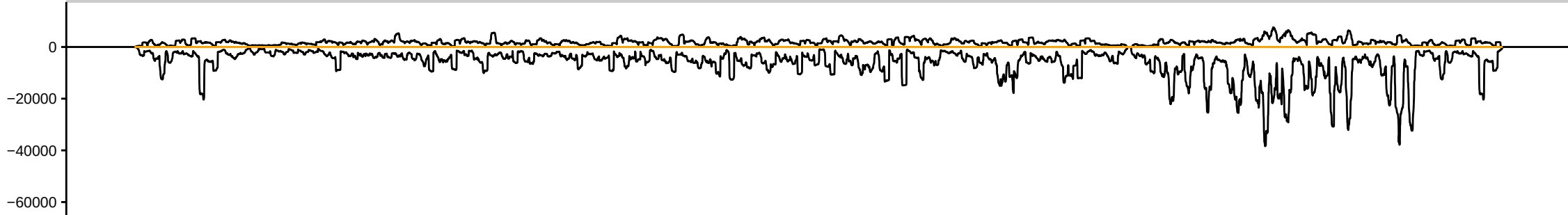
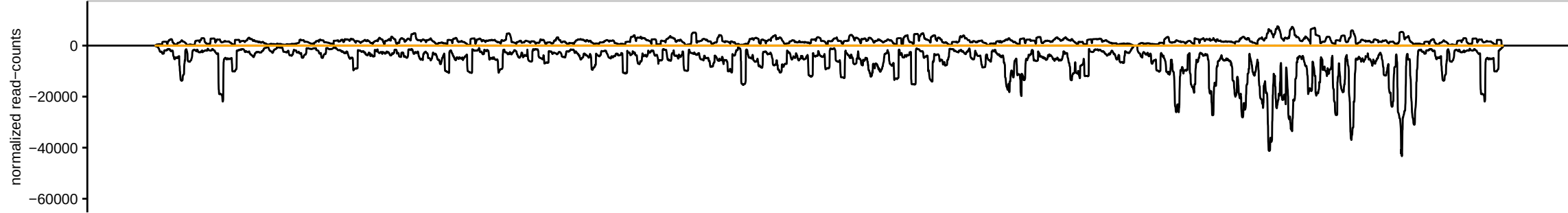


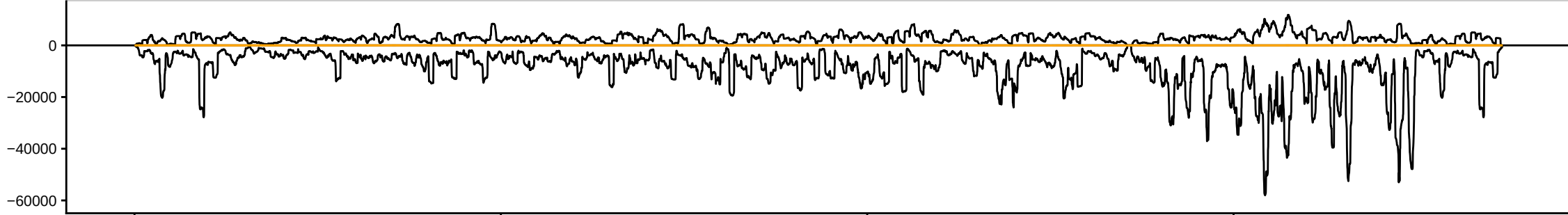
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sRNAseq_HsBam_nos_ControlKD_00hAHS_repC_283081_sRBC_CAGTG



sRNAseq_HsBam_nos_ControlKD_00hAHS_repD_283082_sRBC_GTCAG



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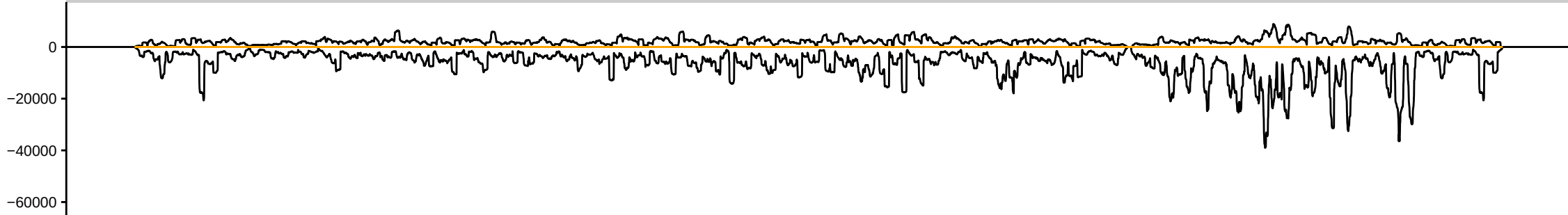
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4000

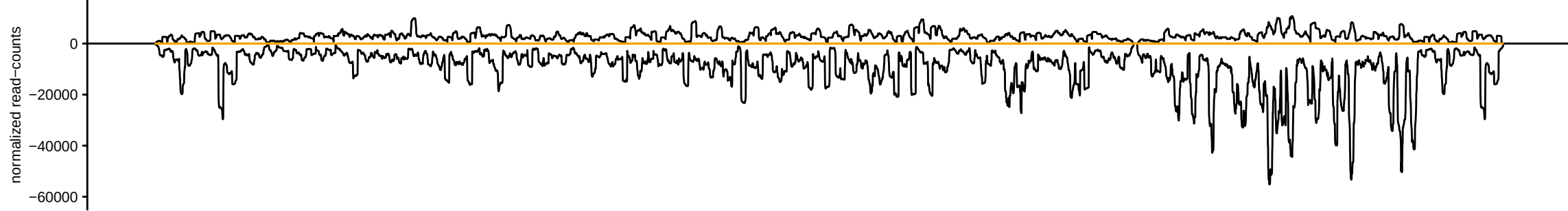
6000

position on gypsy

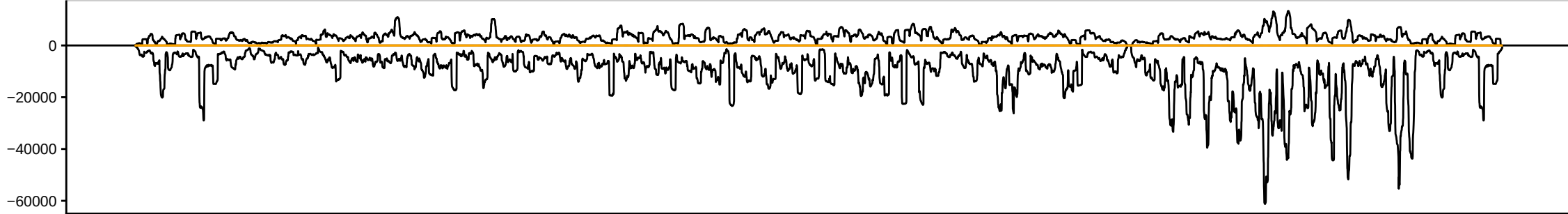
sRNAseq_HsBam_nos_ControlKD_68hAHS_repB_283080_sRBC_TGACT



sRNAseq_HsBam_nos_ControlKD_68hAHS_repC_283081_sRBC_AGCAA



sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT



0

2000

4000

6000

position on gypsy

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repB_283080_sRBC_CGTTC

0
-20000
-40000
-60000

normalized read-counts

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repC_283081_sRBC_GGTAT

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0
-20000
-40000
-60000

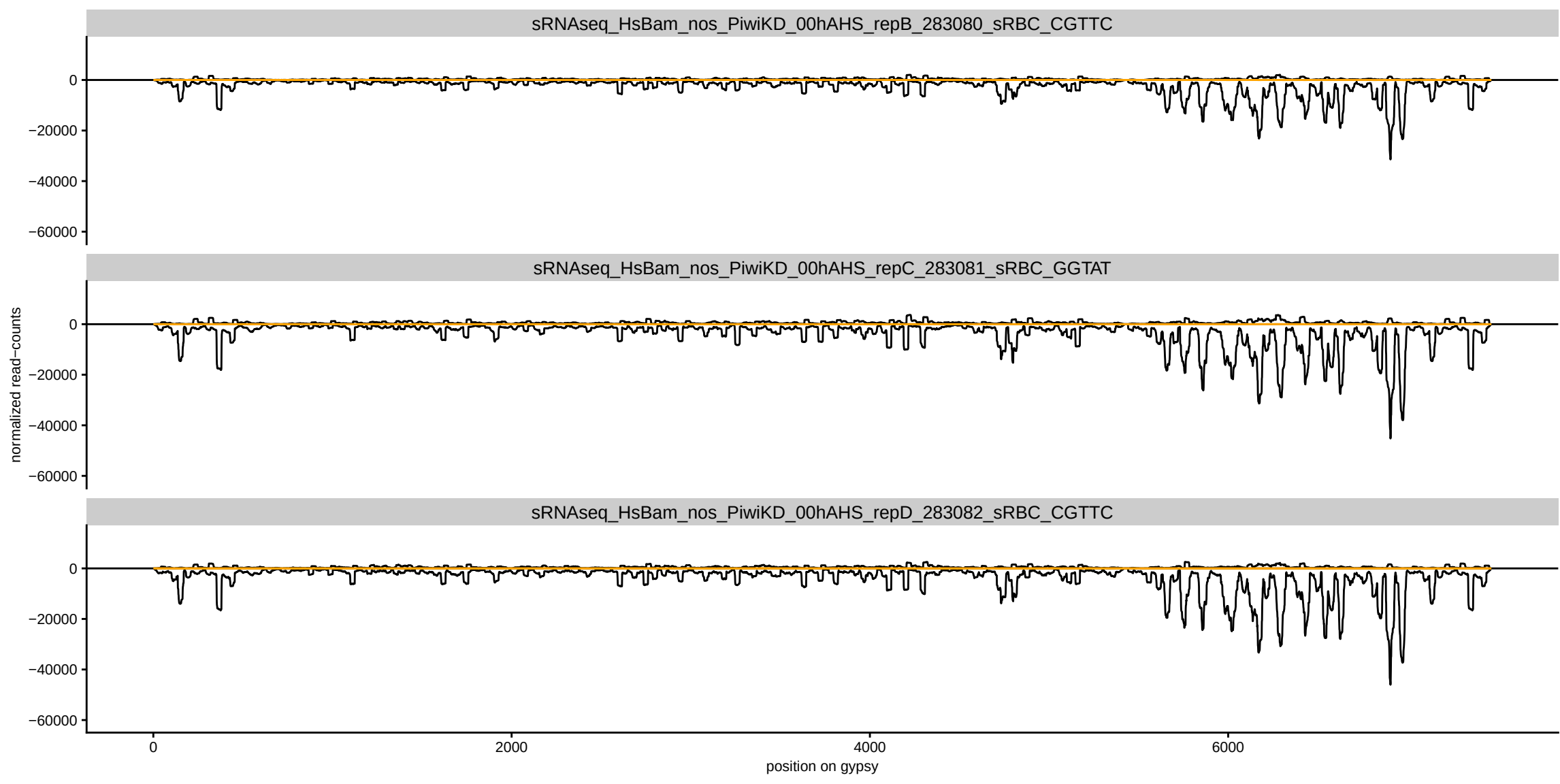
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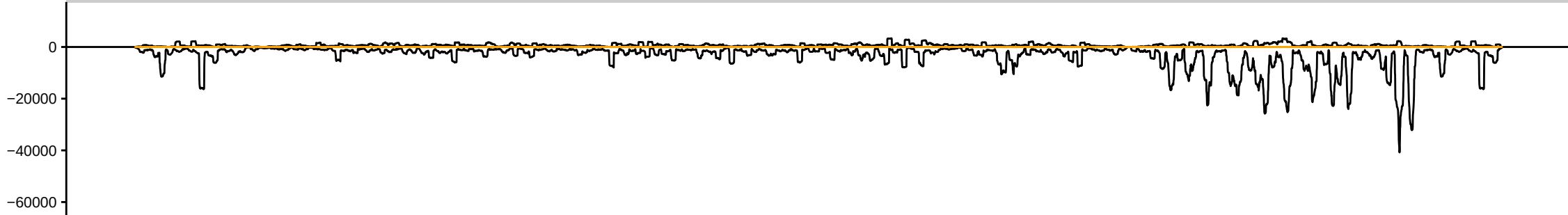
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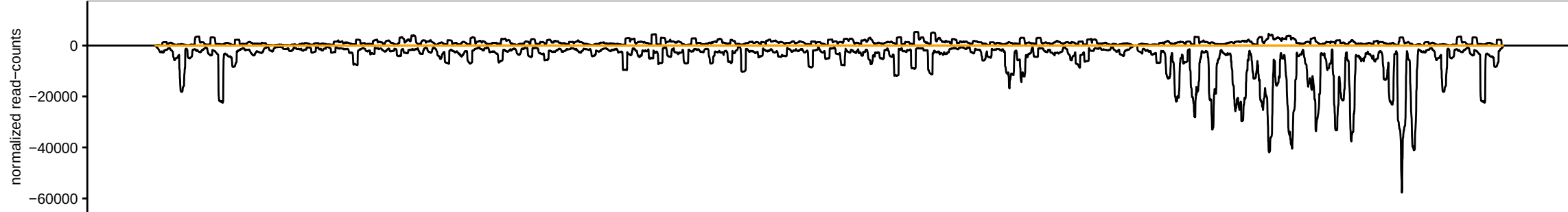
position on gypsy



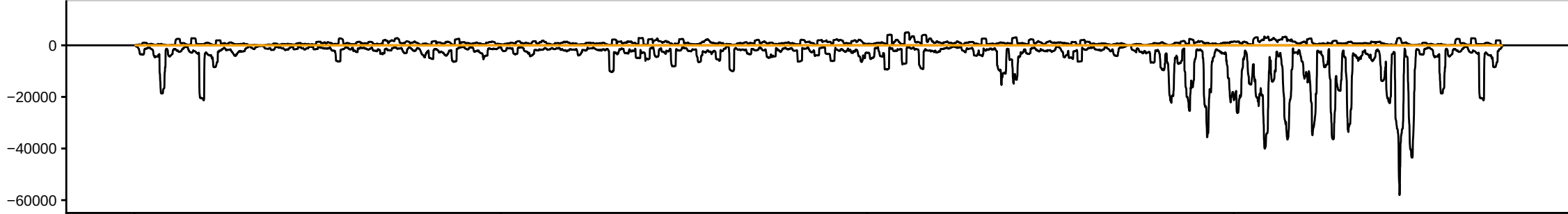
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sRNAseq_HsBam_nos_PiwiKD_68hAHS_repD_283082_sRBC_ATGGA



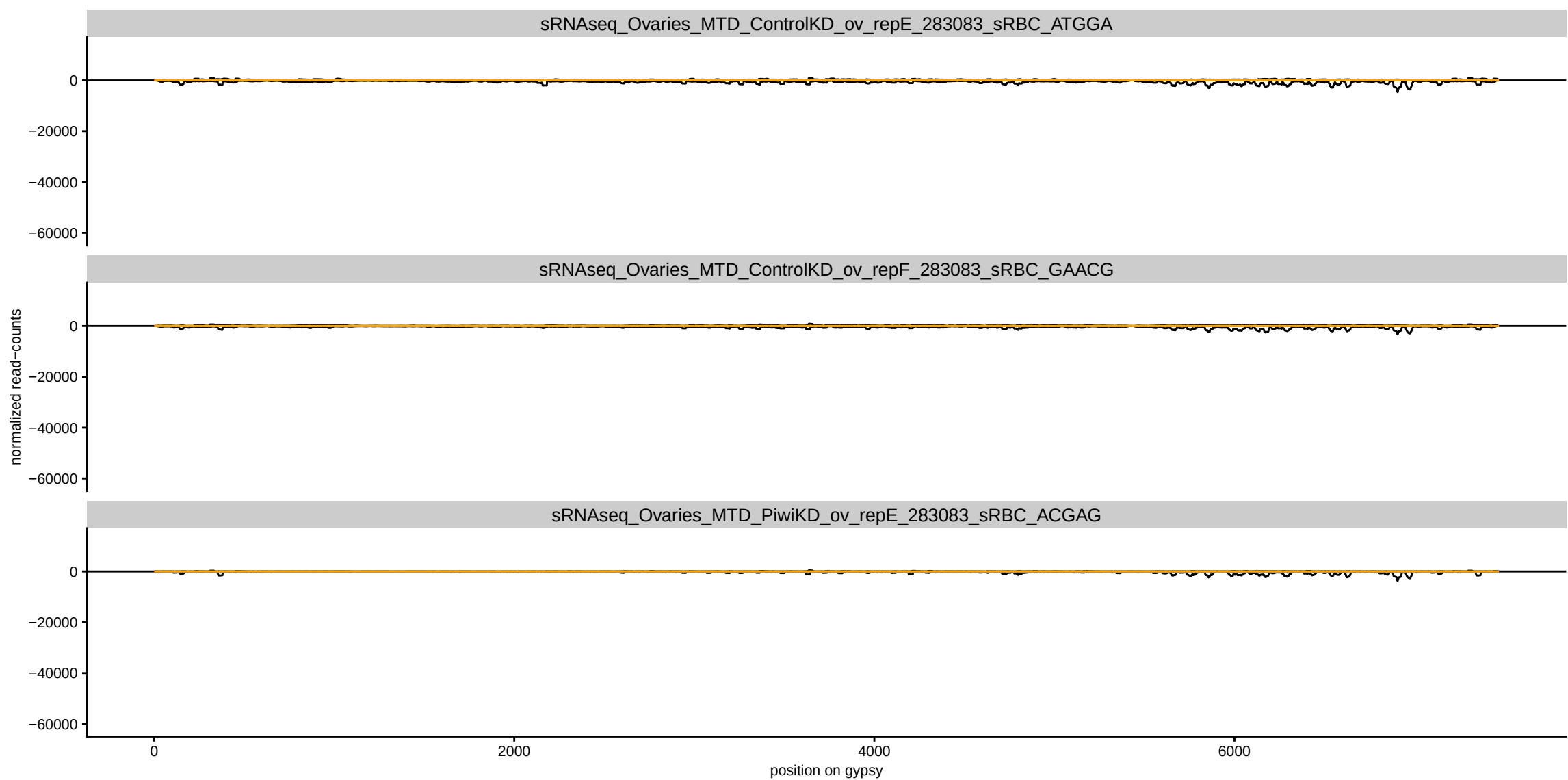
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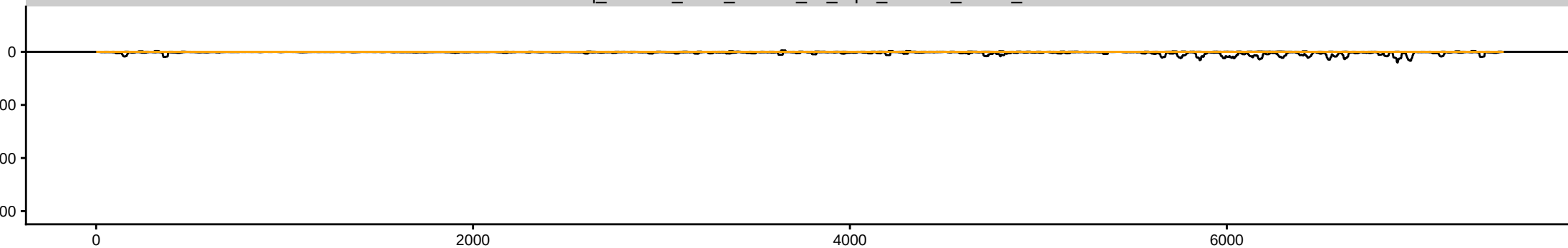
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6000

position on gypsy



normalized read-counts



position on gypsy