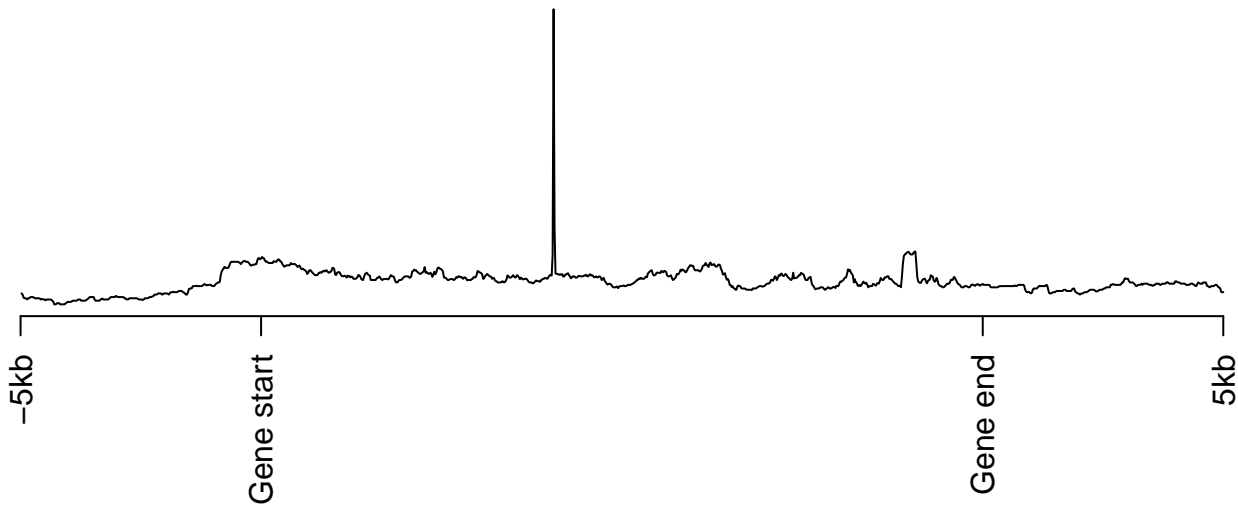


K562TPA15min PositiveSingle E.644 scCLTdegenNuc641

Average coverage per gene

0.004
0.001



K562TPA15min PositiveSingle E.644 scCLTdegenNuc642

Average coverage per gene

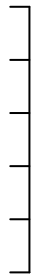
0.0020
0.0005

-5kb

Gene start

Gene end

5kb



K562TPA15min PositiveSingle E.644 scCLTdegenNuc643

Average coverage per gene

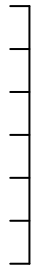
2e-04 5e-04 8e-04

-5kb

Gene start

Gene end

5kb



K562TPA15min PositiveSingle E.651 scCLTdegenNuc639

Average coverage per gene

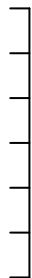
0.0018
0.0012
0.0006

-5kb

Gene start

Gene end

5kb



K562TPA15min PositiveSingle E.651 scCLTdegenNuc640

Average coverage per gene

0.0008
0.0002

-5kb

Gene start

Gene end

5kb



K562TPA15min PositiveMerged E.– K562TPA15minPositiveMerged

Average coverage per gene

0.009
0.006
0.003

-5kb

Gene start

Gene end

5kb

