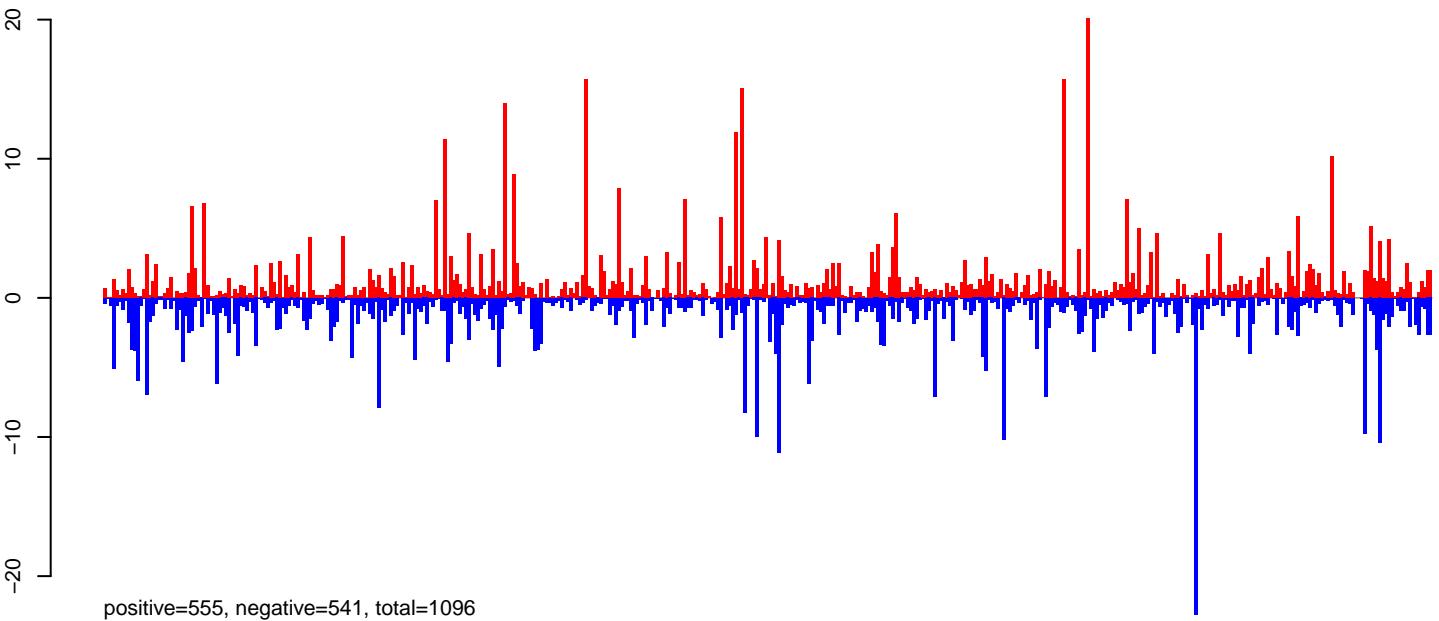
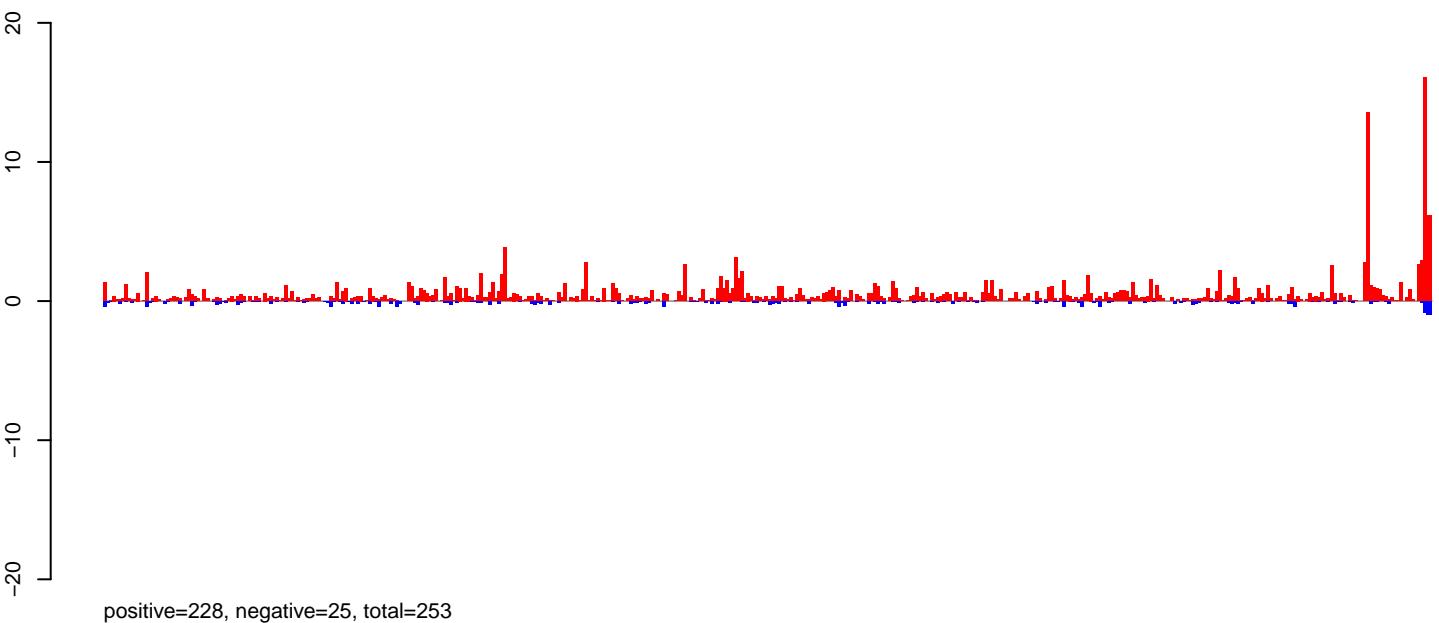


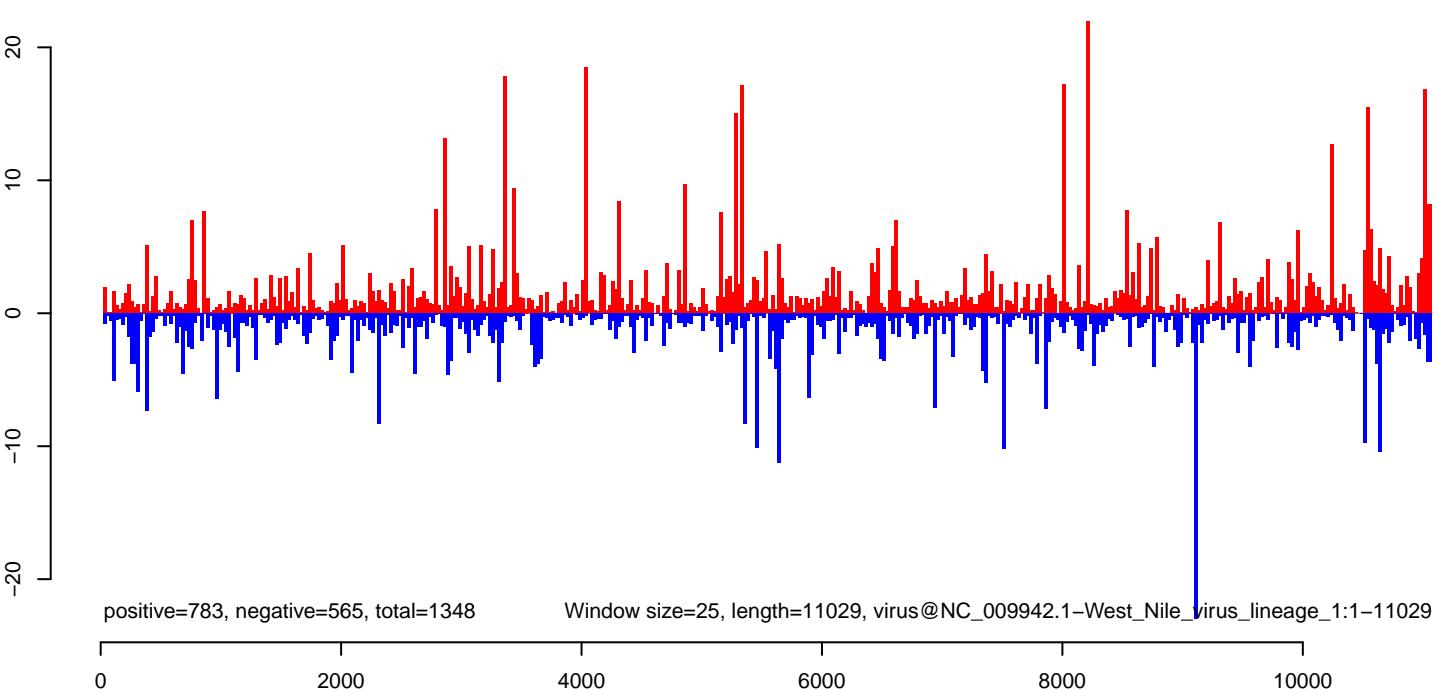
CuQuin_MG_7dpi3.18_23.rep



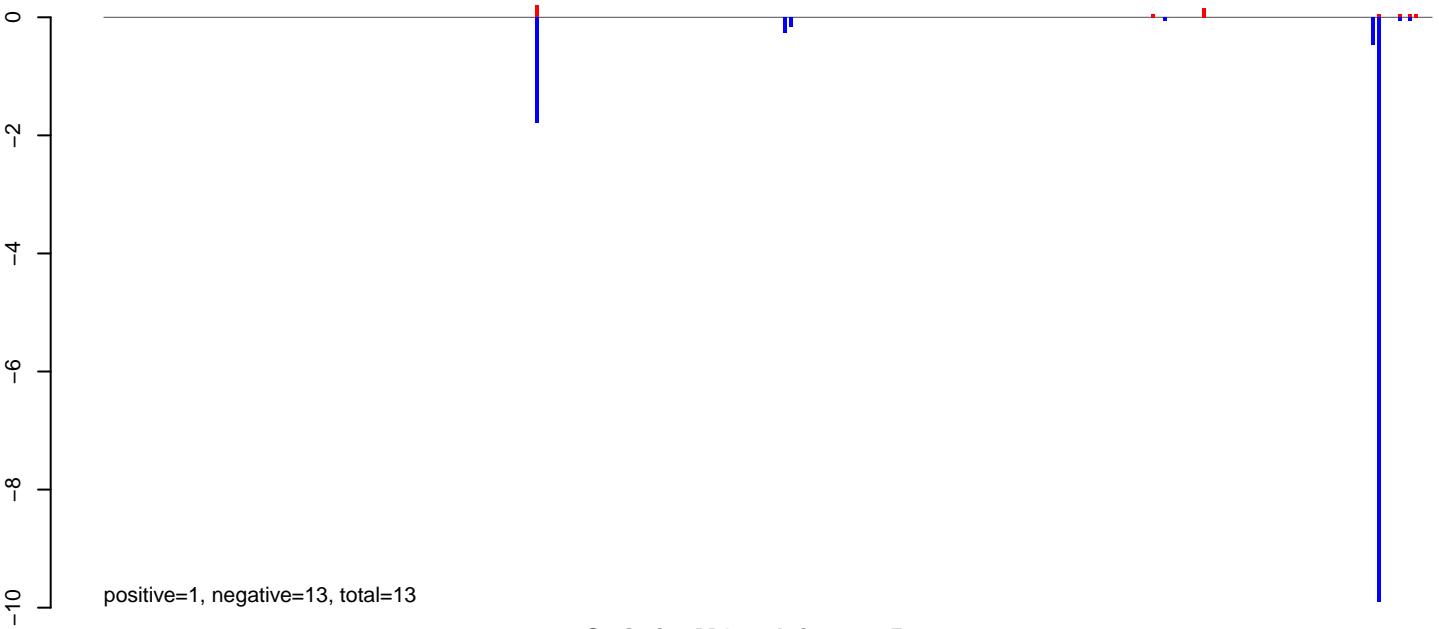
CuQuin_MG_7dpi3.24_35.rep



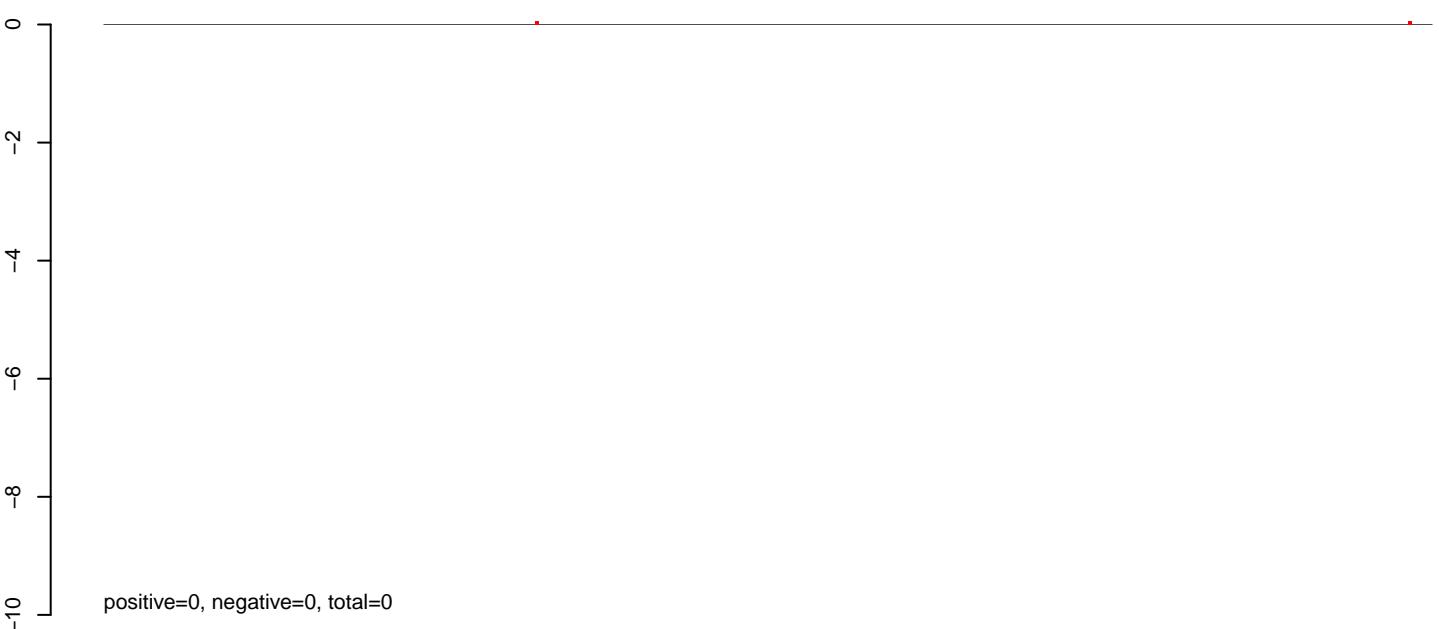
CuQuin_MG_7dpi3.rep



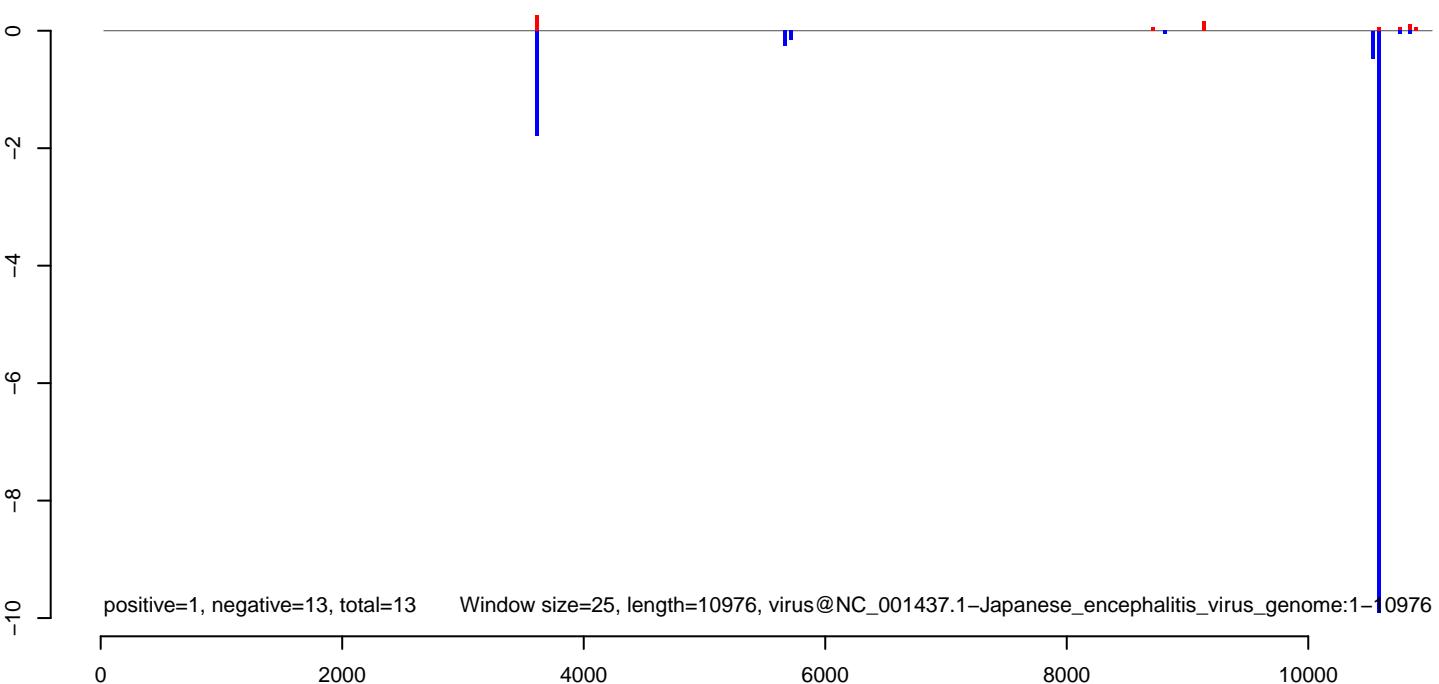
CuQuin_MG_7dpi3.18_23.rep



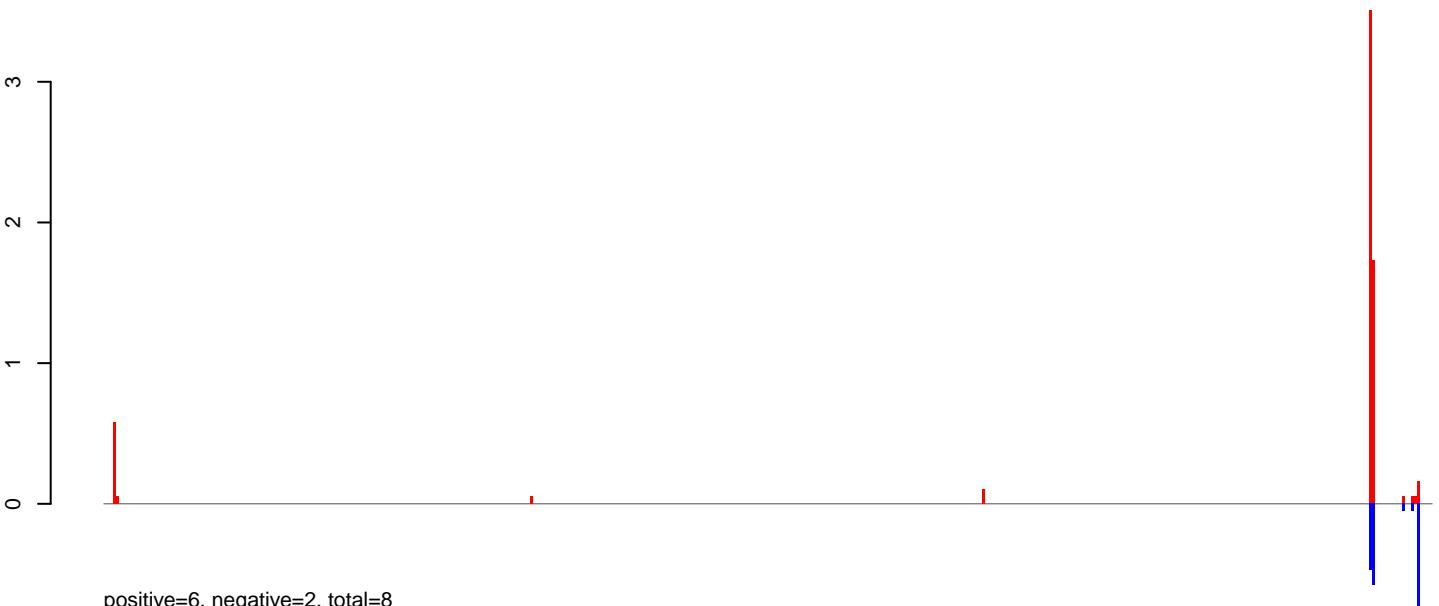
CuQuin_MG_7dpi3.24_35.rep



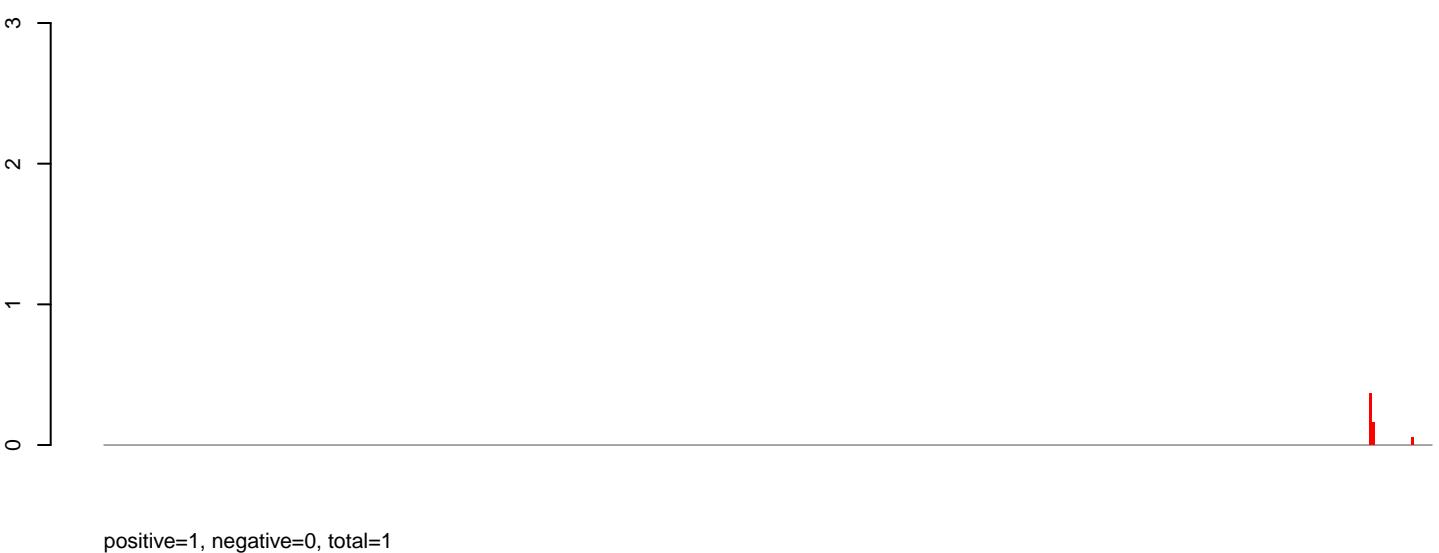
CuQuin_MG_7dpi3.rep



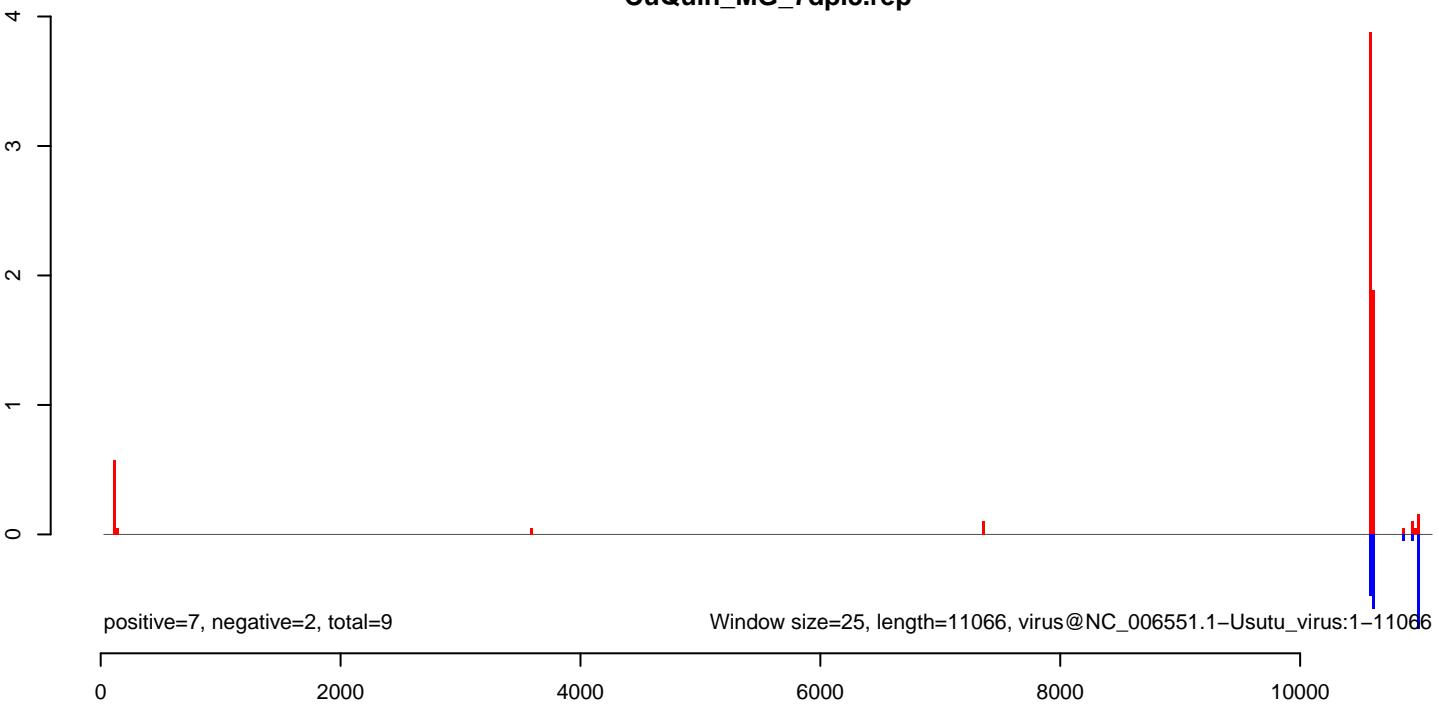
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



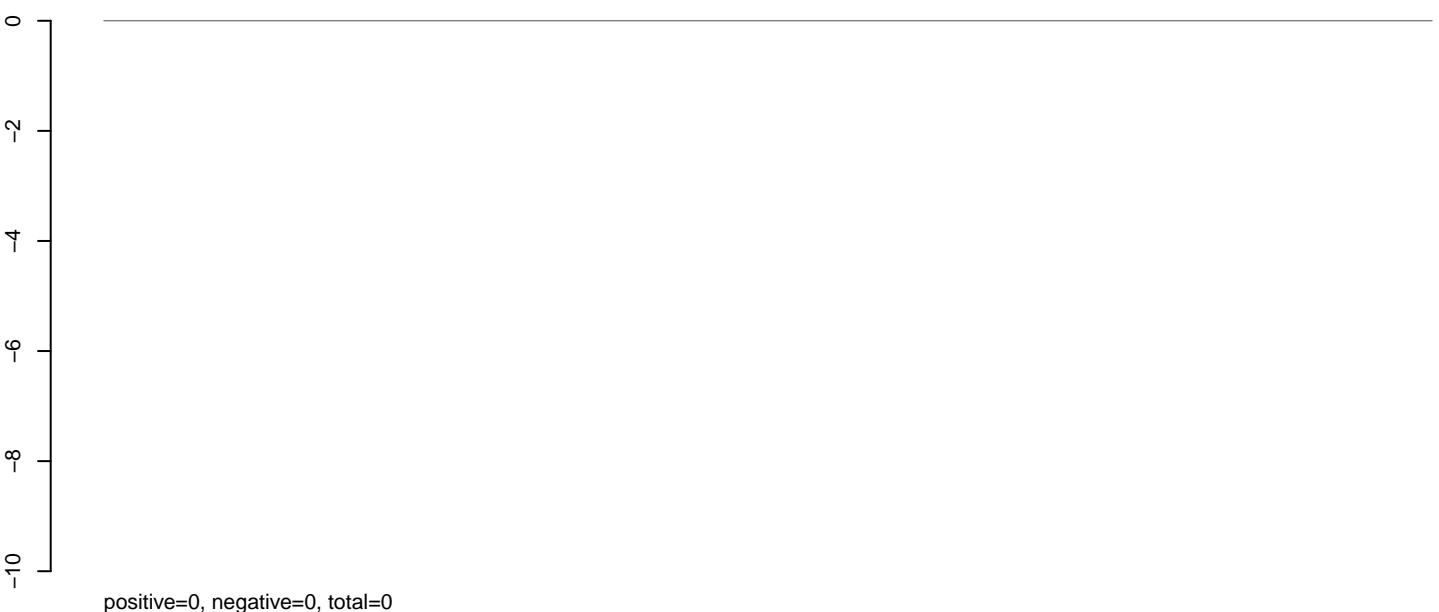
CuQuin_MG_7dpi3.rep



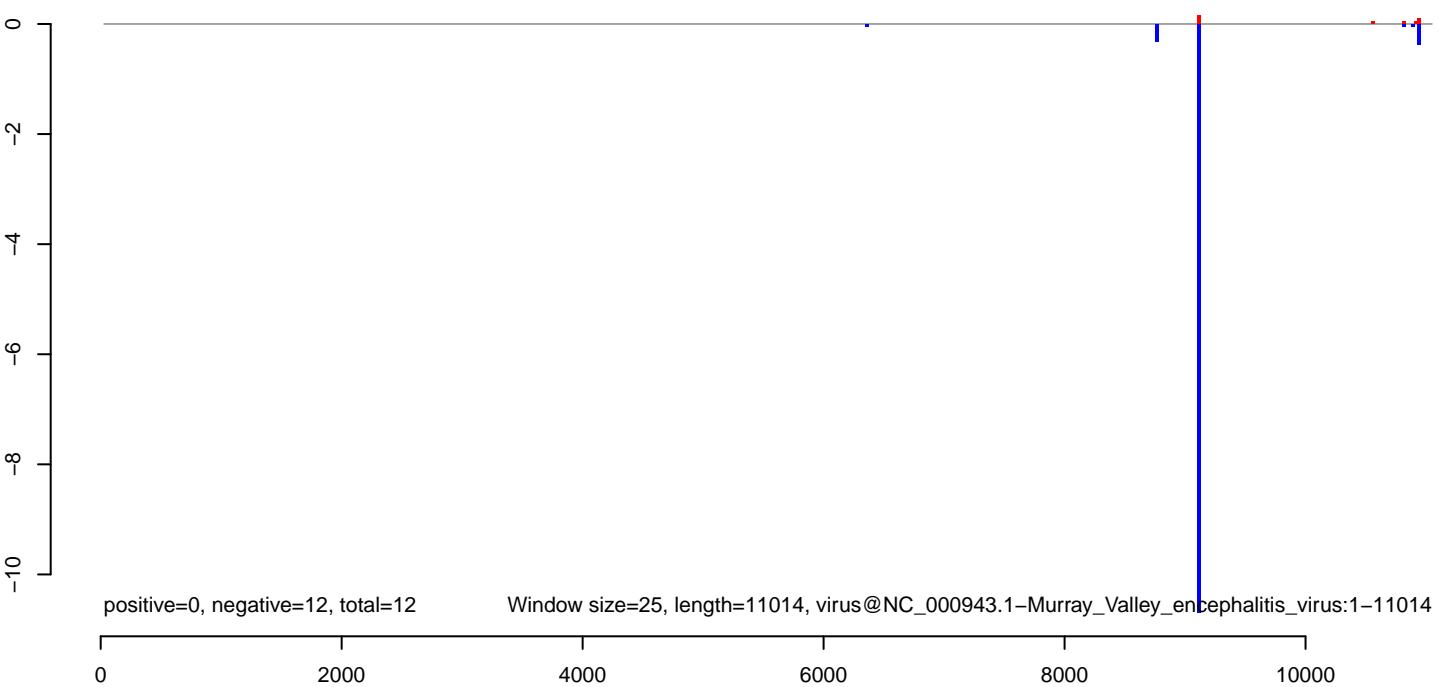
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep

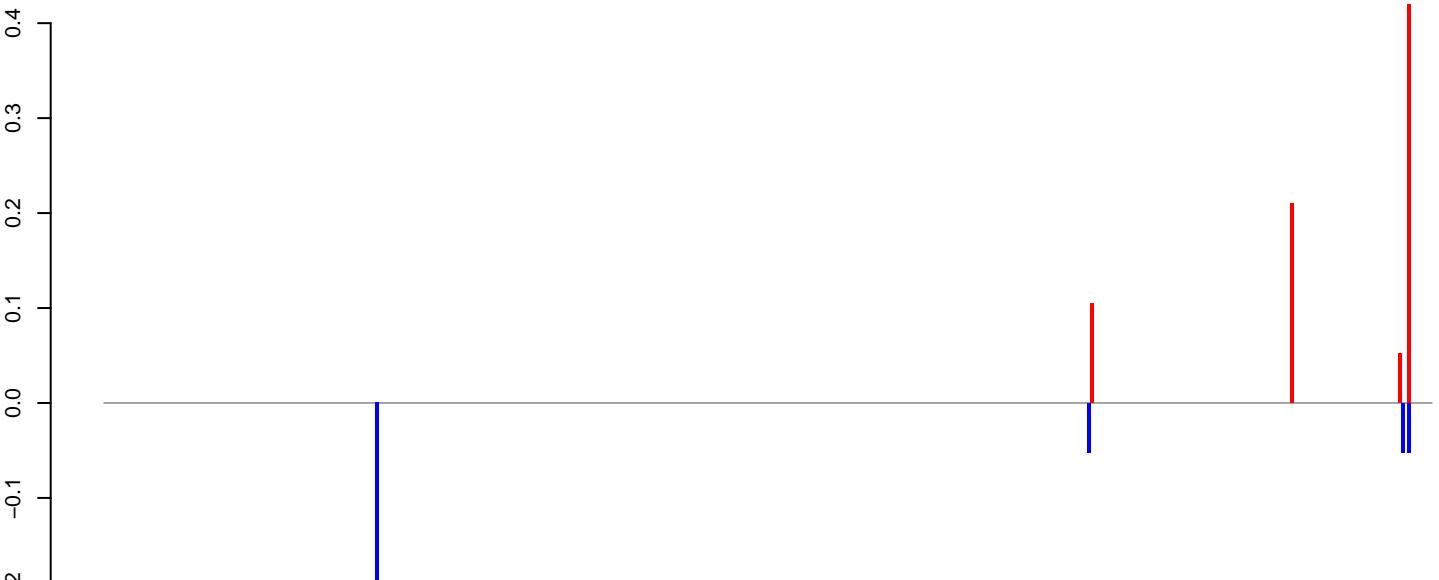


CuQuin_MG_7dpi3.rep



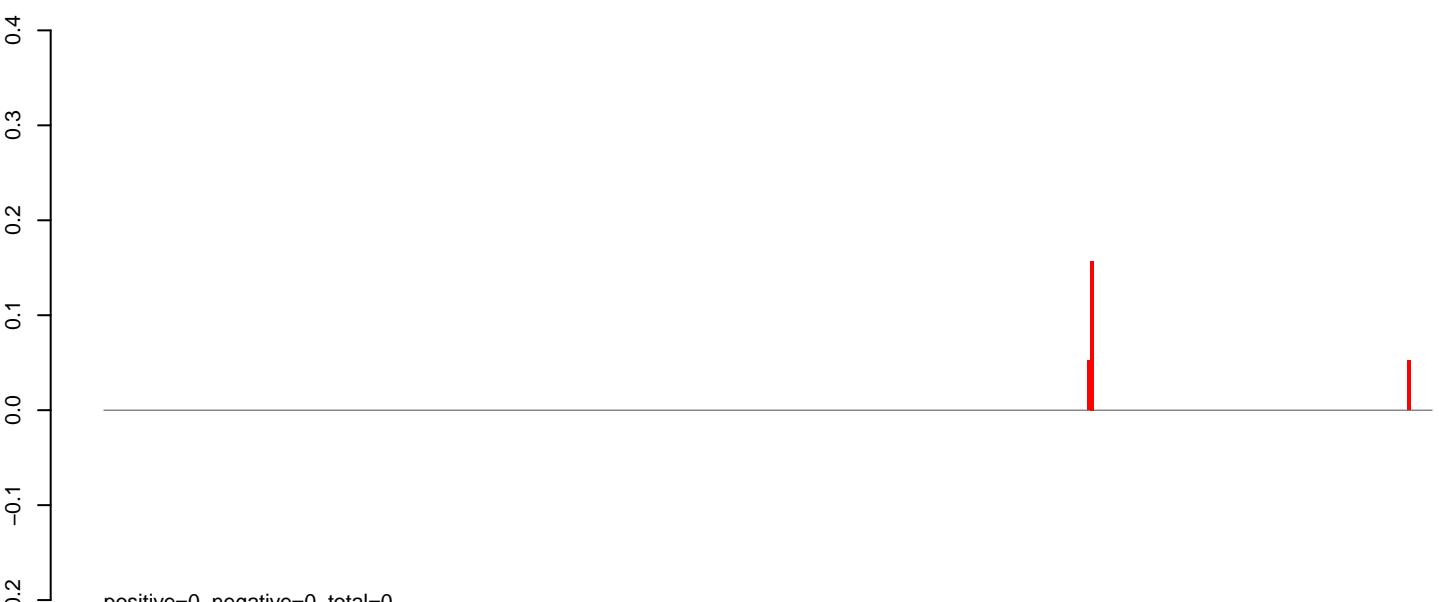
CuQuin_MG_7dpi3.18_23.rep

positive=1, negative=0, total=1



CuQuin_MG_7dpi3.24_35.rep

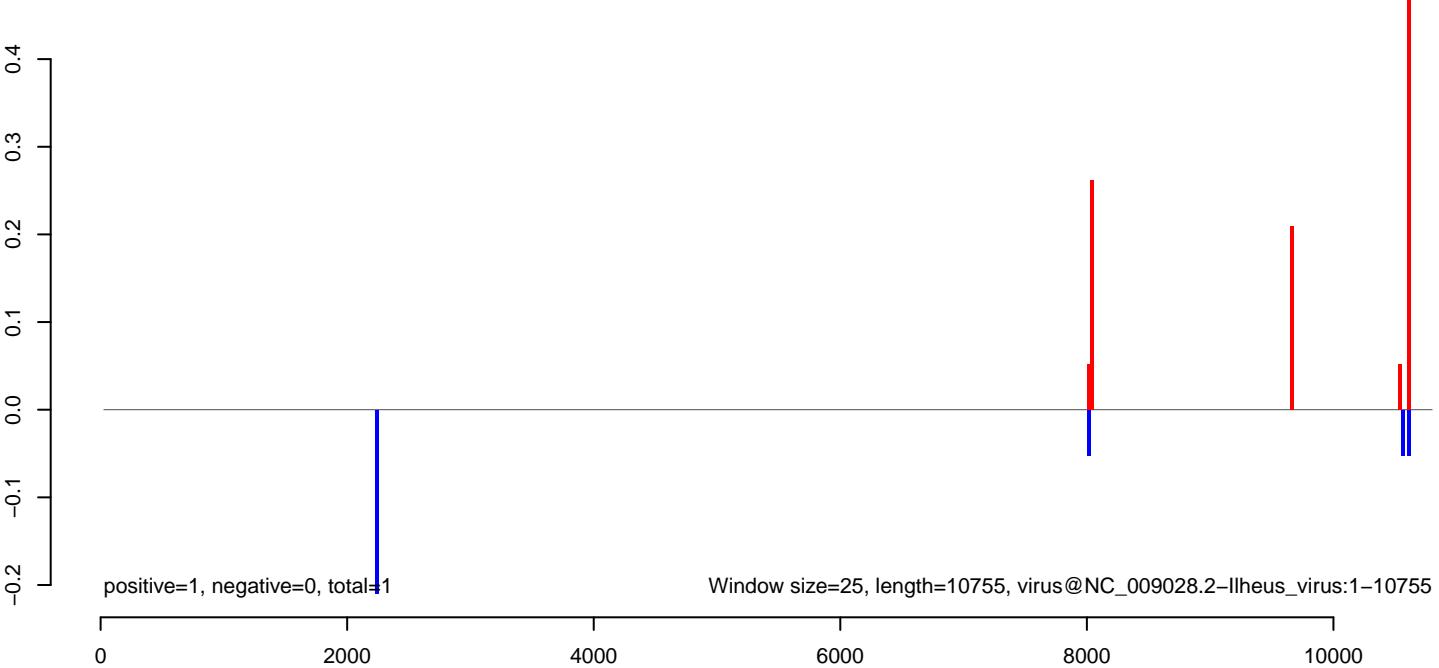
positive=0, negative=0, total=0



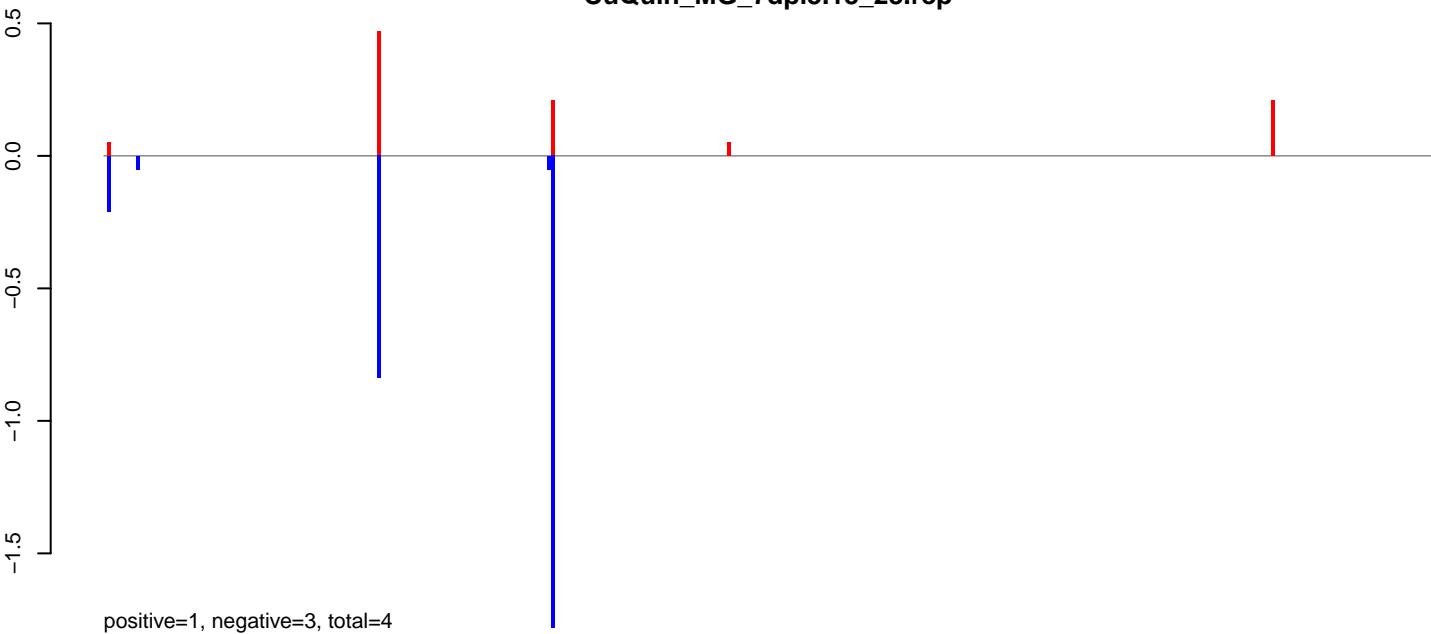
CuQuin_MG_7dpi3.rep

positive=1, negative=0, total=1

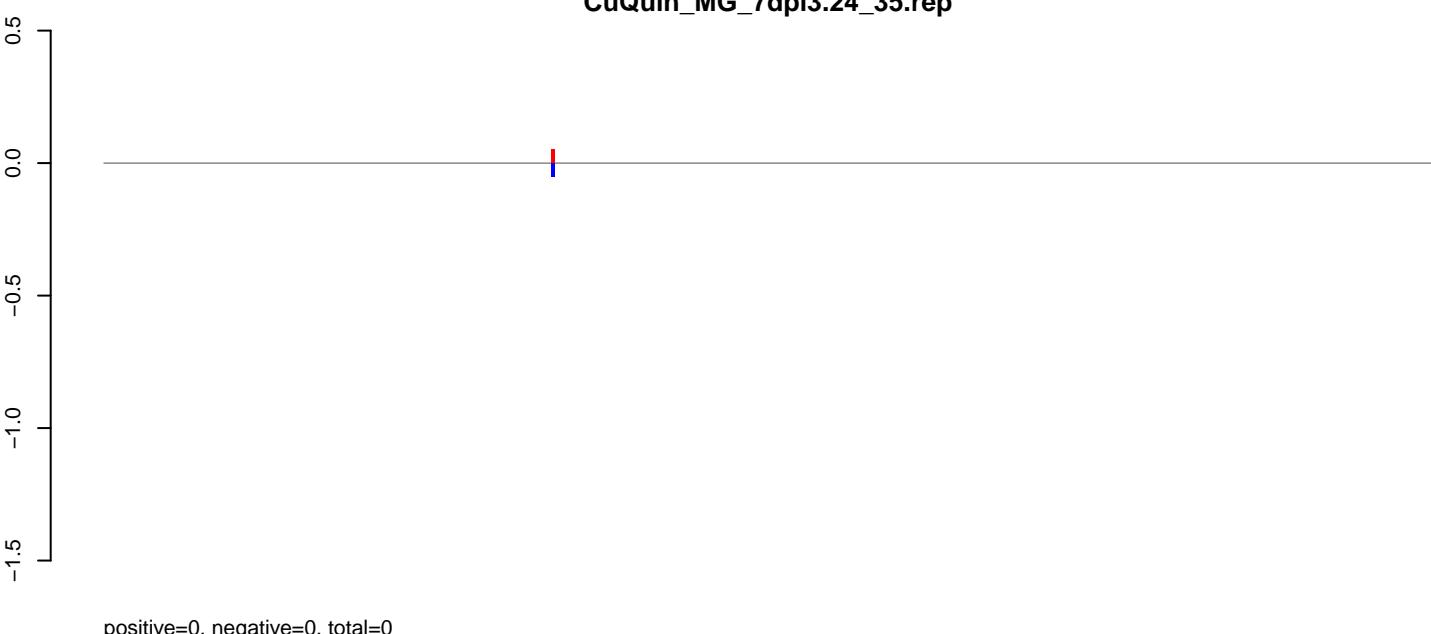
Window size=25, length=10755, virus@NC_009028.2-Ilheus_virus:1-10755



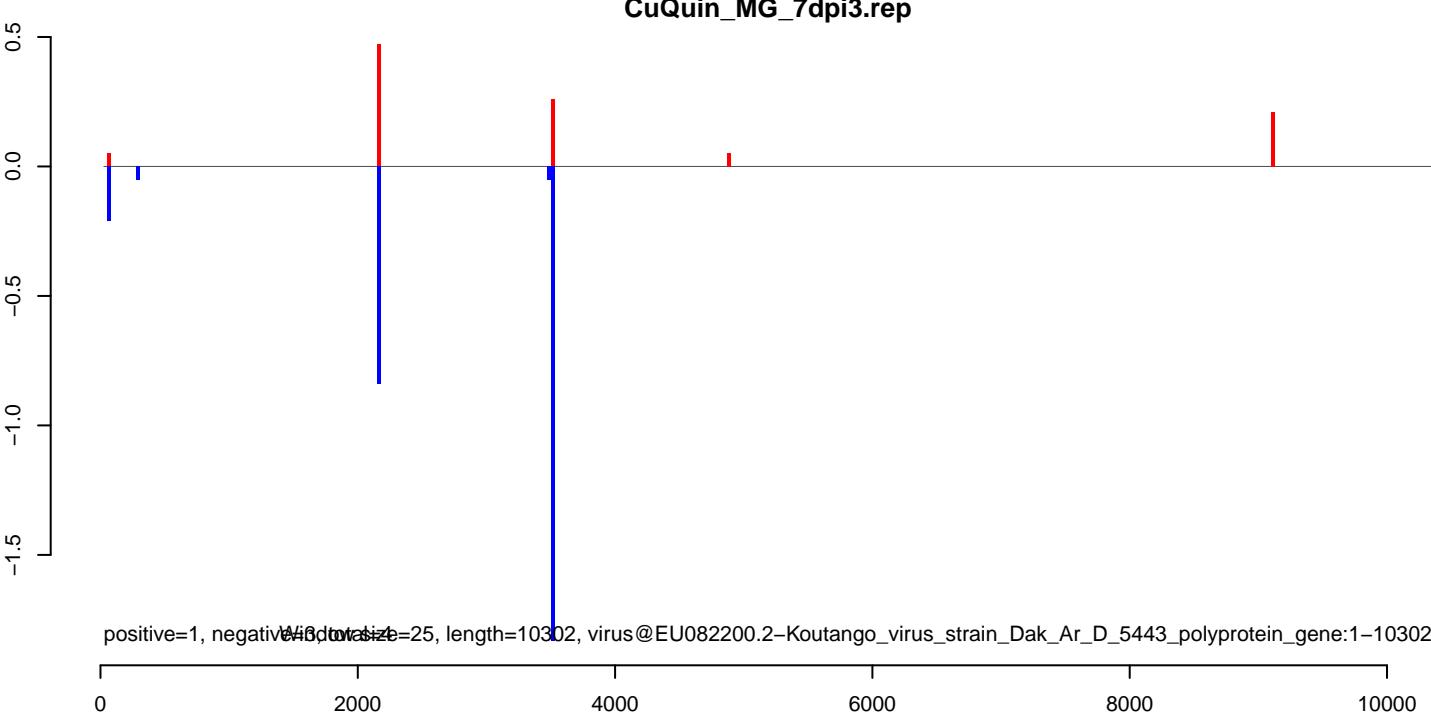
CuQuin_MG_7dpi3.18_23.rep



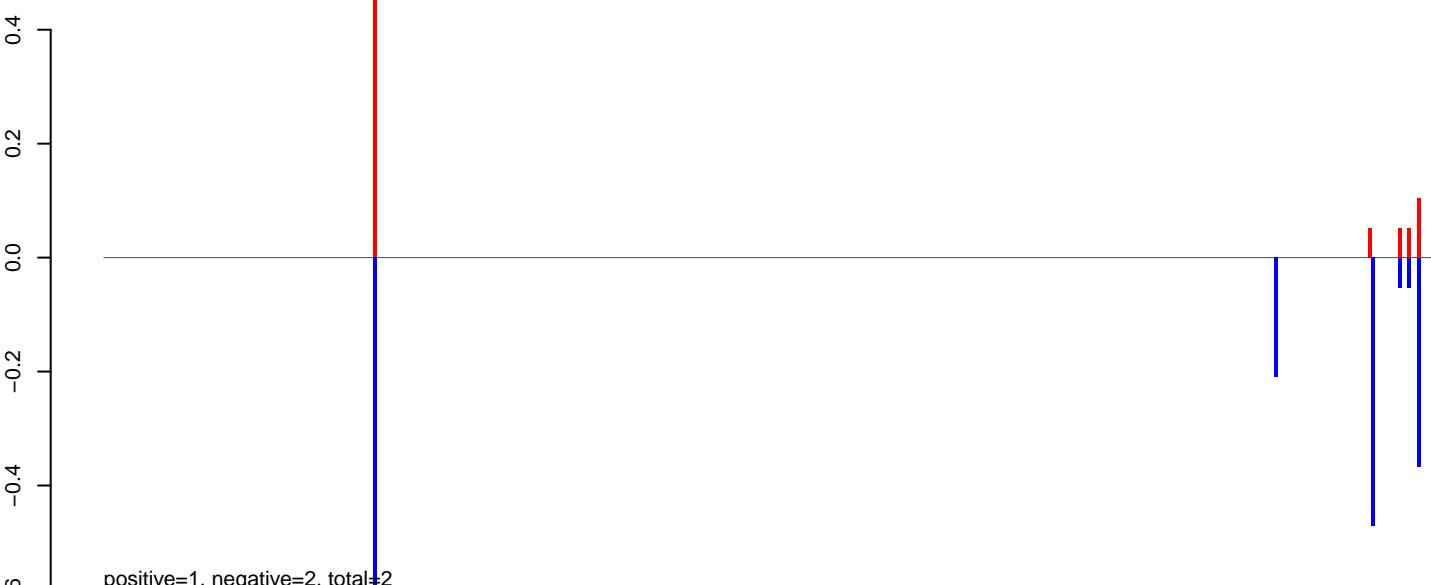
CuQuin_MG_7dpi3.24_35.rep



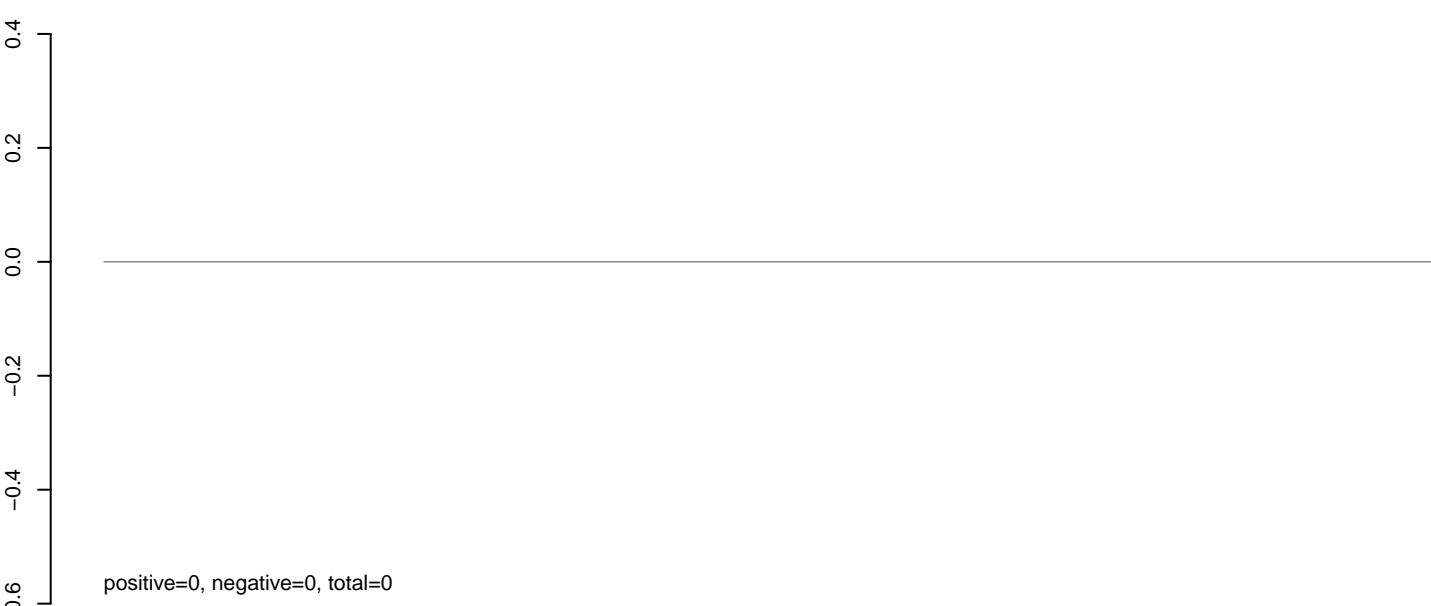
CuQuin_MG_7dpi3.rep



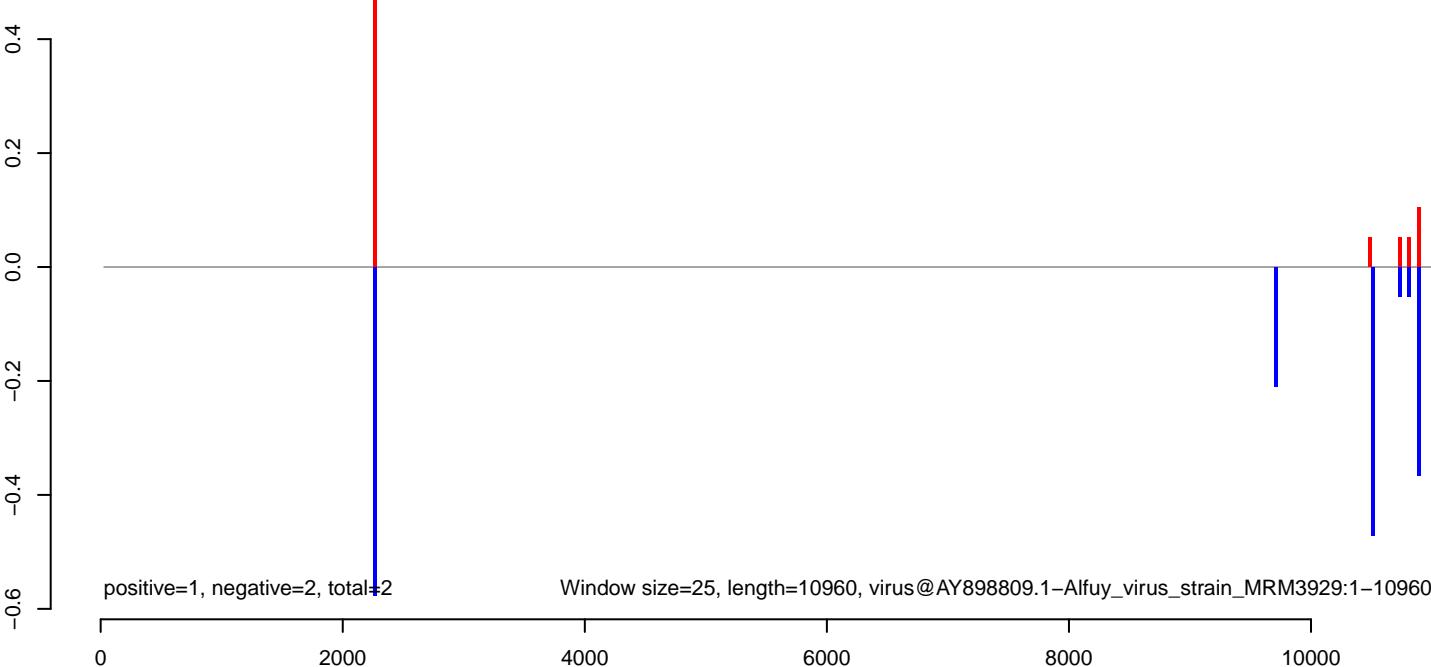
CuQuin_MG_7dpi3.18_23.rep



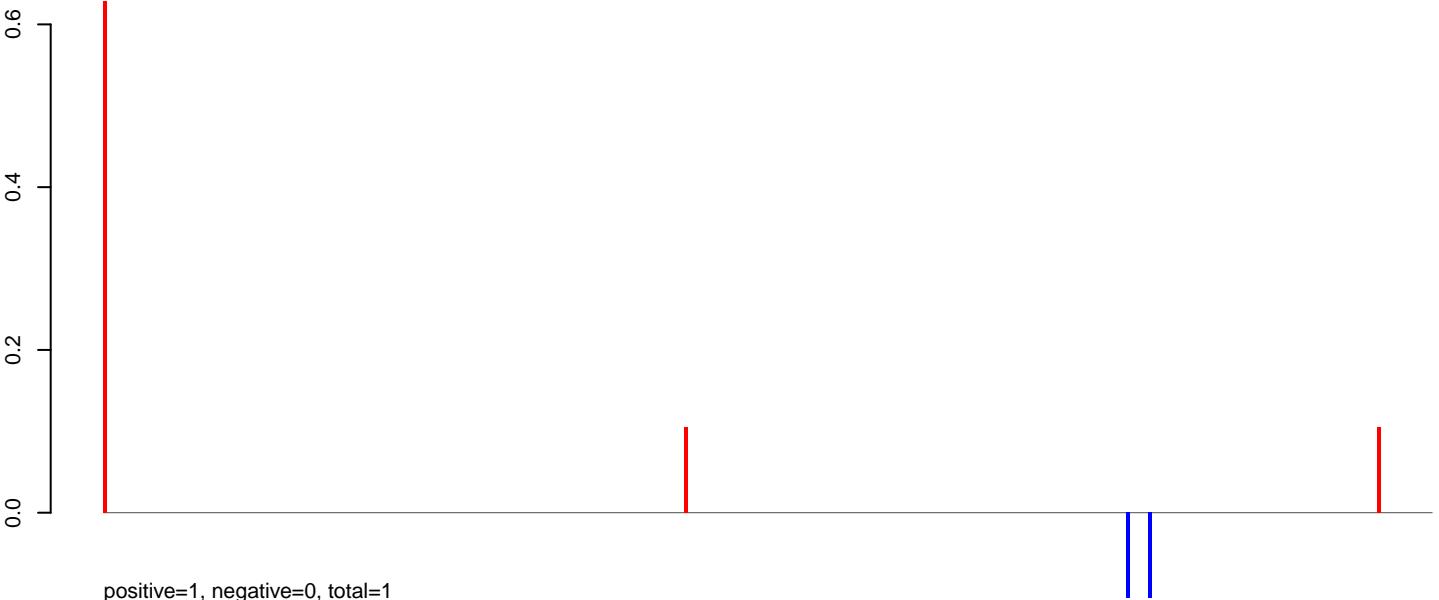
CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



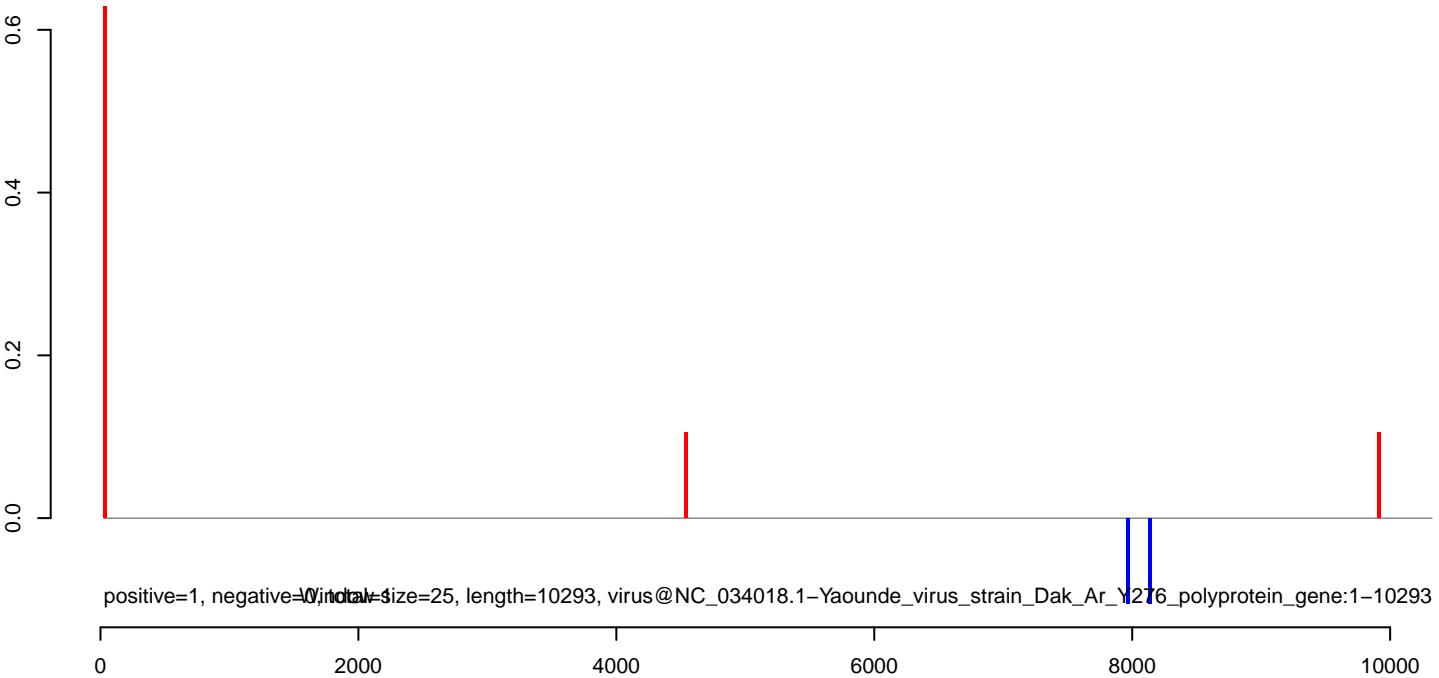
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep

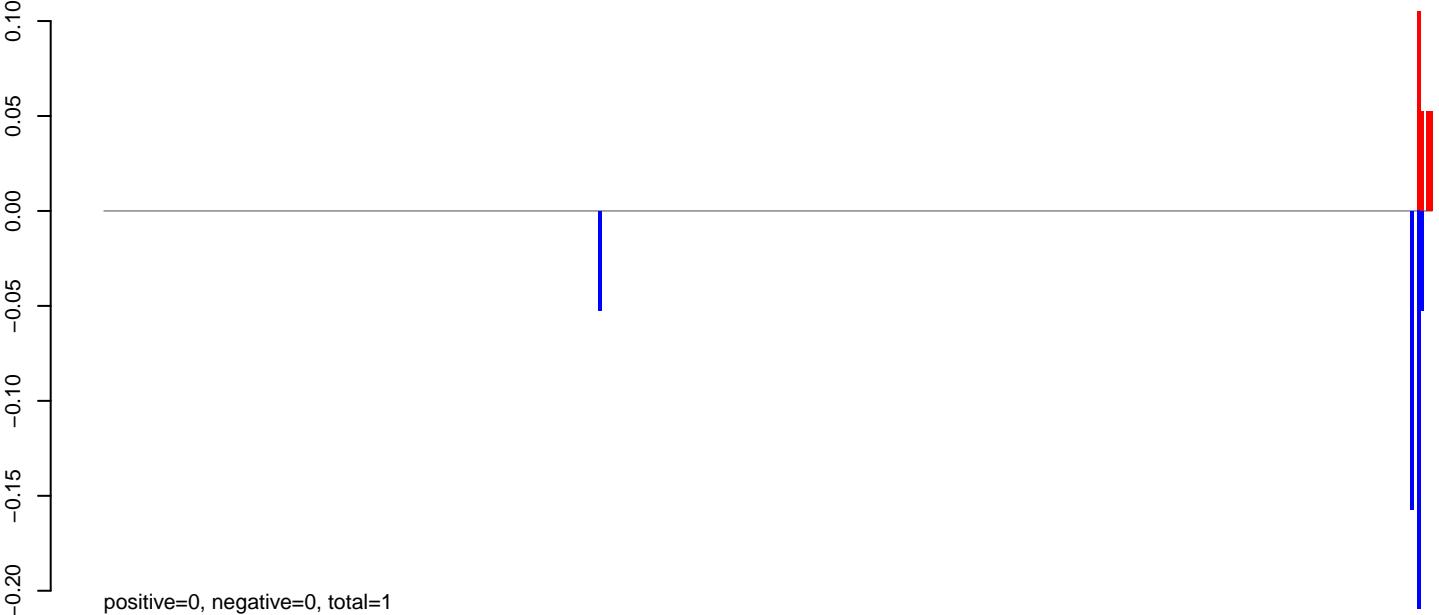


CuQuin_MG_7dpi3.rep



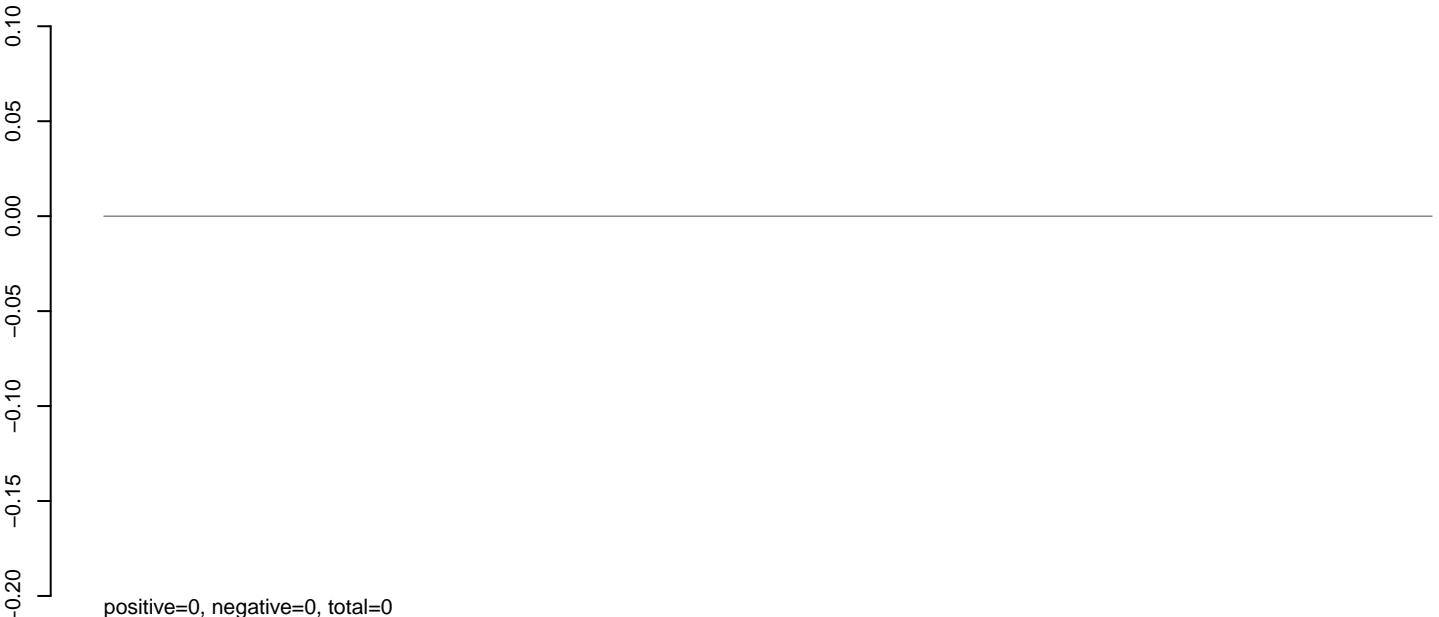
CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=1



CuQuin_MG_7dpi3.24_35.rep

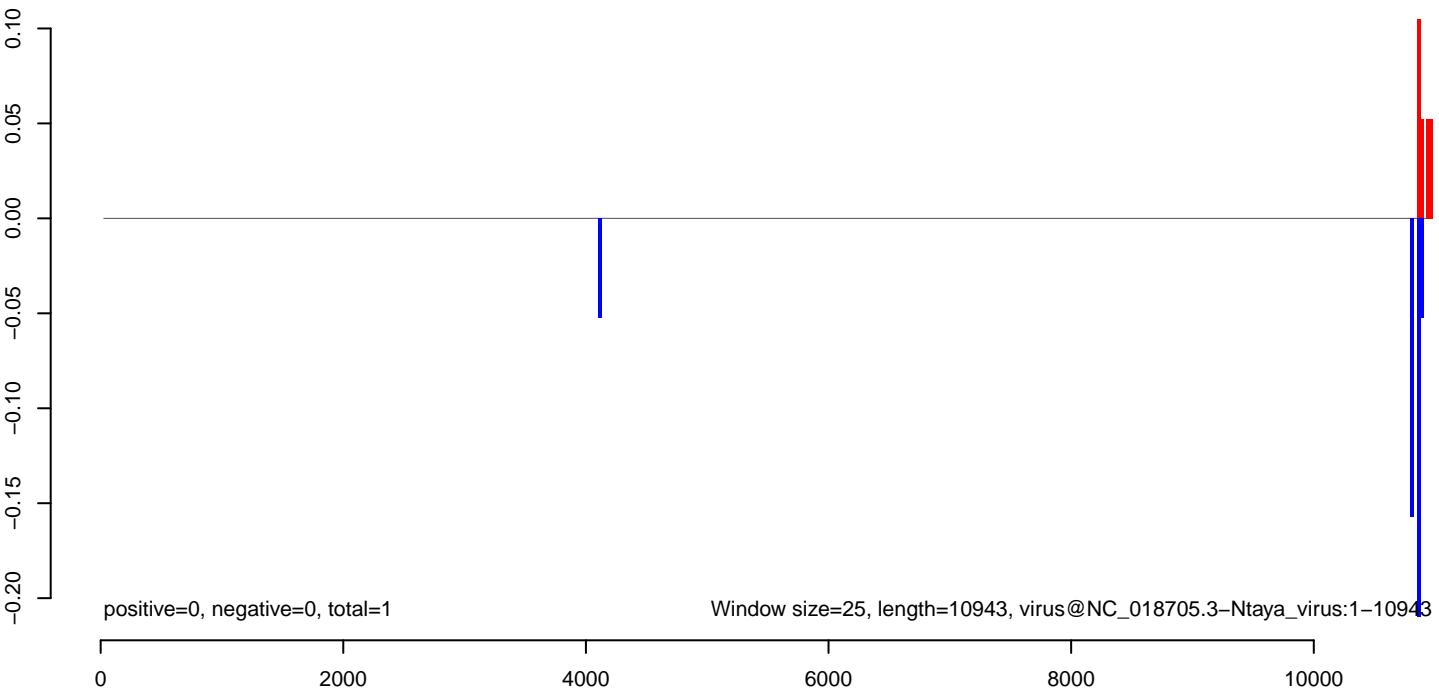
positive=0, negative=0, total=0



CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=1

Window size=25, length=10943, virus@NC_018705.3–Ntaya_virus:1–10943



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=2, total=3

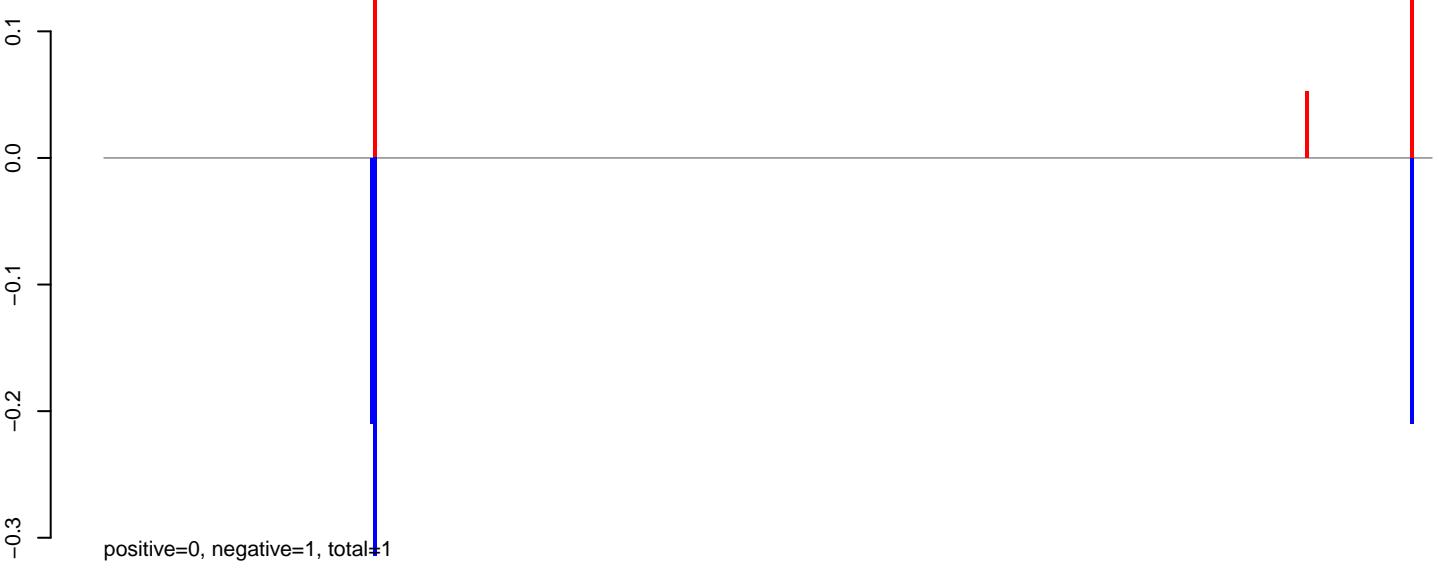
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

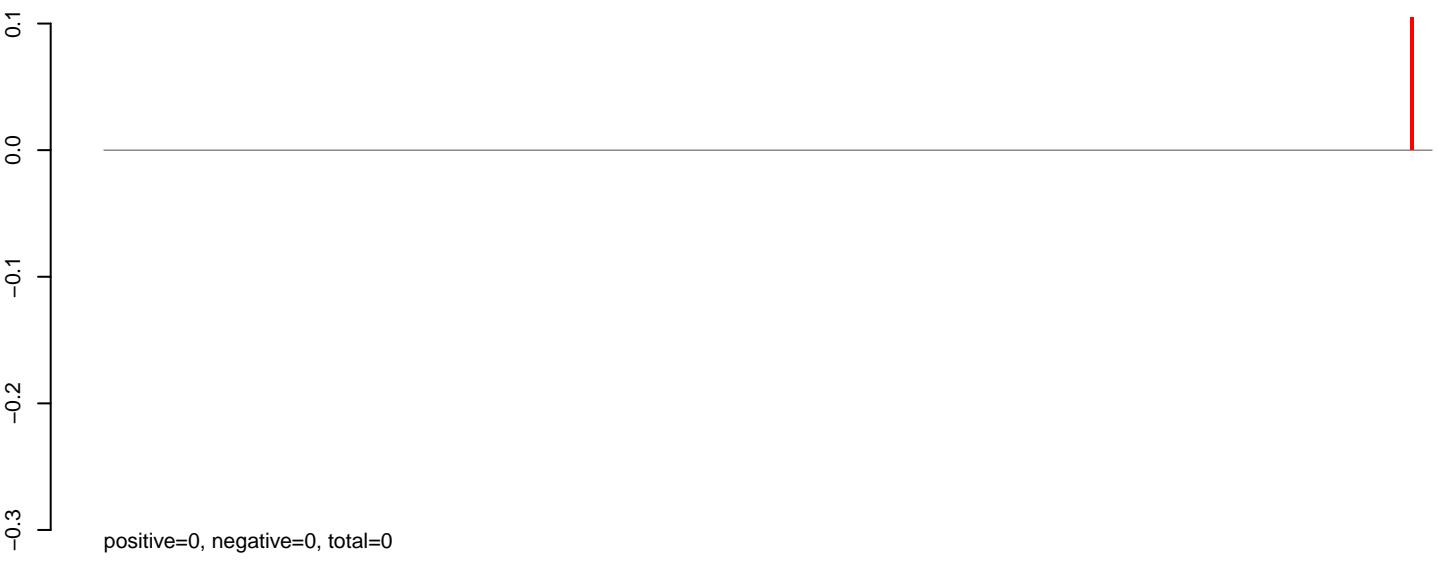
CuQuin_MG_7dpi3.rep

positive=0, negative=25, total=10990, virus@NC_015843.2-Tembusu_virus_flavivirus_polyprotein_flavivirus_polyprotein_gene_gene:1-10990

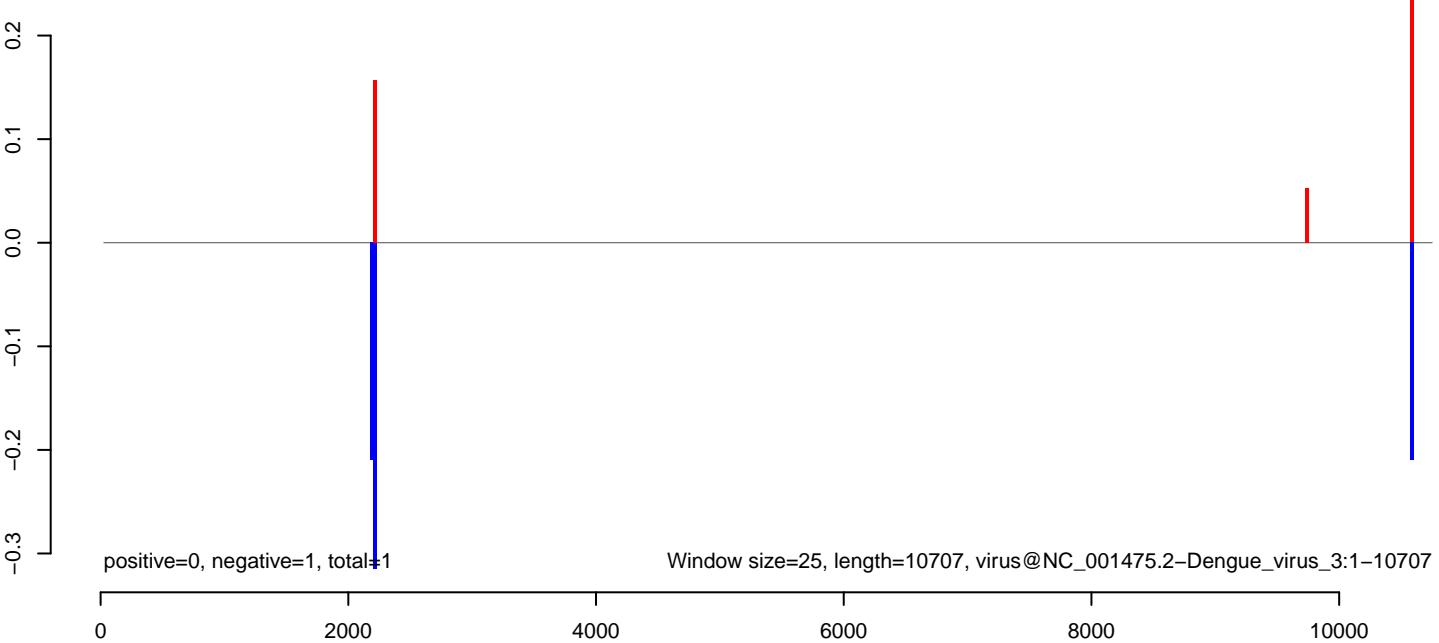
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



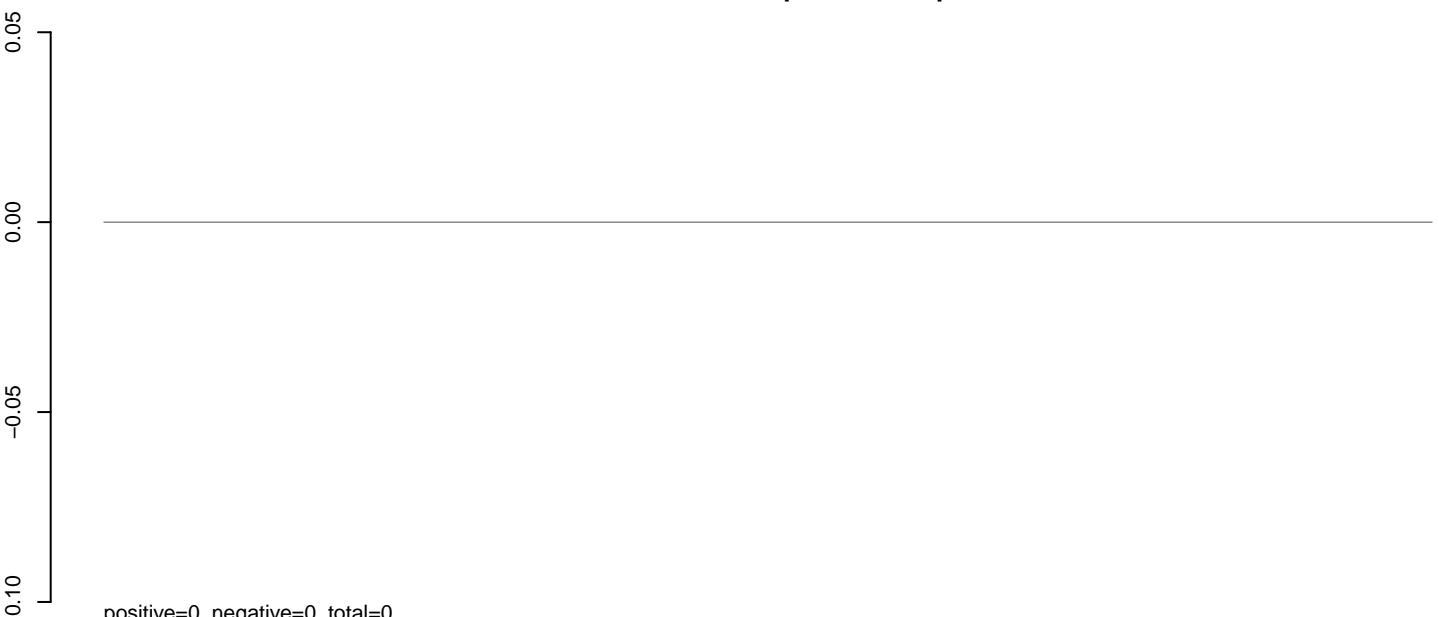
CuQuin_MG_7dpi3.rep



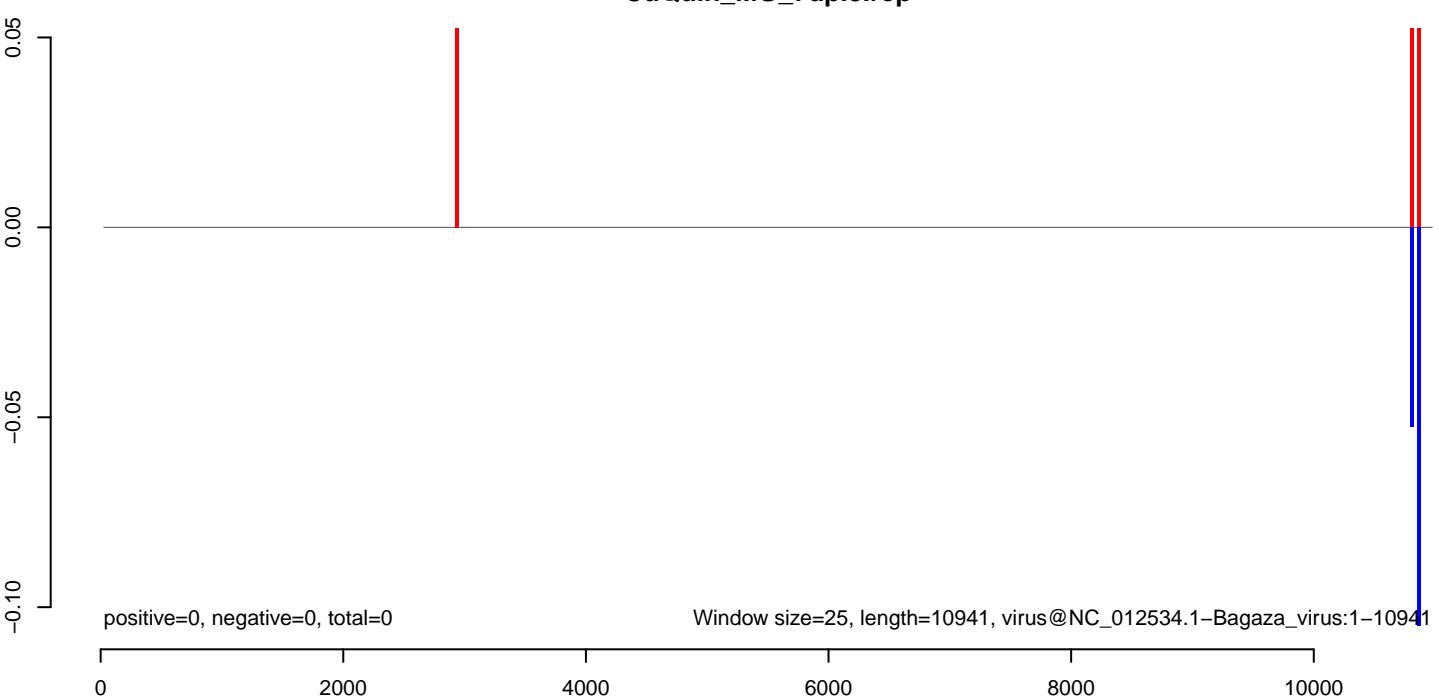
CuQuin_MG_7dpi3.18_23.rep



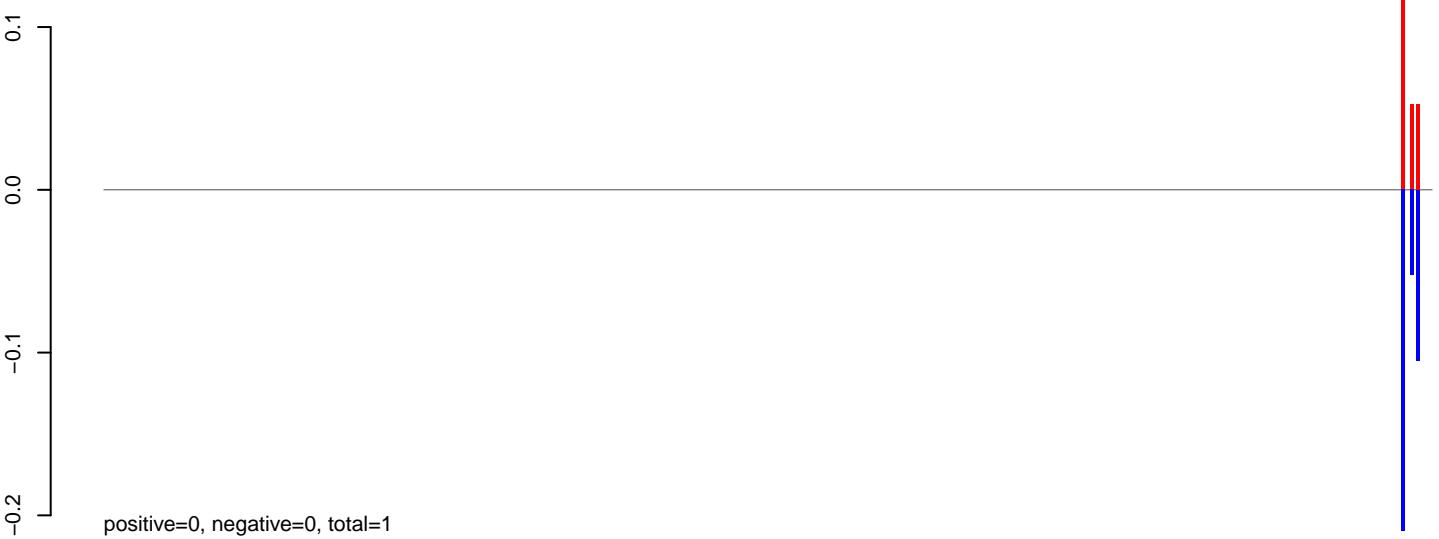
CuQuin_MG_7dpi3.24_35.rep



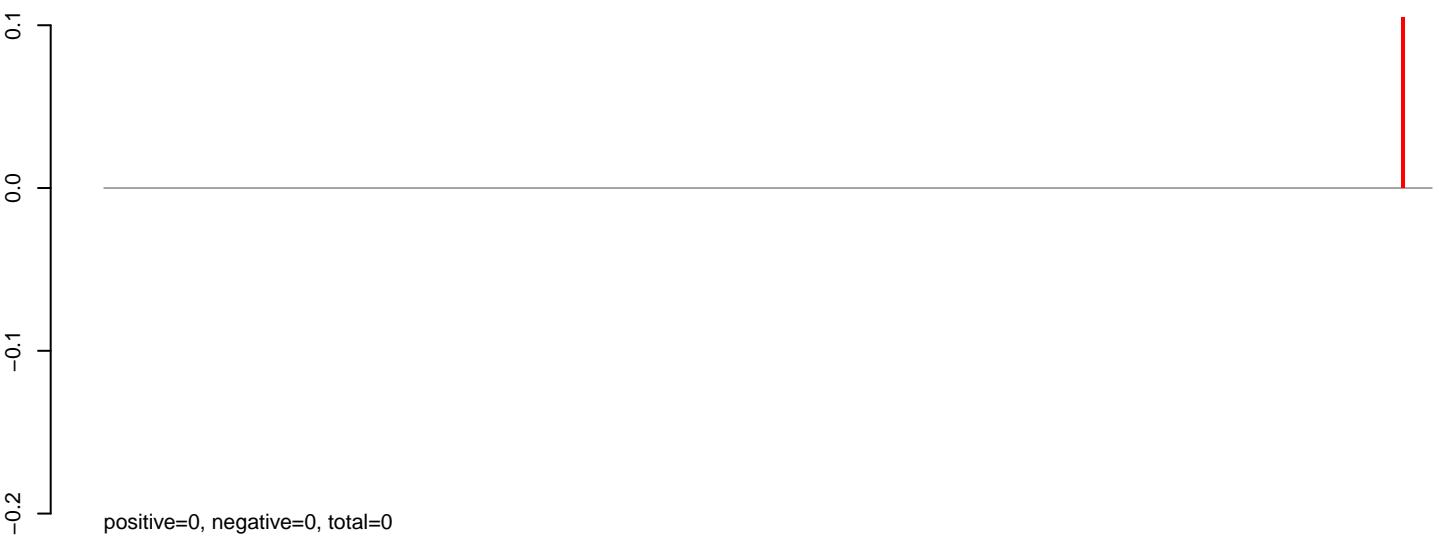
CuQuin_MG_7dpi3.rep



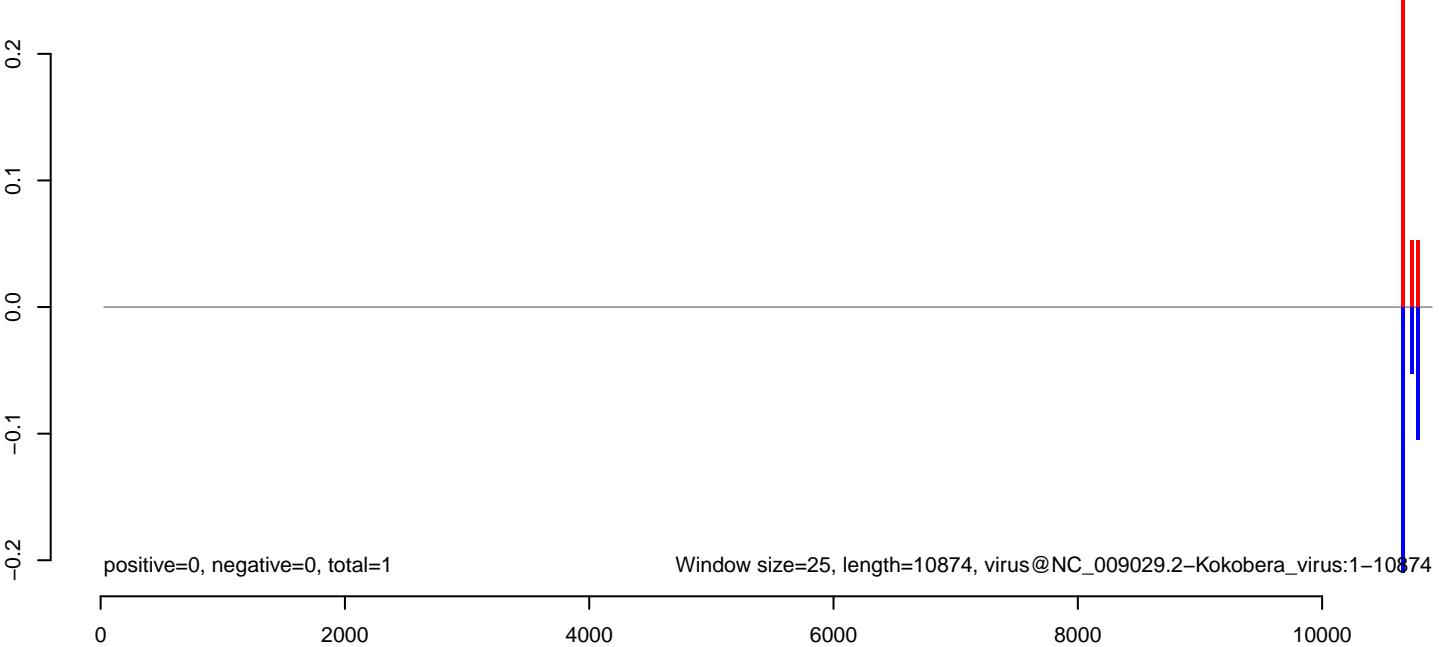
CuQuin_MG_7dpi3.18_23.rep



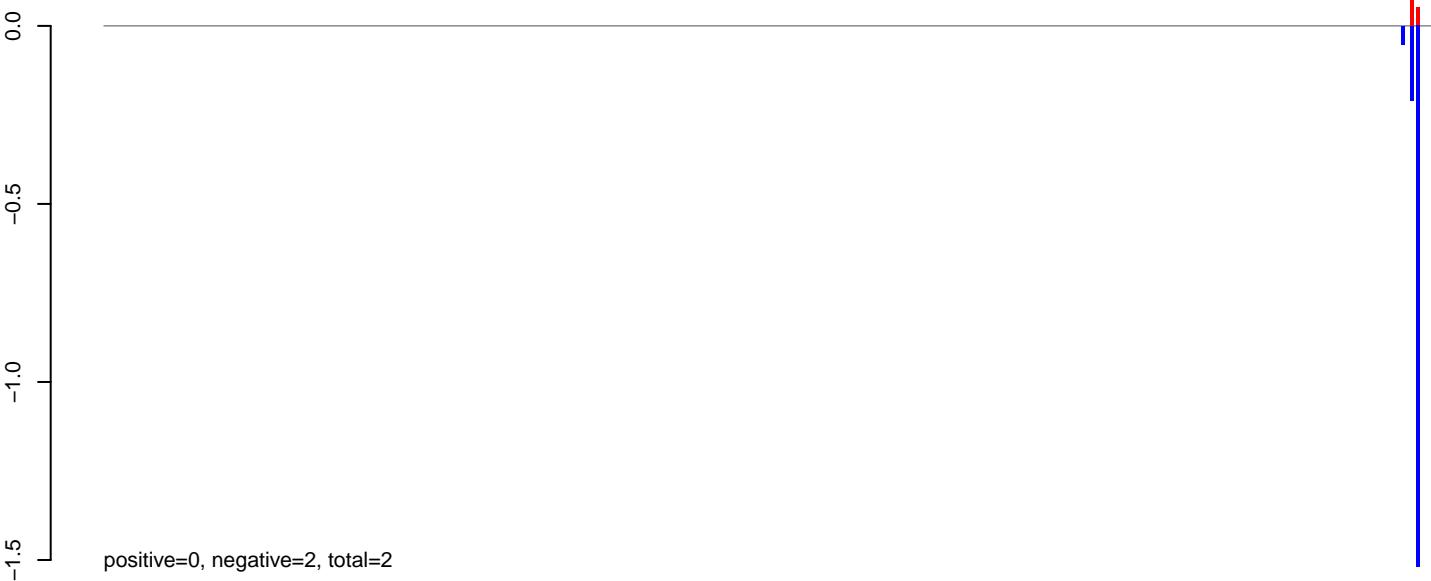
CuQuin_MG_7dpi3.24_35.rep



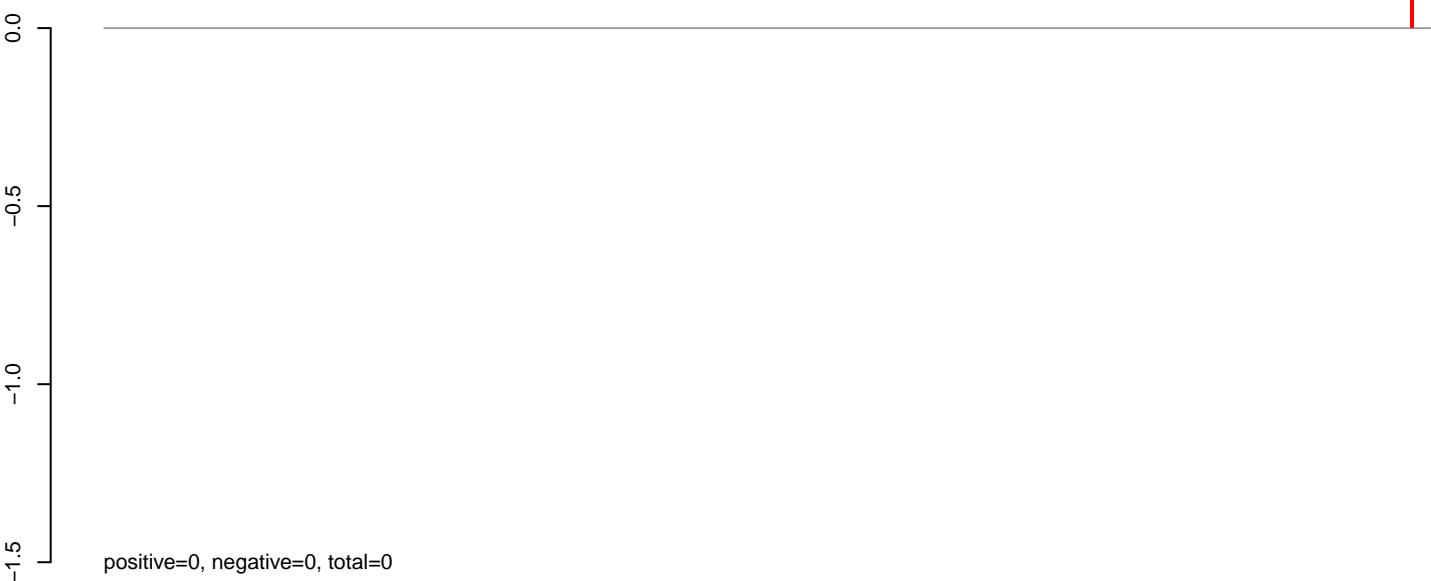
CuQuin_MG_7dpi3.rep



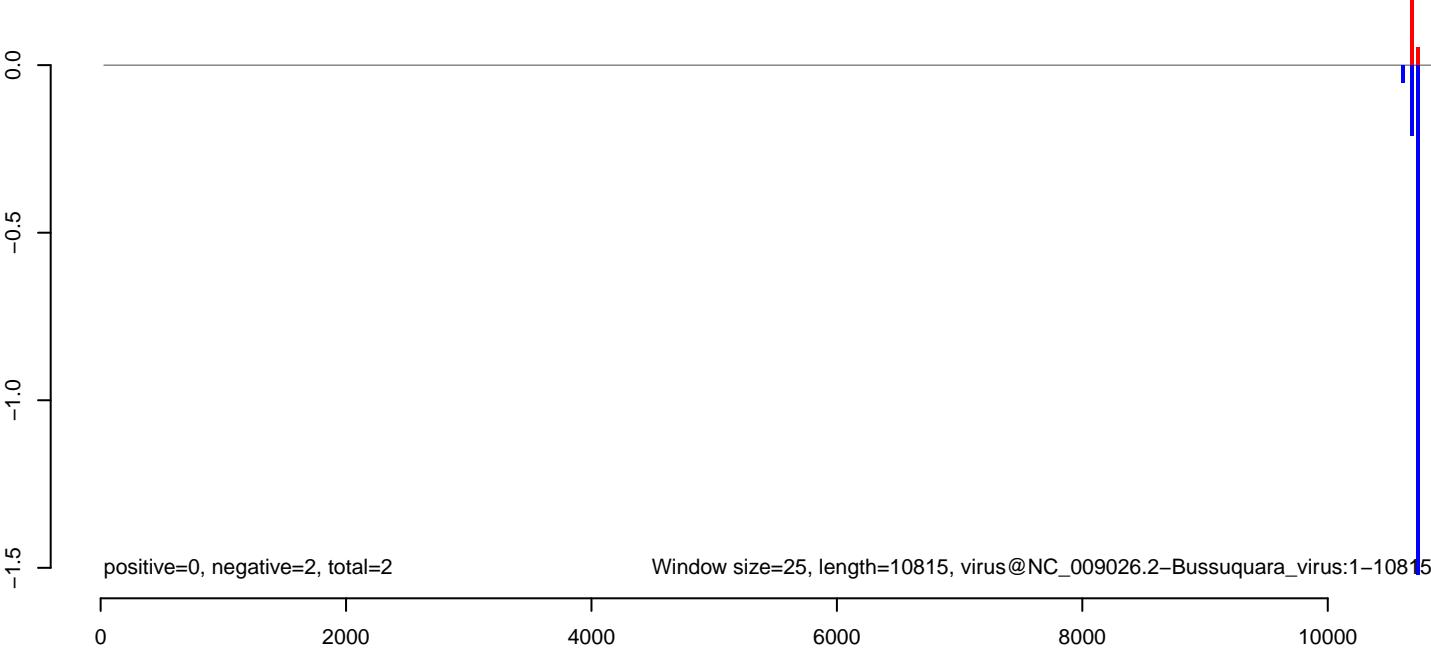
CuQuin_MG_7dpi3.18_23.rep



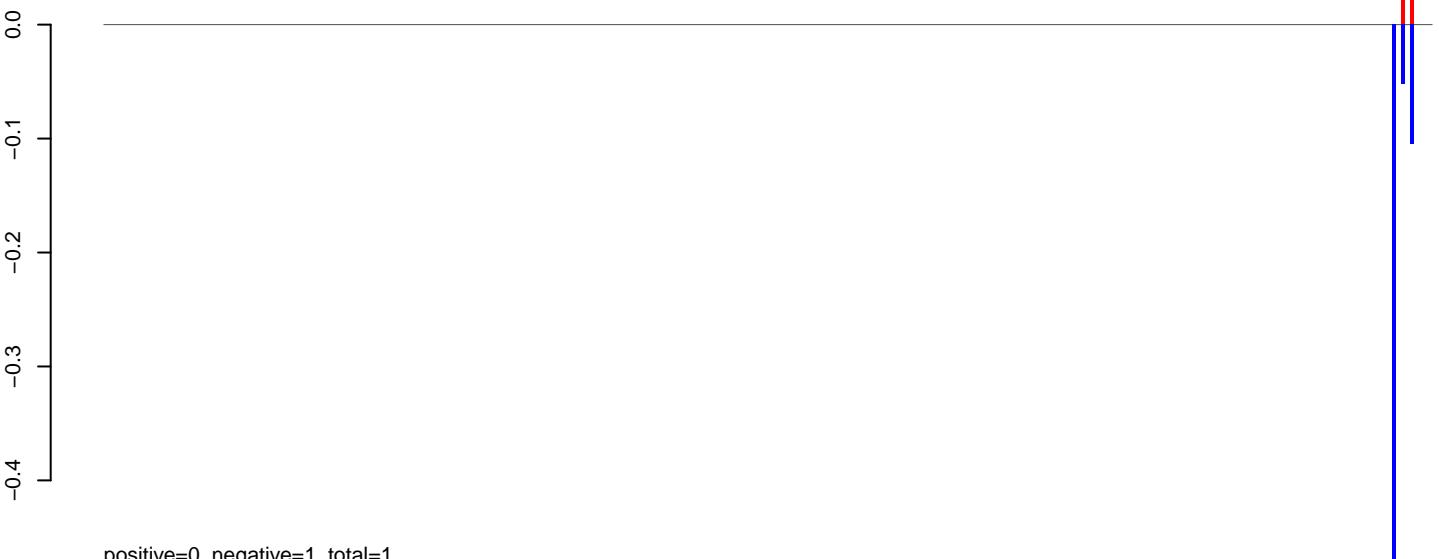
CuQuin_MG_7dpi3.24_35.rep



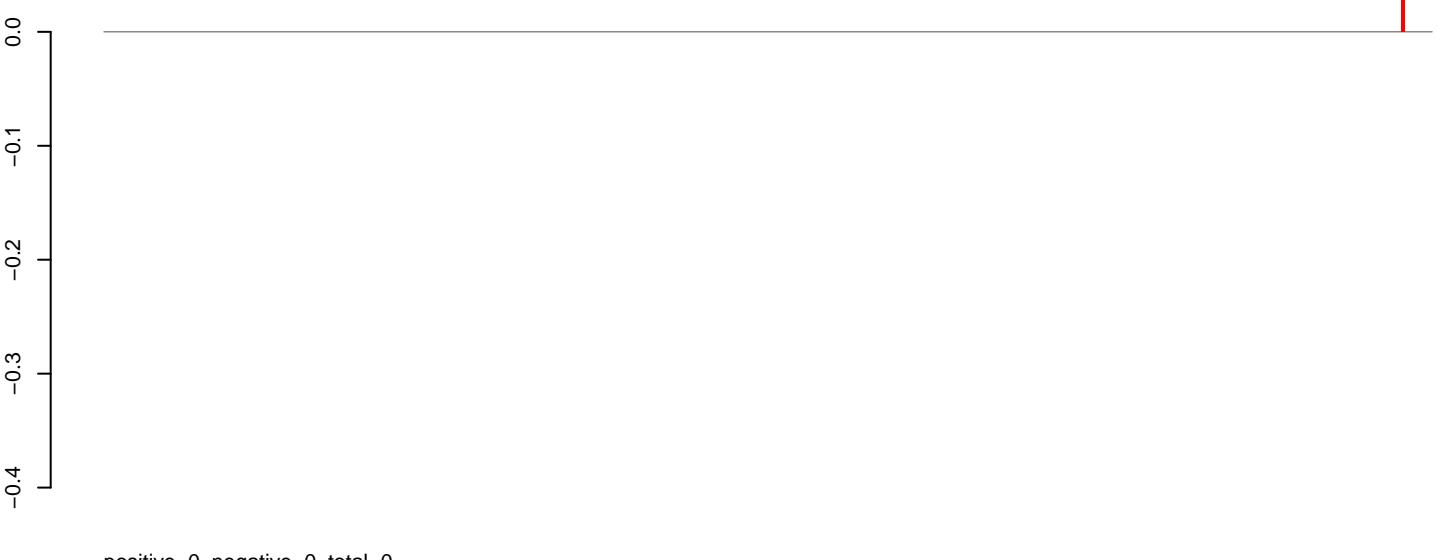
CuQuin_MG_7dpi3.rep



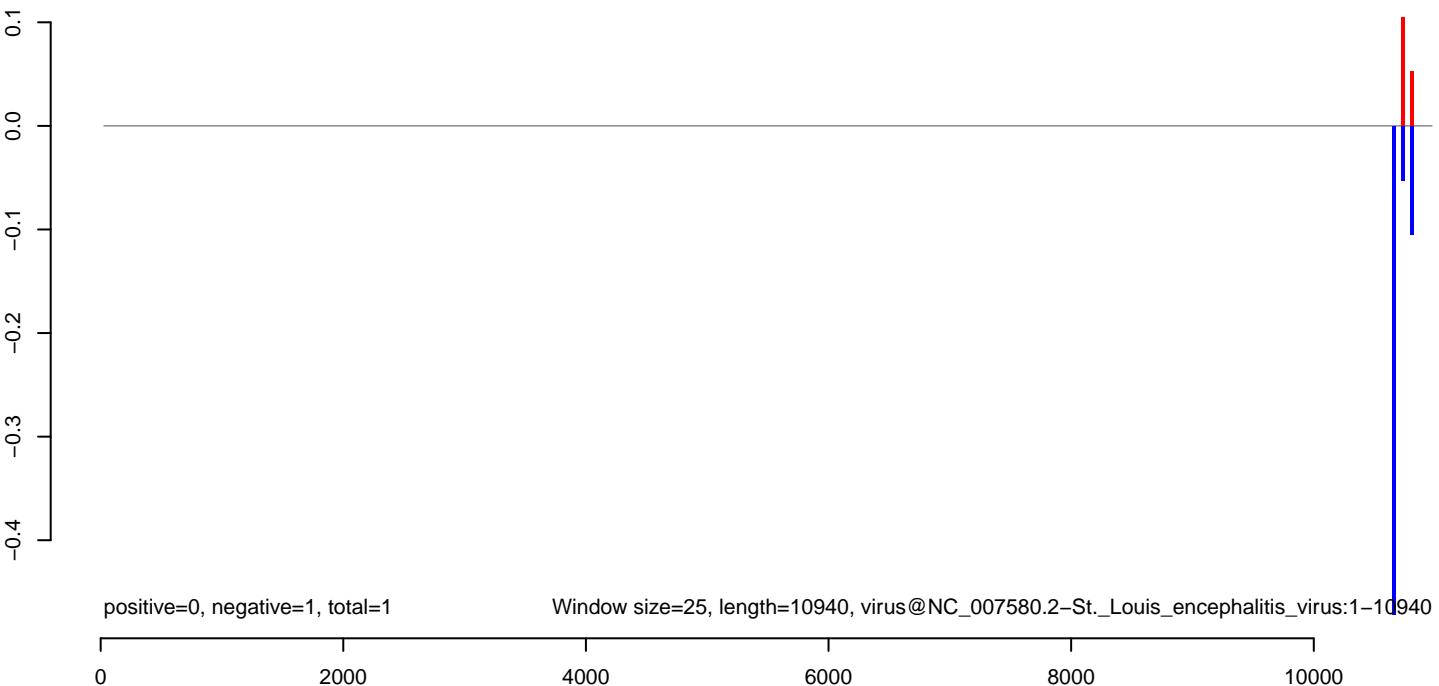
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

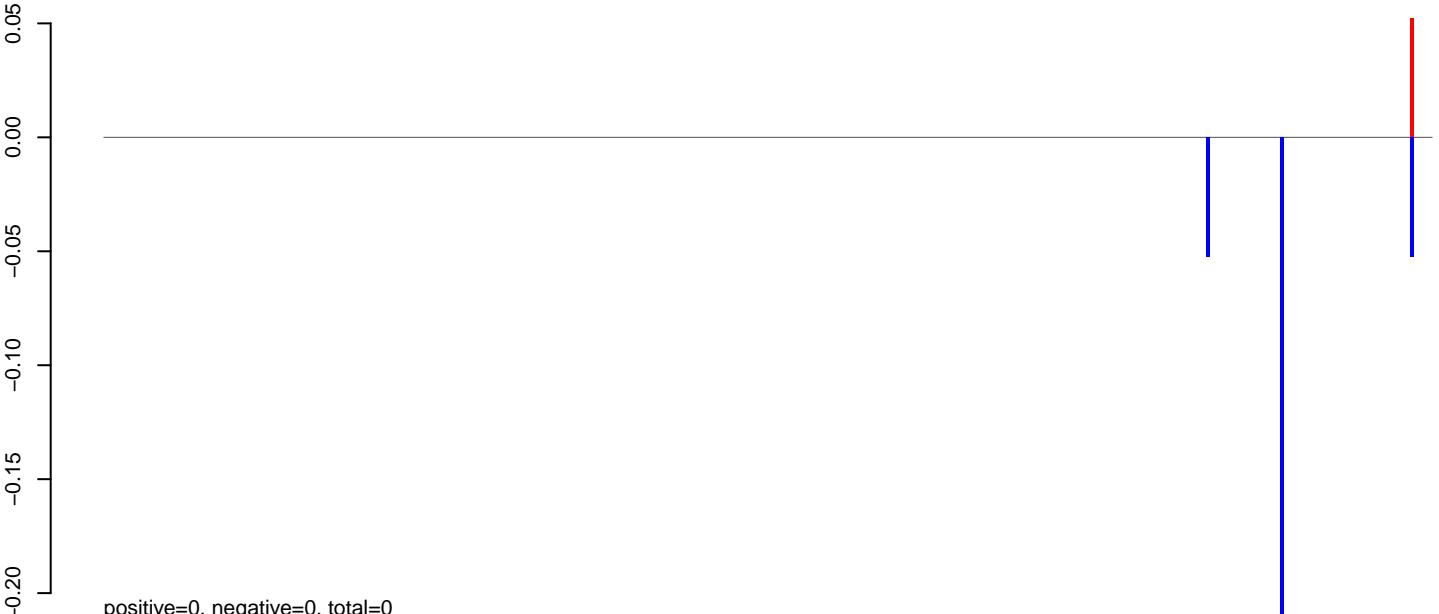
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

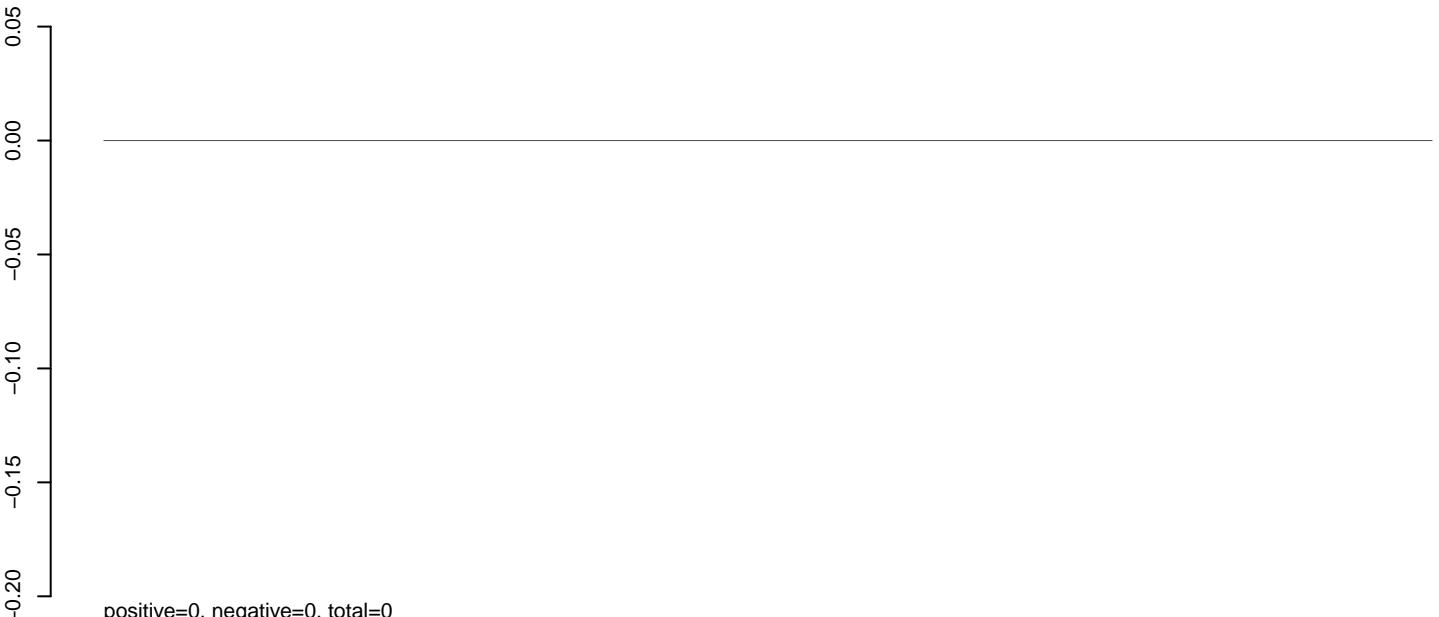
Window size=25, length=10649, virus@NC_002640.1 | Dengue_virus_4:1-10649

0 2000 4000 6000 8000 10000

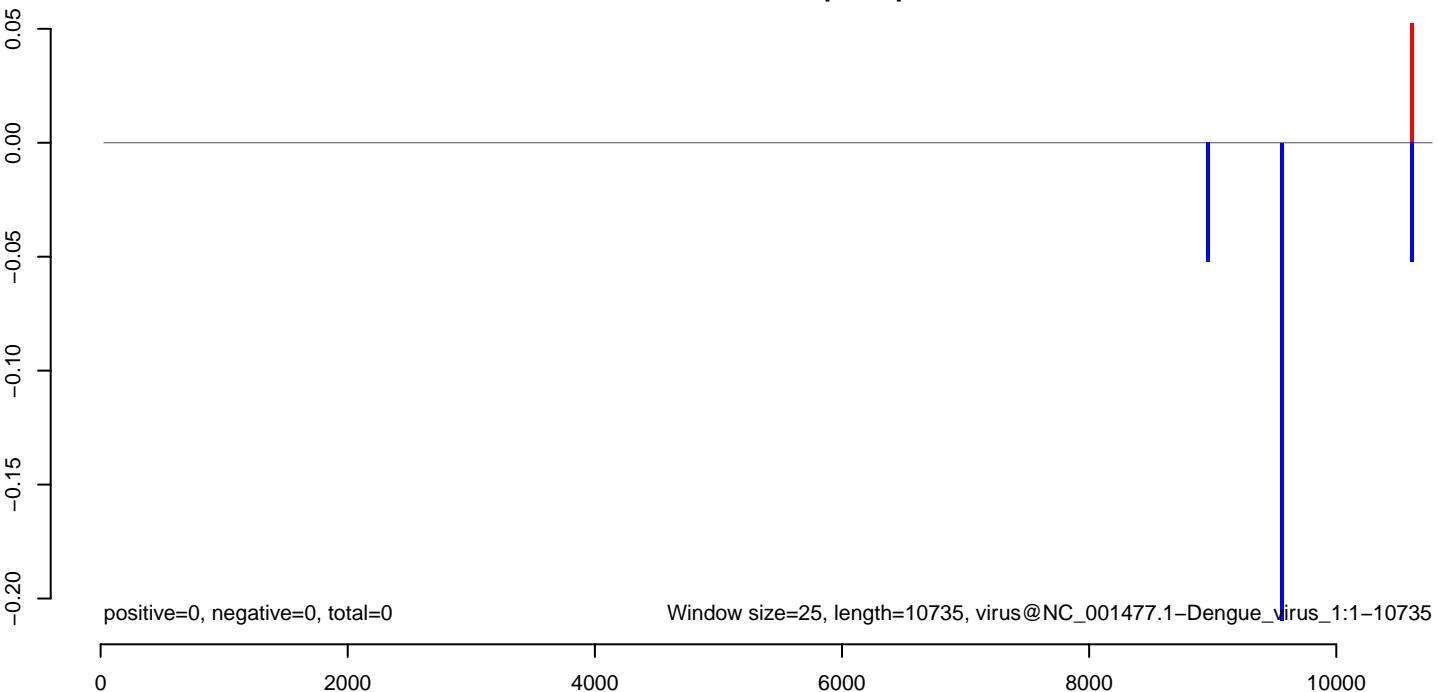
CuQuin_MG_7dpi3.18_23.rep



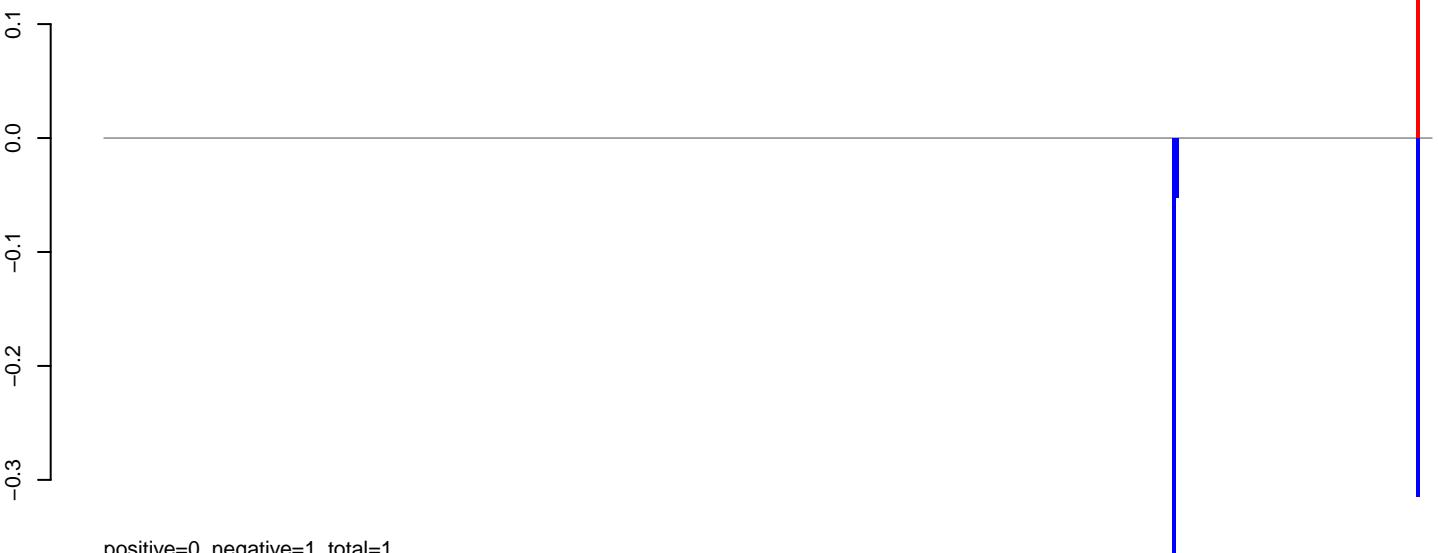
CuQuin_MG_7dpi3.24_35.rep



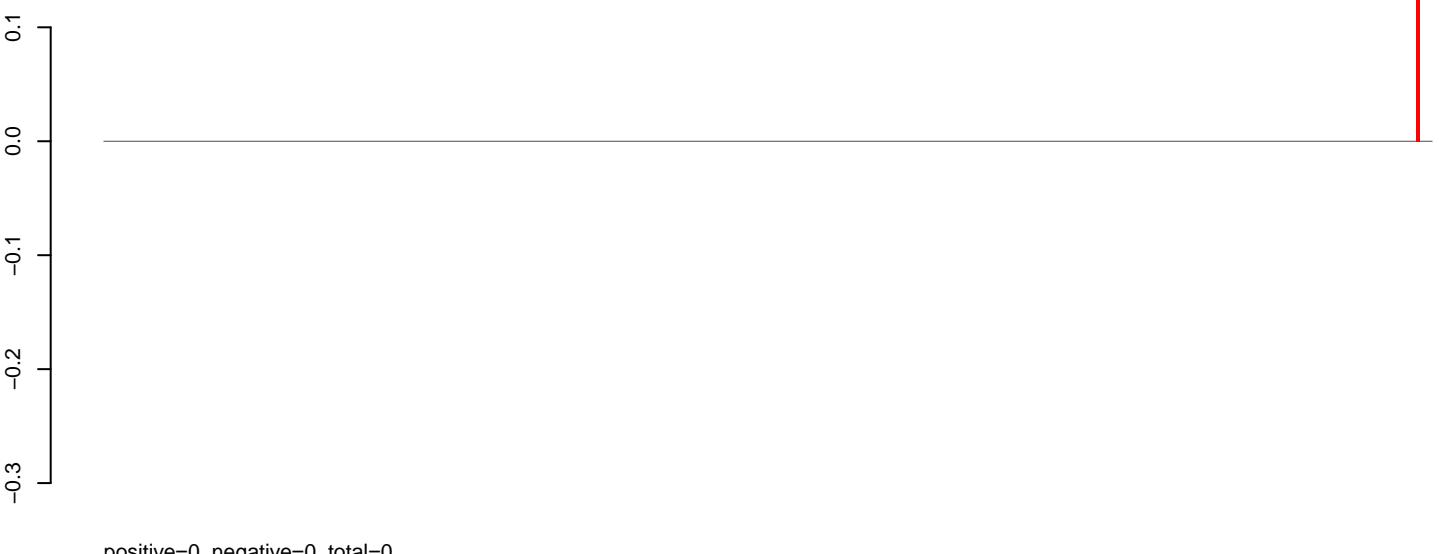
CuQuin_MG_7dpi3.rep



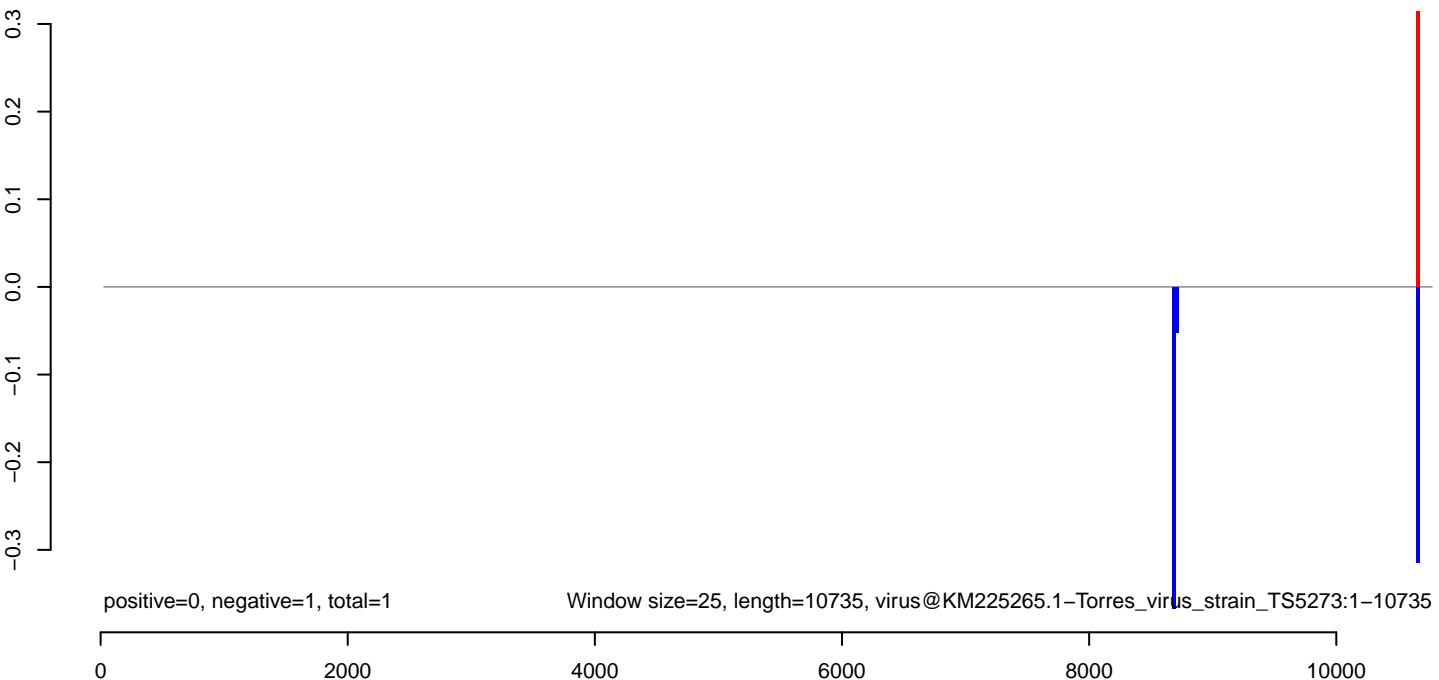
CuQuin_MG_7dpi3.18_23.rep



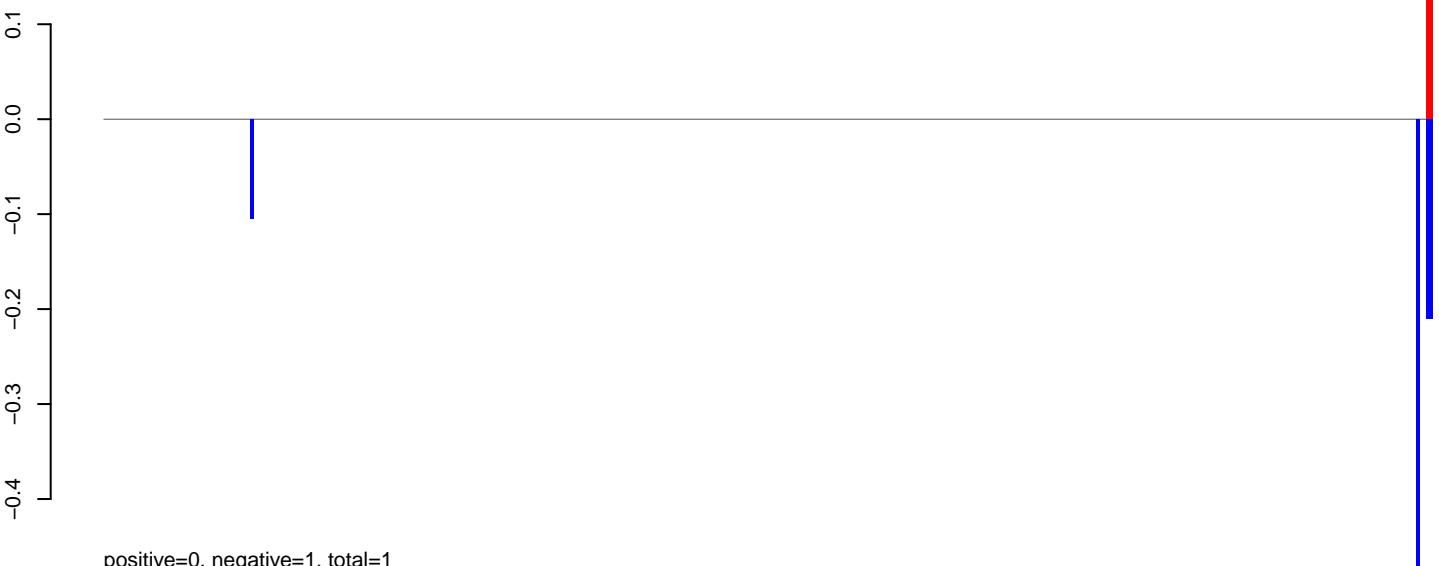
CuQuin_MG_7dpi3.24_35.rep



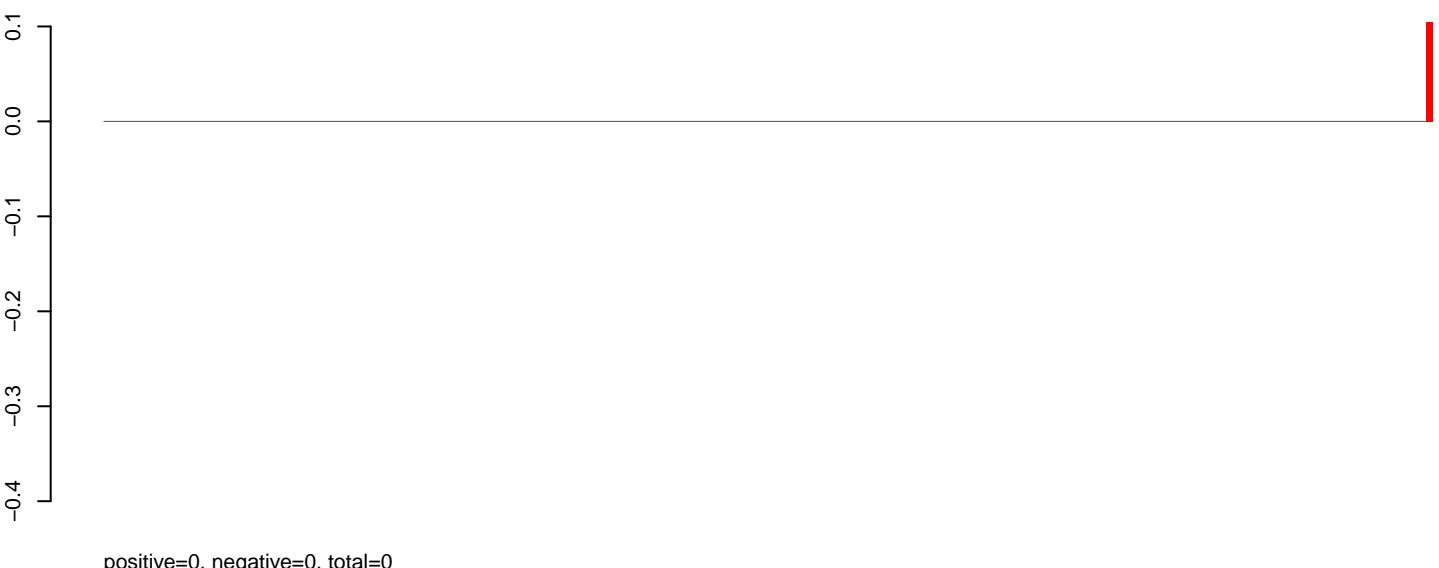
CuQuin_MG_7dpi3.rep



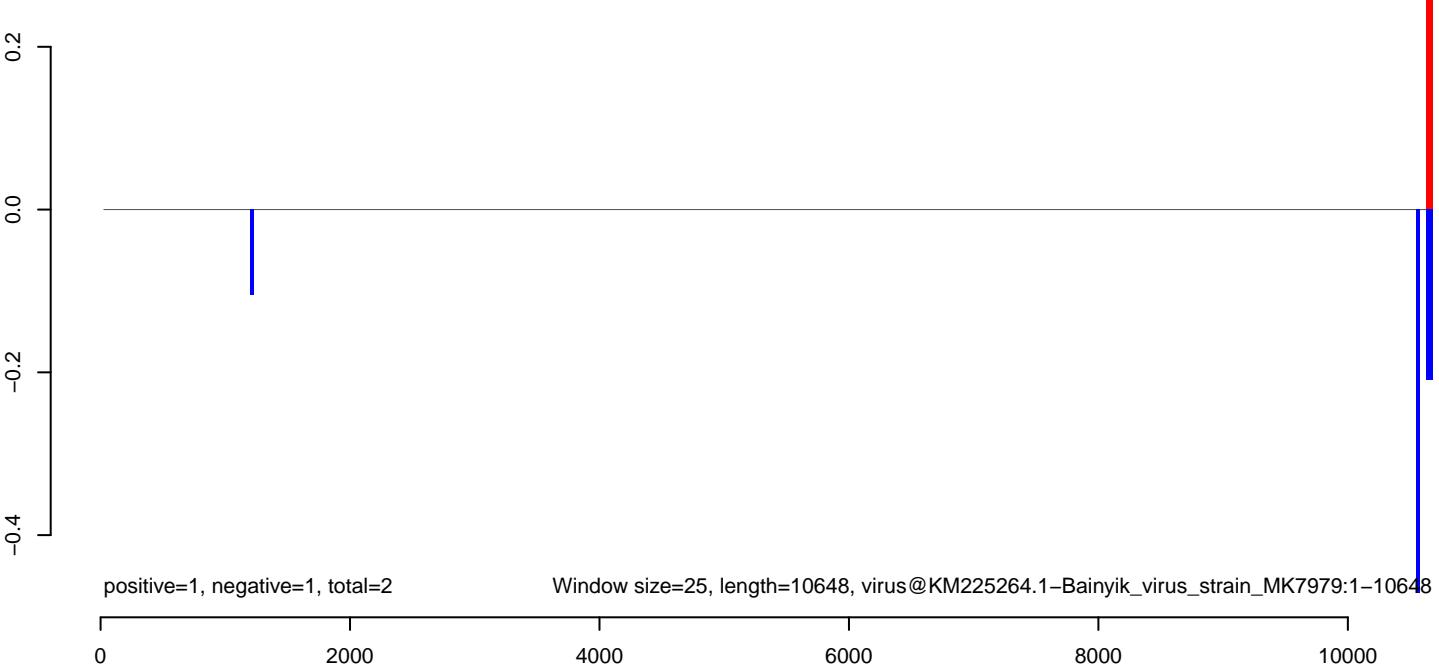
CuQuin_MG_7dpi3.18_23.rep



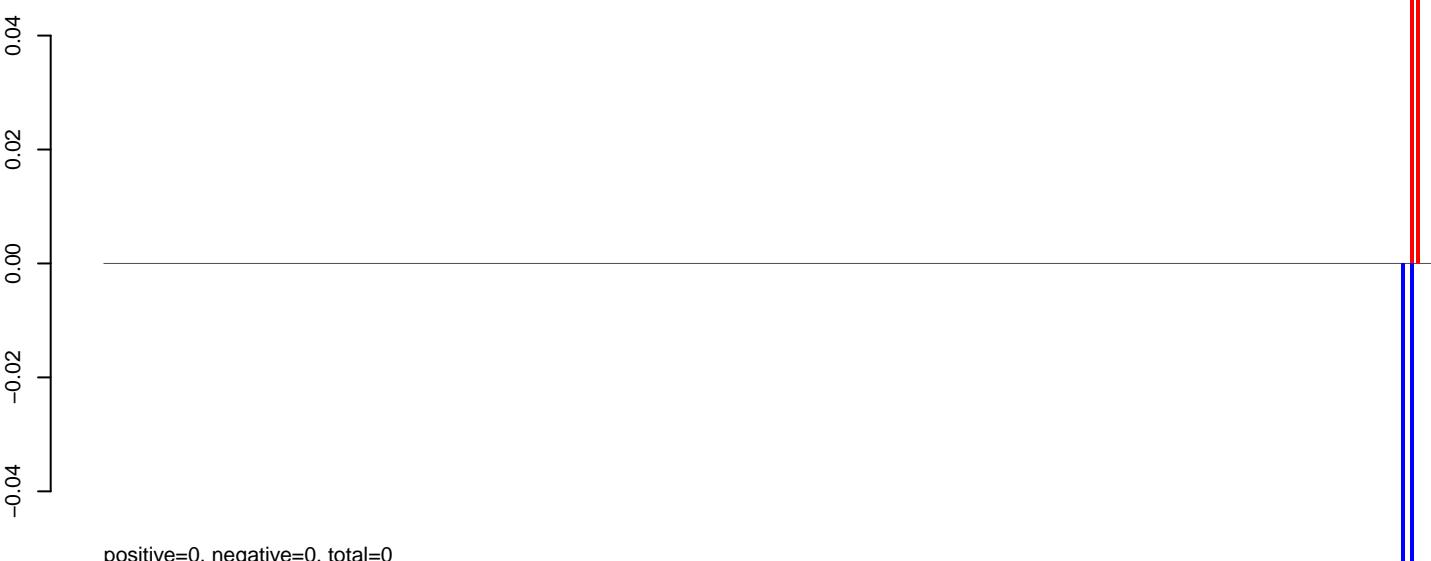
CuQuin_MG_7dpi3.24_35.rep



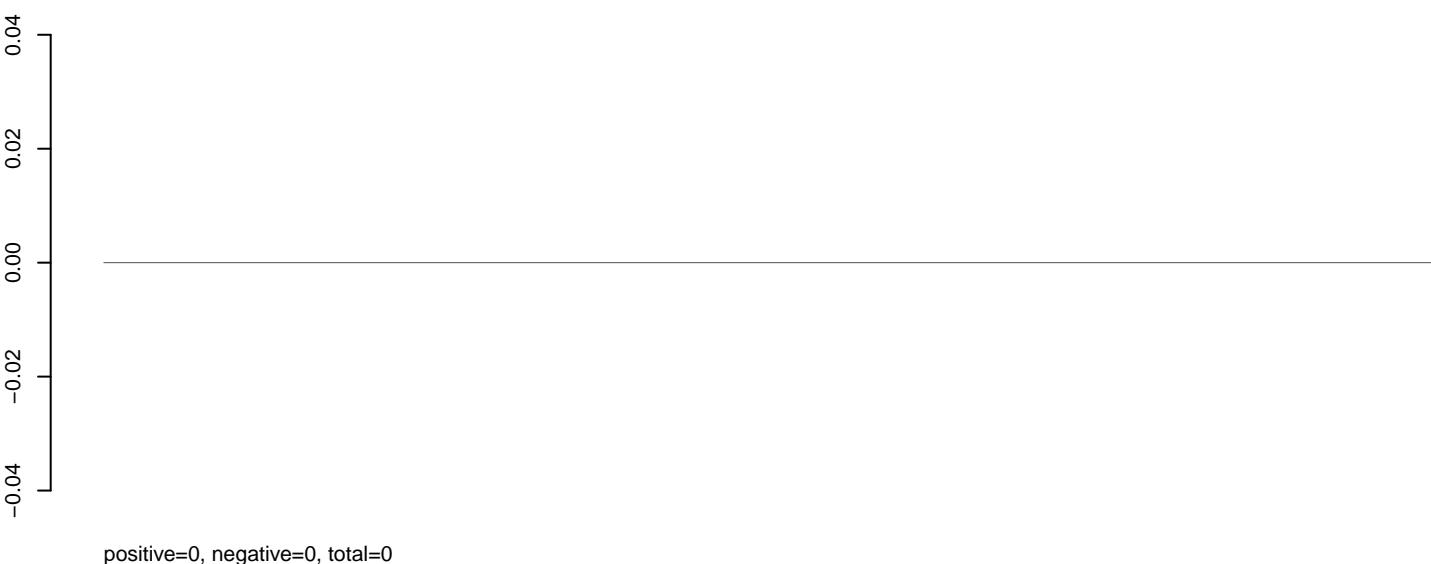
CuQuin_MG_7dpi3.rep



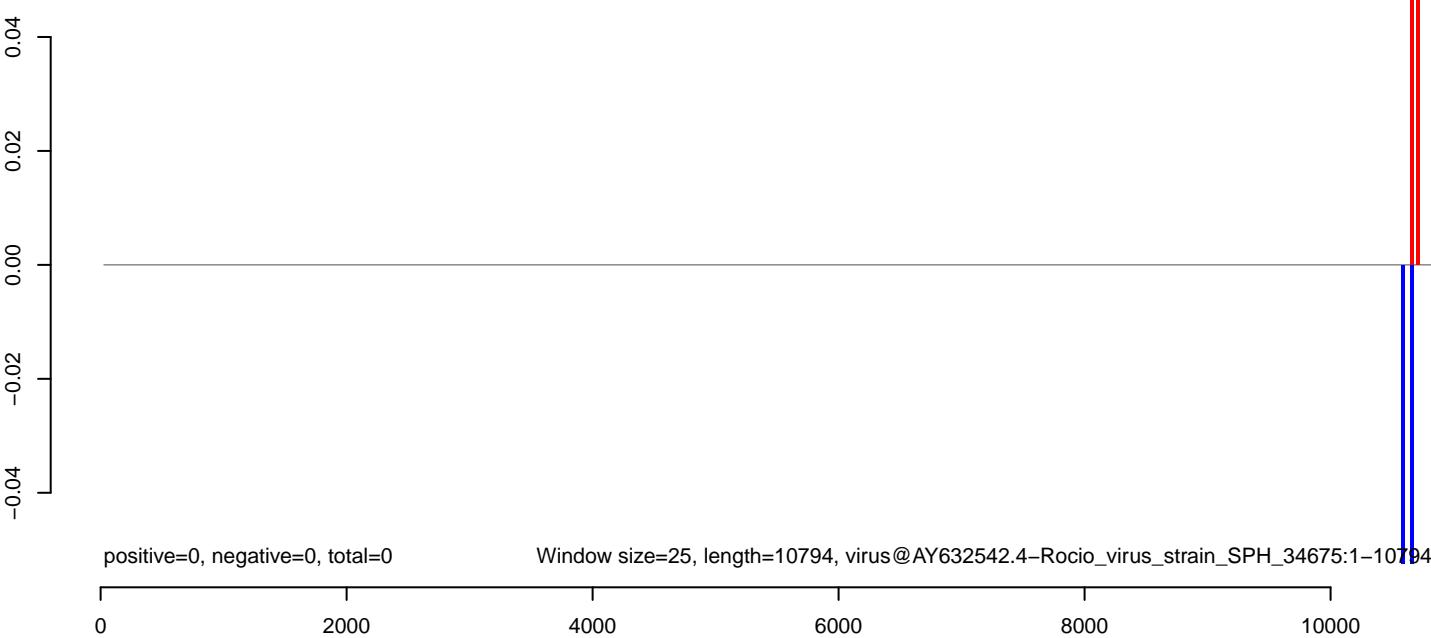
CuQuin_MG_7dpi3.18_23.rep



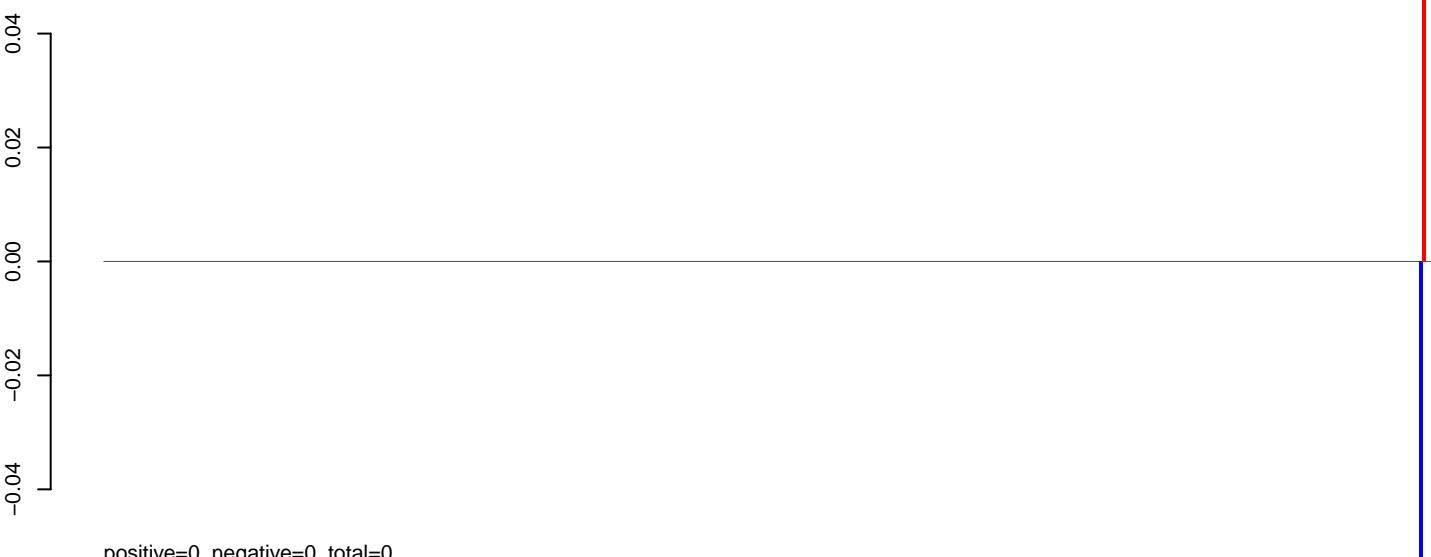
CuQuin_MG_7dpi3.24_35.rep



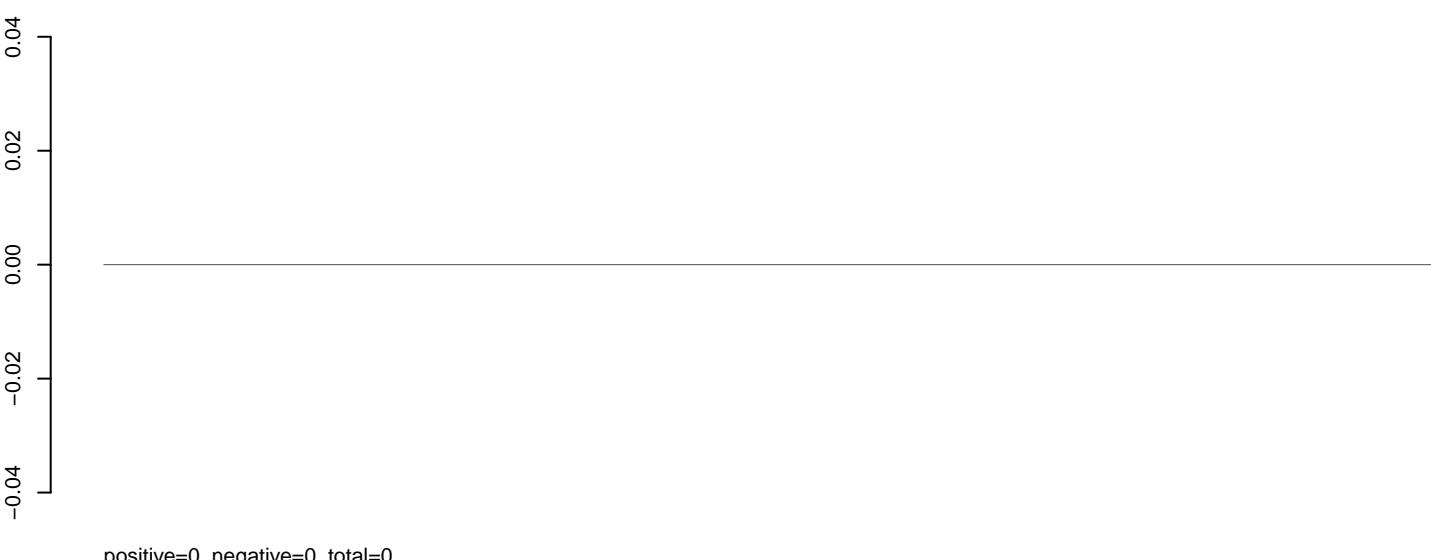
CuQuin_MG_7dpi3.rep



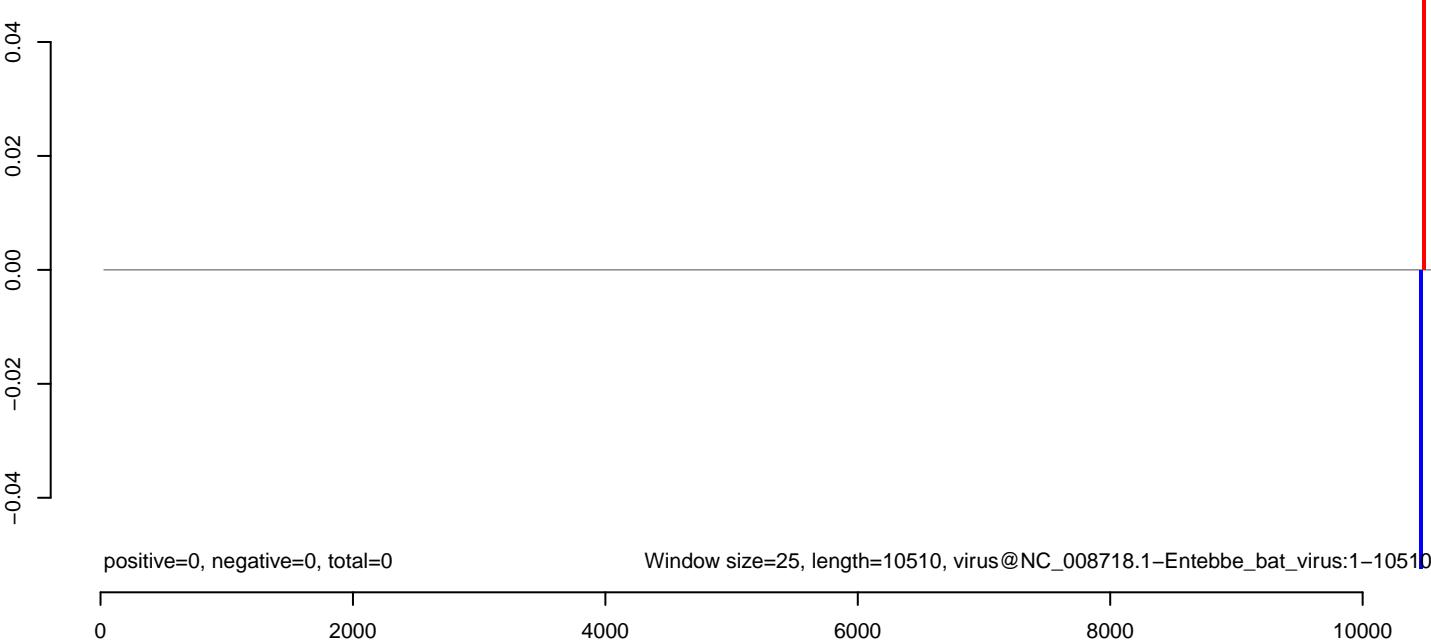
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



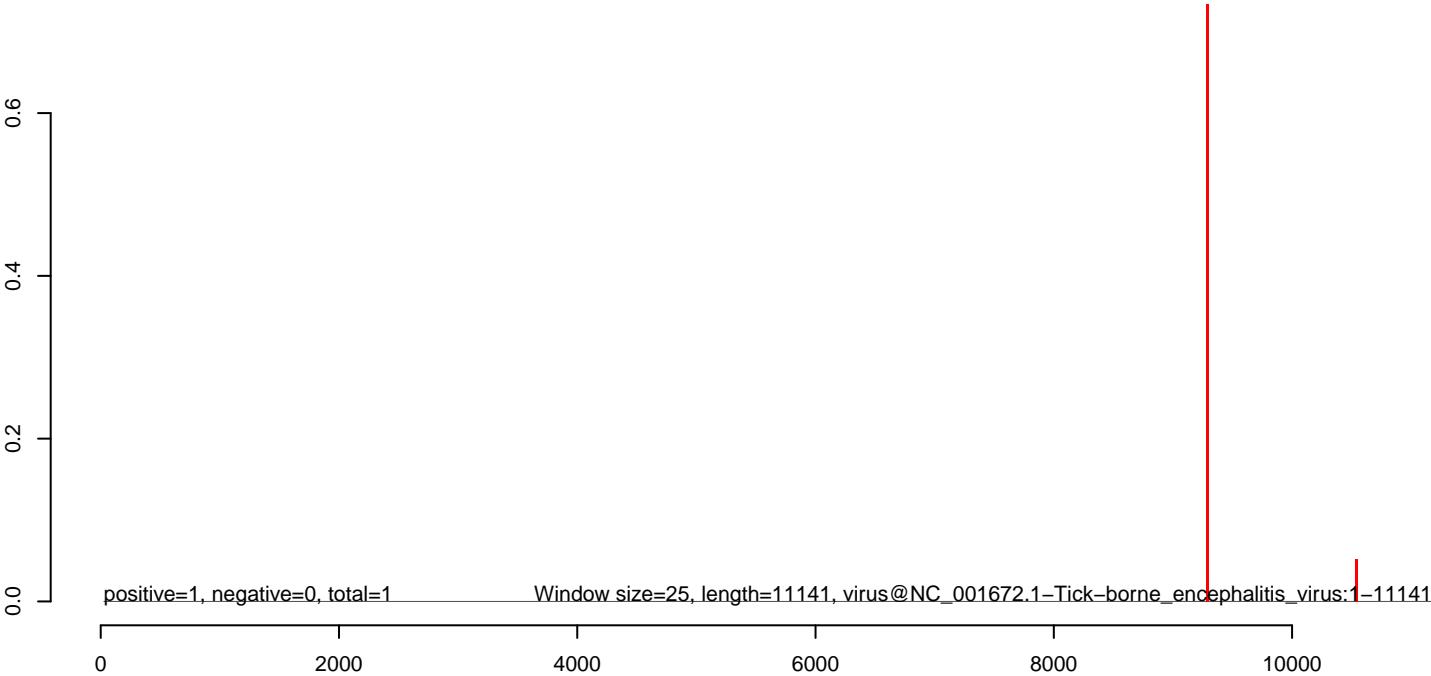
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

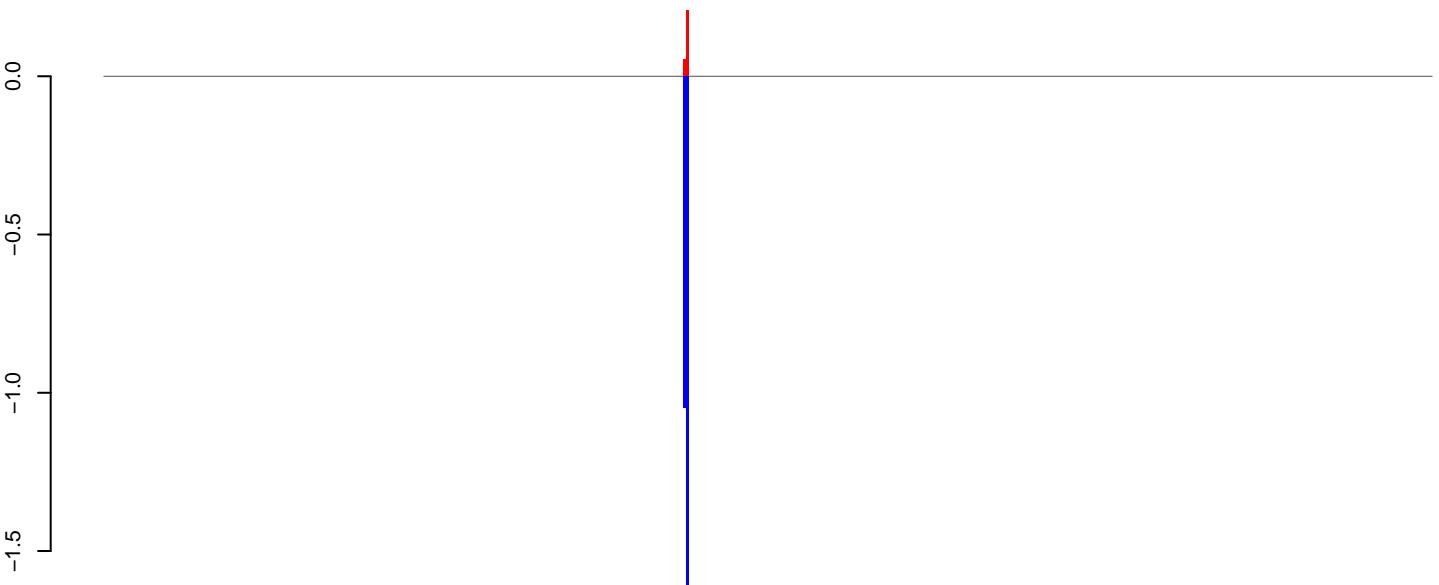
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

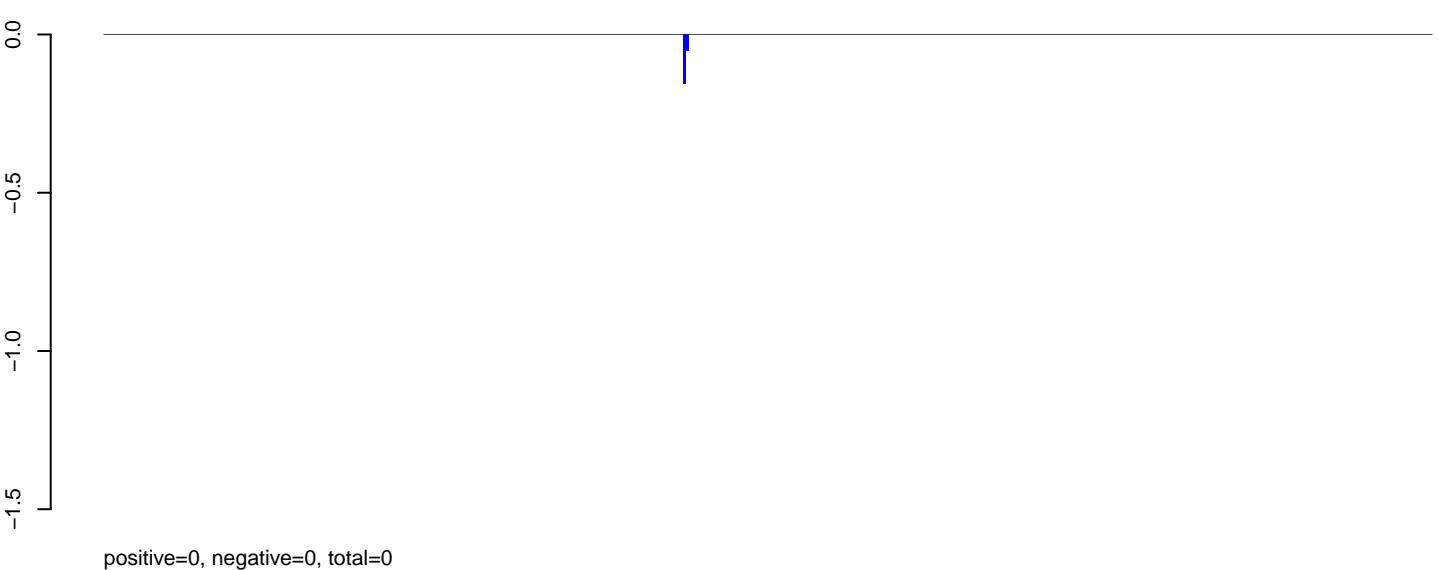
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=10427, virus@MF176249.1_Wuhan_Mosquito_Virus_6_strain_mos172gb65361_segment_1_2_3_4_5:1-10427

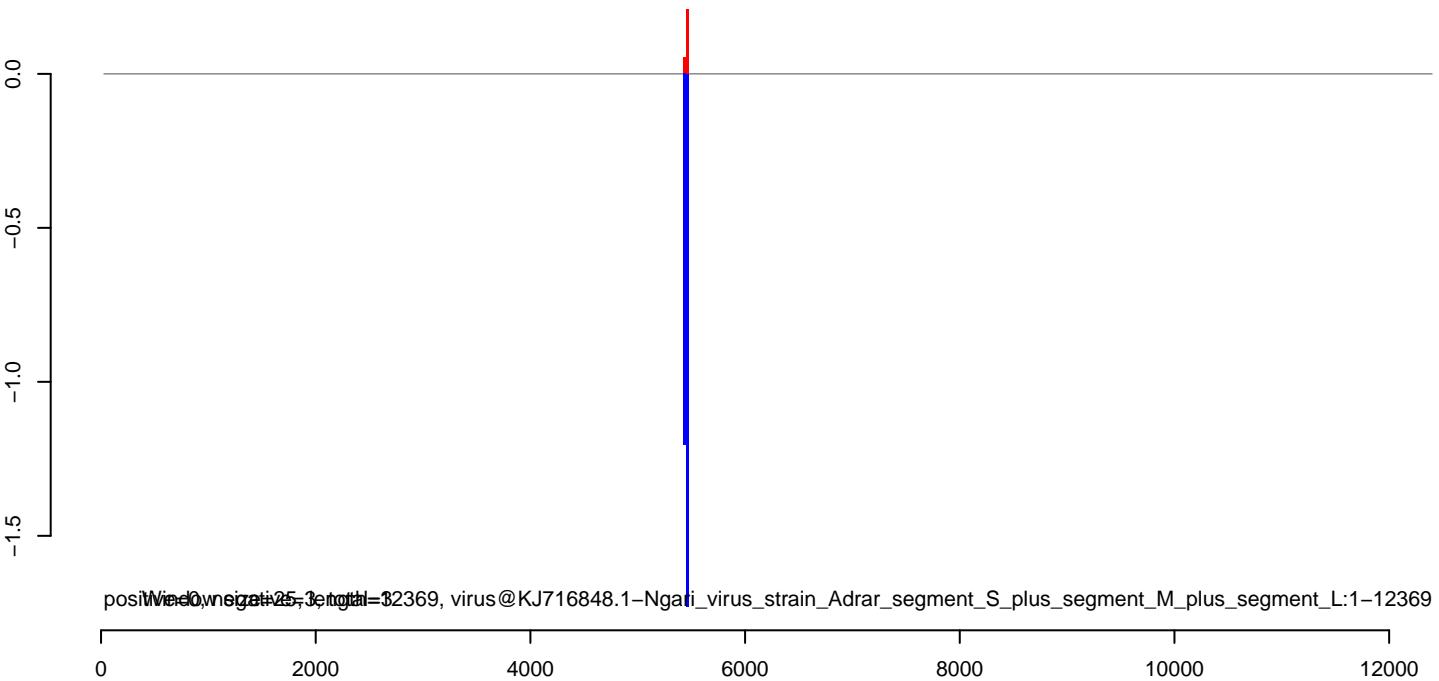
CuQuin_MG_7dpi3.18_23.rep



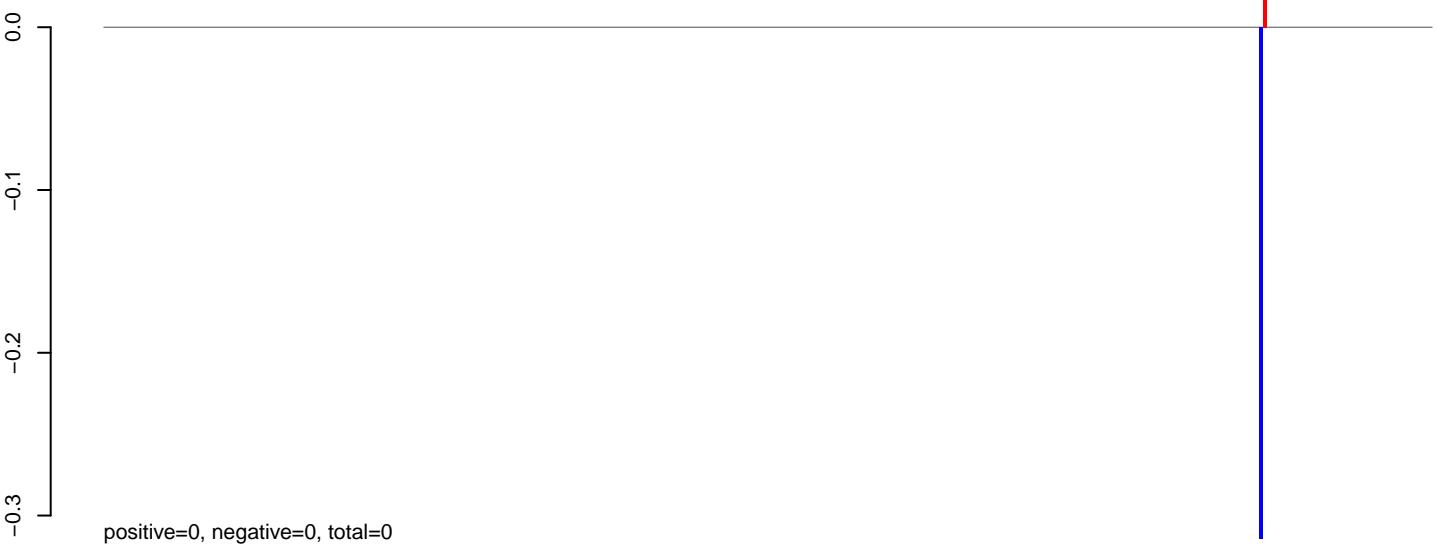
CuQuin_MG_7dpi3.24_35.rep



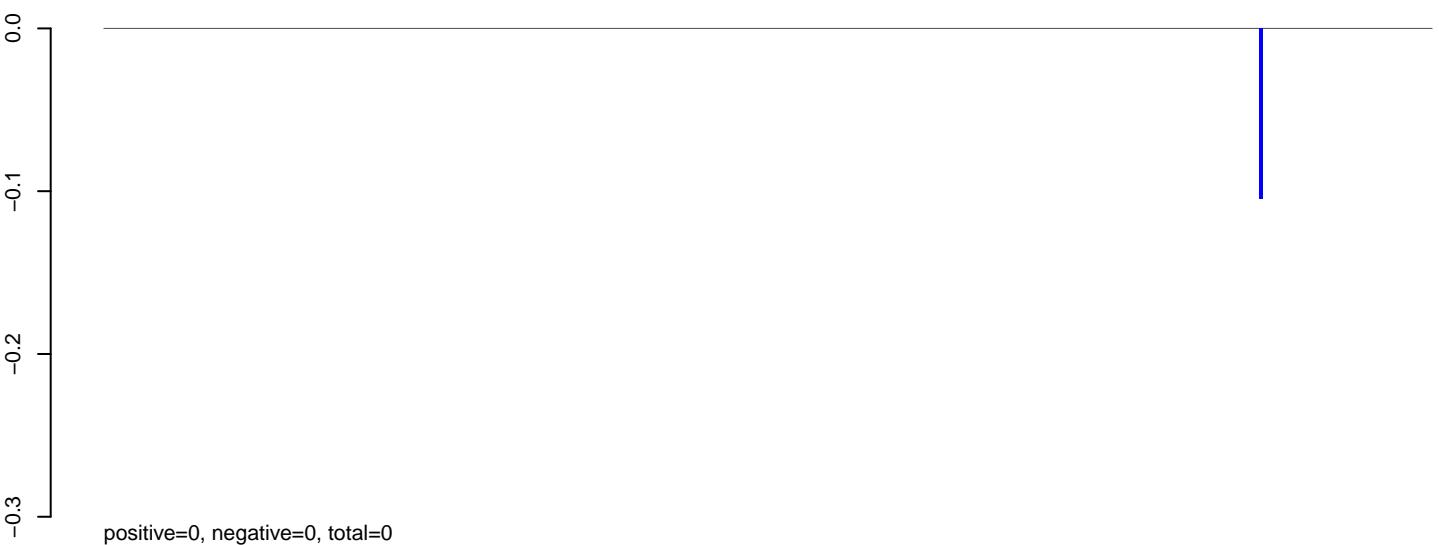
CuQuin_MG_7dpi3.rep



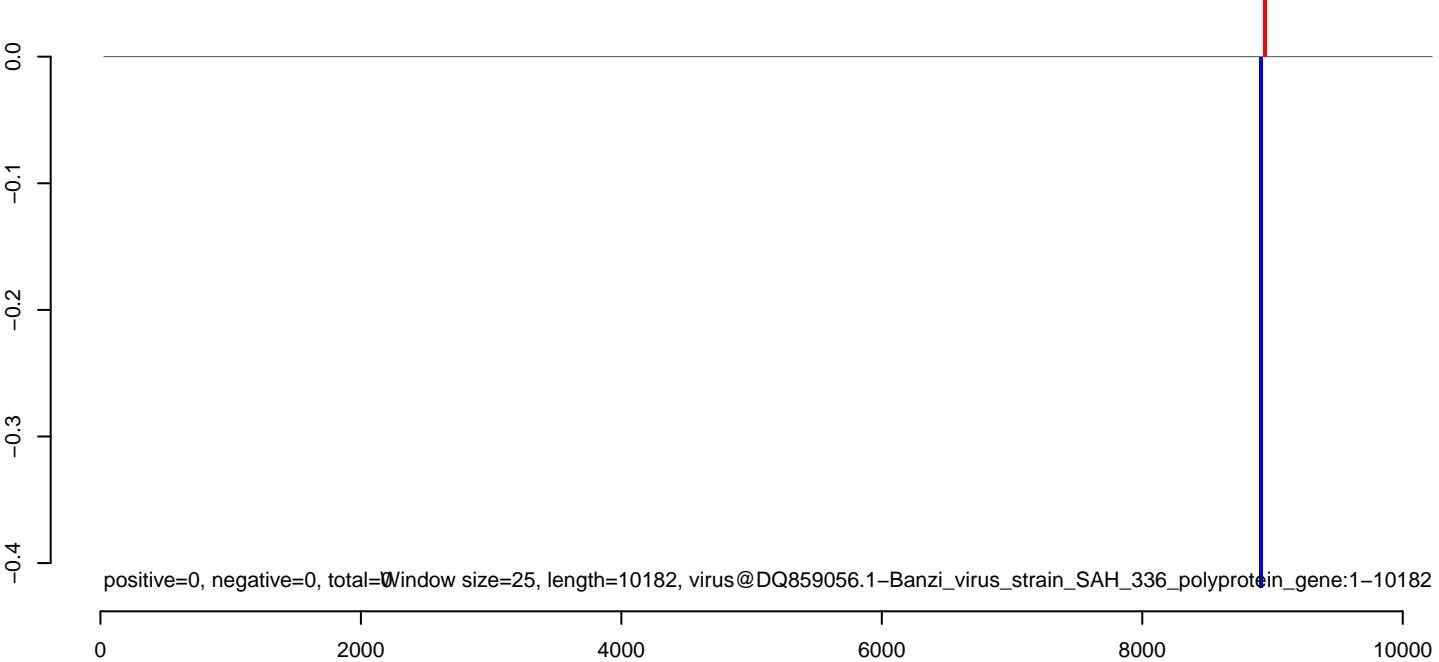
CuQuin_MG_7dpi3.18_23.rep



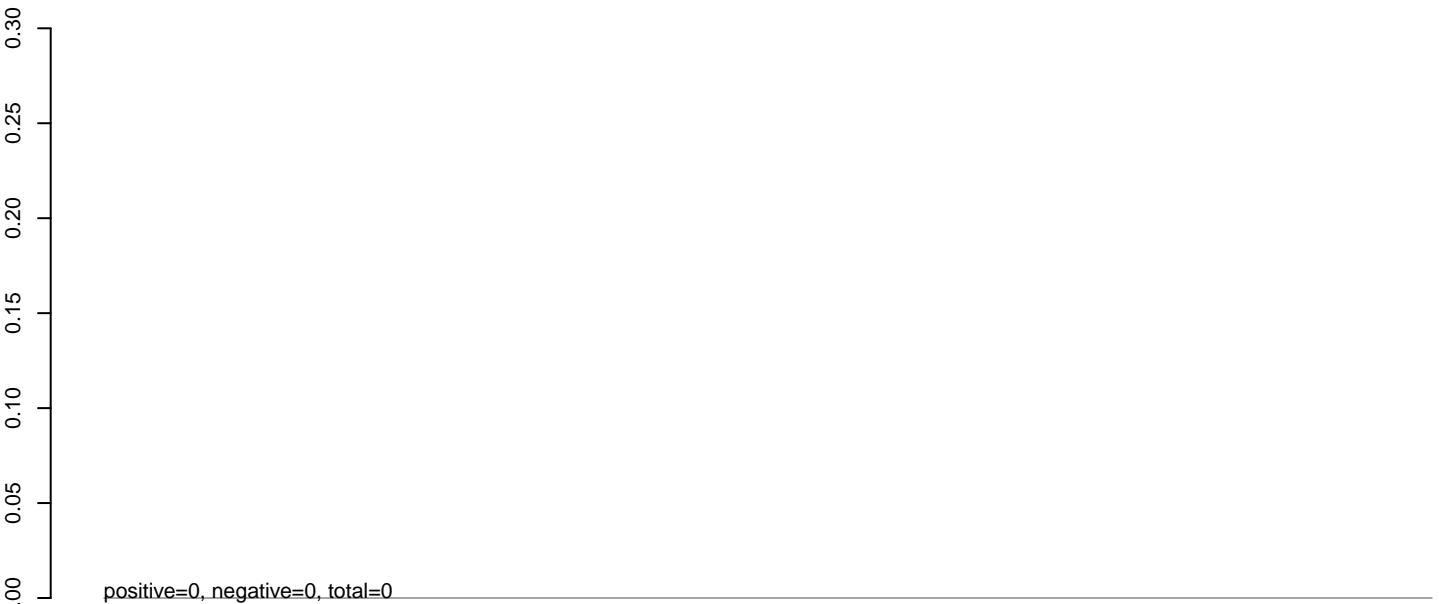
CuQuin_MG_7dpi3.24_35.rep



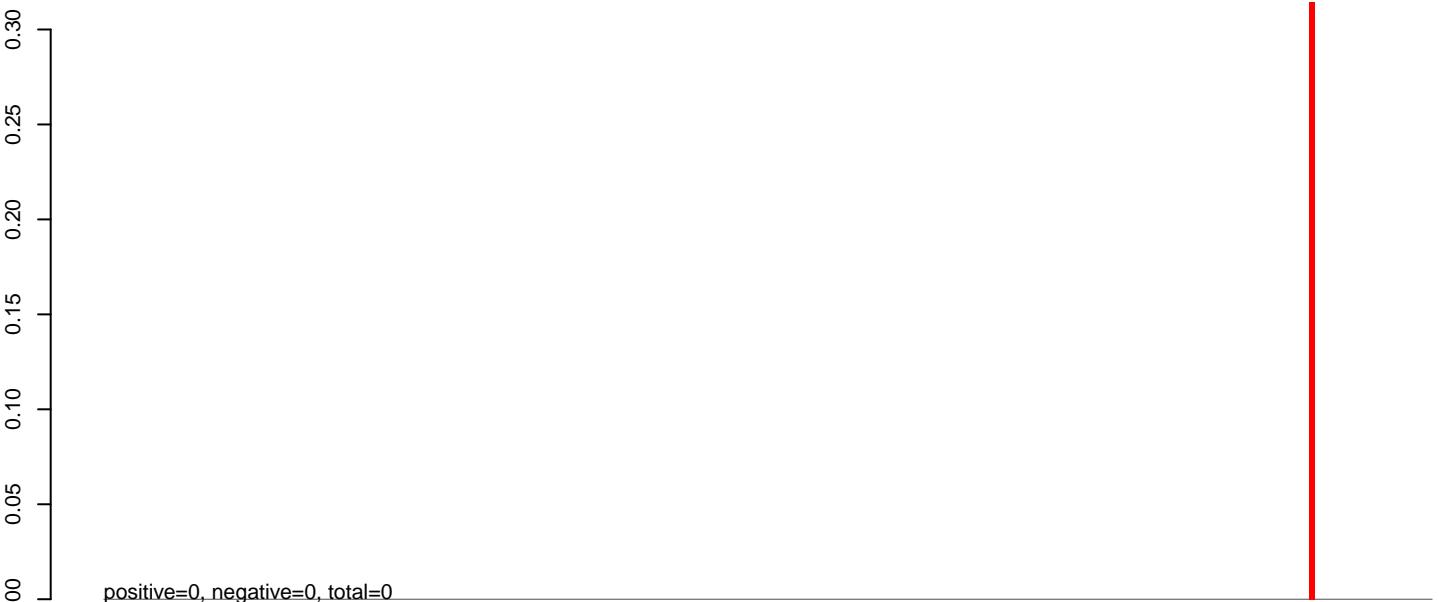
CuQuin_MG_7dpi3.rep



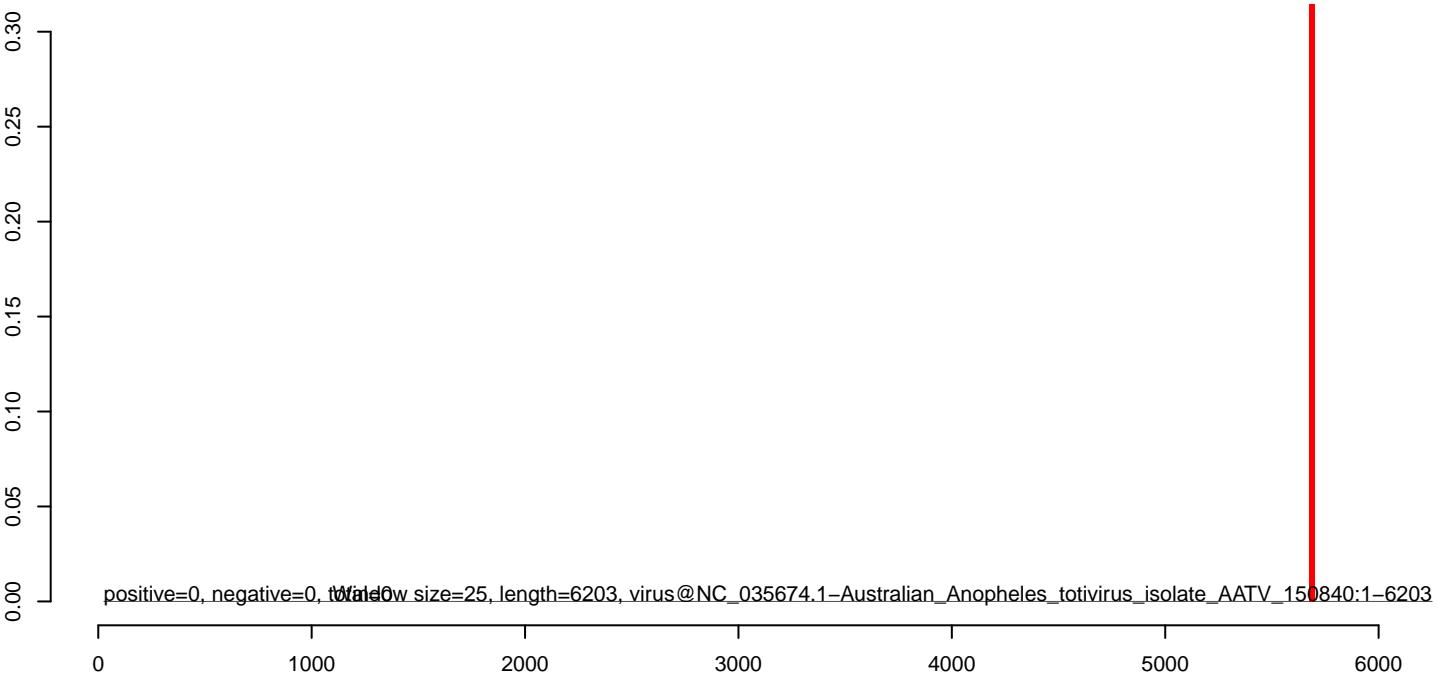
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep



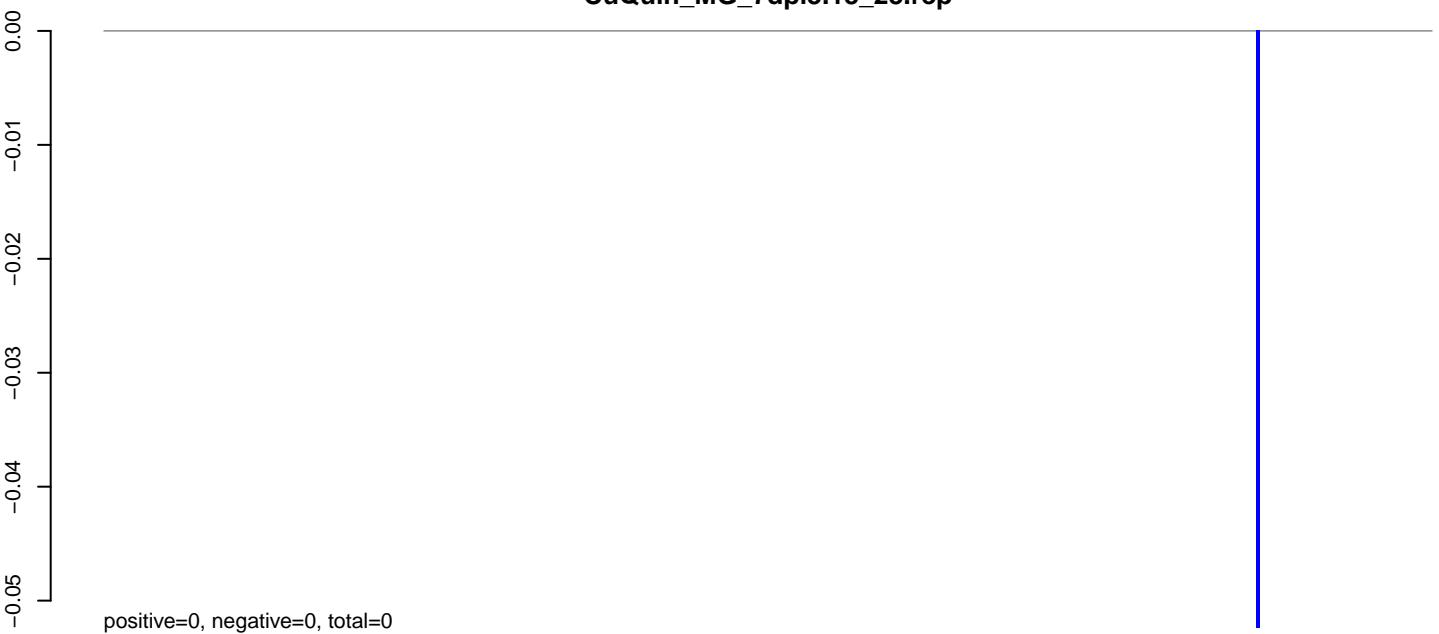
CuQuin_MG_7dpi3.24_35.rep



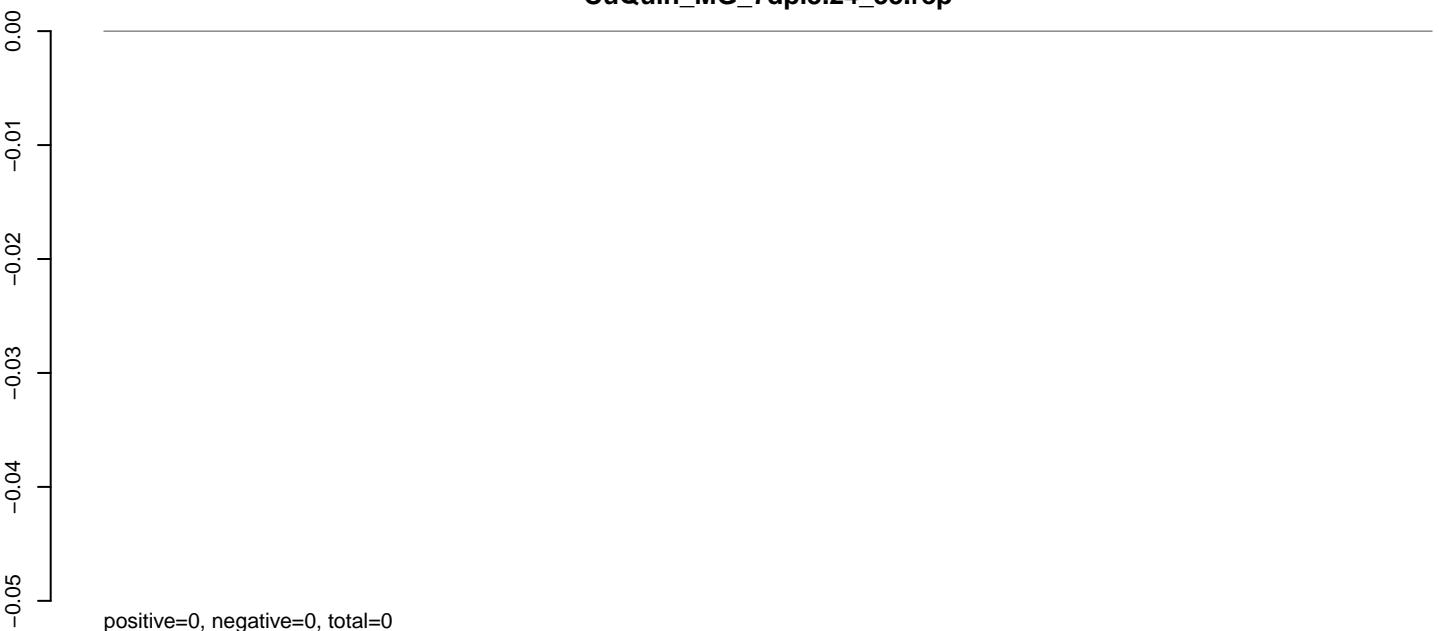
CuQuin_MG_7dpi3.rep



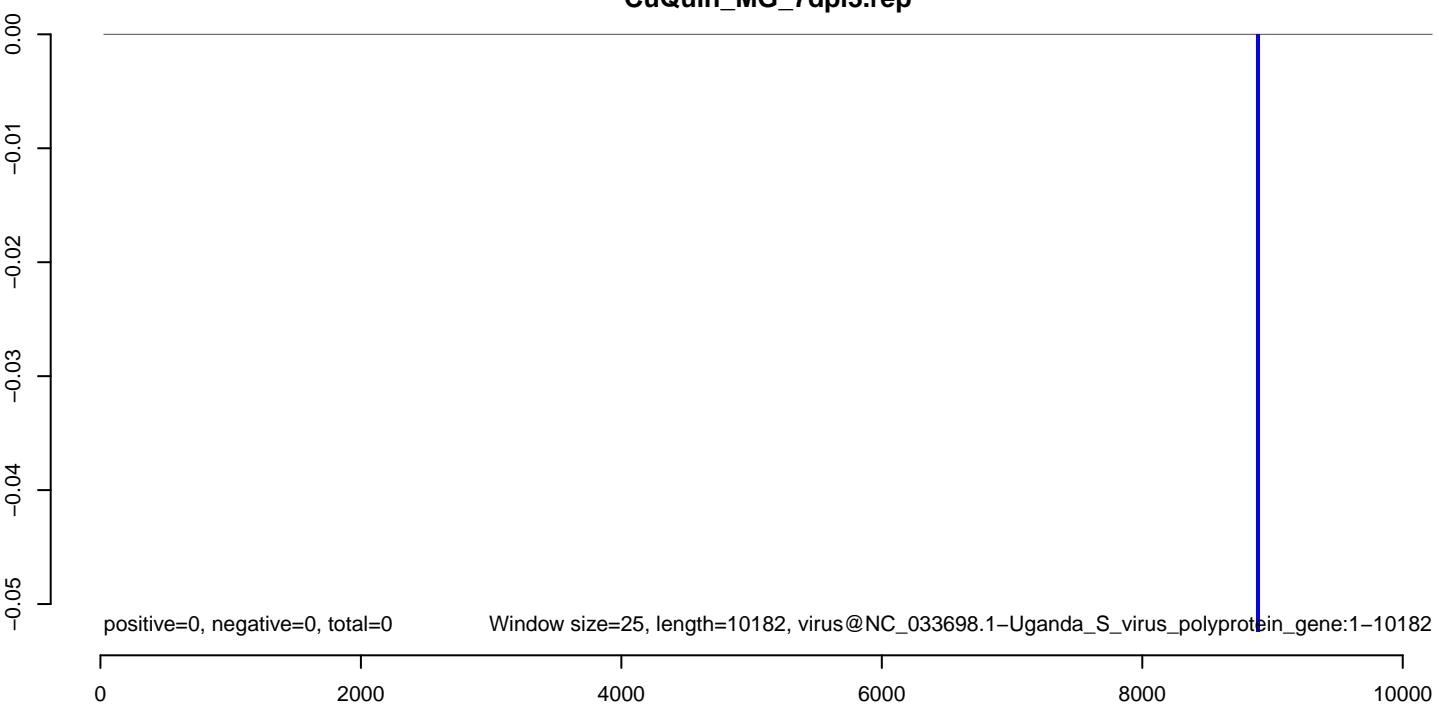
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



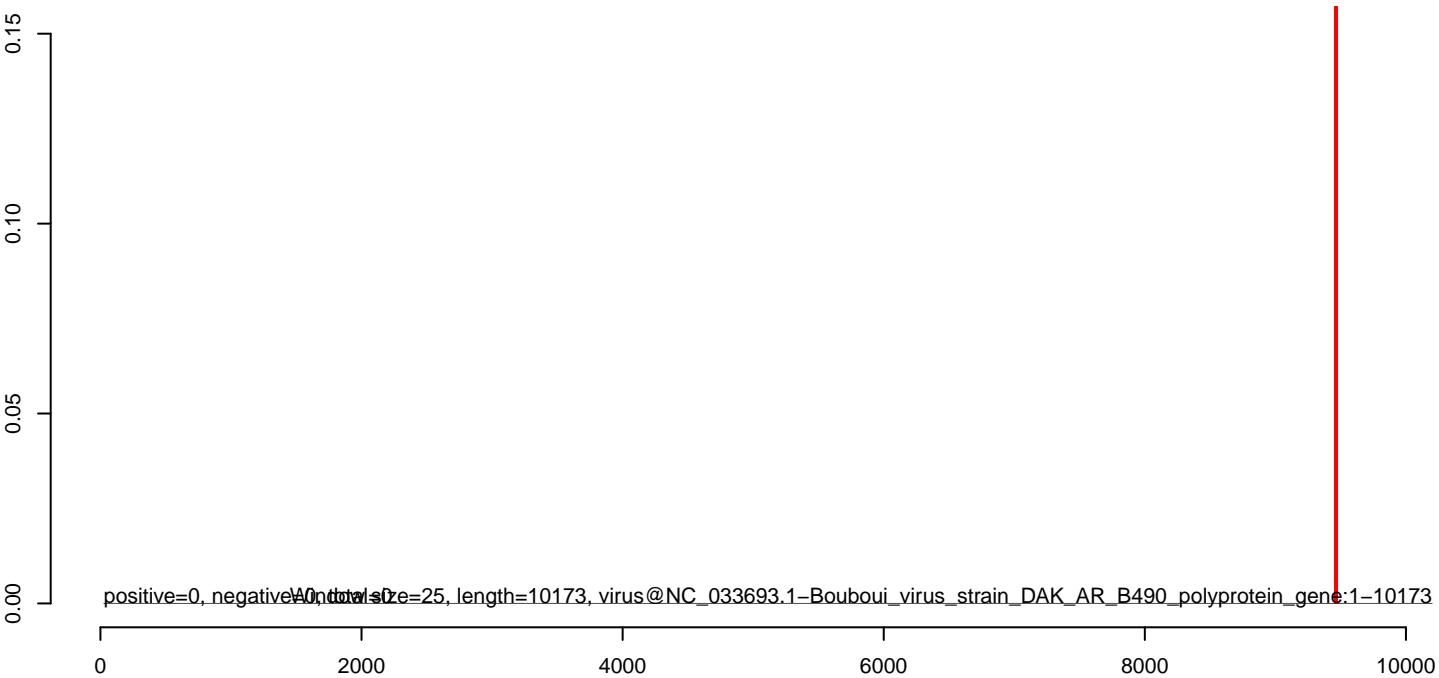
CuQuin_MG_7dpi3.18_23.rep



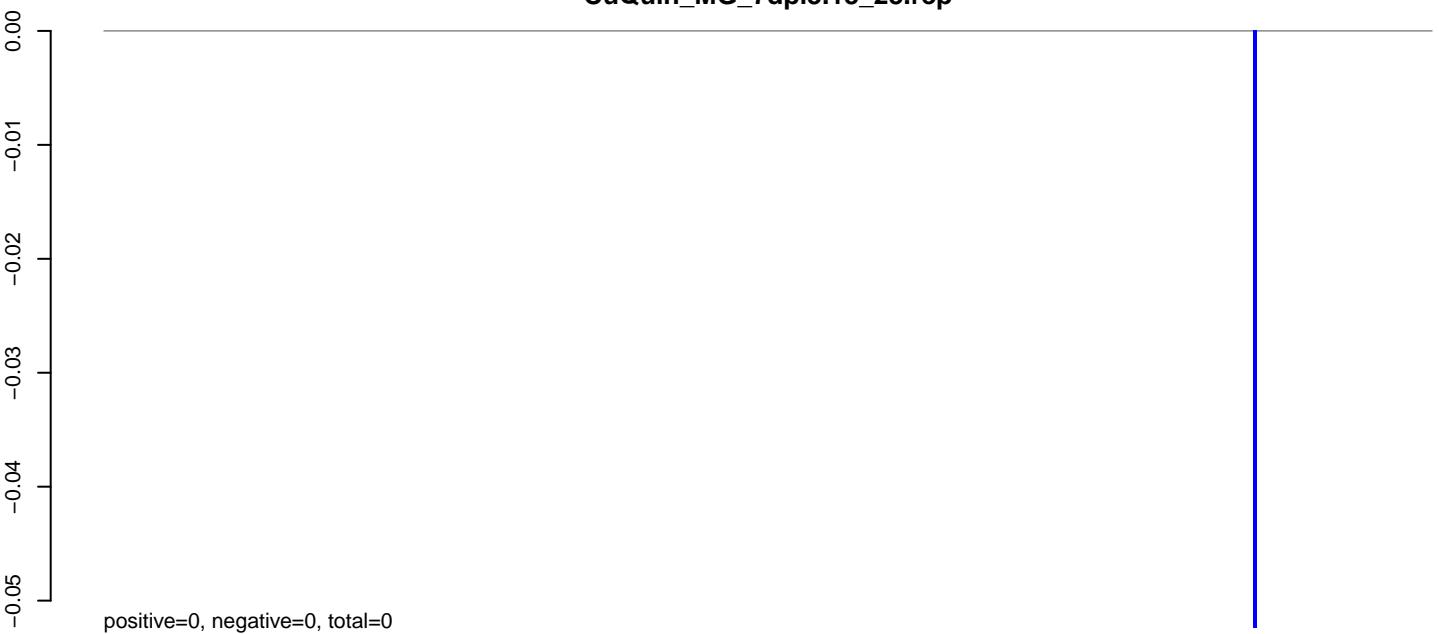
CuQuin_MG_7dpi3.24_35.rep



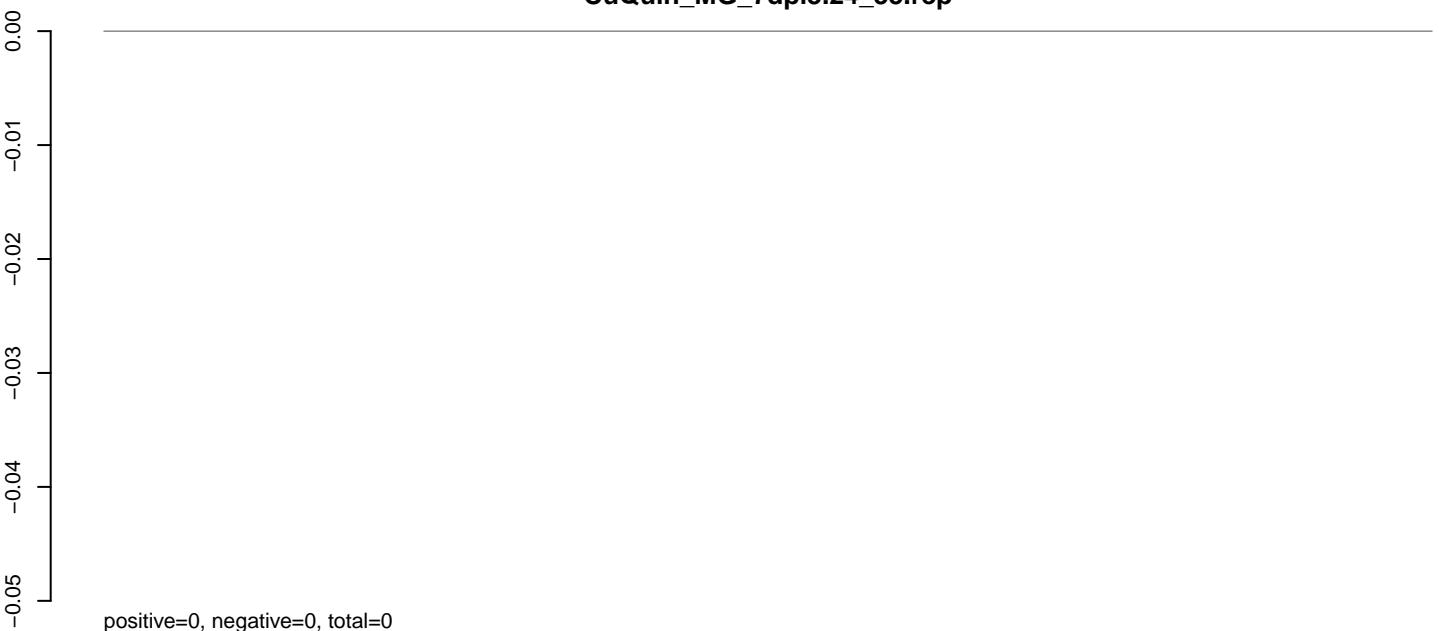
CuQuin_MG_7dpi3.rep



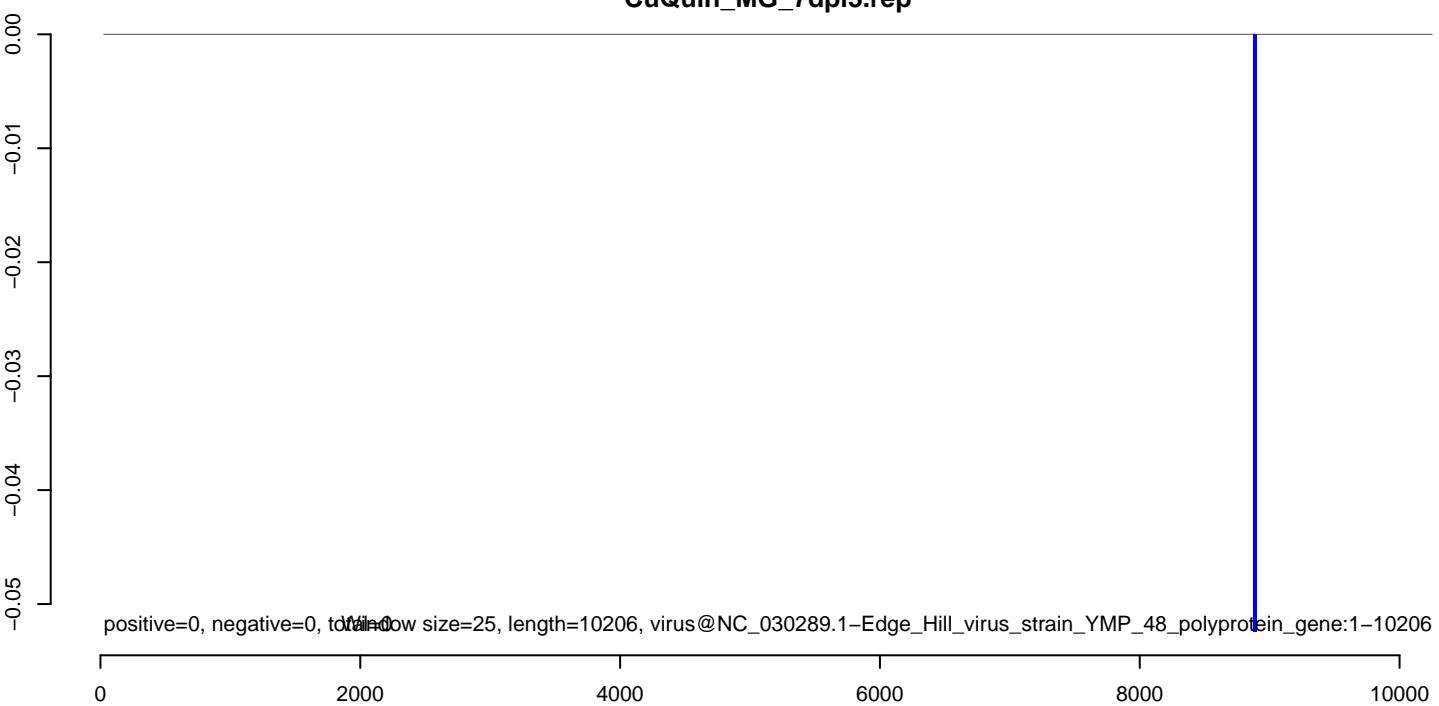
CuQuin_MG_7dpi3.18_23.rep



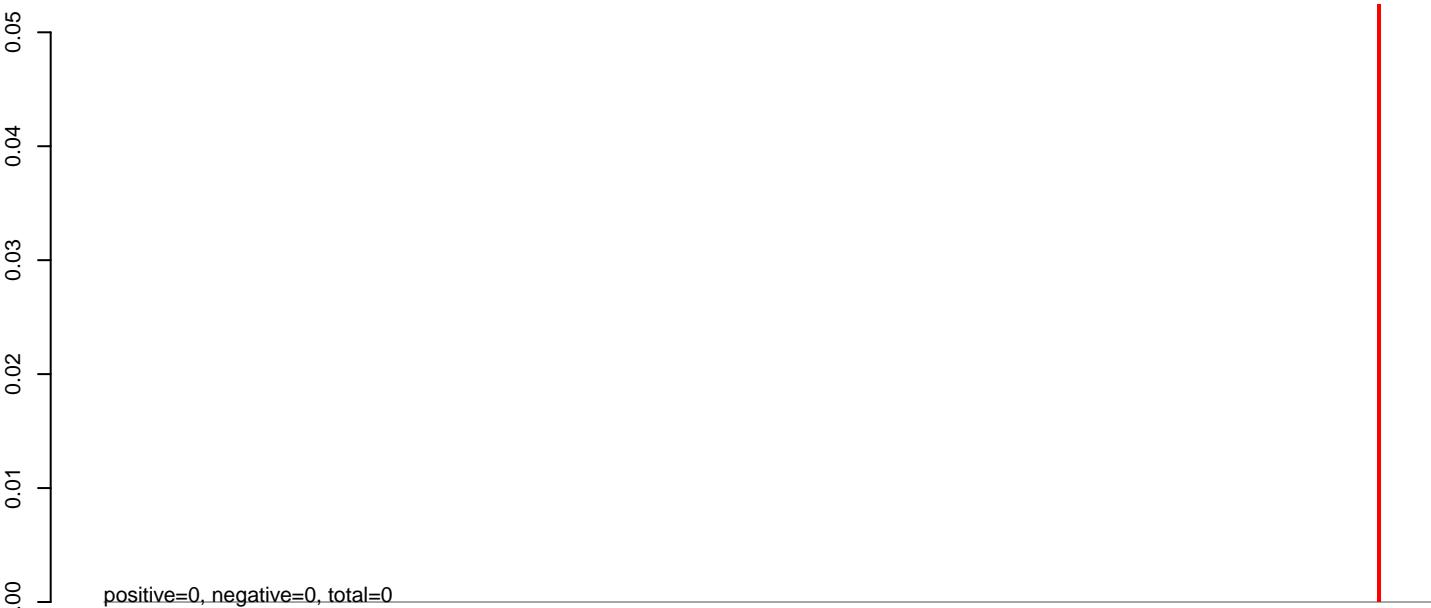
CuQuin_MG_7dpi3.24_35.rep



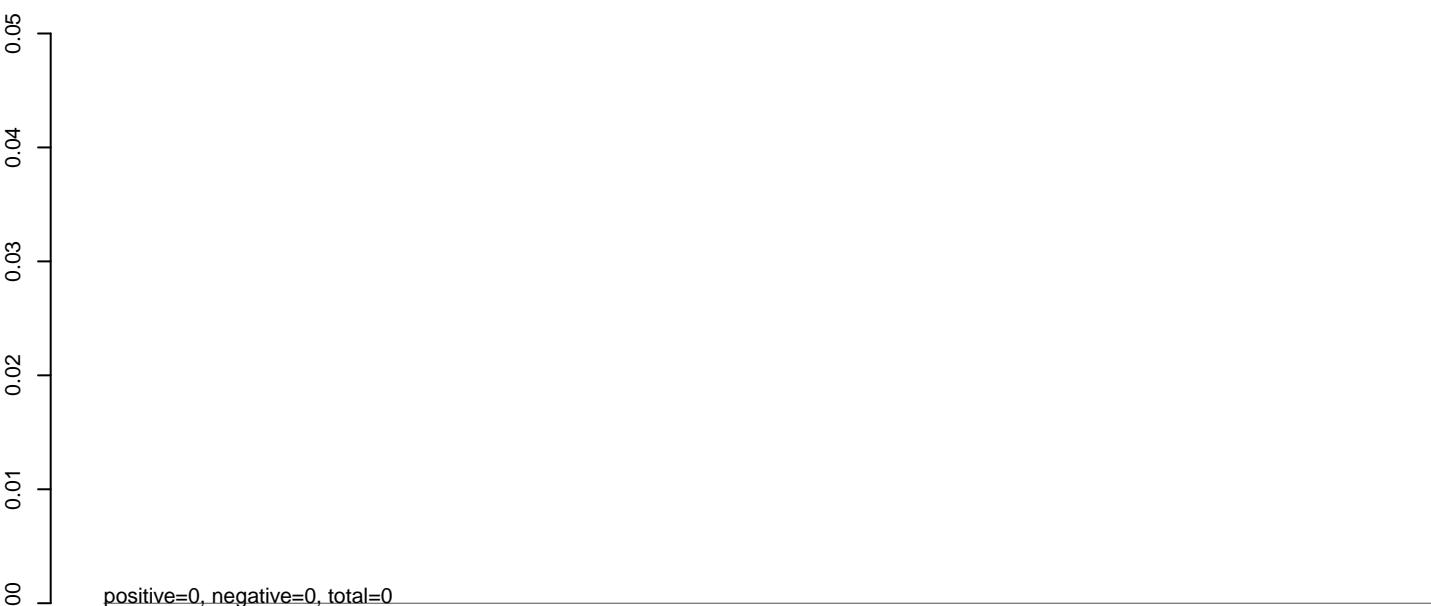
CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep



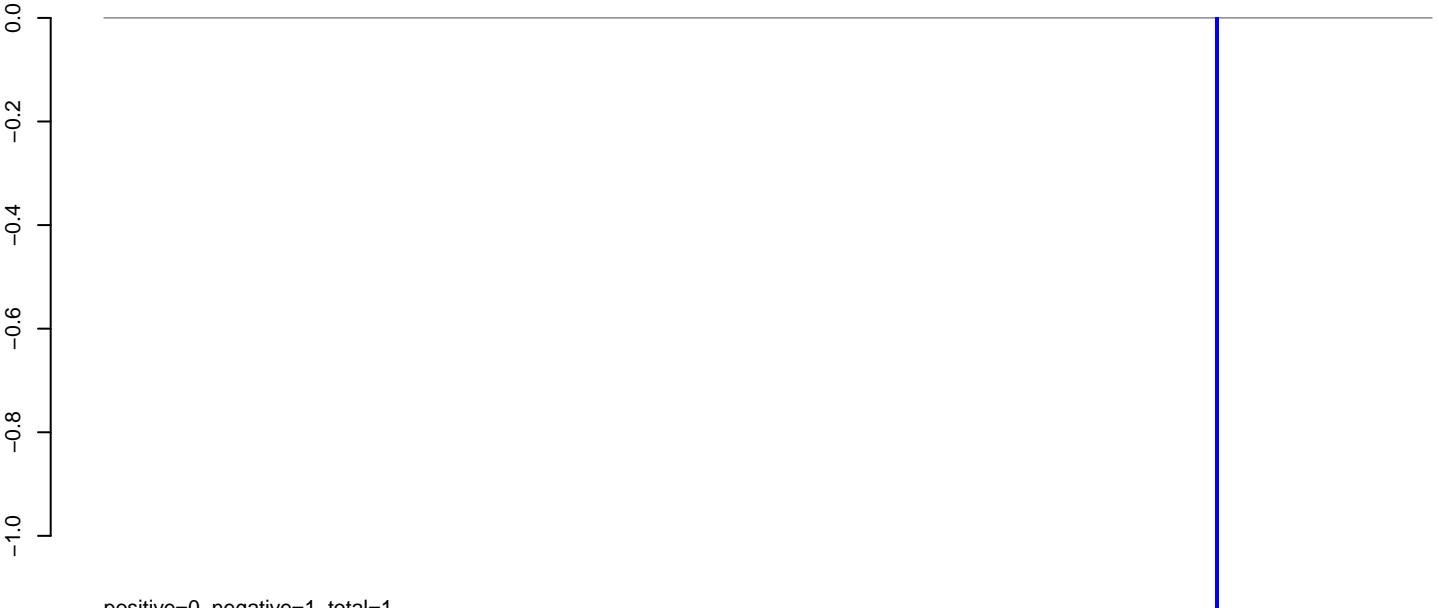
CuQuin_MG_7dpi3.24_35.rep



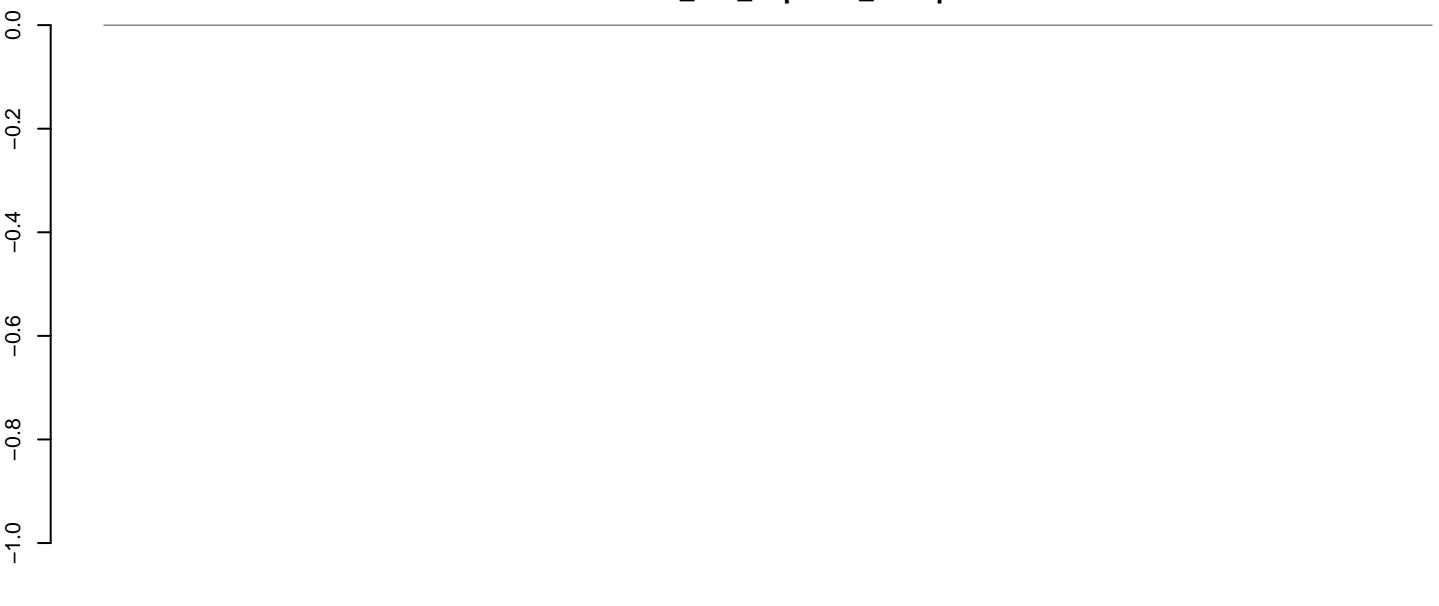
CuQuin_MG_7dpi3.rep



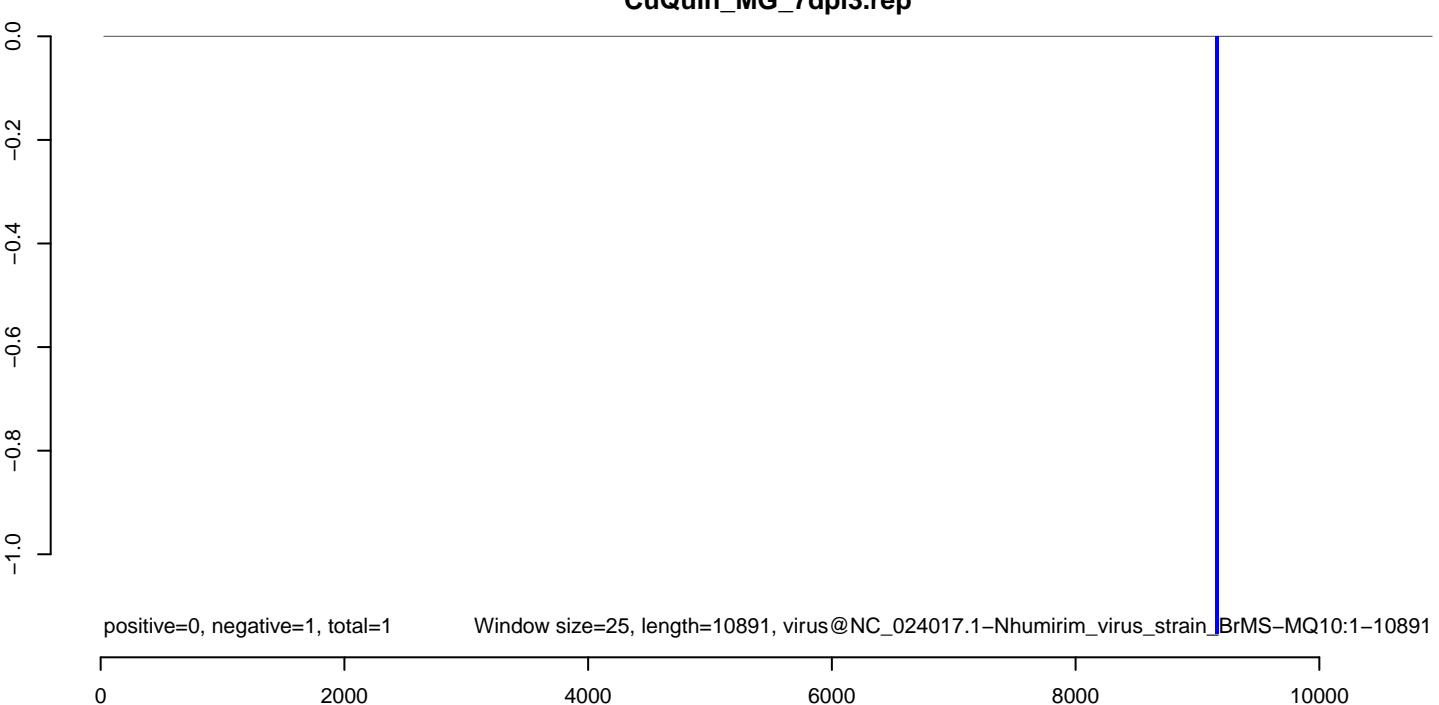
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



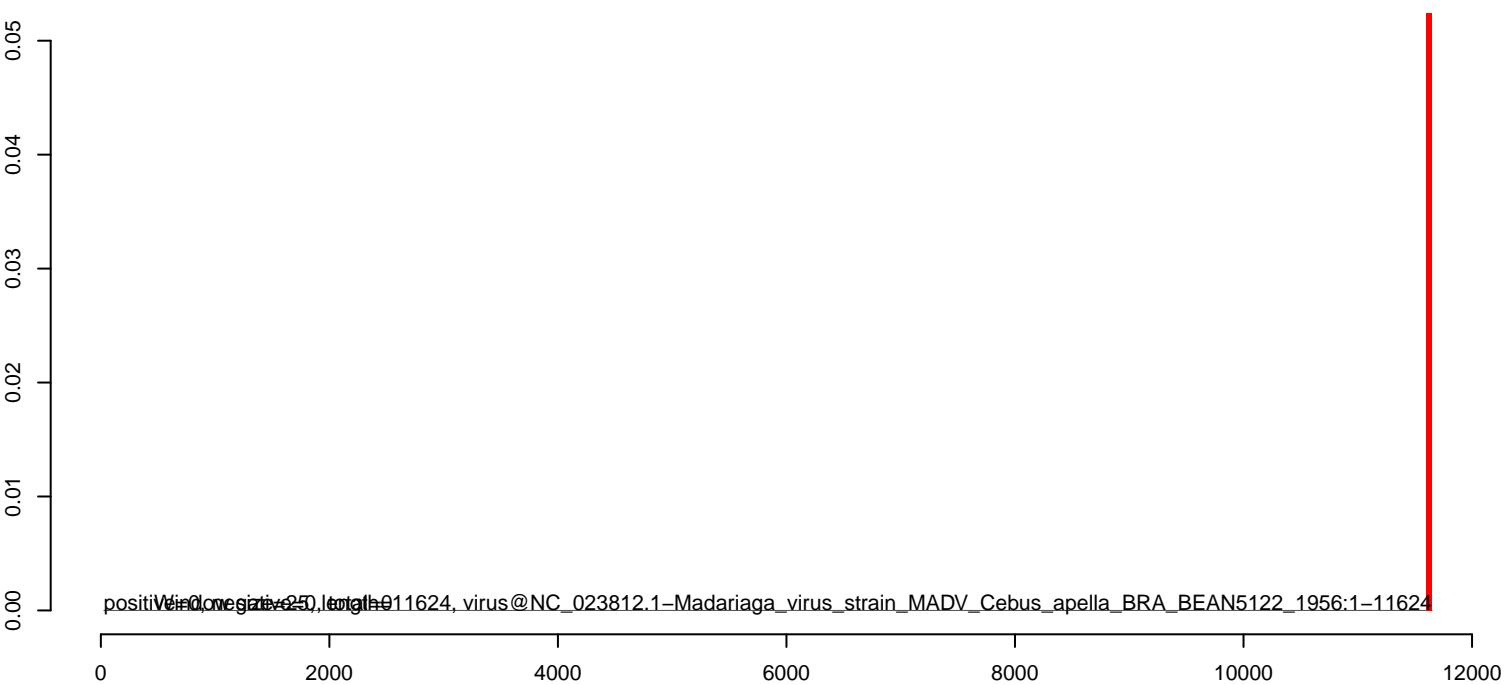
CuQuin_MG_7dpi3.18_23.rep



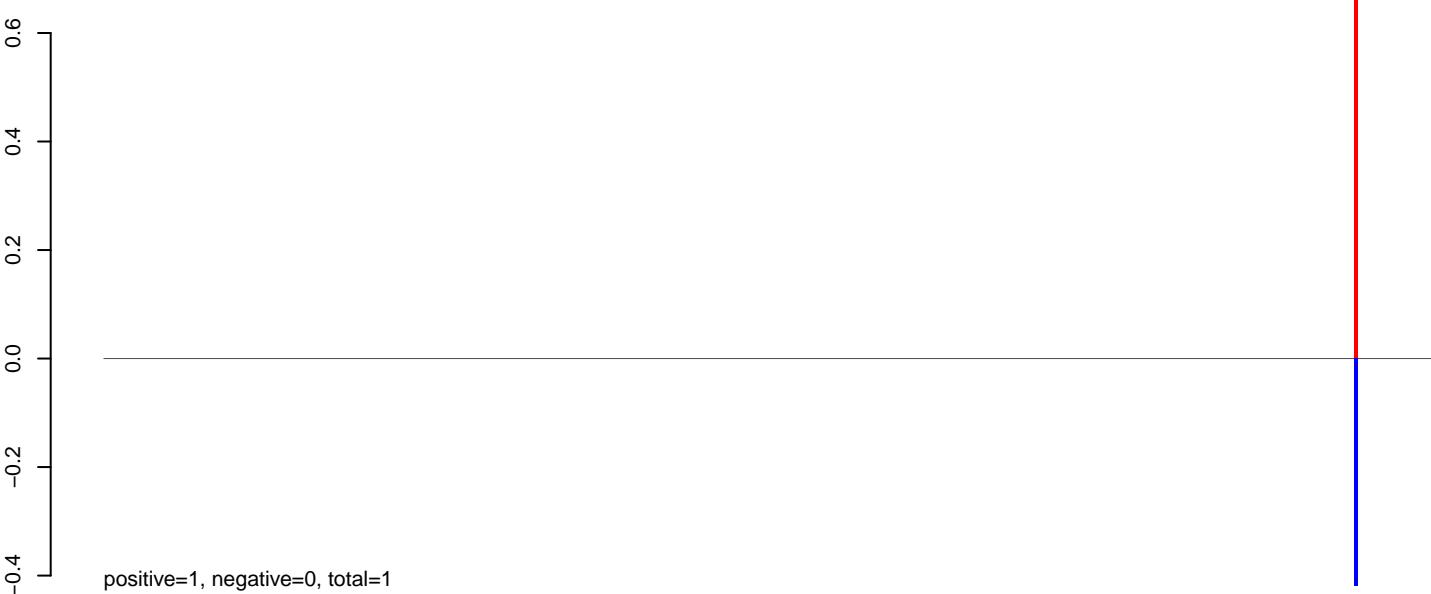
CuQuin_MG_7dpi3.24_35.rep



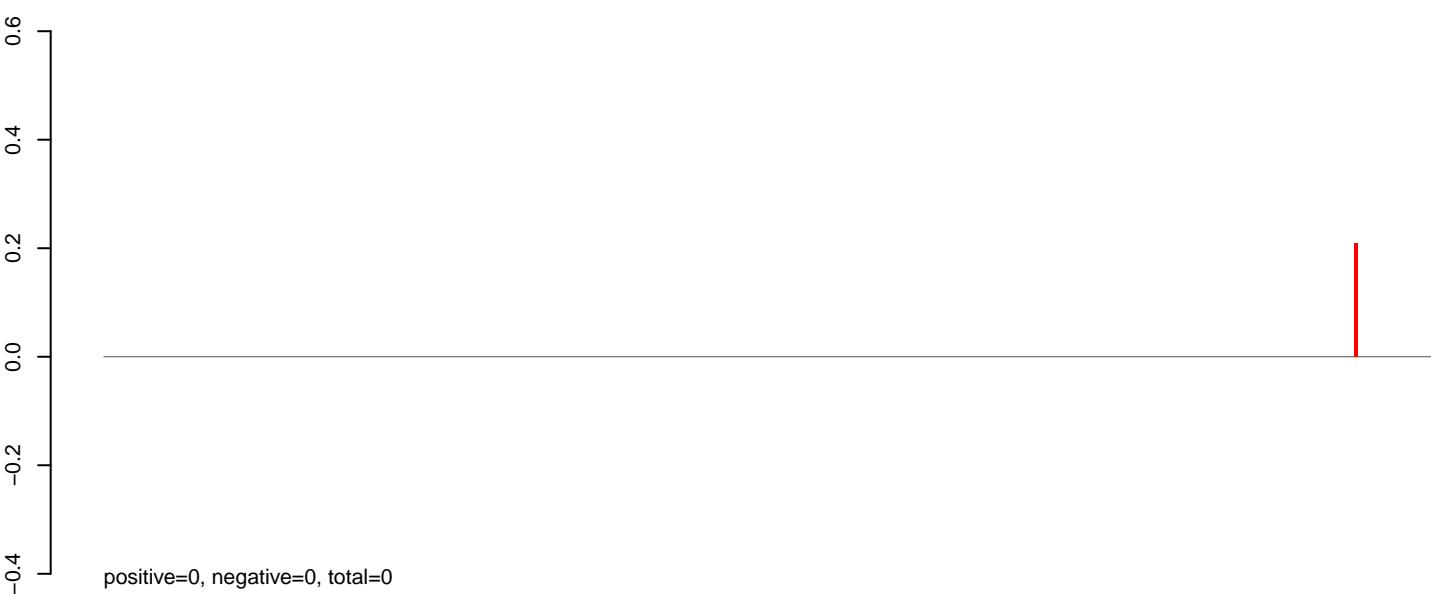
CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

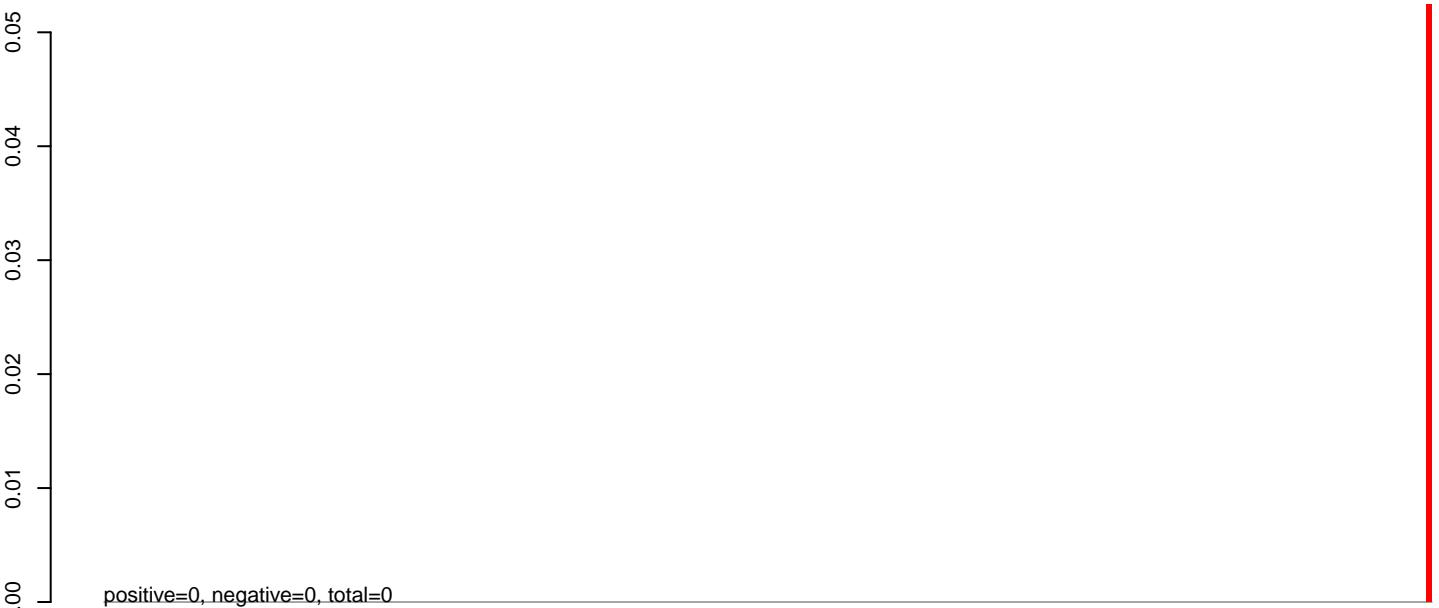
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

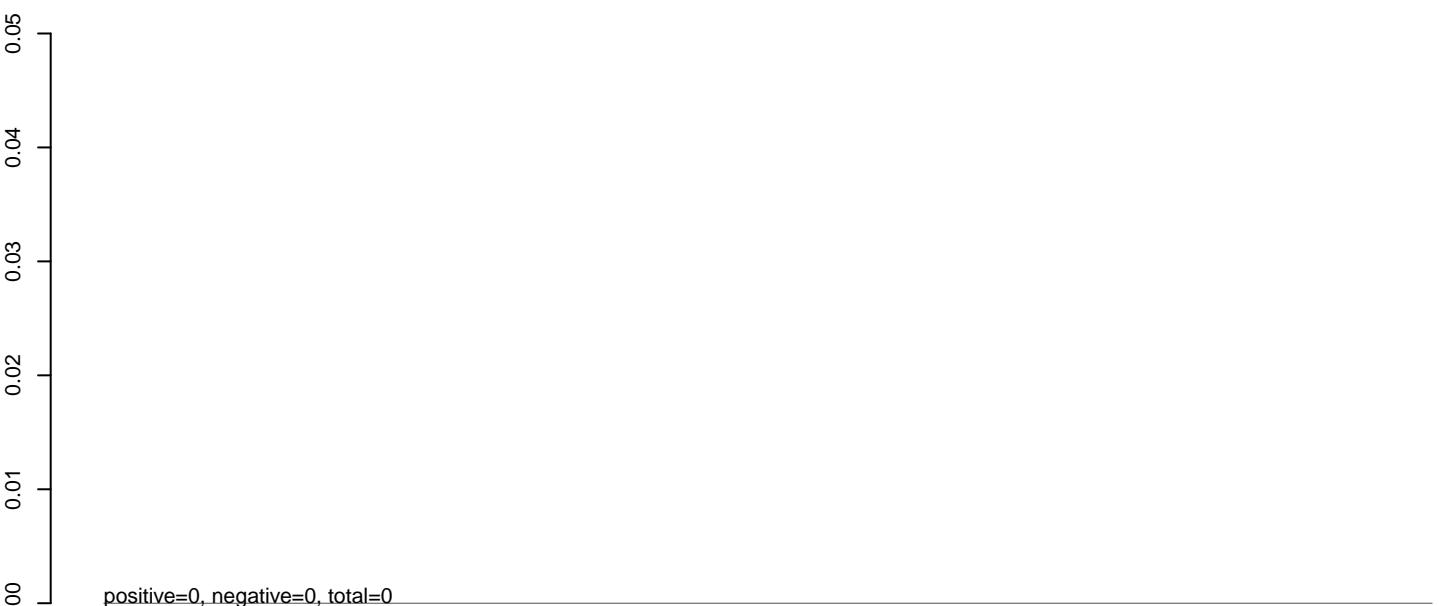
Window size=25, length=11634, virus@NC_018615.1-Eilat_virus_isolate_EO329:1-11634

0 2000 4000 6000 8000 10000 12000

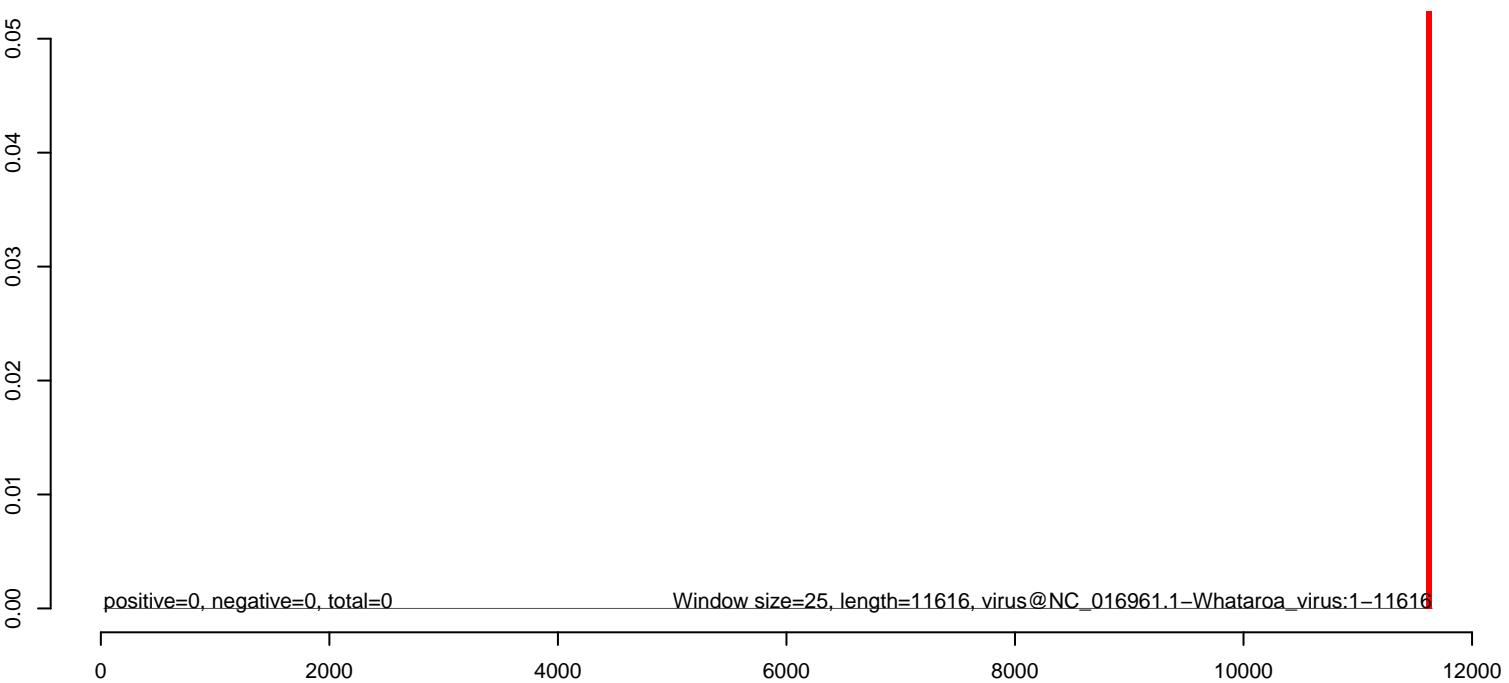
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



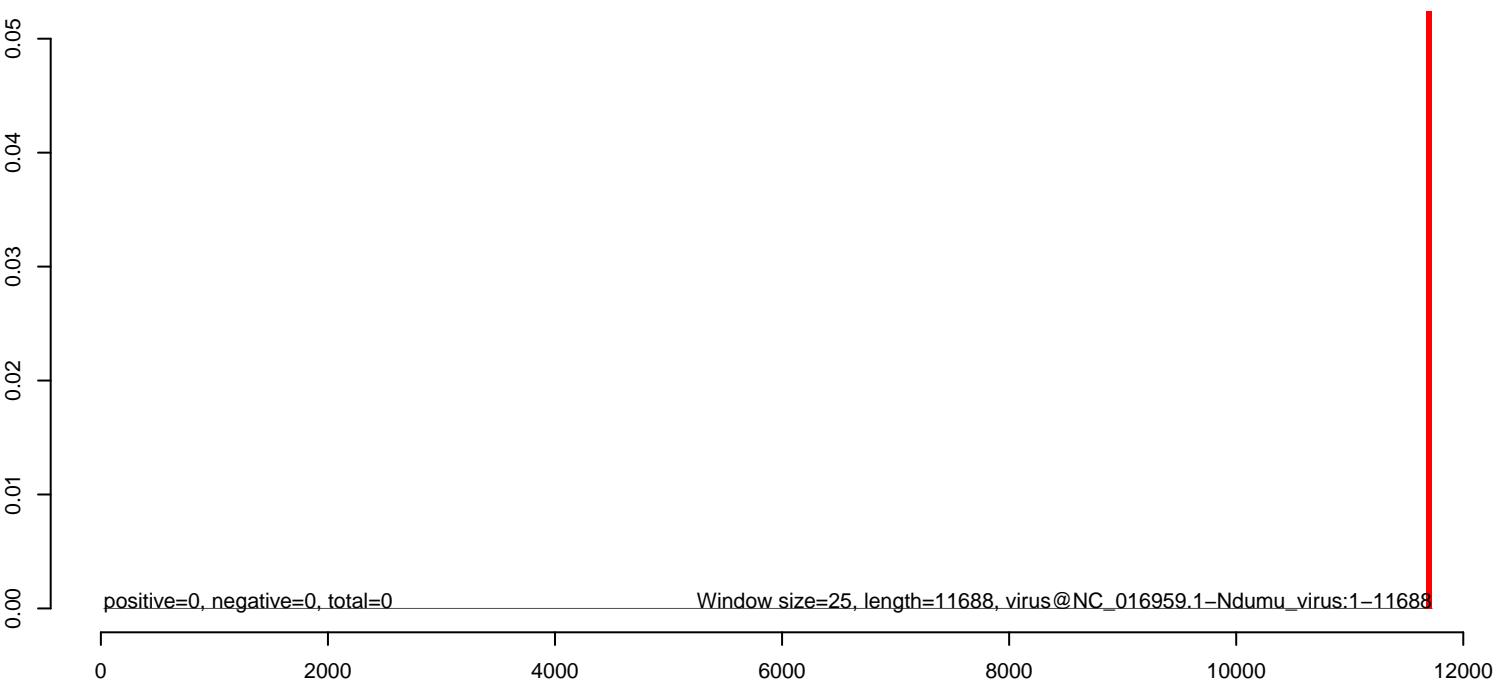
CuQuin_MG_7dpi3.18_23.rep



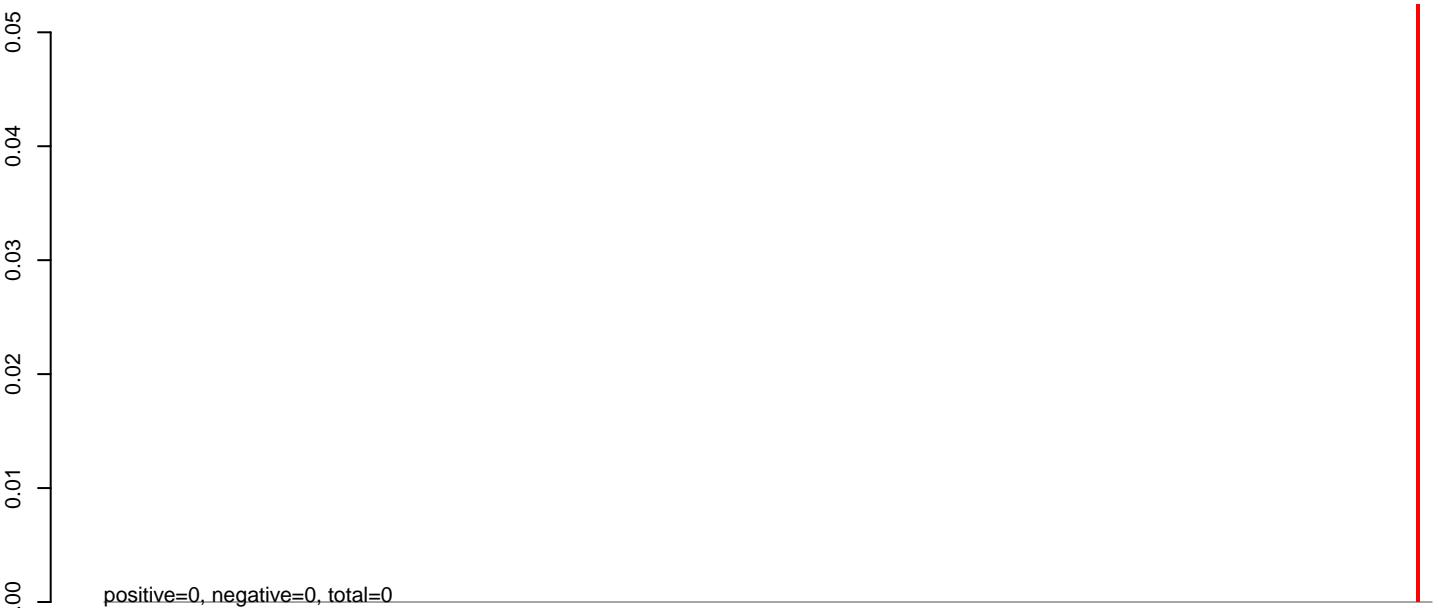
CuQuin_MG_7dpi3.24_35.rep



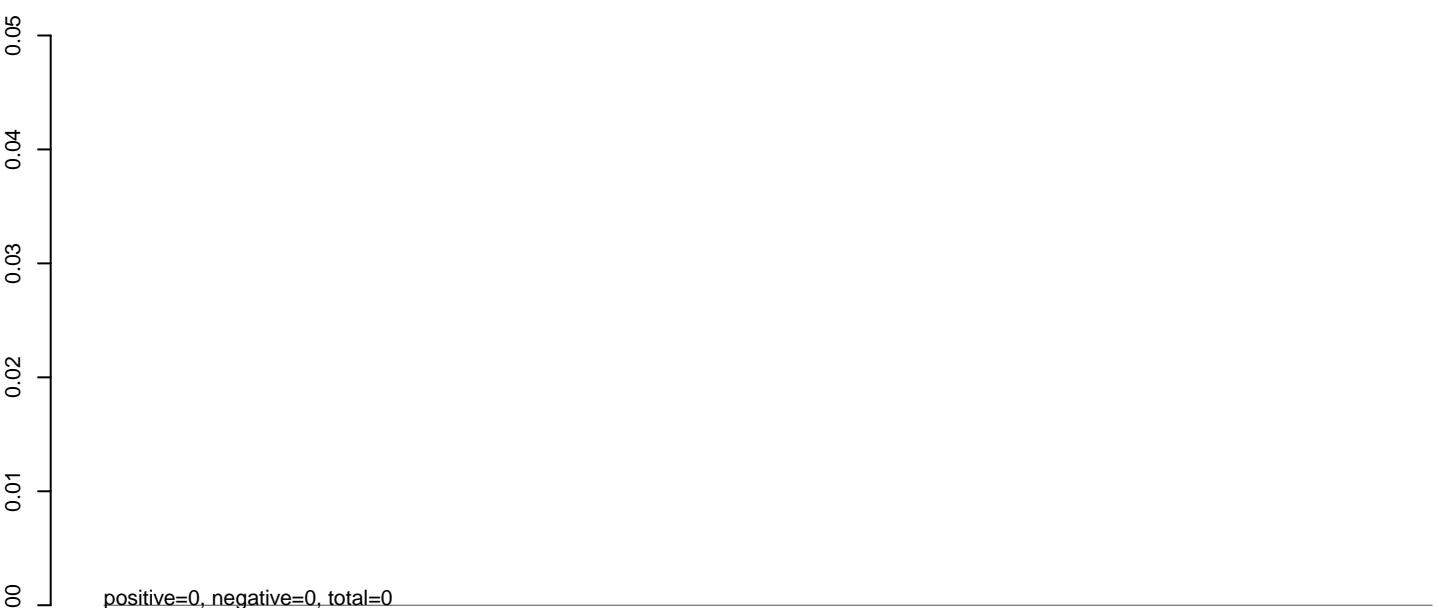
CuQuin_MG_7dpi3.rep



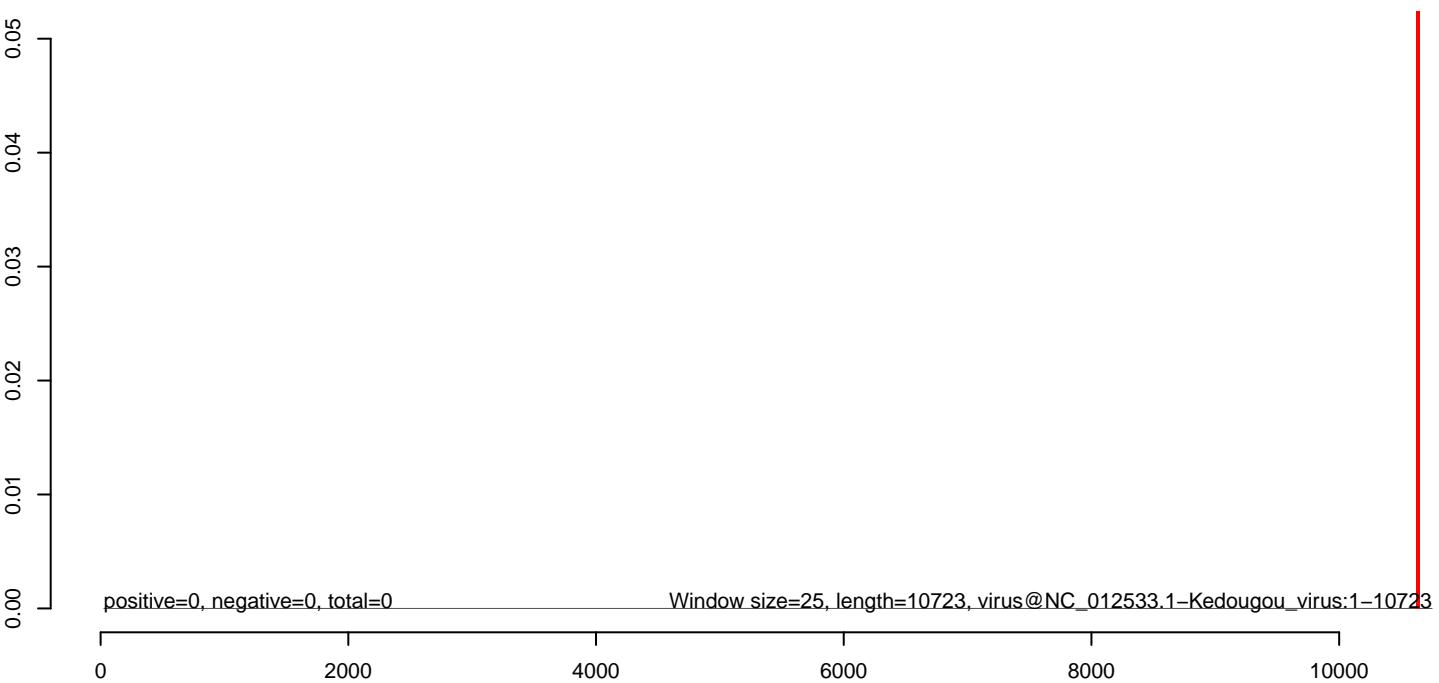
CuQuin_MG_7dpi3.18_23.rep



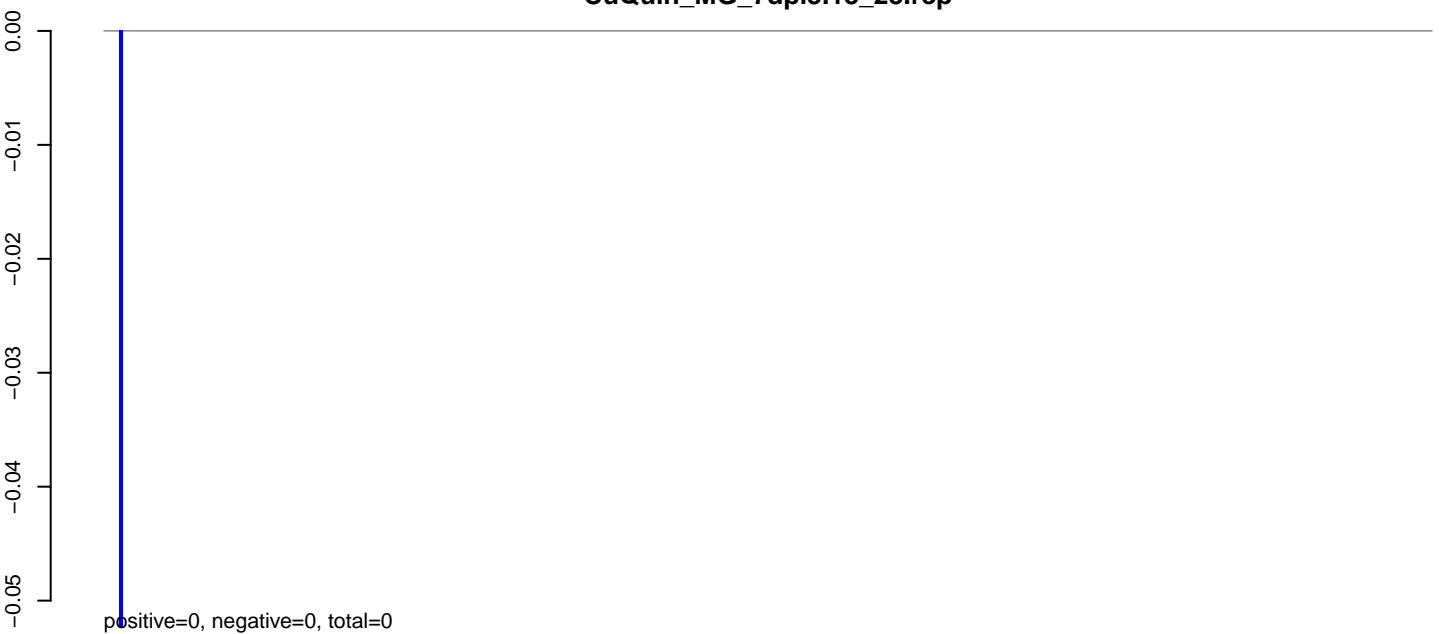
CuQuin_MG_7dpi3.24_35.rep



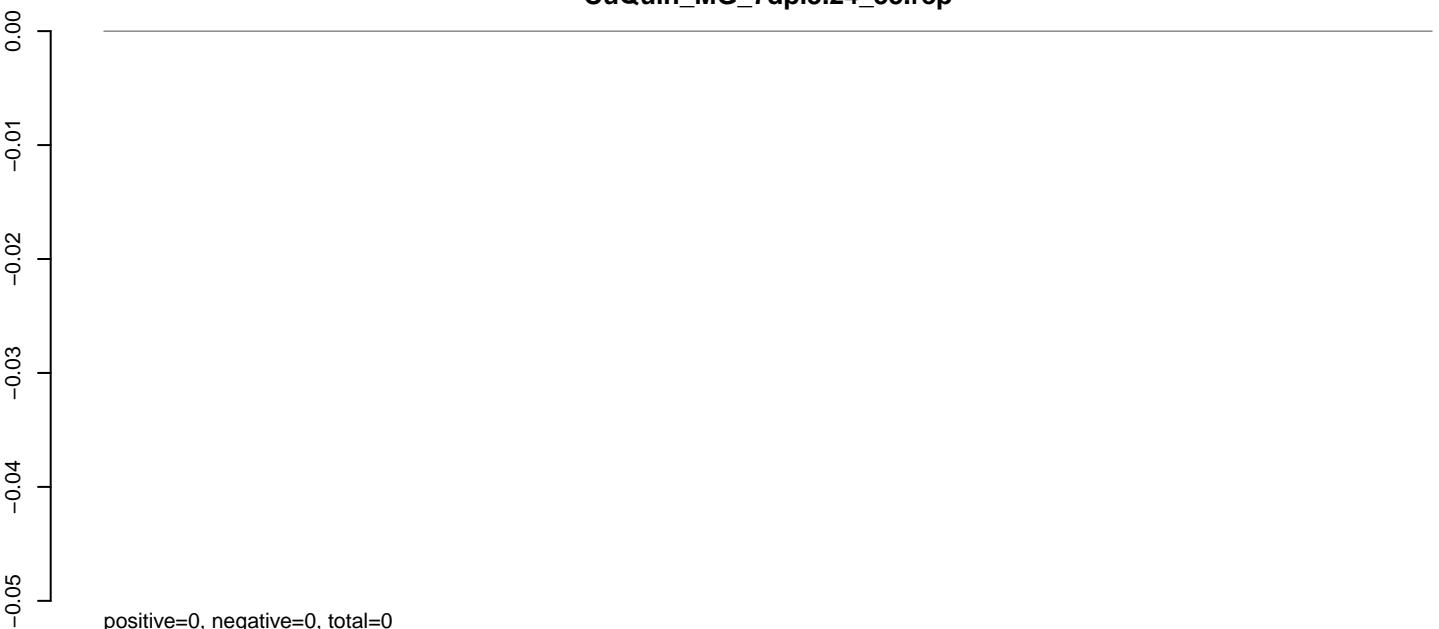
CuQuin_MG_7dpi3.rep



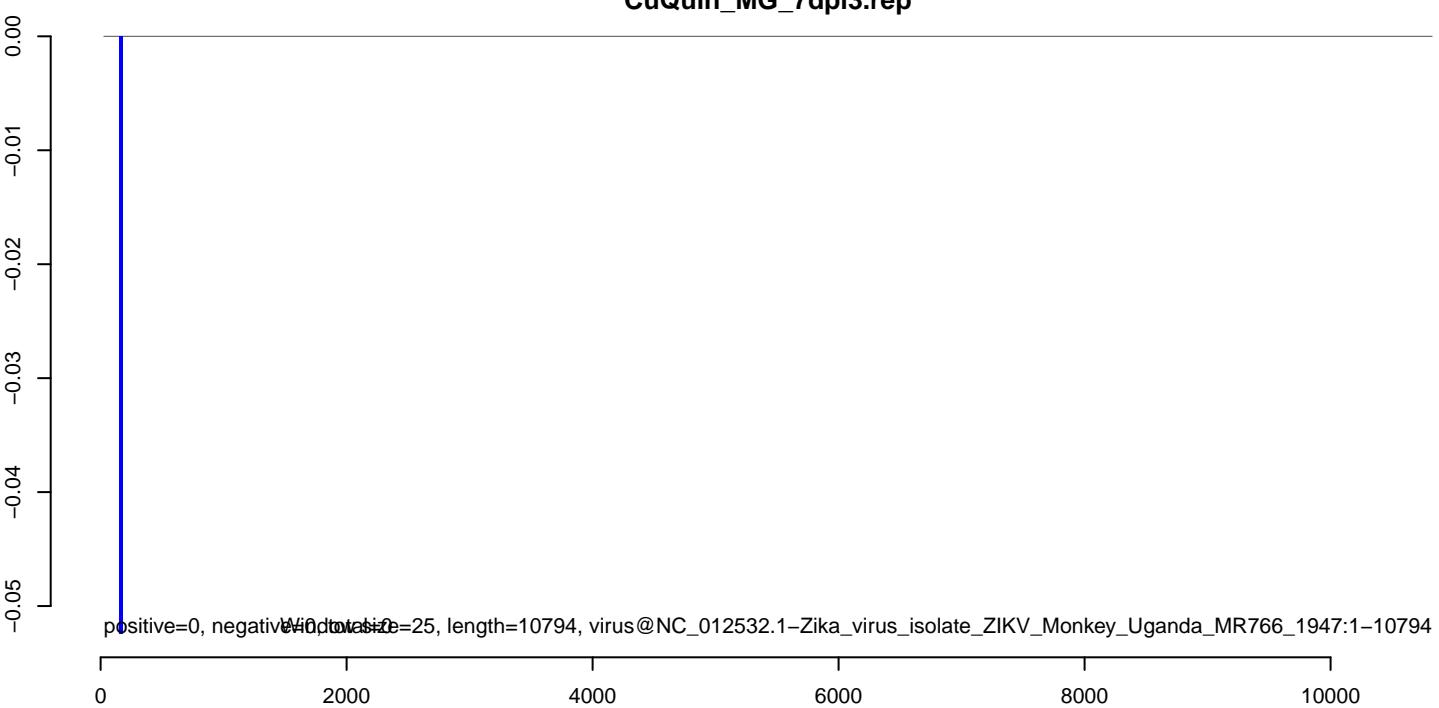
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



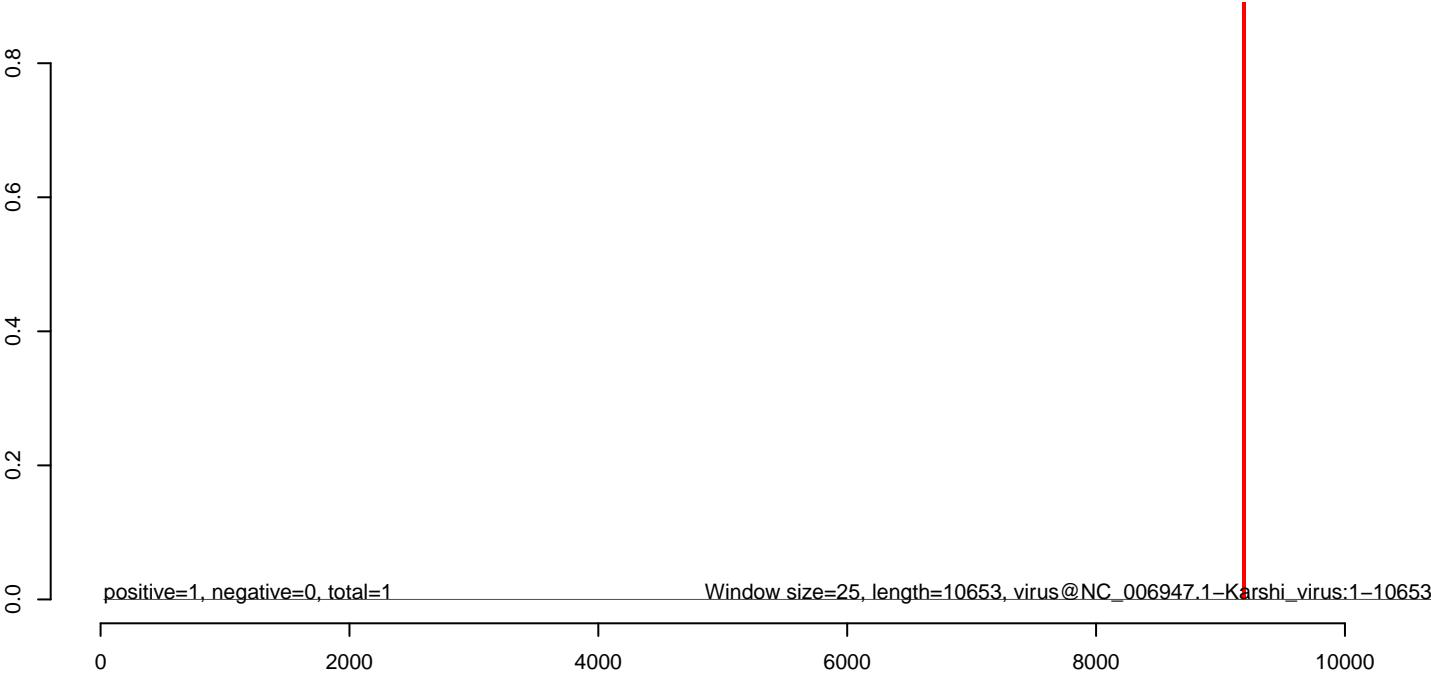
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

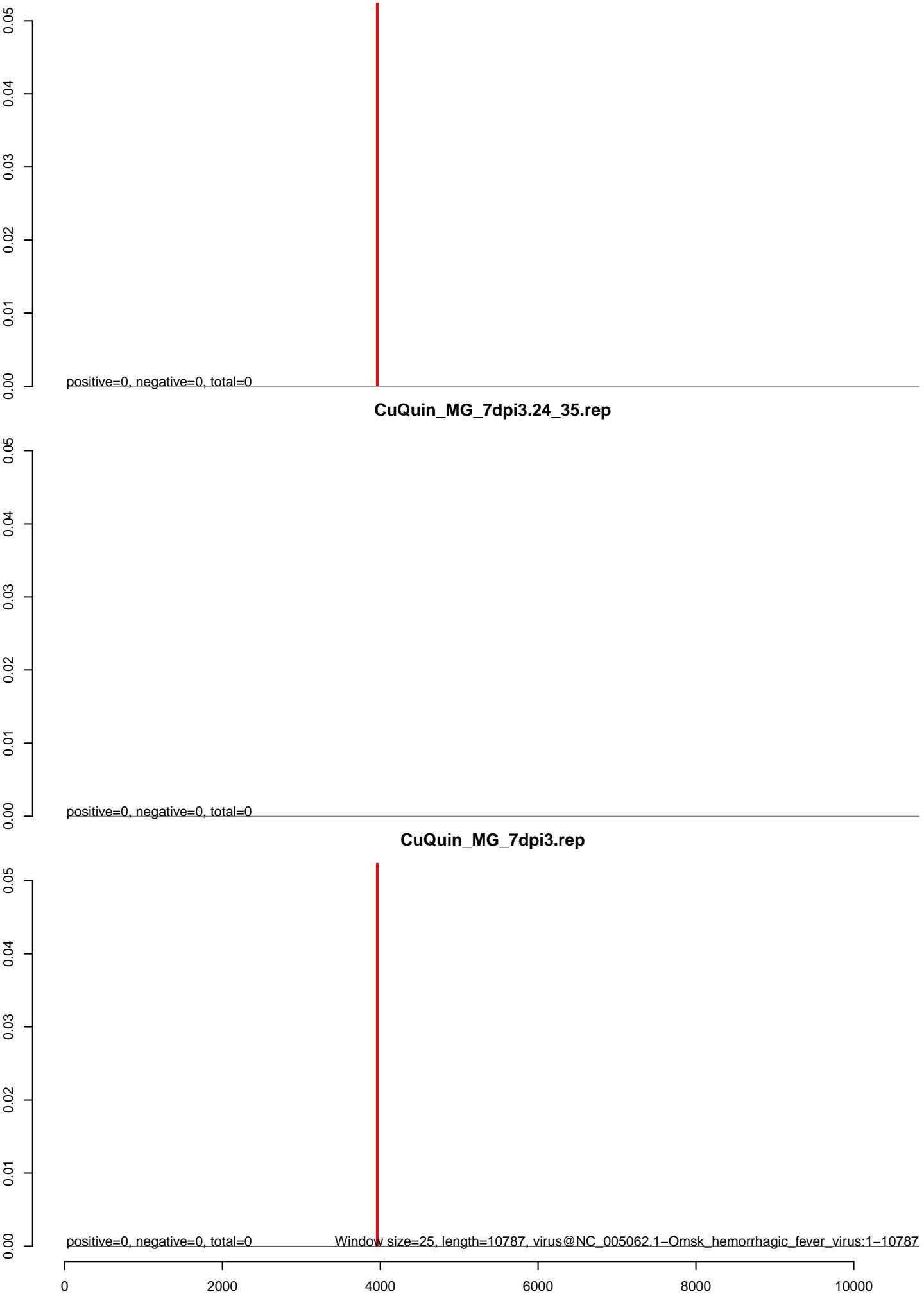
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

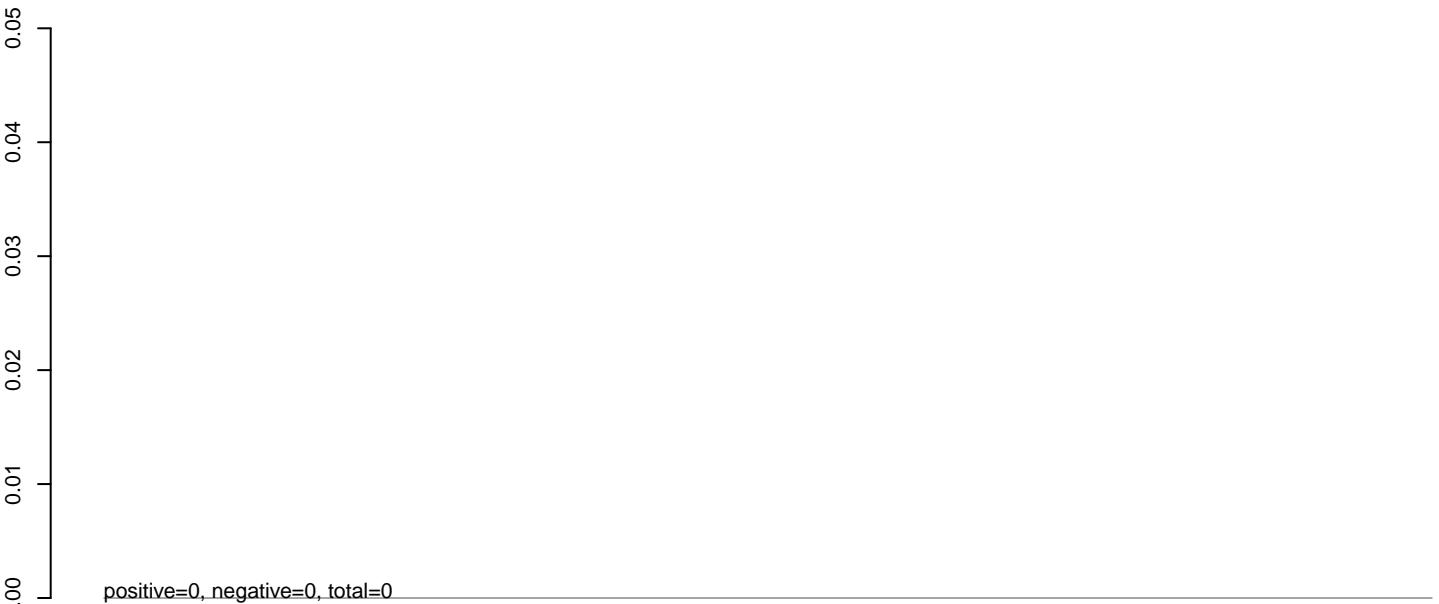
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=10787, virus@NC_005062.1-Omsk_hemorrhagic_fever_virus:1-10787



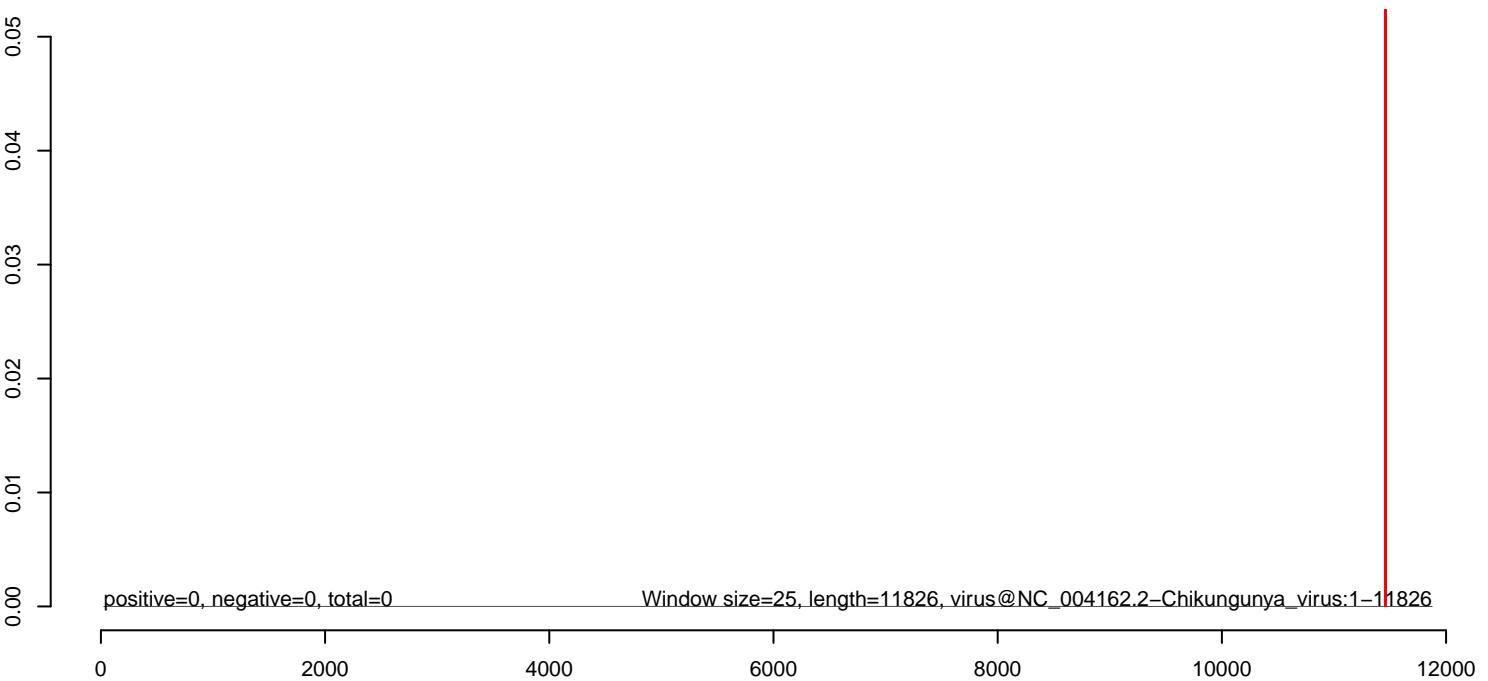
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



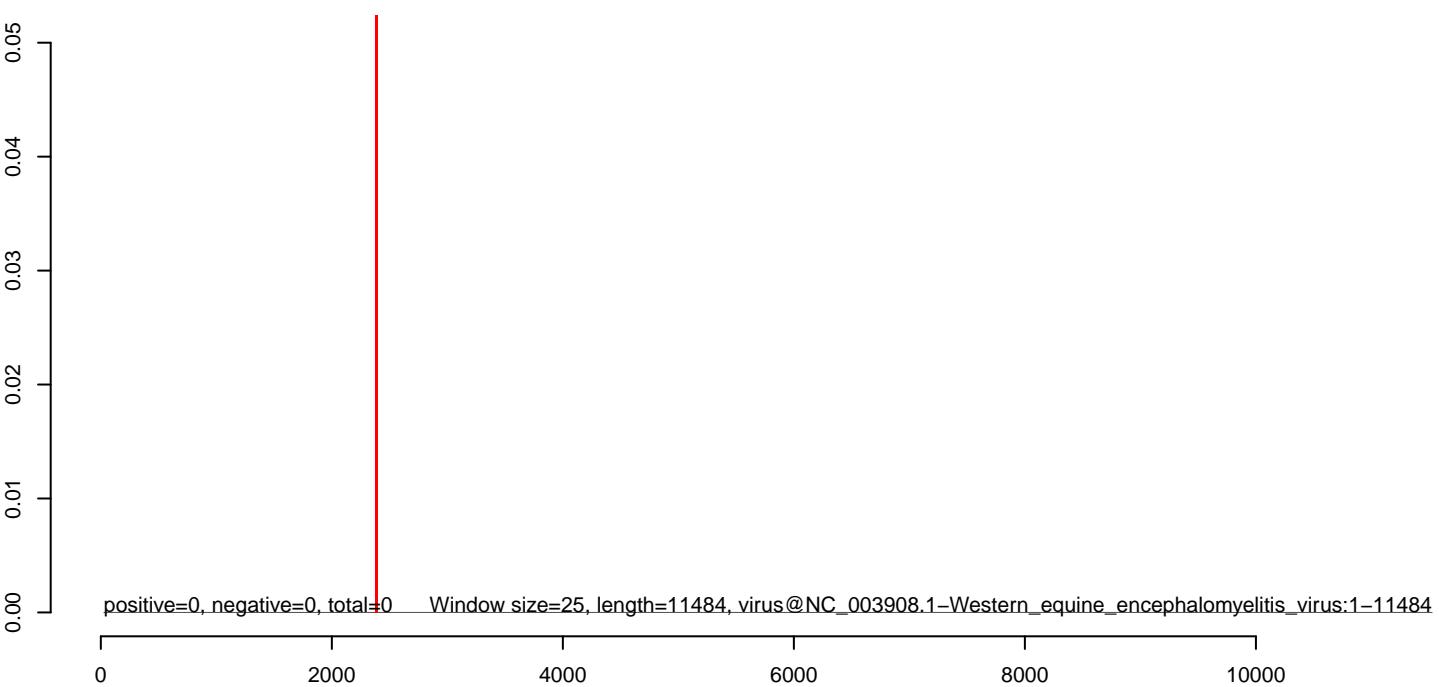
CuQuin_MG_7dpi3.18_23.rep



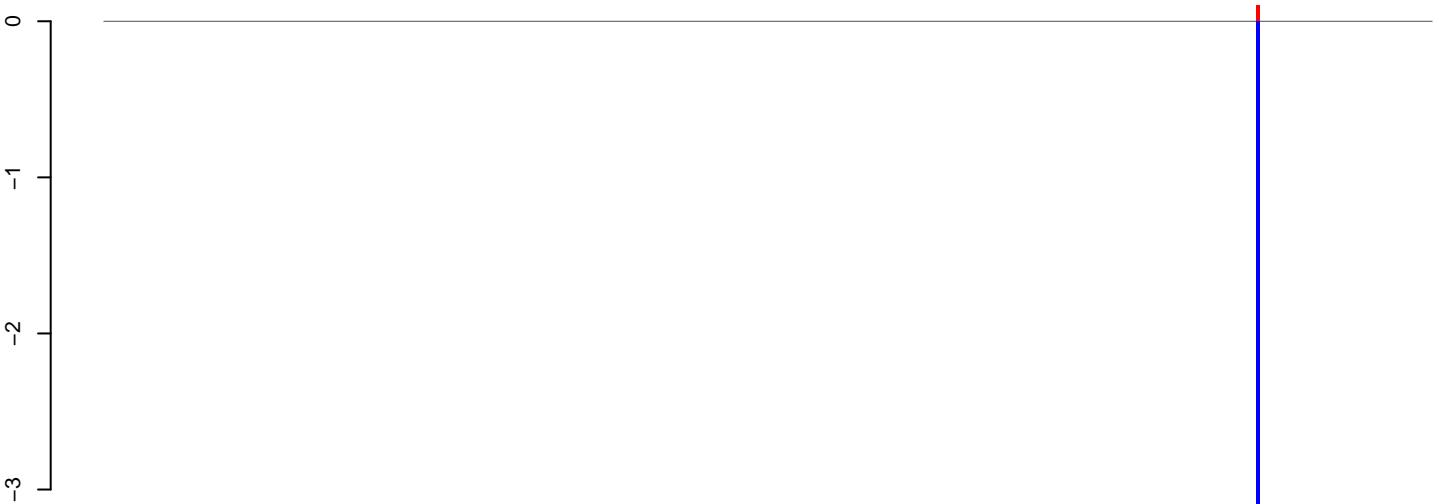
CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep

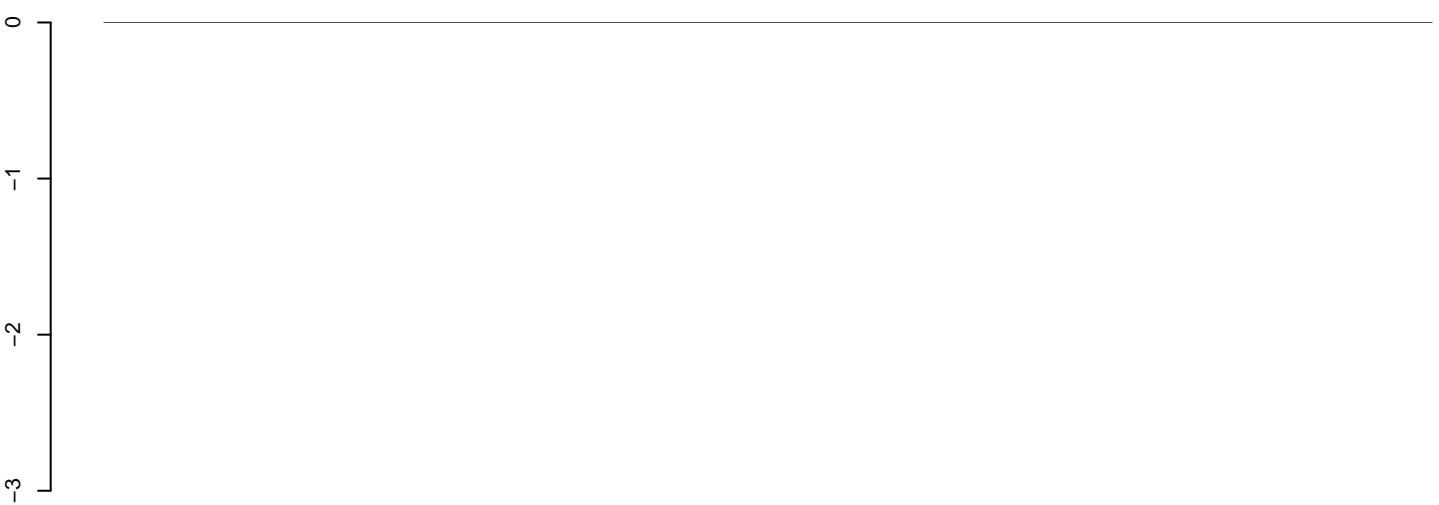


CuQuin_MG_7dpi3.18_23.rep



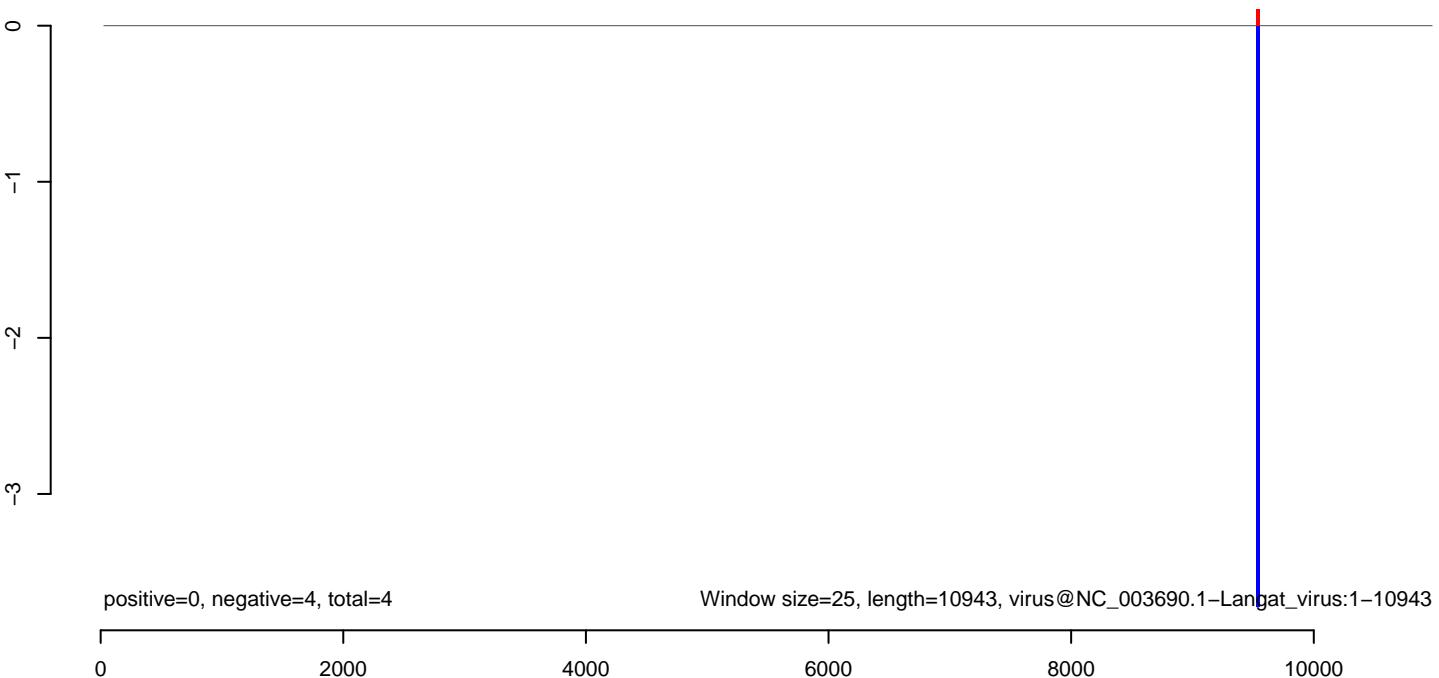
positive=0, negative=4, total=4

CuQuin_MG_7dpi3.24_35.rep



positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep



positive=0, negative=4, total=4

Window size=25, length=10943, virus@NC_003690.1-Langat_virus:1-10943

0 2000 4000 6000 8000 10000

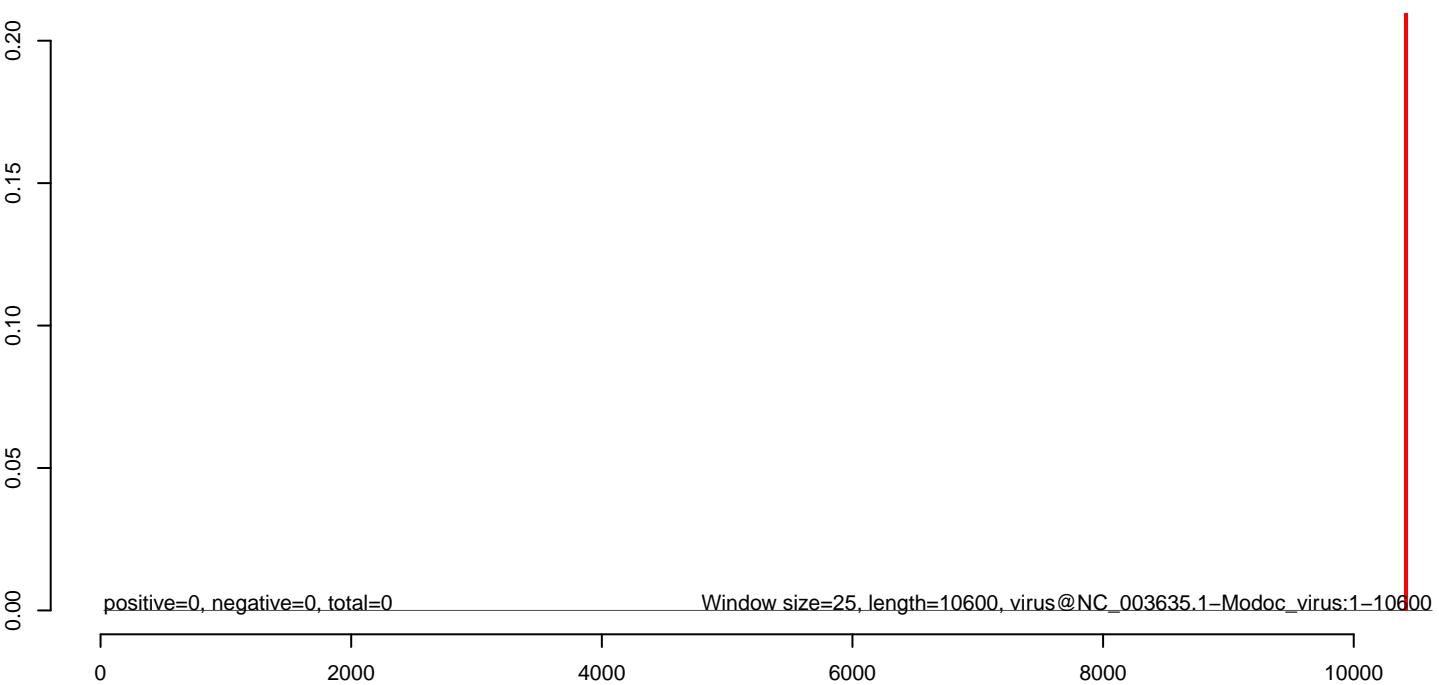
CuQuin_MG_7dpi3.18_23.rep



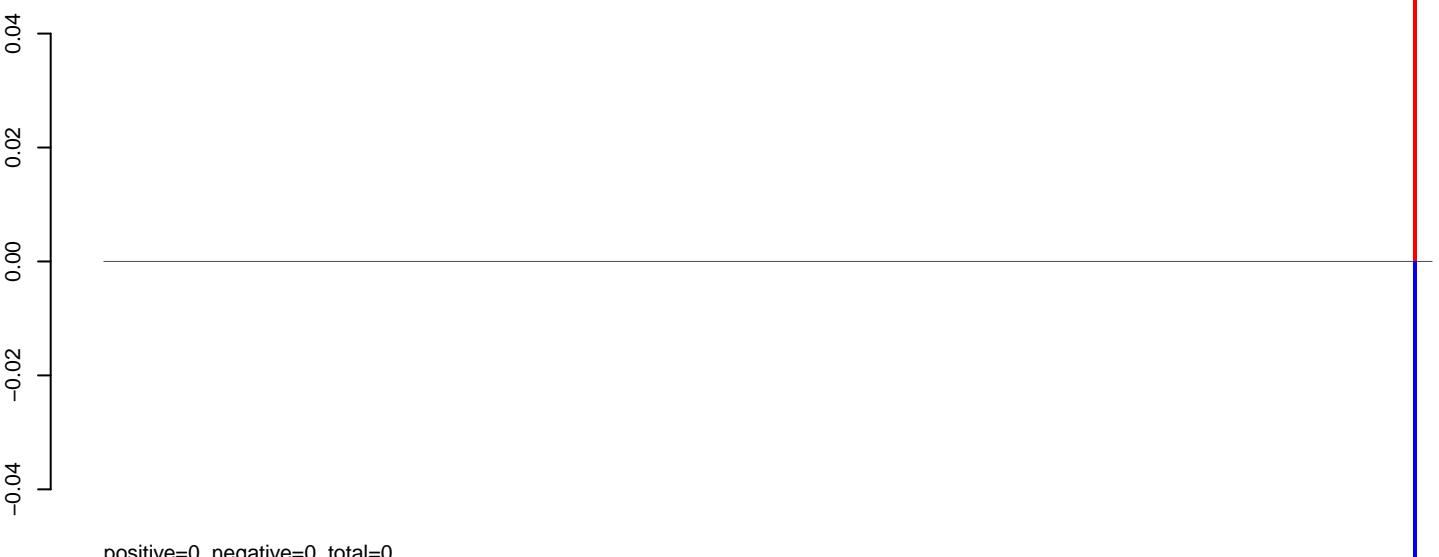
CuQuin_MG_7dpi3.24_35.rep



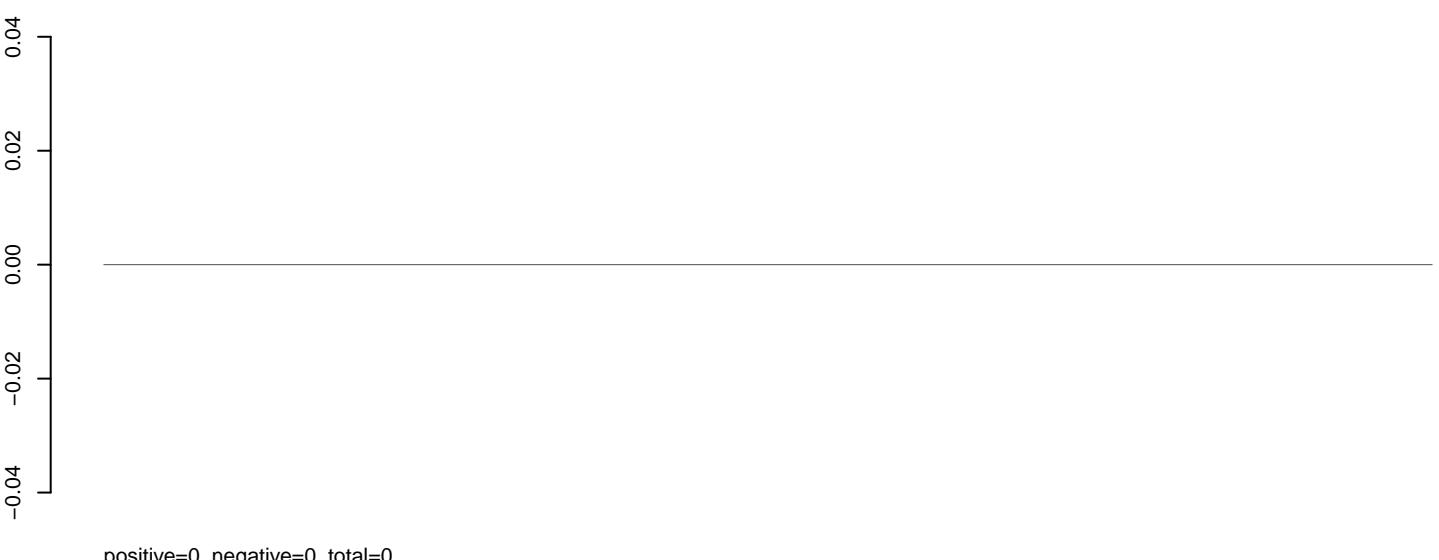
CuQuin_MG_7dpi3.rep



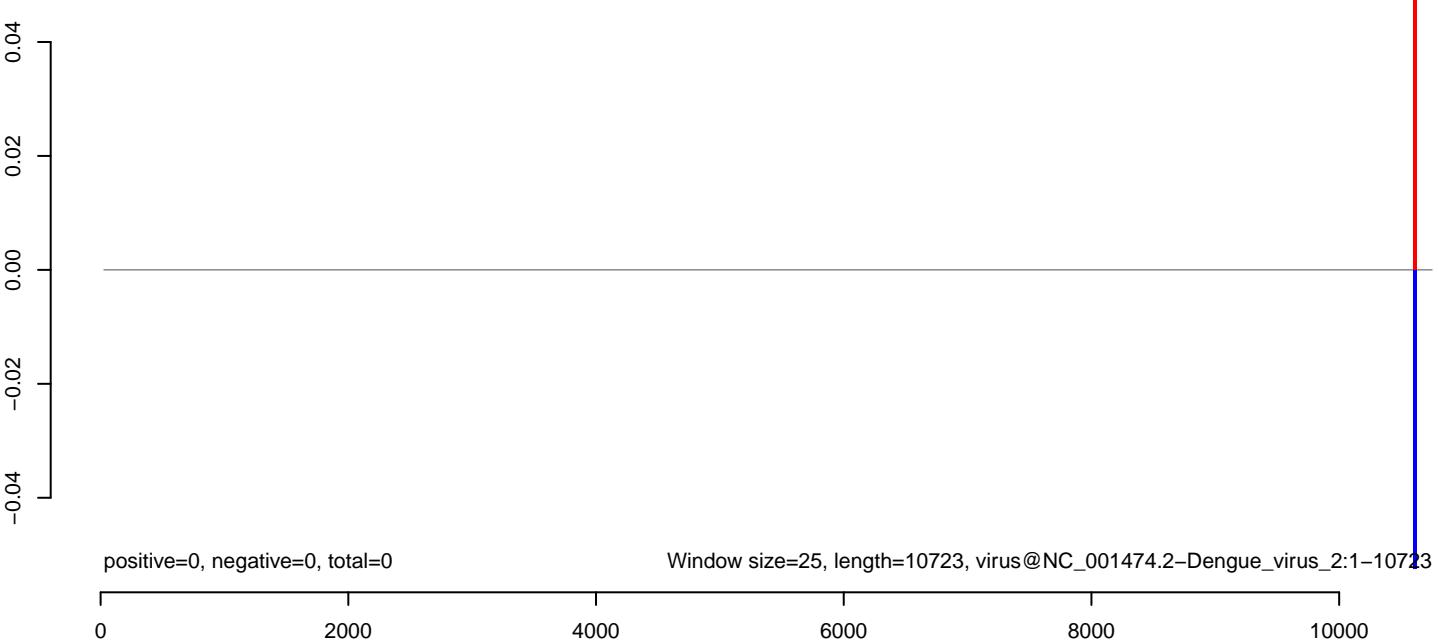
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0



CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0



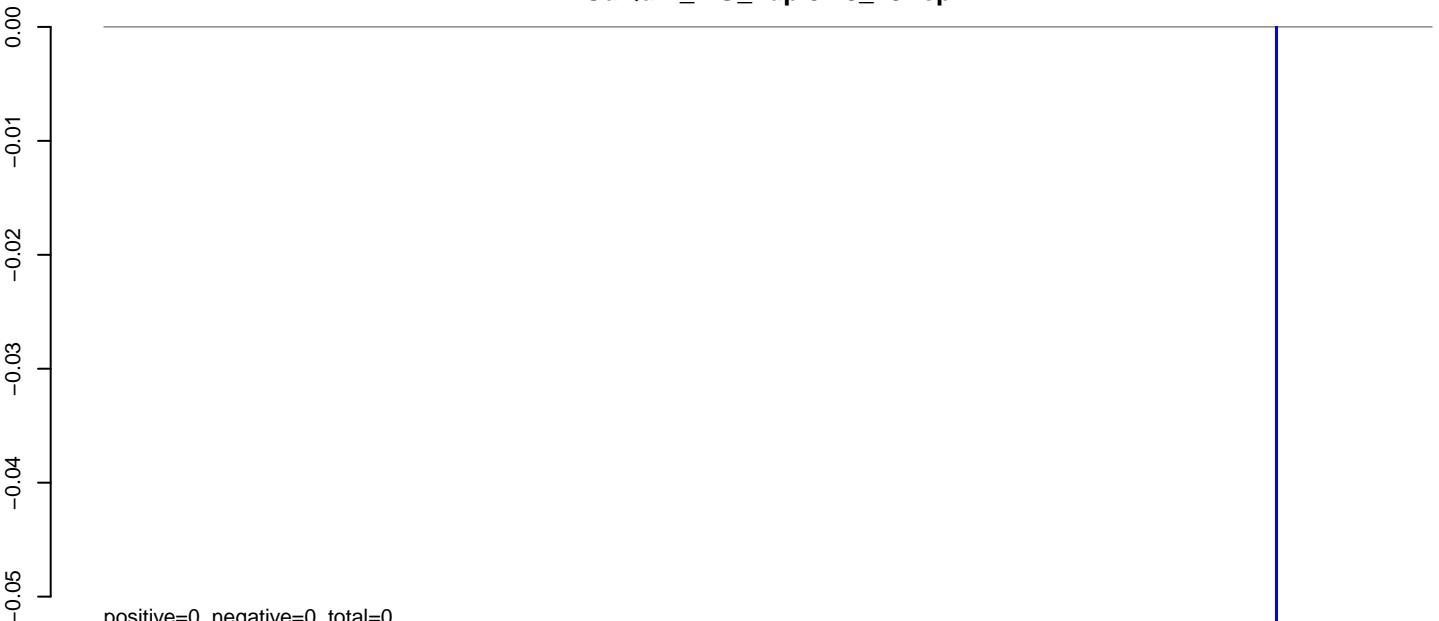
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=6065, virus@MH188054 1-Culex_Y_virus_strain_CYV_Kern_partial_genome:1-6065

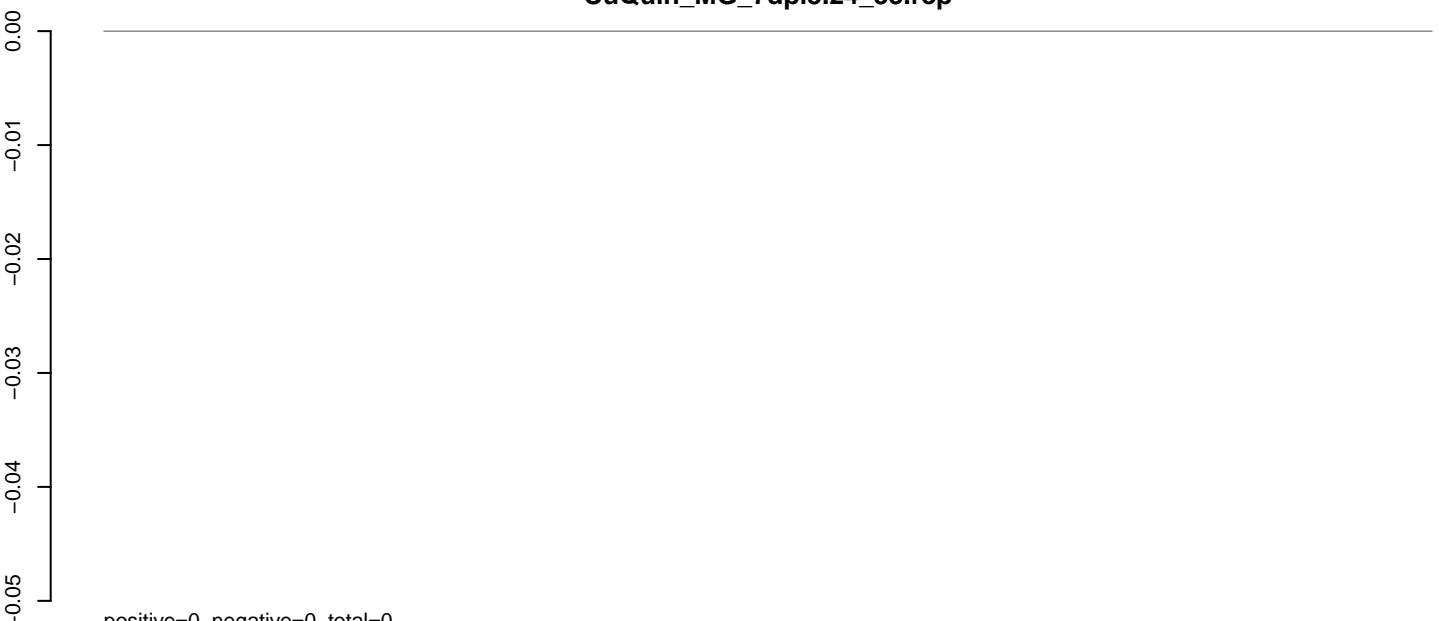


0 1000 2000 3000 4000 5000 6000

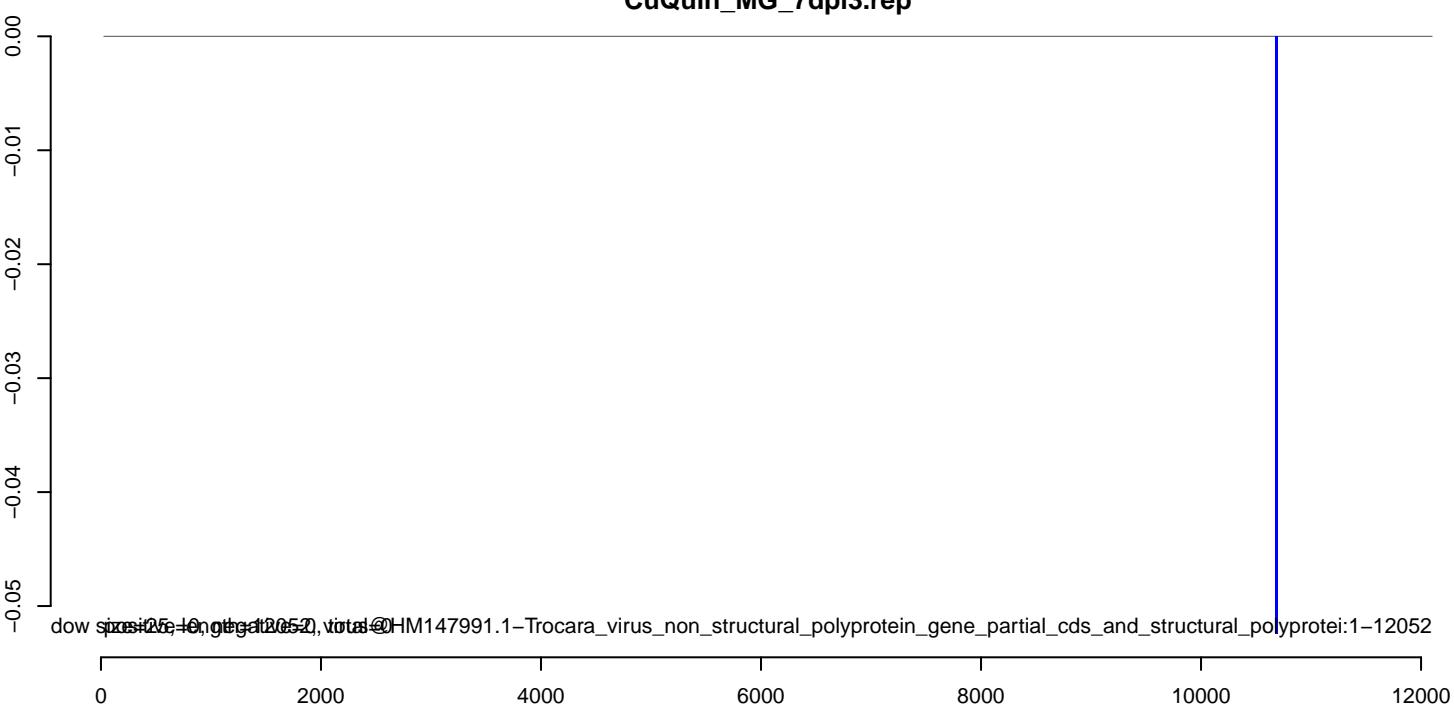
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

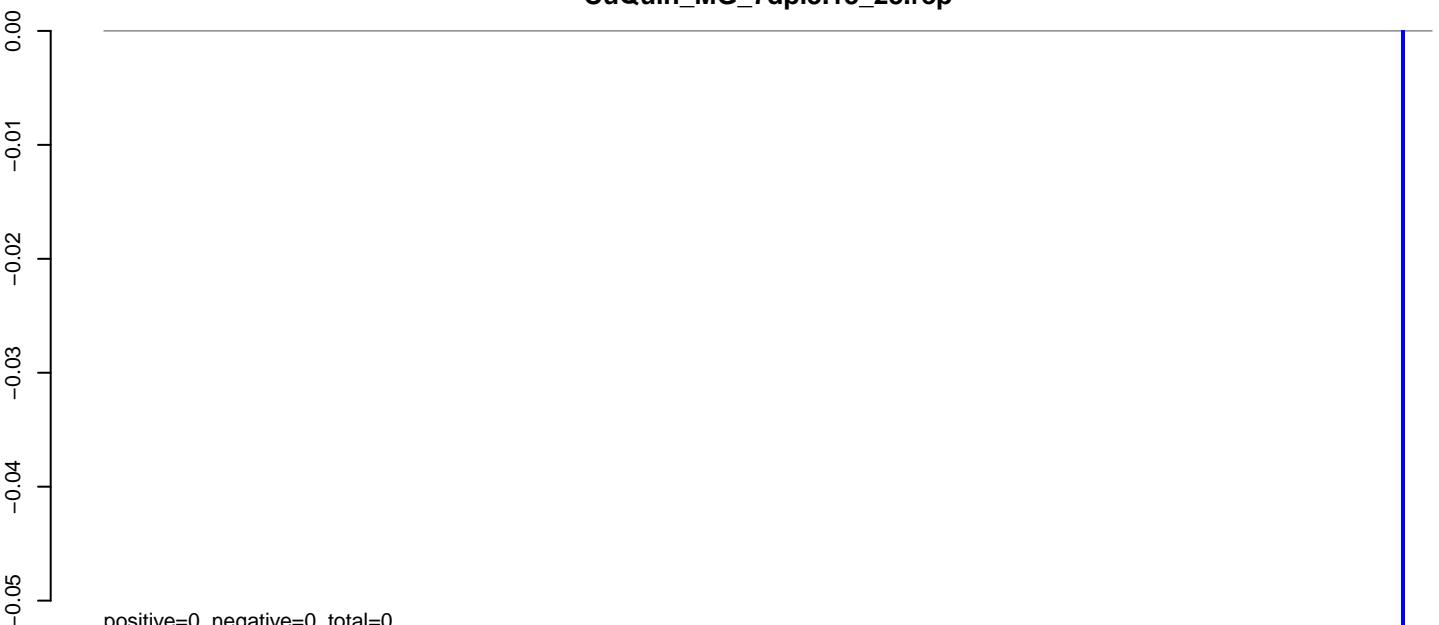
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

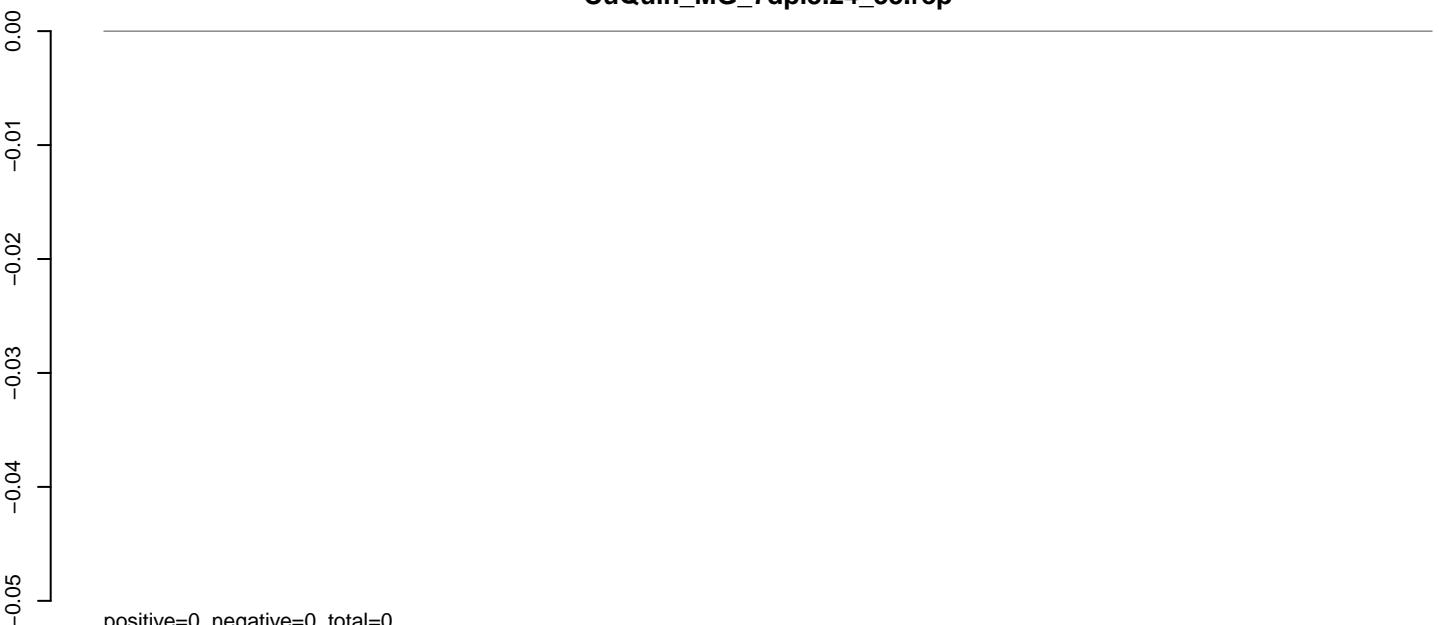
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0, size=25, length=10254, virus@DQ235149.1-Royal_Farm_virus_from_Afghanistan_polyprotein_gene:1-10254

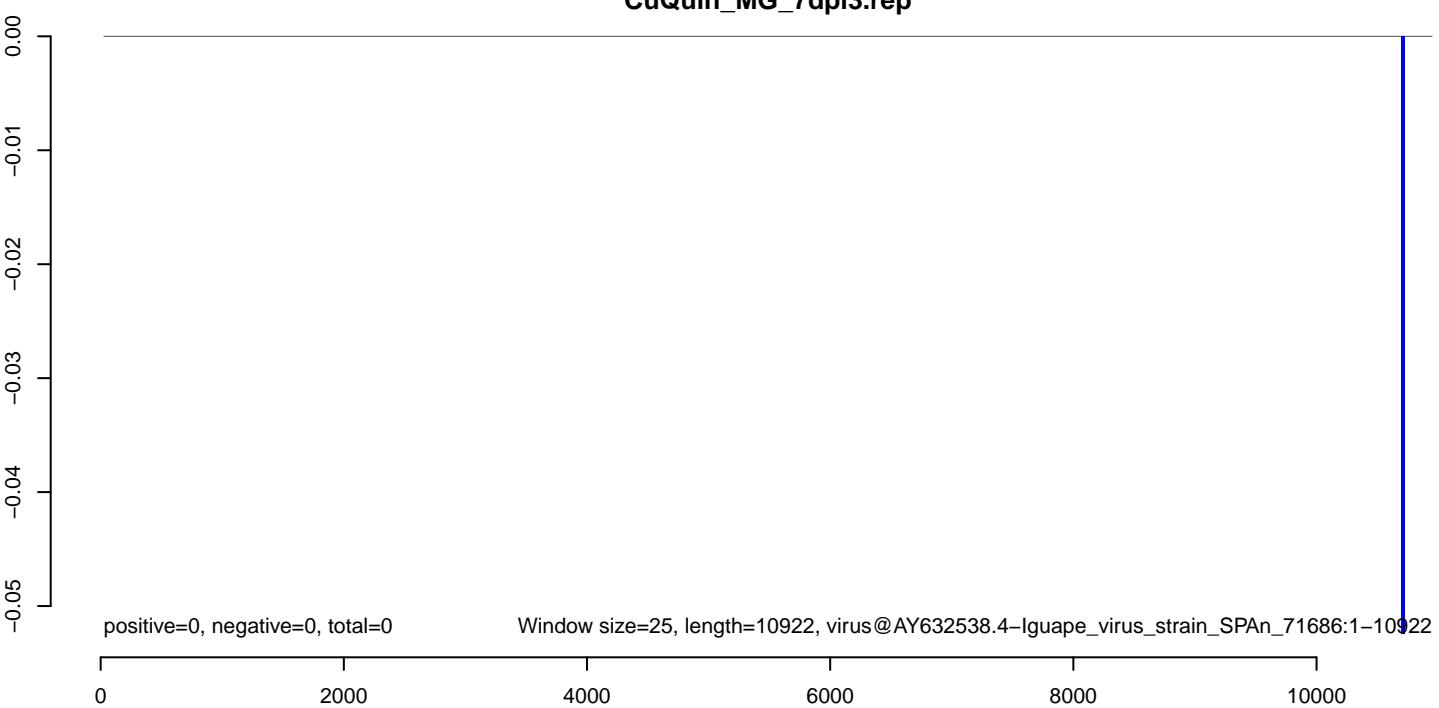
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0



CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

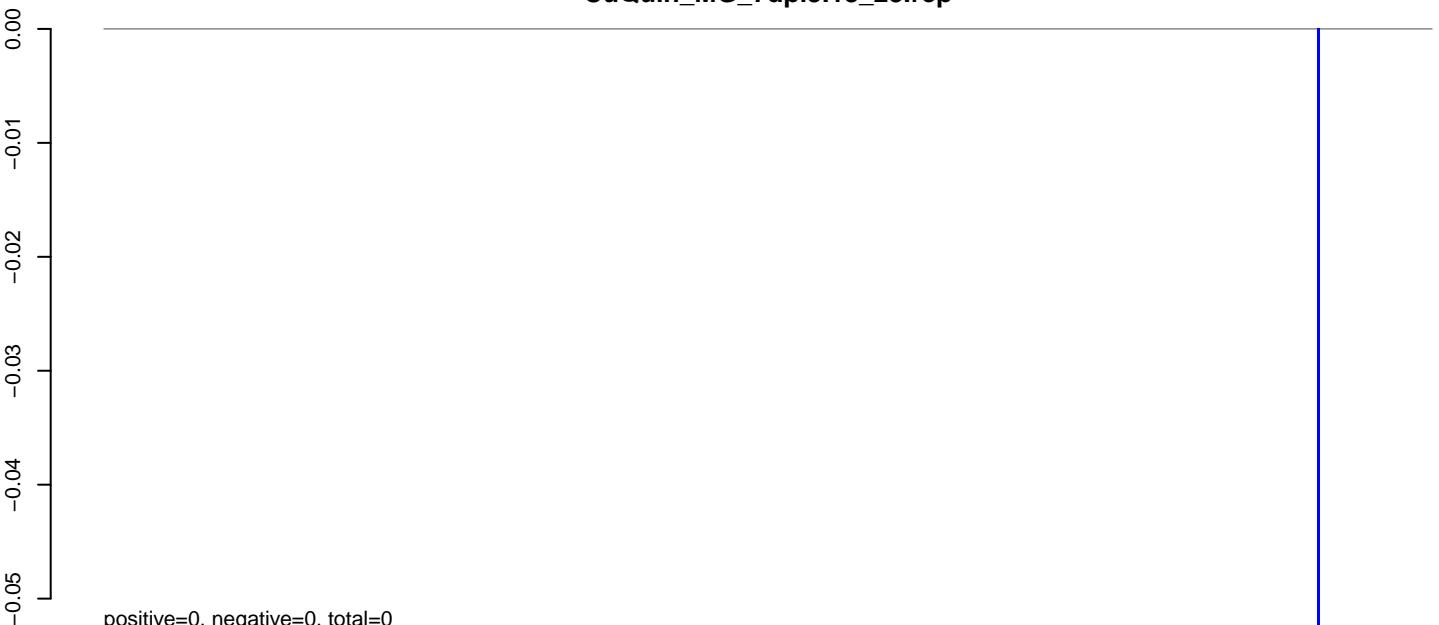


CuQuin_MG_7dpi3.rep

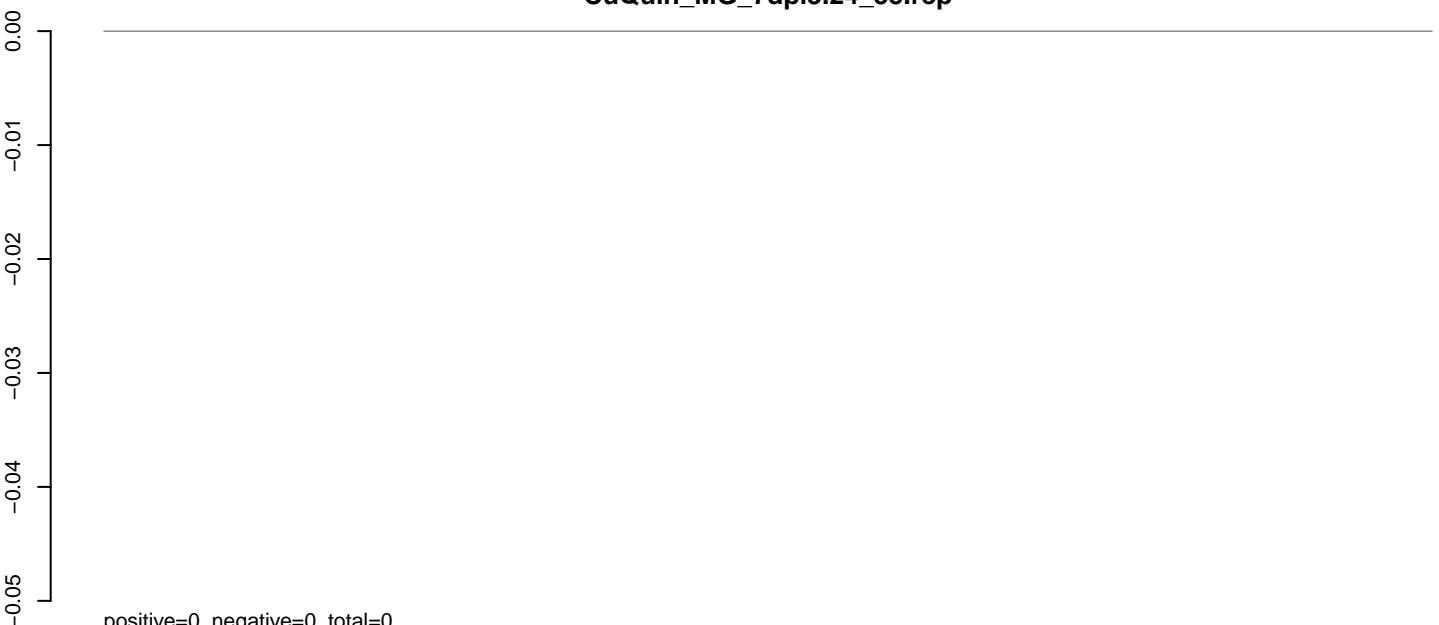
positive=0, negative=0, total=20, length=972, virus@AF411844.1-Elaeis_guineensis_MADS_box_transcription_factor_AGL2-2_mRNA1-972



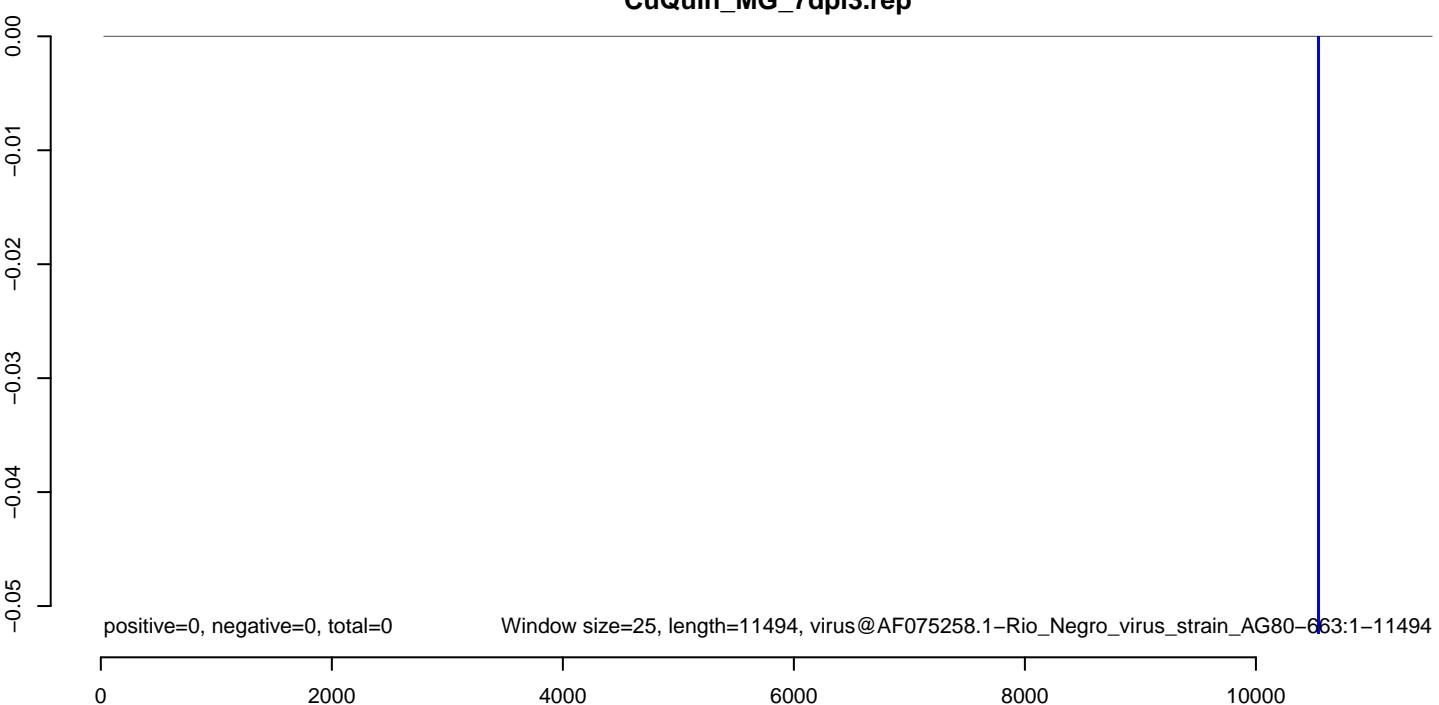
CuQuin_MG_7dpi3.18_23.rep



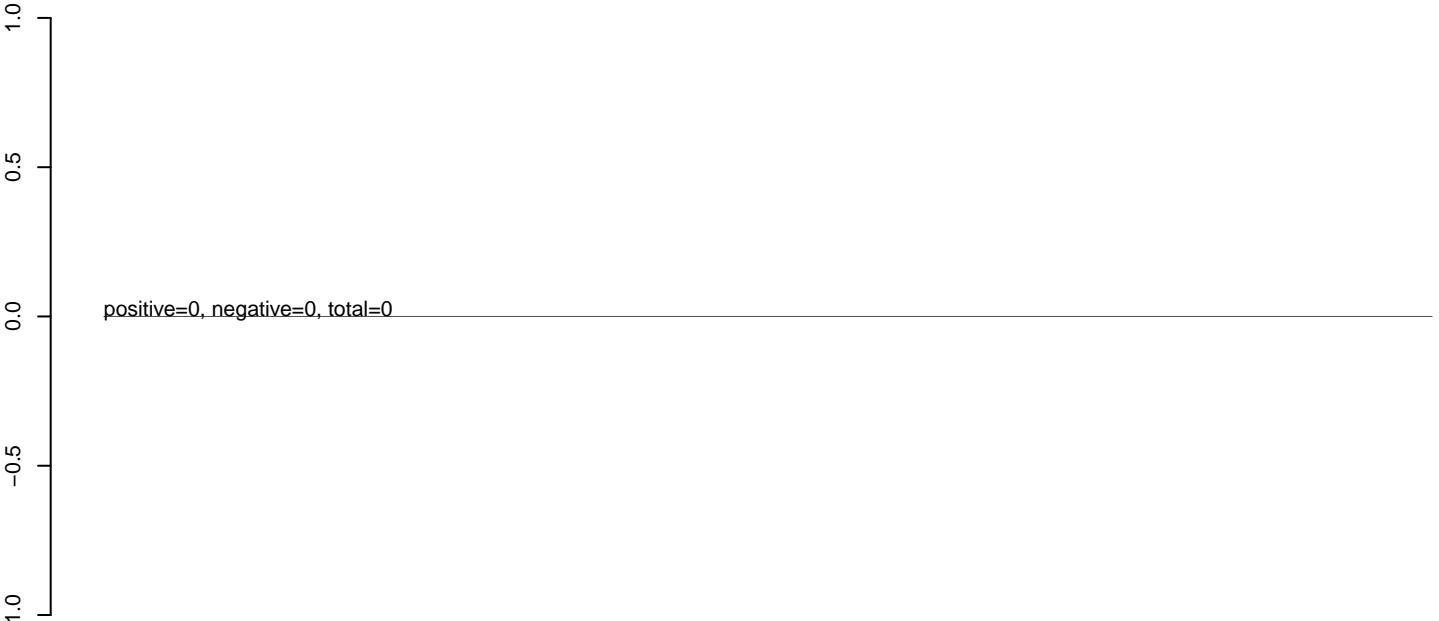
CuQuin_MG_7dpi3.24_35.rep



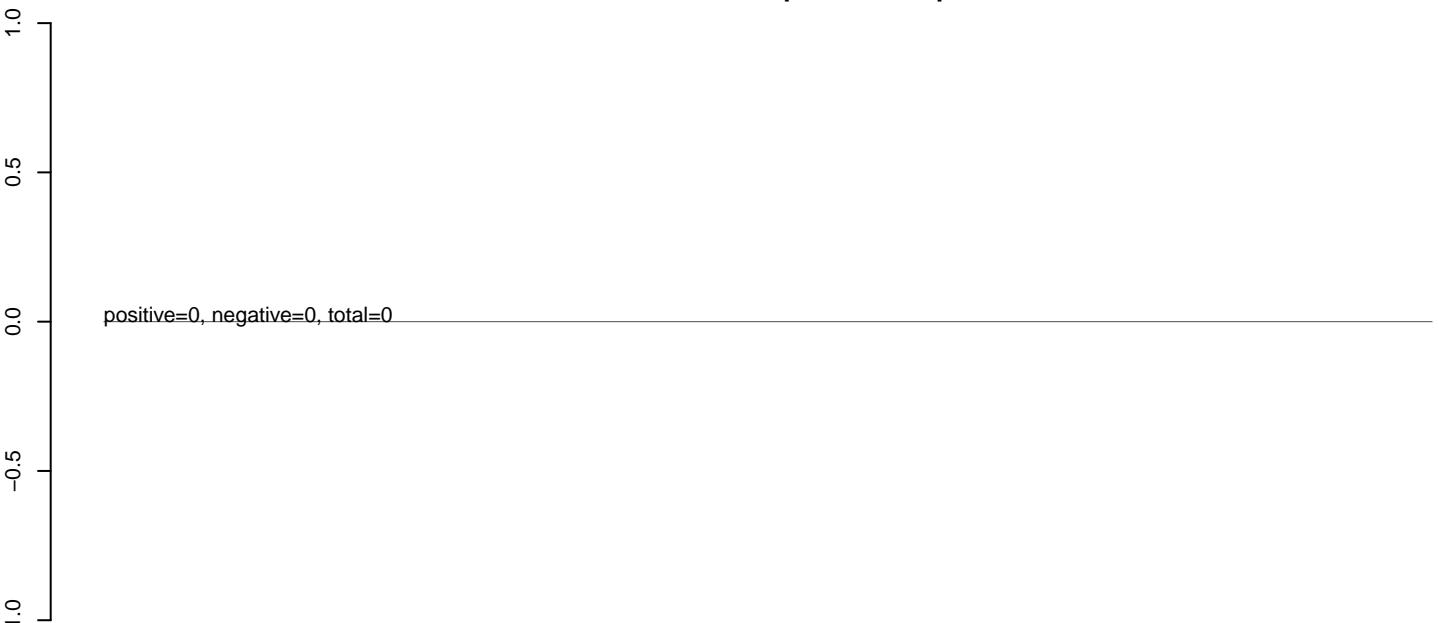
CuQuin_MG_7dpi3.rep



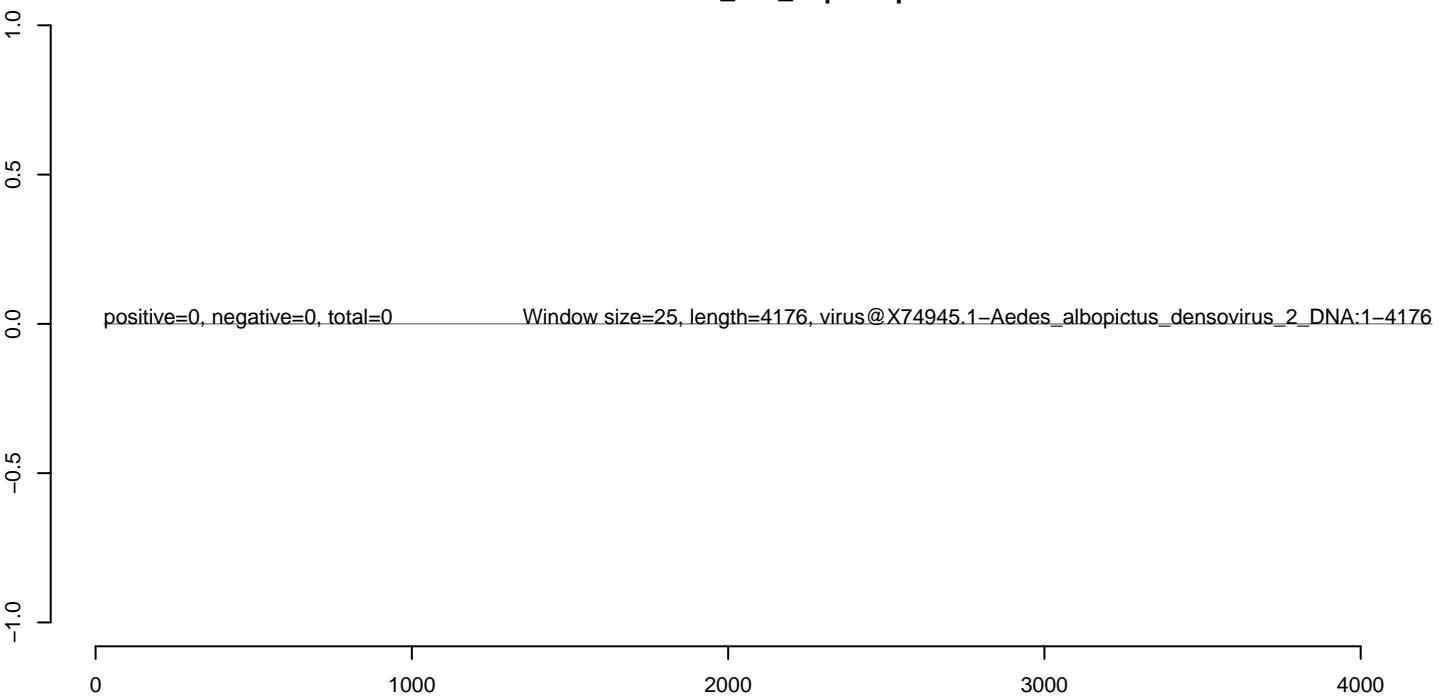
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

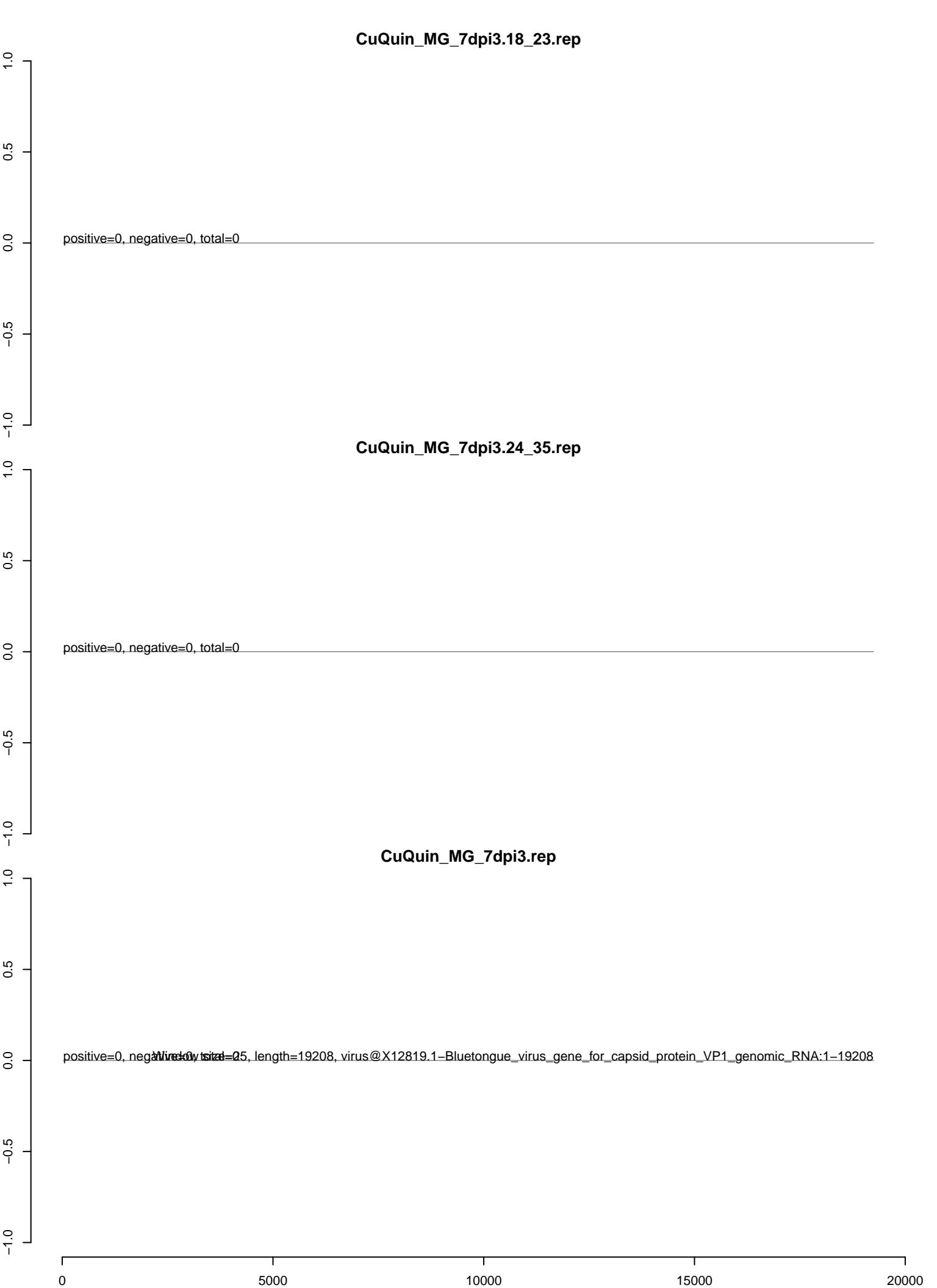
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

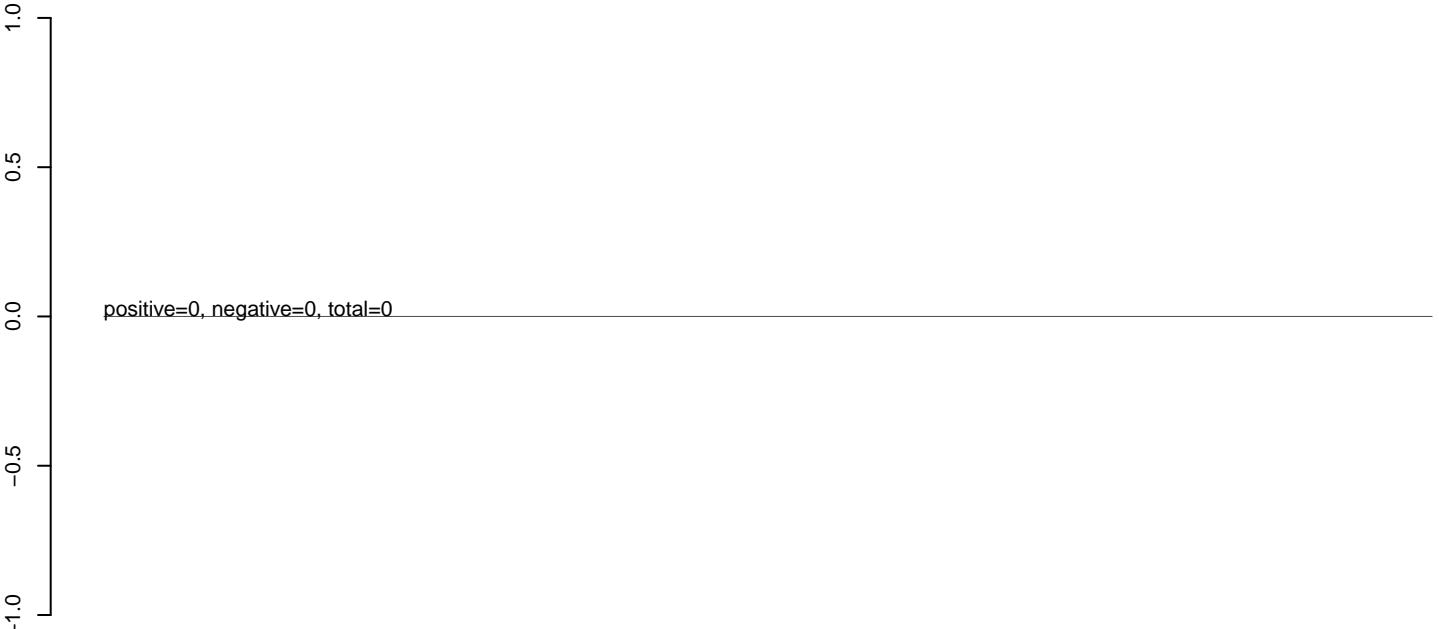
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

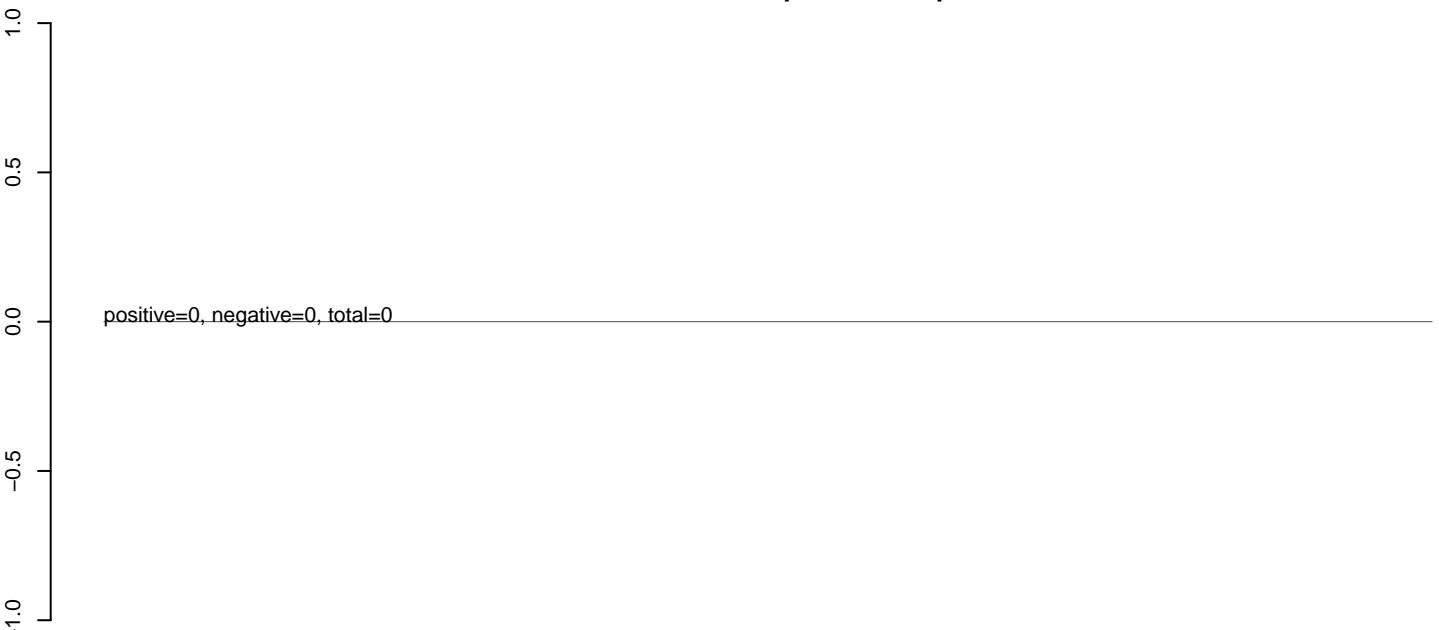
positive=0, negative=0, total=0, virus@X12819.1-Bluetongue_virus_gene_for_capsid_protein_VP1_genomic_RNA:1-19208



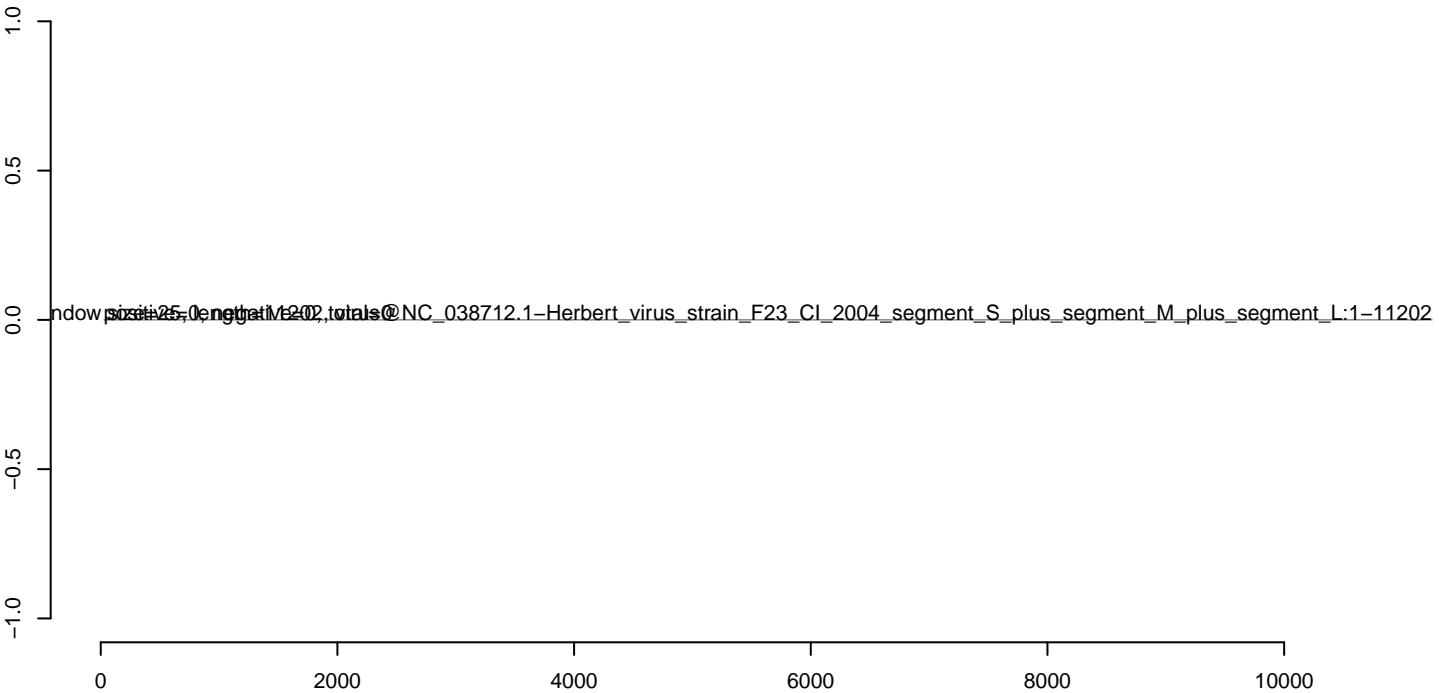
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

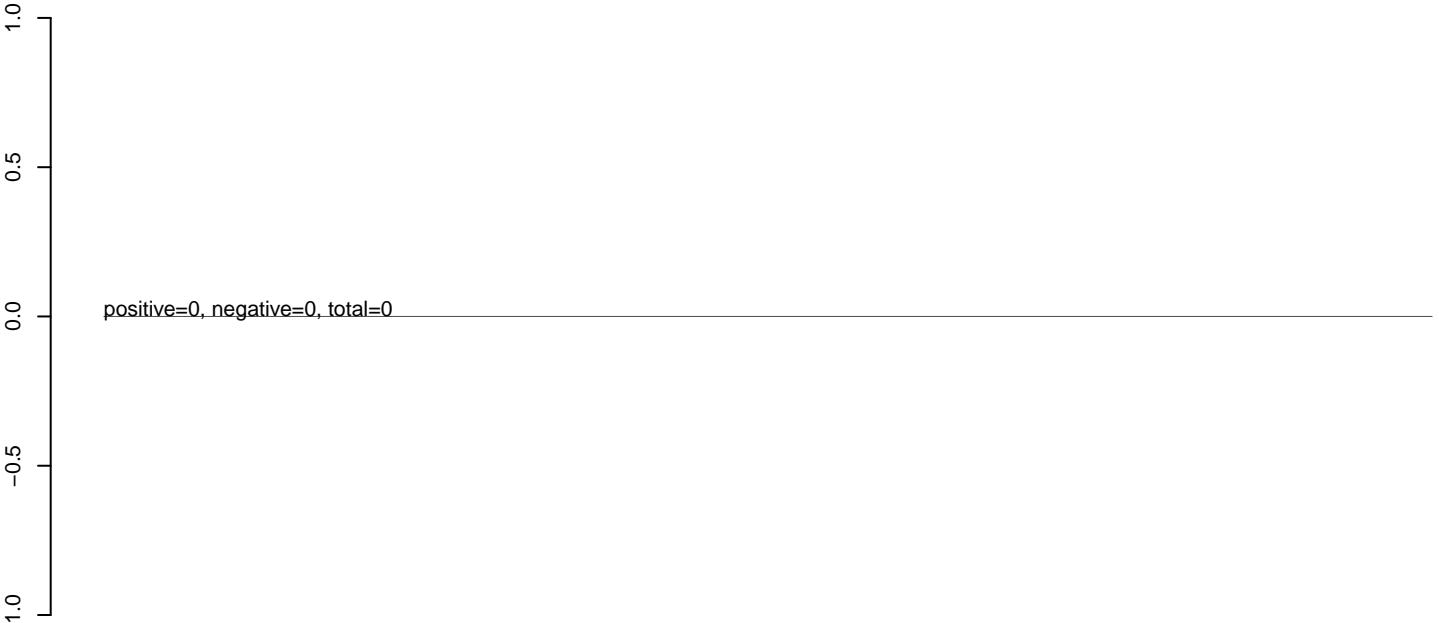
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

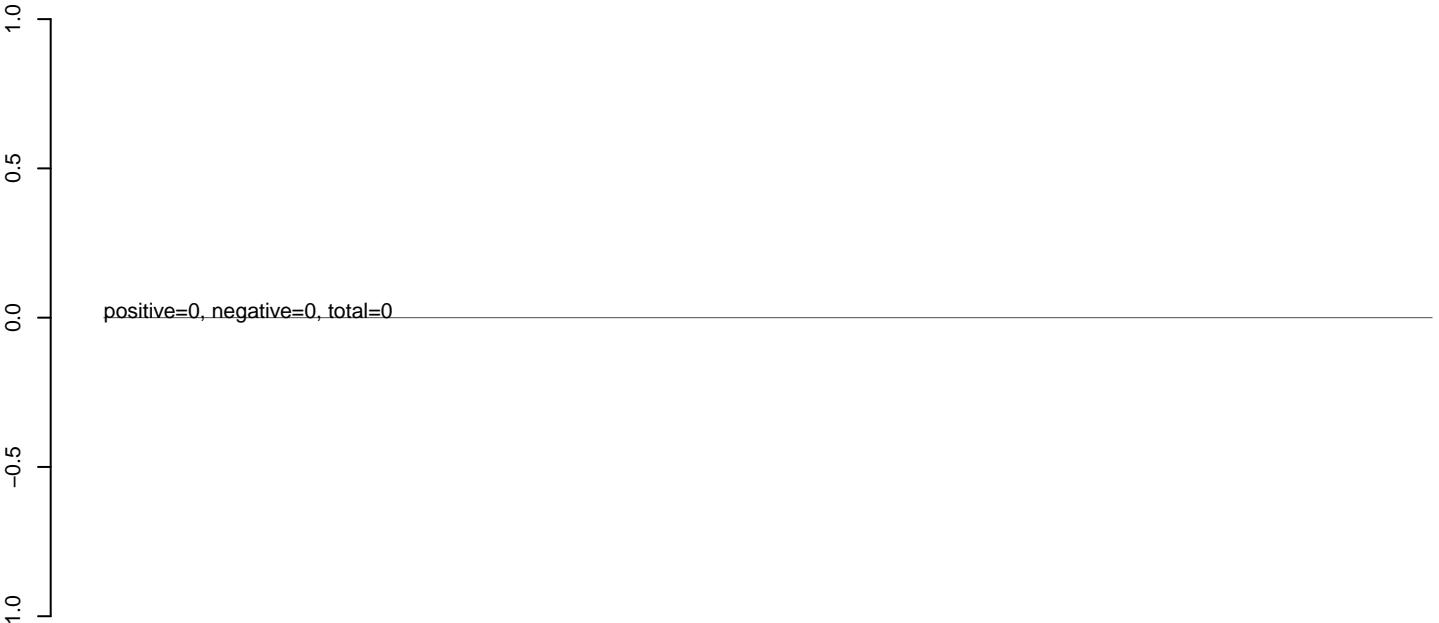
CuQuin_MG_7dpi3.rep

window positive=0, negative=0, total=0 NC_038707.1-Jonchet_virus_strain_B81-CI-2004_segment_S_plus_segment_M_plus_segment_L:1-14098

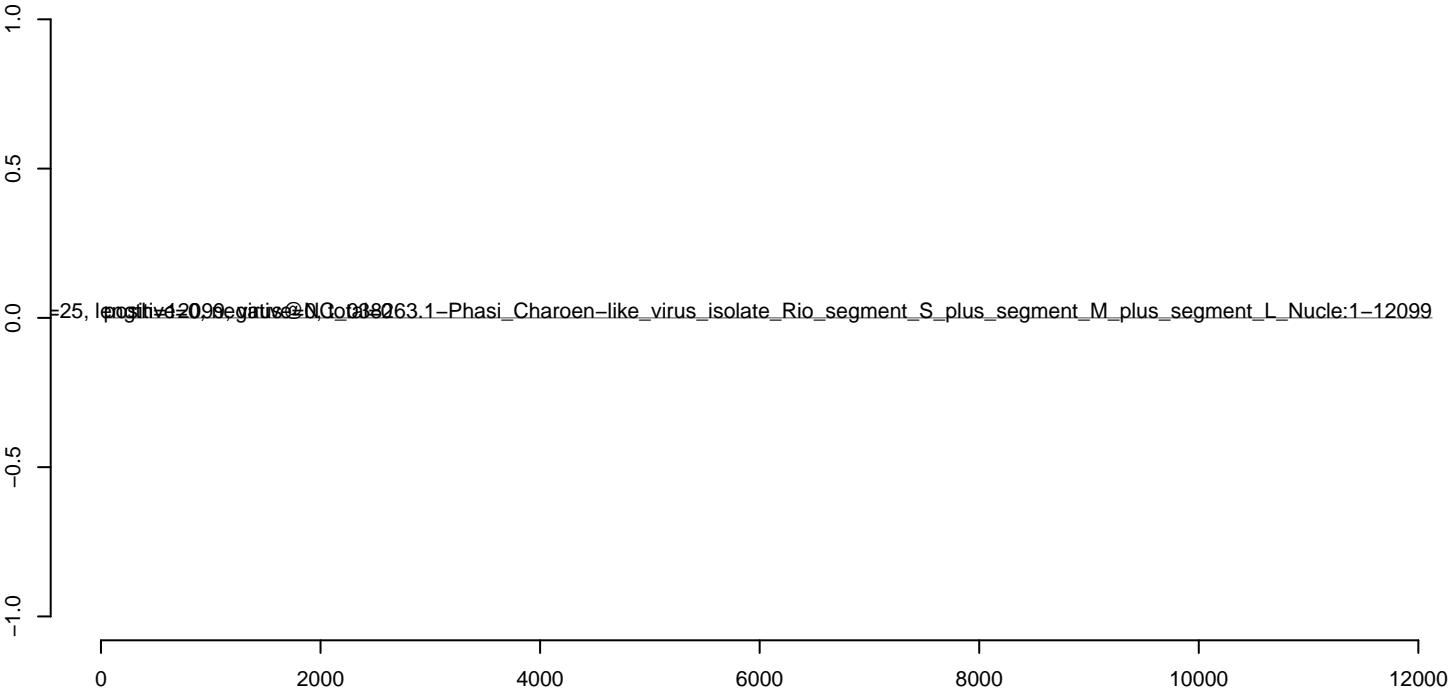
CuQuin_MG_7dpi3.18_23.rep

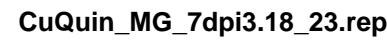


CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep





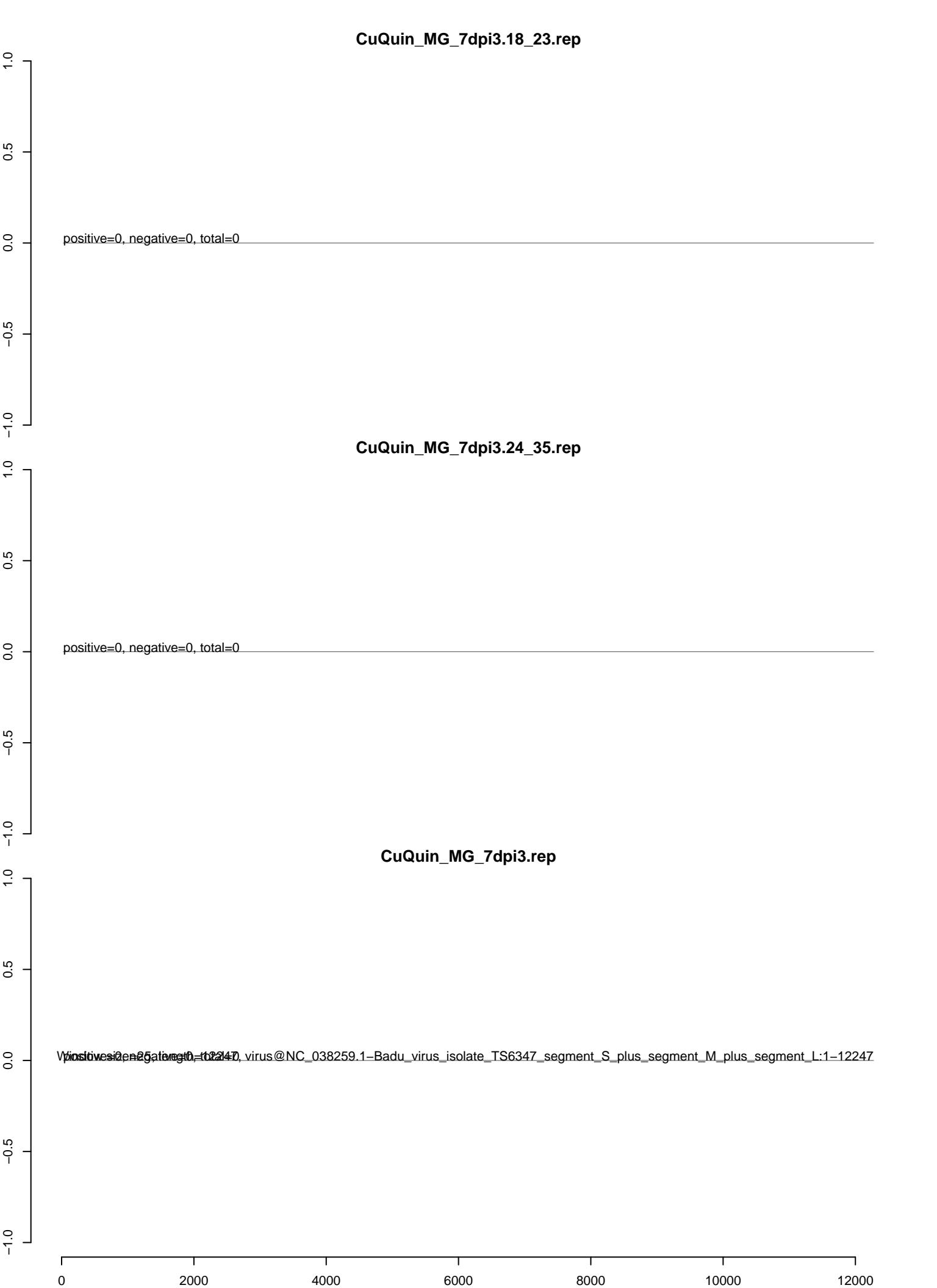
positive=0, negative=0, total=0



positive=0, negative=0, total=0



positive_virus_025_gatctc_0102247, virus@NC_038259.1-Badu_virus_isolate_TS6347_segment_S_plus_segment_M_plus_segment_L:1-12247



CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

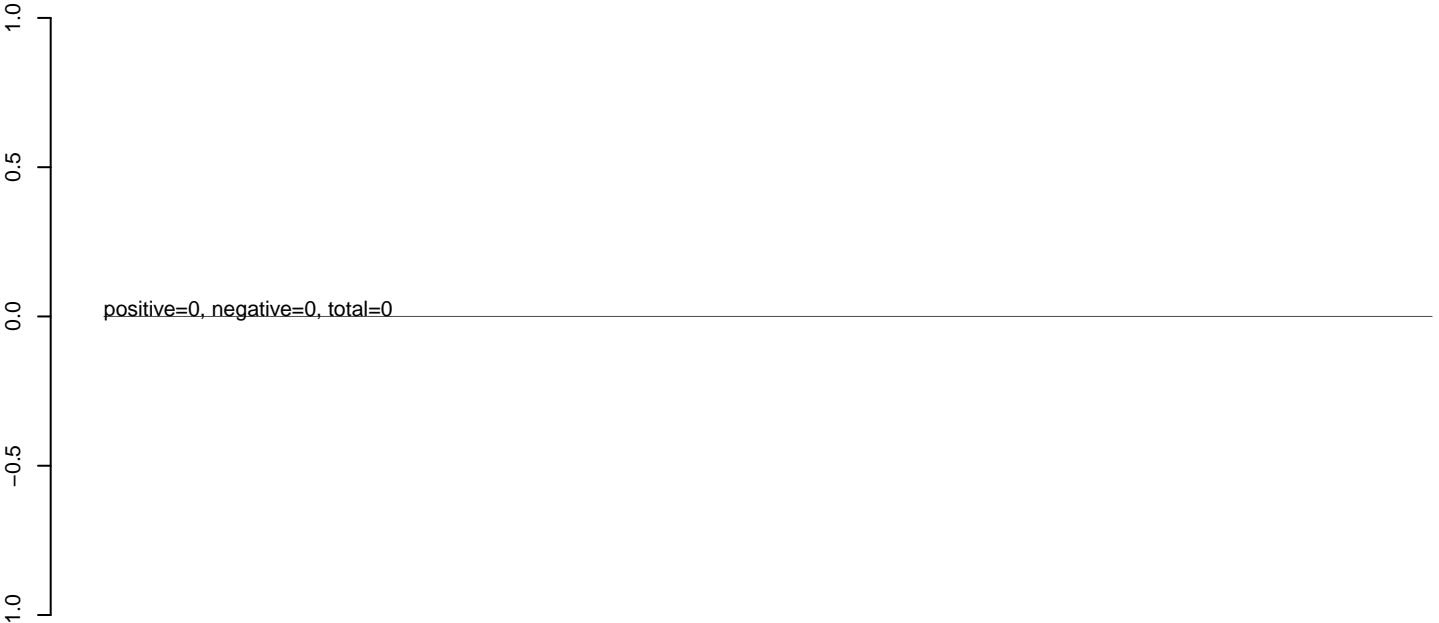
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

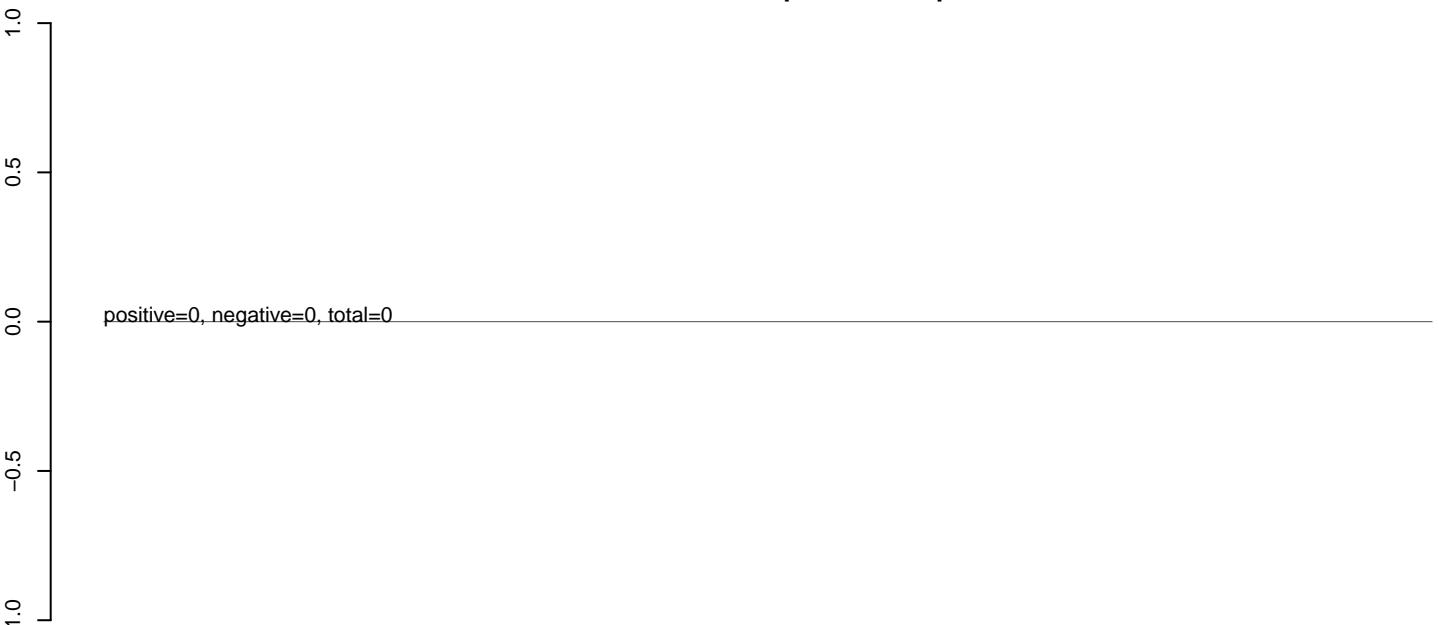
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0
Window size=25, length=11503, virus@NC_035132.1_Culex_rhabdo-like_virus_strain_mosWSB71420:1-11503

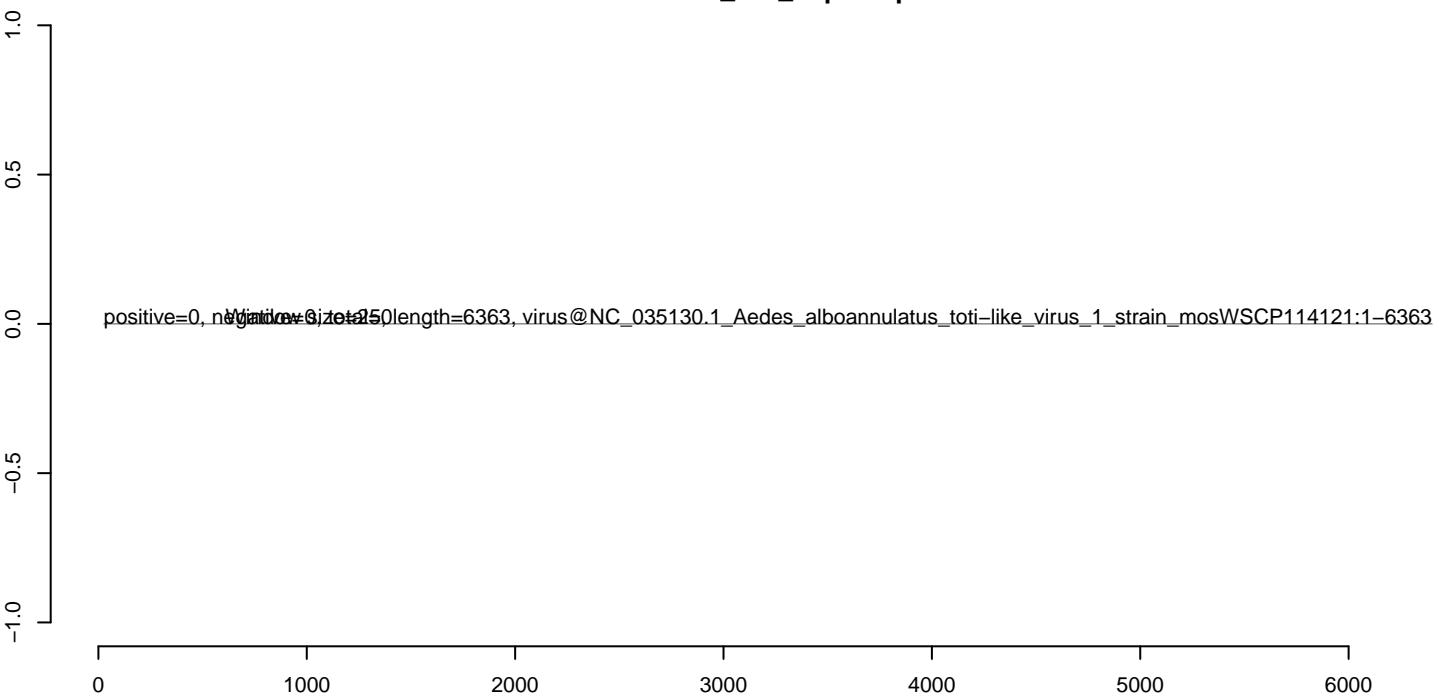
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0
Window size=25, length=9176, virus@NC_035129.1_Culex_negev-like_virus_3_strain_mos172X44875:1-9176

0 2000 4000 6000 8000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0
length=11470, virus@NC_035128.1_Aedes_camptorhynchus_negev-like_virus_strain_mos191CP47896:1–11470

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

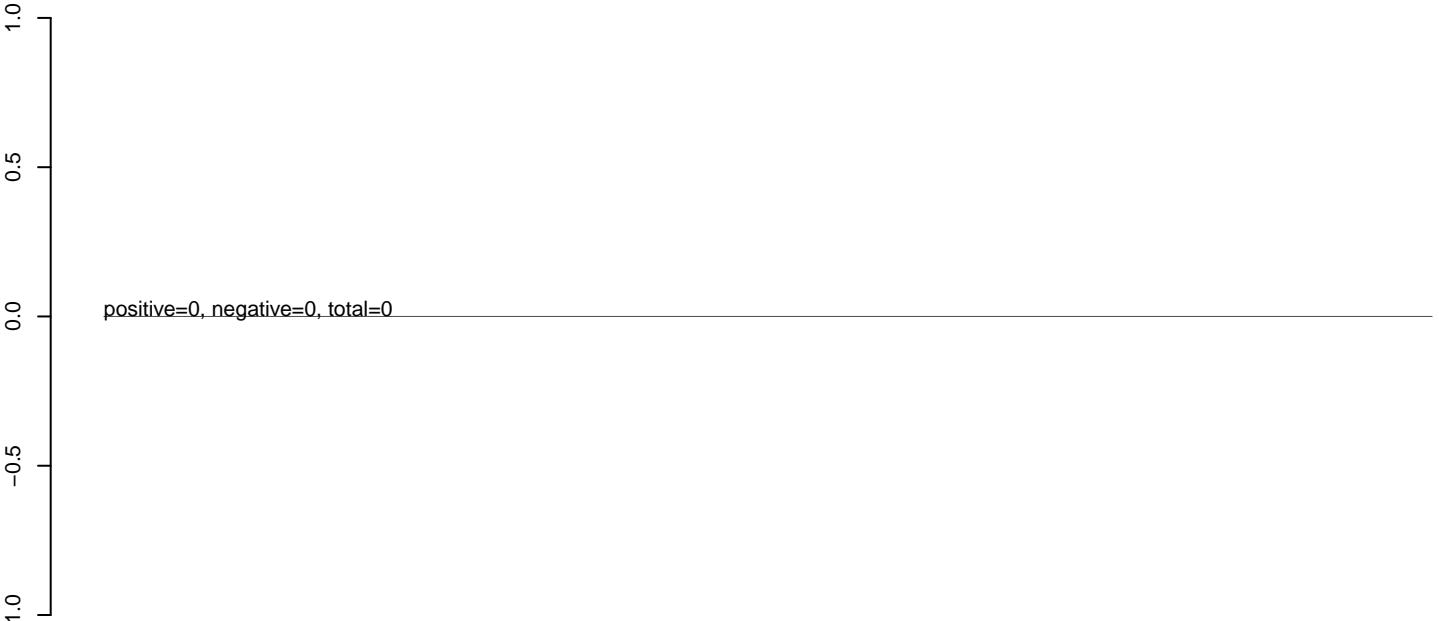
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

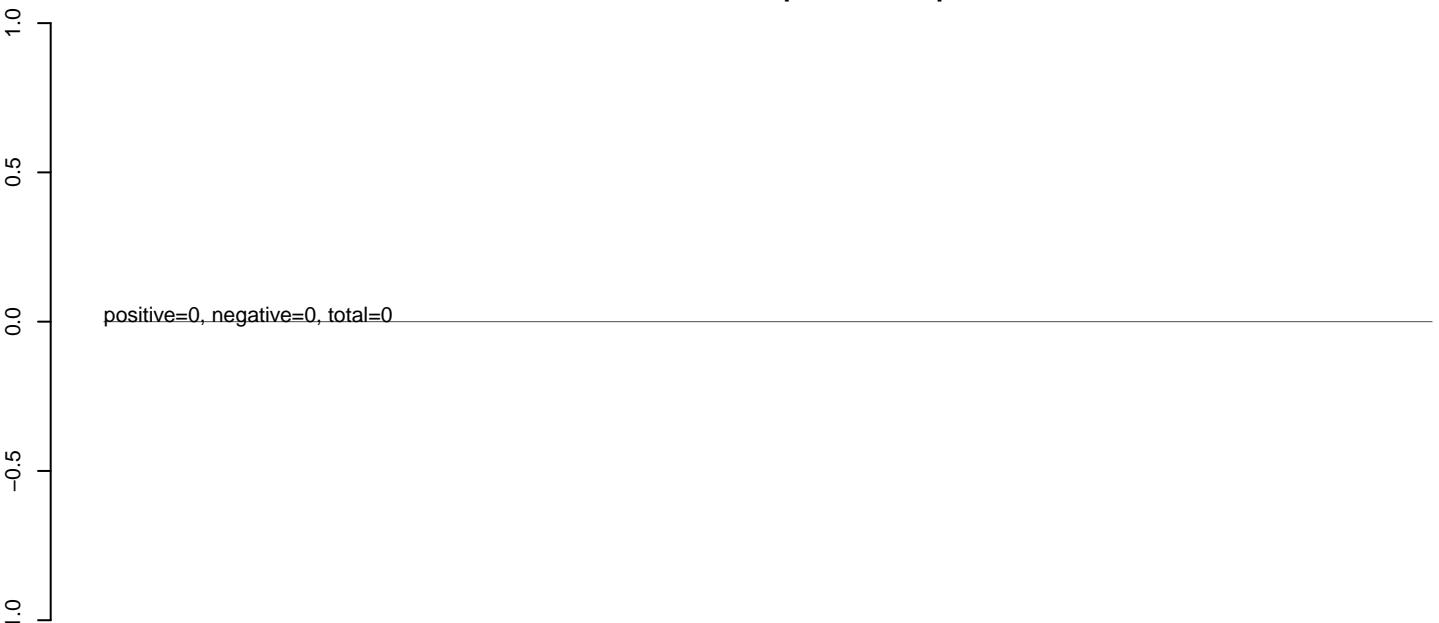
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0, window.size=25, length=2968, virus@NC_035126.1_Wilkie_narna-like_virus_1_strain_mosWSCP70929:1-2968

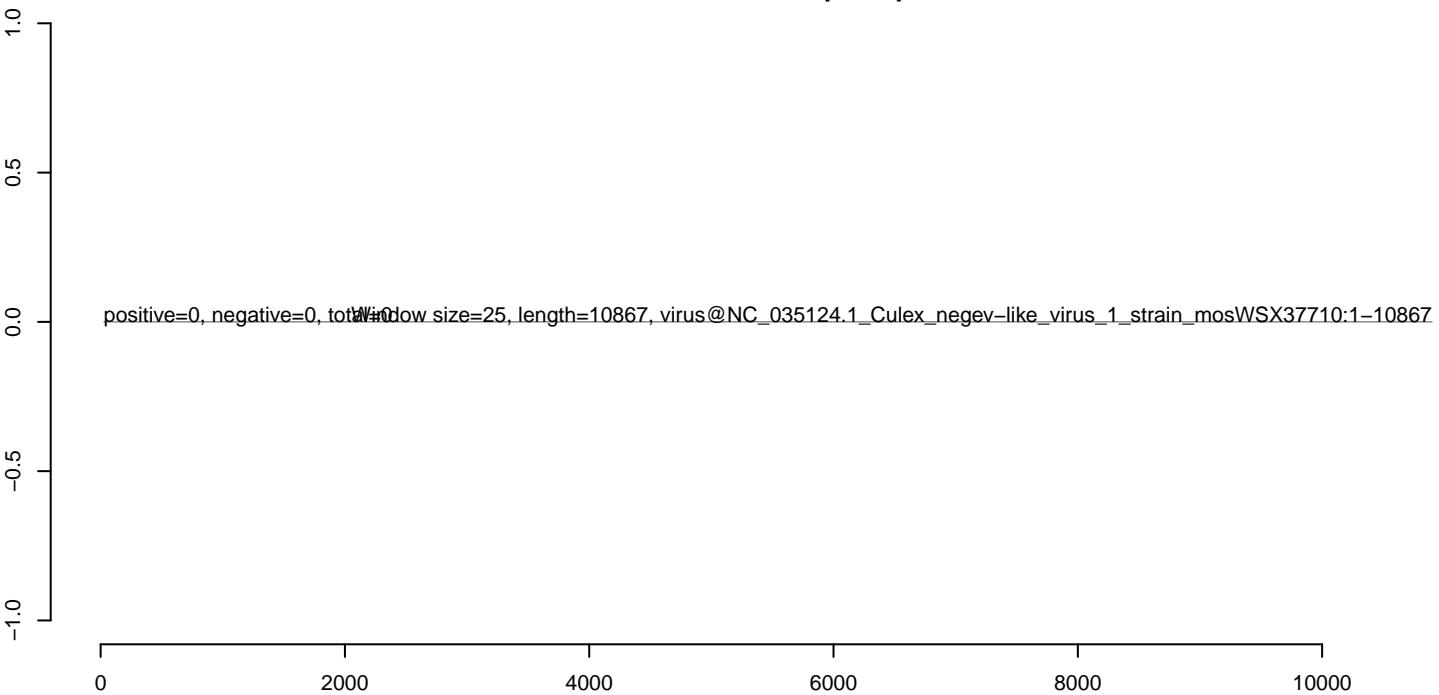
CuQuin_MG_7dpi3.18.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0
Window size=25, length=9324, virus@NC_035123.1_Culex_negev-like_virus_2_strain_mos172X30622:1-9324

0 2000 4000 6000 8000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, ~~Window size=25, length=2857, virus@NC_035121.1_Point-Douro_narna-like_virus_strain_mos172gb29666:1-2857~~

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

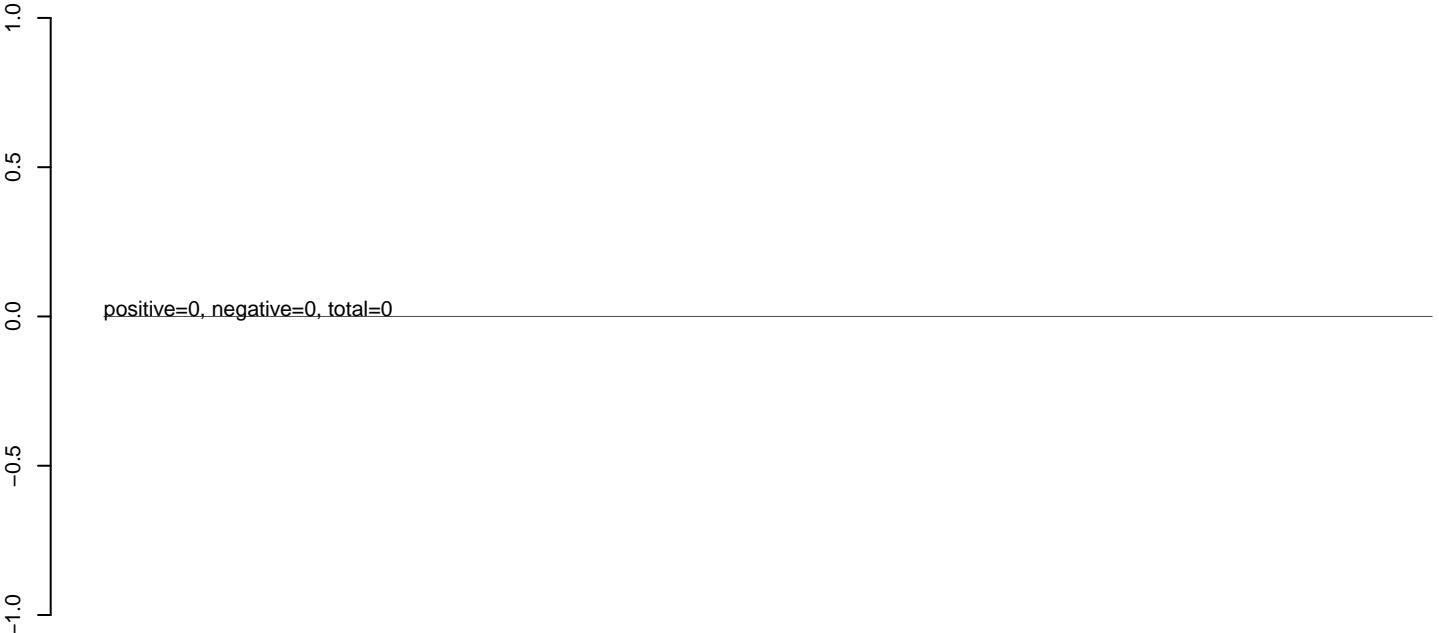
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

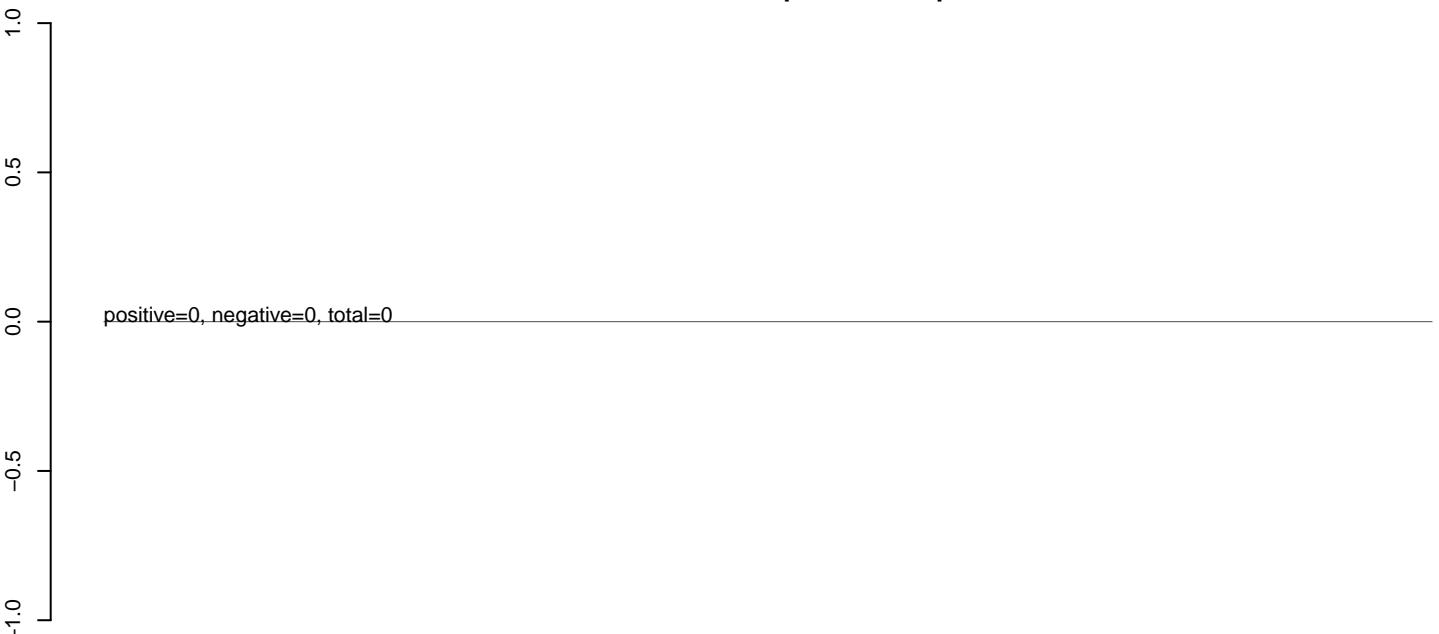
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0, window.size=25, length=3367, virus@NC_035120.1_Wilkie_narna-like_virus_2_strain_mosWSCP85442:1-3367

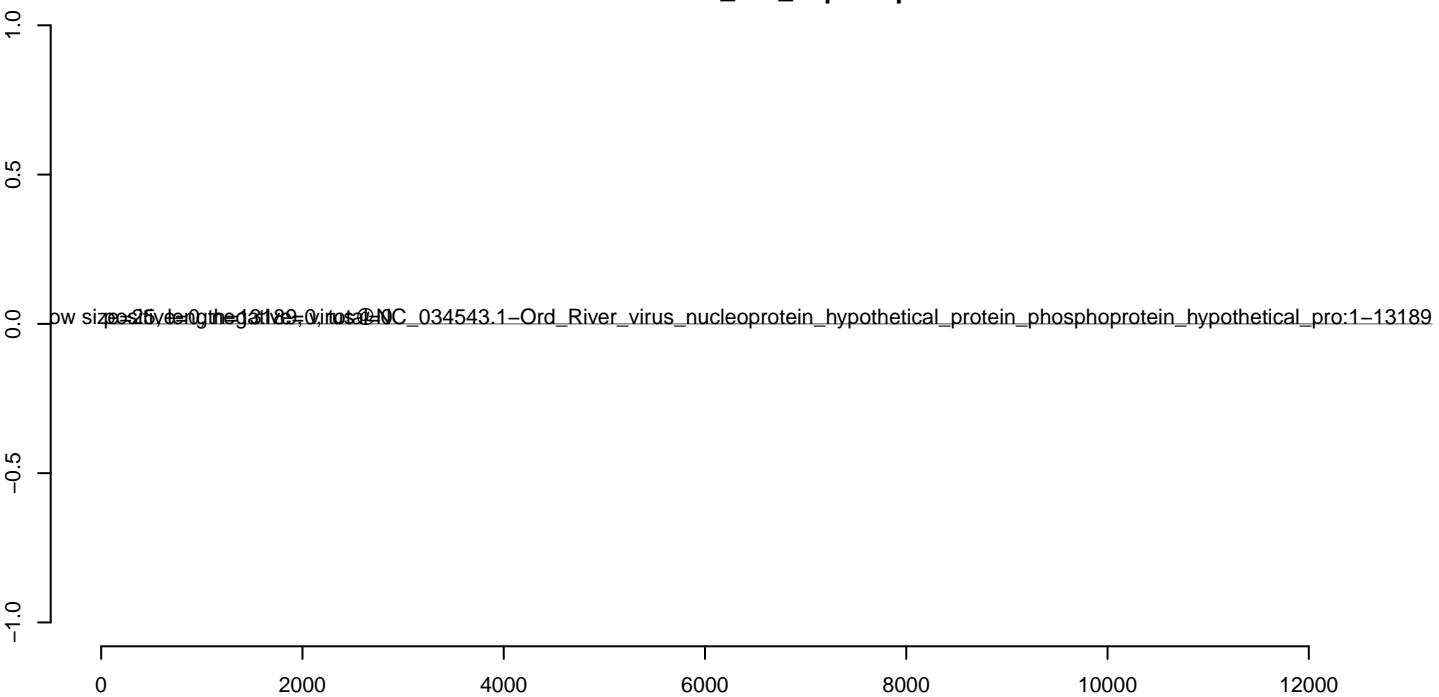
CuQuin_MG_7dpi3.18_23.rep



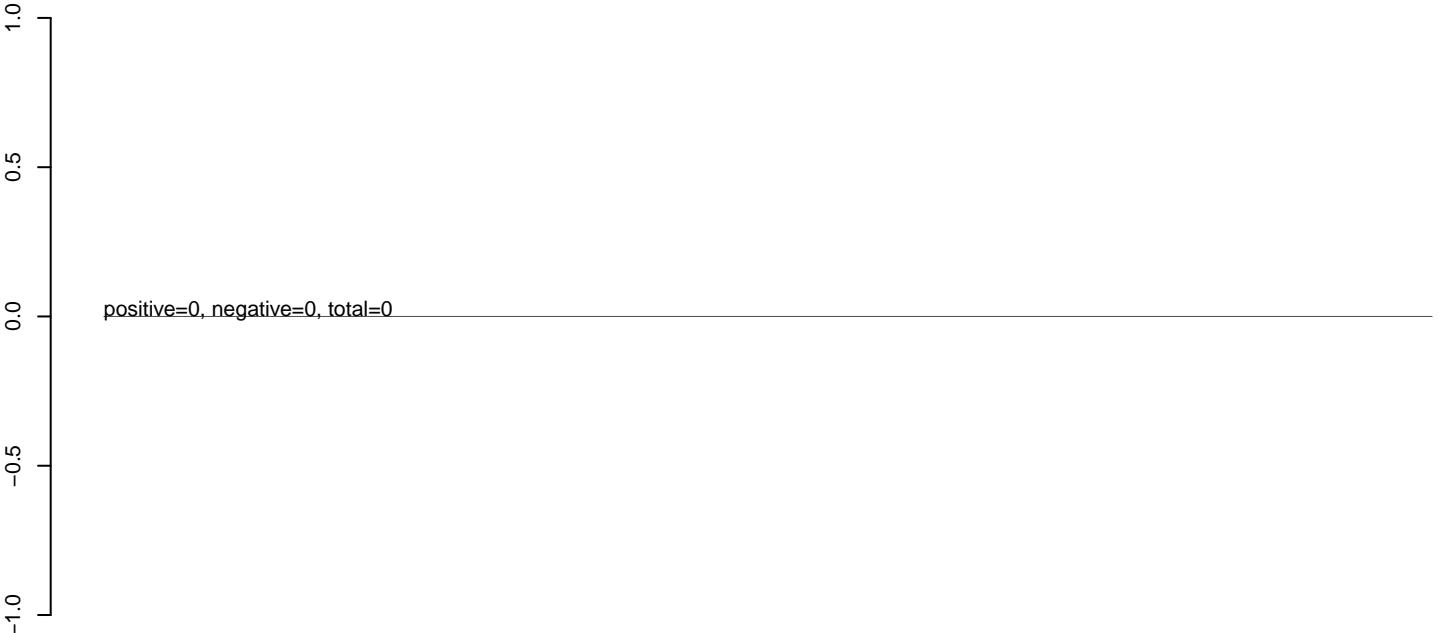
CuQuin_MG_7dpi3.24_35.rep



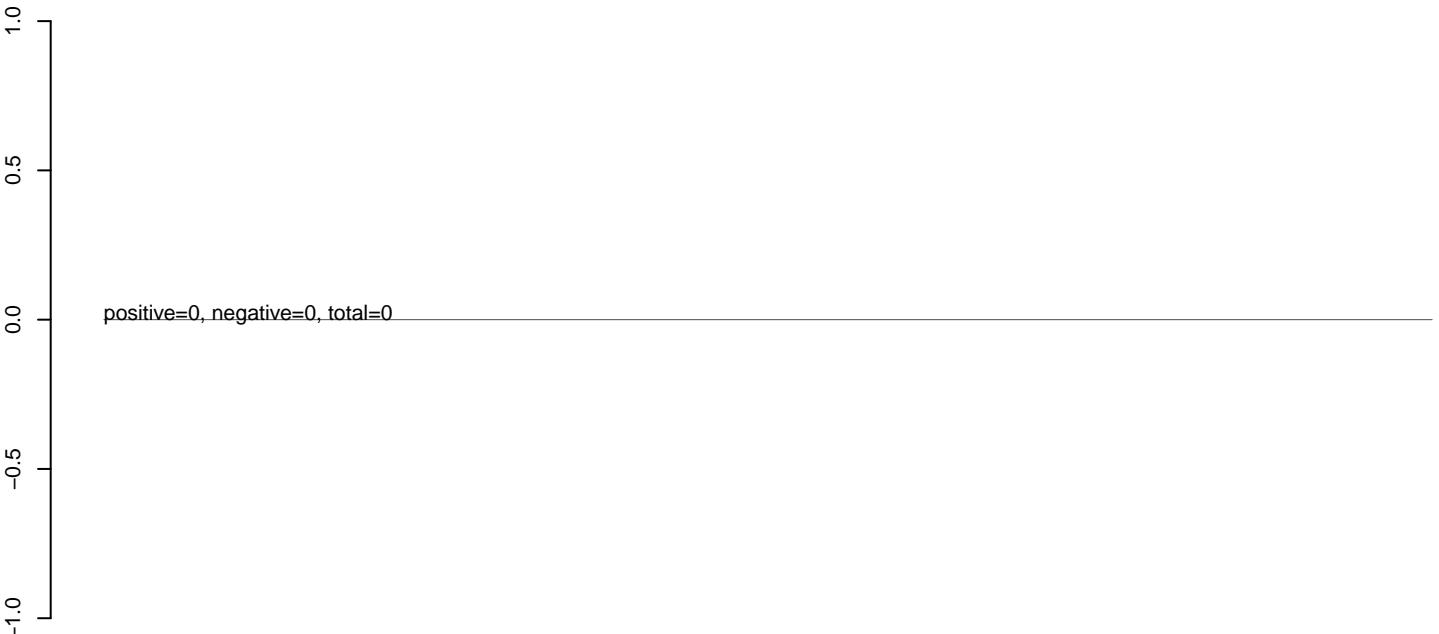
CuQuin_MG_7dpi3.rep



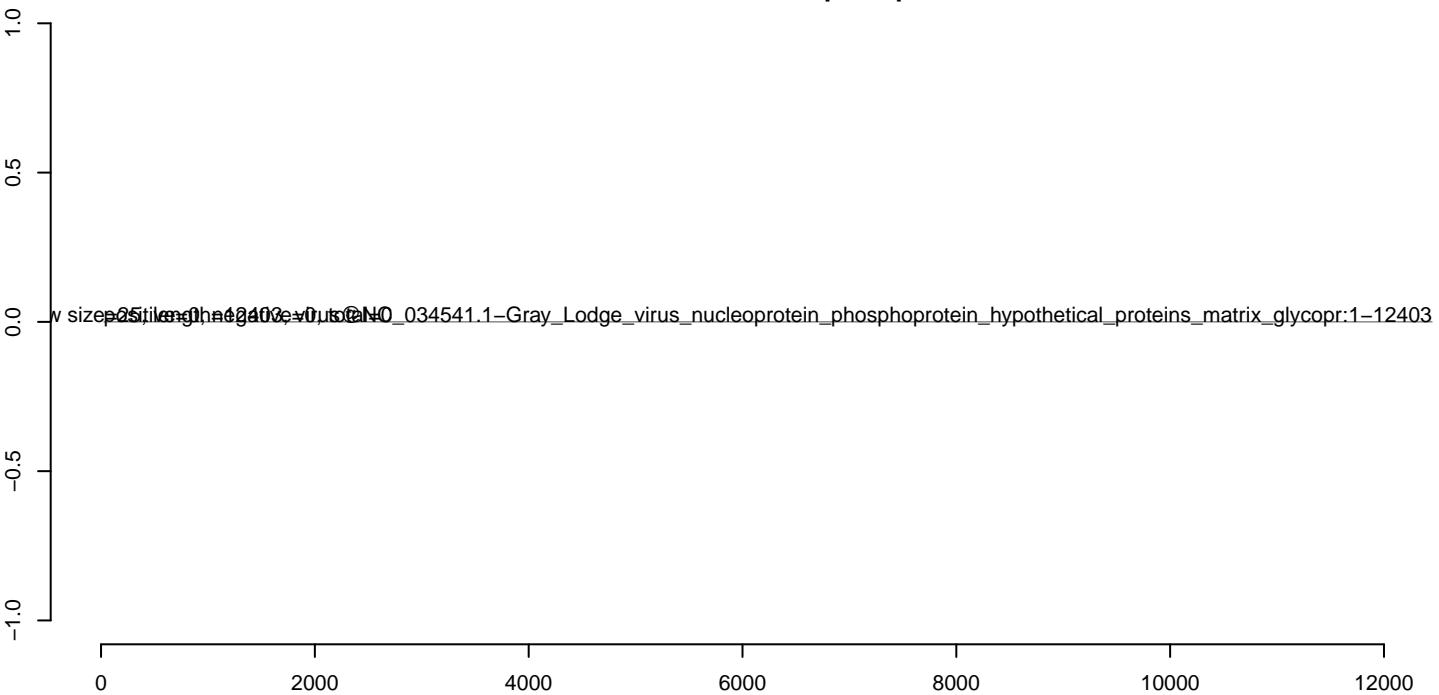
CuQuin_MG_7dpi3.18_23.rep



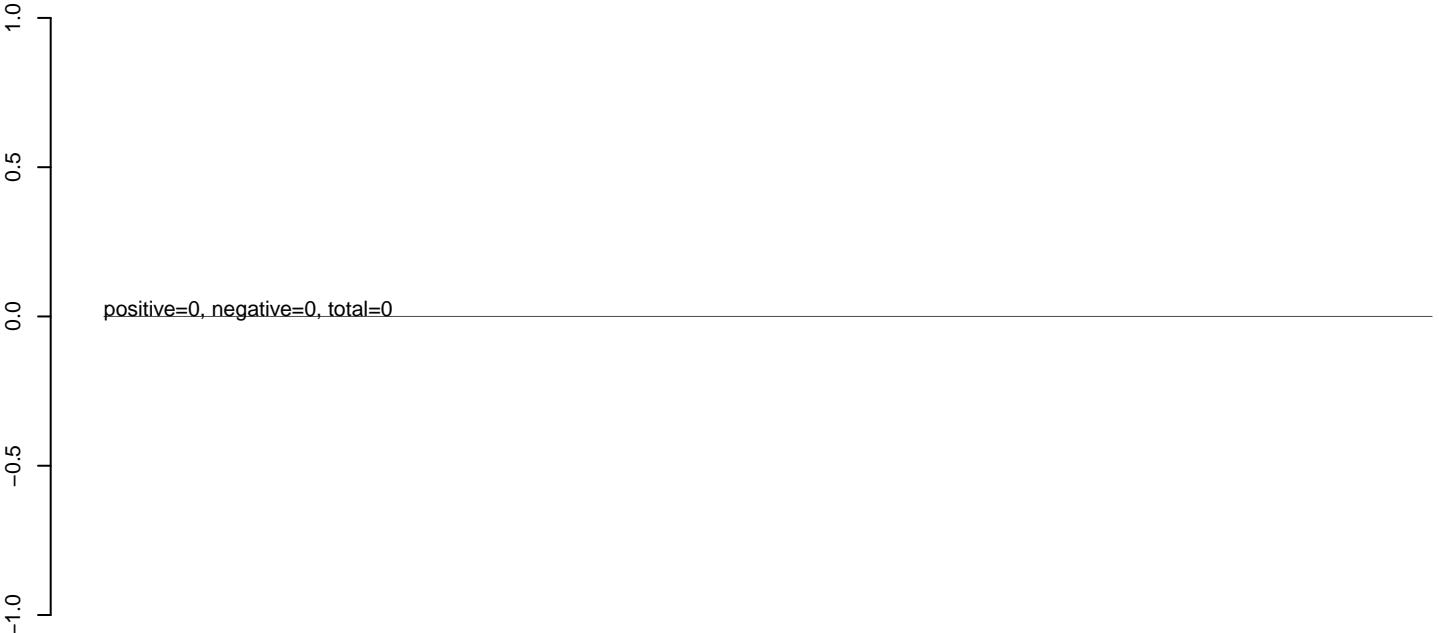
CuQuin_MG_7dpi3.24_35.rep



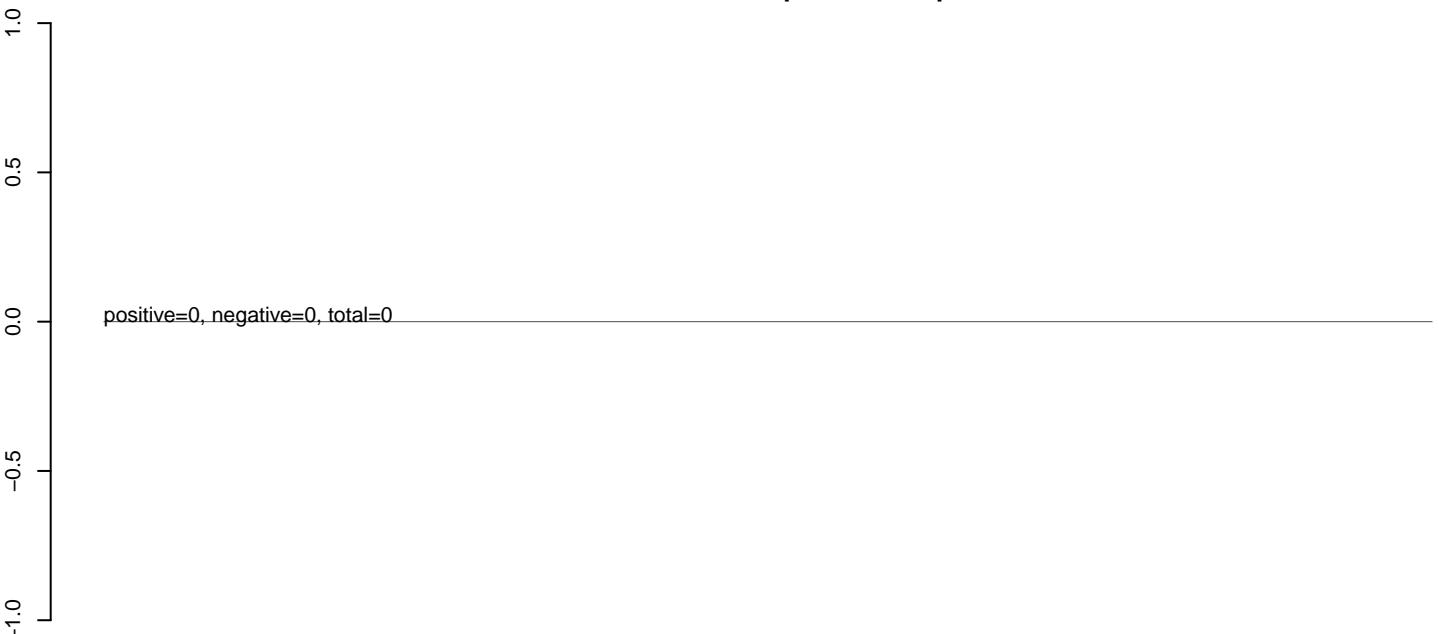
CuQuin_MG_7dpi3.rep



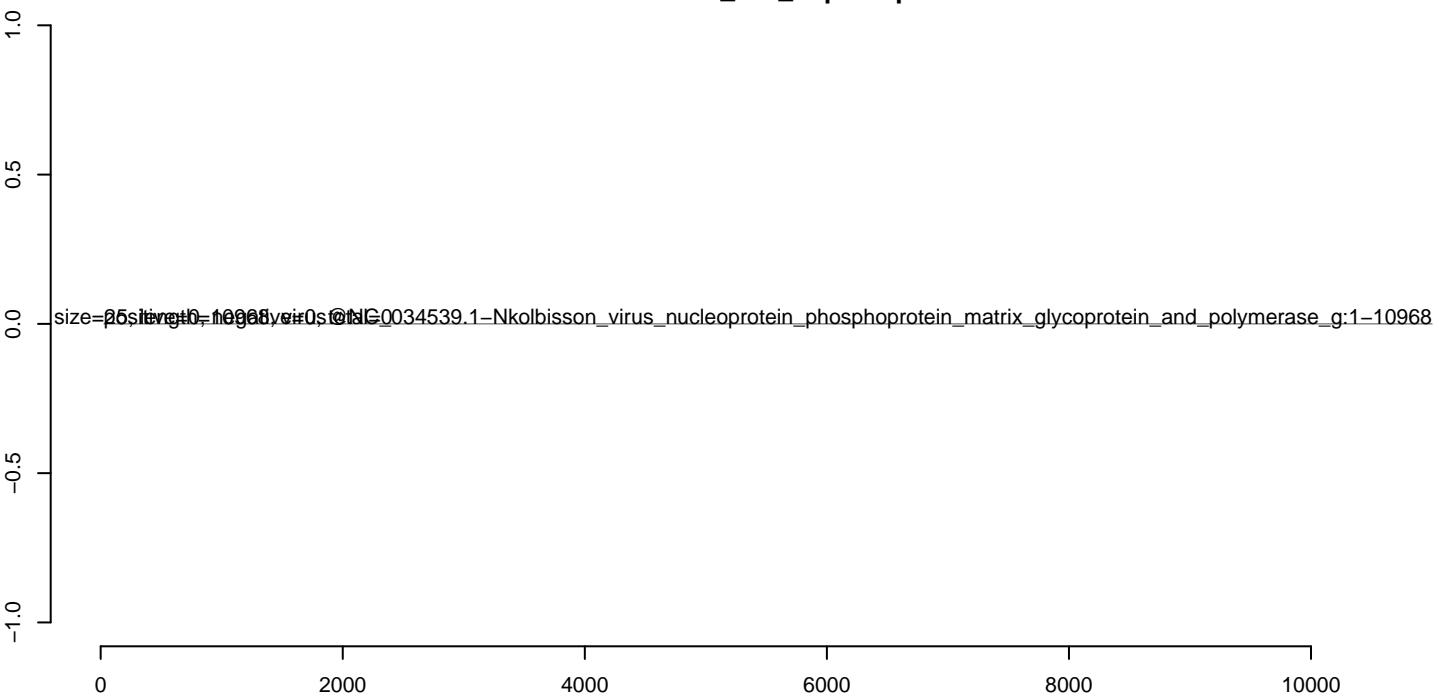
CuQuin_MG_7dpi3.18_23.rep



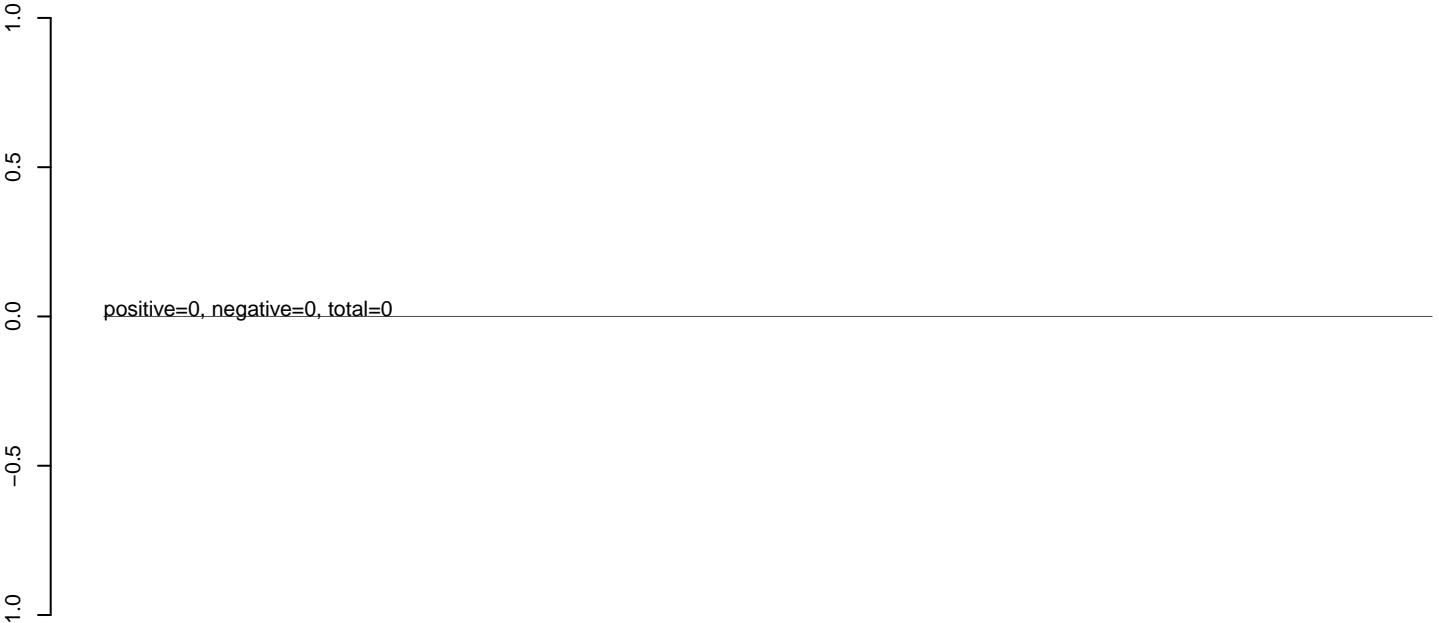
CuQuin_MG_7dpi3.24_35.rep



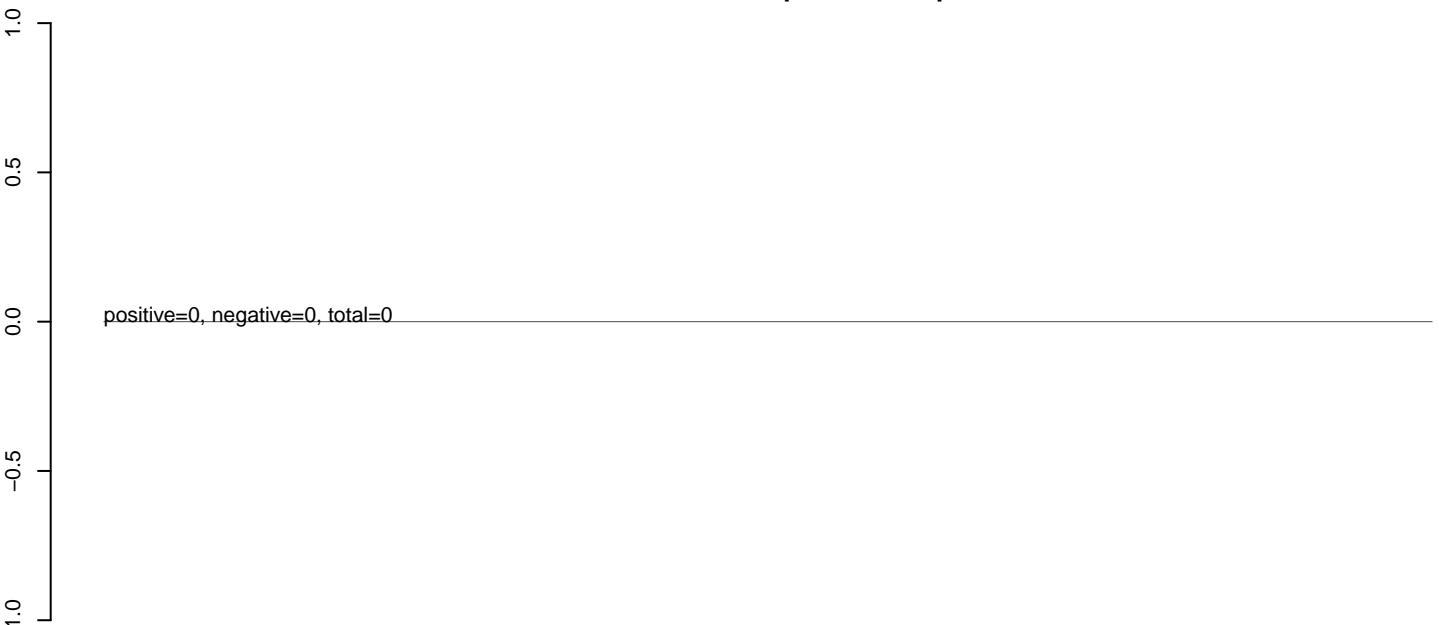
CuQuin_MG_7dpi3.rep



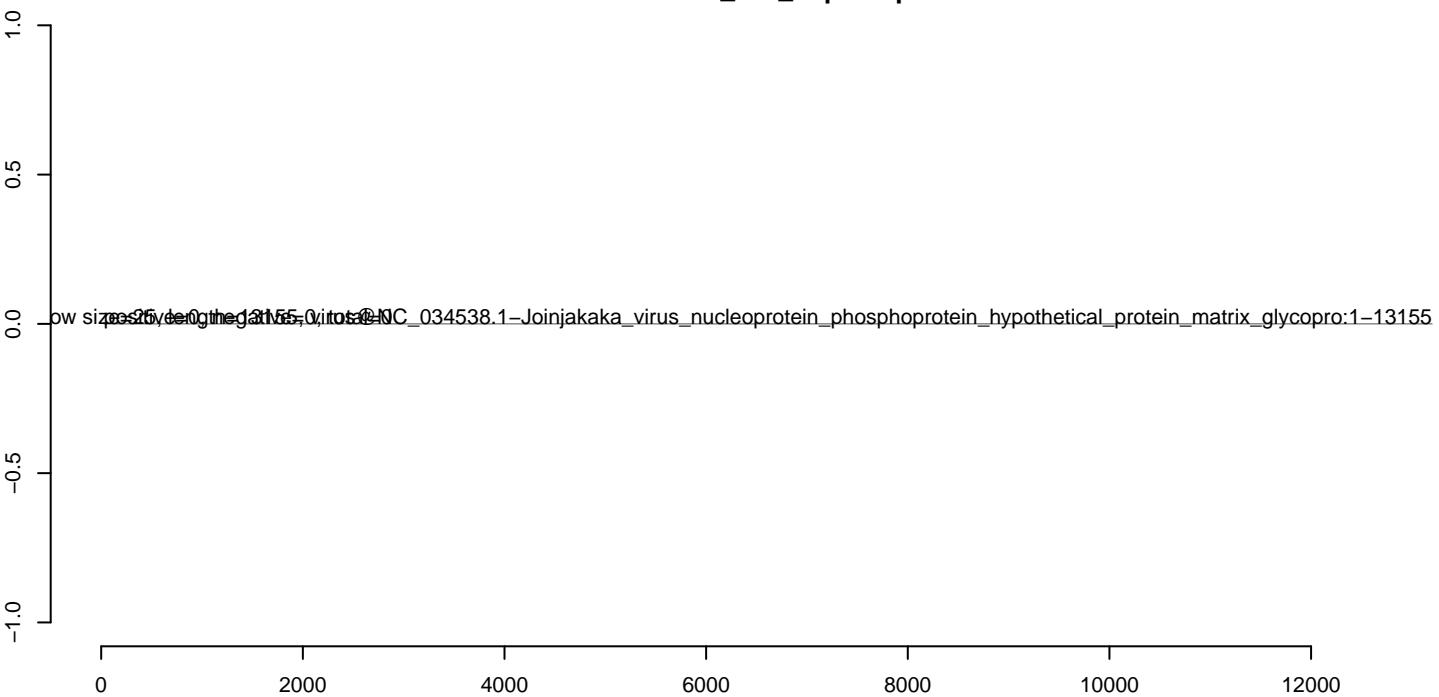
CuQuin_MG_7dpi3.18_23.rep



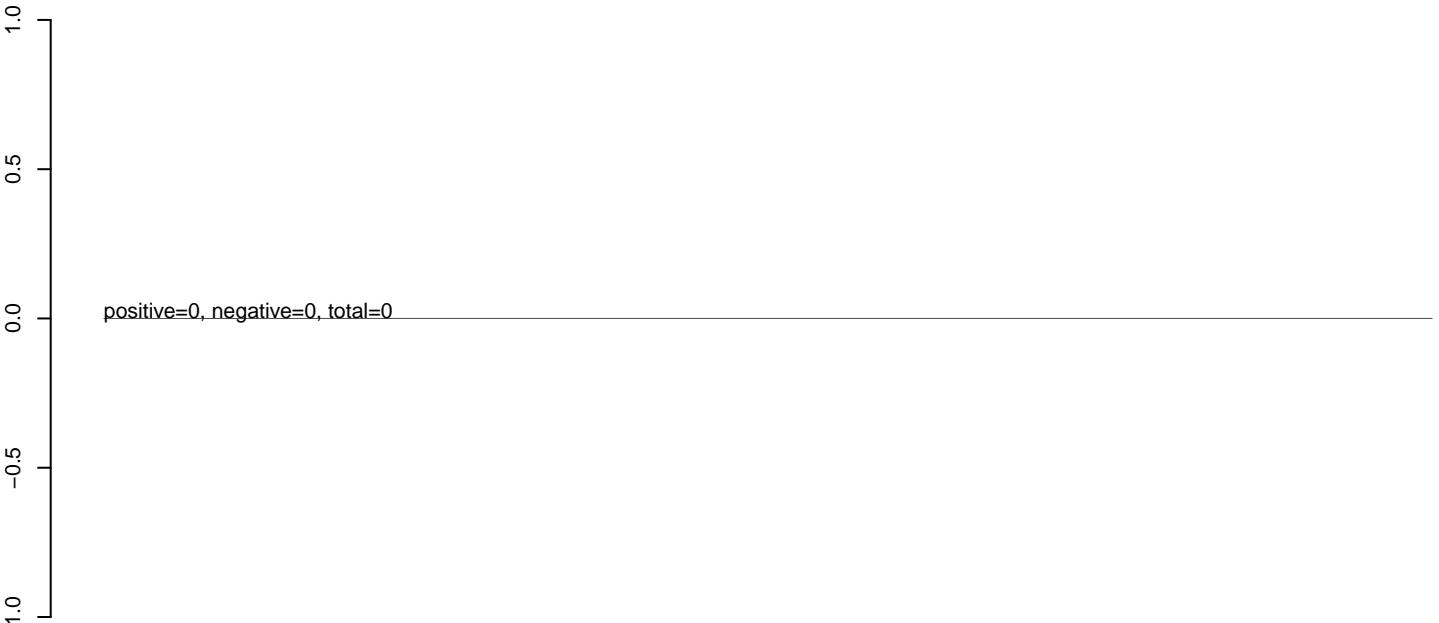
CuQuin_MG_7dpi3.24_35.rep



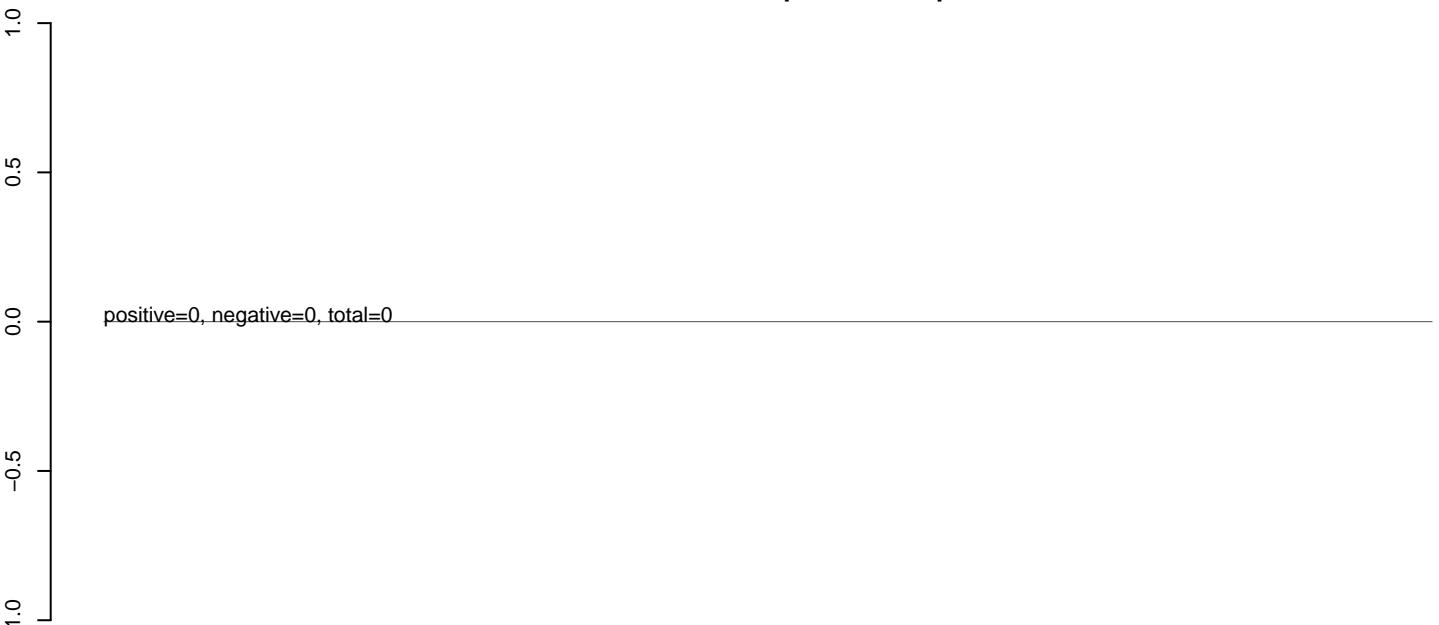
CuQuin_MG_7dpi3.rep



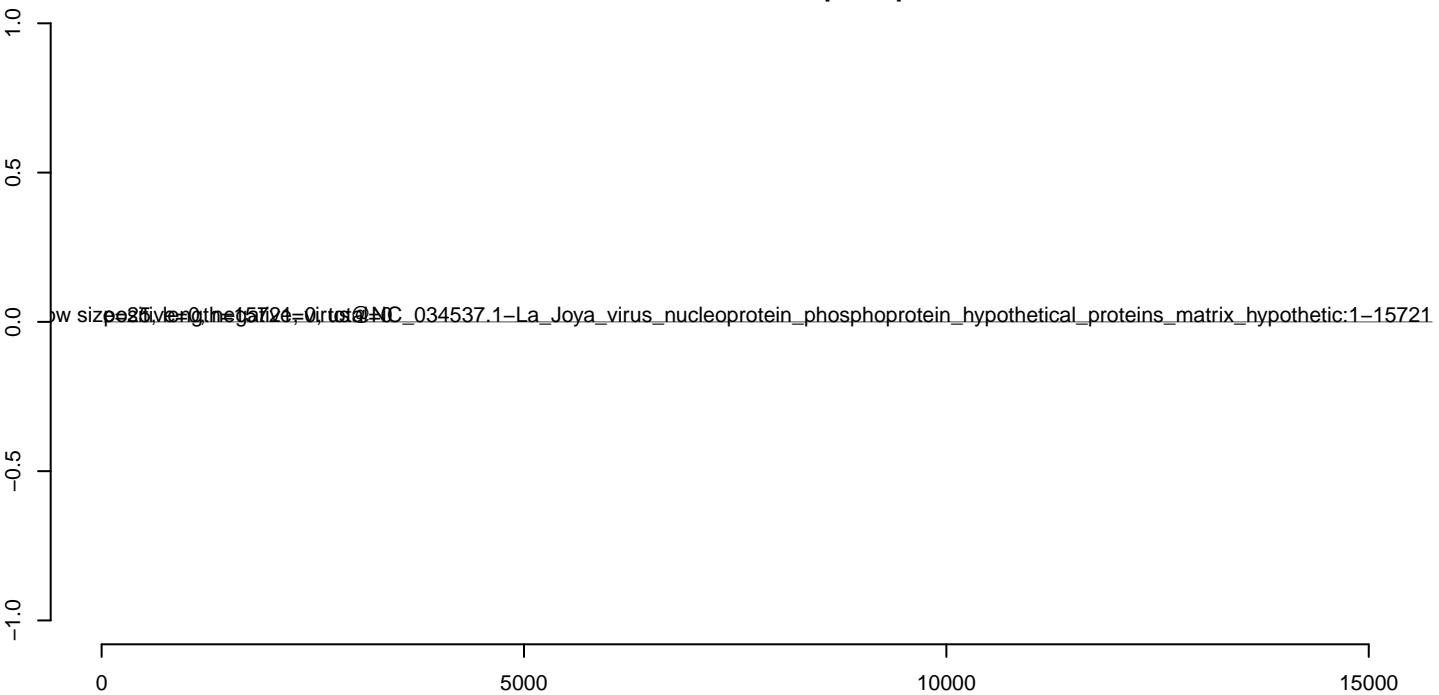
CuQuin_MG_7dpi3.18_23.rep



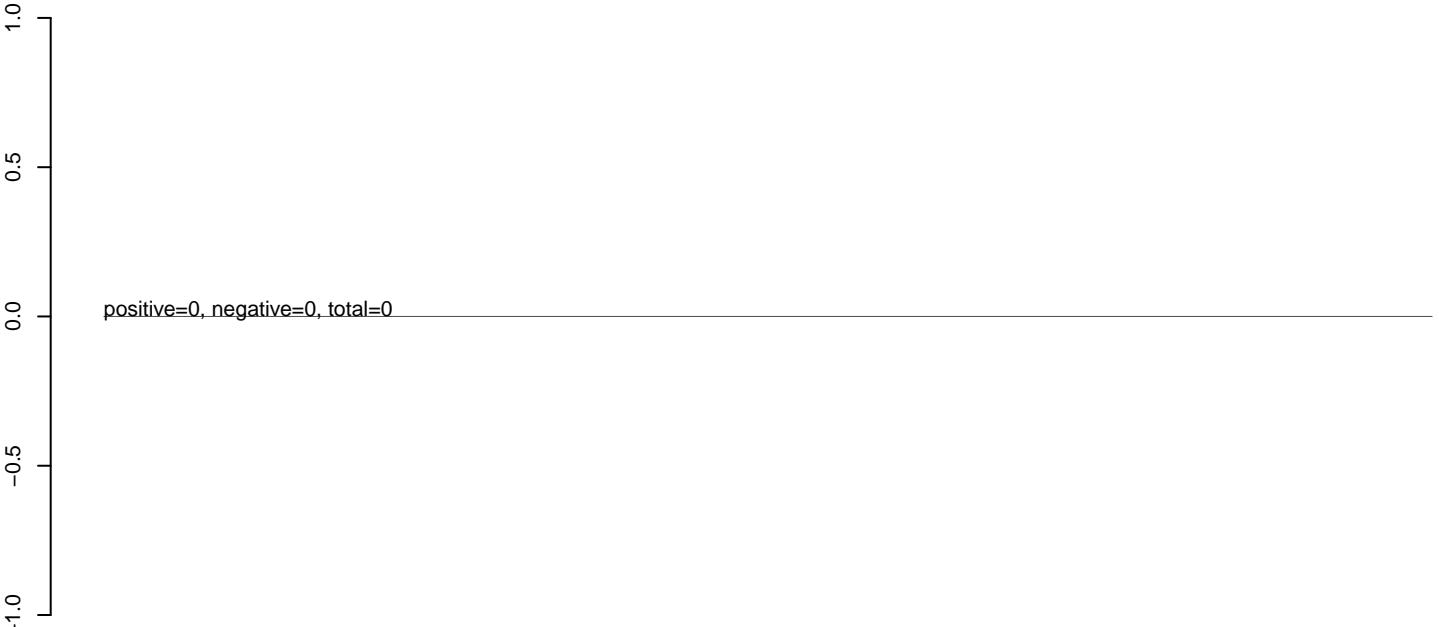
CuQuin_MG_7dpi3.24_35.rep



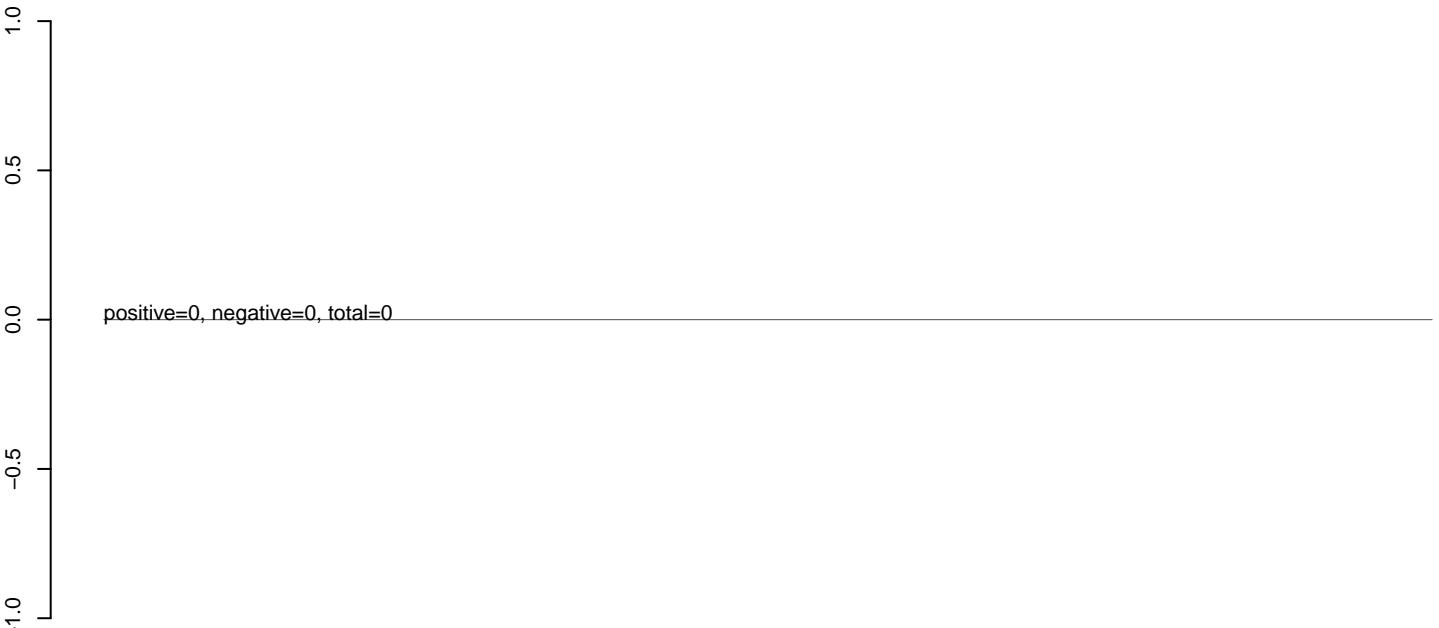
CuQuin_MG_7dpi3.rep



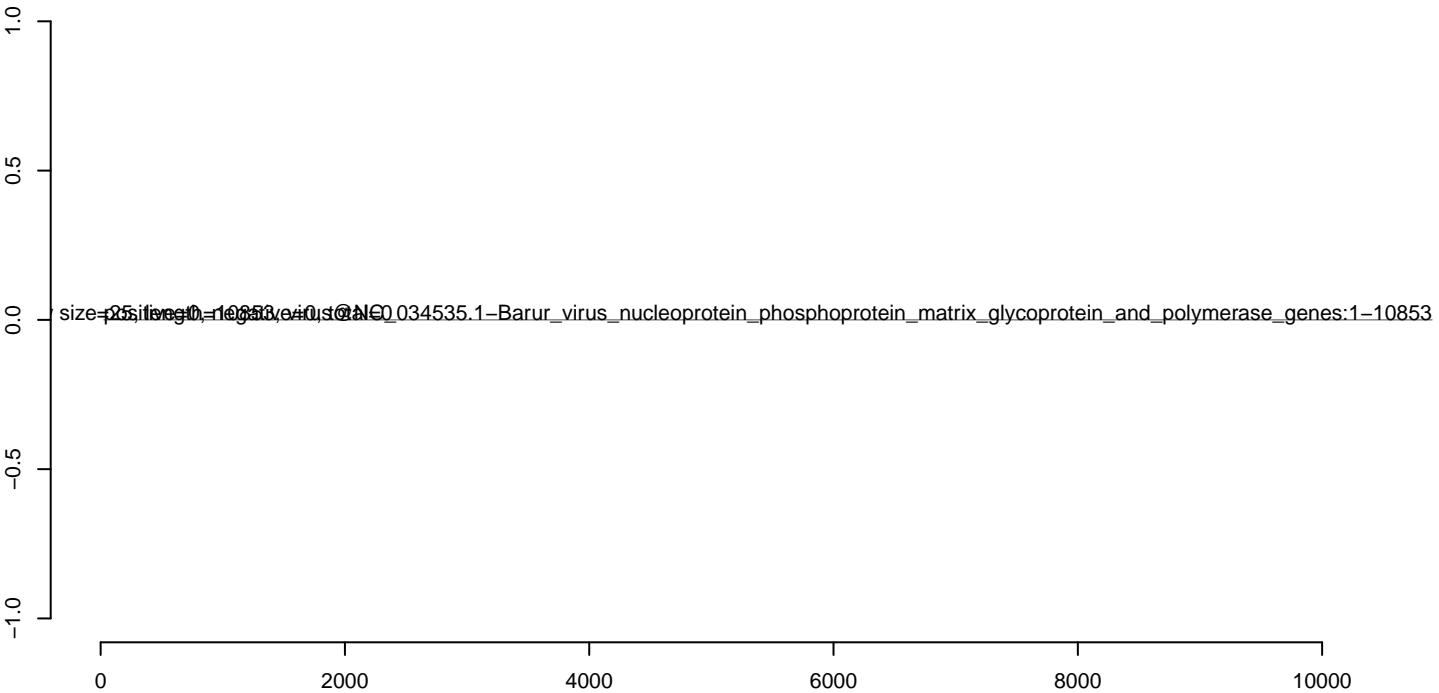
CuQuin_MG_7dpi3.18_23.rep

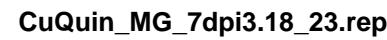


CuQuin_MG_7dpi3.24_35.rep

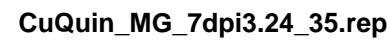


CuQuin_MG_7dpi3.rep





positive=0, negative=0, total=0



positive=0, negative=0, total=0



size=positive|6|18590|virostbl|0034534.1|Rochambeau_virus_nucleoprotein_phosphoprotein_matrix_hypothetical_proteins_glycopr:1-13593

0 2000 4000 6000 8000 10000 12000 14000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

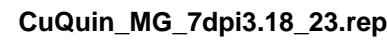
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

low size: 25 bp, no other data, 0 initial NC_034531.1-Manitoba_virus_nucleoprotein_phosphoprotein_hypothetical_proteins_matrix_glycoprot:1-13784

0 2000 4000 6000 8000 10000 12000 14000



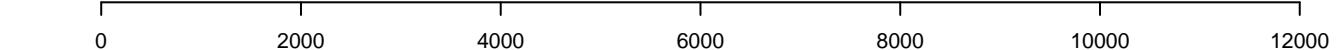
positive=0, negative=0, total=0



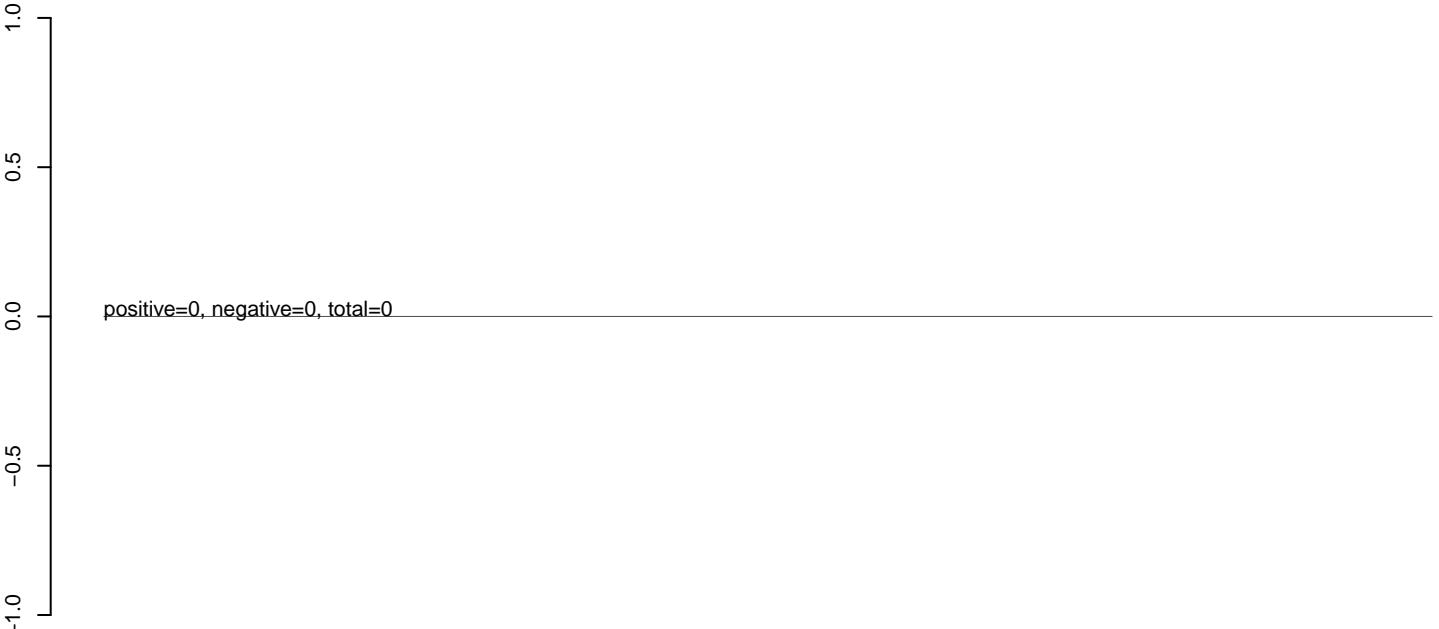
positive=0, negative=0, total=0



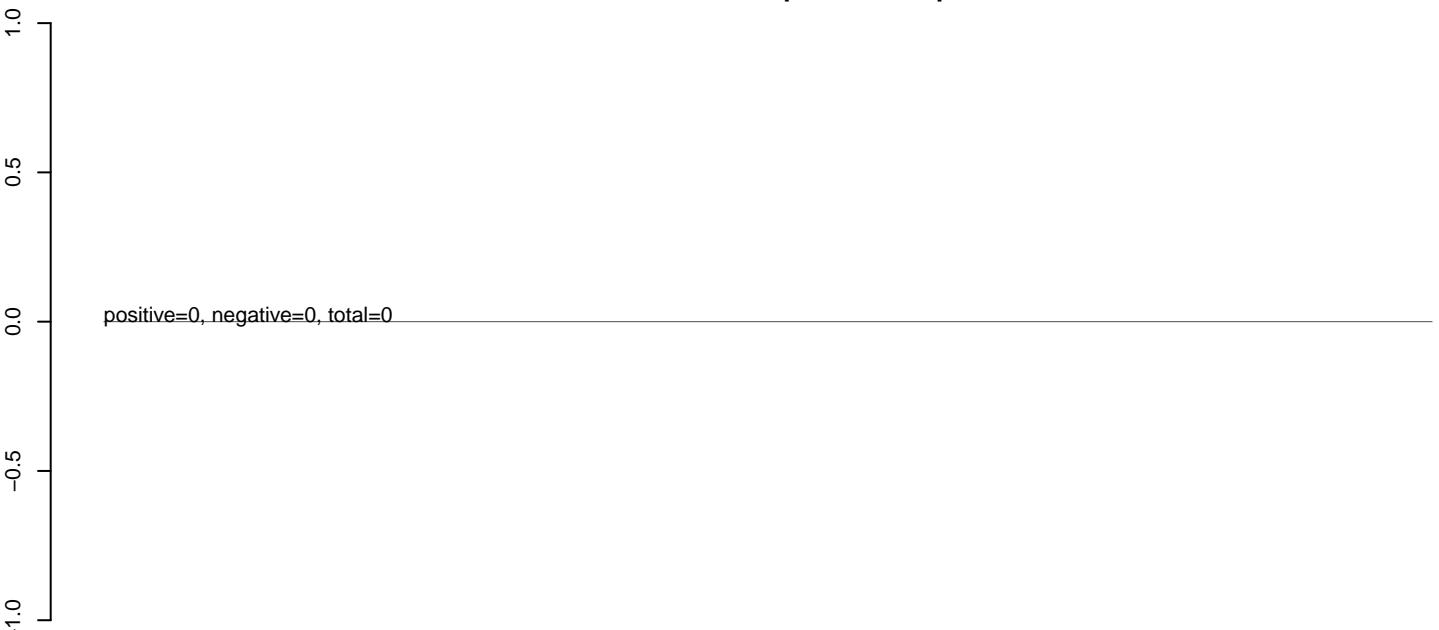
bio size:515 kb on the 13294 - virus NC_034530.1-Marco virus nucleoprotein phosphoprotein matrix glycoprotein hypothetical proteins:1-13294



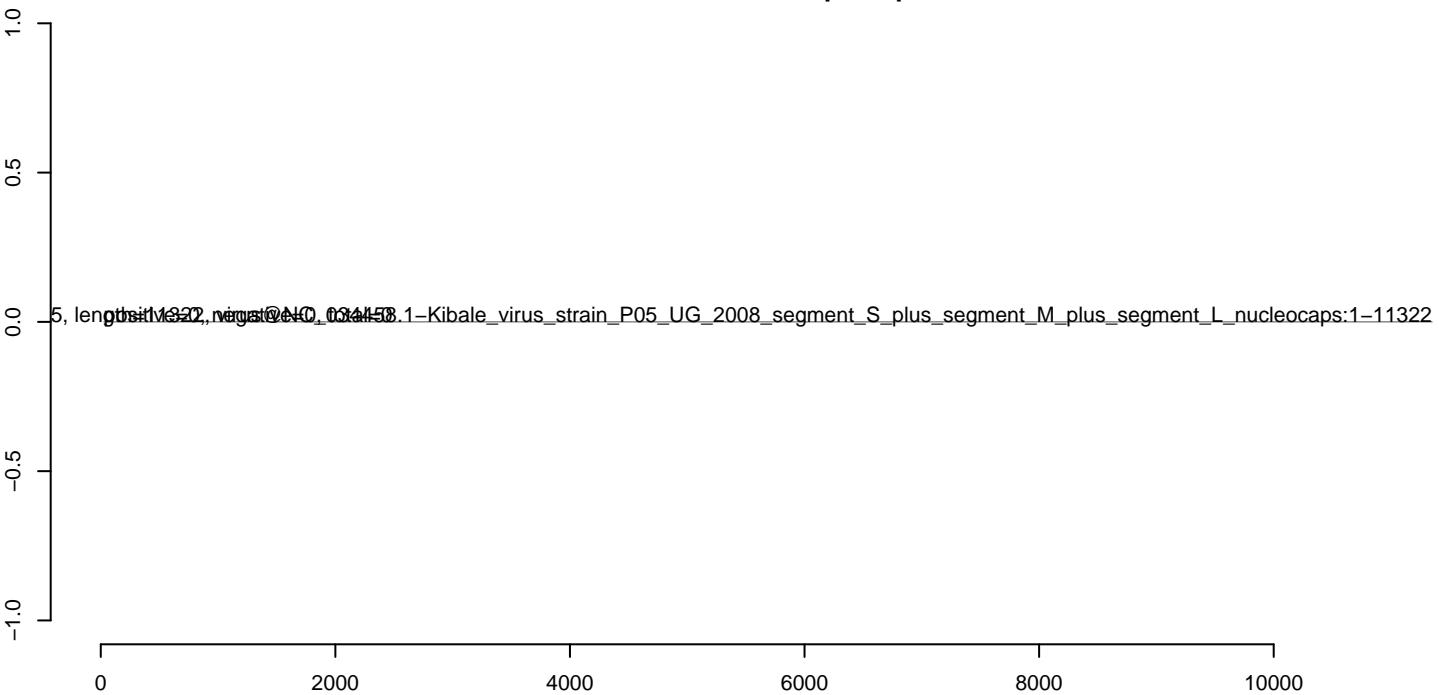
CuQuin_MG_7dpi3.18_23.rep



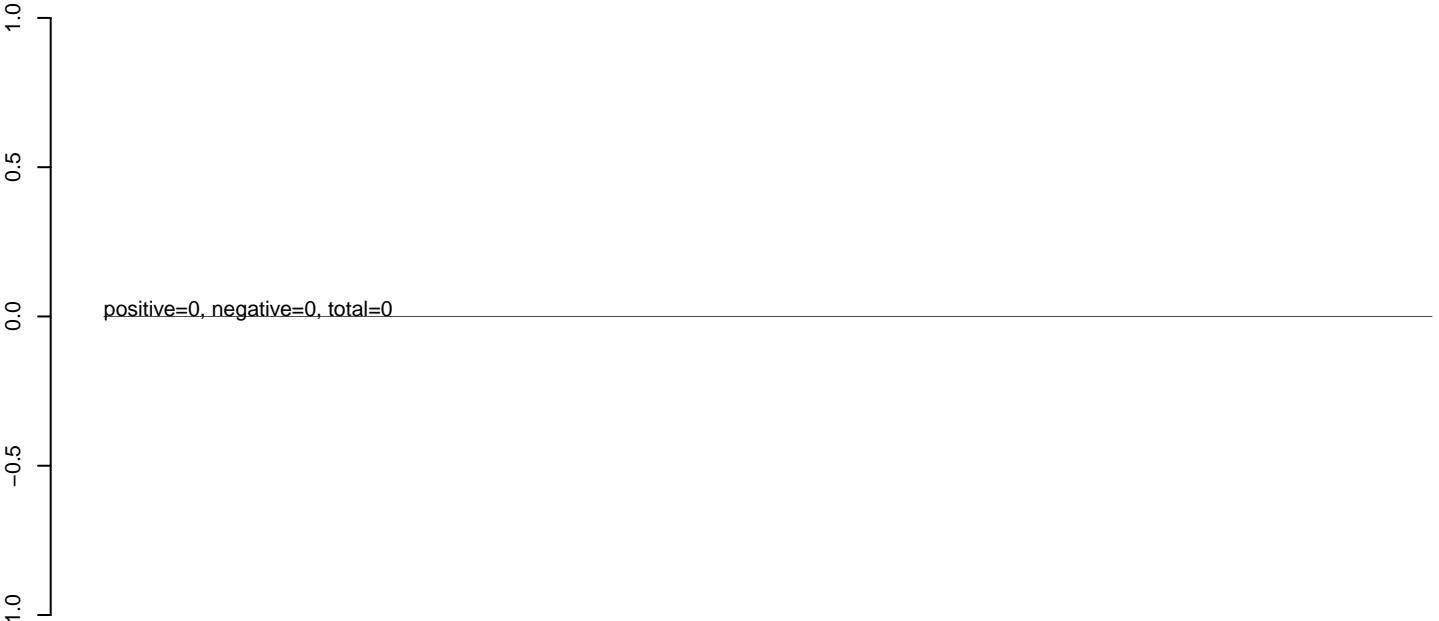
CuQuin_MG_7dpi3.24_35.rep



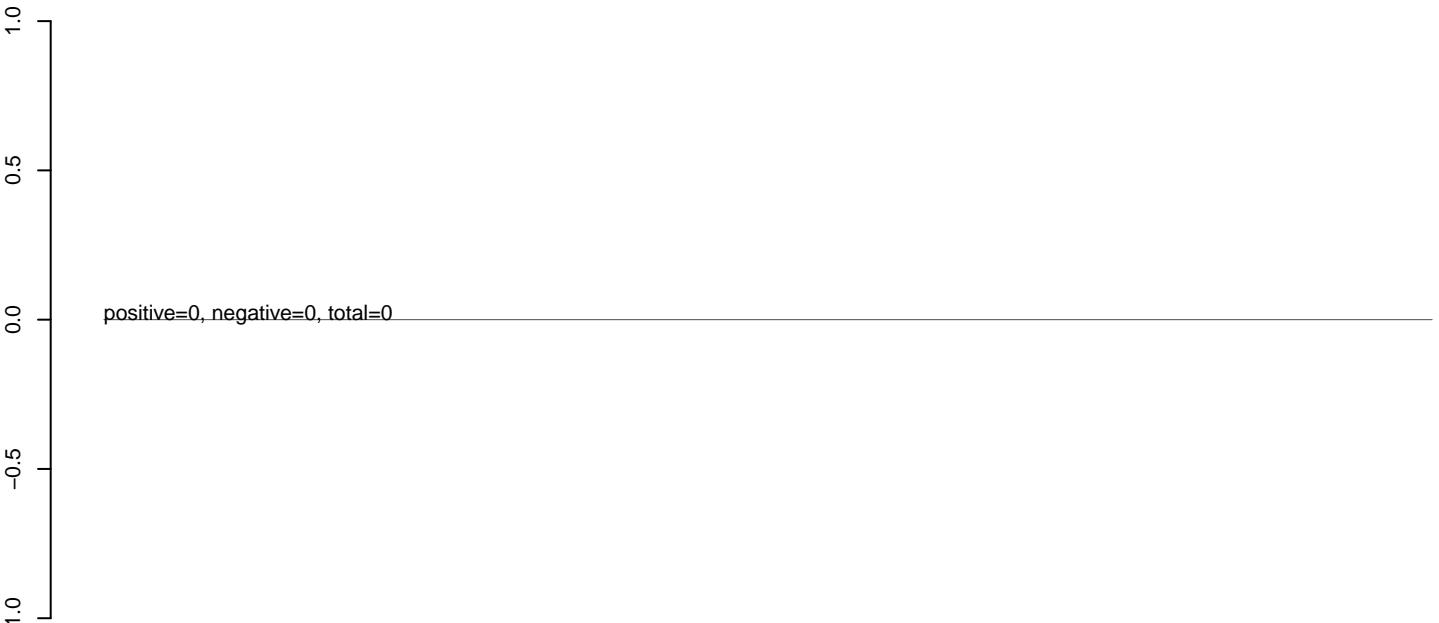
CuQuin_MG_7dpi3.rep



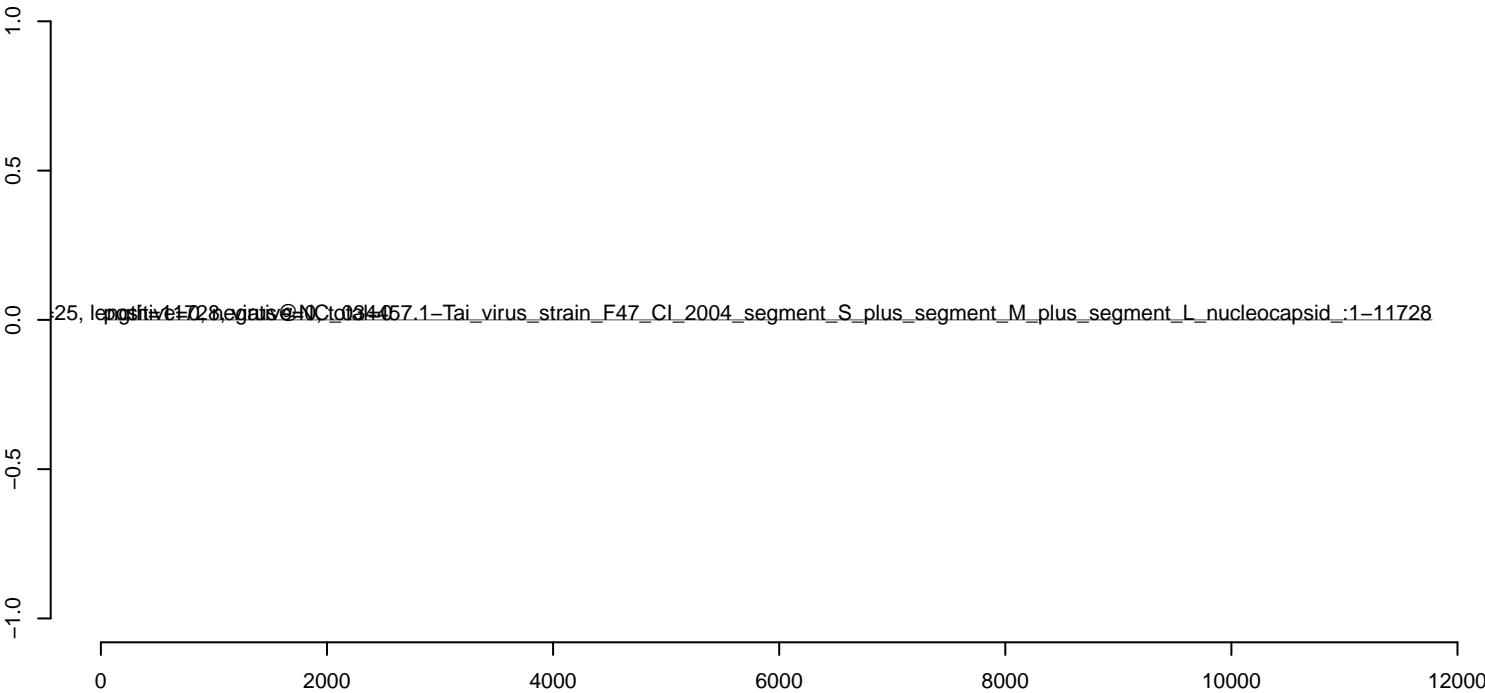
CuQuin_MG_7dpi3.18_23.rep



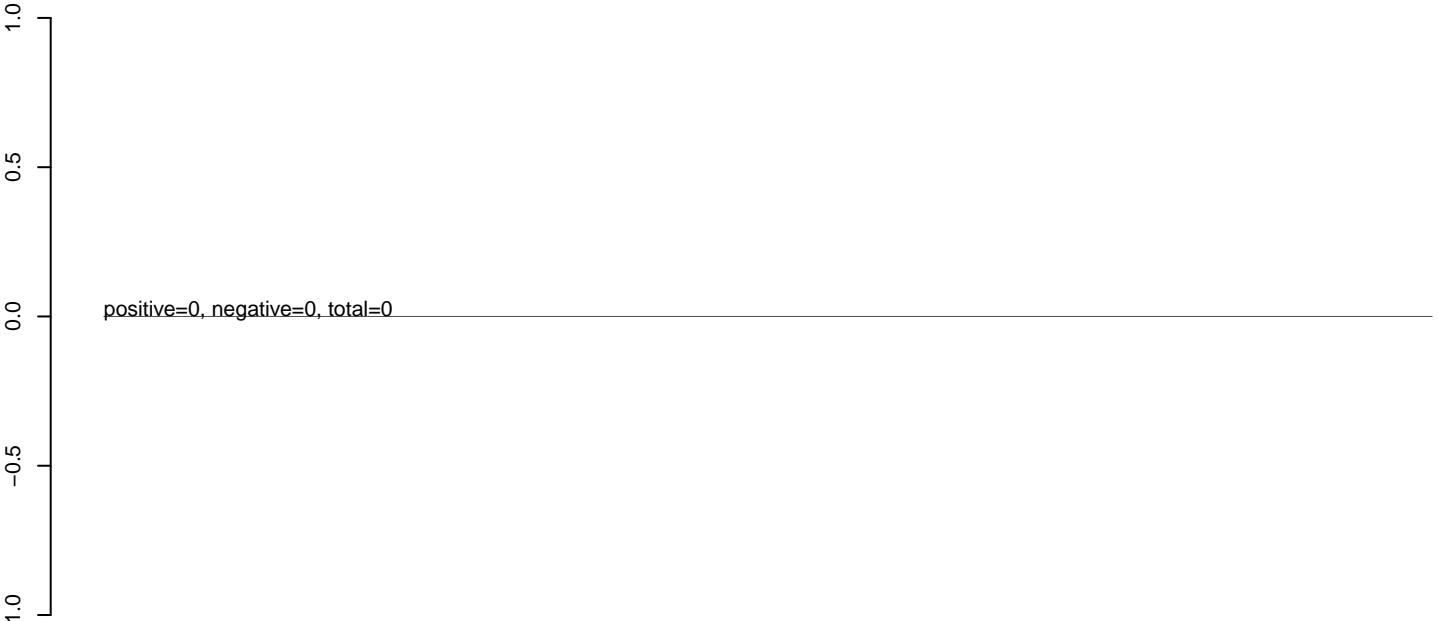
CuQuin_MG_7dpi3.24_35.rep



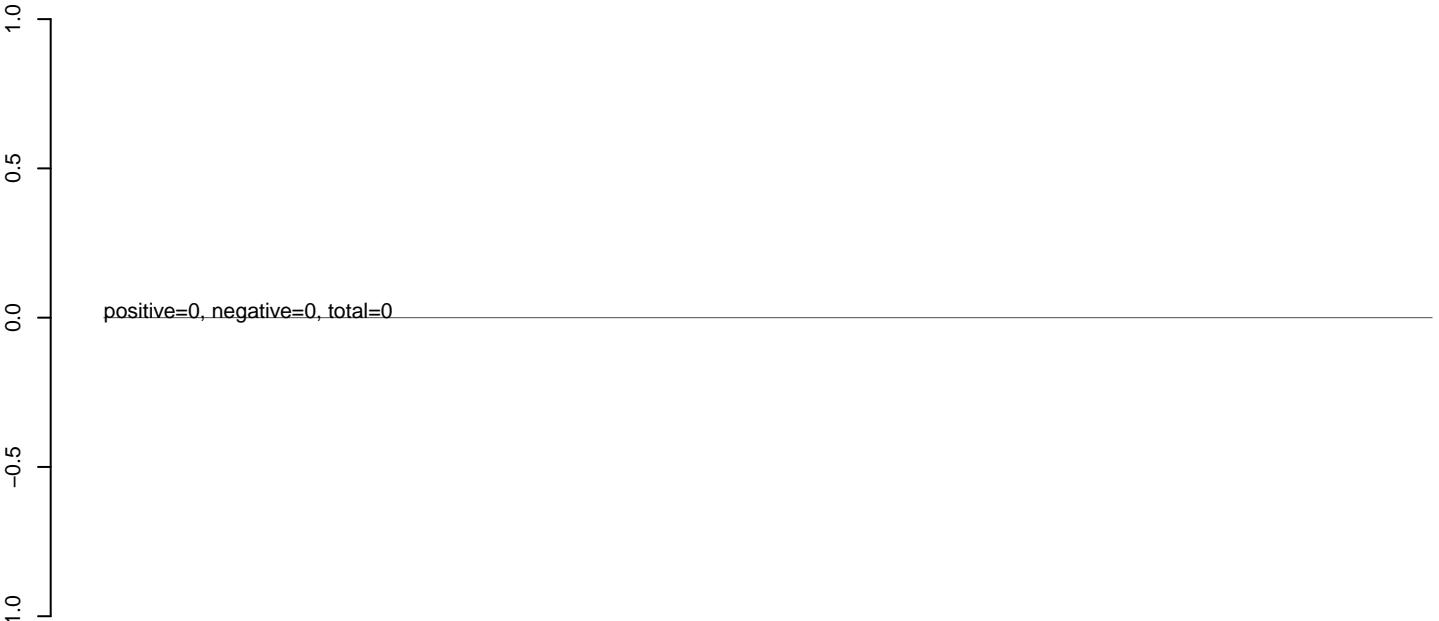
CuQuin_MG_7dpi3.rep



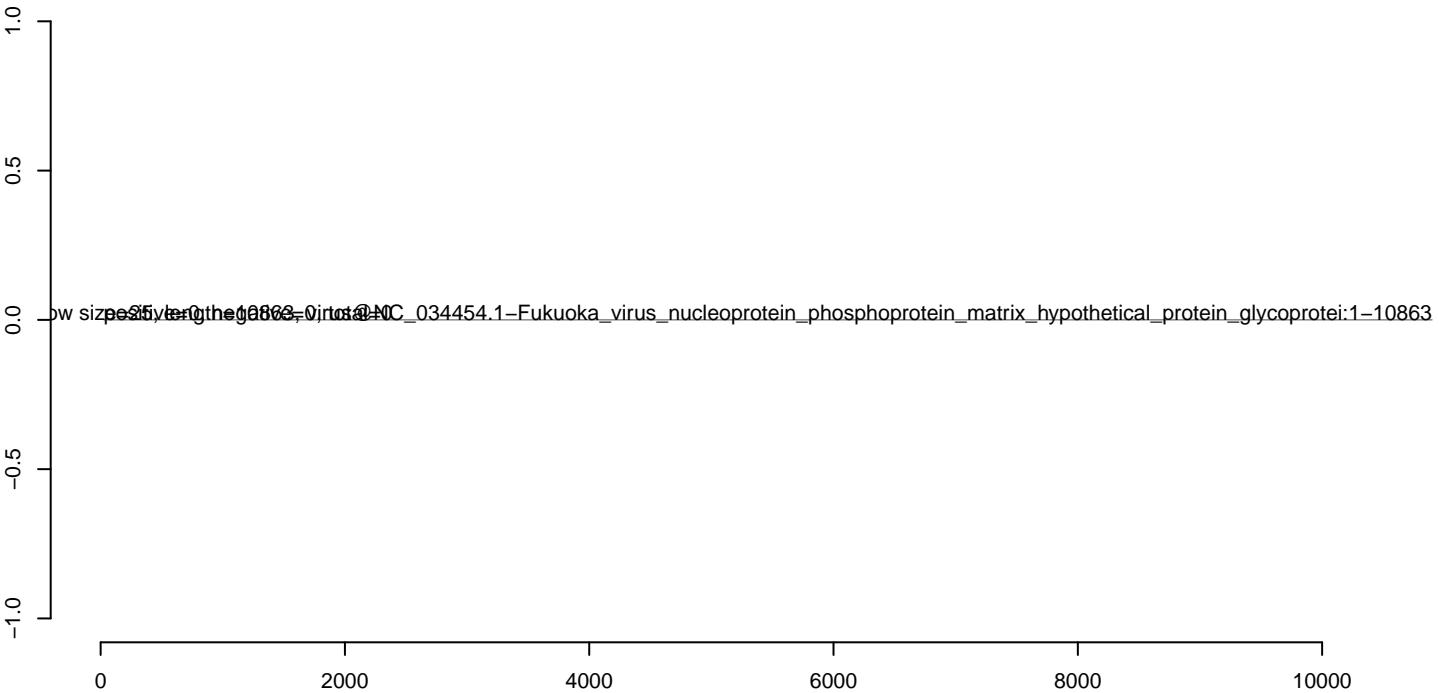
CuQuin_MG_7dpi3.18_23.rep



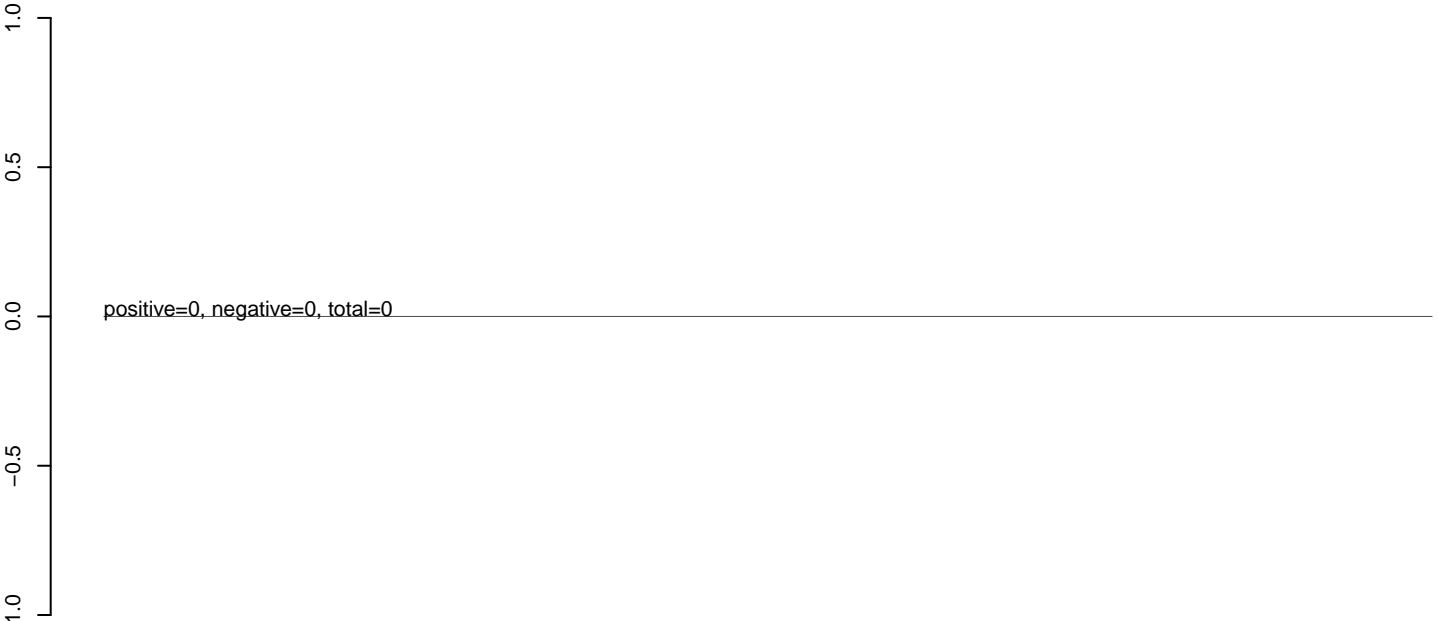
CuQuin_MG_7dpi3.24_35.rep



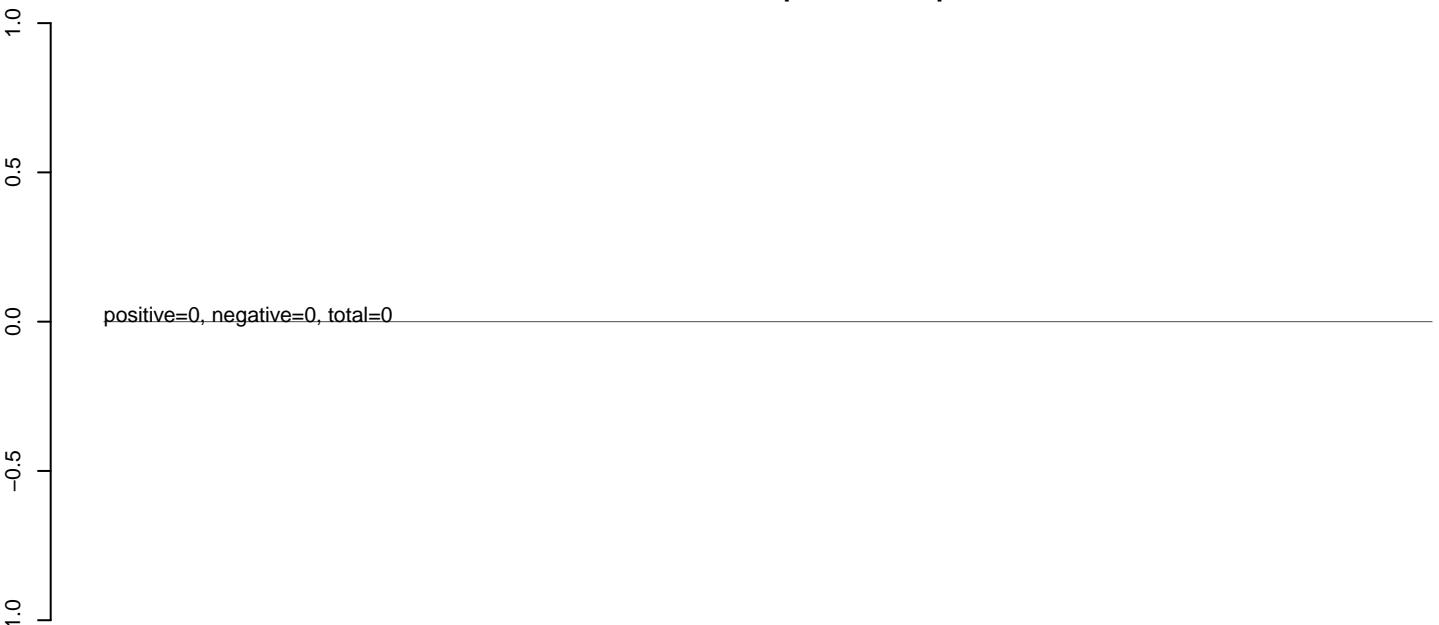
CuQuin_MG_7dpi3.rep



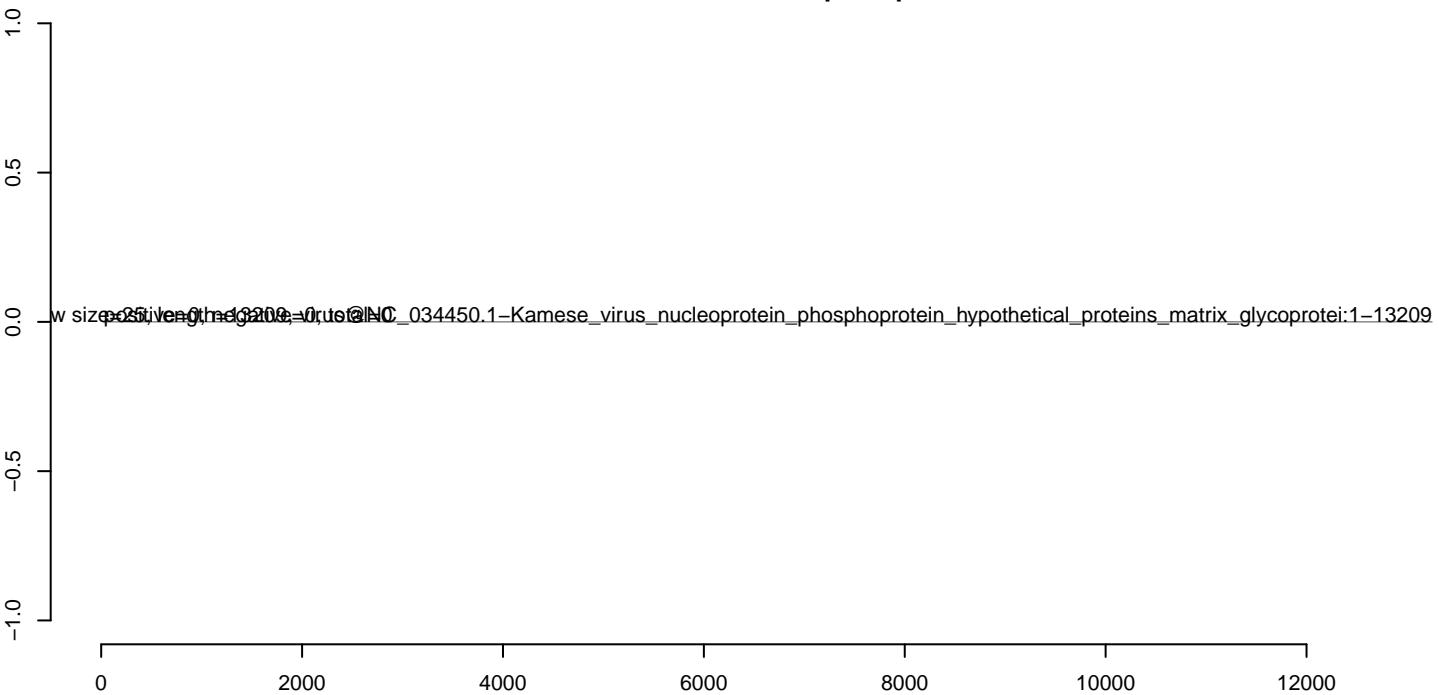
CuQuin_MG_7dpi3.18_23.rep



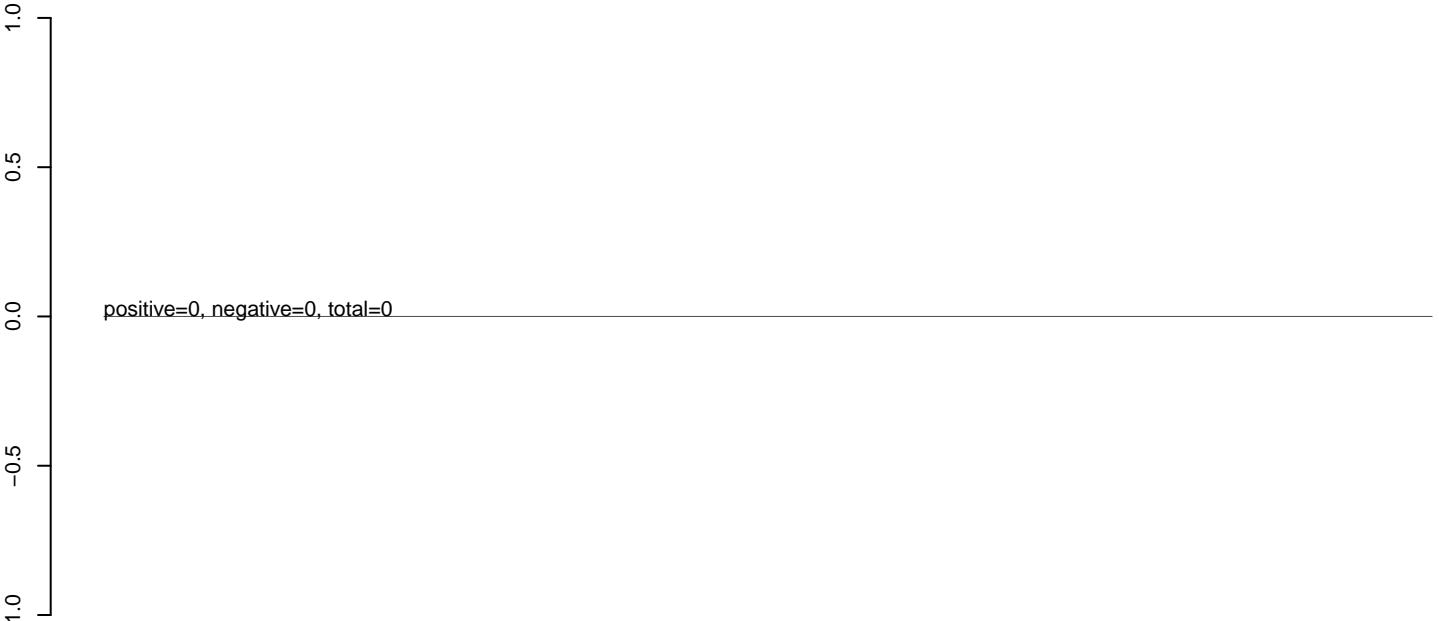
CuQuin_MG_7dpi3.24_35.rep



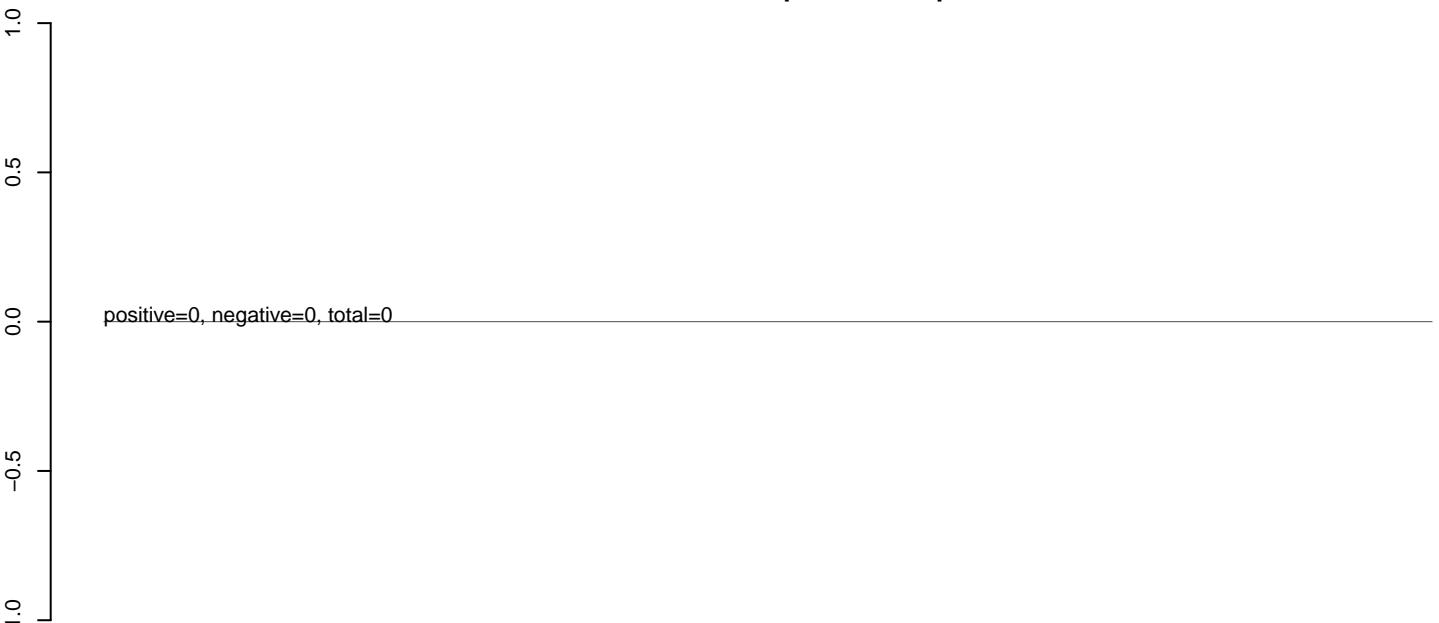
CuQuin_MG_7dpi3.rep



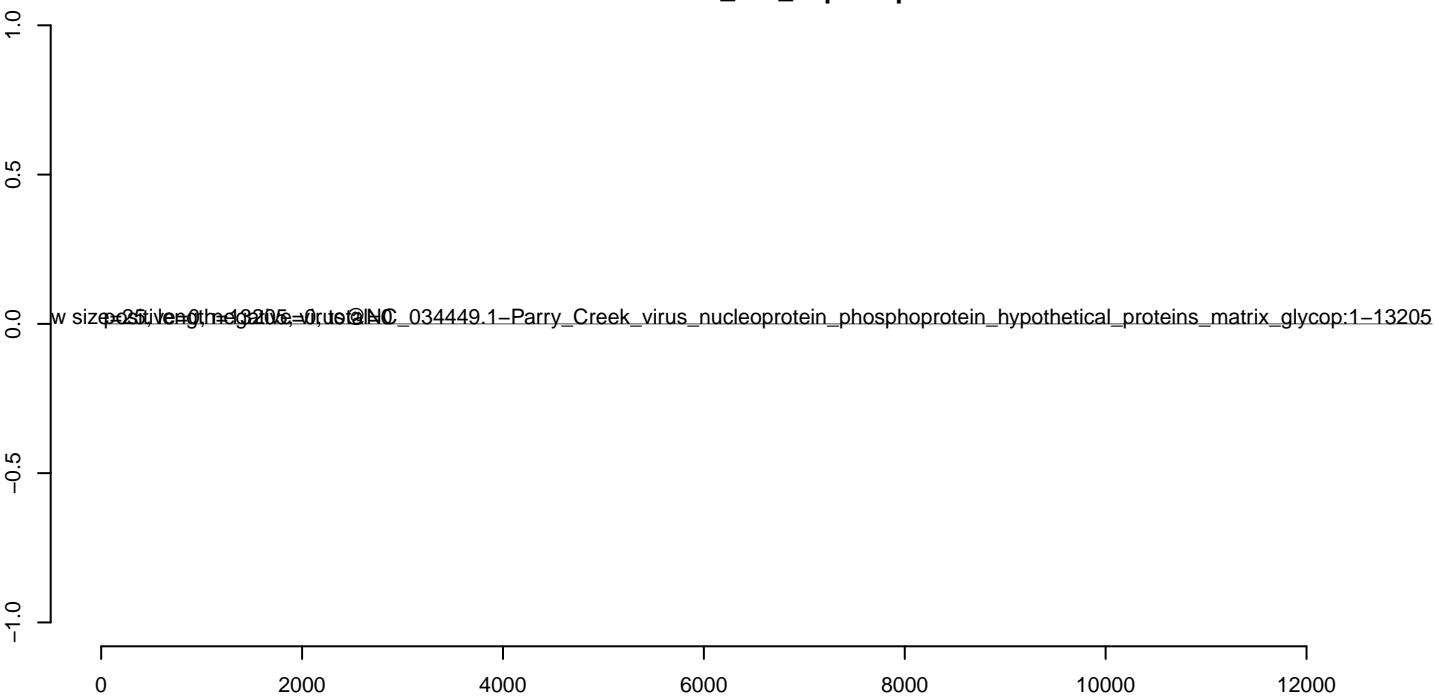
CuQuin_MG_7dpi3.18_23.rep



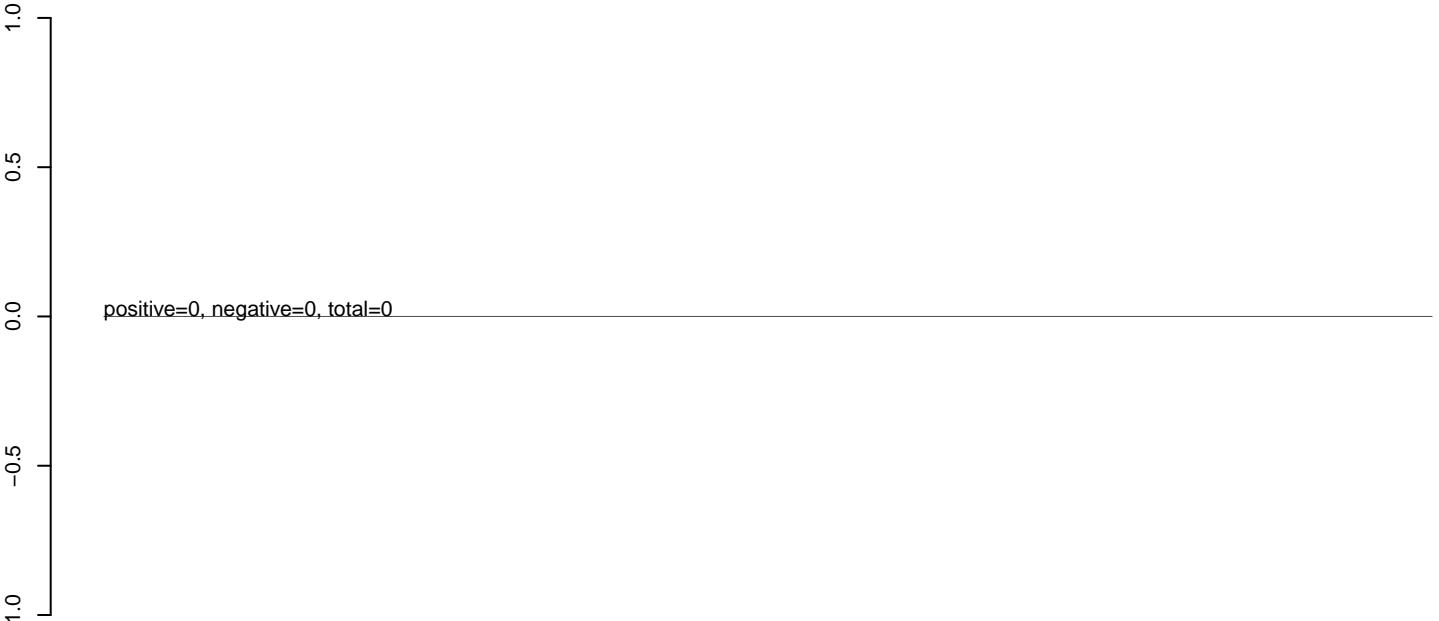
CuQuin_MG_7dpi3.24_35.rep



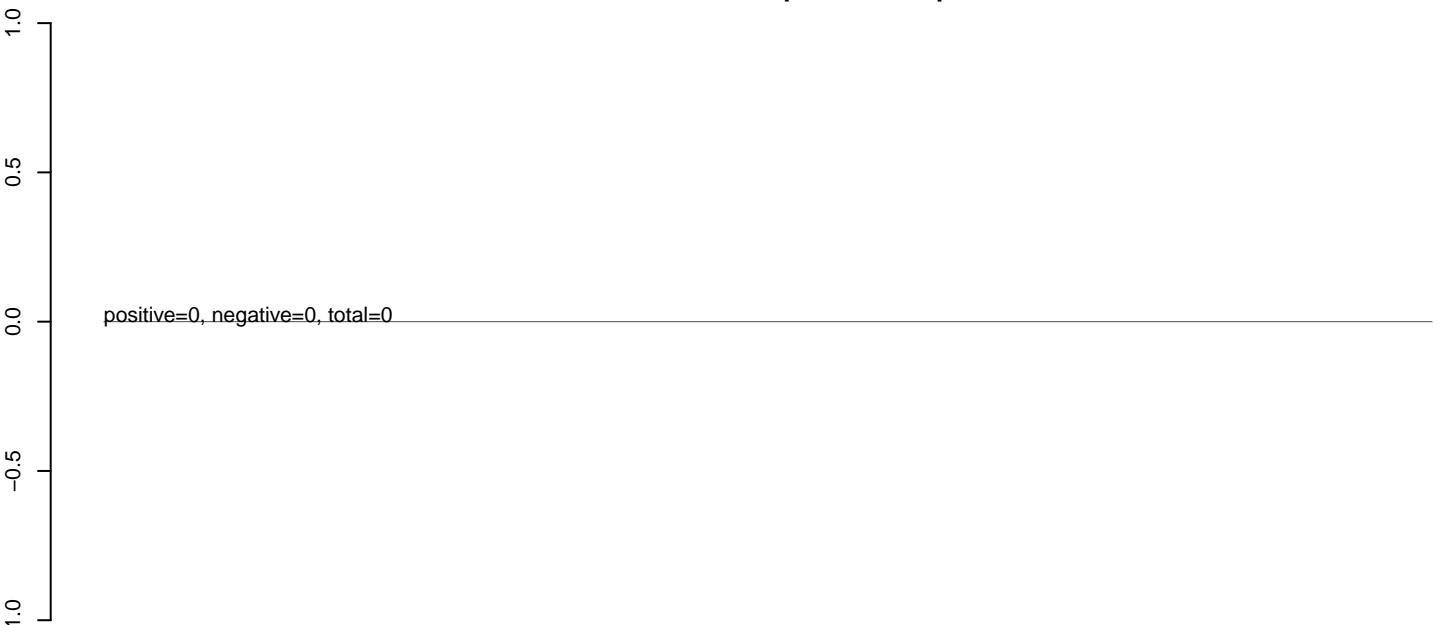
CuQuin_MG_7dpi3.rep



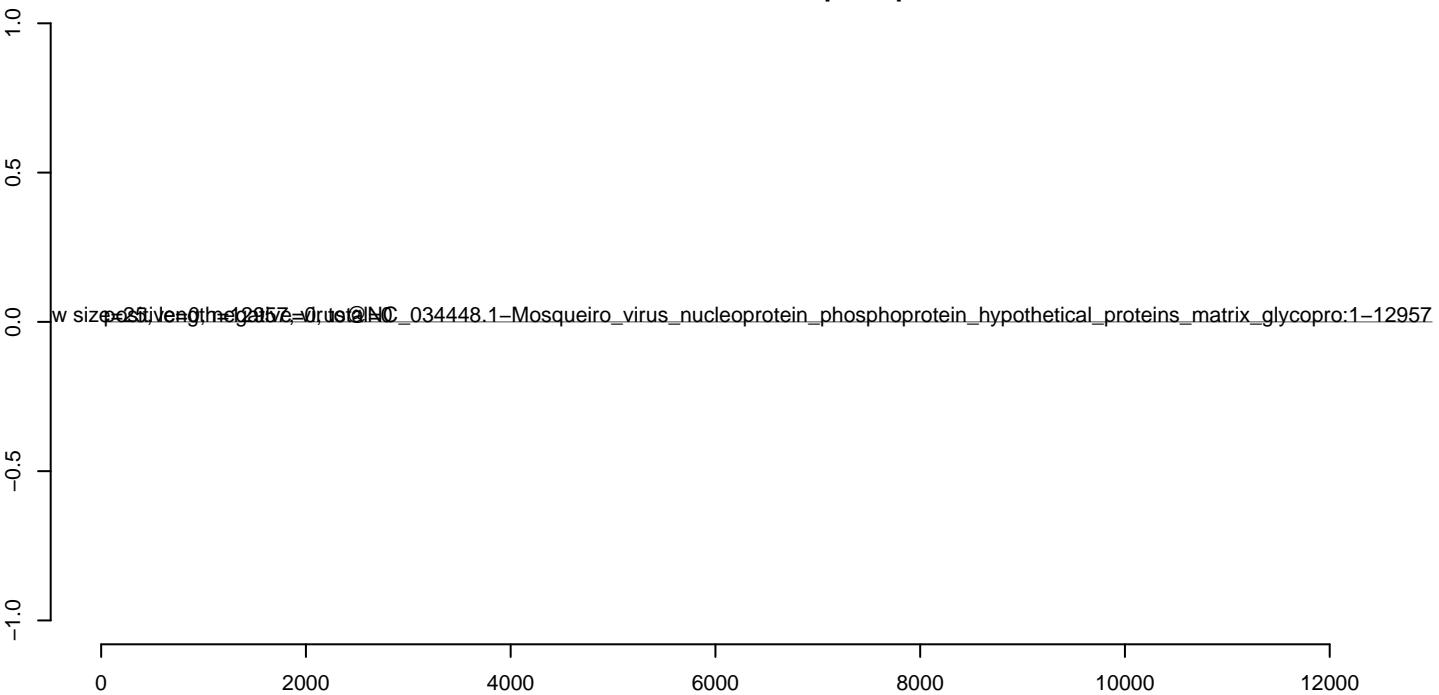
CuQuin_MG_7dpi3.18_23.rep

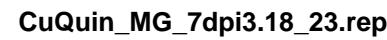


CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep





positive=0, negative=0, total=0



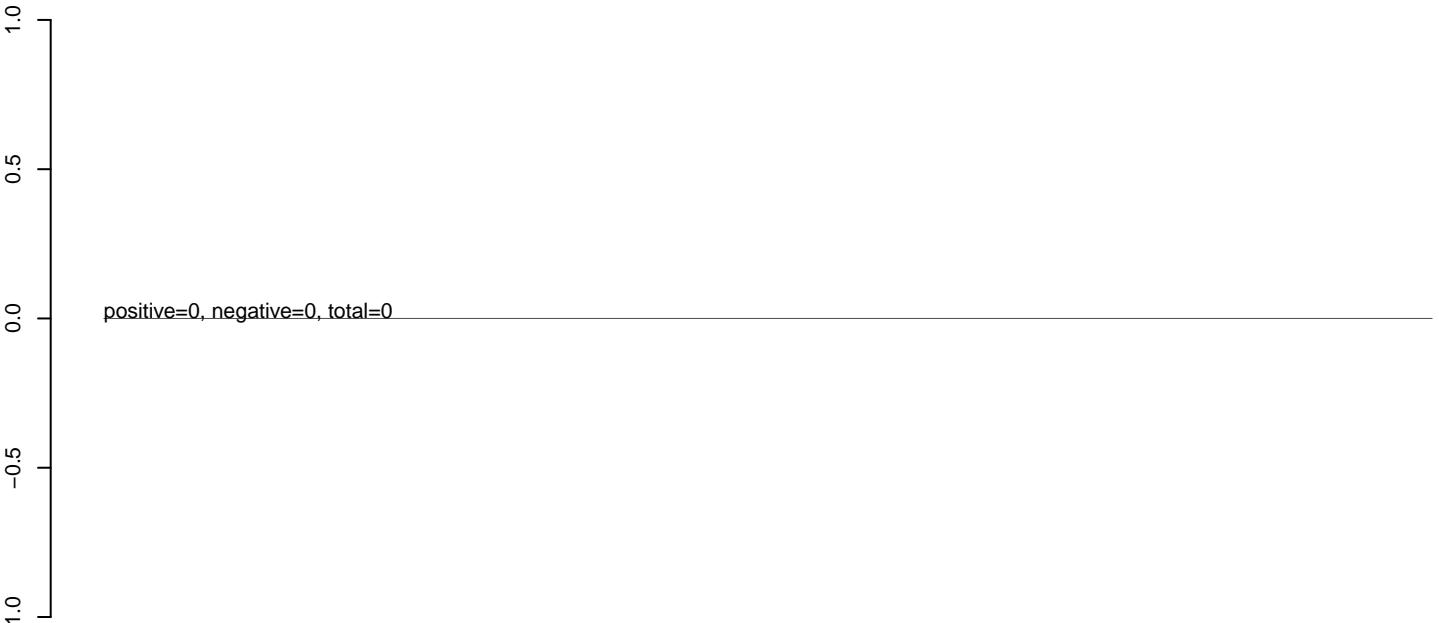
positive=0, negative=0, total=0



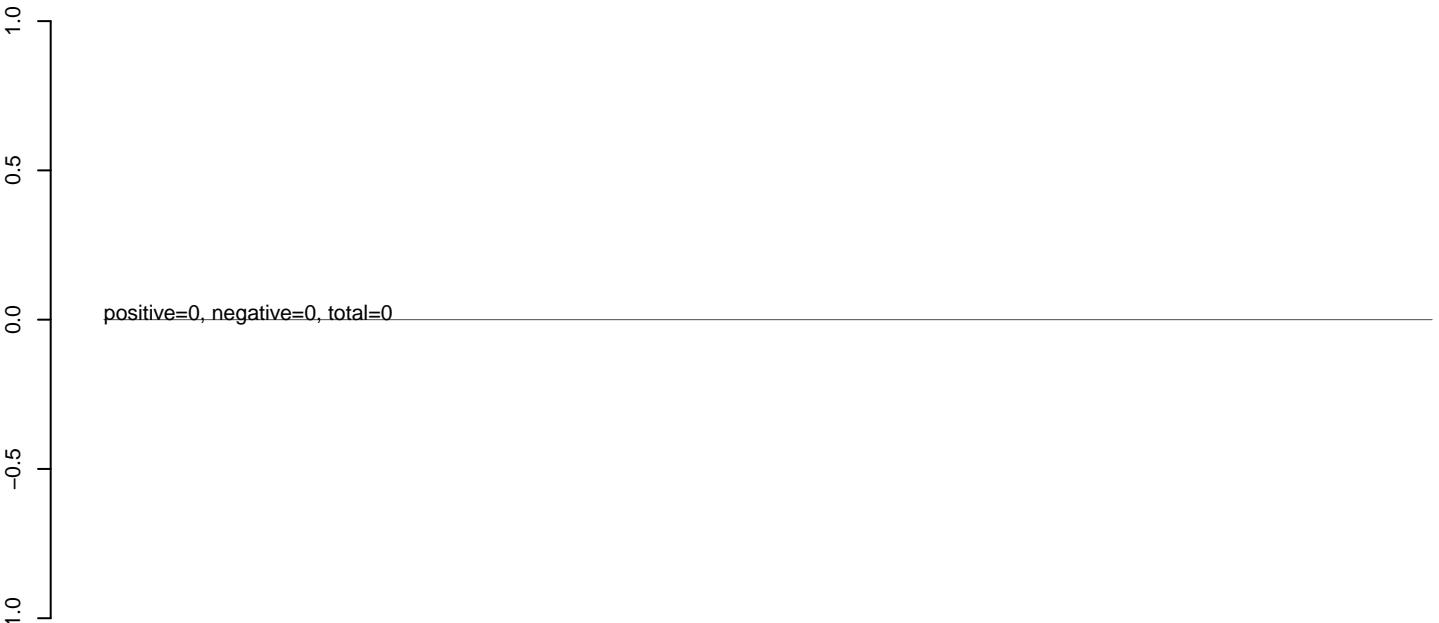
www.ncbi.nlm.nih.gov/blast.ncbi.nlm.nih.gov/Blast.cgi?CMD=Search&QUERY=NC_024447.1; Hert-Pork virus nucleoprotein phosphoprotein hypothetical protein matrix glycoprotein 1; 13104

0 2000 4000 6000 8000 10000 12000

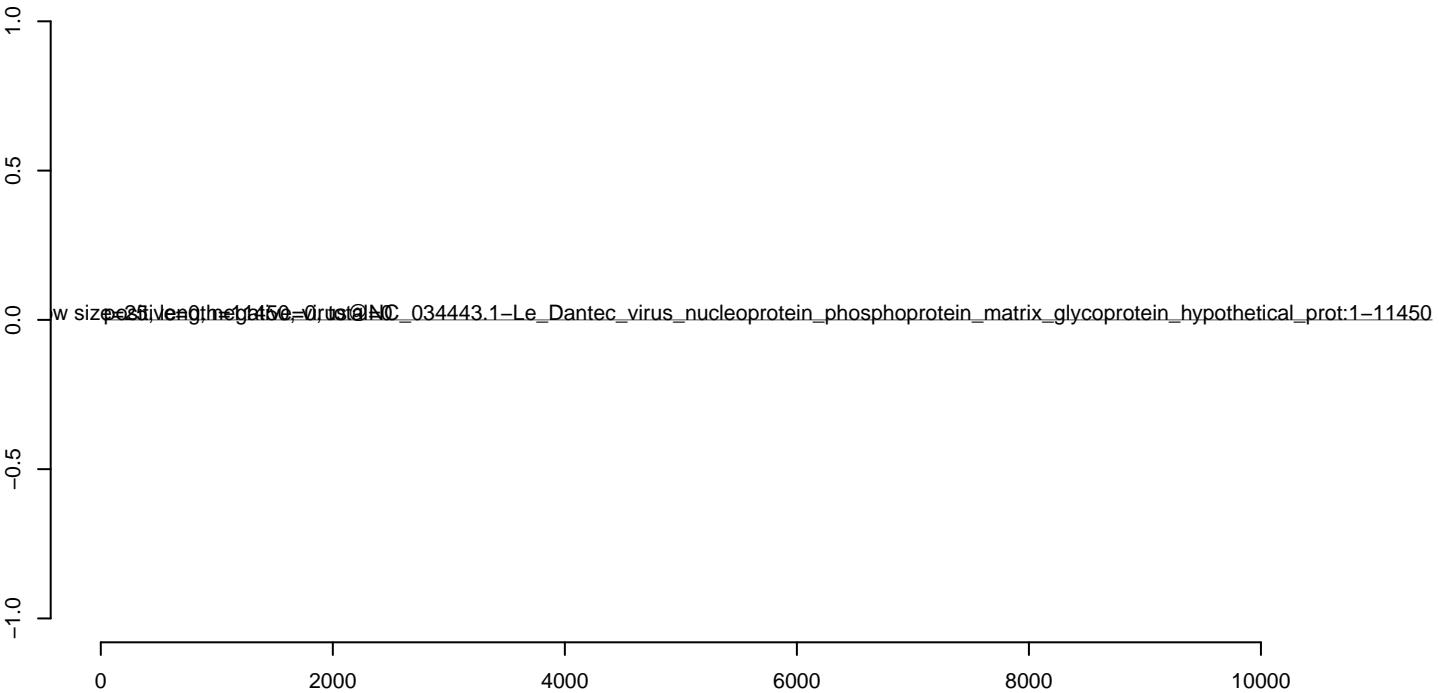
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0
Window size=25, length=10215, virus@NC_033724.1-Kadam_virus_from_Uganda_polyprotein_gene:1-10215

CuQuin_MG_7dpi3.18_23.rep

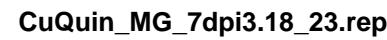
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 window size=25, length=10266, virus@NC_033721.1-Meaban_virus_from_France_polyprotein_gene:1-10266



positive=0, negative=0, total=0



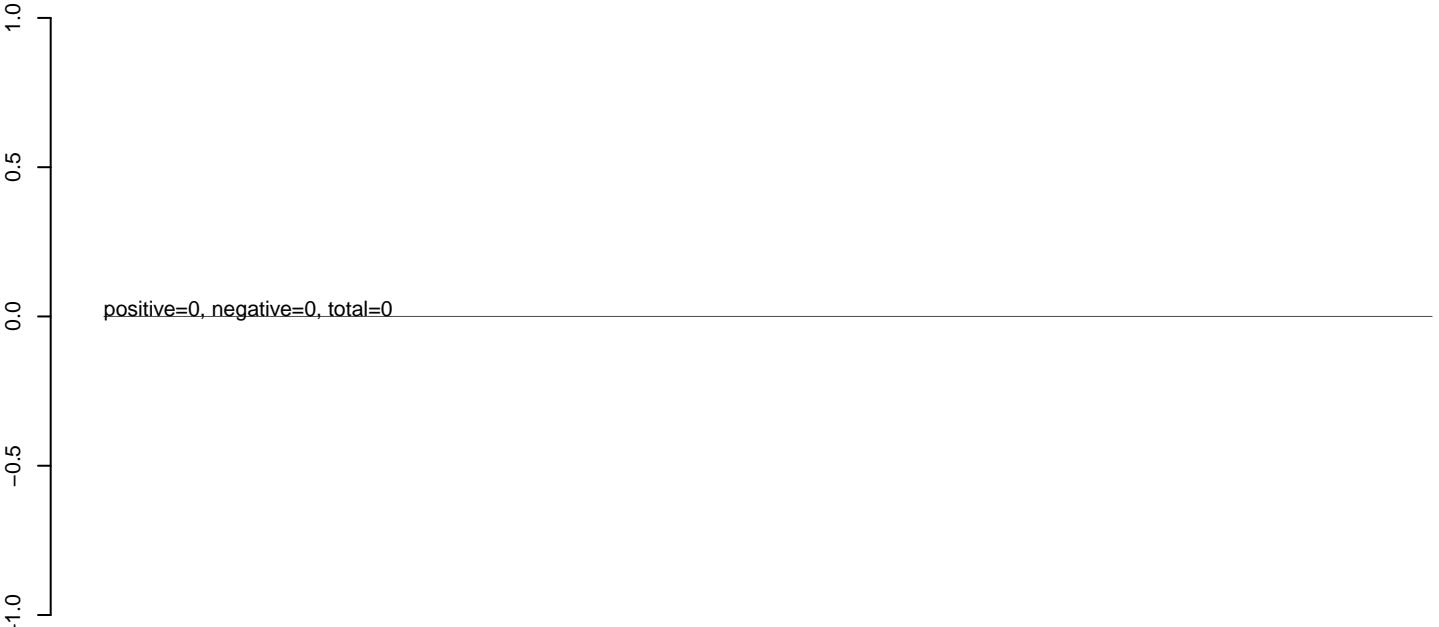
positive=0, negative=0, total=0



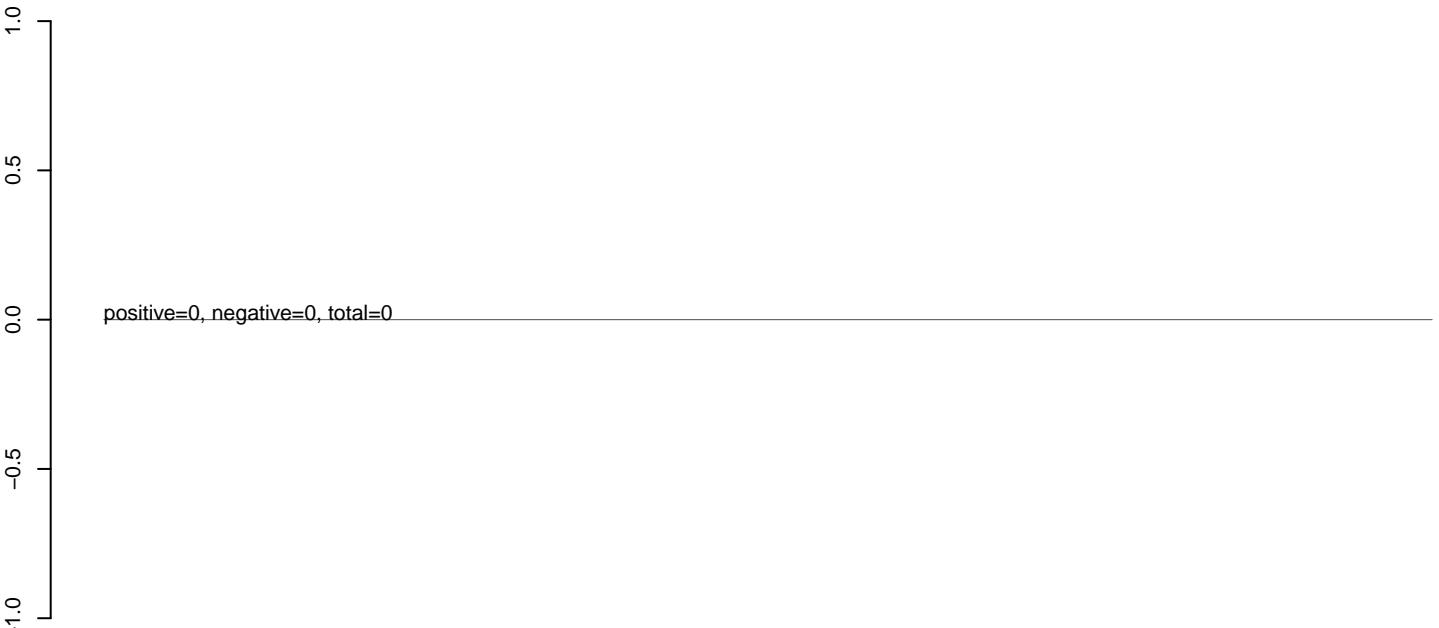
polypeptide size:25.6 kDa; P558. virus@NC_003716.1 Zhejiang mosquito virus 1 strain mosZJ35497 hypothetical protein gene:1-9558

points, single, may

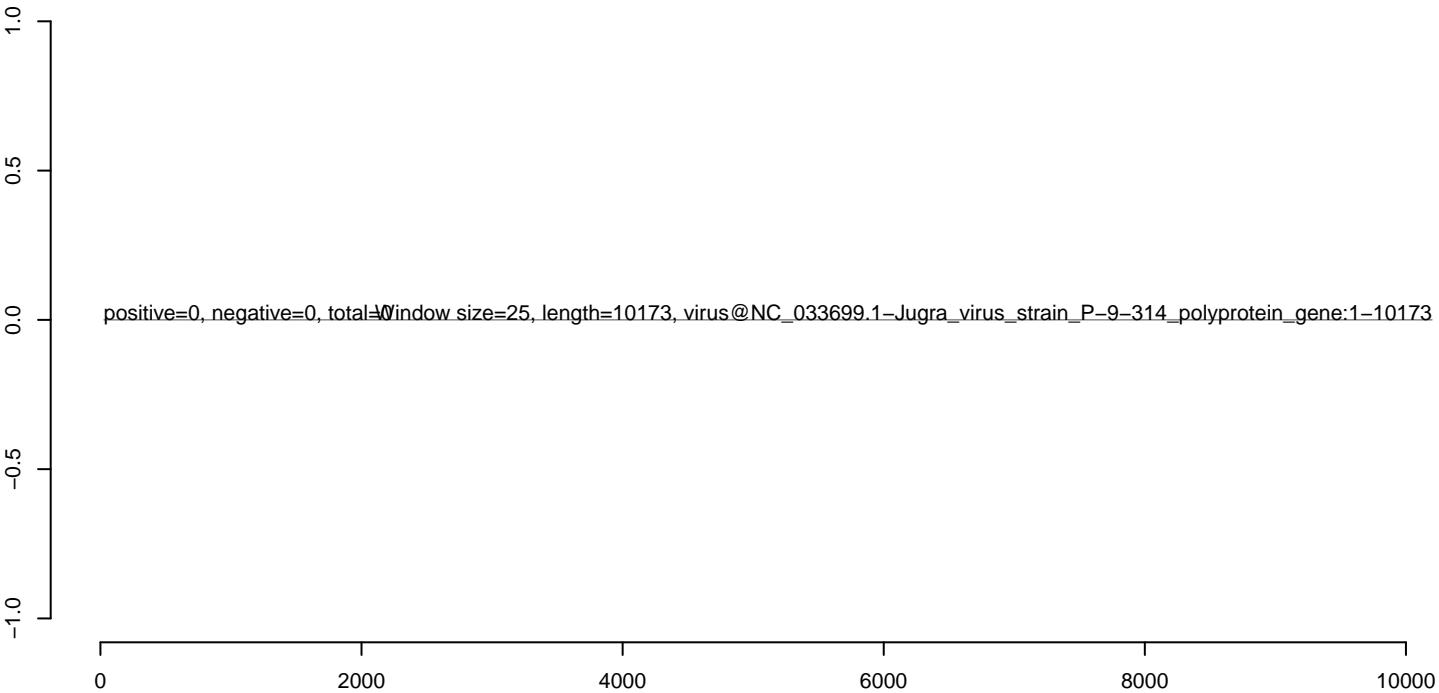
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

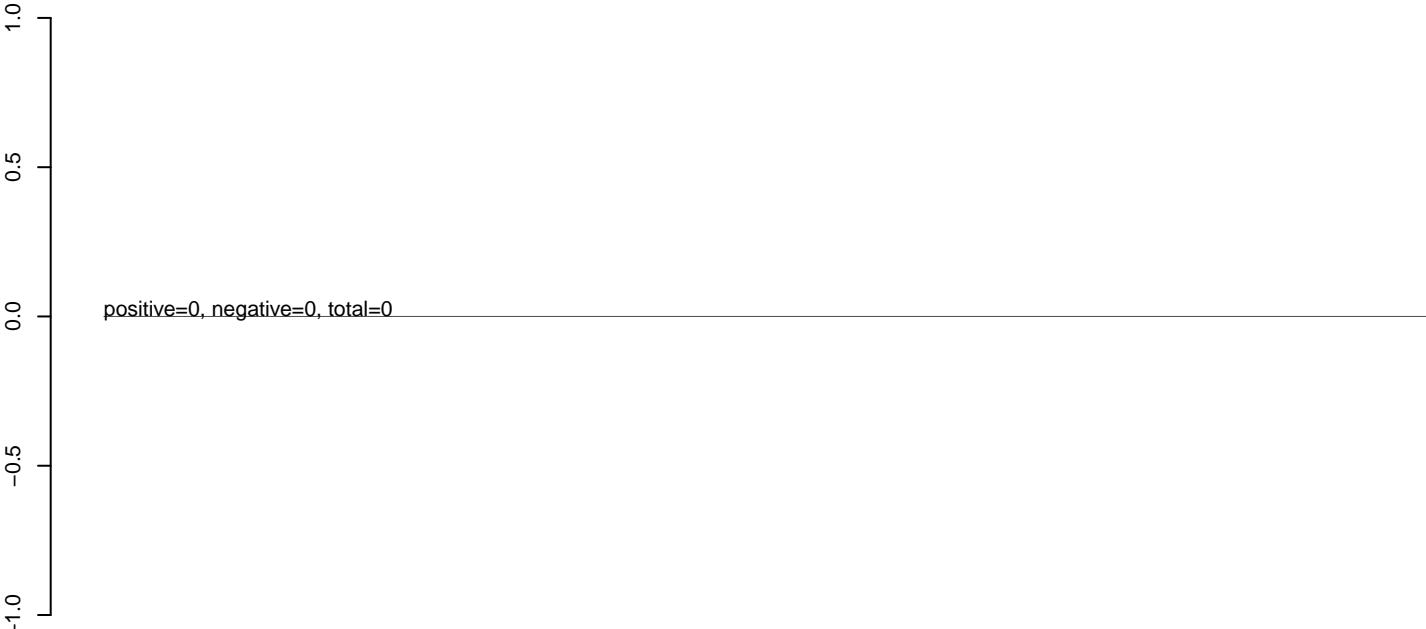
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

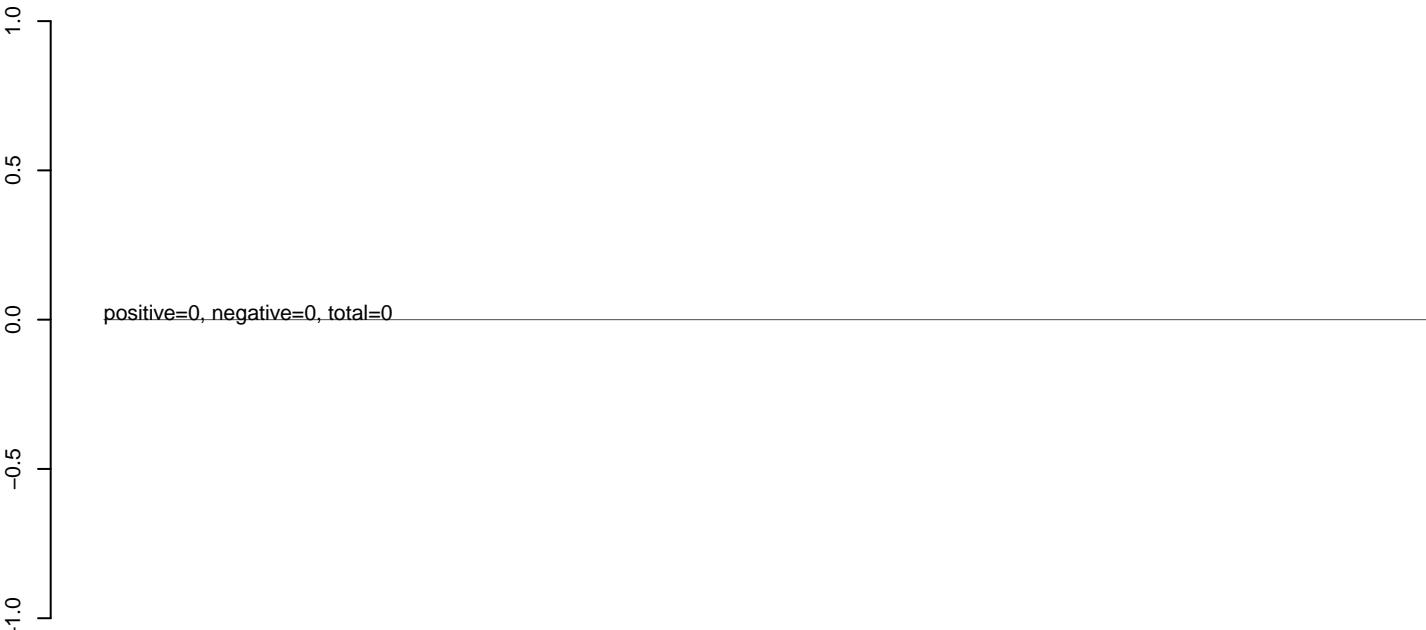
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=25, length=10173, virus@NC_033697.1-Saboya_virus_strain_Dak_AR_D4600_polyprotein_gene:1-10173

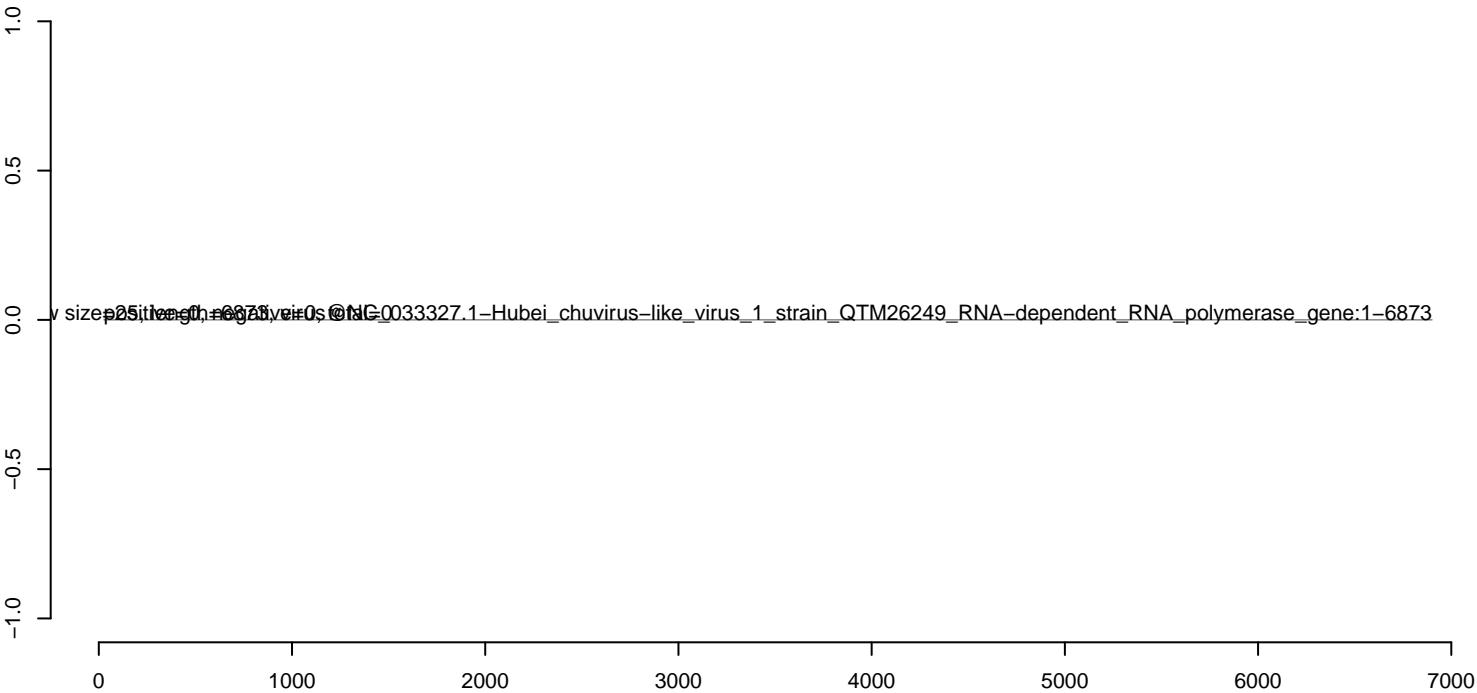
CuQuin_MG_7dpi3.18_23.rep



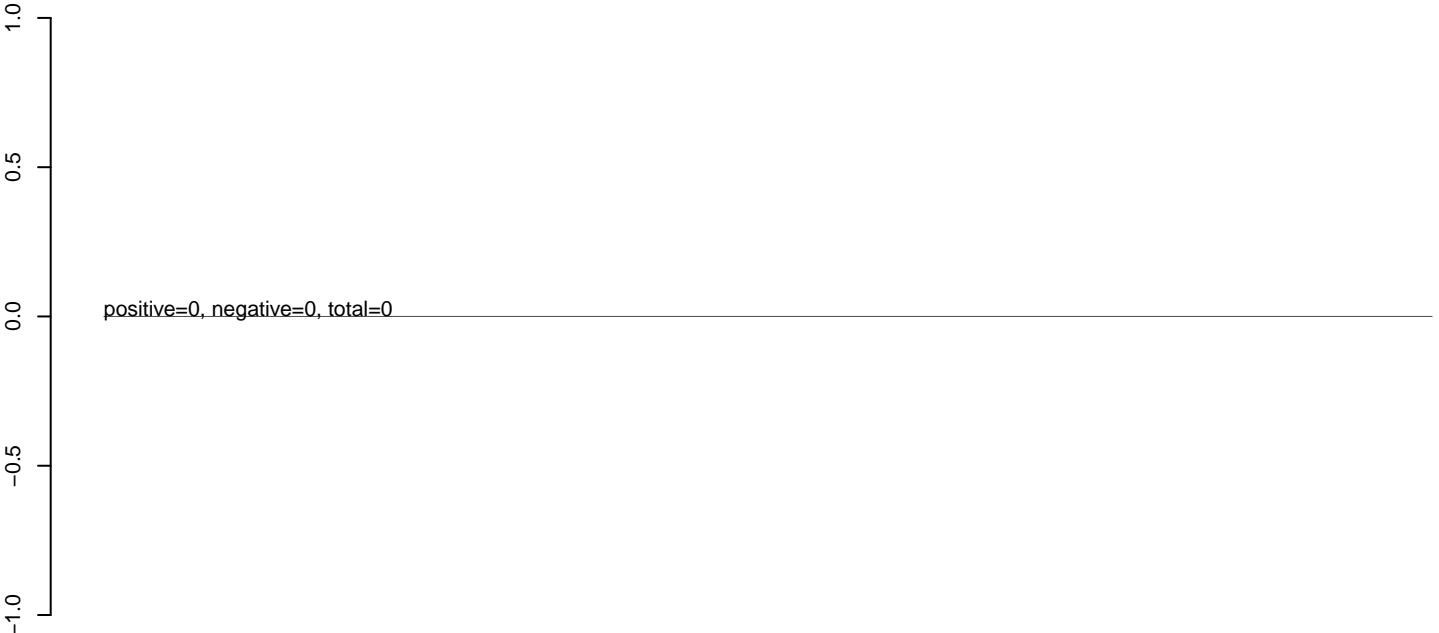
CuQuin_MG_7dpi3.24_35.rep



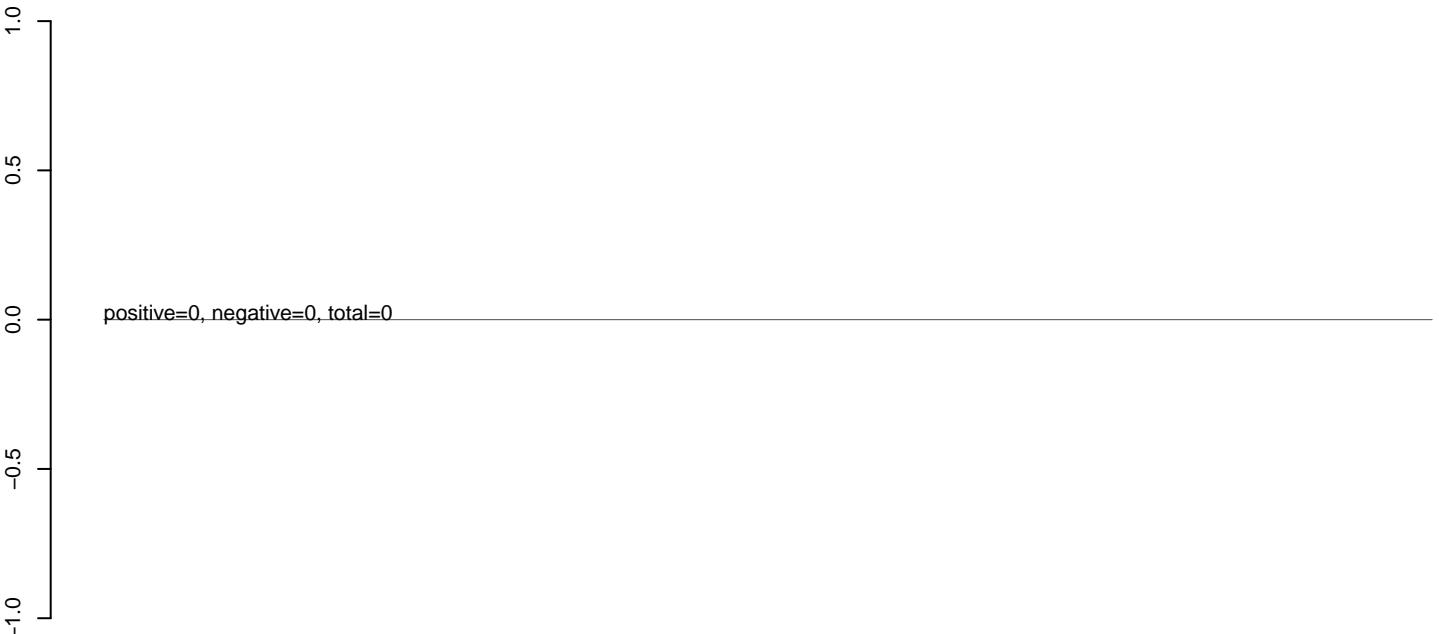
CuQuin_MG_7dpi3.rep



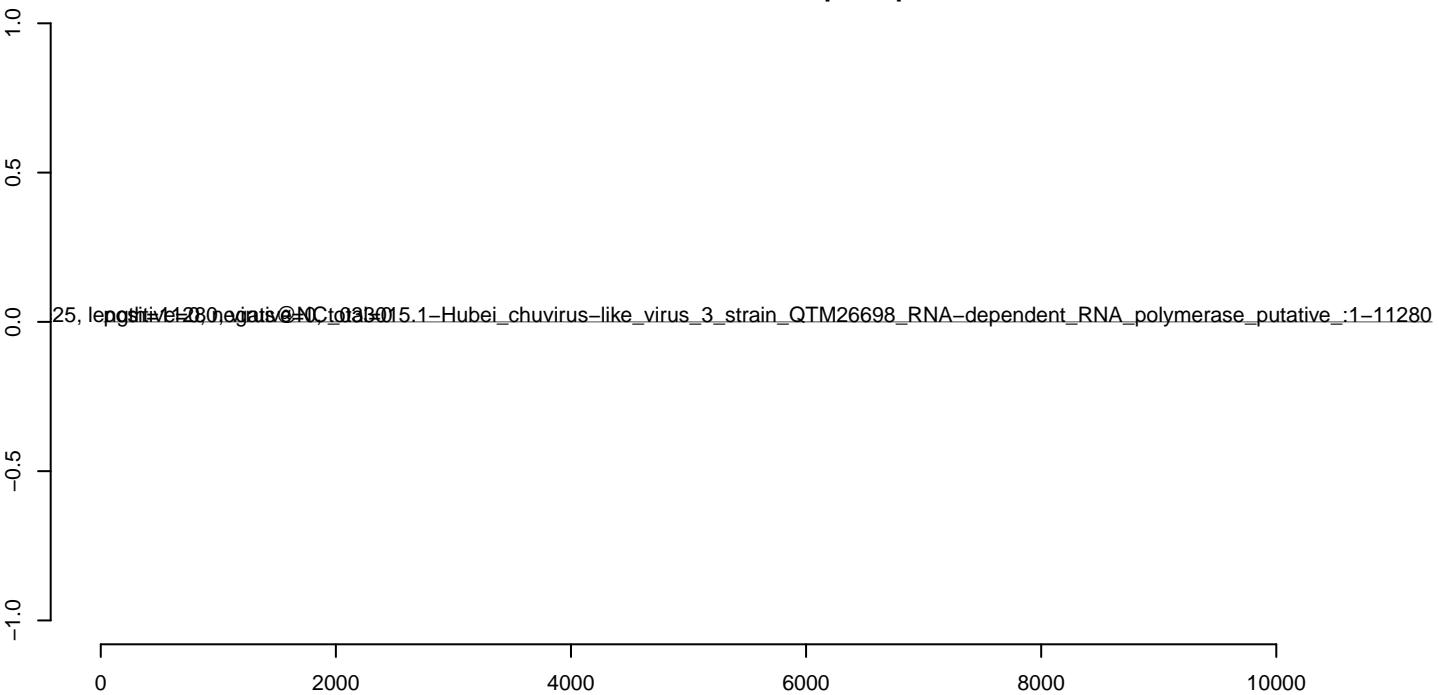
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

size=25 length=3226 positive=0 negative=0
NC_032491.1_Zhejiang_mosquito_virus_3_strain_mosZJ35354_hypothetical_protein_and_RNA-dependent:1-3226

0 500 1000 1500 2000 2500 3000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0, window size=25, length=10869, virus@NC_031957.1-UNVERIFIED_Coot_Bay_virus_strain_EVG5-53:1-10869

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window.size=25, length=11286, virus@NC_031956.1-UNVERIFIED_Balsa_virus_strain_CoB_84:1-11286

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

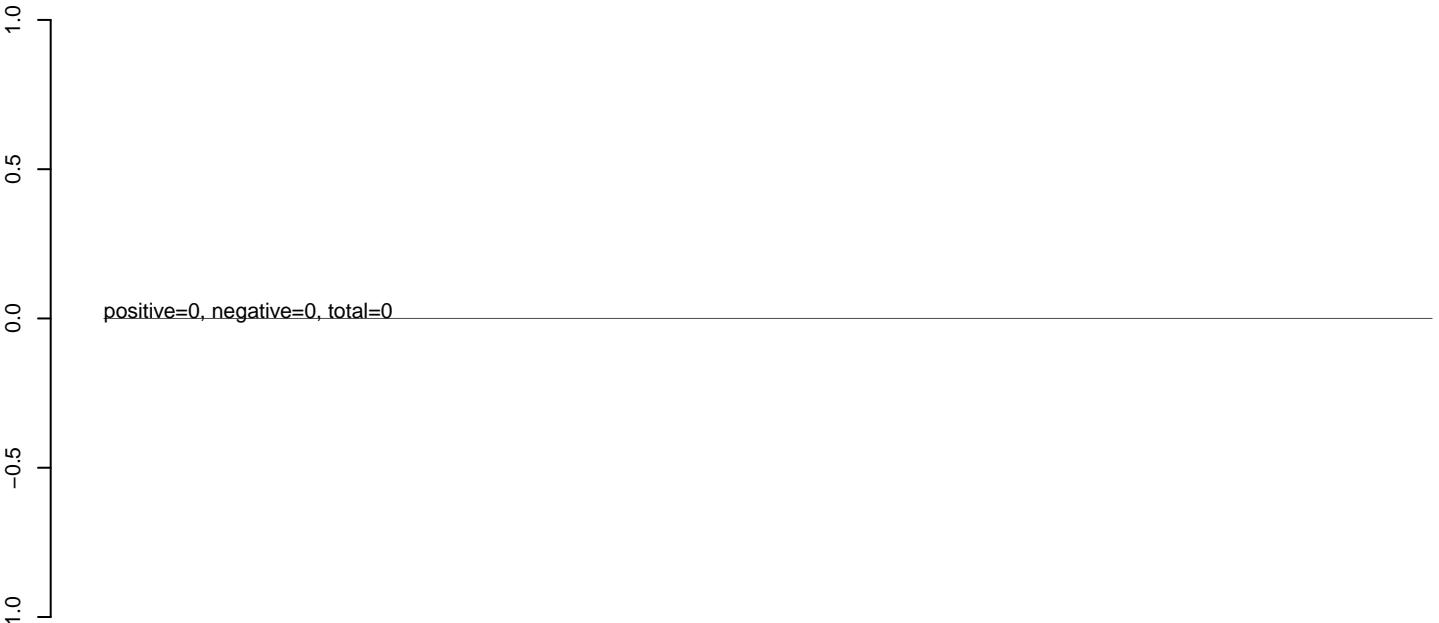
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

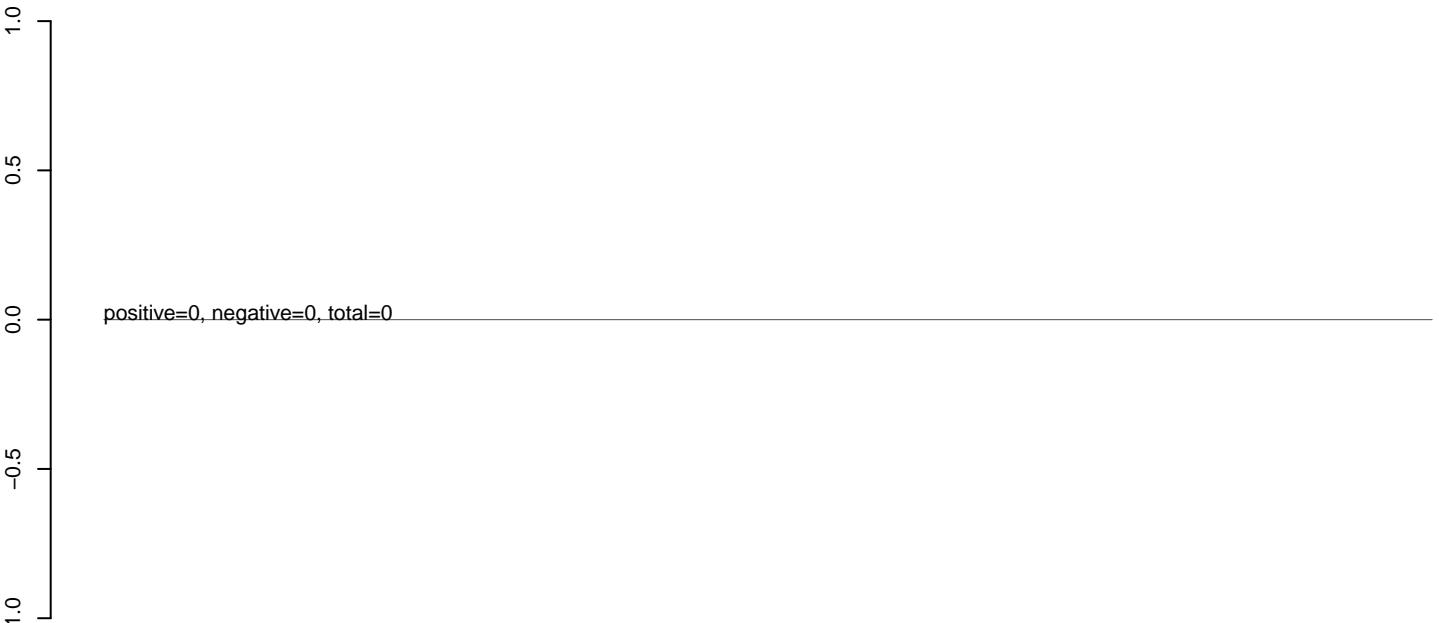
Window size=25, length=10588, virus@NC_031327.1-Anopheles_flavivirus_variant_1:1-10588

0 2000 4000 6000 8000 10000

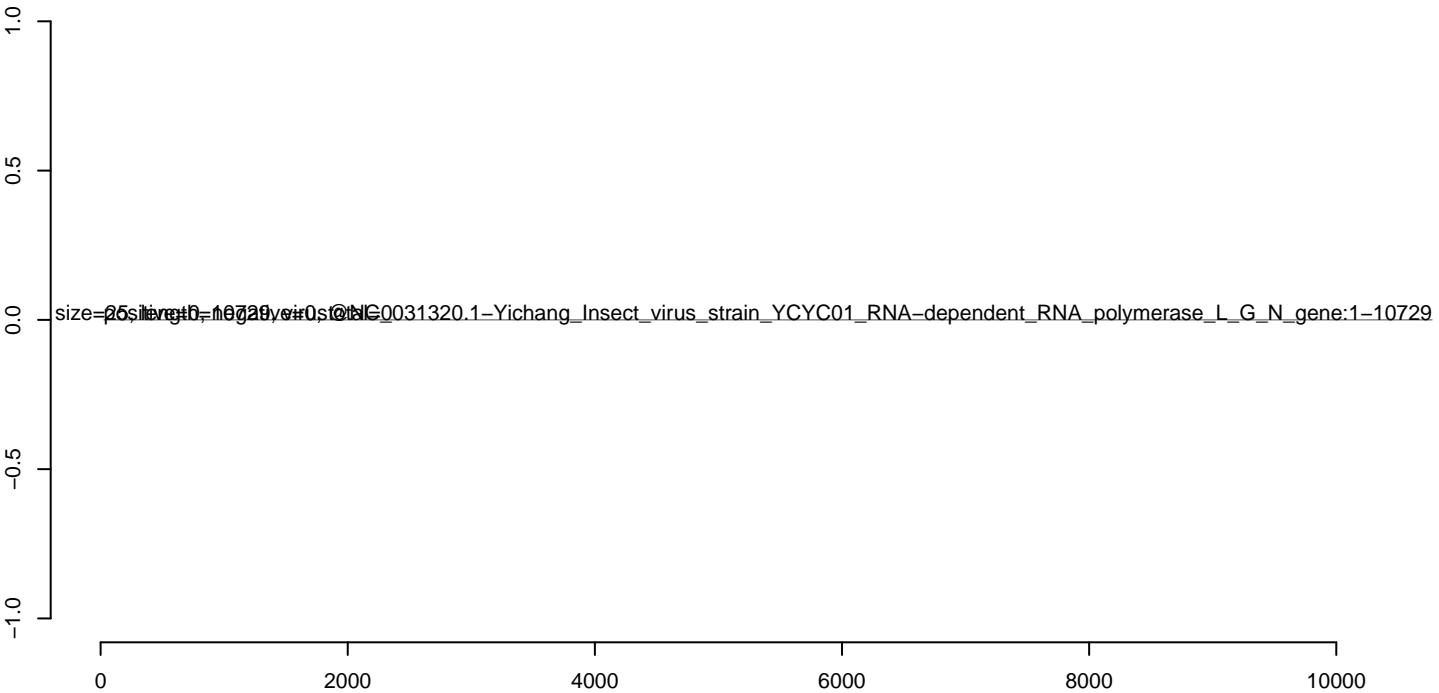
CuQuin_MG_7dpi3.18_23.rep



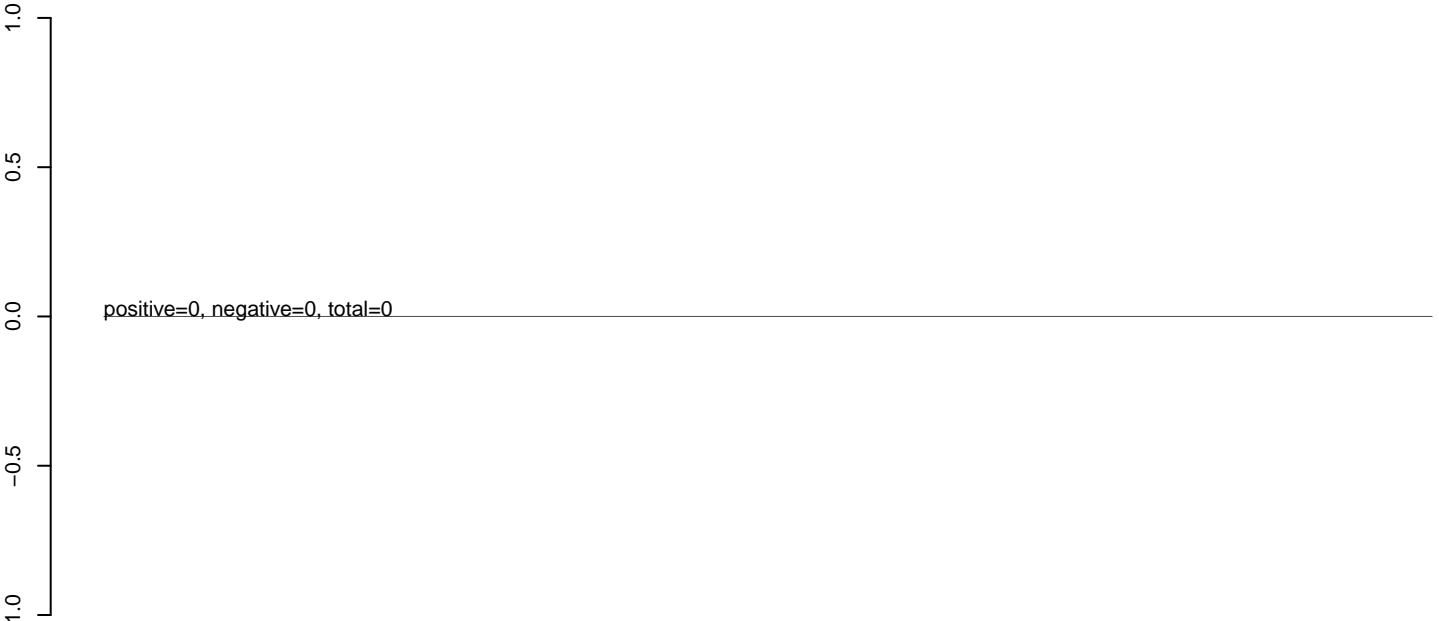
CuQuin_MG_7dpi3.24_35.rep



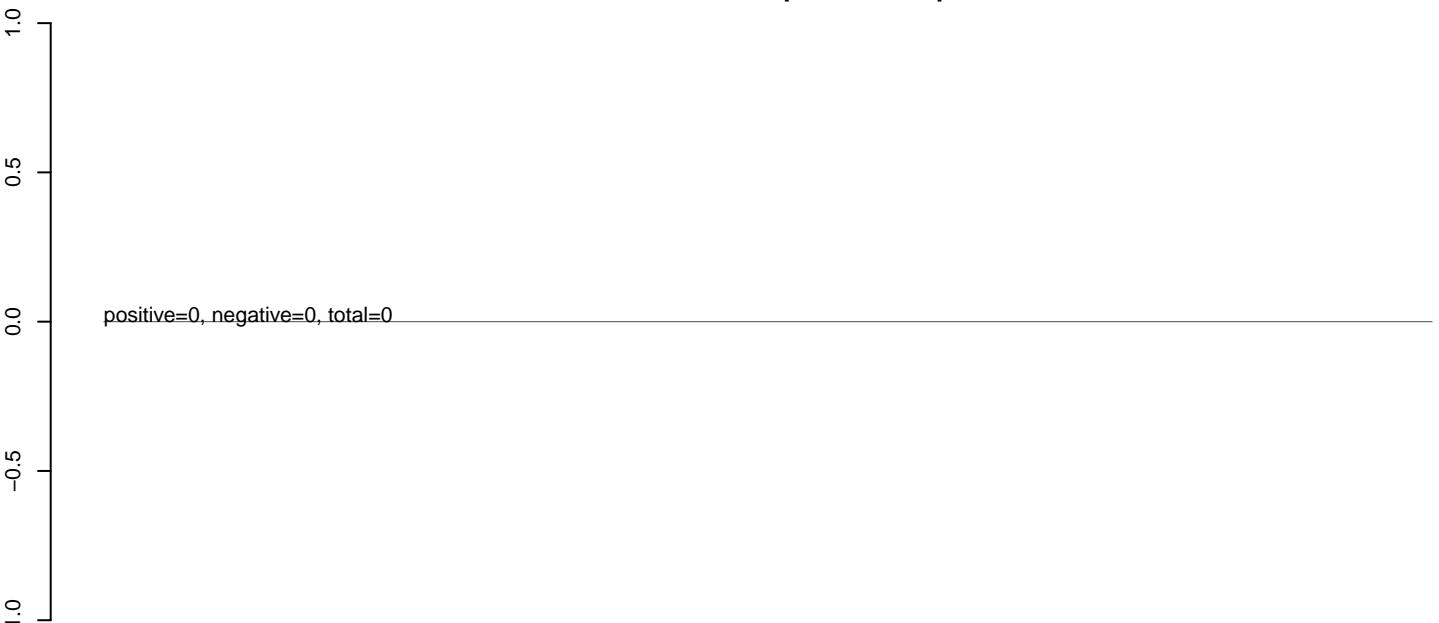
CuQuin_MG_7dpi3.rep



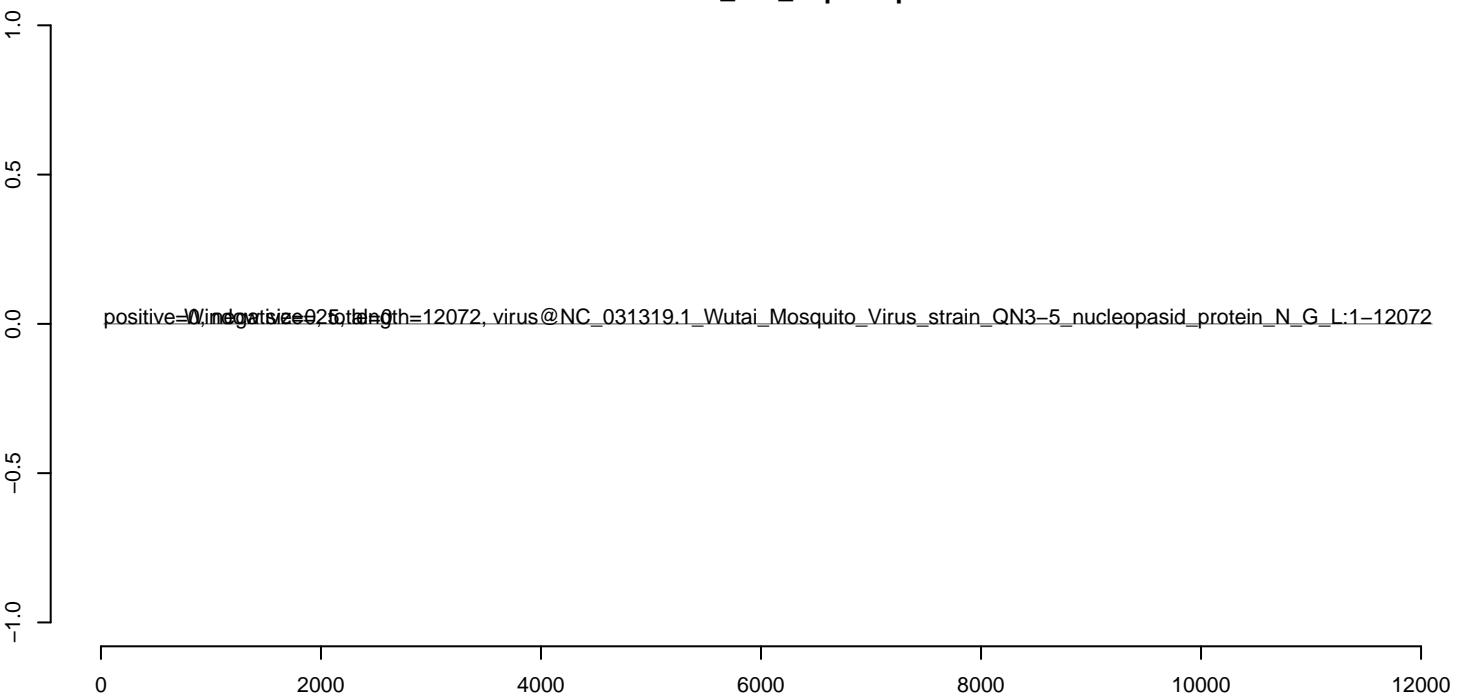
CuQuin_MG_7dpi3.18_23.rep



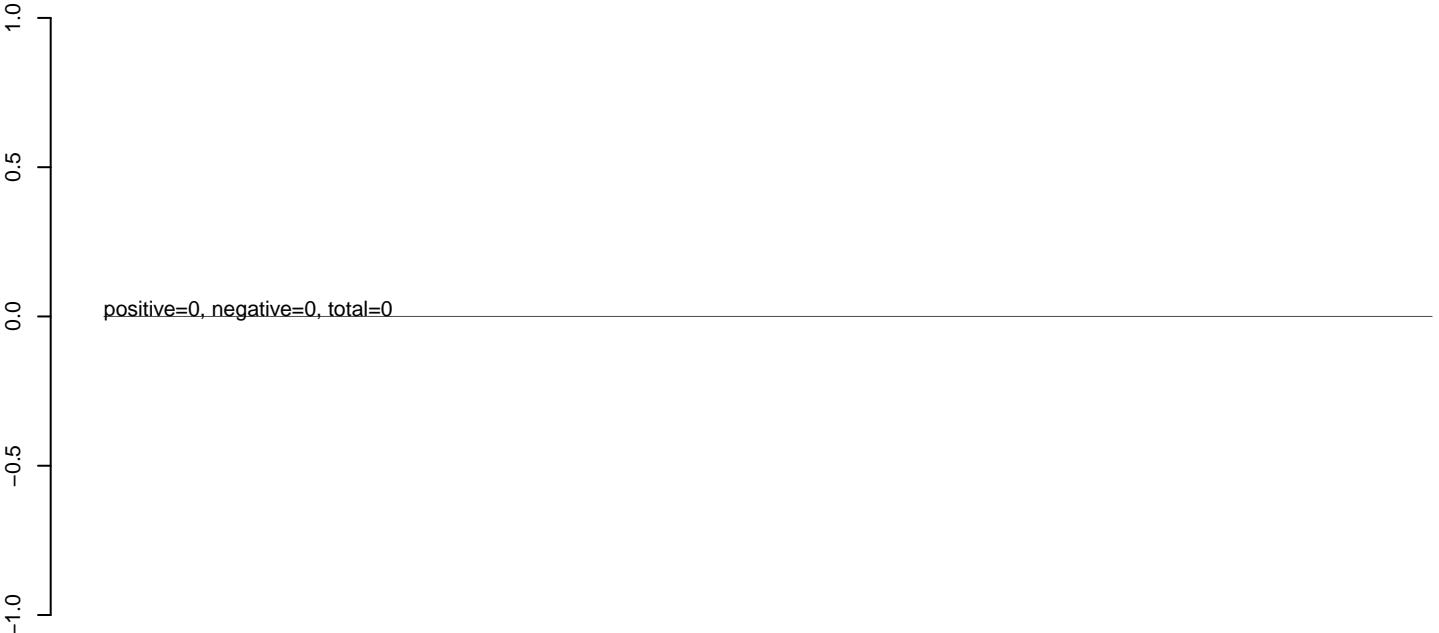
CuQuin_MG_7dpi3.24_35.rep



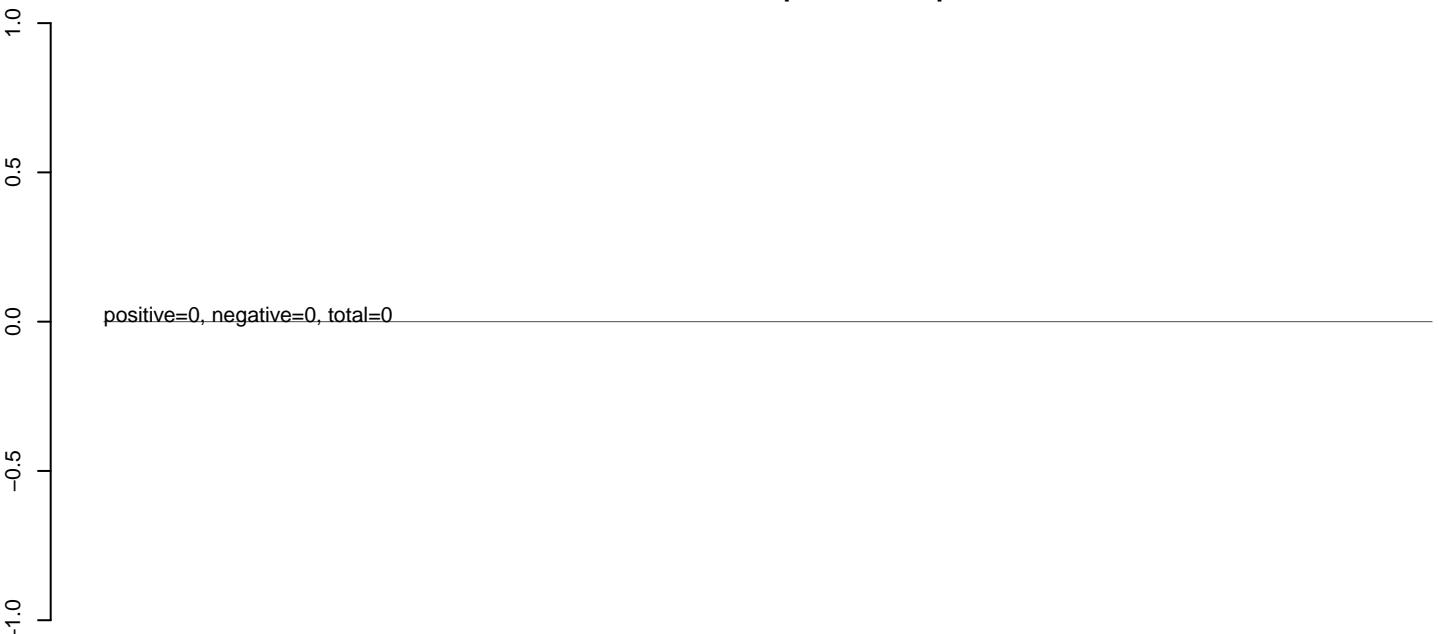
CuQuin_MG_7dpi3.rep



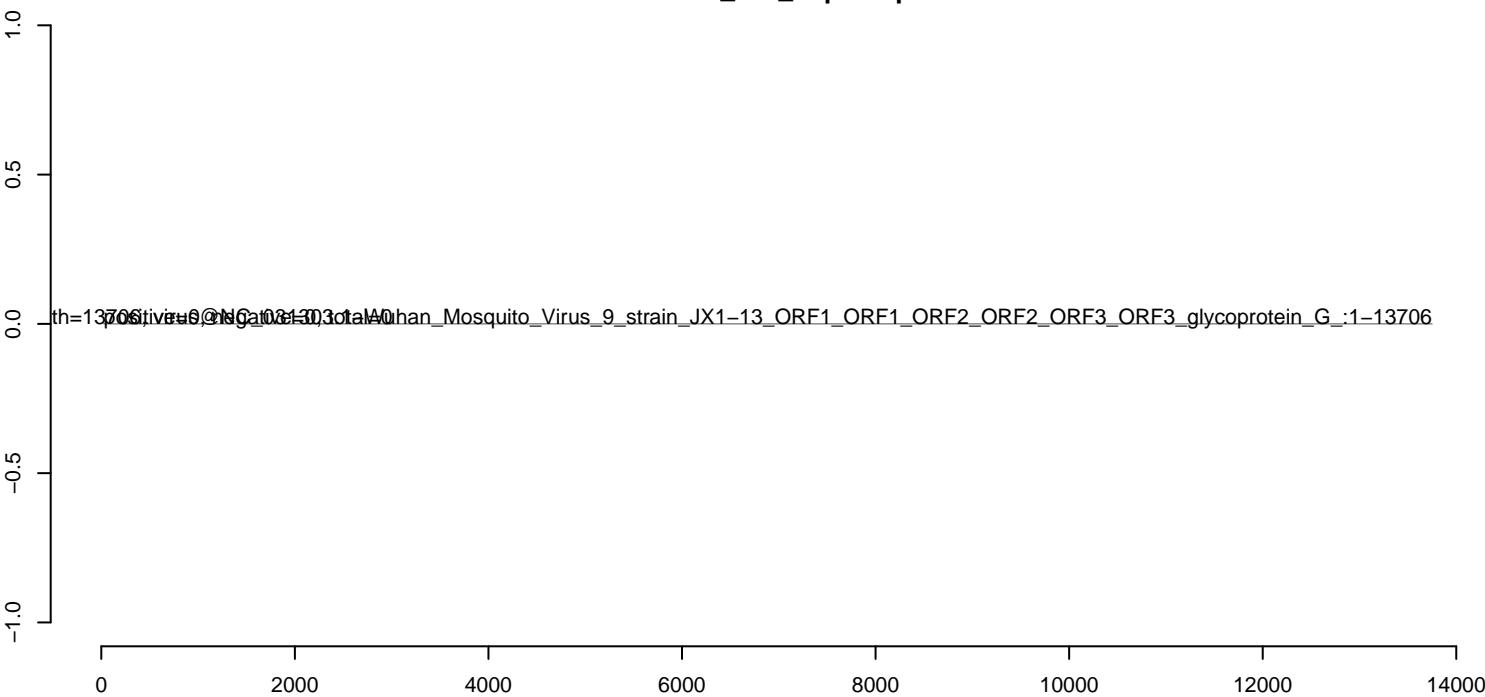
CuQuin_MG_7dpi3.18_23.rep



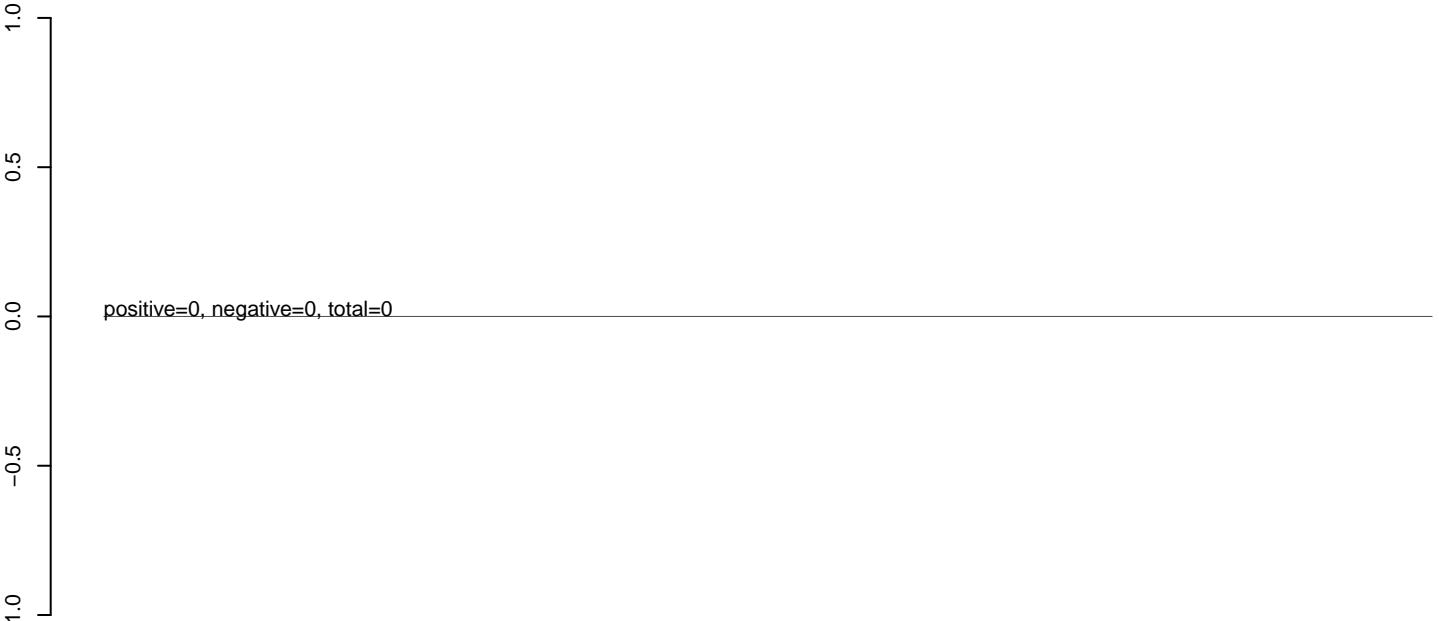
CuQuin_MG_7dpi3.24_35.rep



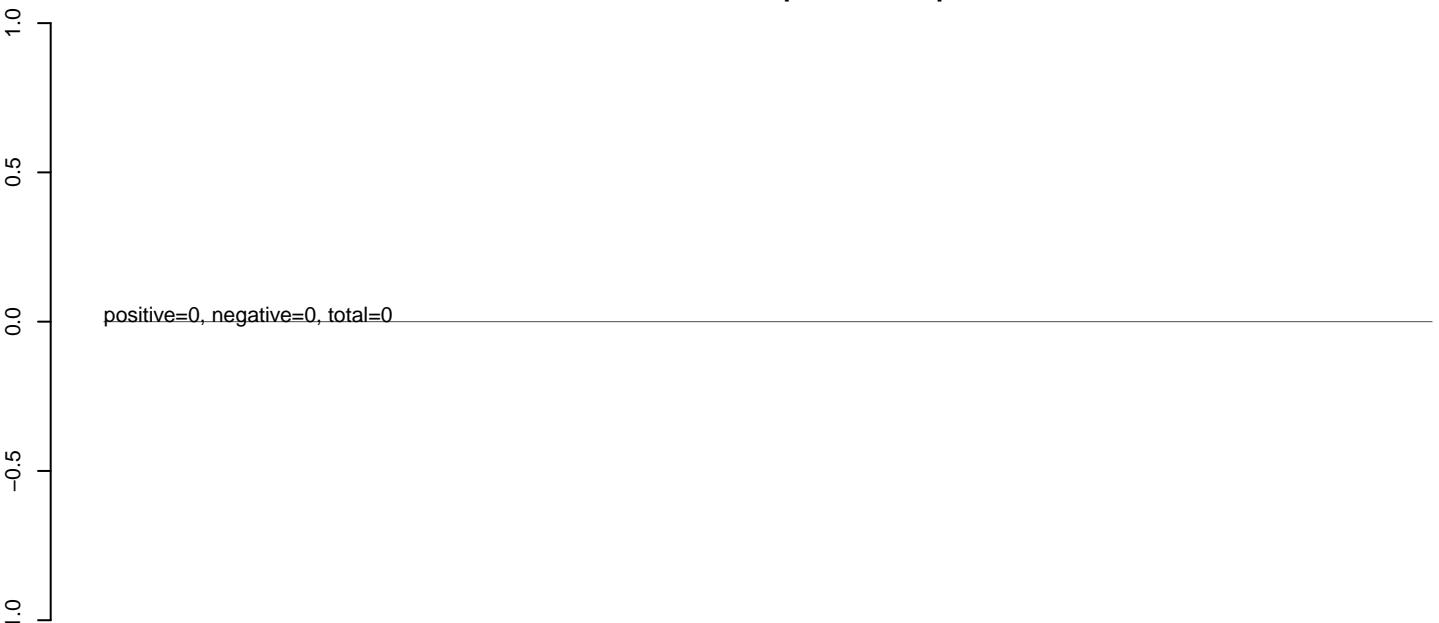
CuQuin_MG_7dpi3.rep



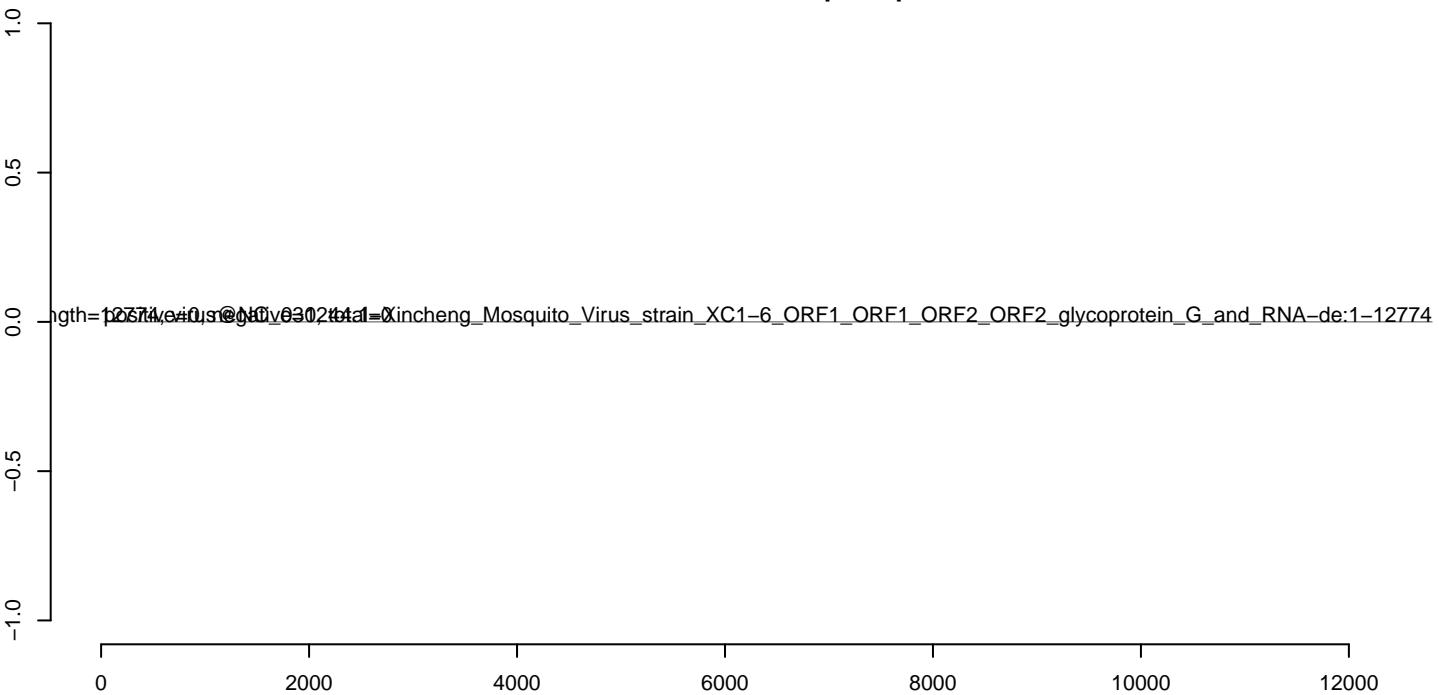
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=10158, virus@NC_030401.1-Hanko_virus_polyprotein_gene:1-10158

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

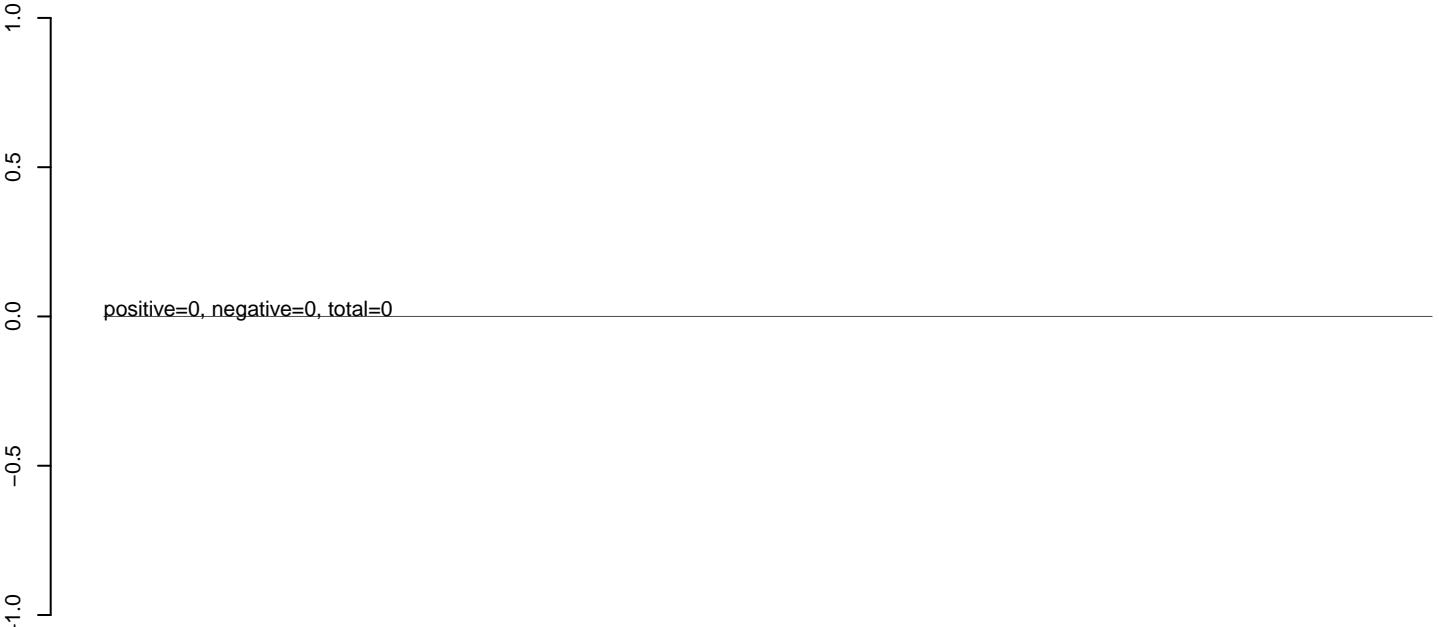
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

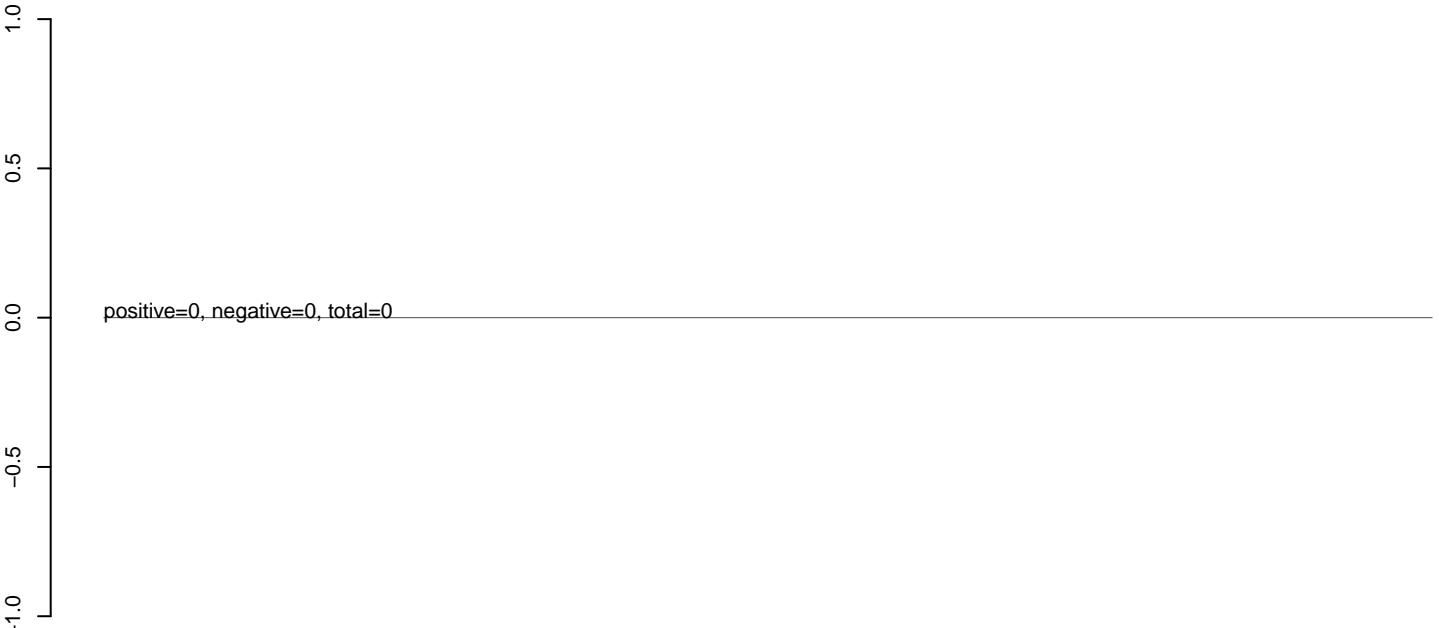
Window size=25, length=10864, virus@NC_030290.1-Culiseta_flavivirus_strain_502-13:1-10864

0 2000 4000 6000 8000 10000

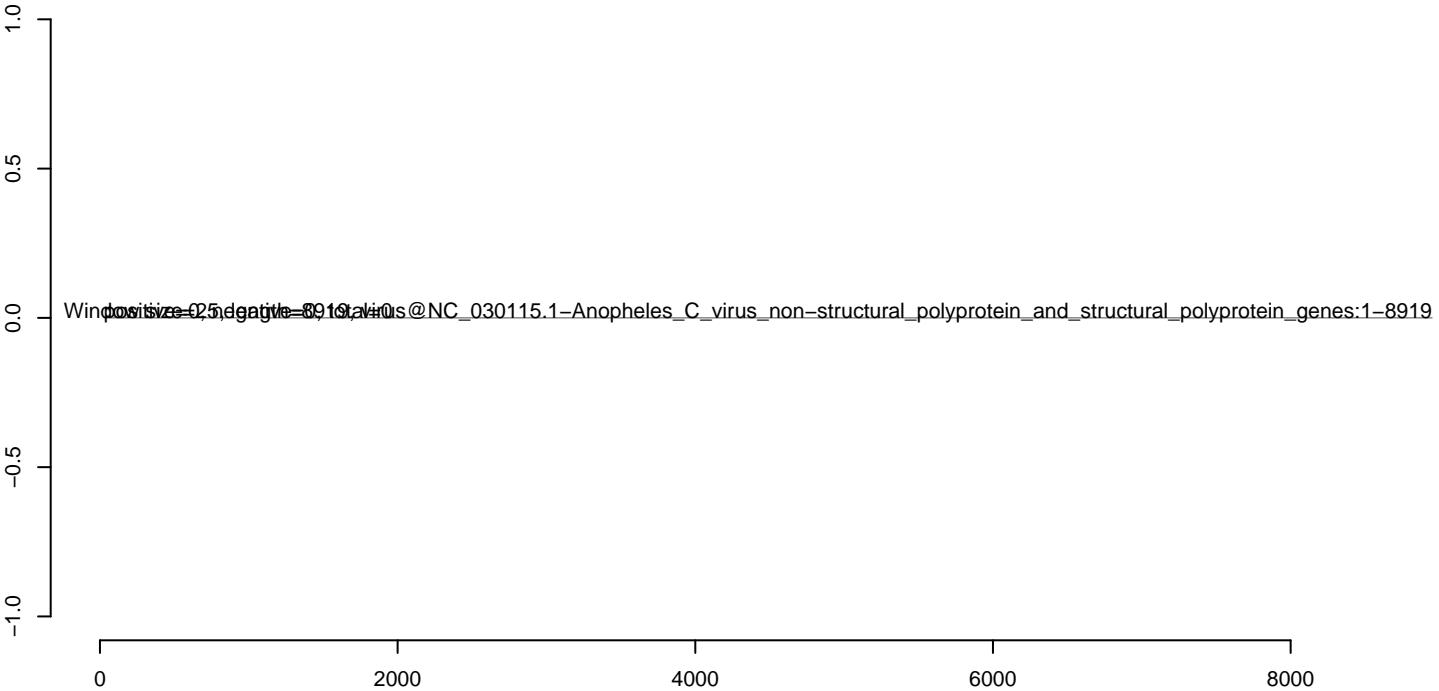
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

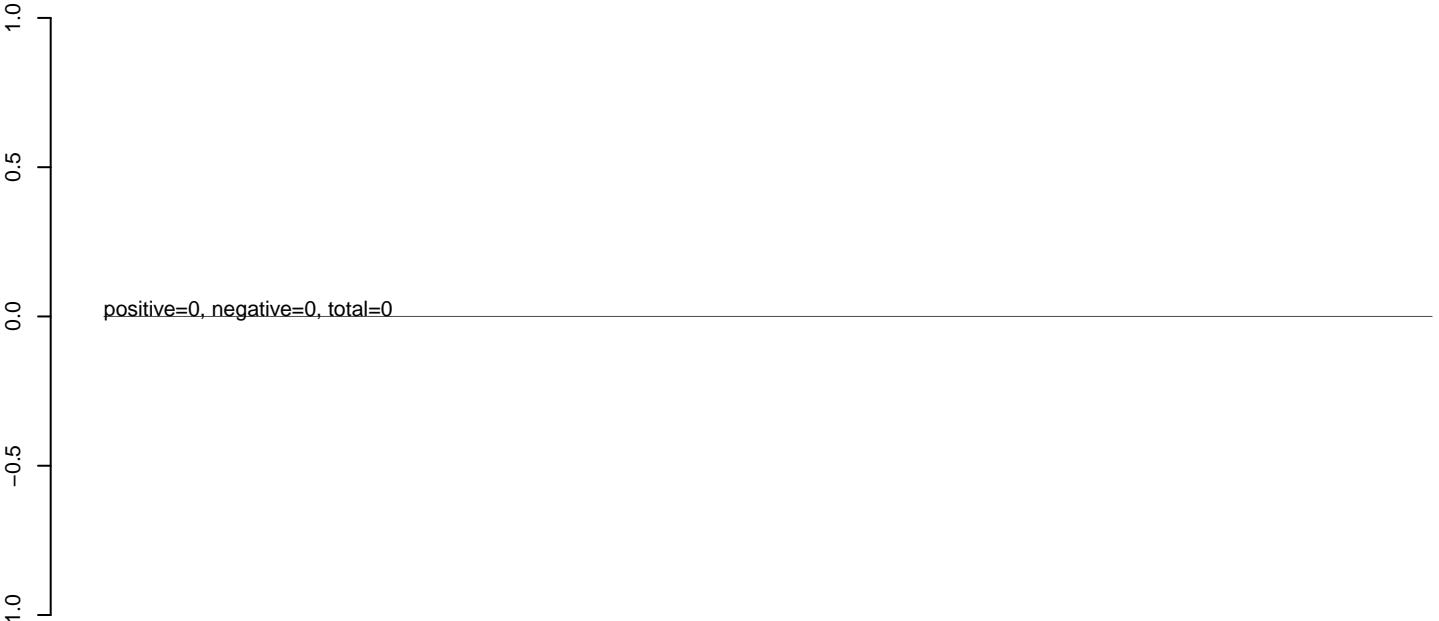
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

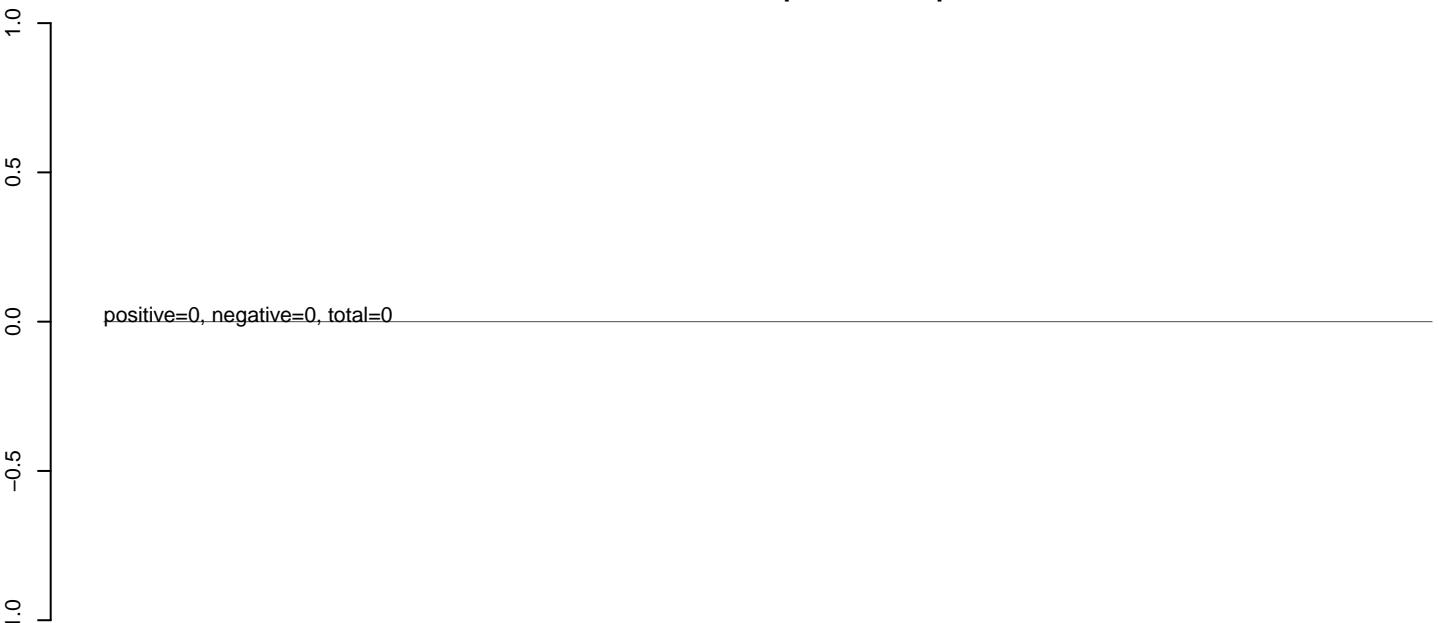
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window.size=25, length=12004, virus@NC_028265.1_Wuhan_Mosquito_Virus_8_strain_XC2-7:1-12004

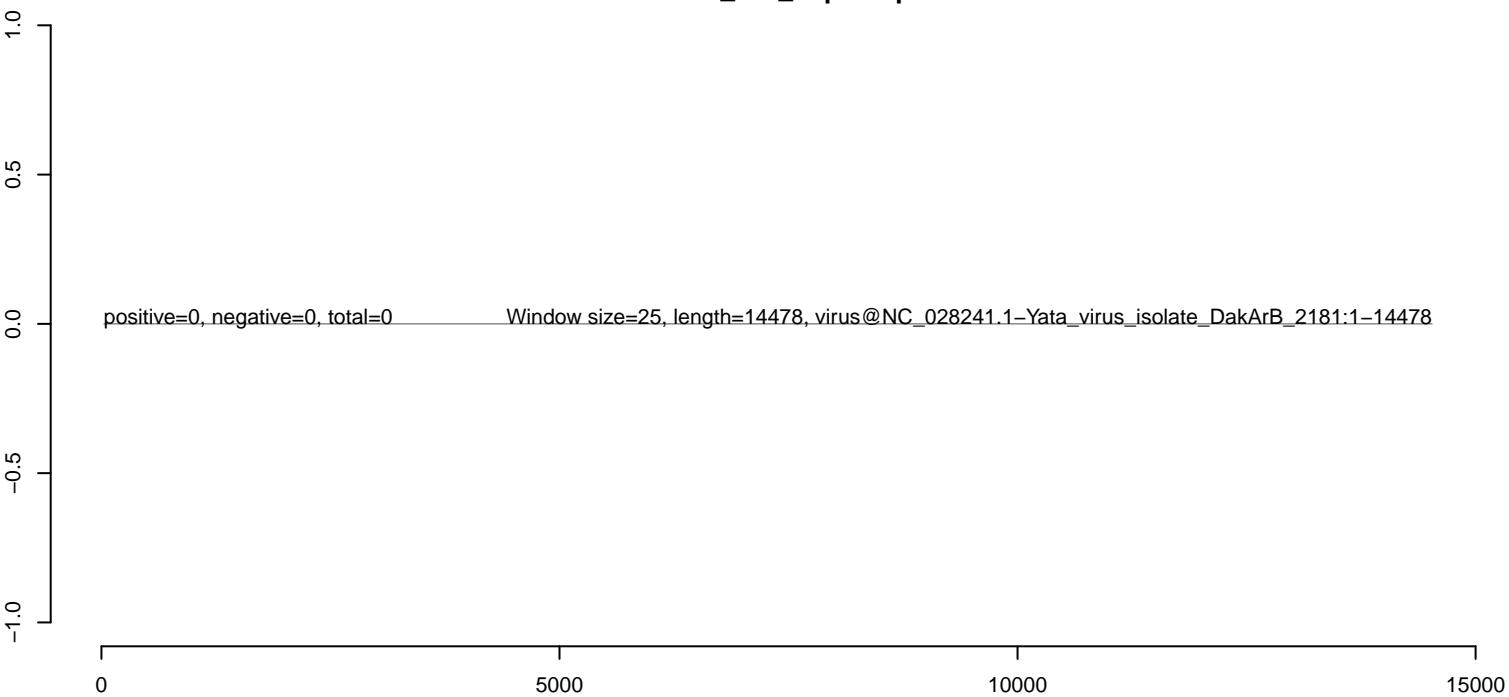
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window.size=25, length=10938, virus@NC_027819.1-Mercadeo_virus_isolate_ER-M10:1-10938

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

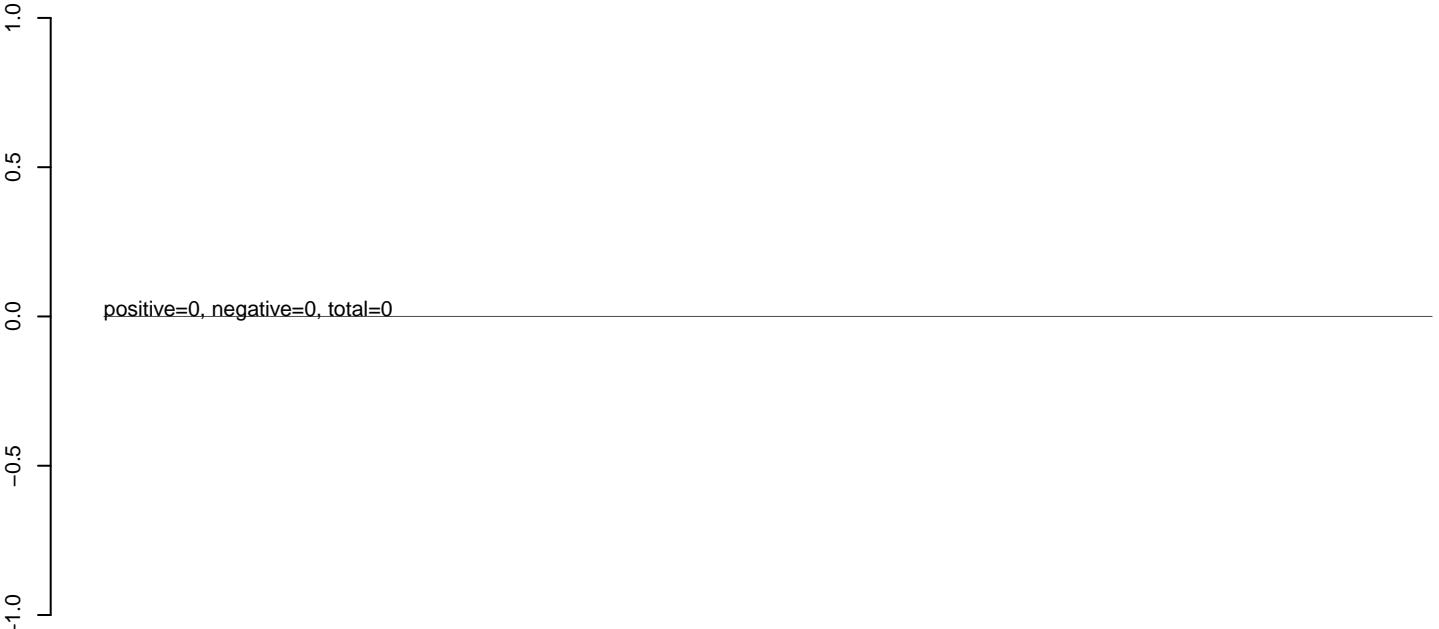
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

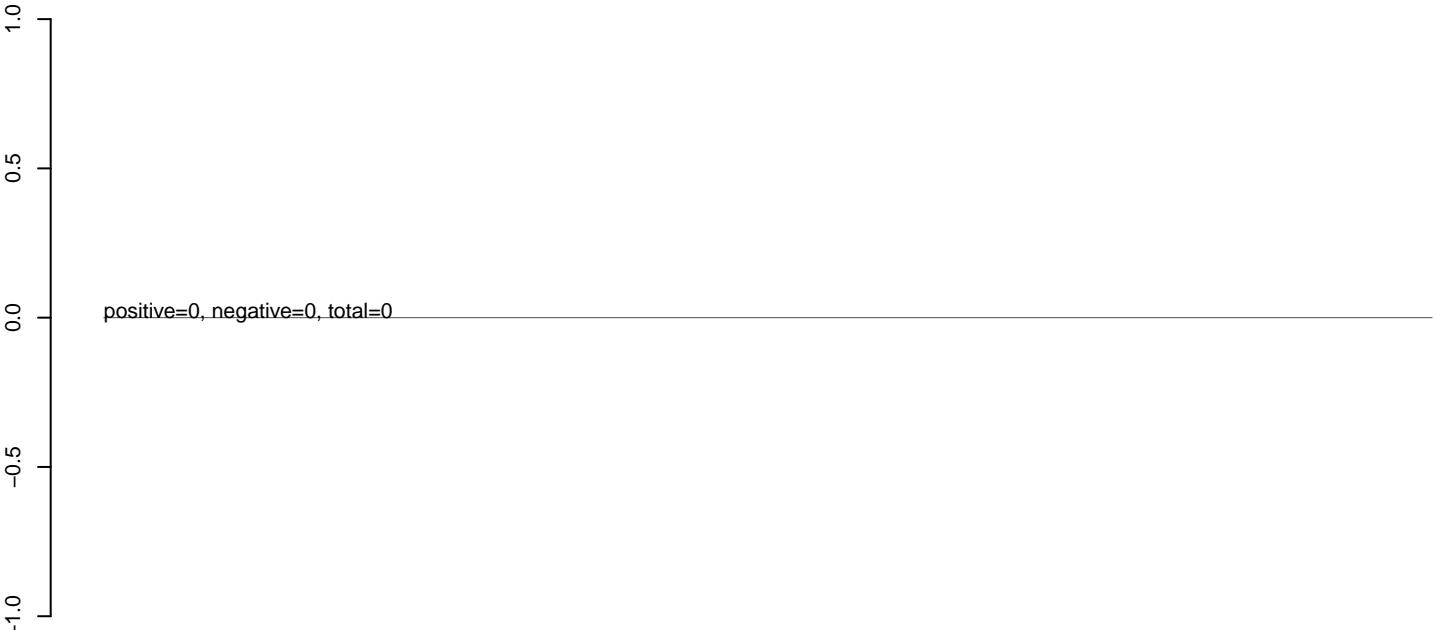
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0|Window size=25, length=10893, virus@NC_027817.1-Parramatta_River_virus_isolate_92-B115745:1-10893

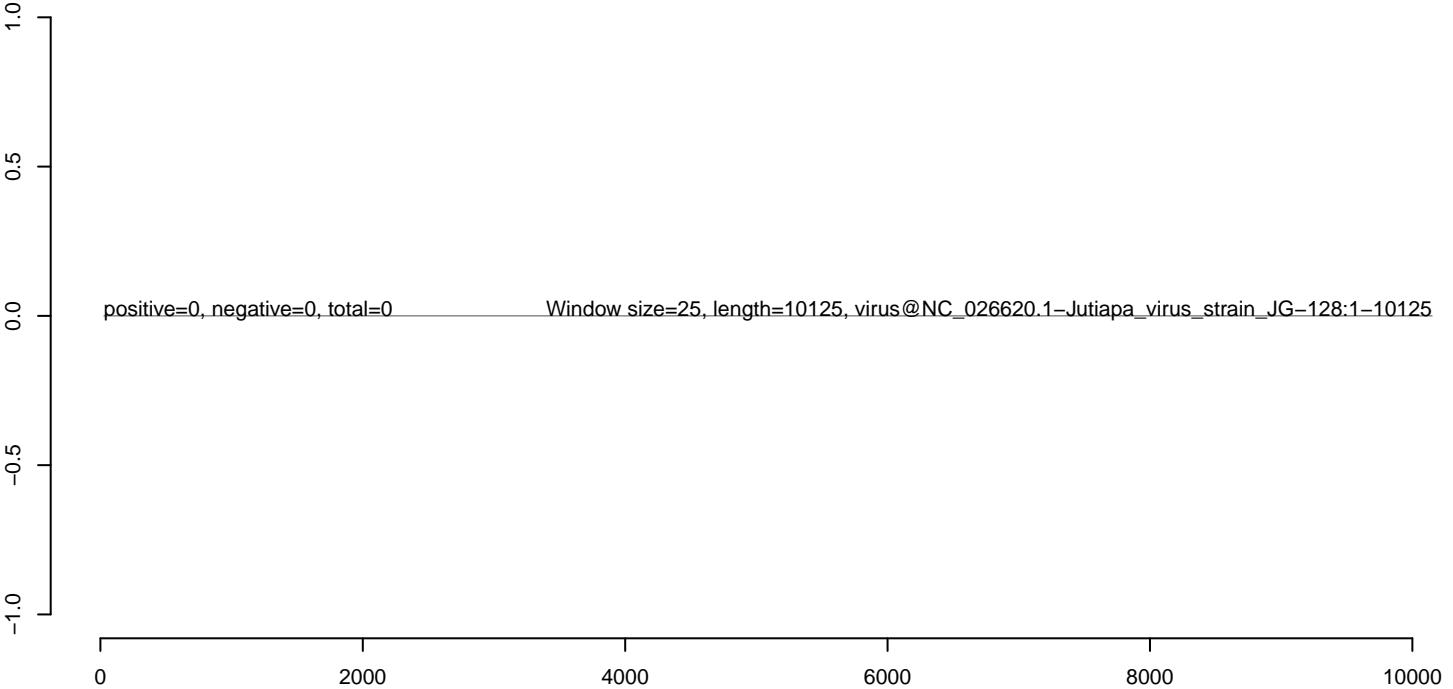
CuQuin_MG_7dpi3.18_23.rep



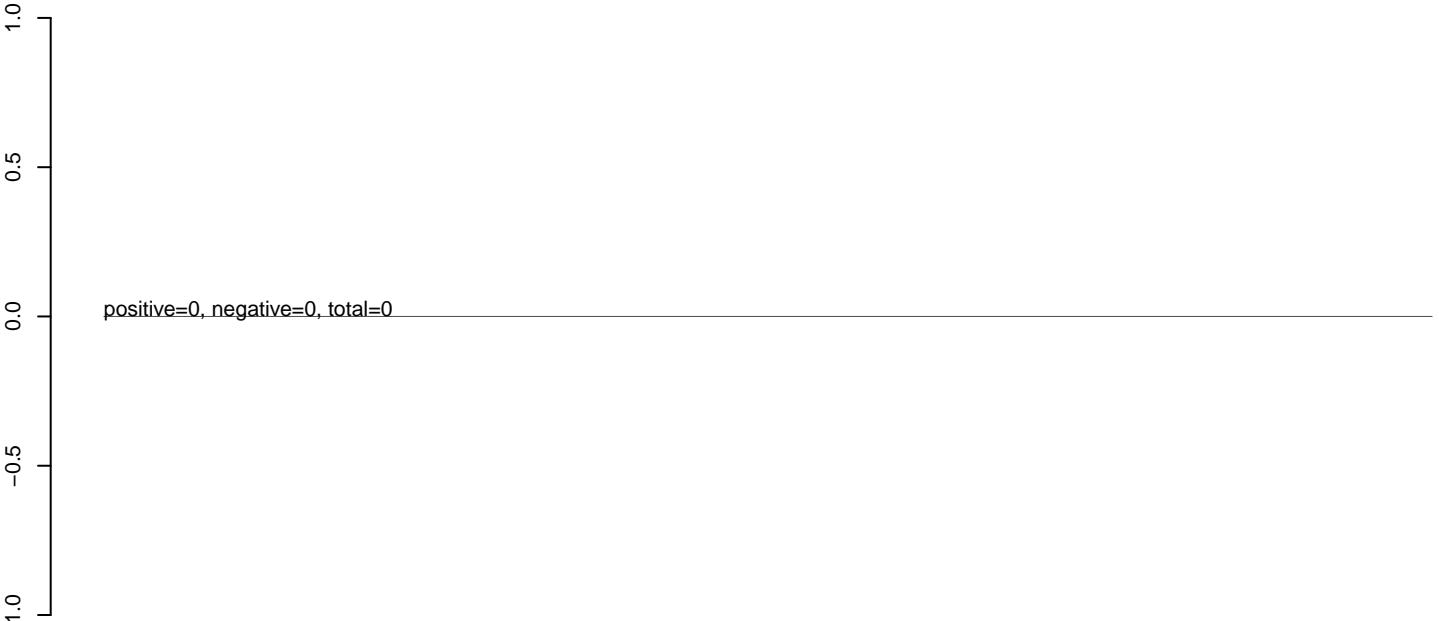
CuQuin_MG_7dpi3.24_35.rep



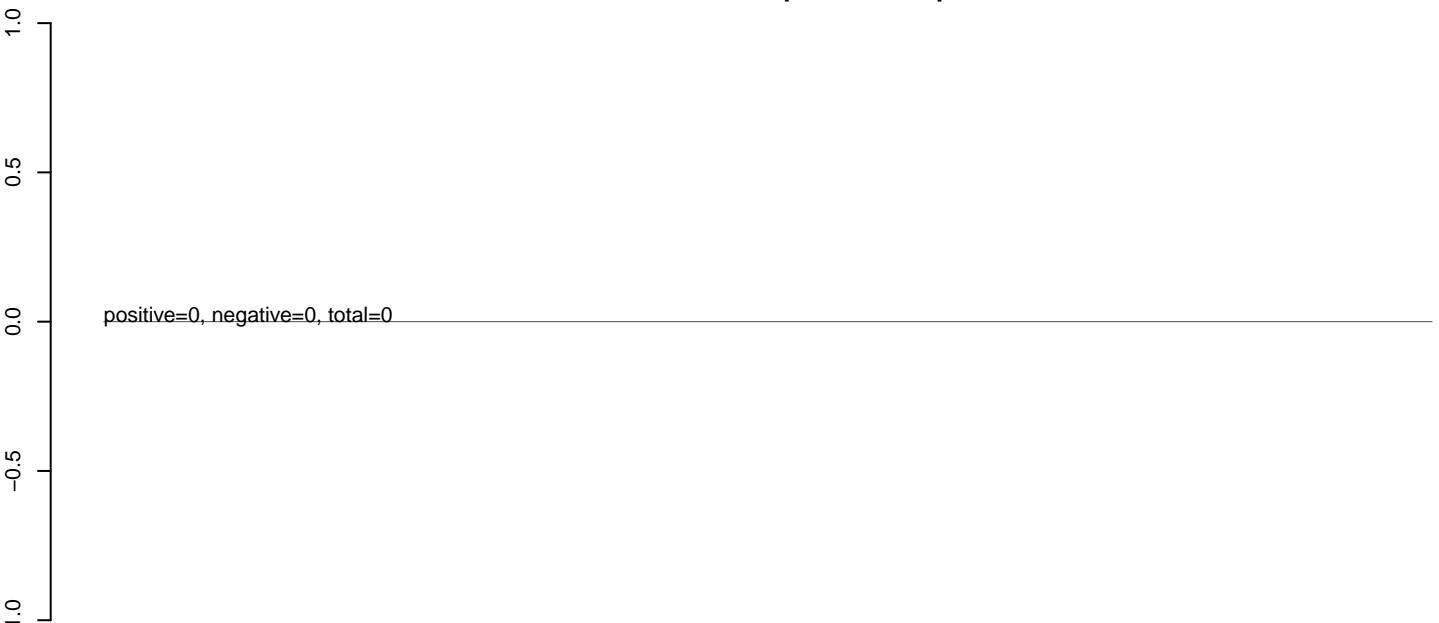
CuQuin_MG_7dpi3.rep



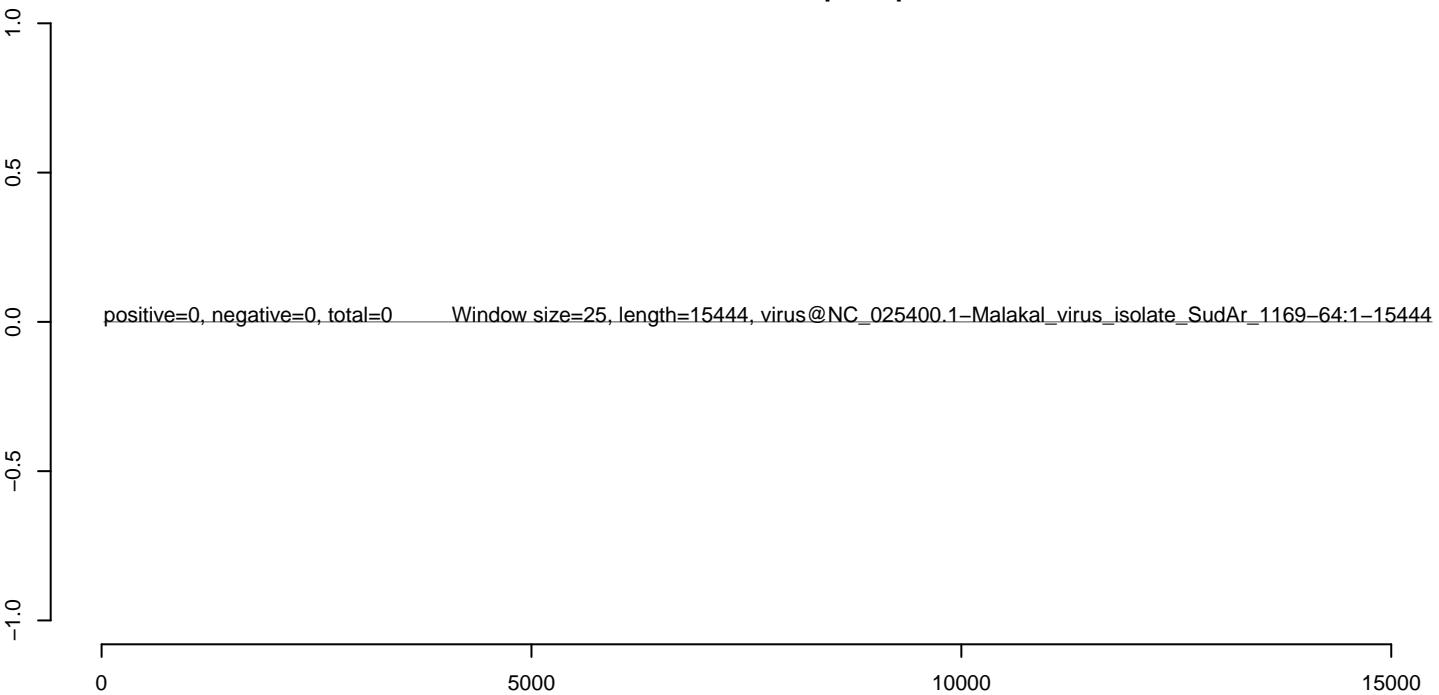
CuQuin_MG_7dpi3.18_23.rep



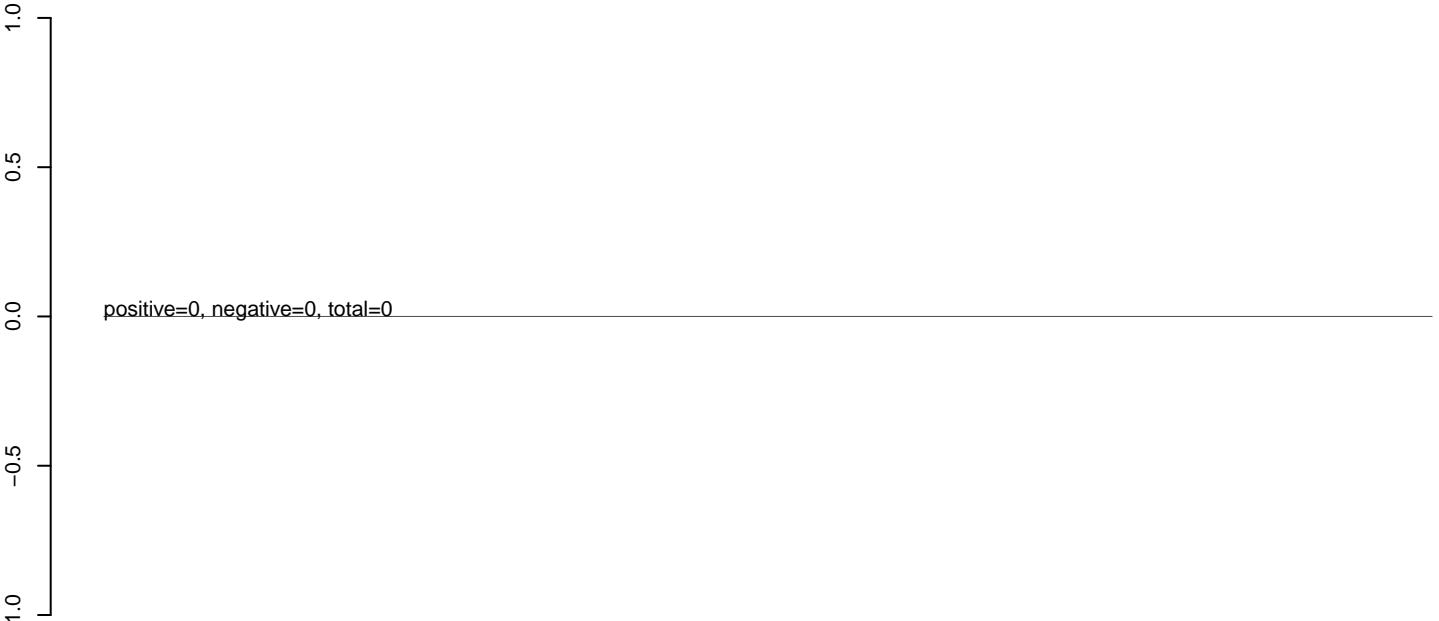
CuQuin_MG_7dpi3.24_35.rep



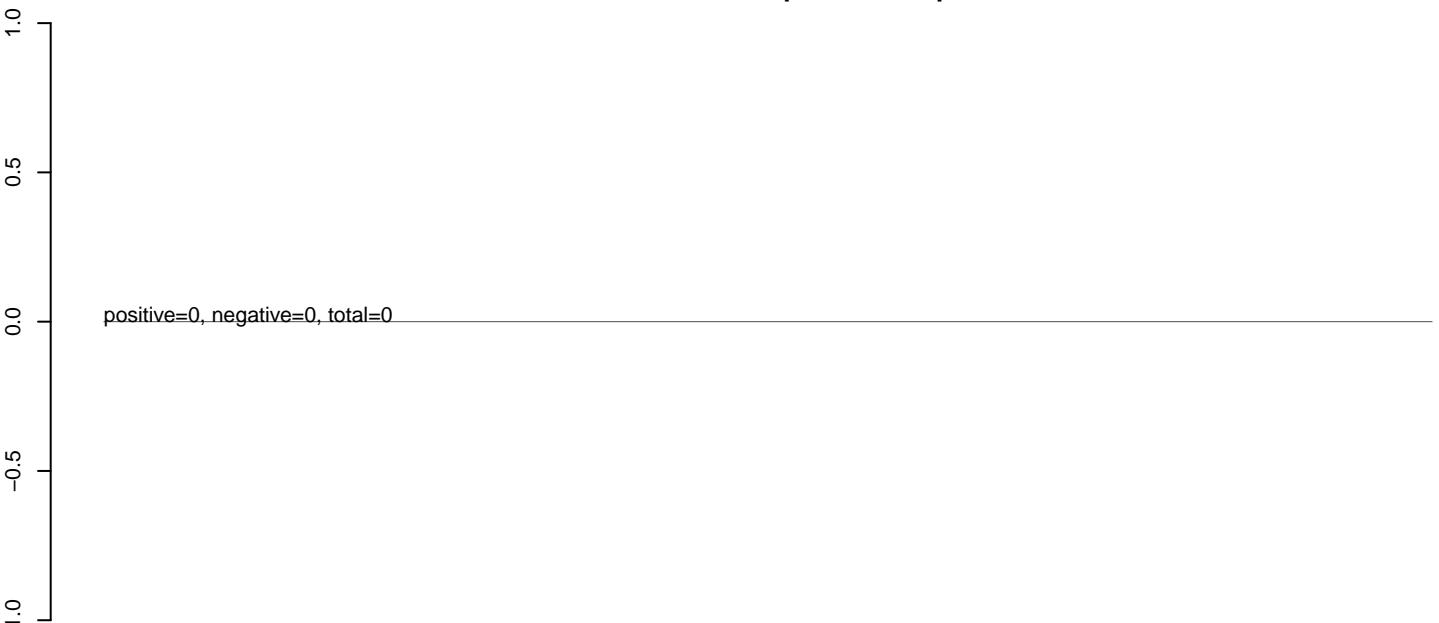
CuQuin_MG_7dpi3.rep



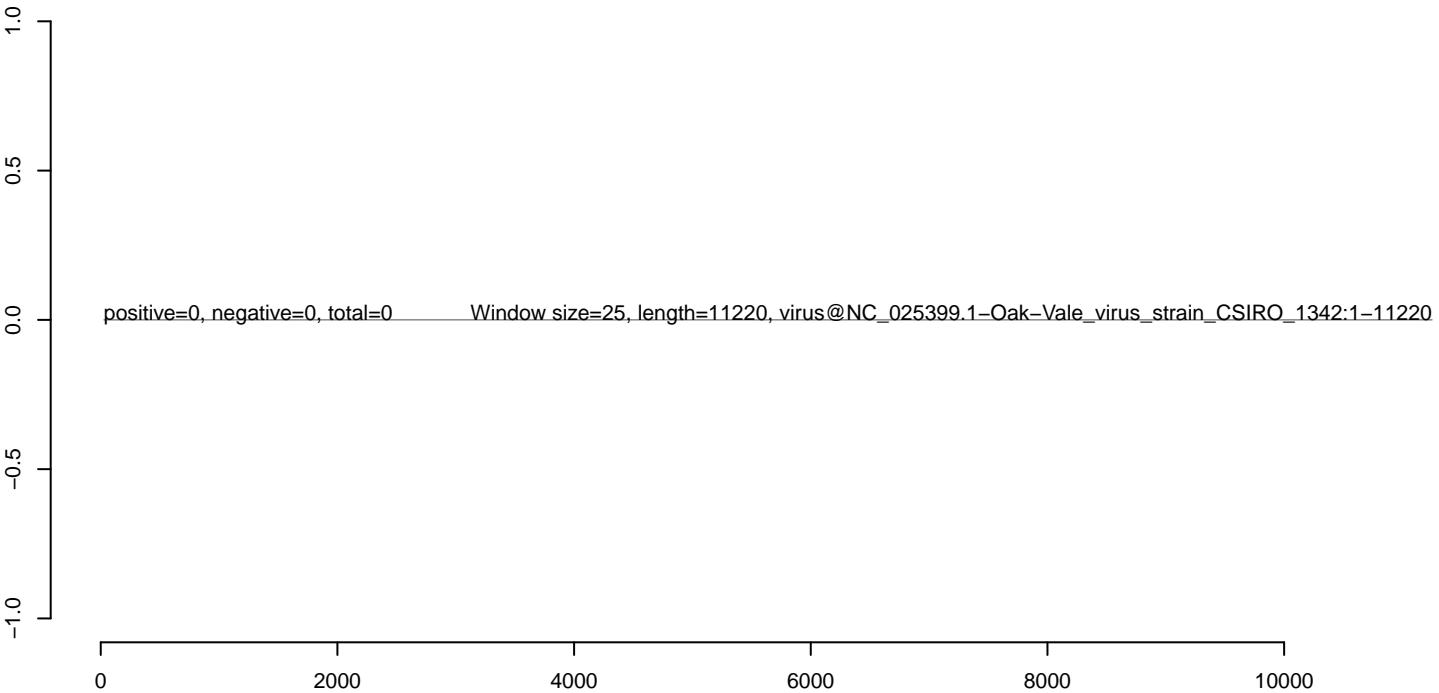
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=11876, virus@NC_025395.1–Puerto_Almendras_virus_isolate_LO-39:1–11876

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 size=25, length=11190, virus@NC_025384.1-Culex_tritaeniorhynchus_rhabdovirus_RNA_strain_TY:1-11190

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

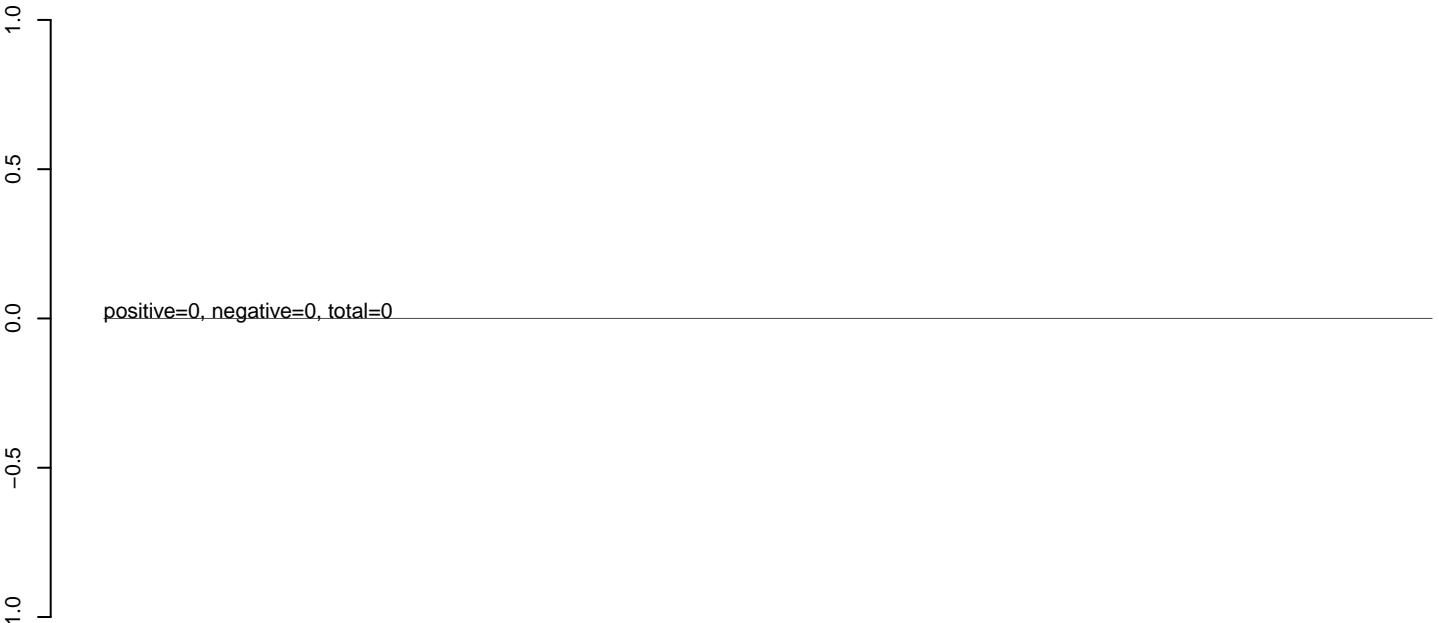
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

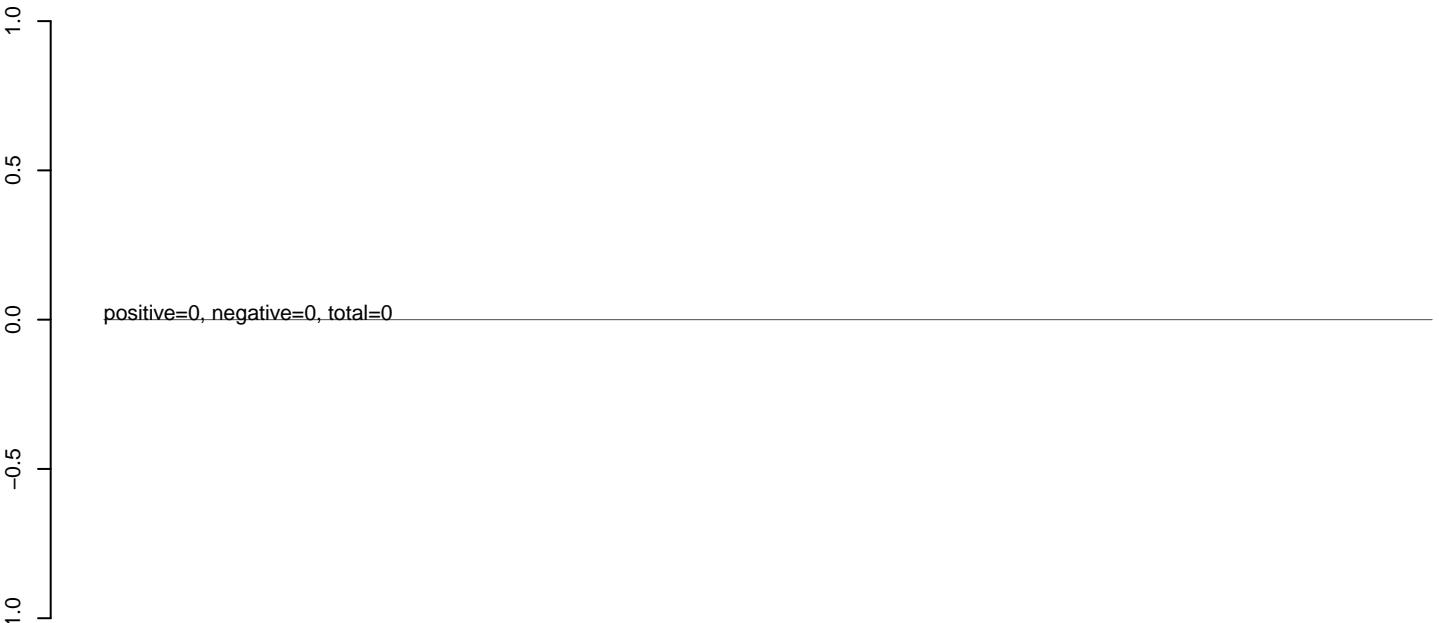
Window size=25, length=11096, virus@NC_025376.1–Grass_carp_rhabdovirus_V76:1–11096

0 2000 4000 6000 8000 10000

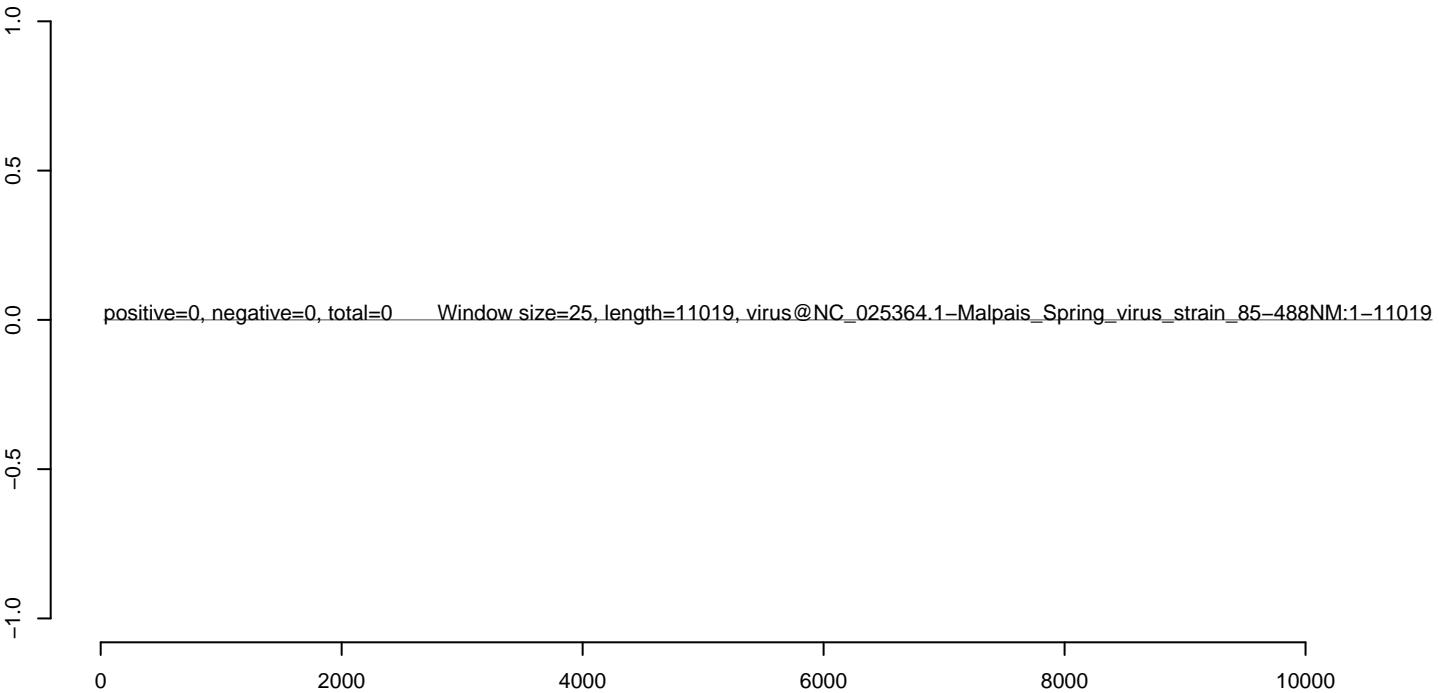
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=12240, virus@NC_025362.1-Xiburema_virus_isolate_XIBV_BE_AR_362159:1-12240

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=11526, virus@NC_025359.1-Moussa_virus_isolate_C23:1-11526

0 2000 4000 6000 8000 10000 12000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=11097, virus@NC_025356.1–Pike_fry_rhabdovirus_isolate_F4:1–11097

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=11550, virus@NC_024887.1-Middleburg_virus_isolate_ArB-8422:1-11550

0 2000 4000 6000 8000 10000 12000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=10320, virus@NC_024806.1-Lammi_virus_isolate_M0719_polyprotein_gene:1-10320

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

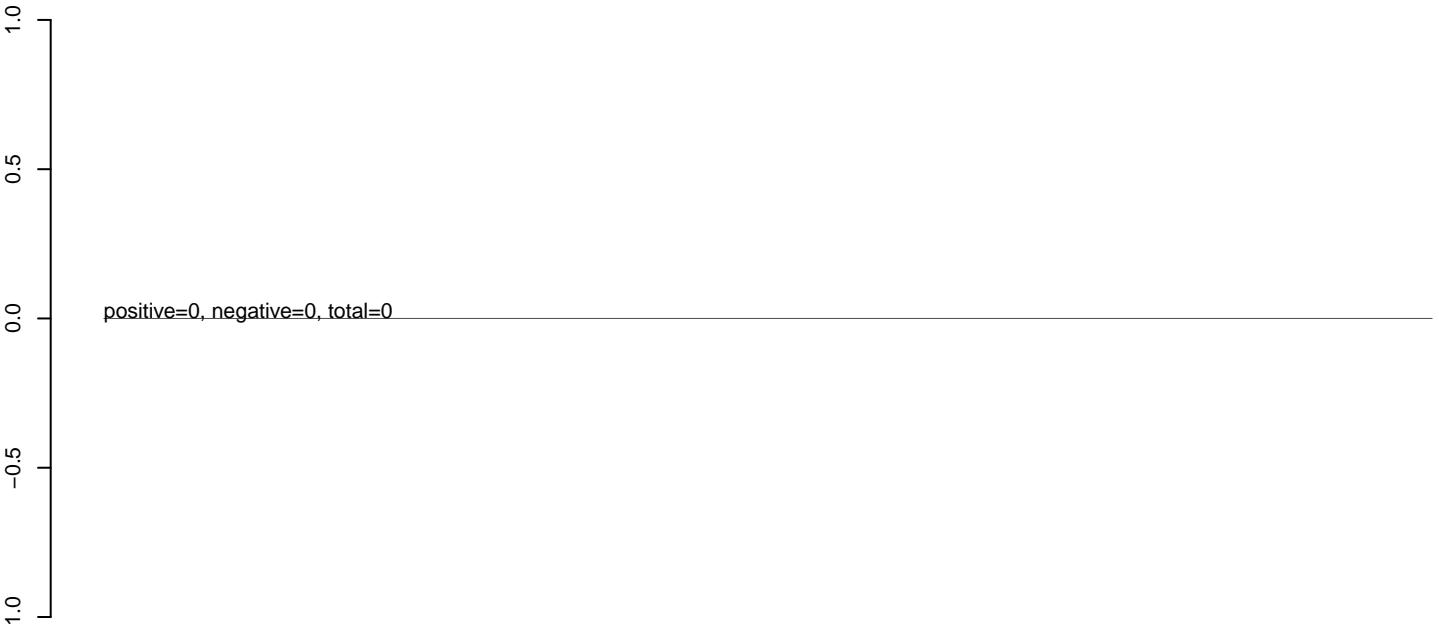
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

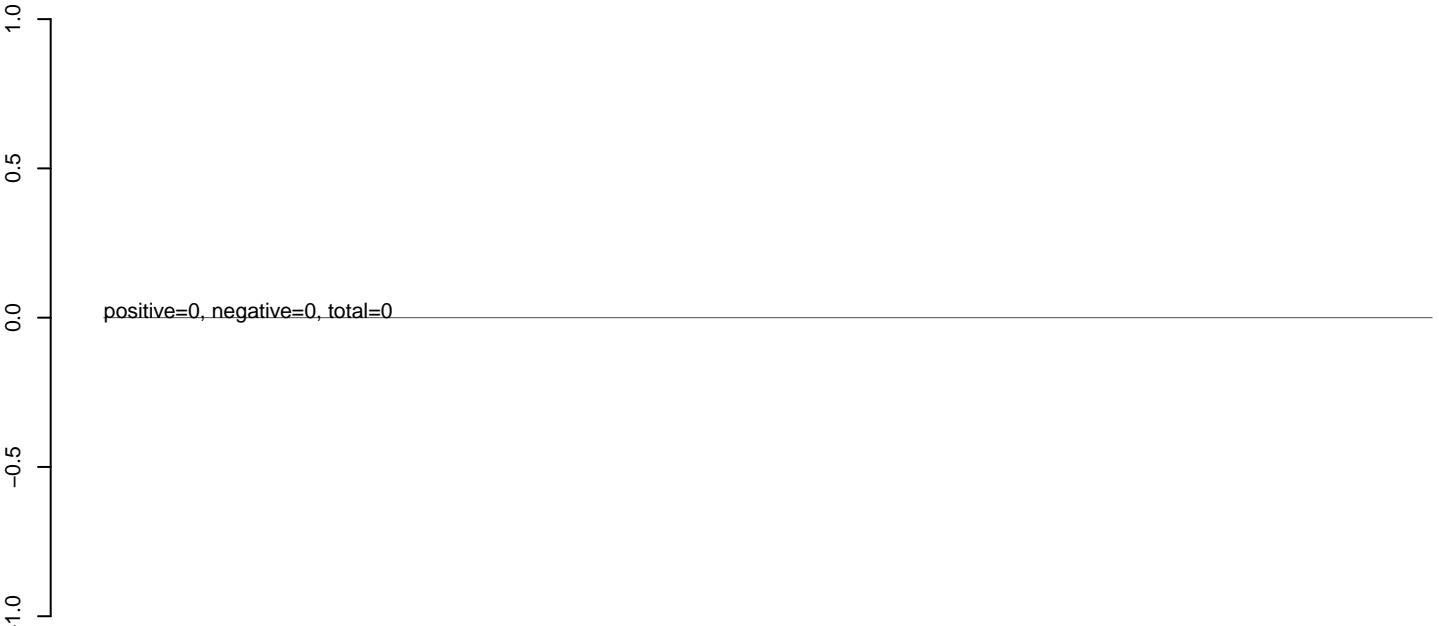
positive=0, negative=0, total=140, window size=25, length=10365, virus@NC_024805.1–llomantsi_virus_isolate_M0724_polyprotein_gene:1–10365

0 2000 4000 6000 8000 10000

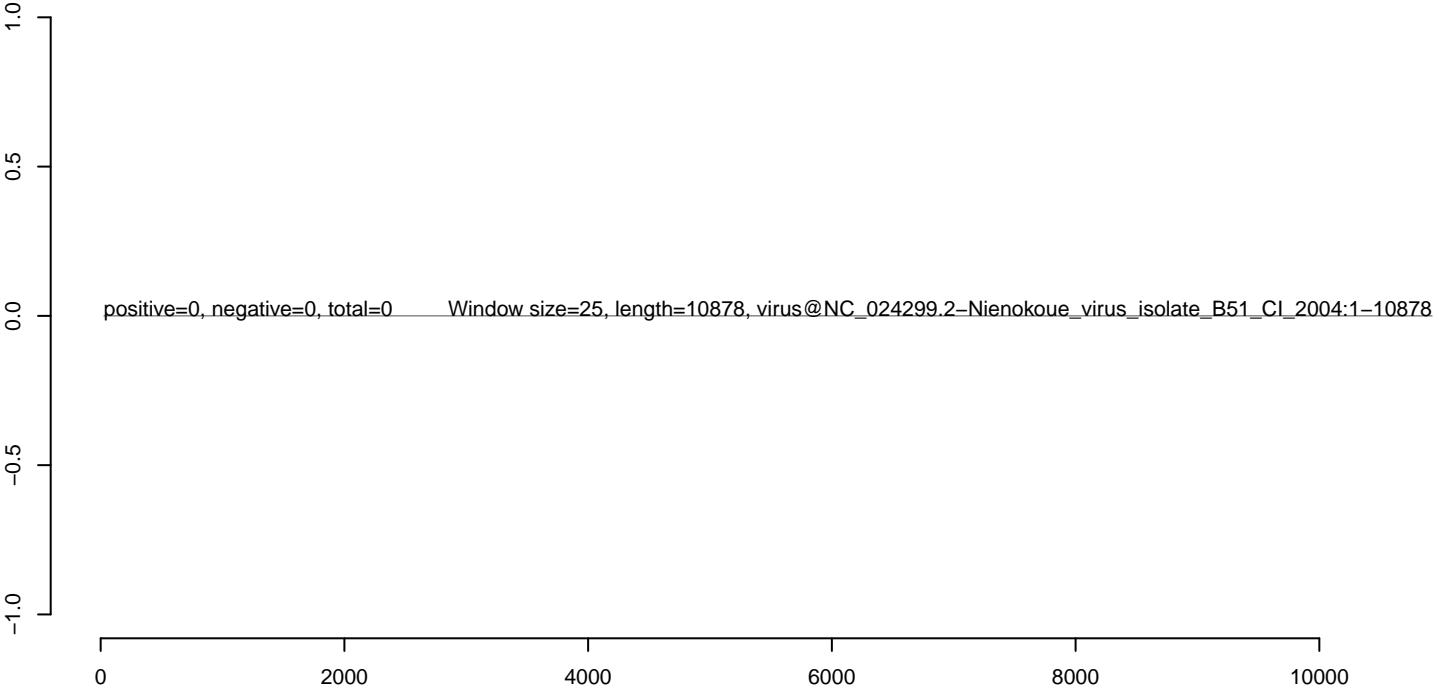
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0, window size=25, length=10673, virus@NC_023424.1-Tyulениевирус_стрain LEIV-6C_polyprotein_gene:1-10673

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

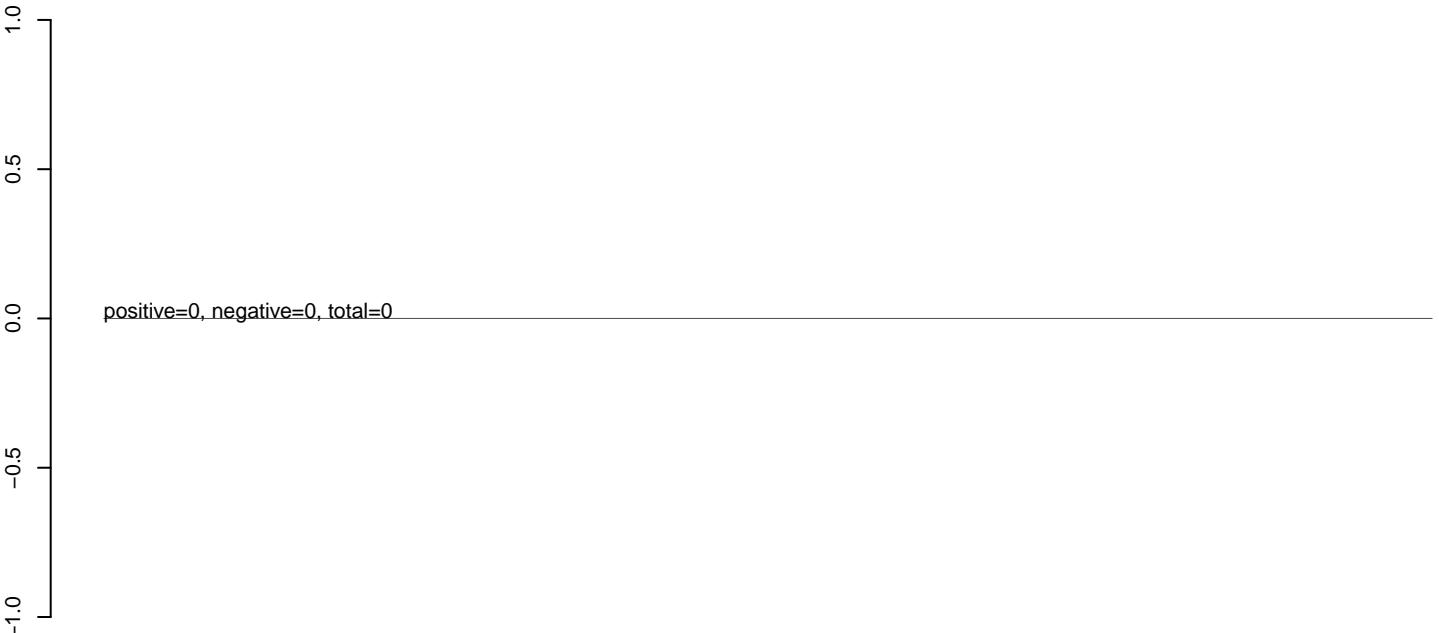
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

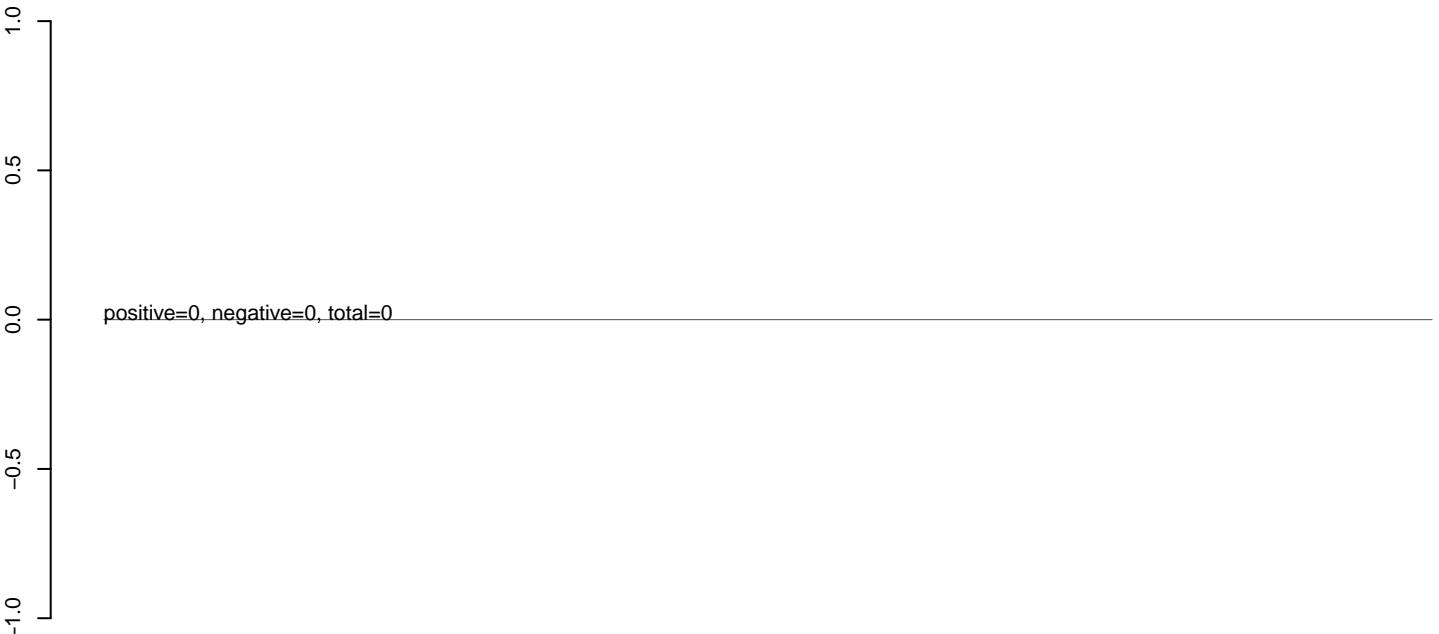
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=10865, virus@NC_021069.1-Mosquito_flavivirus_isolate_LSFaViV-A20-09:1-10865

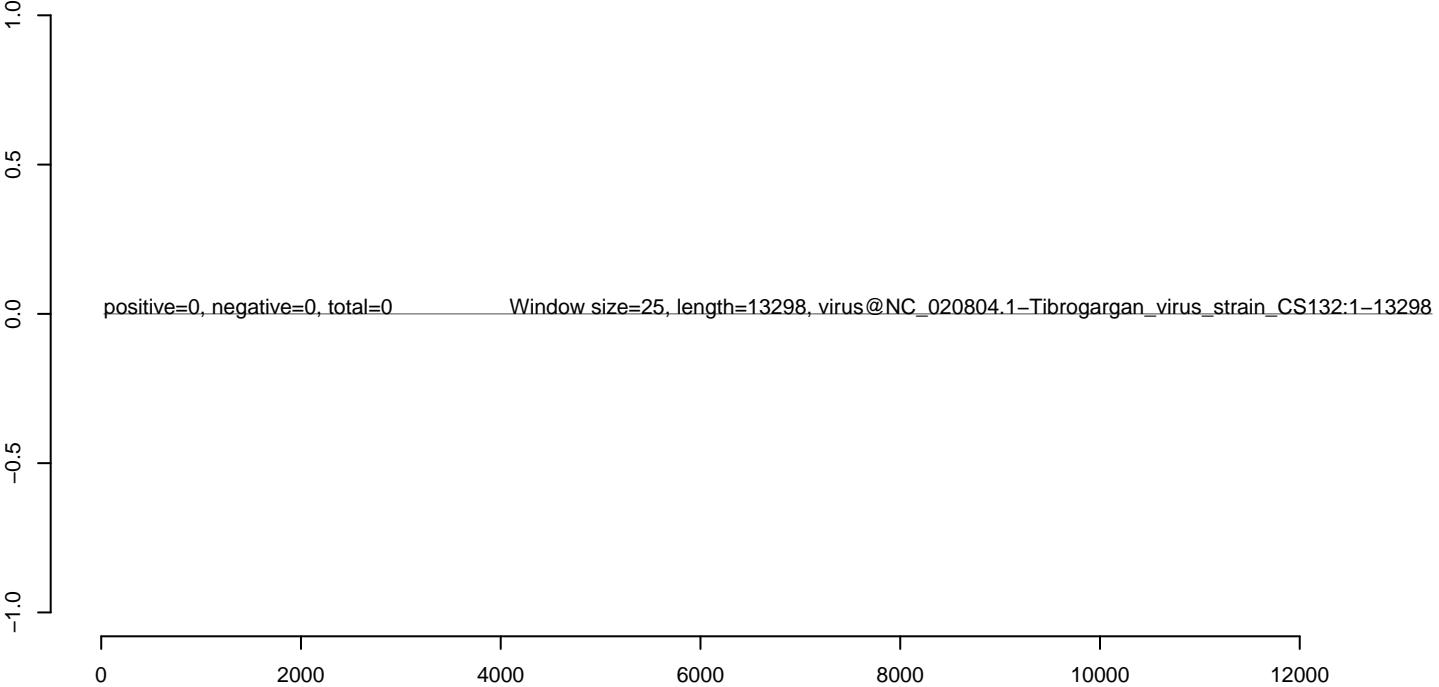
CuQuin_MG_7dpi3.18_23.rep



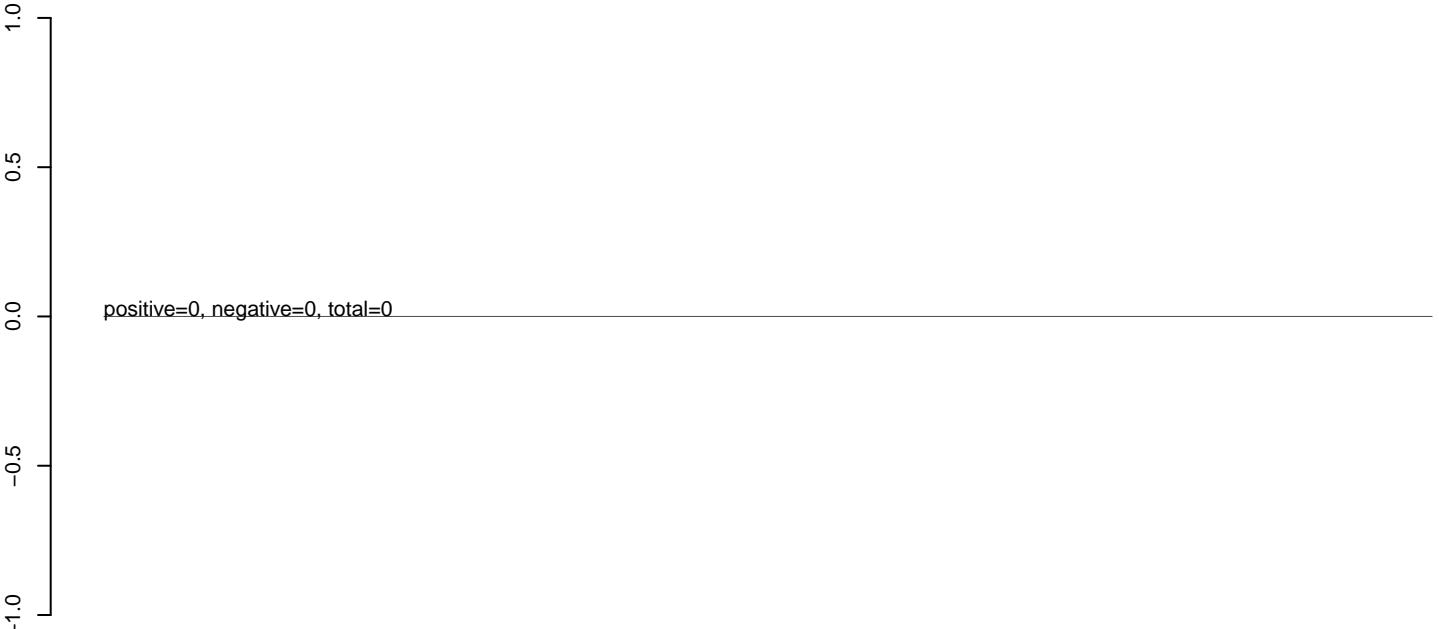
CuQuin_MG_7dpi3.24_35.rep



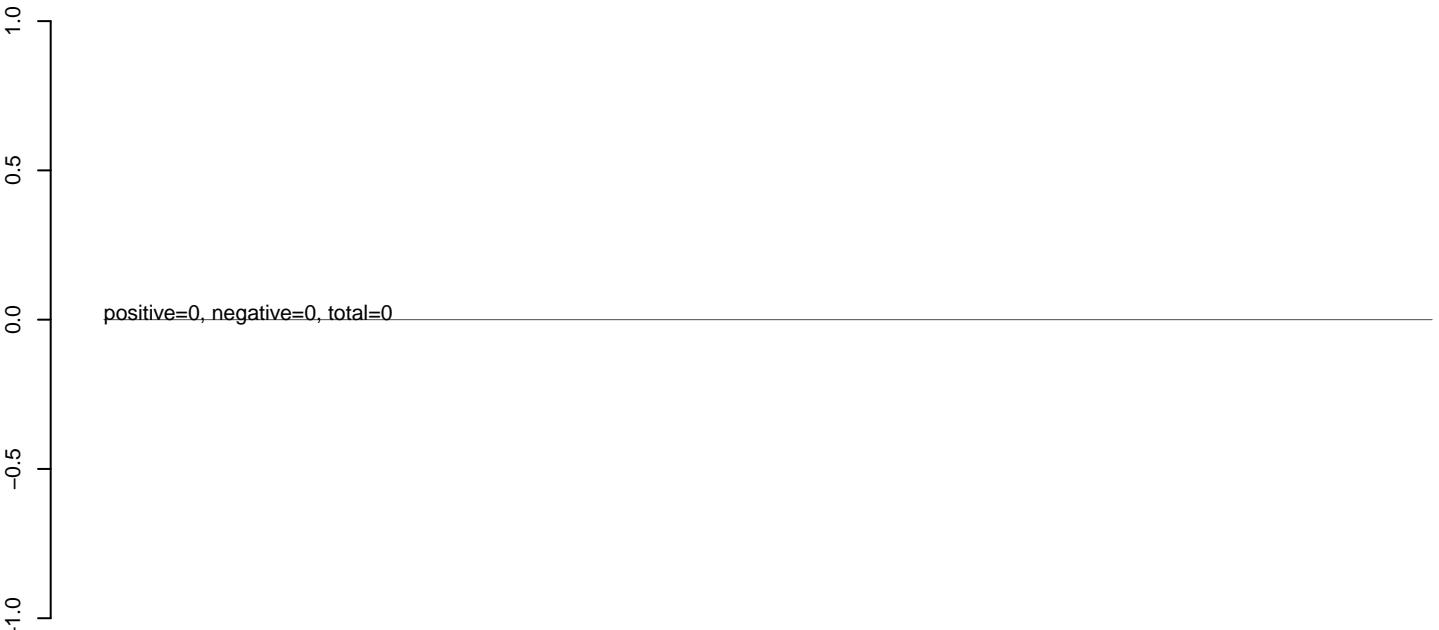
CuQuin_MG_7dpi3.rep



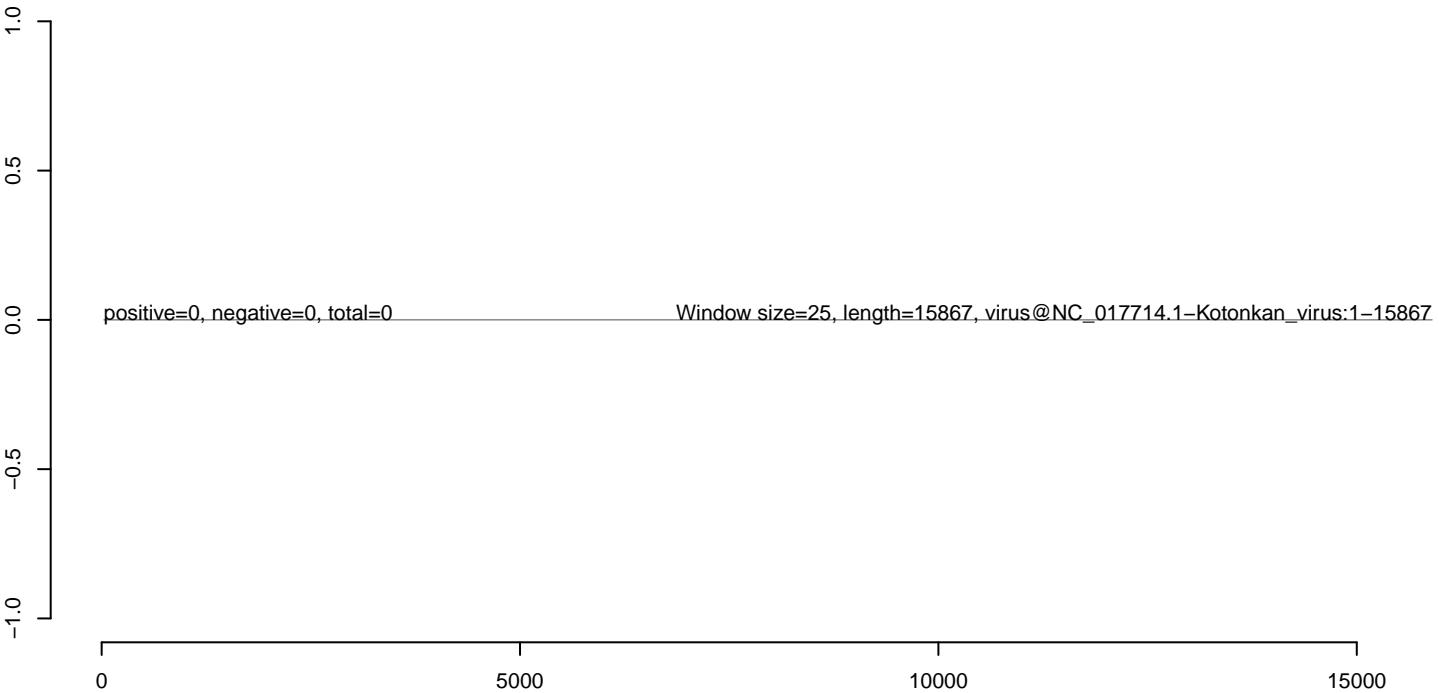
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

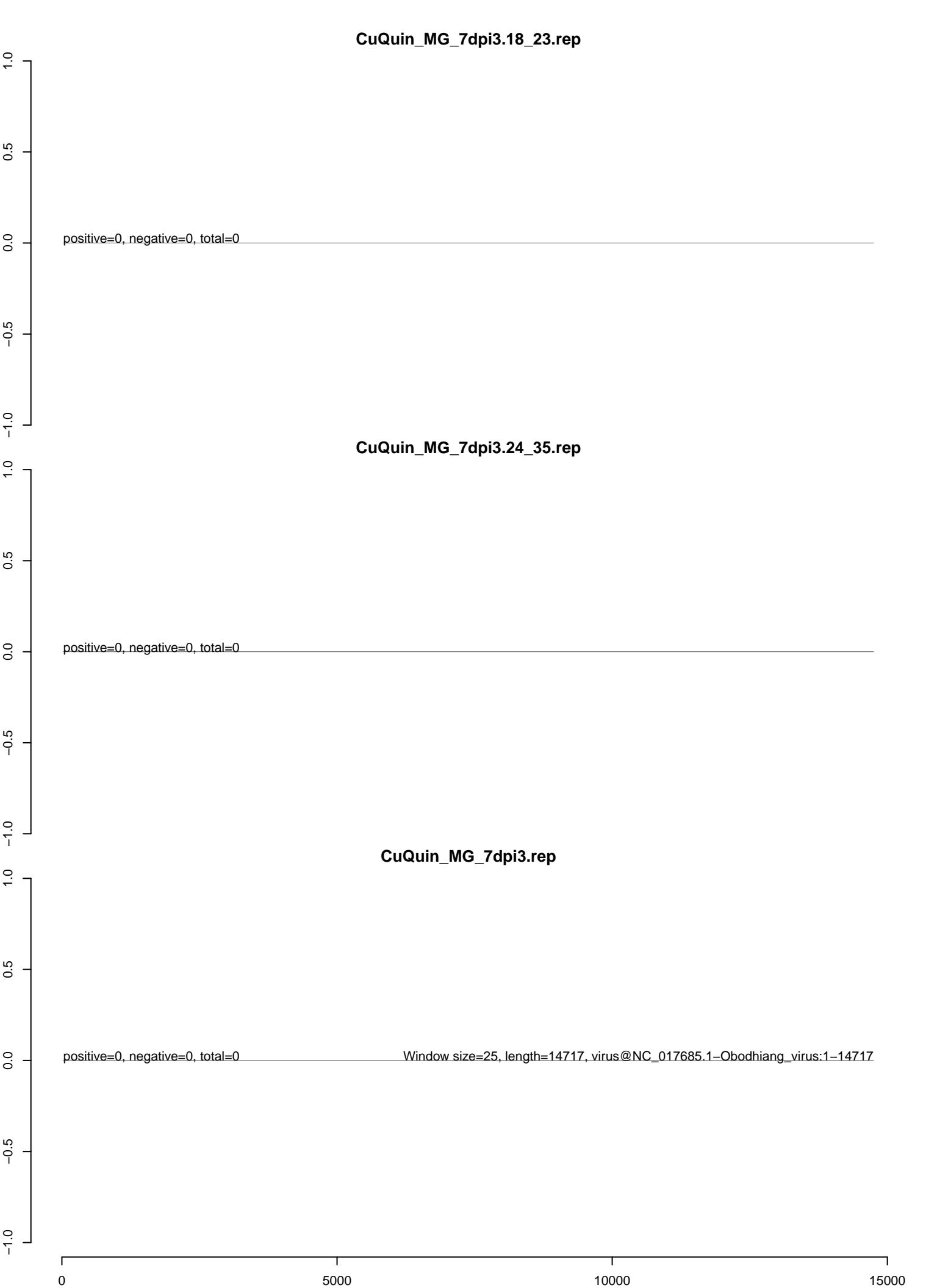
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

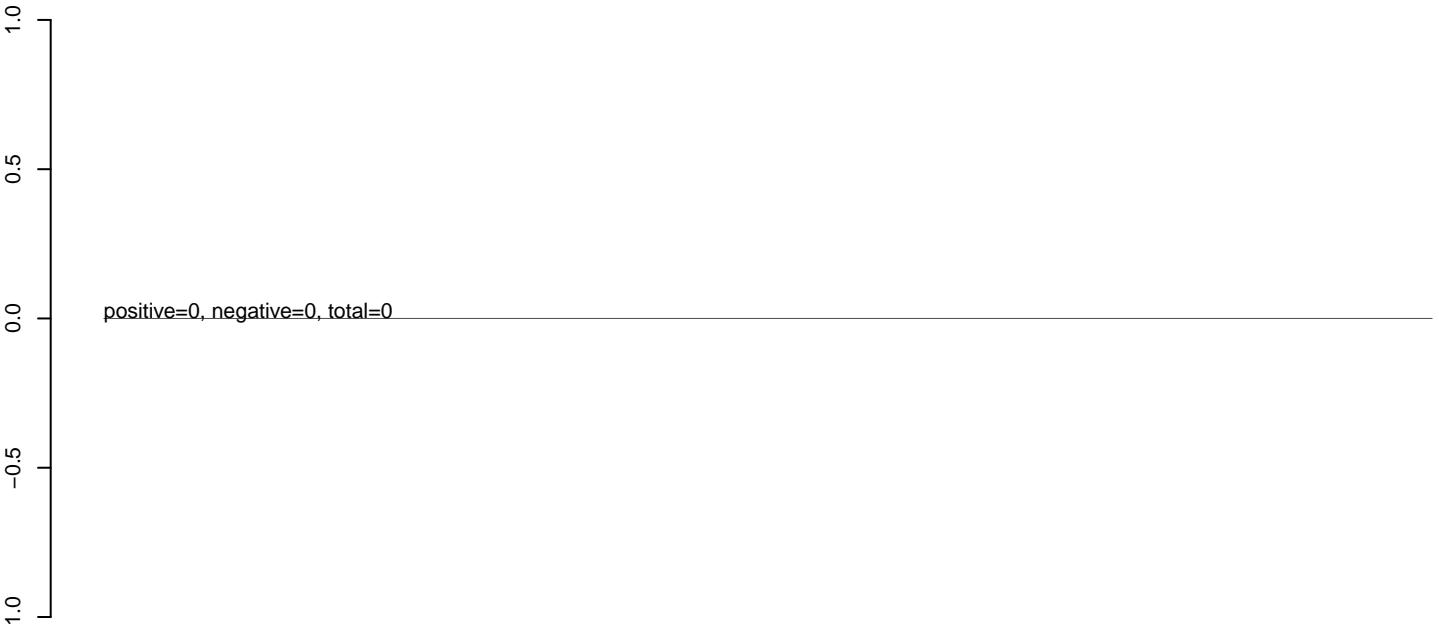
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

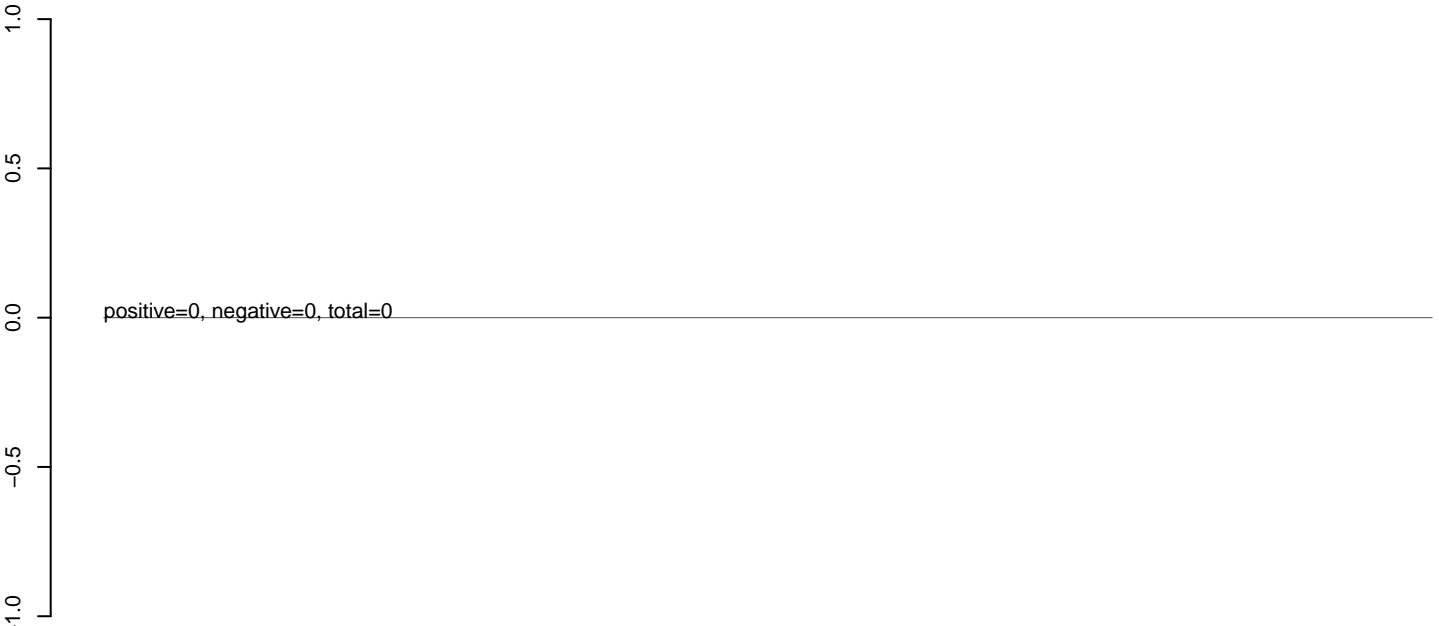
Window size=25, length=14717, virus@NC_017685.1–Obodhiang_virus:1–14717



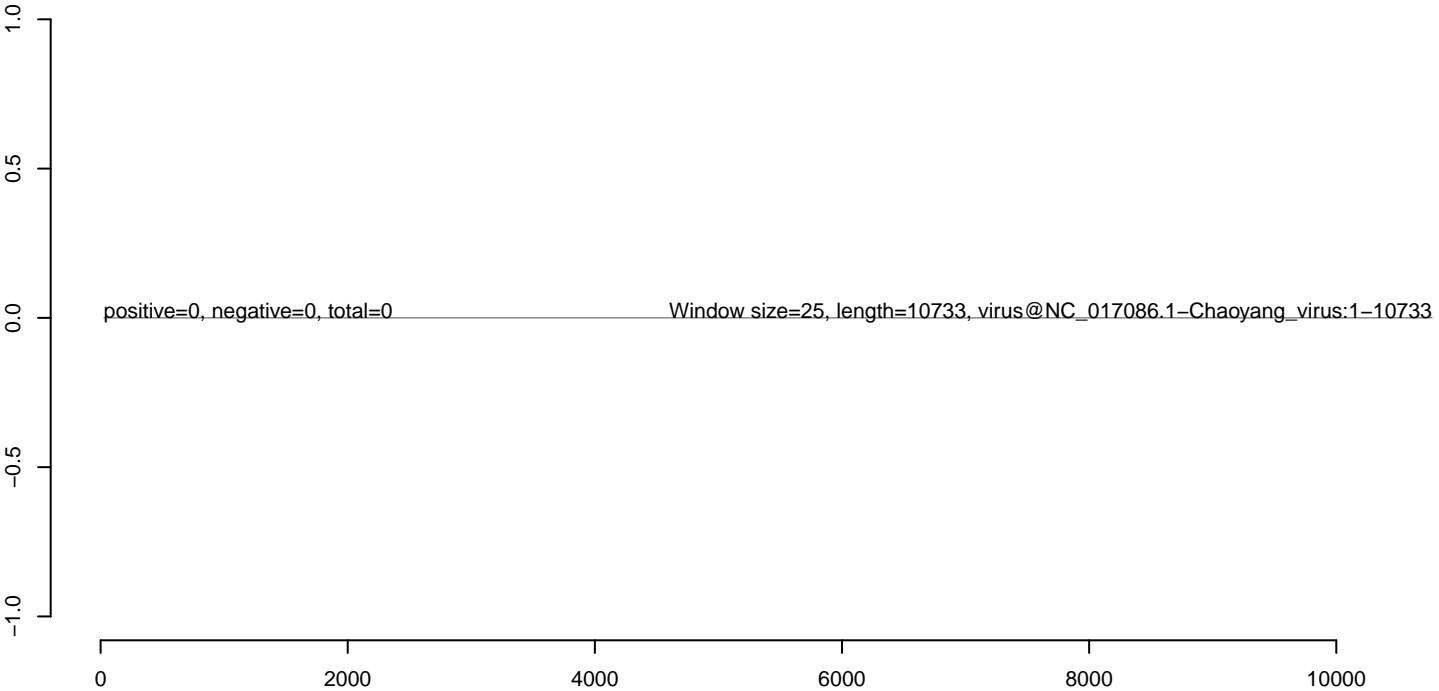
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=10791, virus@NC_016997.1-Donggang_virus:1-10791

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=11877, virus@NC_016962.1-Bebaru_virus:1-11877

0 2000 4000 6000 8000 10000 12000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

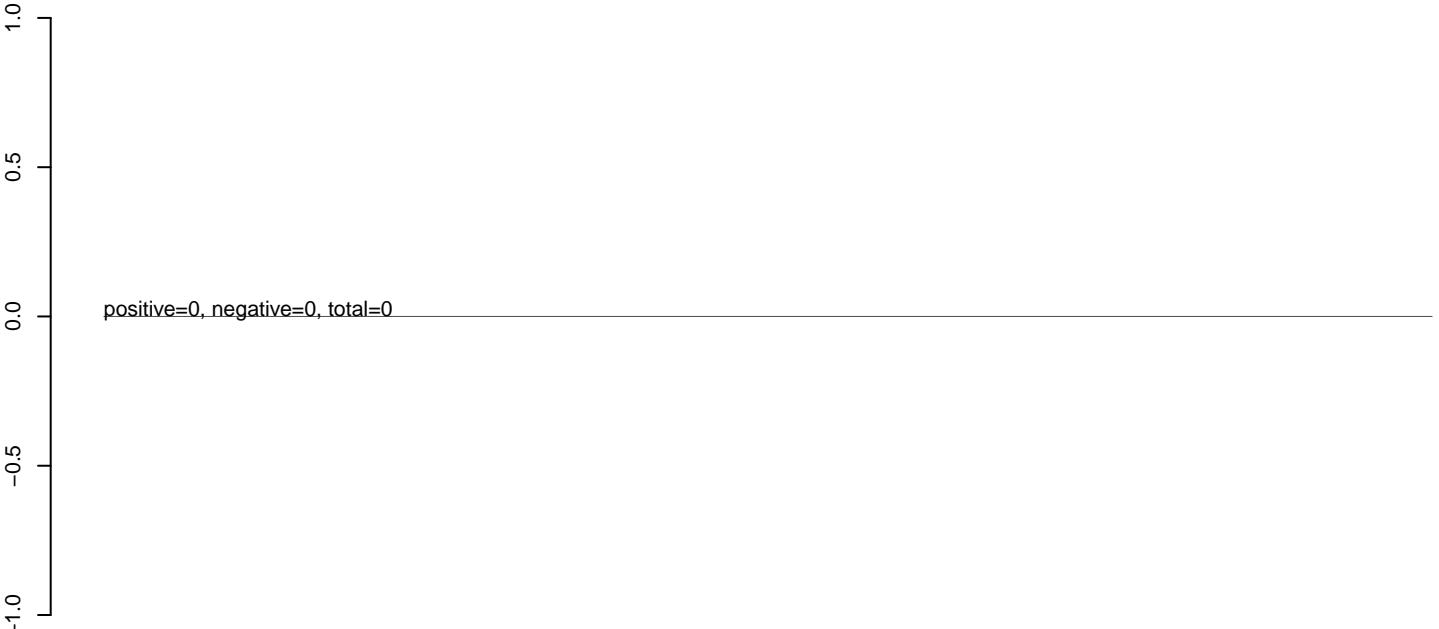
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

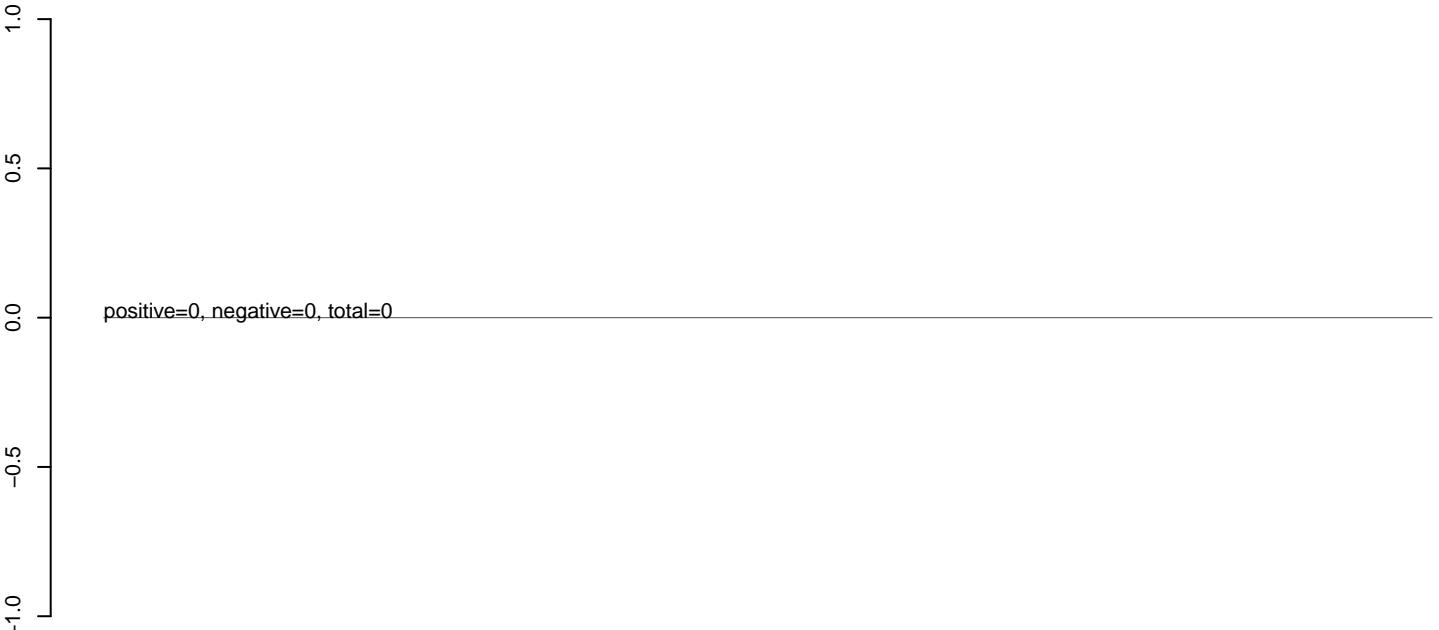
CuQuin_MG_7dpi3.rep

positive=0, negative=25, total=25
Virus@NC_014395.1-Rift_Valley_fever_virus_segment_S_plus_segment_M_plus_segment_L:1-11979

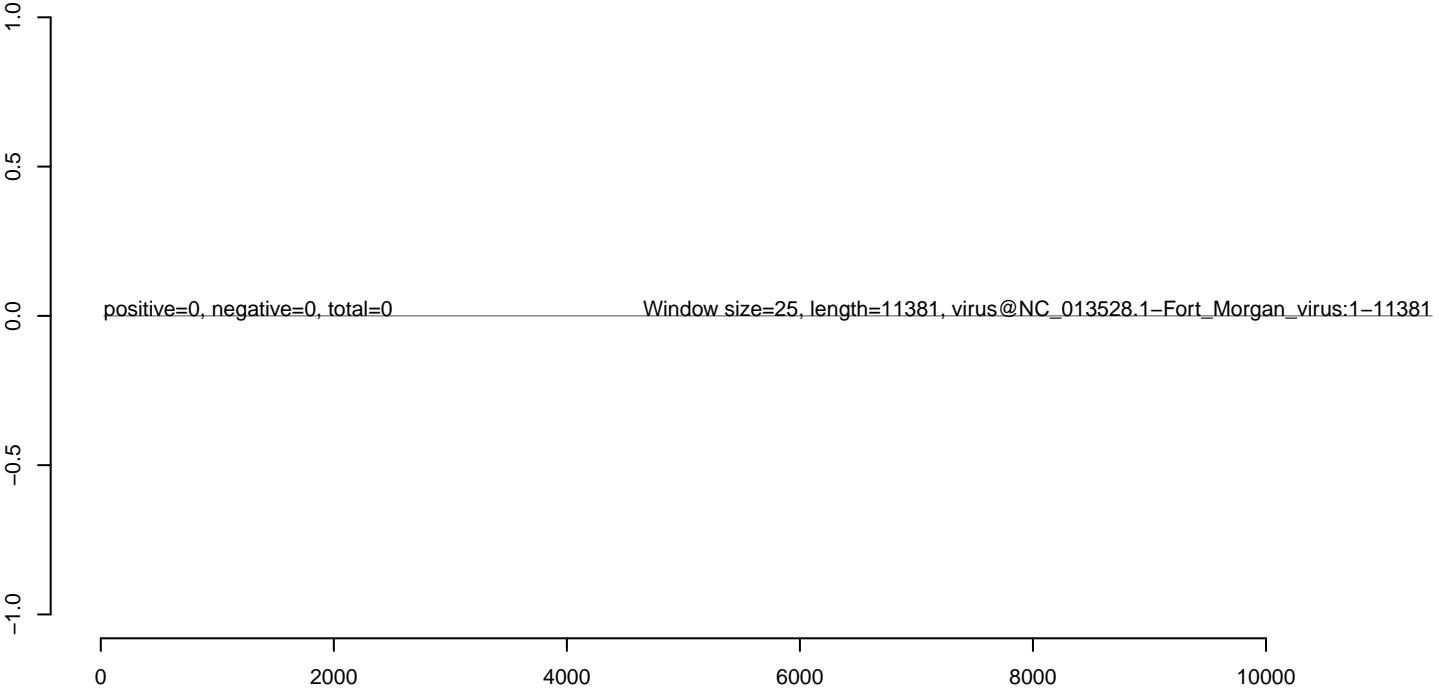
CuQuin_MG_7dpi3.18_23.rep



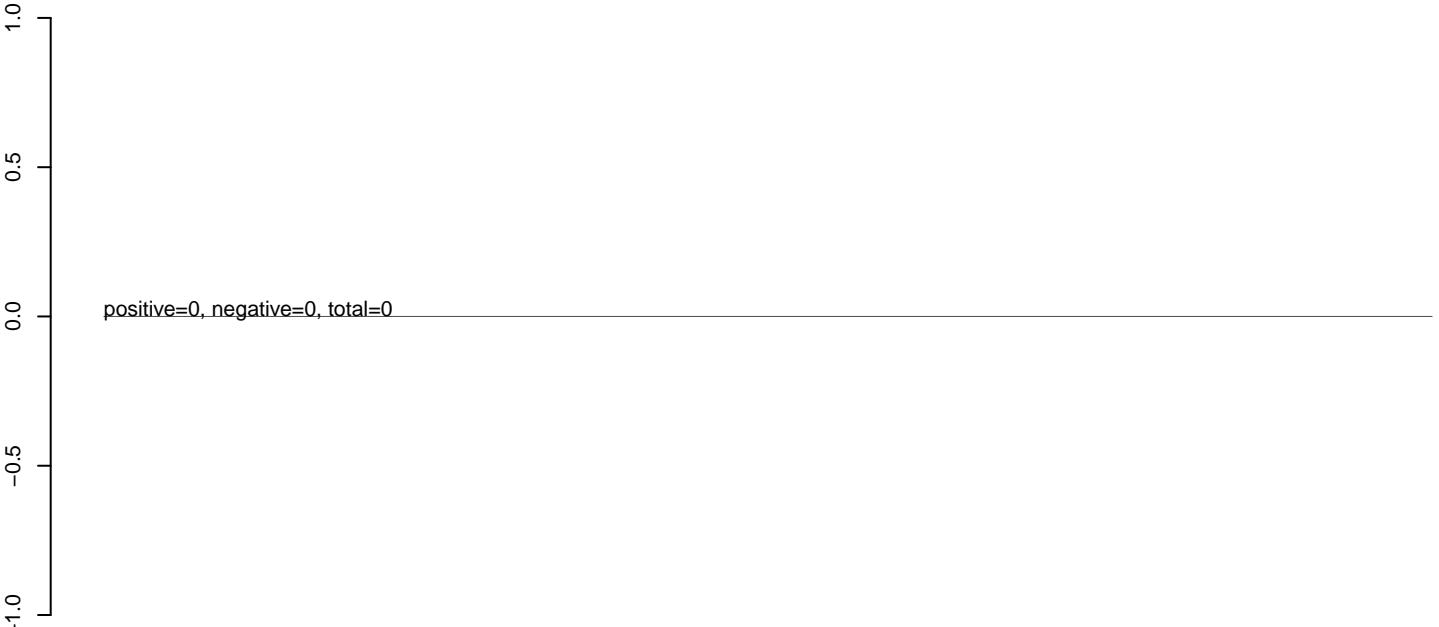
CuQuin_MG_7dpi3.24_35.rep



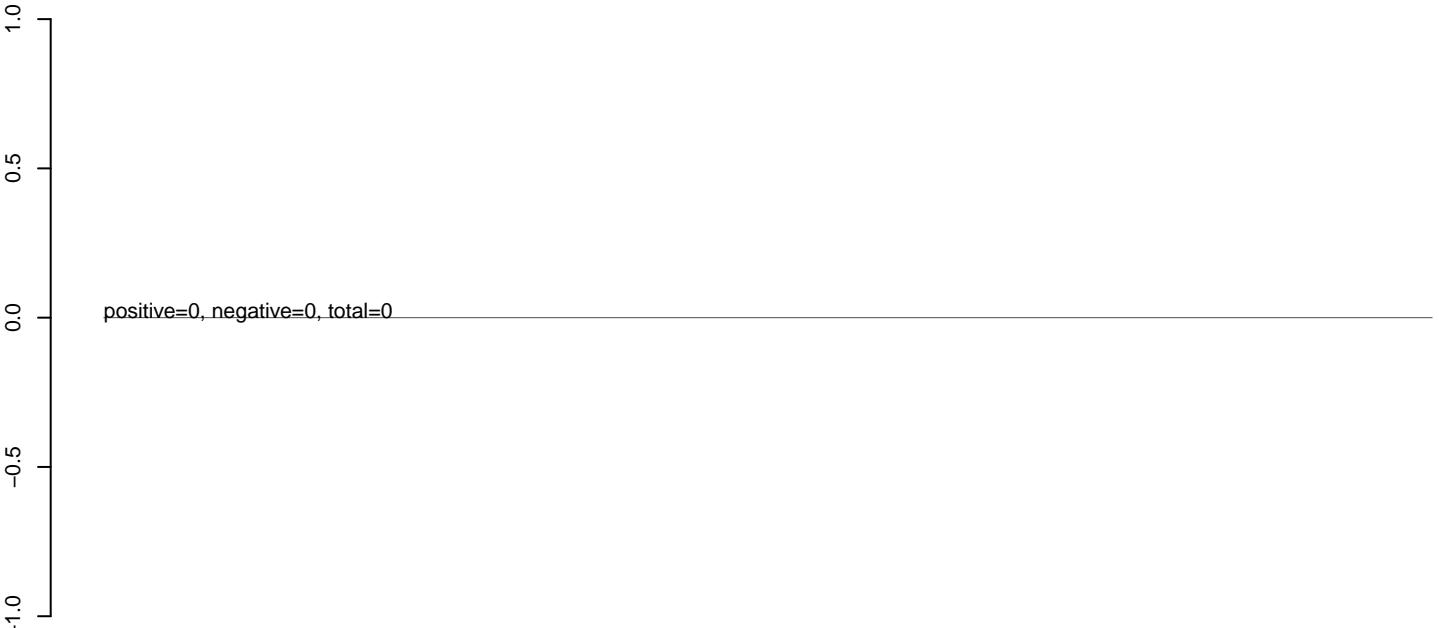
CuQuin_MG_7dpi3.rep



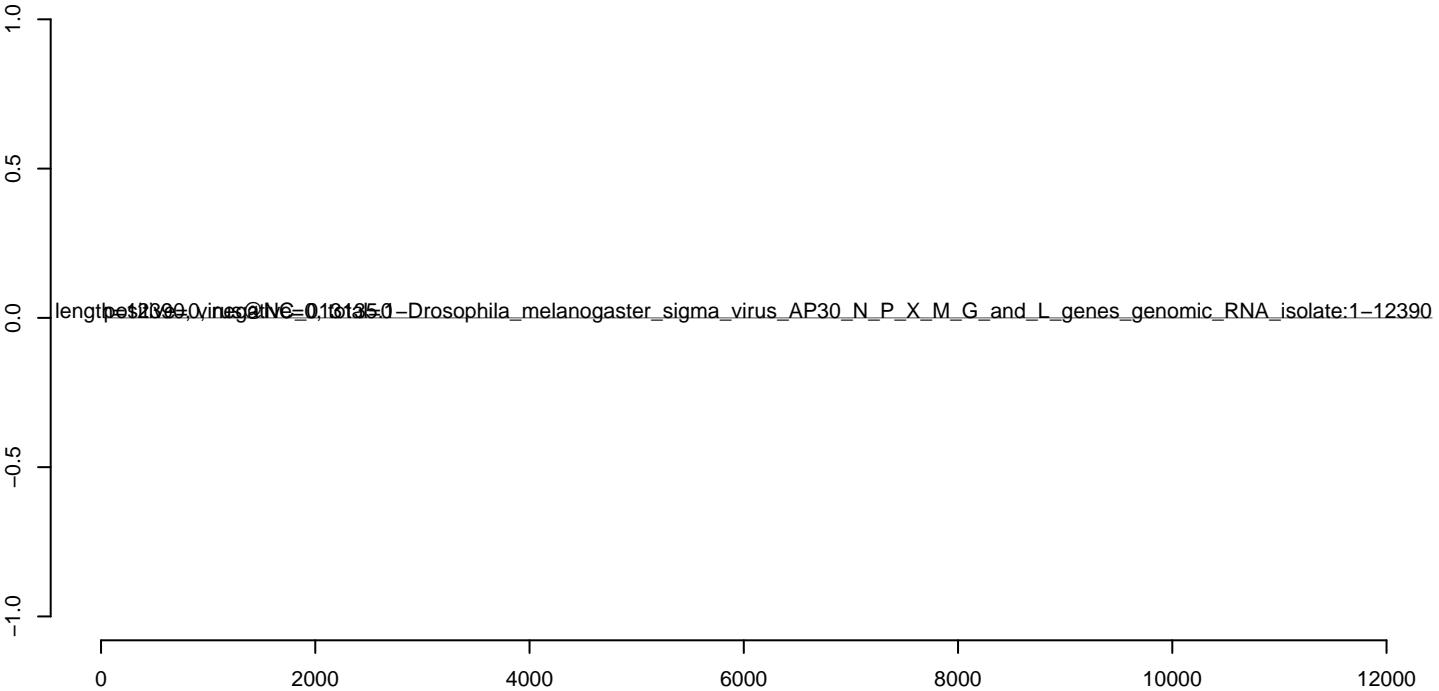
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=4806, virus@NC_012958.1-Drosophila_A_virus:1-4806

0 1000 2000 3000 4000 5000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0, window size=25, length=11064, virus@NC_012932.1-Aedes_flavivirus_genomic_RNA_strain_Narita-21:1-11064

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

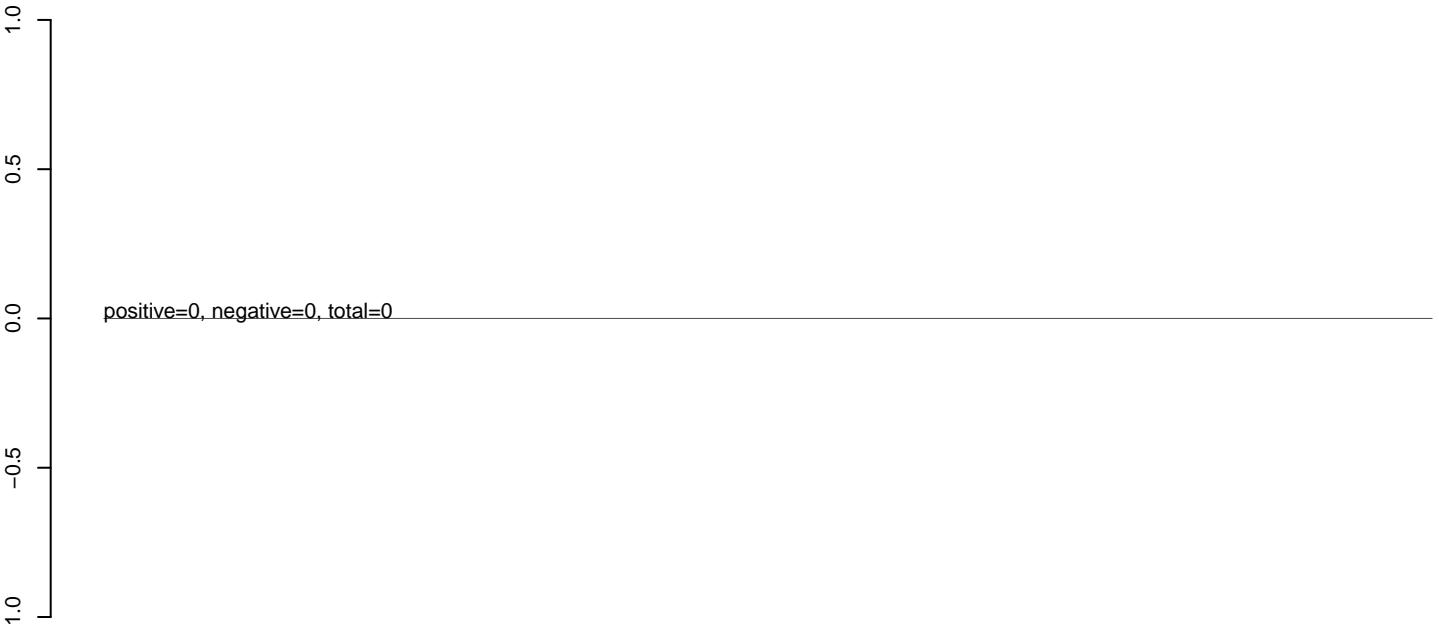
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

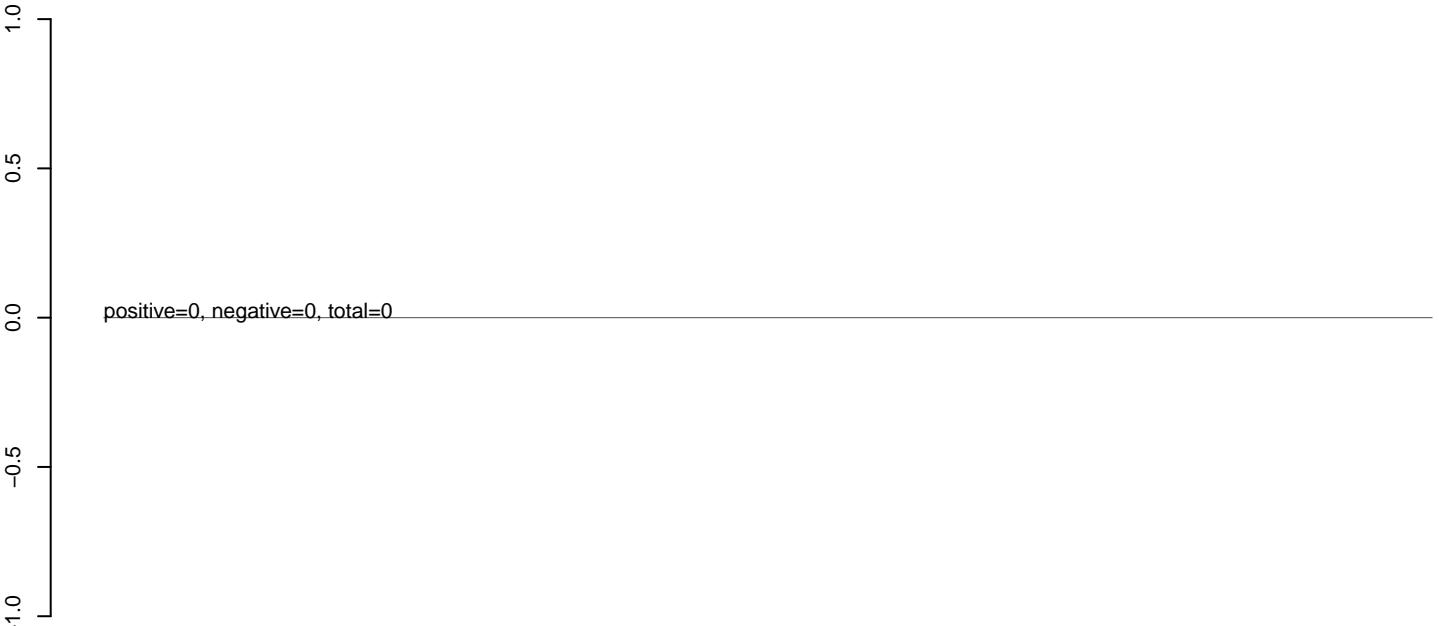
Window size=25, length=10814, virus@NC_012735.1-Wesselsbron_virus:1-10814

0 2000 4000 6000 8000 10000

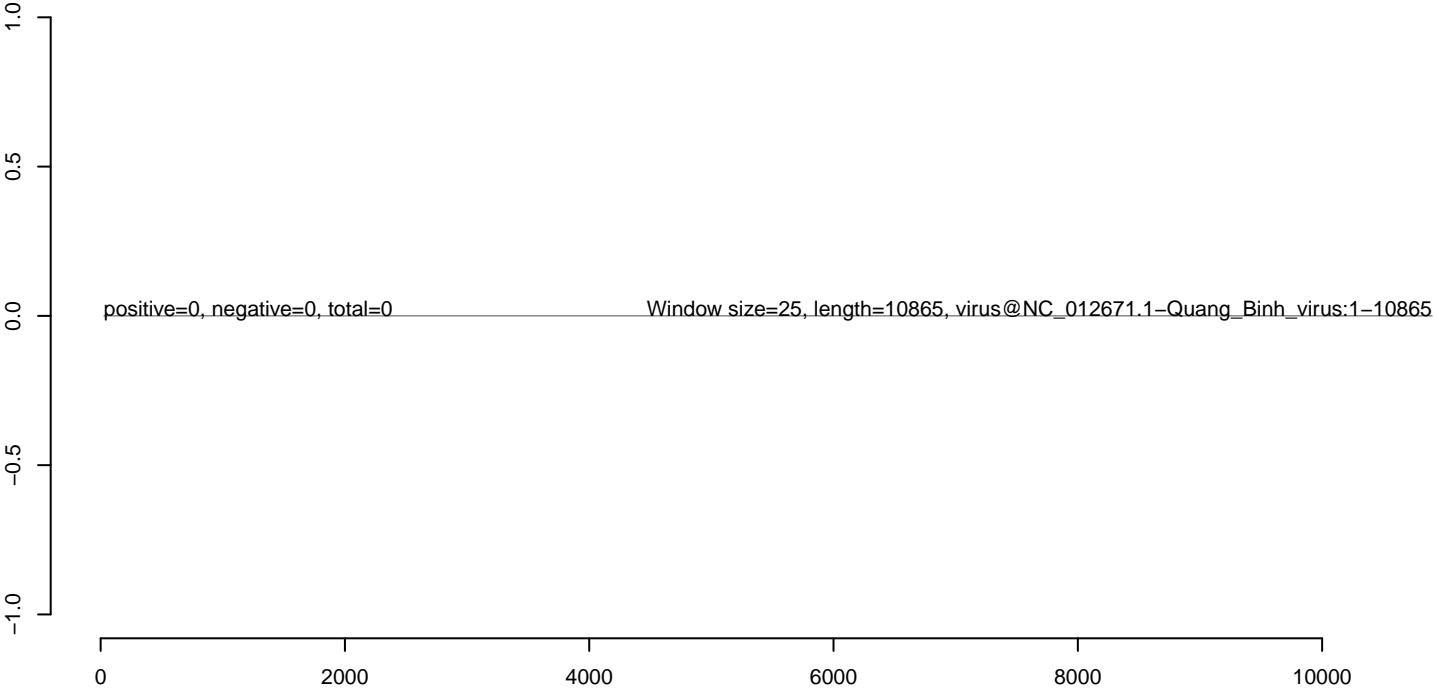
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=3776, virus@NC_012636.1-Aedes_aegypti_densovirus_2_strain_0814616:1-3776

0 1000 2000 3000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

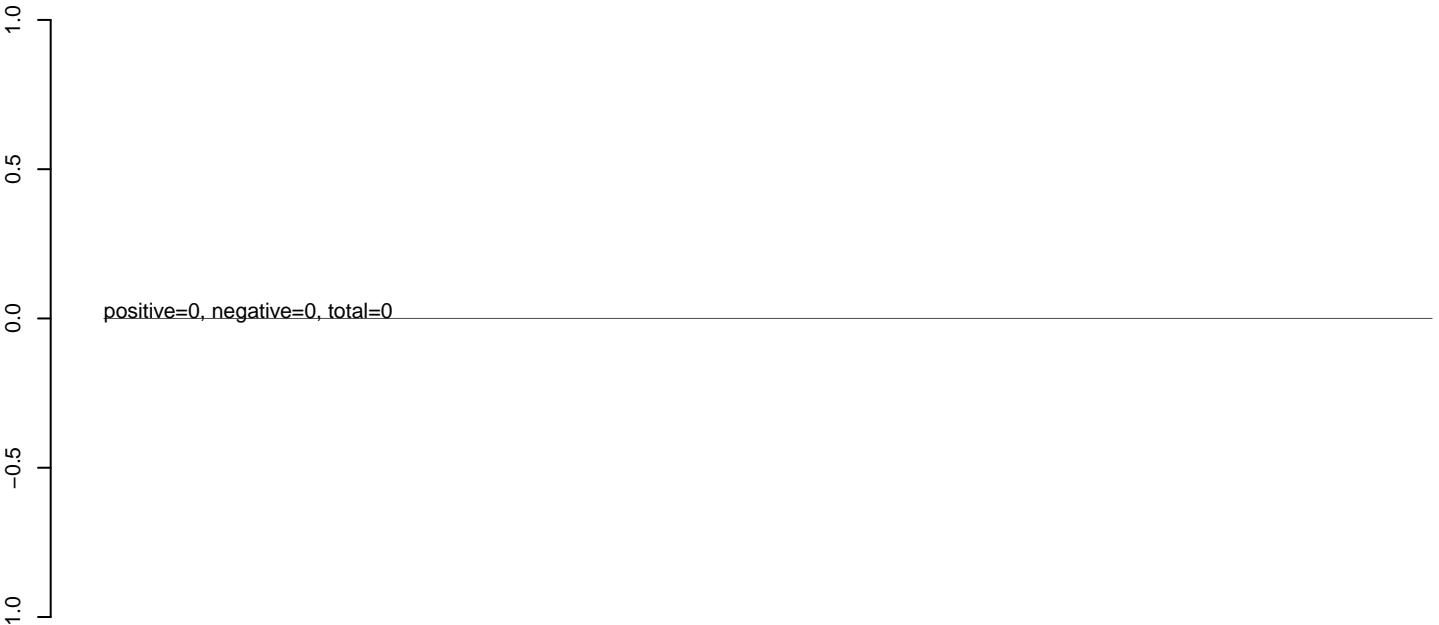
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

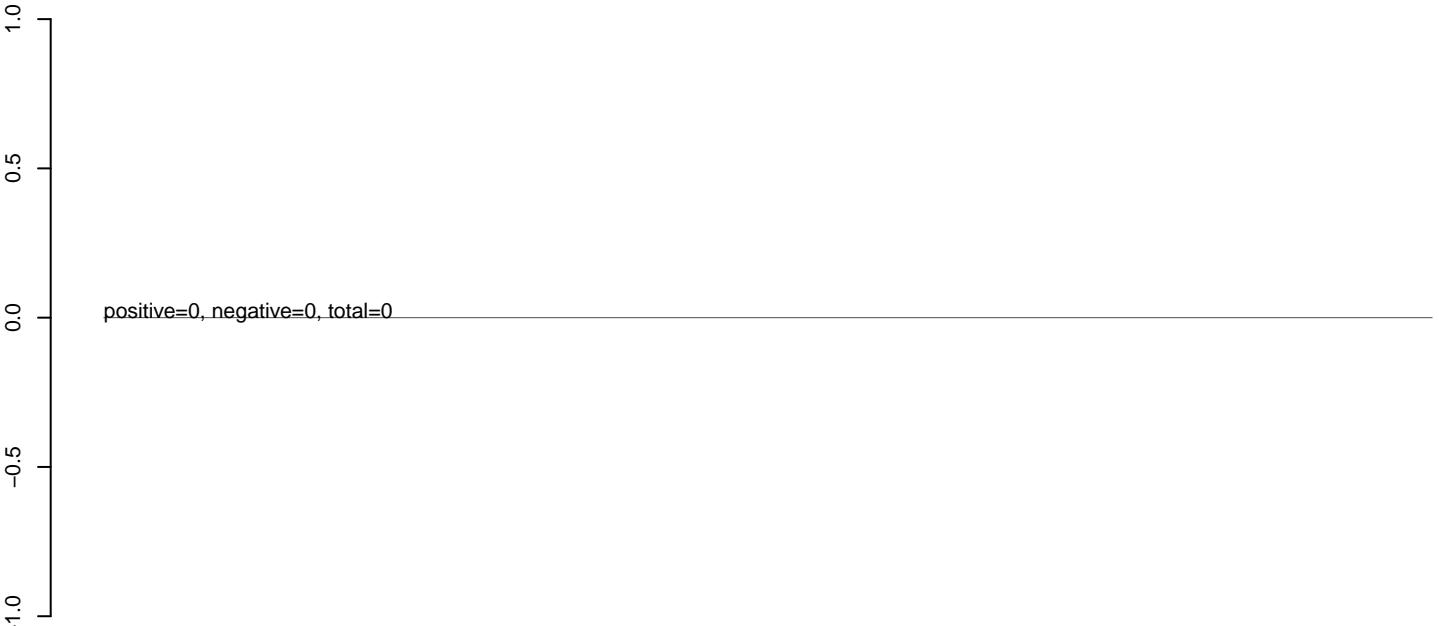
Window size=25, length=11526, virus@NC_012561.1-Highlands_J_virus:1-11526

0 2000 4000 6000 8000 10000 12000

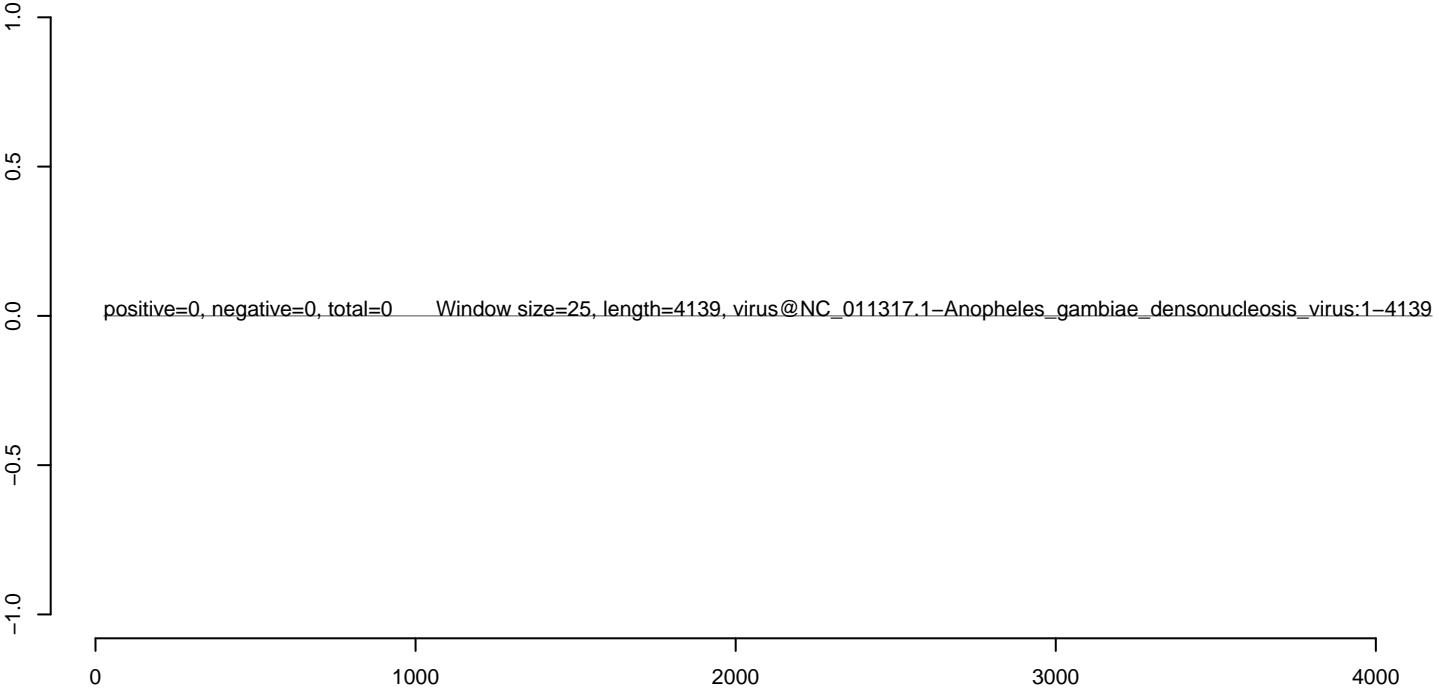
CuQuin_MG_7dpi3.18_23.rep



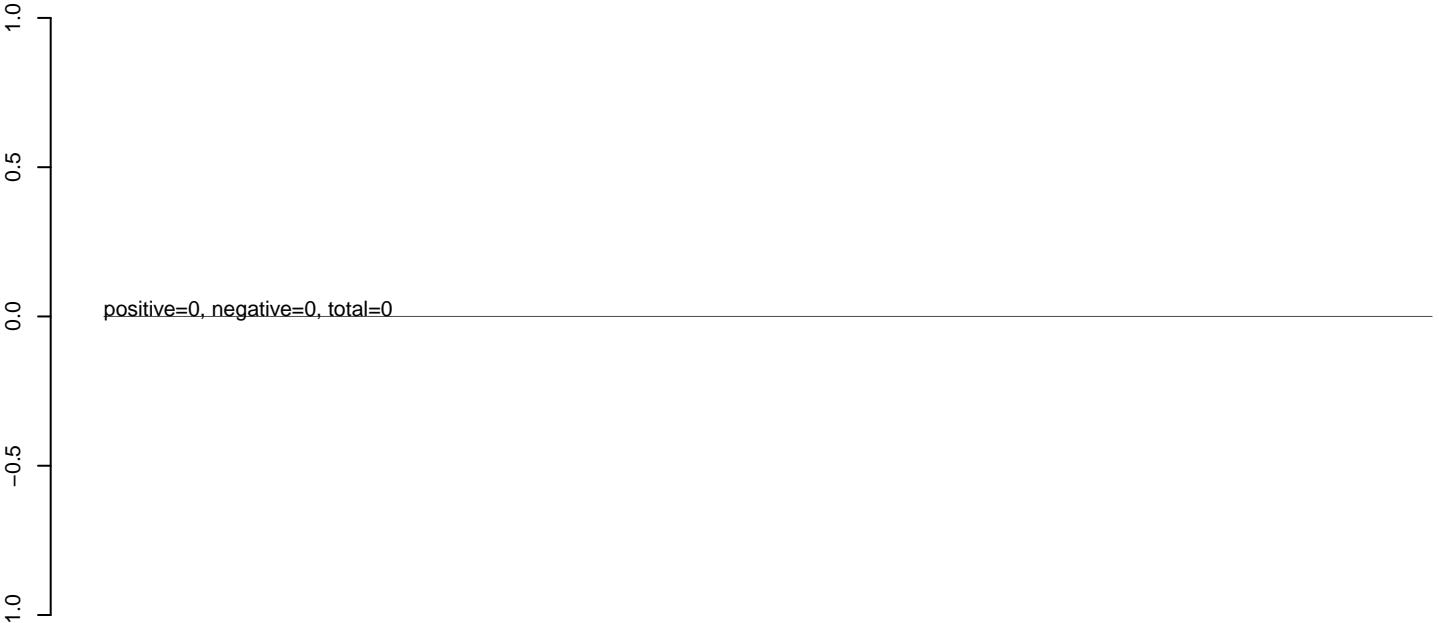
CuQuin_MG_7dpi3.24_35.rep



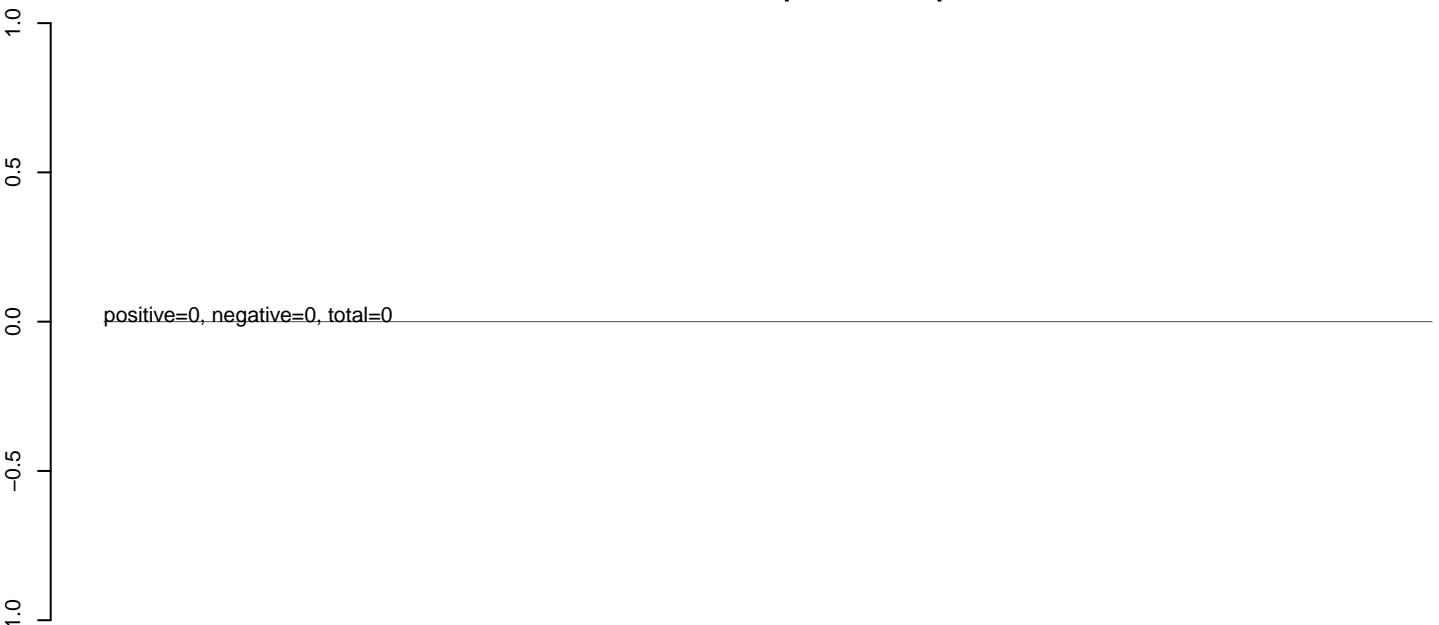
CuQuin_MG_7dpi3.rep



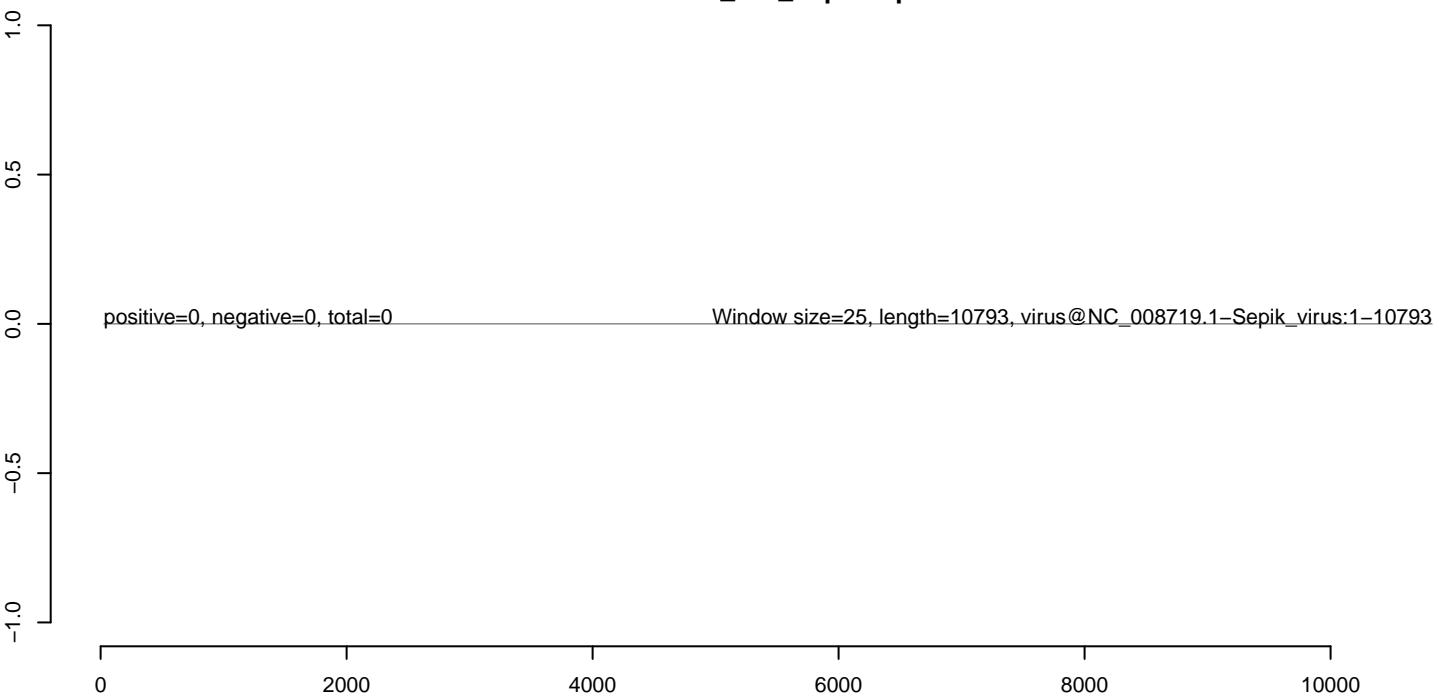
CuQuin_MG_7dpi3.18_23.rep



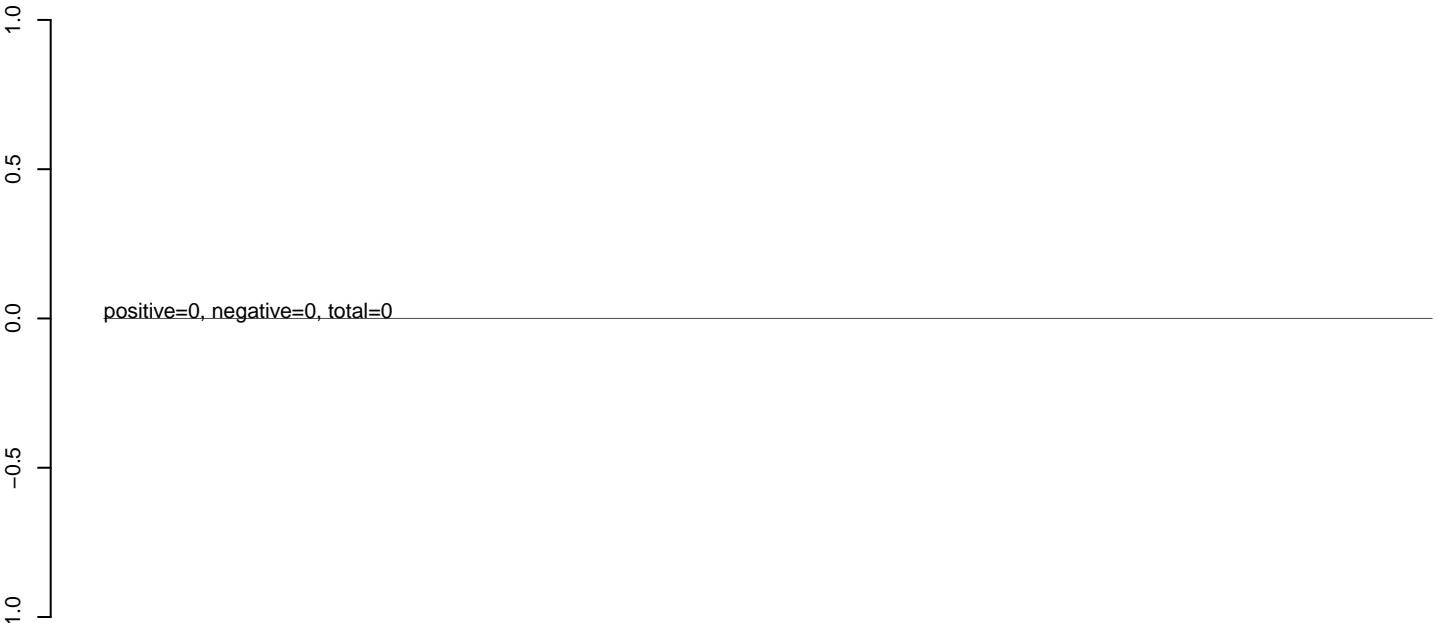
CuQuin_MG_7dpi3.24_35.rep



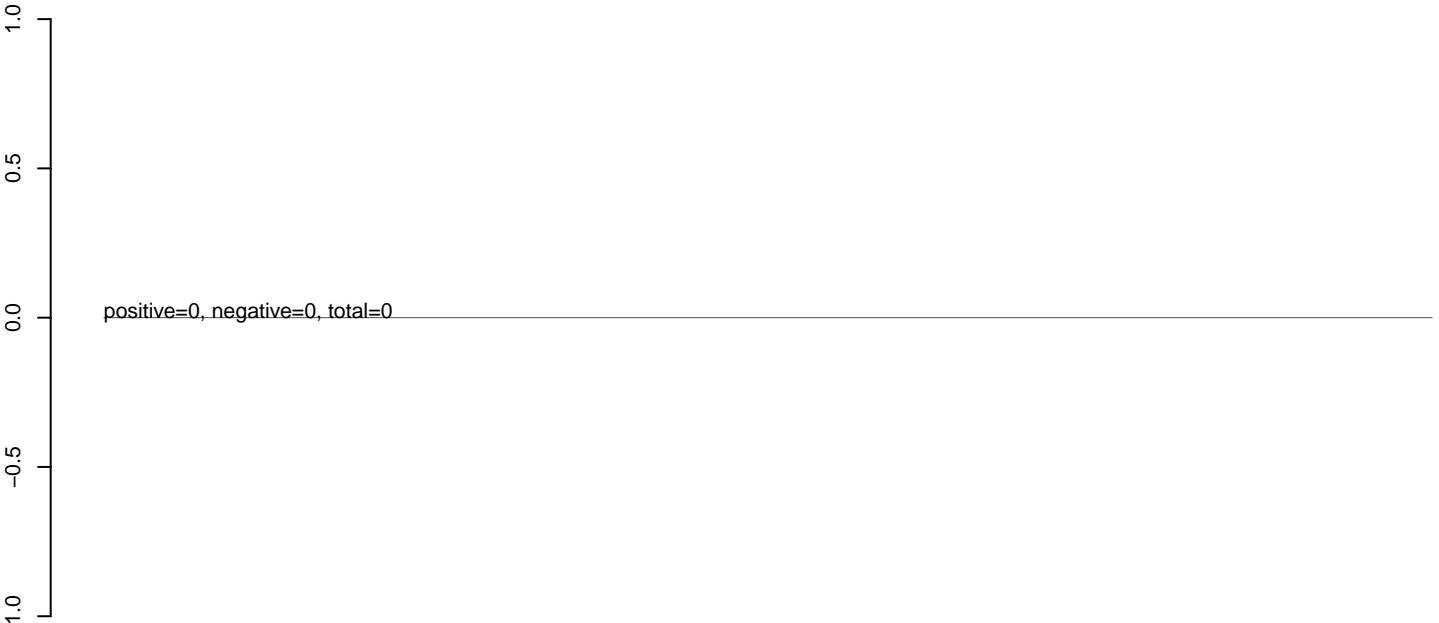
CuQuin_MG_7dpi3.rep



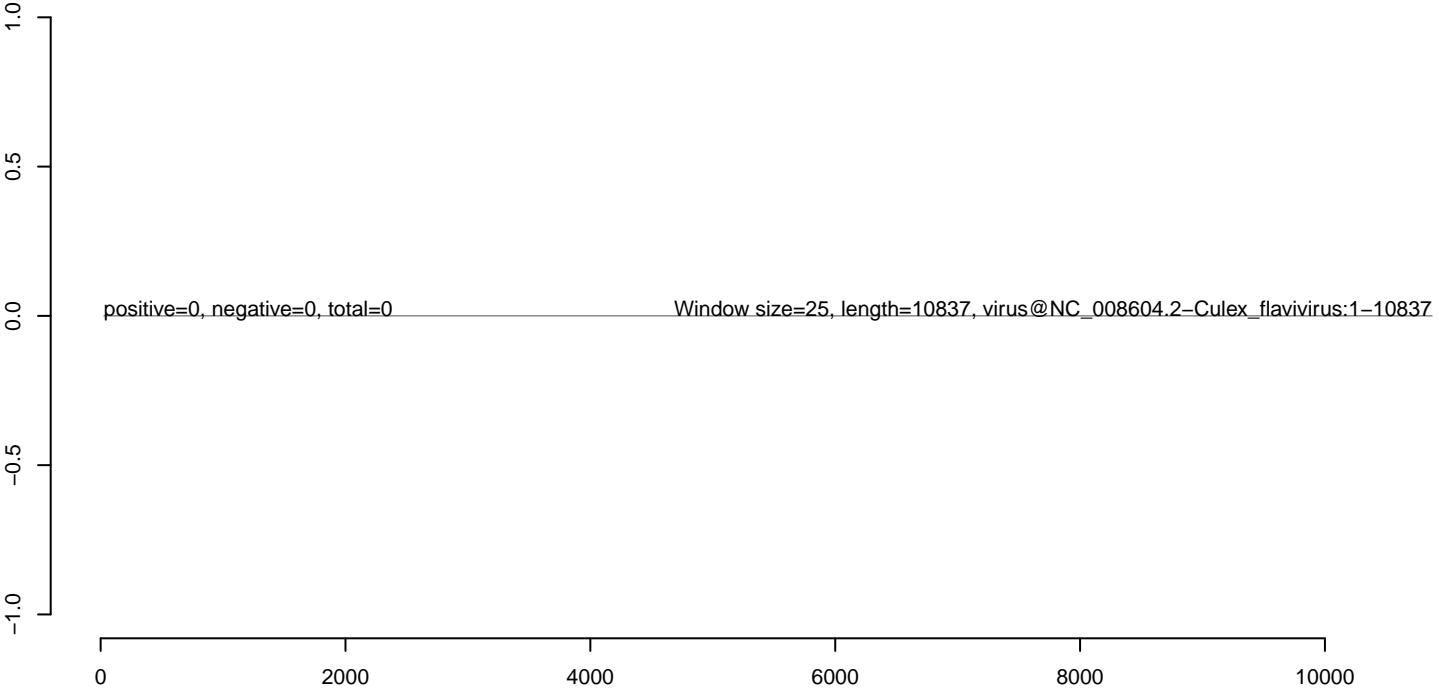
CuQuin_MG_7dpi3.18_23.rep



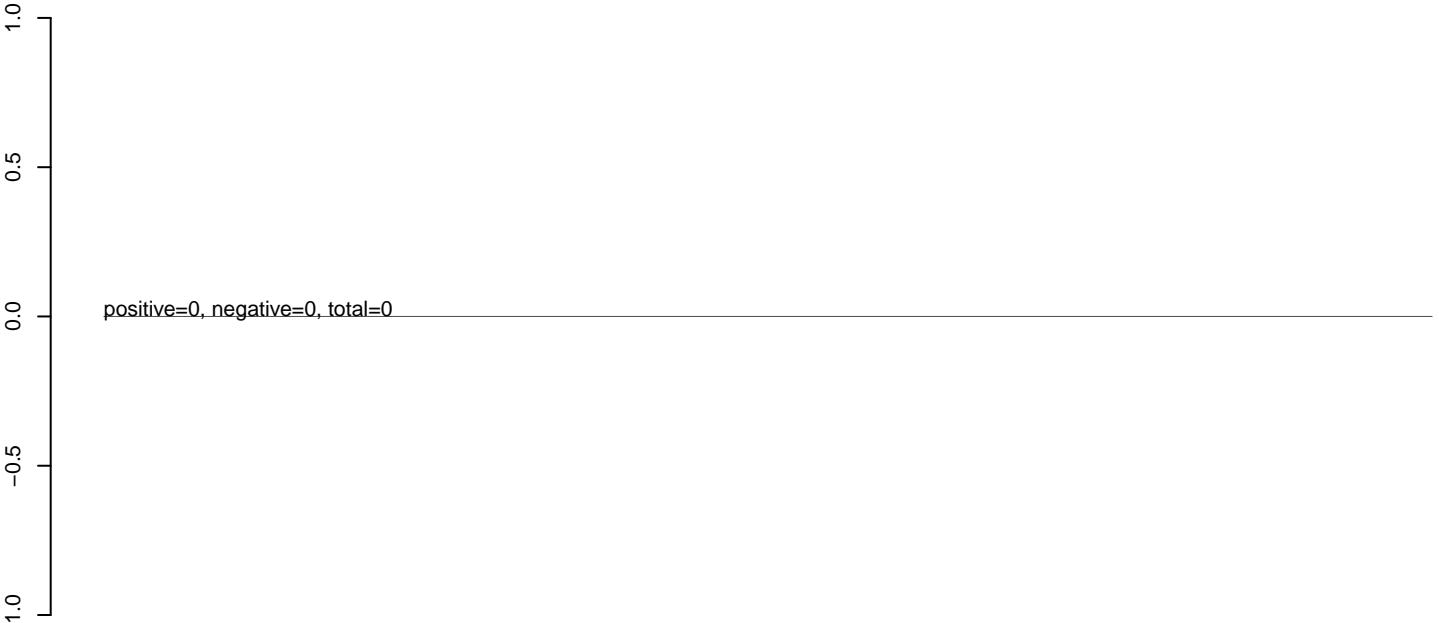
CuQuin_MG_7dpi3.24_35.rep



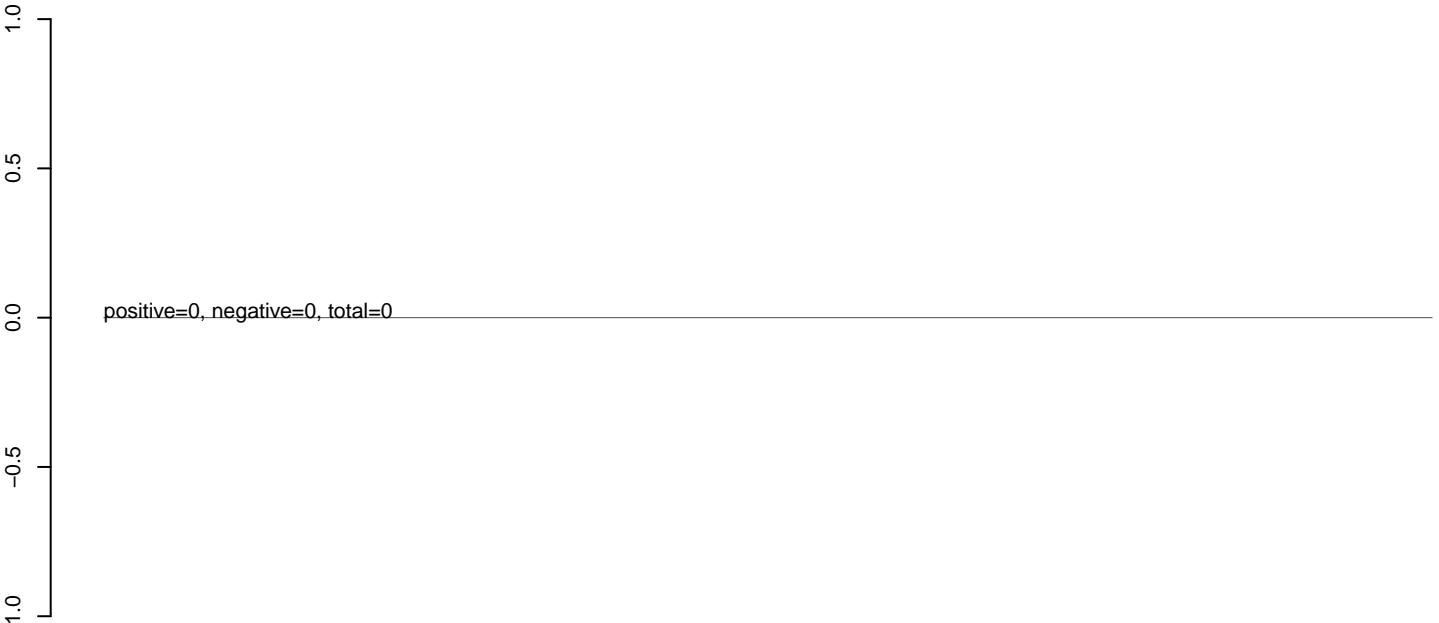
CuQuin_MG_7dpi3.rep



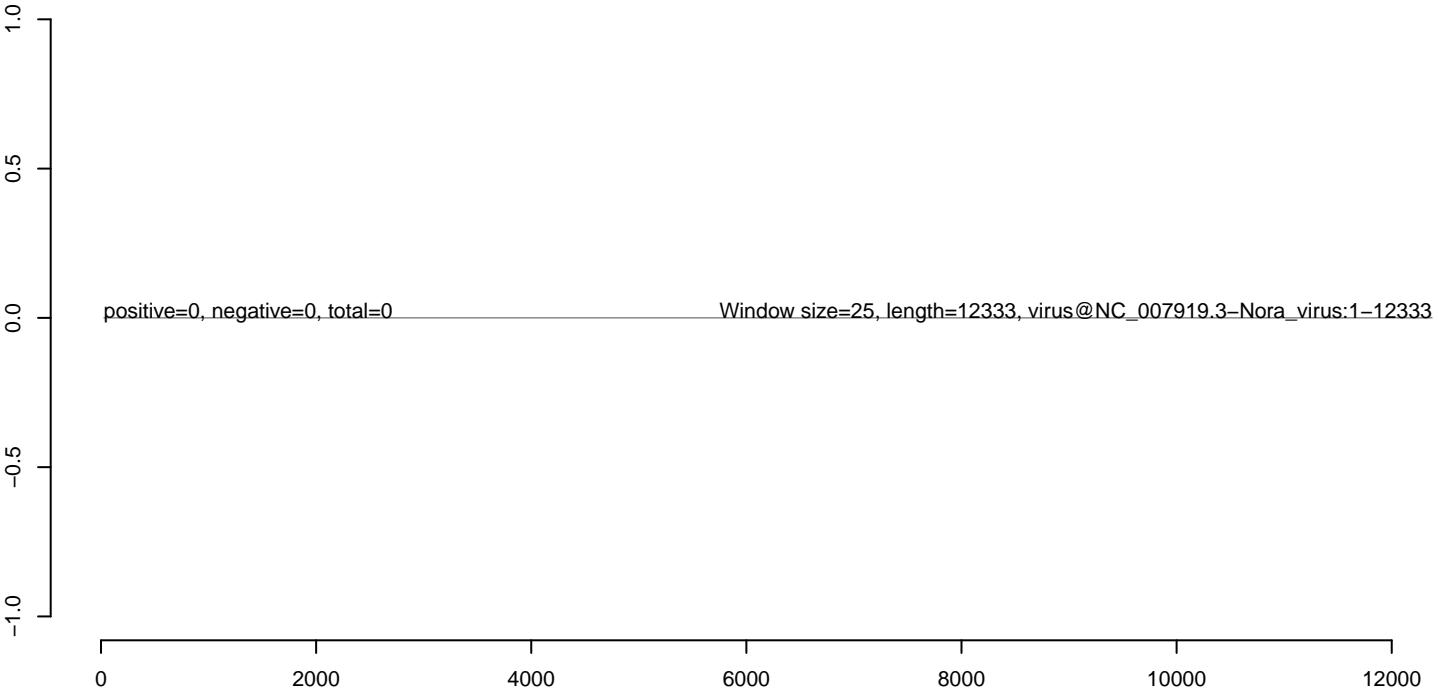
CuQuin_MG_7dpi3.18_23.rep



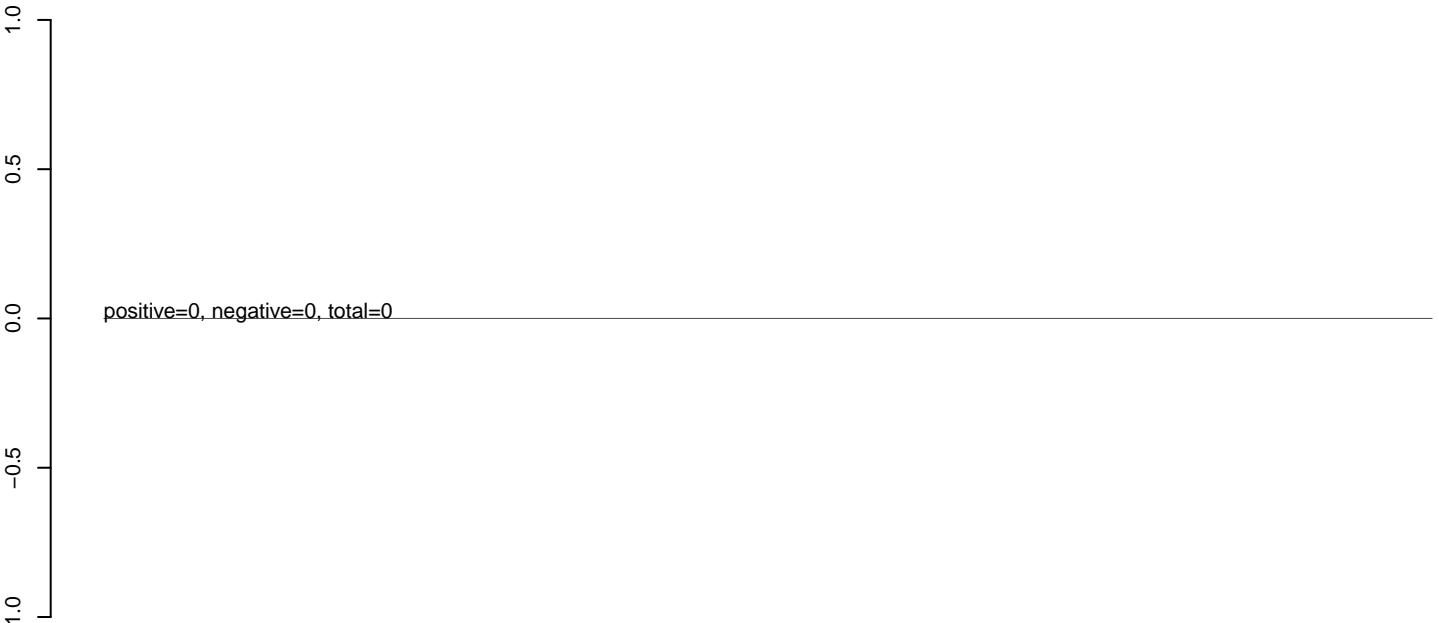
CuQuin_MG_7dpi3.24_35.rep



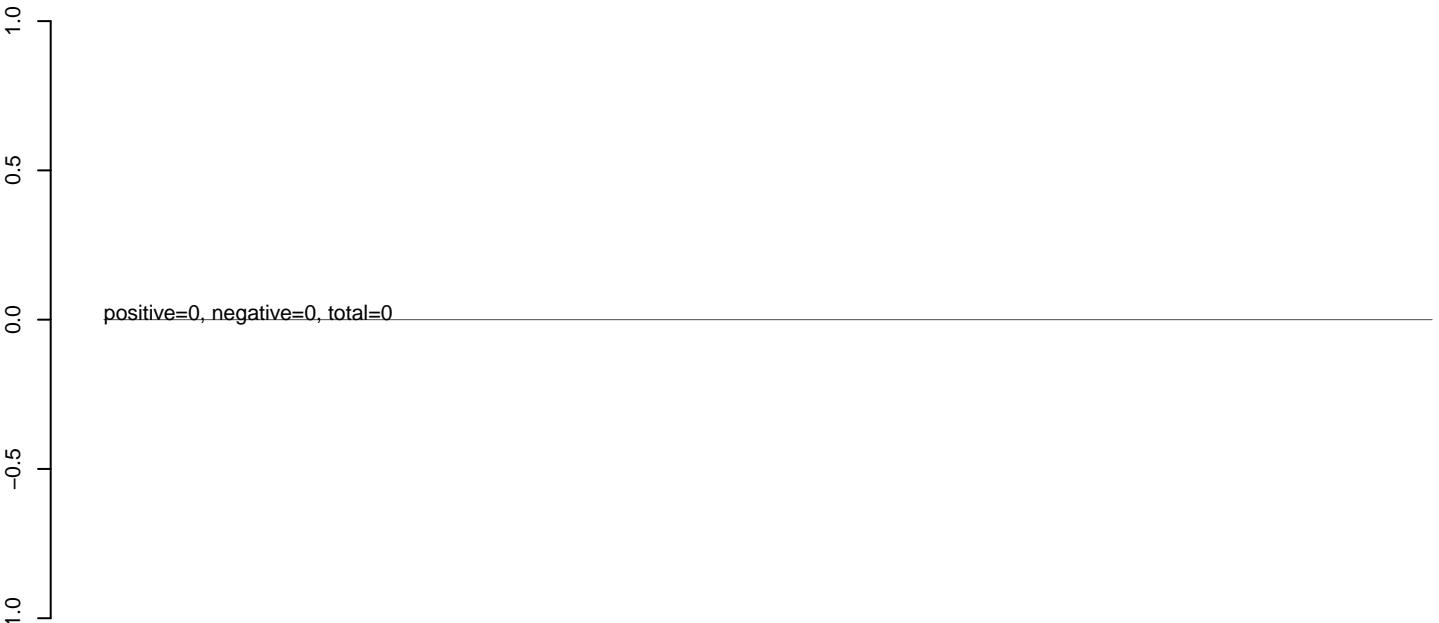
CuQuin_MG_7dpi3.rep



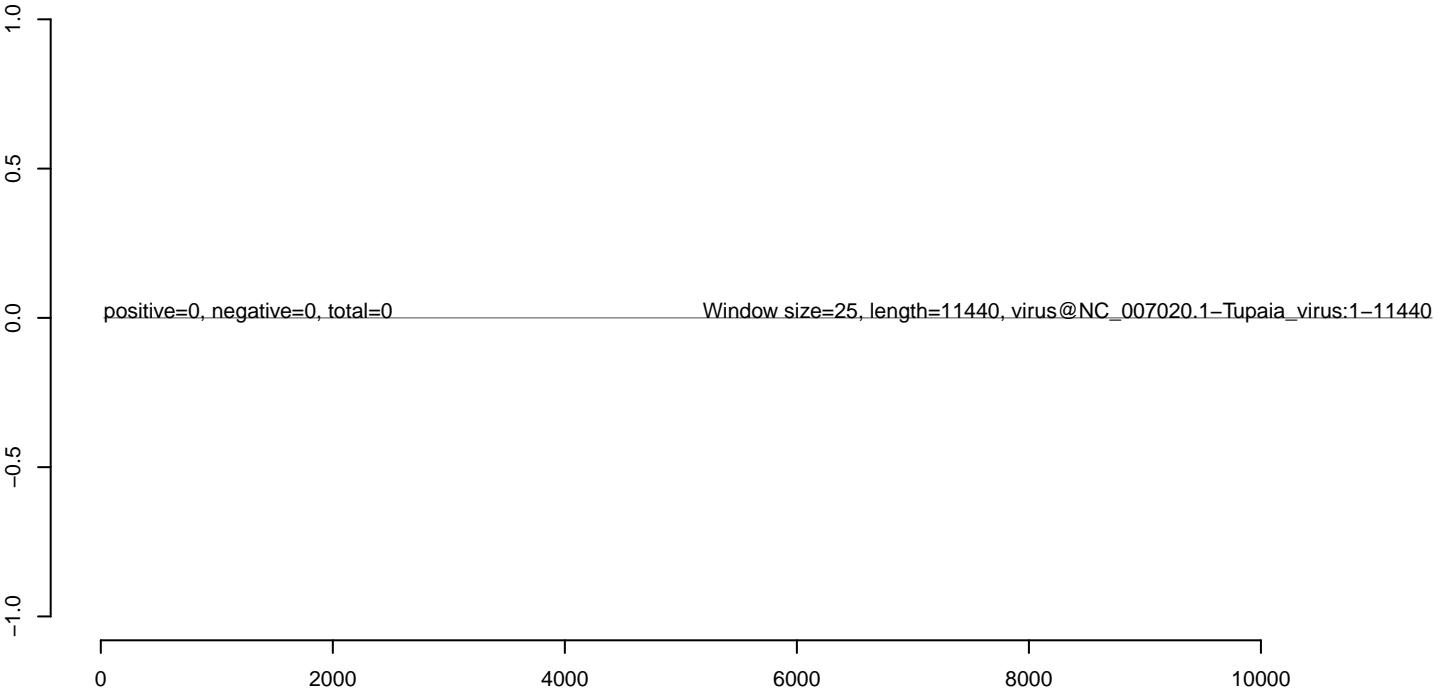
CuQuin_MG_7dpi3.18_23.rep



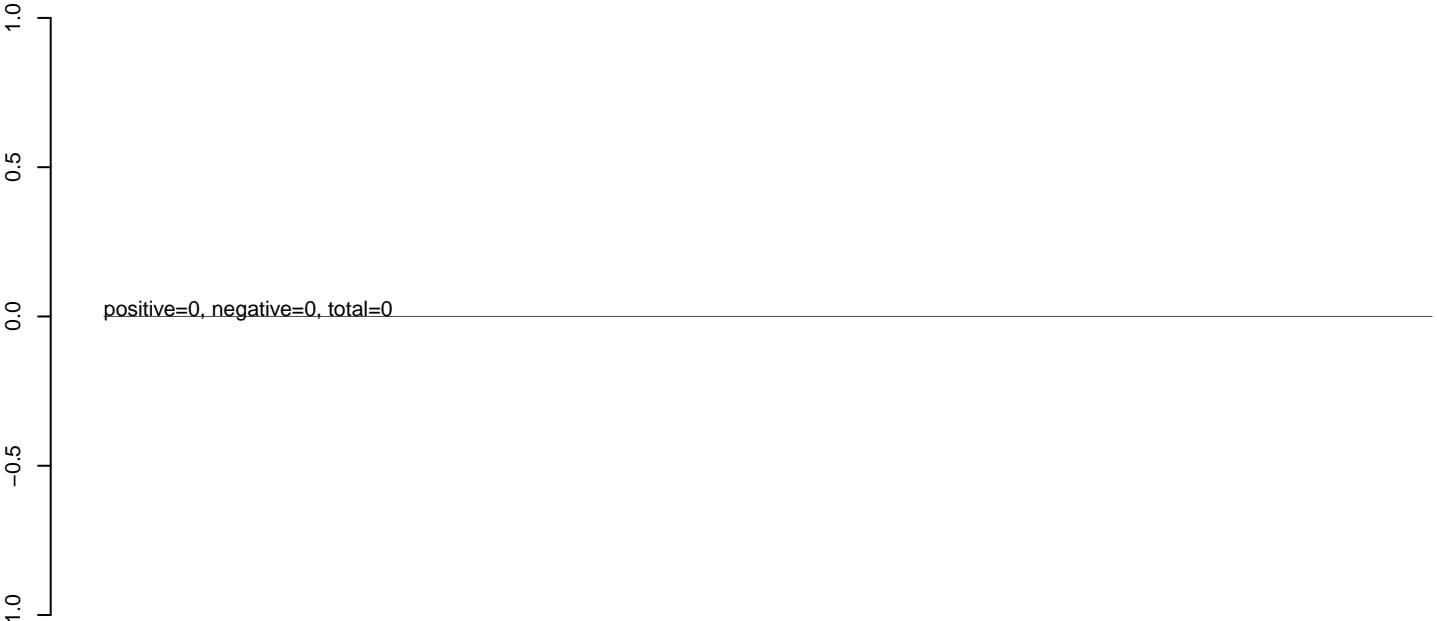
CuQuin_MG_7dpi3.24_35.rep



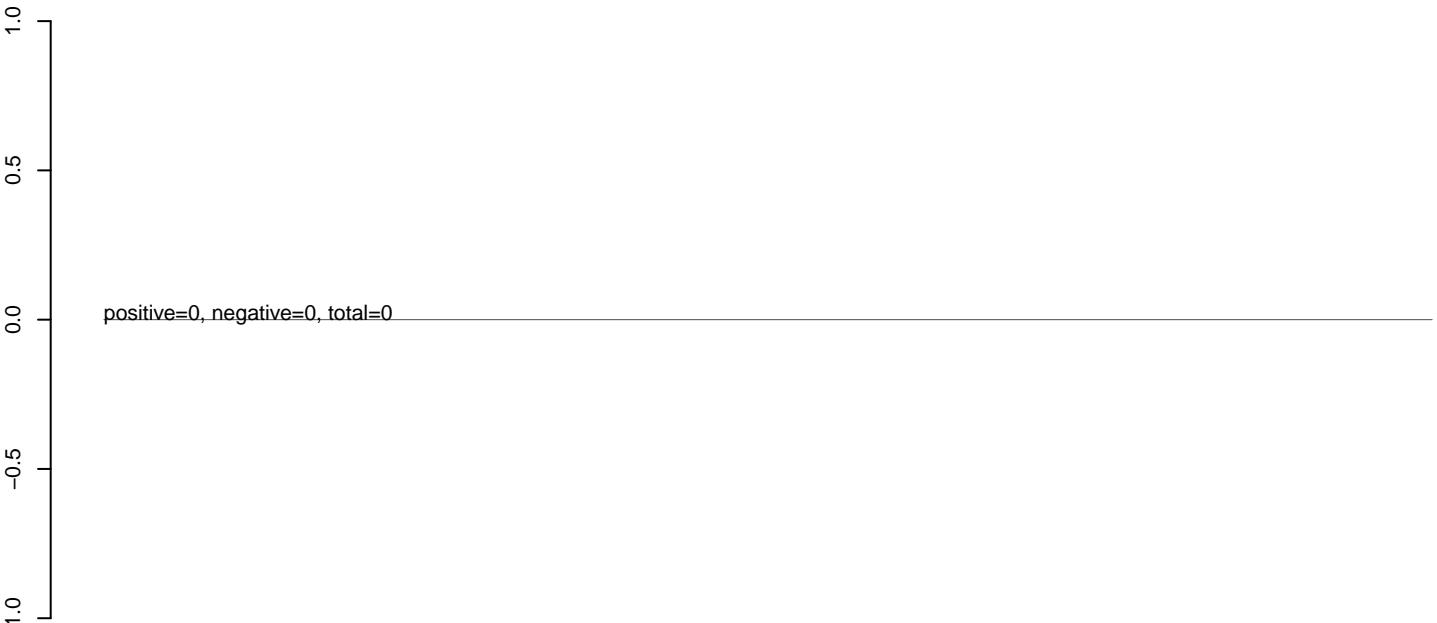
CuQuin_MG_7dpi3.rep



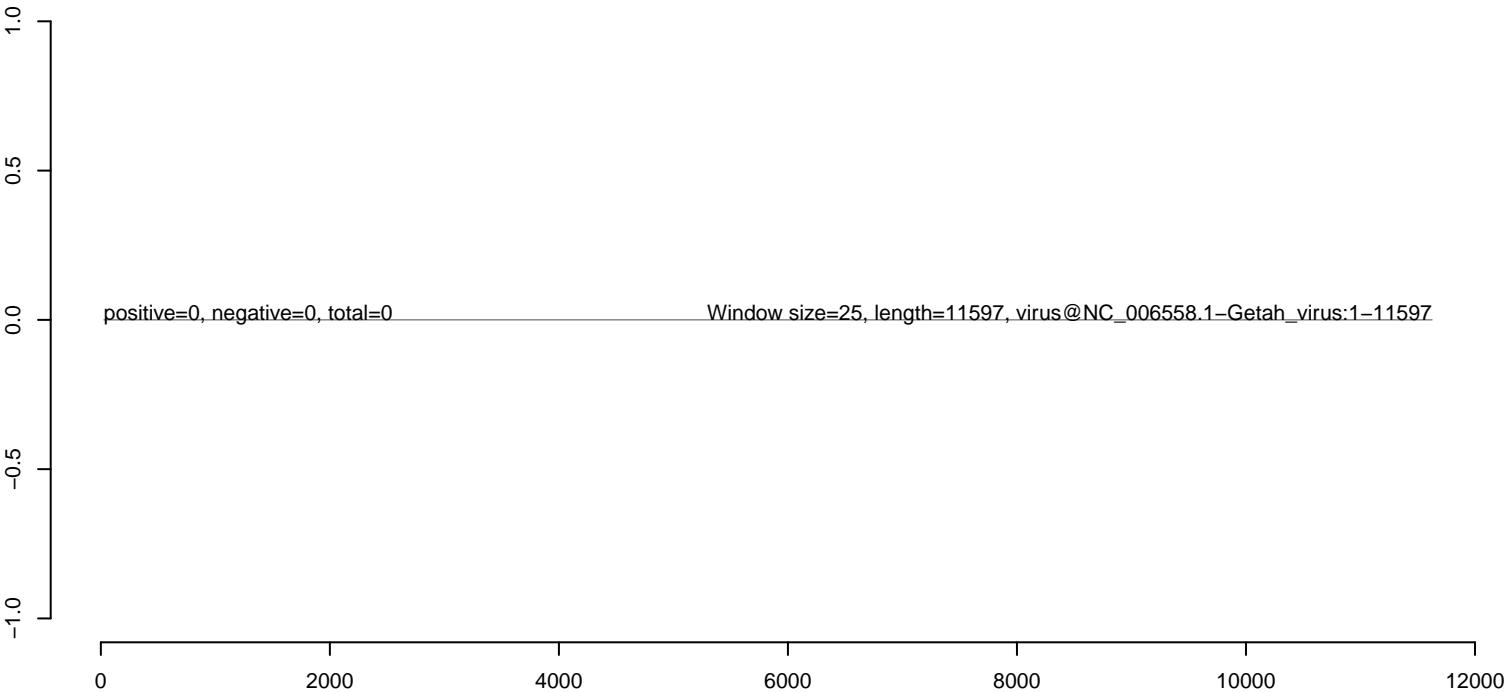
CuQuin_MG_7dpi3.18_23.rep



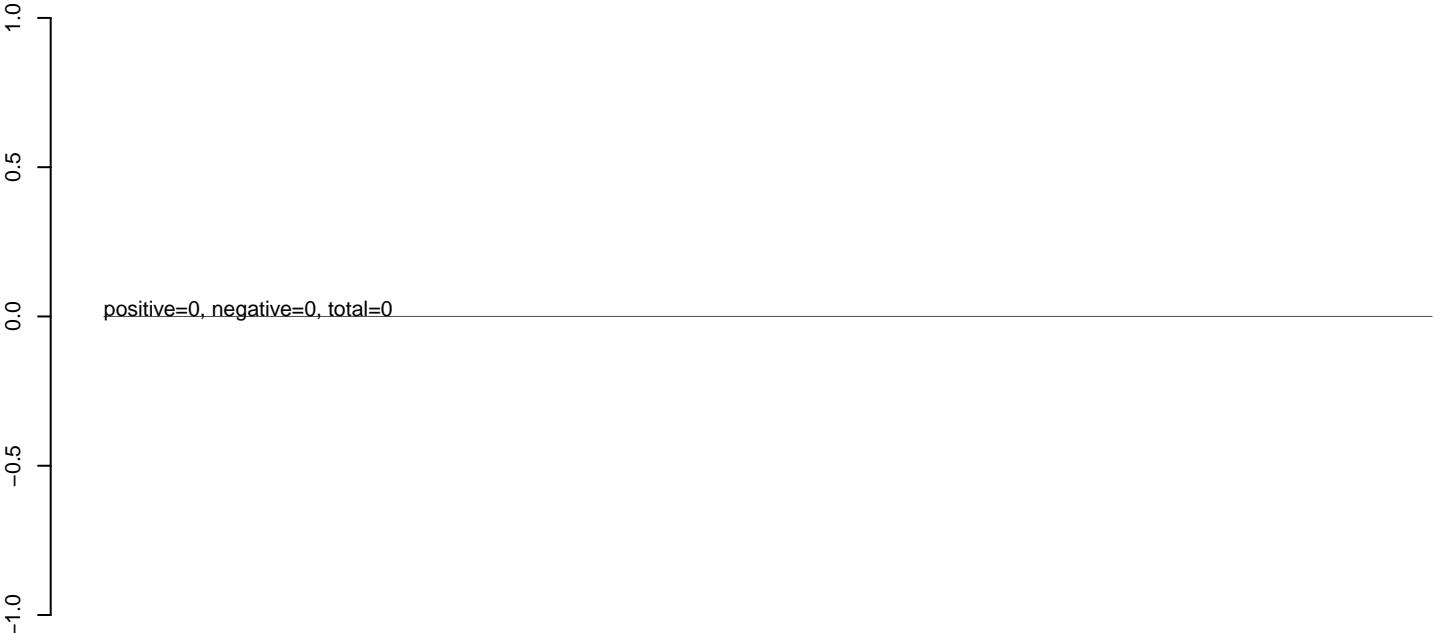
CuQuin_MG_7dpi3.24_35.rep



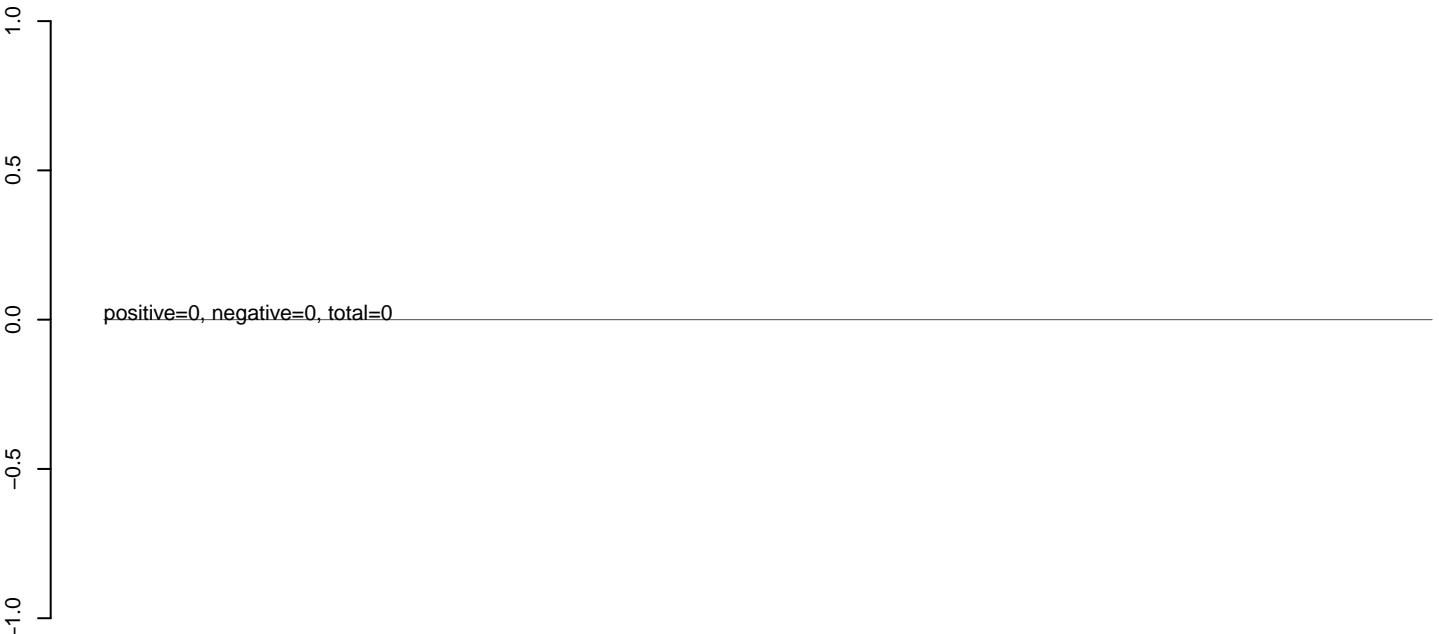
CuQuin_MG_7dpi3.rep



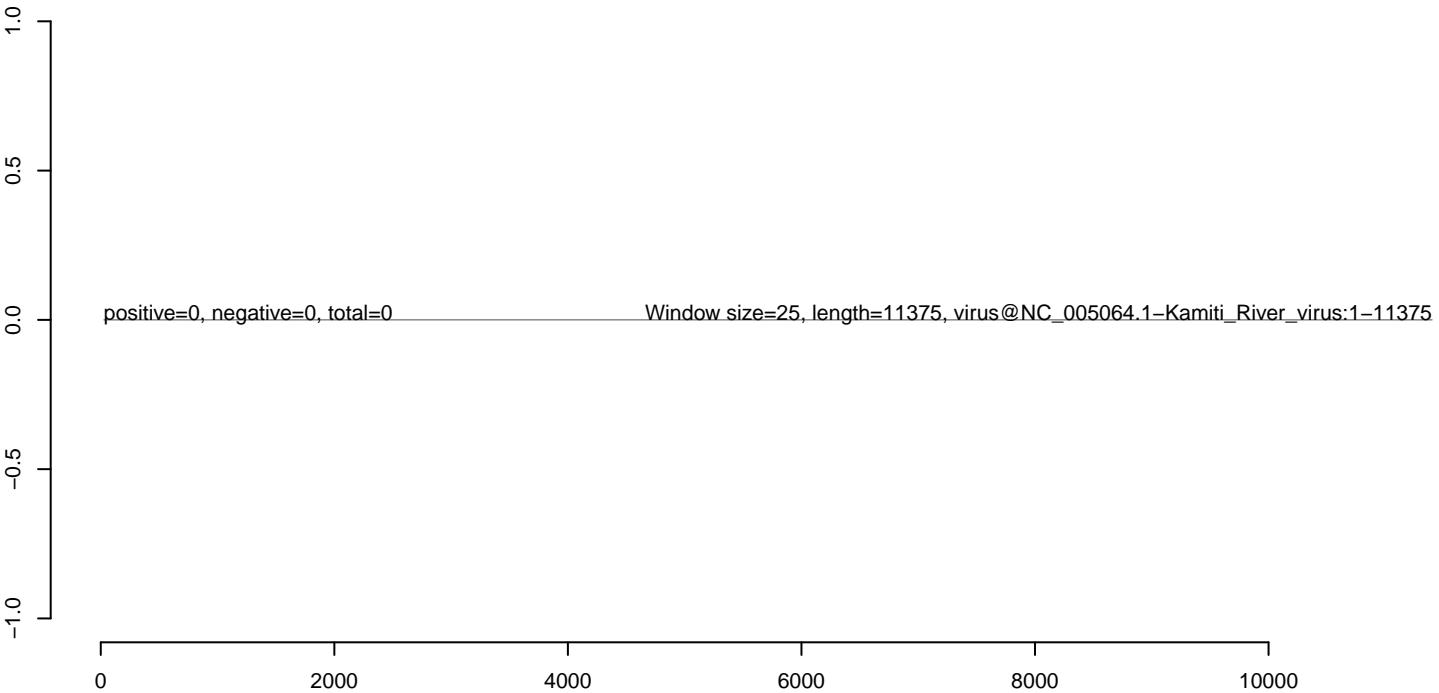
CuQuin_MG_7dpi3.18_23.rep



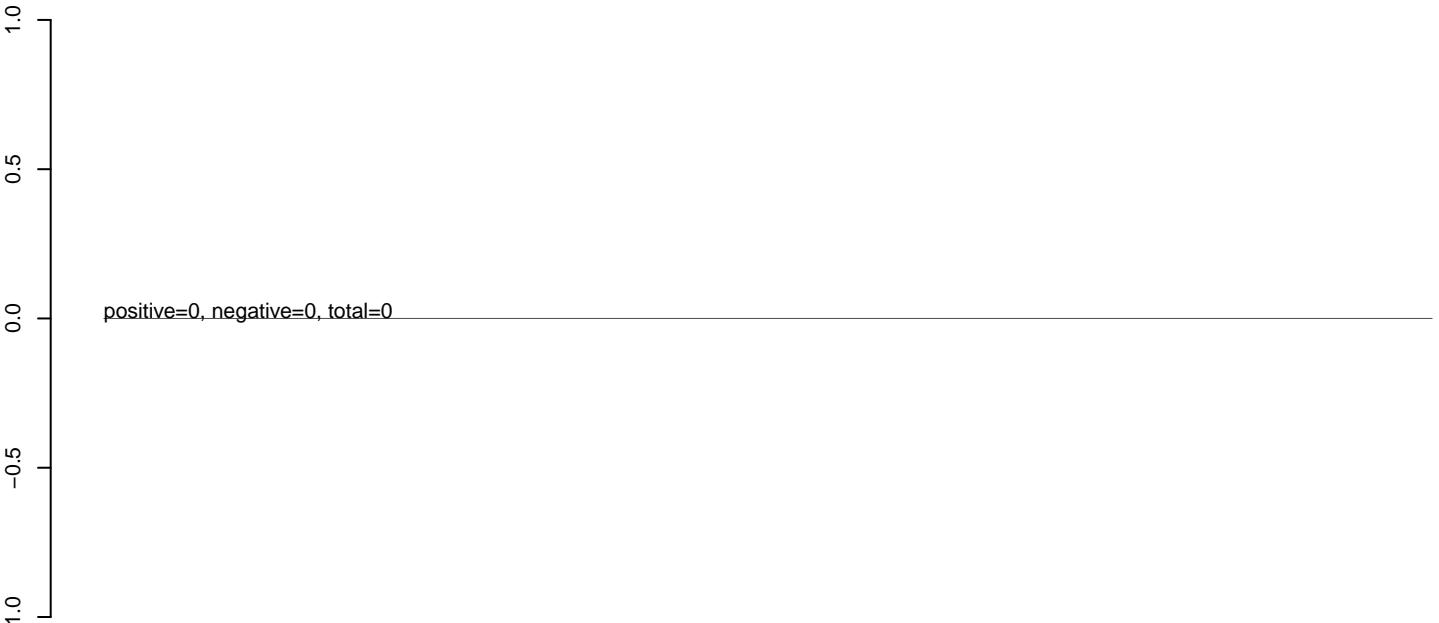
CuQuin_MG_7dpi3.24_35.rep



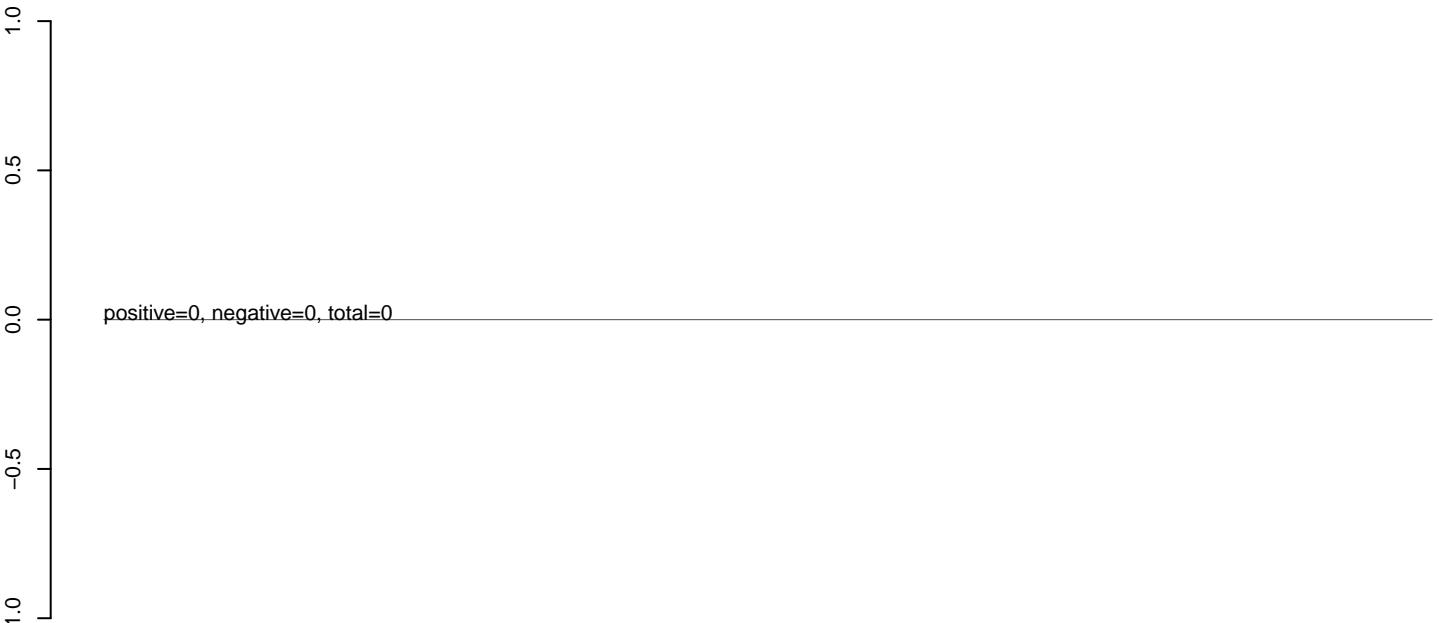
CuQuin_MG_7dpi3.rep



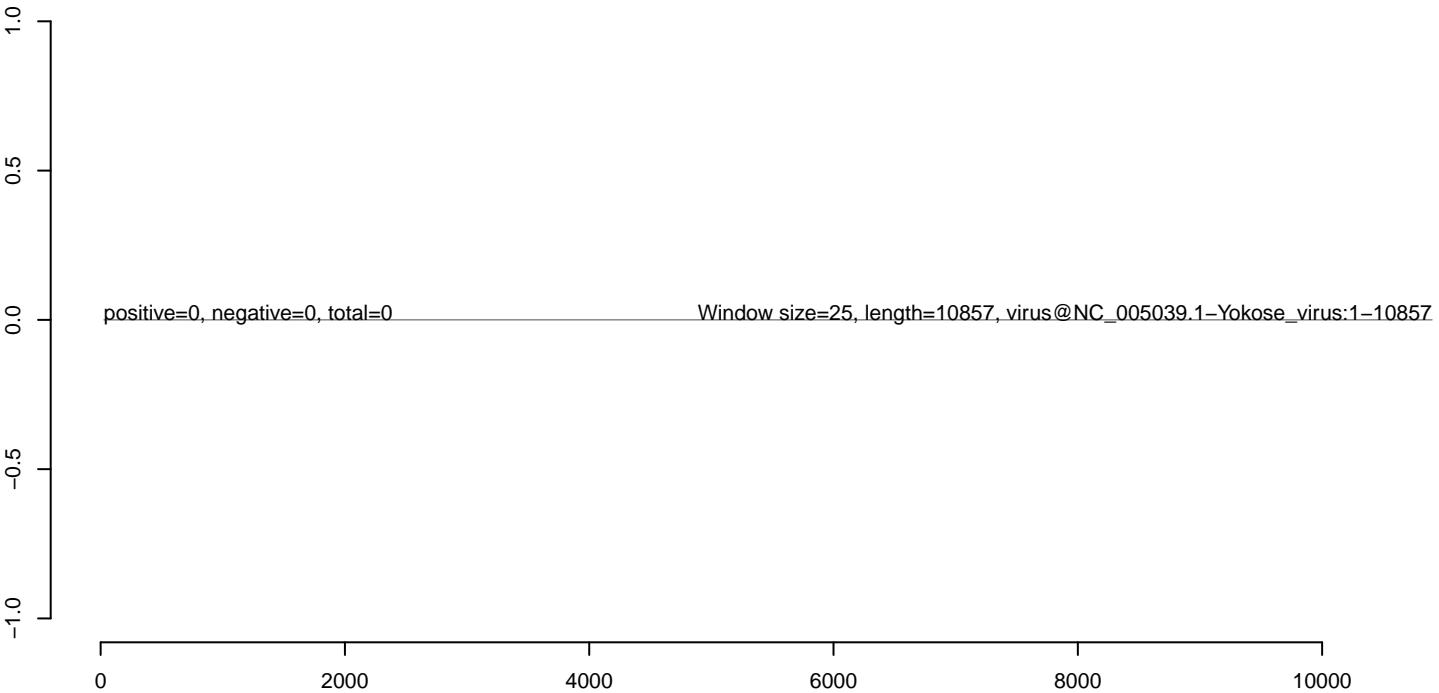
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=10685, virus@NC_004355.1-Alkhurma_virus:1-10685

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

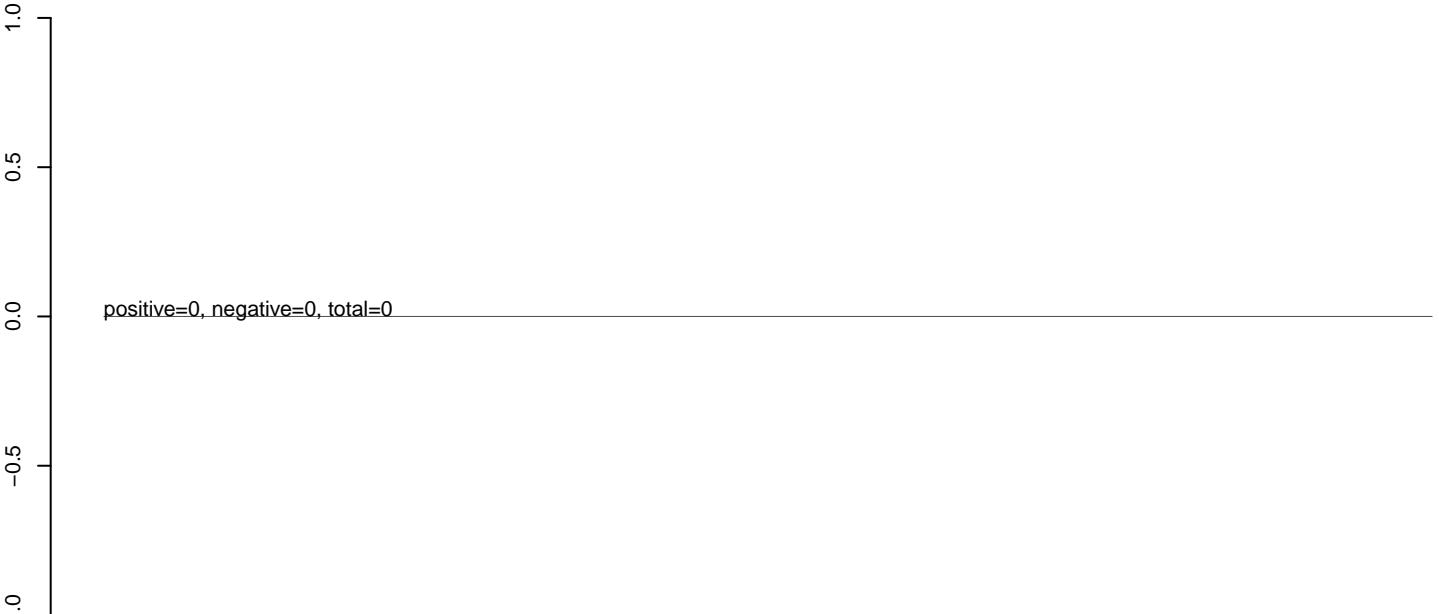
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

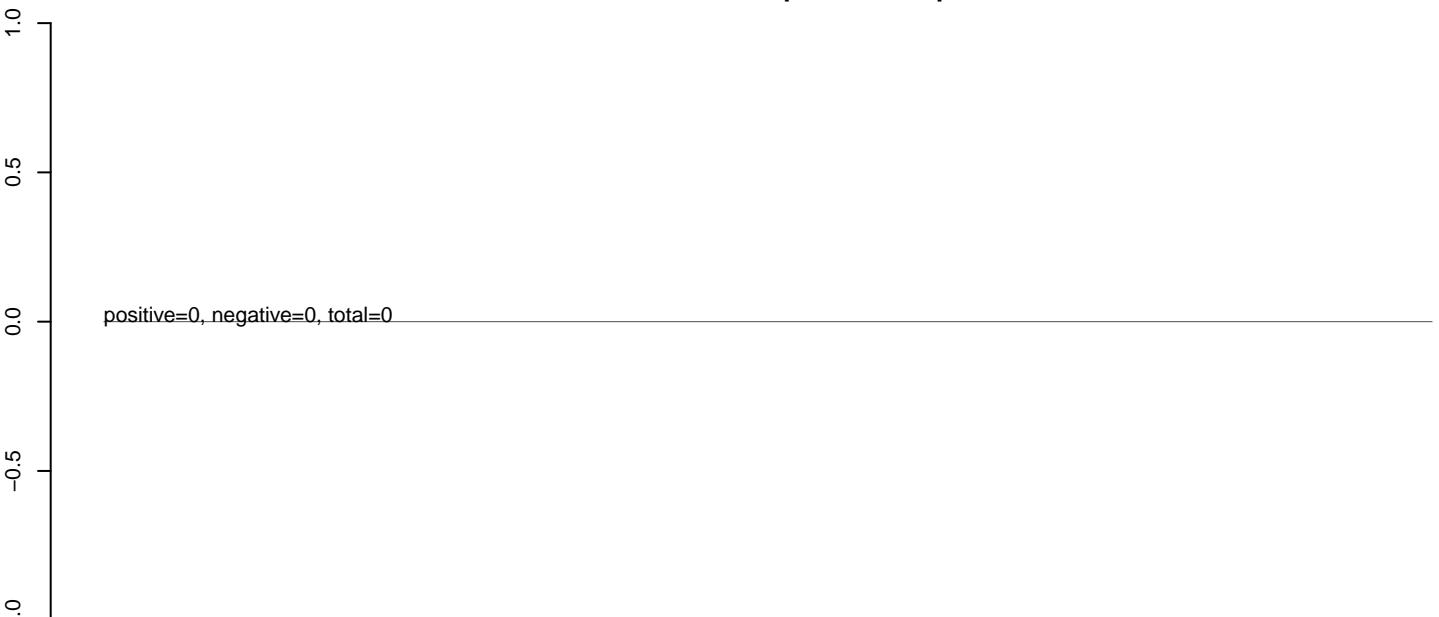
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0/window size=25, length=6603, virus@NC_004177.1-Drosophila_x_virus_segment_A_and_segment_B:1-6603

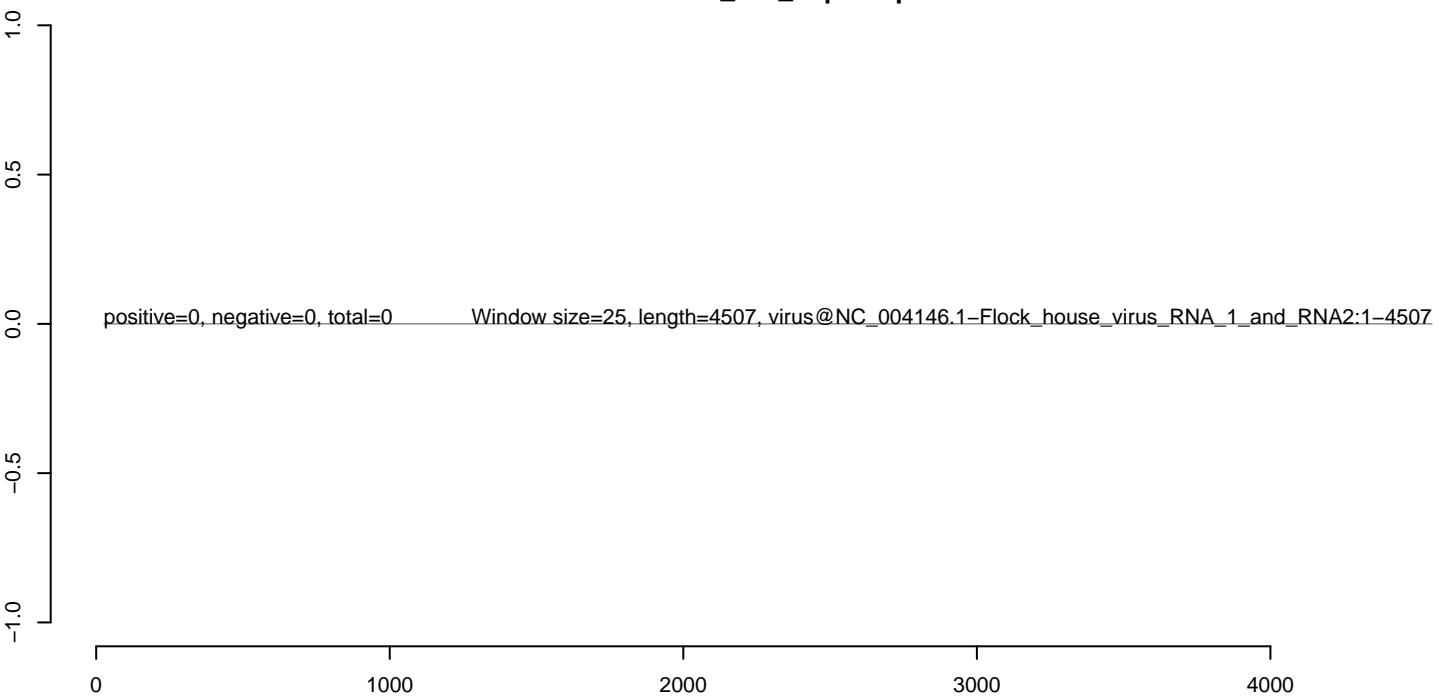
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

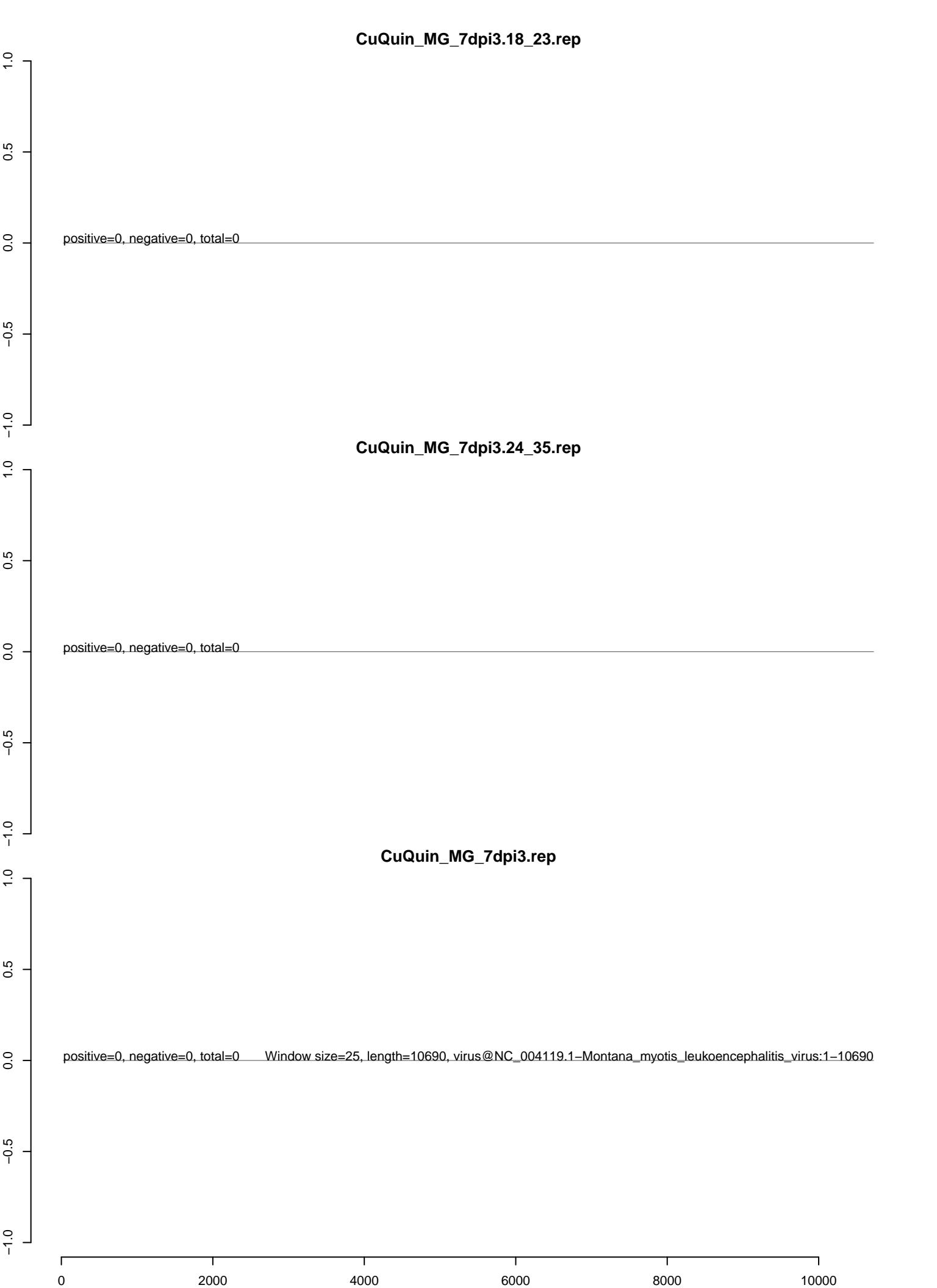
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

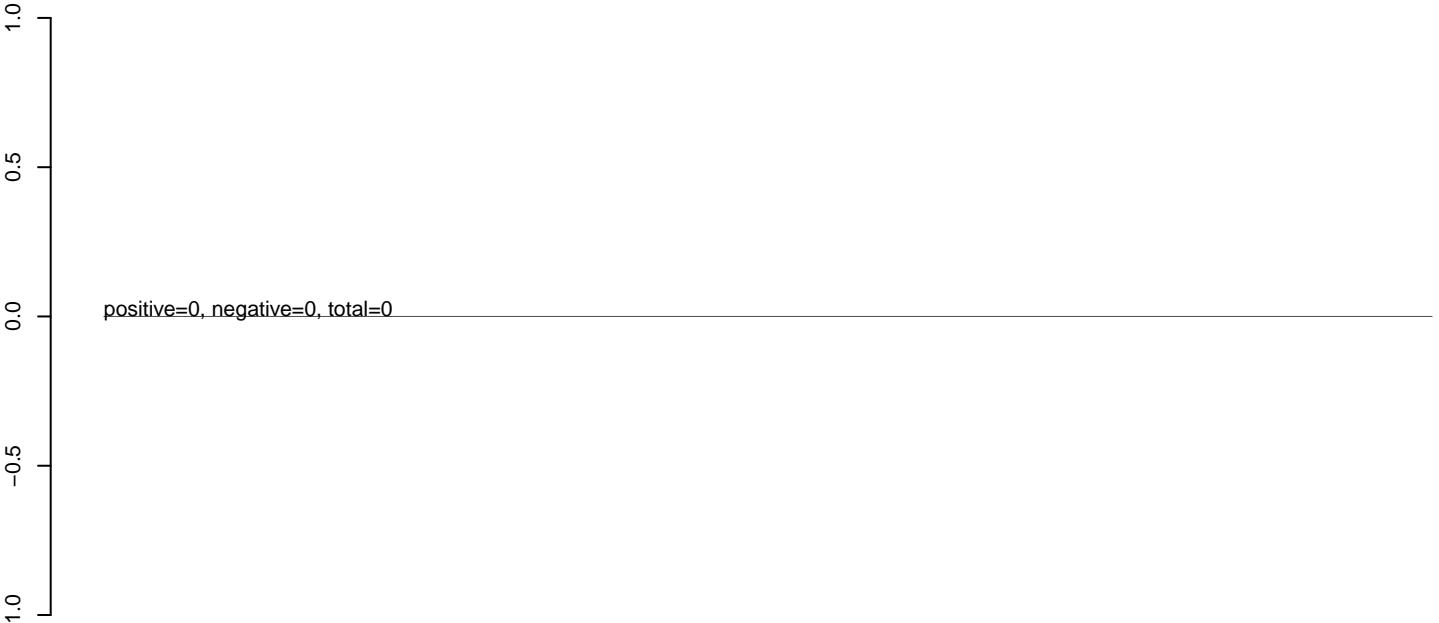
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

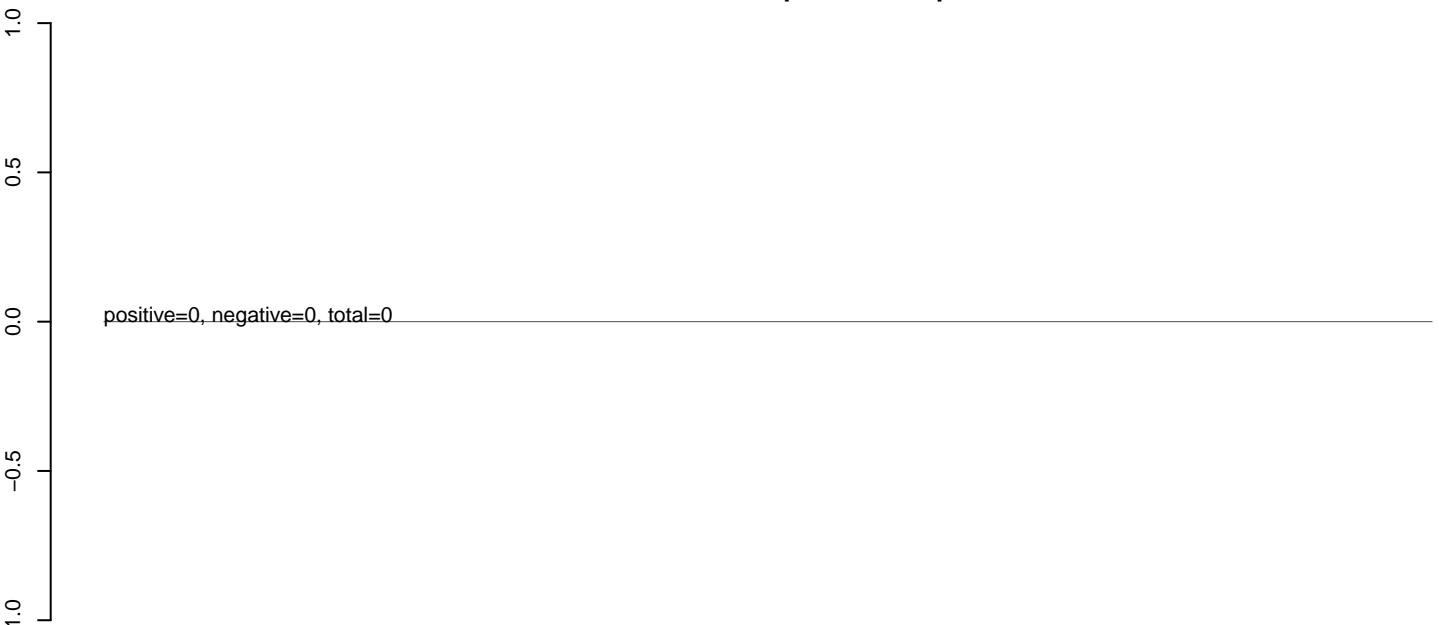
positive=0, negative=0, total=0 Window size=25, length=10690, virus@NC_004119.1-Montana_myotis_leukoencephalitis_virus:1-10690



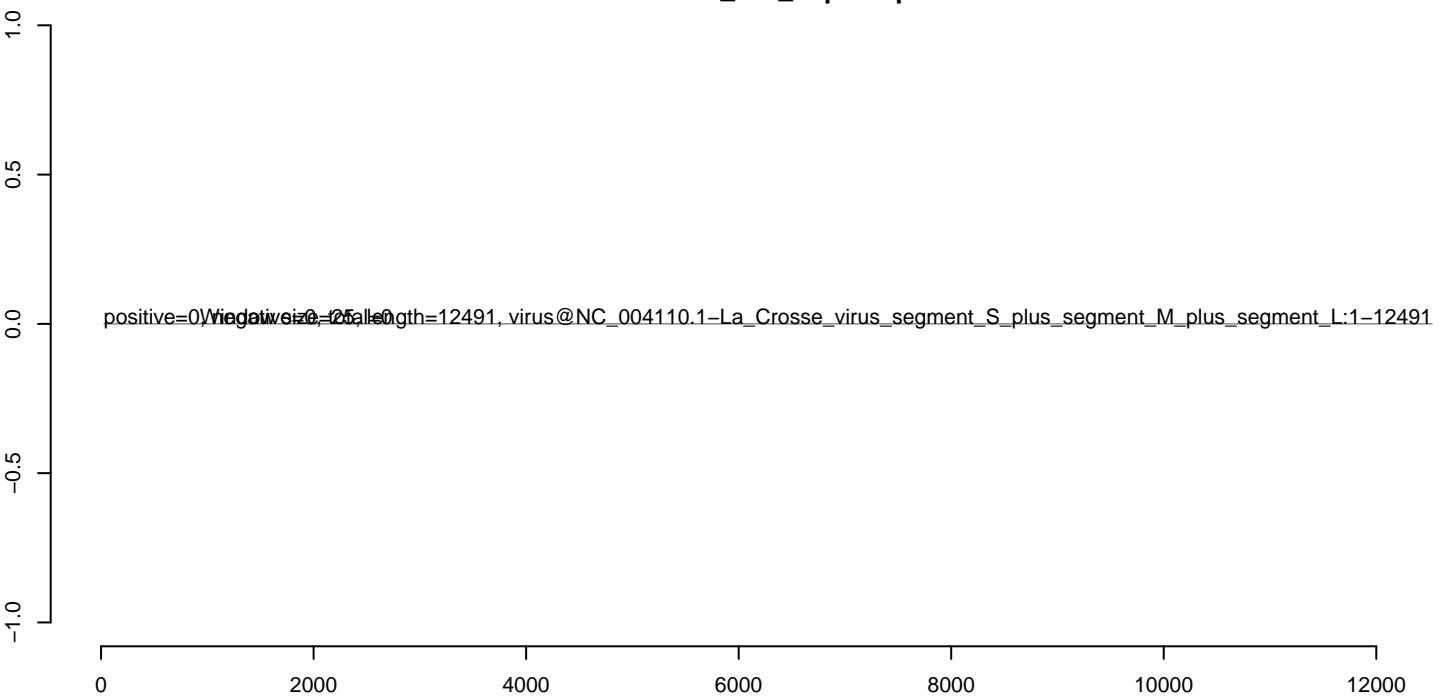
CuQuin_MG_7dpi3.18_23.rep



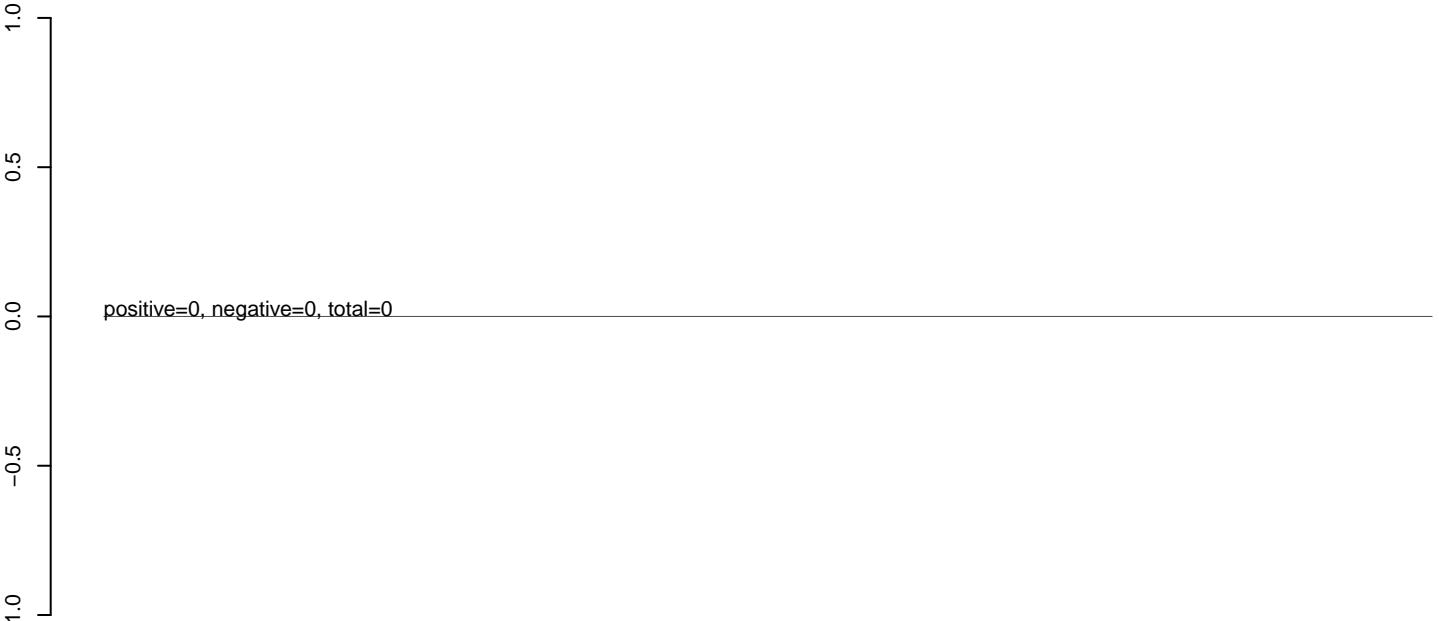
CuQuin_MG_7dpi3.24_35.rep



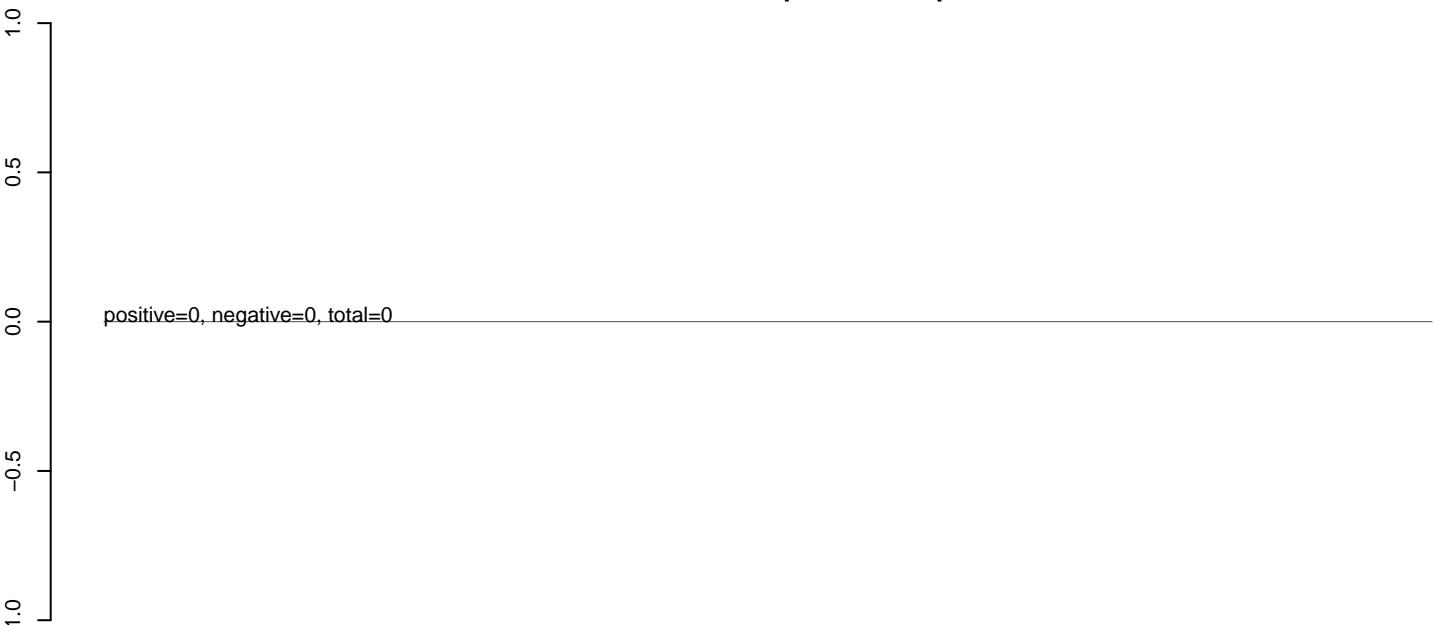
CuQuin_MG_7dpi3.rep



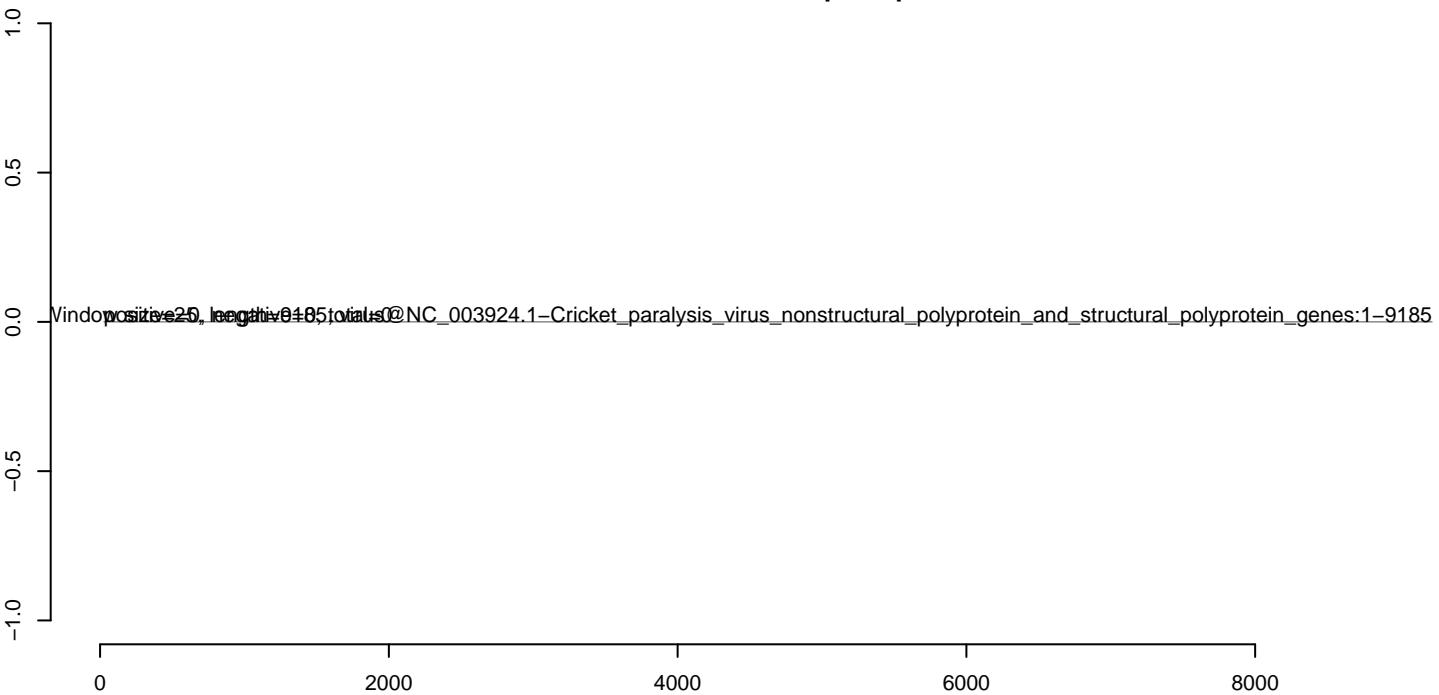
CuQuin_MG_7dpi3.18_23.rep



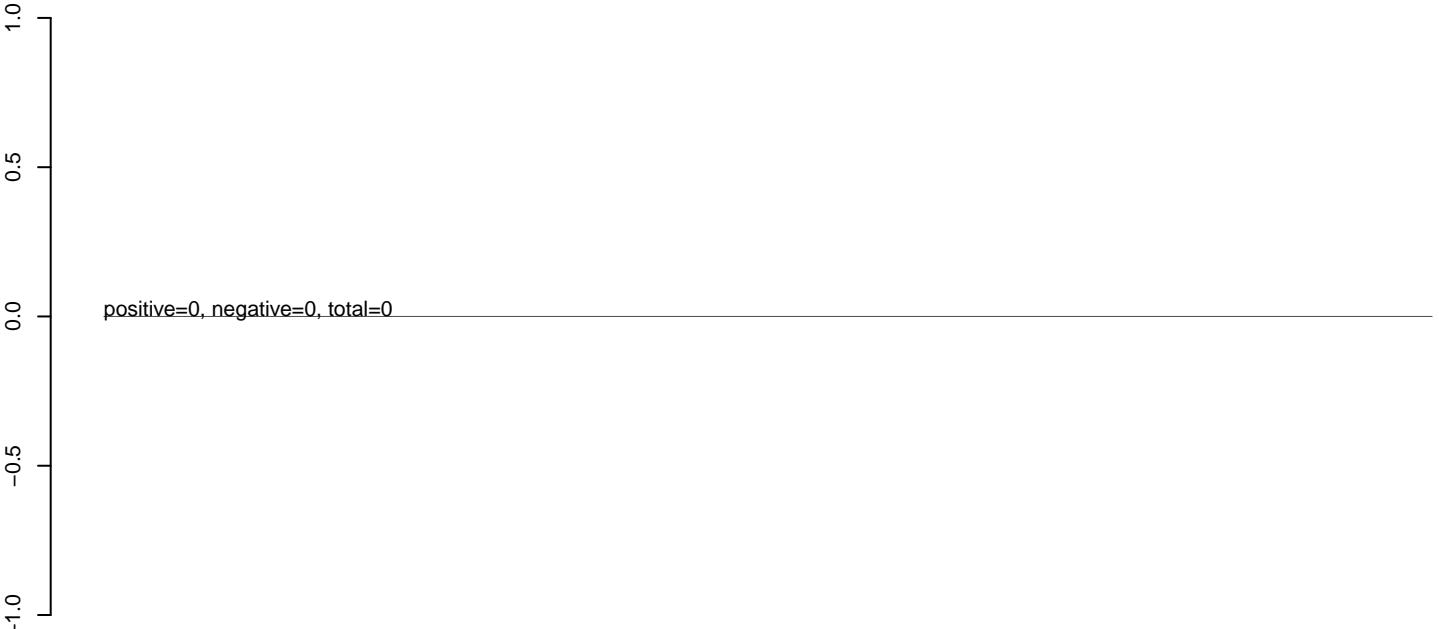
CuQuin_MG_7dpi3.24_35.rep



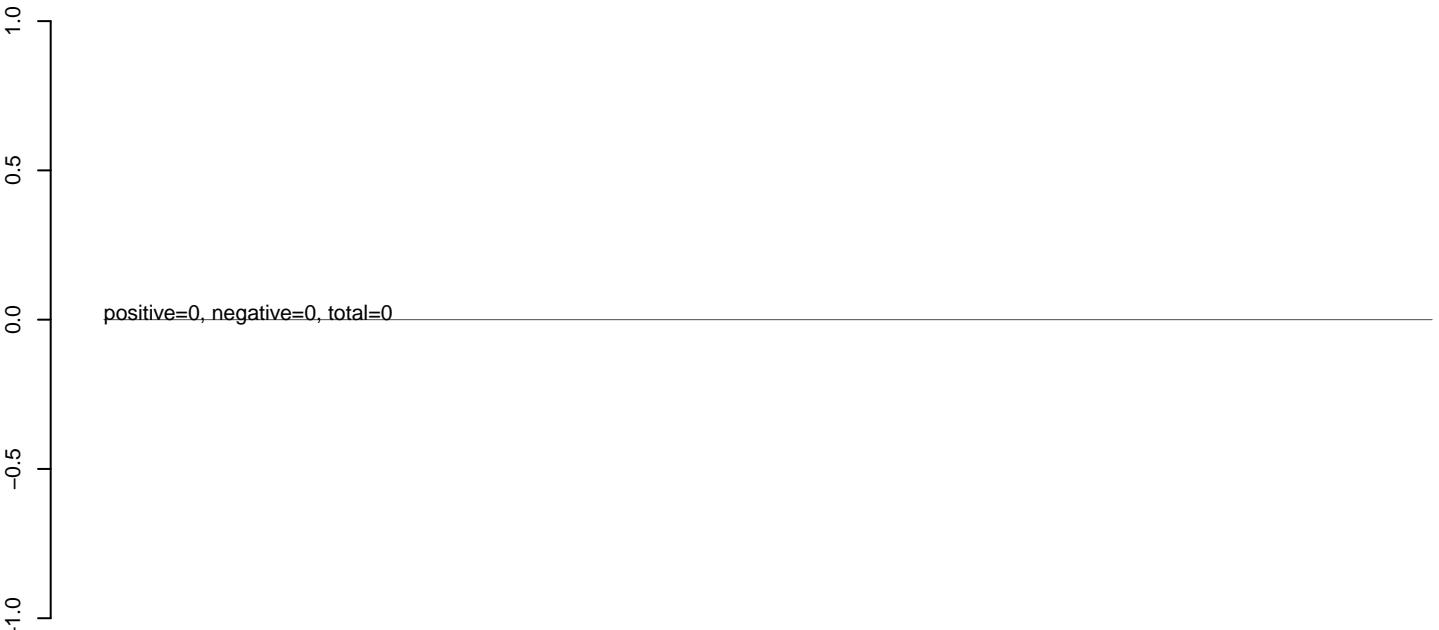
CuQuin_MG_7dpi3.rep



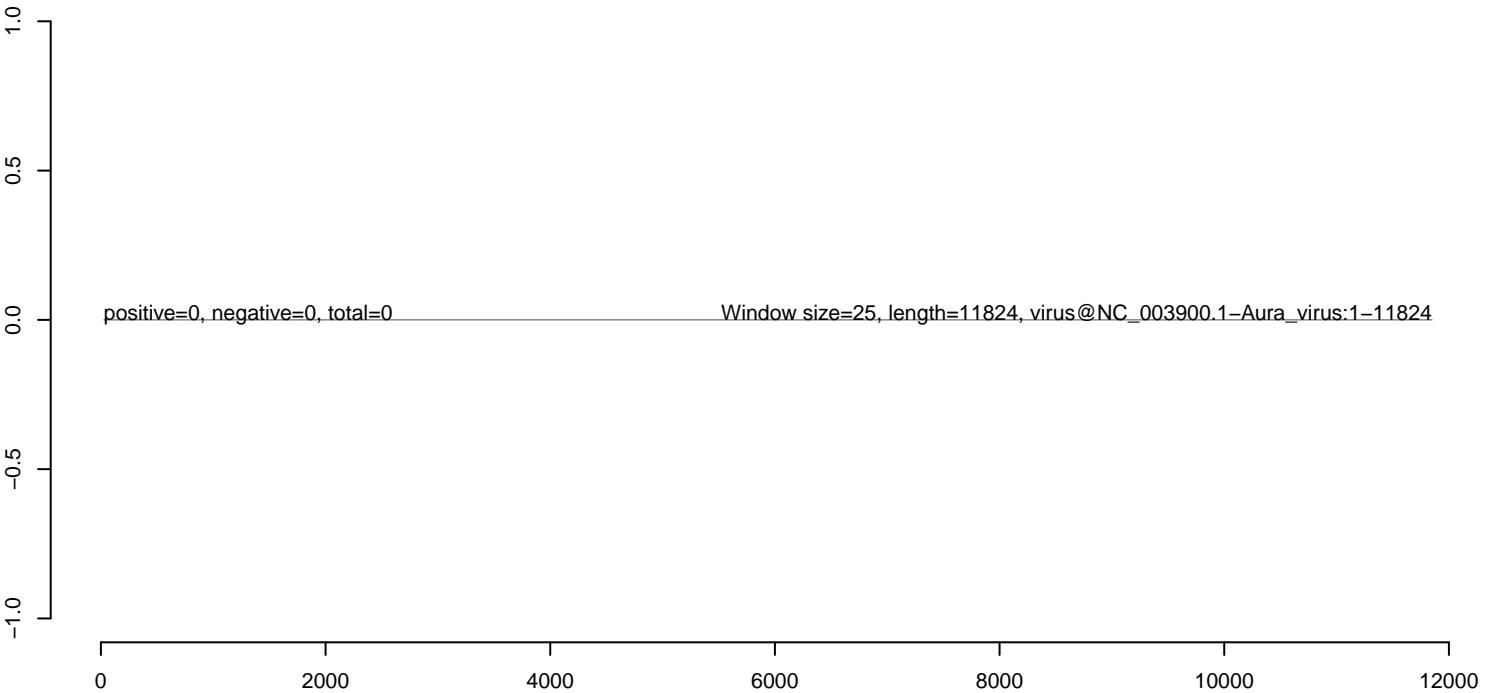
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=11675, virus@NC_003899.1-Eastern_equine_encephalitis_virus:1-11675

0 2000 4000 6000 8000 10000 12000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window.size=25, length=10839, virus@NC_003687.1-Powassan_virus:1-10839

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=10116, virus@NC_003676.1-Apoi_virus_genome:1-10116

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

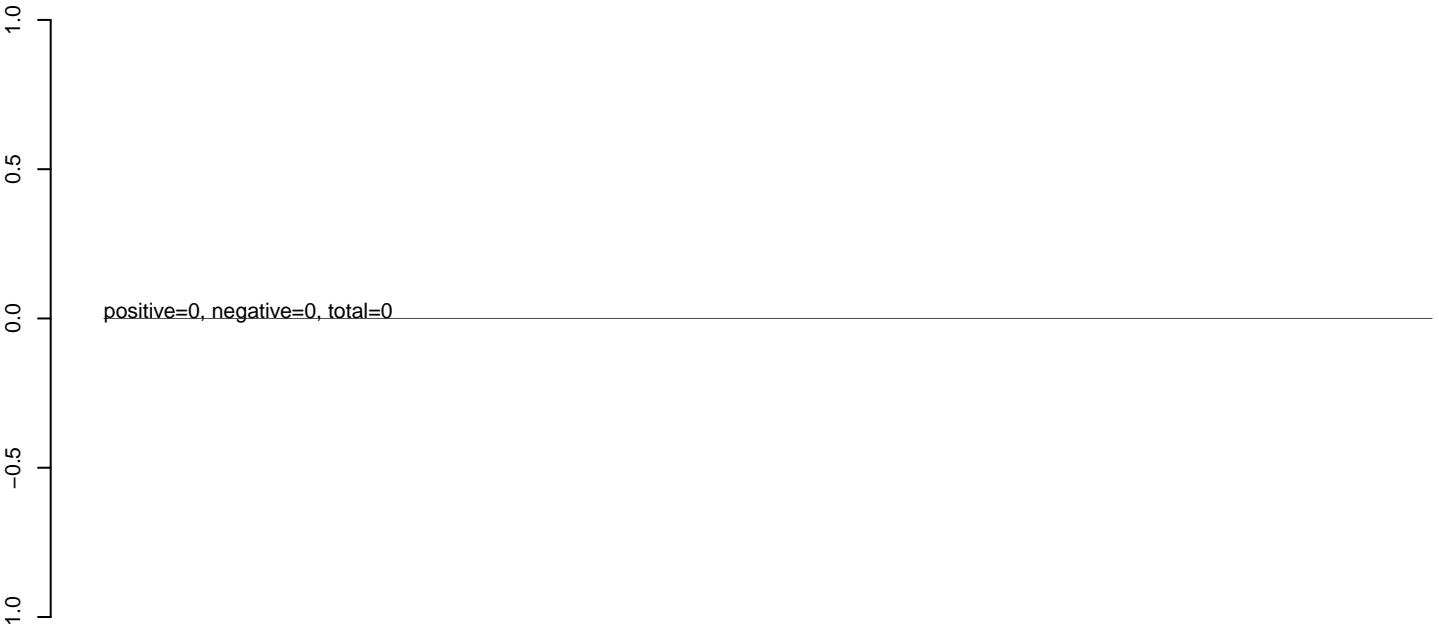
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

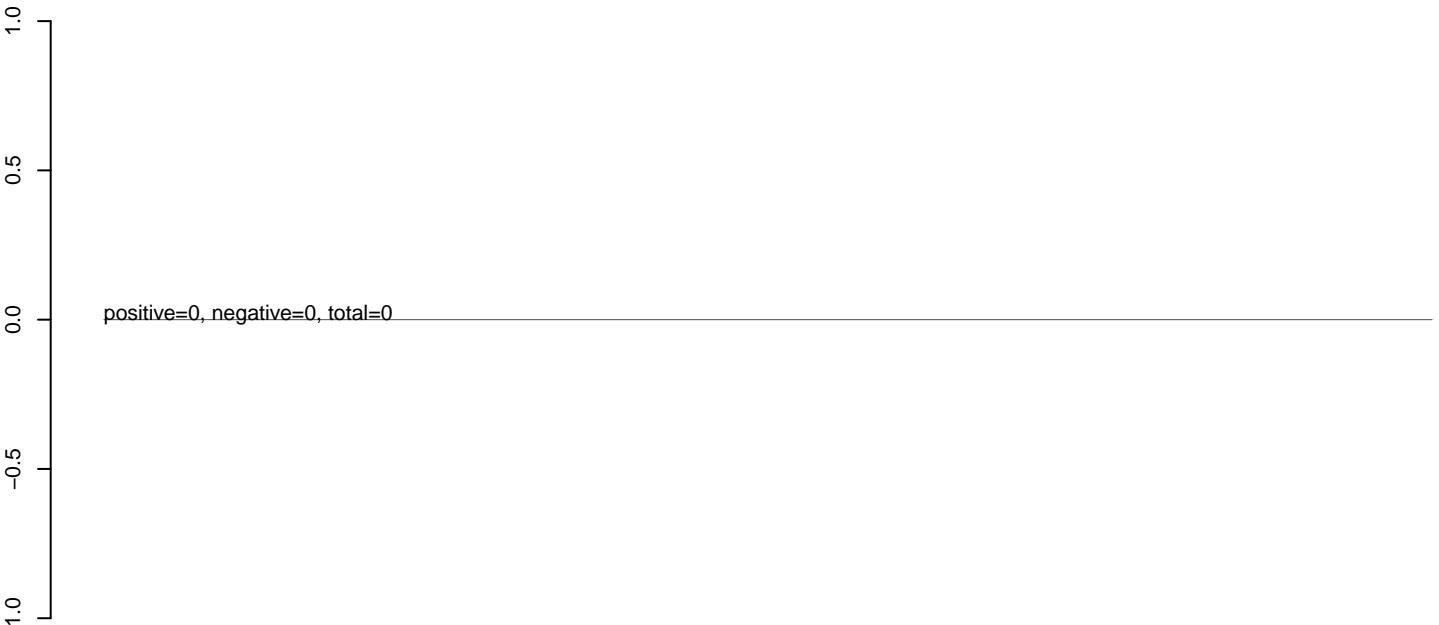
positive=0, negative=0, total=0

Window size=25, length=10140, virus@NC_003675.1–Rio_Bravo_virus_genome:1–10140

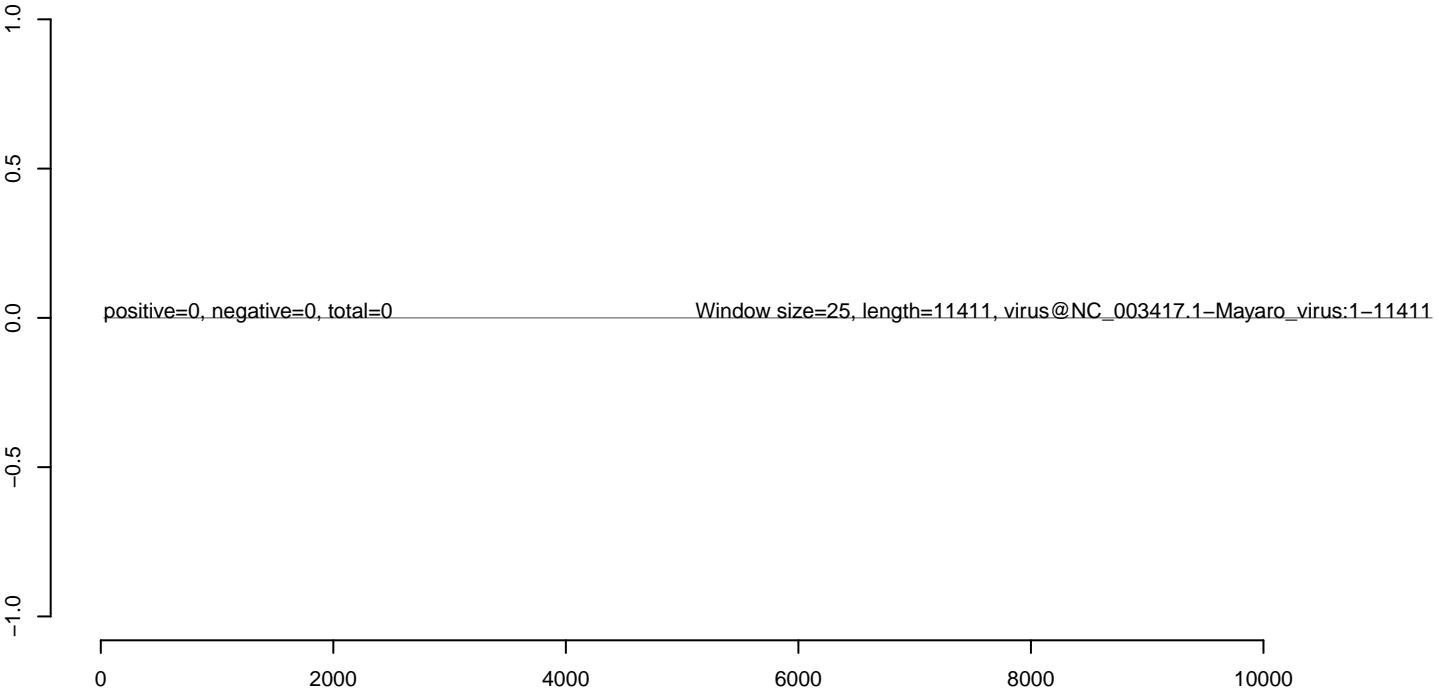
CuQuin_MG_7dpi3.18_23.rep



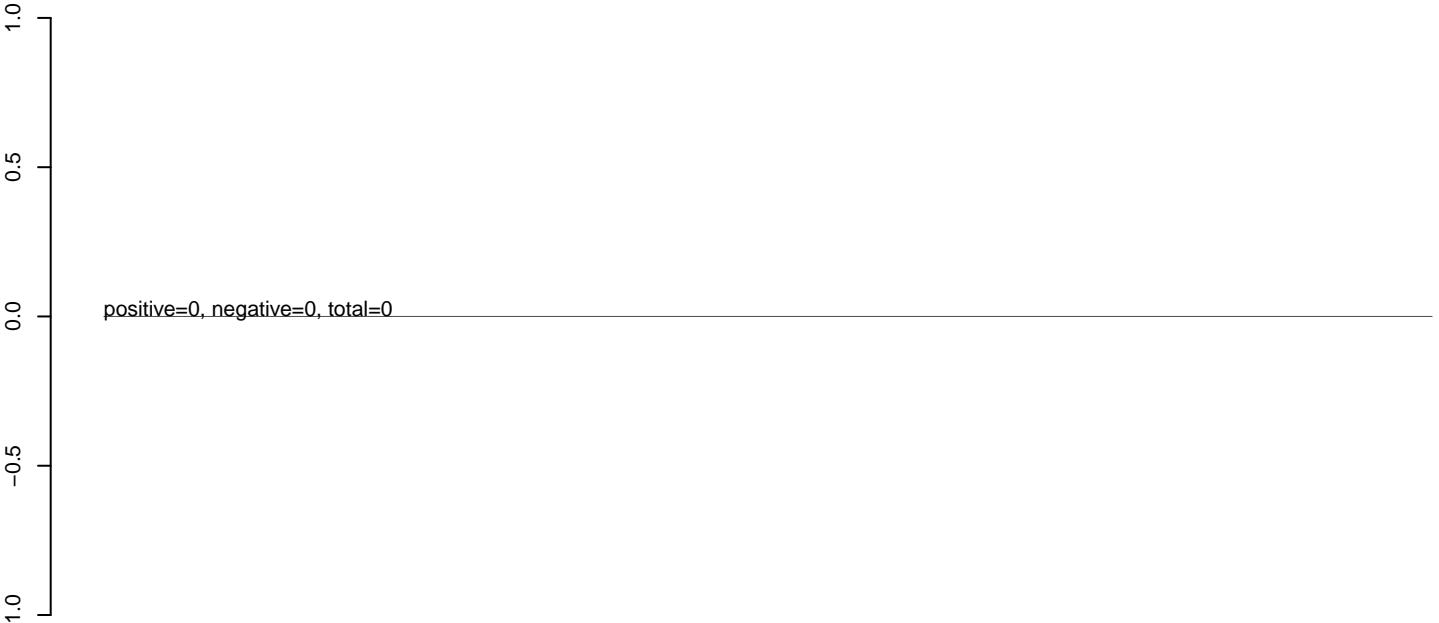
CuQuin_MG_7dpi3.24_35.rep



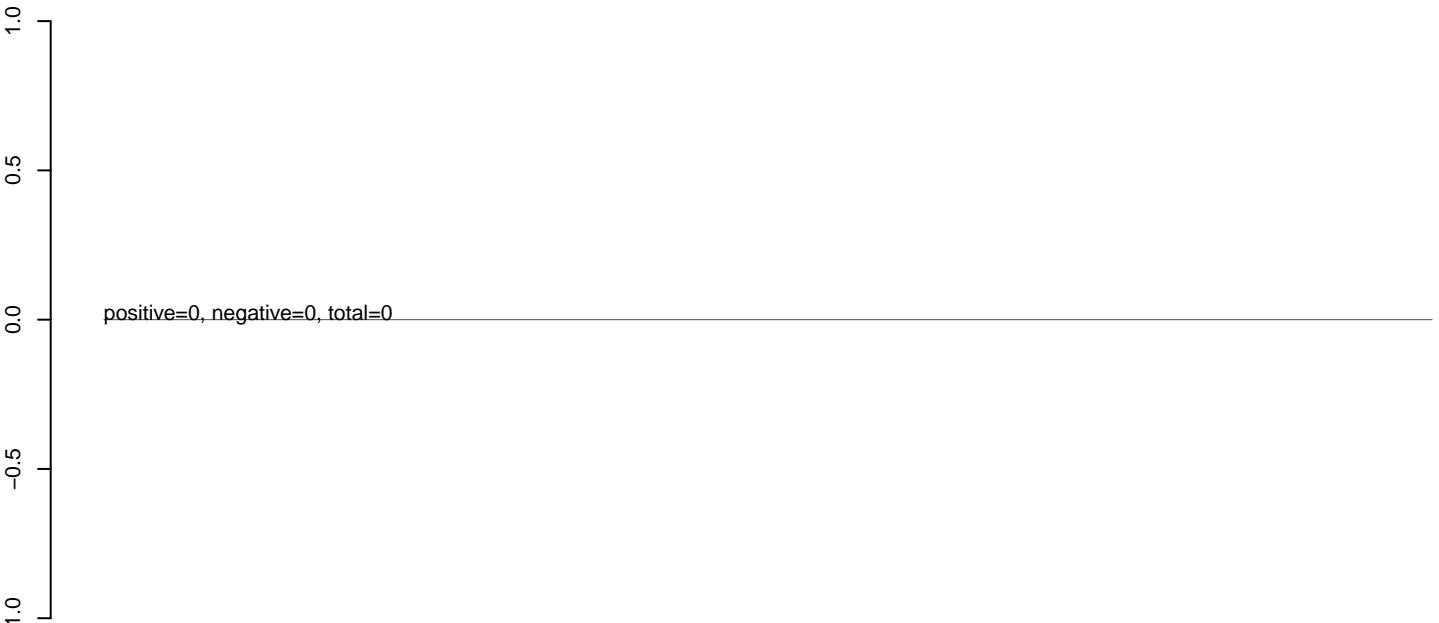
CuQuin_MG_7dpi3.rep



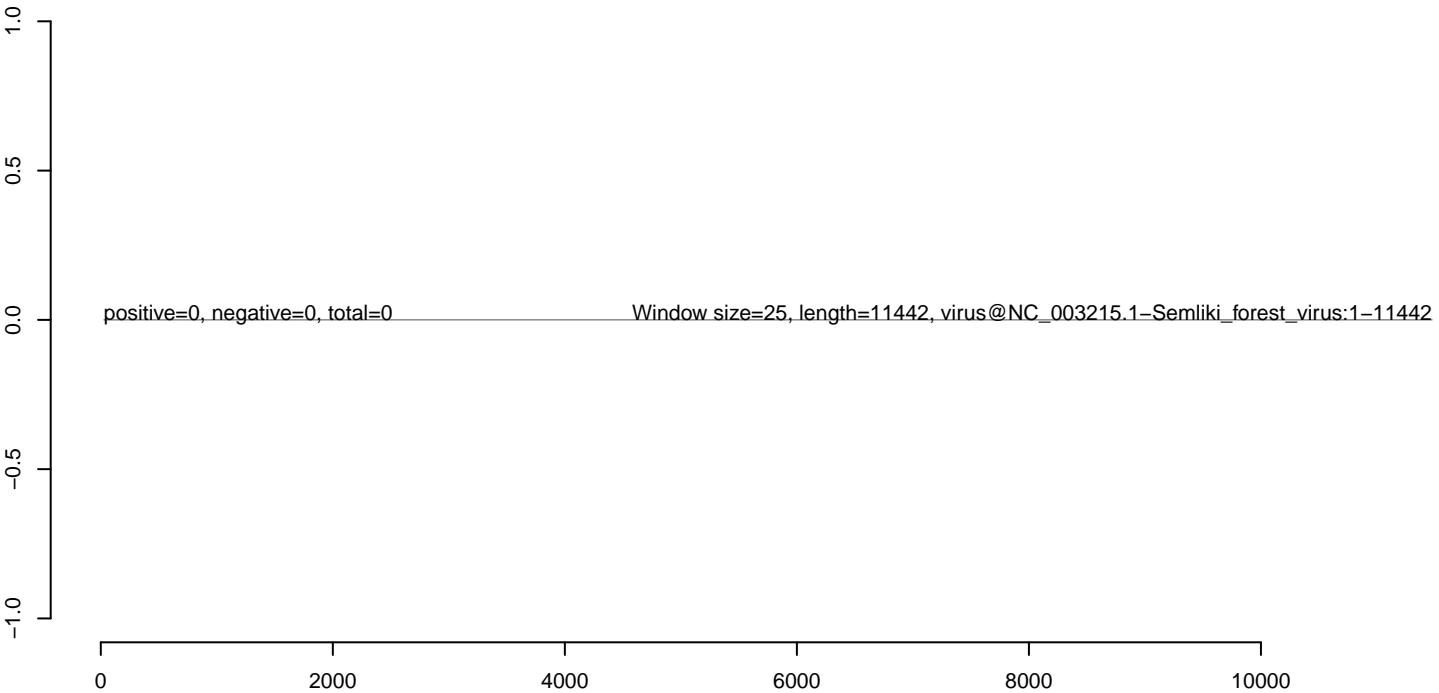
CuQuin_MG_7dpi3.18_23.rep



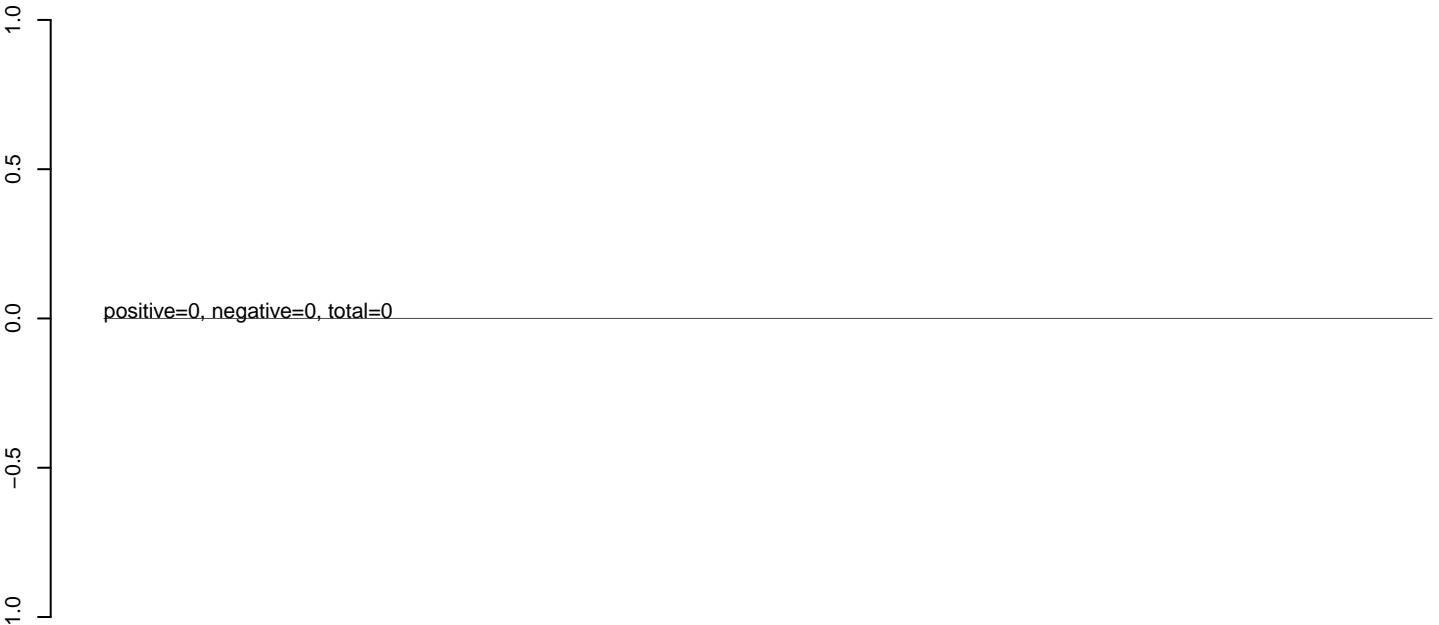
CuQuin_MG_7dpi3.24_35.rep



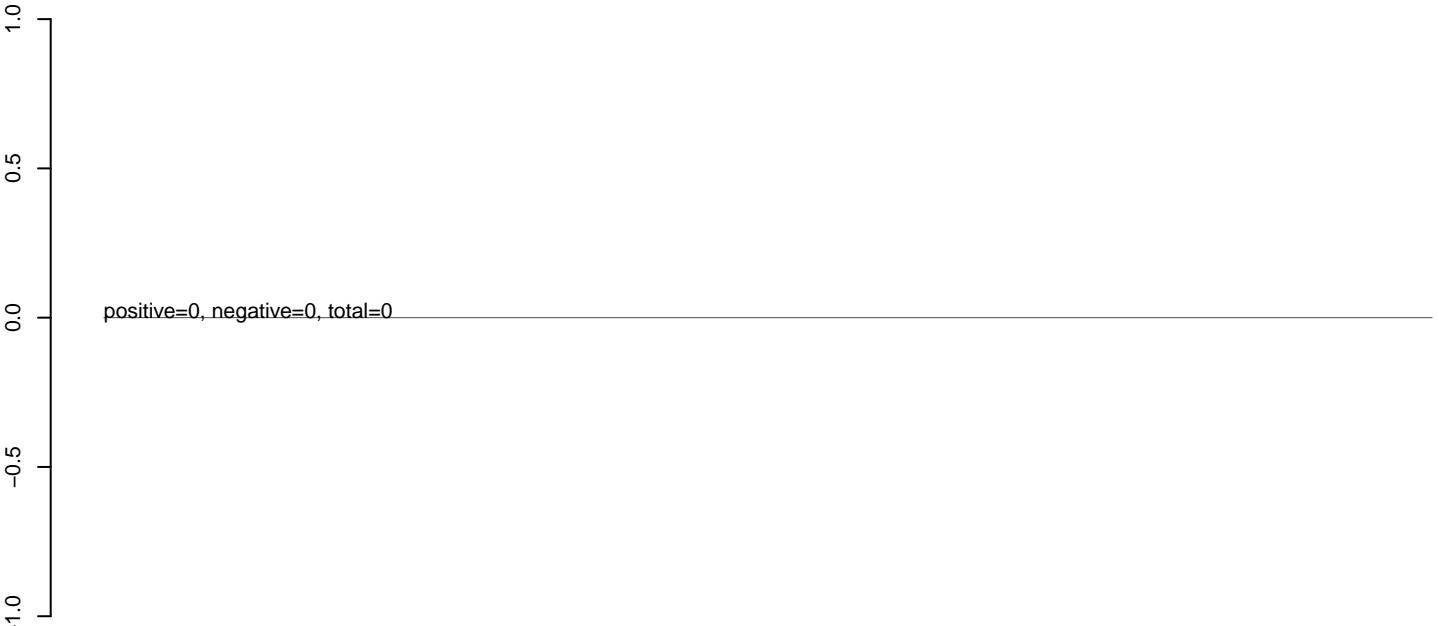
CuQuin_MG_7dpi3.rep



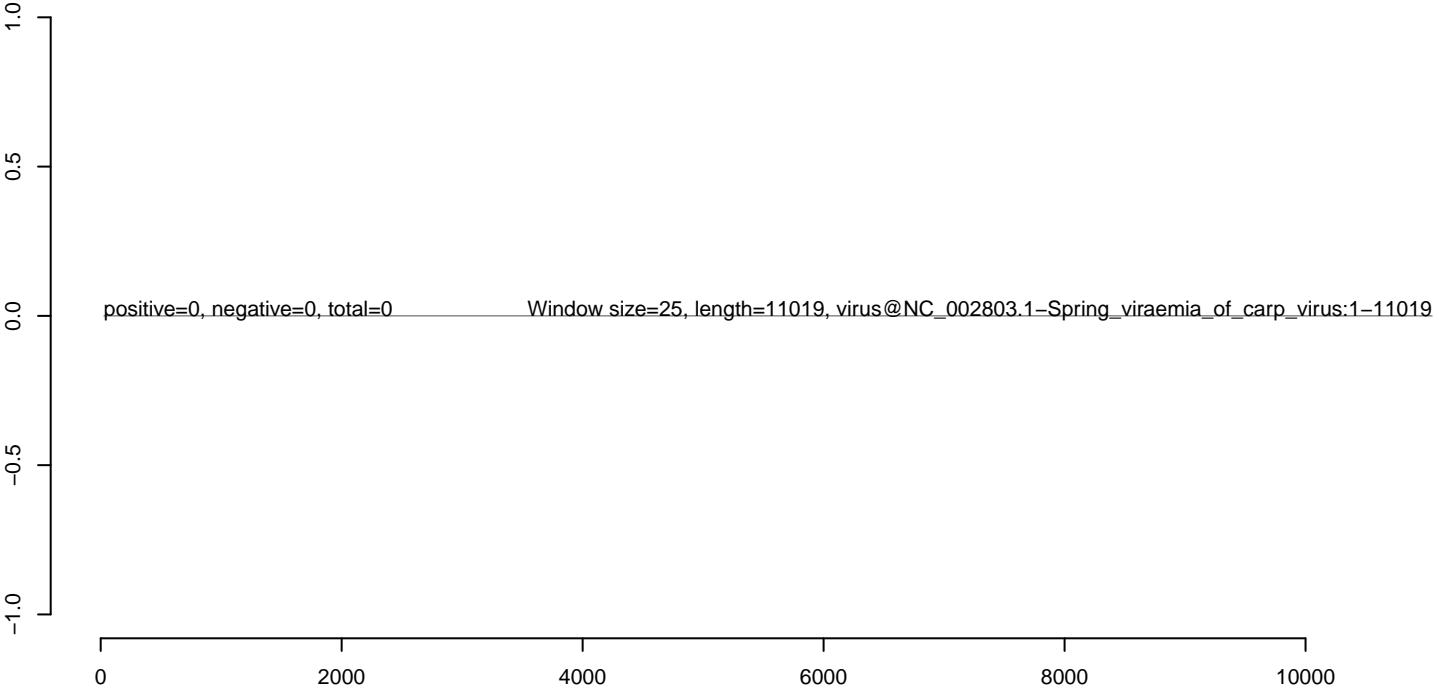
CuQuin_MG_7dpi3.18_23.rep



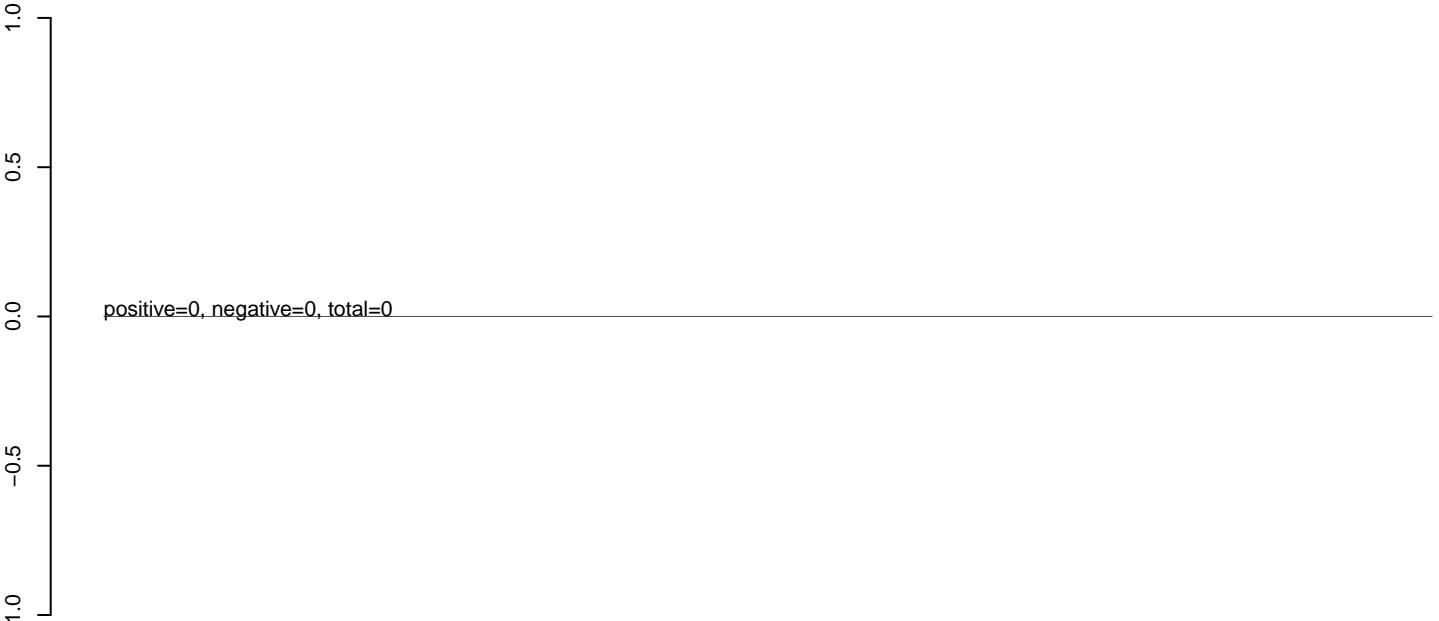
CuQuin_MG_7dpi3.24_35.rep



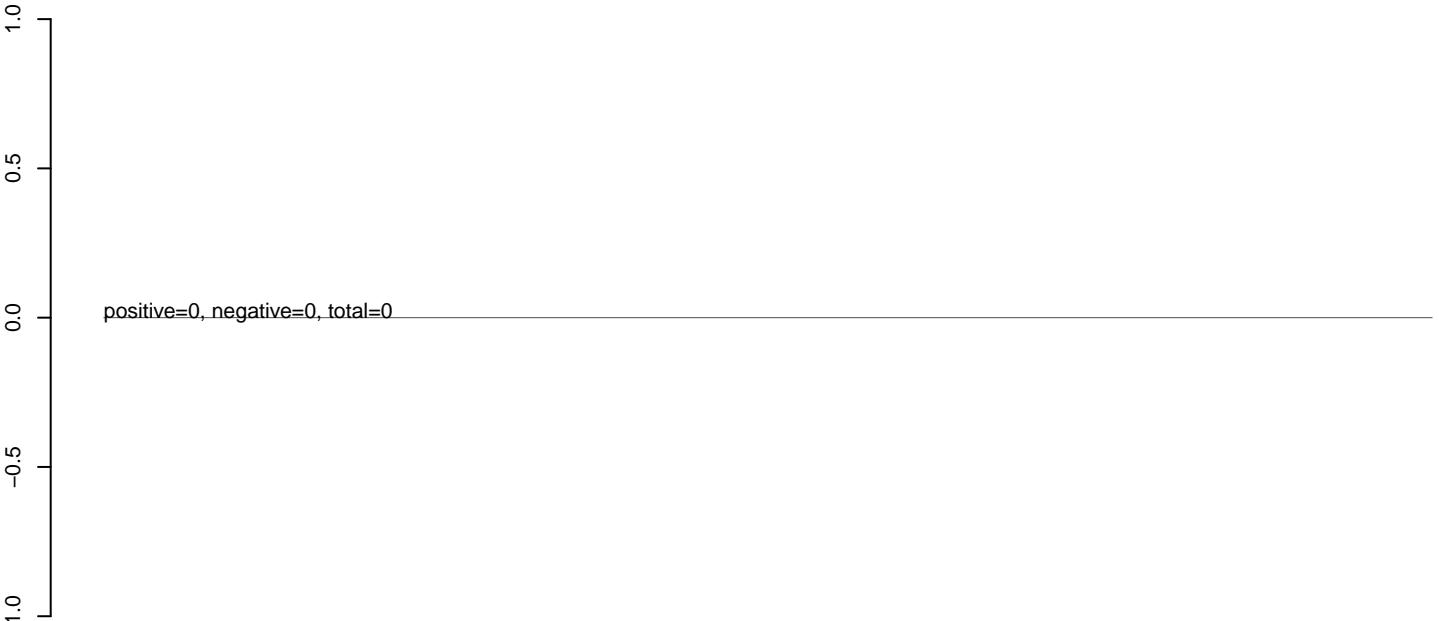
CuQuin_MG_7dpi3.rep



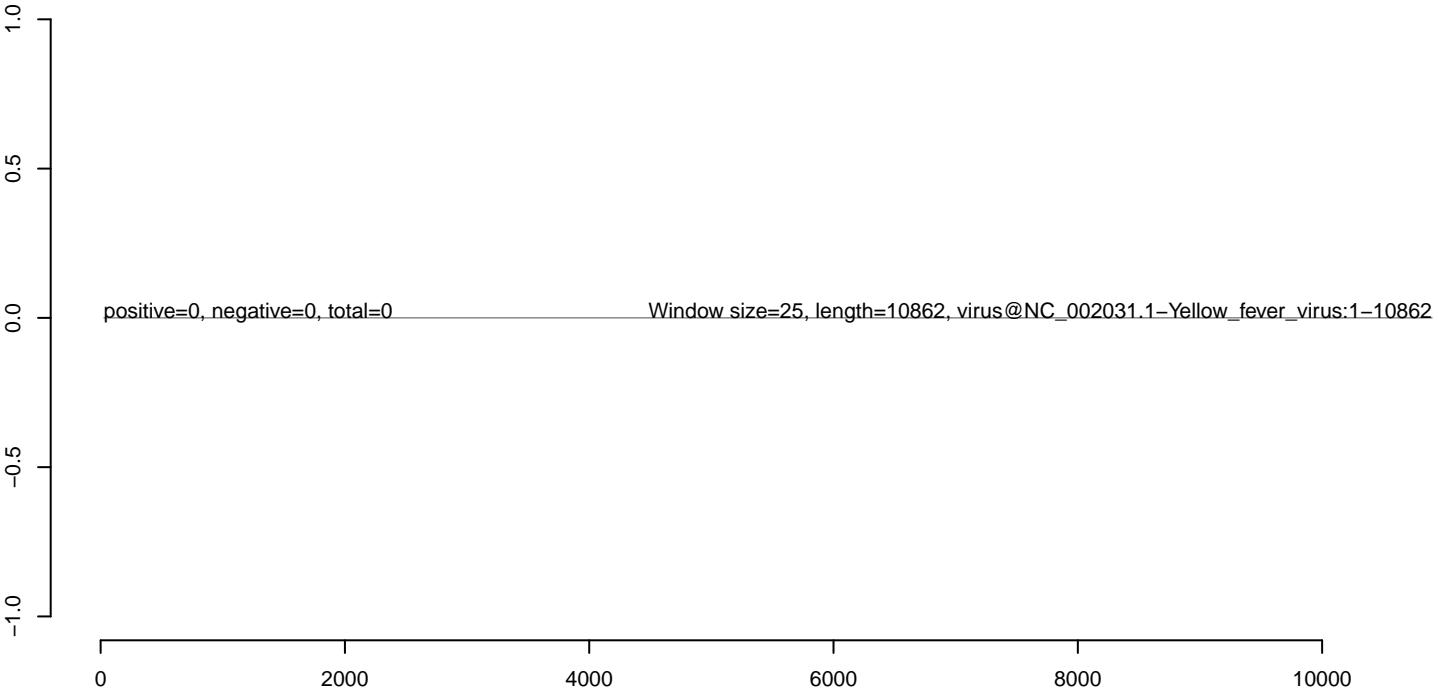
CuQuin_MG_7dpi3.18_23.rep

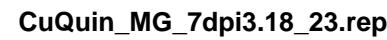


CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep





positive=0, negative=0, total=0

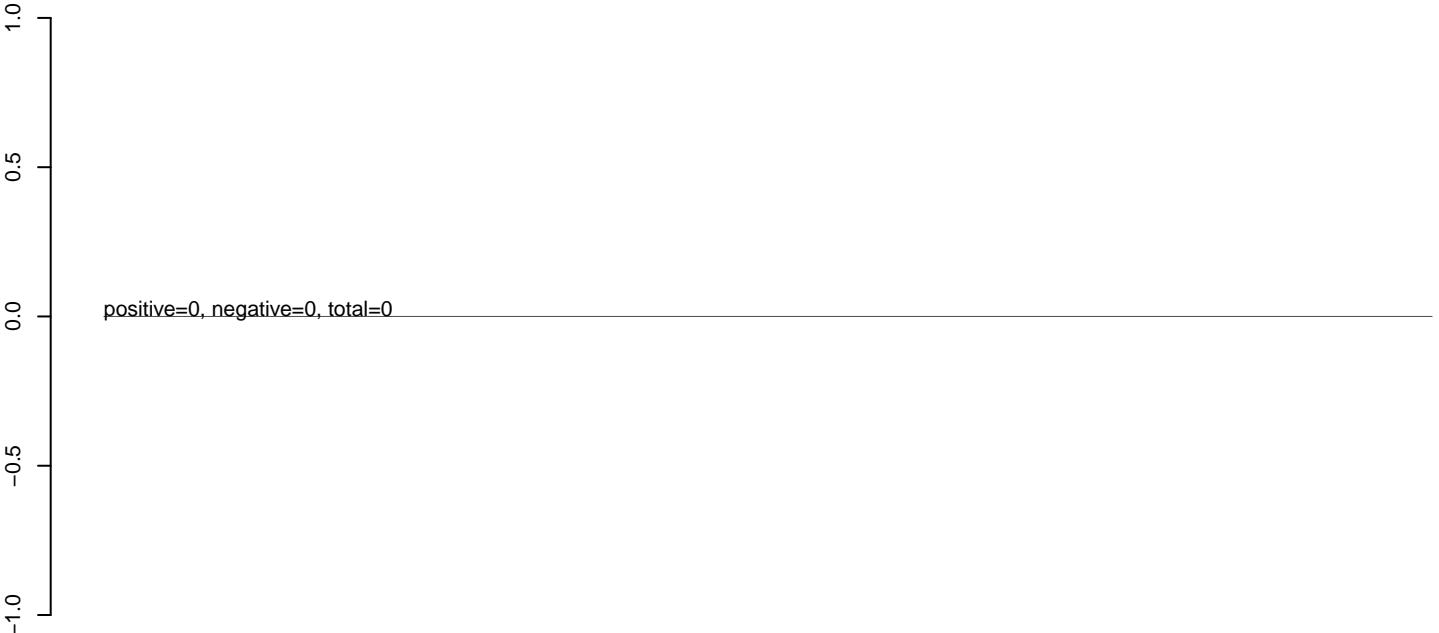


positive=0, negative=0, total=0

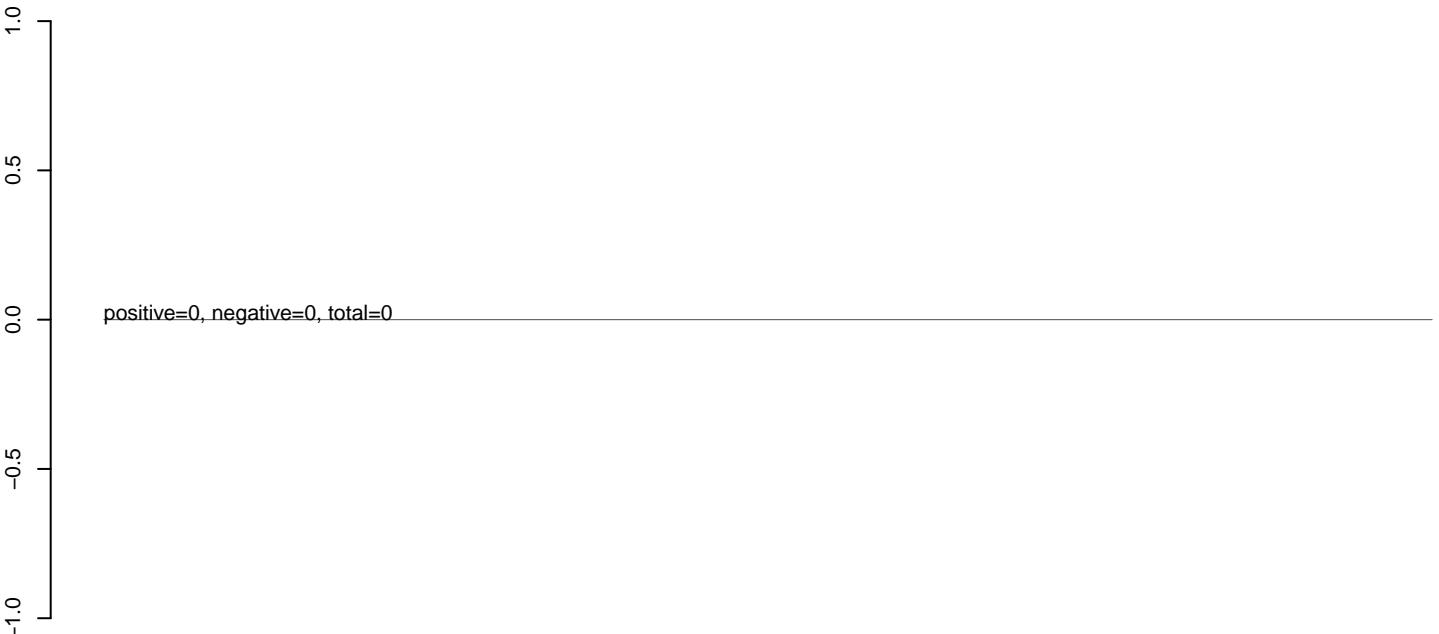


positive=Virus_isolate_025 total=0 h=12294 virus@NC_001927.1-Bunyamwera_virus_segment_S_plus_segment_M_plus_segment_L:1-12294

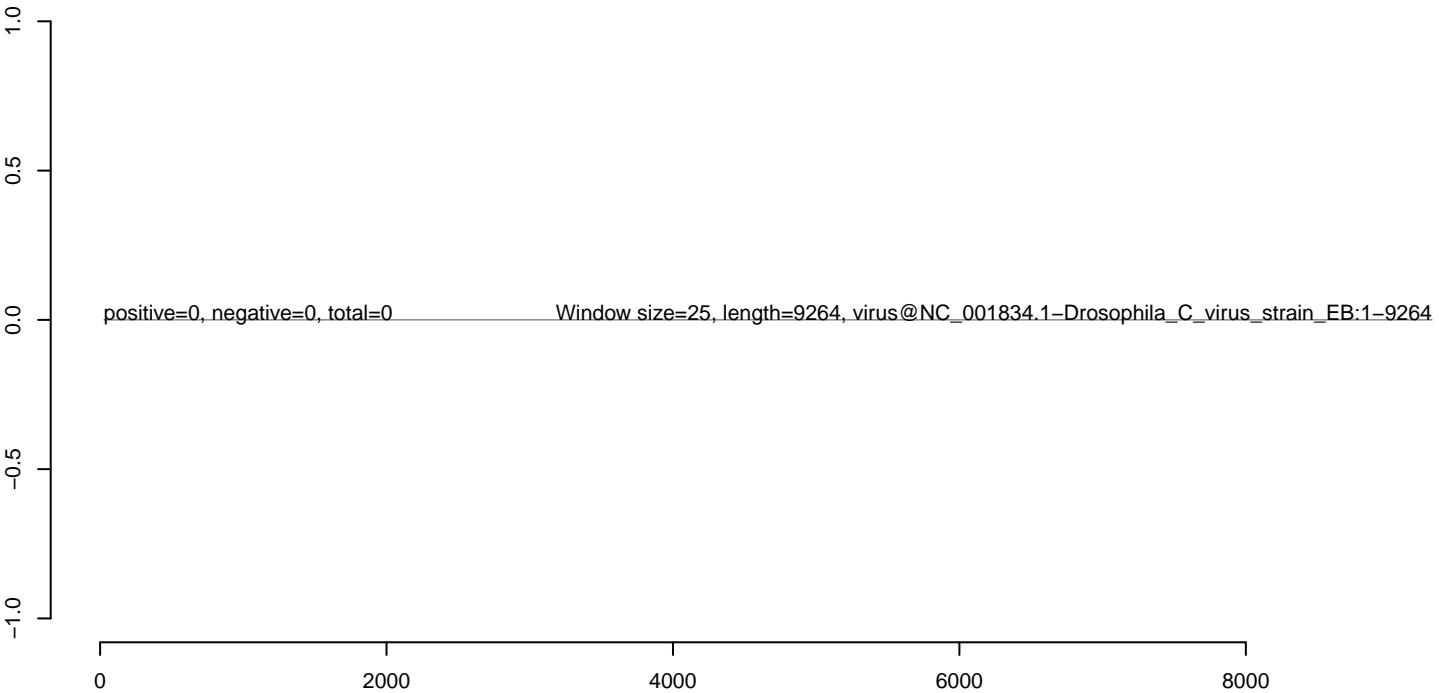
CuQuin_MG_7dpi3.18_23.rep



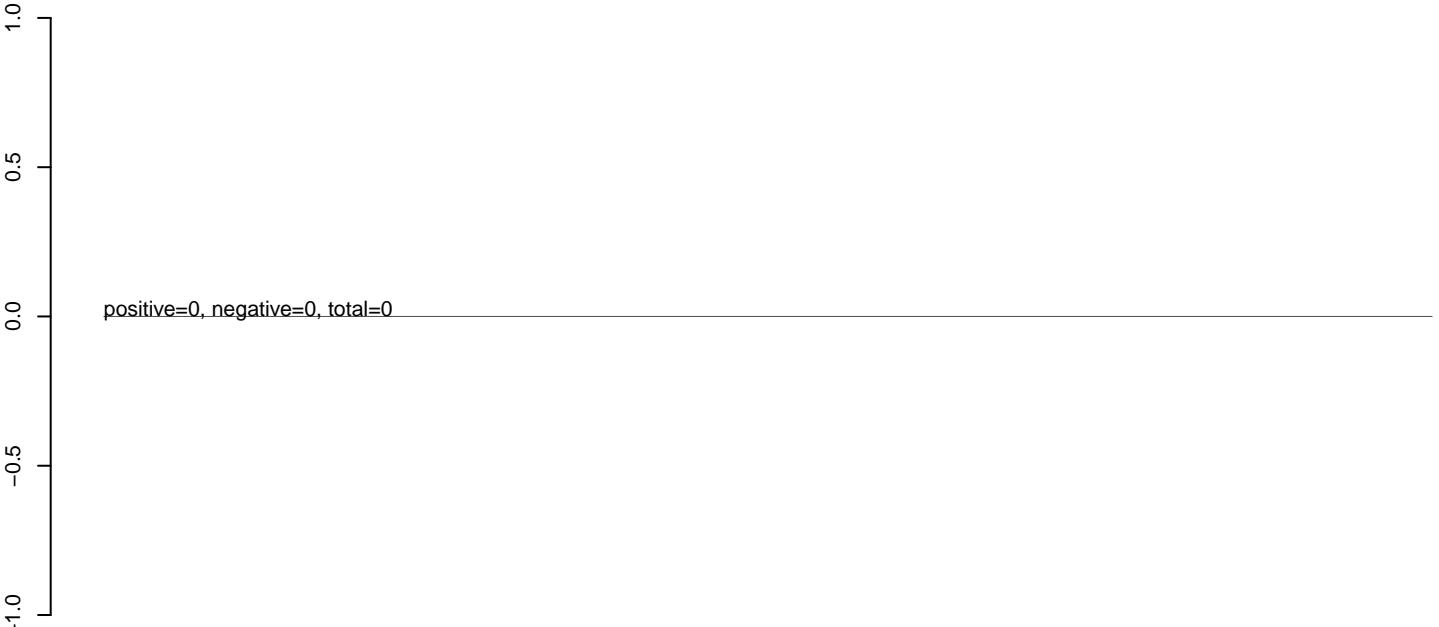
CuQuin_MG_7dpi3.24_35.rep



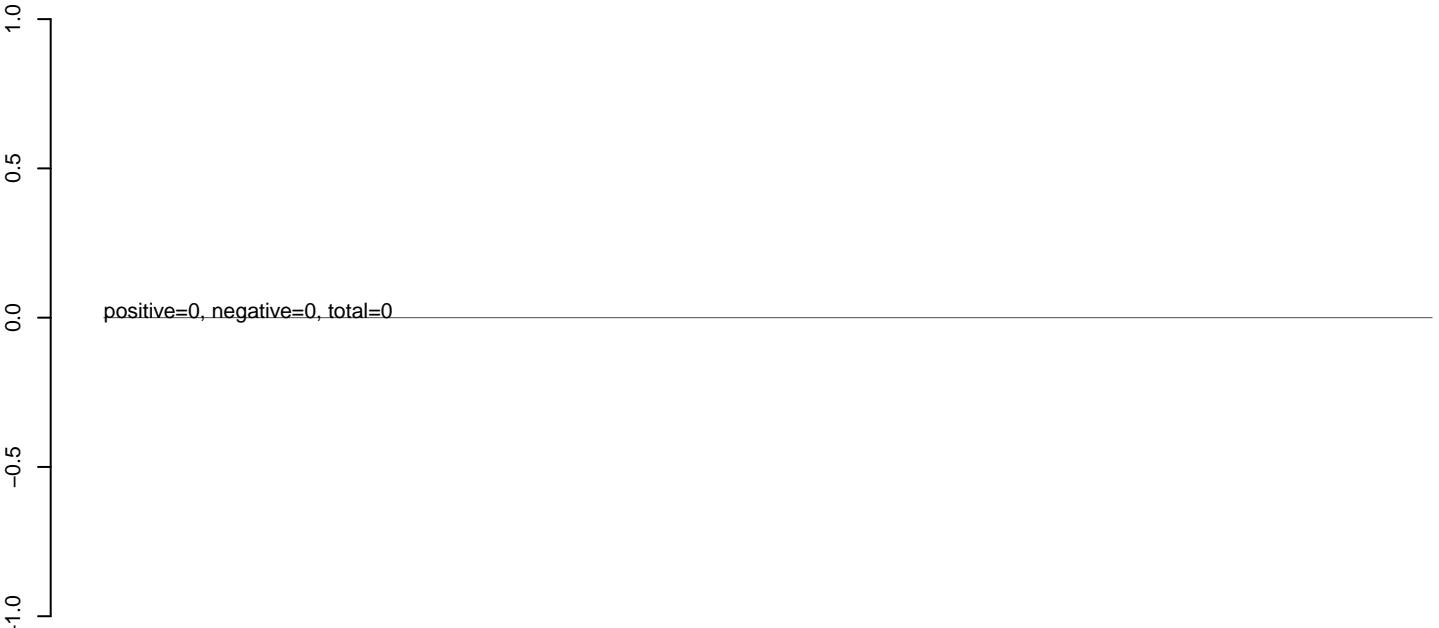
CuQuin_MG_7dpi3.rep



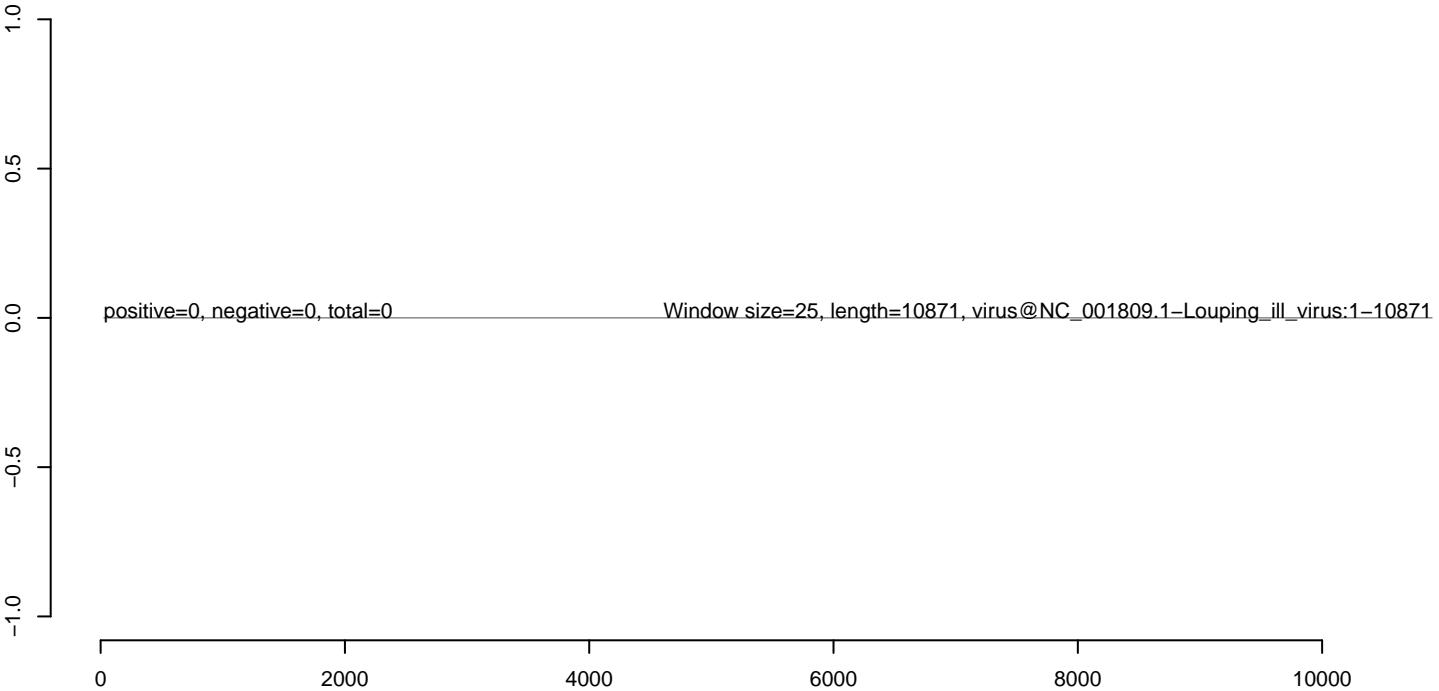
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=11488, virus@NC_001786.1–Barmah_Forest_virus:1–11488

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

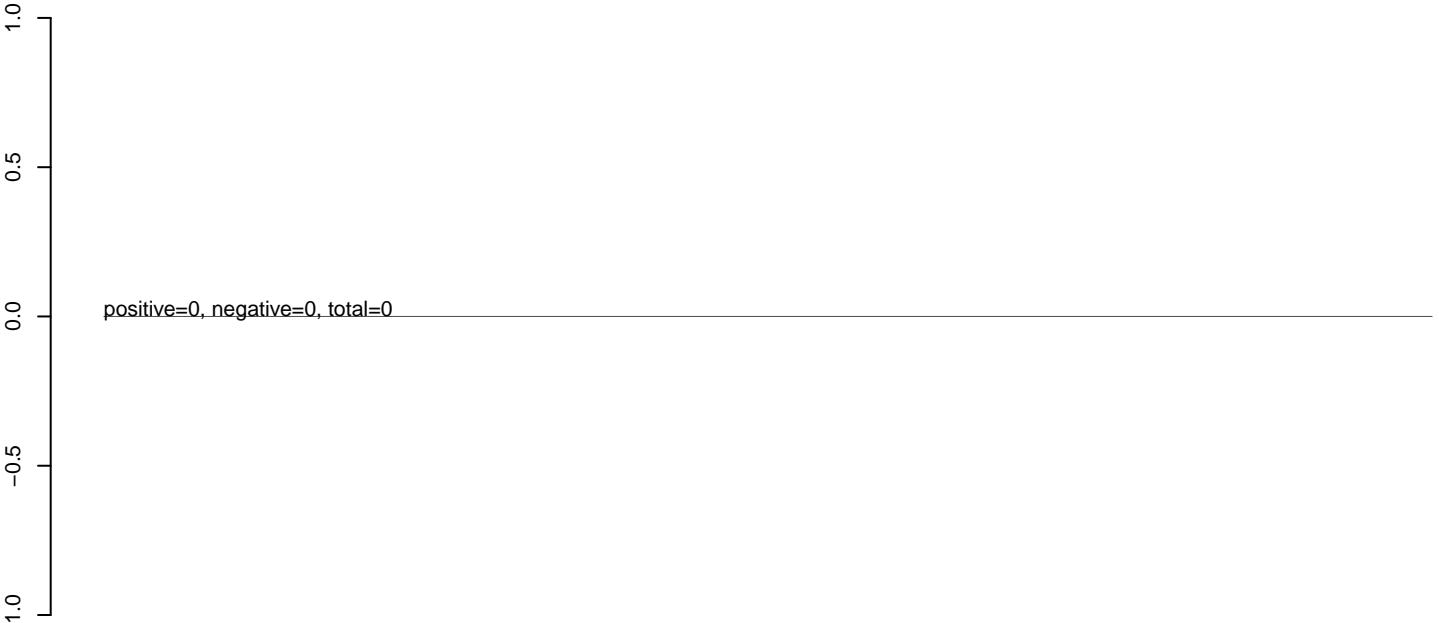
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

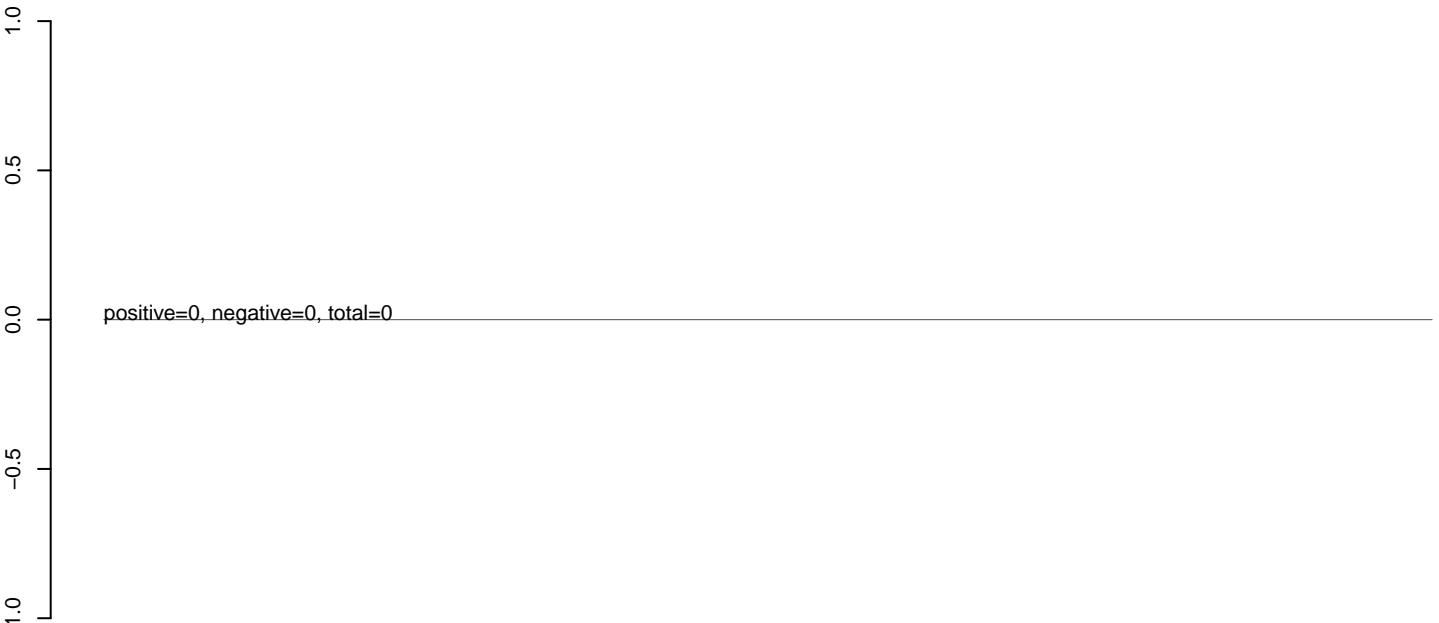
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=10682, virus@NC_001564.2–Cell_fusing_agent_virus_strain_Galveston:1–10682

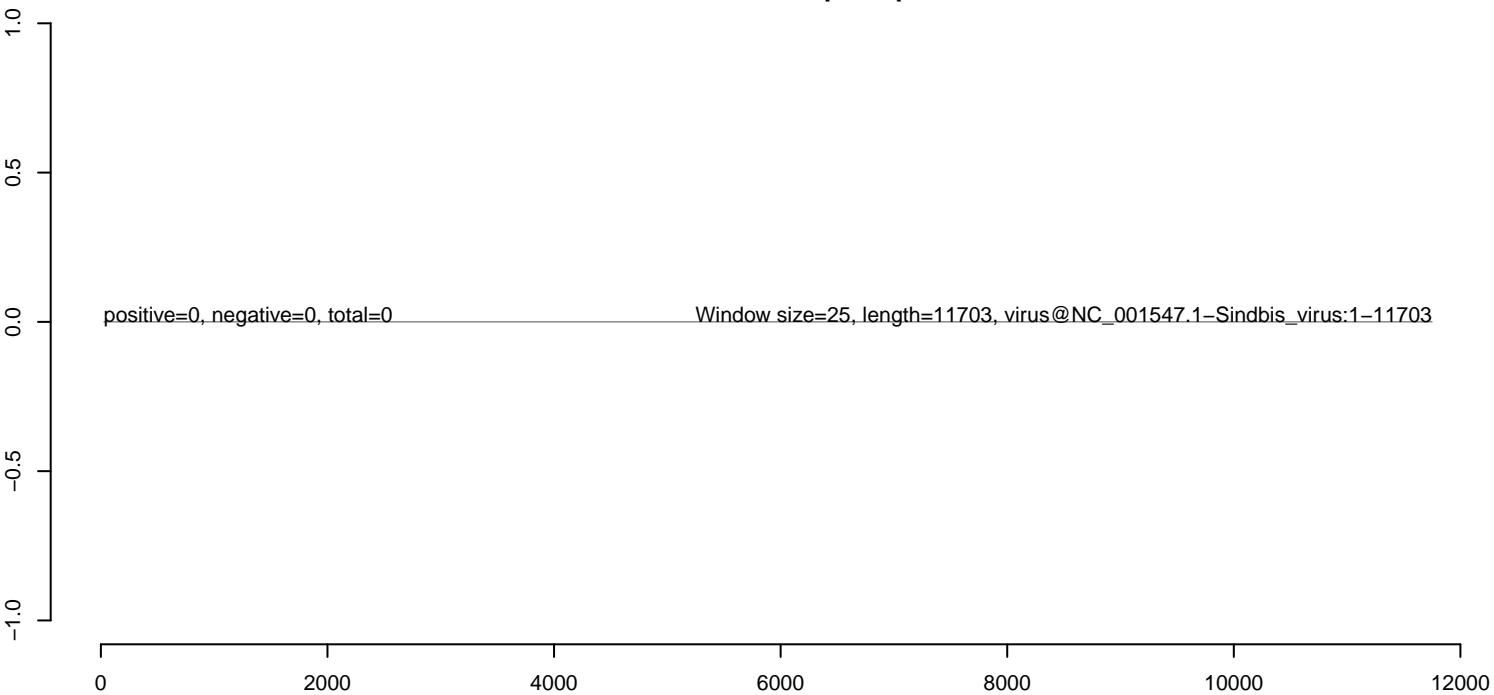
CuQuin_MG_7dpi3.18_23.rep



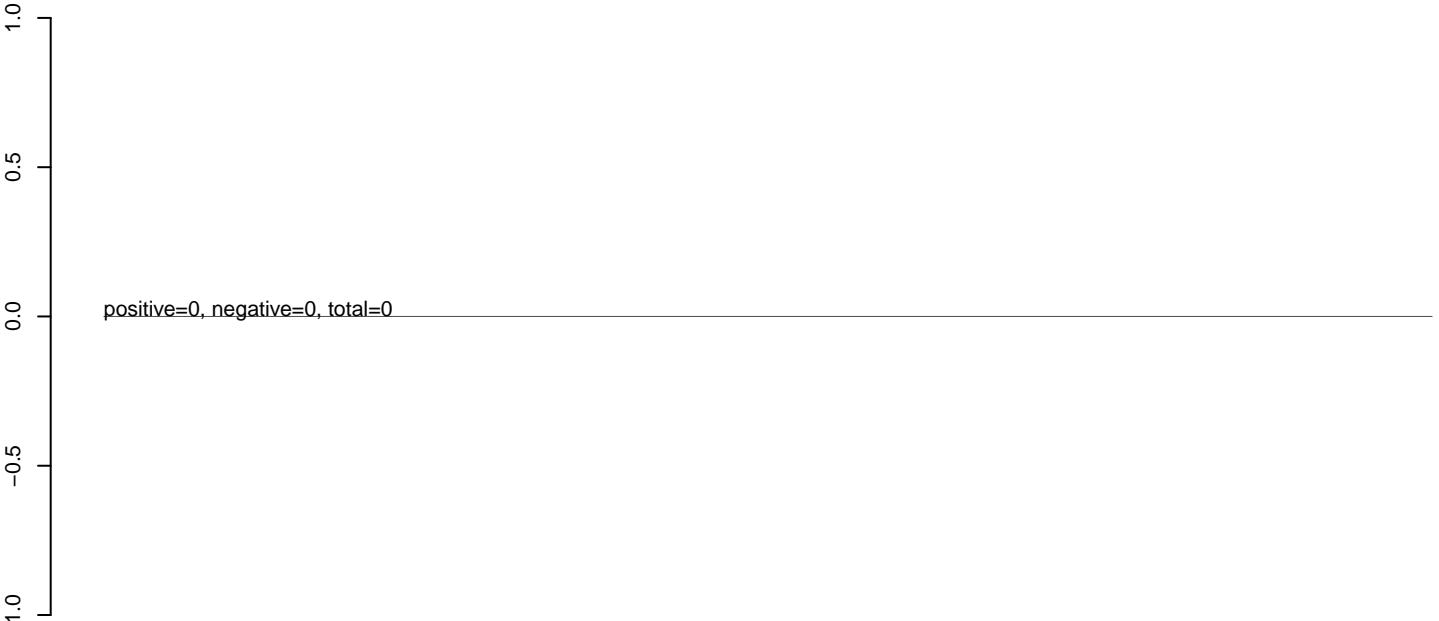
CuQuin_MG_7dpi3.24_35.rep



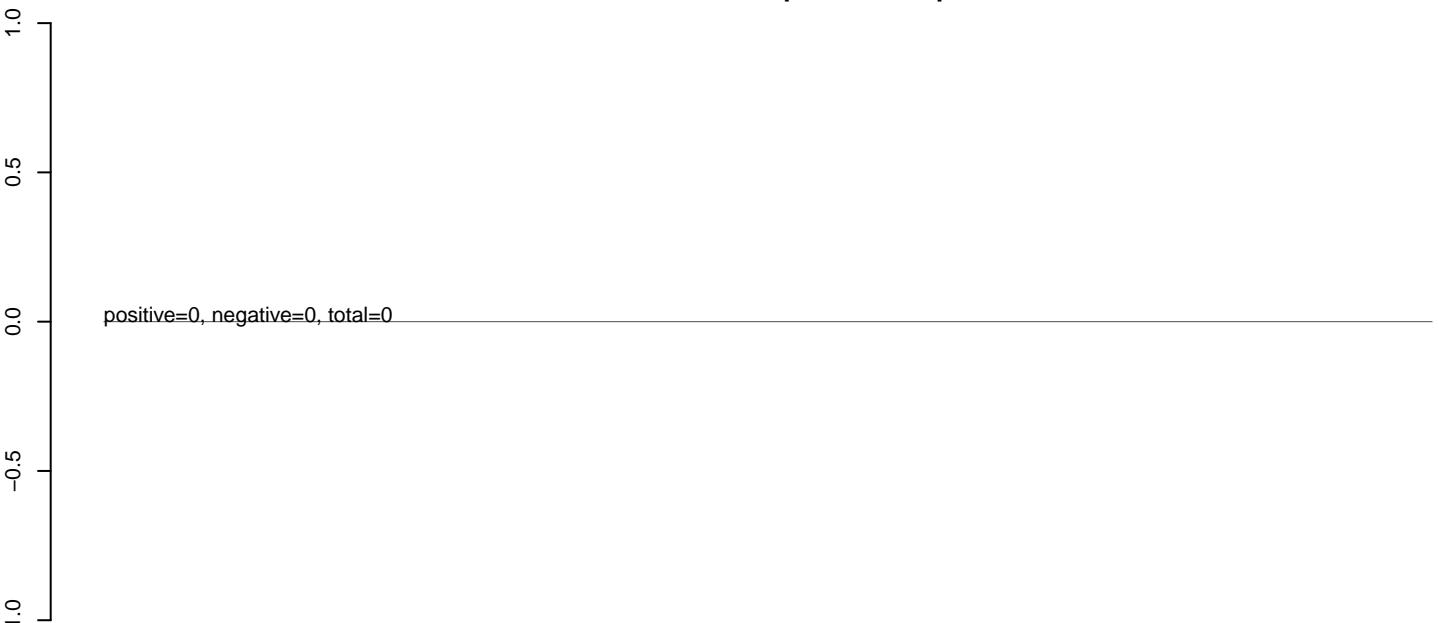
CuQuin_MG_7dpi3.rep



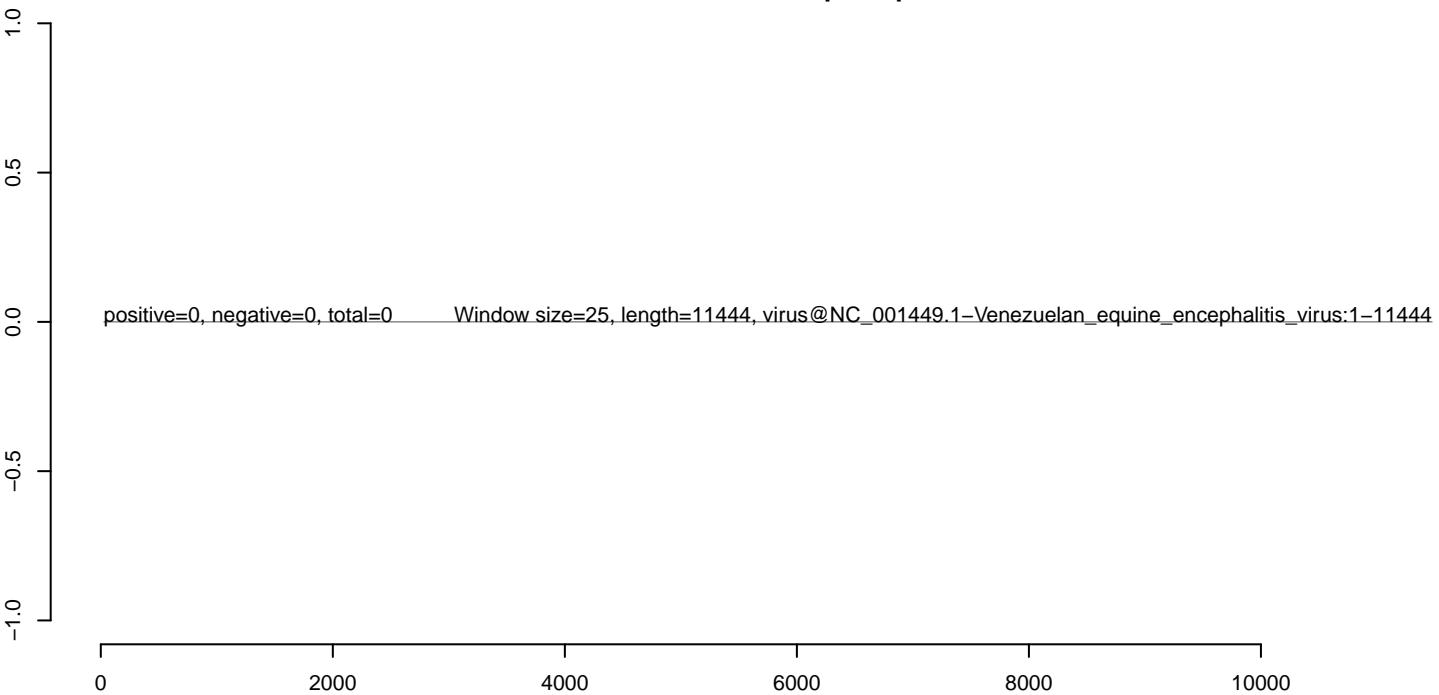
CuQuin_MG_7dpi3.18.rep



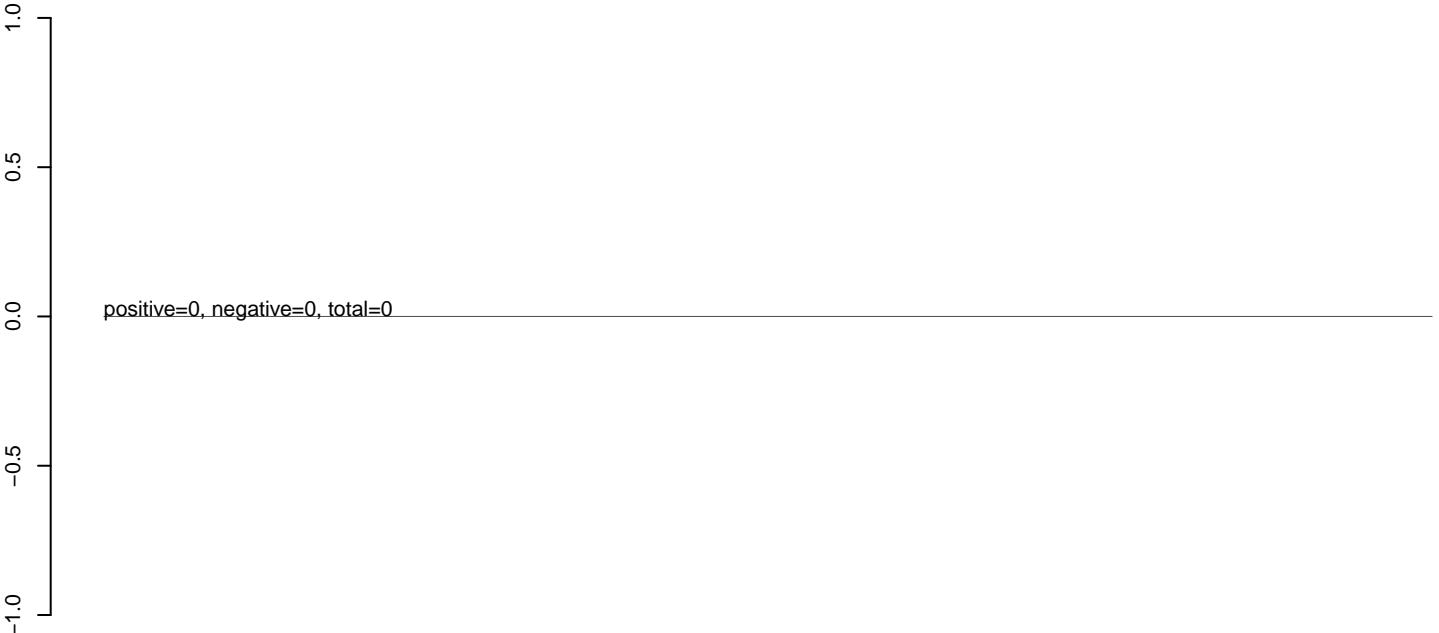
CuQuin_MG_7dpi3.24_35.rep



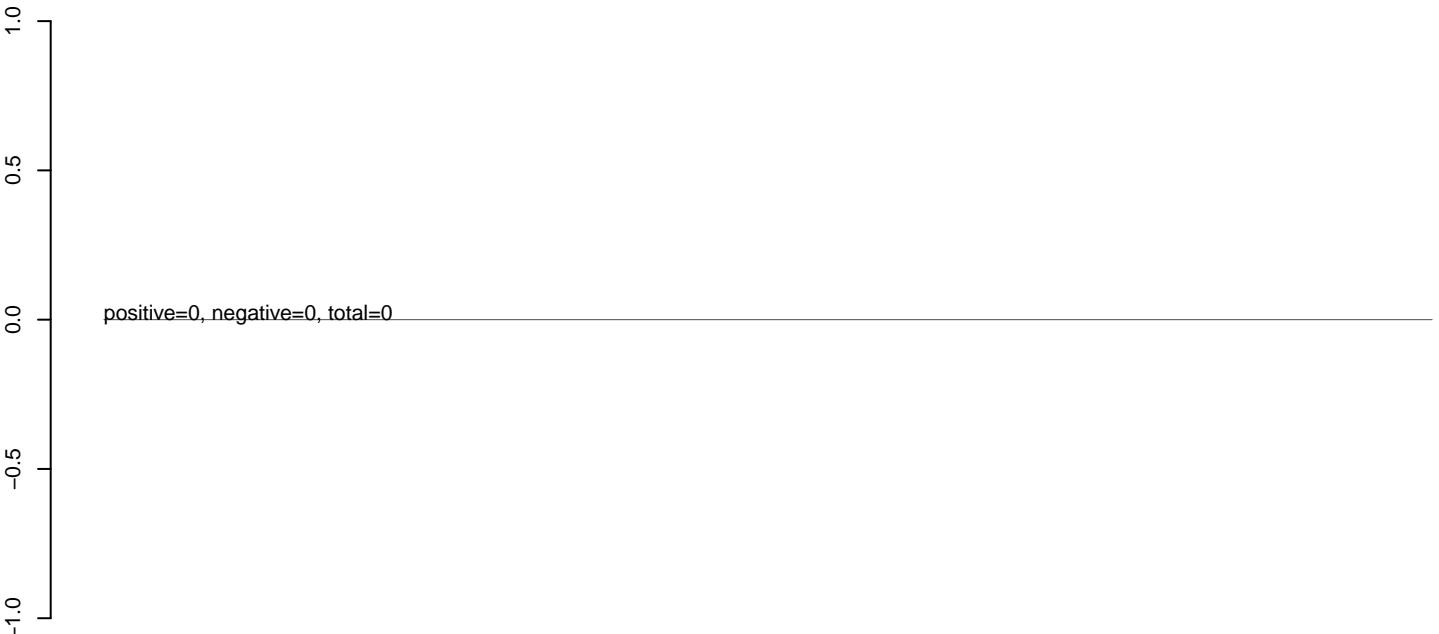
CuQuin_MG_7dpi3.rep



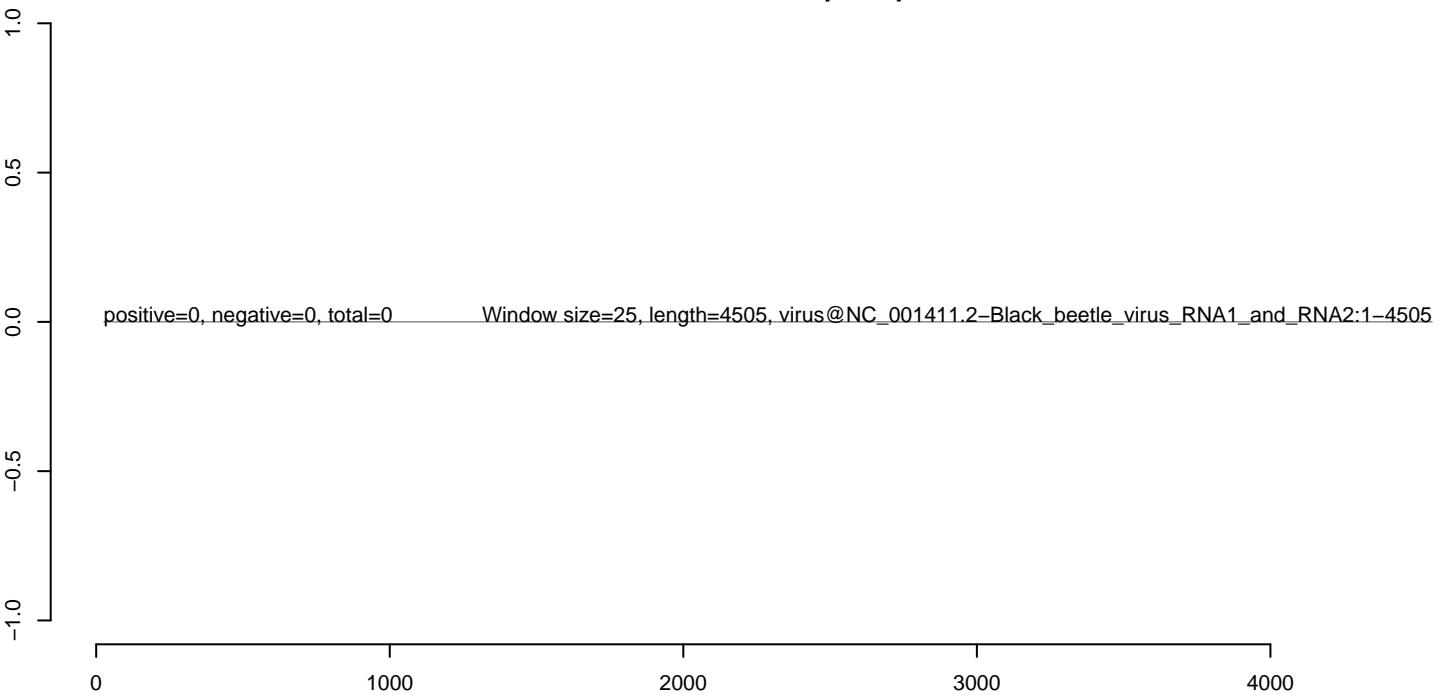
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0
positive_size=25, length=3937, virus@MH550148.1-Aedes_albopictus_densovirus_isolate_NC01_L3_1_trimmed_contig_5889-RC:1-3937

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=12916, virus@MH237595.1-Aedes_aegypti_anphevirus_strain_1a:1-12916

0 2000 4000 6000 8000 10000 12000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=3805, virus@MH188045.1-Culex_densovirus_strain_CDenv_Madera-4:1-3805

0 1000 2000 3000 4000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=3991, virus@MH188044.1-Culex_densovirus_strain_CDenv_Madera-3:1-3991

0 1000 2000 3000 4000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=3157, virus@MH188043.1-Culex_densovirus_strain_CDenv_Madera-2:1-3157

0 500 1000 1500 2000 2500 3000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

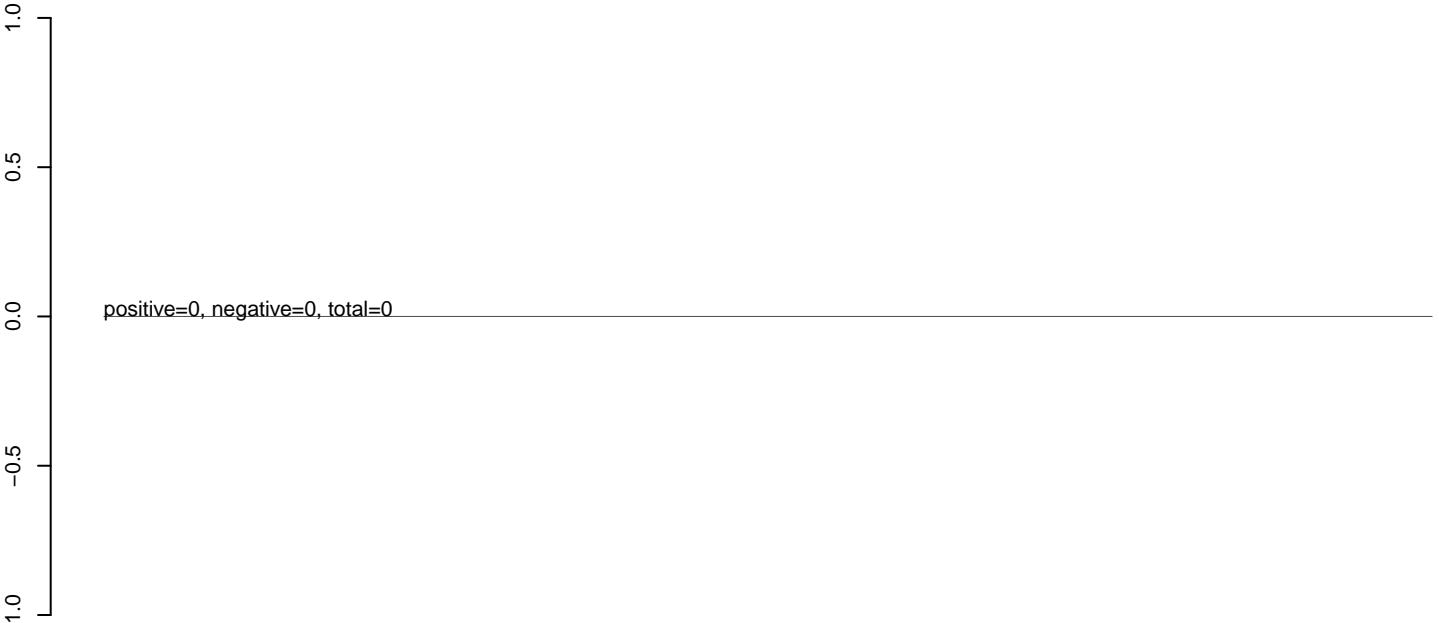
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

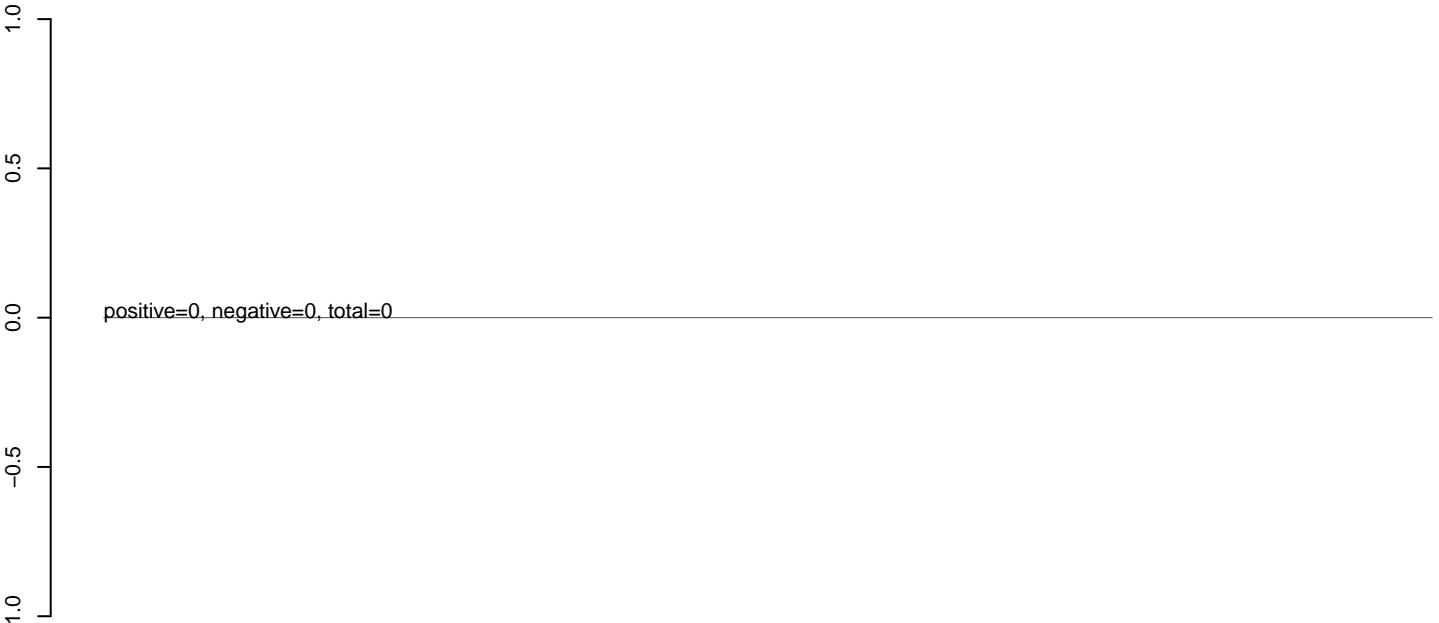
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0
Window size=25, length=12195, virus@MG747536.1_Oropouche_virus_isolate_BeAr_136921_segment_S_M_L_1-12195

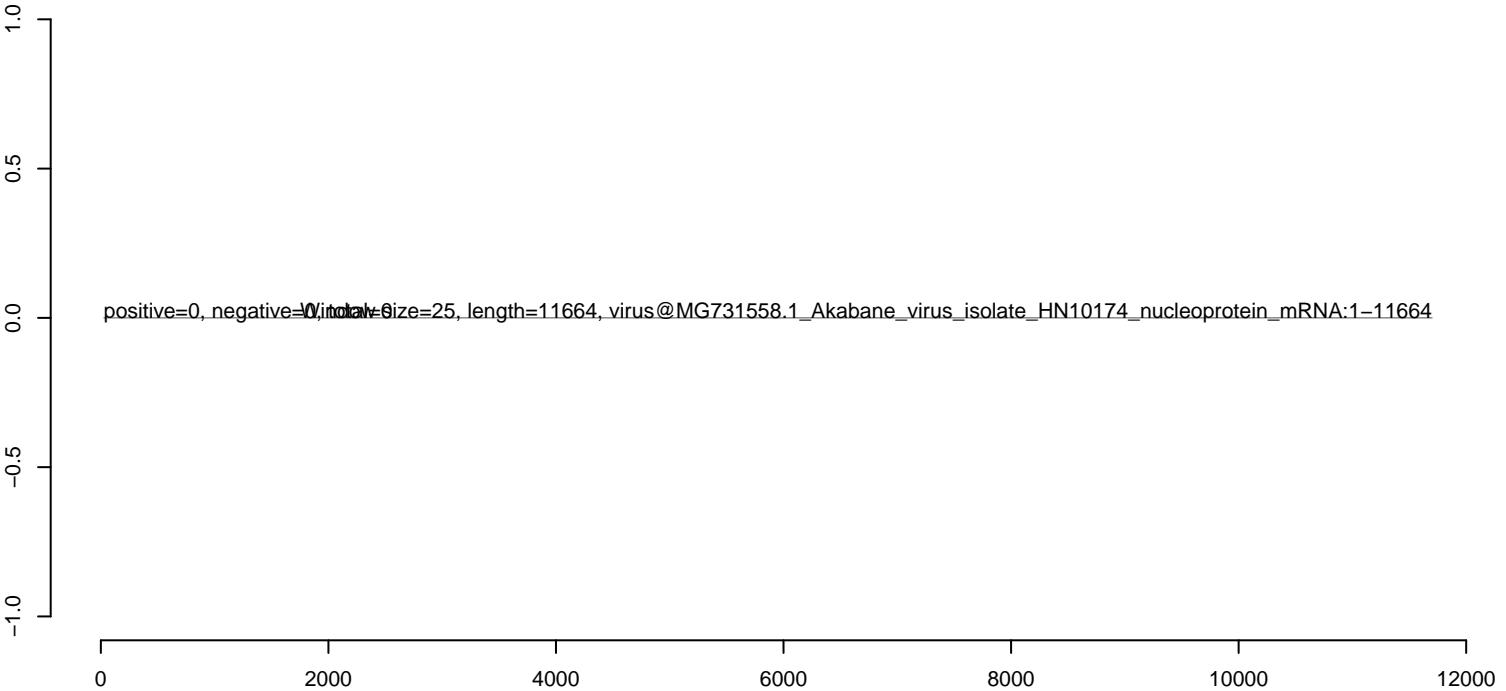
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

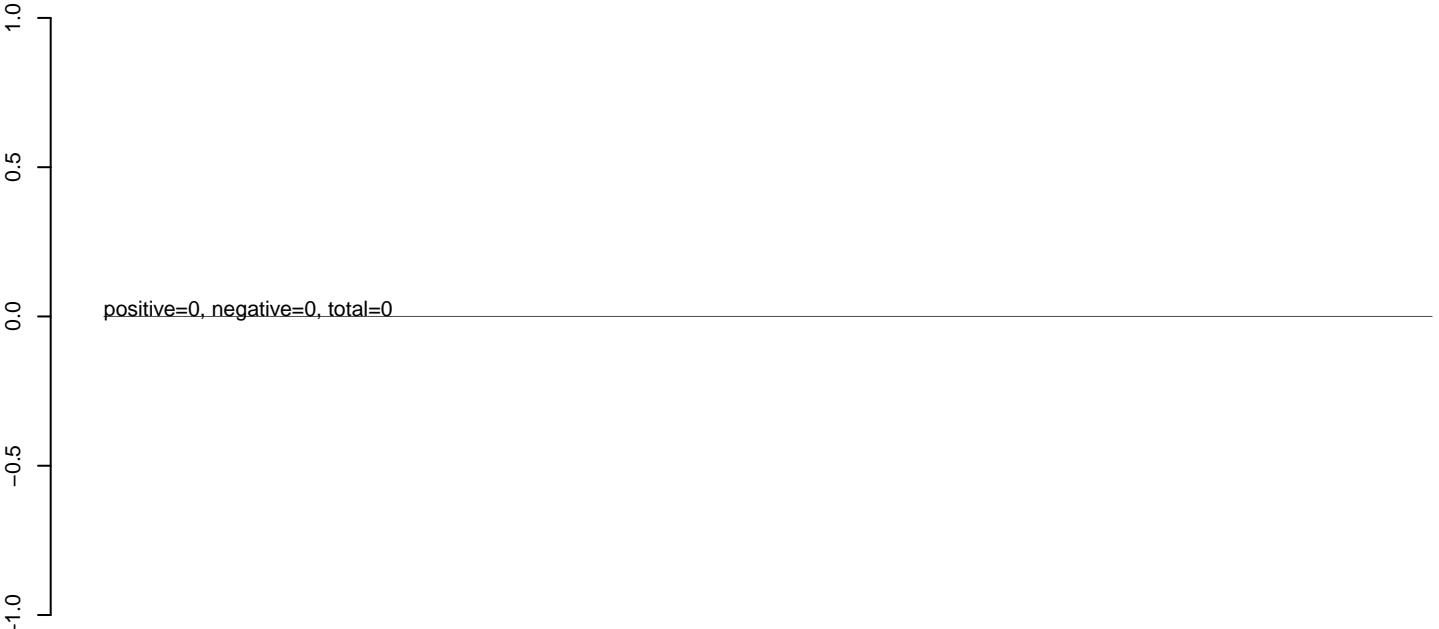
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

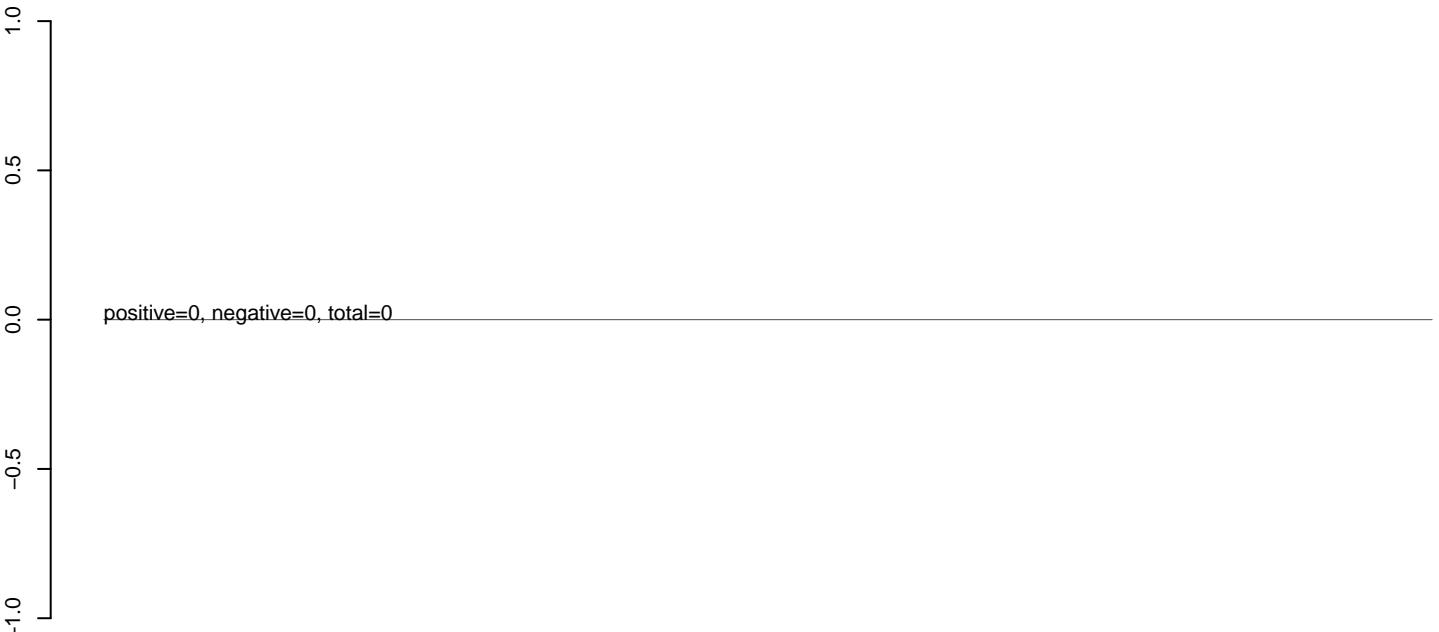
positive=0, negative=0, total=0, length=4245, virus@MF176386.1_Culex_luteo-like_virus_strain_mosWSX49277_segment_1_2:1-4245

0 1000 2000 3000 4000

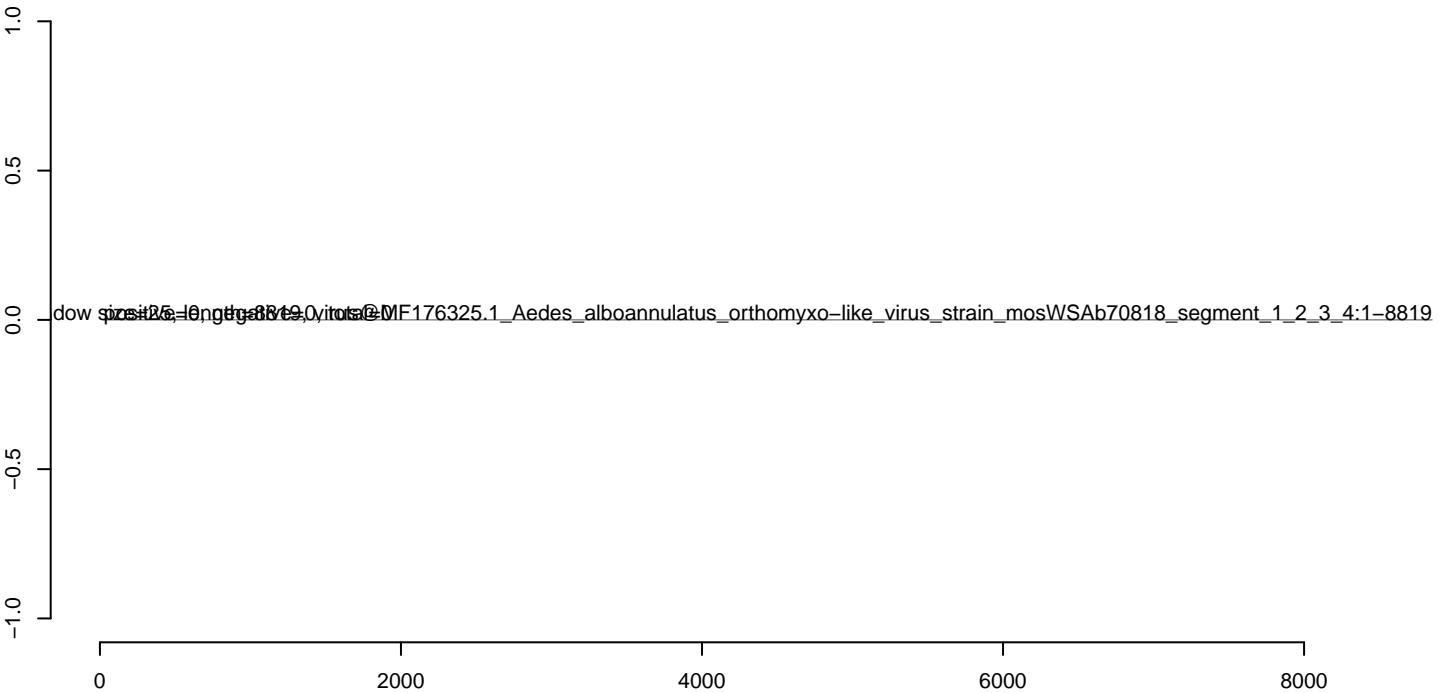
CuQuin_MG_7dpi3.18_23.rep



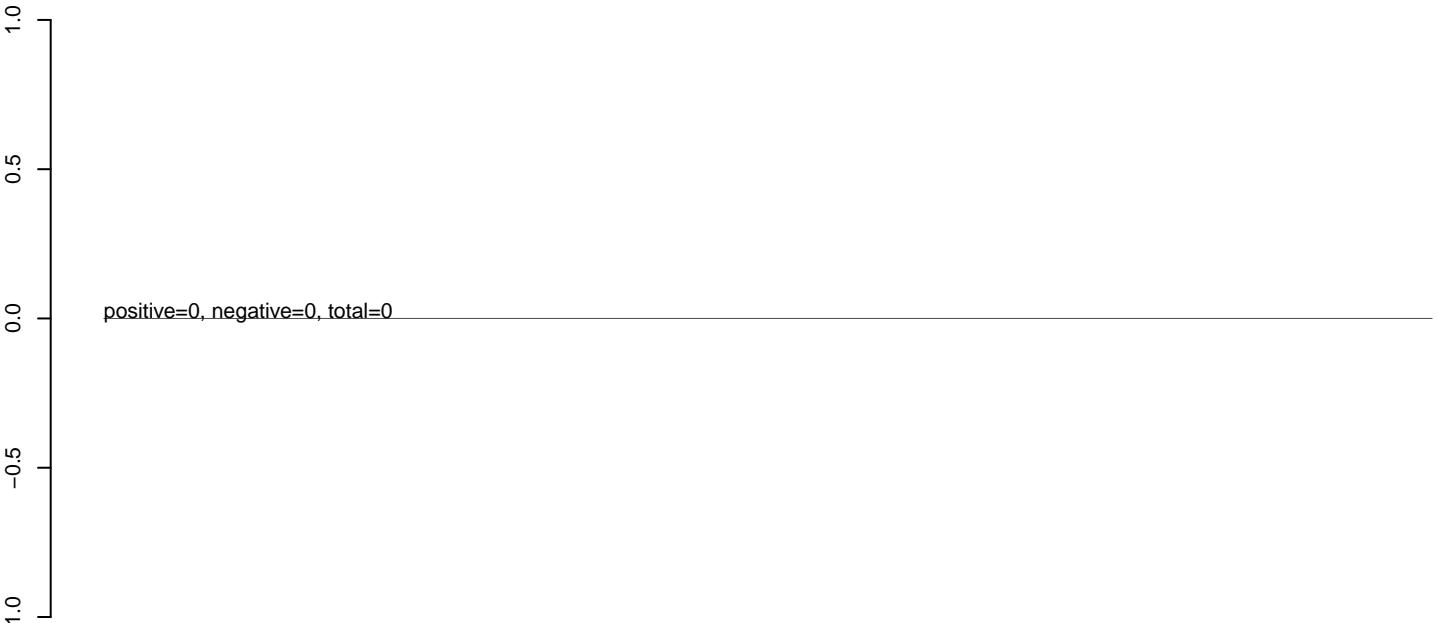
CuQuin_MG_7dpi3.24_35.rep



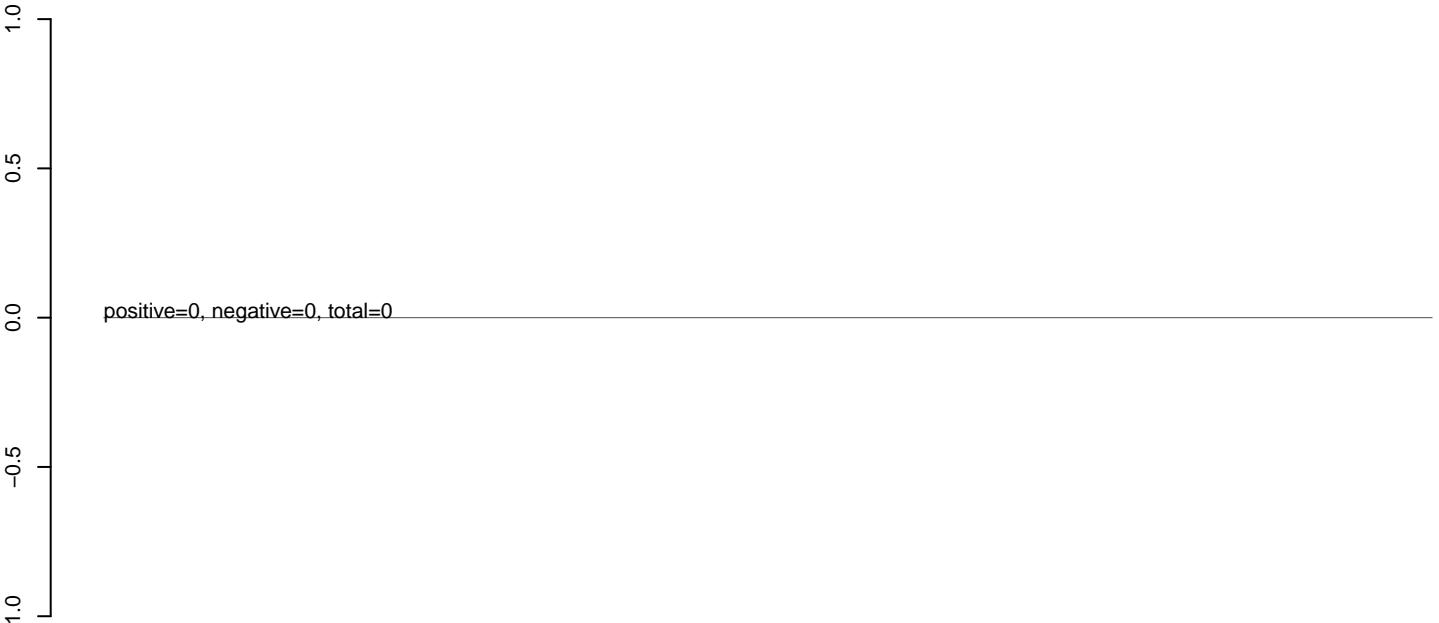
CuQuin_MG_7dpi3.rep



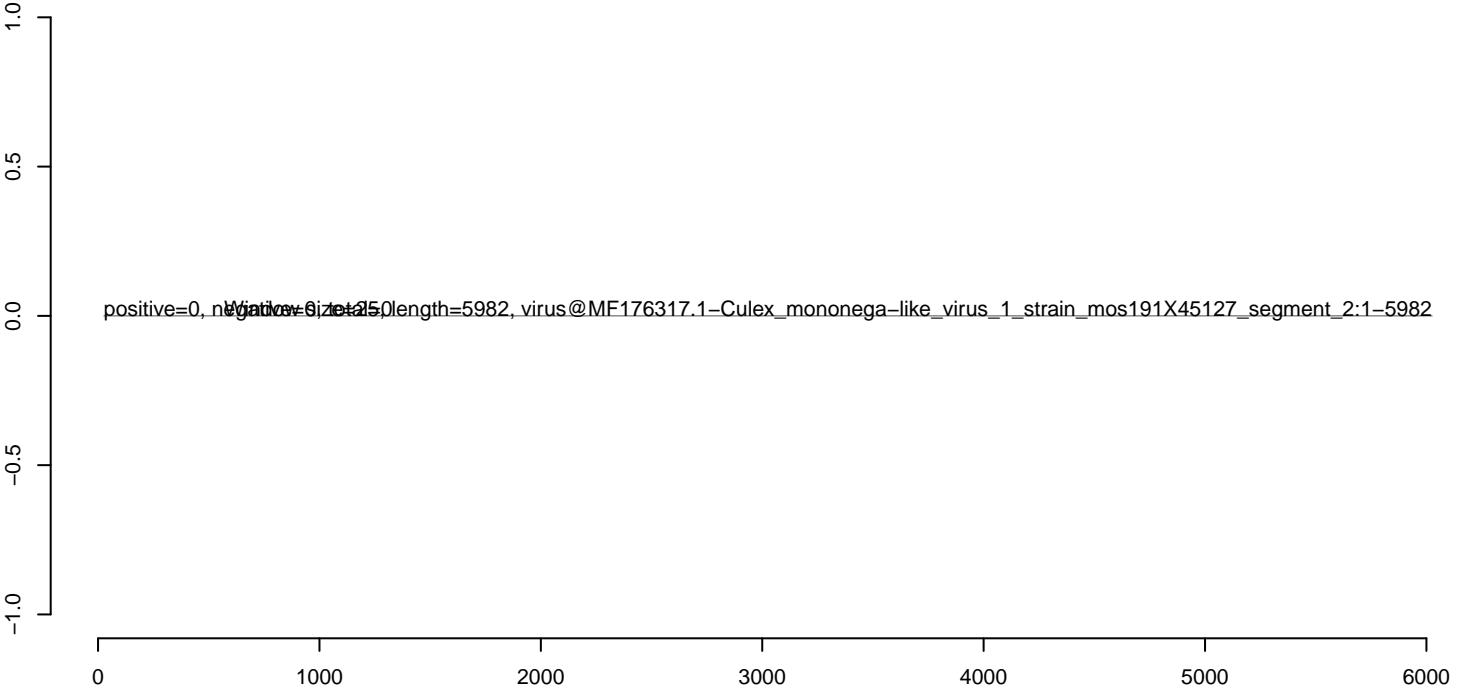
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

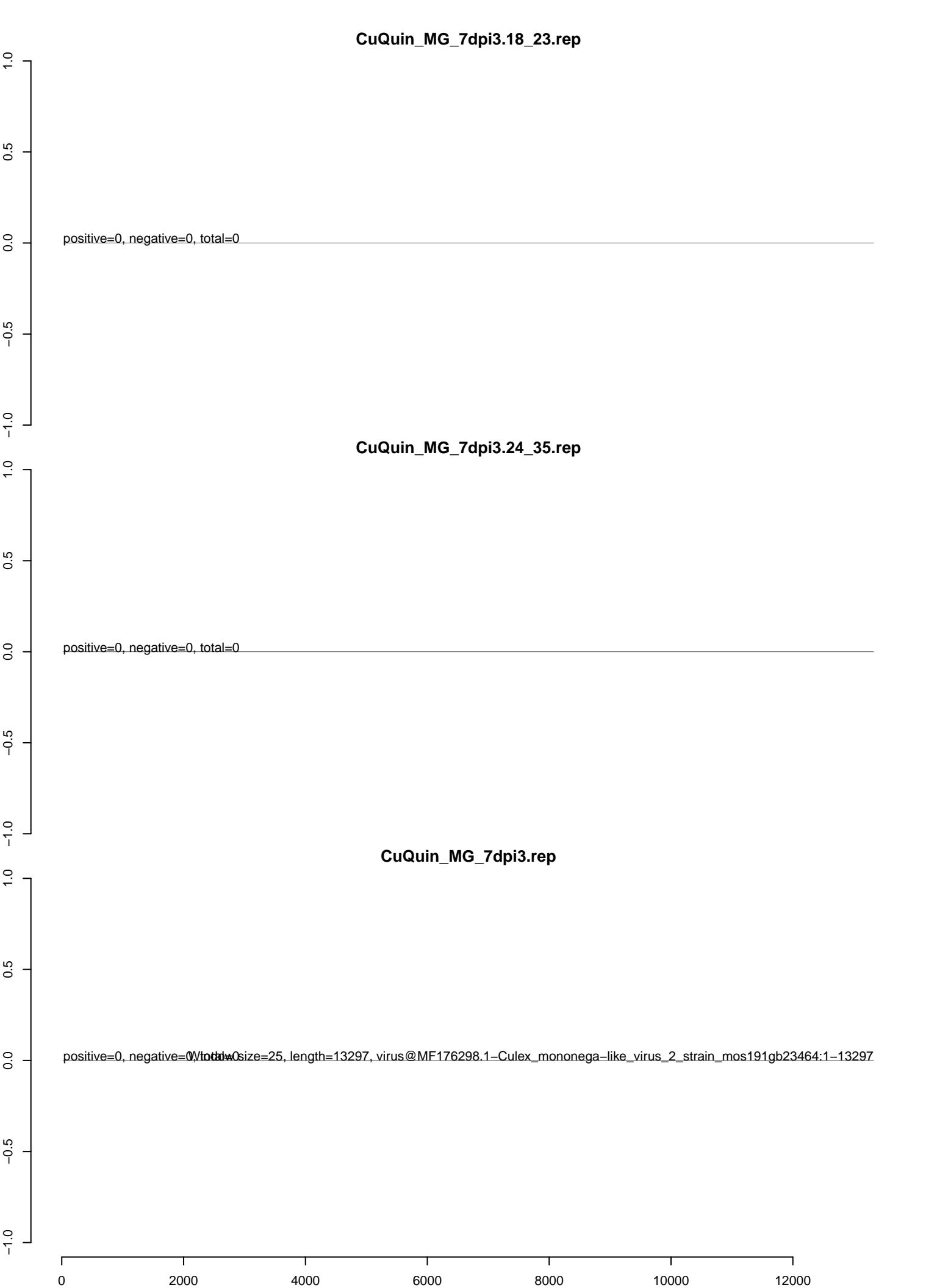
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0, size=25, length=13297, virus@MF176298.1-Culex_mononega-like_virus_2_strain_mos191gb23464:1-13297



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0
Window size=25, length=7347, virus@MF176289.1_Wilkie_qin-like_virus_strain_mos182CP103635_segment_1_2:1-7347

0 2000 4000 6000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

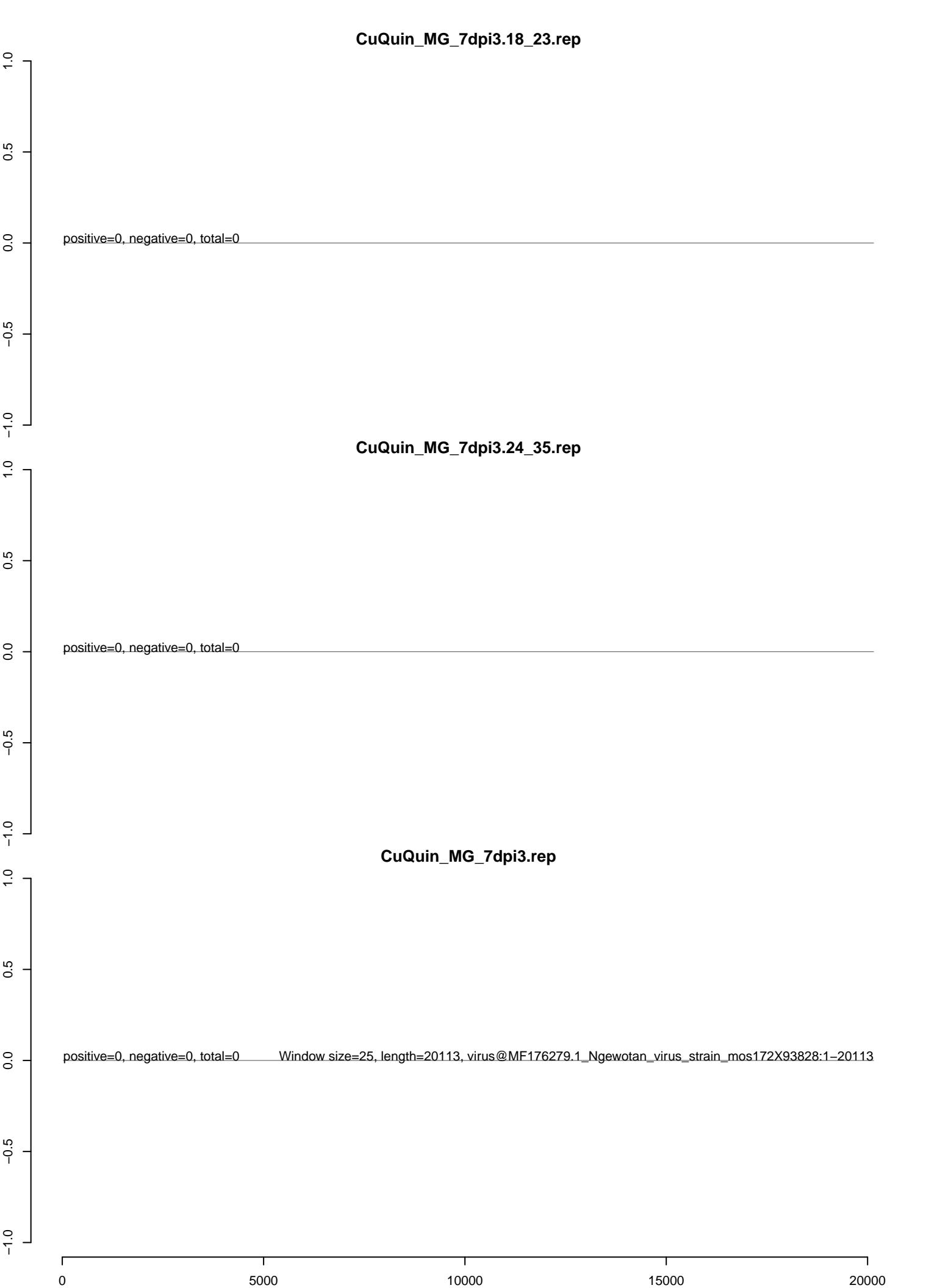
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=20113, virus@MF176279.1_Ngewotan_virus_strain_mos172X93828:1–20113



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=25, total=0, length=10814, virus@MF176267.1_Culex_phasma-like_virus_strain_mos172X35099_segment_S_M_L:1-10814

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=3978, virus@M37899.1-Aedes_aegypti_densovirus_1_genomic_sequence:1-3978

CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

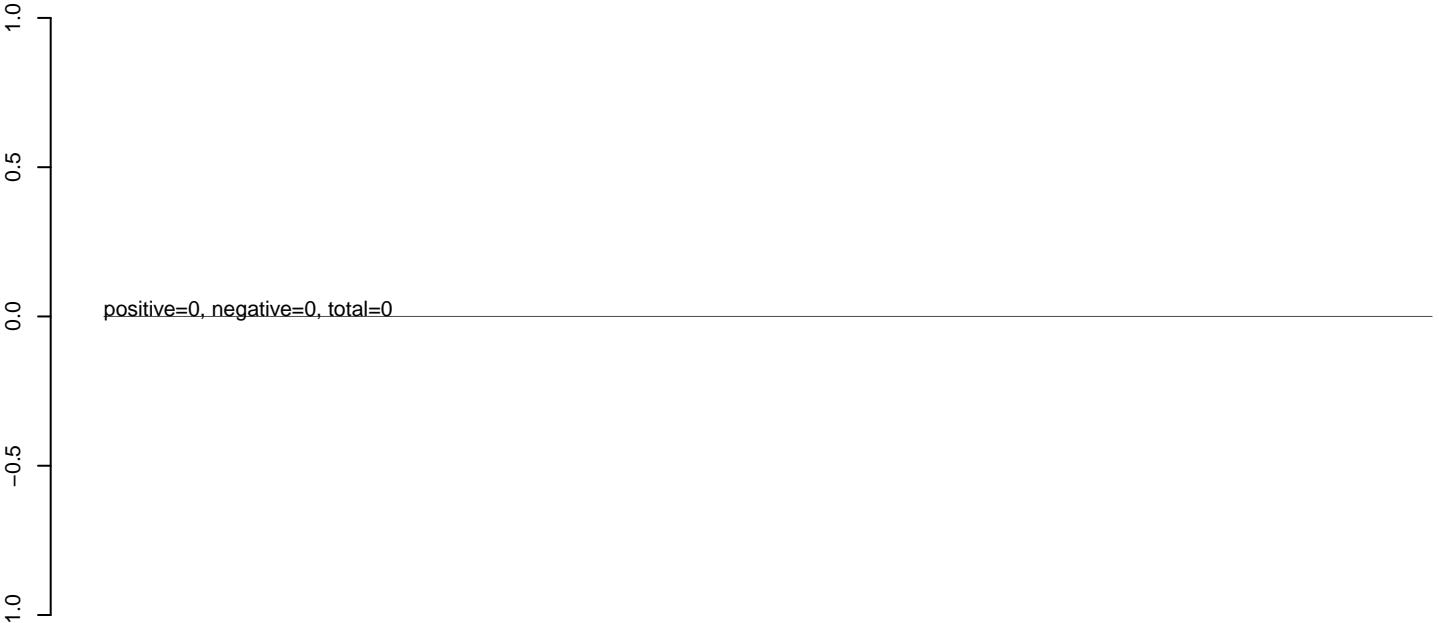
CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

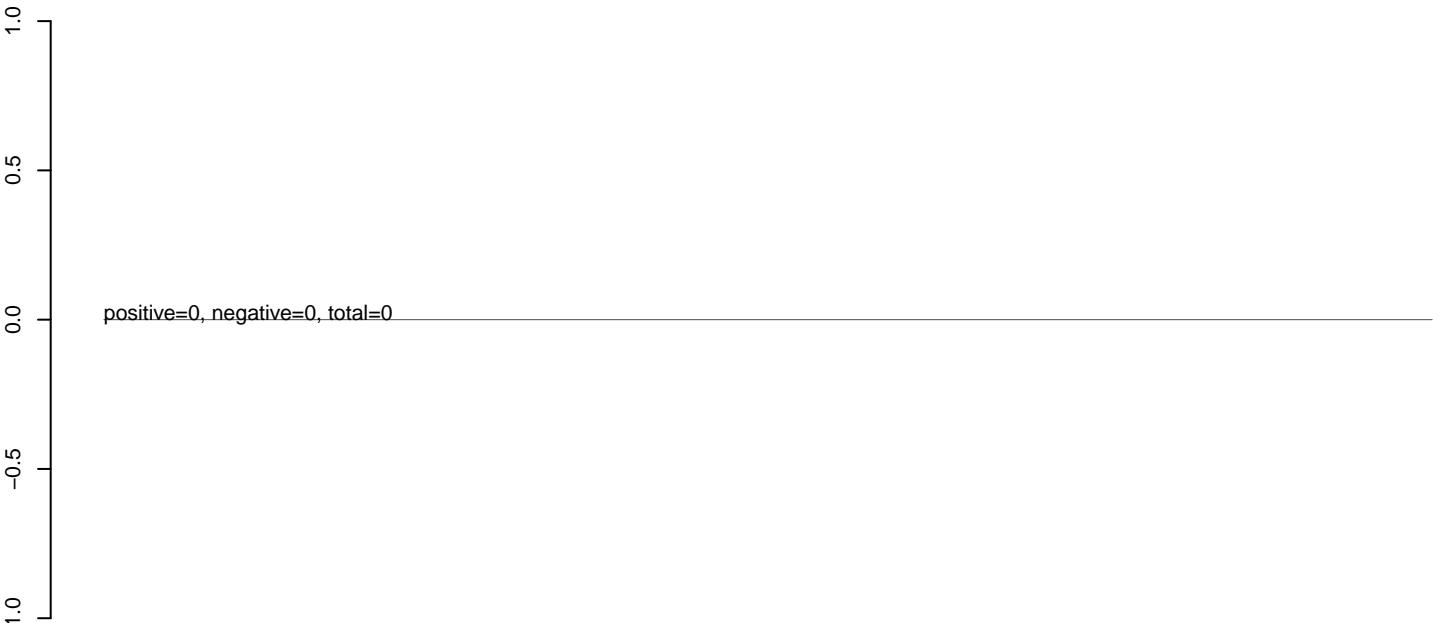
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0, window size=25, length=11835, virus@M20303.1-O-Nyong-nyong_virus_Gulu_strain_complete_genome:1-11835

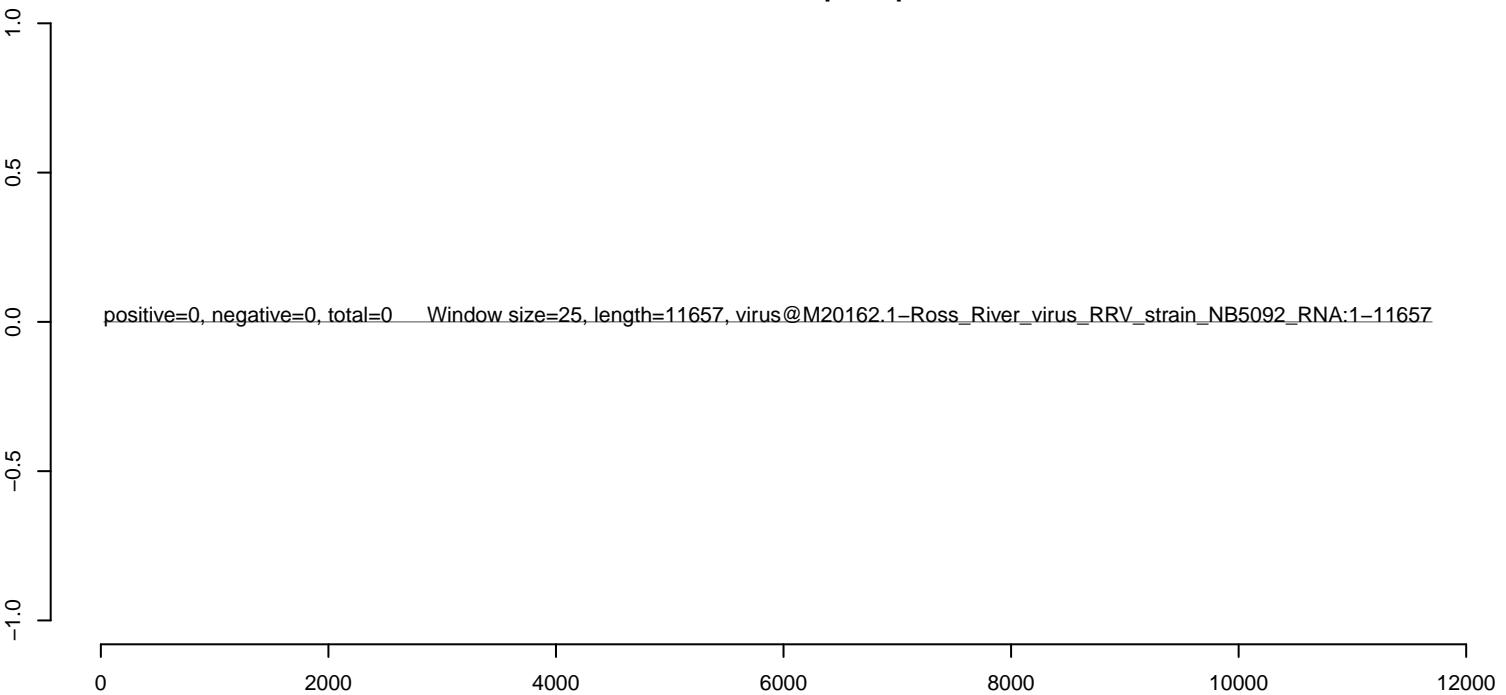
CuQuin_MG_7dpi3.18_23.rep



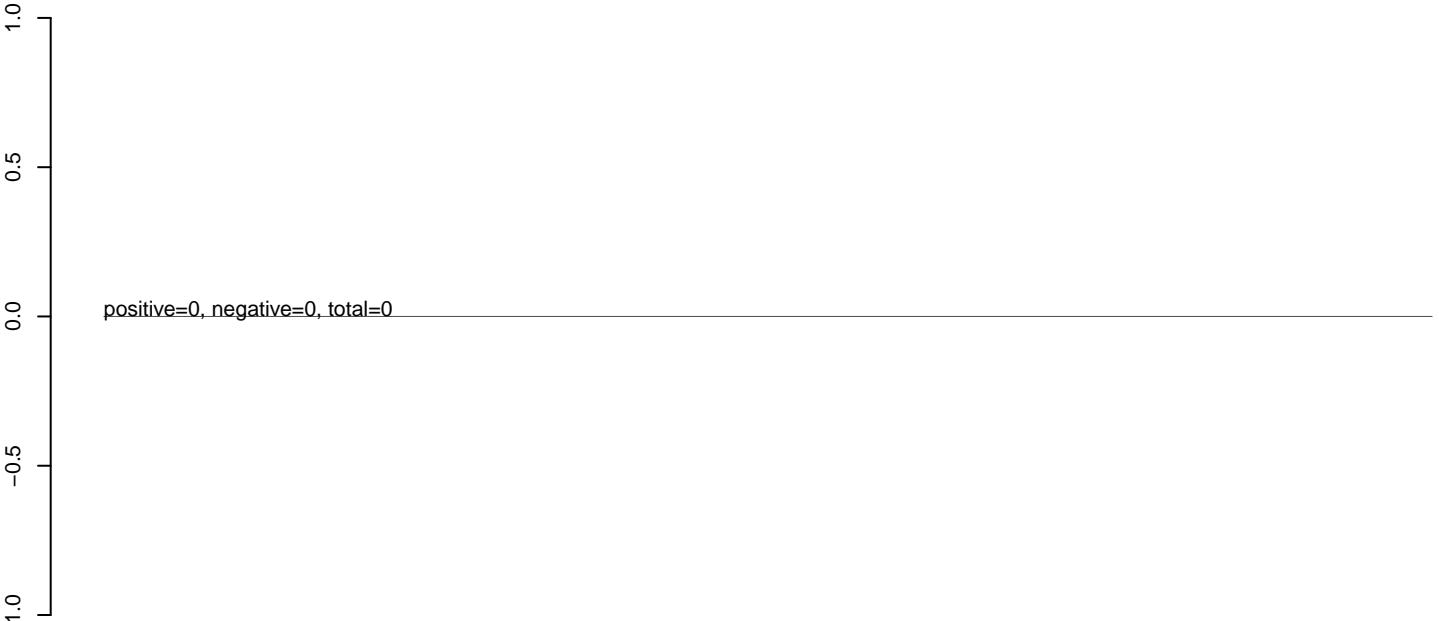
CuQuin_MG_7dpi3.24_35.rep



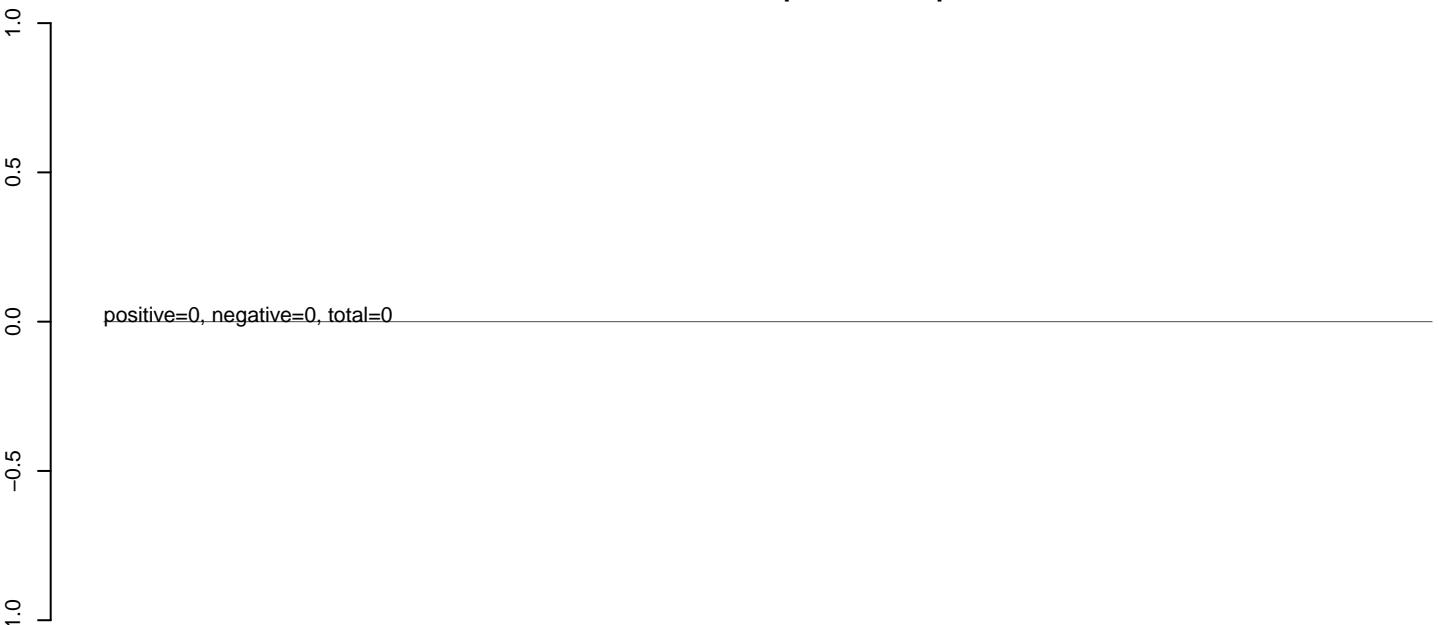
CuQuin_MG_7dpi3.rep



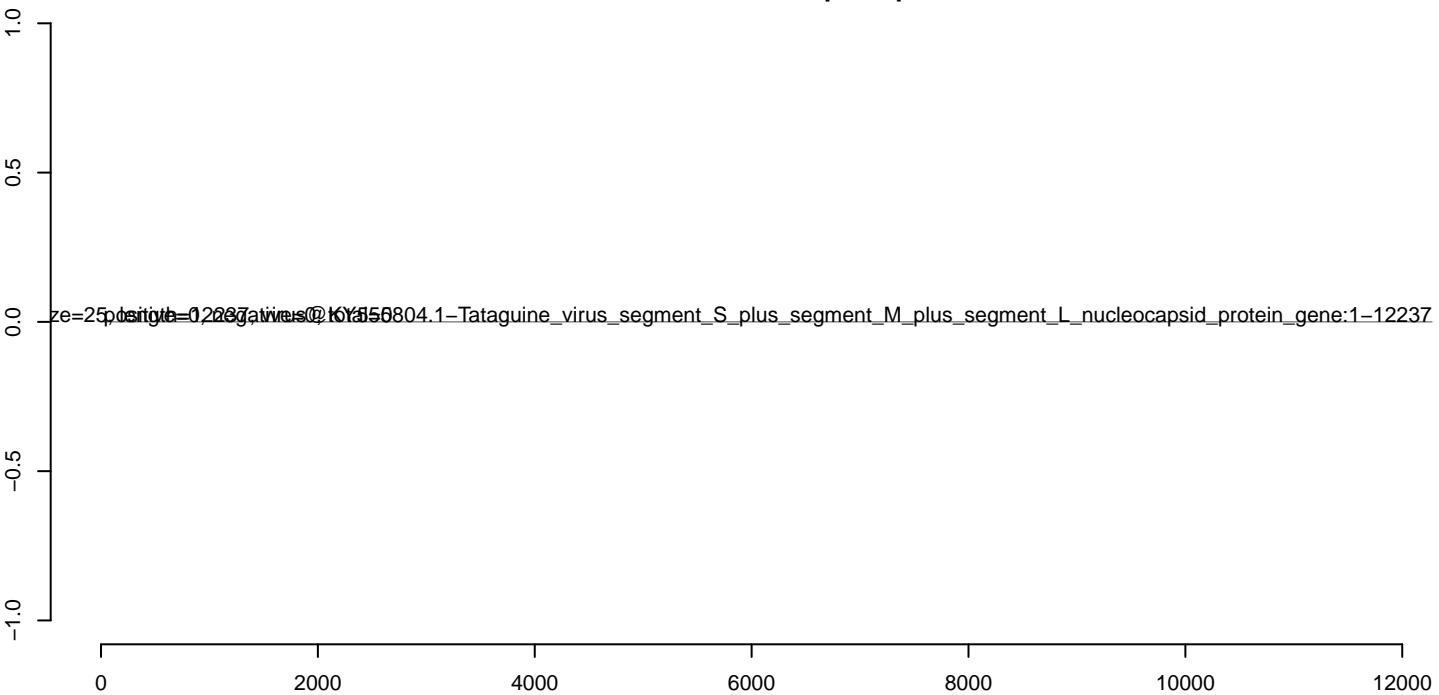
CuQuin_MG_7dpi3.18_23.rep



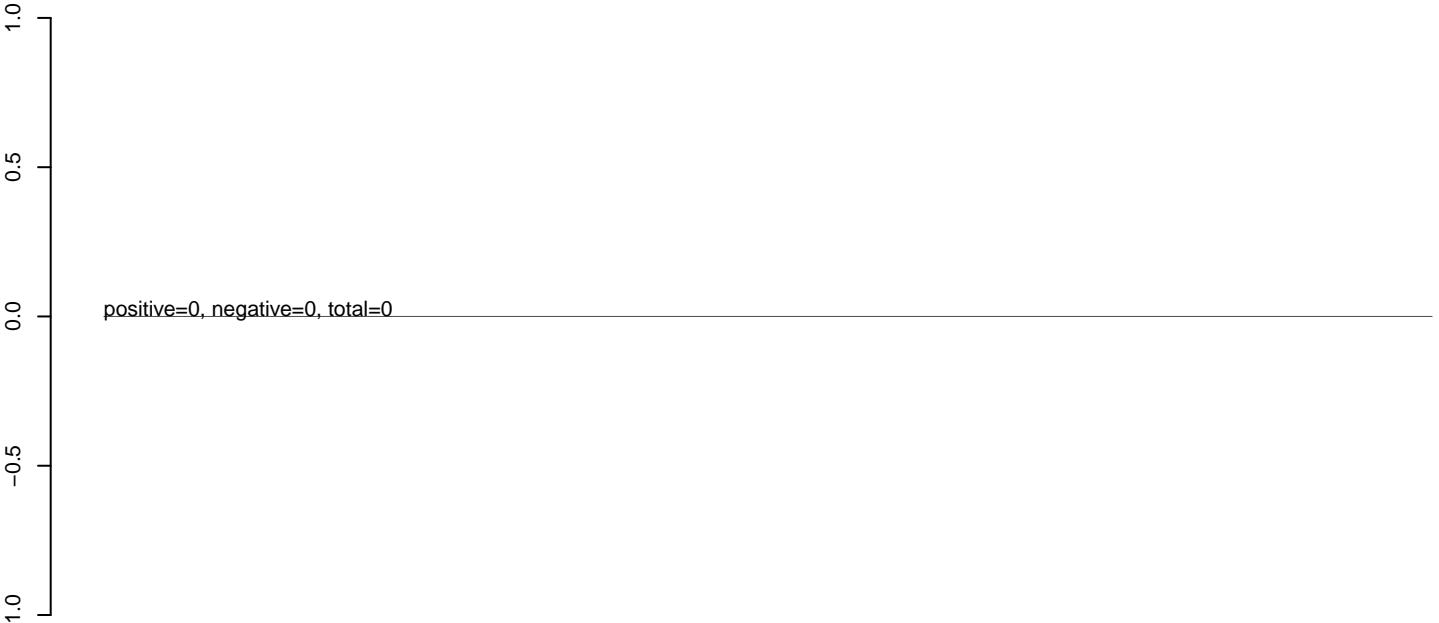
CuQuin_MG_7dpi3.24_35.rep



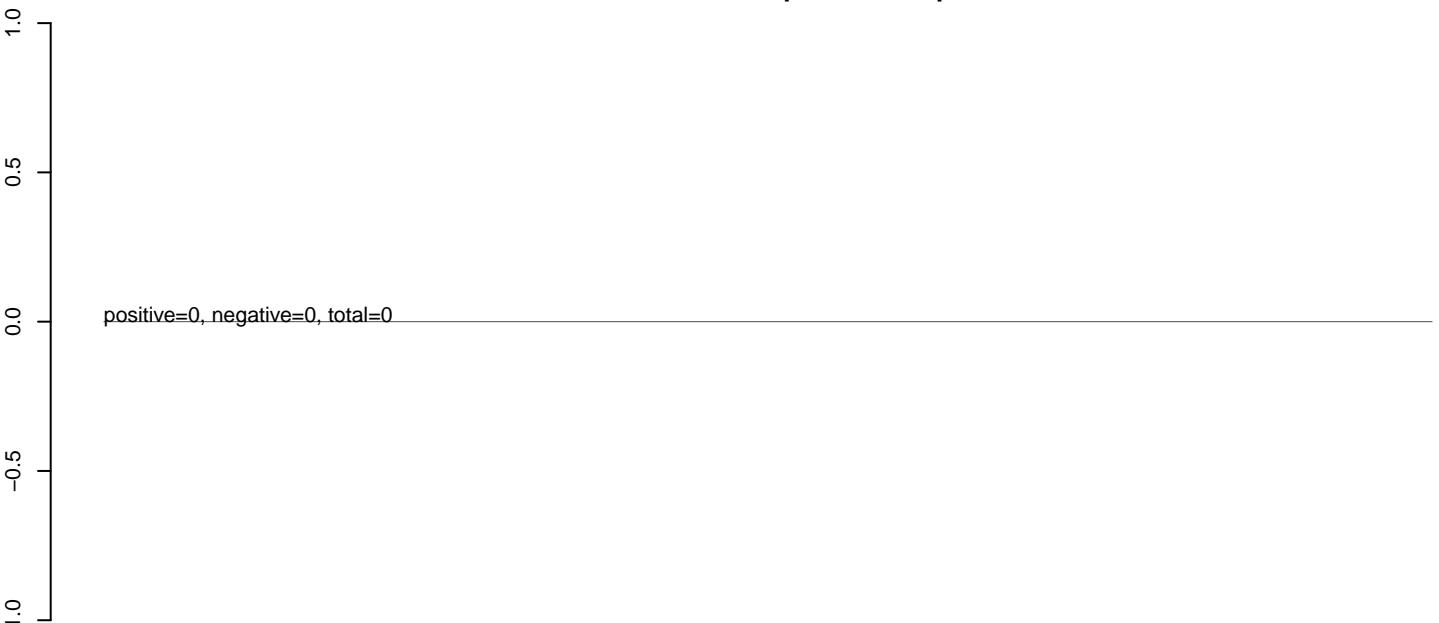
CuQuin_MG_7dpi3.rep



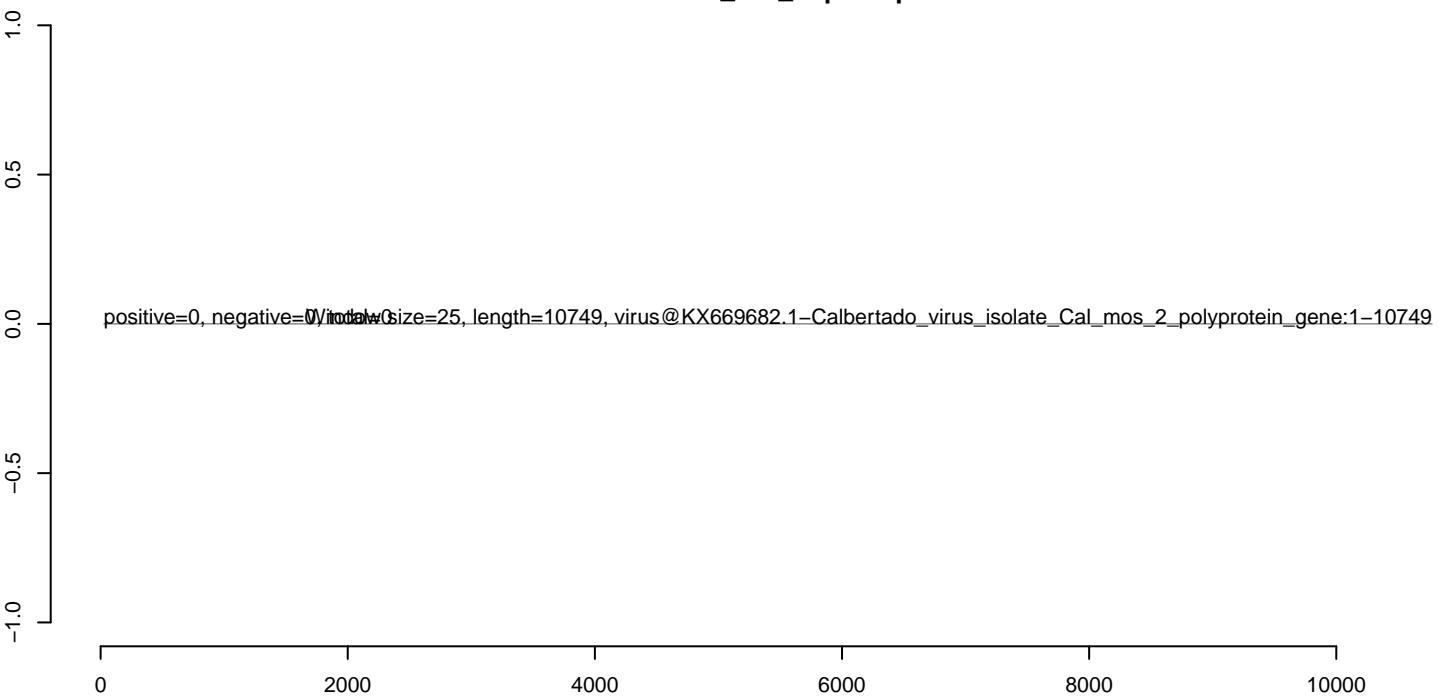
CuQuin_MG_7dpi3.18_23.rep



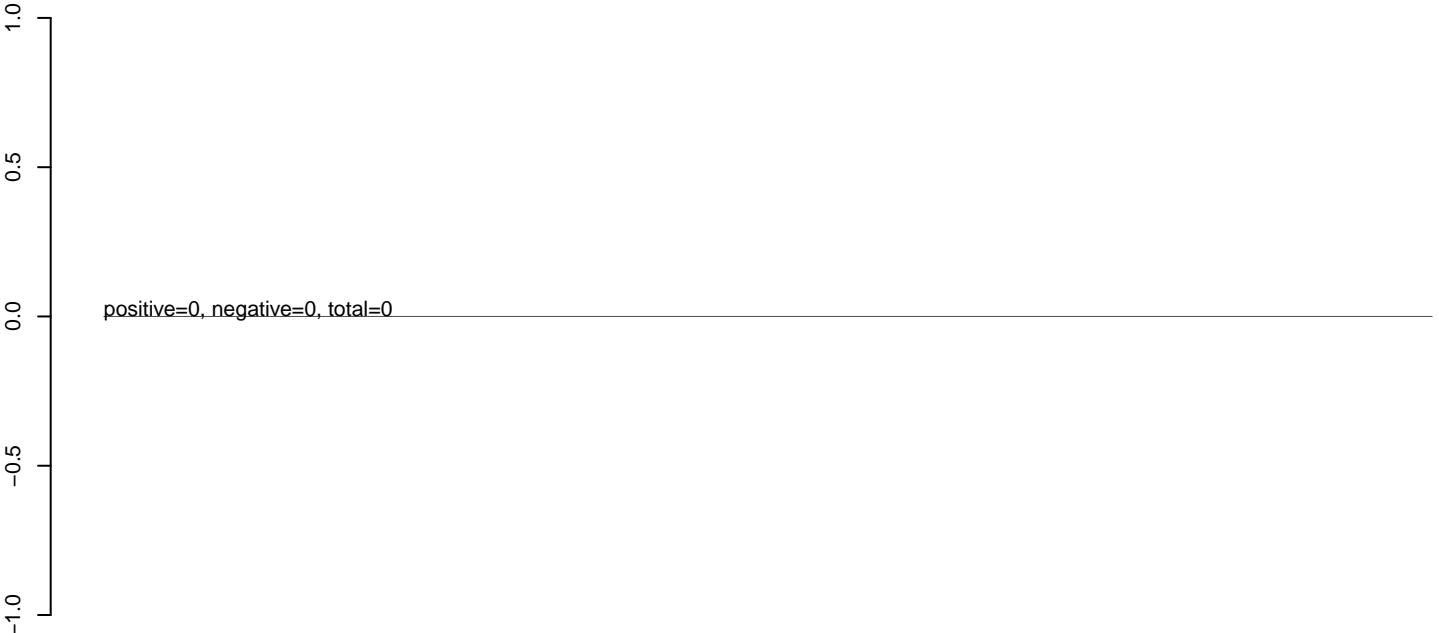
CuQuin_MG_7dpi3.24_35.rep



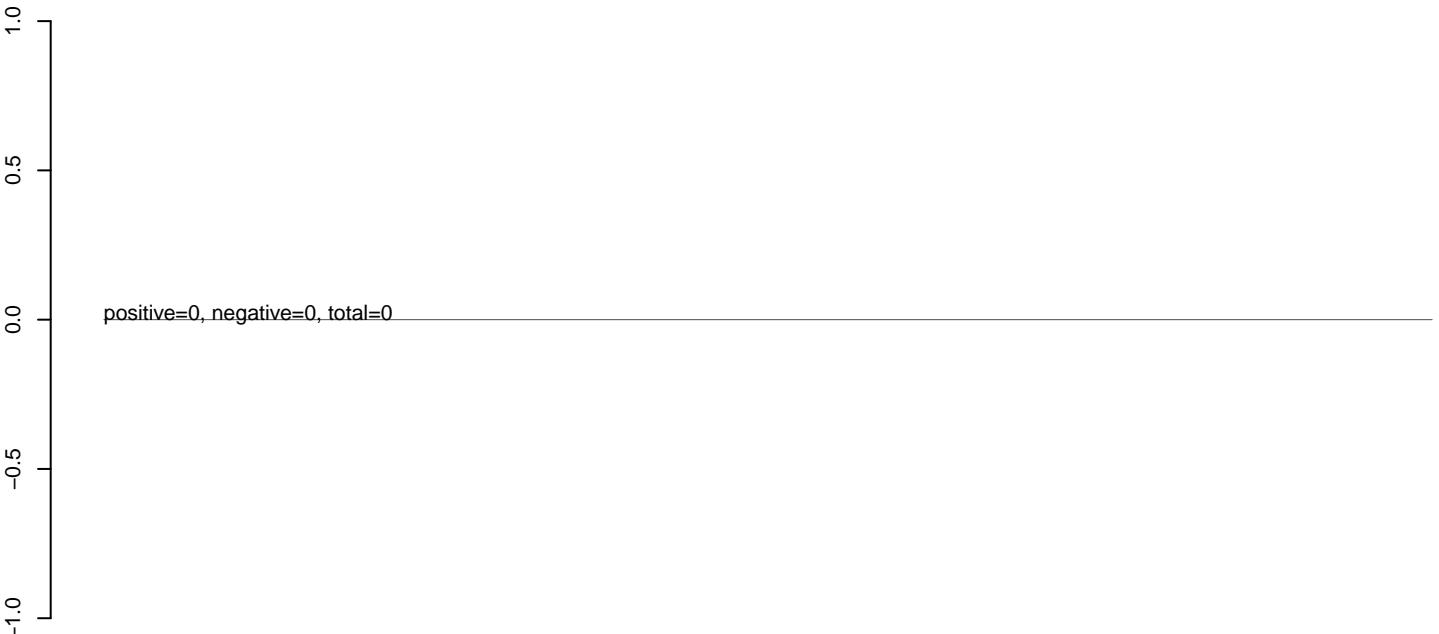
CuQuin_MG_7dpi3.rep



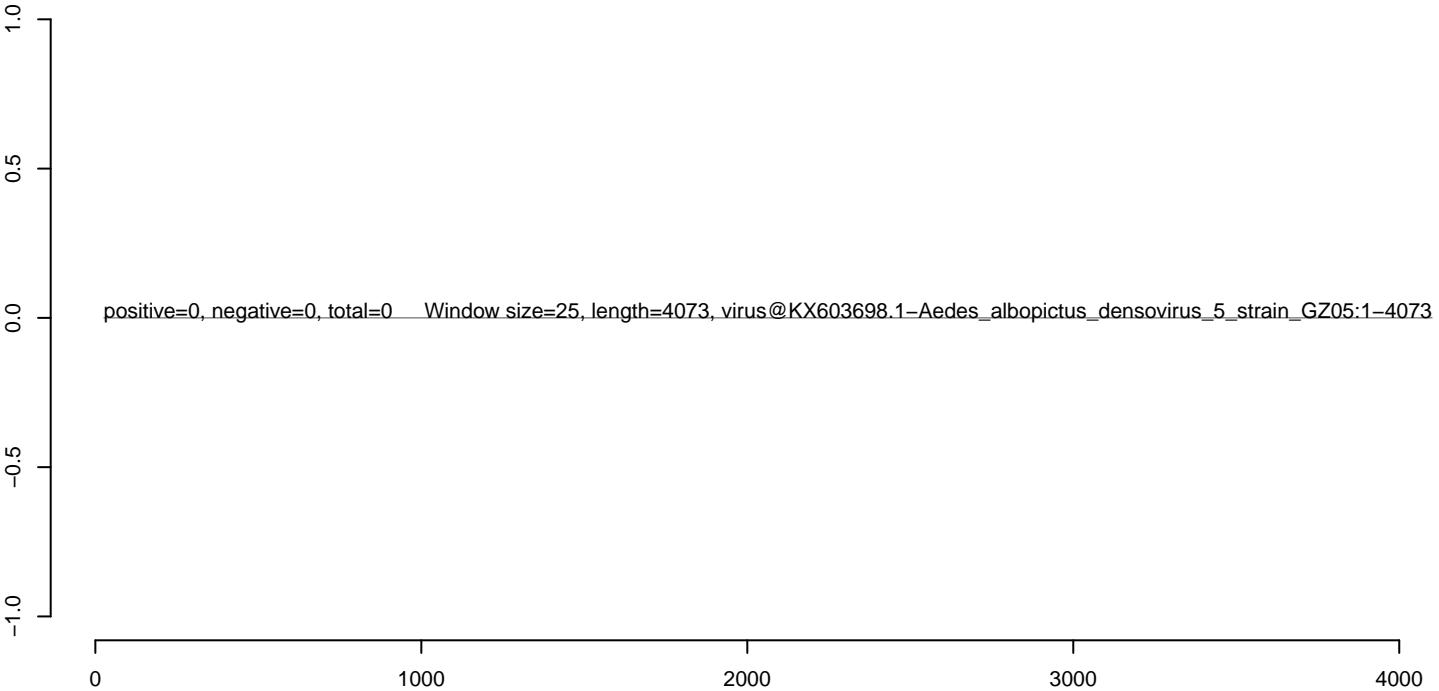
CuQuin_MG_7dpi3.18_23.rep



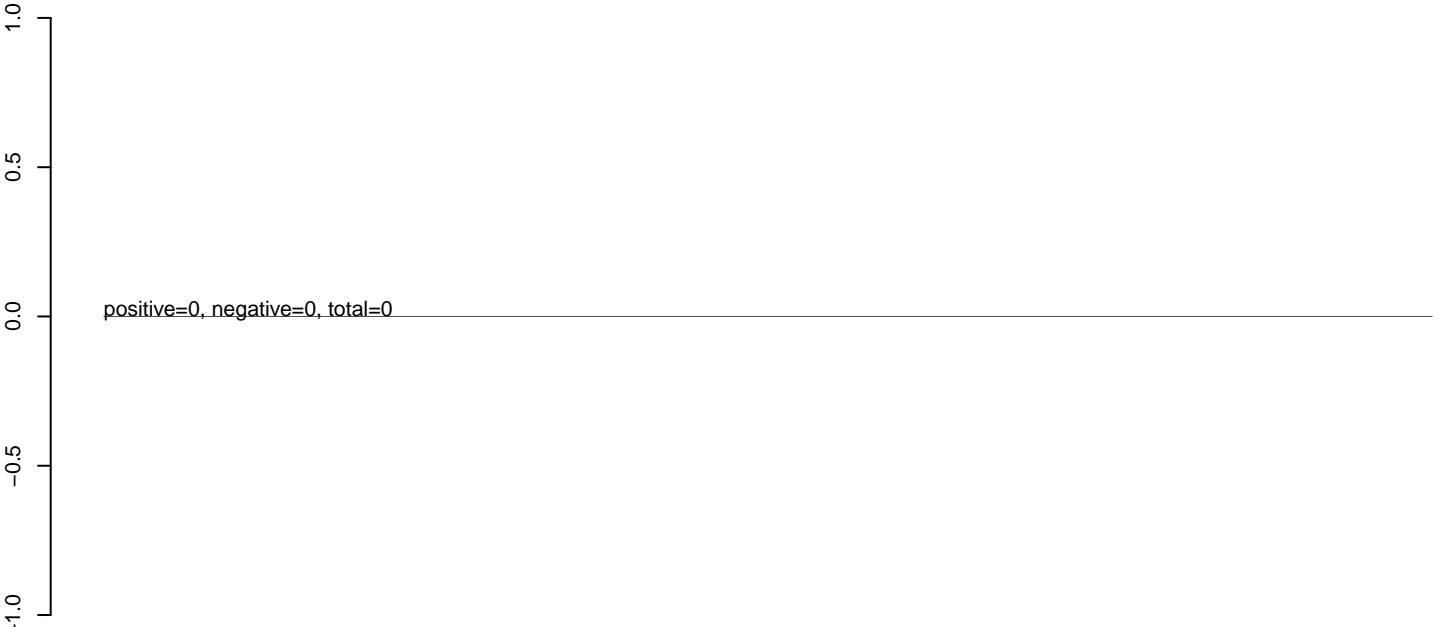
CuQuin_MG_7dpi3.24_35.rep



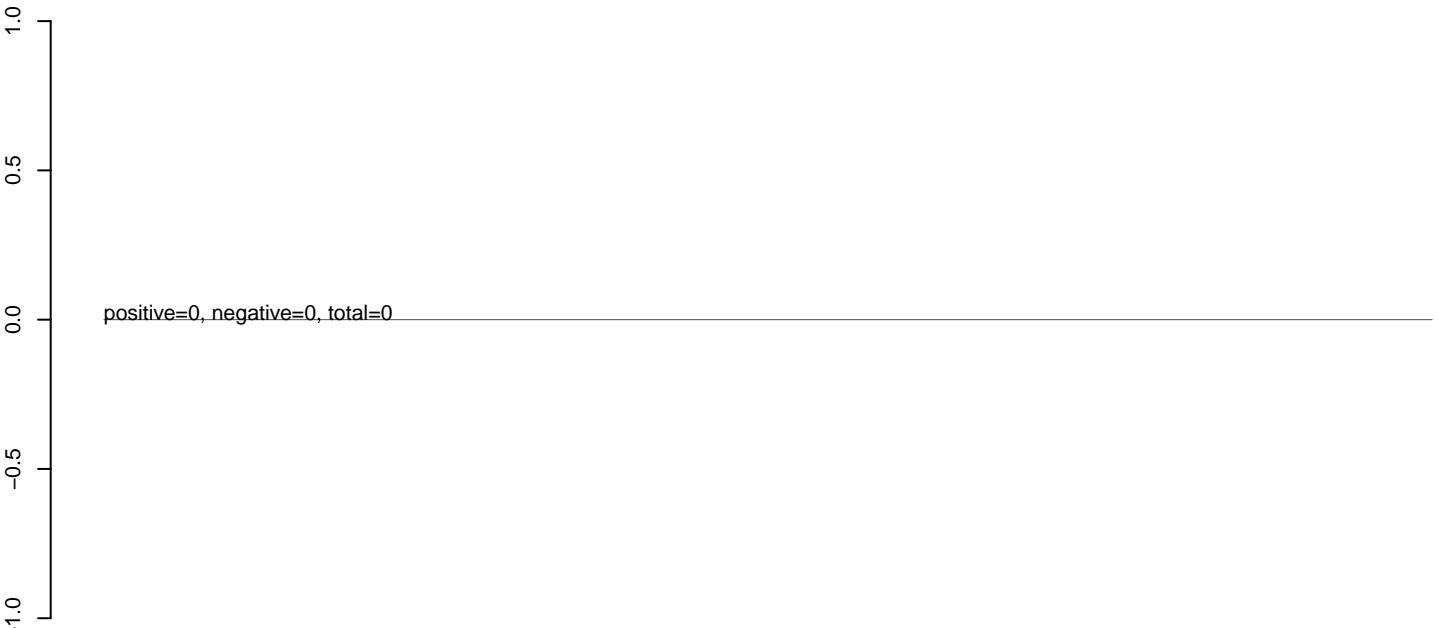
CuQuin_MG_7dpi3.rep



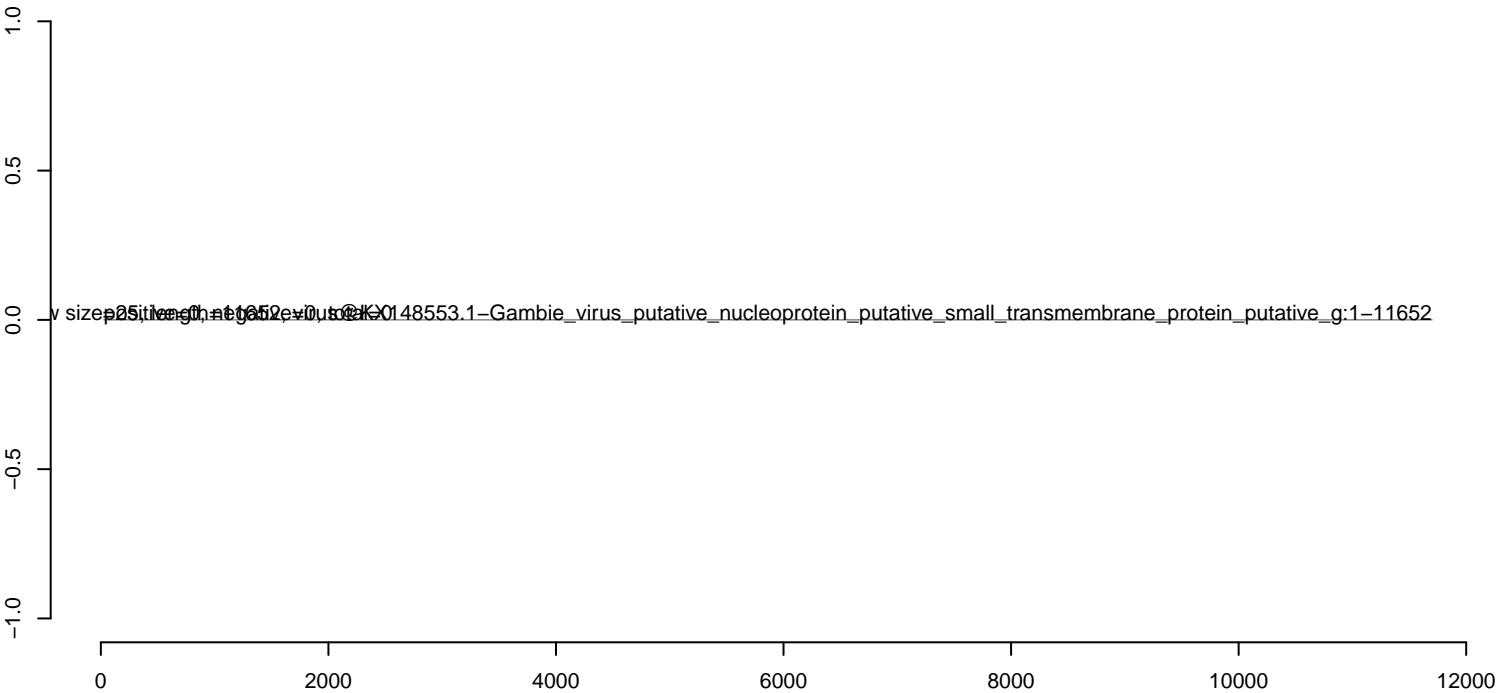
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

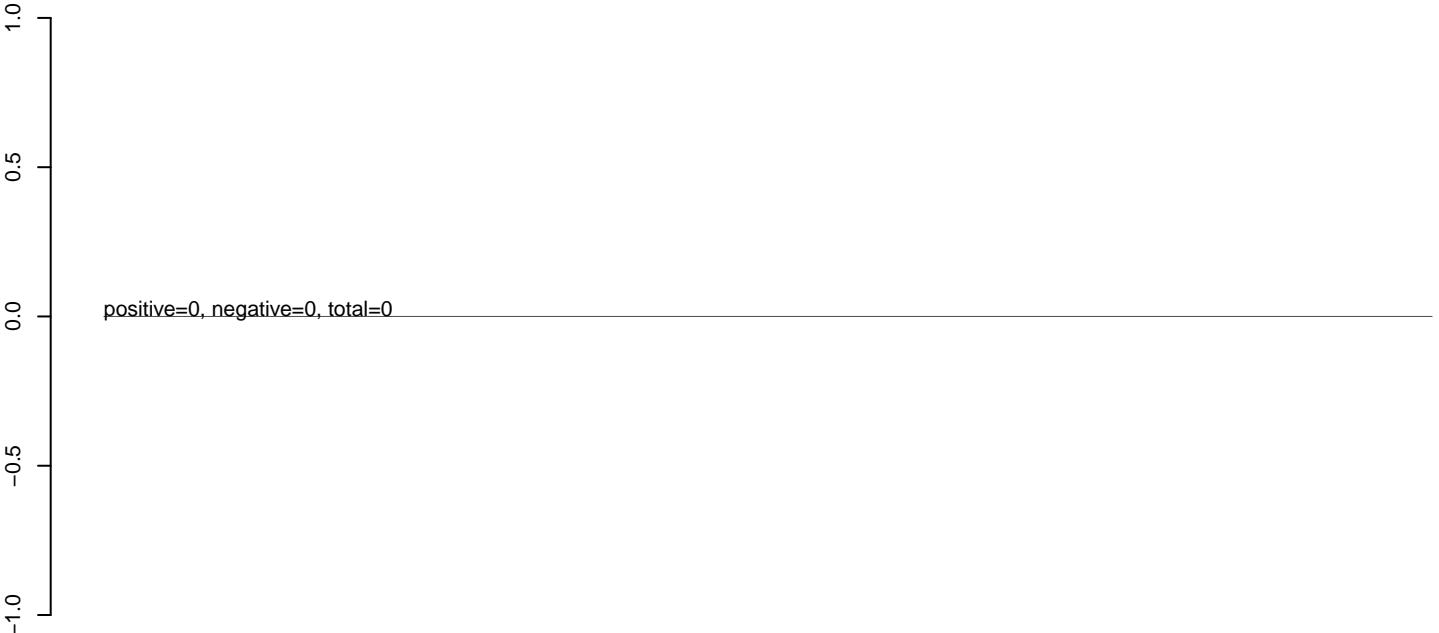
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

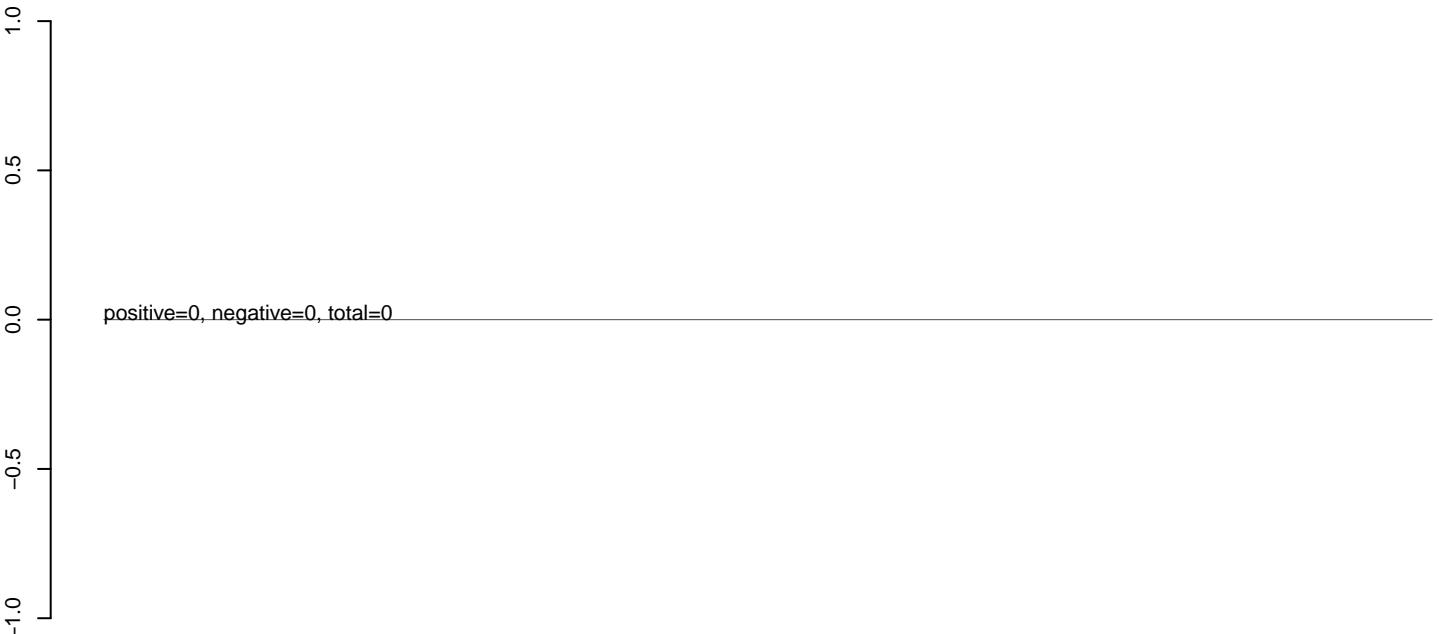
Window size=25, length=11798, virus@KU194360.1_Merida_virus_isolate_MERD-Mex07:1-11798

0 2000 4000 6000 8000 10000 12000

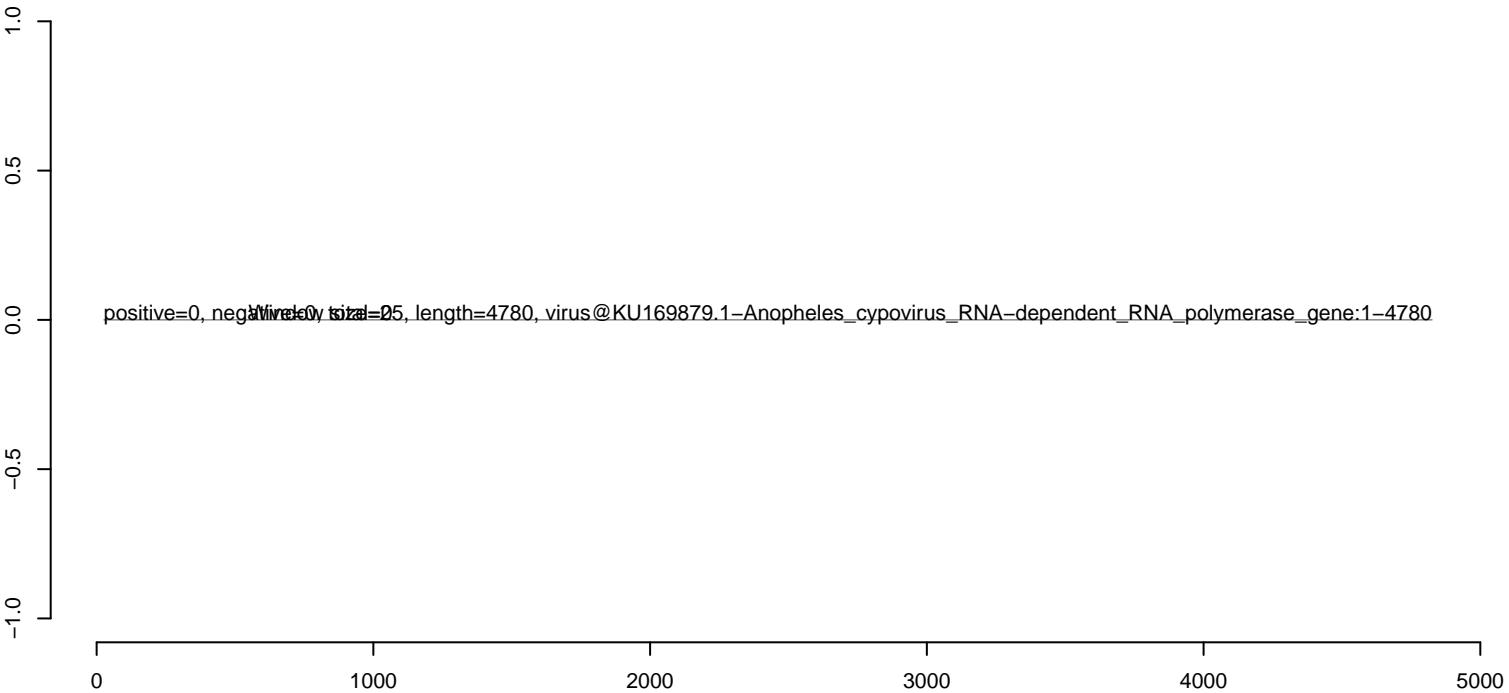
CuQuin_MG_7dpi3.18_23.rep



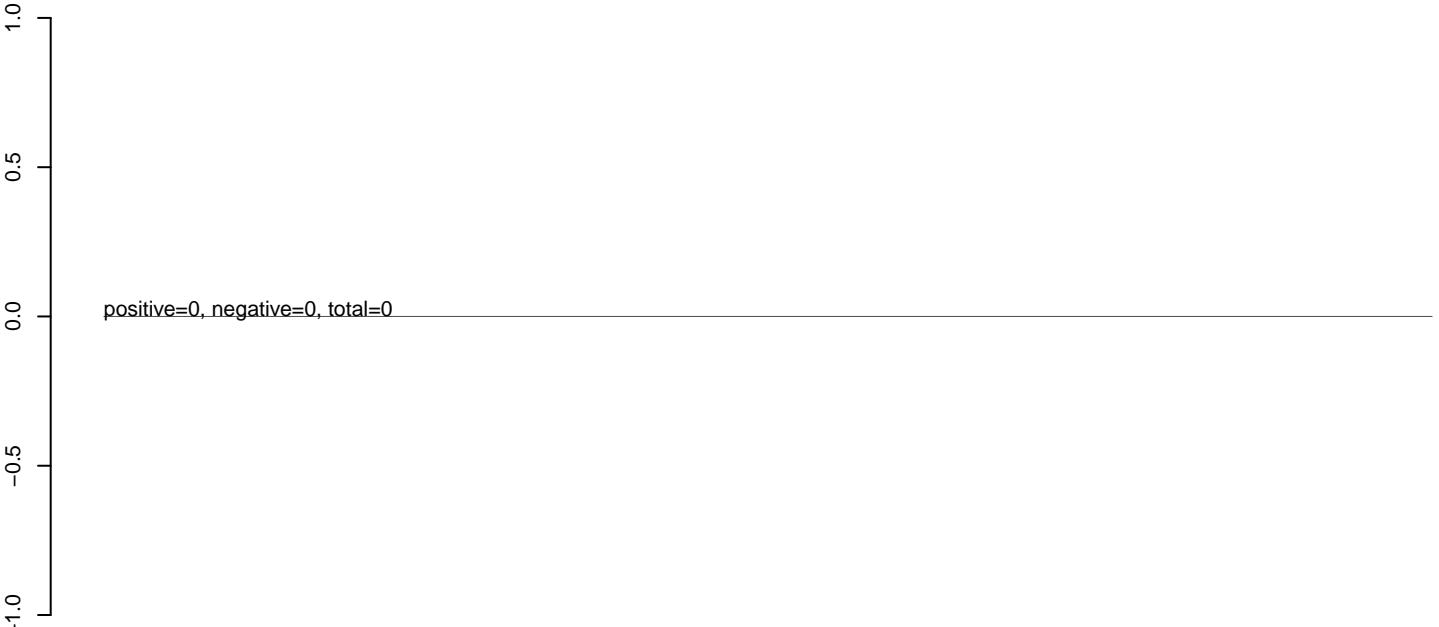
CuQuin_MG_7dpi3.24_35.rep



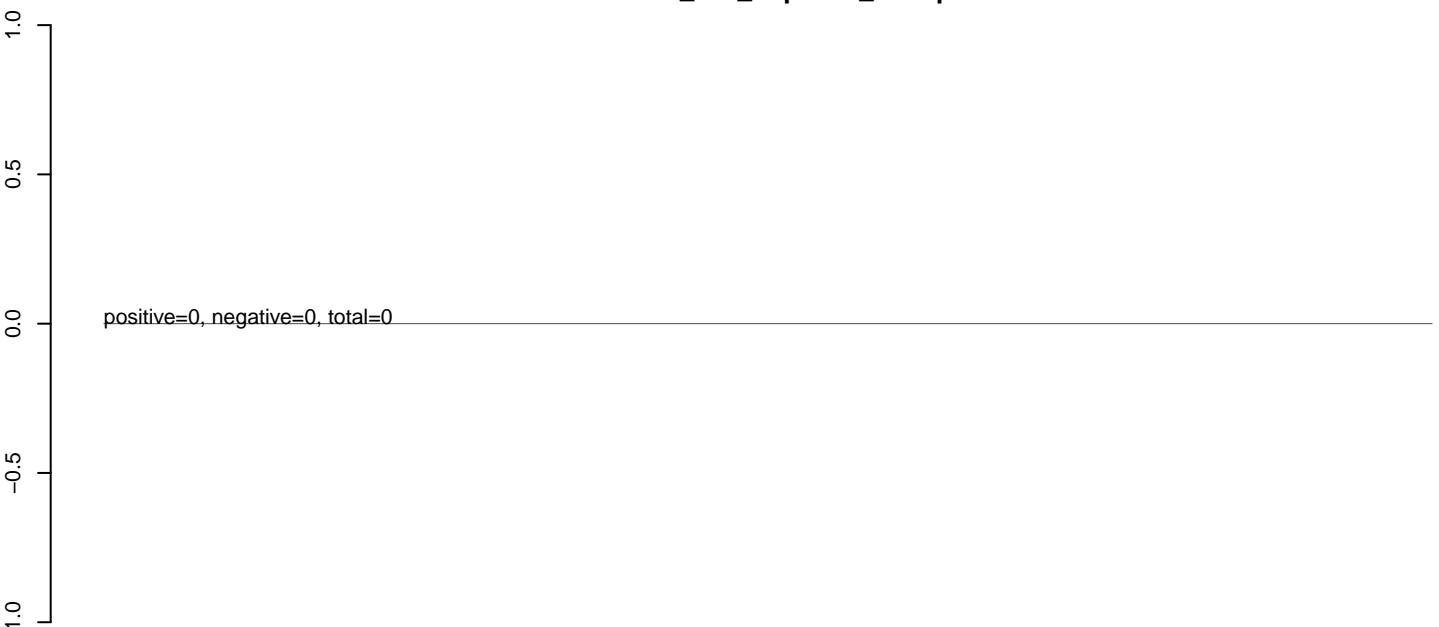
CuQuin_MG_7dpi3.rep



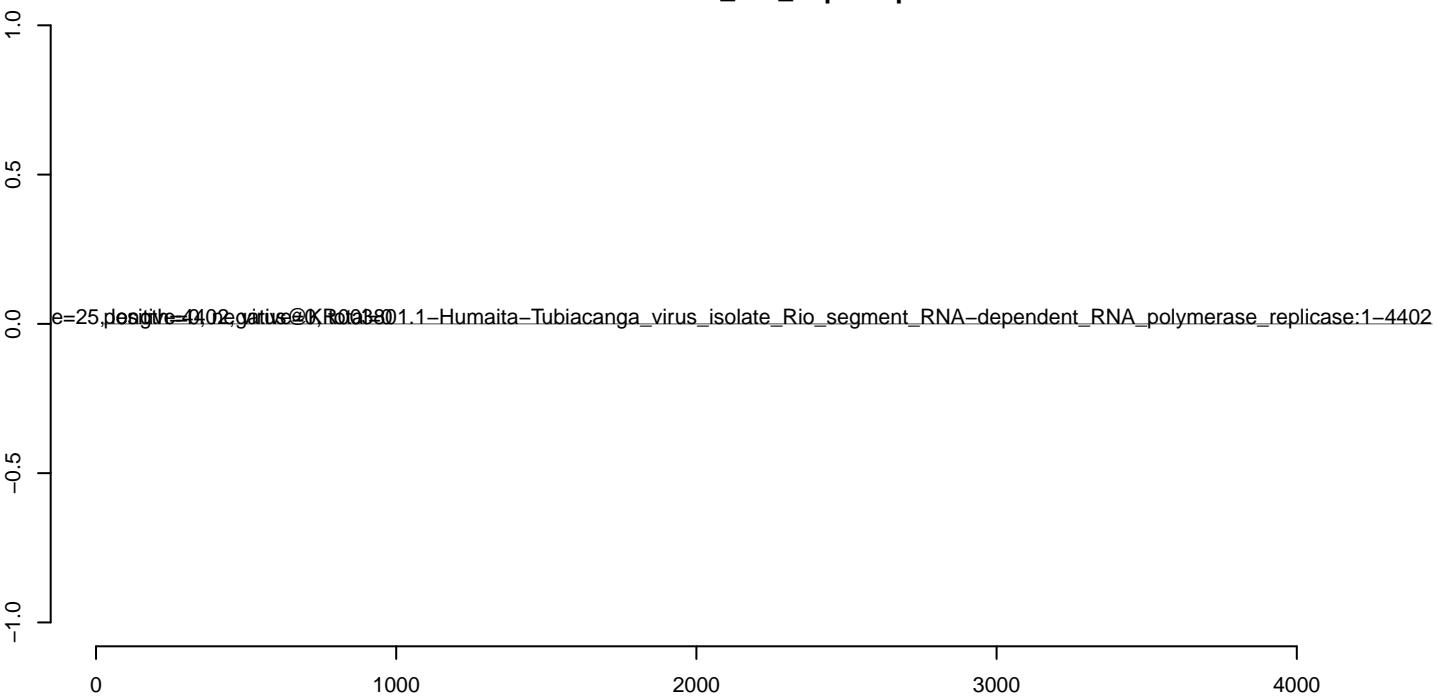
CuQuin_MG_7dpi3.18_23.rep



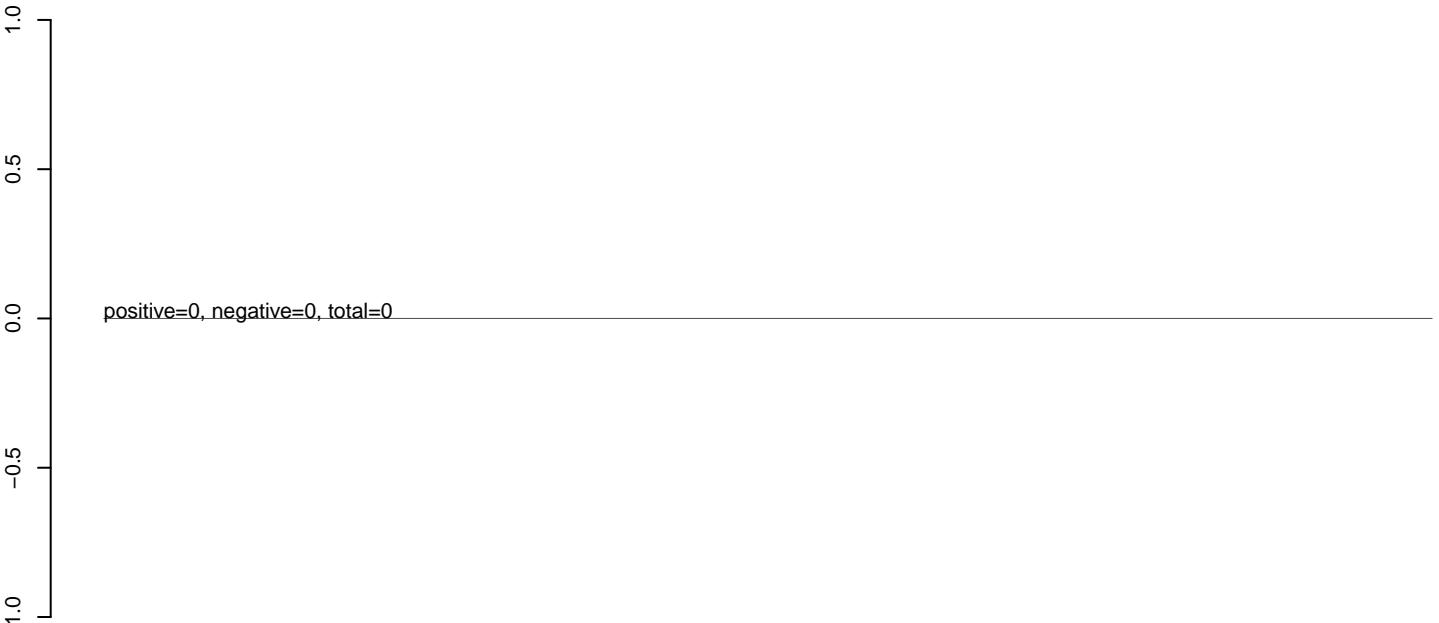
CuQuin_MG_7dpi3.24_35.rep



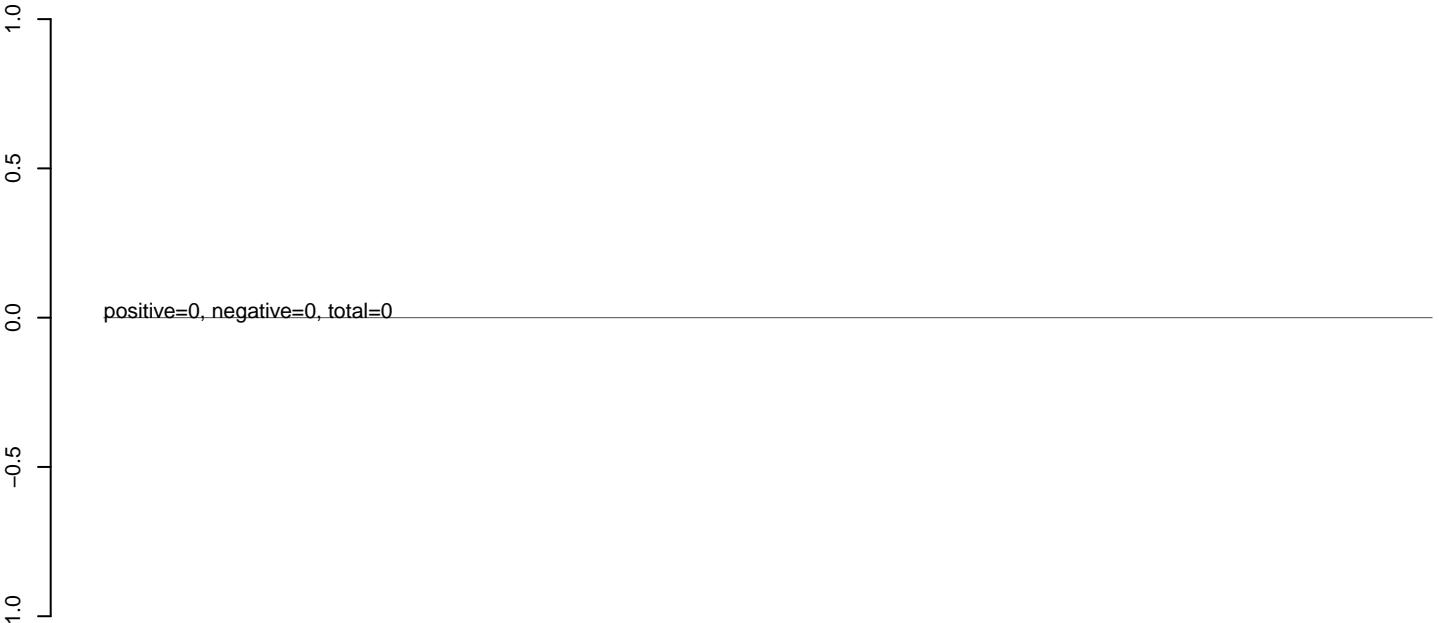
CuQuin_MG_7dpi3.rep



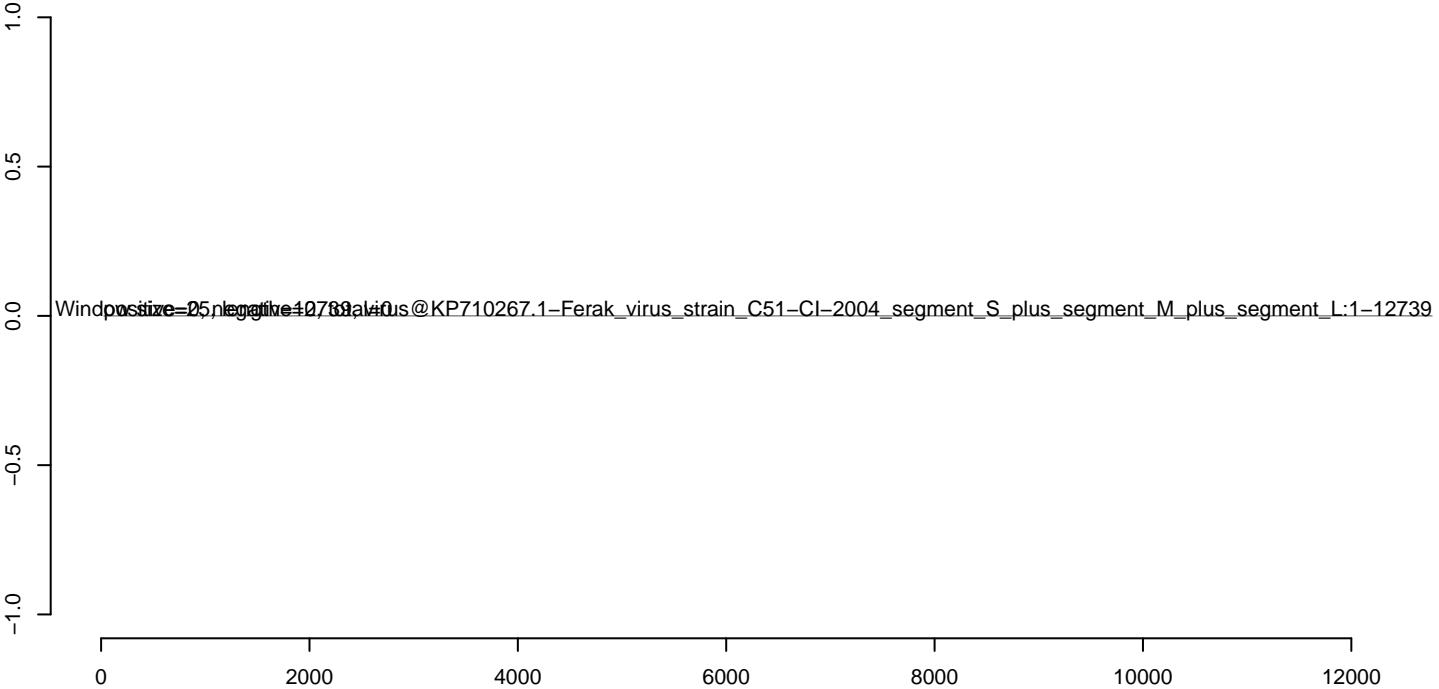
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

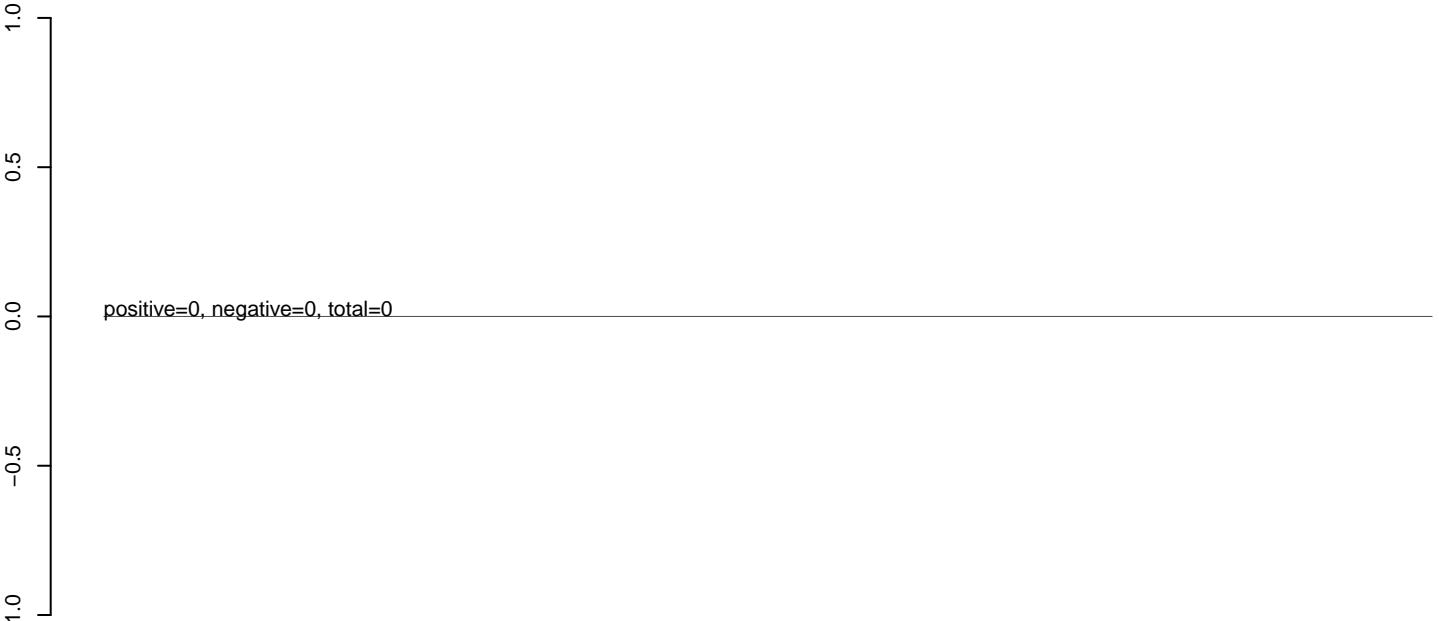
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

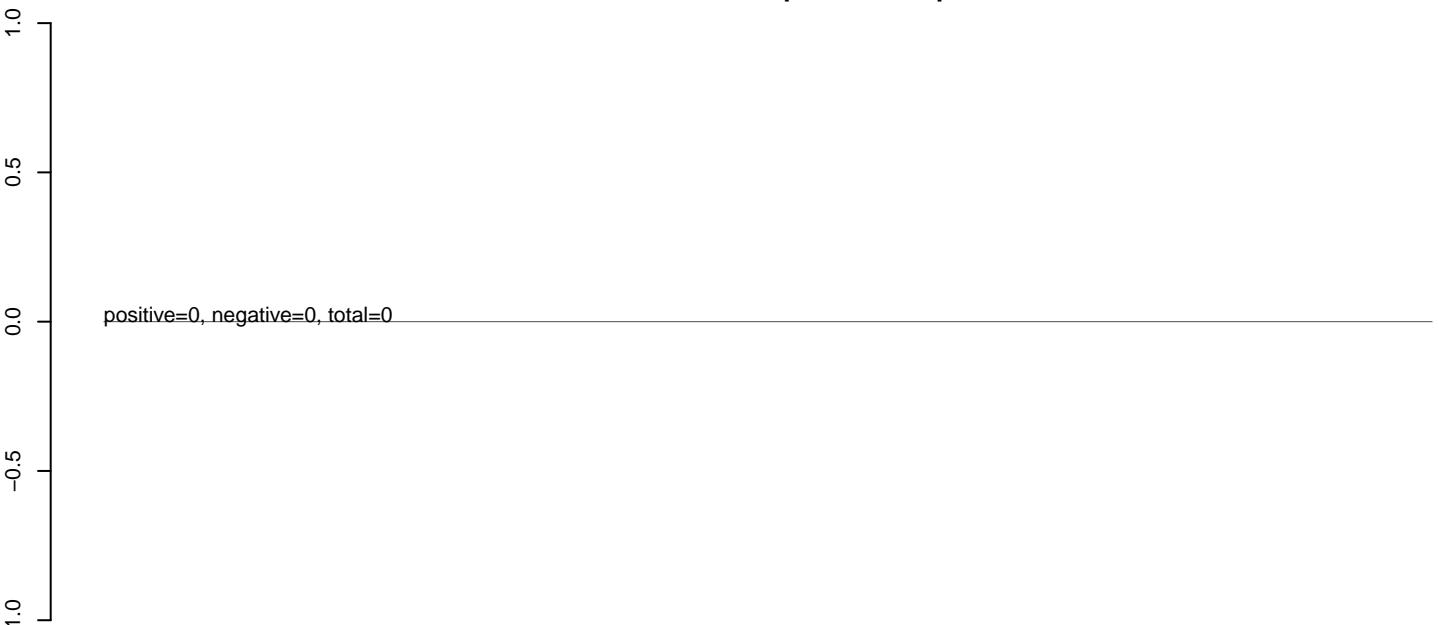
Window size=25, length=10560, virus@KM225263.1–Stratford_virus_strain_C338:1–10560

0 2000 4000 6000 8000 10000

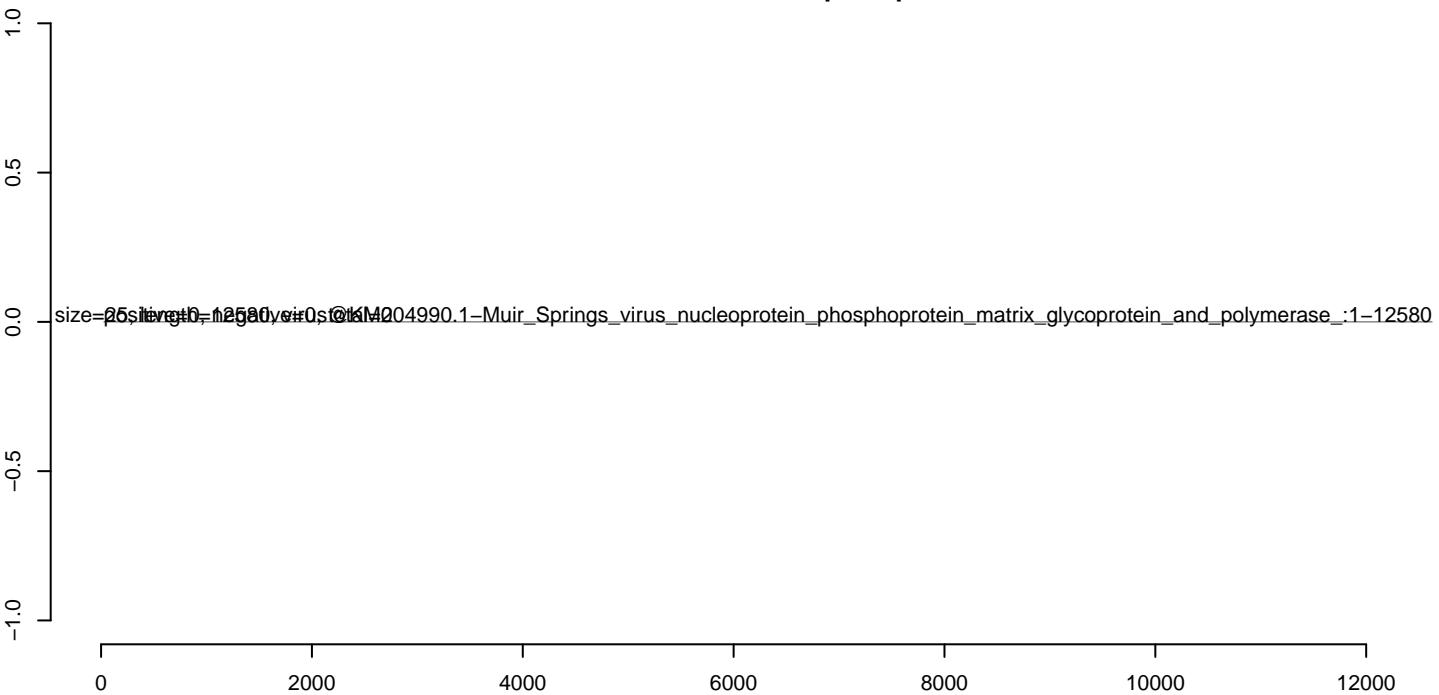
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

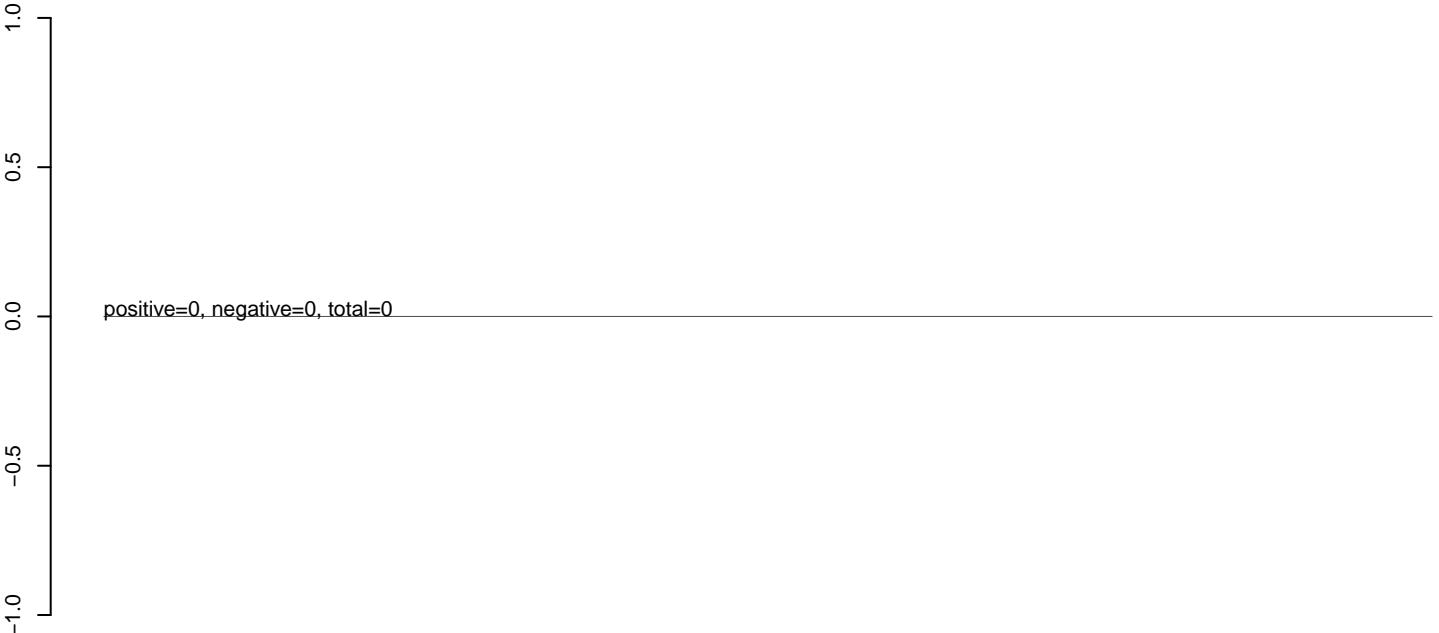
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

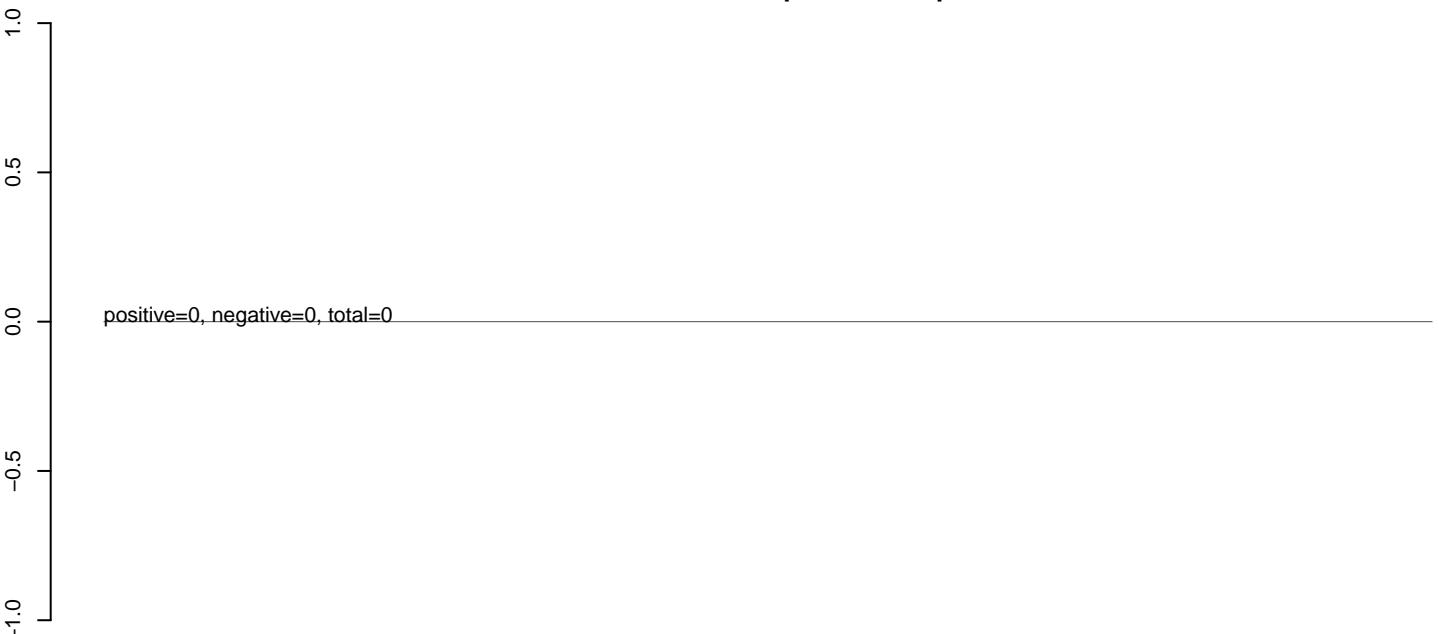
positive=0, negative=0, size=26, length=4773, virus@KJ632942.1-Mosinovirus_strain_C36_Cl-2004_segment_RNA_1_and_RNA_2:1-4773

0 1000 2000 3000 4000

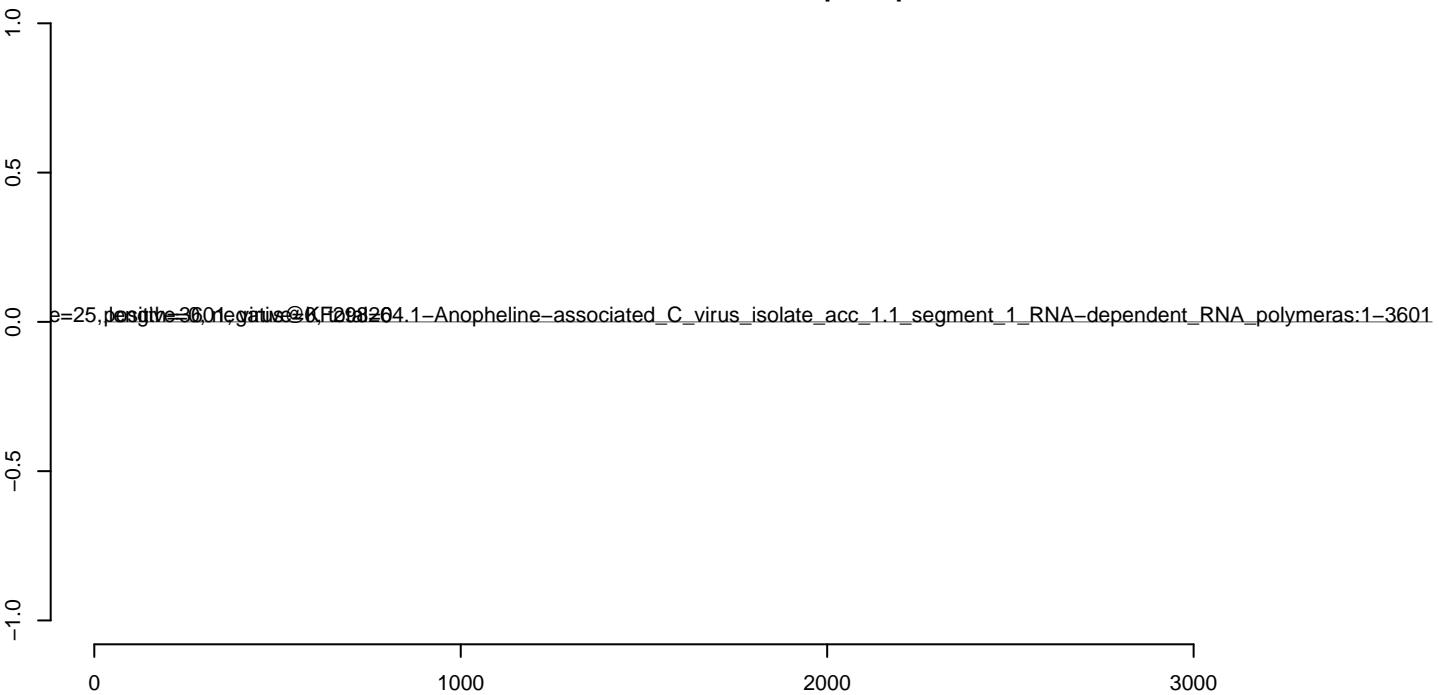
CuQuin_MG_7dpi3.18_23.rep



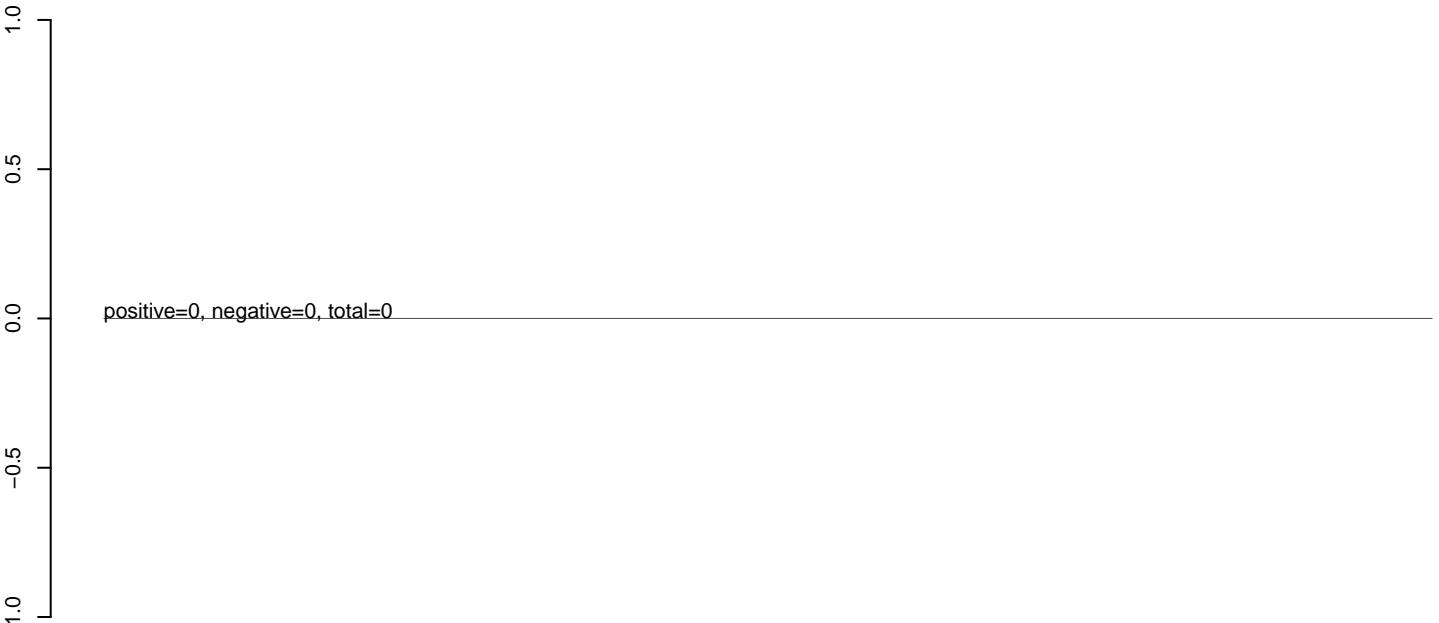
CuQuin_MG_7dpi3.24_35.rep



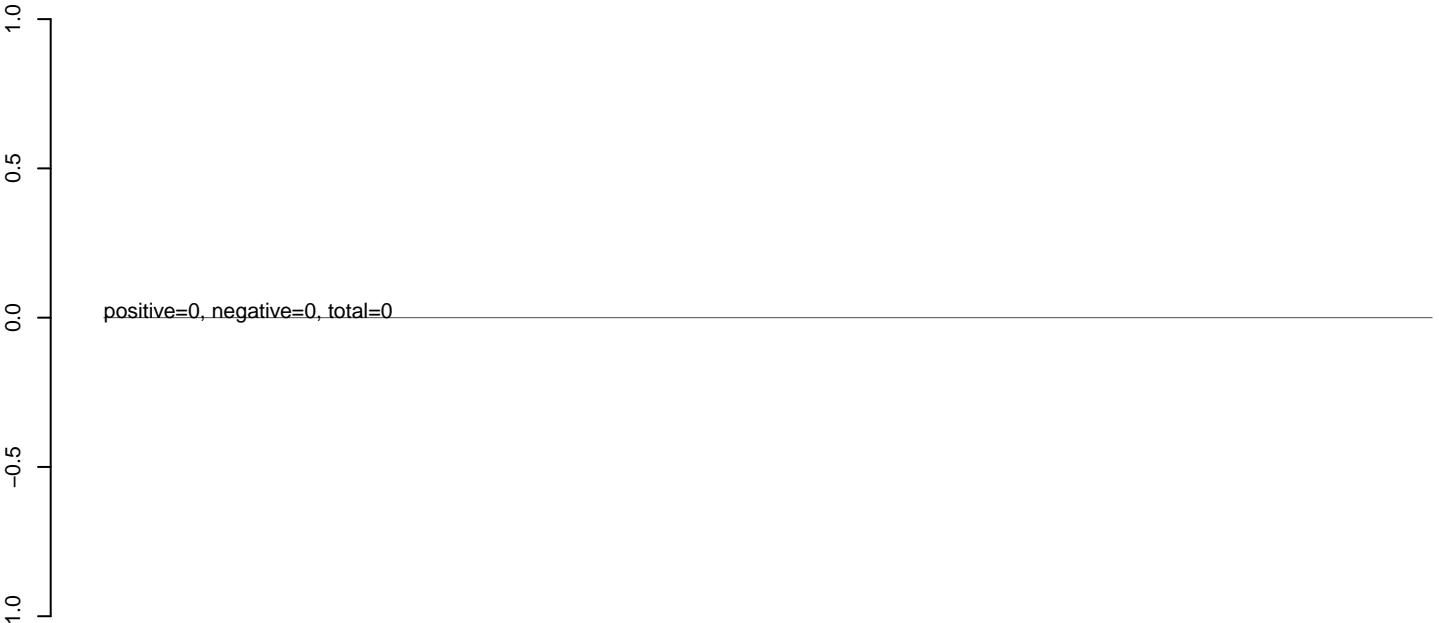
CuQuin_MG_7dpi3.rep



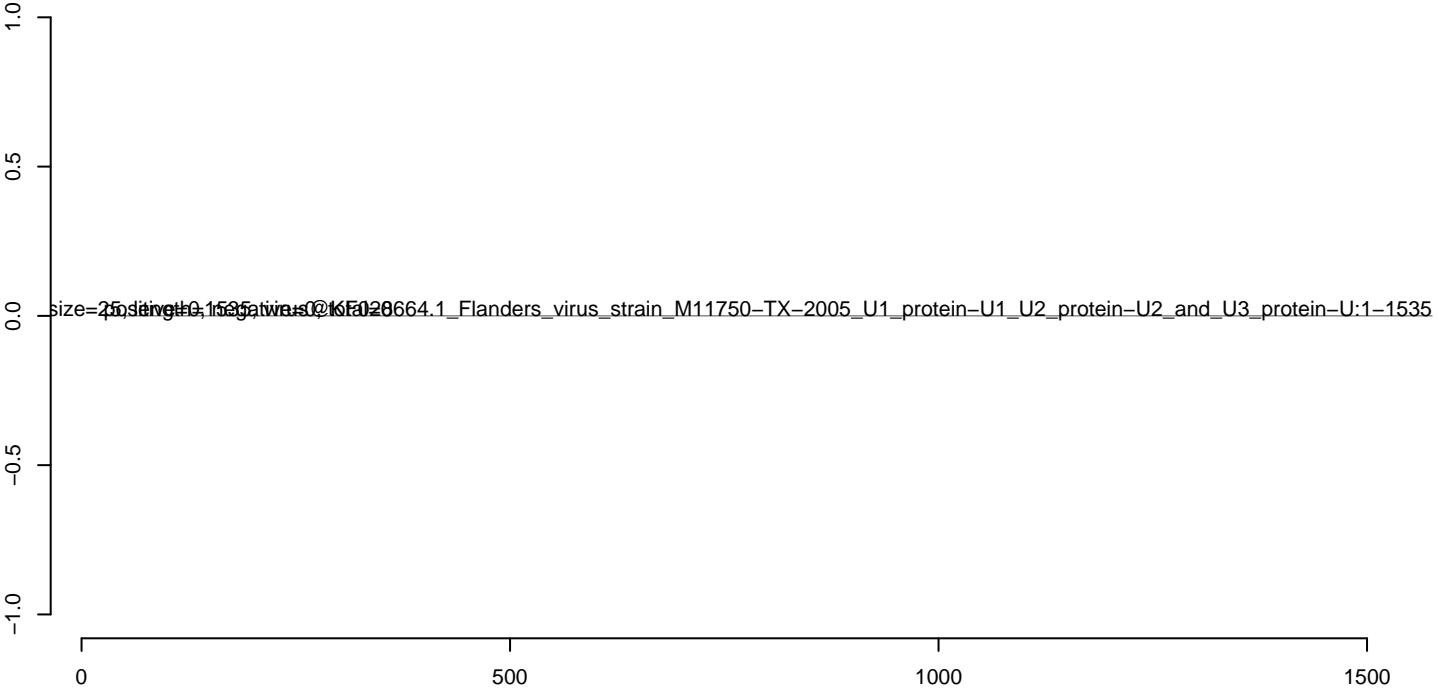
CuQuin_MG_7dpi3.18_23.rep



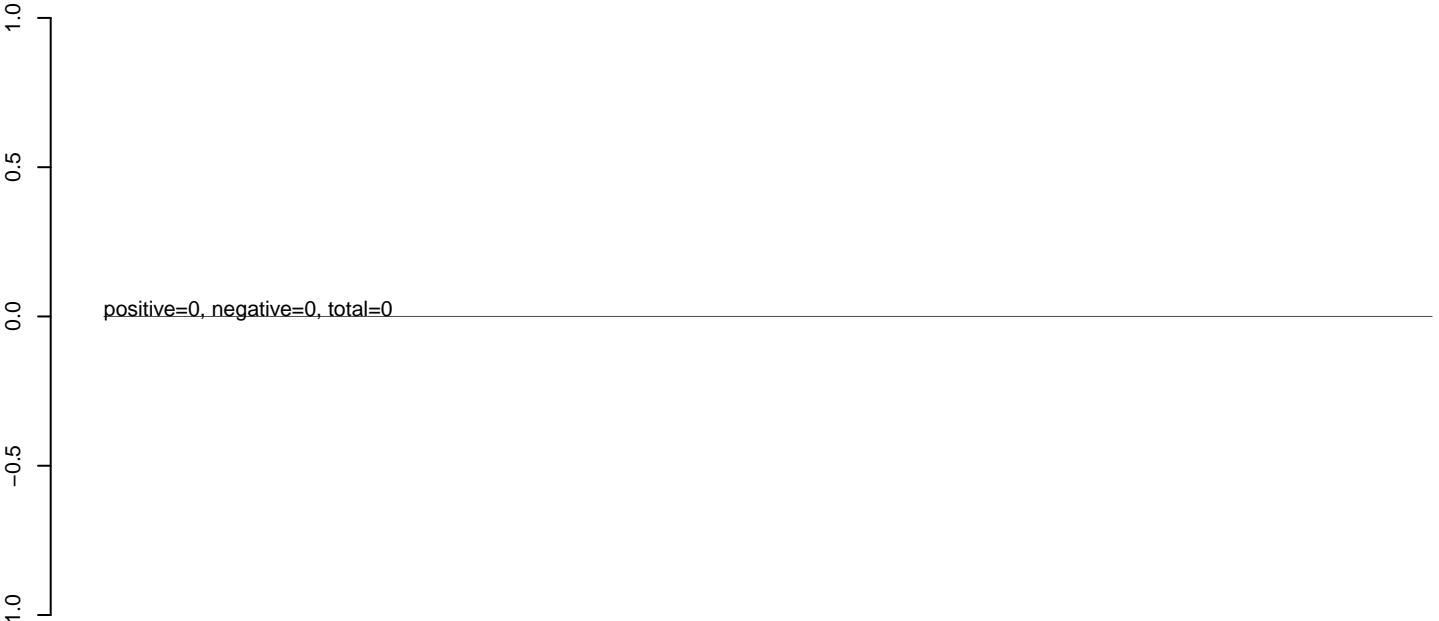
CuQuin_MG_7dpi3.24_35.rep



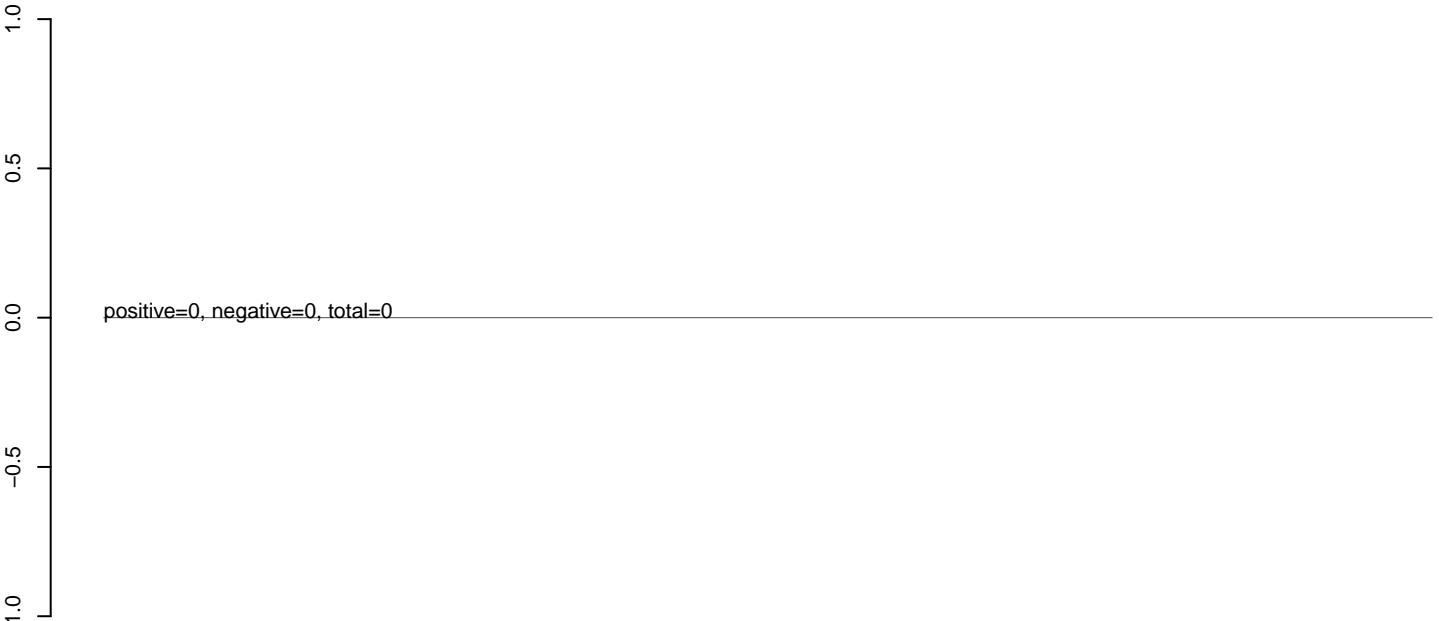
CuQuin_MG_7dpi3.rep



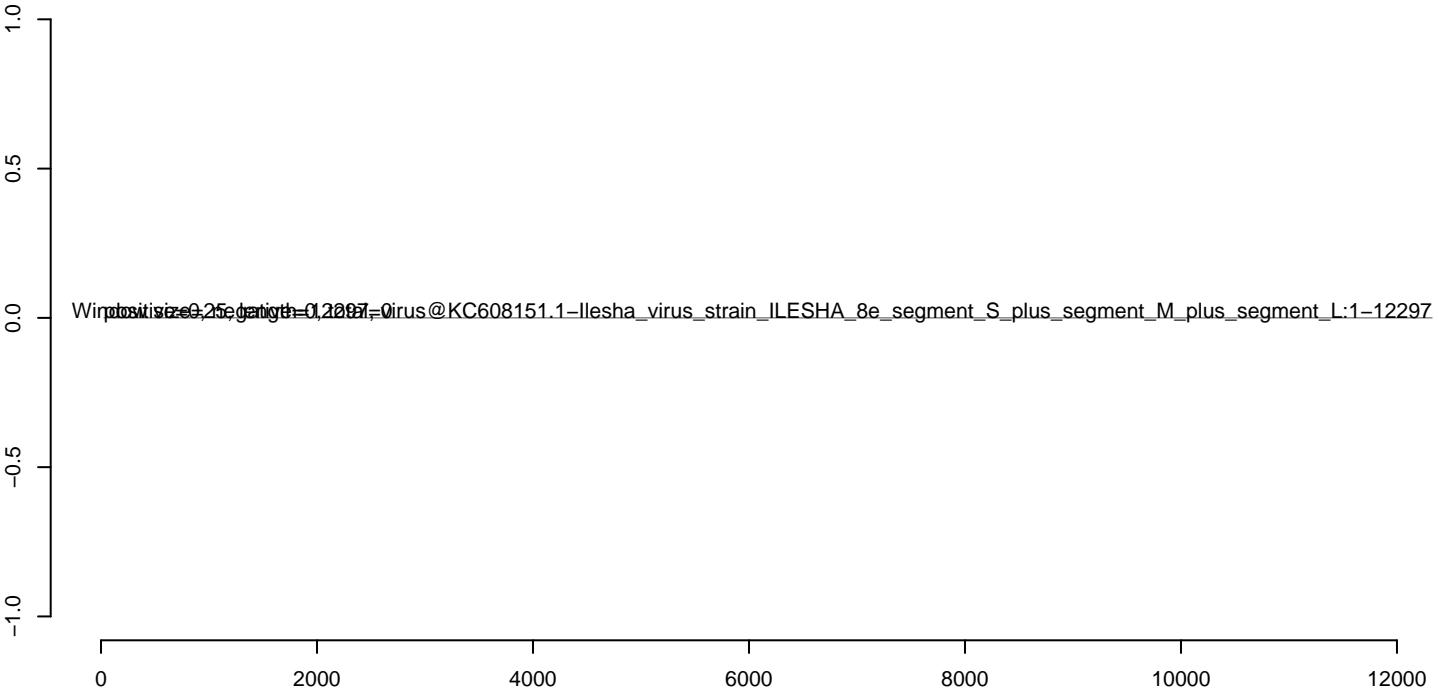
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

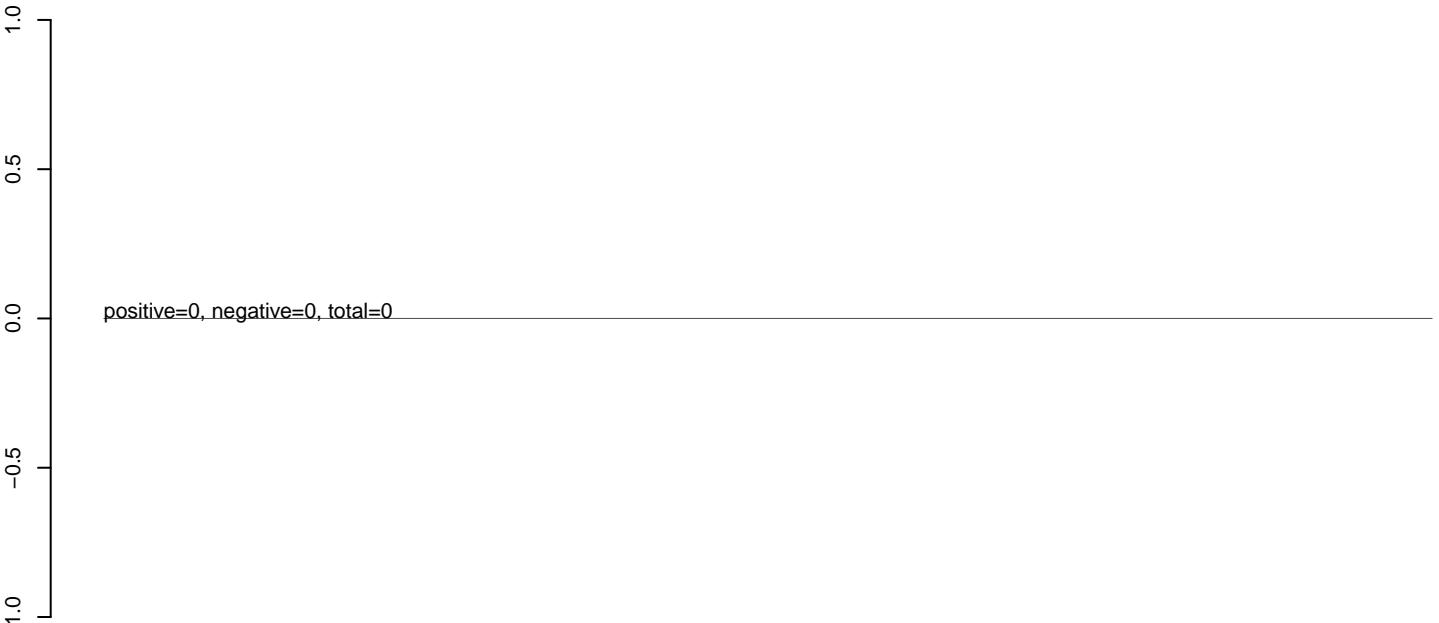
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

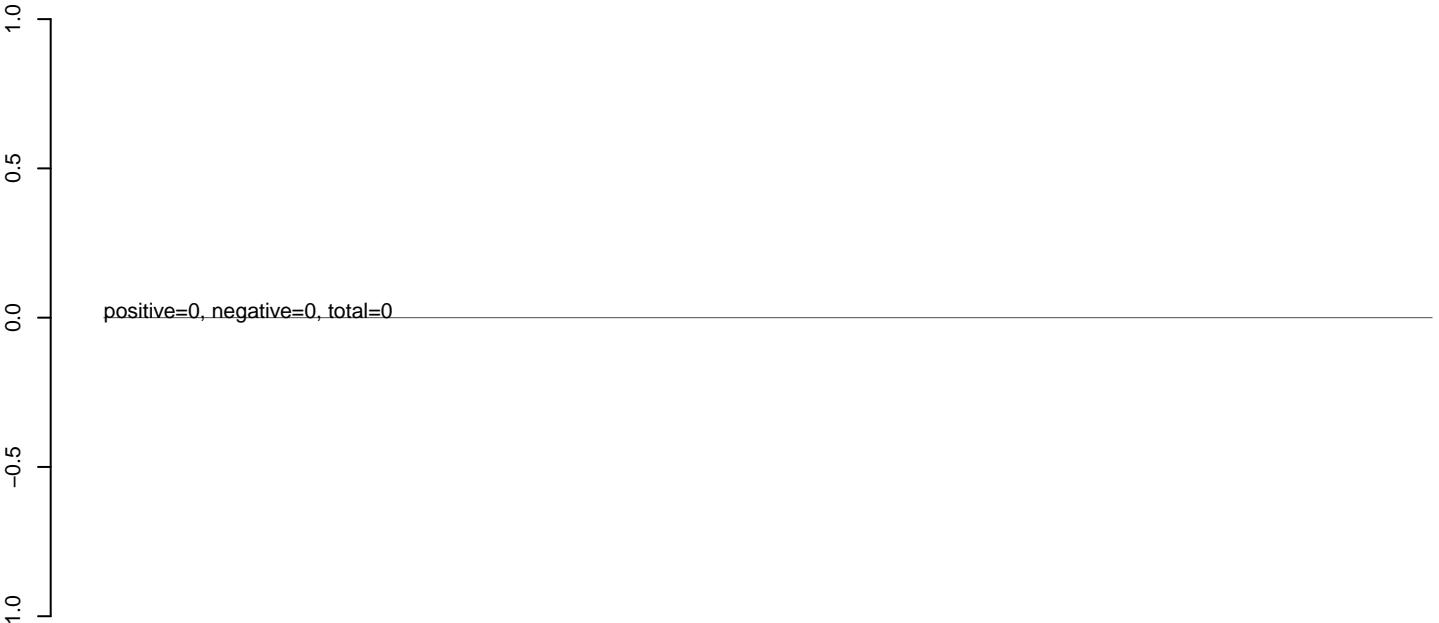
positive=0, negative=0, windowSize=25, length=11079, virus@KC181923.1-Aedes_flavivirus_strain_AEFV-SPFLD-MO-2011-MP6:1-11079

0 2000 4000 6000 8000 10000

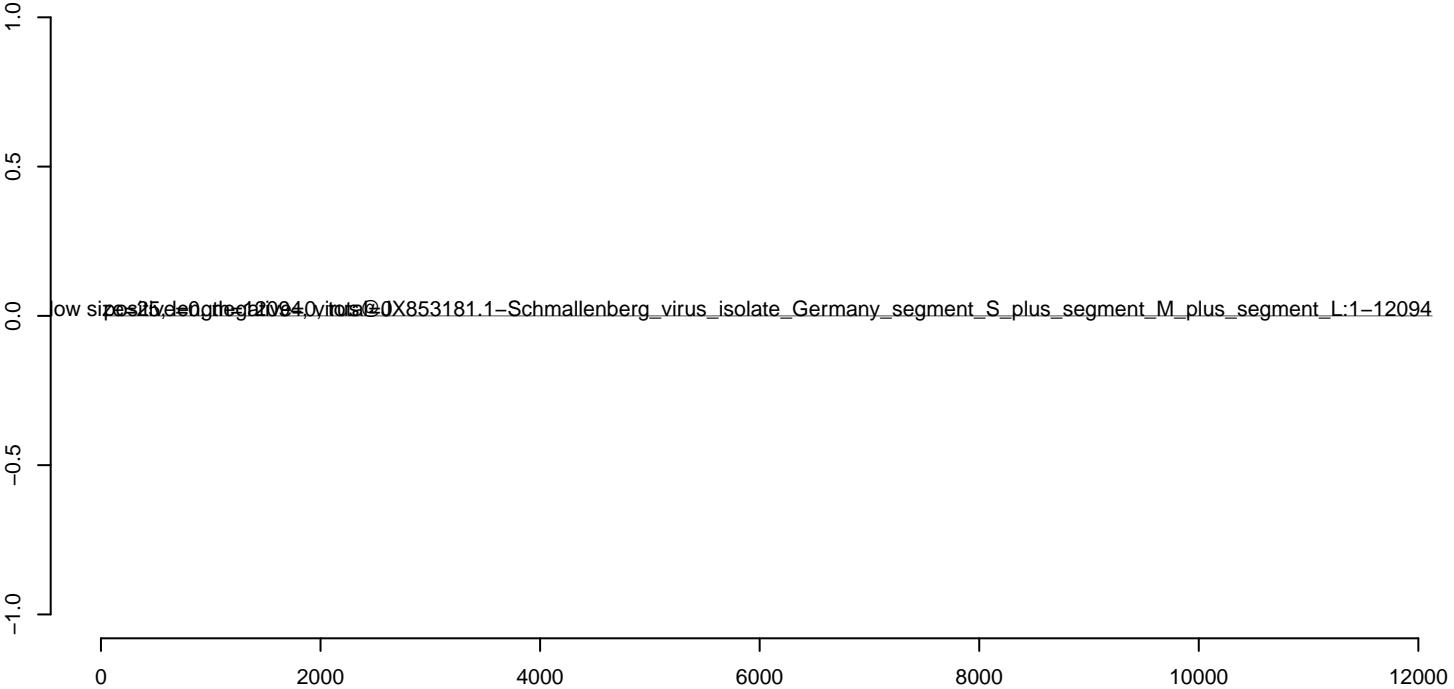
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

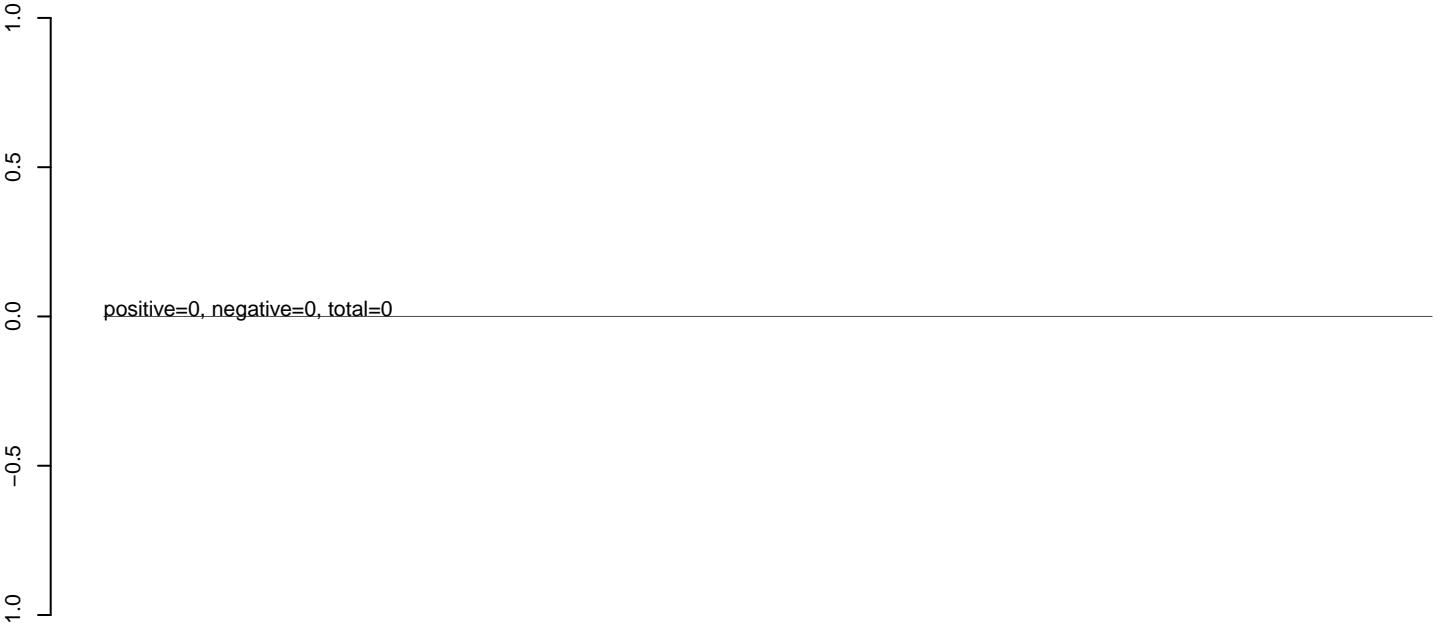
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

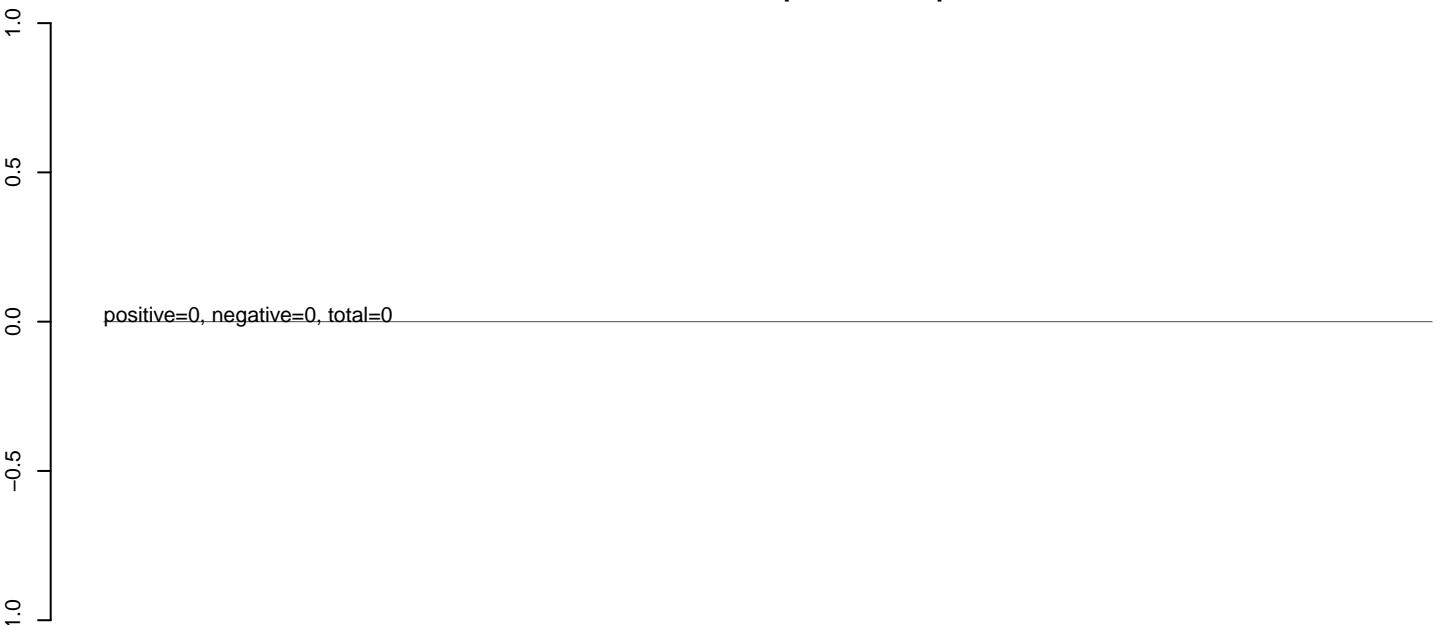
Window size=25, length=10841, virus@JQ308187.1–Culex_flavivirus_isolate_HLD105:1–10841

0 2000 4000 6000 8000 10000

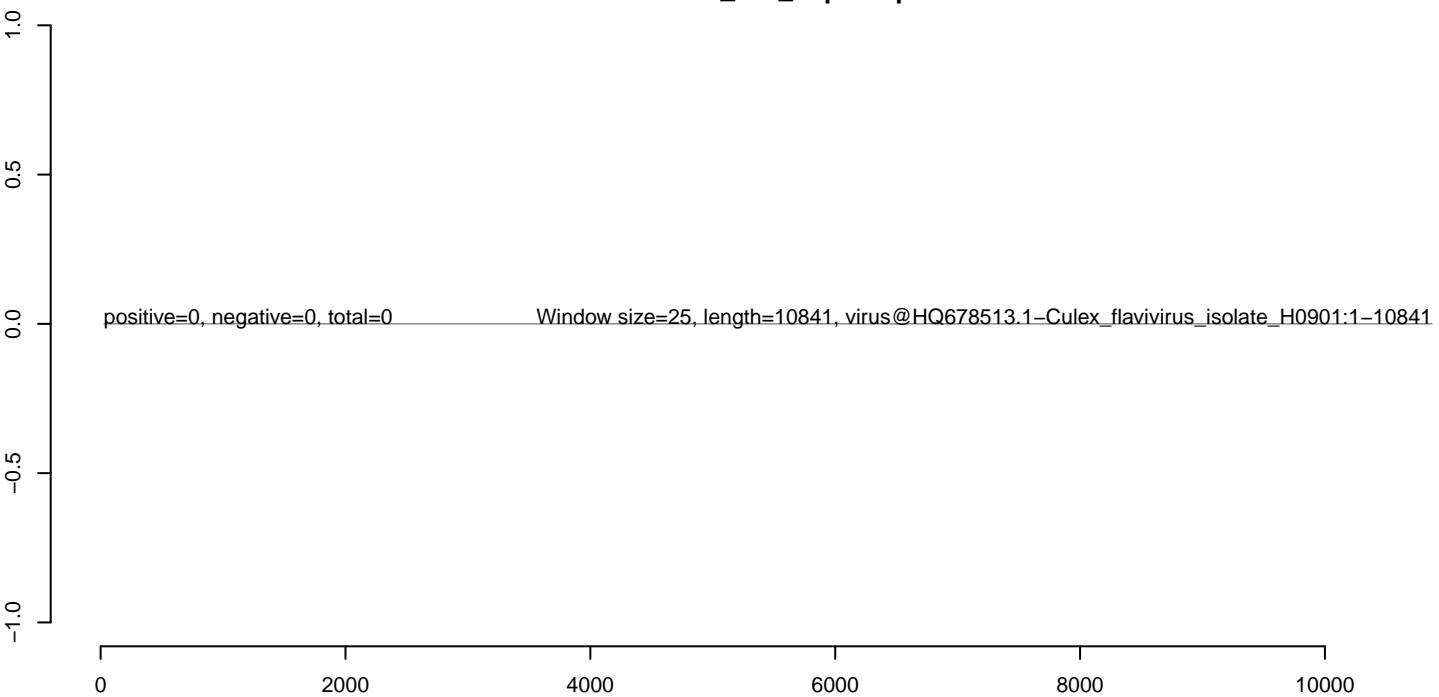
CuQuin_MG_7dpi3.18_23.rep



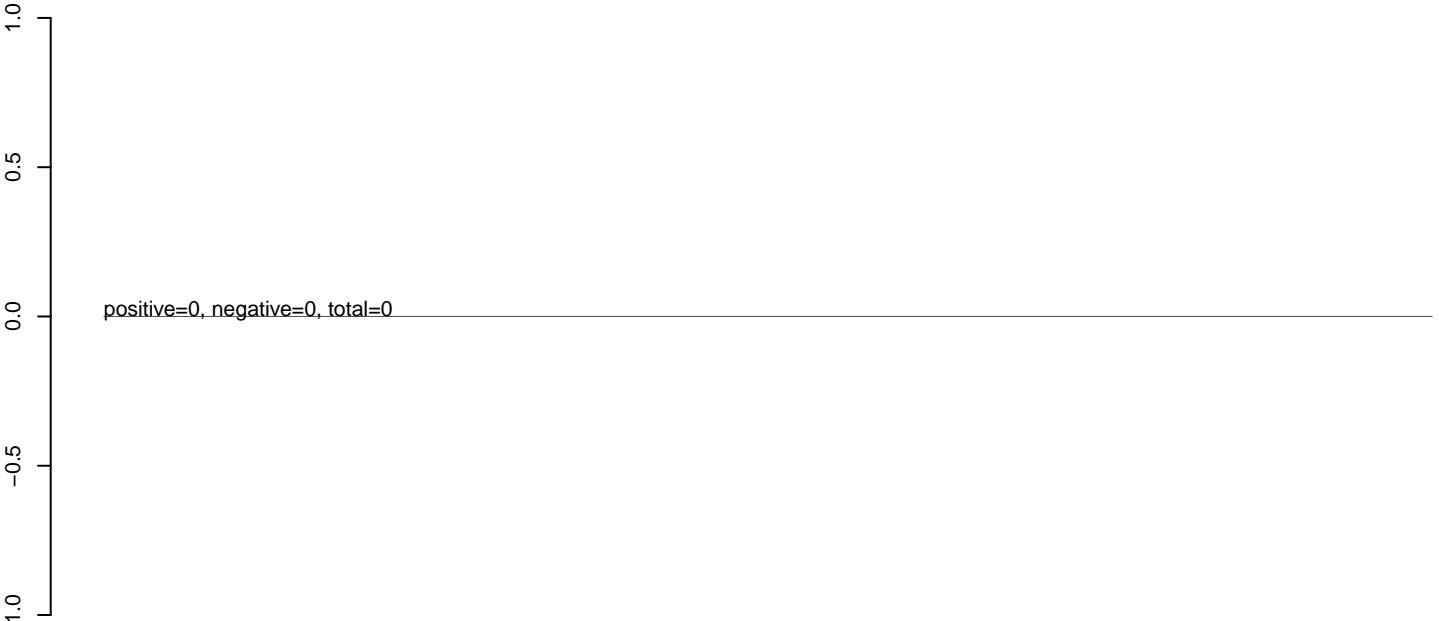
CuQuin_MG_7dpi3.24_35.rep



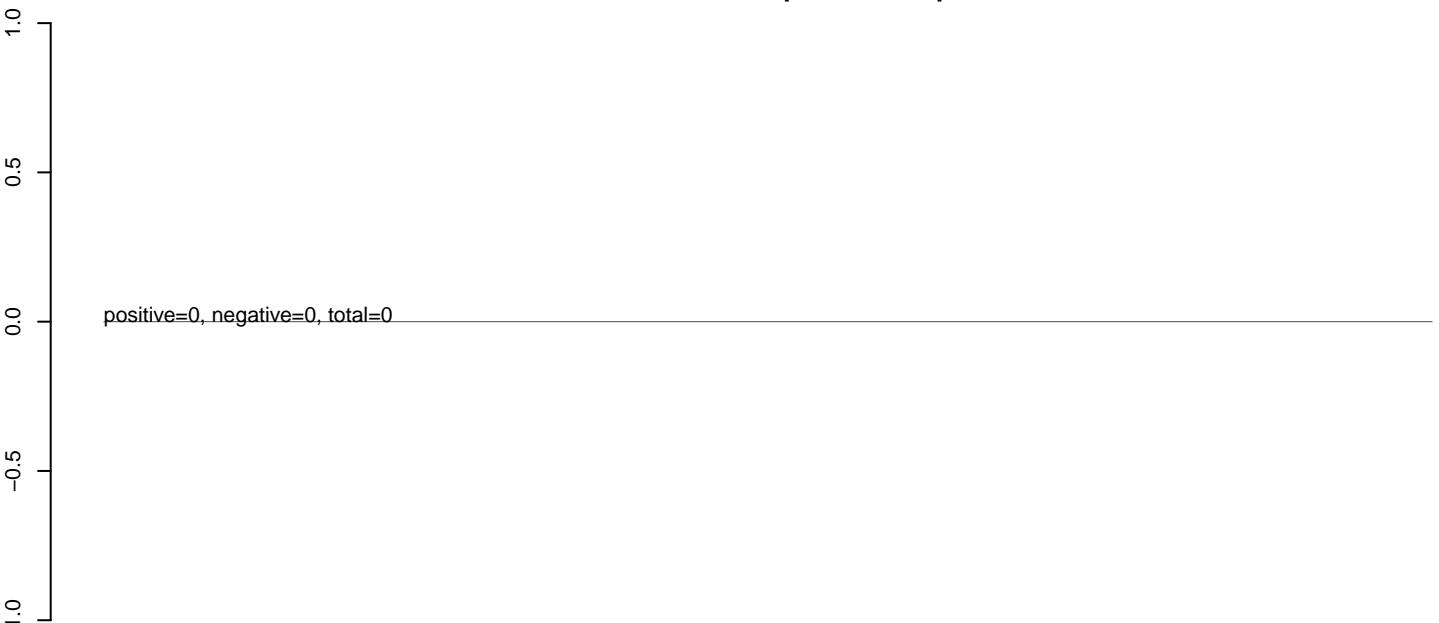
CuQuin_MG_7dpi3.rep



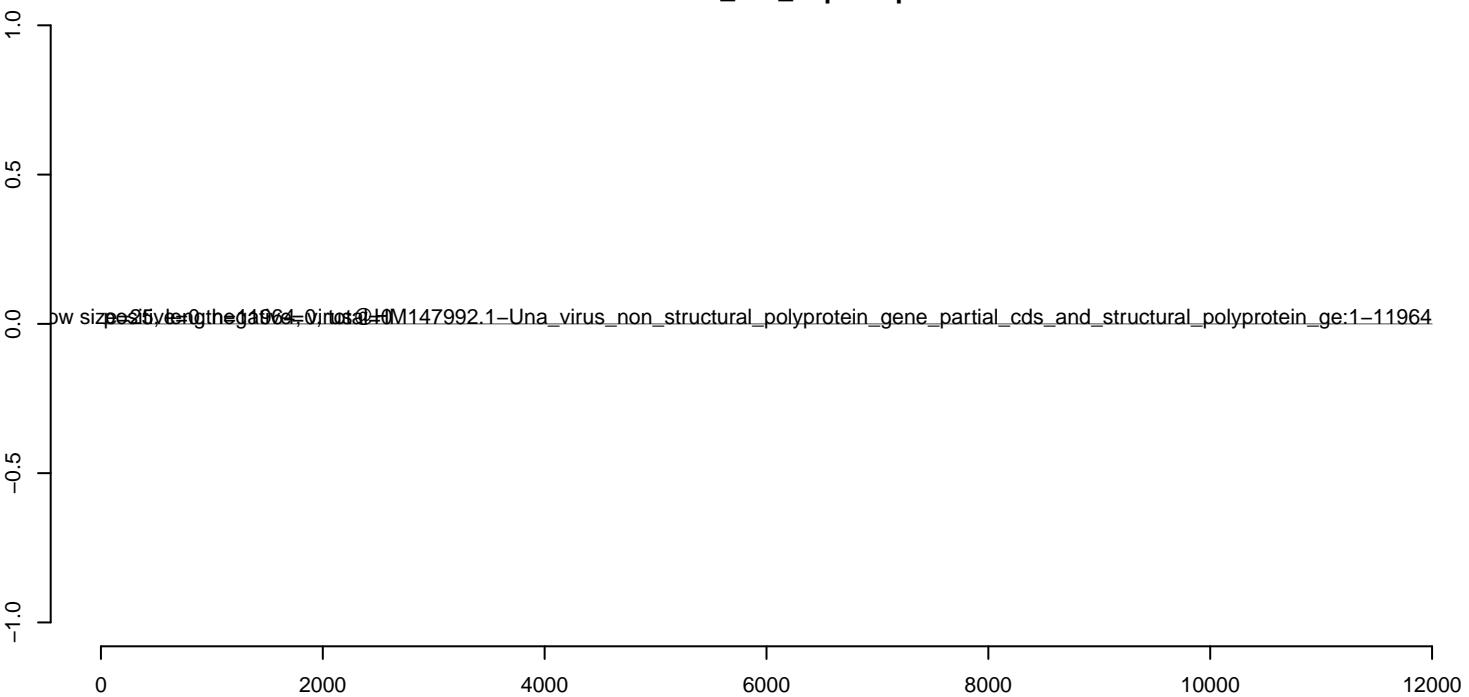
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

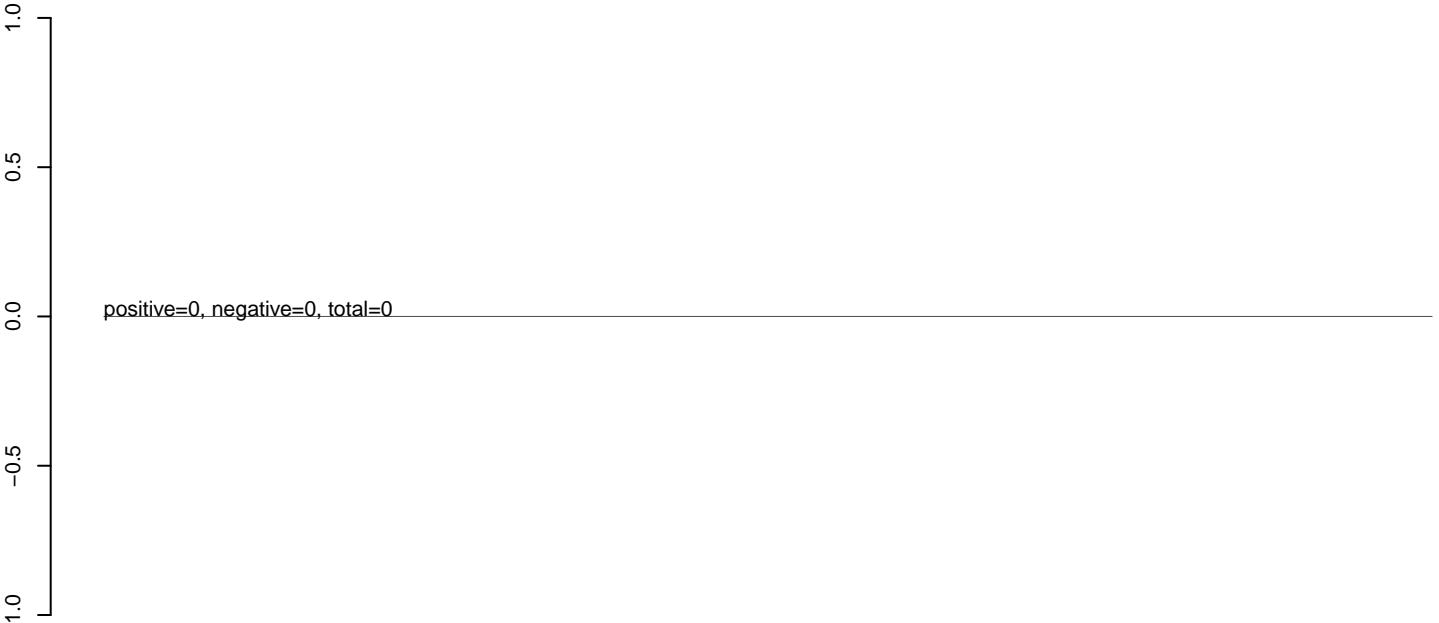
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

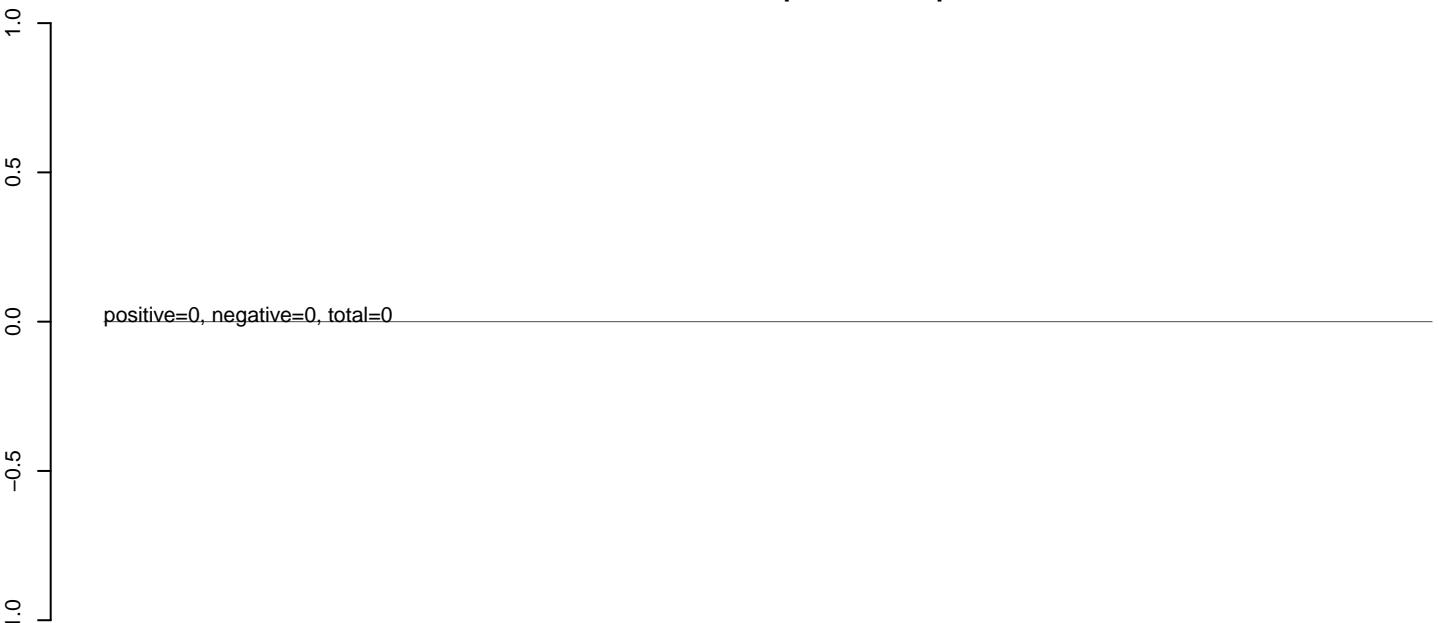
positive=0, negative=0, total=0, window size=25, length=10774, virus@HM055369.1-Kyasanur_forest_disease_virus_strain_KFD_P_9605:1-10774

0 2000 4000 6000 8000 10000

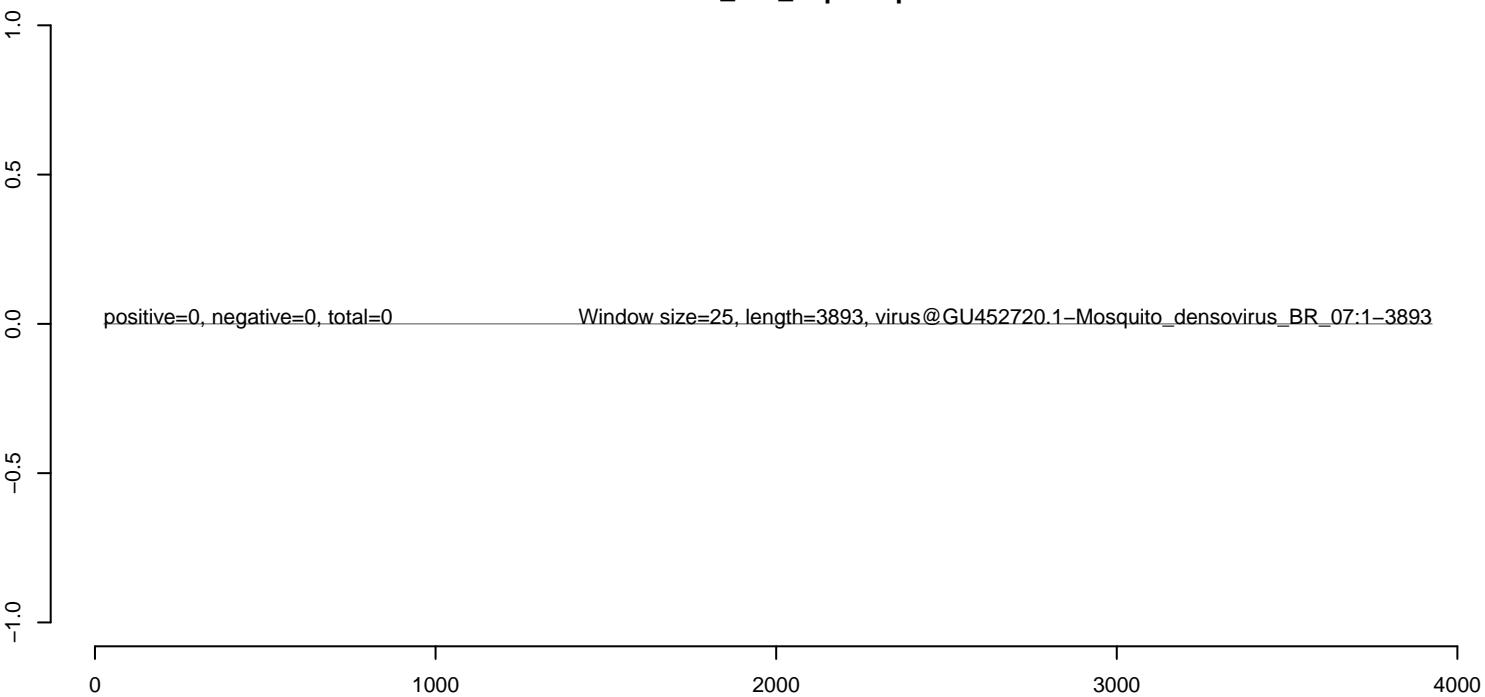
CuQuin_MG_7dpi3.18_23.rep



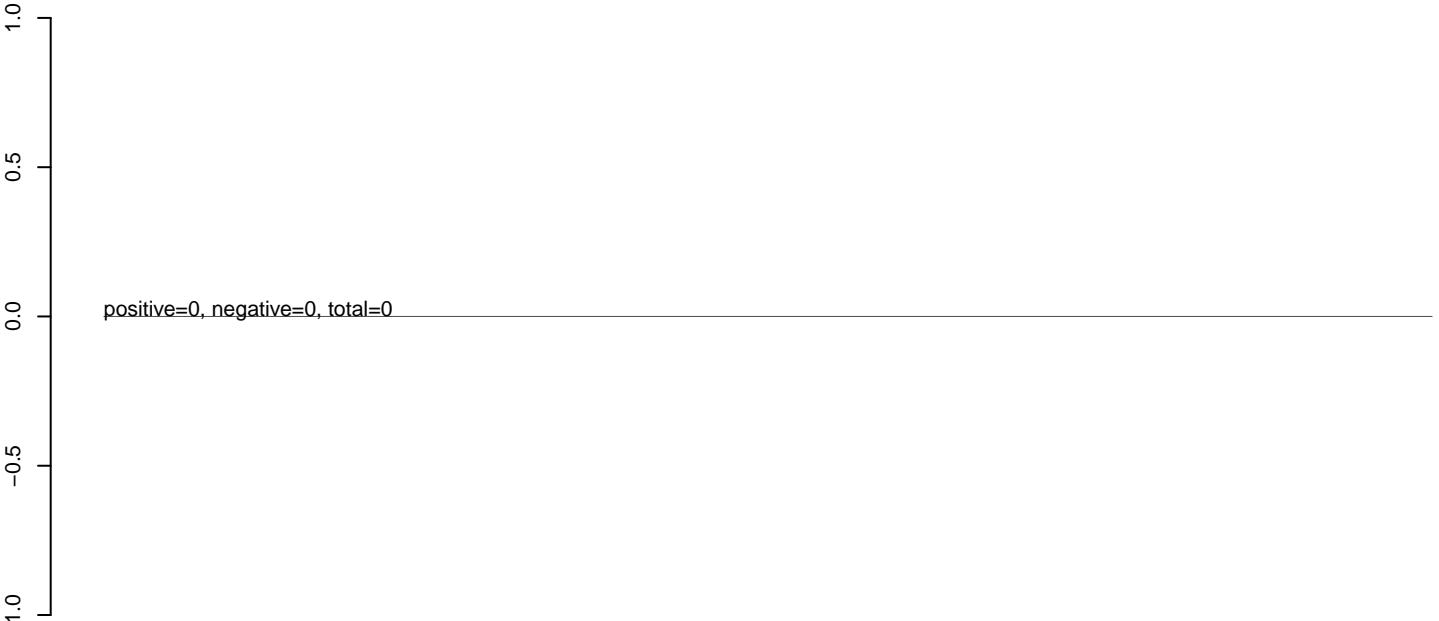
CuQuin_MG_7dpi3.24_35.rep



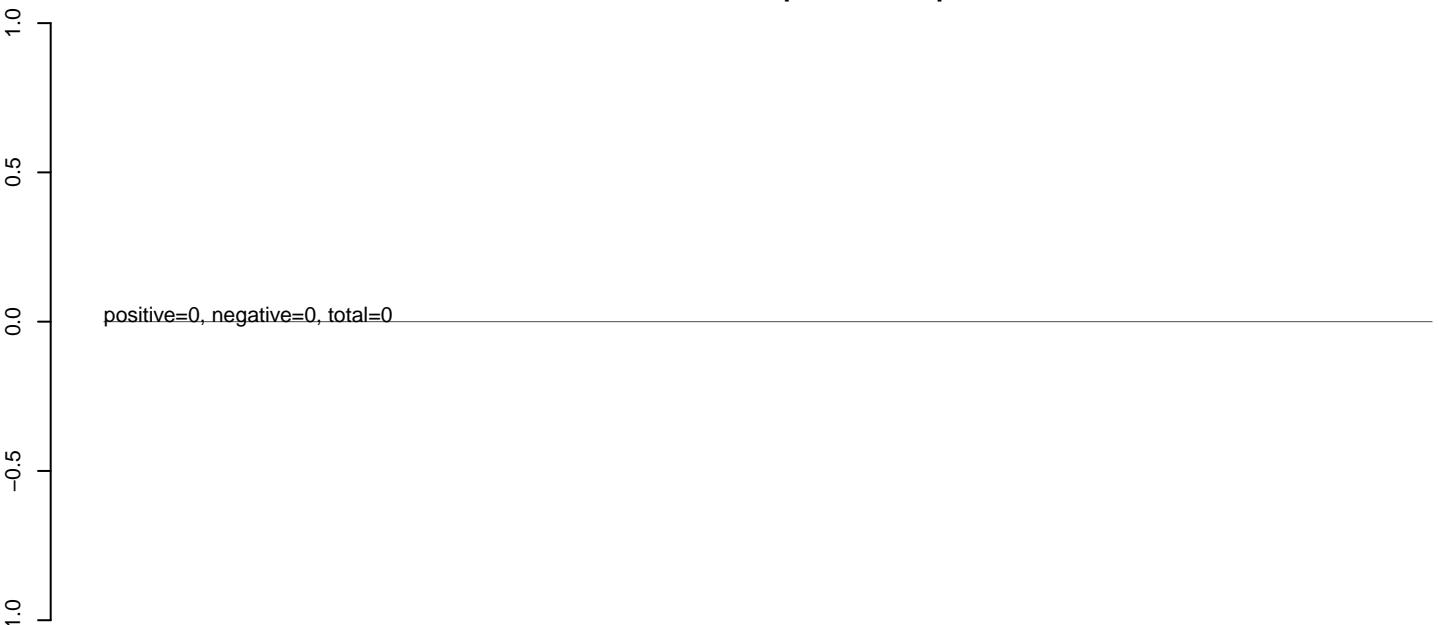
CuQuin_MG_7dpi3.rep



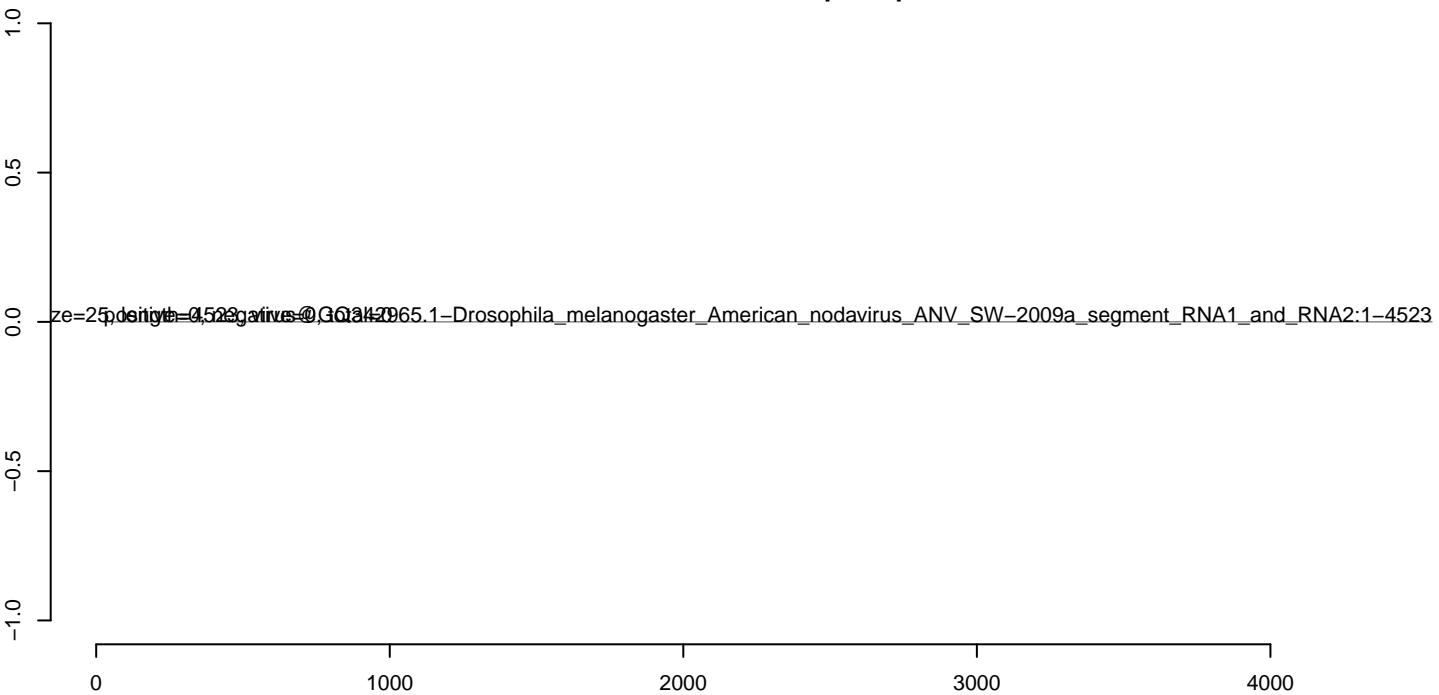
CuQuin_MG_7dpi3.18_23.rep



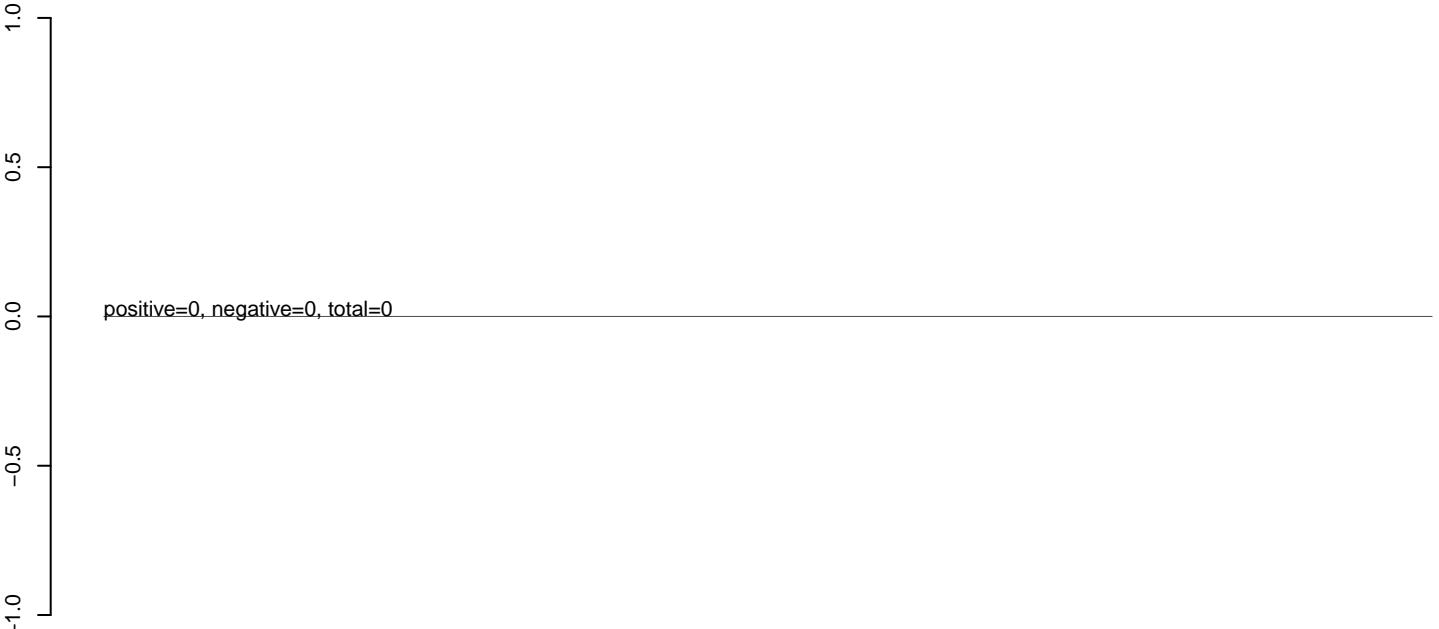
CuQuin_MG_7dpi3.24_35.rep



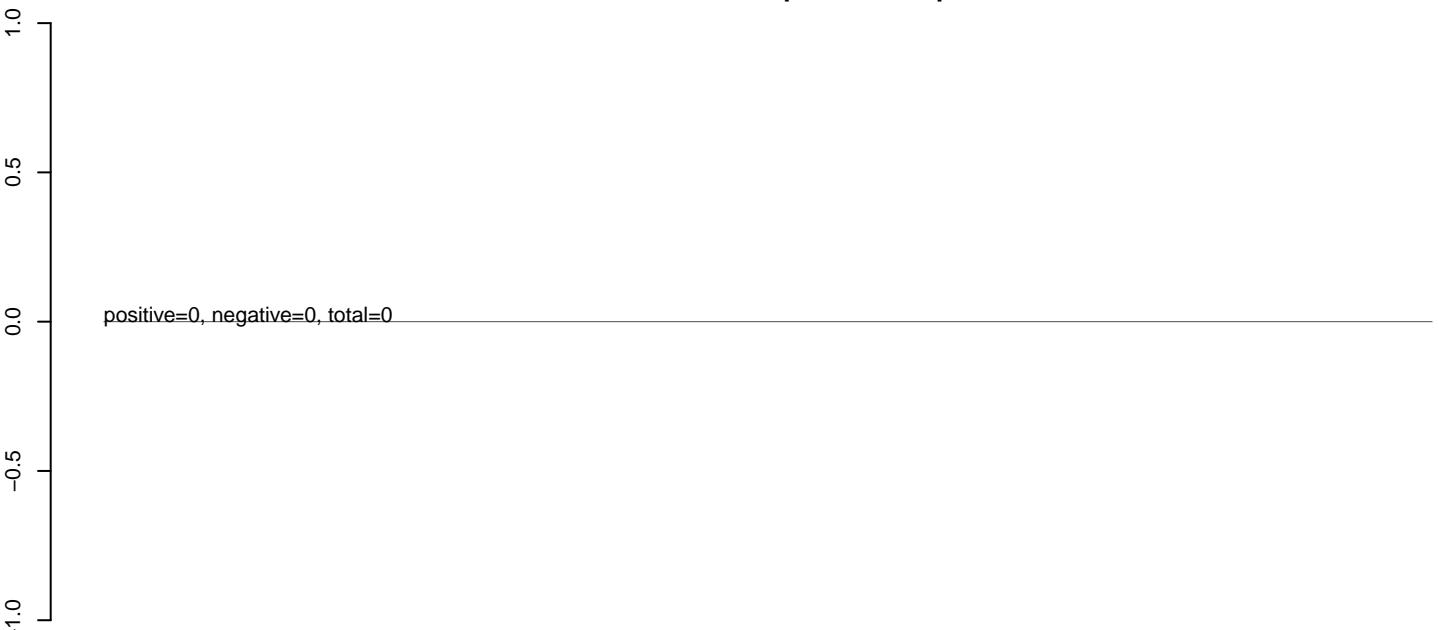
CuQuin_MG_7dpi3.rep



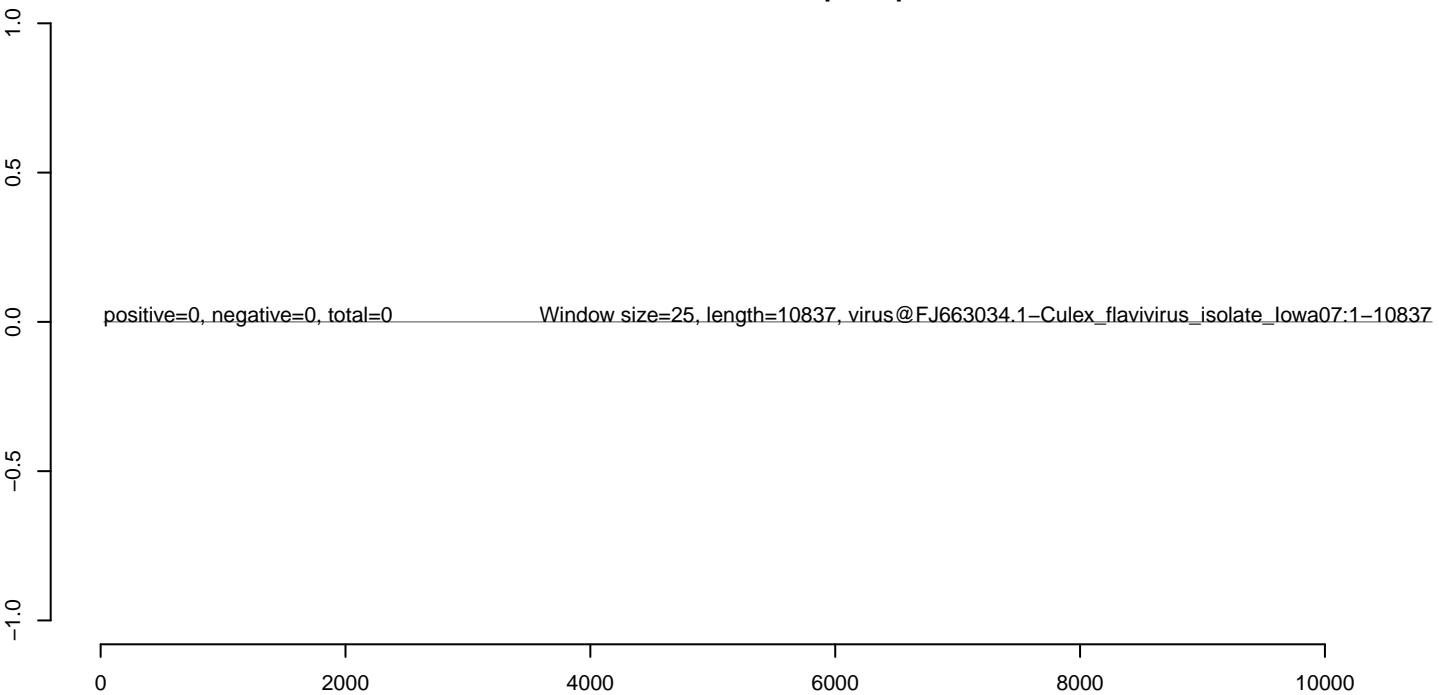
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=10837, virus@FJ502995.1-Culex_flavivirus_strain_HOU24518:1-10837

0 2000 4000 6000 8000 10000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

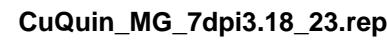
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=10837, virus@EU879060.1-Culex_flavivirus_strain_CxFV-Mex07:1-10837

0 2000 4000 6000 8000 10000



positive=0, negative=0, total=0



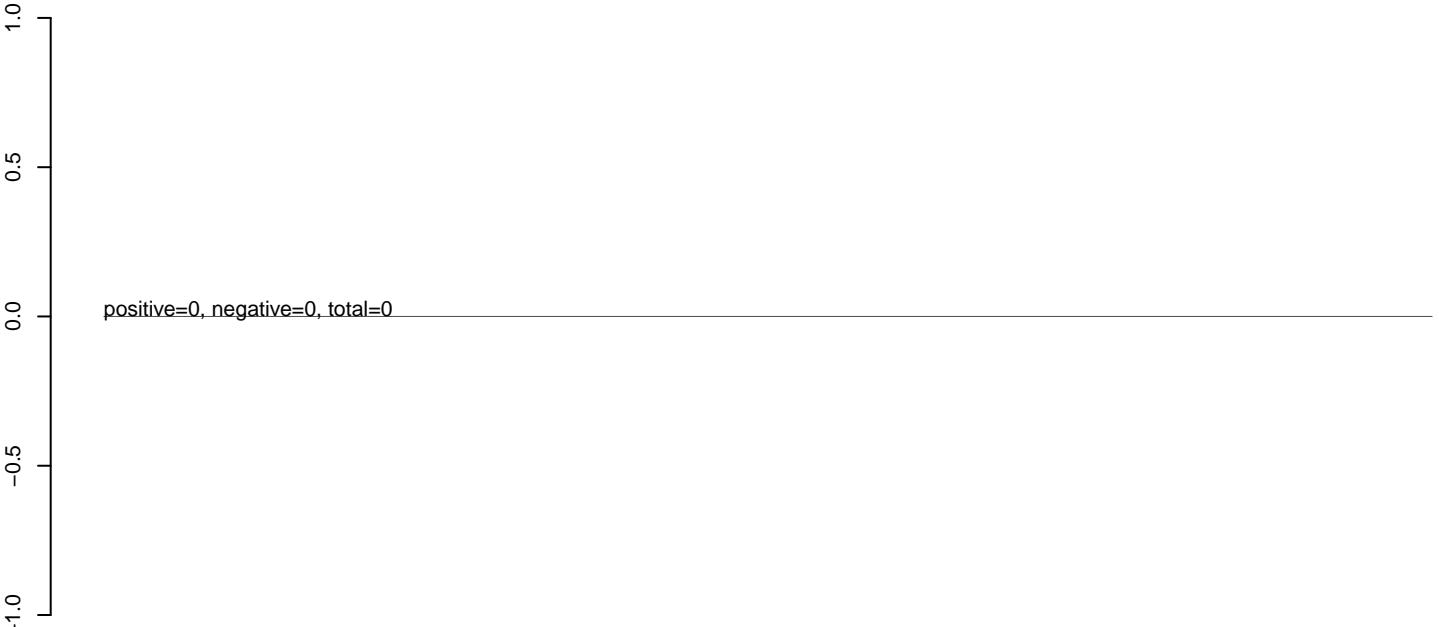
positive=0, negative=0, total=0



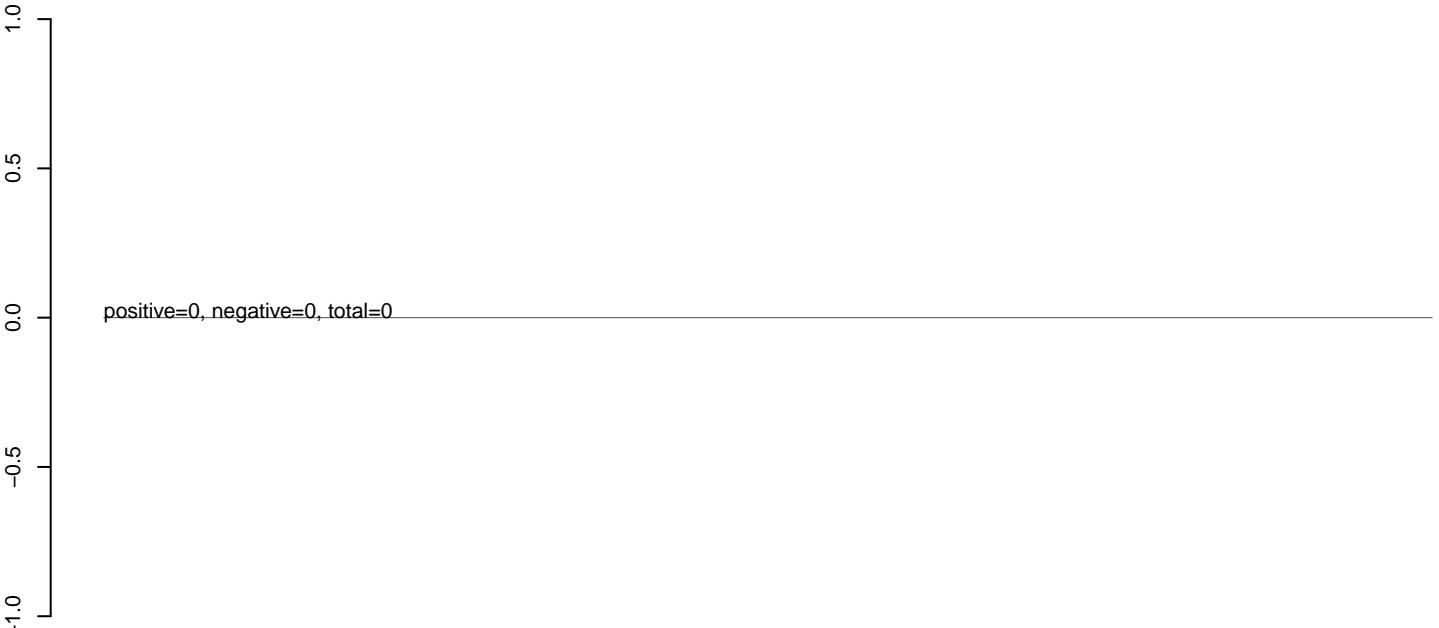
Cx_ow_cxpxz3f5ydo66tho26109-0-0-0@E5E5372756.1_Culex_pipiens-pallens_denguevirus-strain_IZ_16_pro-structural-protein-1_NS1_pro-ctrxi1_3498

0 500 1000 1500 2000 2500 3000 3500

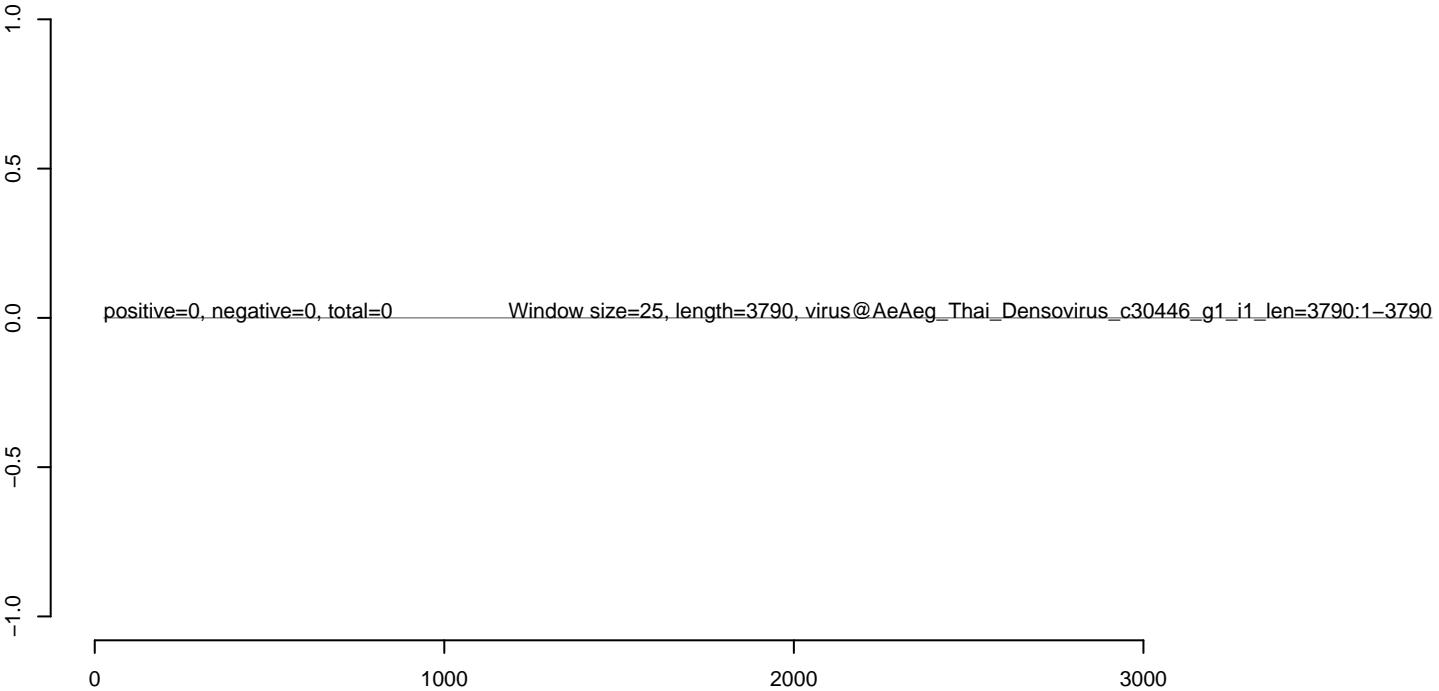
CuQuin_MG_7dpi3.18_23.rep



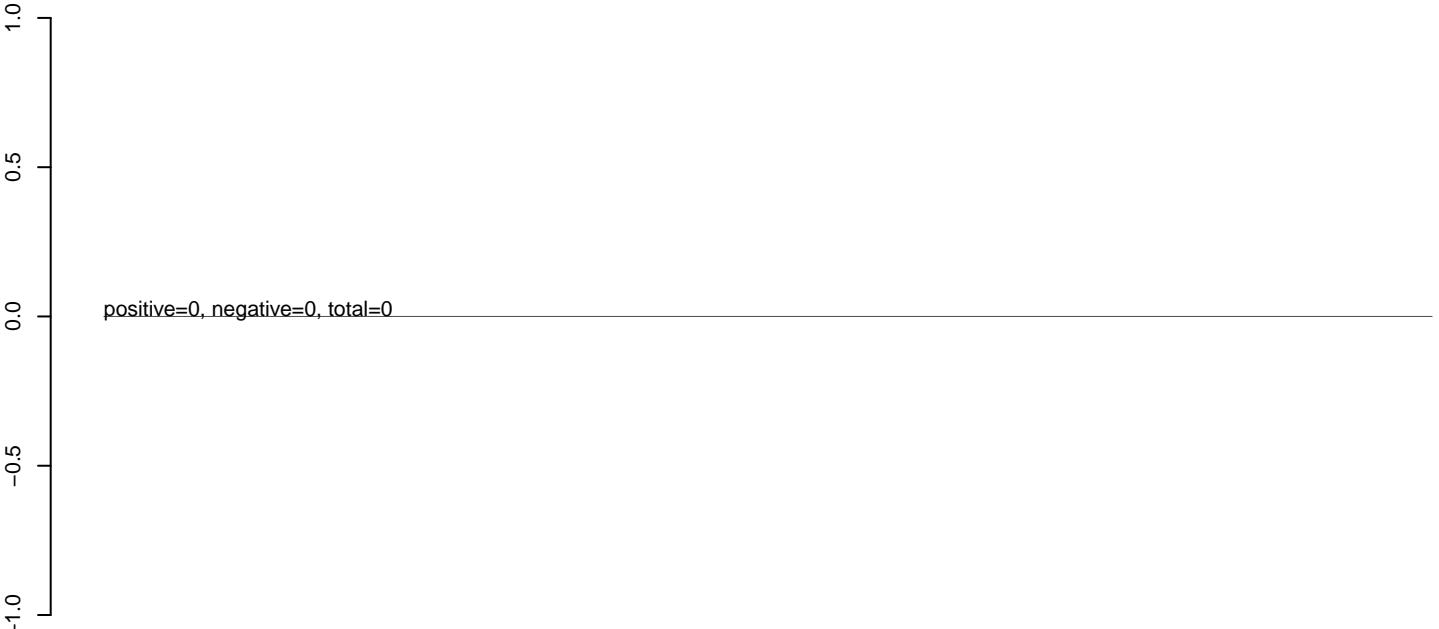
CuQuin_MG_7dpi3.24_35.rep



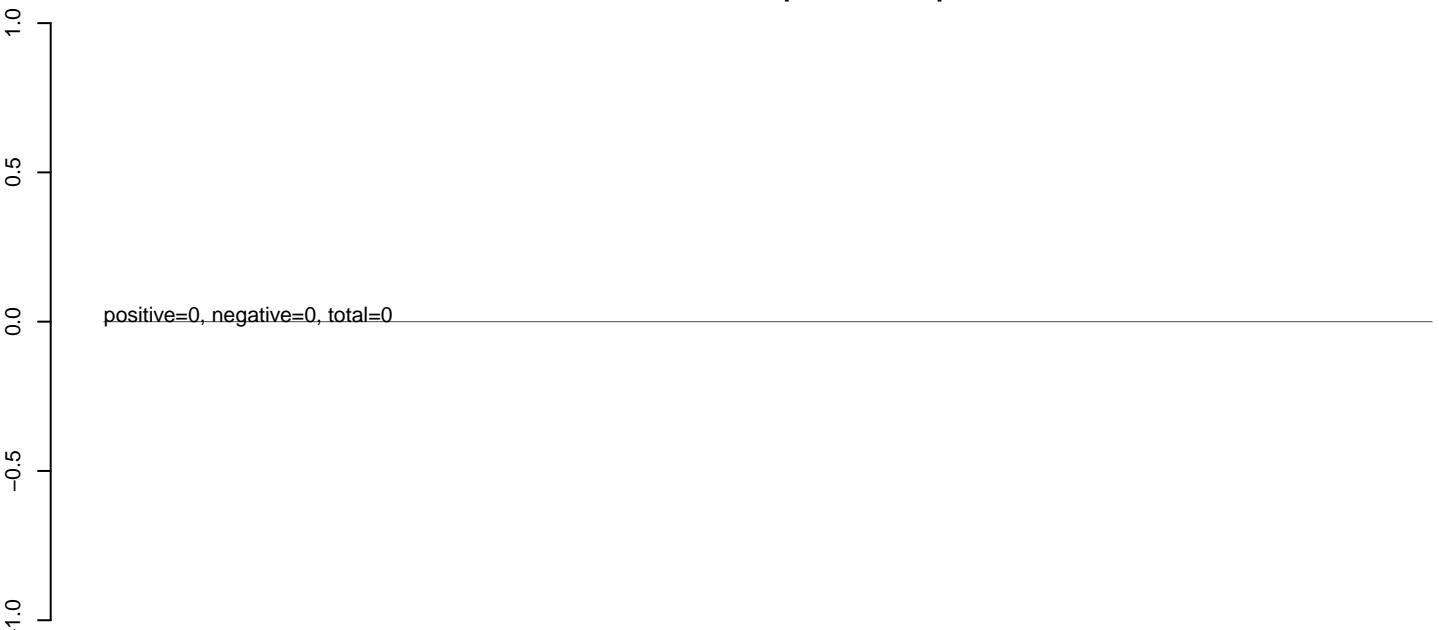
CuQuin_MG_7dpi3.rep



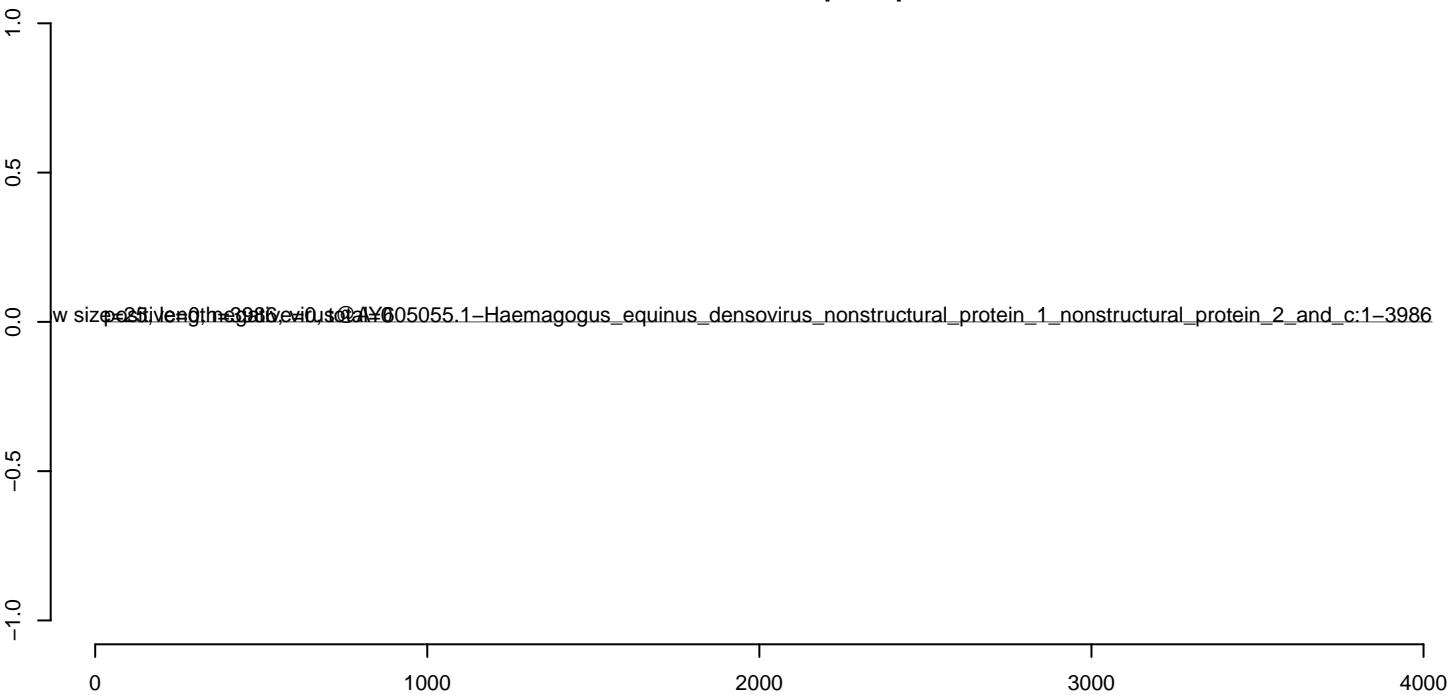
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

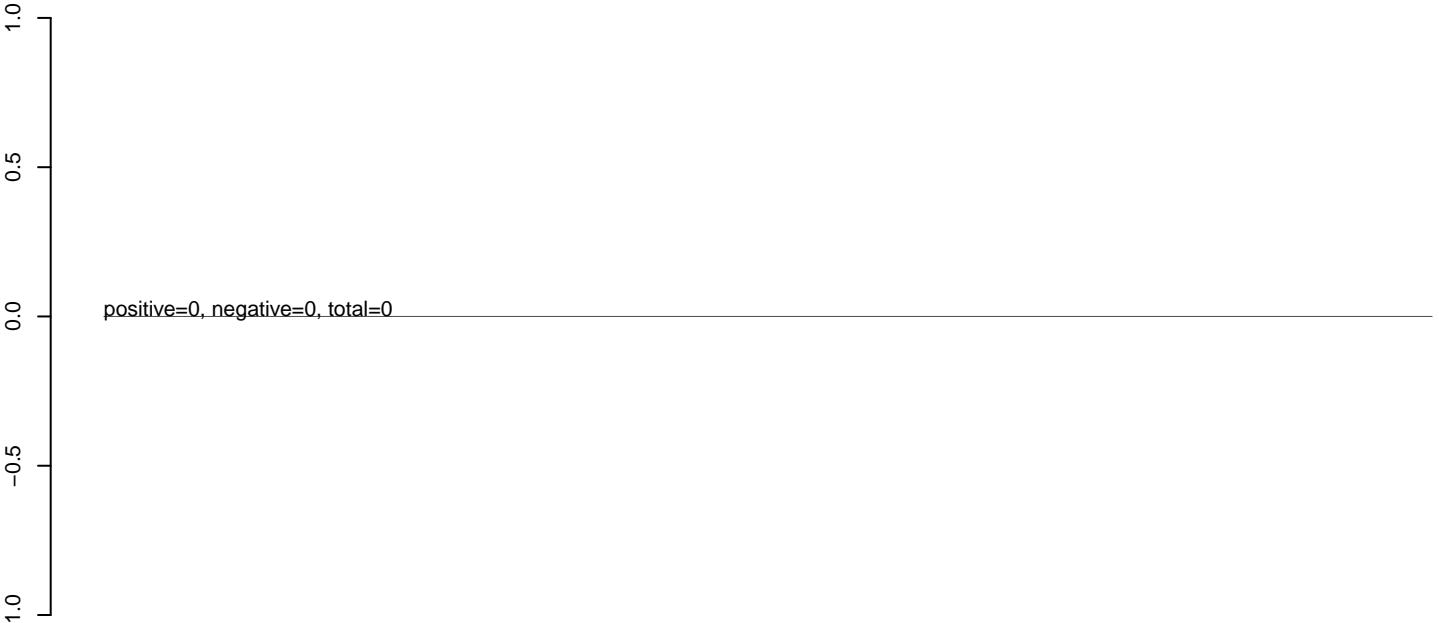
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

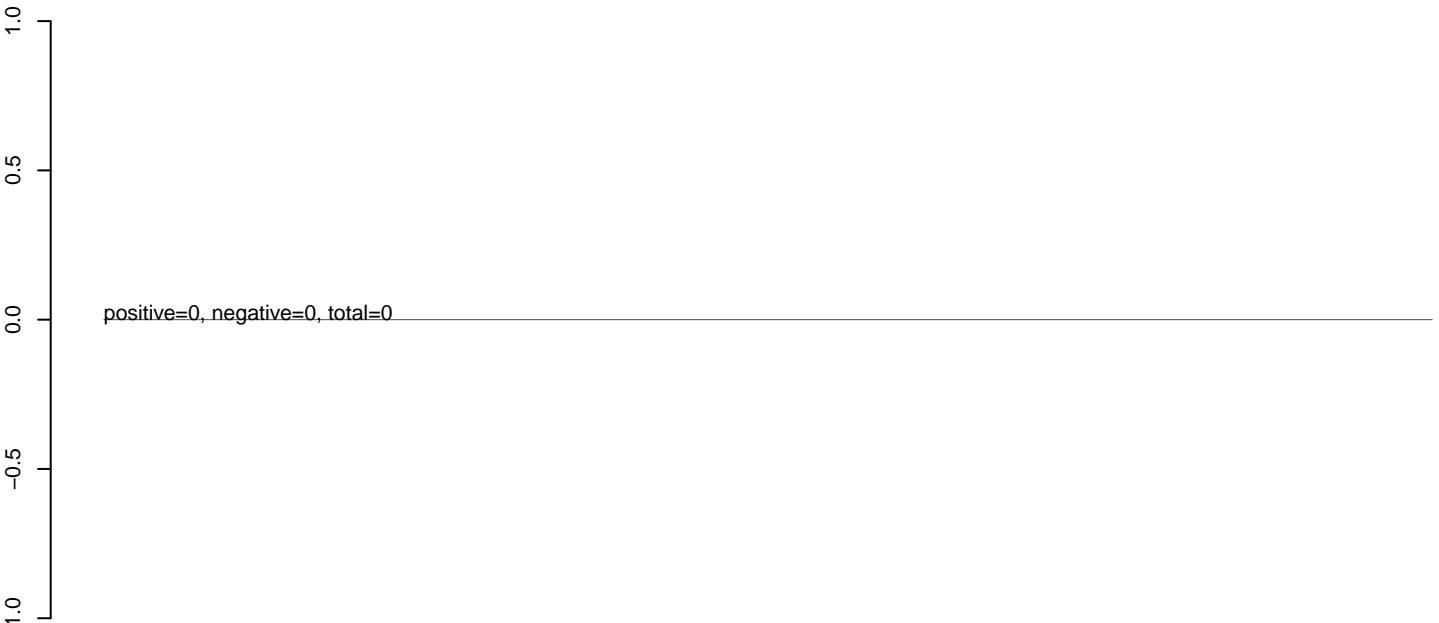
Window size=25, length=2965, virus@AY347953.2-Aedes_aegypti_genomic_sequence:1-2965

0 500 1000 1500 2000 2500 3000

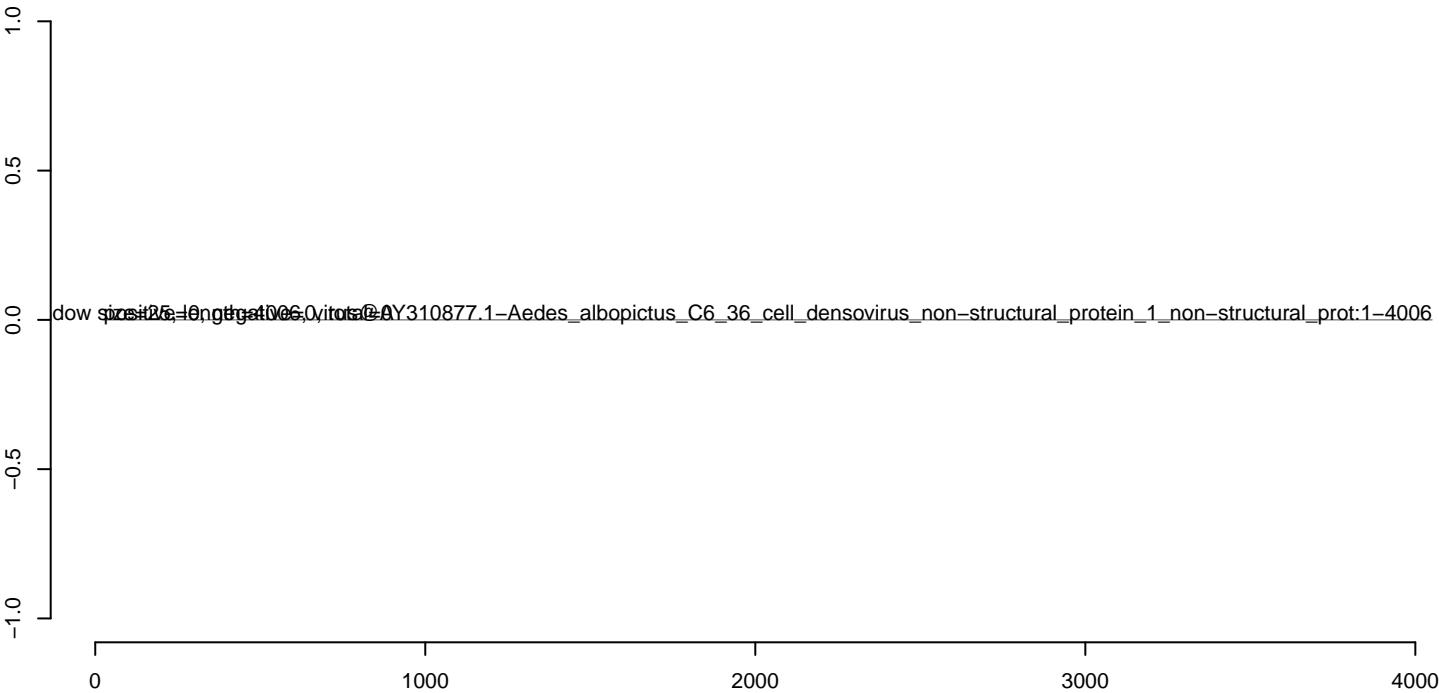
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep

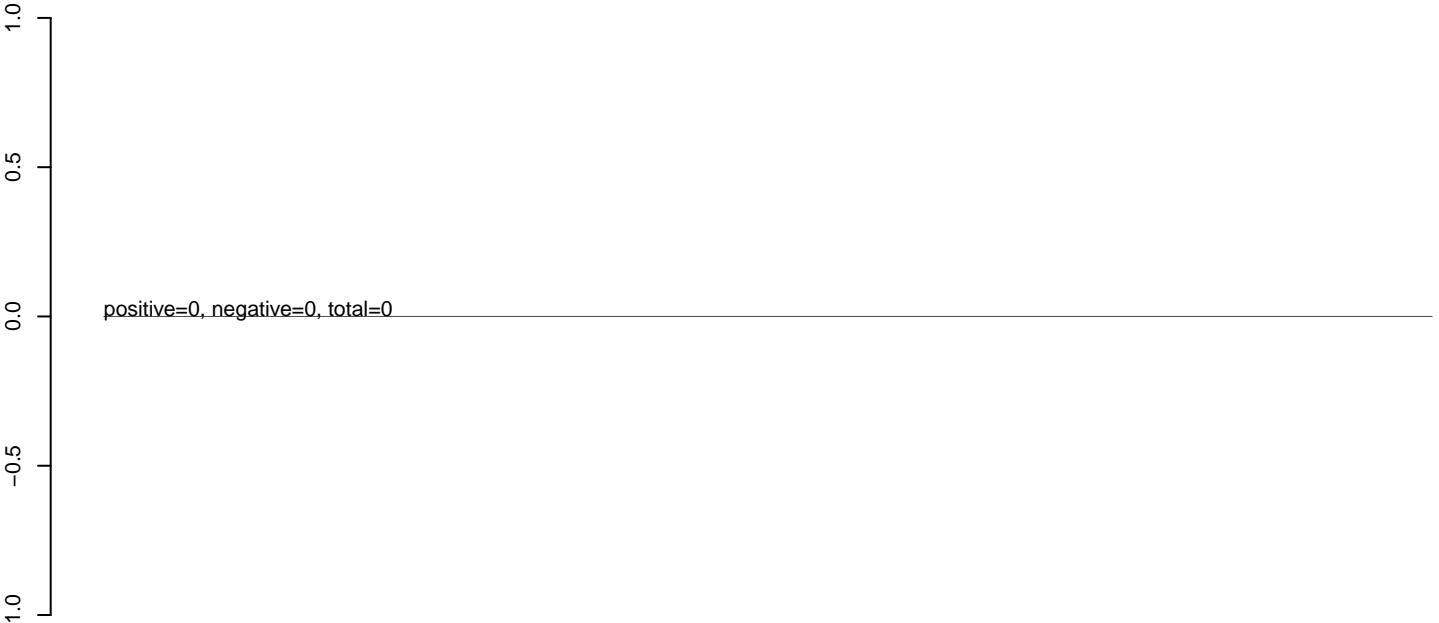


CuQuin_MG_7dpi3.rep

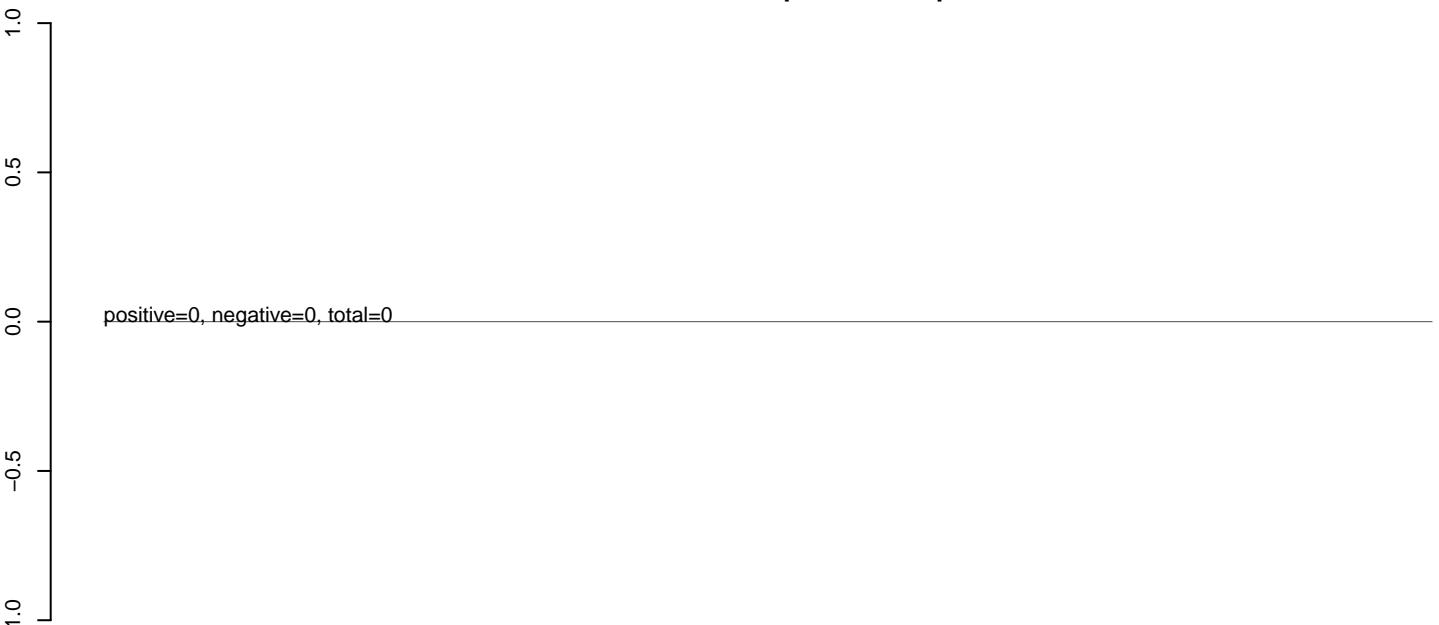


dow size#25, length#1006, 0, info@AY310877.1-Aedes_albopictus_C6_36_cell_densovirus_non-structural_protein_1_non-structural_prot:1-4006

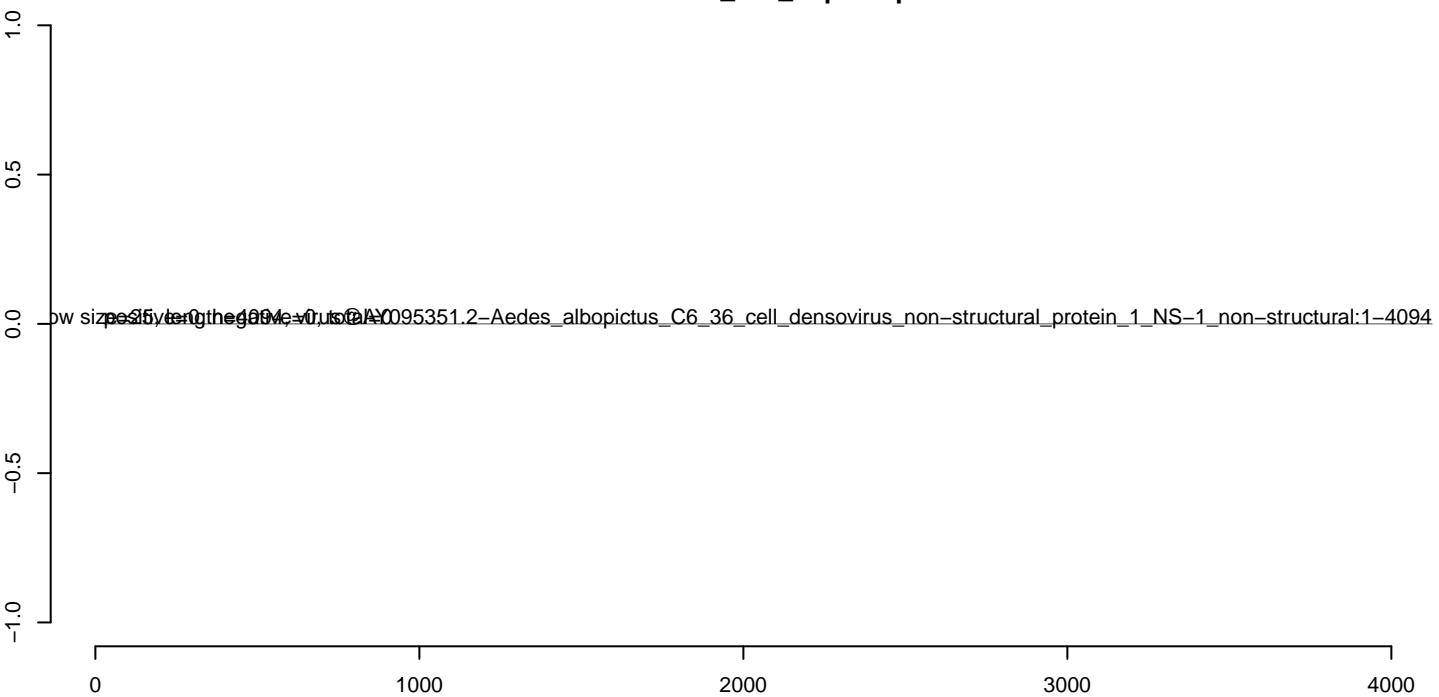
CuQuin_MG_7dpi3.18_23.rep



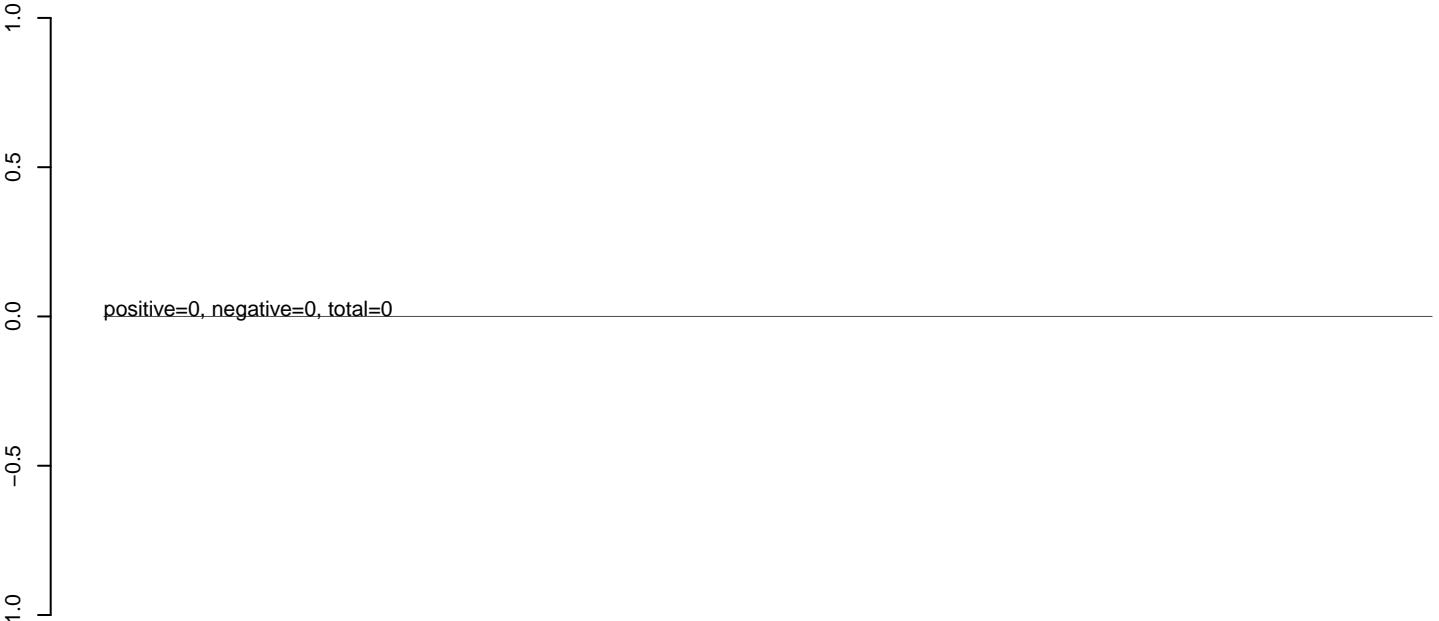
CuQuin_MG_7dpi3.24_35.rep



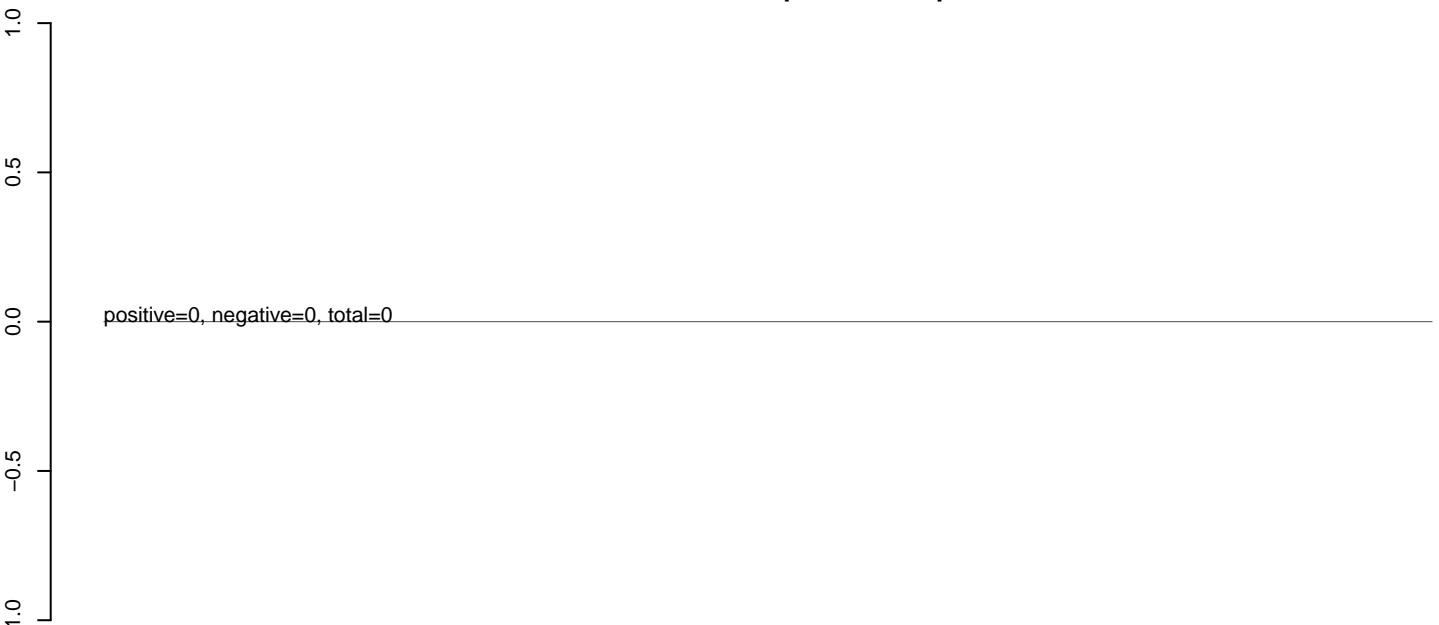
CuQuin_MG_7dpi3.rep



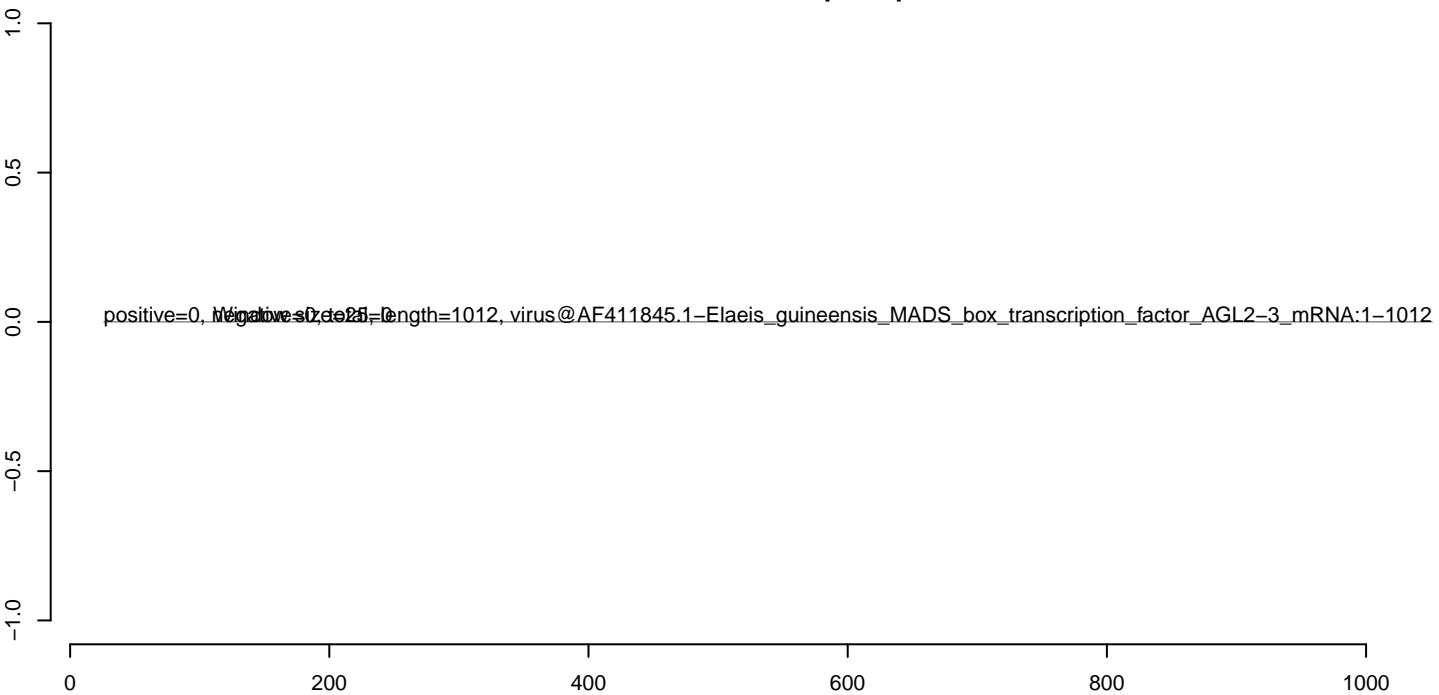
CuQuin_MG_7dpi3.18_23.rep



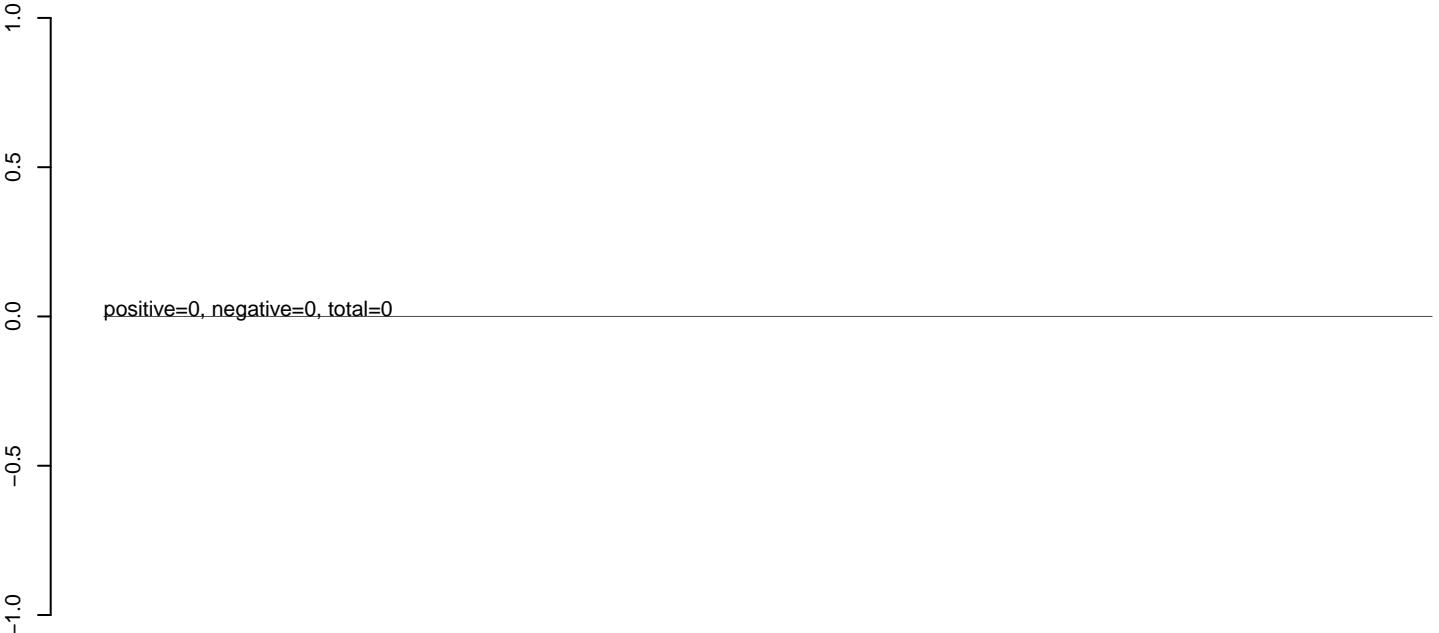
CuQuin_MG_7dpi3.24_35.rep



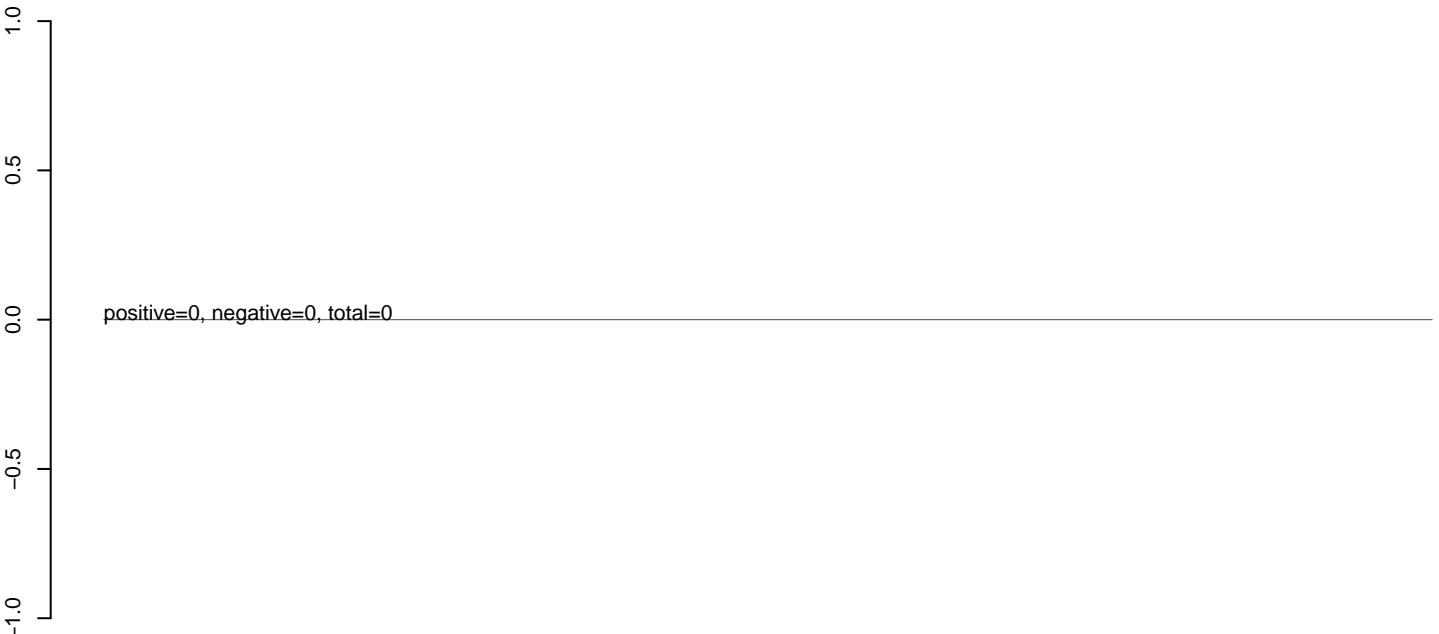
CuQuin_MG_7dpi3.rep



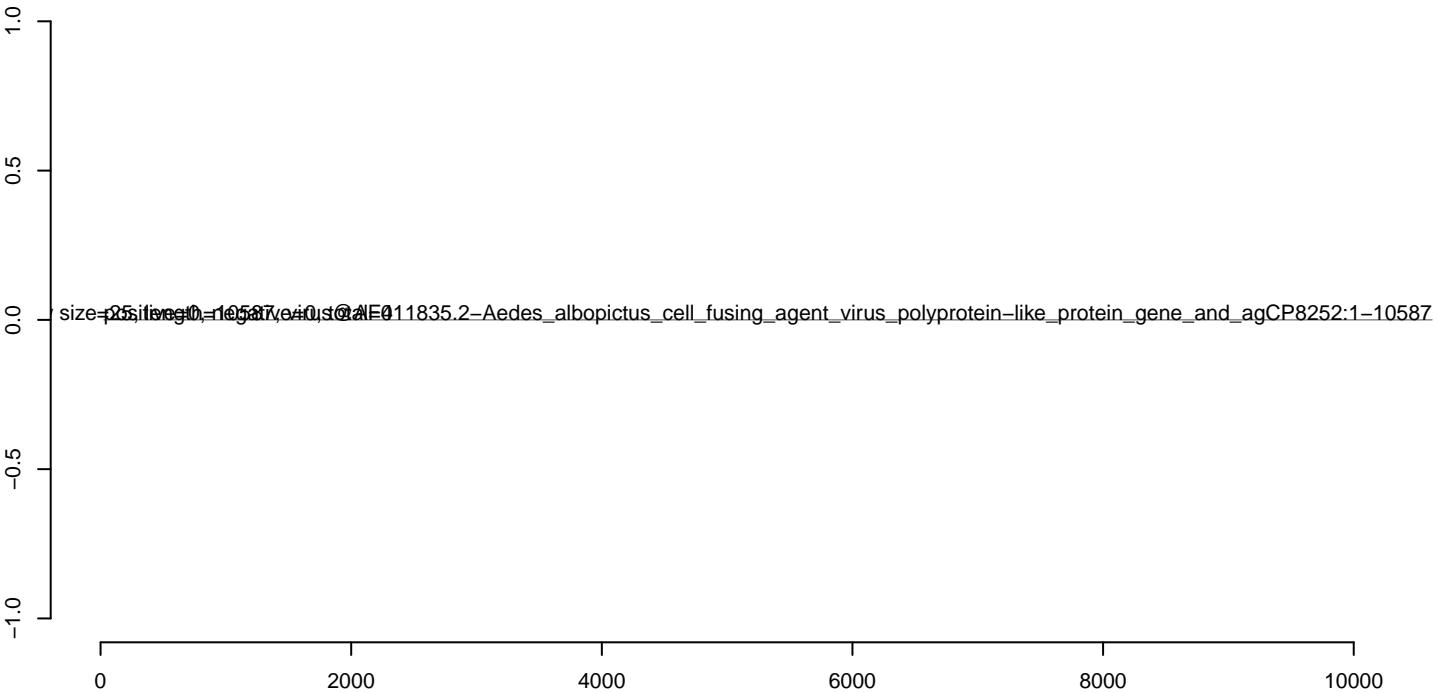
CuQuin_MG_7dpi3.18_23.rep



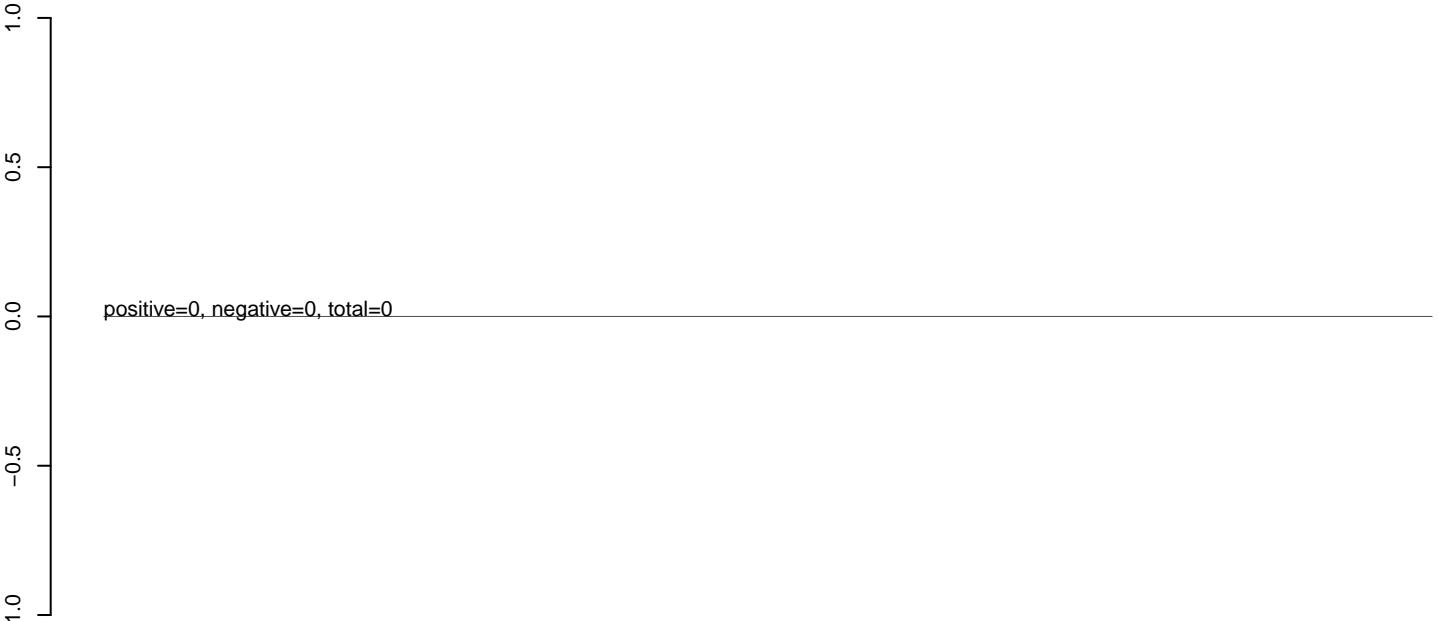
CuQuin_MG_7dpi3.24_35.rep



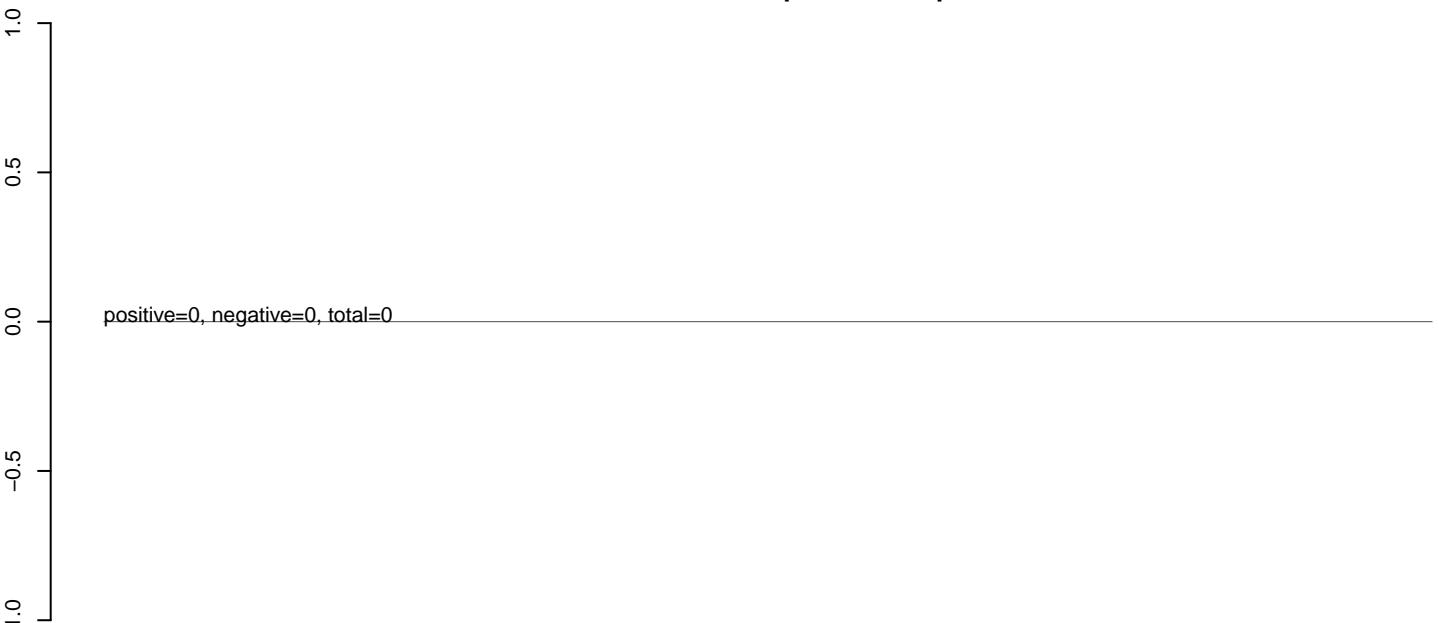
CuQuin_MG_7dpi3.rep



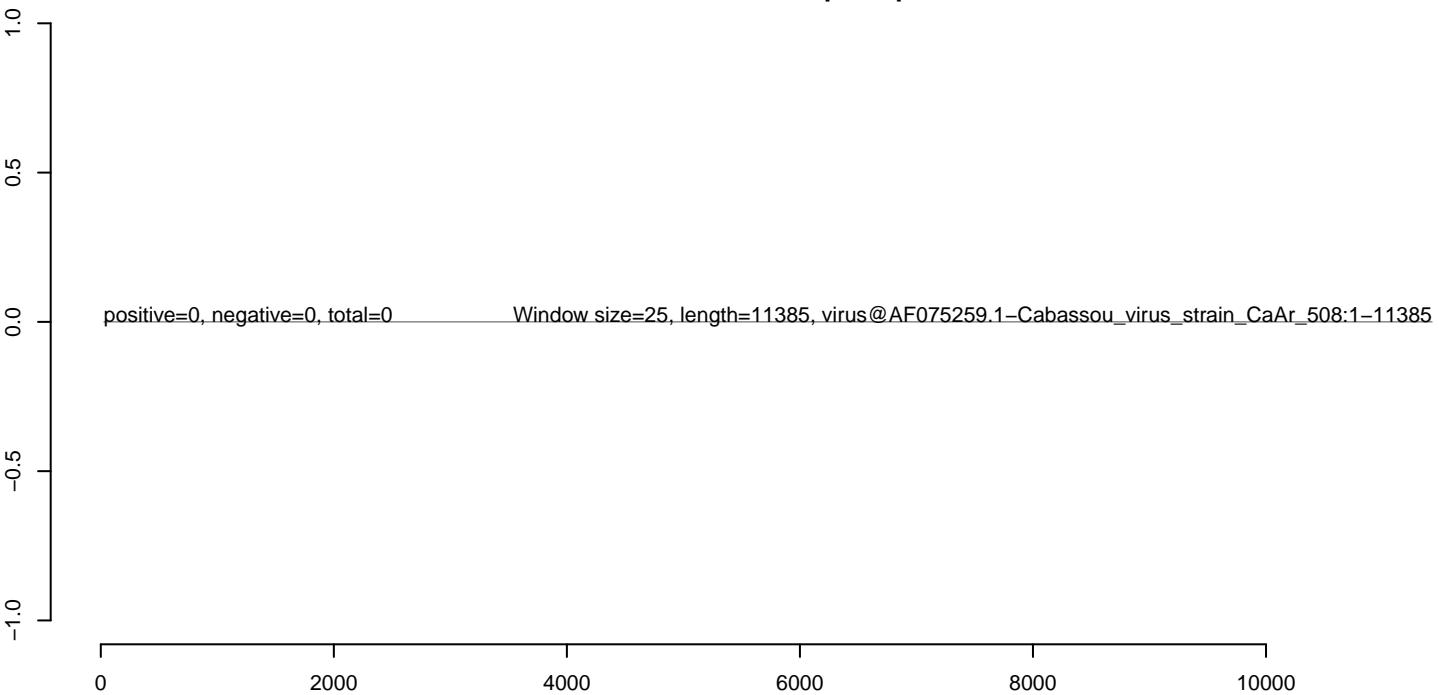
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

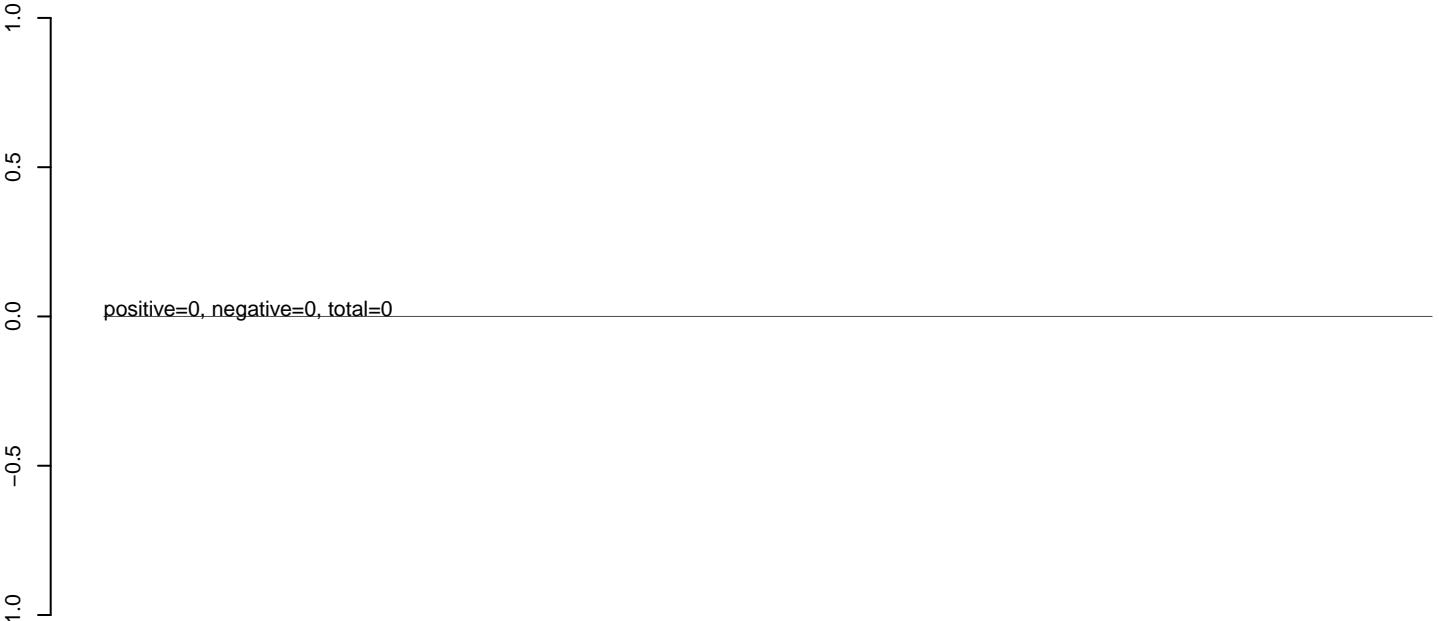
CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

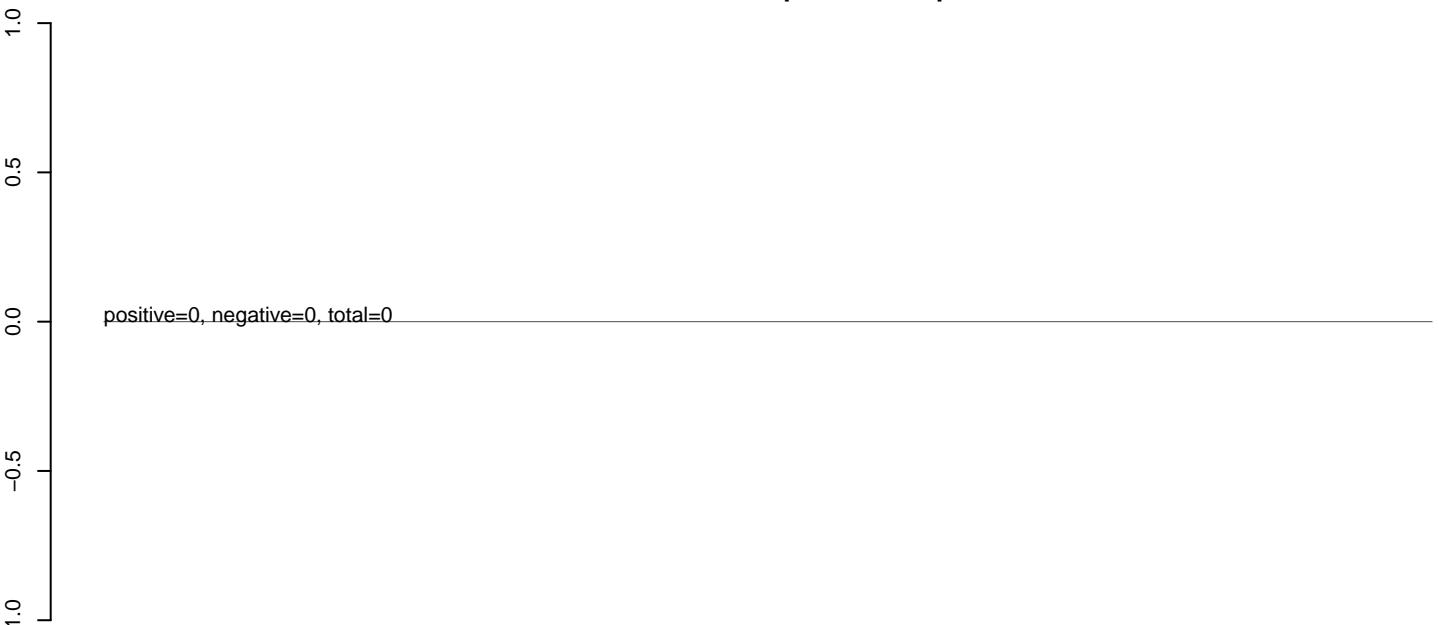
Window size=25, length=11465, virus@AF075257.1-Mosso_das_Pedras_virus:1-11465

0 2000 4000 6000 8000 10000

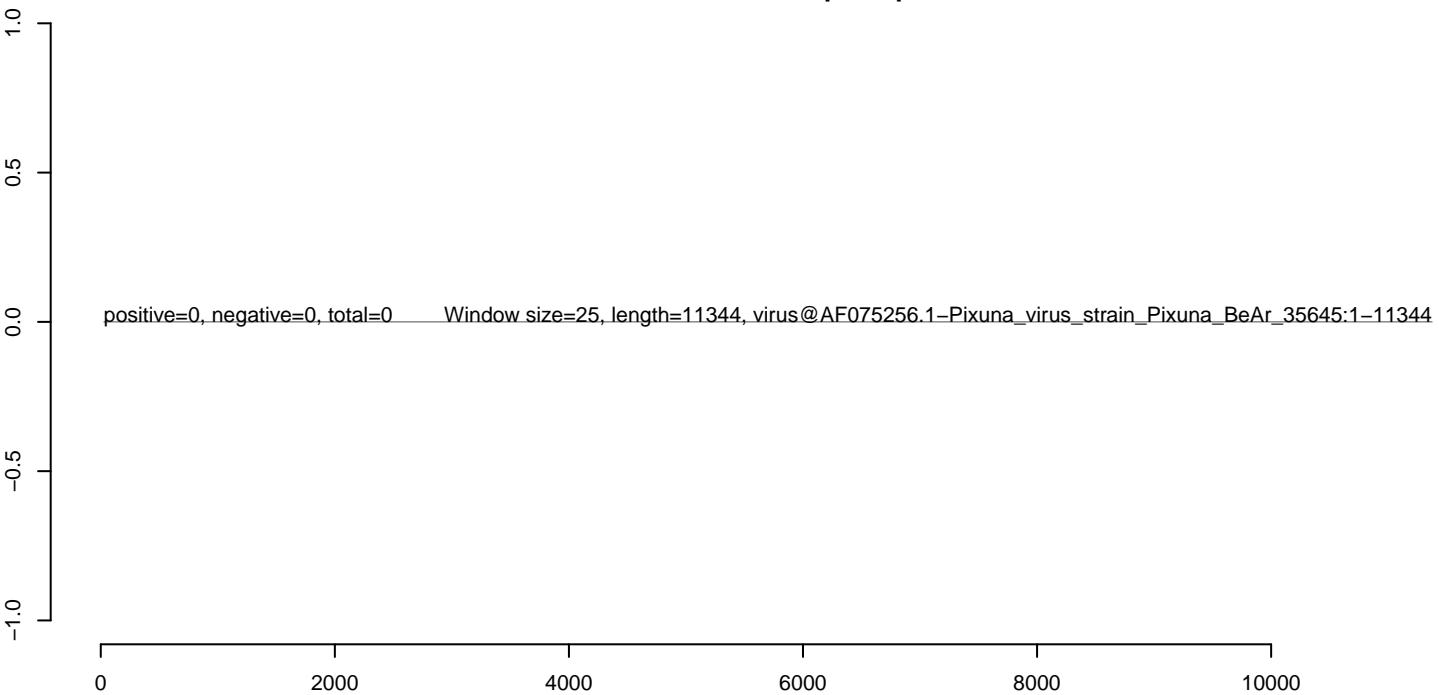
CuQuin_MG_7dpi3.18.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0

Window size=25, length=11530, virus@AF075254.1-Tonate_virus_strain_Tonate_CaAn_410d:1-11530

0 2000 4000 6000 8000 10000 12000

CuQuin_MG_7dpi3.18.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

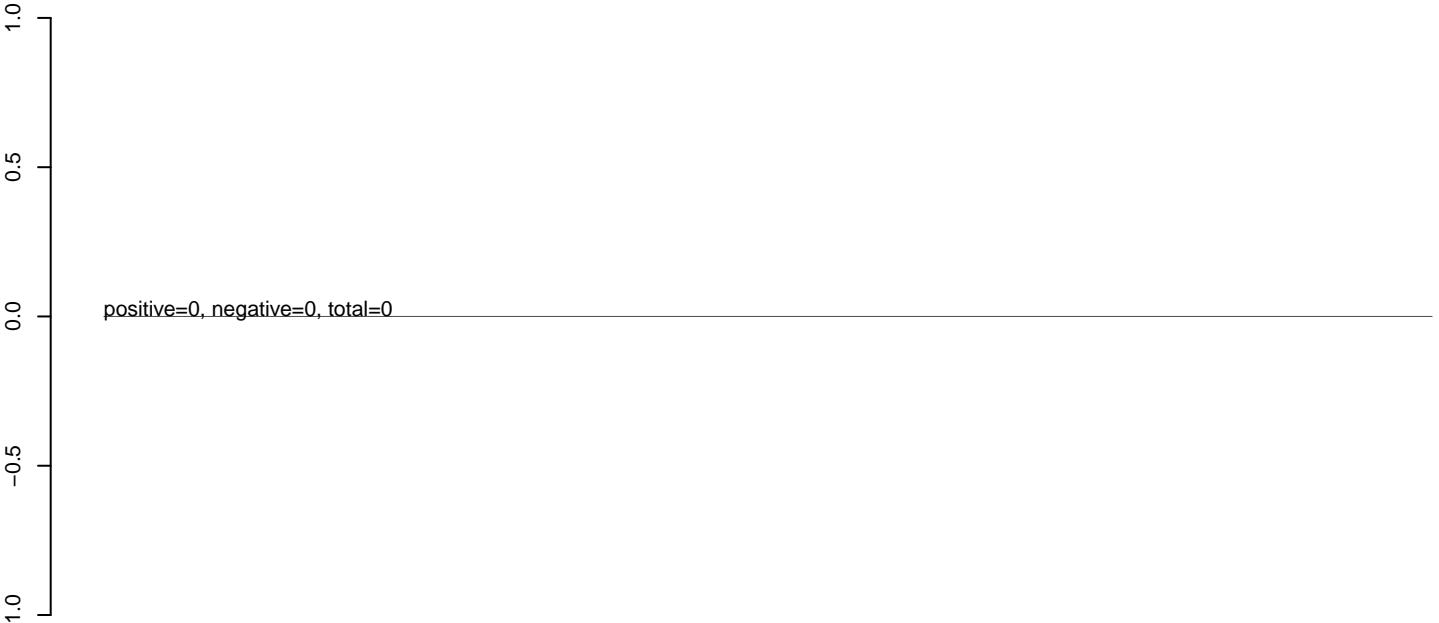
positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

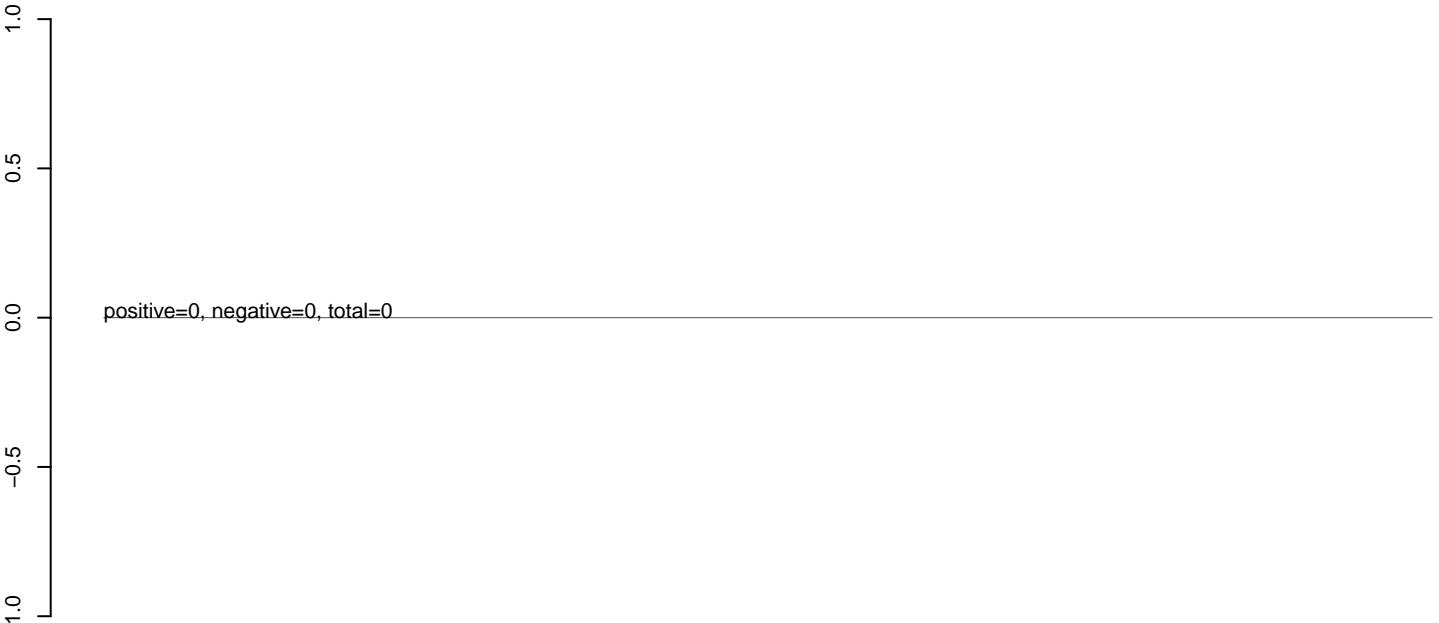
positive=0, negative=0, total=0 Window size=25, length=11391, virus@AF075253.1-Mucambo_virus_strain_Mucambo_BeAn_8:1-11391

0 2000 4000 6000 8000 10000

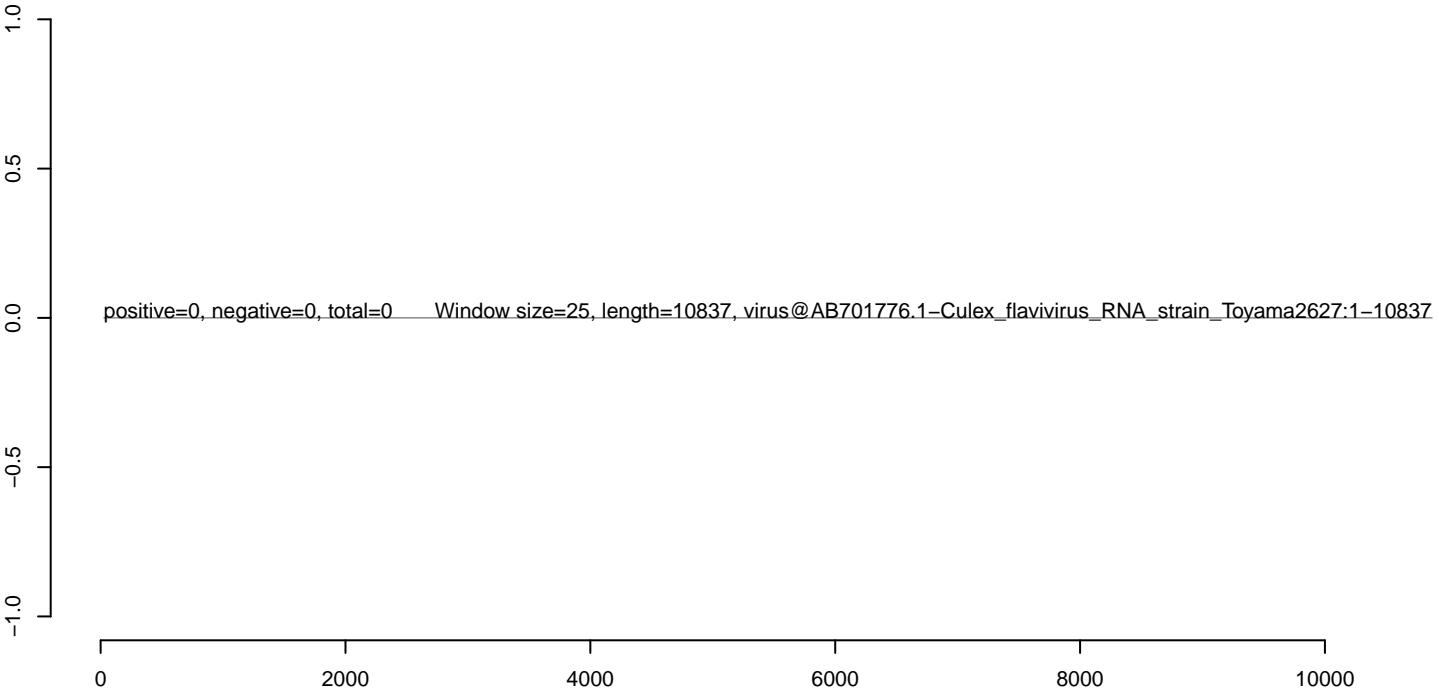
CuQuin_MG_7dpi3.18_23.rep



CuQuin_MG_7dpi3.24_35.rep



CuQuin_MG_7dpi3.rep



CuQuin_MG_7dpi3.18_23.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.24_35.rep

positive=0, negative=0, total=0

CuQuin_MG_7dpi3.rep

positive=0, negative=0, total=0 Window size=25, length=10837, virus@AB262759.2-Culex_flavivirus_genomic_RNA_strain_Tokyo:1-10837