

# MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc429

Average coverage per gene

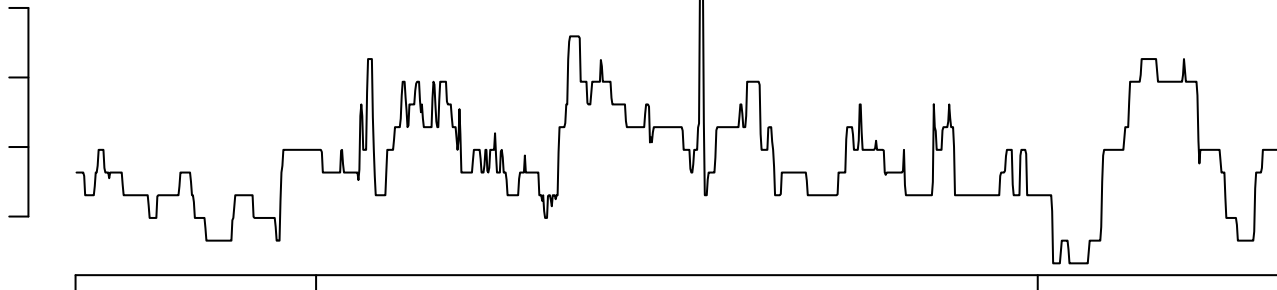
1e-04 3e-04

-5kb

Gene start

Gene end

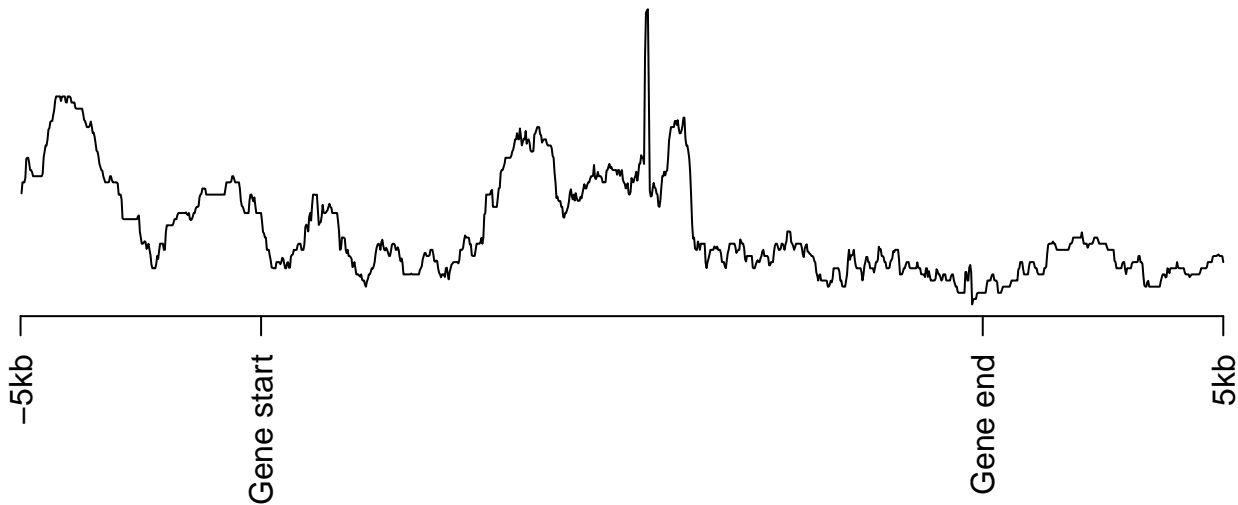
5kb



# MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc430

Average coverage per gene

0.0015  
0.0005



# MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc431

Average coverage per gene

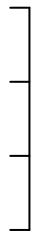
0.0020 0.0030

-5kb

Gene start

Gene end

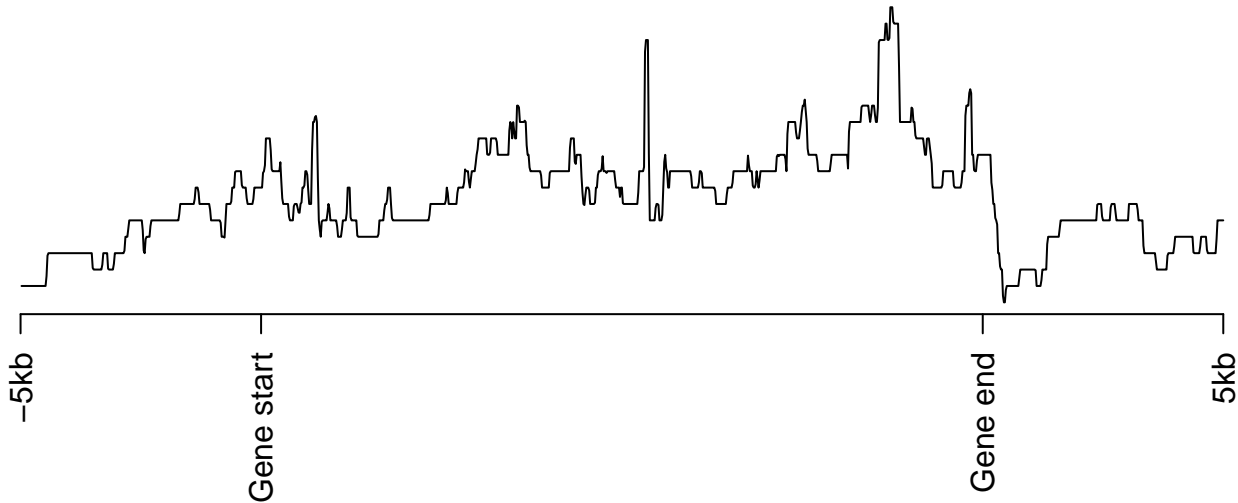
5kb



# MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc433

Average coverage per gene

1e-04  
4e-04



# MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc886

Average coverage per gene

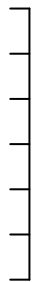
1e-04  
4e-04  
7e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc891

Average coverage per gene

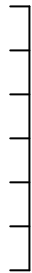
3e-04 6e-04 9e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc892

Average coverage per gene

2e-04

5e-04

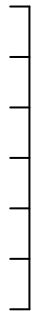
8e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture NegCtrlPooled E.597 scCLTdegenNuc471

Average coverage per gene

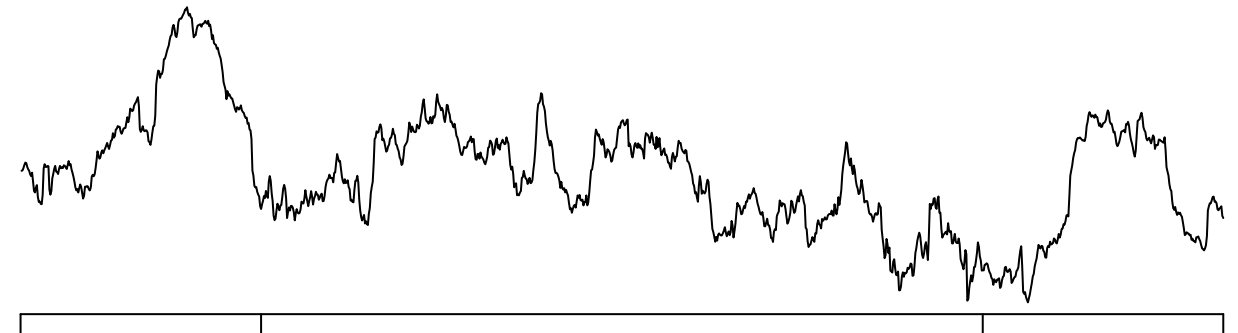
0.16 0.18 0.20

-5kb

Gene start

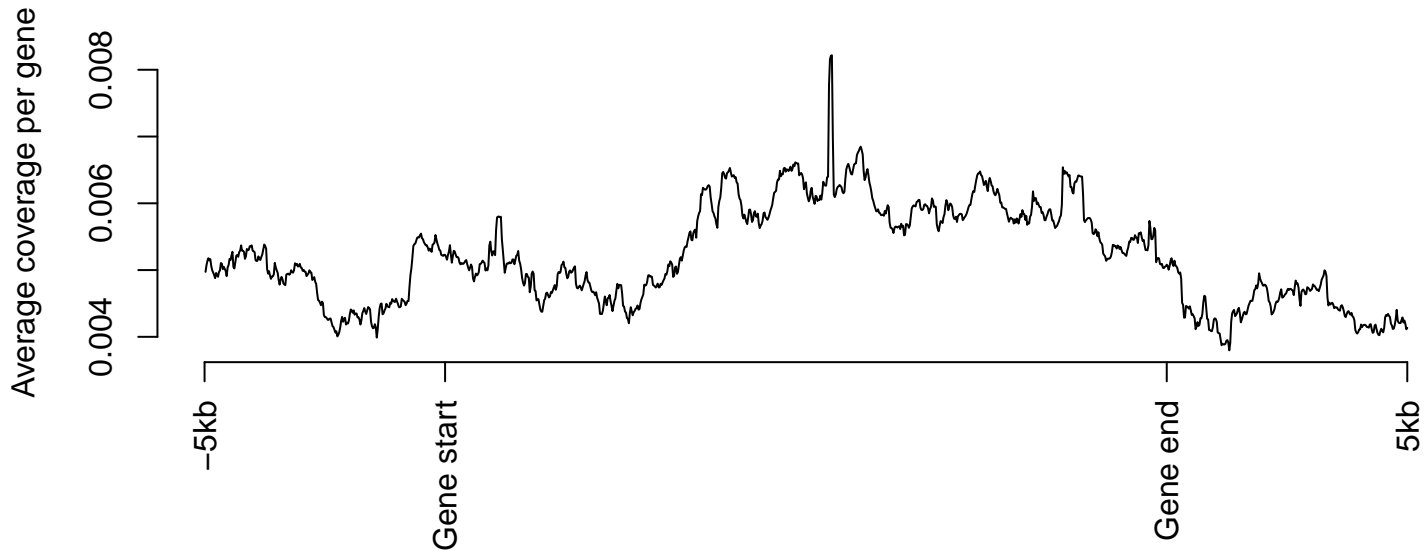
Gene end

5kb





# MouseNeuronCulture NegCtrlMerged E.– MouseNeuronCultureNegCtrlMerged



# MouseNeuronCulture NegCtrlAll E.– MouseNeuronCultureNegCtrlAll

Average coverage per gene

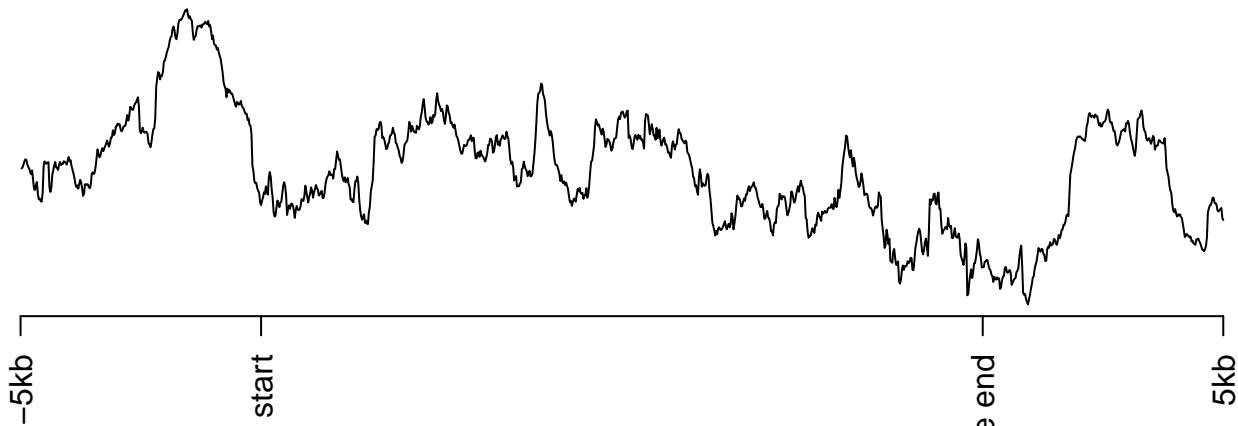
0.16 0.18 0.20

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc387

Average coverage per gene

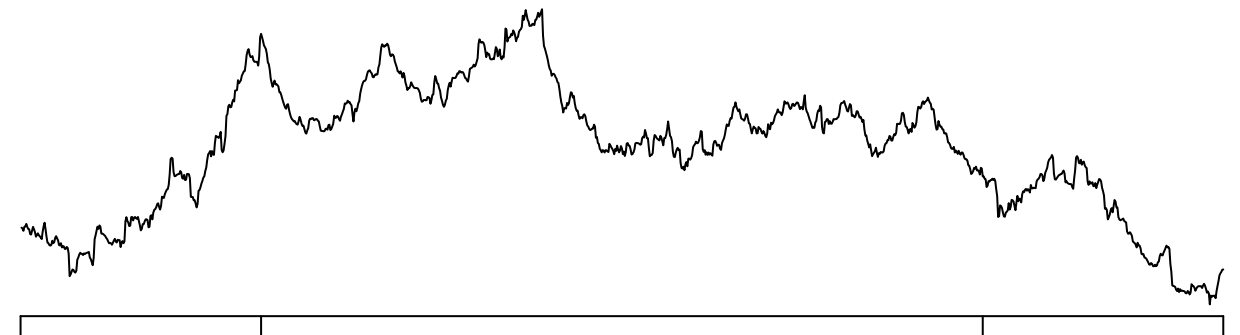
0.005  
0.003

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc388

Average coverage per gene

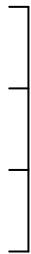
0.0030  
0.0020

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc390

Average coverage per gene

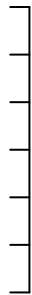
3e-04  
6e-04  
9e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc391

Average coverage per gene

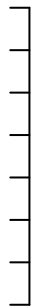
0.0030 0.0045 0.0060

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc392

Average coverage per gene

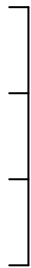
0.0020  
0.0030

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc393

Average coverage per gene

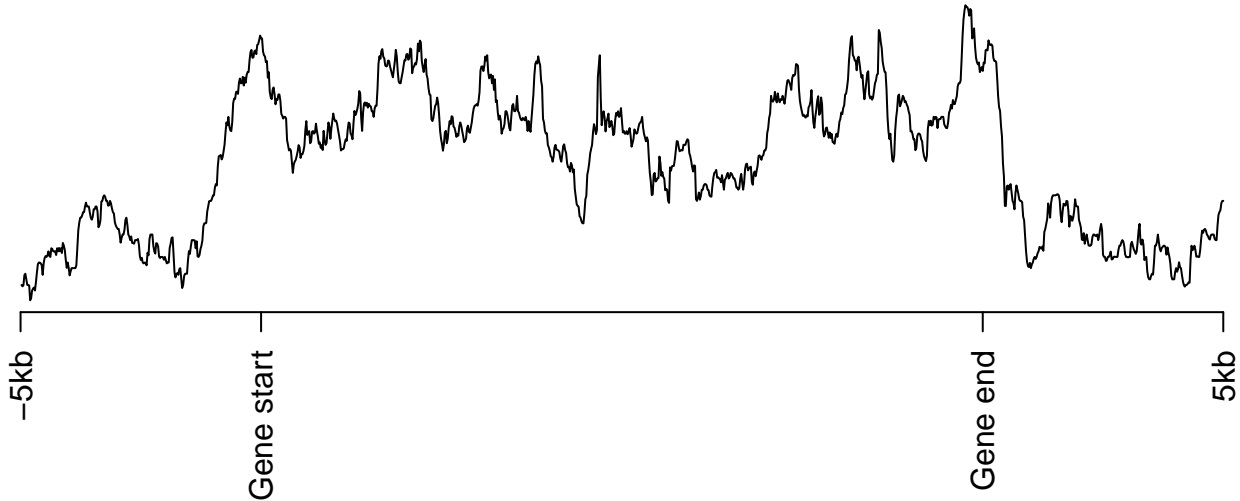
0.0030  
0.0020

-5kb

Gene start

Gene end

5kb





# MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc501

Average coverage per gene

4e-04  
8e-04

-5kb

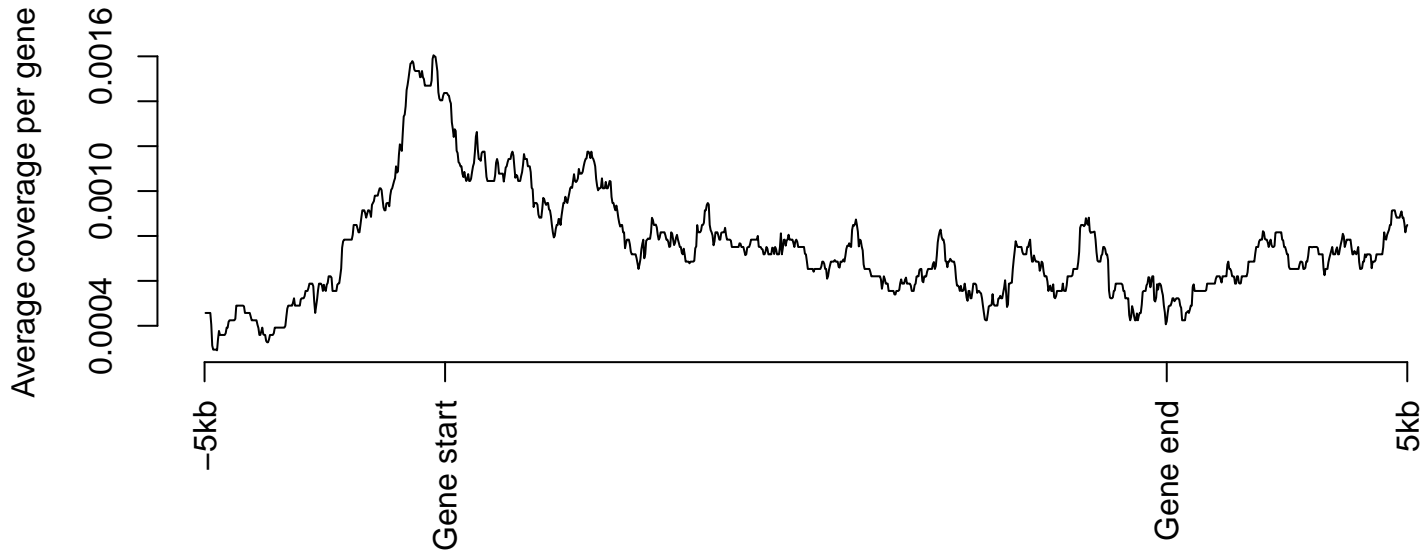
Gene start

Gene end

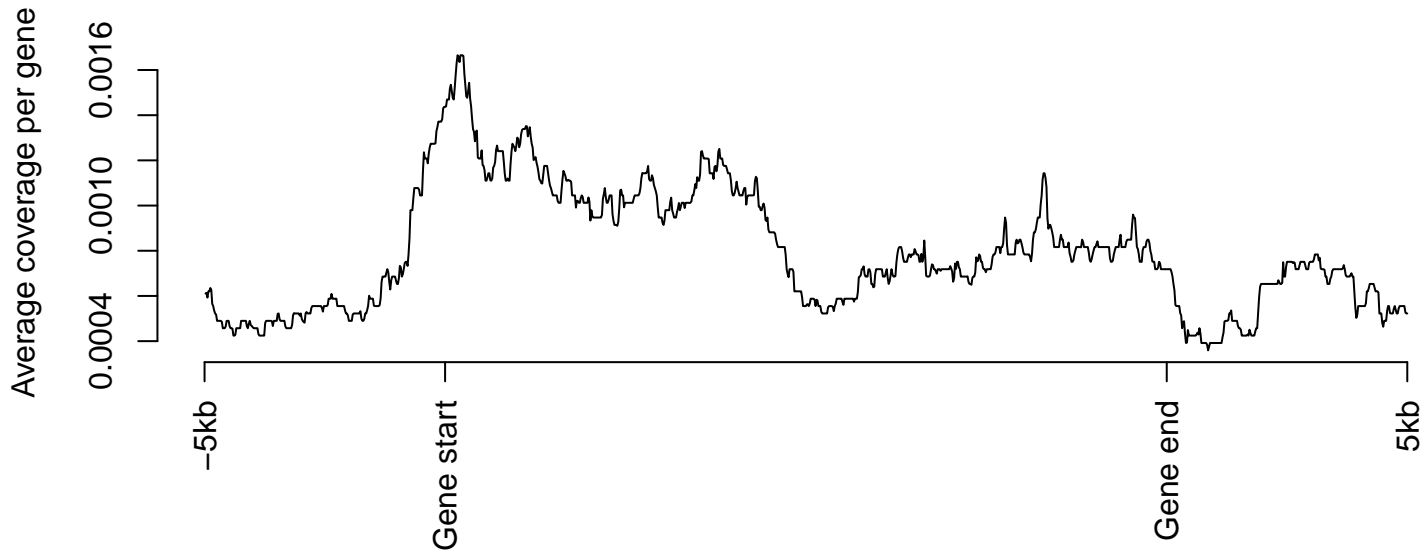
5kb



# MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc502

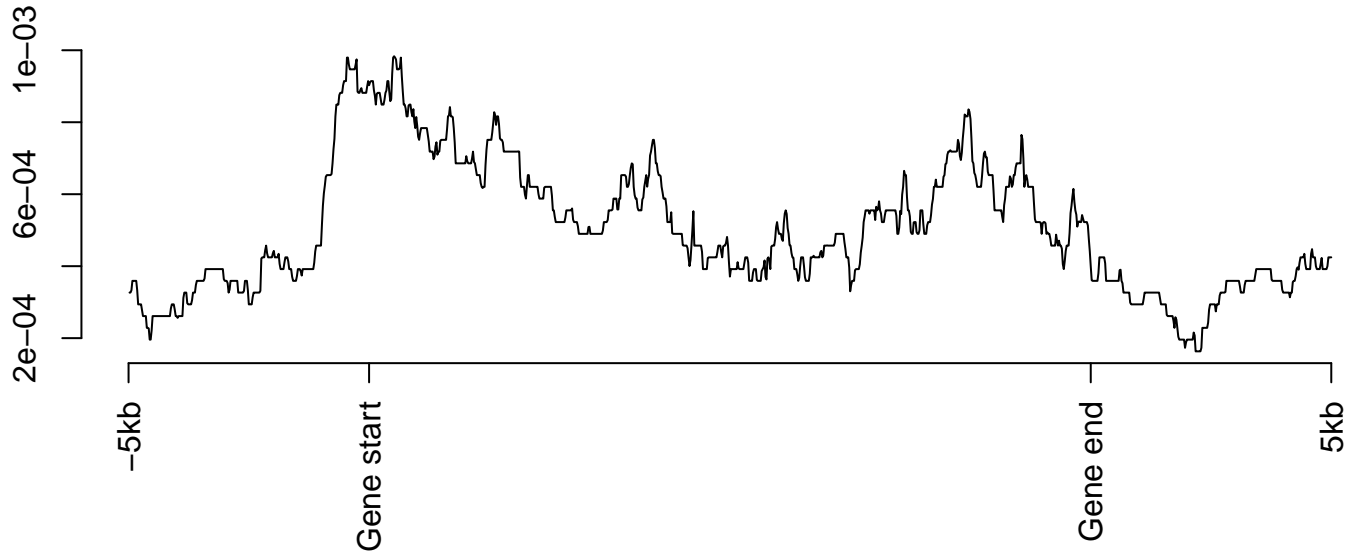


# MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc503



# MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc504

Average coverage per gene



# MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc505

Average coverage per gene

0.0015  
0.0005

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc510

Average coverage per gene

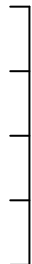
1e-04 3e-04 5e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc511

Average coverage per gene

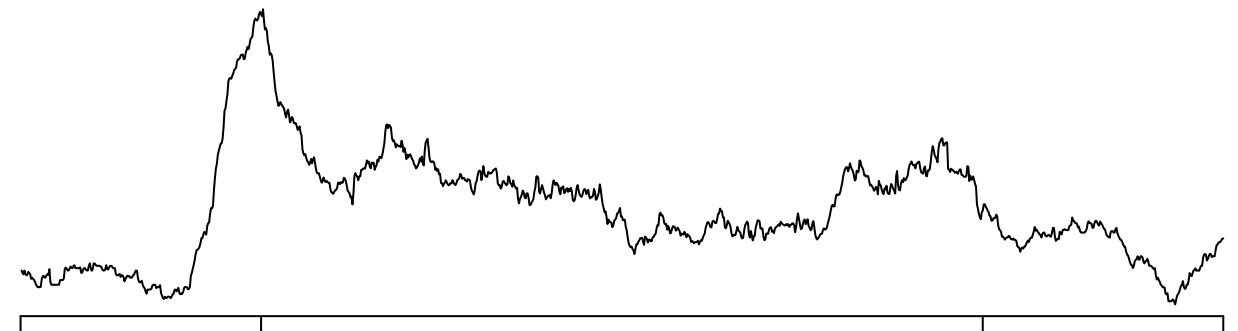
0.004  
0.002

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc512

Average coverage per gene

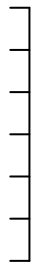
0.0015 0.0030 0.0045

-5kb

Gene start

Gene end

5kb

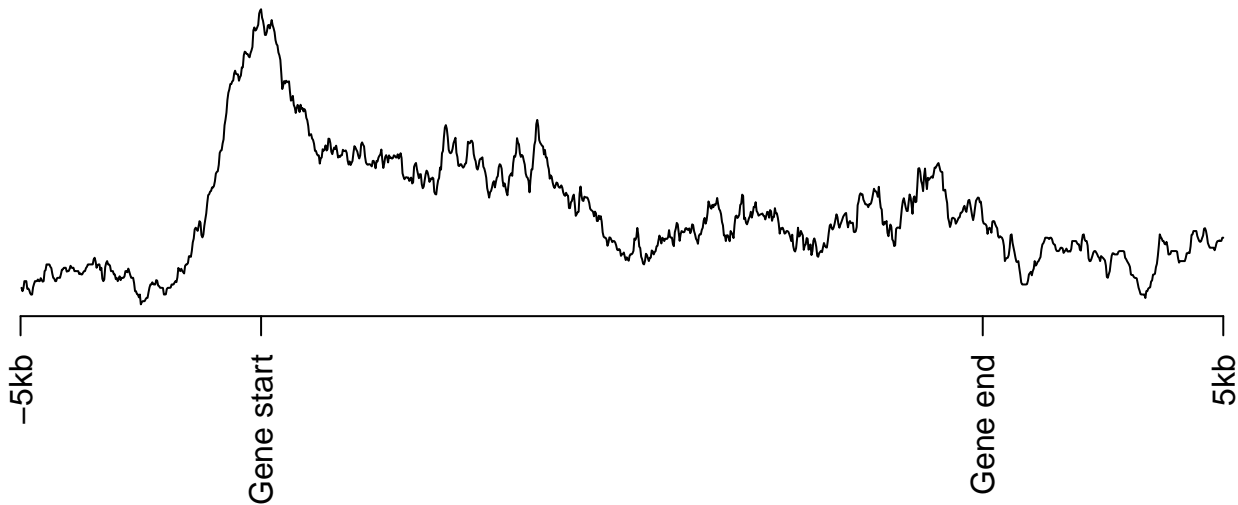




# MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc513

Average coverage per gene

0.0015 0.0030



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc521

Average coverage per gene

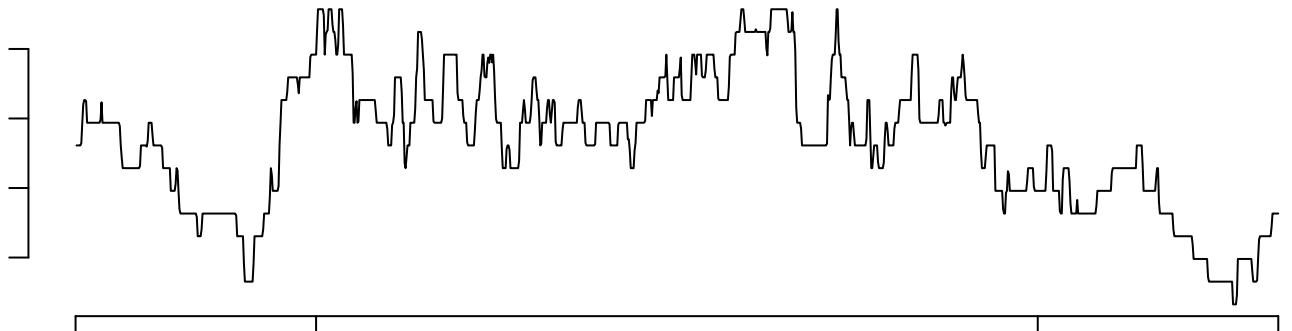
1e-04 3e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc522

Average coverage per gene

1e-04  
3e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc523

Average coverage per gene

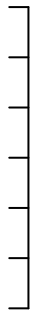
0e+00 3e-04 6e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc524

Average coverage per gene

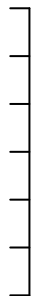
0e+00 3e-04 6e-04

-5kb

Gene start

Gene end

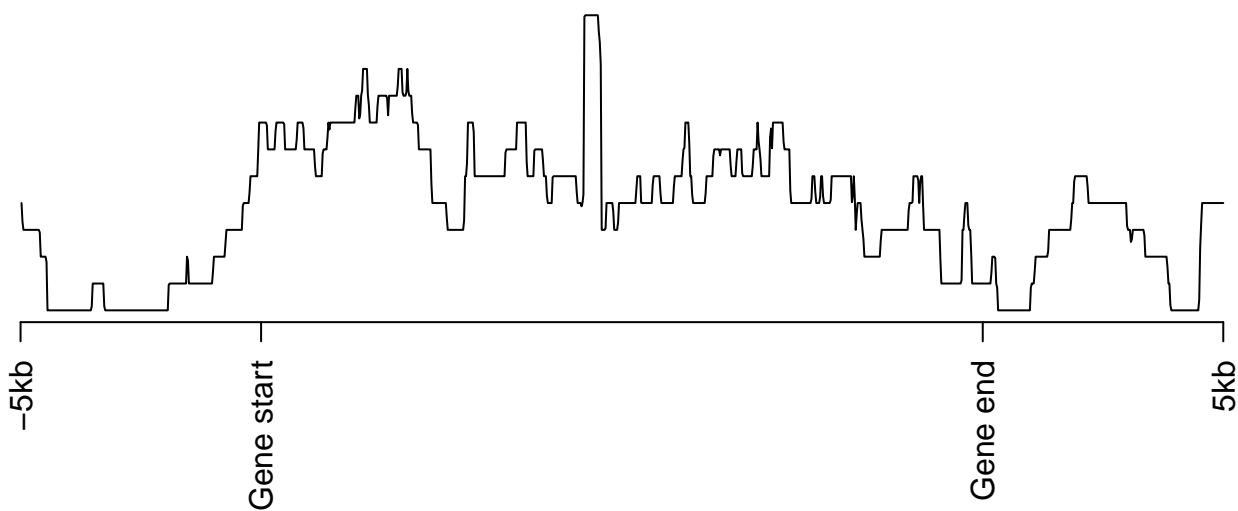
5kb



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc525

Average coverage per gene

0.00005 0.00025



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc526

Average coverage per gene

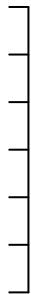
7e-04  
4e-04  
1e-04

-5kb

Gene start

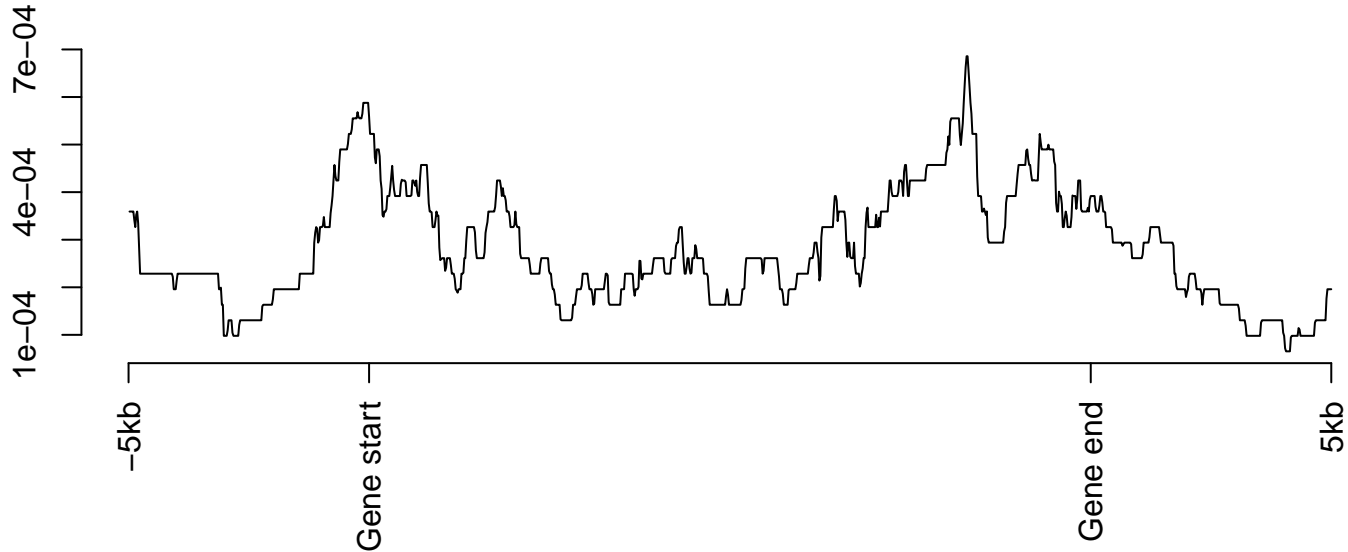
Gene end

5kb



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc527

Average coverage per gene





# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc528

Average coverage per gene

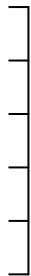
0e+00 6e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc529

Average coverage per gene

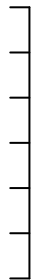
1e-04 4e-04 7e-04

-5kb

Gene start

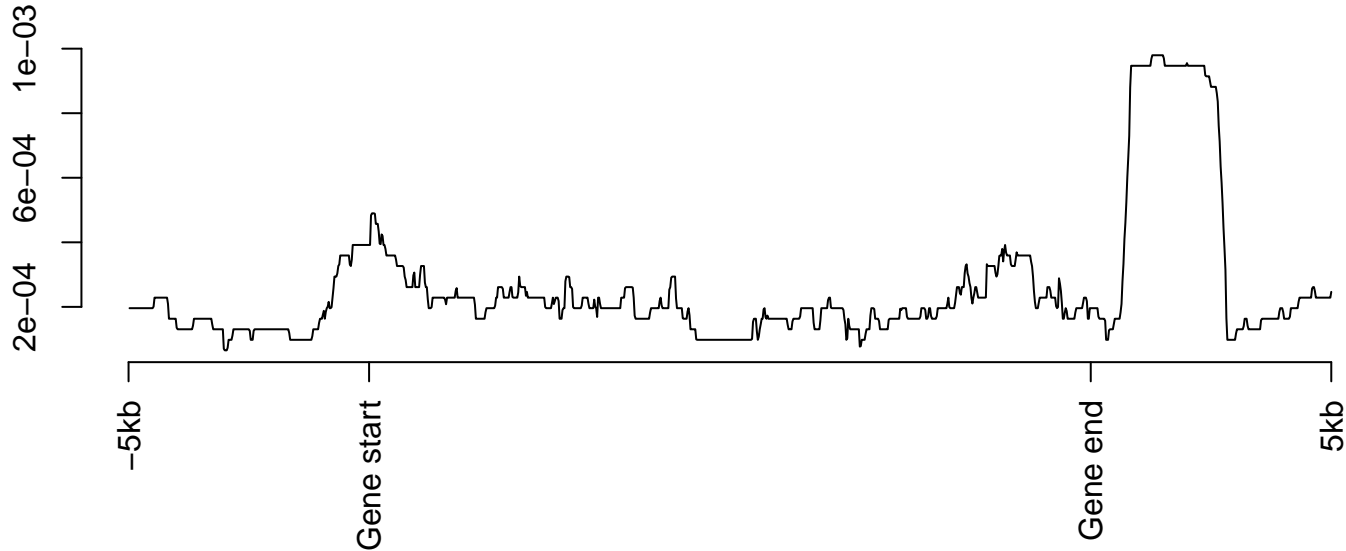
Gene end

5kb



# MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc530

Average coverage per gene



# MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc531

Average coverage per gene

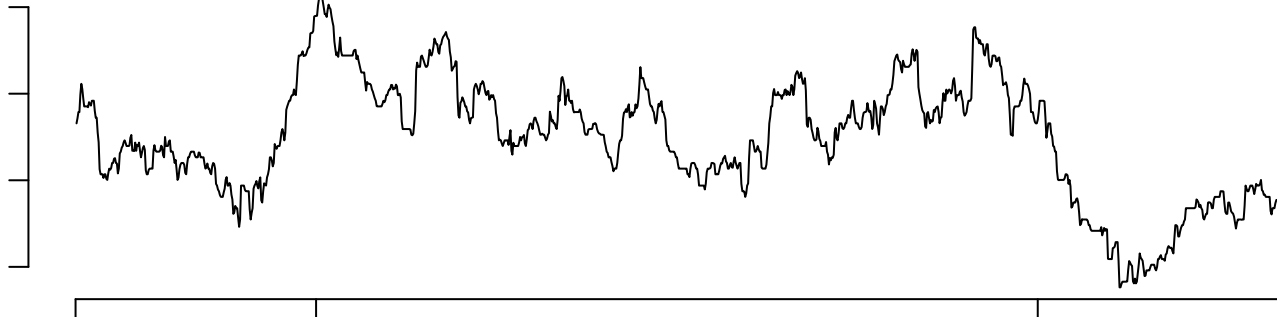
0.0020  
0.0010

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc532

Average coverage per gene

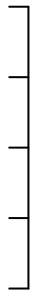
0.0030  
0.0020  
0.0010

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc533

Average coverage per gene

0.0010  
0.0004

-5kb

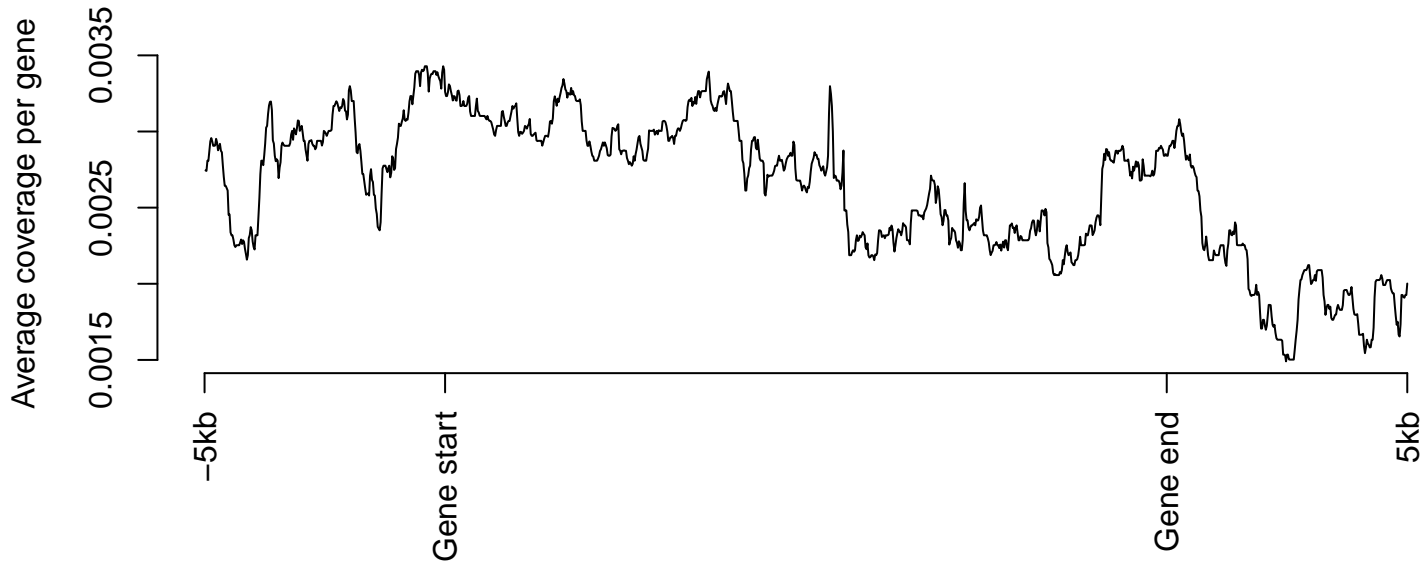
Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc534



# MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc535

Average coverage per gene

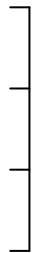
0.010  
0.006

-5kb

Gene start

Gene end

5kb

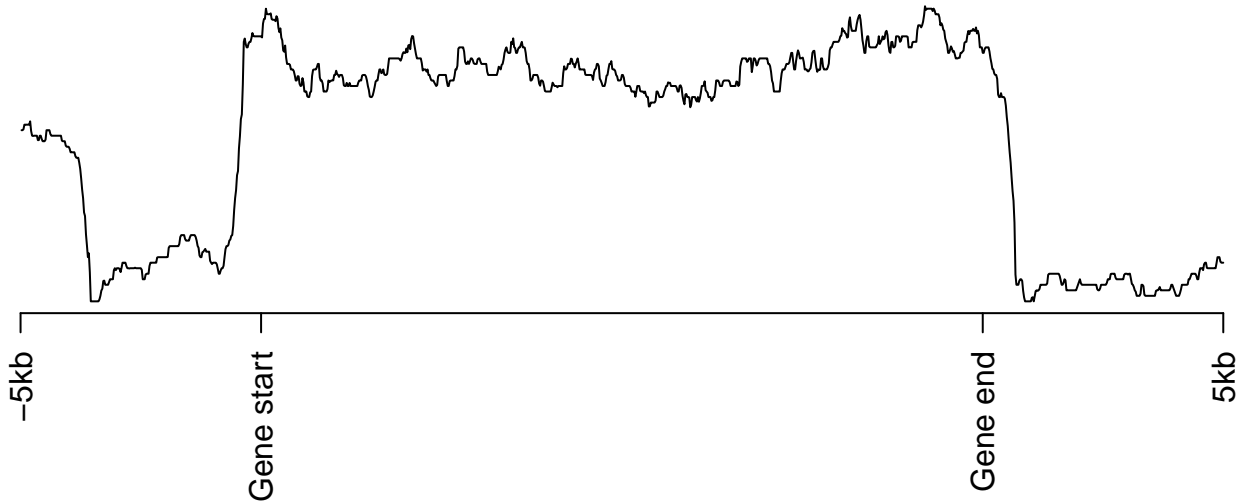




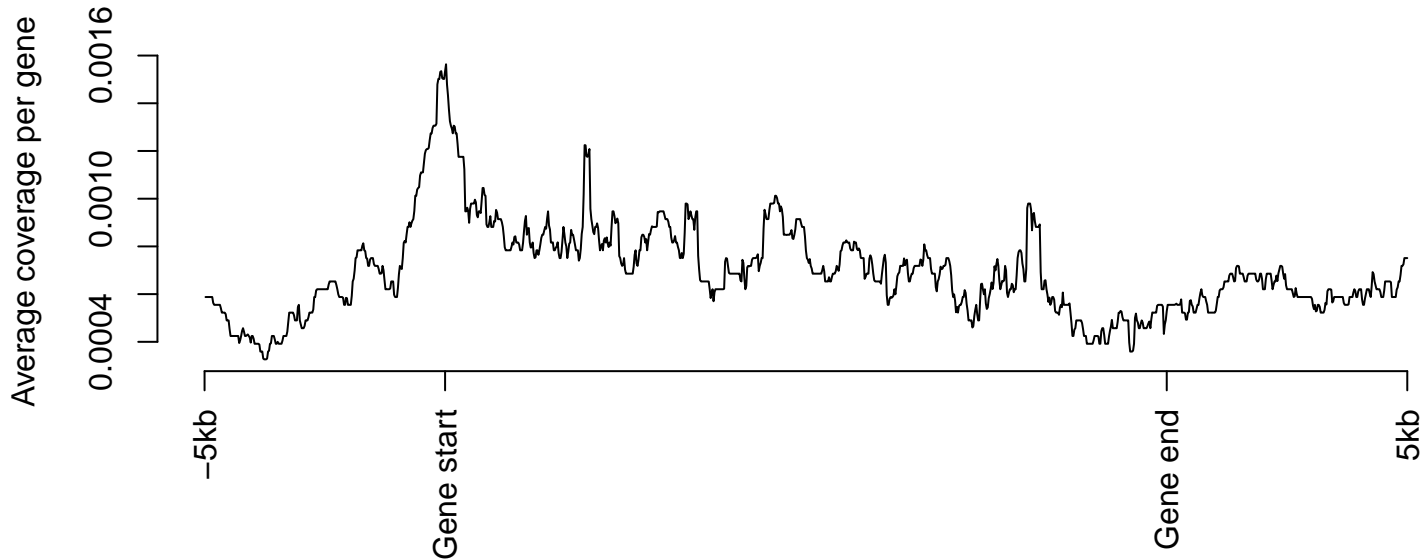
# MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc536

Average coverage per gene

0.0015  
0.0005



# MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc537



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc542

Average coverage per gene

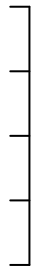
1e-04 3e-04 5e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc543

Average coverage per gene

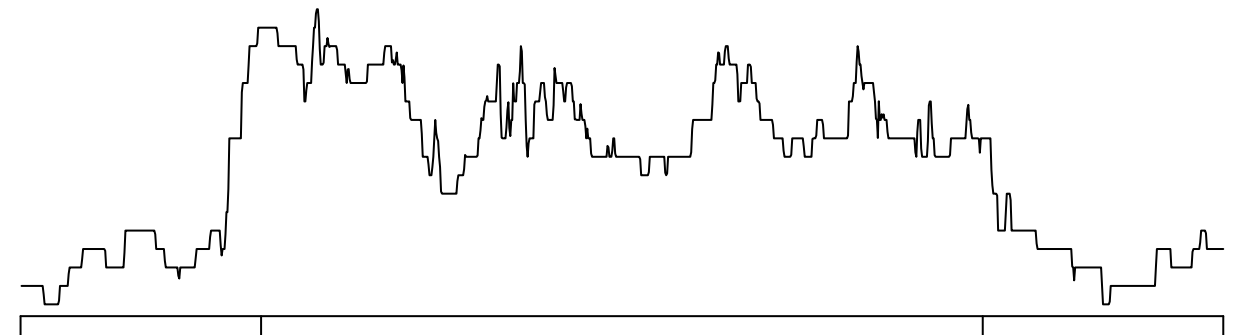
1e-04  
4e-04

-5kb

Gene start

Gene end

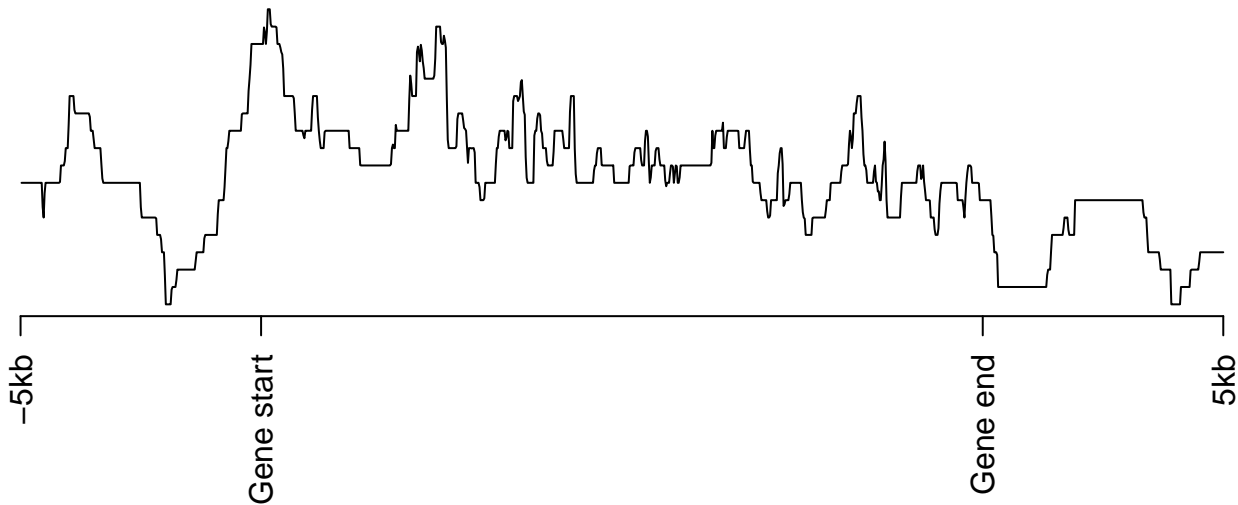
5kb



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc544

Average coverage per gene

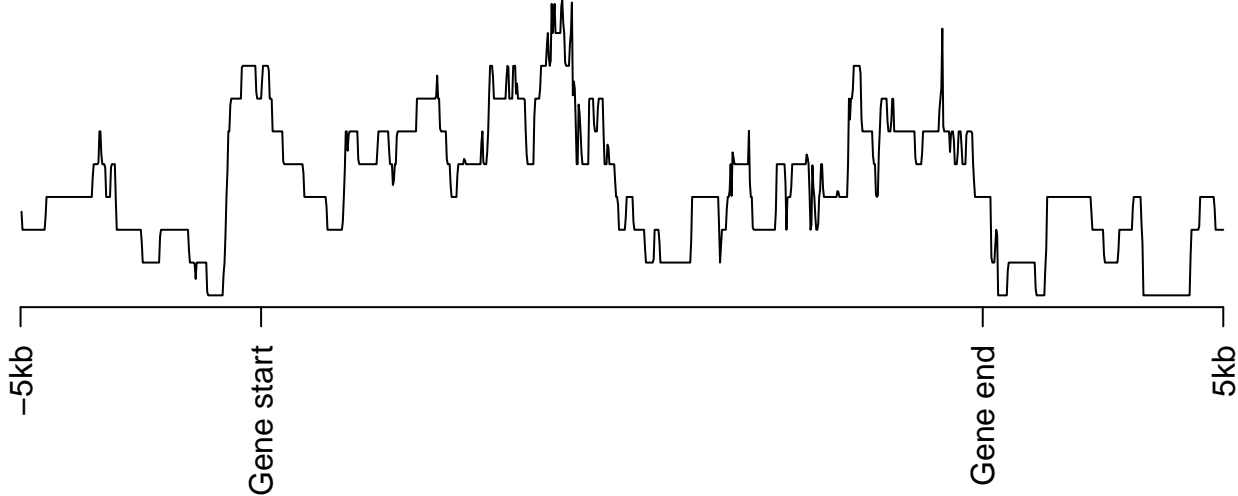
0e+00 3e-04



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc545

Average coverage per gene

0.00010 0.00025



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc546

Average coverage per gene

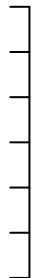
1e-04  
4e-04  
7e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc547

Average coverage per gene

0e+00 3e-04

-5kb

Gene start

Gene end

5kb





# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc548

Average coverage per gene

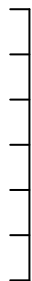
0.00000  
0.00020

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc549

Average coverage per gene

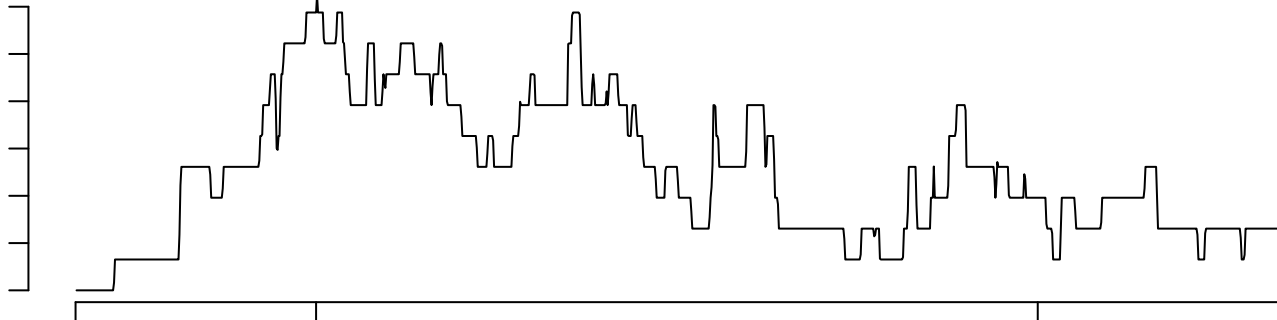
0.00020  
0.00000

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc550

Average coverage per gene

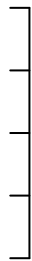
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc551

Average coverage per gene

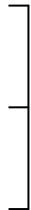
0.0020  
0.0010

-5kb

Gene start

Gene end

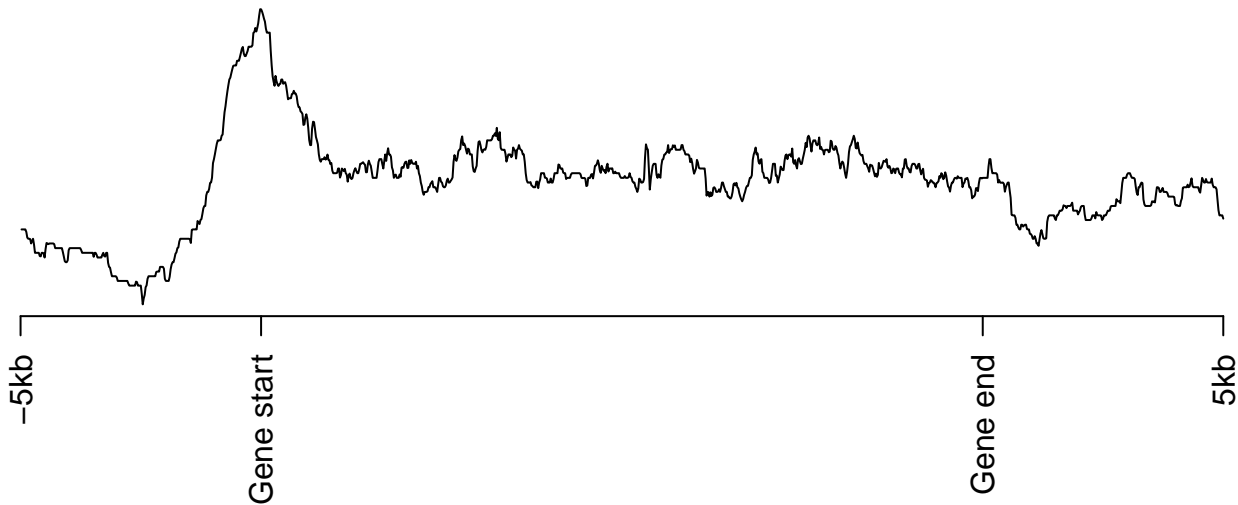
5kb



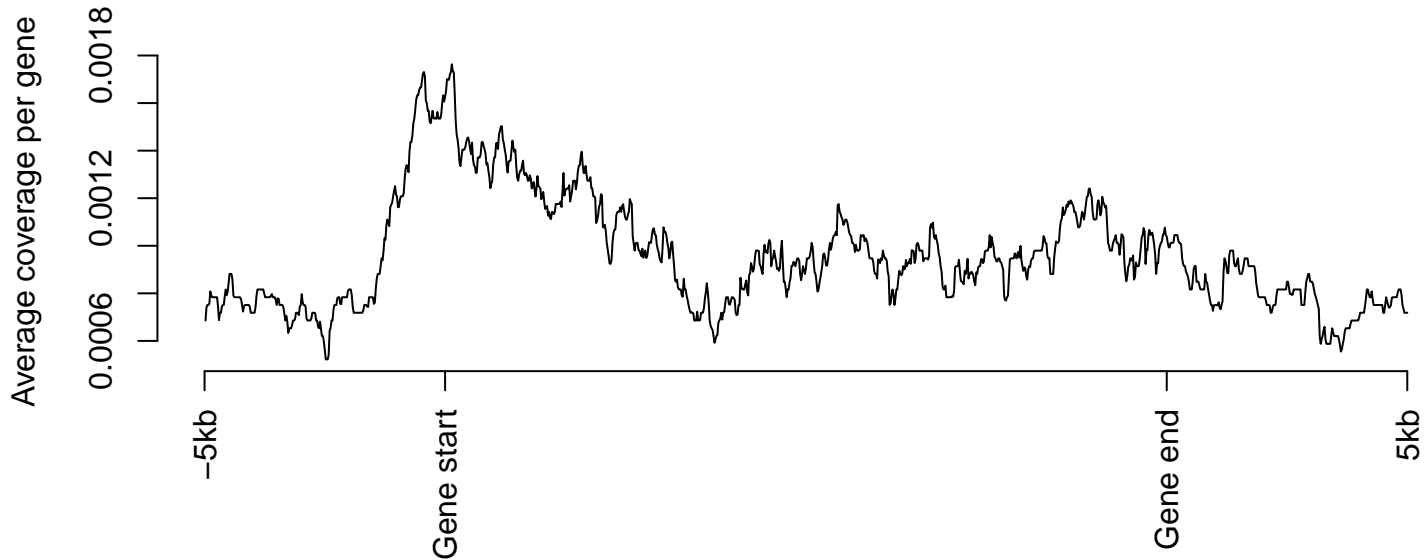
# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc552

Average coverage per gene

0.0005 0.0015



# MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc553



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc554

Average coverage per gene

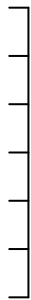
0.0004 0.0010 0.0016

-5kb

Gene start

Gene end

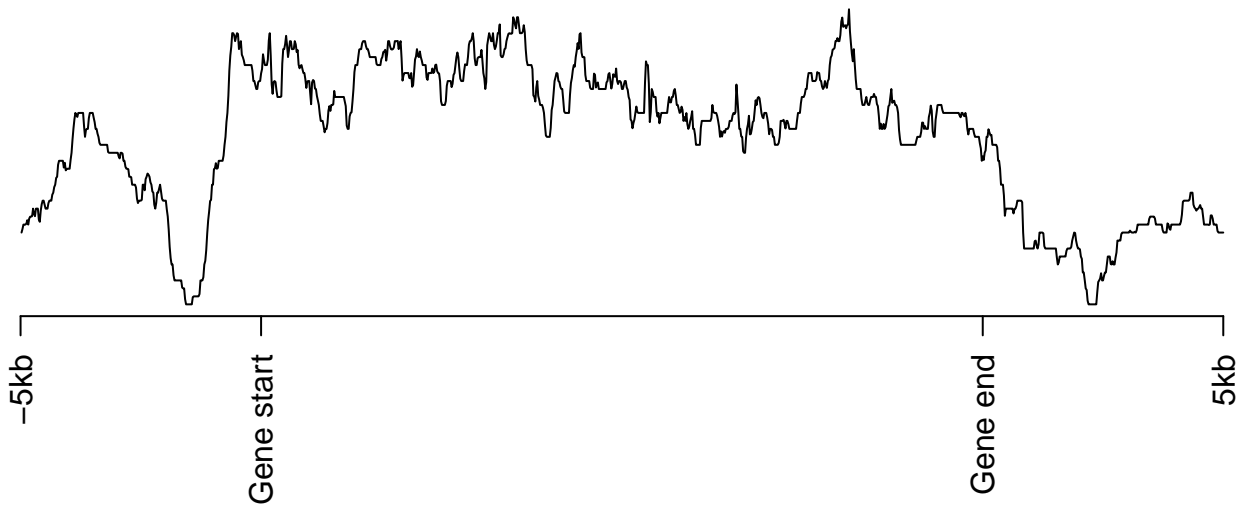
5kb



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc555

Average coverage per gene

0.0004 0.0010





# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc556

Average coverage per gene

0.0015  
0.0005

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc557

Average coverage per gene

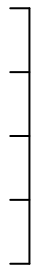
0.0010 0.0020 0.0030

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc558

Average coverage per gene

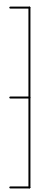
0.0020  
0.0010

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc559

Average coverage per gene

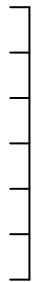
0.00025  
0.00005

-5kb

Gene start

Gene end

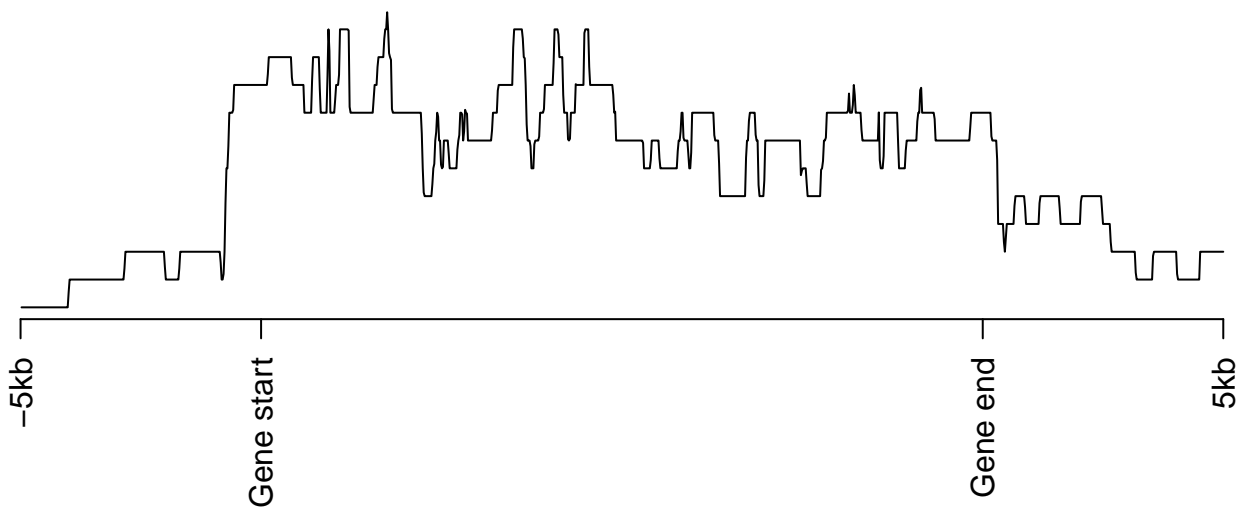
5kb



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc561

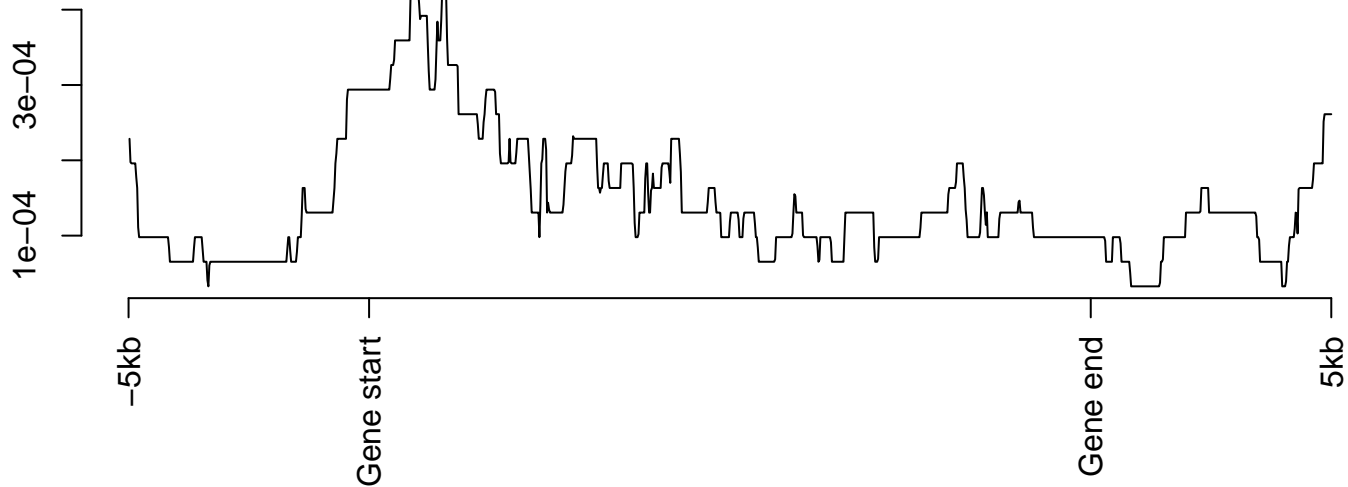
Average coverage per gene

0.00000 0.00020



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc562

Average coverage per gene



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc563

Average coverage per gene

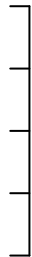
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc564

Average coverage per gene

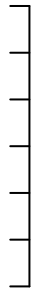
7e-04  
4e-04  
1e-04

-5kb

Gene start

Gene end

5kb





# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc565

Average coverage per gene

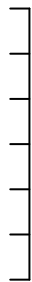
1e-04  
4e-04  
7e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc566

Average coverage per gene

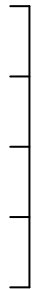
0e+00 2e-04 4e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc567

Average coverage per gene

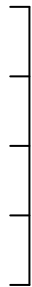
1e-04  
3e-04  
5e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc568

Average coverage per gene

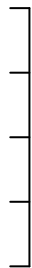
0e+00 2e-04 4e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc569

Average coverage per gene

0.00000 0.00015

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc570

Average coverage per gene

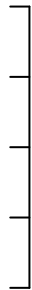
0e+00 4e-04 8e-04

-5kb

Gene start

Gene end

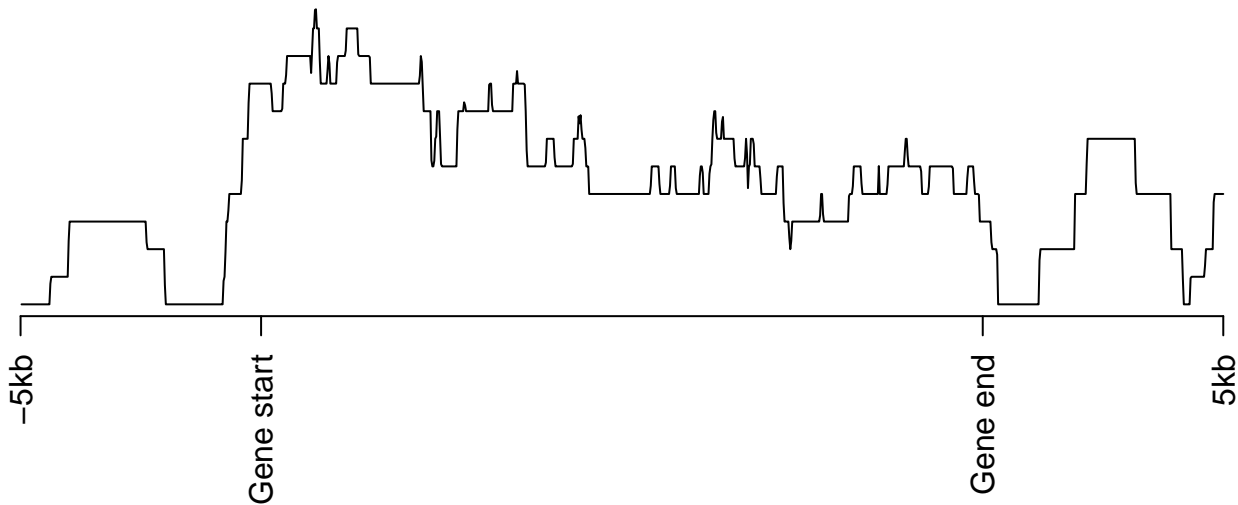
5kb



# MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc571

Average coverage per gene

0.00000 0.00020



# MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc572

Average coverage per gene

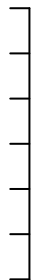
0.00000  
0.00020

-5kb

Gene start

Gene end

5kb

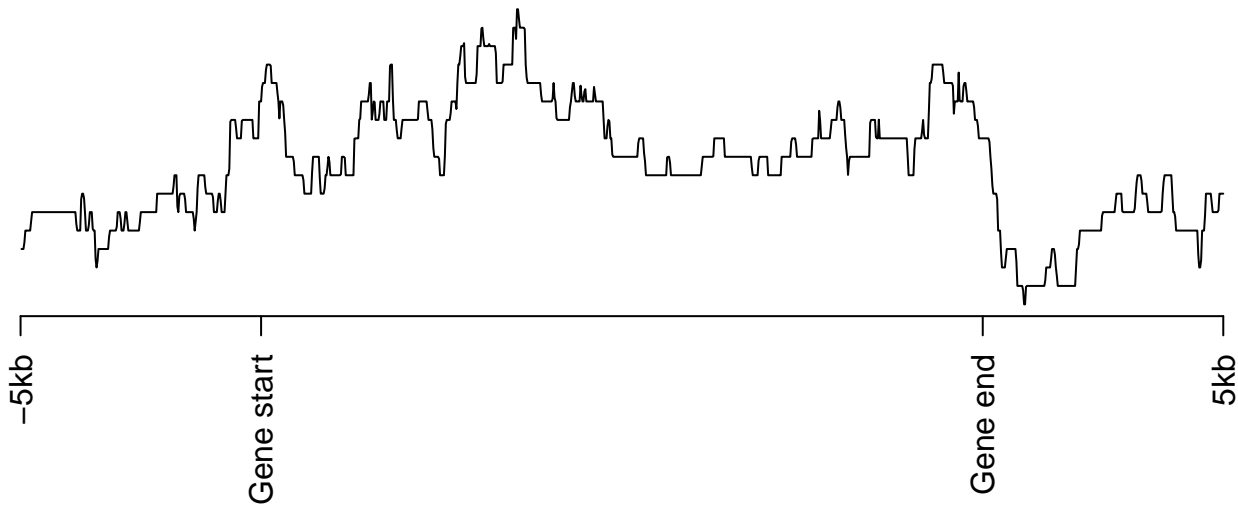




# MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc573

Average coverage per gene

1e-04  
4e-04



# MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc576

Average coverage per gene

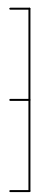
0.0030  
0.0020

-5kb

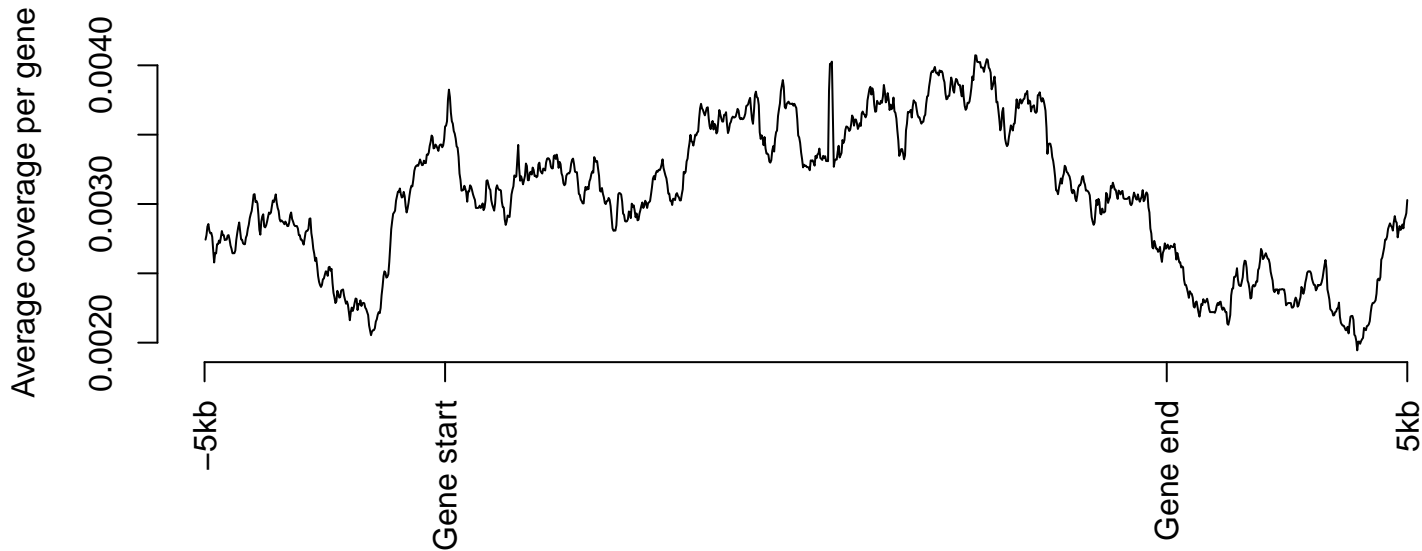
Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc601



# MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc602

Average coverage per gene

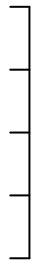
0.0025 0.0035 0.0045

-5kb

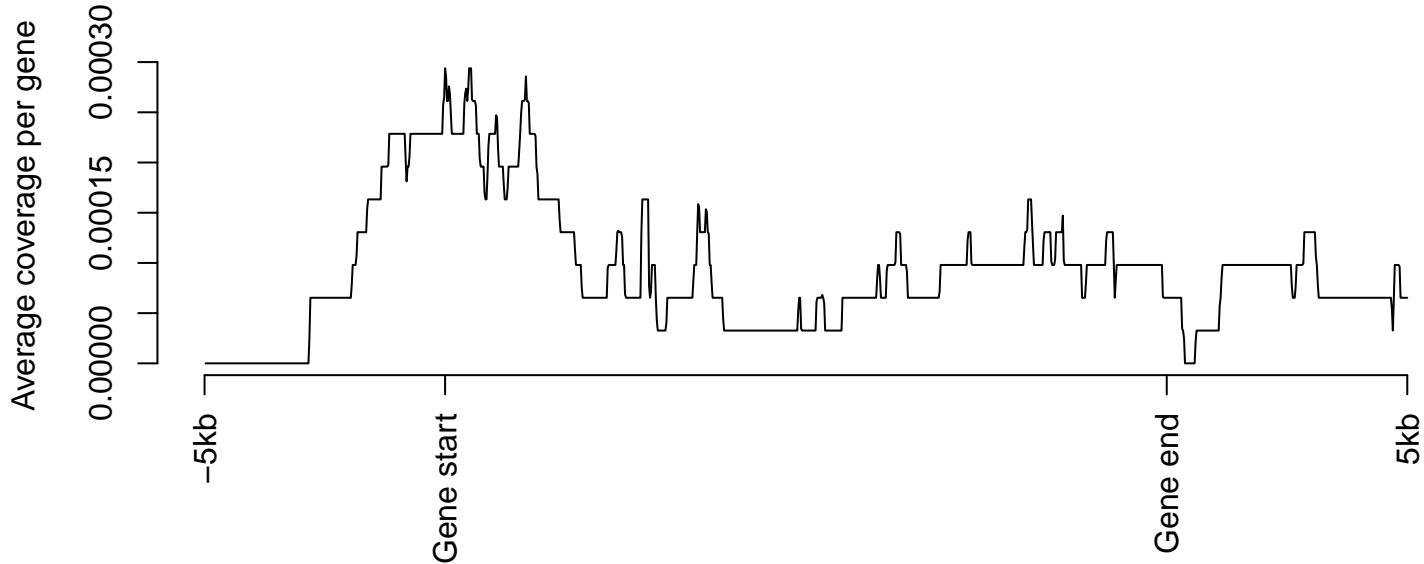
Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc577b



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc578b

Average coverage per gene

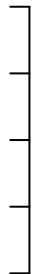
0e+00 2e-04 4e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc579b

Average coverage per gene

0e+00

2e-04

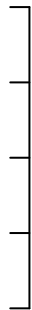
4e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc580b

Average coverage per gene

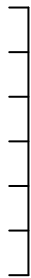
0.0065  
0.0050  
0.0035

-5kb

Gene start

Gene end

5kb





# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc581b

Average coverage per gene

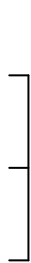
0.0015 0.0025

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc582b

Average coverage per gene

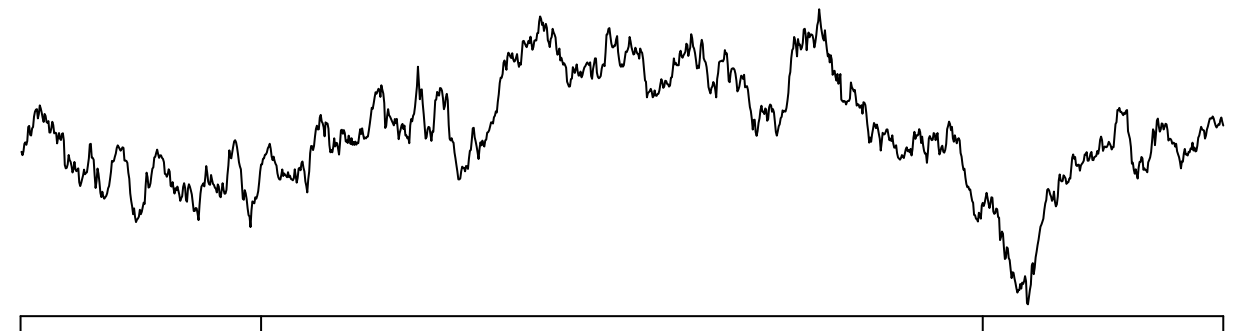
0.0040 0.0055

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc583b

Average coverage per gene

0.0006 0.0012

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc584b

Average coverage per gene

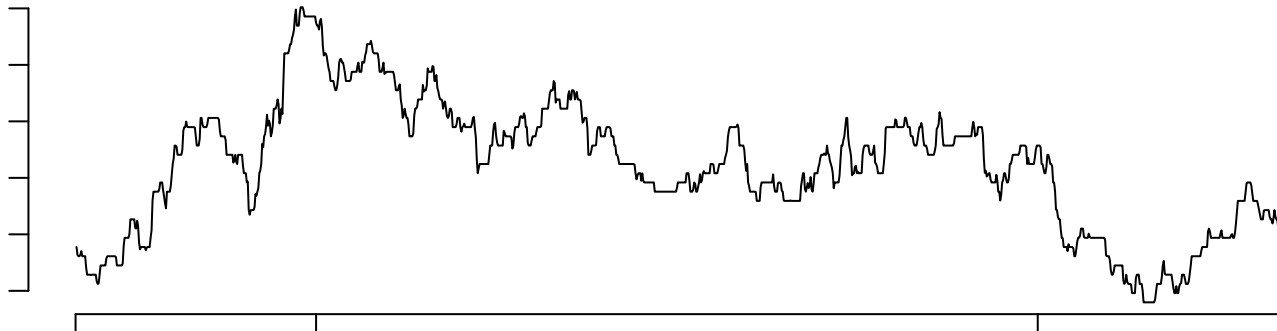
0.0004 0.0010

-5kb

Gene start

Gene end

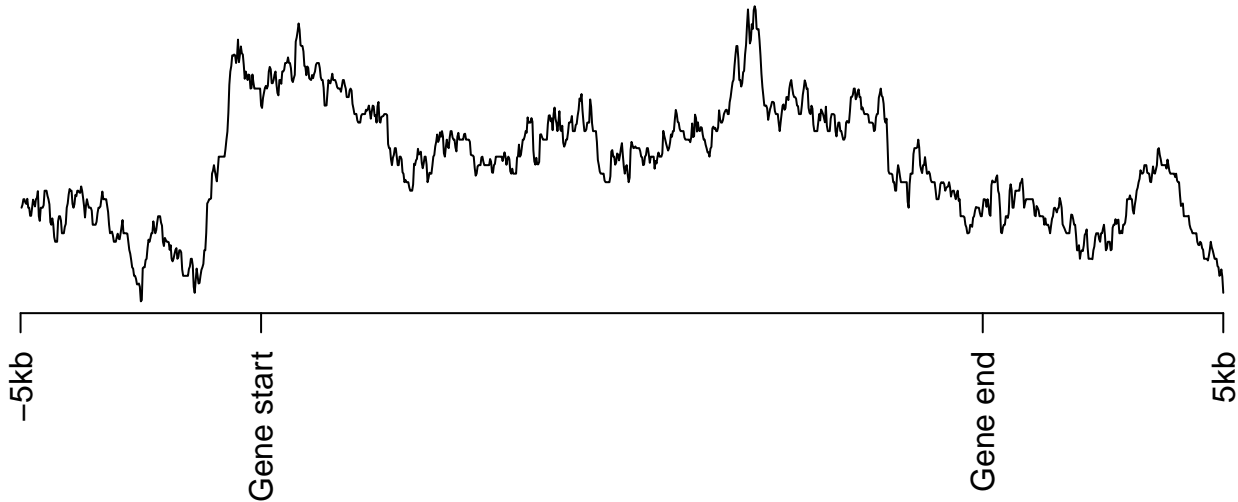
5kb



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc585b

Average coverage per gene

0.0012 0.0018



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc586b

Average coverage per gene

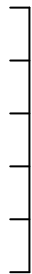
0.0010 0.0025

-5kb

Gene start

Gene end

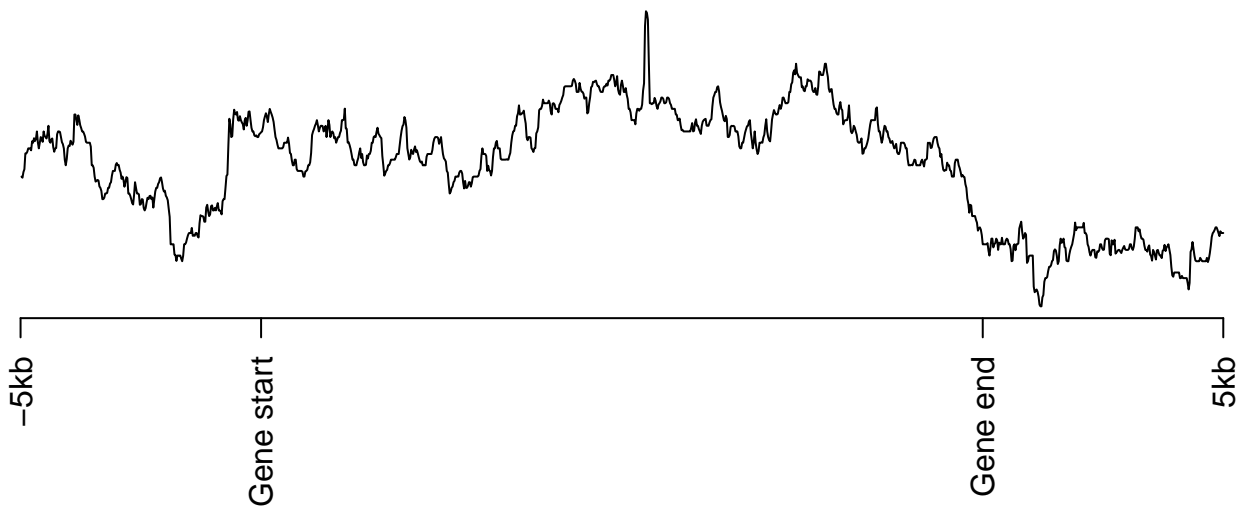
5kb



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc587b

Average coverage per gene

0.0025  
0.0015



# MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc588b

Average coverage per gene

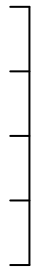
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

5kb





# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc589b

Average coverage per gene

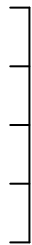
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

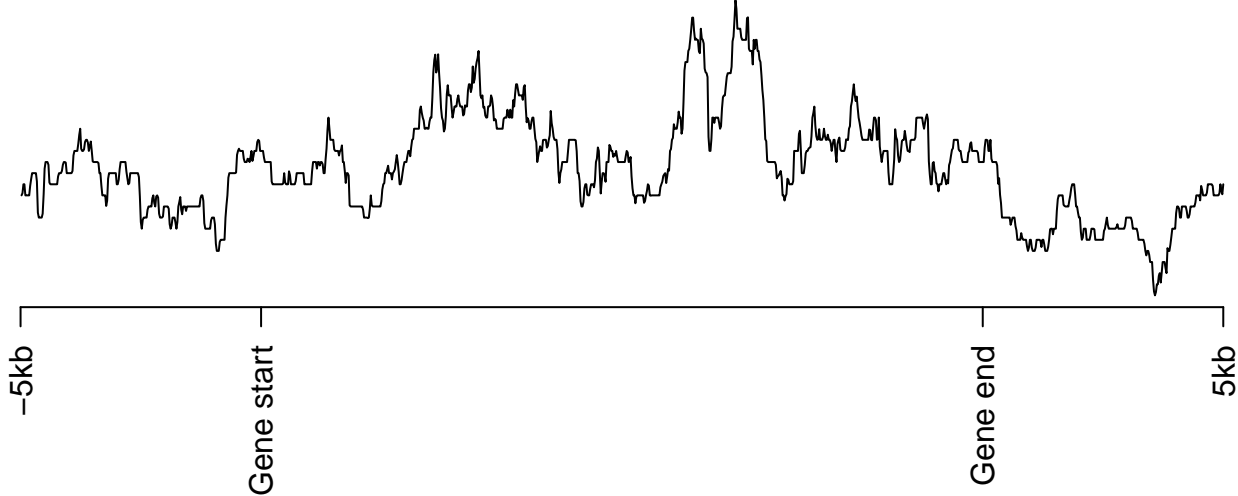
5kb



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc590b

Average coverage per gene

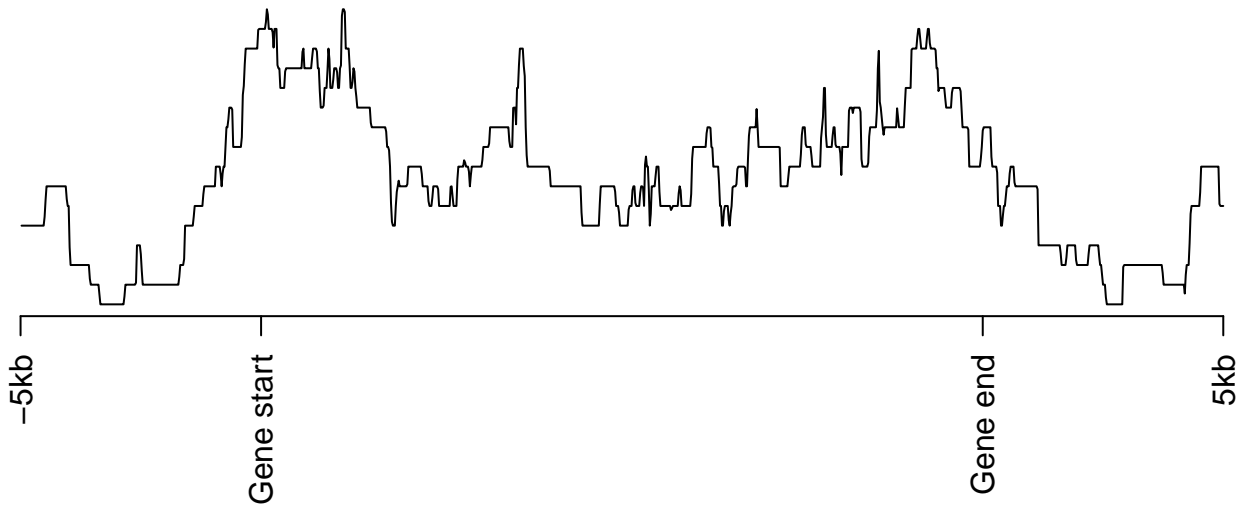
0.0004 0.0008 0.0012



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc591b

Average coverage per gene

1e-04 3e-04 5e-04



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc592b

Average coverage per gene

0.0008  
0.0002

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc593b

Average coverage per gene

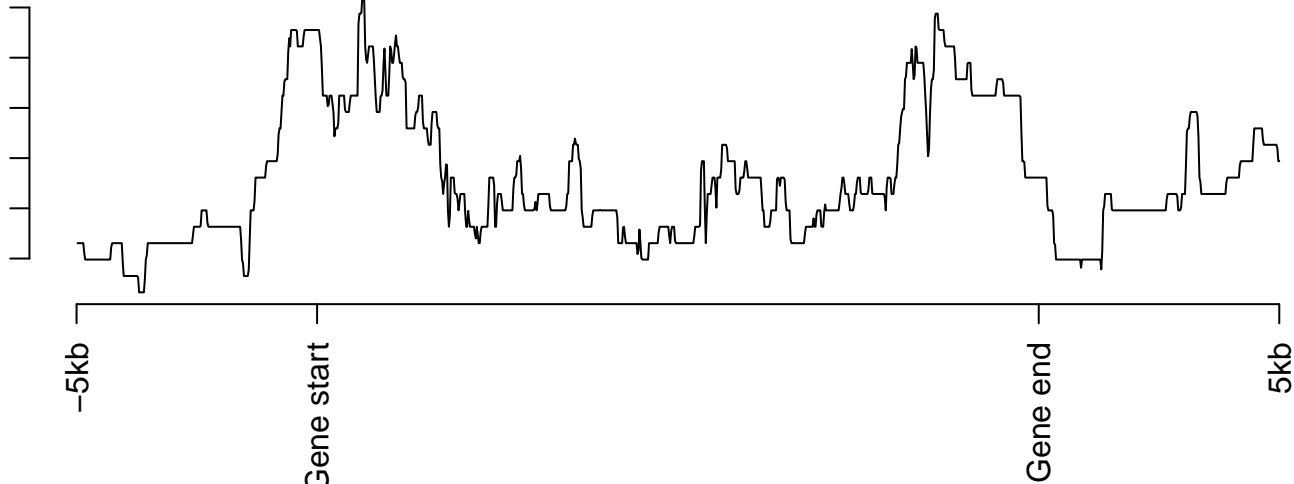
1e-04  
4e-04

-5kb

Gene start

Gene end

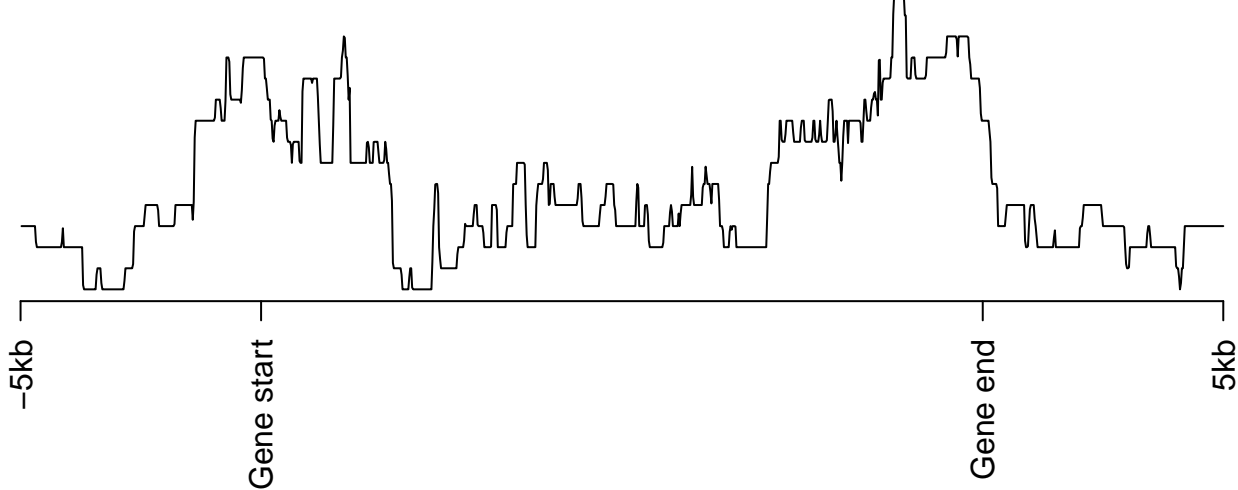
5kb



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc594b

Average coverage per gene

1e-04  
3e-04  
5e-04



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc595b

Average coverage per gene

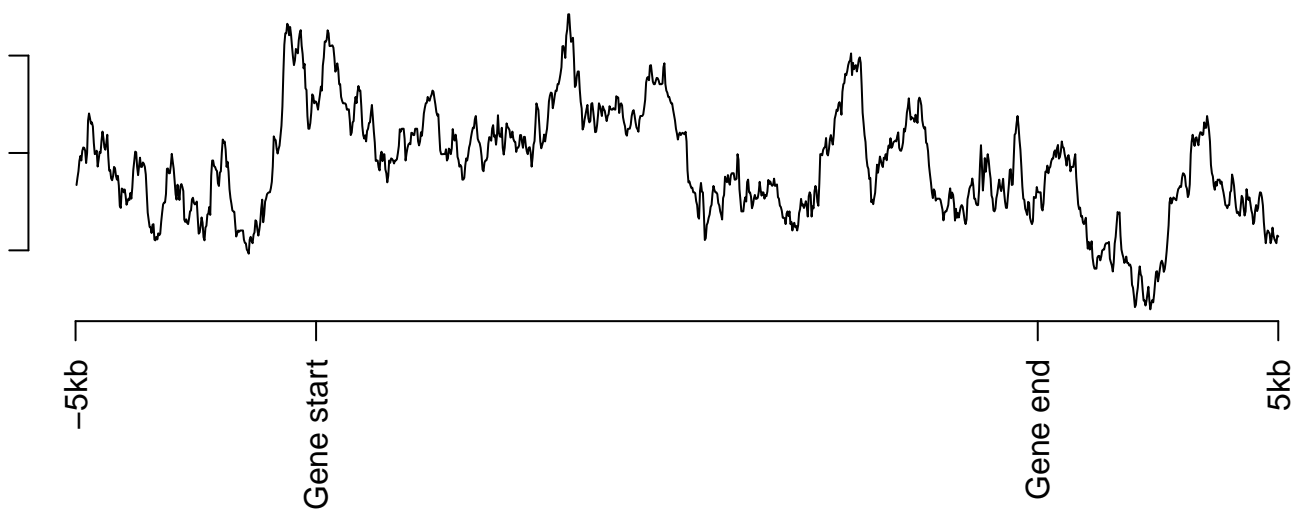
0.0040  
0.0030

-5kb

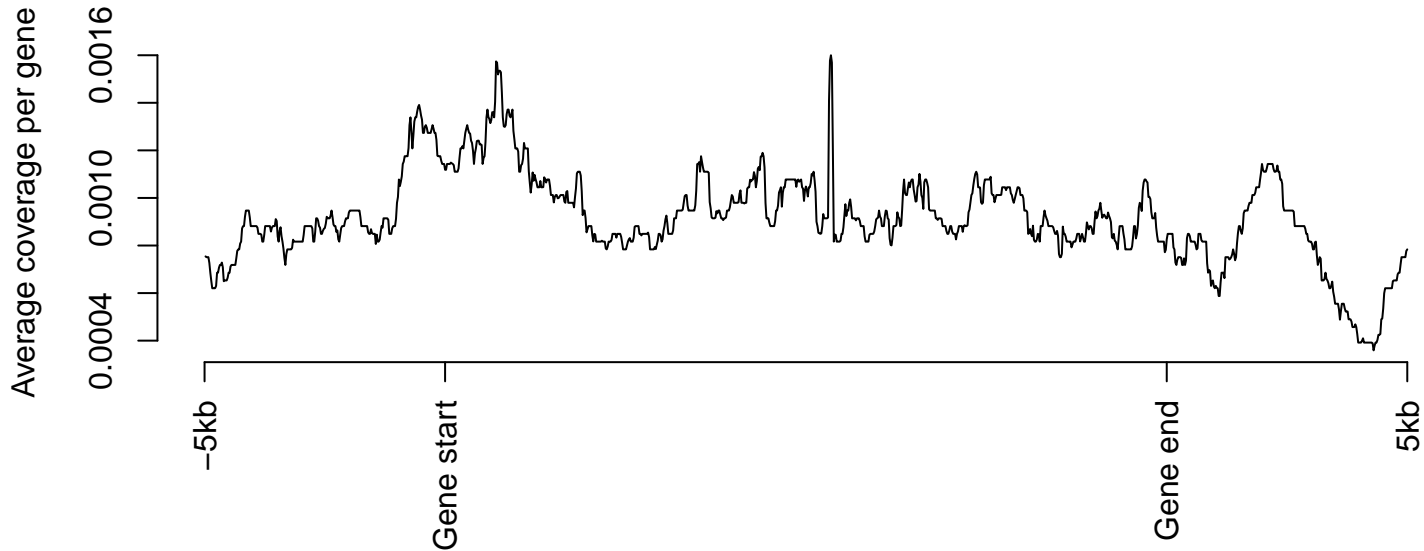
Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc596b





# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc597b

Average coverage per gene

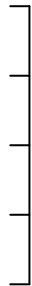
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc598b

Average coverage per gene

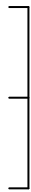
0.0020  
0.0010

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc599b

Average coverage per gene

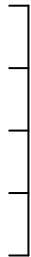
0.0030 0.0040 0.0050

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc600b

Average coverage per gene

0.0015 0.0025

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc779

Average coverage per gene

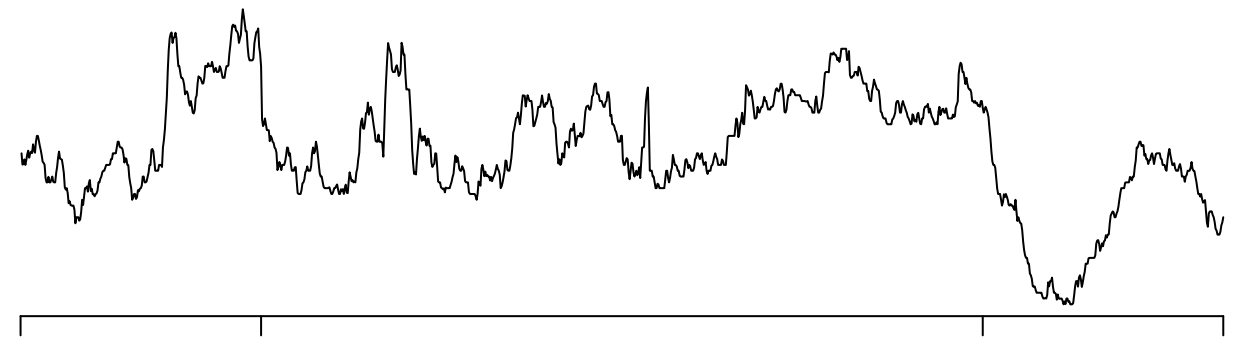
0.0020  
0.0010

-5kb

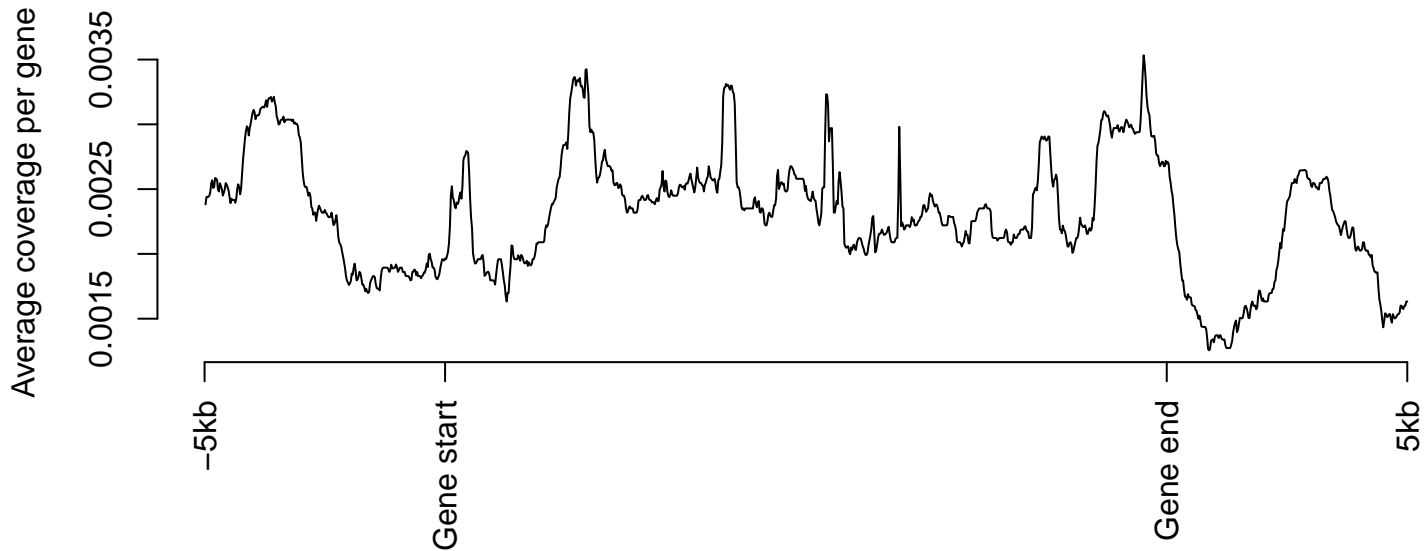
Gene start

Gene end

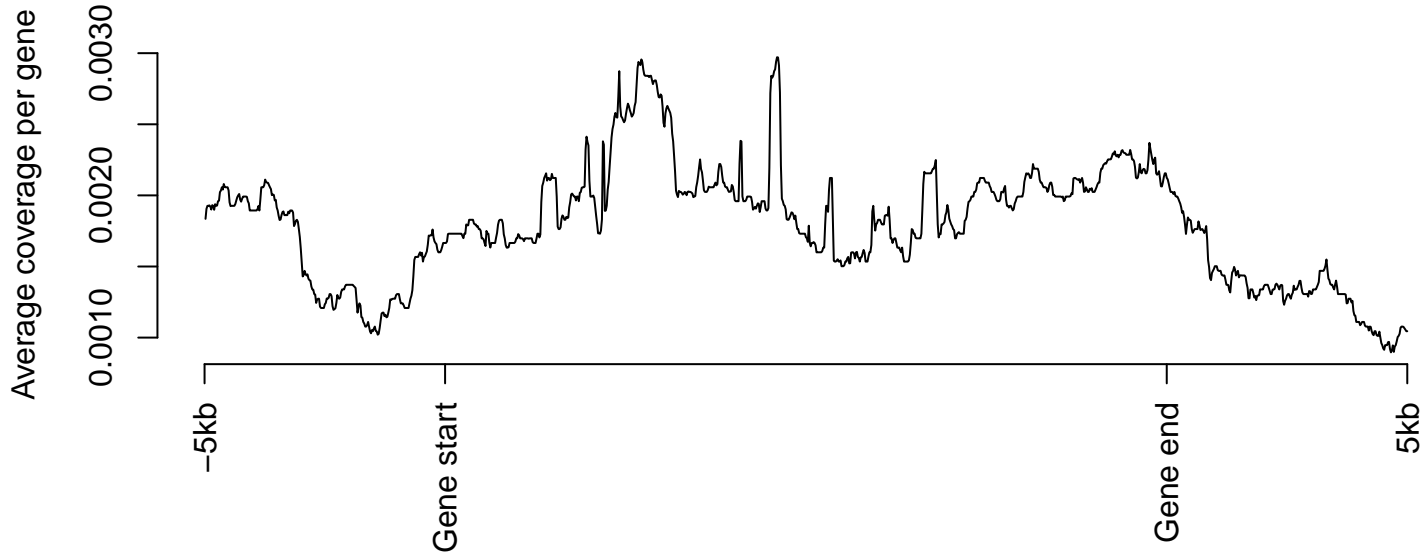
5kb



# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc780



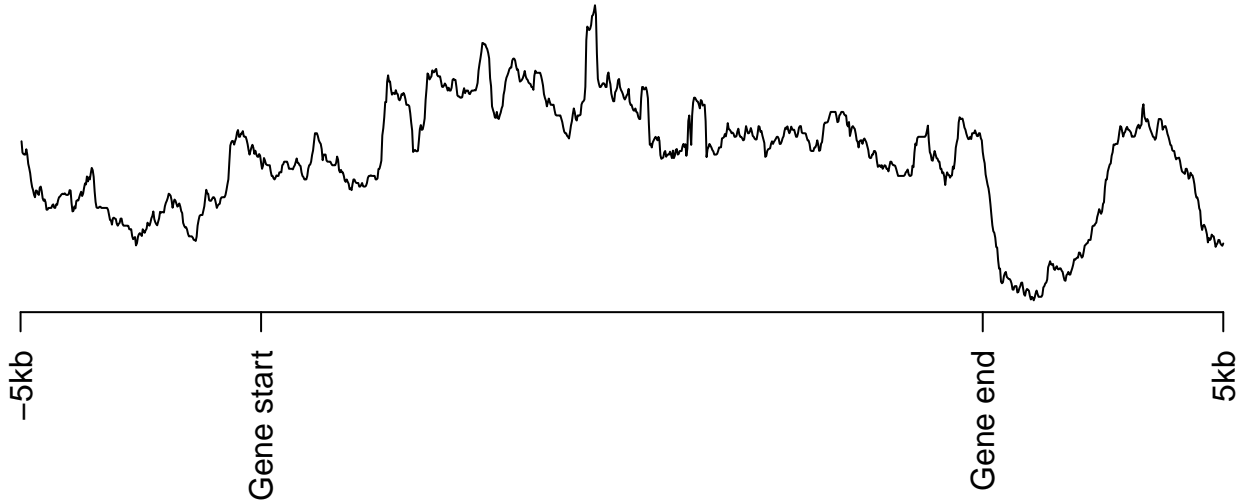
# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc781



# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc782

Average coverage per gene

0.0030  
0.0015





# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc783

Average coverage per gene

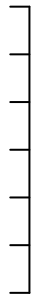
0.0015 0.0030 0.0045

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc784

Average coverage per gene

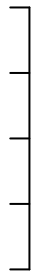
0.0010 0.0020 0.0030

-5kb

Gene start

Gene end

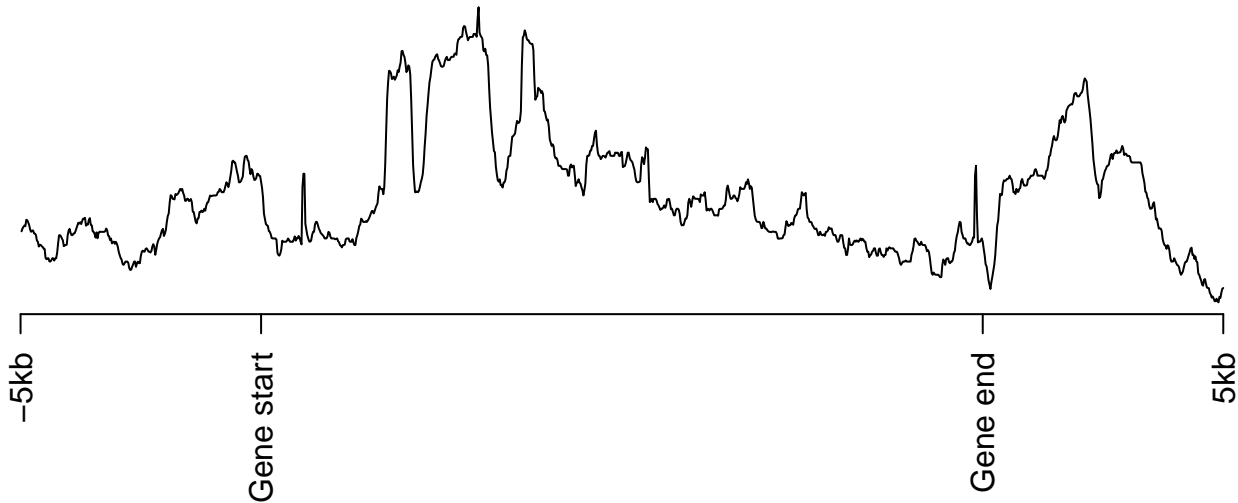
5kb



# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc785

Average coverage per gene

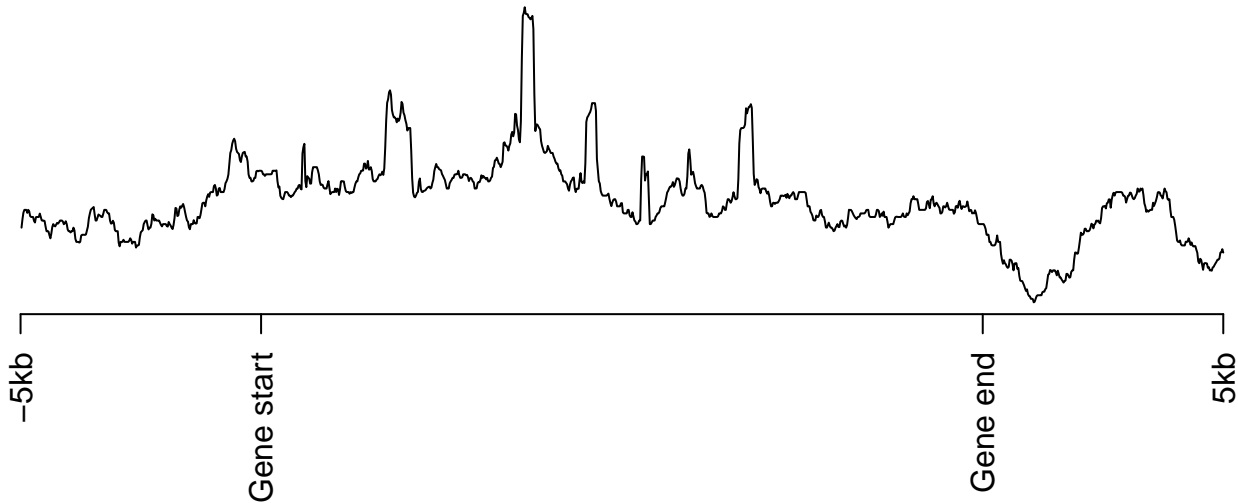
0.0015 0.0030



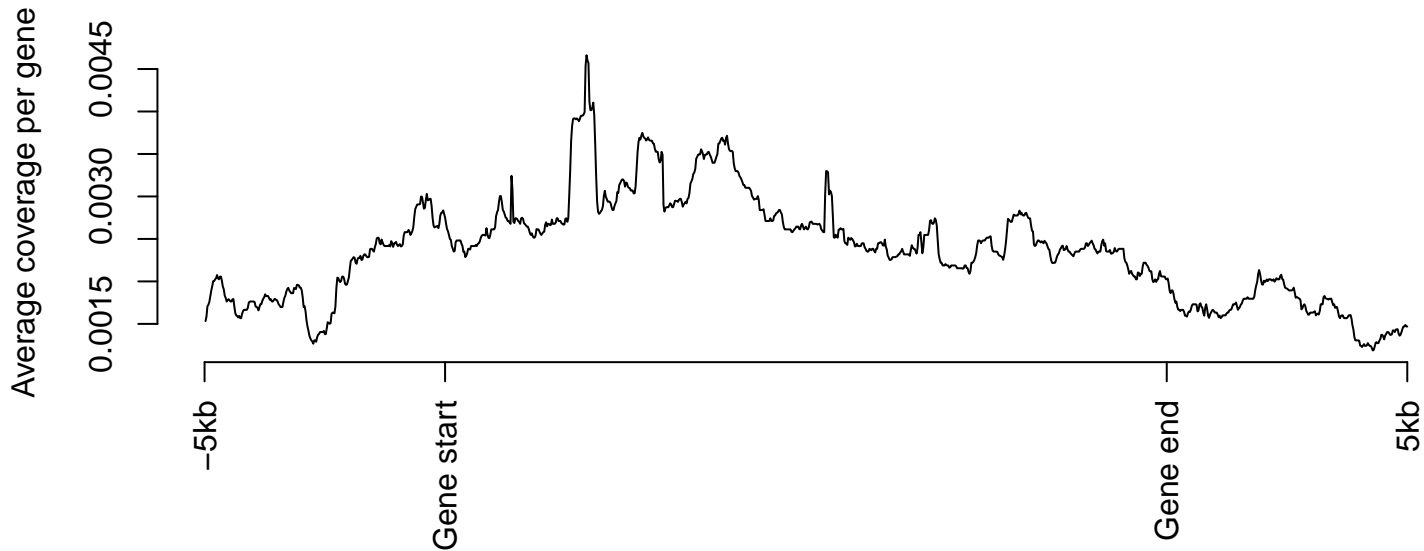
# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc786

Average coverage per gene

0.0010 0.0025



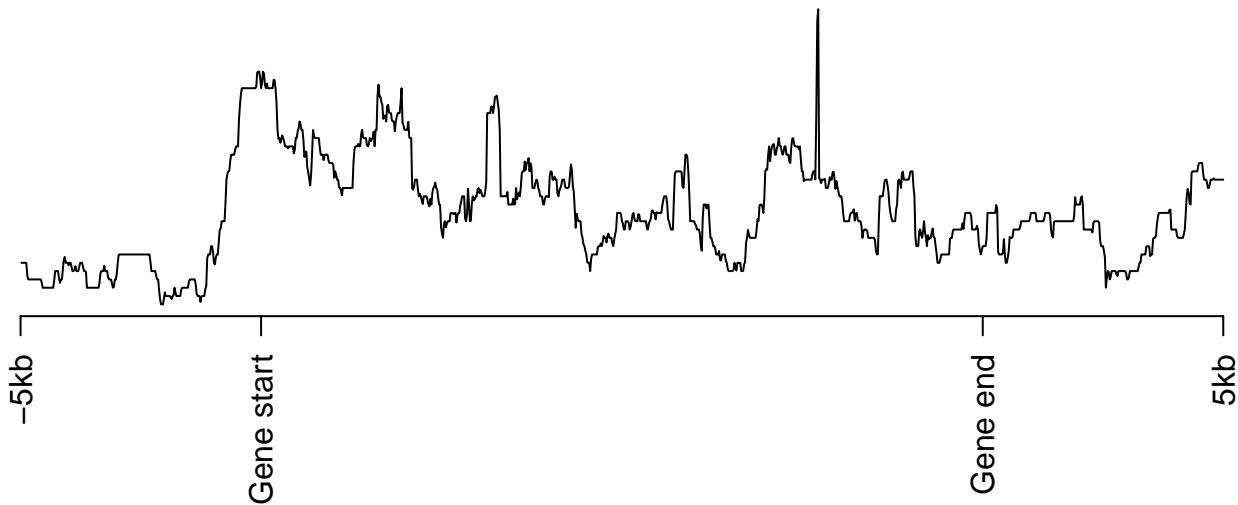
# MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc787



# MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc788

Average coverage per gene

0.0010  
0.0004



# MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc789

Average coverage per gene

2e-04  
6e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc790

Average coverage per gene

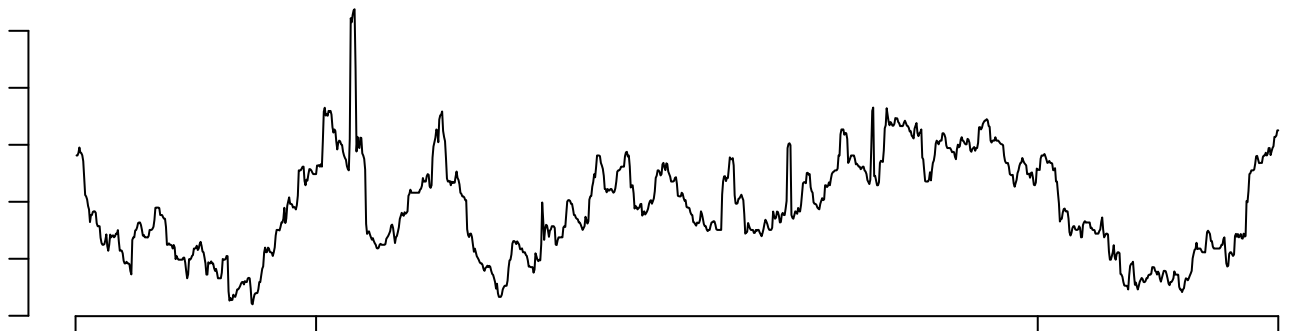
0.0030  
0.0015

-5kb

Gene start

Gene end

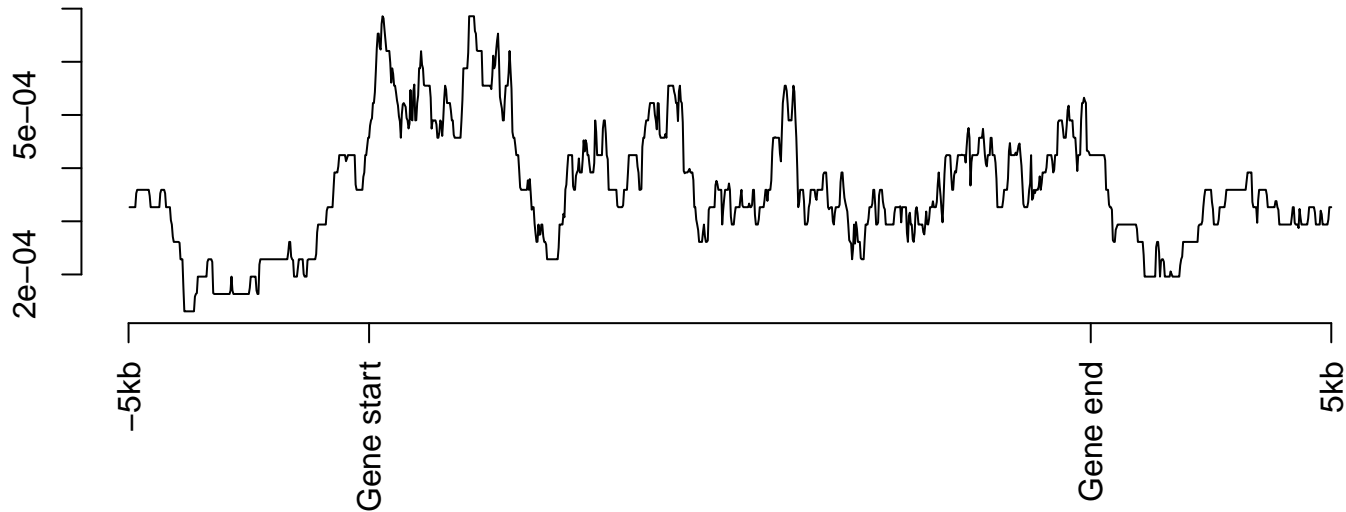
5kb





# MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc791

Average coverage per gene



# MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc792

Average coverage per gene

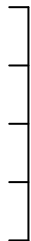
0.0010  
0.0004

-5kb

Gene start

Gene end

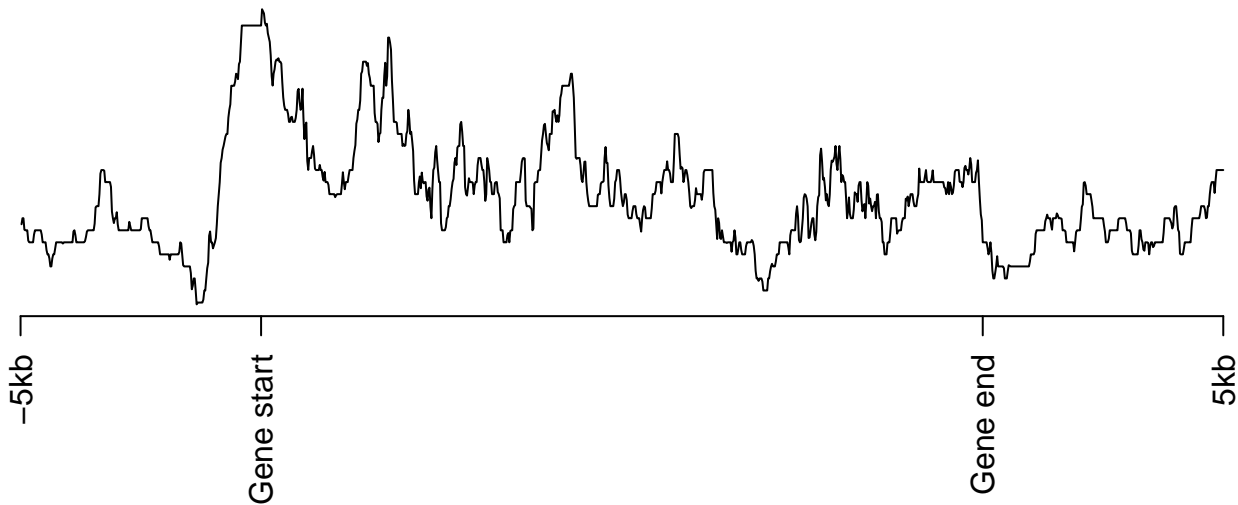
5kb



# MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc793

Average coverage per gene

4e-04  
8e-04



# MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc794

Average coverage per gene

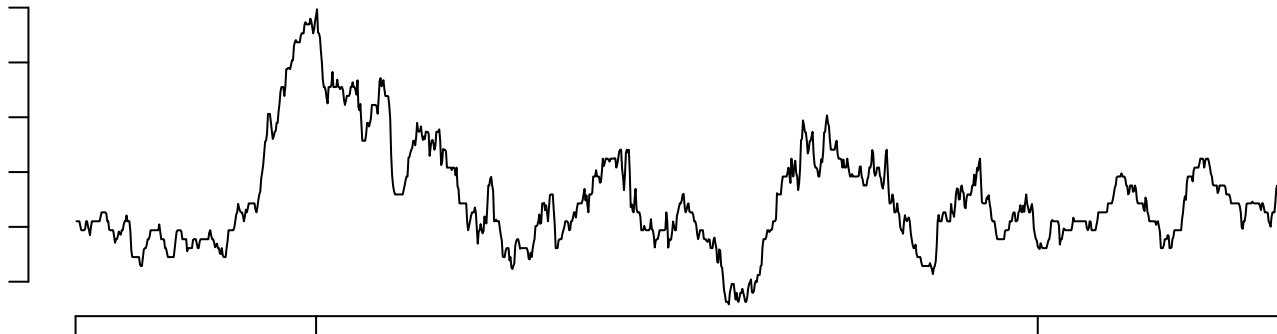
0.0010  
0.0004

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc795

Average coverage per gene

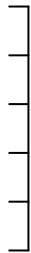
0.0004 0.0010

-5kb

Gene start

Gene end

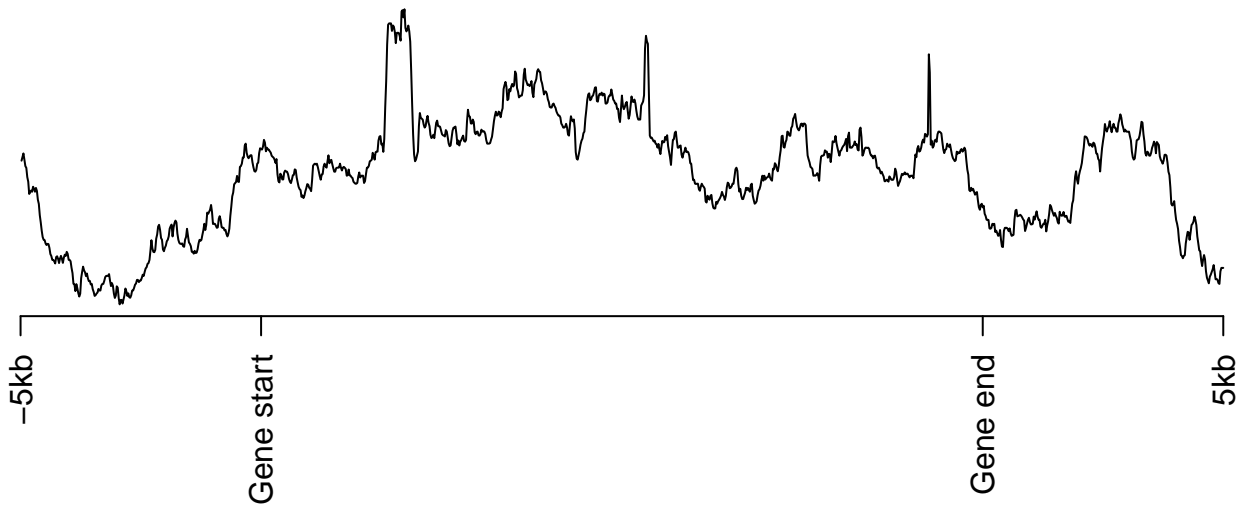
5kb



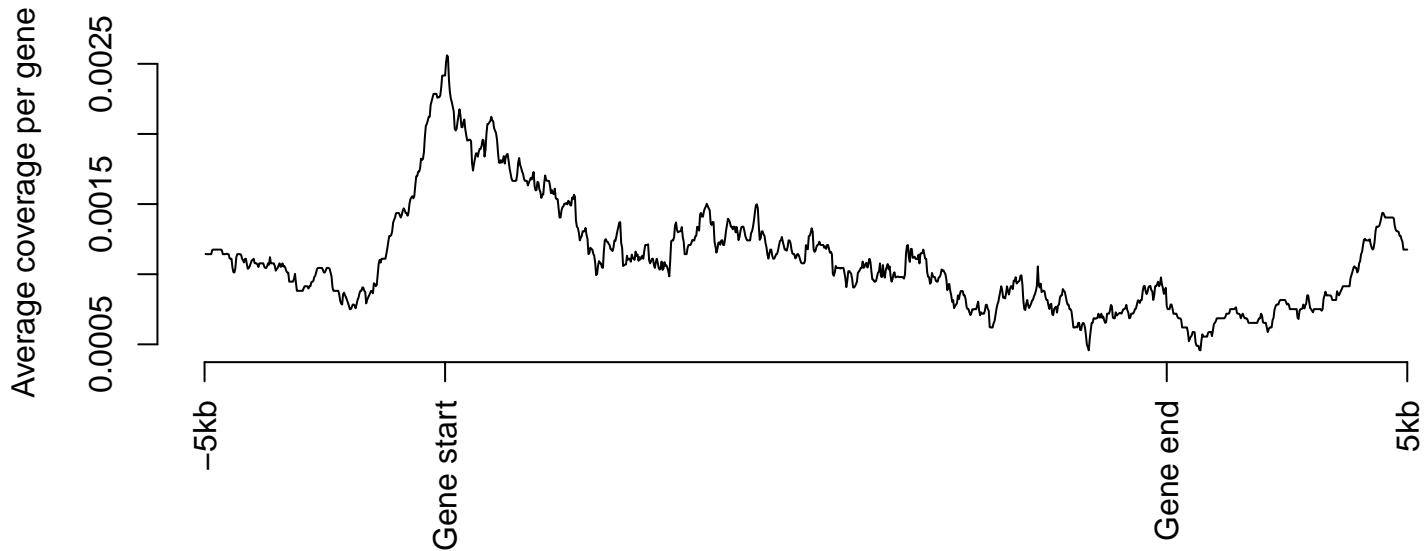
# MouseNeuronCulture PositiveSingle E.669 scCLTdegenNuc808

Average coverage per gene

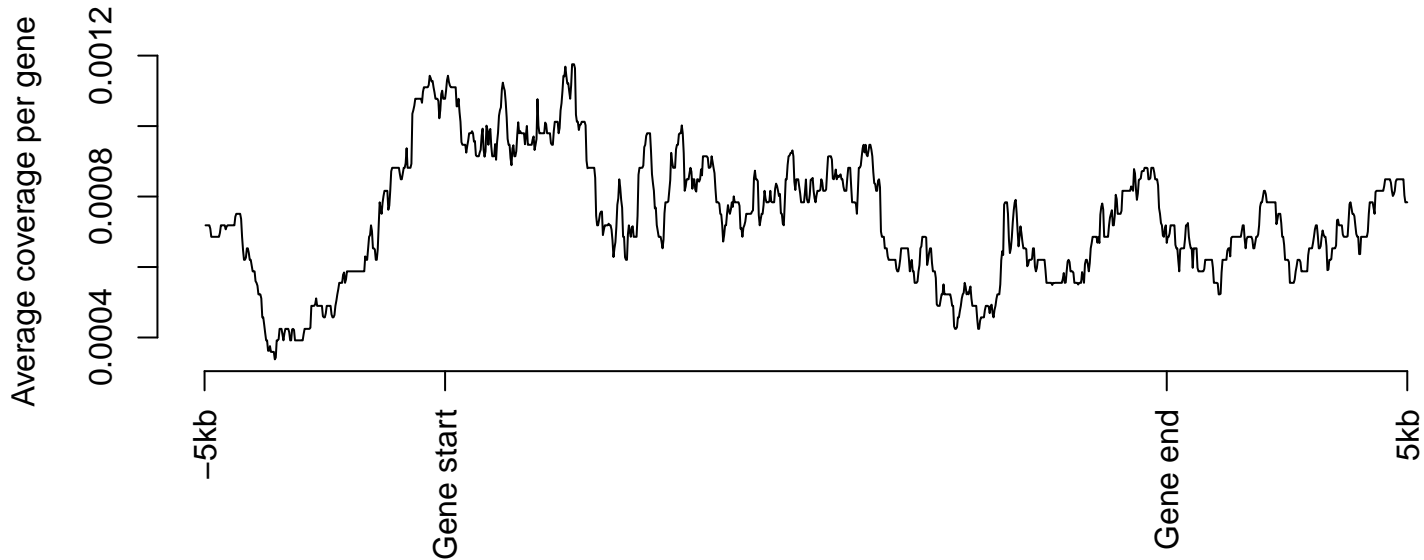
0.0040  
0.0025



# MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc797



# MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc798





# MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc799

Average coverage per gene

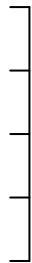
0.0015 0.0025 0.0035

-5kb

Gene start

Gene end

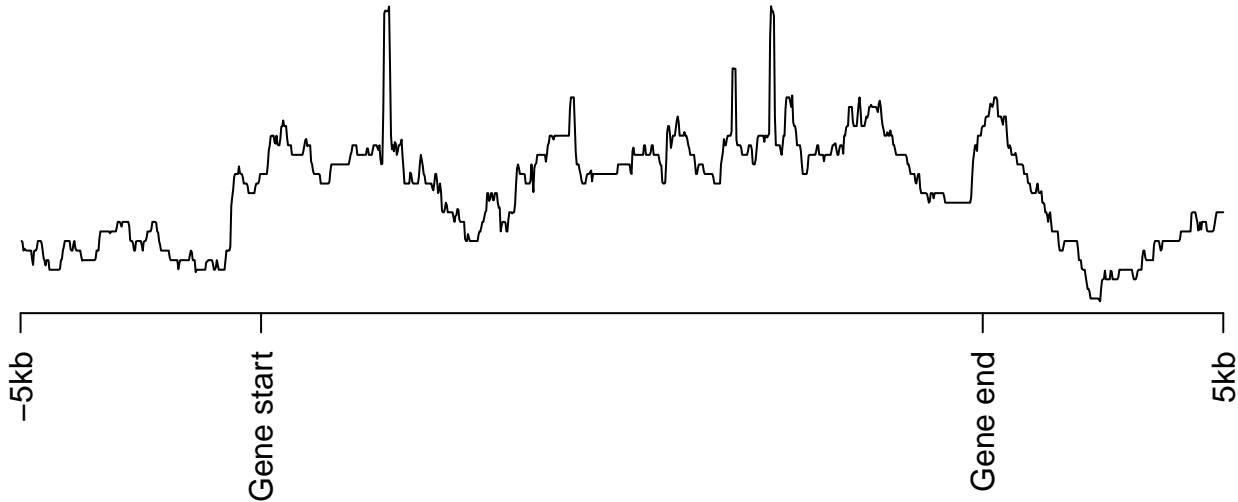
5kb



# MouseNeuronCulture PositiveSingle E.671 scCLTdegenNuc803

Average coverage per gene

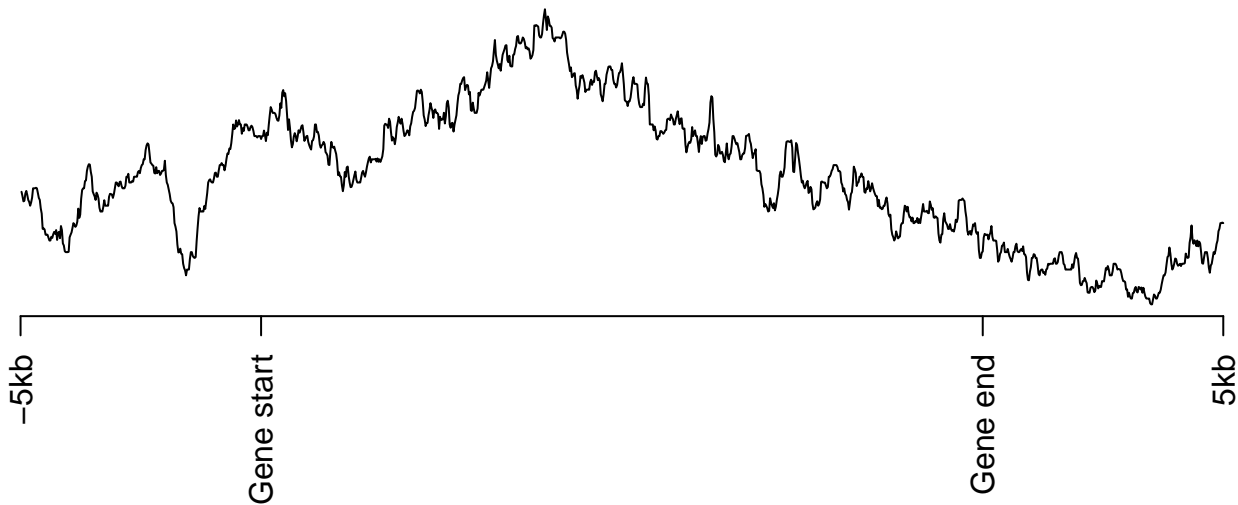
0.0008  
0.0002



# MouseNeuronCulture PositiveSingle E.671 scCLTdegenNuc807

Average coverage per gene

0.0015 0.0025



# MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc965

Average coverage per gene

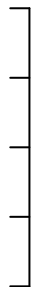
0e+00 2e-04 4e-04

-5kb

Gene start

Gene end

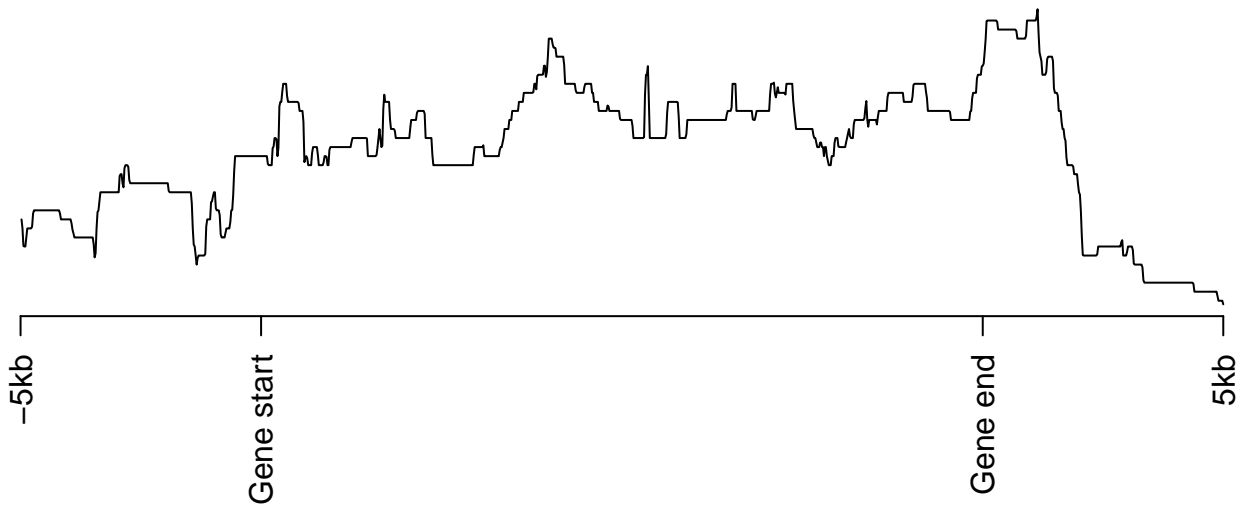
5kb



# MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc966

Average coverage per gene

0e+00 6e-04



# MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc967

Average coverage per gene

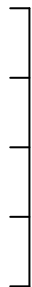
0e+00 4e-04 8e-04

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc968

Average coverage per gene

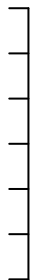
0.00000 0.00020

-5kb

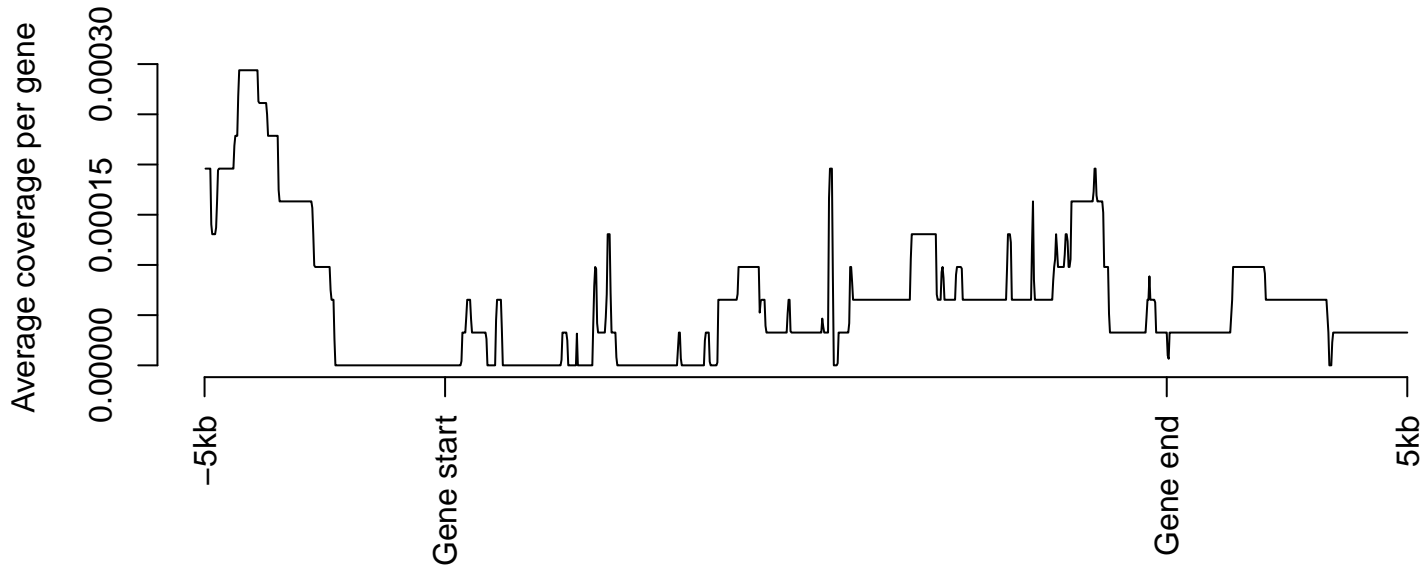
Gene start

Gene end

5kb



# MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc969





# MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc970

Average coverage per gene

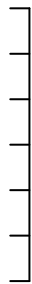
0.0018  
0.0012  
0.0006

-5kb

Gene start

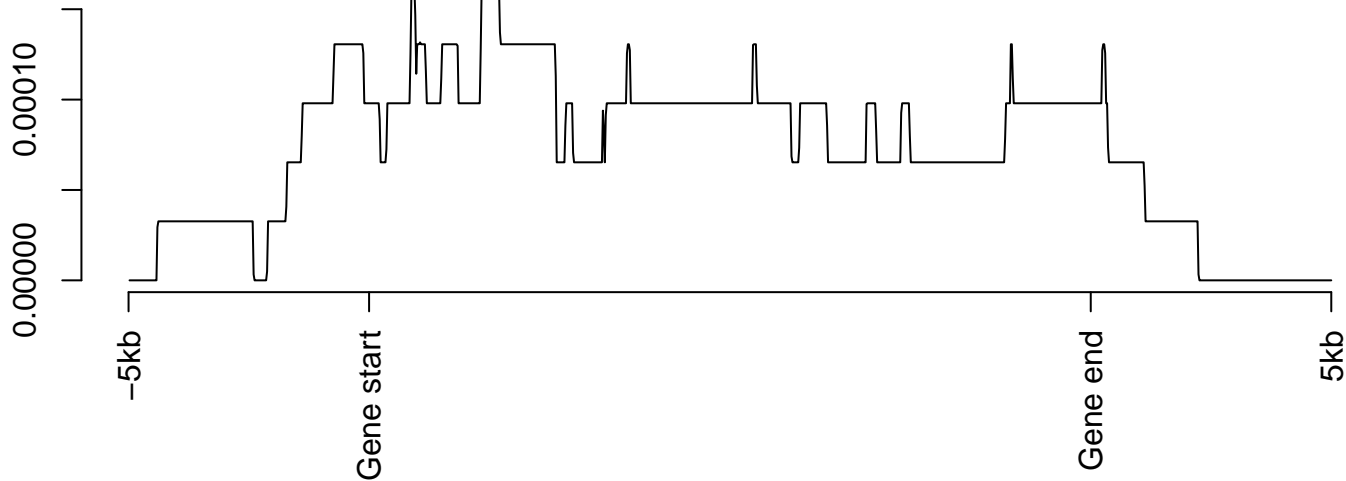
Gene end

5kb



# MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1063

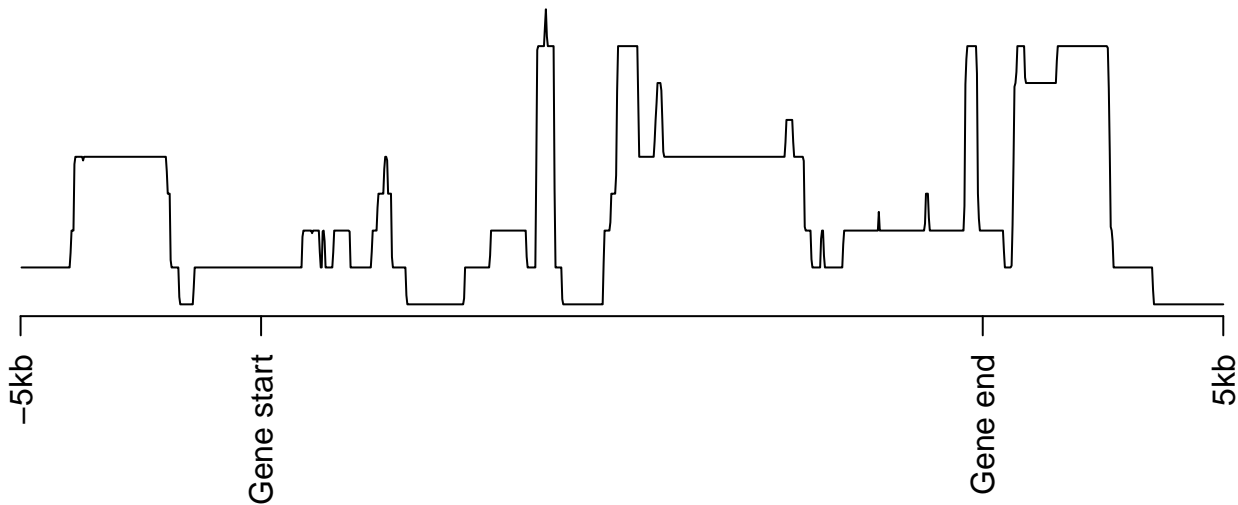
Average coverage per gene



# MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1064

Average coverage per gene

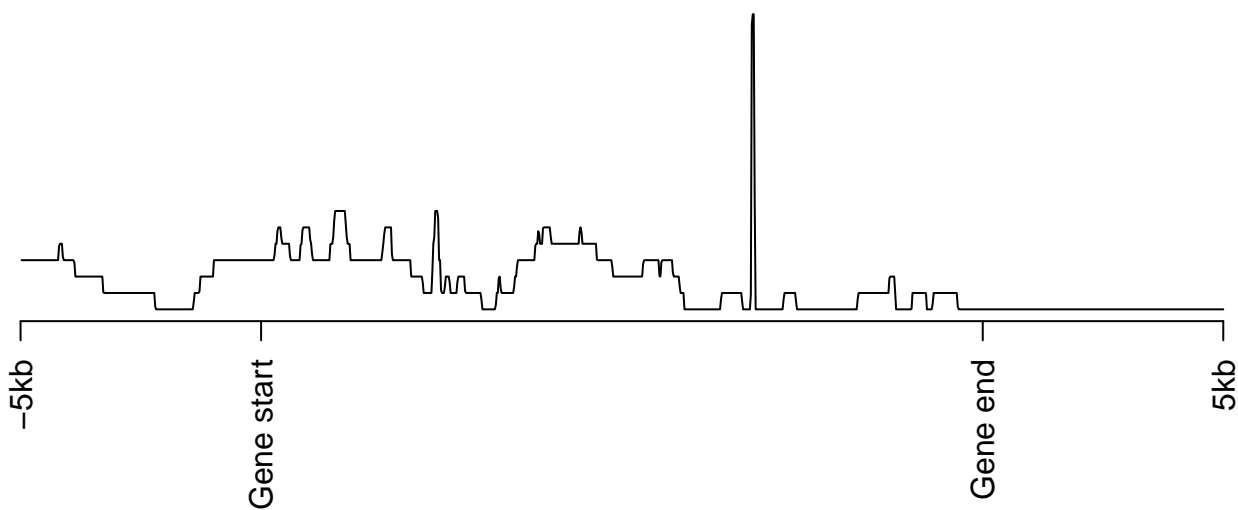
0.00000 0.00015



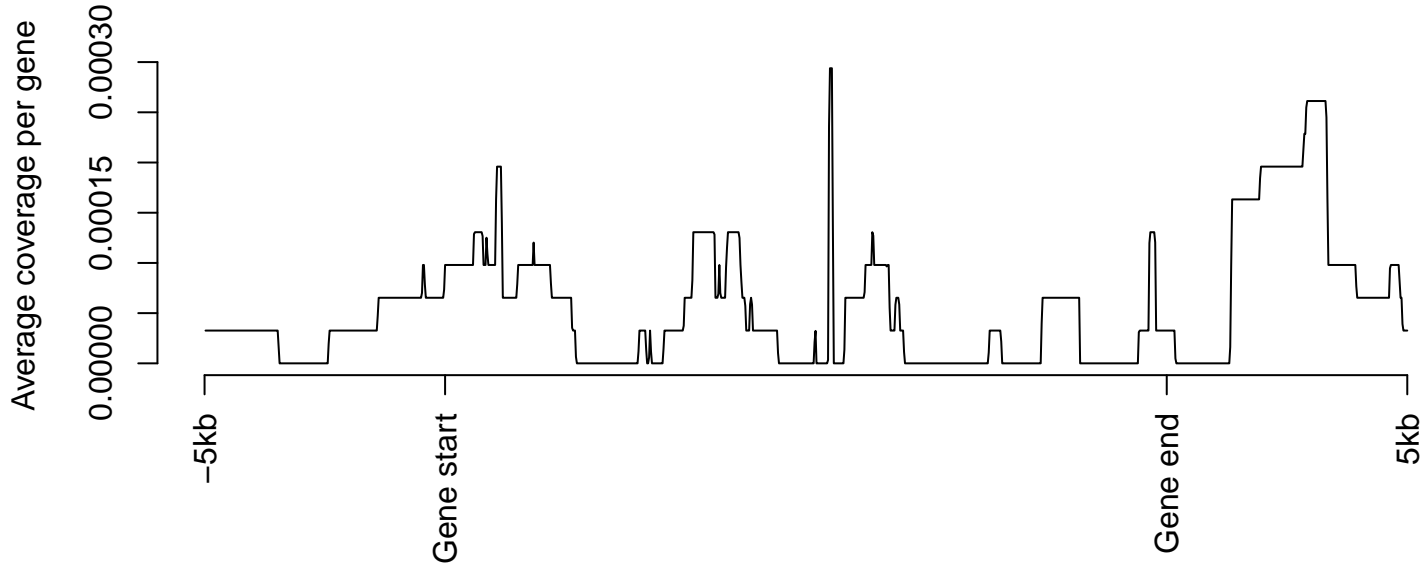
# MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1065

Average coverage per gene

0e+00 3e-04 6e-04

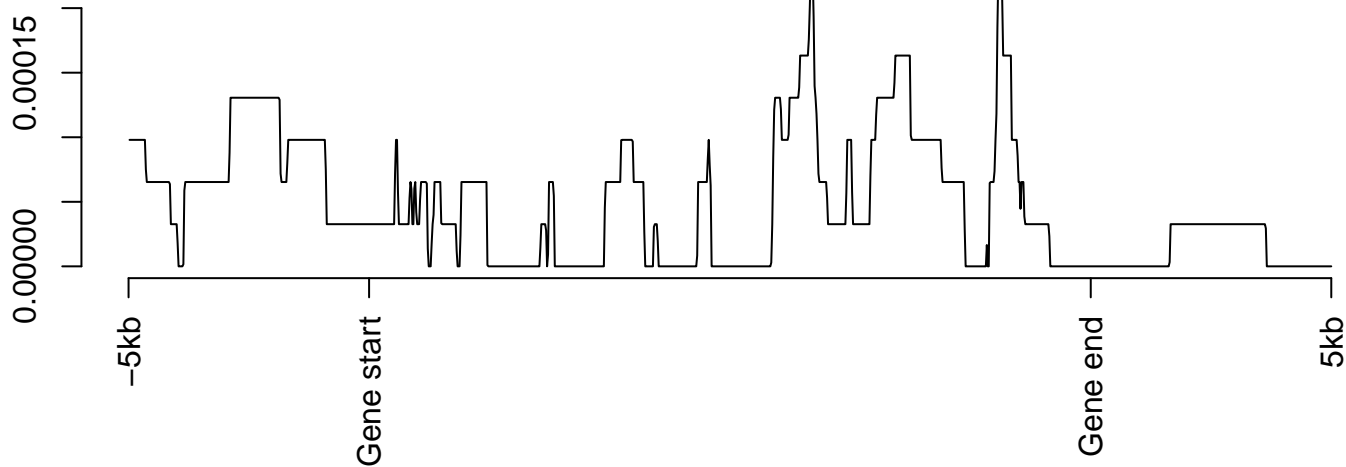


# MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1068



# MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1066

Average coverage per gene



# MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1067

Average coverage per gene

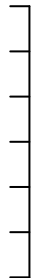
0e+00 3e-05 6e-05

-5kb

Gene start

Gene end

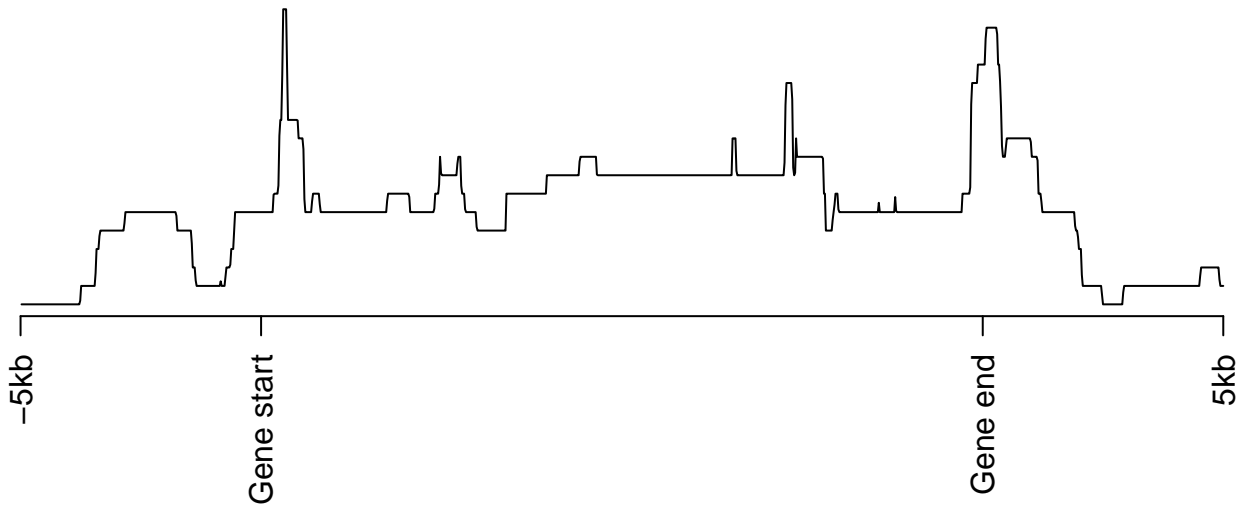
5kb



# MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1095

Average coverage per gene

0e+00 3e-04





# MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1096

Average coverage per gene

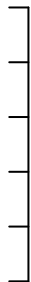
0e+00 6e-04

-5kb

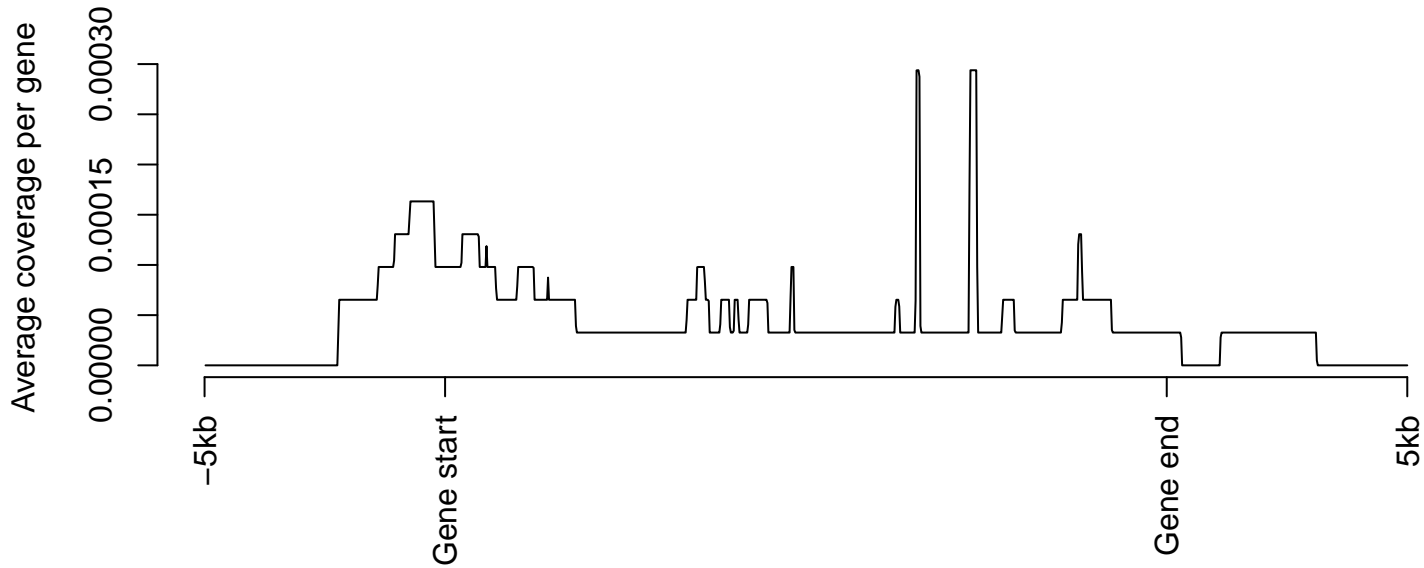
Gene start

Gene end

5kb



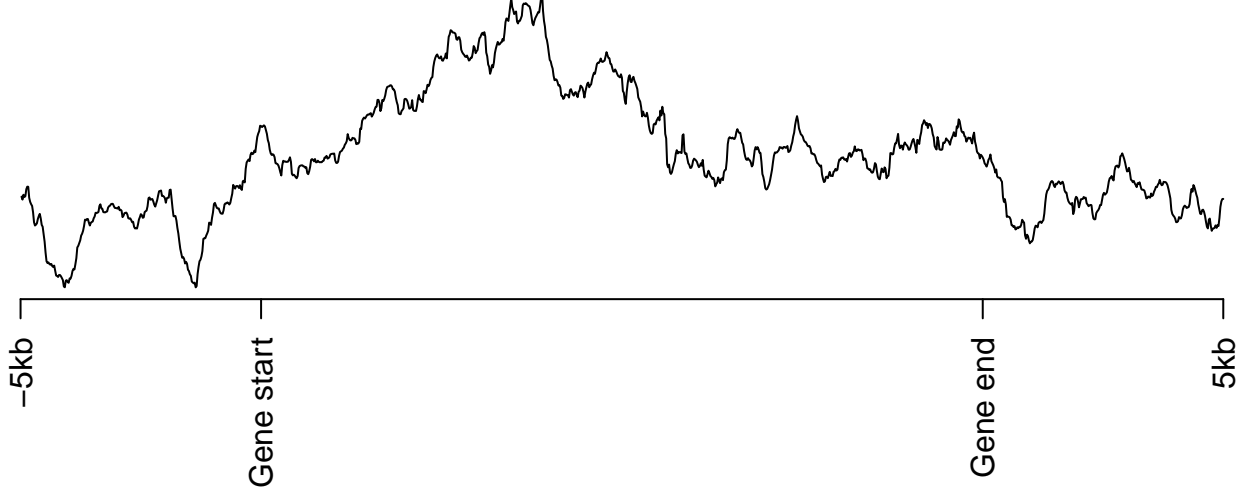
# MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1097



# MouseNeuronCulture PositivePooled E.584 scCLTdegenNuc415

Average coverage per gene

0.07 0.09 0.11



# MouseNeuronCulture PositivePooled E.643 scCLTdegenNuc654

Average coverage per gene

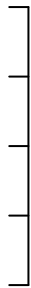
0.10  
0.08  
0.06

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositivePooled E.669 scCLTdegenNuc809

Average coverage per gene

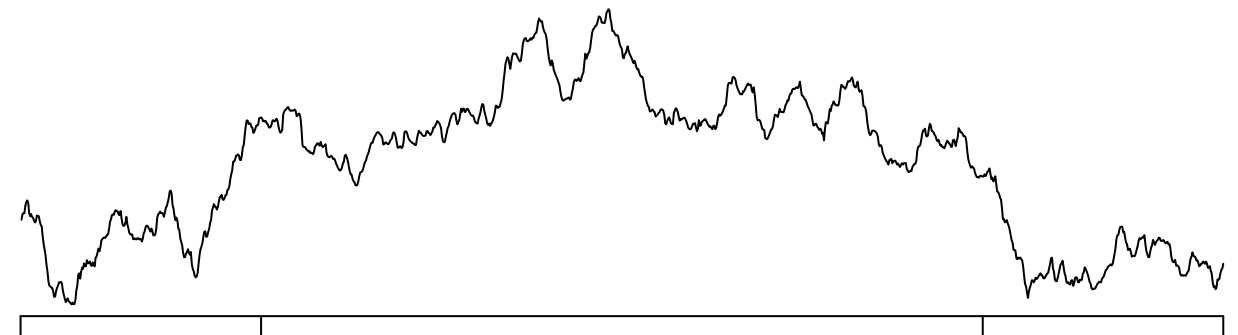
0.40  
0.30

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveMerged E.– MouseNeuronCulturePositiveMerged

Average coverage per gene

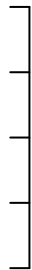
0.12 0.16 0.20

-5kb

Gene start

Gene end

5kb



# MouseNeuronCulture PositiveAll E.– MouseNeuronCulturePositiveAll

Average coverage per gene

0.55 0.65 0.75 0.85

-5kb

Gene start

Gene end

5kb

