

# K562TPA2hr PositiveSingle E.644 scCLTdegenNuc644

Average coverage per gene

0.0010 0.0020

-5kb

TSS

5kb



# K562TPA2hr PositiveSingle E.644 scCLTdegenNuc645

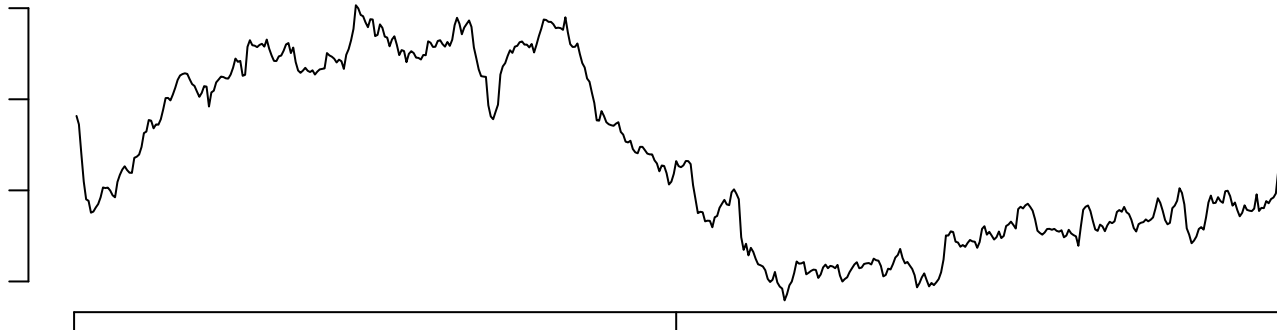
Average coverage per gene

0.0035  
0.0025

-5kb

TSS

5kb



# K562TPA2hr PositiveSingle E.651 scCLTdegenNuc646

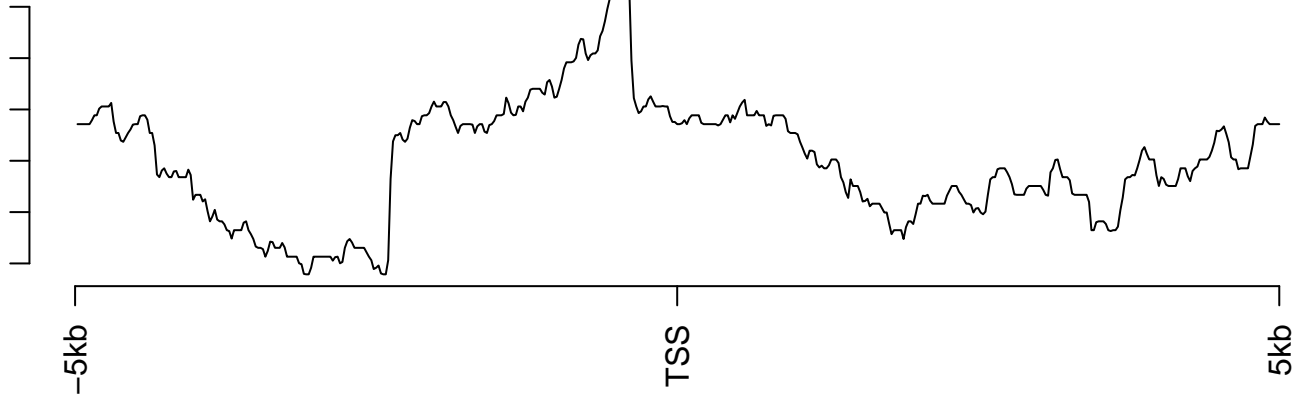
Average coverage per gene

7e-04  
4e-04

-5kb

TSS

5kb



# K562TPA2hr PositiveSingle E.651 scCLTdegenNuc647

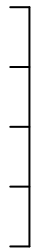
Average coverage per gene

0.011 0.013 0.015

-5kb

TSS

5kb



# K562TPA2hr PositiveSingle E.651 scCLTdegenNuc648

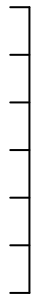
Average coverage per gene

1e-04  
4e-04  
7e-04

-5kb

TSS

5kb



# K562TPA2hr PositiveSingle E.651 scCLTdegenNuc649

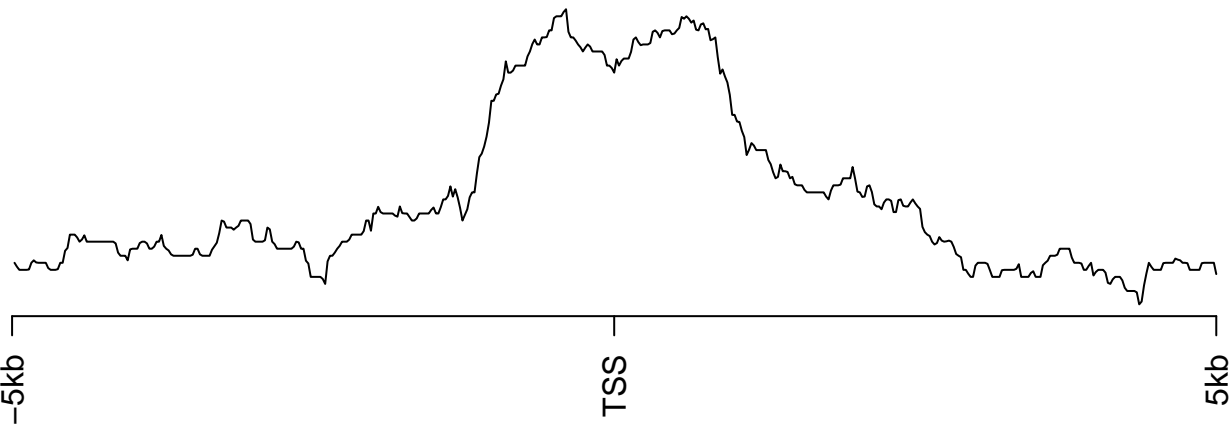
Average coverage per gene

4e-04  
8e-04

-5kb

TSS

5kb



# K562TPA2hr PositiveSingle E.651 scCLTdegenNuc650

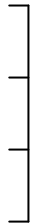
Average coverage per gene

6e-04 8e-04

-5kb

TSS

5kb



# K562TPA2hr PositiveSingle E.673 scCLTdegenNuc922\_b

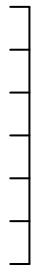
Average coverage per gene

0.00005 0.00025

-5kb

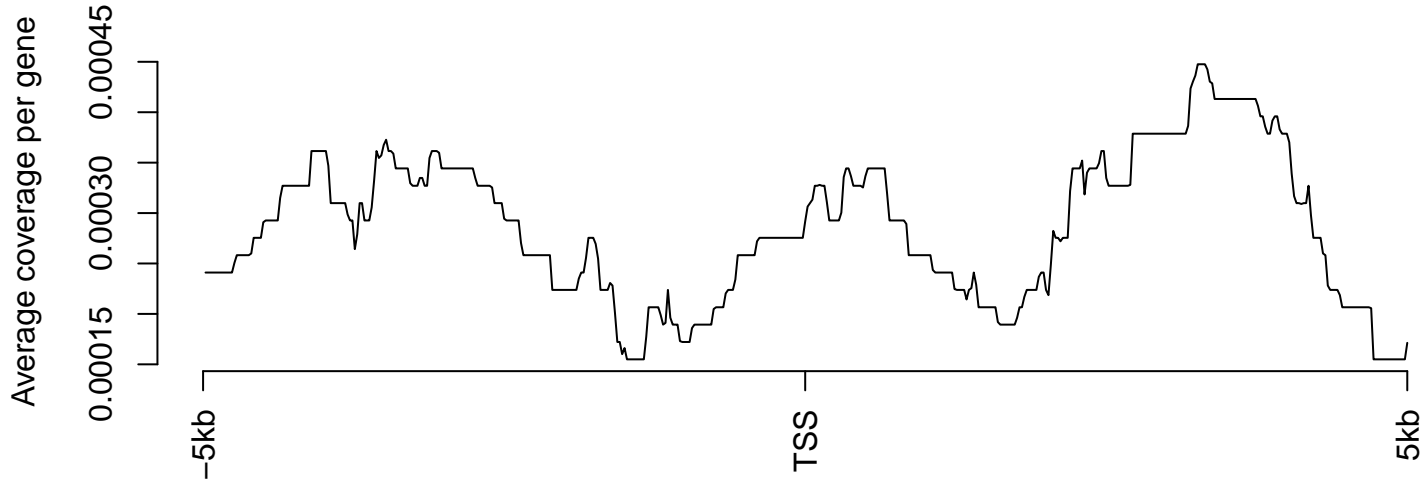
TSS

5kb





**K562TPA2hr PositiveSingle E.673 scCLTdegenNuc923\_b**



# K562TPA2hr PositiveSingle E.673 scCLTdegenNuc927\_b

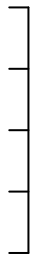
Average coverage per gene

0.0010  
0.0004

-5kb

TSS

5kb



**K562TPA2hr PositiveSingle E.673 scCLTdegenNuc928\_b**



# K562TPA2hr PositiveSingle E.673 scCLTdegenNuc930\_b

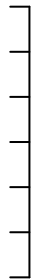
Average coverage per gene

0.00000 0.00020

-5kb

TSS

5kb



# K562TPA2hr PositiveSingle E.673 scCLTdegenNuc931\_b

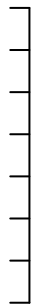
Average coverage per gene

1e-04 4e-04 7e-04

-5kb

TSS

5kb



# K562TPA2hr PositiveMerged E.– K562TPA2hrPositiveMerged

Average coverage per gene

0.021 0.023 0.025

-5kb

TSS

5kb

