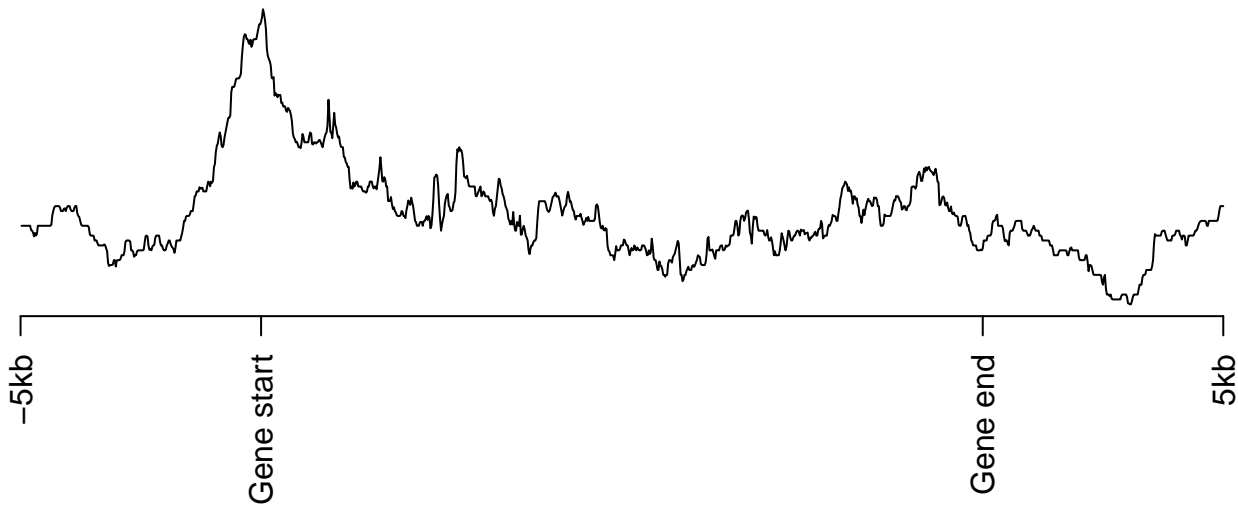


# MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc603

Average coverage per gene

0.0015  
0.0005



# MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc604

Average coverage per gene

0.0010  
0.0004

-5kb

Gene start

Gene end

5kb



# MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc605

Average coverage per gene

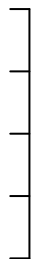
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

5kb



# MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc606

Average coverage per gene

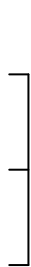
0.0015  
0.0005

-5kb

Gene start

Gene end

5kb



# MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc607

Average coverage per gene

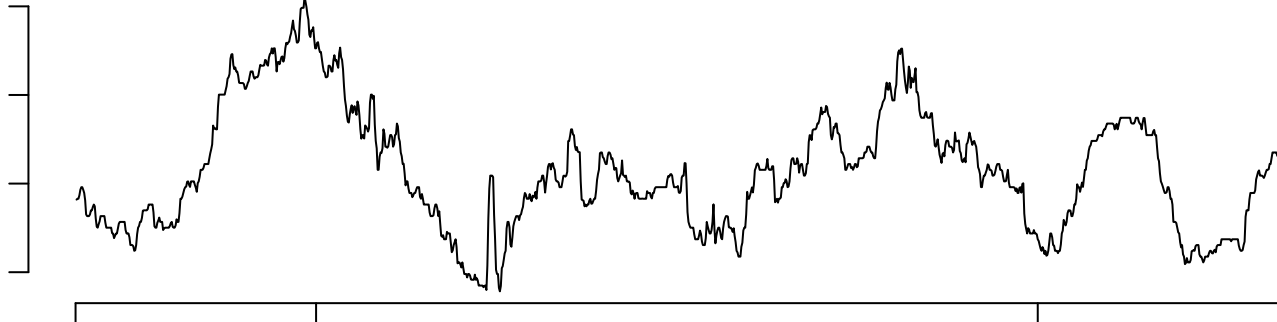
0.0015  
0.0005

-5kb

Gene start

Gene end

5kb



# MouseInterneuronSlice PositiveSingle E.642 scCLTdegenNuc608b

Average coverage per gene

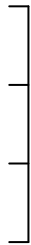
2e-04  
6e-04

-5kb

Gene start

Gene end

5kb



# MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc621

Average coverage per gene

0.0020  
0.0010

-5kb

Gene start

Gene end

5kb



# MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc622

Average coverage per gene

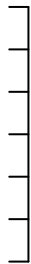
0.013  
0.010  
0.007

-5kb

Gene start

Gene end

5kb

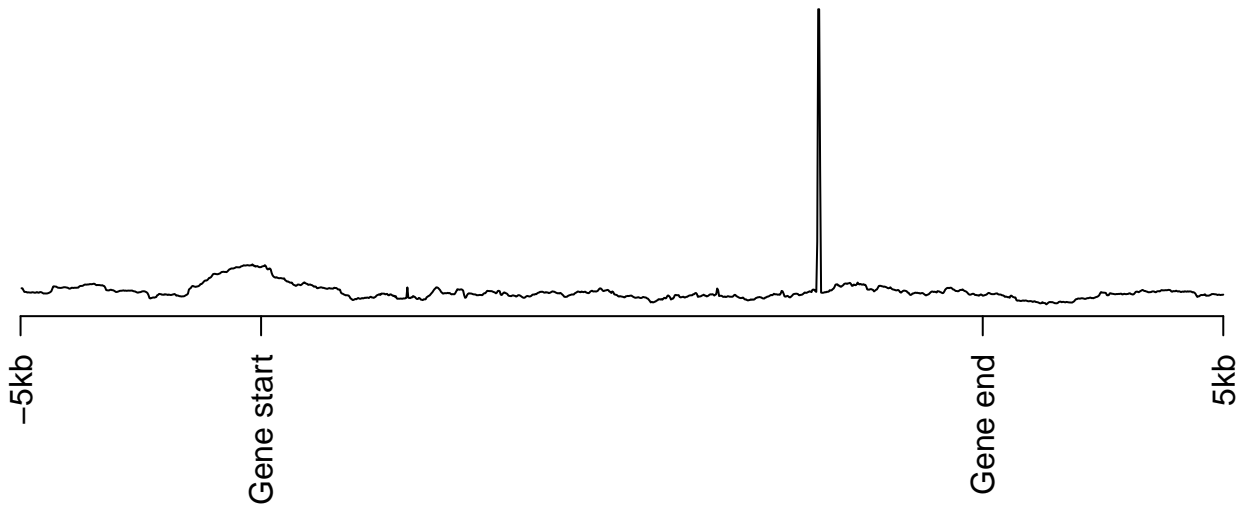




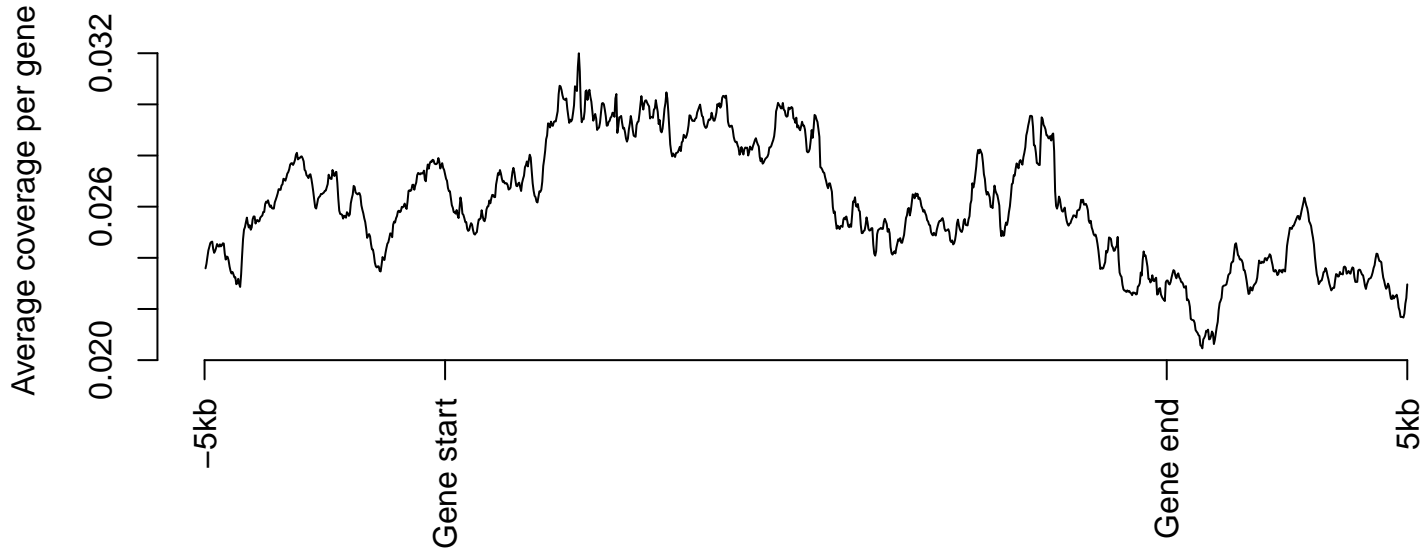
# MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc623

Average coverage per gene

0.01  
0.03



# MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc624



# MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc625

Average coverage per gene

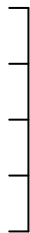
0.0014  
0.0008

-5kb

Gene start

Gene end

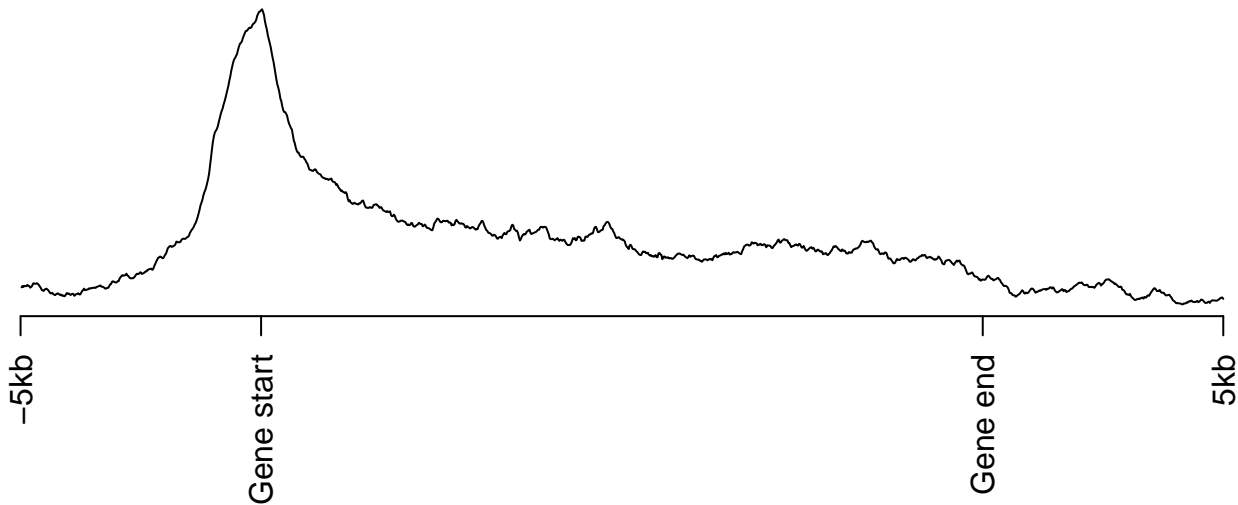
5kb



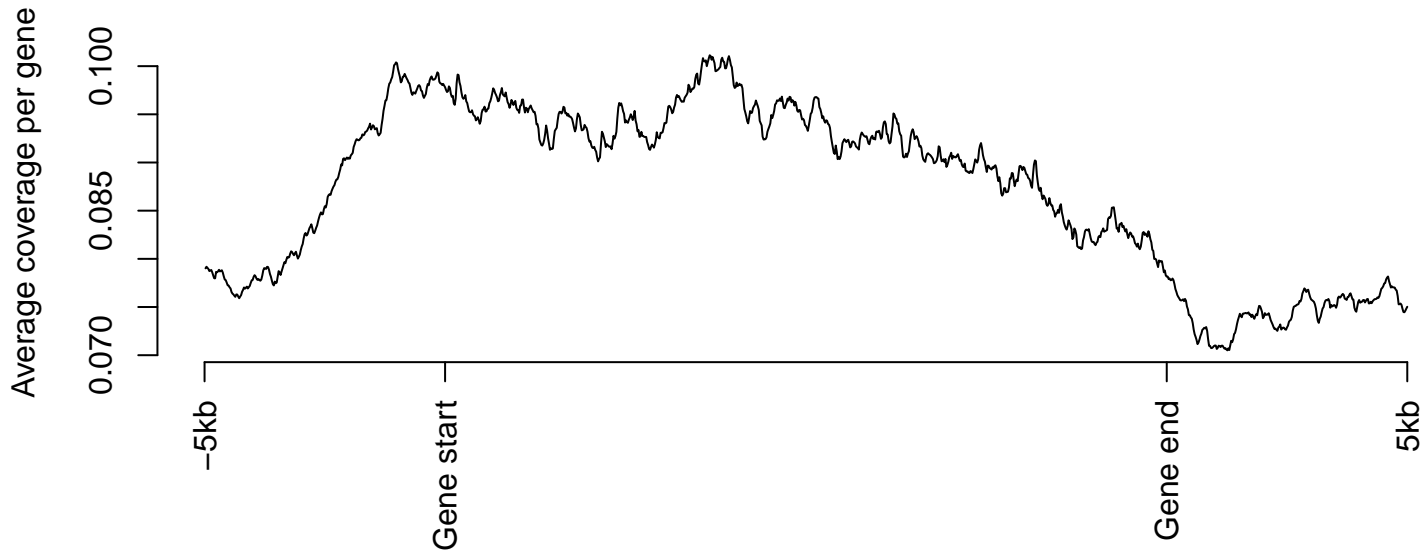
# MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc626

Average coverage per gene

0.18  
0.14  
0.10



# MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc627



# MouseInterneuronSlice PositiveSingle E.653 scCLTdegenNuc663

Average coverage per gene

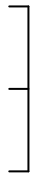
0.0020  
0.0010

-5kb

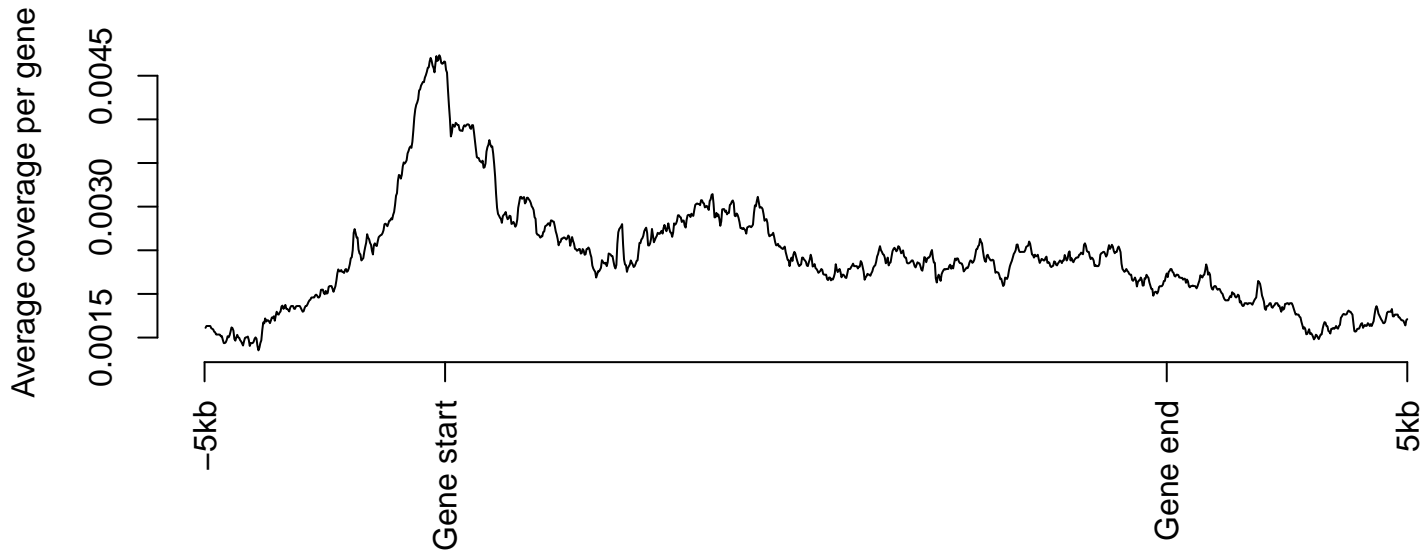
Gene start

Gene end

5kb



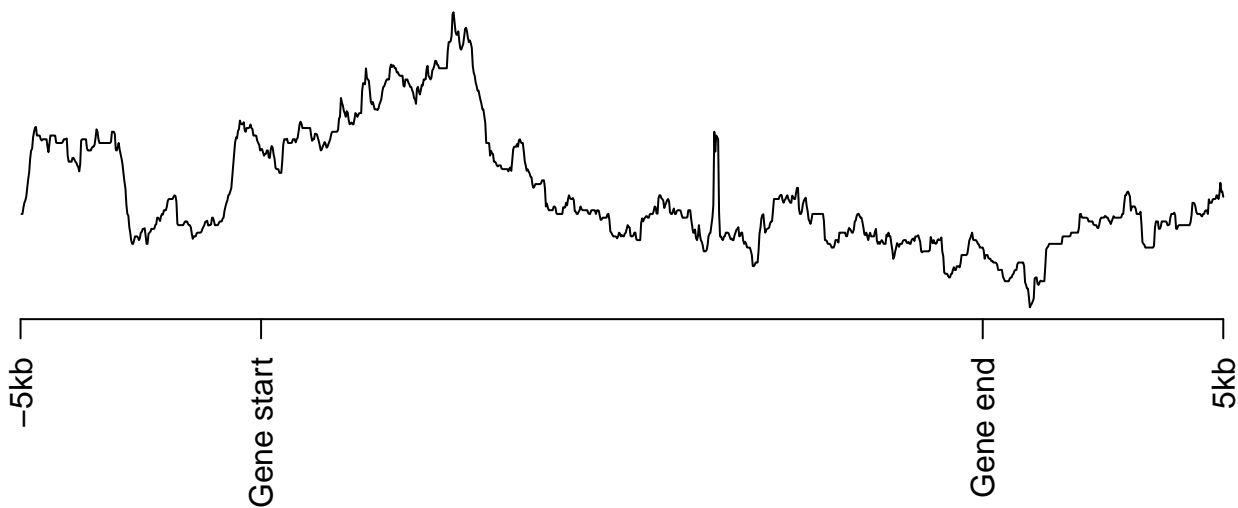
# MouseInterneuronSlice PositiveSingle E.653 scCLTdegenNuc664



# MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc697

Average coverage per gene

0.0020  
0.0005

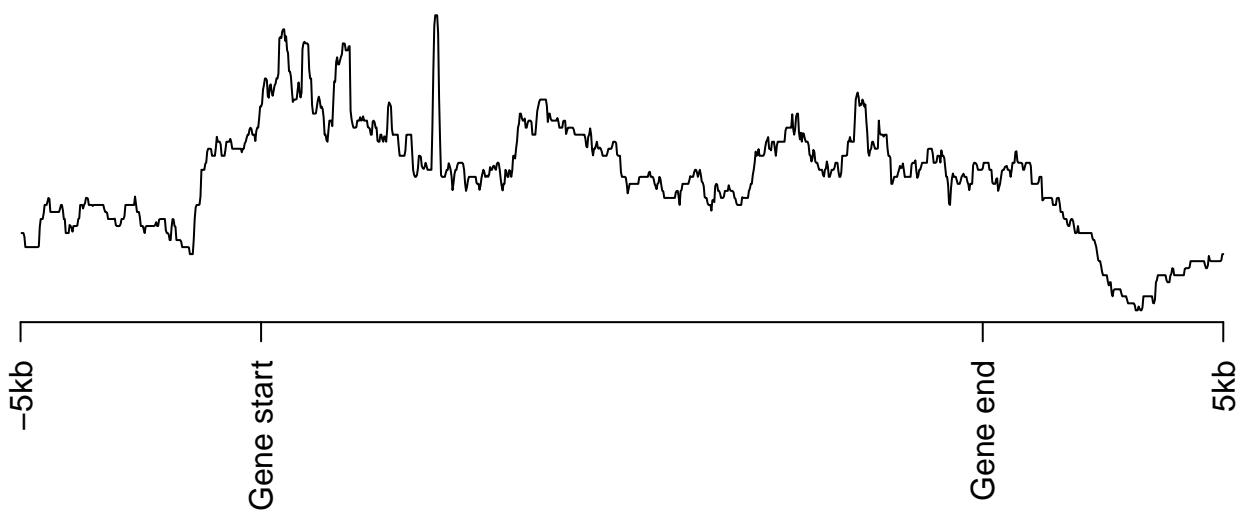




# MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc698

Average coverage per gene

0.0002 0.0008 0.0014



# MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc699

Average coverage per gene

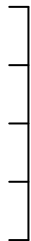
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

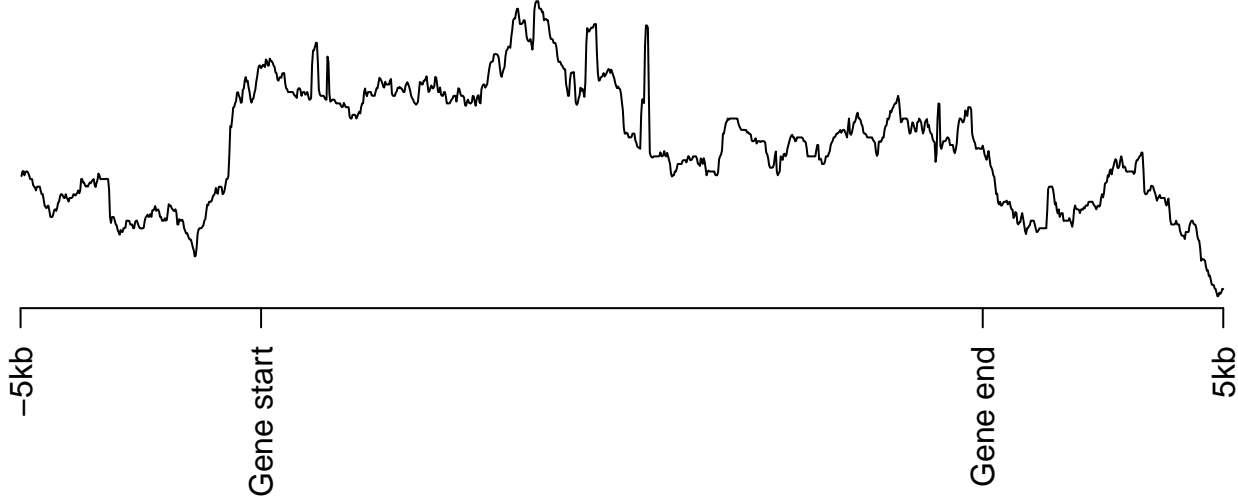
5kb



# MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc776

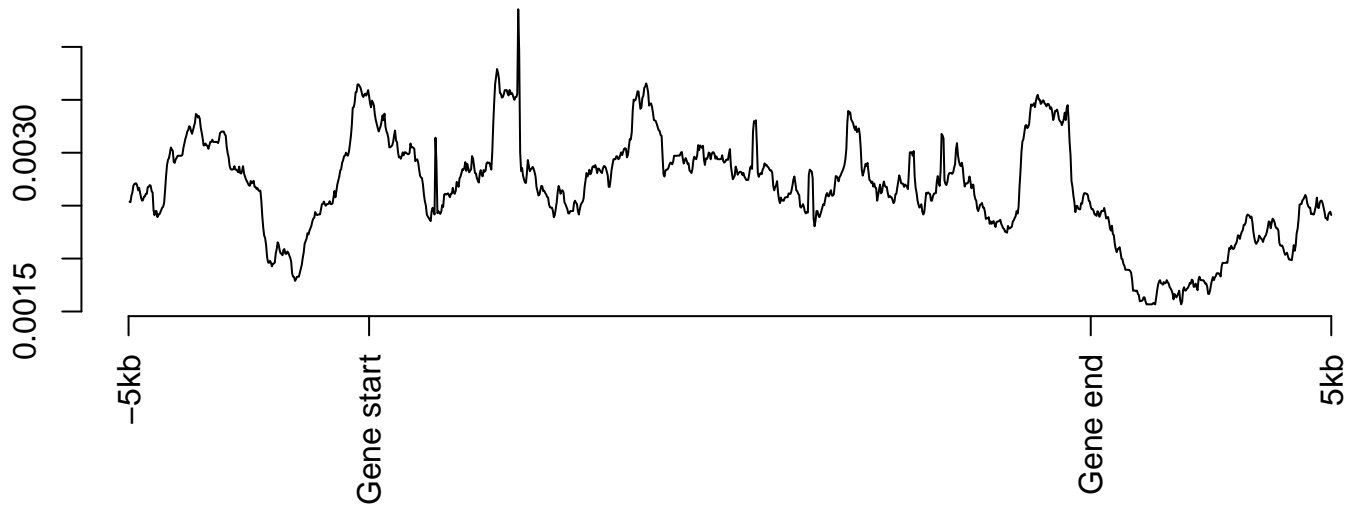
Average coverage per gene

0.0010 0.0025



# MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc777

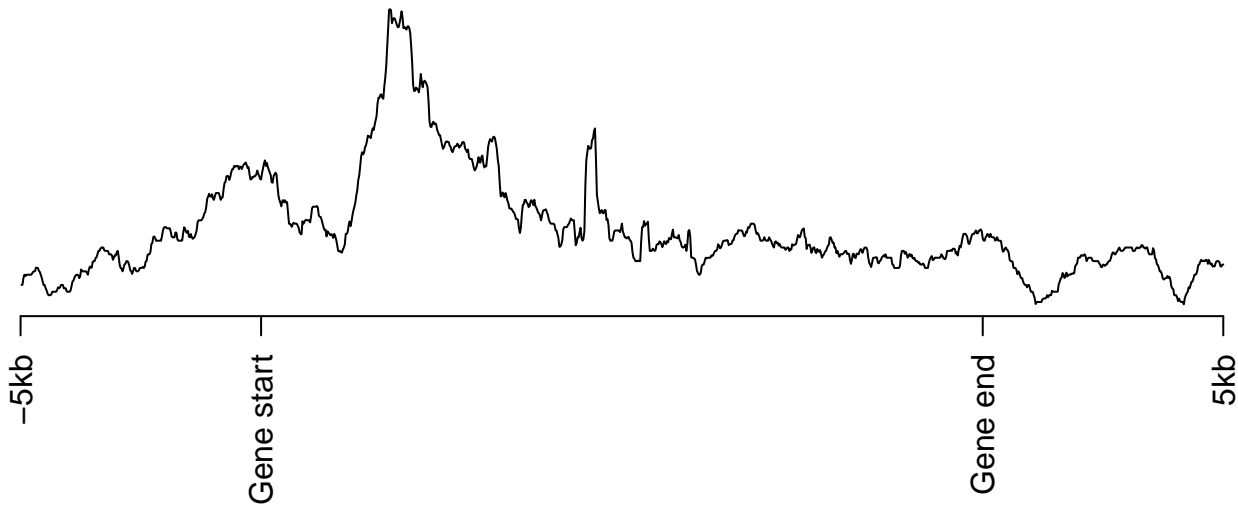
Average coverage per gene



# MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc778

Average coverage per gene

0.0010 0.0025



# MouseInterneuronSlice PositiveMerged E.– MouseInterneuronSlicePositiveMerged

Average coverage per gene

0.25 0.30 0.35

-5kb

Gene start

Gene end

5kb

