

K562TPA15min PositiveSingle E.644 scCLTdegenNuc641

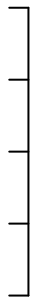
Average coverage per gene

0.0004 0.0008 0.0012

-5kb

TSS

5kb



K562TPA15min PositiveSingle E.644 scCLTdegenNuc642

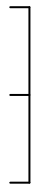
Average coverage per gene

0.0020
0.0010

-5kb

TSS

5kb



K562TPA15min PositiveSingle E.644 scCLTdegenNuc643

Average coverage per gene

3e-04
5e-04

-5kb

TSS

5kb



K562TPA15min PositiveSingle E.651 scCLTdegenNuc639

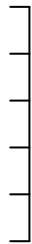
Average coverage per gene

0.0008 0.0014

-5kb

TSS

5kb



K562TPA15min PositiveSingle E.651 scCLTdegenNuc640

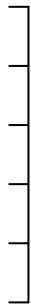
Average coverage per gene

0.0008
0.0002

-5kb

TSS

5kb



K562TPA15min PositiveMerged E.– K562TPA15minPositiveMerged

Average coverage per gene

0.0030 0.0045

-5kb

TSS

5kb

