

HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc473

Average coverage per Exon

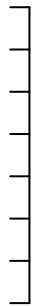
0.00145 0.00165

-500bp

Exon start

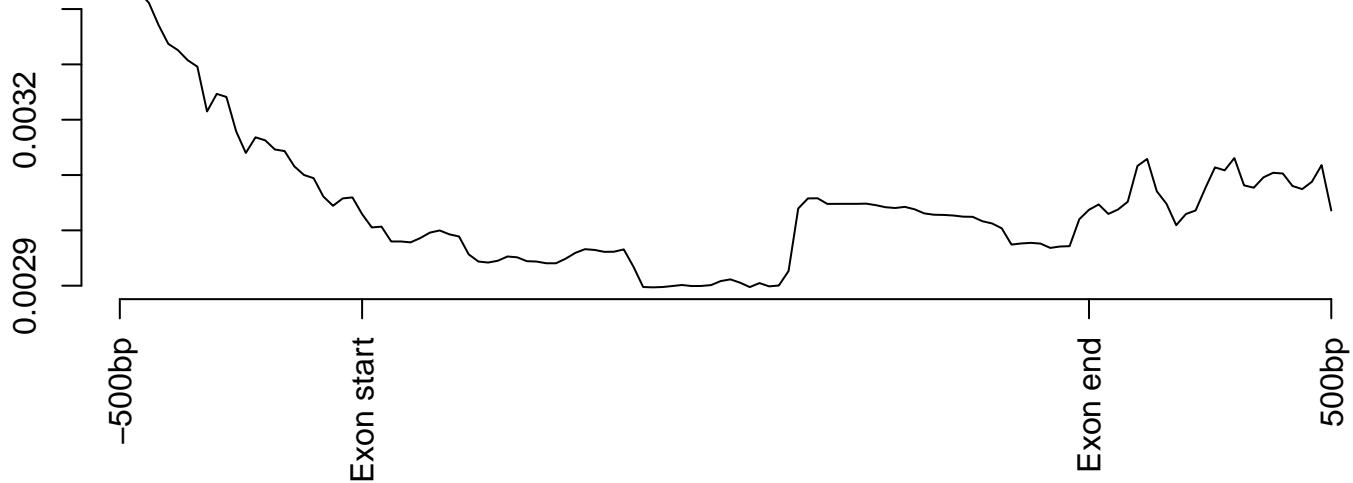
Exon end

500bp

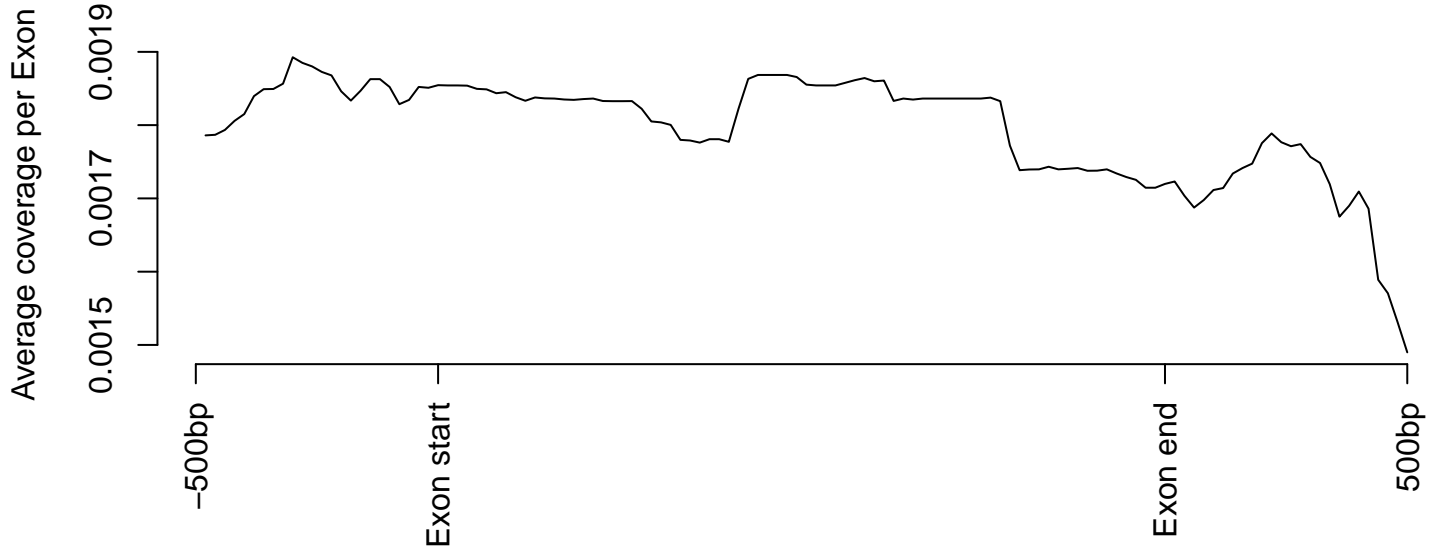


HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc474

Average coverage per Exon



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc475



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc476

Average coverage per Exon

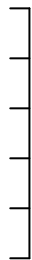
0.00145 0.00160

-500bp

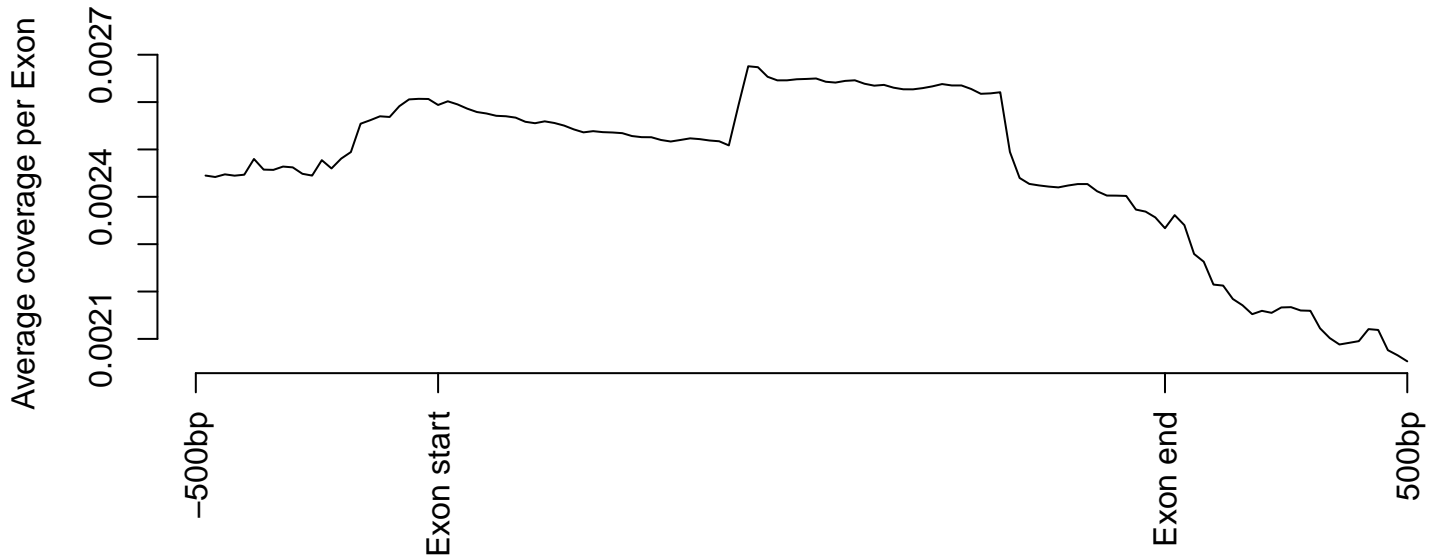
Exon start

Exon end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc477



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc478

Average coverage per Exon

0.0023 0.0026

-500bp

Exon start

Exon end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc480

Average coverage per Exon

0.0027 0.0029

-500bp

Exon start

Exon end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc481

Average coverage per Exon

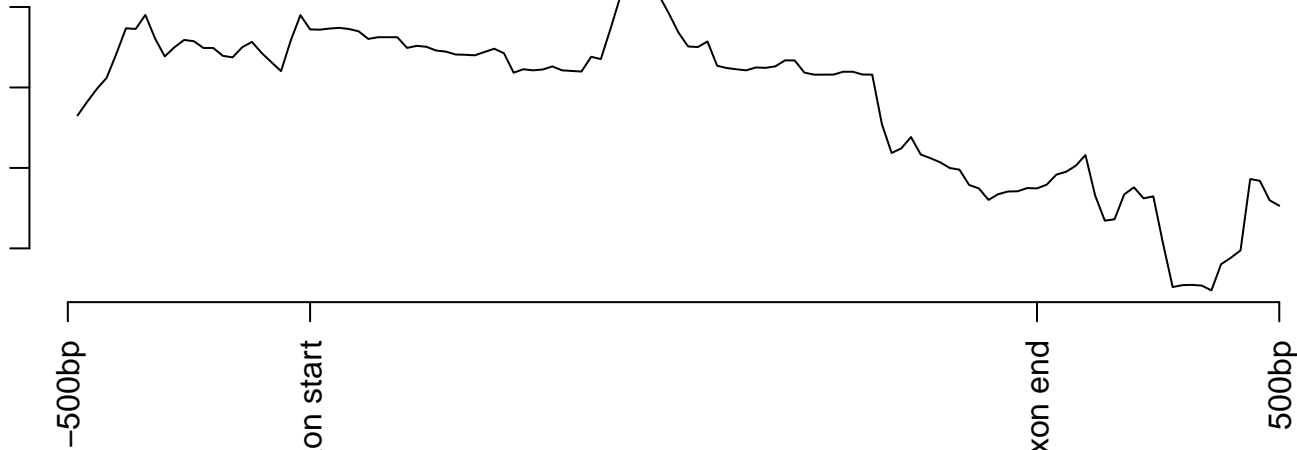
0.0023
0.0021

-500bp

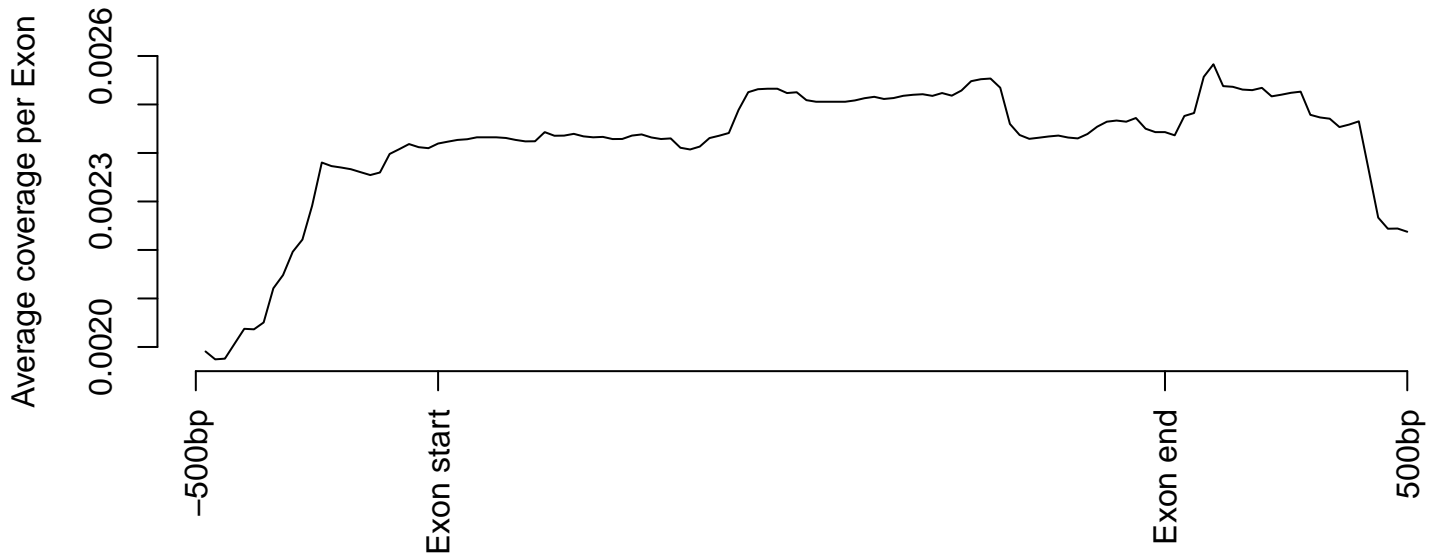
Exon start

Exon end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc482



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc483

Average coverage per Exon

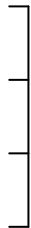
0.0017 0.0019

-500bp

Exon start

Exon end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc484

Average coverage per Exon

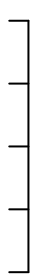
0.00165
0.00150

-500bp

Exon start

Exon end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc485

Average coverage per Exon

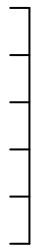
0.00125 0.00145

-500bp

Exon start

Exon end

500bp



HumanNeuronCulture PositiveMerged E.– HumanNeuronCulturePositiveMerged

