

sRNAseq_HsBam_nos_ControlKD_00hAHS_repB_283080_sRBC_GTCAG

20000
10000
0
-10000

0

2500

5000

7500

10000

position on gypsy12

sRNAseq_HsBam_nos_ControlKD_00hAHS_repC_283081_sRBC_CAGTG

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10000
0
-10000

0

2500

5000

7500

10000

position on gypsy12

sRNAseq_HsBam_nos_ControlKD_00hAHS_repD_283082_sRBC_GTCAG

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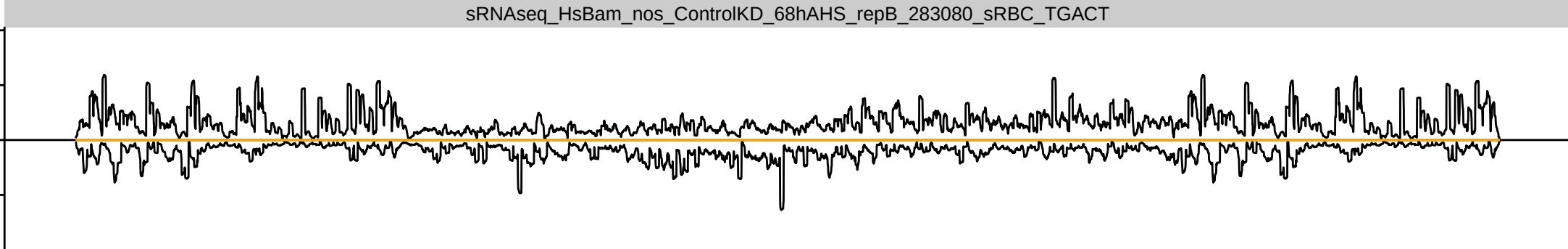
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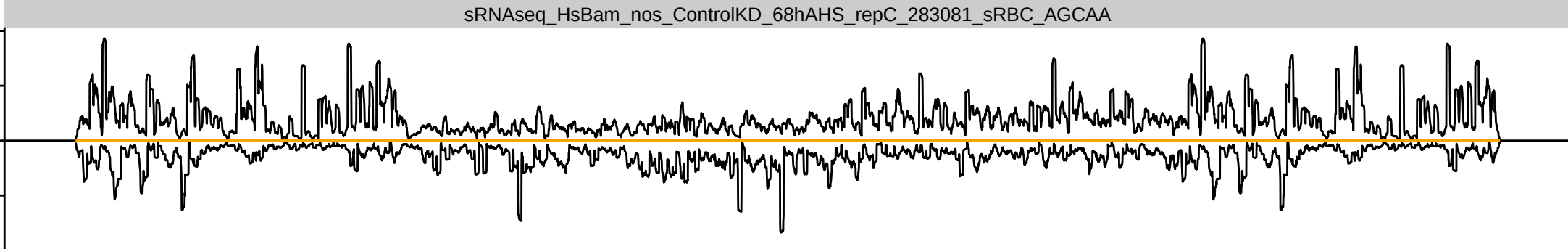
position on gypsy12

normalized read-counts

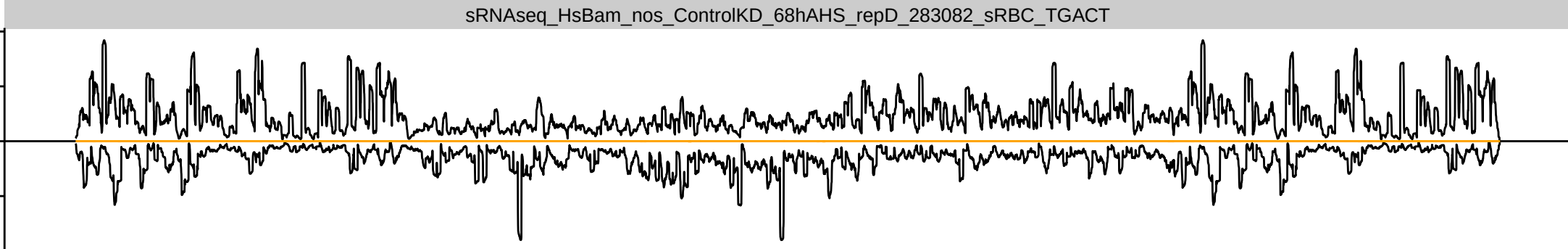
sRNAseq_HsBam_nos_ControlKD_68hAHS_repB_283080_sRBC_TGACT

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sRNAseq_HsBam_nos_ControlKD_68hAHS_repC_283081_sRBC_AGCAA

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0
-10000
normalized read-counts

sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT

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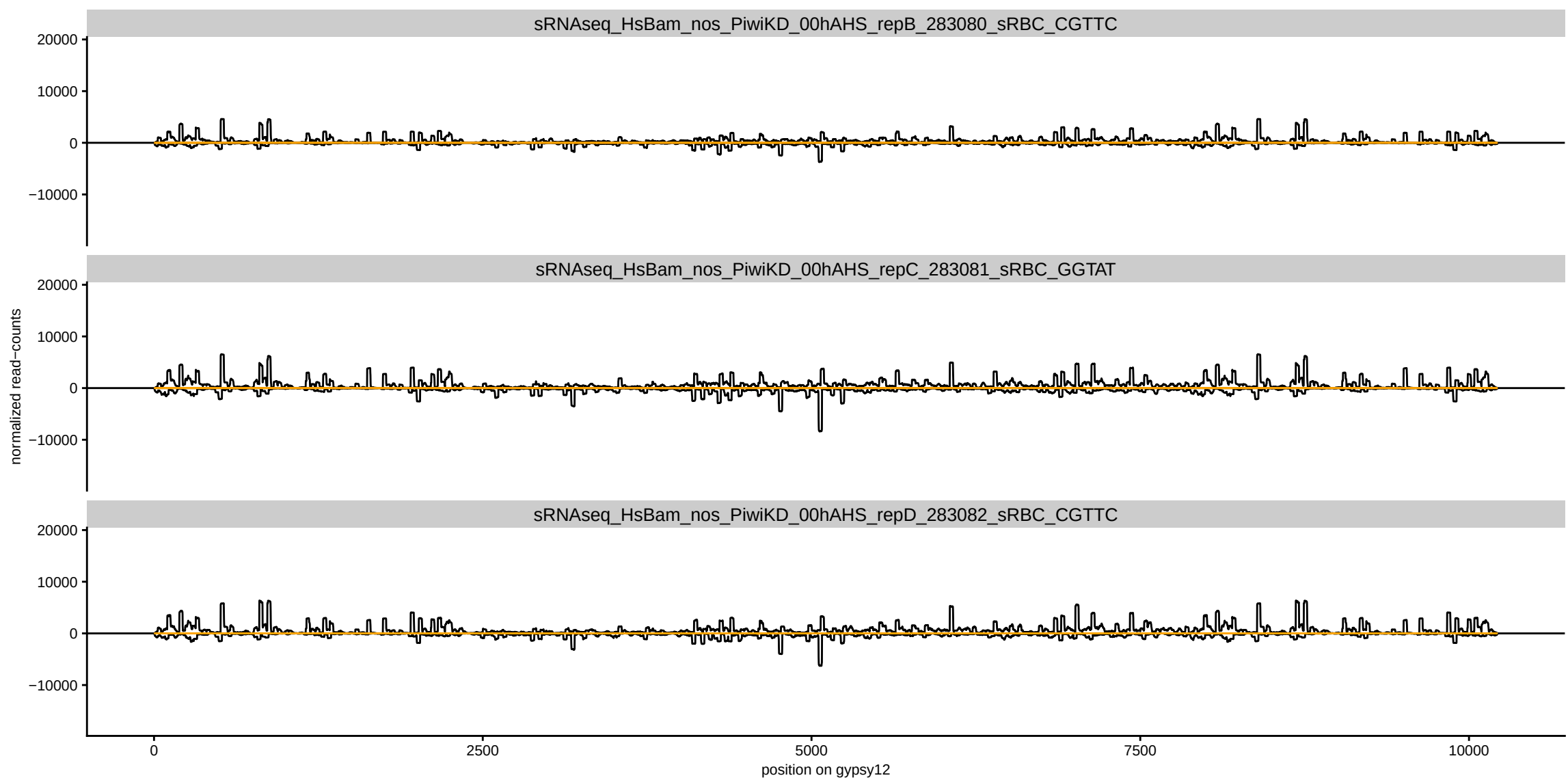
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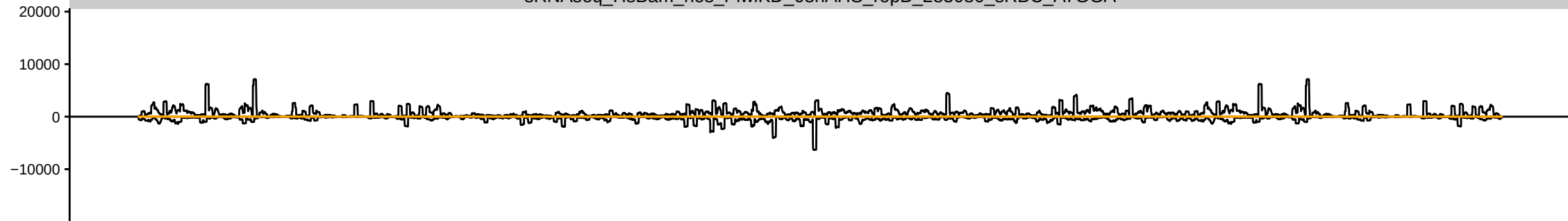
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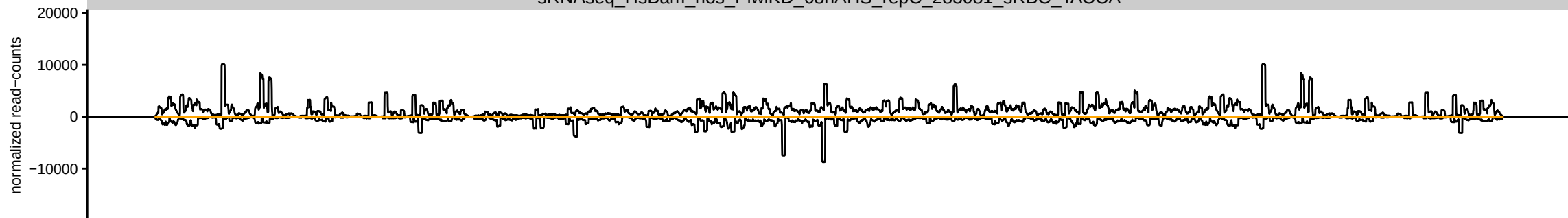
position on gypsy12



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

