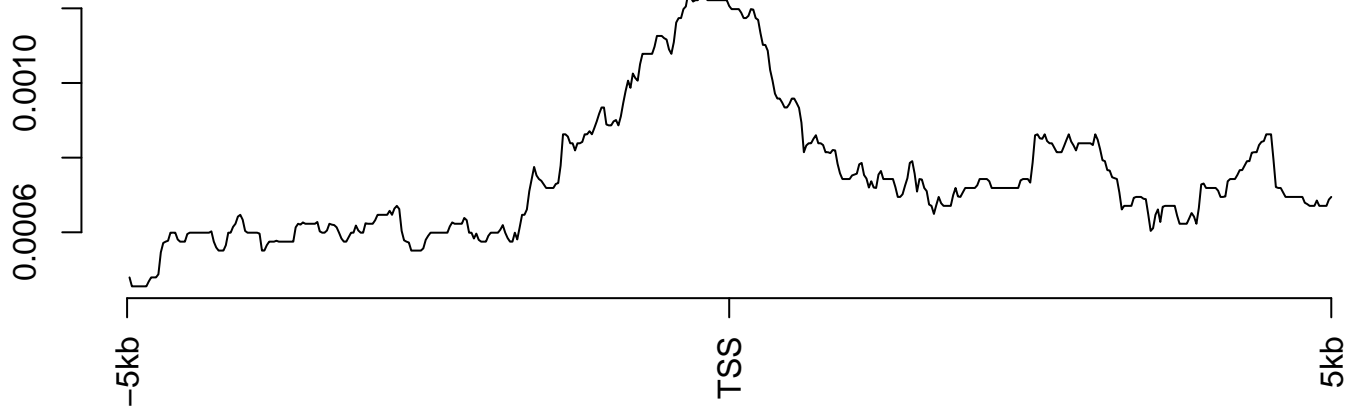


MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc603



MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc604

Average coverage per gene



MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc605

Average coverage per gene

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

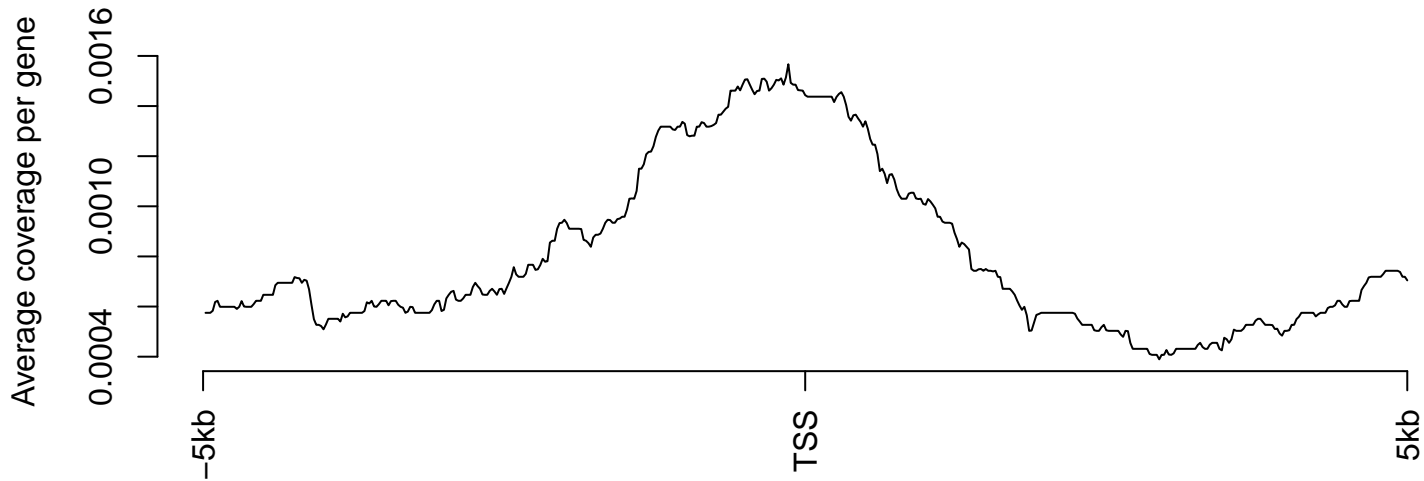
-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc606



MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc607

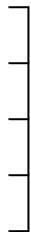
Average coverage per gene

0.0014
0.0008

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.642 scCLTdegenNuc608b

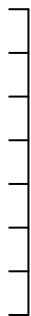
Average coverage per gene

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

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc621

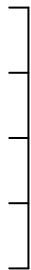
Average coverage per gene

0.0010 0.0020 0.0030

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc622

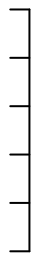
Average coverage per gene

0.007 0.010

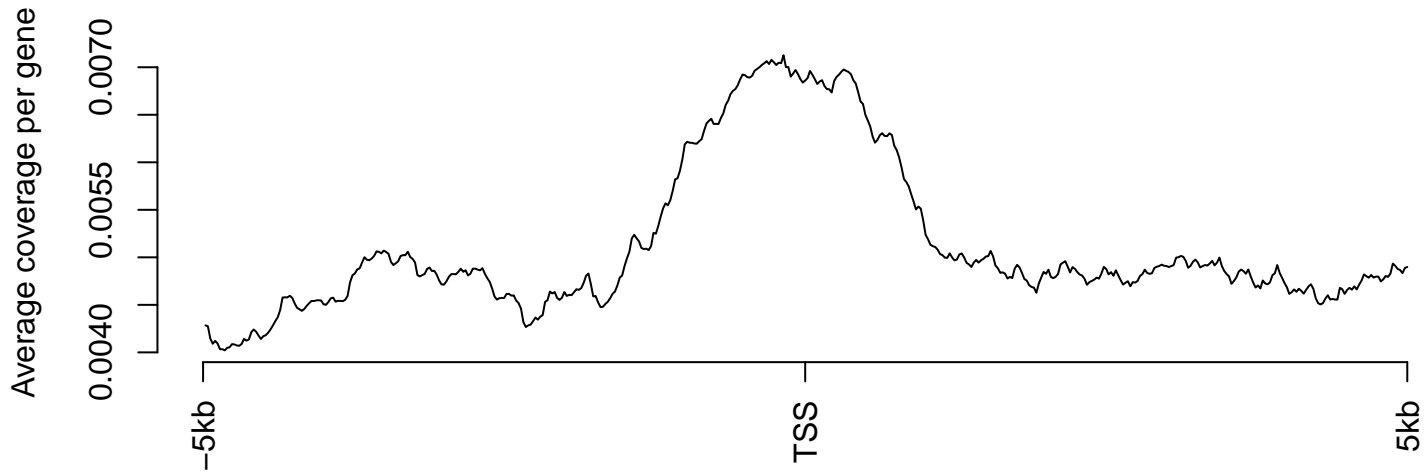
-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc623



MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc624

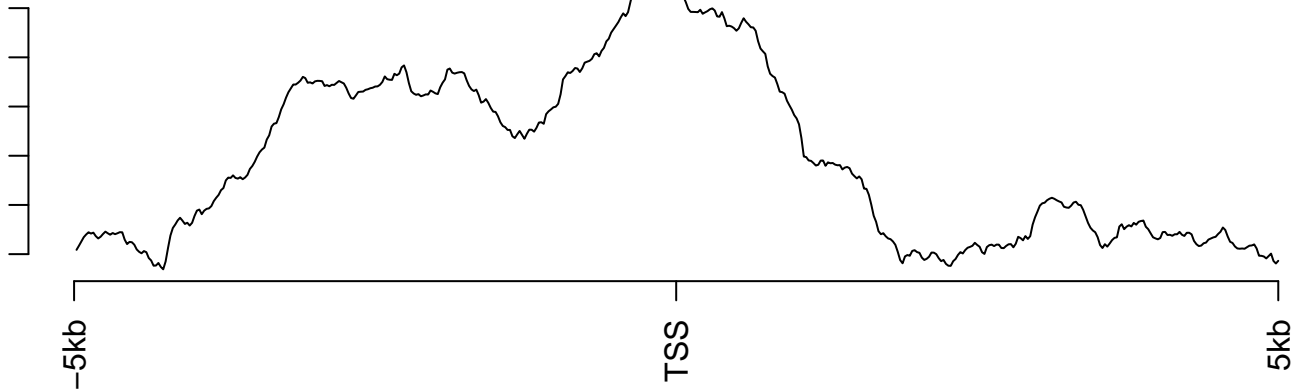
Average coverage per gene

0.020
0.026

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc625

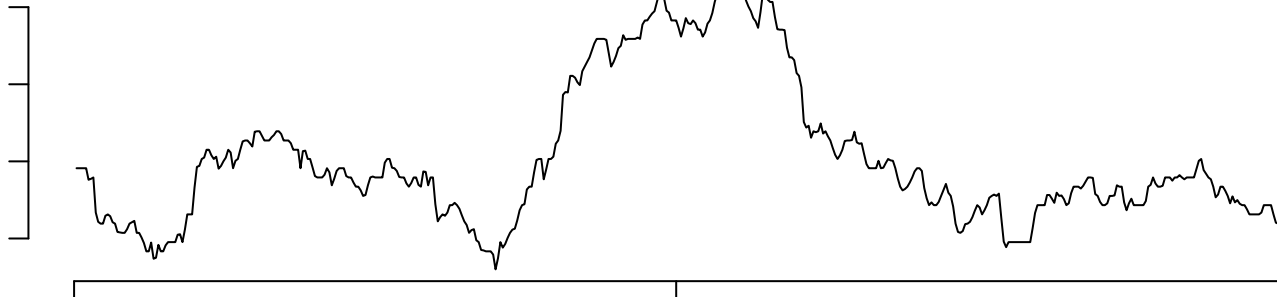
Average coverage per gene

0.0008 0.0012

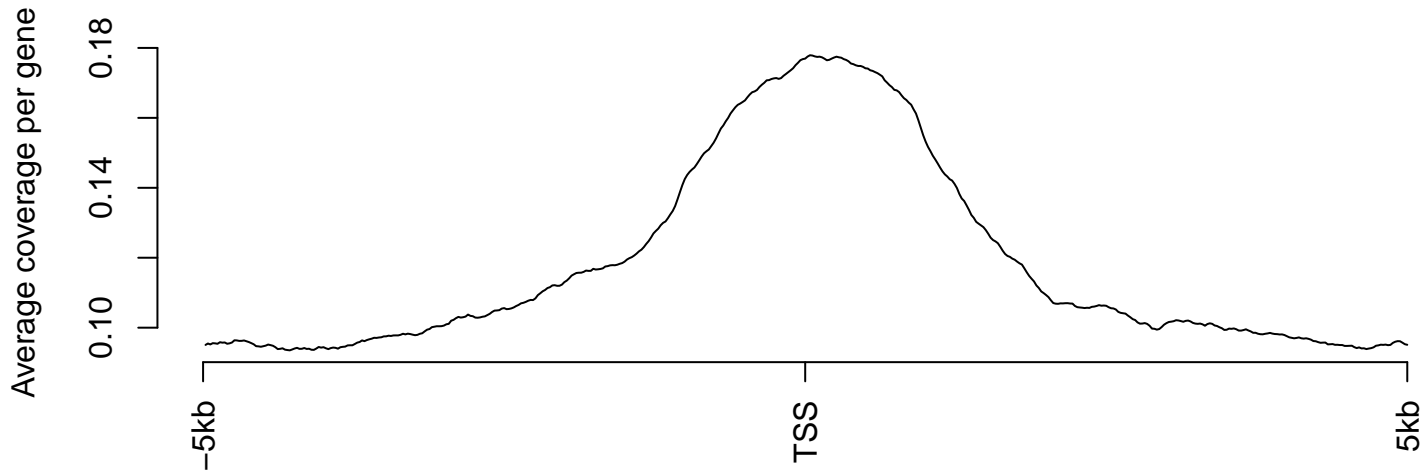
-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc626



MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc627

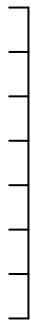
Average coverage per gene

0.065 0.080 0.095

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.653 scCLTdegenNuc663

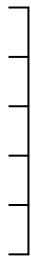
Average coverage per gene

0.0008 0.0014

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.653 scCLTdegenNuc664

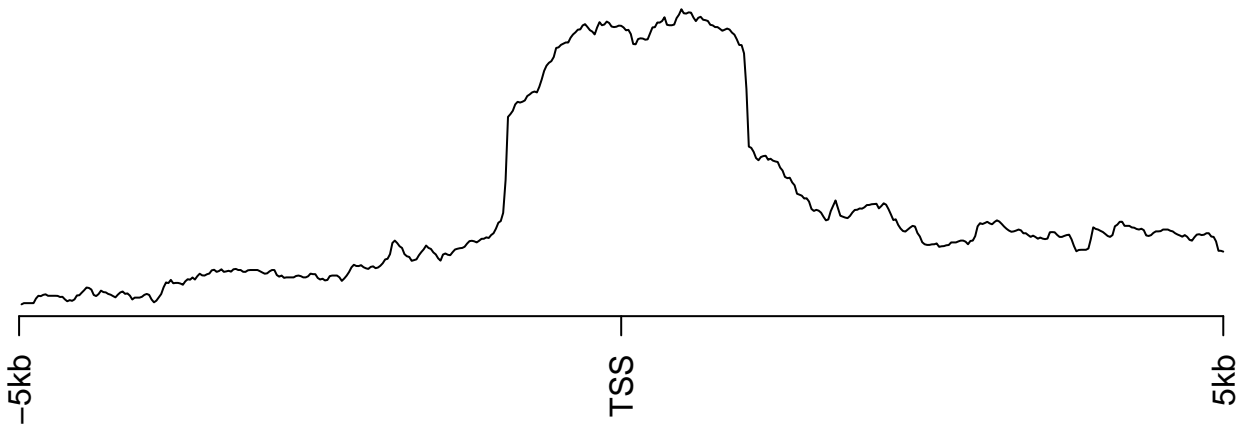
Average coverage per gene

0.002
0.004

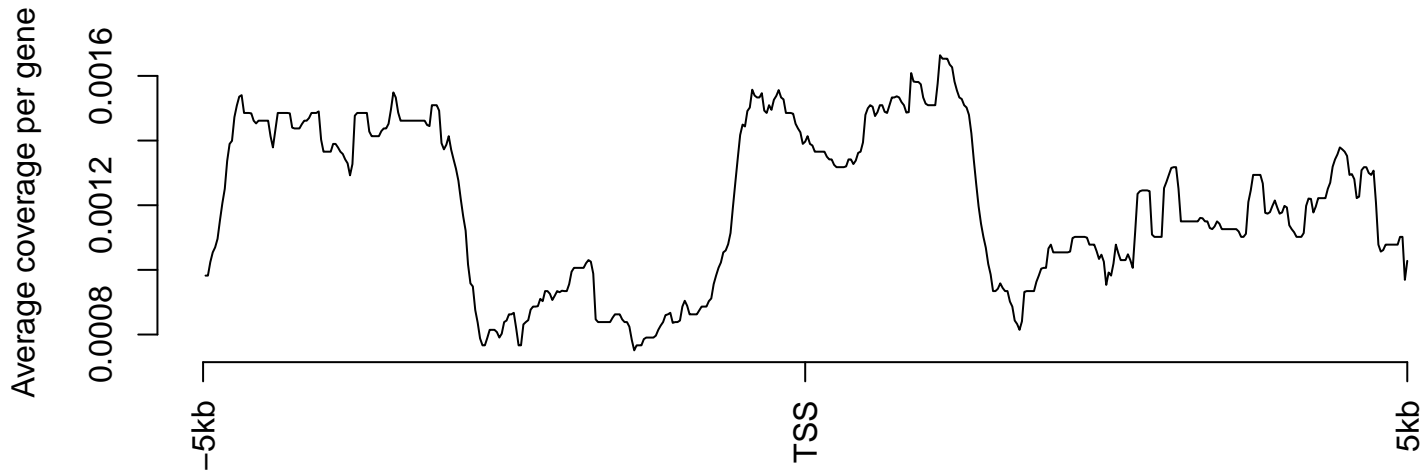
-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc697



MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc698

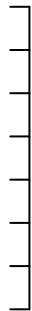
Average coverage per gene

0.0004 0.0007 0.0010

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc699

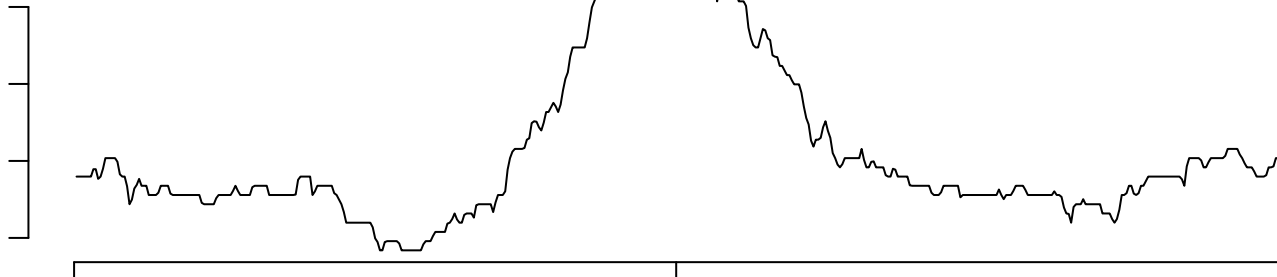
Average coverage per gene

2e-04
6e-04

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc776

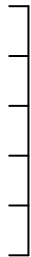
Average coverage per gene

0.0010 0.0016

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc777

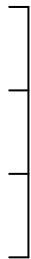
Average coverage per gene

0.0025
0.0015

-5kb

TSS

5kb



MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc778

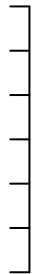
Average coverage per gene

0.0008 0.0014 0.0020

-5kb

TSS

5kb



MouseInterneuronSlice PositiveMerged E.– MouseInterneuronSlicePositiveMerged

