

# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc735

Average coverage per Exon

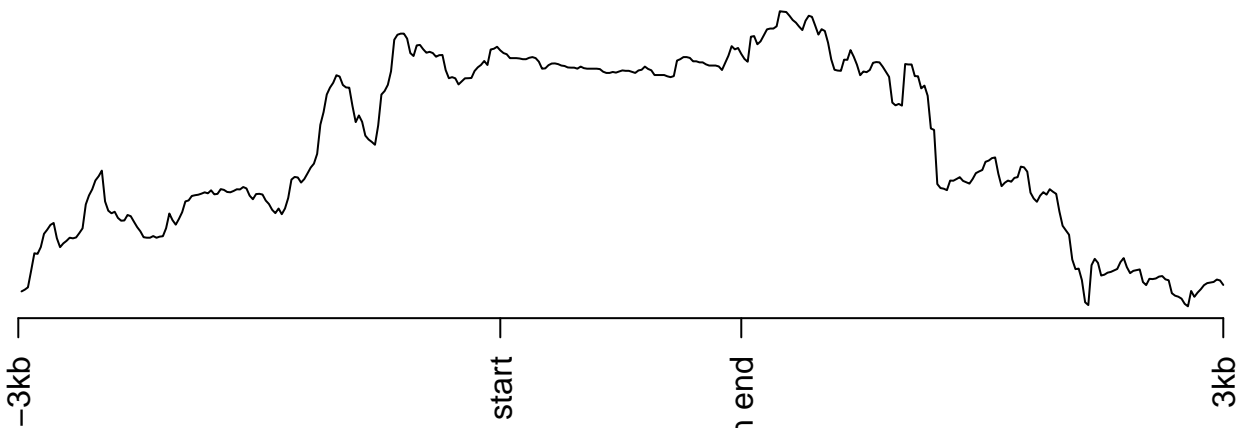
0.0010  
0.0007

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc736

Average coverage per Exon

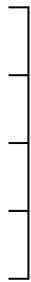
0.0004 0.0008 0.0012

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc737

Average coverage per Exon

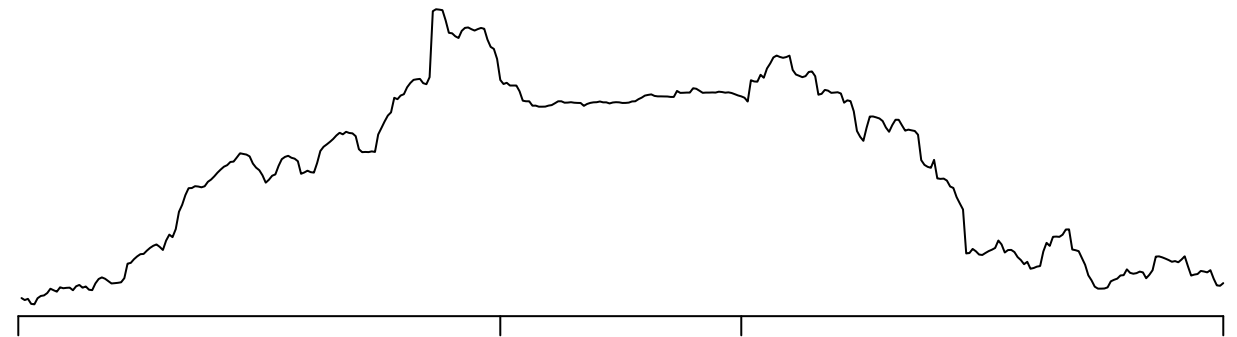
0.0016  
0.0010

-3kb

Exon start

Exon end

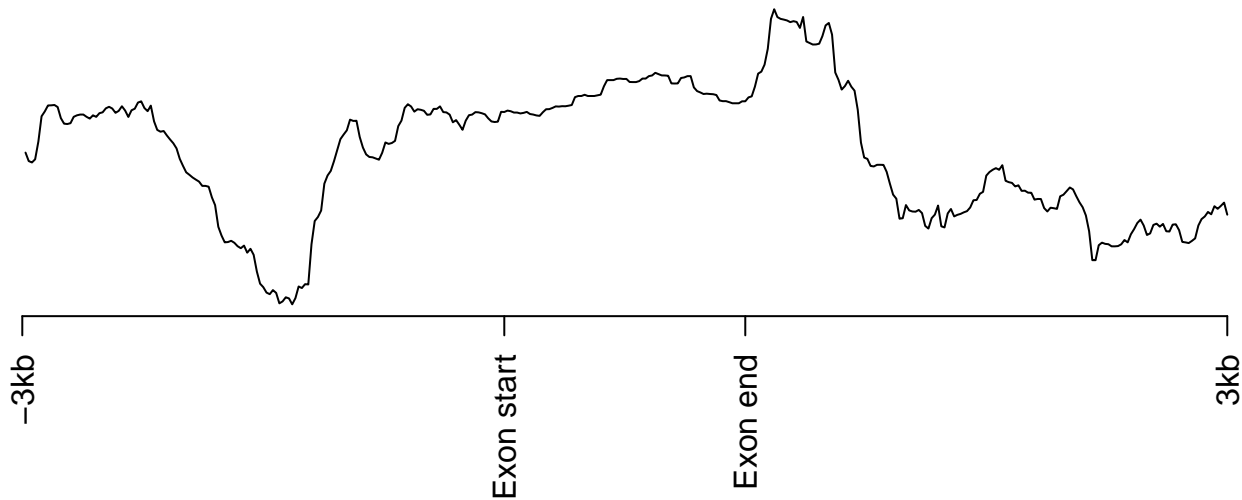
3kb



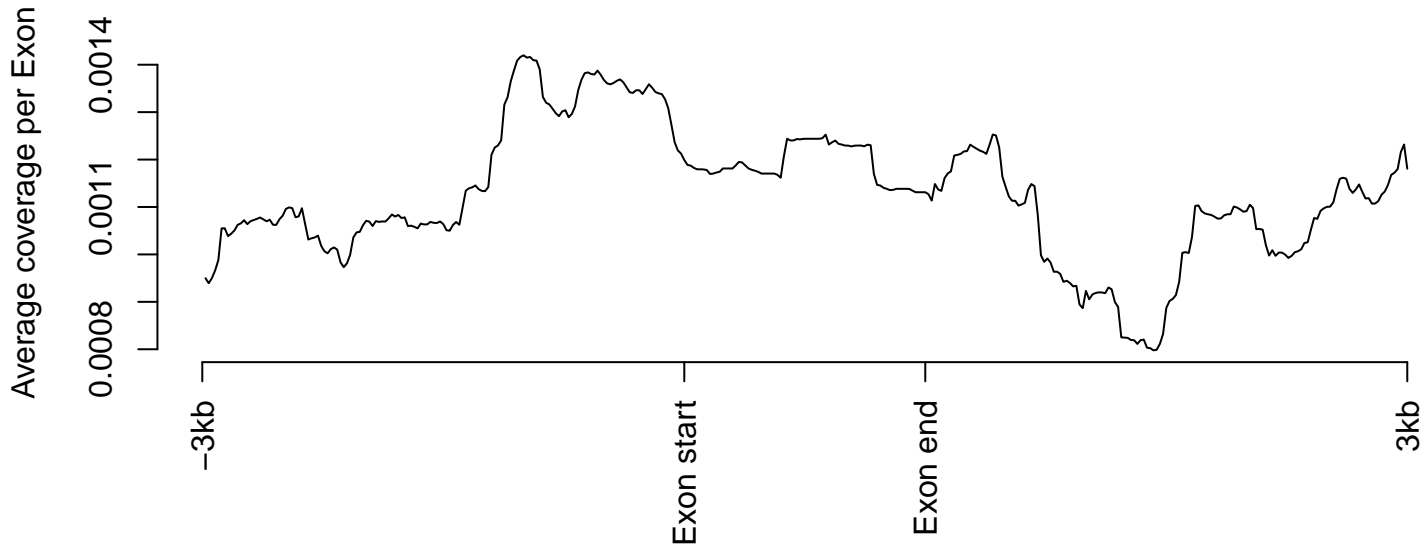
# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc738

Average coverage per Exon

0.0016  
0.0012



# HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc739



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc740

Average coverage per Exon

0.035

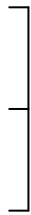
0.025

-3kb

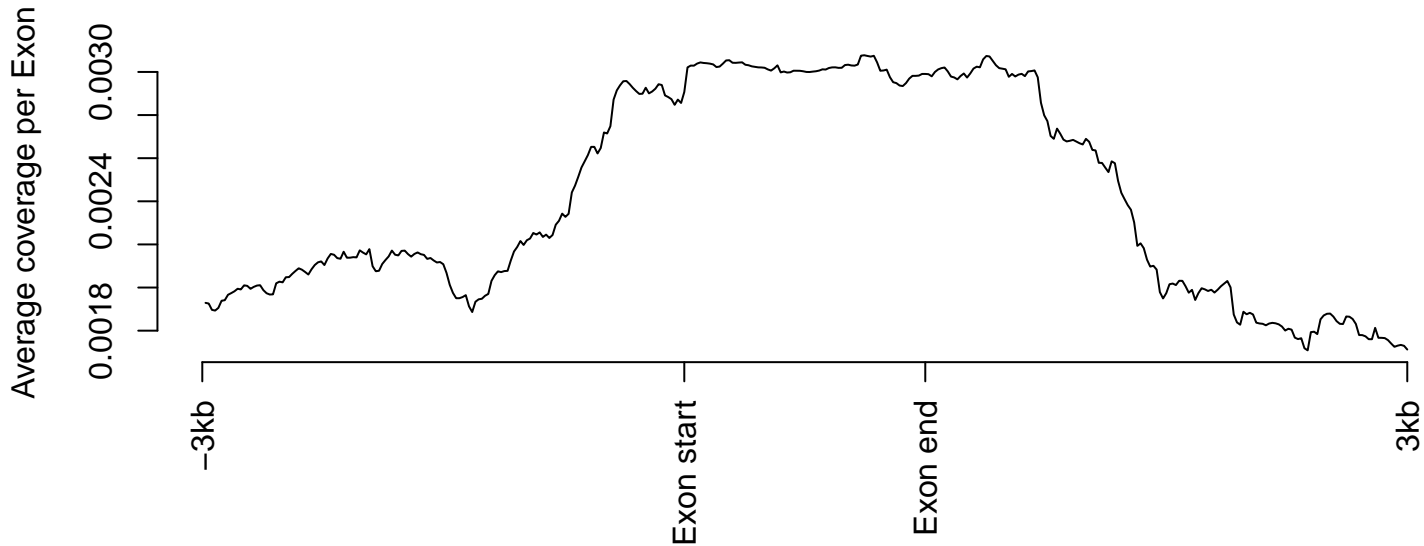
Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc741



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc742

Average coverage per Exon

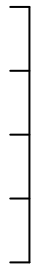
0.030  
0.026  
0.022

-3kb

Exon start

Exon end

3kb





# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc743

Average coverage per Exon

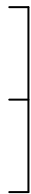
0.0025 0.0035

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc744

Average coverage per Exon

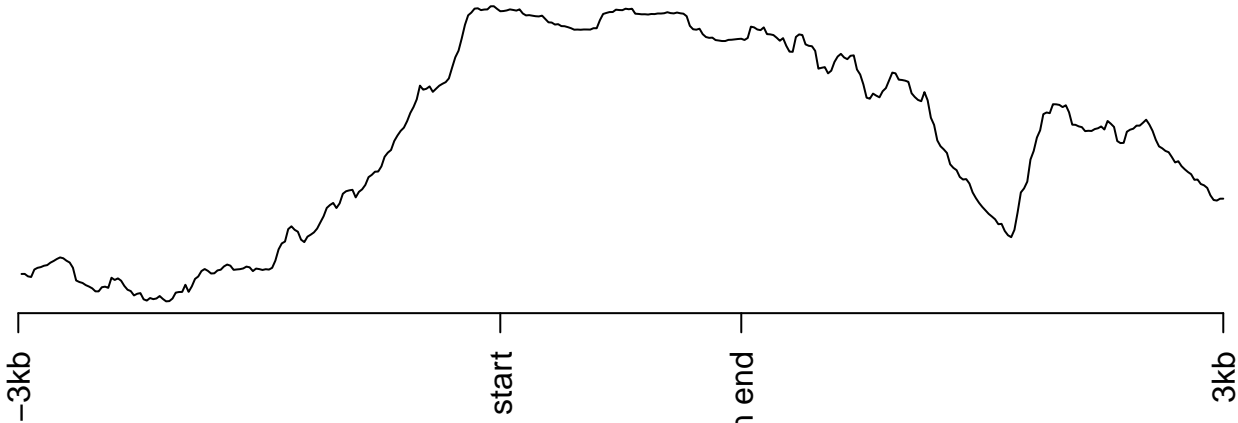
0.0025 0.0035

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc745

Average coverage per Exon

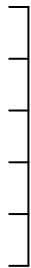
0.0018  
0.0012

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc746

Average coverage per Exon

0.0010 0.0016

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc747

Average coverage per Exon

0.0012 0.0016

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc748

Average coverage per Exon

0.0006 0.0012

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc749

Average coverage per Exon

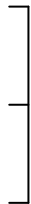
0.0015 0.0025

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc750

Average coverage per Exon

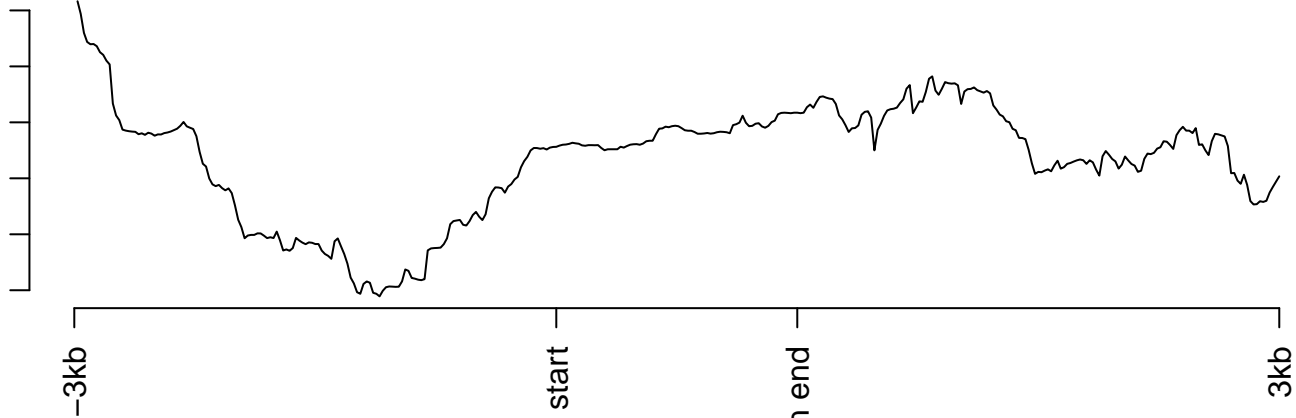
0.0018  
0.0012

-3kb

Exon start

Exon end

3kb





# HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc751

Average coverage per Exon

0.0025 0.0035

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc752

Average coverage per Exon

0.006 0.008 0.010

-3kb

Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc753

Average coverage per Exon

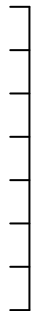
0.0005 0.0020 0.0035

-3kb

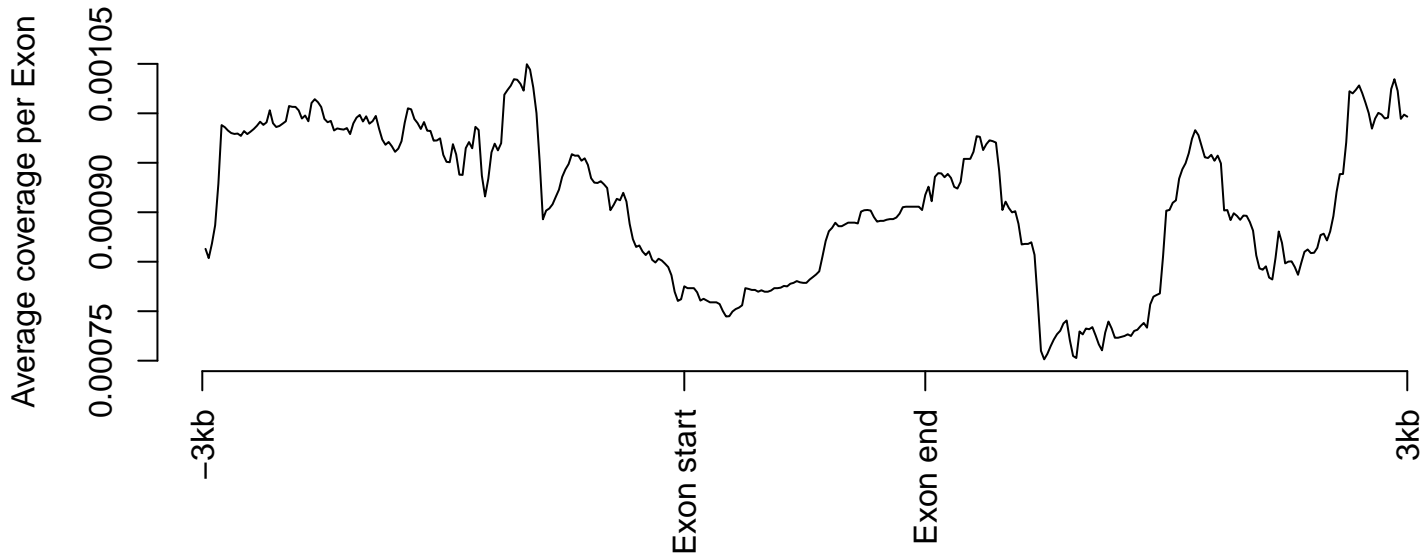
Exon start

Exon end

3kb



# HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc754



# HumanInterneuronCulture PositiveMerged E.– HumanInterneuronCulturePositiveMerged

Average coverage per Exon

0.095  
0.085

-3kb

Exon start

Exon end

3kb

