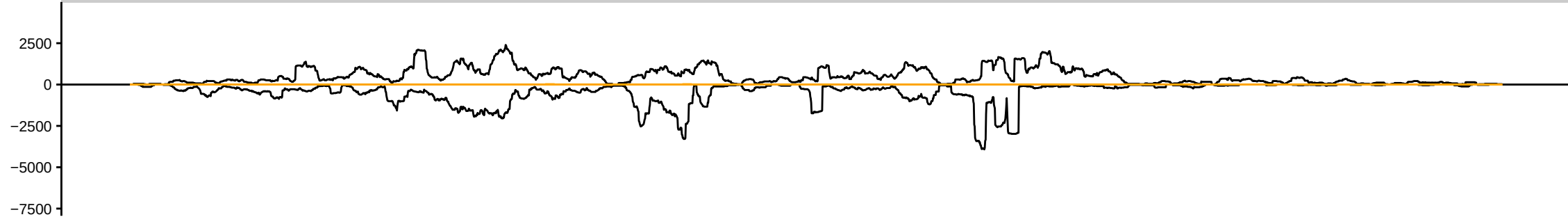
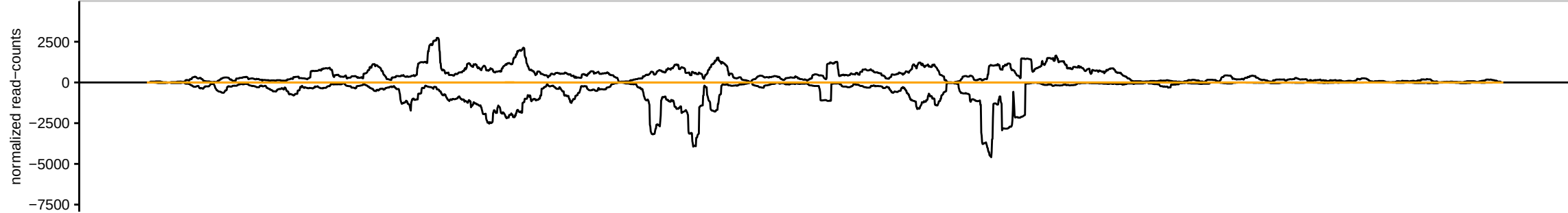


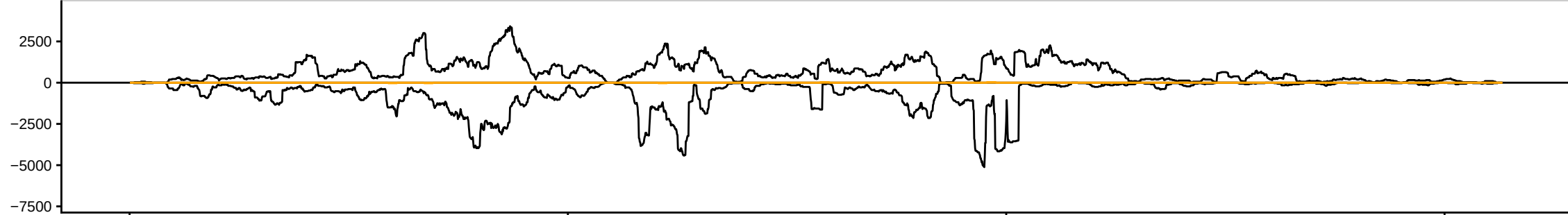
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sRNAseq_HsBam_nos_ControlKD_00hAHS_repC_283081_sRBC_CAGTG



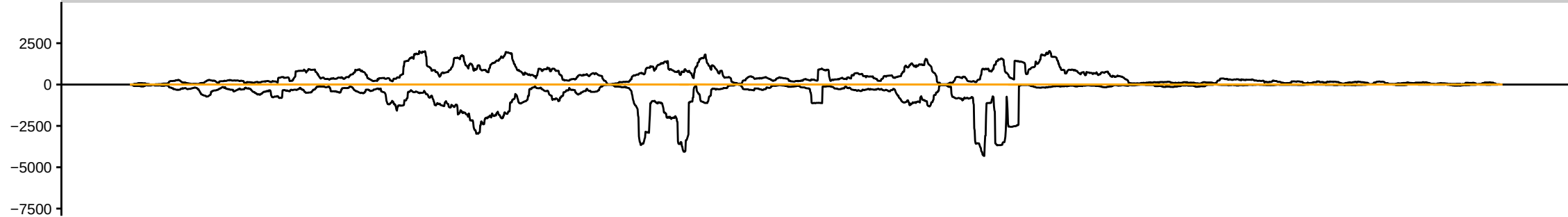
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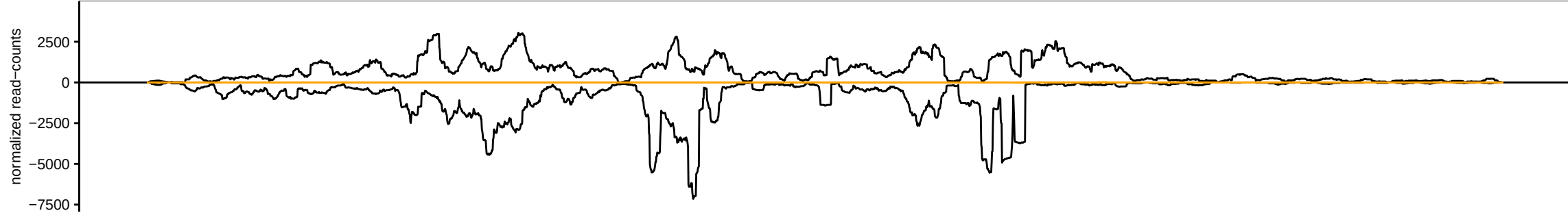
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position on Fw3

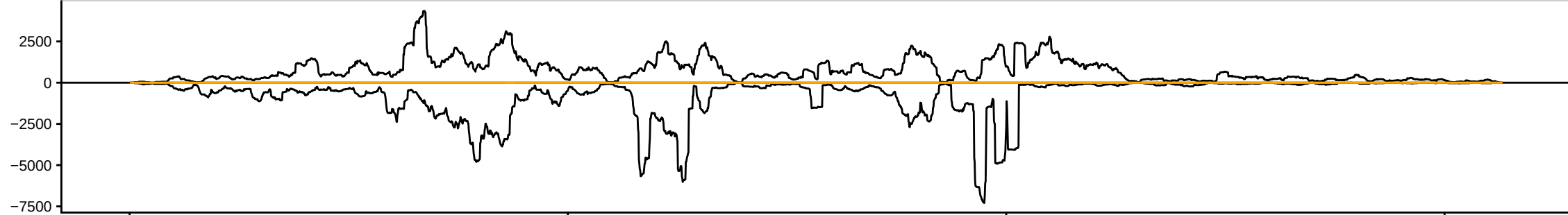
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sRNAseq_HsBam_nos_ControlKD_68hAHS_repC_283081_sRBC_AGCAA



sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT



0 1000 2000 3000

position on Fw3

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repB_283080_sRBC_CGTTC

2500
0
-2500
-5000
-7500

normalized read-counts

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repC_283081_sRBC_GGTAT

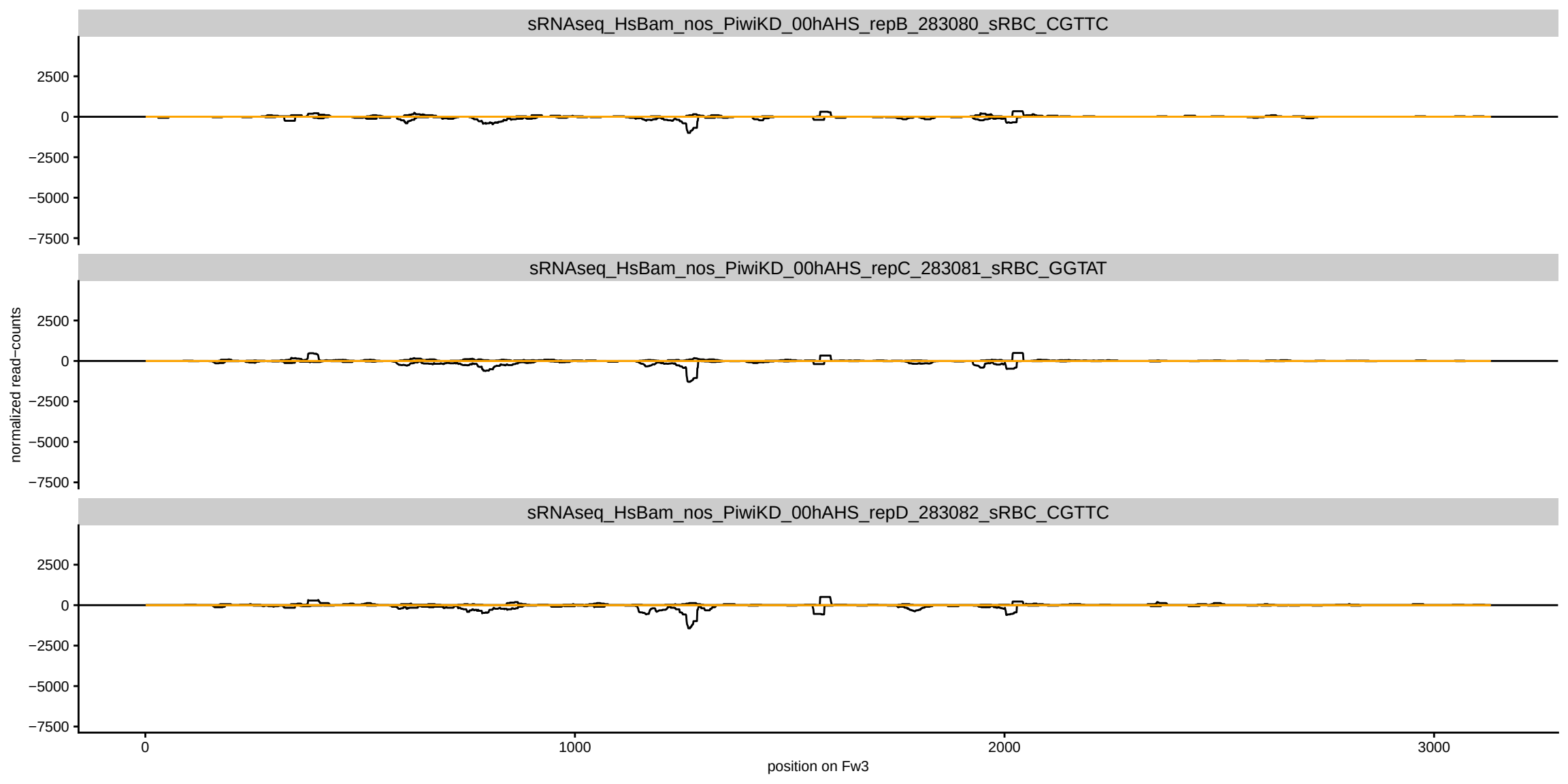
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-2500
-5000
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sRNAseq_HsBam_nos_PiwiKD_00hAHS_repD_283082_sRBC_CGTTC

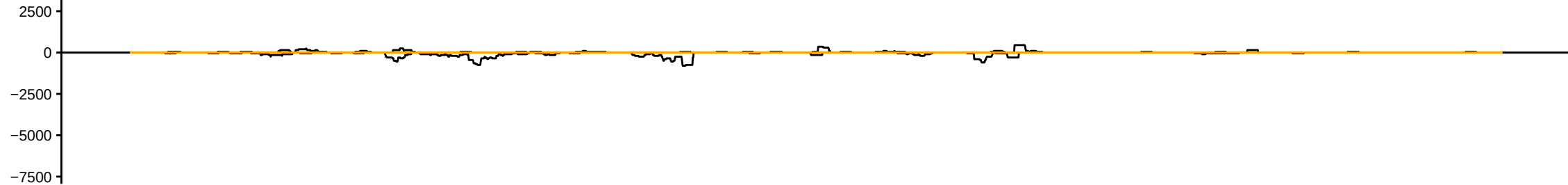
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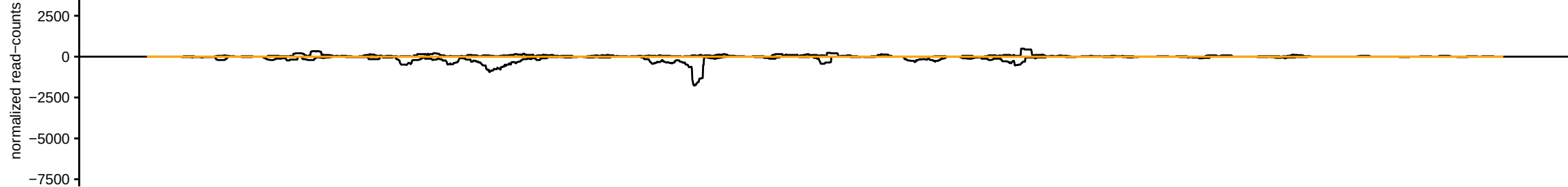
position on Fw3



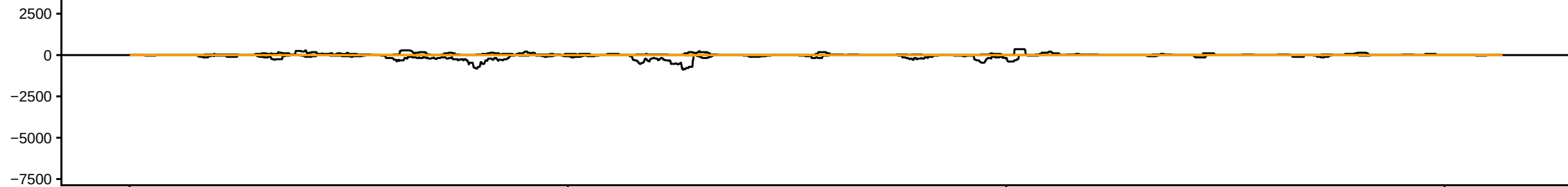
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sRNAseq_HsBam_nos_PiwiKD_68hAHS_repC_283081_sRBC_TACCA



sRNAseq_HsBam_nos_PiwiKD_68hAHS_repD_283082_sRBC_ATGGA



position on Fw3

