

HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc473

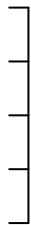
Average coverage per gene

0.0018
0.0012

-5kb

TSS

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc474

Average coverage per gene

0.0025 0.0035

-5kb

TSS

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc475

Average coverage per gene

0.0008 0.0014

-5kb

TSS

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc476

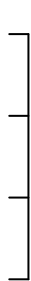
Average coverage per gene

0.0018 0.0022

-5kb

TSS

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc477

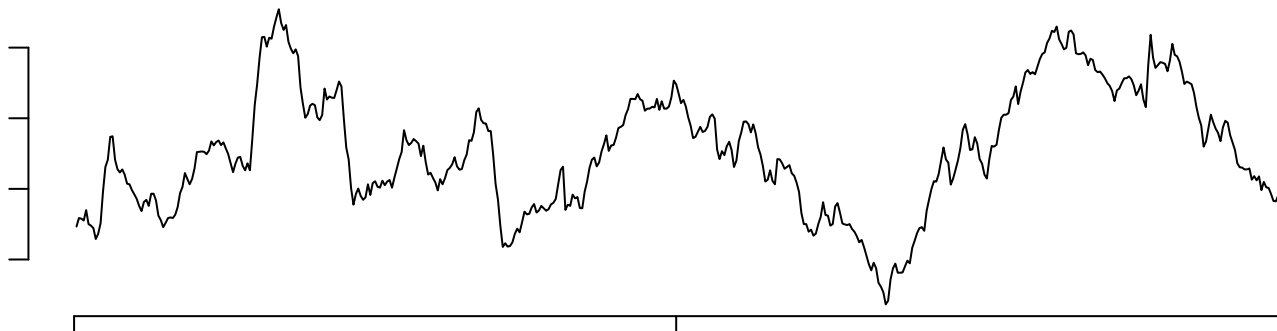
Average coverage per gene

0.0020 0.0024

-5kb

TSS

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc478

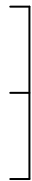
Average coverage per gene

0.0020 0.0030

-5kb

TSS

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc480

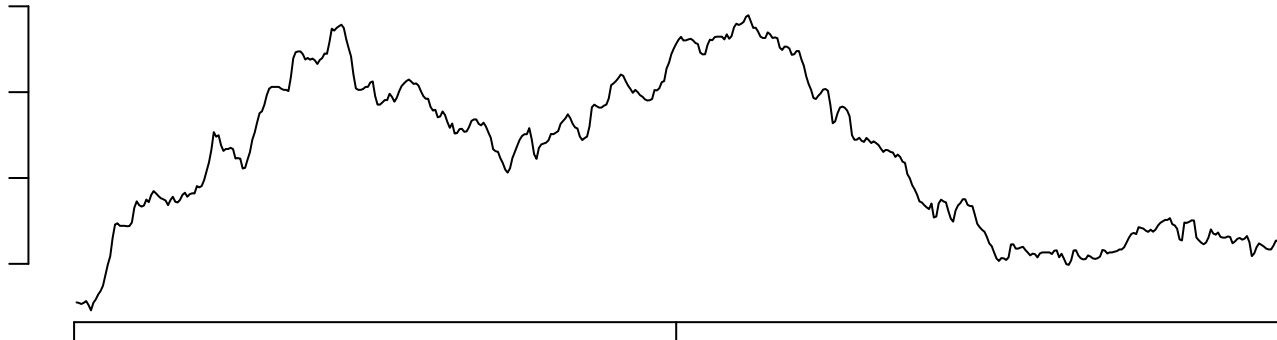
Average coverage per gene

0.0015
0.0025

-5kb

TSS

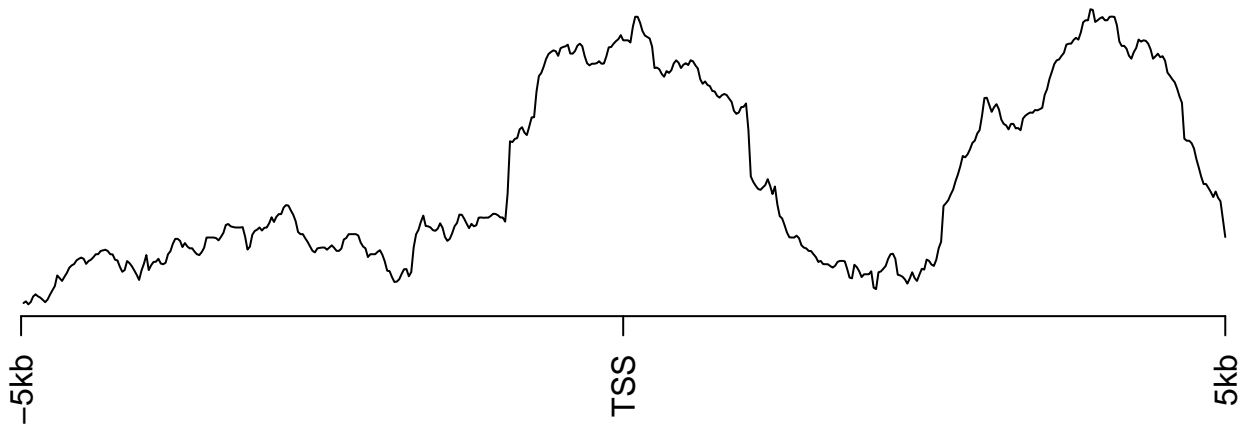
5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc481

Average coverage per gene

0.0015 0.0025



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc482

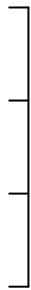
Average coverage per gene

0.0010
0.0020

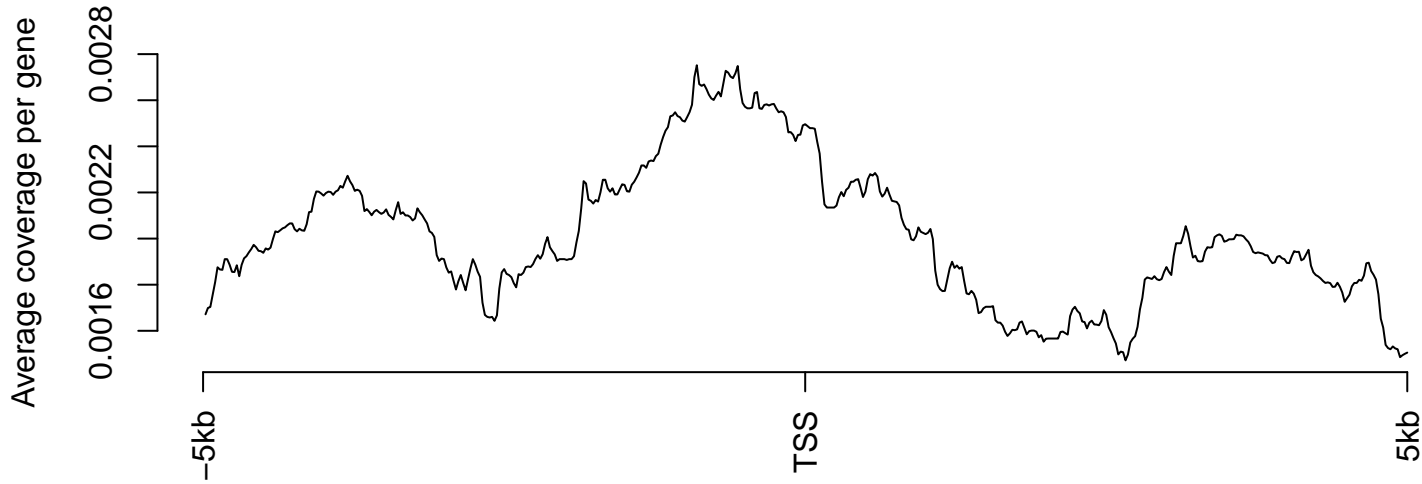
-5kb

TSS

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc483



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc484

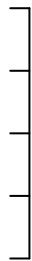
Average coverage per gene

0.0018
0.0014
0.0010

-5kb

TSS

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc485

Average coverage per gene

0.0008 0.0011 0.0014

-5kb

TSS

5kb



HumanNeuronCulture PositiveMerged E.– HumanNeuronCulturePositiveMerged

Average coverage per gene

0.018 0.022 0.026

-5kb

TSS

5kb

