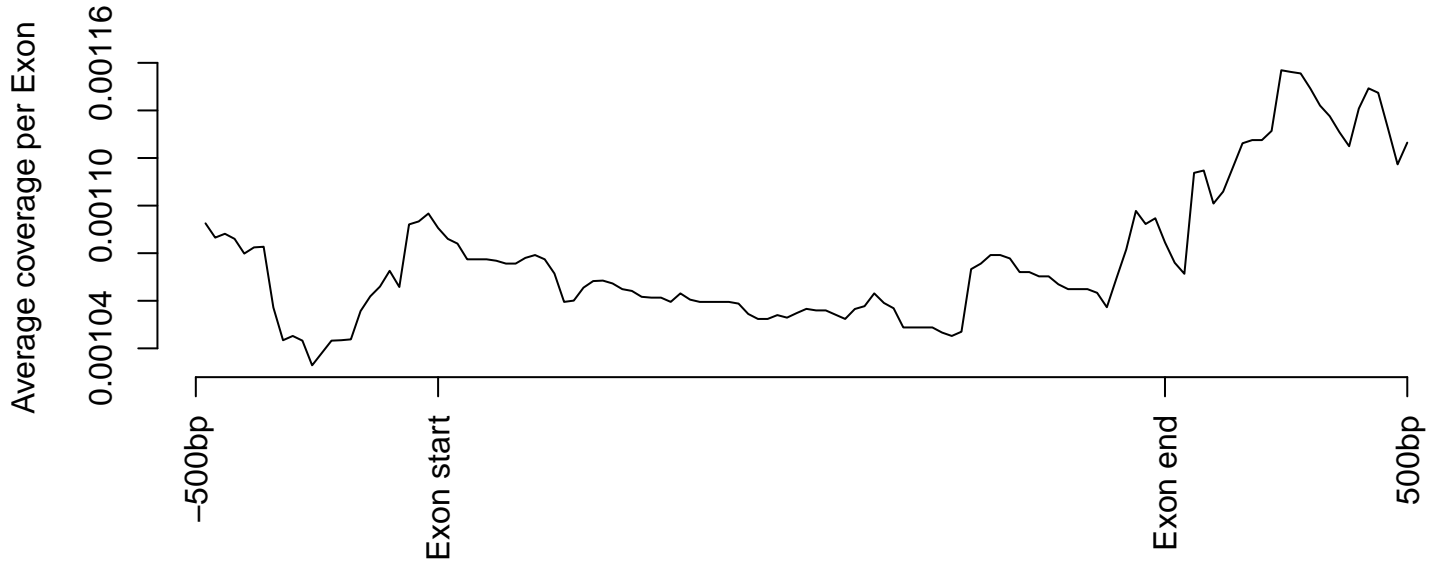


# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc735



# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc736

Average coverage per Exon

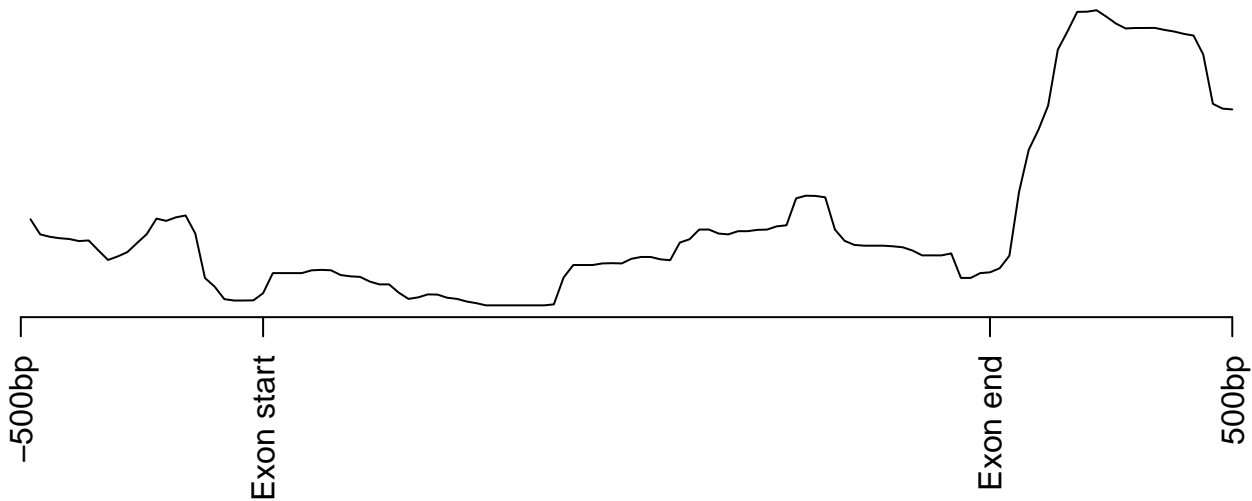
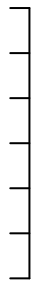
0.00120  
0.00100

-500bp

Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc737

Average coverage per Exon

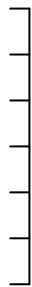
0.00170  
0.00150

-500bp

Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc738

Average coverage per Exon

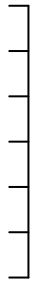
0.00160 0.00180

-500bp

Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc739

Average coverage per Exon

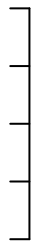
0.00130  
0.00115

-500bp

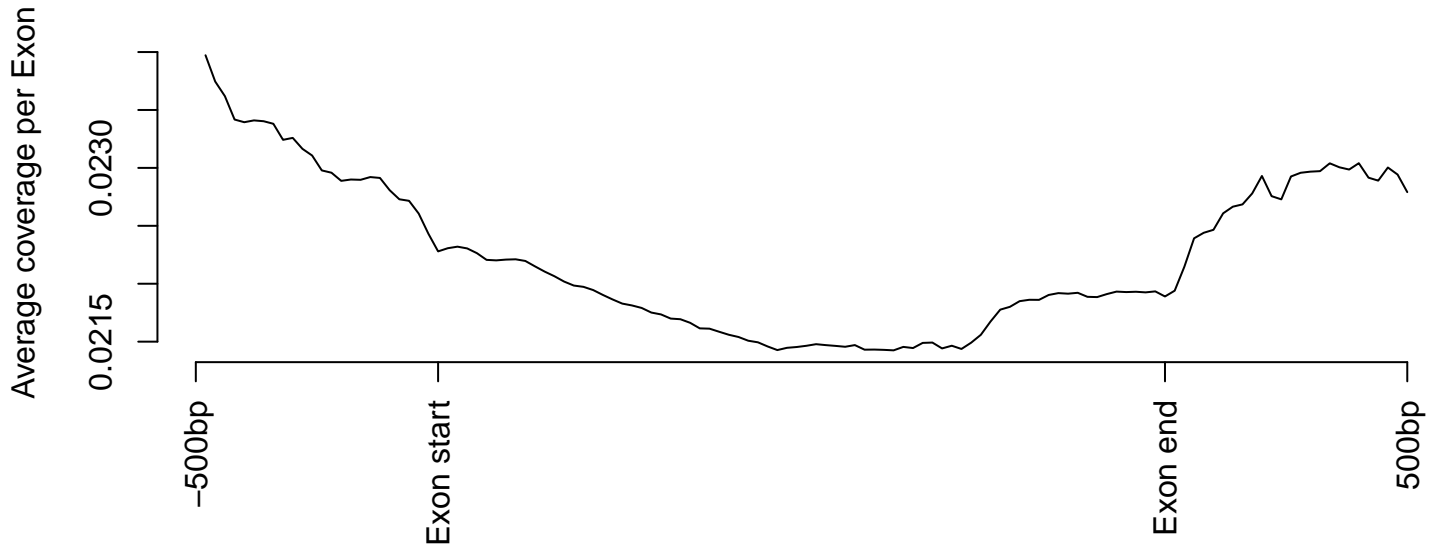
Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc740



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc741

Average coverage per Exon

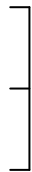
0.0028 0.0030

-500bp

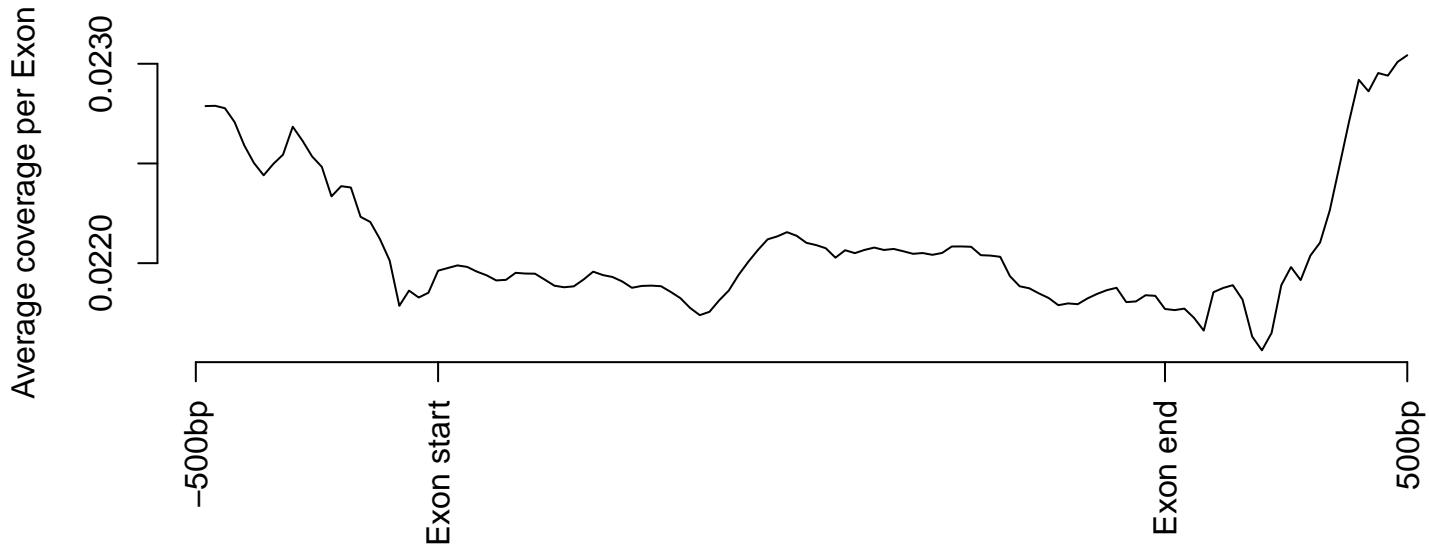
Exon start

Exon end

500bp

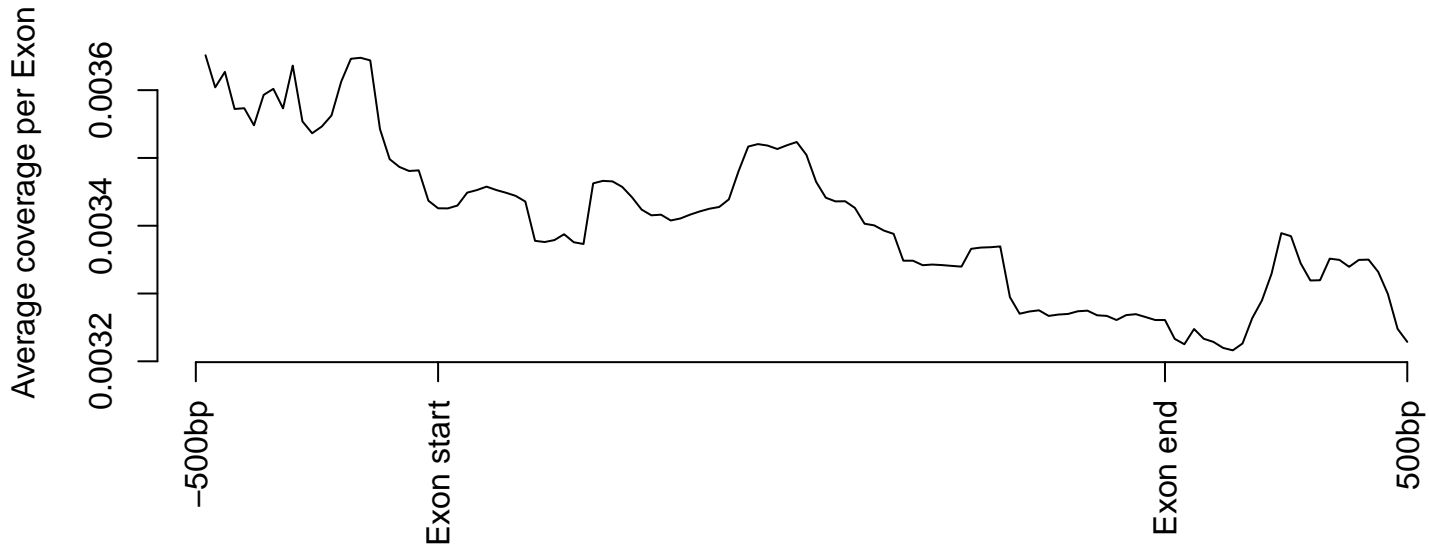


# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc742





# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc743



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc744

Average coverage per Exon

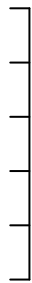
0.0035 0.0038

-500bp

Exon start

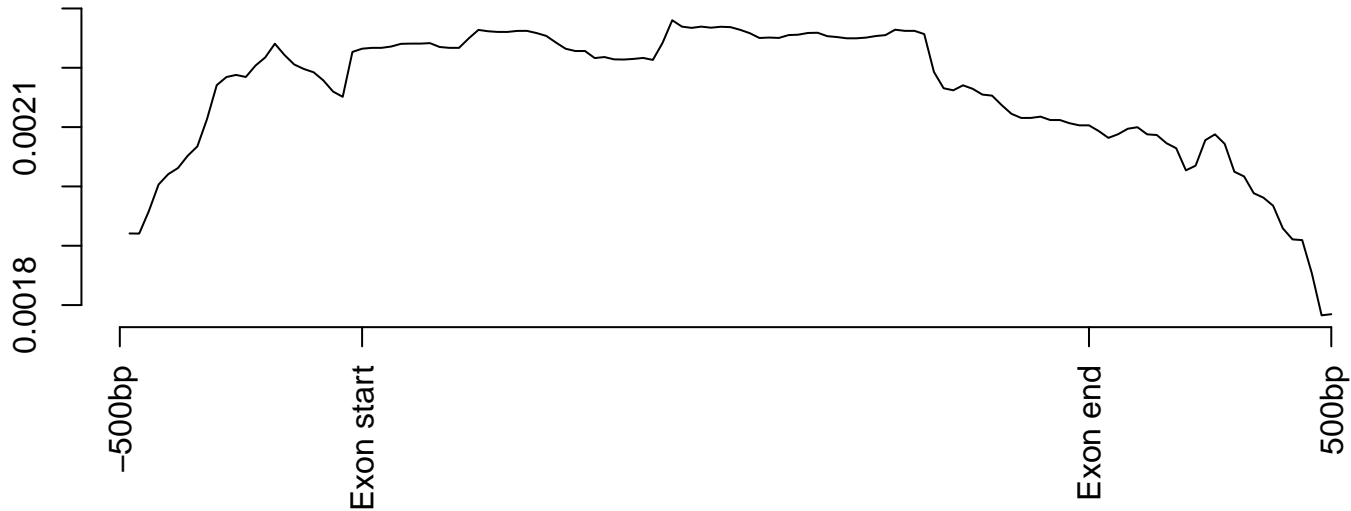
Exon end

500bp

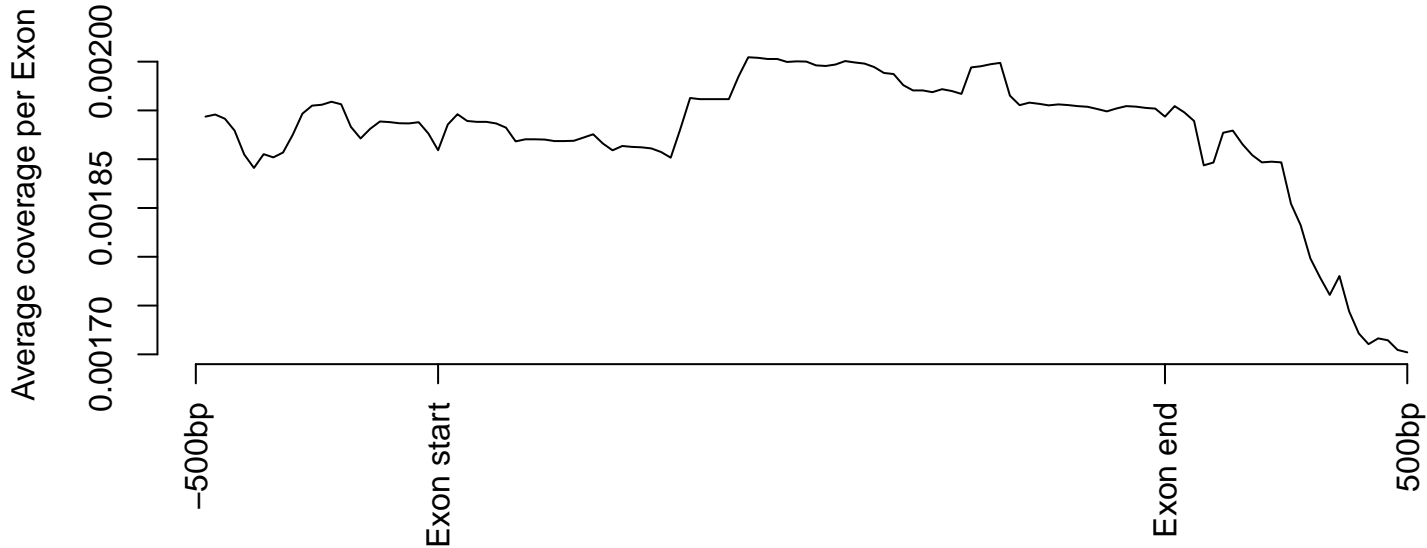


# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc745

Average coverage per Exon



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc746



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc747

Average coverage per Exon

0.0016 0.0018

-500bp

Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc748

Average coverage per Exon

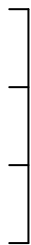
0.0013 0.0015

-500bp

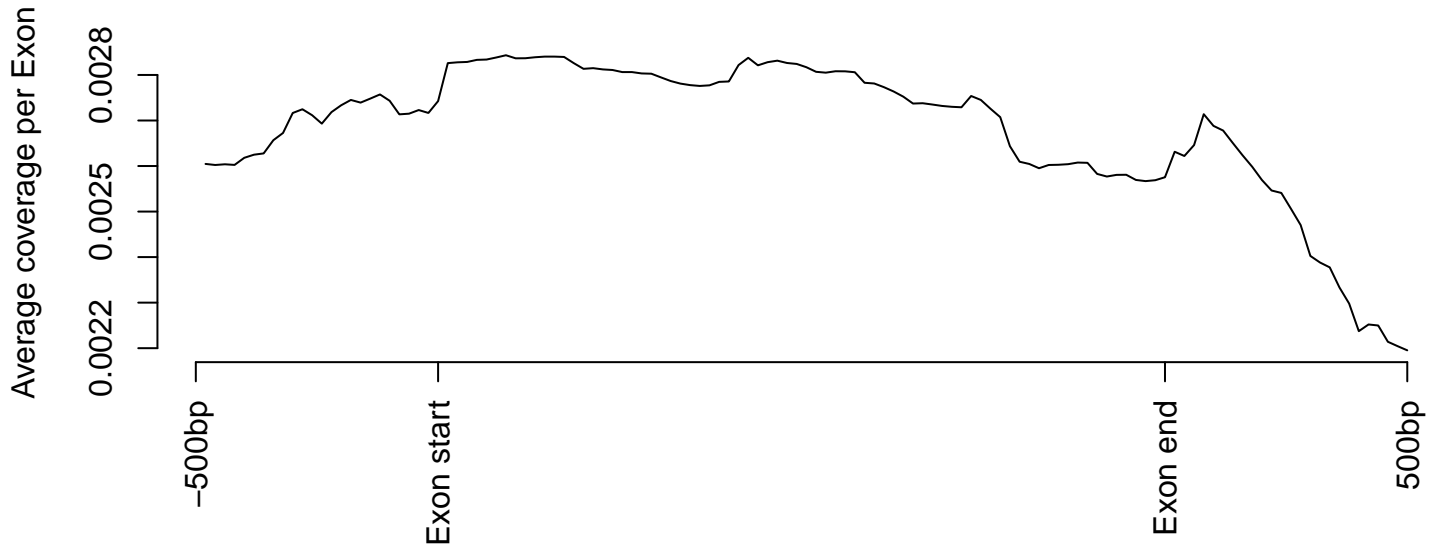
Exon start

Exon end

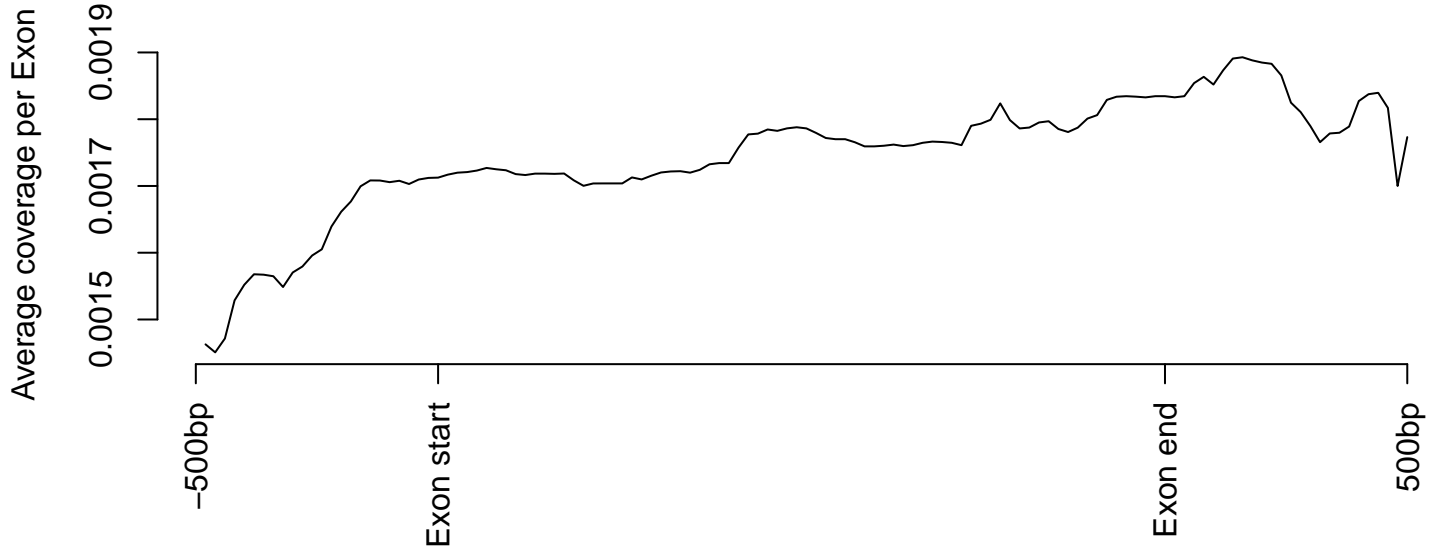
500bp



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc749



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc750





# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc751

Average coverage per Exon

0.0024

0.0021

-500bp

Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc752

Average coverage per Exon

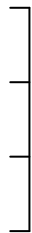
0.0060 0.0070

-500bp

Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc753

Average coverage per Exon

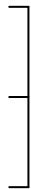
0.0010 0.0020

-500bp

Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc754

Average coverage per Exon

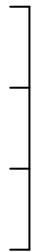
0.00080 0.00090

-500bp

Exon start

Exon end

500bp



# HumanInterneuronCulture PositiveMerged E.– HumanInterneuronCulturePositiveMerged

Average coverage per Exon

