

Average coverage per intergenic region

MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc429

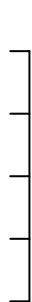
0e+00 2e-04 4e-04

-5kb

intergenic start

intergenic end

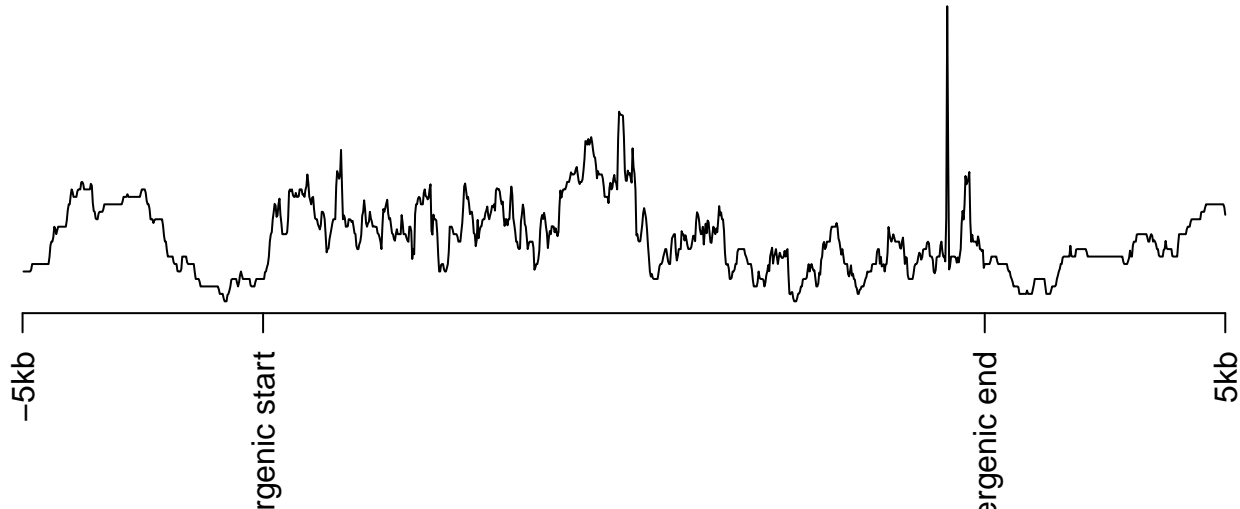
5kb



MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc430

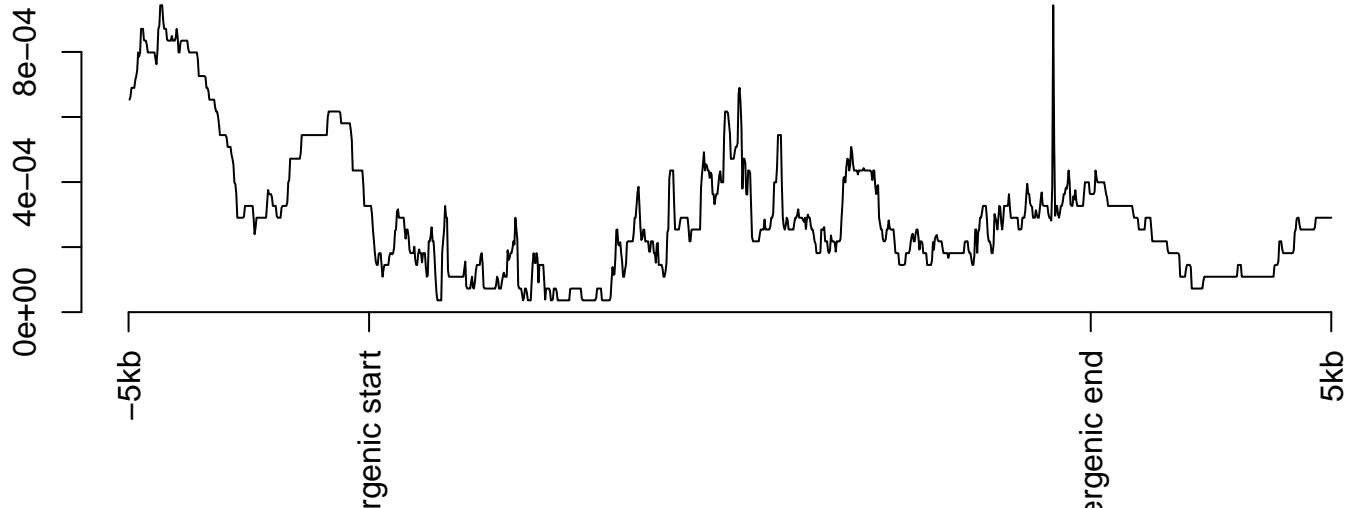
Average coverage per intergenic region

0.0002 0.0010



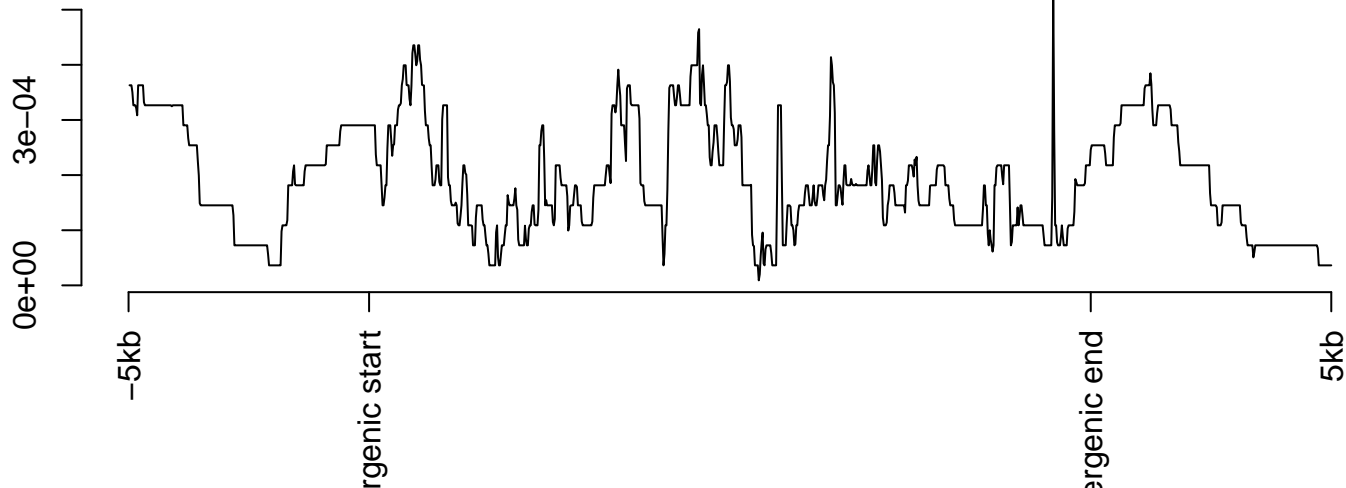
Average coverage per intergenic region

MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc431



Average coverage per intergenic region

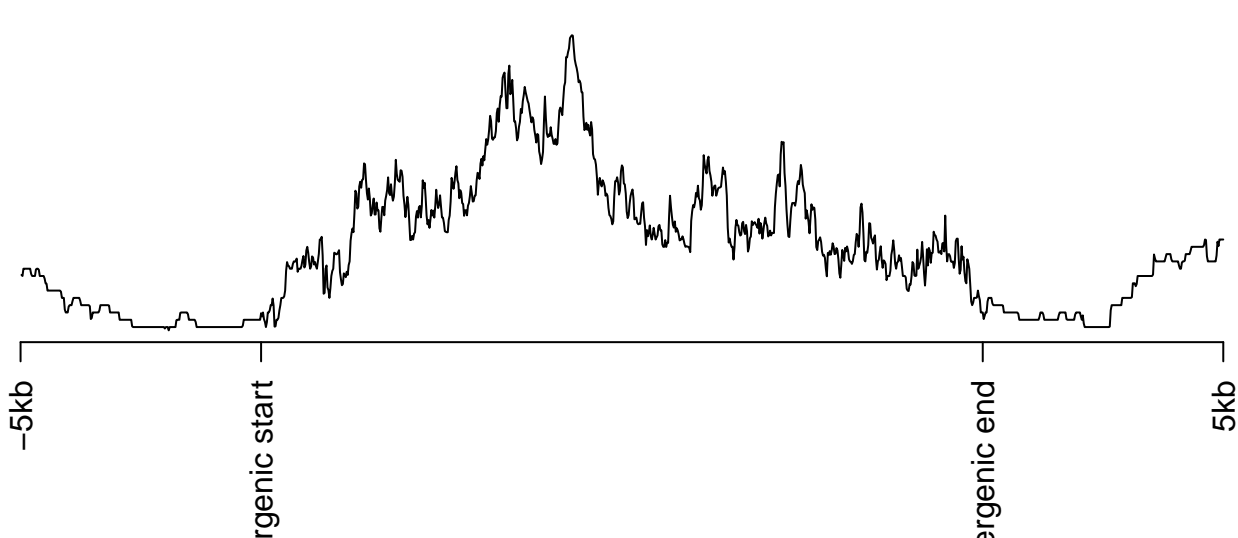
MouseNeuronCulture NegCtrlSingle E.584 scCLTdegenNuc433



Average coverage per intergenic region

MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc886

0.0015
0.0005



Average coverage per intergenic region

MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc891

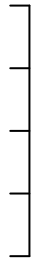
1e-04 3e-04 5e-04

-5kb

intergenic start

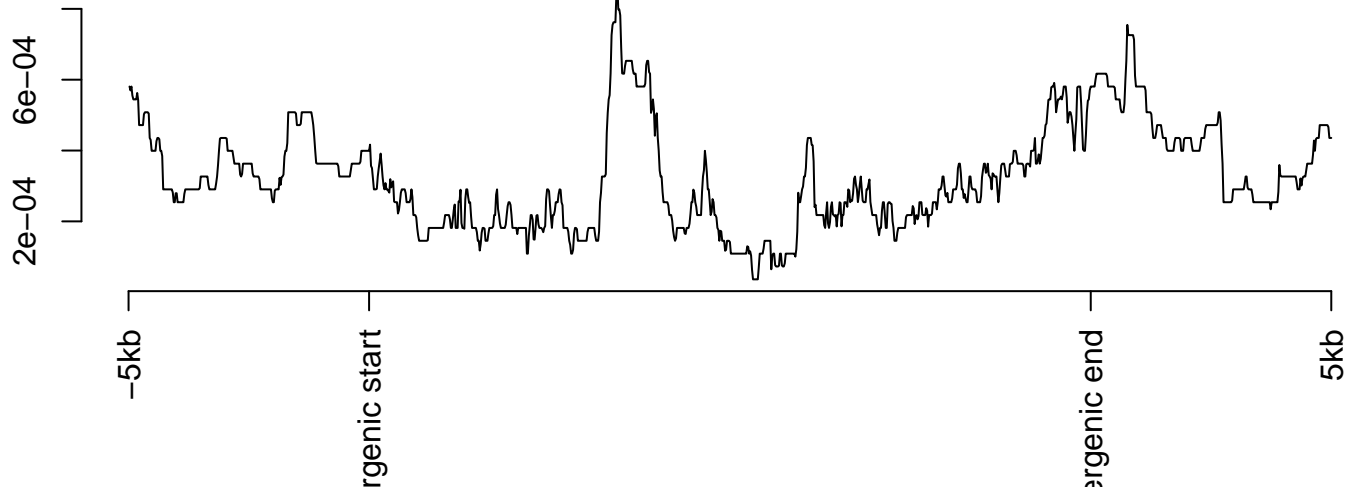
intergenic end

5kb



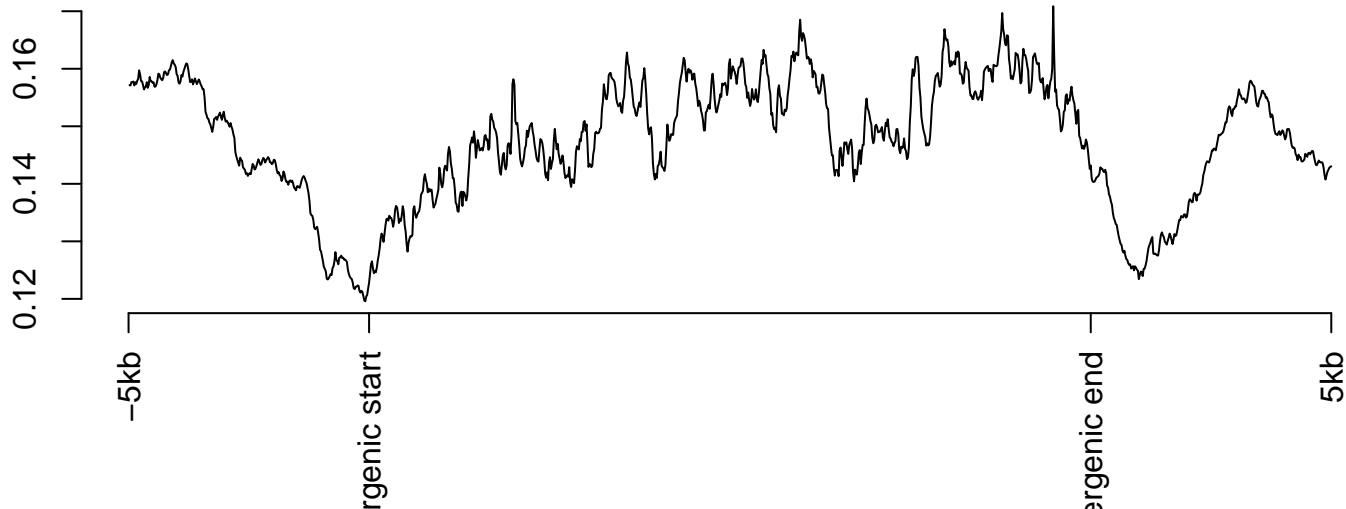
Average coverage per intergenic region

MouseNeuronCulture NegCtrlSingle E.670 scCLTdegenNuc892



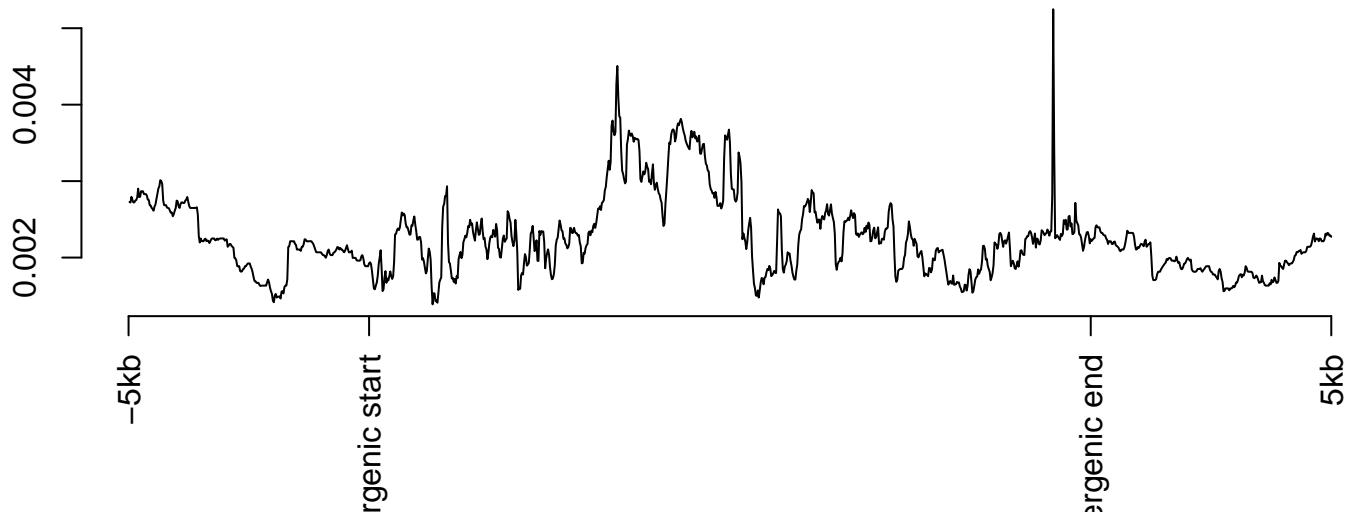
Average coverage per intergenic region

MouseNeuronCulture NegCtrlPooled E.597 scCLTdegenNuc471



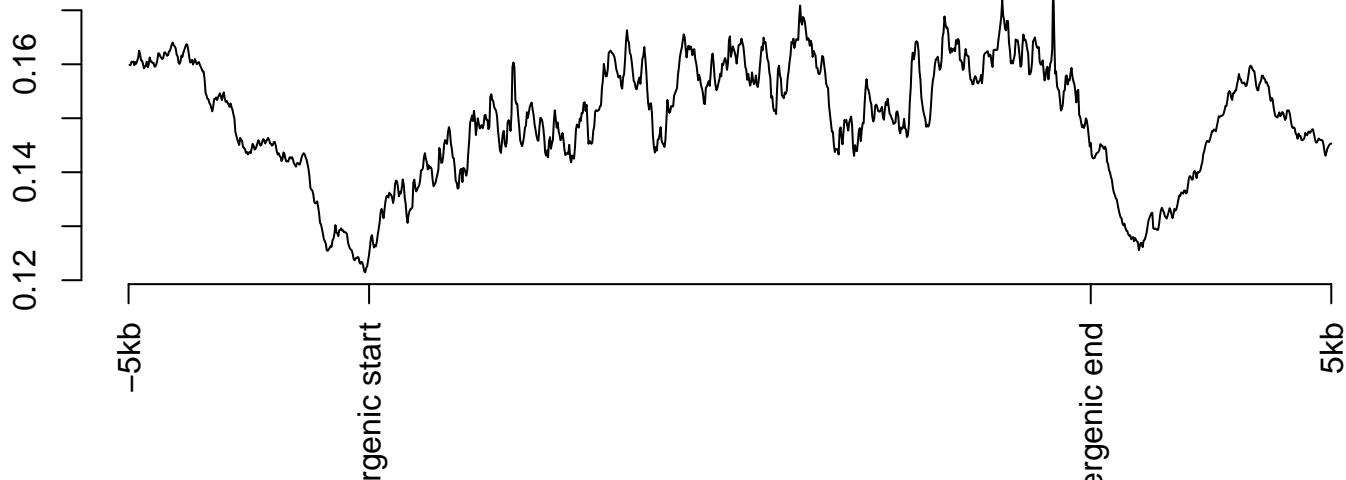
Average coverage per intergenic region

MouseNeuronCulture NegCtrlMerged E.– MouseNeuronCultureNegCtrlMerged



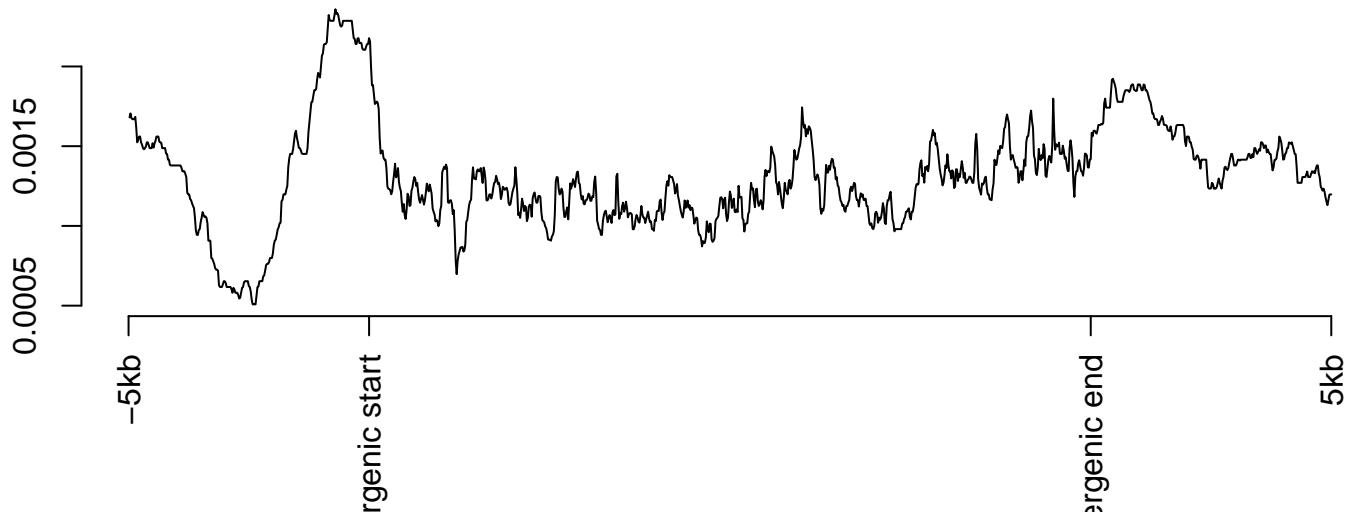
Average coverage per intergenic region

MouseNeuronCulture NegCtrlAll E.– MouseNeuronCultureNegCtrlAll



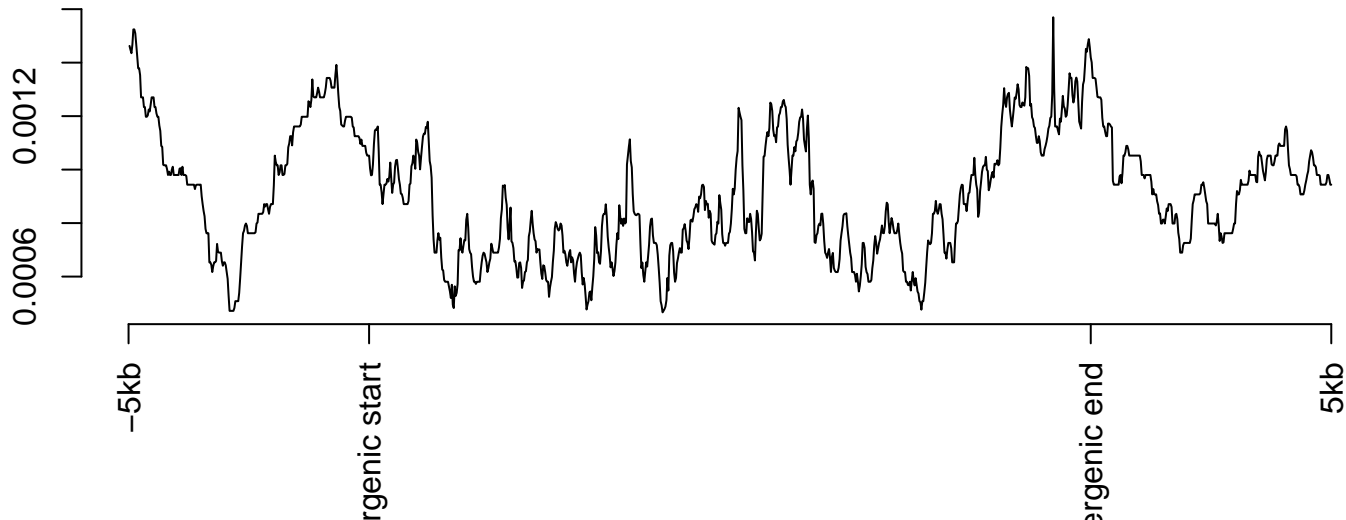
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc387



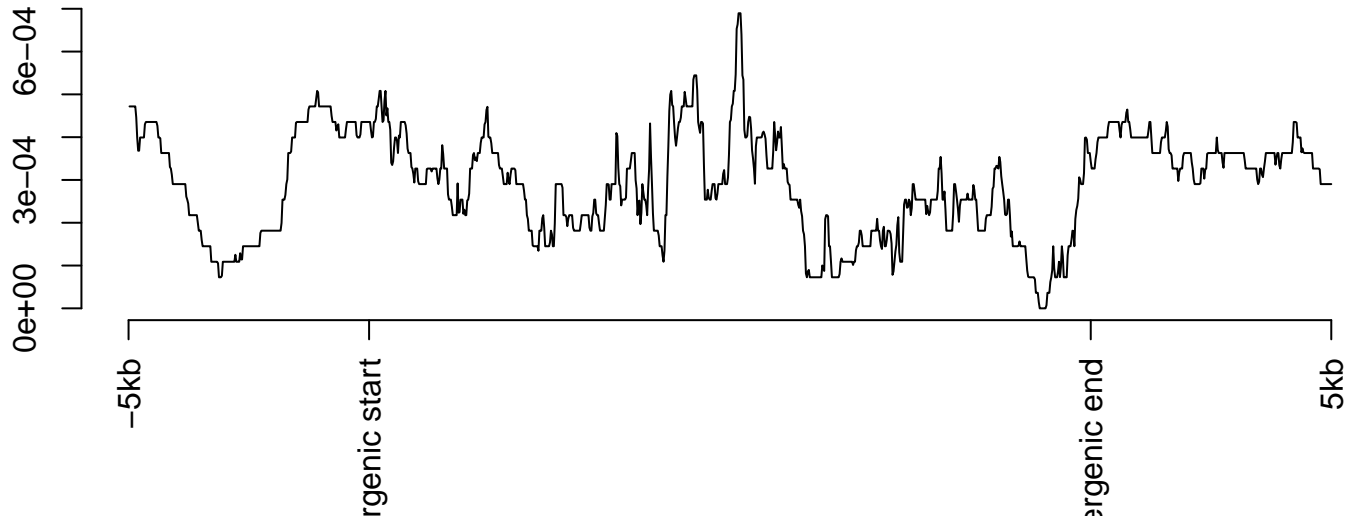
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc388



Average coverage per intergenic region

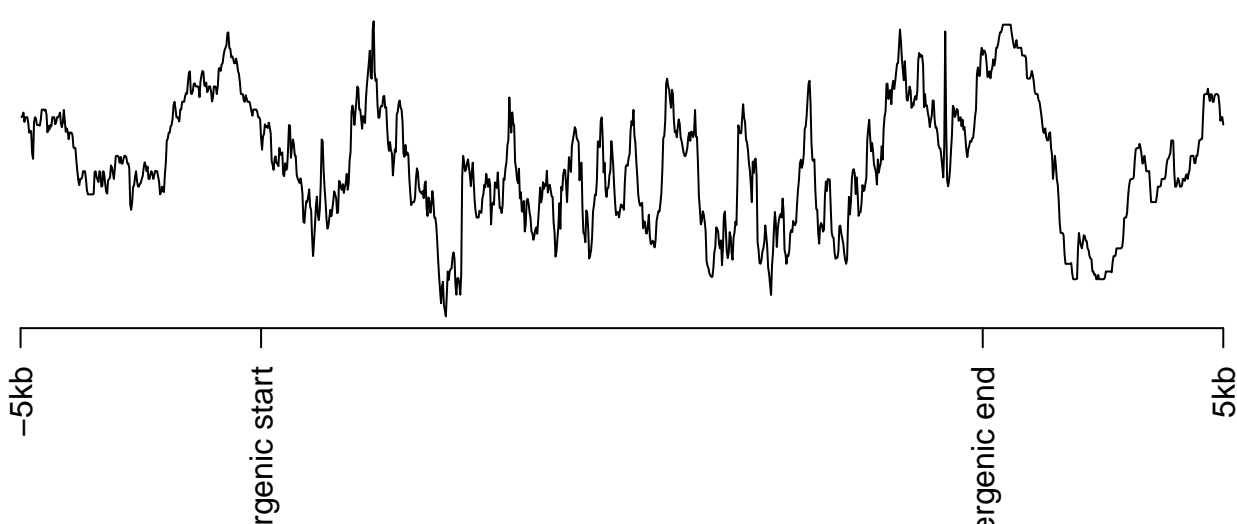
MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc390



Average coverage per intergenic region

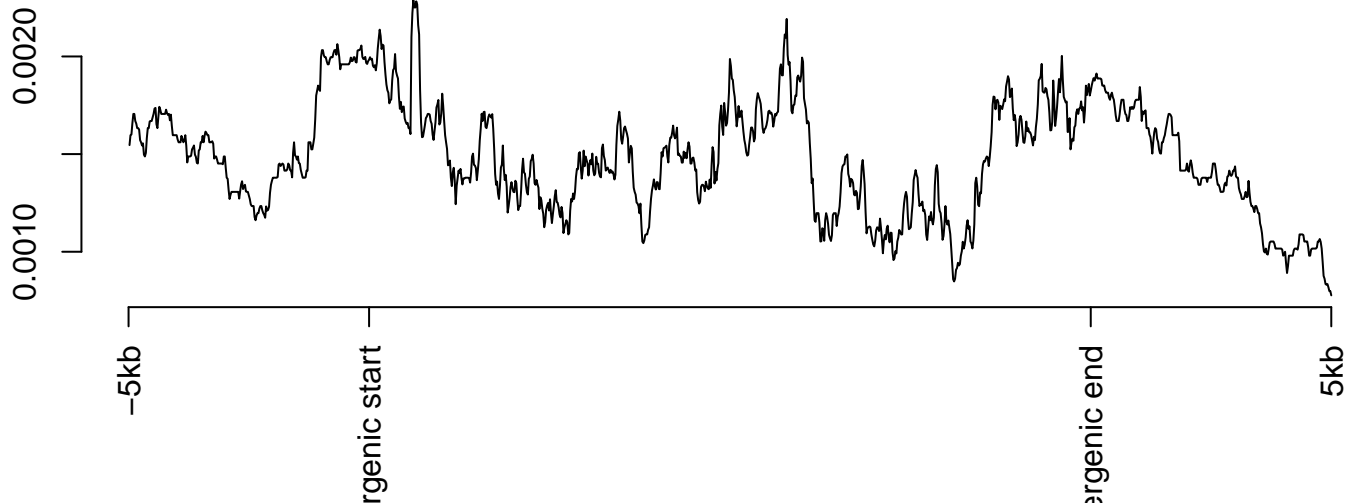
MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc391

0.0008 0.0014 0.0020



Average coverage per intergenic region

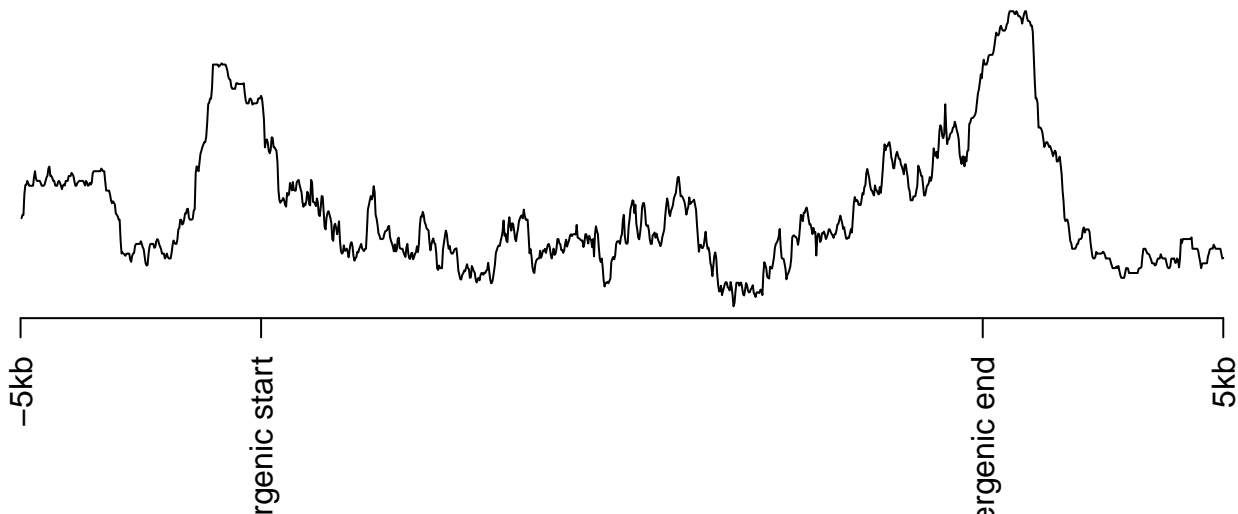
MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc392



MouseNeuronCulture PositiveSingle E.584 scCLTdegenNuc393

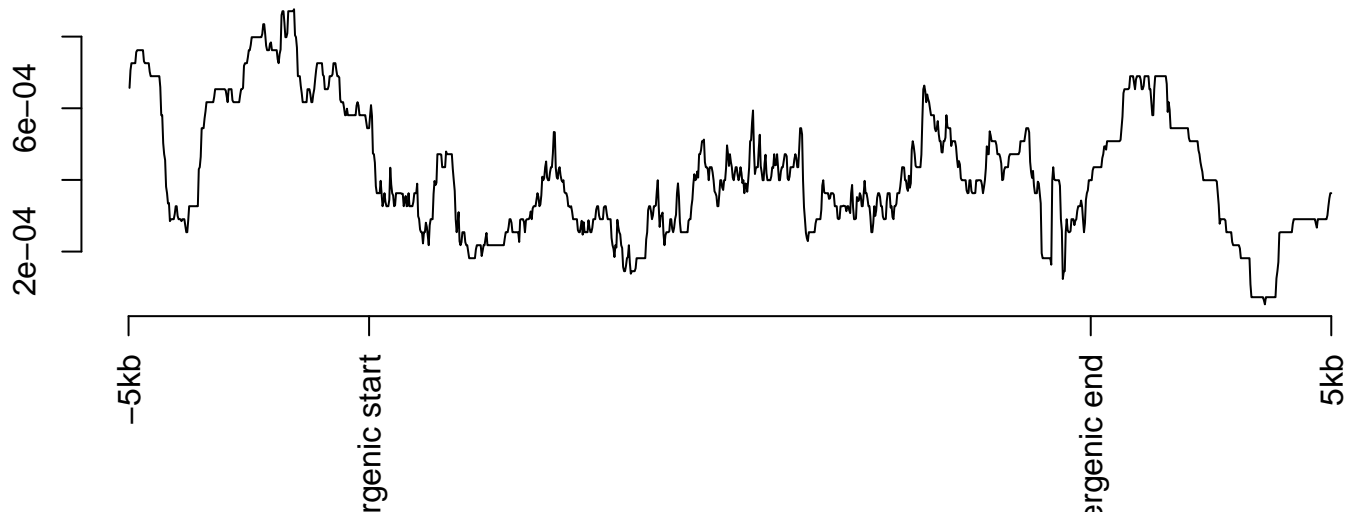
Average coverage per intergenic region

0.0010 0.0020



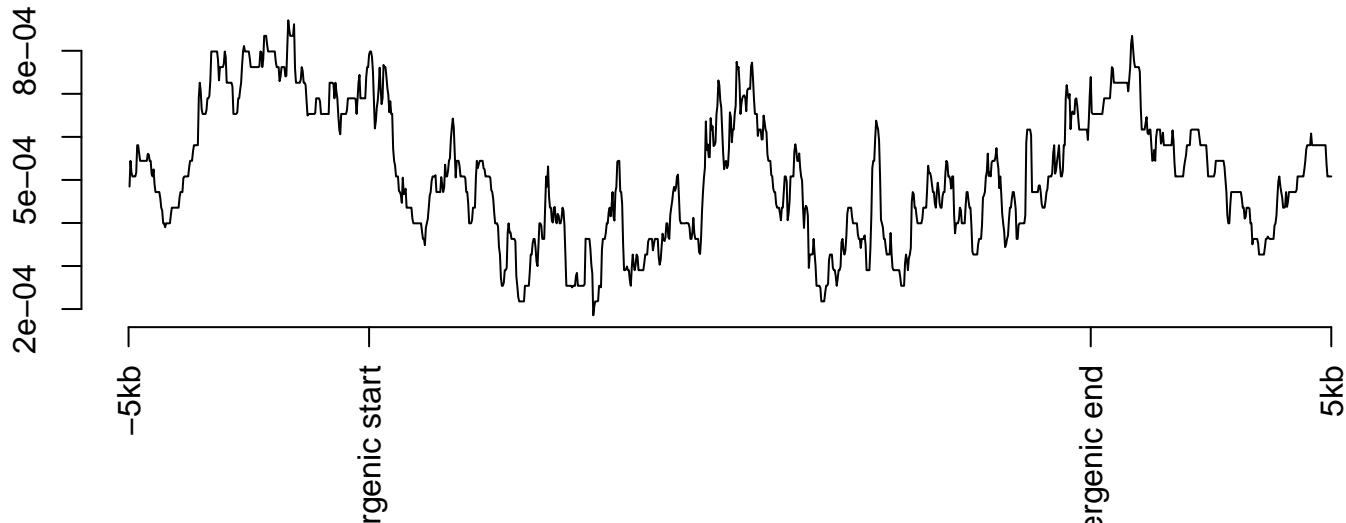
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc501



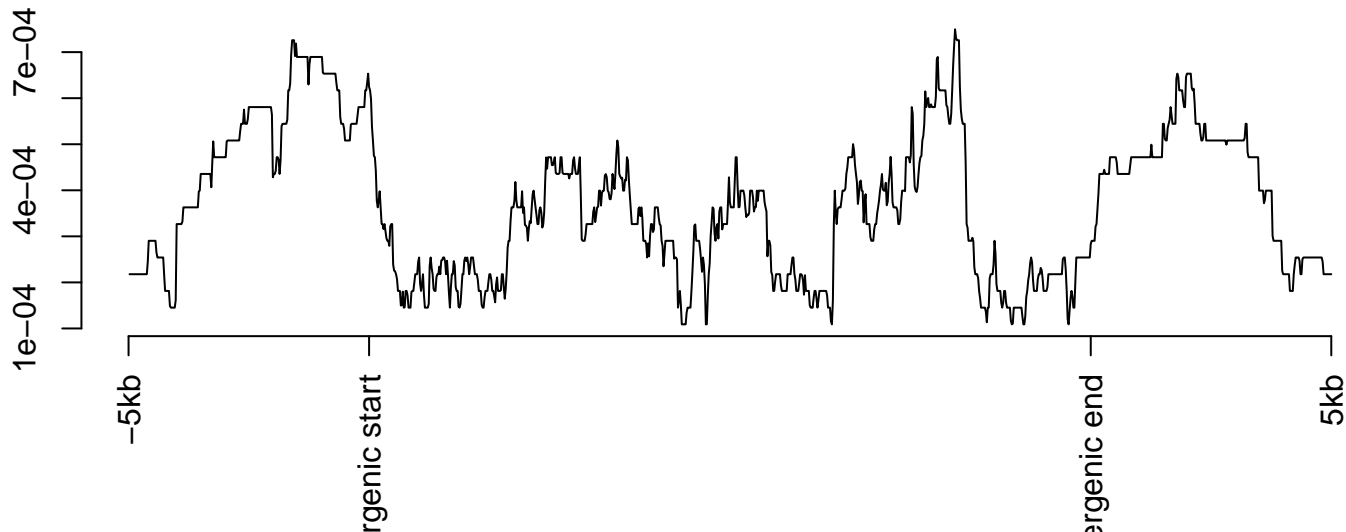
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc502



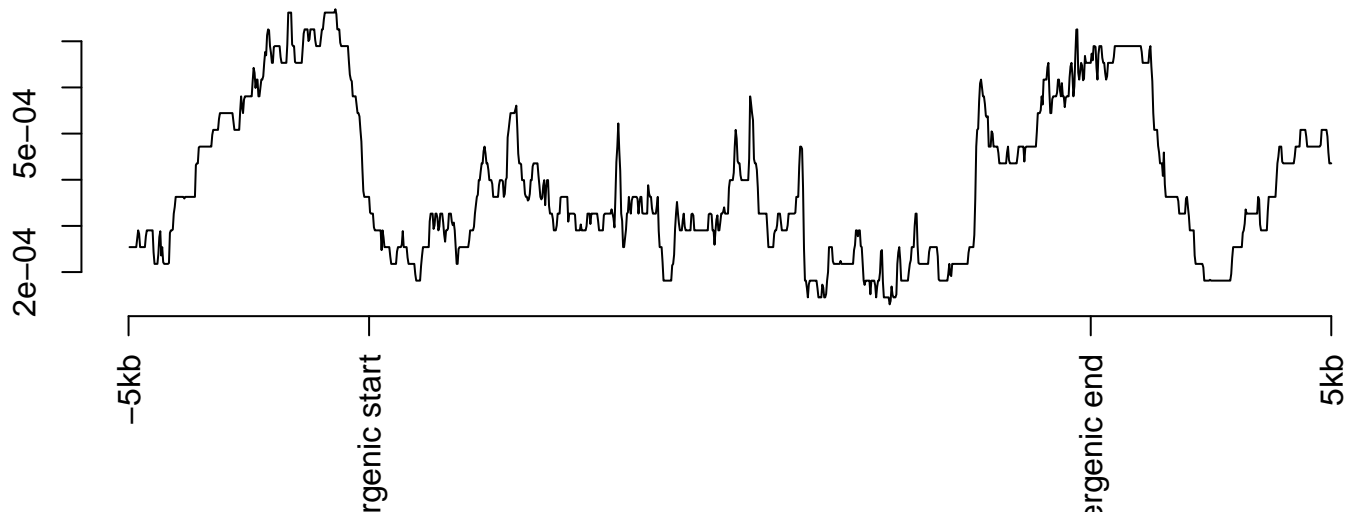
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc503



Average coverage per intergenic region

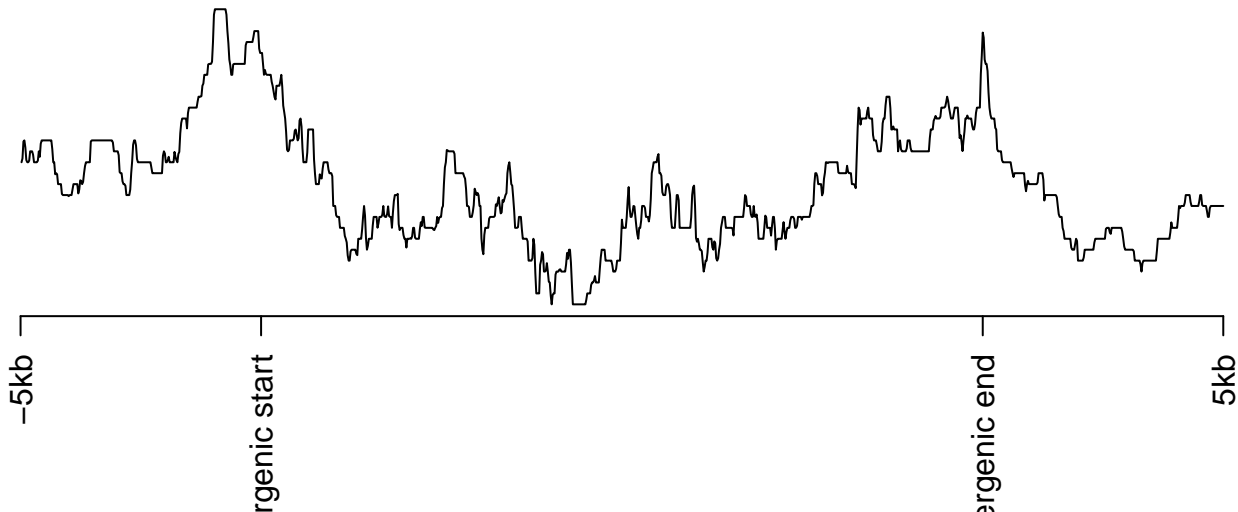
MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc504



Average coverage per intergenic region

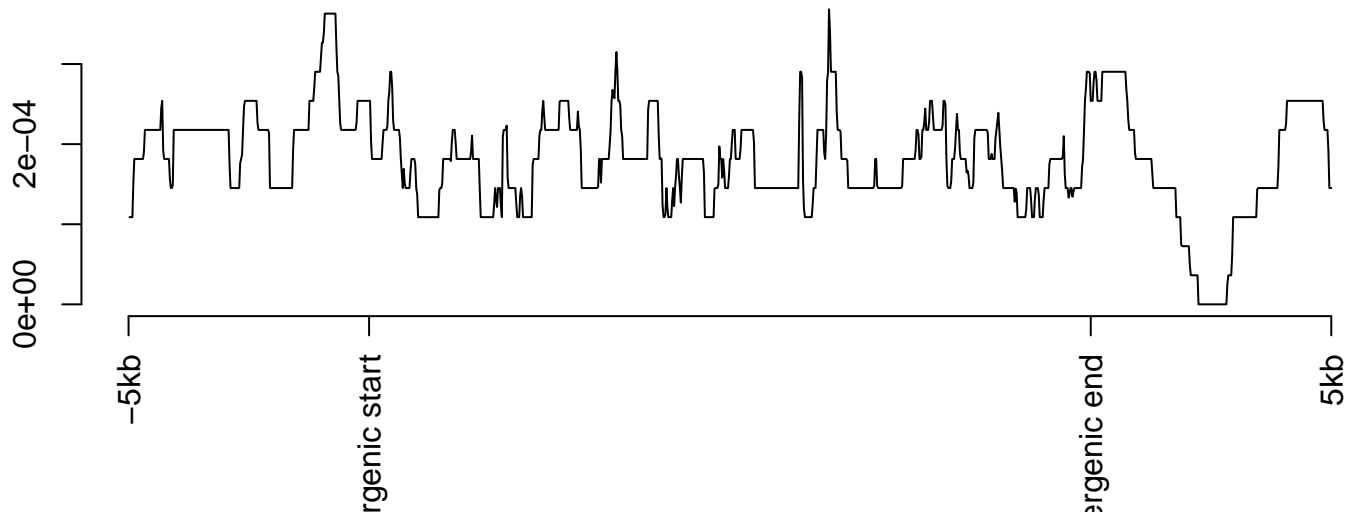
MouseNeuronCulture PositiveSingle E.619 scCLTdegenNuc505

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



Average coverage per intergenic region

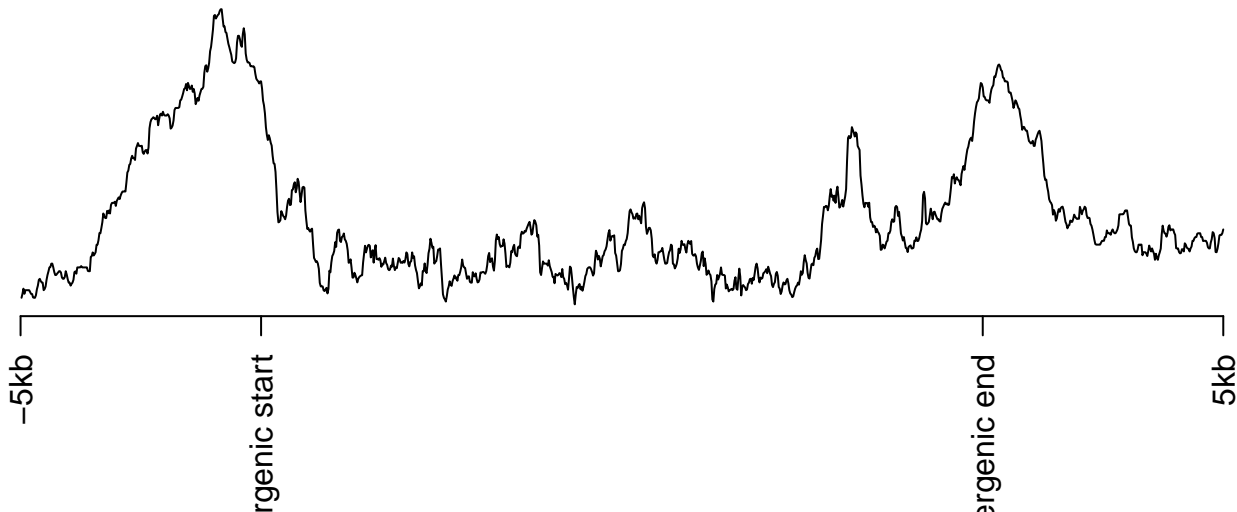
MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc510



MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc511

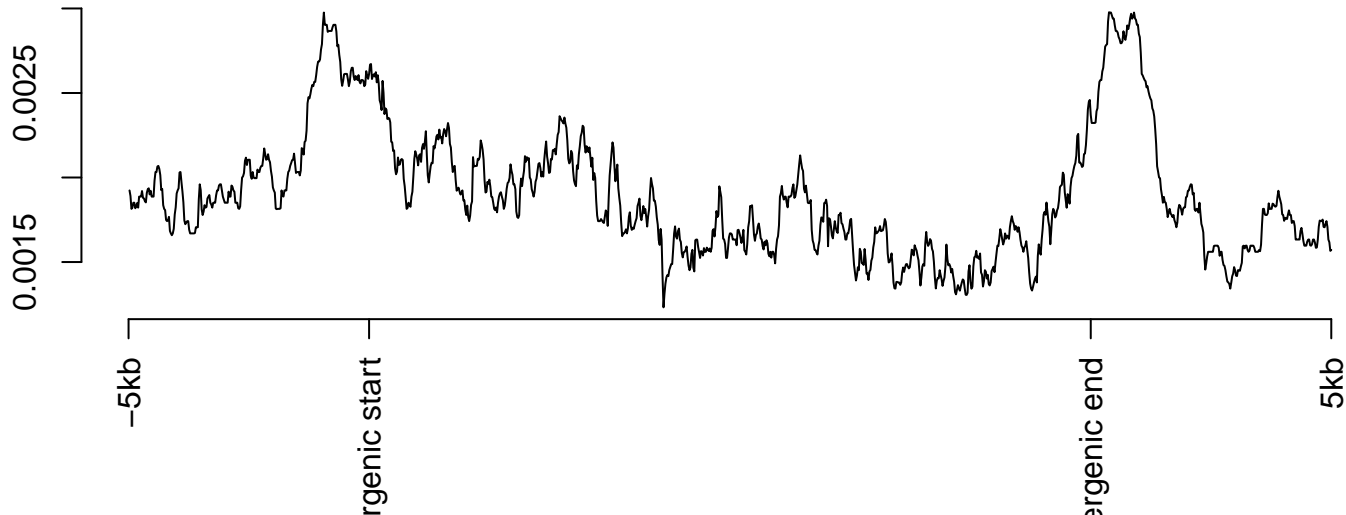
Average coverage per intergenic region

0.0030
0.0015



Average coverage per intergenic region

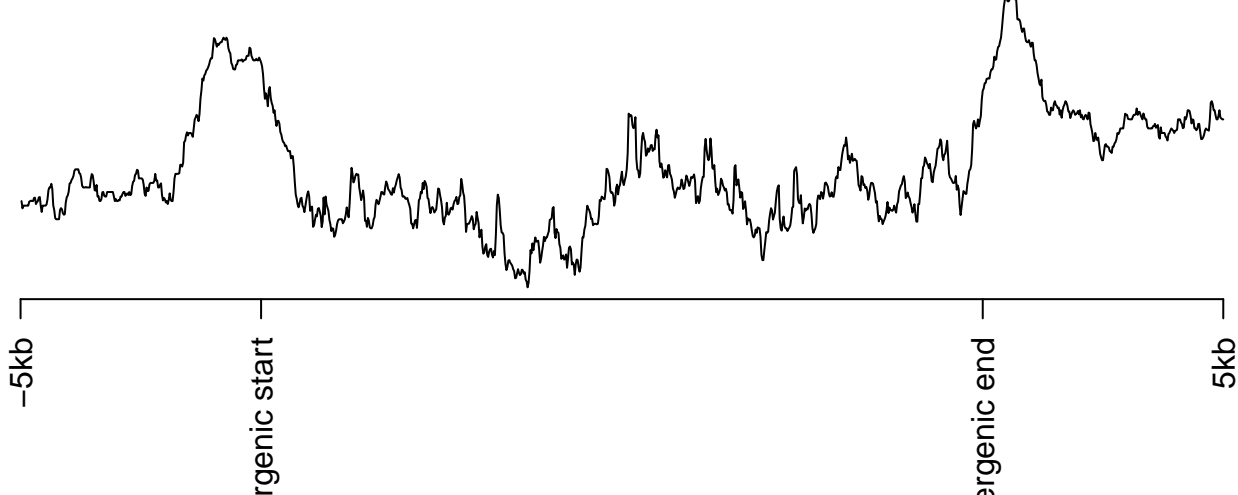
MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc512



Average coverage per intergenic region

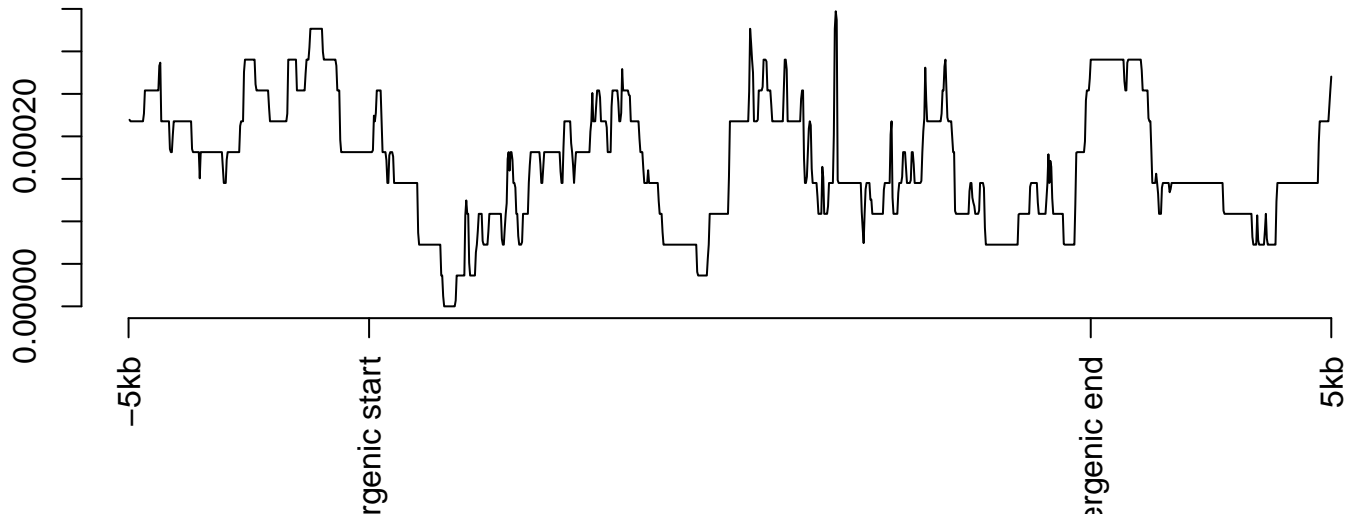
MouseNeuronCulture PositiveSingle E.622 scCLTdegenNuc513

0.0010 0.0020 0.0030



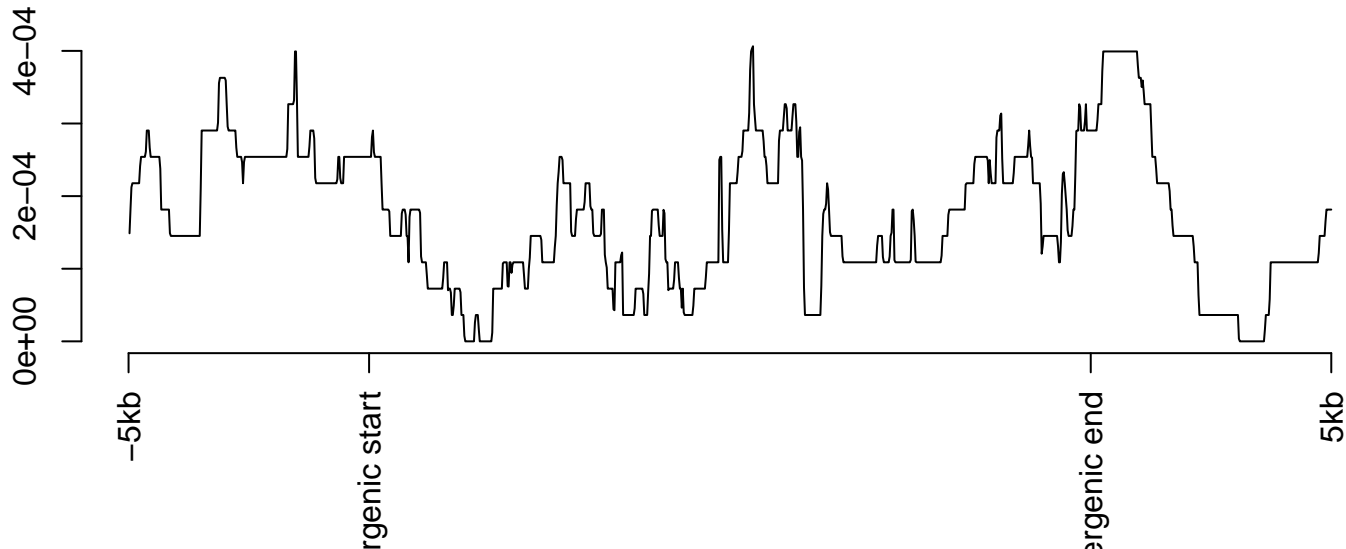
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc521



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc522



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc523

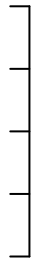
1e-04
3e-04
5e-04

-5kb

intergenic start

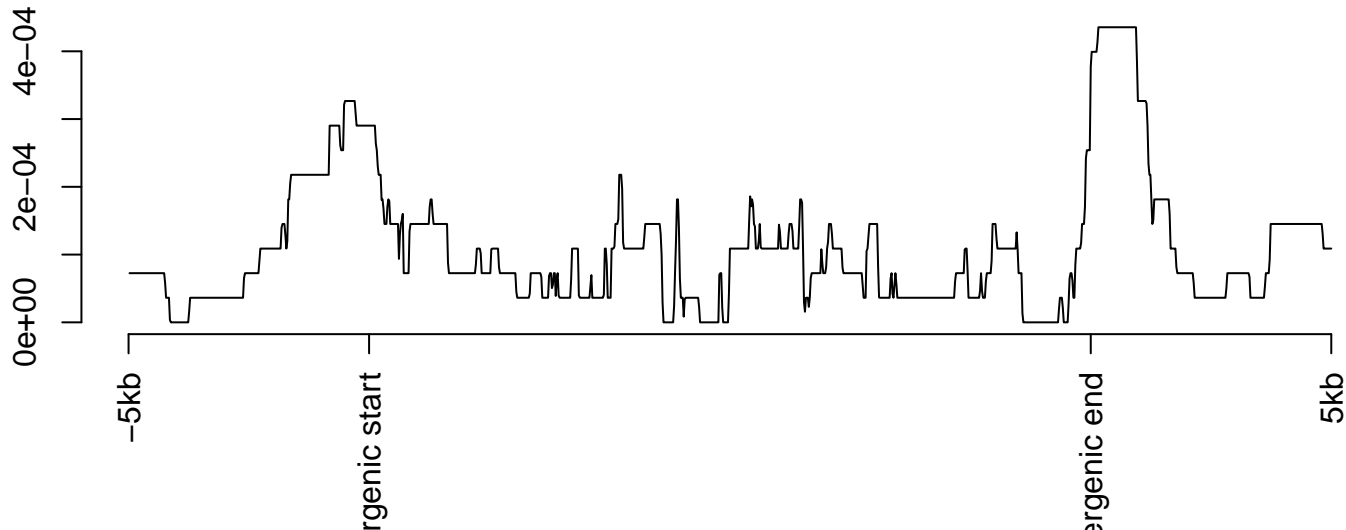
intergenic end

5kb



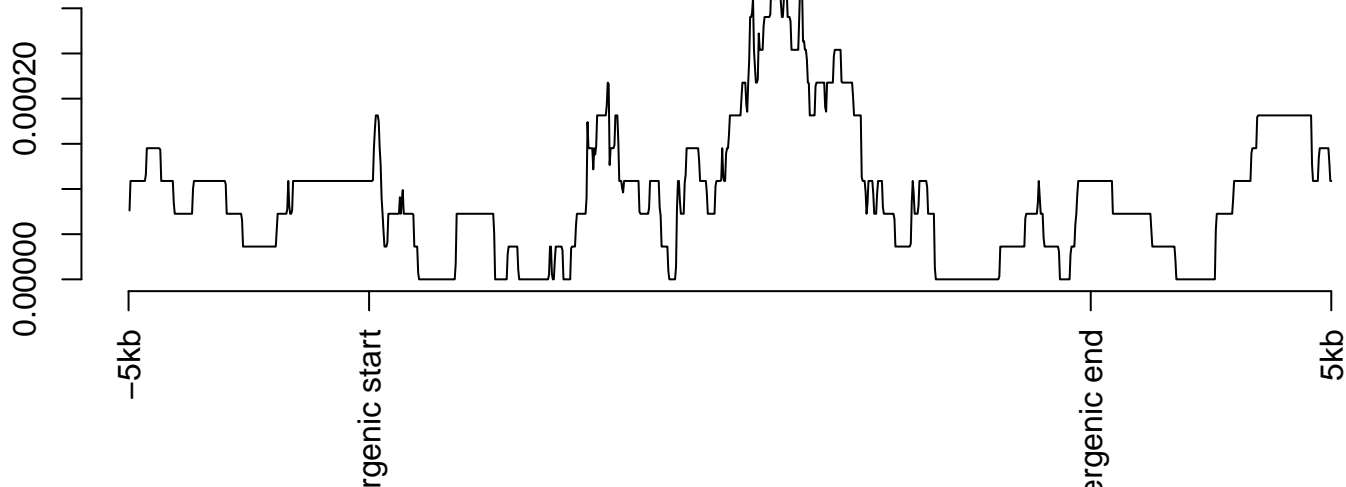
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc524



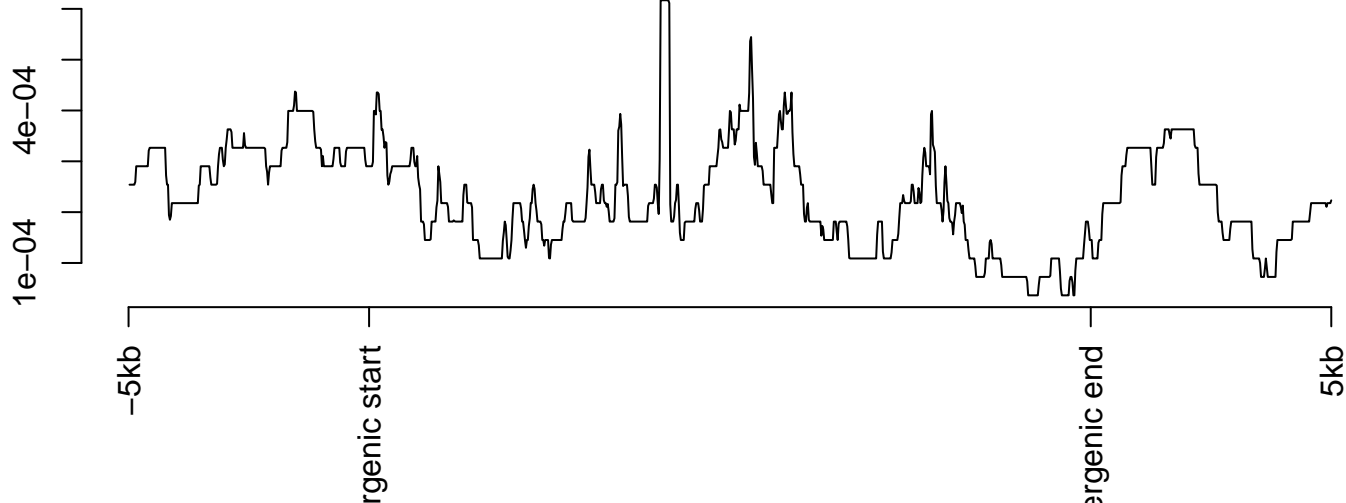
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc525



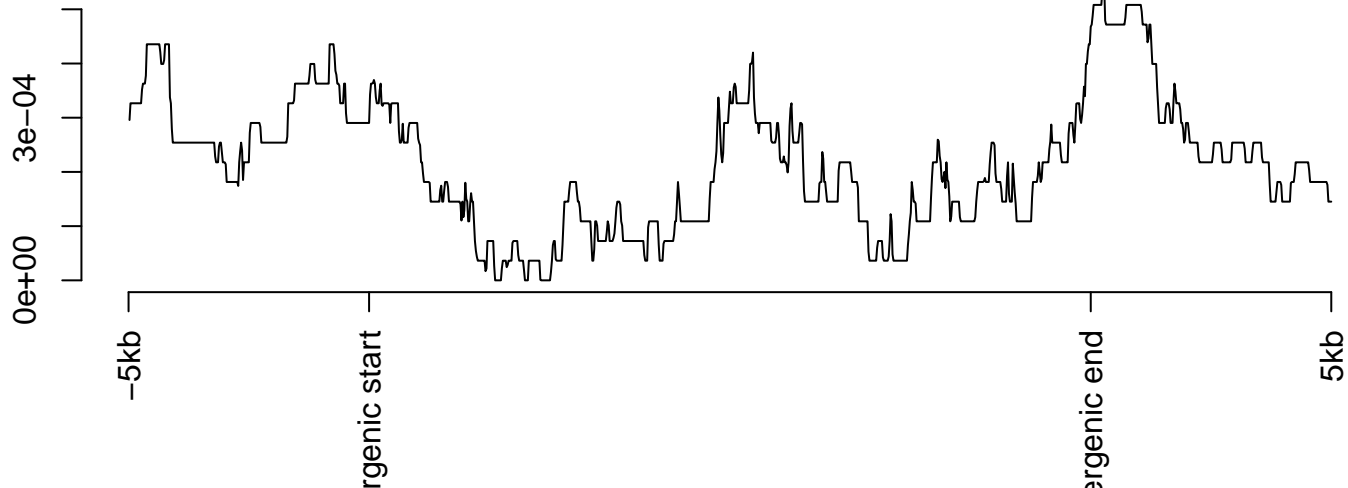
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc526



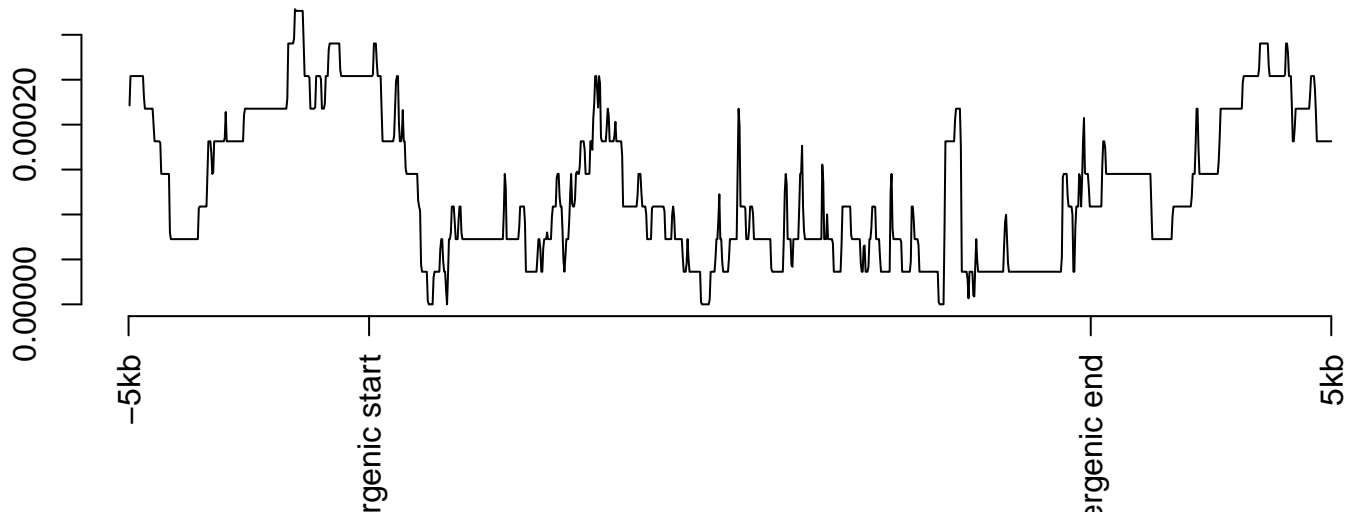
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc527



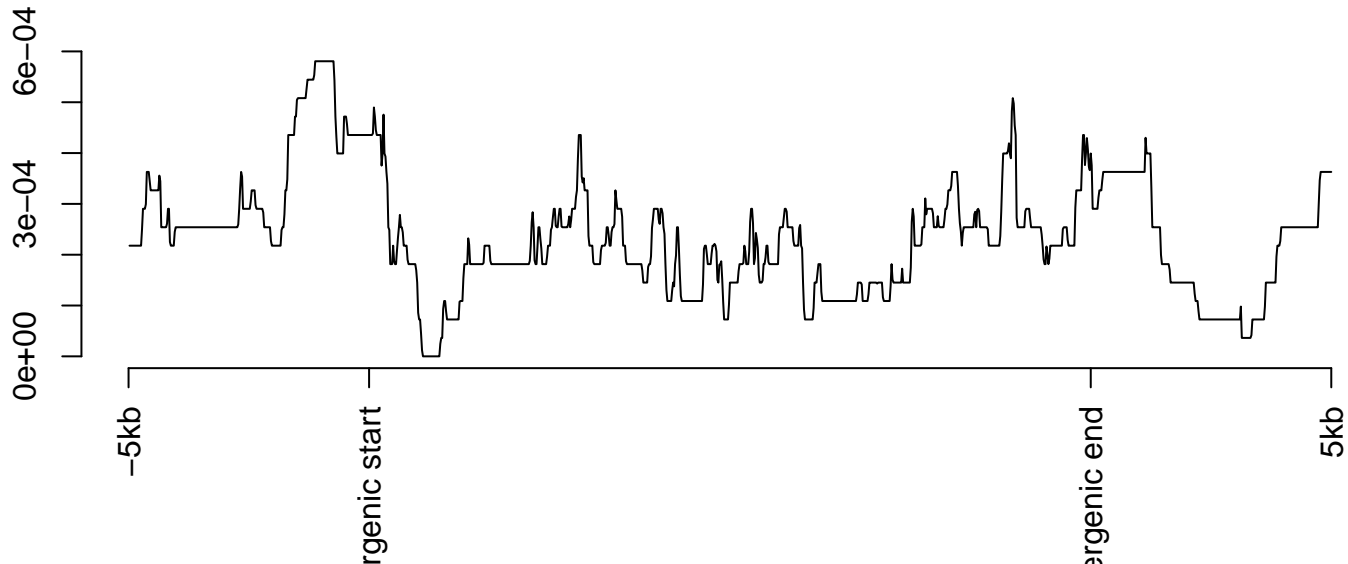
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc528



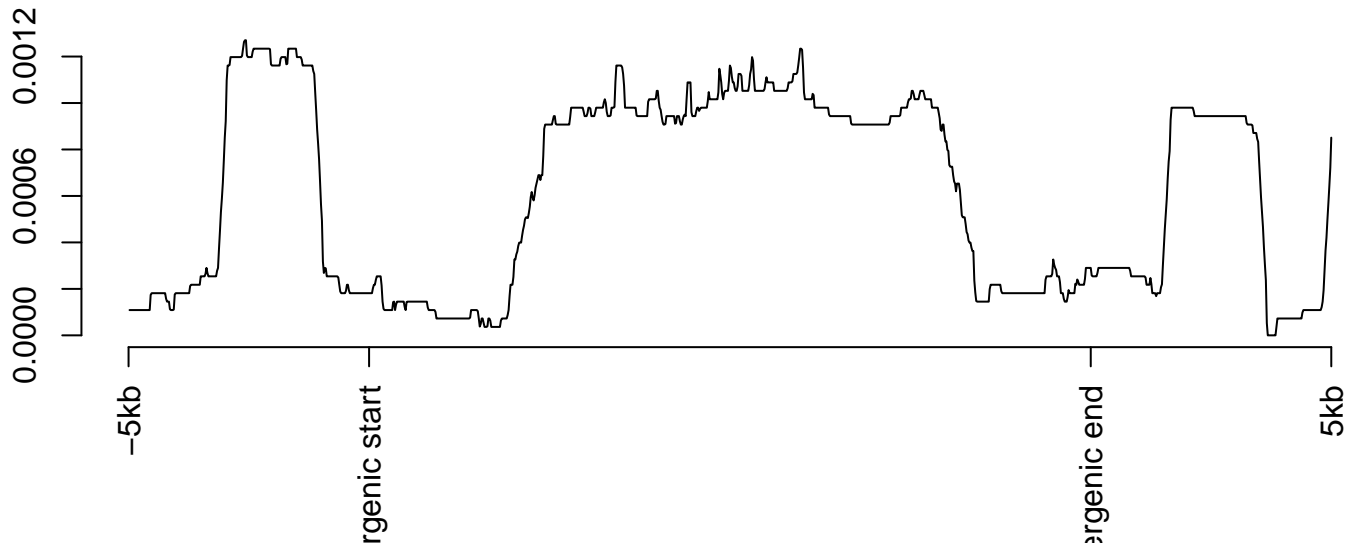
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc529



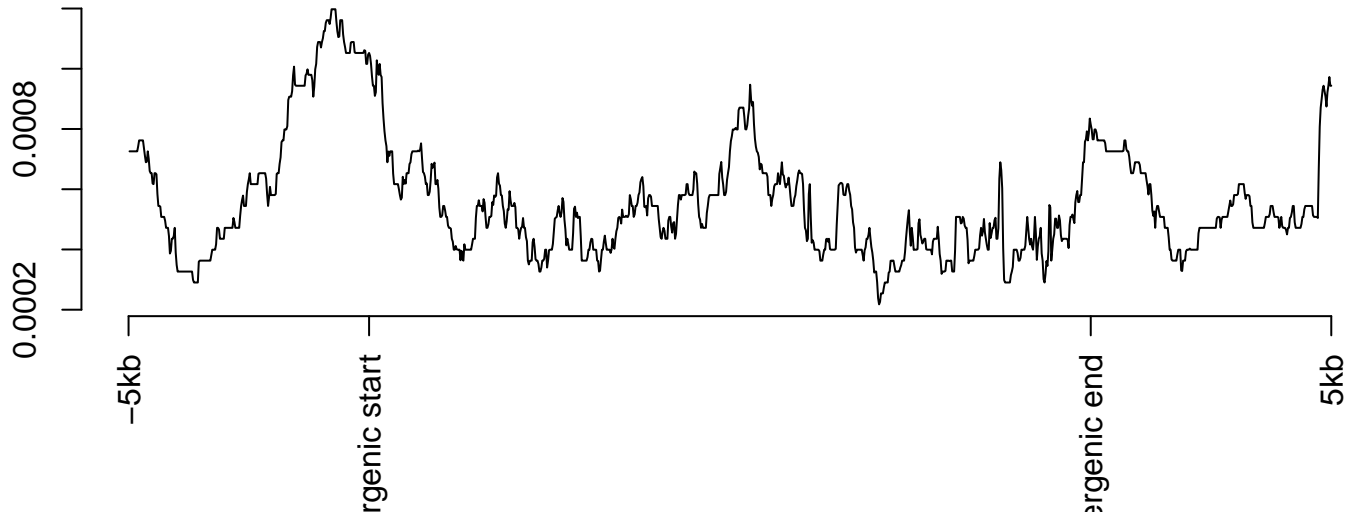
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.623 scCLTdegenNuc530



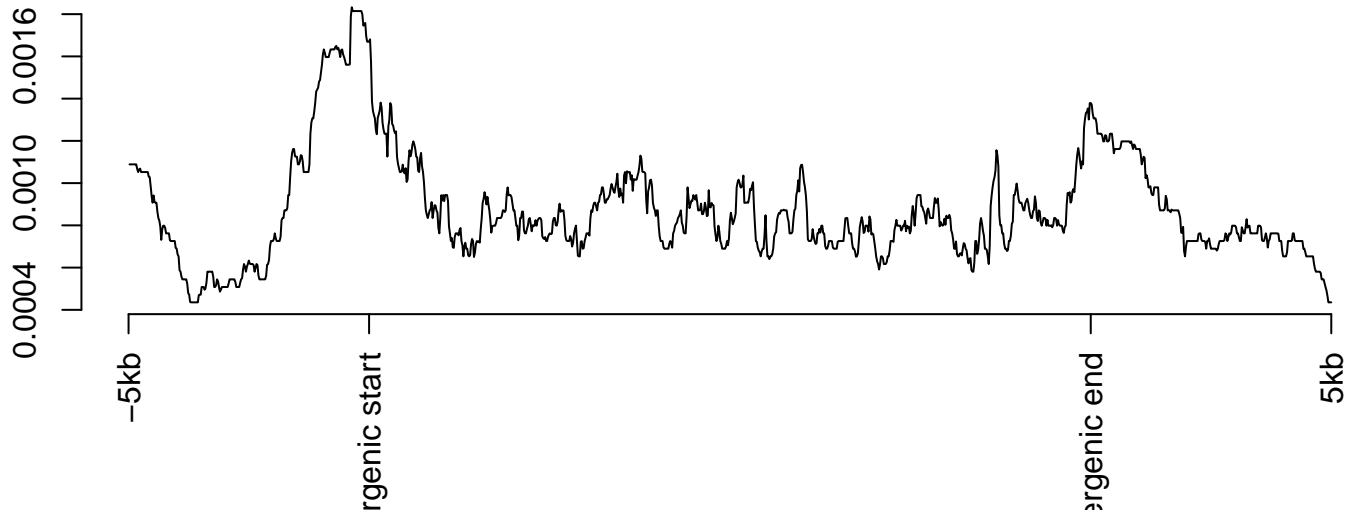
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc531



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc532



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc533

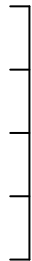
2e-04 6e-04 1e-03

-5kb

intergenic start

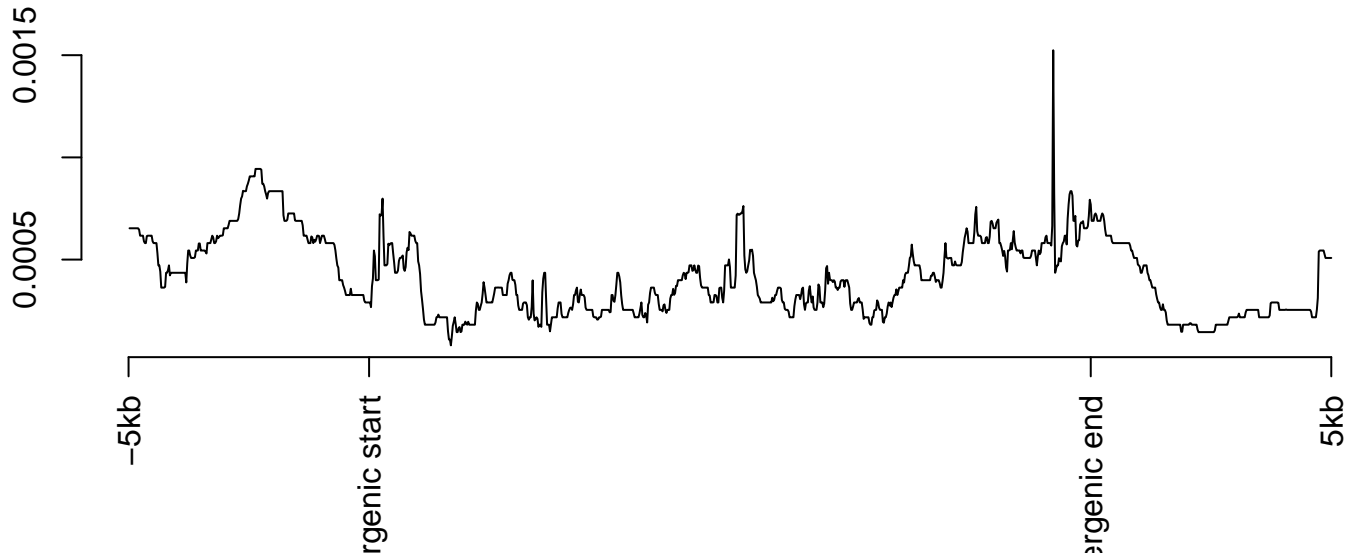
intergenic end

5kb



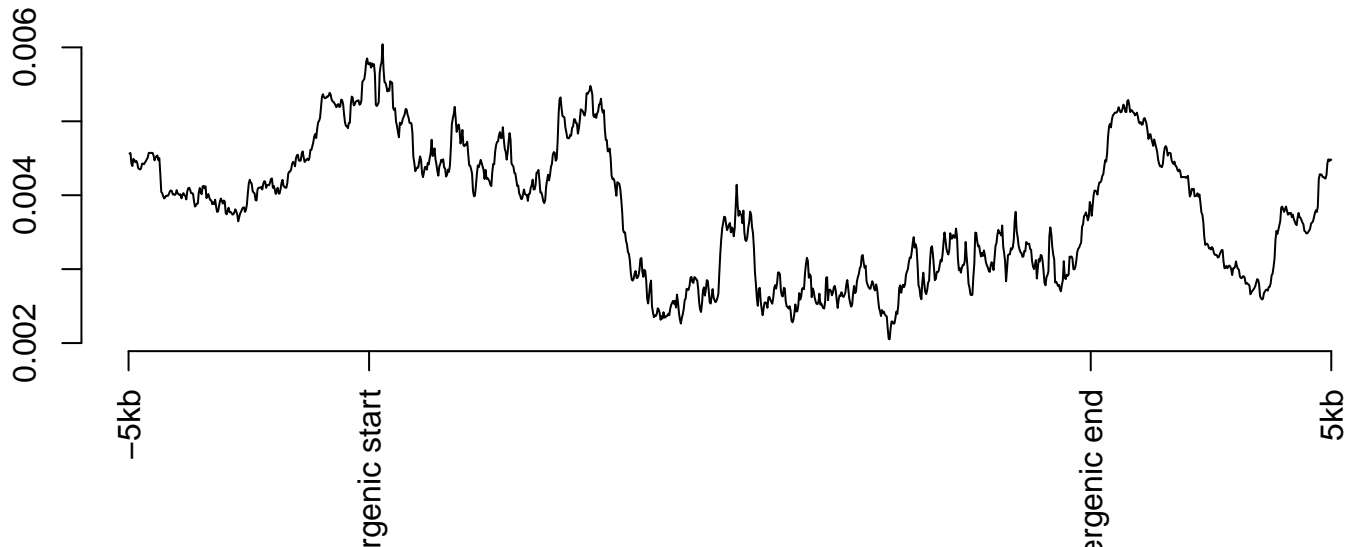
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc534



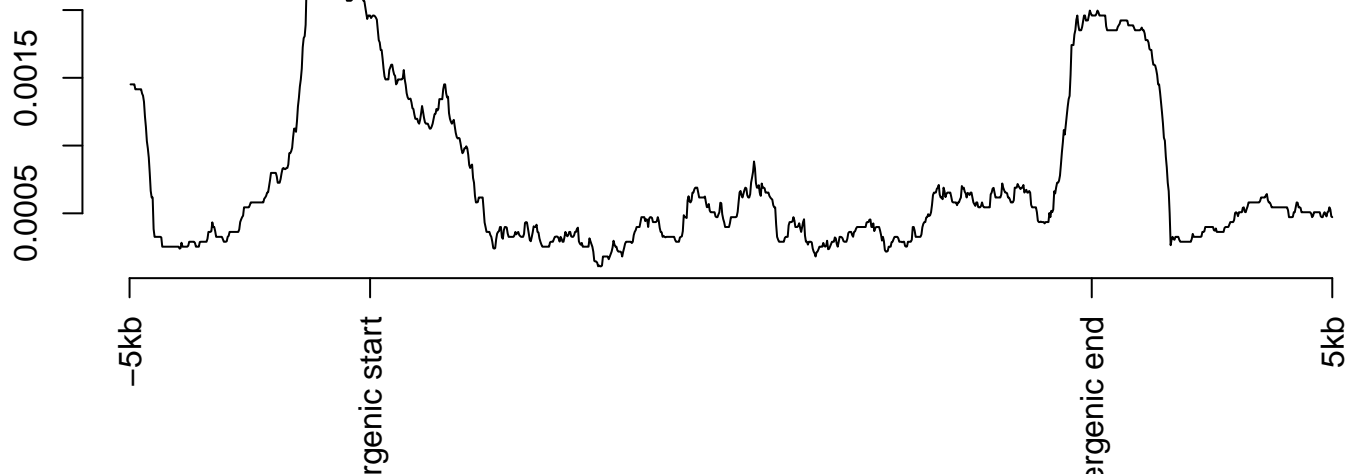
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc535



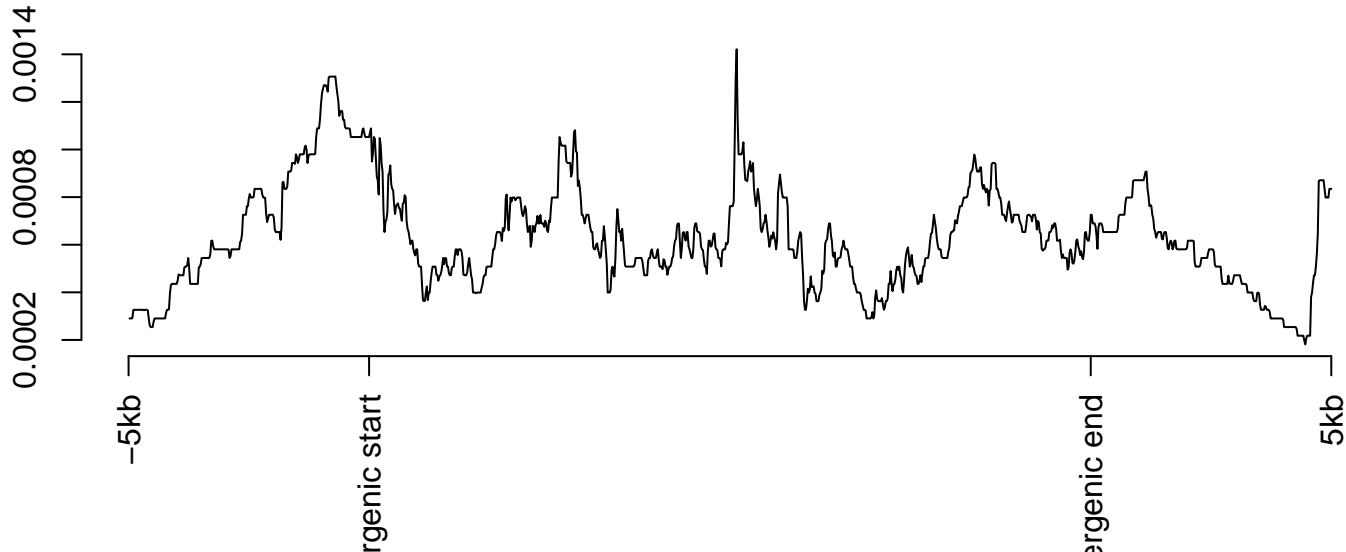
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc536



Average coverage per intergenic region

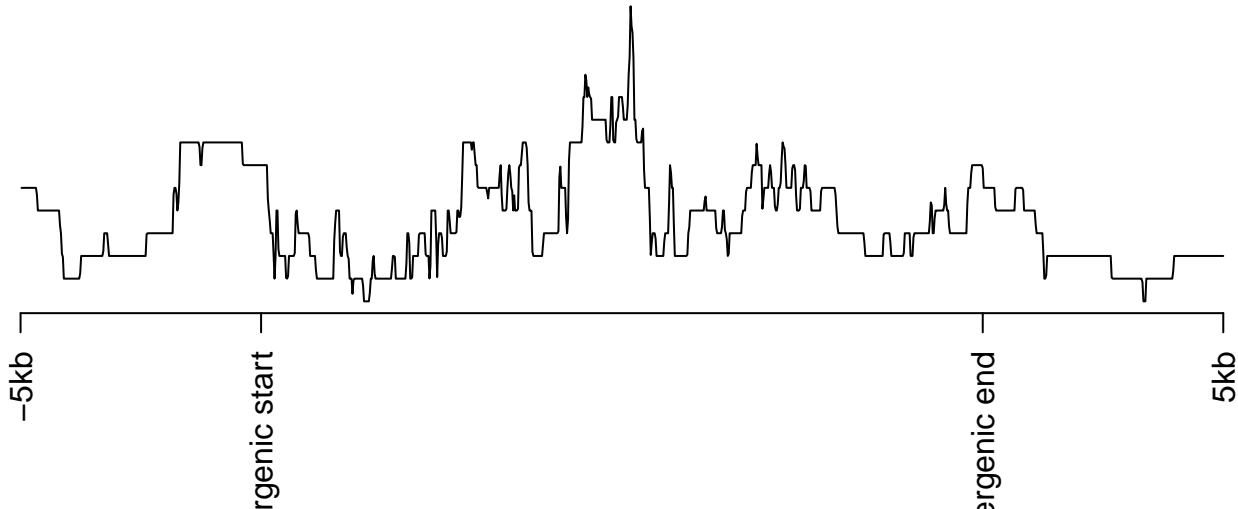
MouseNeuronCulture PositiveSingle E.624 scCLTdegenNuc537



Average coverage per intergenic region

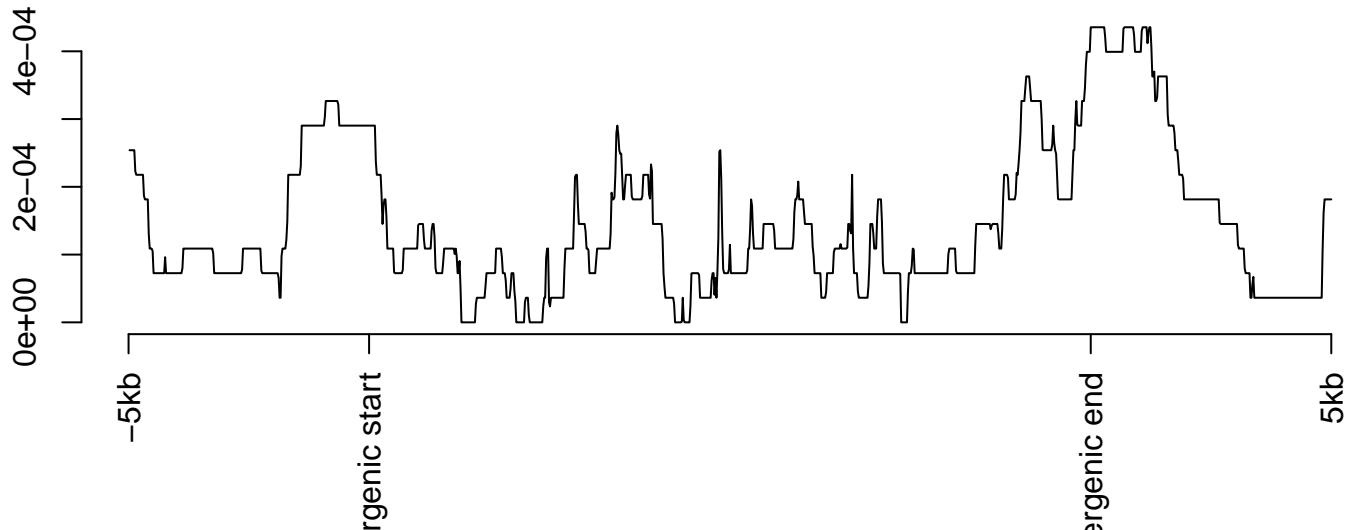
MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc542

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



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc543



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc544

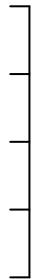
0e+00 2e-04 4e-04

-5kb

intergenic start

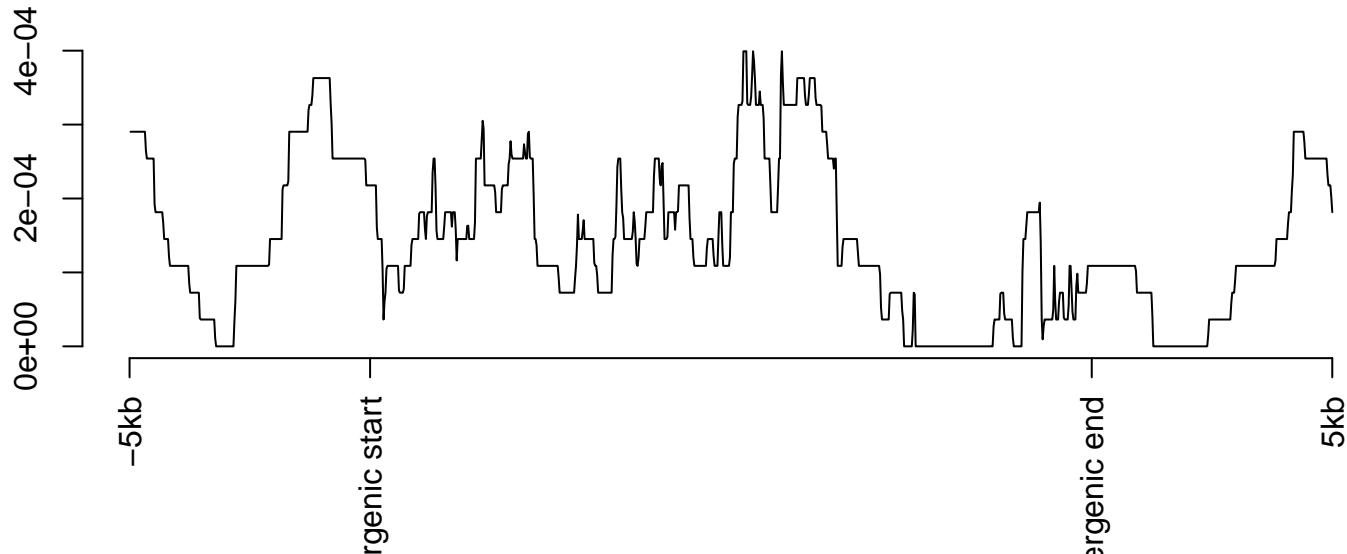
intergenic end

5kb



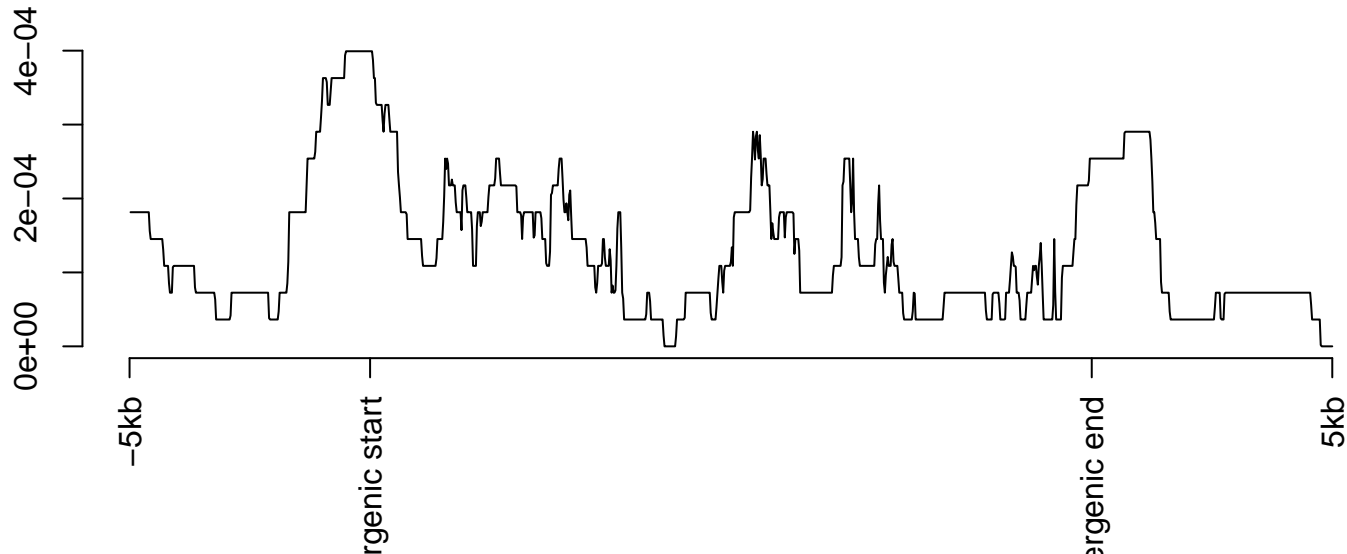
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc545



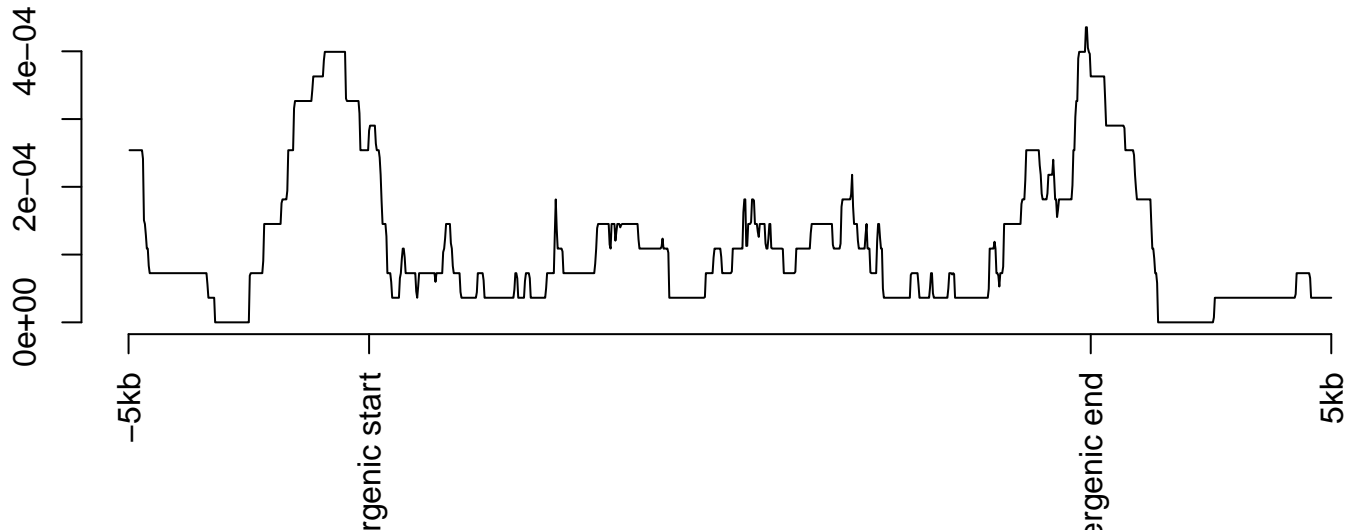
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc546



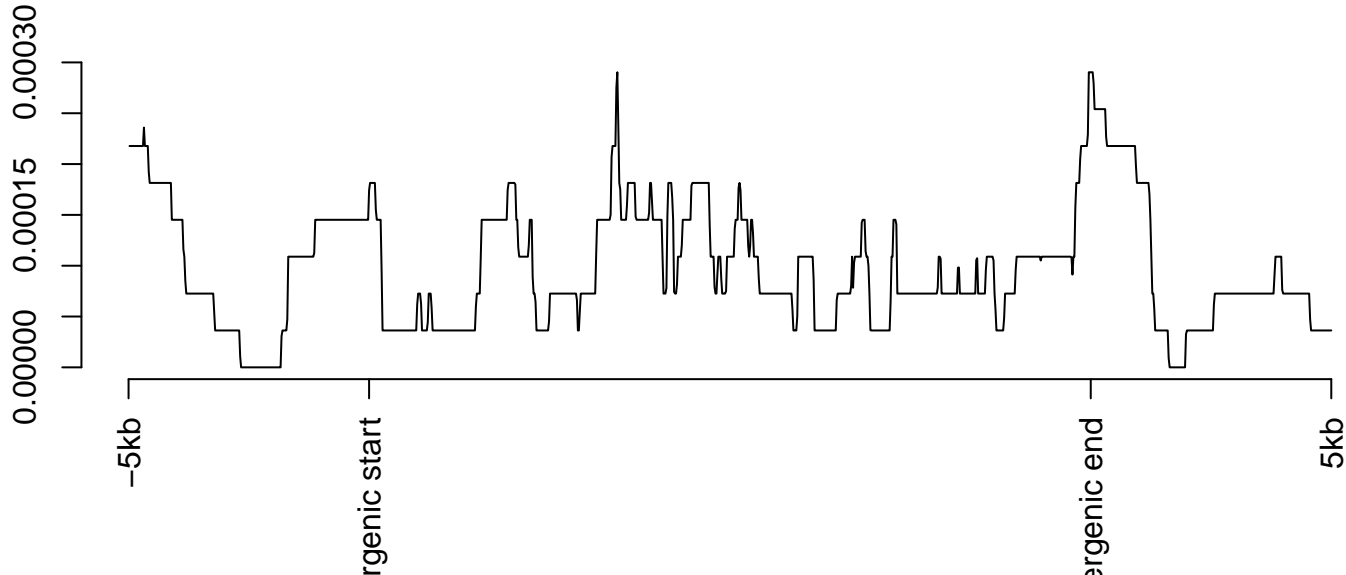
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc547



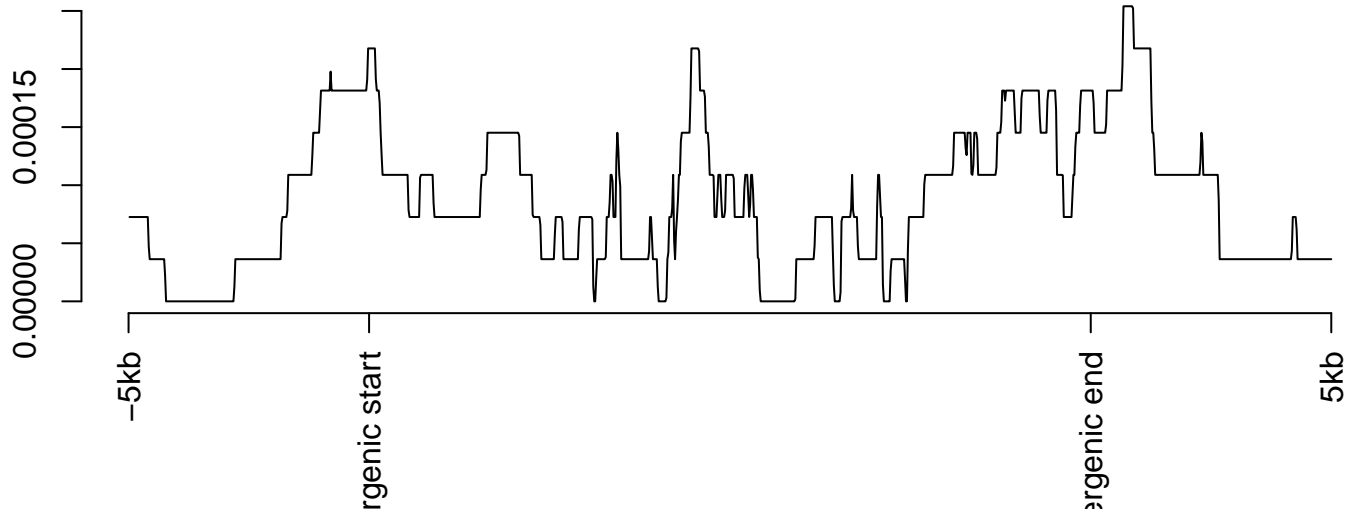
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc548



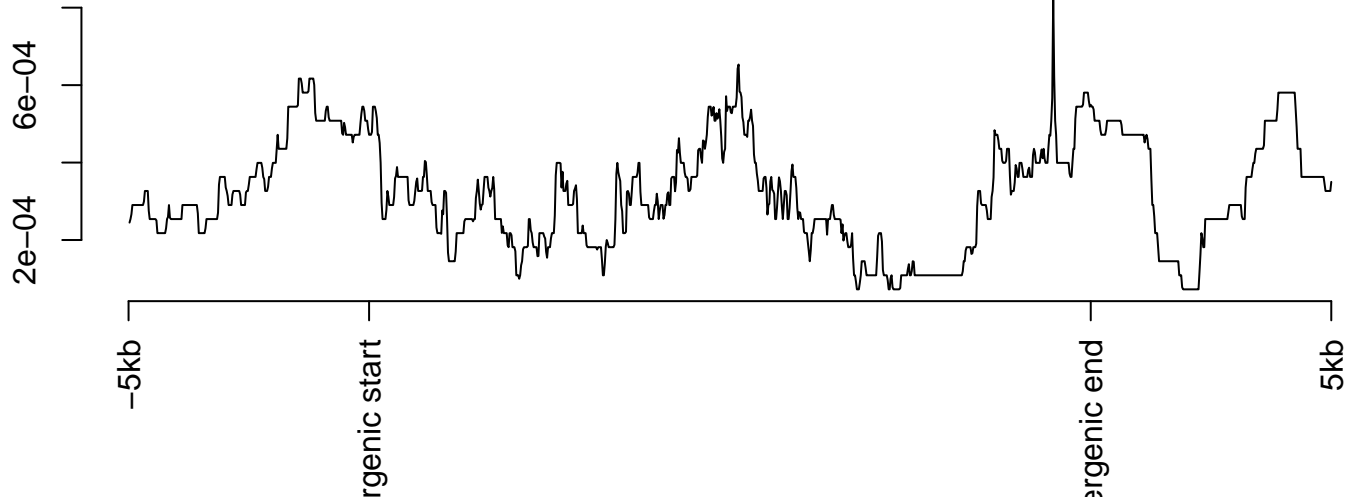
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc549



Average coverage per intergenic region

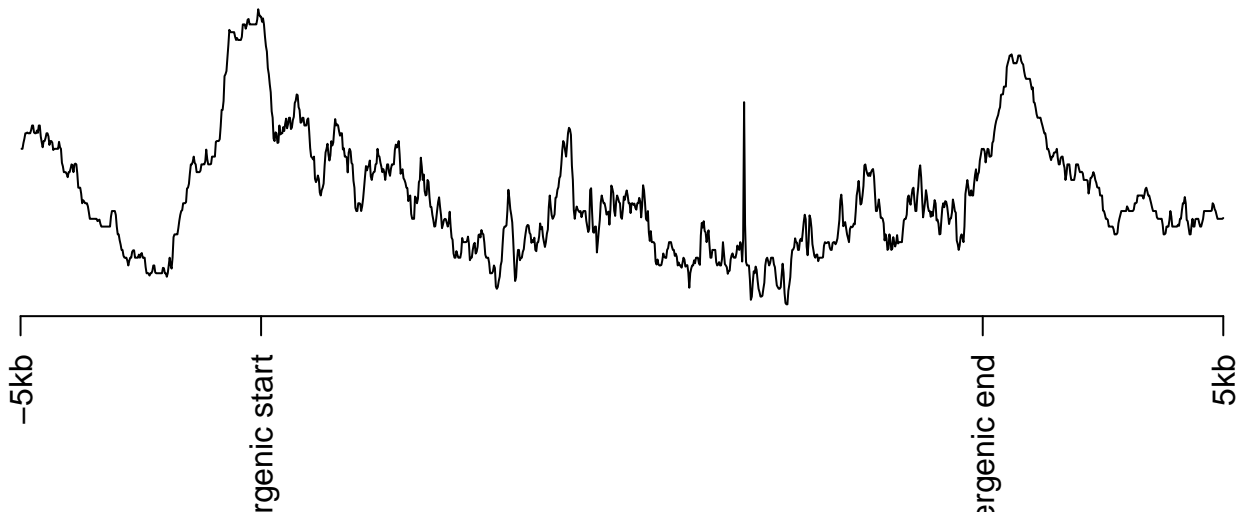
MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc550



MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc551

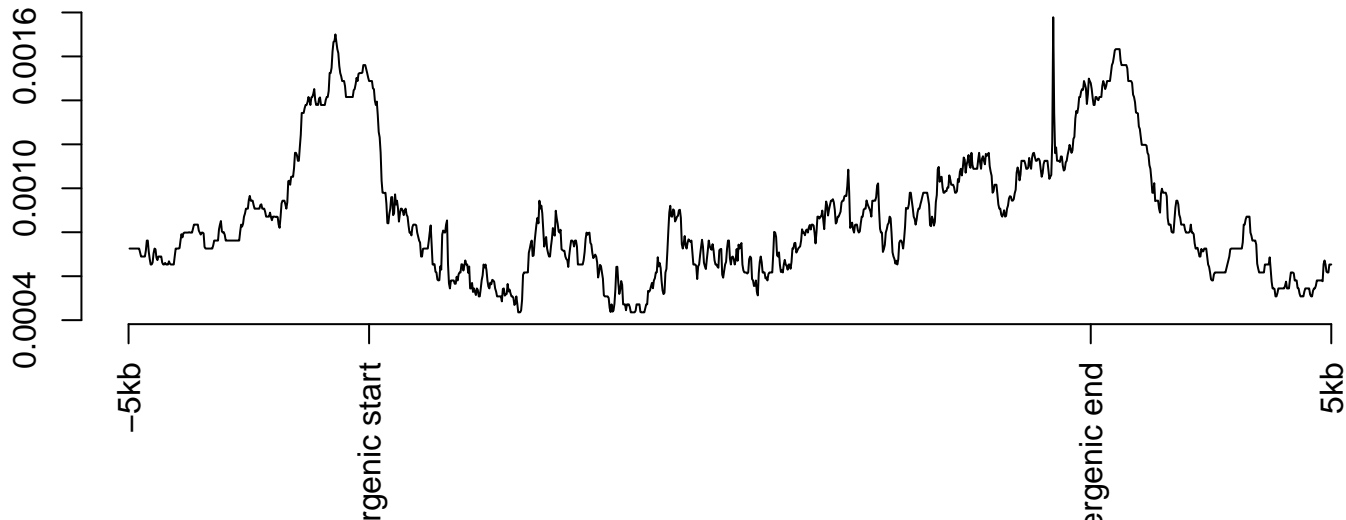
Average coverage per intergenic region

0.0004 0.0010 0.0016



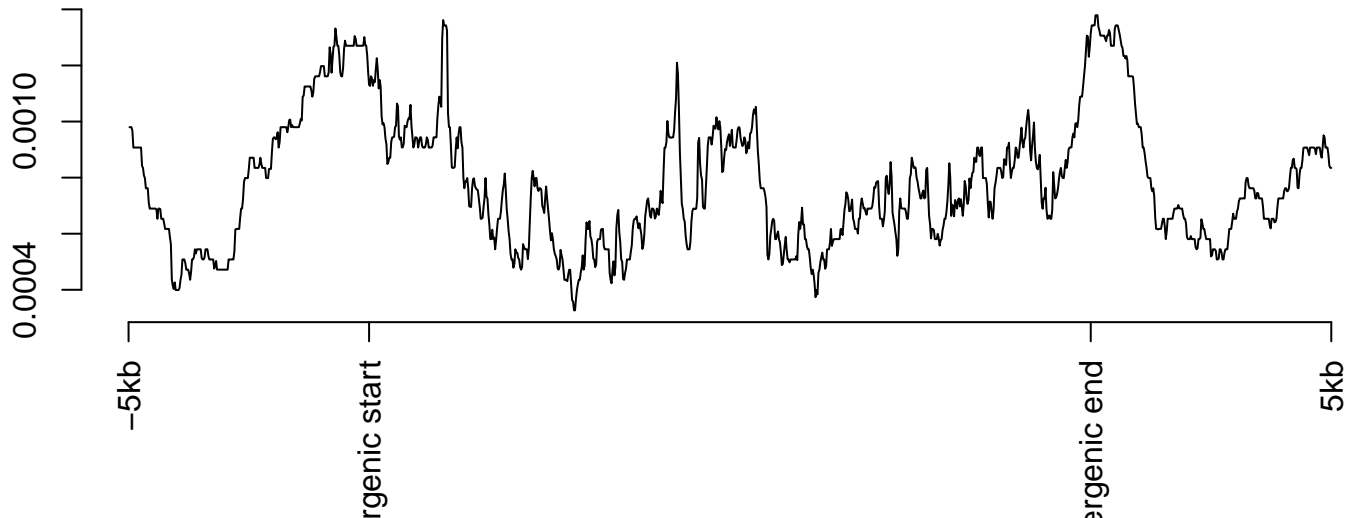
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc552



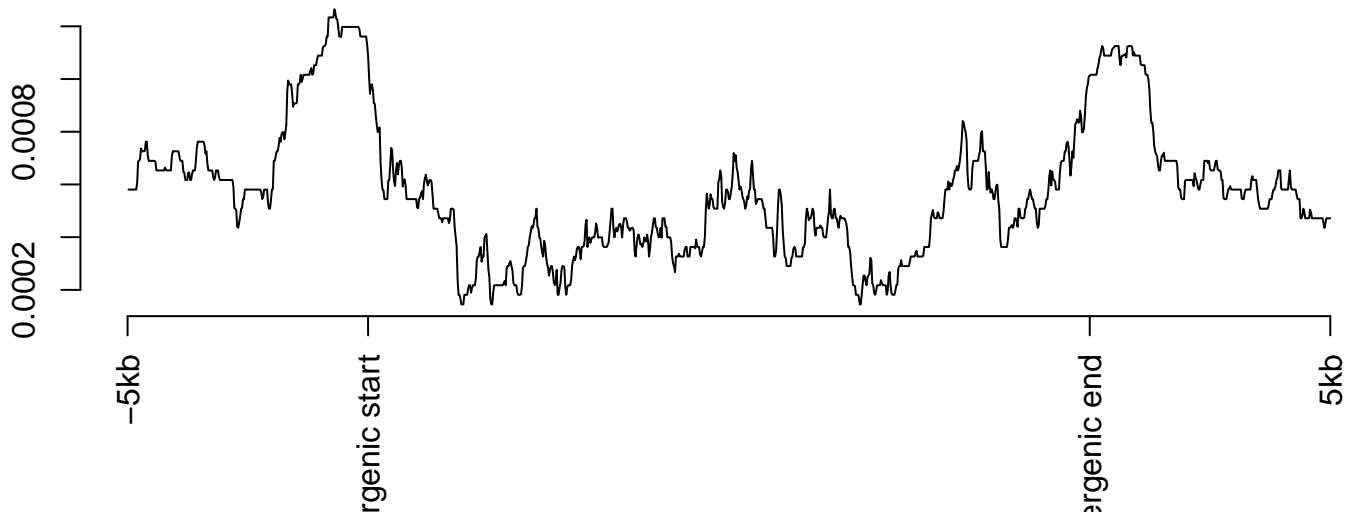
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.627 scCLTdegenNuc553



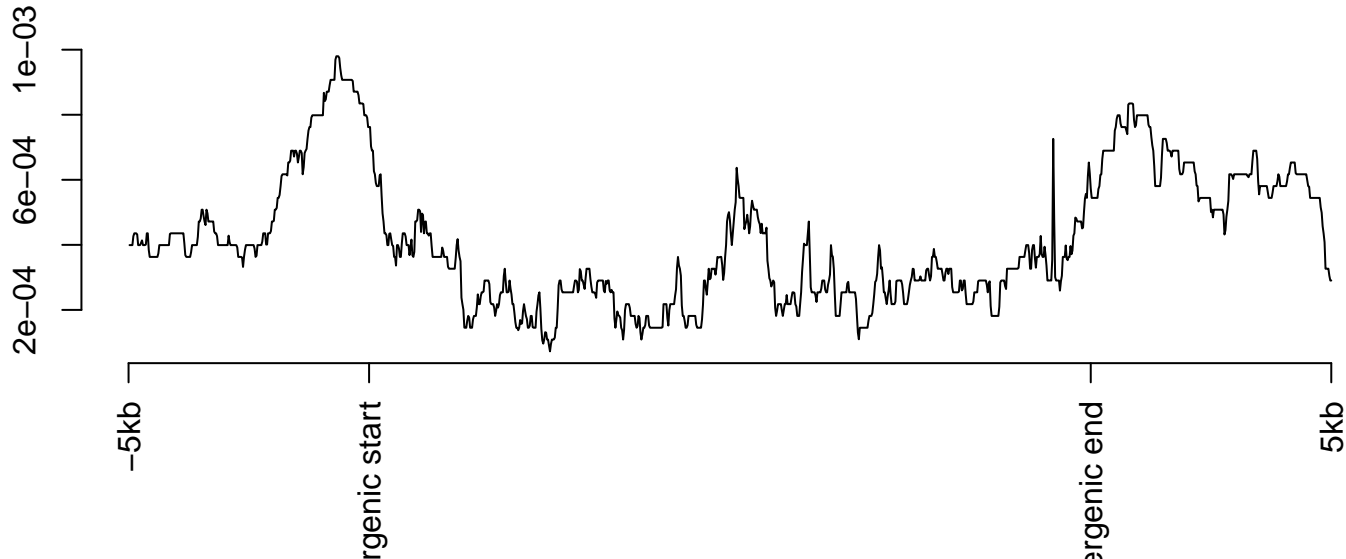
MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc554

Average coverage per intergenic region



Average coverage per intergenic region

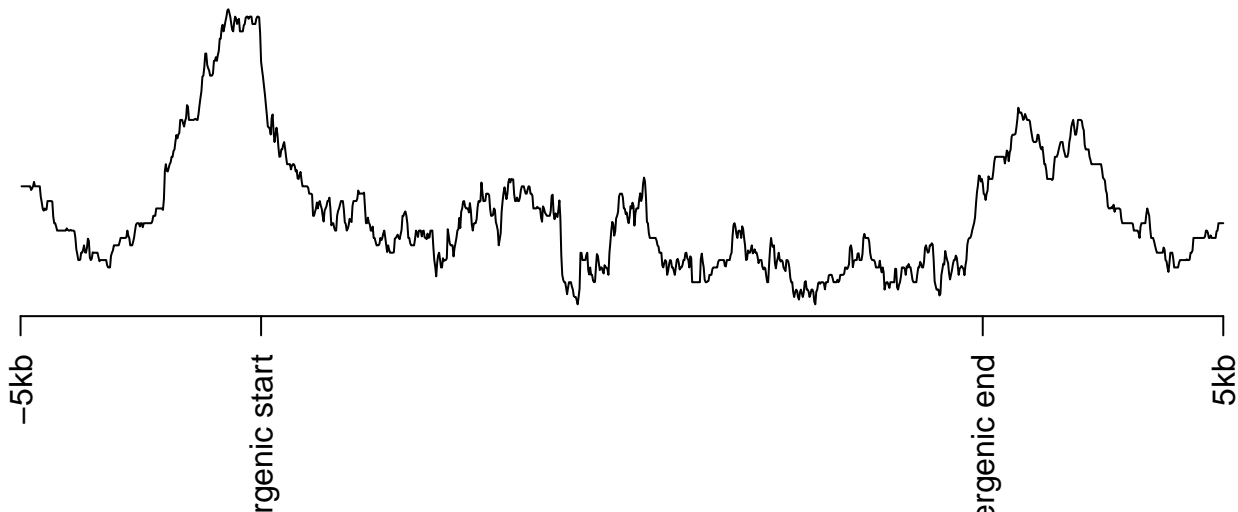
MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc555



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc556

Average coverage per intergenic region

0.0015
0.0005



MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc557

Average coverage per intergenic region

0.0015
0.0005

-5kb

intergenic start

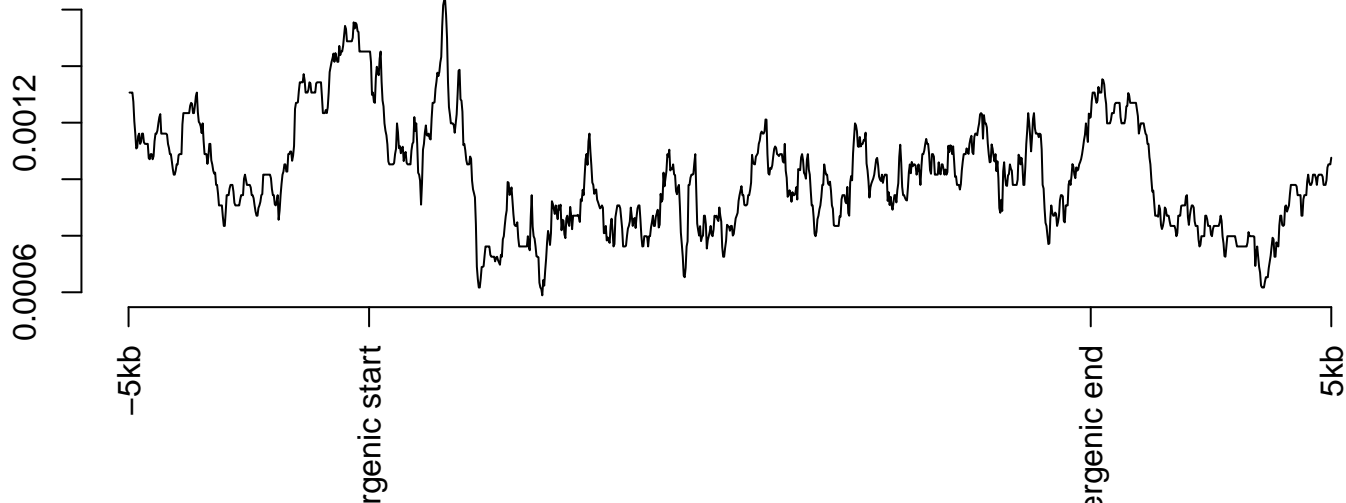
intergenic end

5kb



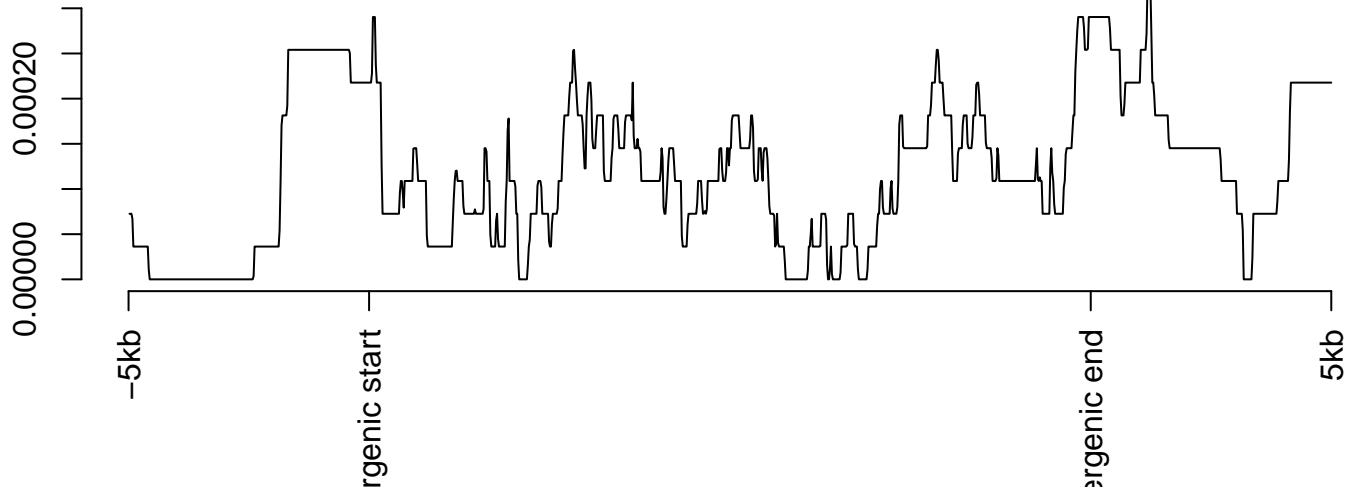
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc558



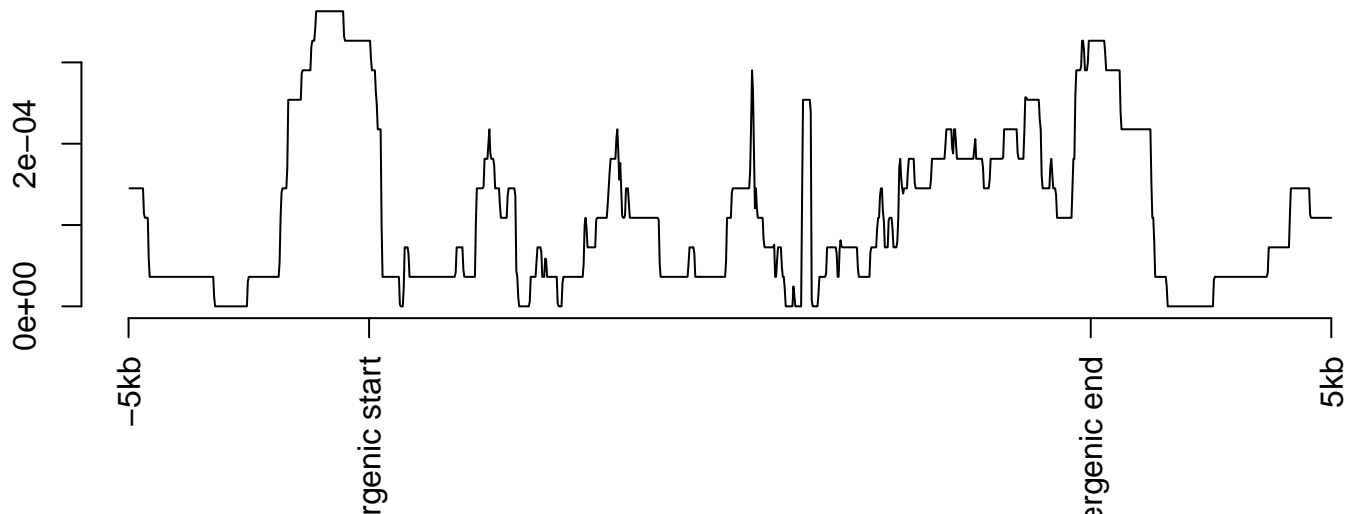
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc559



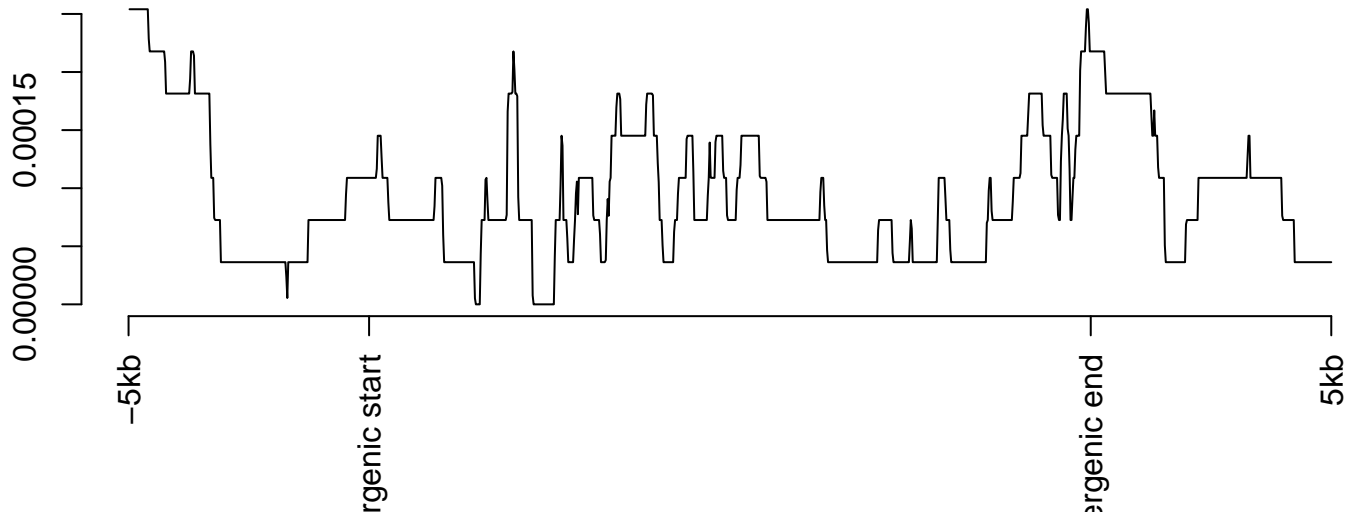
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc561



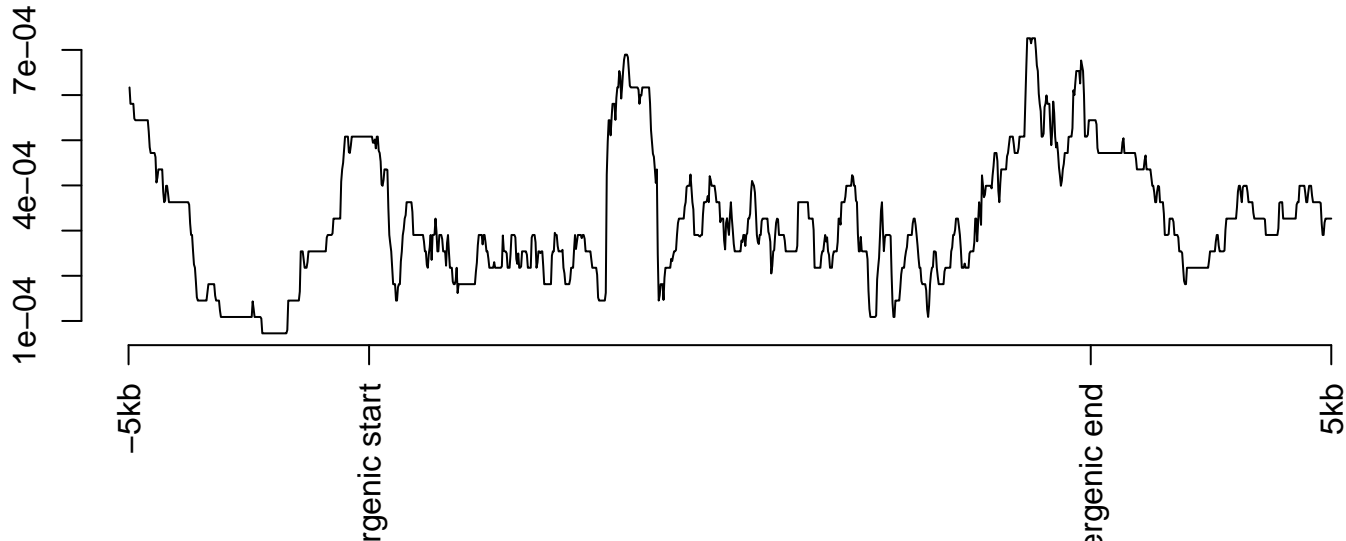
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc562



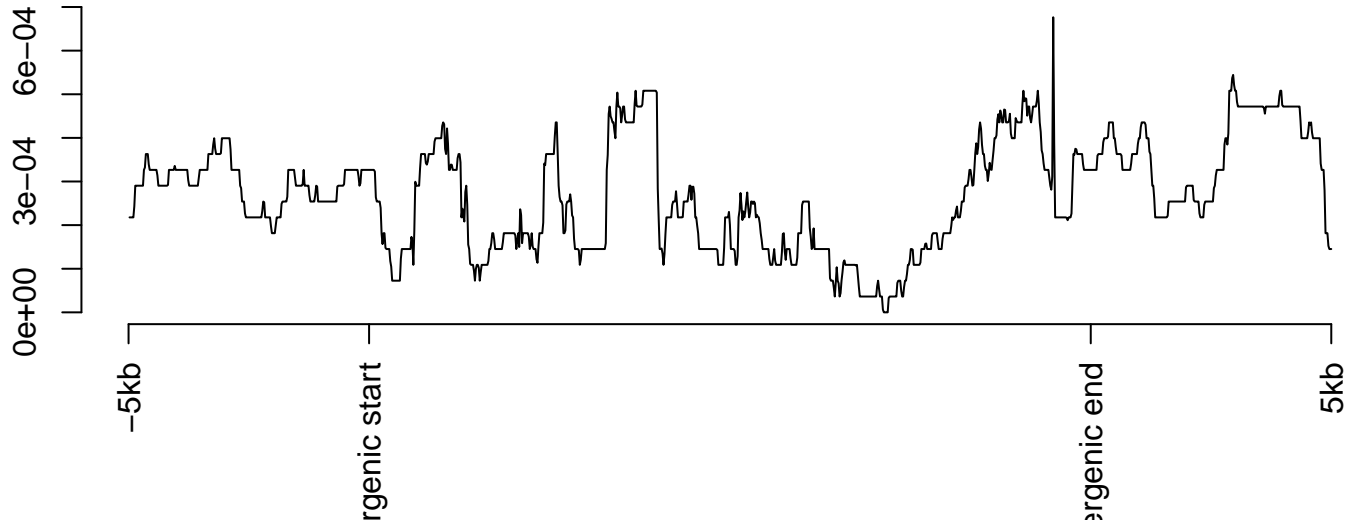
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc563



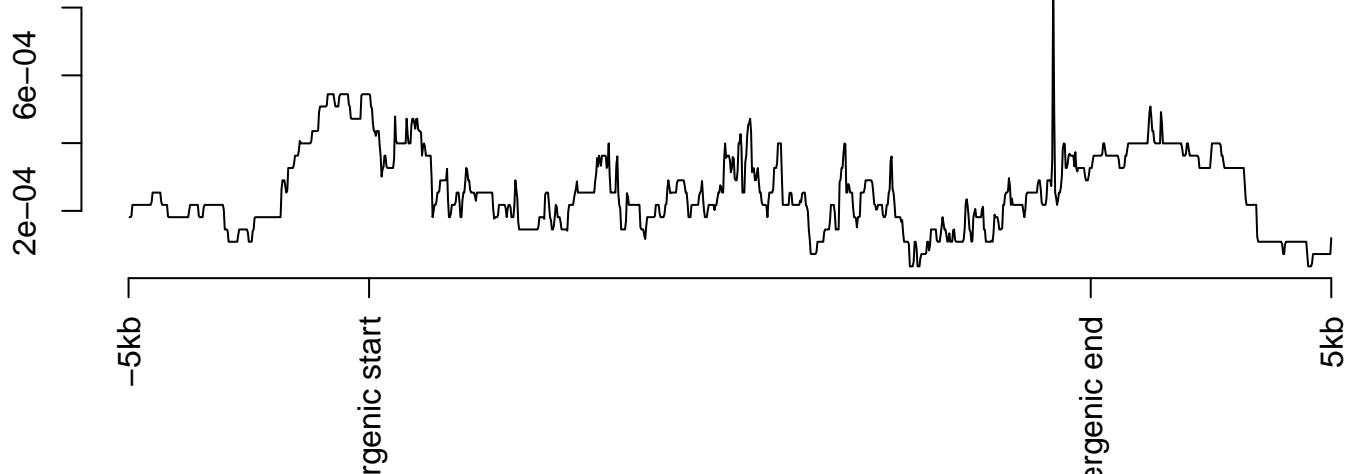
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc564



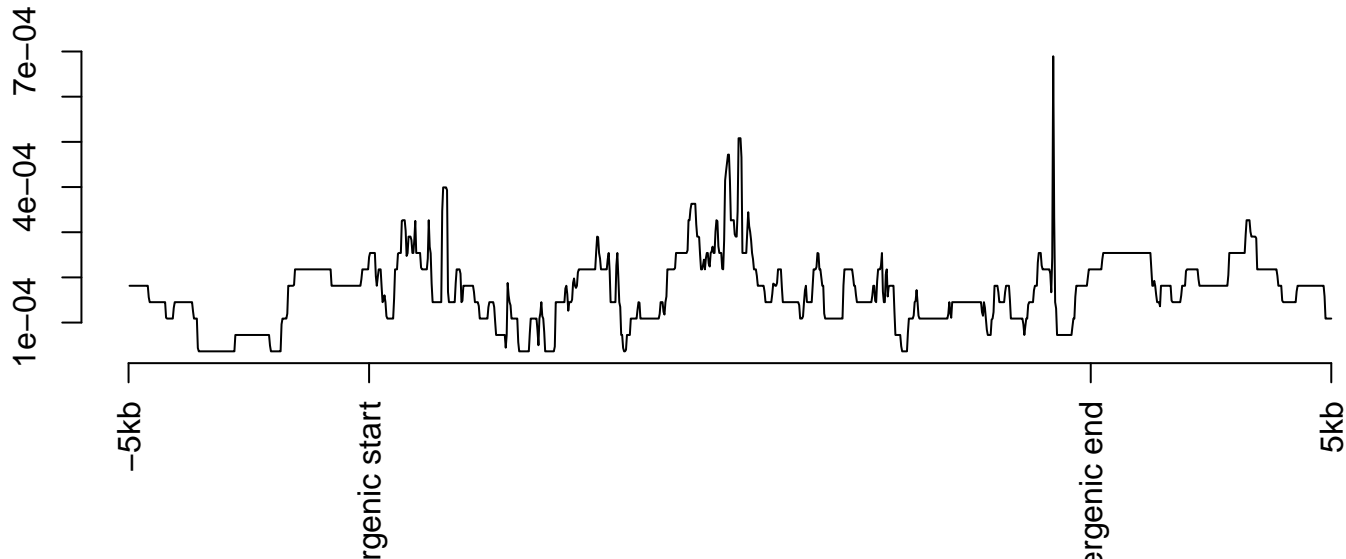
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc565



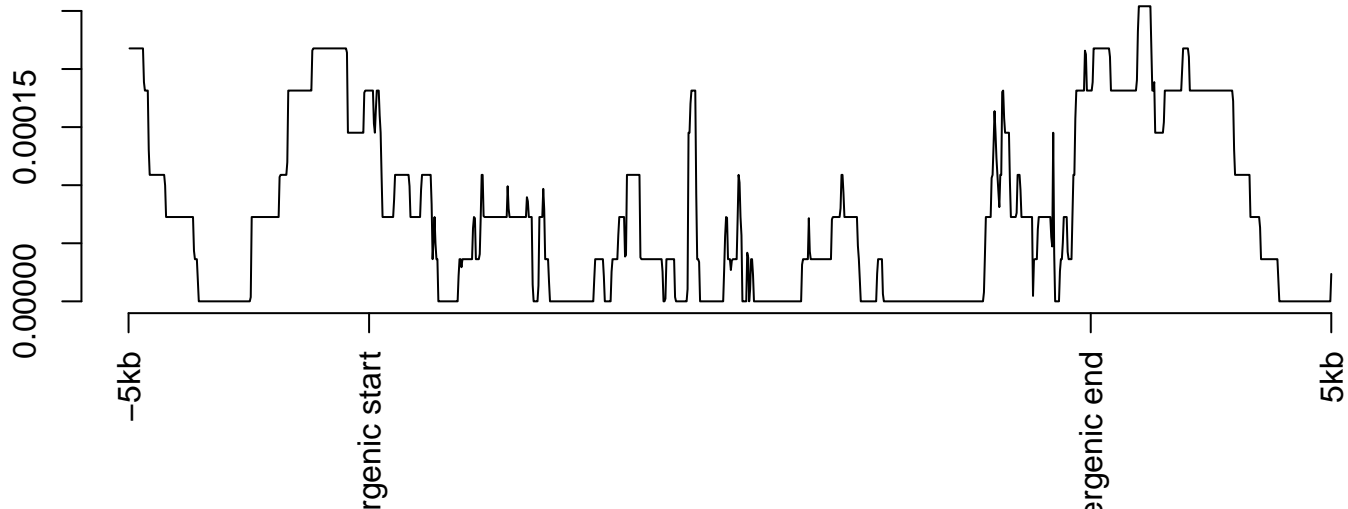
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.628 scCLTdegenNuc566



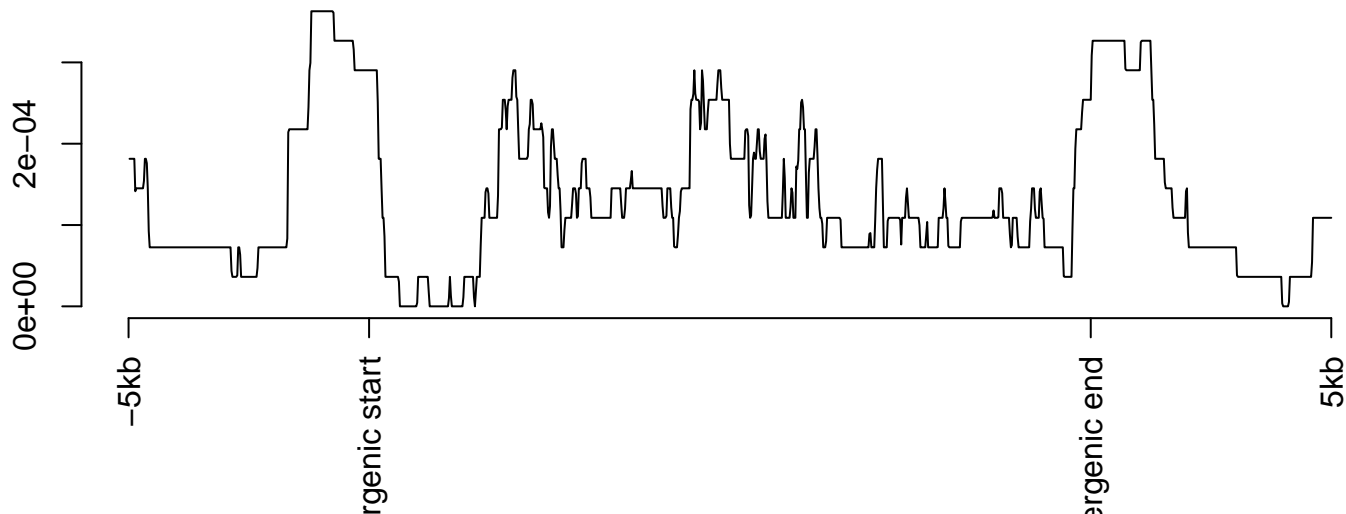
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc567



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc568



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc569

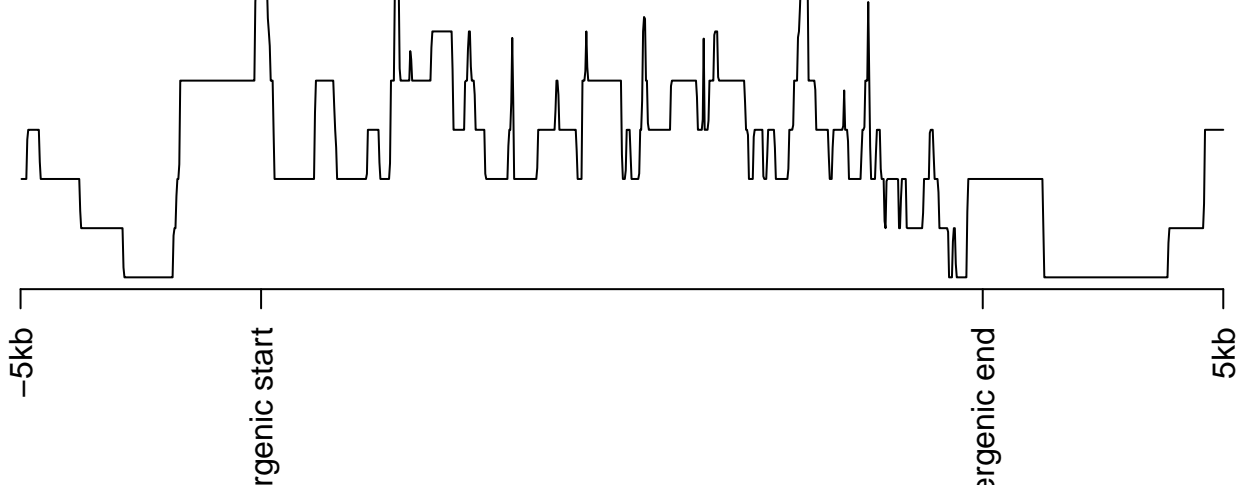
0.00015
0.00000

-5kb

intergenic start

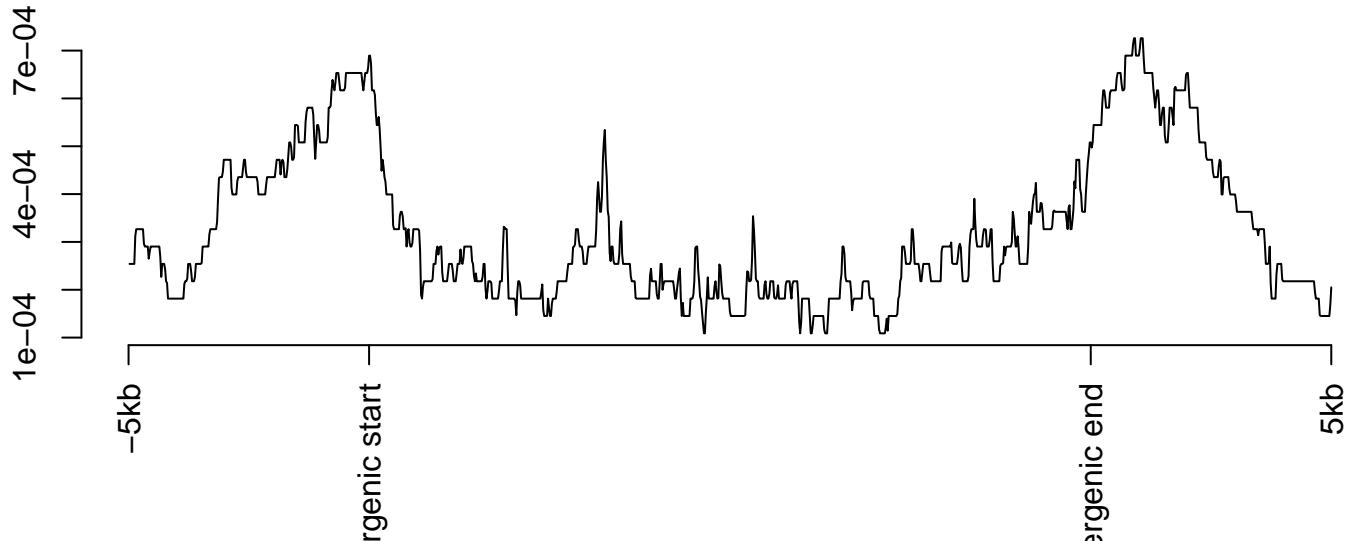
intergenic end

5kb



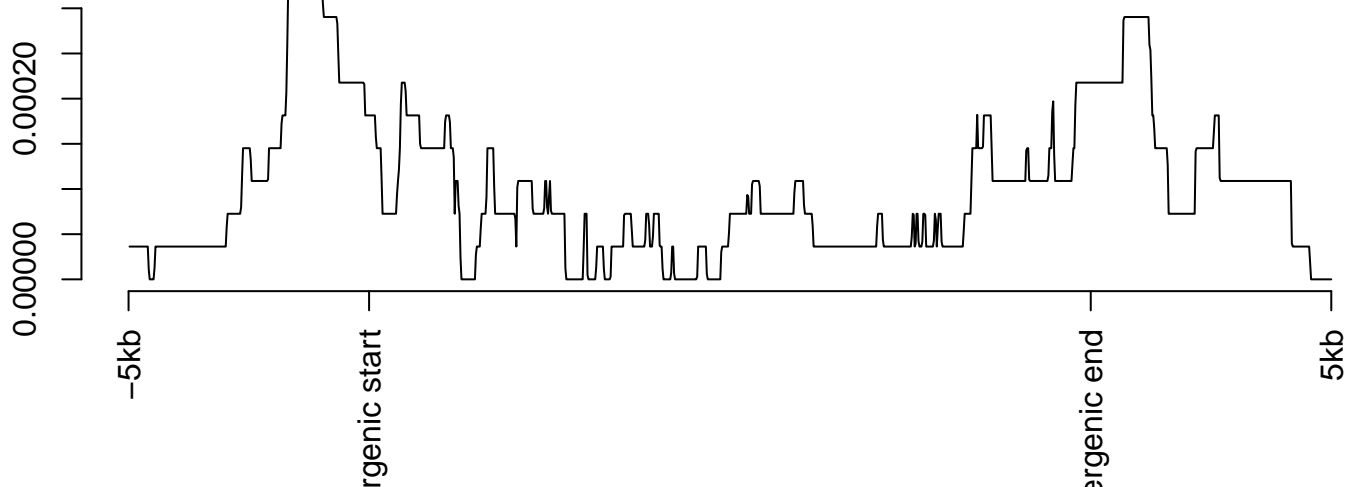
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc570



Average coverage per intergenic region

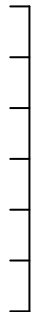
MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc571



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc572

0.00030
0.00015
0.00000



-5kb

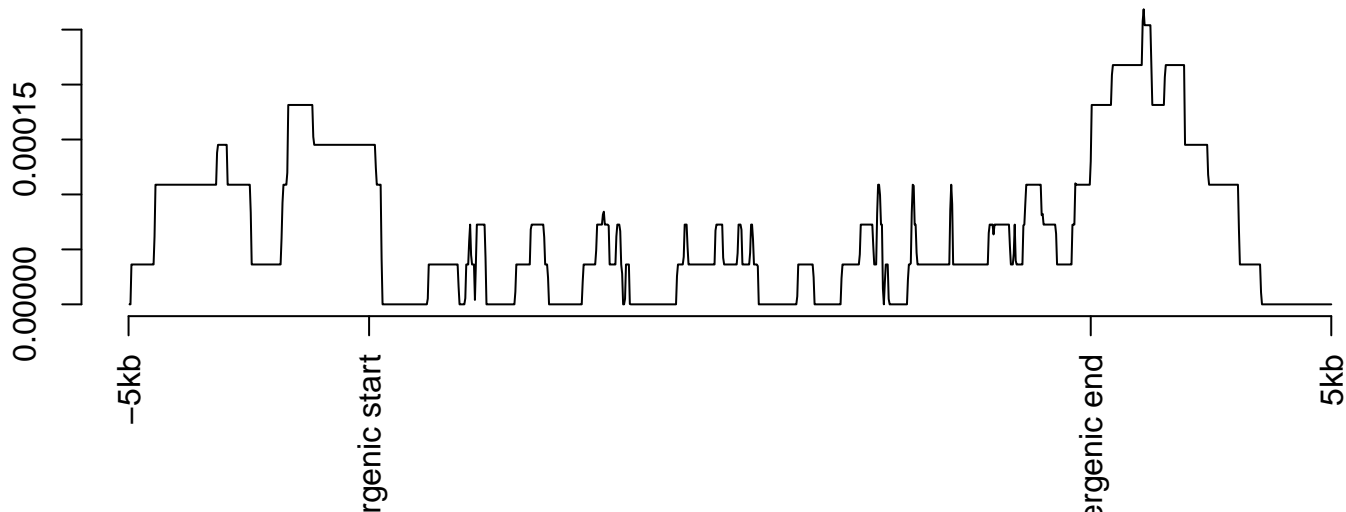
intergenic start

intergenic end

5kb

MouseNeuronCulture PositiveSingle E.629 scCLTdegenNuc573

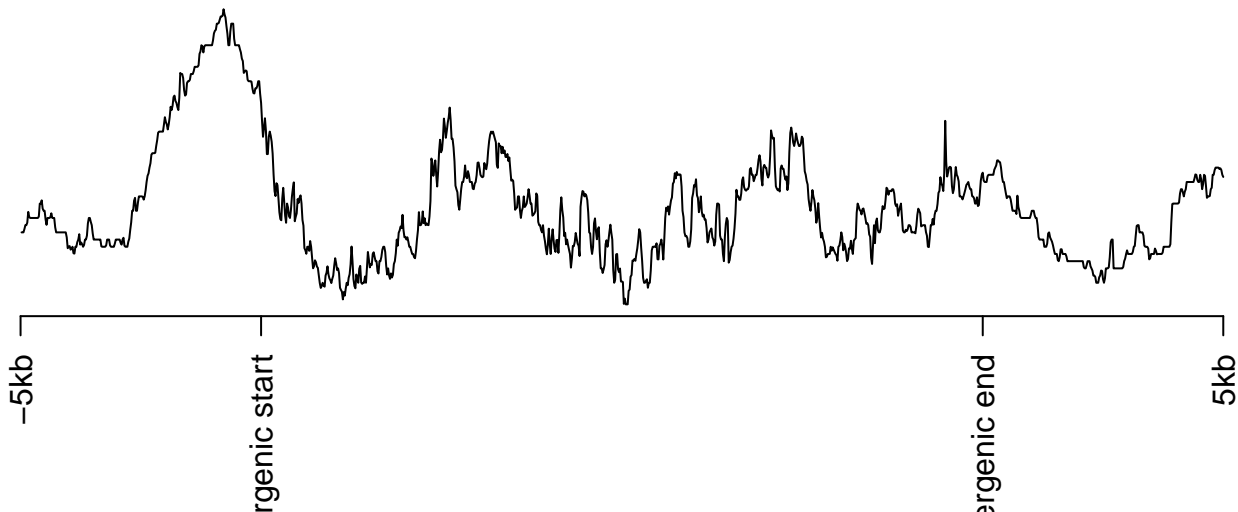
Average coverage per intergenic region



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc576

0.0015
0.0005



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc601

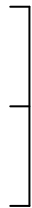
0.0020
0.0010

-5kb

intergenic start

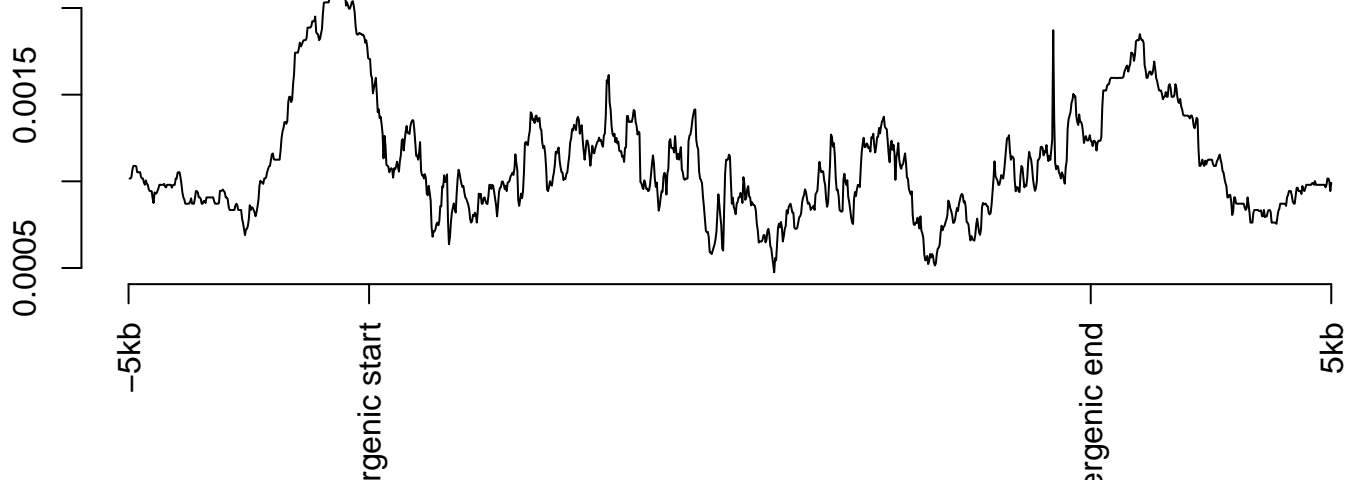
intergenic end

5kb



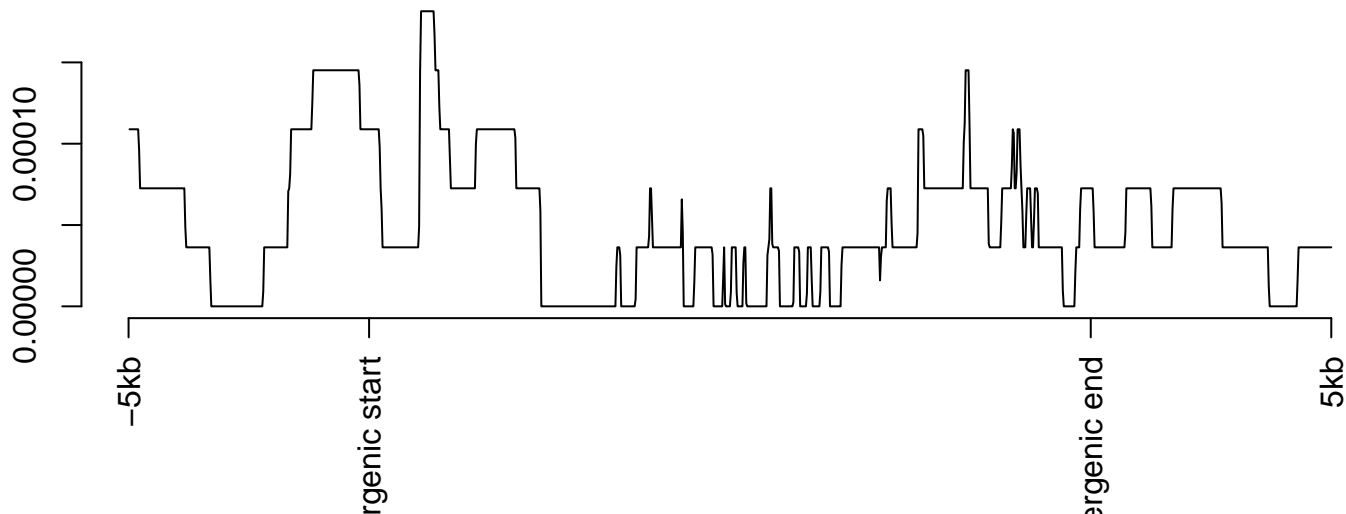
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.643 scCLTdegenNuc602



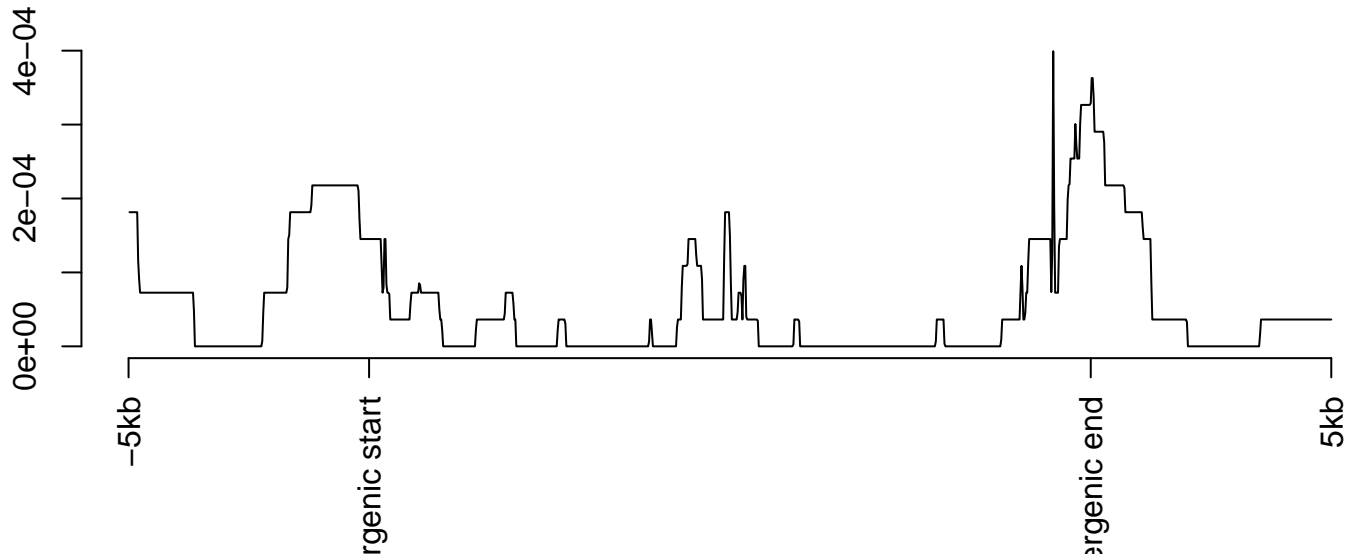
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc577b



Average coverage per intergenic region

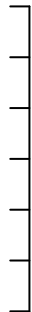
MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc578b



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc579b

0.00030
0.00015
0.00000



-5kb

intergenic start

intergenic end

5kb

Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc580b

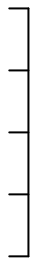
0.0020
0.0005

-5kb

intergenic start

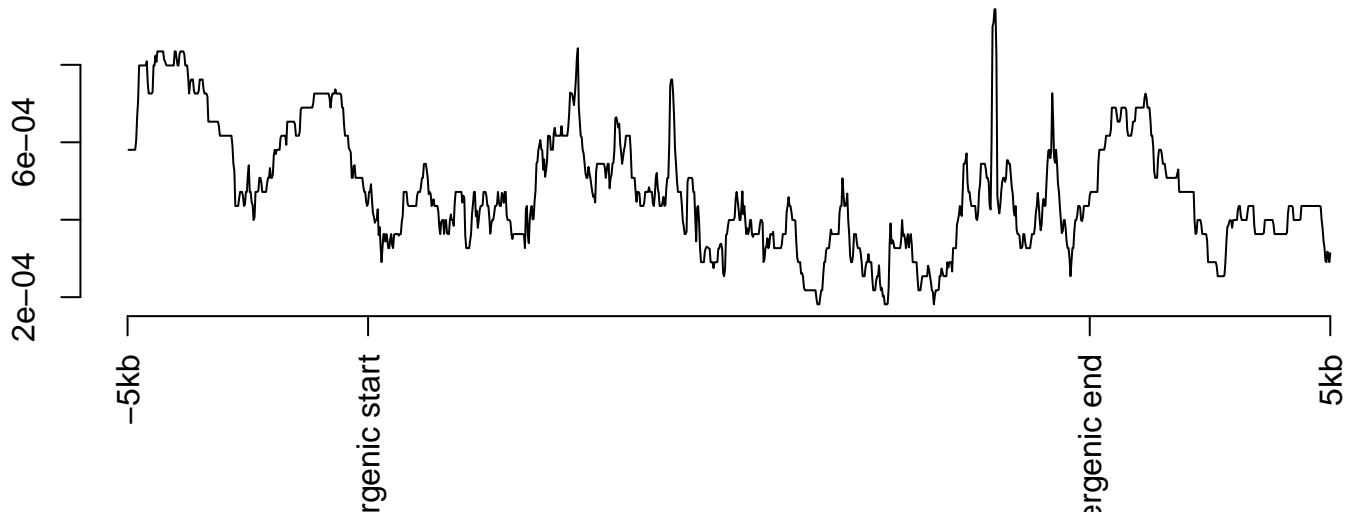
intergenic end

5kb



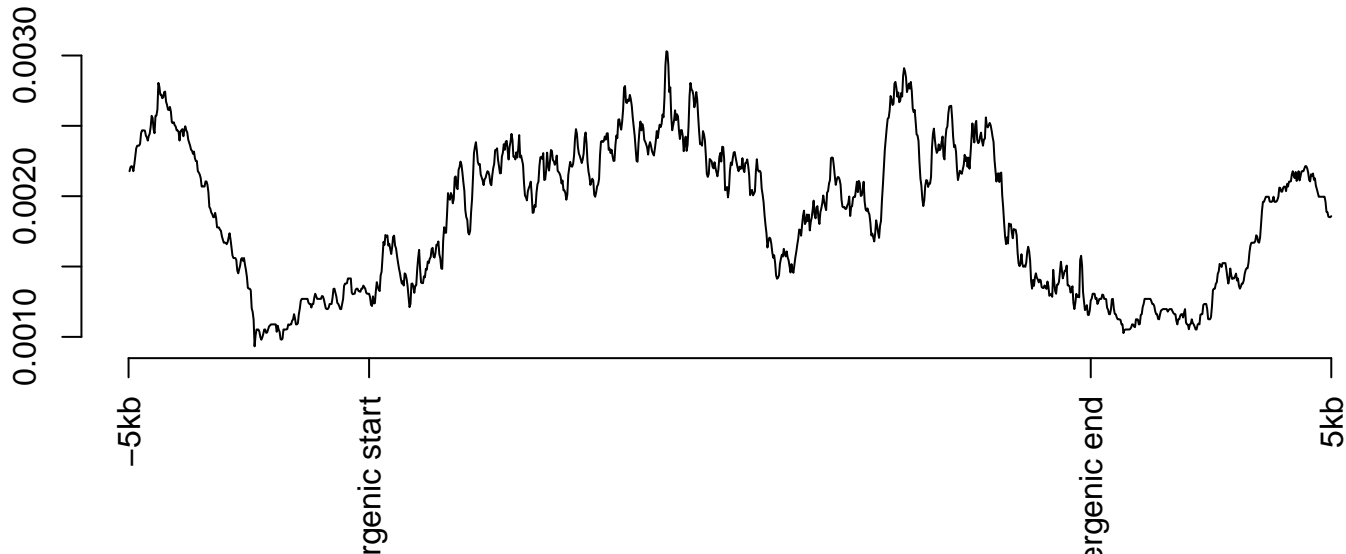
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc581b



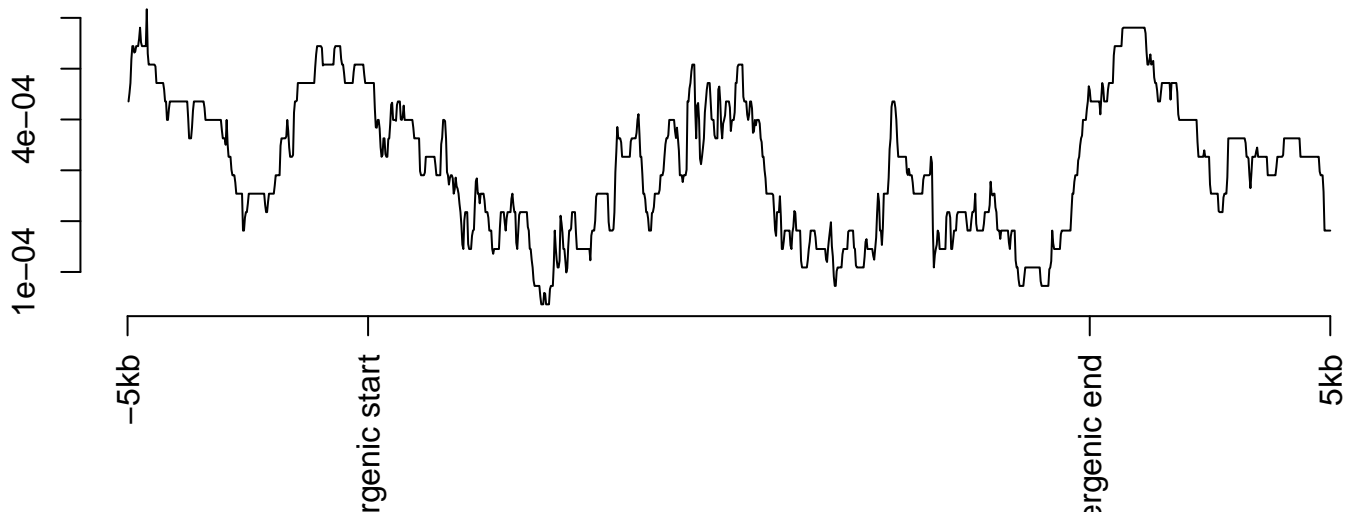
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc582b



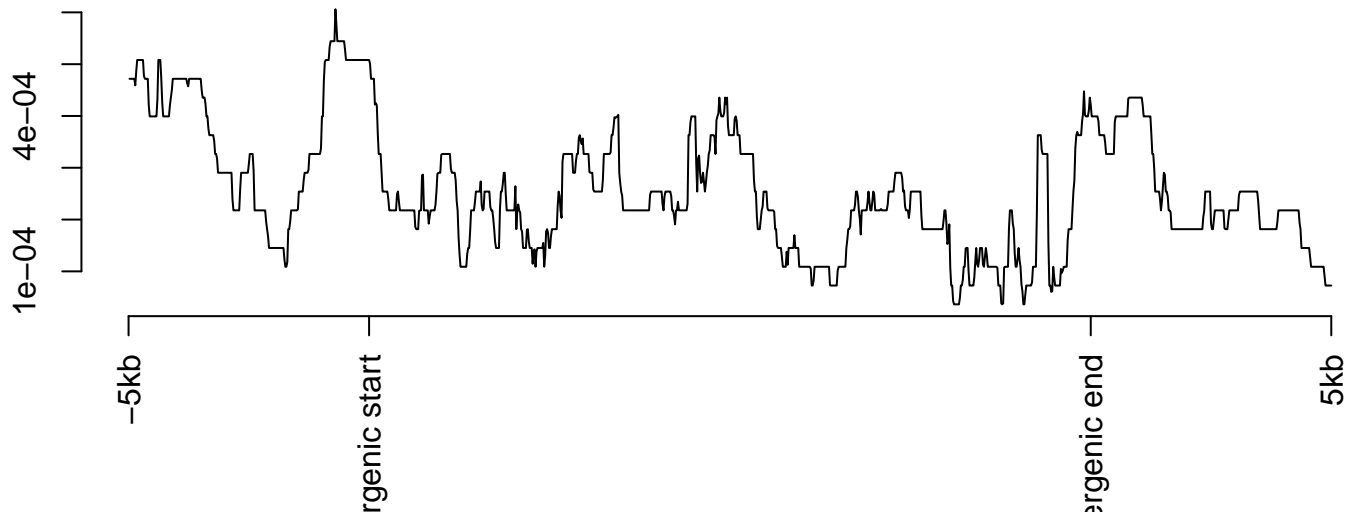
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc583b



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc584b



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc585b

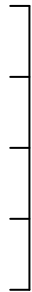
1e-03
6e-04
2e-04

-5kb

intergenic start

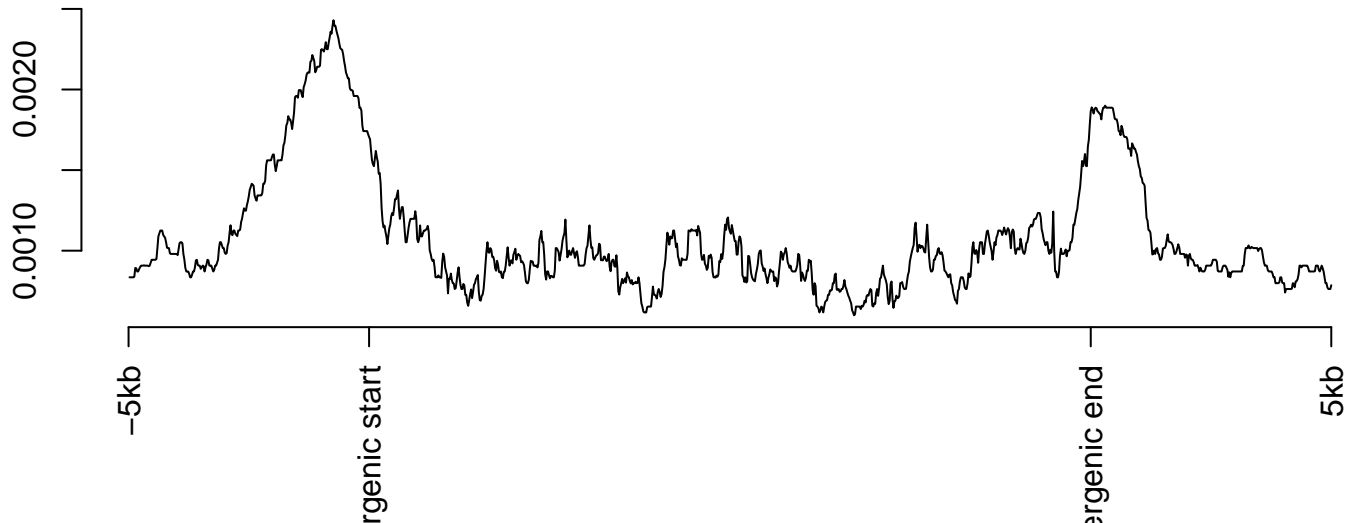
intergenic end

5kb



Average coverage per intergenic region

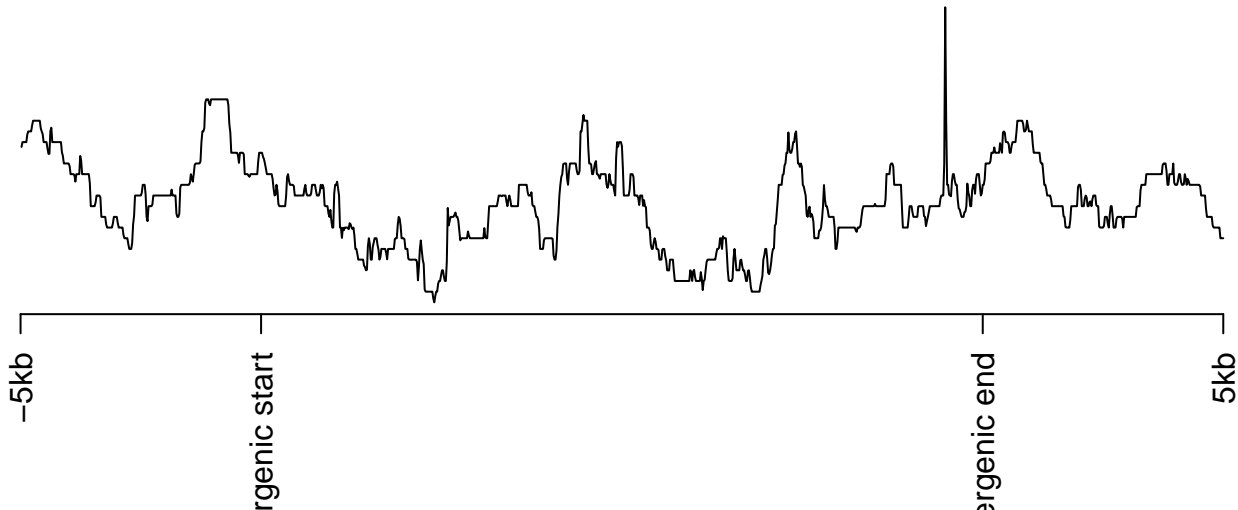
MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc586b



Average coverage per intergenic region

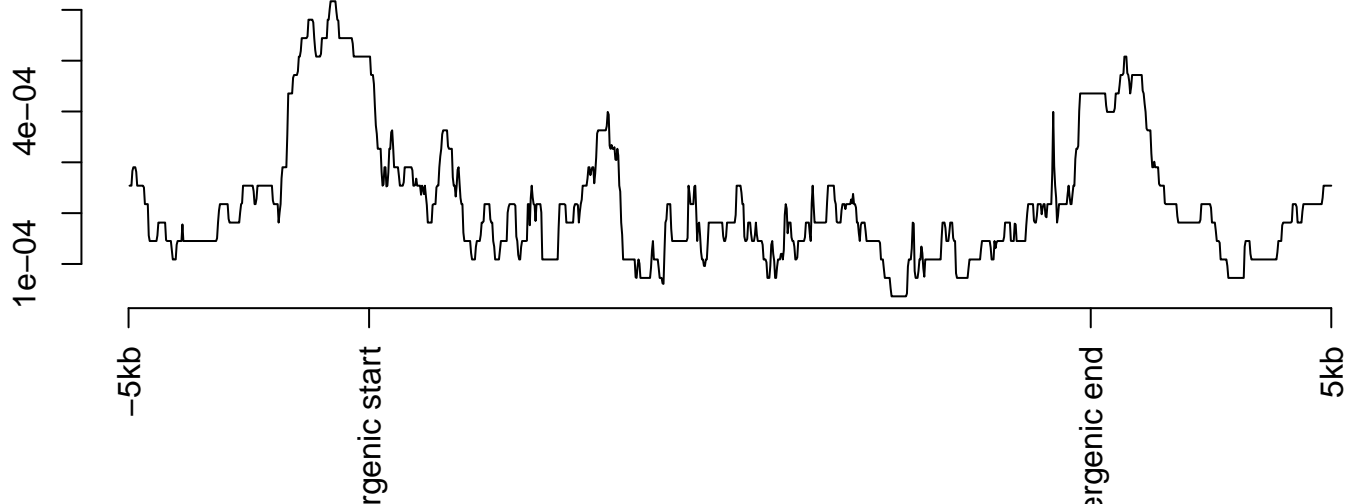
MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc587b

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



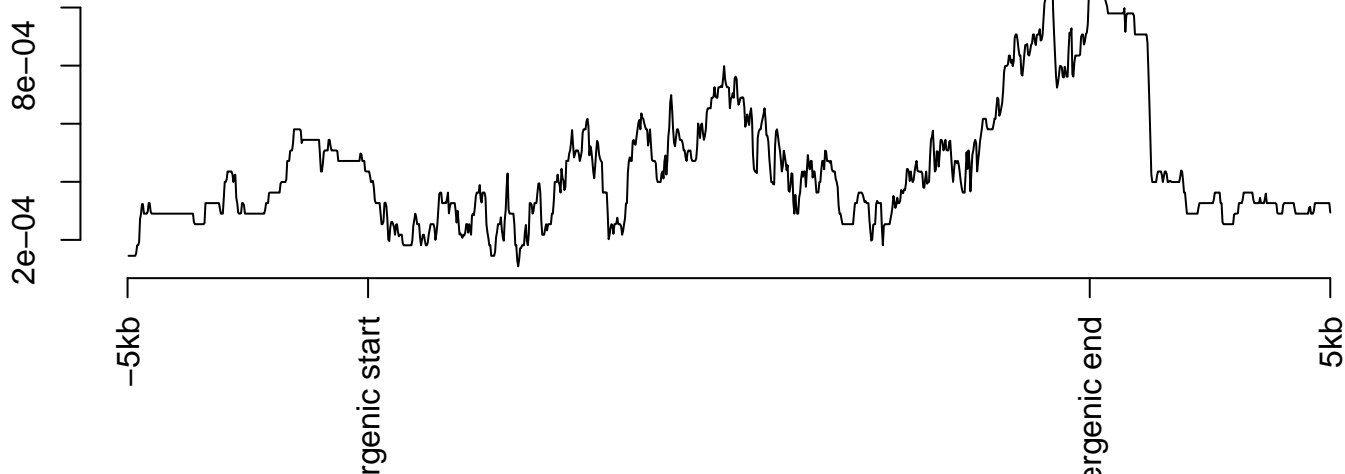
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.648 scCLTdegenNuc588b



Average coverage per intergenic region

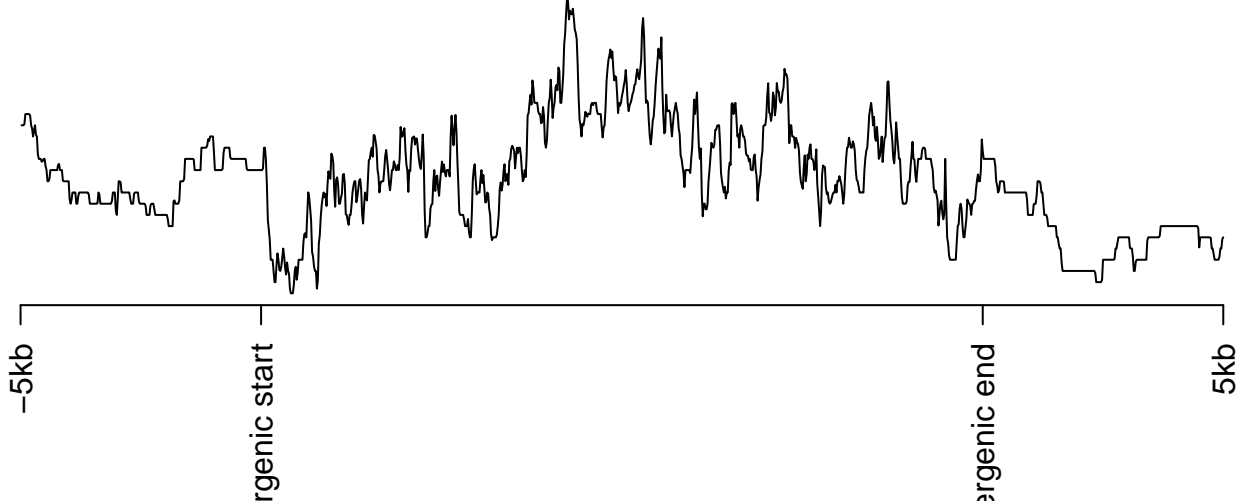
MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc589b



Average coverage per intergenic region

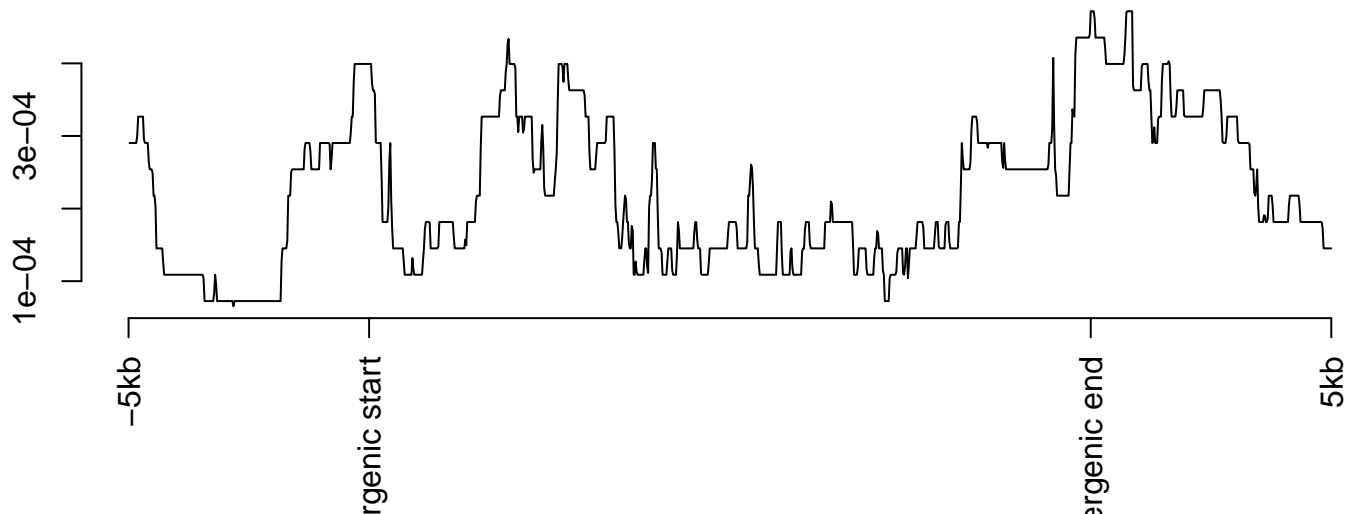
MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc590b

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



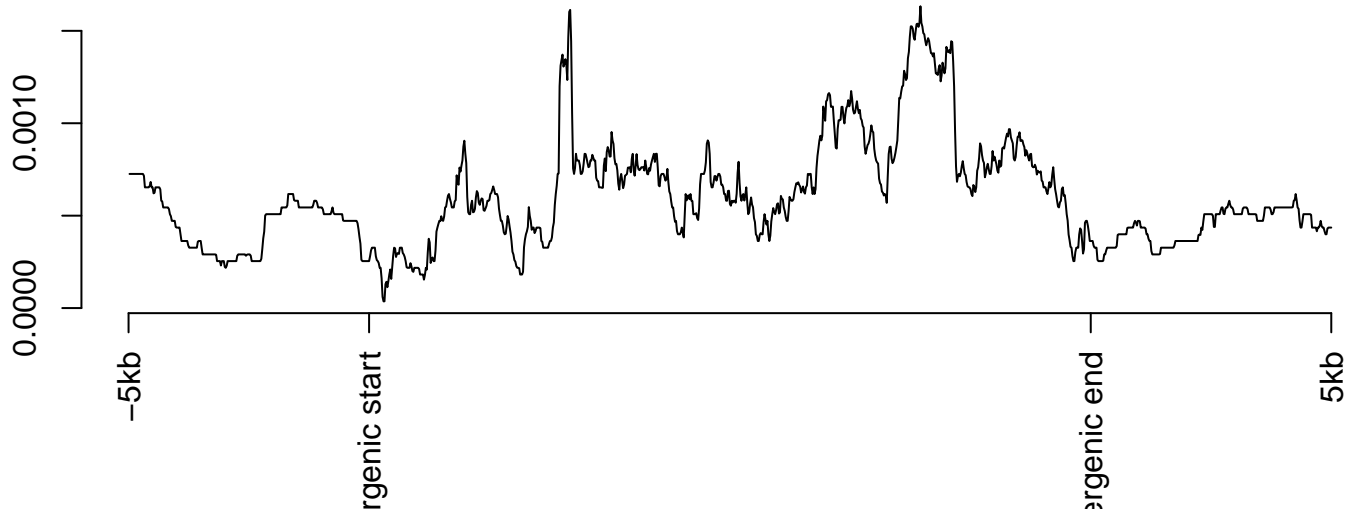
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc591b



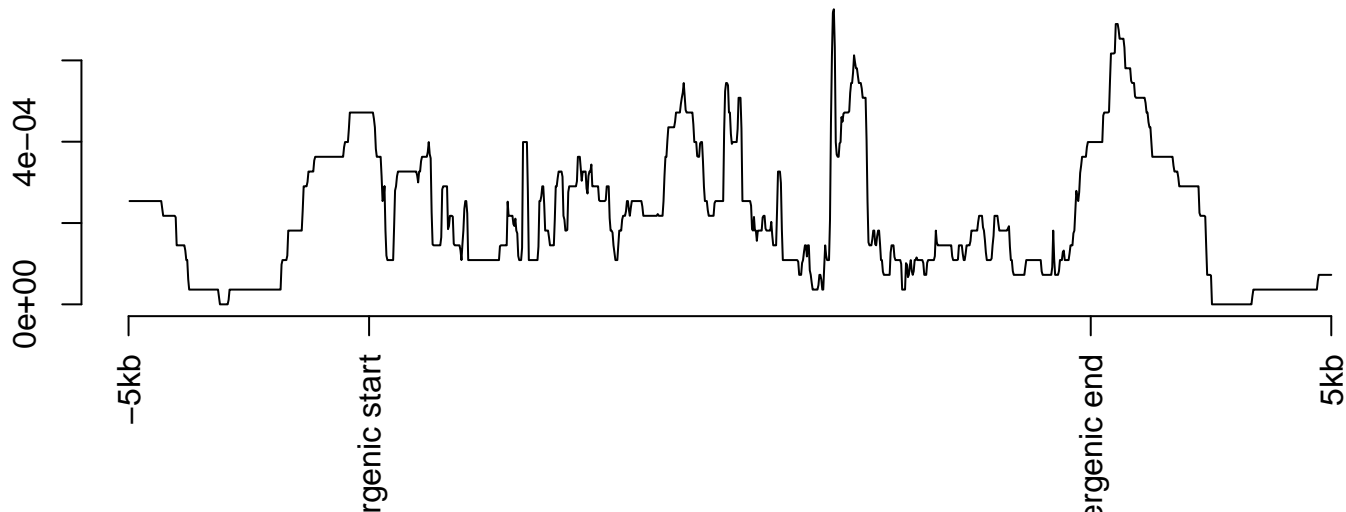
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc592b



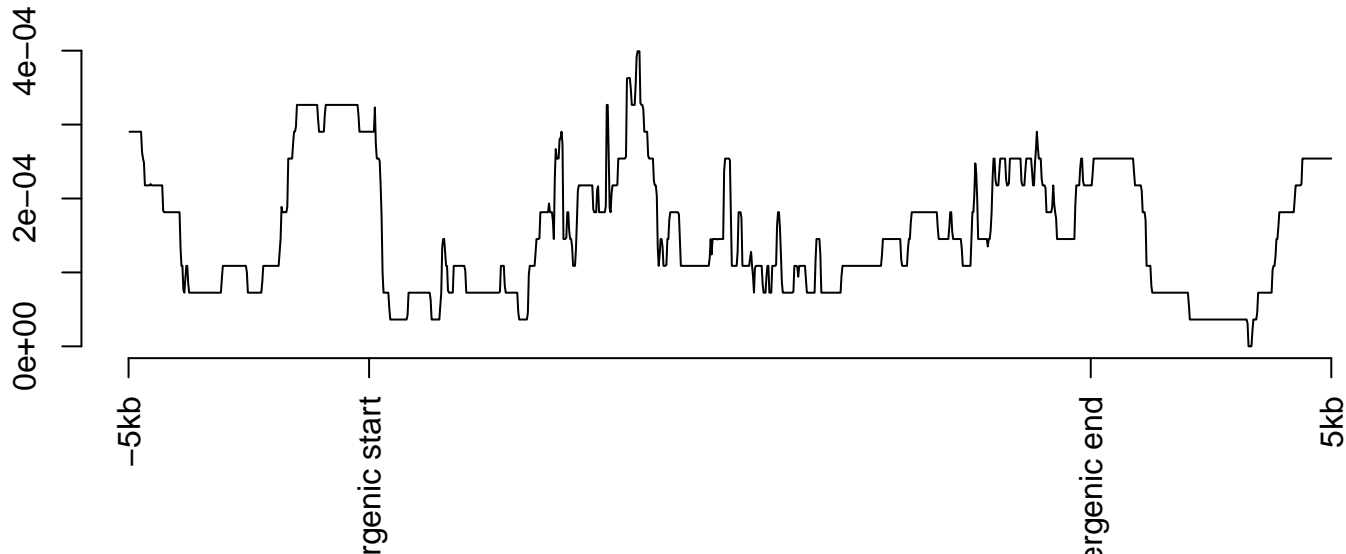
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc593b



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc594b



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc595b

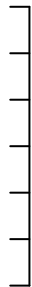
0.0004 0.0010 0.0016

-5kb

intergenic start

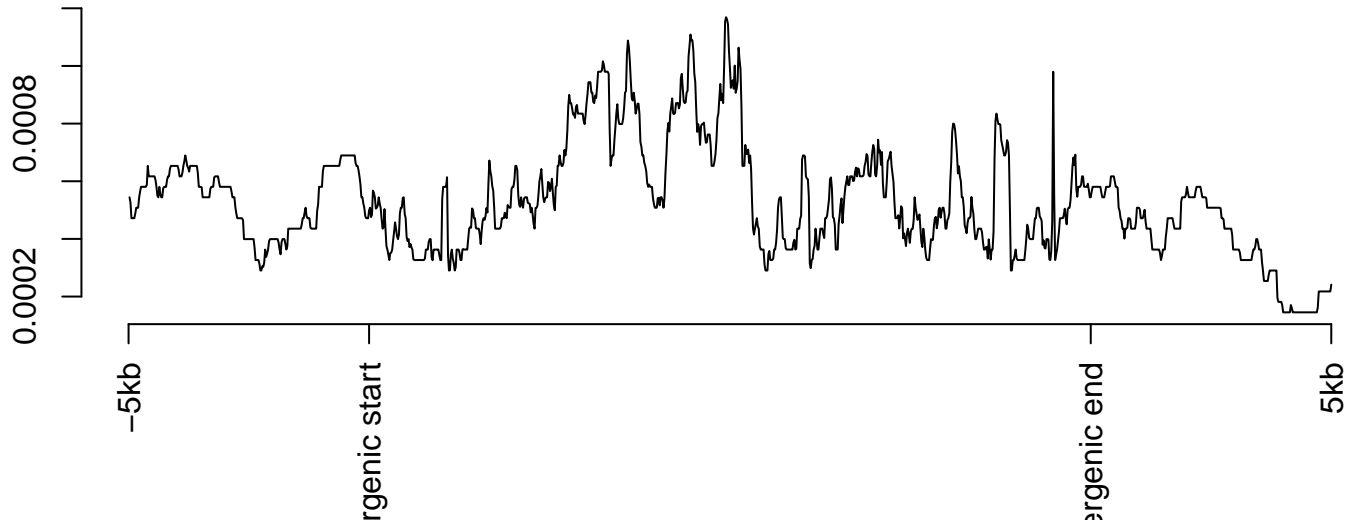
intergenic end

5kb



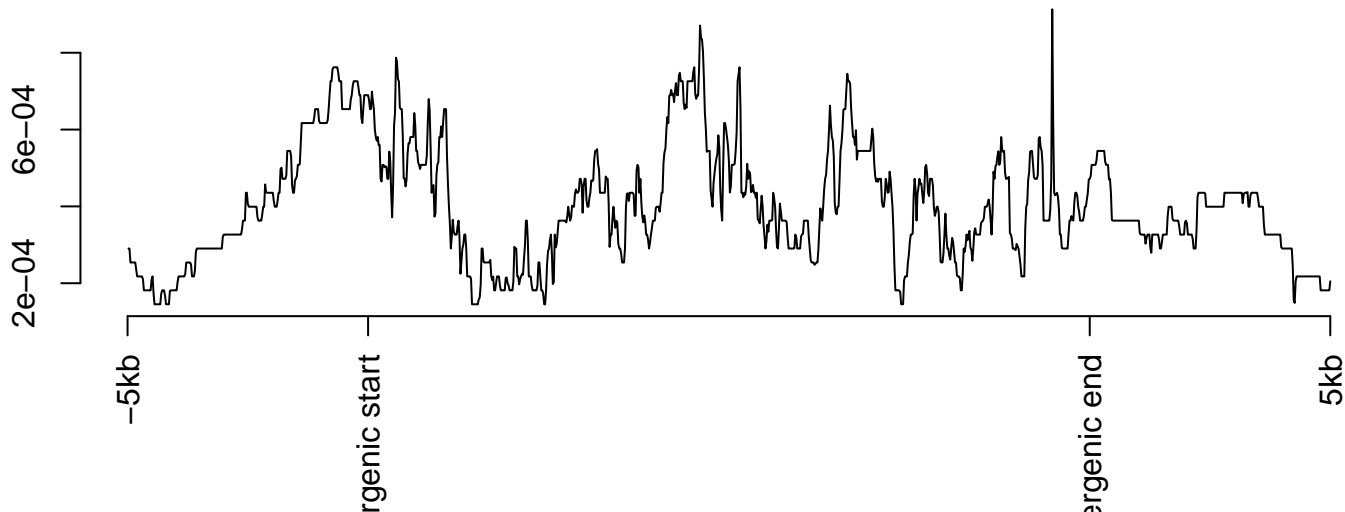
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc596b



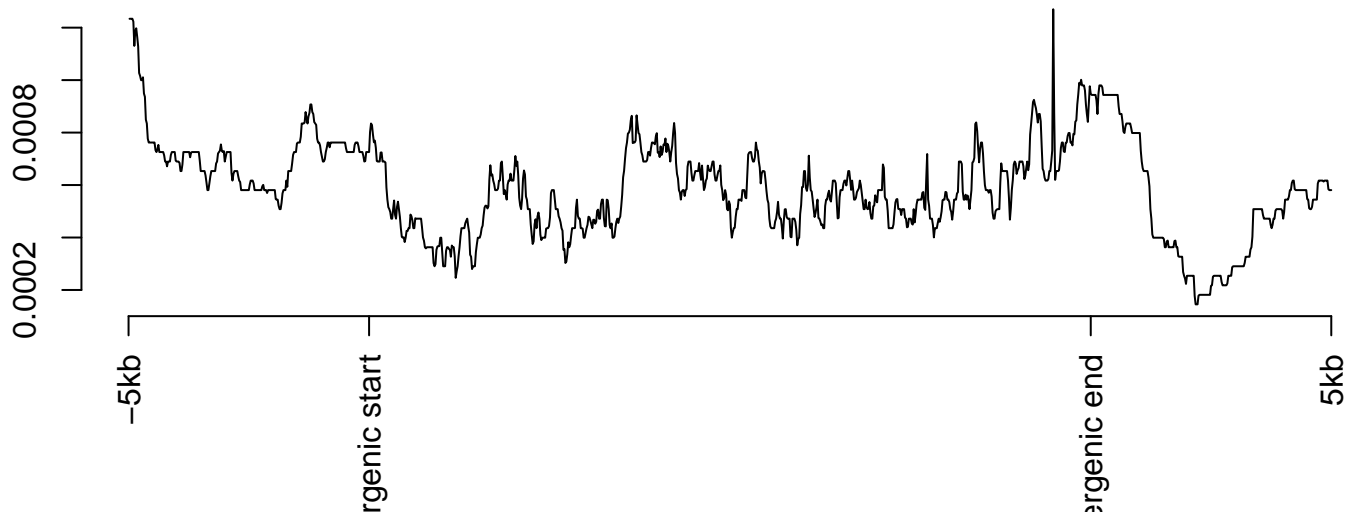
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc597b



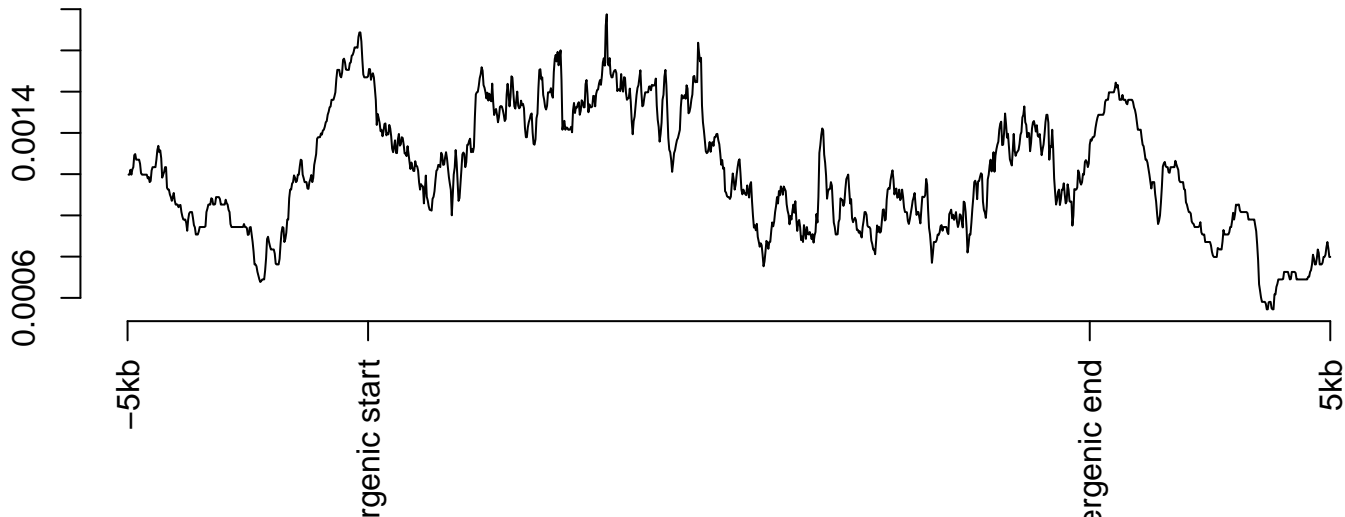
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc598b



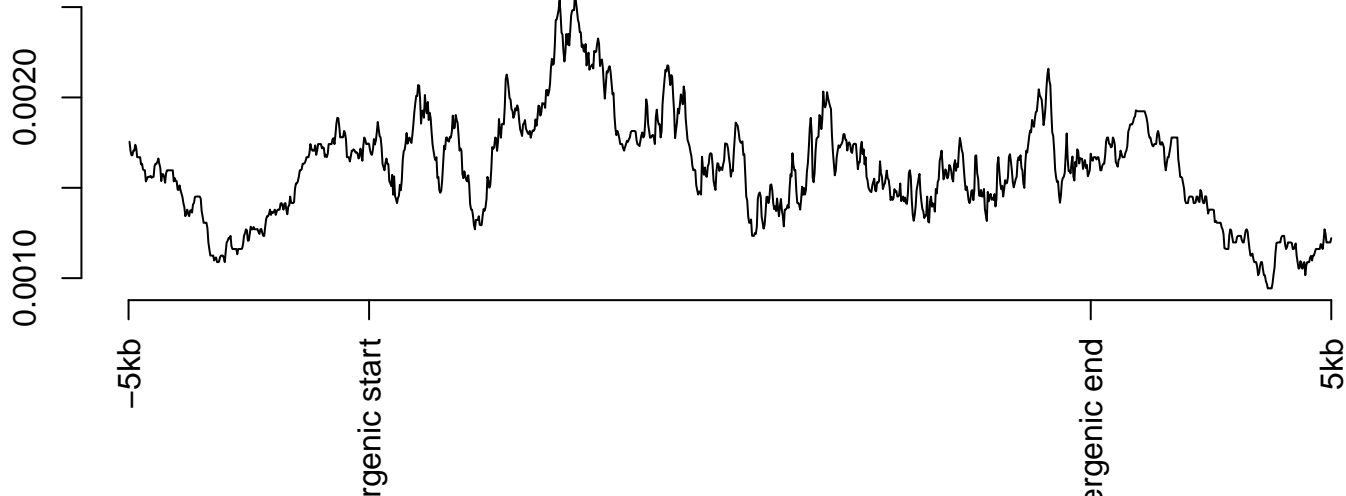
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc599b



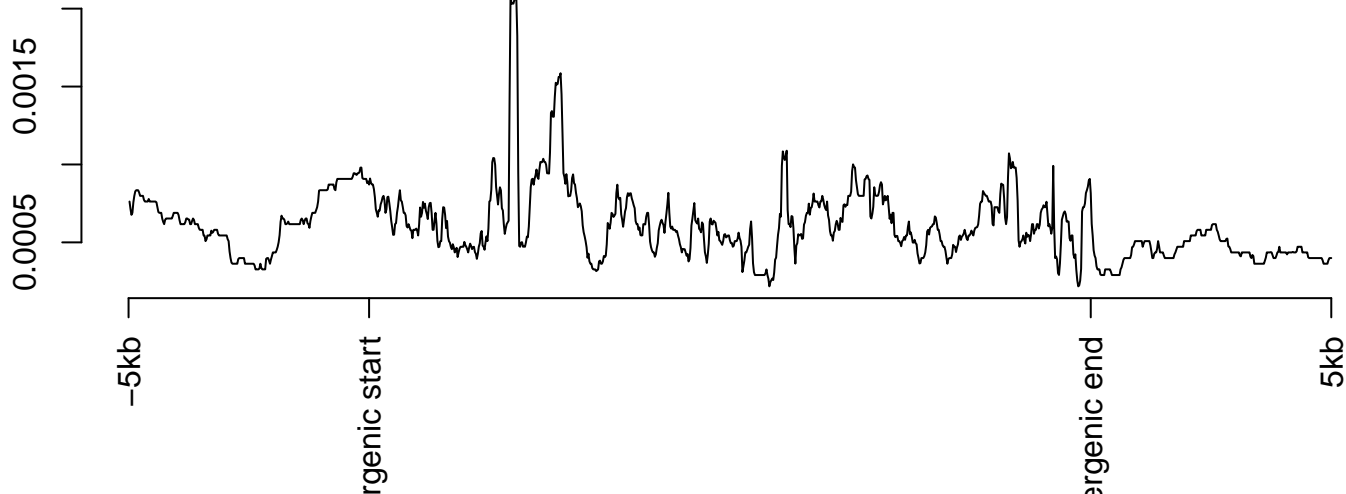
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.649 scCLTdegenNuc600b



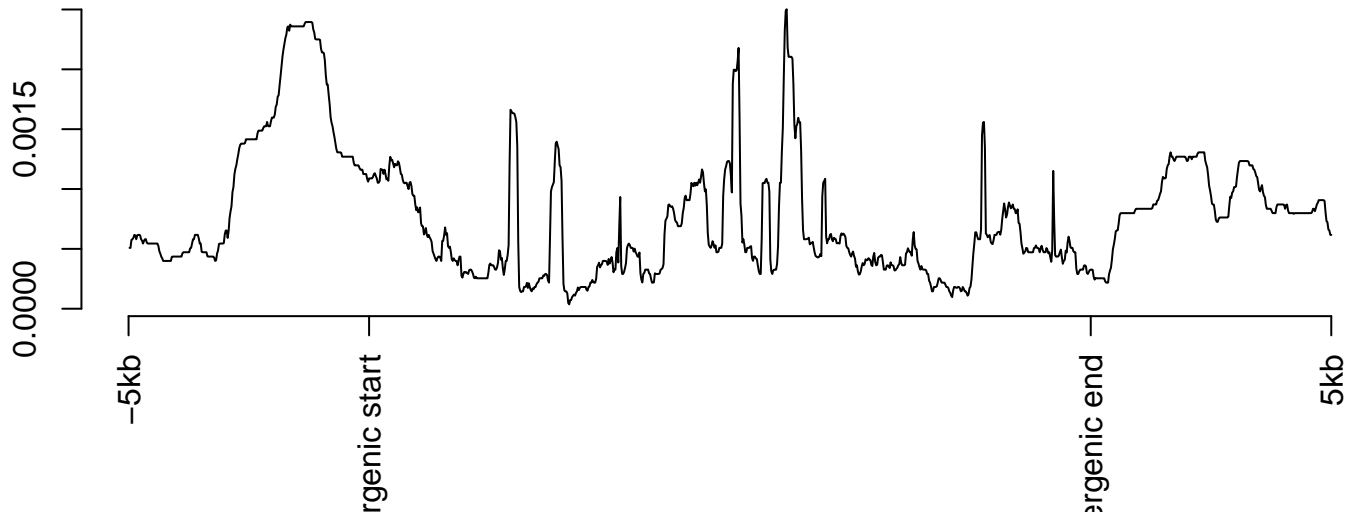
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc779



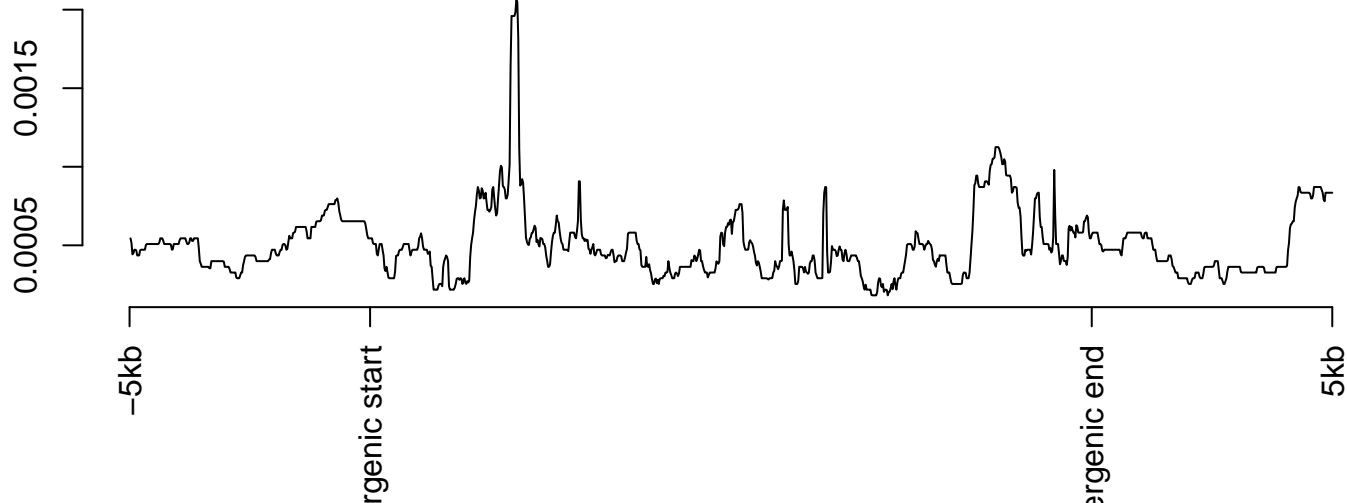
MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc780

Average coverage per intergenic region



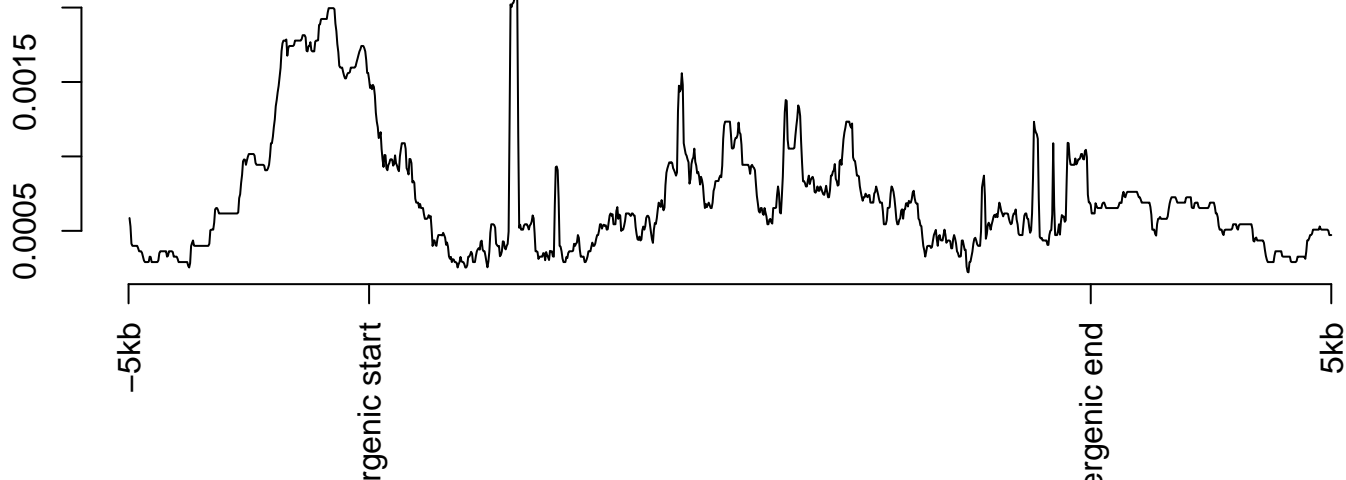
MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc781

Average coverage per intergenic region



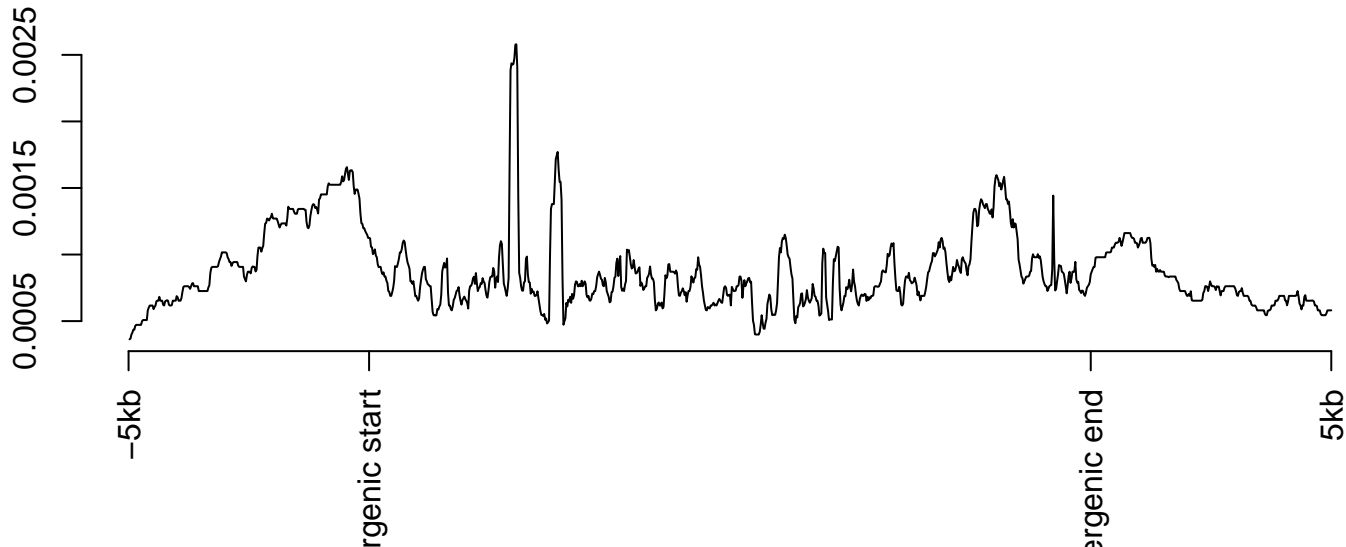
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc782



Average coverage per intergenic region

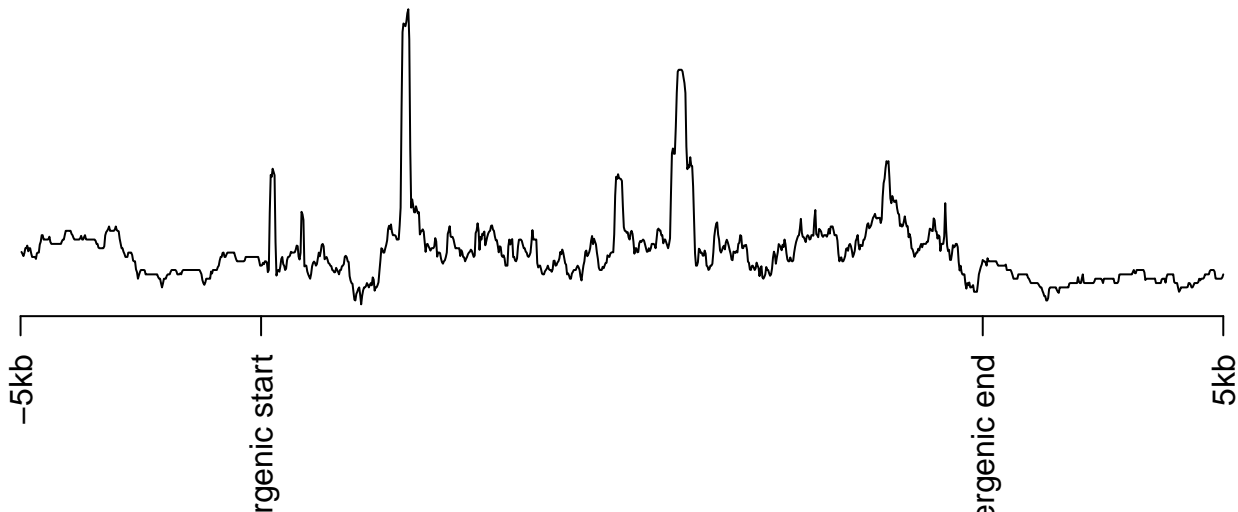
MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc783



MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc784

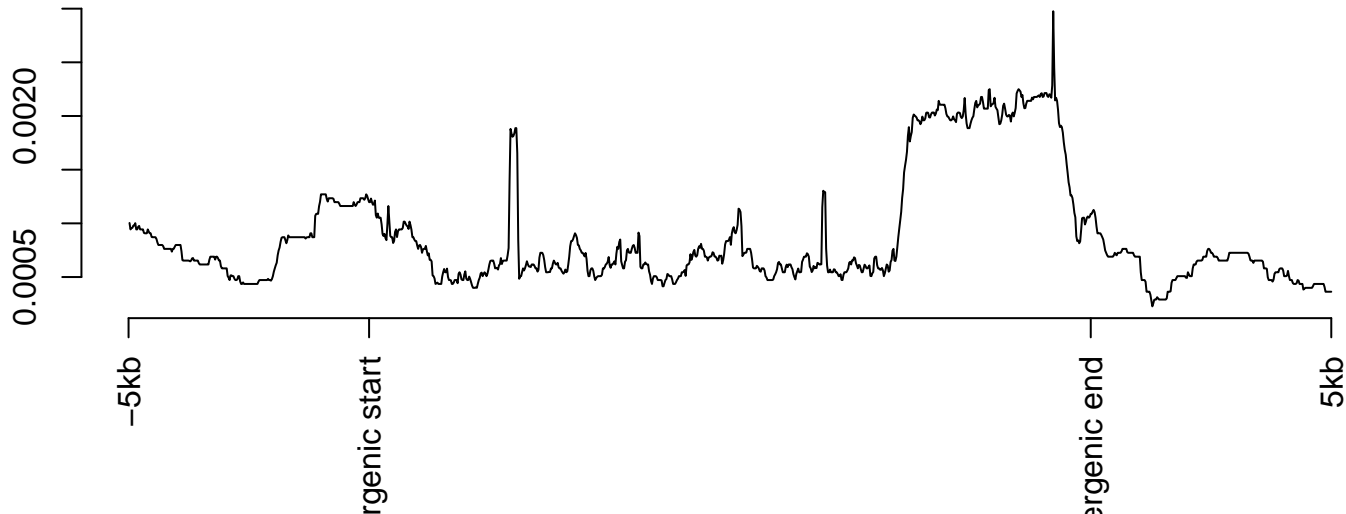
Average coverage per intergenic region

0.0020
0.0005



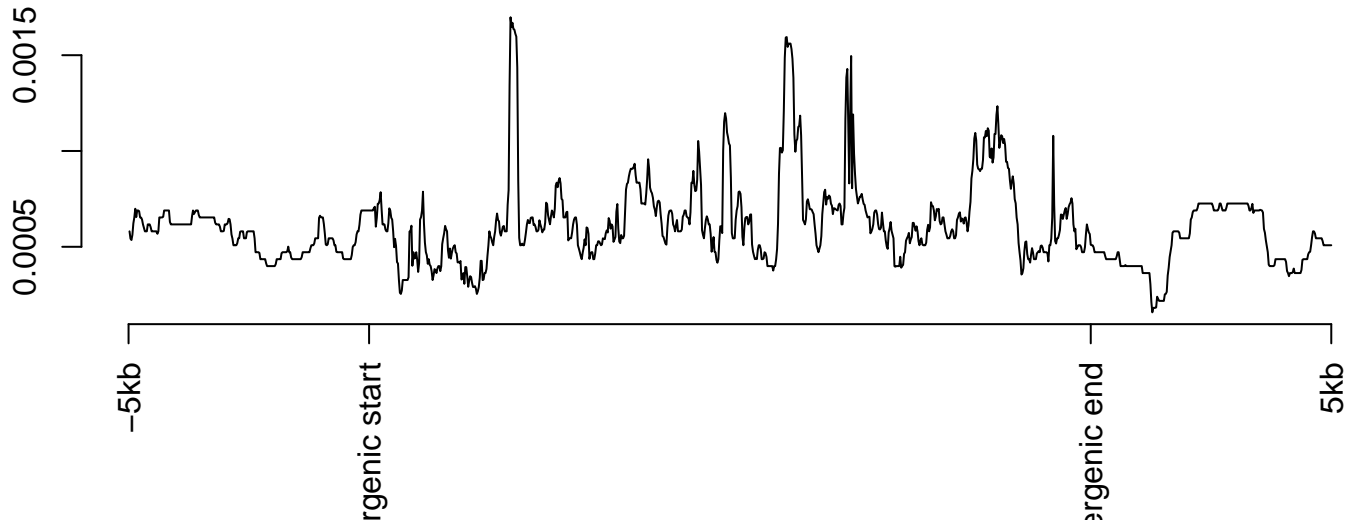
MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc785

Average coverage per intergenic region



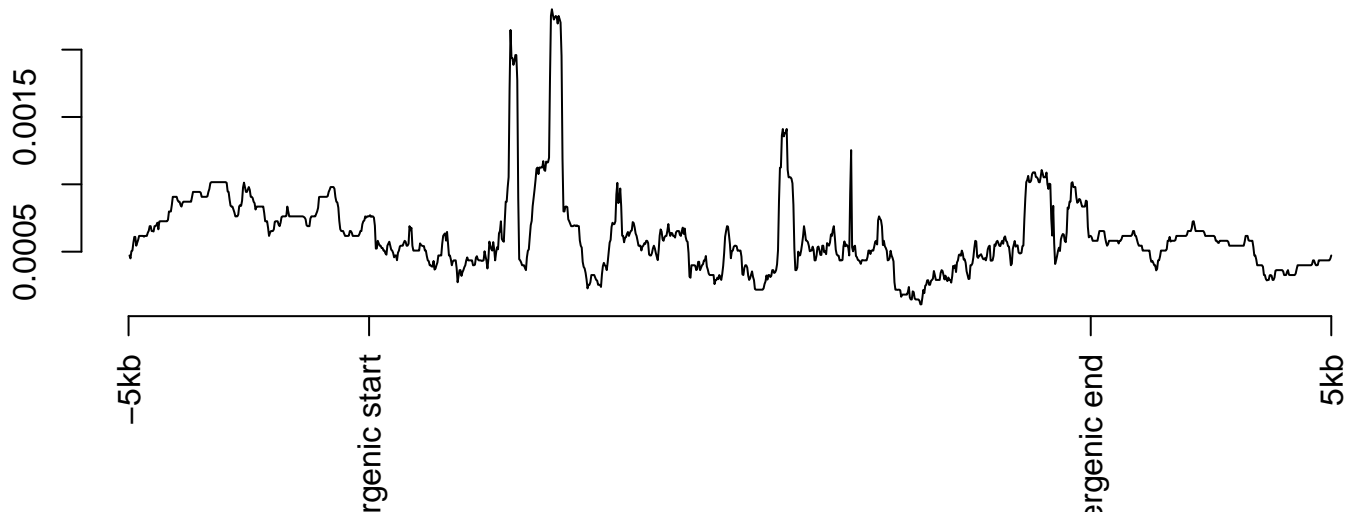
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc786



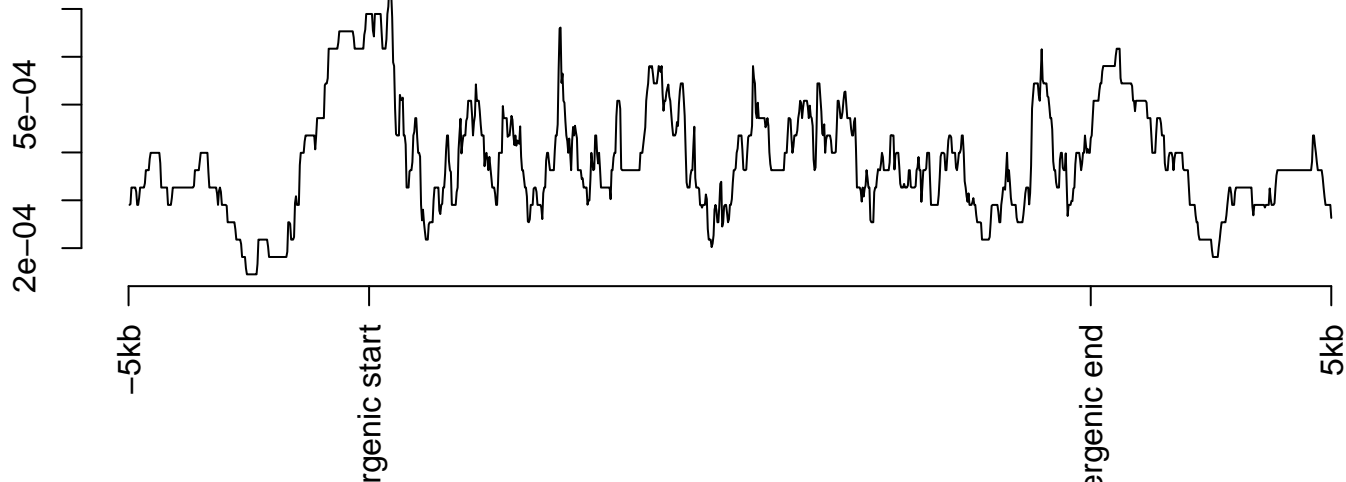
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.666 scCLTdegenNuc787



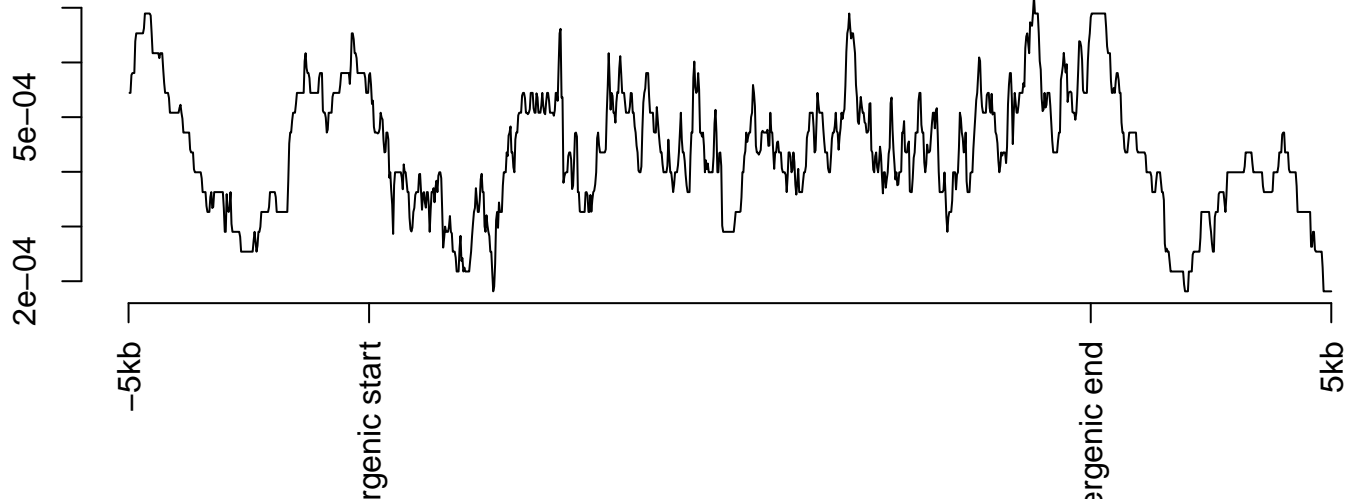
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc788



Average coverage per intergenic region

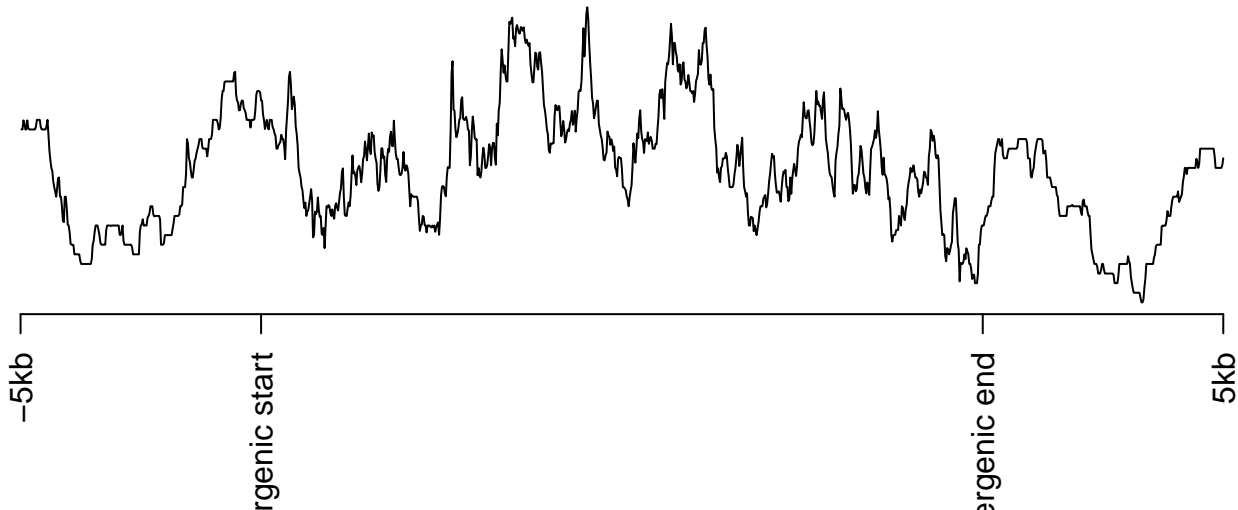
MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc789



Average coverage per intergenic region

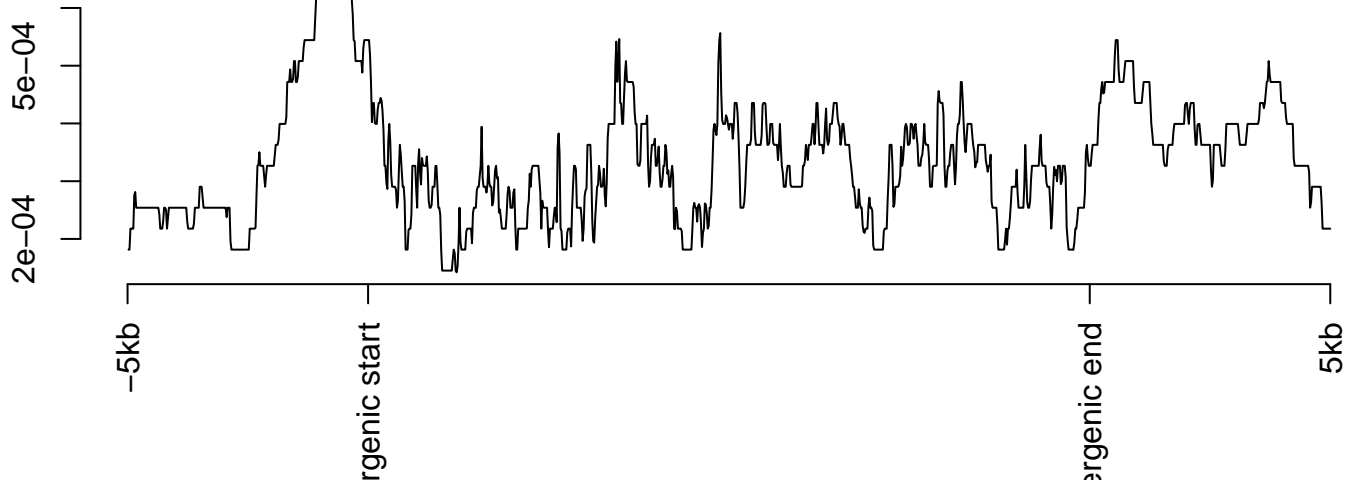
MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc790

0.0004 0.0010



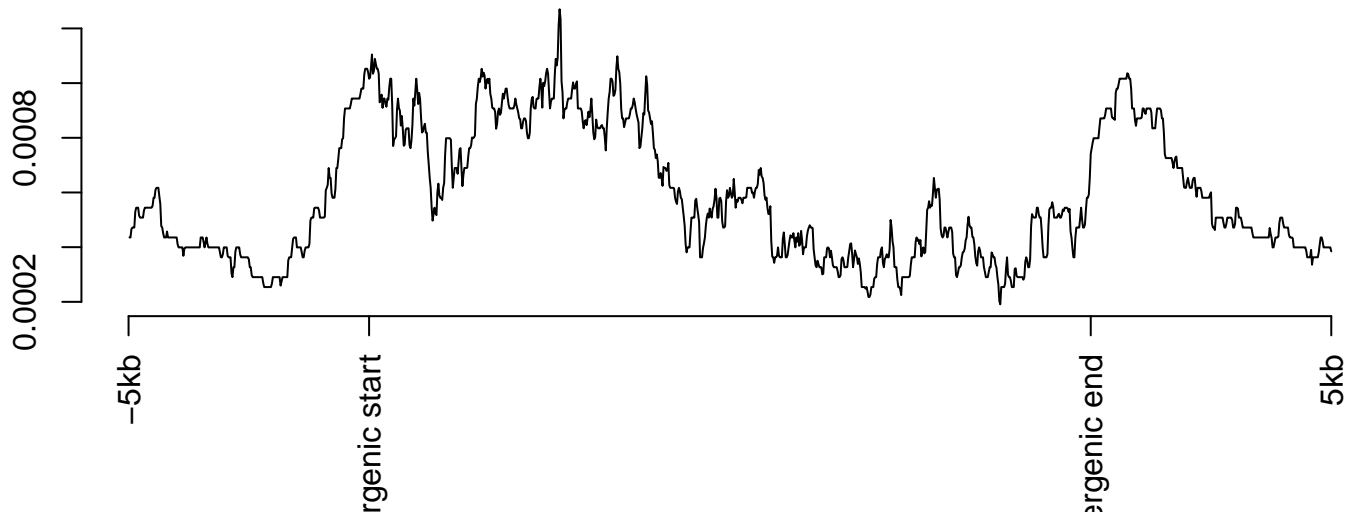
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc791



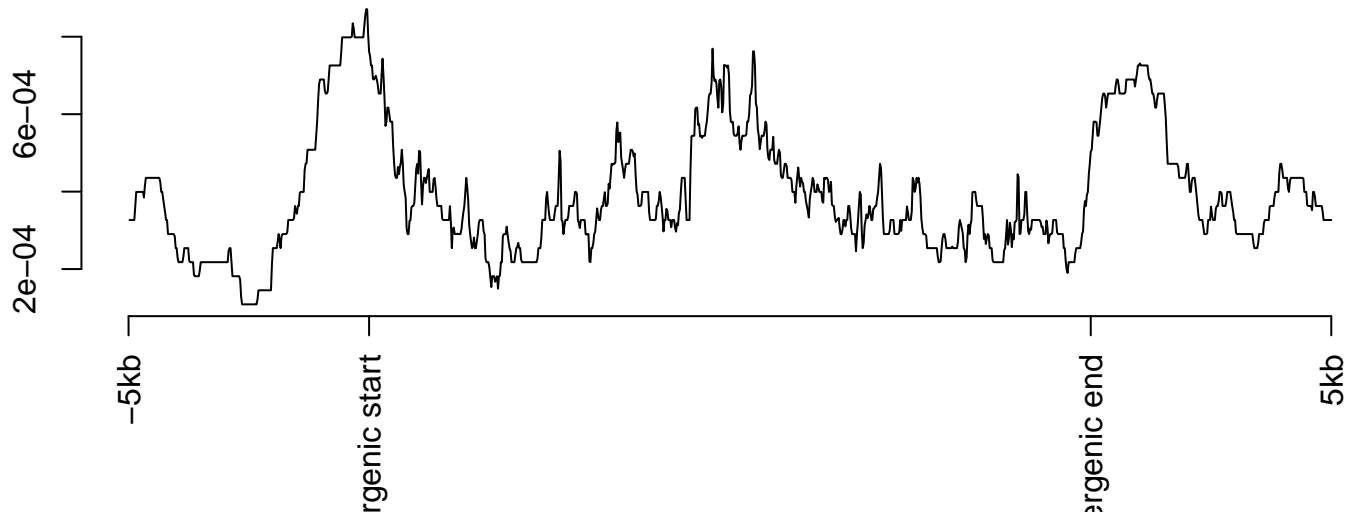
MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc792

Average coverage per intergenic region



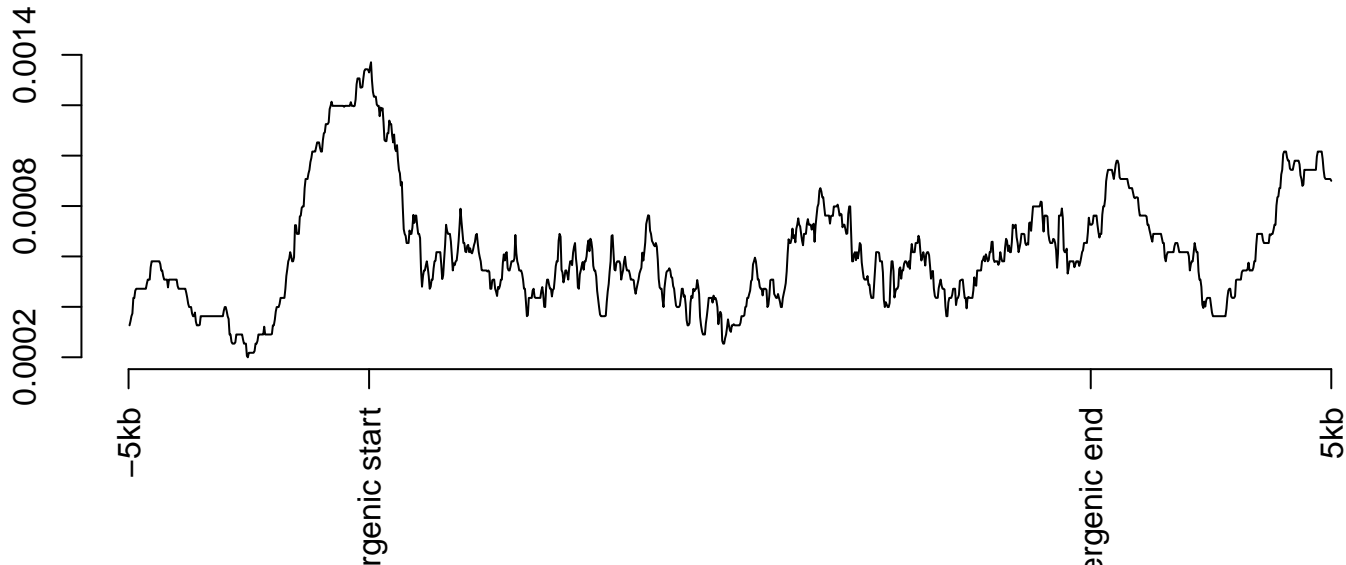
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc793



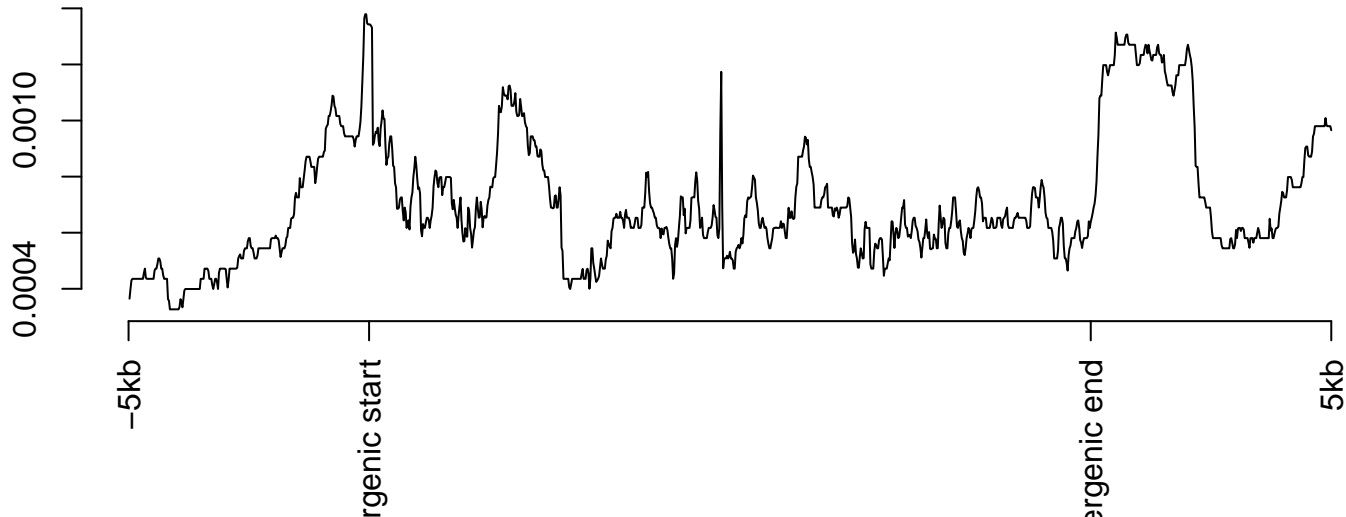
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc794



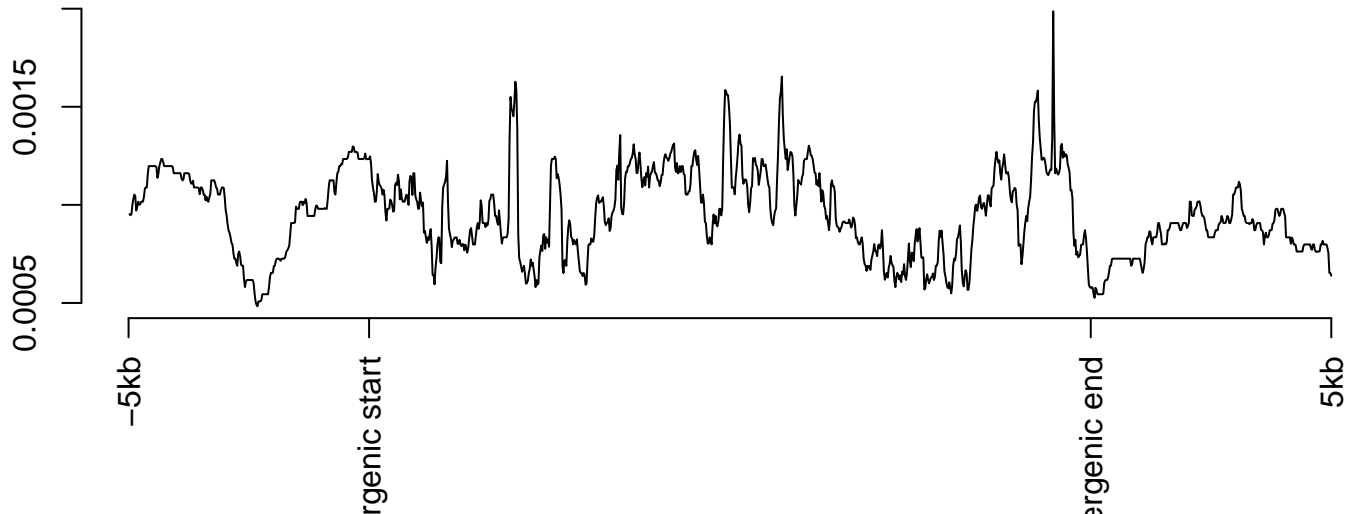
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.668 scCLTdegenNuc795



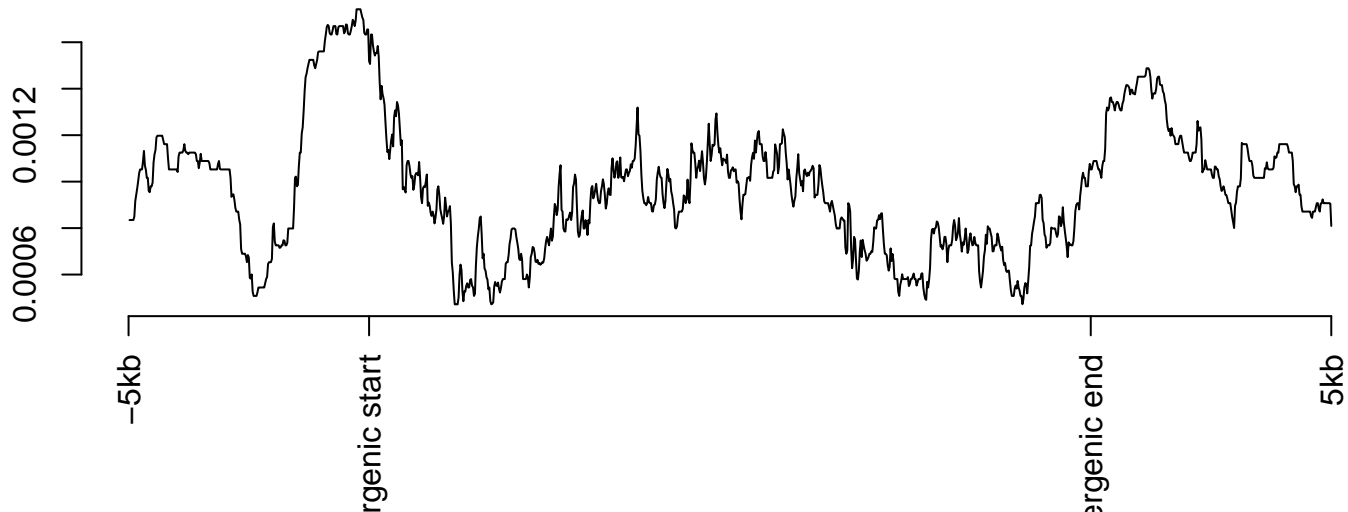
MouseNeuronCulture PositiveSingle E.669 scCLTdegenNuc808

Average coverage per intergenic region



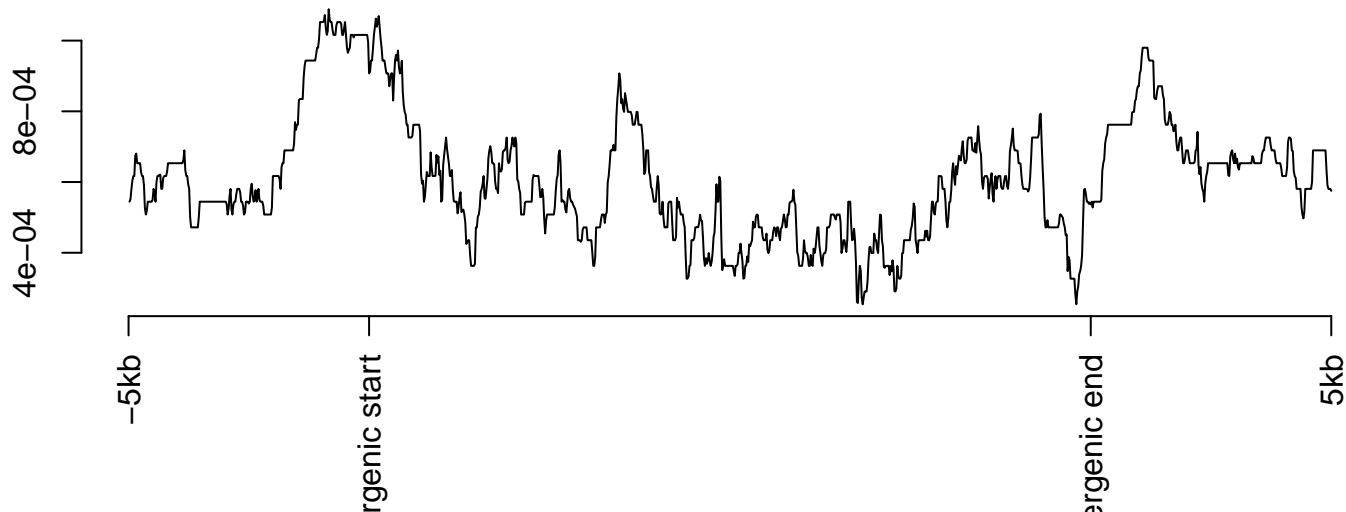
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc797



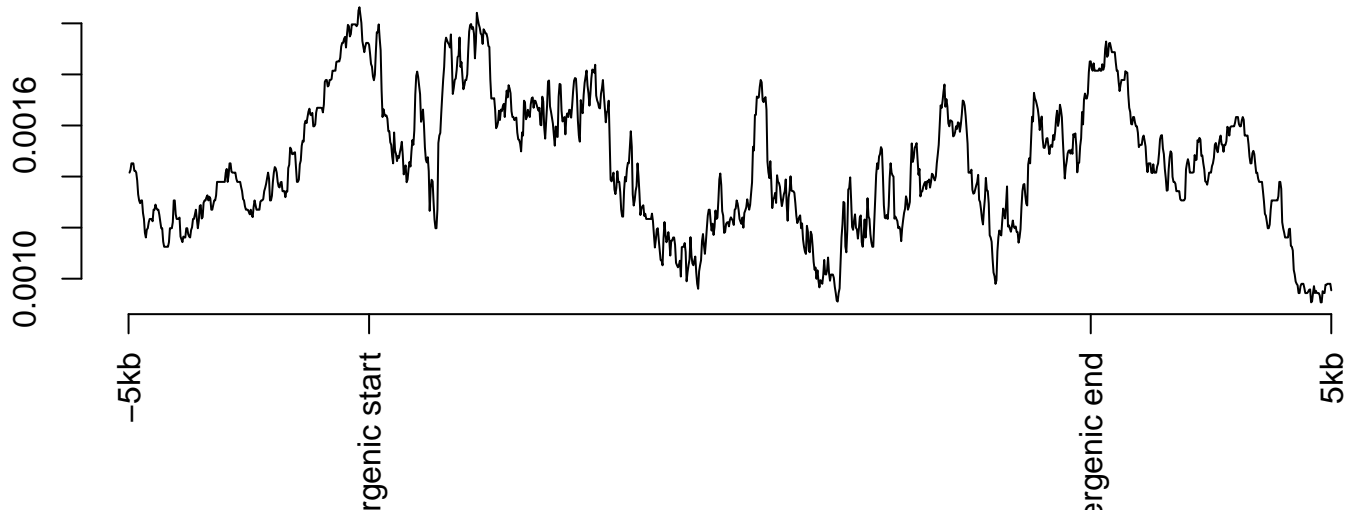
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc798



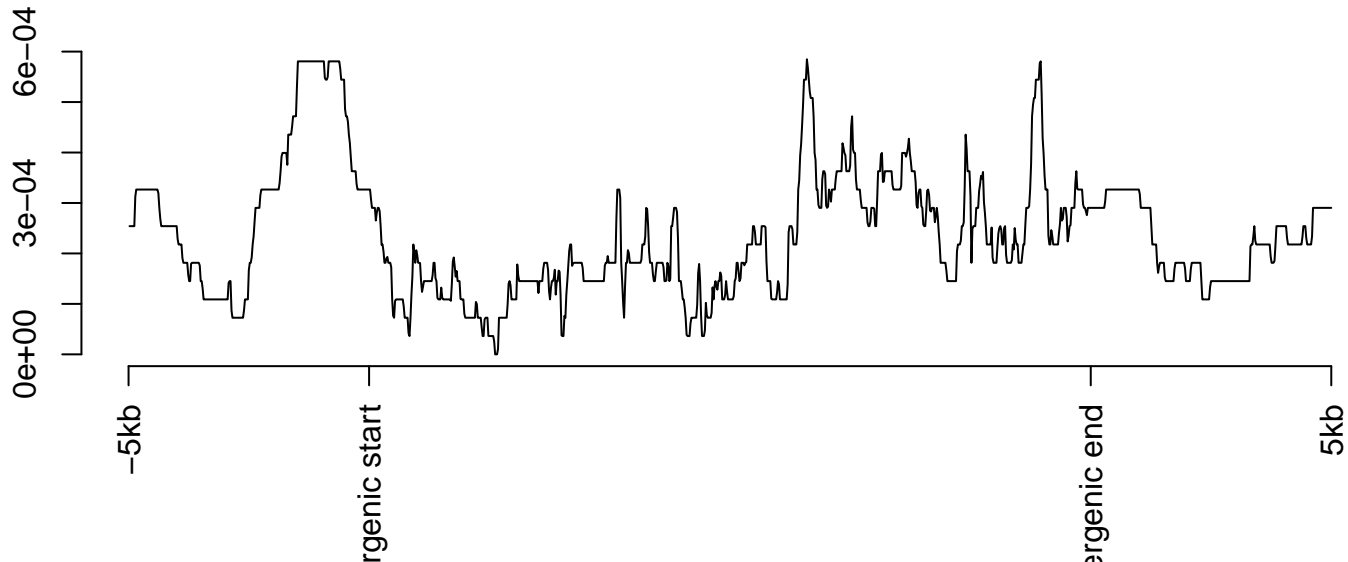
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.670 scCLTdegenNuc799



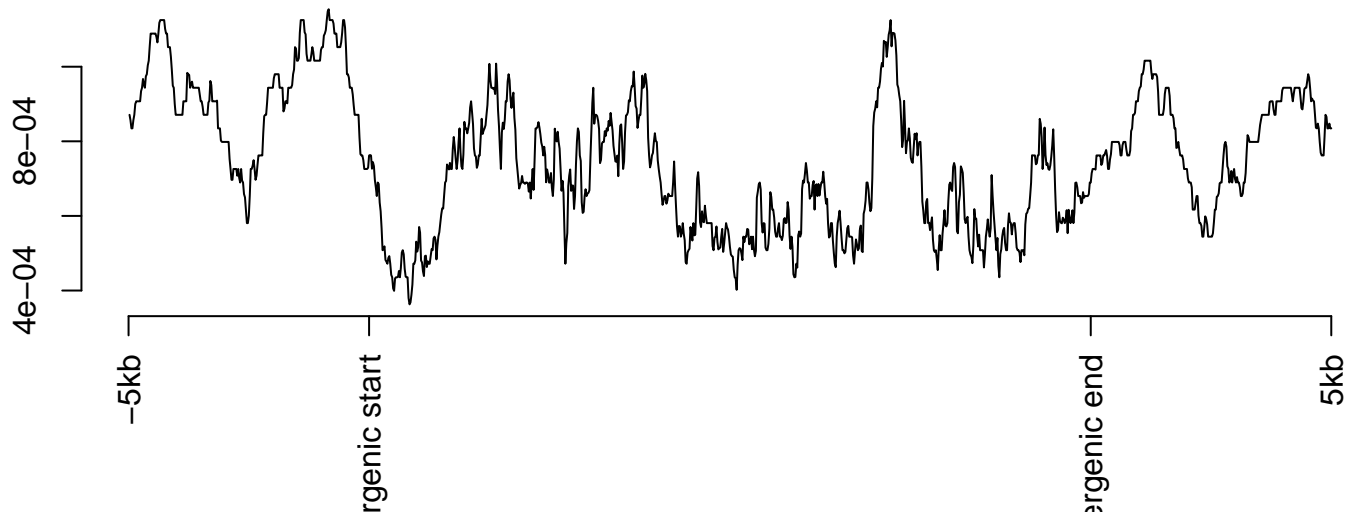
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.671 scCLTdegenNuc803



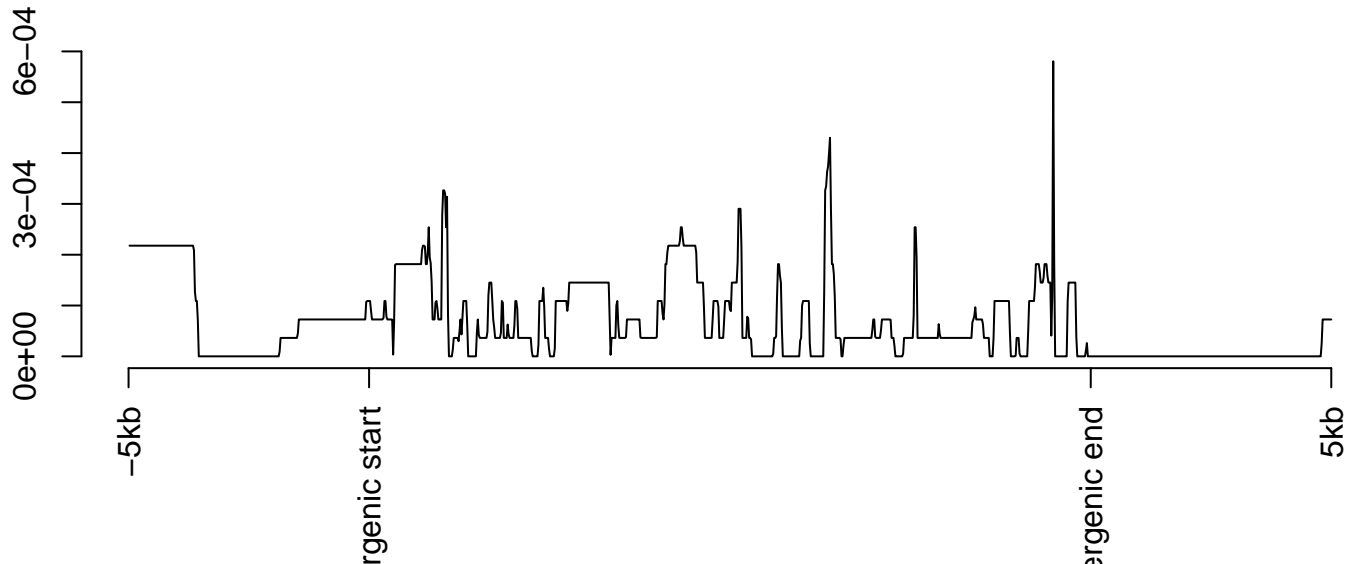
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.671 scCLTdegenNuc807



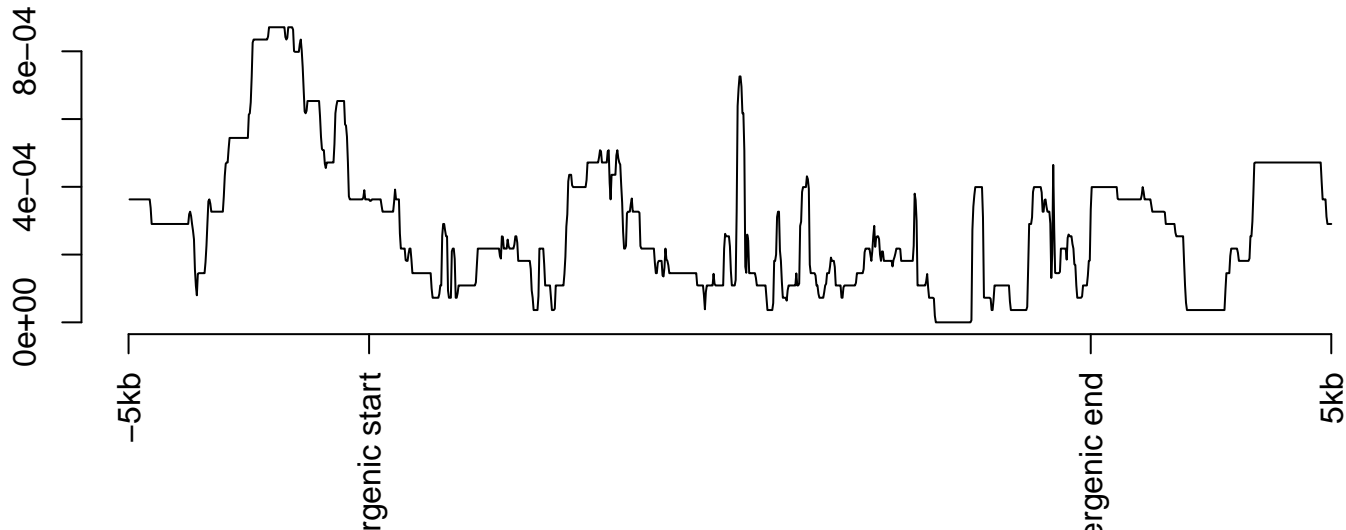
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc965



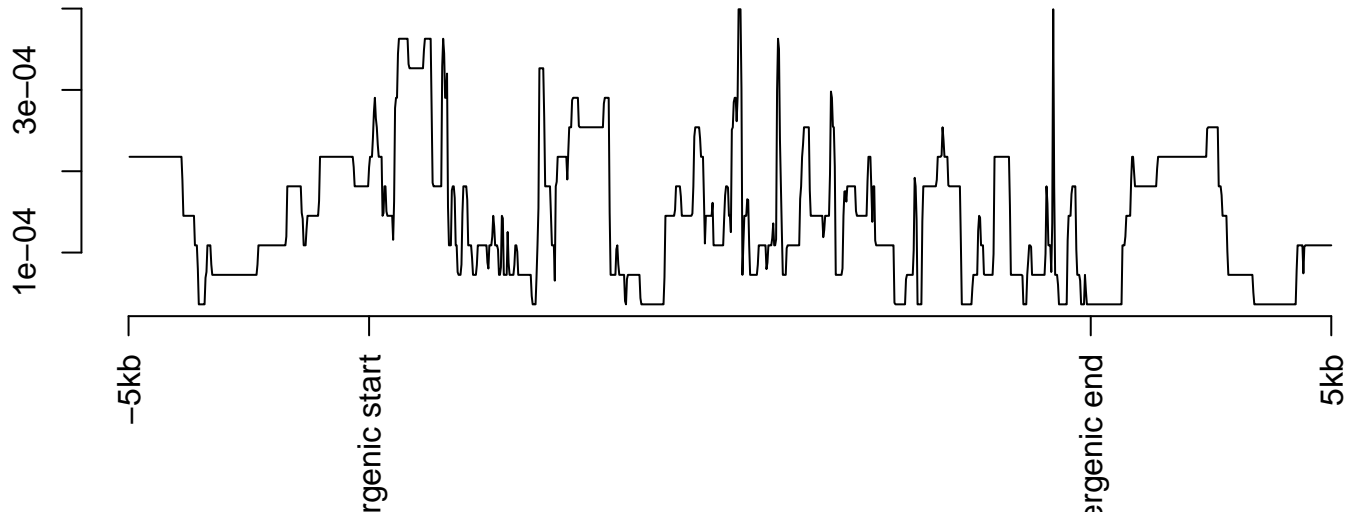
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc966



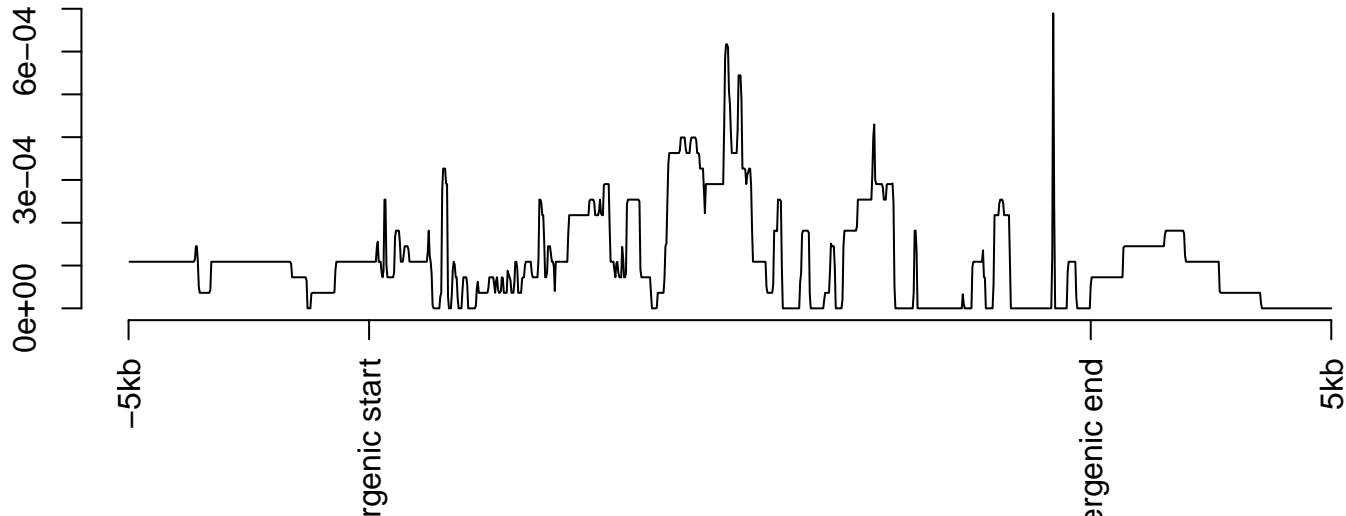
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc967



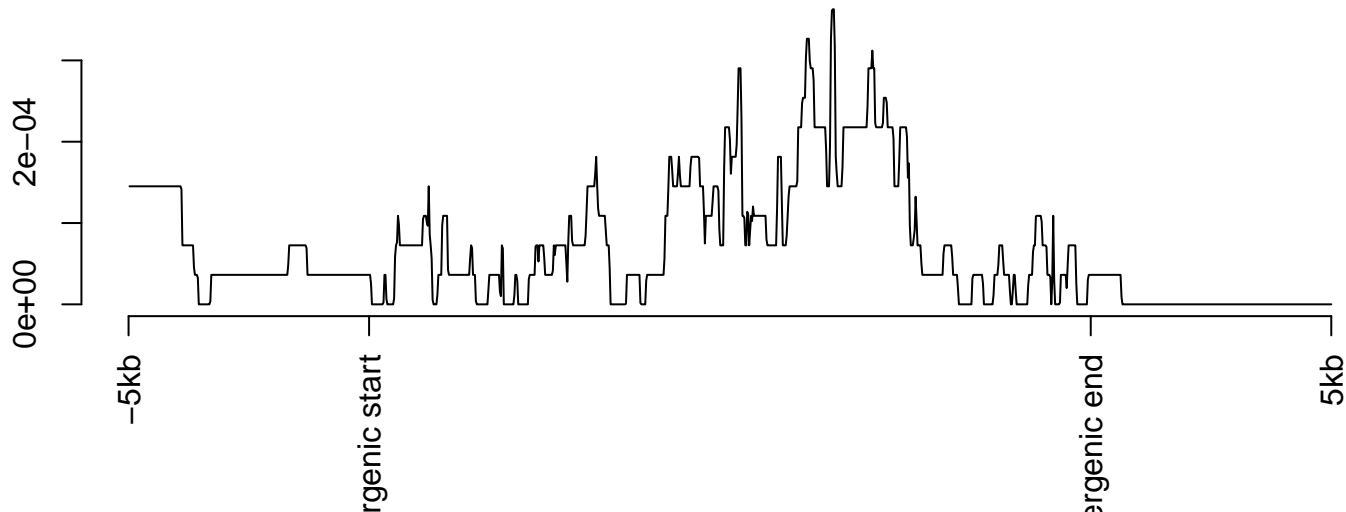
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc968



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc969



Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.681 scCLTdegenNuc970

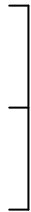
0.0020
0.0010

-5kb

intergenic start

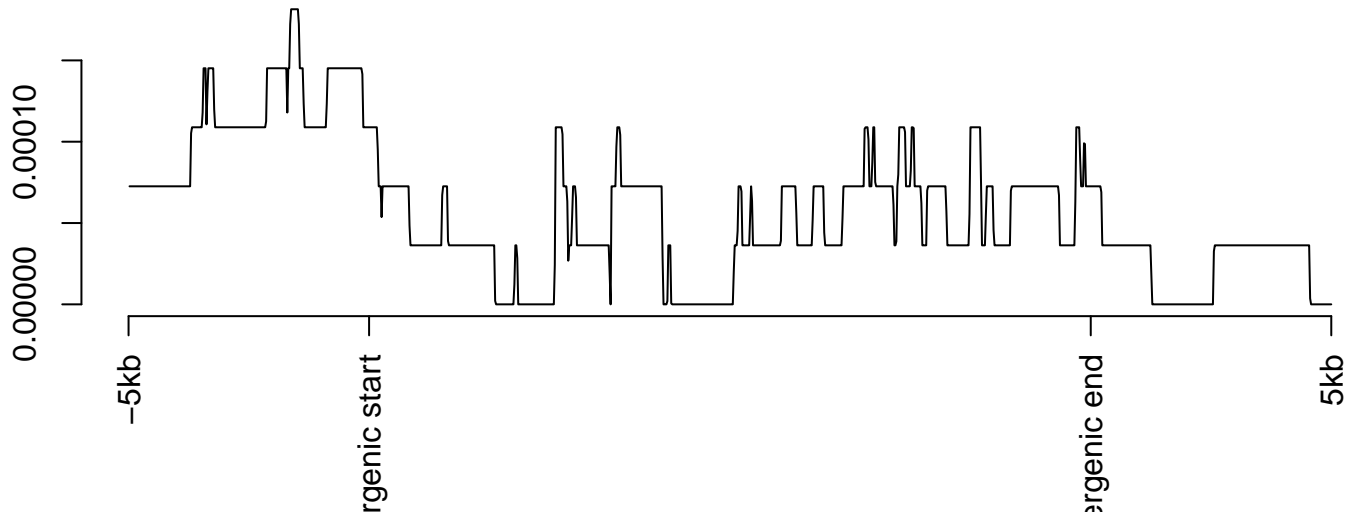
intergenic end

5kb



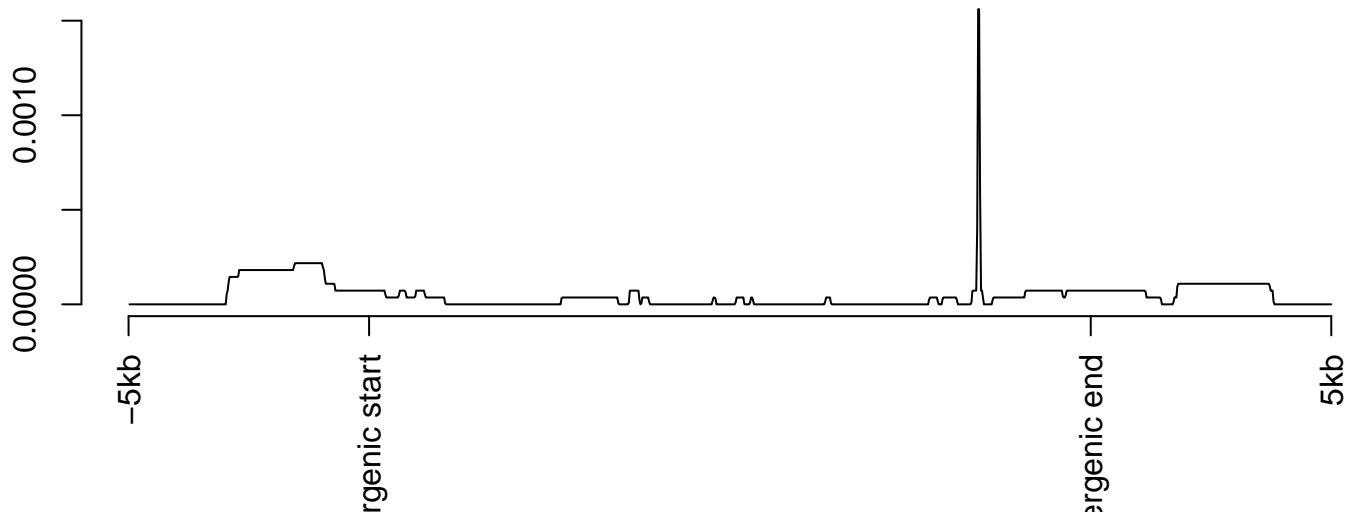
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1063



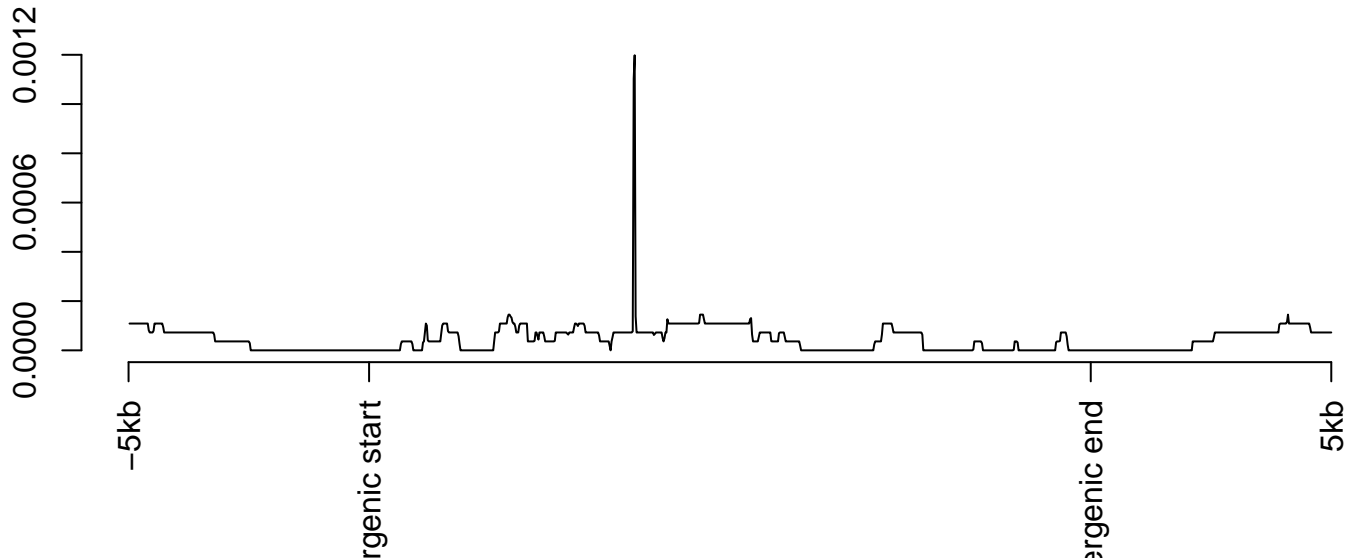
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1064



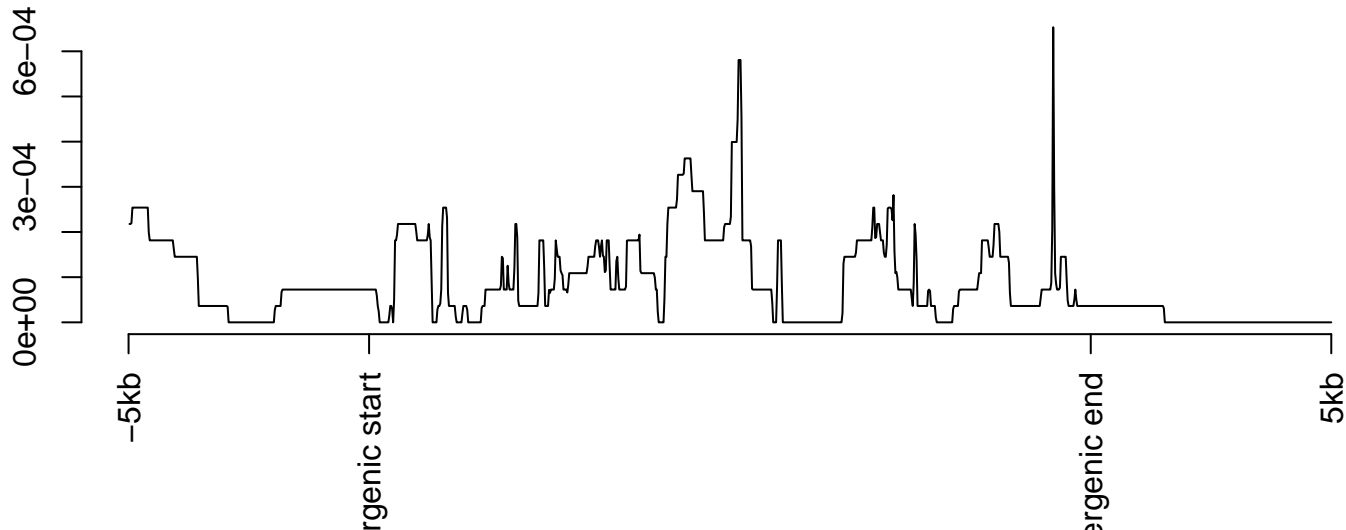
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1065



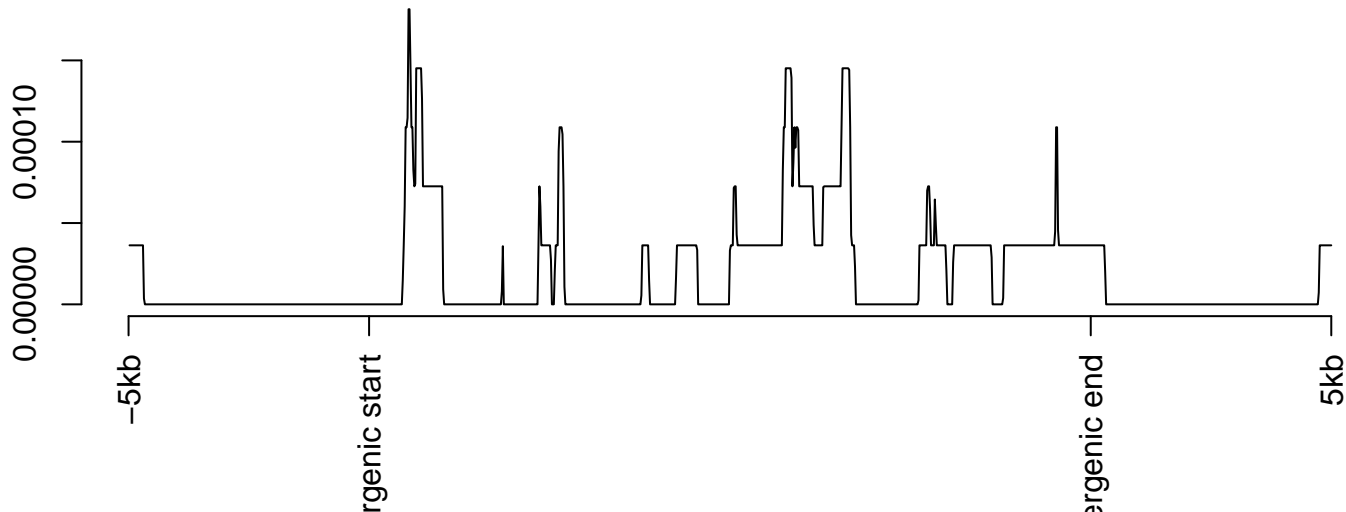
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.692 scCLTdegenNuc1068



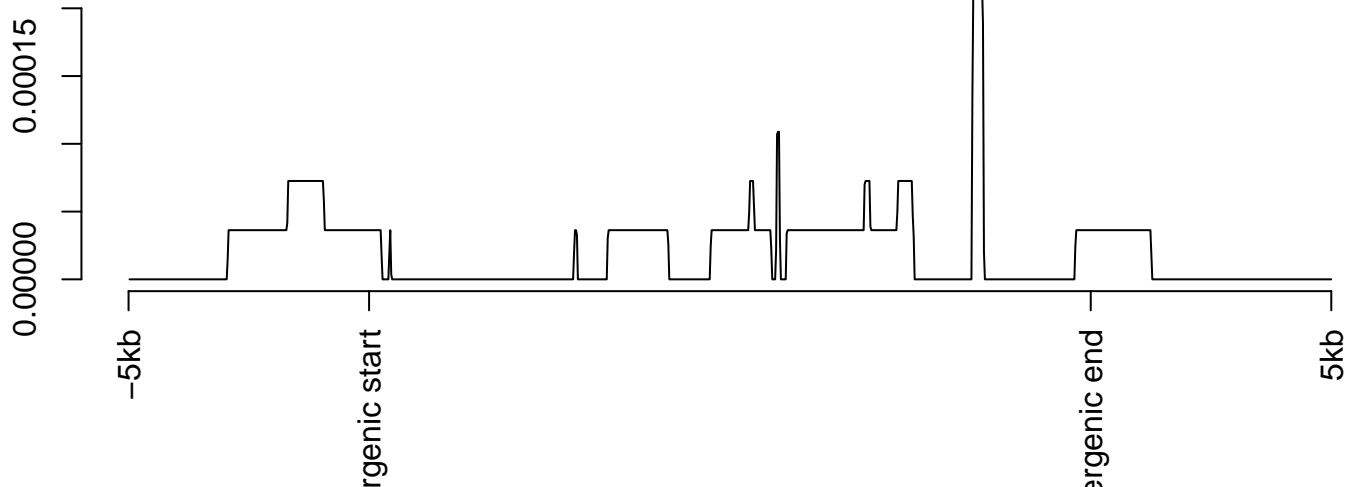
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1066



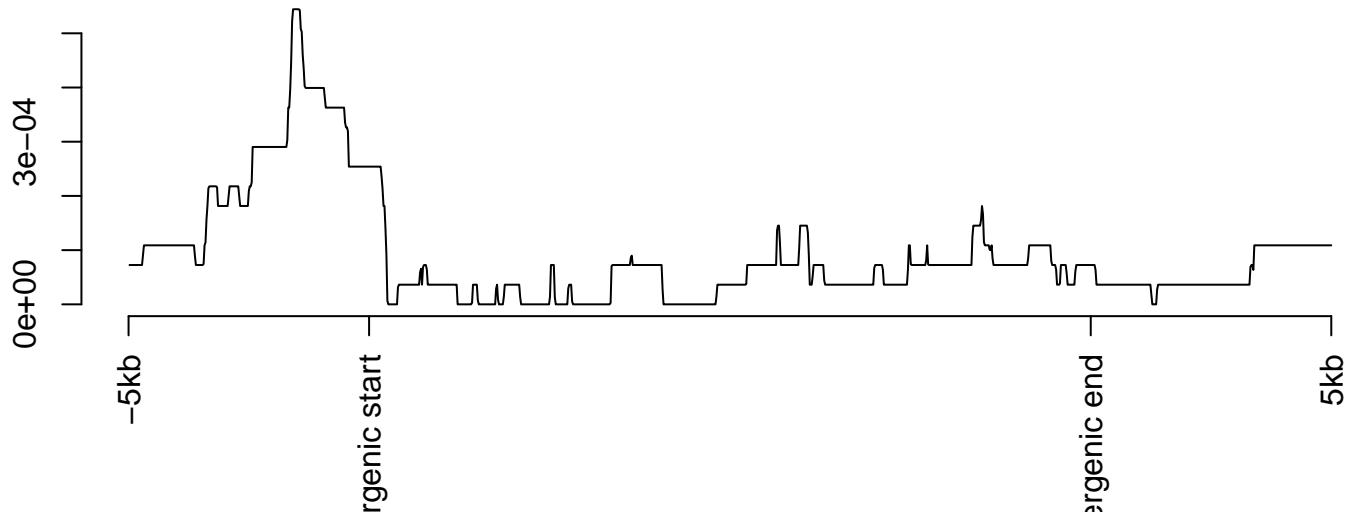
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1067



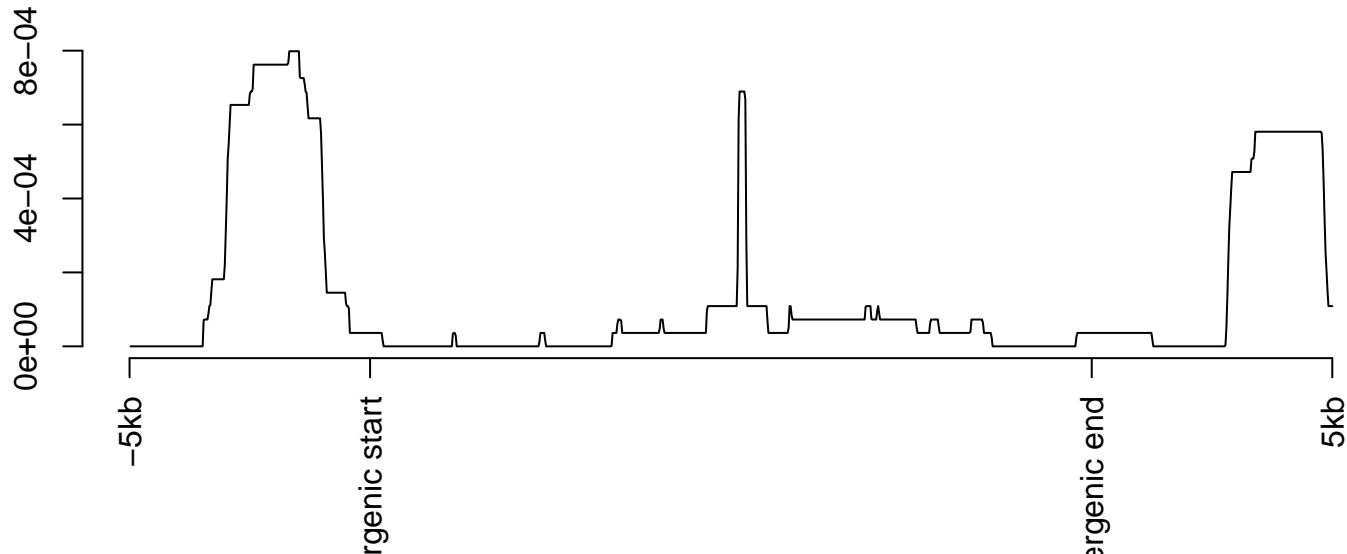
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1095



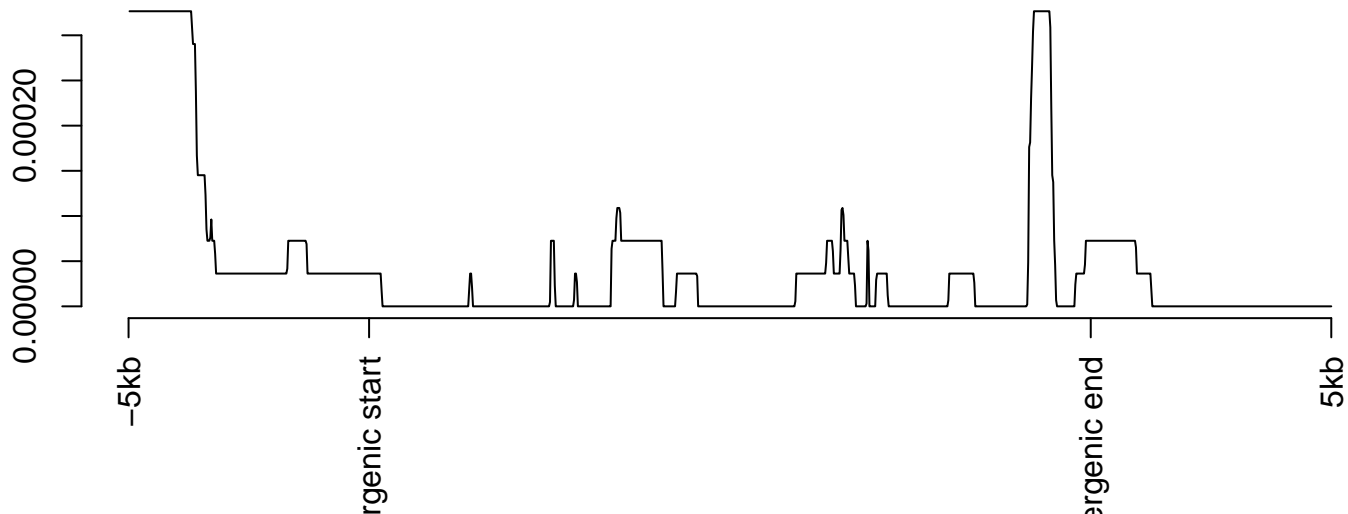
Average coverage per intergenic region

MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1096



Average coverage per intergenic region

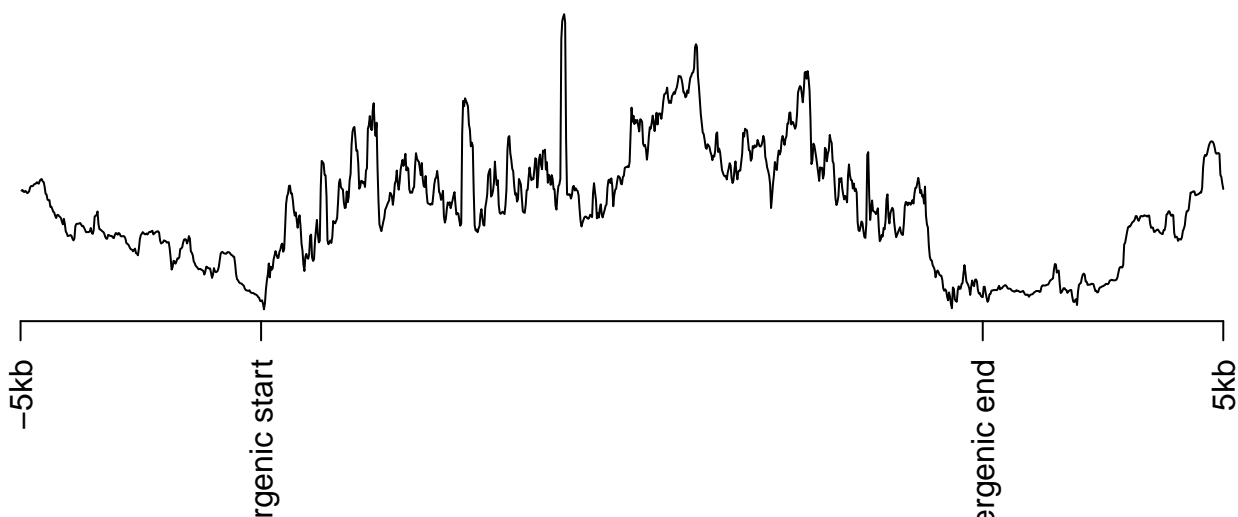
MouseNeuronCulture PositiveSingle E.697 scCLTdegenNuc1097



Average coverage per intergenic region

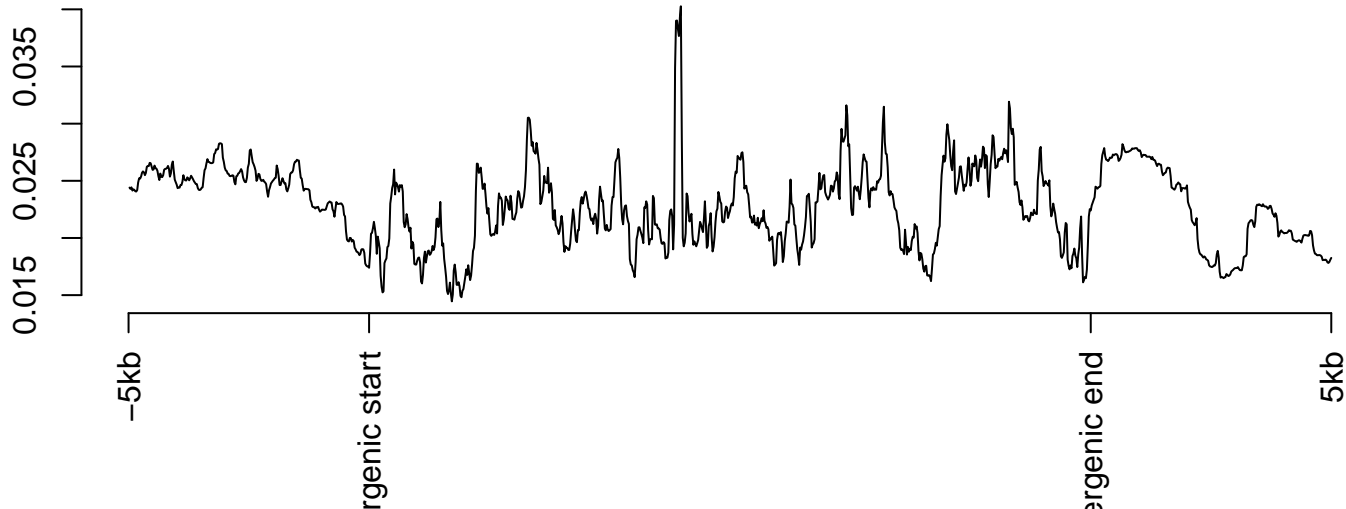
MouseNeuronCulture PositivePooled E.584 scCLTdegenNuc415

0.015 0.025 0.035



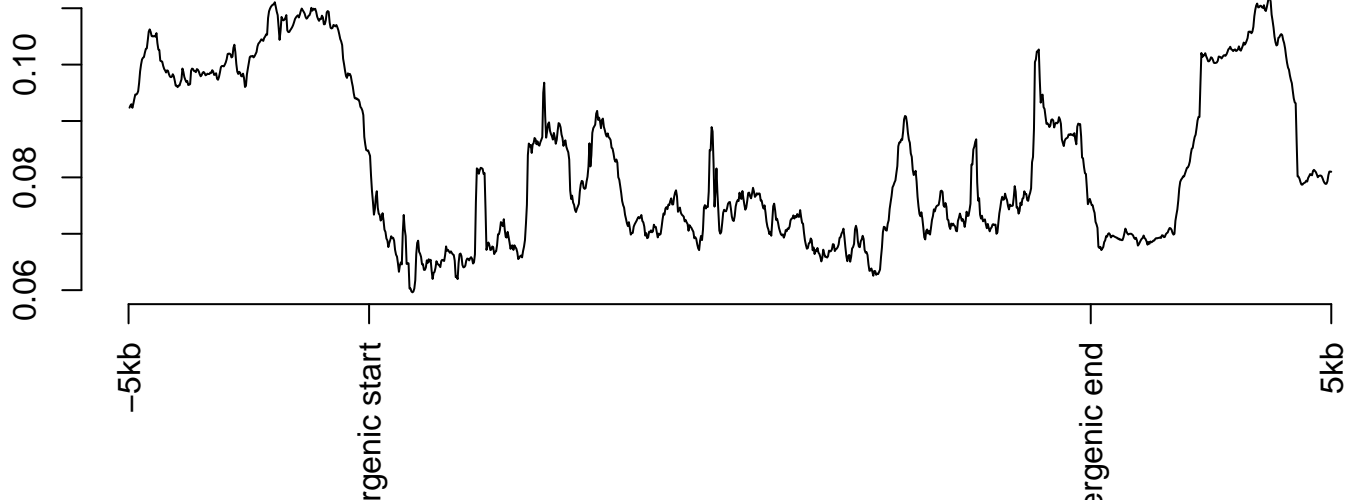
Average coverage per intergenic region

MouseNeuronCulture PositivePooled E.643 scCLTdegenNuc654



Average coverage per intergenic region

MouseNeuronCulture PositivePooled E.669 scCLTdegenNuc809



Average coverage per intergenic region

MouseNeuronCulture PositiveMerged E.– MouseNeuronCulturePositiveMerged

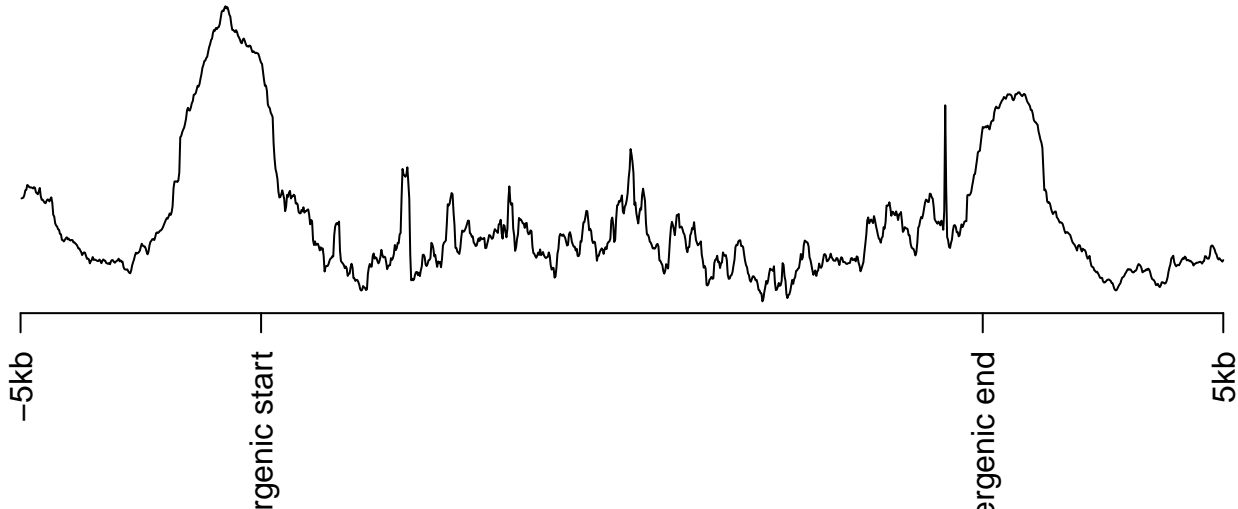
0.10
0.08
0.06

-5kb

intergenic start

intergenic end

5kb



Average coverage per intergenic region

MouseNeuronCulture PositiveAll E.– MouseNeuronCulturePositiveAll

