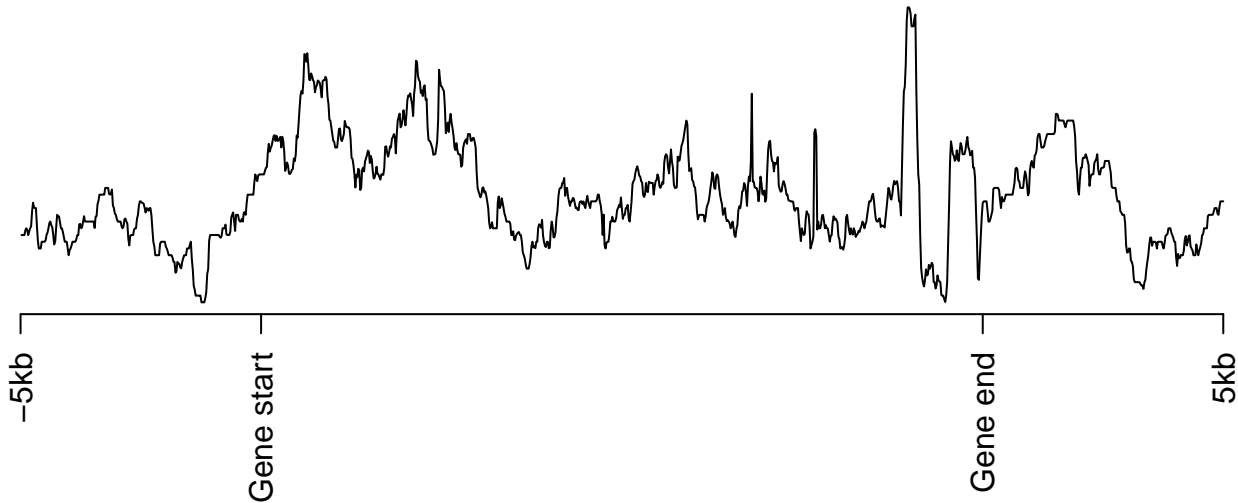


HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc735

Average coverage per gene

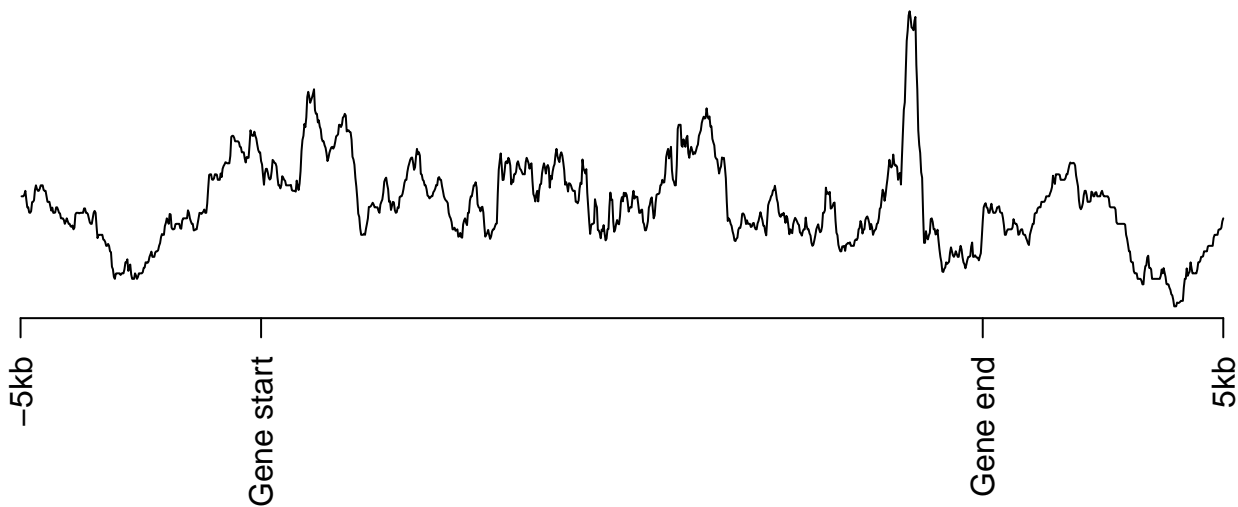
0.0006 0.0012



HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc736

Average coverage per gene

0.0006 0.0012 0.0018



HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc737

Average coverage per gene

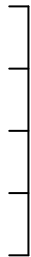
0.0014
0.0008

-5kb

Gene start

Gene end

5kb



HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc738

Average coverage per gene

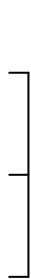
0.0020
0.0010

-5kb

Gene start

Gene end

5kb



HumanInterneuronCulture PositiveSingle E.658 scCLTdegenNuc739

Average coverage per gene

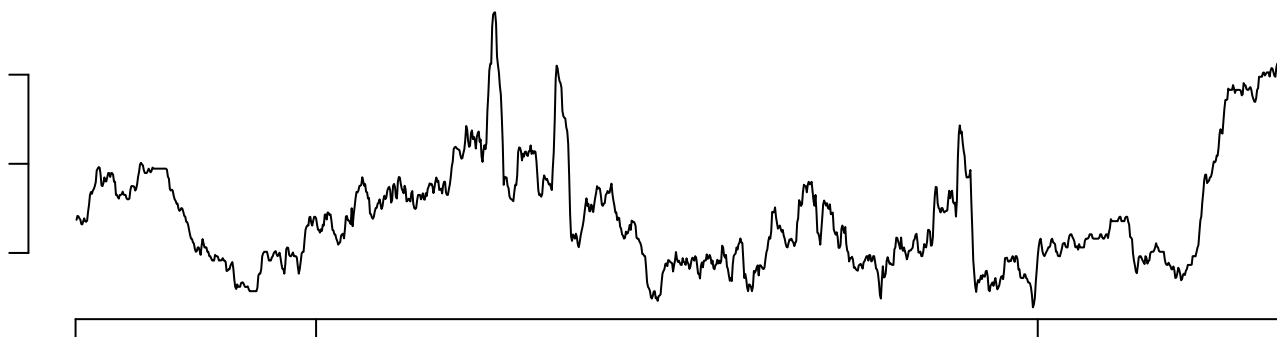
0.0020
0.0010

-5kb

Gene start

Gene end

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc740

Average coverage per gene

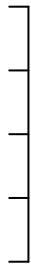
0.035 0.045 0.055

-5kb

Gene start

Gene end

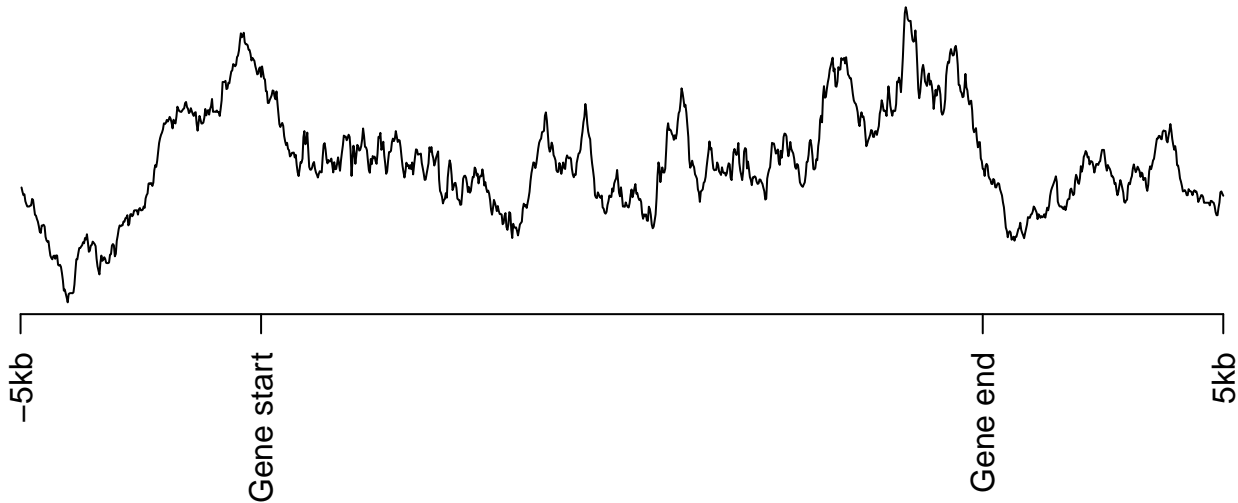
5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc741

Average coverage per gene

0.0015 0.0025



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc742

Average coverage per gene

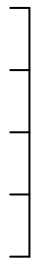
0.08
0.06
0.04

-5kb

Gene start

Gene end

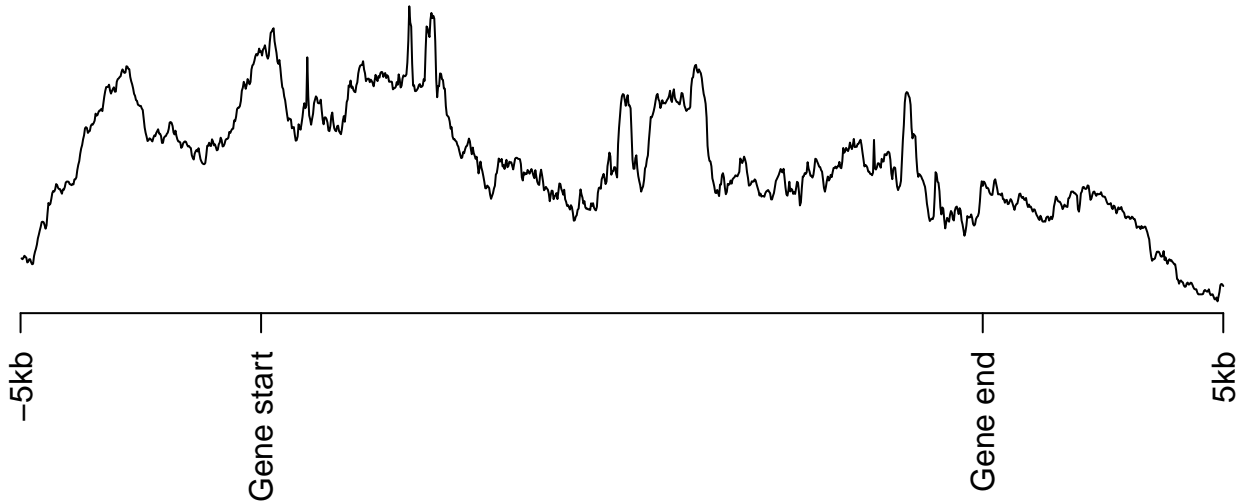
5kb



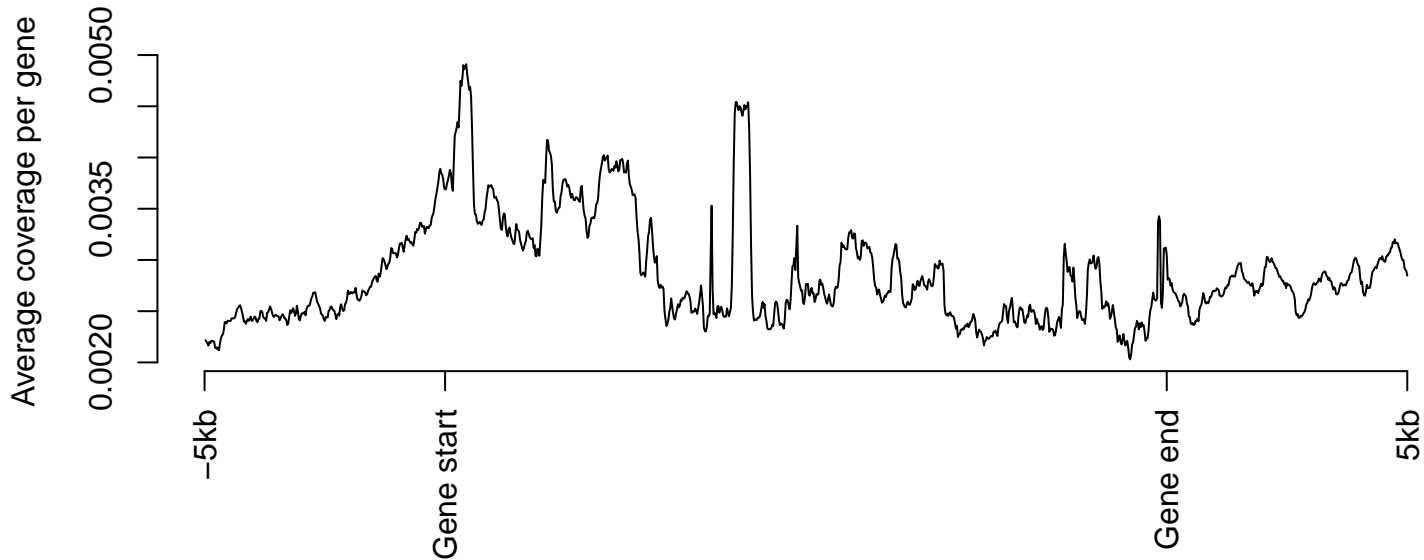
HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc743

Average coverage per gene

0.0030
0.0015



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc744



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc745

Average coverage per gene

0.0020
0.0010

-5kb

Gene start

Gene end

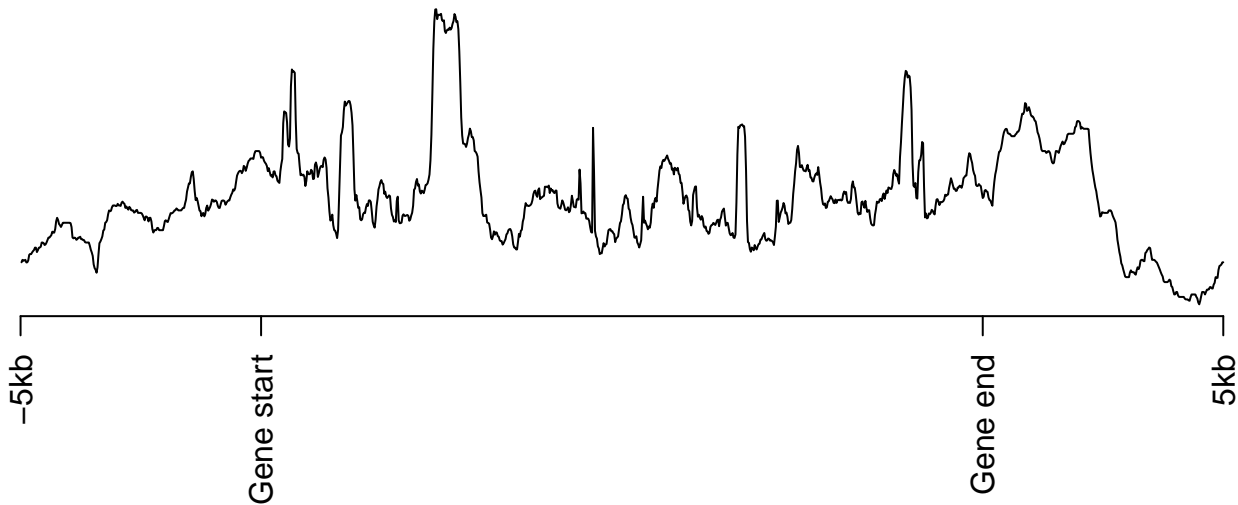
5kb



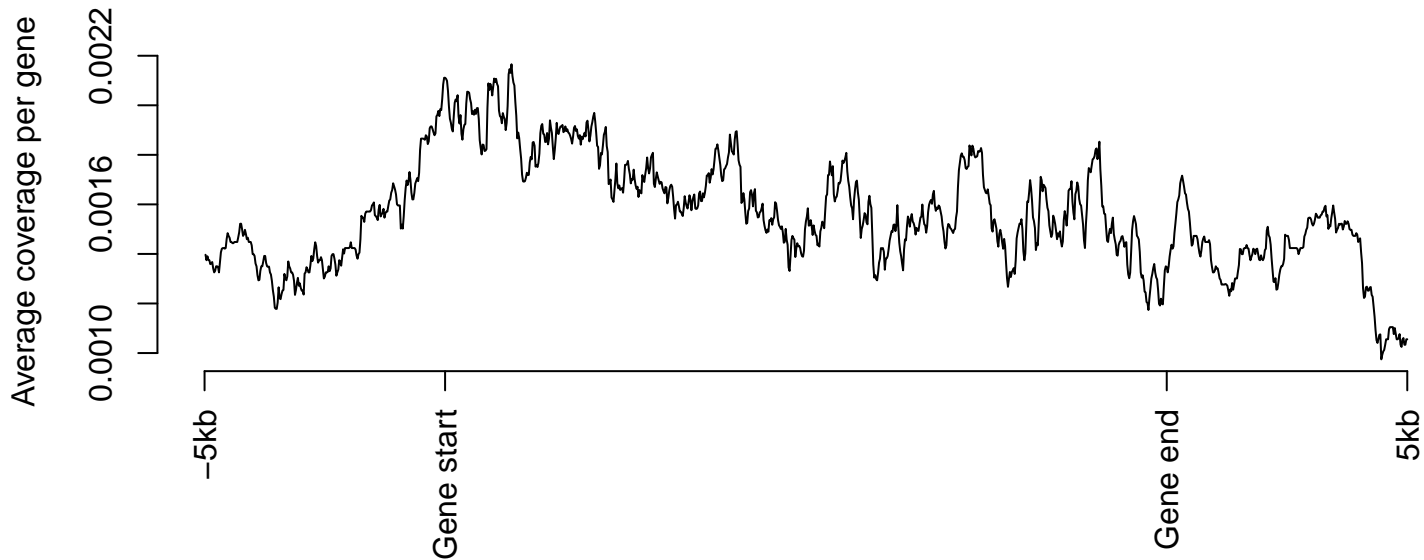
HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc746

Average coverage per gene

0.0010 0.0025



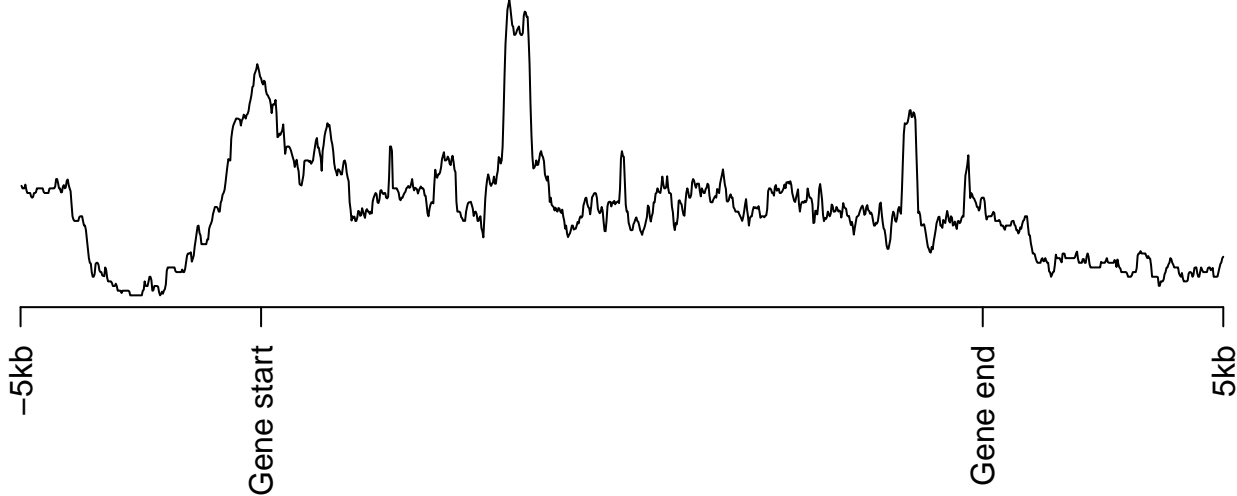
HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc747



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc748

Average coverage per gene

0.0015
0.0005



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc749

Average coverage per gene

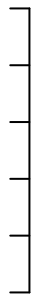
0.0010 0.0025

-5kb

Gene start

Gene end

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc750

Average coverage per gene

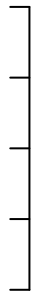
0.0030
0.0020
0.0010

-5kb

Gene start

Gene end

5kb



HumanInterneuronCulture PositiveSingle E.663 scCLTdegenNuc751

Average coverage per gene

0.0035
0.0025

-5kb

Gene start

Gene end

5kb



HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc752

Average coverage per gene

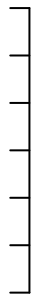
0.012
0.009
0.006

-5kb

Gene start

Gene end

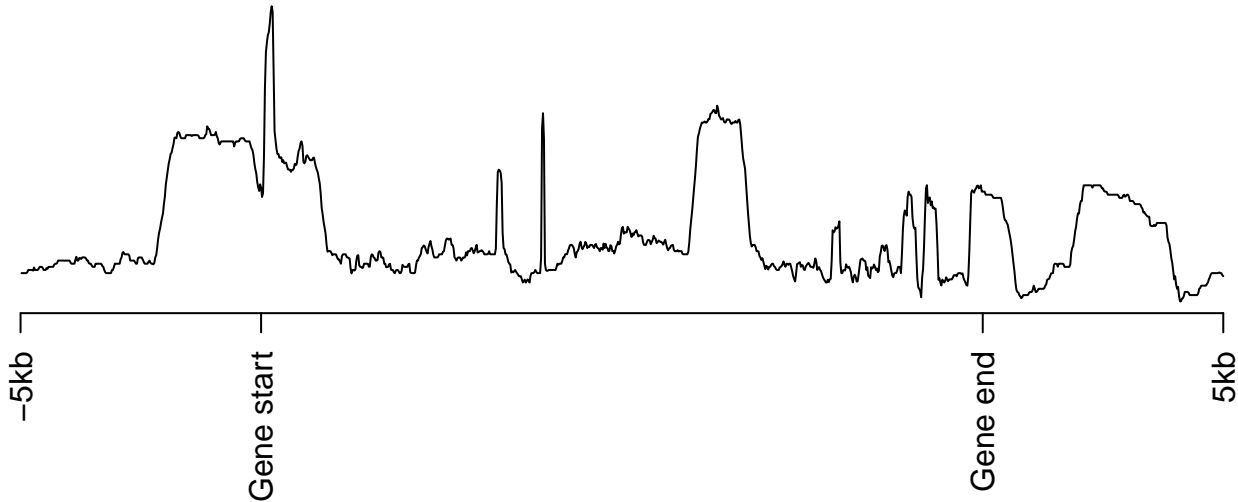
5kb



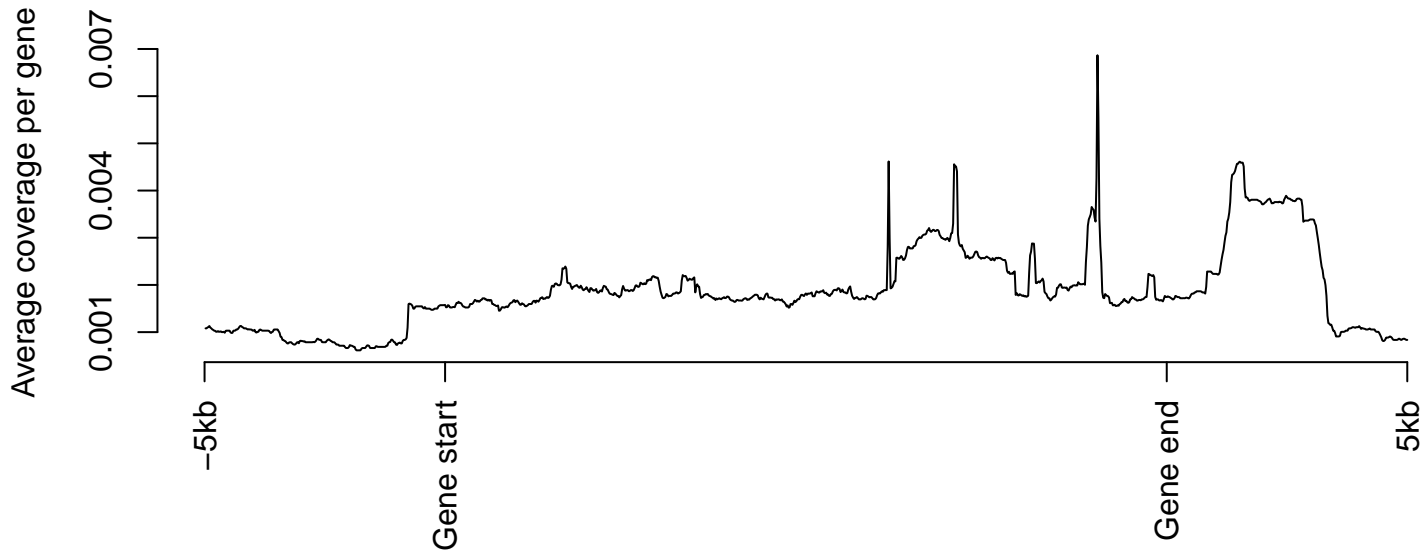
HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc753

Average coverage per gene

0.0025
0.0015
0.0005



HumanInterneuronCulture PositiveSingle E.664 scCLTdegenNuc754



HumanInterneuronCulture PositiveMerged E. – HumanInterneuronCulturePositiveMerged

