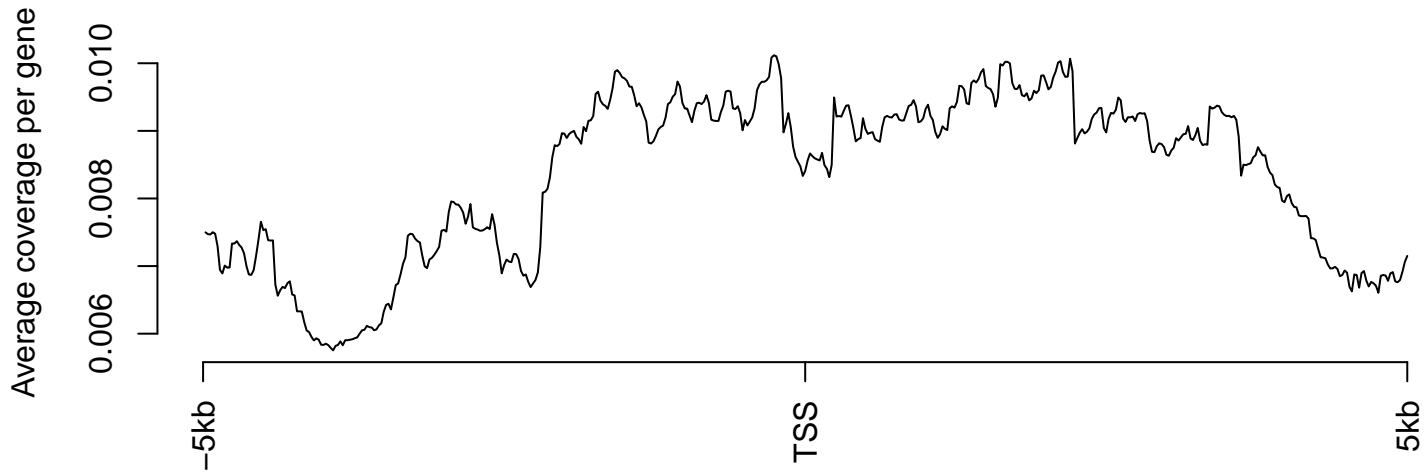


MouseNeuronSlice NegCtrlPooled E.597 scCLTdegenNuc470



MouseNeuronSlice PositiveSingle E.589 scCLTdegenNuc458

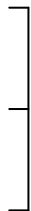
Average coverage per gene

0.030
0.020

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.589 scCLTdegenNuc459

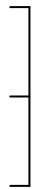
Average coverage per gene

0.035
0.025

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.589 scCLTdegenNuc460

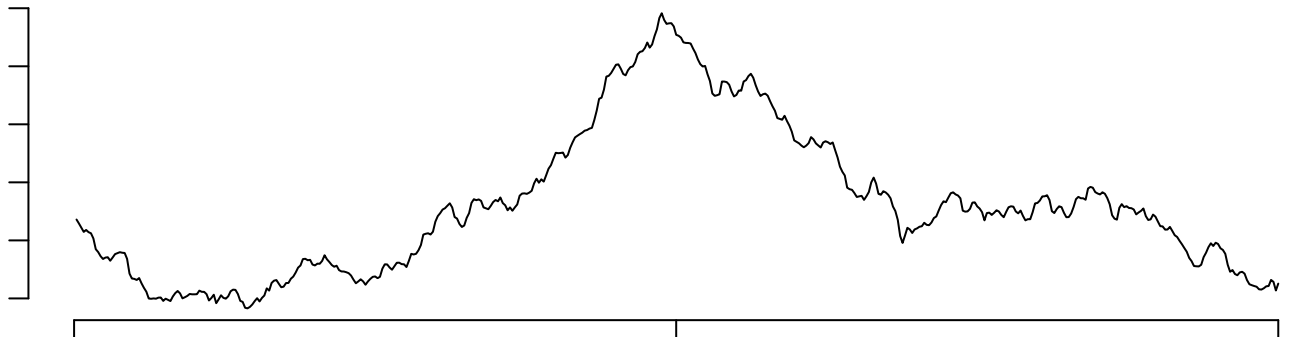
Average coverage per gene

0.006 0.008 0.010

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.591 scCLTdegenNuc465

Average coverage per gene

0.022 0.026 0.030

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.591 scCLTdegenNuc466

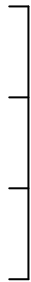
Average coverage per gene

0.045
0.035

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.591 scCLTdegenNuc467

Average coverage per gene

0.034 0.040 0.046

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.591 scCLTdegenNuc468

Average coverage per gene

0.070
0.060

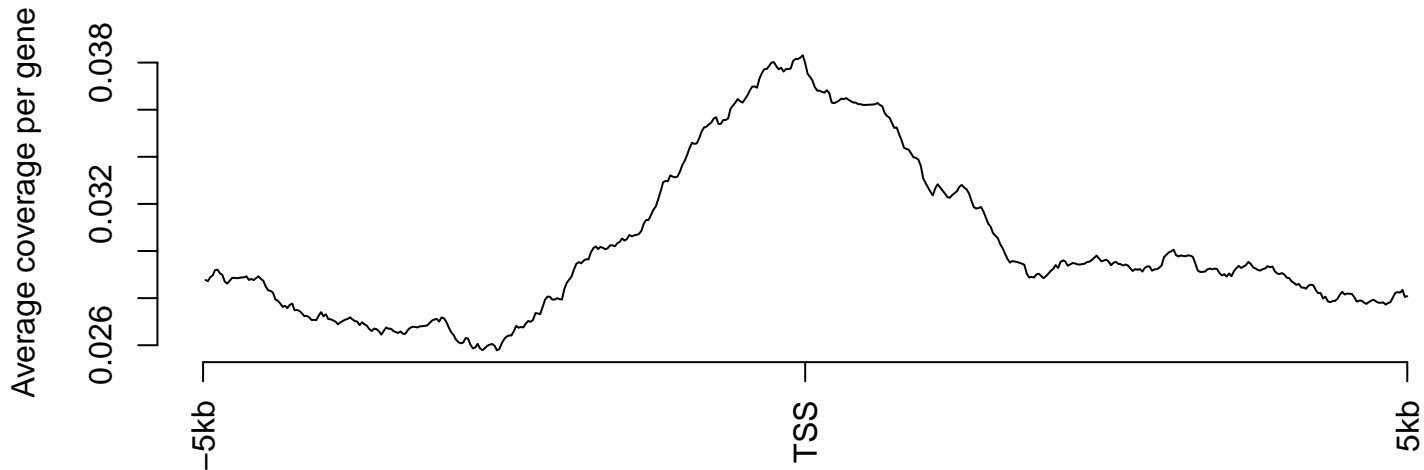
-5kb

TSS

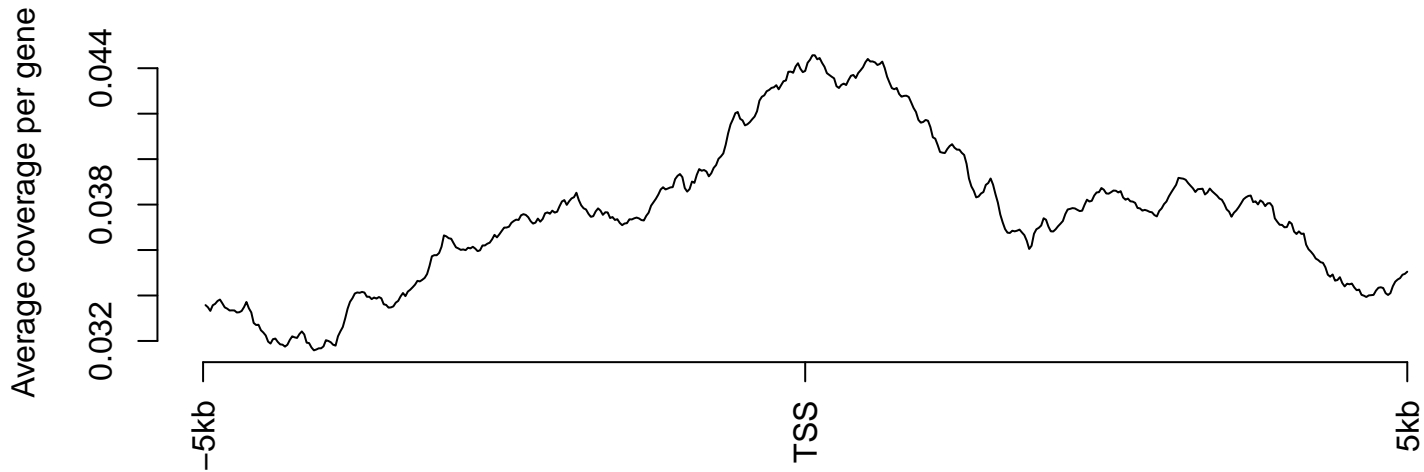
5kb



MouseNeuronSlice PositiveSingle E.597 scCLTdegenNuc469



MouseNeuronSlice PositiveSingle E.619 scCLTdegenNuc495



MouseNeuronSlice PositiveSingle E.622 scCLTdegenNuc515

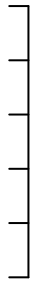
Average coverage per gene

0.006 0.008 0.010

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.622 scCLTdegenNuc516

Average coverage per gene

0.0015 0.0025 0.0035

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.622 scCLTdegenNuc517

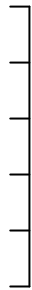
Average coverage per gene

0.006 0.008 0.010

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.622 scCLTdegenNuc518

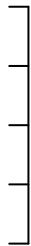
Average coverage per gene

0.0045
0.0030

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc609b

Average coverage per gene

0.0035 0.0050

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc610b

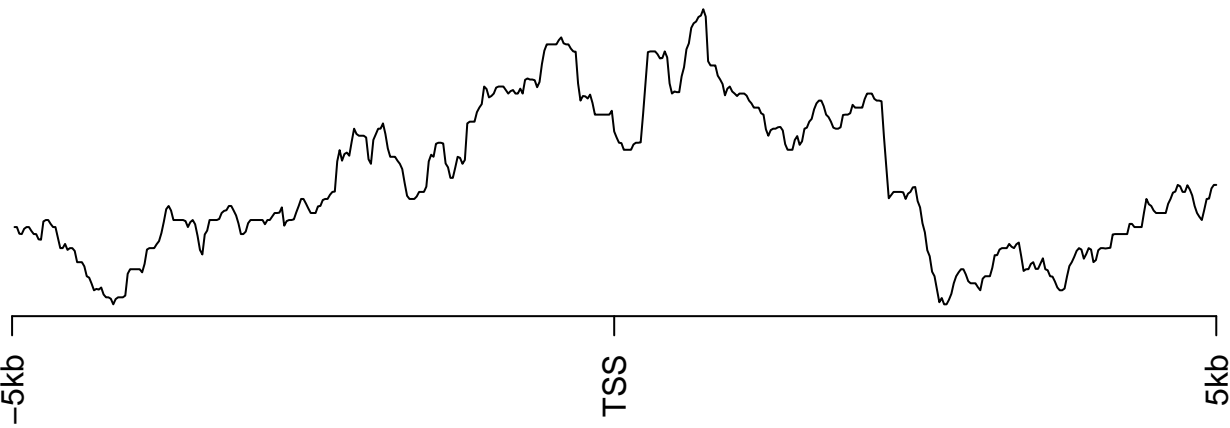
Average coverage per gene

0.0014
0.0008

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc612b

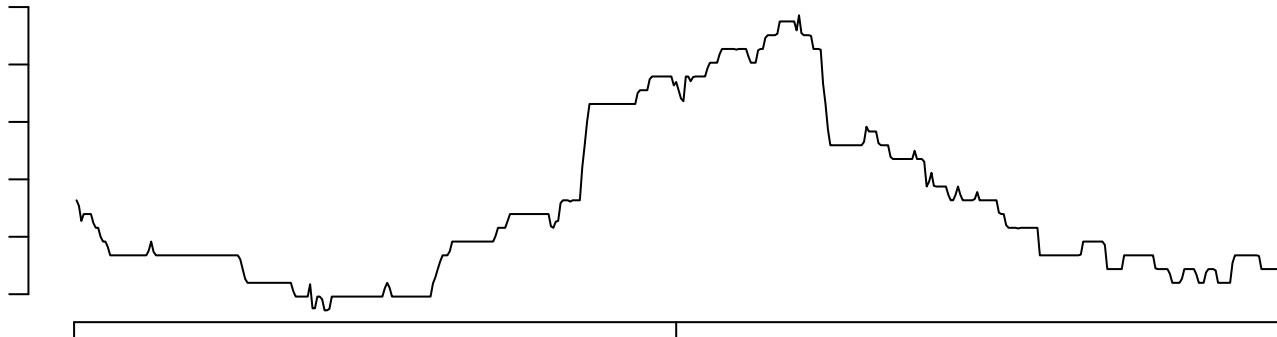
Average coverage per gene

1e-04
4e-04

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc613b

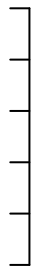
Average coverage per gene

1e-04
4e-04

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc614b

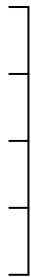
Average coverage per gene

0.0004 0.0008 0.0012

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc615b

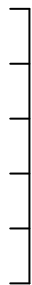
Average coverage per gene

0.013 0.015 0.017

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc616b

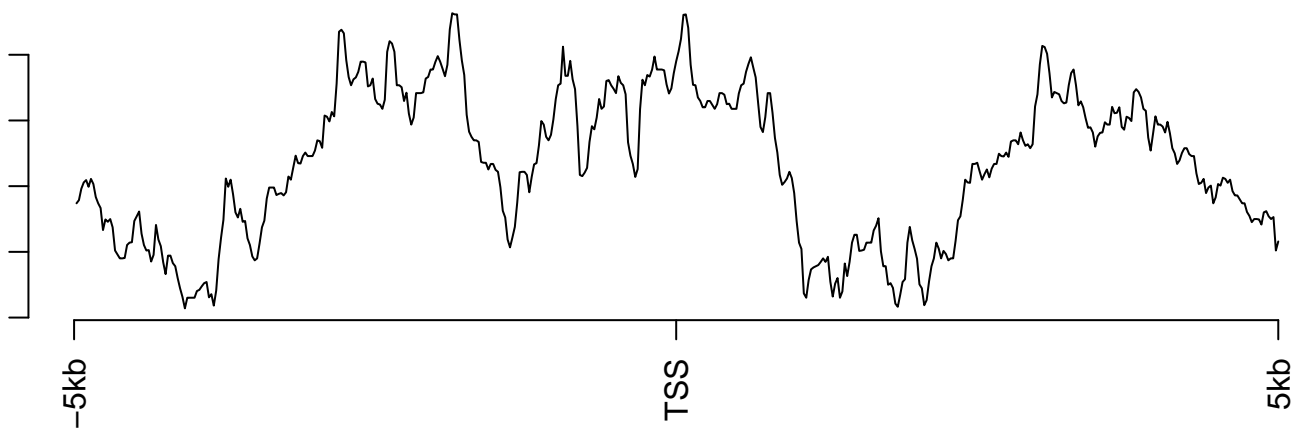
Average coverage per gene

0.0028
0.0024
0.0020

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc617b

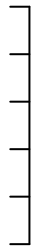
Average coverage per gene

0.0040 0.0055

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc618b

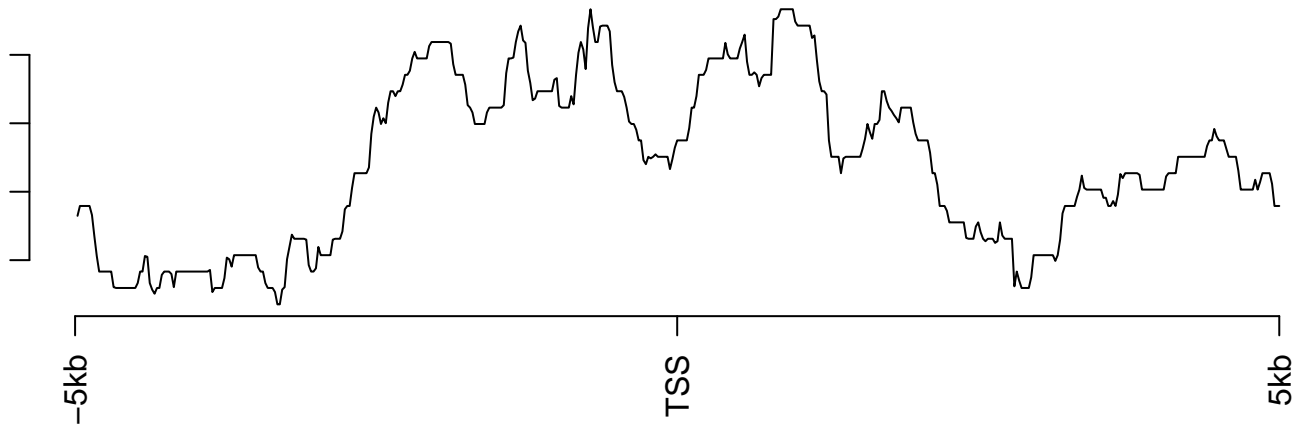
Average coverage per gene

4e-04 6e-04

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc619b

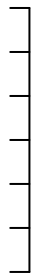
Average coverage per gene

0.00040
0.00020

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.643 scCLTdegenNuc620

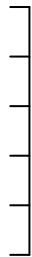
Average coverage per gene

0.0009
0.0006

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.670 scCLTdegenNuc840

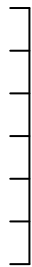
Average coverage per gene

0.0035
0.0020
0.0005

-5kb

TSS

5kb



MouseNeuronSlice PositiveSingle E.670 scCLTdegenNuc843

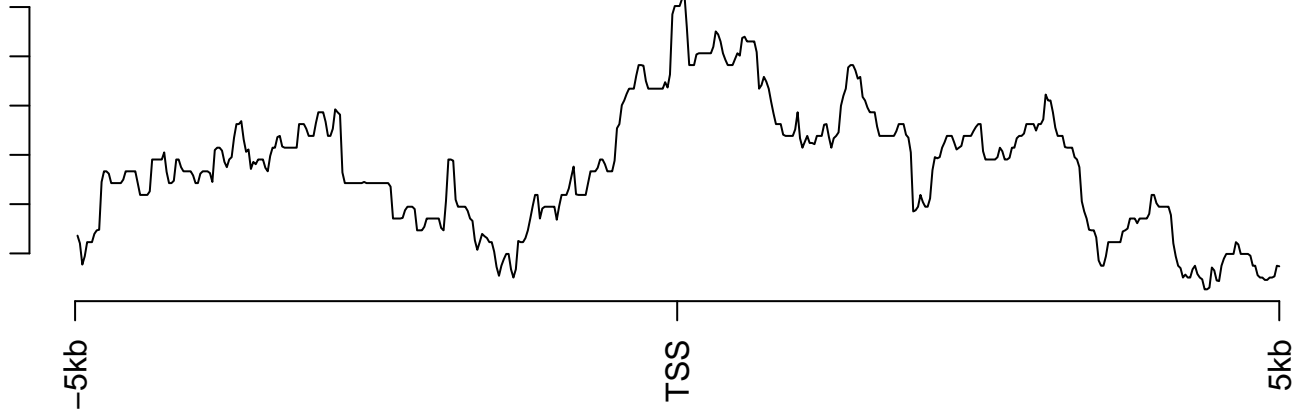
Average coverage per gene

0.0009
0.0006

-5kb

TSS

5kb



MouseNeuronSlice PositivePooled E.584 scCLTdegenNuc413

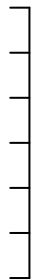
Average coverage per gene

0.08 0.10 0.12 0.14

-5kb

TSS

5kb



MouseNeuronSlice PositivePooled E.589 scCLTdegenNuc462

Average coverage per gene

0.014 0.020 0.026

-5kb

TSS

5kb



MouseNeuronSlice PositiveMerged E.– MouseNeuronSlicePositiveMerged

Average coverage per gene

0.32 0.36 0.40

-5kb

TSS

5kb



MouseNeuronSlice PositiveAll E.– MouseNeuronSlicePositiveAll

