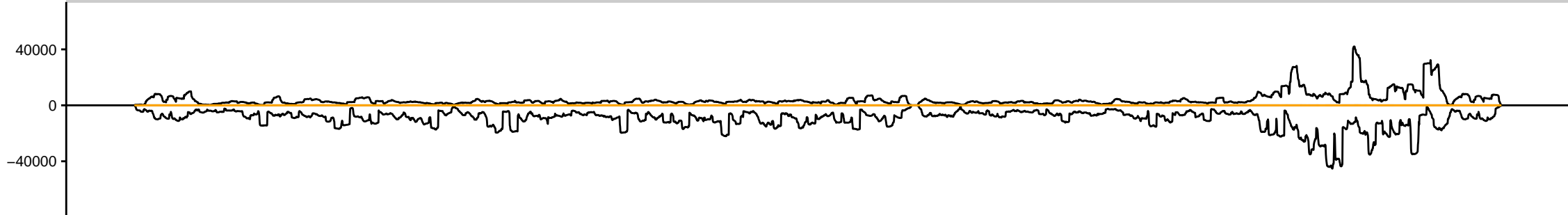
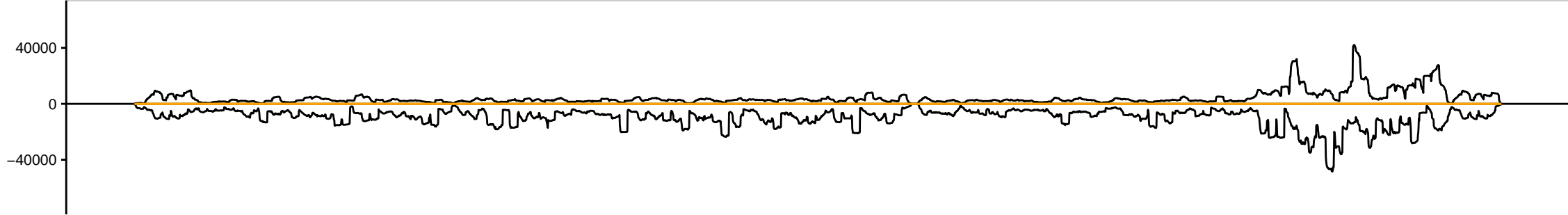


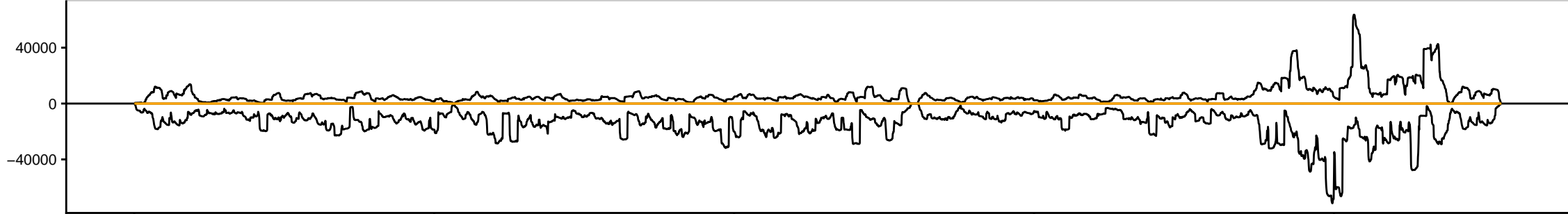
sRNAseq\_HsBam\_nos\_ControlKD\_00hAHS\_repB\_283080\_sRBC\_GTCAG



sRNAseq\_HsBam\_nos\_ControlKD\_00hAHS\_repC\_283081\_sRBC\_CAGTG

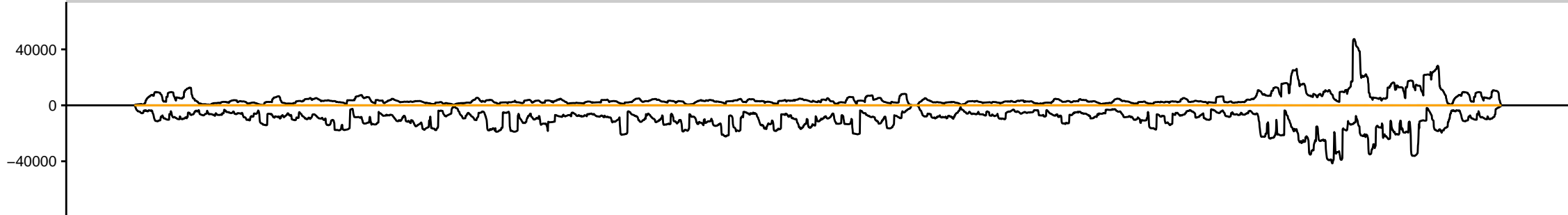


sRNAseq\_HsBam\_nos\_ControlKD\_00hAHS\_repD\_283082\_sRBC\_GTCAG

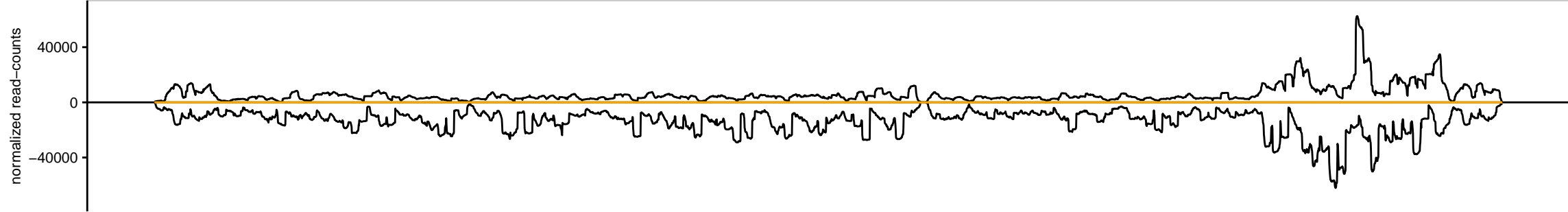


position on Dm88

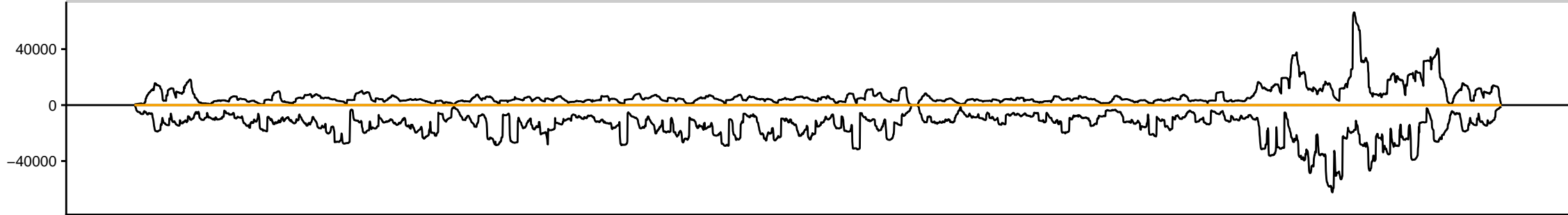
sRNAseq\_HsBam\_nos\_ControlKD\_68hAHS\_repB\_283080\_sRBC\_TGACT



sRNAseq\_HsBam\_nos\_ControlKD\_68hAHS\_repC\_283081\_sRBC\_AGCAA

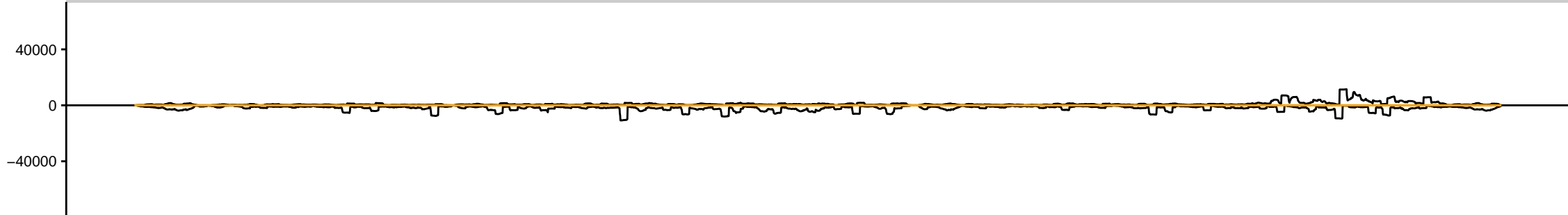


sRNAseq\_HsBam\_nos\_ControlKD\_68hAHS\_repD\_283082\_sRBC\_TGACT

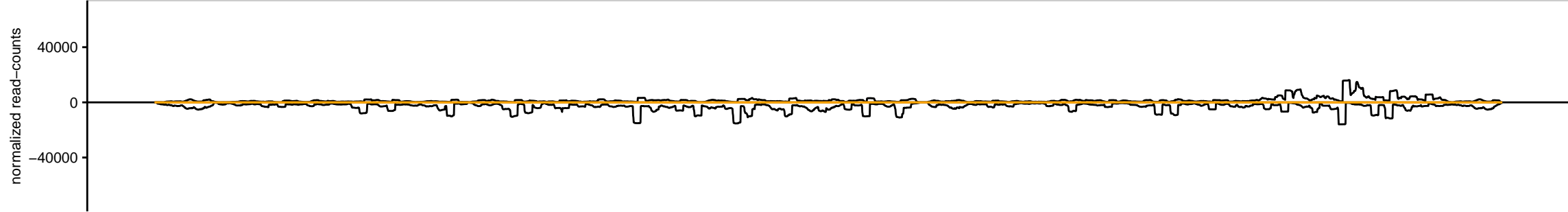


position on Dm88

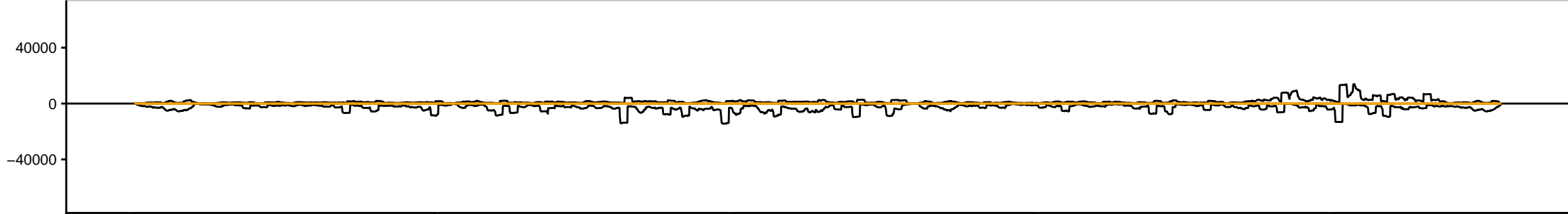
sRNAseq\_HsBam\_nos\_PiwiKD\_00hAHS\_repB\_283080\_sRBC\_CGTTTC



sRNAseq\_HsBam\_nos\_PiwiKD\_00hAHS\_repC\_283081\_sRBC\_GGTAT



sRNAseq\_HsBam\_nos\_PiwiKD\_00hAHS\_repD\_283082\_sRBC\_CGTTTC



0

1000

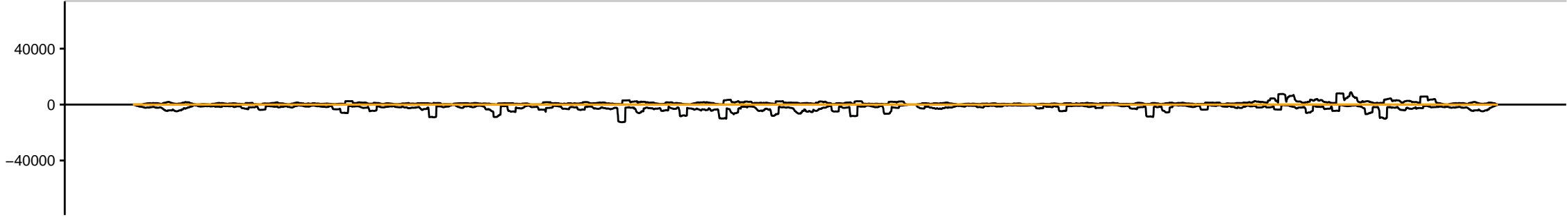
2000

3000

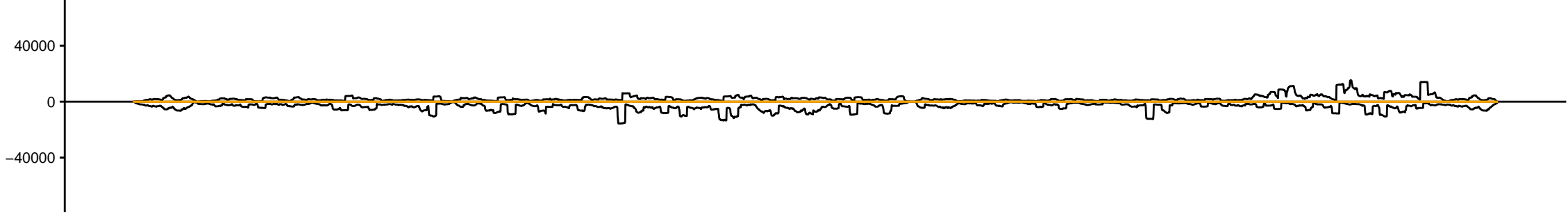
4000

position on Dm88

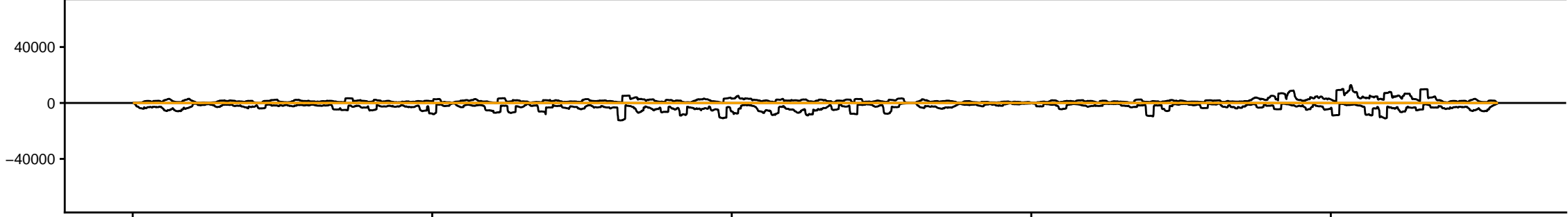
sRNAseq\_HsBam\_nos\_PiwiKD\_68hAHS\_repB\_283080\_sRBC\_ATGGA



sRNAseq\_HsBam\_nos\_PiwiKD\_68hAHS\_repC\_283081\_sRBC\_TACCA



sRNAseq\_HsBam\_nos\_PiwiKD\_68hAHS\_repD\_283082\_sRBC\_ATGGA



0

1000

2000

3000

4000

position on Dm88

sRNAseq\_Ovaries\_MTD\_ControlKD\_ov\_repE\_283083\_sRBC\_ATGGA

40000

0

-40000

sRNAseq\_Ovaries\_MTD\_ControlKD\_ov\_repF\_283083\_sRBC\_GAACG

40000

0

-40000

sRNAseq\_Ovaries\_MTD\_PiwiKD\_ov\_repE\_283083\_sRBC\_ACGAG

40000

0

-40000

0

1000

2000

3000

4000

position on Dm88

normalized read-counts

normalized read-counts

40000  
0  
-40000

0

1000

2000

3000

4000

position on Dm88

