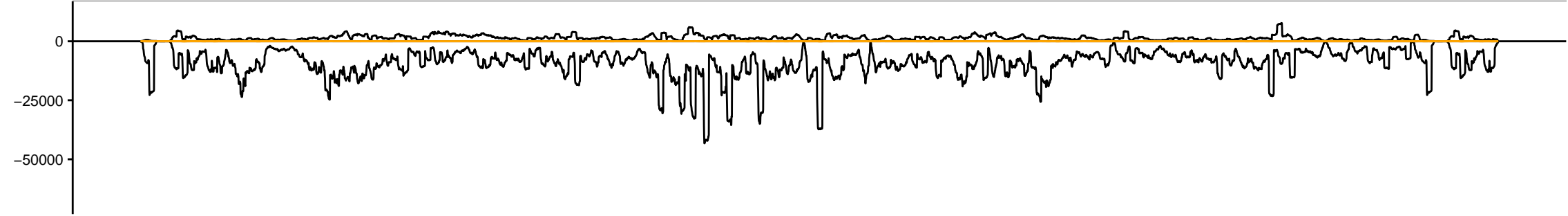
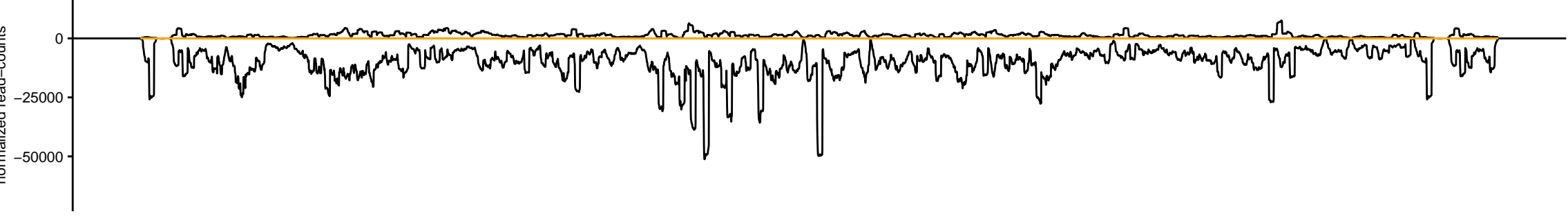


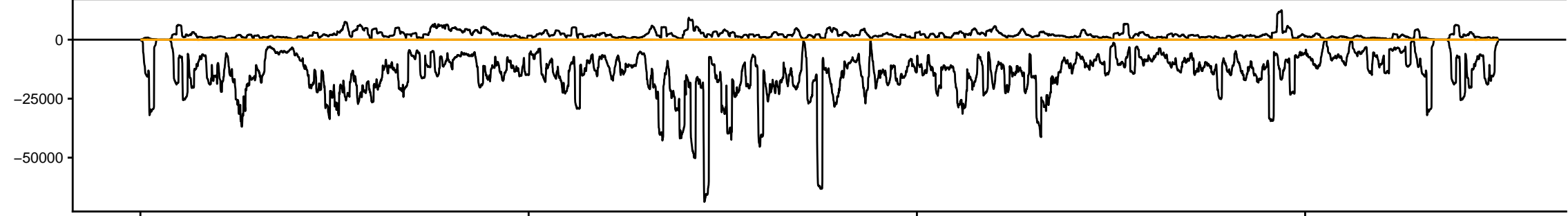
sRNAseq_HsBam_nos_ControlKD_00hAHS_repB_283080_sRBC_GTCAG



sRNAseq_HsBam_nos_ControlKD_00hAHS_repC_283081_sRBC_CAGTG

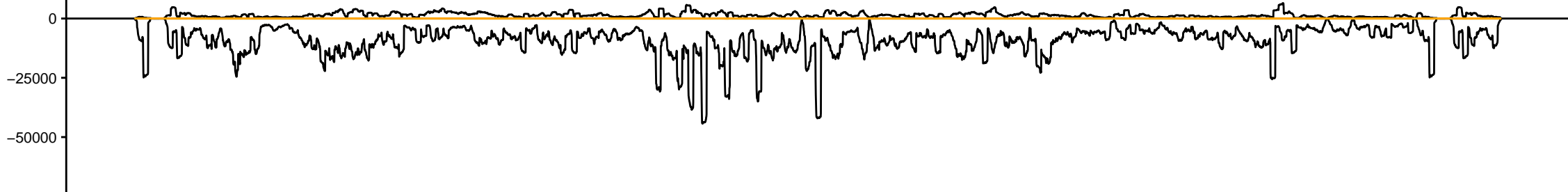


sRNAseq_HsBam_nos_ControlKD_00hAHS_repD_283082_sRBC_GTCAG

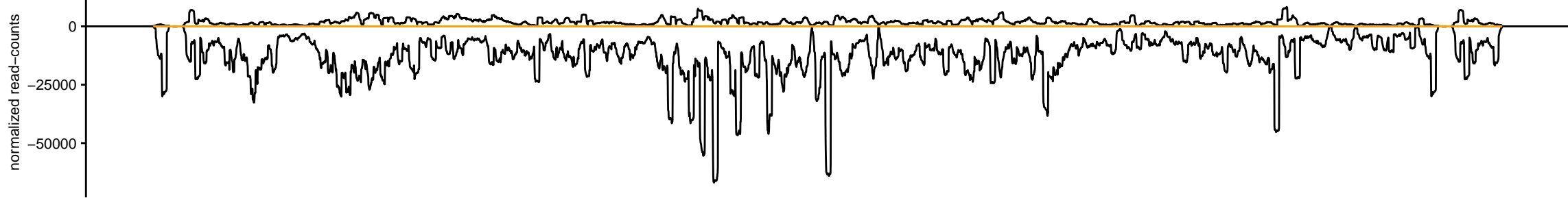


position on 297

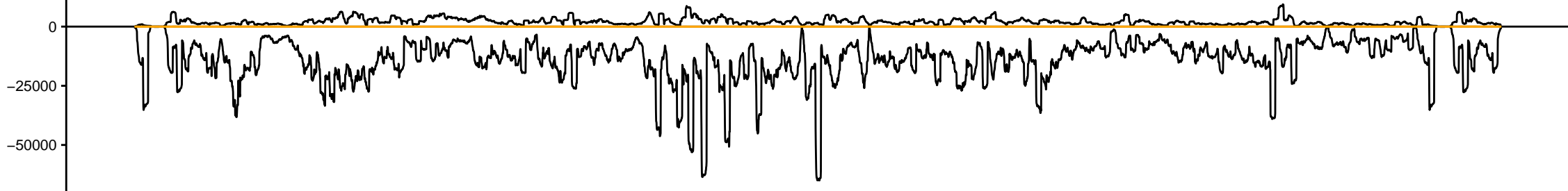
sRNAseq_HsBam_nos_ControlKD_68hAHS_repB_283080_sRBC_TGACT



sRNAseq_HsBam_nos_ControlKD_68hAHS_repC_283081_sRBC_AGCAA

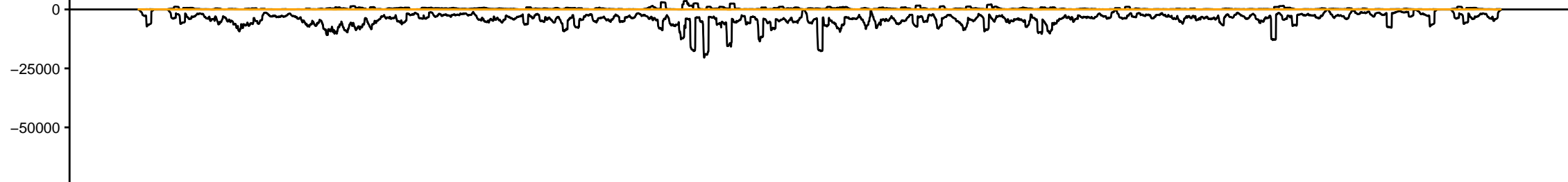


sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT

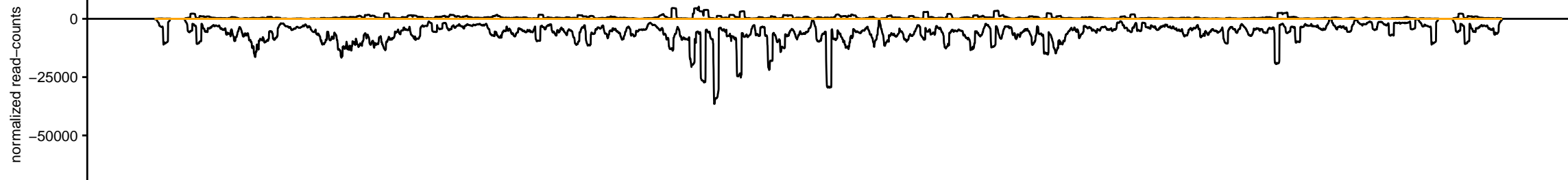


position on 297

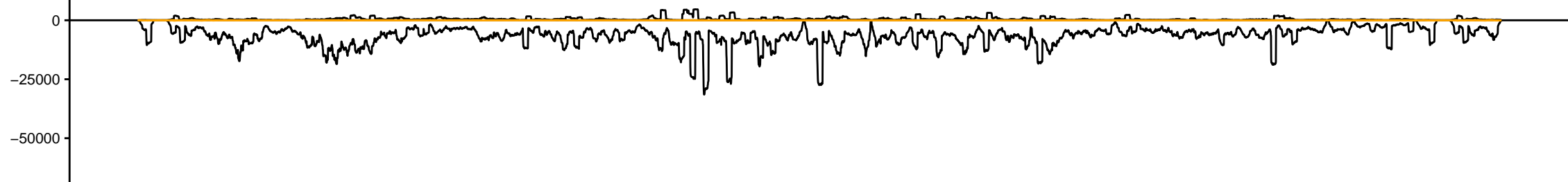
sRNAseq_HsBam_nos_PiwiKD_00hAHS_repB_283080_sRBC_CGTTTC



sRNAseq_HsBam_nos_PiwiKD_00hAHS_repC_283081_sRBC_GGTAT



sRNAseq_HsBam_nos_PiwiKD_00hAHS_repD_283082_sRBC_CGTTTC



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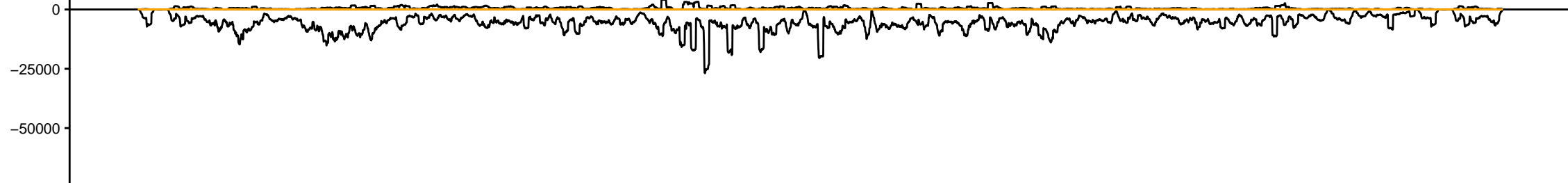
2000

4000

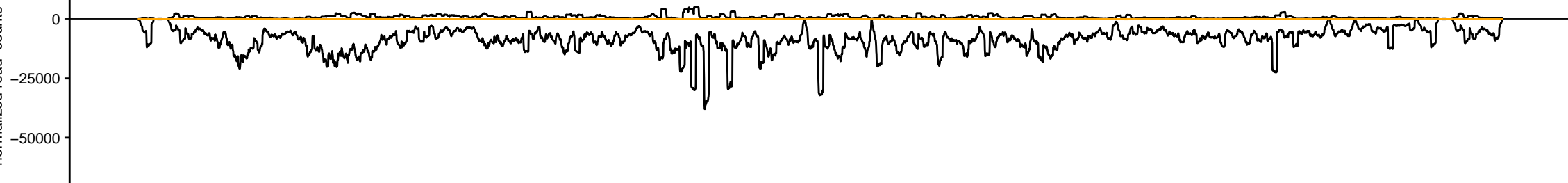
6000

position on 297

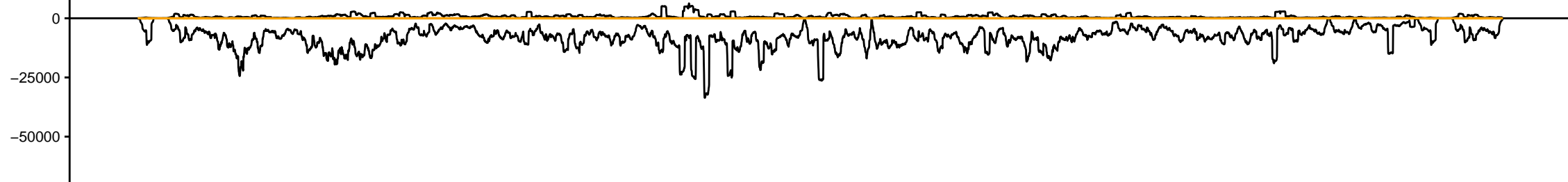
sRNAseq_HsBam_nos_PiwiKD_68hAHS_repB_283080_sRBC_ATGGA



sRNAseq_HsBam_nos_PiwiKD_68hAHS_repC_283081_sRBC_TACCA

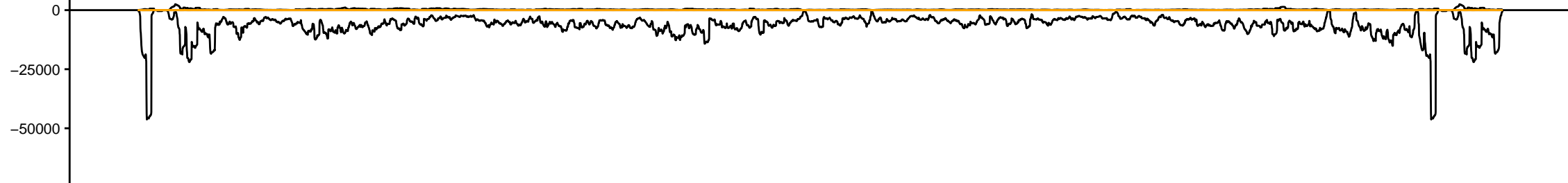


sRNAseq_HsBam_nos_PiwiKD_68hAHS_repD_283082_sRBC_ATGGA

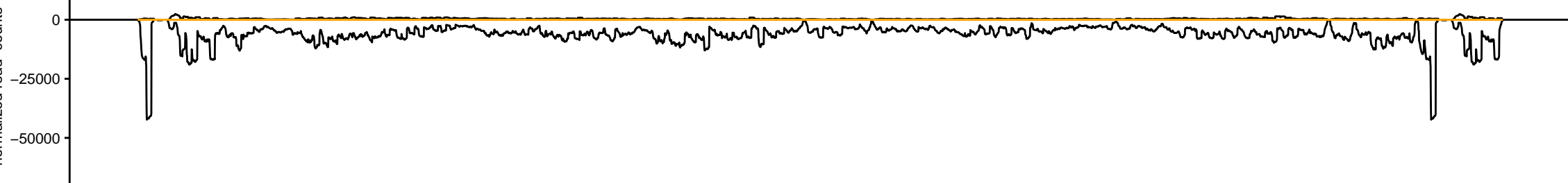


position on 297

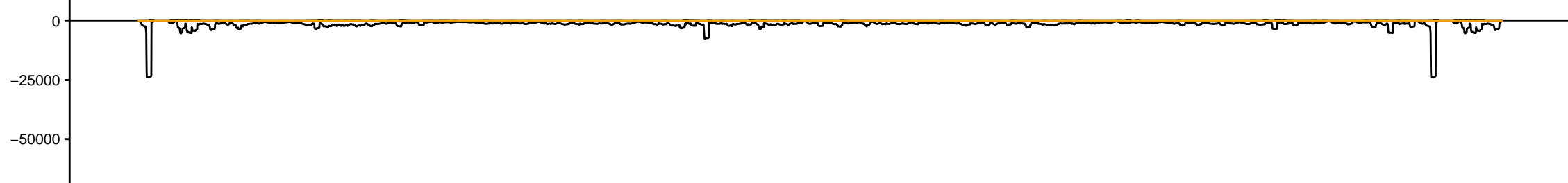
sRNAseq_Ovaries_MTD_ControlKD_ov_repE_283083_sRBC_ATGGA



sRNAseq_Ovaries_MTD_ControlKD_ov_repF_283083_sRBC_GAACG



sRNAseq_Ovaries_MTD_PiwiKD_ov_repE_283083_sRBC_ACGAG



position on 297

normalized read-counts

