

Average coverage per intergenic region

MouseAstroCulture NegCtrlSingle E.586 scCLTdegenNuc435

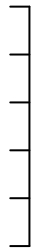
0.0004 0.0010

-5kb

intergenic start

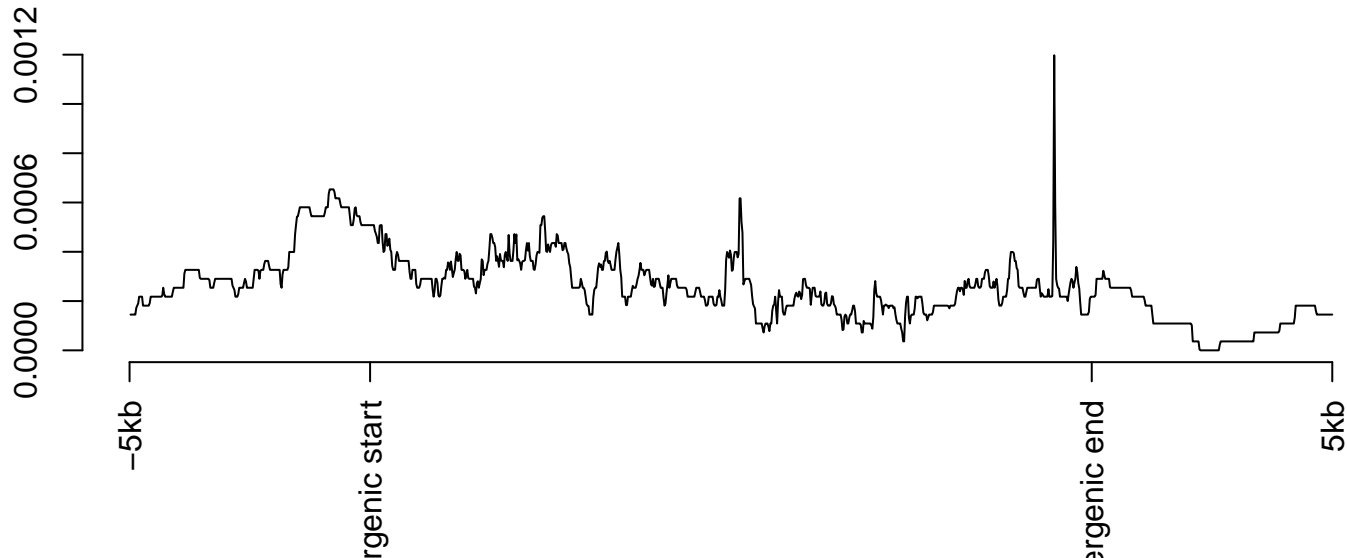
intergenic end

5kb



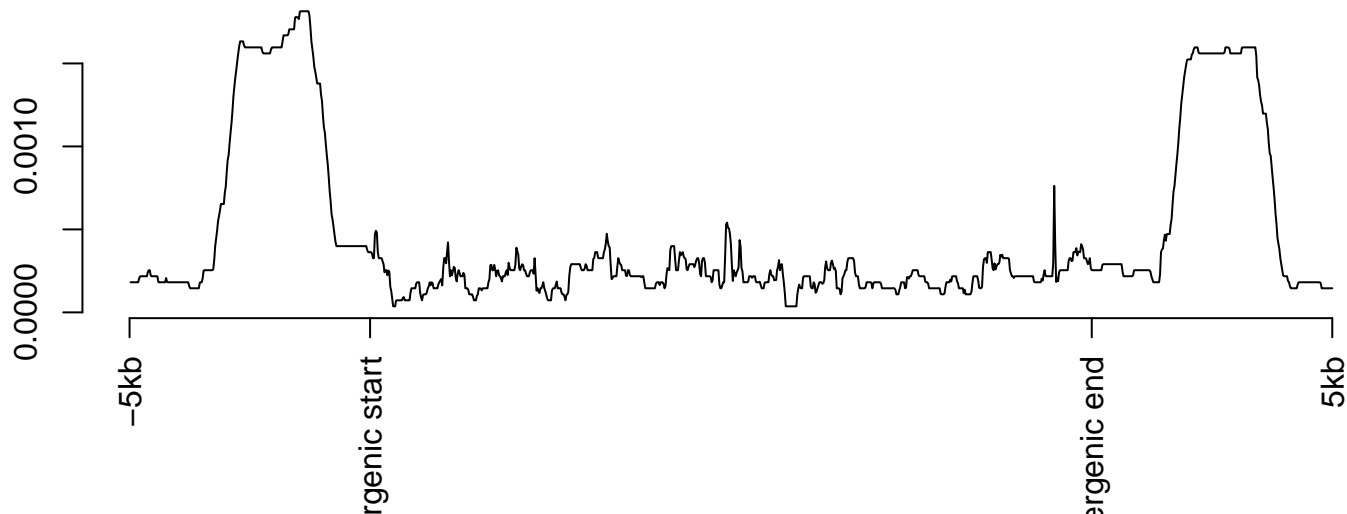
Average coverage per intergenic region

MouseAstroCulture NegCtrlSingle E.586 scCLTdegenNuc436



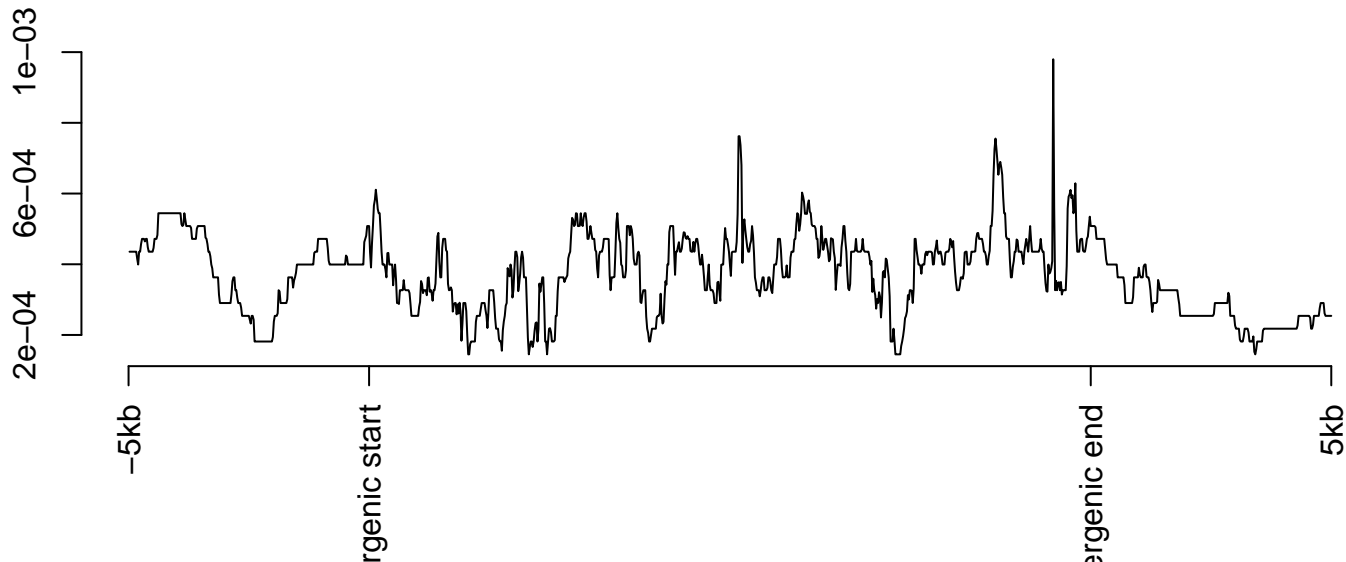
Average coverage per intergenic region

MouseAstroCulture NegCtrlSingle E.586 scCLTdegenNuc437



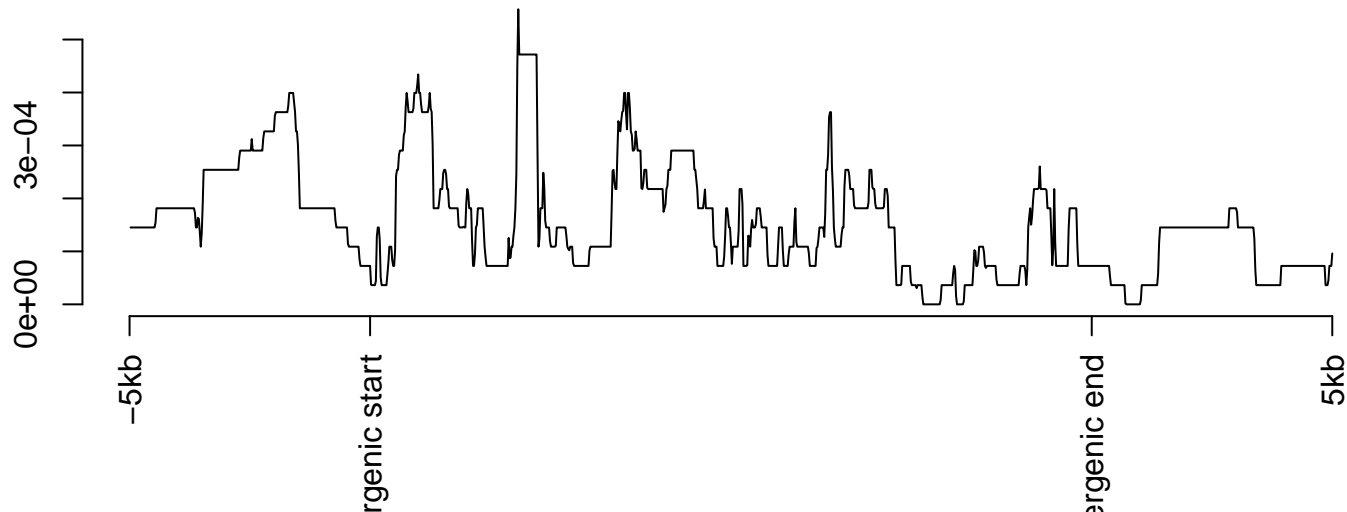
Average coverage per intergenic region

MouseAstroCulture NegCtrlSingle E.586 scCLTdegenNuc438



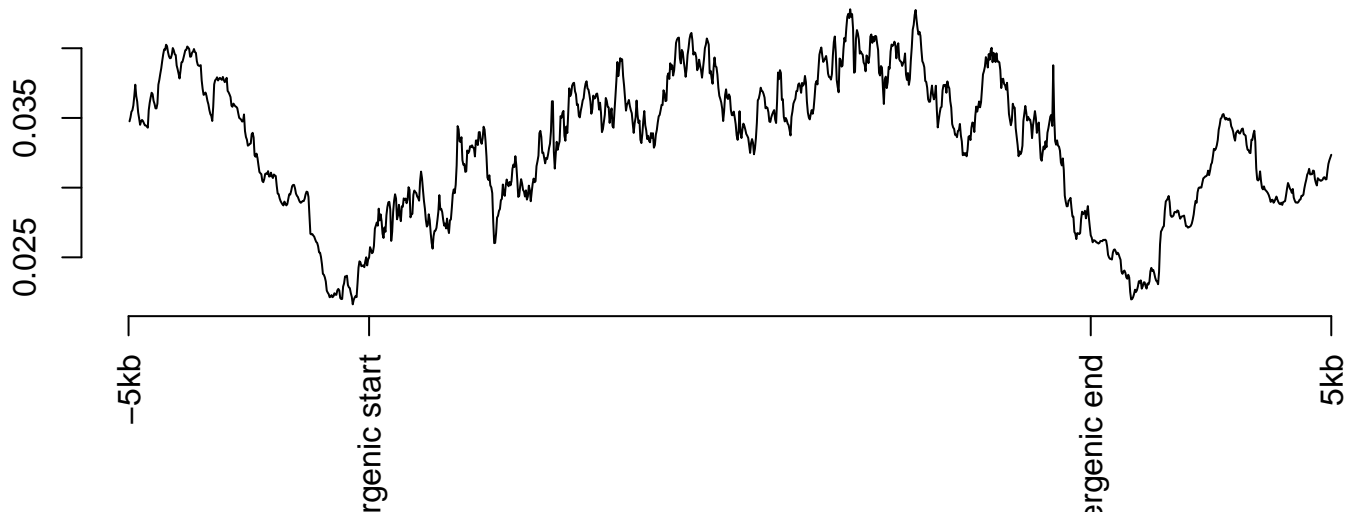
Average coverage per intergenic region

MouseAstroCulture NegCtrlSingle E.679 scCLTdegenNuc964



Average coverage per intergenic region

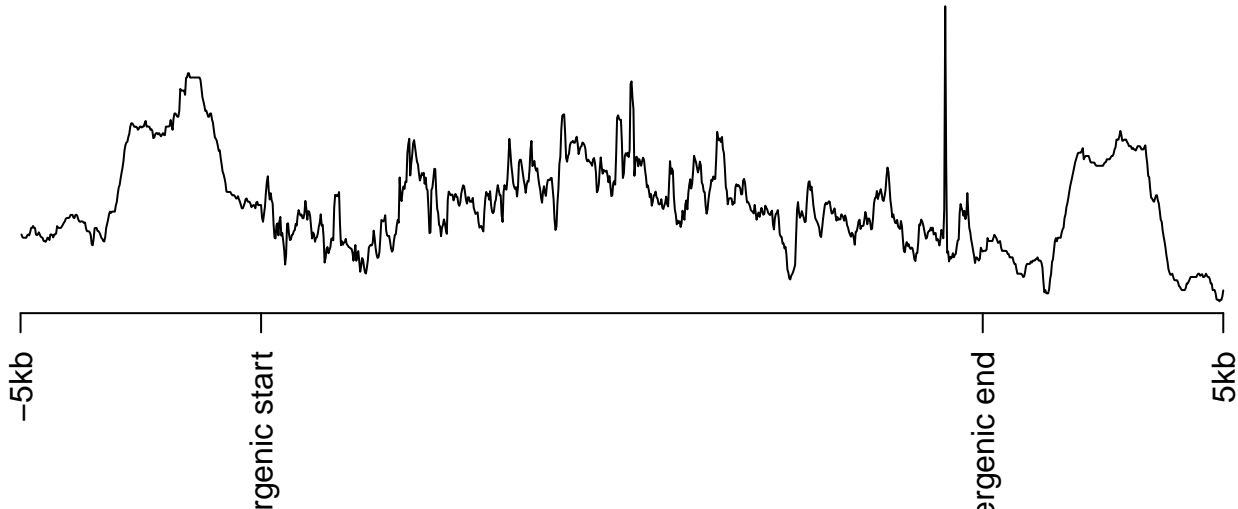
MouseAstroCulture NegCtrlPooled E.597 scCLTdegenNuc472



Average coverage per intergenic region

MouseAstroCulture NegCtrlMerged E.– MouseAstroCultureNegCtrlMerged

0.0040
0.0025
0.0010



Average coverage per intergenic region

MouseAstroCulture NegCtrlAll E.– MouseAstroCultureNegCtrlAll

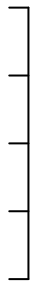
0.045
0.035
0.025

-5kb

intergenic start

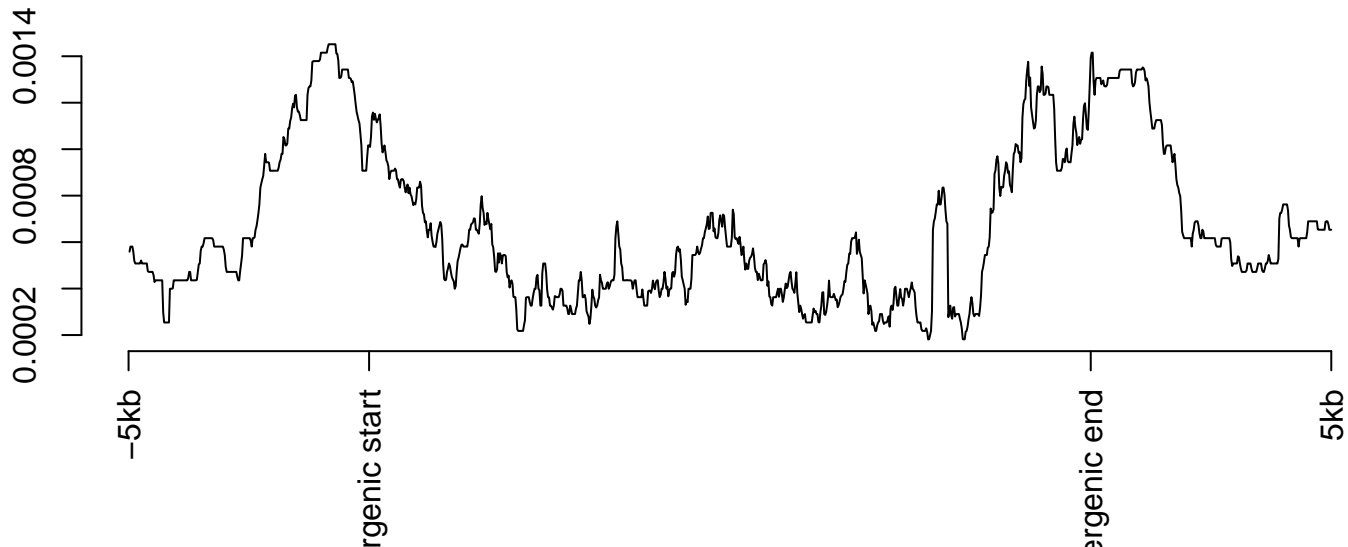
intergenic end

5kb



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.586 scCLTdegenNuc394



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.586 scCLTdegenNuc395

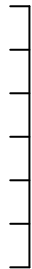
0.0002 0.0008 0.0014

-5kb

intergenic start

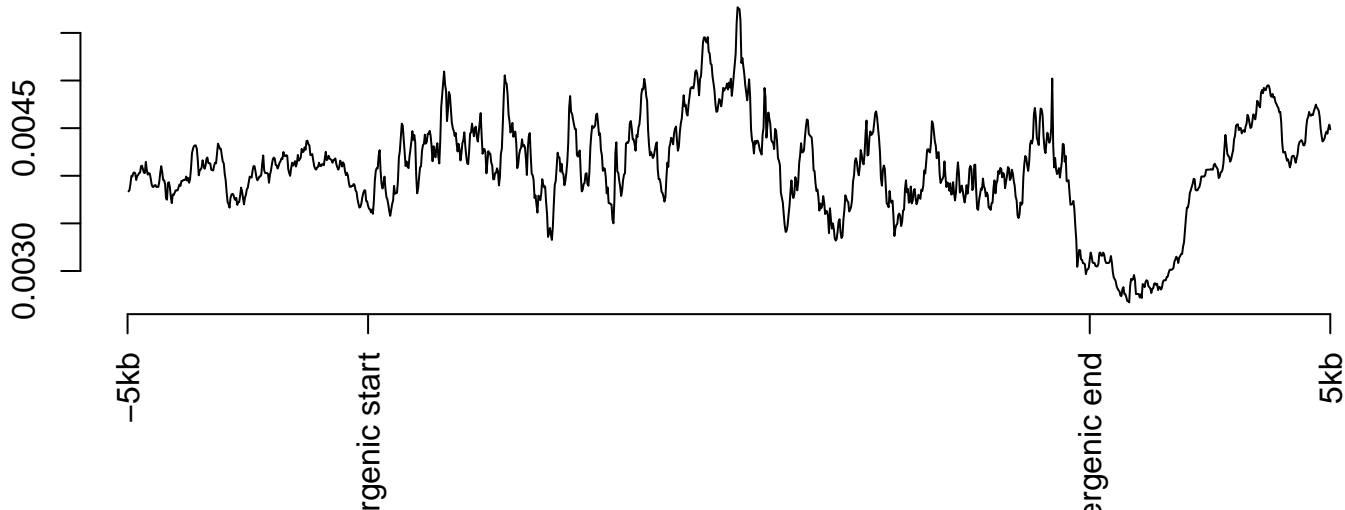
intergenic end

5kb



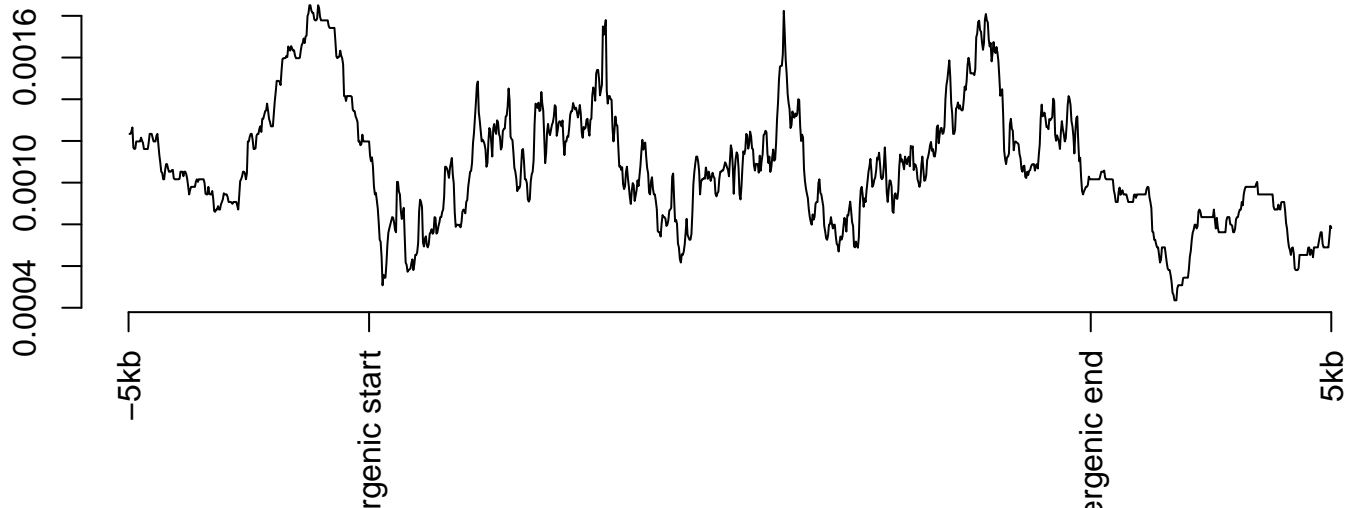
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.586 scCLTdegenNuc397



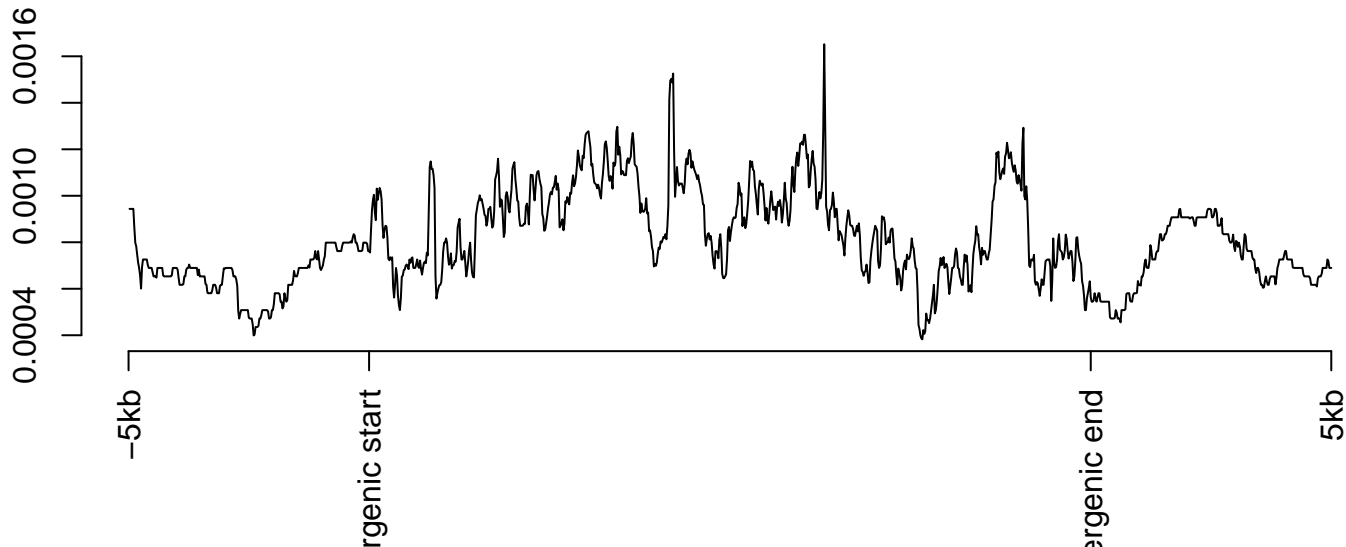
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.586 scCLTdegenNuc398



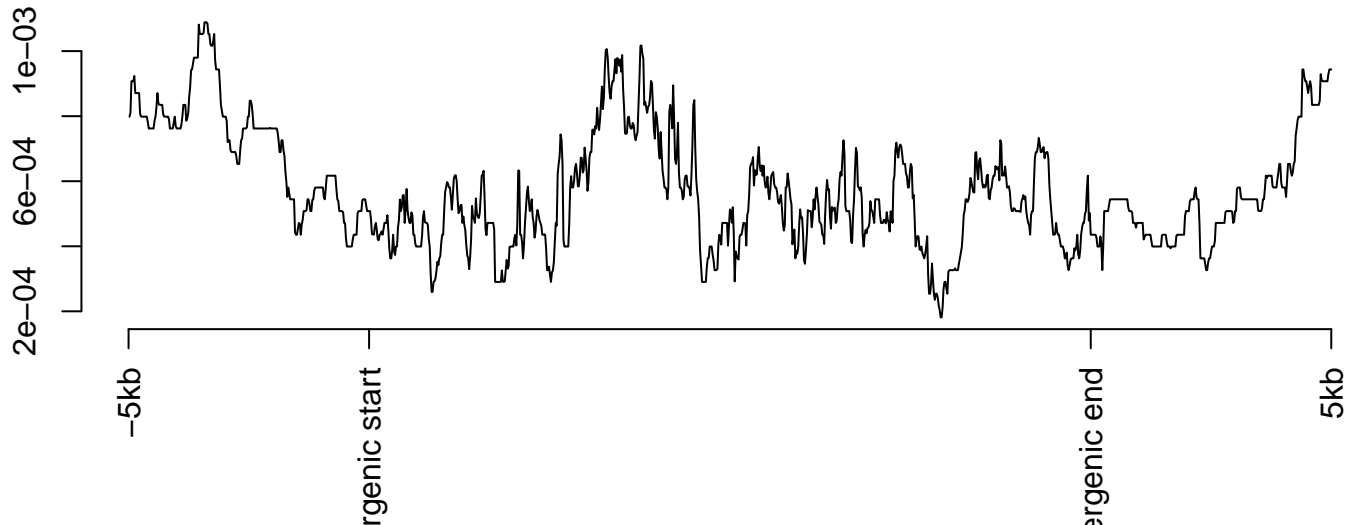
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.586 scCLTdegenNuc399



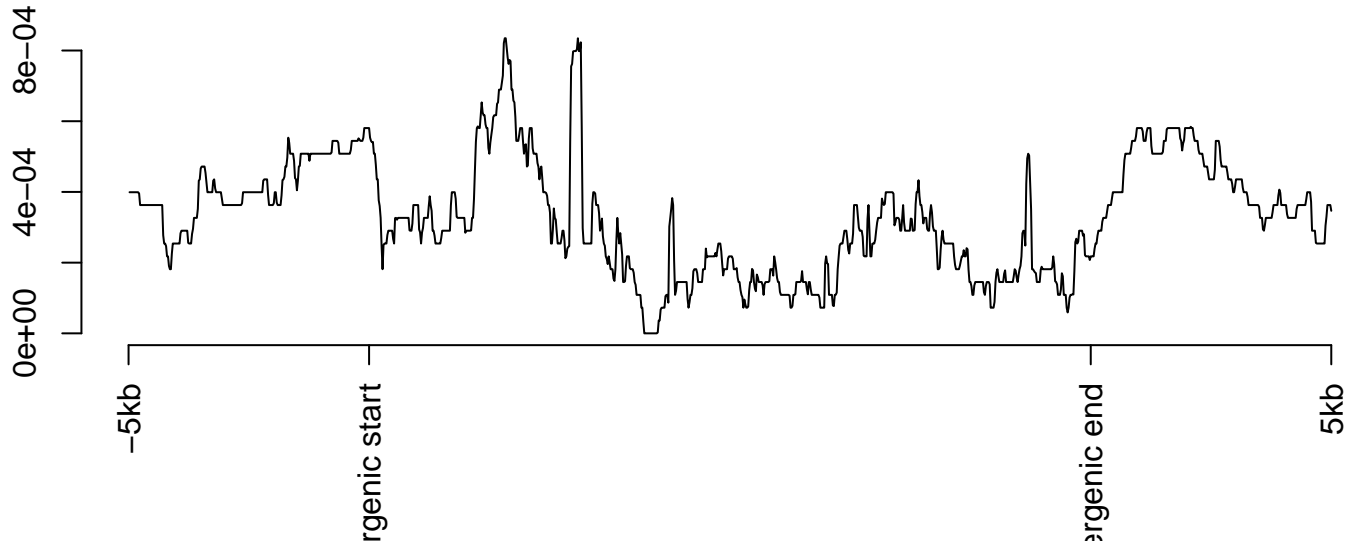
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.586 scCLTdegenNuc400



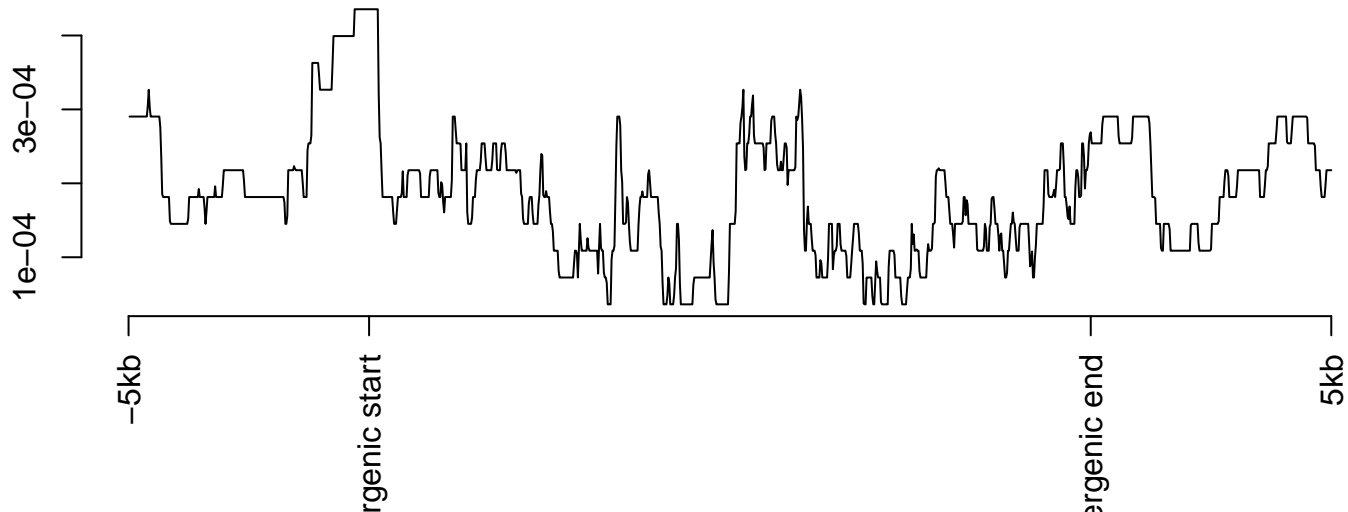
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.586 scCLTdegenNuc401



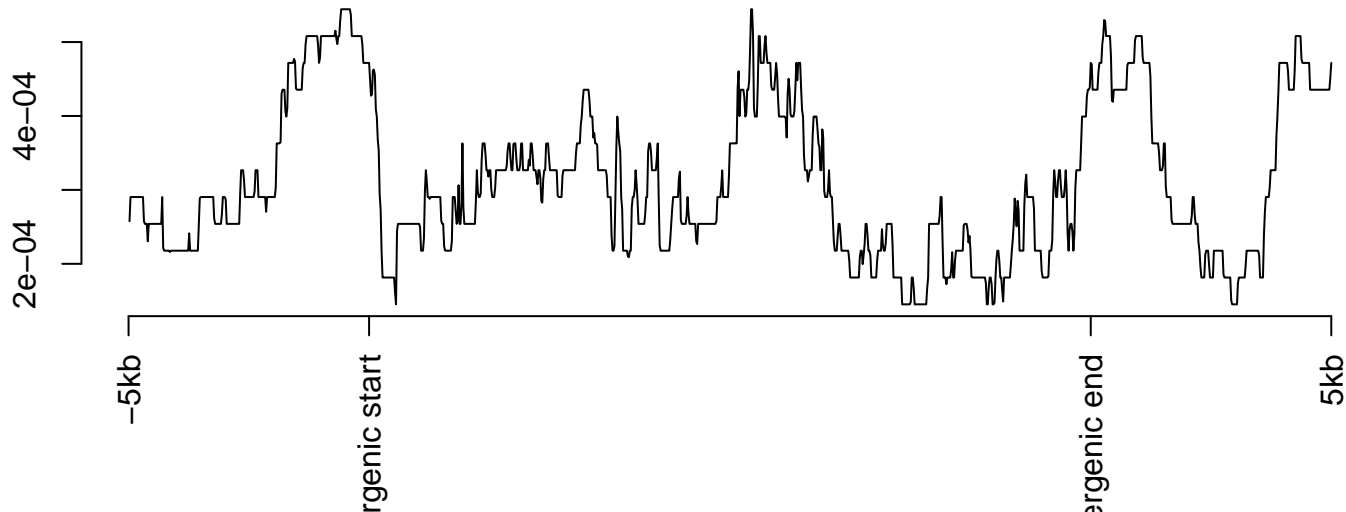
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.619 scCLTdegenNuc496



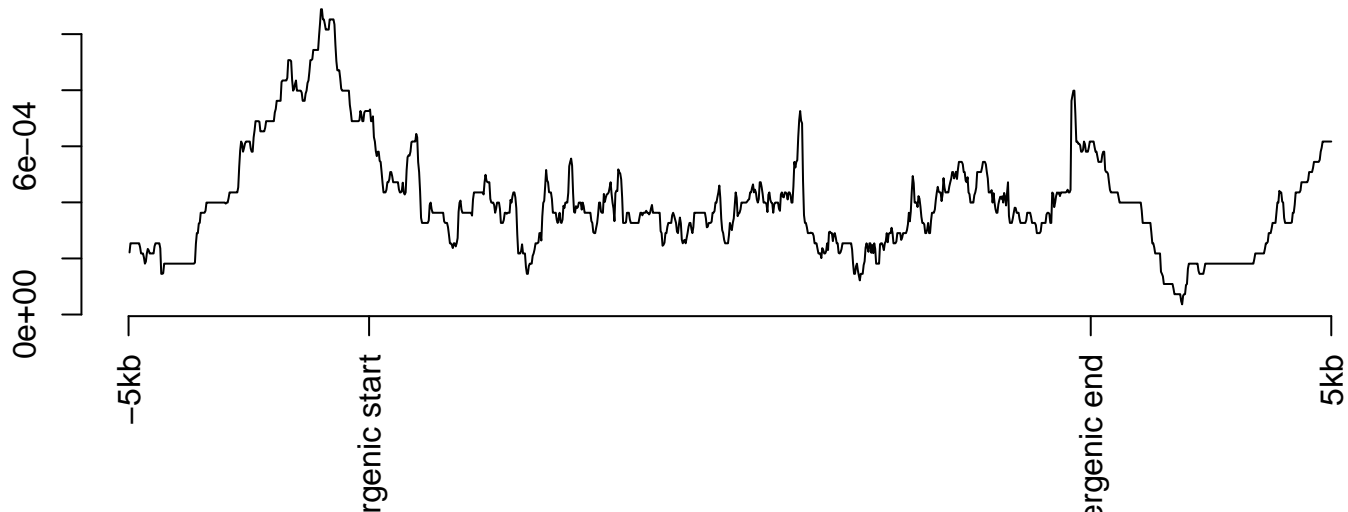
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.619 scCLTdegenNuc497



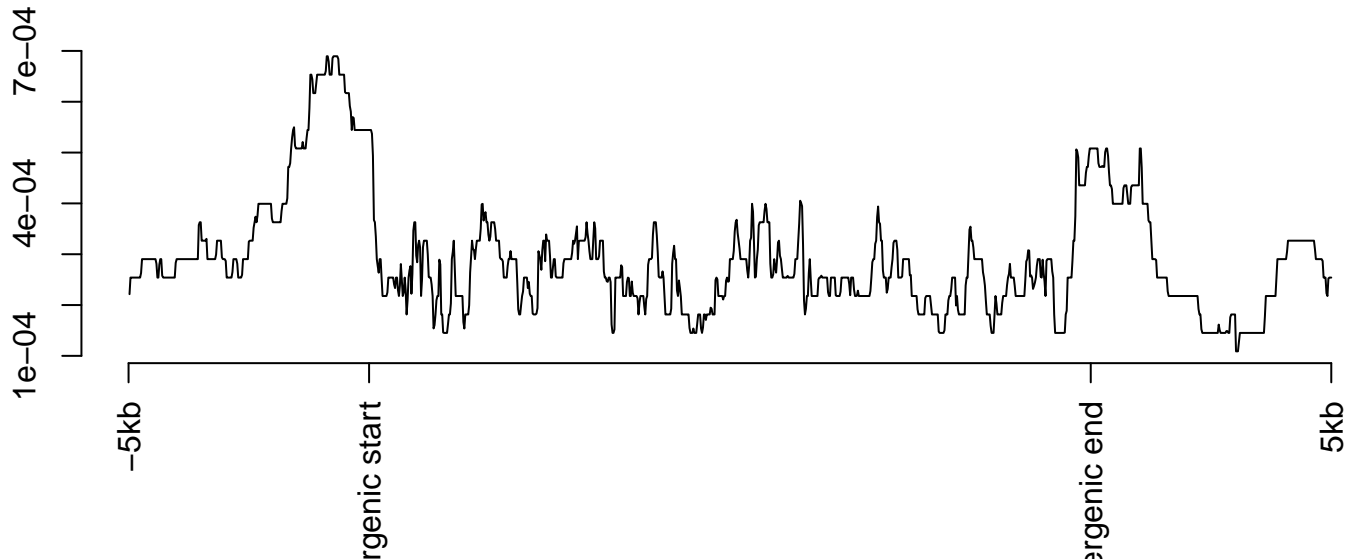
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.619 scCLTdegenNuc498



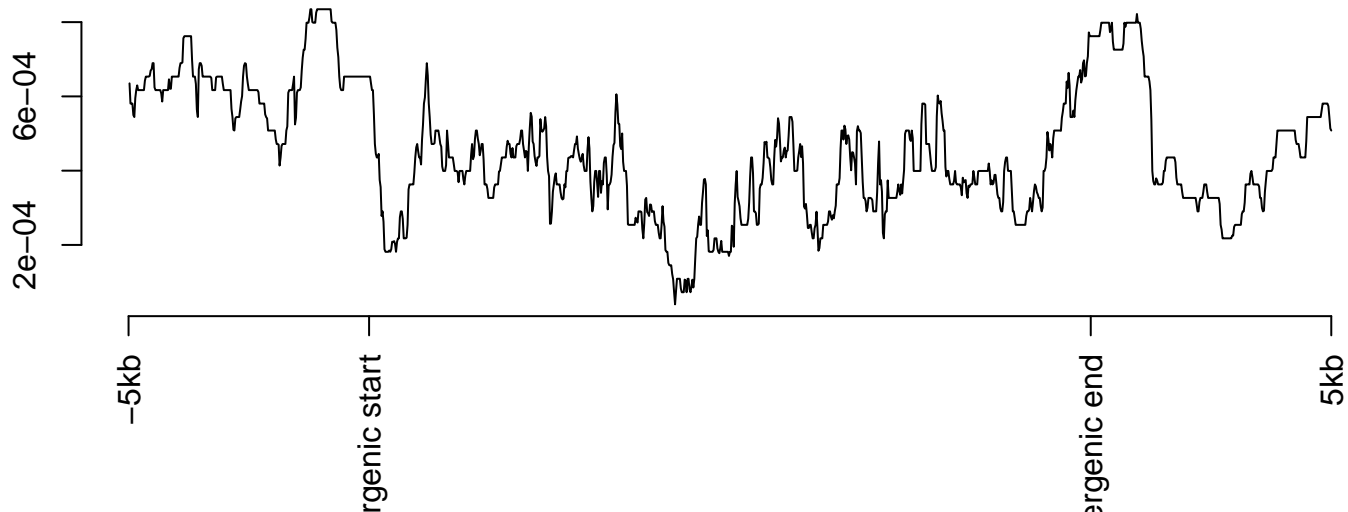
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.619 scCLTdegenNuc499



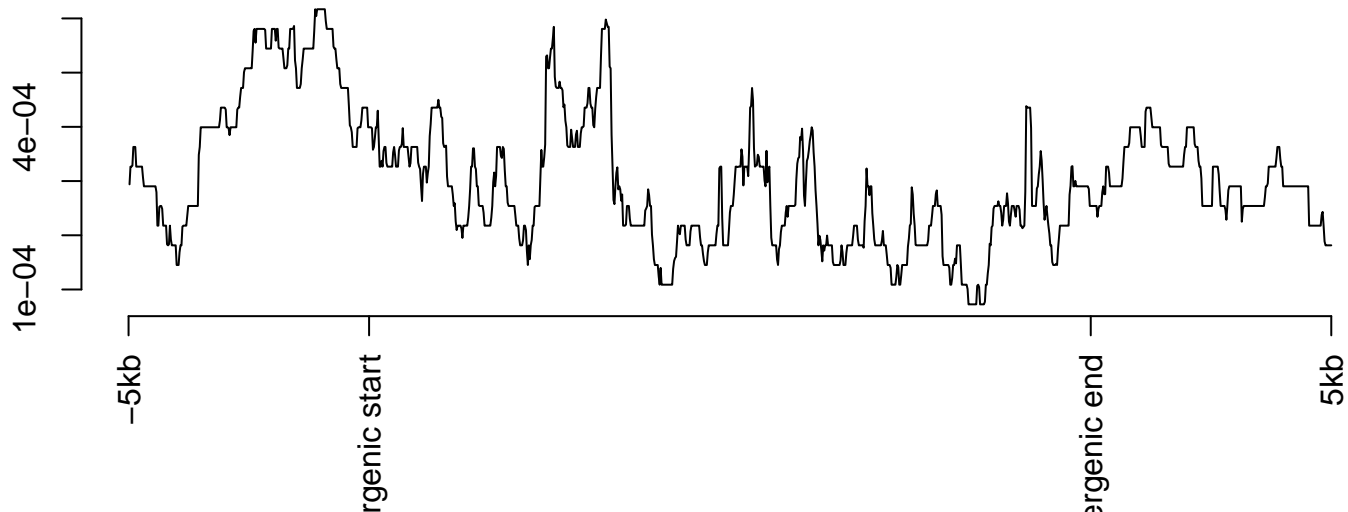
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.619 scCLTdegenNuc500



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.619 scCLTdegenNuc506



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.622 scCLTdegenNuc507

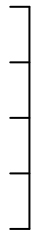
2e-04
5e-04

-5kb

intergenic start

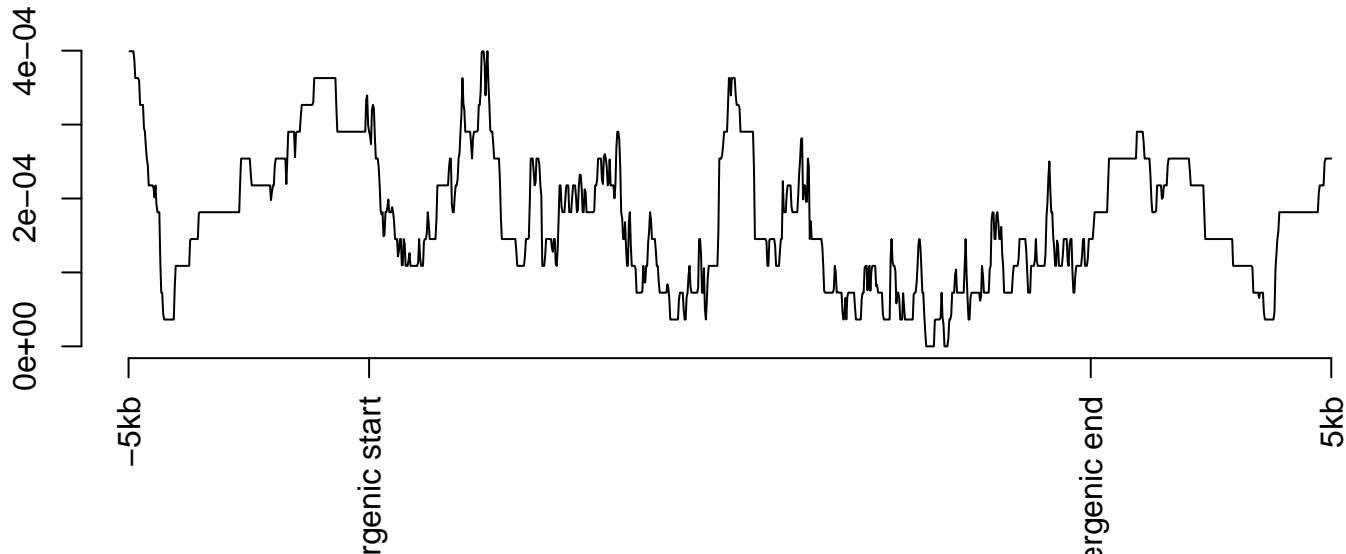
intergenic end

5kb



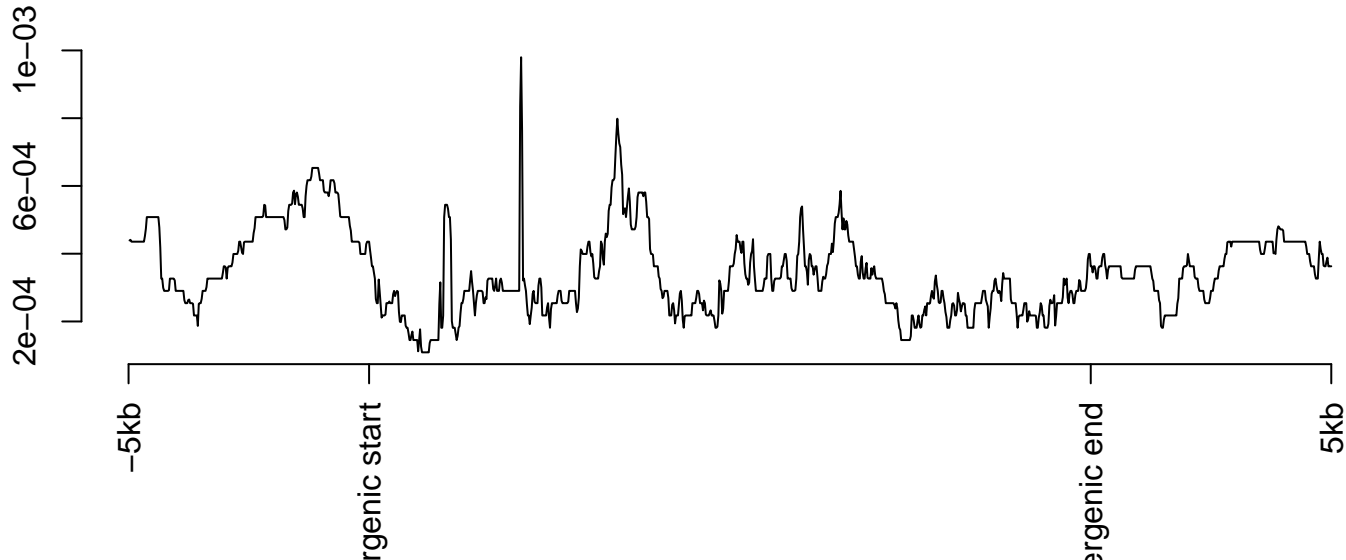
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.622 scCLTdegenNuc508



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.622 scCLTdegenNuc509



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.622 scCLTdegenNuc514

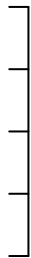
0.0010 0.0020 0.0030

-5kb

intergenic start

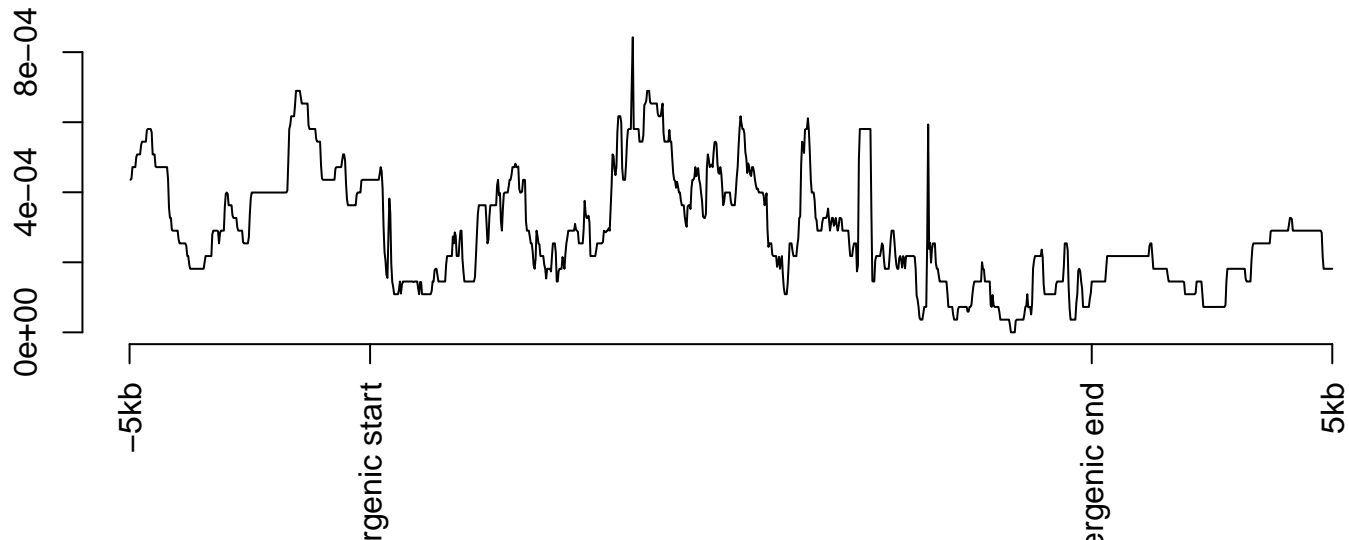
intergenic end

5kb



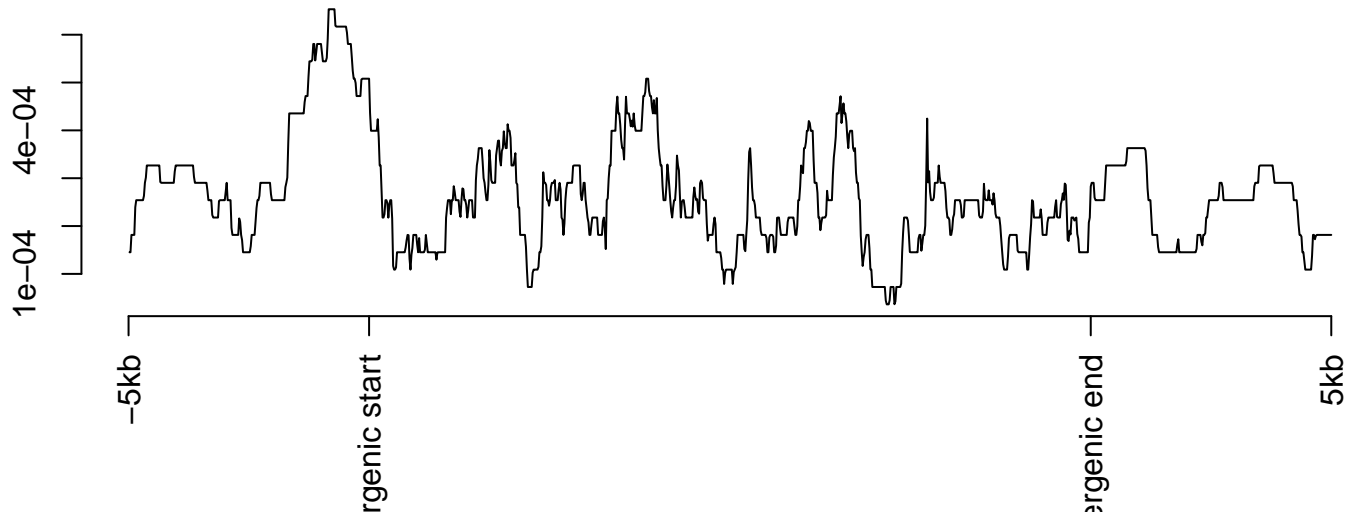
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.667 scCLTdegenNuc812



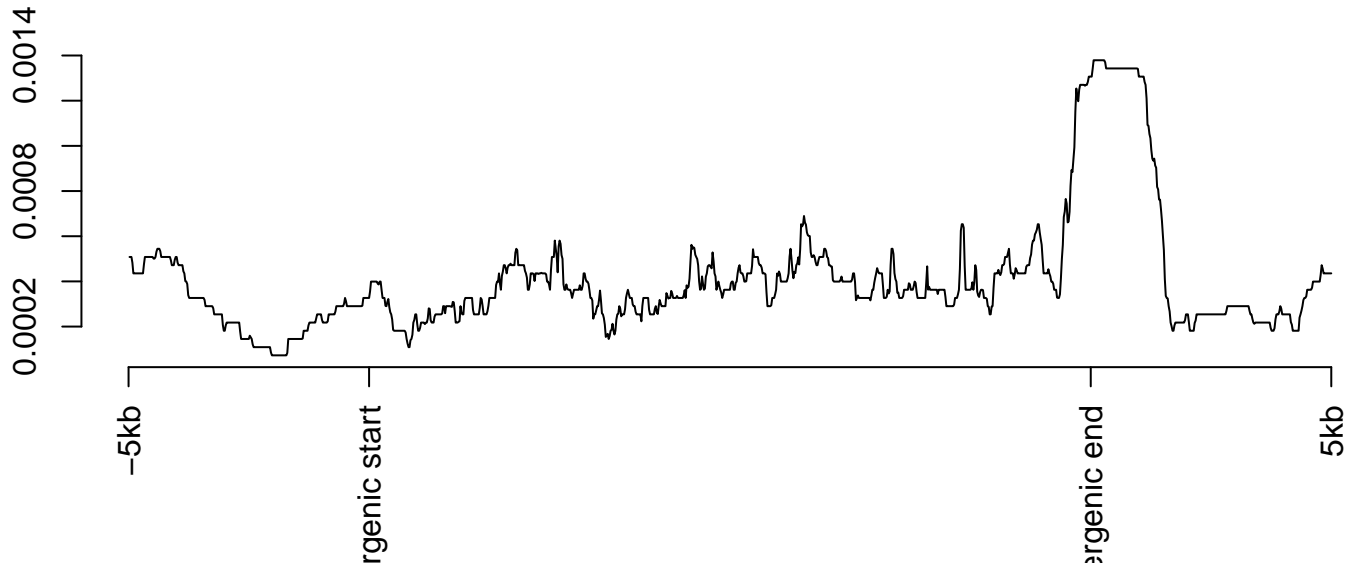
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.667 scCLTdegenNuc813



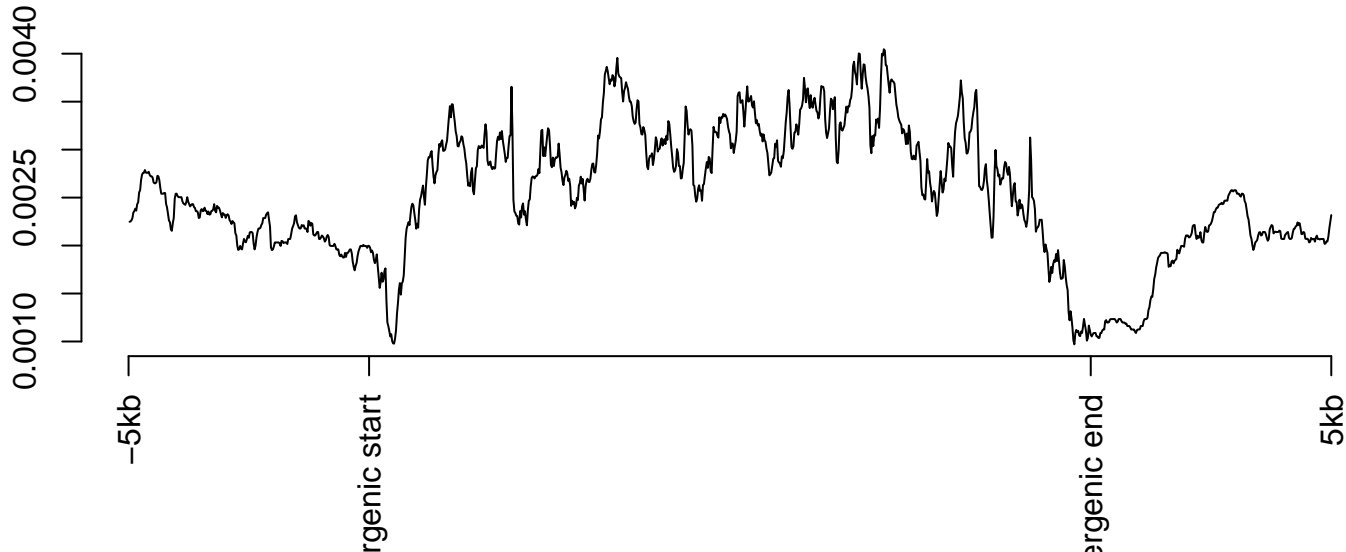
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.667 scCLTdegenNuc814



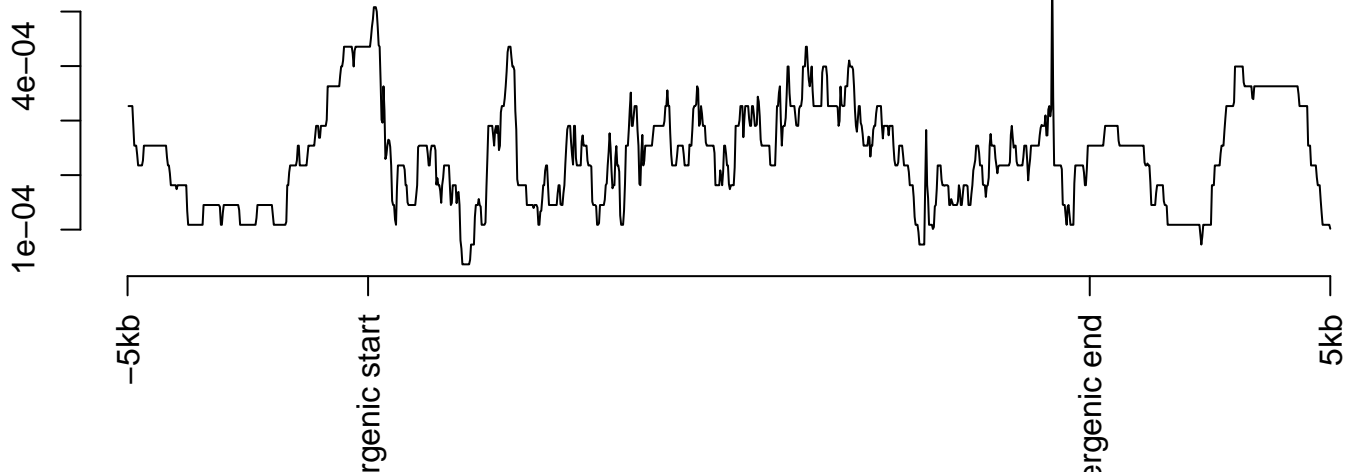
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.667 scCLTdegenNuc815



Average coverage per intergenic region

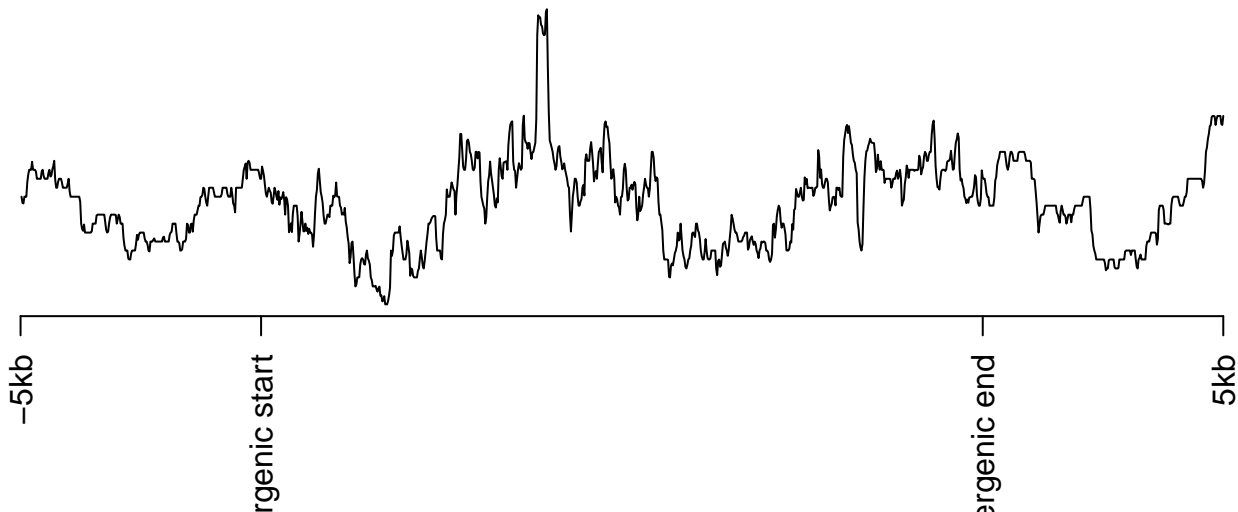
MouseAstroCulture PositiveSingle E.667 scCLTdegenNuc816



MouseAstroCulture PositiveSingle E.667 scCLTdegenNuc819

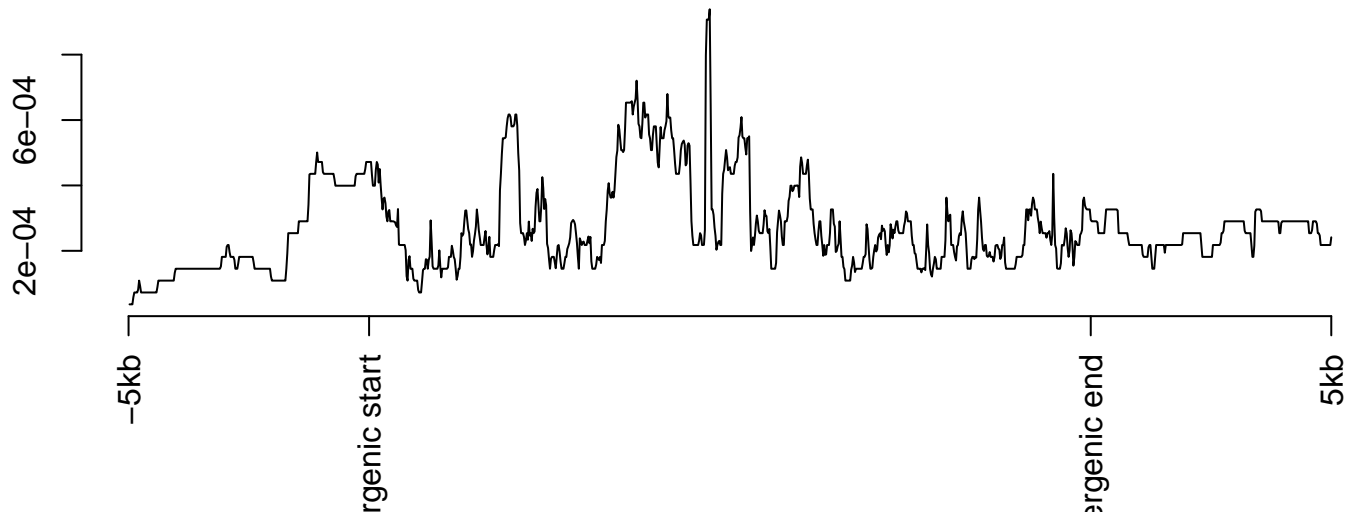
Average coverage per intergenic region

0.0010
0.0004



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.667 scCLTdegenNuc820



MouseAstroCulture PositiveSingle E.670 scCLTdegenNuc835

Average coverage per intergenic region

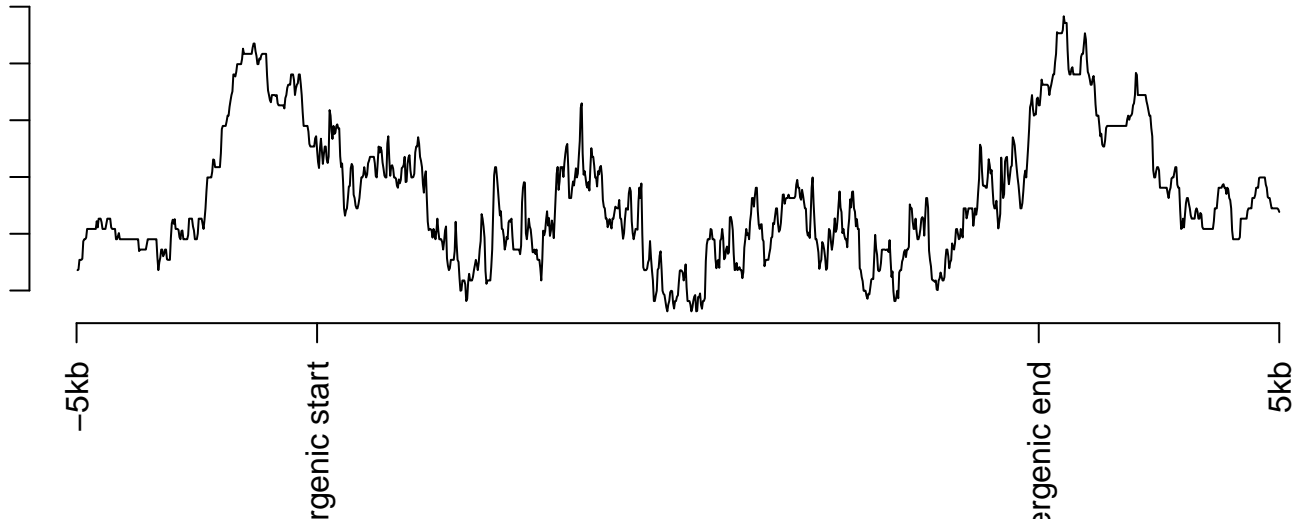
0.0010
0.0004

-5kb

intergenic start

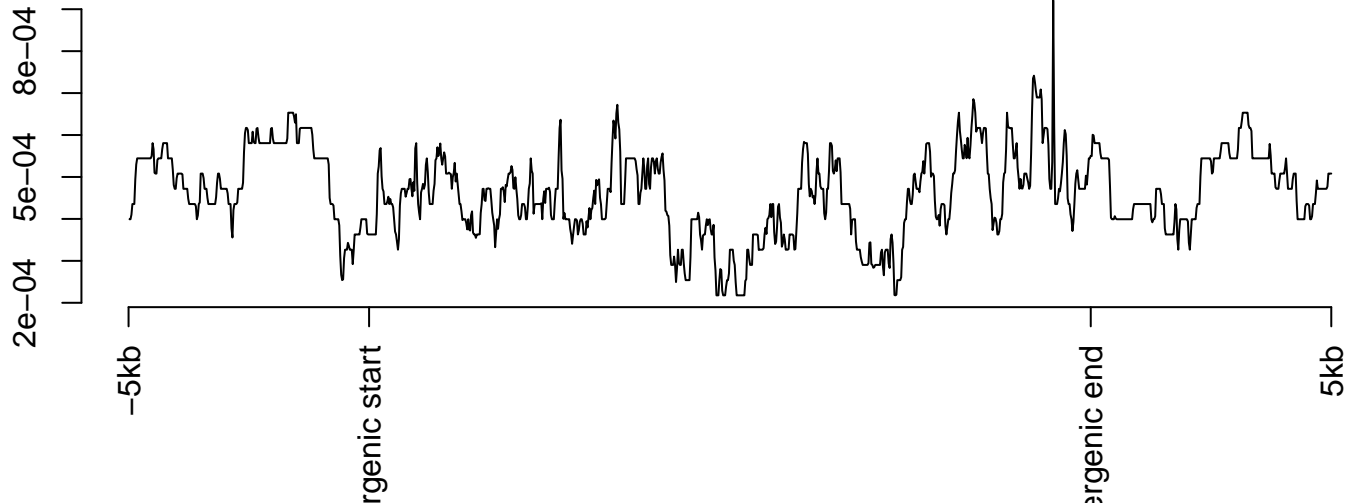
intergenic end

5kb



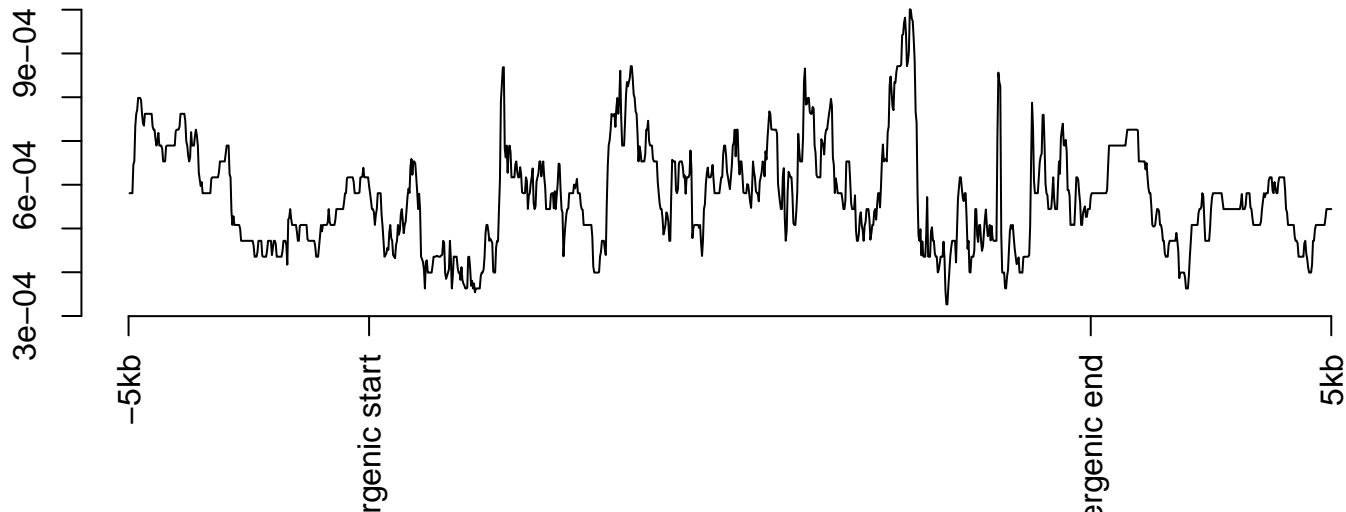
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.670 scCLTdegenNuc836



Average coverage per intergenic region

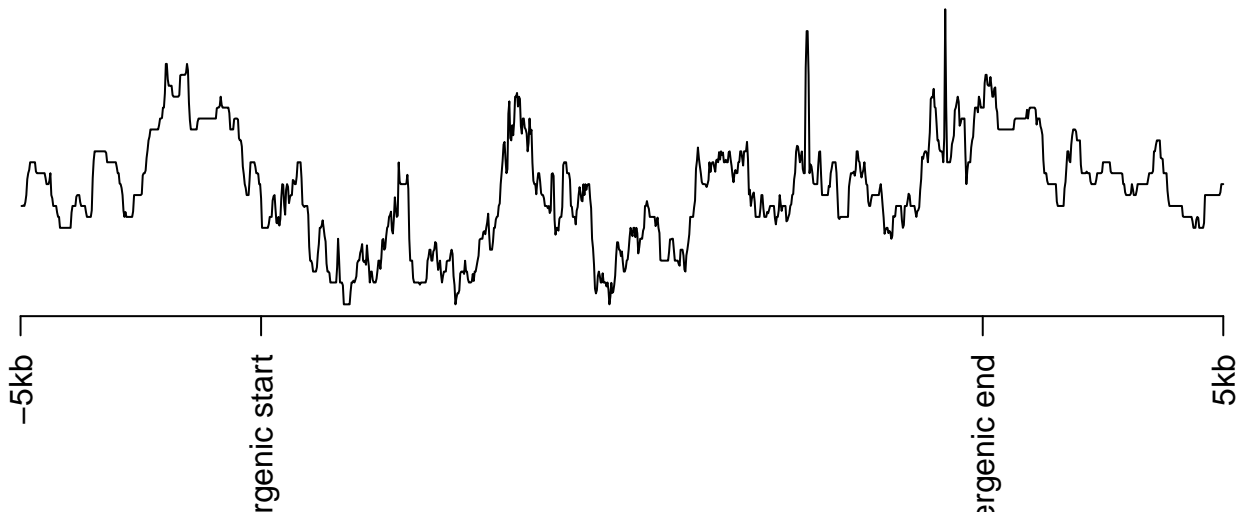
MouseAstroCulture PositiveSingle E.670 scCLTdegenNuc837



Average coverage per intergenic region

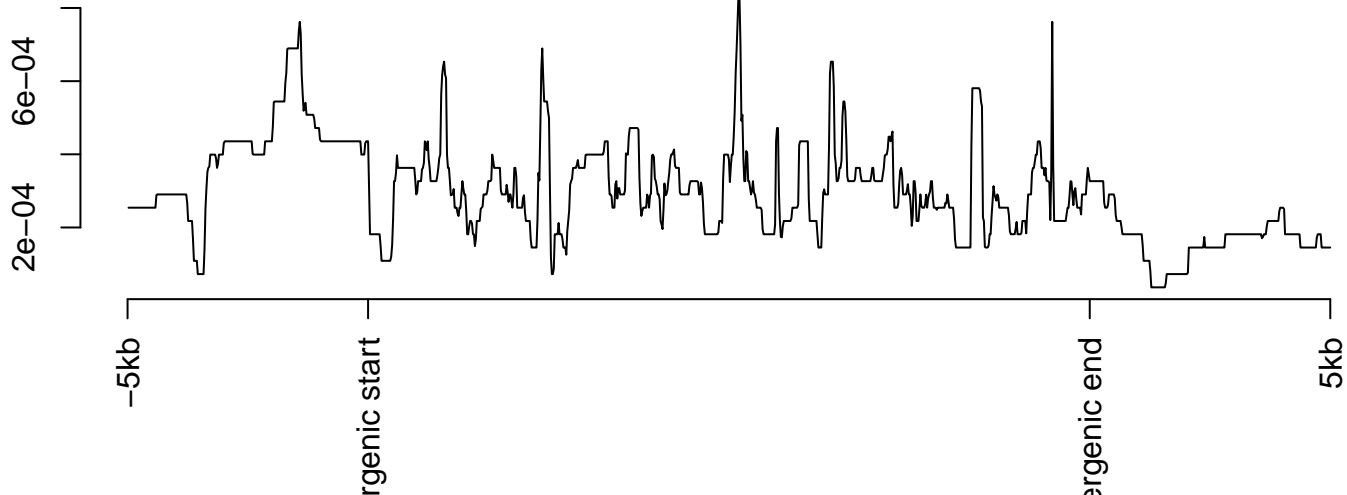
MouseAstroCulture PositiveSingle E.670 scCLTdegenNuc838

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



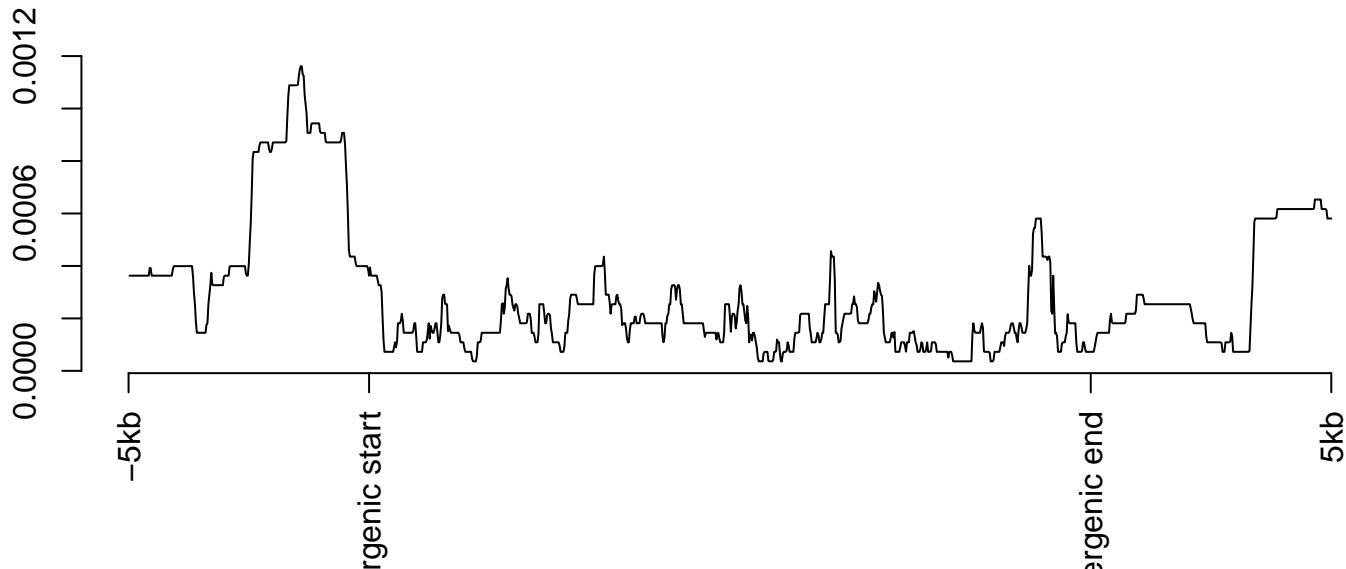
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.679 scCLTdegenNuc958



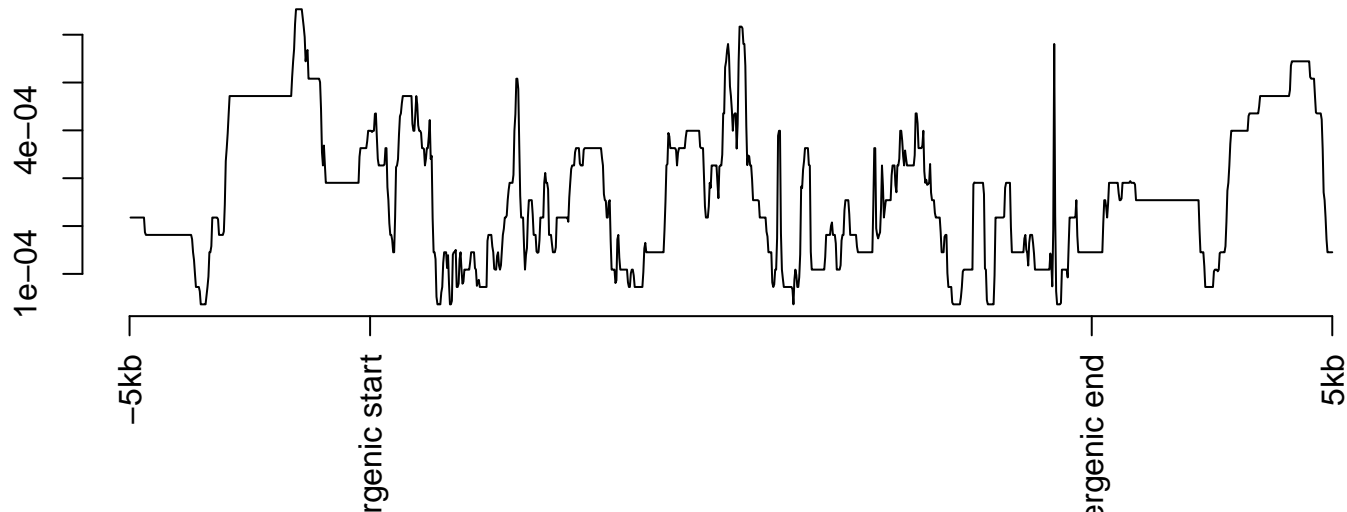
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.679 scCLTdegenNuc959



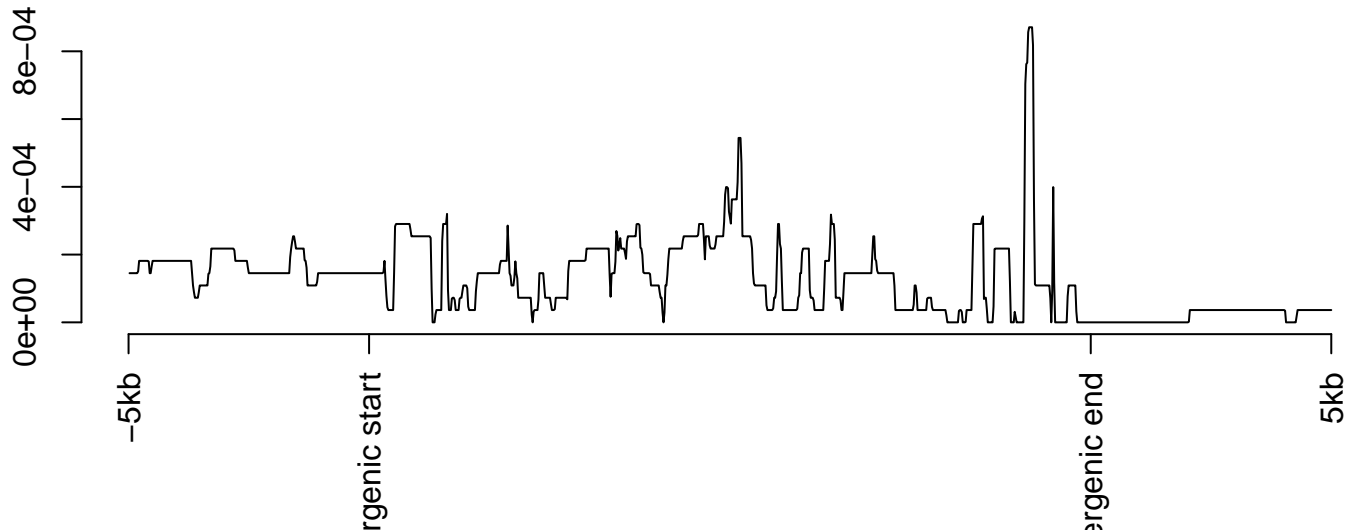
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.679 scCLTdegenNuc960



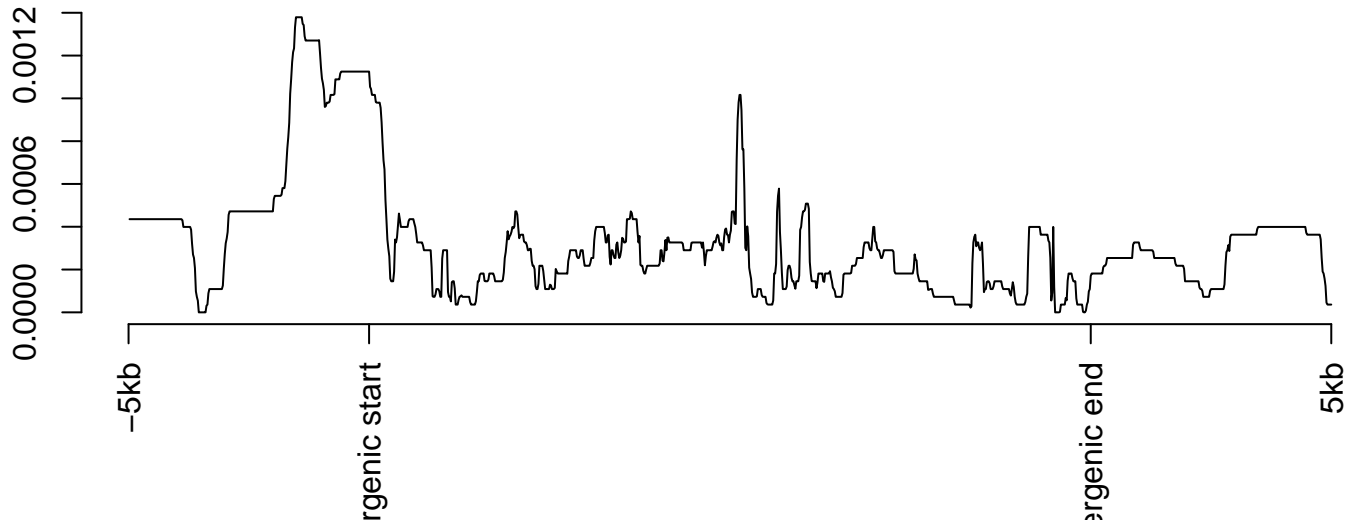
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.679 scCLTdegenNuc961



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.679 scCLTdegenNuc962



Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.679 scCLTdegenNuc963

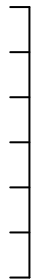
0.0014
0.0008
0.0002

-5kb

intergenic start

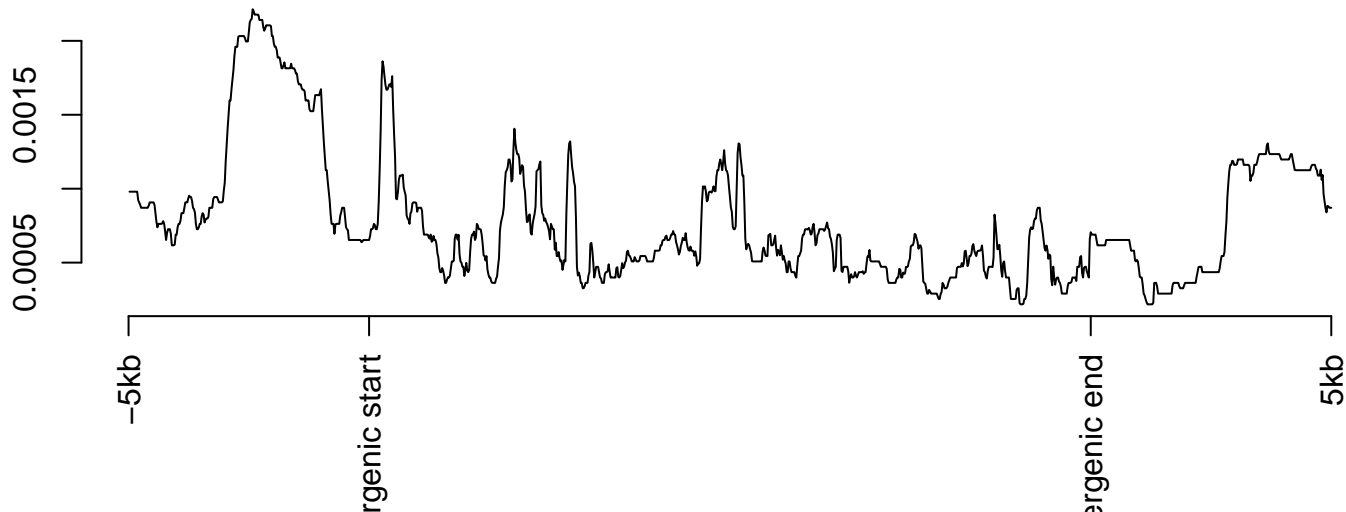
intergenic end

5kb



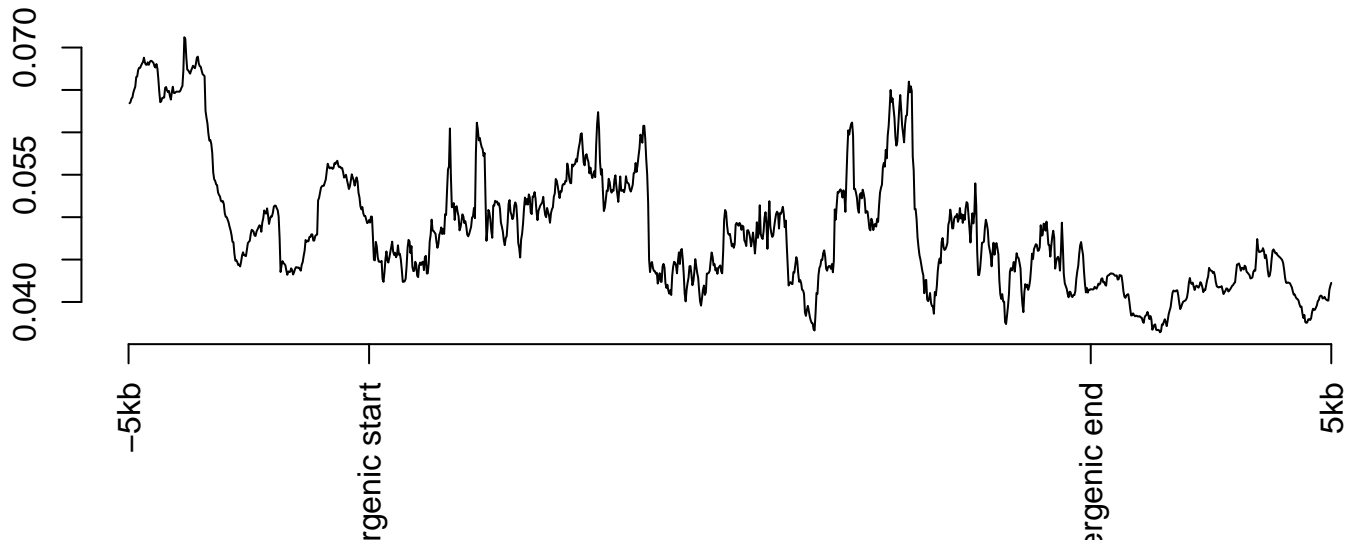
Average coverage per intergenic region

MouseAstroCulture PositiveSingle E.697 scCLTdegenNuc1094



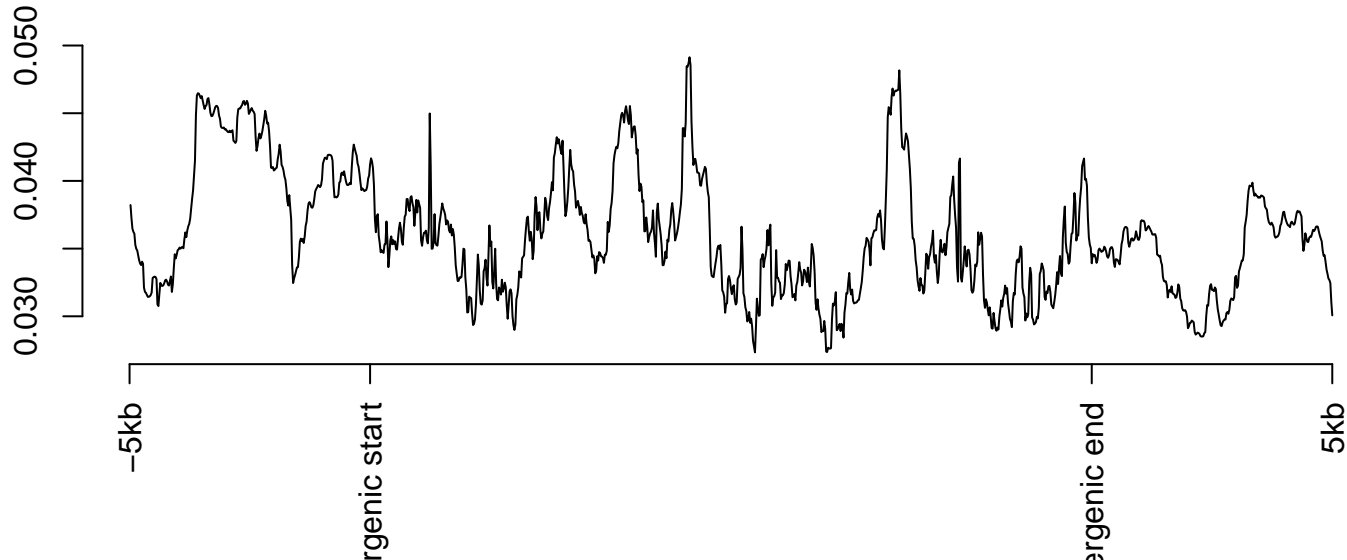
Average coverage per intergenic region

MouseAstroCulture PositivePooled E.586 scCLTdegenNuc414



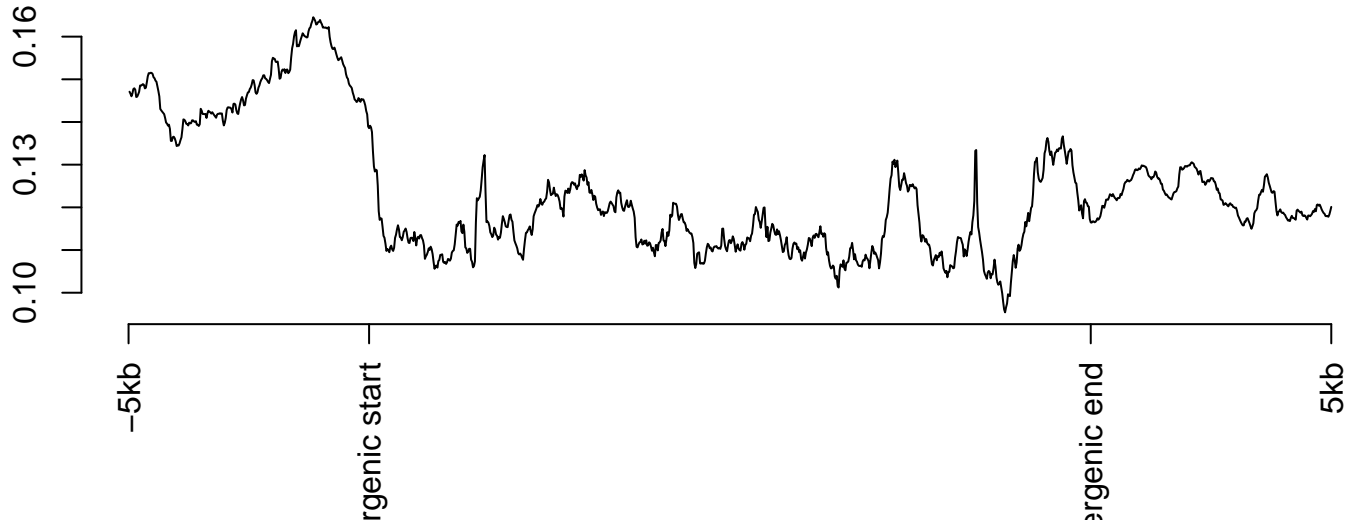
Average coverage per intergenic region

MouseAstroCulture PositivePooled E.643 scCLTdegenNuc655



Average coverage per intergenic region

MouseAstroCulture PositivePooled E.667 scCLTdegenNuc821



Average coverage per intergenic region

MouseAstroCulture PositiveMerged E.– MouseAstroCulturePositiveMerged

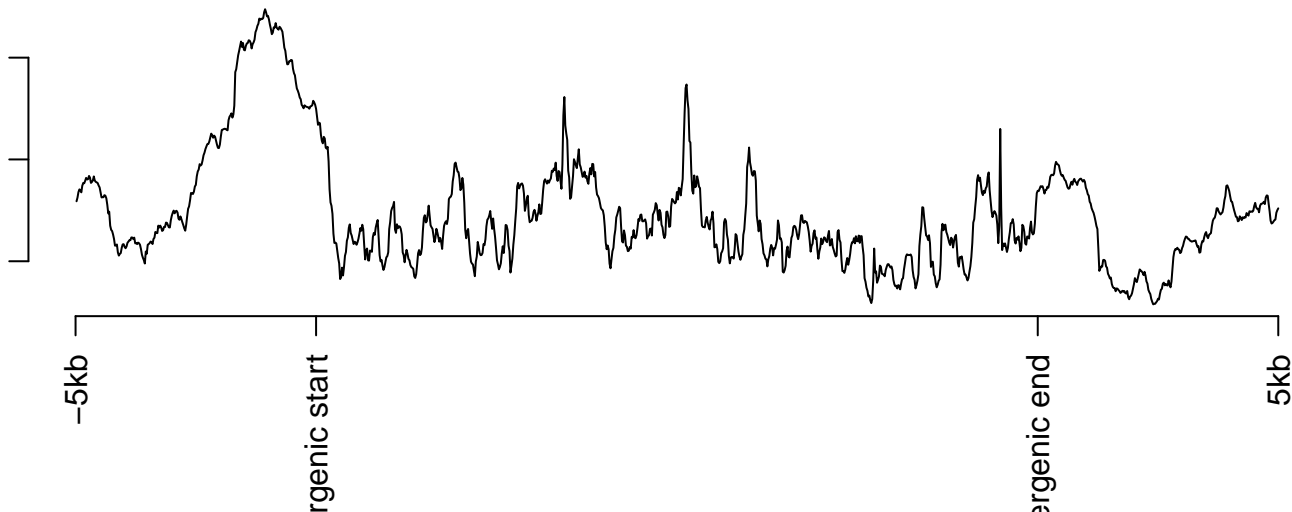
0.030
0.020

-5kb

intergenic start

intergenic end

5kb



Average coverage per intergenic region

MouseAstroCulture PositiveAll E.– MouseAstroCulturePositiveAll

0.28
0.24
0.20

-5kb

intergenic start

intergenic end

5kb

