

K562TPA2hr PositiveSingle E.644 scCLTdegenNuc644

Average coverage per gene

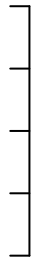
0.0010 0.0020 0.0030

-5kb

Gene start

Gene end

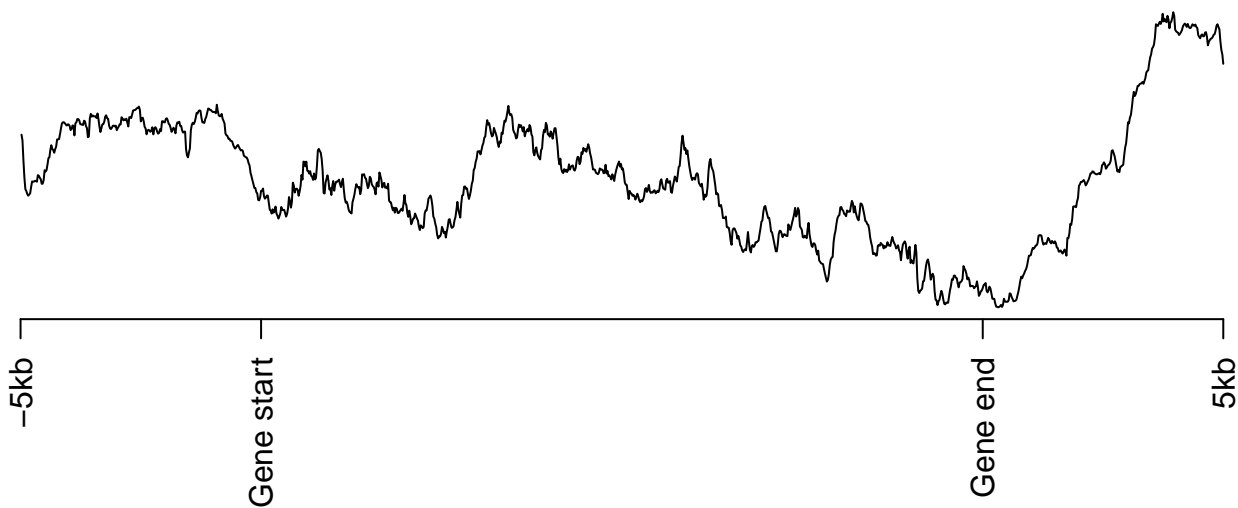
5kb



K562TPA2hr PositiveSingle E.644 scCLTdegenNuc645

Average coverage per gene

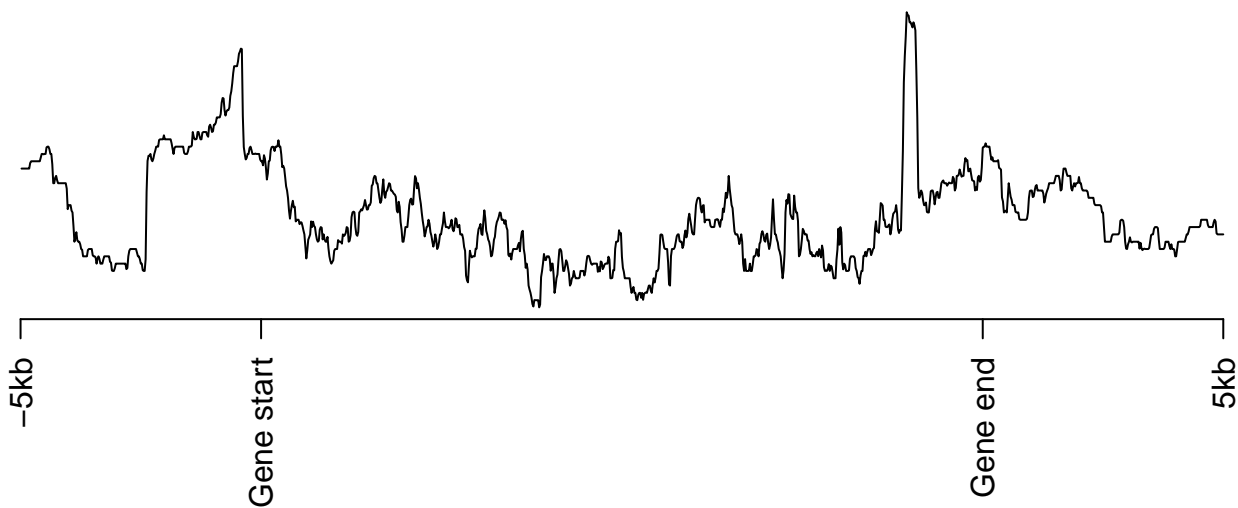
0.0015 0.0030 0.0045



K562TPA2hr PositiveSingle E.651 scCLTdegenNuc646

Average coverage per gene

0.0008
0.0002



K562TPA2hr PositiveSingle E.651 scCLTdegenNuc647

Average coverage per gene

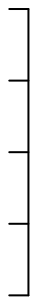
0.016
0.012
0.008

-5kb

Gene start

Gene end

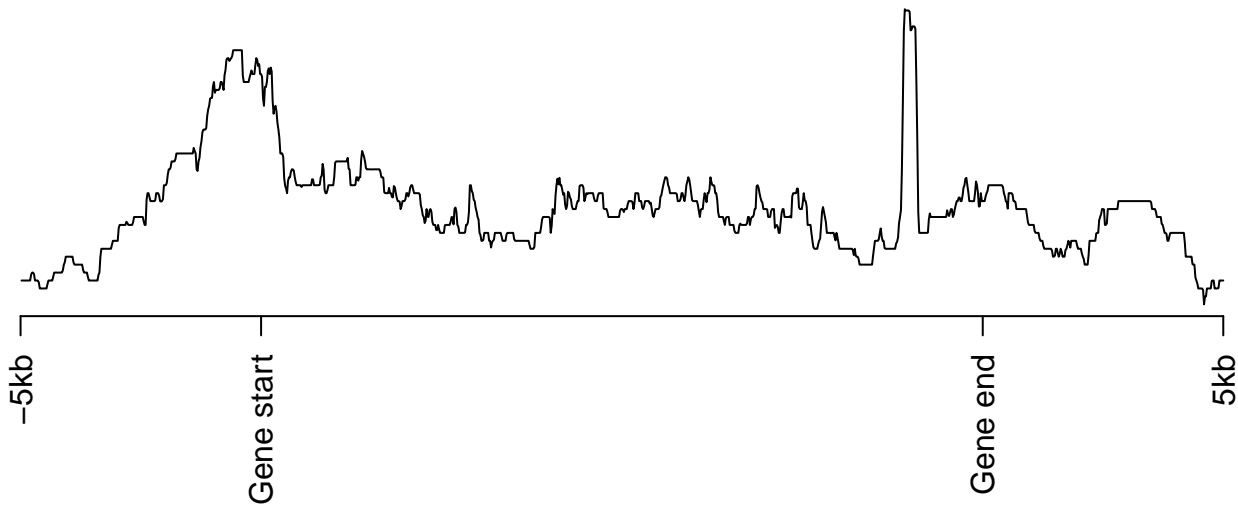
5kb



K562TPA2hr PositiveSingle E.651 scCLTdegenNuc648

Average coverage per gene

2e-04 6e-04



K562TPA2hr PositiveSingle E.651 scCLTdegenNuc649

Average coverage per gene

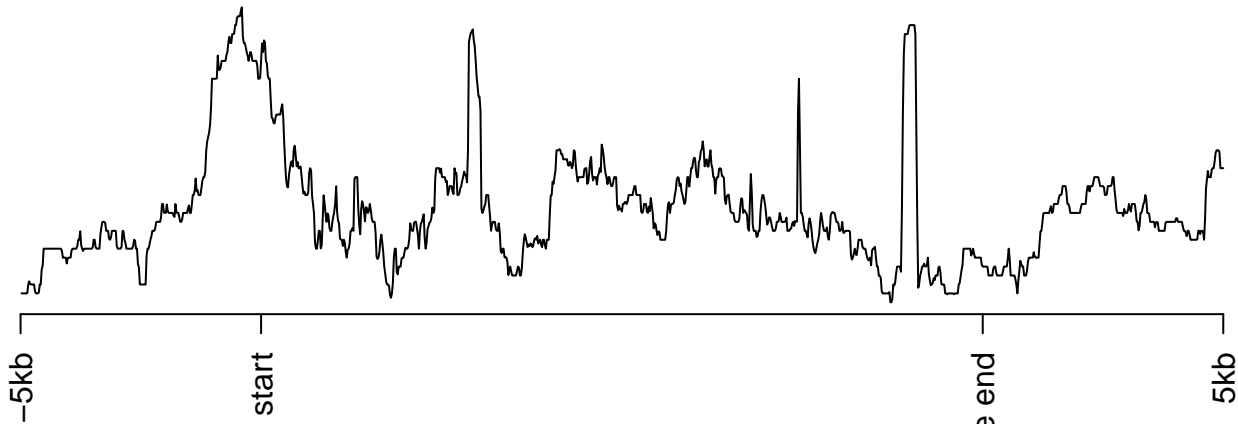
4e-04
8e-04

-5kb

Gene start

Gene end

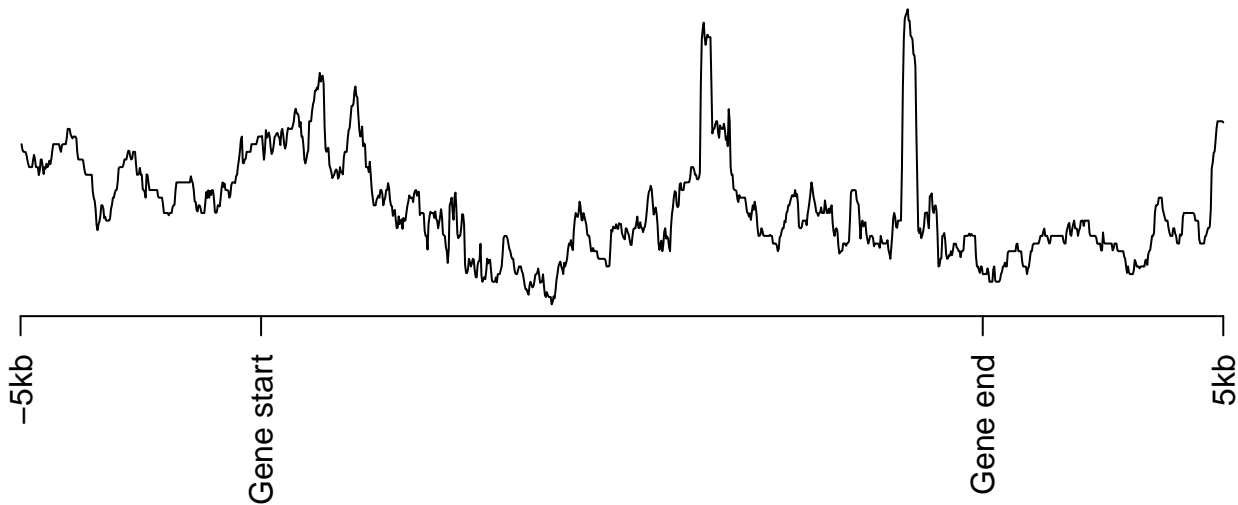
5kb



K562TPA2hr PositiveSingle E.651 scCLTdegenNuc650

Average coverage per gene

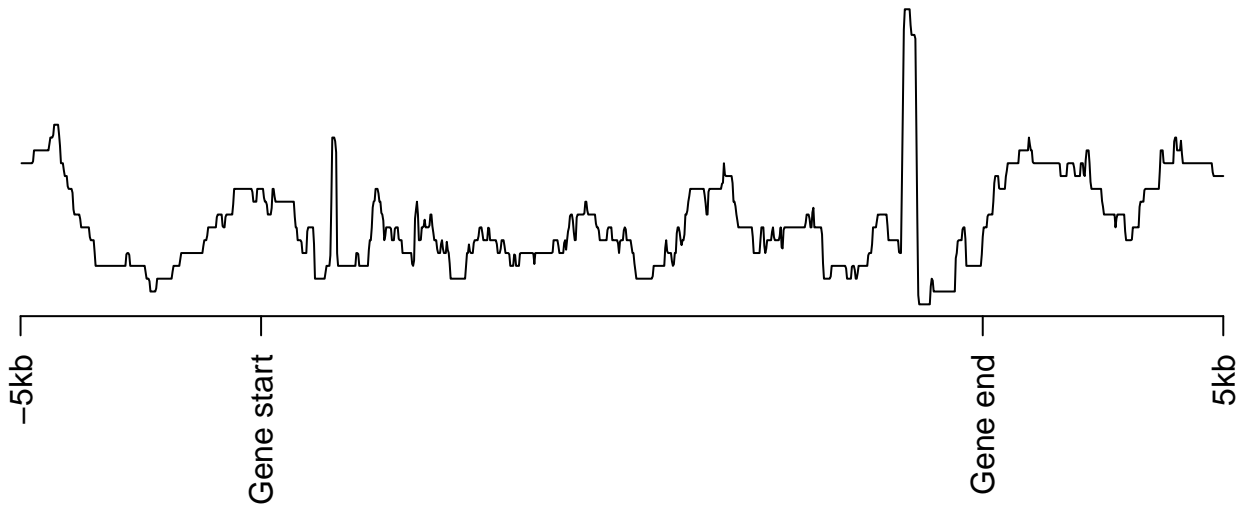
0.0004 0.0008 0.0012



K562TPA2hr PositiveSingle E.673 scCLTdegenNuc922_b

Average coverage per gene

0e+00 3e-04



K562TPA2hr PositiveSingle E.673 scCLTdegenNuc923_b

Average coverage per gene

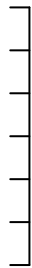
1e-04
4e-04
7e-04

-5kb

Gene start

Gene end

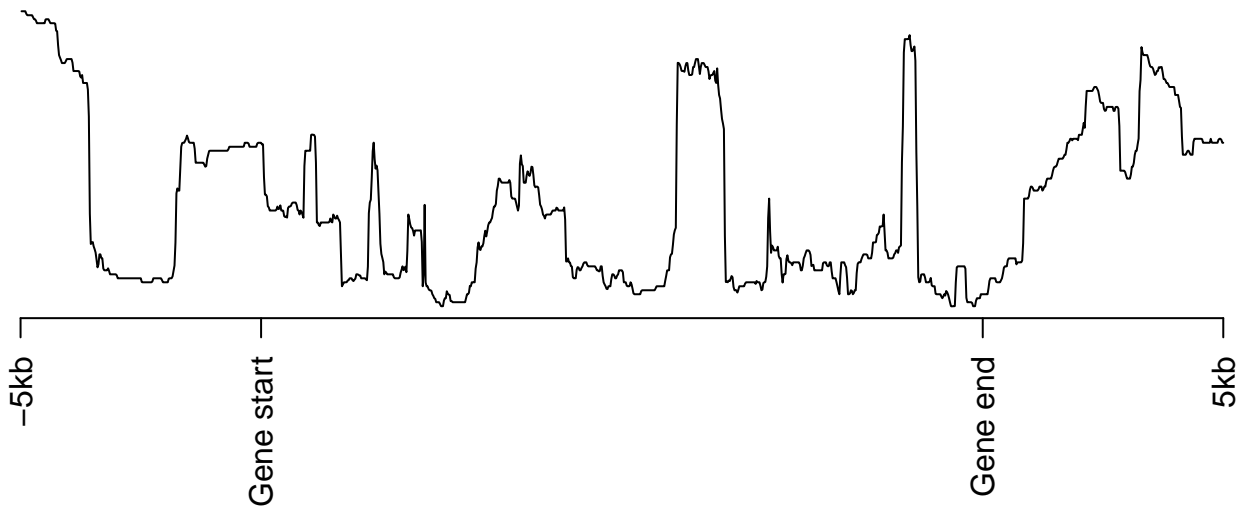
5kb



K562TPA2hr PositiveSingle E.673 scCLTdegenNuc927_b

Average coverage per gene

0.0015
0.0005



K562TPA2hr PositiveSingle E.673 scCLTdegenNuc928_b

Average coverage per gene

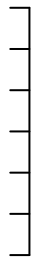
0.0012
0.0004

-5kb

Gene start

Gene end

5kb



K562TPA2hr PositiveSingle E.673 scCLTdegenNuc930_b

Average coverage per gene

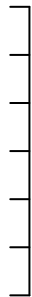
0e+00 3e-04 6e-04

-5kb

Gene start

Gene end

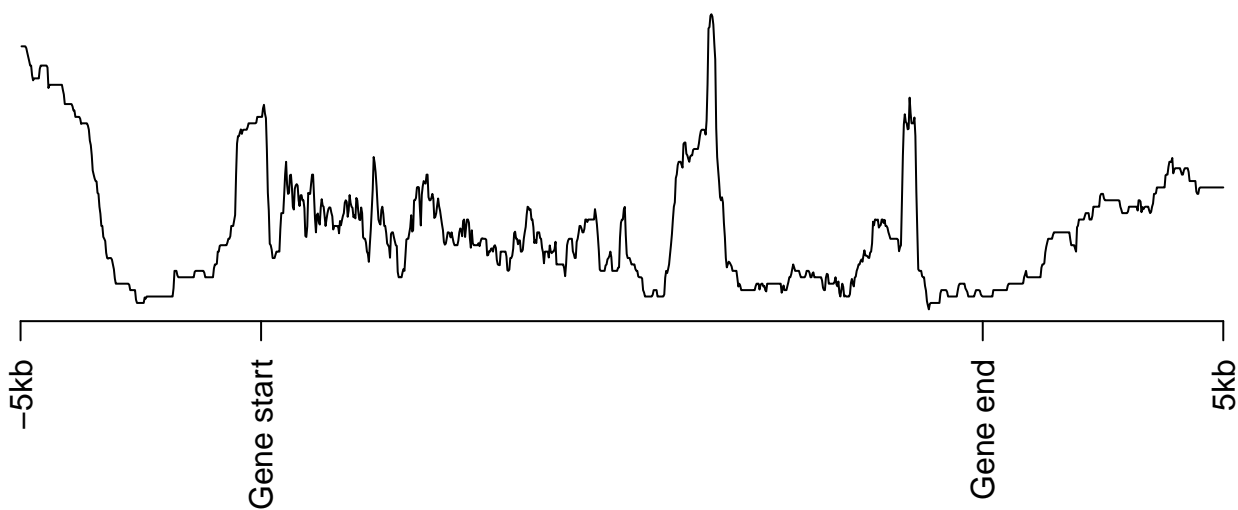
5kb



K562TPA2hr PositiveSingle E.673 scCLTdegenNuc931_b

Average coverage per gene

0.0008
0.0002



K562TPA2hr PositiveMerged E.– K562TPA2hrPositiveMerged

Average coverage per gene

0.026
0.020
0.014

-5kb

Gene start

Gene end

5kb

