

K562TPA1hr PositiveSingle E.644 scCLTdegenNuc631

Average coverage per gene

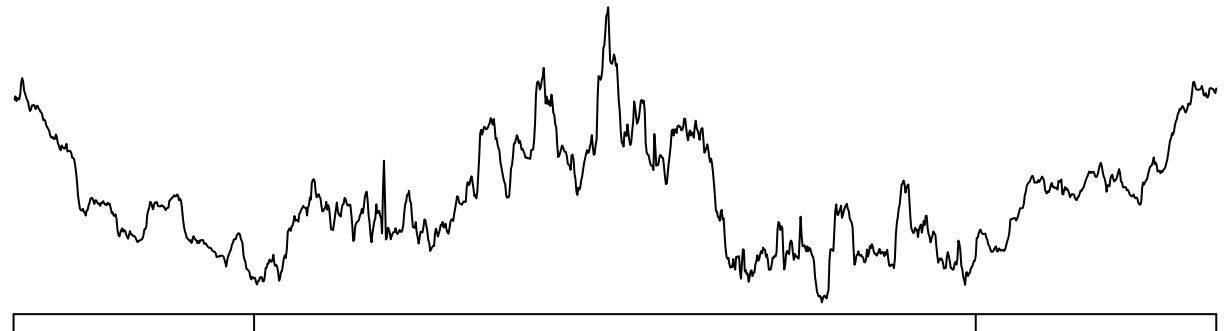
0.004
0.002

-5kb

Gene start

Gene end

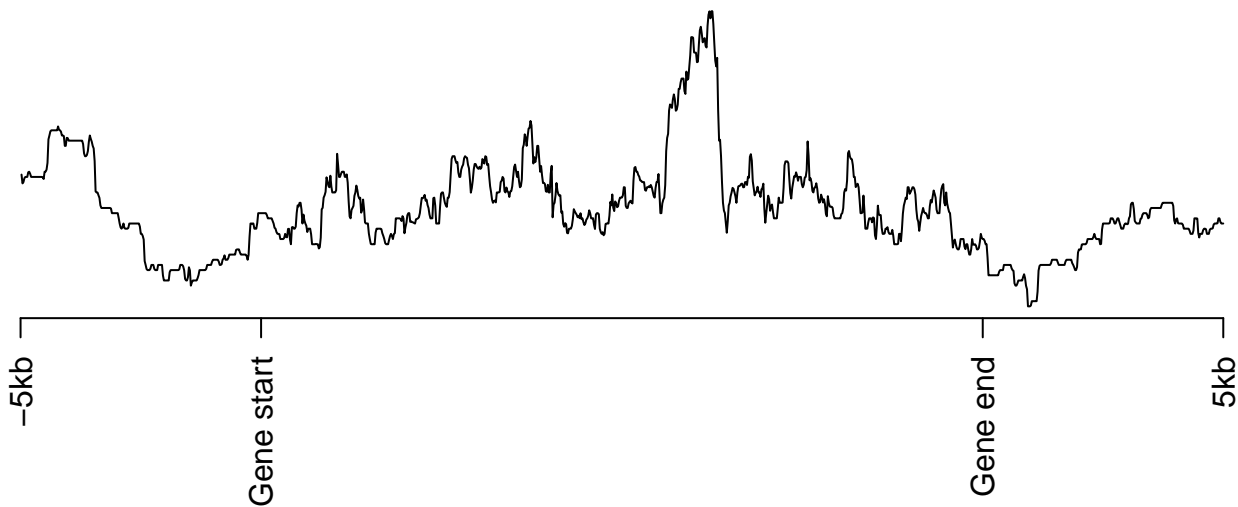
5kb



K562TPA1hr PositiveSingle E.644 scCLTdegenNuc633

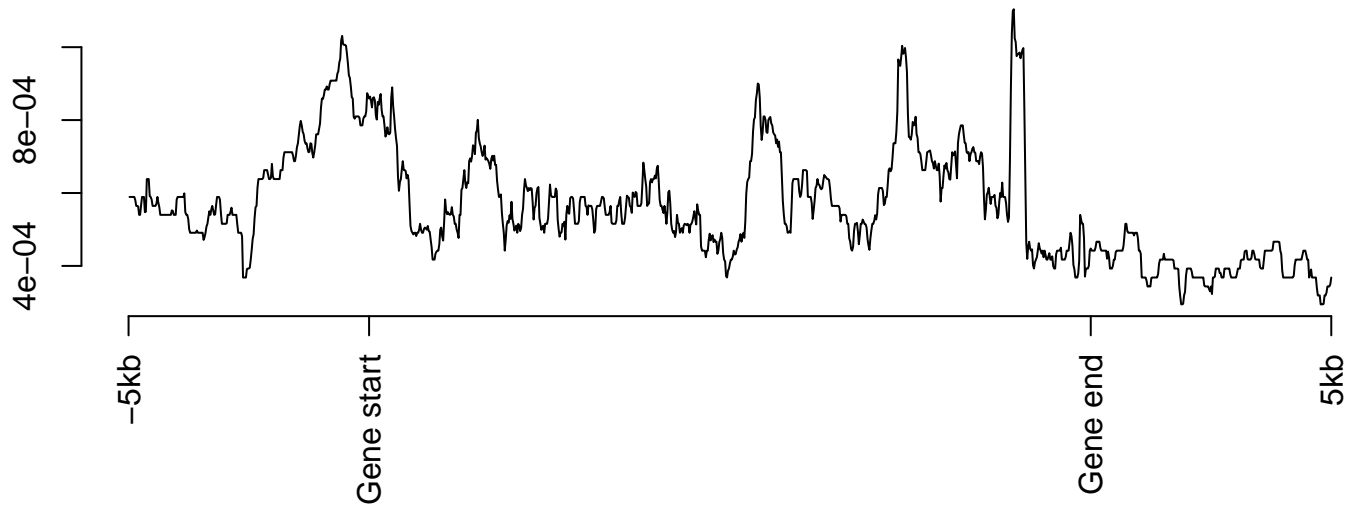
Average coverage per gene

0.0002 0.0008 0.0014



K562TPA1hr PositiveSingle E.651 scCLTdegenNuc651

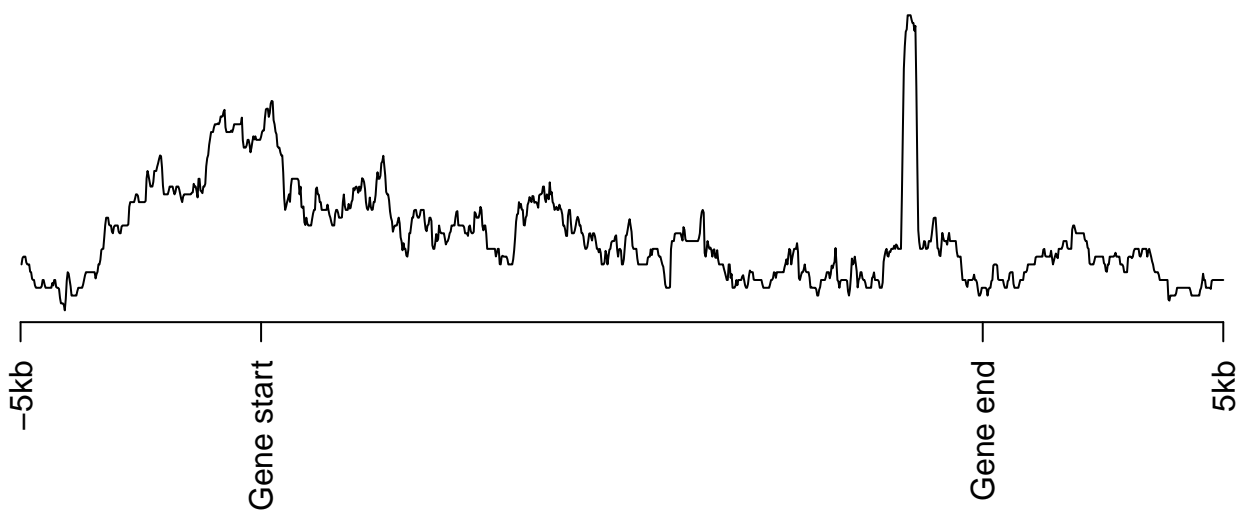
Average coverage per gene



K562TPA1hr PositiveSingle E.651 scCLTdegenNuc652

Average coverage per gene

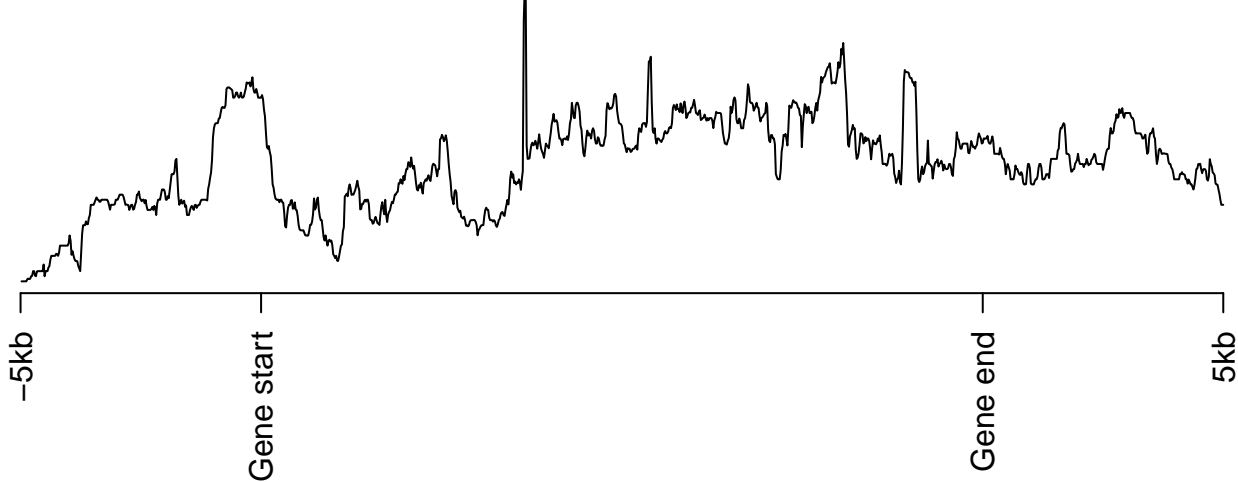
0.0004 0.0008 0.0012



K562TPA1hr PositiveSingle E.652 scCLTdegenNuc653

Average coverage per gene

0.0006 0.0012 0.0018



K562TPA1hr PositiveMerged E.– K562TPA1hrPositiveMerged

Average coverage per gene

0.008
0.006
0.004

-5kb

Gene start

Gene end

5kb

