

HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc473

Average coverage per gene

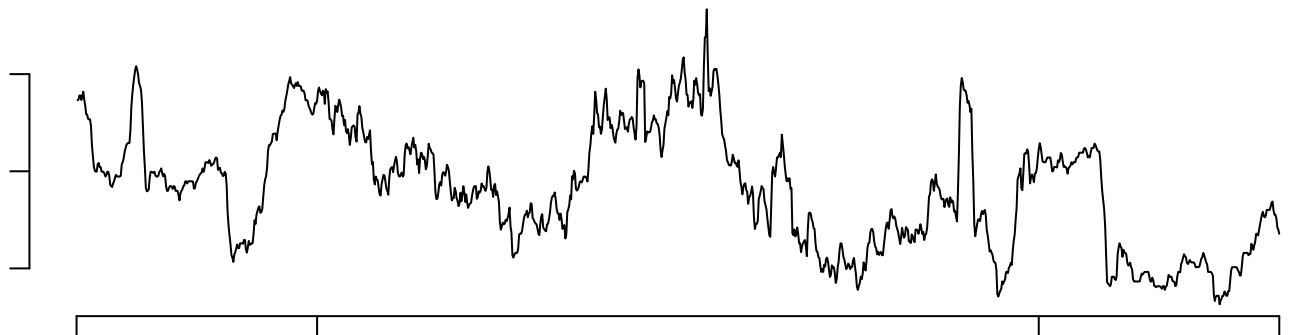
0.0020
0.0010

-5kb

Gene start

Gene end

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc474

Average coverage per gene

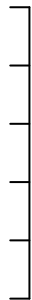
0.0030
0.0015

-5kb

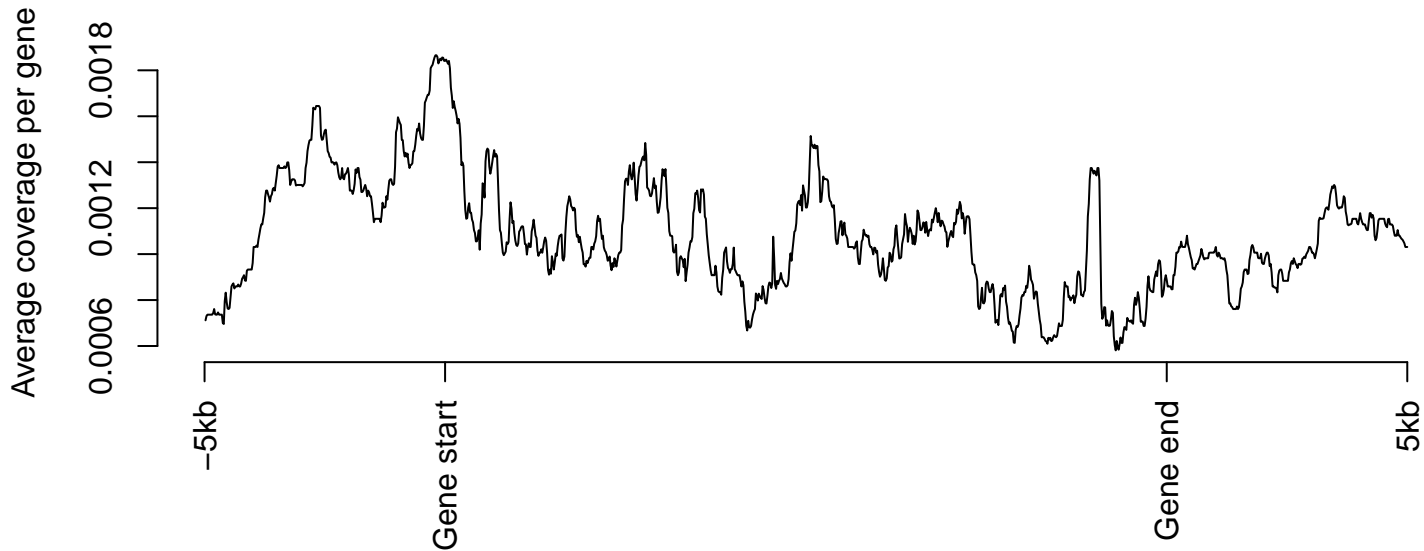
Gene start

Gene end

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc475



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc476

Average coverage per gene

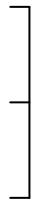
0.0025
0.0015

-5kb

Gene start

Gene end

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc477

Average coverage per gene

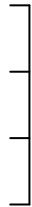
0.0015 0.0025

-5kb

Gene start

Gene end

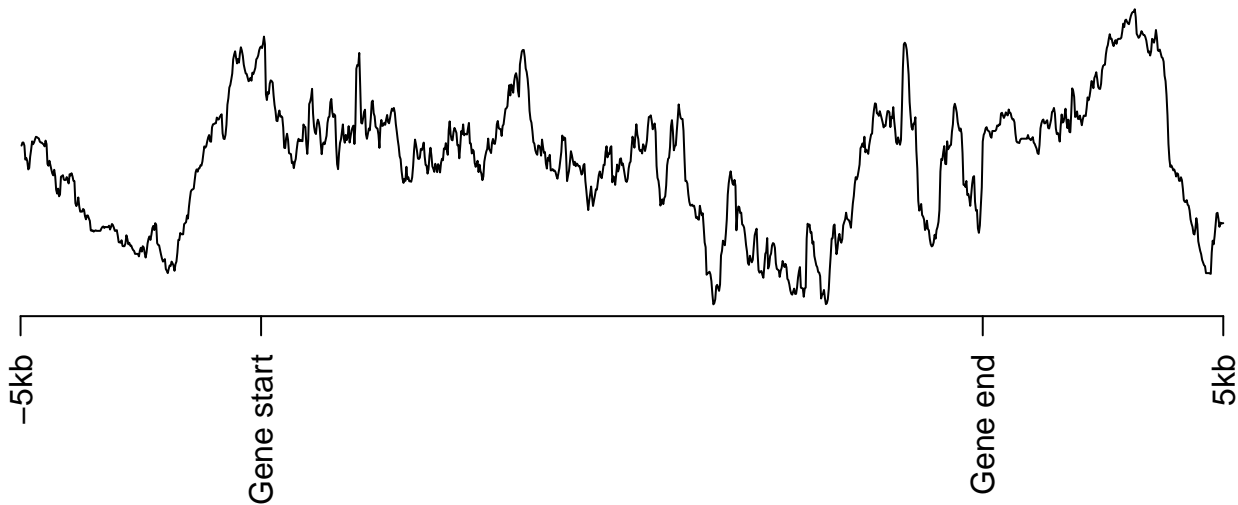
5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc478

Average coverage per gene

0.0015 0.0025



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc480

Average coverage per gene

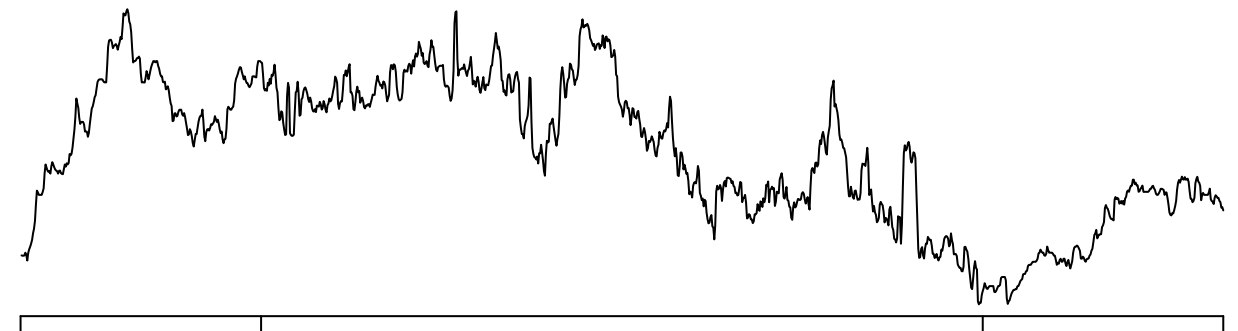
0.0010
0.0025

-5kb

Gene start

Gene end

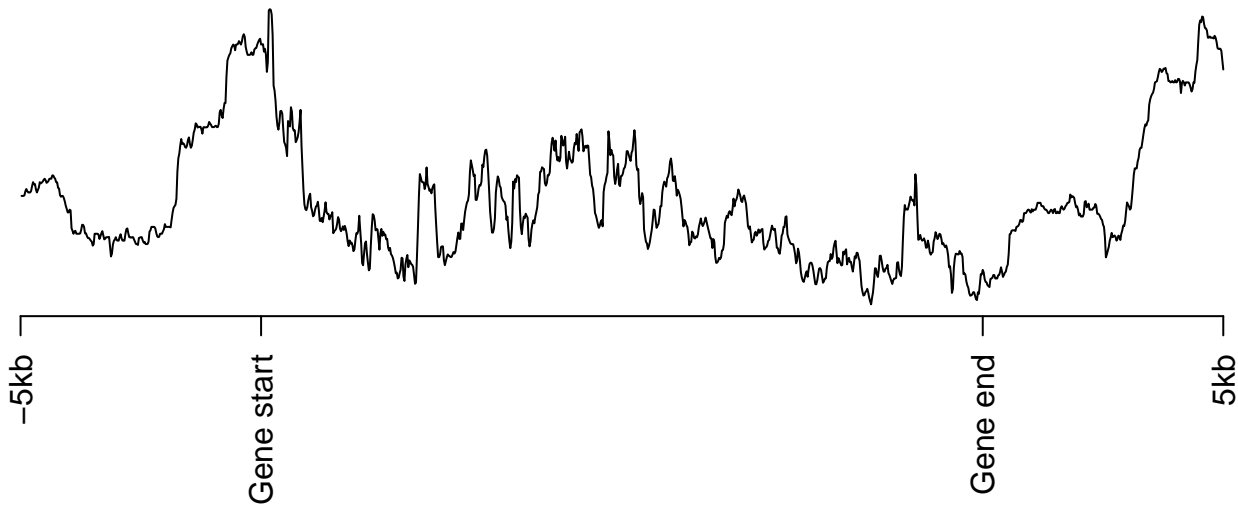
5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc481

Average coverage per gene

0.0010 0.0020



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc482

Average coverage per gene

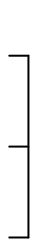
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0.0010

-5kb

Gene start

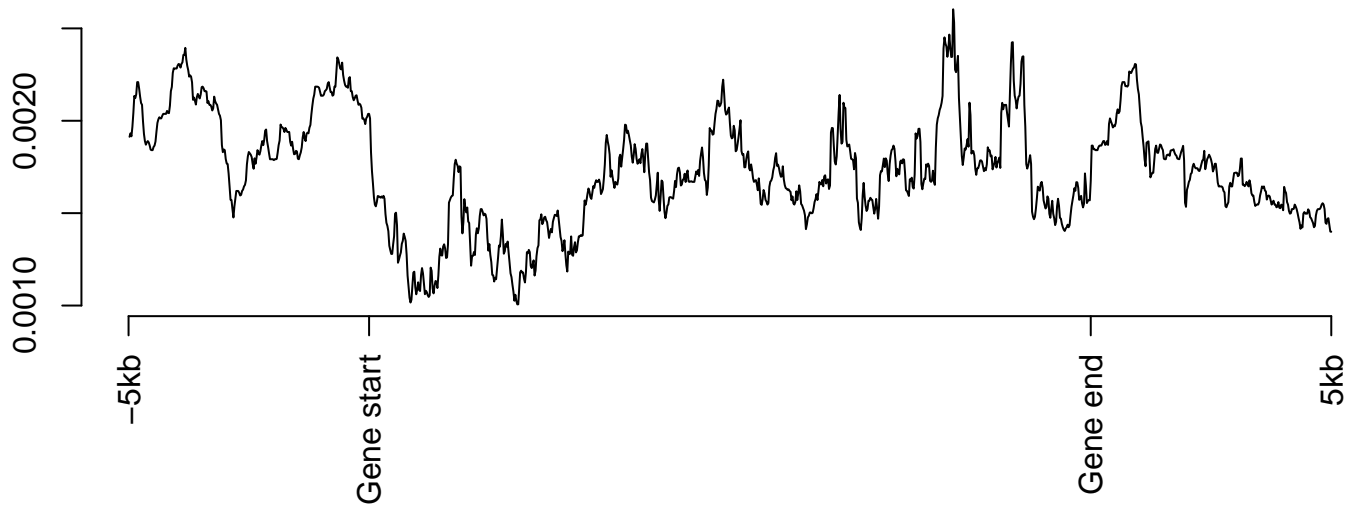
Gene end

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc483

Average coverage per gene



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc484

Average coverage per gene

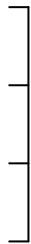
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-5kb

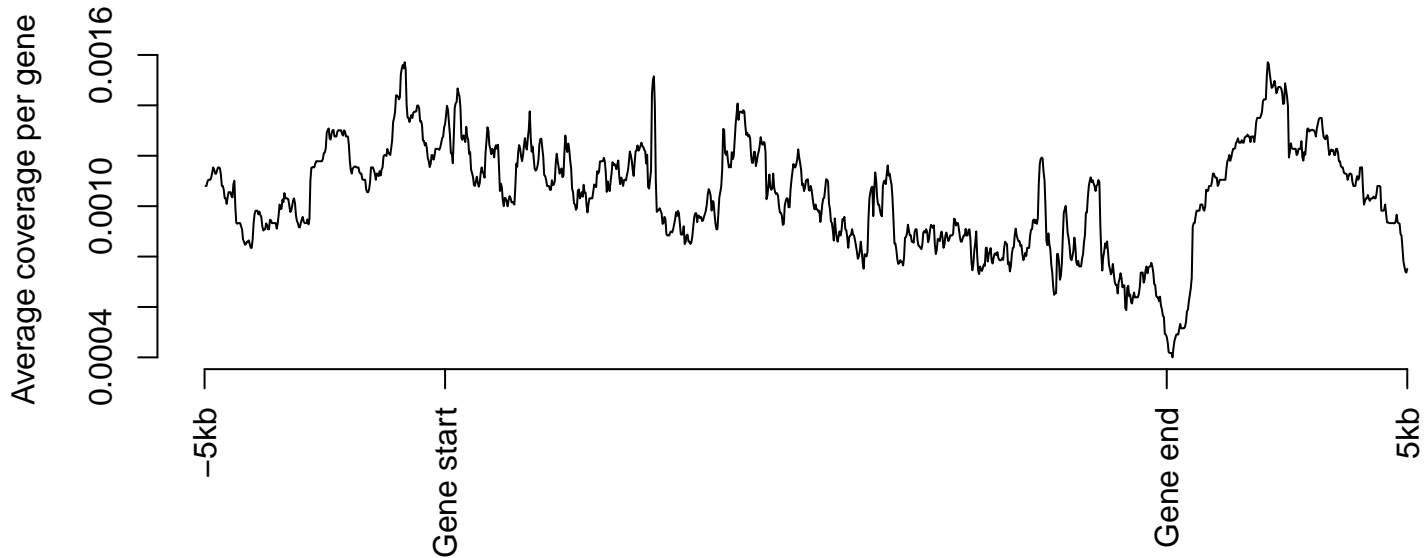
Gene start

Gene end

5kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc485



HumanNeuronCulture PositiveMerged E.– HumanNeuronCulturePositiveMerged

Average coverage per gene

0.022
0.016

-5kb

Gene start

Gene end

5kb

