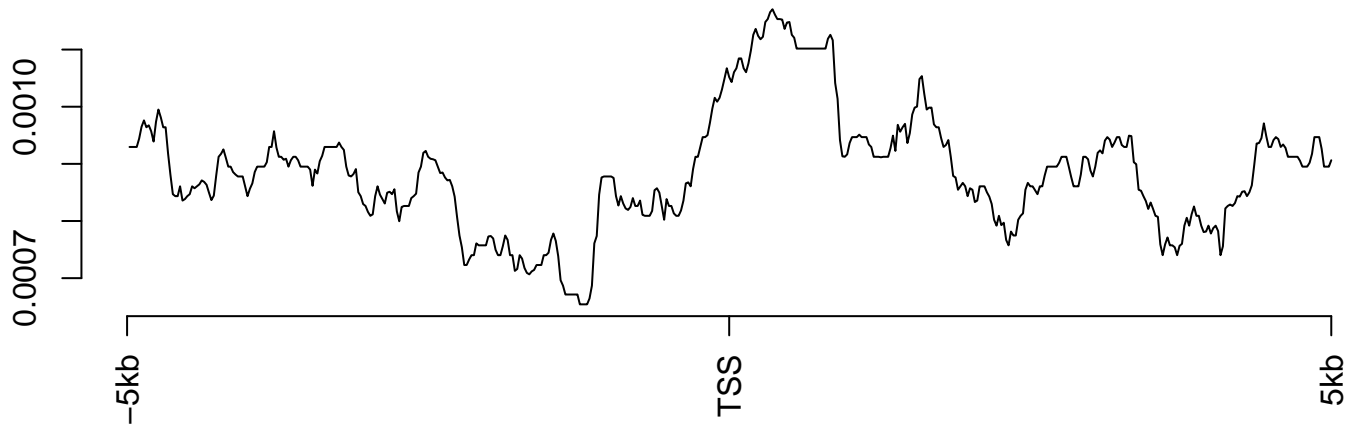


HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc735

Average coverage per gene



HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc736

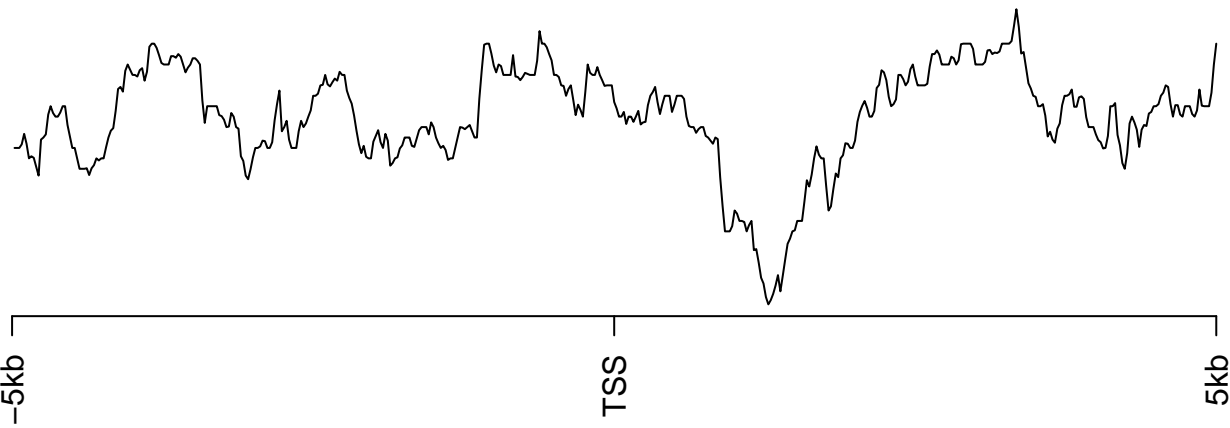
Average coverage per gene

0.0010
0.0007

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc737

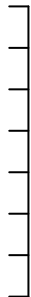
Average coverage per gene

0.0012
0.0008

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc738

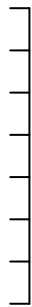
Average coverage per gene

0.0010 0.0016 0.0022

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc739

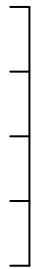
Average coverage per gene

0.0018
0.0014
0.0010

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc740

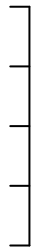
Average coverage per gene

0.024 0.028 0.032

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc741

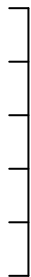
Average coverage per gene

0.0016 0.0022

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc742

Average coverage per gene

0.024 0.030

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc743

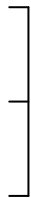
Average coverage per gene

0.0020
0.0030

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc744

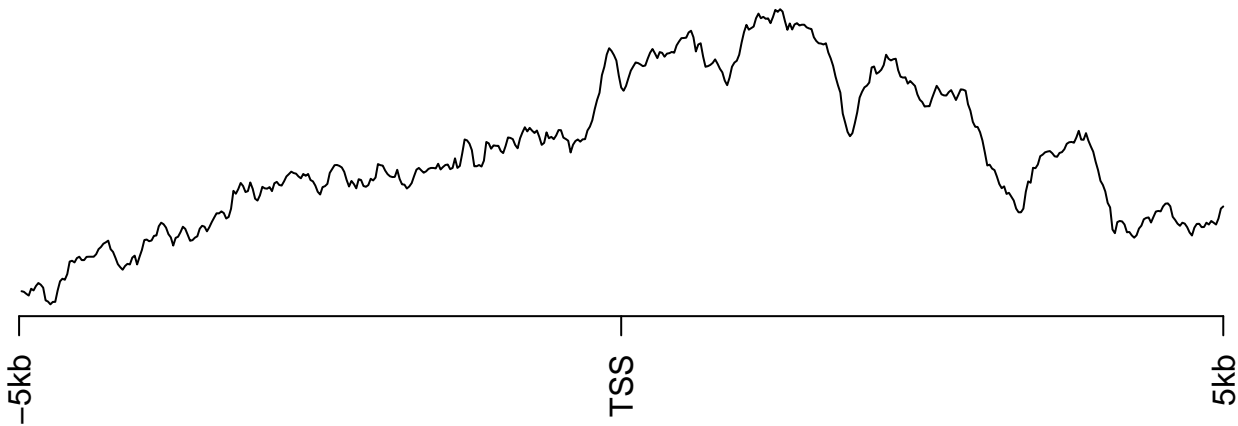
Average coverage per gene

0.0020
0.0030

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc745

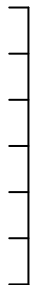
Average coverage per gene

0.0008 0.0014 0.0020

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc746

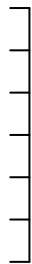
Average coverage per gene

0.0012 0.0018 0.0024

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc747

Average coverage per gene

0.0014 0.0018

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc748

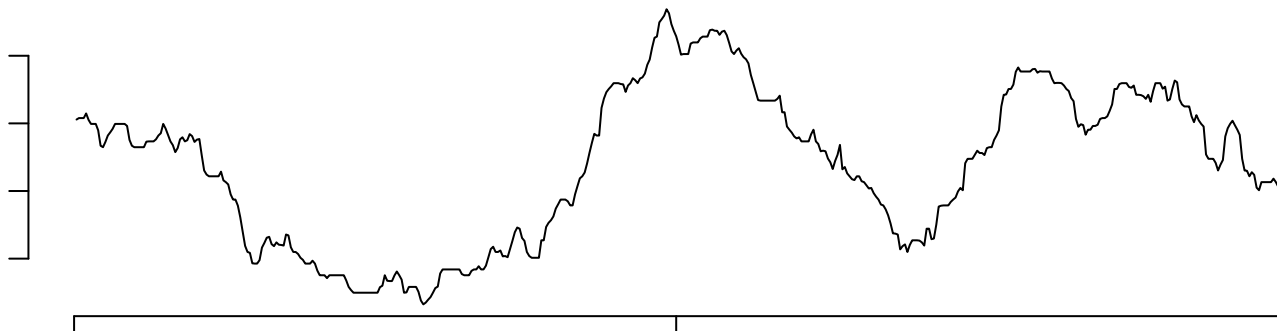
Average coverage per gene

0.0006 0.0010

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc749

Average coverage per gene

0.0015 0.0025

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc750

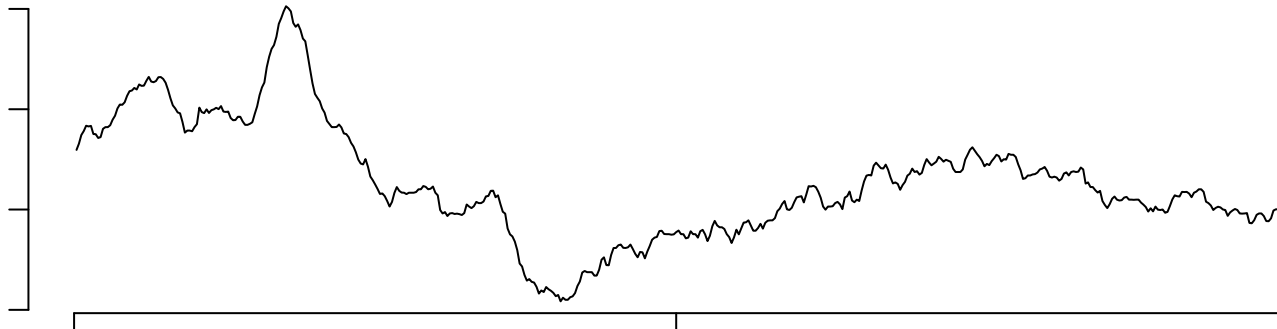
Average coverage per gene

0.0020
0.0010

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc751

Average coverage per gene

0.0032
0.0024

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc752

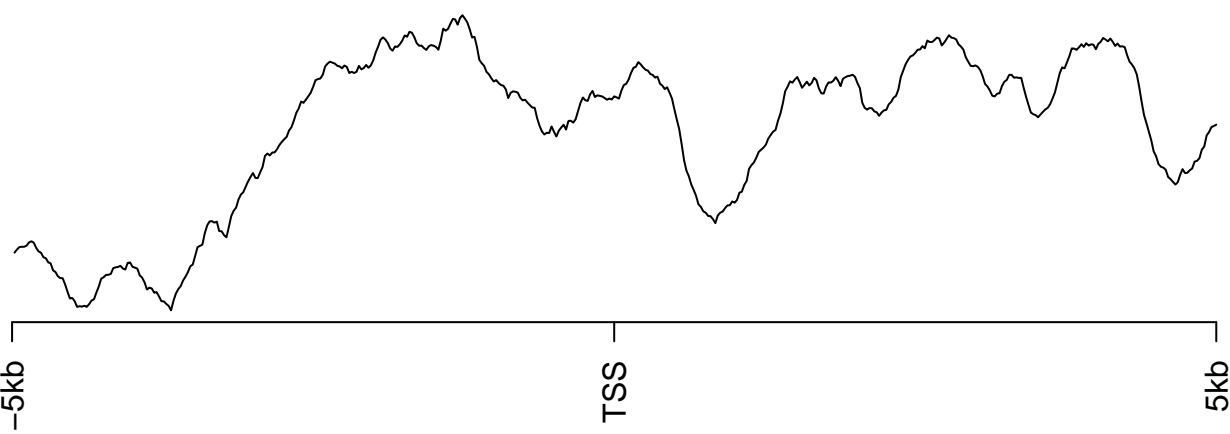
Average coverage per gene

0.011
0.009

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc753

Average coverage per gene

0.0004 0.0010 0.0016

-5kb

TSS

5kb



HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc754

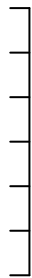
Average coverage per gene

0.0007 0.0010 0.0013

-5kb

TSS

5kb



HumanInterneuronCulture PositiveMerged E.– HumanInterneuronCulturePositiveMerged

Average coverage per gene

0.100
0.090

-5kb

TSS

5kb

