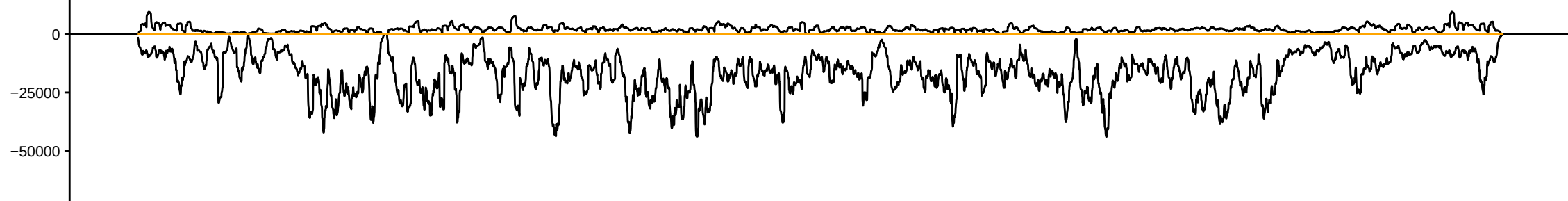
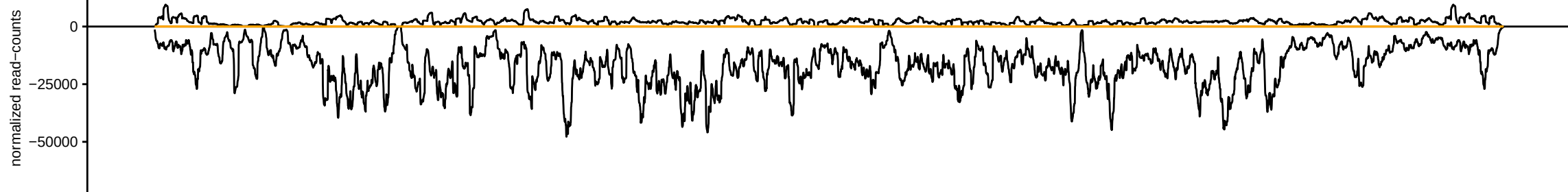


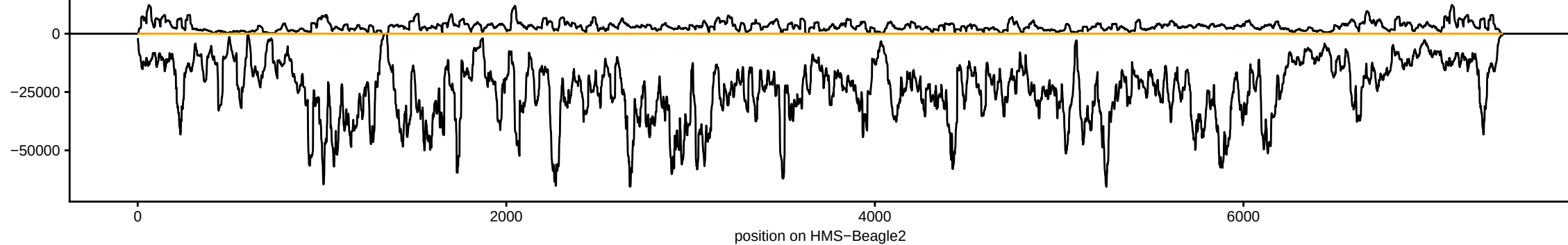
sRNAseq_HsBam_nos_ControlKD_00hAHS_repB_283080_sRBC_GTCAG



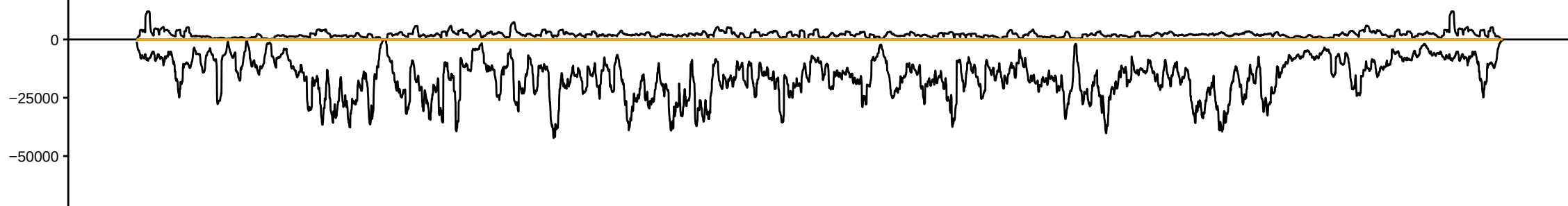
sRNAseq_HsBam_nos_ControlKD_00hAHS_repC_283081_sRBC_CAGTG



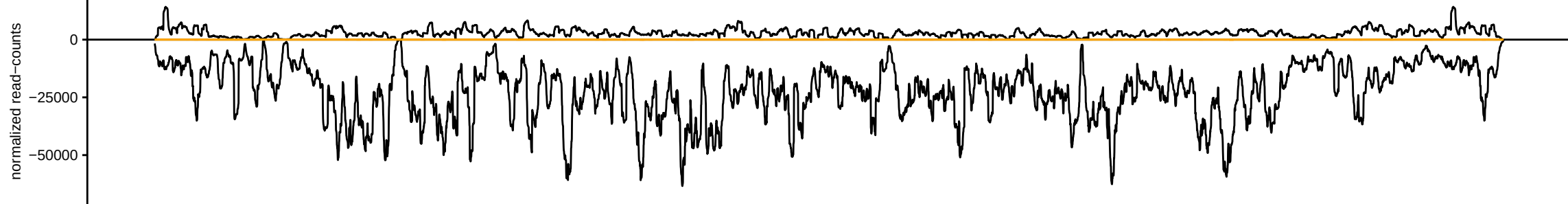
sRNAseq_HsBam_nos_ControlKD_00hAHS_repD_283082_sRBC_GTCAG



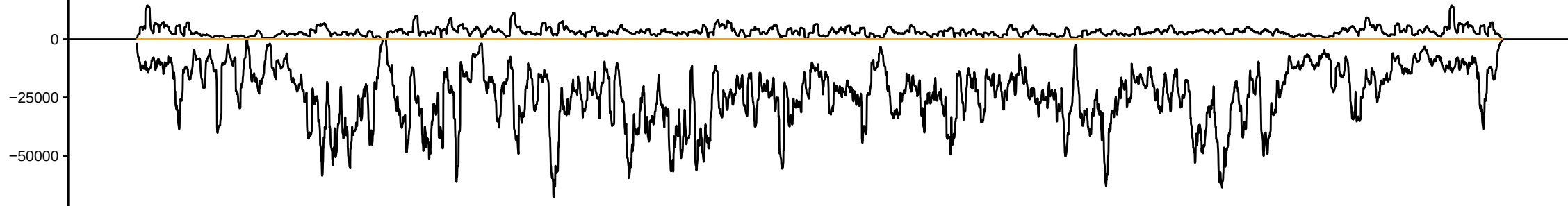
sRNAseq_HsBam_nos_ControlKD_68hAHS_repB_283080_sRBC_TGACT



sRNAseq_HsBam_nos_ControlKD_68hAHS_repC_283081_sRBC_AGCAA



sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT



position on HMS-Beagle2

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repB_283080_sRBC_CGTTTC

0
-25000
-50000

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repC_283081_sRBC_GGTAT

normalized read-counts

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repD_283082_sRBC_CGTTTC

0
-25000
-50000

position on HMS-Beagle2

sRNAseq_HsBam_nos_PiwiKD_68hAHS_repB_283080_sRBC_ATGGA

0
-25000
-50000

sRNAseq_HsBam_nos_PiwiKD_68hAHS_repC_283081_sRBC_TACCA

normalized read-counts

0
-25000
-50000

sRNAseq_HsBam_nos_PiwiKD_68hAHS_repD_283082_sRBC_ATGGA

0
-25000
-50000

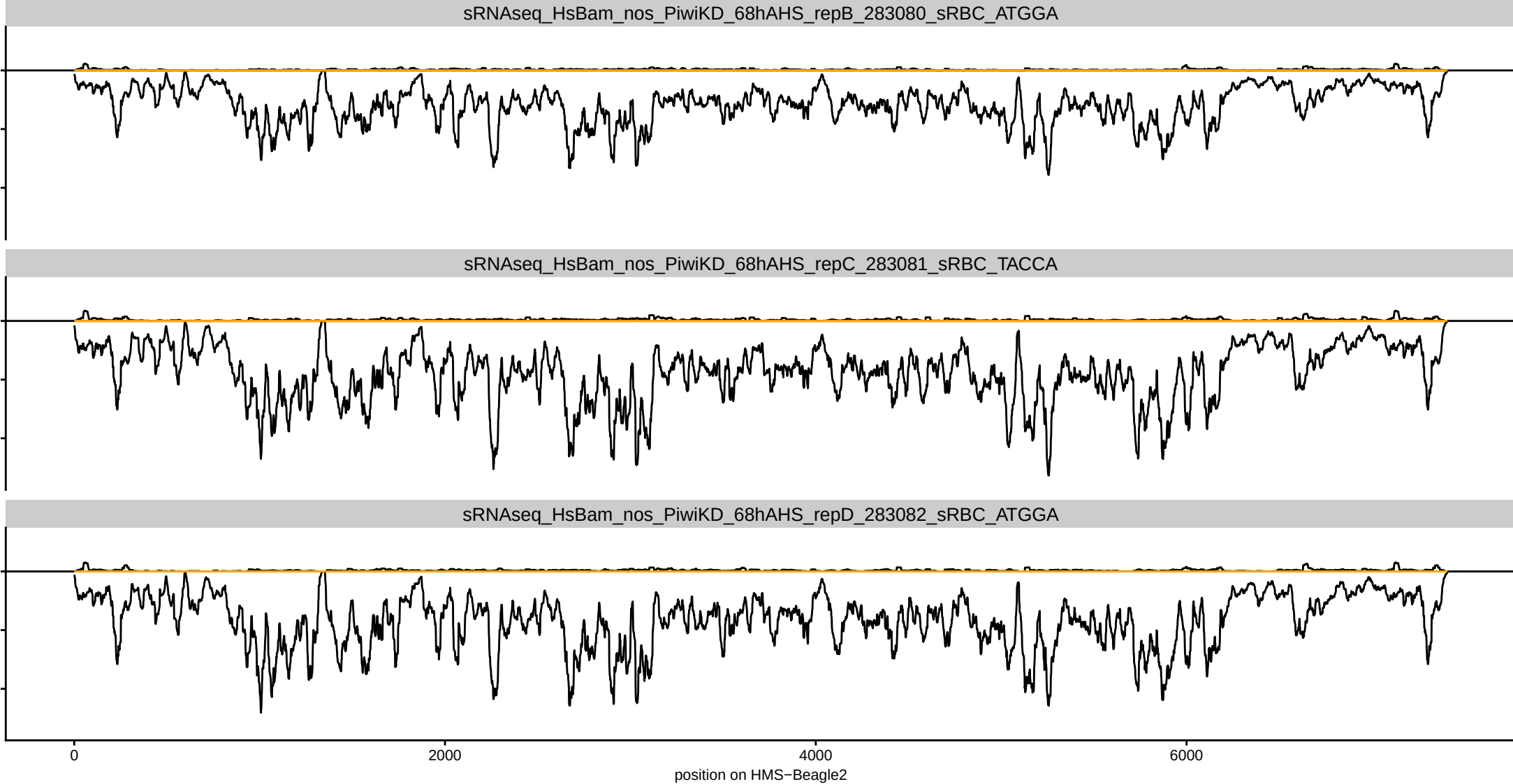
position on HMS-Beagle2

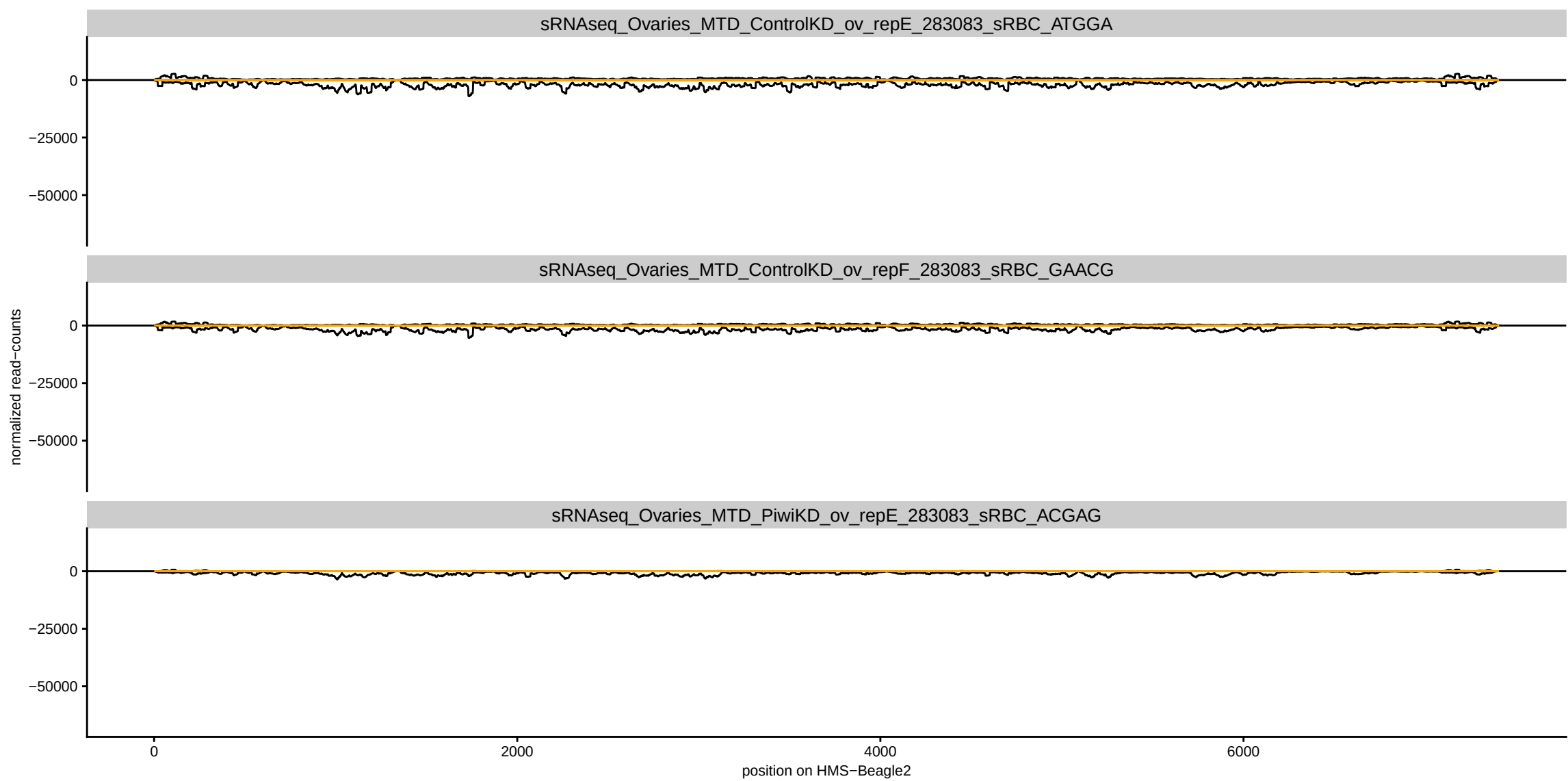
0

2000

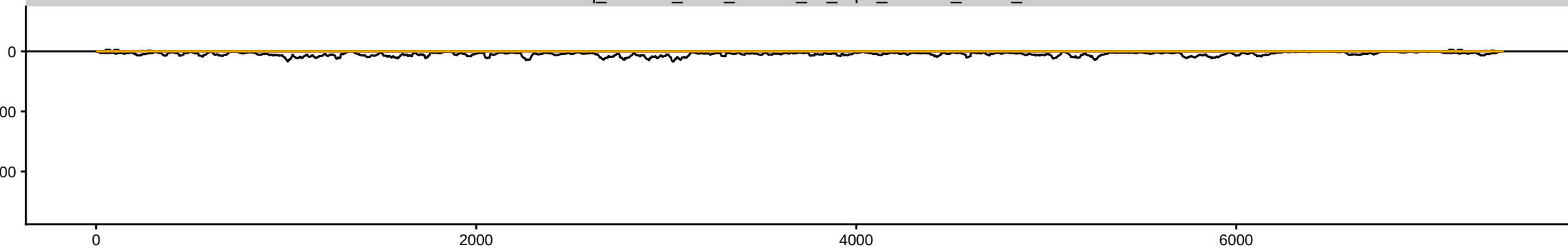
4000

6000





normalized read-counts



position on HMS-Beagle2