

HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc473

Average coverage per Intron

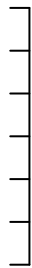
0.0018
0.0012
0.0006

-3kb

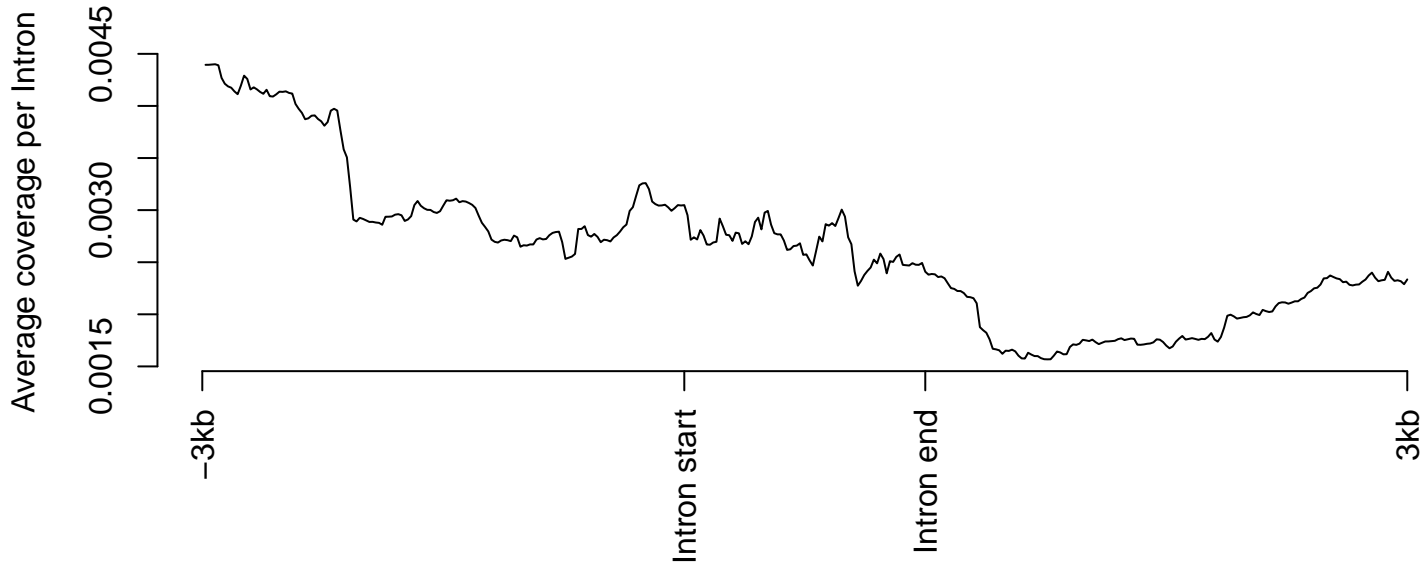
Intron start

Intron end

3kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc474



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc475

Average coverage per Intron

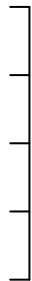
0.0014
0.0010
0.0006

-3kb

Intron start

Intron end

3kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc476

Average coverage per Intron

0.0018
0.0012

-3kb

Intron start

Intron end

3kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc477

Average coverage per Intron

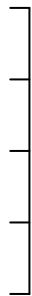
0.0035
0.0025
0.0015

-3kb

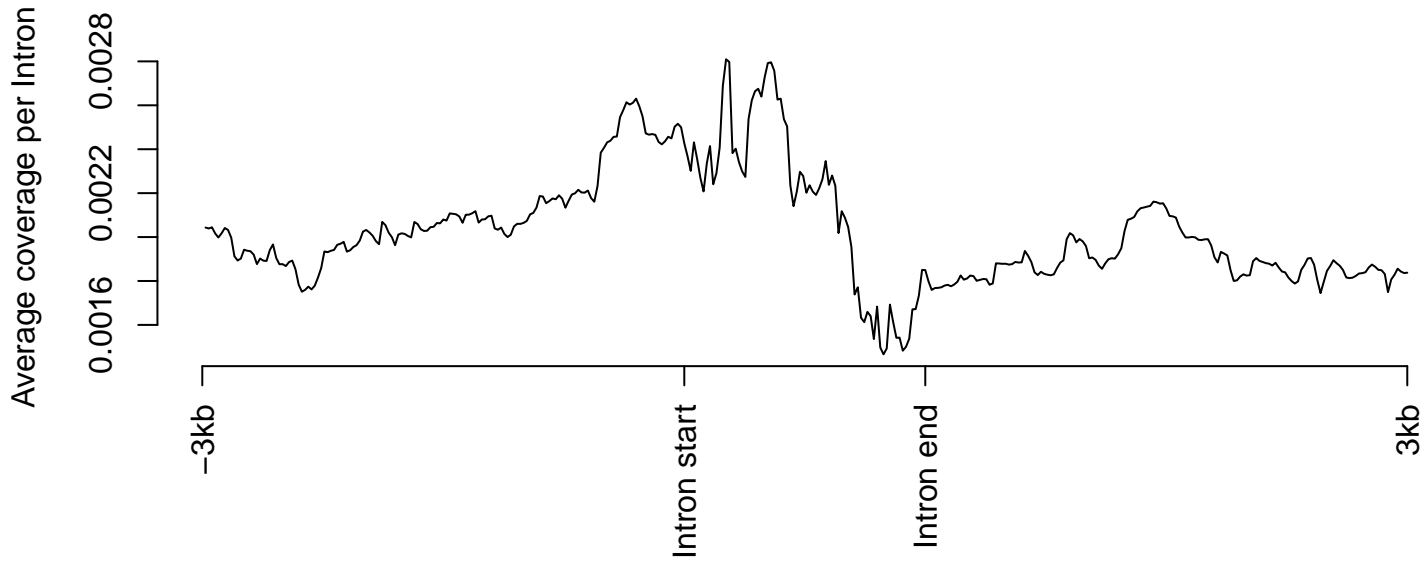
Intron start

Intron end

3kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc478



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc480

Average coverage per Intron

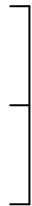
0.0015
0.0025

-3kb

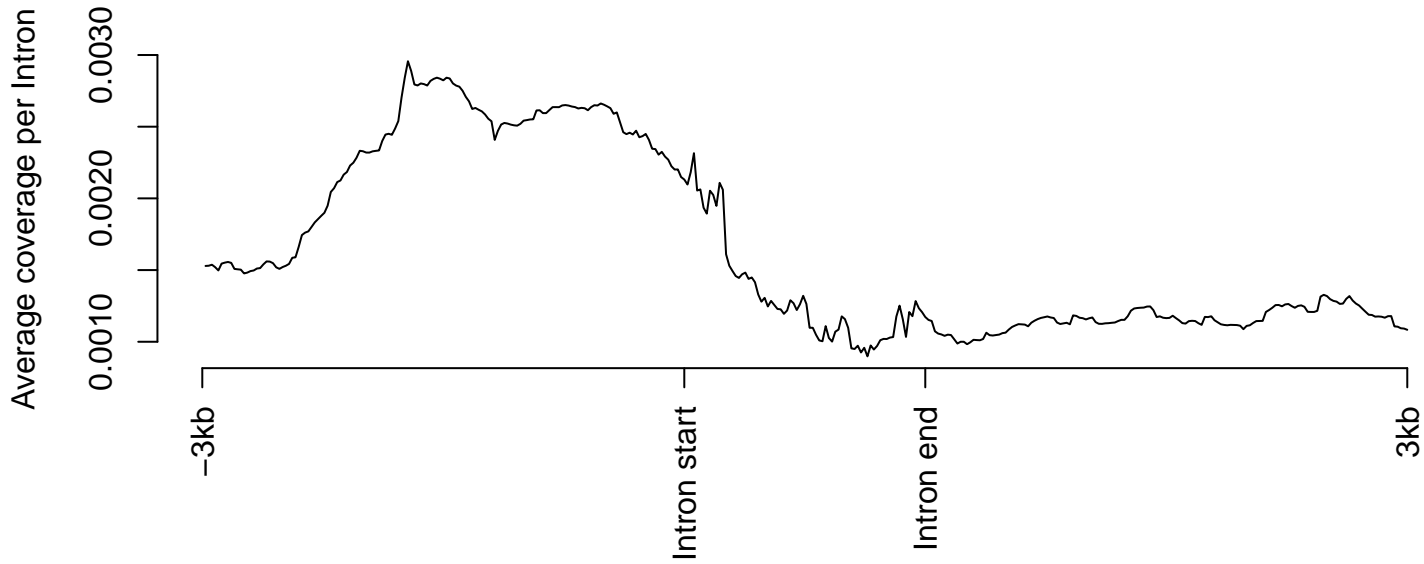
Intron start

Intron end

3kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc481



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc482

Average coverage per Intron

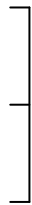
0.0020
0.0010

-3kb

Intron start

Intron end

3kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc483

Average coverage per Intron

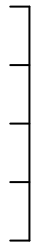
0.0016
0.0010

-3kb

Intron start

Intron end

3kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc484

Average coverage per Intron

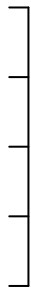
0.0008 0.0012 0.0016

-3kb

Intron start

Intron end

3kb



HumanNeuronCulture PositiveSingle E.602 scCLTdegenNuc485

Average coverage per Intron

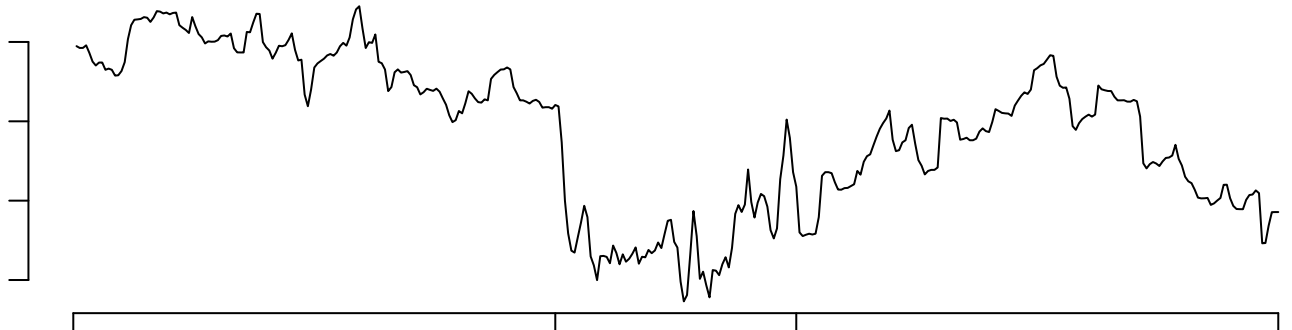
0.0012
0.0008

-3kb

Intron start

Intron end

3kb



HumanNeuronCulture PositiveMerged E.– HumanNeuronCulturePositiveMerged

Average coverage per Intron

0.024
0.020
0.016

-3kb

Intron start

Intron end

3kb

