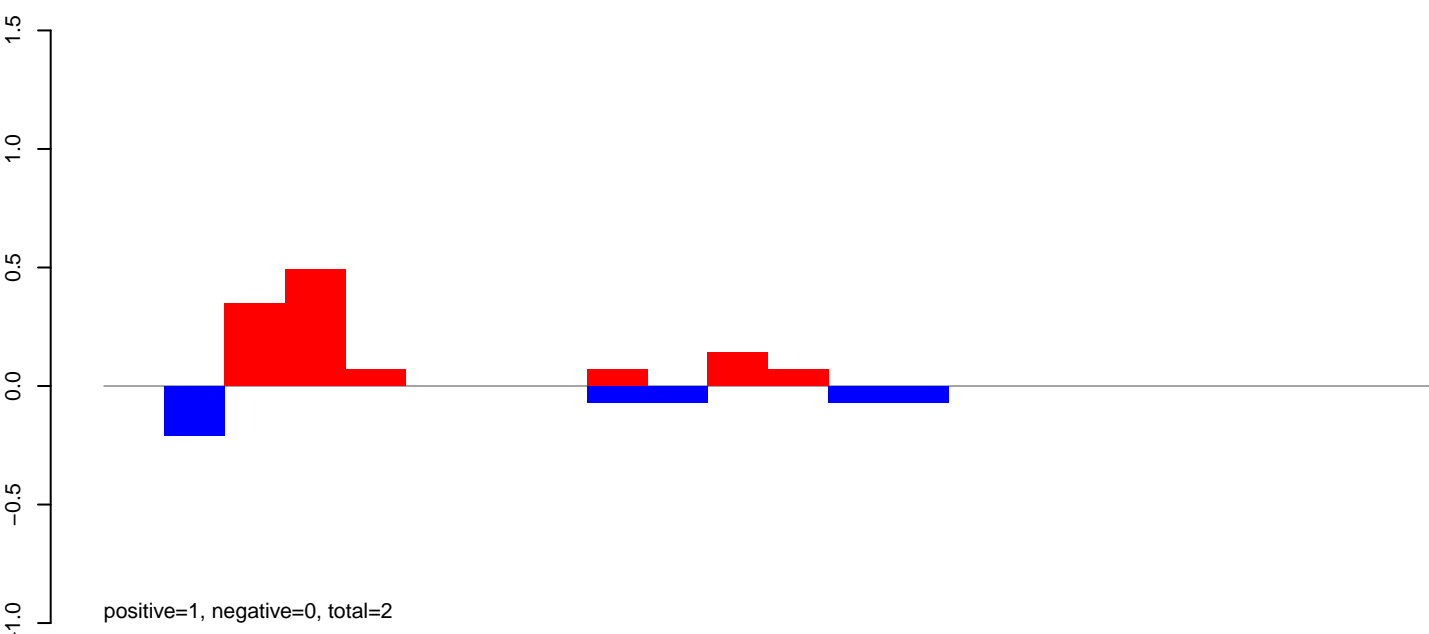
CuQuin_Mcarc_NL.rep



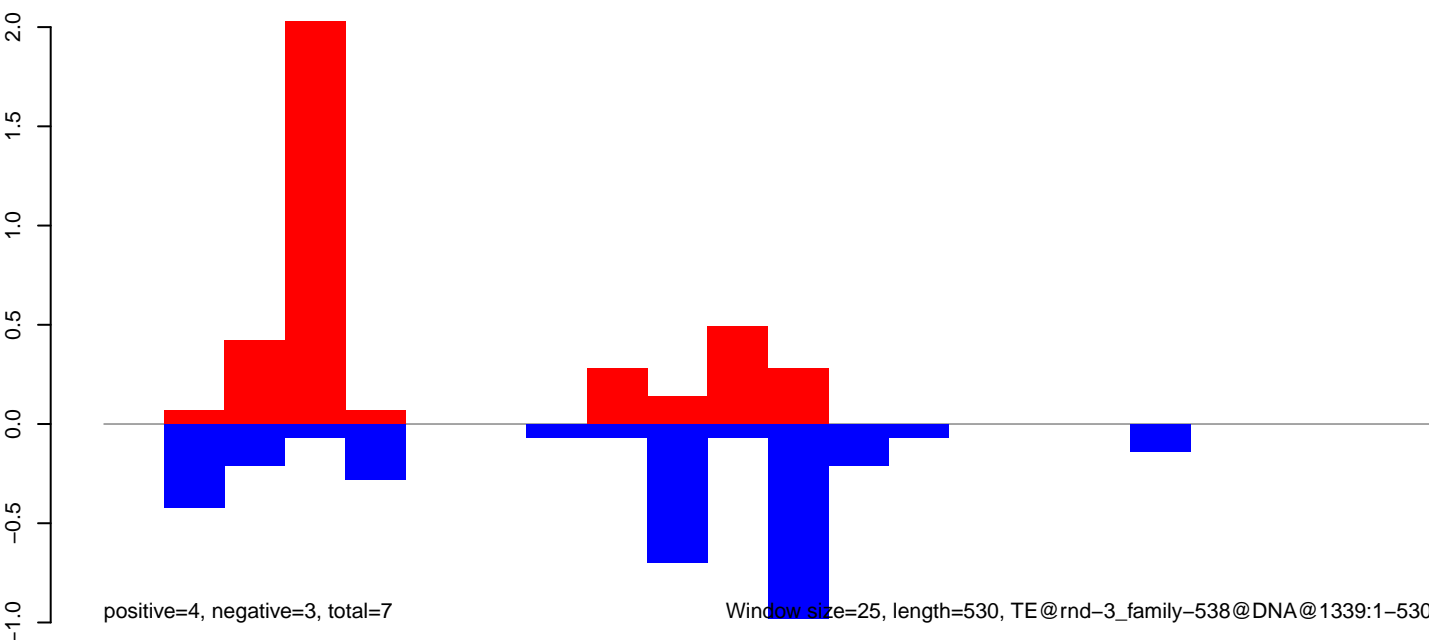
CuQuin_Mcarc_NL.18_23.rep



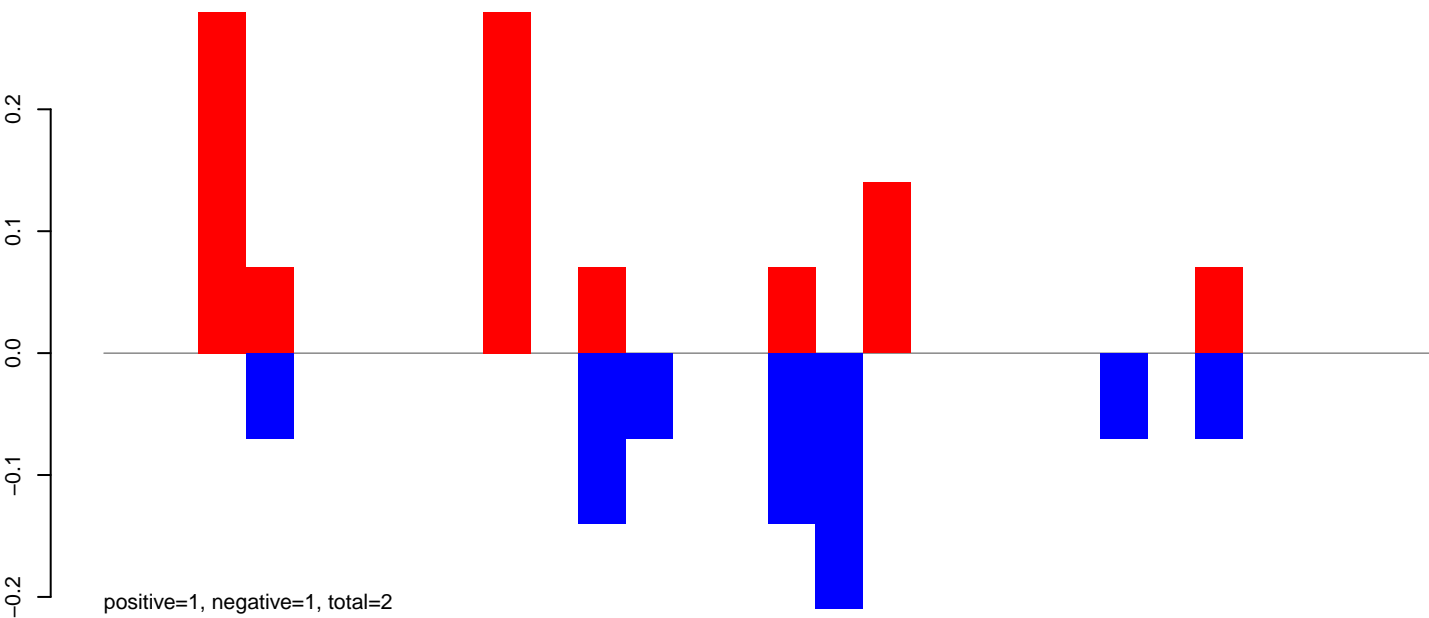
CuQuin_Mcarc_NL.24_35.rep



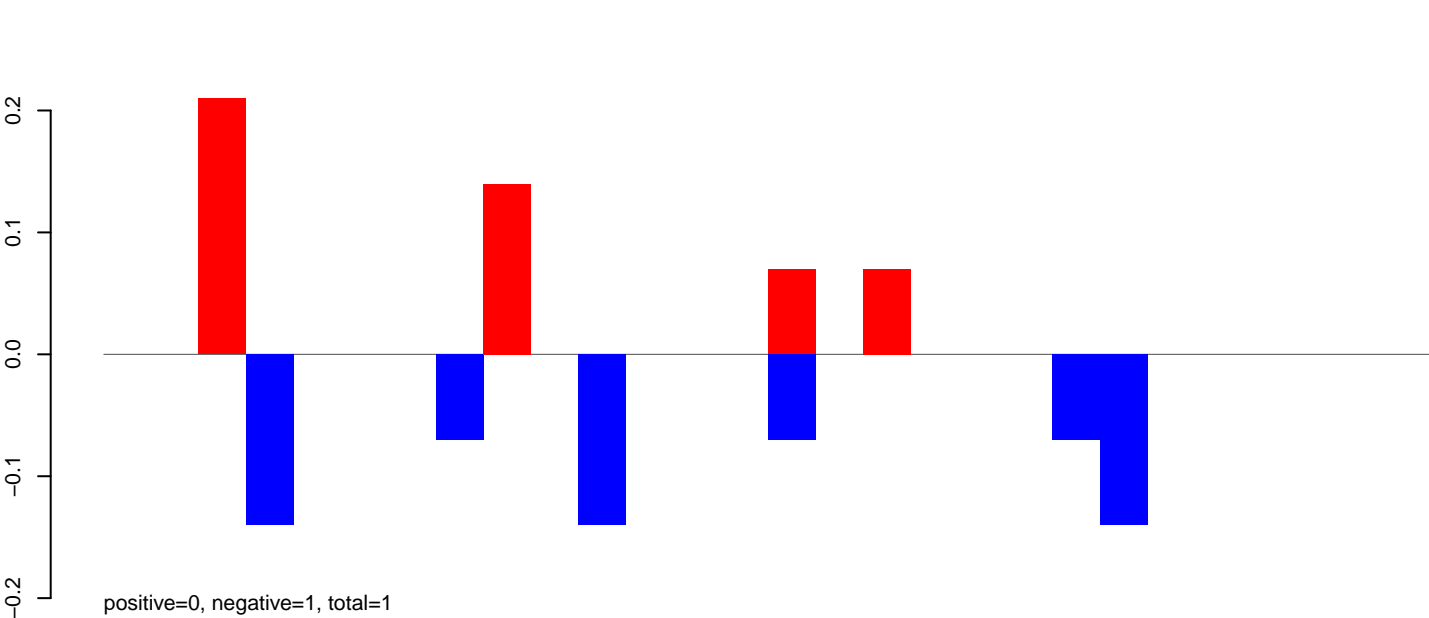
CuQuin_Mcarc_NL.rep



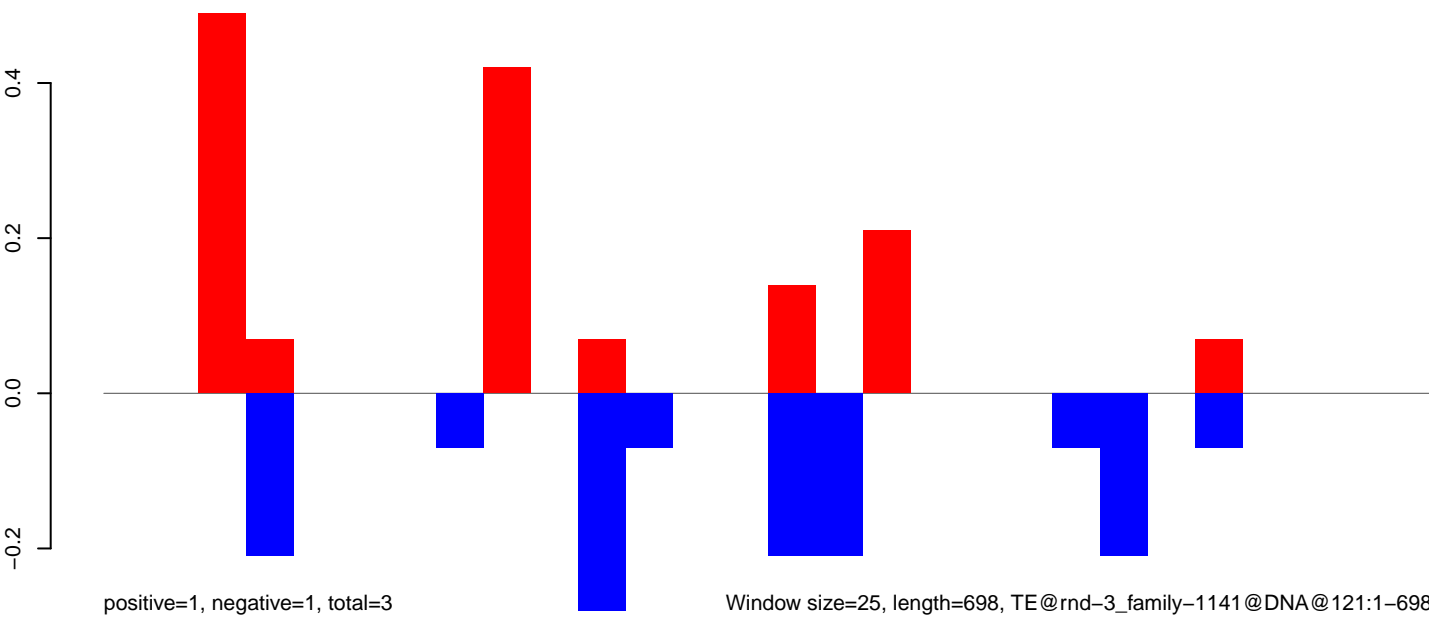
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

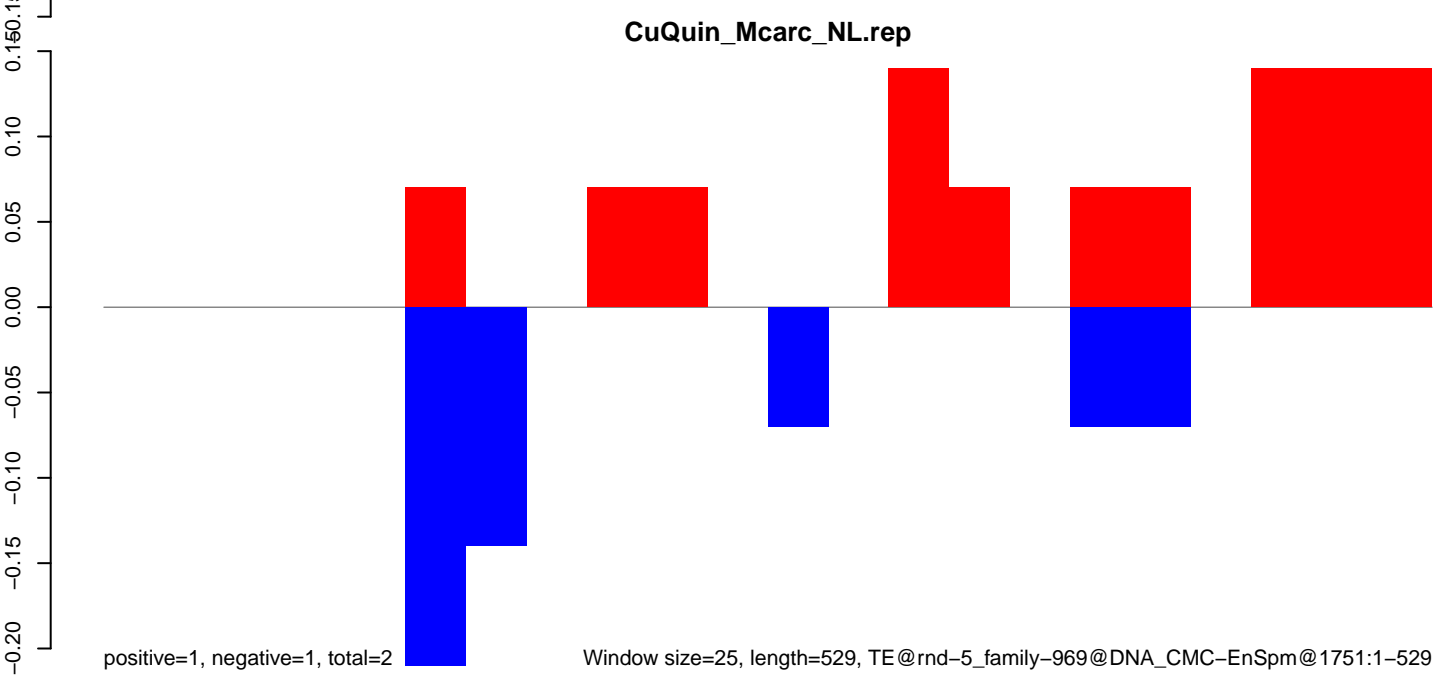
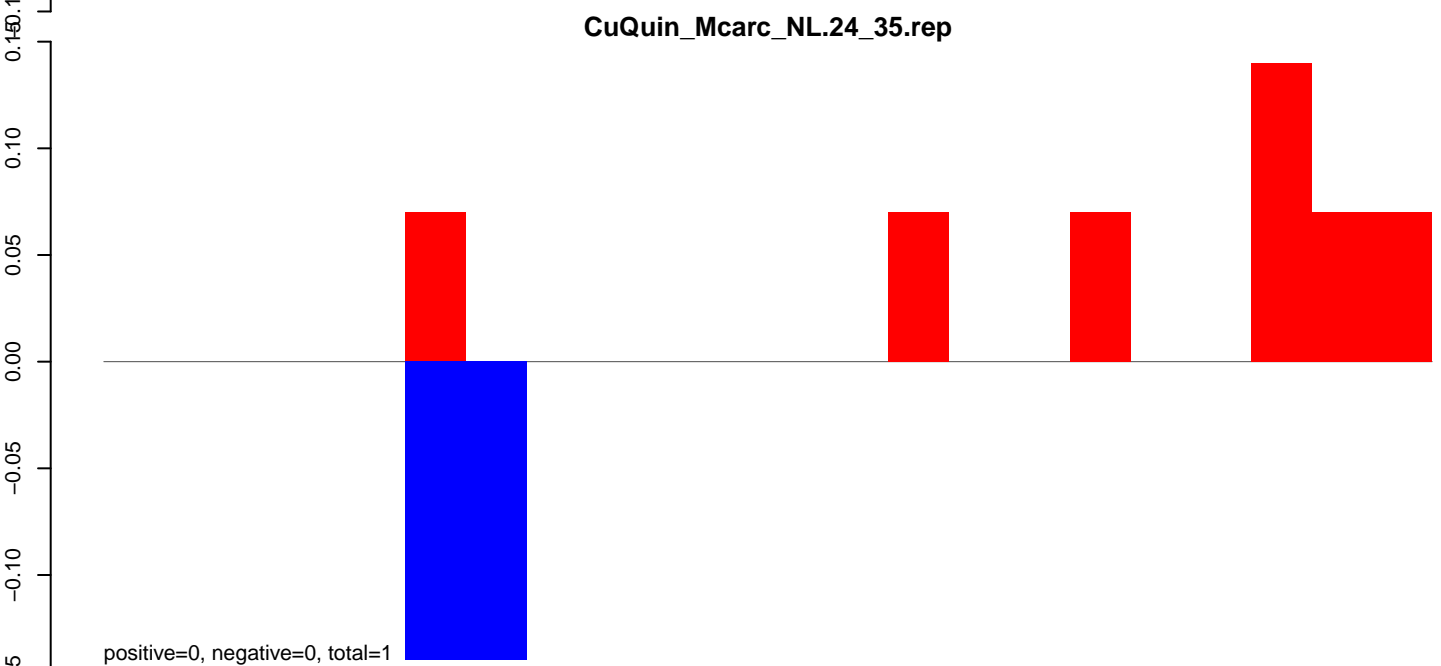
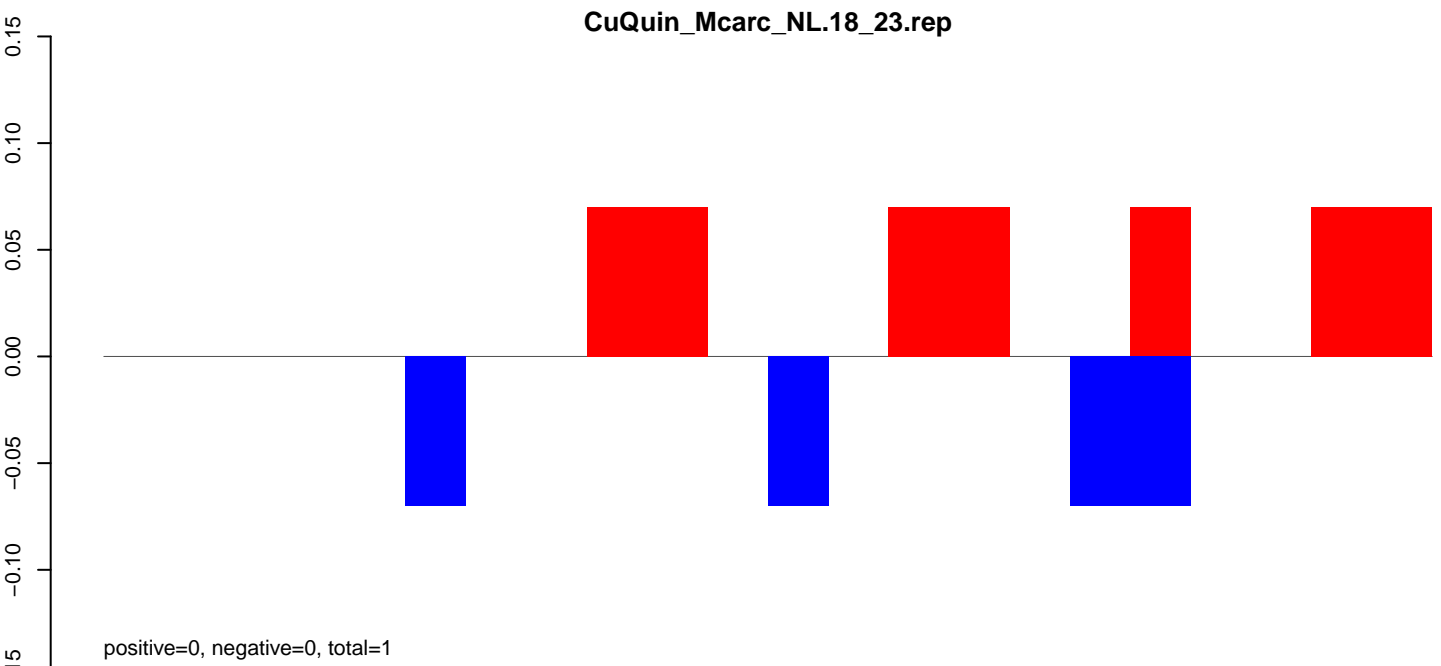


CuQuin_Mcarc_NL.rep

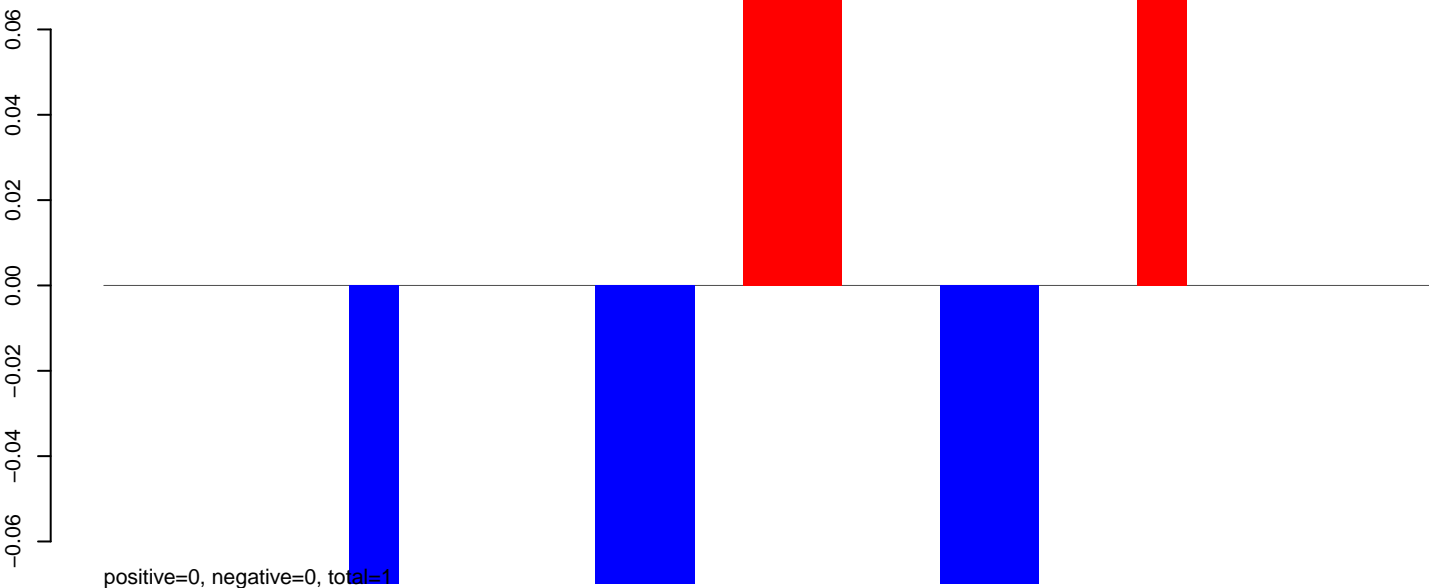


positive=1, negative=1, total=3

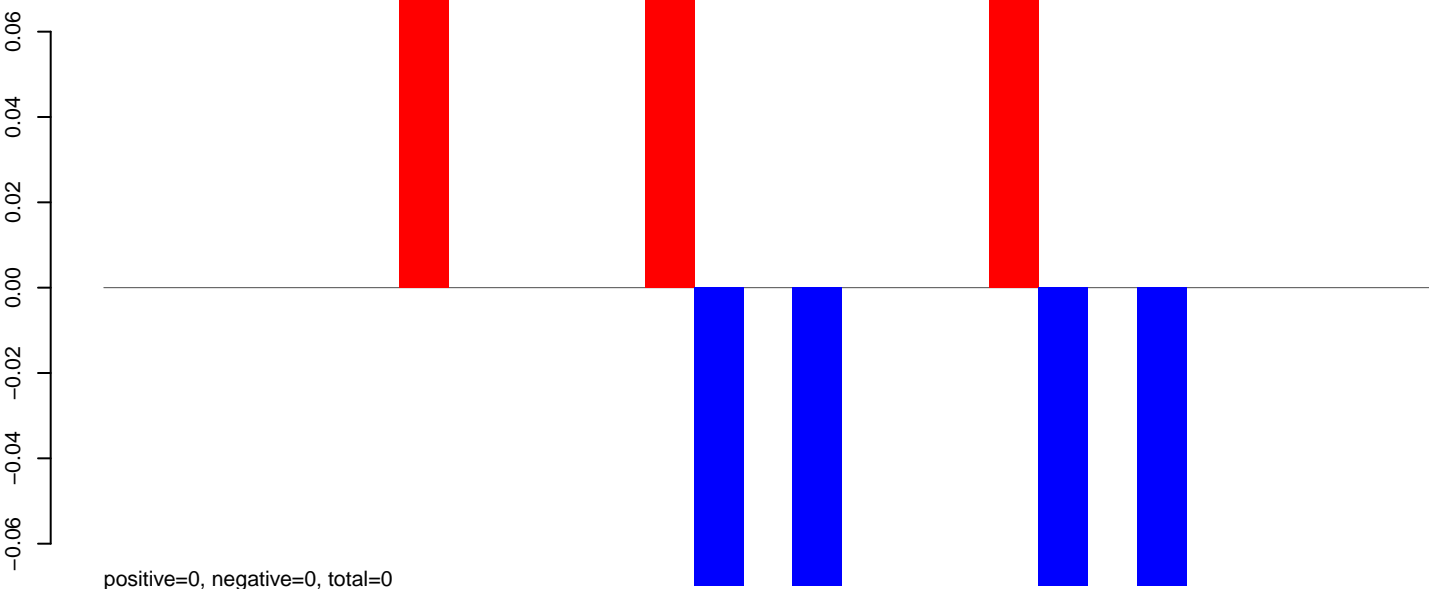
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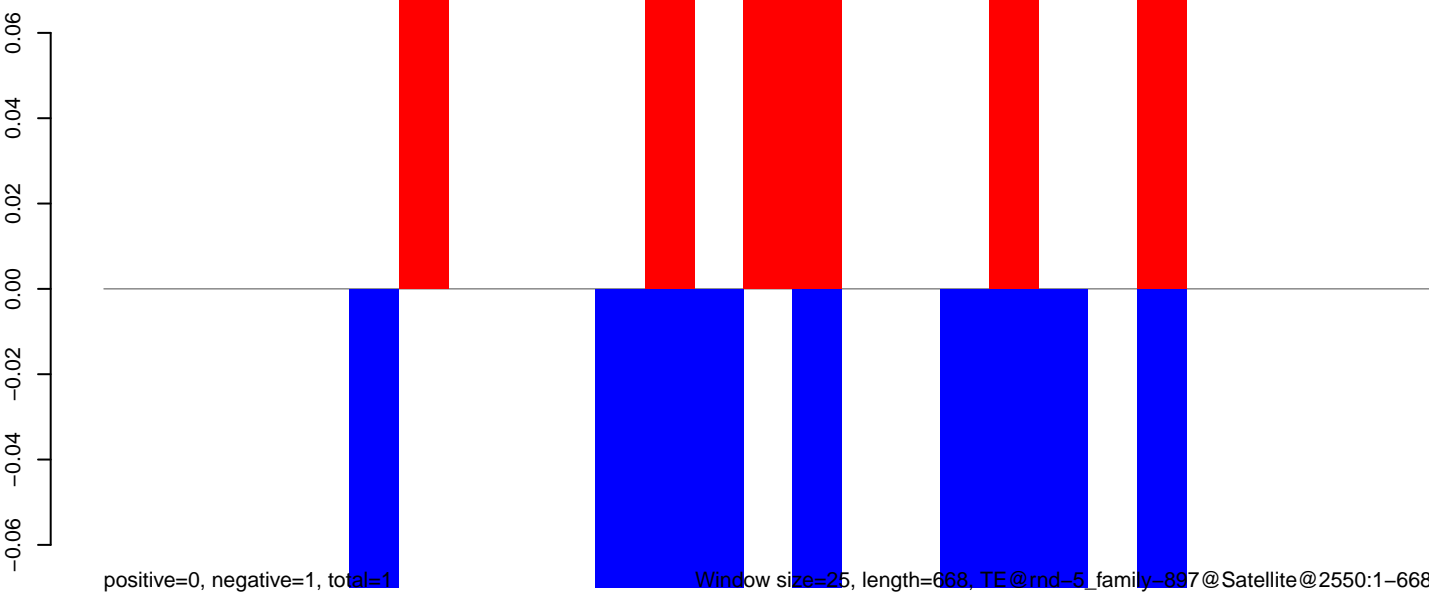
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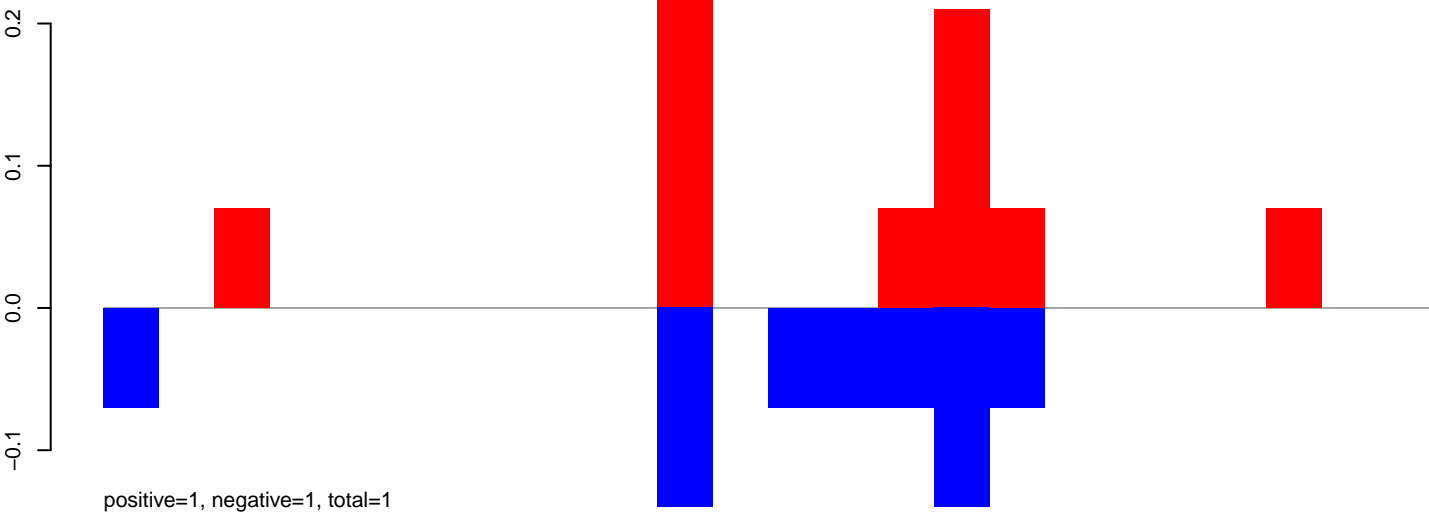
CuQuin_Mcarc_NL.24_35.rep



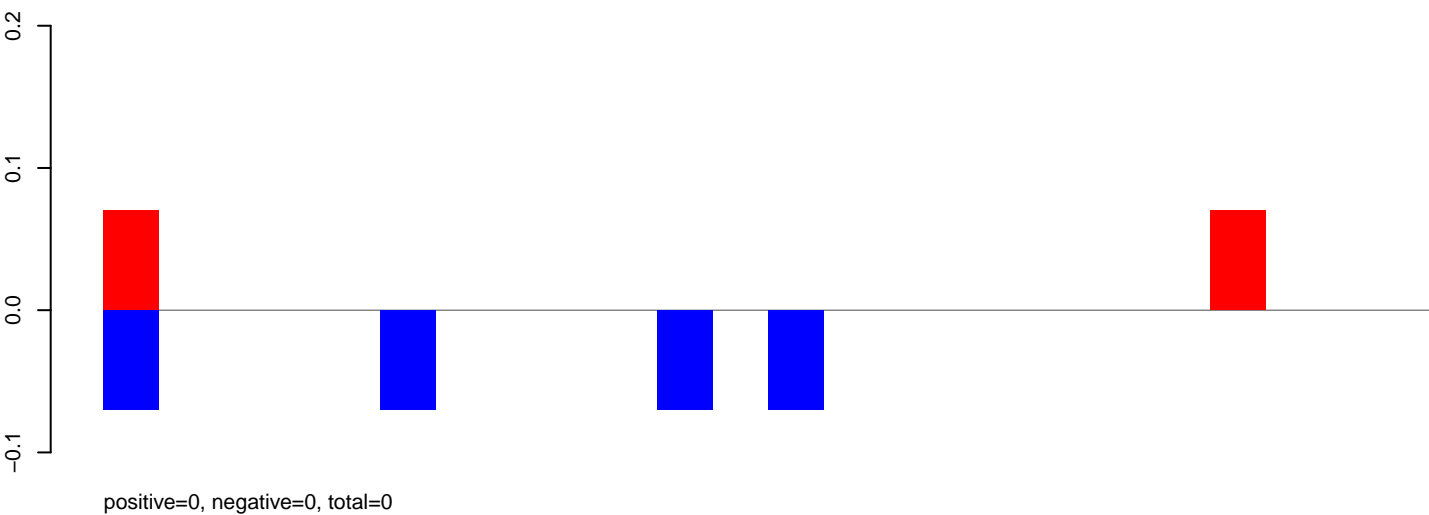
CuQuin_Mcarc_NL.rep



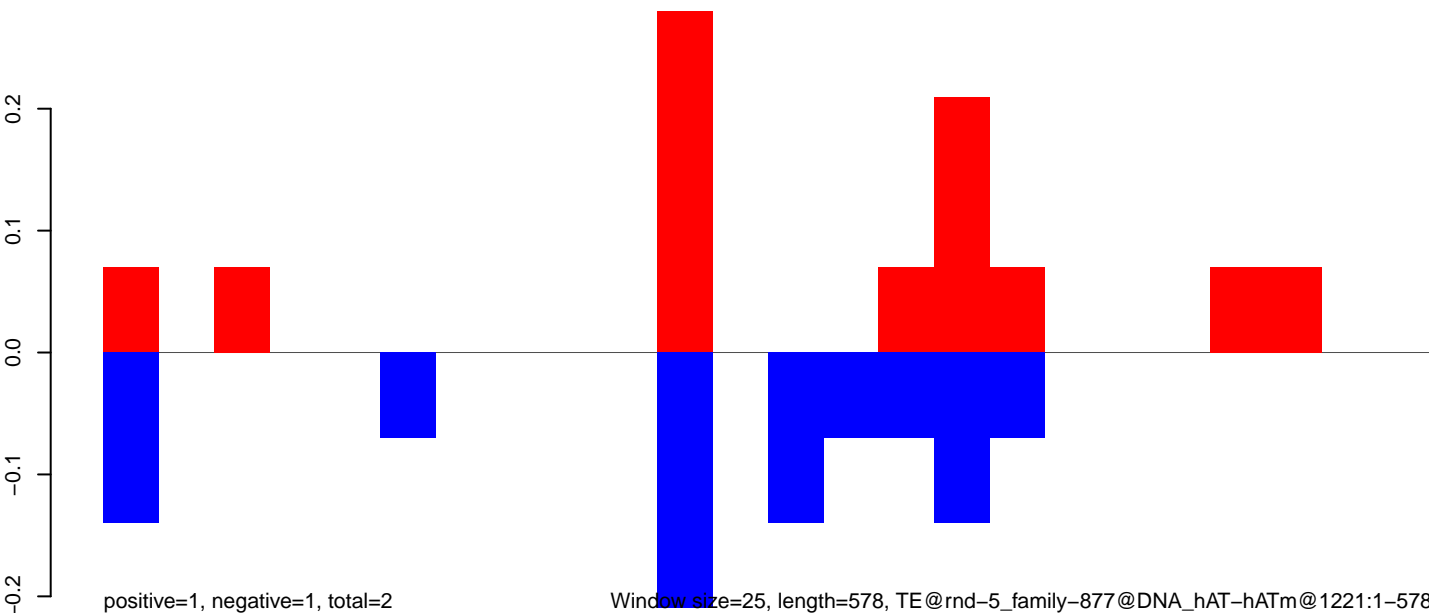
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CuQuin_Mcarc_NL.24_35.rep

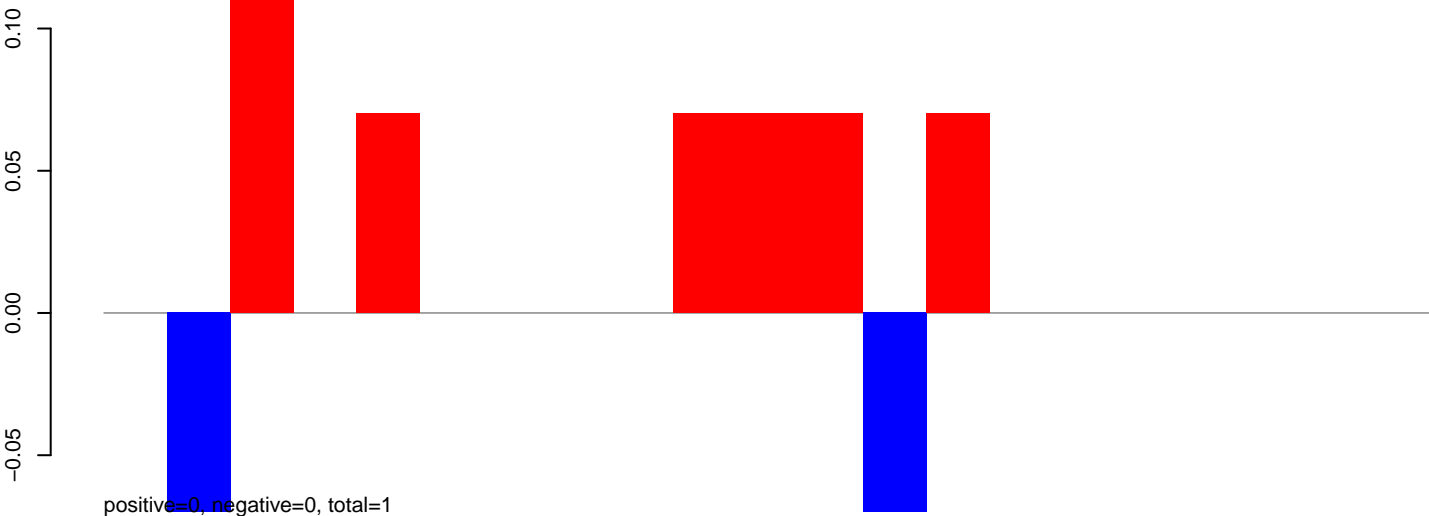


CuQuin_Mcarc_NL.rep

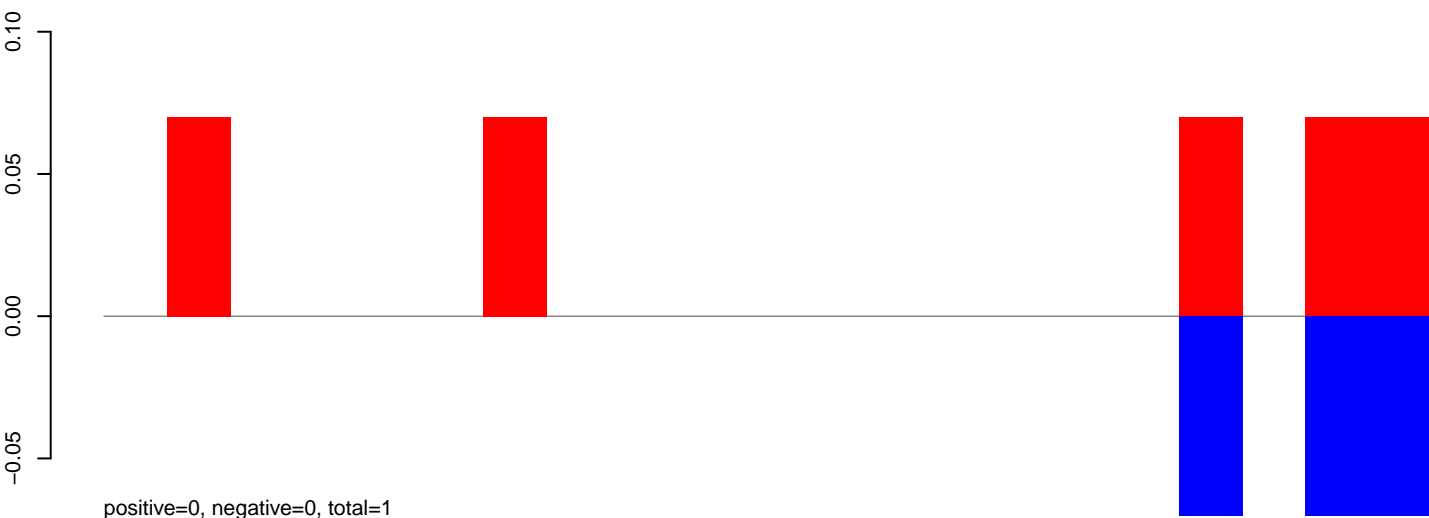


Window size=25, length=578, TE@rnd-5_family-877@DNA_hAT-hATm@1221:1-578

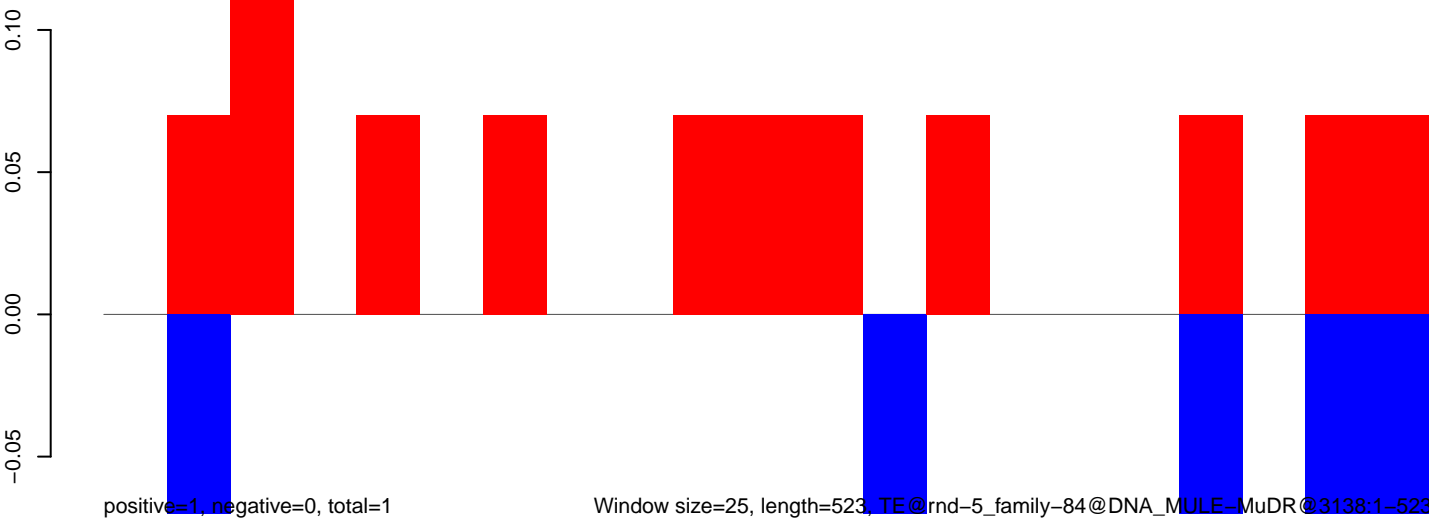
CuQuin_Mcarc_NL.18_23.rep

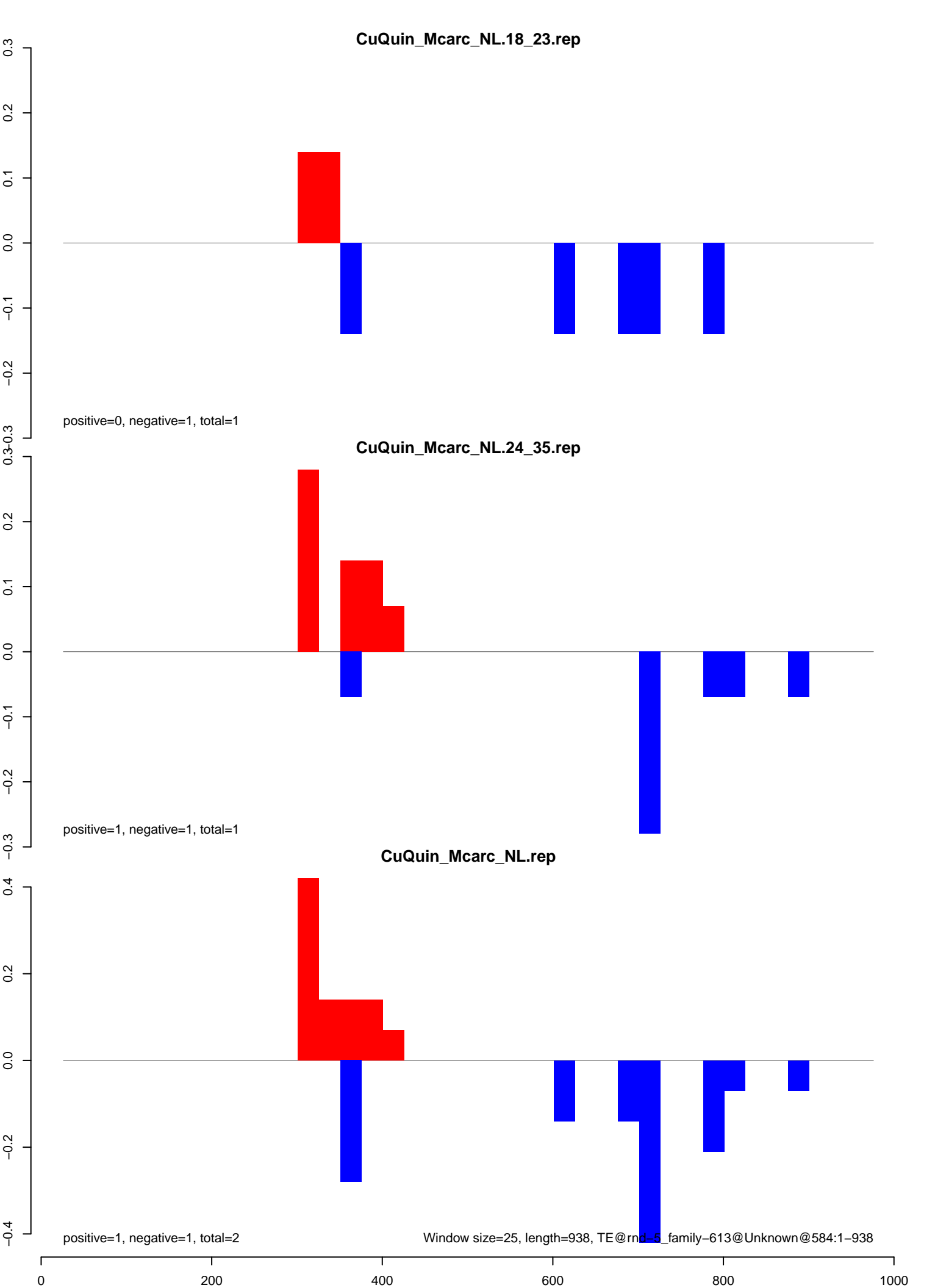


CuQuin_Mcarc_NL.24_35.rep

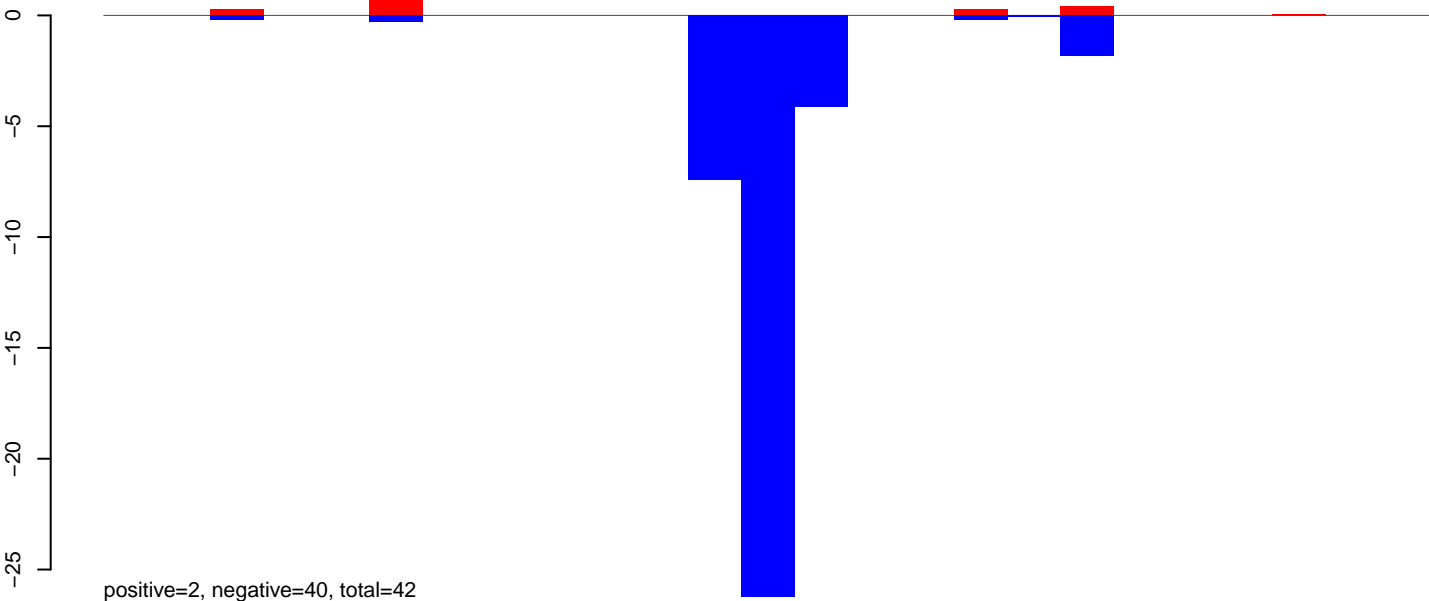


CuQuin_Mcarc_NL.rep

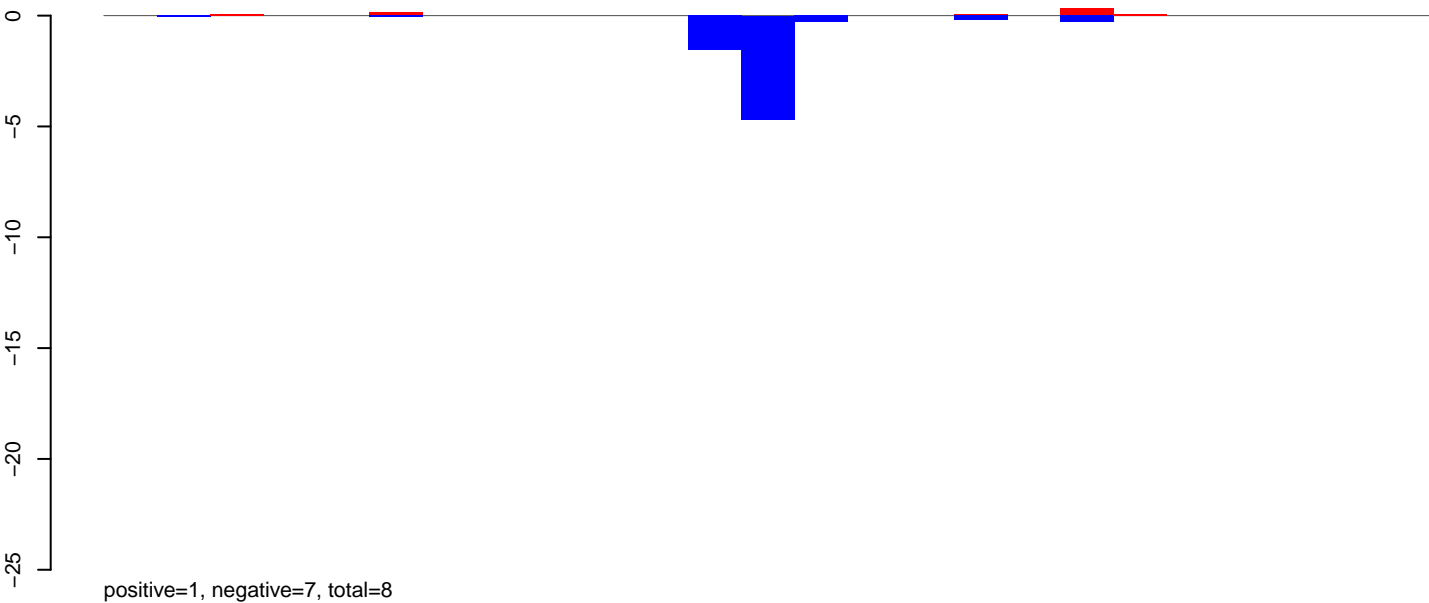




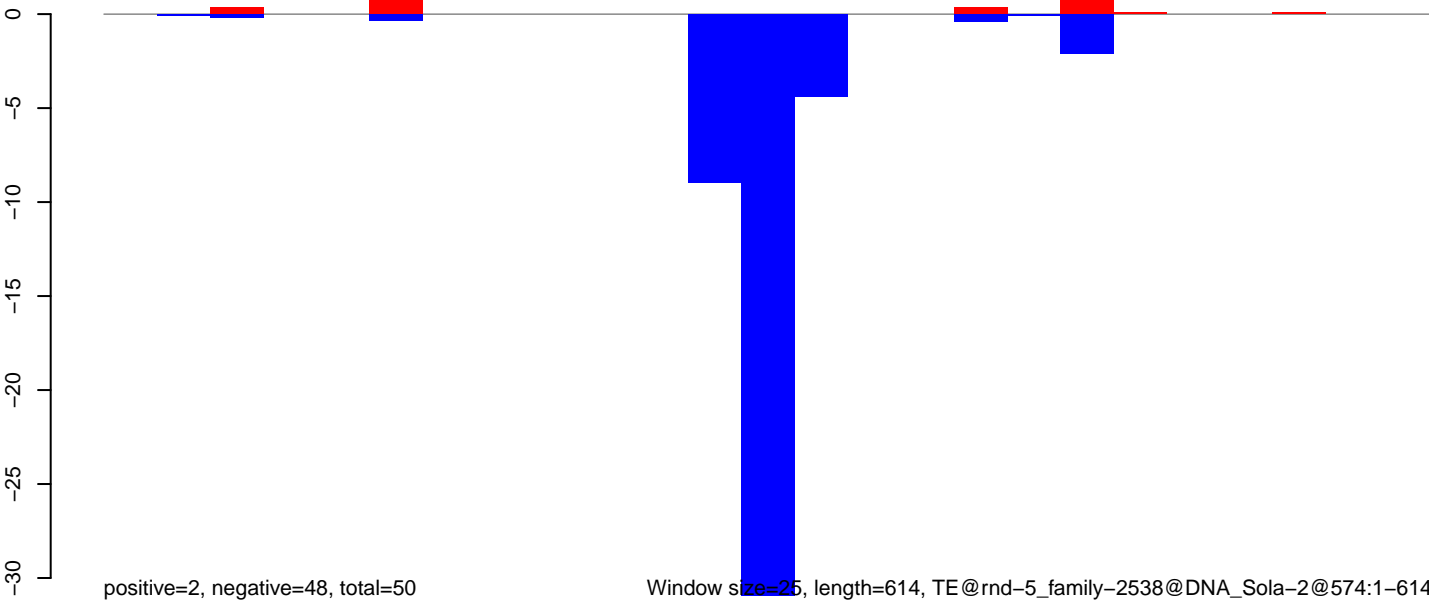
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

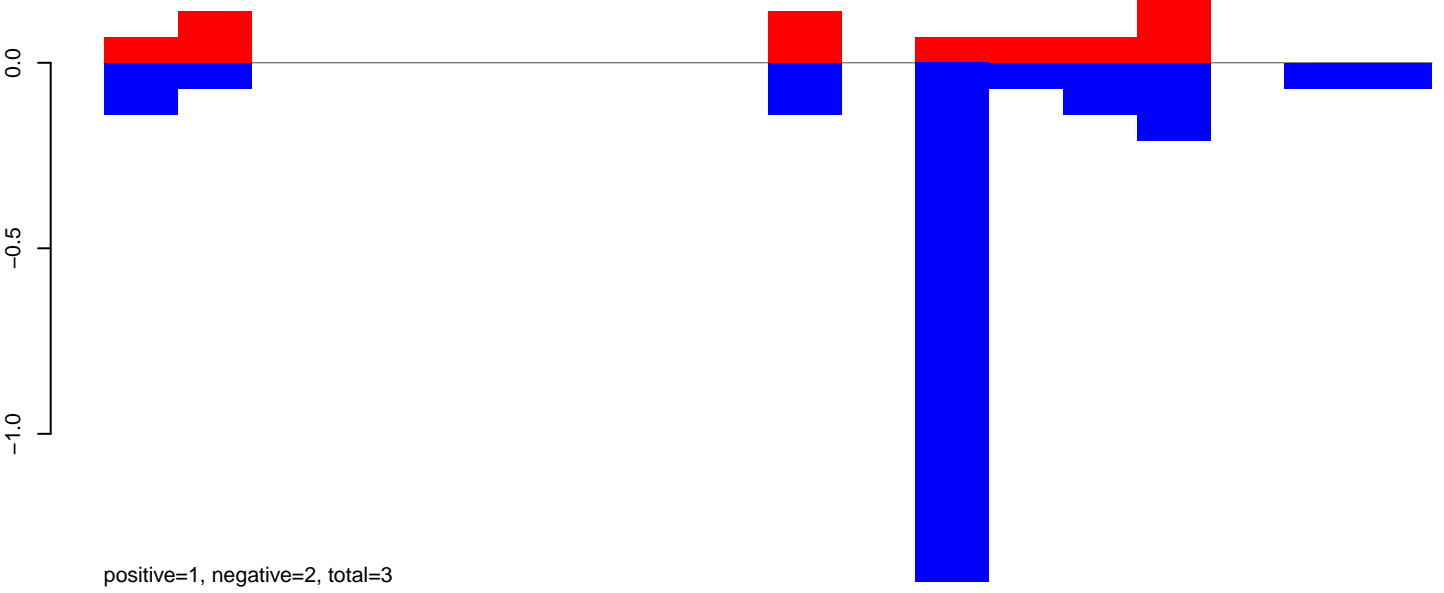


CuQuin_Mcarc_NL.rep

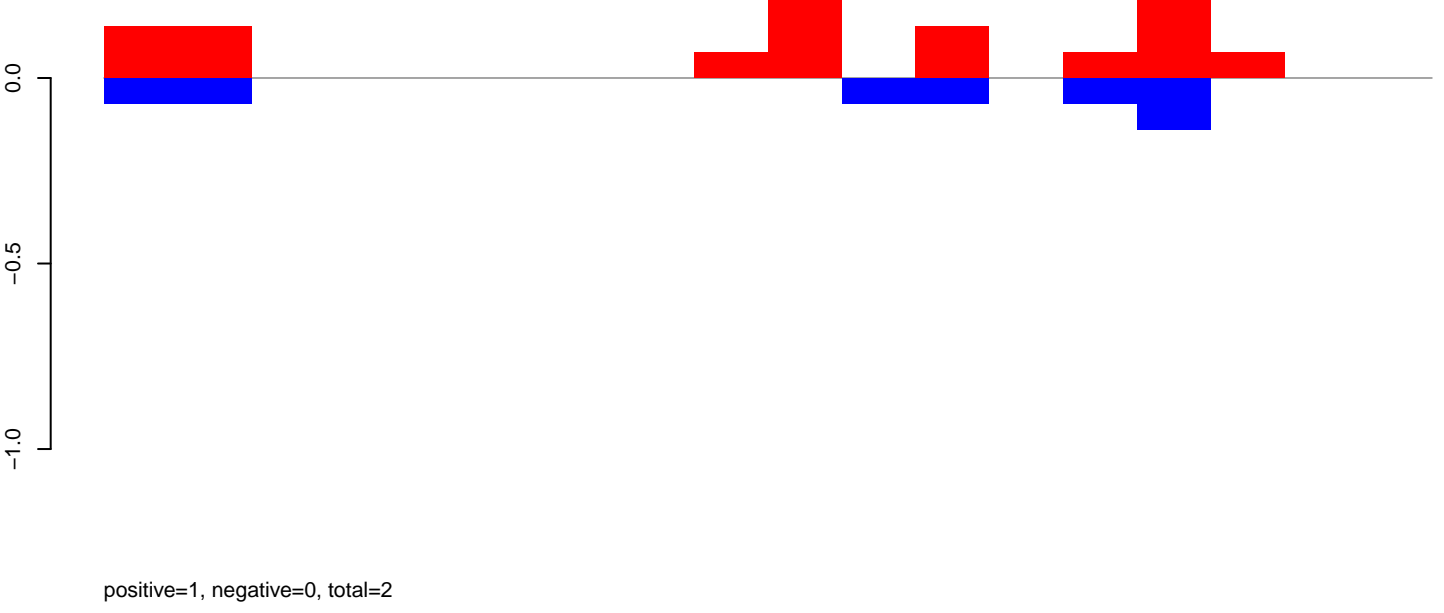


Window size=25, length=614, TE@rnd-5_family-2538@DNA_Sola-2@574:1-614

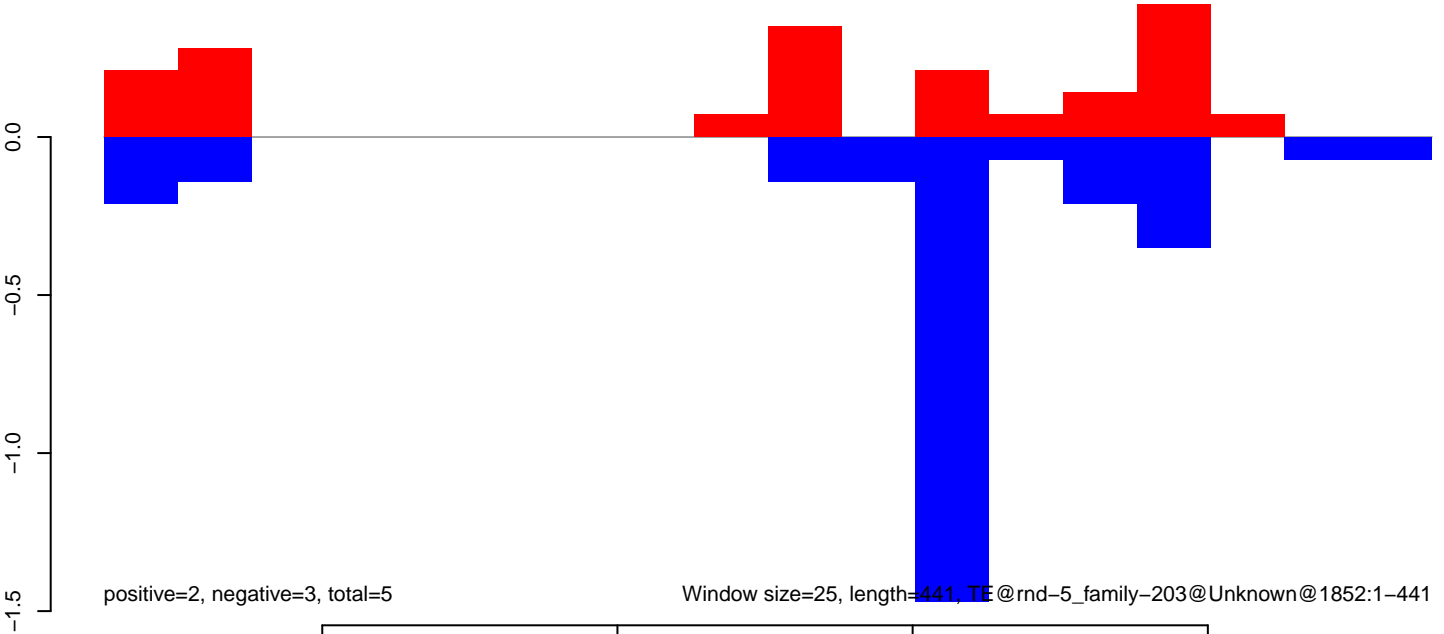
CuQuin_Mcarc_NL.18_23.rep



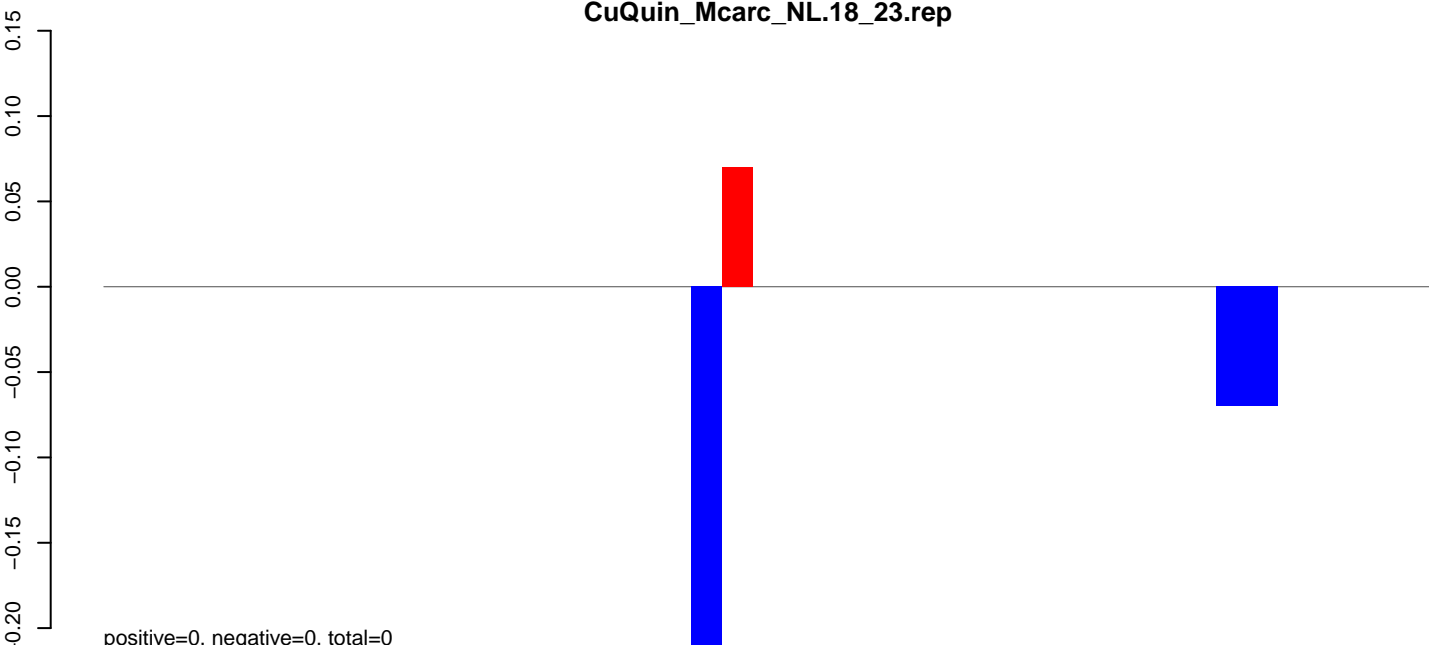
CuQuin_Mcarc_NL.24_35.rep



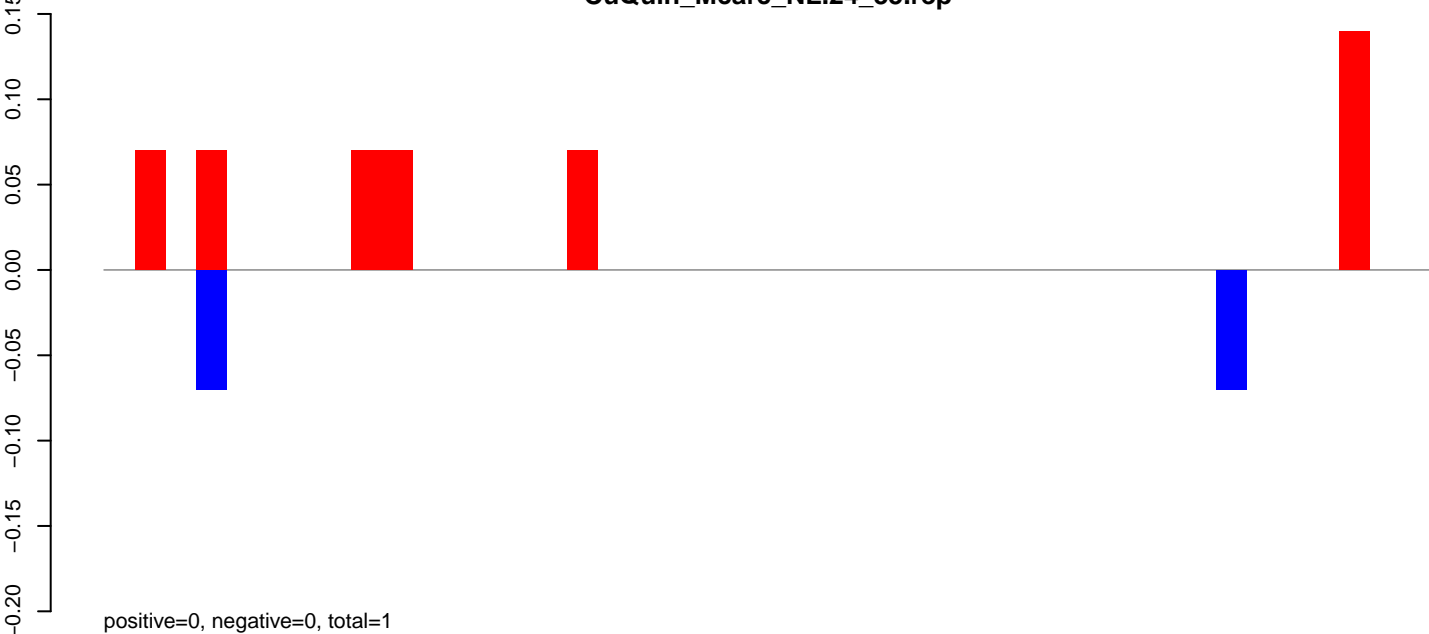
CuQuin_Mcarc_NL.rep



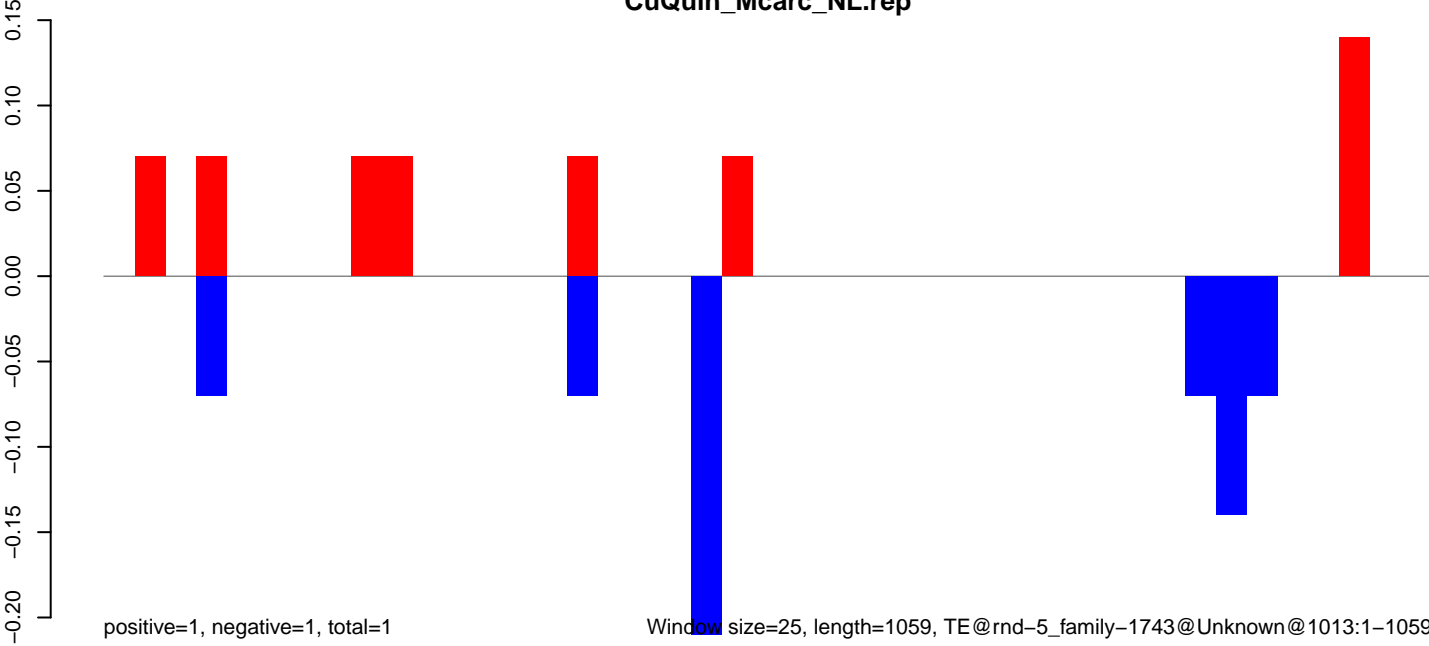
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

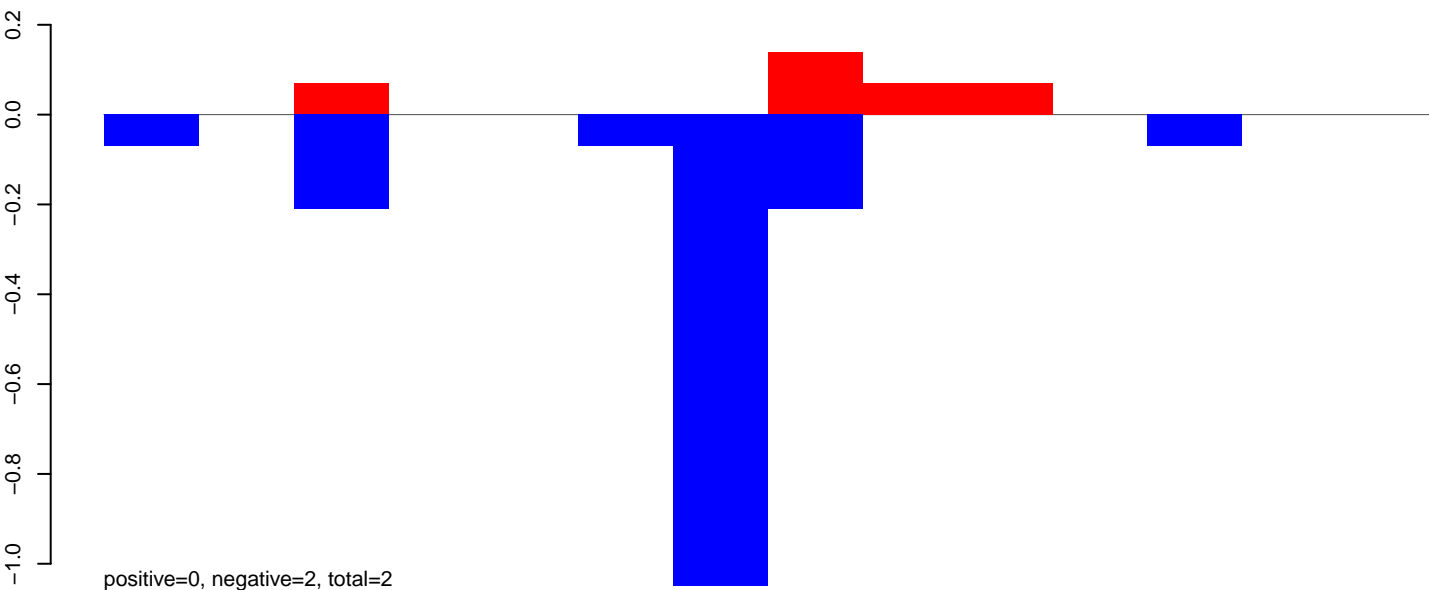


CuQuin_Mcarc_NL.rep

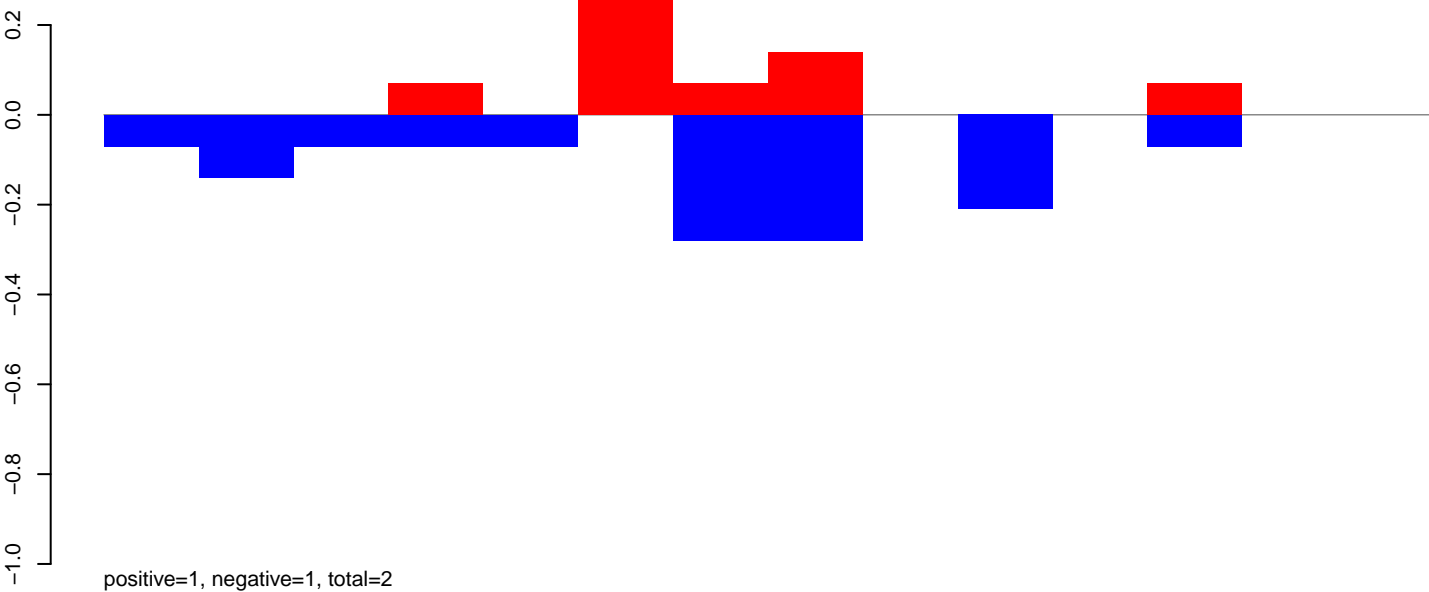


Window size=25, length=1059, TE@rnd-5_family-1743@Unknown@1013:1-1059

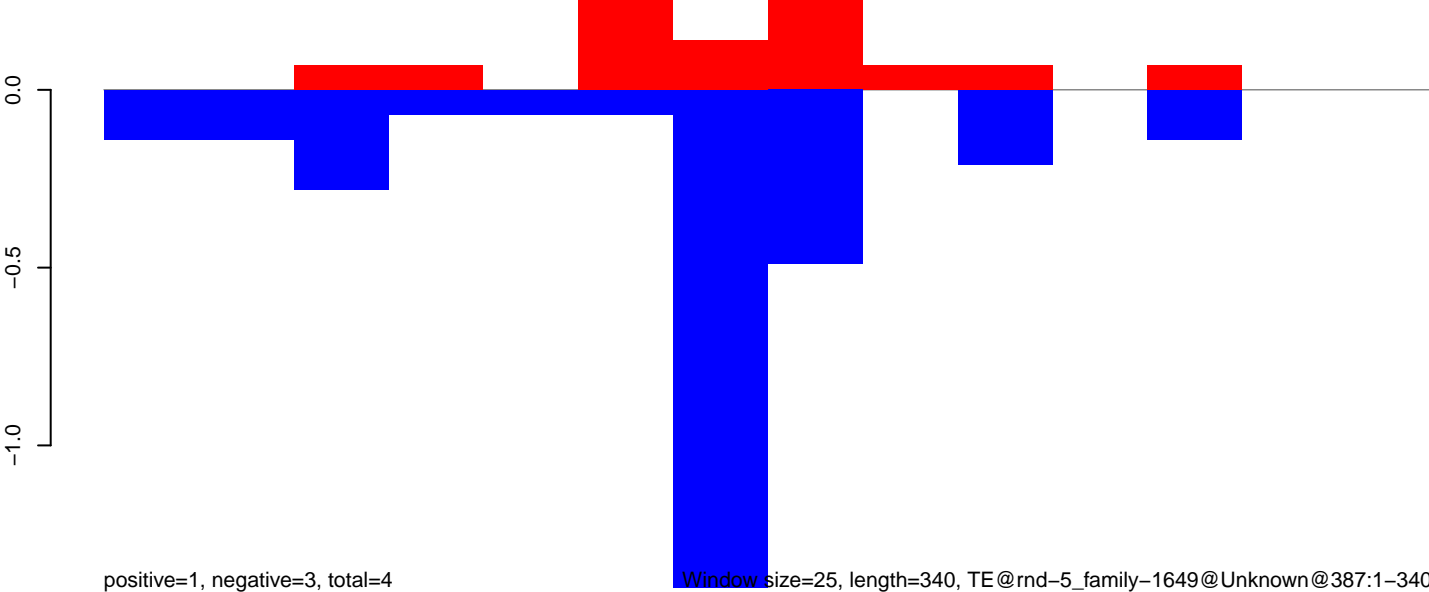
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

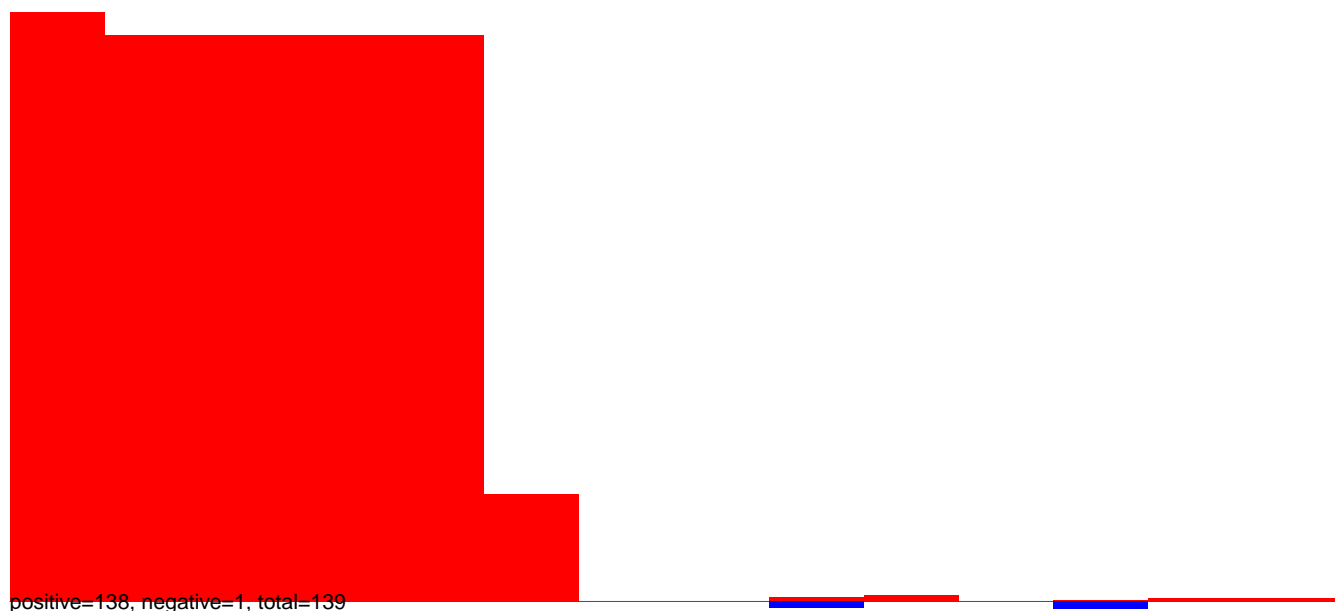


CuQuin_Mcarc_NL.rep

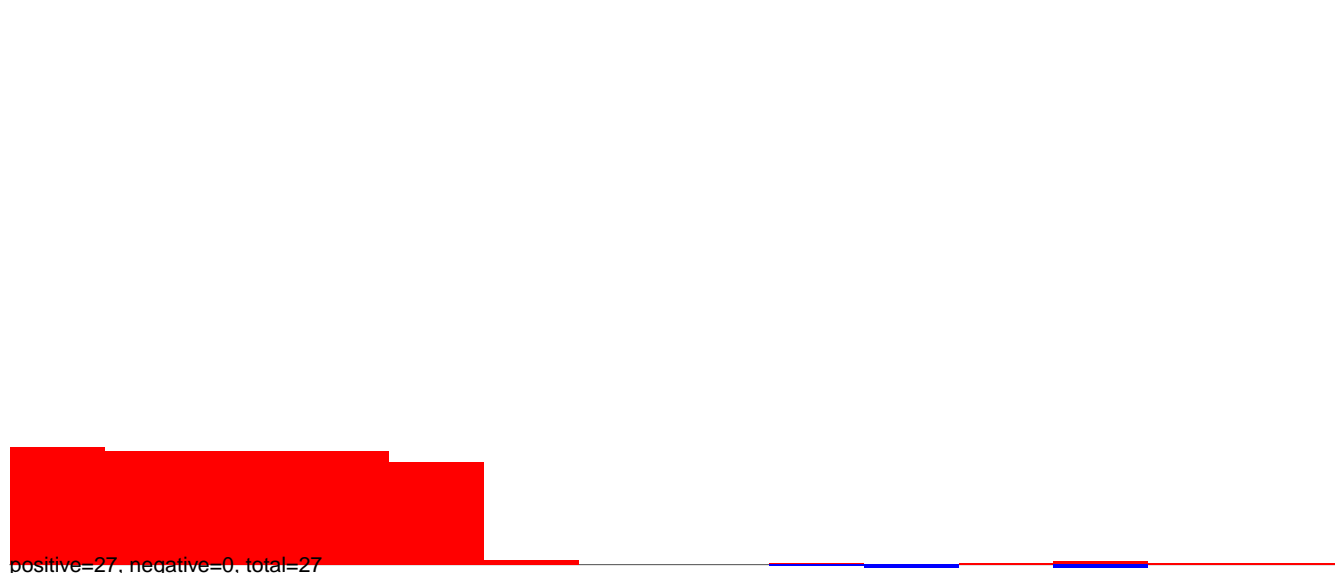


Window size=25, length=340, TE@rnd-5_family-1649@Unknown@387:1-340

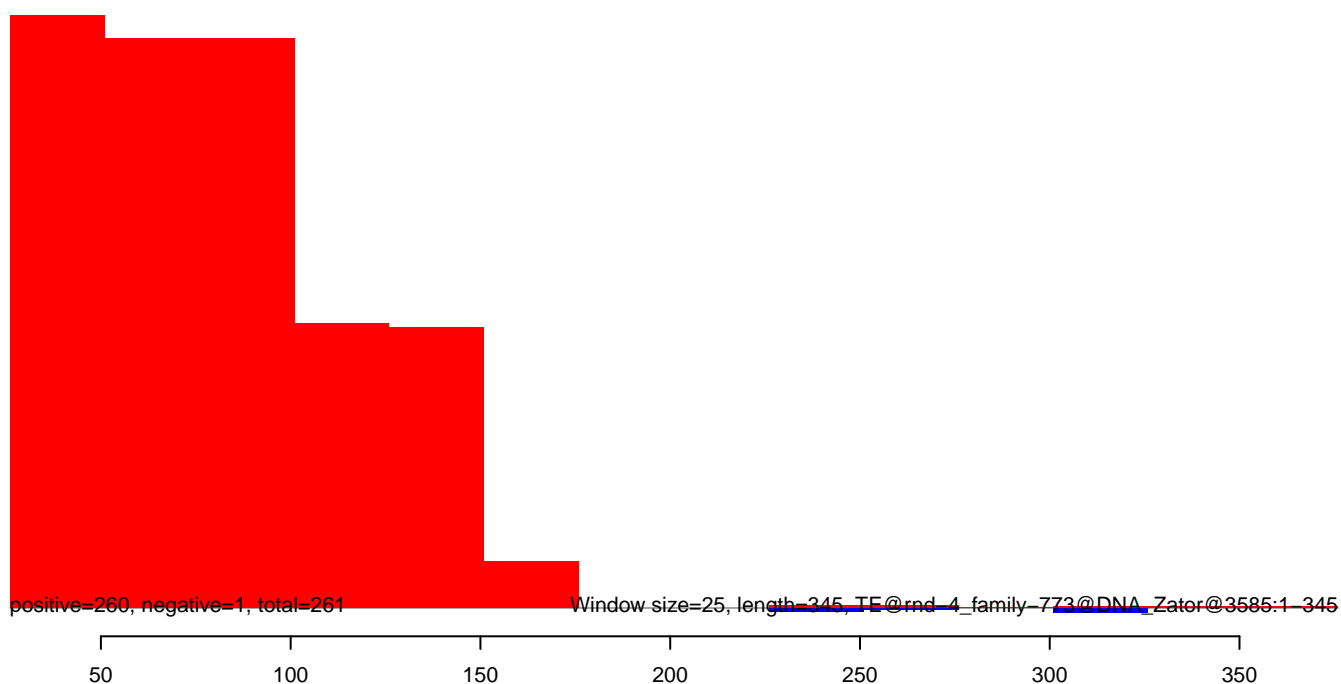
CuQuin_Mcarc_NL.18_23.rep



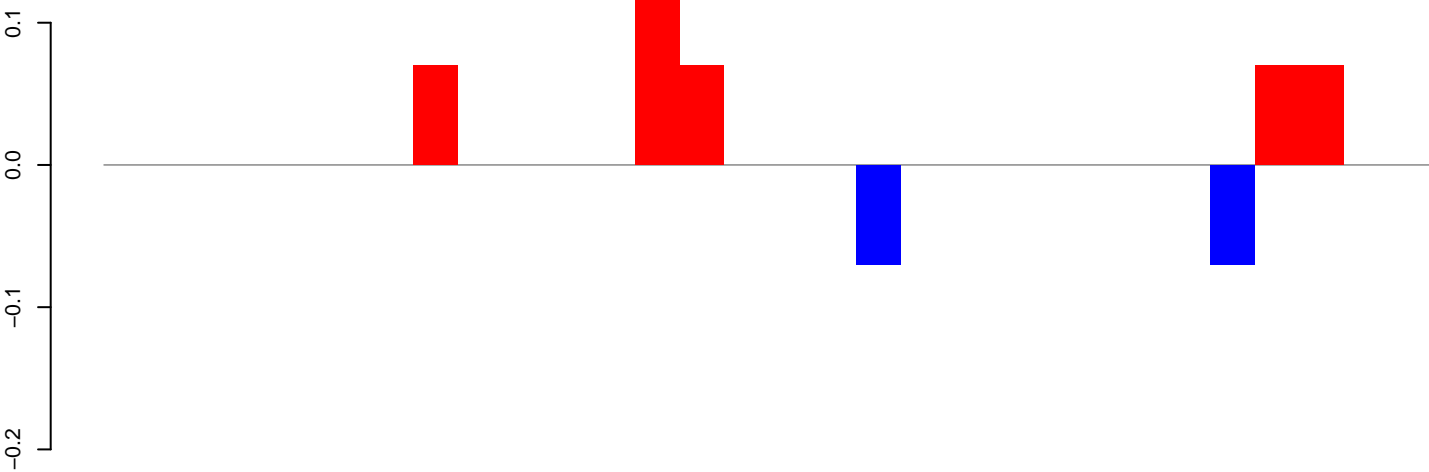
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

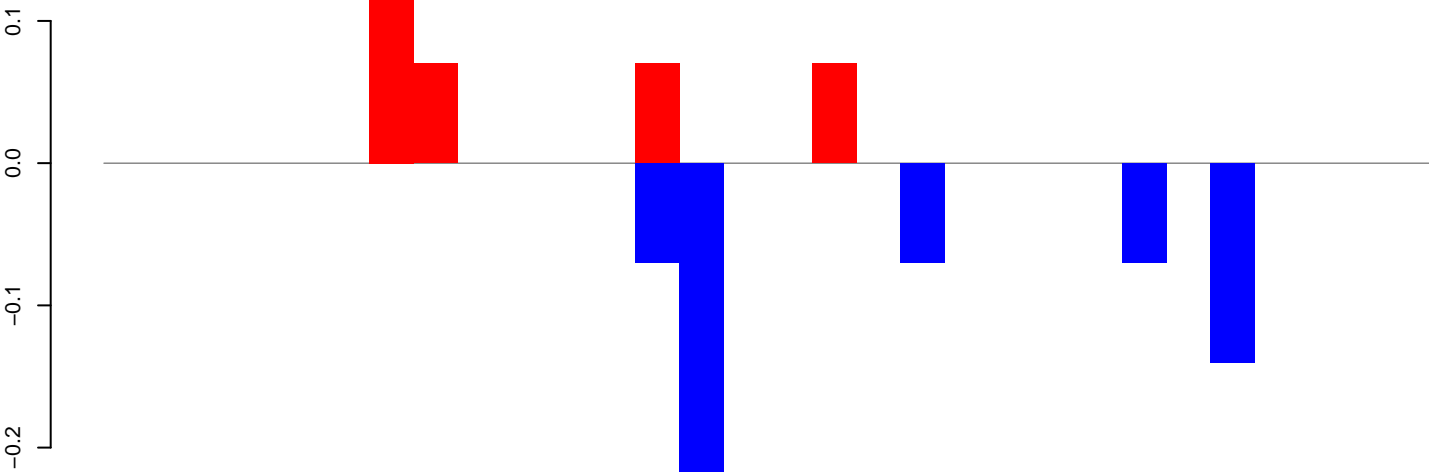


CuQuin_Mcarc_NL.18_23.rep



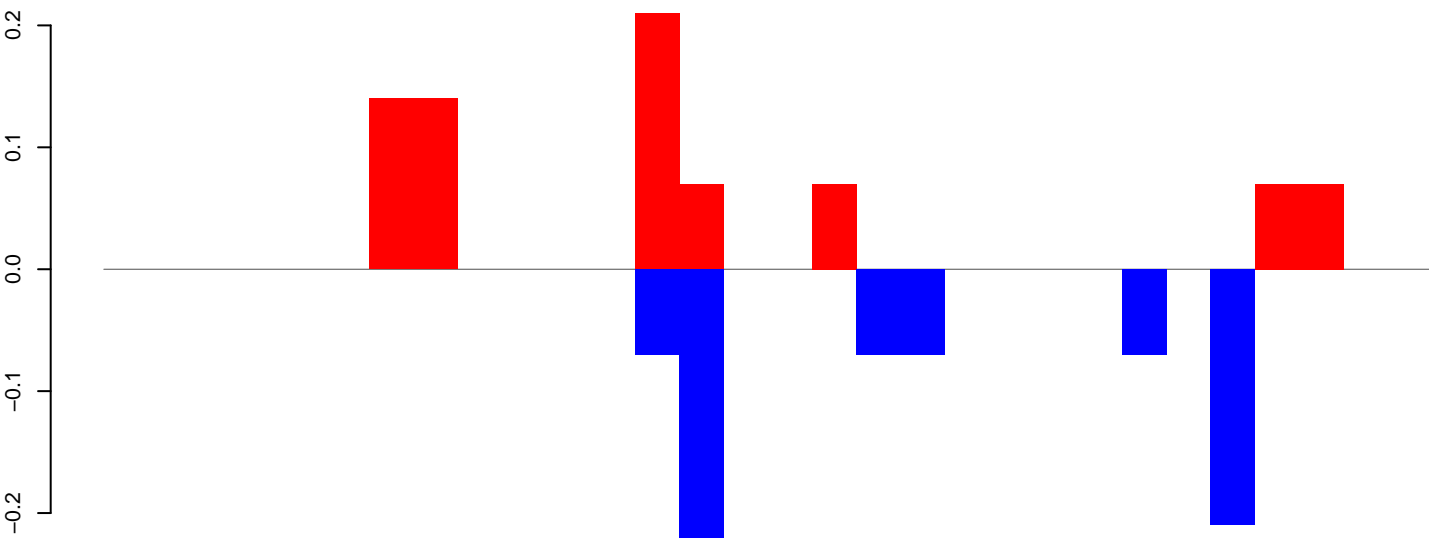
positive=0, negative=0, total=1

CuQuin_Mcarc_NL.24_35.rep



positive=0, negative=1, total=1

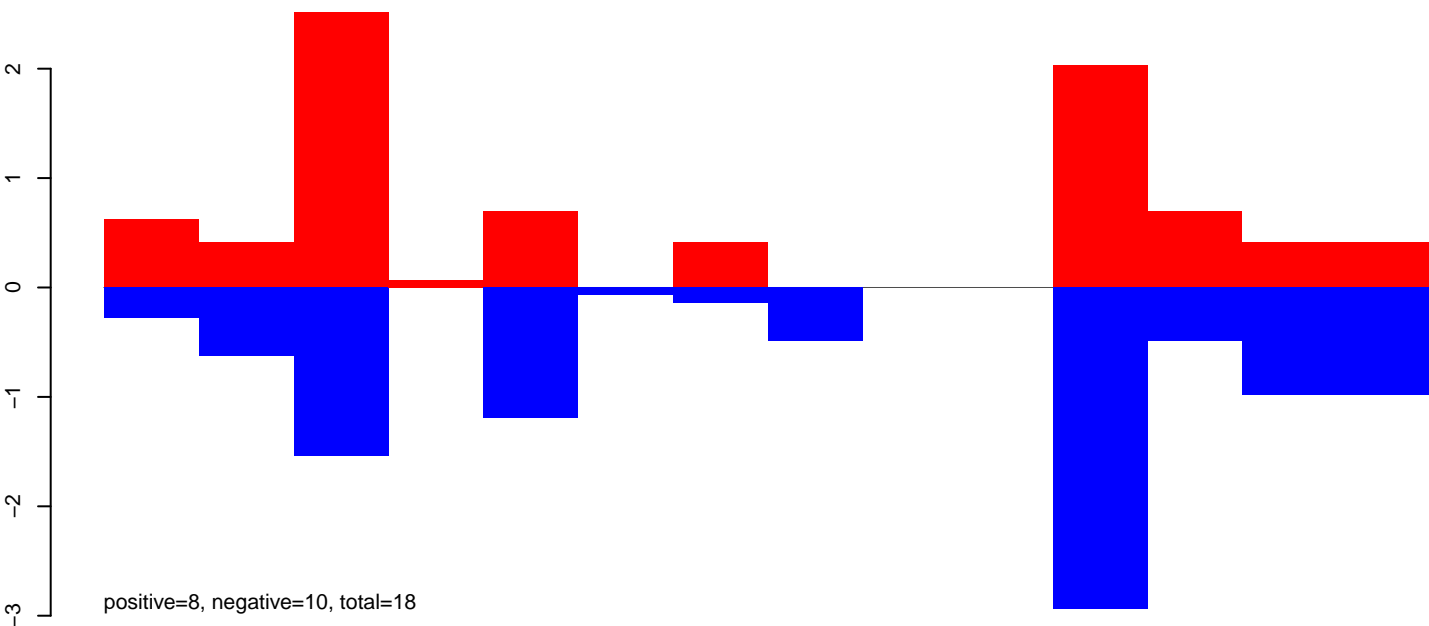
CuQuin_Mcarc_NL.rep



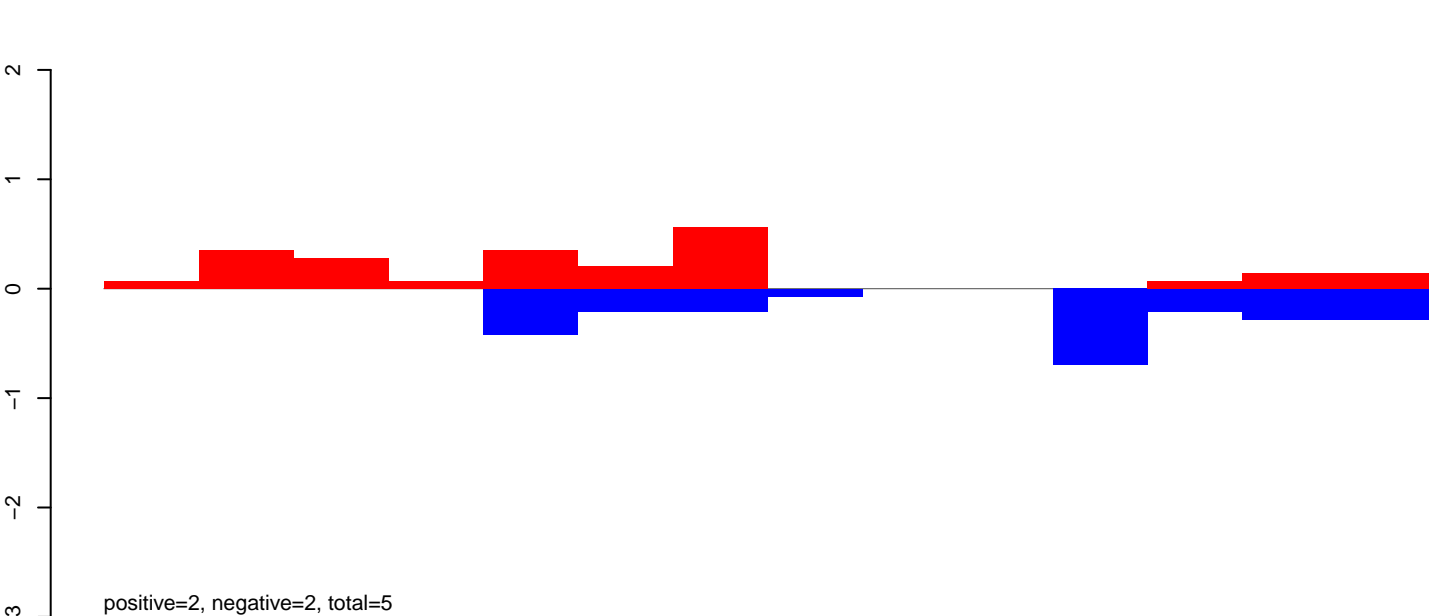
positive=1, negative=1, total=2

Window size=25, length=746, TE@rnd-4_family-615@DNA_Sola-1@1603:1-746

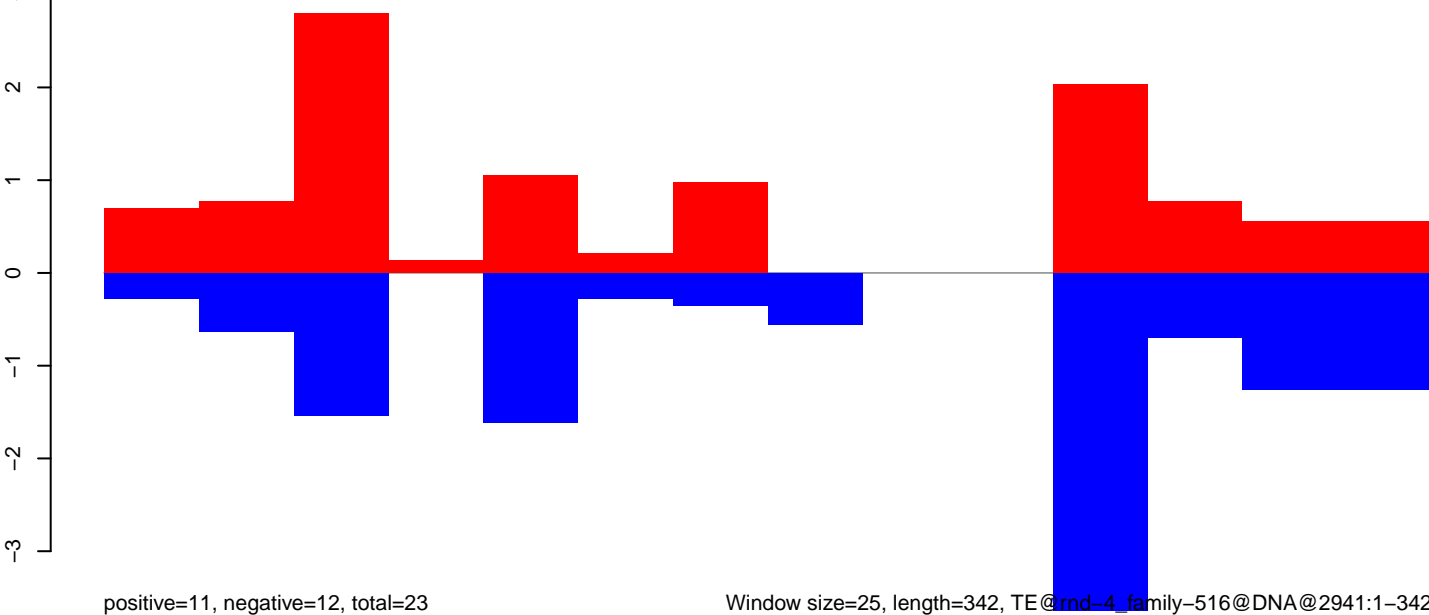
CuQuin_Mcarc_NL.18_23.rep



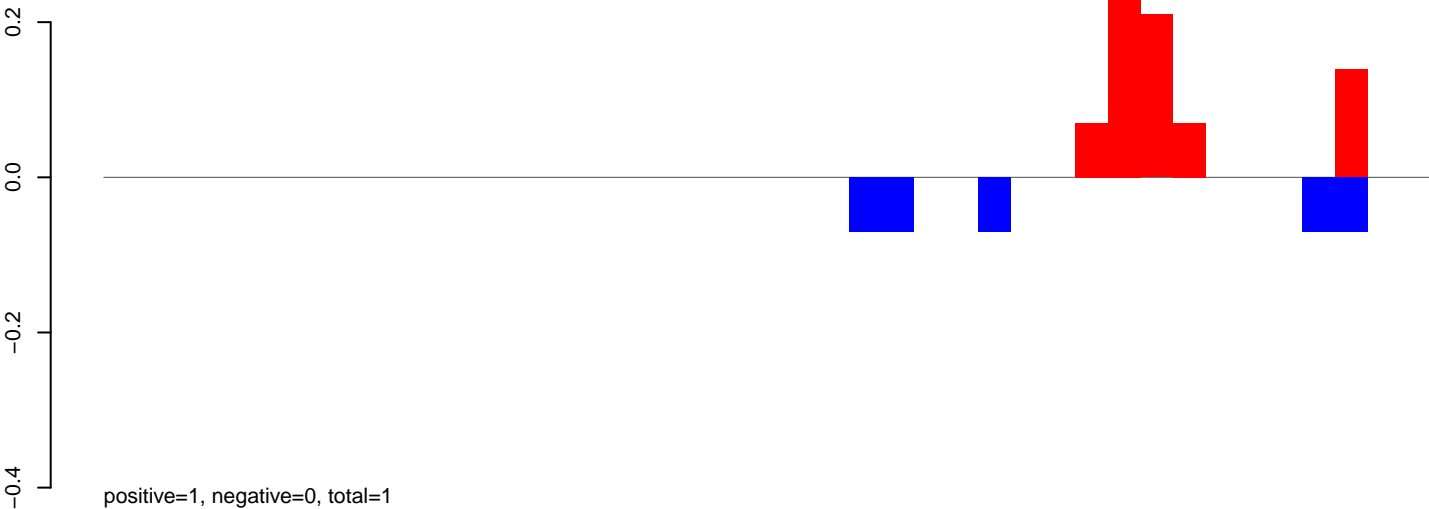
CuQuin_Mcarc_NL.24_35.rep



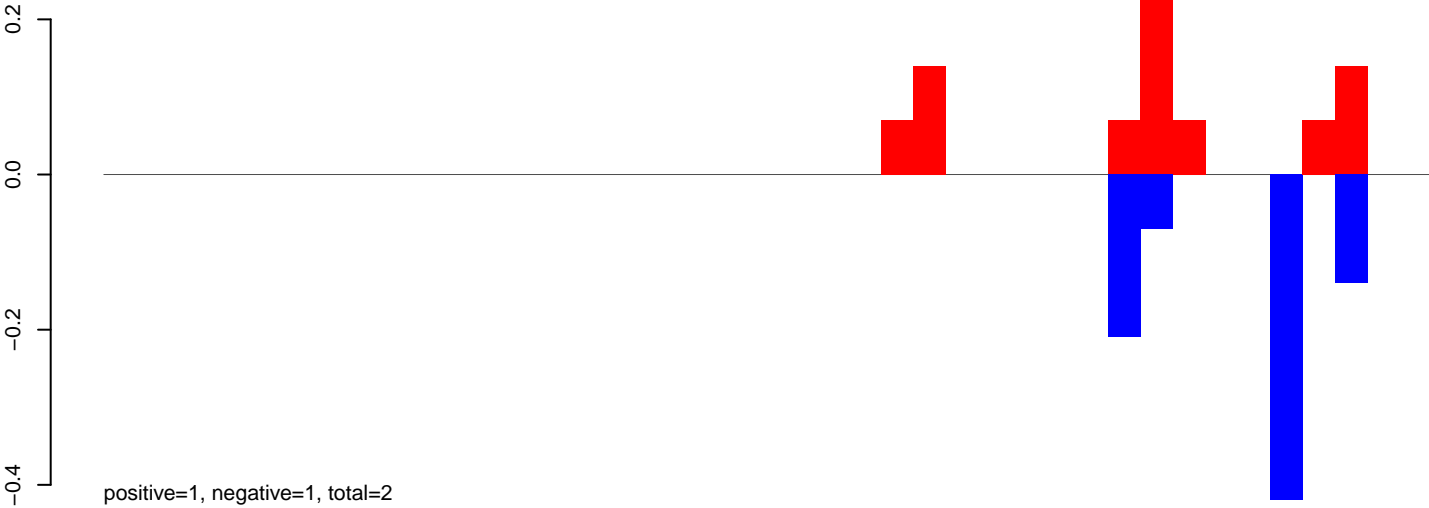
CuQuin_Mcarc_NL.rep



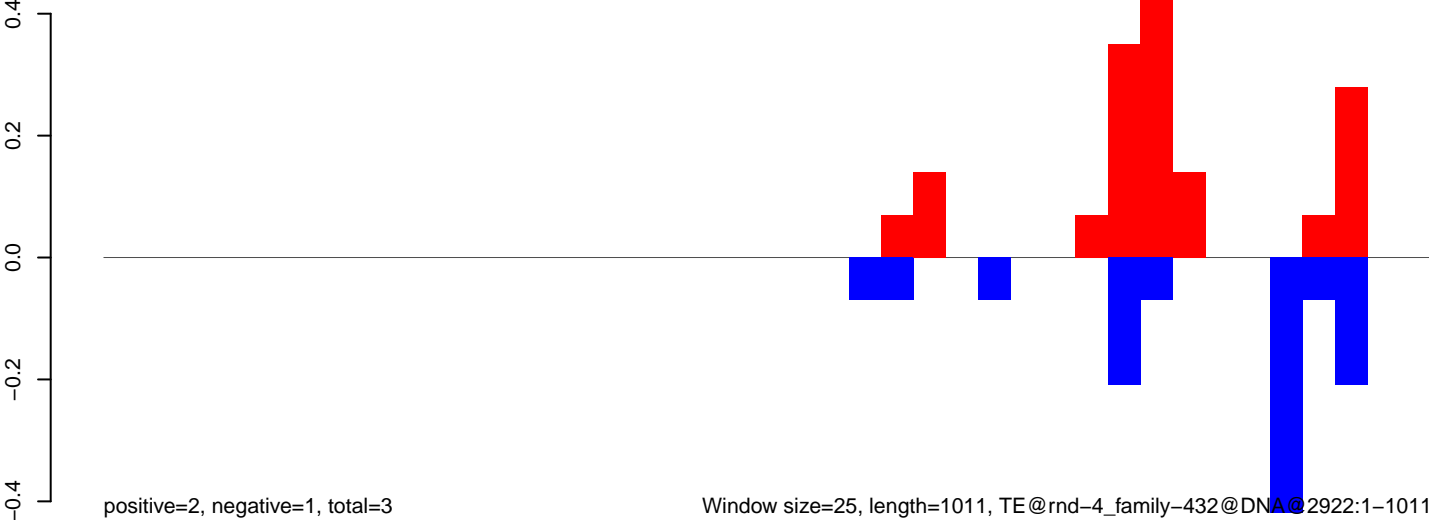
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

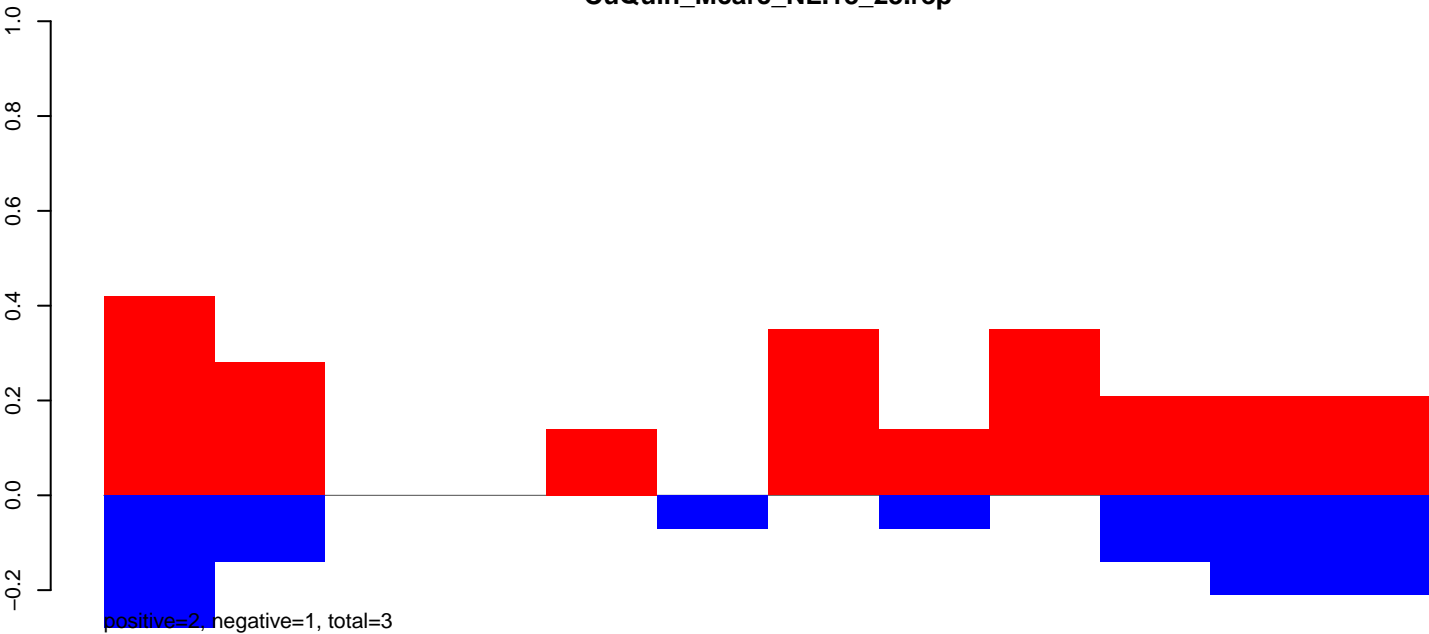


CuQuin_Mcarc_NL.rep

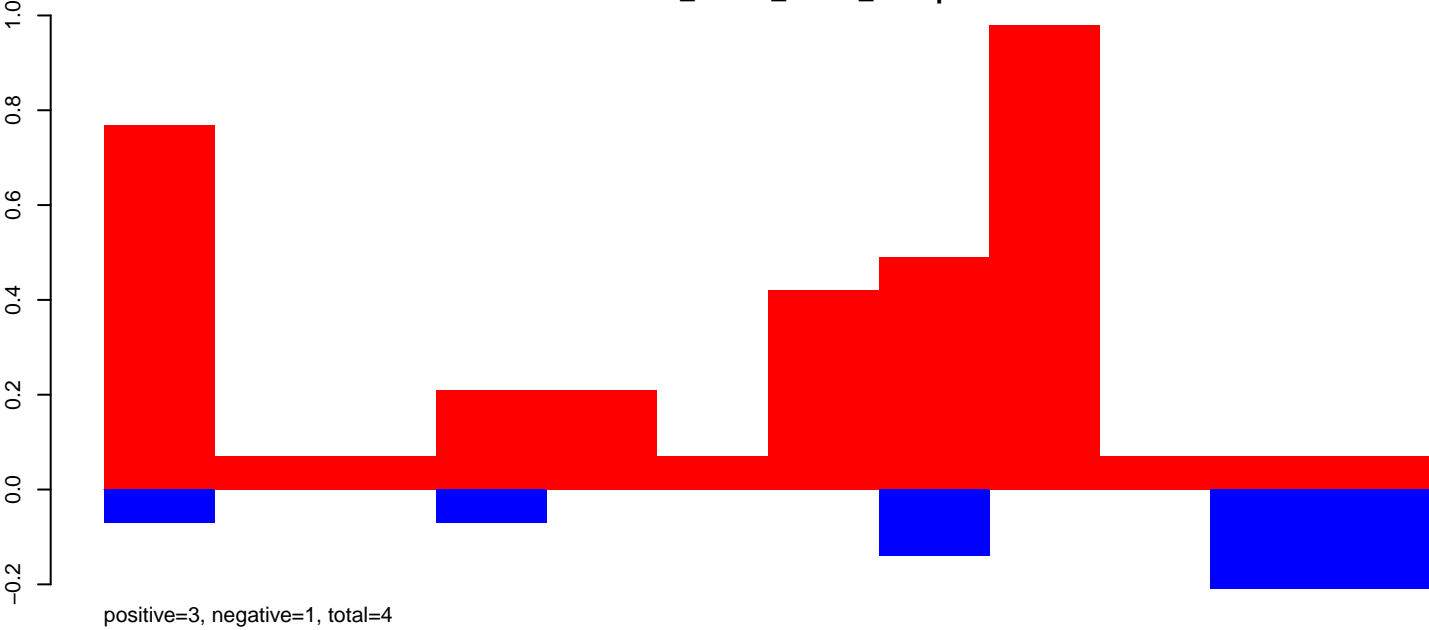


Window size=25, length=1011, TE@rnd-4_family-432@DNA@2922:1-1011

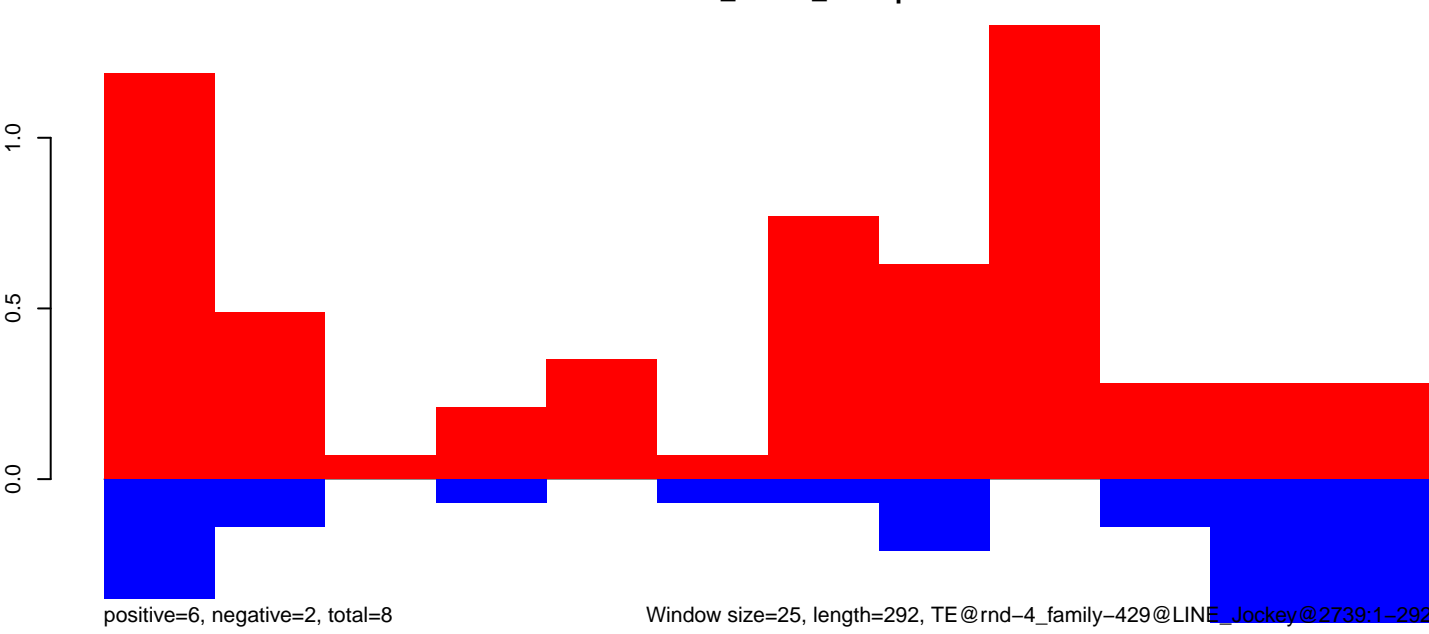
CuQuin_Mcarc_NL.18_23.rep



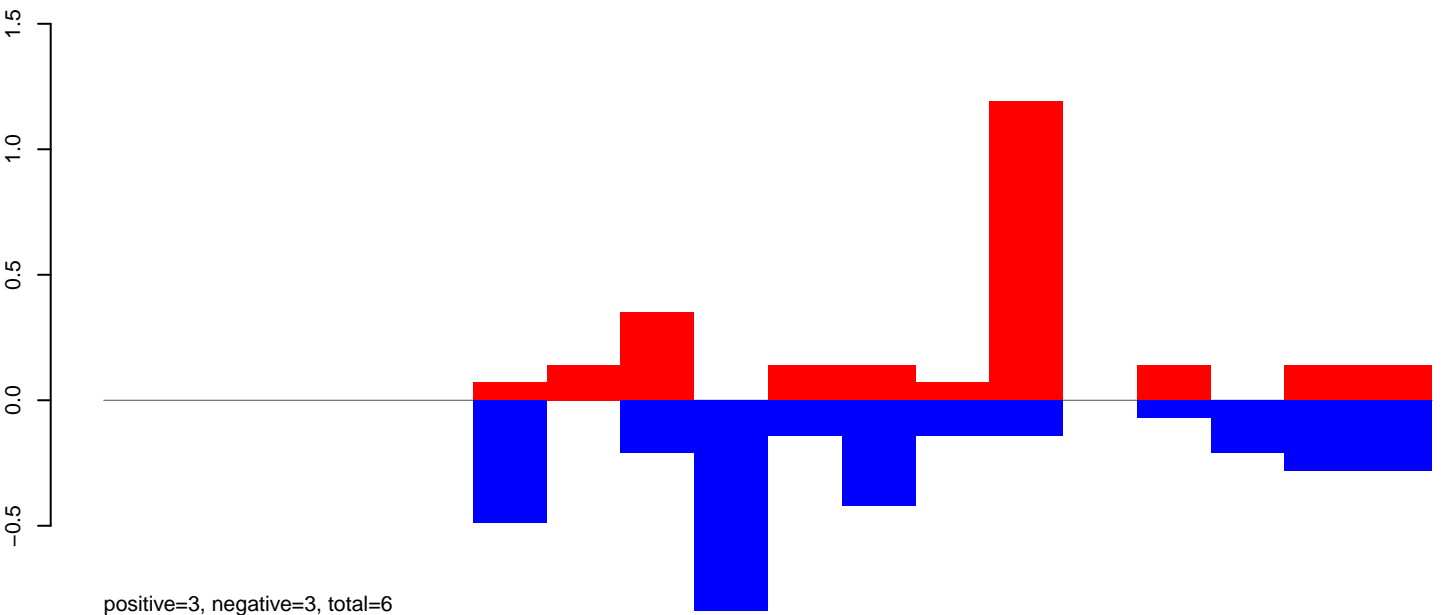
CuQuin_Mcarc_NL.24_35.rep



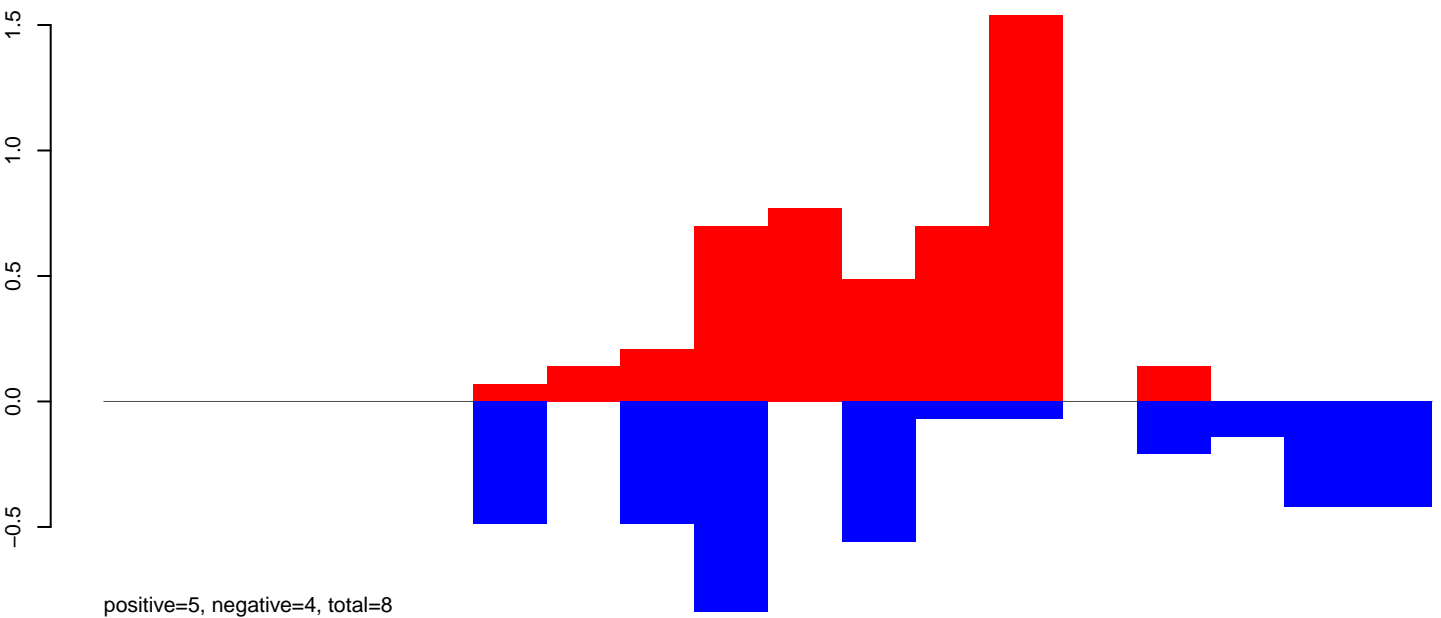
CuQuin_Mcarc_NL.rep



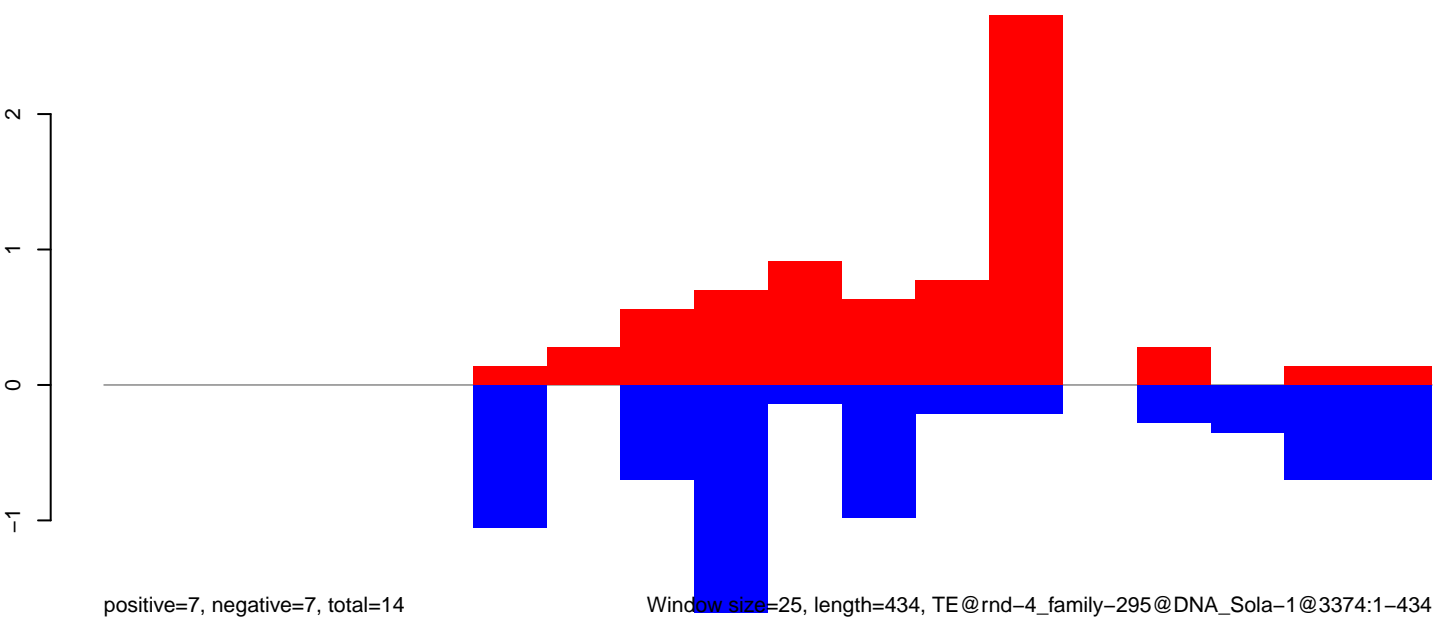
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=434, TE@rnd-4_family-295@DNA_Sola-1@3374:1-434

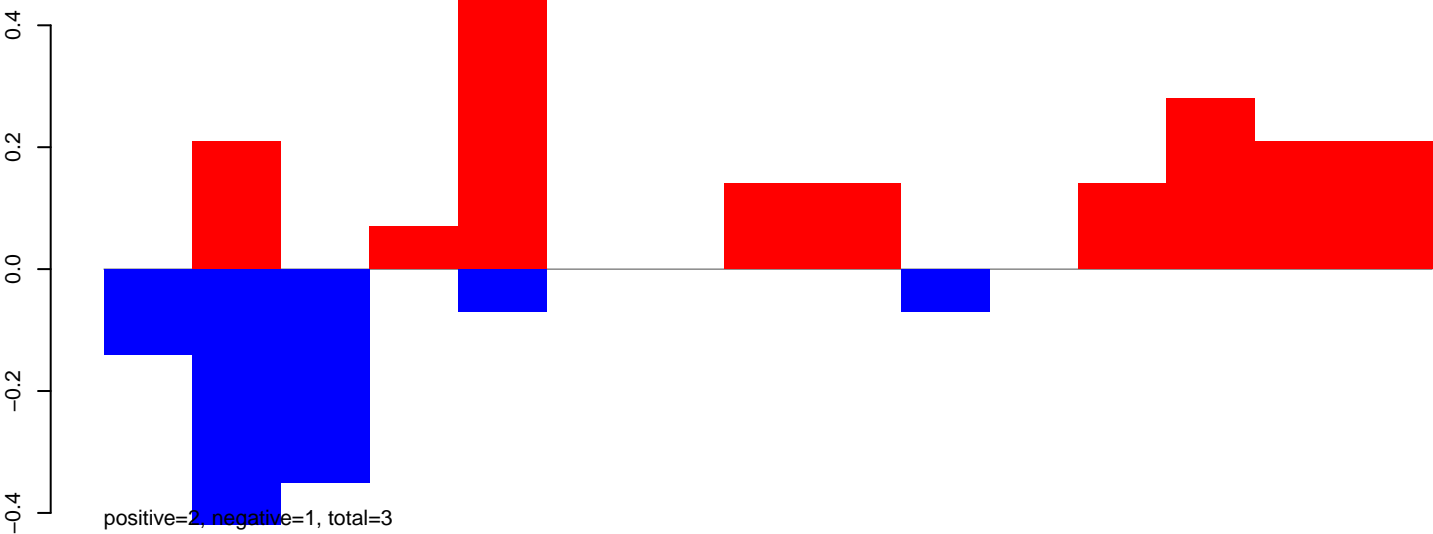
100

200

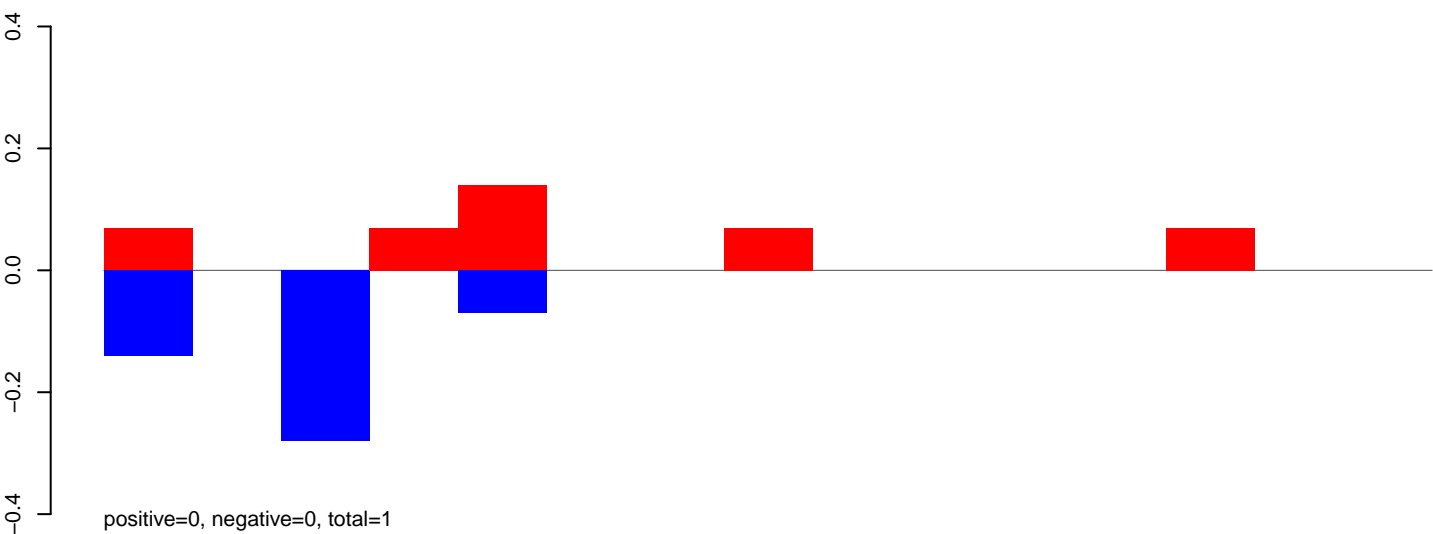
300

400

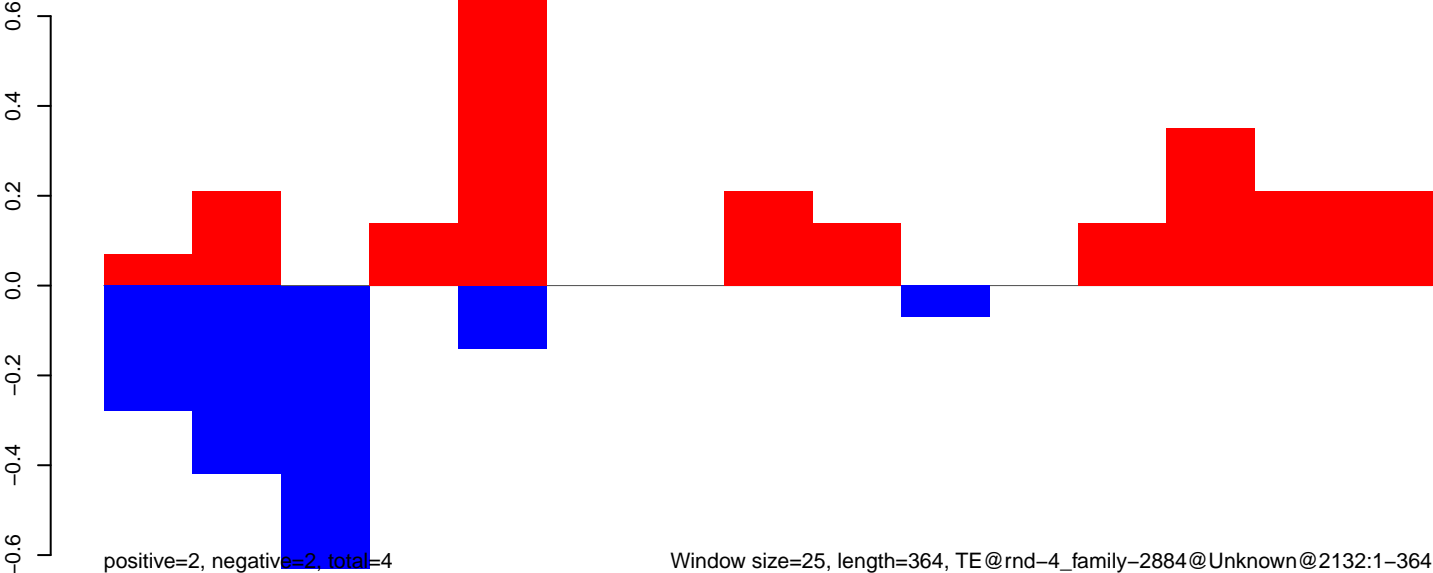
CuQuin_Mcarc_NL.18_23.rep



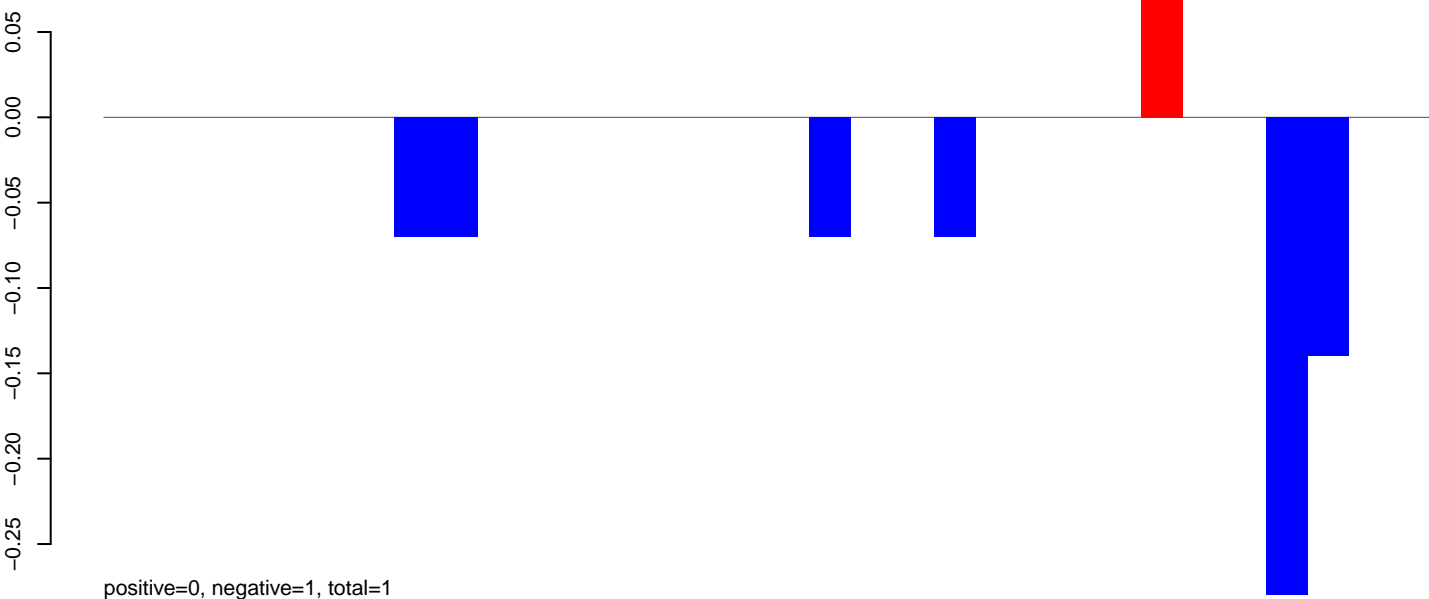
CuQuin_Mcarc_NL.24_35.rep



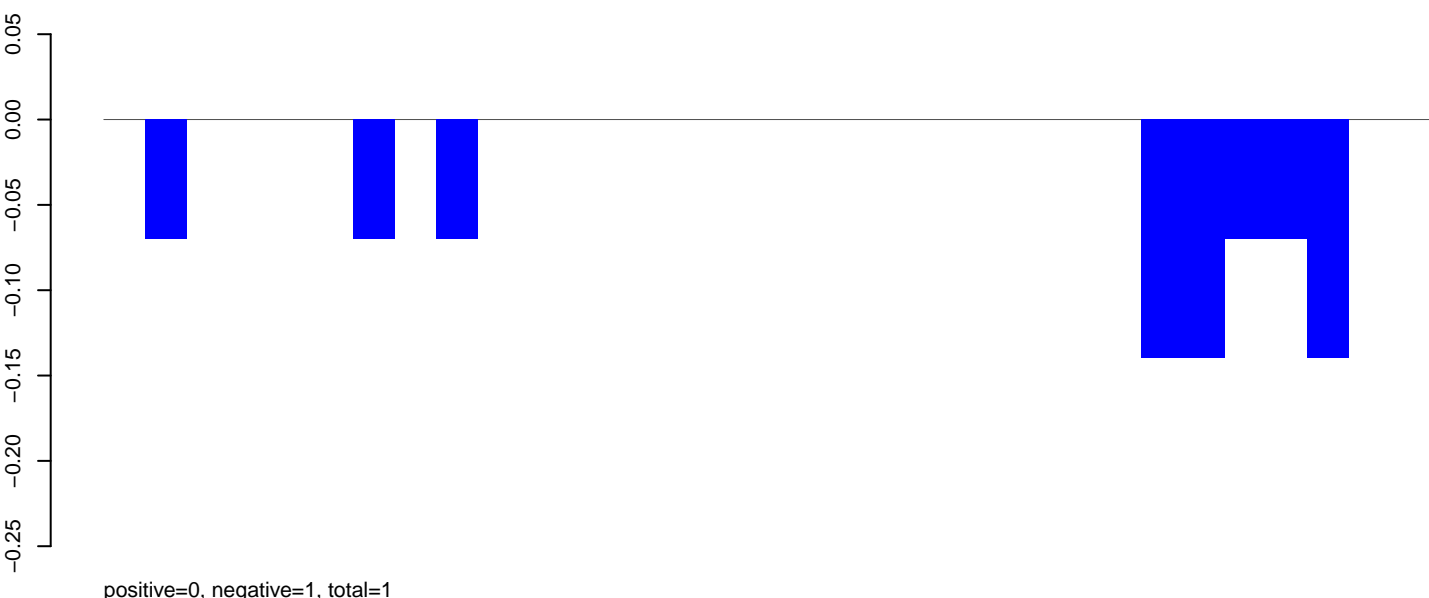
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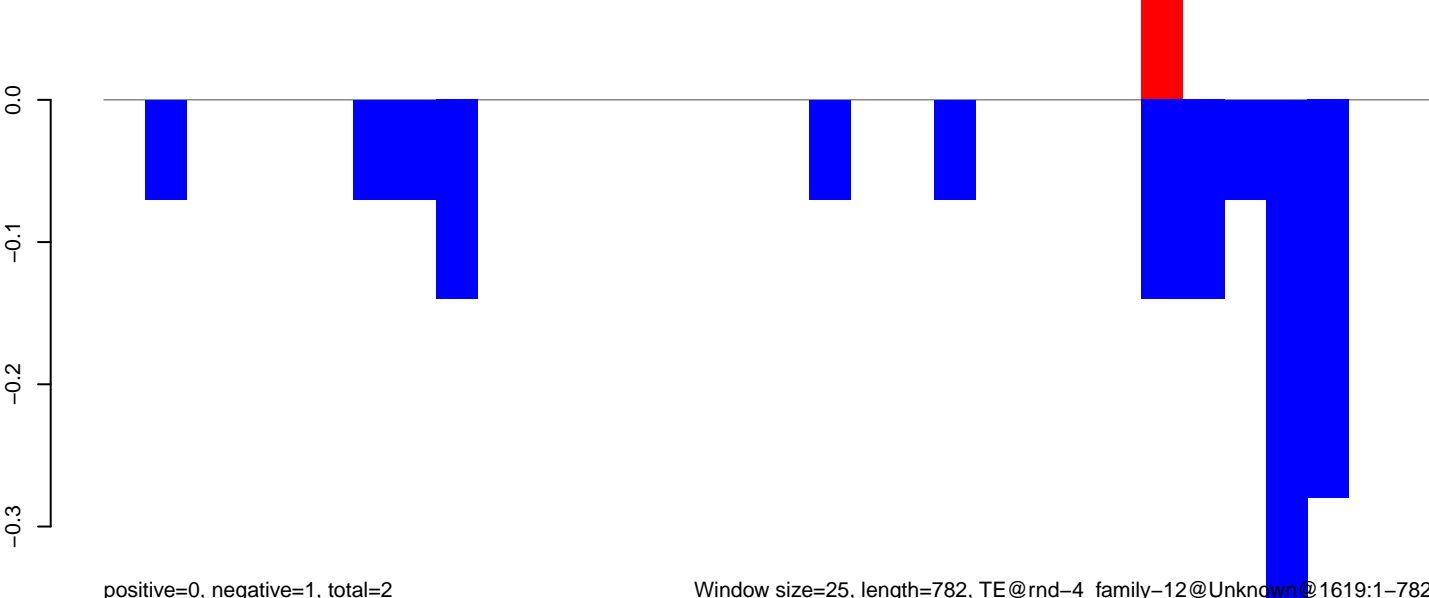
CuQuin_Mcarc_NL.18_23.rep



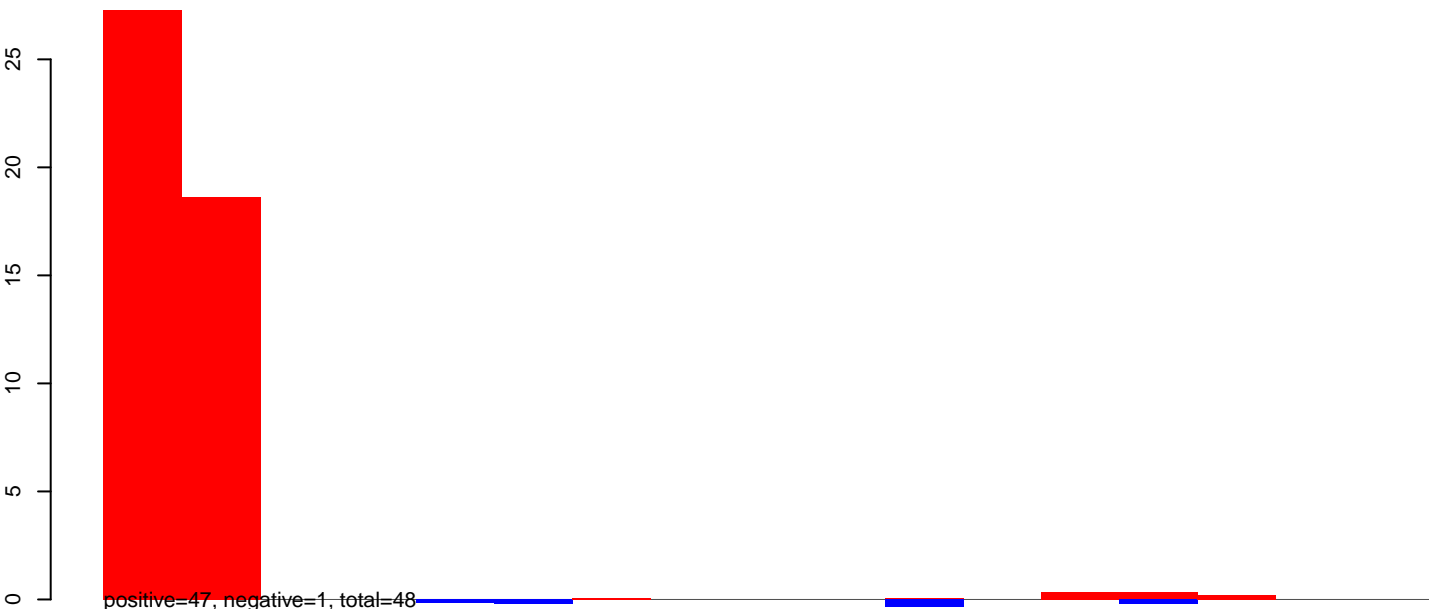
CuQuin_Mcarc_NL.24_35.rep



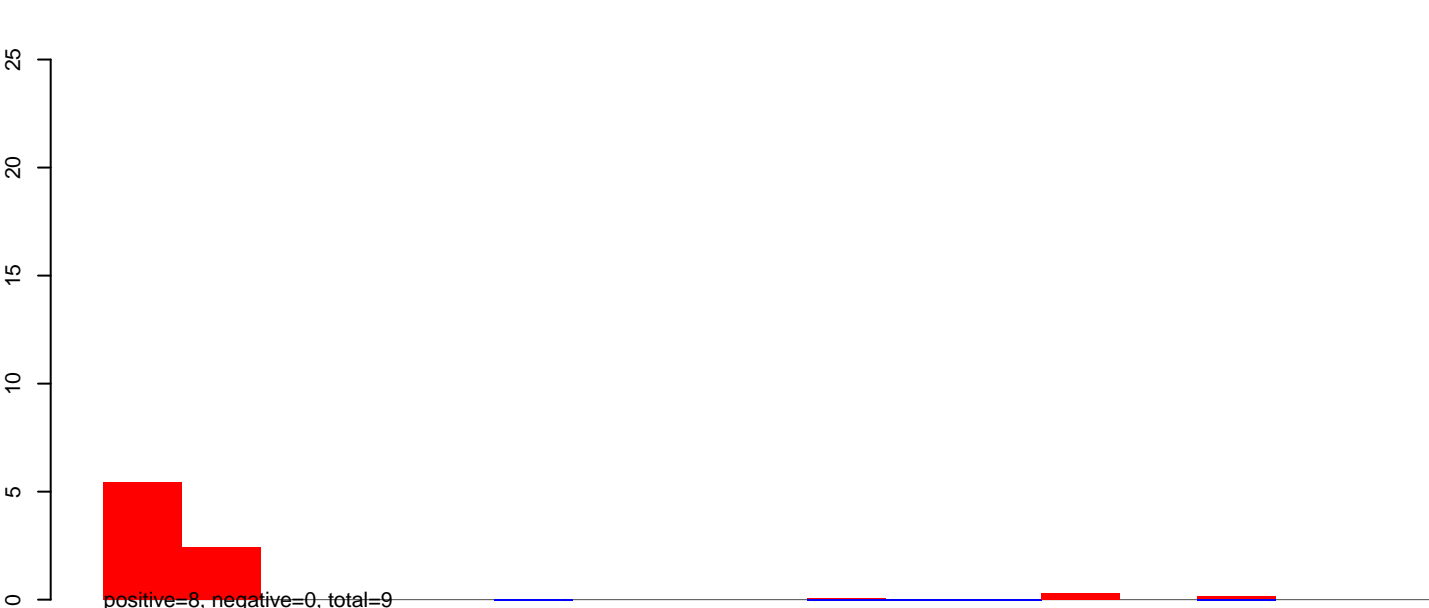
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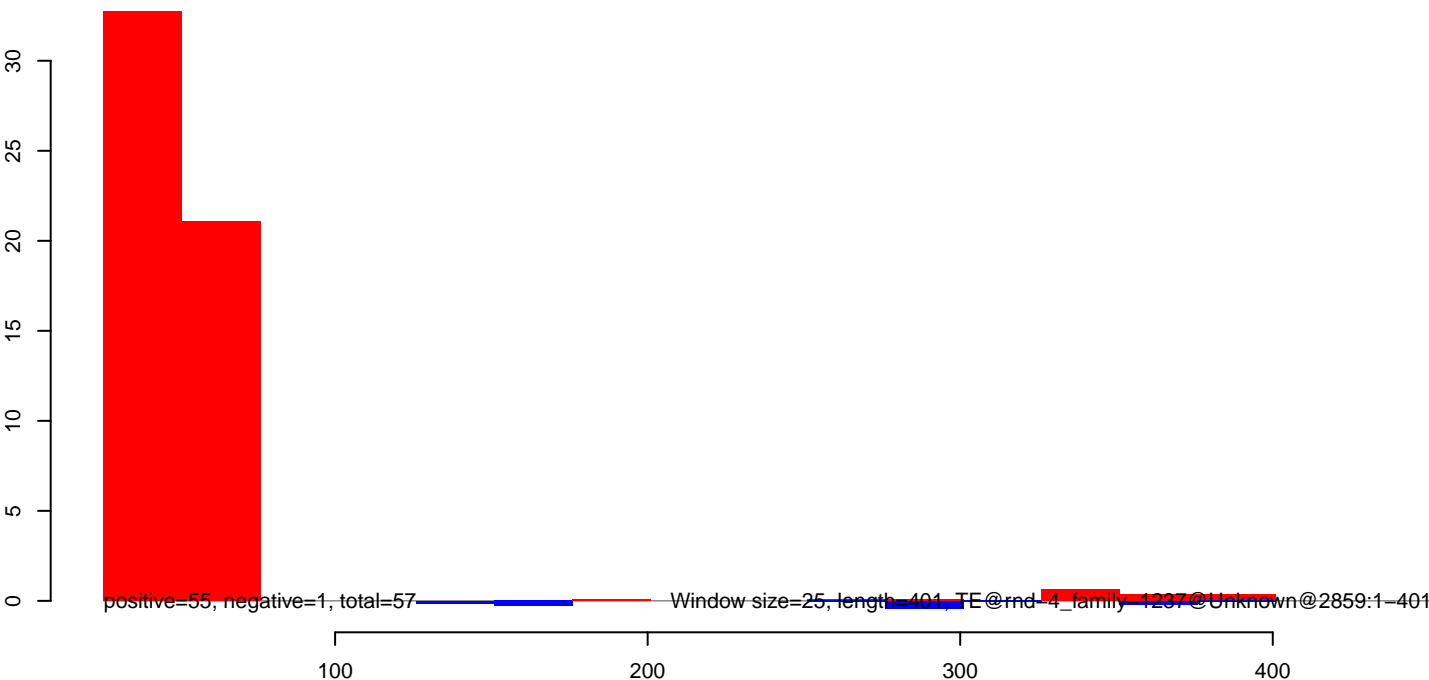
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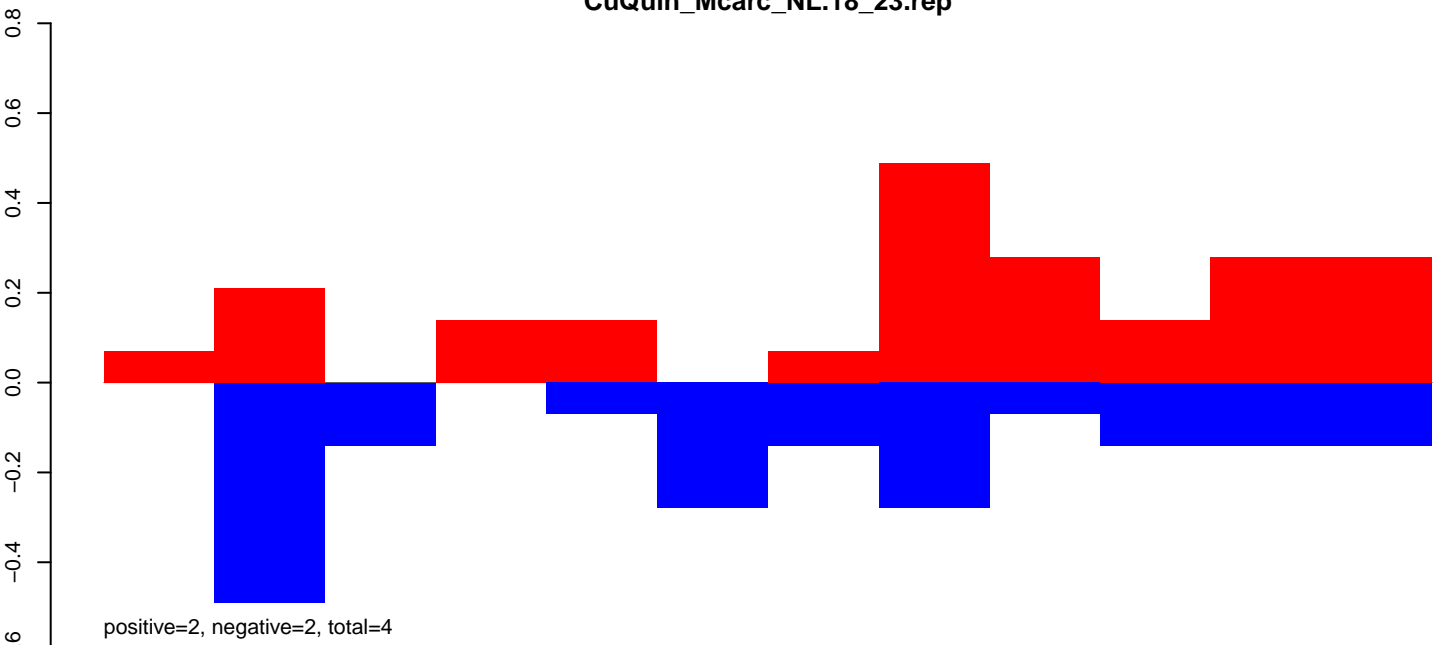
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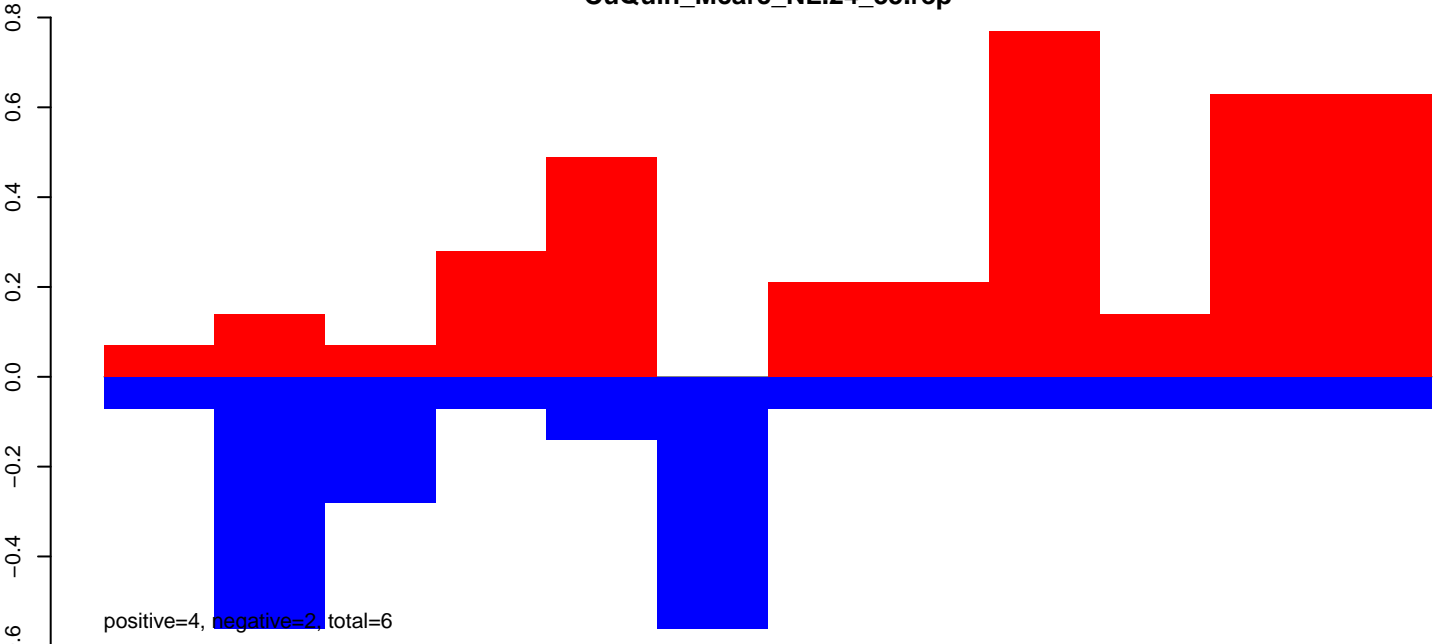
CuQuin_Mcarc_NL.rep



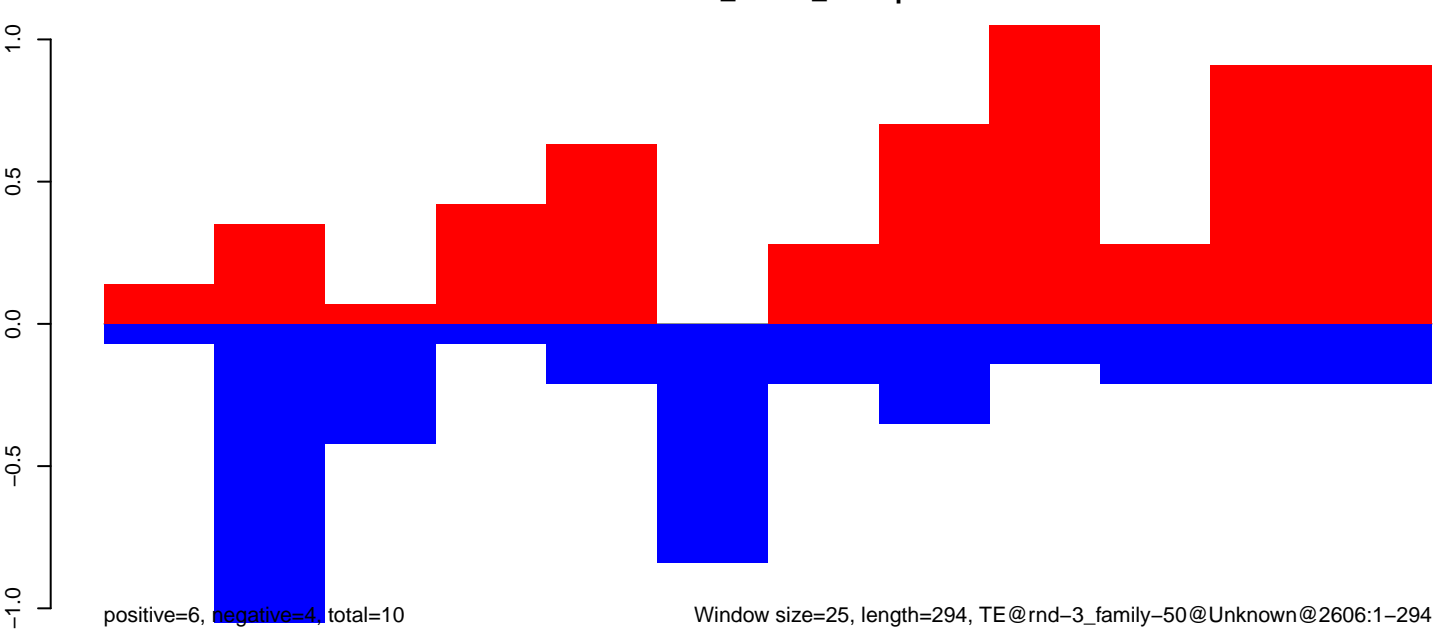
CuQuin_Mcarc_NL.18_23.rep



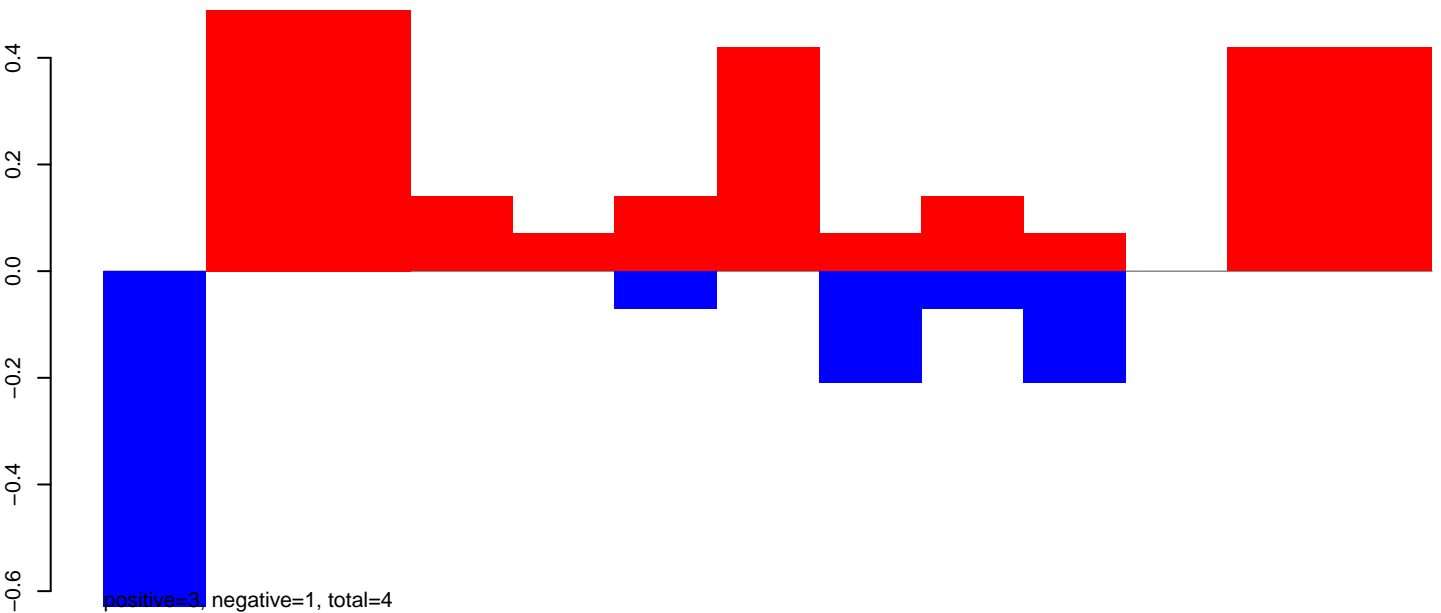
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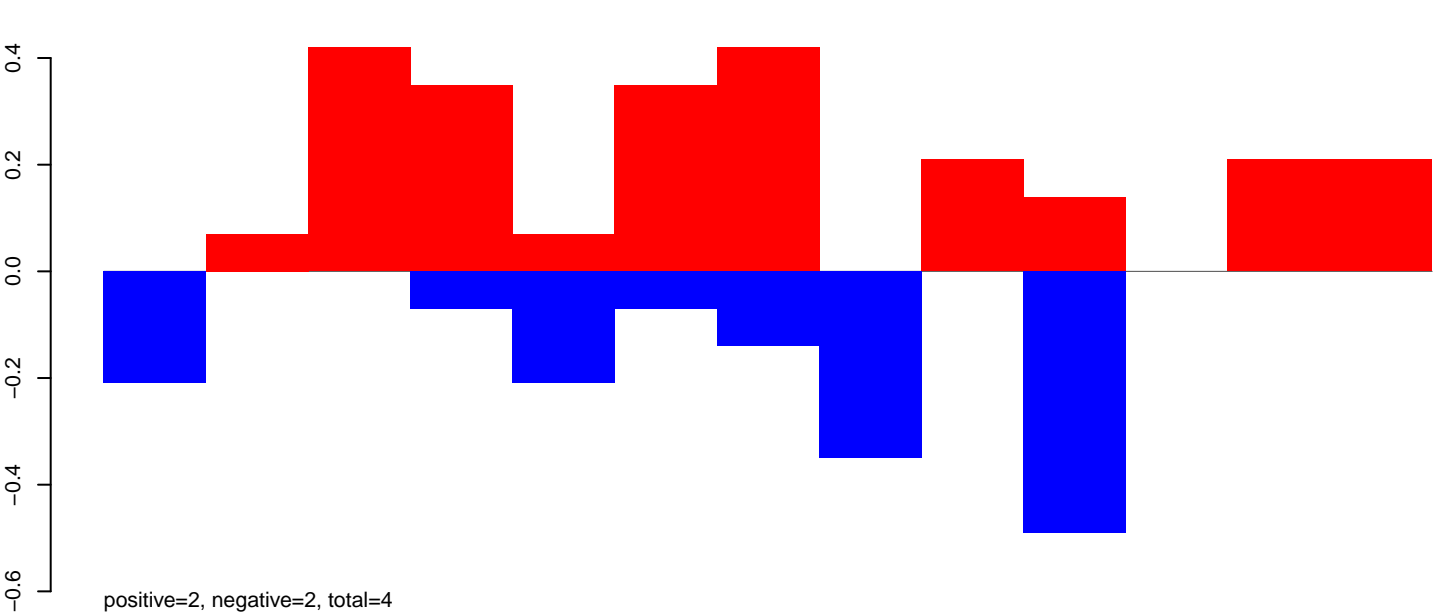
CuQuin_Mcarc_NL.rep



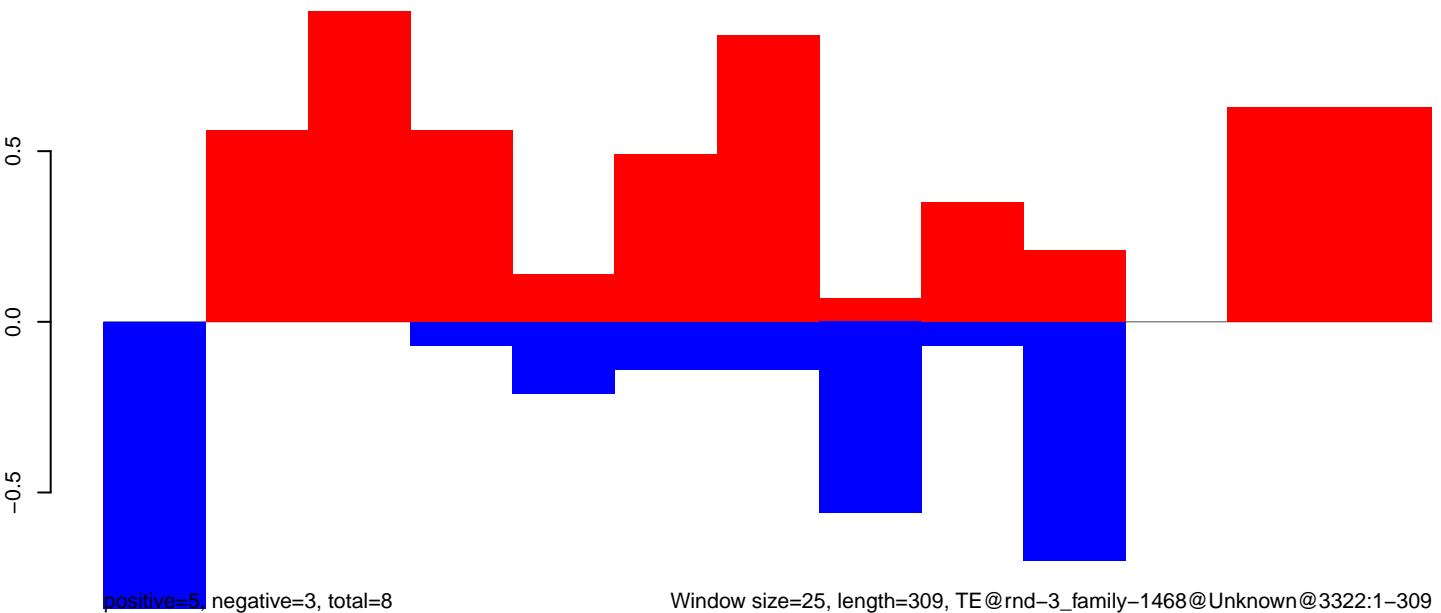
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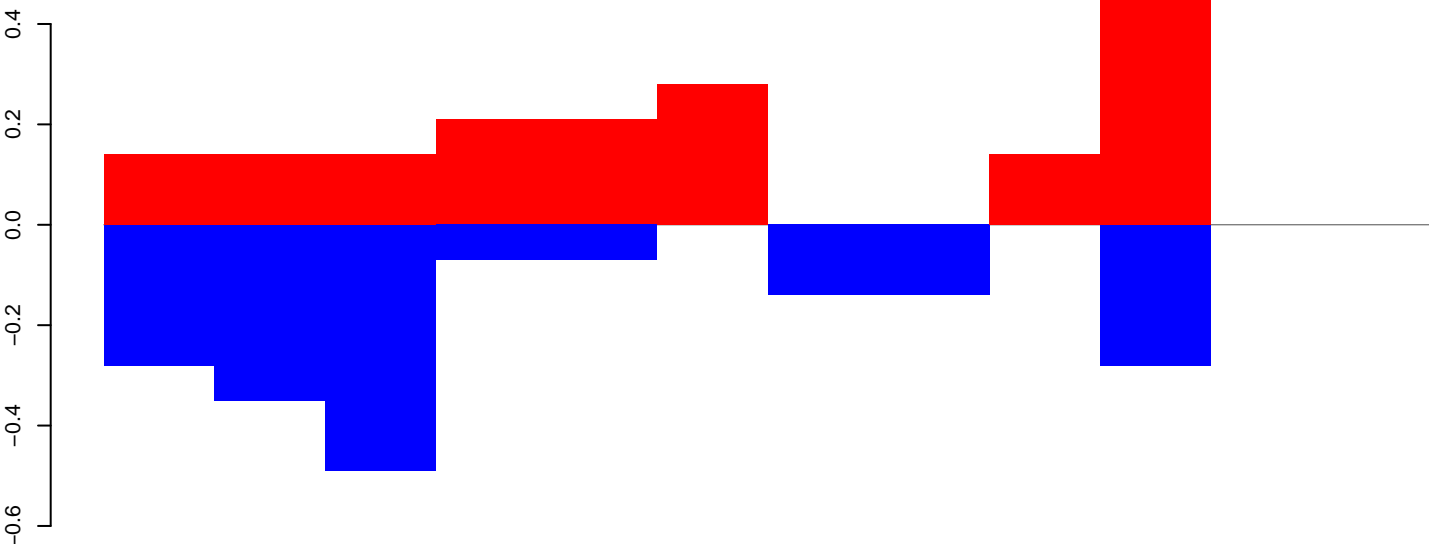
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CuQuin_Mcarc_NL.rep

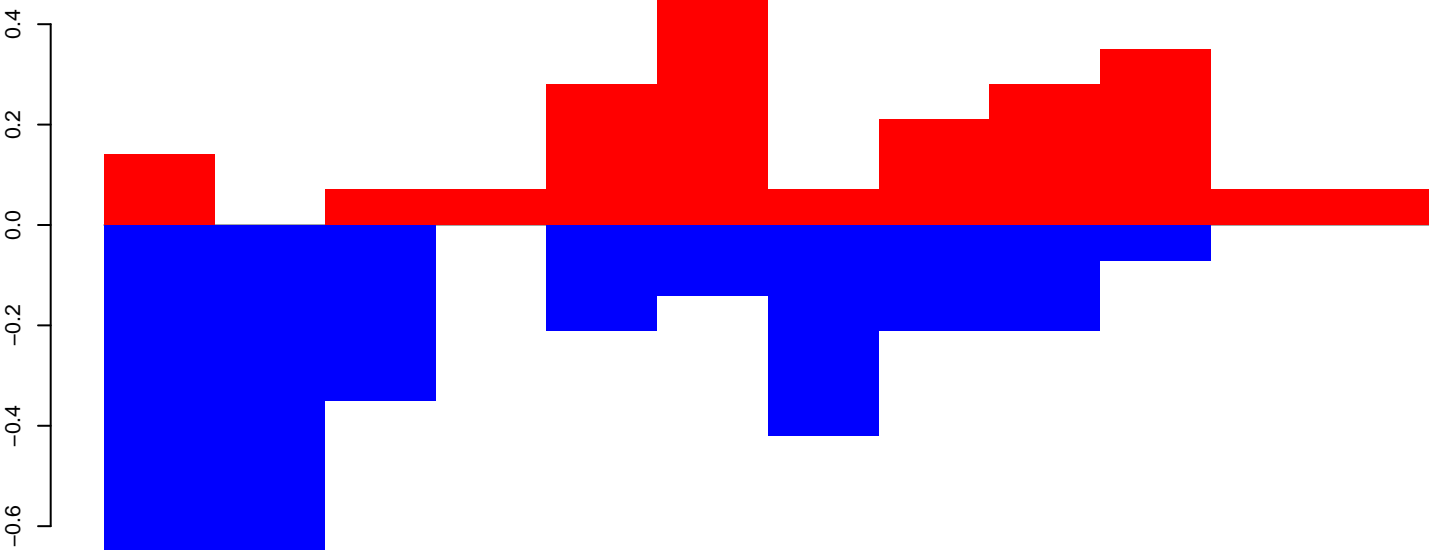


CuQuin_Mcarc_NL.18_23.rep



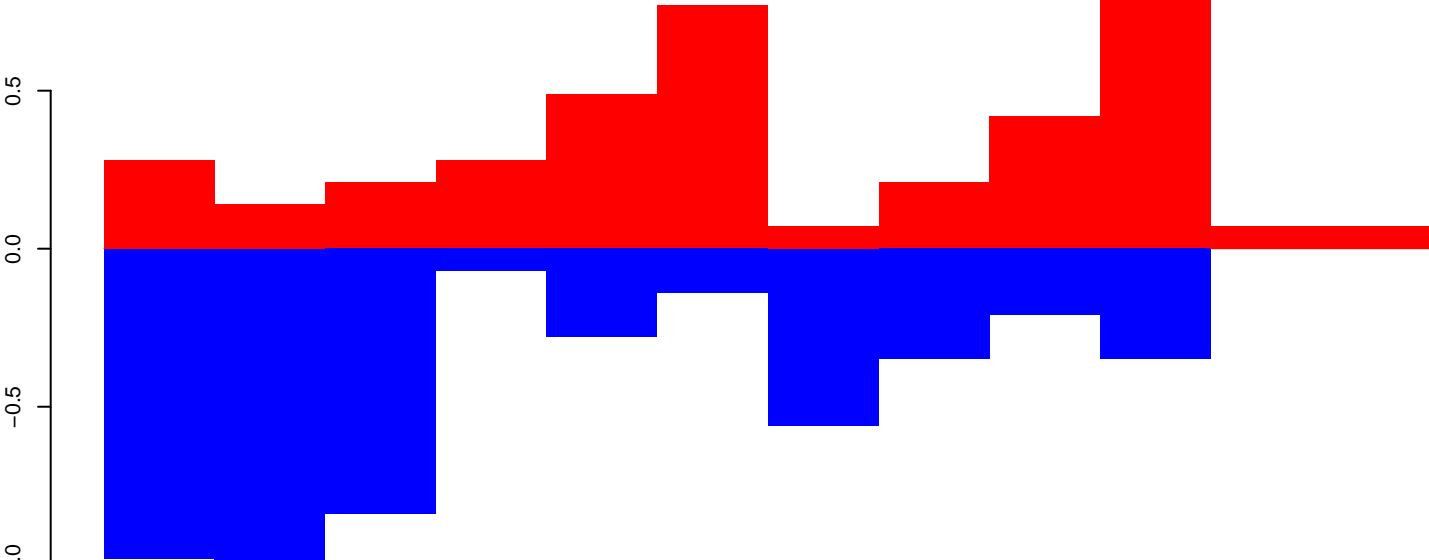
positive=2, negative=2, total=4

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=3, total=5

CuQuin_Mcarc_NL.rep

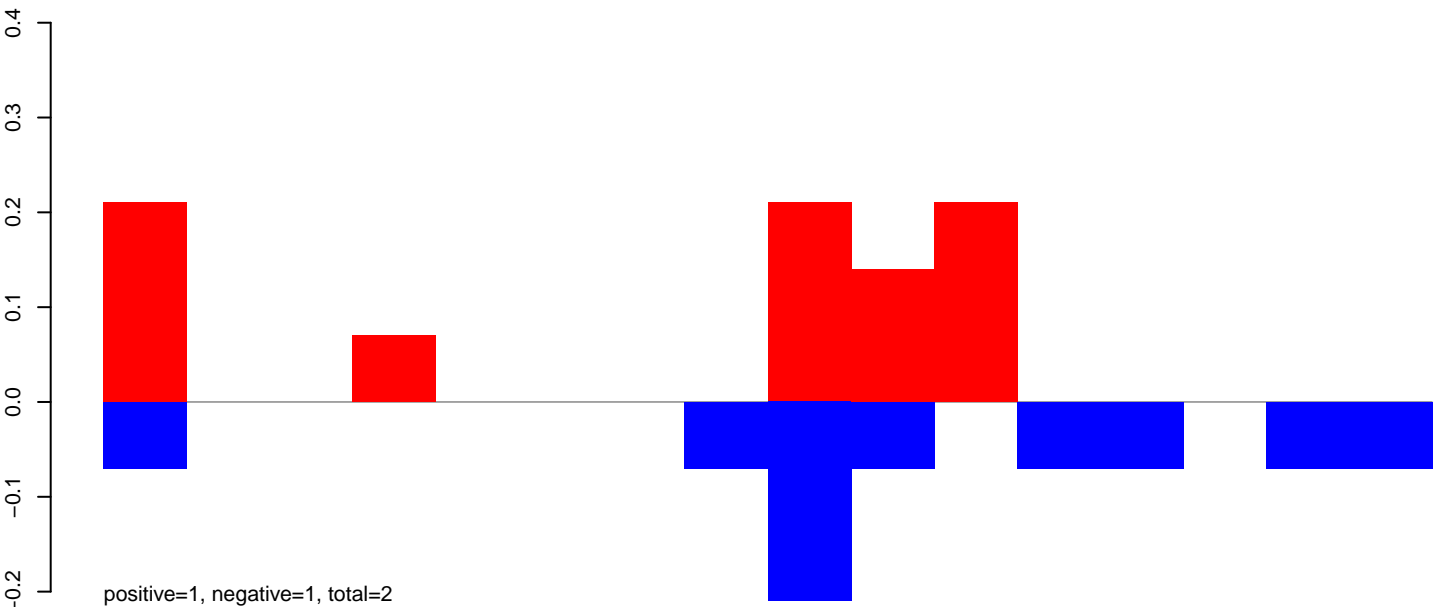


positive=4, negative=5, total=9

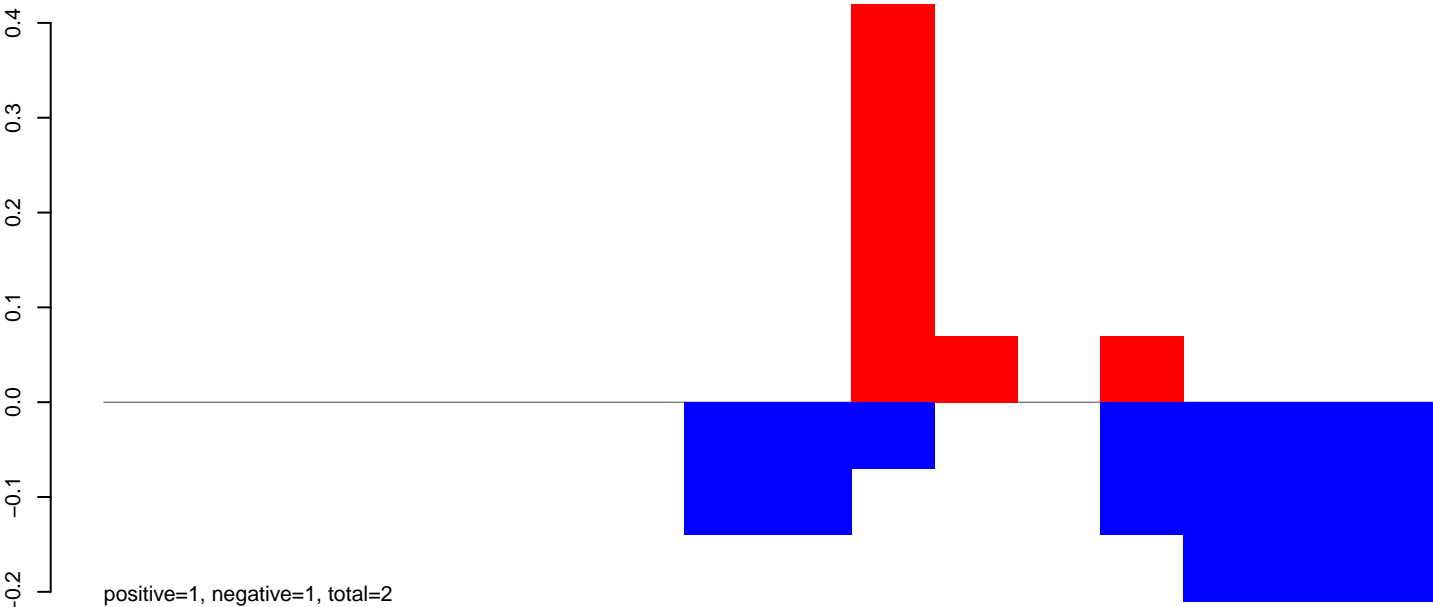
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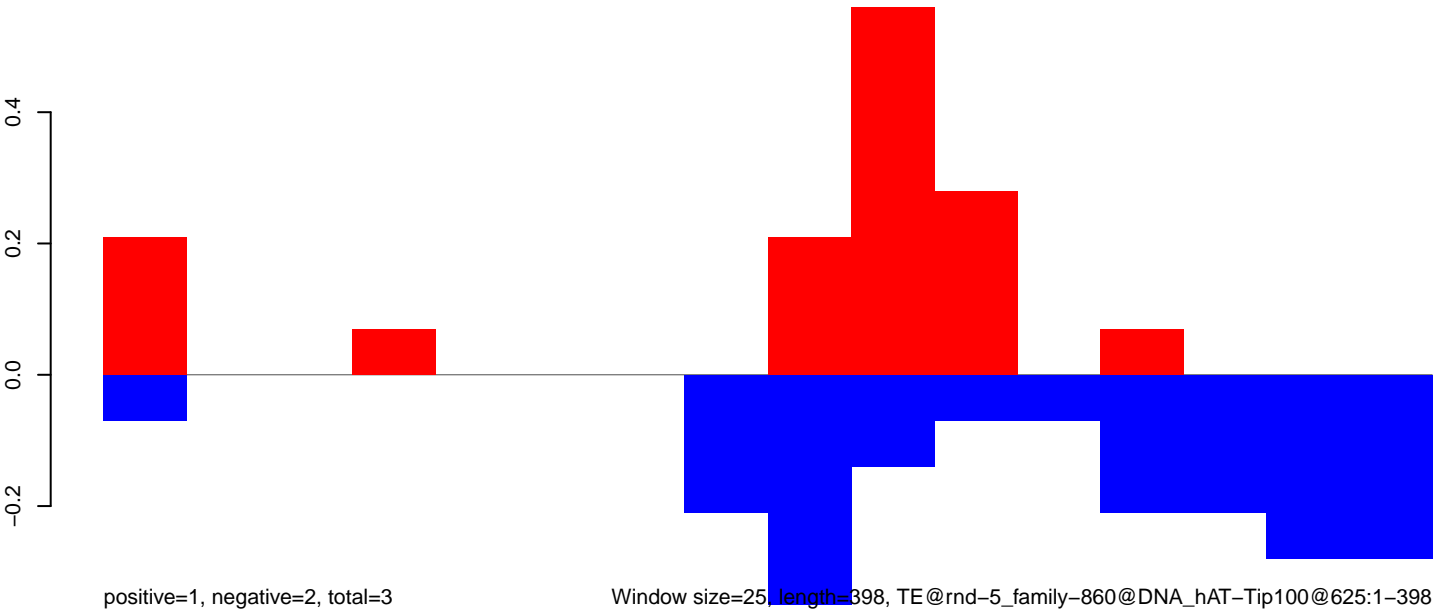
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



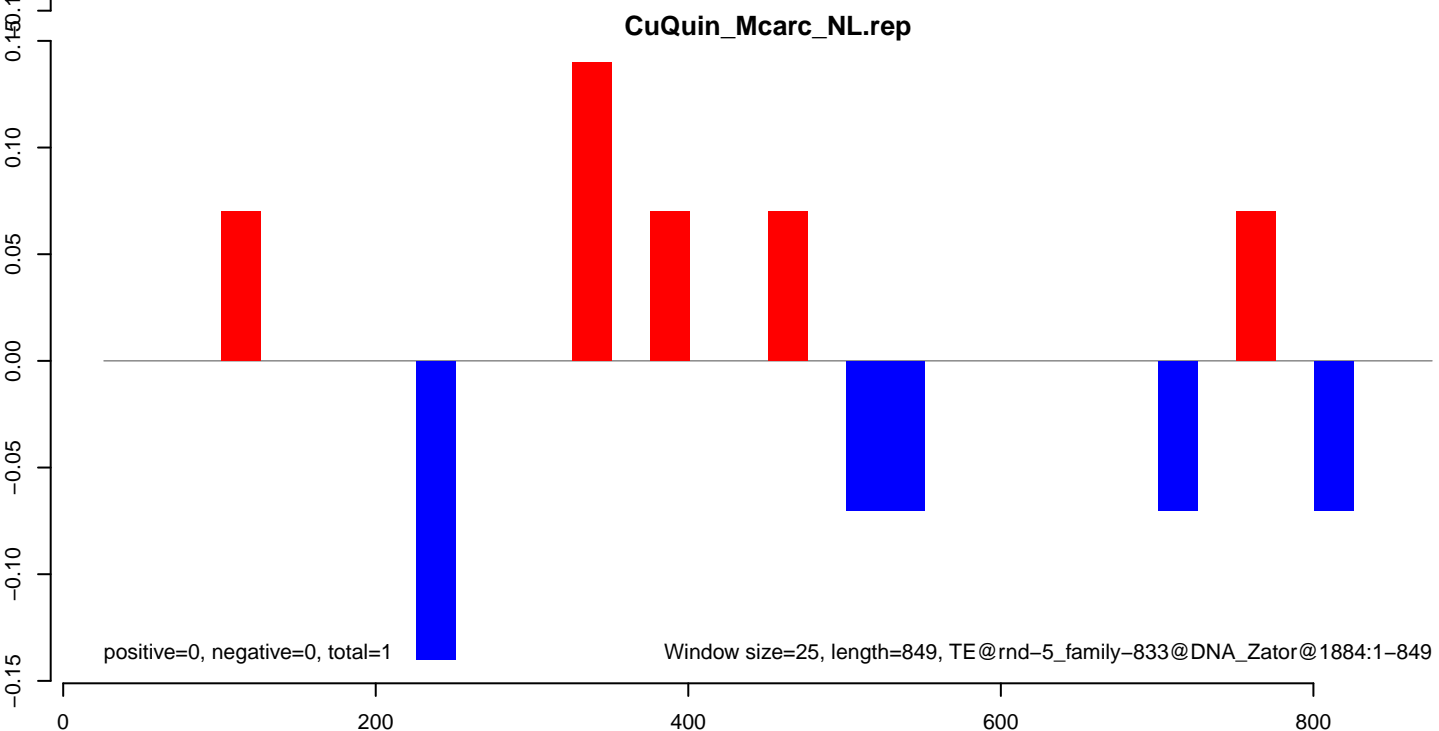
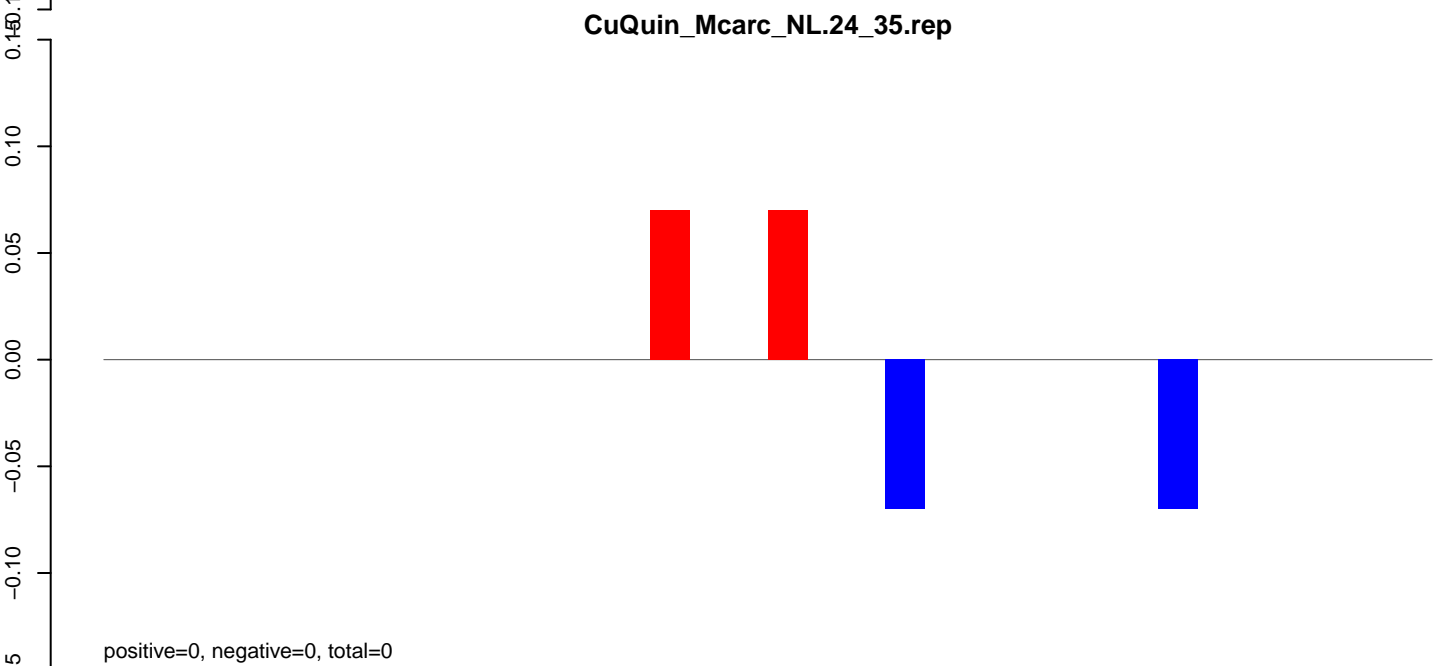
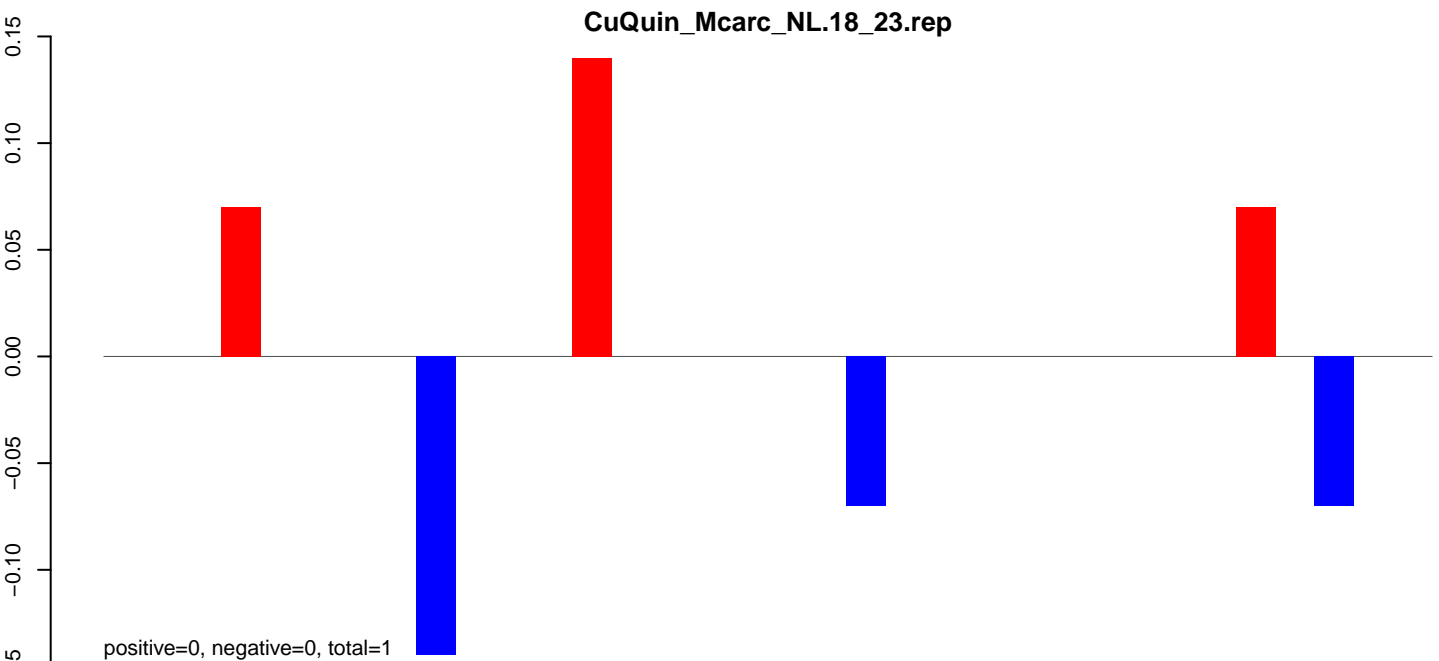
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100

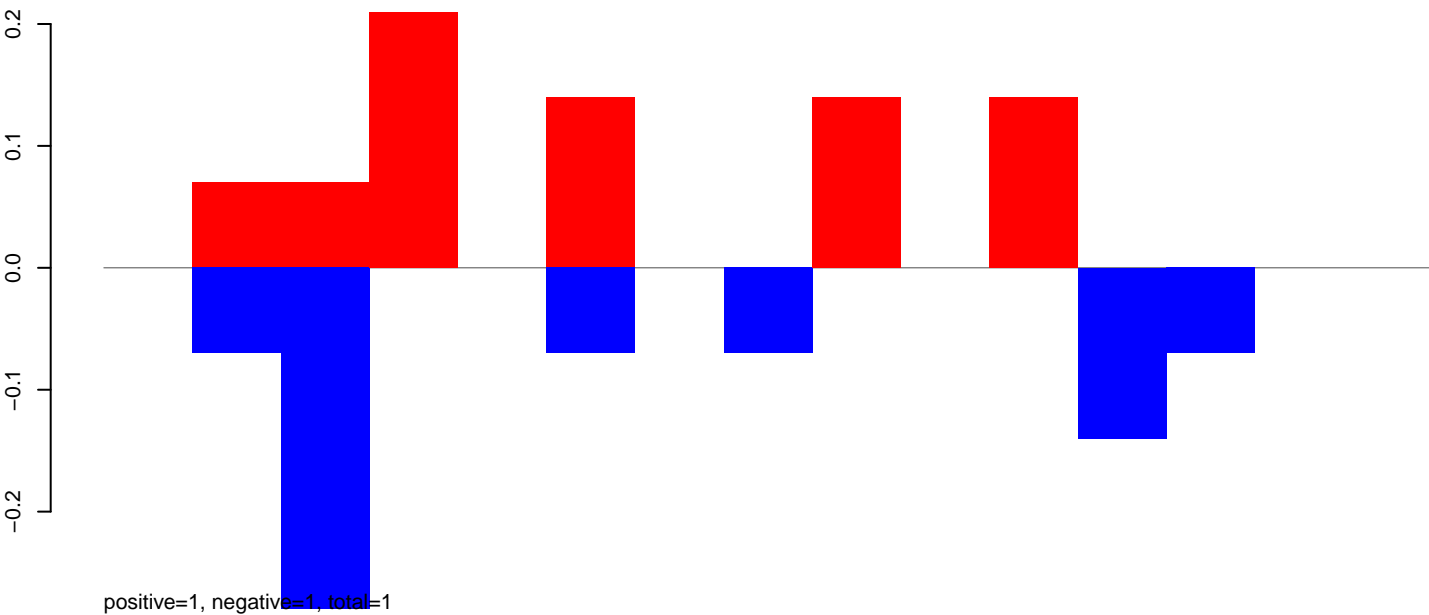
200

300

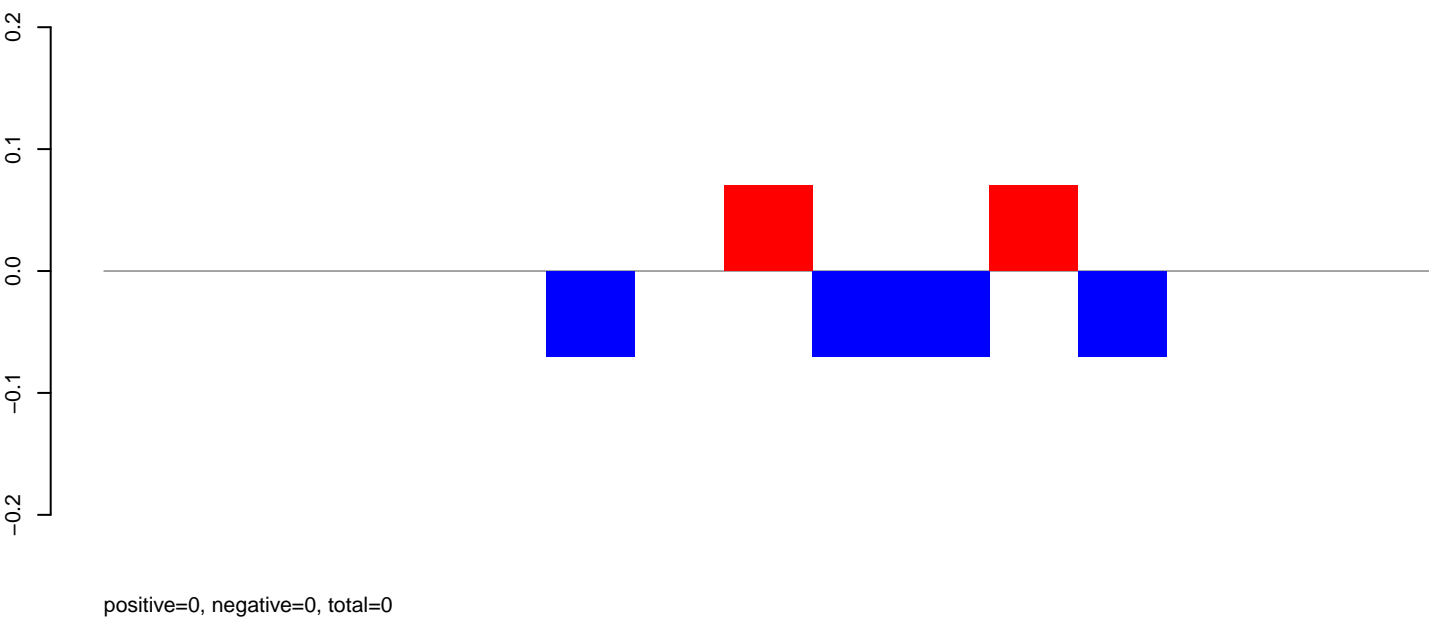
400



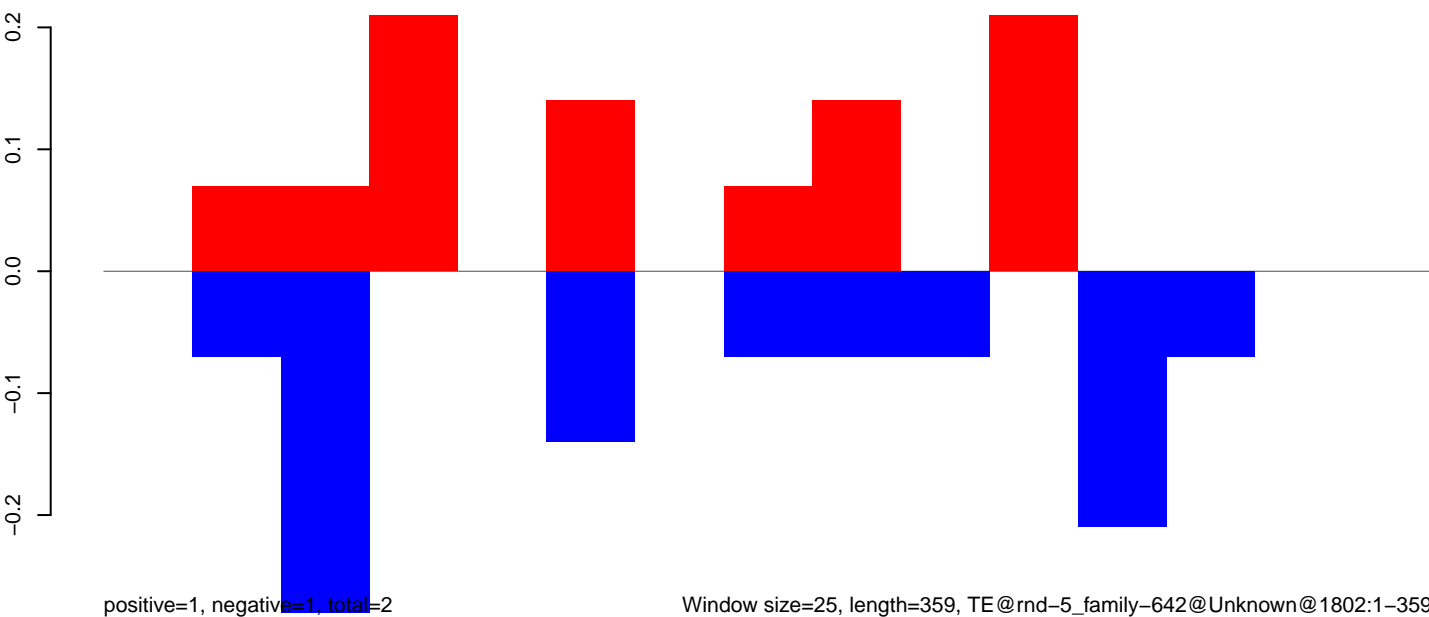
CuQuin_Mcarc_NL.18_23.rep



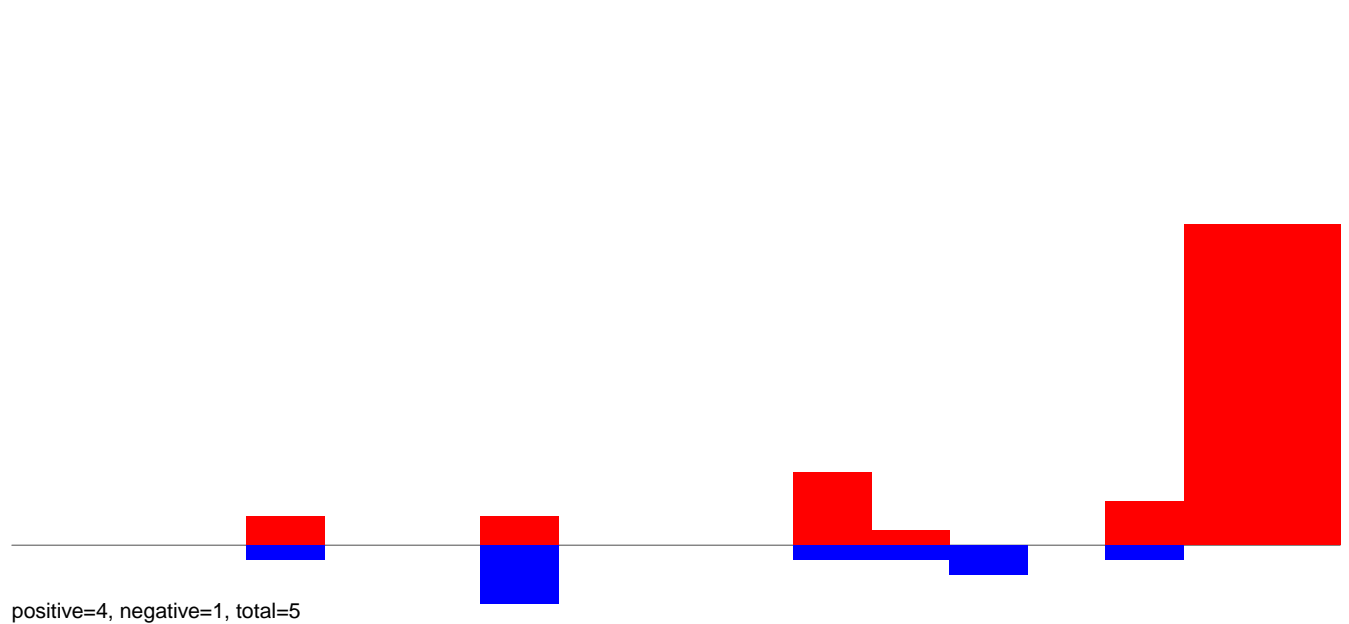
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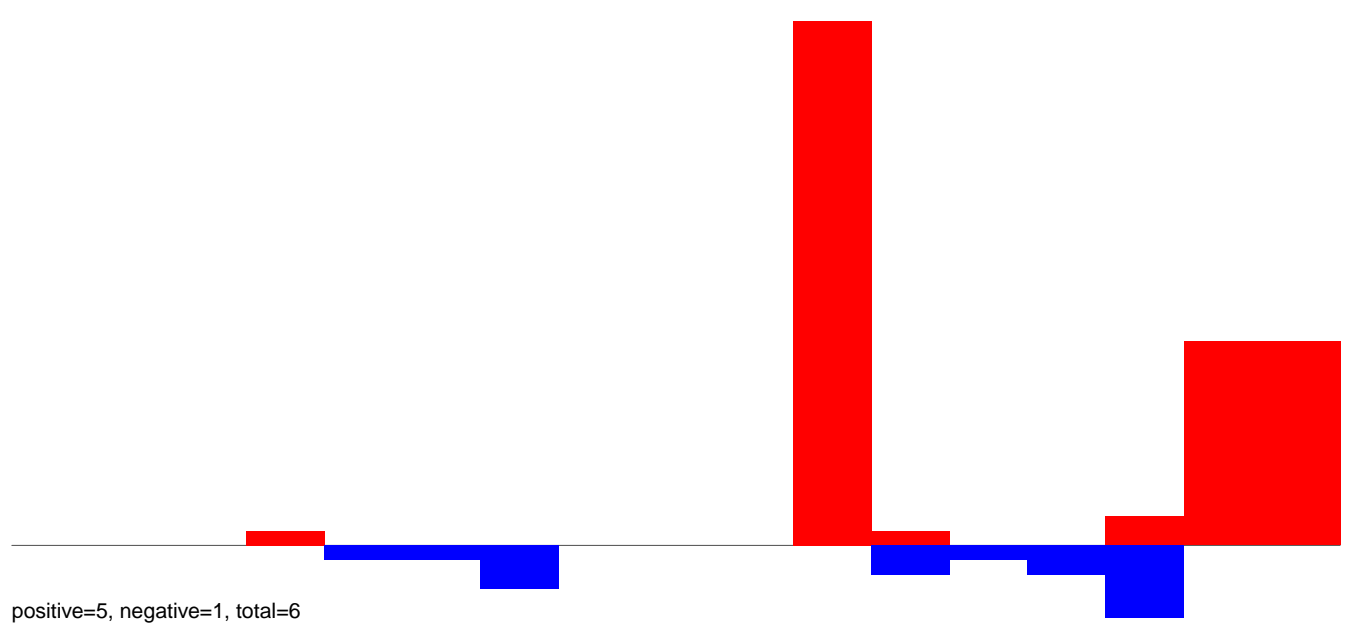
CuQuin_Mcarc_NL.rep



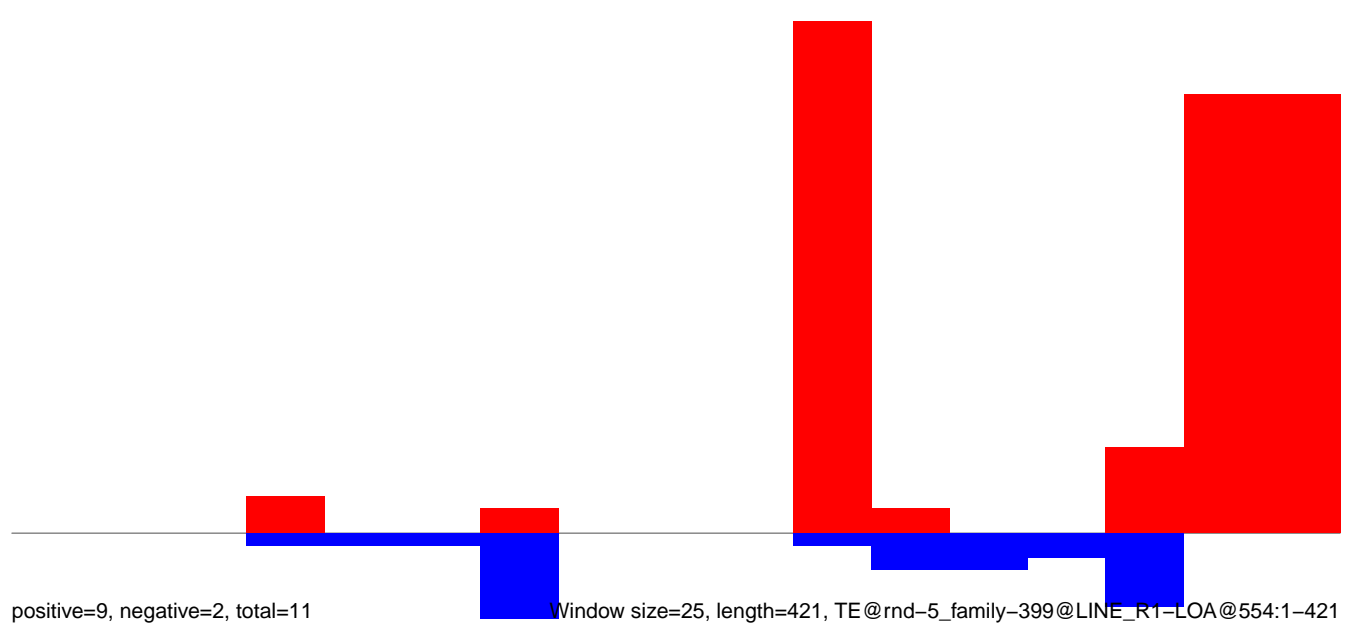
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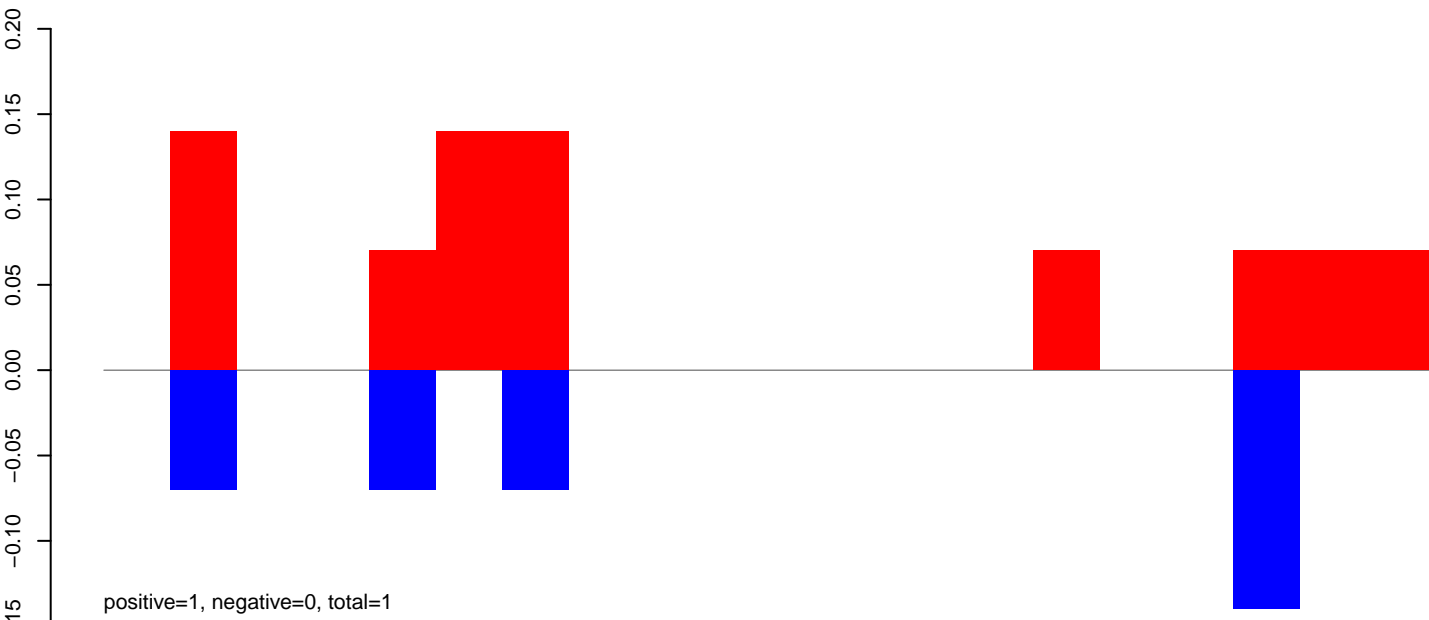
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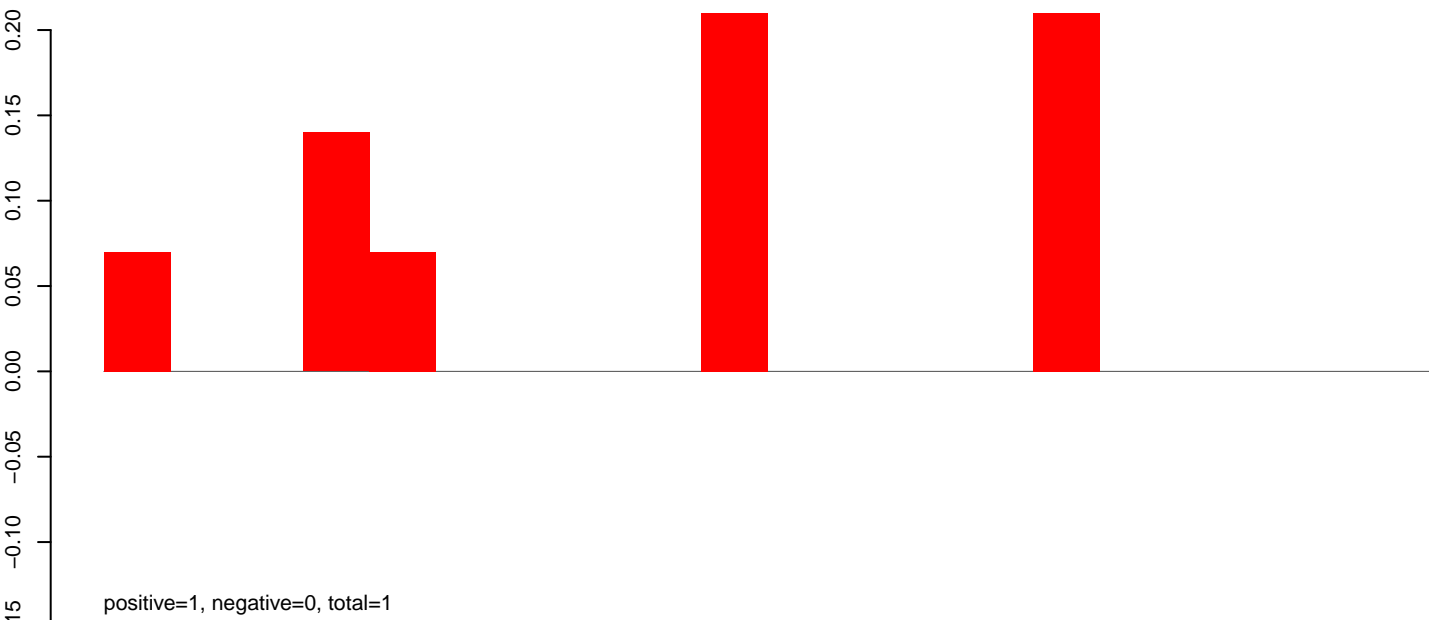
CuQuin_Mcarc_NL.rep



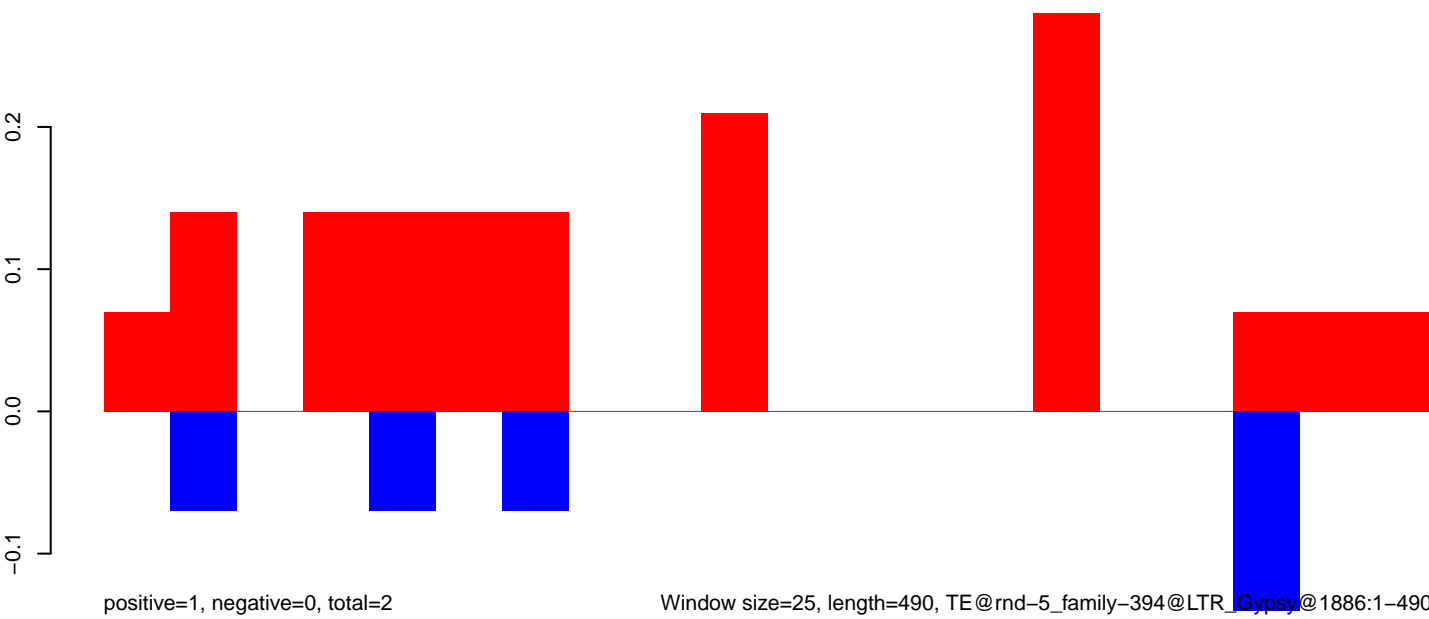
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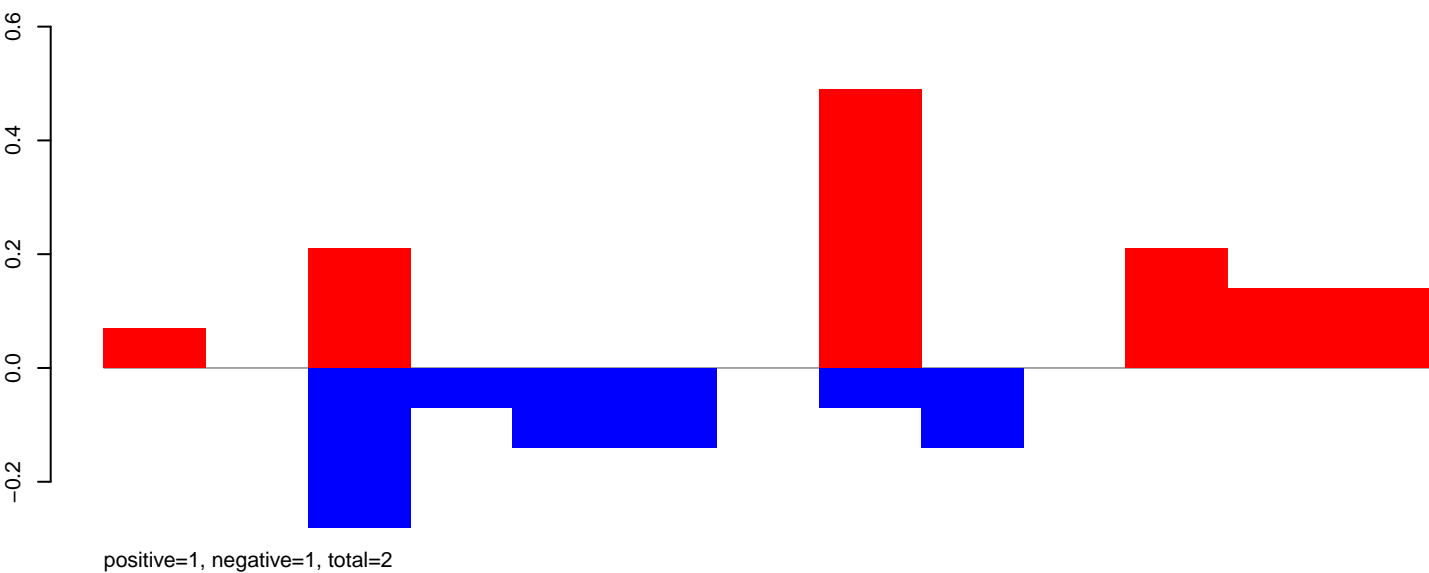
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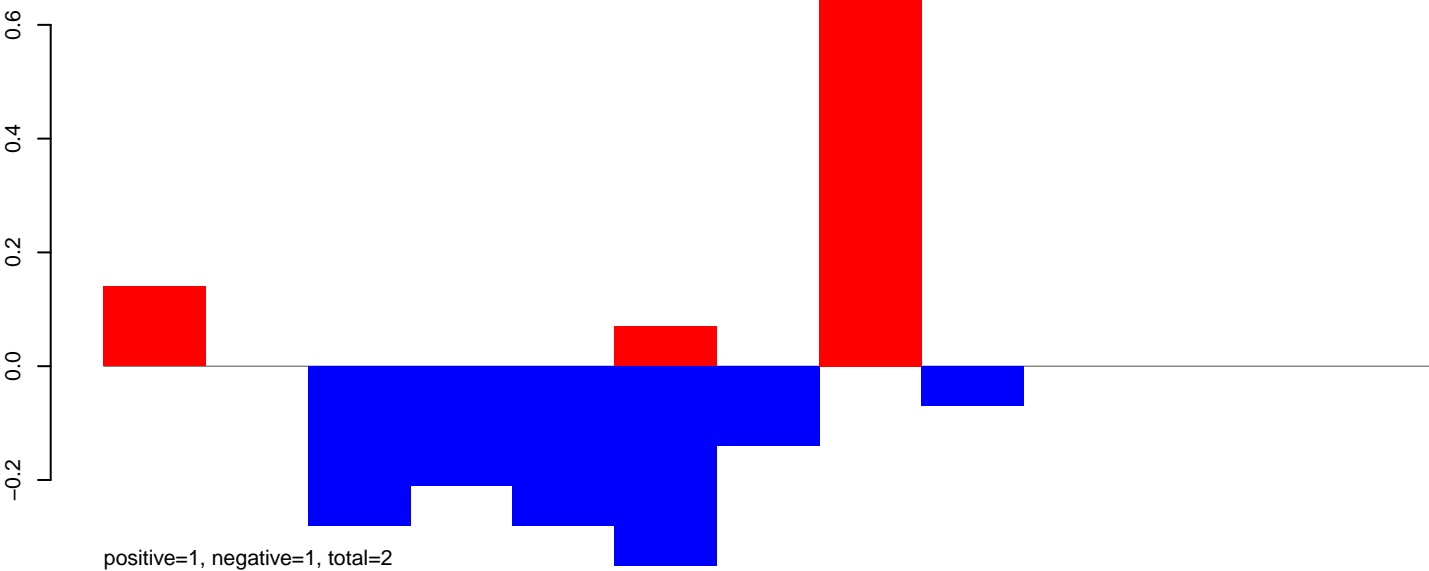
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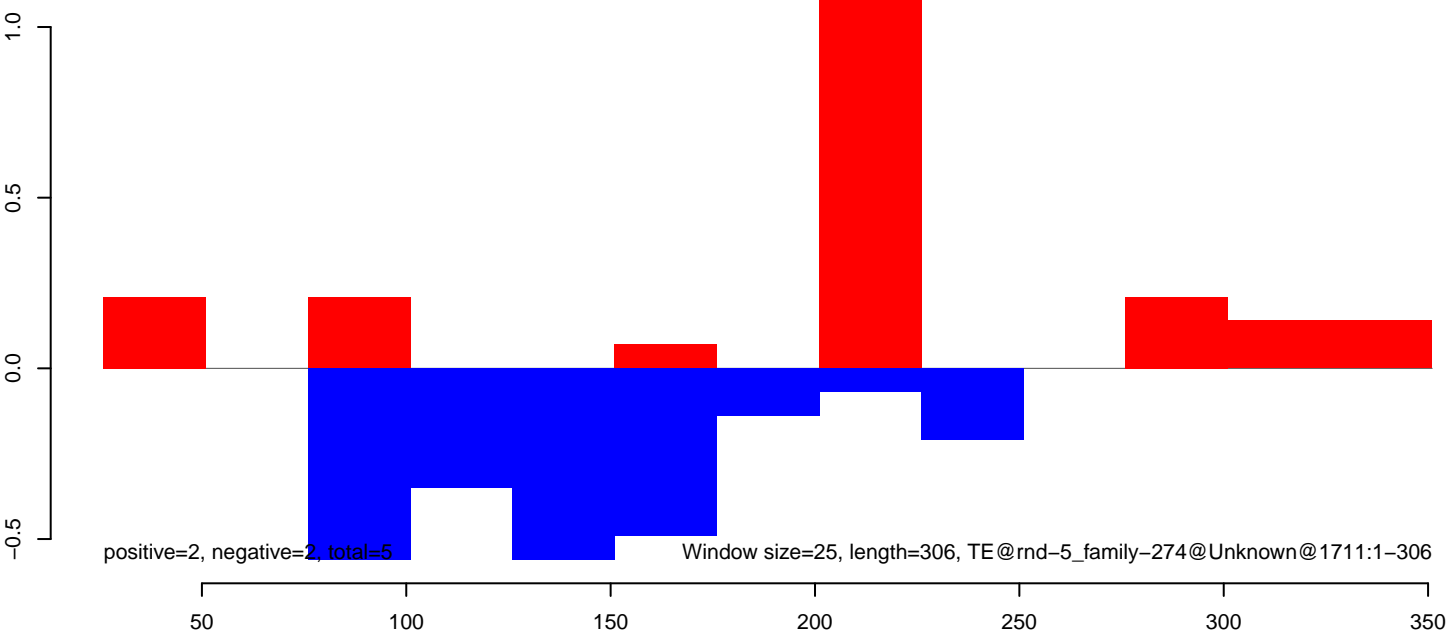
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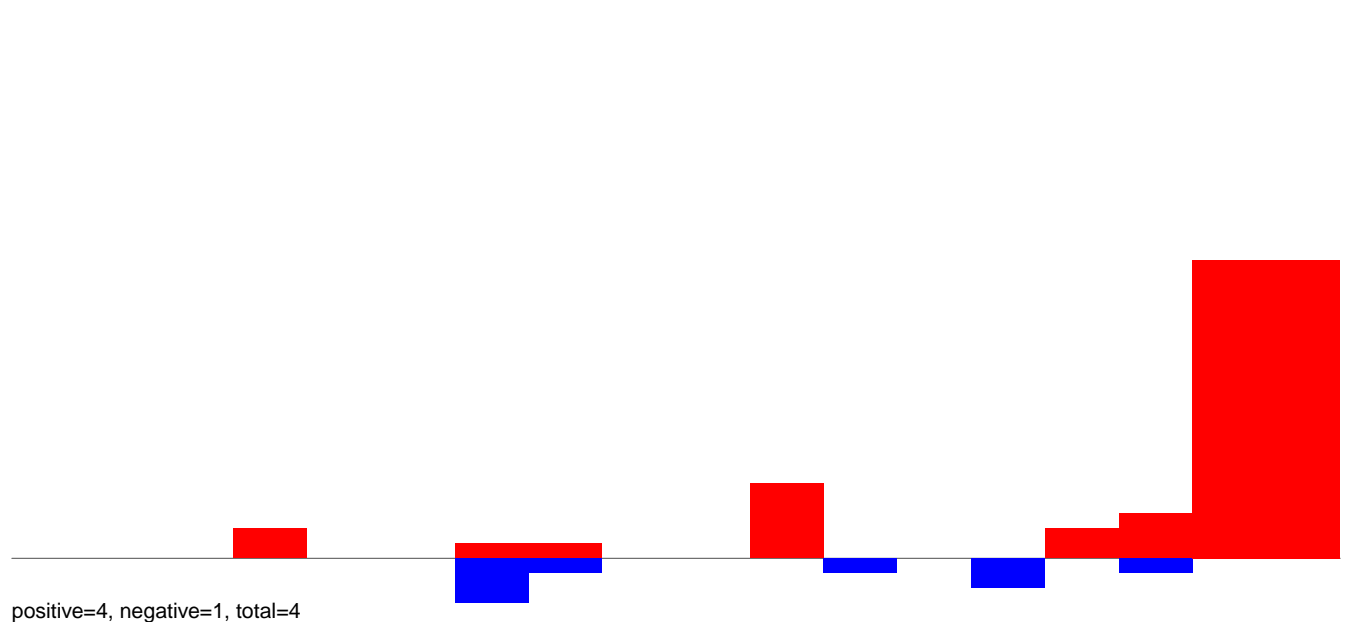
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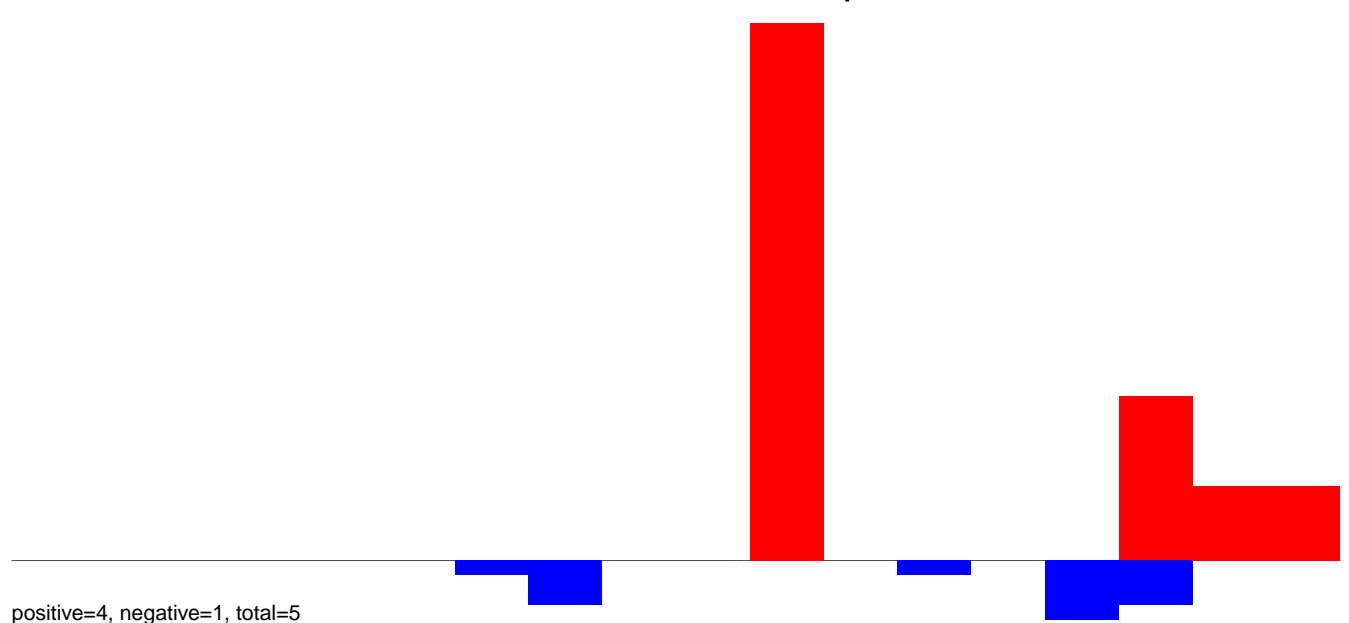
CuQuin_Mcarc_NL.rep



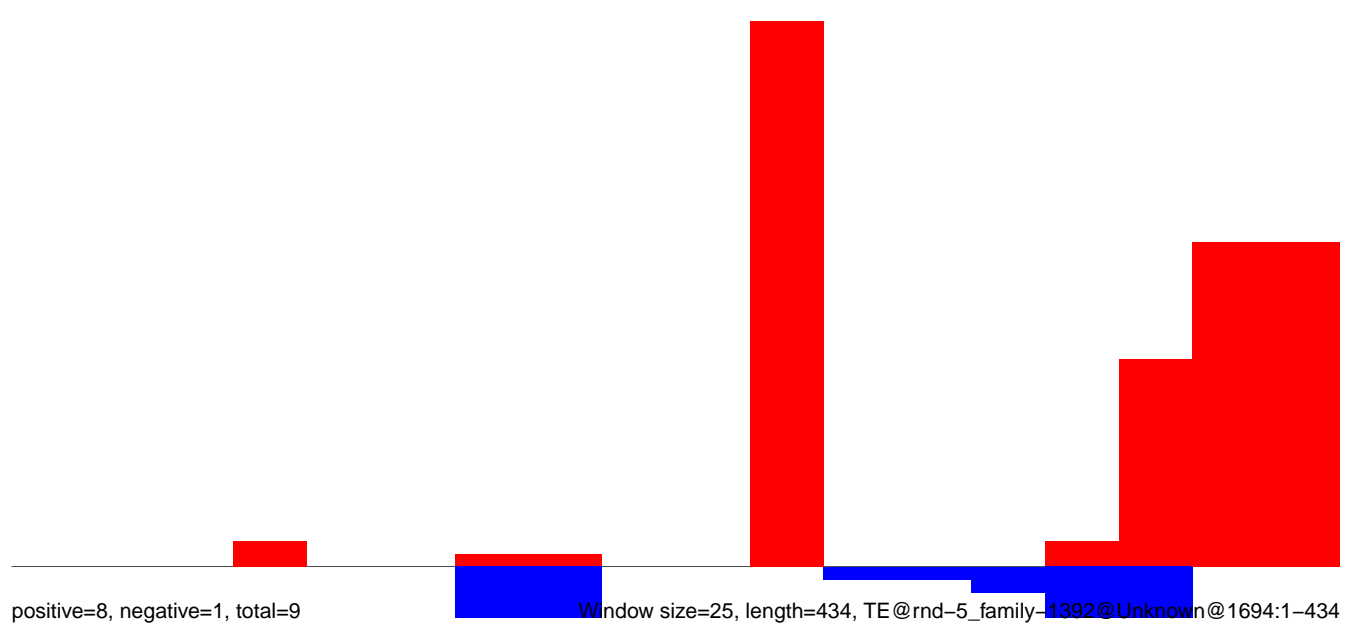
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CuQuin_Mcarc_NL.24_35.rep

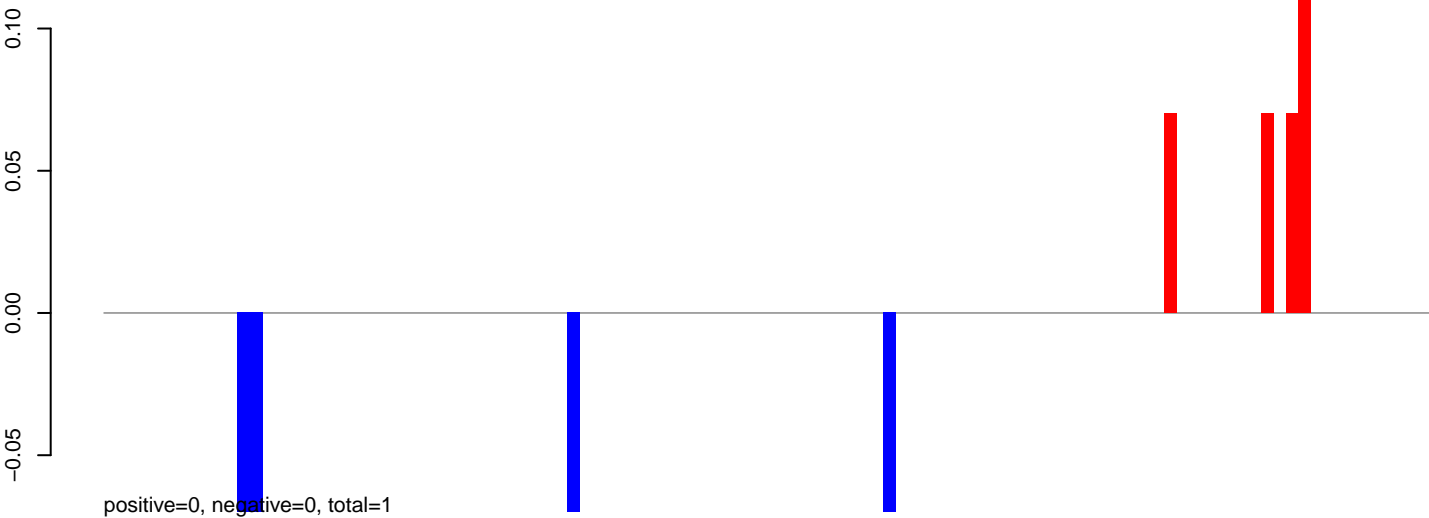


CuQuin_Mcarc_NL.rep

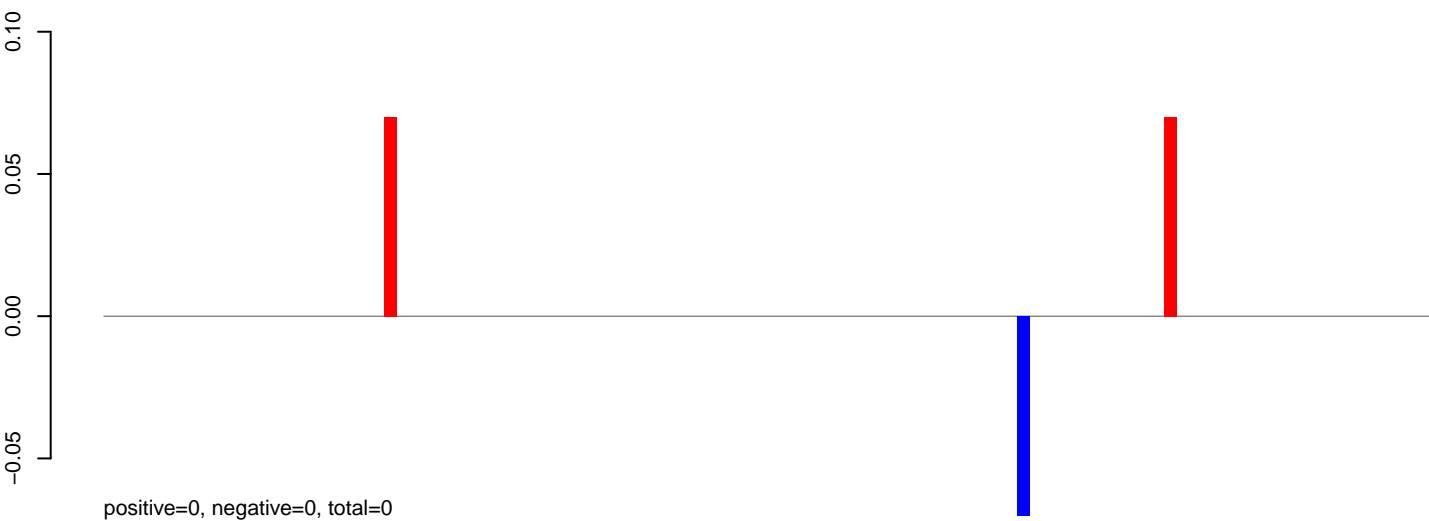


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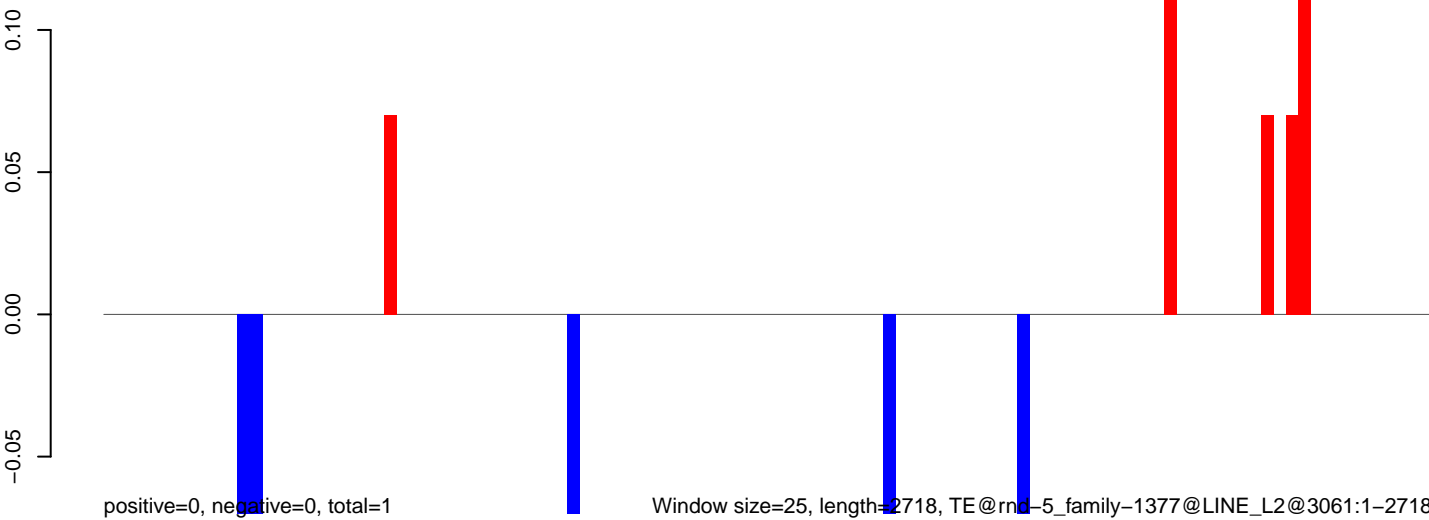
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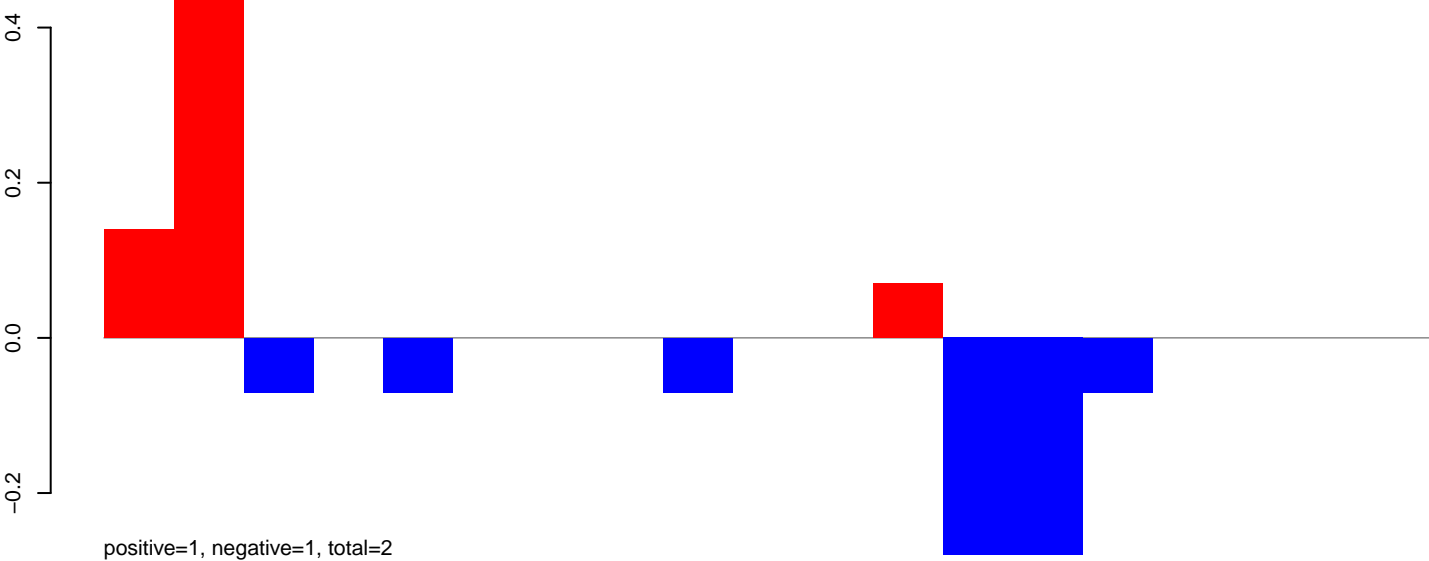
CuQuin_Mcarc_NL.24_35.rep



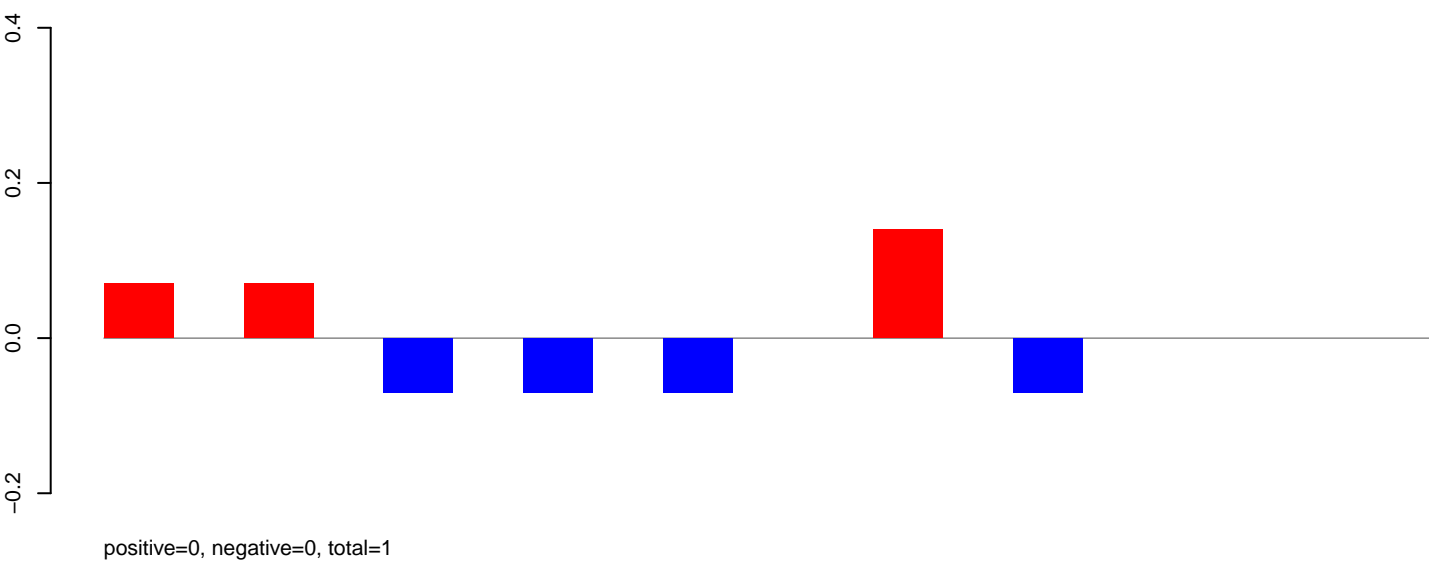
CuQuin_Mcarc_NL.rep



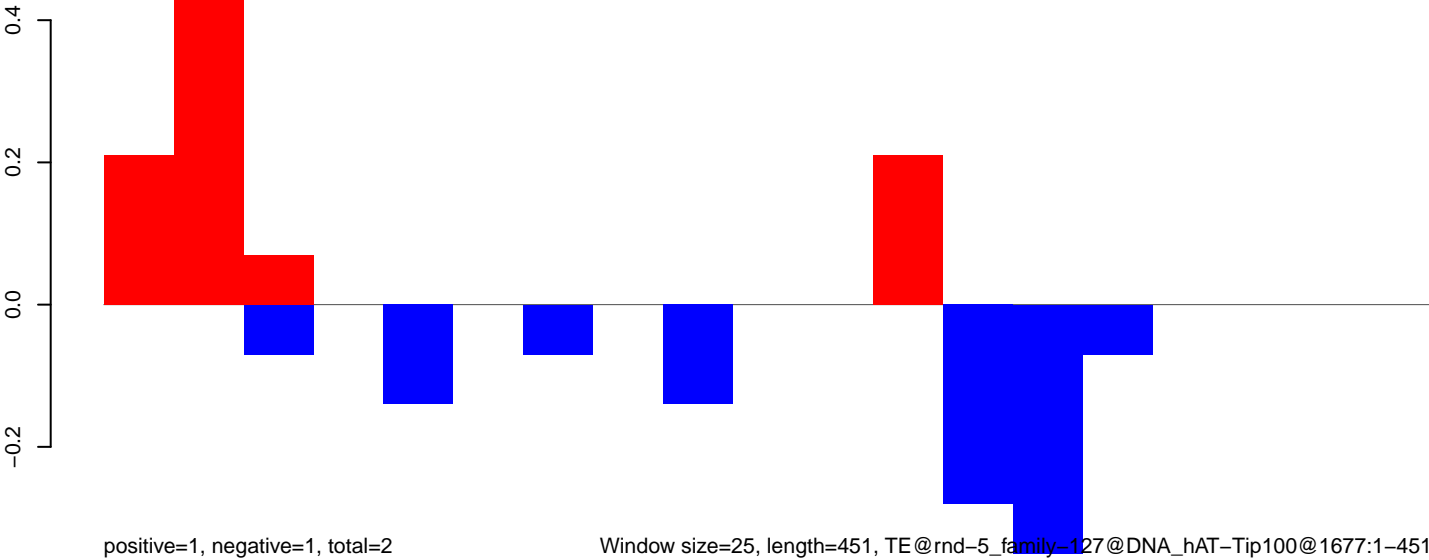
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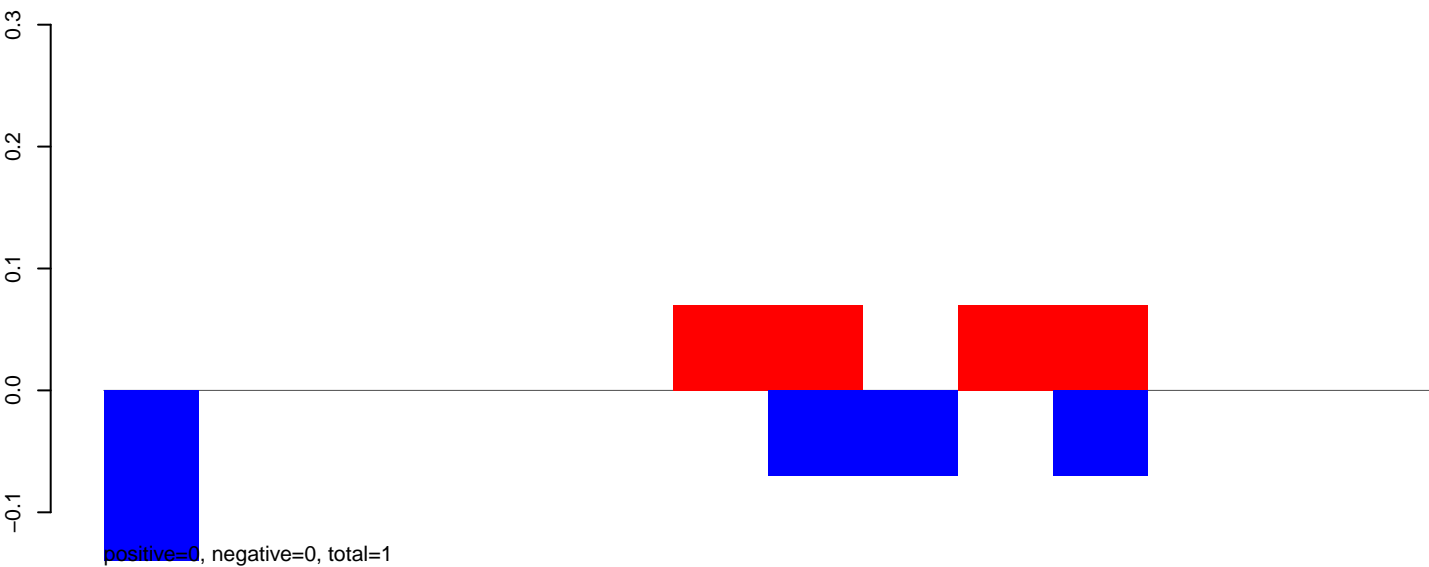
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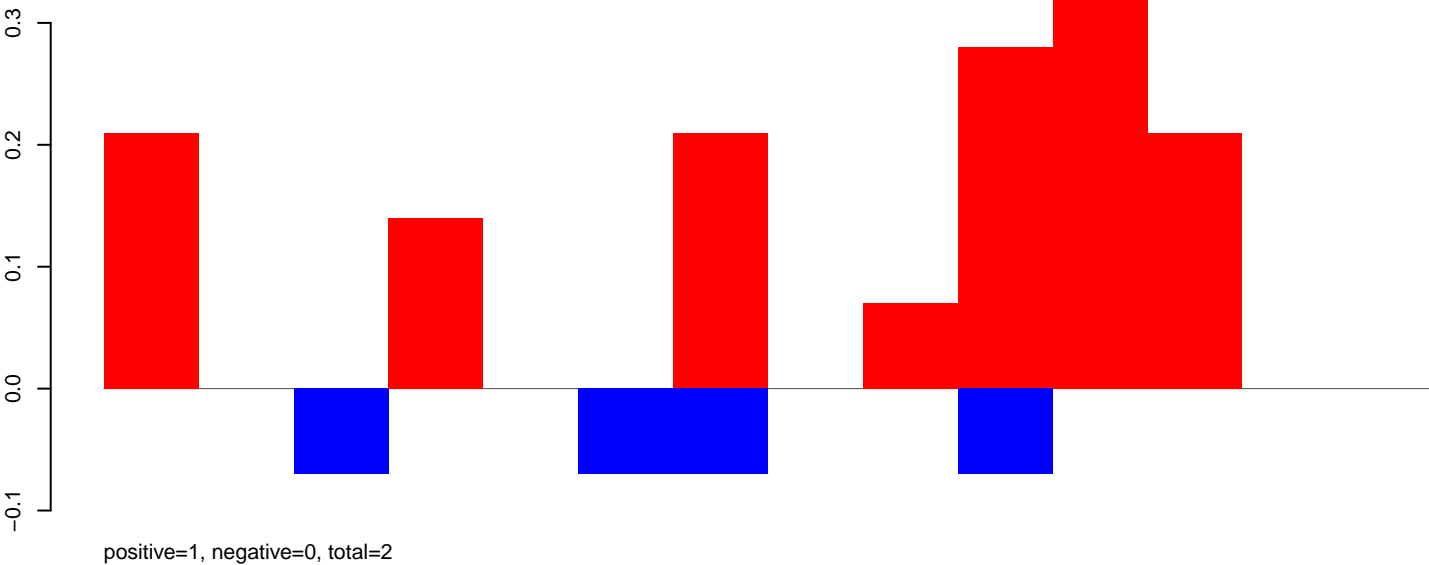
CuQuin_Mcarc_NL.rep



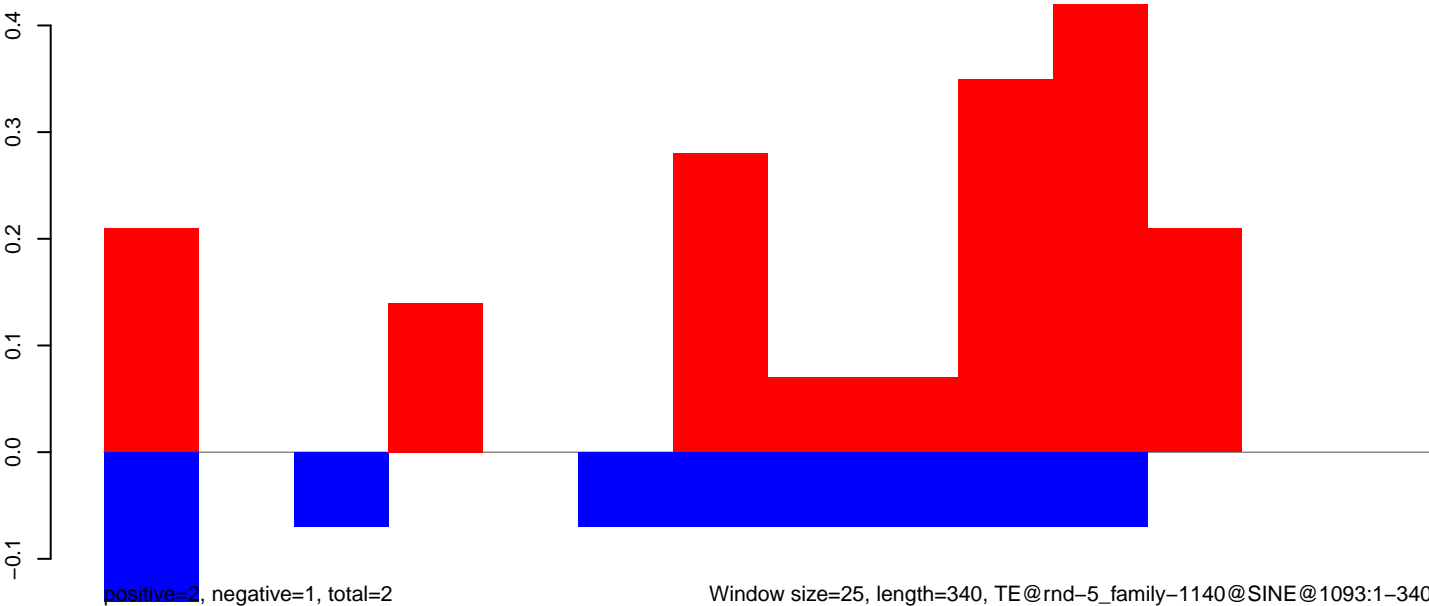
CuQuin_Mcarc_NL.18_23.rep

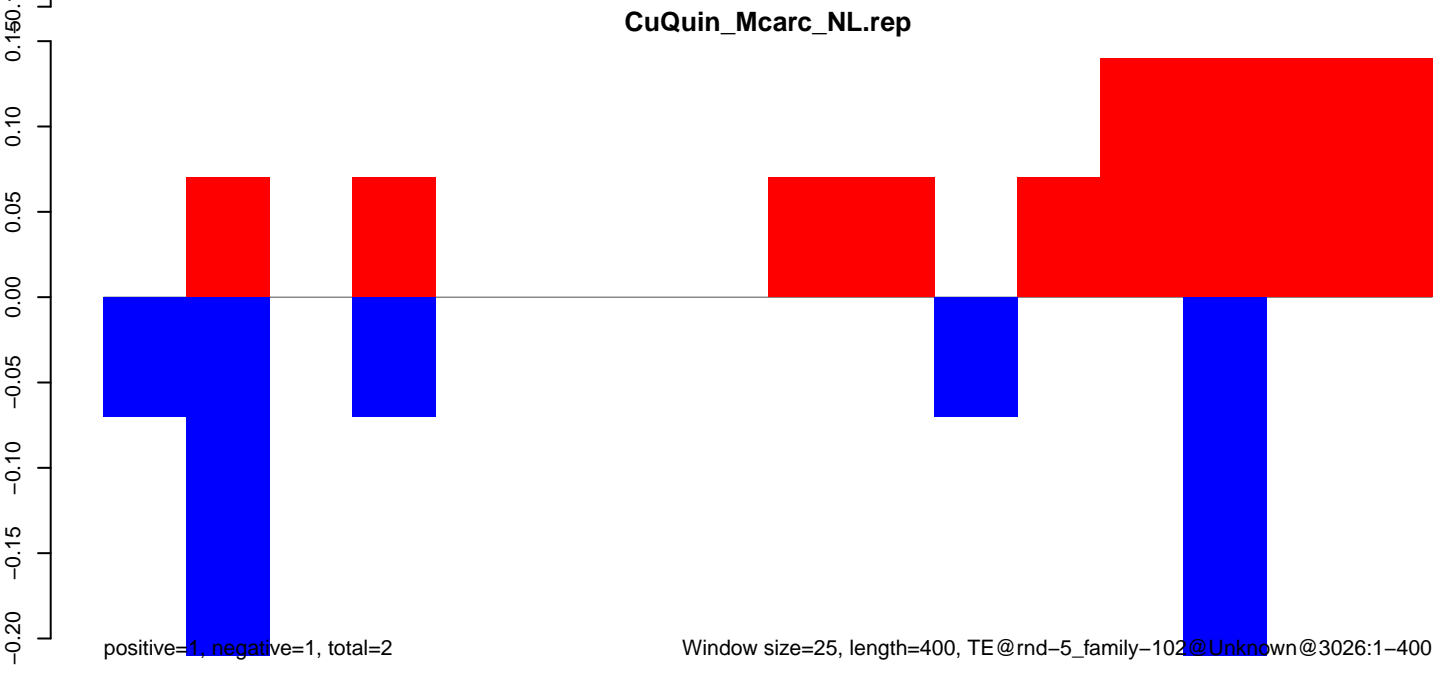
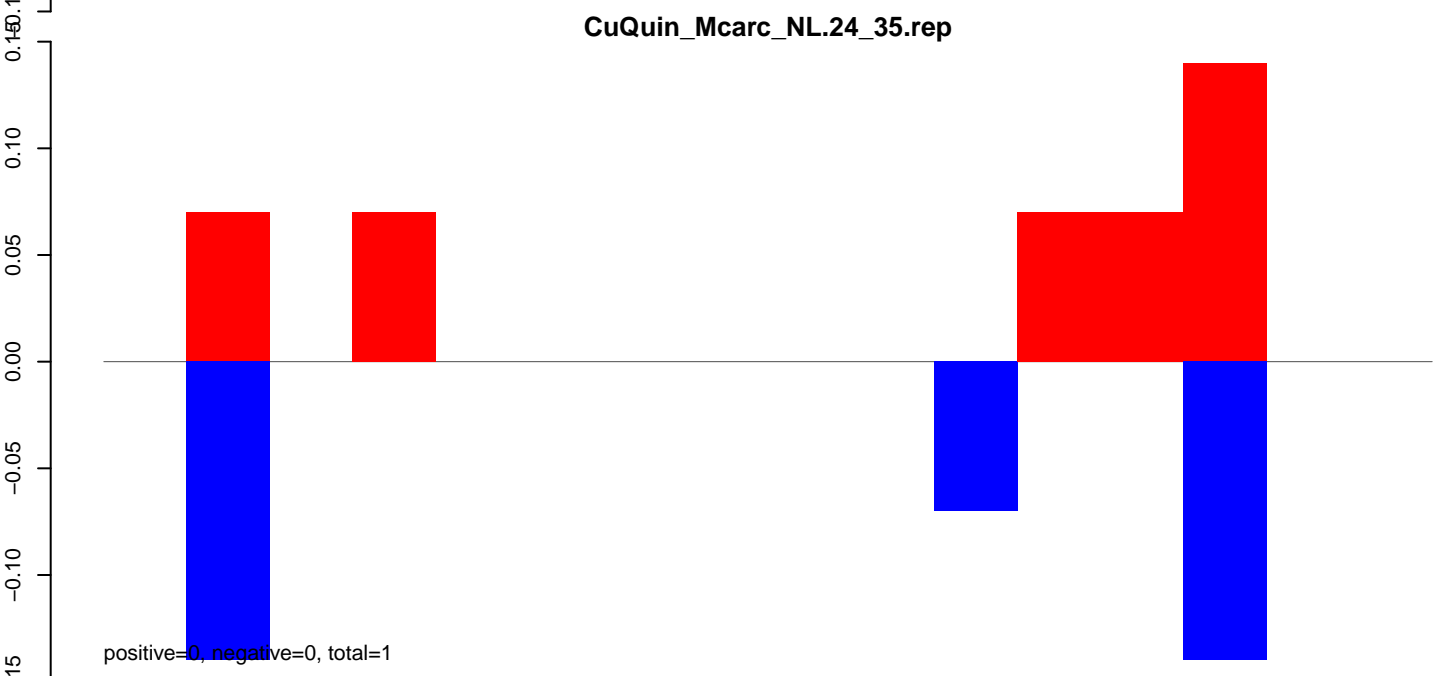
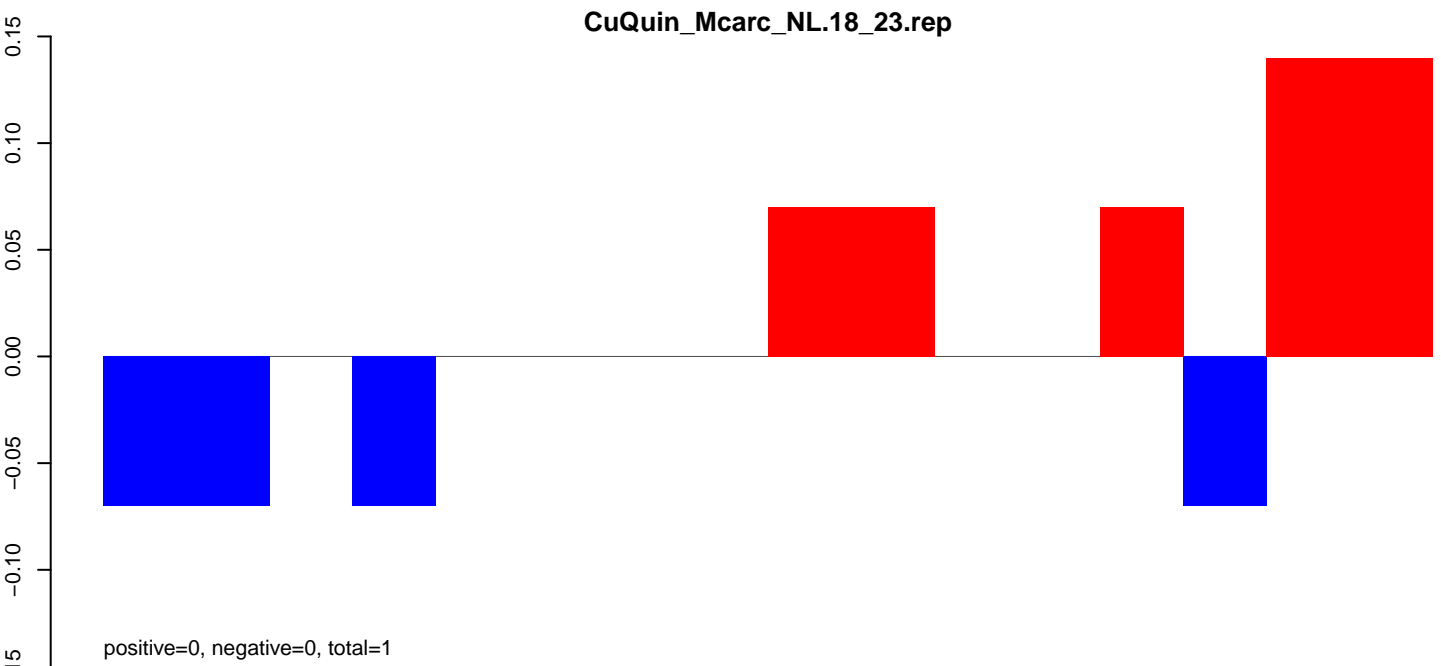


CuQuin_Mcarc_NL.24_35.rep

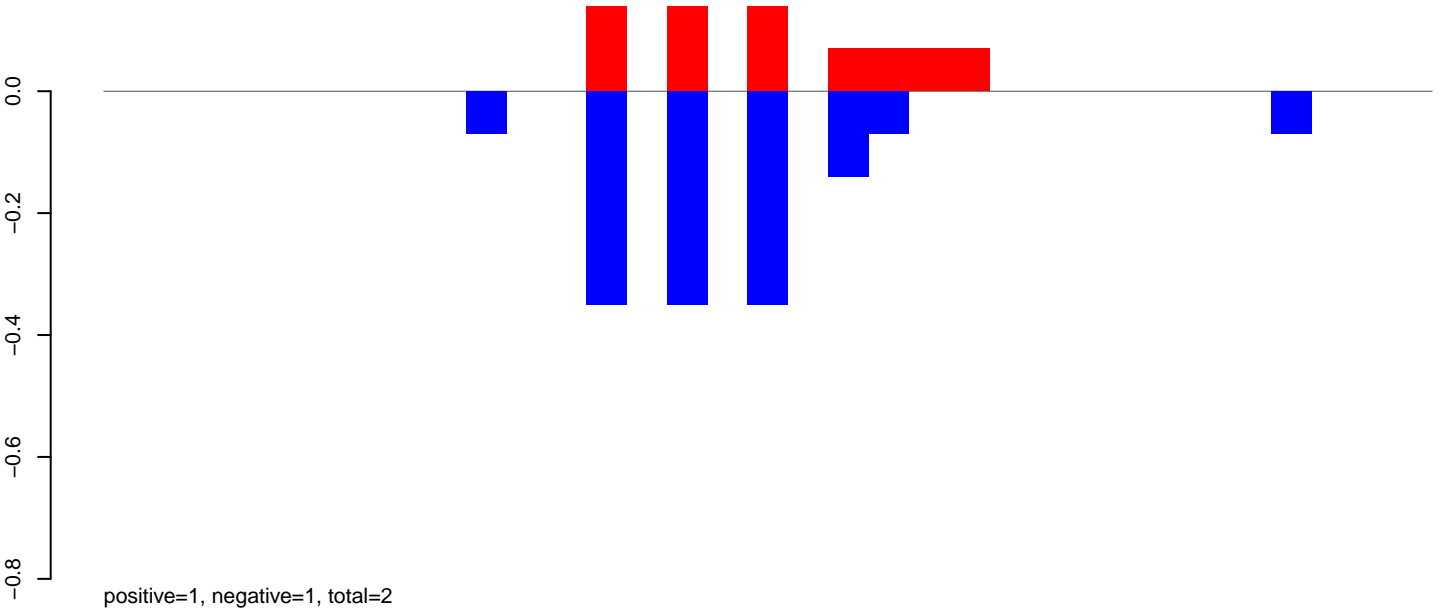


CuQuin_Mcarc_NL.rep

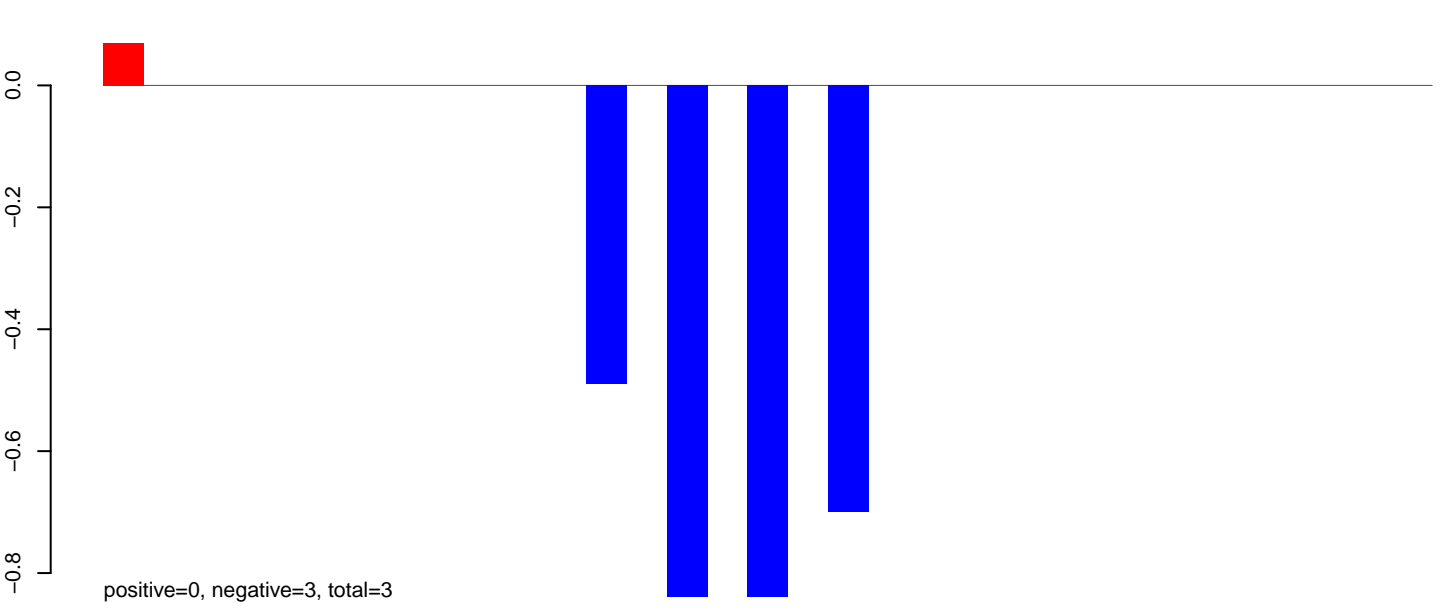




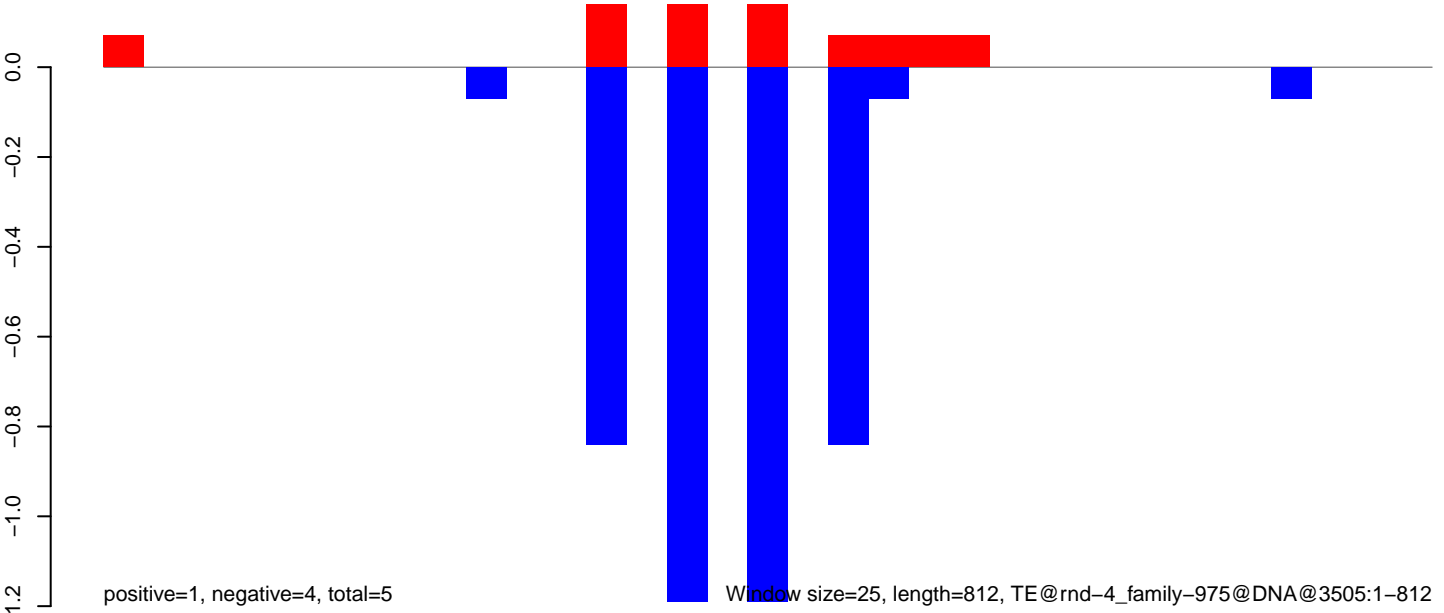
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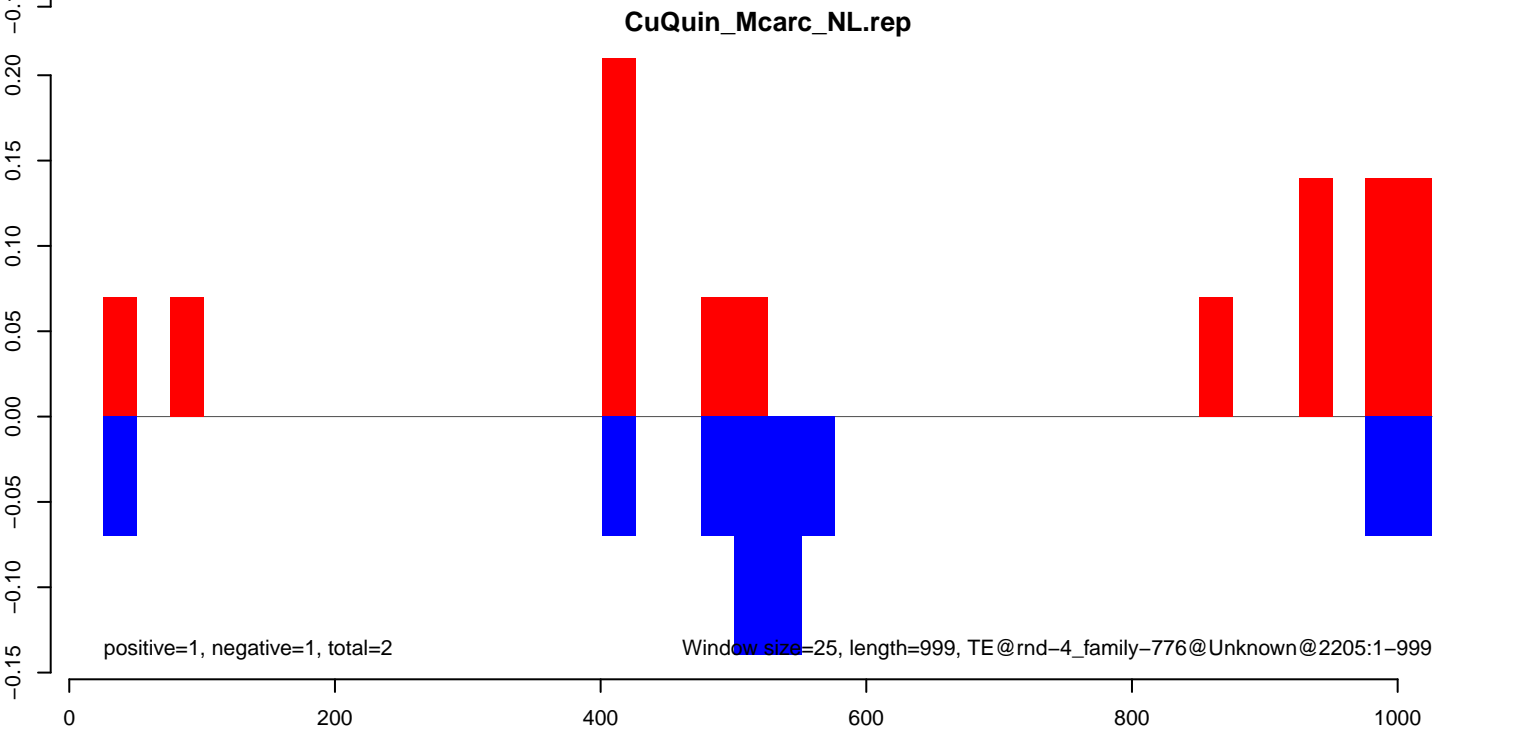
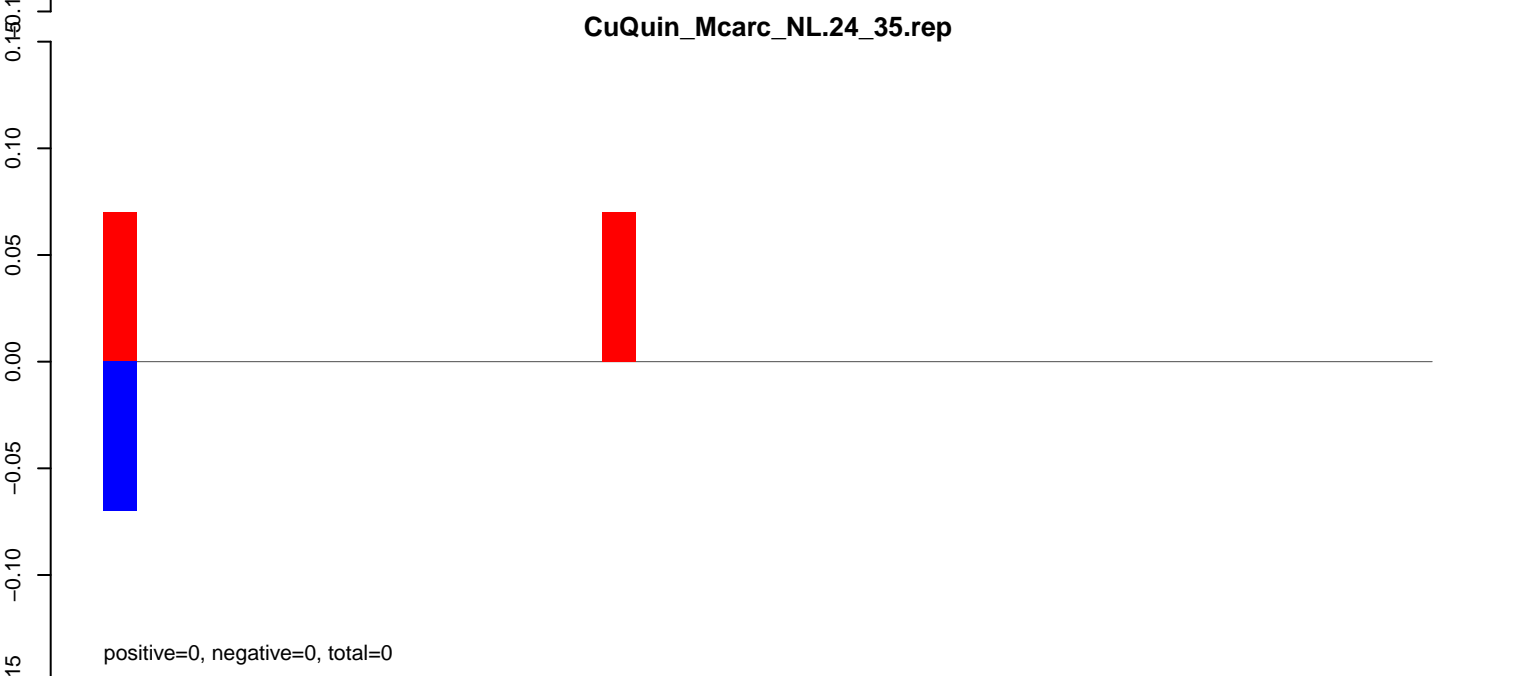
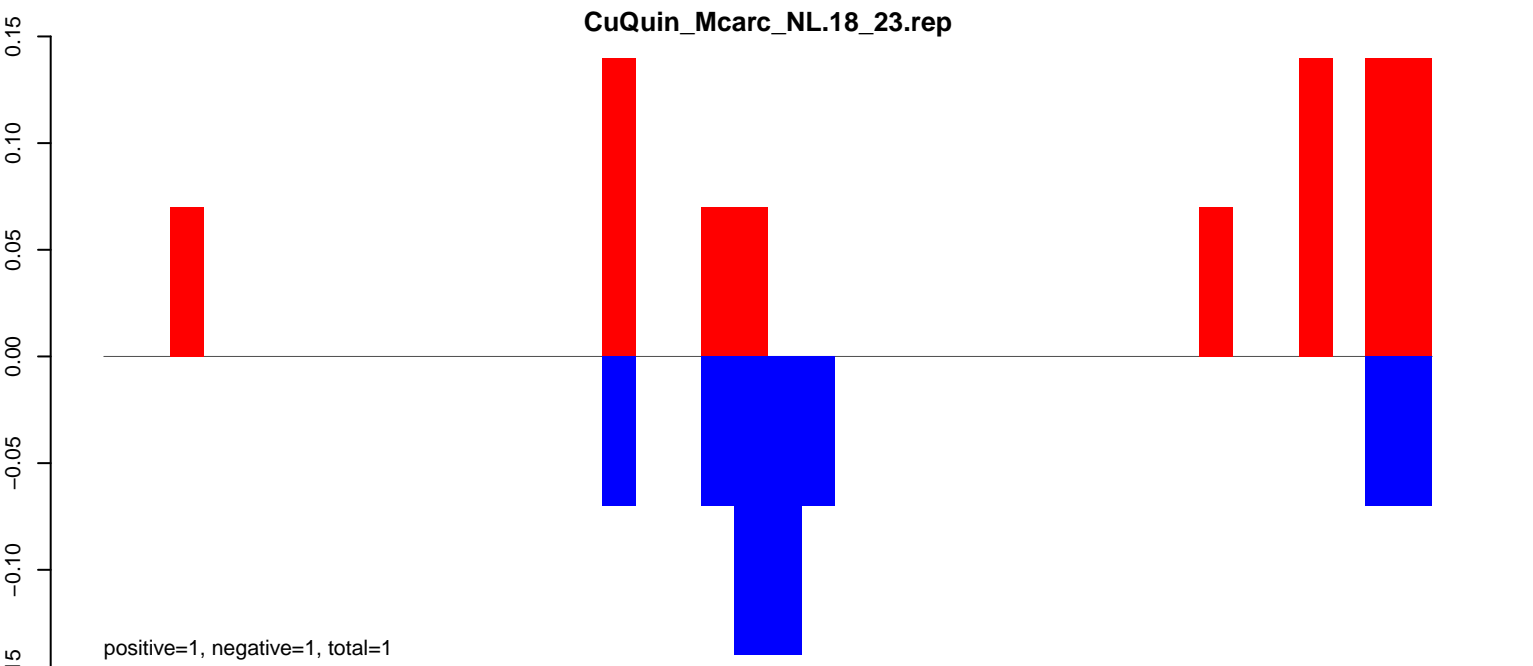
CuQuin_Mcarc_NL.24_35.rep



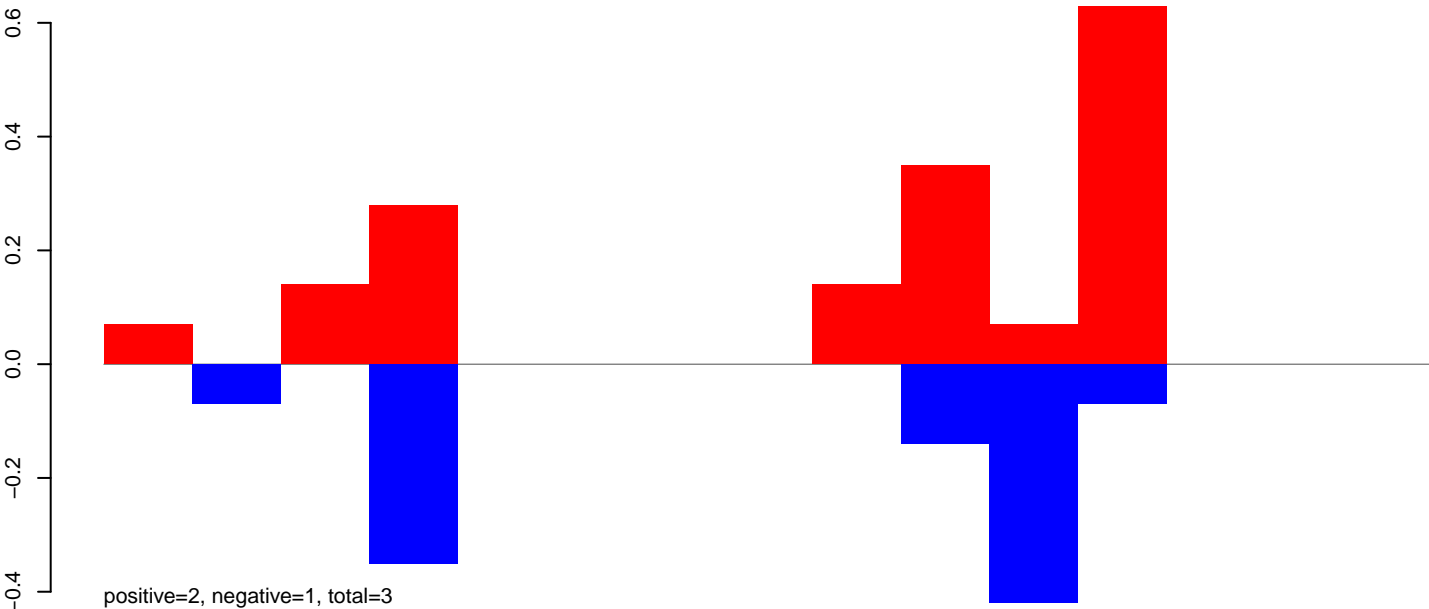
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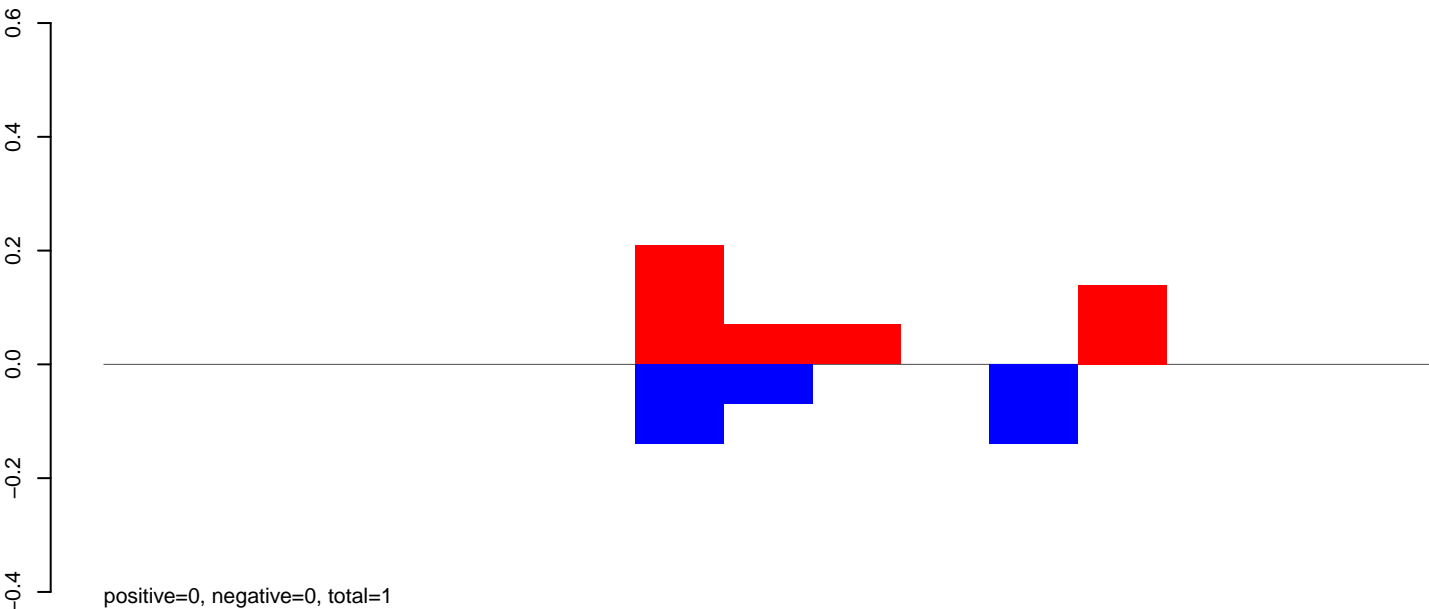
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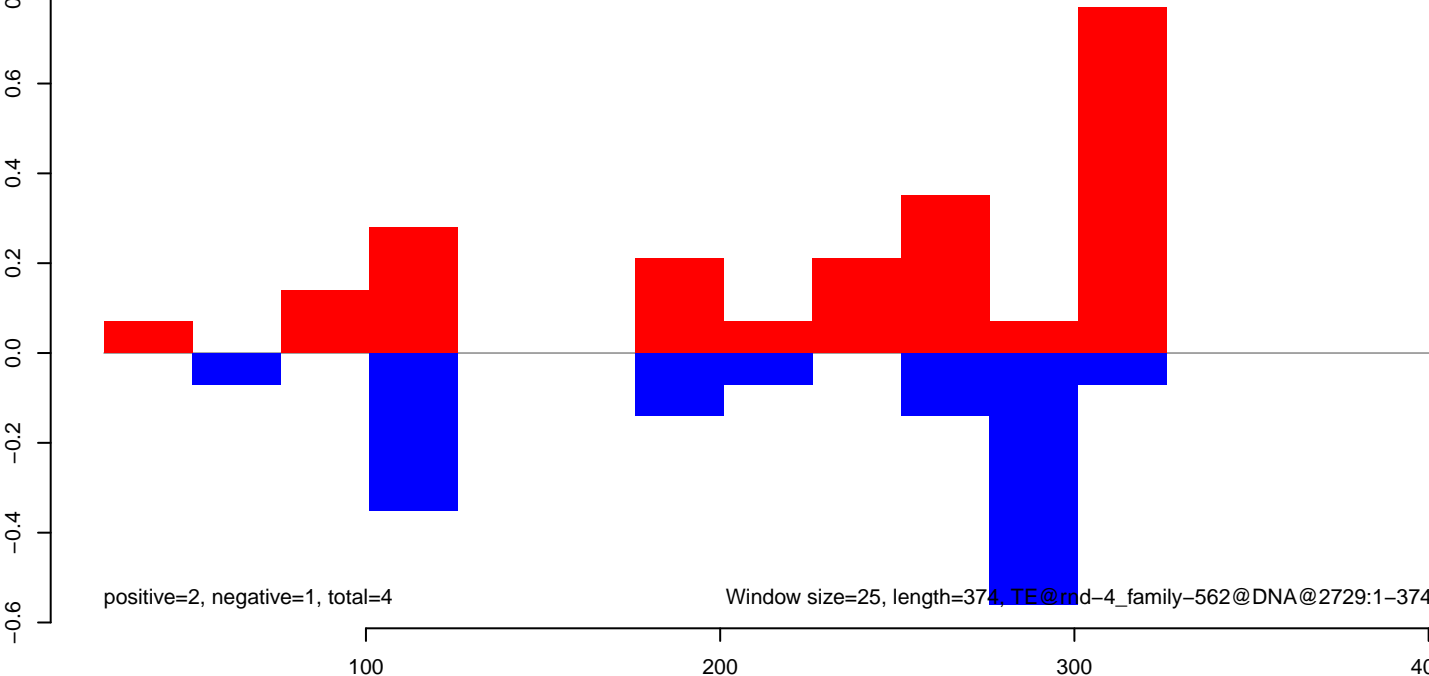
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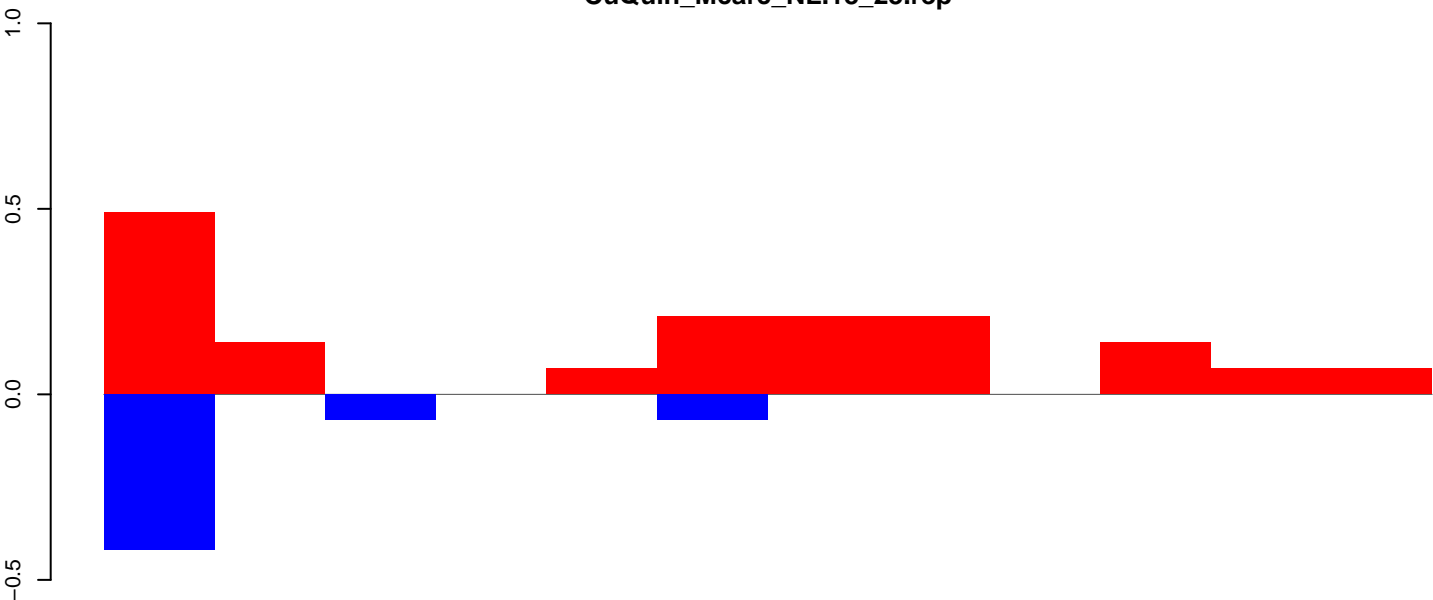
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

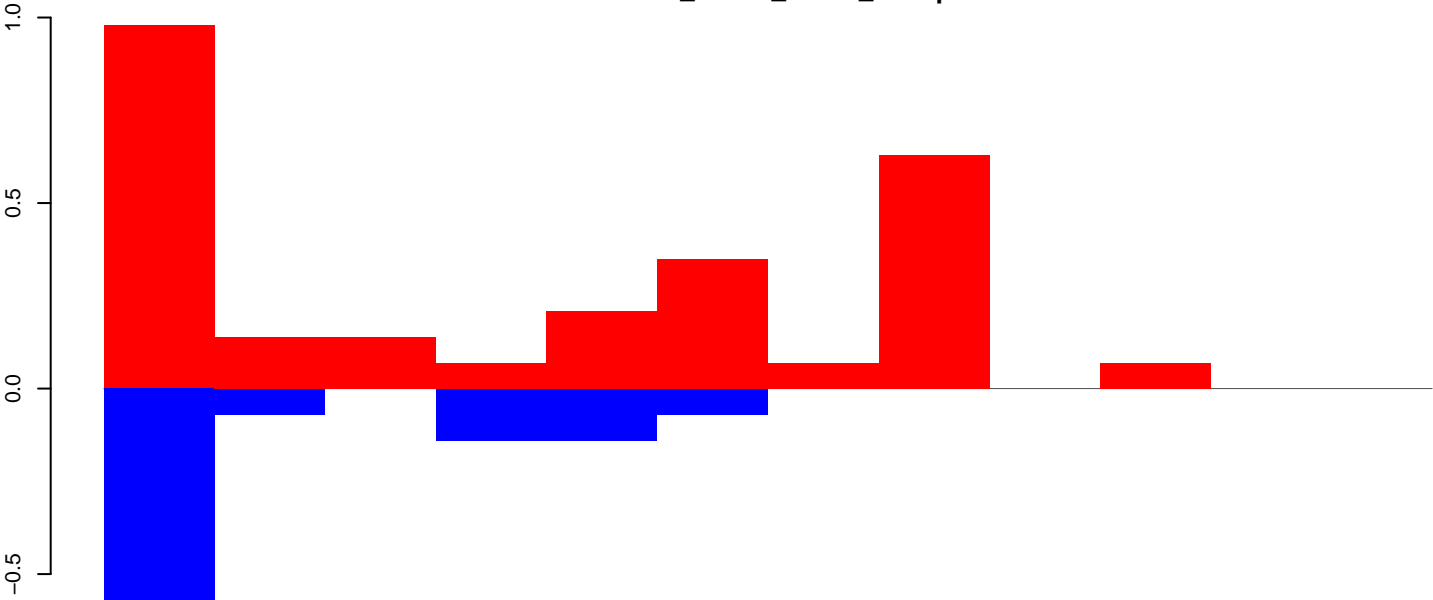


CuQuin_Mcarc_NL.18_23.rep



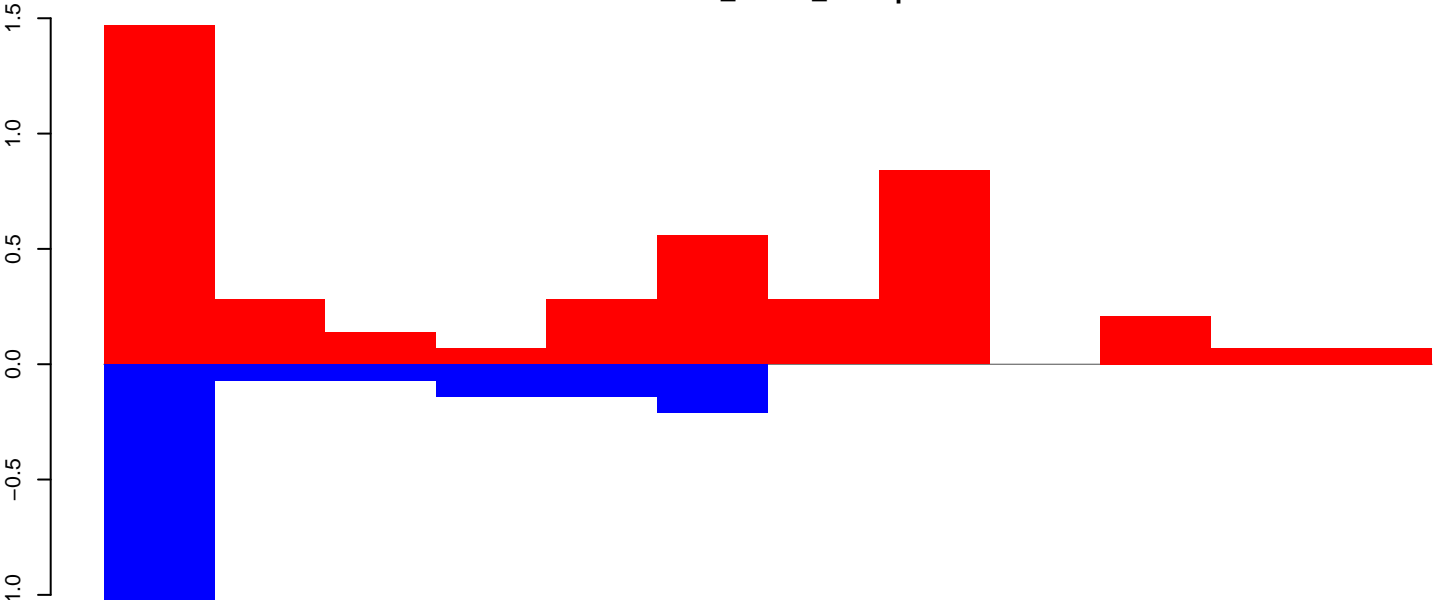
positive=2, negative=1, total=2

CuQuin_Mcarc_NL.24_35.rep



positive=3, negative=1, total=4

CuQuin_Mcarc_NL.rep

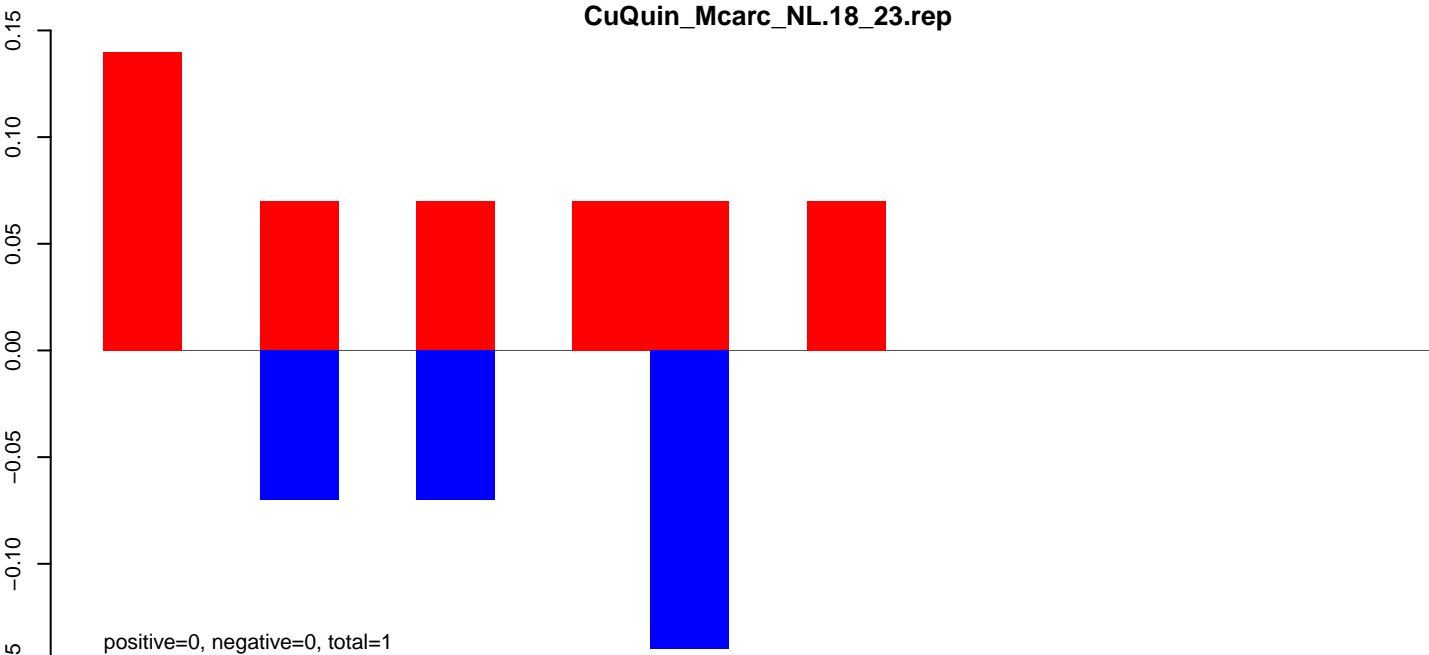


positive=4, negative=2, total=6

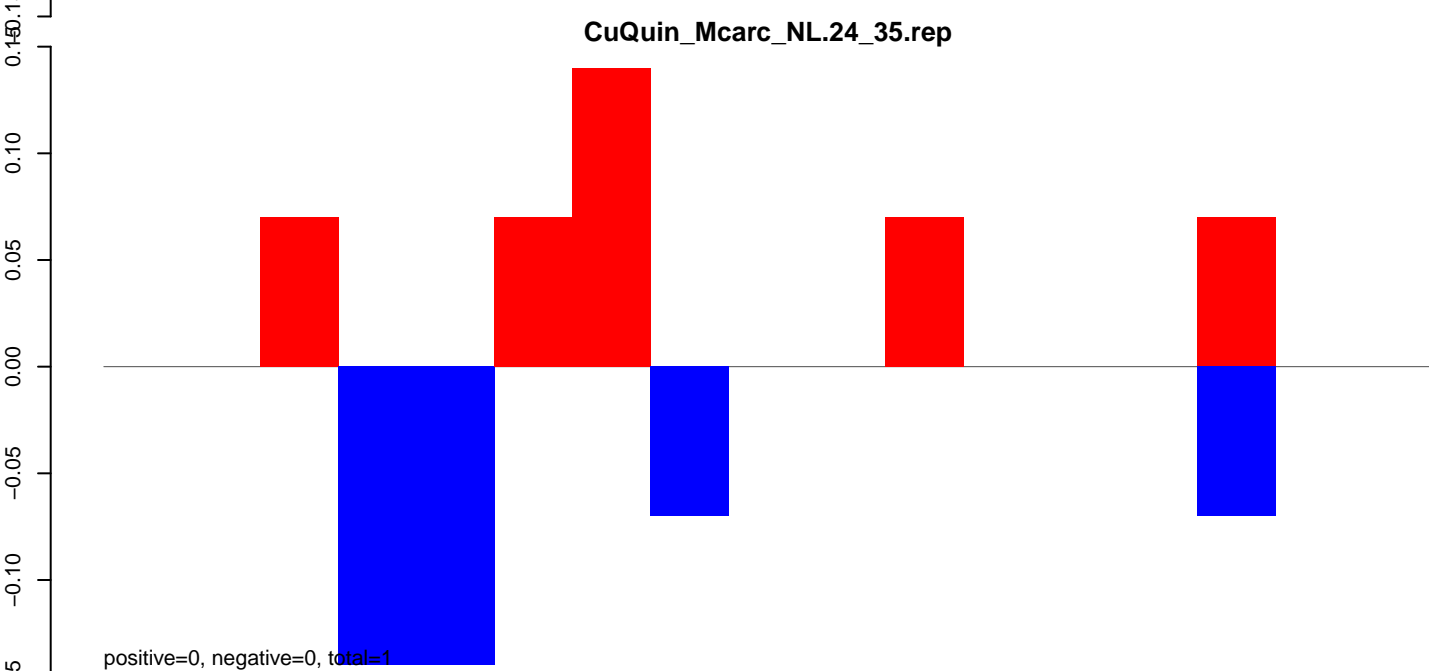
Window size=25, length=277, TE@rnd-4_family-51@Unknown@970:1-277

50 100 150 200 250 300

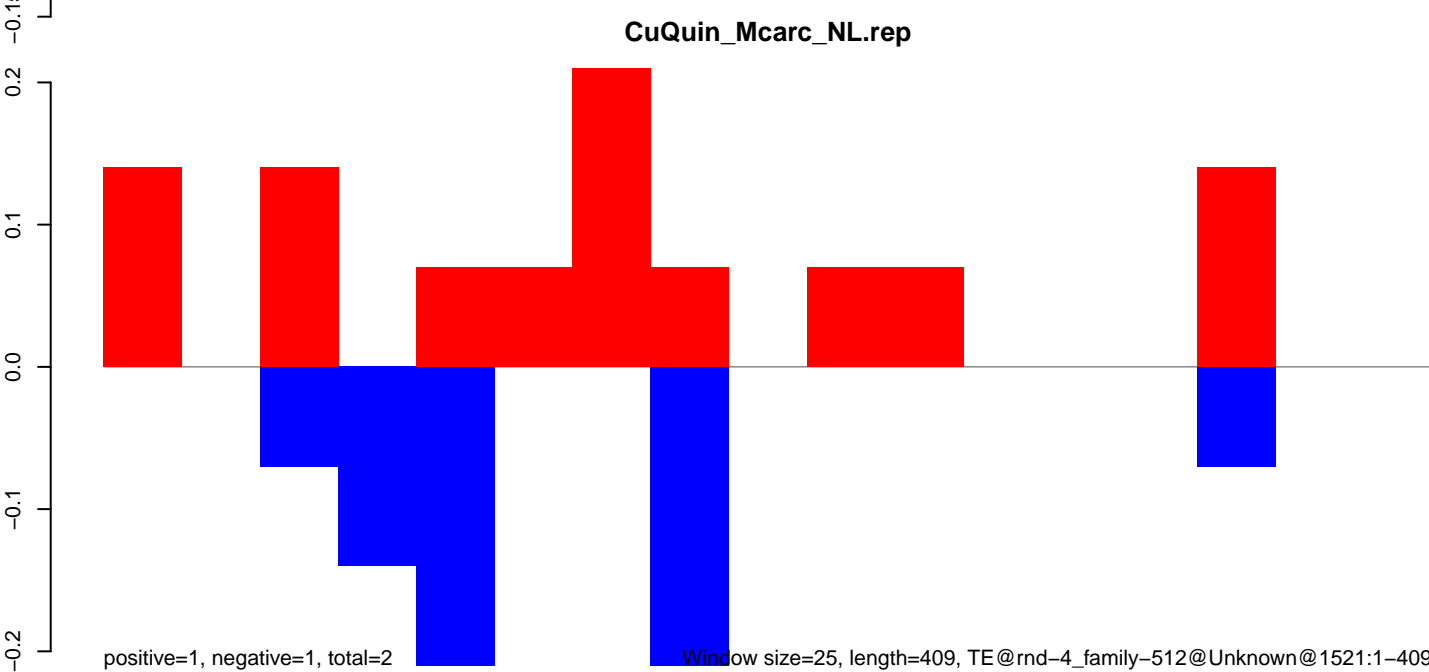
CuQuin_Mcarc_NL.18_23.rep



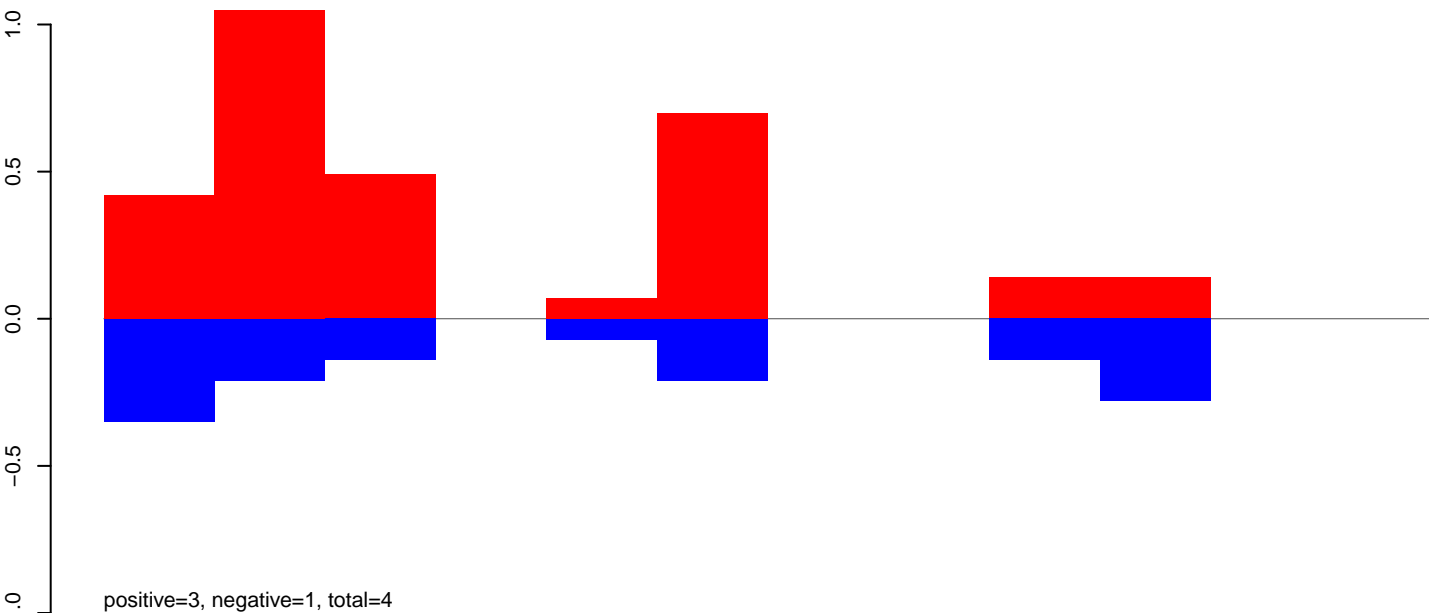
CuQuin_Mcarc_NL.24_35.rep



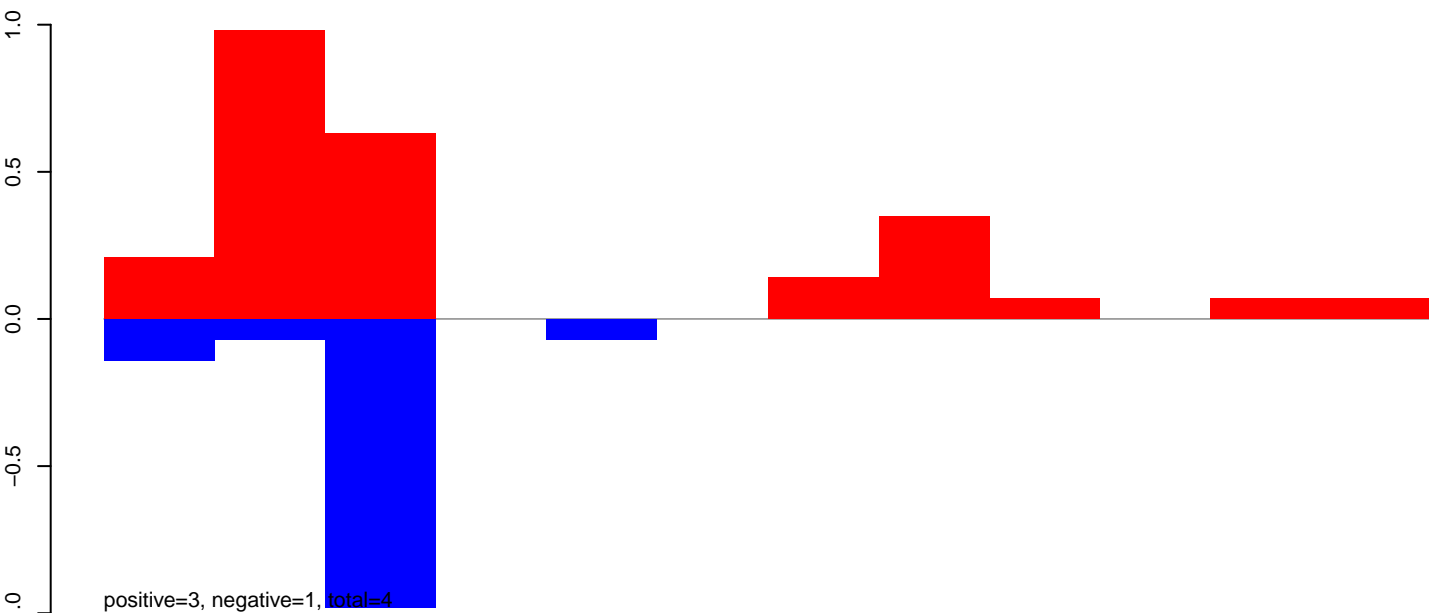
CuQuin_Mcarc_NL.rep



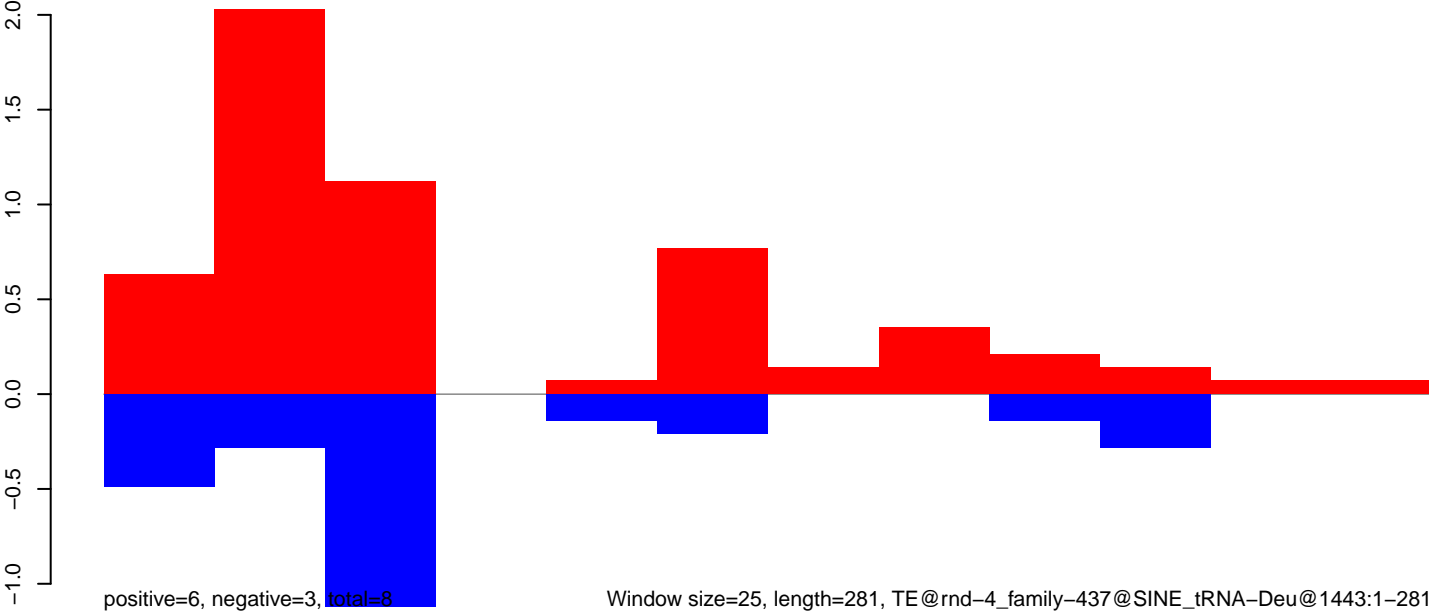
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

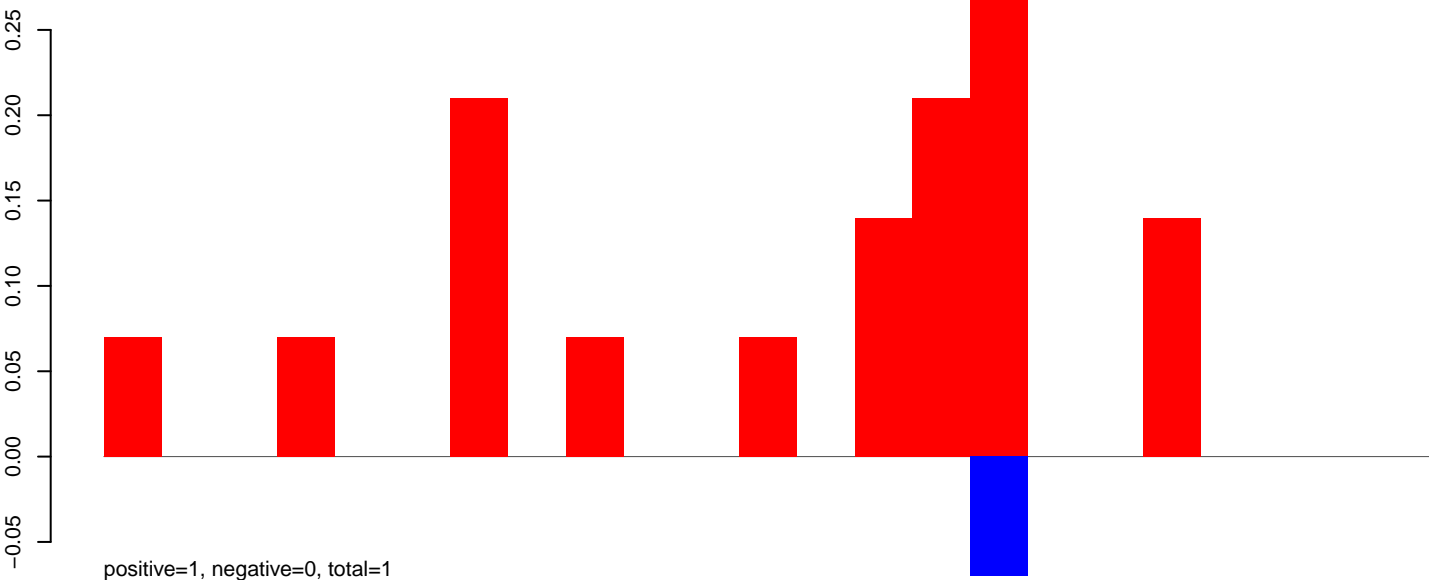


CuQuin_Mcarc_NL.rep

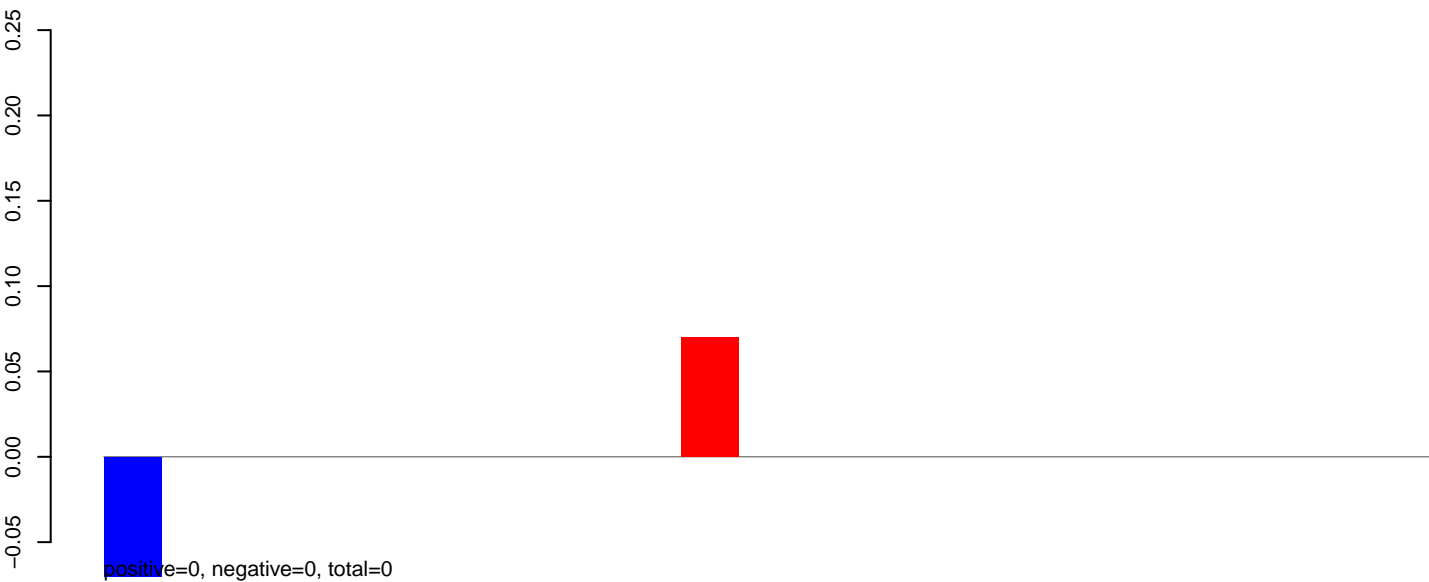


Window size=25, length=281, TE@rnd-4_family-437@SINE_tRNA-Deu@1443:1-281

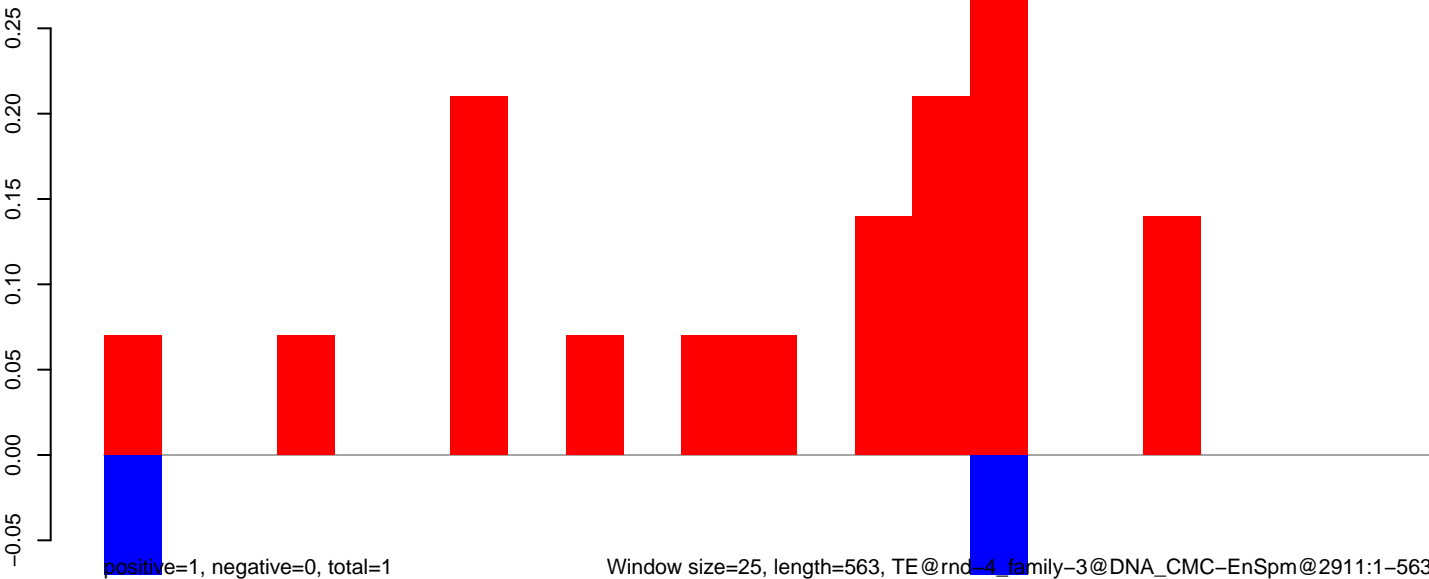
CuQuin_Mcarc_NL.18_23.rep



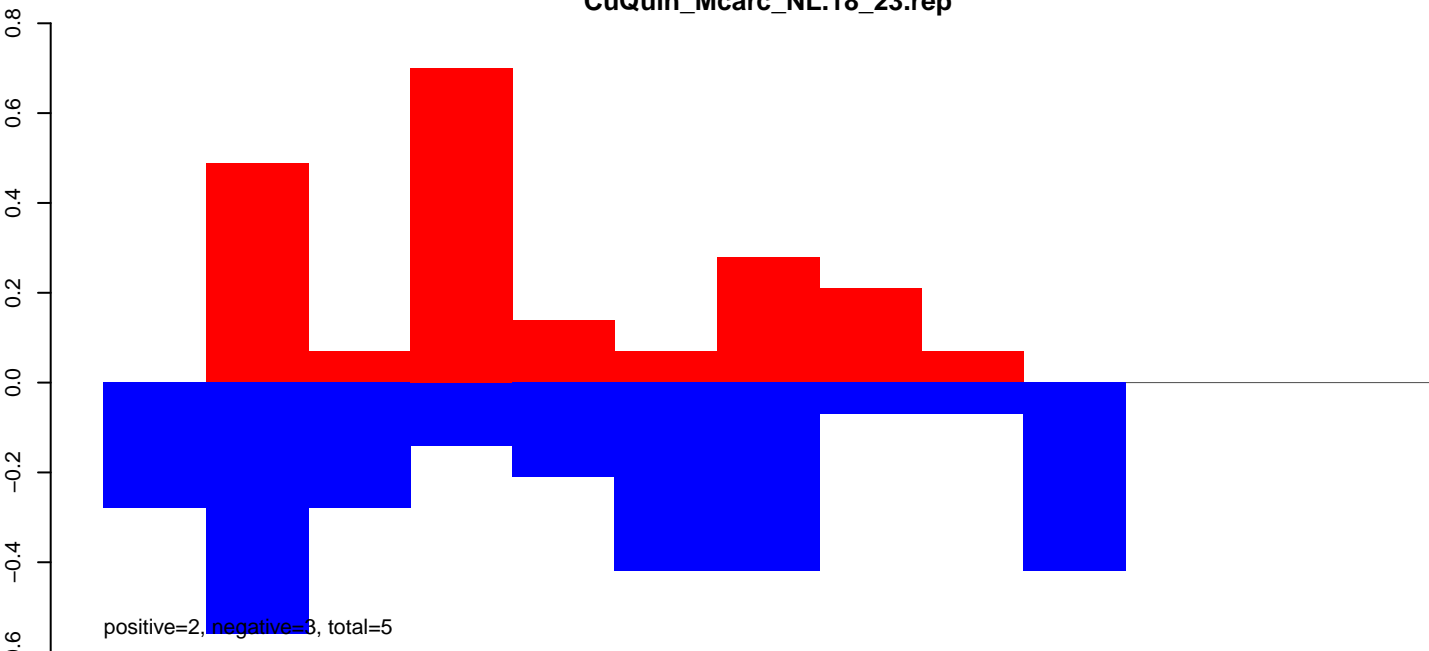
CuQuin_Mcarc_NL.24_35.rep



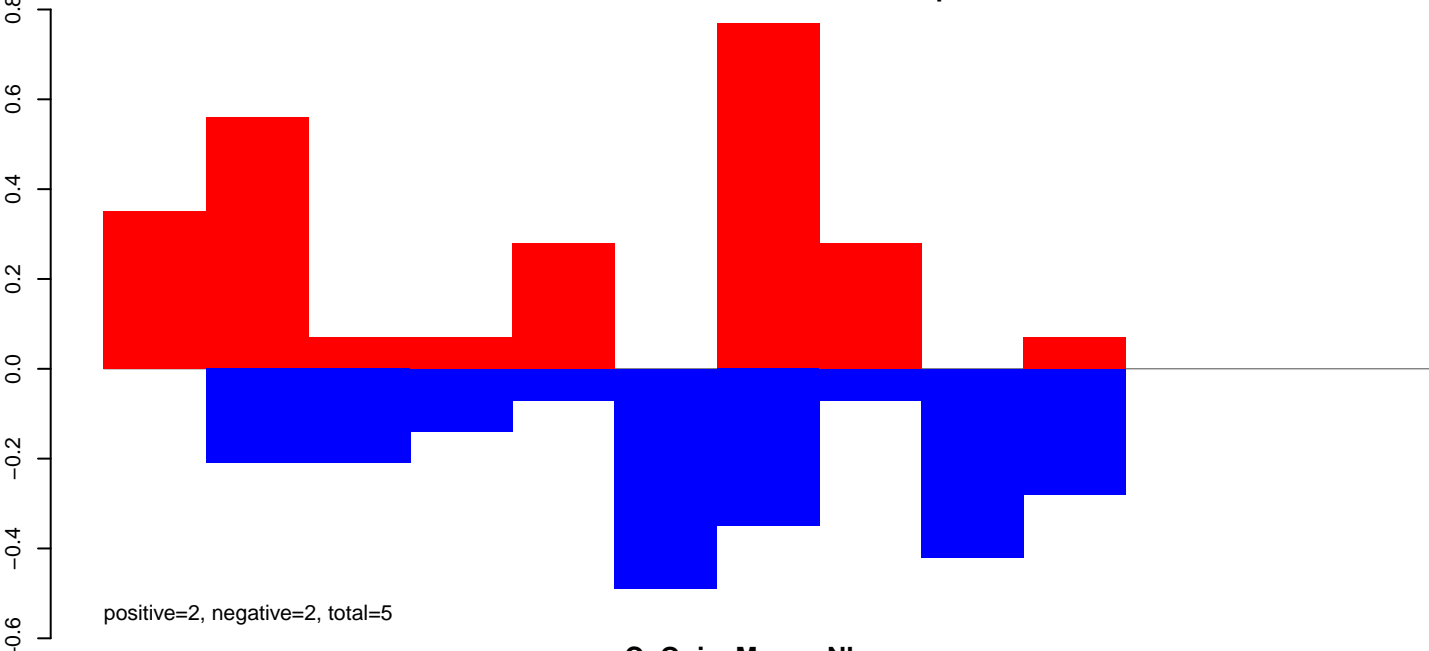
CuQuin_Mcarc_NL.rep



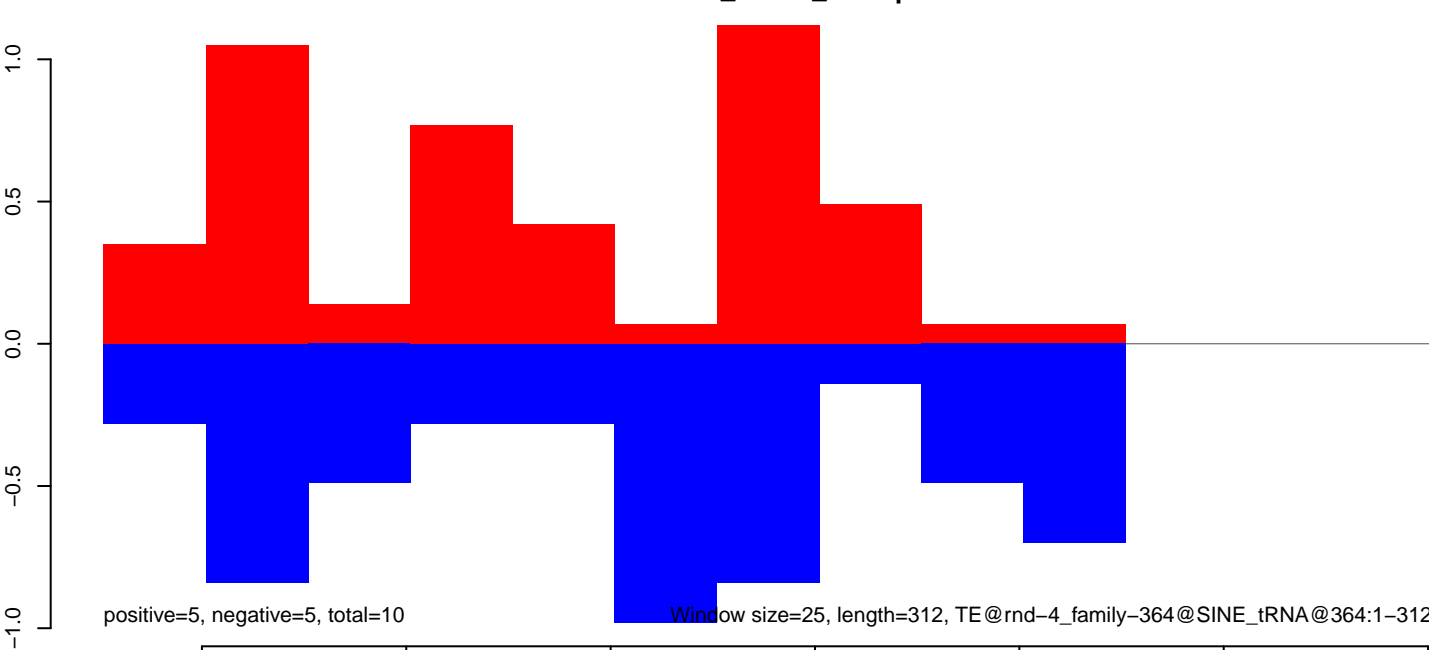
CuQuin_Mcarc_NL.18_23.rep



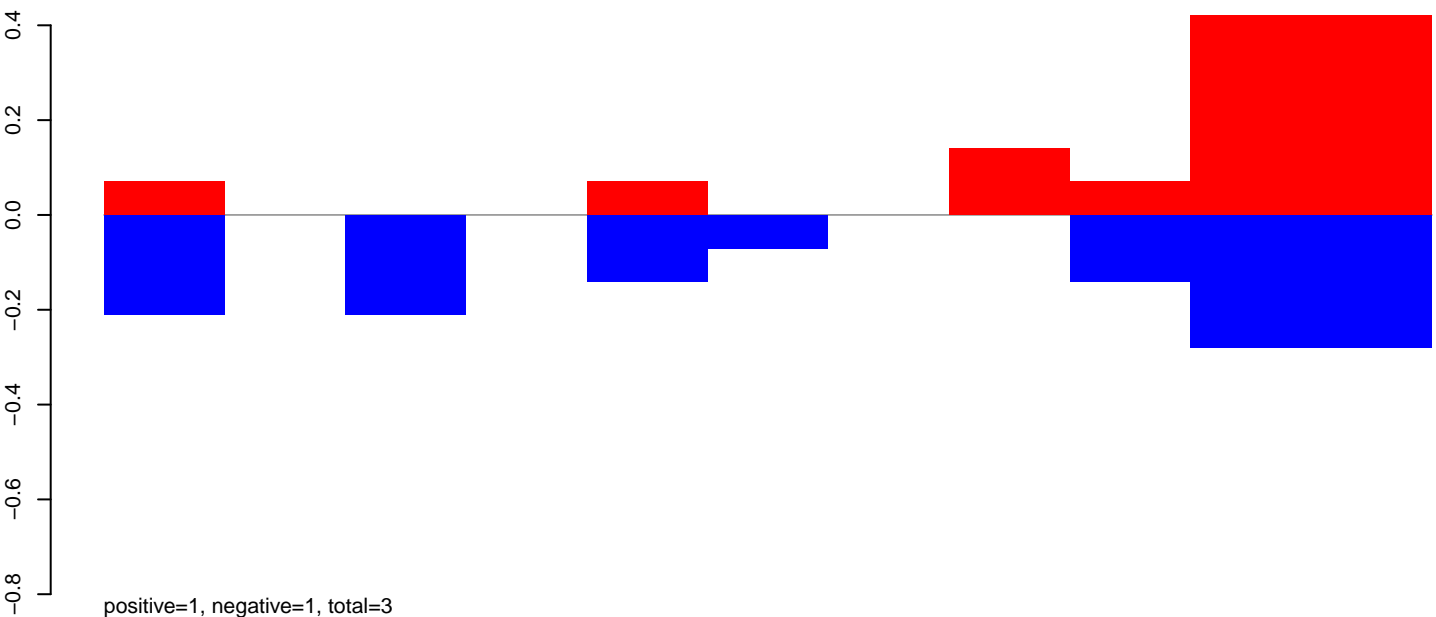
CuQuin_Mcarc_NL.24_35.rep



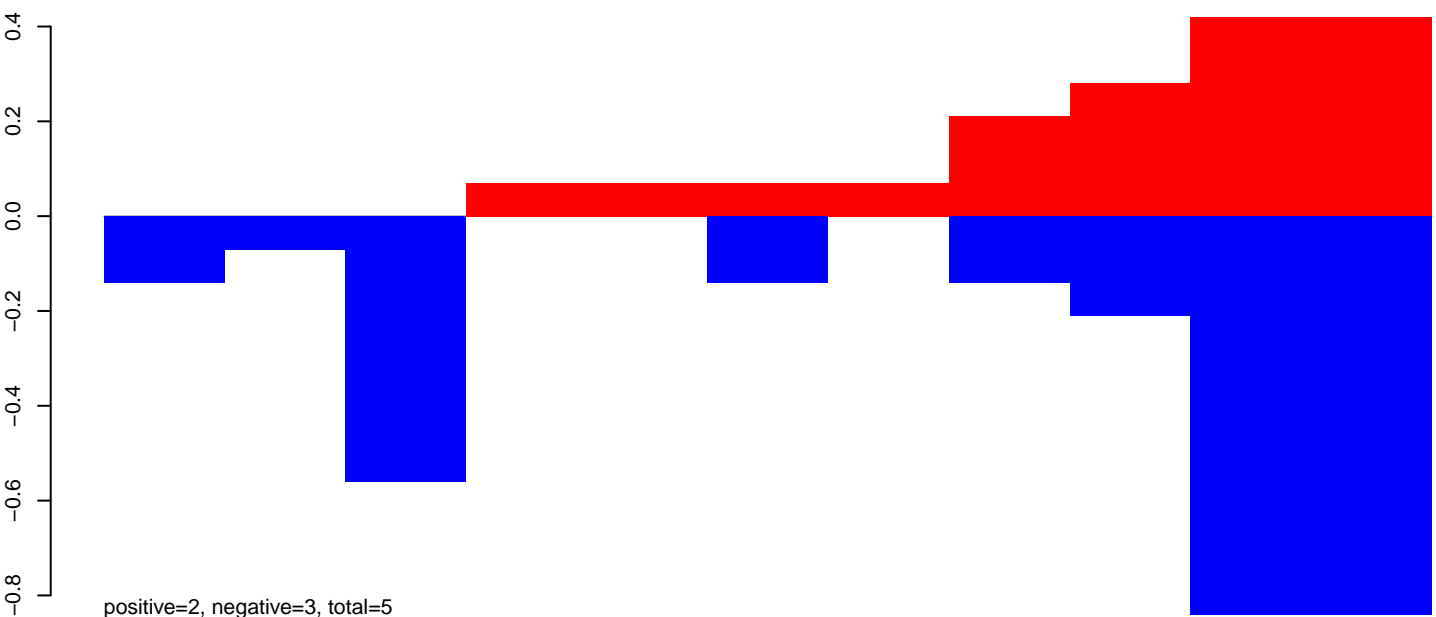
CuQuin_Mcarc_NL.rep



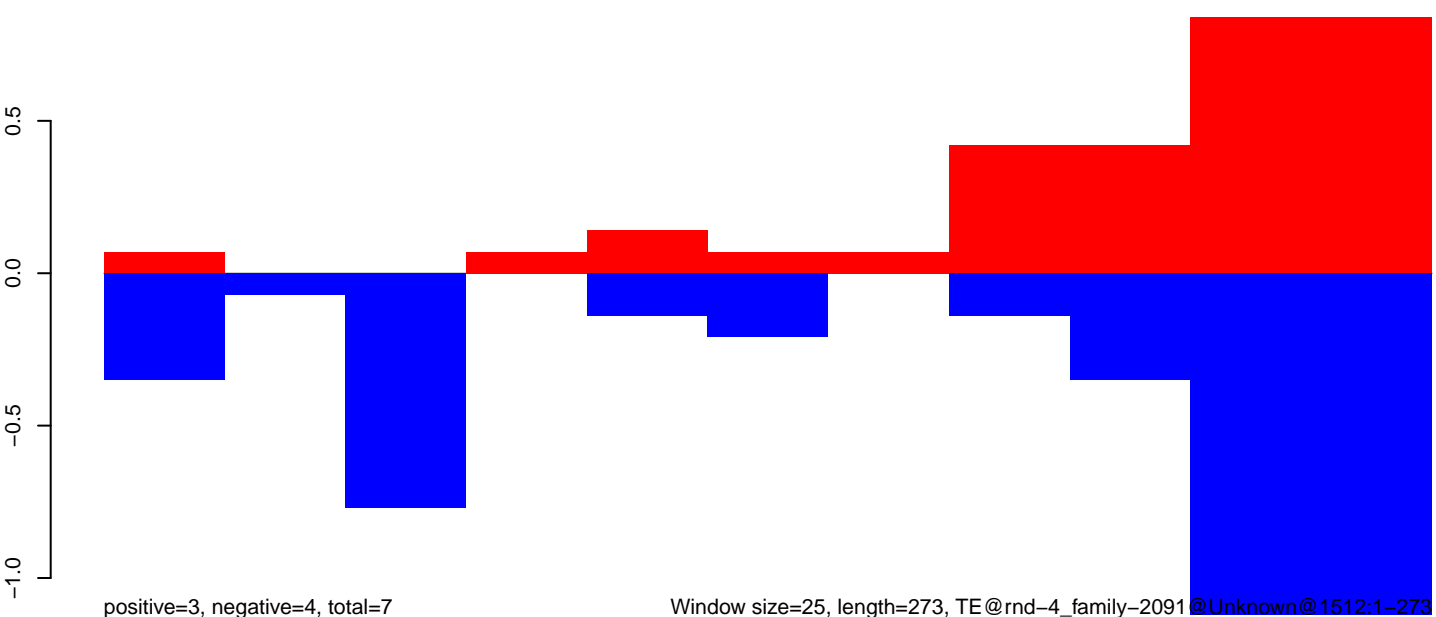
CuQuin_Mcarc_NL.18_23.rep



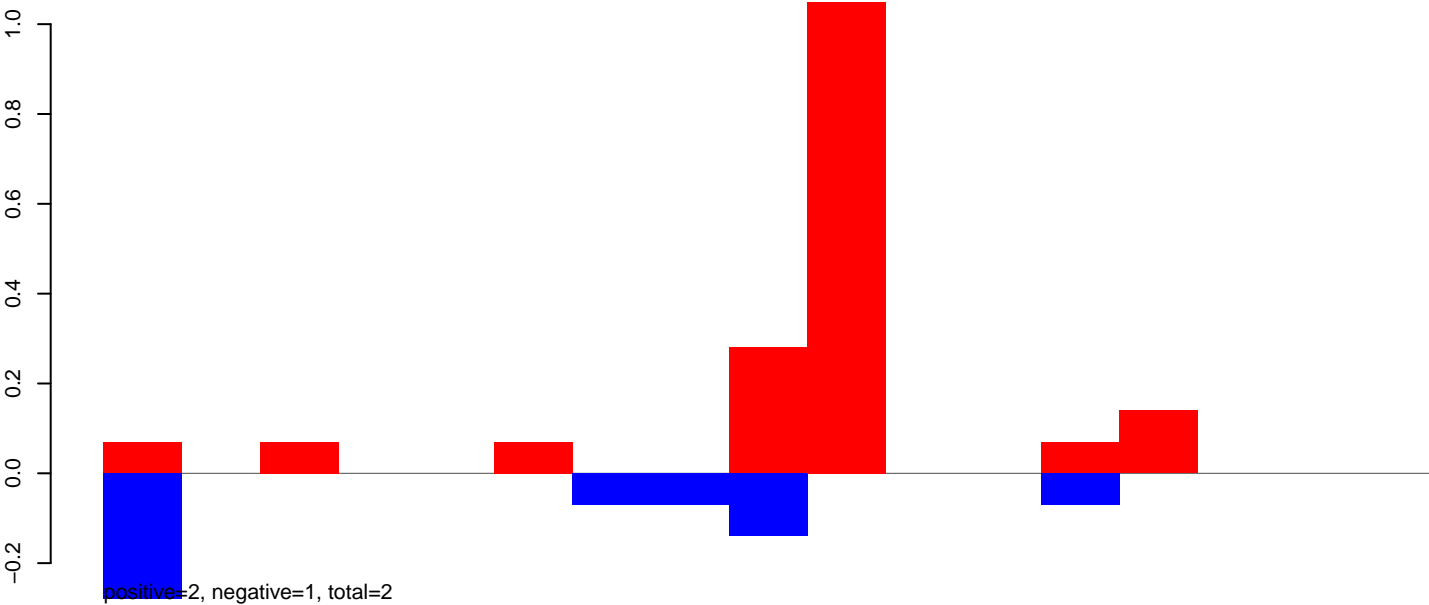
CuQuin_Mcarc_NL.24_35.rep



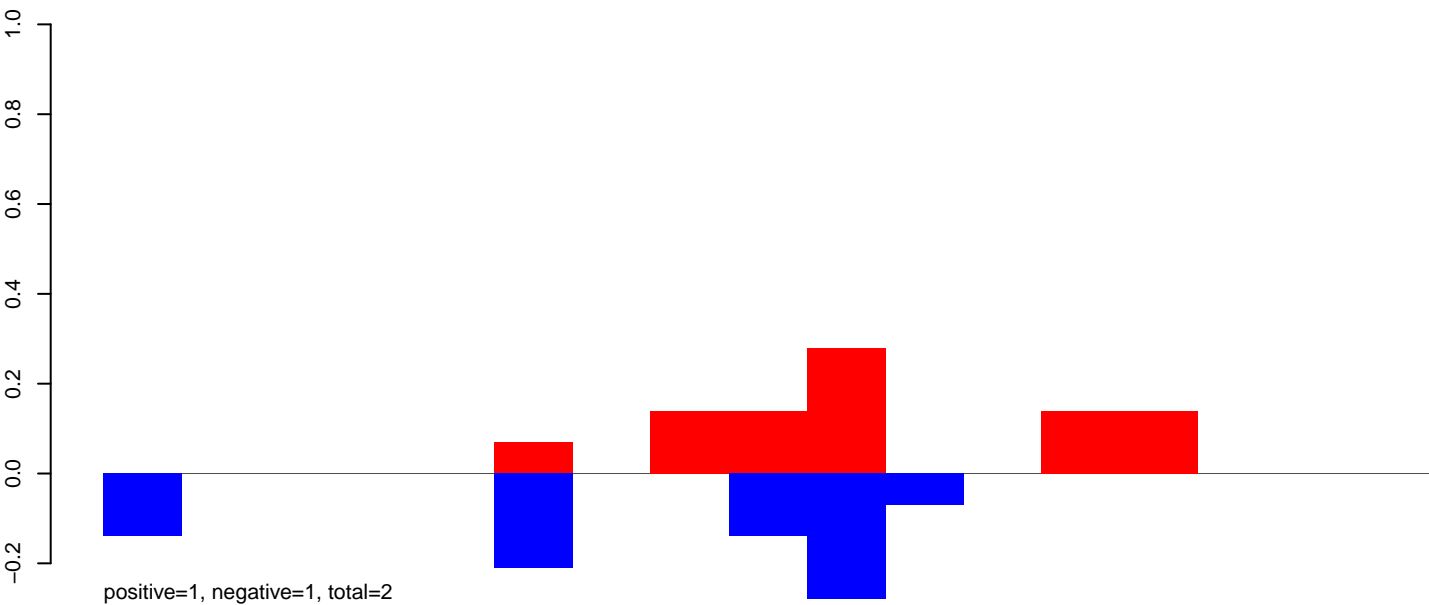
CuQuin_Mcarc_NL.rep



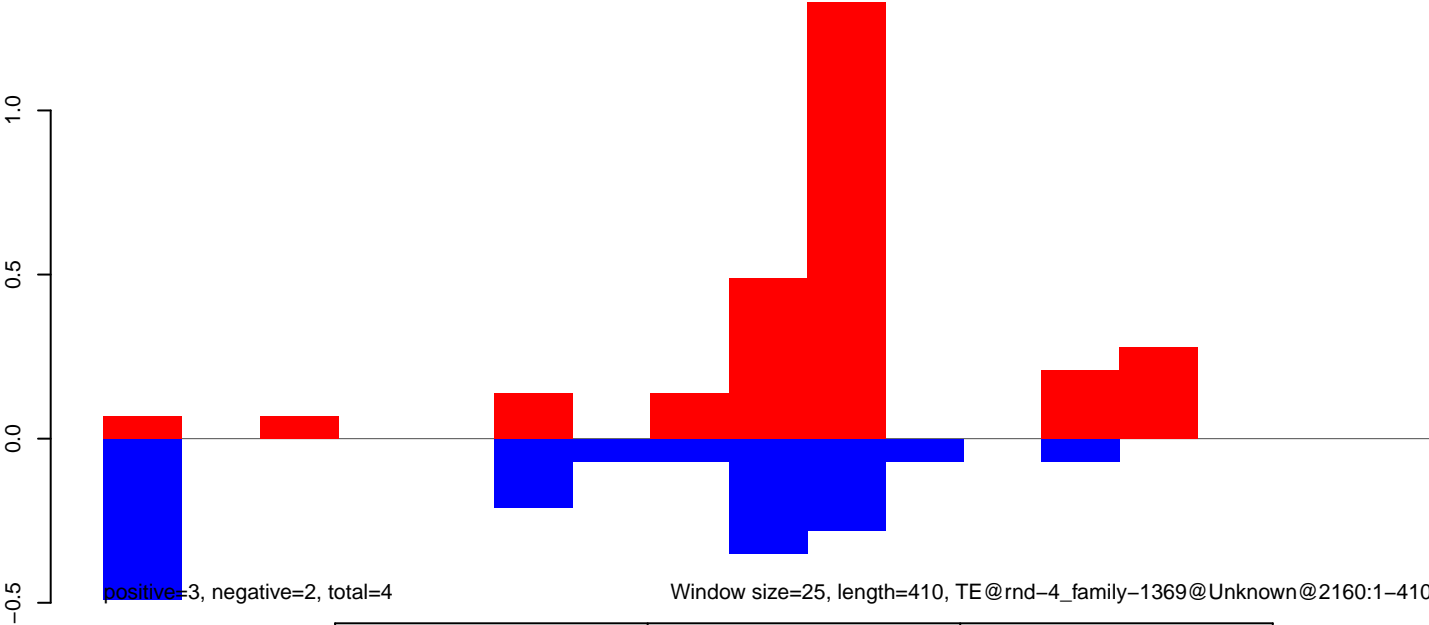
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

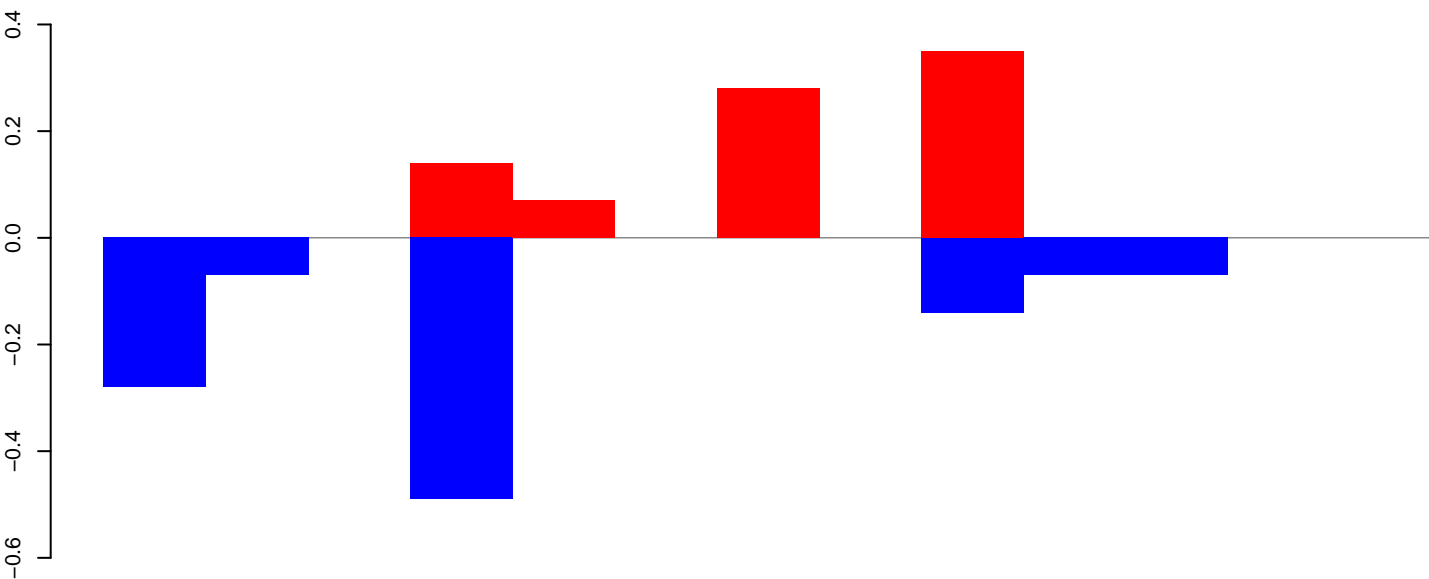


CuQuin_Mcarc_NL.rep



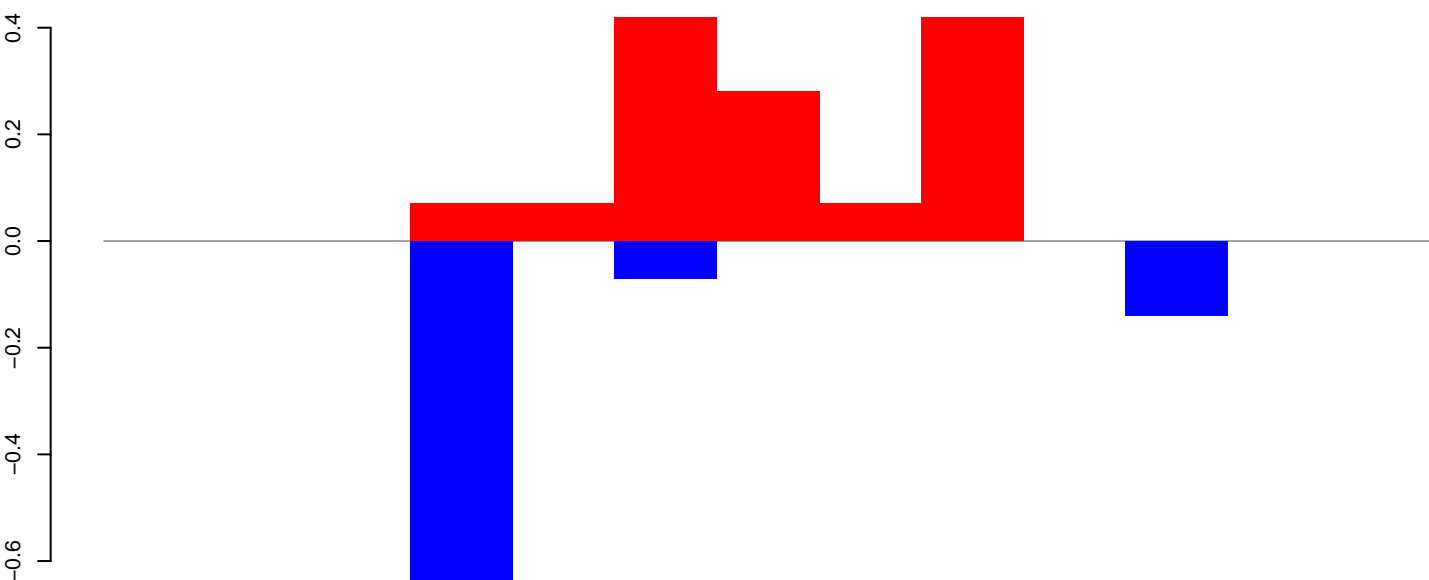
Window size=25, length=410, TE@rnd-4_family-1369@Unknown@2160:1-410

CuQuin_Mcarc_NL.18_23.rep



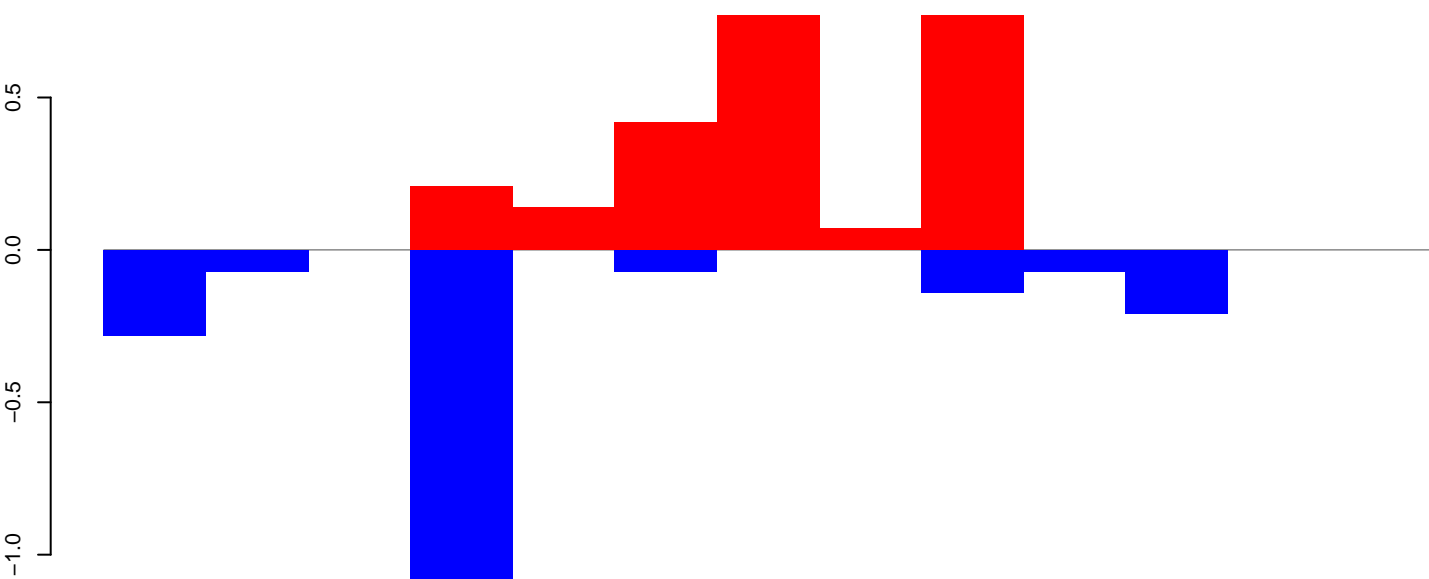
positive=1, negative=1, total=2

CuQuin_Mcarc_NL.24_35.rep



positive=1, negative=1, total=2

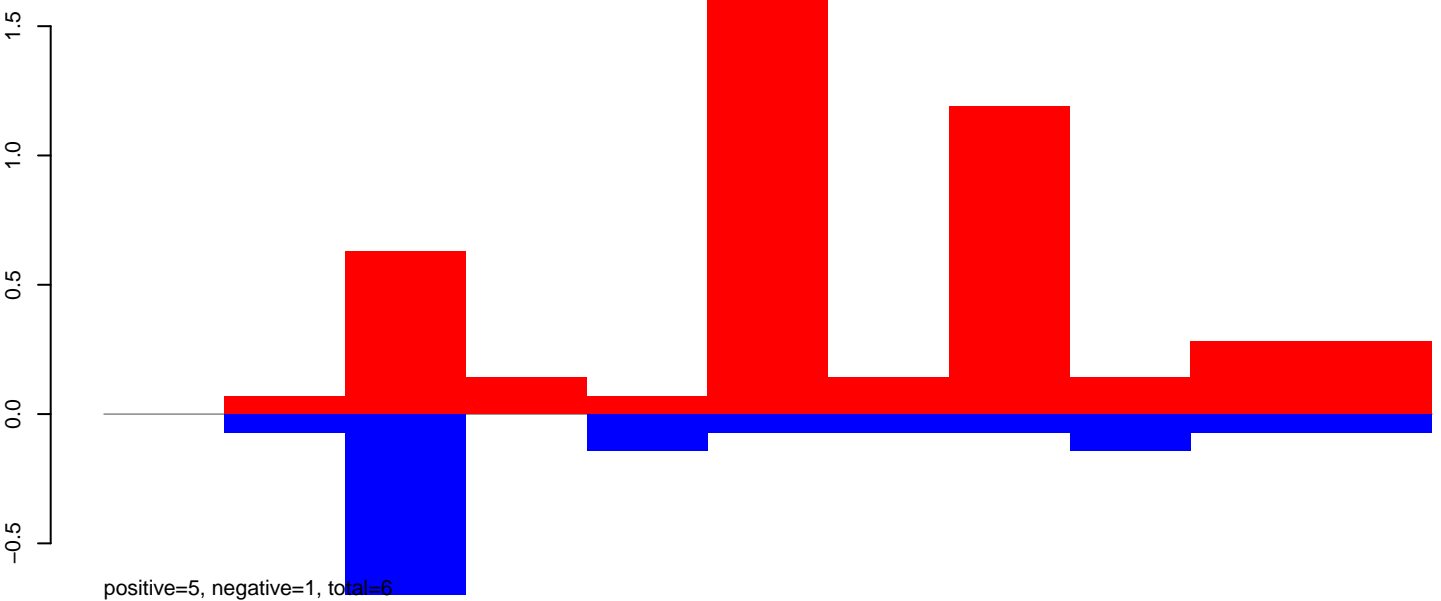
CuQuin_Mcarc_NL.rep



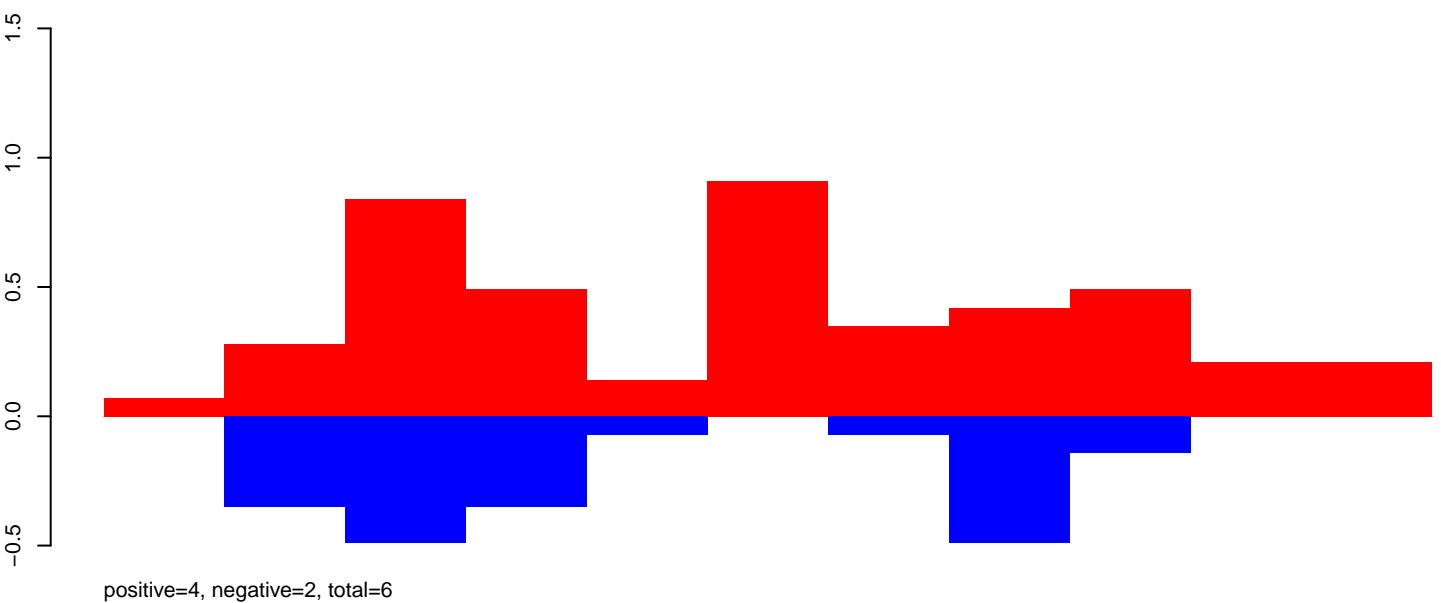
positive=2, negative=2, total=4

Window size=25, length=304, TE@rnd-4_family-1272@Unknown@2155:1-304

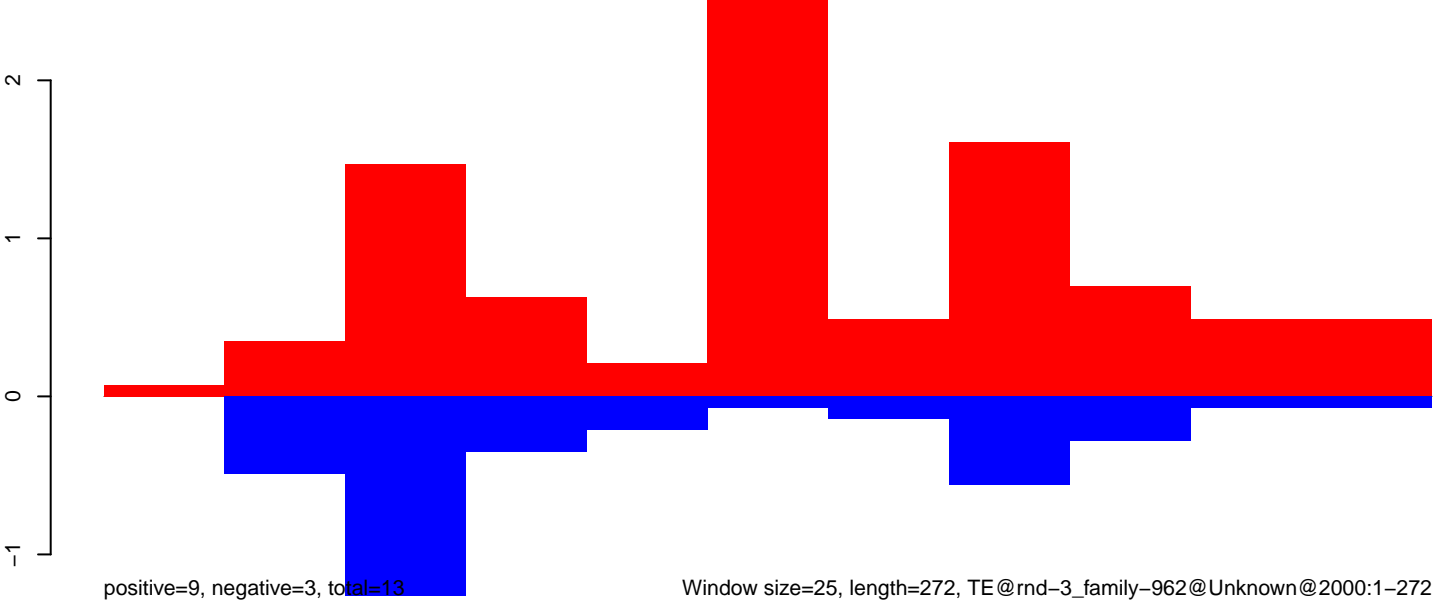
CuQuin_Mcarc_NL.18_23.rep



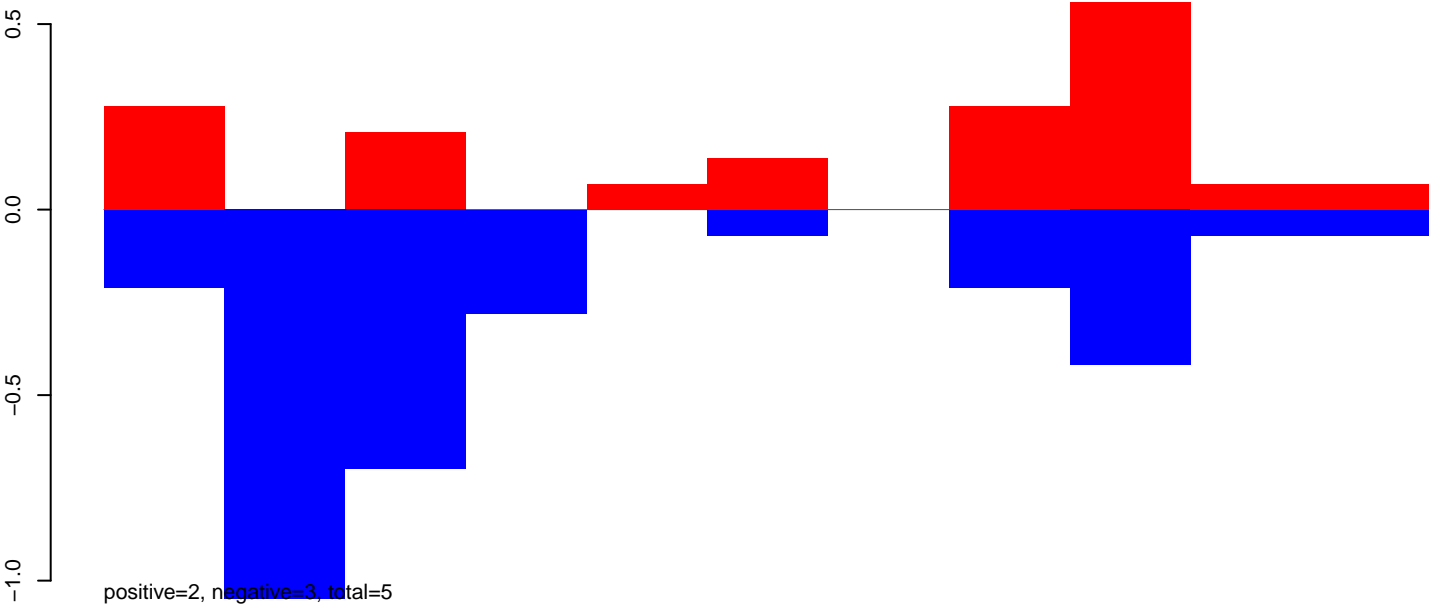
CuQuin_Mcarc_NL.24_35.rep



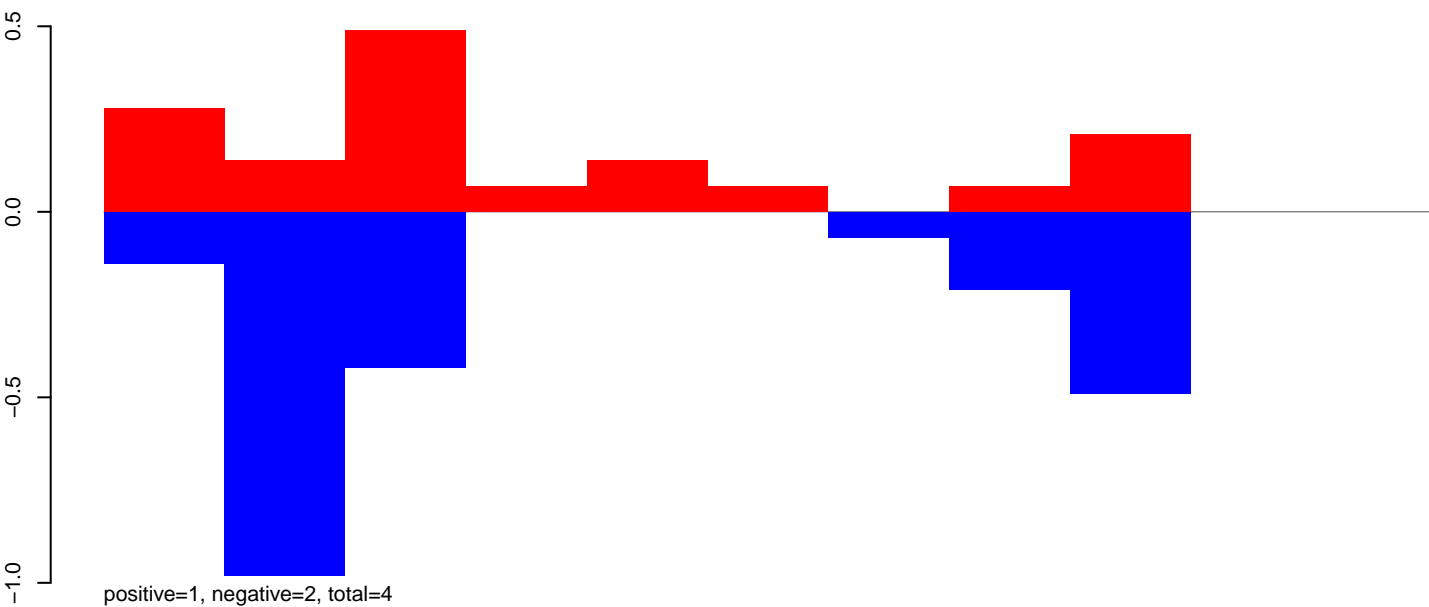
CuQuin_Mcarc_NL.rep



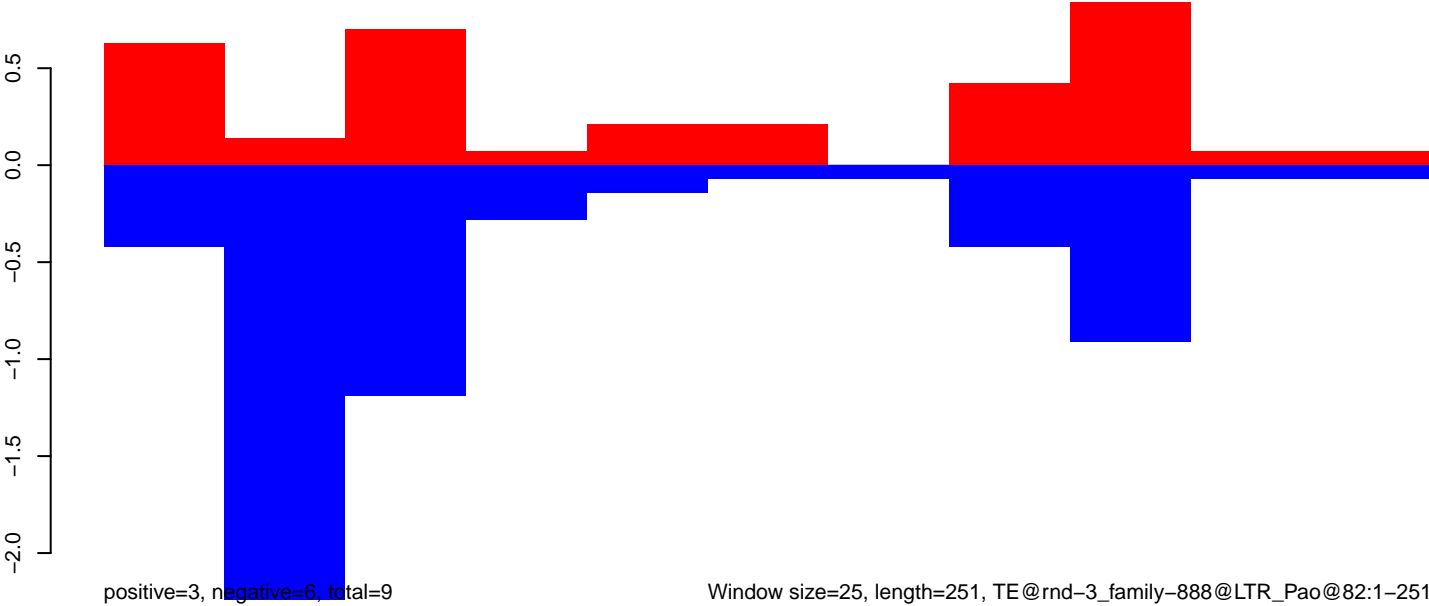
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=251, TE@rnd-3_family-888@LTR_Pao@82:1-251

50

100

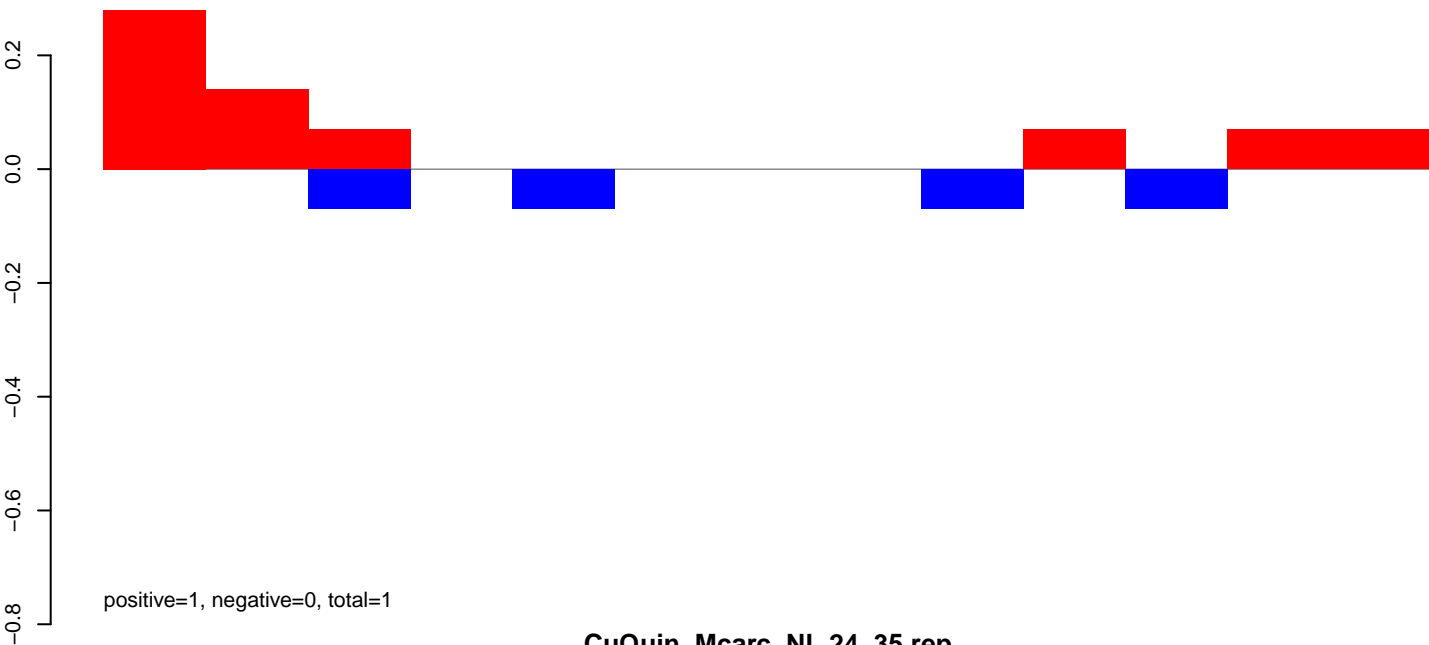
150

200

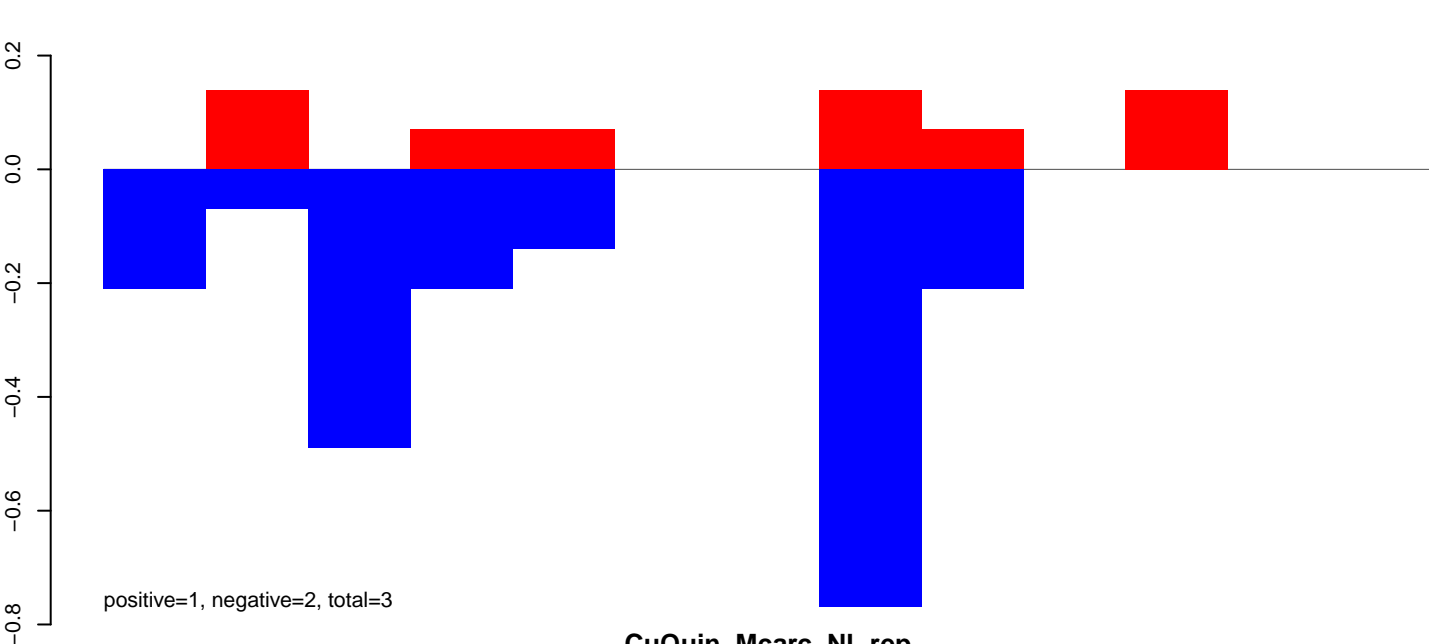
250

300

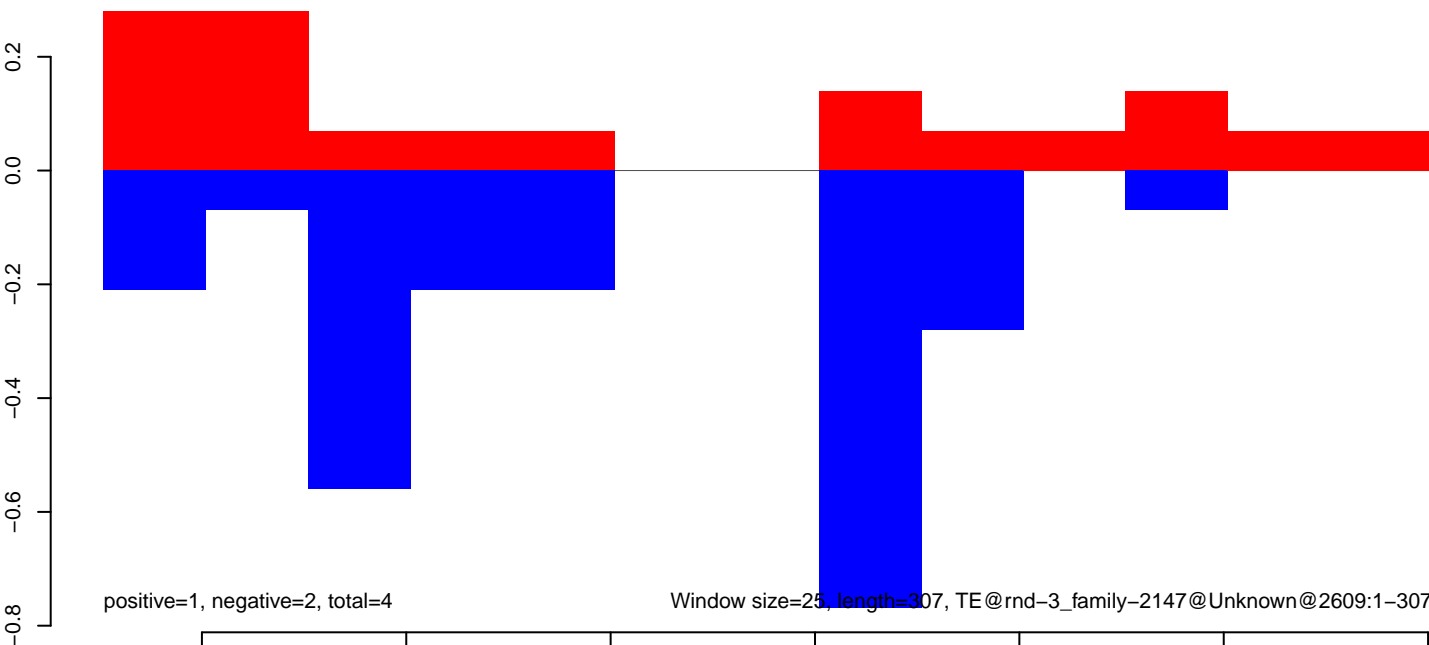
CuQuin_Mcarc_NL.18_23.rep



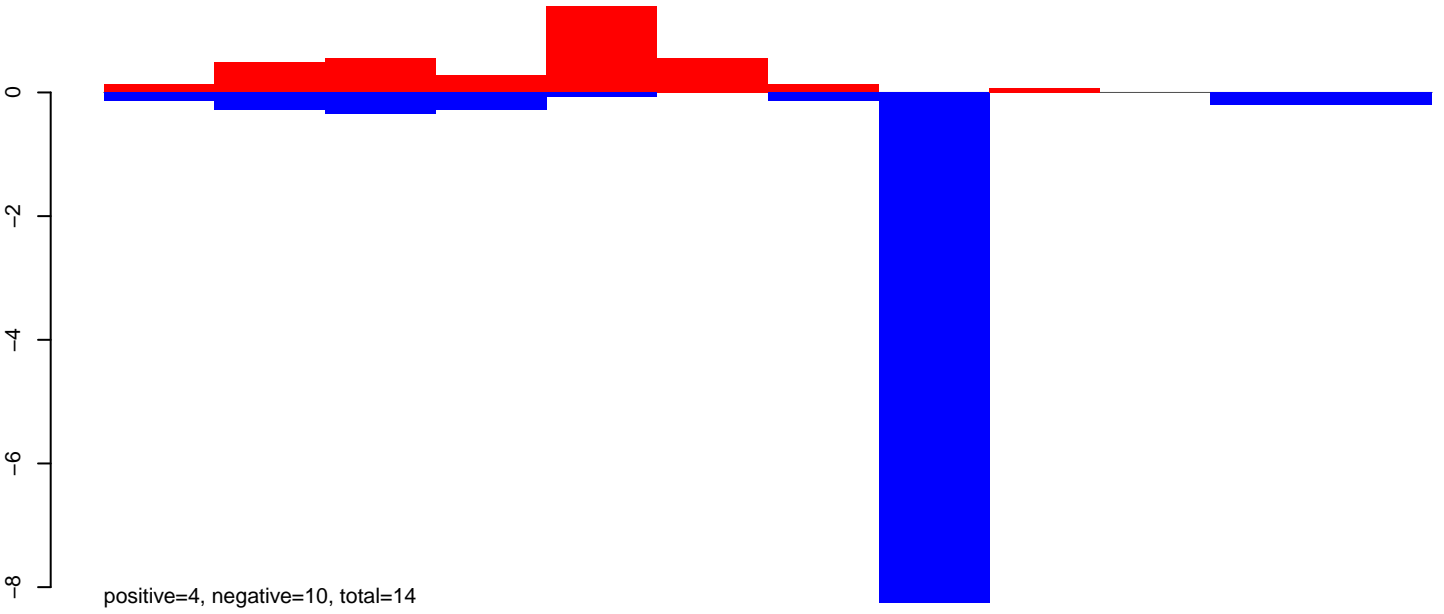
CuQuin_Mcarc_NL.24_35.rep



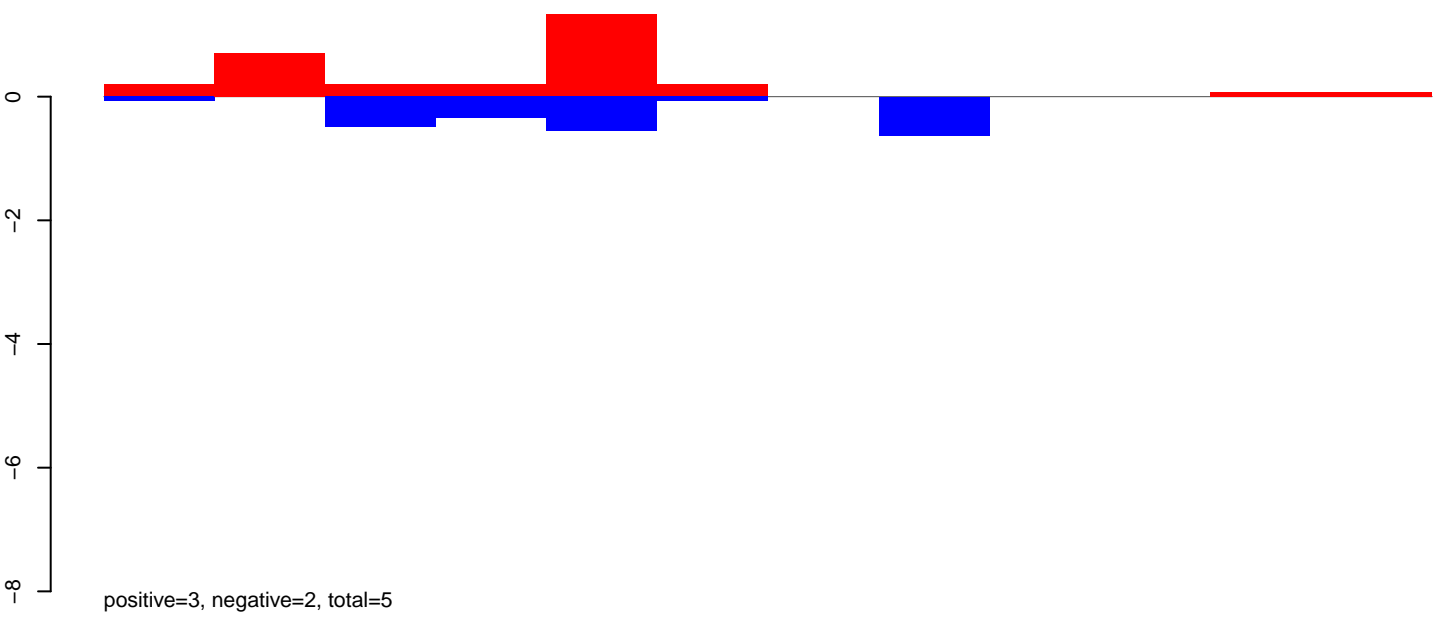
CuQuin_Mcarc_NL.rep



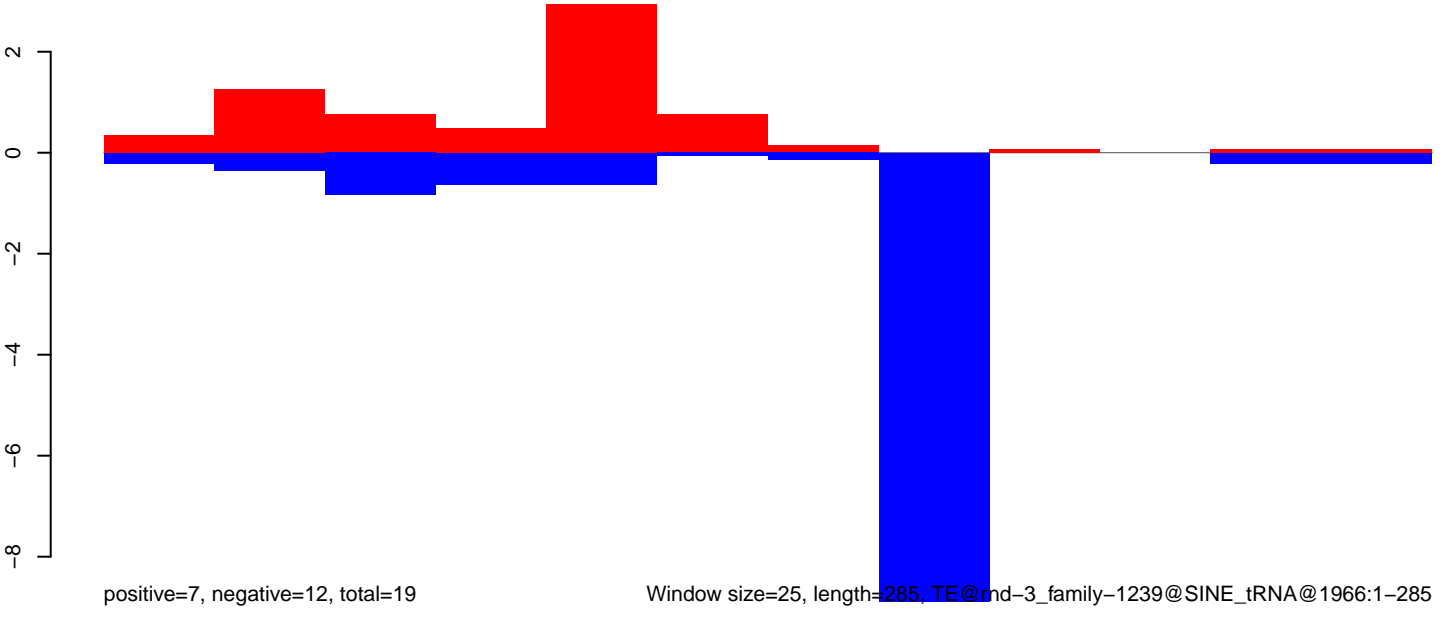
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

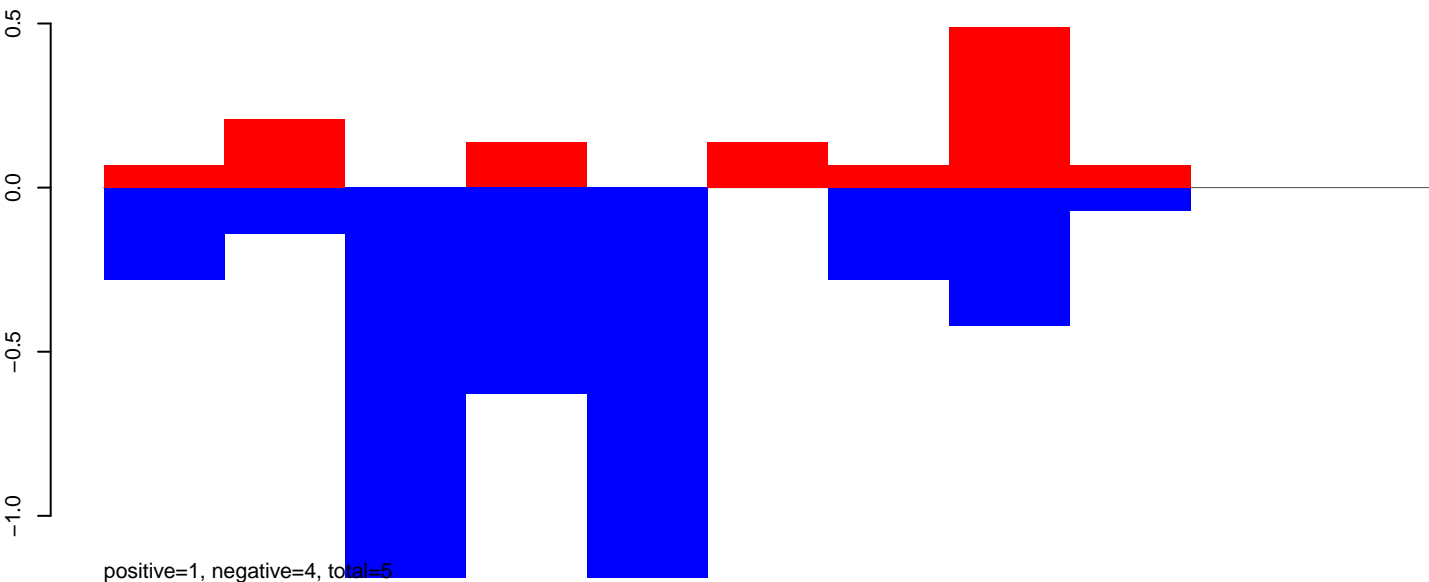


CuQuin_Mcarc_NL.rep

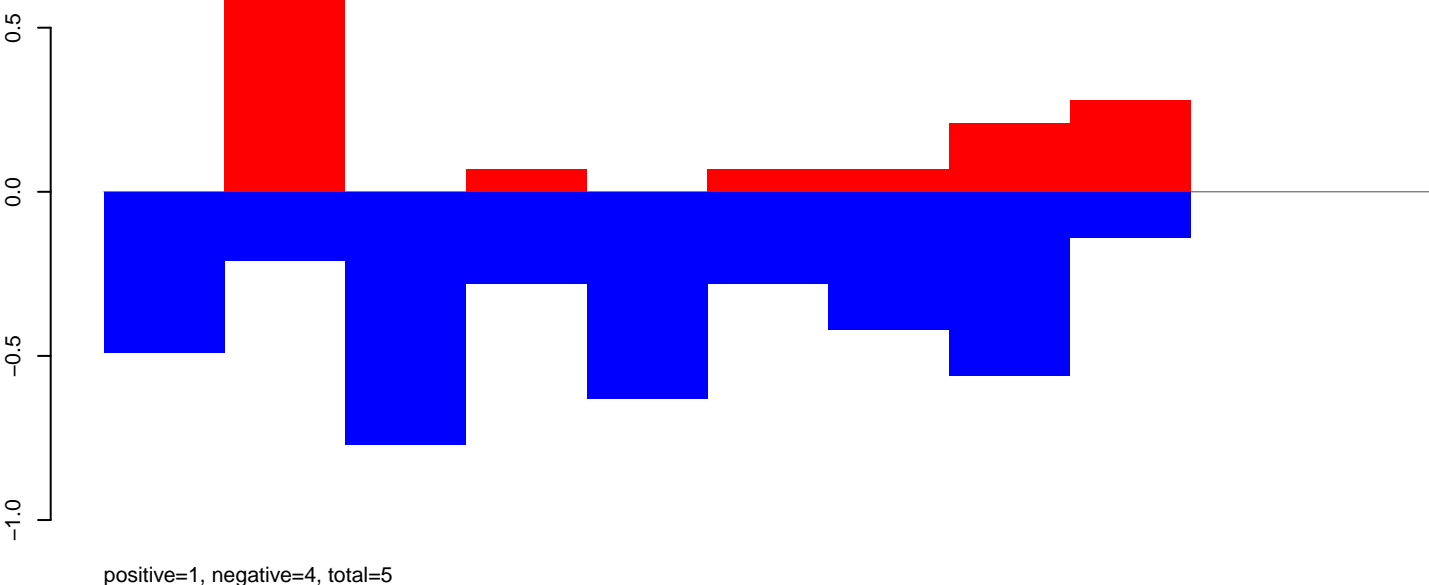


Window size=25, length=285, TE@rmd-3_family-1239@SINE_tRNA@1966:1-285

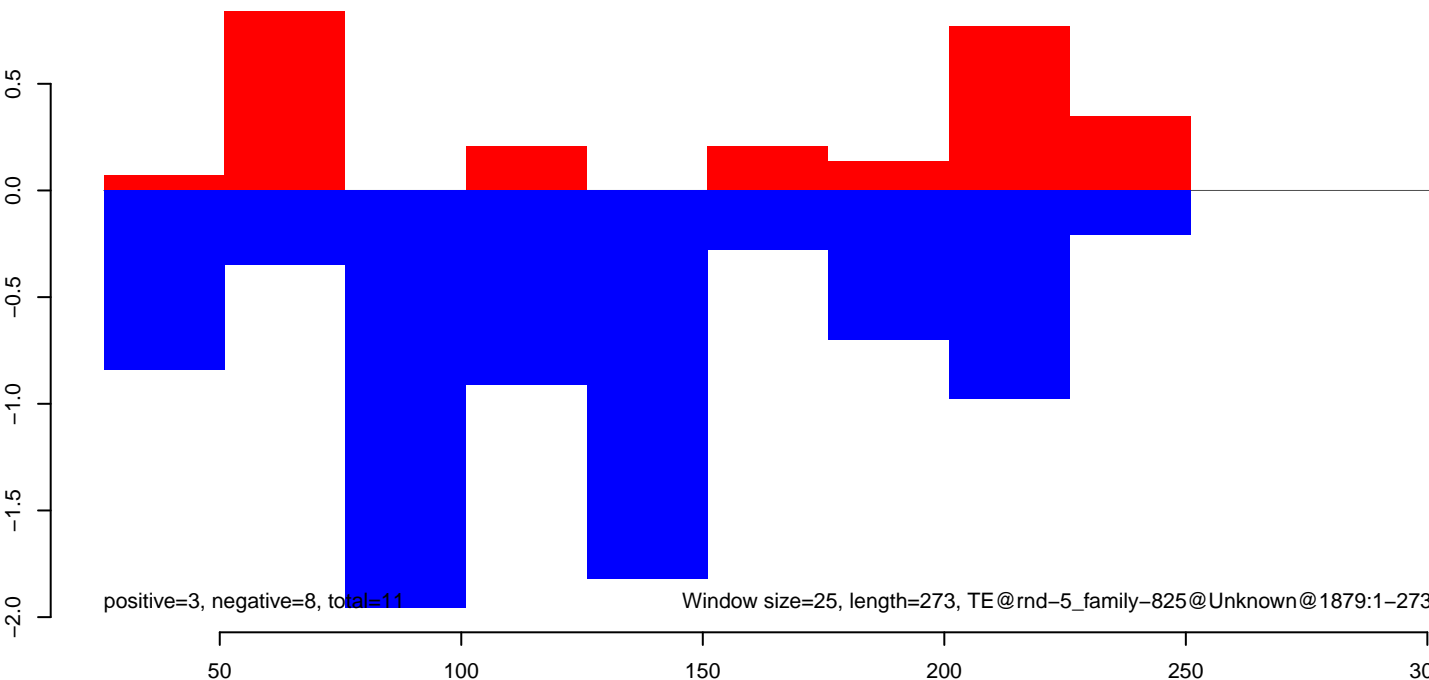
CuQuin_Mcarc_NL.18_23.rep



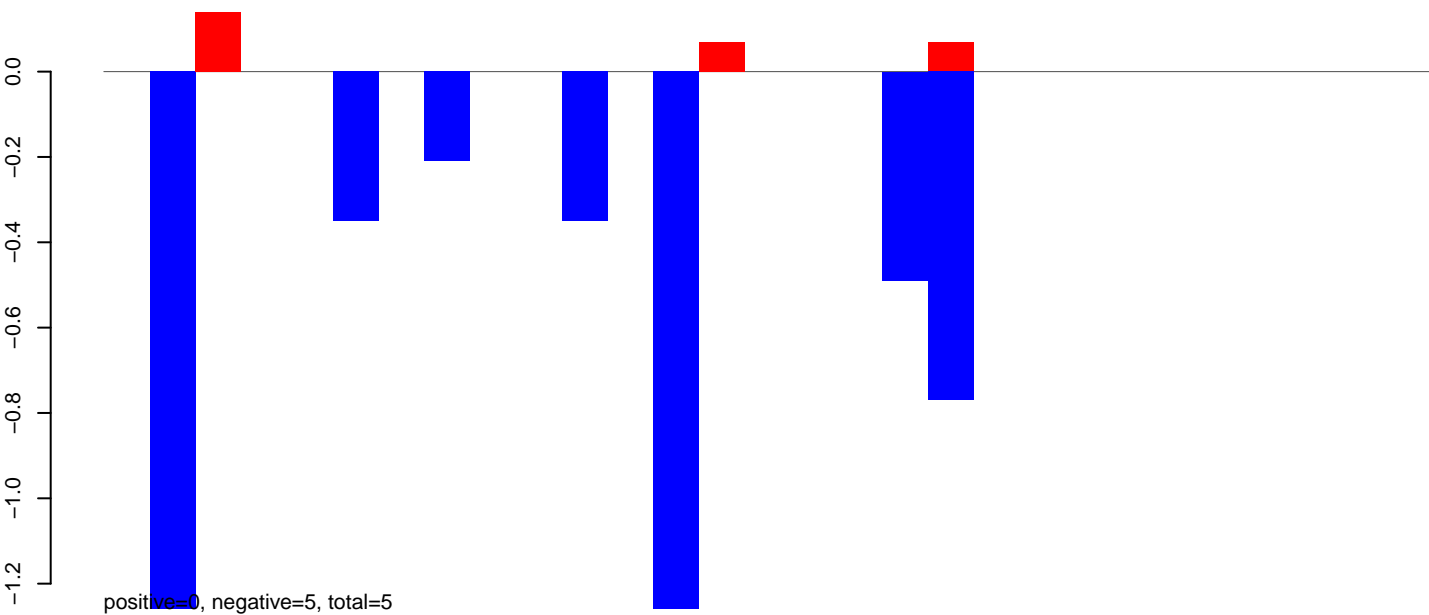
CuQuin_Mcarc_NL.24_35.rep



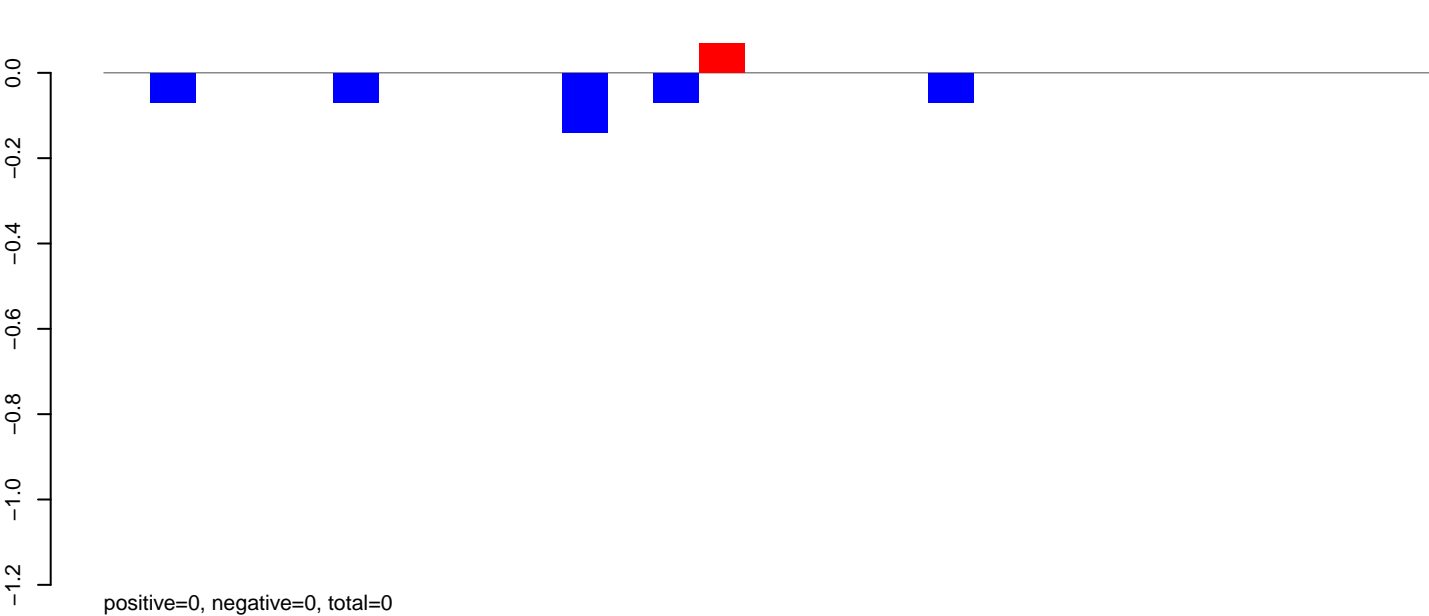
CuQuin_Mcarc_NL.rep



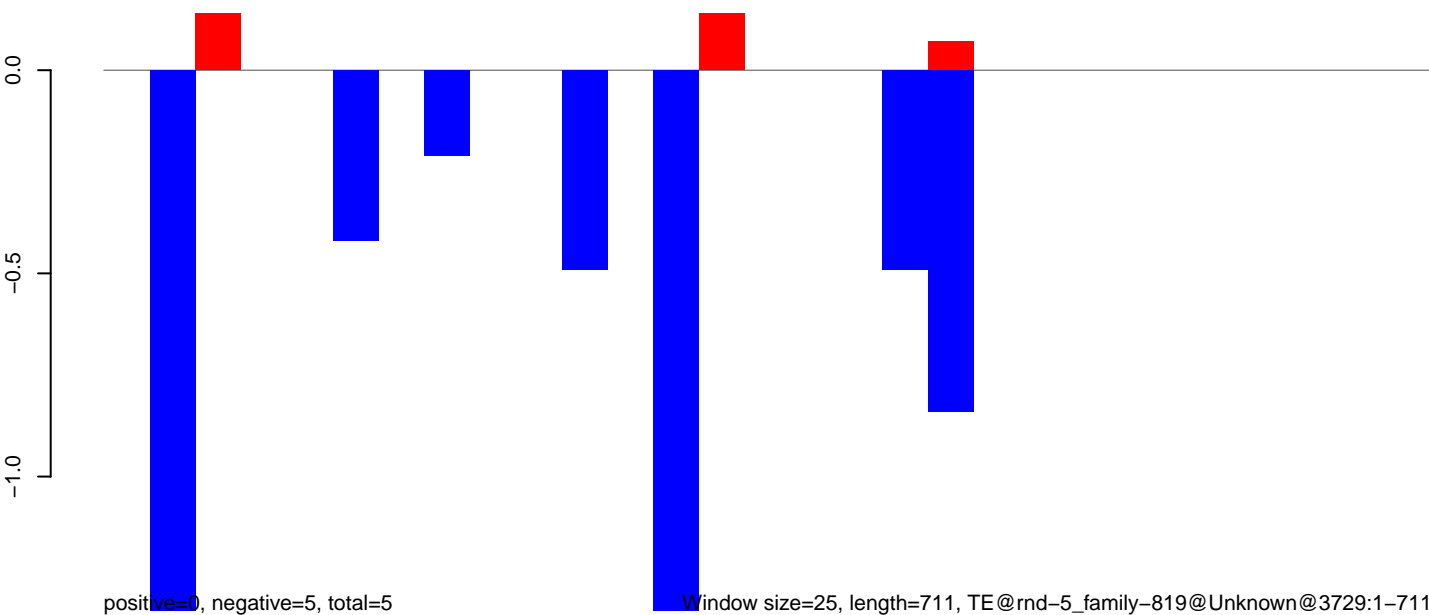
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

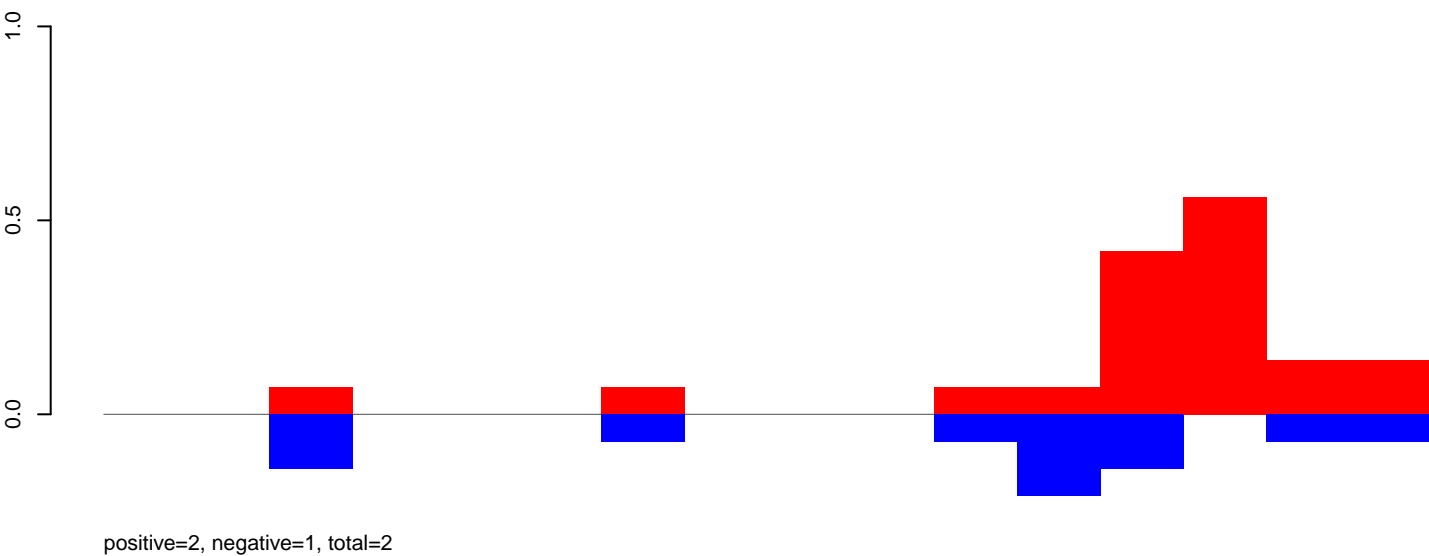


CuQuin_Mcarc_NL.rep

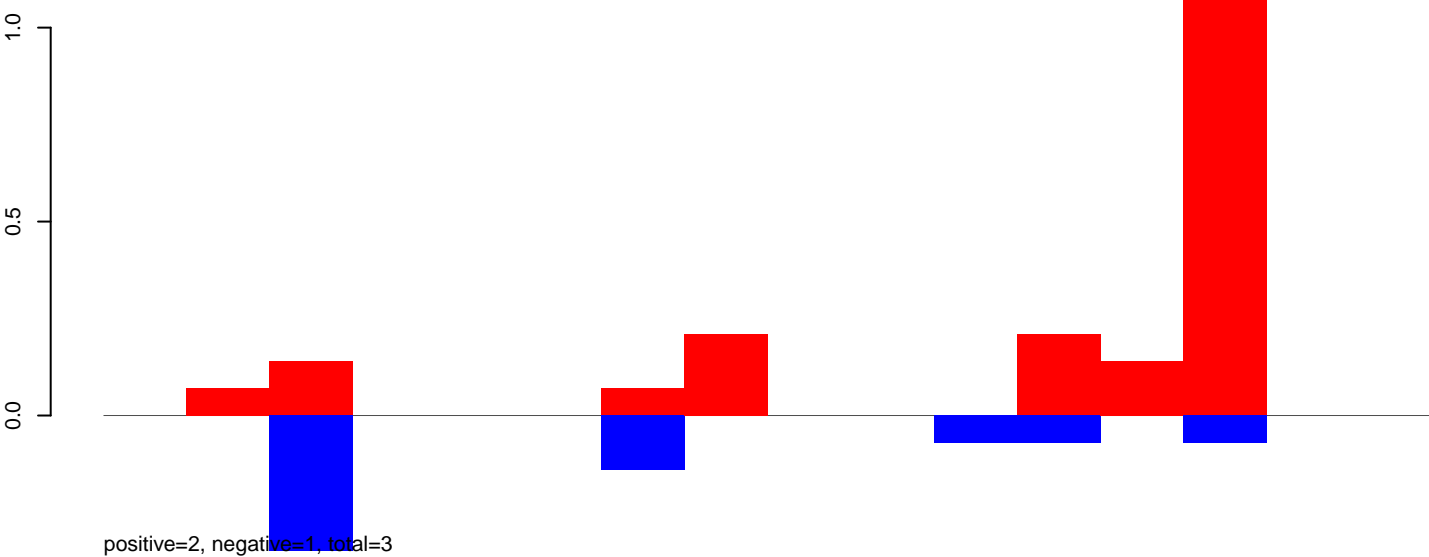


Window size=25, length=711, TE@rnd-5_family-819@Unknown@3729:1-711

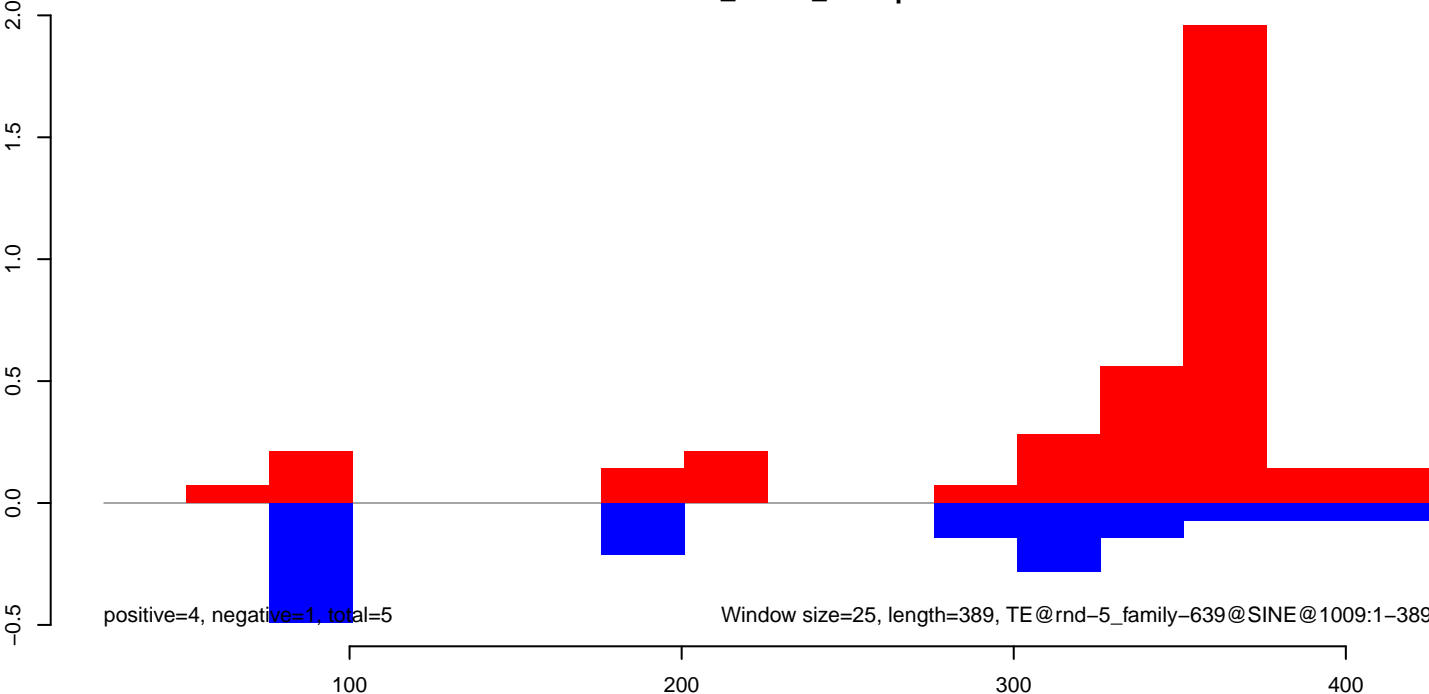
CuQuin_Mcarc_NL.18_23.rep



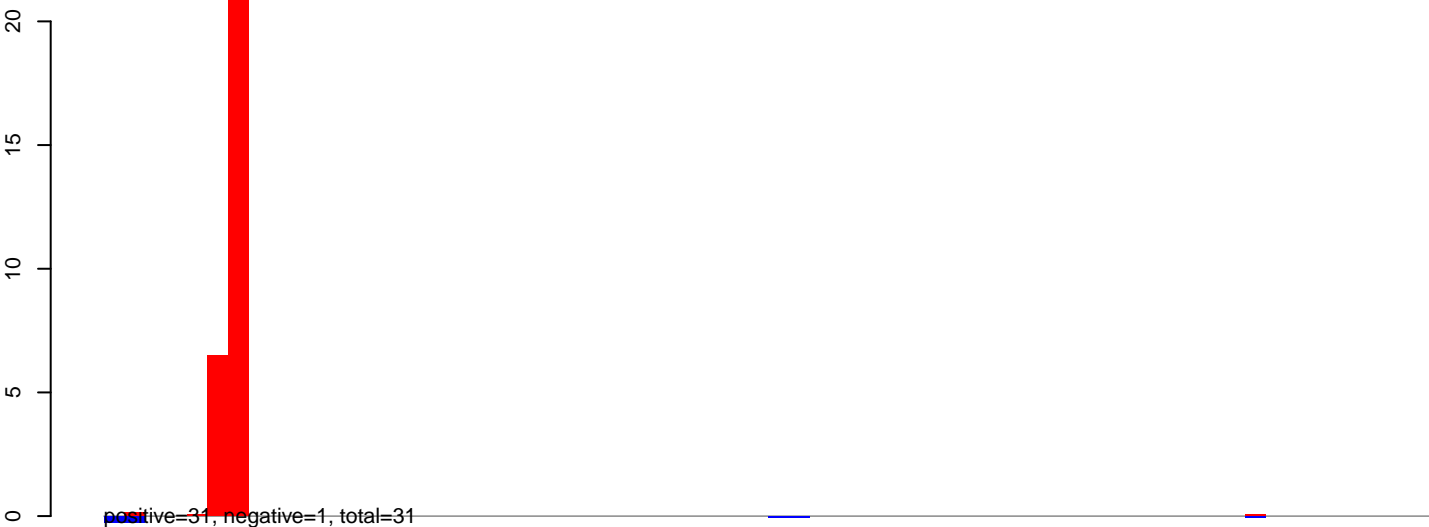
CuQuin_Mcarc_NL.24_35.rep



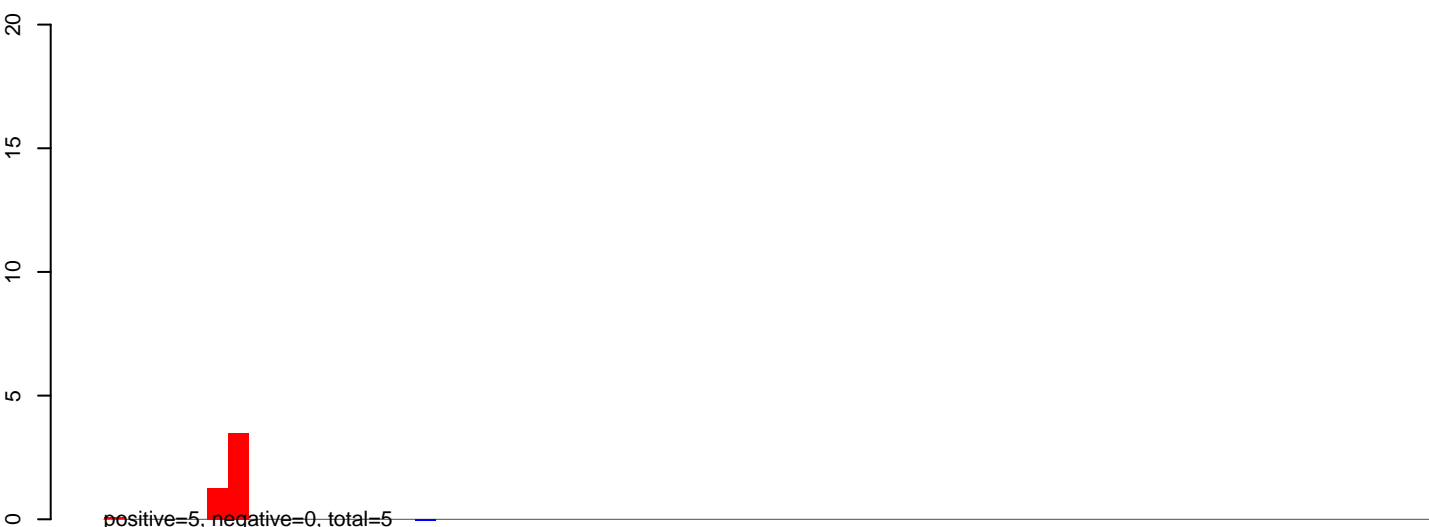
CuQuin_Mcarc_NL.rep



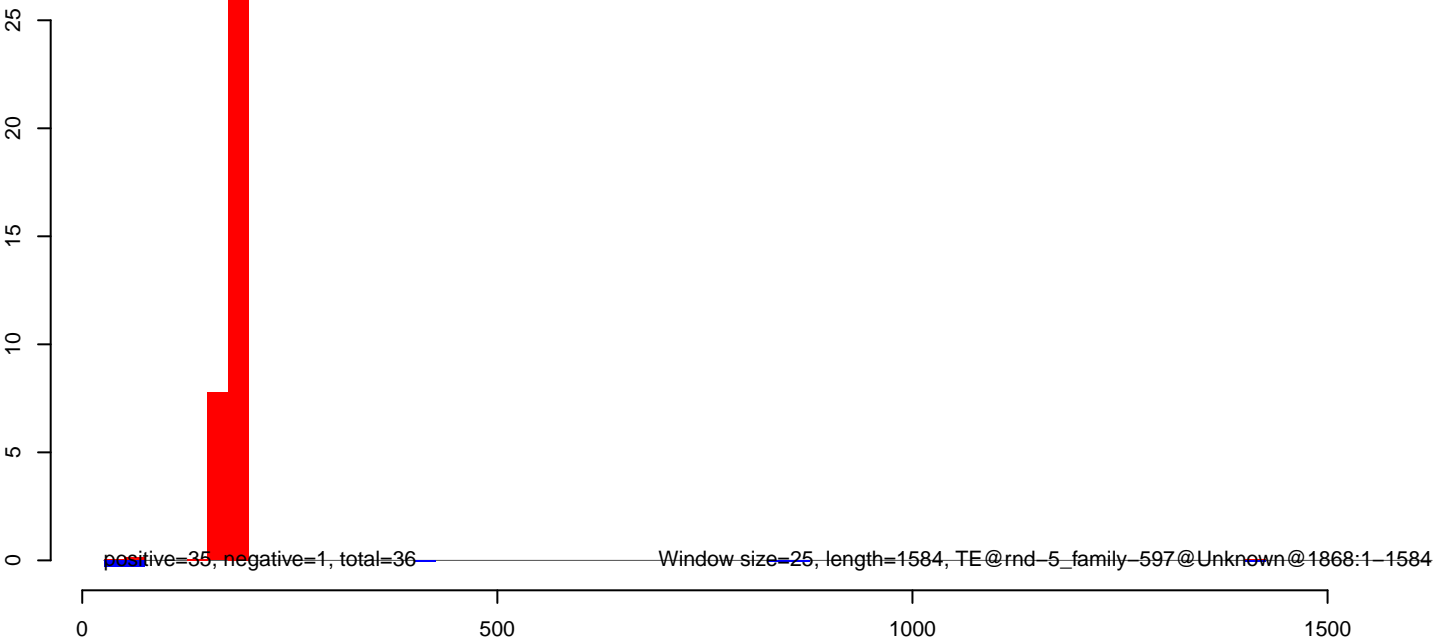
CuQuin_Mcarc_NL.18_23.rep



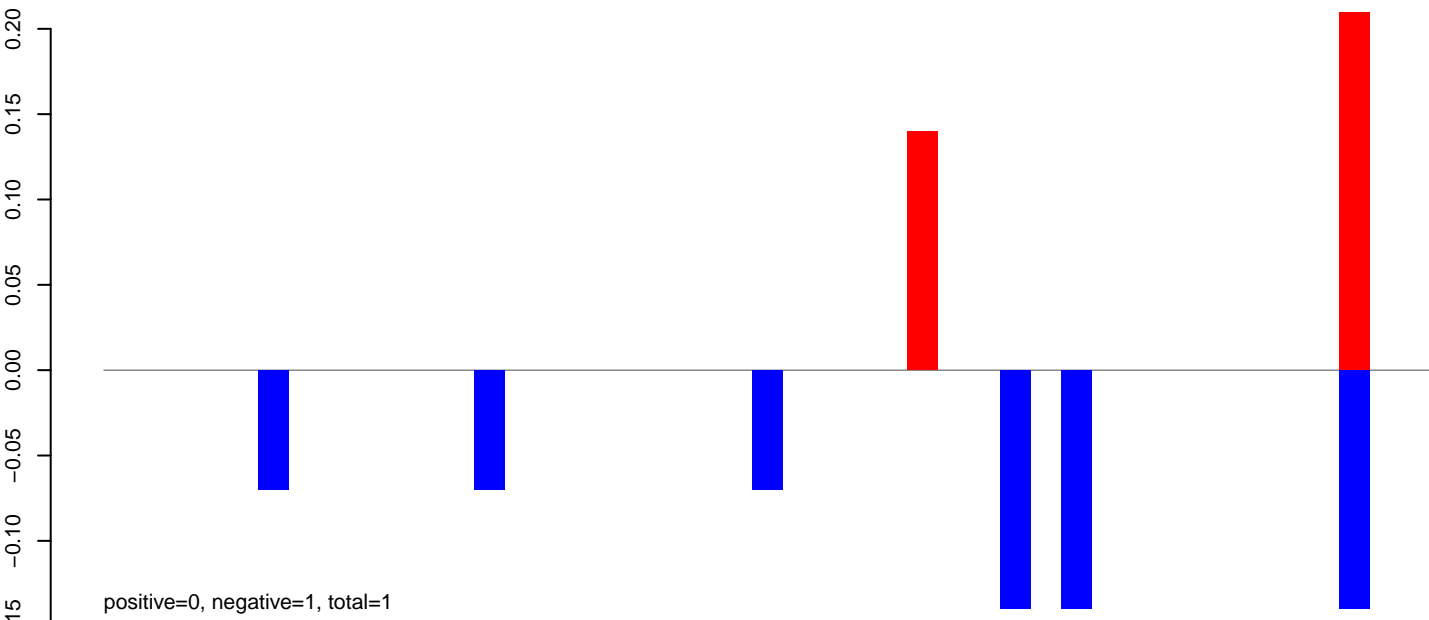
CuQuin_Mcarc_NL.24_35.rep



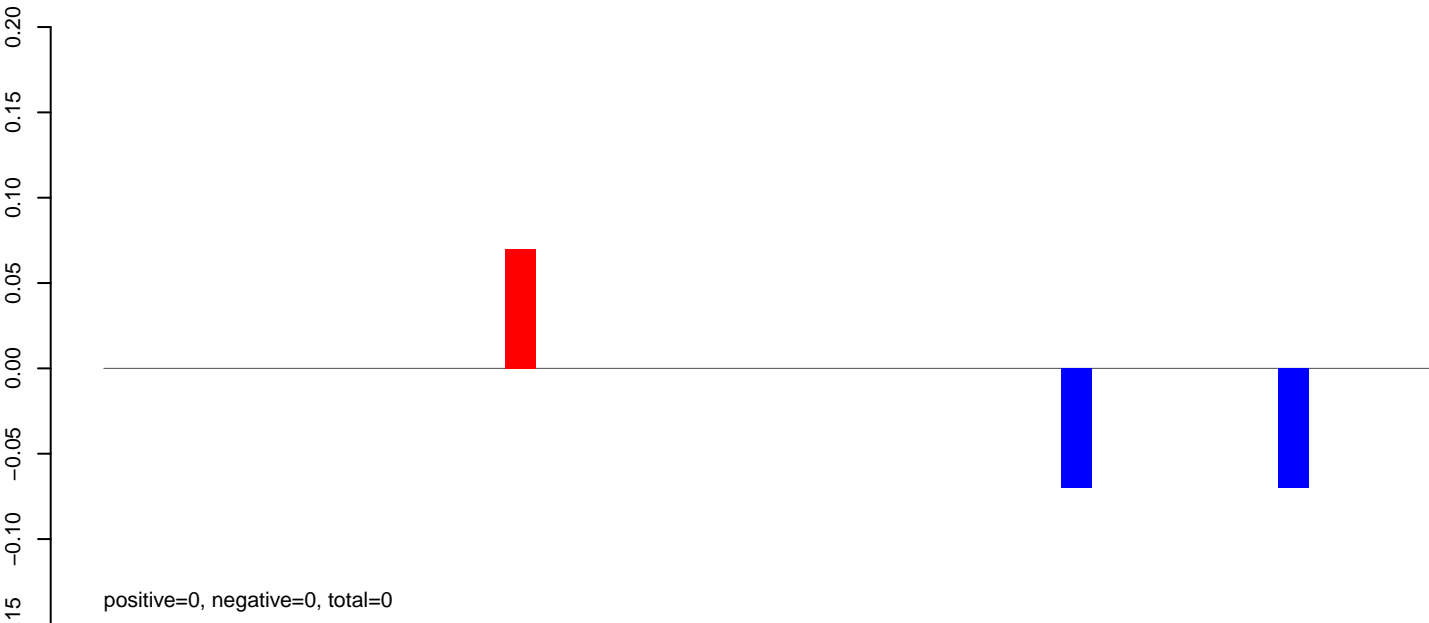
CuQuin_Mcarc_NL.rep



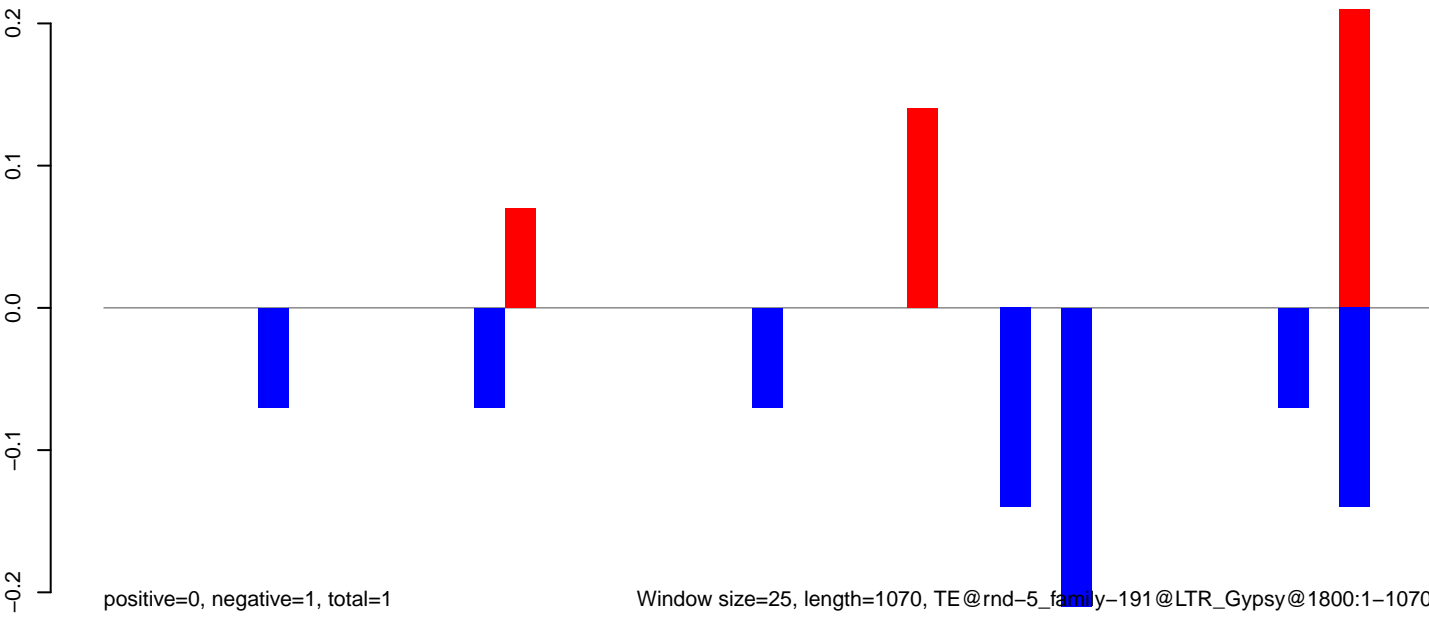
CuQuin_Mcarc_NL.18_23.rep



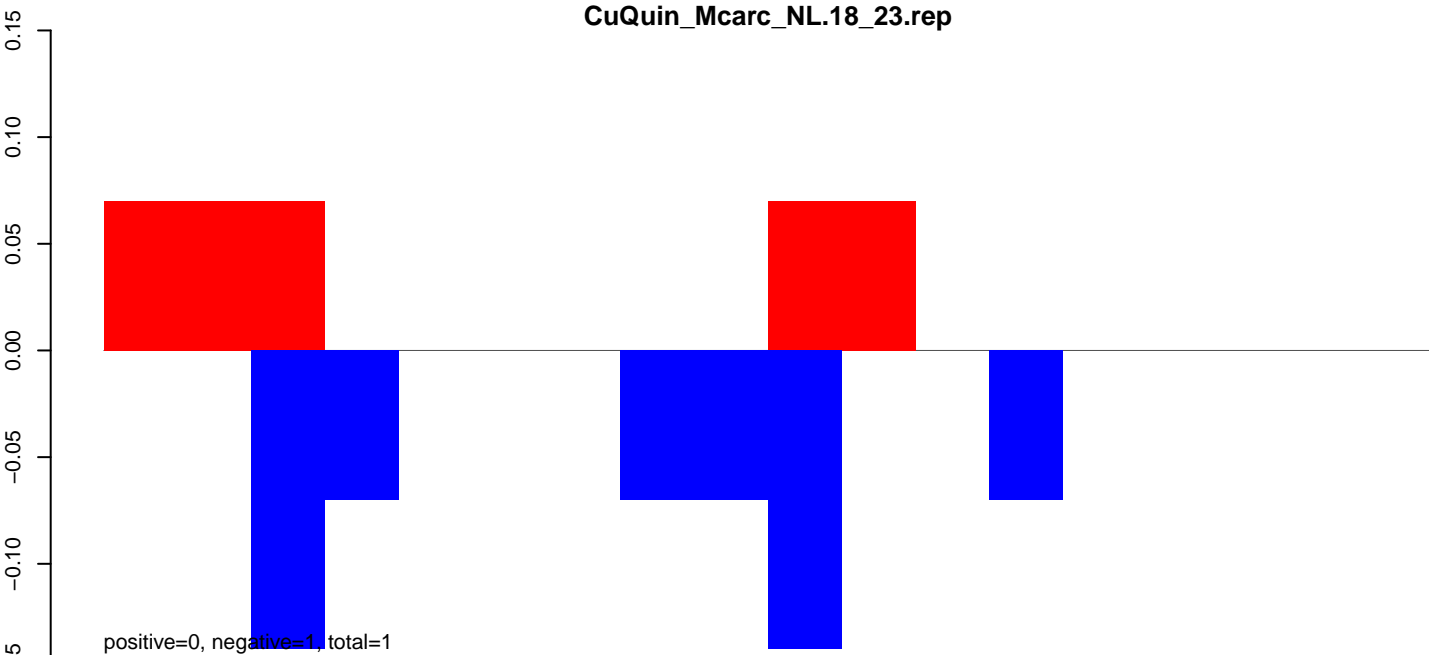
CuQuin_Mcarc_NL.24_35.rep



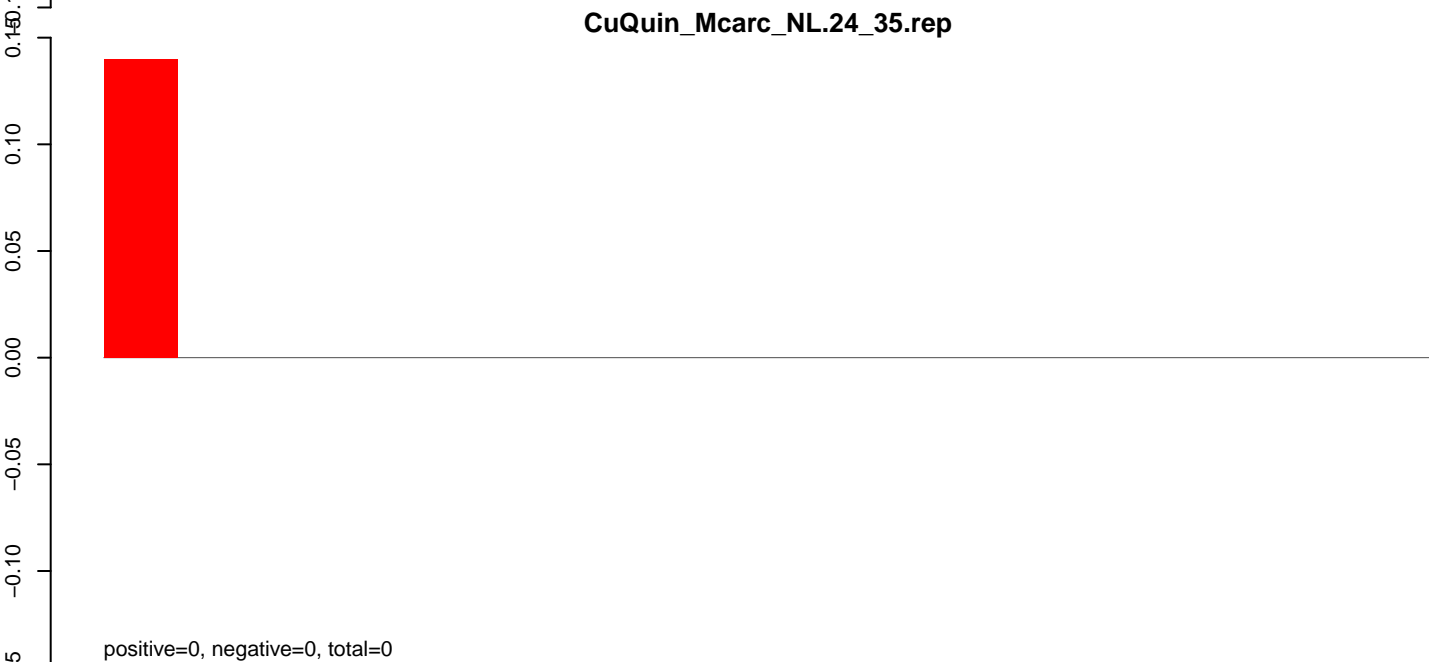
CuQuin_Mcarc_NL.rep



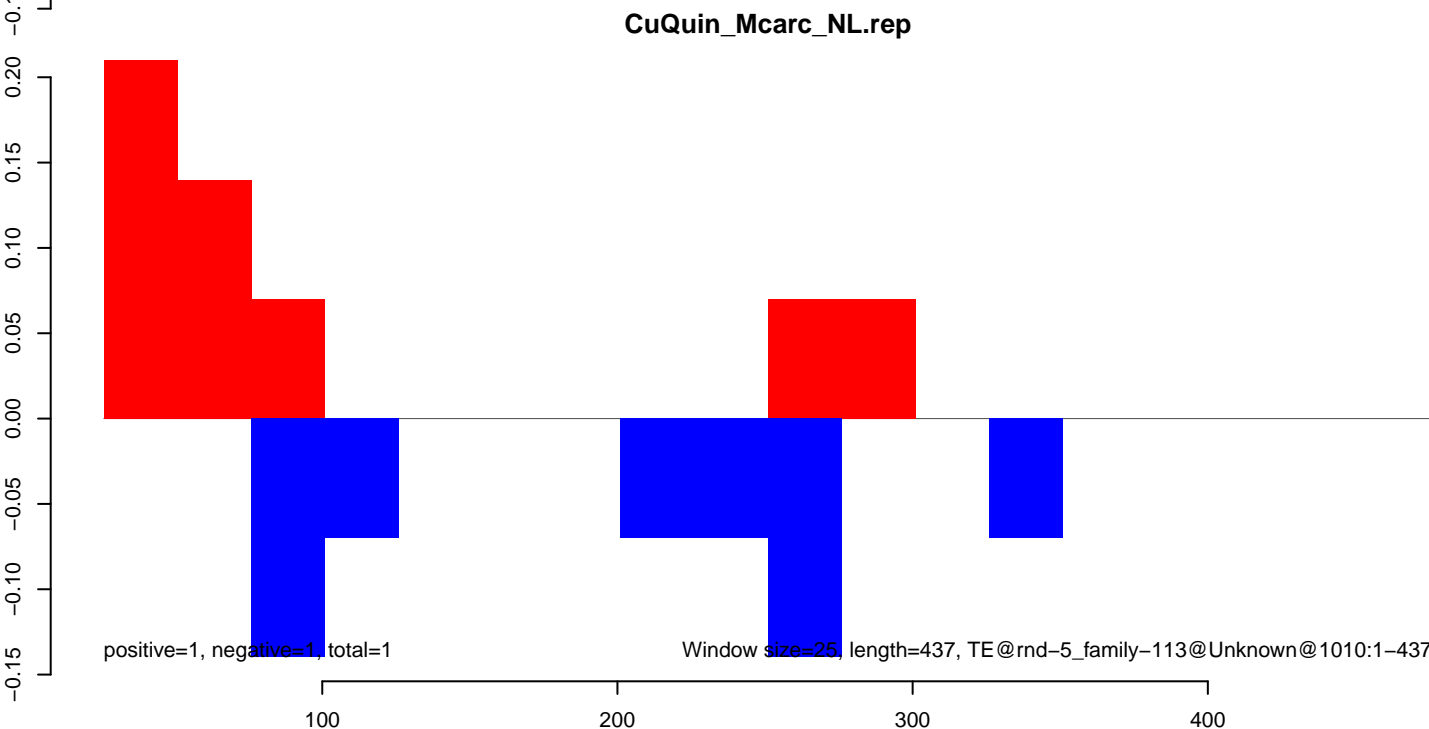
CuQuin_Mcarc_NL.18_23.rep

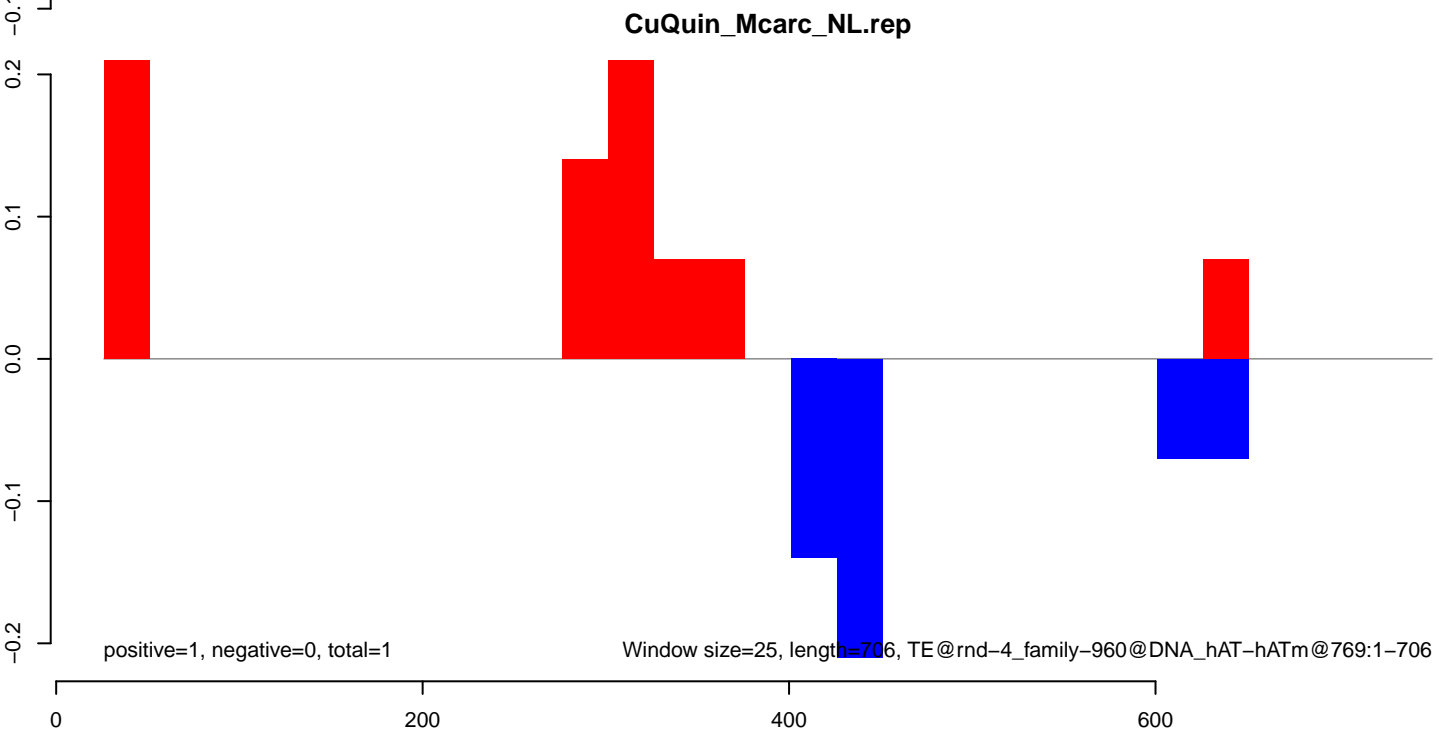
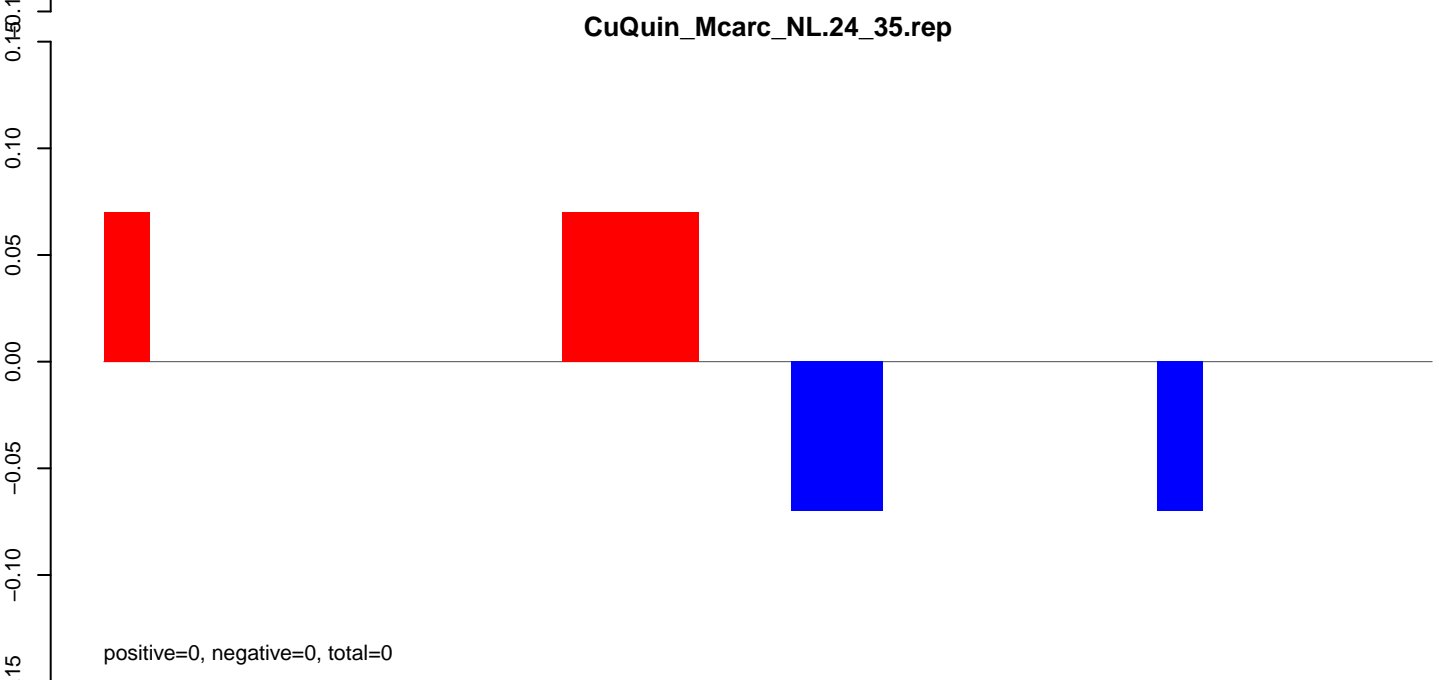
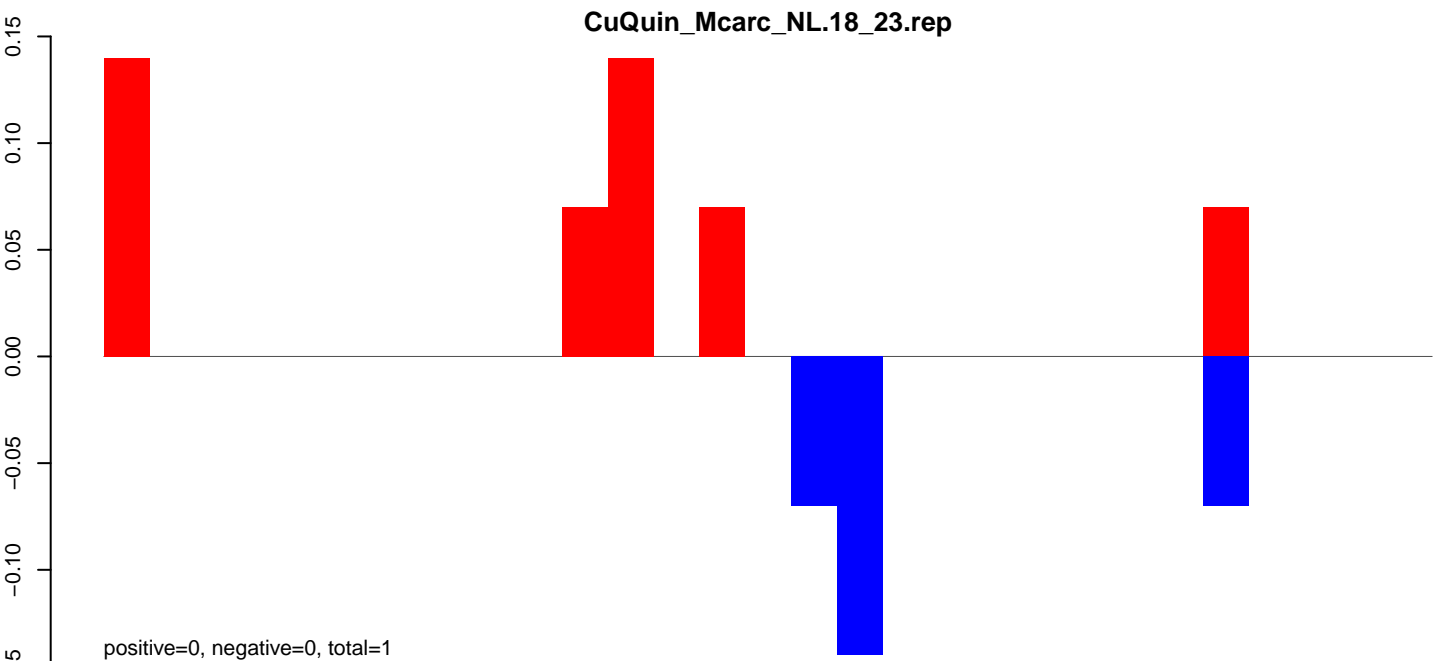


CuQuin_Mcarc_NL.24_35.rep

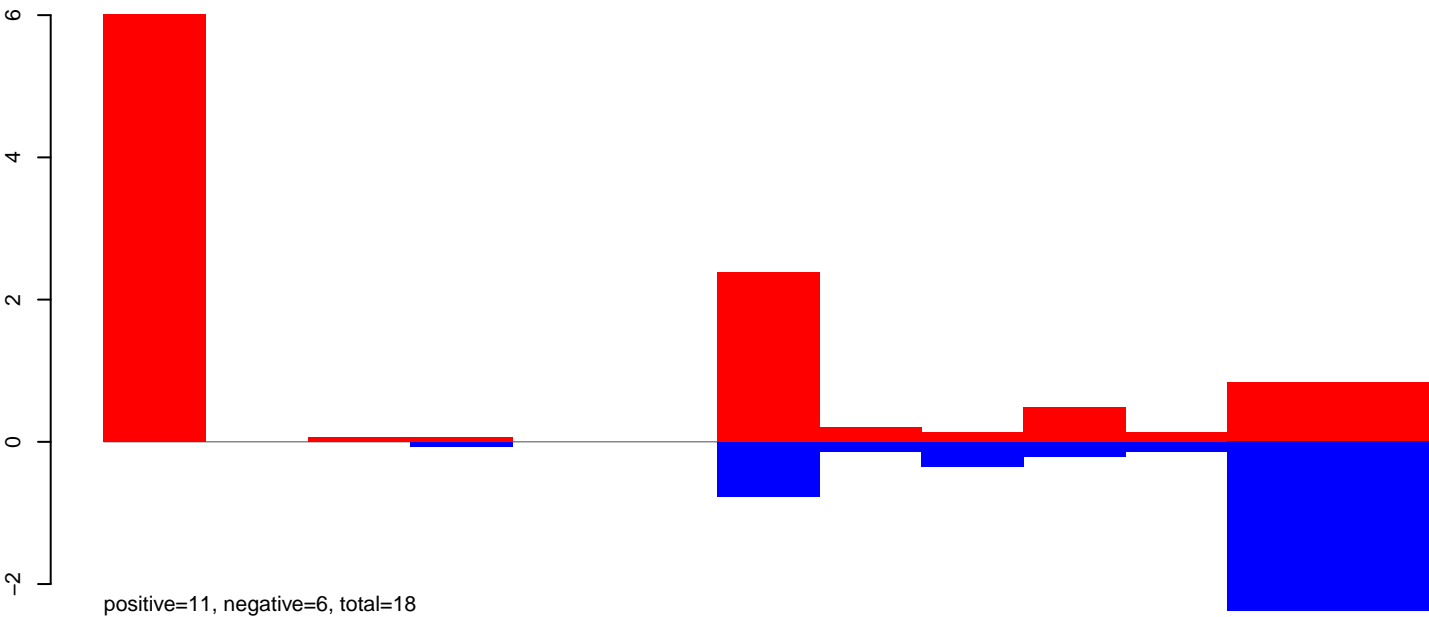


CuQuin_Mcarc_NL.rep

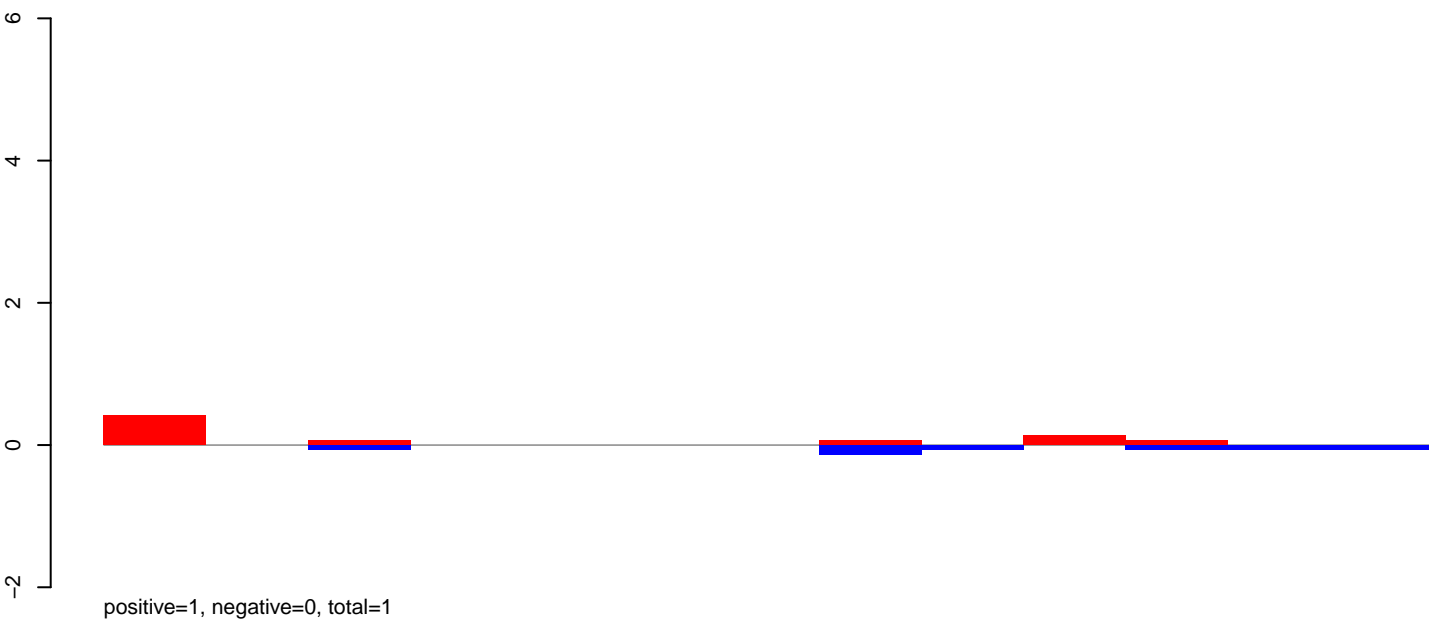




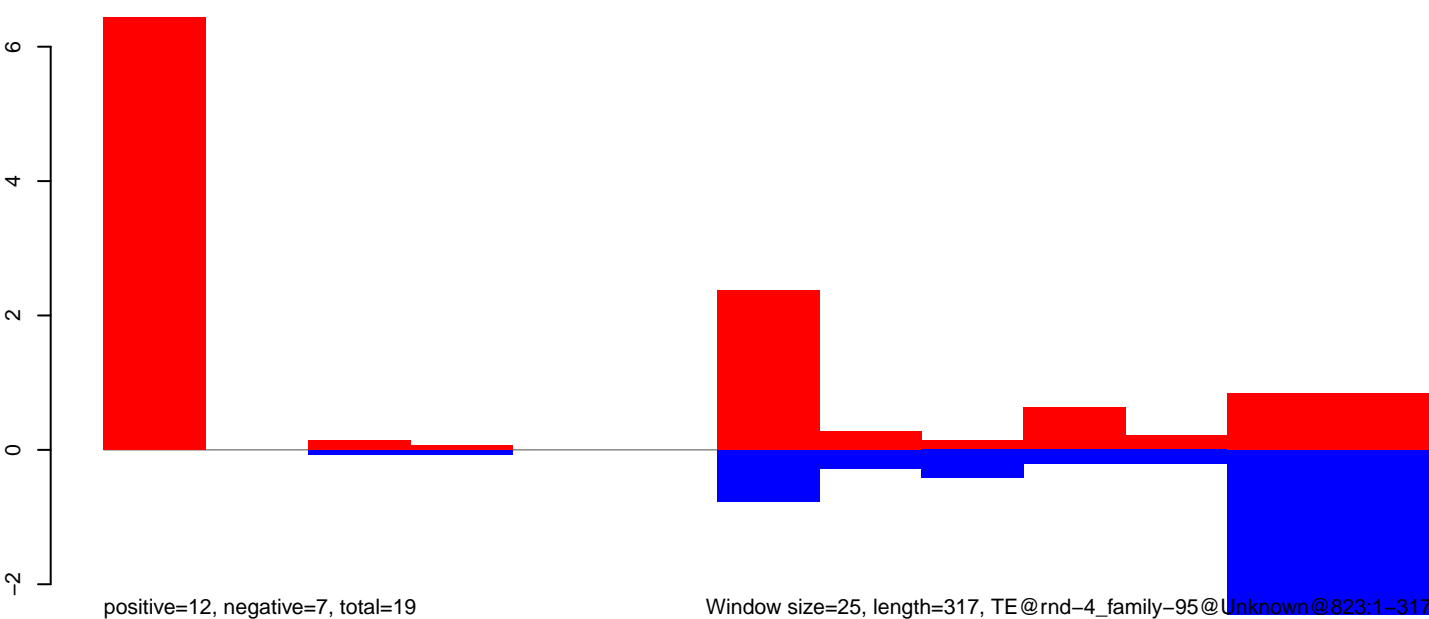
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

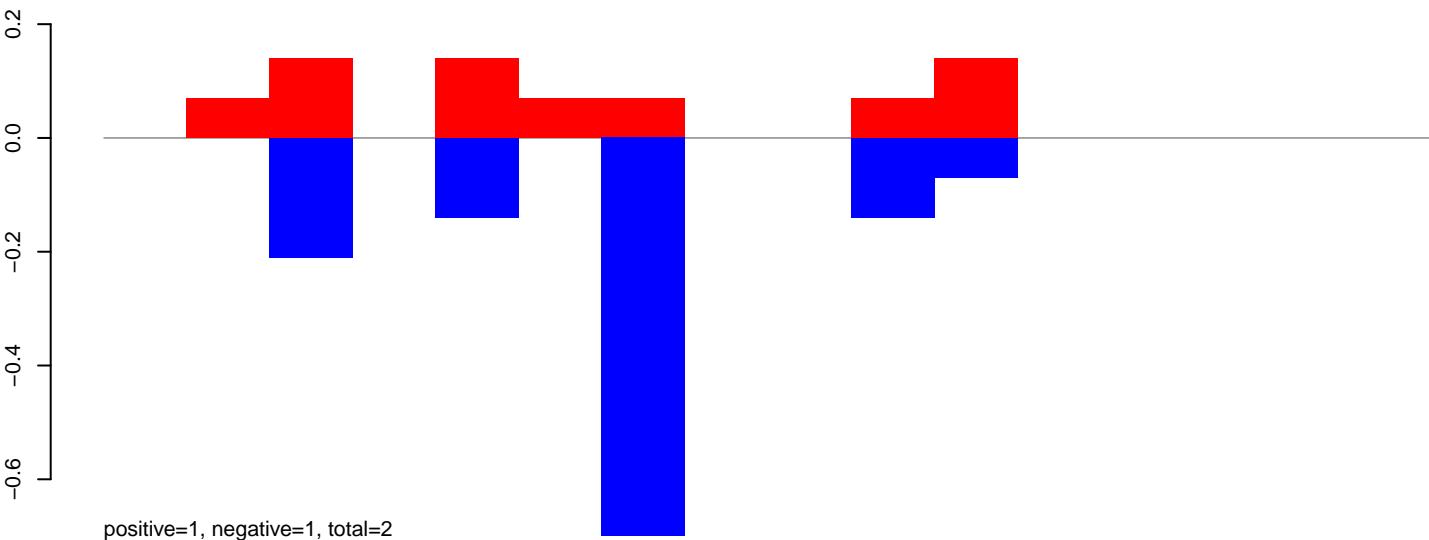


CuQuin_Mcarc_NL.rep

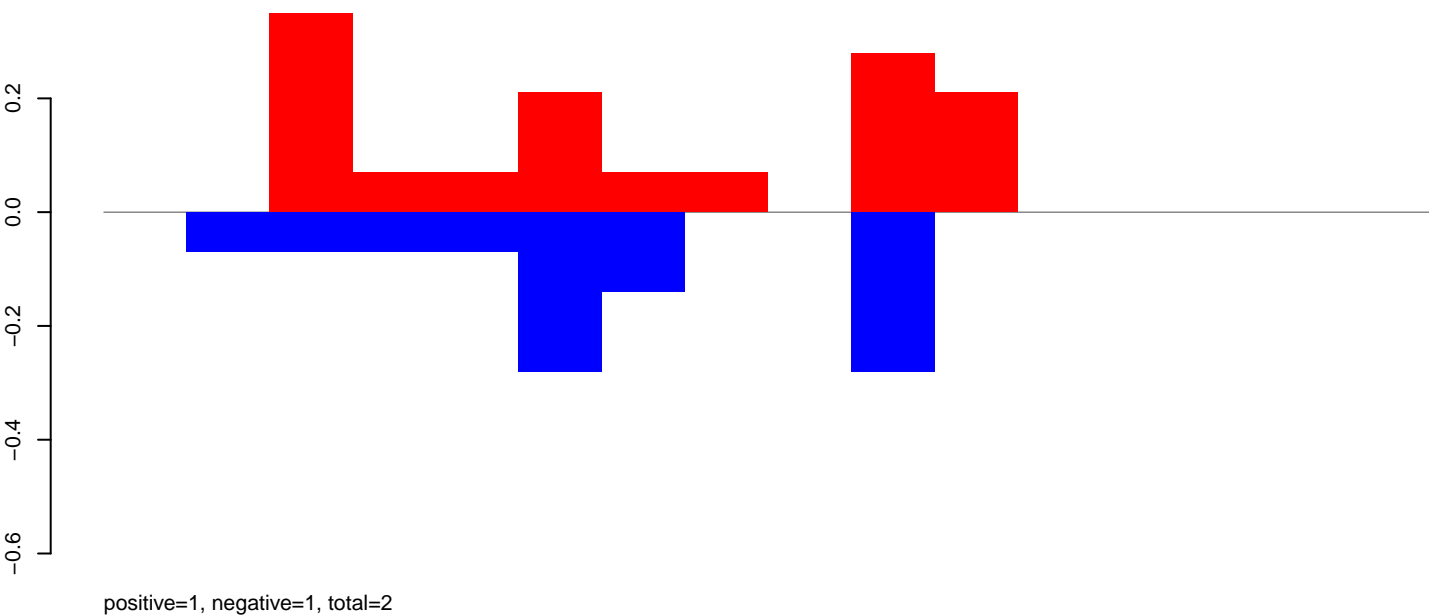


Window size=25, length=317, TE@rnd-4_family-95@Unknown@823:1-317

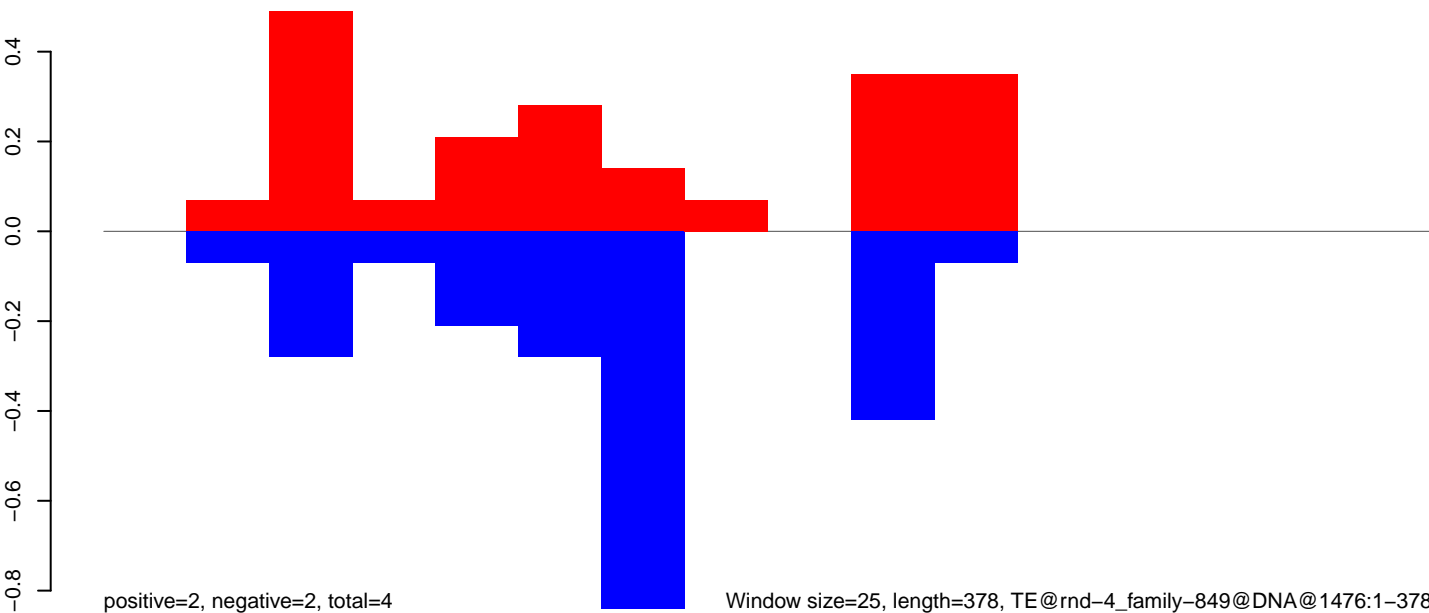
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=378, TE@rnd-4_family-849@DNA@1476:1-378

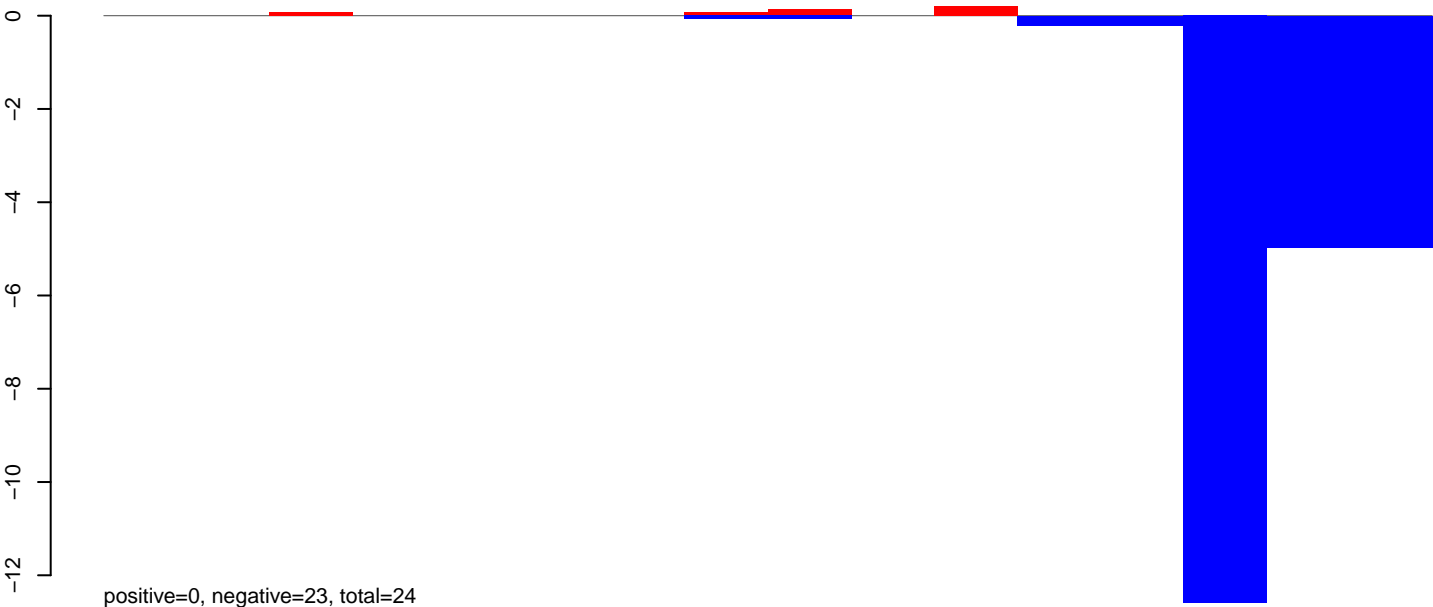
100

200

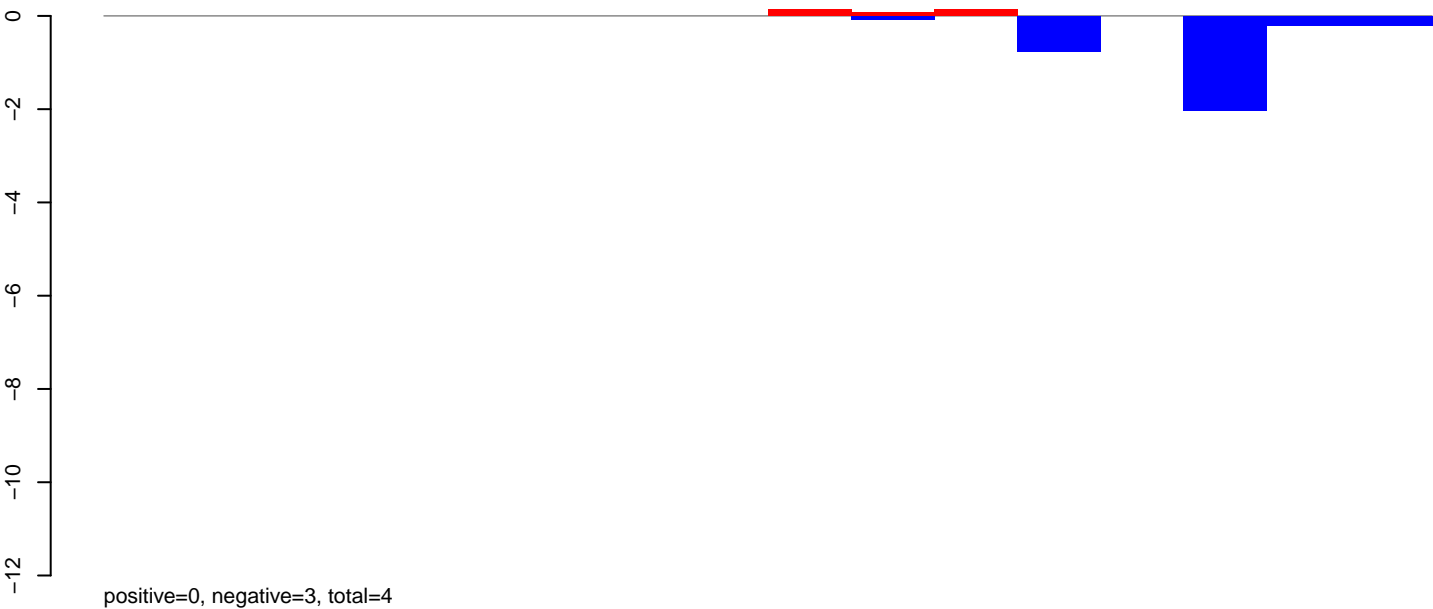
300

400

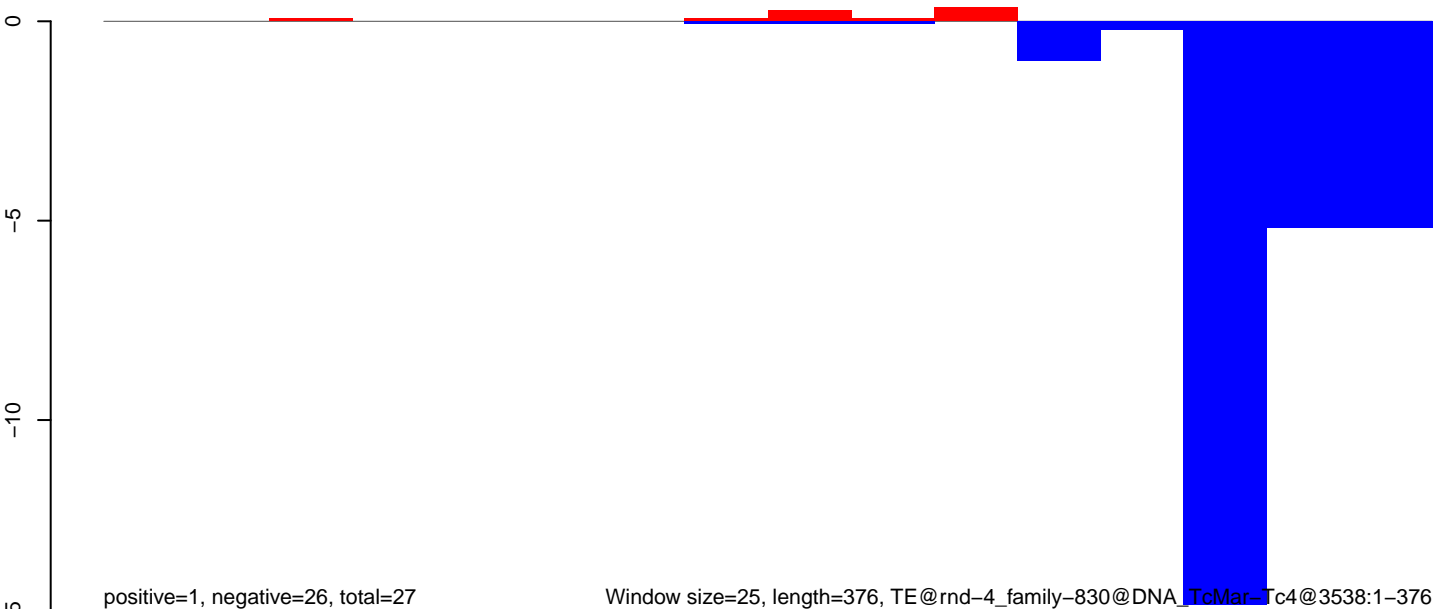
CuQuin_Mcarc_NL.18_23.rep



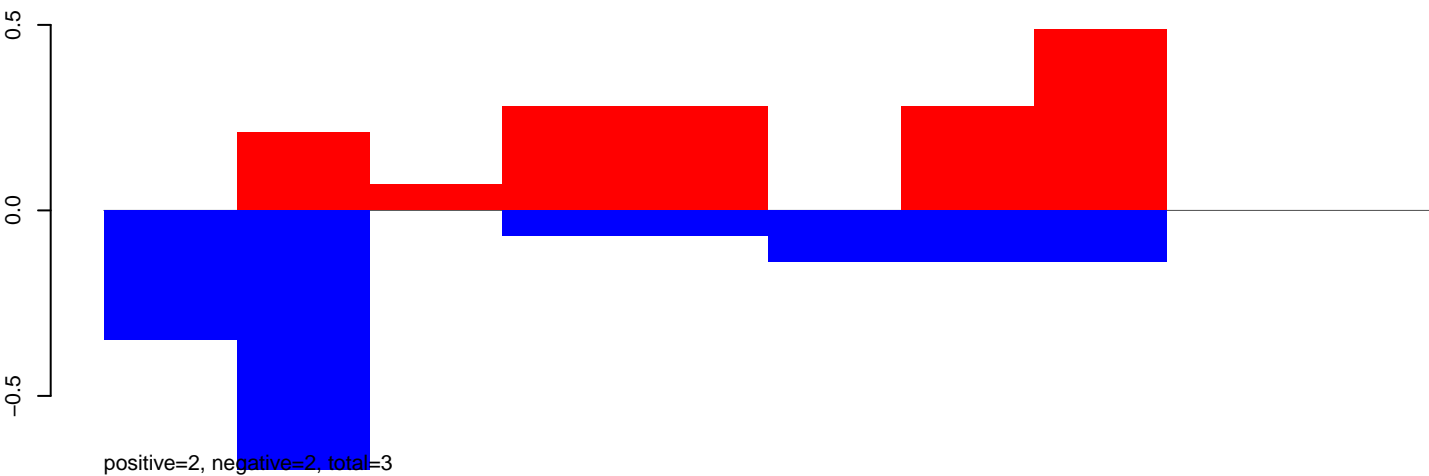
CuQuin_Mcarc_NL.24_35.rep



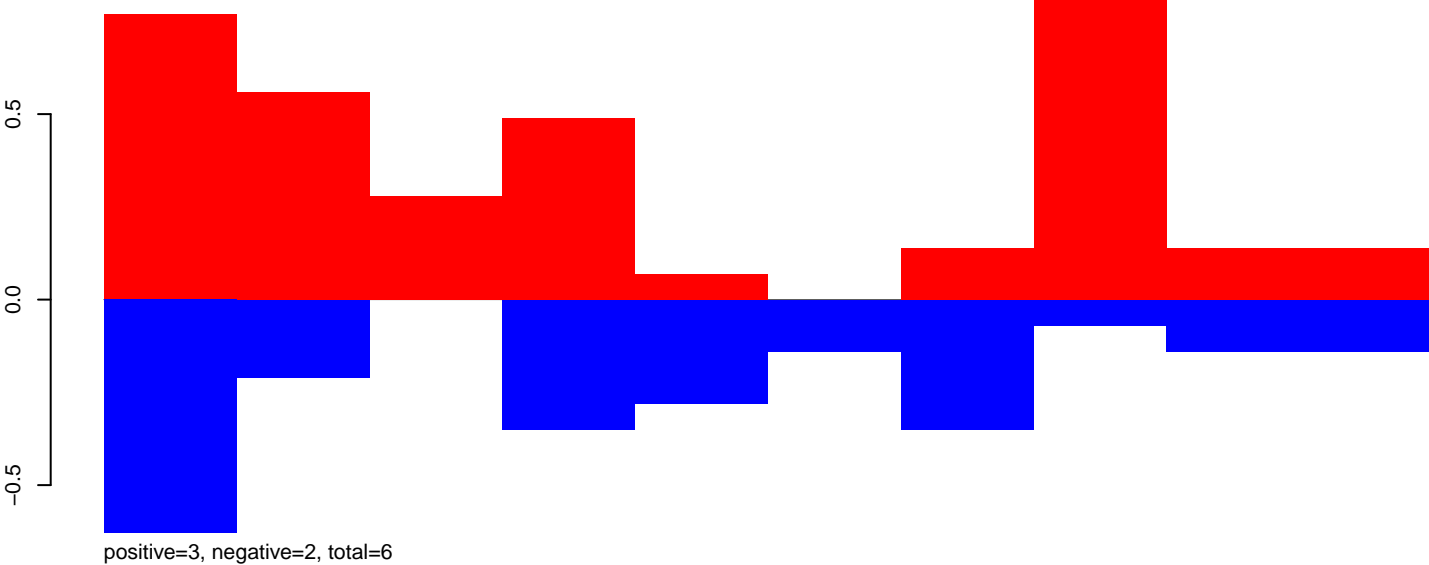
CuQuin_Mcarc_NL.rep



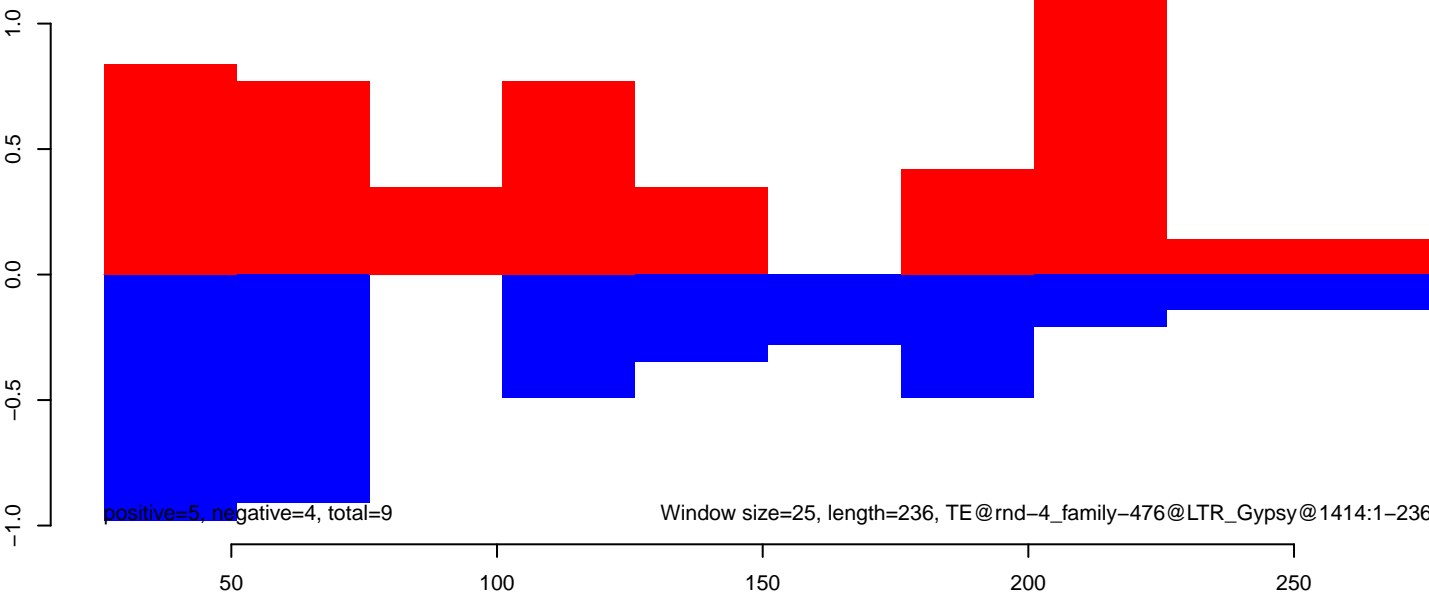
CuQuin_Mcarc_NL.18_23.rep



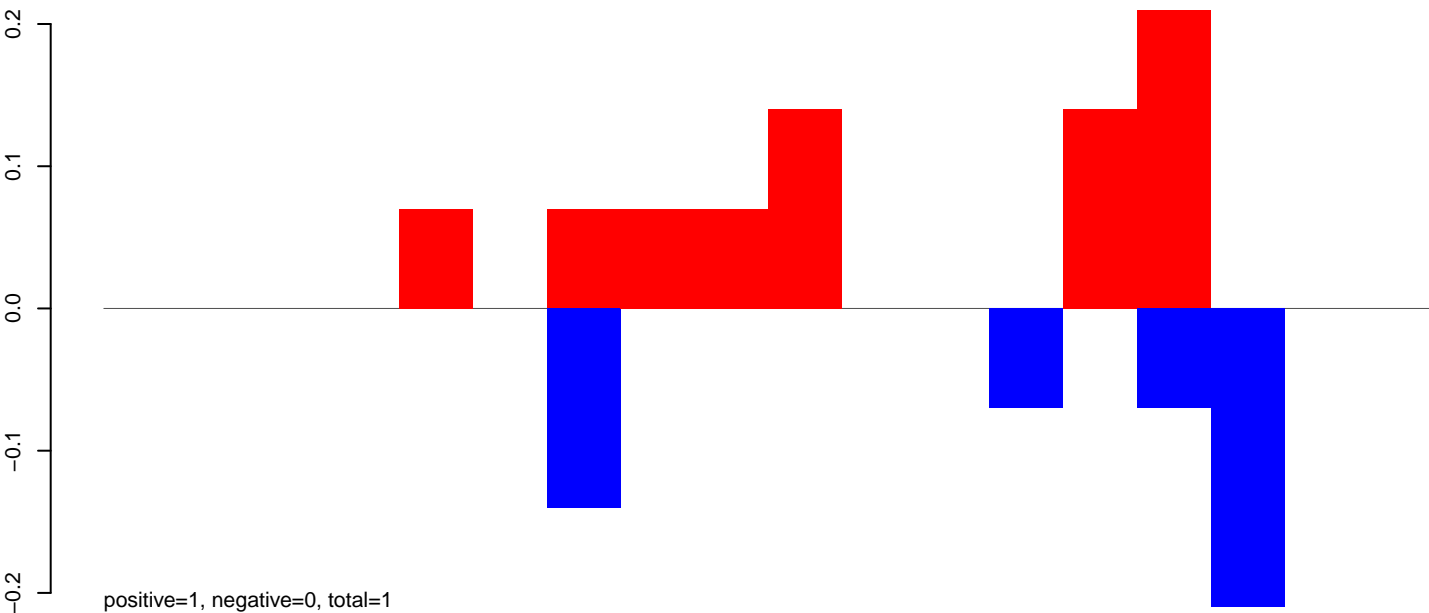
CuQuin_Mcarc_NL.24_35.rep



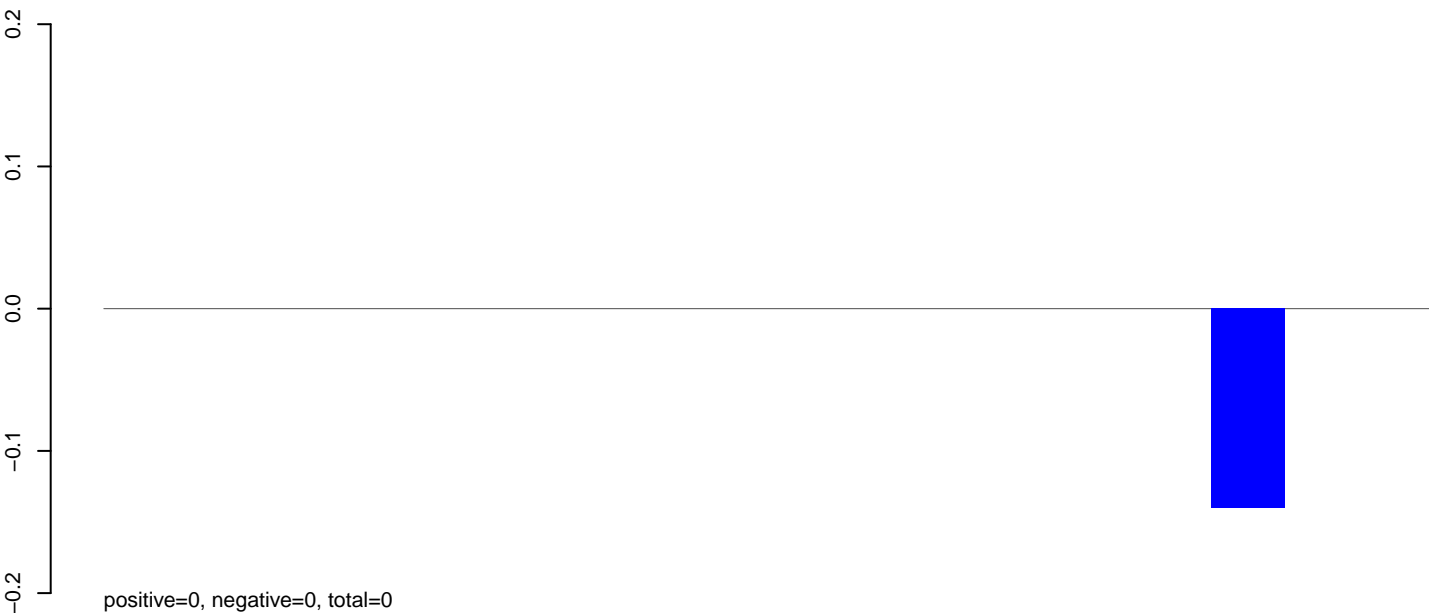
CuQuin_Mcarc_NL.rep



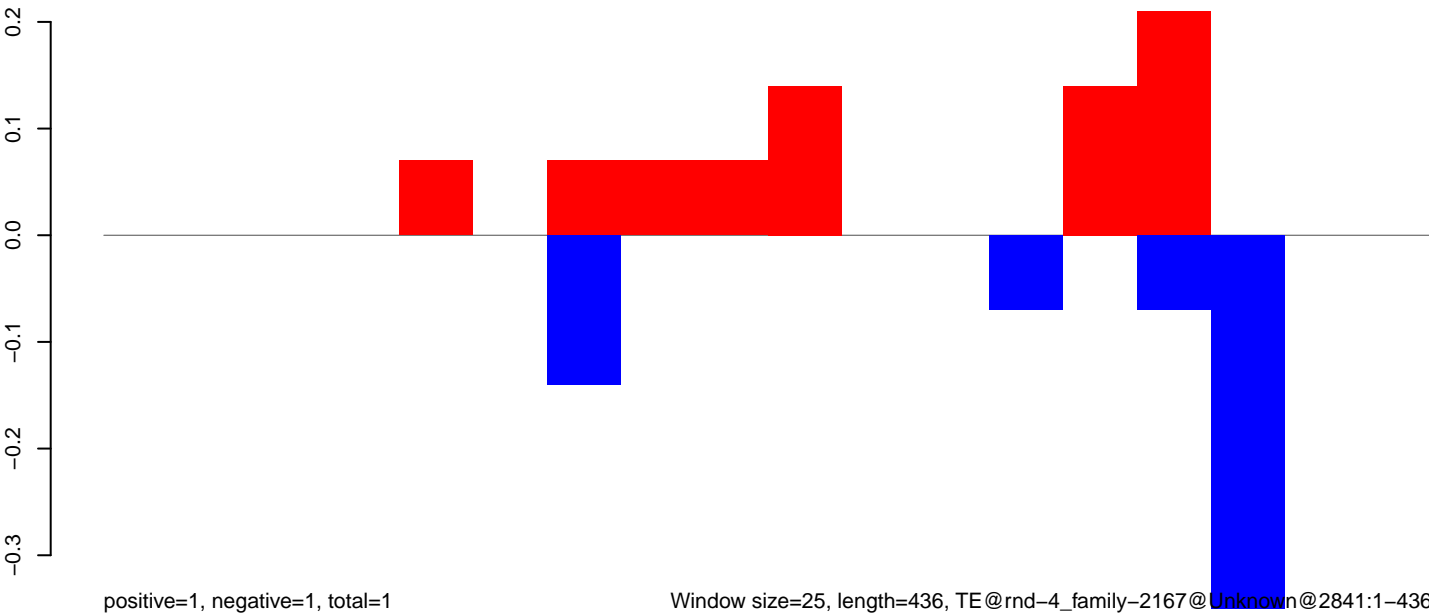
CuQuin_Mcarc_NL.18_23.rep



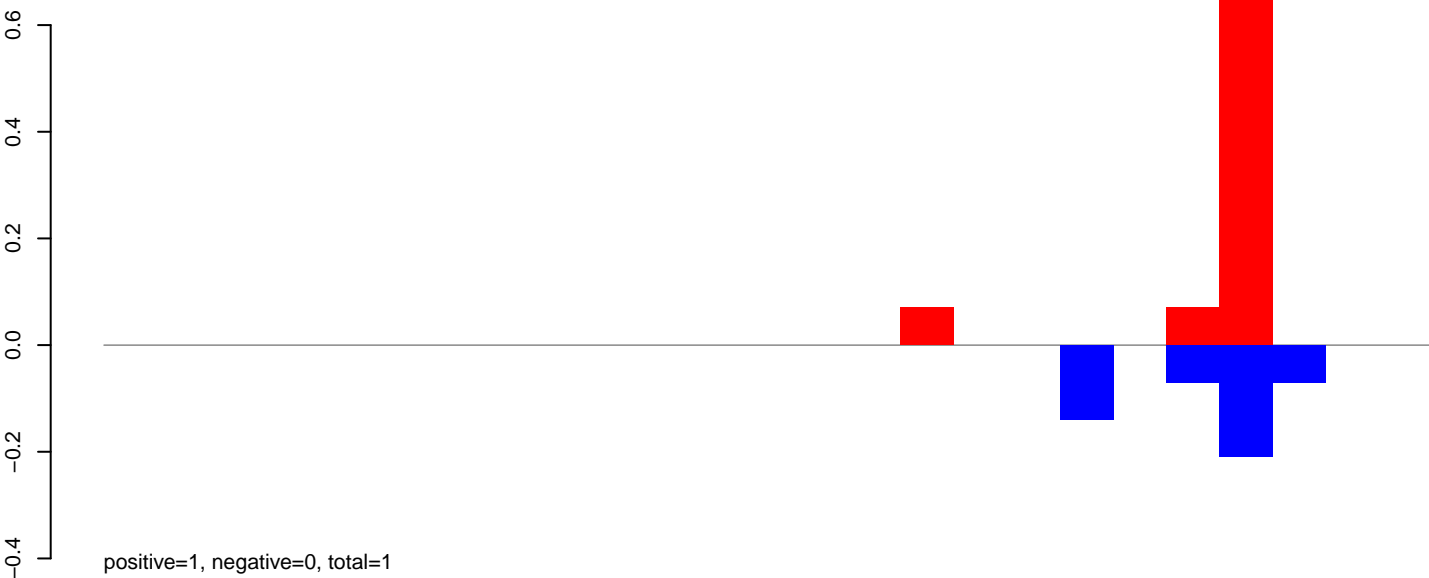
CuQuin_Mcarc_NL.24_35.rep



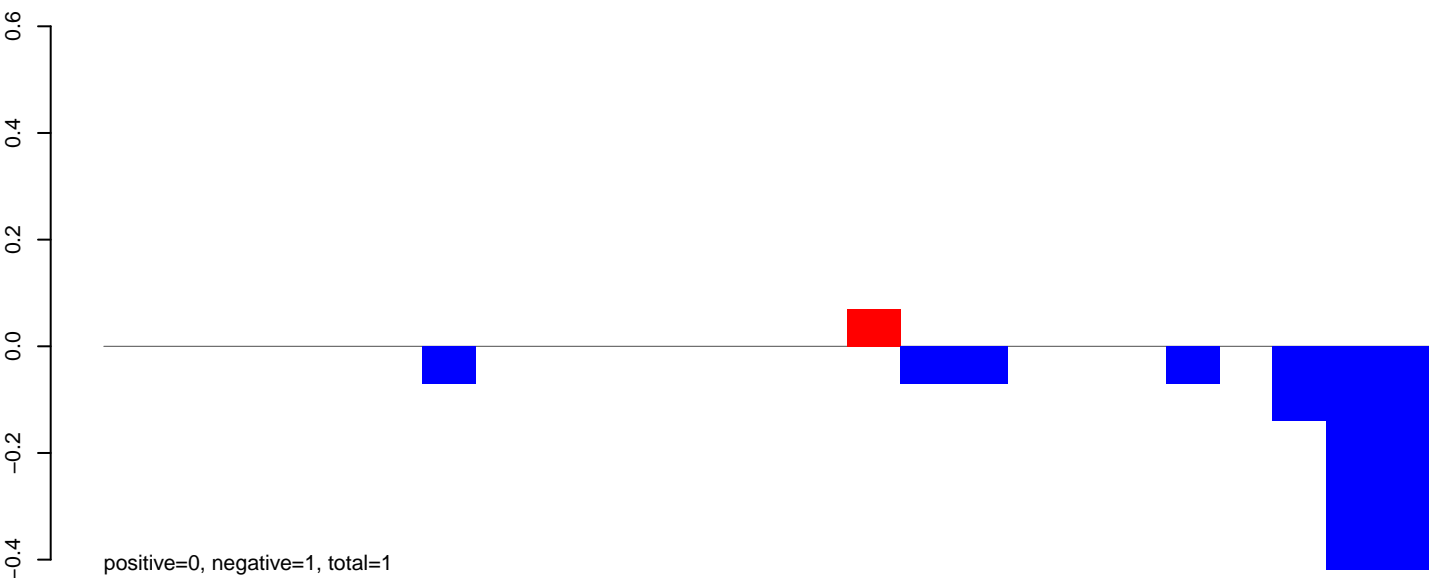
CuQuin_Mcarc_NL.rep



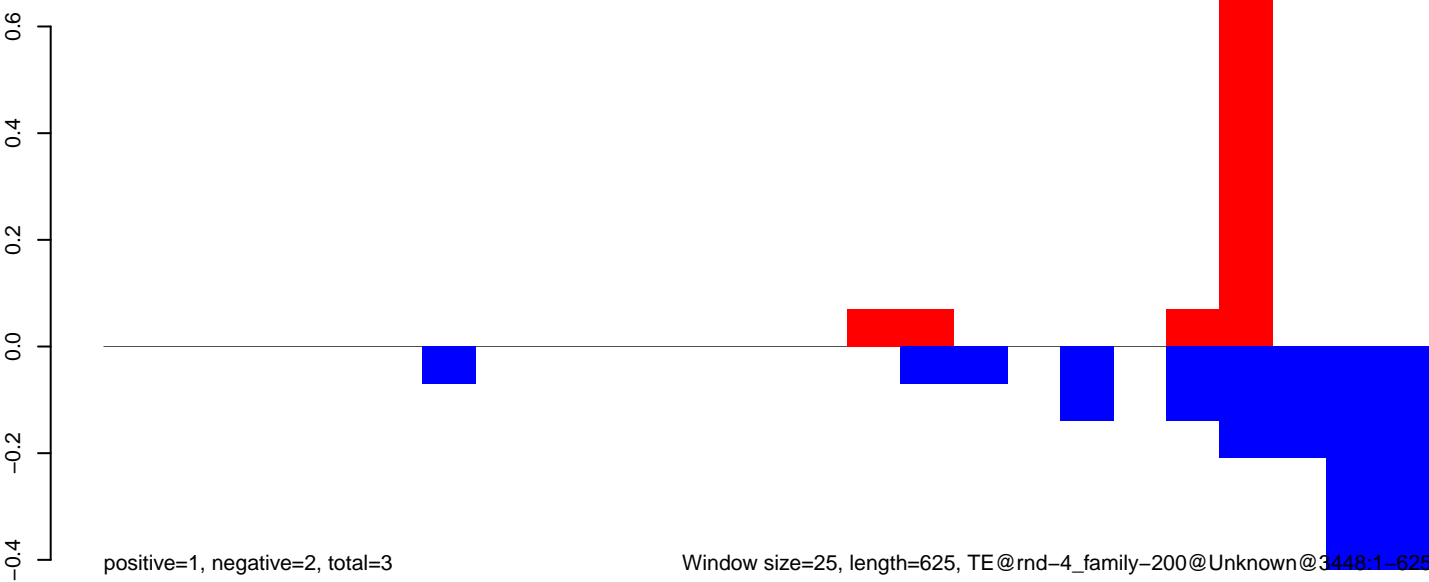
CuQuin_Mcarc_NL.18_23.rep



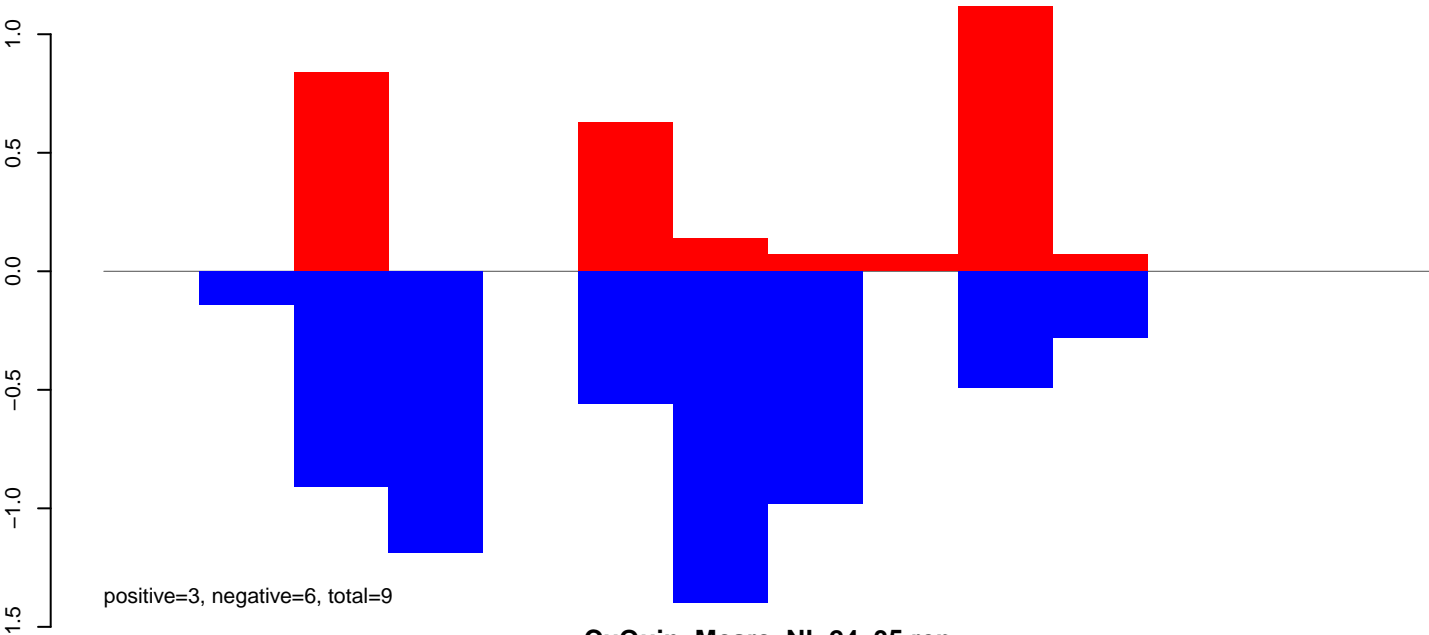
CuQuin_Mcarc_NL.24_35.rep



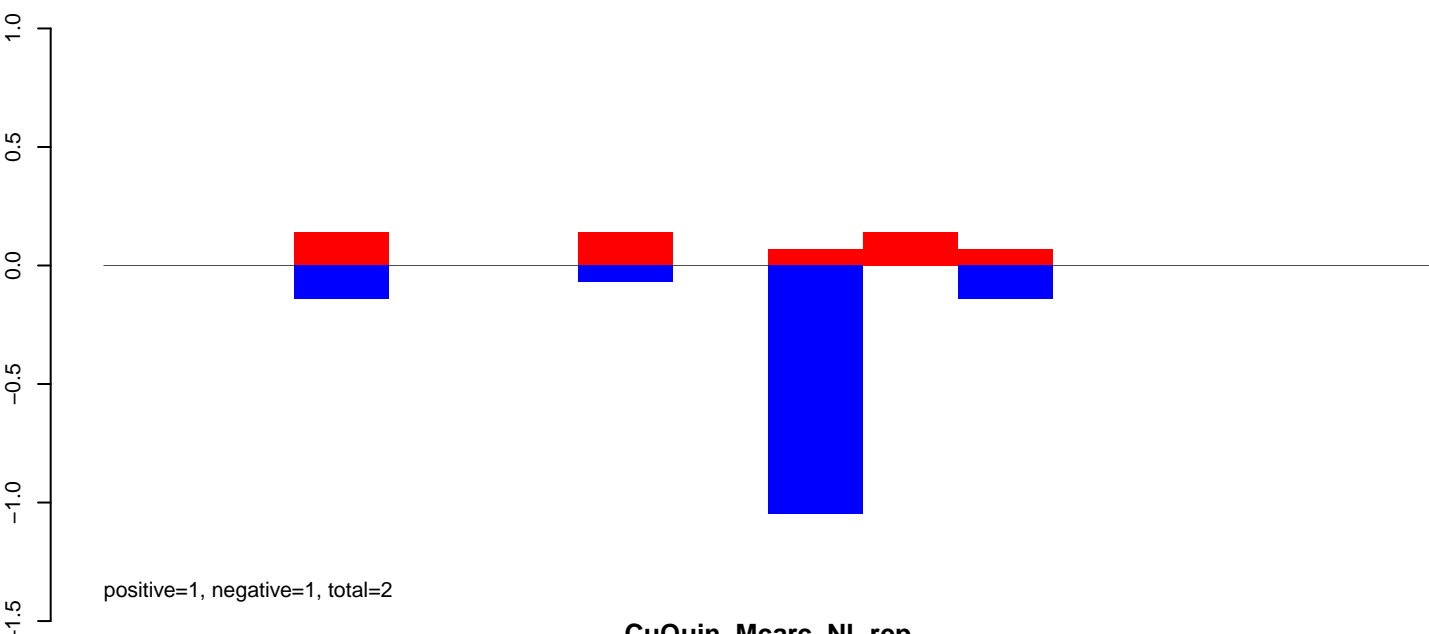
CuQuin_Mcarc_NL.rep



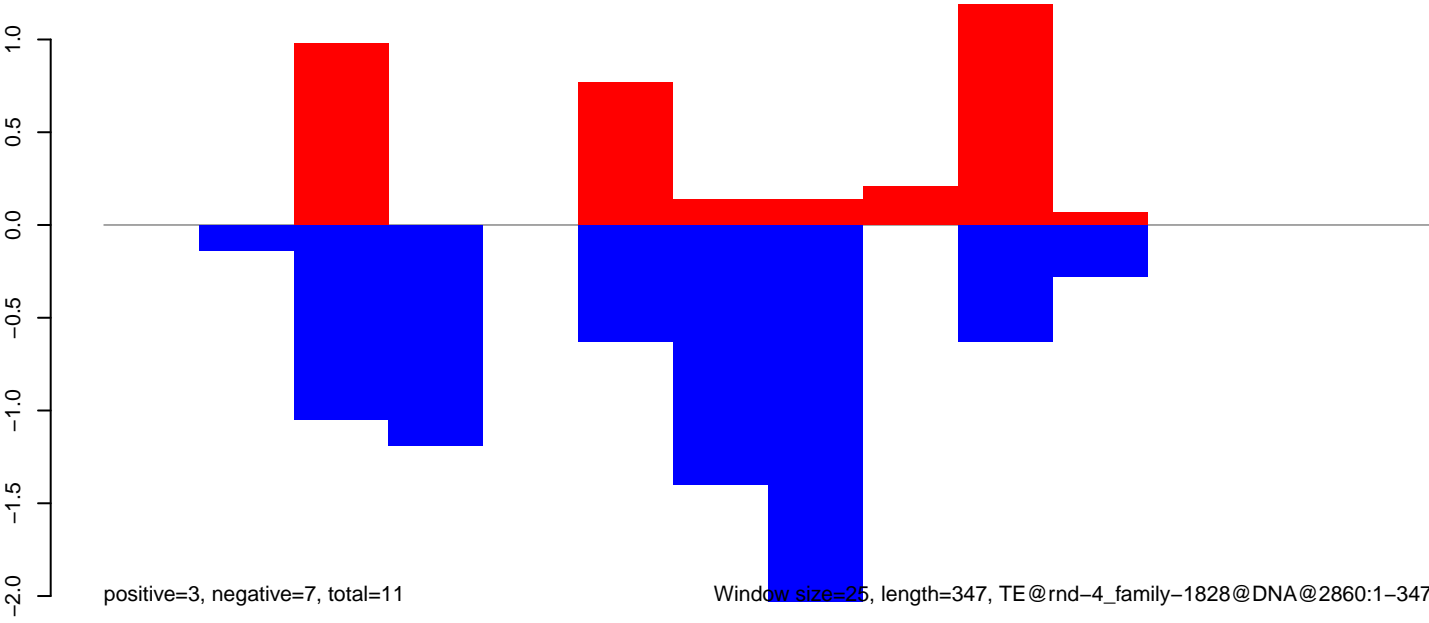
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

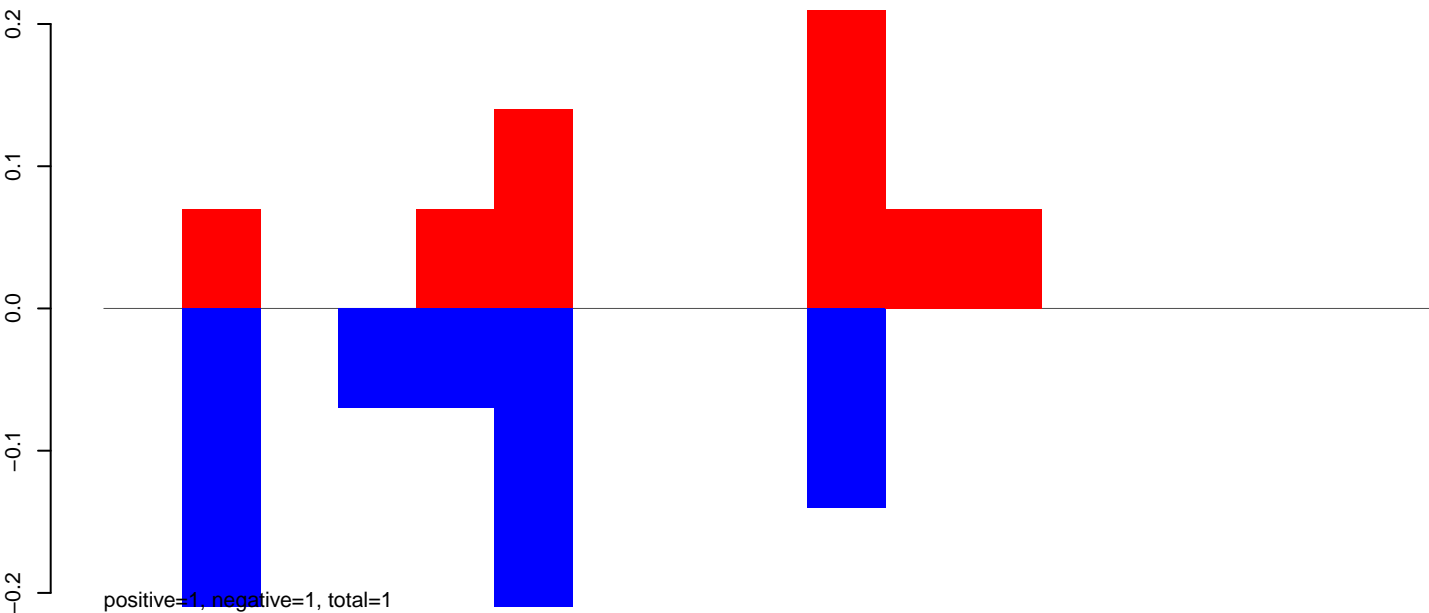


CuQuin_Mcarc_NL.rep

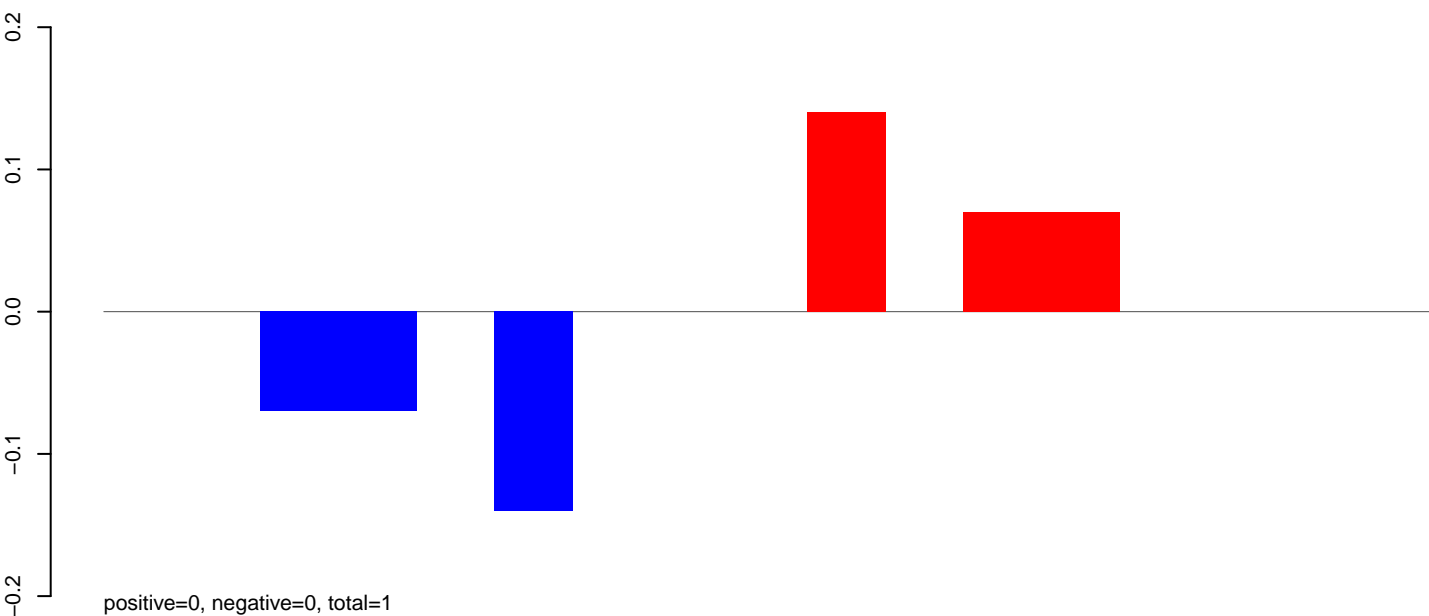


Window size=25, length=347, TE@rnd-4_family-1828@DNA@2860:1-347

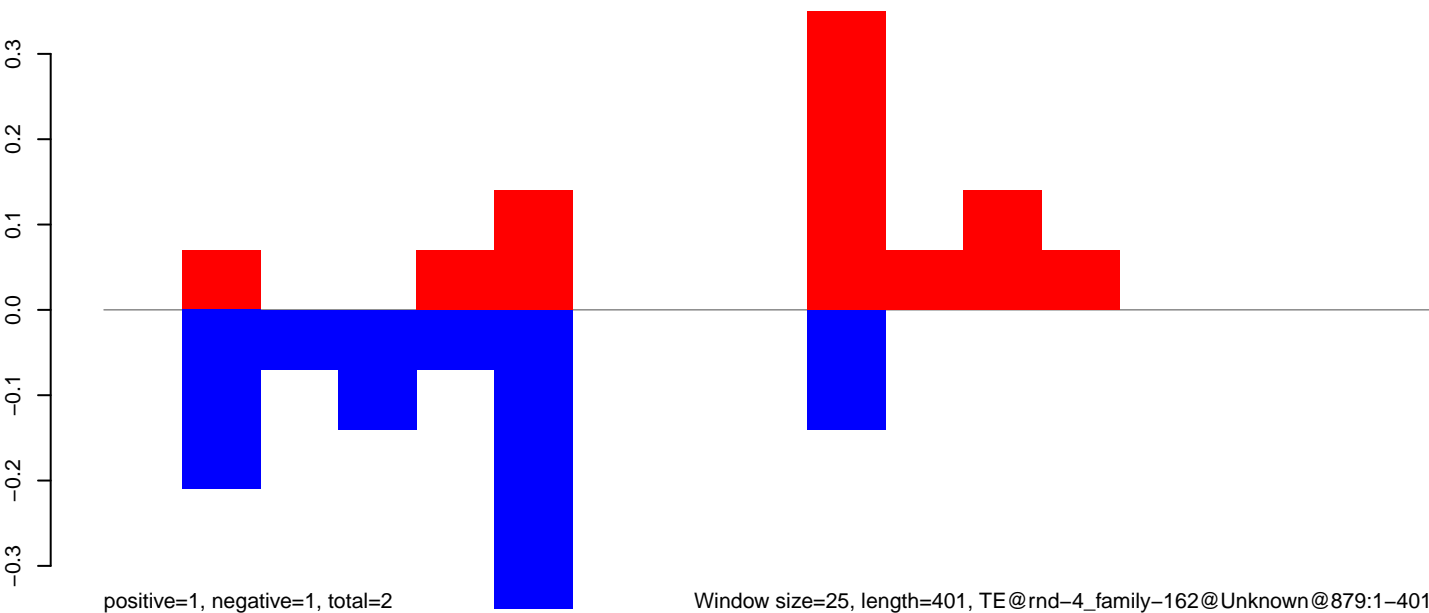
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=401, TE@rnd-4_family-162@Unknown@879:1-401

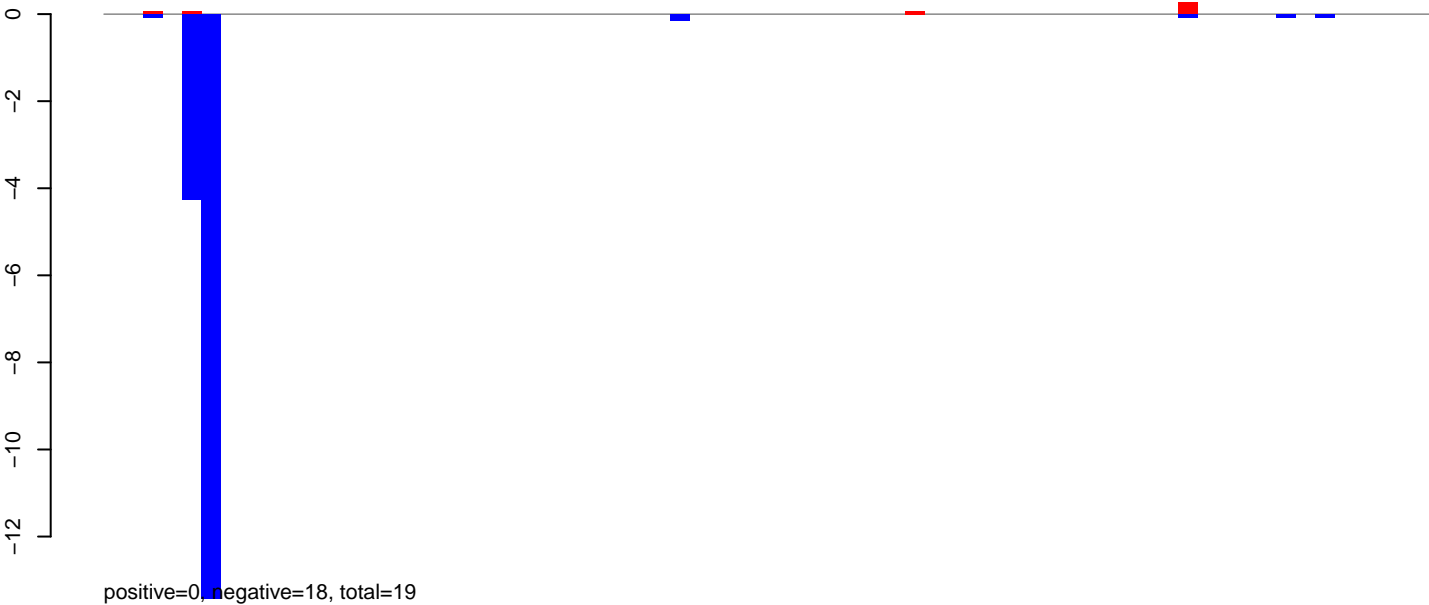
100

200

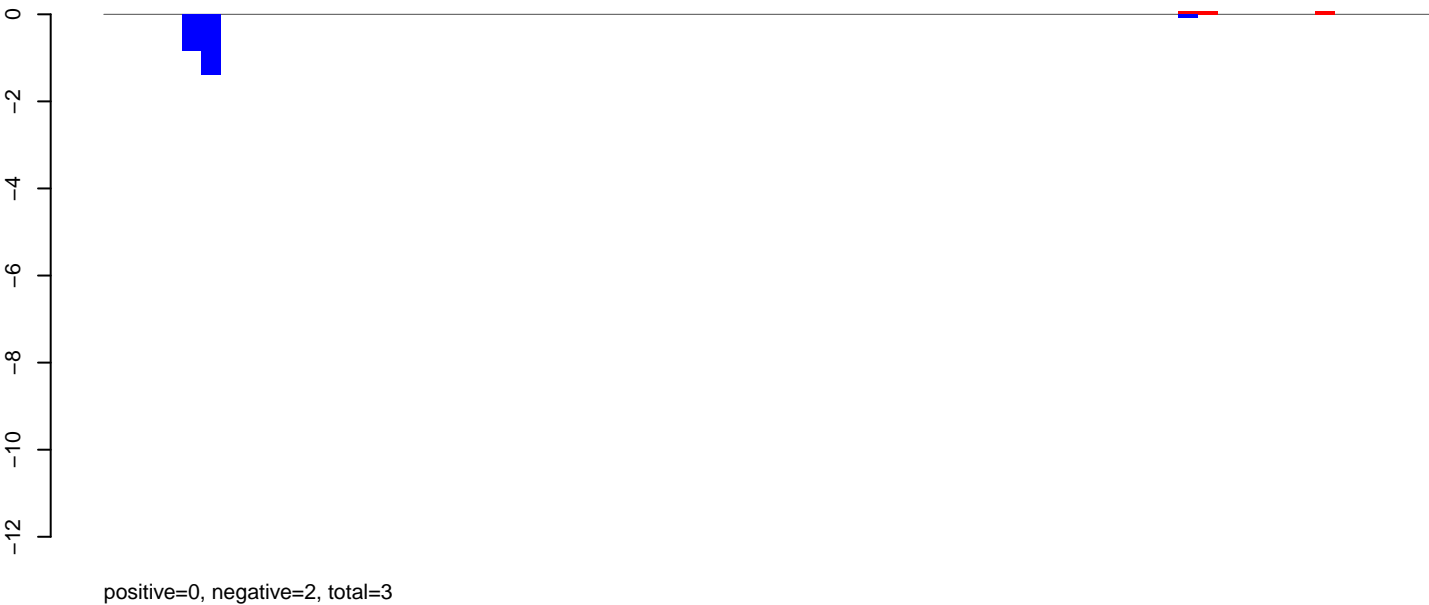
300

400

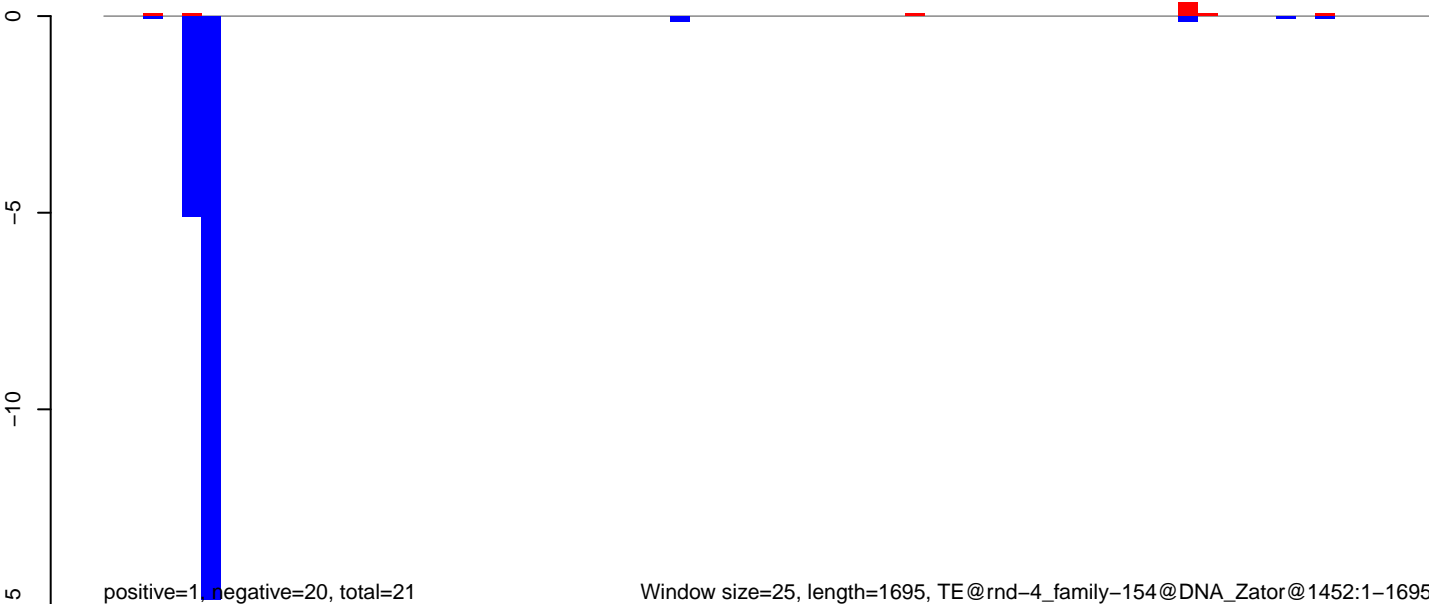
CuQuin_Mcarc_NL.18_23.rep



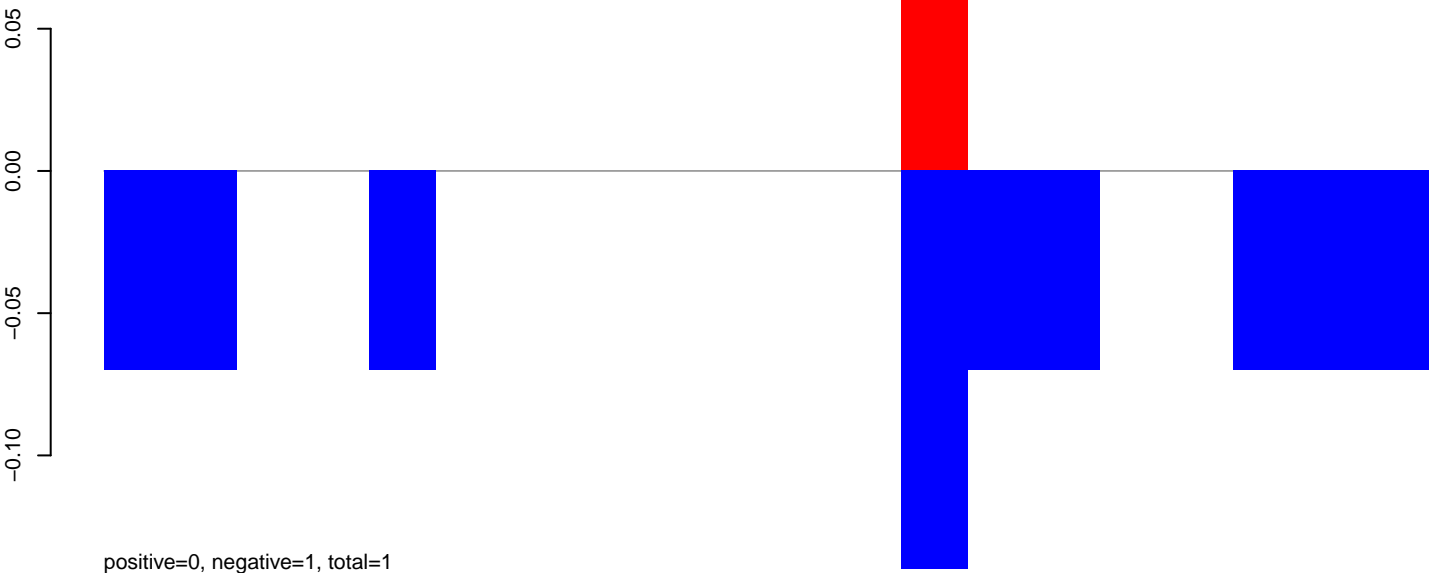
CuQuin_Mcarc_NL.24_35.rep



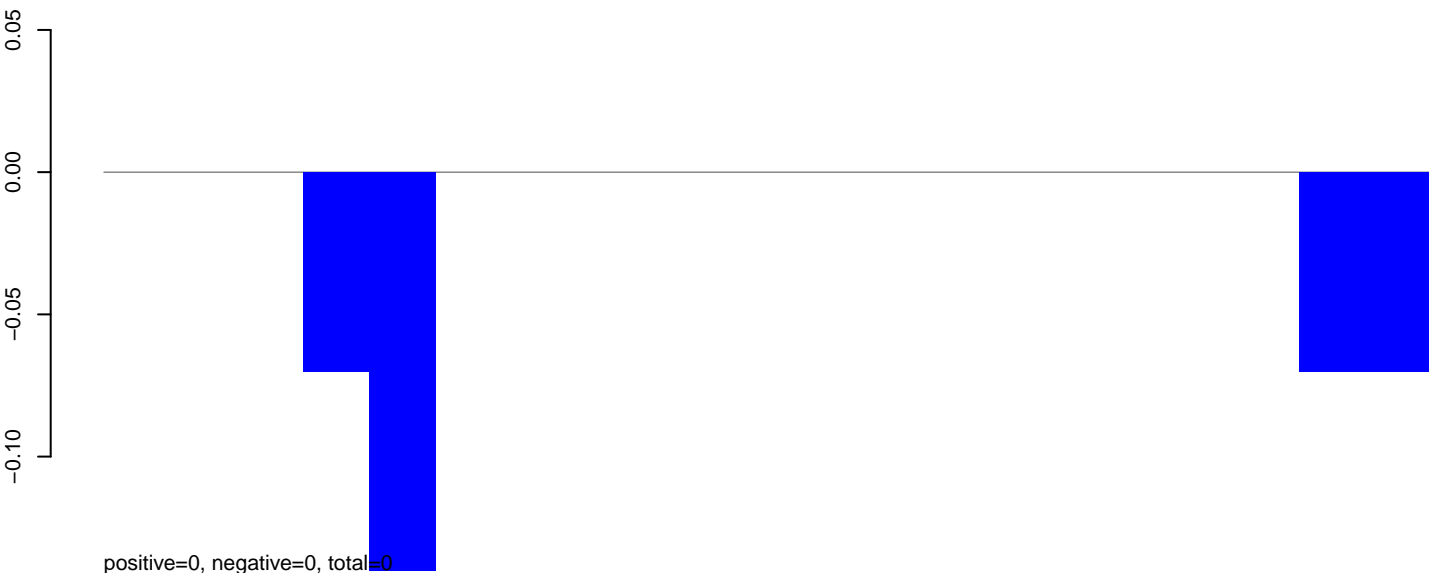
CuQuin_Mcarc_NL.rep



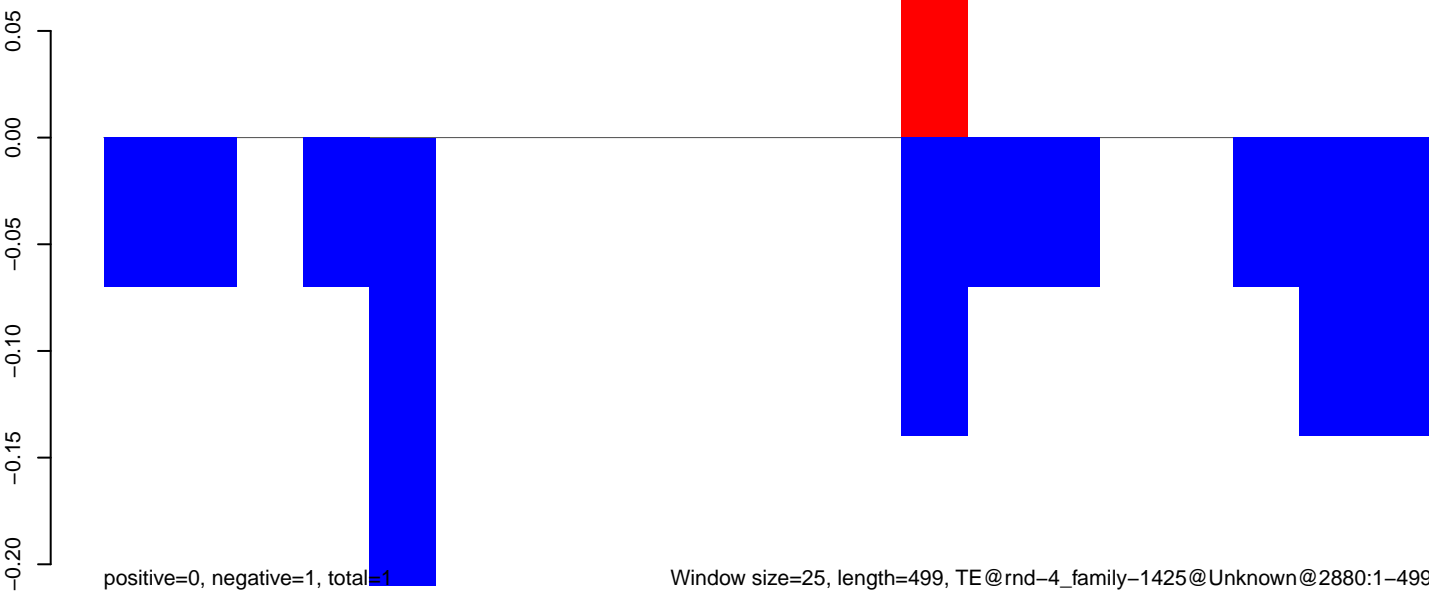
CuQuin_Mcarc_NL.18_23.rep



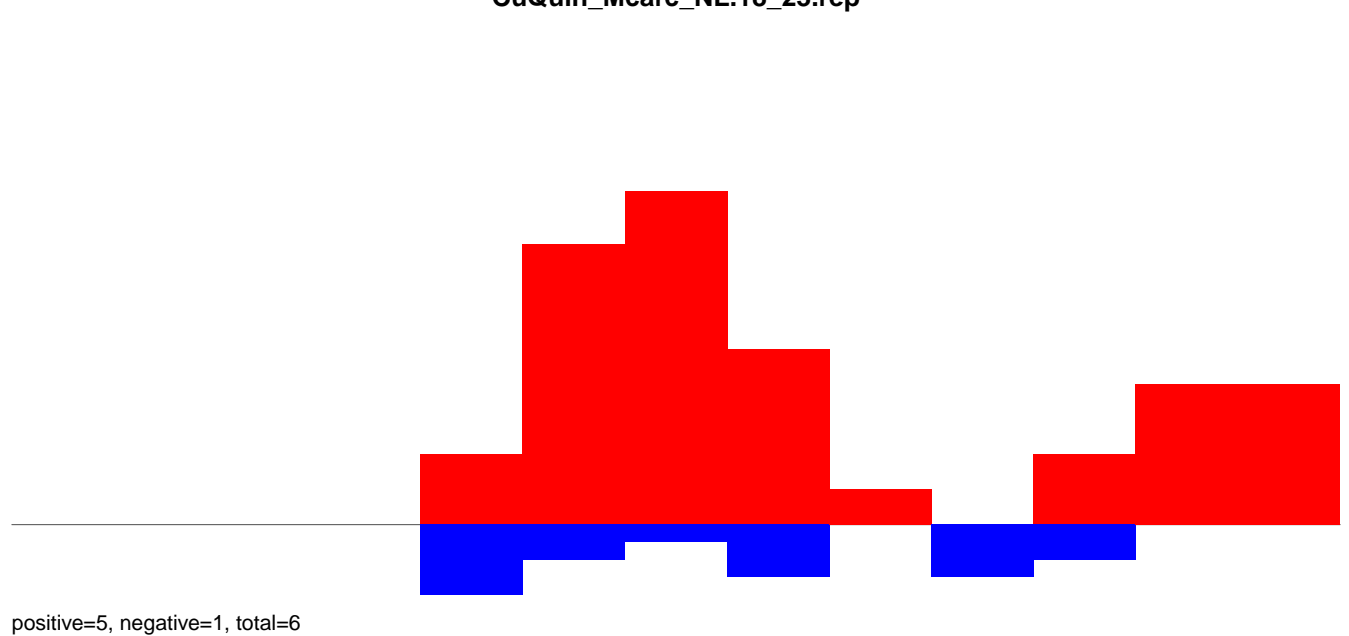
CuQuin_Mcarc_NL.24_35.rep



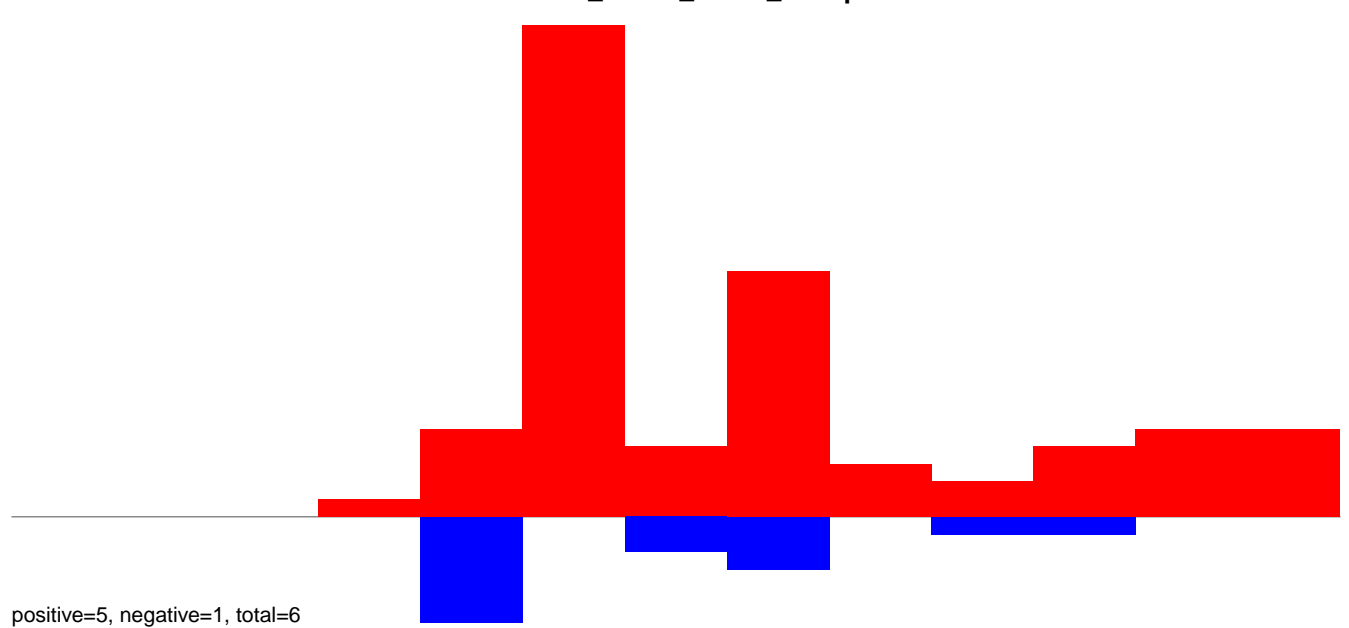
CuQuin_Mcarc_NL.rep



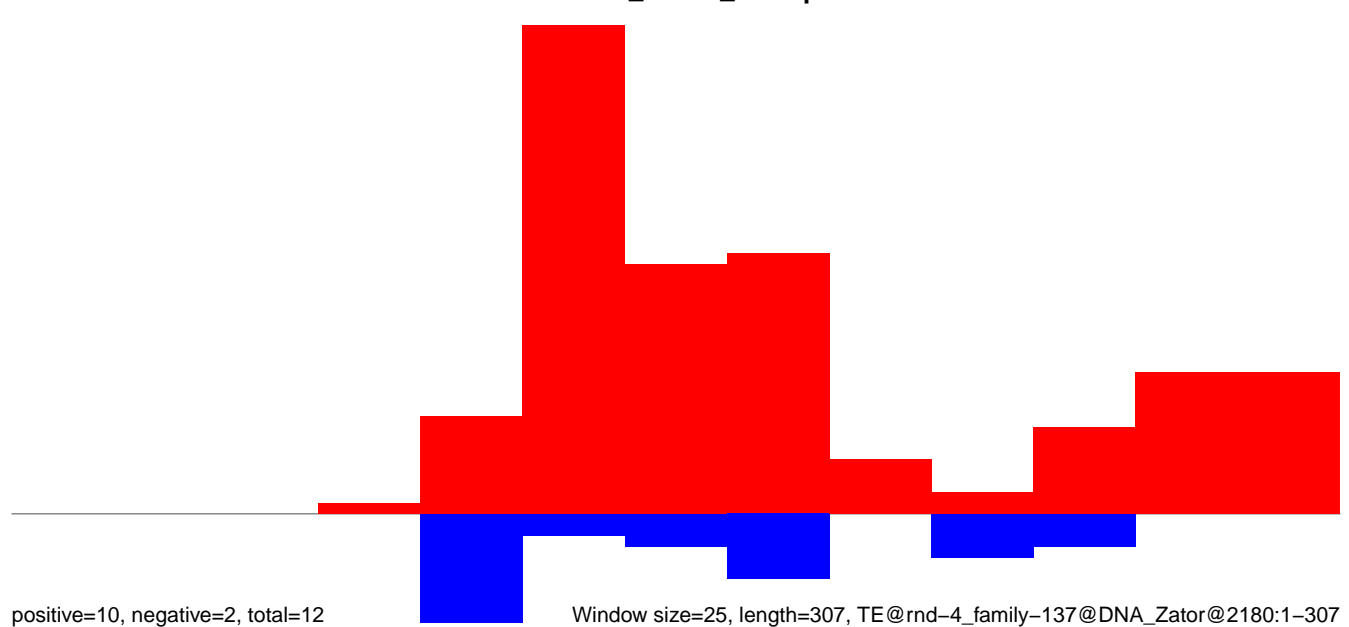
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



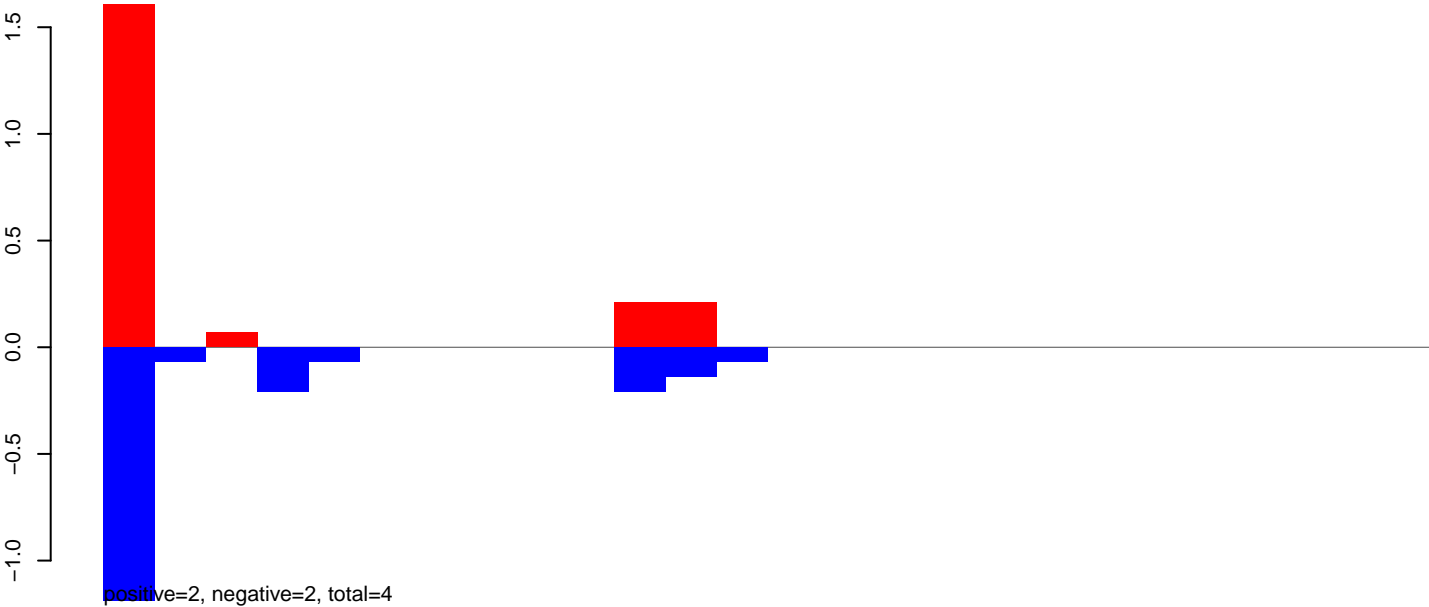
CuQuin_Mcarc_NL.rep



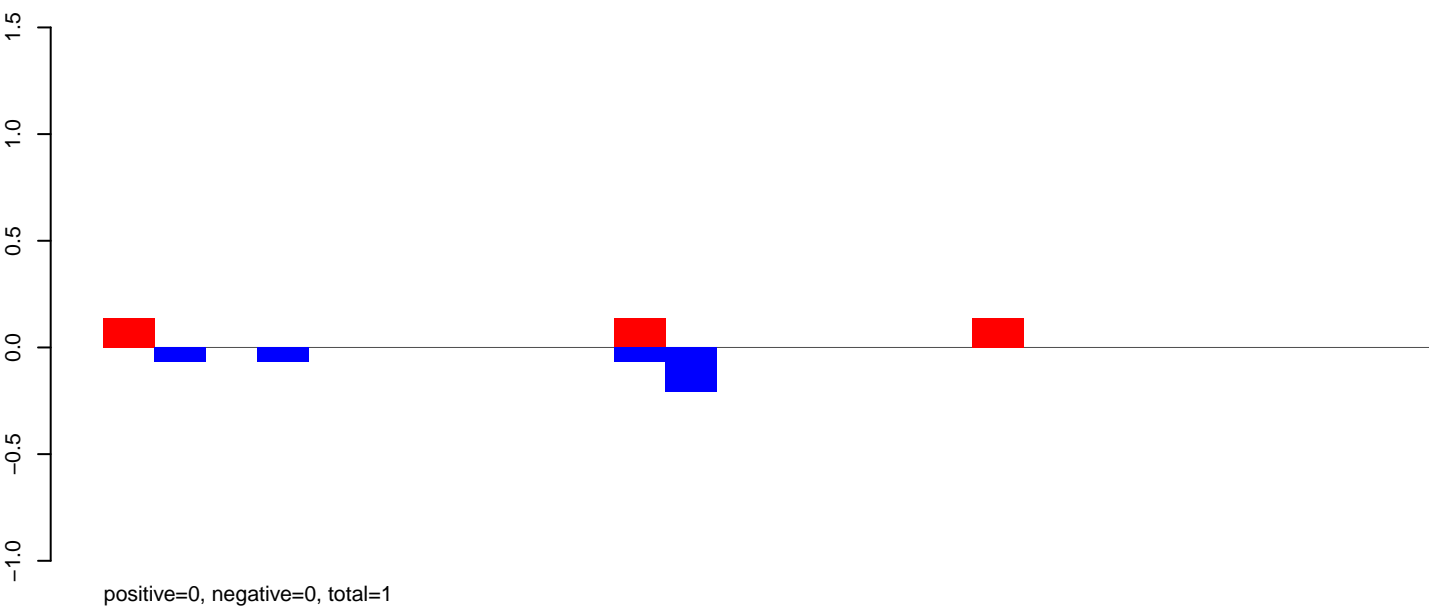
Window size=25, length=307, TE@rnd-4_family-137@DNA_Zator@2180:1-307

50 100 150 200 250 300 350

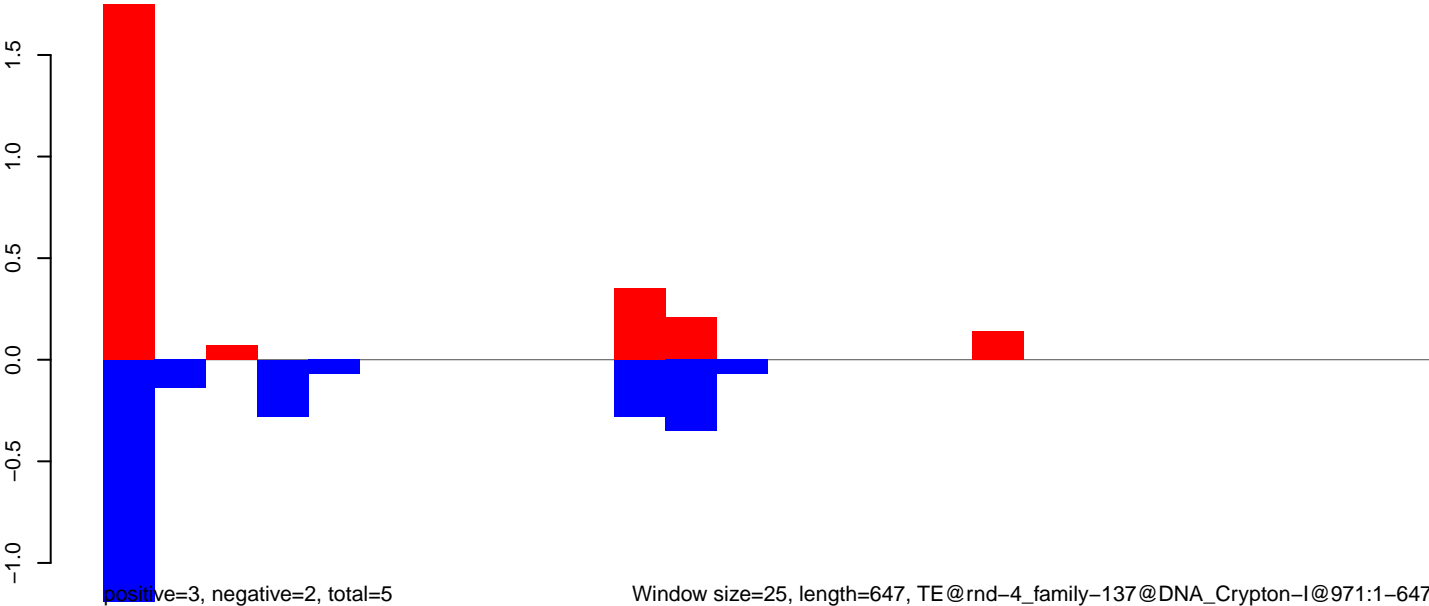
CuQuin_Mcarc_NL.18_23.rep



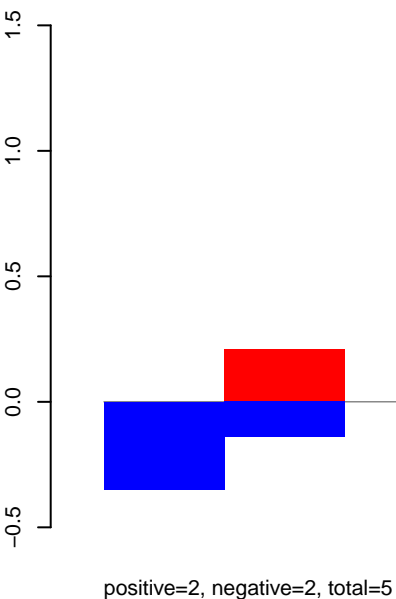
CuQuin_Mcarc_NL.24_35.rep



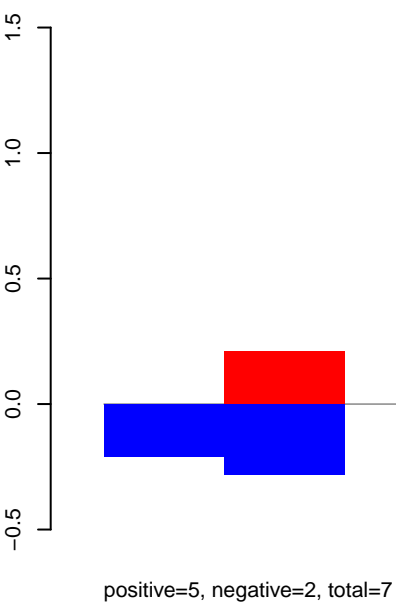
CuQuin_Mcarc_NL.rep



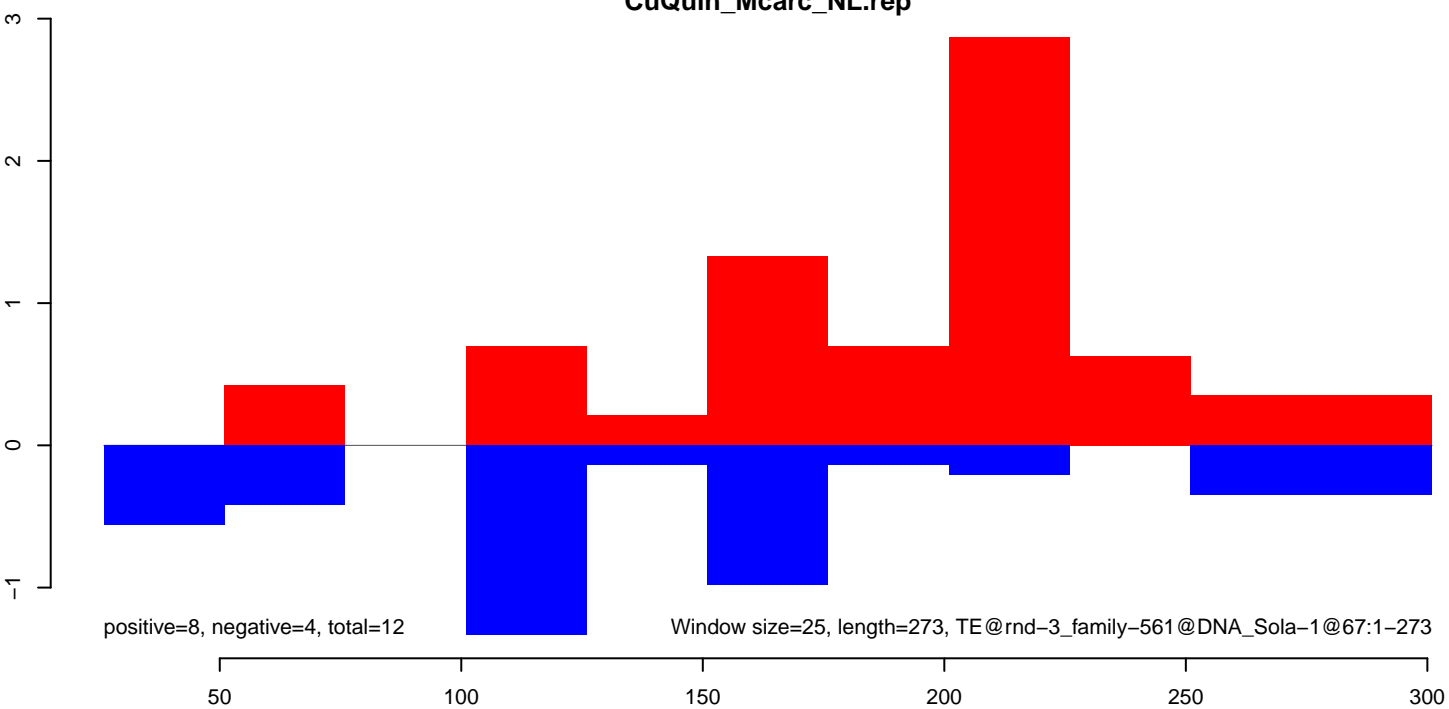
CuQuin_Mcarc_NL.18_23.rep

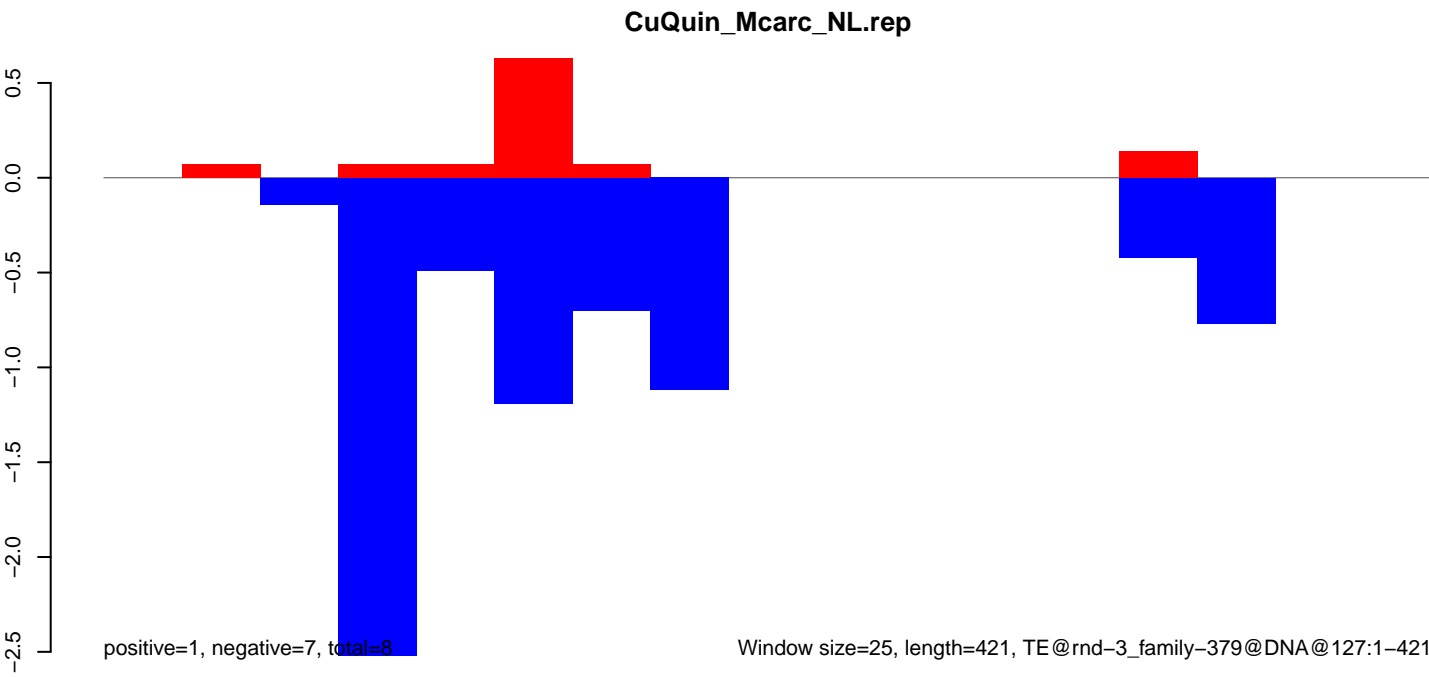
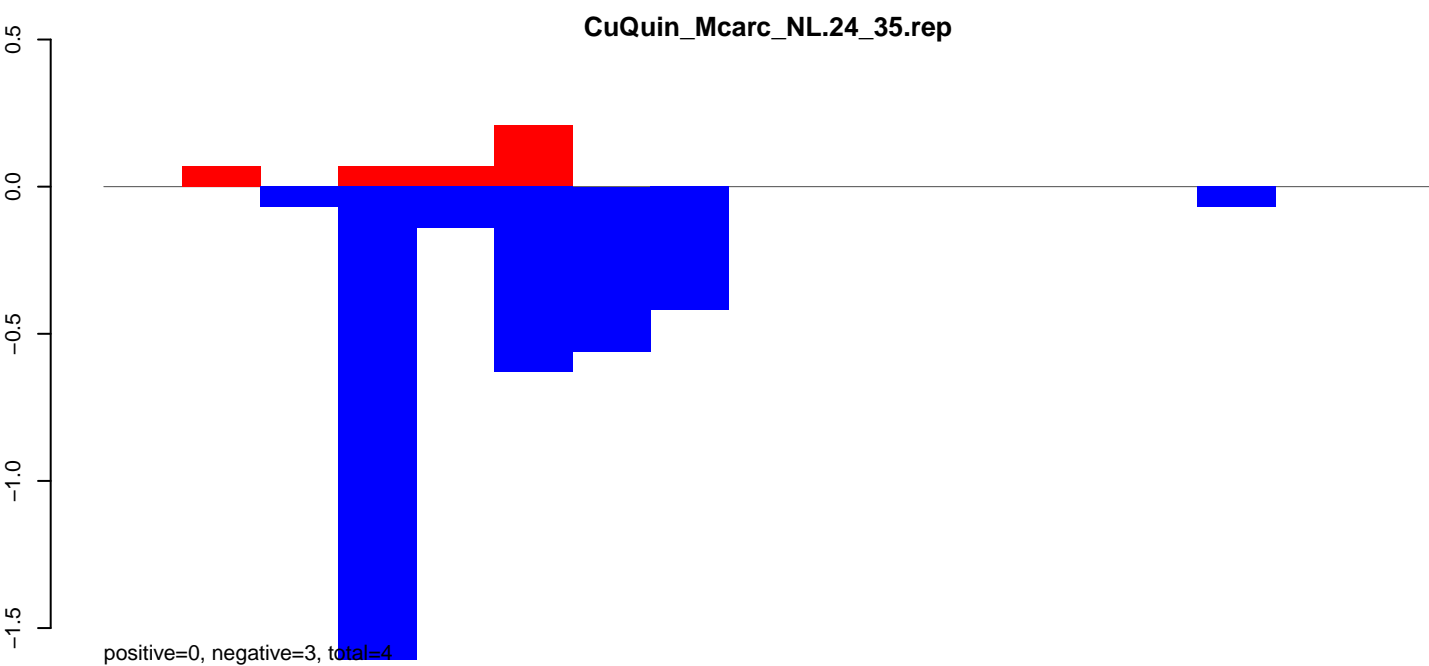
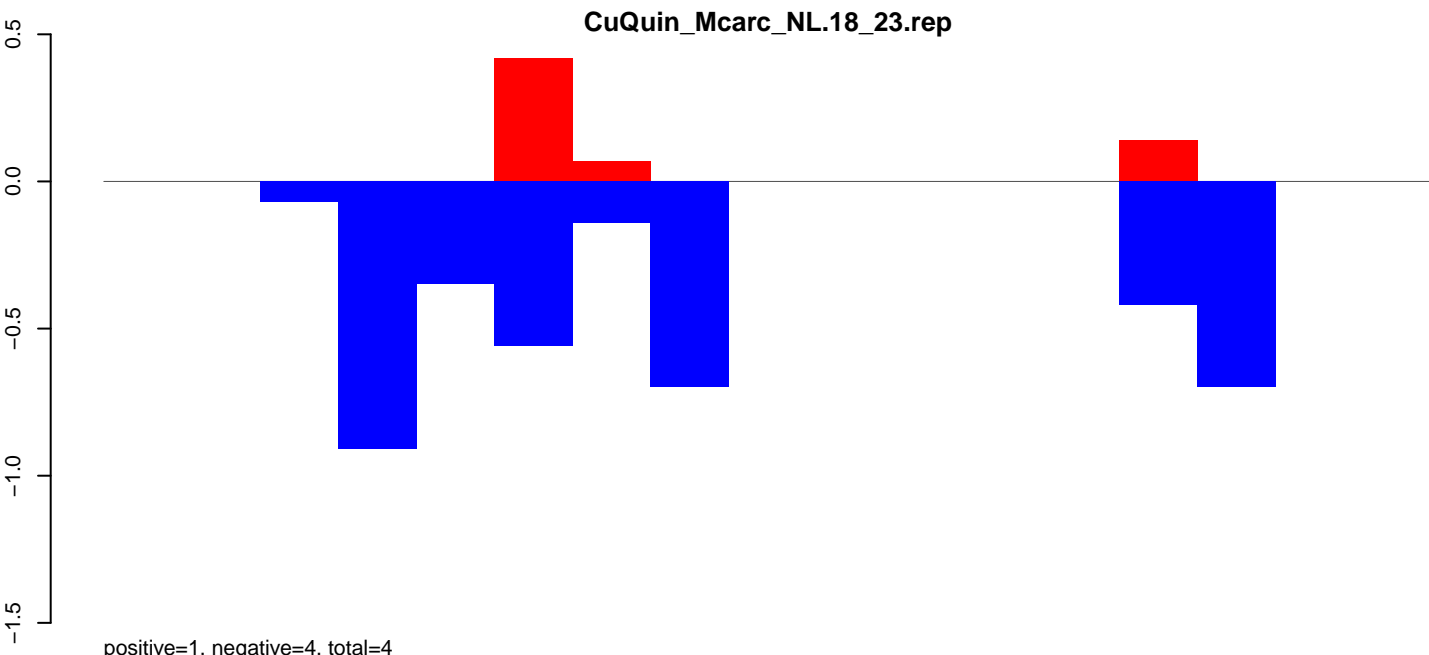


CuQuin_Mcarc_NL.24_35.rep

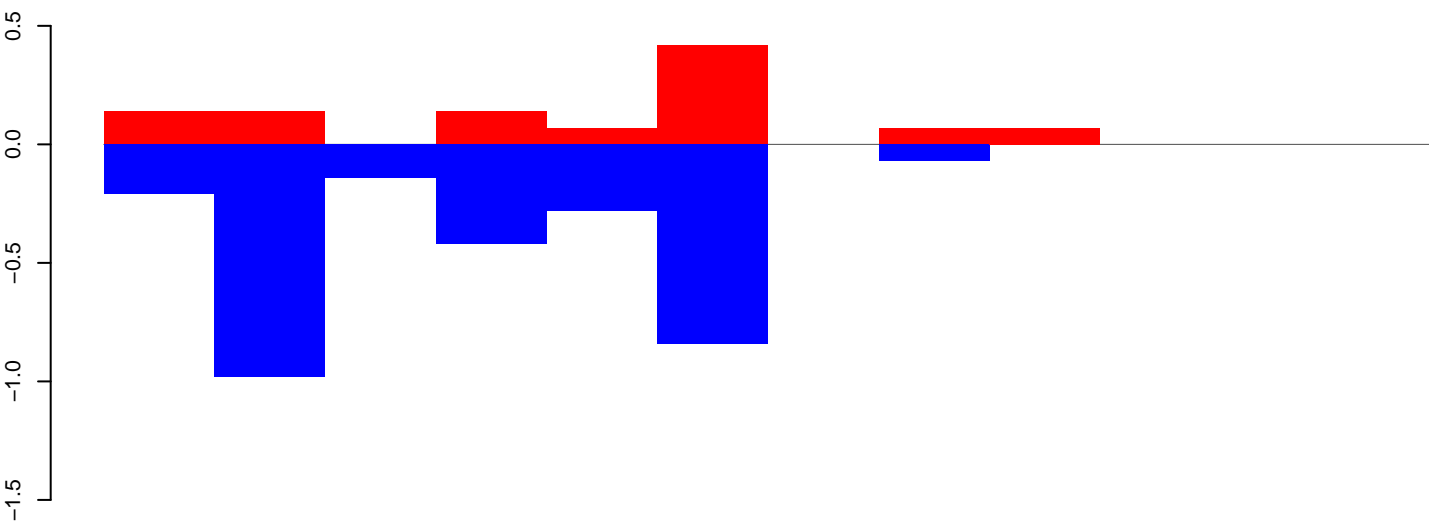


CuQuin_Mcarc_NL.rep



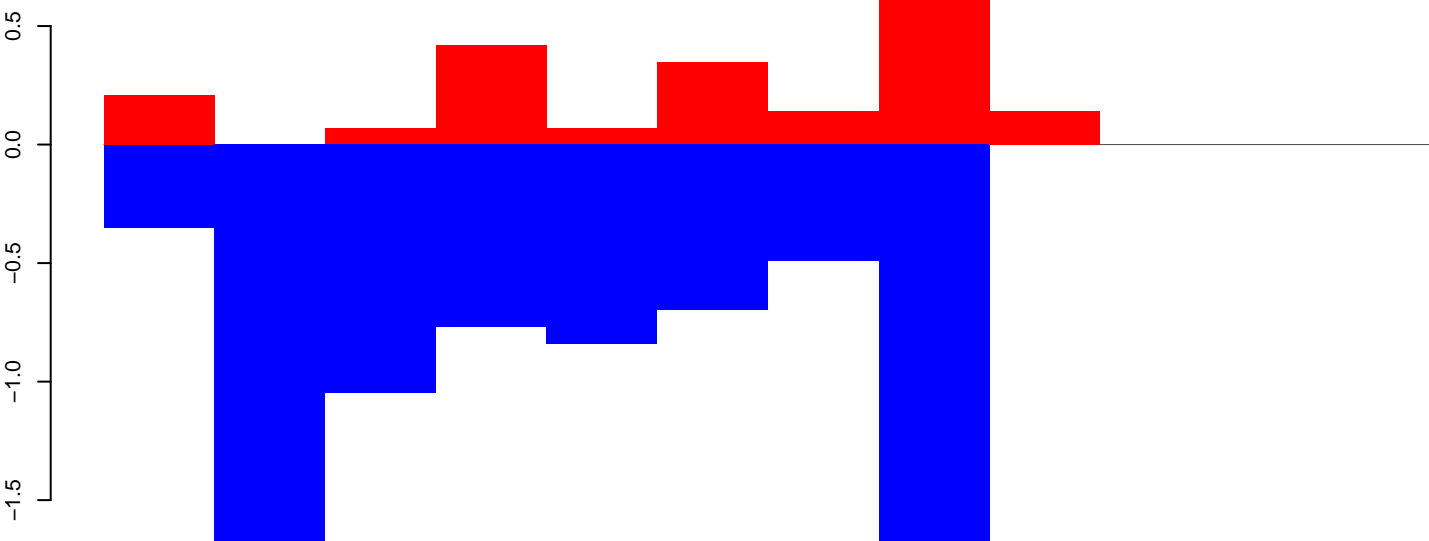


CuQuin_Mcarc_NL.18_23.rep



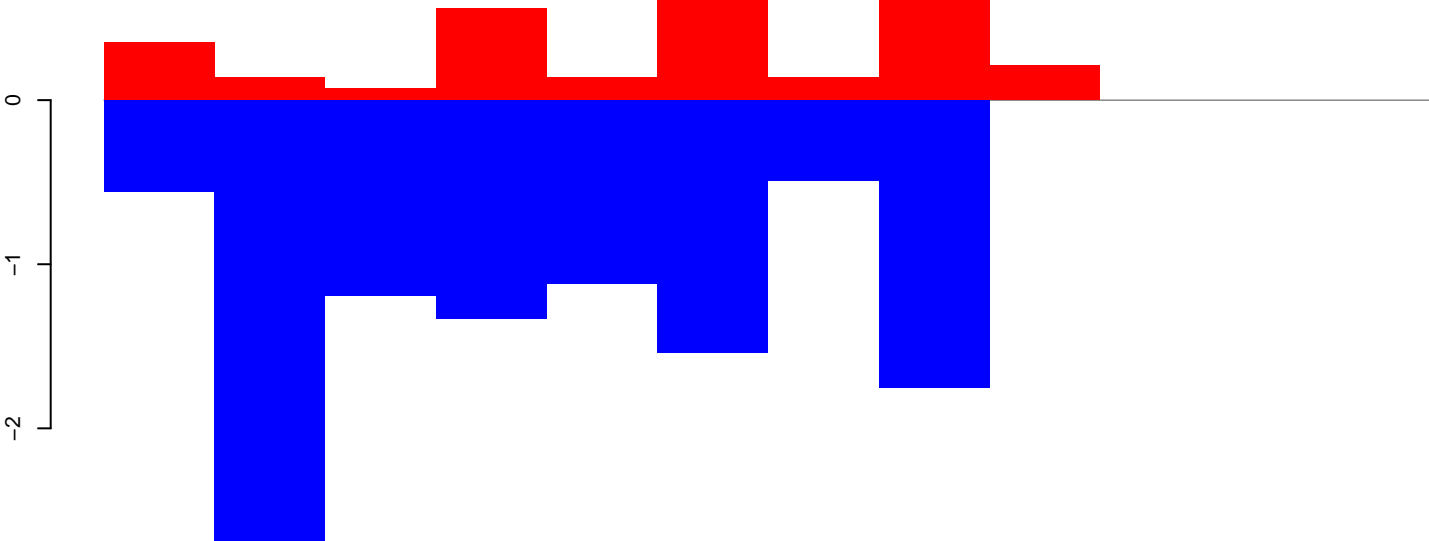
positive=1, negative=3, total=4

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=8, total=10

CuQuin_Mcarc_NL.rep

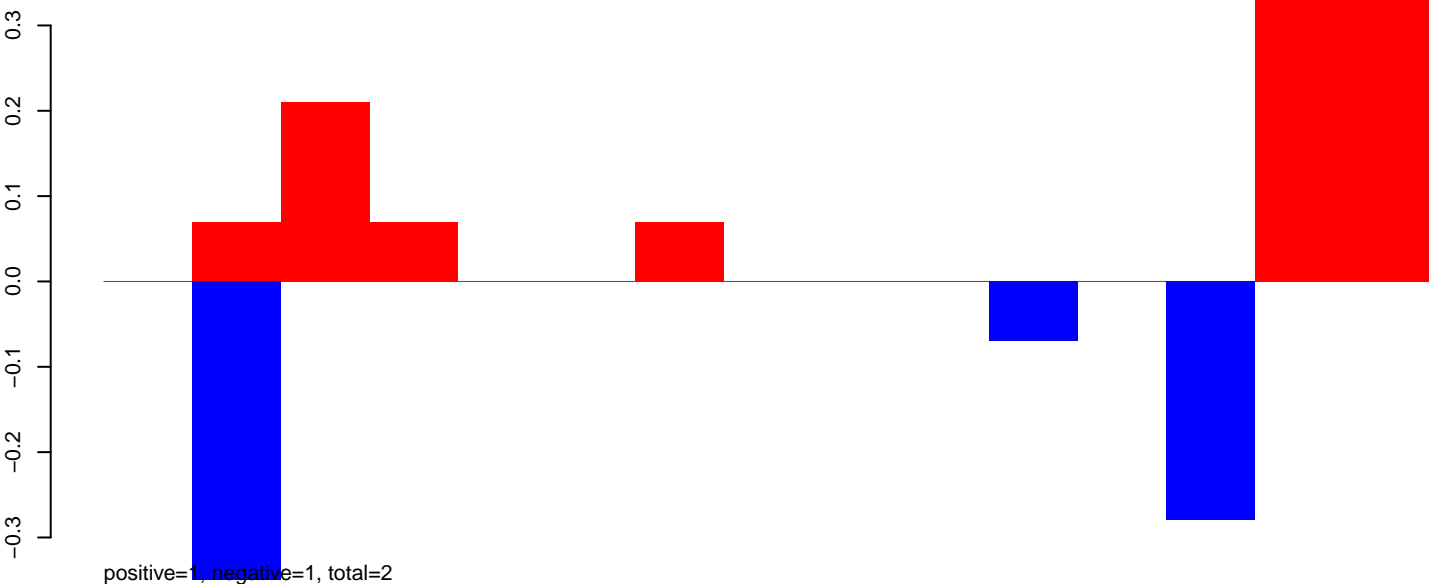


positive=3, negative=11, total=14

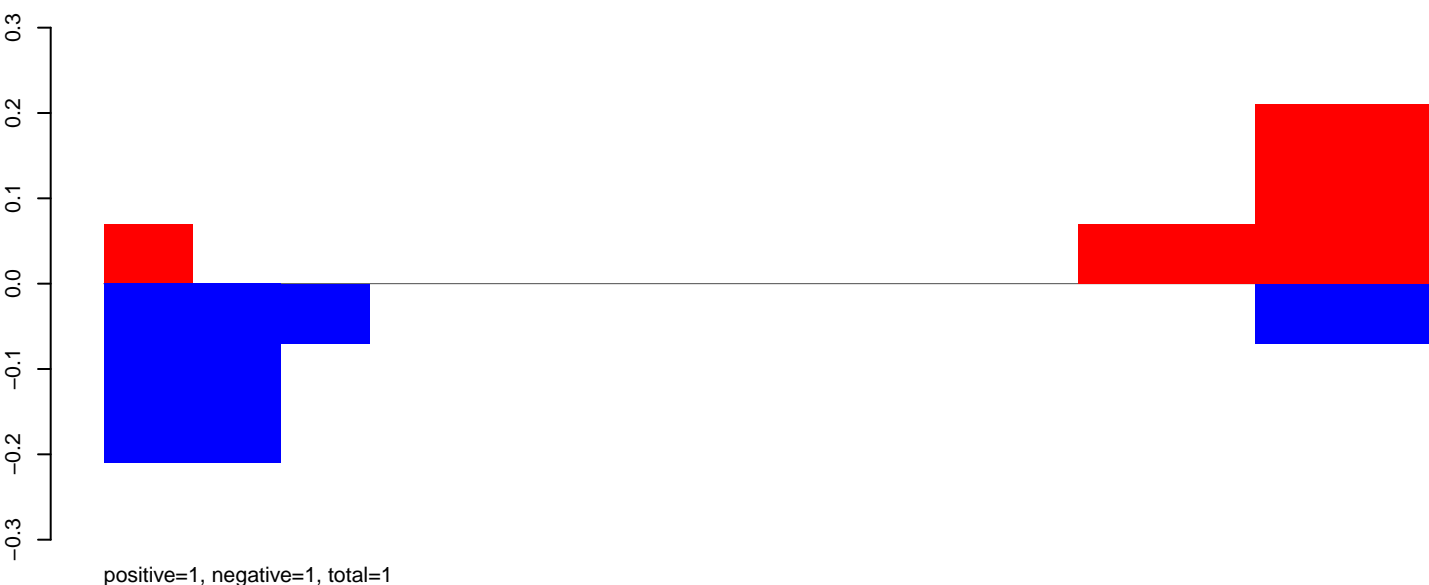
Window size=25, length=279, TE@rnd-3_family-36@Unknown@3340:1-279

50 100 150 200 250 300

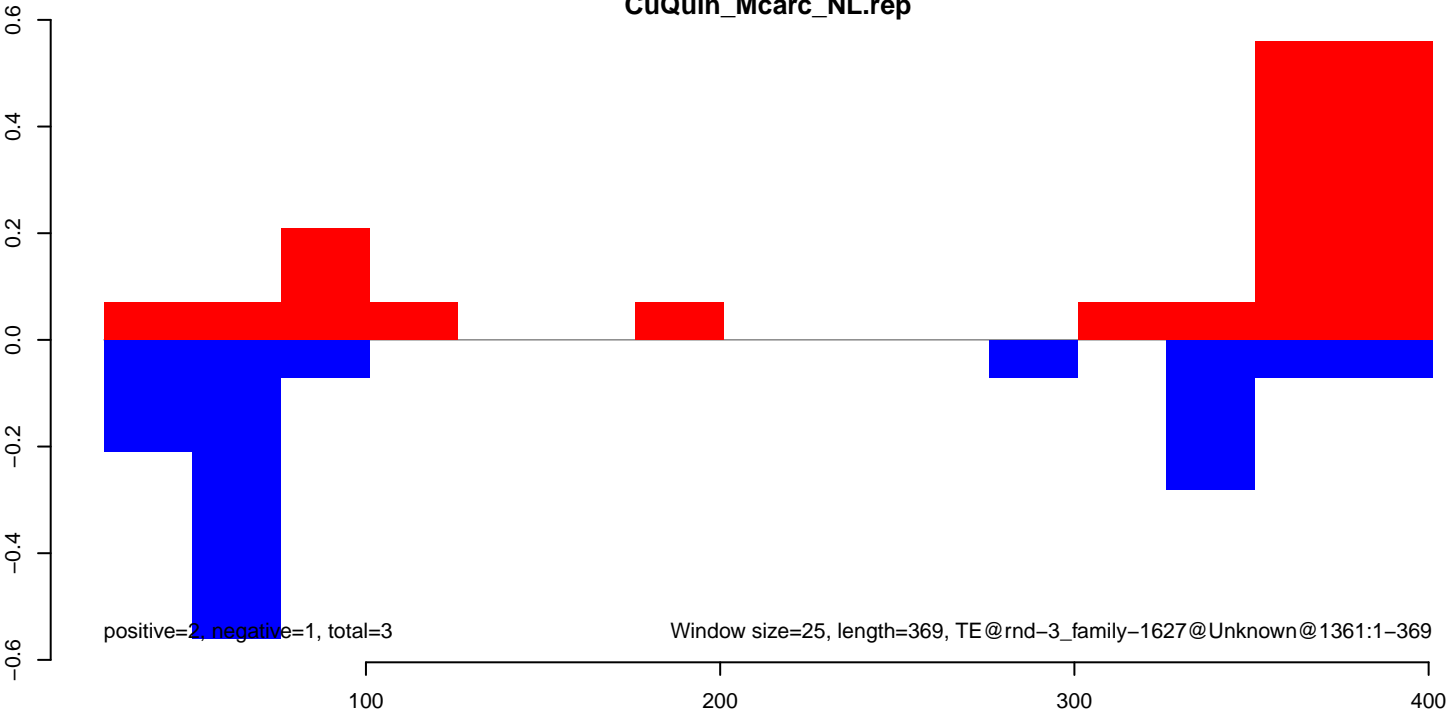
CuQuin_Mcarc_NL.18_23.rep



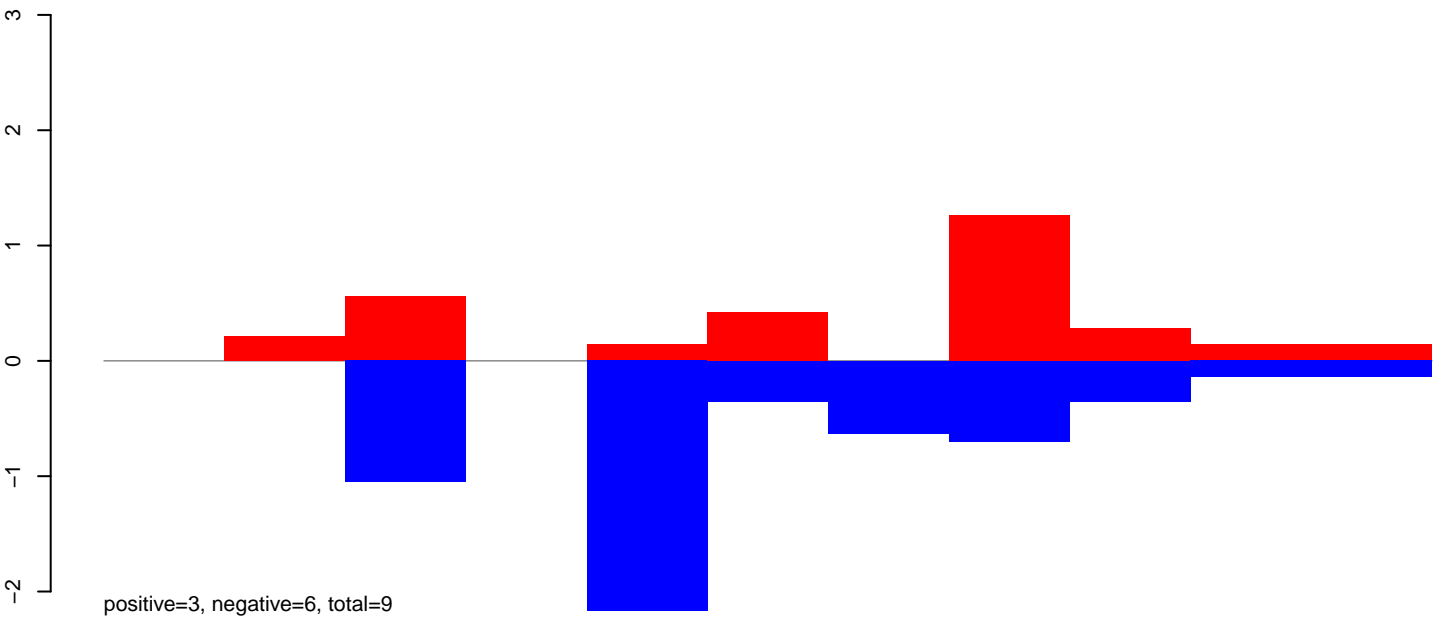
CuQuin_Mcarc_NL.24_35.rep



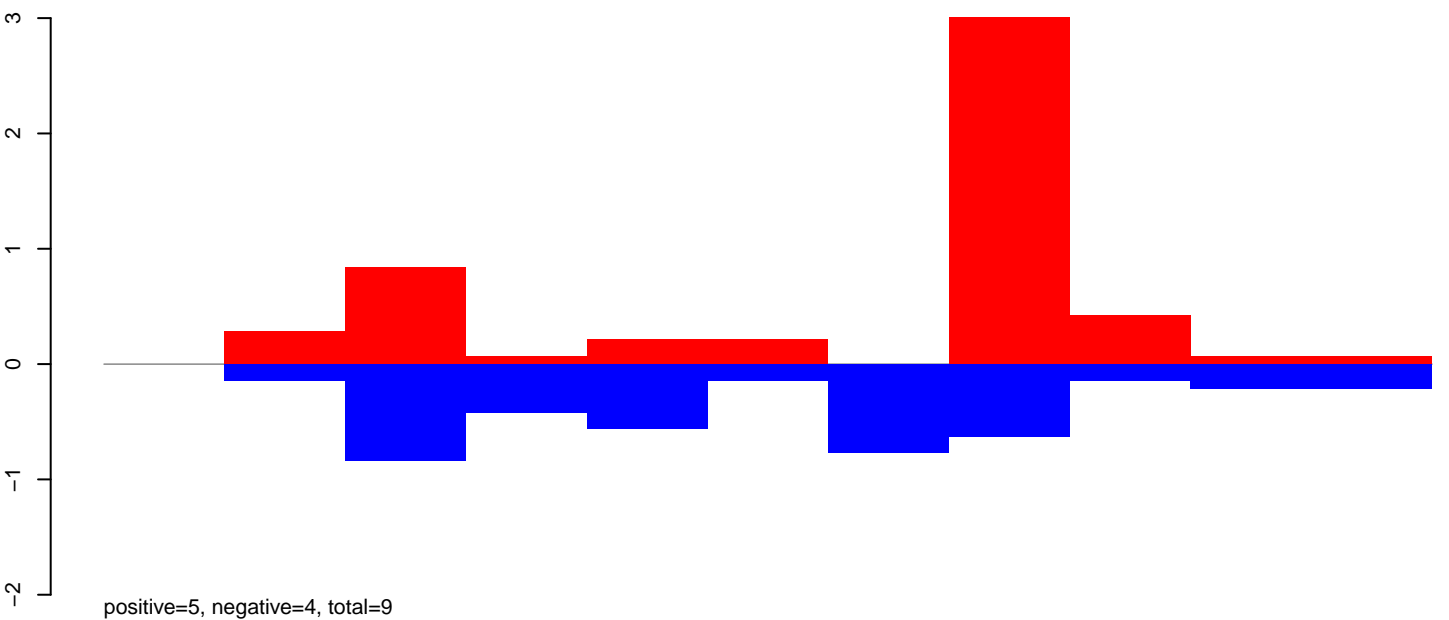
CuQuin_Mcarc_NL.rep



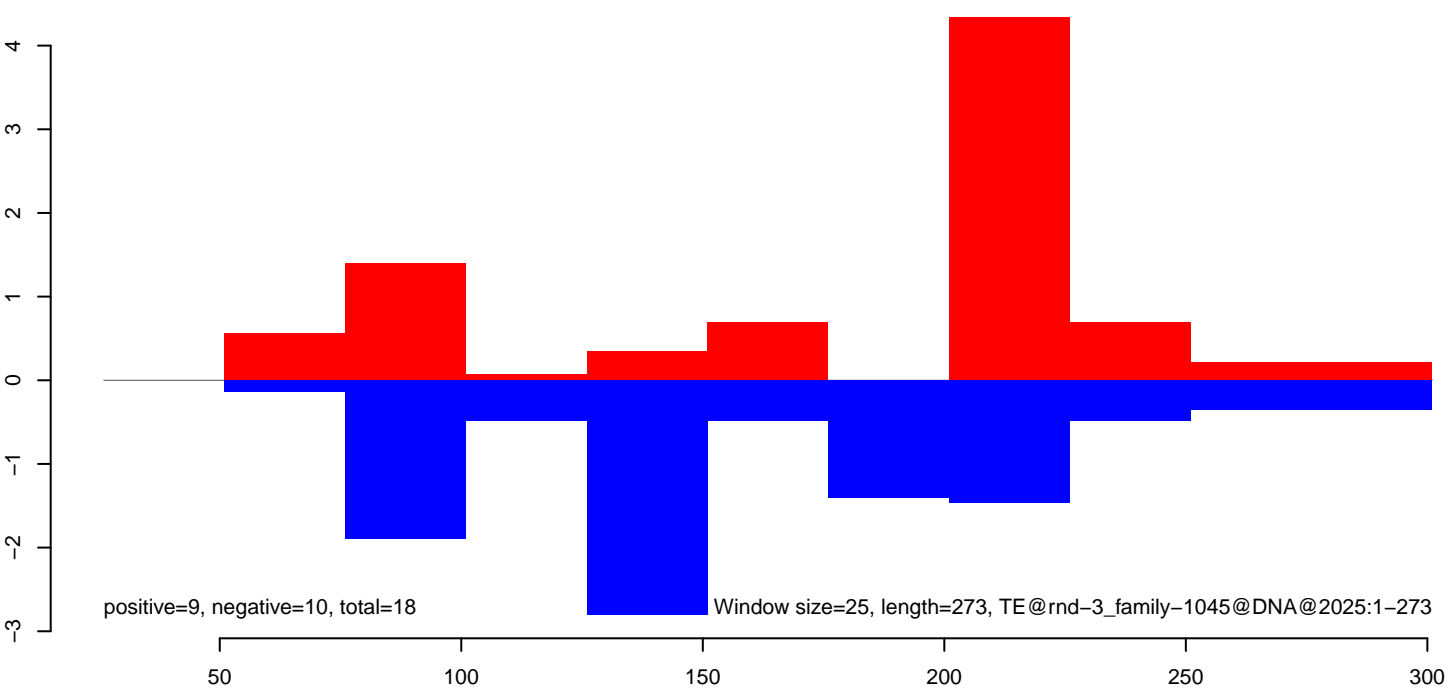
CuQuin_Mcarc_NL.18_23.rep

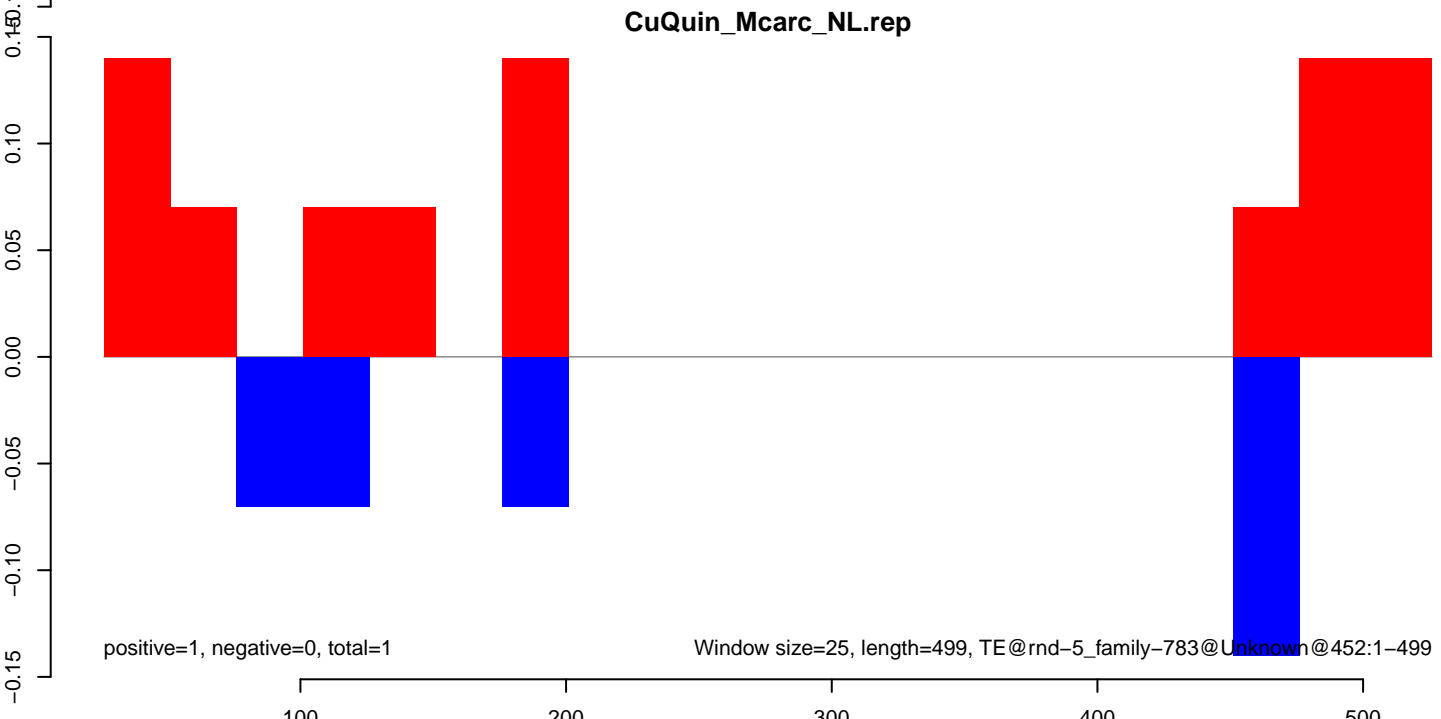
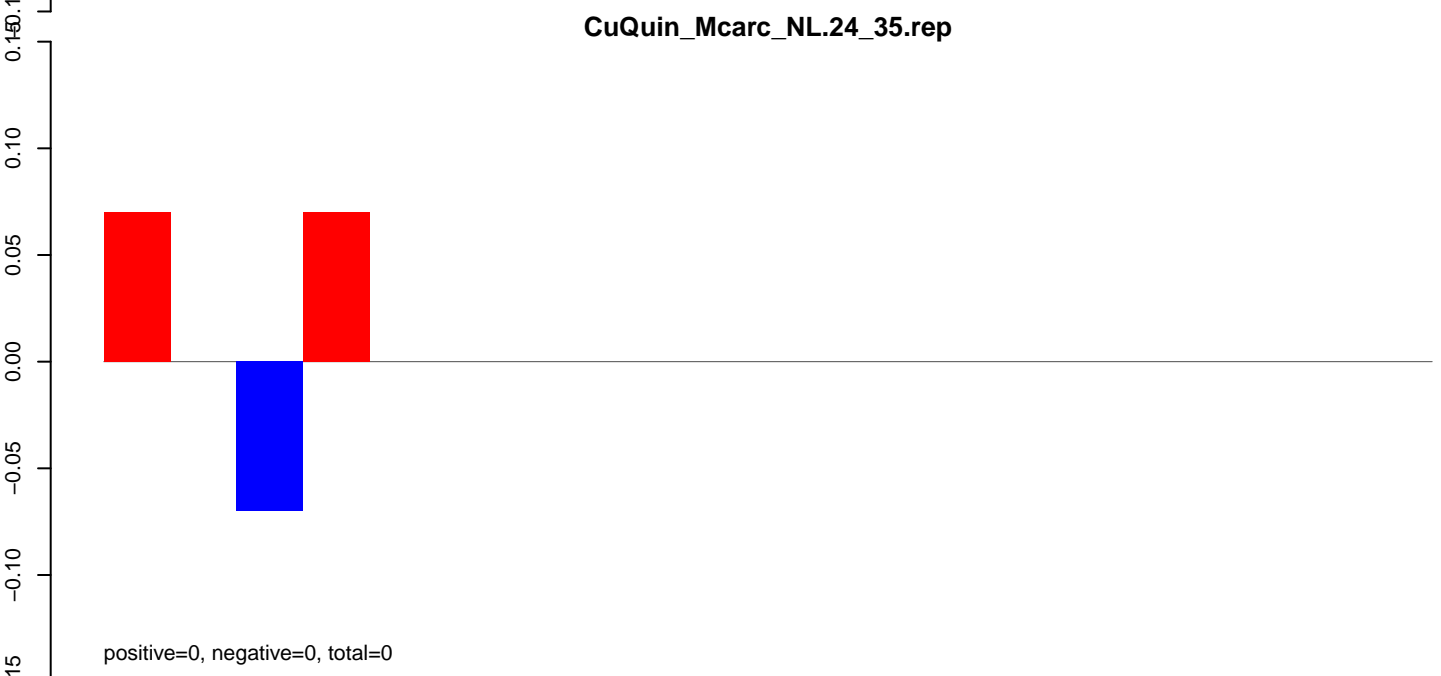
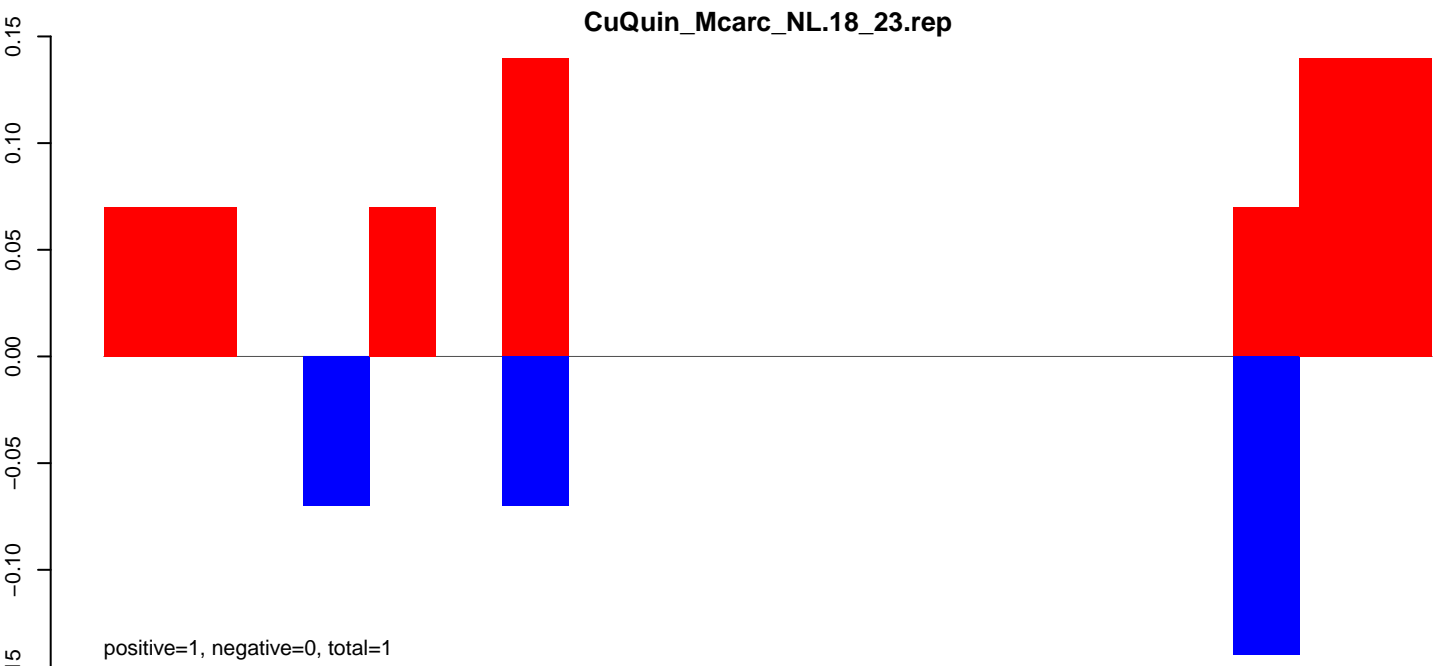


CuQuin_Mcarc_NL.24_35.rep

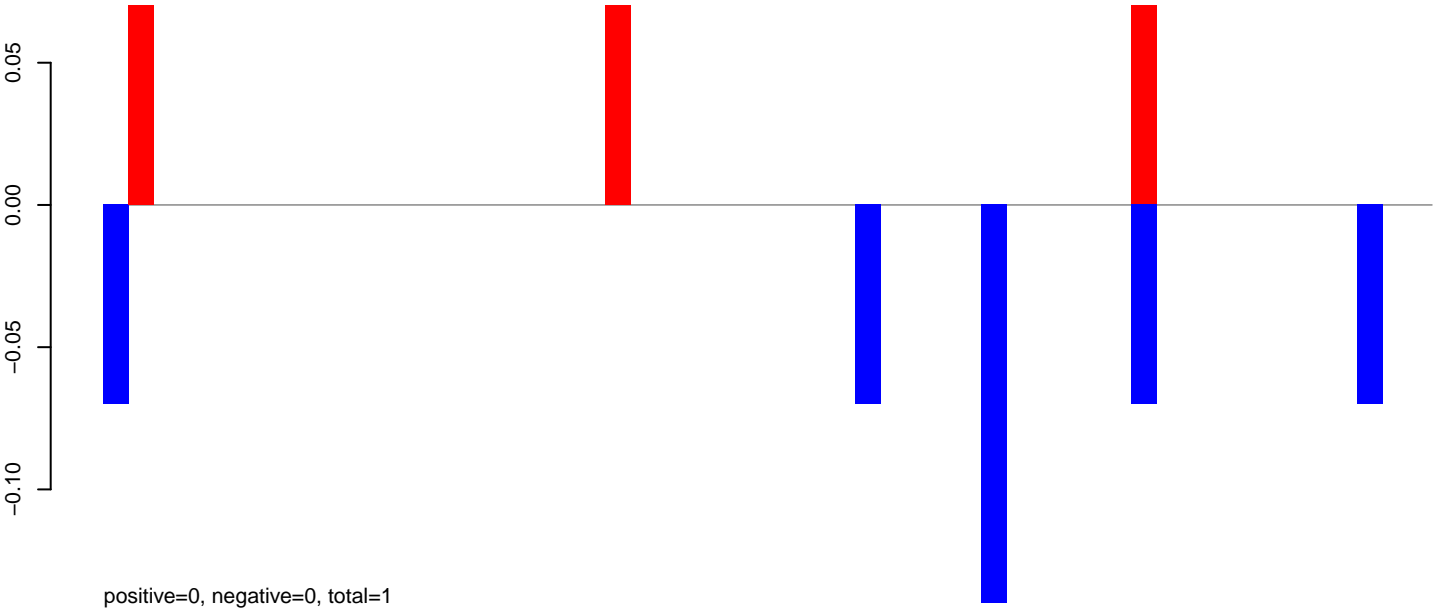


CuQuin_Mcarc_NL.rep

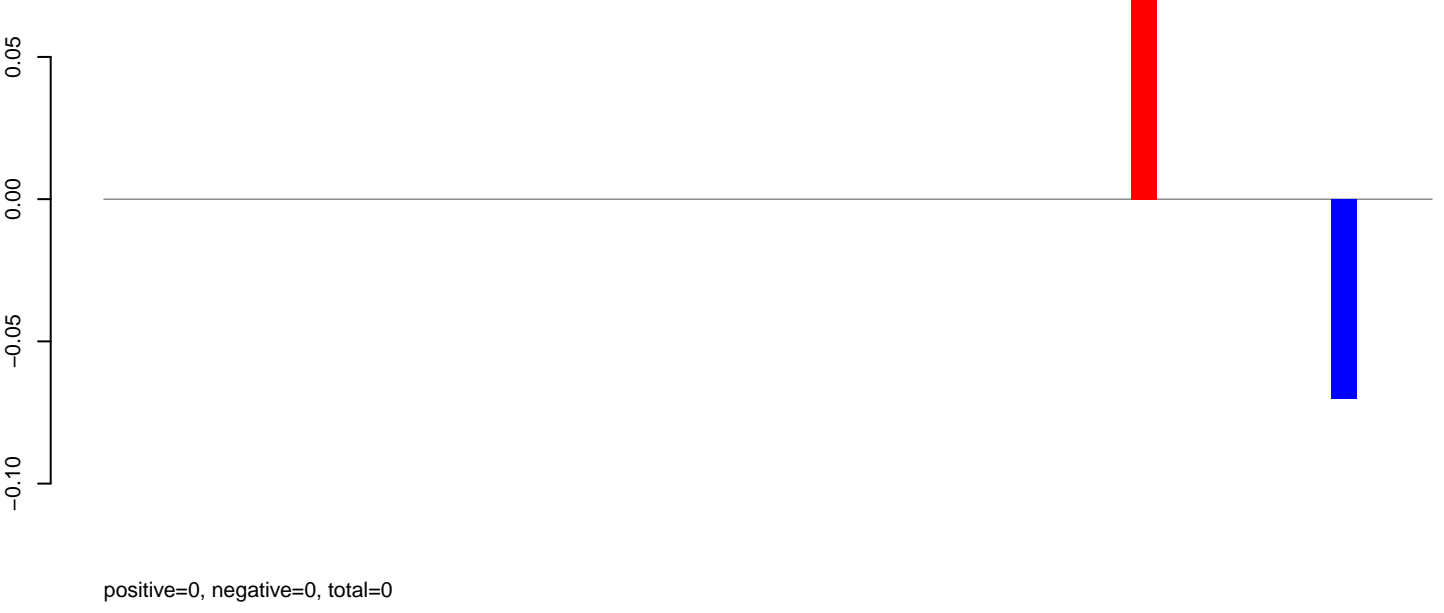




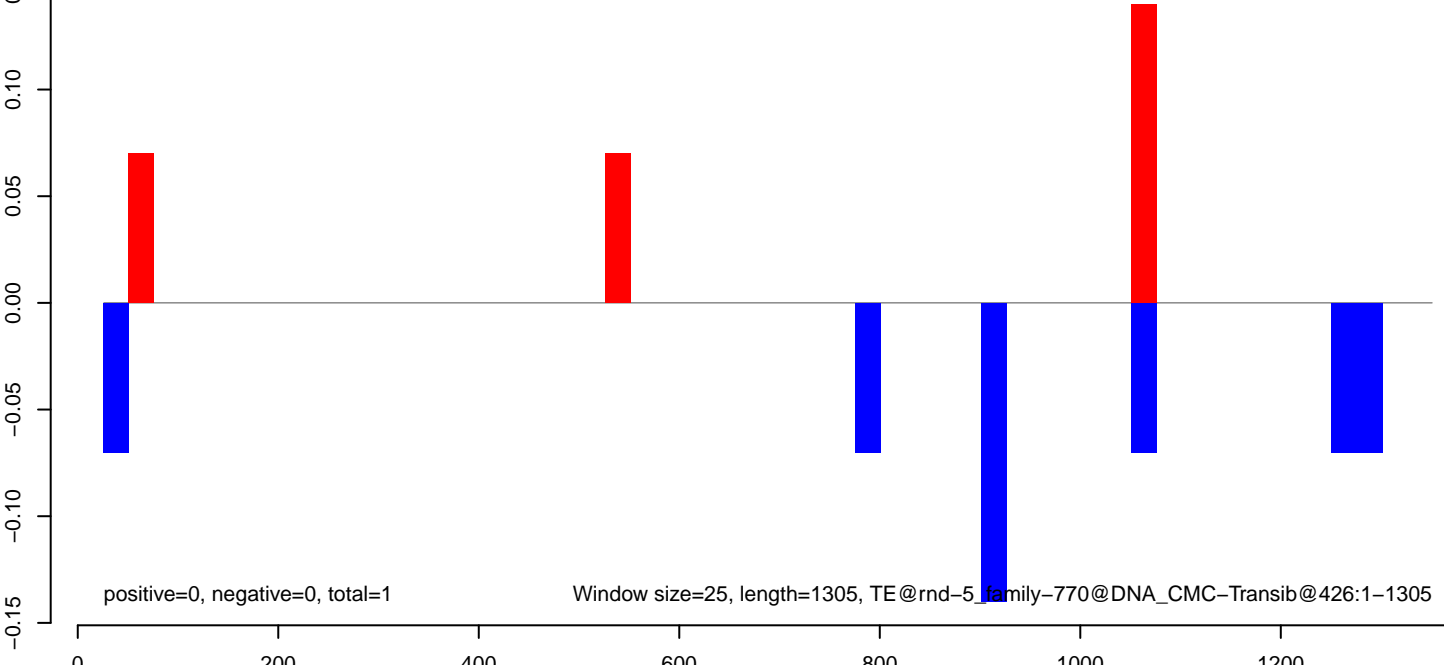
CuQuin_Mcarc_NL.18_23.rep



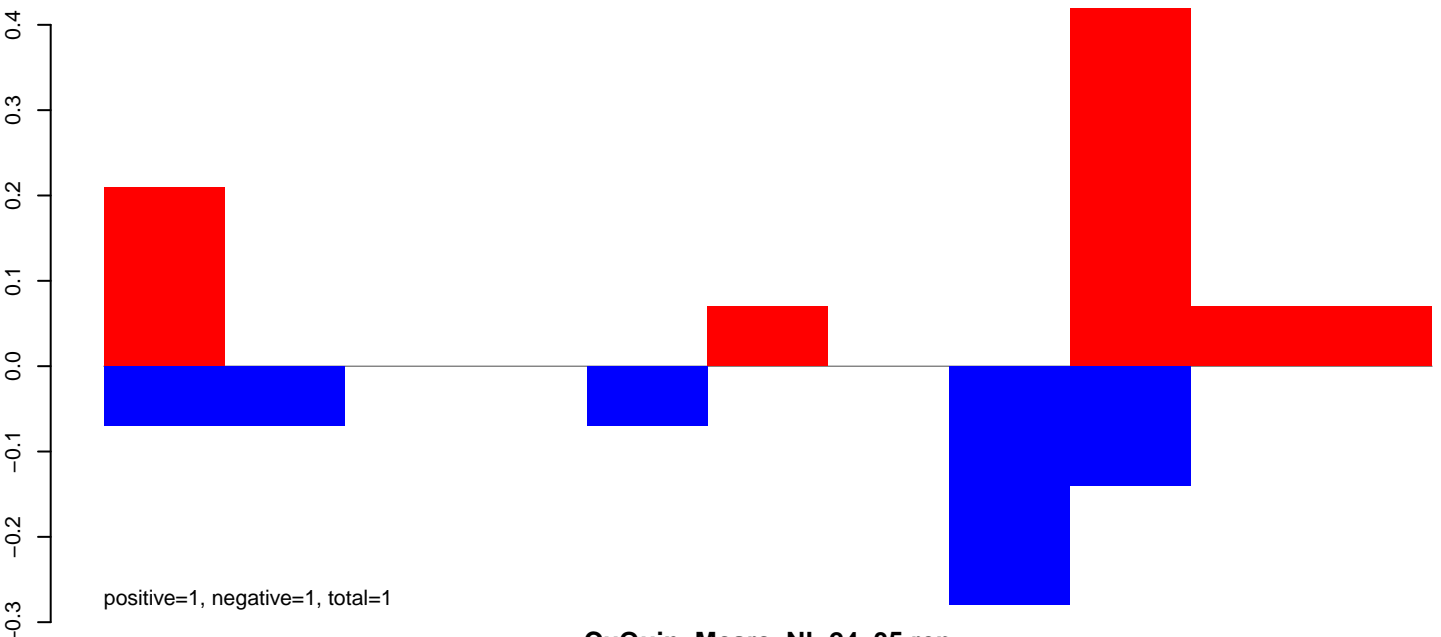
CuQuin_Mcarc_NL.24_35.rep



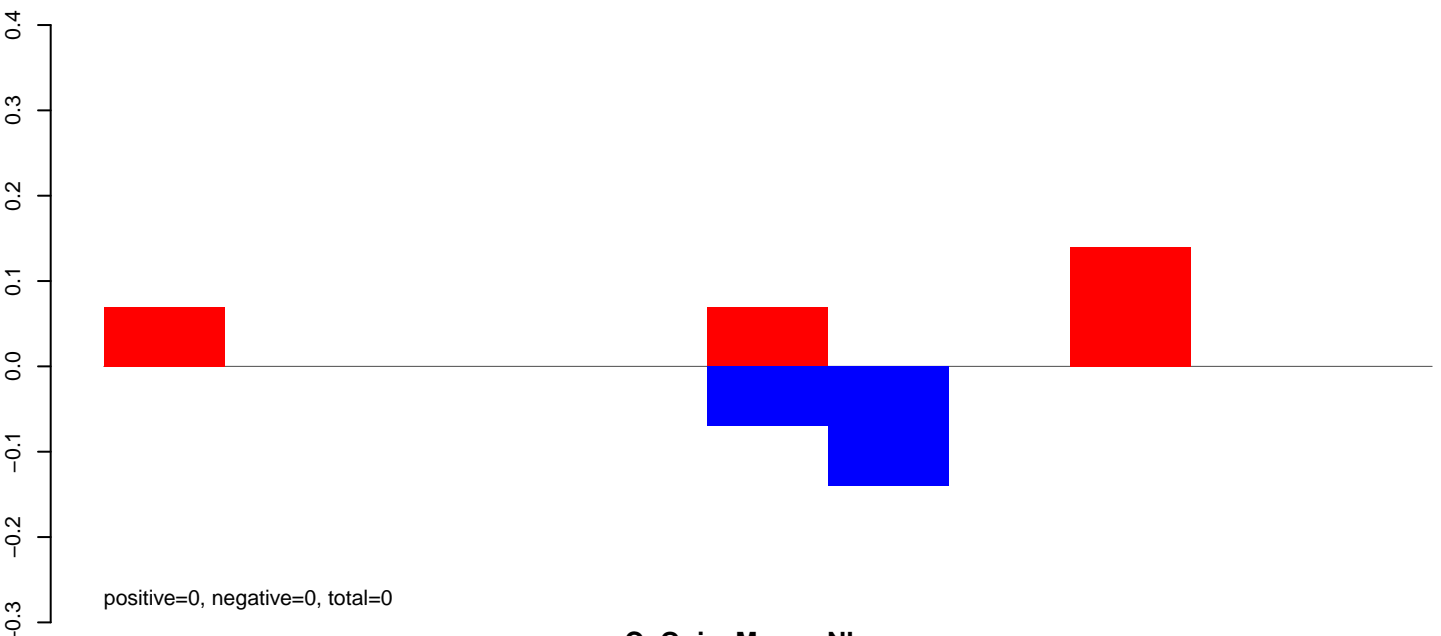
CuQuin_Mcarc_NL.rep



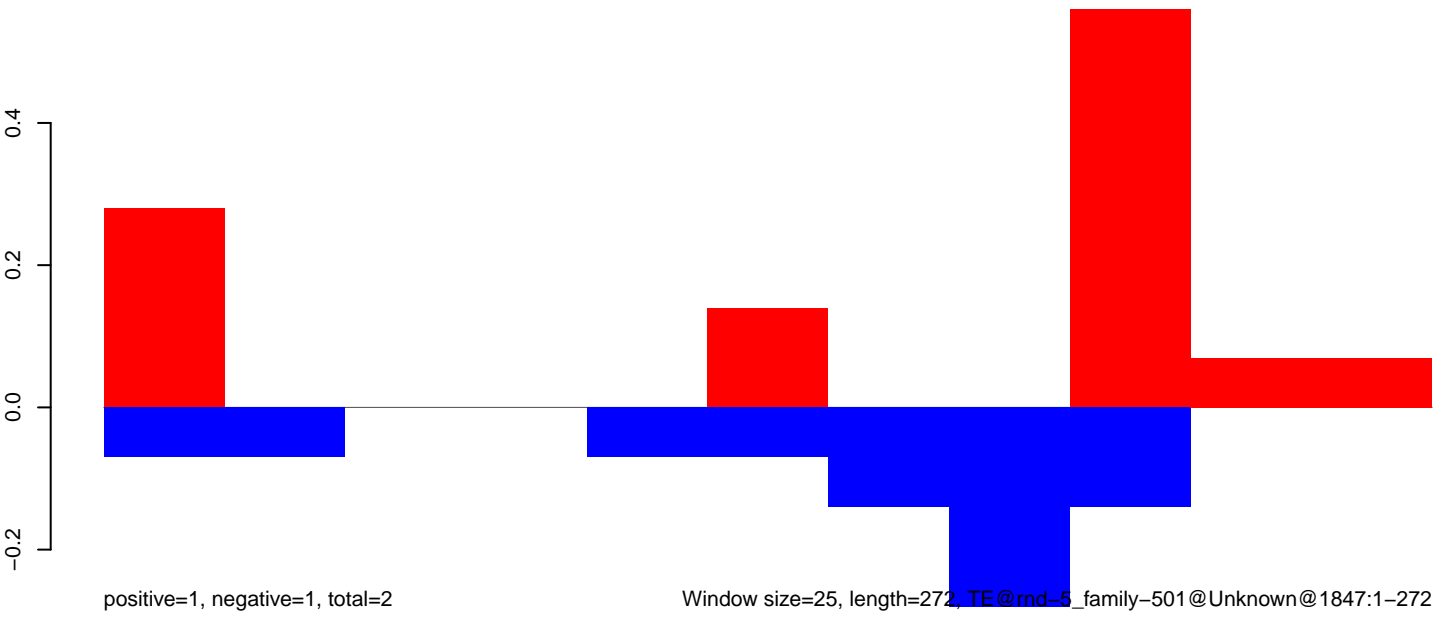
CuQuin_Mcarc_NL.18_23.rep



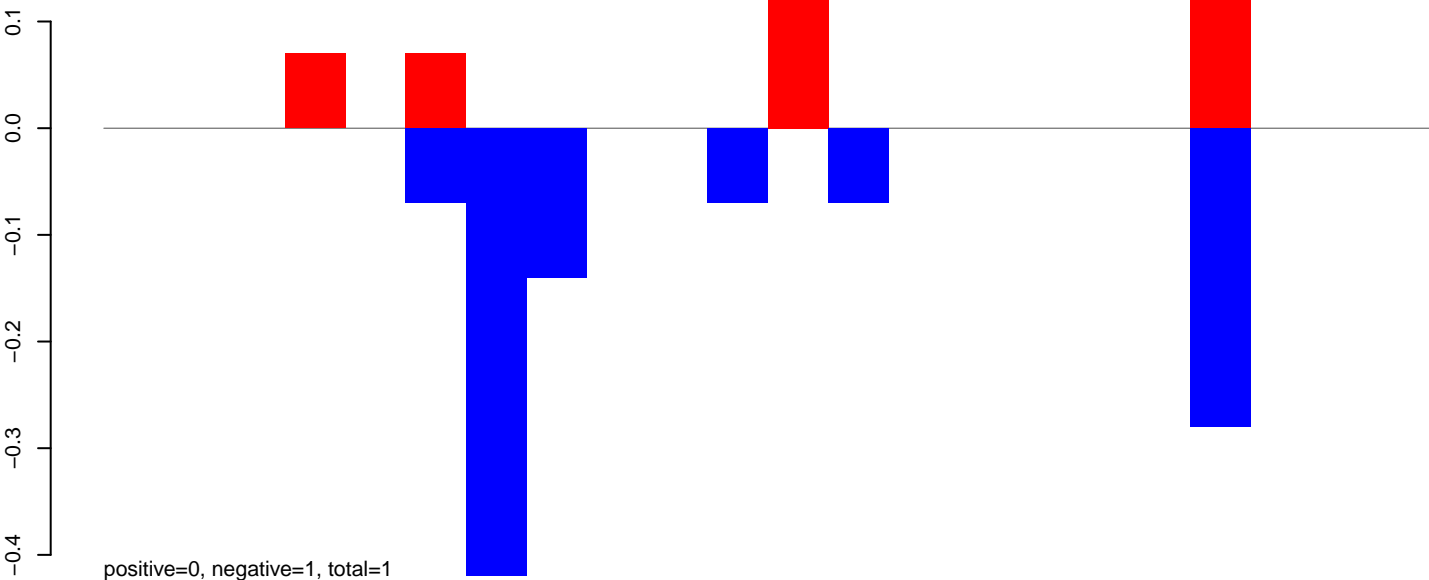
CuQuin_Mcarc_NL.24_35.rep



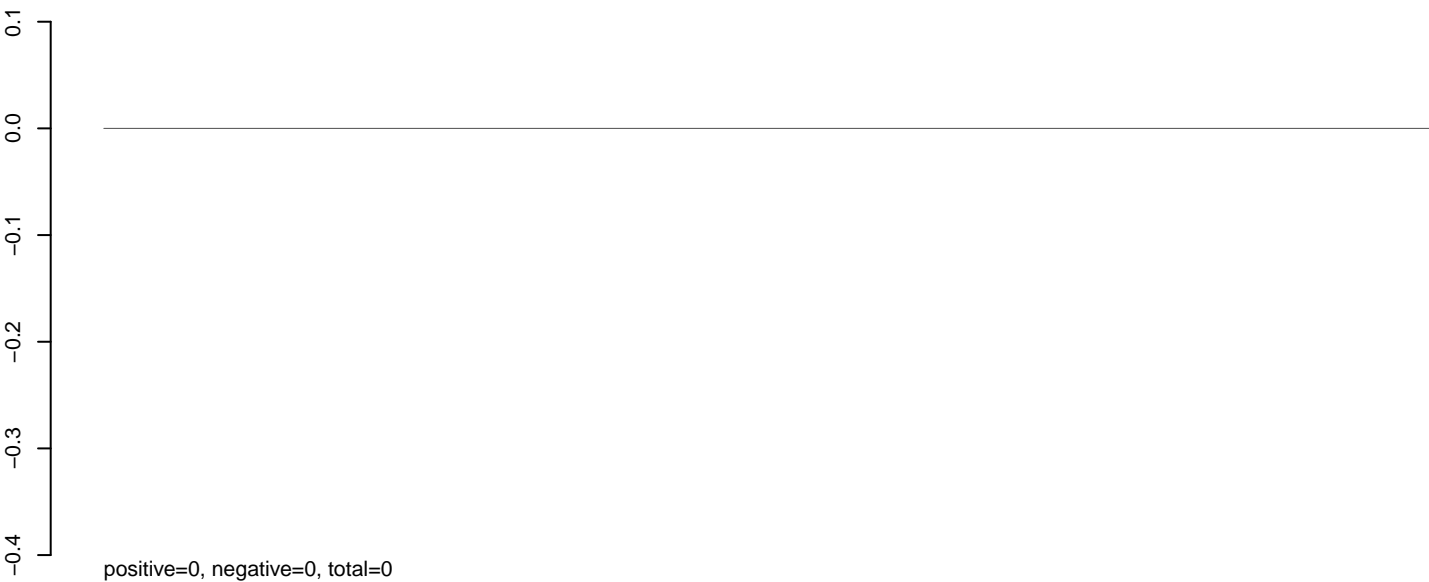
CuQuin_Mcarc_NL.rep



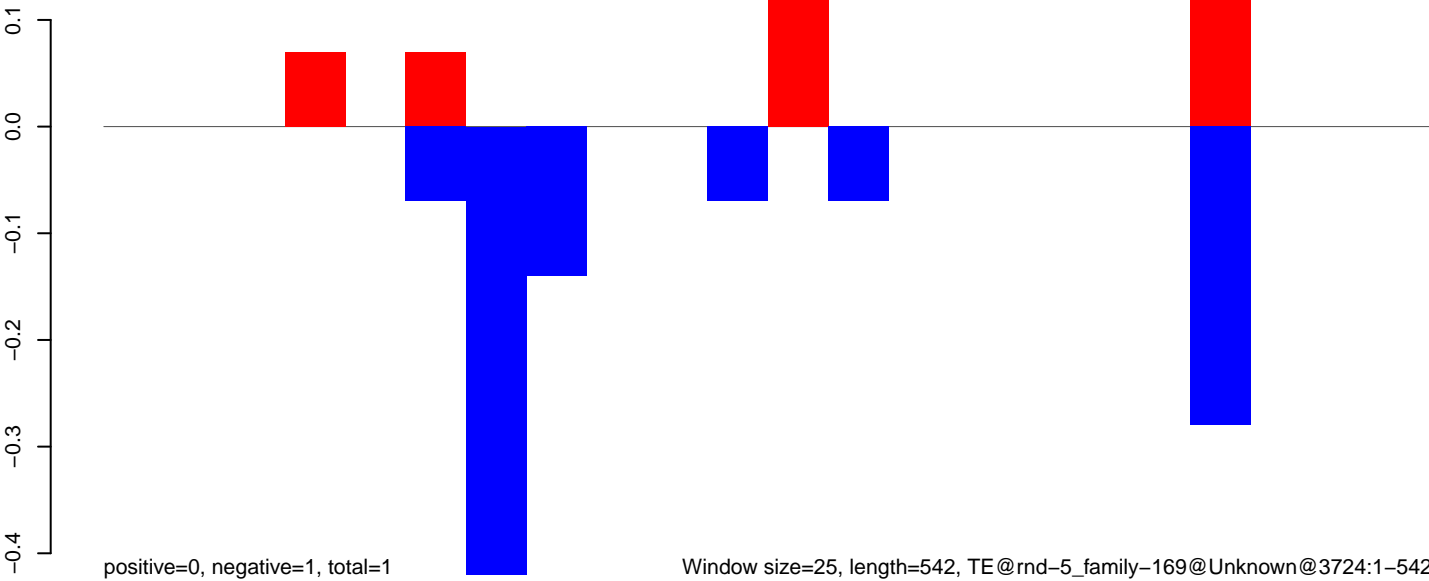
CuQuin_Mcarc_NL.18_23.rep



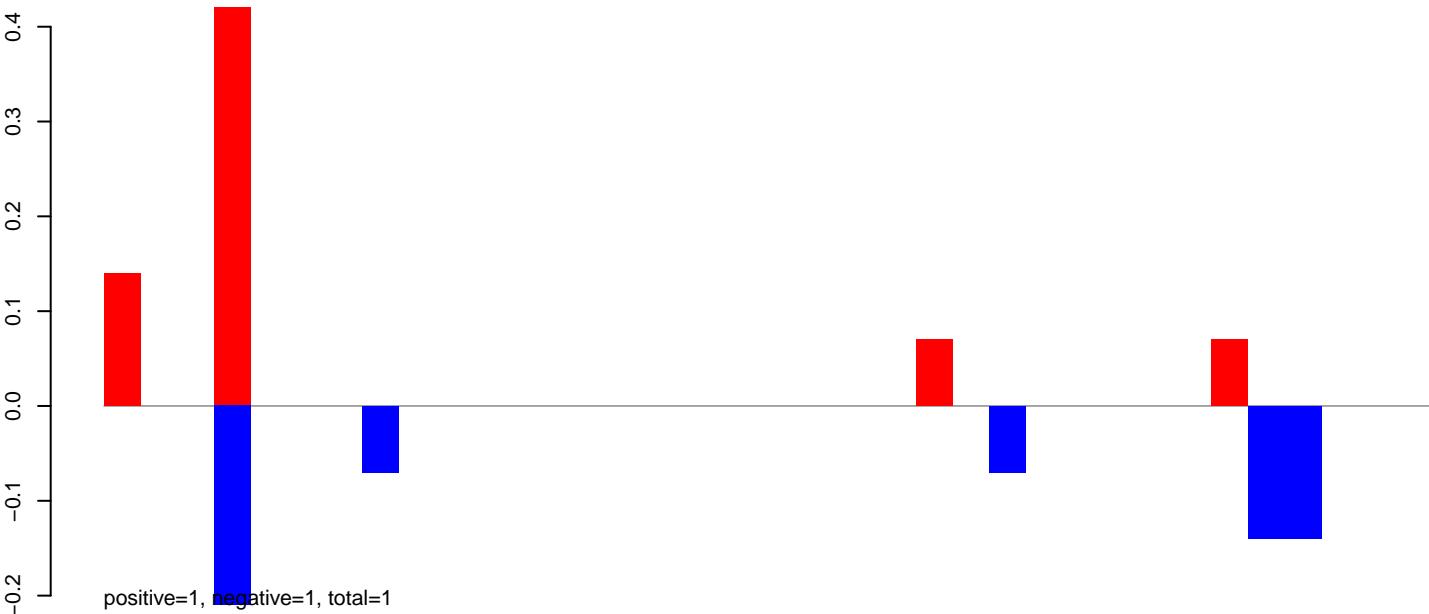
CuQuin_Mcarc_NL.24_35.rep



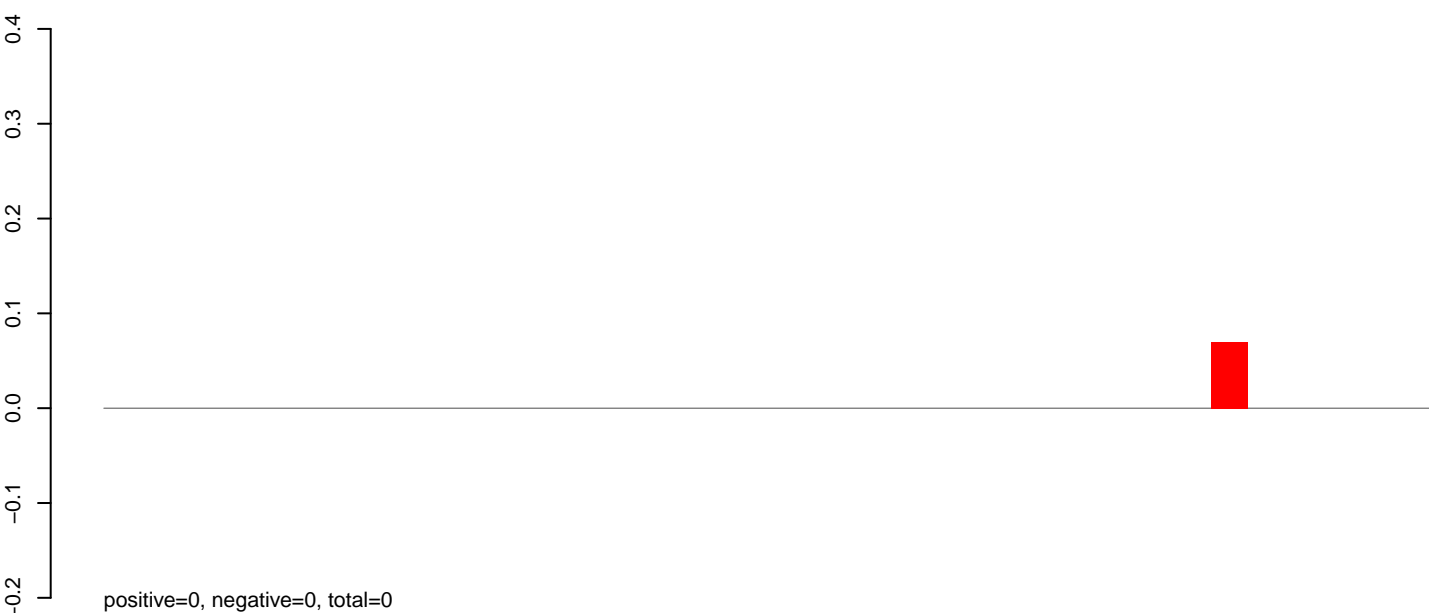
CuQuin_Mcarc_NL.rep



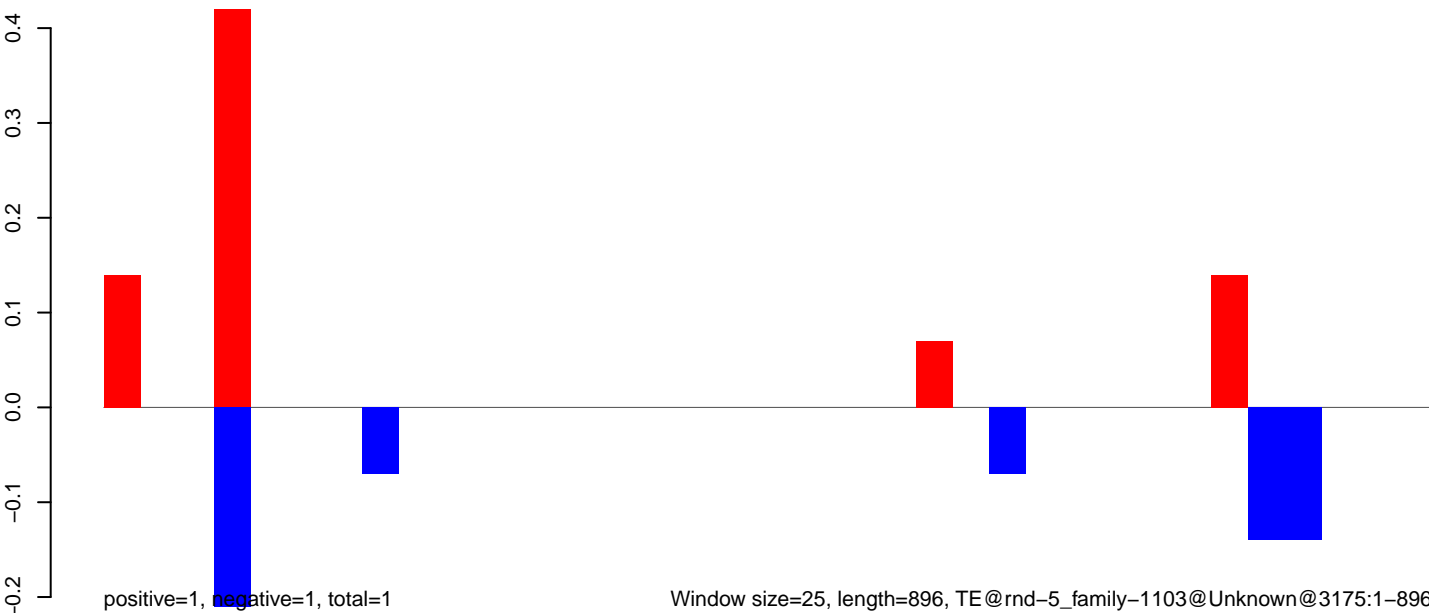
CuQuin_Mcarc_NL.18_23.rep



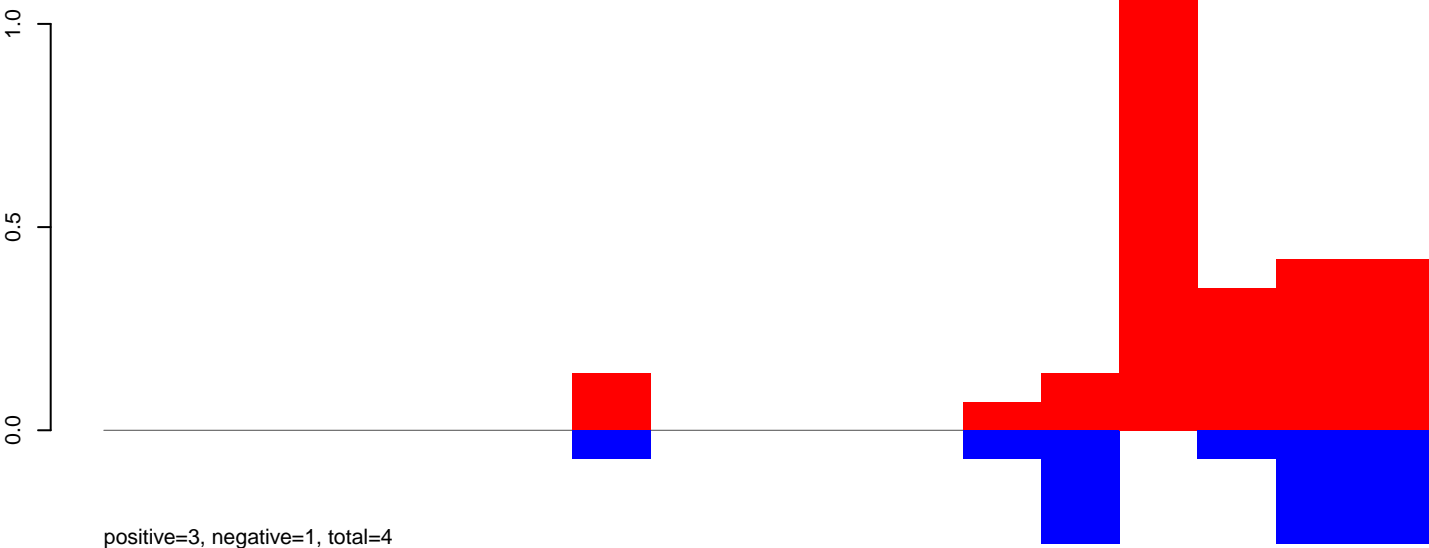
CuQuin_Mcarc_NL.24_35.rep



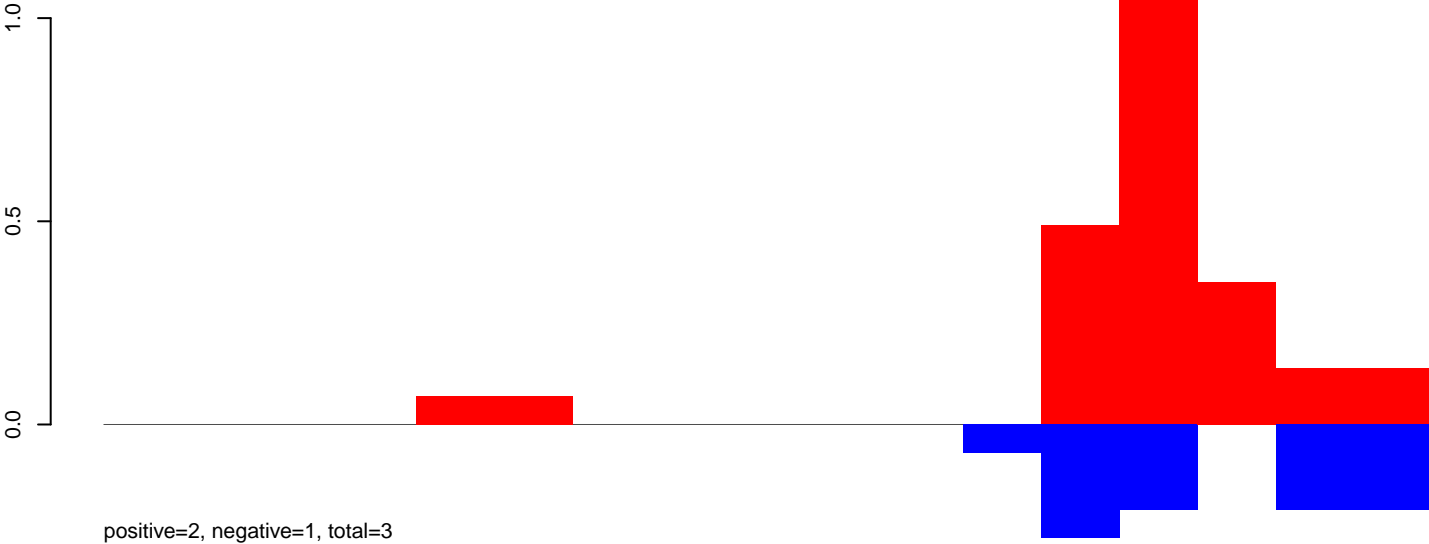
CuQuin_Mcarc_NL.rep



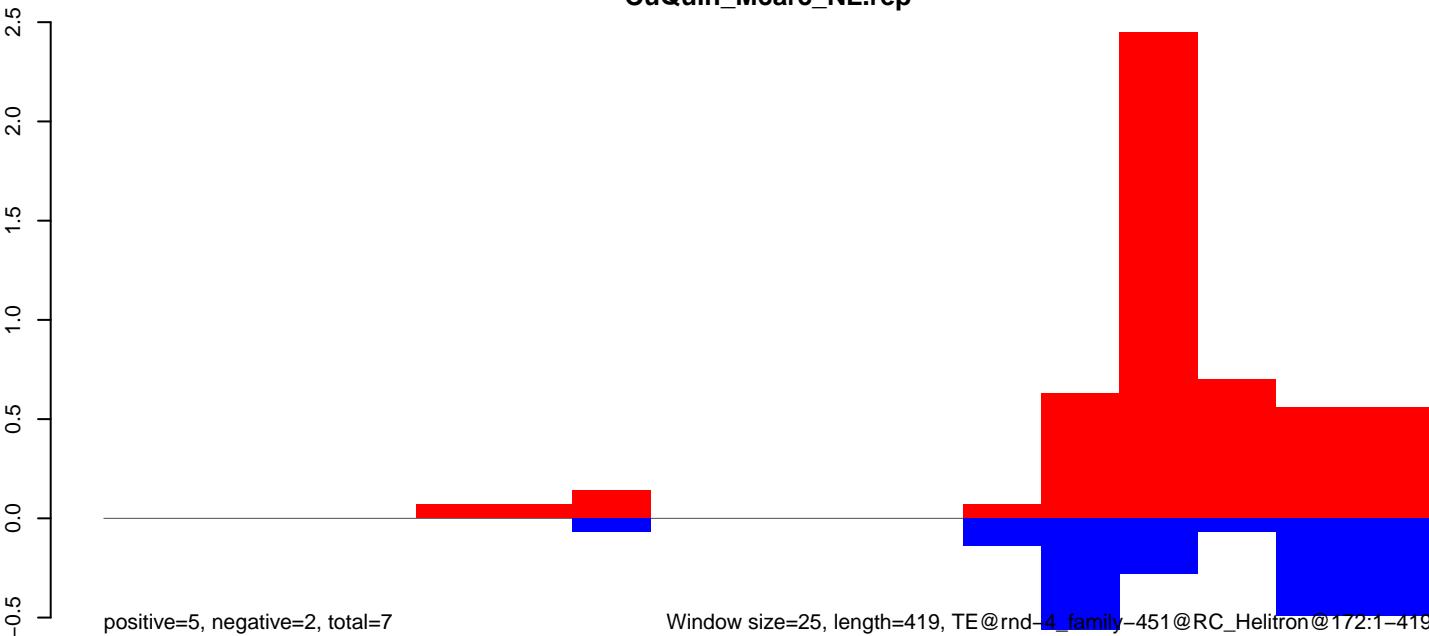
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

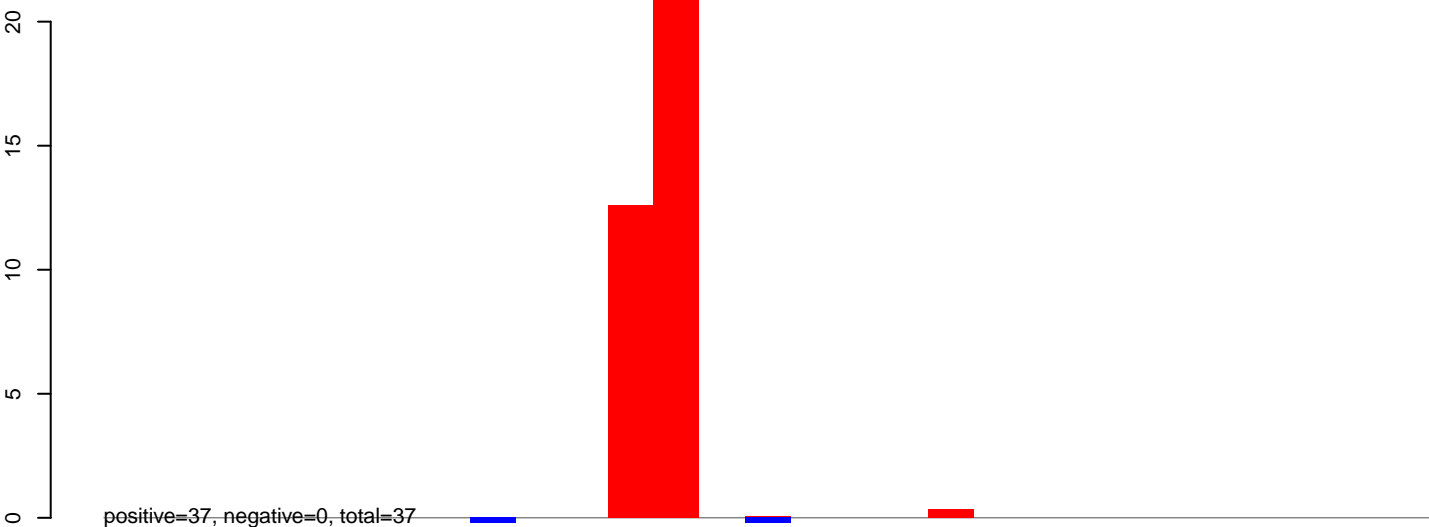


CuQuin_Mcarc_NL.rep

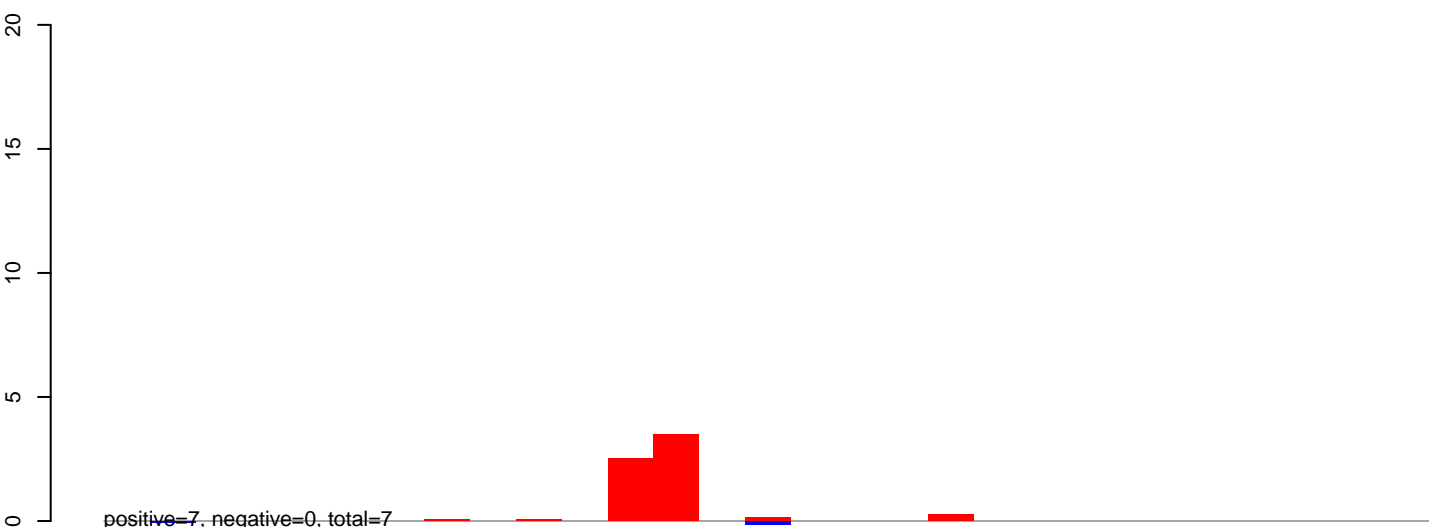


Window size=25, length=419, TE@rnd-4_family-451@RC_Helitron@172:1-419

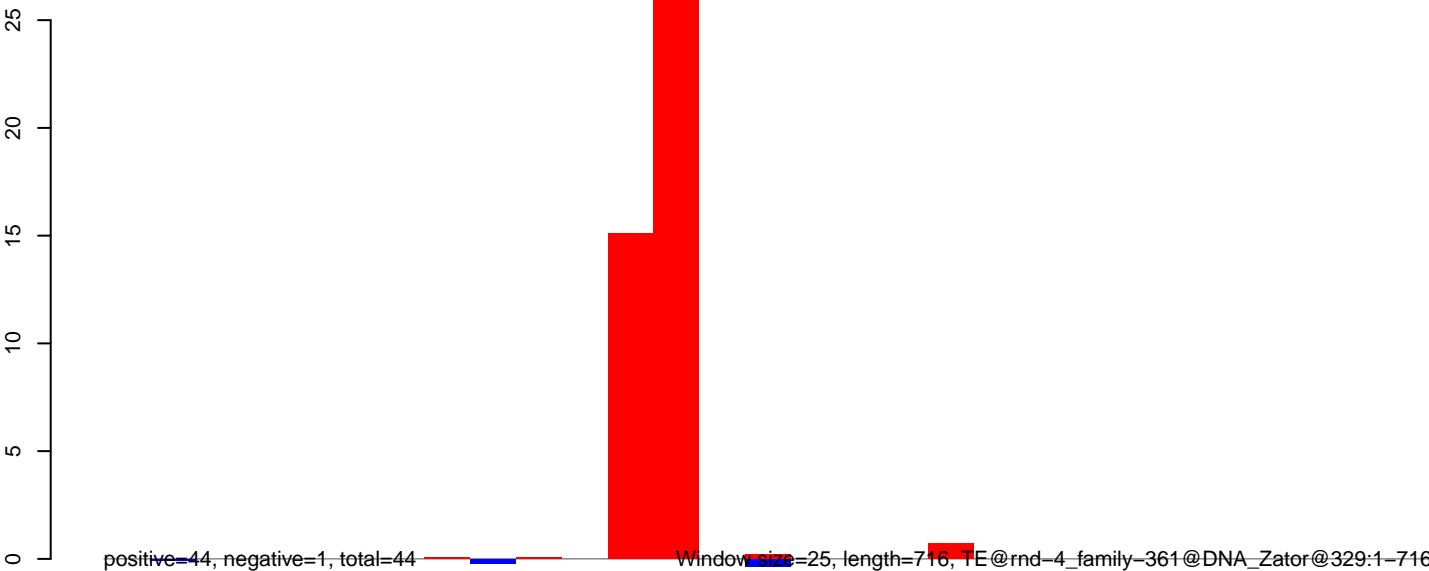
CuQuin_Mcarc_NL.18_23.rep



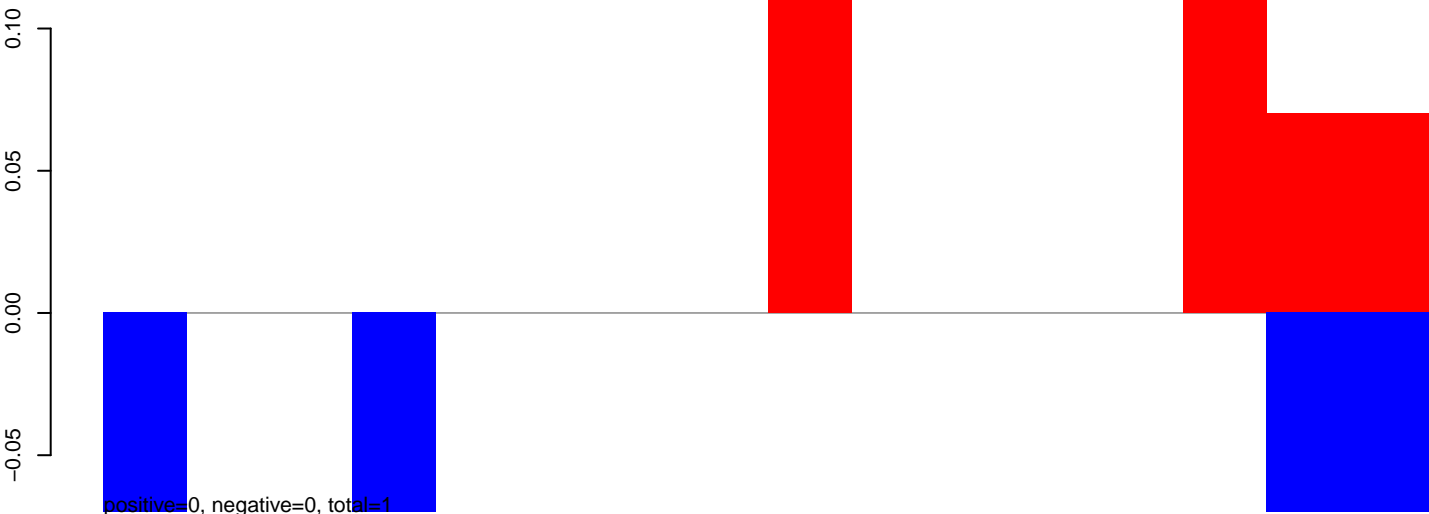
CuQuin_Mcarc_NL.24_35.rep



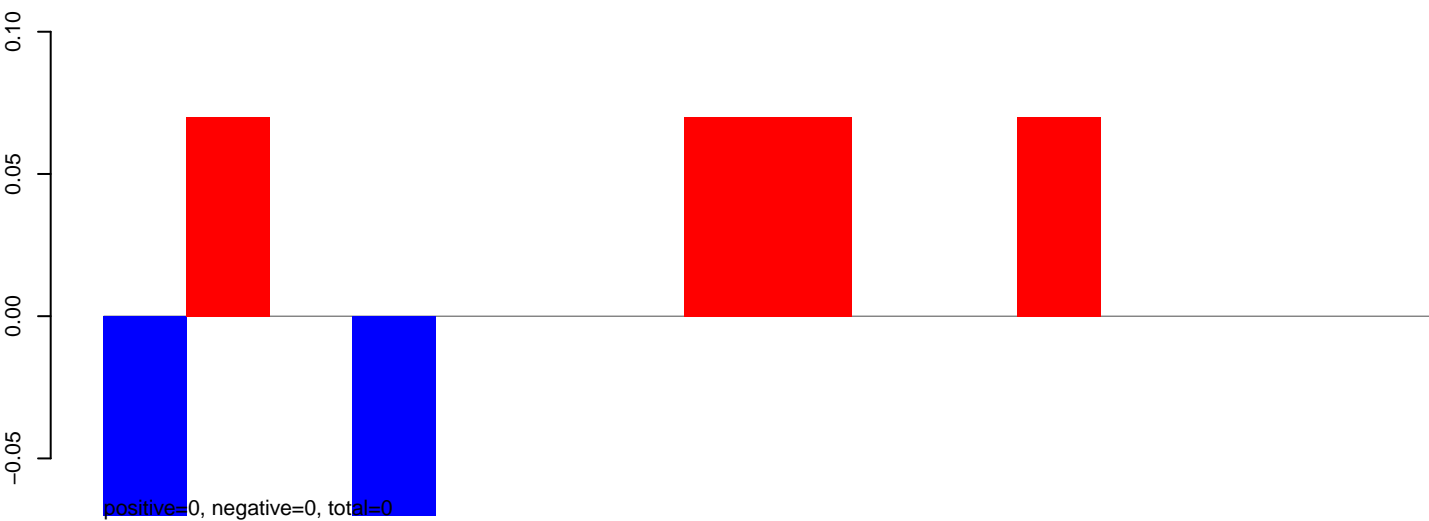
CuQuin_Mcarc_NL.rep



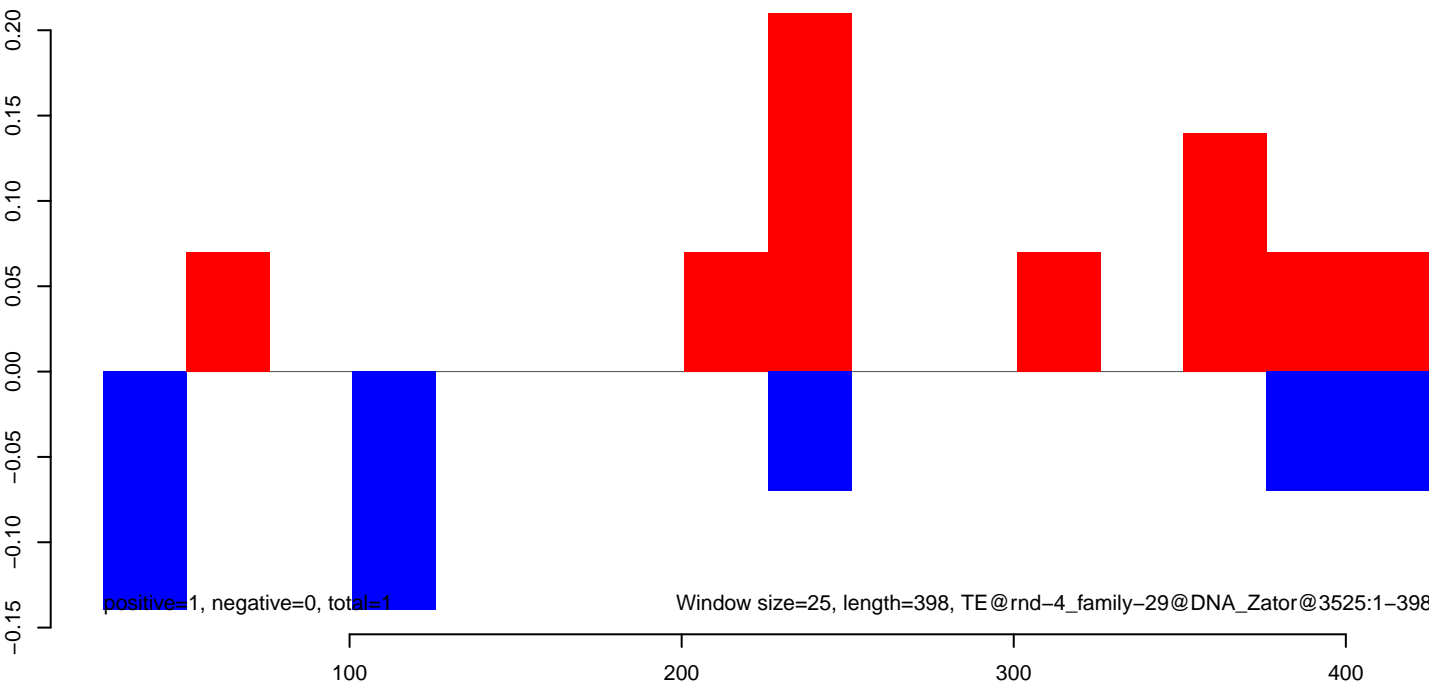
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

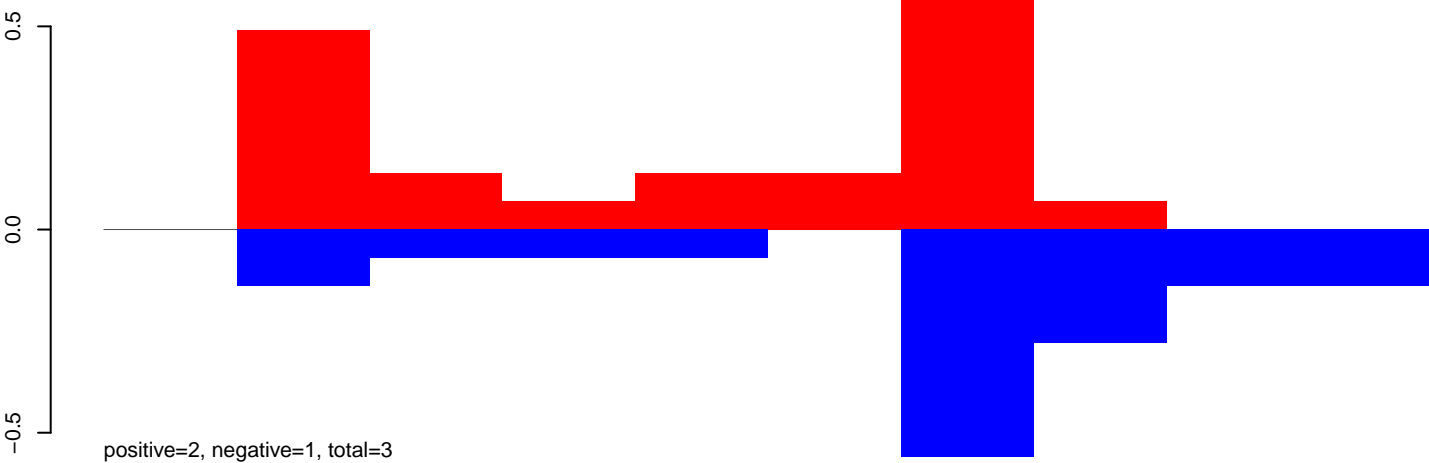


CuQuin_Mcarc_NL.rep

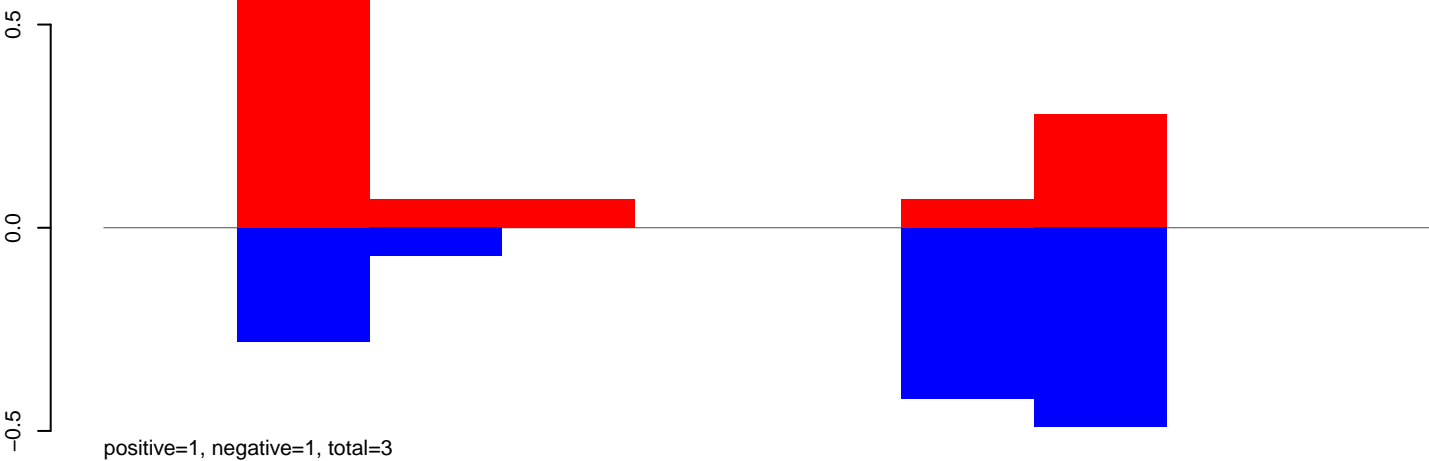


Window size=25, length=398, TE@rnd-4_family-29@DNA_Zator@3525:1-398

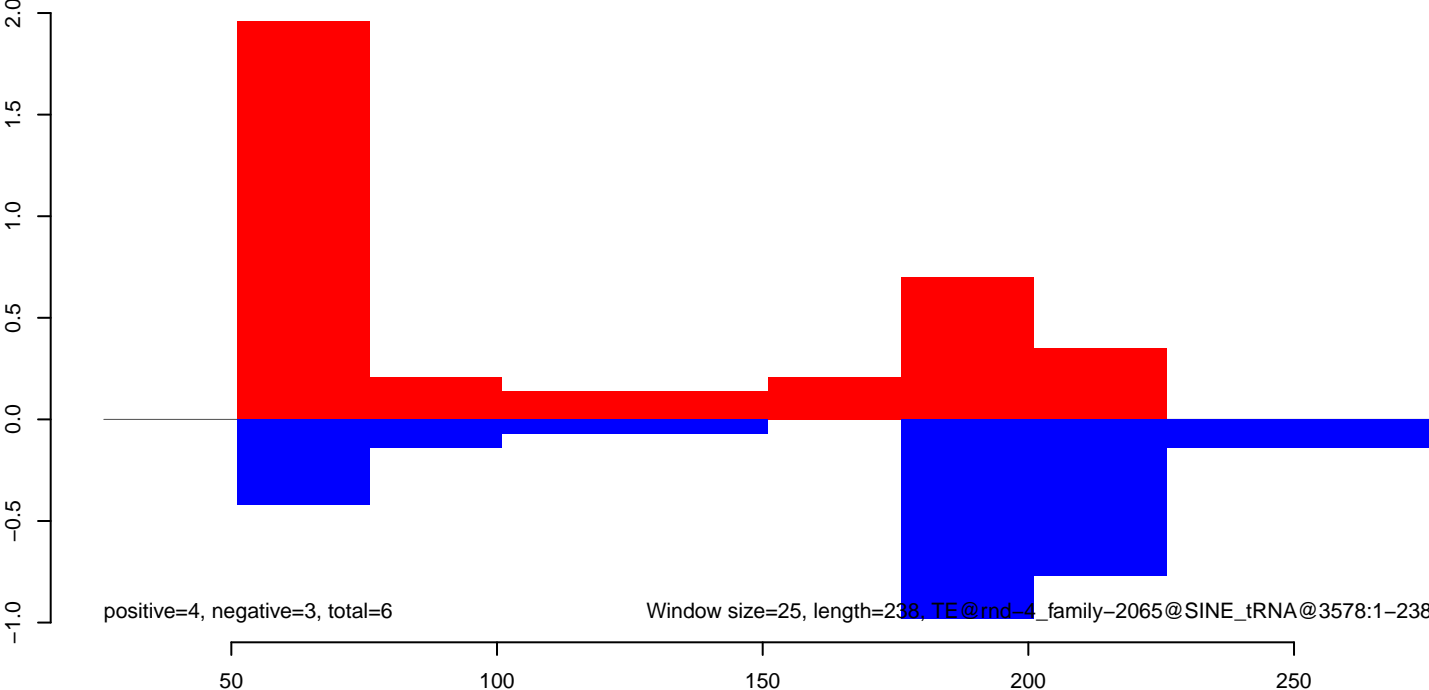
CuQuin_Mcarc_NL.18_23.rep



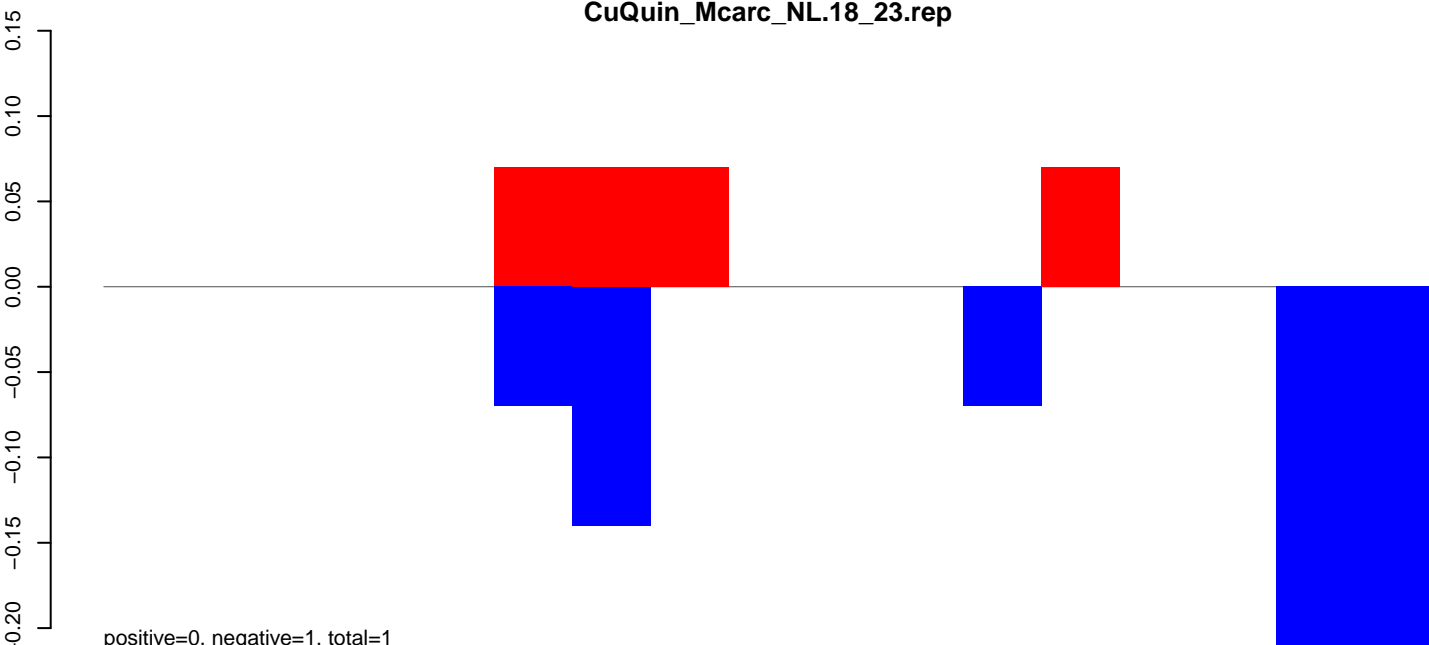
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

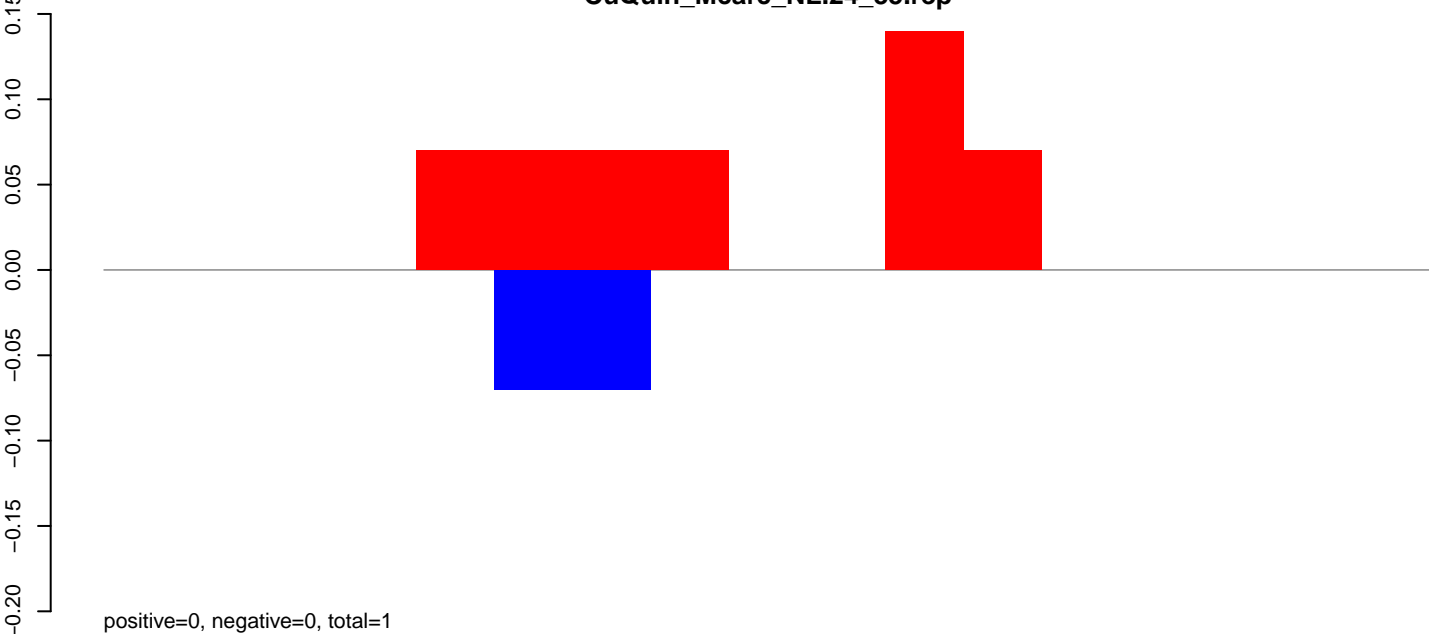


CuQuin_Mcarc_NL.18_23.rep



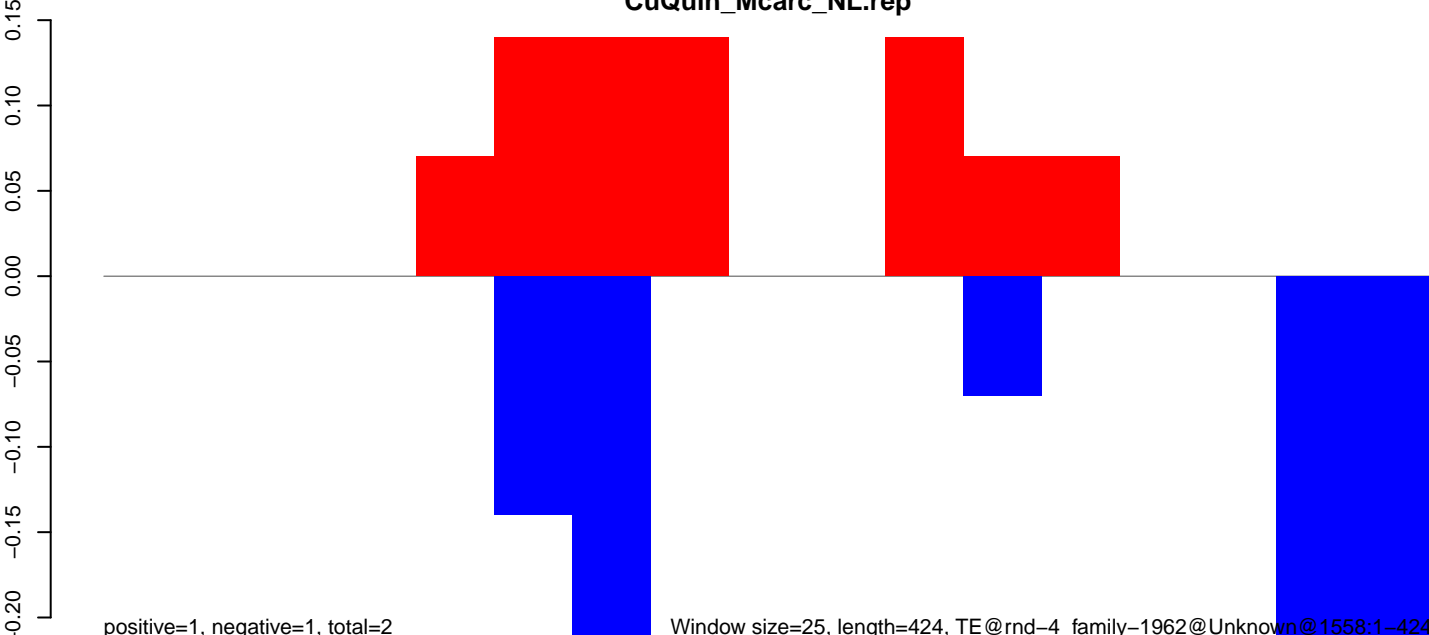
positive=0, negative=1, total=1

CuQuin_Mcarc_NL.24_35.rep



positive=0, negative=0, total=1

CuQuin_Mcarc_NL.rep



positive=1, negative=1, total=2

Window size=25, length=424, TE@rnd-4_family-1962@Unknown@1558:1-424

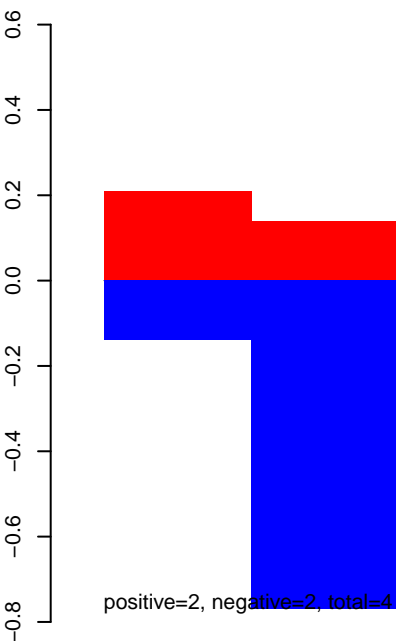
100

200

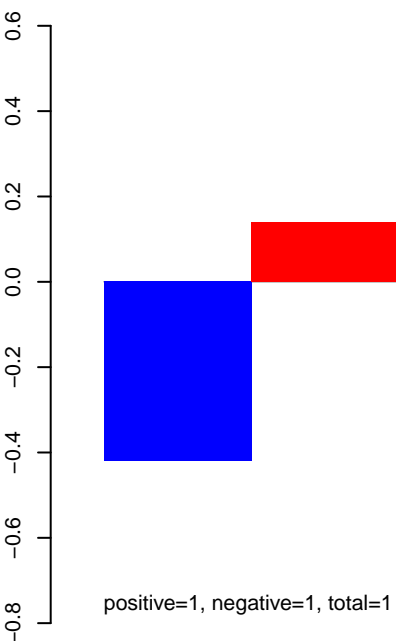
300

400

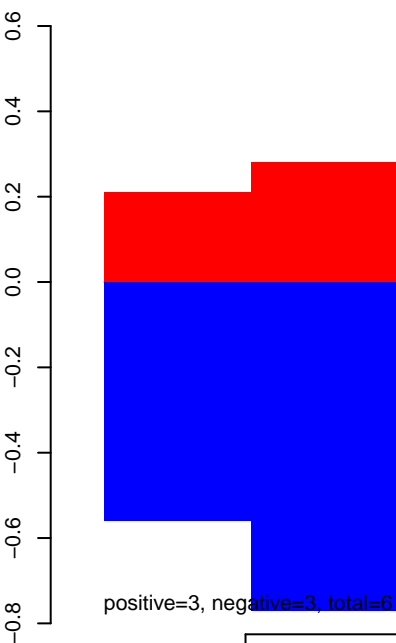
CuQuin_Mcarc_NL.18_23.rep



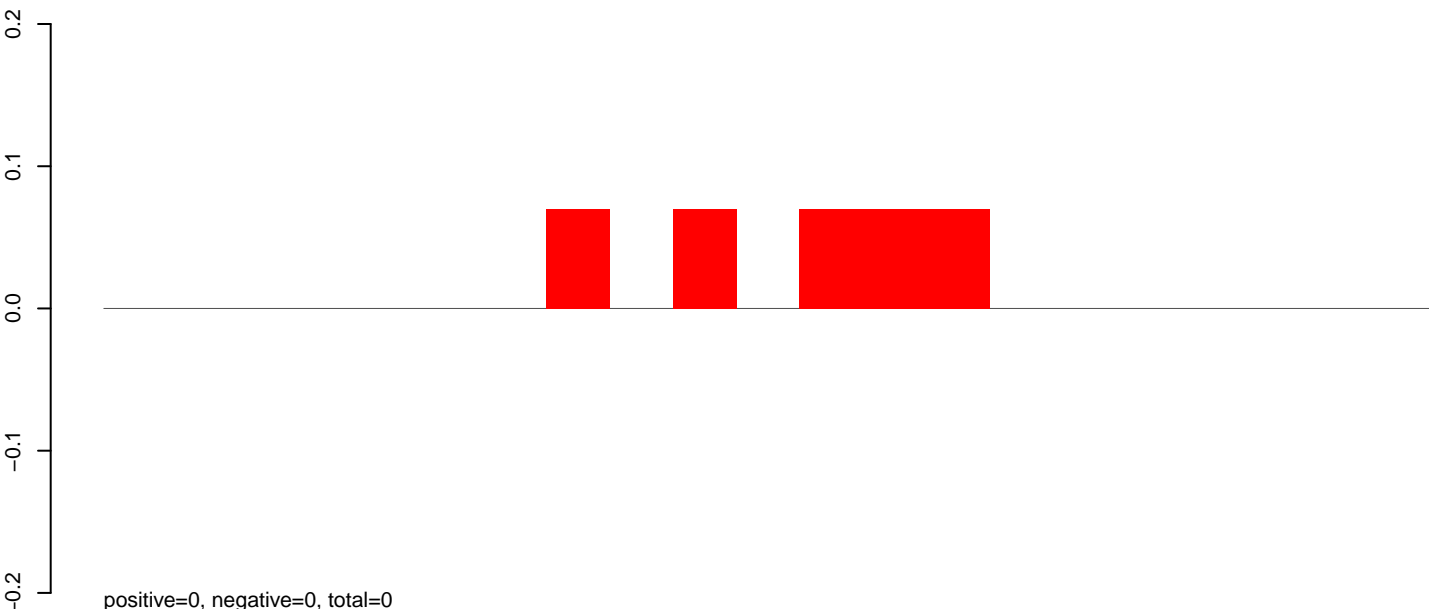
CuQuin_Mcarc_NL.24_35.rep



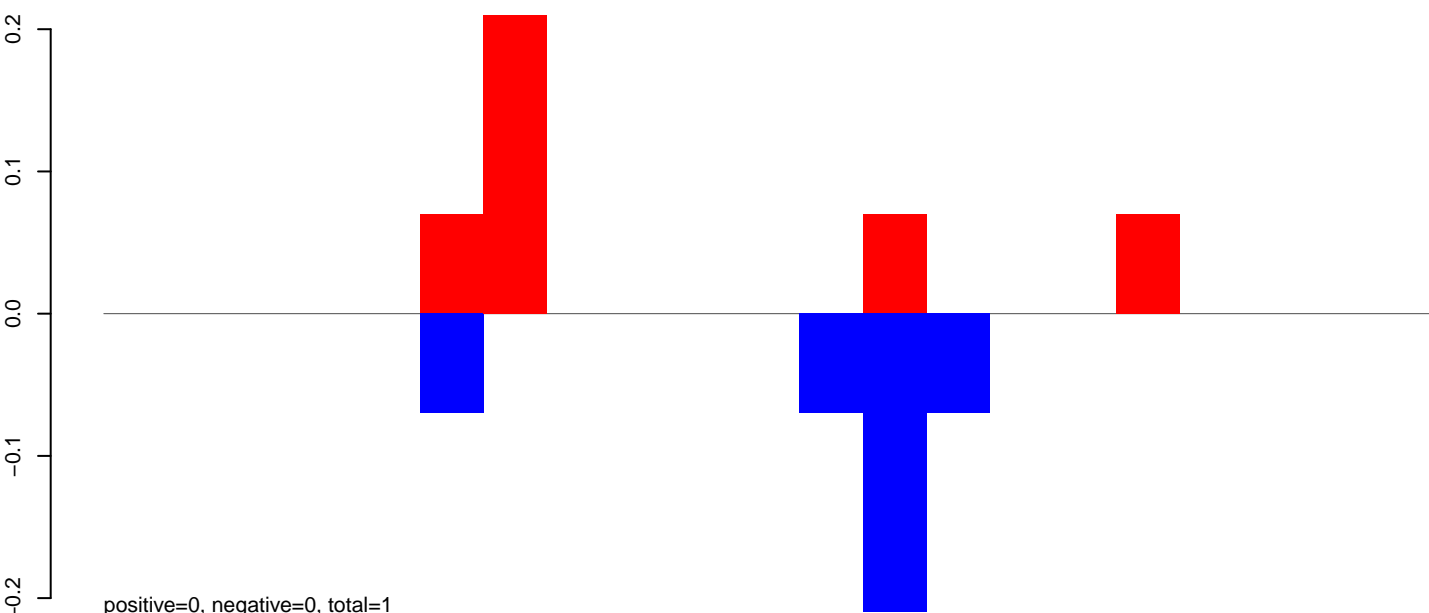
CuQuin_Mcarc_NL.rep



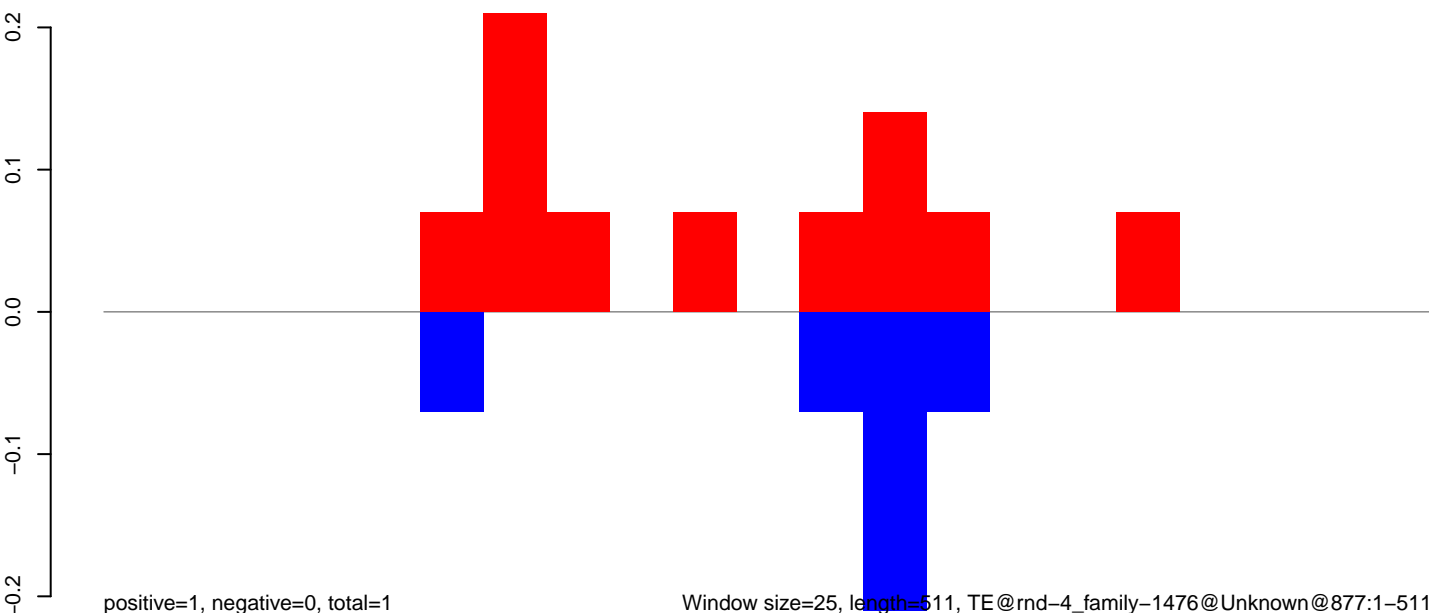
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

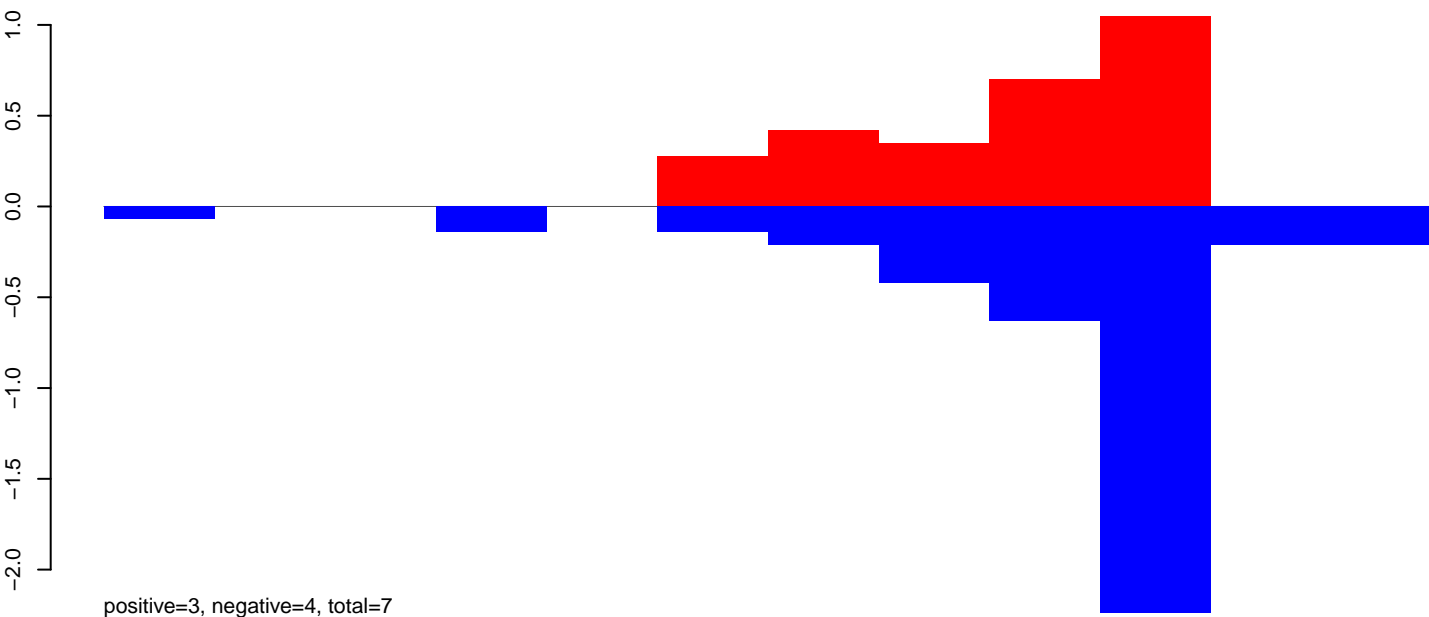


CuQuin_Mcarc_NL.rep

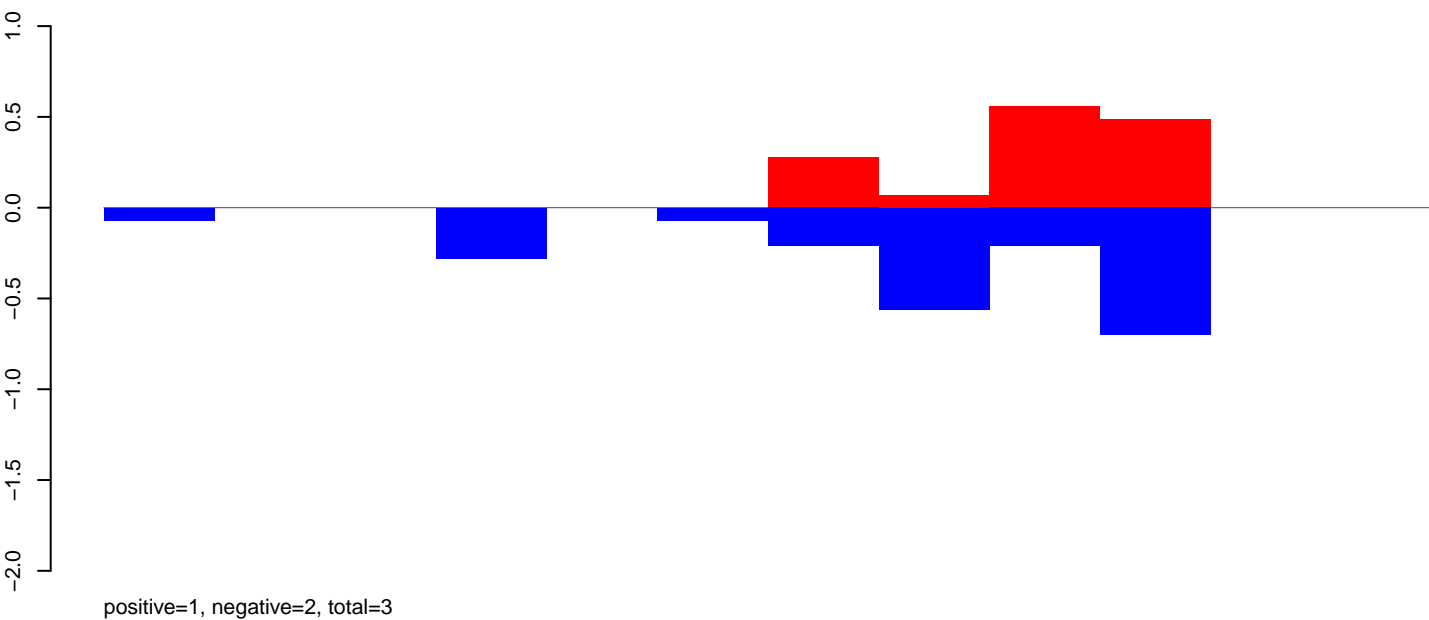


Window size=25, length=511, TE@rnd-4_family-1476@Unknown@877:1-511

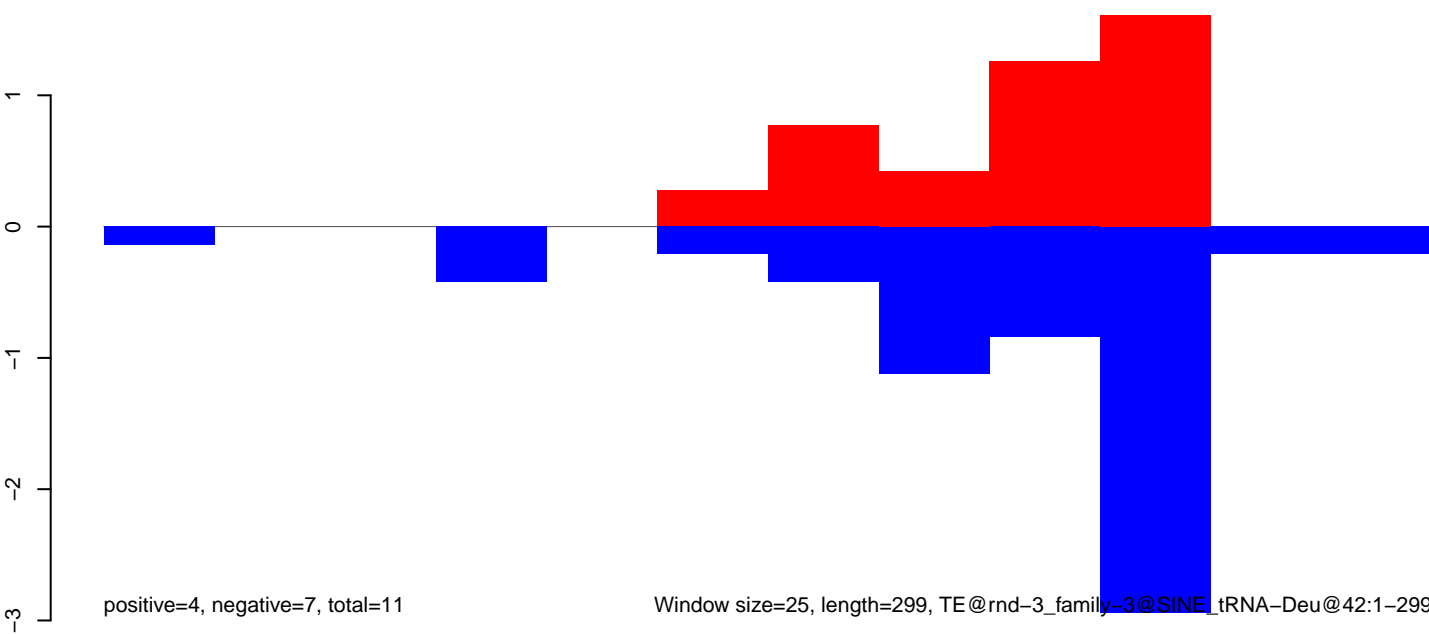
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

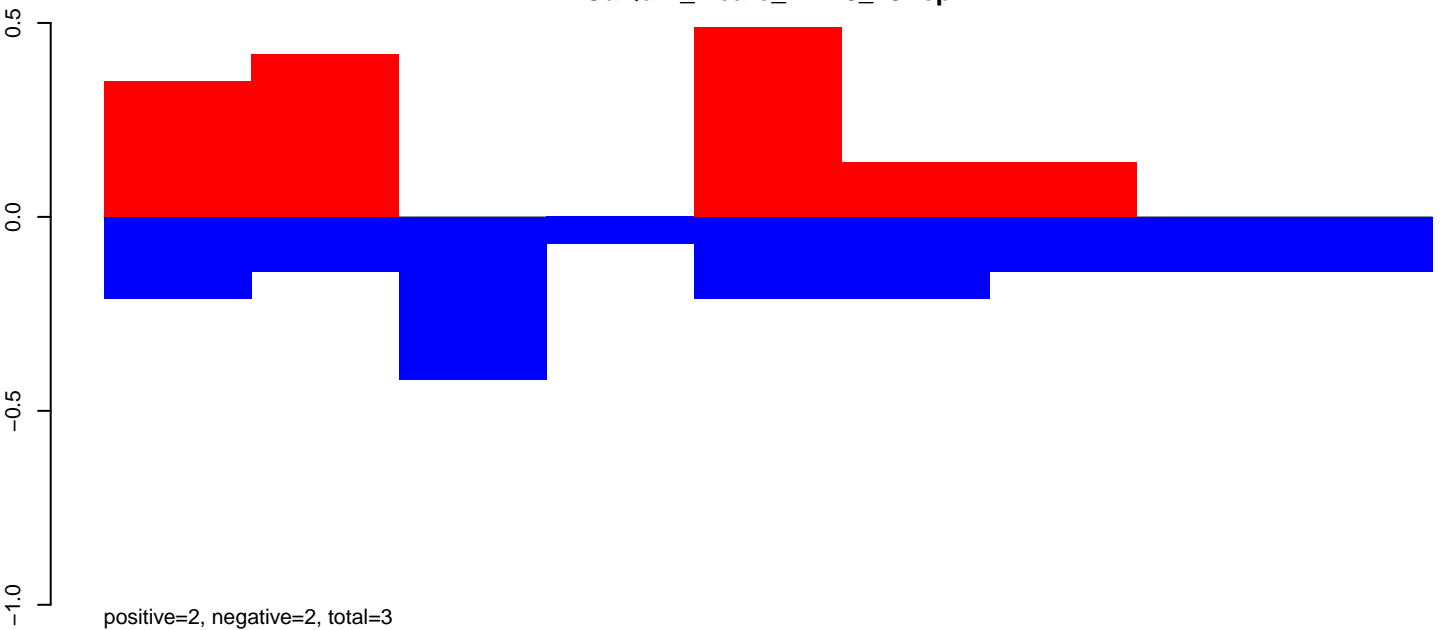


CuQuin_Mcarc_NL.rep

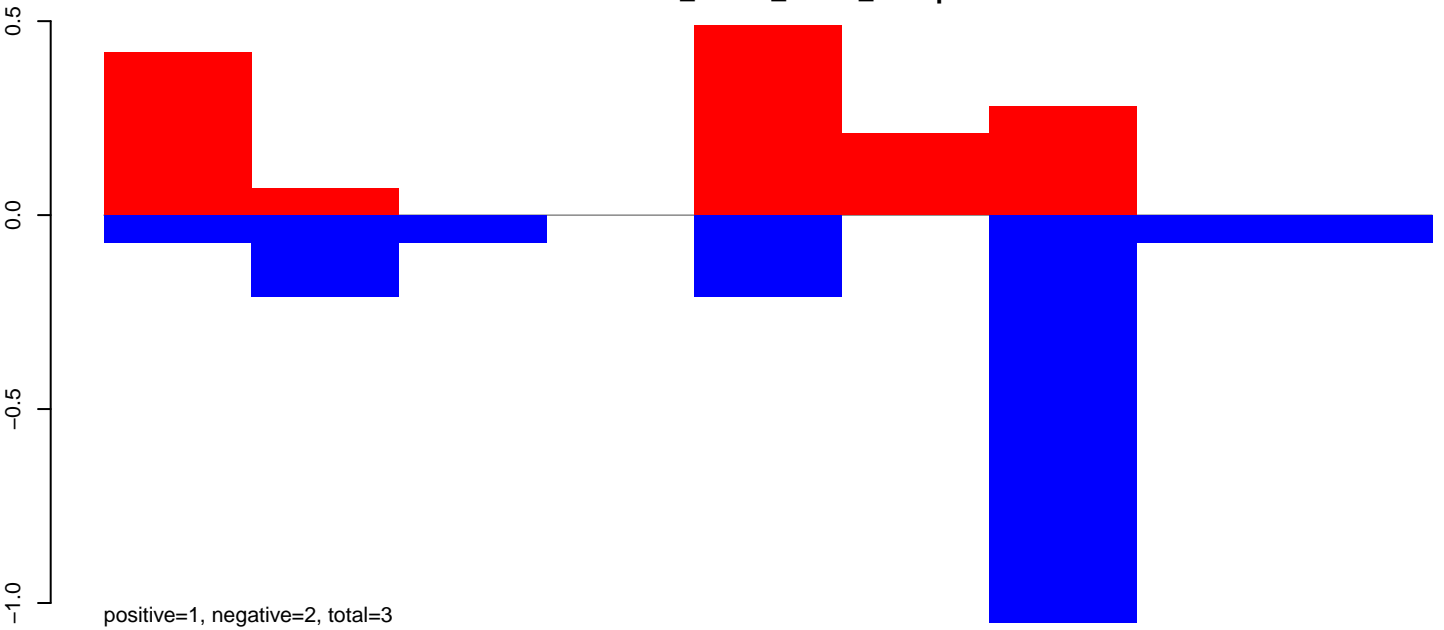


Window size=25, length=299, TE@rnd-3_family-3@SINE_tRNA-Deu@42:1-299

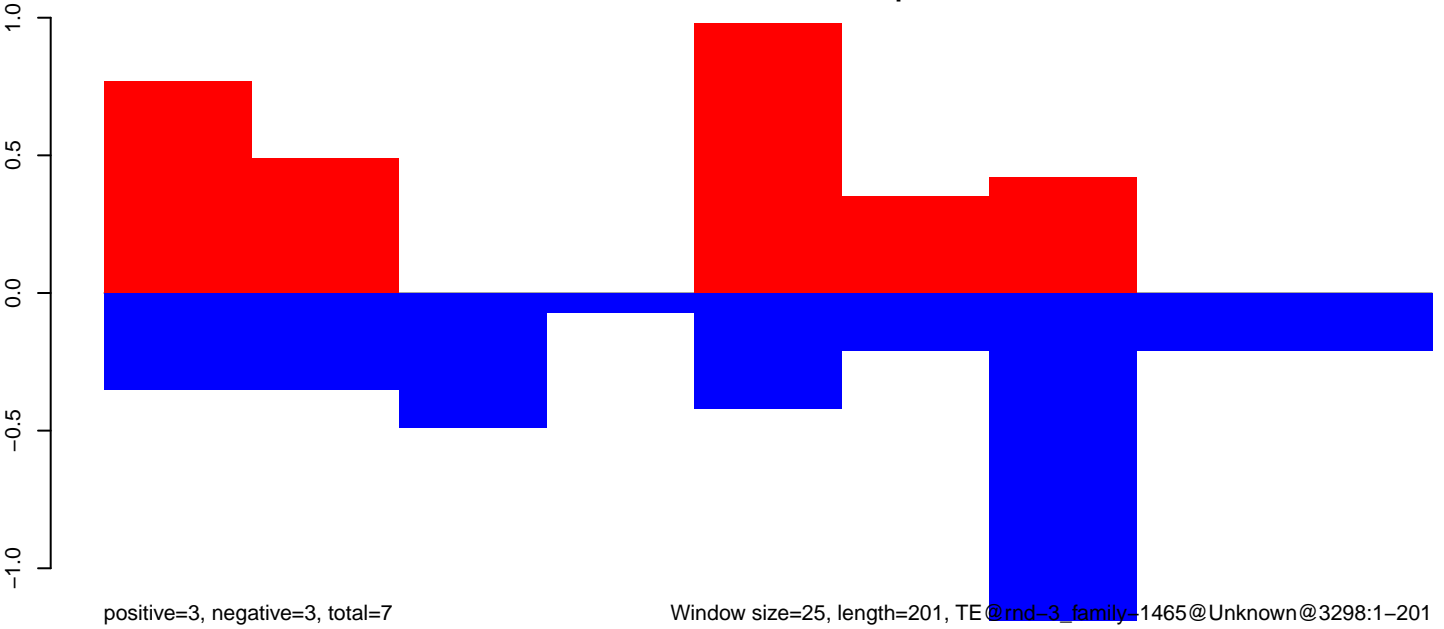
CuQuin_Mcarc_NL.18_23.rep



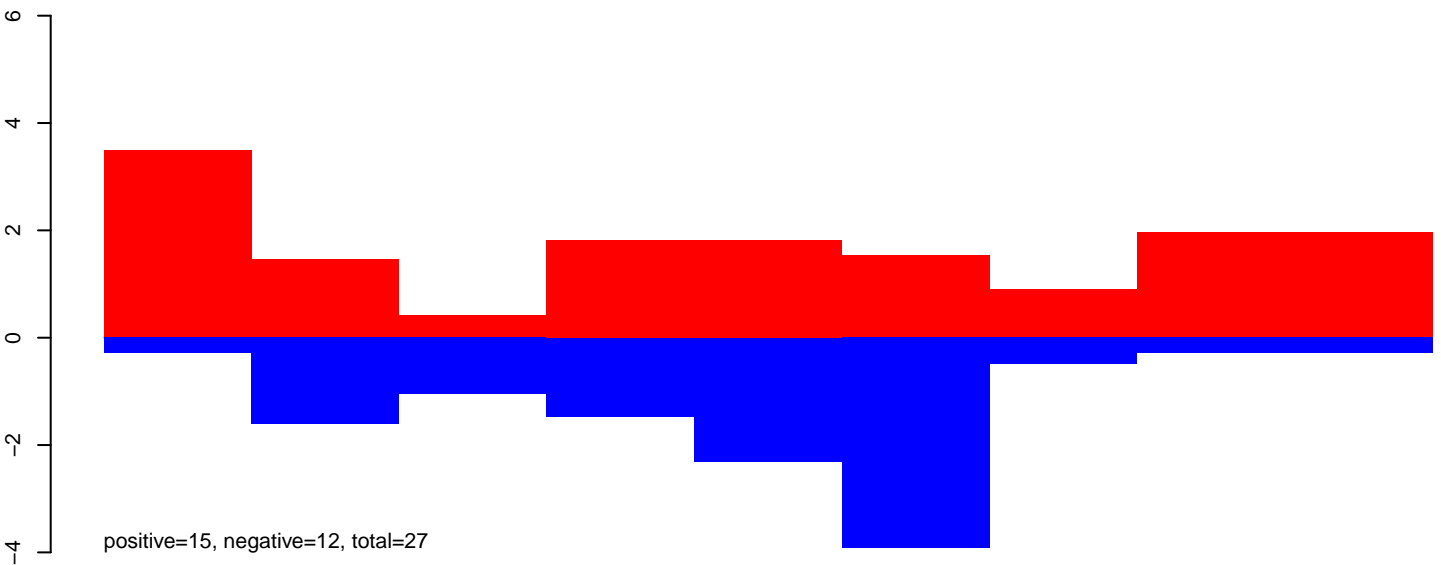
CuQuin_Mcarc_NL.24_35.rep



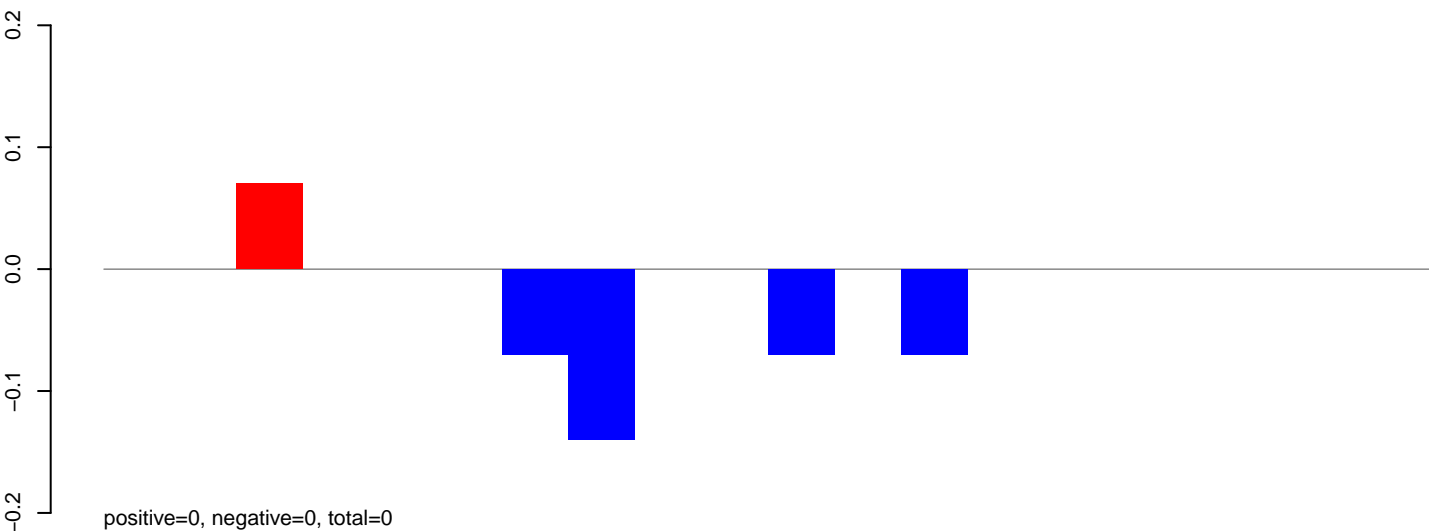
CuQuin_Mcarc_NL.rep



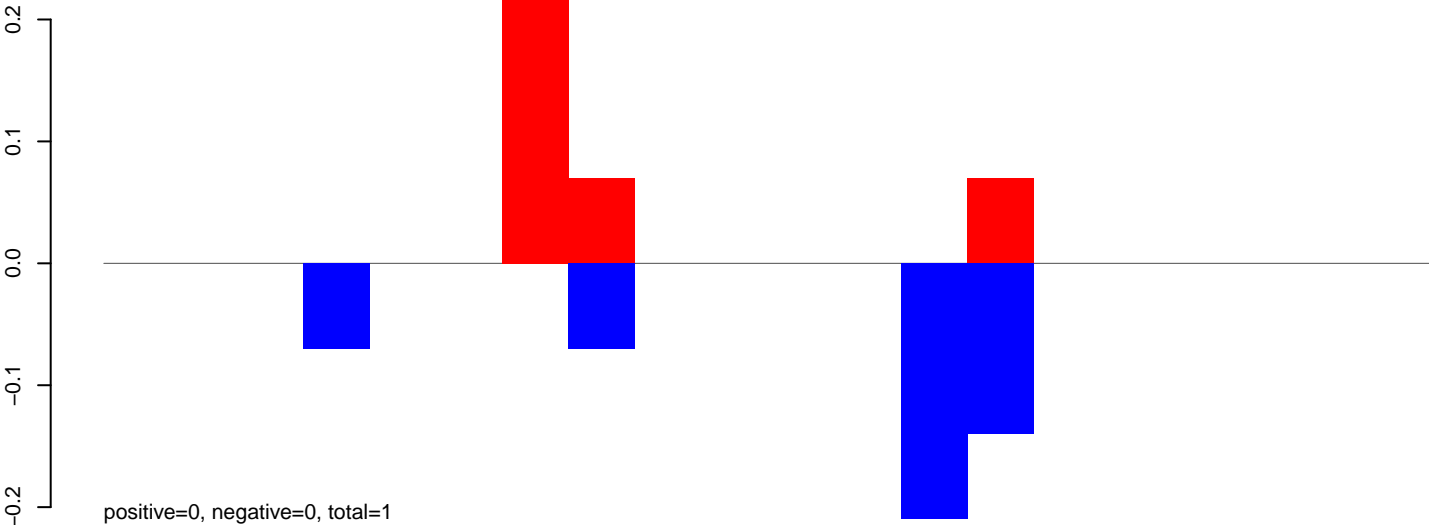
CuQuin_Mcarc_NL.18_23.rep



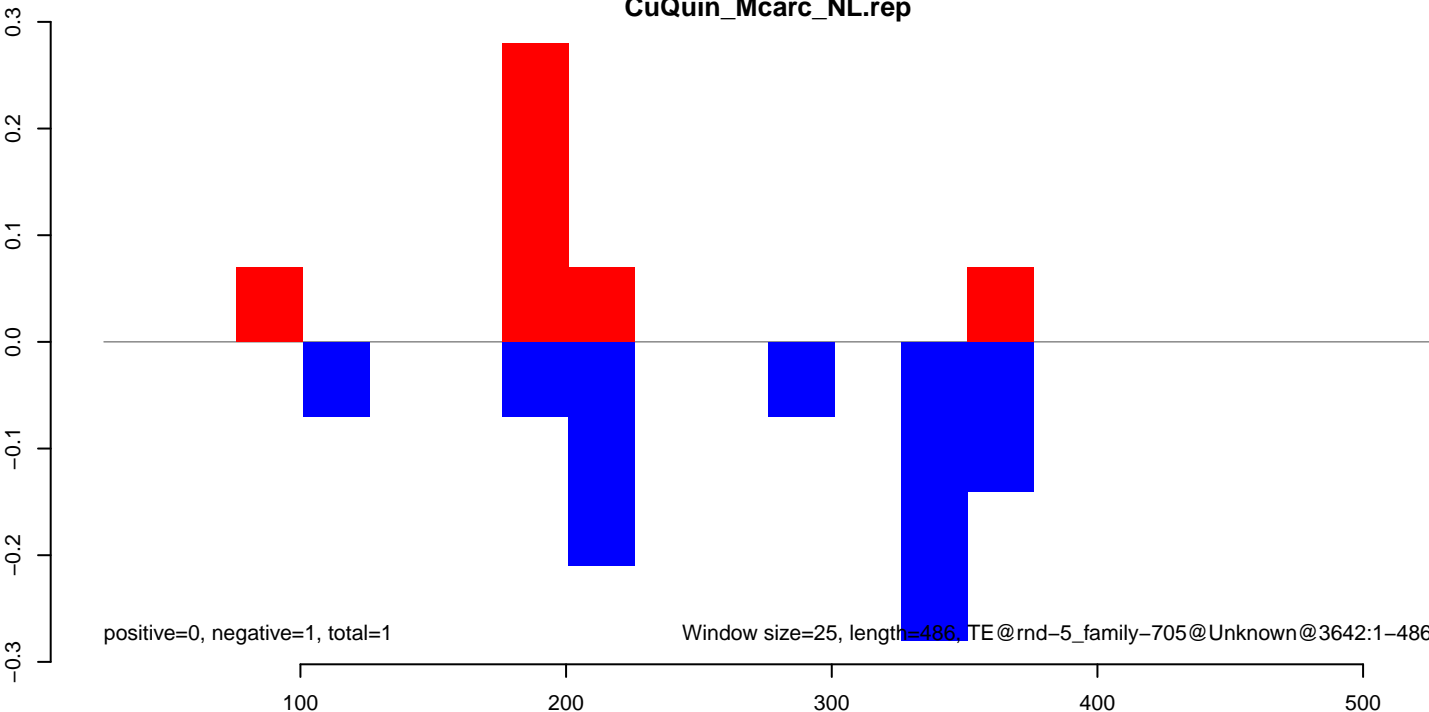
CuQuin_Mcarc_NL.18_23.rep



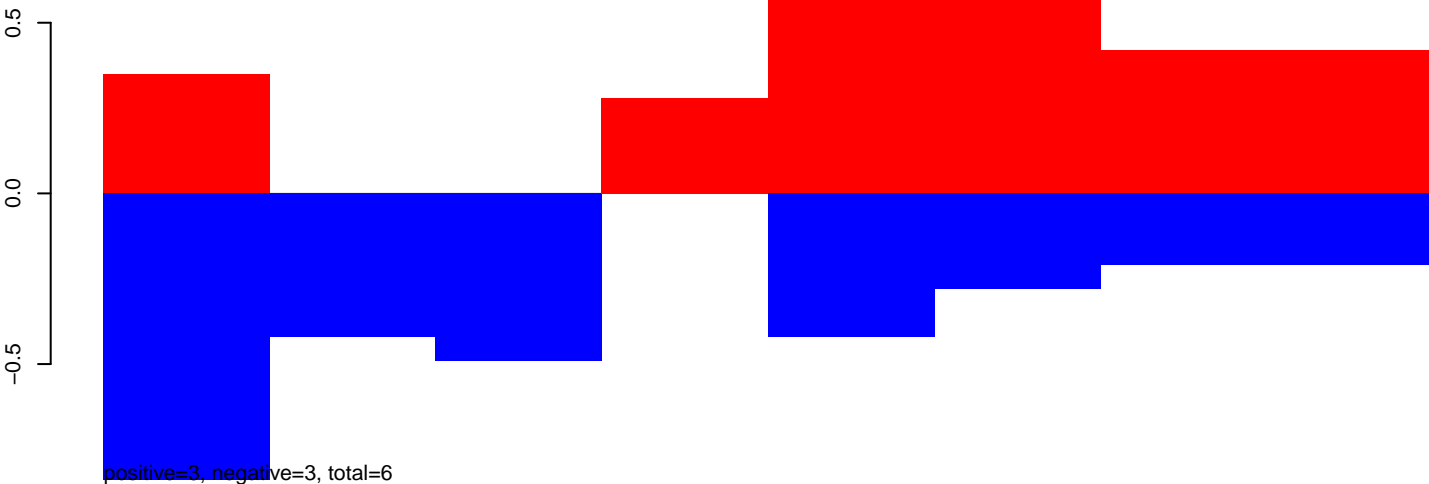
CuQuin_Mcarc_NL.24_35.rep



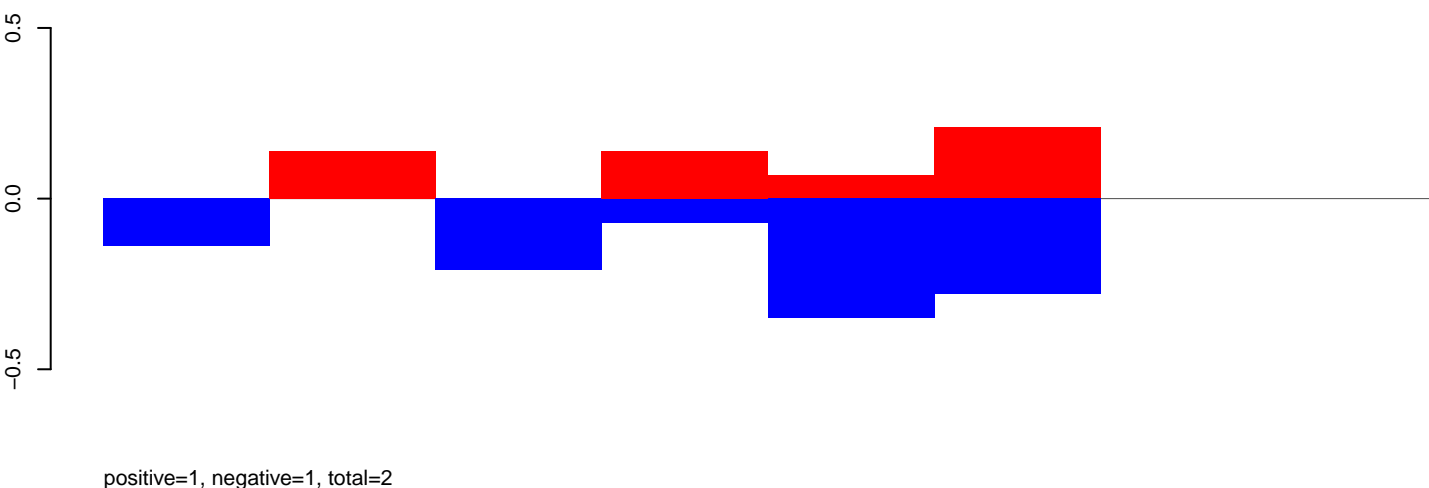
CuQuin_Mcarc_NL.rep



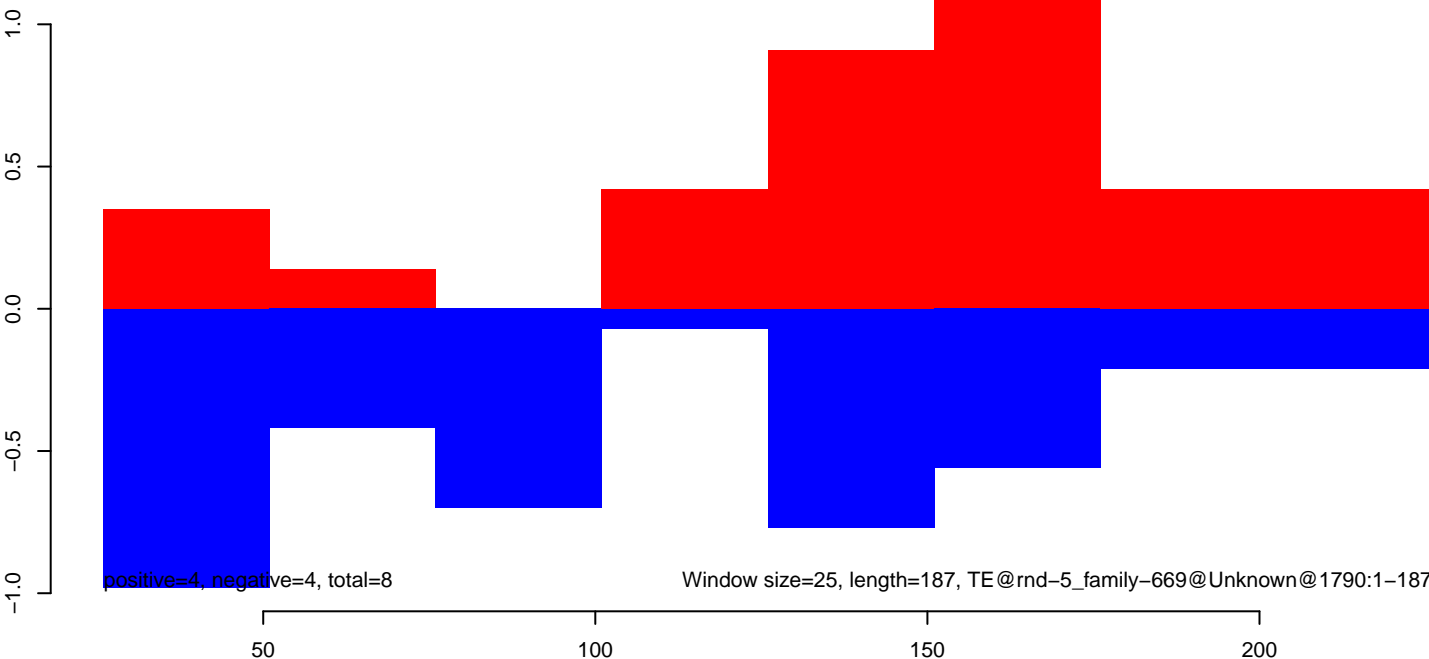
CuQuin_Mcarc_NL.18_23.rep



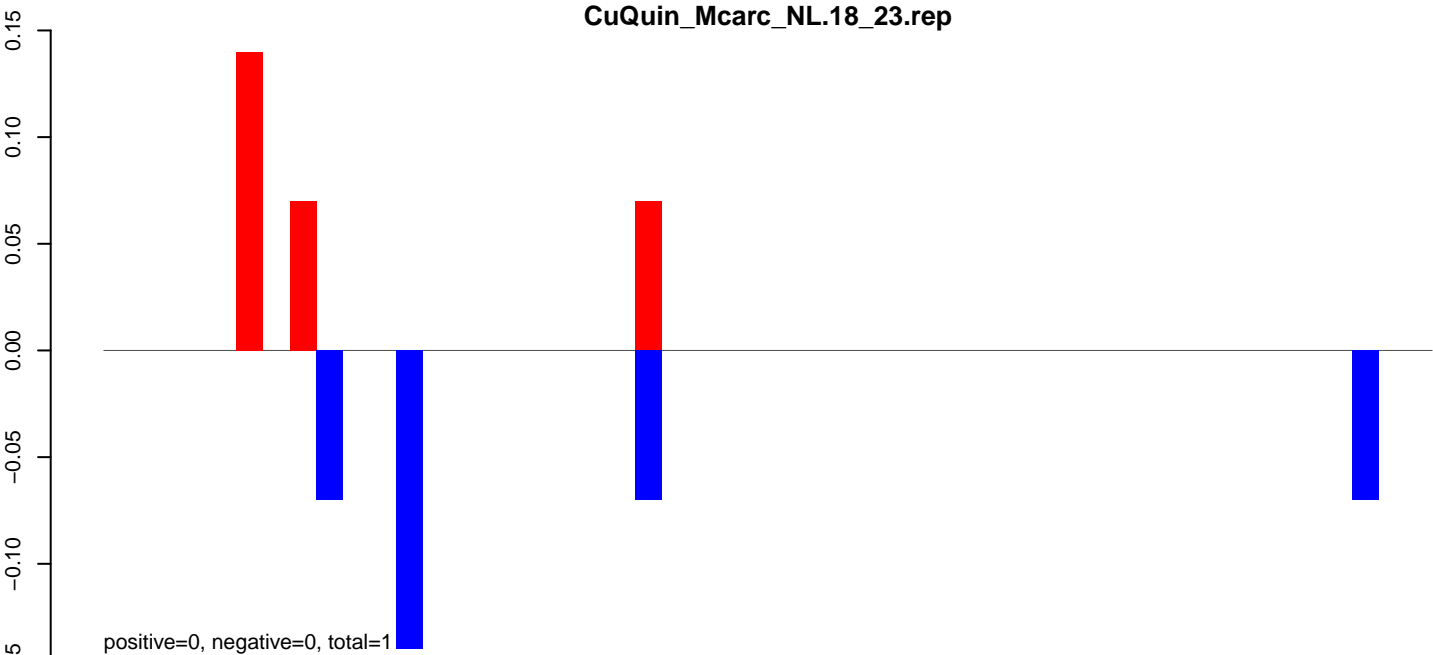
CuQuin_Mcarc_NL.24_35.rep



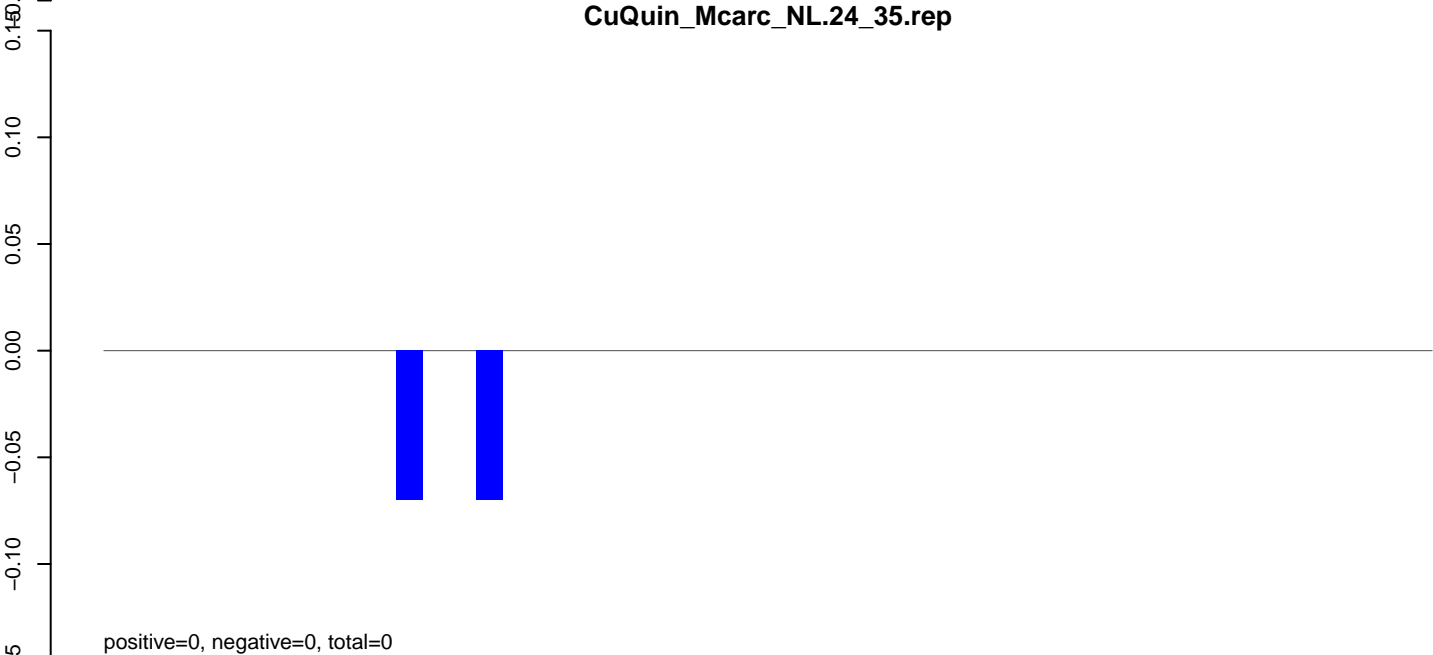
CuQuin_Mcarc_NL.rep



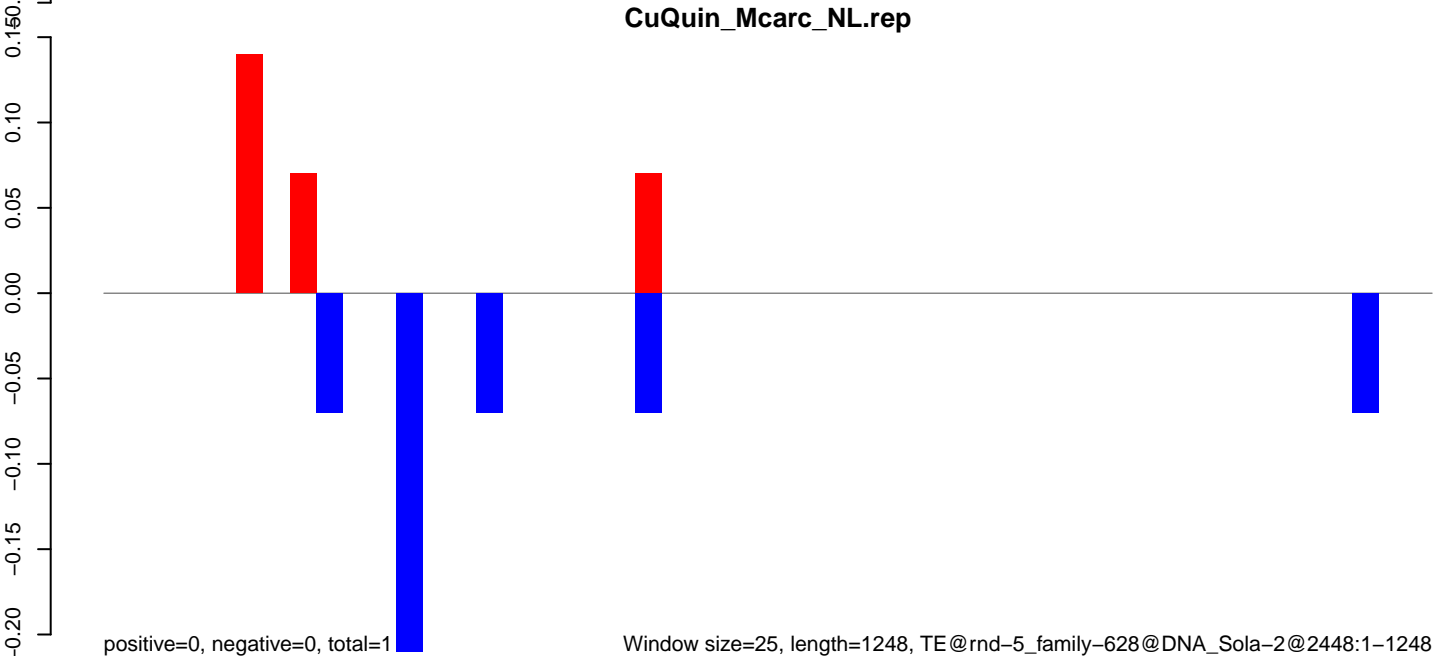
CuQuin_Mcarc_NL.18_23.rep



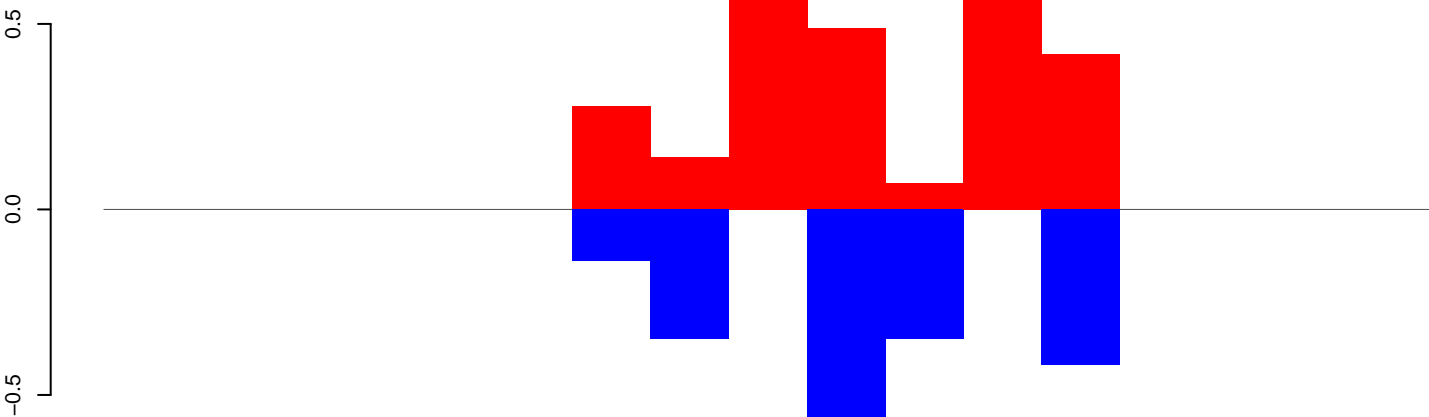
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

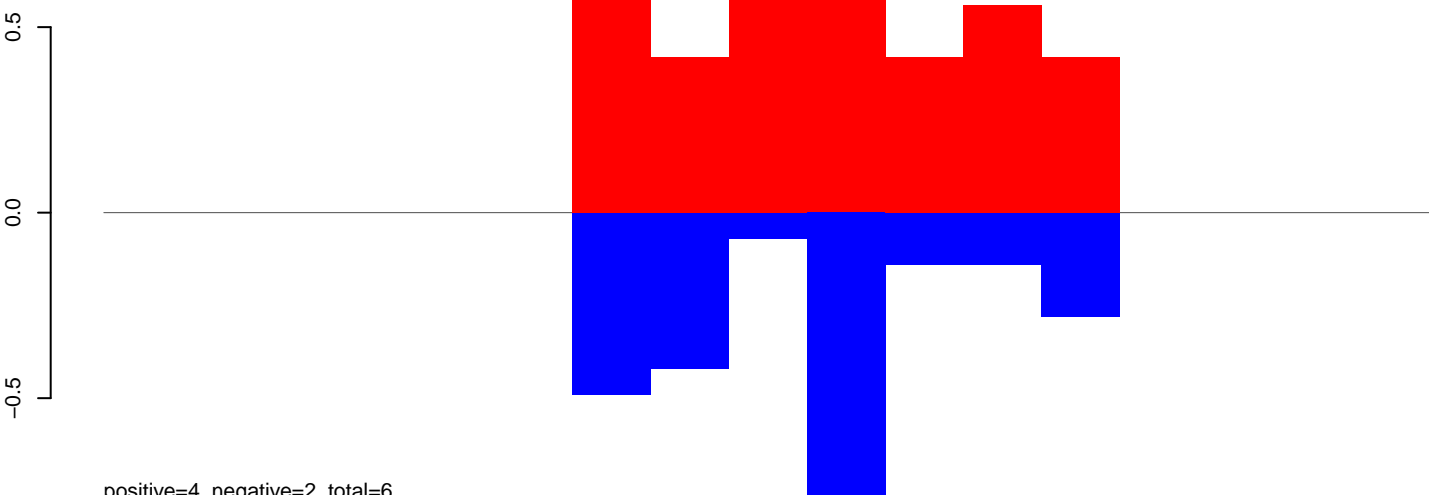


CuQuin_Mcarc_NL.18_23.rep



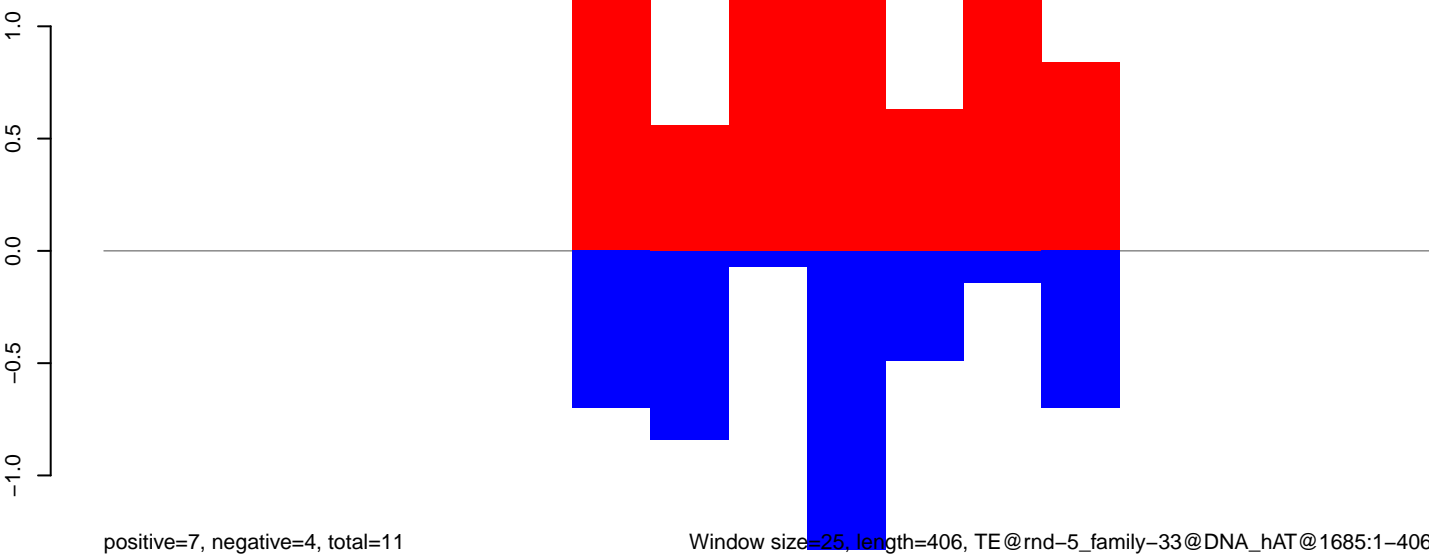
positive=3, negative=2, total=5

CuQuin_Mcarc_NL.24_35.rep



positive=4, negative=2, total=6

CuQuin_Mcarc_NL.rep



positive=7, negative=4, total=11

Window size=25, length=406, TE@rnd-5_family-33@DNA_hAT@1685:1-406

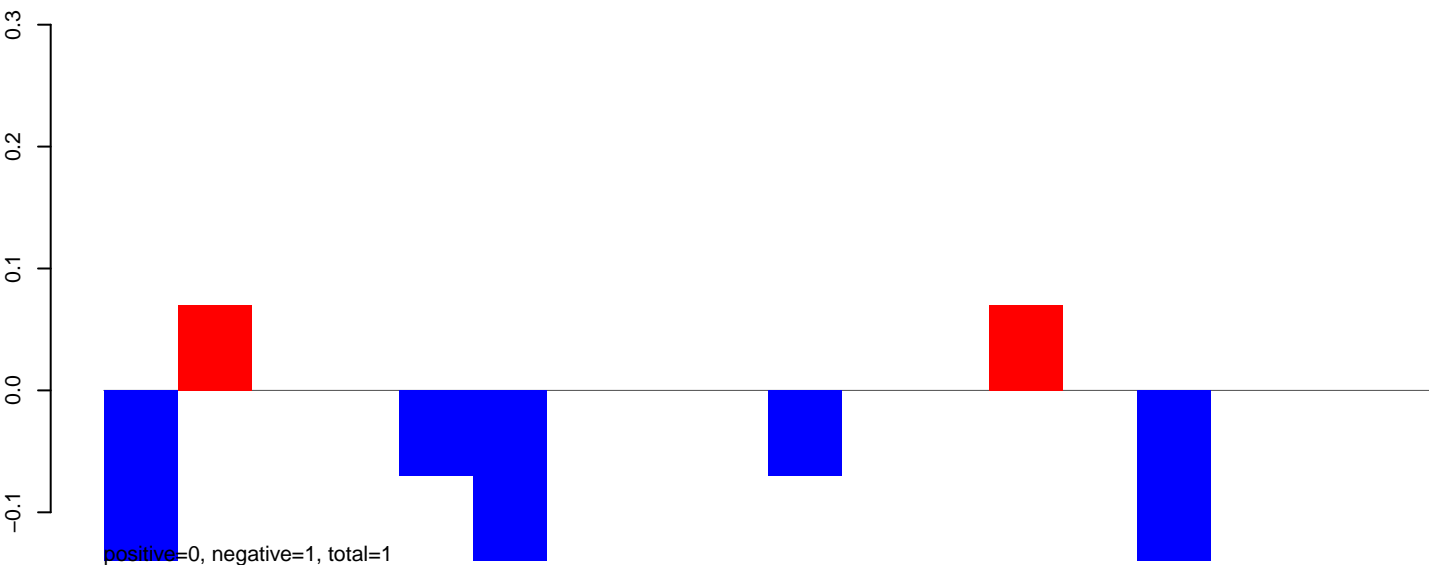
100

200

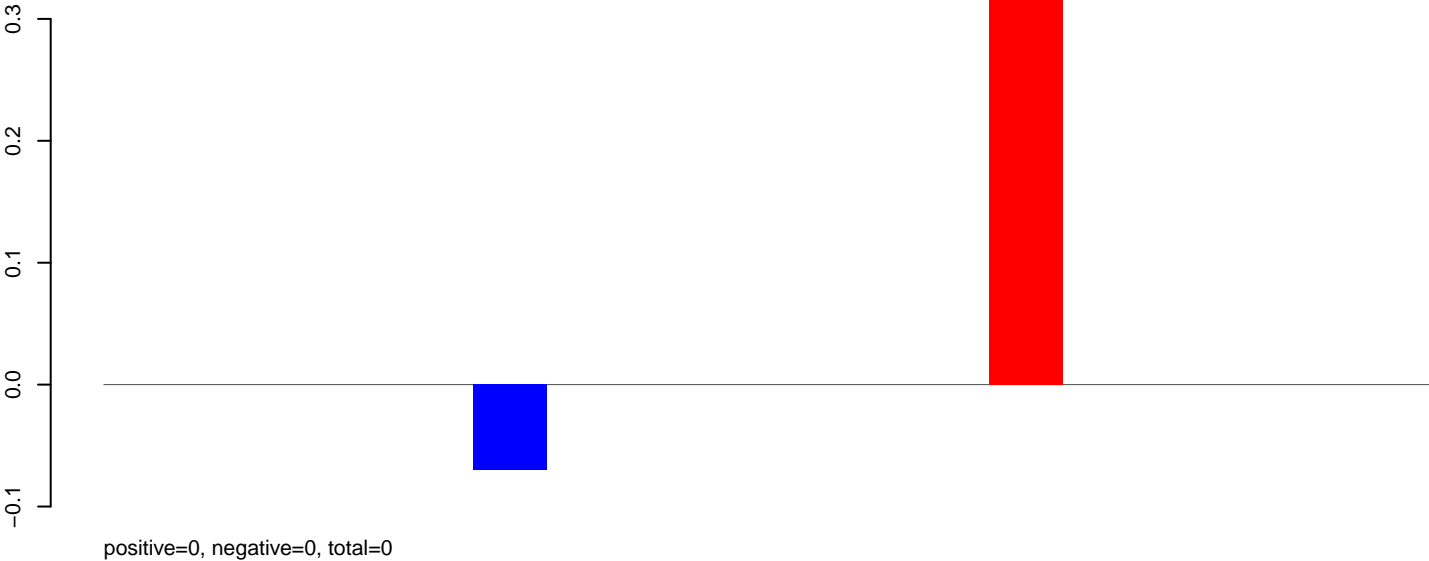
300

400

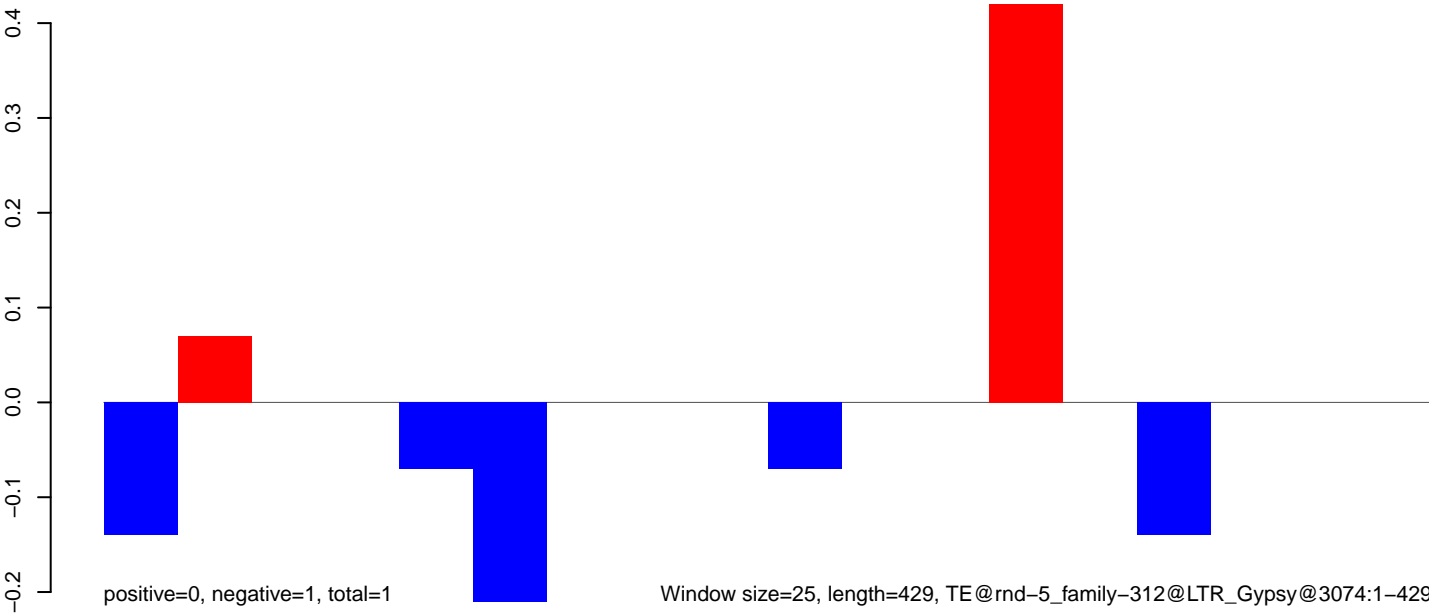
CuQuin_Mcarc_NL.18_23.rep



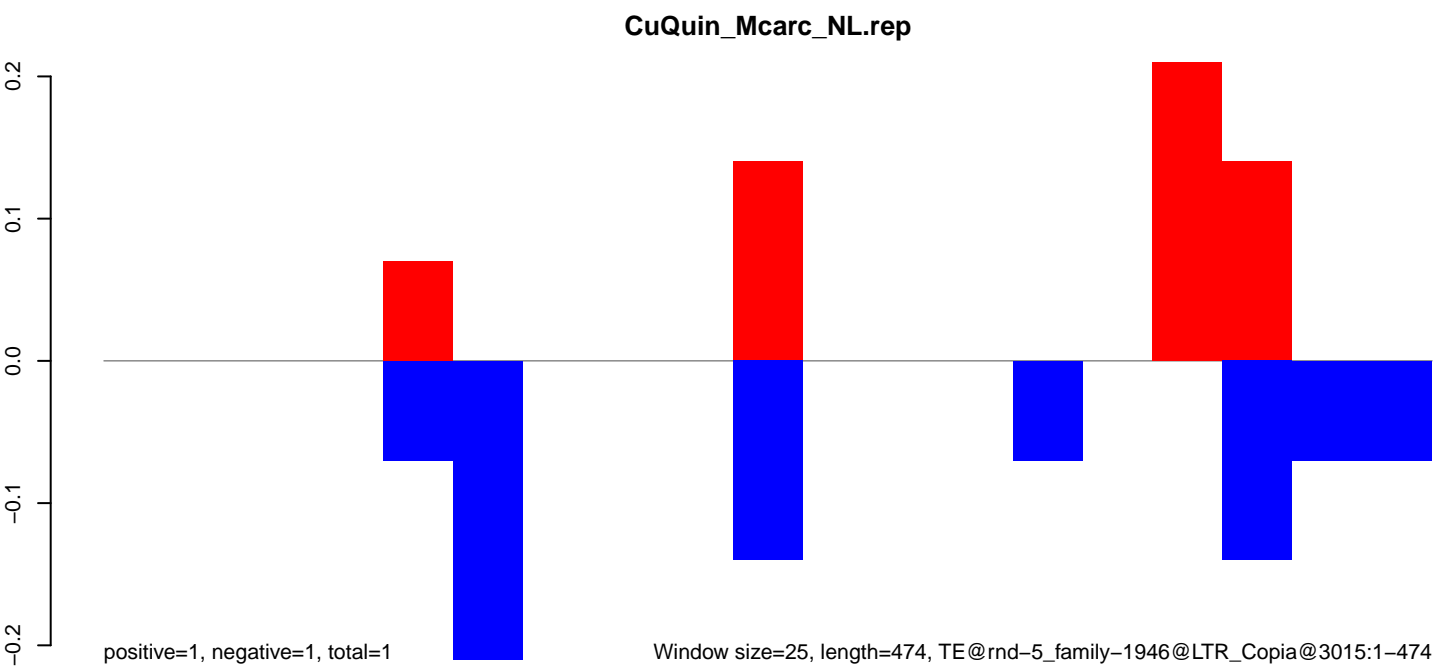
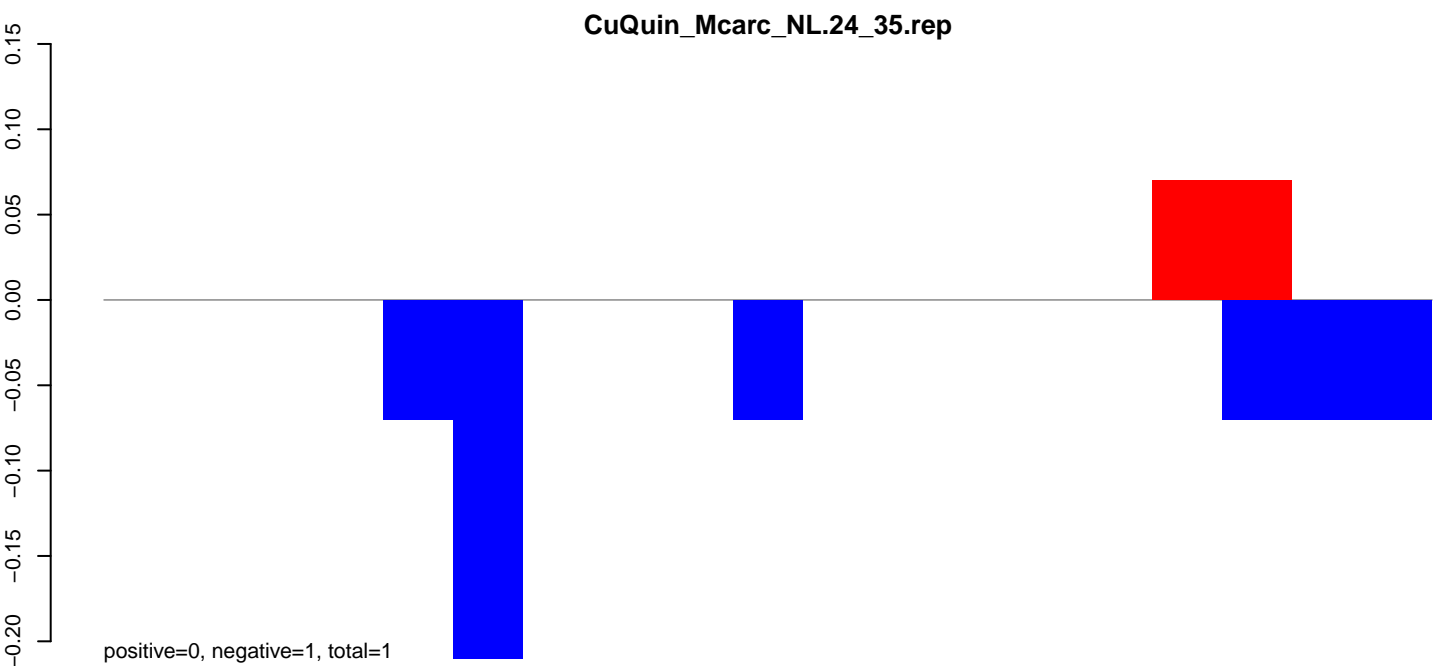
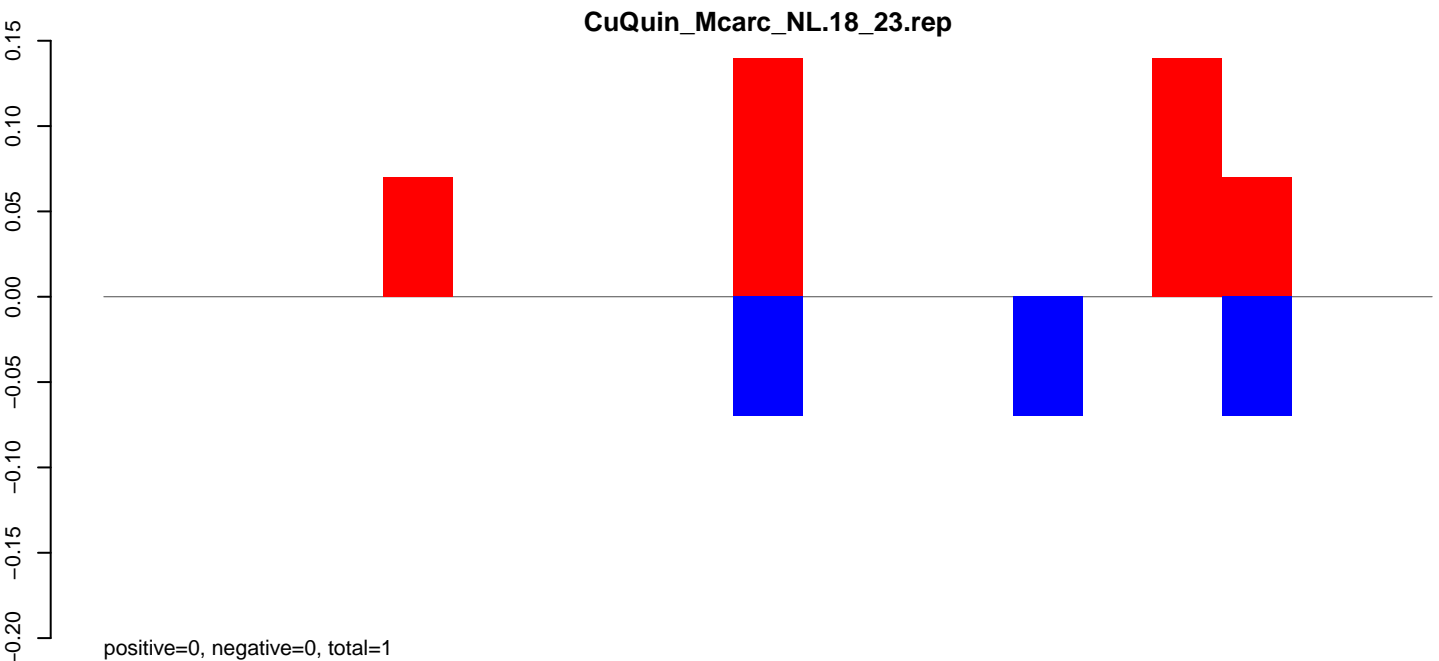
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

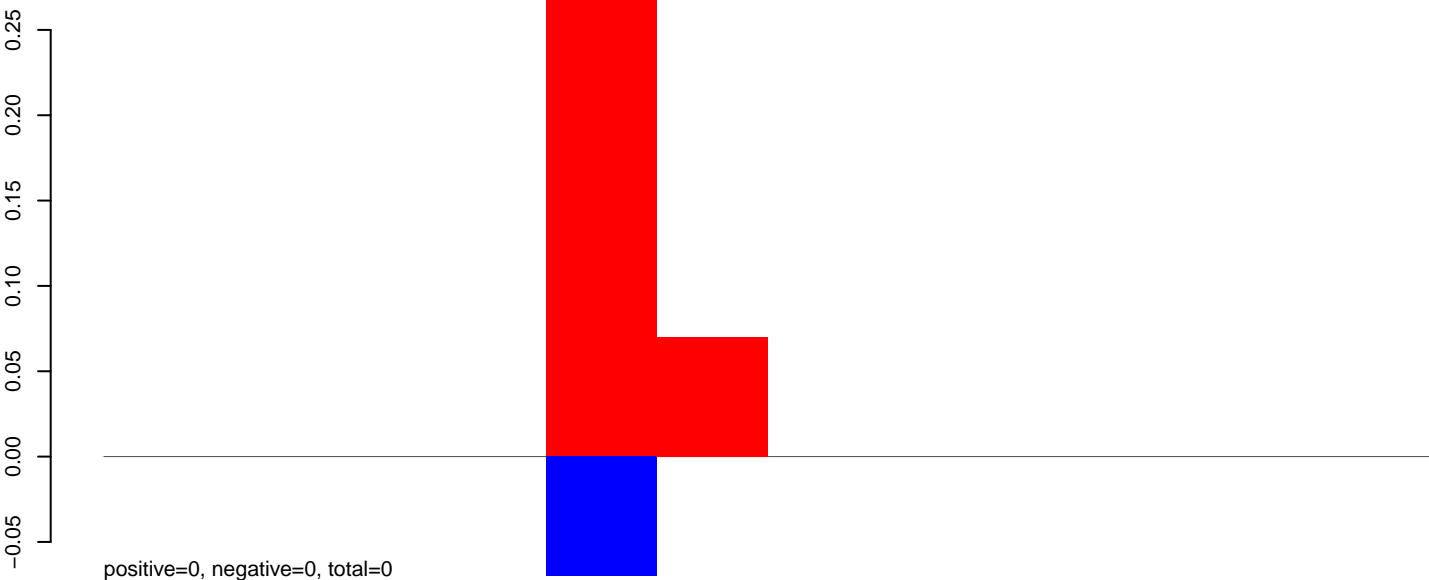


Window size=25, length=429, TE@rnd-5_family-312@LTR_Gypsy@3074:1-429

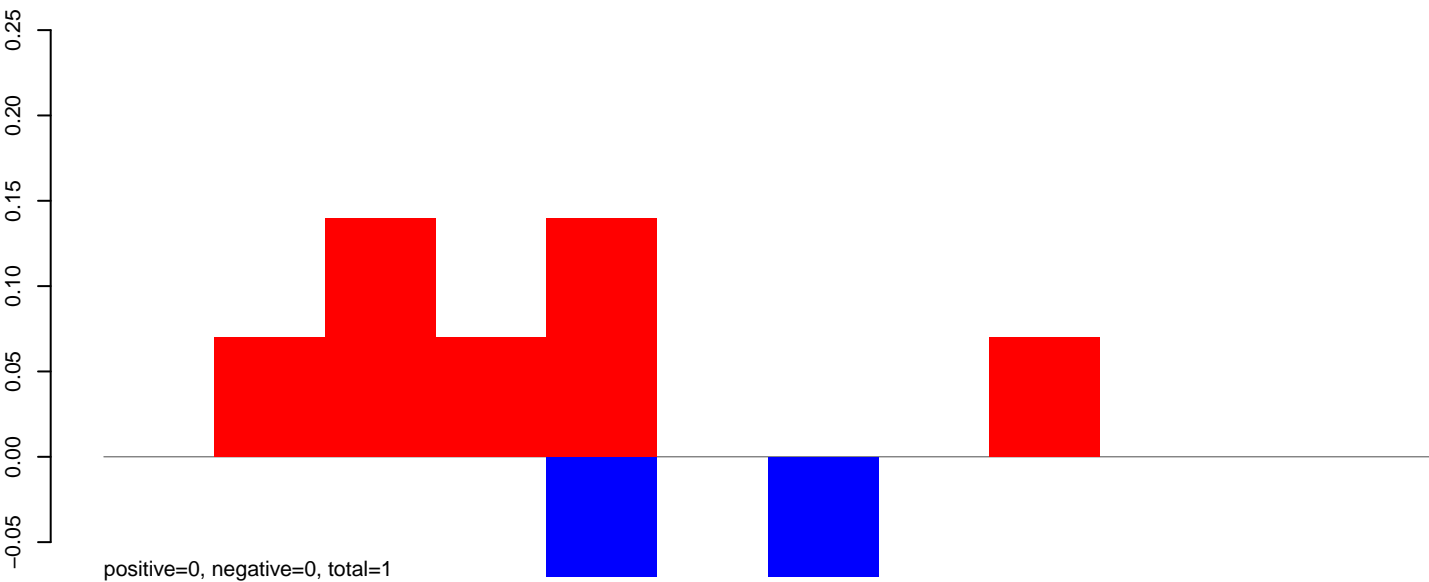


100 200 300 400 500

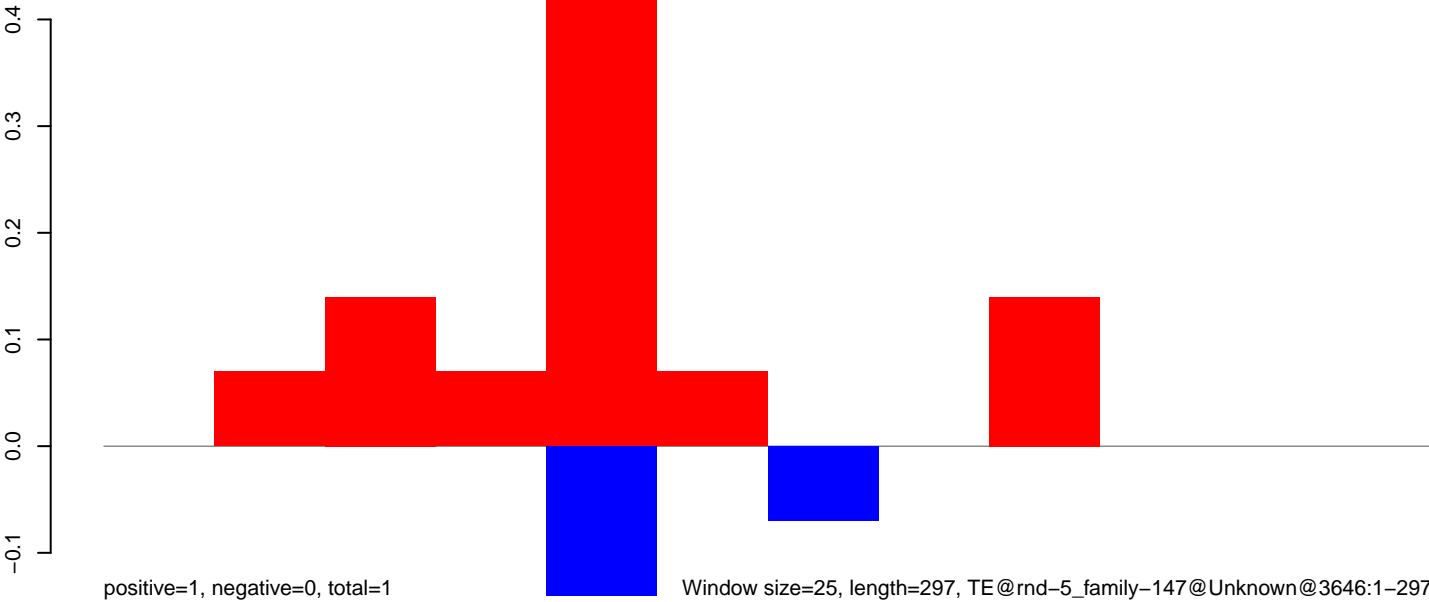
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



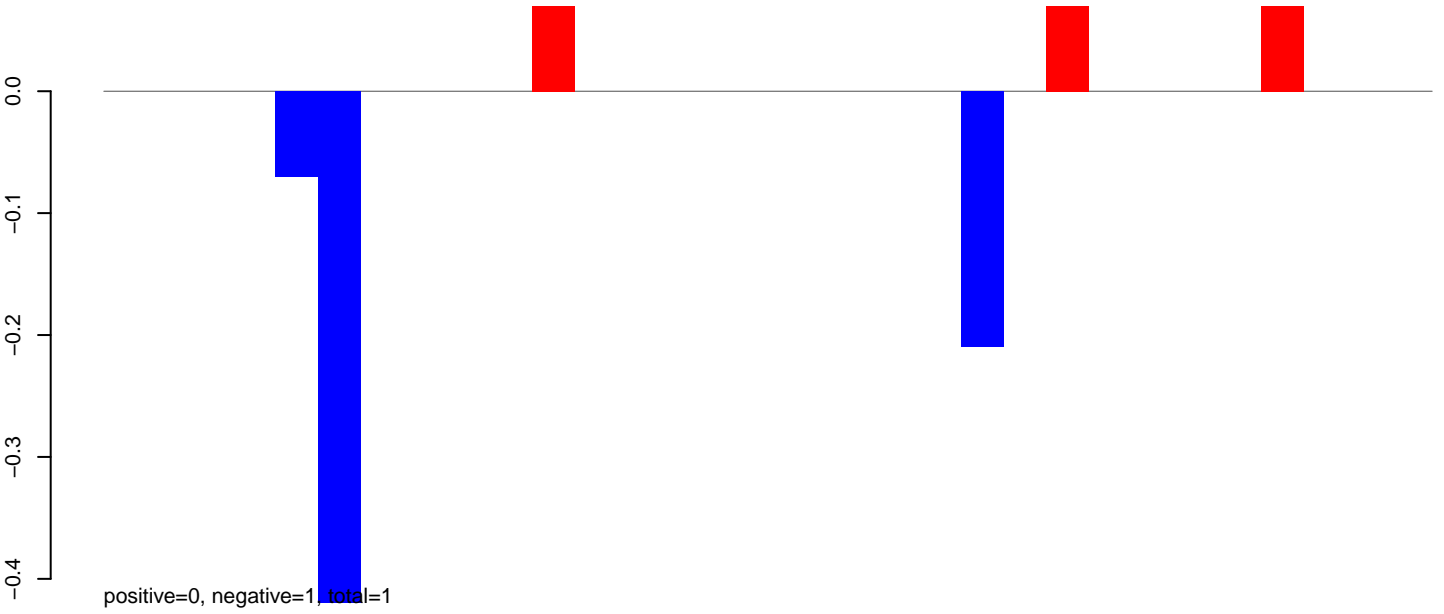
CuQuin_Mcarc_NL.rep



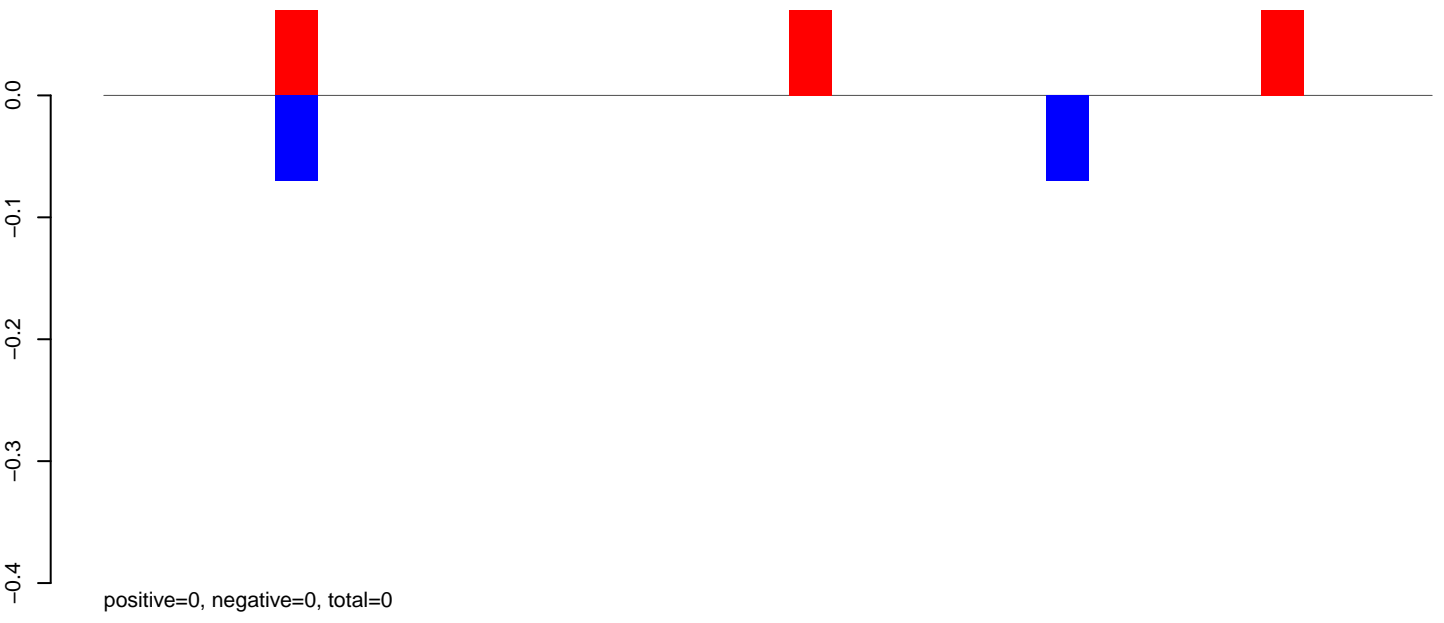
Window size=25, length=297, TE@rnd-5_family-147@Unknown@3646:1-297

50 100 150 200 250 300

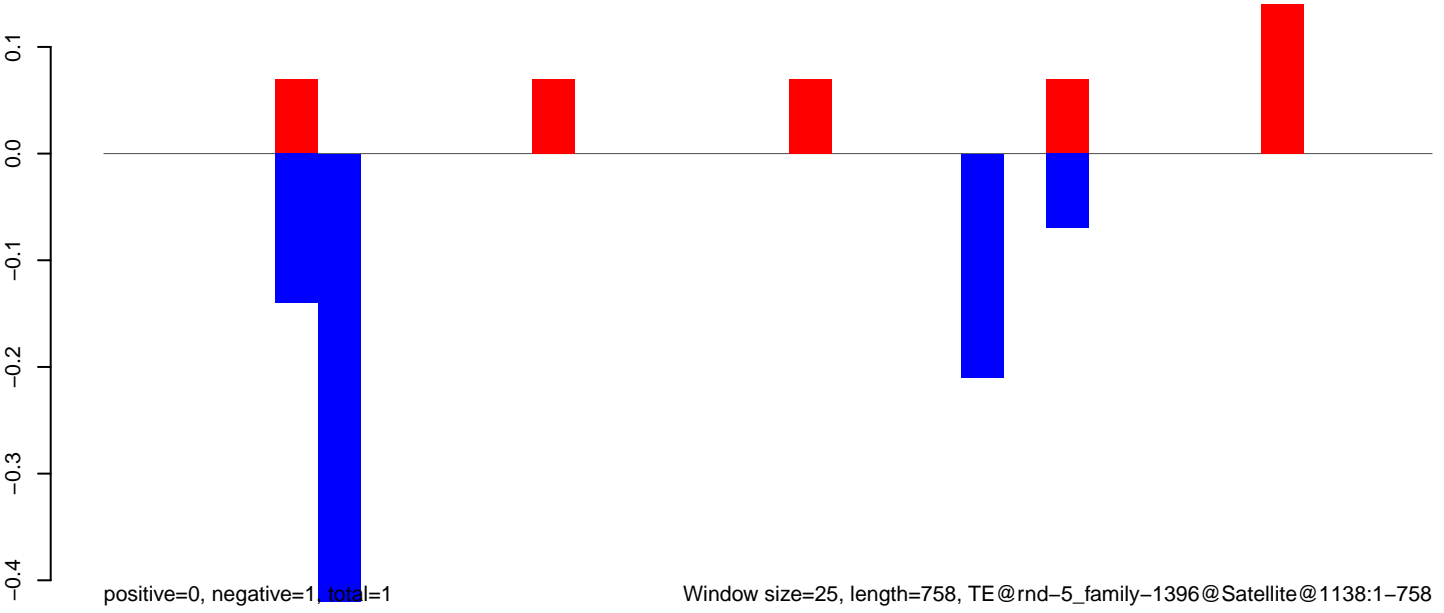
CuQuin_Mcarc_NL.18_23.rep



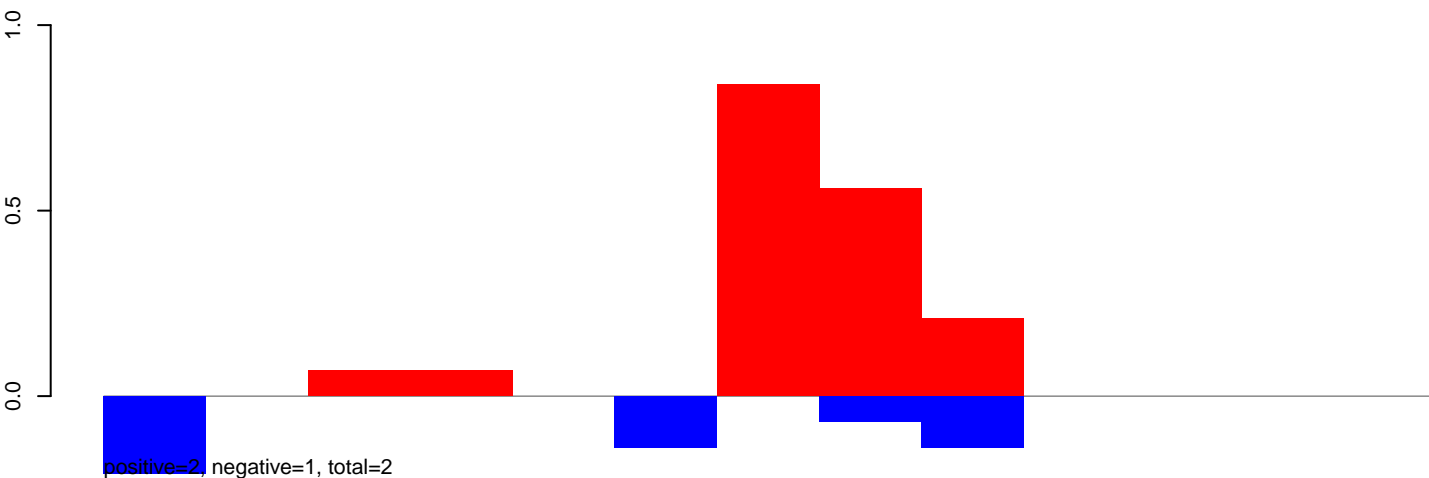
CuQuin_Mcarc_NL.24_35.rep



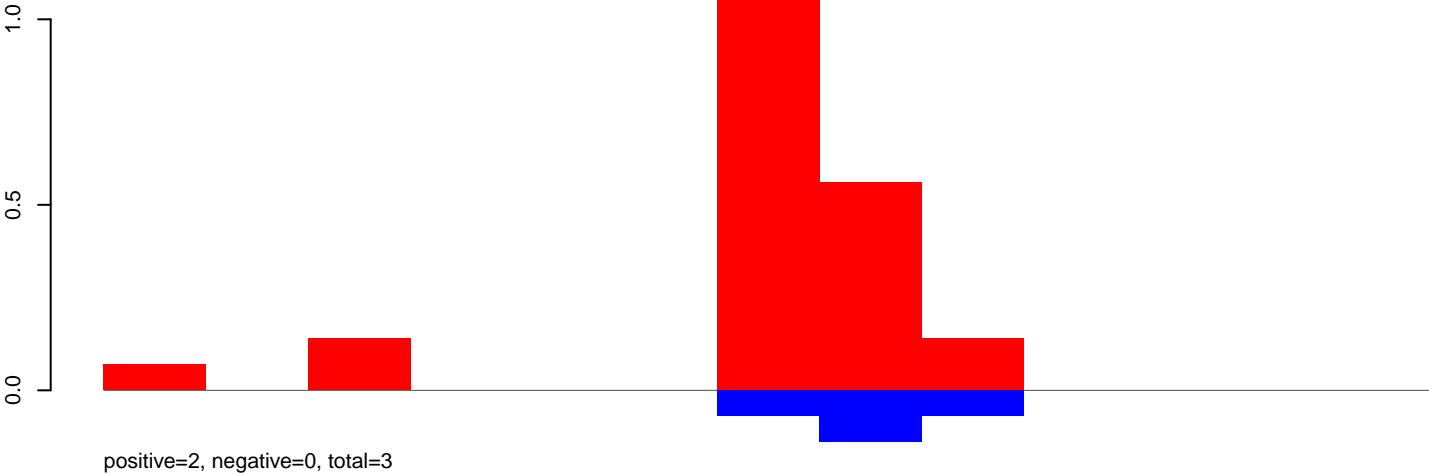
CuQuin_Mcarc_NL.rep



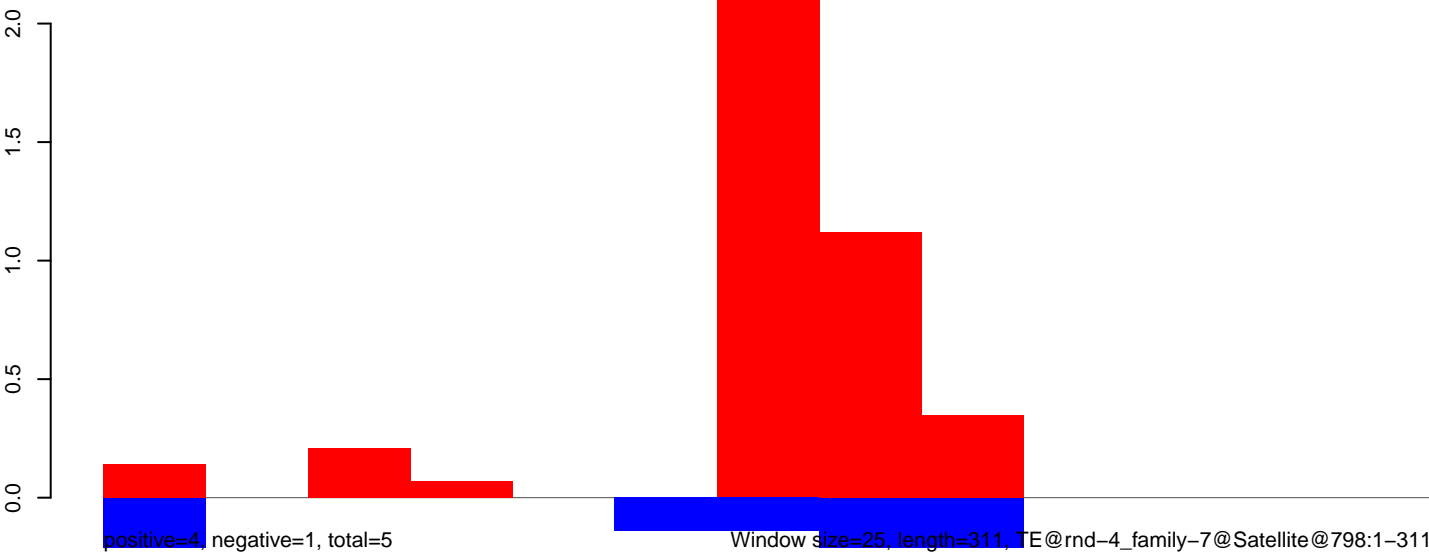
CuQuin_Mcarc_NL.18_23.rep



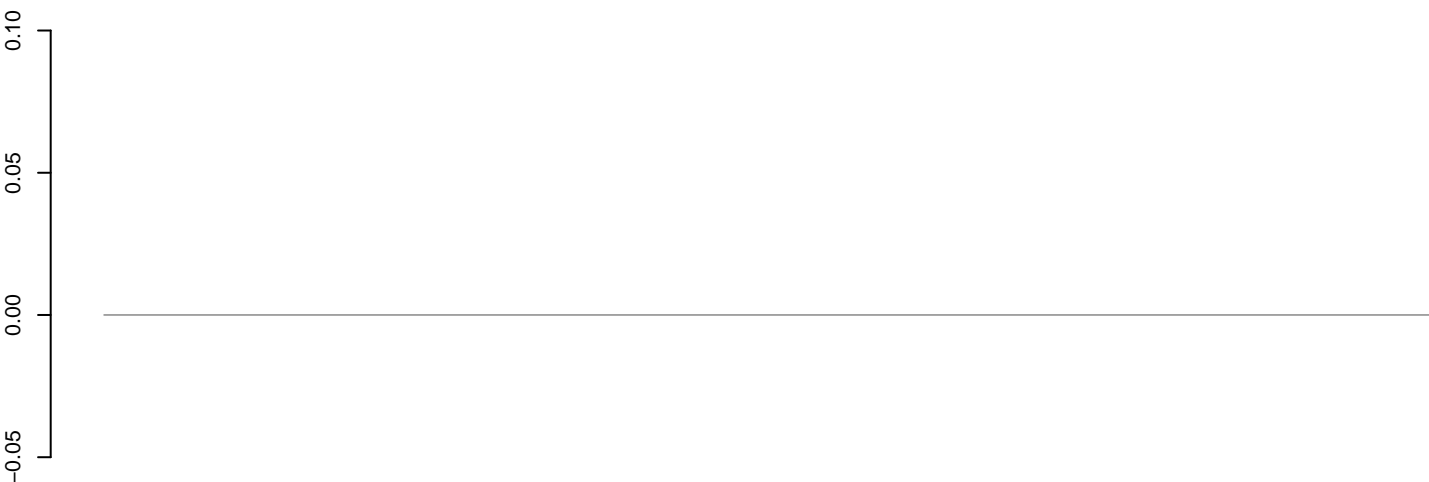
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

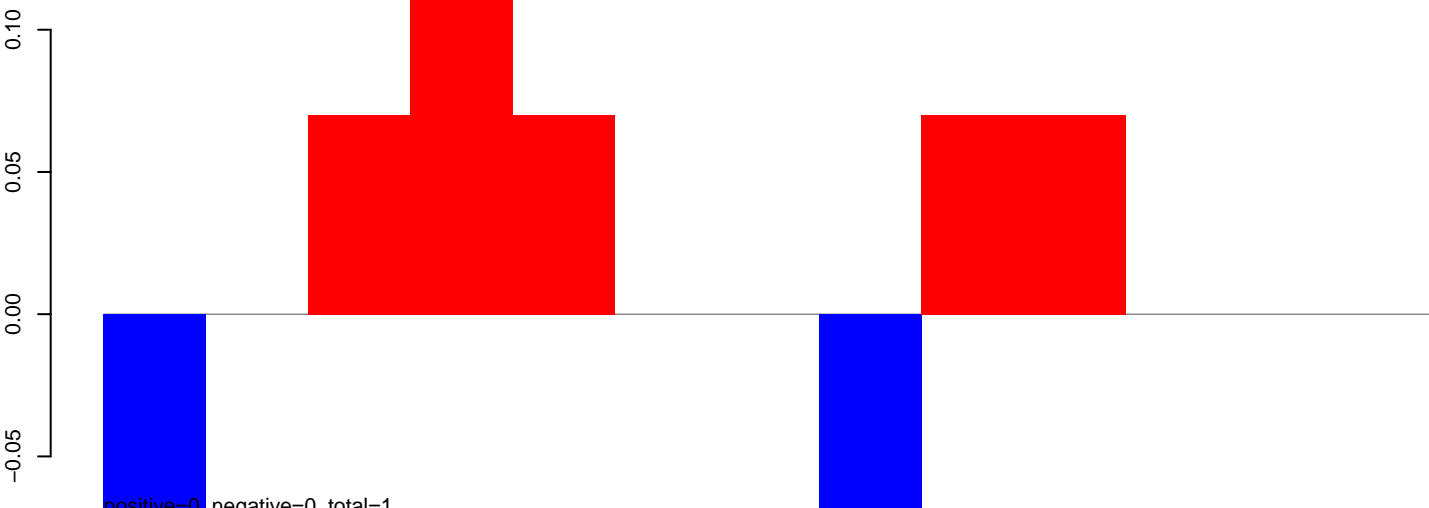


CuQuin_Mcarc_NL.18_23.rep



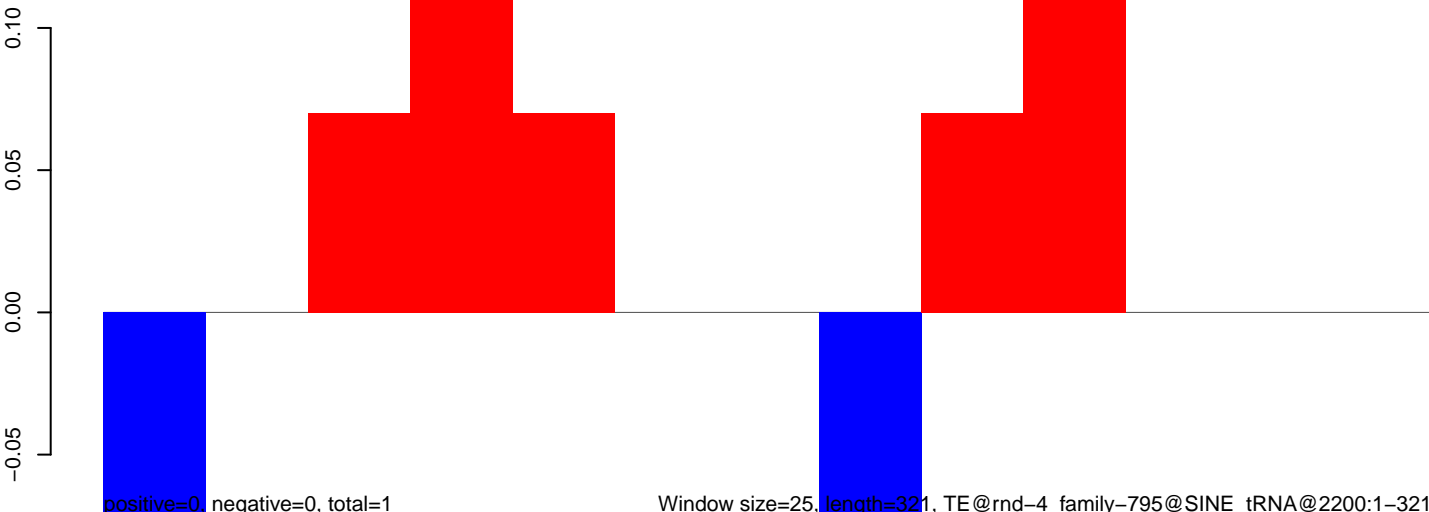
positive=0, negative=0, total=0

CuQuin_Mcarc_NL.24_35.rep



positive=0, negative=0, total=1

CuQuin_Mcarc_NL.rep

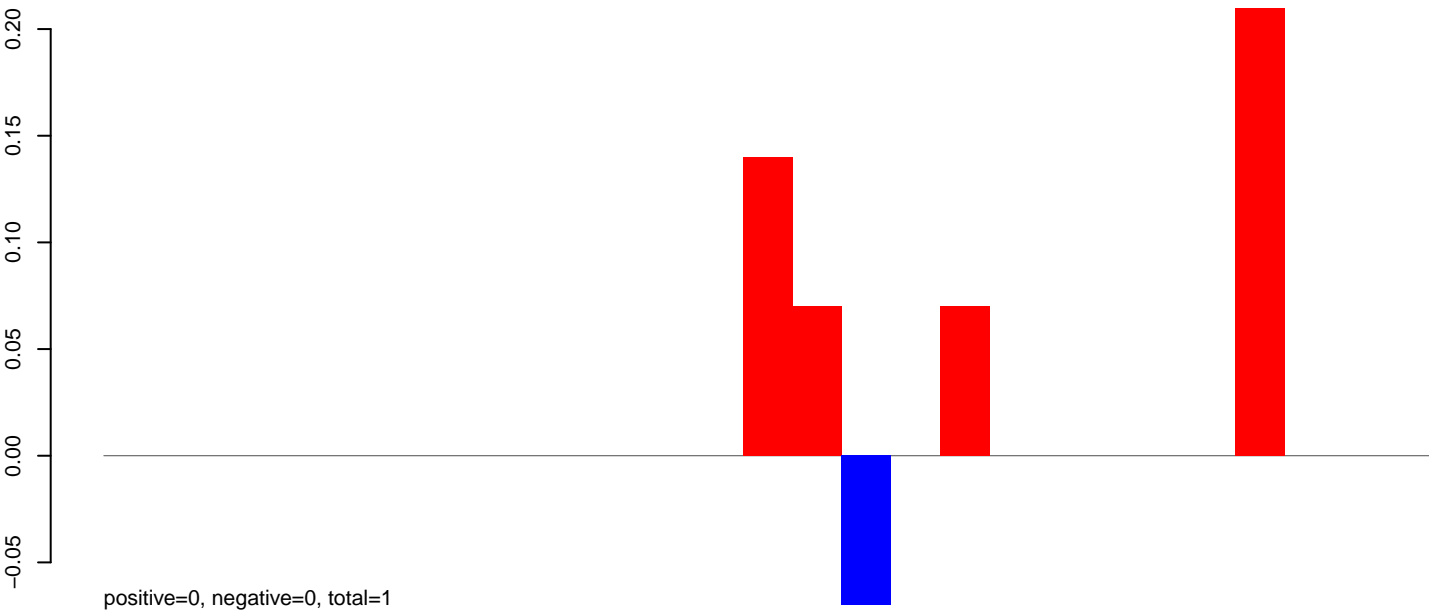


positive=0, negative=0, total=1

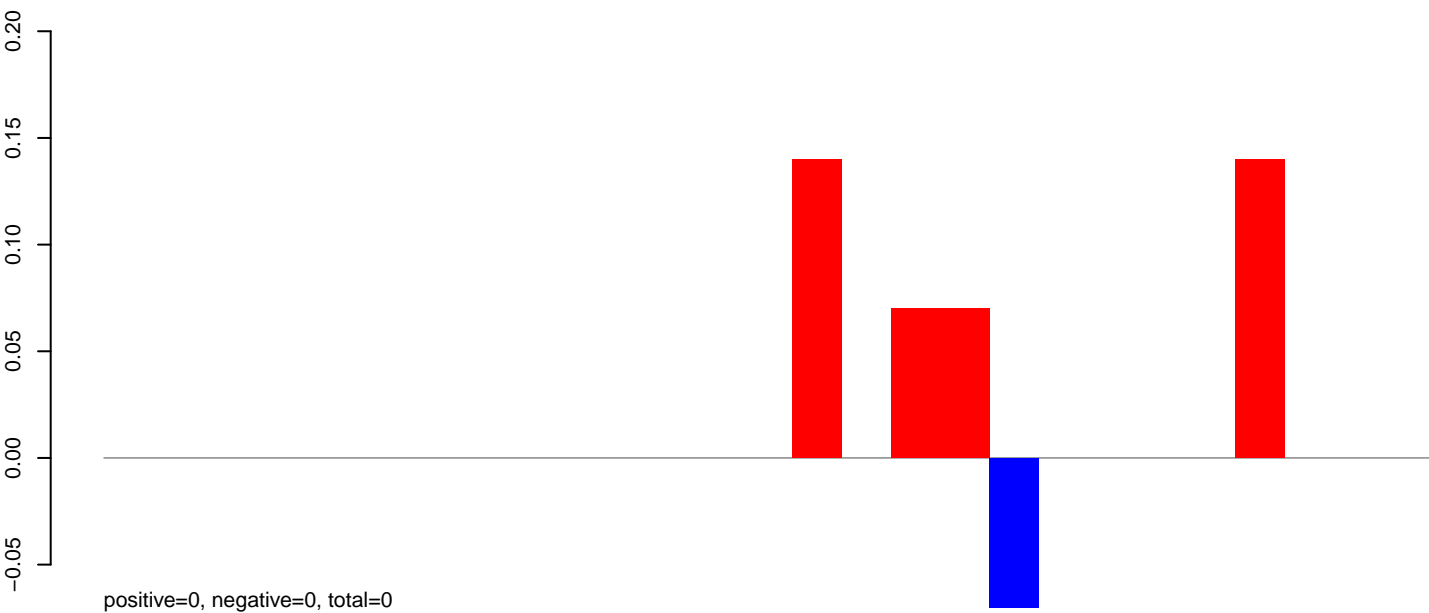
Window size=25, length=321, TE@rnd-4_family-795@SINE_tRNA@2200:1-321

50 100 150 200 250 300 350

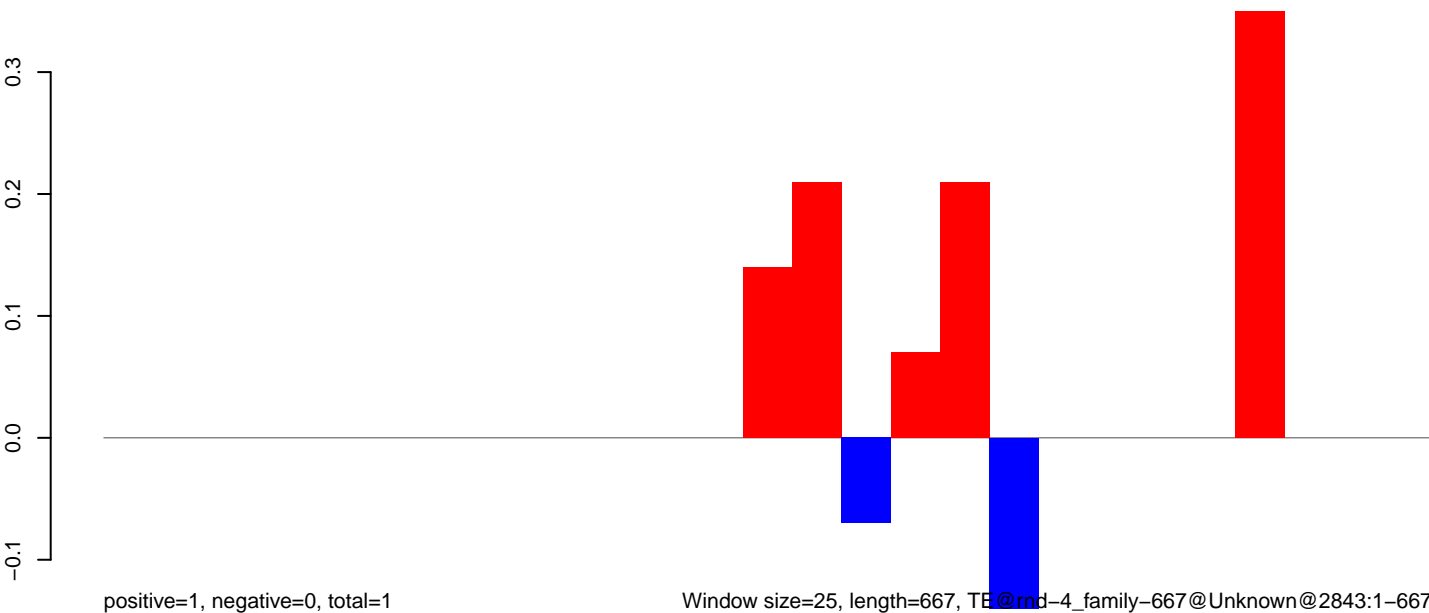
CuQuin_Mcarc_NL.18_23.rep



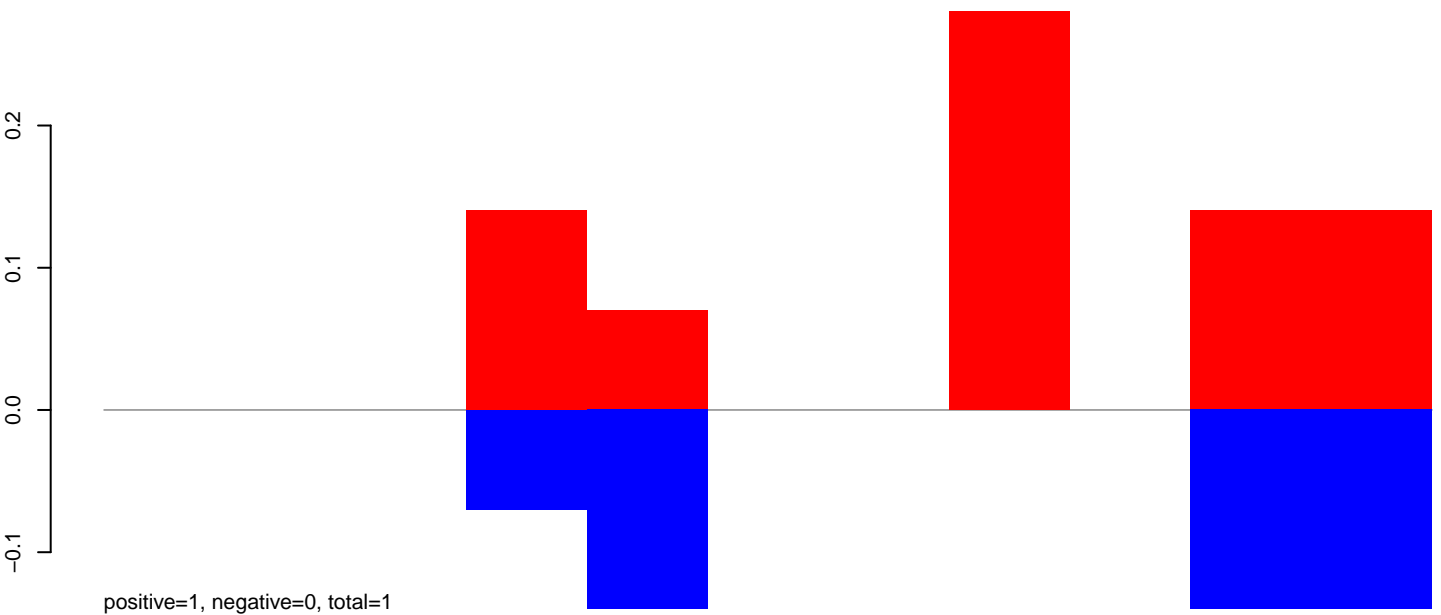
CuQuin_Mcarc_NL.24_35.rep



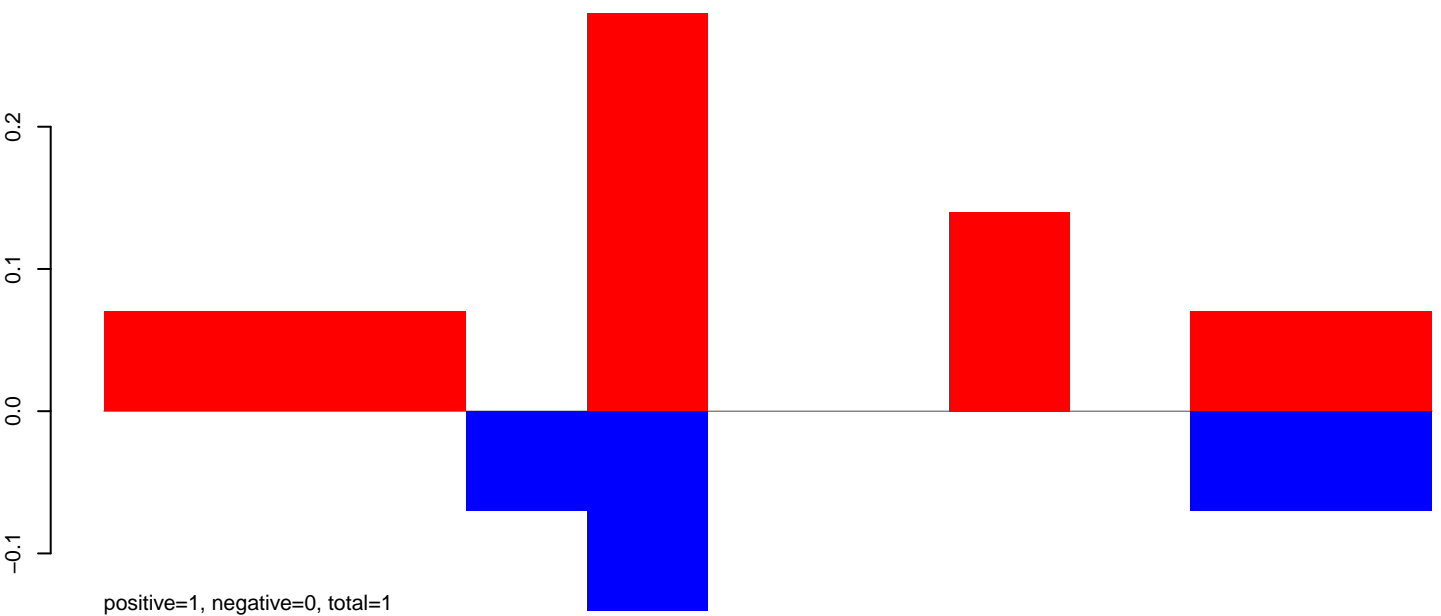
CuQuin_Mcarc_NL.rep



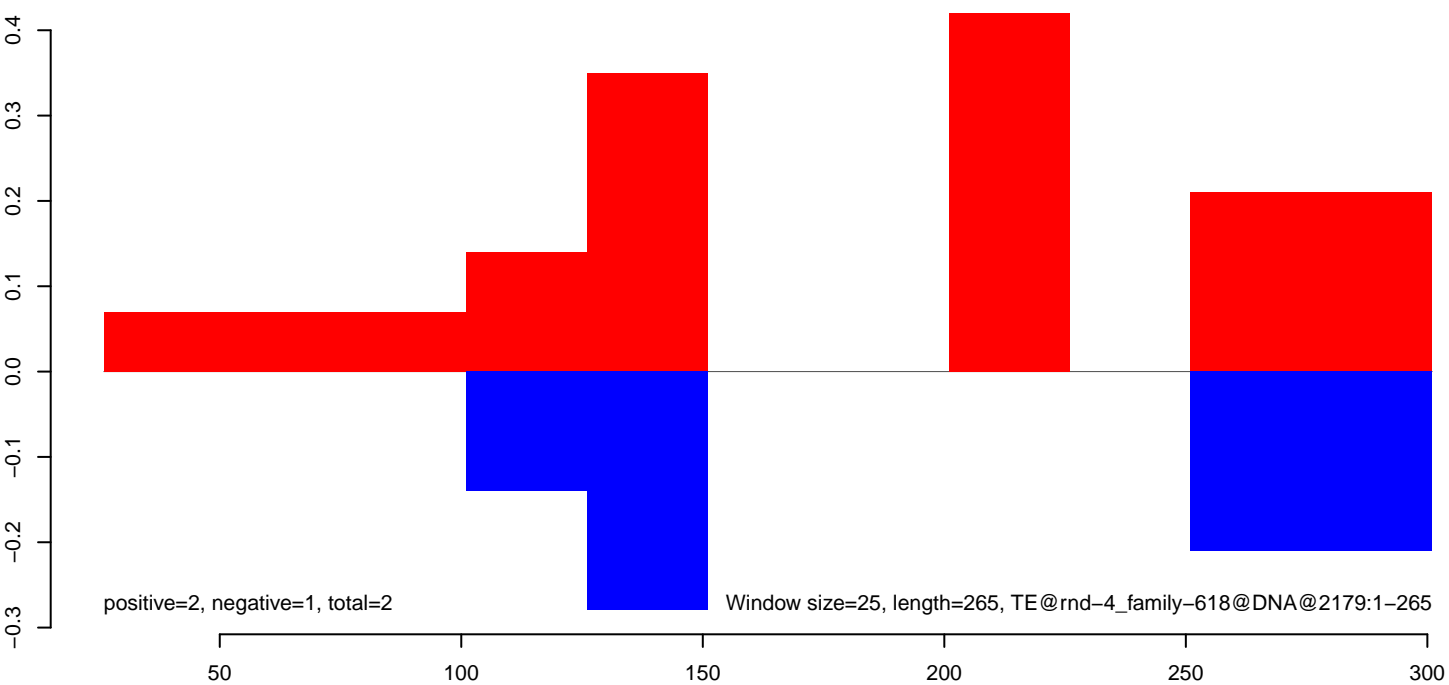
CuQuin_Mcarc_NL.18_23.rep



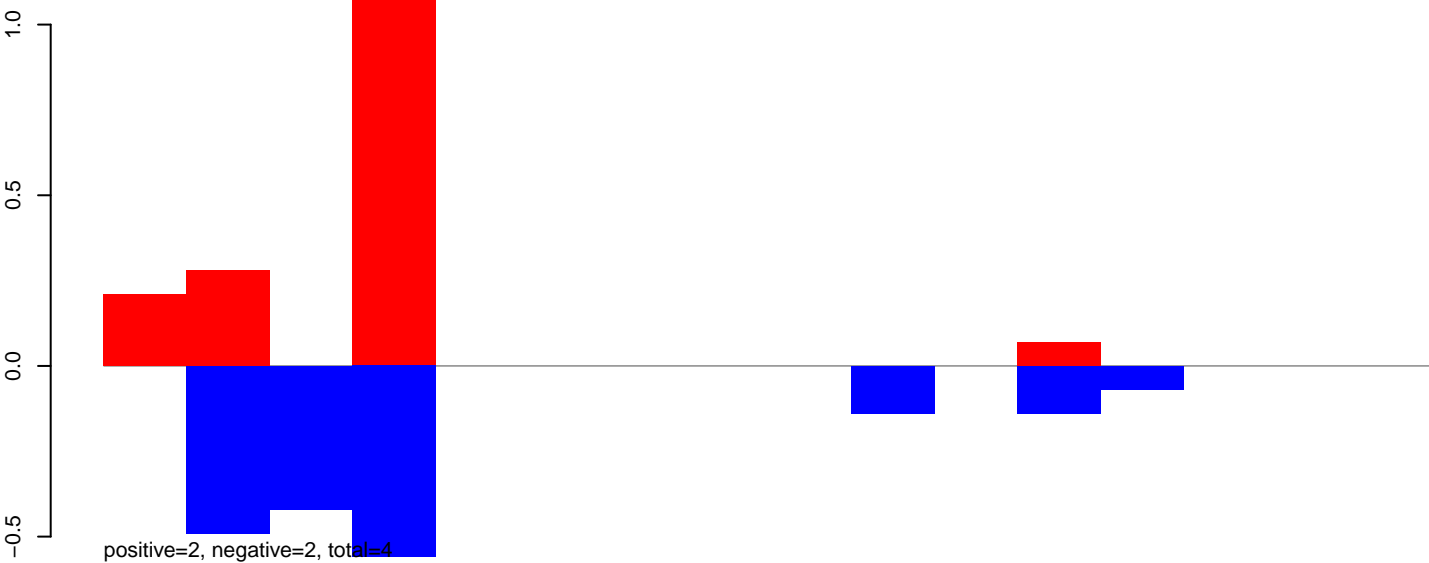
CuQuin_Mcarc_NL.24_35.rep



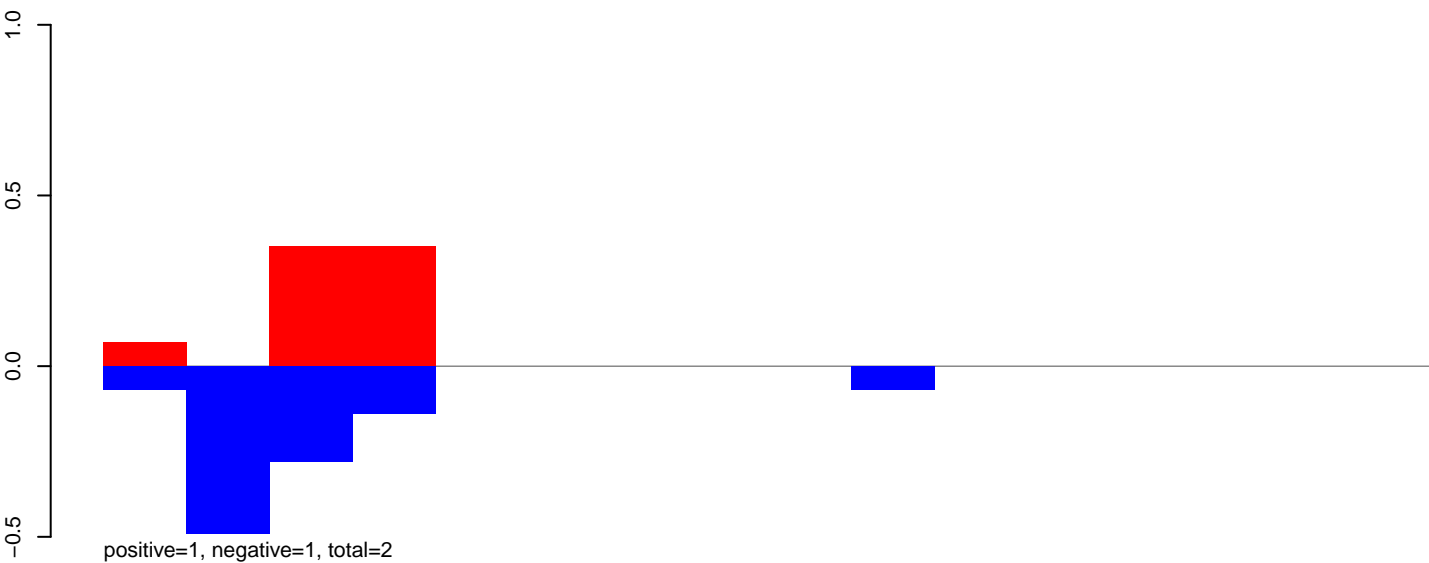
CuQuin_Mcarc_NL.rep



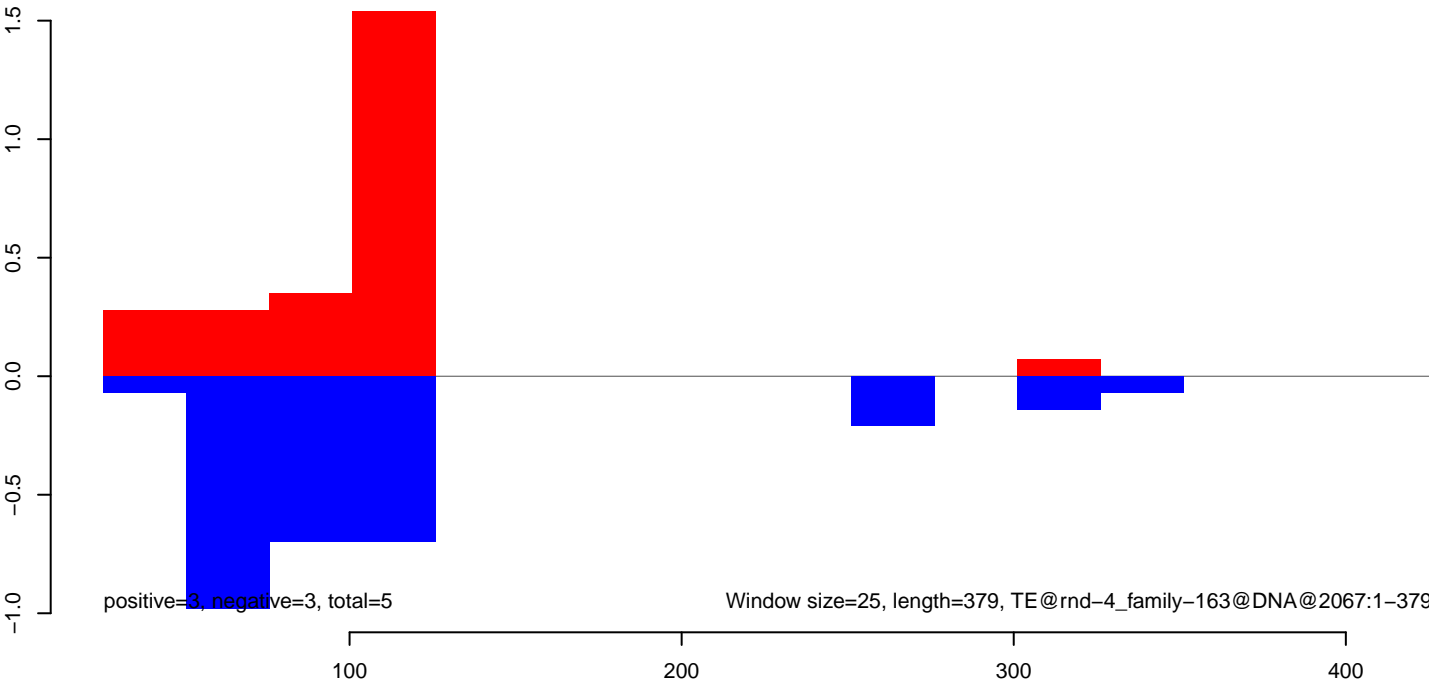
CuQuin_Mcarc_NL.18_23.rep



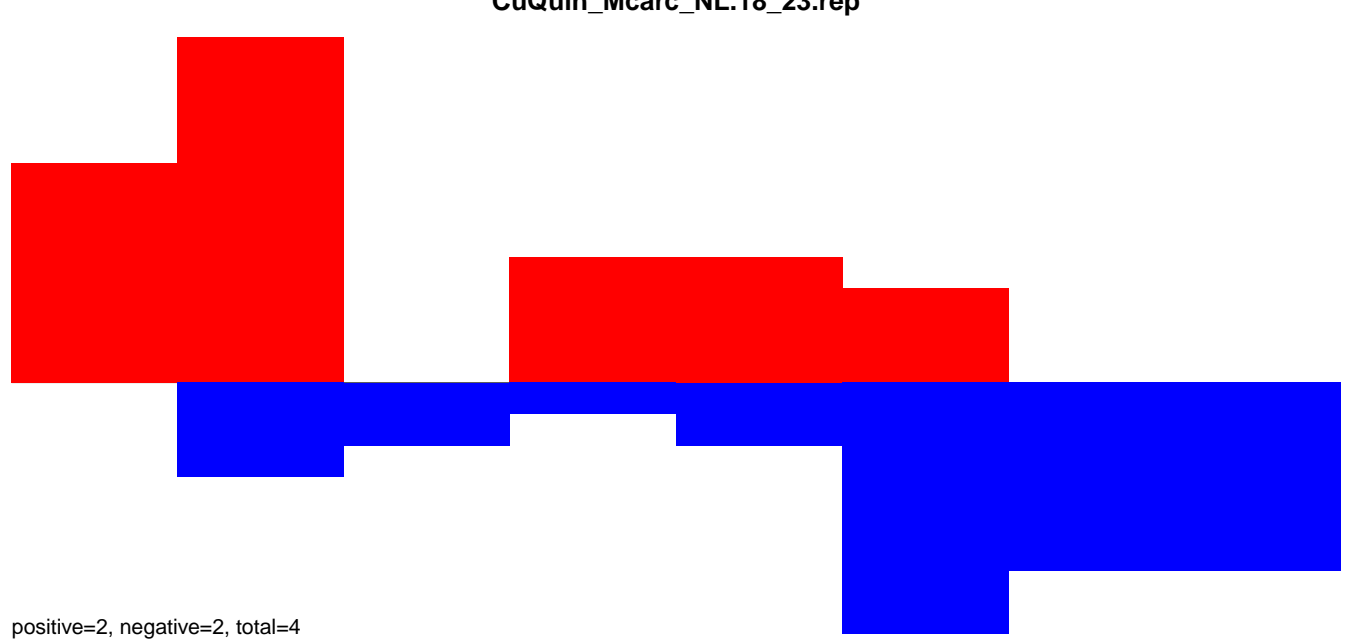
CuQuin_Mcarc_NL.24_35.rep



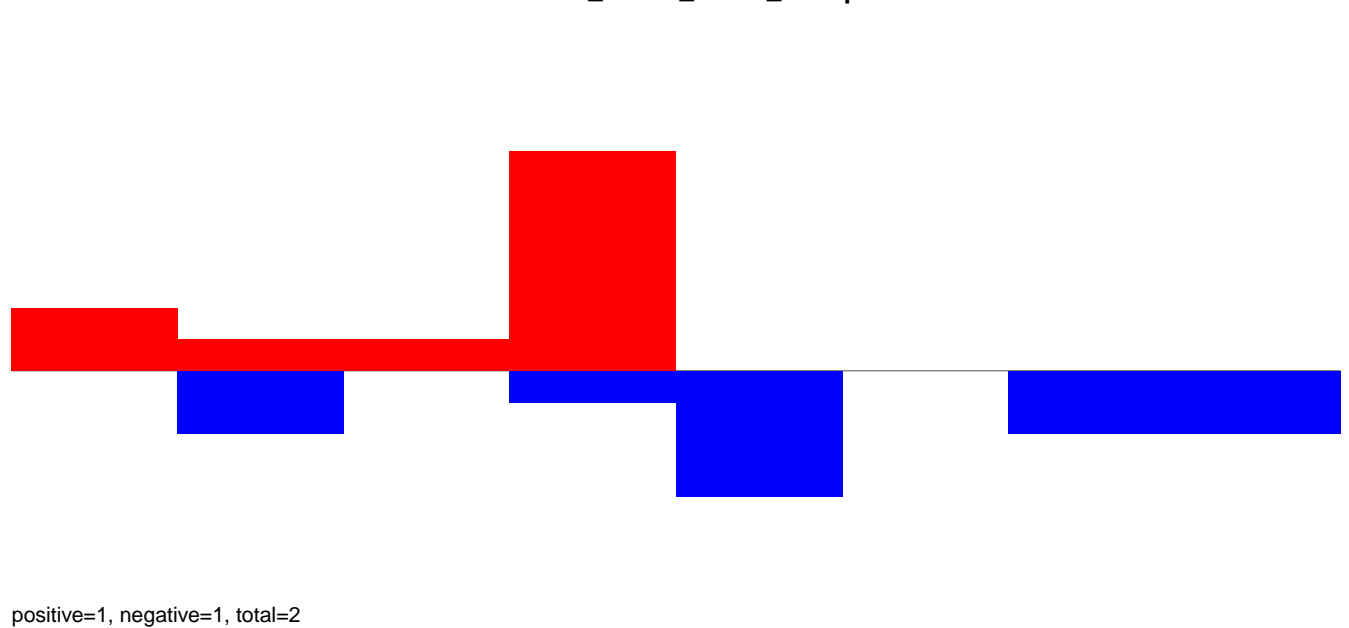
CuQuin_Mcarc_NL.rep



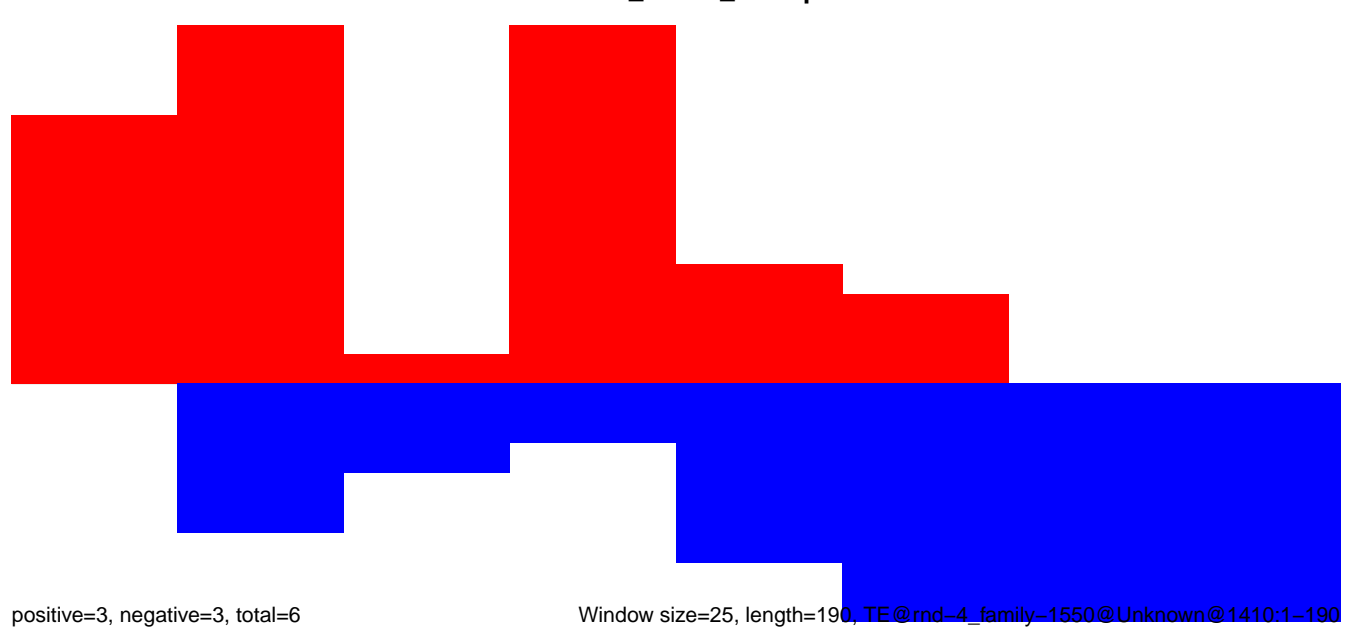
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

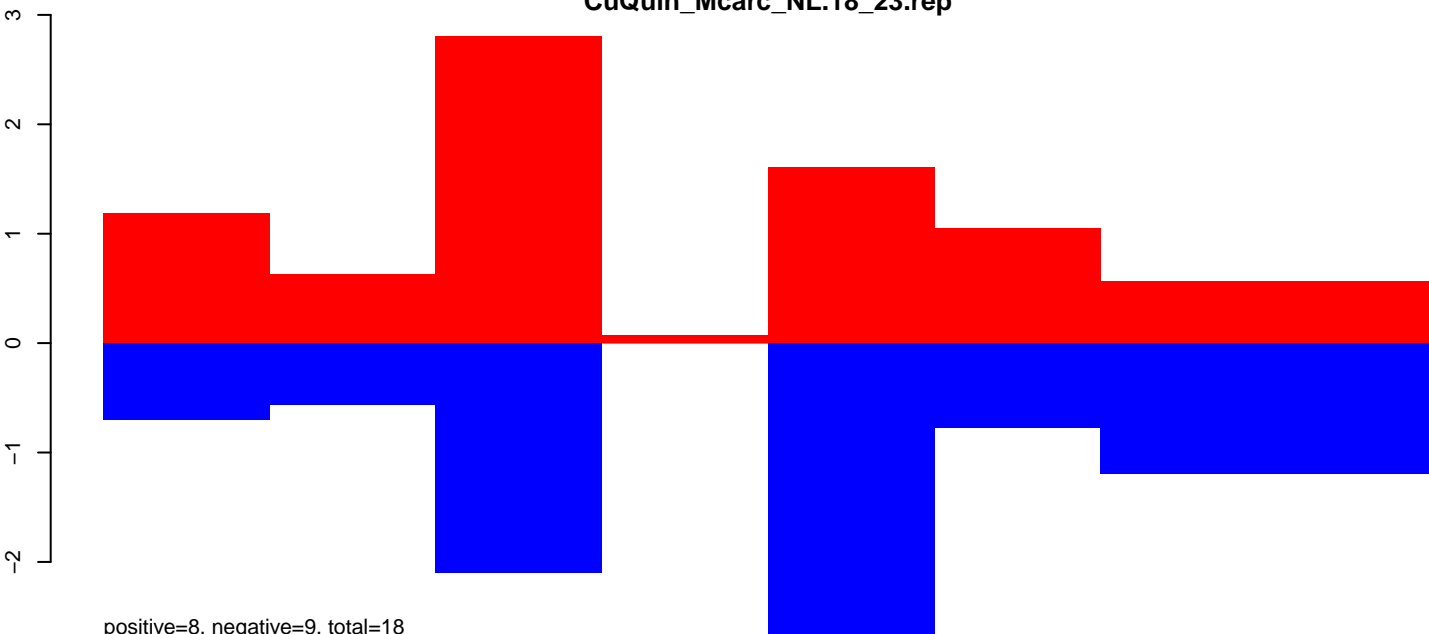


CuQuin_Mcarc_NL.rep



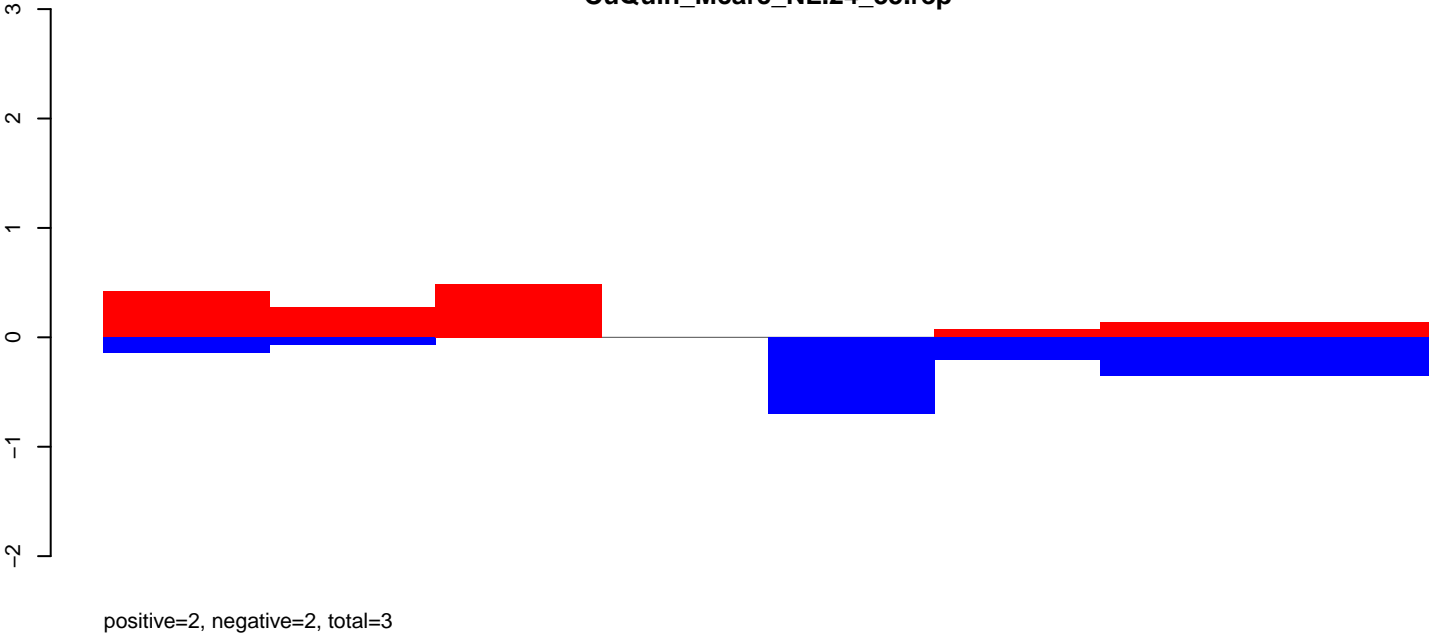
Window size=25, length=190, TE@rnd-4_family-1550@Unknown@1410:1-190

CuQuin_Mcarc_NL.18_23.rep



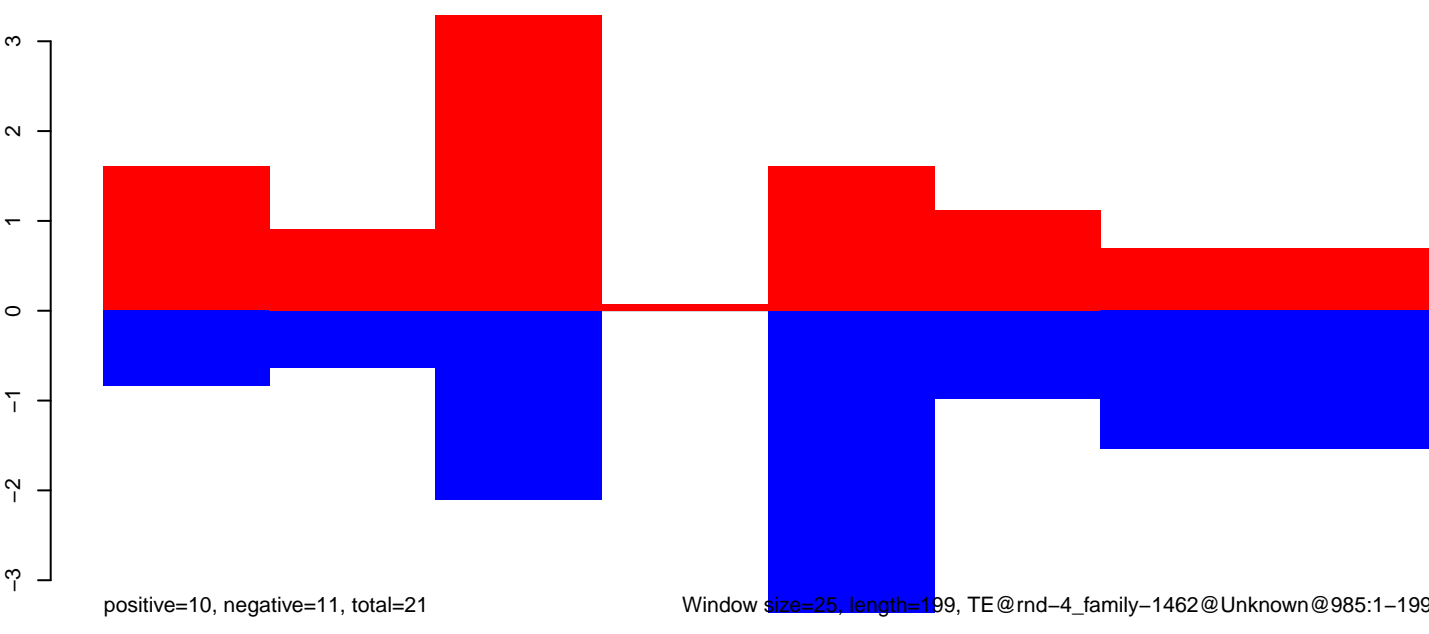
positive=8, negative=9, total=18

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=2, total=3

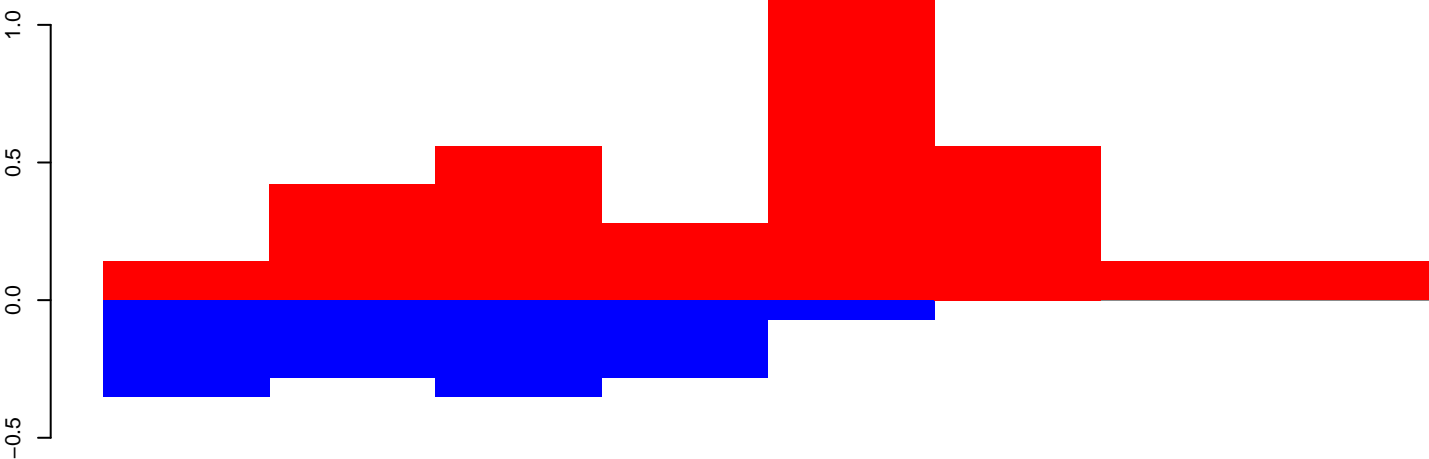
CuQuin_Mcarc_NL.rep



positive=10, negative=11, total=21

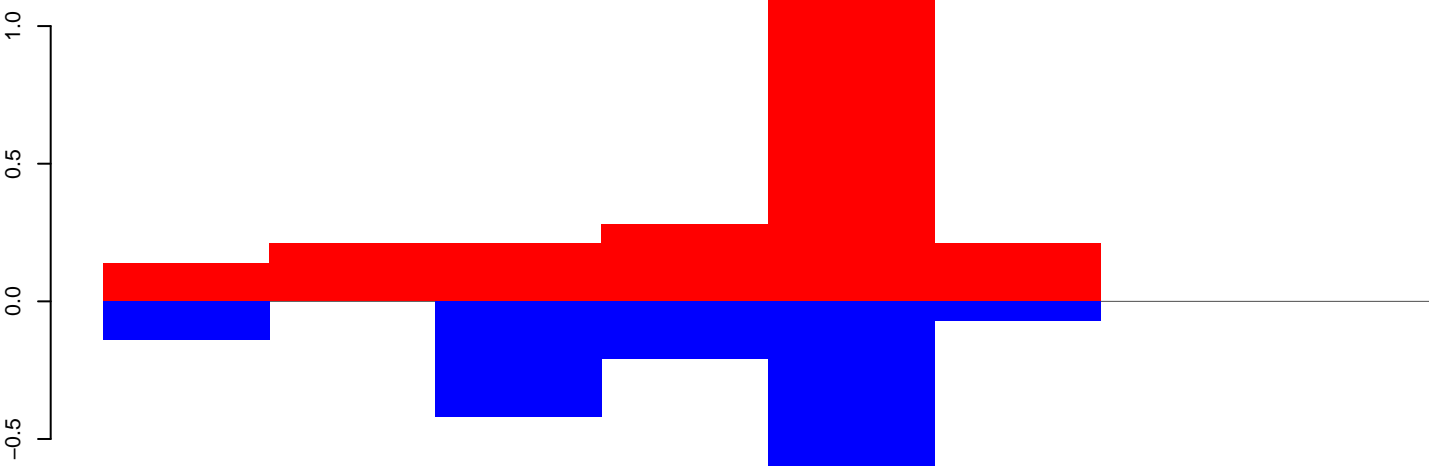
Window size=25, length=199, TE@rnd

CuQuin_Mcarc_NL.18_23.rep



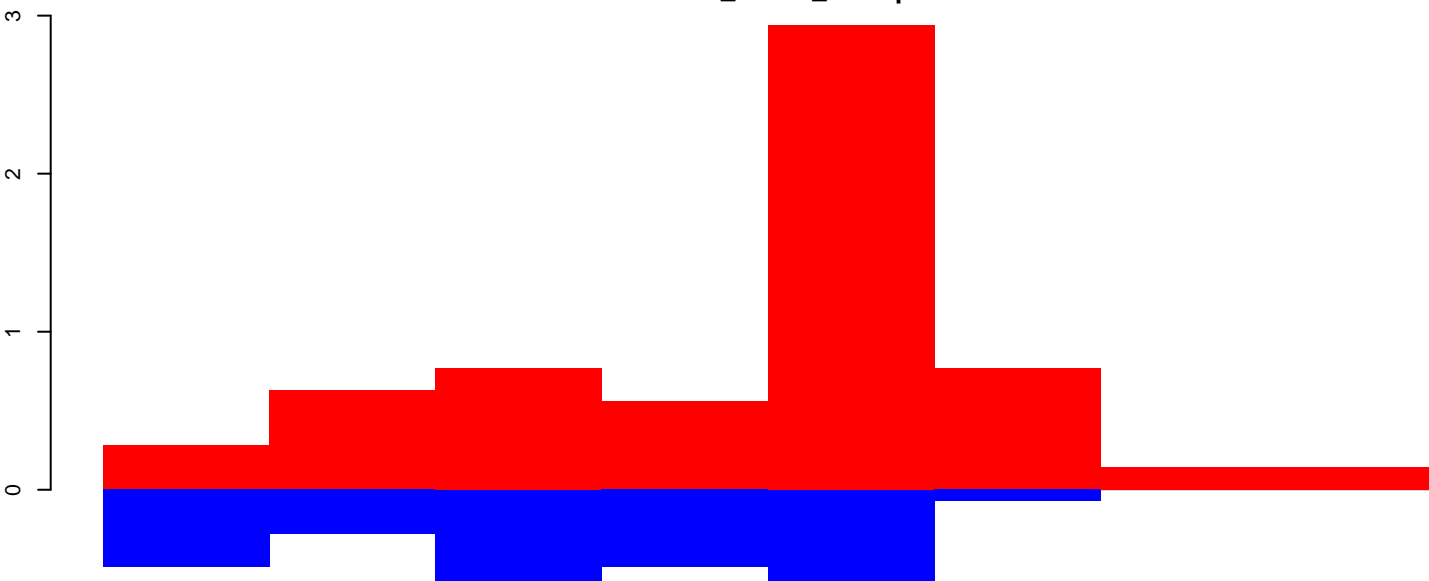
positive=4, negative=1, total=5

CuQuin_Mcarc_NL.24_35.rep



positive=2, negative=2, total=4

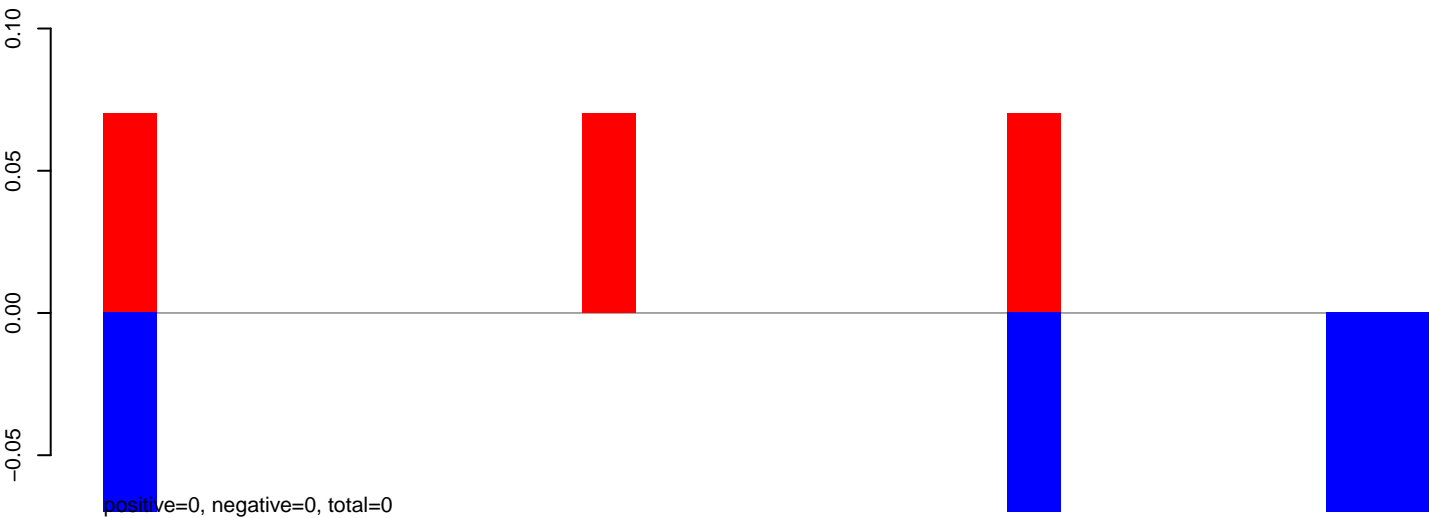
CuQuin_Mcarc_NL.rep



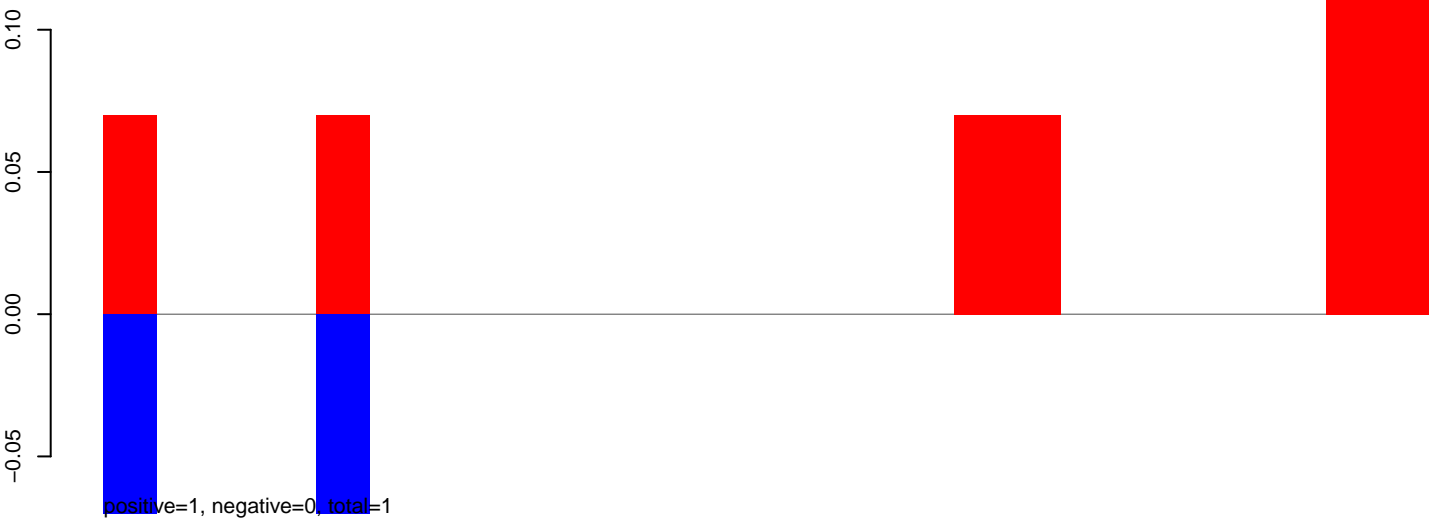
positive=6, negative=3, total=9

Window size=25, length=185, TE@rnd-3_family-1876@Unknown@727:1-185

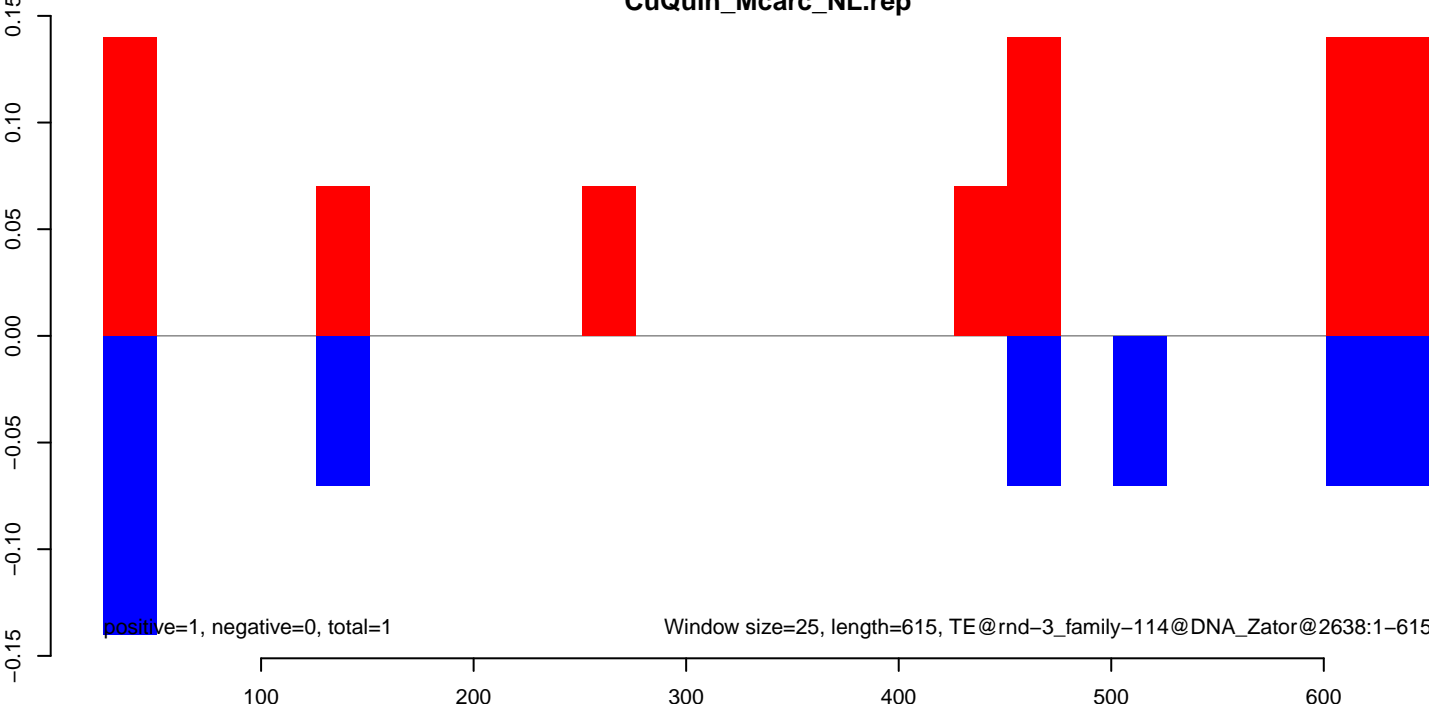
CuQuin_Mcarc_NL.18_23.rep



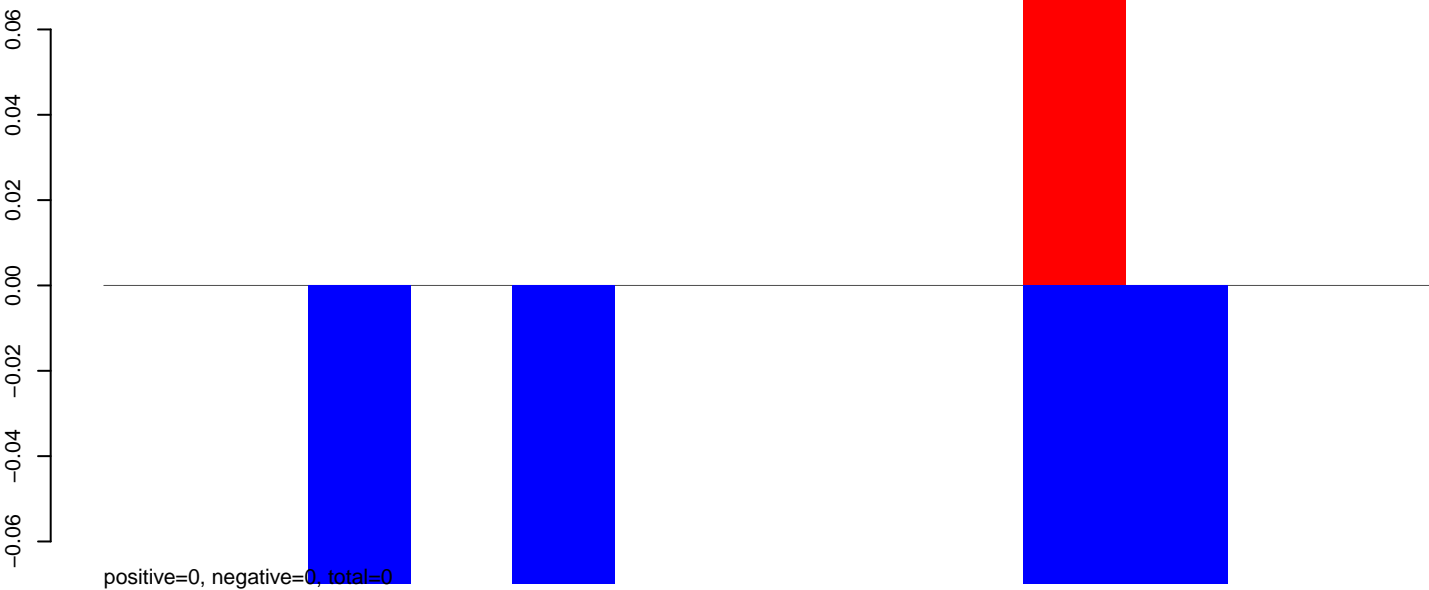
CuQuin_Mcarc_NL.24_35.rep



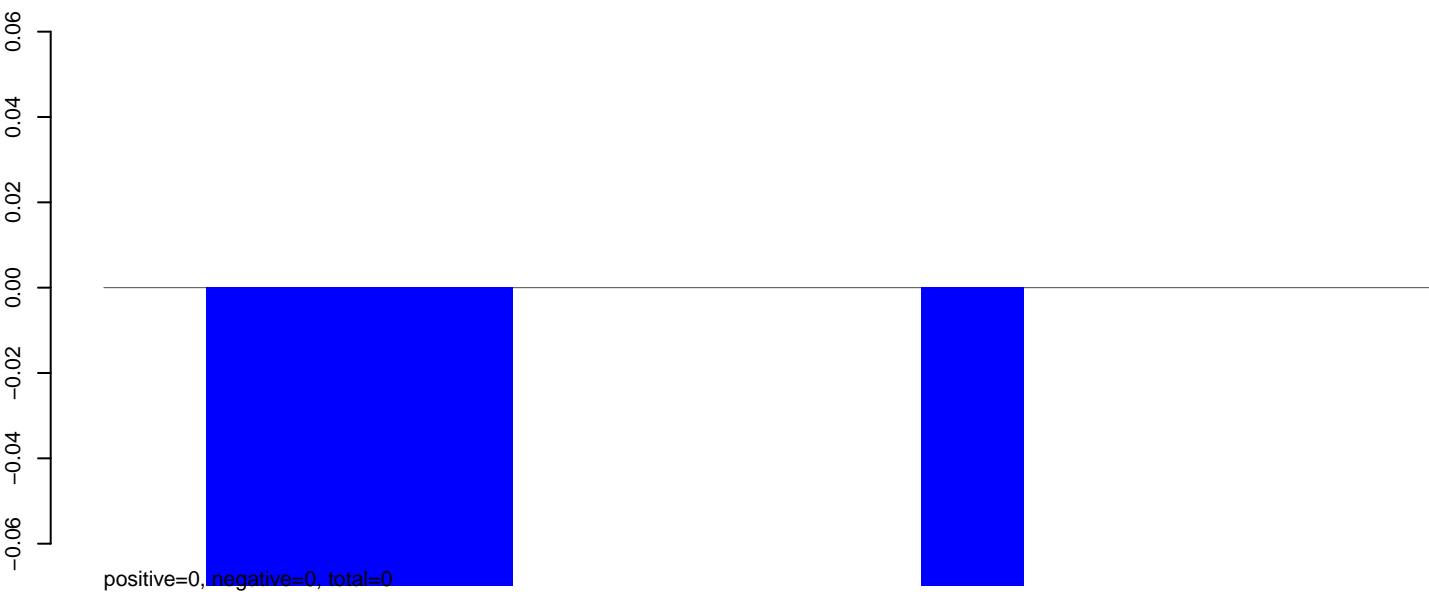
CuQuin_Mcarc_NL.rep



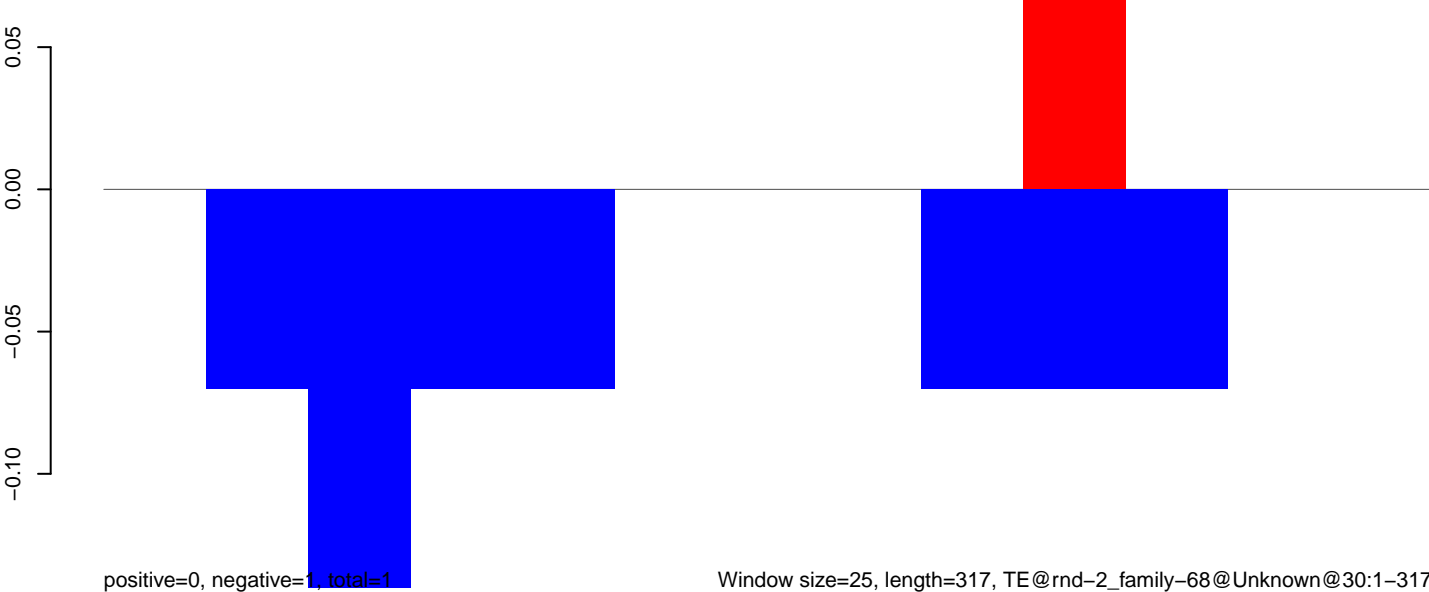
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



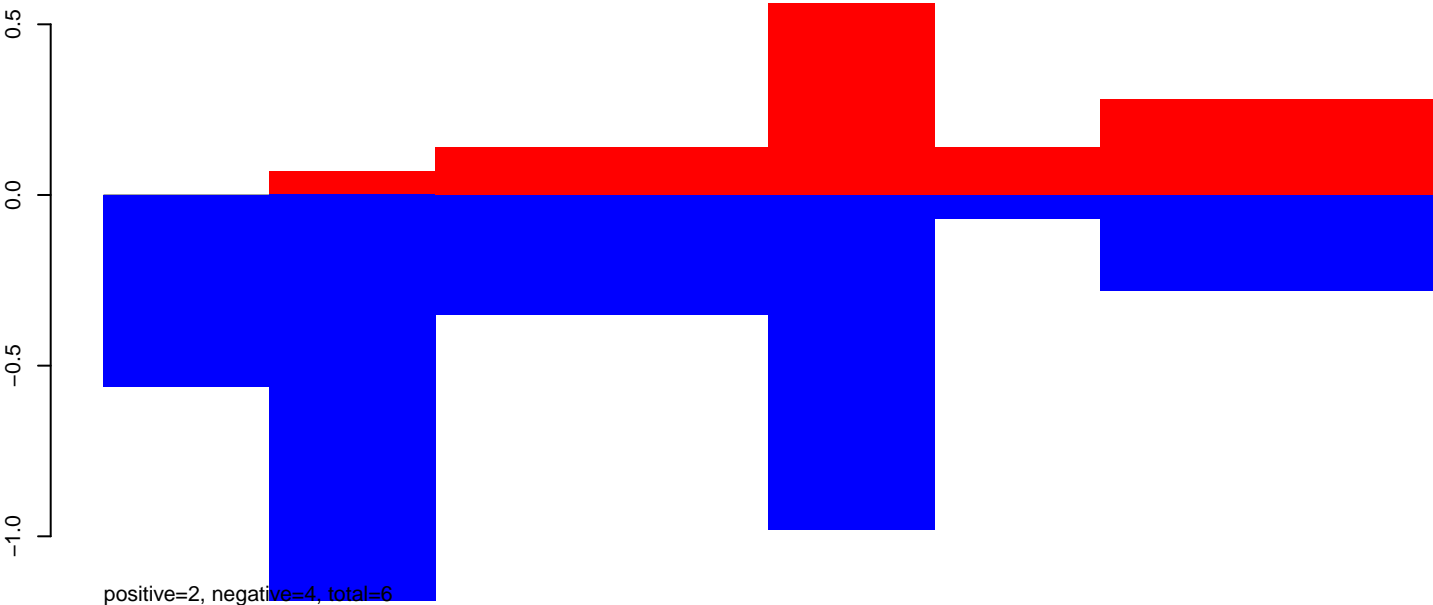
CuQuin_Mcarc_NL.rep



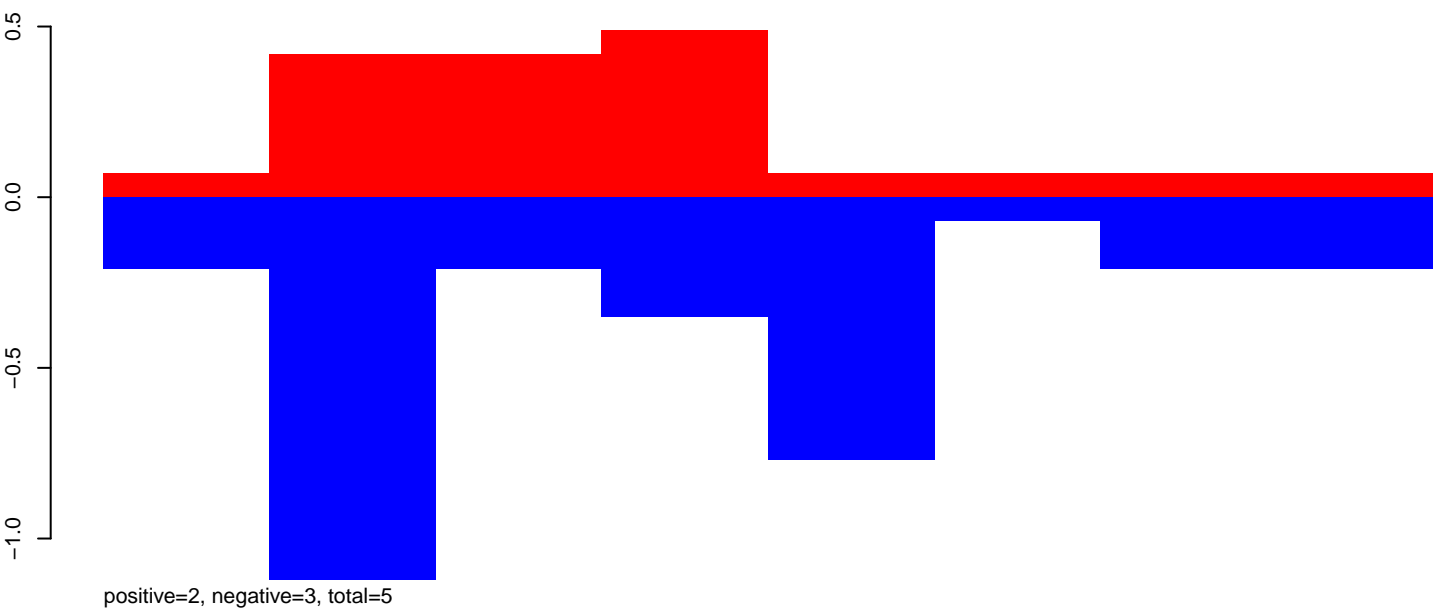
Window size=25, length=317, TE@rnd-2_family-68@Unknown@30:1-317

50 100 150 200 250 300 350

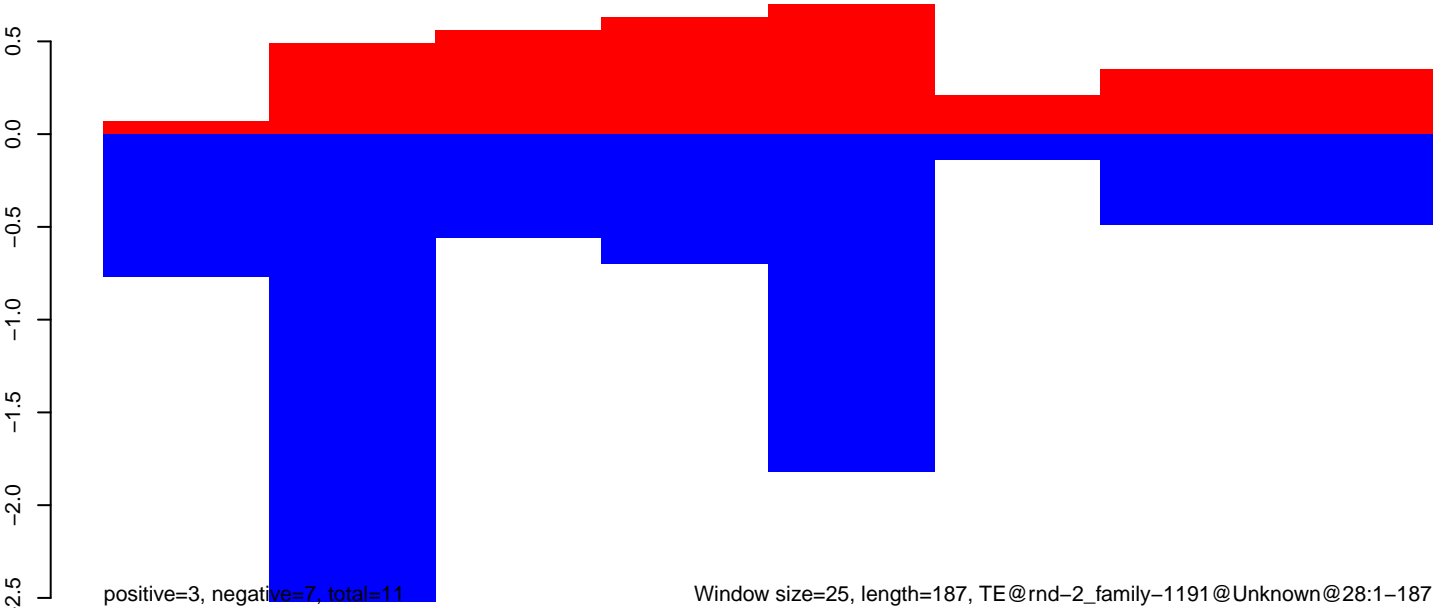
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=187, TE@rnd-2_family-1191@Unknown@28:1-187

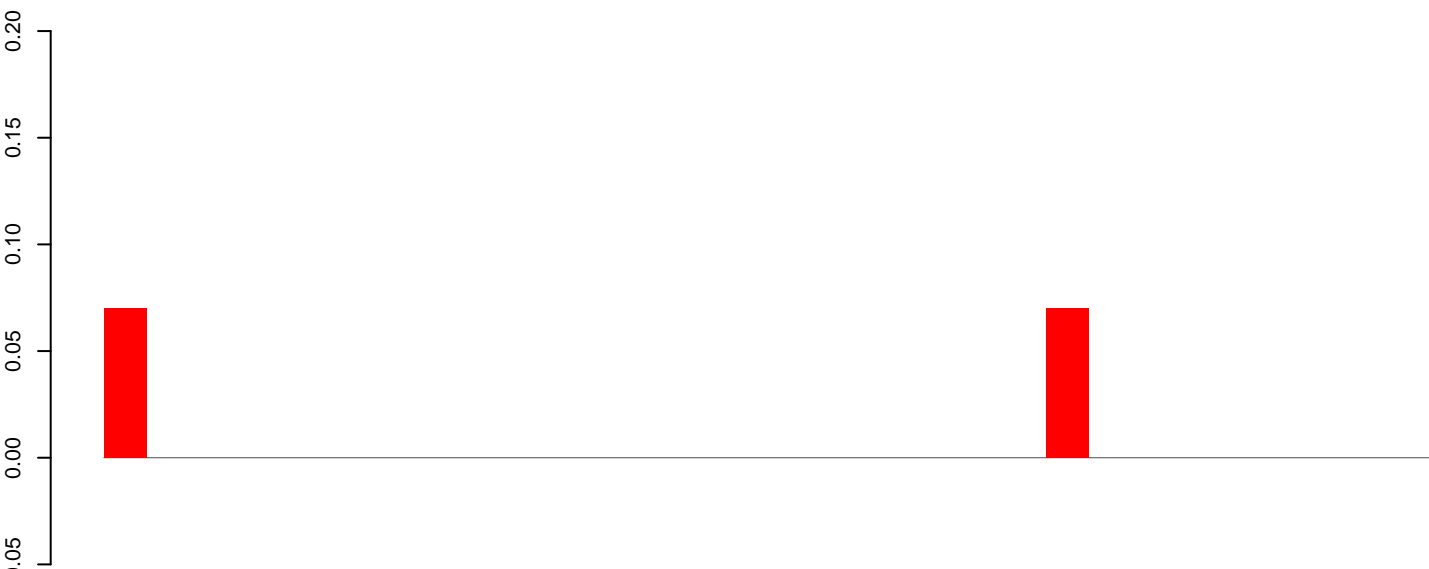
50

100

150

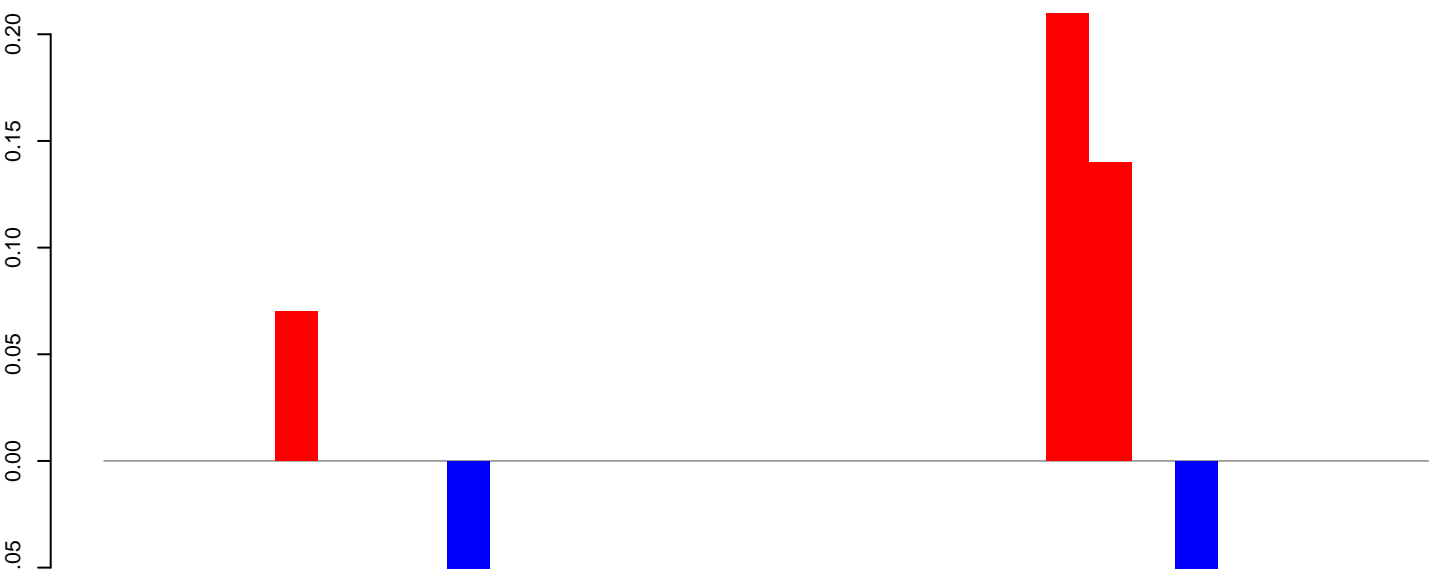
200

CuQuin_Mcarc_NL.18_23.rep



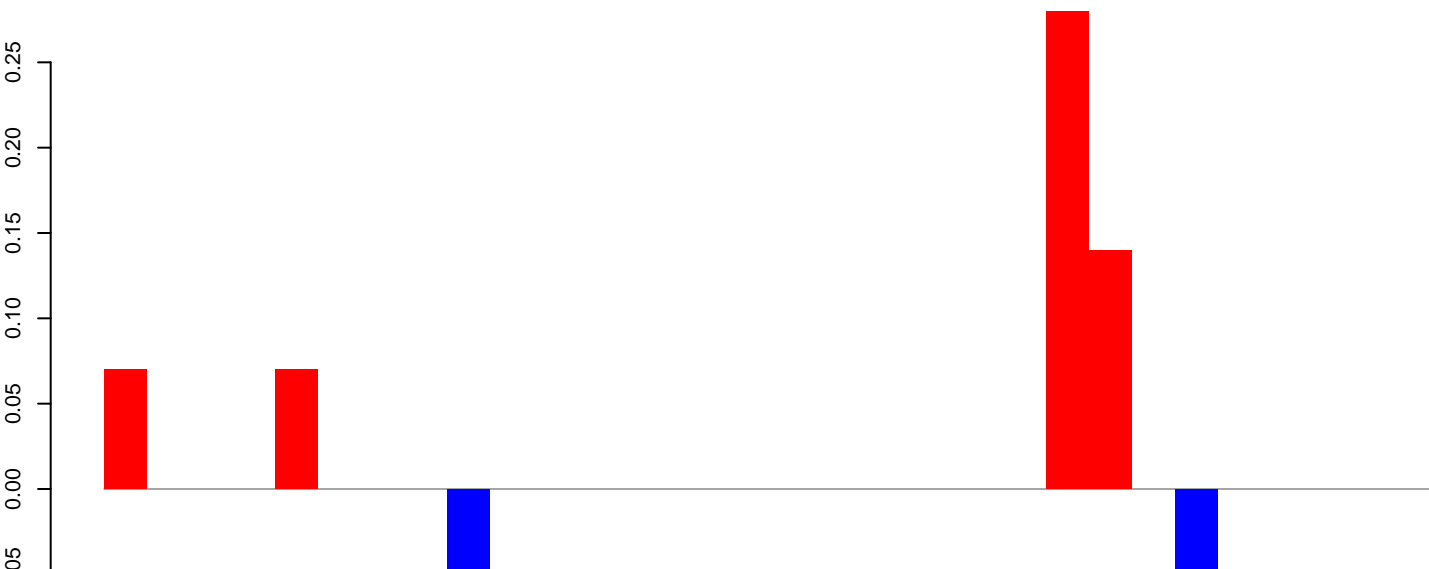
positive=0, negative=0, total=0

CuQuin_Mcarc_NL.24_35.rep



positive=0, negative=0, total=1

CuQuin_Mcarc_NL.rep

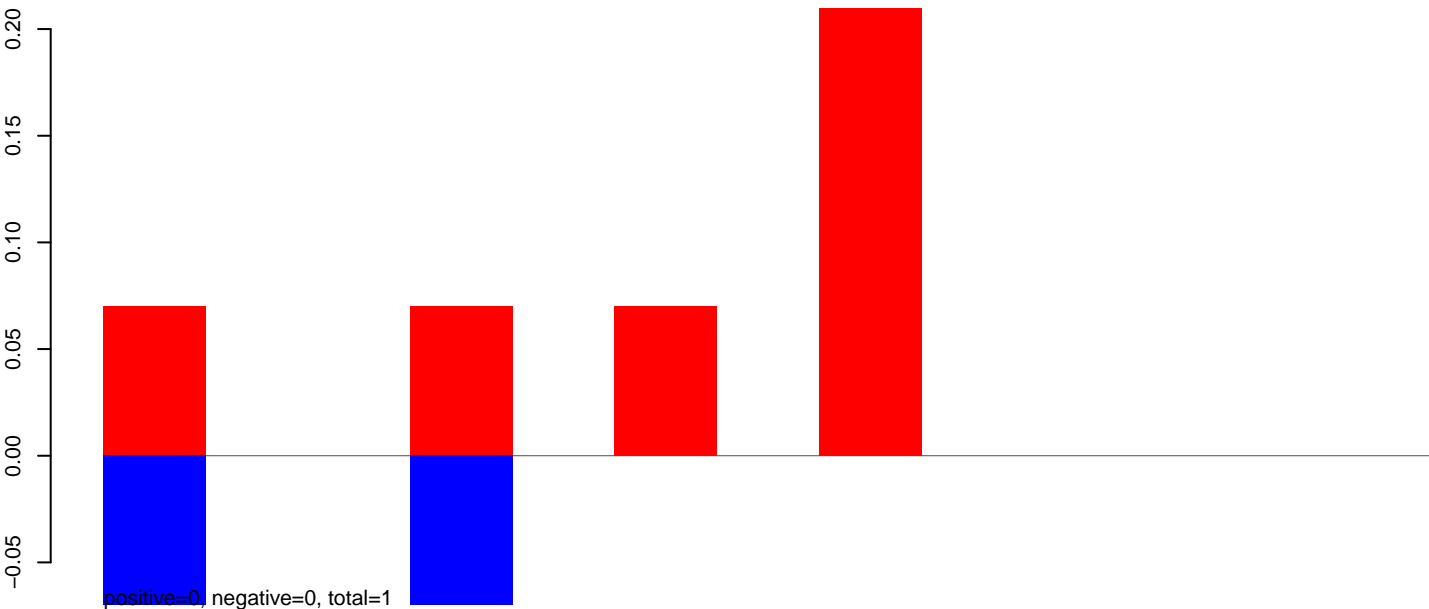


positive=1, negative=0, total=1

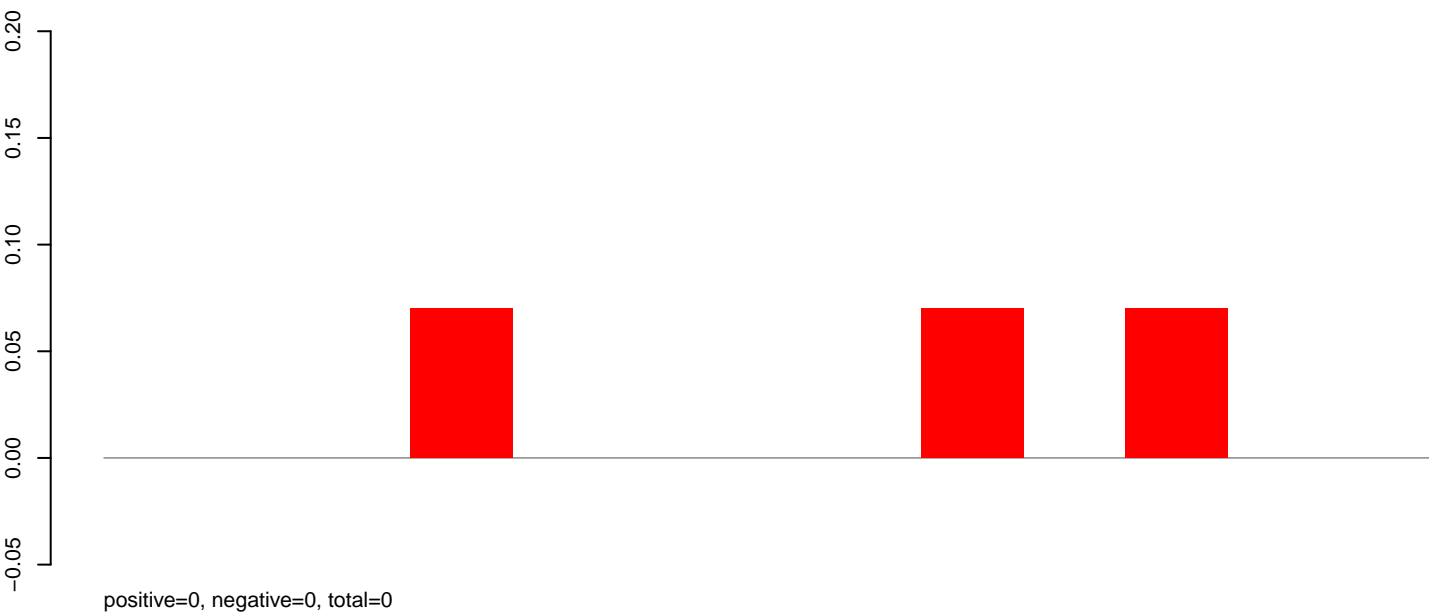
Window size=25, length=766, TE@rnd-5_family-975@Unknown@1849:1-766

0 200 400 600 800

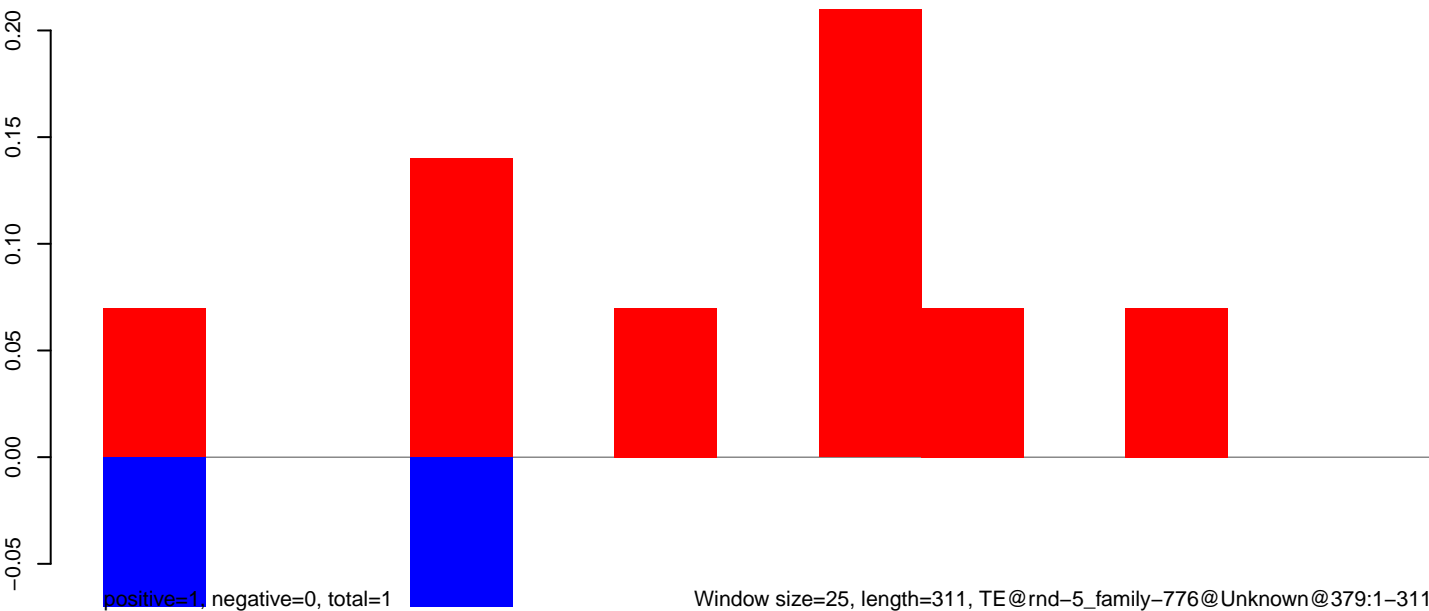
CuQuin_Mcarc_NL.18_23.rep



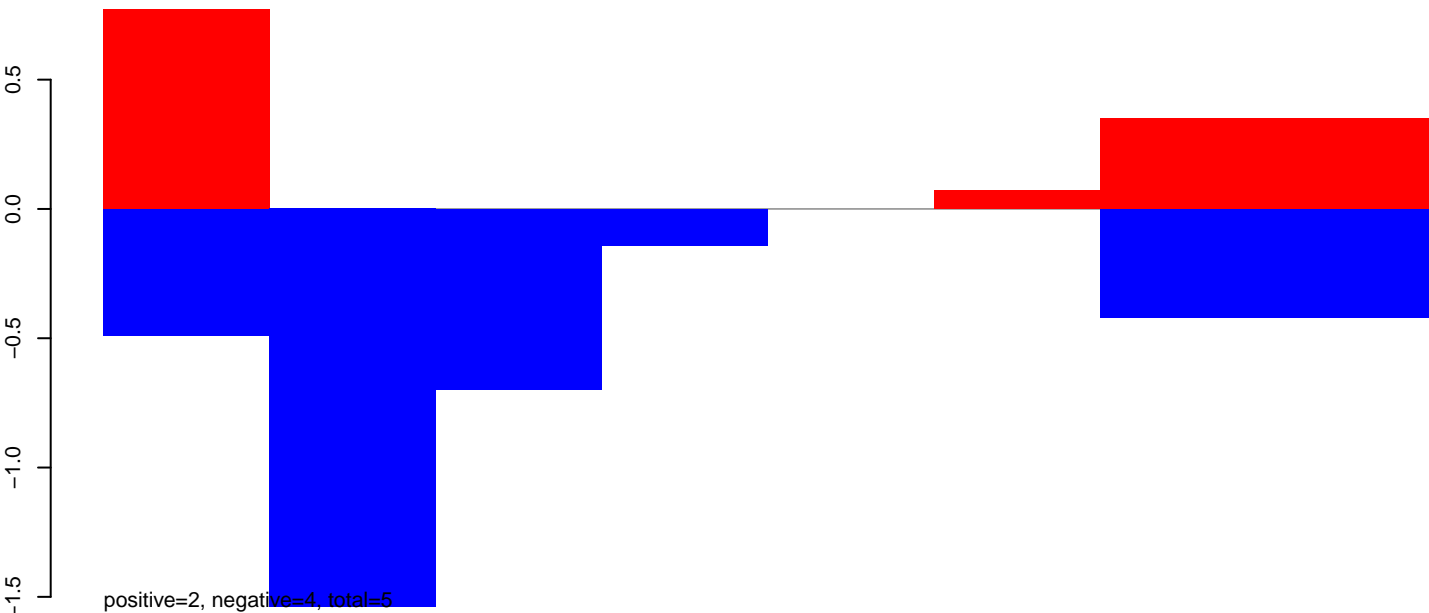
CuQuin_Mcarc_NL.24_35.rep



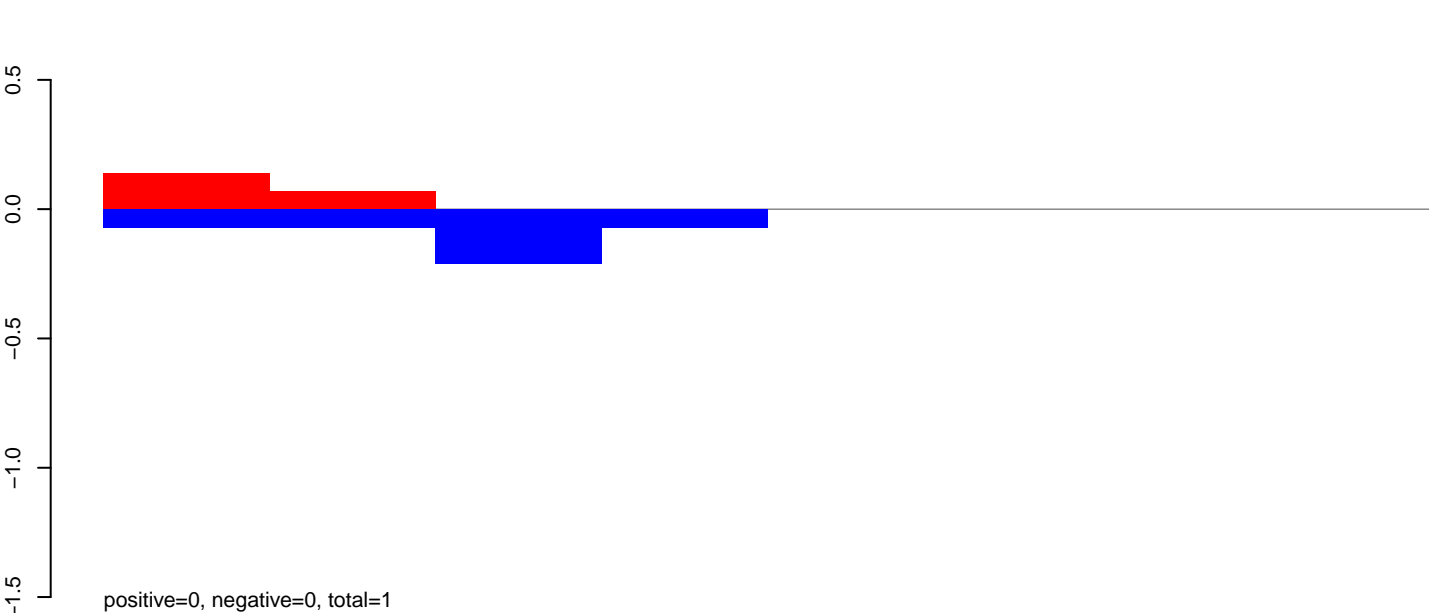
CuQuin_Mcarc_NL.rep



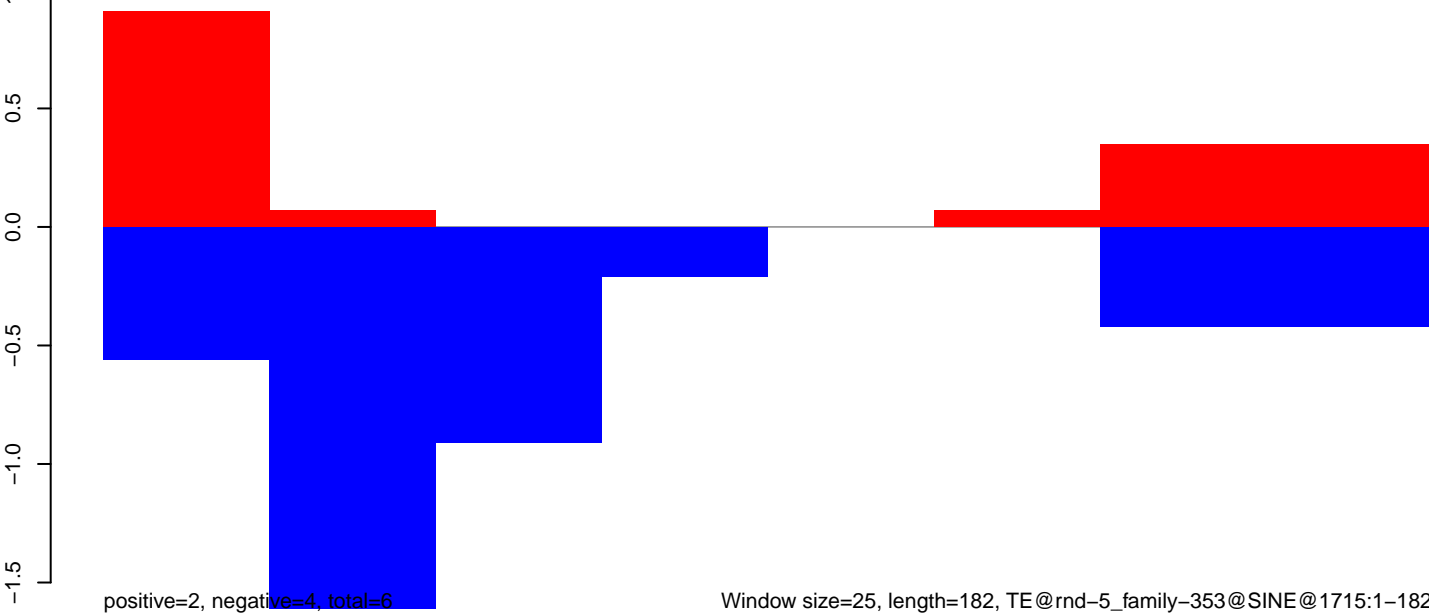
CuQuin_Mcarc_NL.18_23.rep



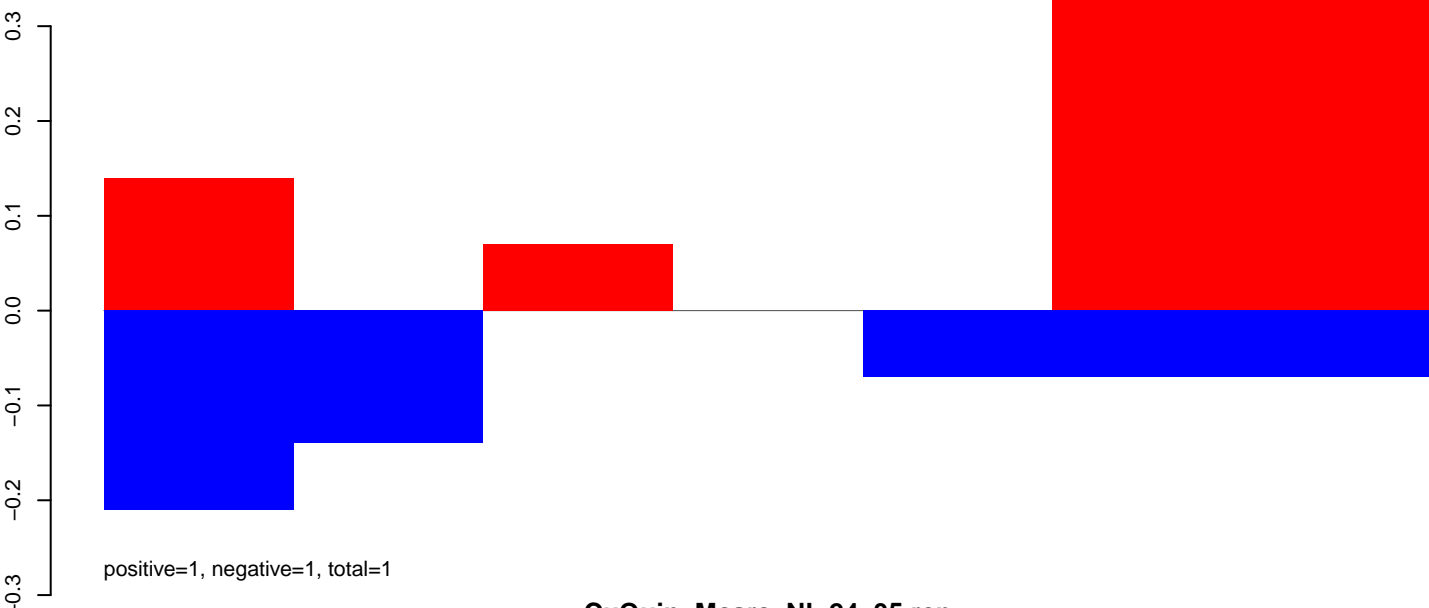
CuQuin_Mcarc_NL.24_35.rep



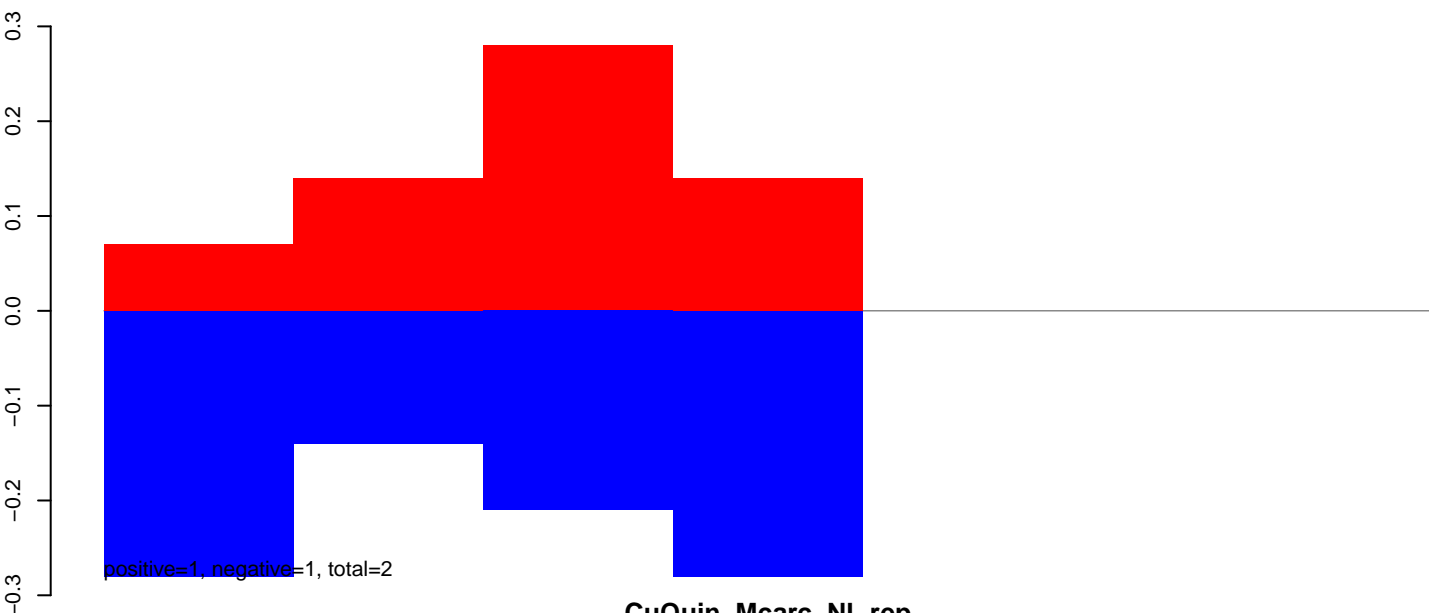
CuQuin_Mcarc_NL.rep



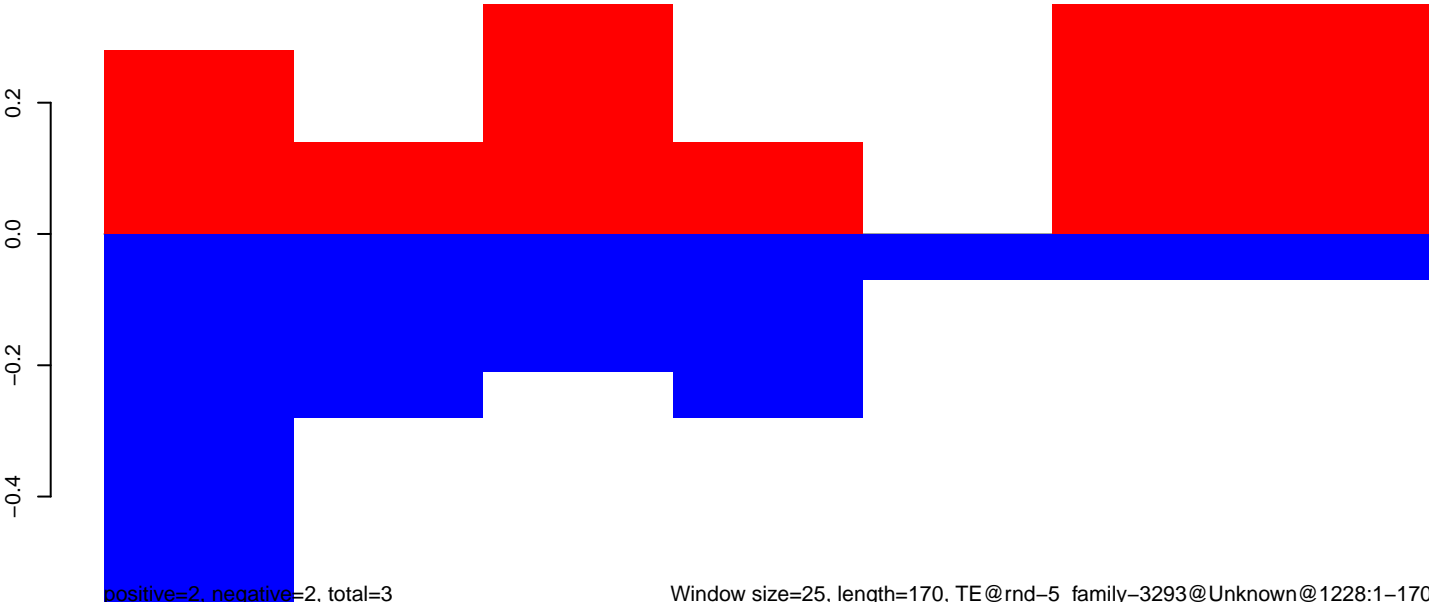
CuQuin_Mcarc_NL.18_23.rep



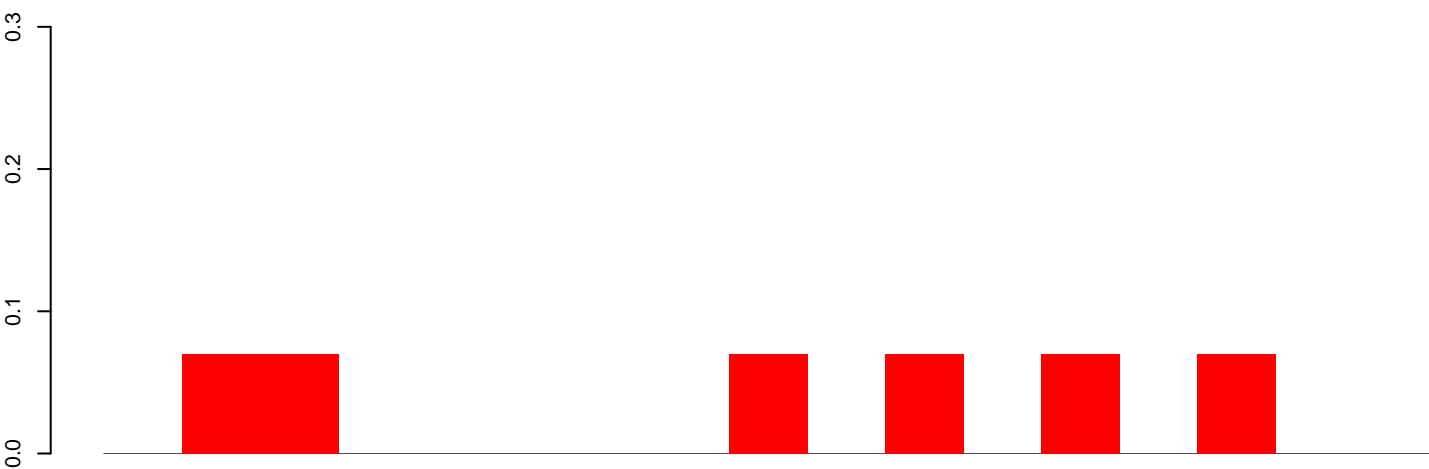
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

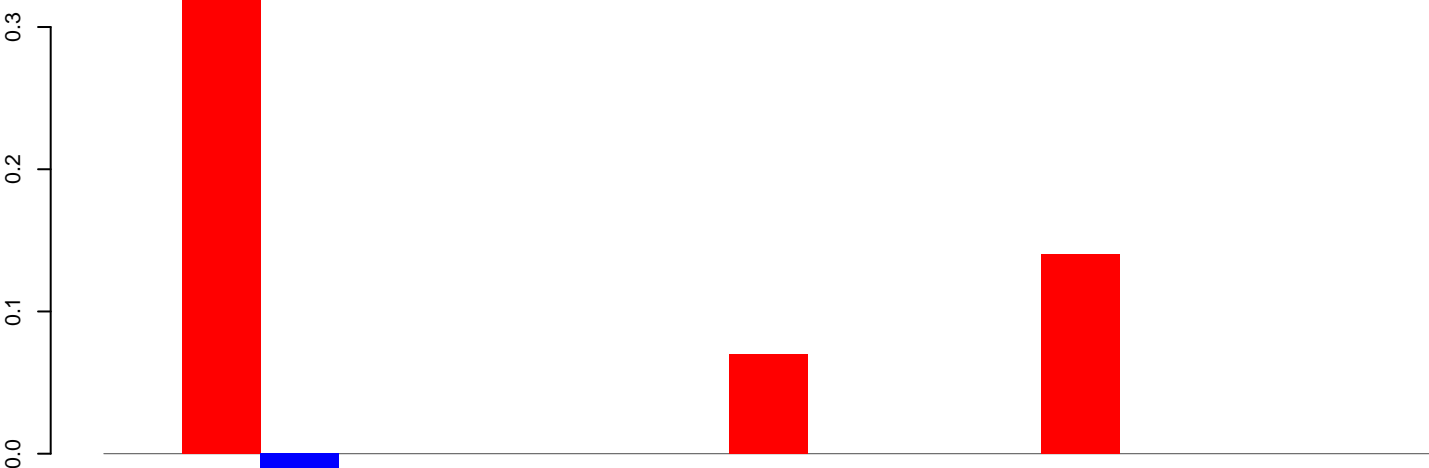


CuQuin_Mcarc_NL.18_23.rep



positive=0, negative=0, total=0

CuQuin_Mcarc_NL.24_35.rep



positive=1, negative=0, total=1

CuQuin_Mcarc_NL.rep



positive=2, negative=0, total=2

Window size=25, length=417, TE@rnd-5_family-2527@Unknown@3119:1-417

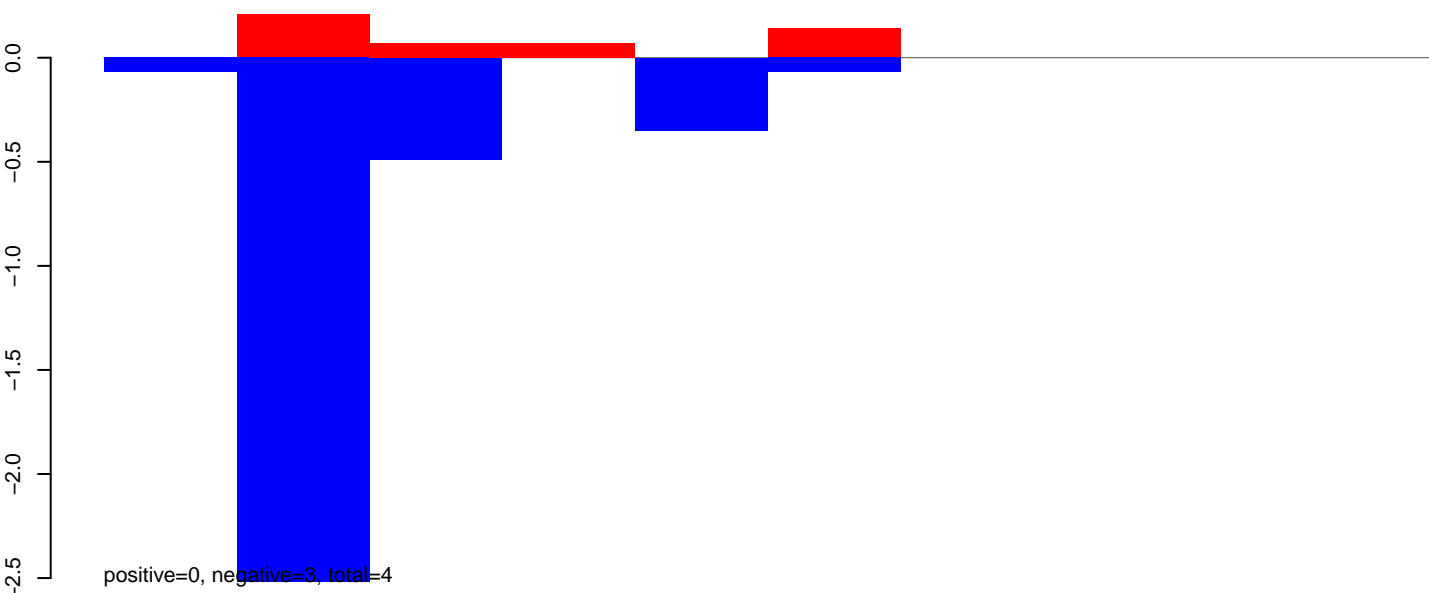
100

200

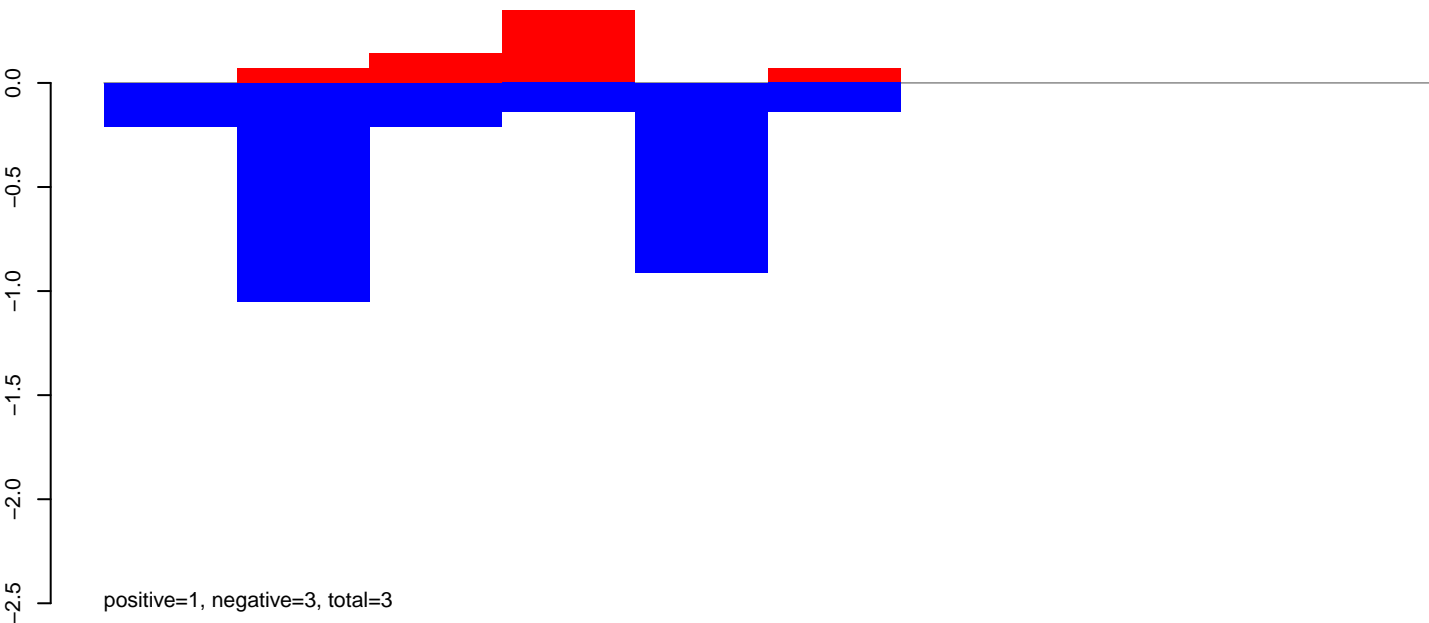
300

400

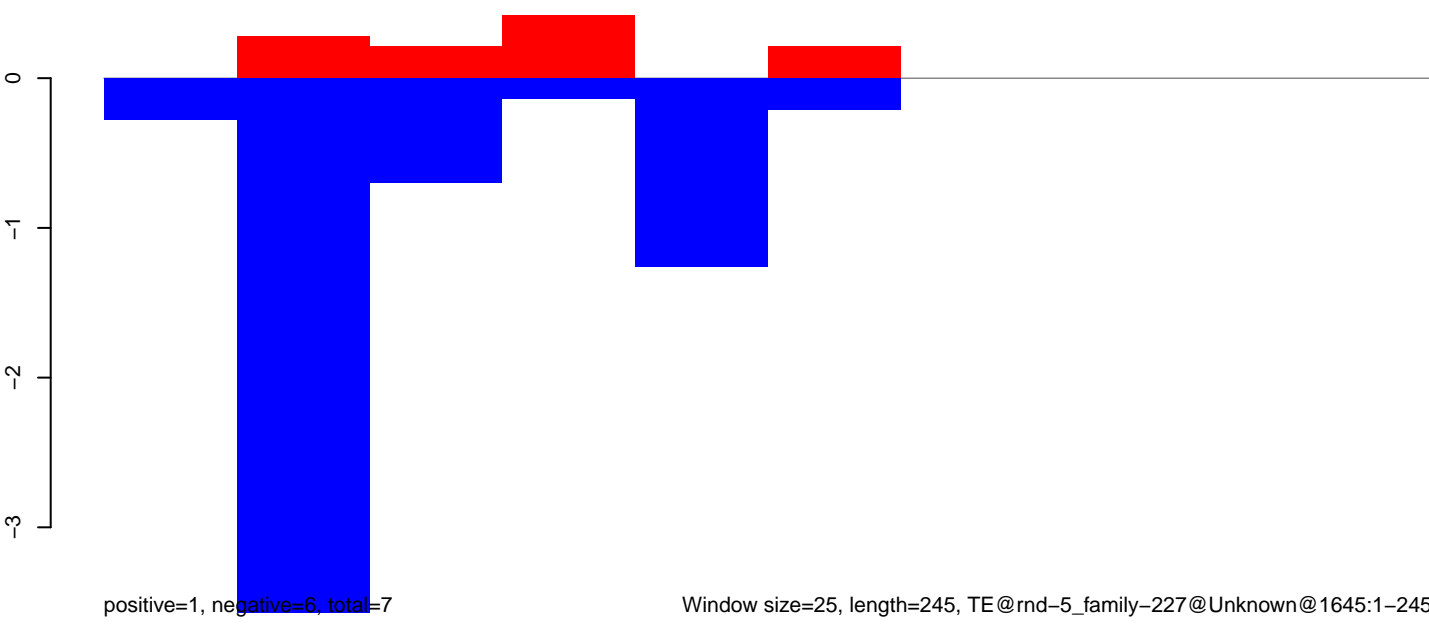
CuQuin_Mcarc_NL.18_23.rep



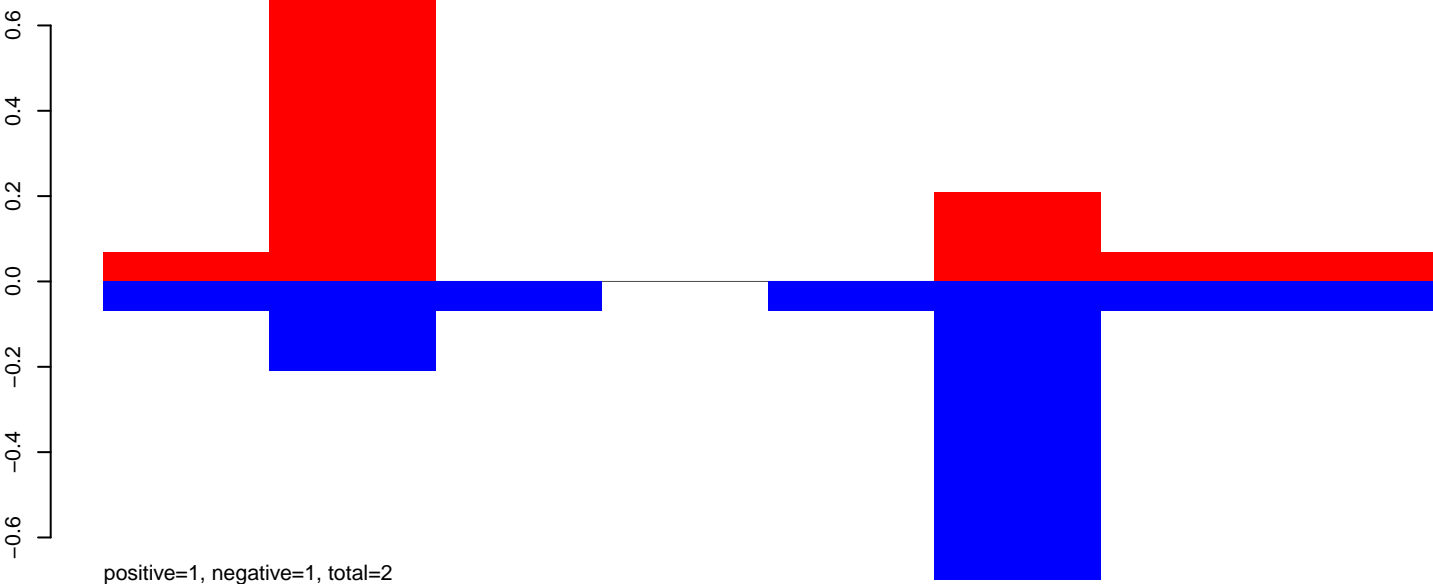
CuQuin_Mcarc_NL.24_35.rep



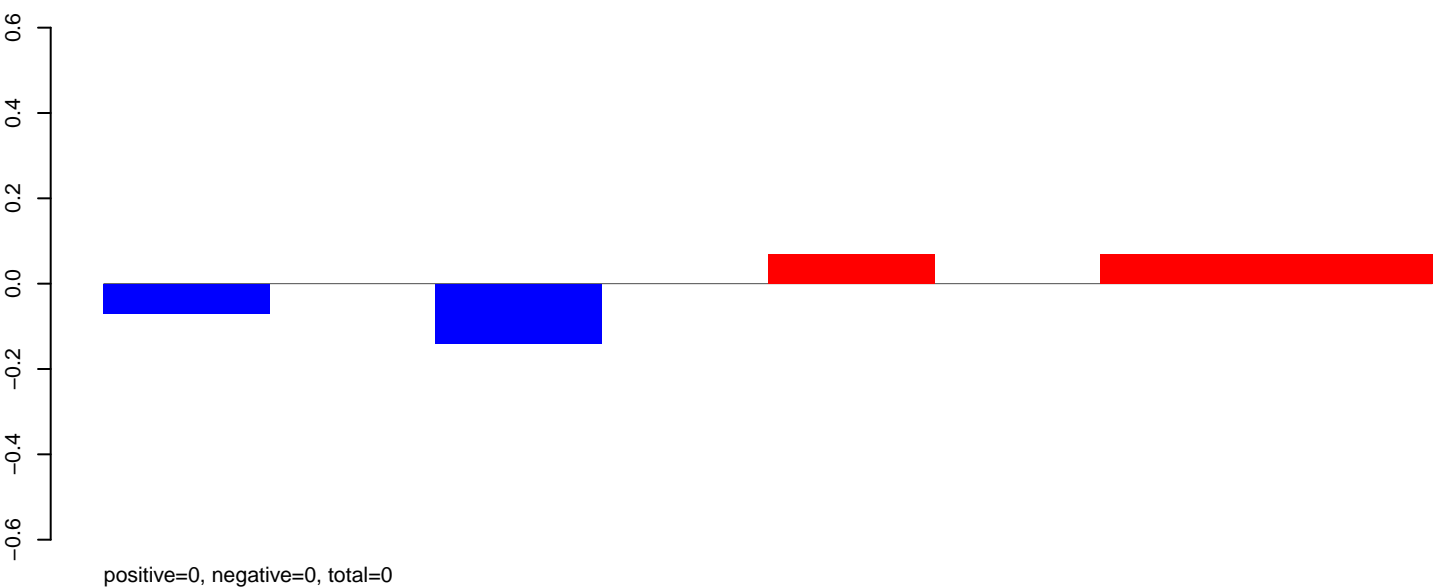
CuQuin_Mcarc_NL.rep



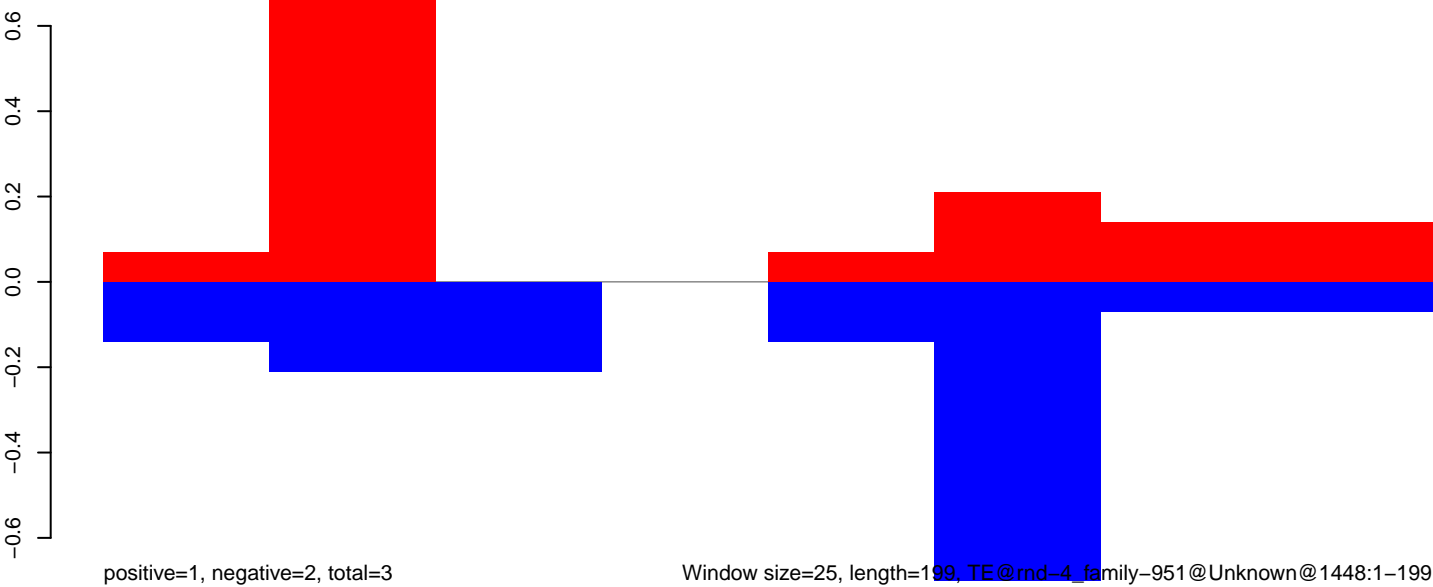
CuQuin_Mcarc_NL.18_23.rep



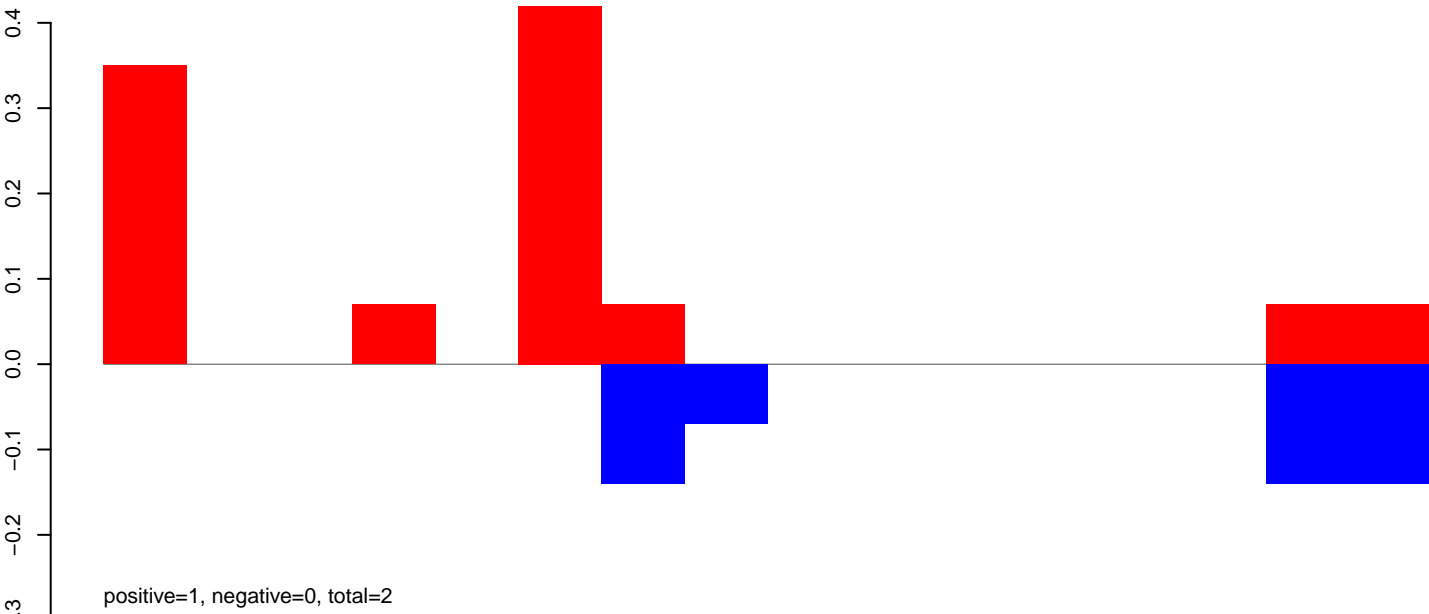
CuQuin_Mcarc_NL.24_35.rep



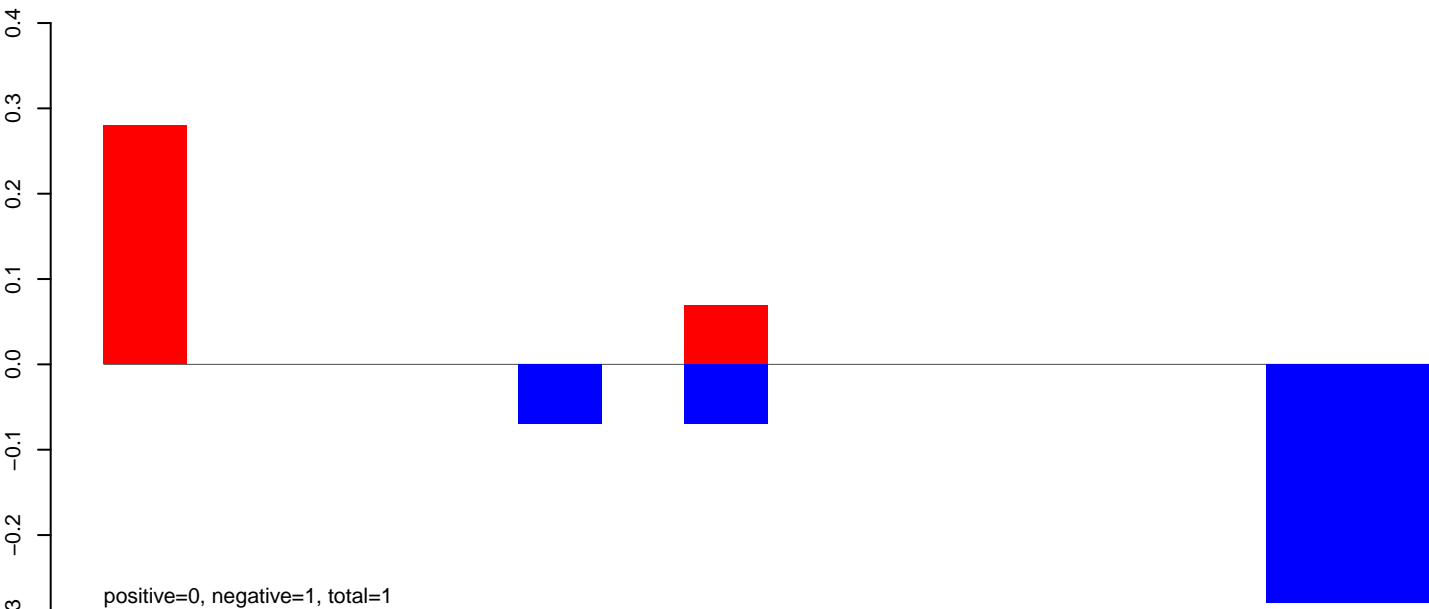
CuQuin_Mcarc_NL.rep



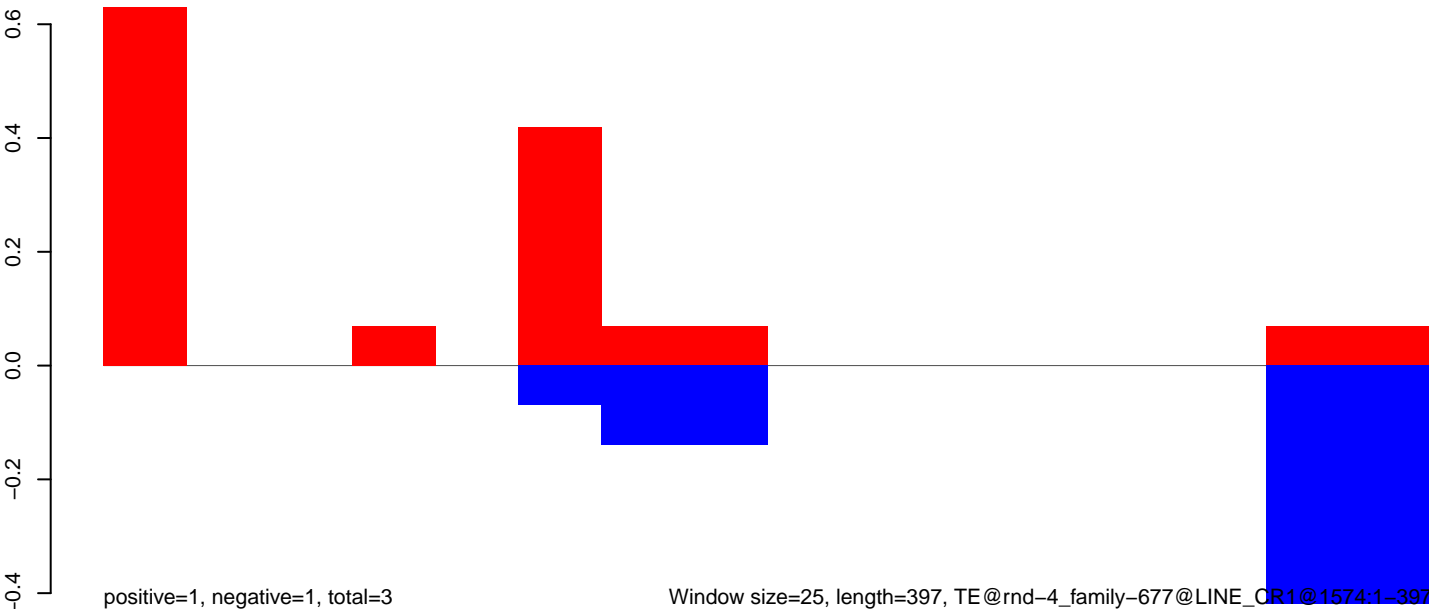
CuQuin_Mcarc_NL.18_23.rep



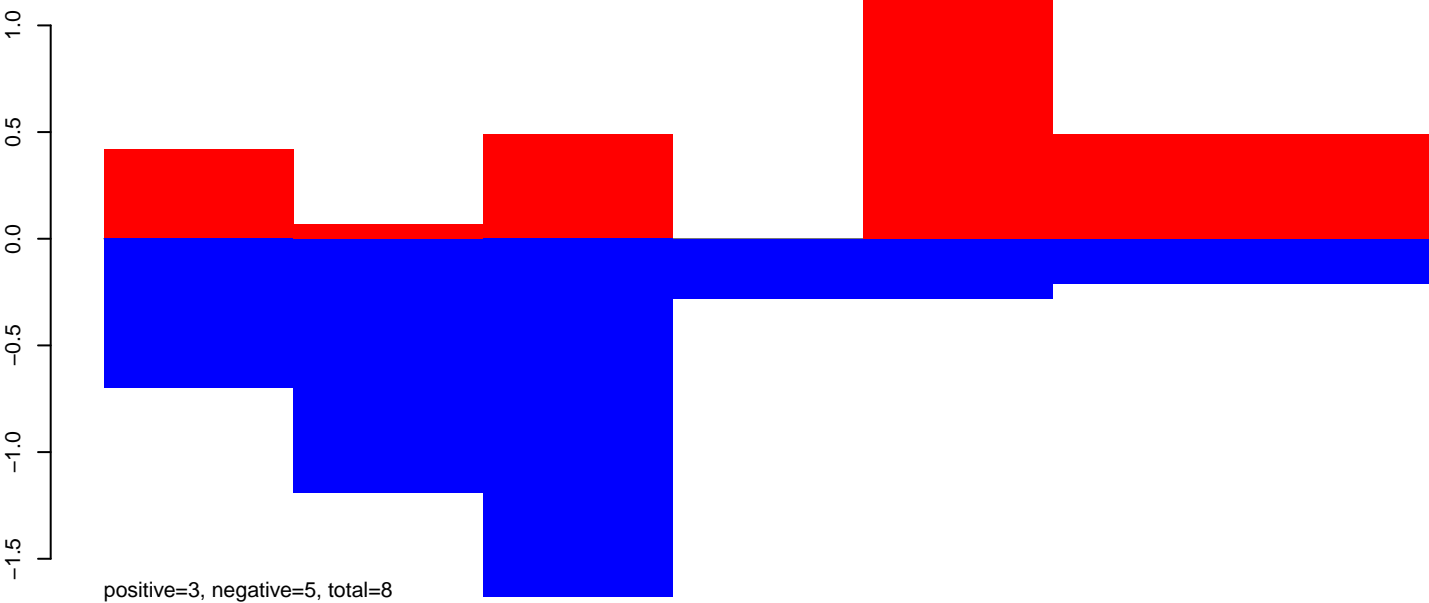
CuQuin_Mcarc_NL.24_35.rep



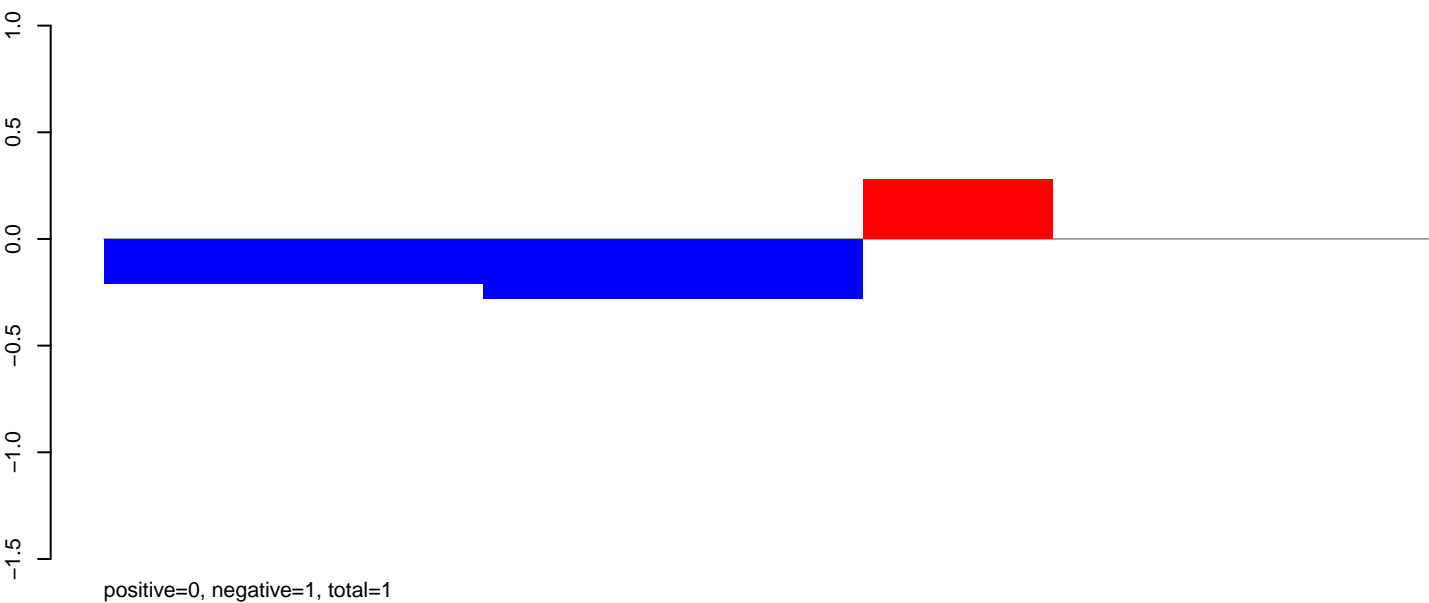
CuQuin_Mcarc_NL.rep



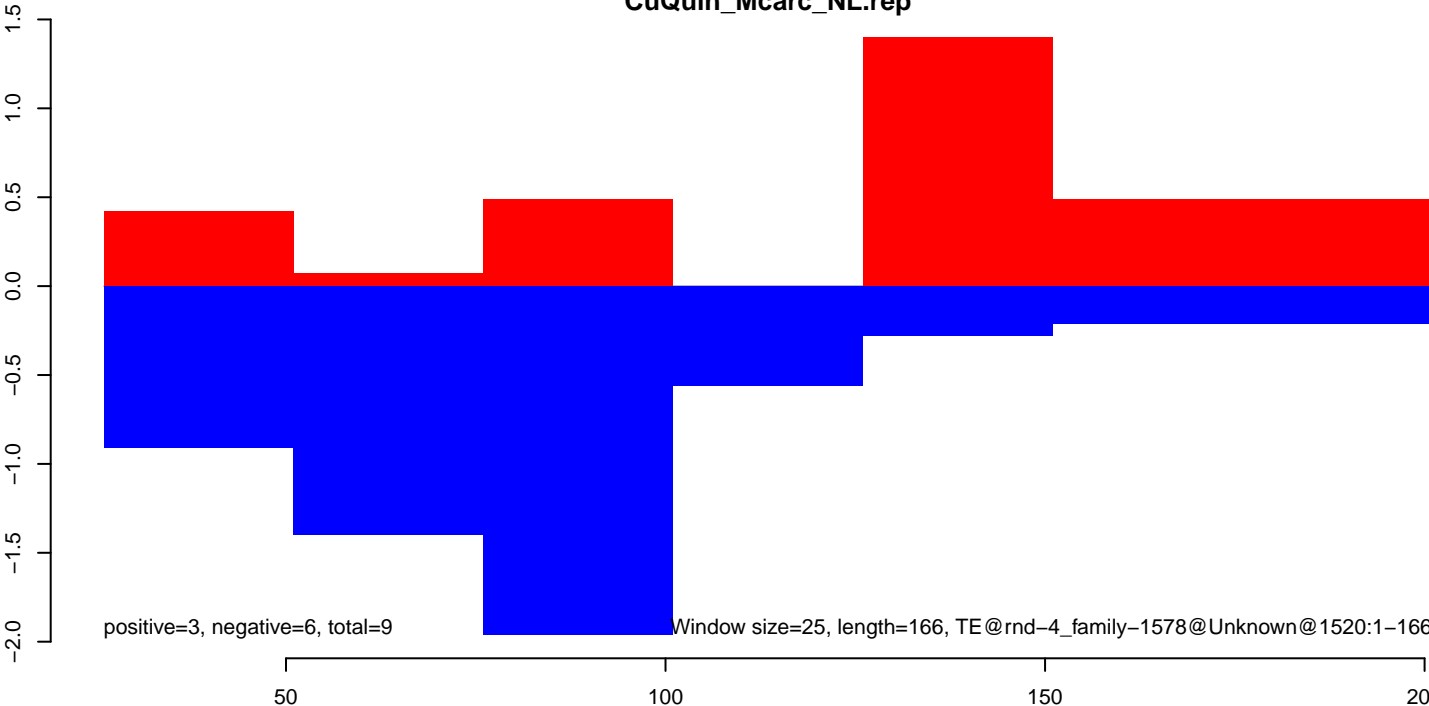
CuQuin_Mcarc_NL.18_23.rep



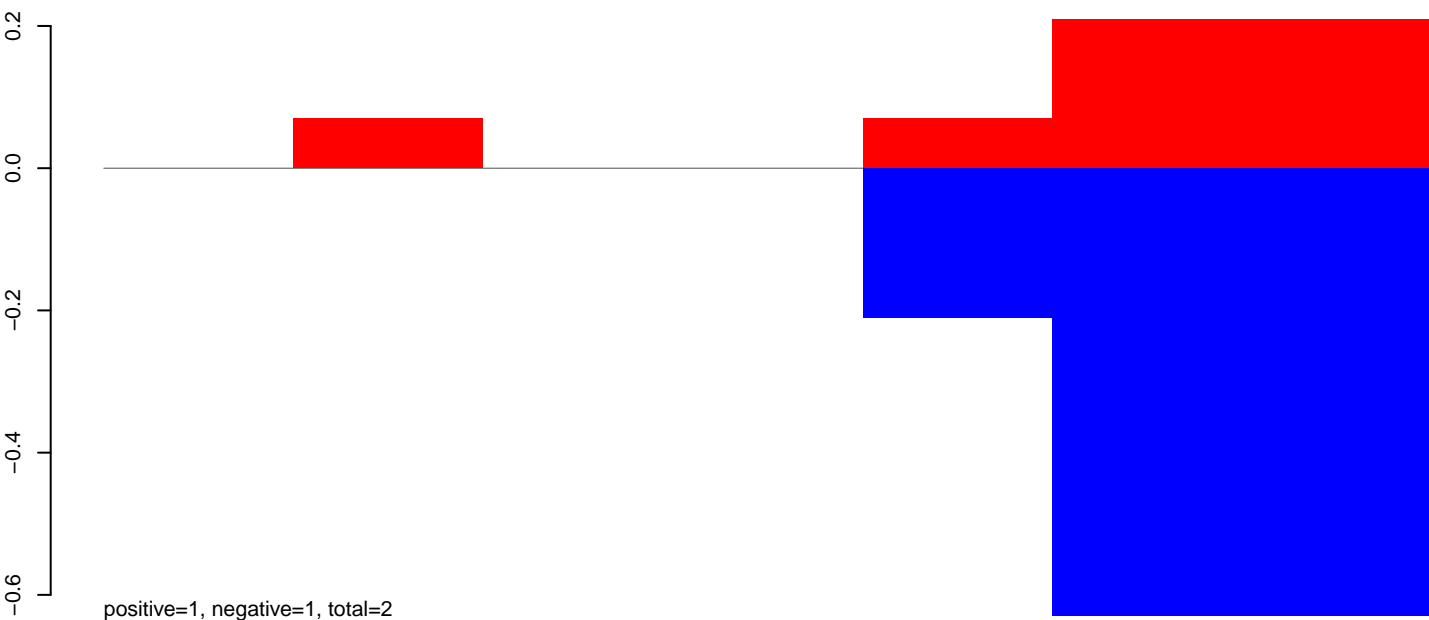
CuQuin_Mcarc_NL.24_35.rep



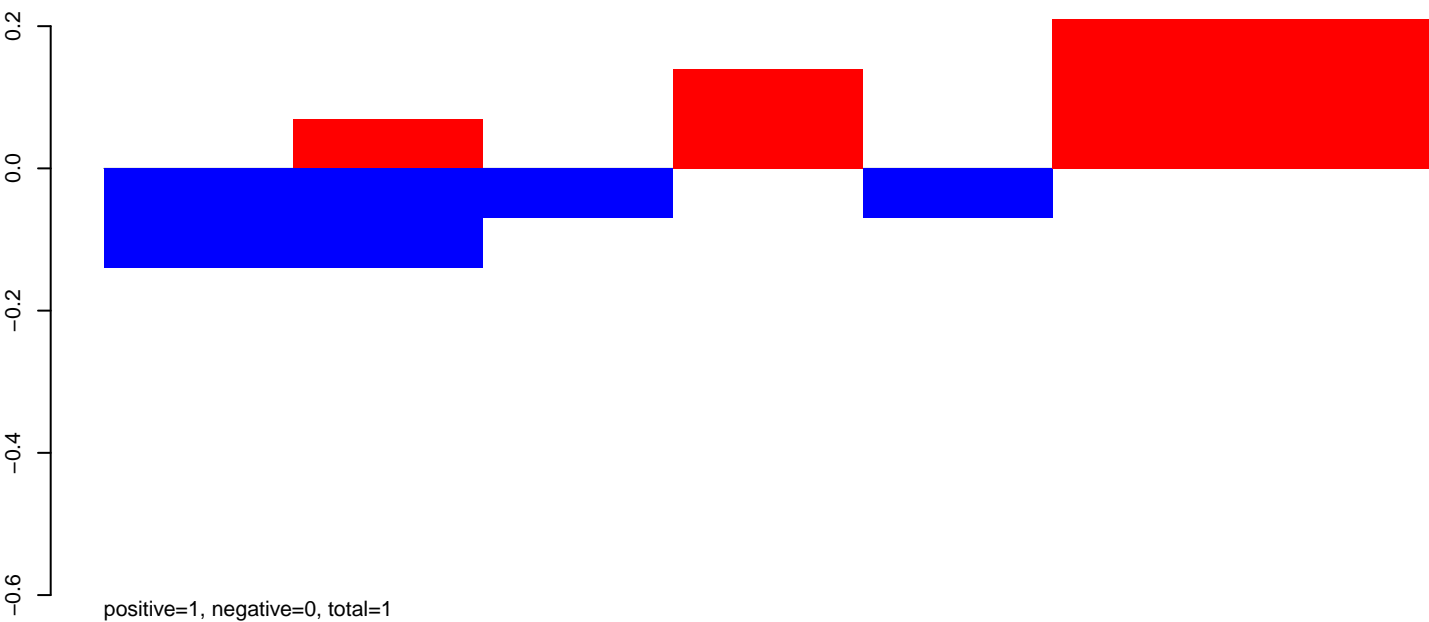
CuQuin_Mcarc_NL.rep



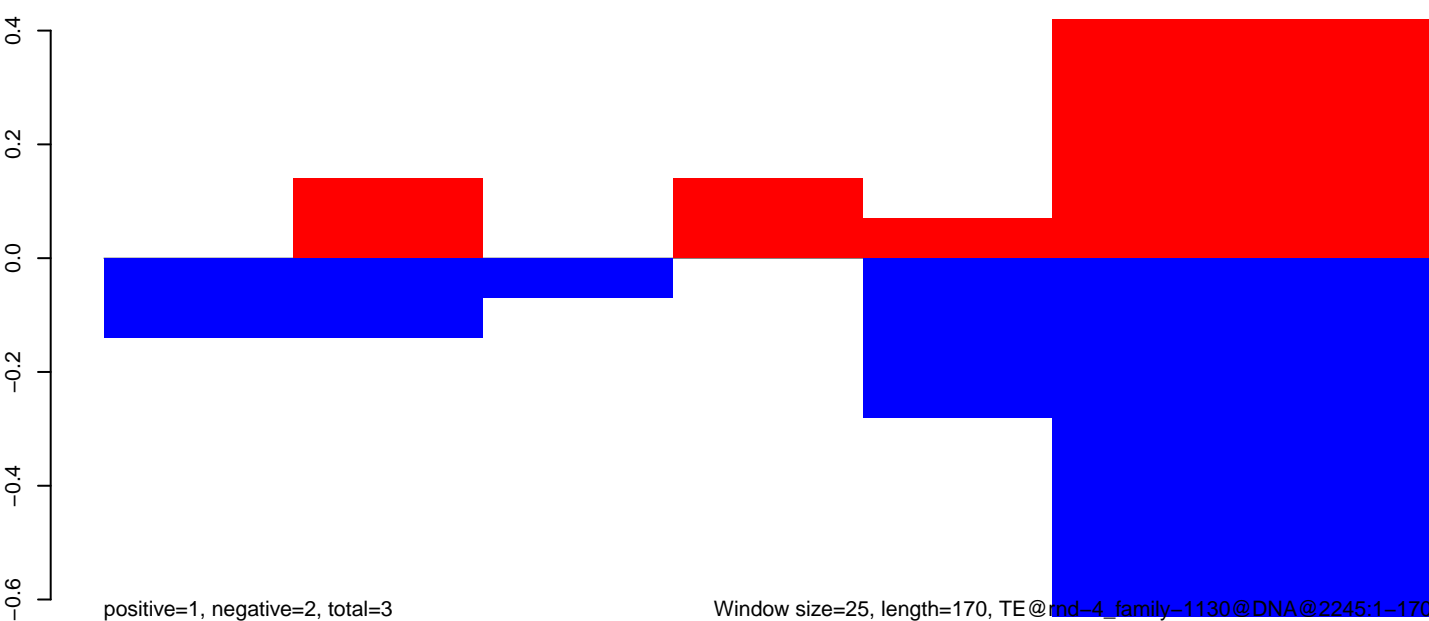
CuQuin_Mcarc_NL.18_23.rep



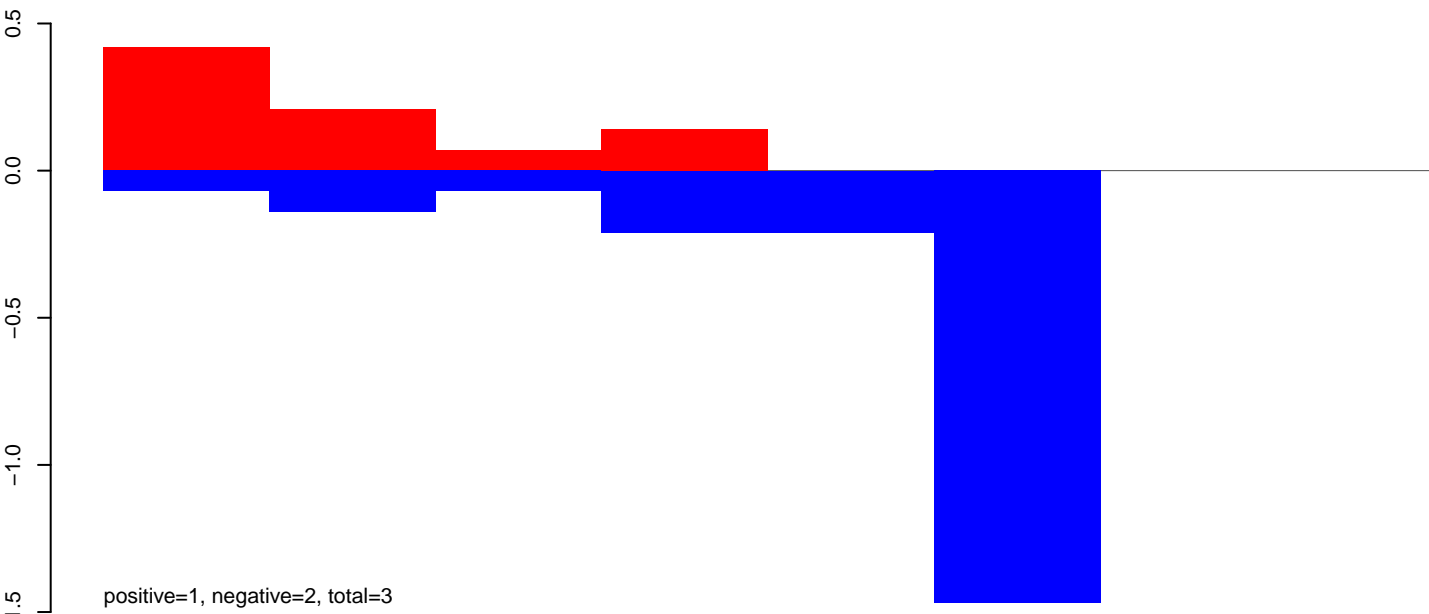
CuQuin_Mcarc_NL.24_35.rep



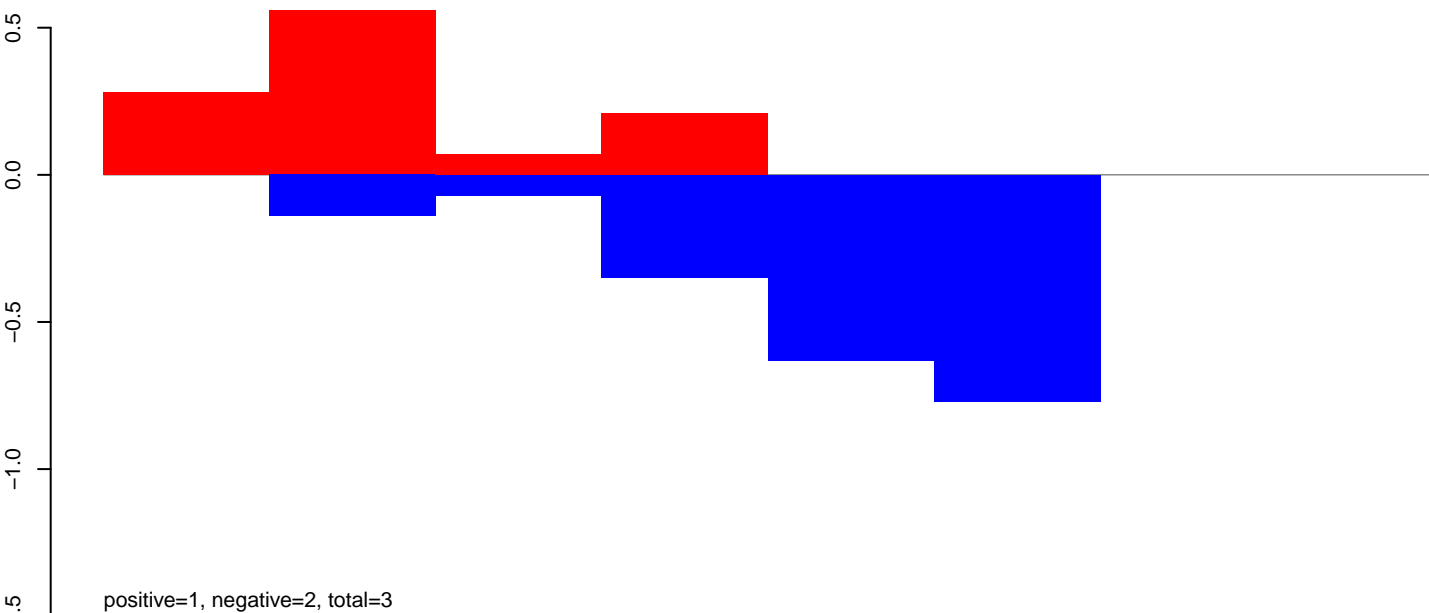
CuQuin_Mcarc_NL.rep



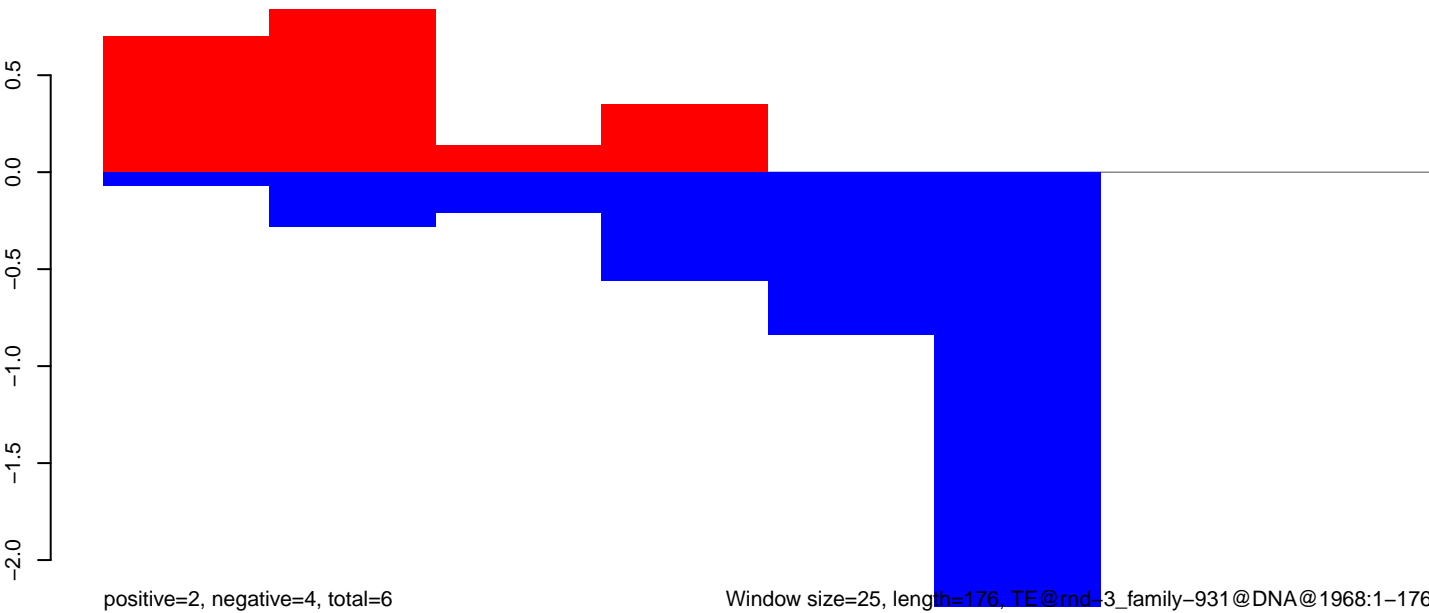
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

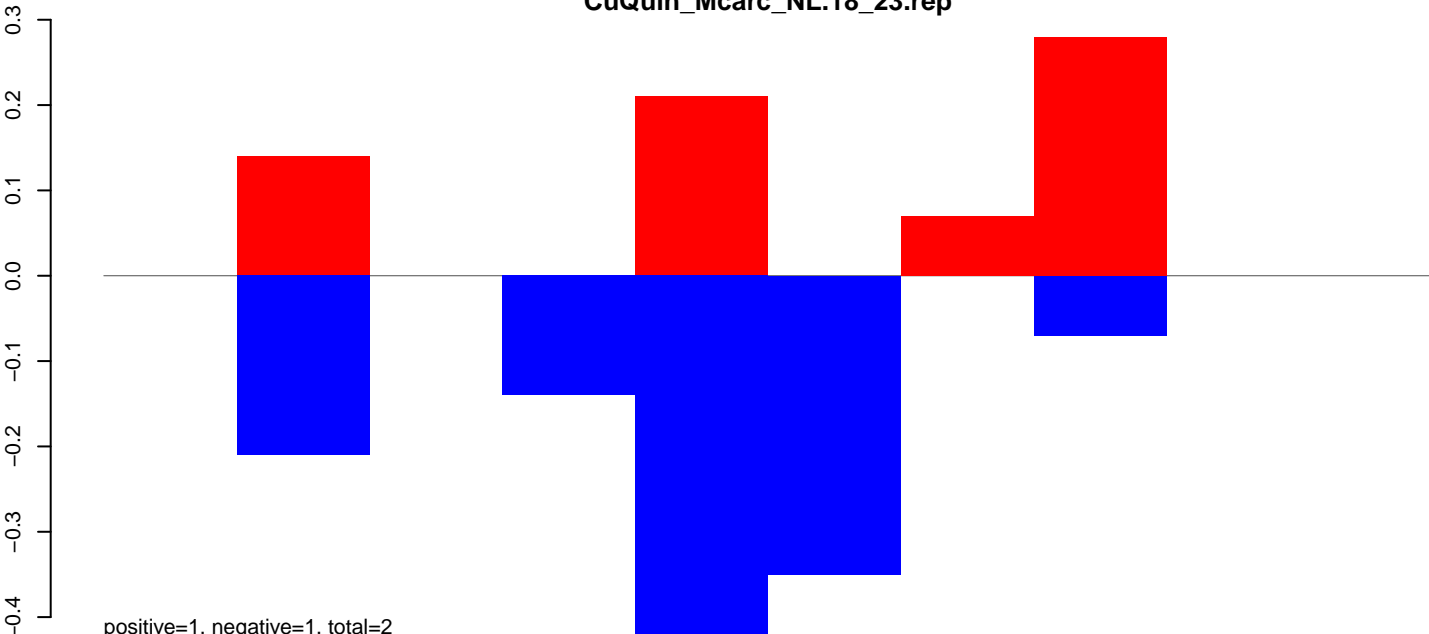


CuQuin_Mcarc_NL.rep

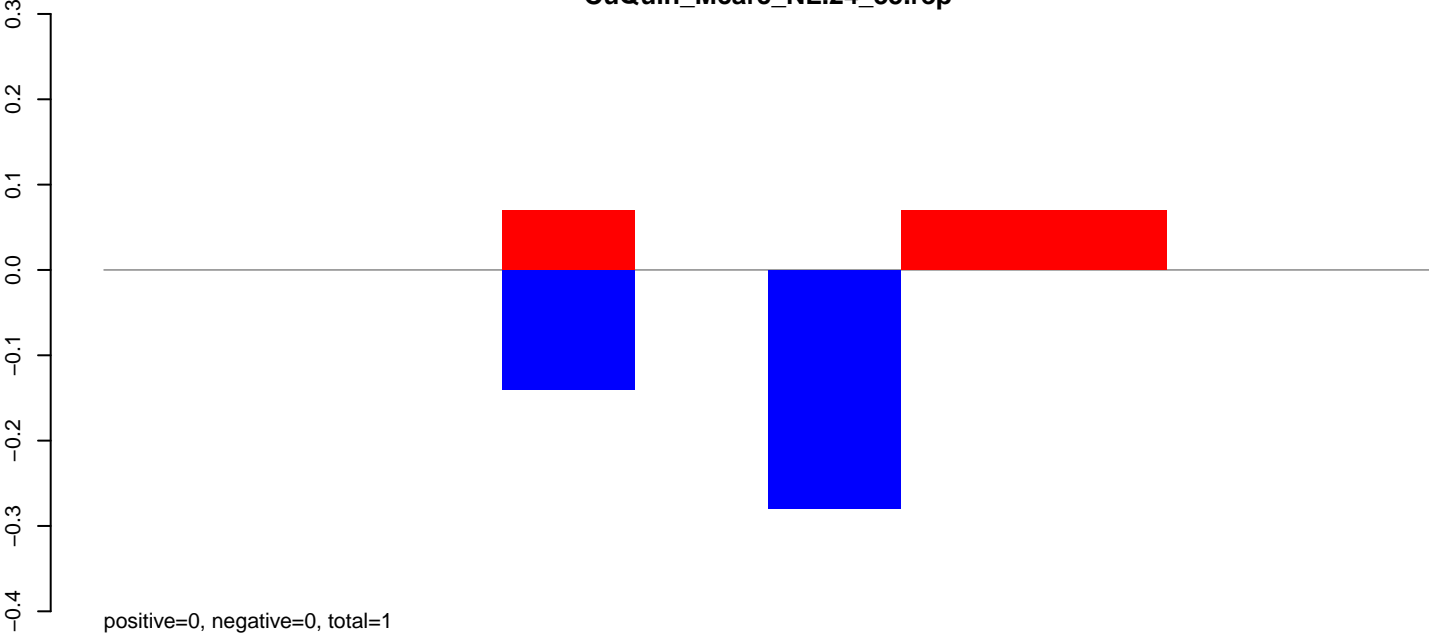


Window size=25, length=176, TE@rnd_3_family-931@DNA@1968:1-176

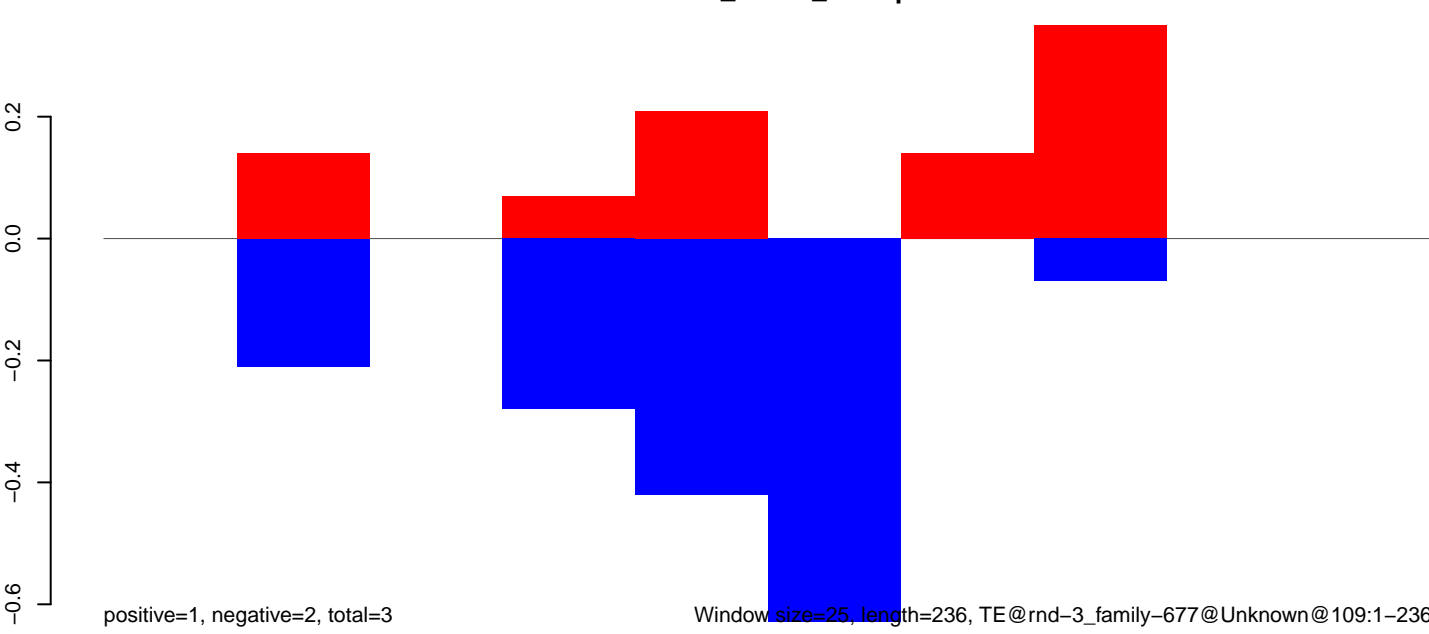
CuQuin_Mcarc_NL.18_23.rep



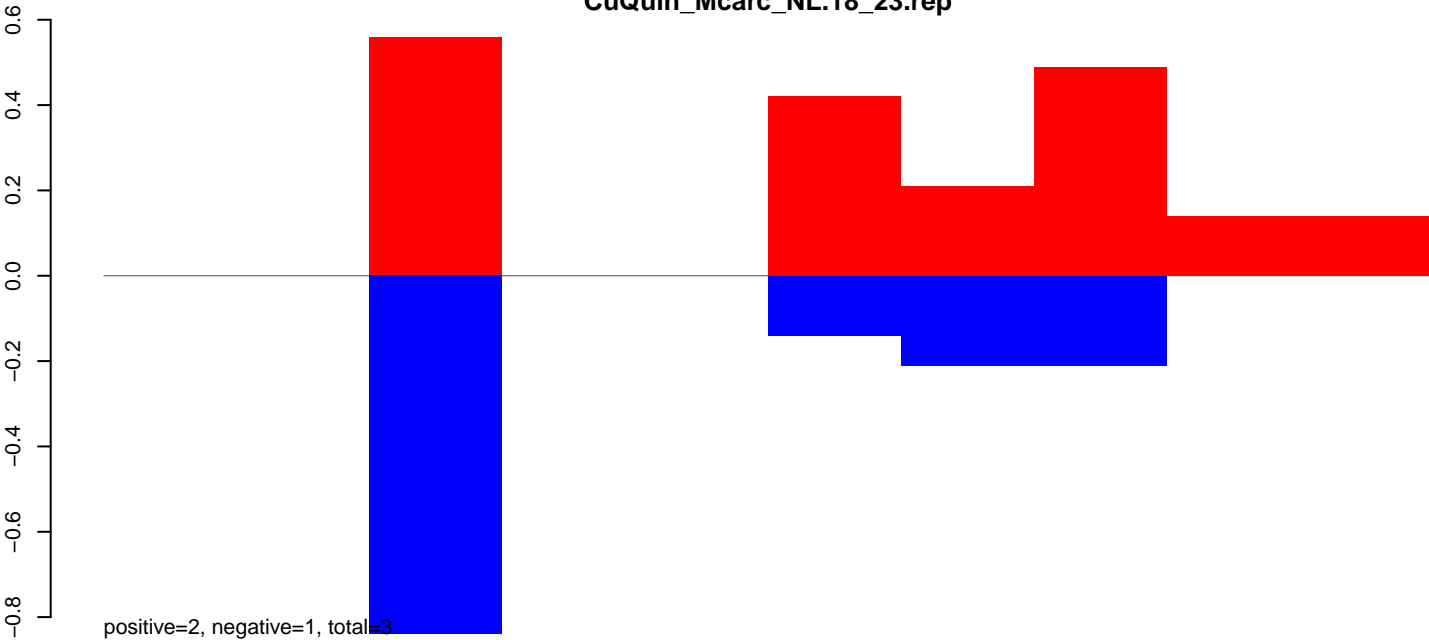
CuQuin_Mcarc_NL.24_35.rep



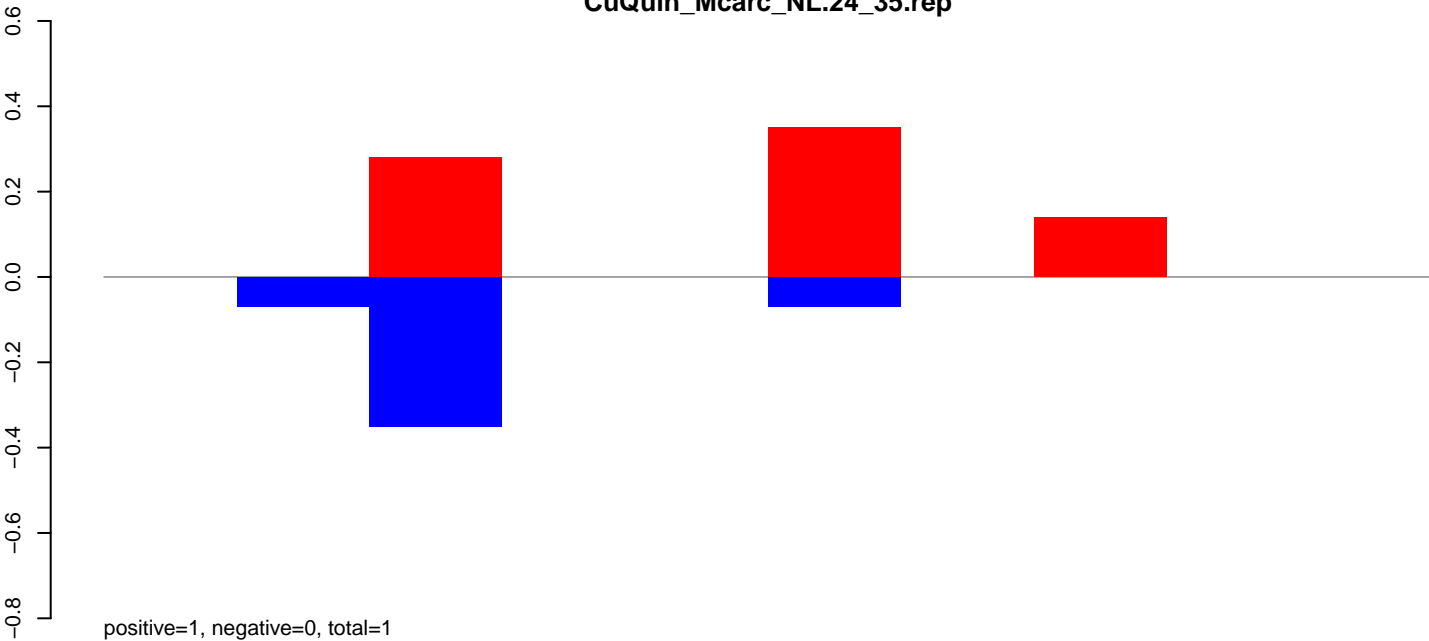
CuQuin_Mcarc_NL.rep



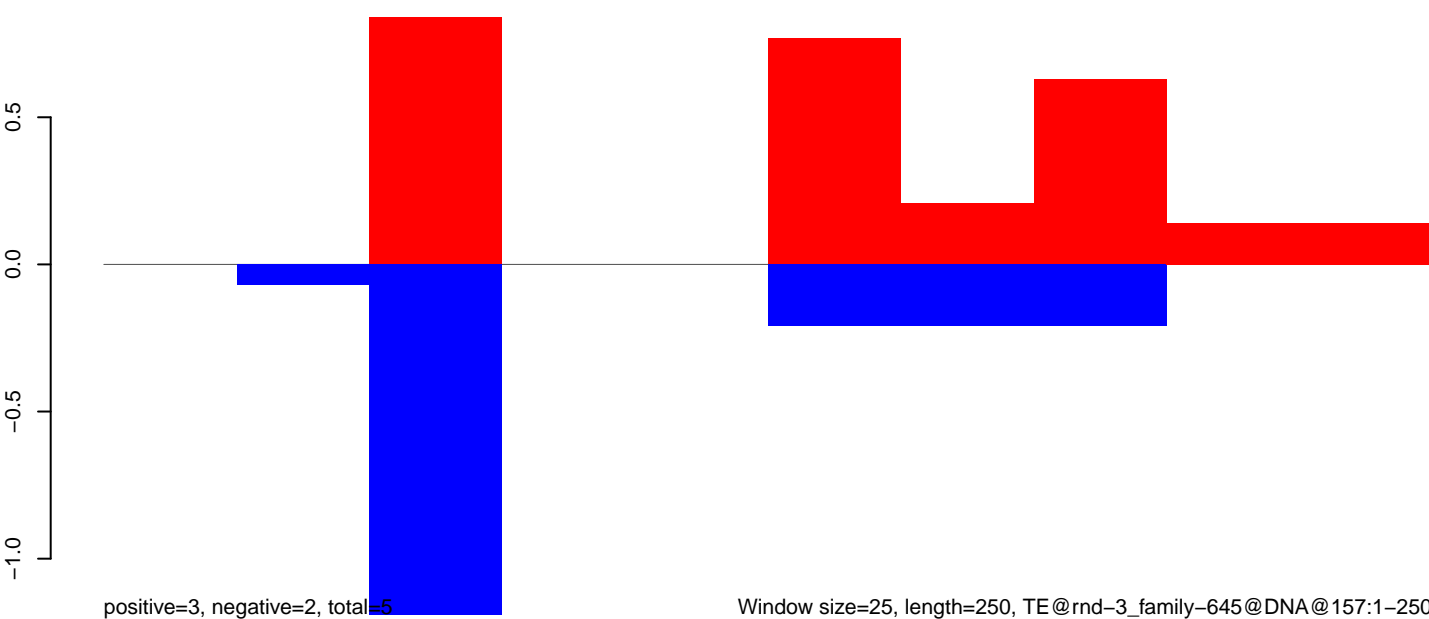
CuQuin_Mcarc_NL.18_23.rep



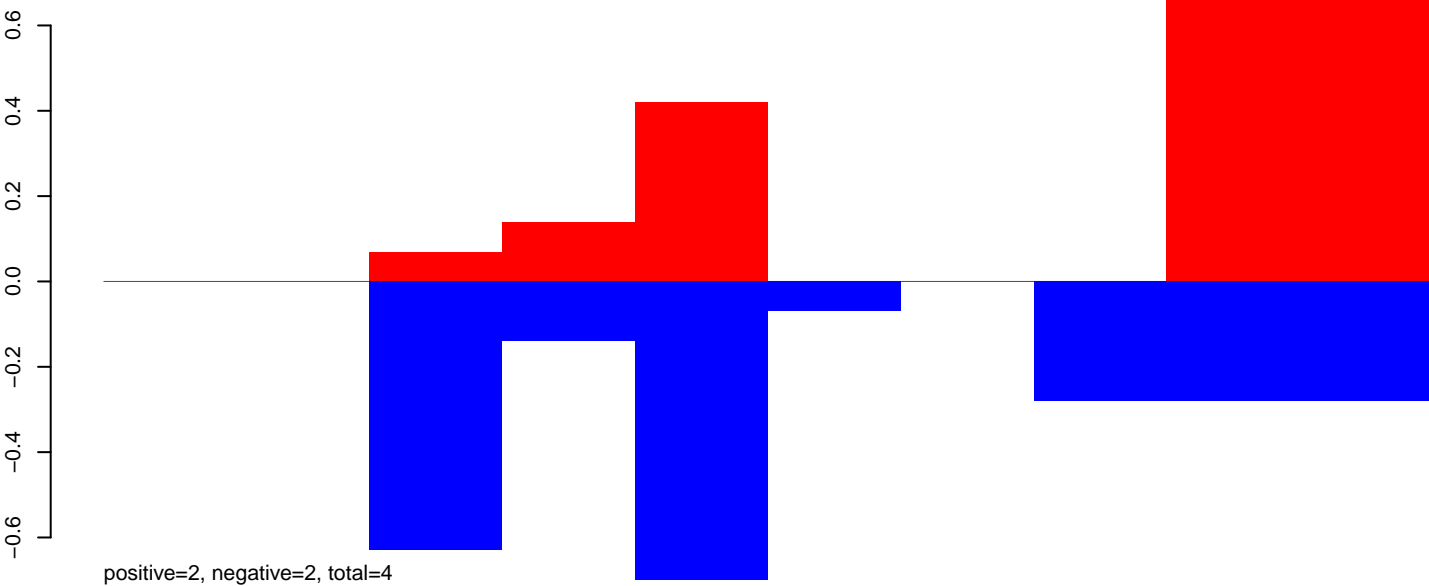
CuQuin_Mcarc_NL.24_35.rep



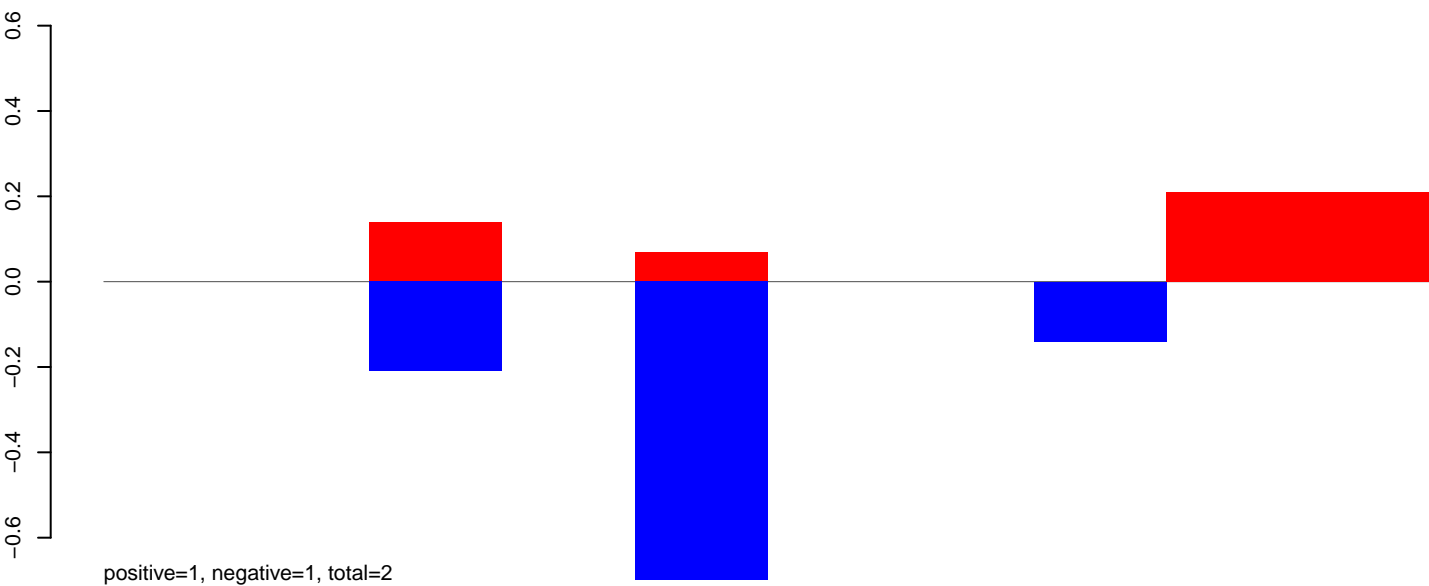
CuQuin_Mcarc_NL.rep



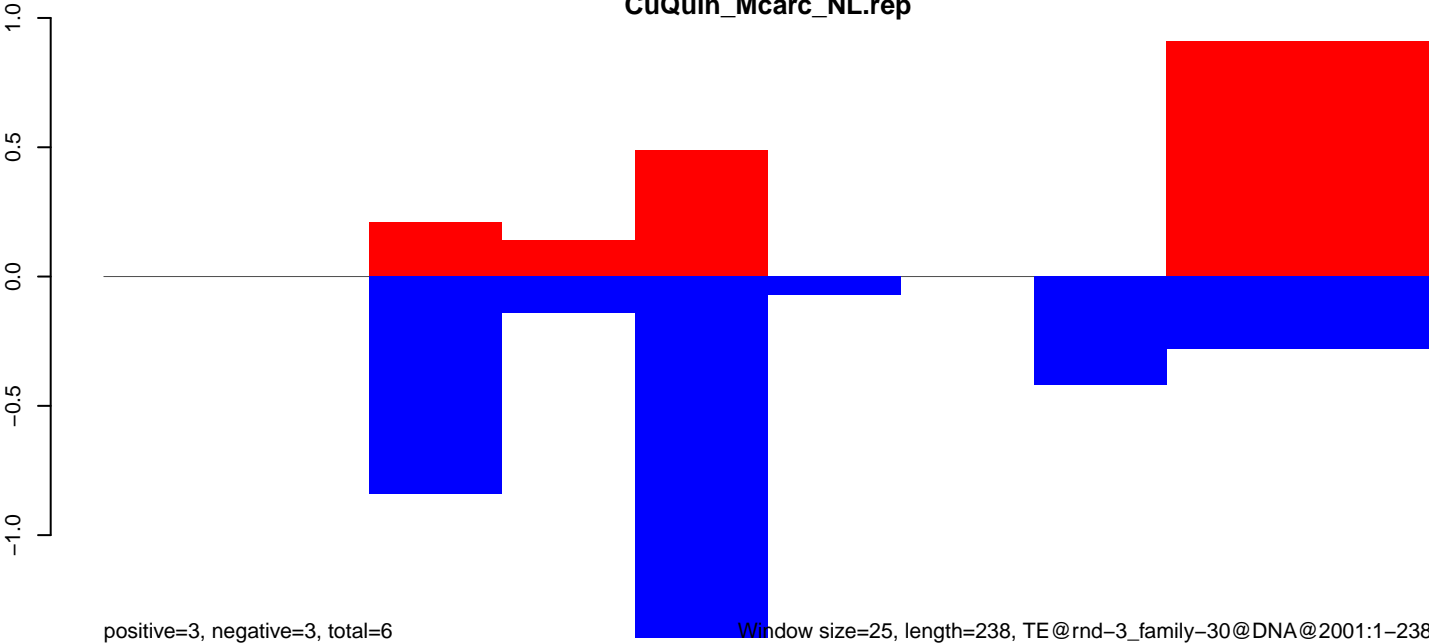
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



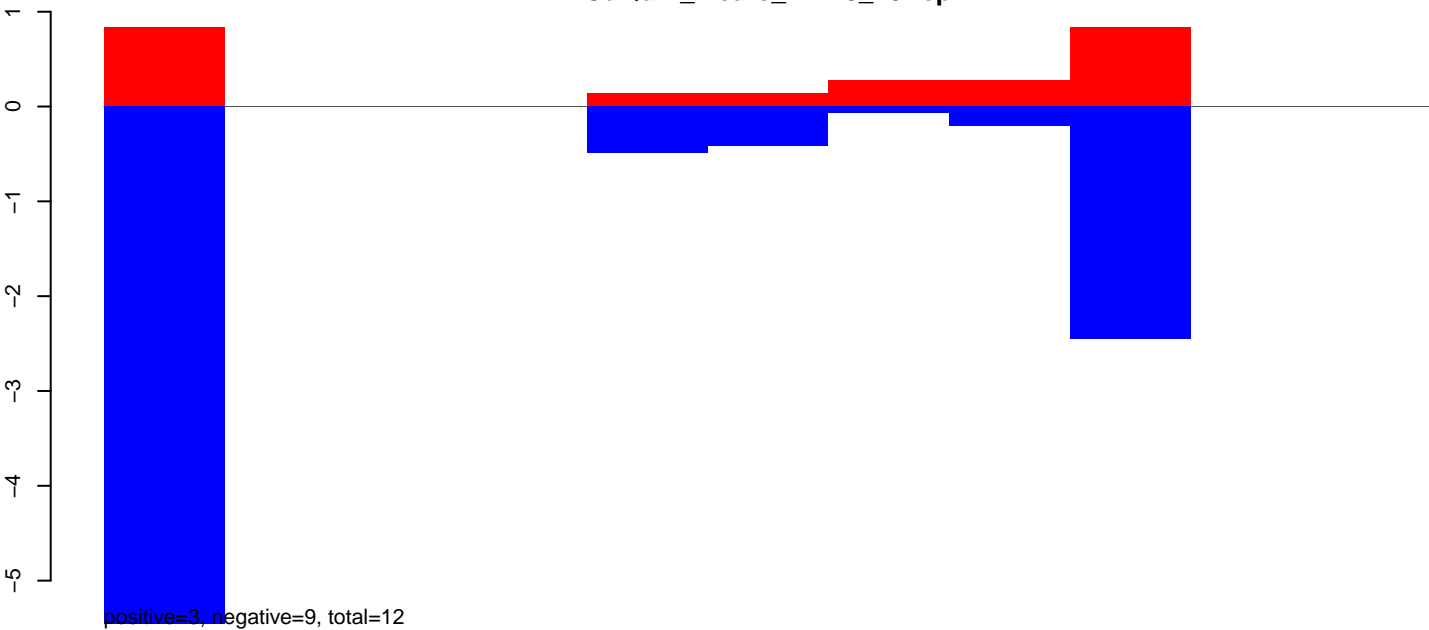
CuQuin_Mcarc_NL.rep



Window size=25, length=238, TE@rnd-3_family-30@DNA@2001:1-238

50 100 150 200 250

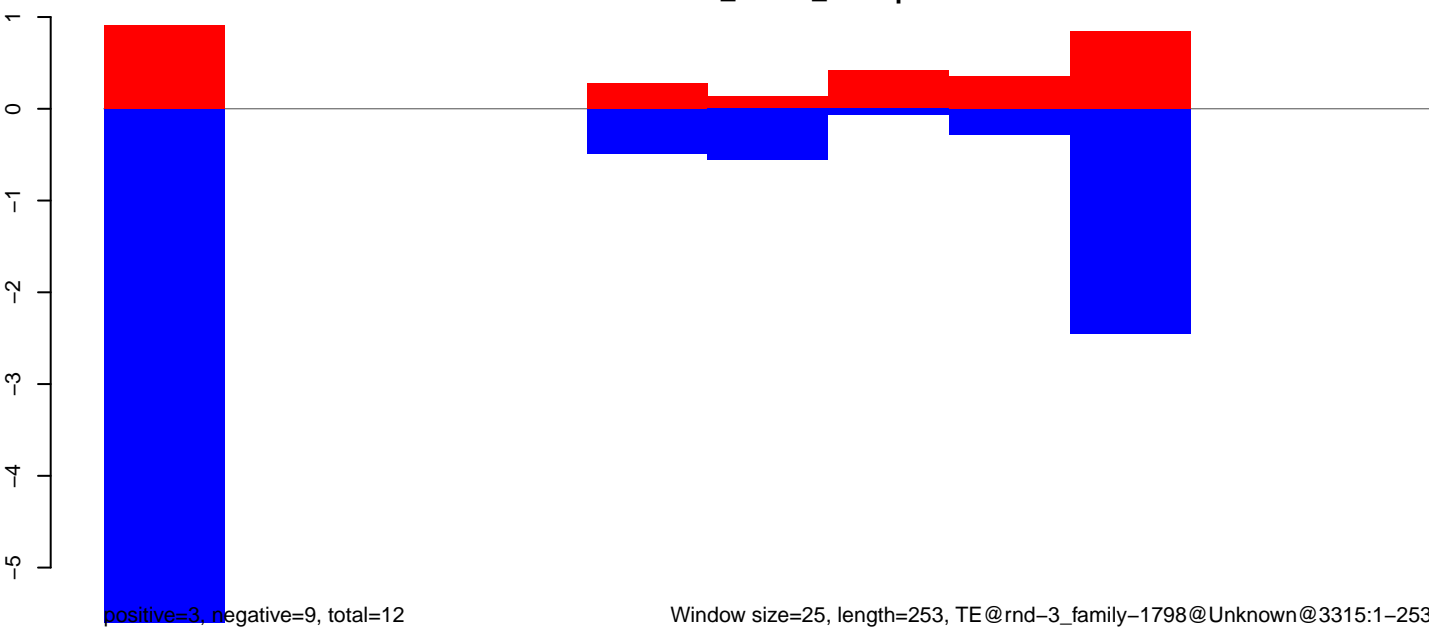
CuQuin_Mcarc_NL.18_23.rep



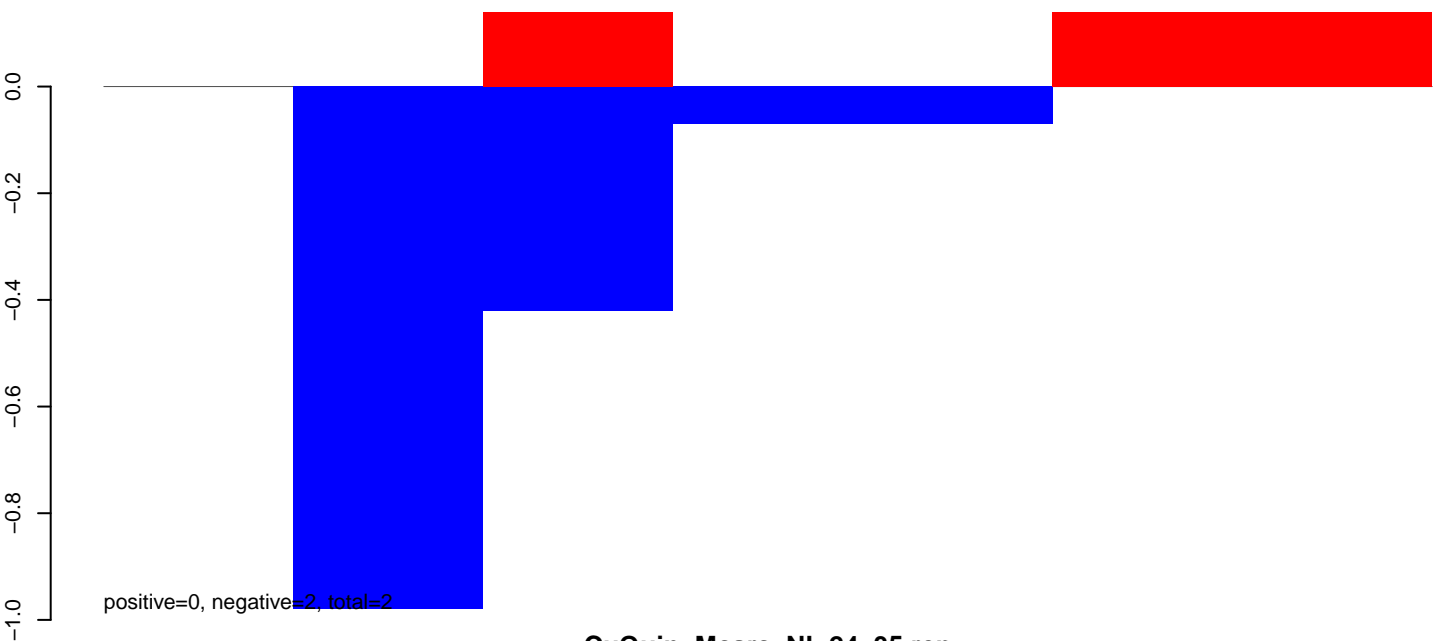
CuQuin_Mcarc_NL.24_35.rep



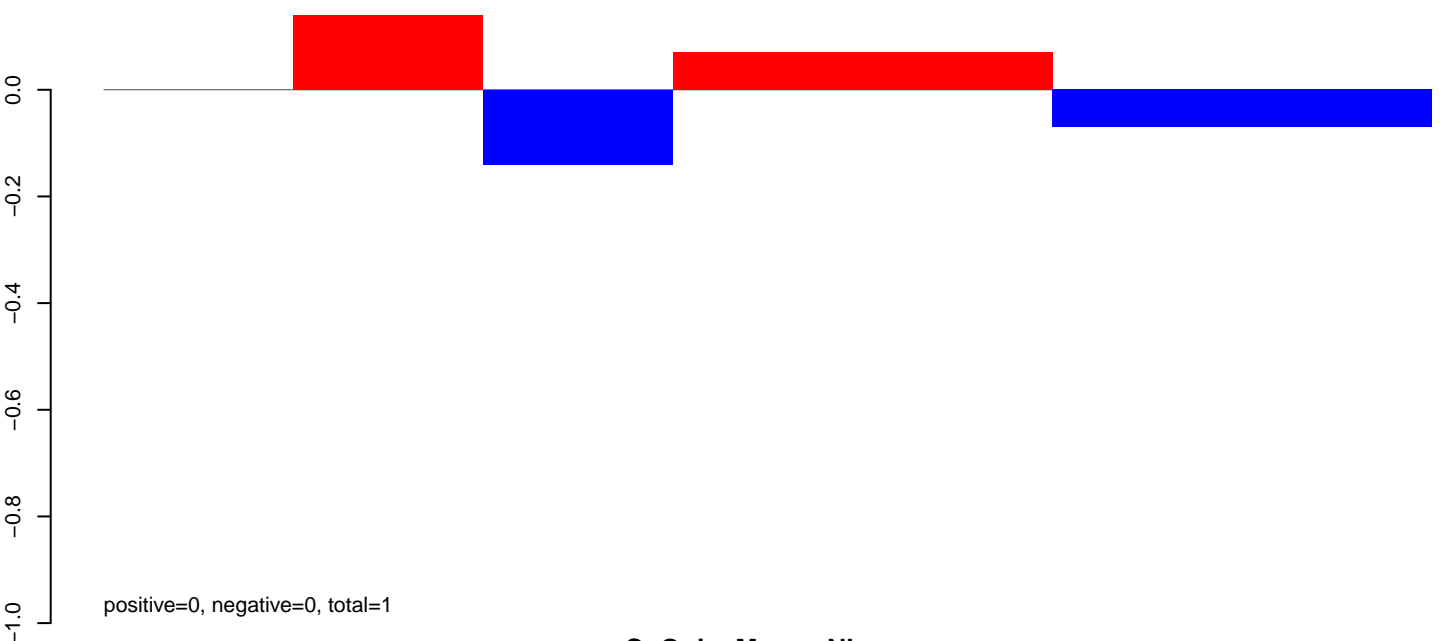
CuQuin_Mcarc_NL.rep



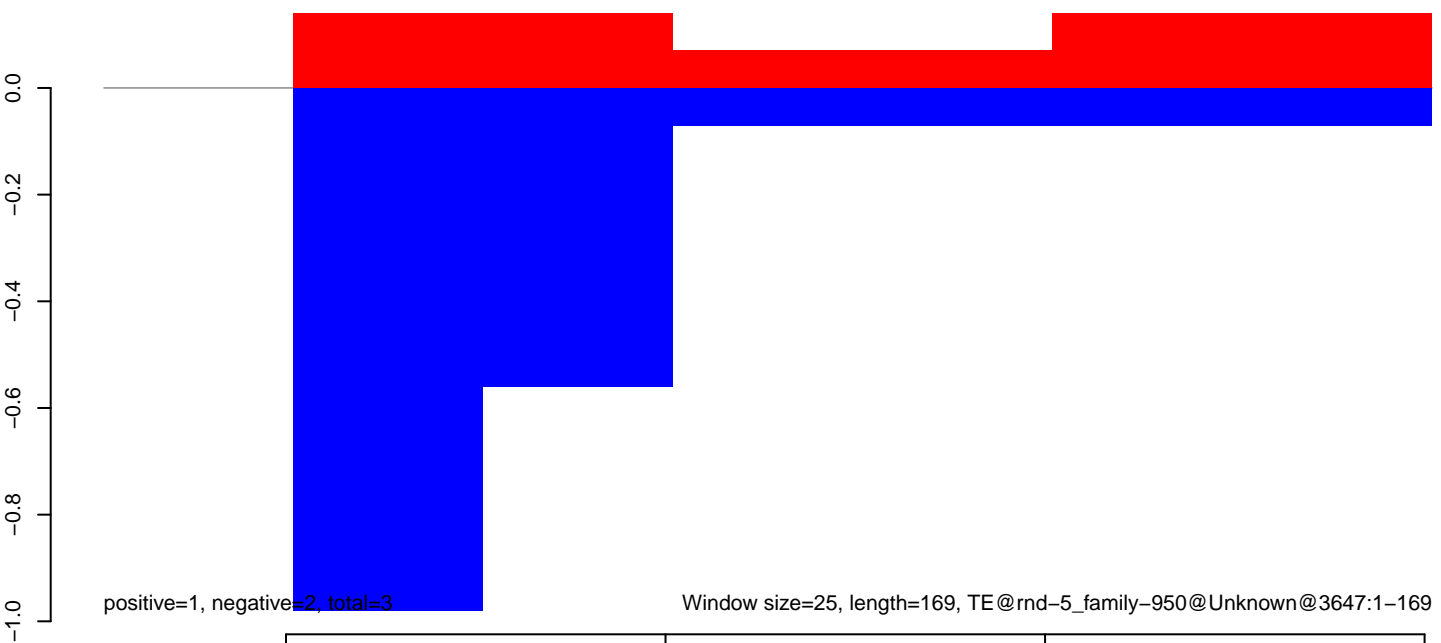
CuQuin_Mcarc_NL.18_23.rep



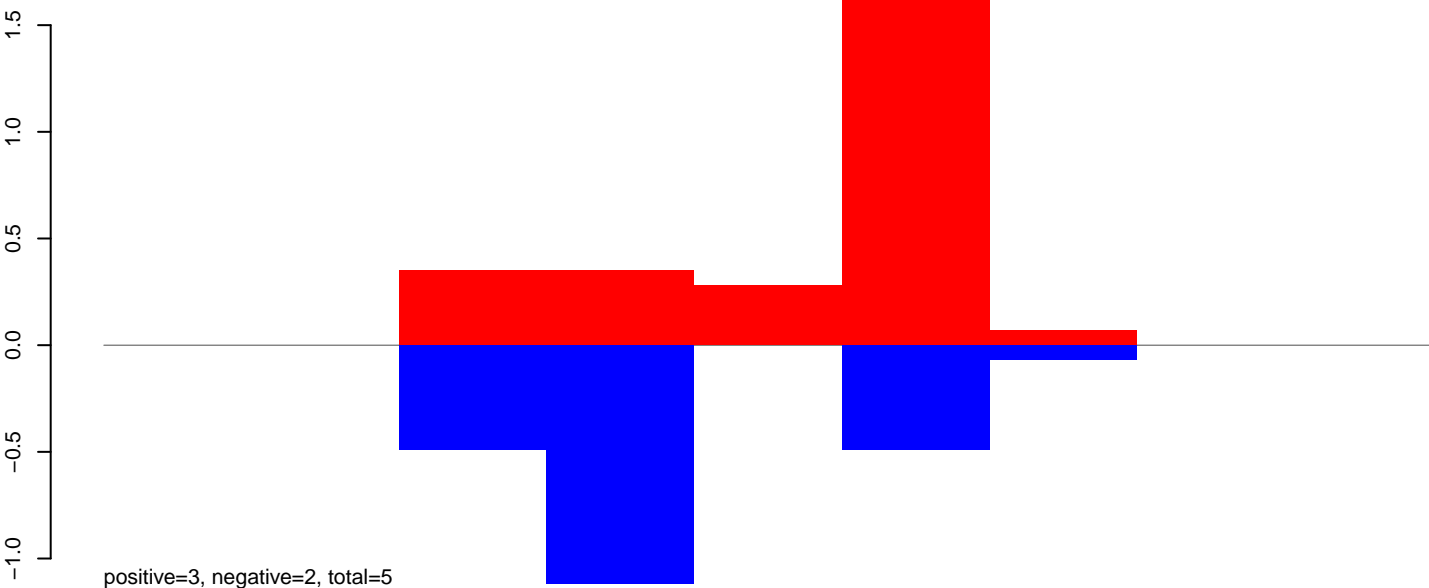
CuQuin_Mcarc_NL.24_35.rep



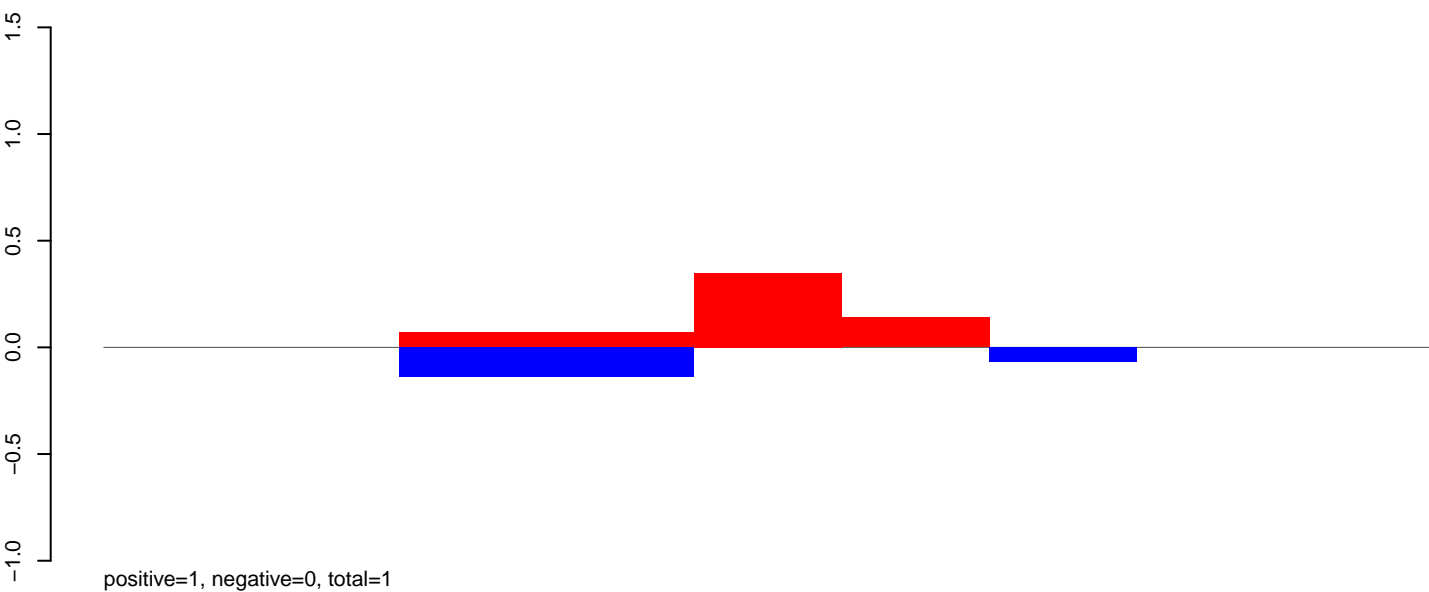
CuQuin_Mcarc_NL.rep



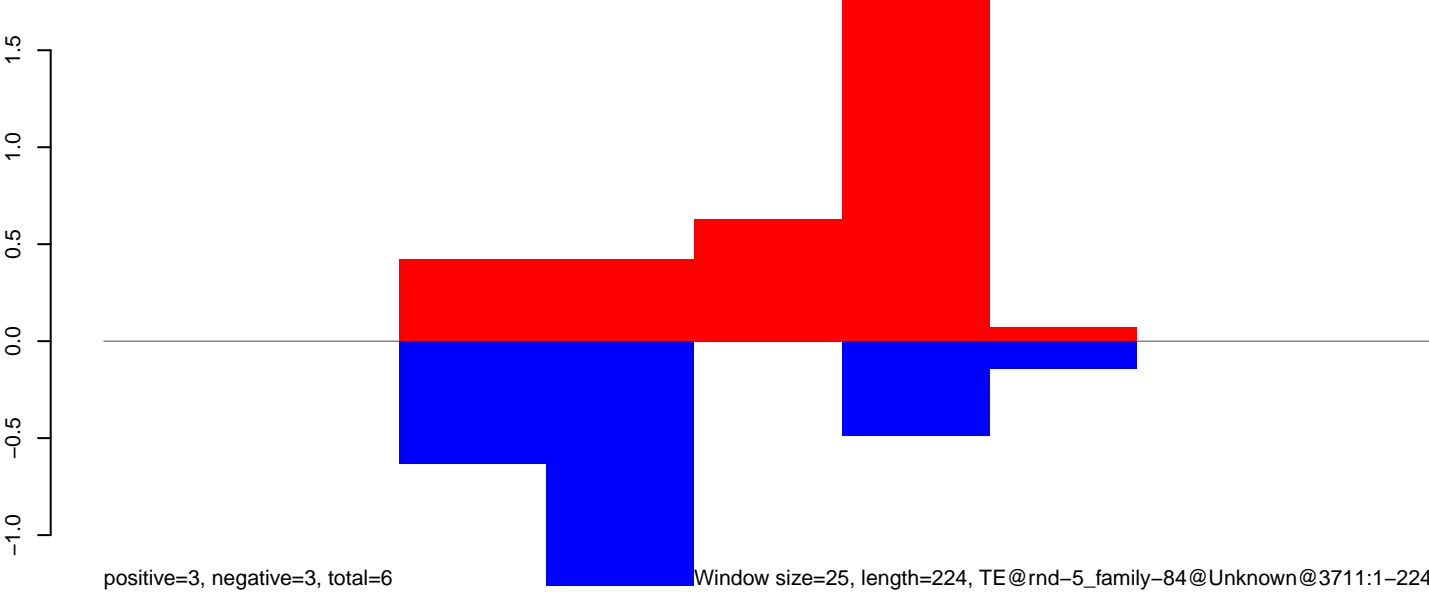
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



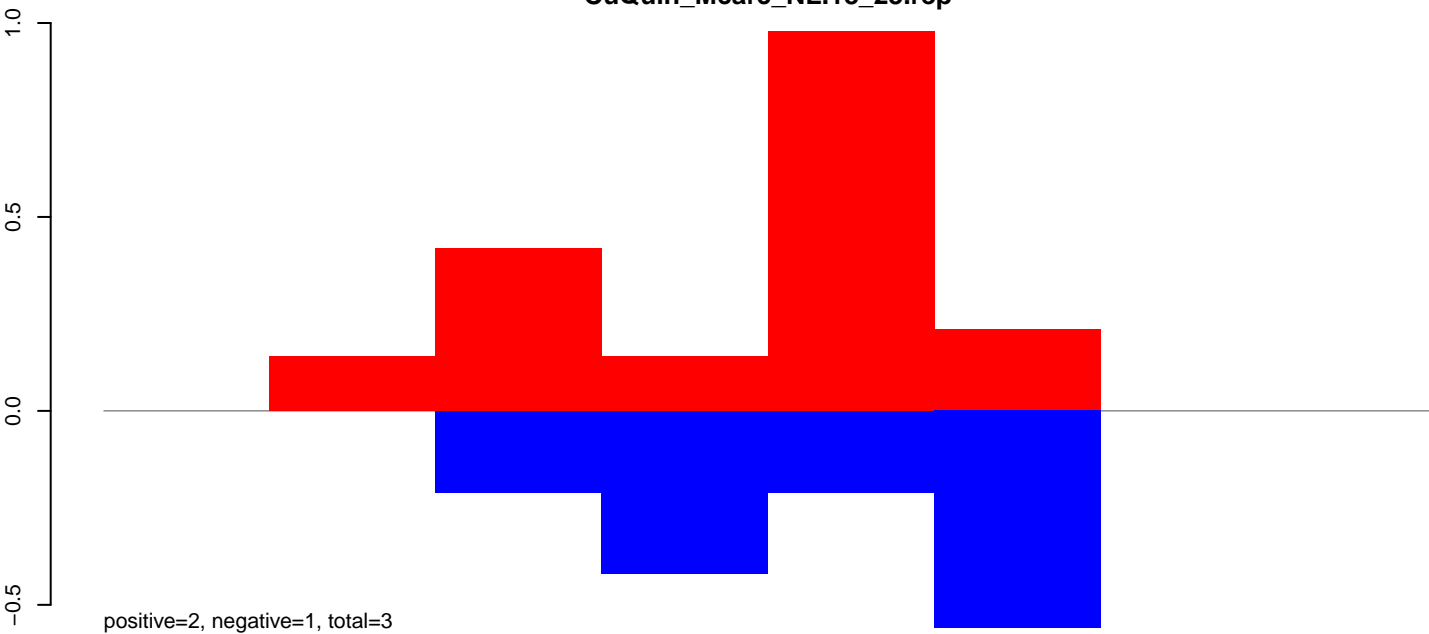
CuQuin_Mcarc_NL.rep



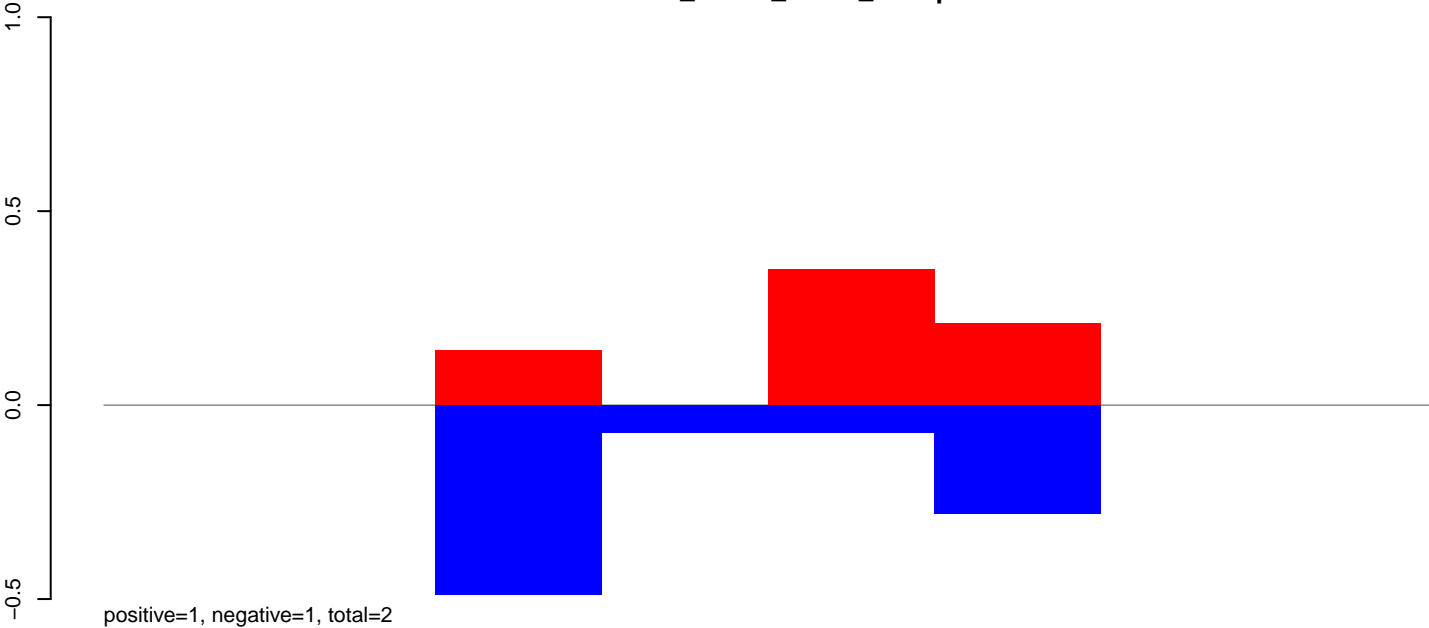
Window size=25, length=224, TE@rnd-5_family-84@Unknown@3711:1-224

50 100 150 200 250

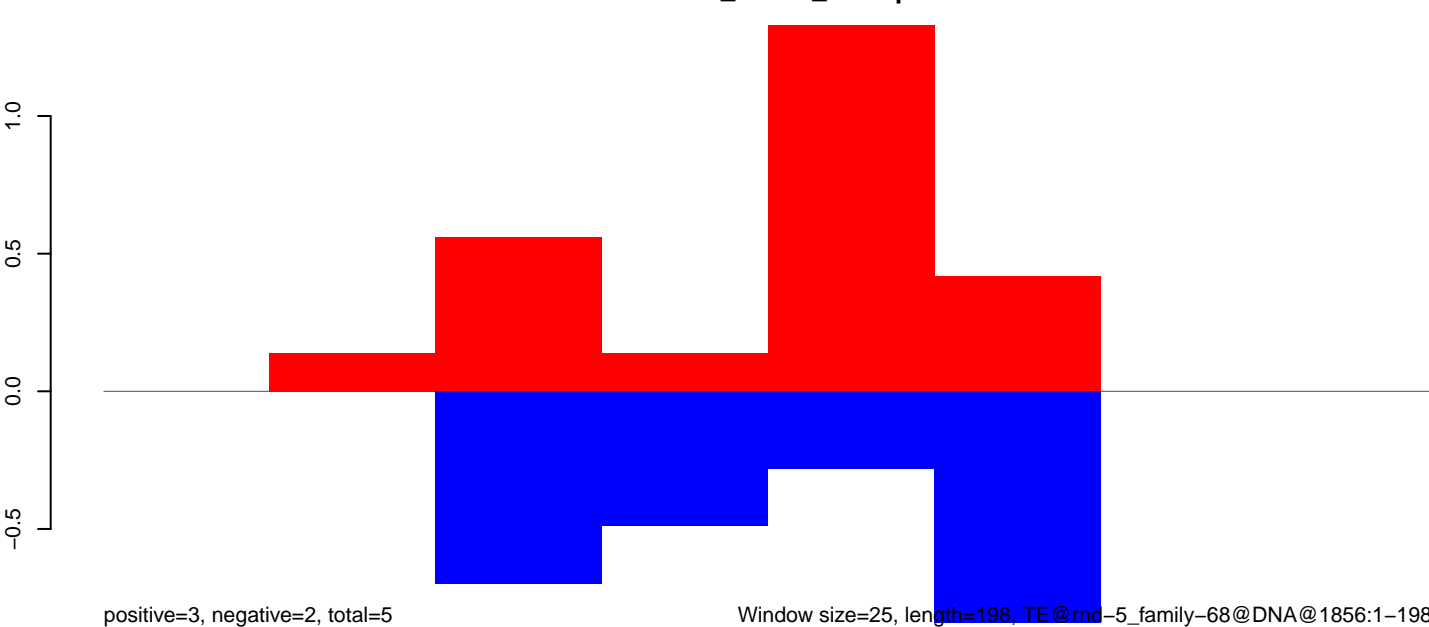
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



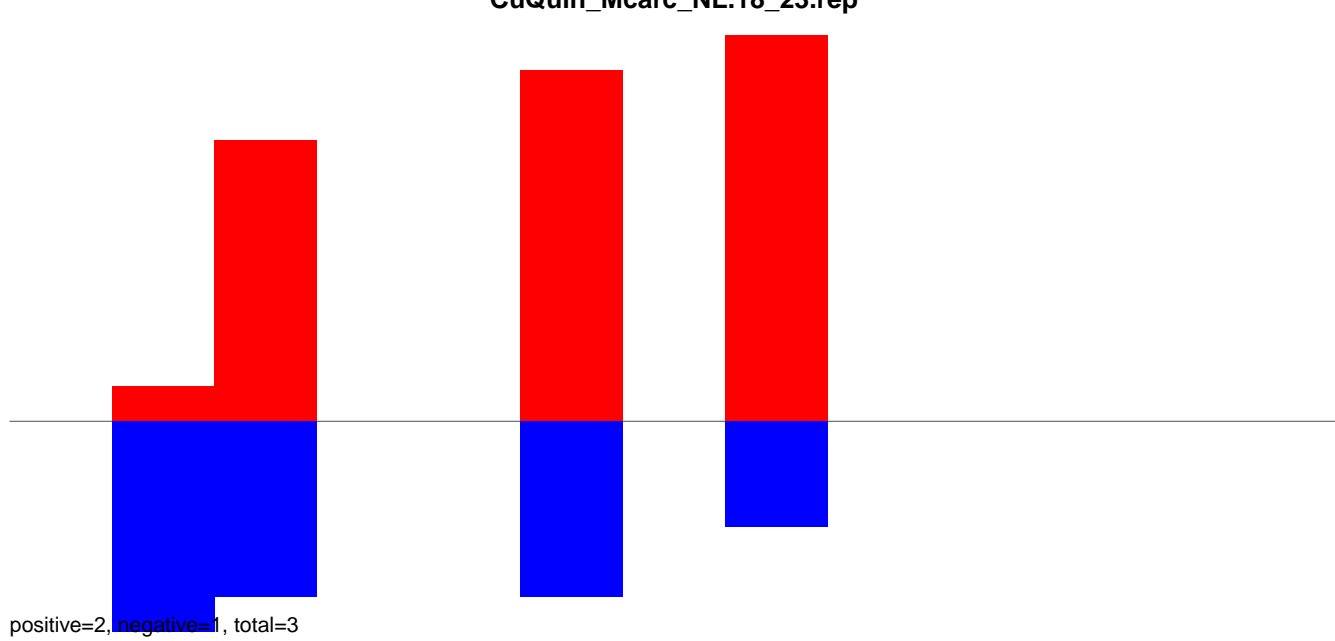
CuQuin_Mcarc_NL.rep



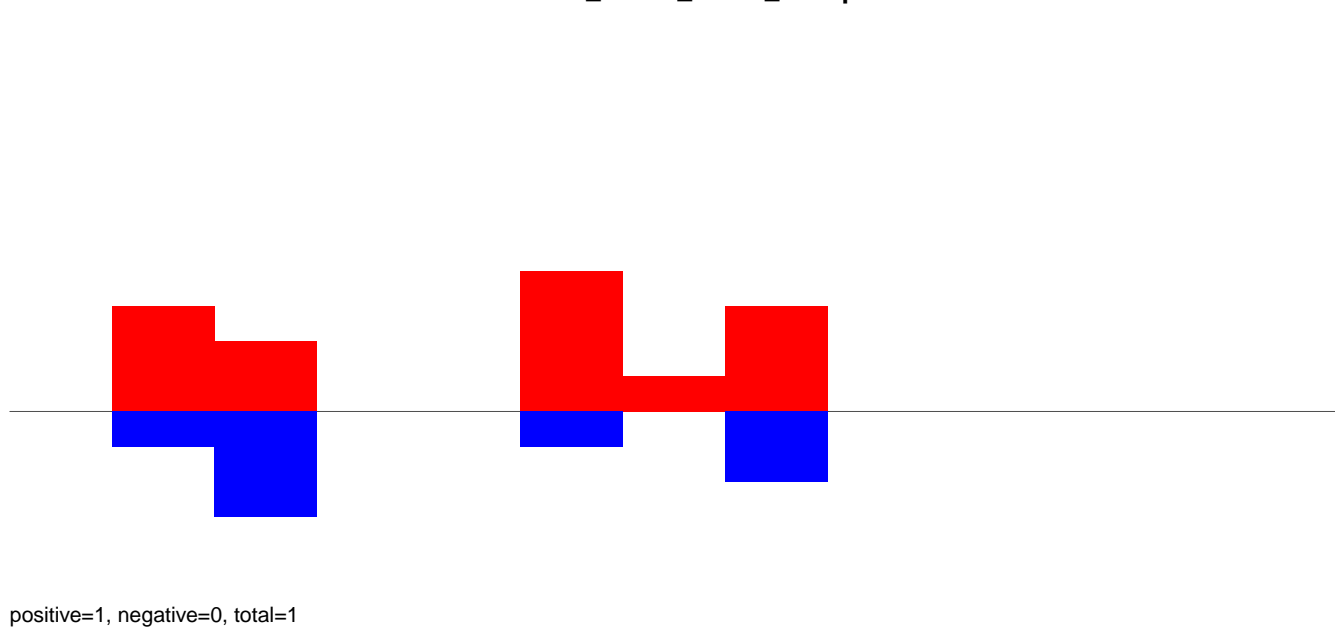
Window size=25, length=198, TE@mc-5_family-68@DNA@1856:1-198

50 100 150 200

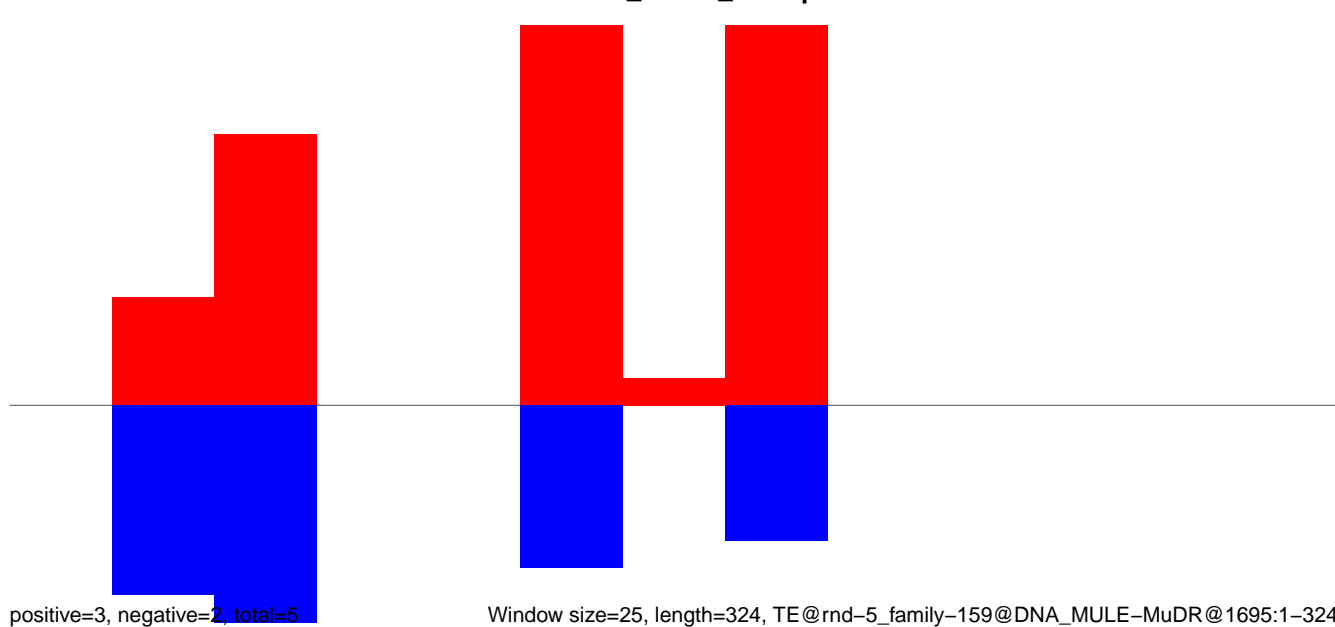
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

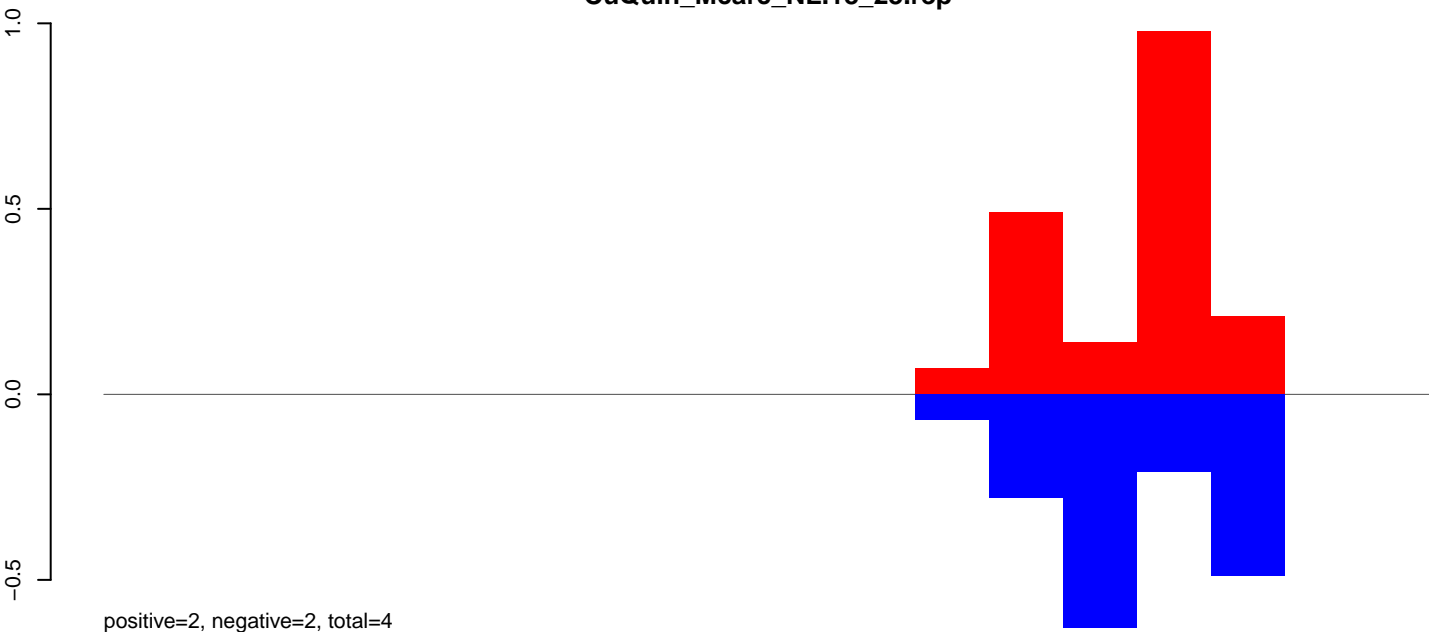


CuQuin_Mcarc_NL.rep

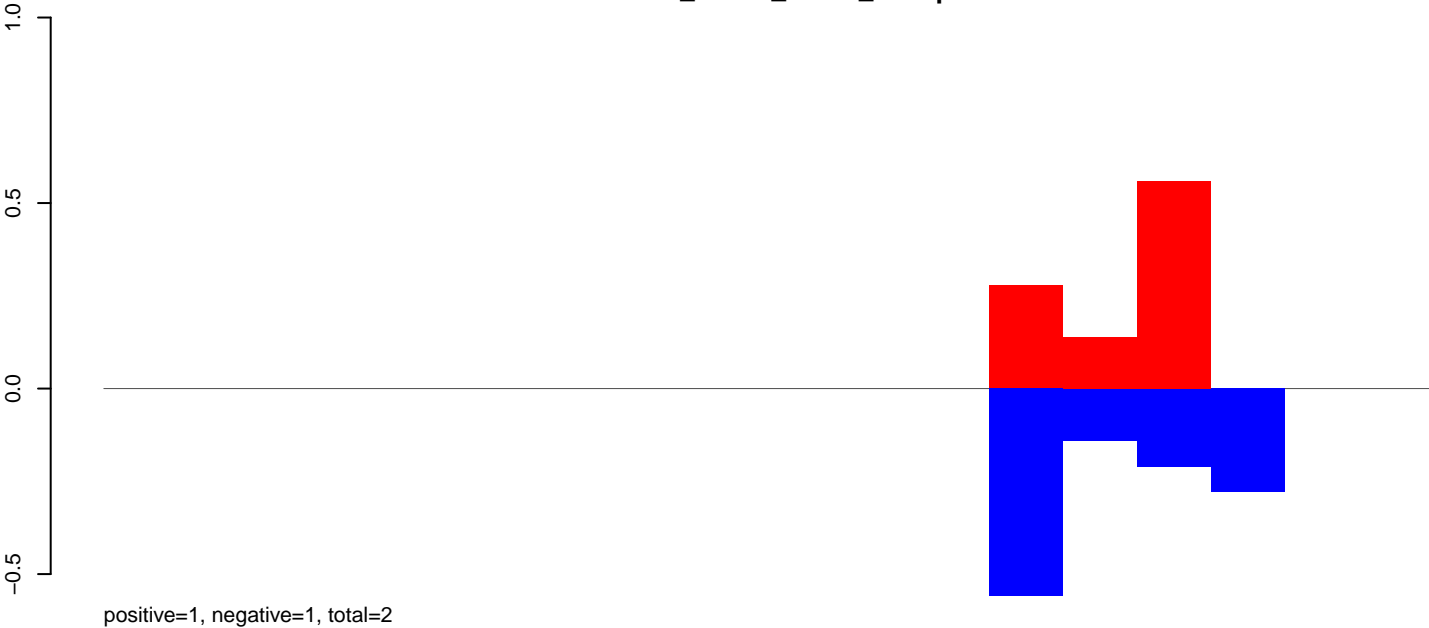


Window size=25, length=324, TE@rnd-5_family-159@DNA_MULE-MuDR@1695:1-324

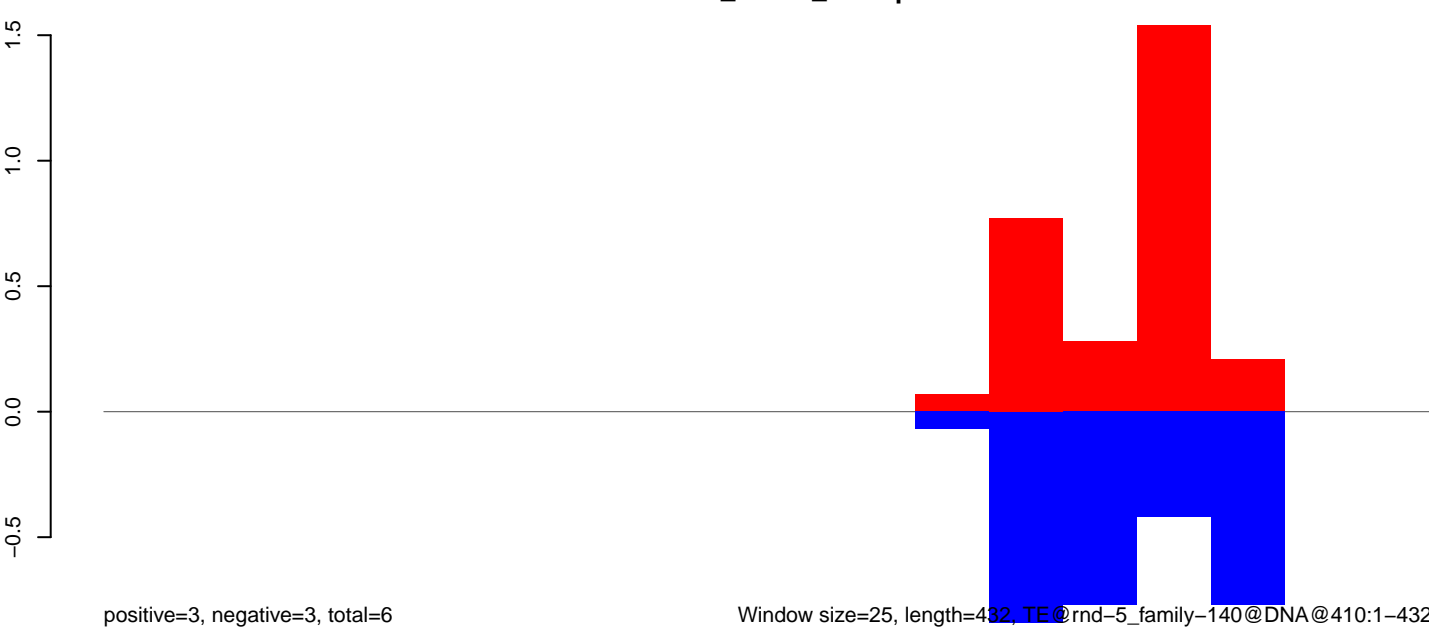
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=432, TE@rnd-5_family-140@DNA@410:1-432

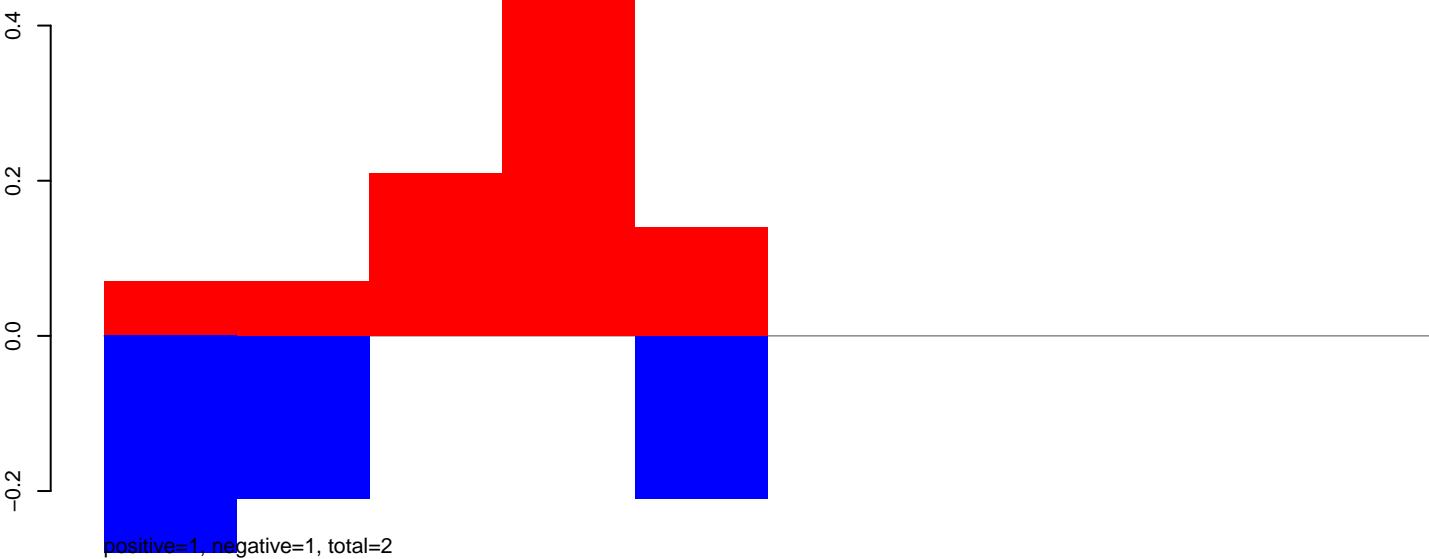
100

200

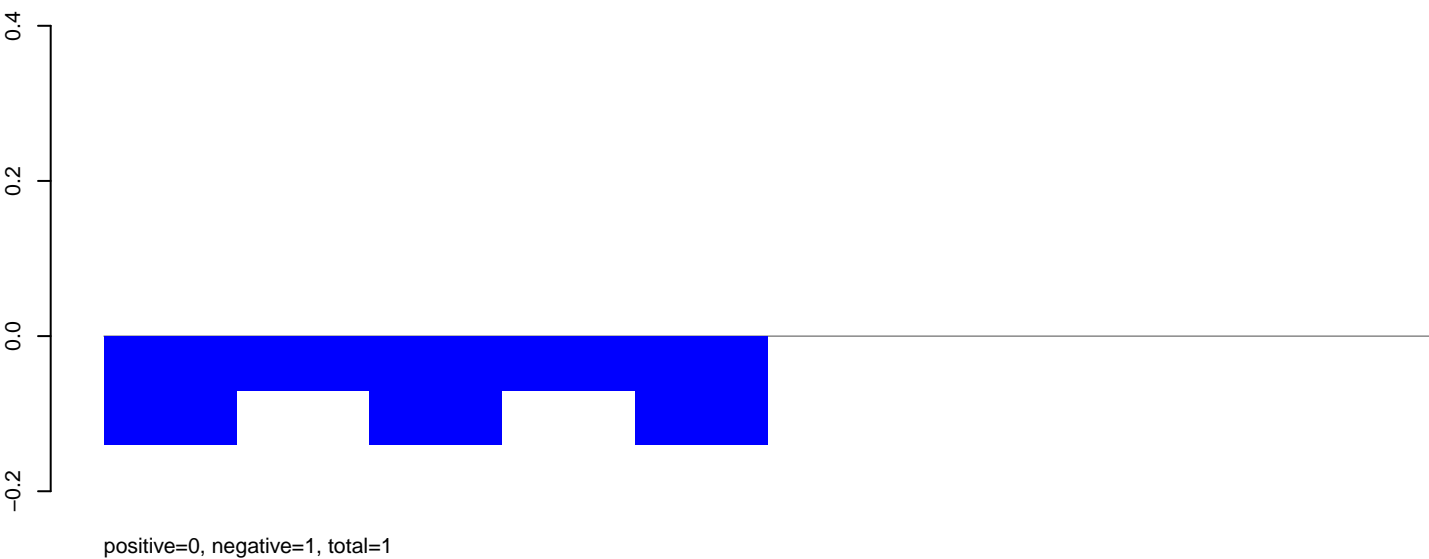
300

400

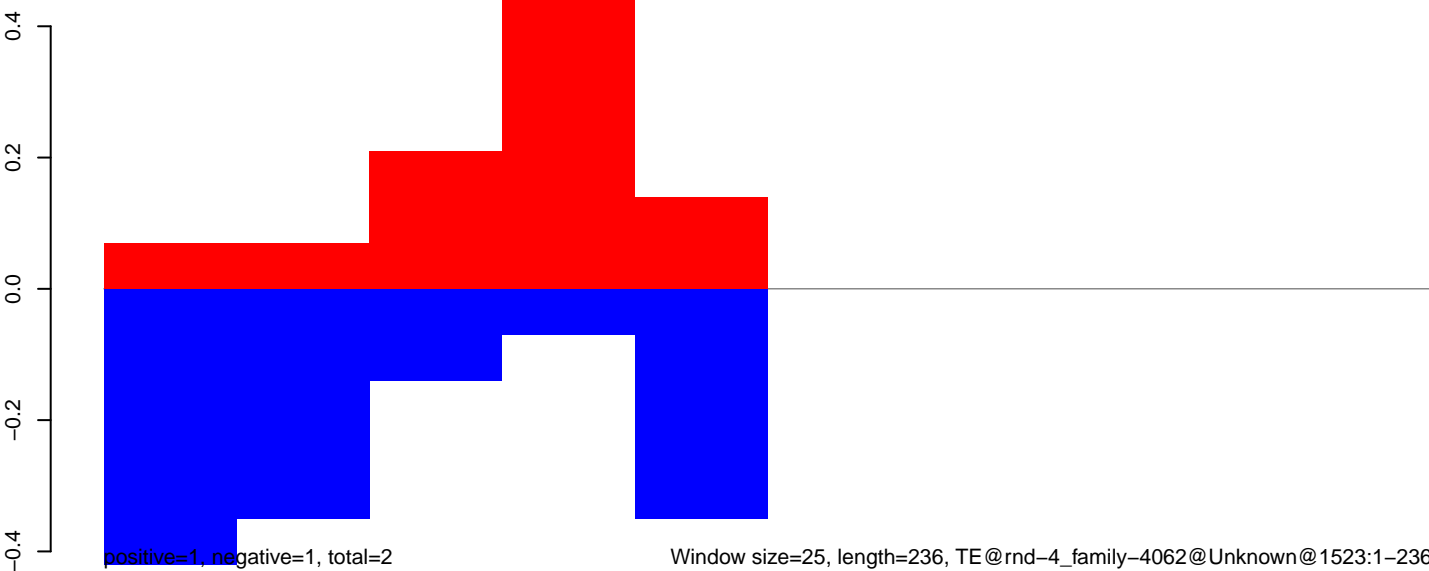
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

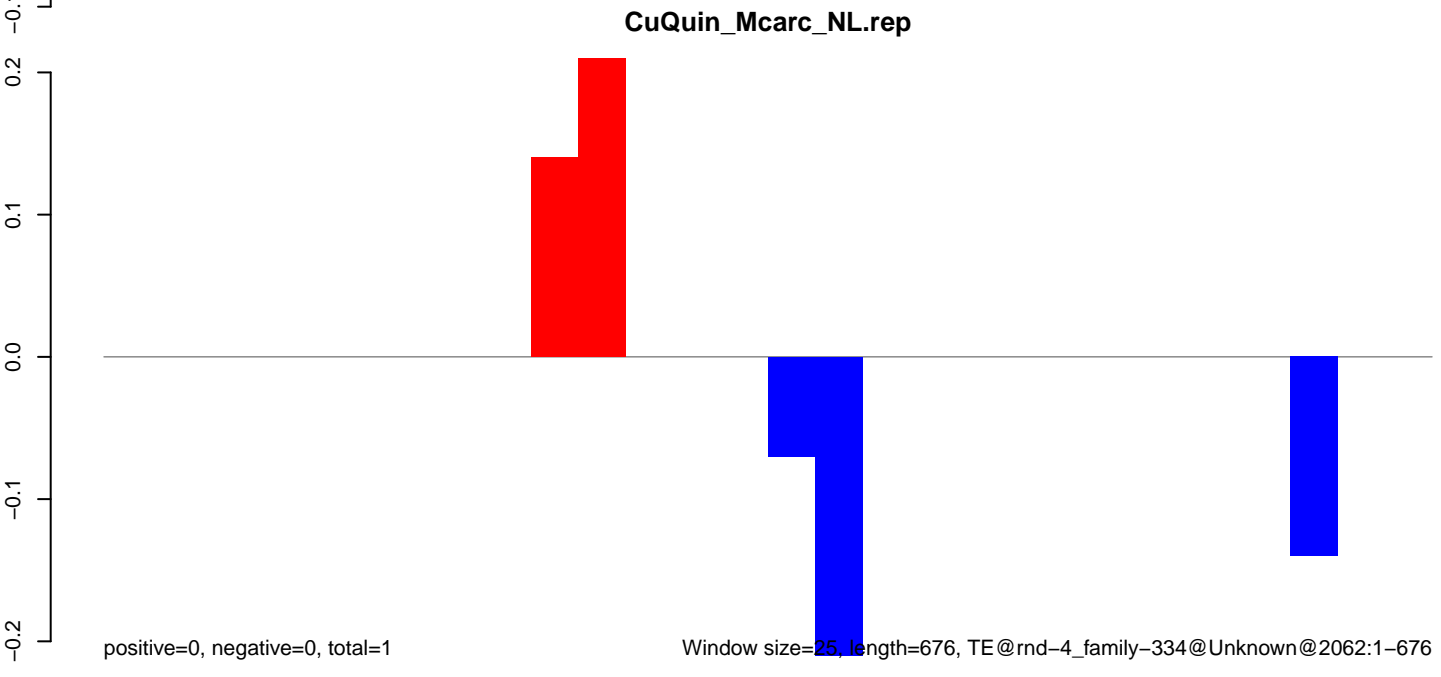
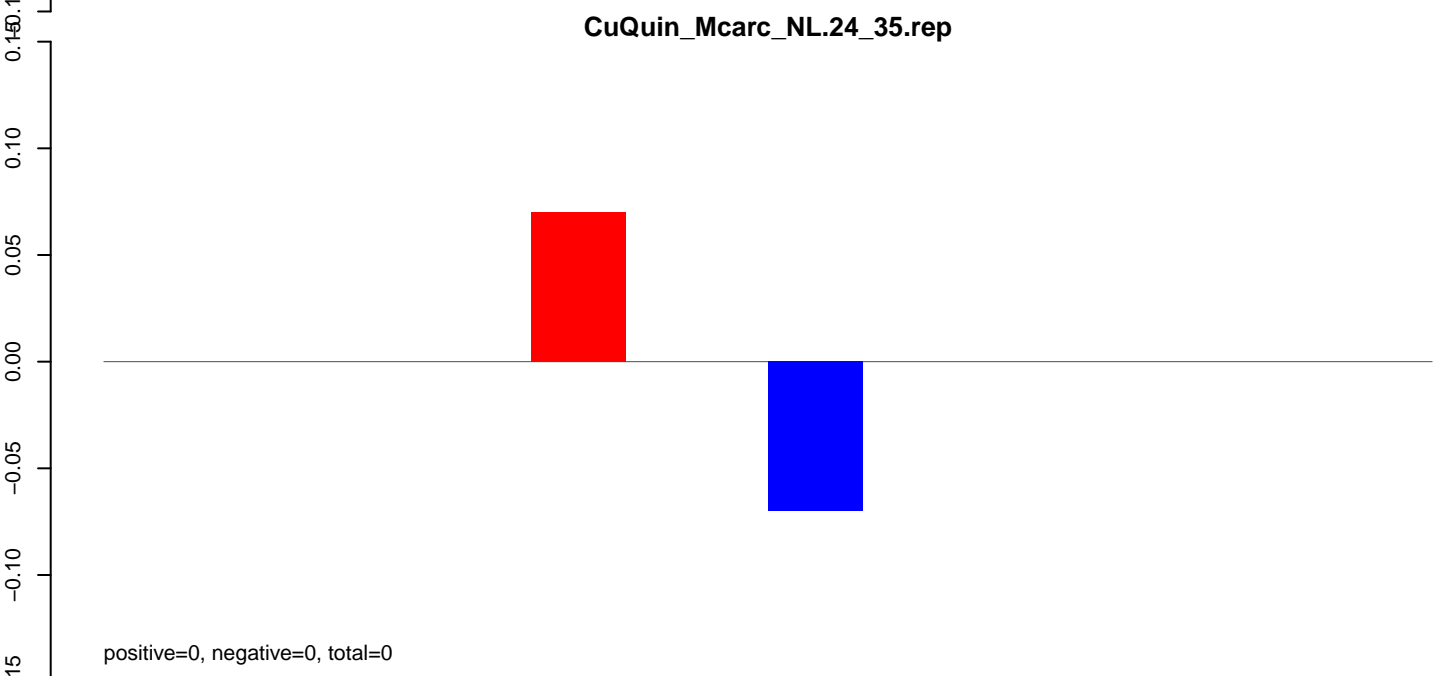
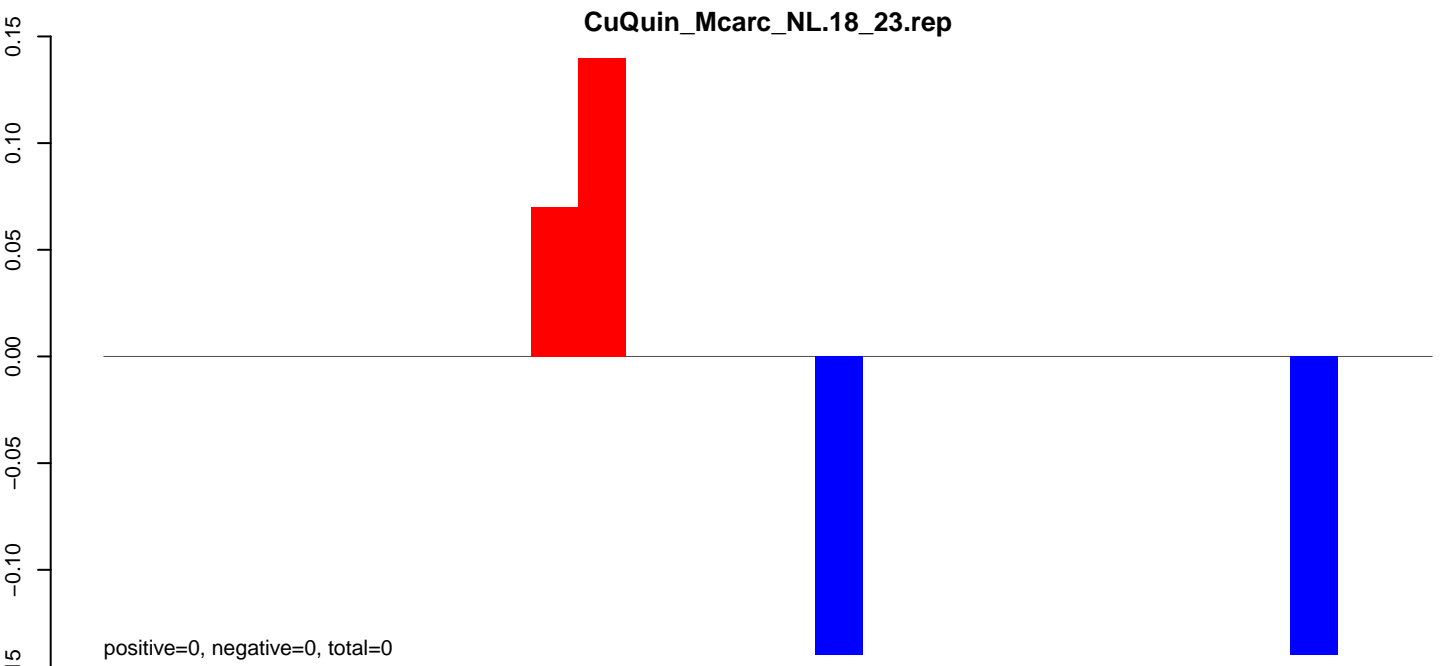


CuQuin_Mcarc_NL.rep



Window size=25, length=236, TE@rnd-4_family-4062@Unknown@1523:1-236

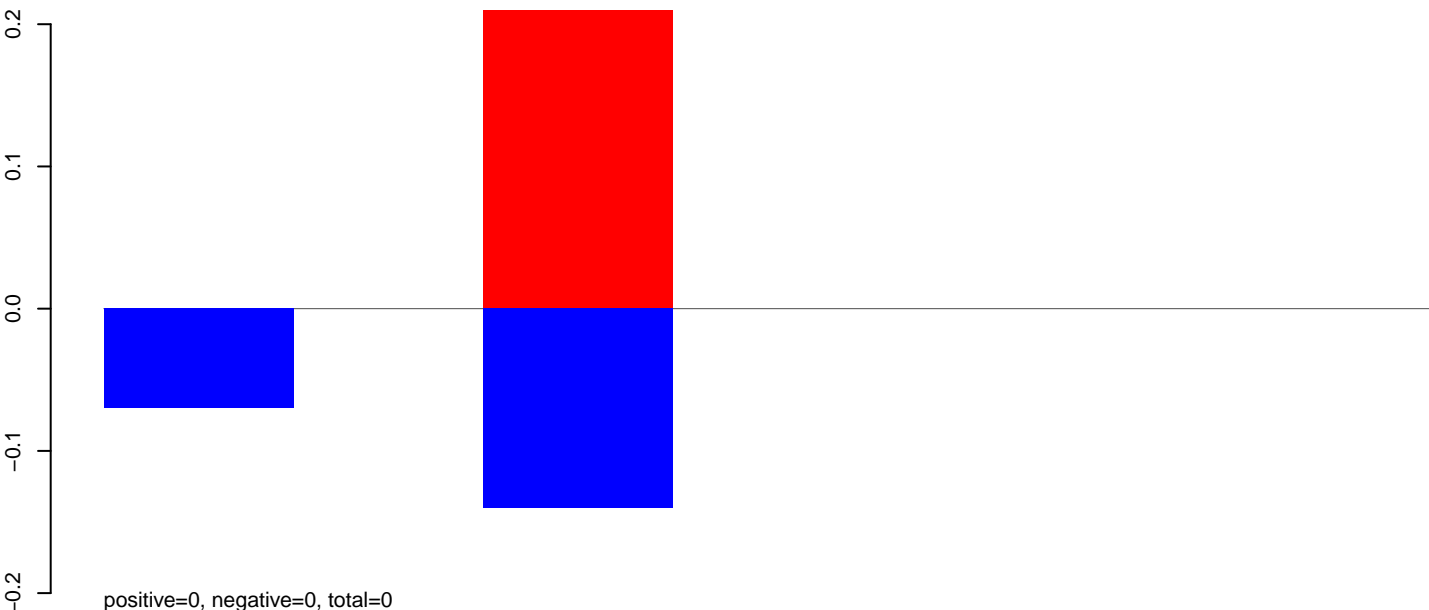
50 100 150 200 250



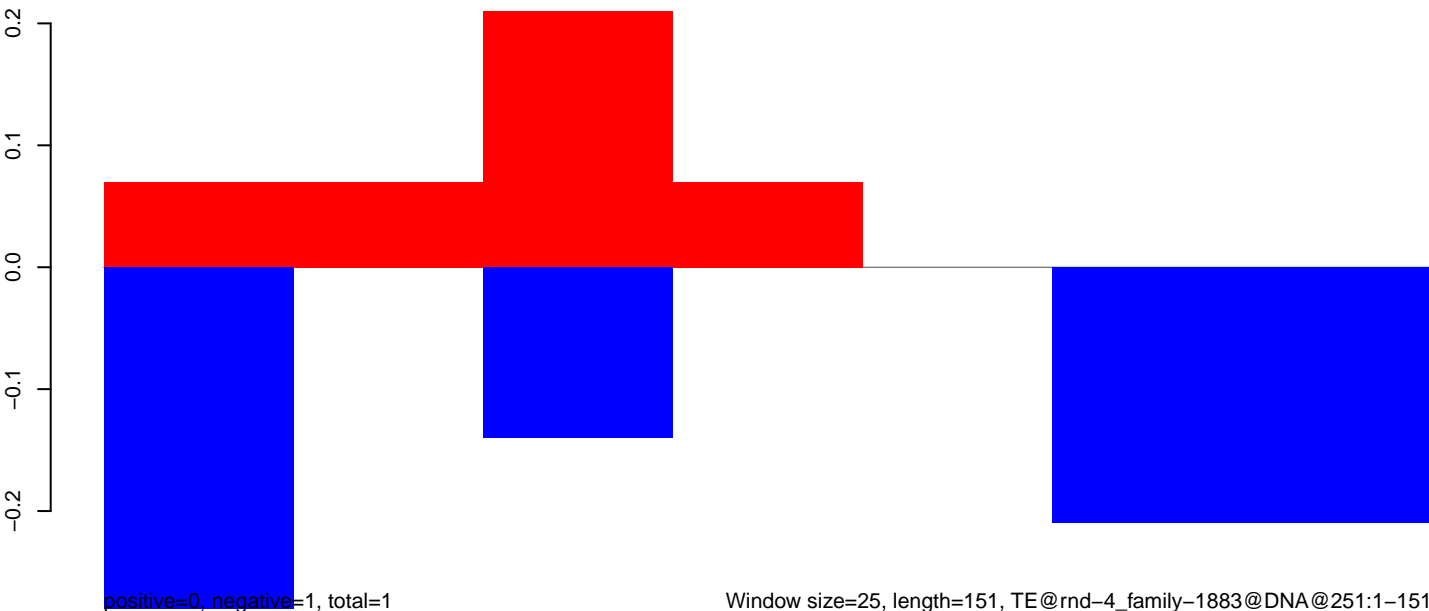
CuQuin_Mcarc_NL.18_23.rep



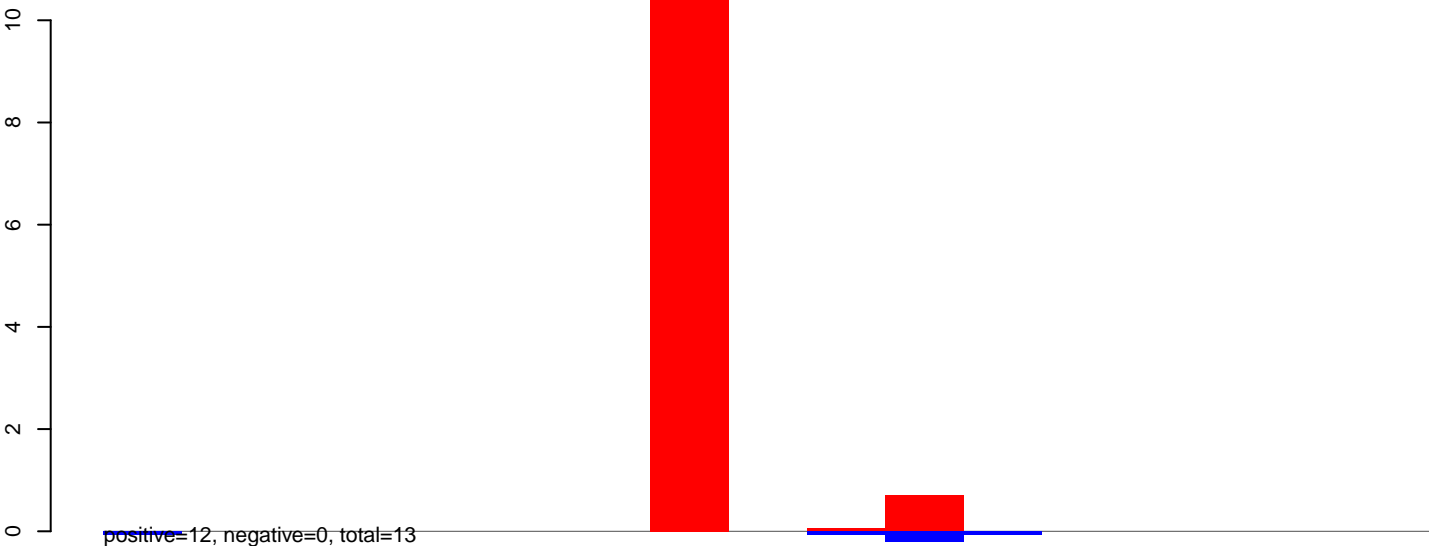
CuQuin_Mcarc_NL.24_35.rep



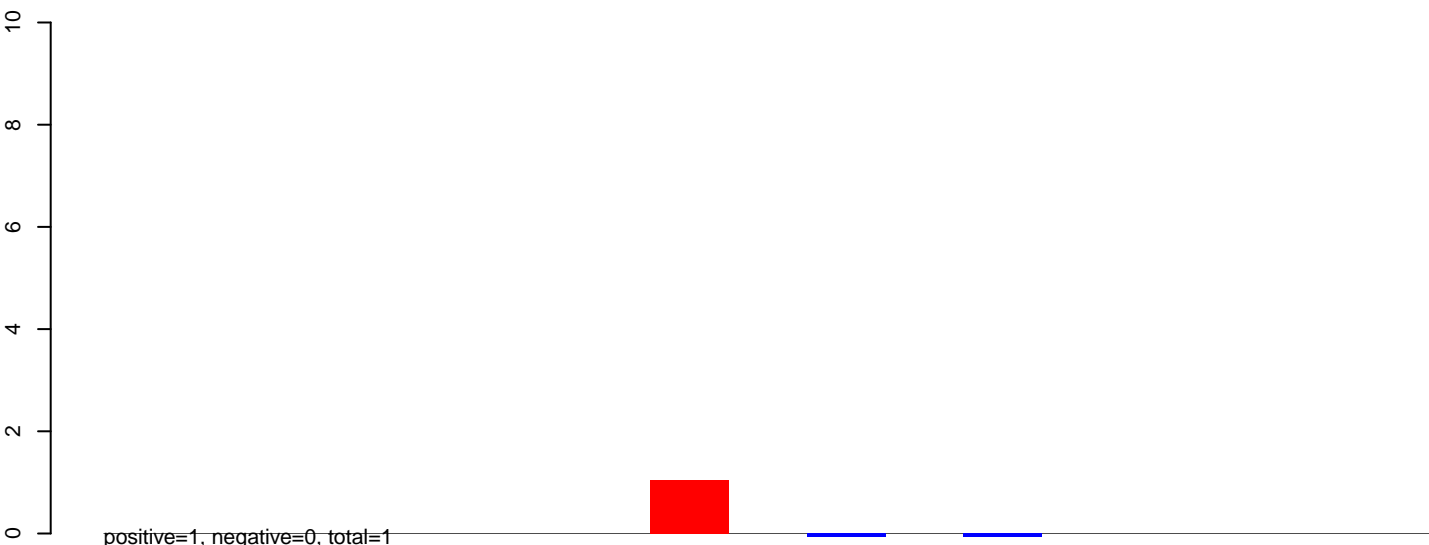
CuQuin_Mcarc_NL.rep



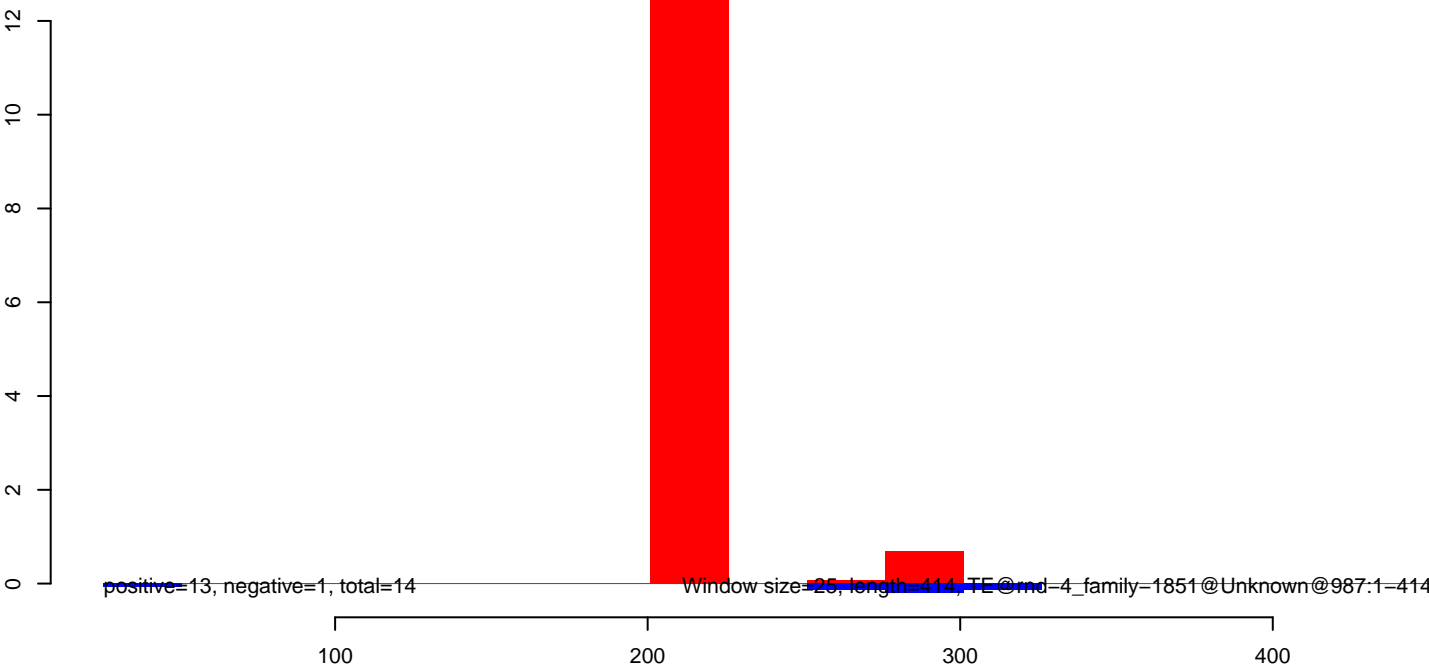
CuQuin_Mcarc_NL.18_23.rep



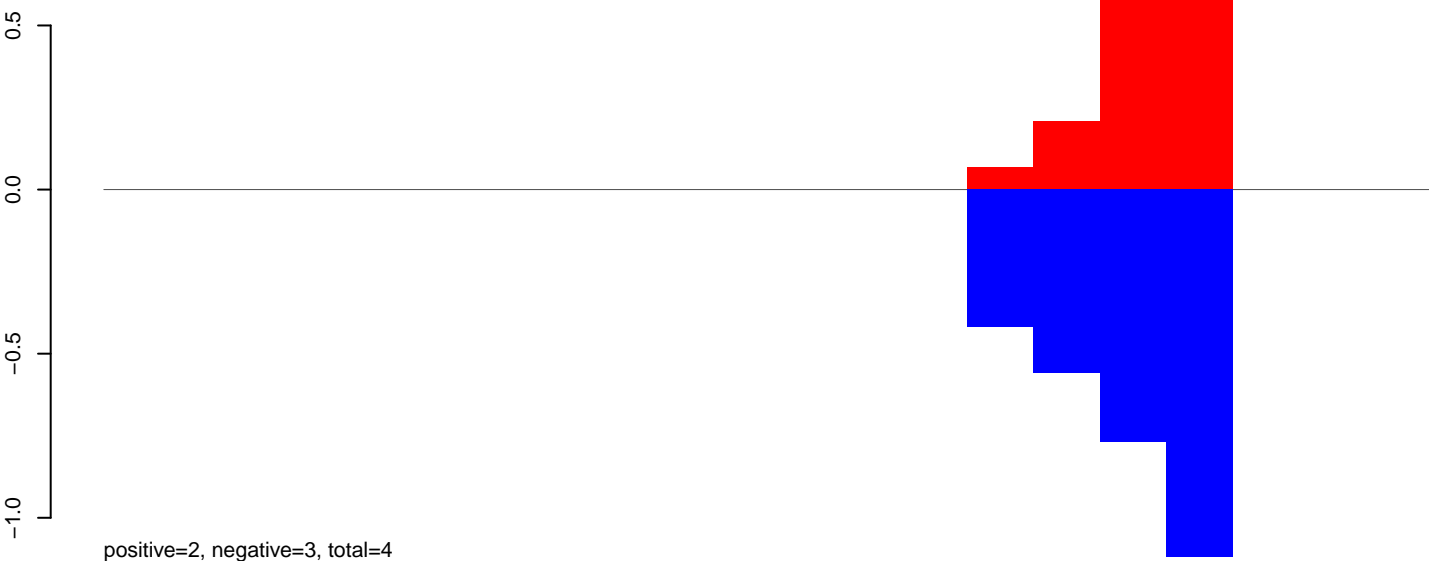
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



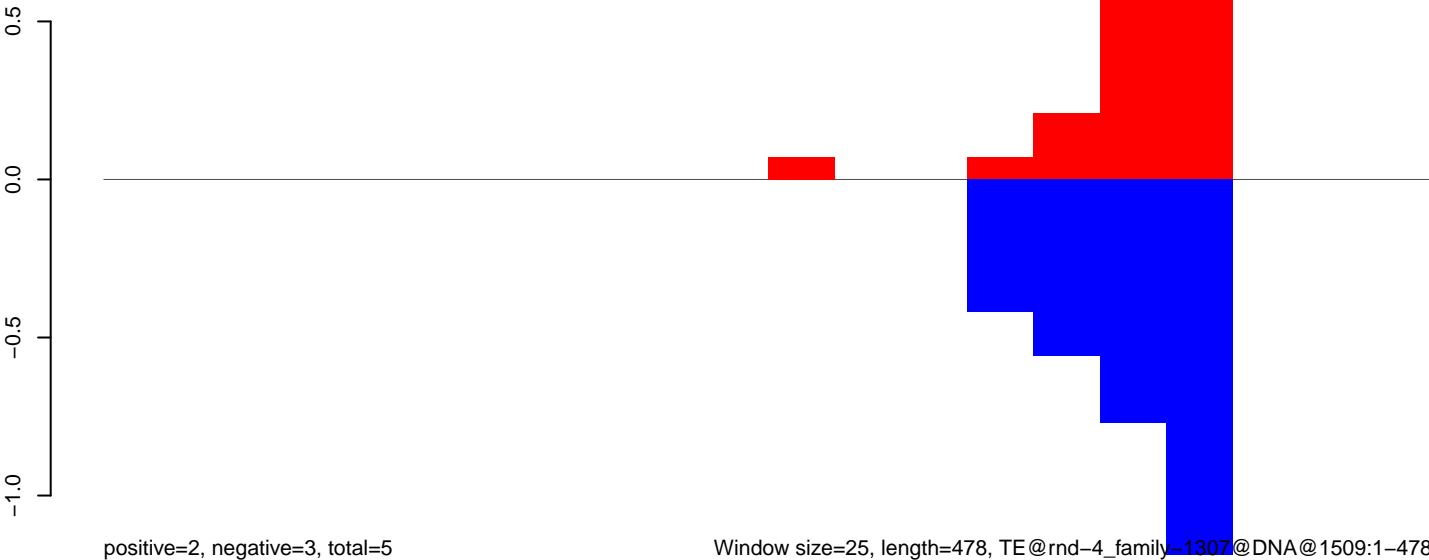
CuQuin_Mcarc_NL.18_23.rep



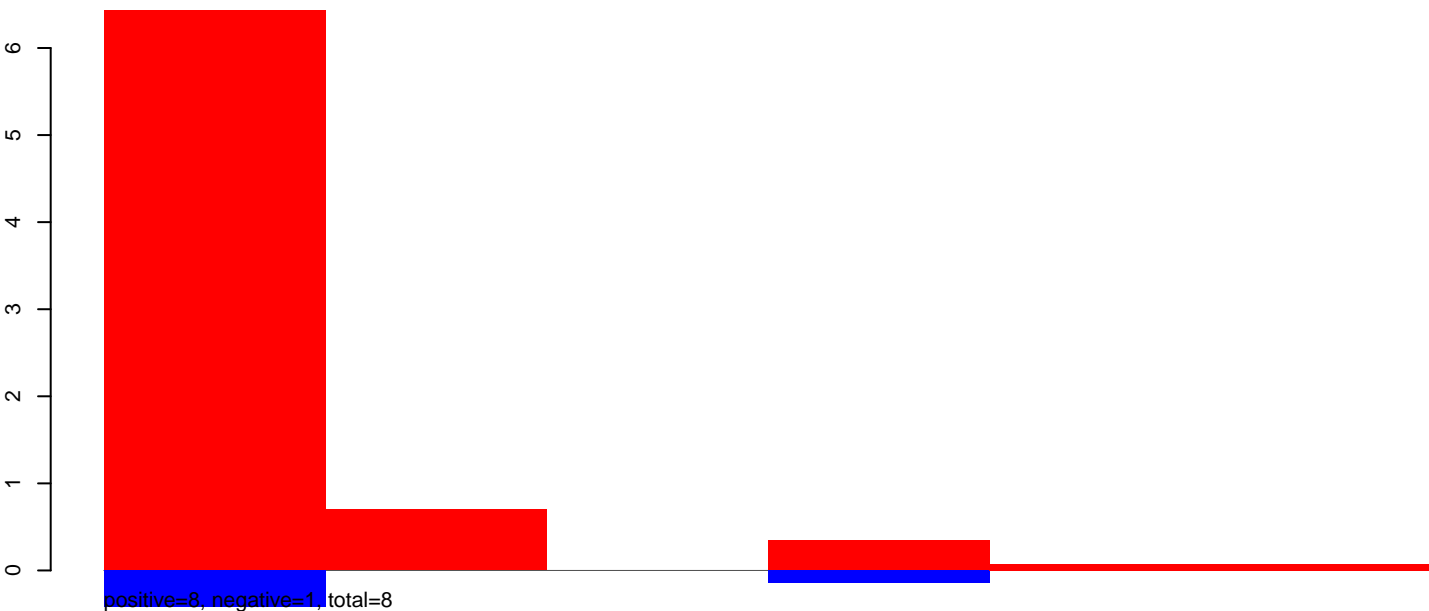
CuQuin_Mcarc_NL.24_35.rep



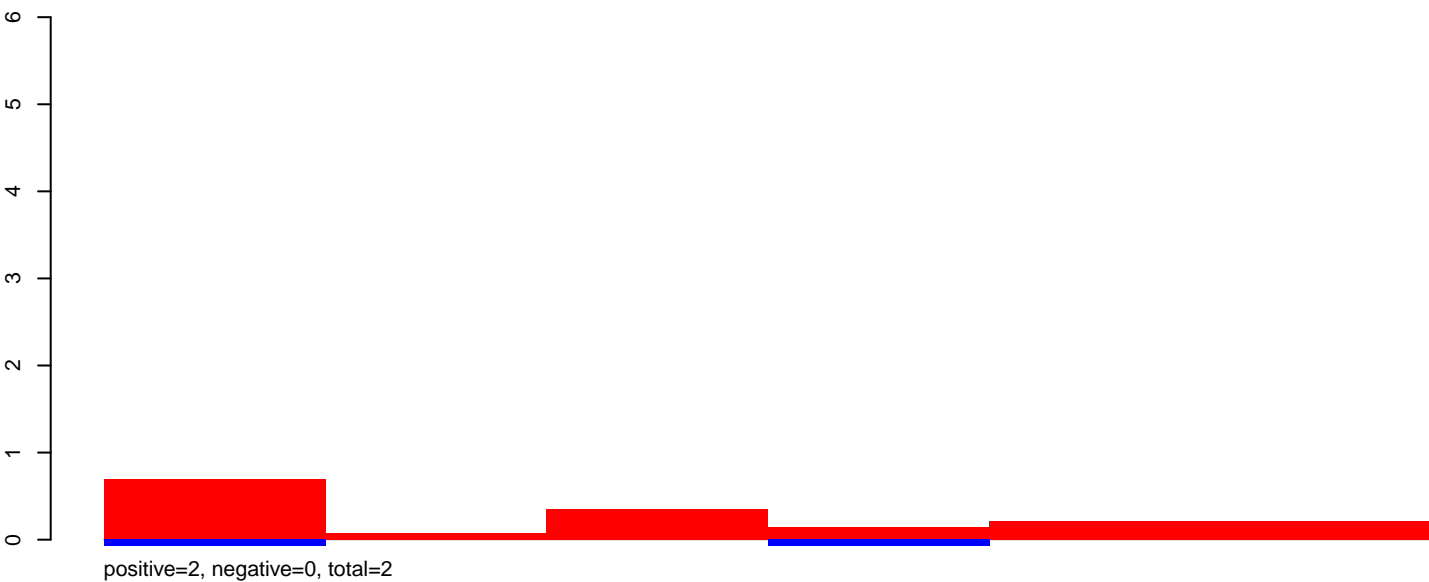
CuQuin_Mcarc_NL.rep



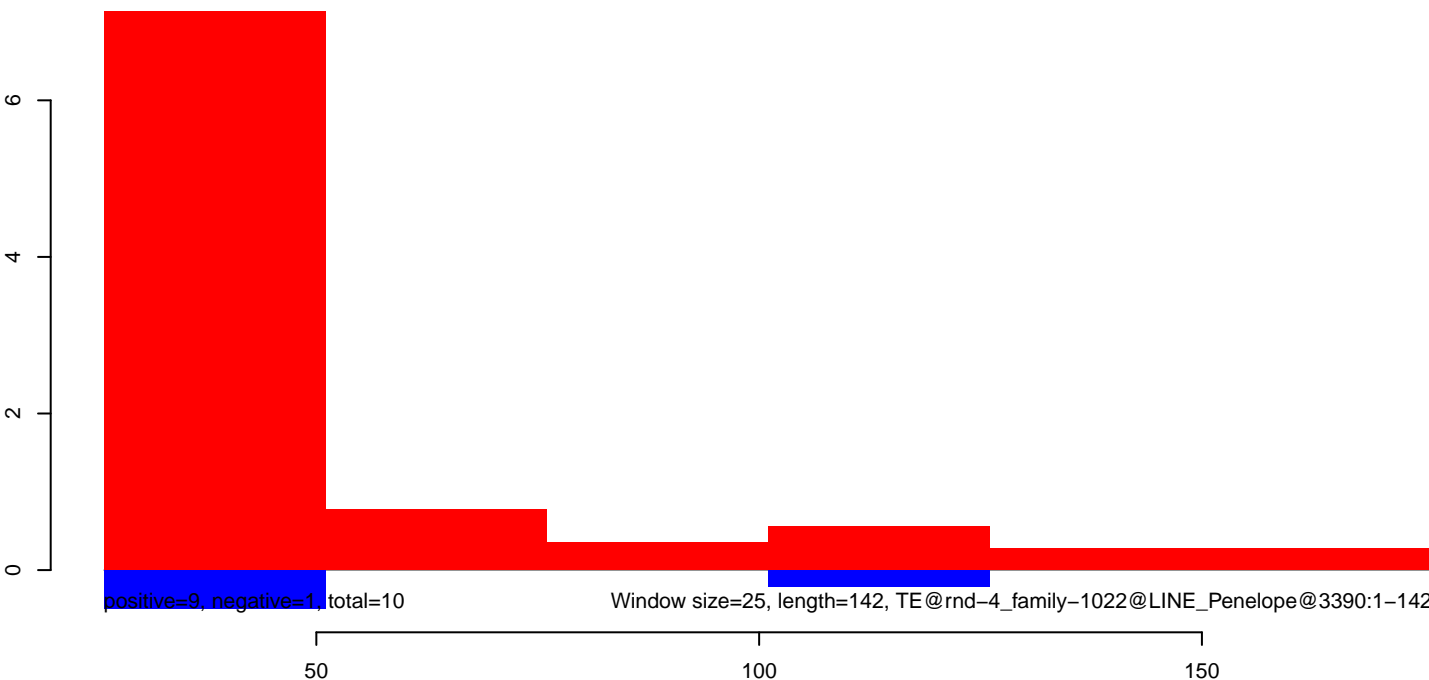
CuQuin_Mcarc_NL.18_23.rep



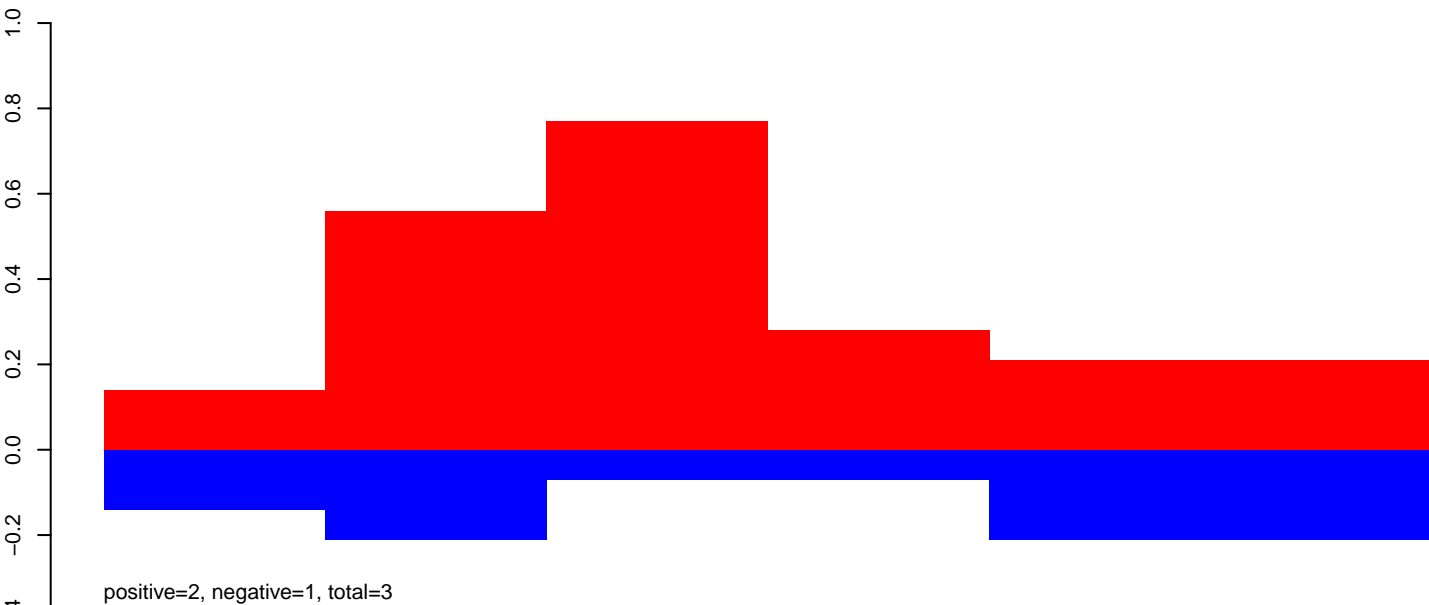
CuQuin_Mcarc_NL.24_35.rep



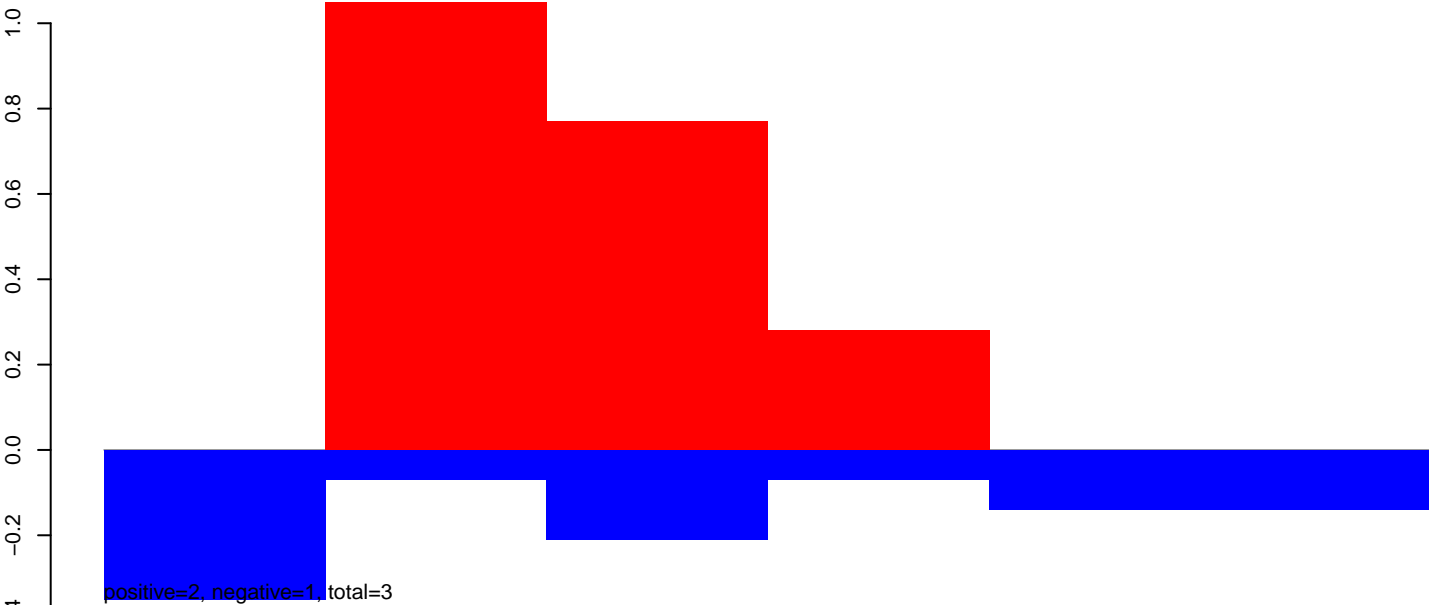
CuQuin_Mcarc_NL.rep



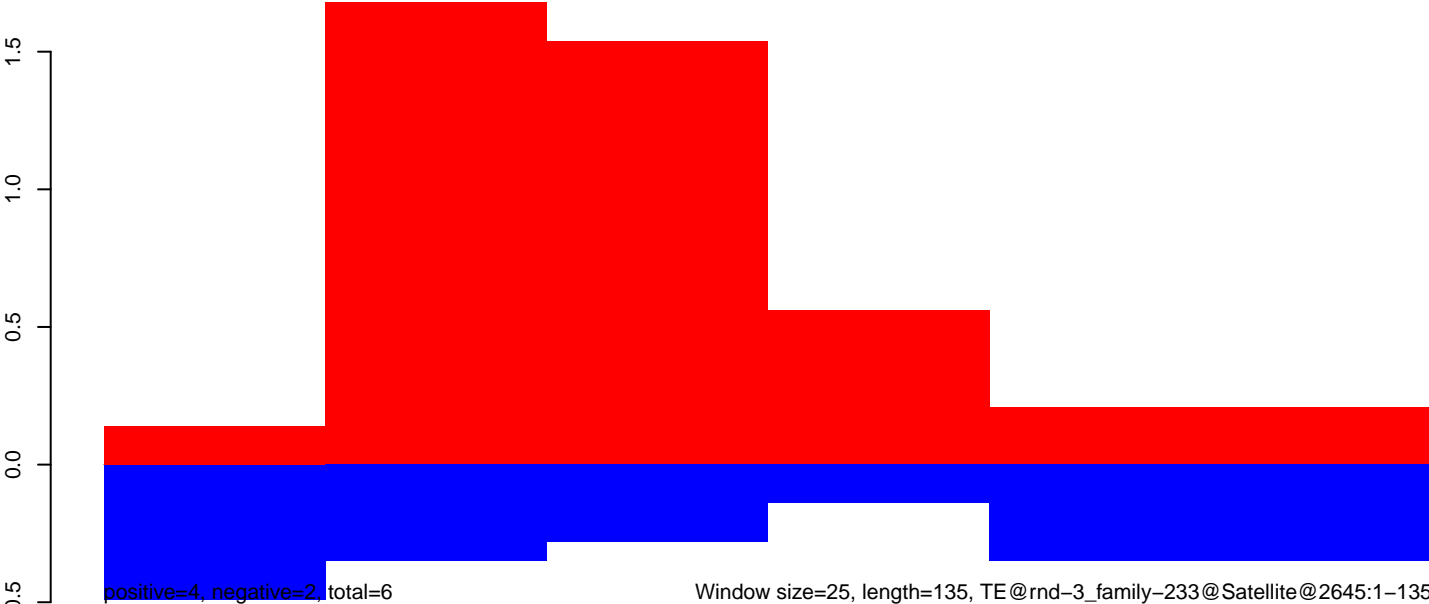
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep

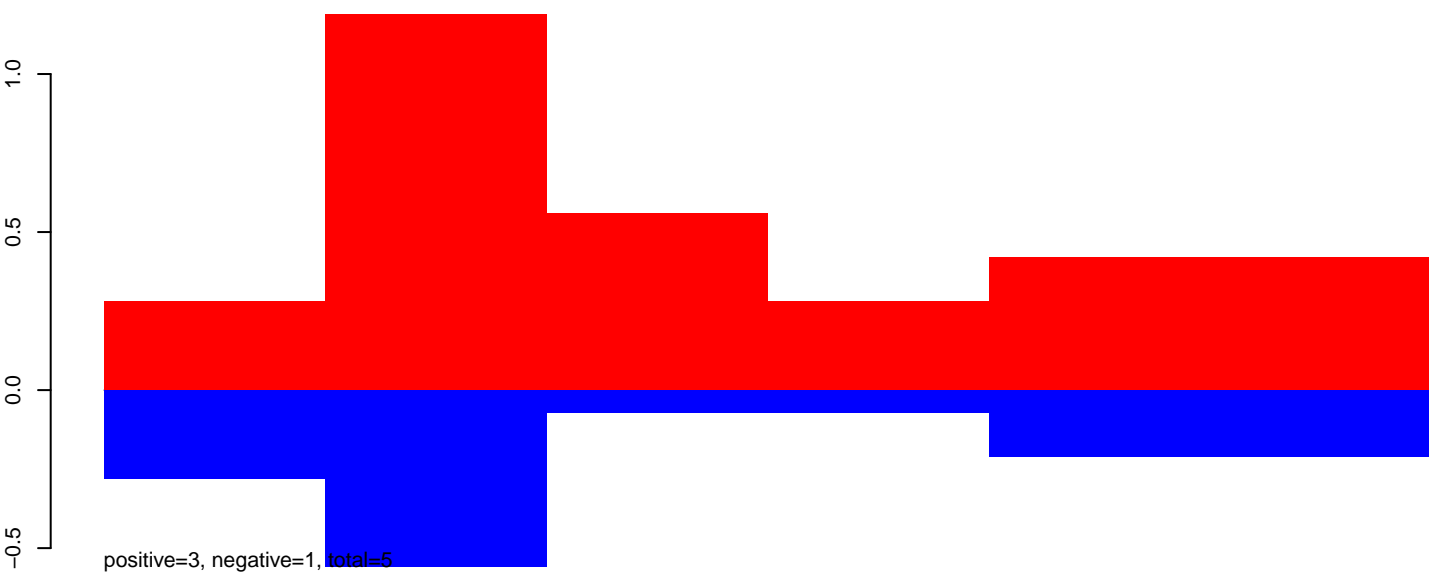


CuQuin_Mcarc_NL.rep

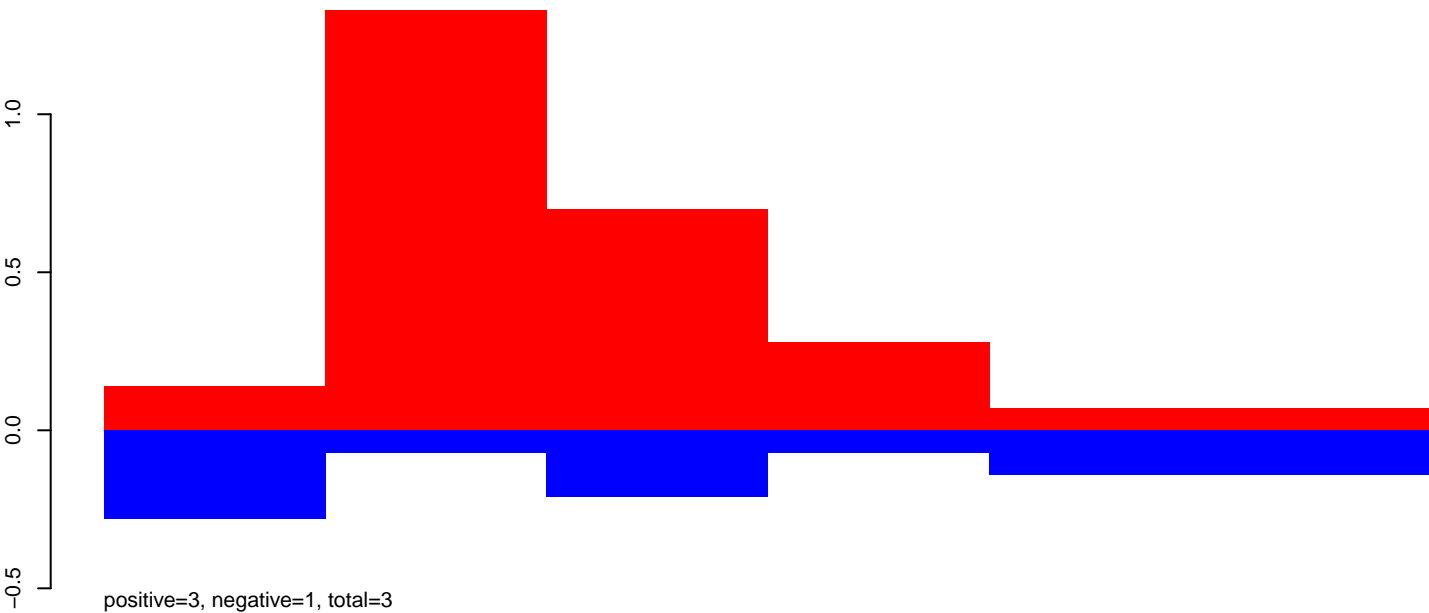


Window size=25, length=135, TE@rnd-3_family-233@Satellite@2645:1-135

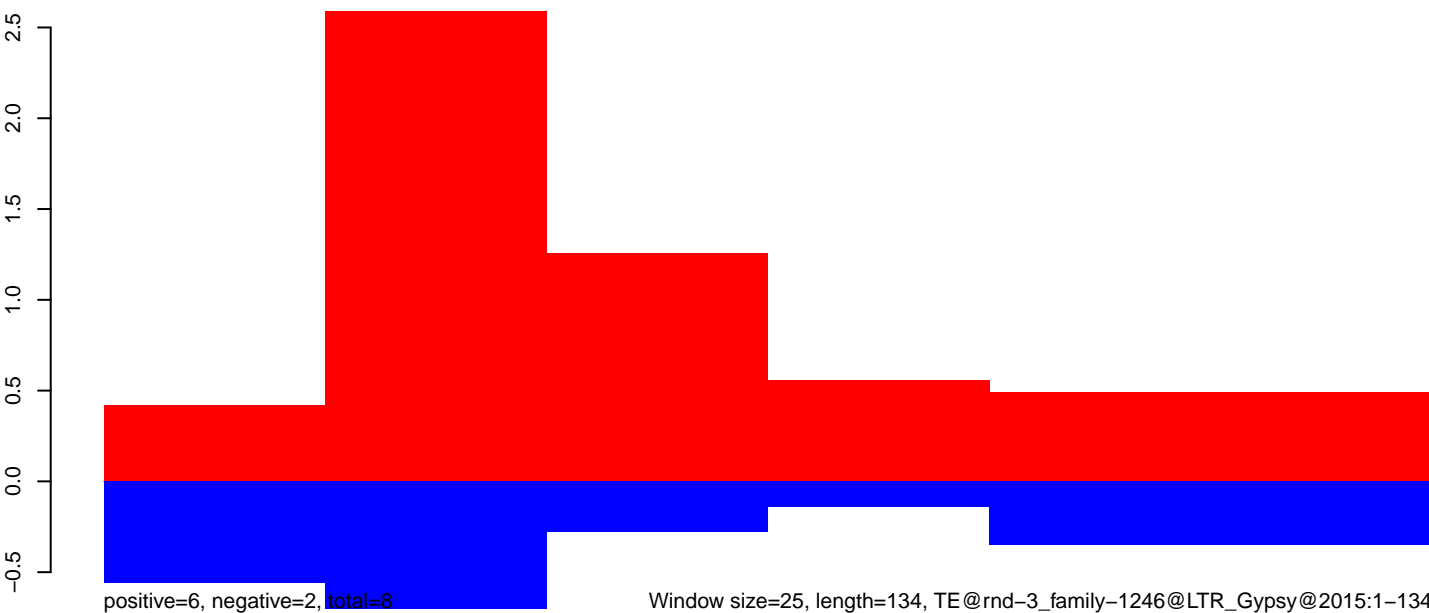
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



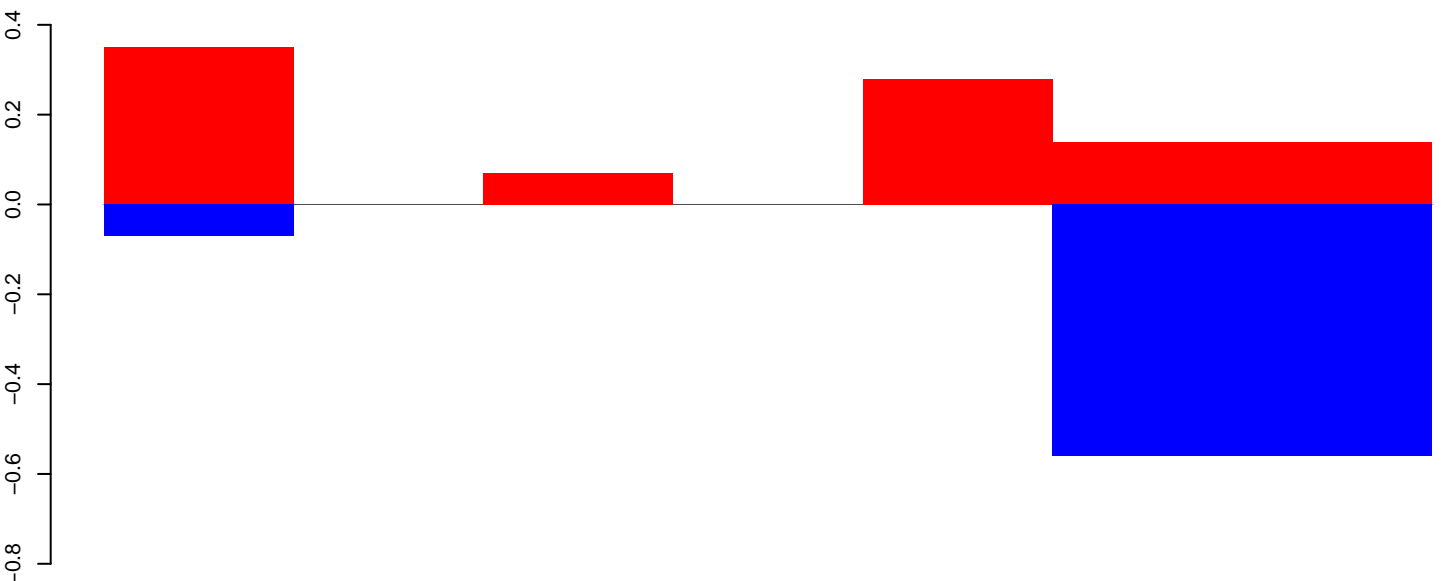
Window size=25, length=134, TE@rnd-3_family-1246@LTR_Gypsy@2015:1-134

50

100

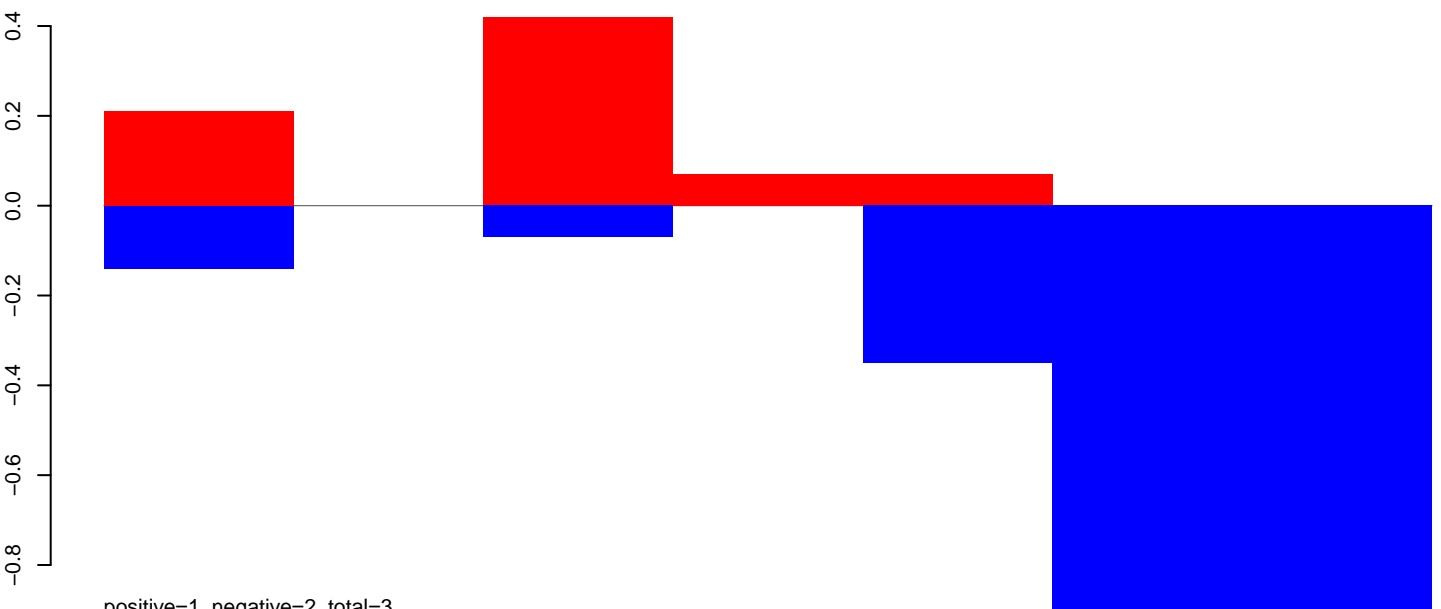
150

CuQuin_Mcarc_NL.18_23.rep



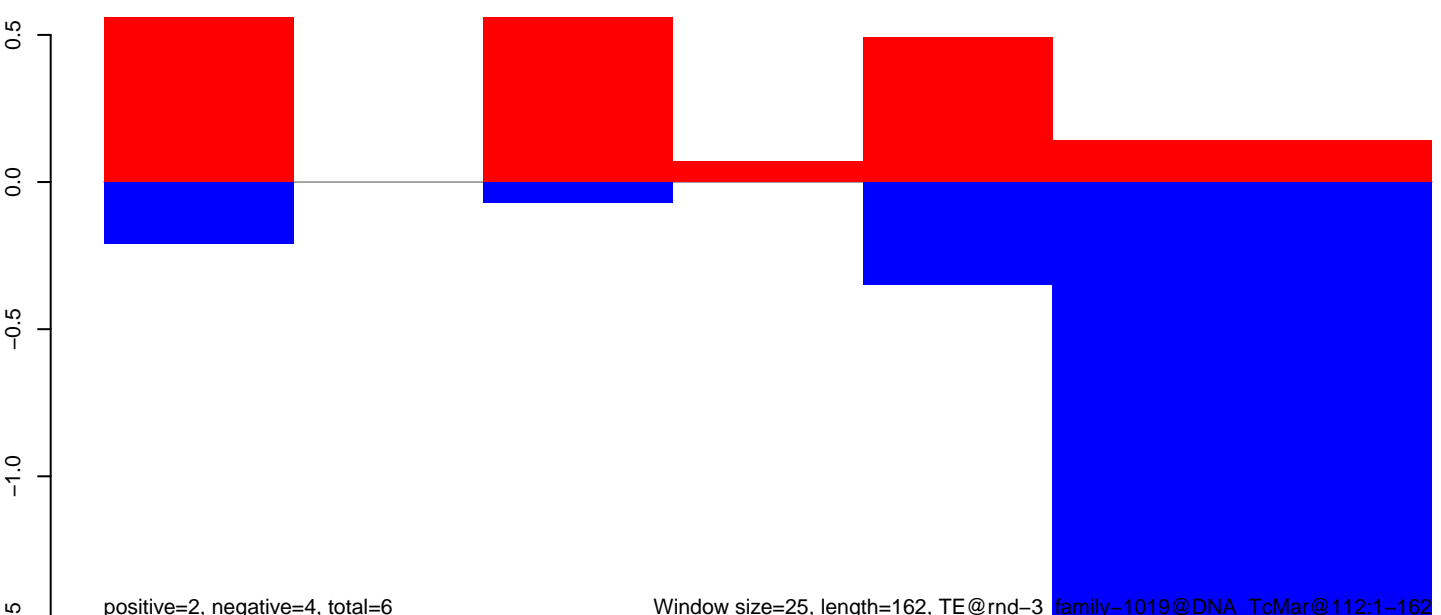
positive=1, negative=1, total=2

CuQuin_Mcarc_NL.24_35.rep



positive=1, negative=2, total=3

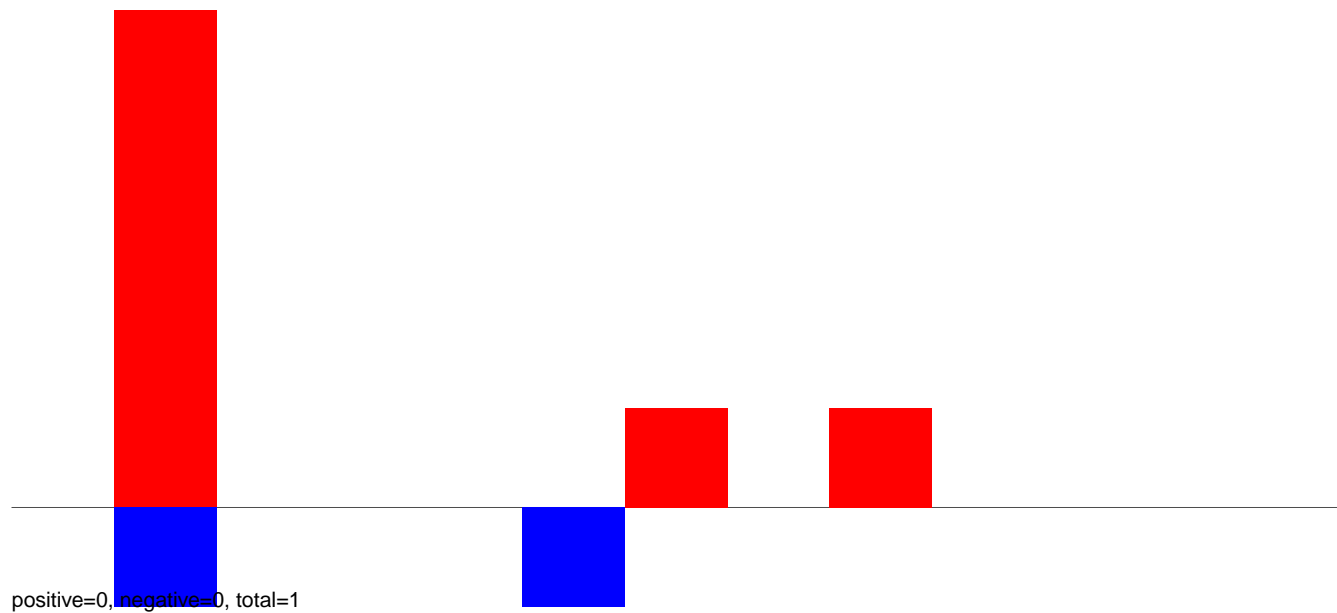
CuQuin_Mcarc_NL.rep



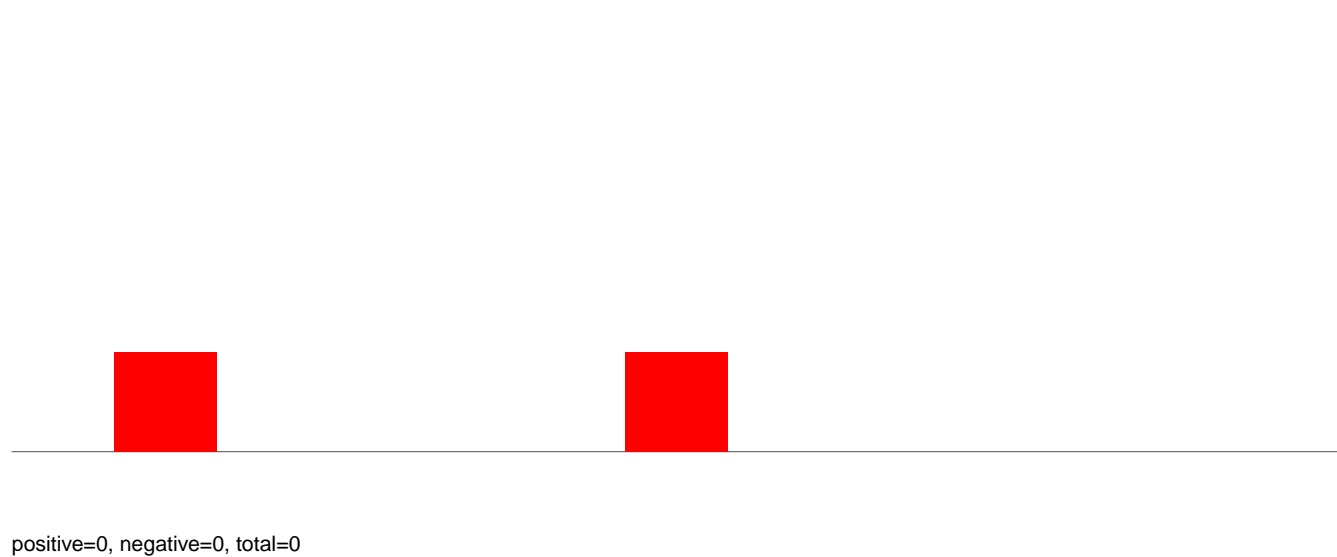
positive=2, negative=4, total=6

Window size=25, length=162, TE@rnd-3_family-1019@DNA_TcMar@112:1-162

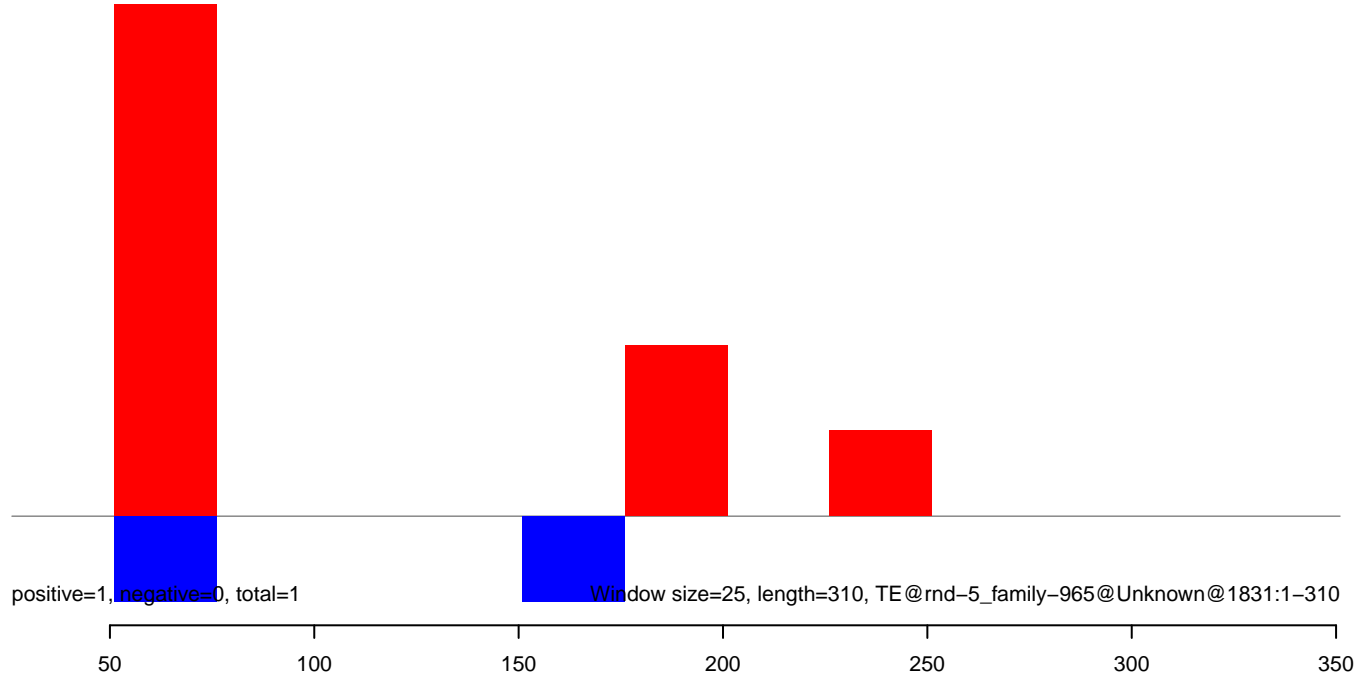
CuQuin_Mcarc_NL.18_23.rep



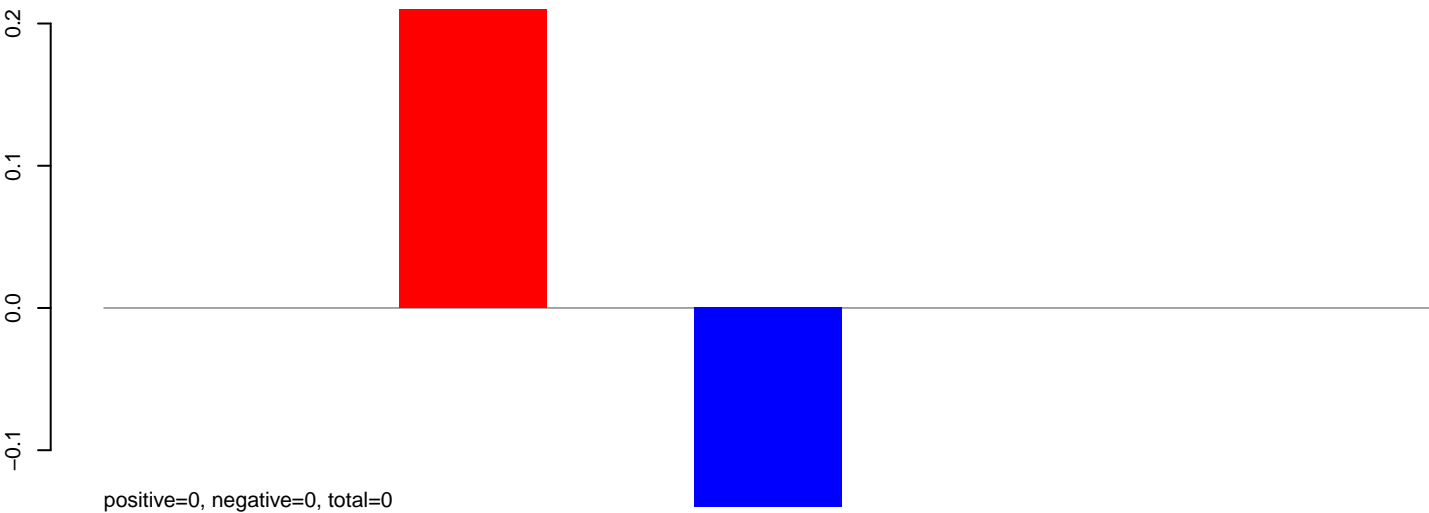
CuQuin_Mcarc_NL.24_35.rep



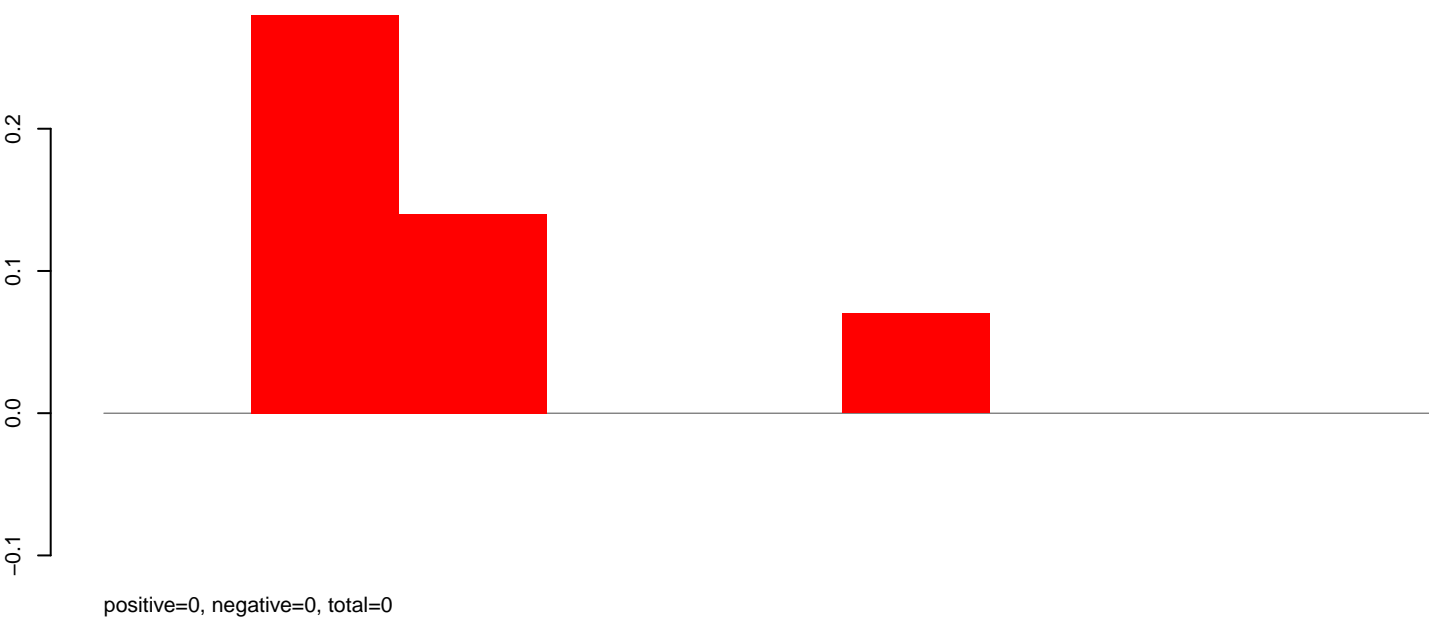
CuQuin_Mcarc_NL.rep



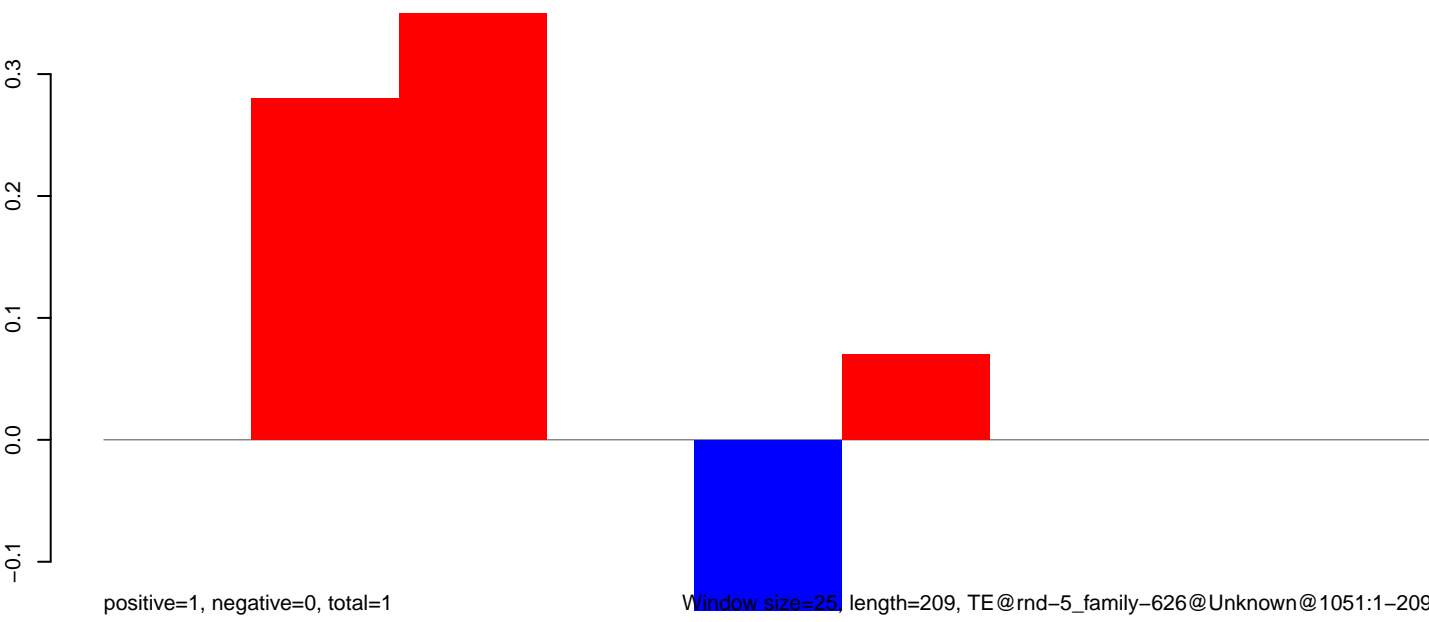
CuQuin_Mcarc_NL.18_23.rep



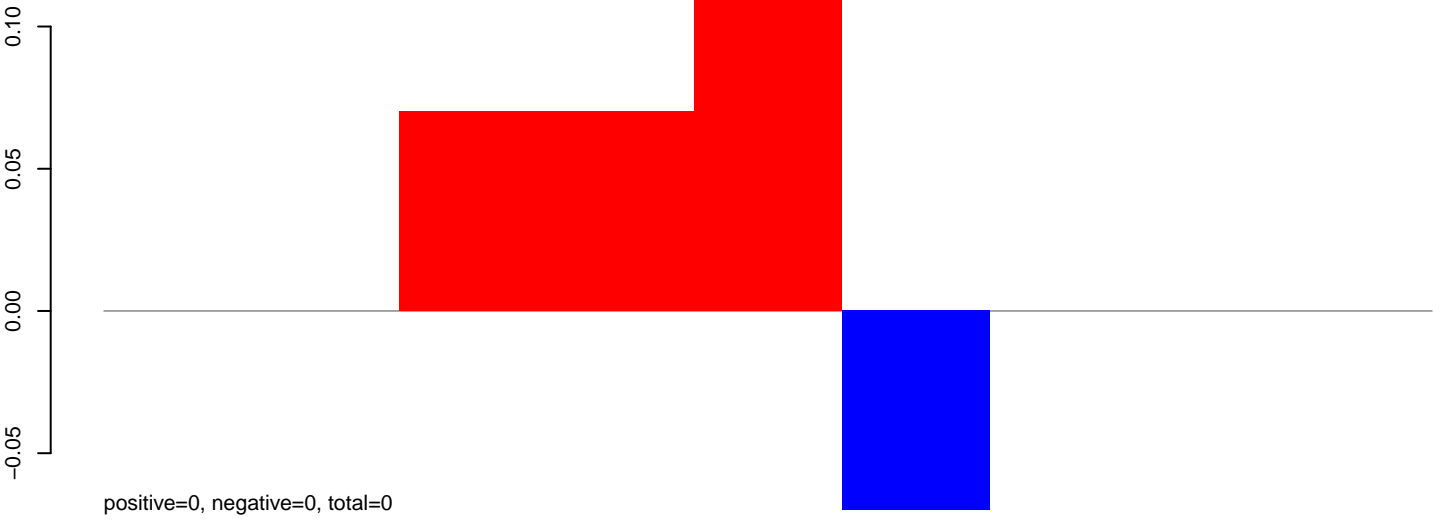
CuQuin_Mcarc_NL.24_35.rep



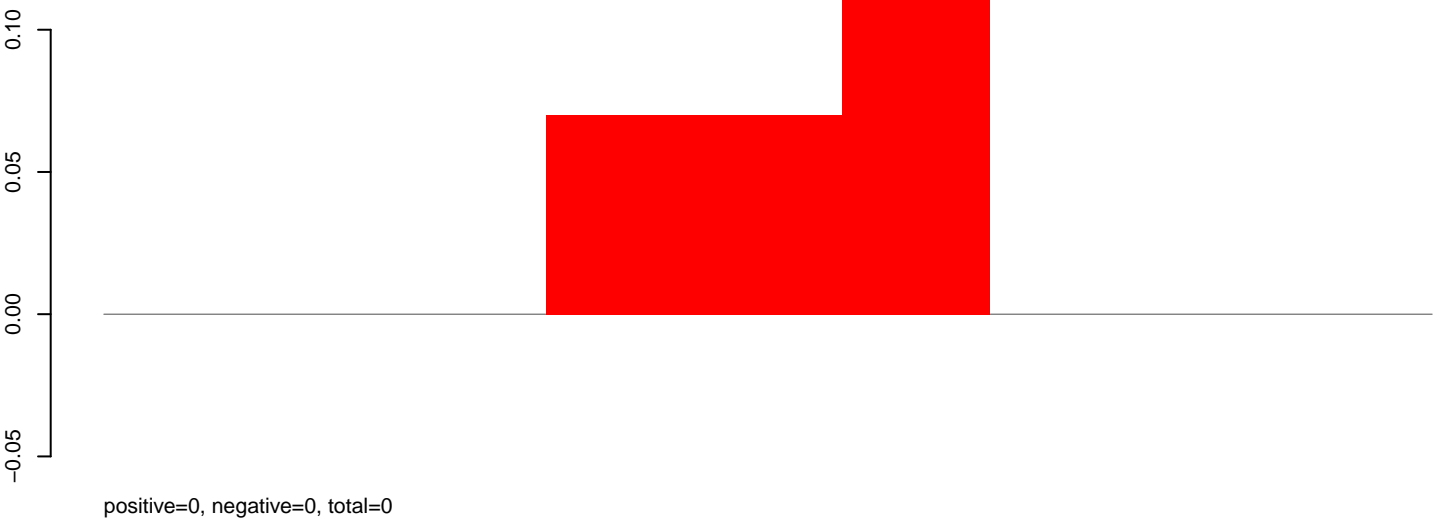
CuQuin_Mcarc_NL.rep



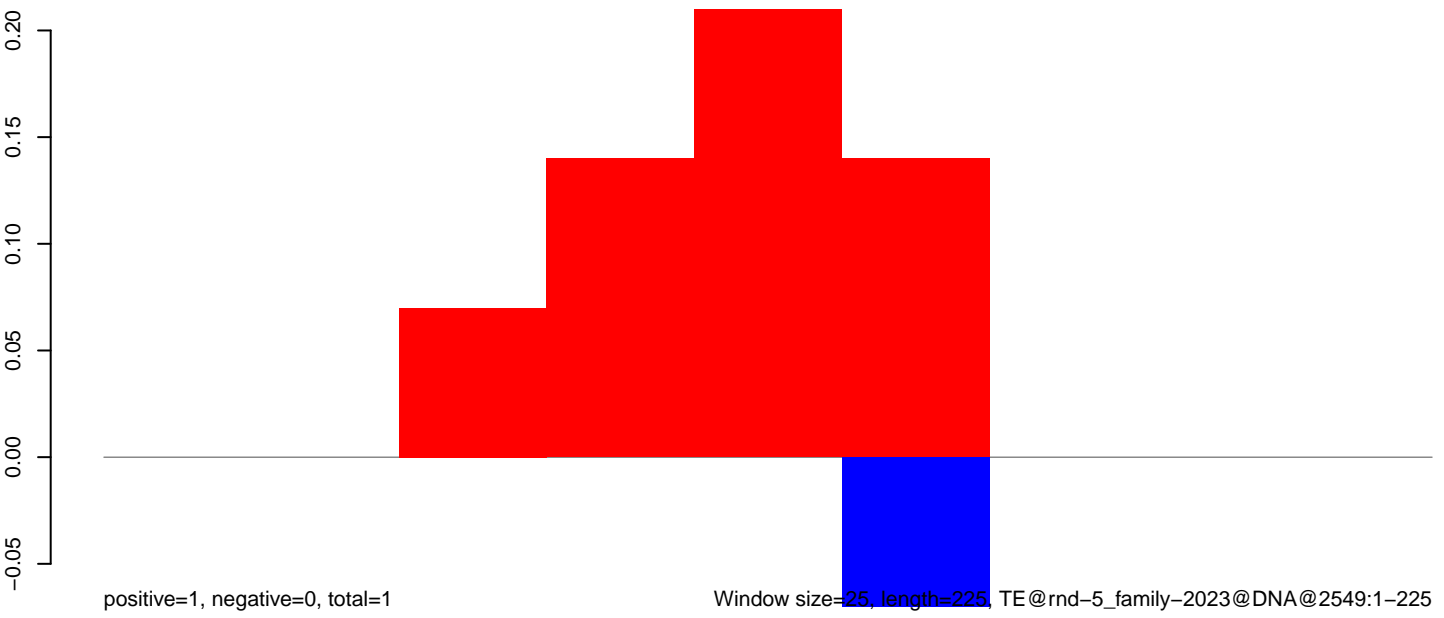
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



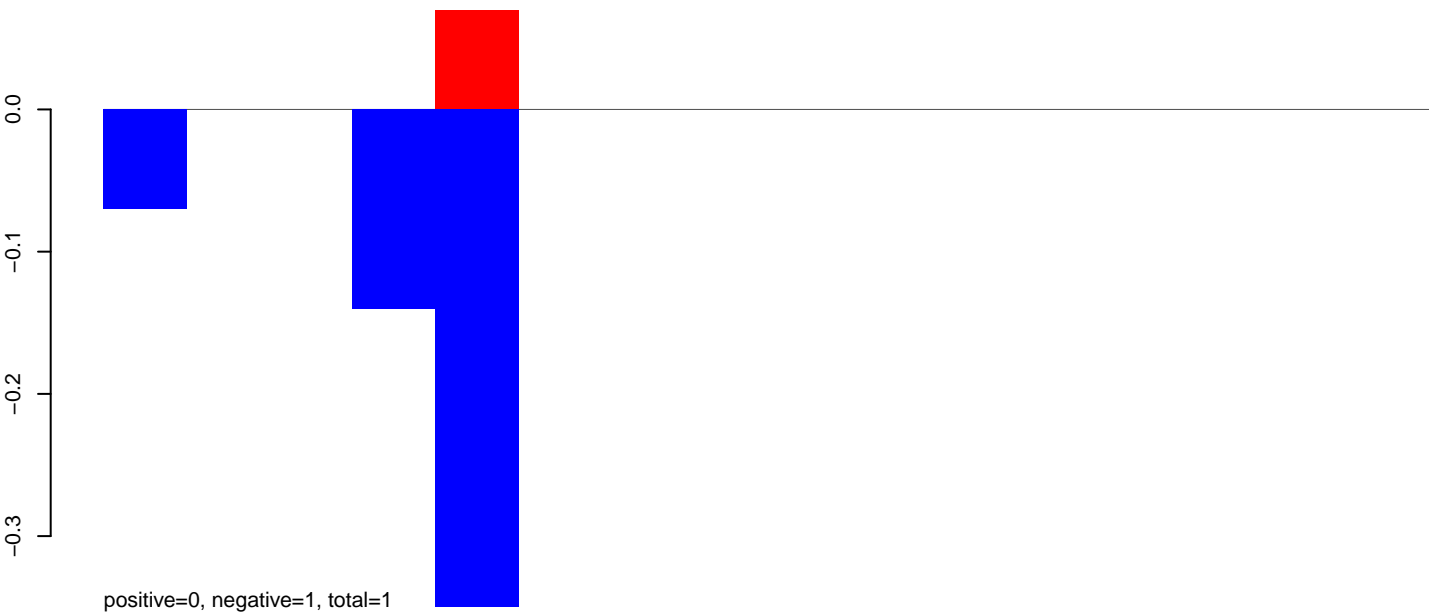
CuQuin_Mcarc_NL.rep



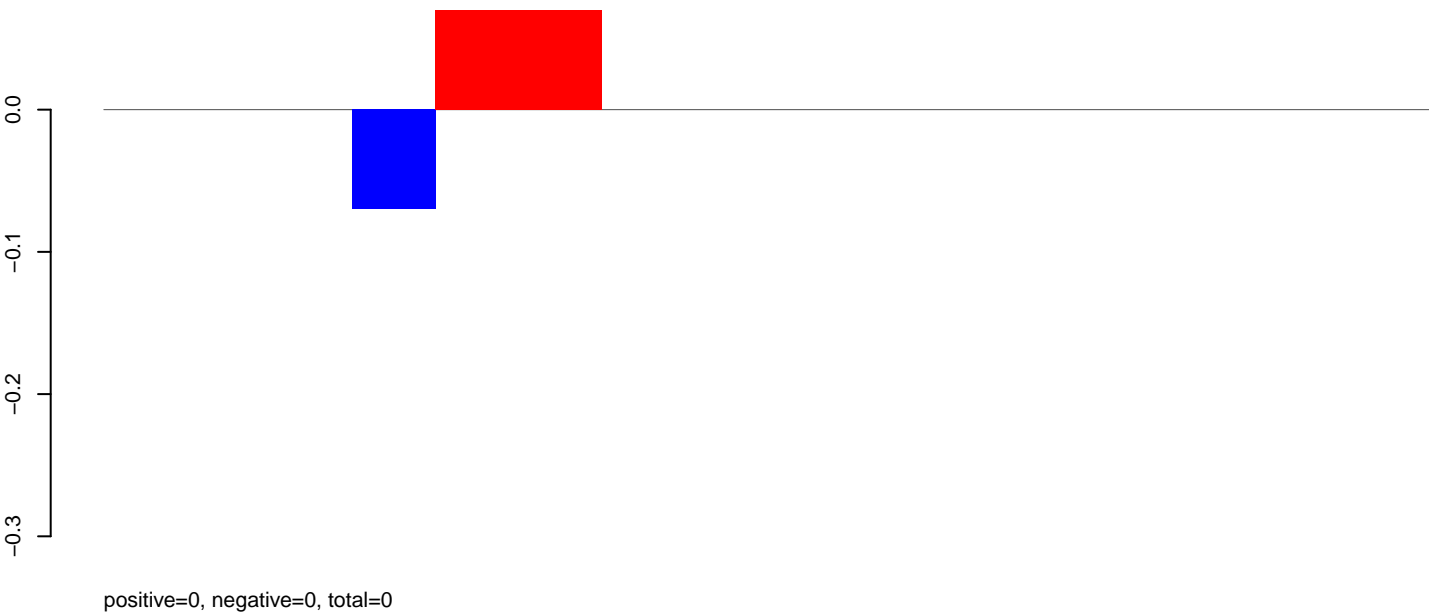
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50 100 150 200 250

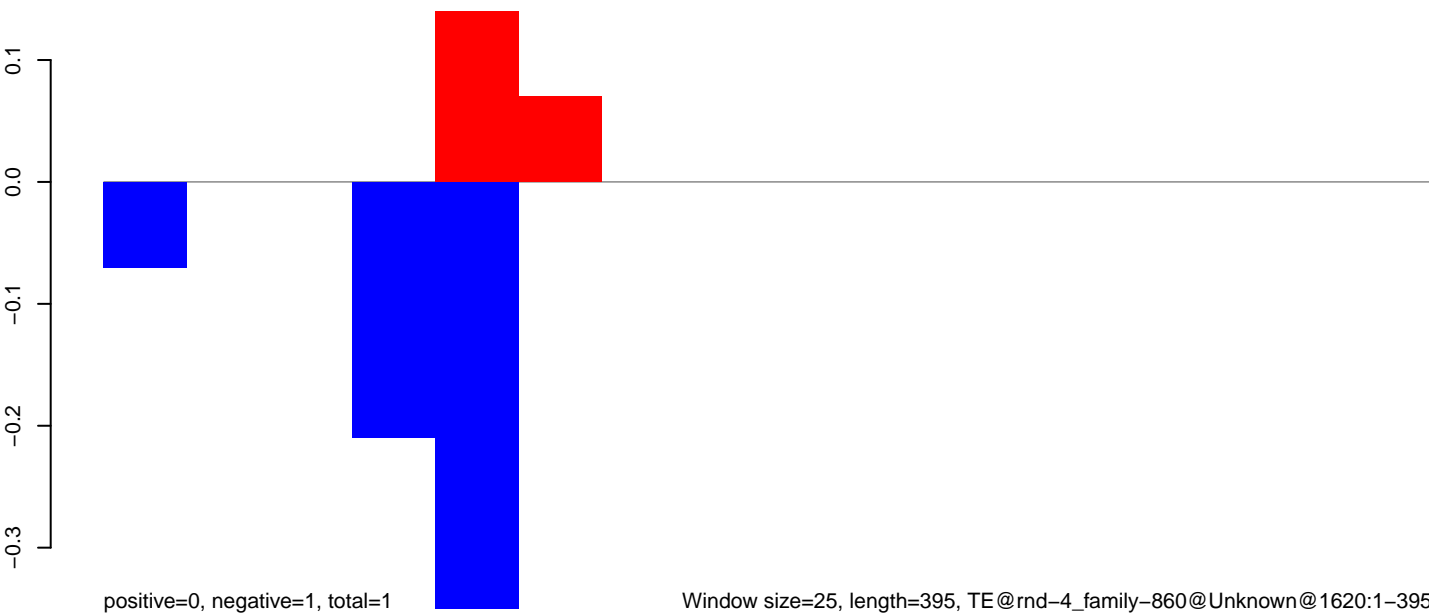
CuQuin_Mcarc_NL.18_23.rep



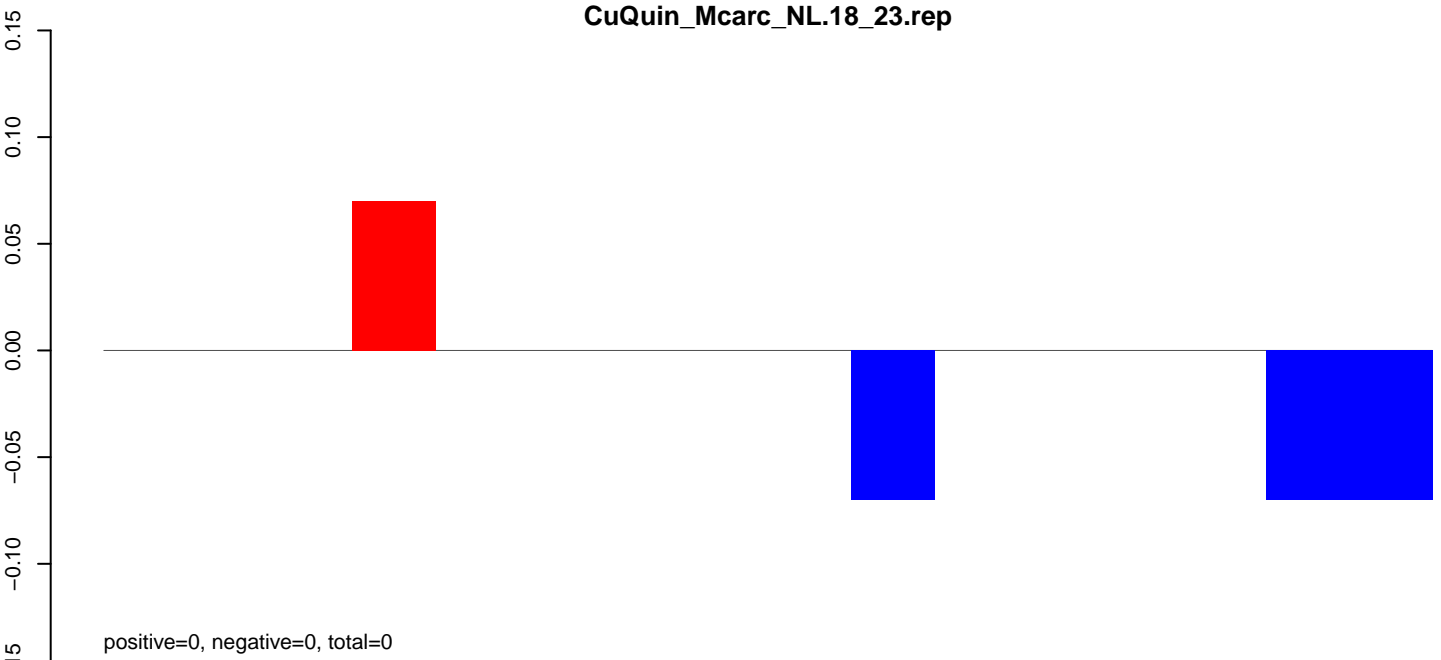
CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

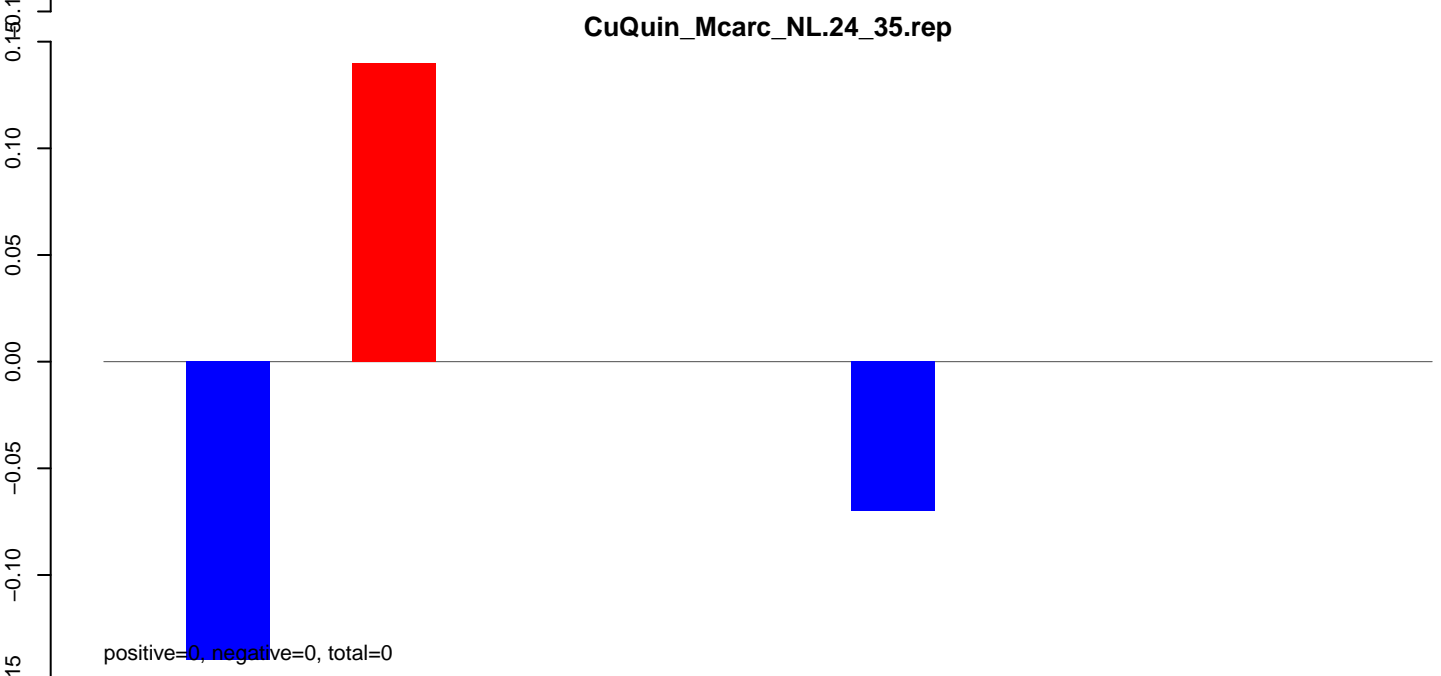


CuQuin_Mcarc_NL.18_23.rep



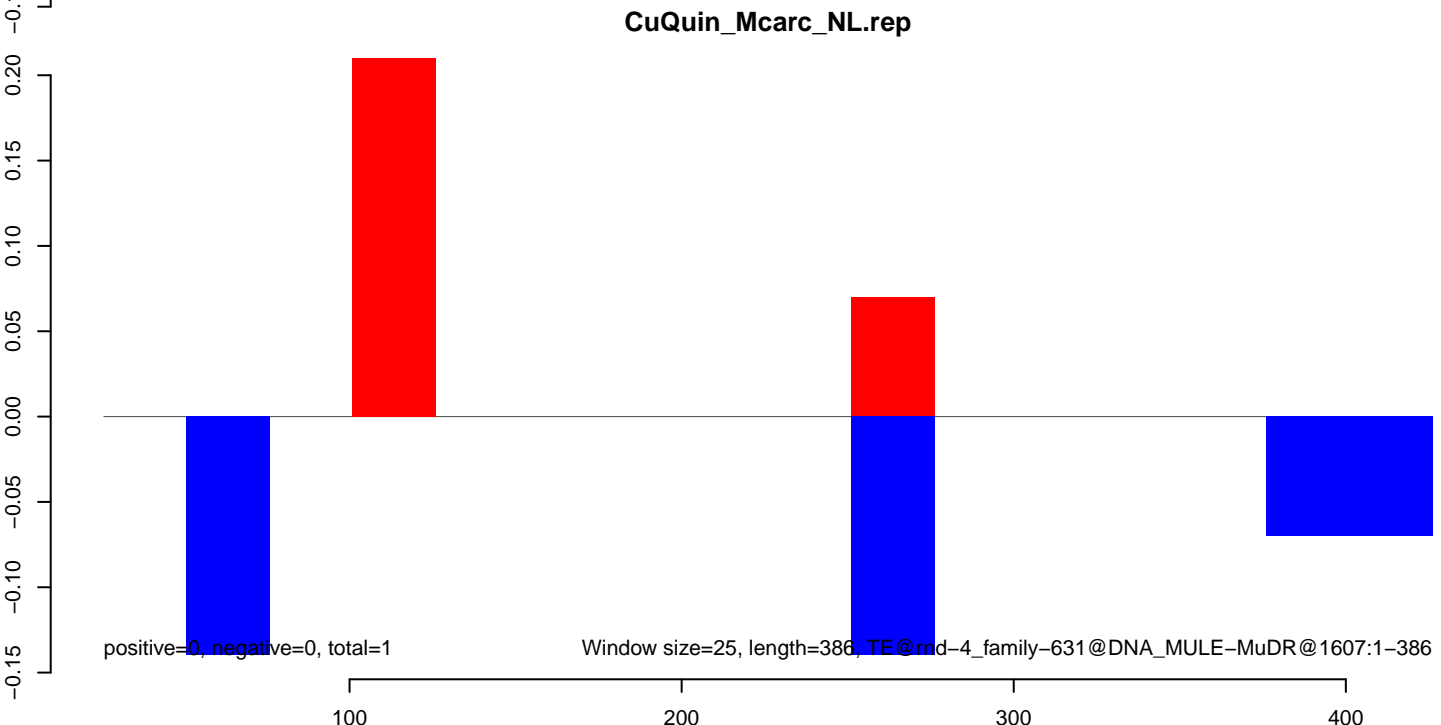
positive=0, negative=0, total=0

CuQuin_Mcarc_NL.24_35.rep



positive=0, negative=0, total=0

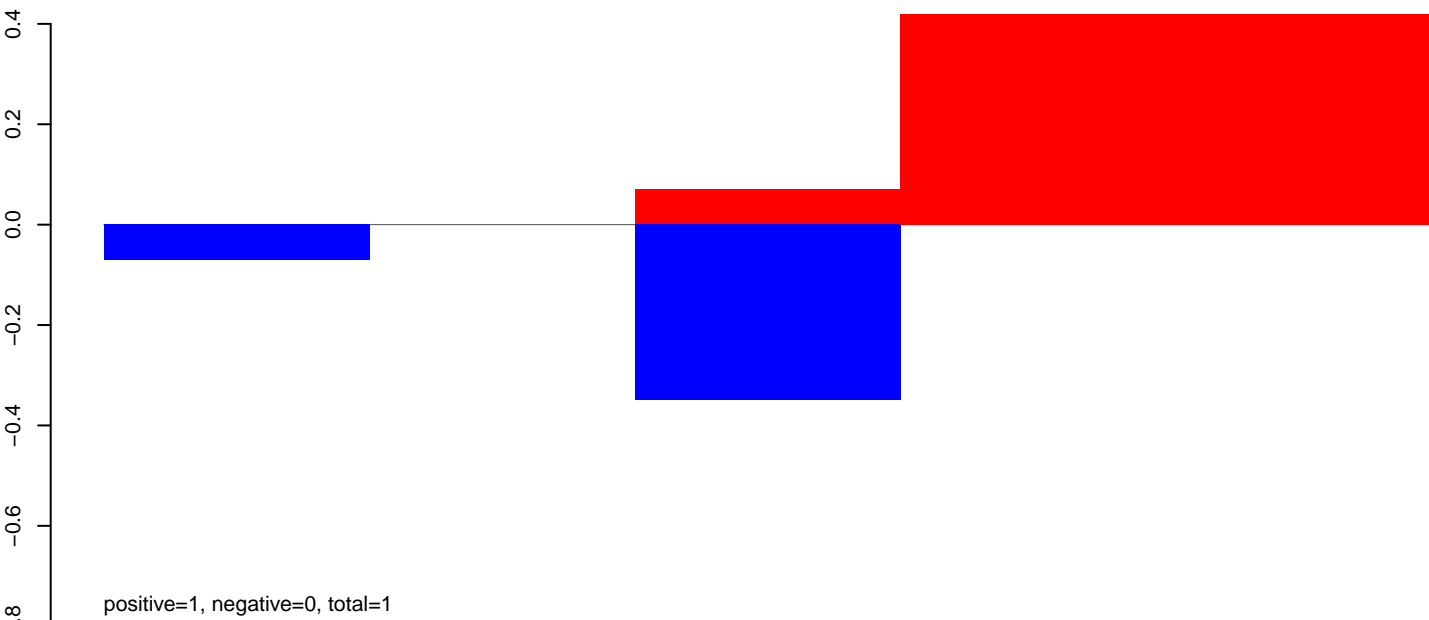
CuQuin_Mcarc_NL.rep



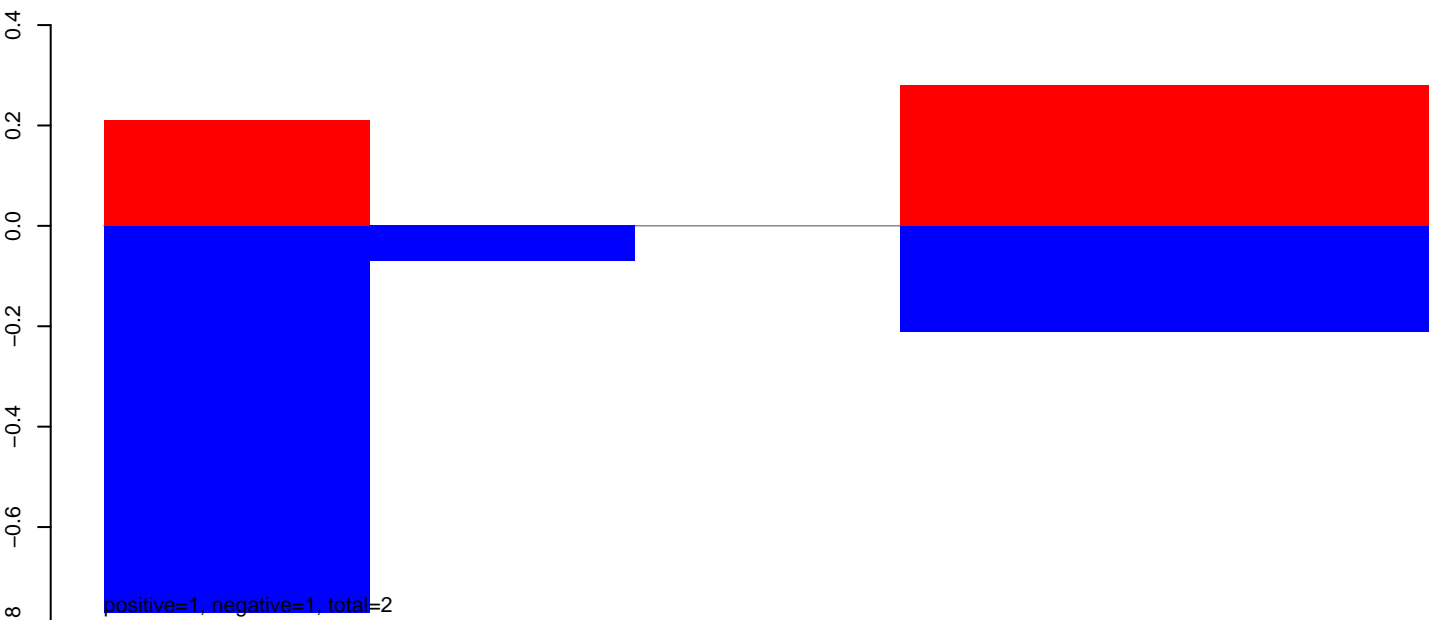
positive=0, negative=0, total=1

Window size=25, length=386, TE@rrnd-4_family-631@DNA_MULE-MuDR@1607:1-386

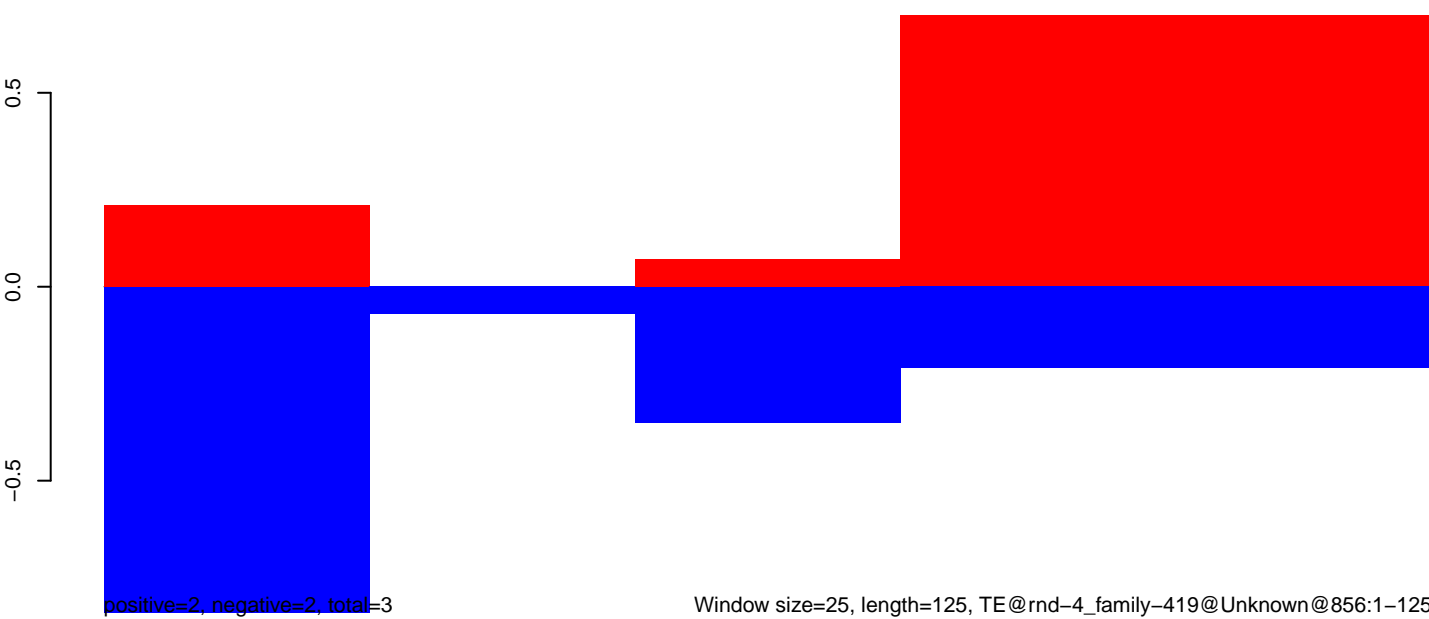
CuQuin_Mcarc_NL.18_23.rep



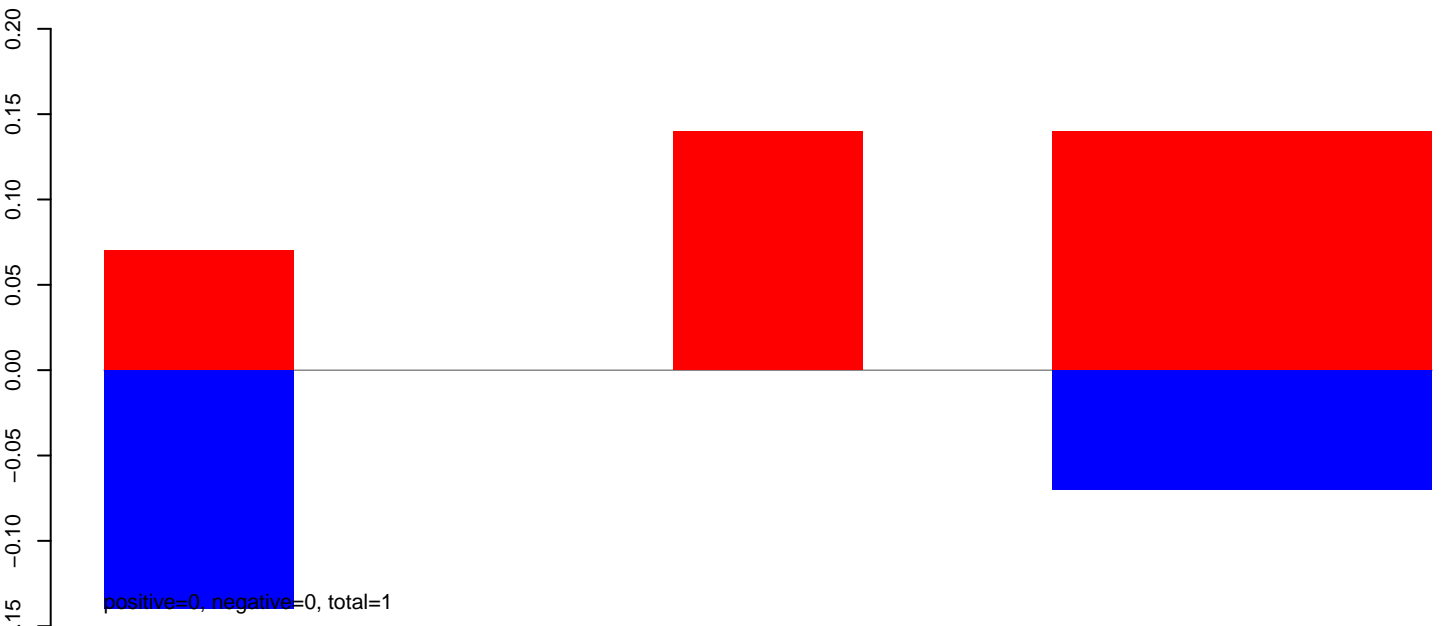
CuQuin_Mcarc_NL.24_35.rep



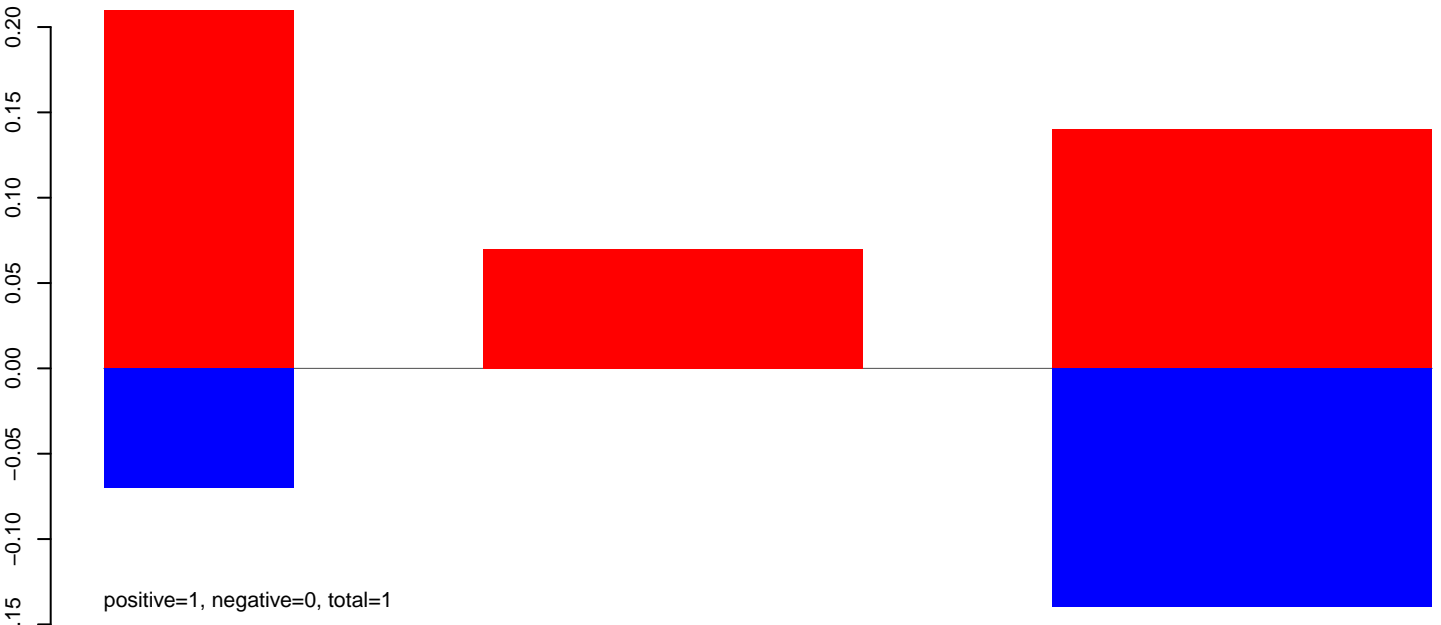
CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep



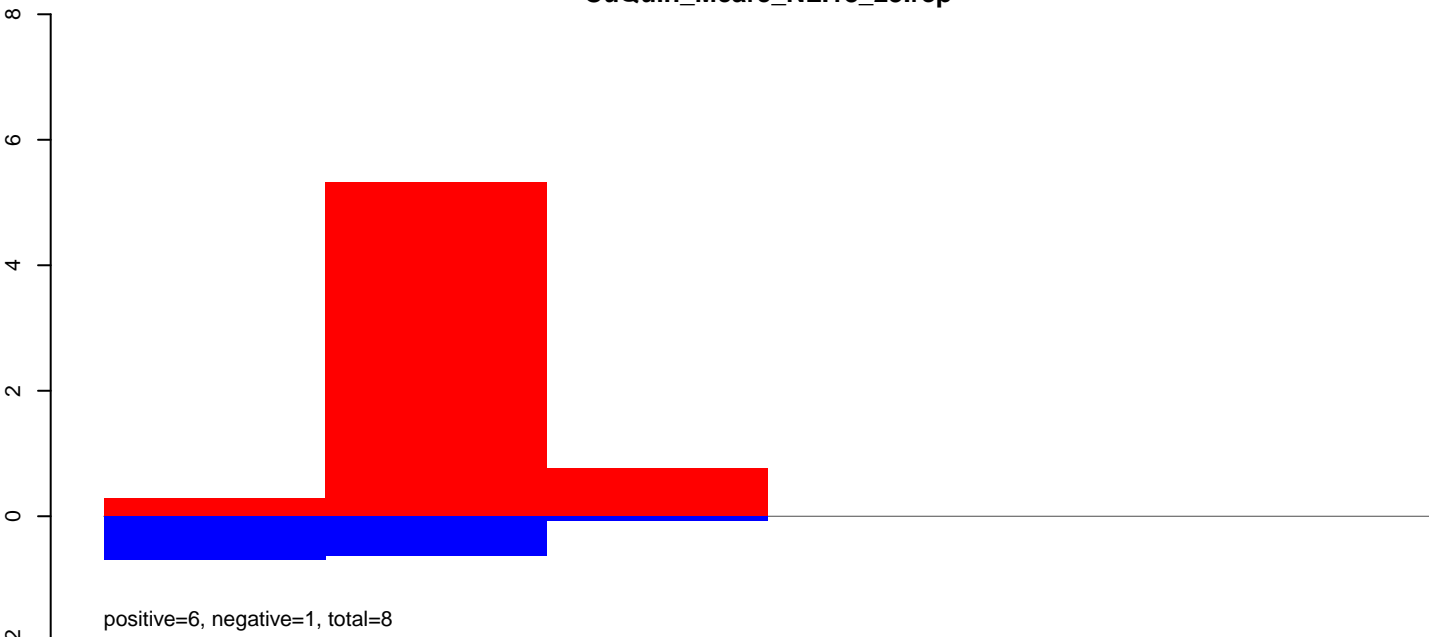
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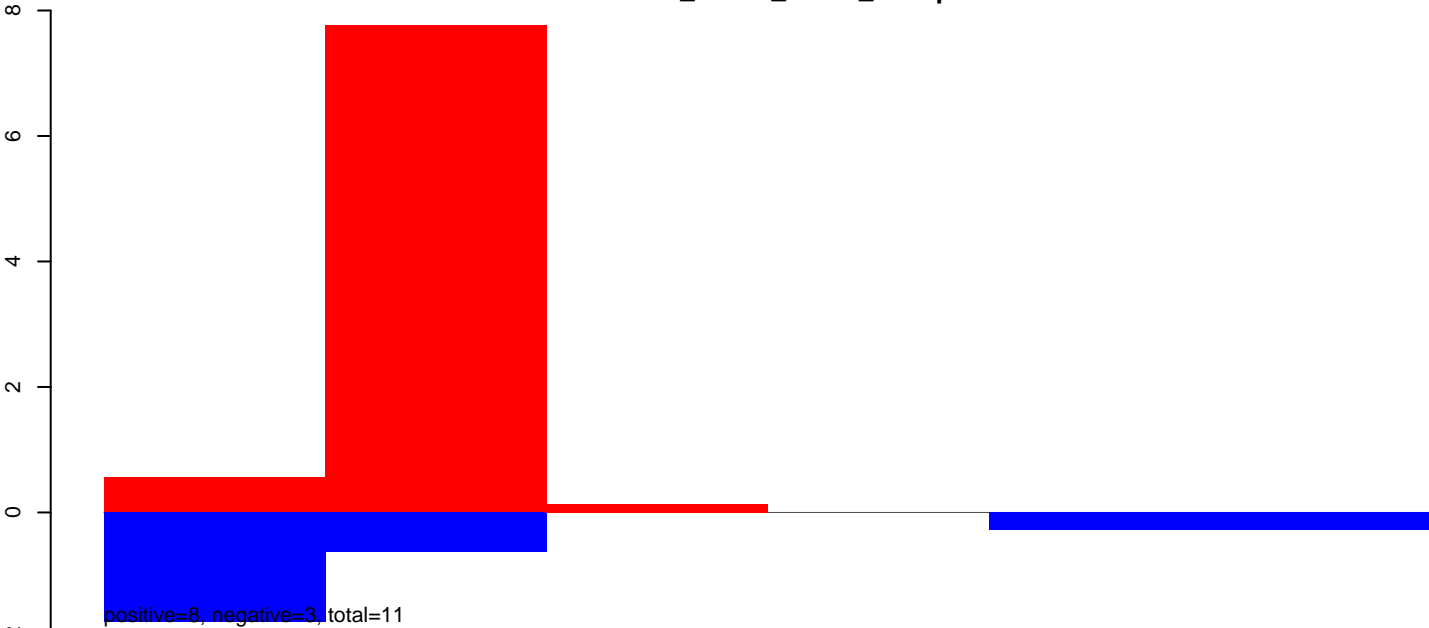
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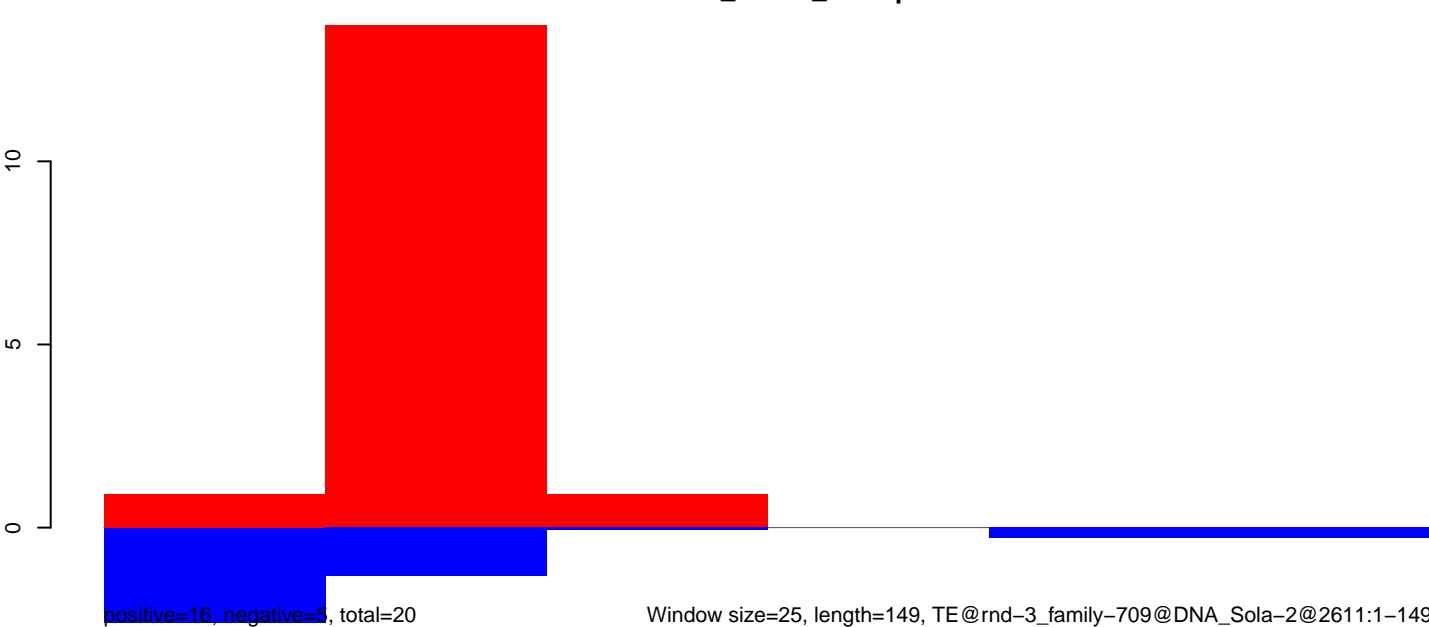
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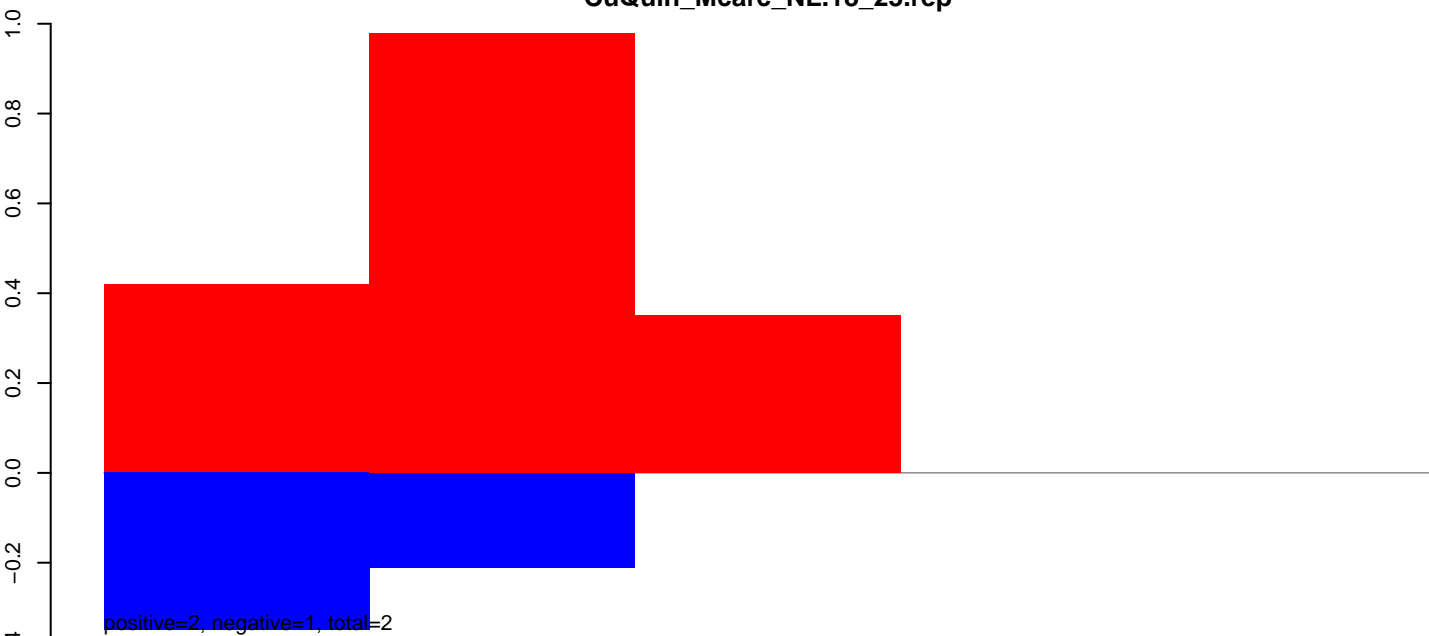
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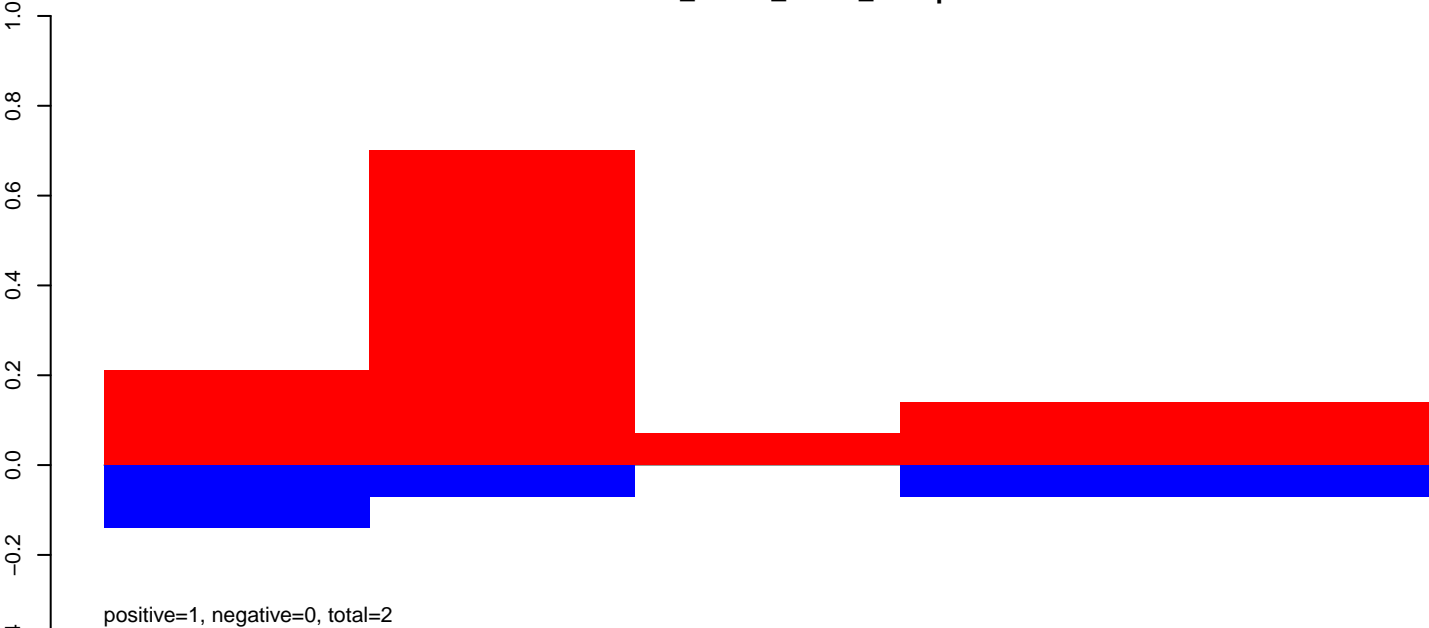
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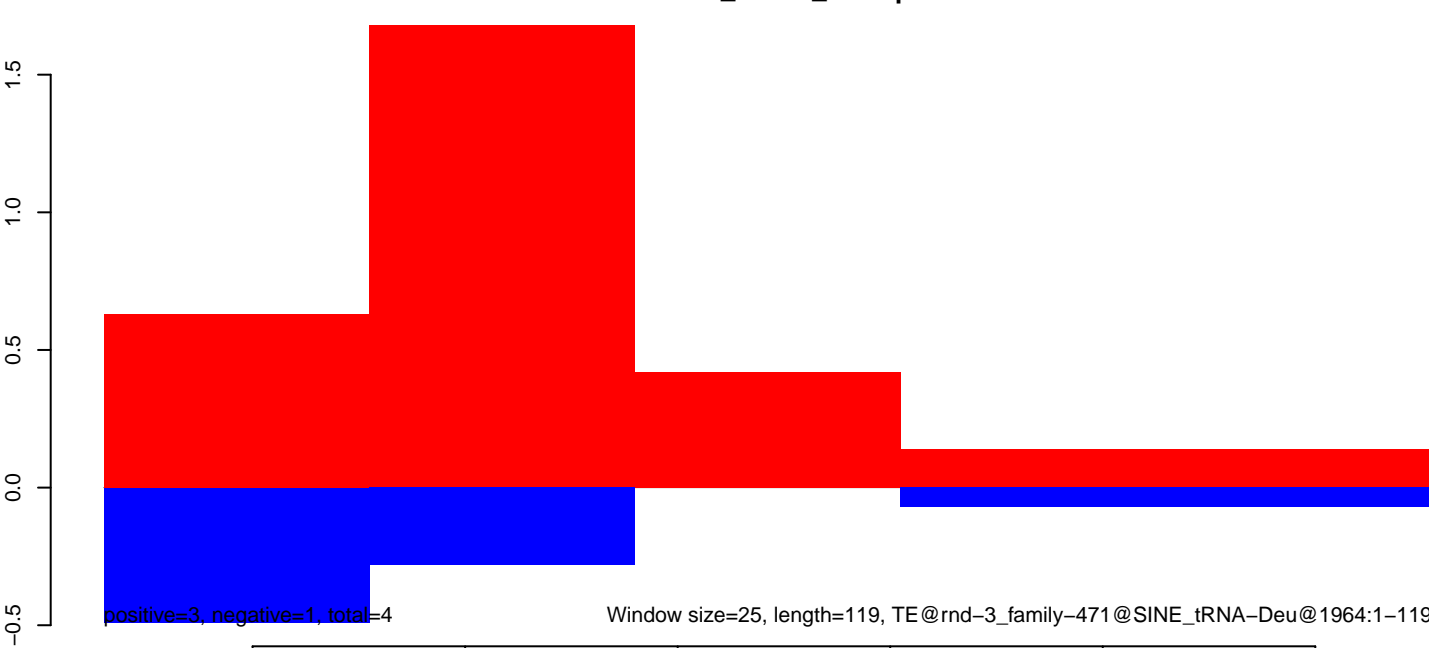
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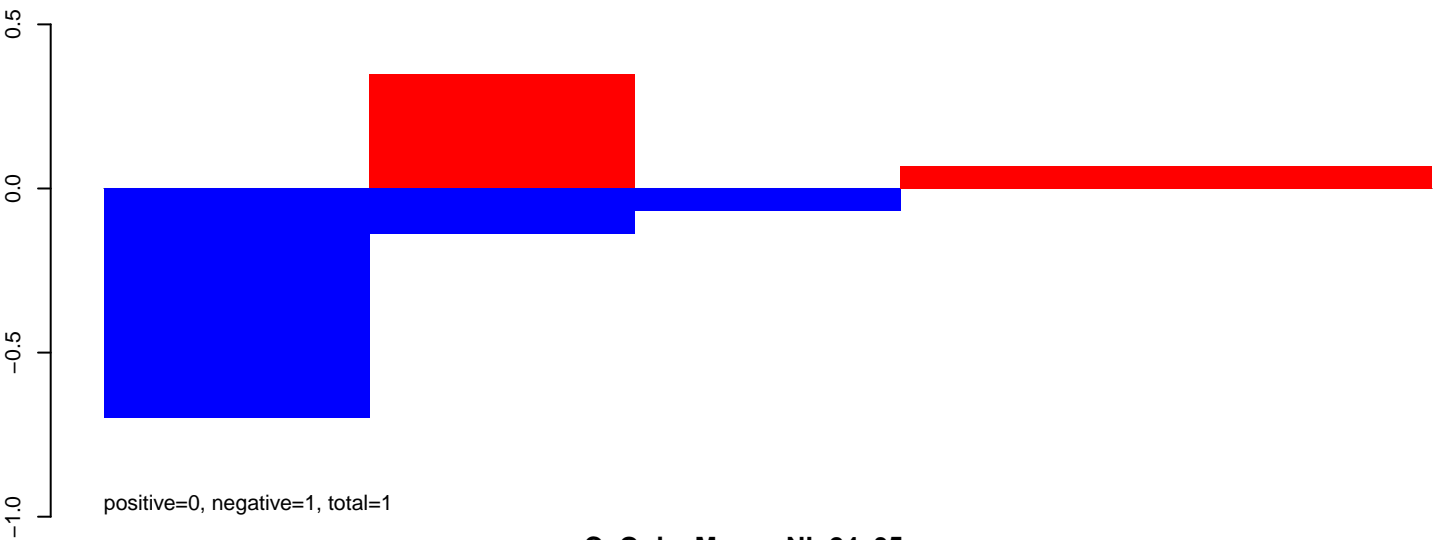
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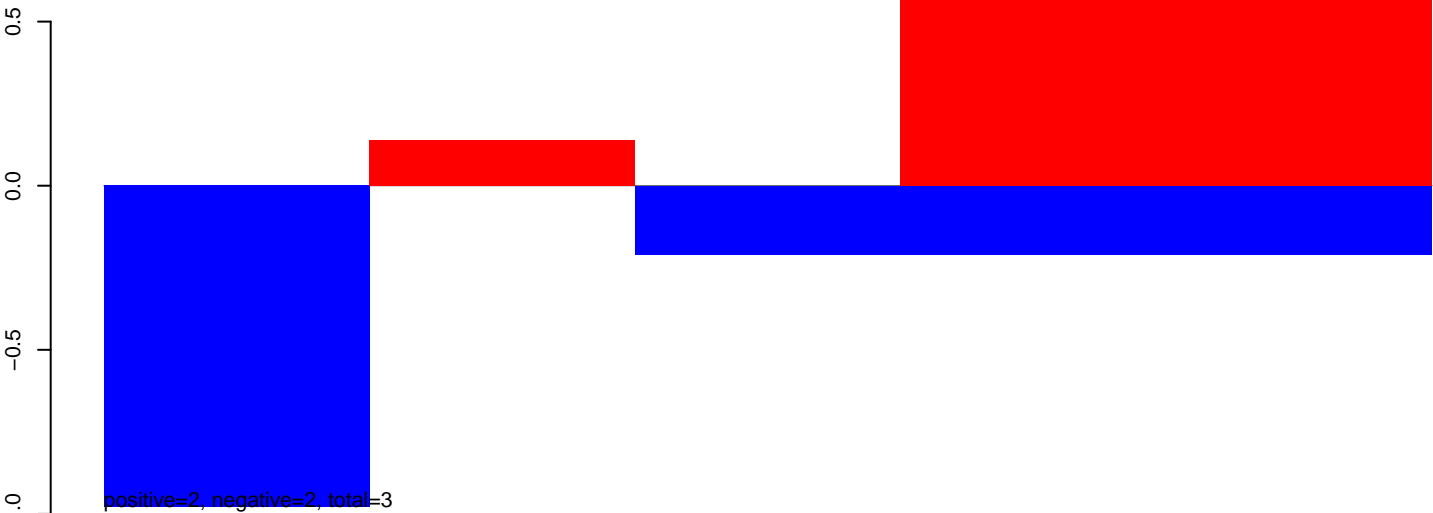
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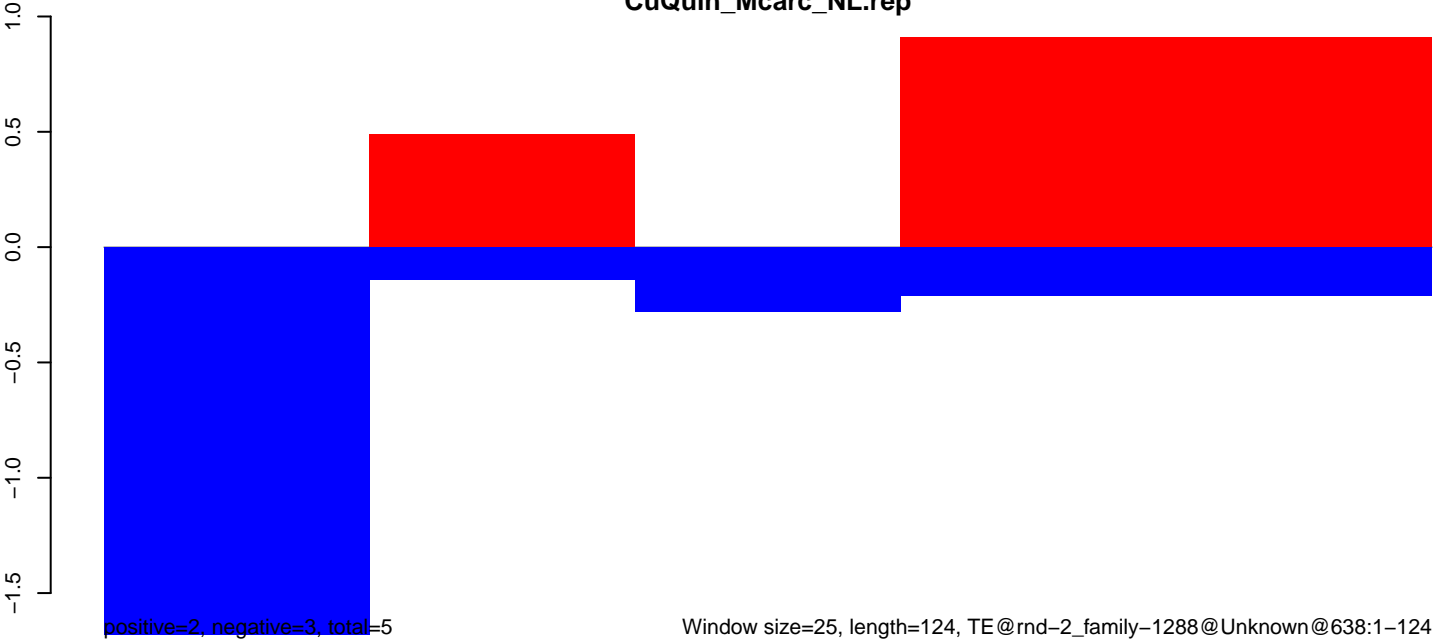
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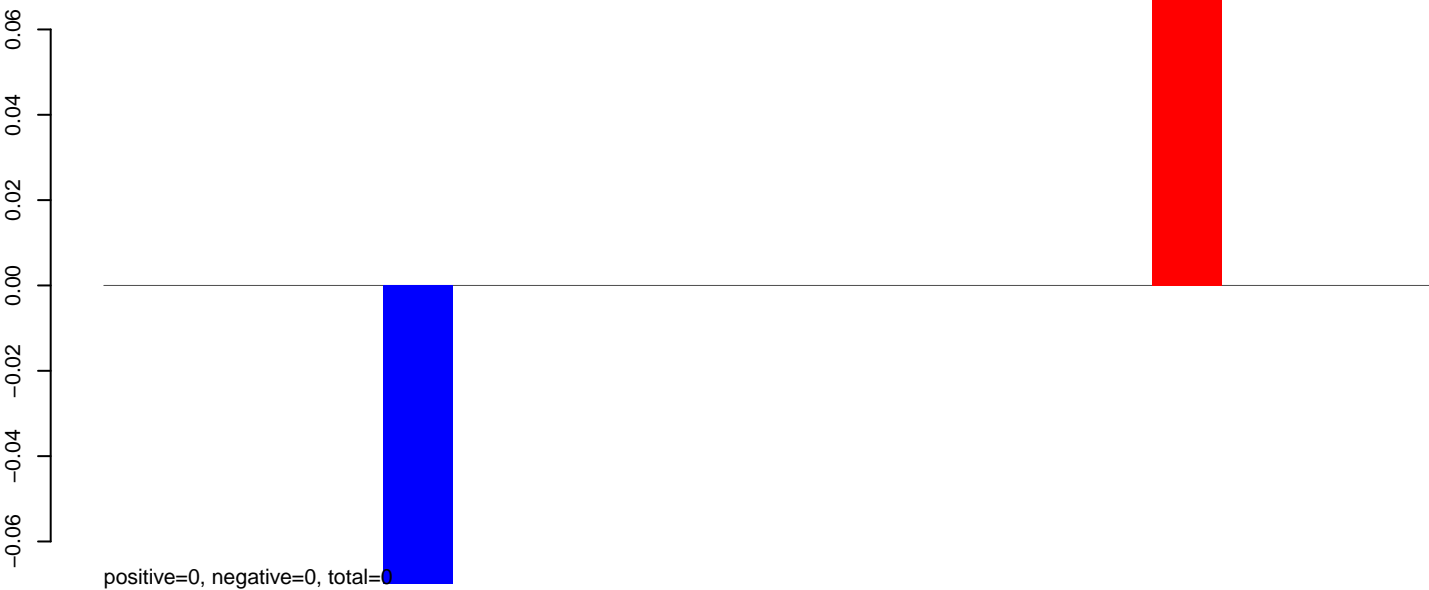
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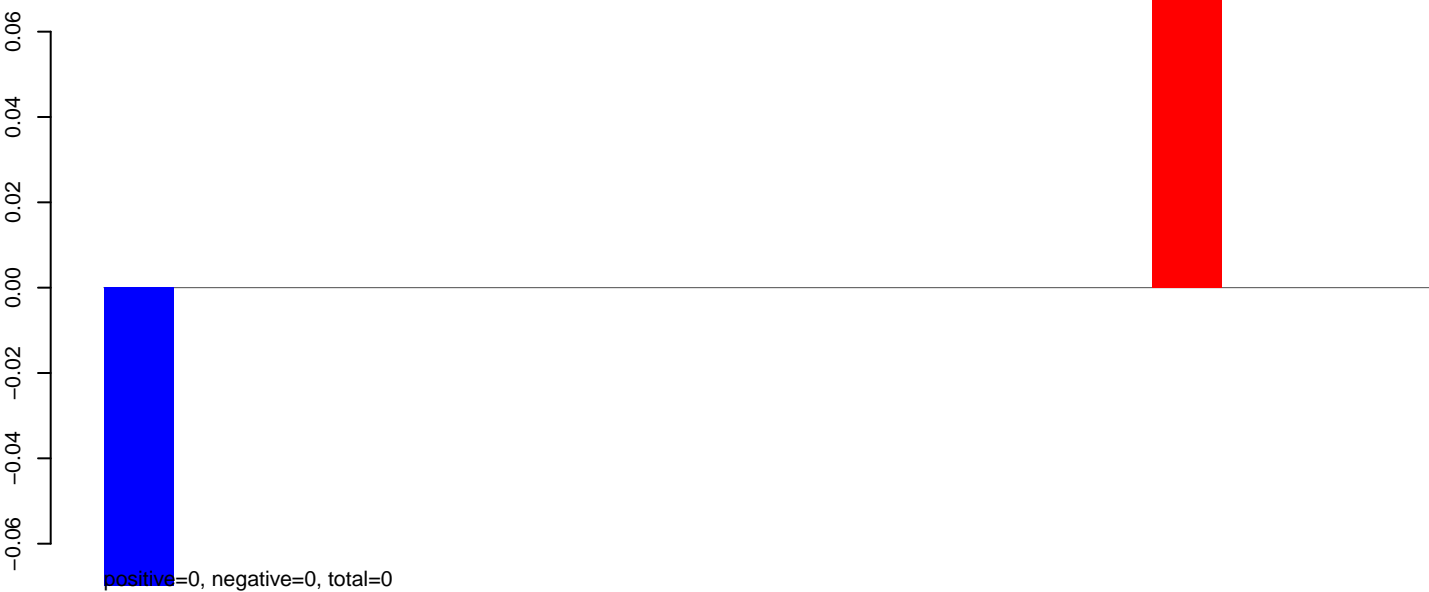
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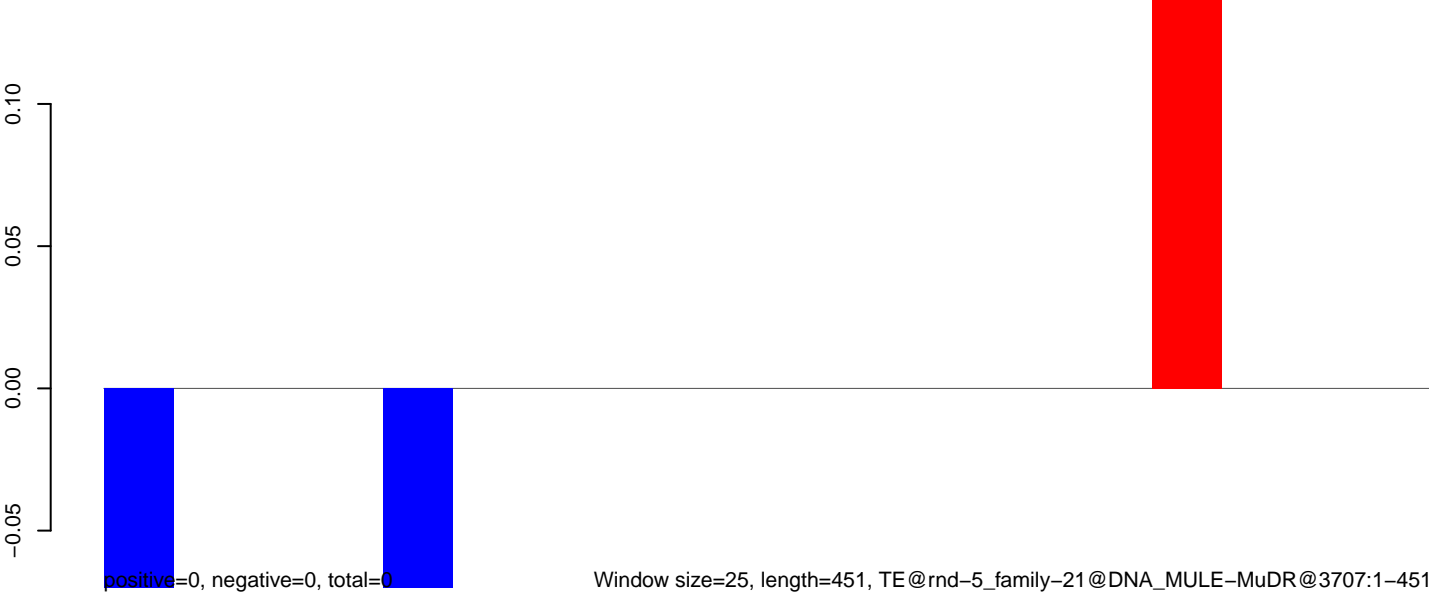
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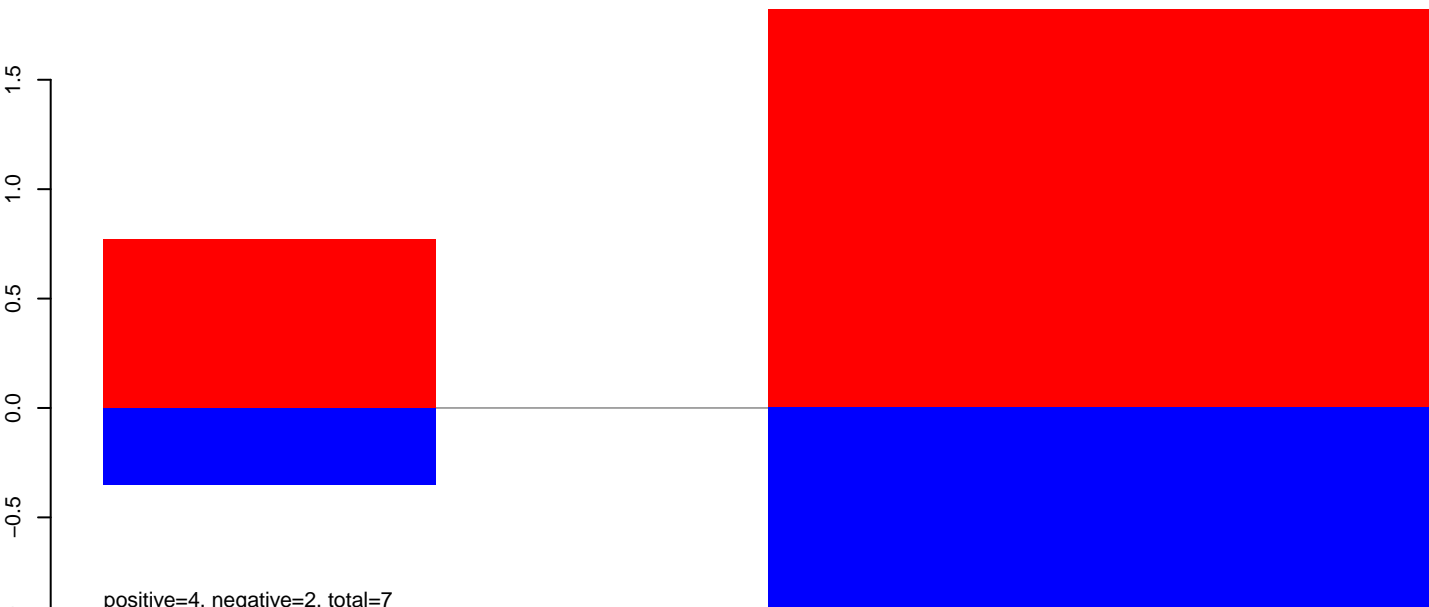
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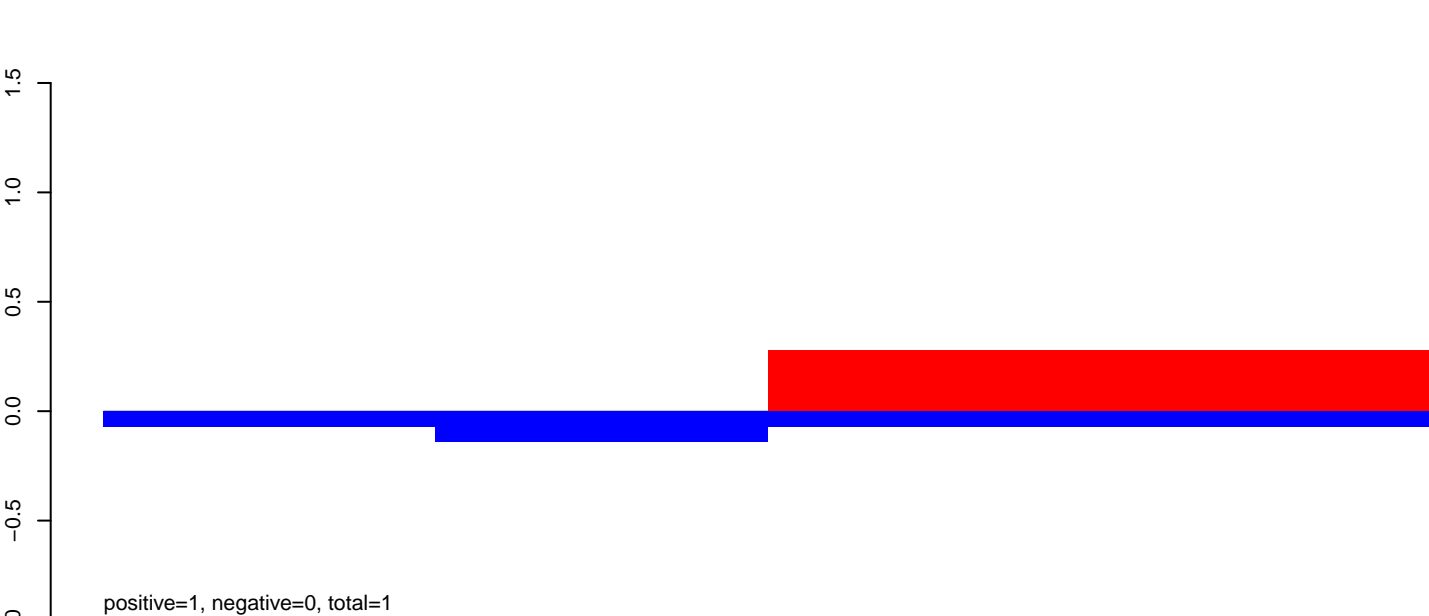
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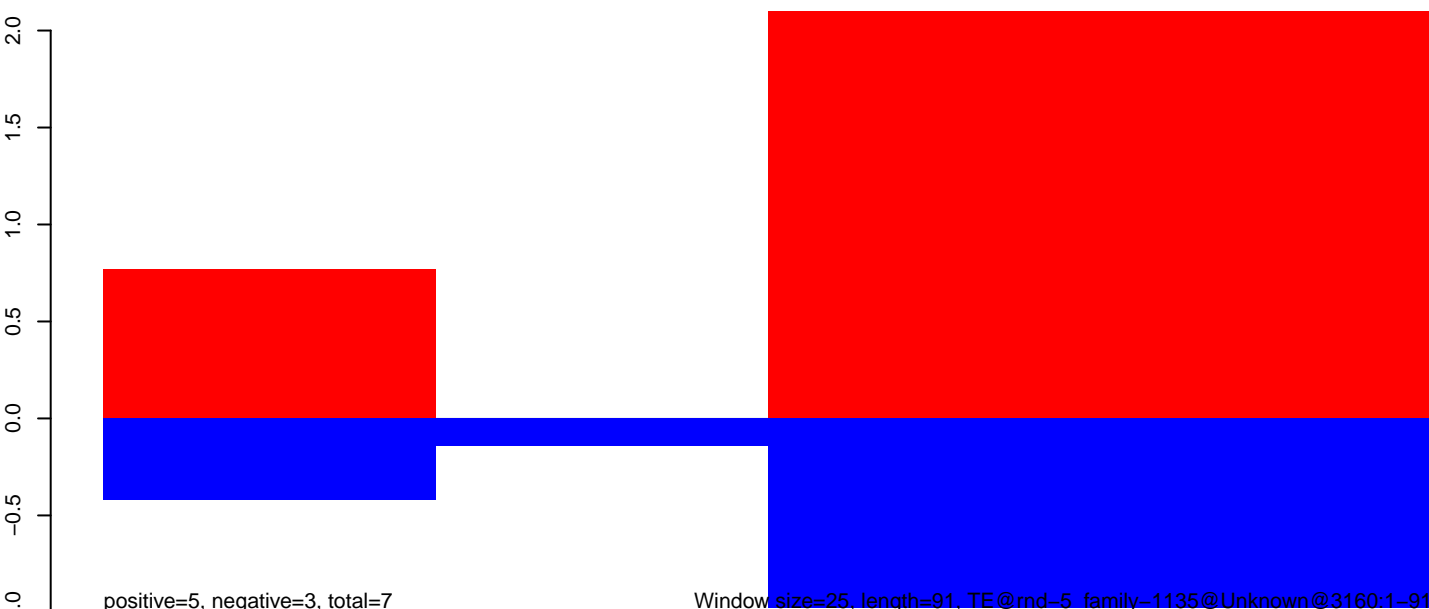
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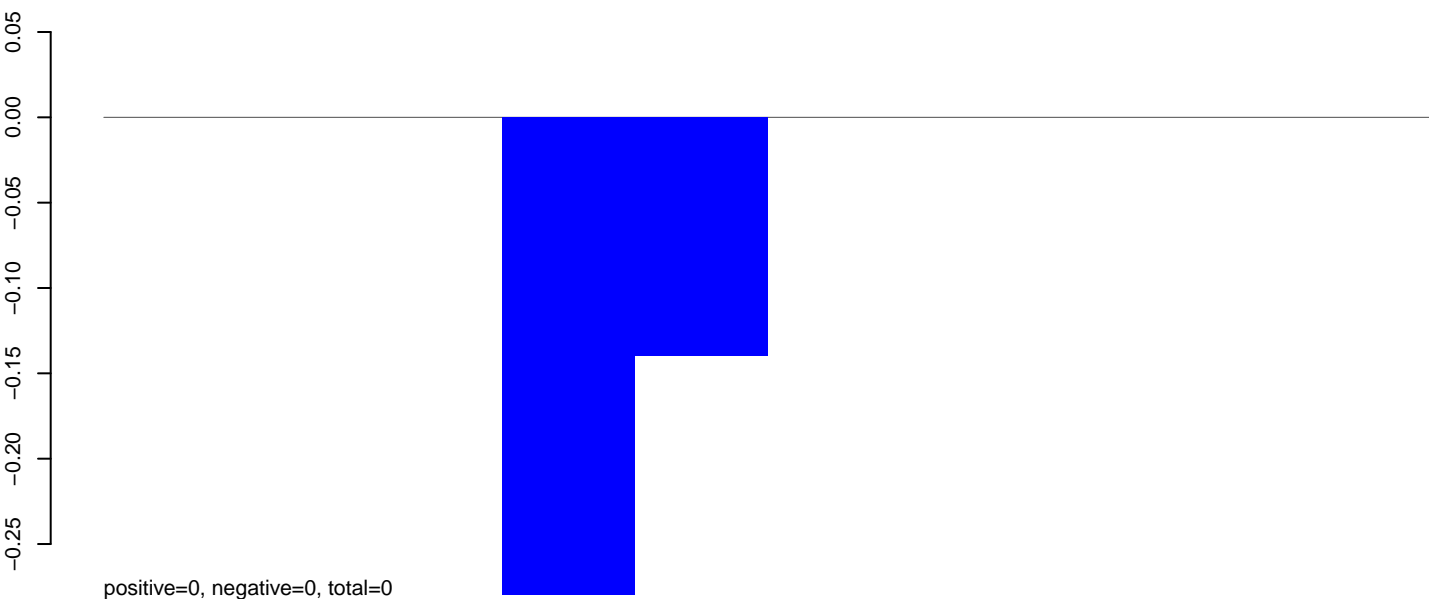
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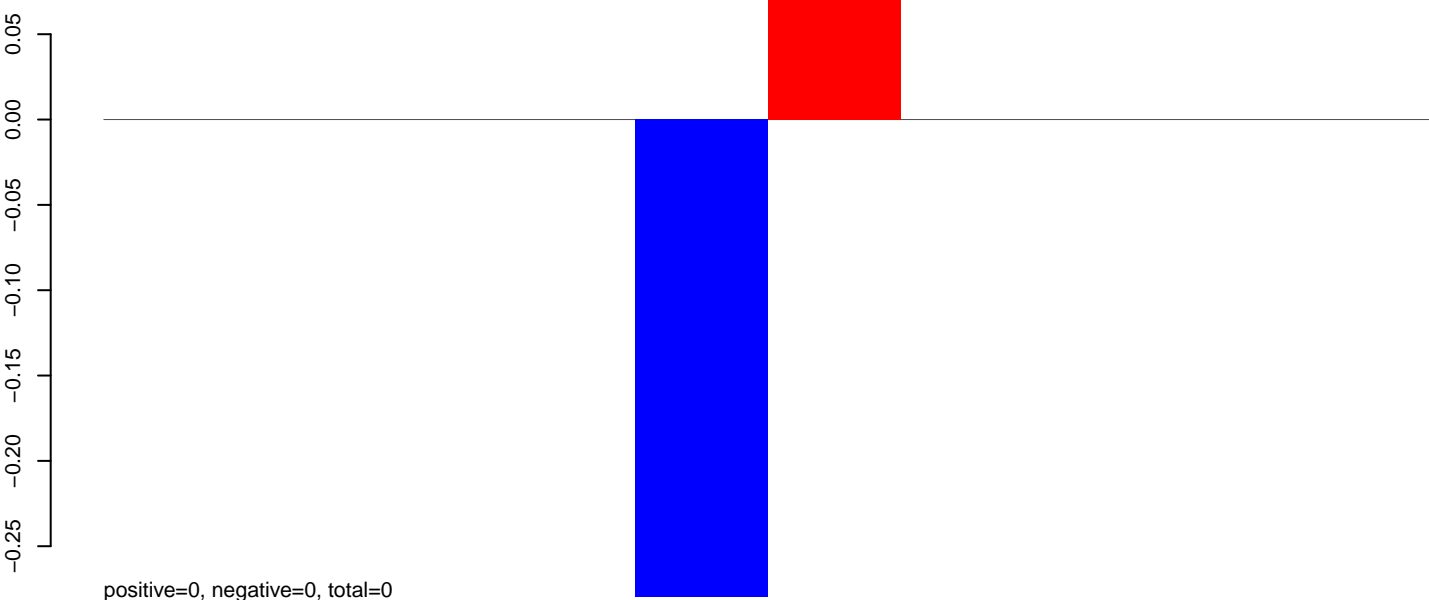
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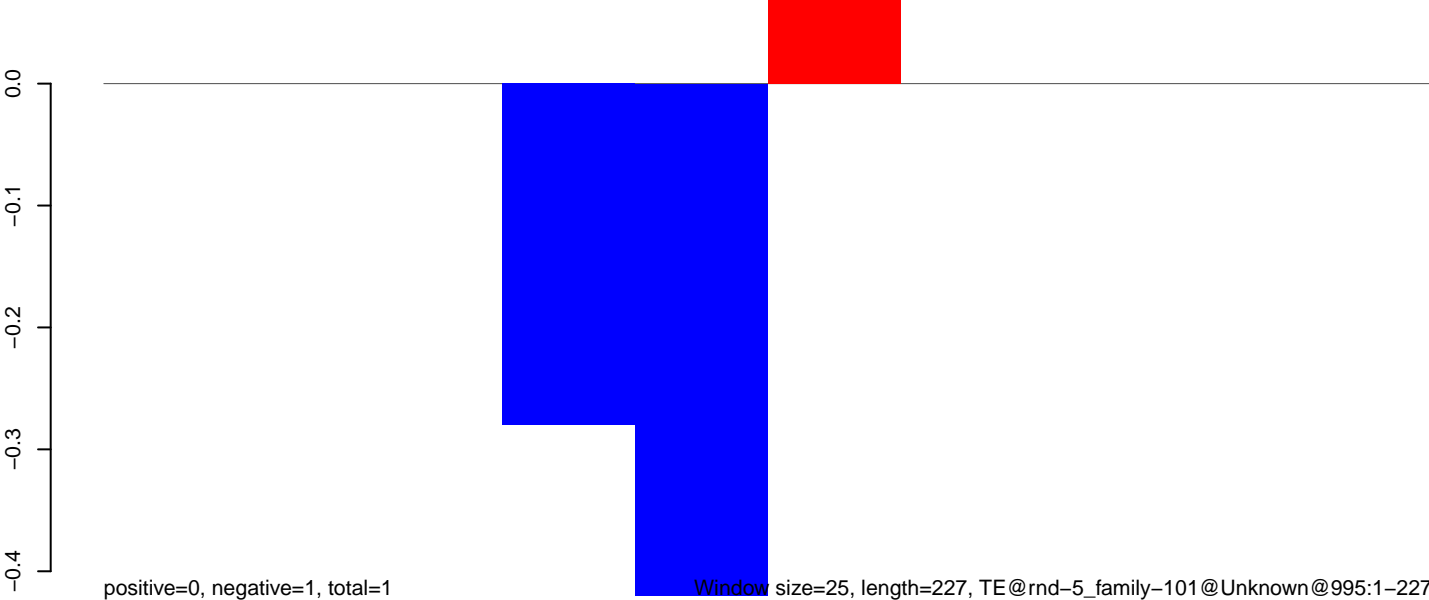
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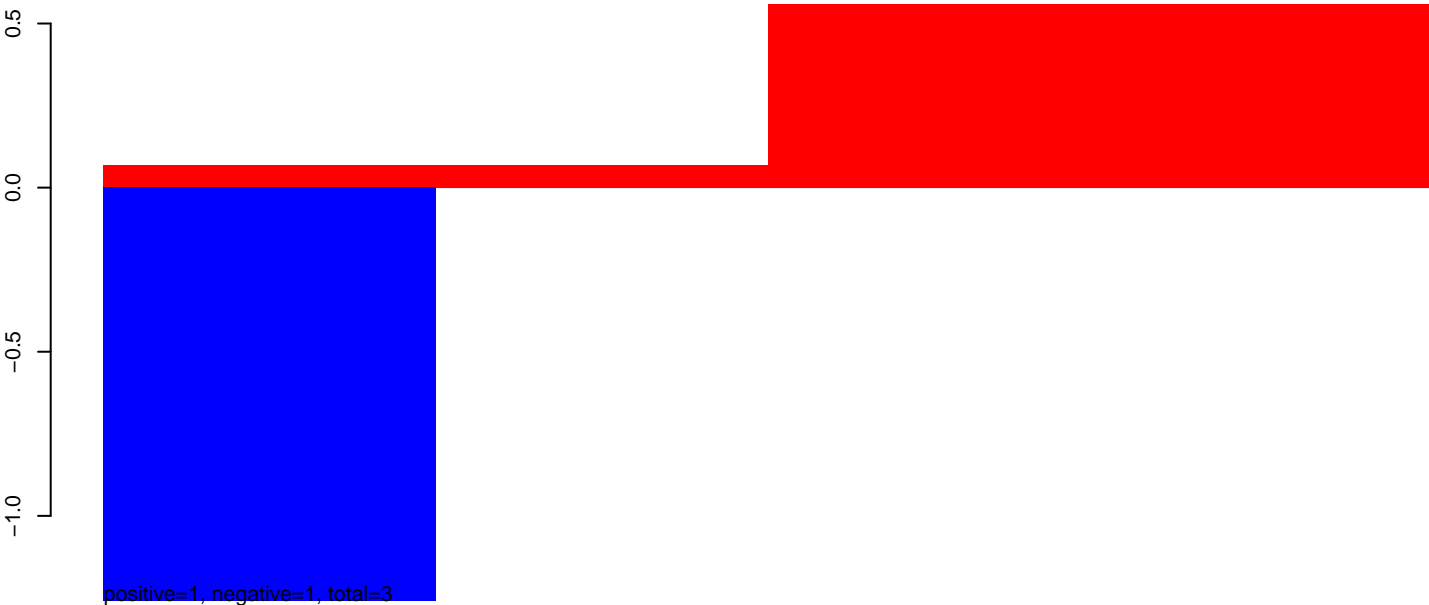
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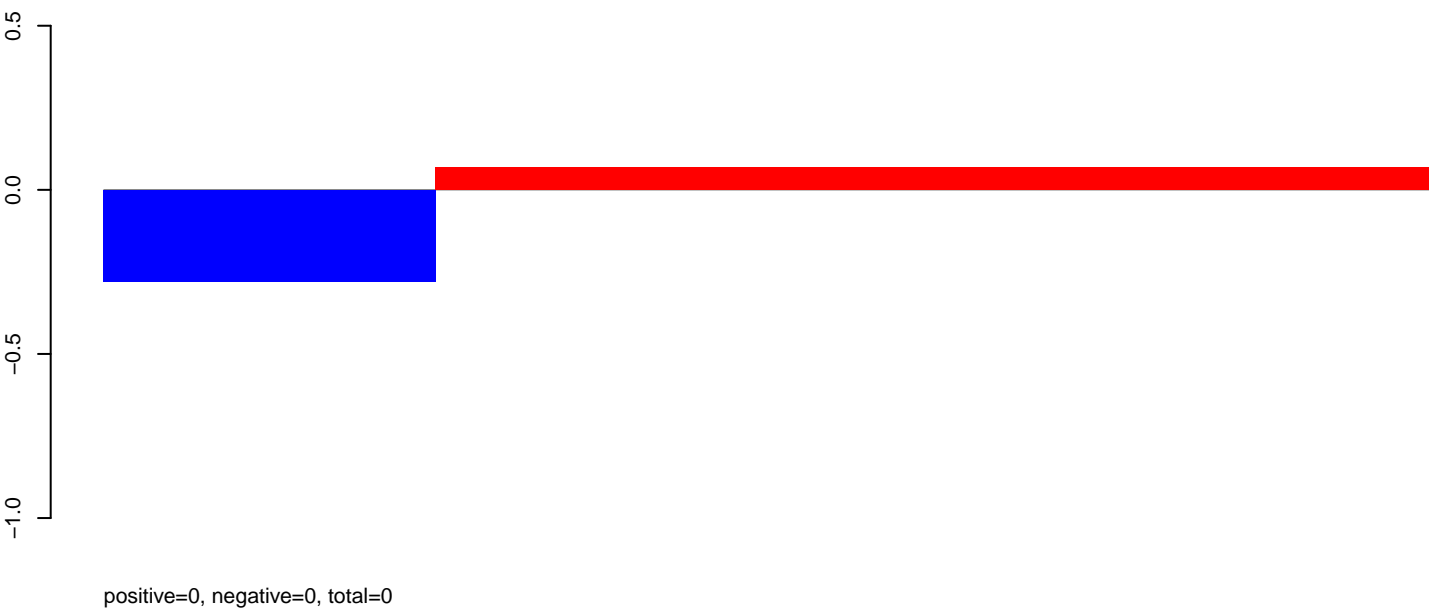
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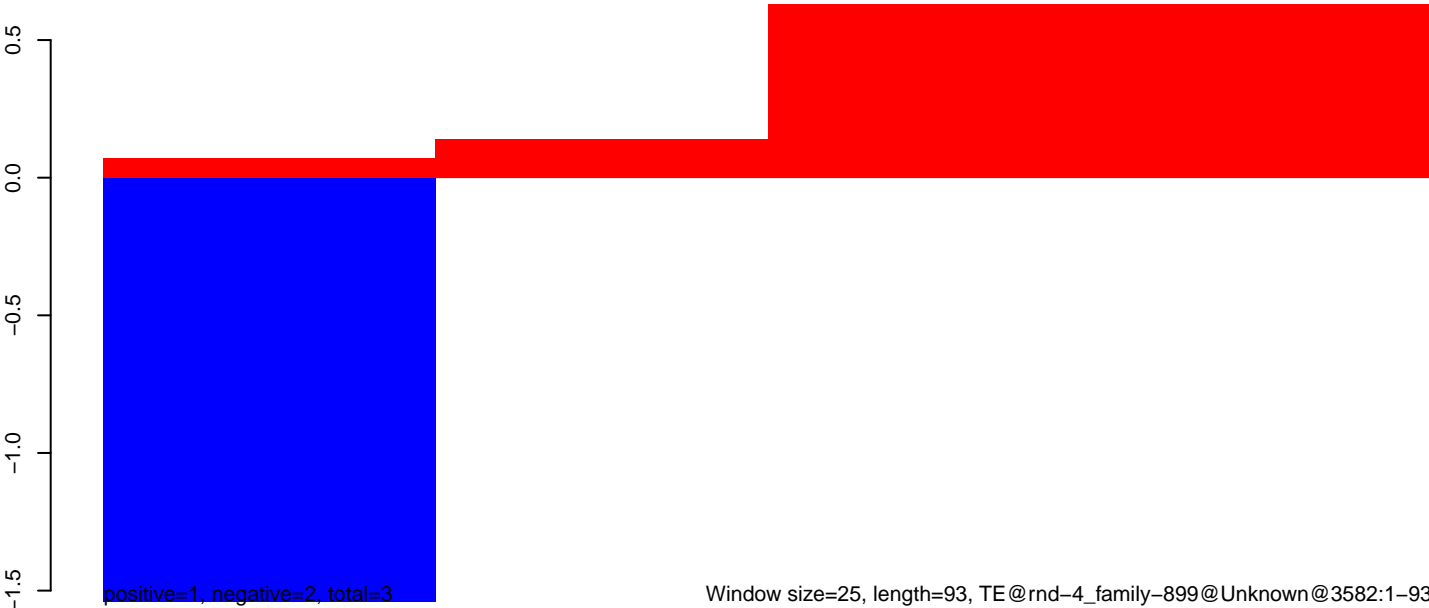
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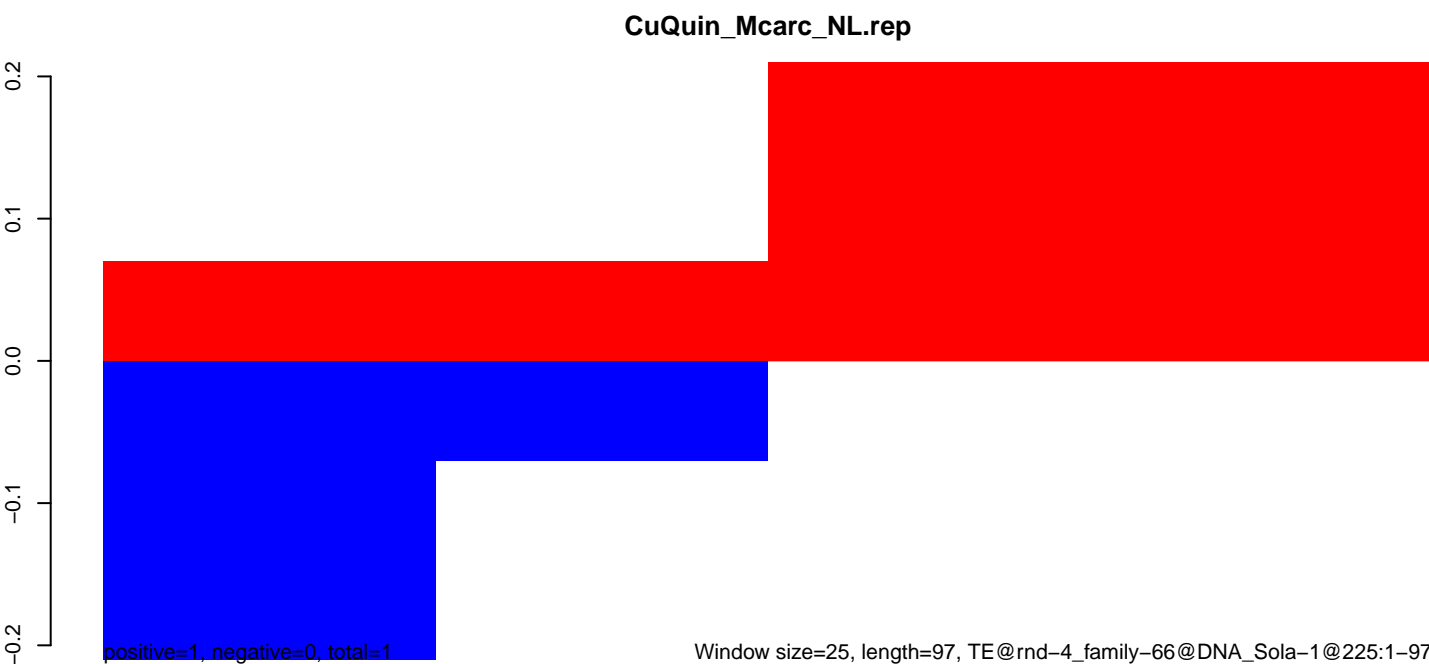
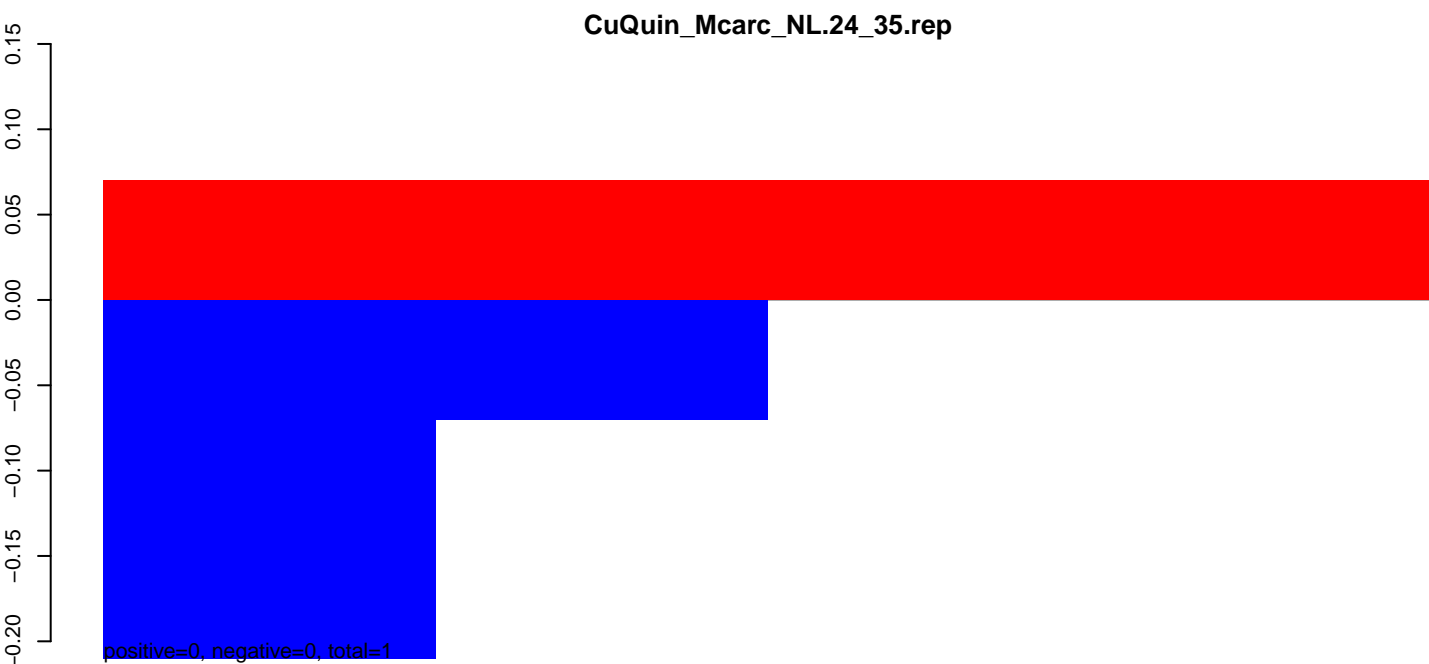
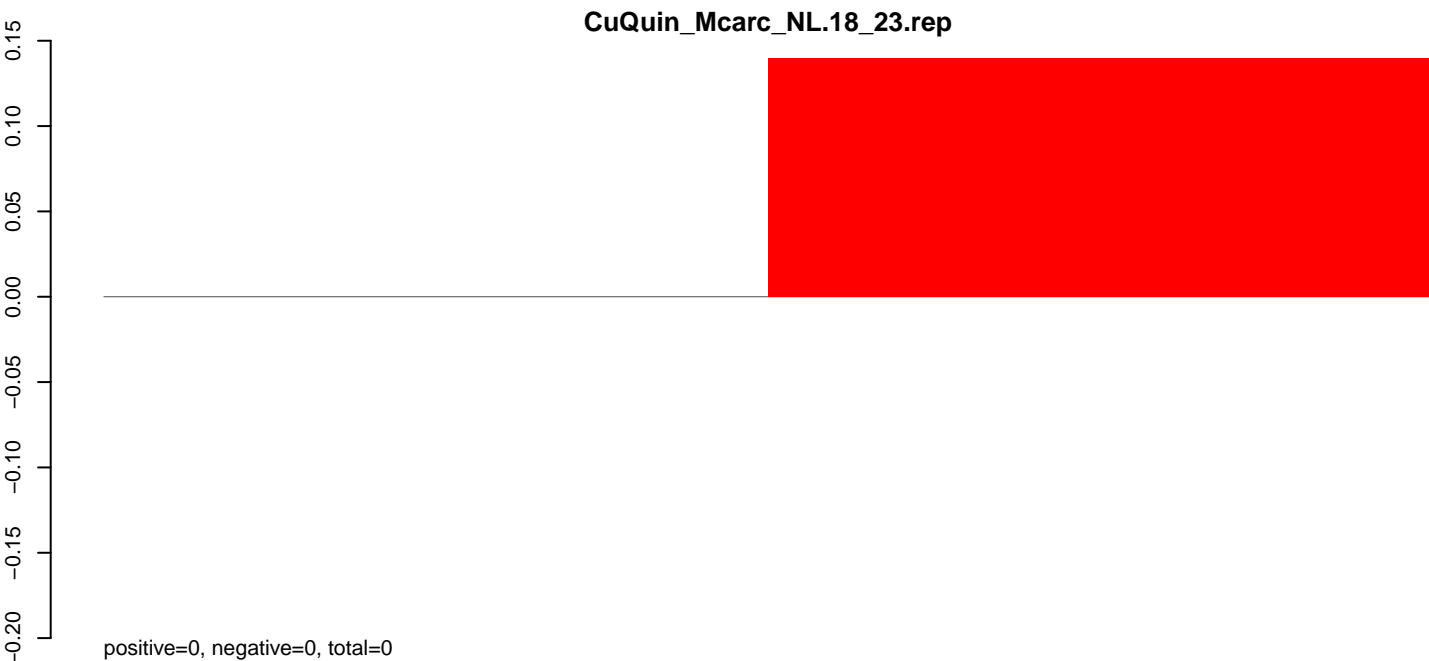


CuQuin_Mcarc_NL.24_35.rep

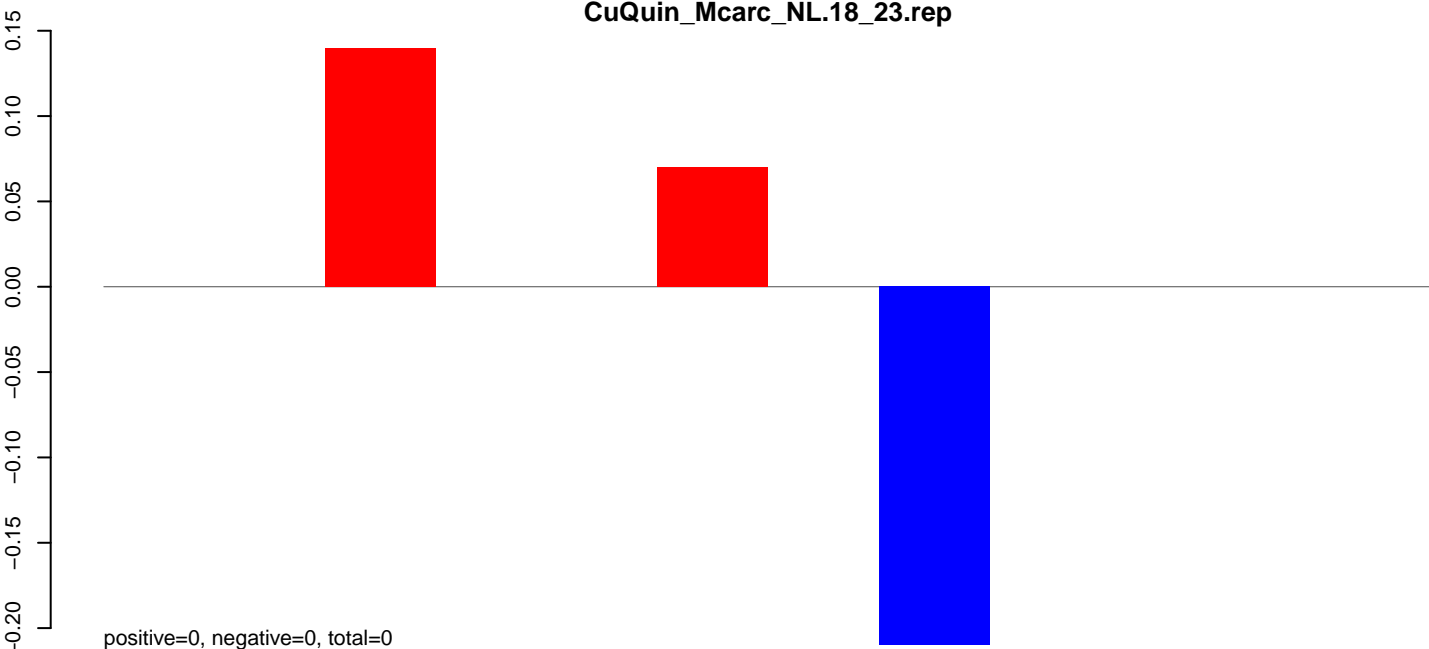


CuQuin_Mcarc_NL.rep

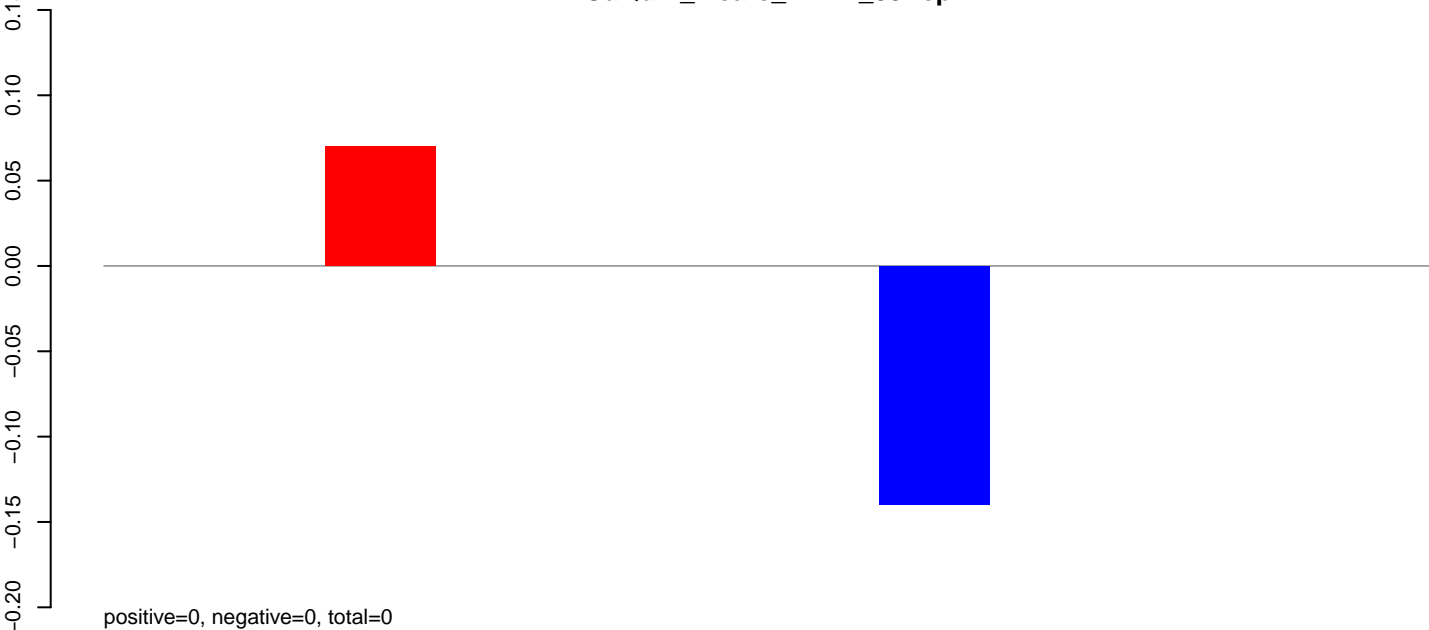




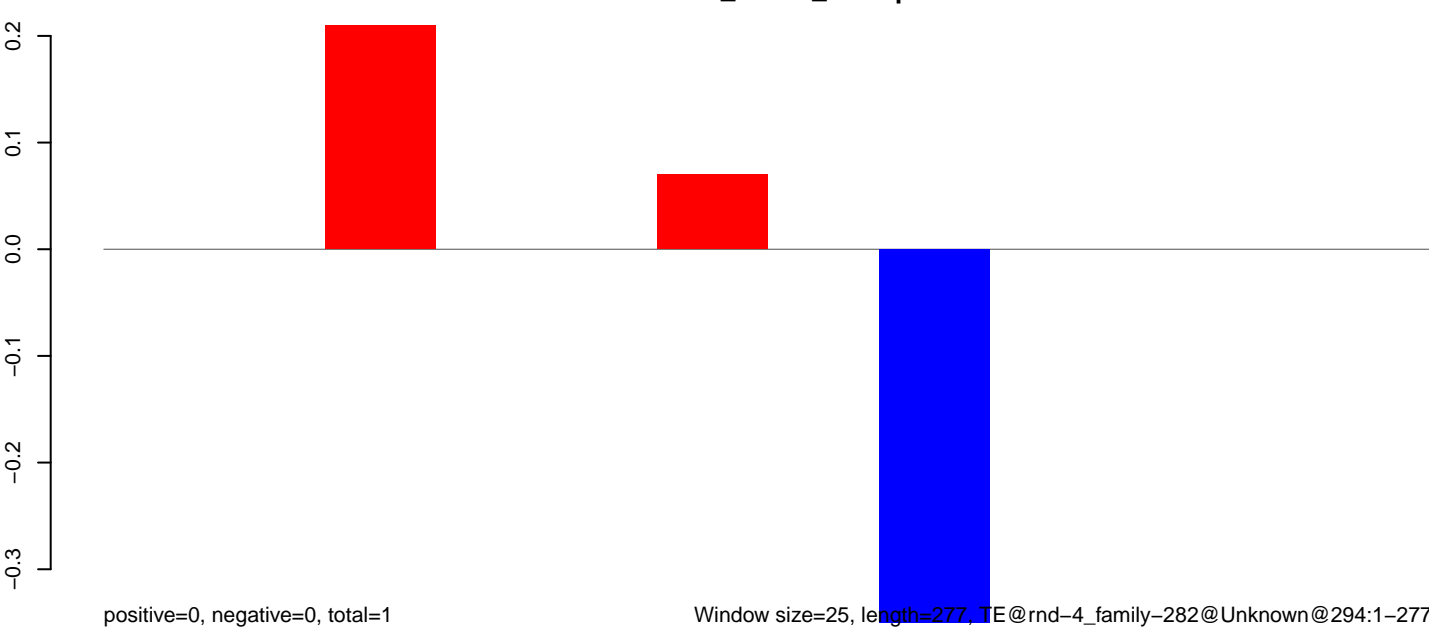
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CuQuin_Mcarc_NL.24_35.rep



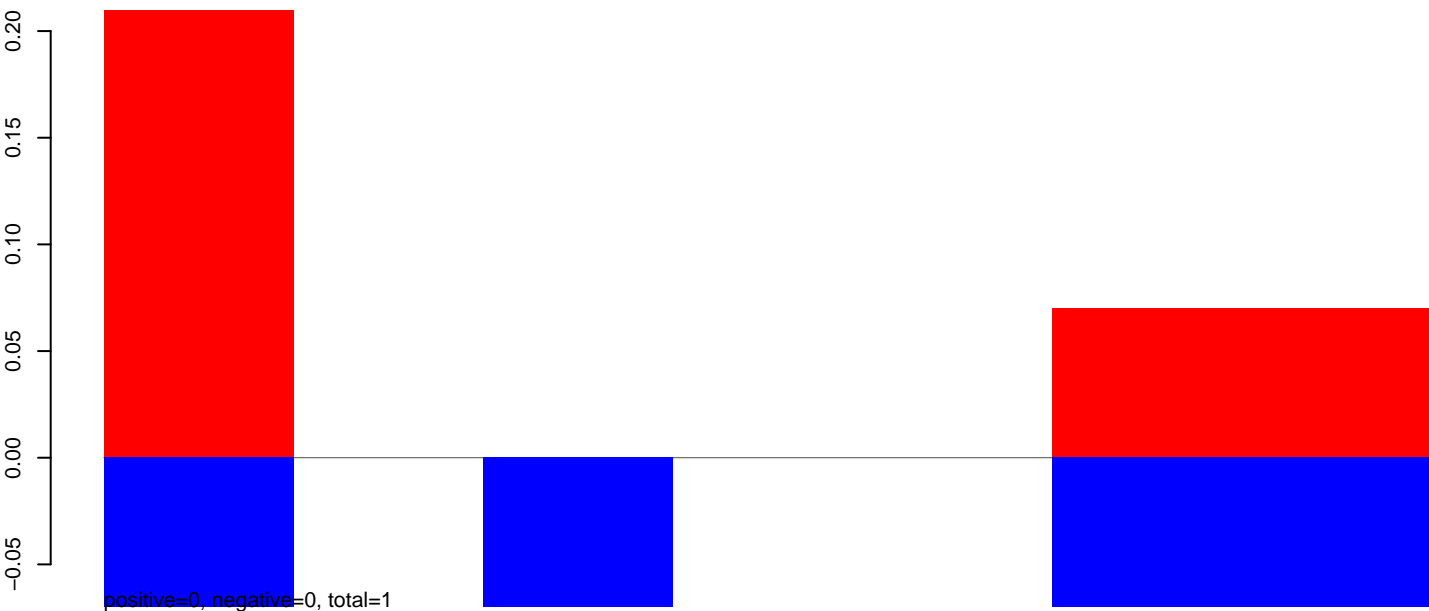
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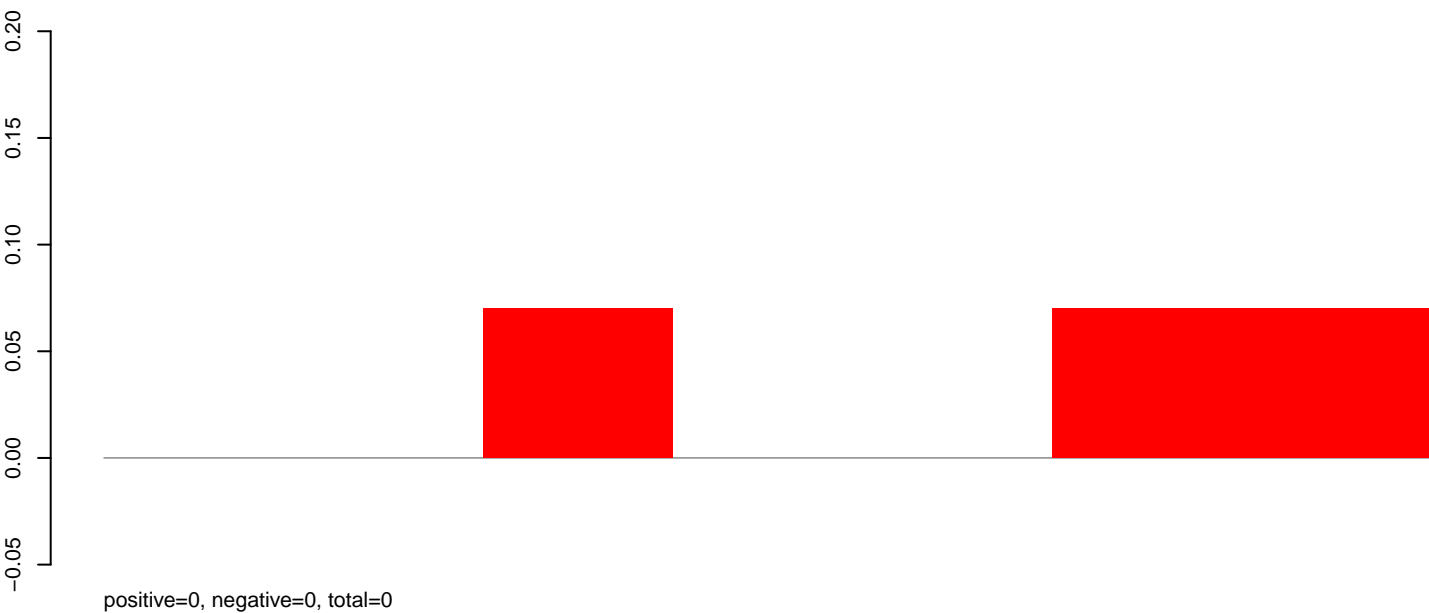
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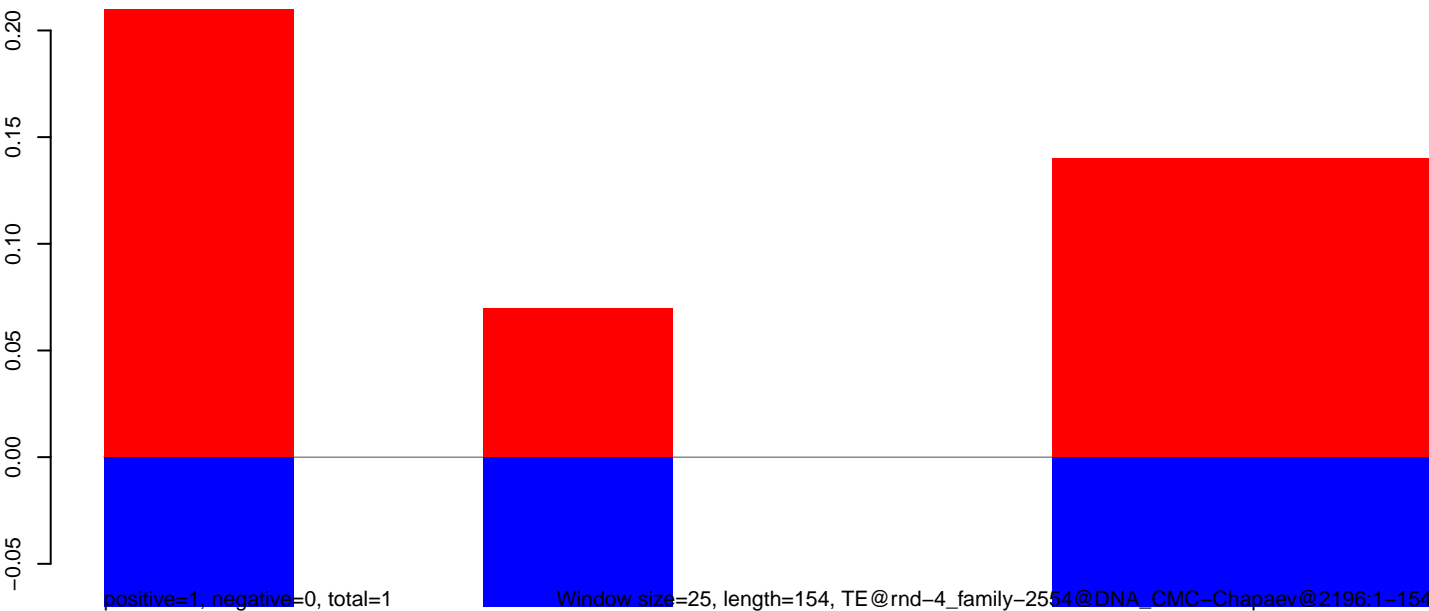
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CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



Window size=25, length=154, TE@rnd-4_family-2554@DNA_CMC-Chapaev@2196:1-154

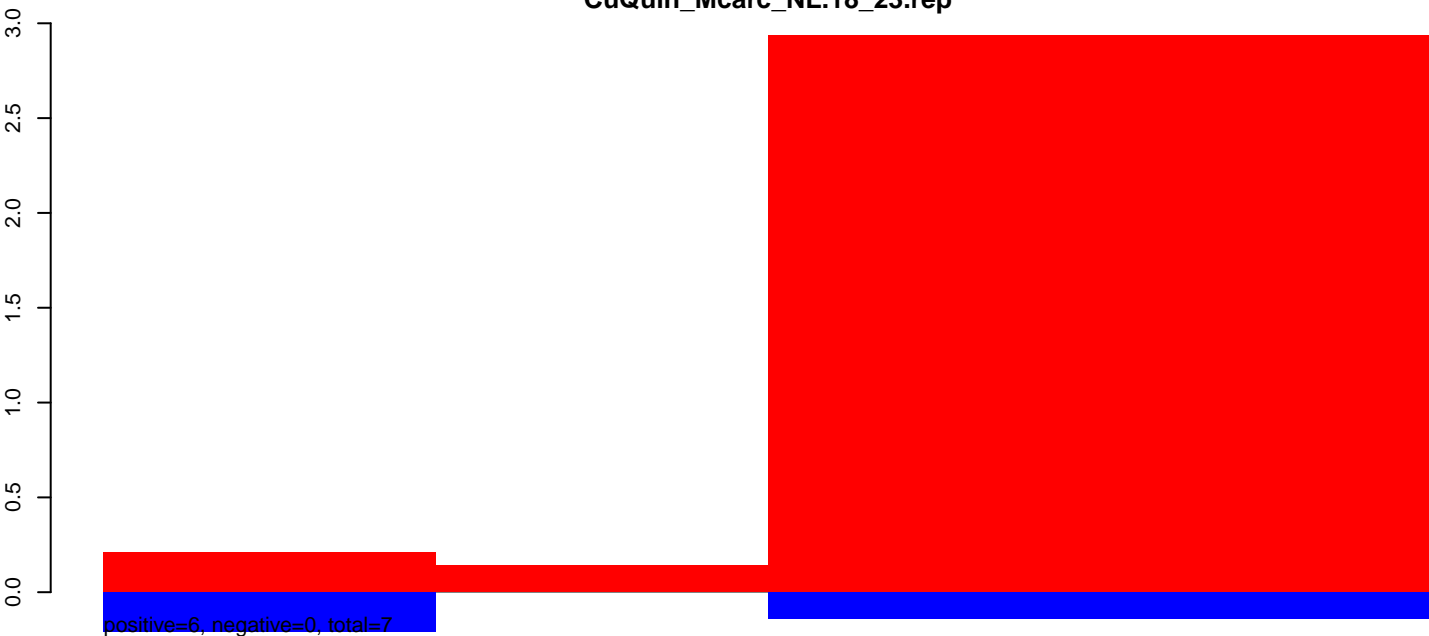
50

100

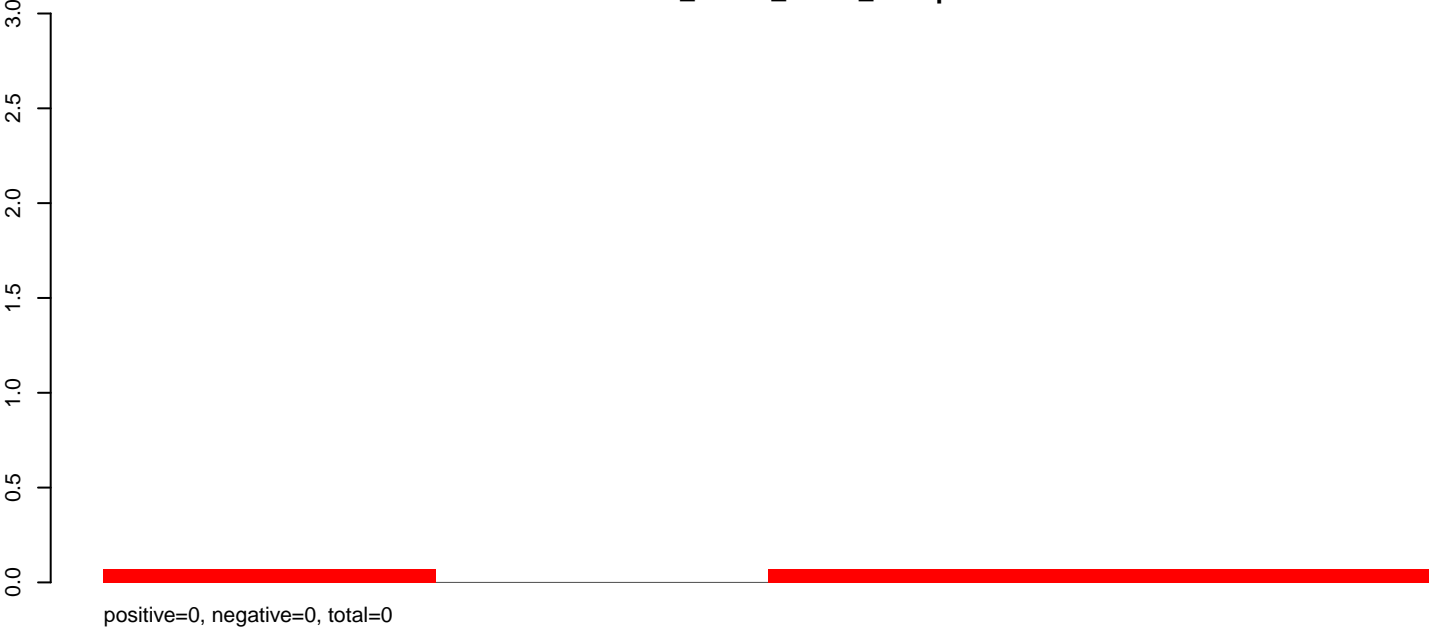
150

200

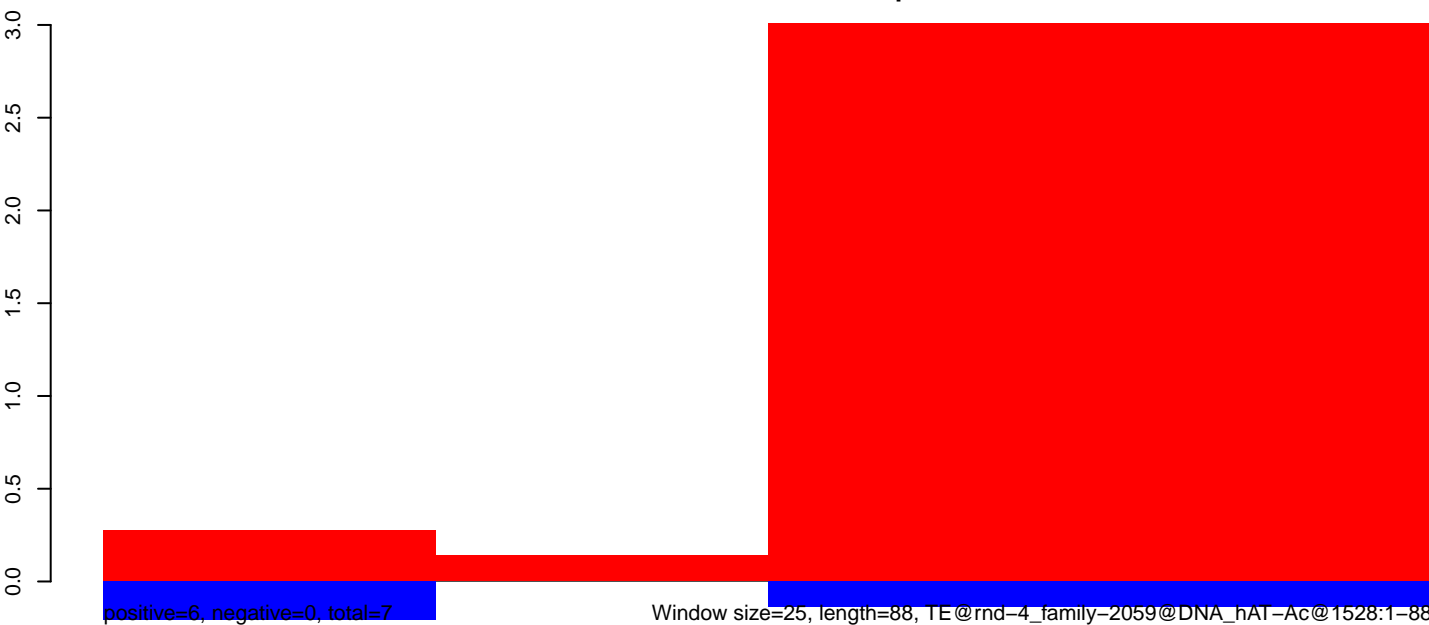
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CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



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40

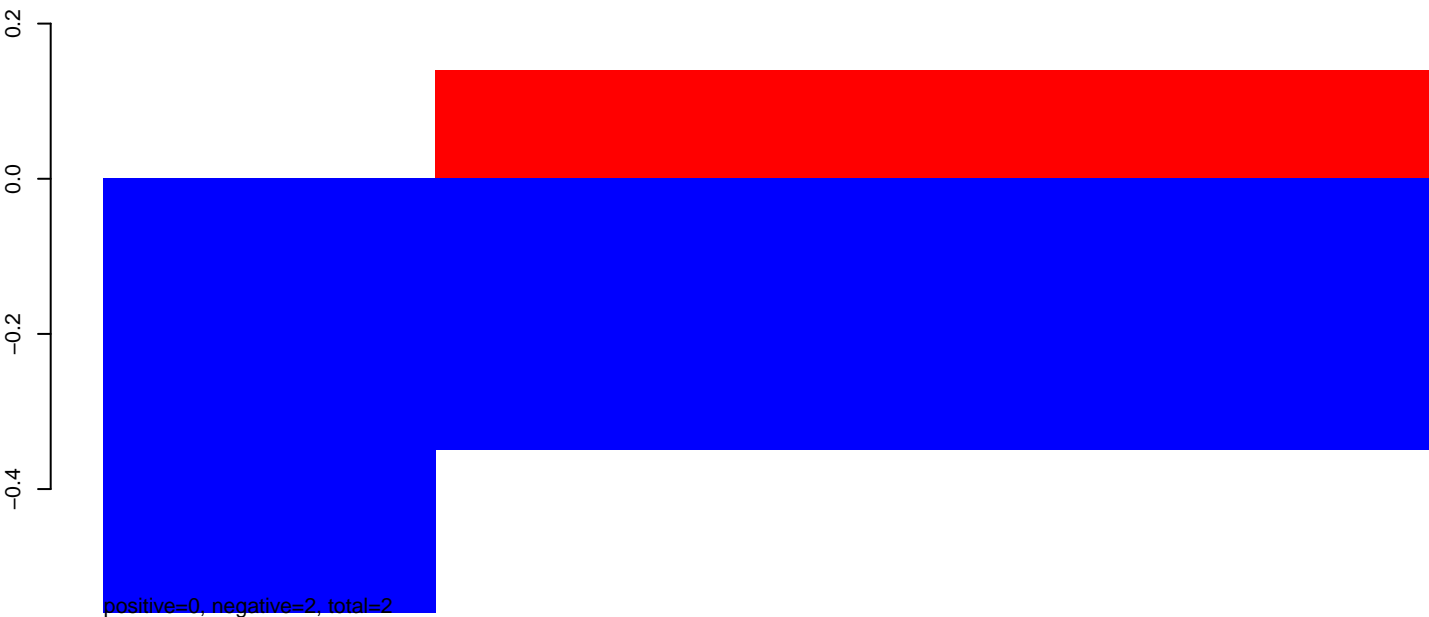
60

80

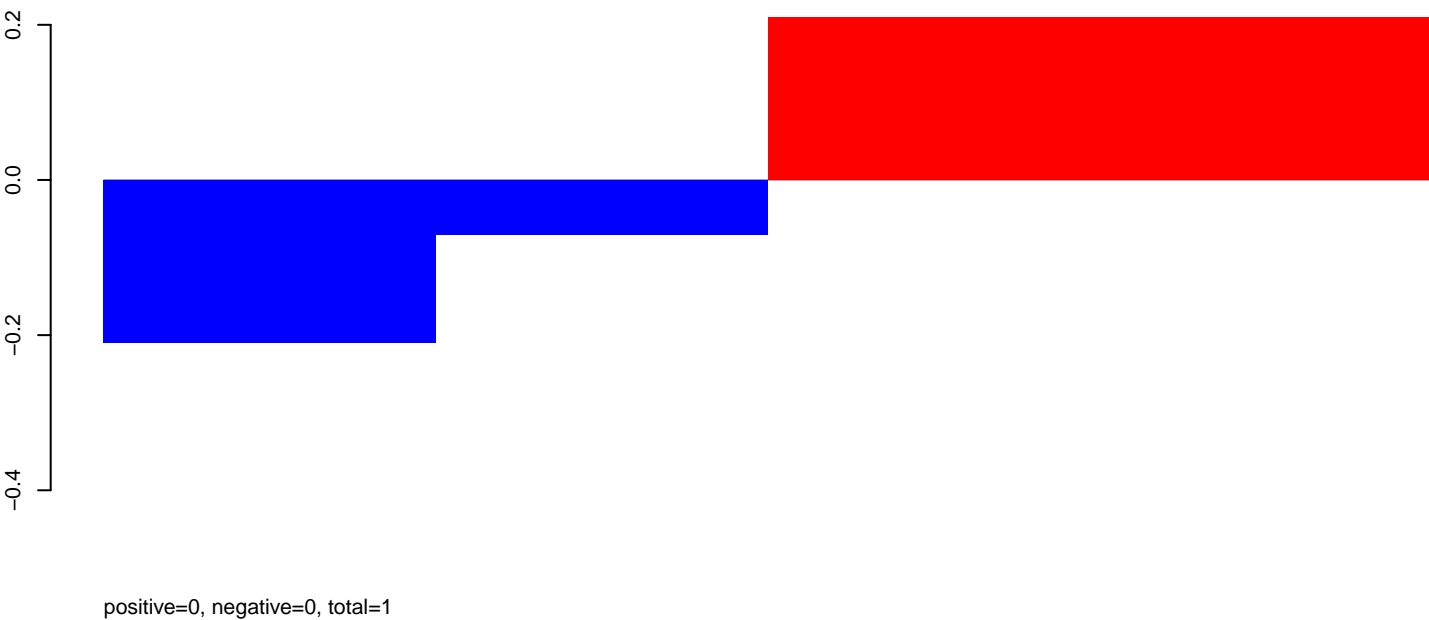
100

120

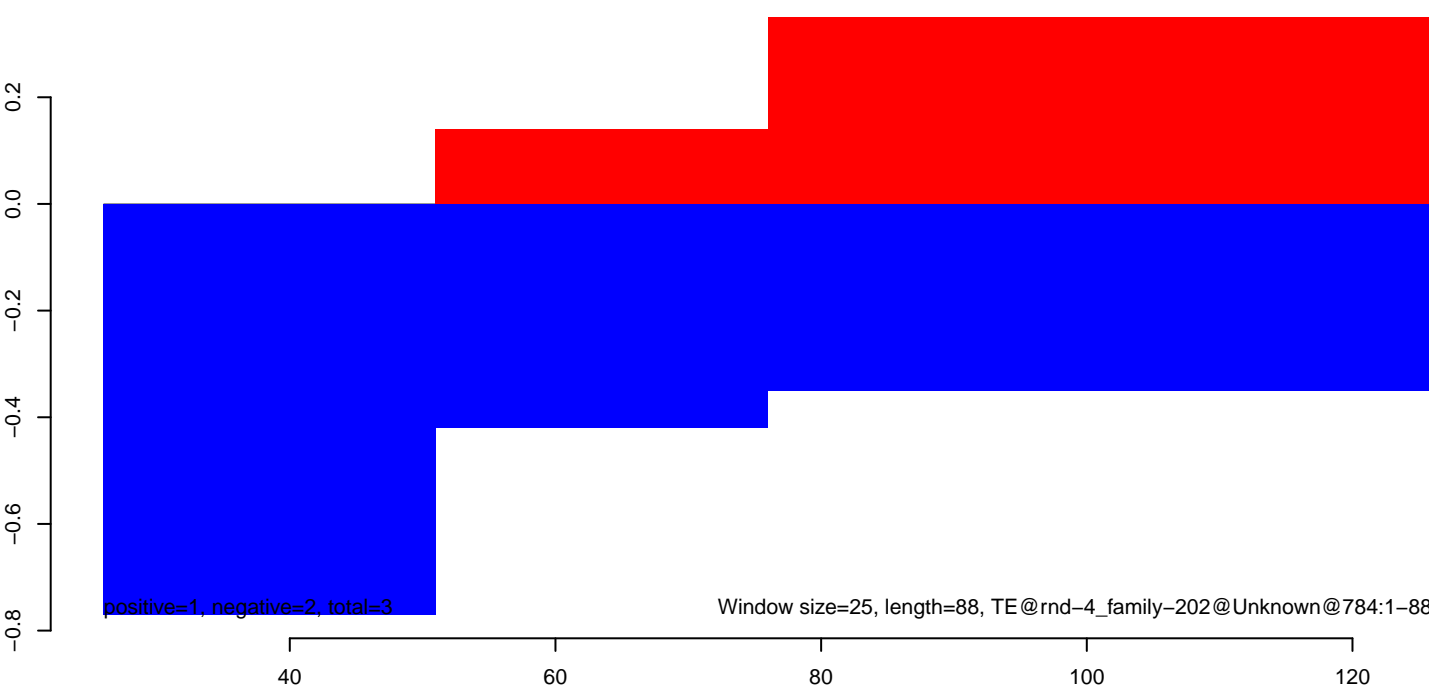
CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10
-0.12
-0.14

positive=0, negative=0, total=0

CuQuin_Mcarc_NL.24_35.rep

0.00
-0.02
-0.04
-0.06
-0.08
-0.10
-0.12
-0.14

positive=0, negative=0, total=0

CuQuin_Mcarc_NL.rep

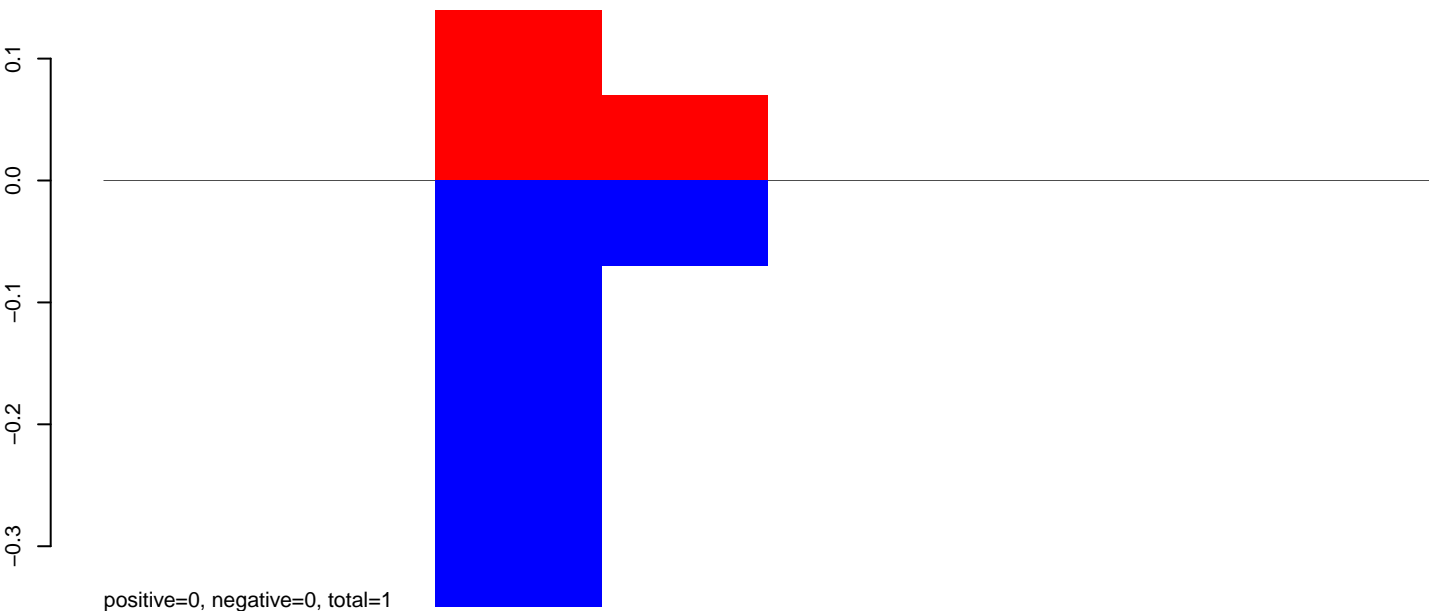
0.00
-0.02
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-0.10
-0.12
-0.14

positive=0, negative=0, total=0

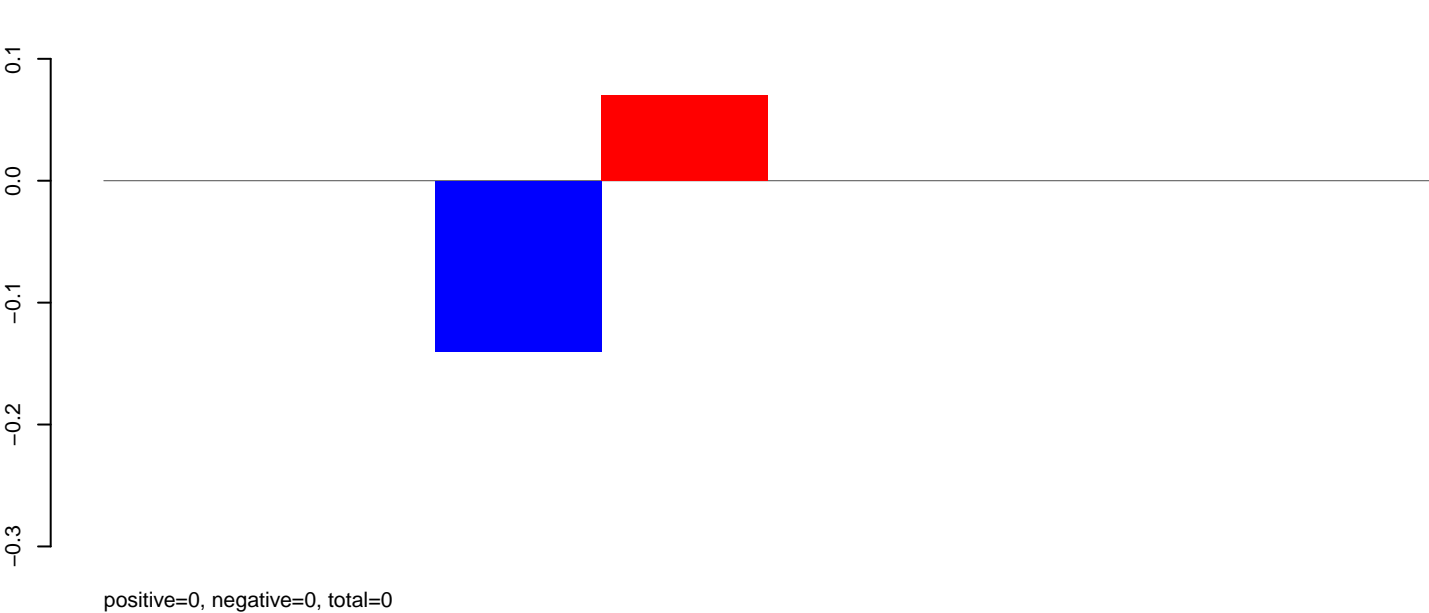
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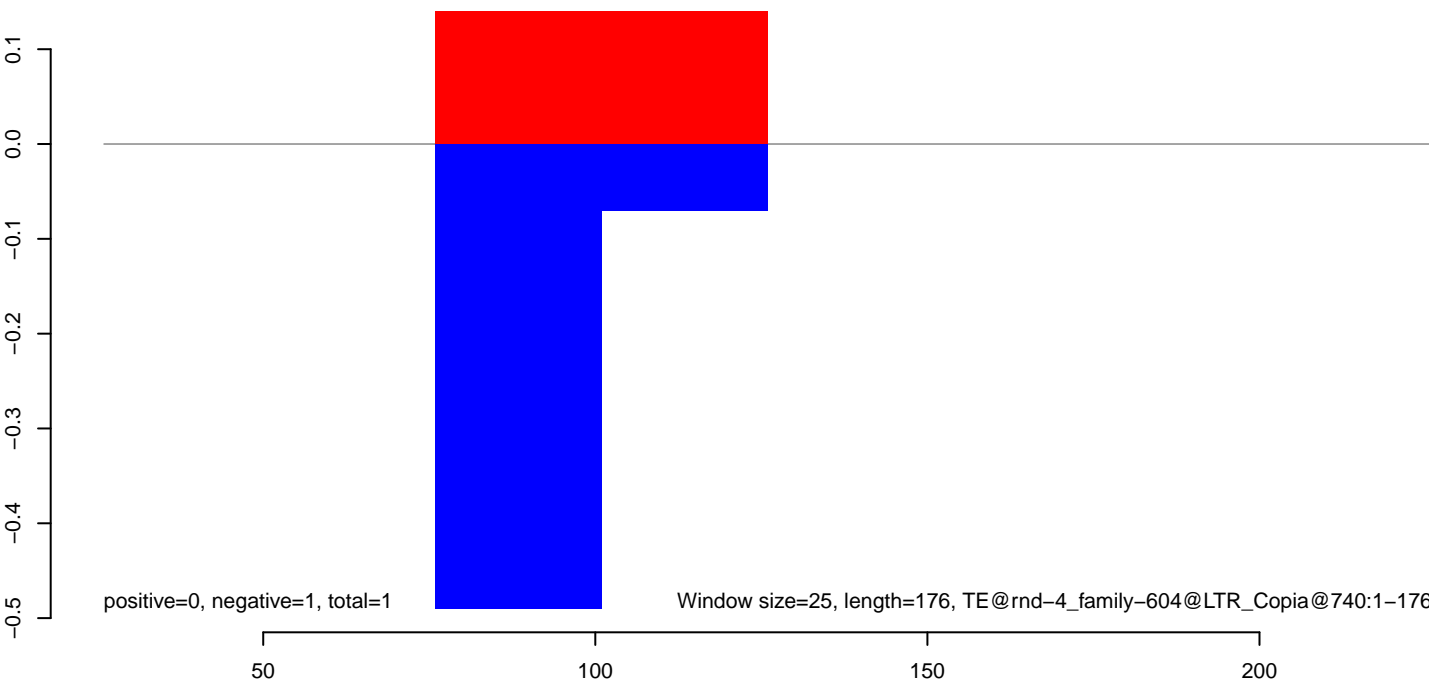
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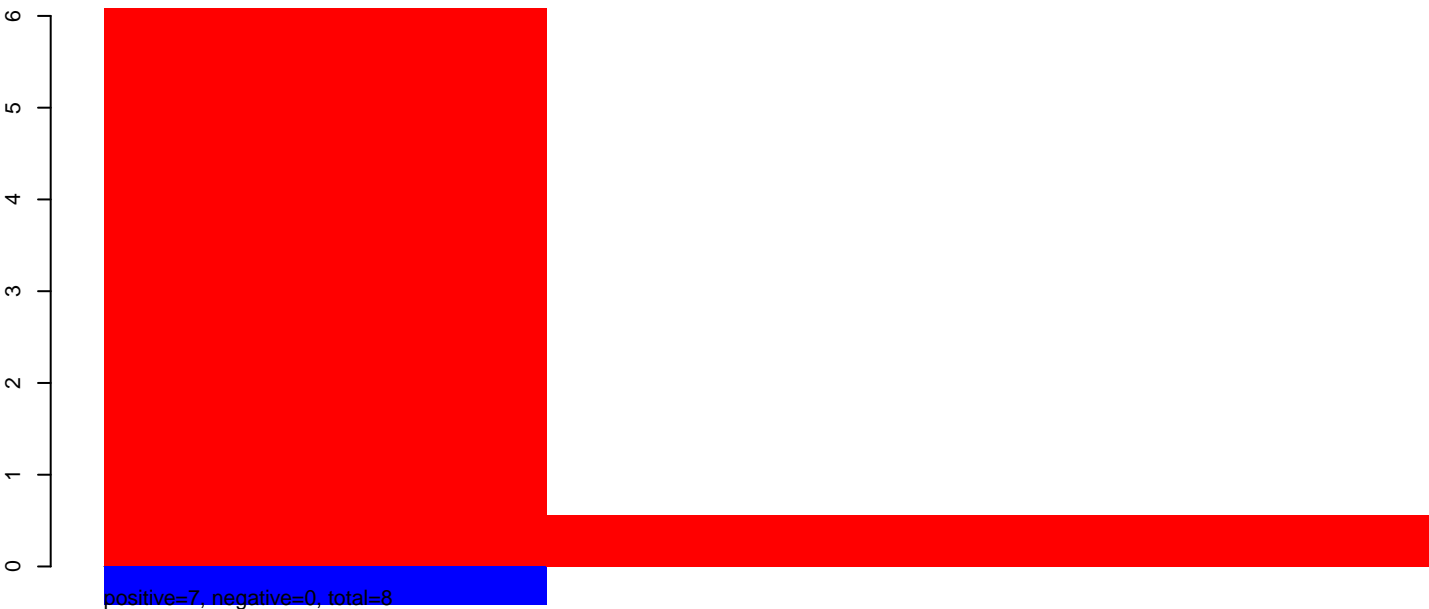
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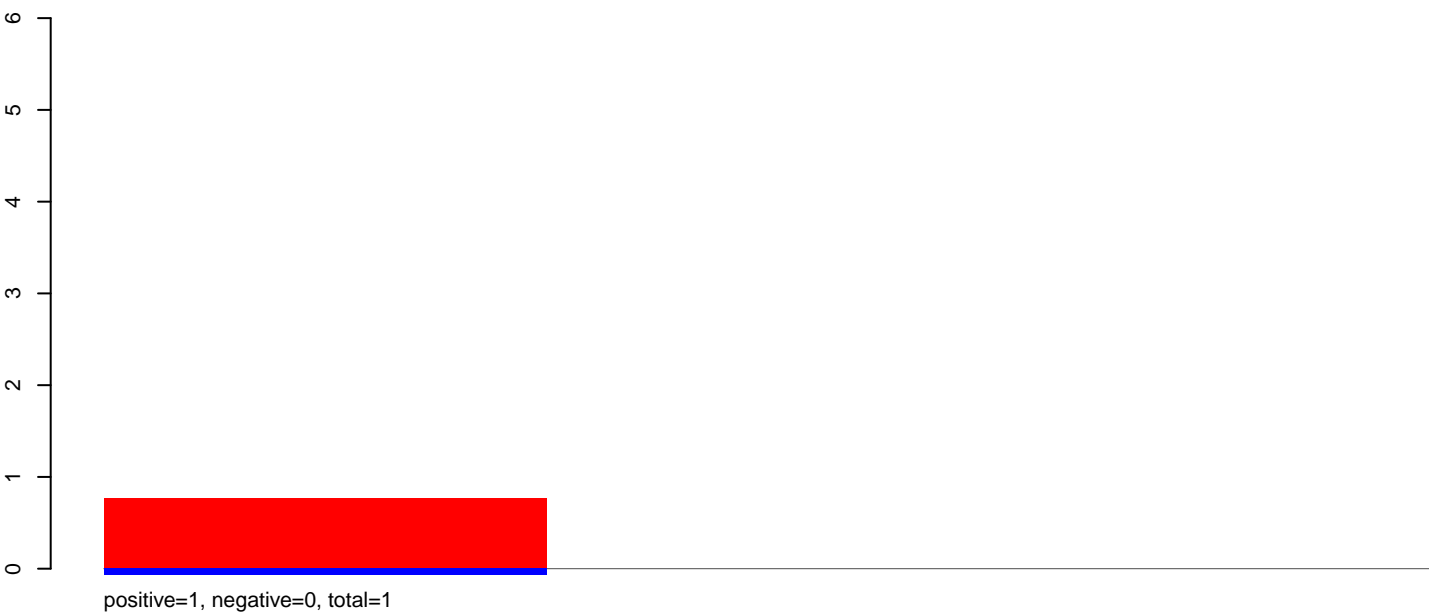
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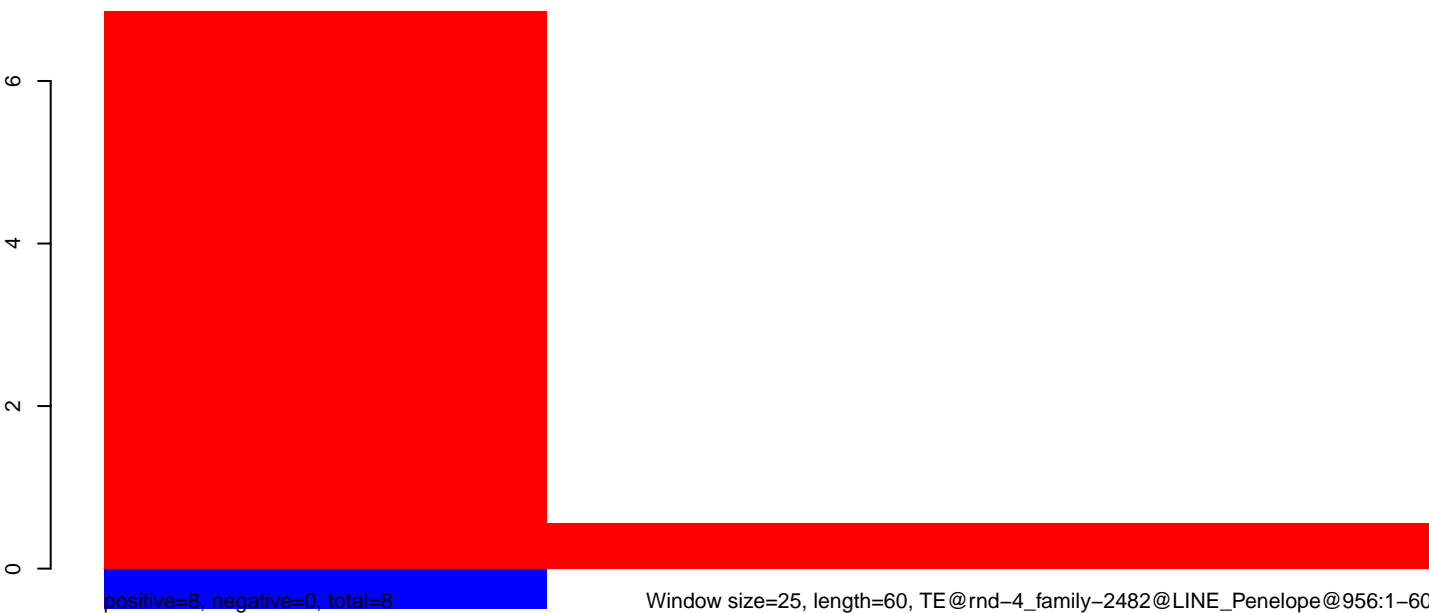
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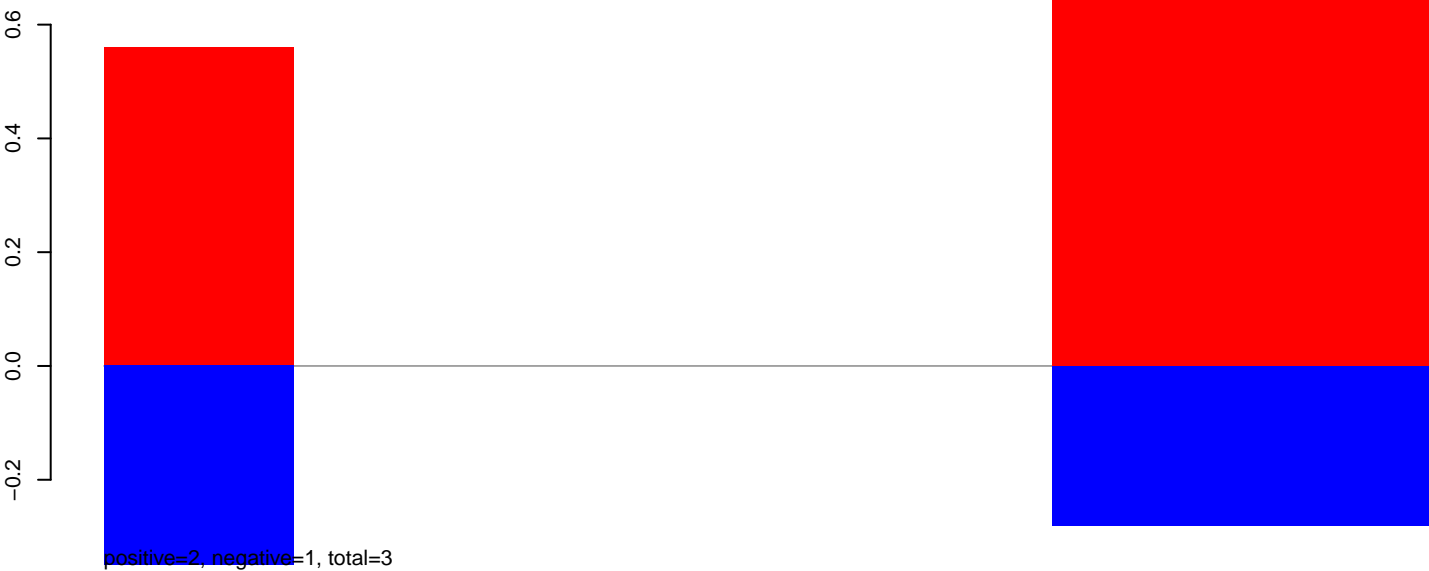
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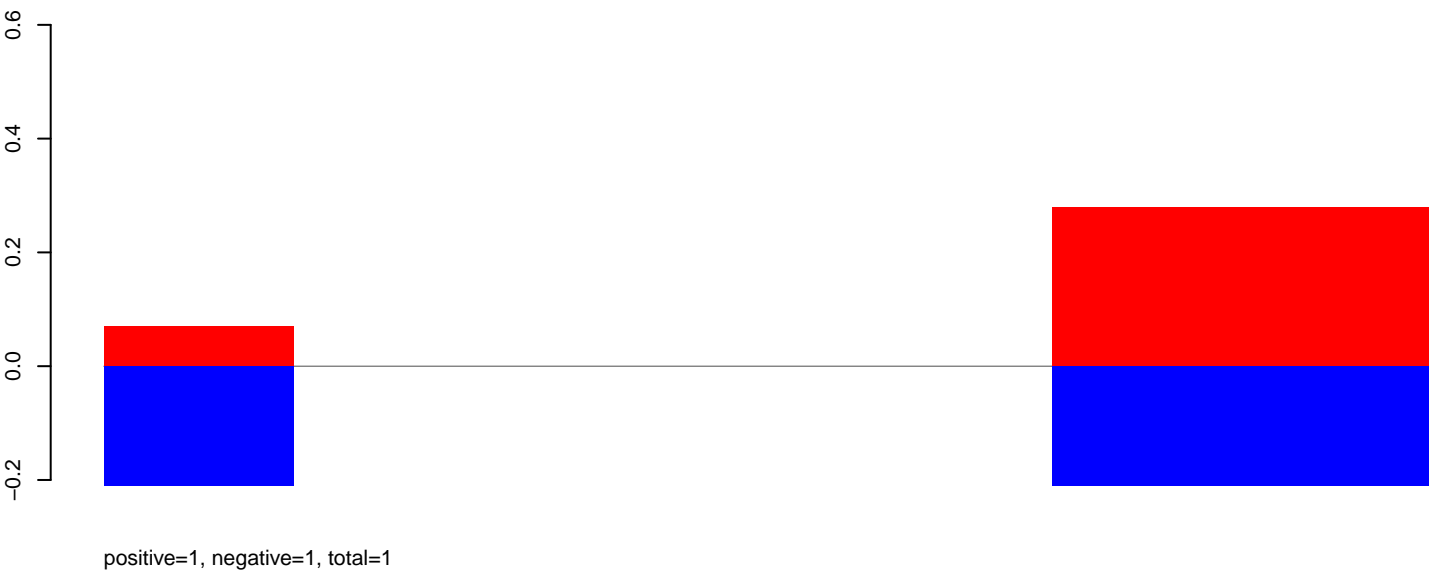
CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



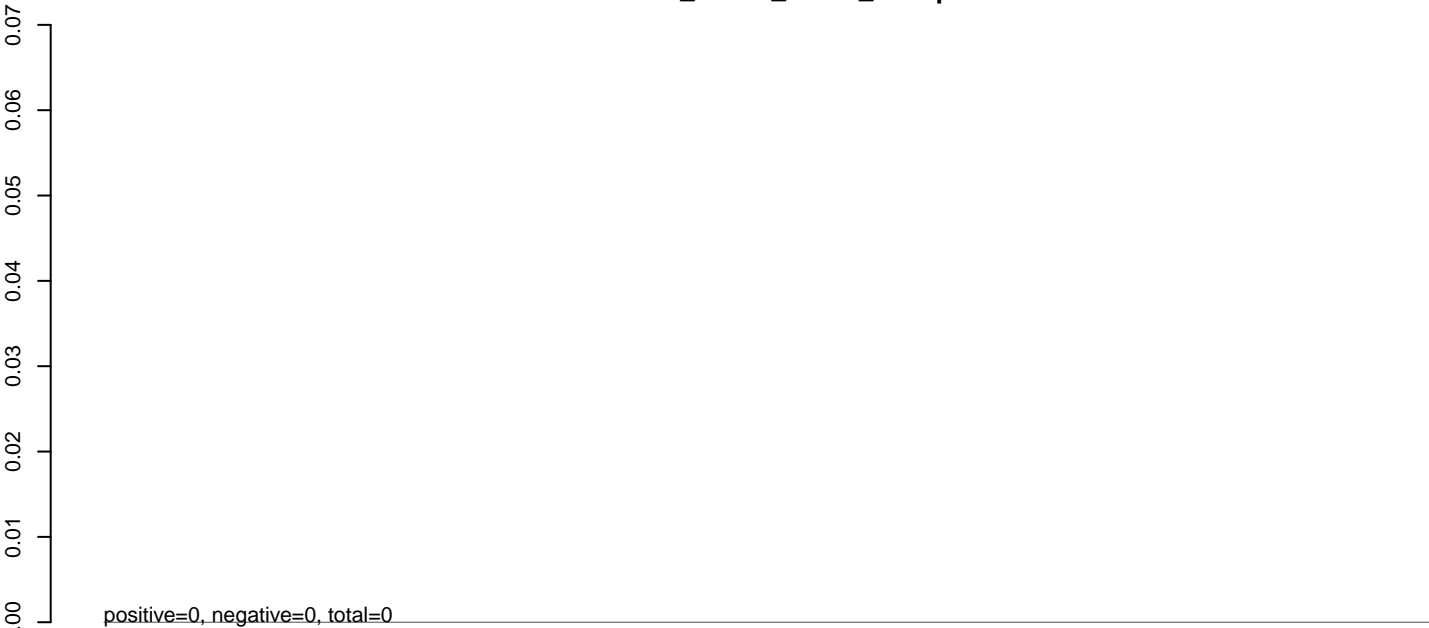
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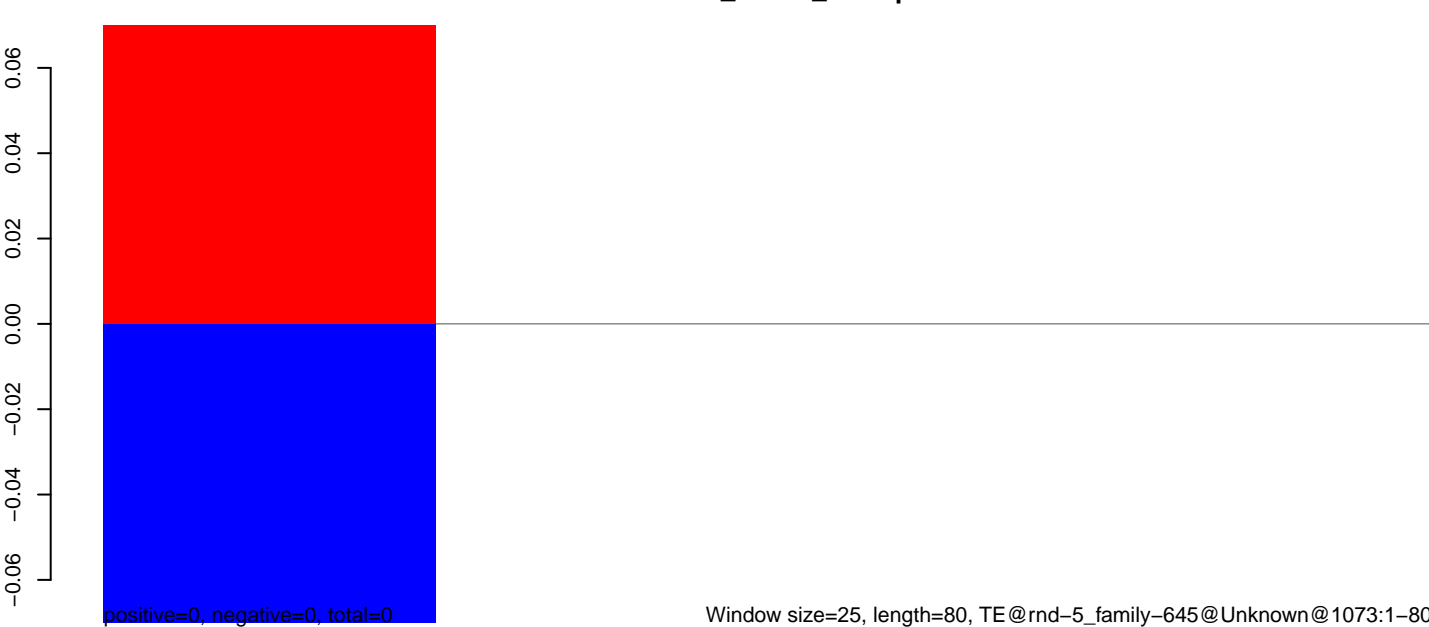
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CuQuin_Mcarc_NL.24_35.rep



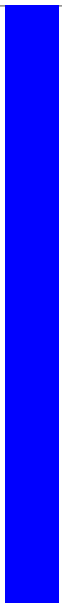
CuQuin_Mcarc_NL.rep



CuQuin_Mcarc_NL.18_23.rep

0.00
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-0.02
-0.03
-0.04
-0.05
-0.06
-0.07

positive=0, negative=0, total=0



CuQuin_Mcarc_NL.24_35.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06
-0.07

positive=0, negative=0, total=0

CuQuin_Mcarc_NL.rep

0.00
-0.01
-0.02
-0.03
-0.04
-0.05
-0.06
-0.07

positive=0, negative=0, total=0

Window size=25, length=620, TE@rnd-5_family-3807@Unknown@2547:1-620



100 200 300 400 500 600

CuQuin_Mcarc_NL.18_23.rep



CuQuin_Mcarc_NL.24_35.rep



CuQuin_Mcarc_NL.rep

