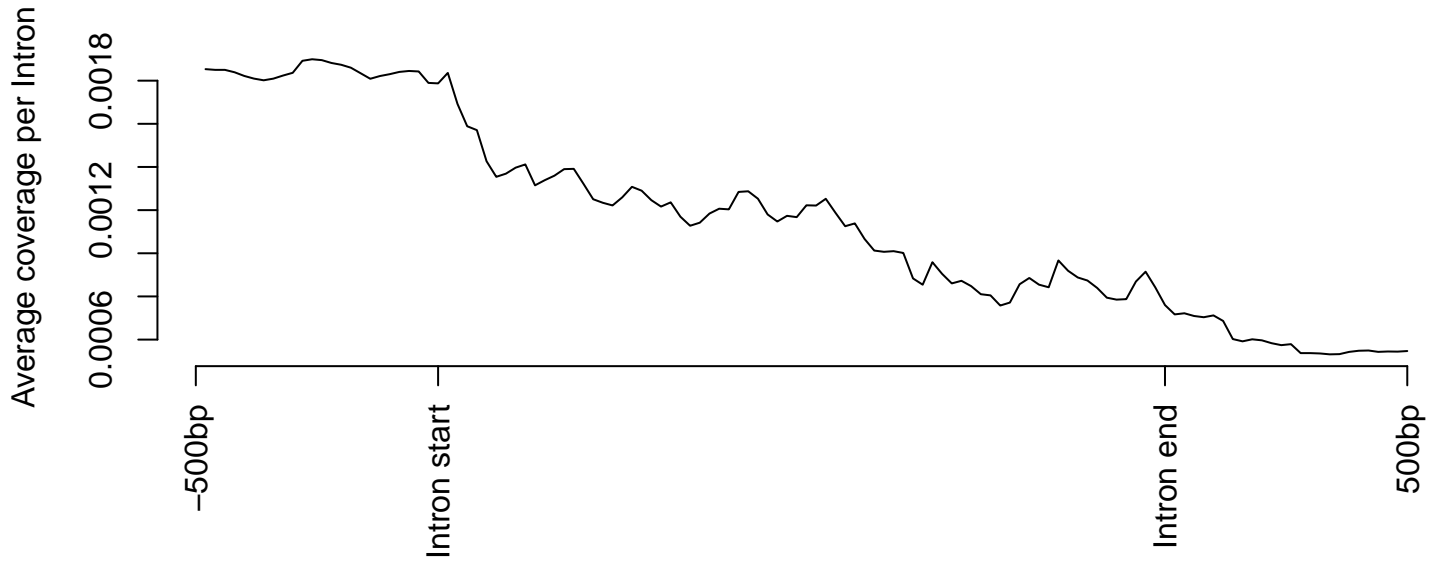


HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc473



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc474

Average coverage per Intron

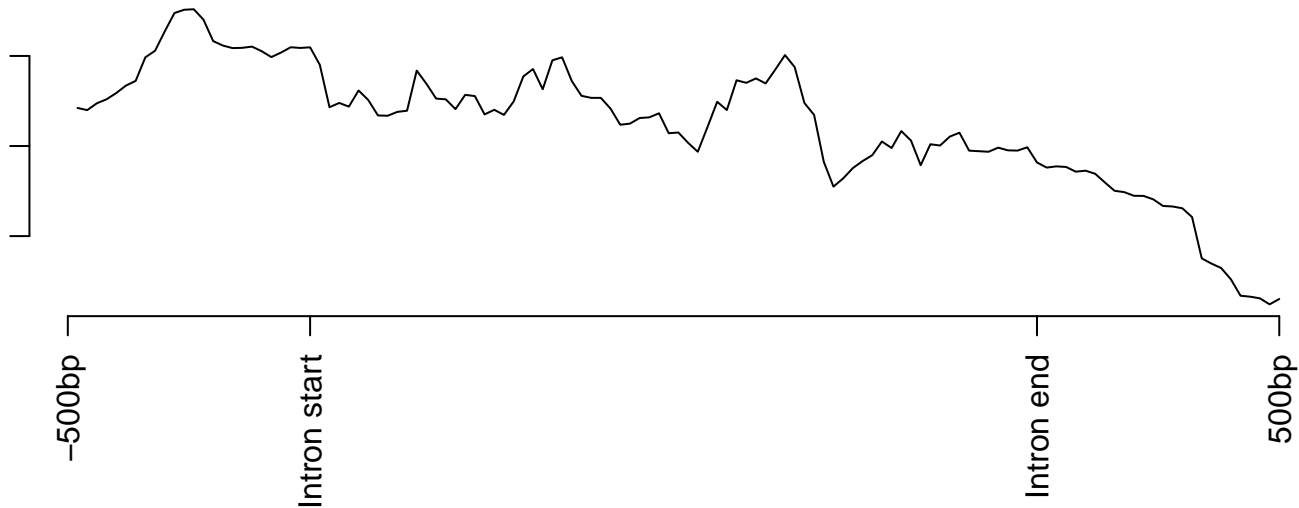
0.0030
0.0020

-500bp

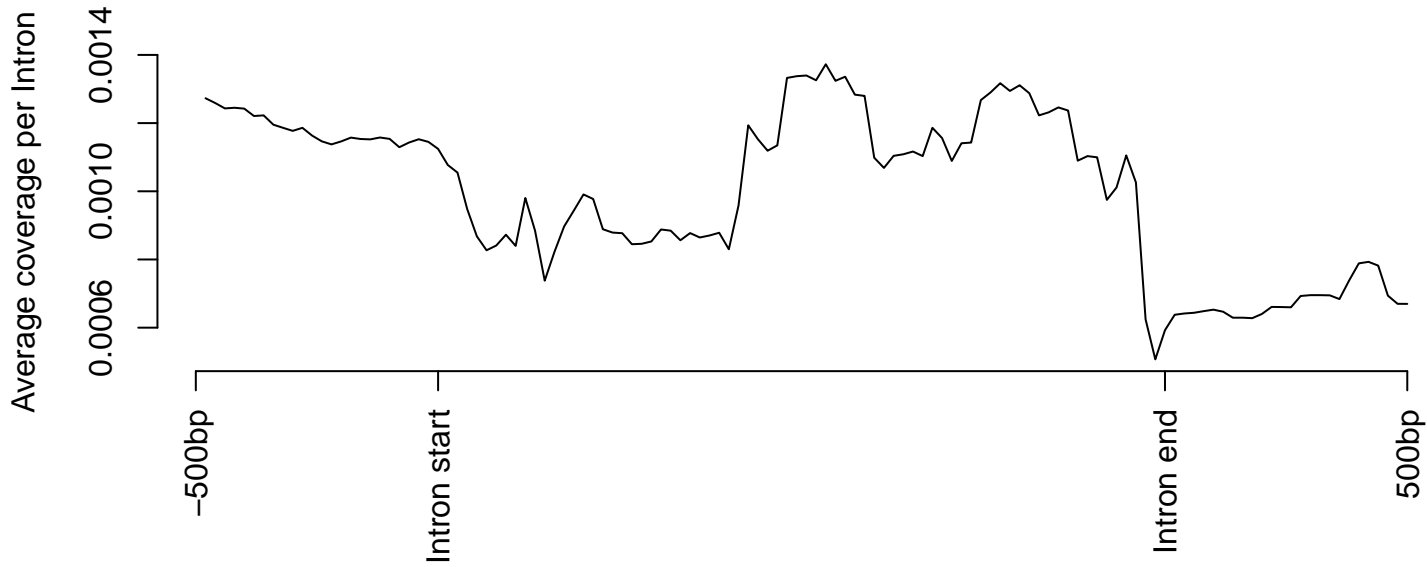
Intron start

Intron end

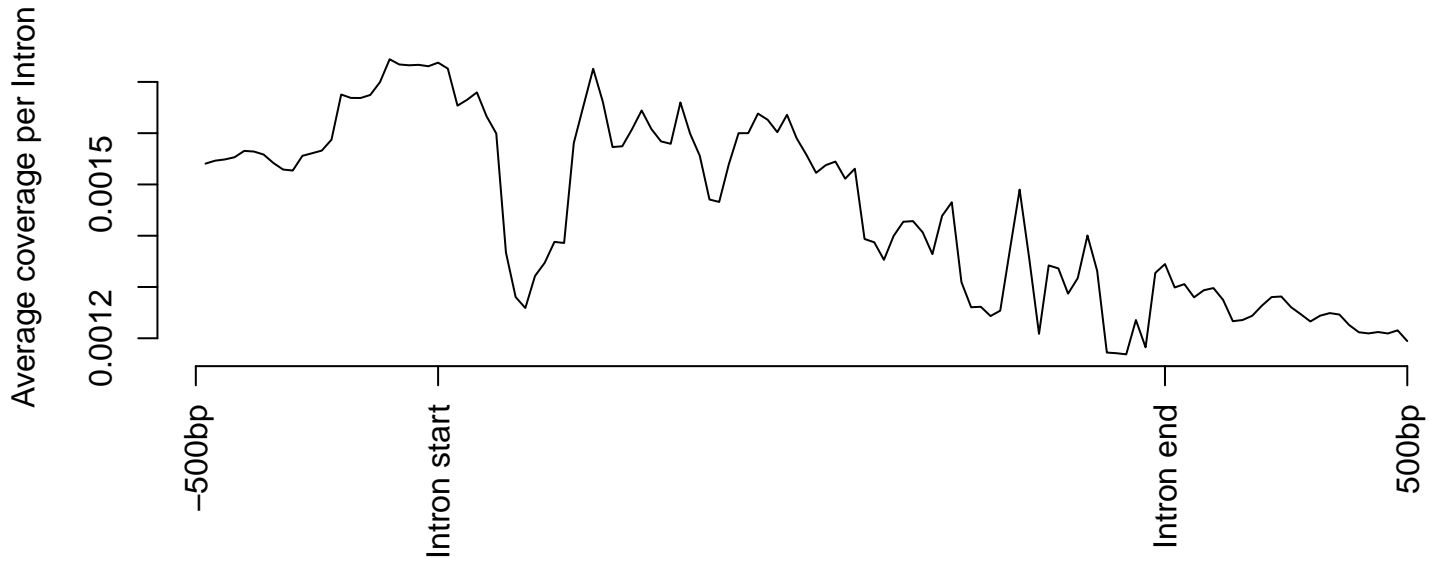
500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc475



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc476



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc477

Average coverage per Intron

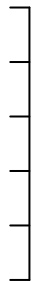
0.0016 0.0022

-500bp

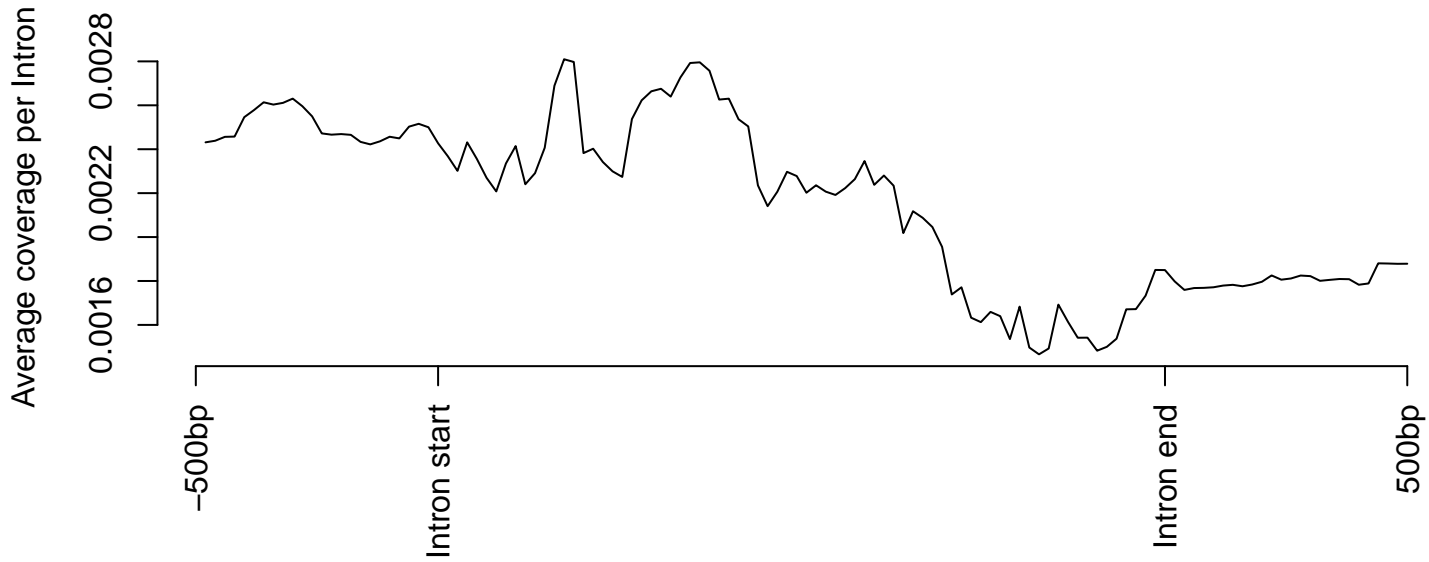
Intron start

Intron end

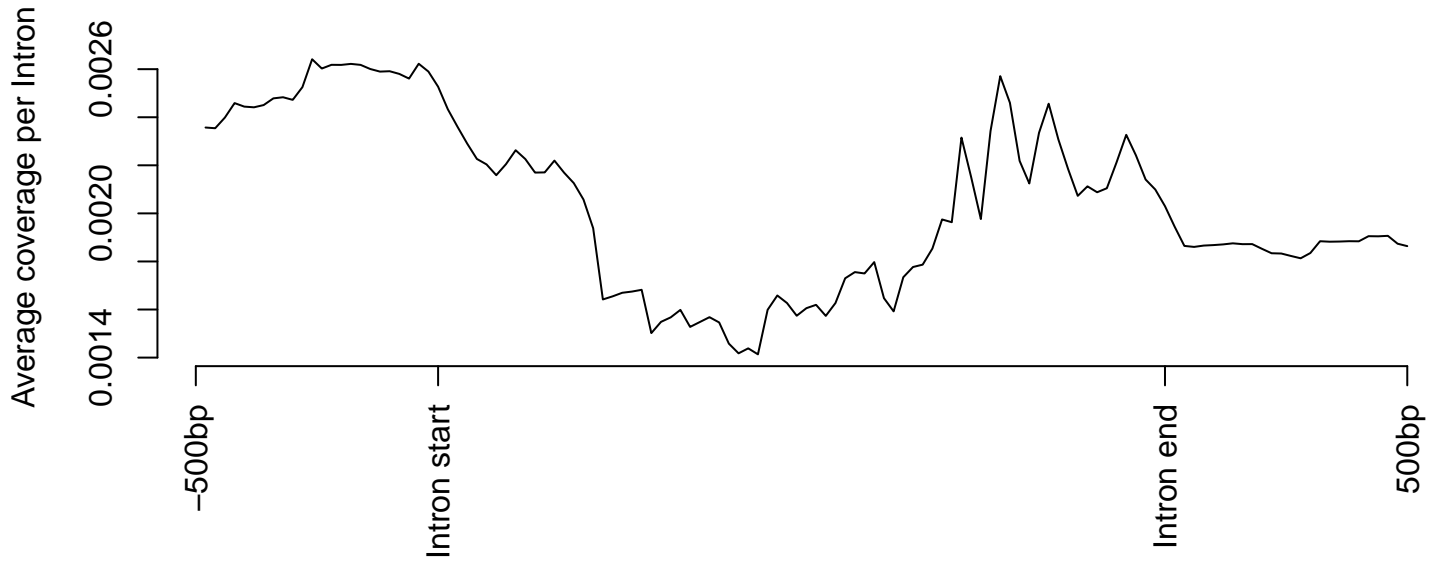
500bp



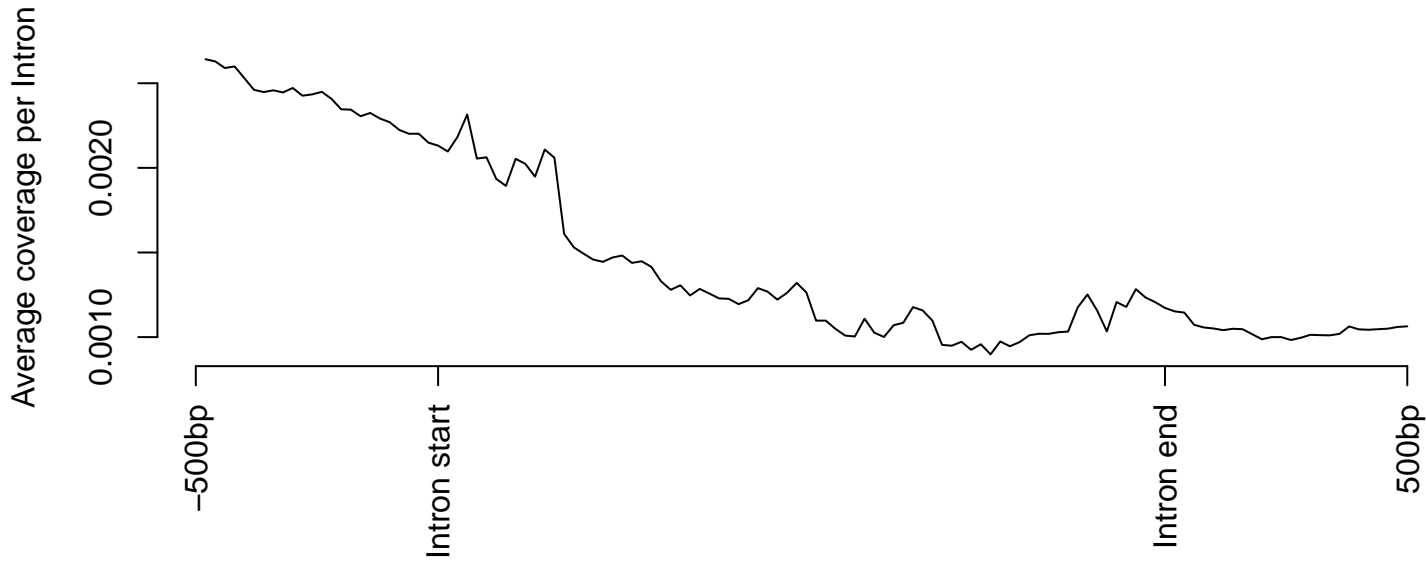
HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc478



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc480



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc481



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc482

Average coverage per Intron

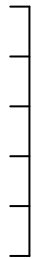
0.0016
0.0010

-500bp

Intron start

Intron end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc483

Average coverage per Intron

0.0009 0.0013

-500bp

Intron start

Intron end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc484

Average coverage per Intron

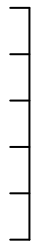
0.0014
0.0011

-500bp

Intron start

Intron end

500bp



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc485

Average coverage per Intron

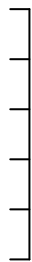
0.0008 0.0011

-500bp

Intron start

Intron end

500bp



HumanNeuronCulture PositiveMerged E.– HumanNeuronCulturePositiveMerged

