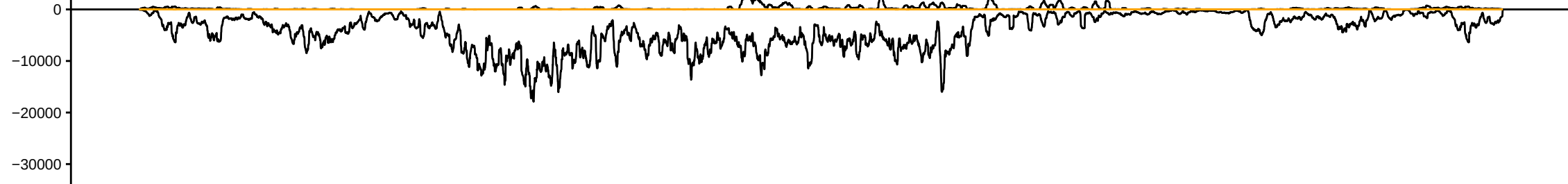
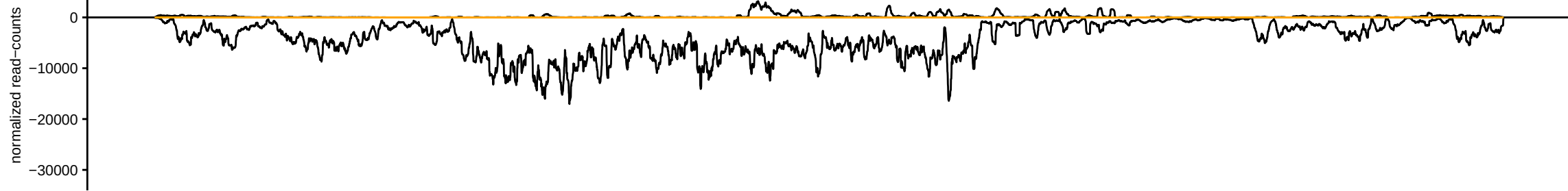


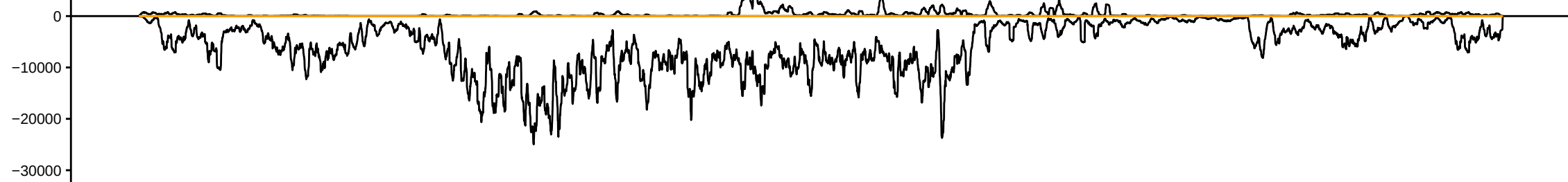
sRNAseq_HsBam_nos_ControlKD_00hAHS_repB_283080_sRBC_GTCAG



sRNAseq_HsBam_nos_ControlKD_00hAHS_repC_283081_sRBC_CAGTG



sRNAseq_HsBam_nos_ControlKD_00hAHS_repD_283082_sRBC_GTCAG



0

2000

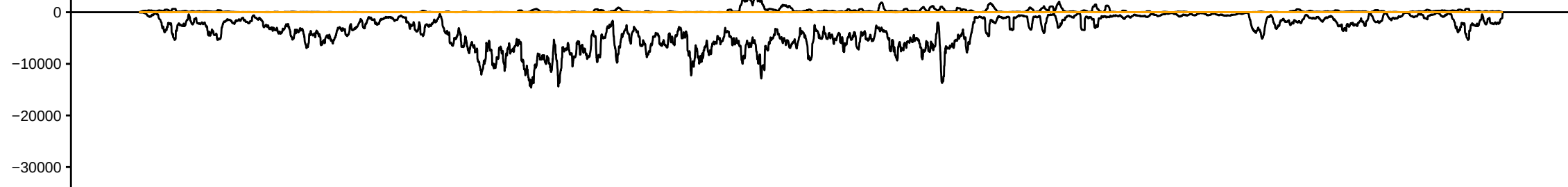
4000

6000

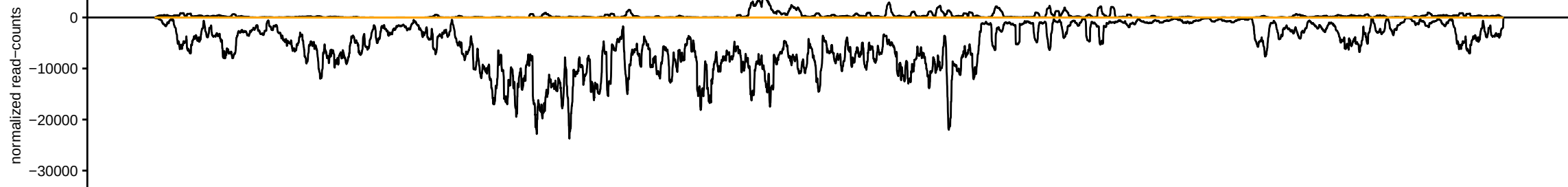
8000

position on gypsy5

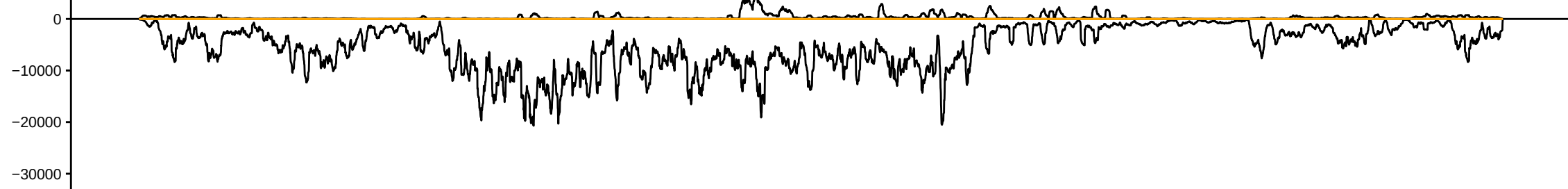
sRNAseq_HsBam_nos_ControlKD_68hAHS_repB_283080_sRBC_TGACT



sRNAseq_HsBam_nos_ControlKD_68hAHS_repC_283081_sRBC_AGCAA



sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT



0

2000

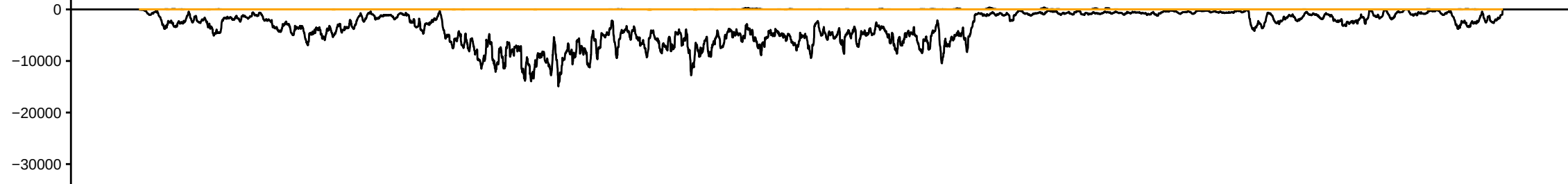
4000

6000

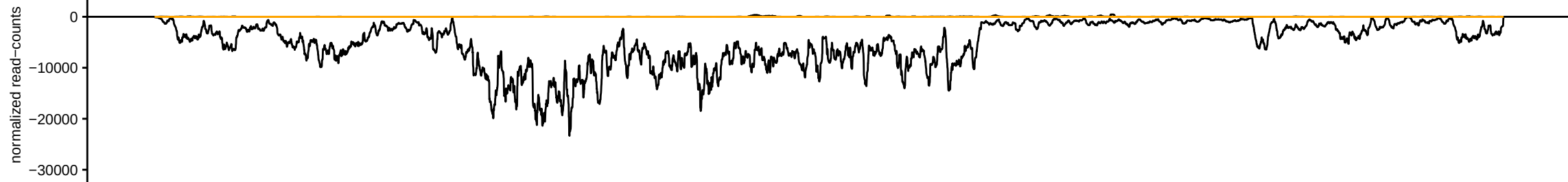
8000

position on gypsy5

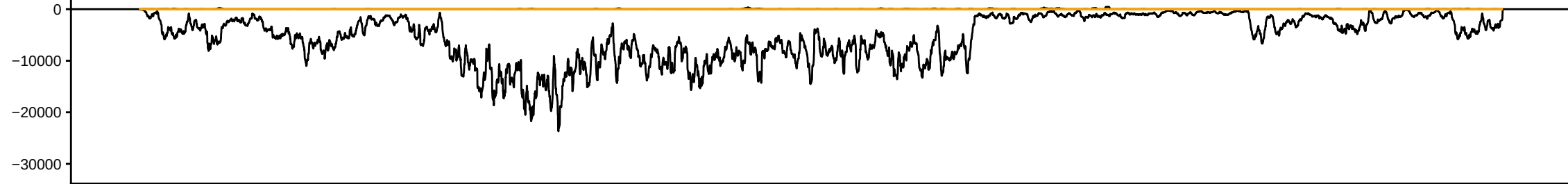
sRNAseq_HsBam_nos_PiwiKD_00hAHS_repB_283080_sRBC_CGTTC



sRNAseq_HsBam_nos_PiwiKD_00hAHS_repC_283081_sRBC_GGTAT



sRNAseq_HsBam_nos_PiwiKD_00hAHS_repD_283082_sRBC_CGTTC



0

2000

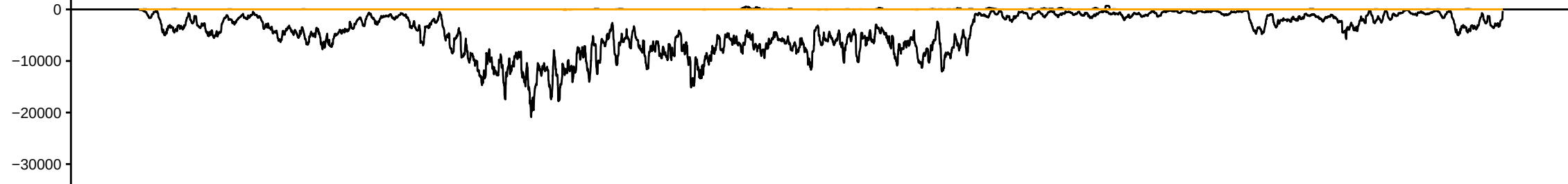
4000

6000

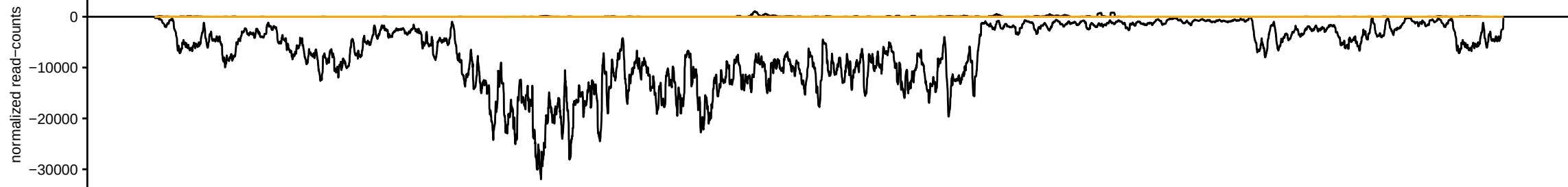
8000

position on gypsy5

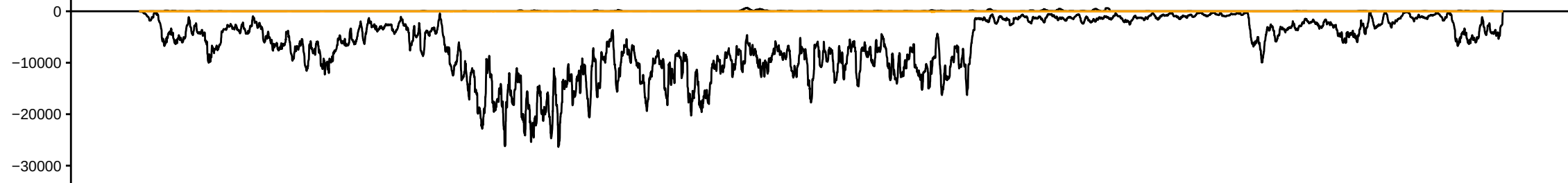
sRNAseq_HsBam_nos_PiwiKD_68hAHS_repB_283080_sRBC_ATGGA



sRNAseq_HsBam_nos_PiwiKD_68hAHS_repC_283081_sRBC_TACCA



sRNAseq_HsBam_nos_PiwiKD_68hAHS_repD_283082_sRBC_ATGGA



0

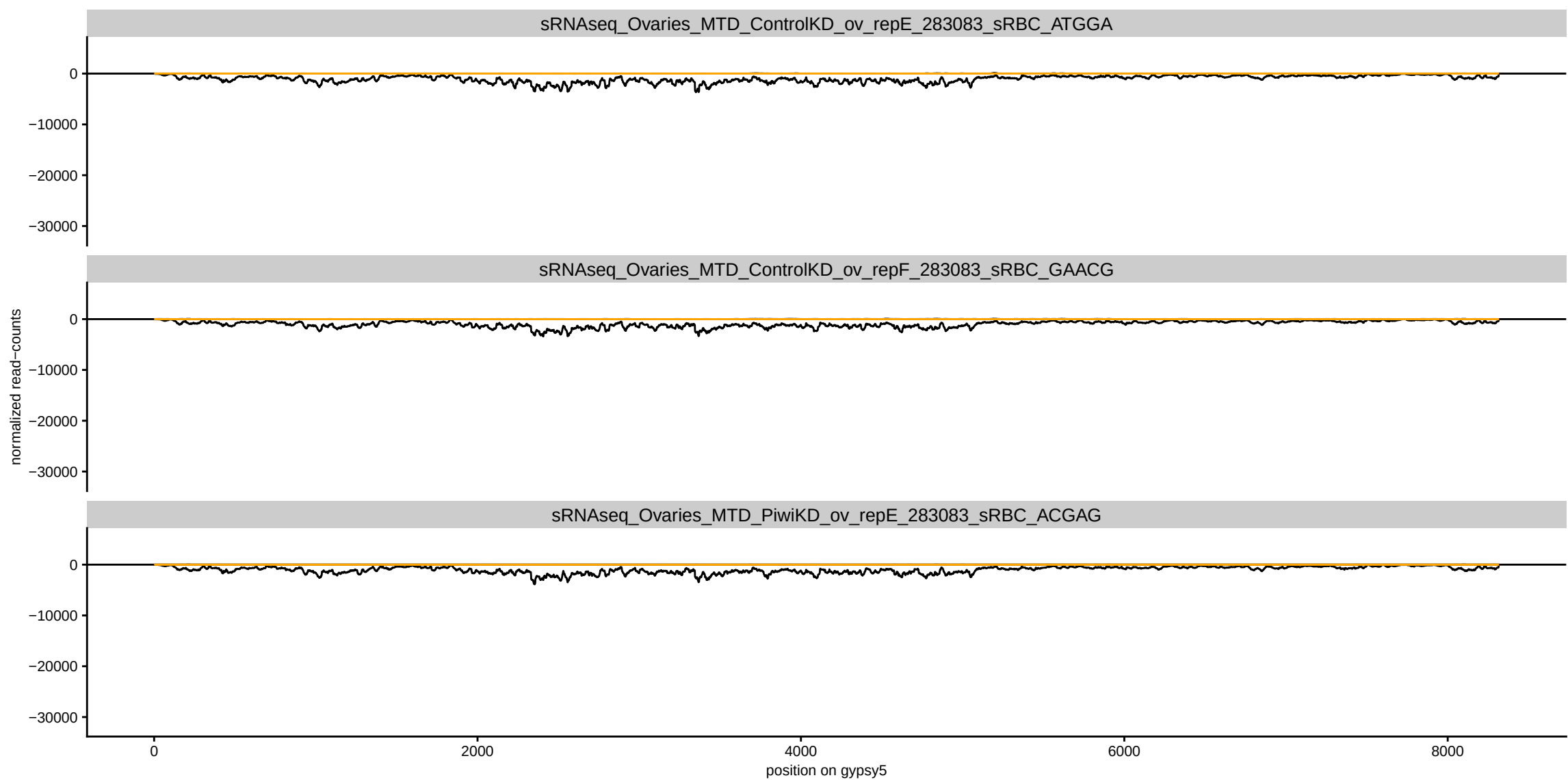
2000

4000

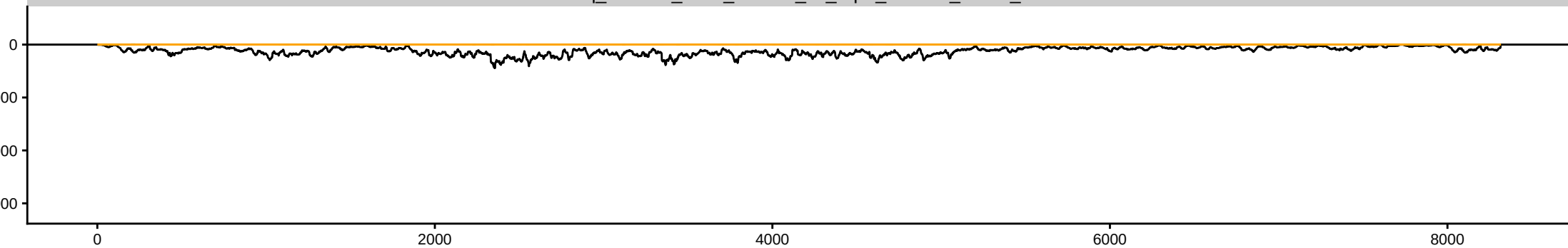
6000

8000

position on gypsy5



normalized read-counts



position on gypsy5