

# K562TPA1hr PositiveSingle E.644 scCLTdegenNuc631

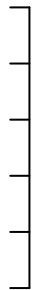
Average coverage per gene

0.0035  
0.0020

-5kb

TSS

5kb



# K562TPA1hr PositiveSingle E.644 scCLTdegenNuc633

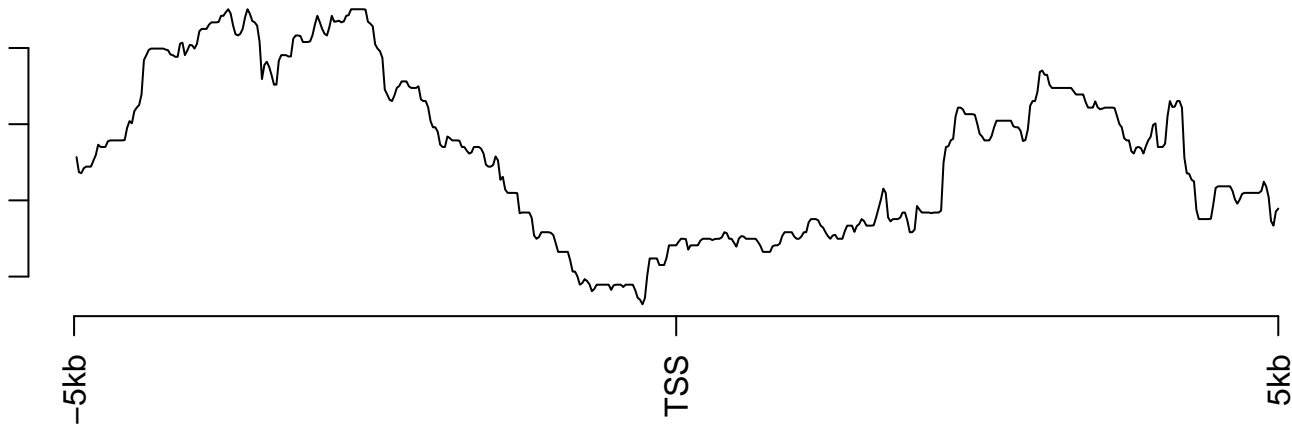
Average coverage per gene

8e-04  
4e-04

-5kb

TSS

5kb



# K562TPA1hr PositiveSingle E.651 scCLTdegenNuc651

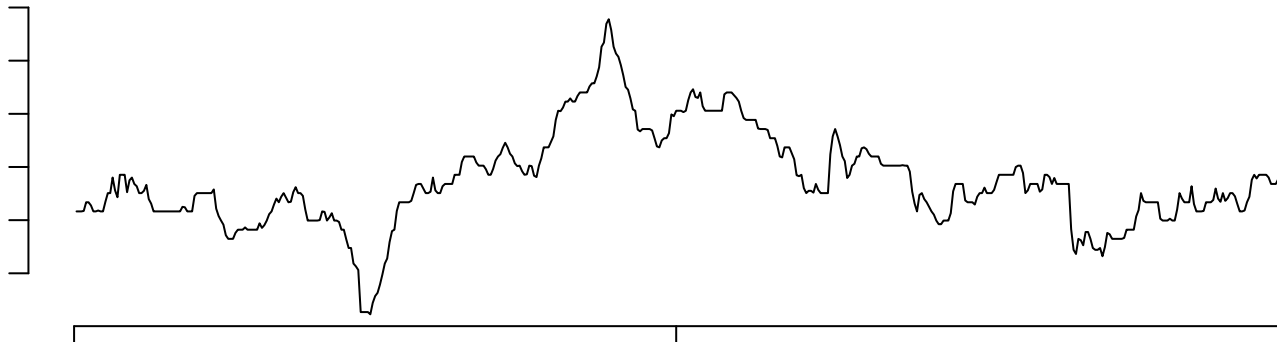
Average coverage per gene

4e-04  
7e-04

-5kb

TSS

5kb



# K562TPA1hr PositiveSingle E.651 scCLTdegenNuc652

Average coverage per gene

2e-04 5e-04 8e-04

-5kb

TSS

5kb



# K562TPA1hr PositiveSingle E.652 scCLTdegenNuc653

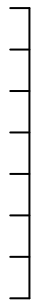
Average coverage per gene

0.0006 0.0014

-5kb

TSS

5kb



# K562TPA1hr PositiveMerged E.– K562TPA1hrPositiveMerged

Average coverage per gene

0.0065  
0.0050

-5kb

TSS

5kb

