

MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc429

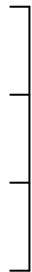
Average coverage per gene

0.00015
0.00005

-5kb

TSS

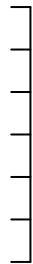
5kb



MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc430

Average coverage per gene

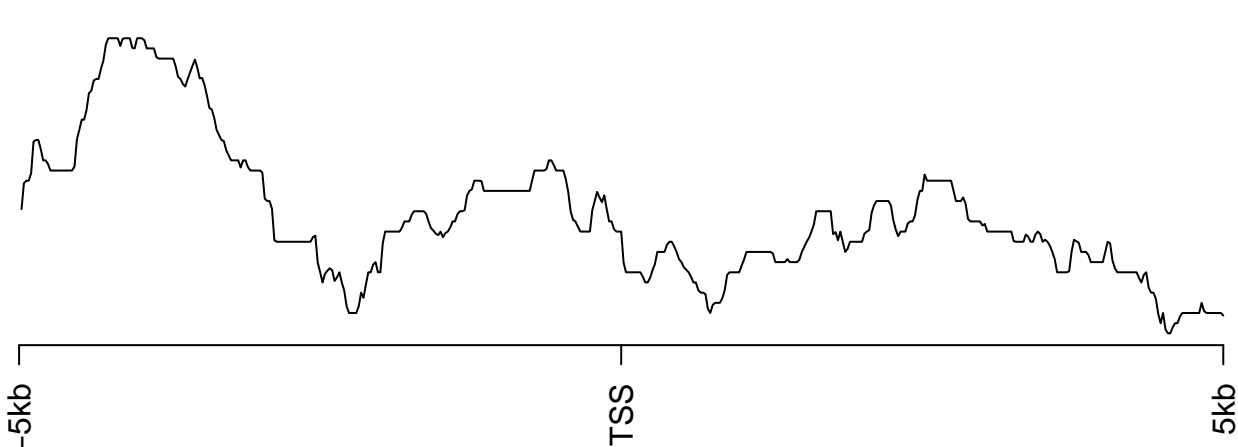
1e-03
7e-04
4e-04



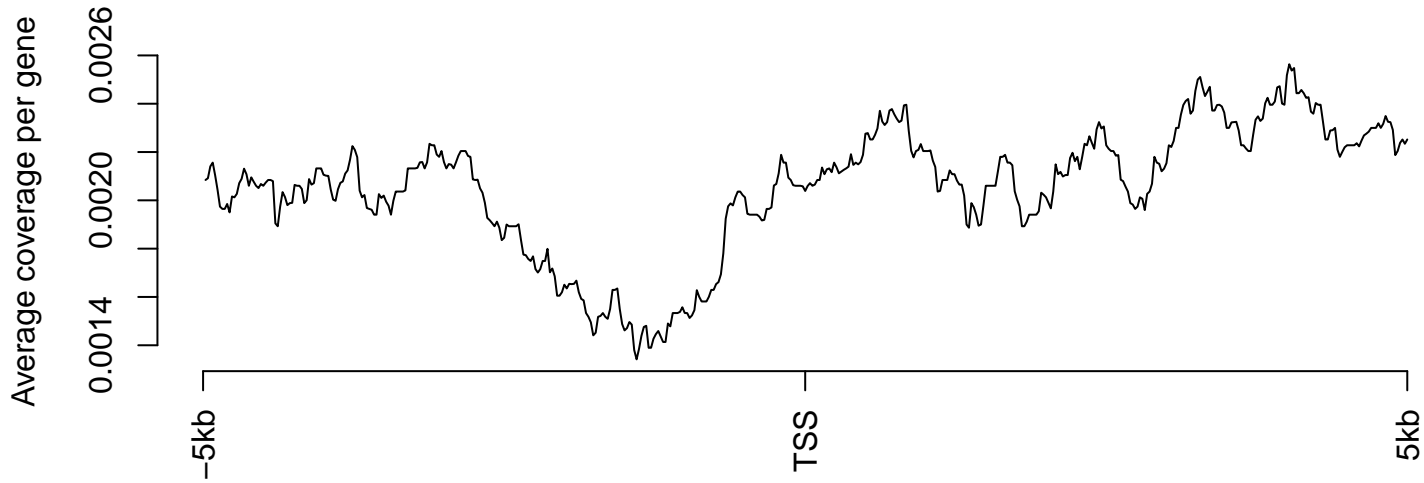
-5kb

TSS

5kb



MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc431



MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc433

Average coverage per gene

0.00010 0.00020

-5kb

TSS

5kb



MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc886

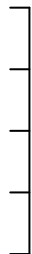
Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc891

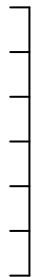
Average coverage per gene

2e-04 5e-04 8e-04

-5kb

TSS

5kb



MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc892

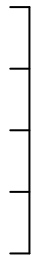
Average coverage per gene

2e-04 4e-04 6e-04

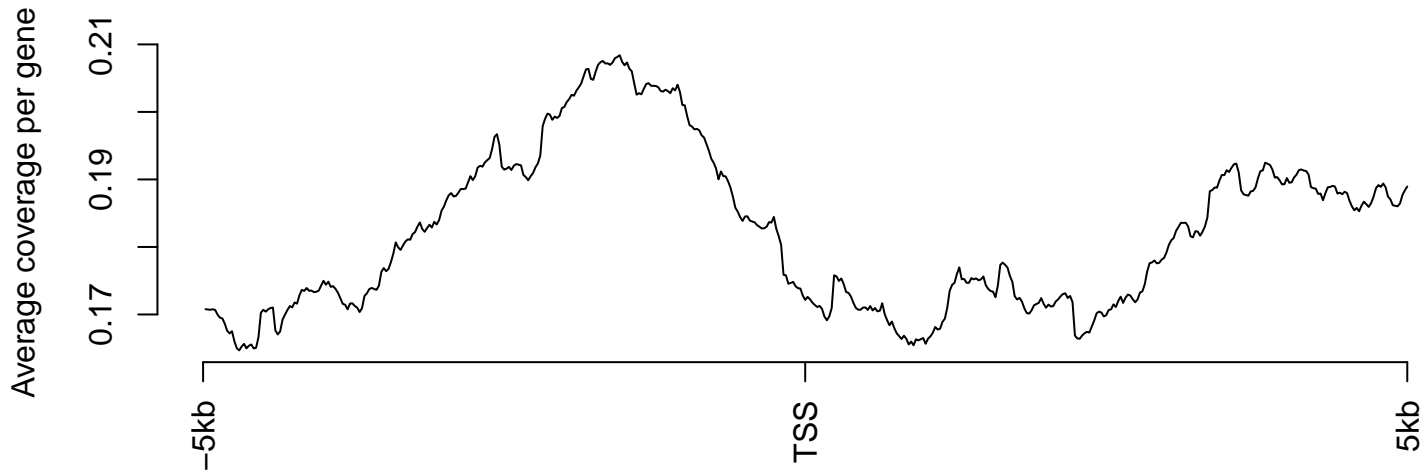
-5kb

TSS

5kb



MouseNeuronCulture NegCtrlPooled E.597 scCLTdegenNuc471



MouseNeuronCulture NegCtrlMerged E.– MouseNeuronCultureNegCtrlMerged

Average coverage per gene

0.0036 0.0042

-5kb

TSS

5kb



MouseNeuronCulture NegCtrlAll E. – MouseNeuronCultureNegCtrlAll

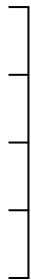
Average coverage per gene

0.17 0.19 0.21

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc387

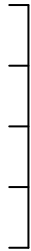
Average coverage per gene

0.0025 0.0040

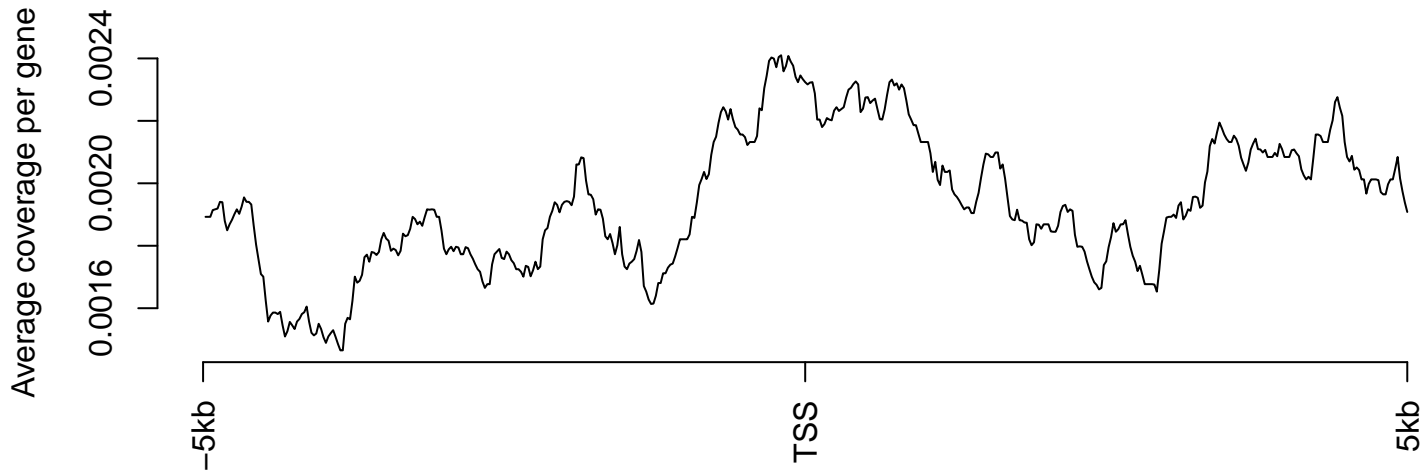
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc388



MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc390

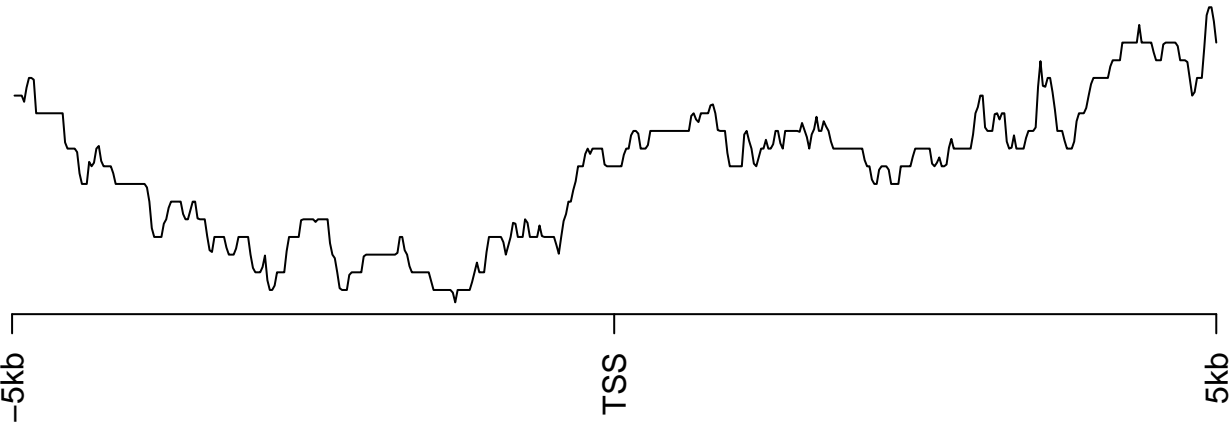
Average coverage per gene

3e-04
5e-04

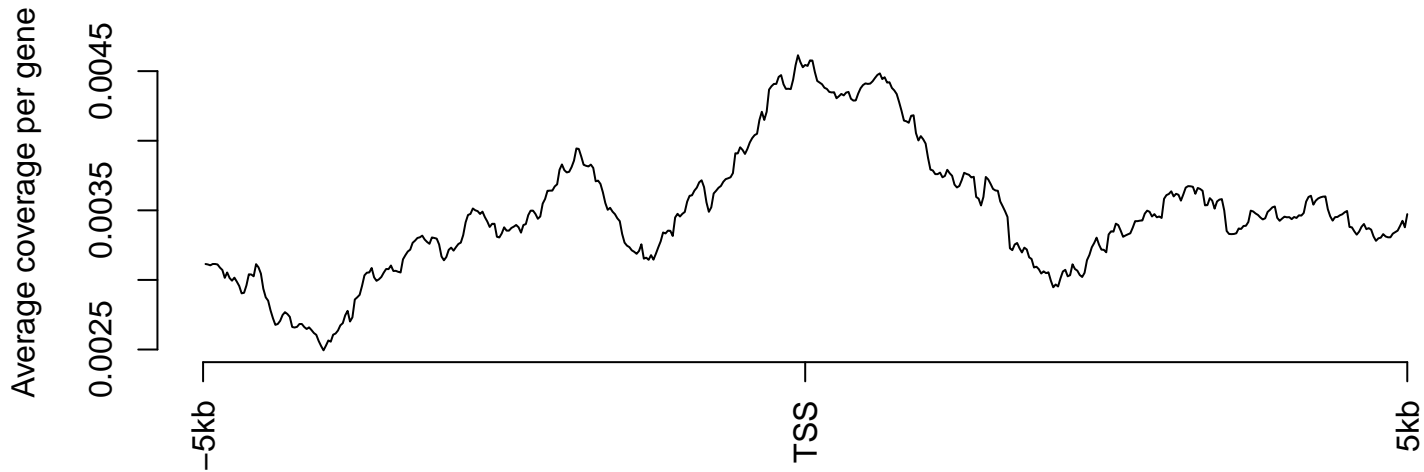
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc391



MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc392

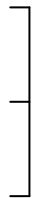
Average coverage per gene

0.0030
0.0020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc393

Average coverage per gene

0.0015 0.0025

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc501

Average coverage per gene

4e-04
6e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc502

Average coverage per gene

0.0002 0.0008

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc503

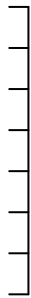
Average coverage per gene

0.0004 0.0008

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc504

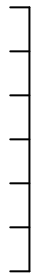
Average coverage per gene

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

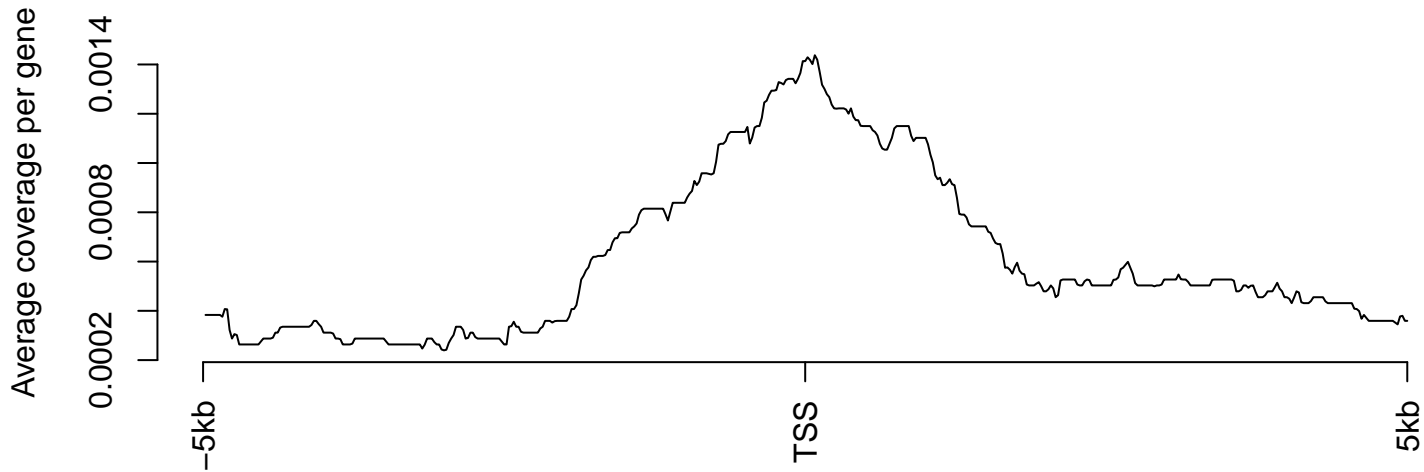
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc505



MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc510

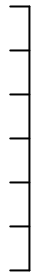
Average coverage per gene

0.00025
0.00005

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc511

Average coverage per gene

0.0015 0.0030 0.0045

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc512

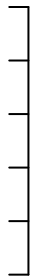
Average coverage per gene

0.0015 0.0030

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc513

Average coverage per gene

0.0015 0.0030

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc521

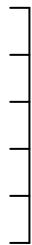
Average coverage per gene

0.00030
0.00010

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc522

Average coverage per gene

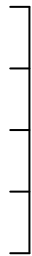
0.00025

0.00010

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc523

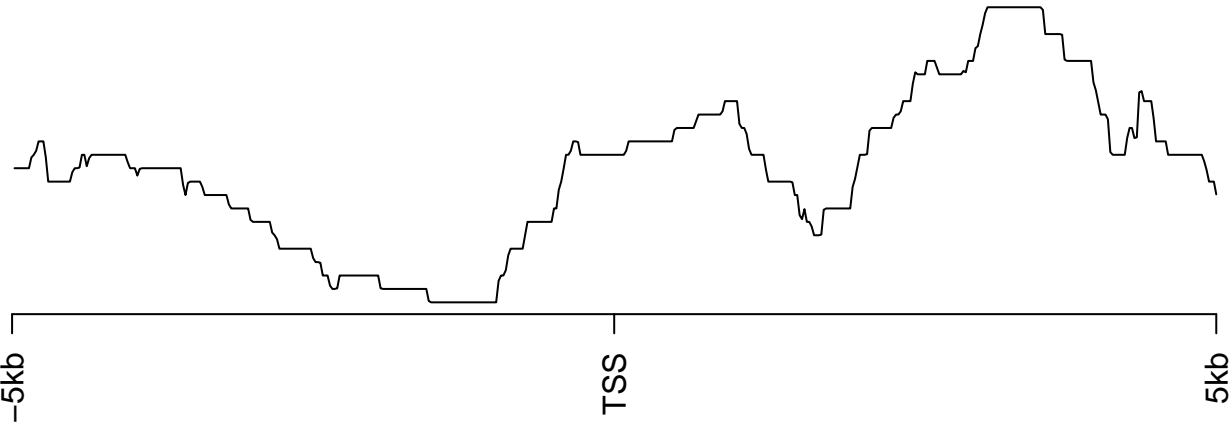
Average coverage per gene

4e-04
1e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc524

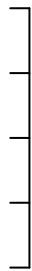
Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc525

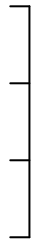
Average coverage per gene

0.00005 0.00015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc526

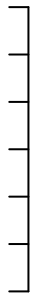
Average coverage per gene

0.00030
0.00010

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc527

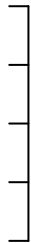
Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc528

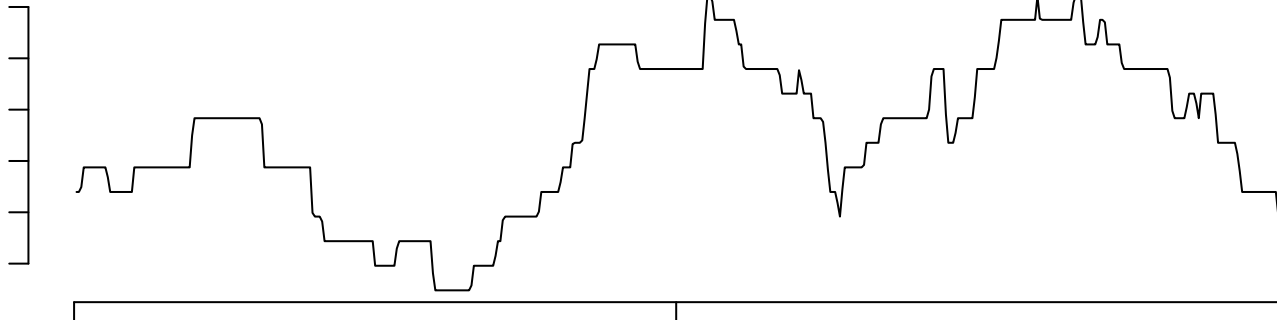
Average coverage per gene

0.00005 0.00020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc529

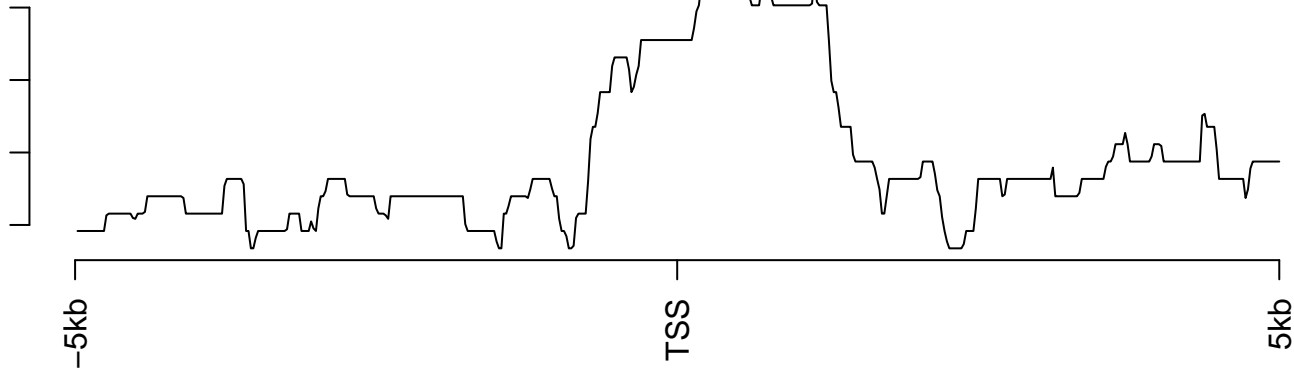
Average coverage per gene

2e-04
4e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc530

Average coverage per gene

0.0008
0.0000

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc531

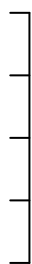
Average coverage per gene

0.0010 0.0014 0.0018

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc532

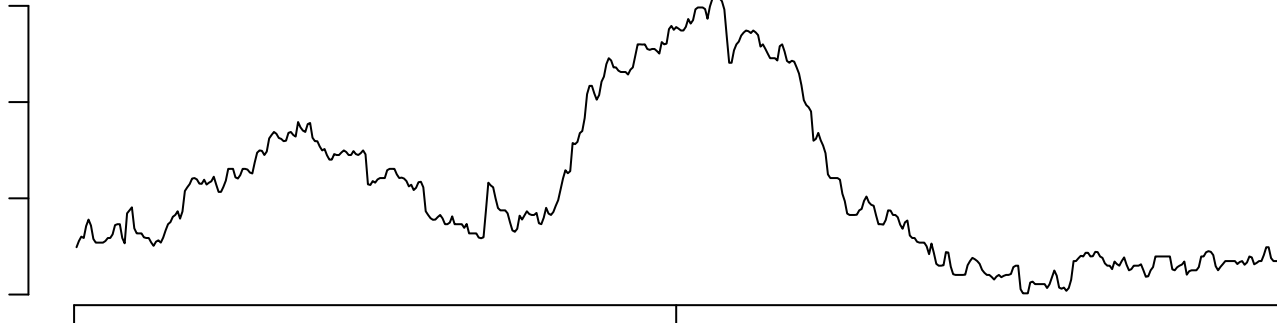
Average coverage per gene

0.0020
0.0010

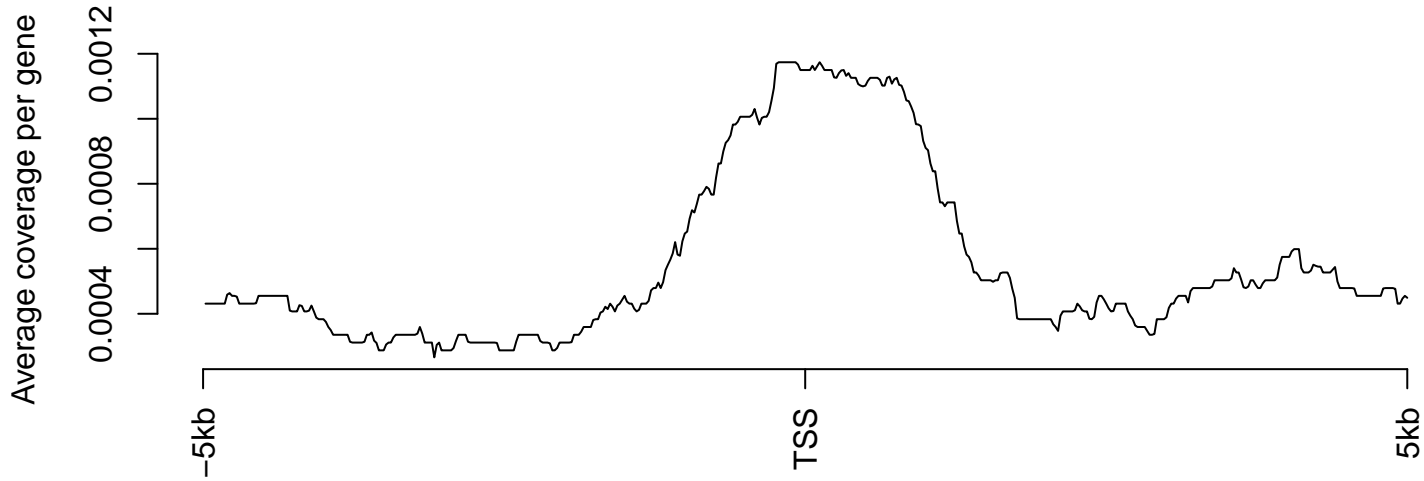
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc533



MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc534

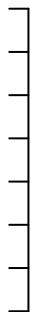
Average coverage per gene

0.0012 0.0018 0.0024

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc535

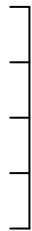
Average coverage per gene

0.006 0.008 0.010

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc536

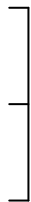
Average coverage per gene

0.0015
0.0005

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc537

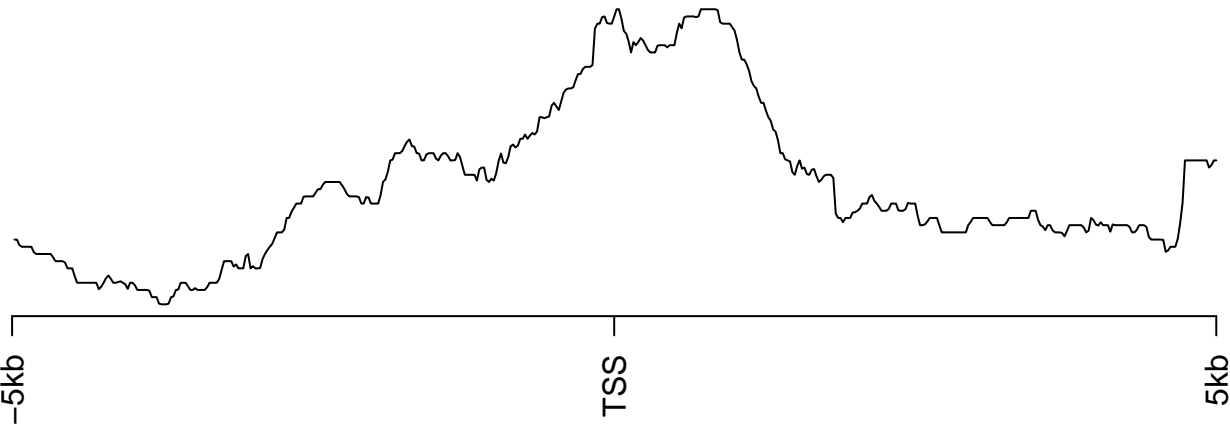
Average coverage per gene

0.0010
0.0004

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc542

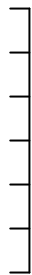
Average coverage per gene

0.00025
0.00005

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc543

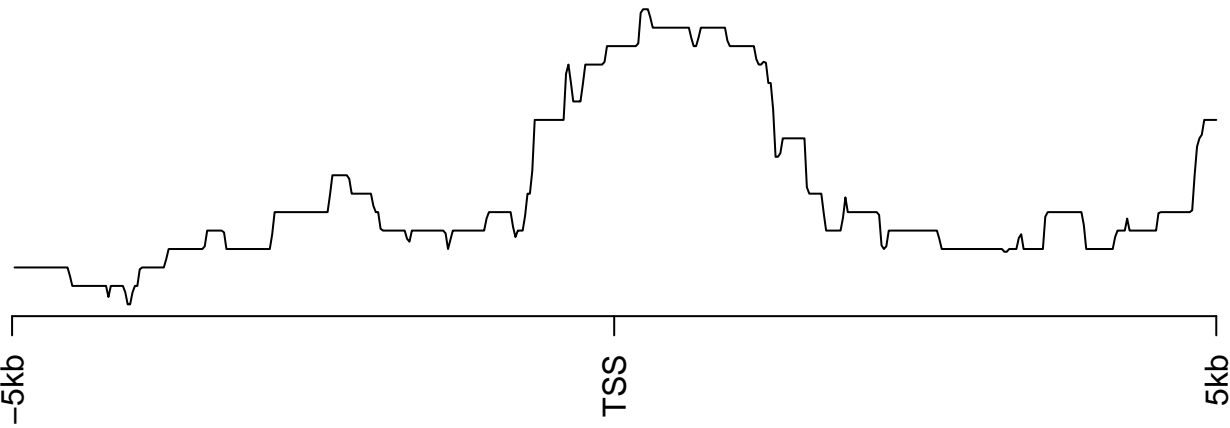
Average coverage per gene

1e-04
3e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc544

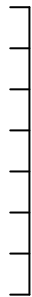
Average coverage per gene

0.00025
0.00005

-5kb

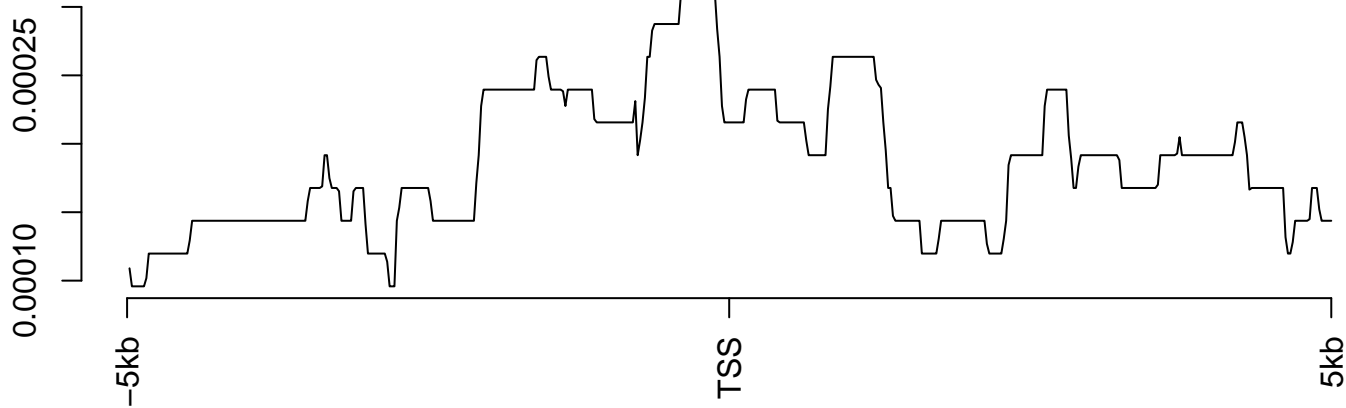
TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc545

Average coverage per gene



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc546

Average coverage per gene

1e-04 3e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc547

Average coverage per gene

0.00000 0.00020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc548

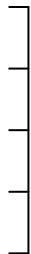
Average coverage per gene

0.00000
0.00015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc549

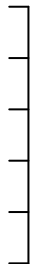
Average coverage per gene

0.00000 0.00015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc550

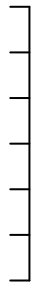
Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc551

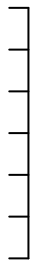
Average coverage per gene

0.0008 0.0014 0.0020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc552

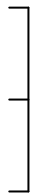
Average coverage per gene

0.0015
0.0005

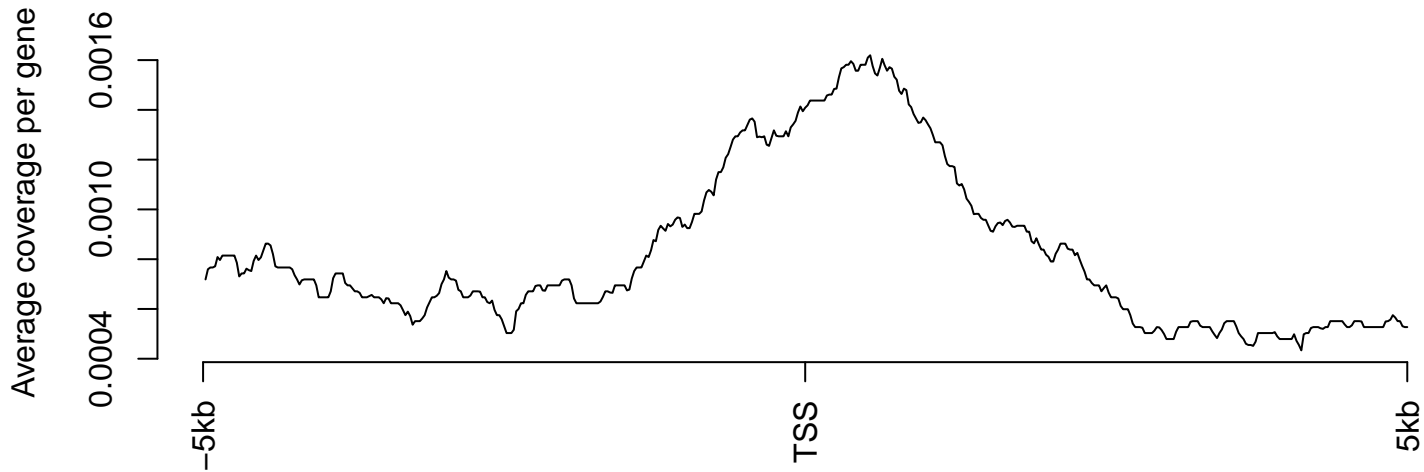
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc553



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc554

Average coverage per gene

0.0004 0.0010

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc555

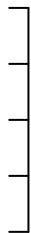
Average coverage per gene

0.0010
0.0004

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc556

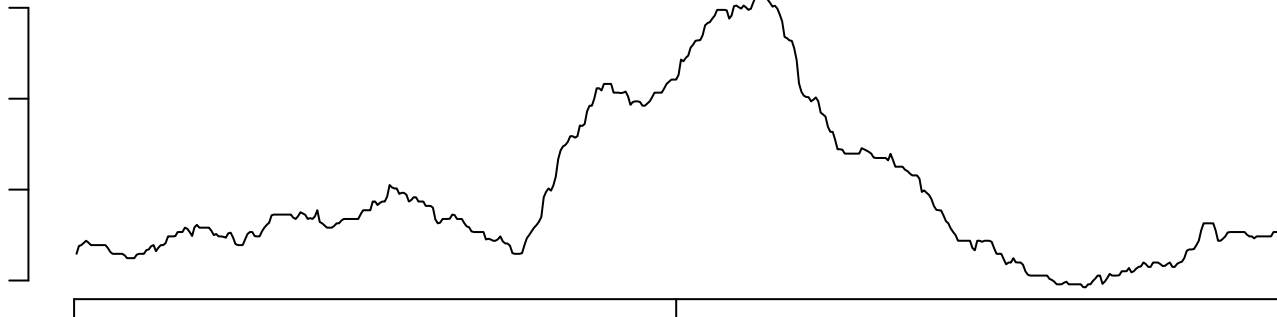
Average coverage per gene

0.0005
0.0015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc557

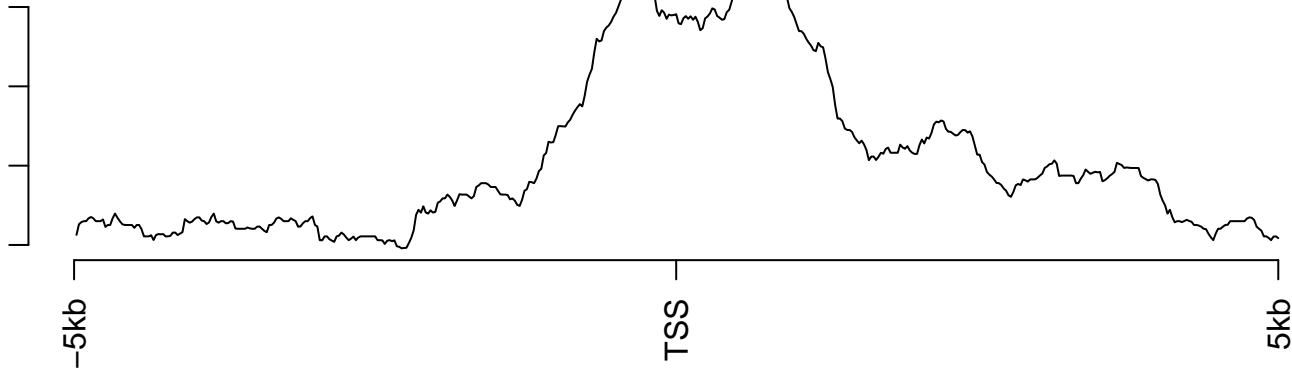
Average coverage per gene

0.0010 0.0020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc558

Average coverage per gene

0.0010 0.0016

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc559

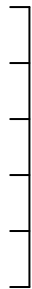
Average coverage per gene

0.00005 0.00020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc561

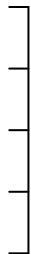
Average coverage per gene

0.00000 0.00015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc562

Average coverage per gene

0.00005 0.00015

-5kb

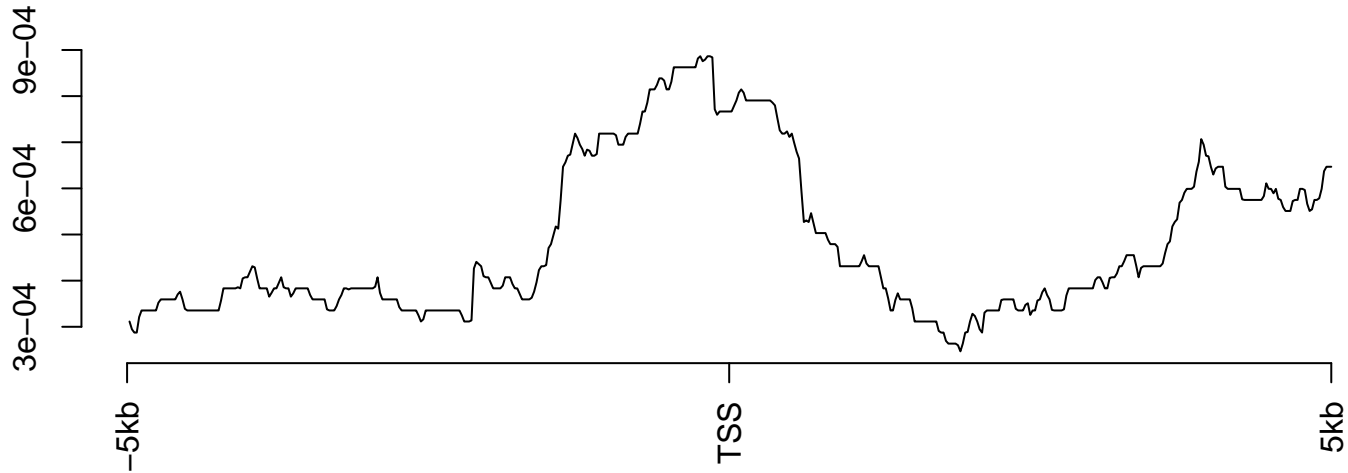
TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc563

Average coverage per gene



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc564

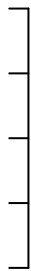
Average coverage per gene

2e-04 4e-04 6e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc565

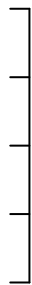
Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc566

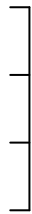
Average coverage per gene

0.00025
0.00010

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc567

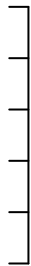
Average coverage per gene

0.00010 0.00025

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc568

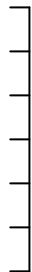
Average coverage per gene

0.00000
0.00020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc569

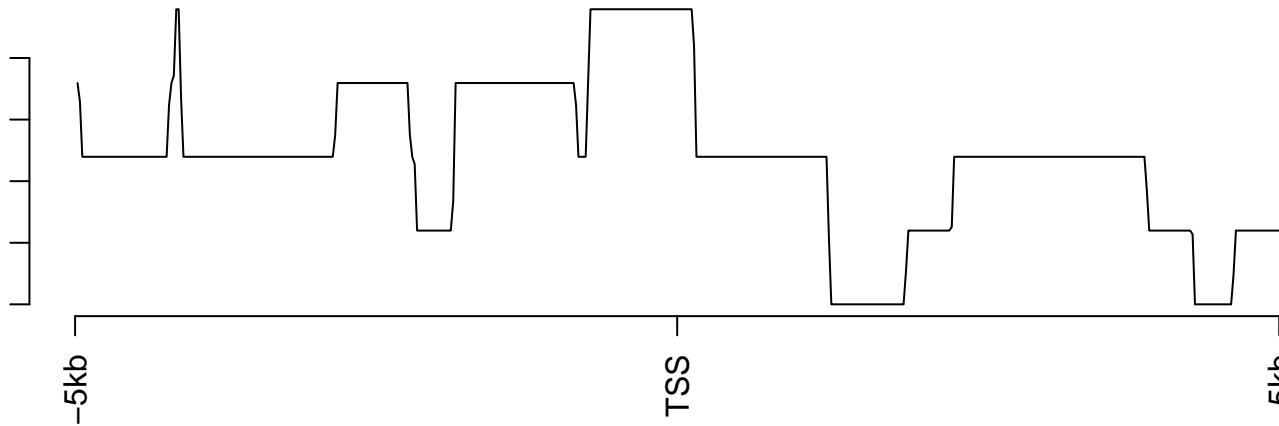
Average coverage per gene

0e+00 4e-05 8e-05

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc570

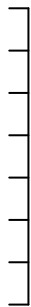
Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc571

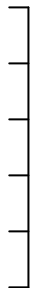
Average coverage per gene

0.00000 0.00015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc572

Average coverage per gene

0.00000 0.00015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc573

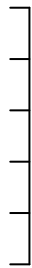
Average coverage per gene

0.00015 0.00030

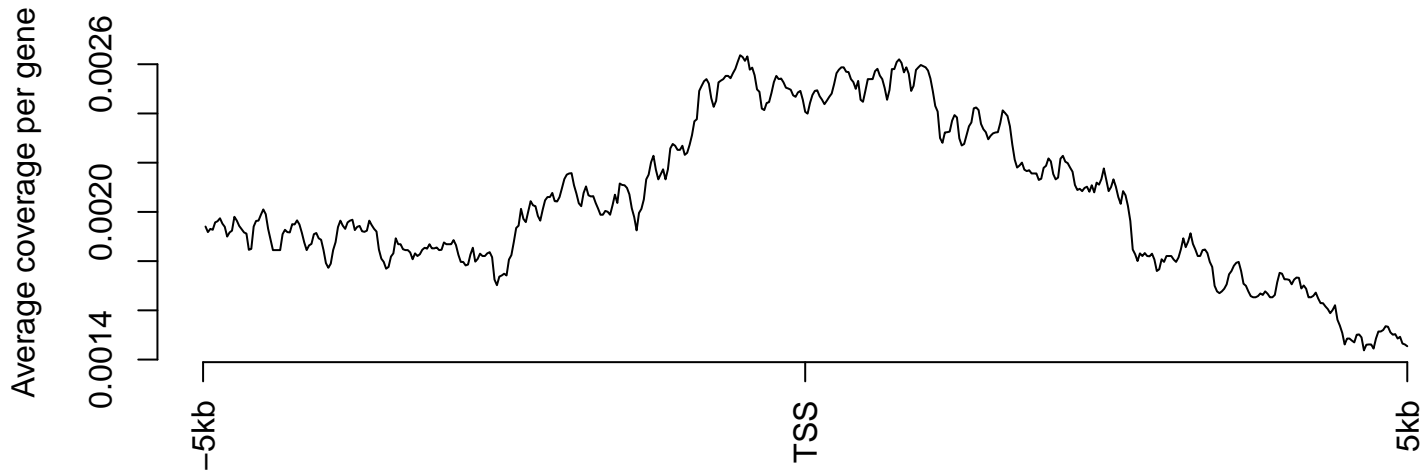
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc576



MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc601

Average coverage per gene

0.0018 0.0026

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc602

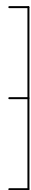
Average coverage per gene

0.0025 0.0035

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc577b

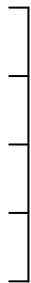
Average coverage per gene

0.00015
0.00000

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc578b

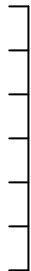
Average coverage per gene

0.00000 0.00020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc579b

Average coverage per gene

0.00000 0.00015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc580b

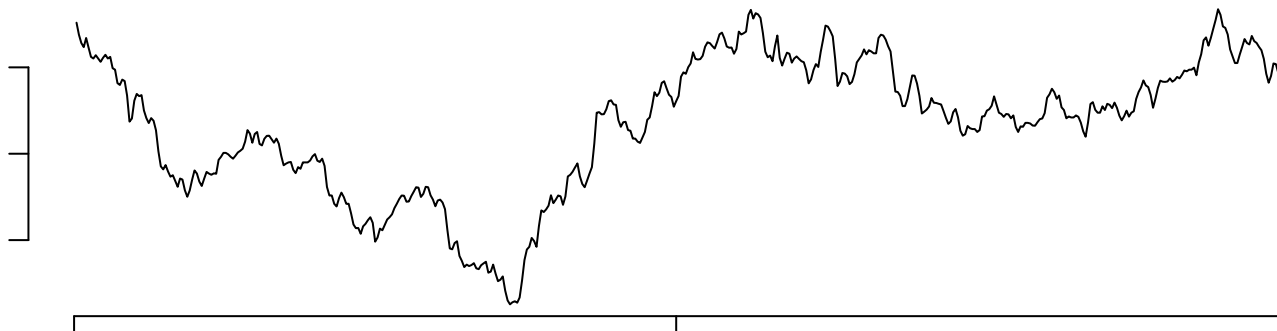
Average coverage per gene

0.0040
0.0030

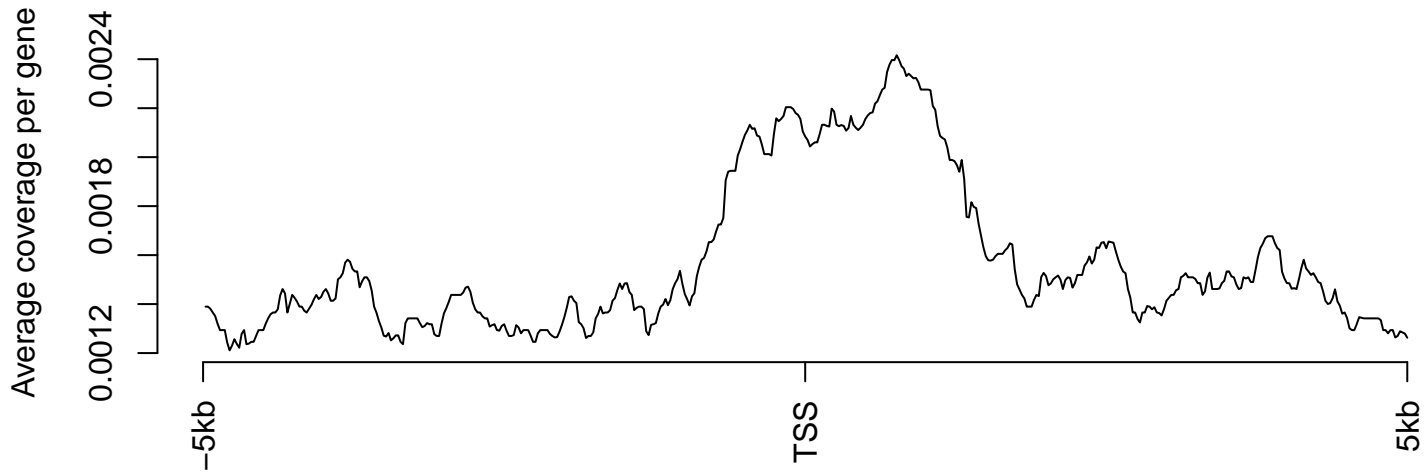
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc581b



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc582b

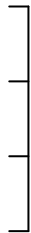
Average coverage per gene

0.0035 0.0045

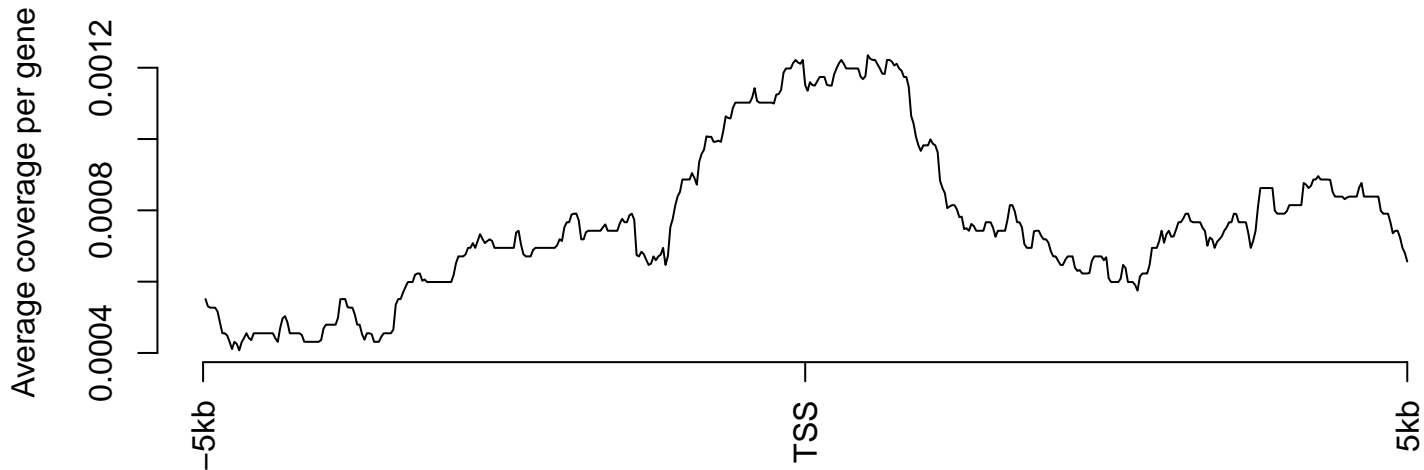
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc583b



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc584b

Average coverage per gene

4e-04
8e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc585b

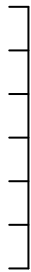
Average coverage per gene

0.0009 0.0012 0.0015

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc586b



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc587b

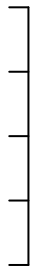
Average coverage per gene

0.0012 0.0016 0.0020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc588b

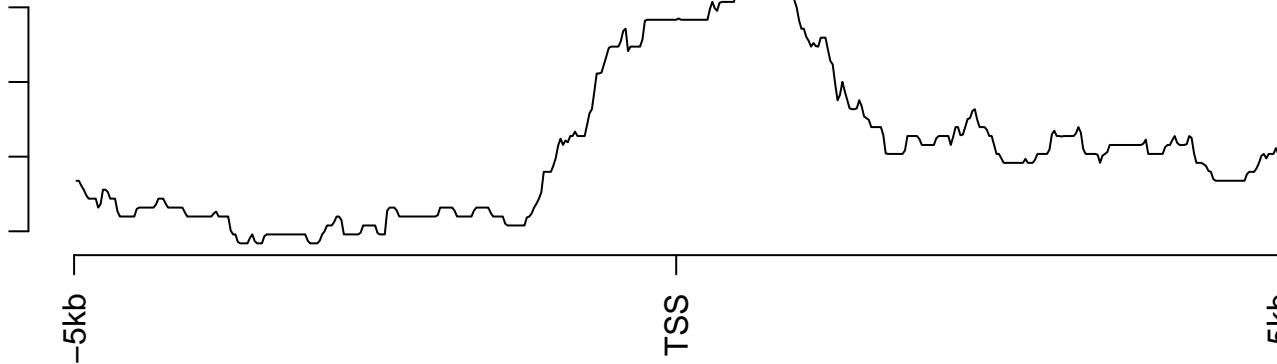
Average coverage per gene

2e-04
6e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc589b

Average coverage per gene

3e-04

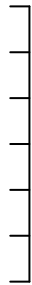
6e-04

9e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc590b

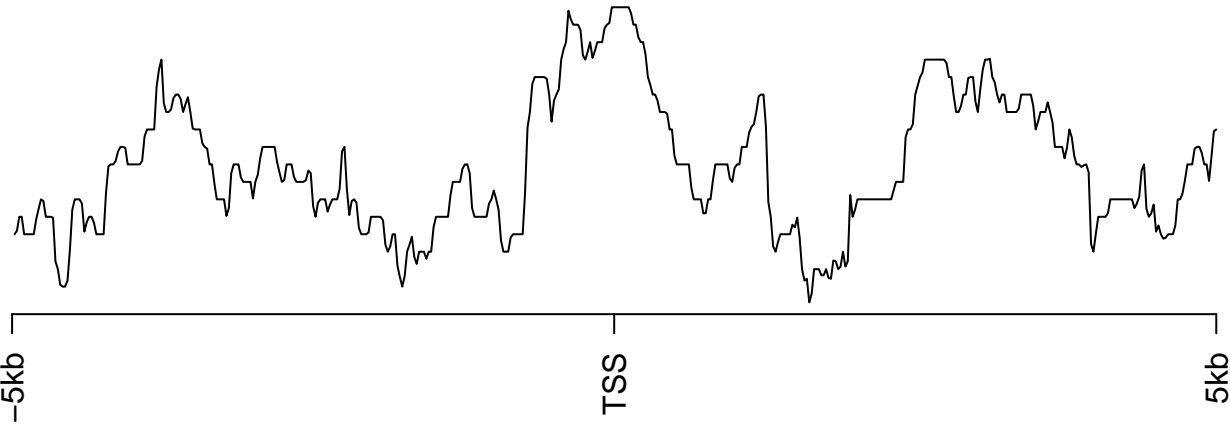
Average coverage per gene

6e-04 8e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc591b

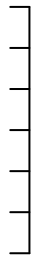
Average coverage per gene

0.00030
0.00010

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc592b

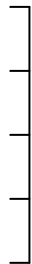
Average coverage per gene

4e-04 6e-04 8e-04

-5kb

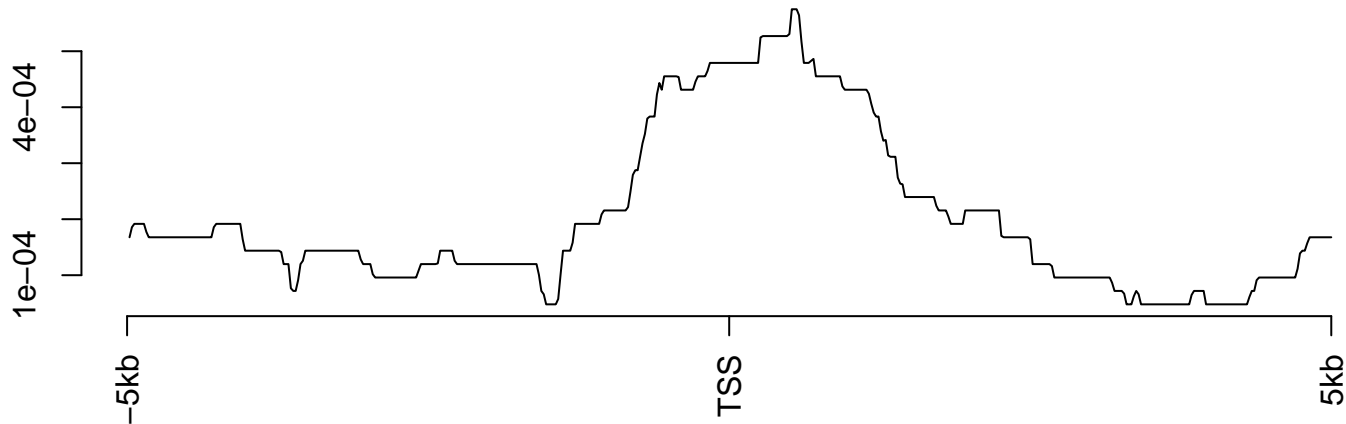
TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc593b

Average coverage per gene



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc594b

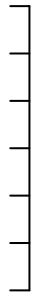
Average coverage per gene

0.00005
0.00025

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc595b

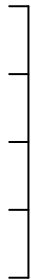
Average coverage per gene

0.0024 0.0028 0.0032

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc596b

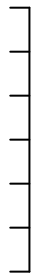
Average coverage per gene

0.0005 0.0008 0.0011

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc597b

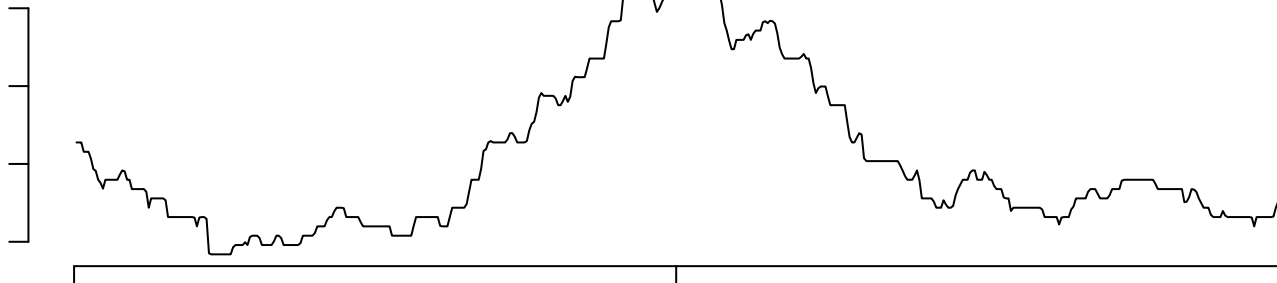
Average coverage per gene

2e-04
6e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc598b

Average coverage per gene

0.0008 0.0012

-5kb

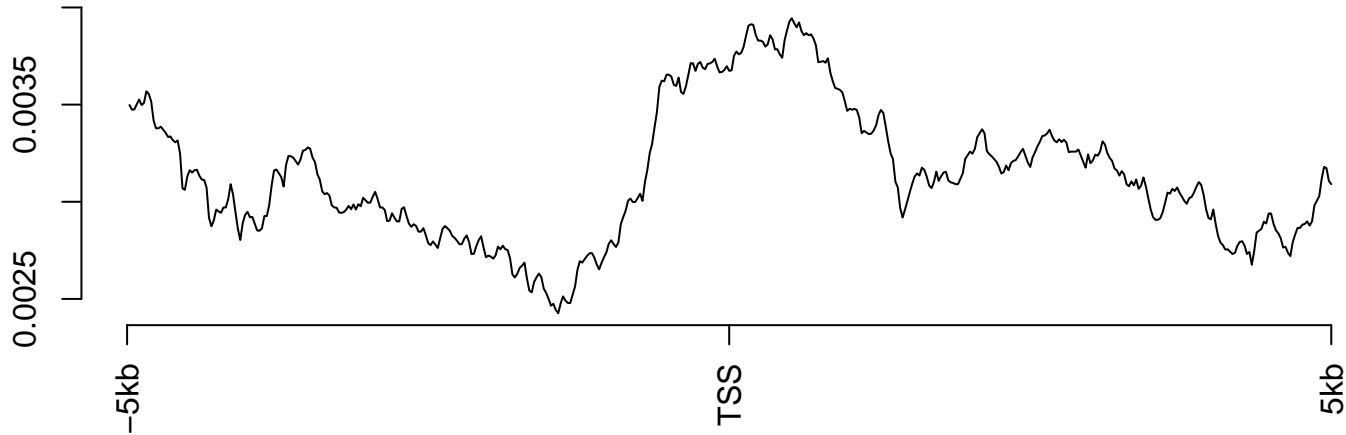
TSS

5kb



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc599b

Average coverage per gene



MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc600b

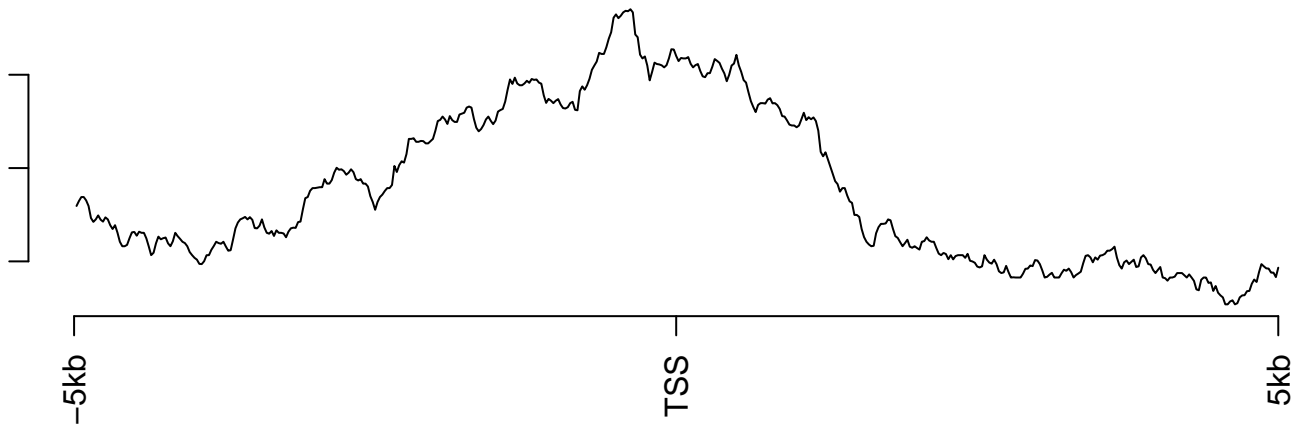
Average coverage per gene

0.0015 0.0025

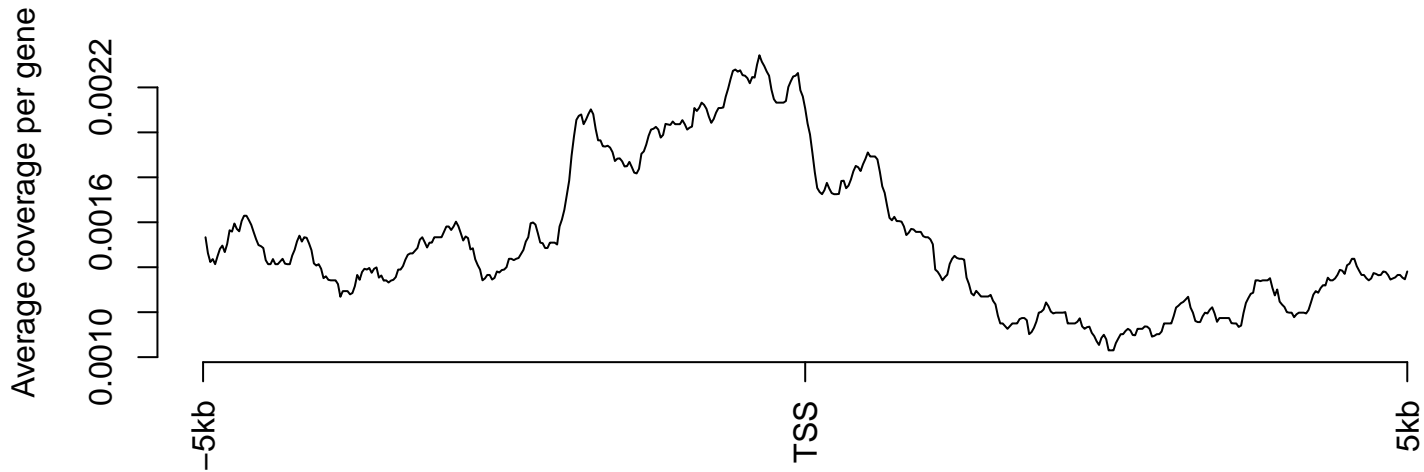
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc779



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc780

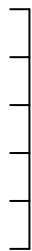
Average coverage per gene

0.0014 0.0020

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc781

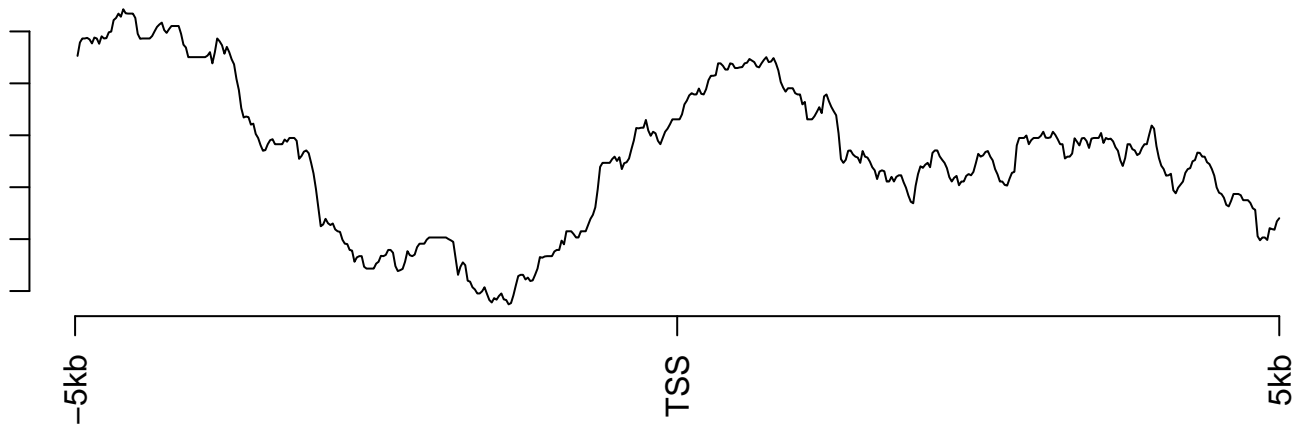
Average coverage per gene

0.0014
0.0008

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc782

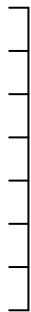
Average coverage per gene

0.0014 0.0020 0.0026

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc783

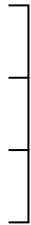
Average coverage per gene

0.0015 0.0025

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc784

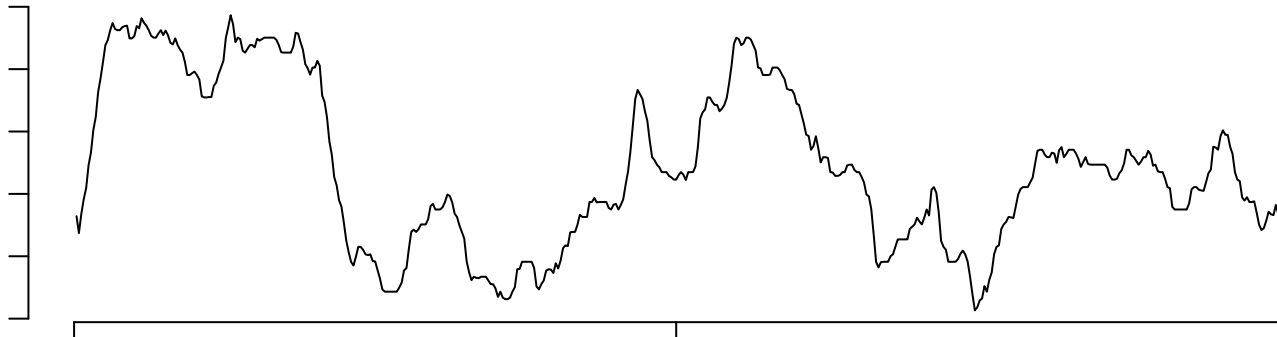
Average coverage per gene

0.0008 0.0012 0.0016

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc785

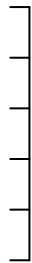
Average coverage per gene

0.0016 0.0022

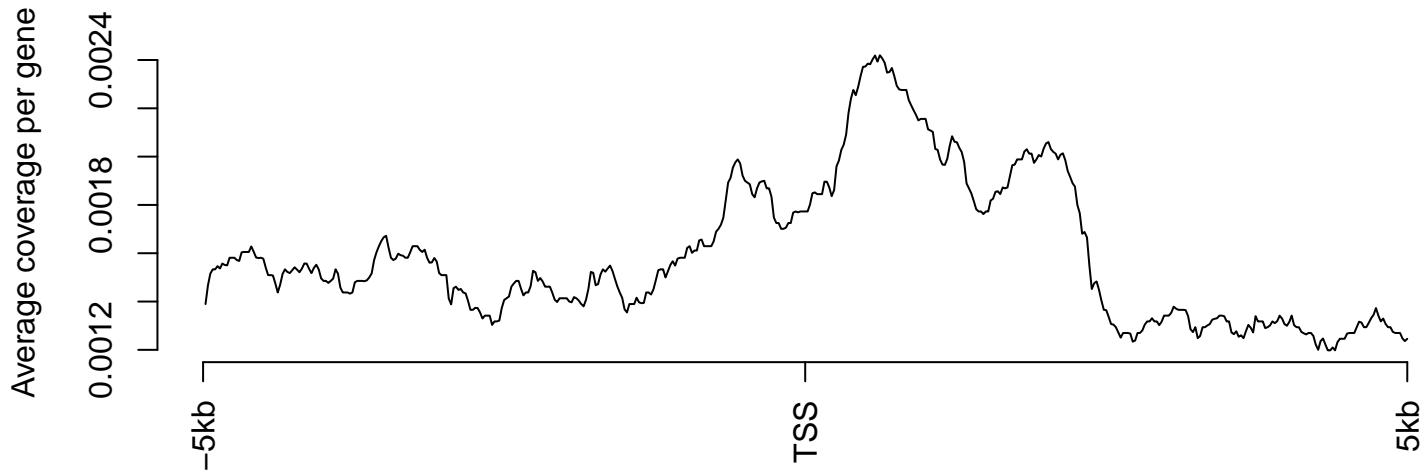
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc786



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc787

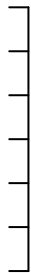
Average coverage per gene

0.0012 0.0018 0.0024

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc788

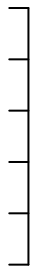
Average coverage per gene

3e-04 6e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc789

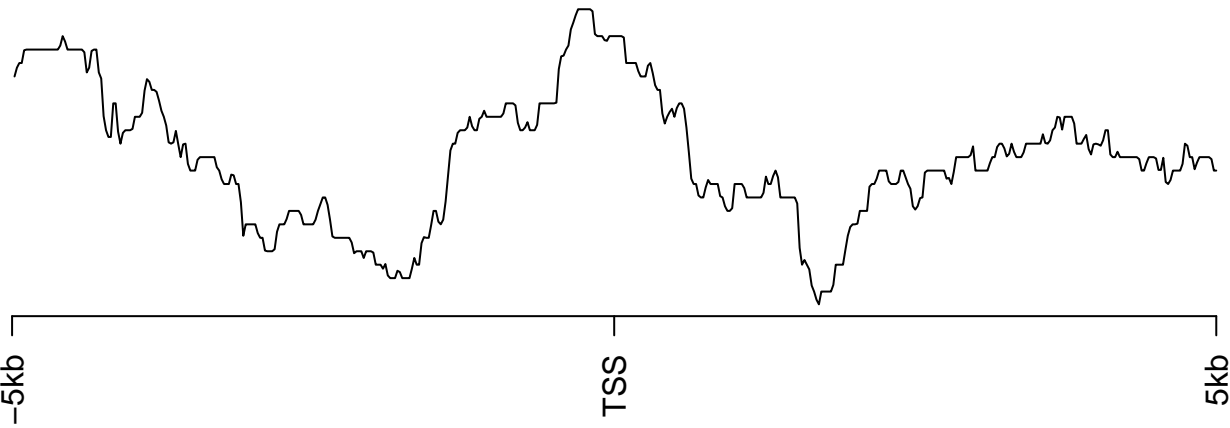
Average coverage per gene

7e-04
4e-04

-5kb

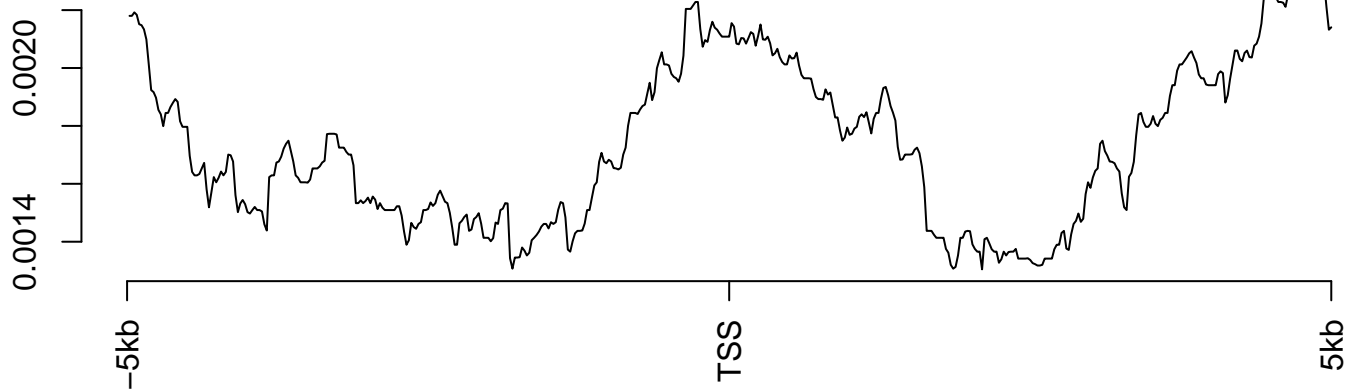
TSS

5kb



MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc790

Average coverage per gene



MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc791

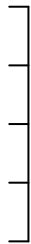
Average coverage per gene

2e-04 4e-04 6e-04

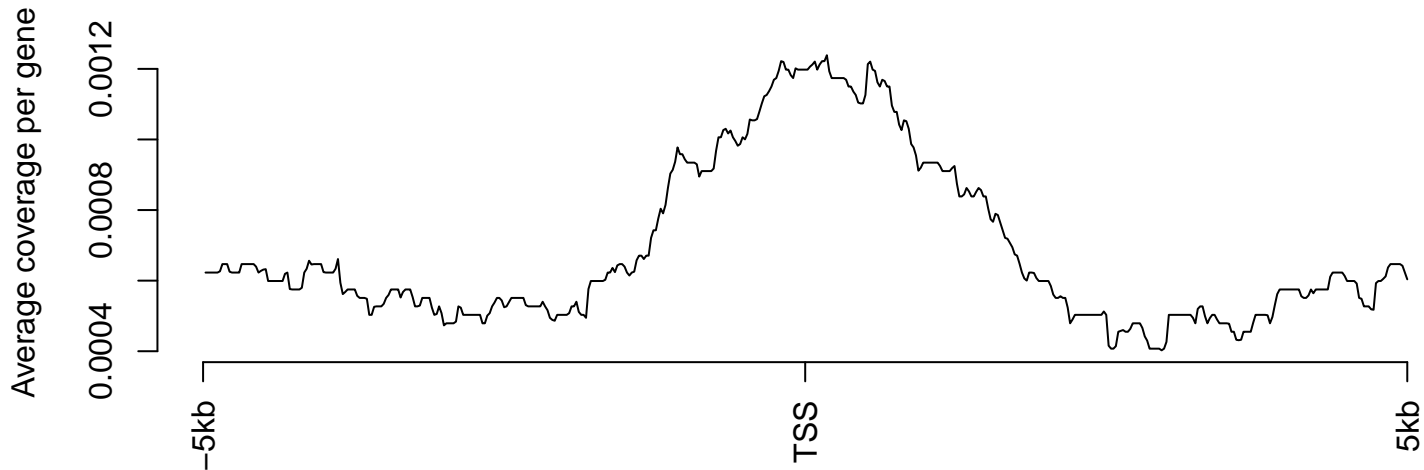
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc792



MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc793

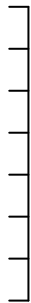
Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc794

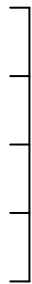
Average coverage per gene

0.0006 0.0010 0.0014

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc795

Average coverage per gene

0.0004 0.0010

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.669 scCLTdegenNuc808

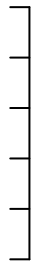
Average coverage per gene

0.0020 0.0026

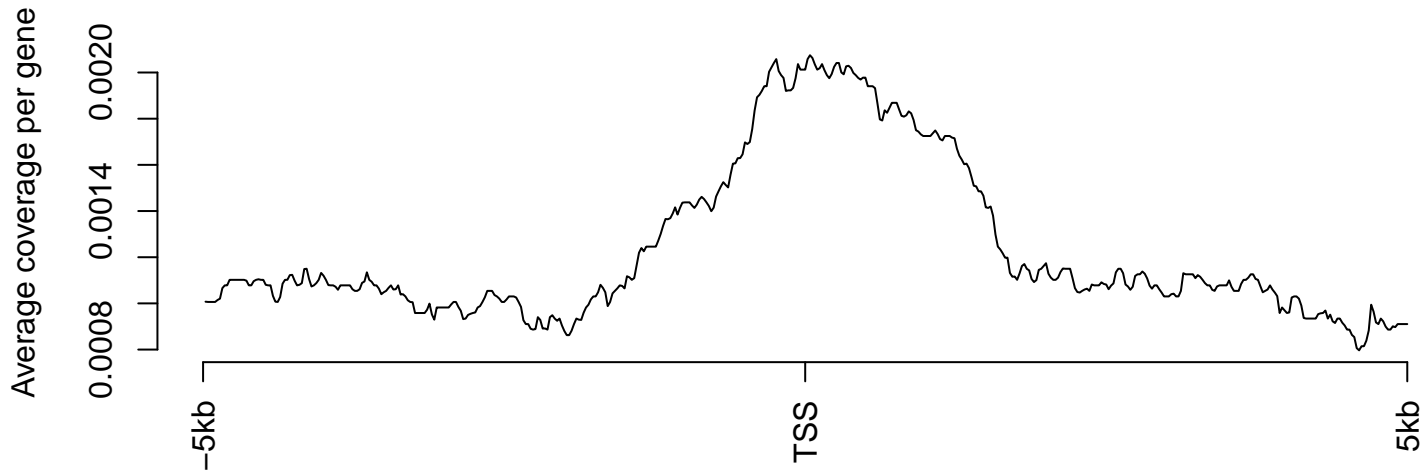
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc797



MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc798

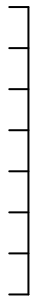
Average coverage per gene

0.0009
0.0005

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc799

Average coverage per gene

0.0015 0.0025

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.671 scCLTdegenNuc803

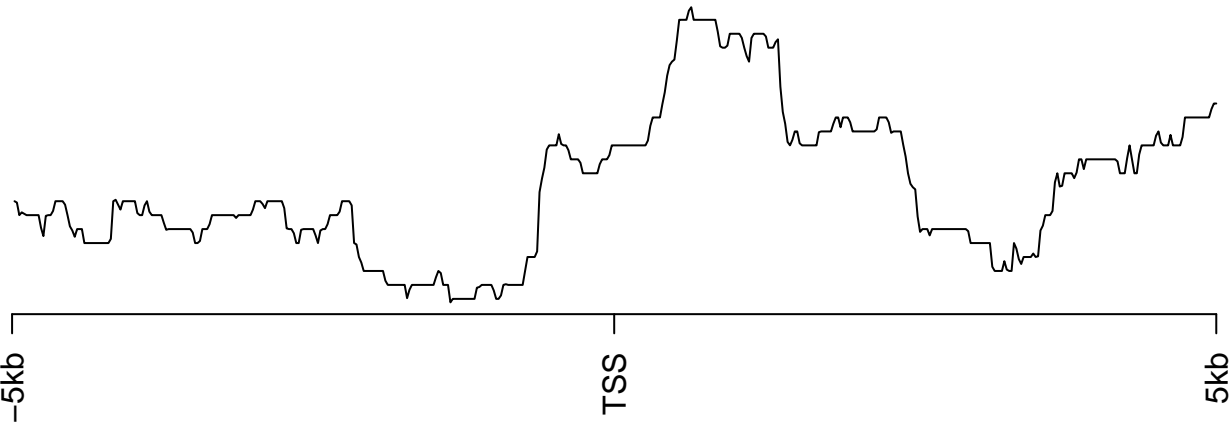
Average coverage per gene

3e-04
6e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.671 scCLTdegenNuc807

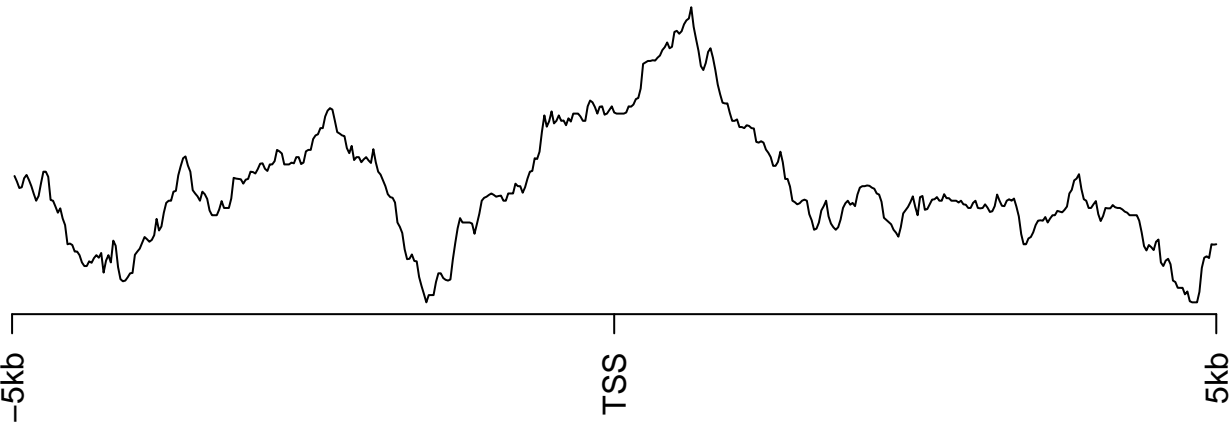
Average coverage per gene

0.0020
0.0014

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc965

Average coverage per gene

0.00010
0.00000

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc966

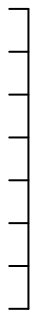
Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc967

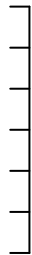
Average coverage per gene

0.00025
0.00005

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc968

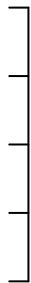
Average coverage per gene

0.00015
0.00000

-5kb

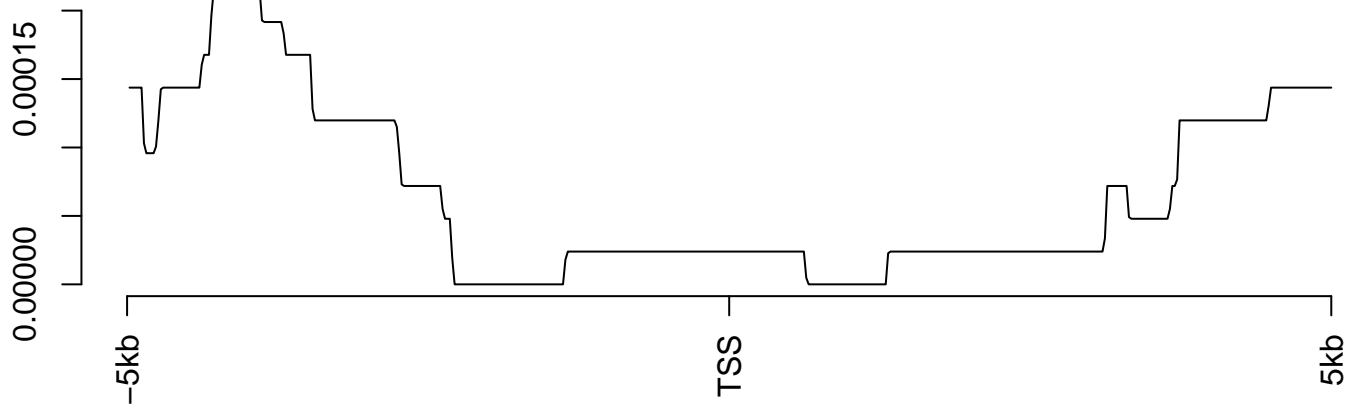
TSS

5kb



MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc969

Average coverage per gene



MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc970

Average coverage per gene

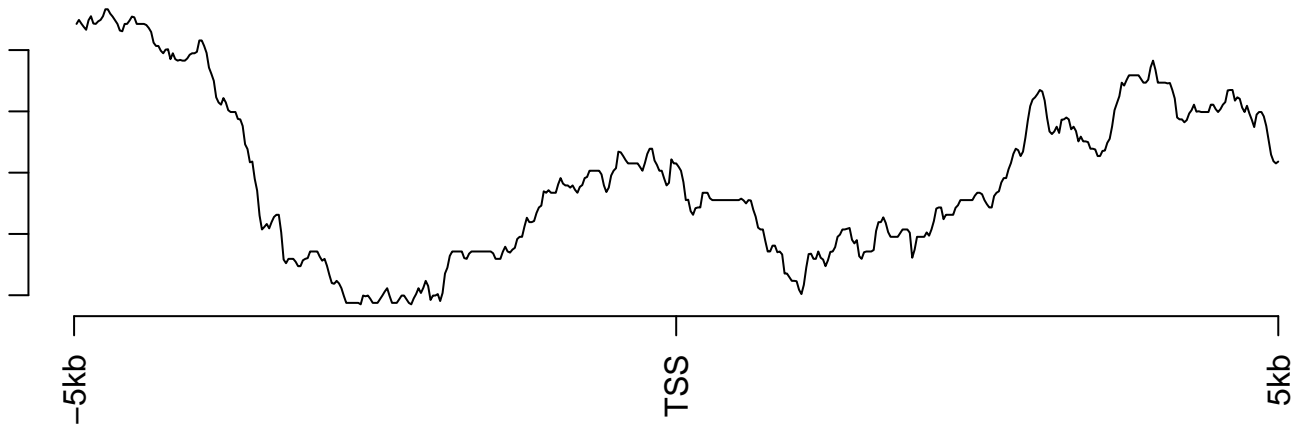
0.0012

0.0006

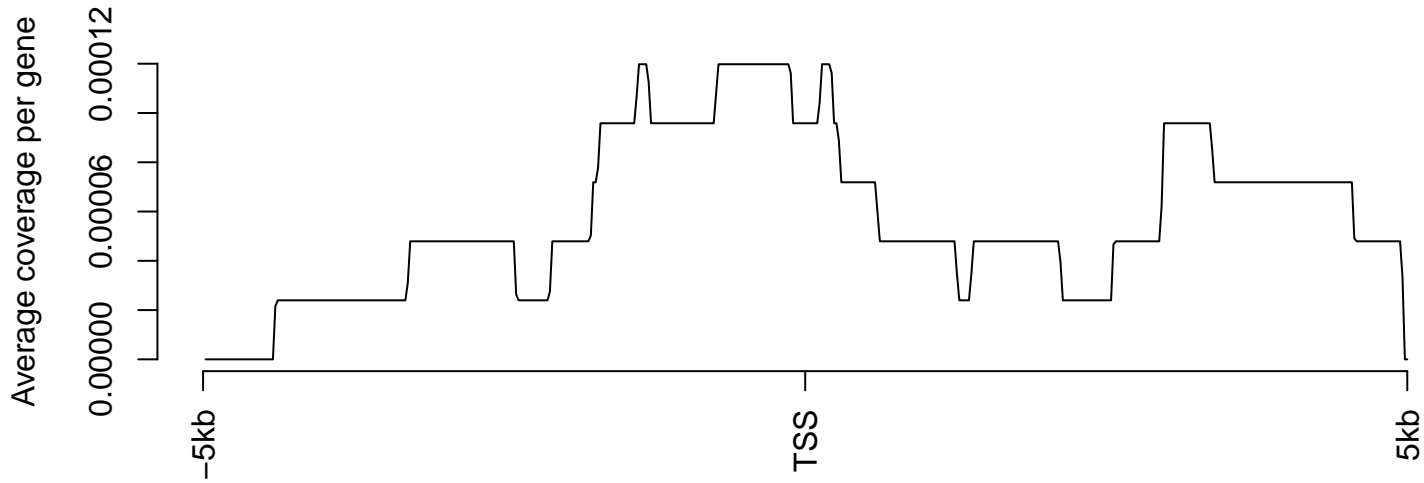
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1063



MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1064

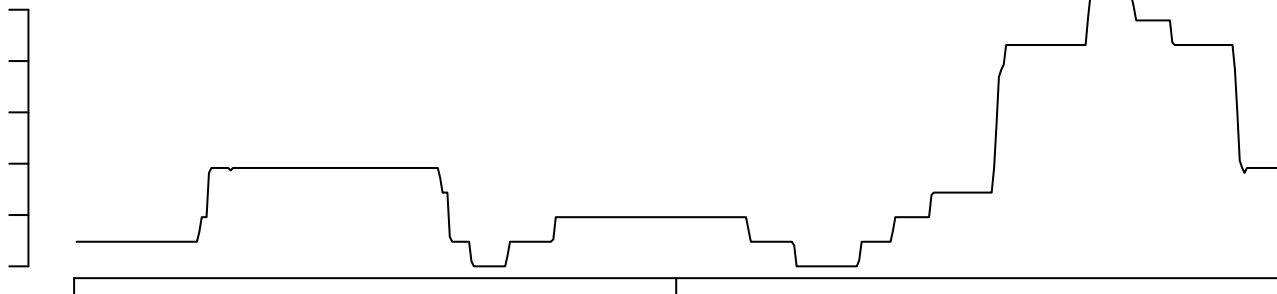
Average coverage per gene

0.00000 0.00015

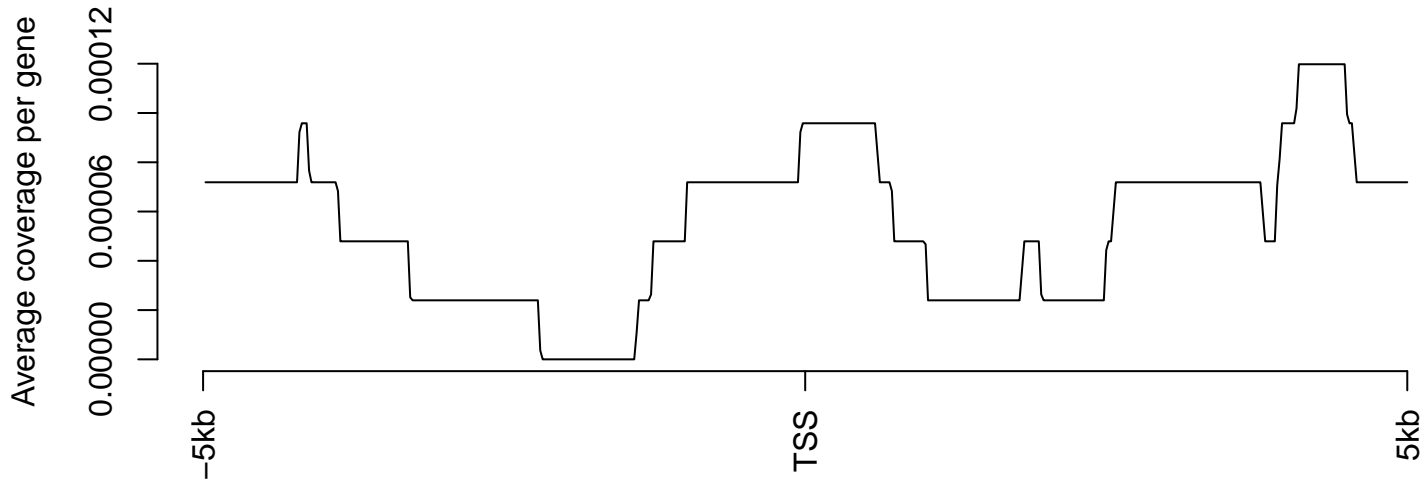
-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1065



MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1068

Average coverage per gene

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

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1066

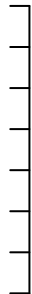
Average coverage per gene

0.00000 0.00008

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1067

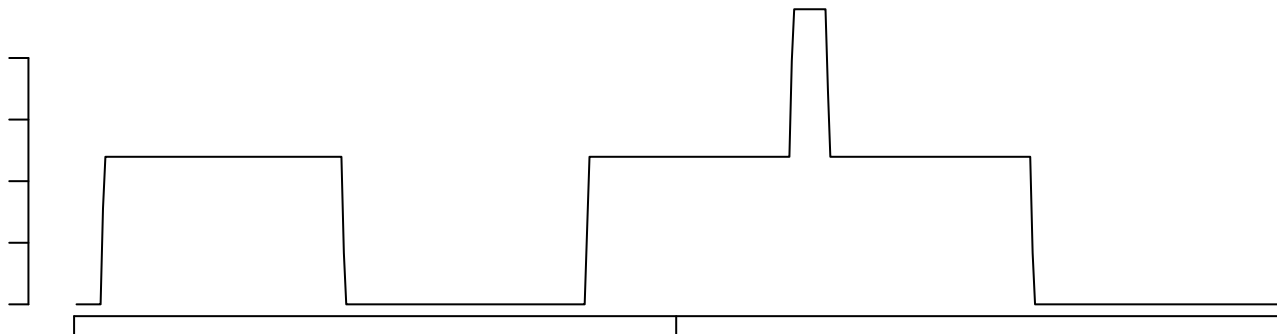
Average coverage per gene

0e+00 2e-05 4e-05

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1095

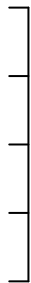
Average coverage per gene

0.00015
0.00000

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1096

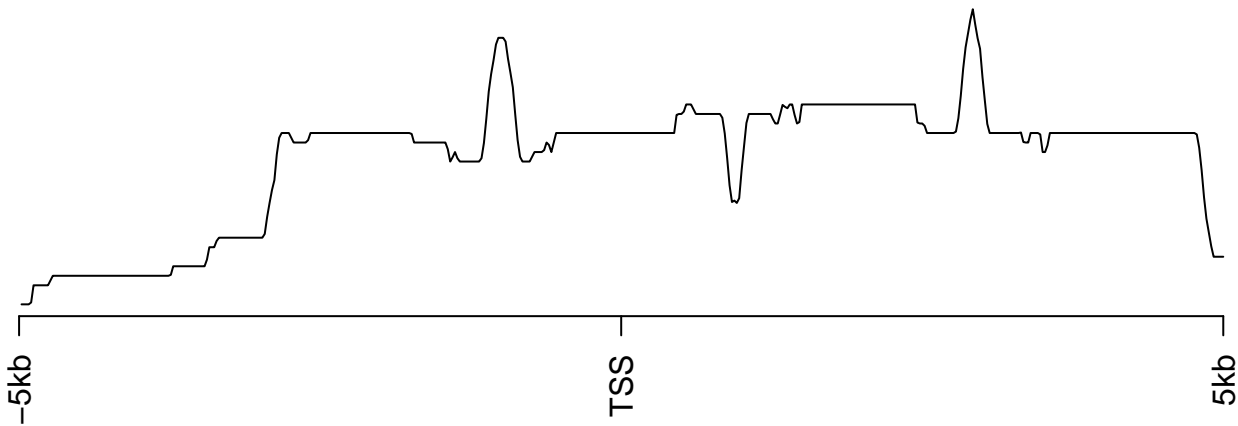
Average coverage per gene

0e+00
4e-04

-5kb

TSS

5kb



MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1097

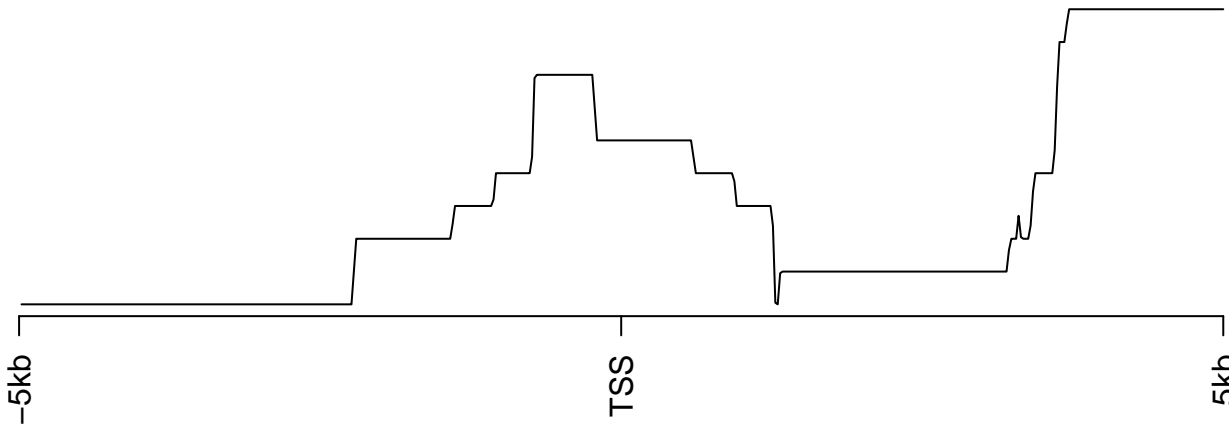
Average coverage per gene

0.00015
0.00000

-5kb

TSS

5kb



MouseNeuronCulture PositivePooled E.584 scCLTdegenNuc415

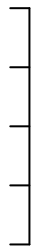
Average coverage per gene

0.055 0.065 0.075

-5kb

TSS

5kb



MouseNeuronCulture PositivePooled E.643 scCLTdegenNuc654

Average coverage per gene

0.065
0.055

-5kb

TSS

5kb



MouseNeuronCulture PositivePooled E.669 scCLTdegenNuc809

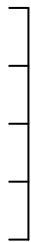
Average coverage per gene

0.24 0.28 0.32

-5kb

TSS

5kb



MouseNeuronCulture PositiveMerged E.– MouseNeuronCulturePositiveMerged

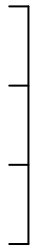
Average coverage per gene

0.10
0.14

-5kb

TSS

5kb



MouseNeuronCulture PositiveAll E.– MouseNeuronCulturePositiveAll

