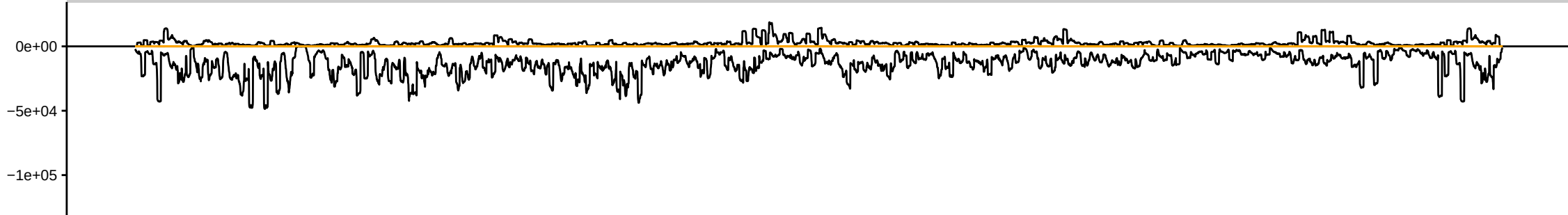
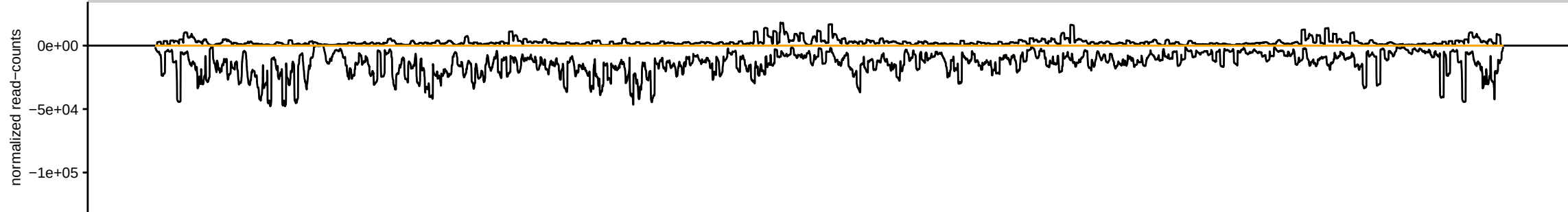


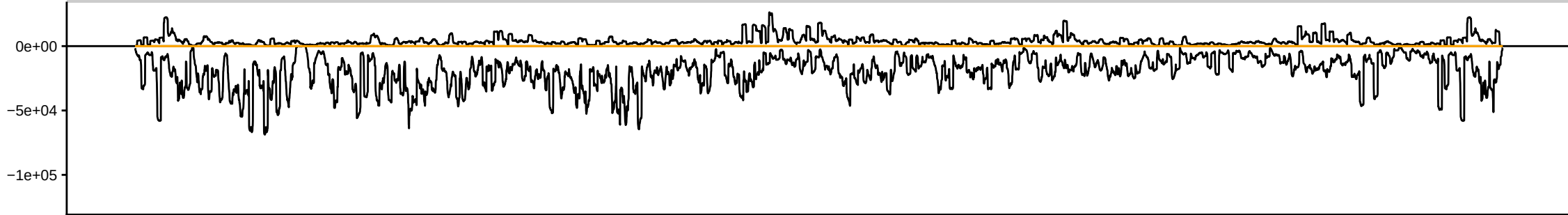
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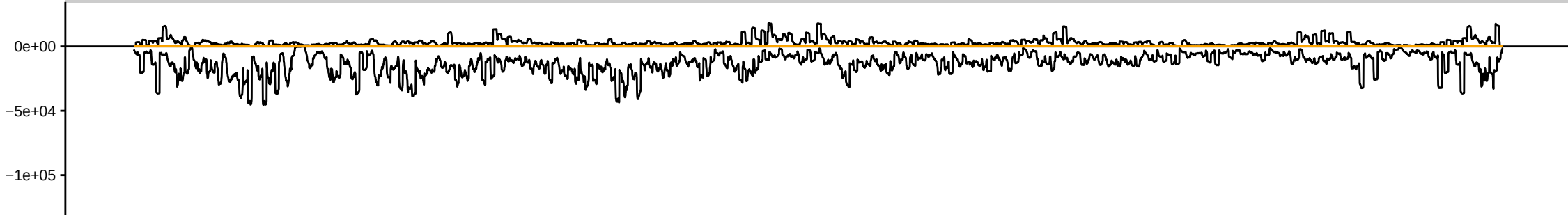
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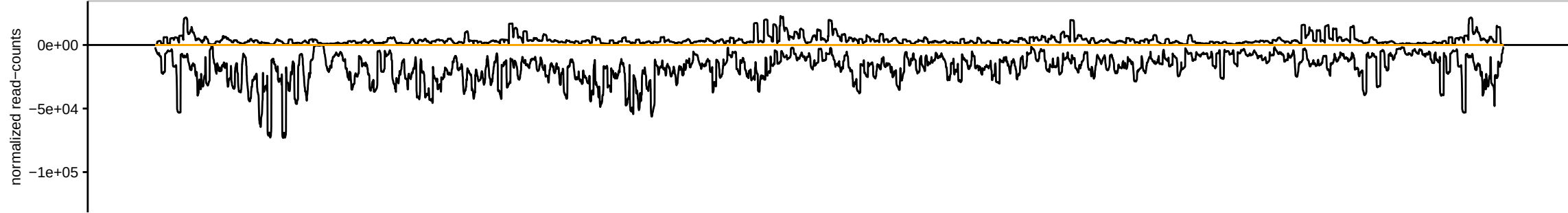
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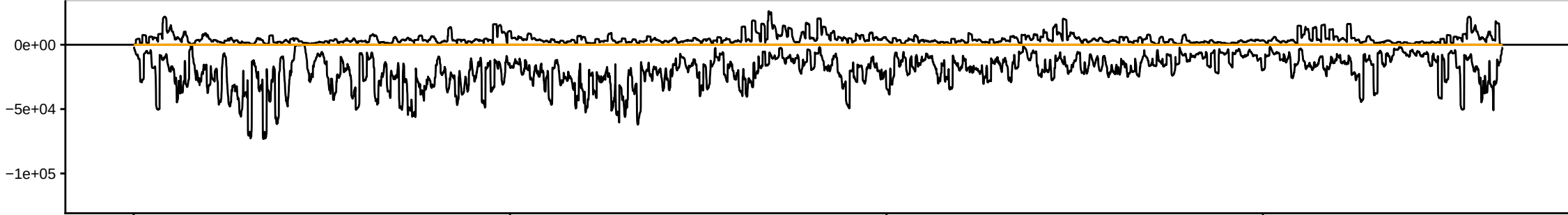
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sRNAseq_HsBam_nos_ControlKD_68hAHS_repD_283082_sRBC_TGACT



0

2500

5000

7500

position on roo

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repB_283080_sRBC_CGTTC

0e+00

-5e+04

-1e+05

normalized read-counts

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repC_283081_sRBC_GGTAT

0e+00

-5e+04

-1e+05

sRNAseq_HsBam_nos_PiwiKD_00hAHS_repD_283082_sRBC_CGTTC

0e+00

-5e+04

-1e+05

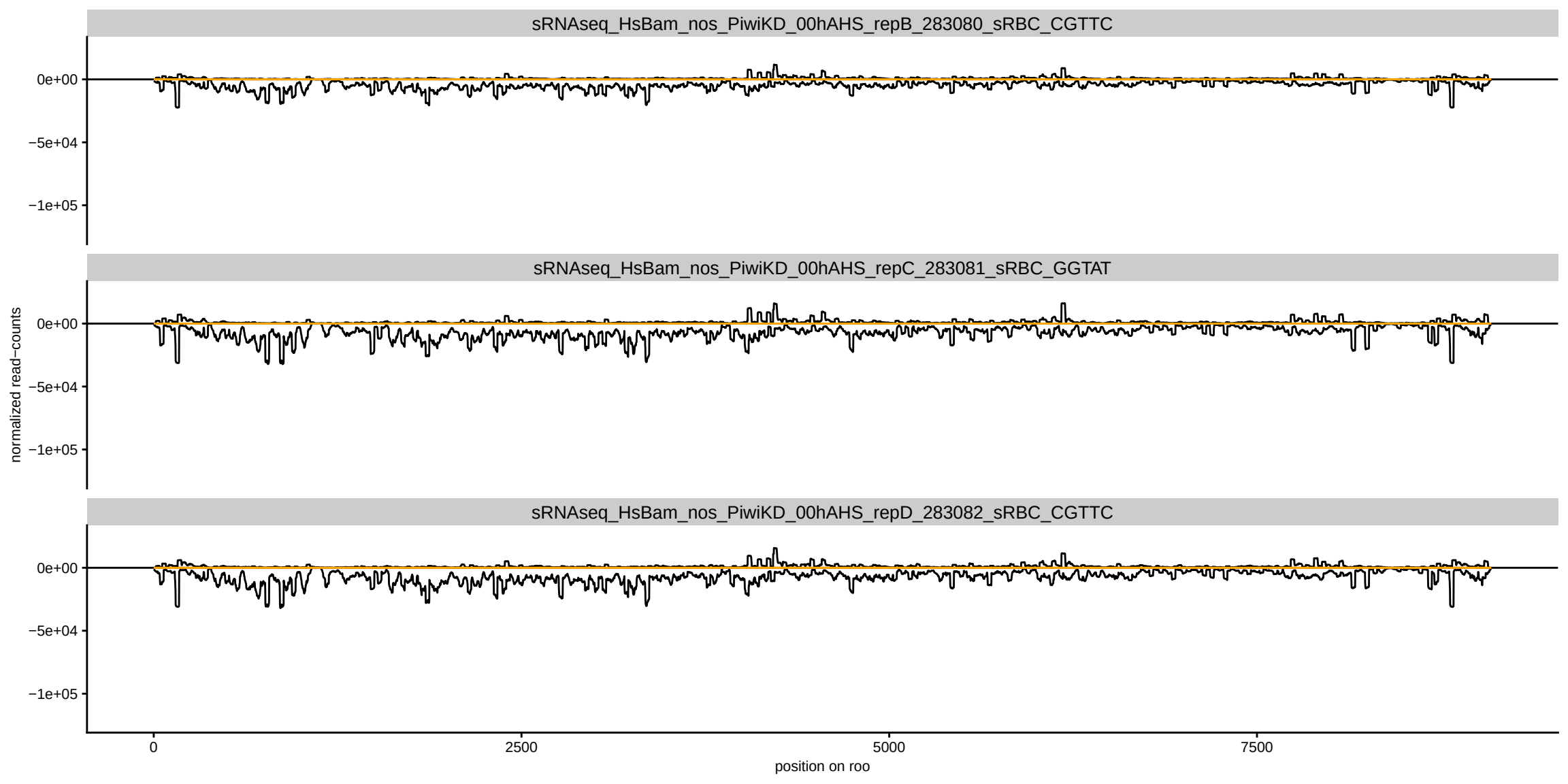
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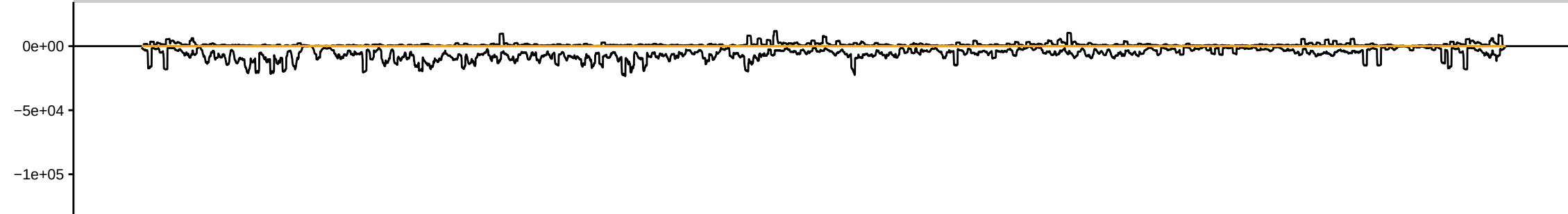
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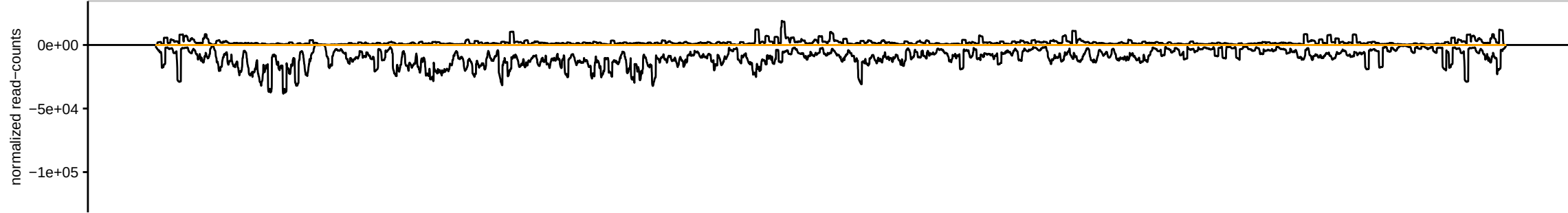
position on roo



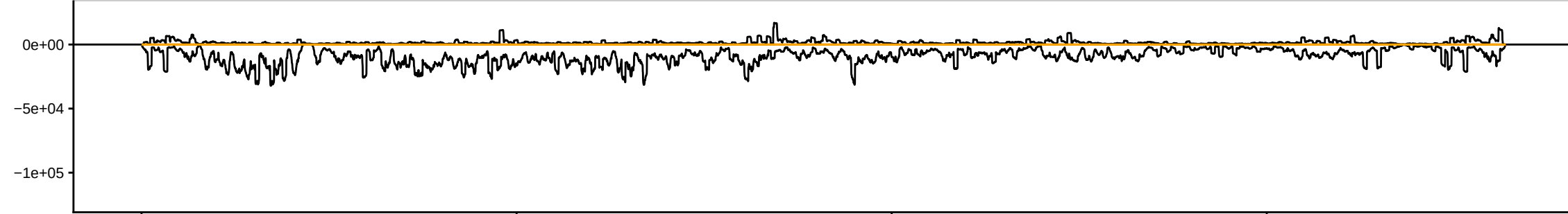
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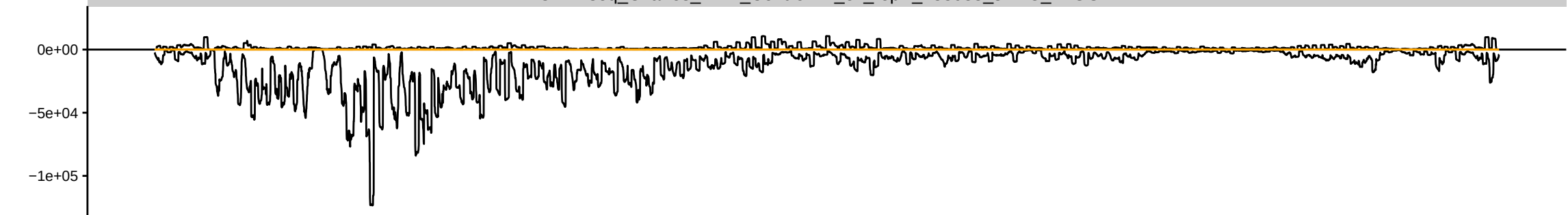
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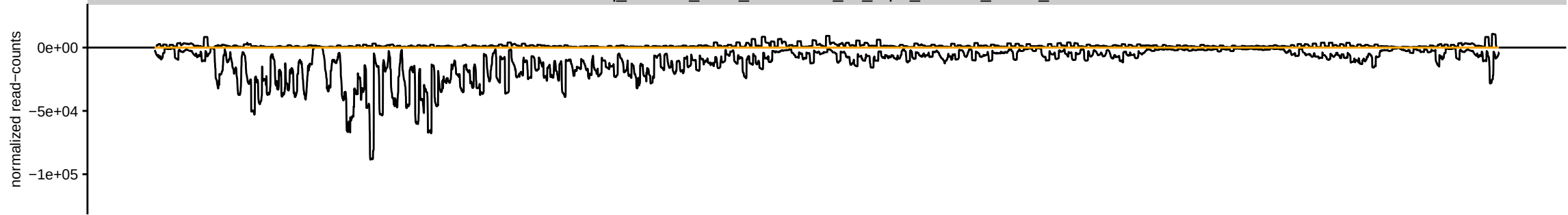
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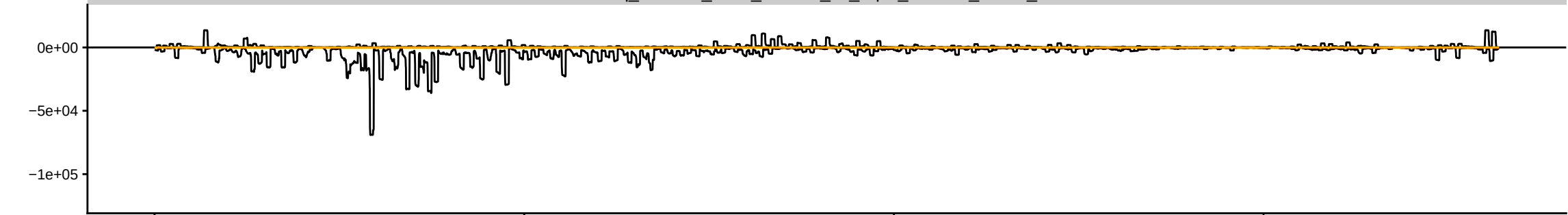
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sRNAseq_Ovaries_MTD_ControlKD_ov_repF_283083_sRBC_GAACG



sRNAseq_Ovaries_MTD_PiwiKD_ov_repE_283083_sRBC_ACGAG



position on roo

