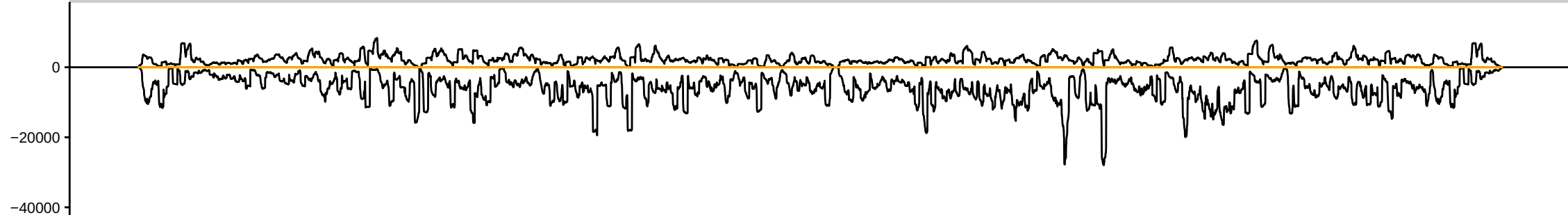
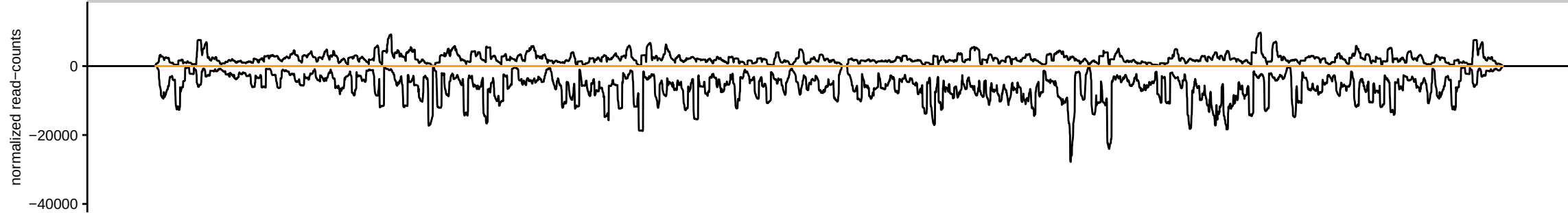


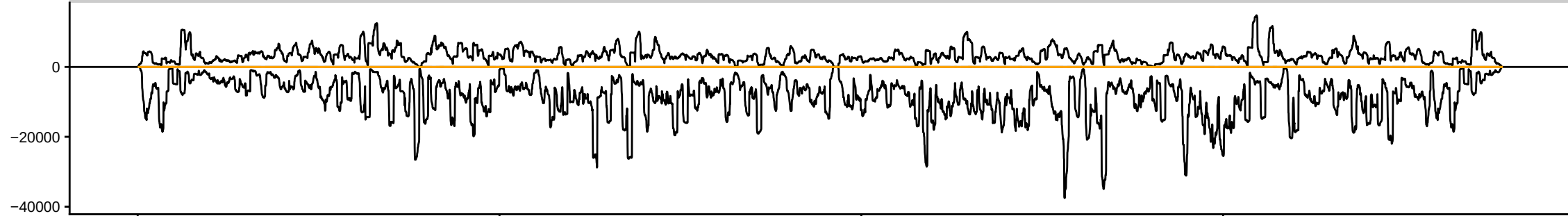
sRNAseq_HsBam_nos_ControlKD_00hAHS_repB_283080_sRBC_GTCAG



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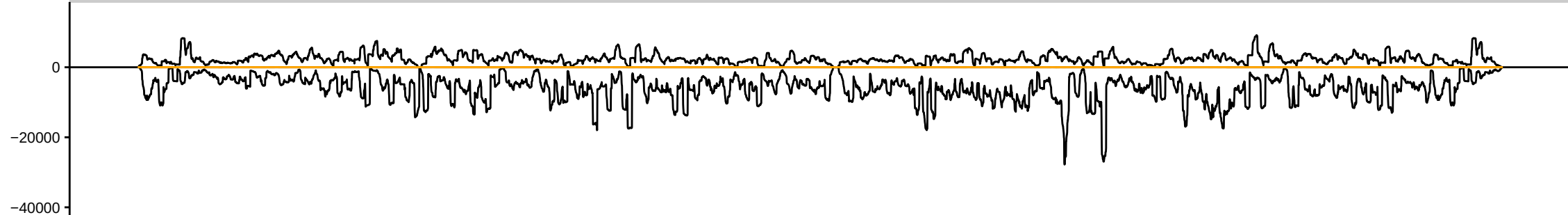


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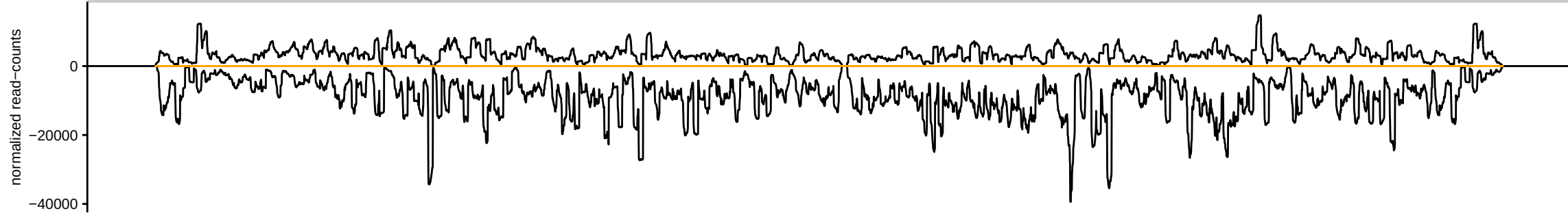


position on springer

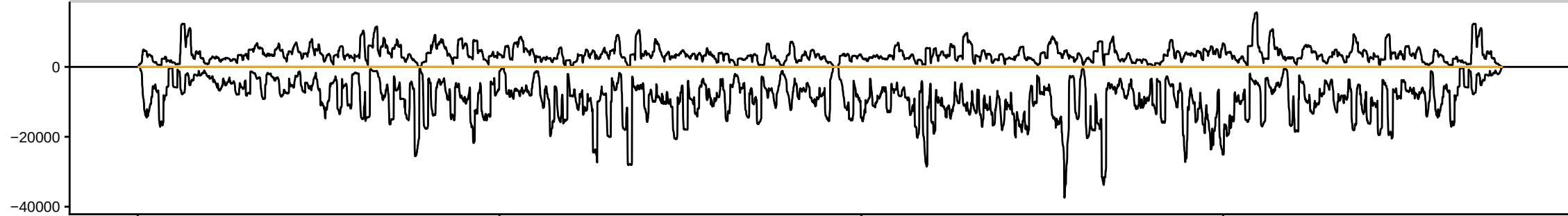
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sRNAseq_HsBam_nos_ControlKD_68hAHS_repC_283081_sRBC_AGCAA

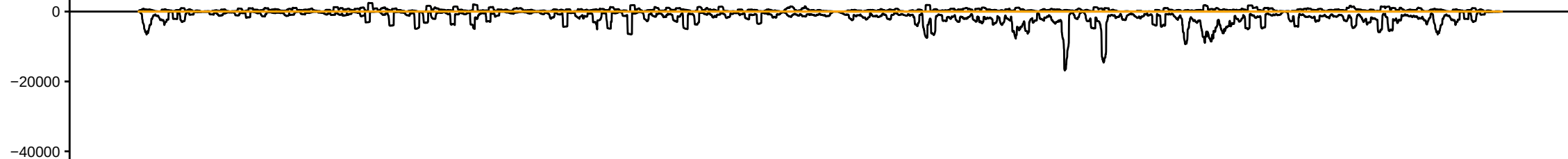


sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT

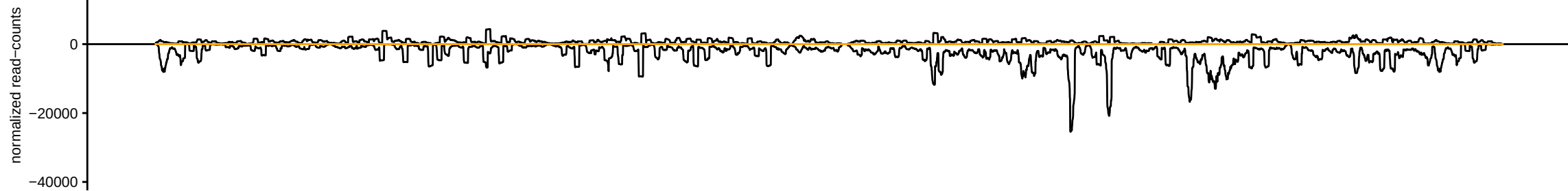


position on springer

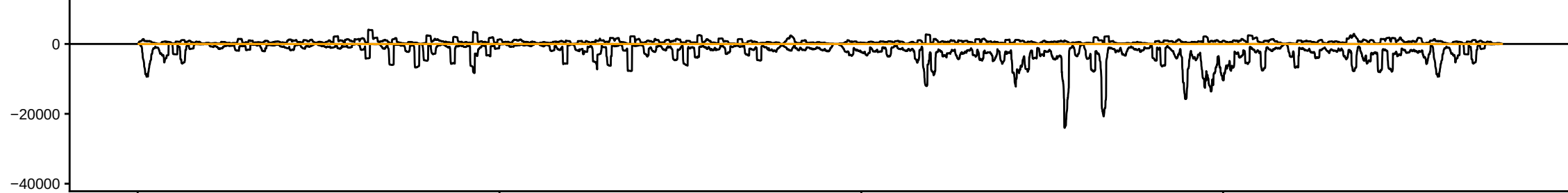
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sRNAseq_HsBam_nos_PiwiKD_00hAHS_repC_283081_sRBC_GGTAT

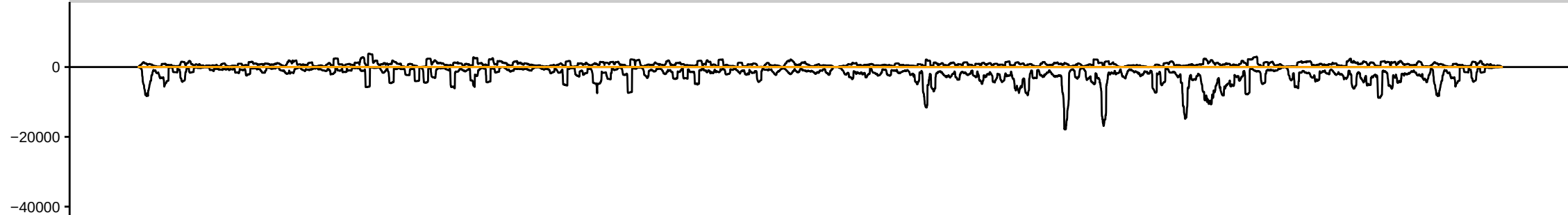


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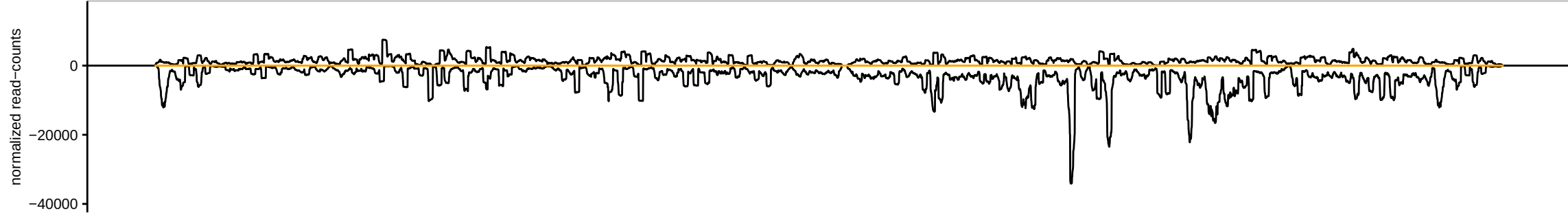


position on springer

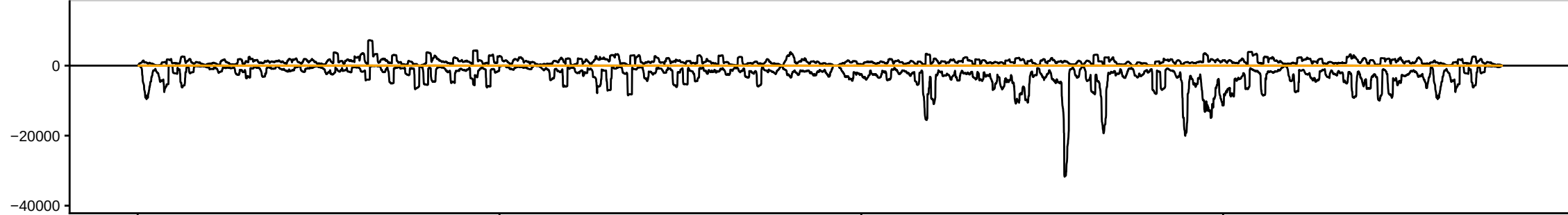
sRNAseq_HsBam_nos_PiwiKD_68hAHS_repB_283080_sRBC_ATGGA



sRNAseq_HsBam_nos_PiwiKD_68hAHS_repC_283081_sRBC_TACCA



sRNAseq_HsBam_nos_PiwiKD_68hAHS_repD_283082_sRBC_ATGGA



position on springer

sRNAseq_Ovaries_MTD_ControlKD_ov_repE_283083_sRBC_ATGGA

0
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-40000

normalized read-counts

sRNAseq_Ovaries_MTD_ControlKD_ov_repF_283083_sRBC_GAACG

0
-20000
-40000

sRNAseq_Ovaries_MTD_PiwiKD_ov_repE_283083_sRBC_ACGAG

0
-20000
-40000

0

2000

4000

6000

position on springer

