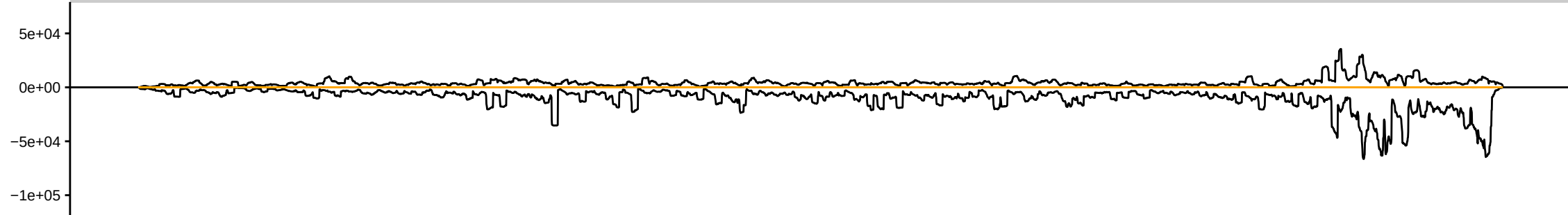
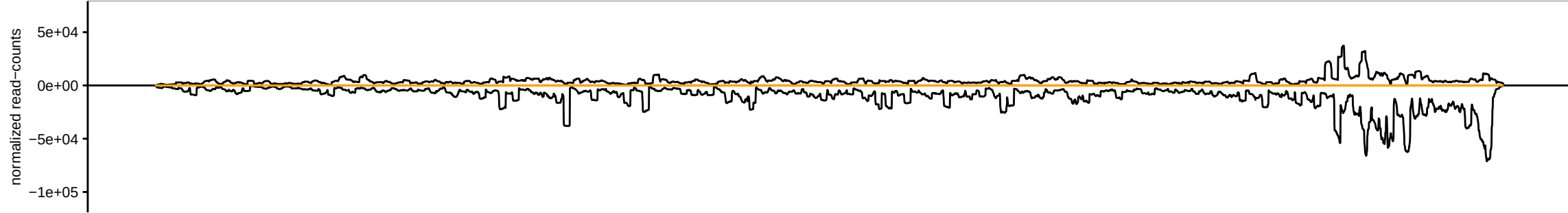


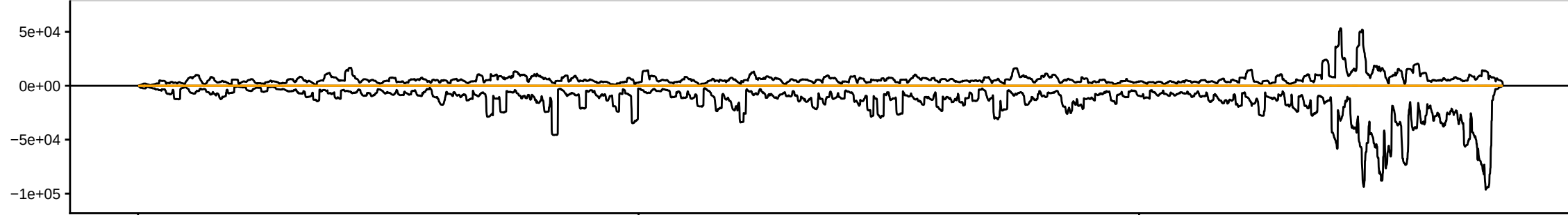
sRNAseq_HsBam_nos_ControlKD_00hAHS_repB_283080_sRBC_GTCAG



sRNAseq_HsBam_nos_ControlKD_00hAHS_repC_283081_sRBC_CAGTG



sRNAseq_HsBam_nos_ControlKD_00hAHS_repD_283082_sRBC_GTCAG



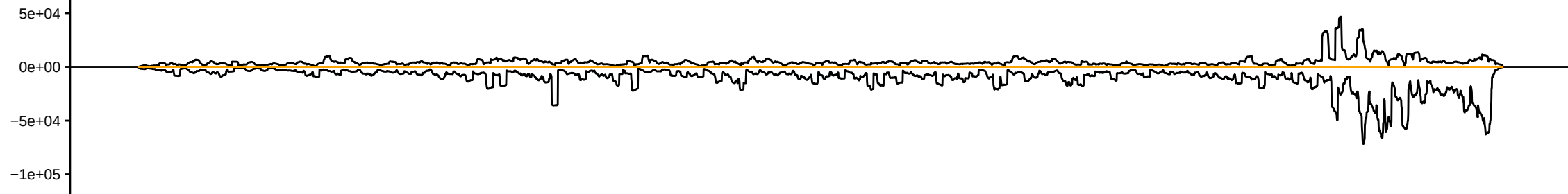
0

2000

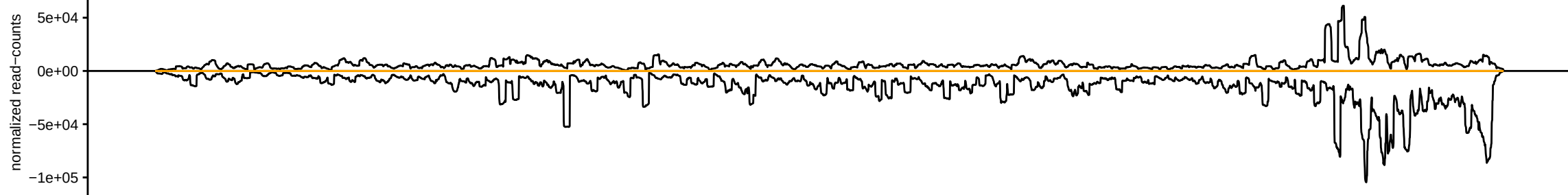
4000

position on baggins

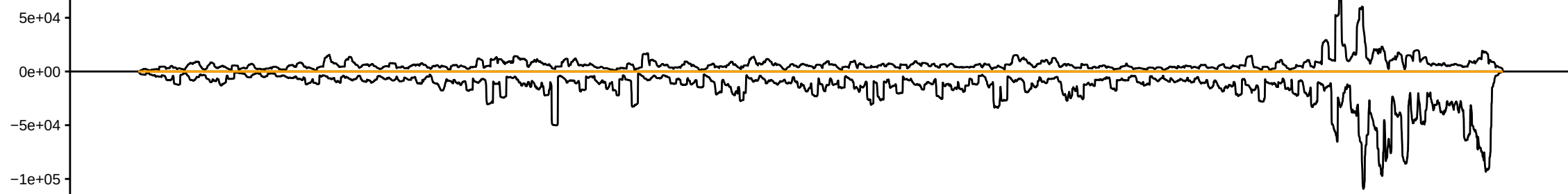
sRNAseq_HsBam_nos_ControlKD_68hAHS_repB_283080_sRBC_TGACT



sRNAseq_HsBam_nos_ControlKD_68hAHS_repC_283081_sRBC_AGCAA



sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT



0

2000

4000

position on baggins

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repB_283080_sRBC_CGTTC

5e+04
0e+00
-5e+04
-1e+05

normalized read-counts

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repC_283081_sRBC_GGTAT

5e+04
0e+00
-5e+04
-1e+05

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repD_283082_sRBC_CGTTC

5e+04
0e+00
-5e+04
-1e+05

0

2000

4000

position on baggins

