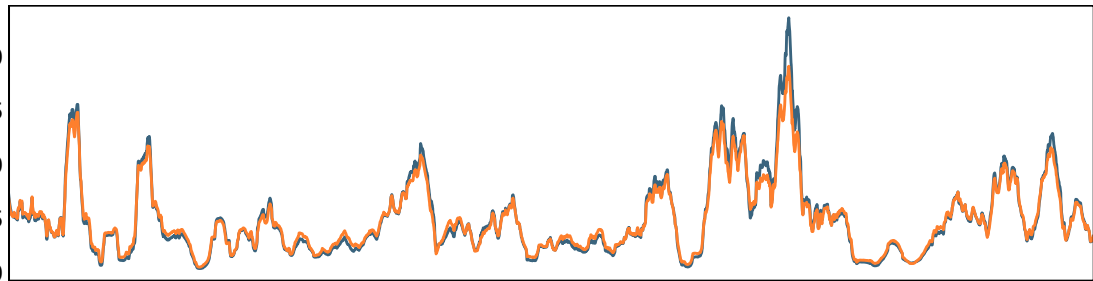


Avg. H3K27me3 CPM

20
15
10
5
0



cGAS_KO

Chromatin
State



2e+07

4e+07

6e+07

Chr 19 coordinate