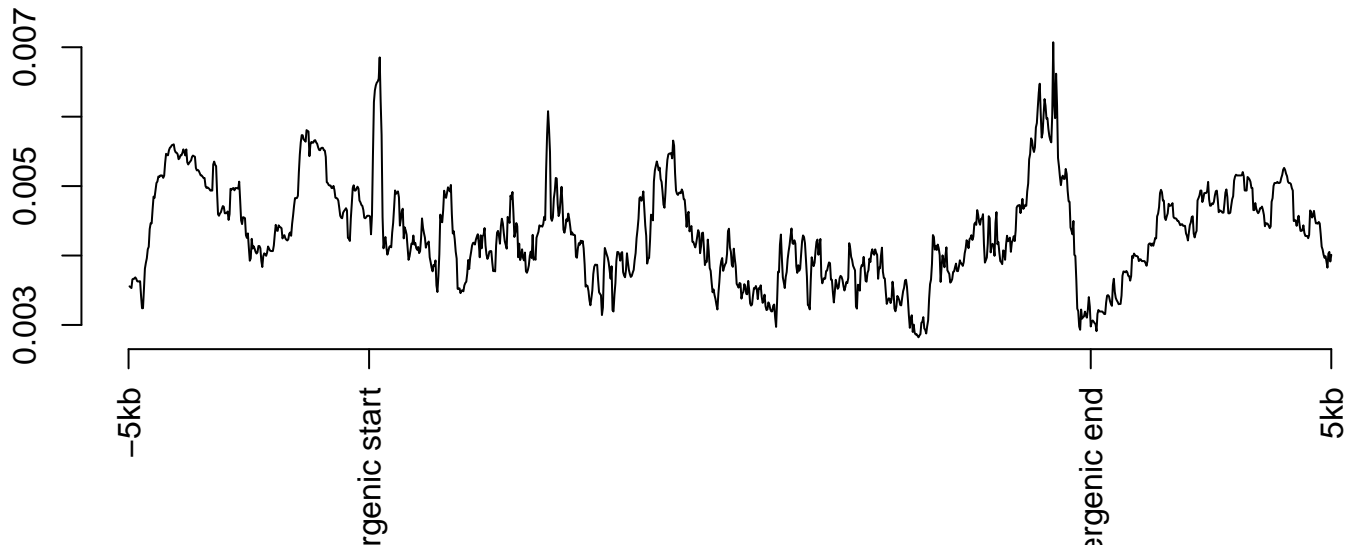


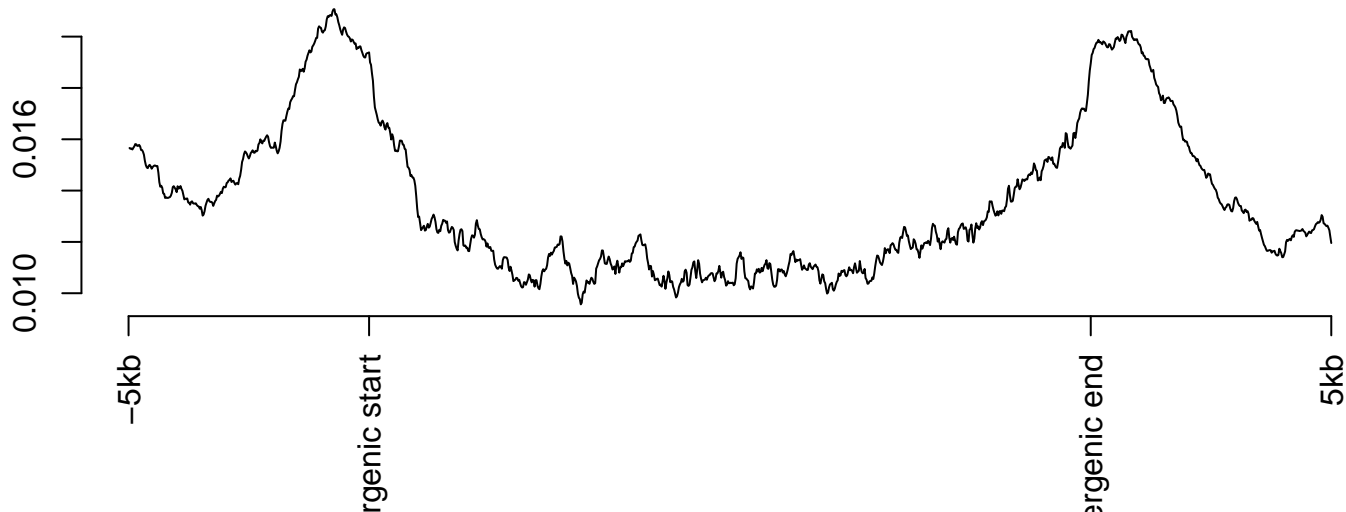
Average coverage per intergenic region

MouseNeuronSlice NegCtrlPooled E.597 scCLTdegenNuc470



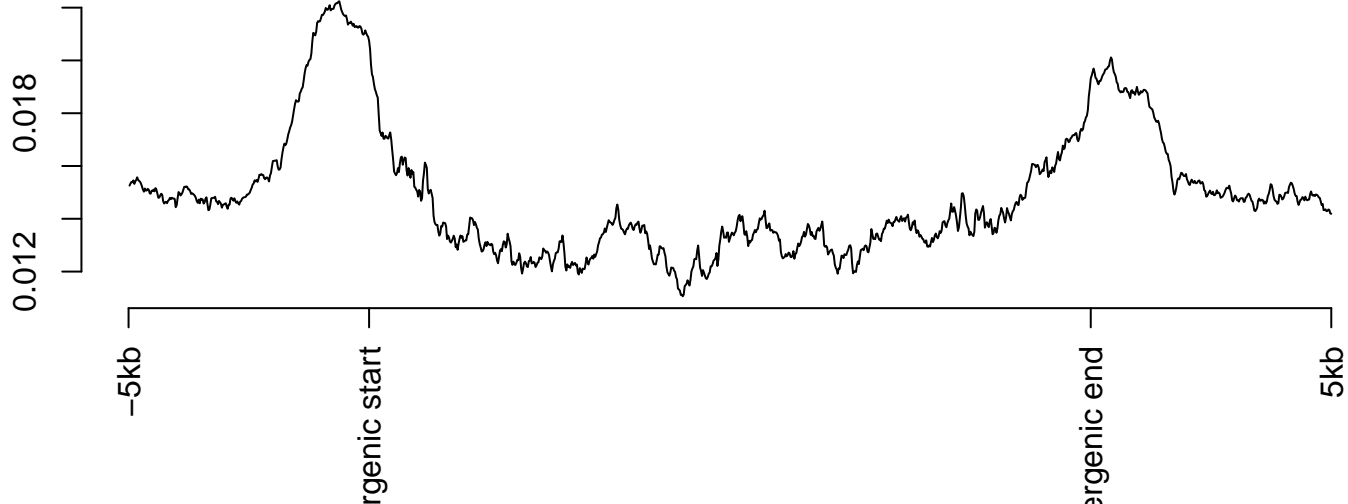
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.589 scCLTdegenNuc458



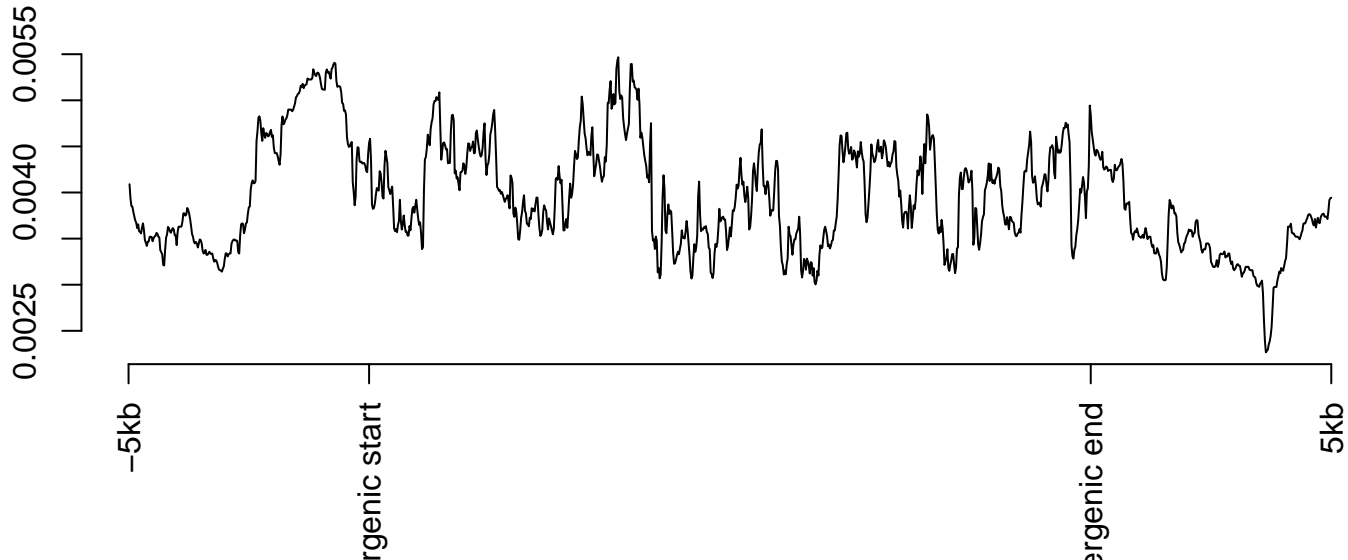
MouseNeuronSlice PositiveSingle E.589 scCLTdegenNuc459

Average coverage per intergenic region



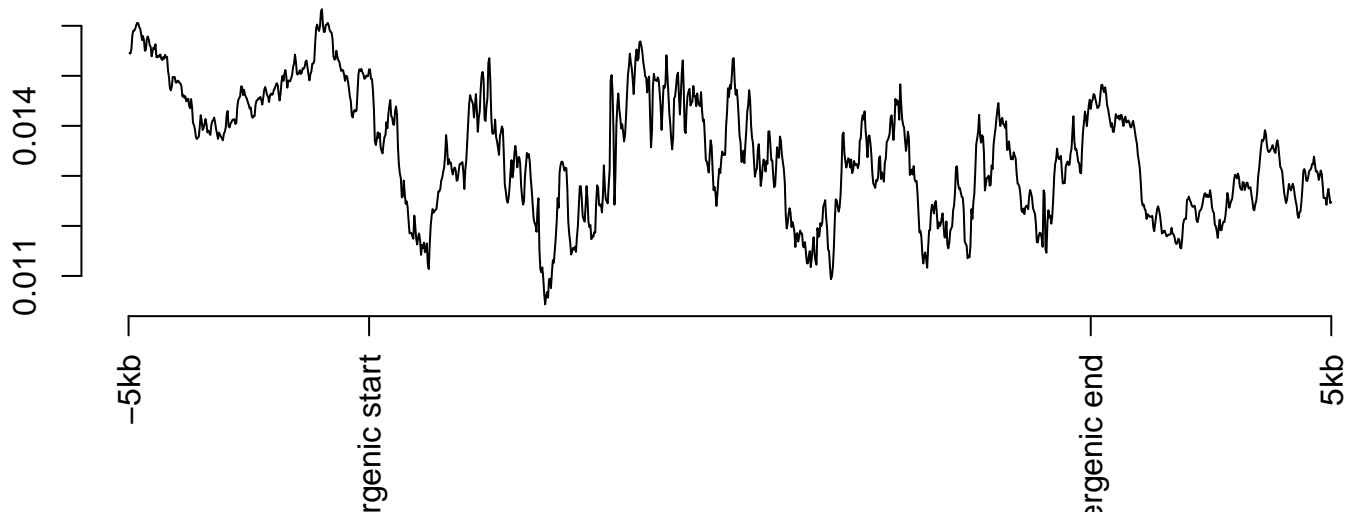
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.589 scCLTdegenNuc460



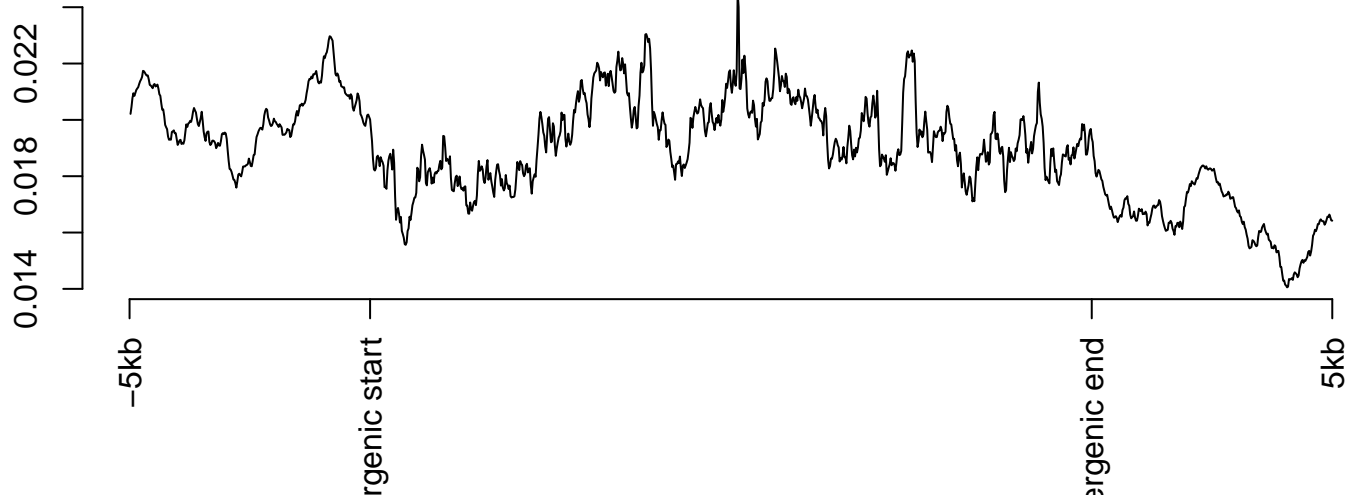
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.591 scCLTdegenNuc465



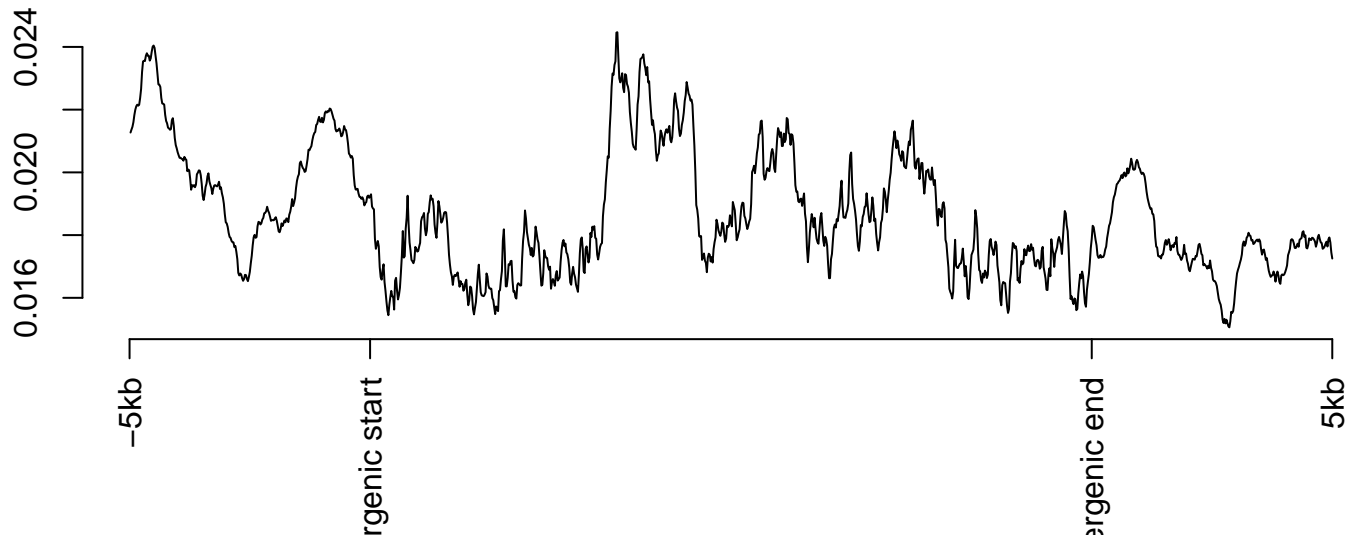
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.591 scCLTdegenNuc466



Average coverage per intergenic region

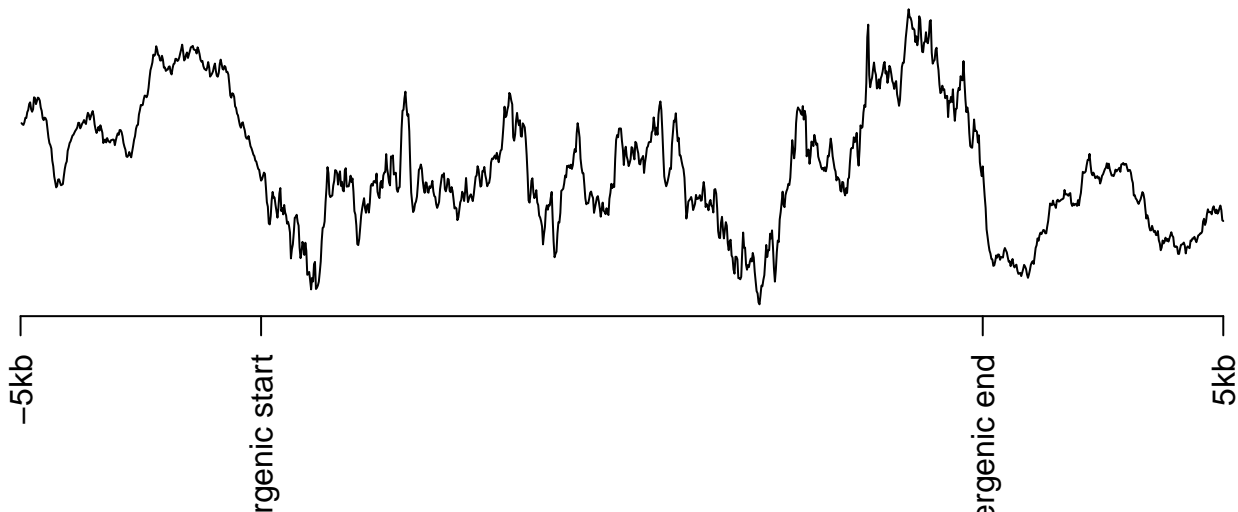
MouseNeuronSlice PositiveSingle E.591 scCLTdegenNuc467



Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.591 scCLTdegenNuc468

0.040
0.034



Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.597 scCLTdegenNuc469

0.014 0.017 0.020

-5kb

intergenic start

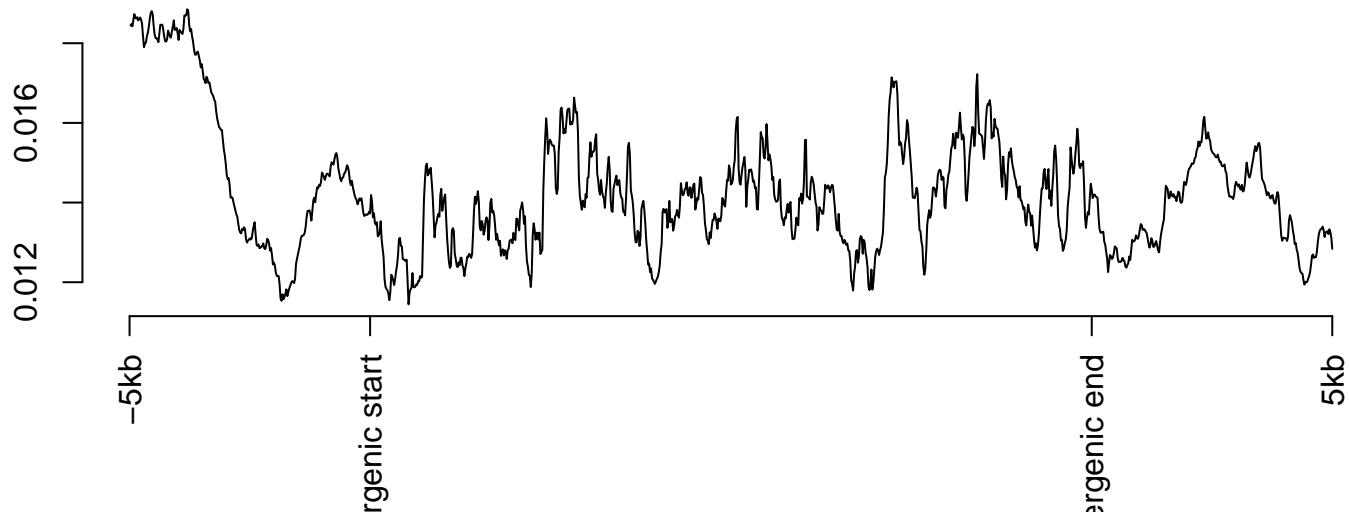
intergenic end

5kb



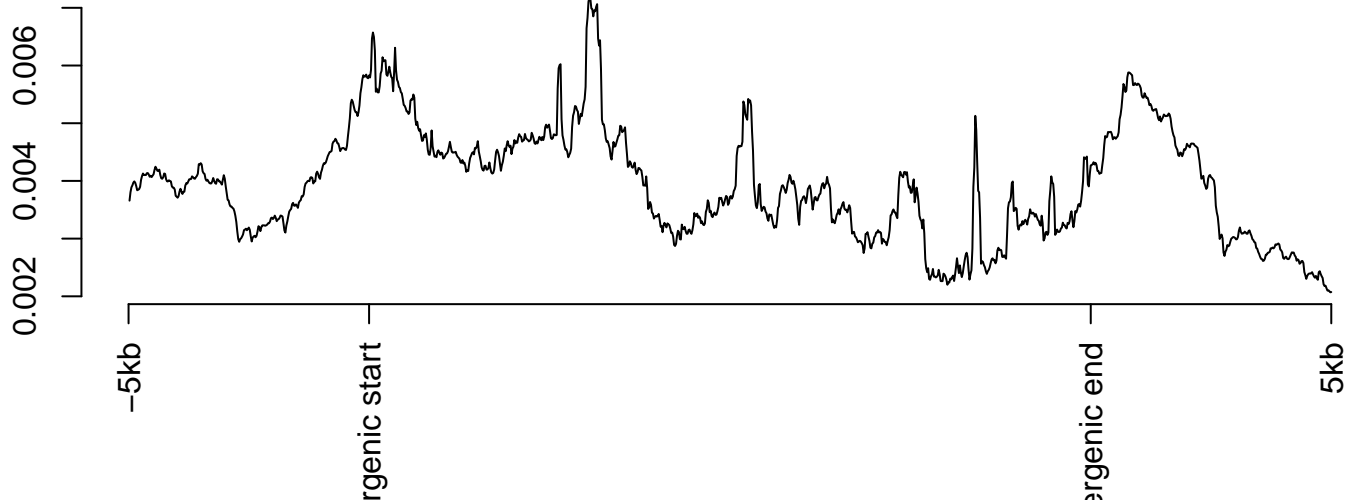
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.619 scCLTdegenNuc495



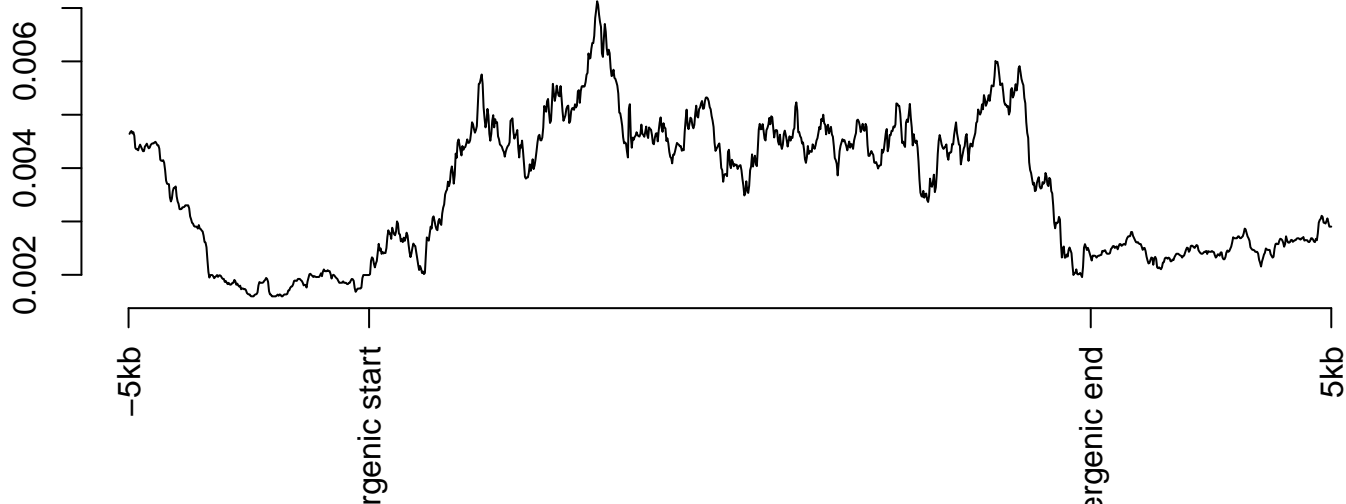
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.622 scCLTdegenNuc515



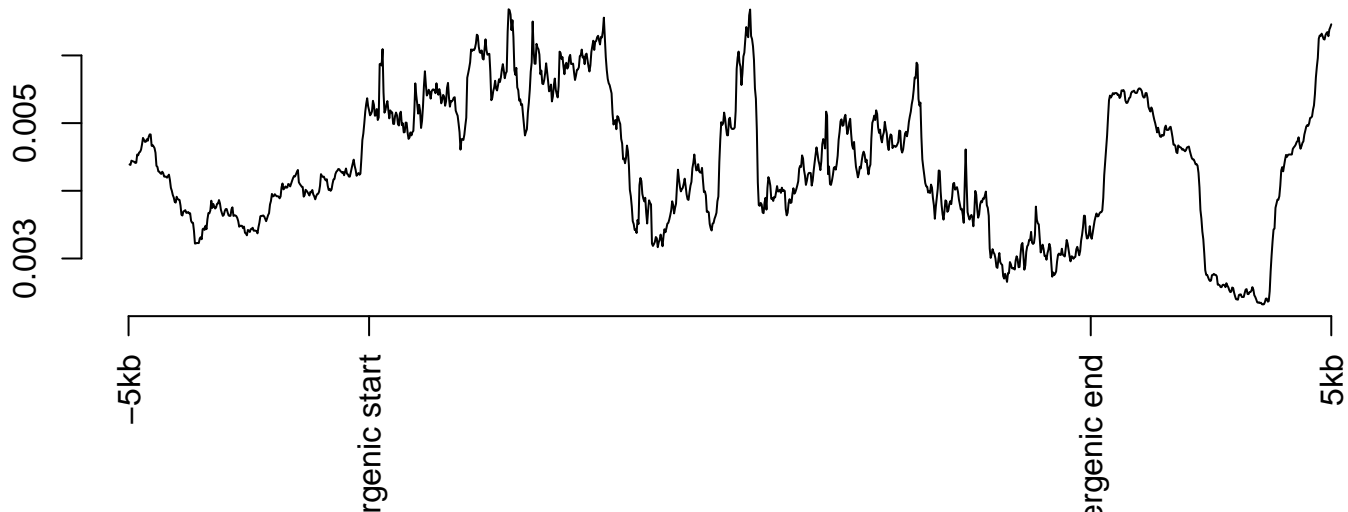
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.622 scCLTdegenNuc516



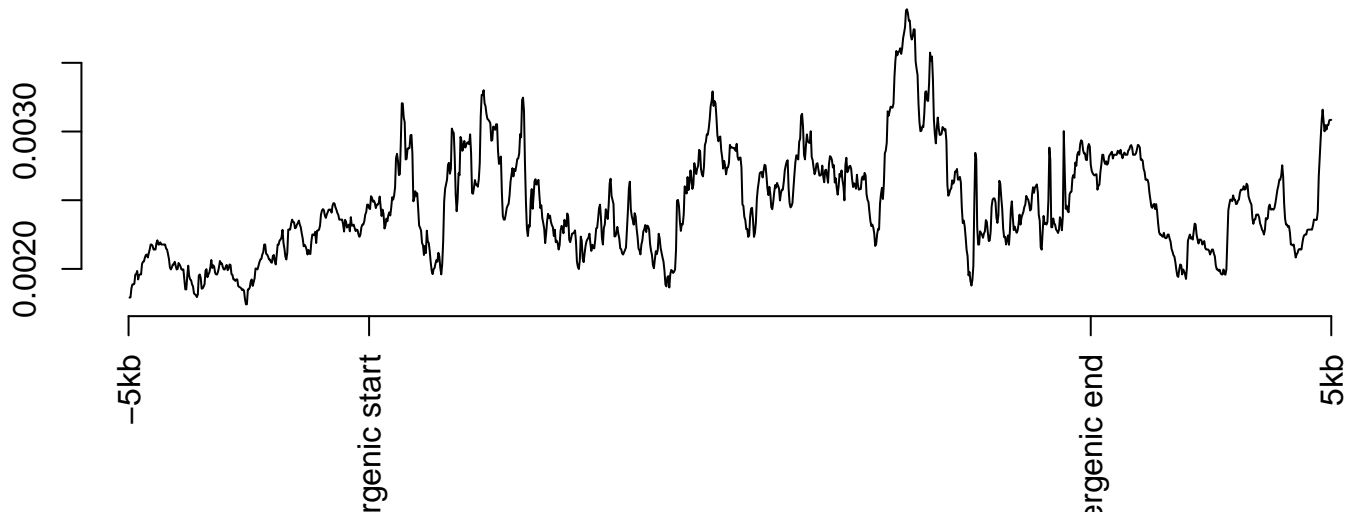
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.622 scCLTdegenNuc517



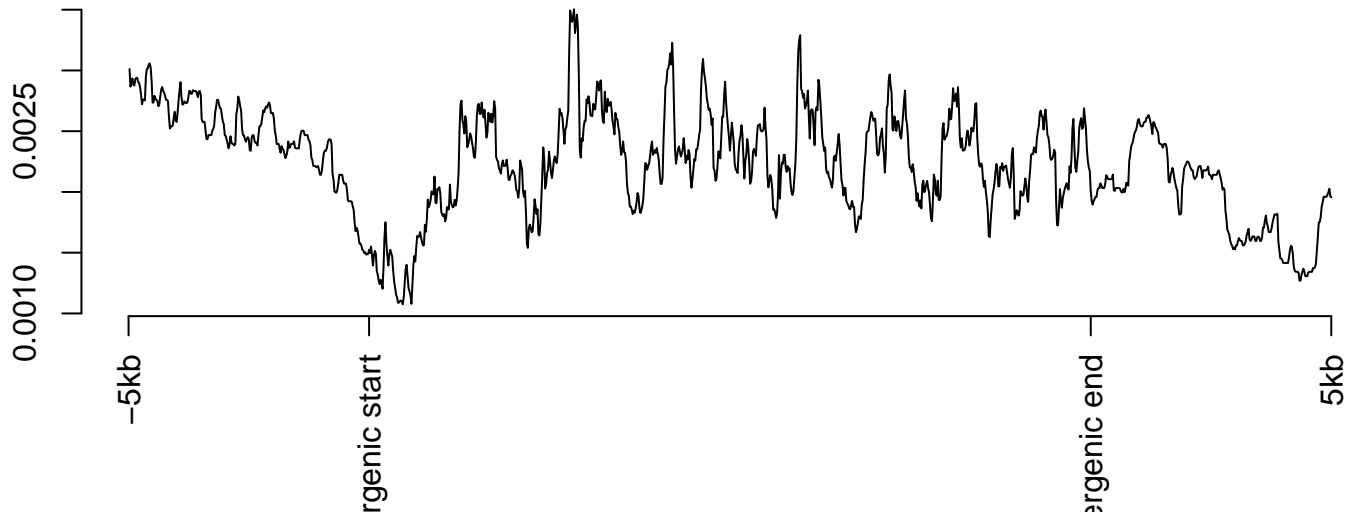
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.622 scCLTdegenNuc518



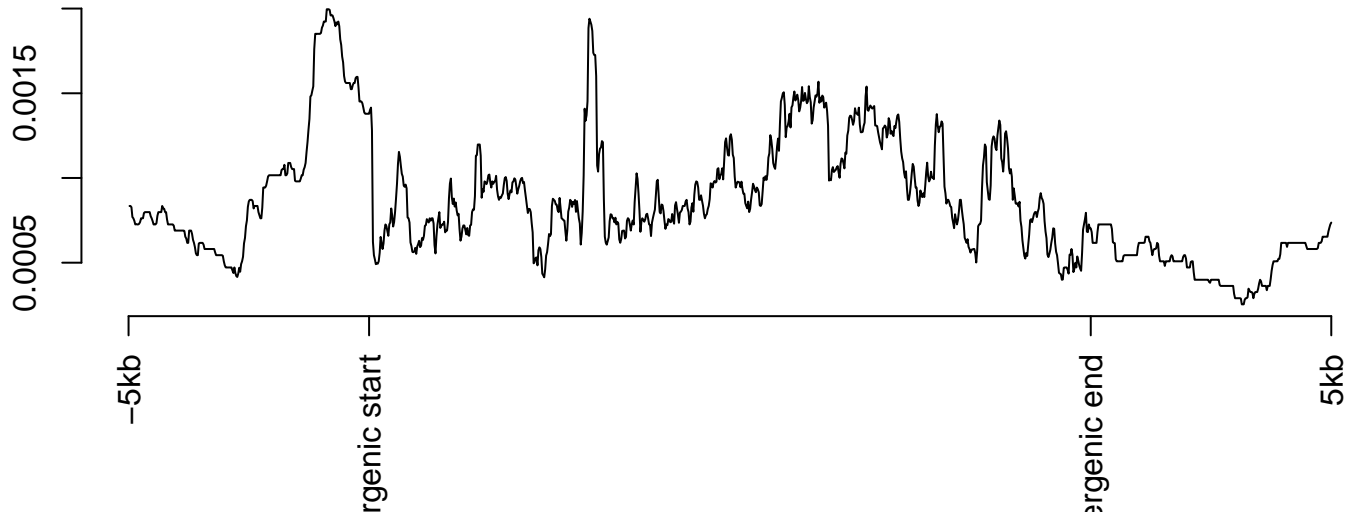
MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc609b

Average coverage per intergenic region



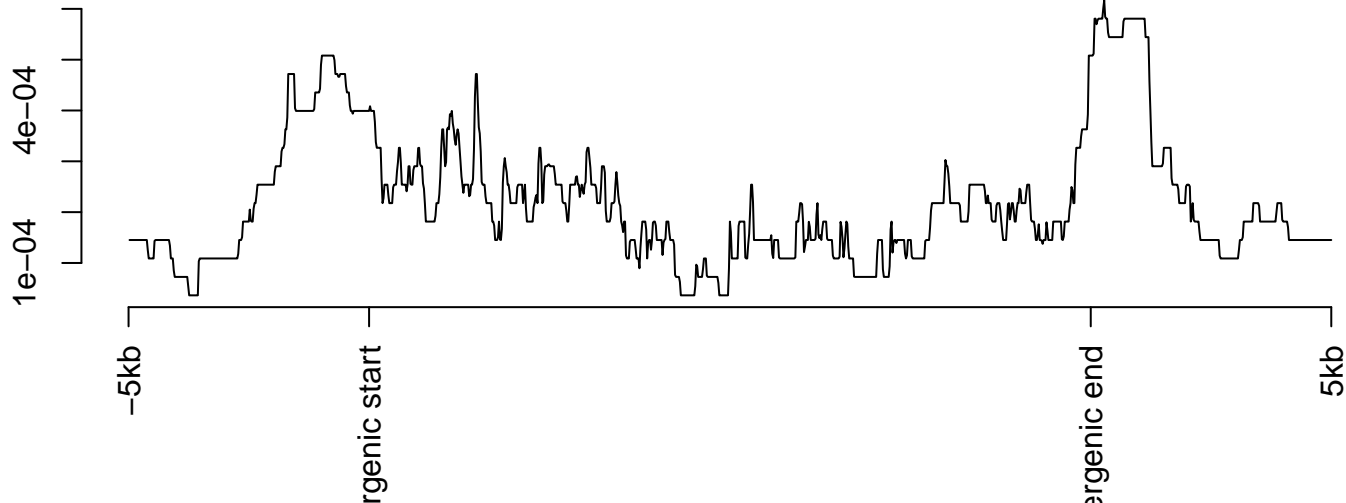
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc610b



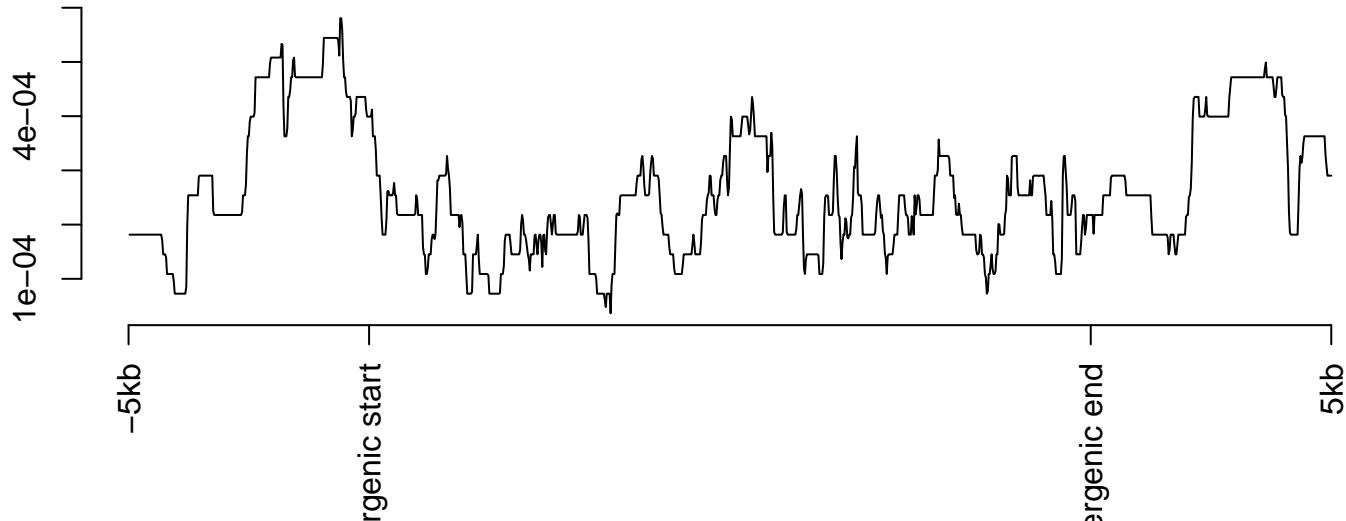
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc612b



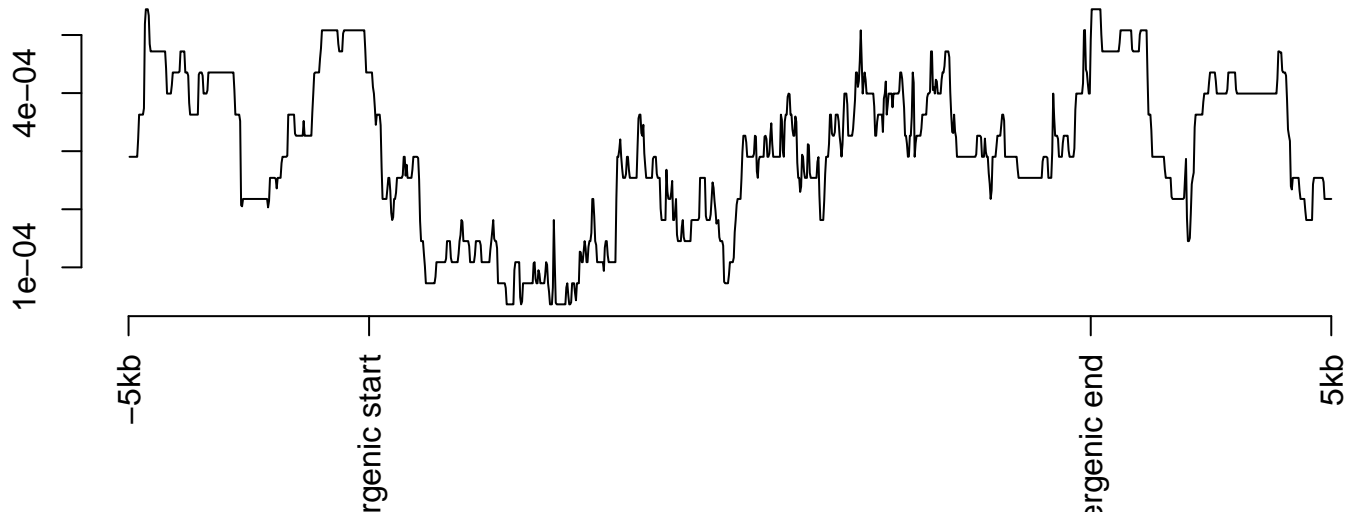
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc613b



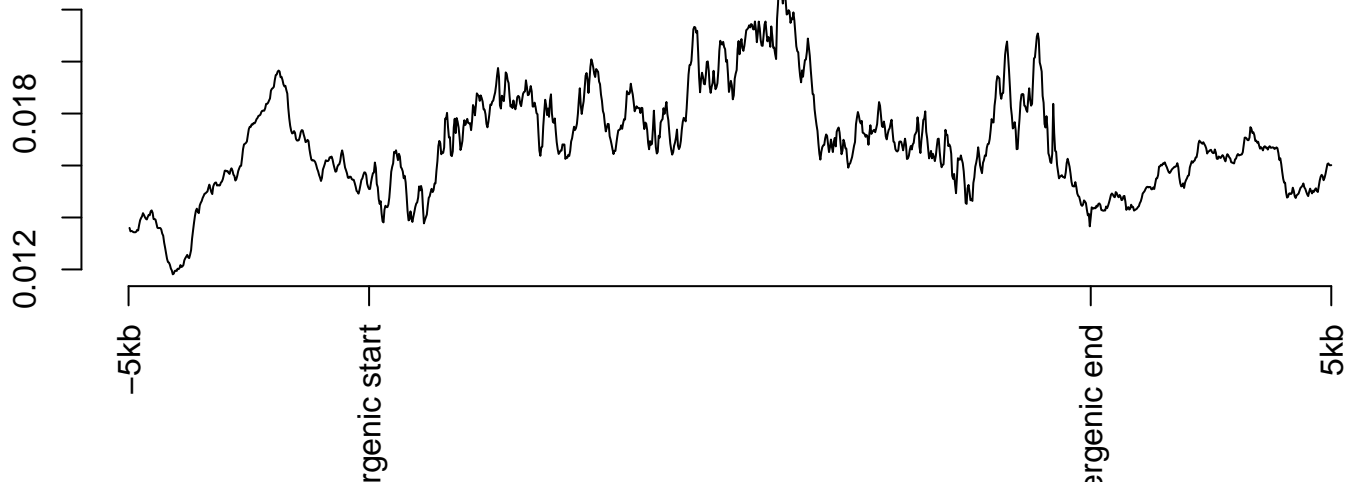
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc614b



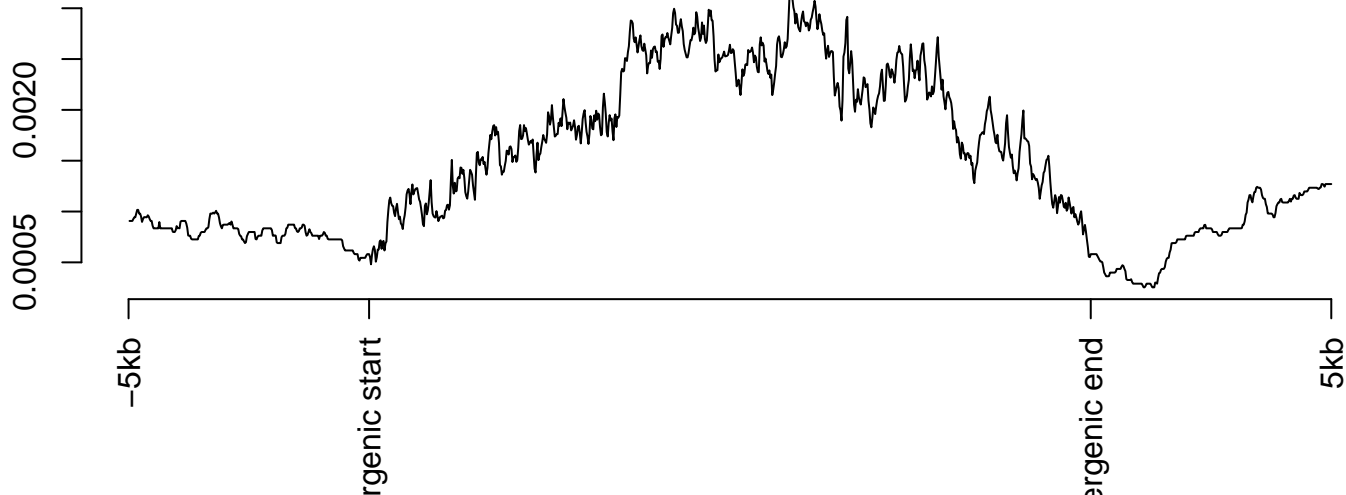
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc615b



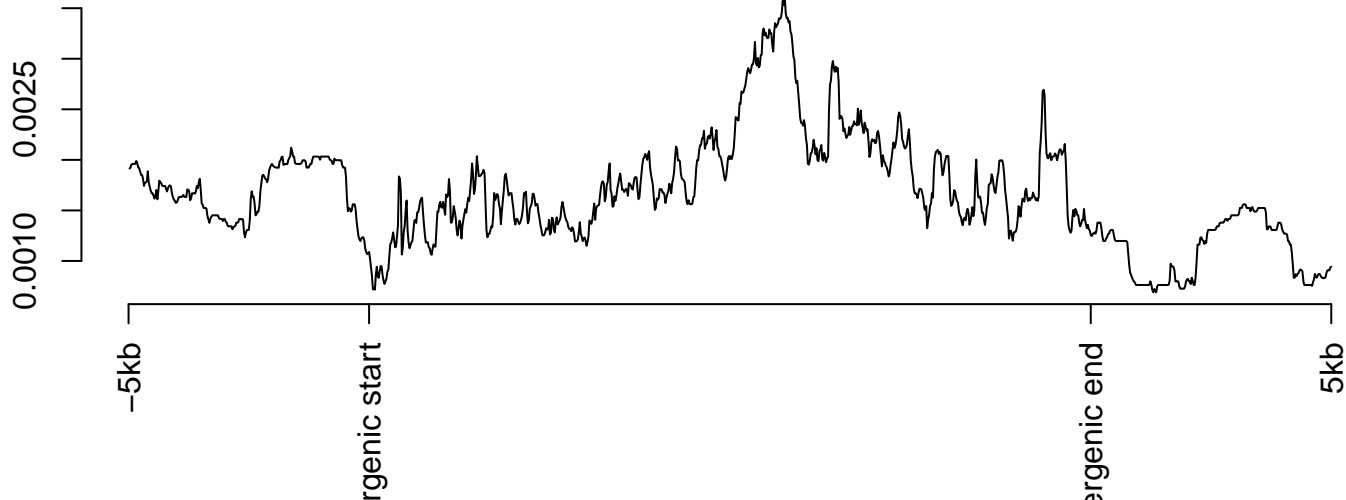
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc616b



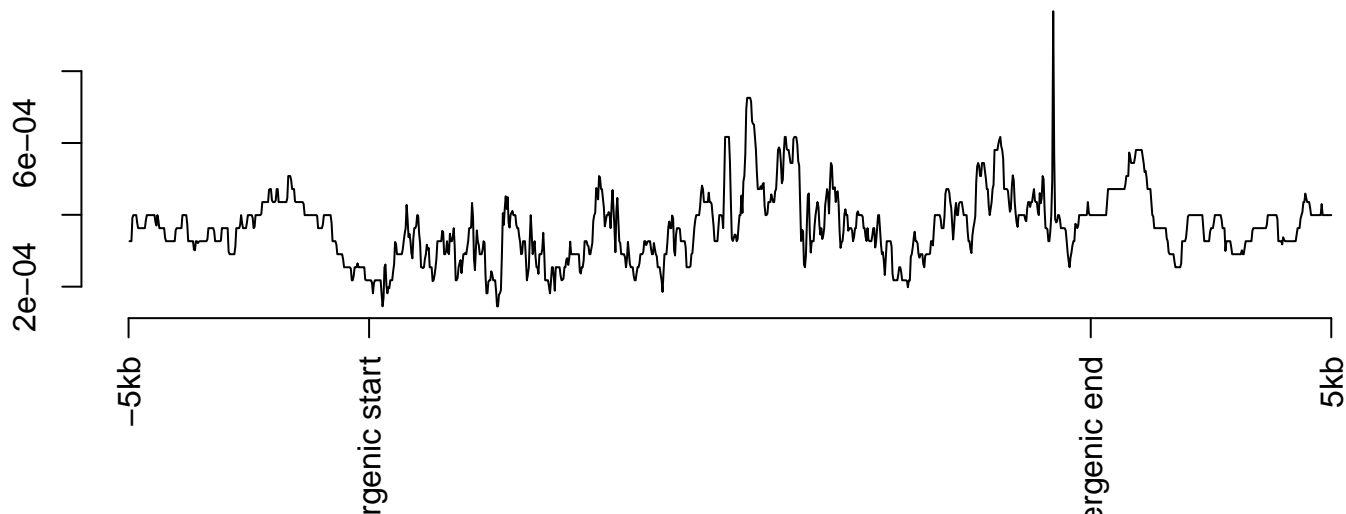
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc617b



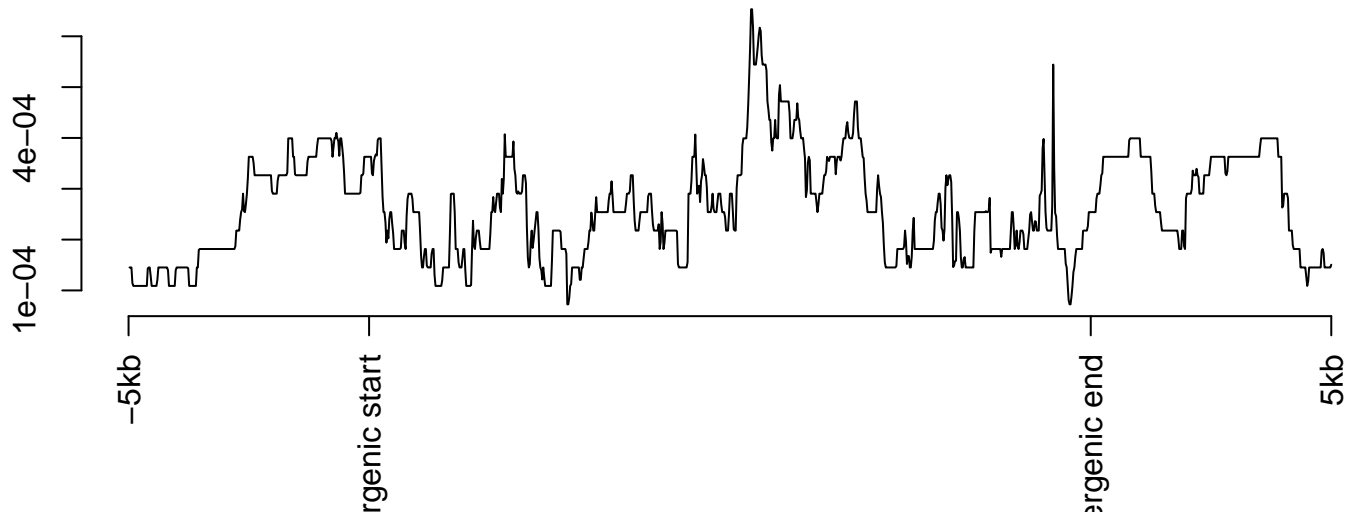
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc618b



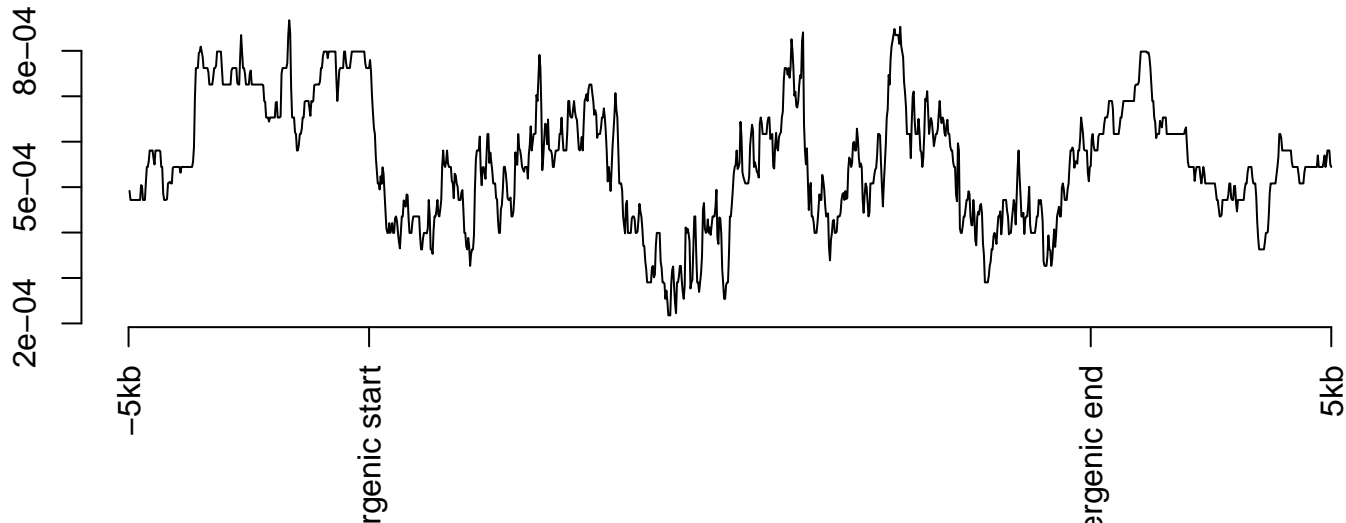
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.642 scCLTdegenNuc619b



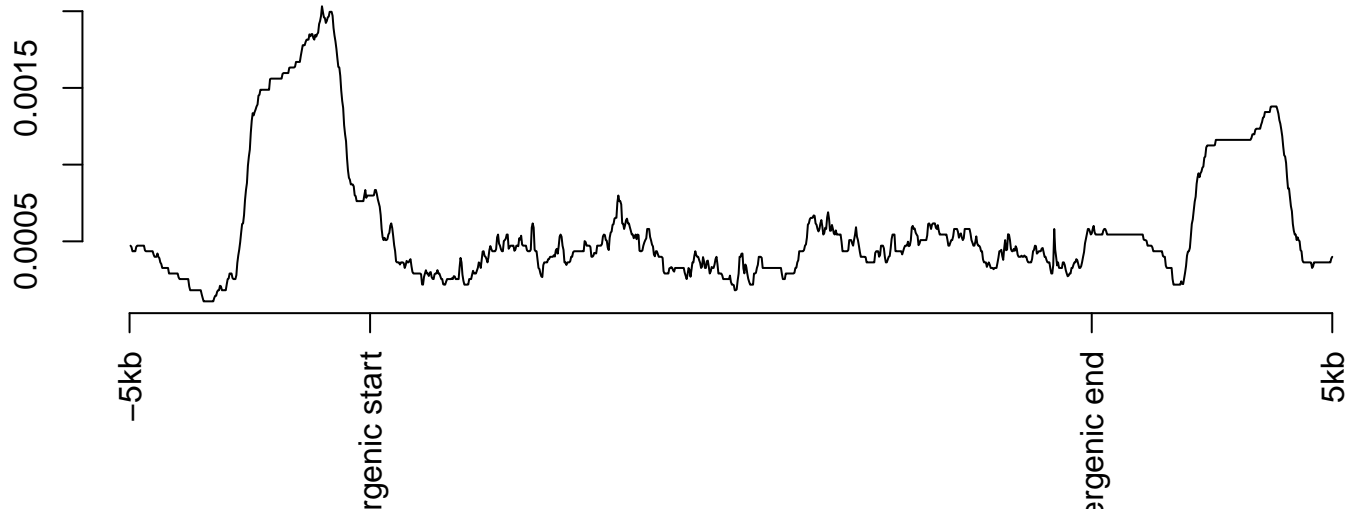
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.643 scCLTdegenNuc620



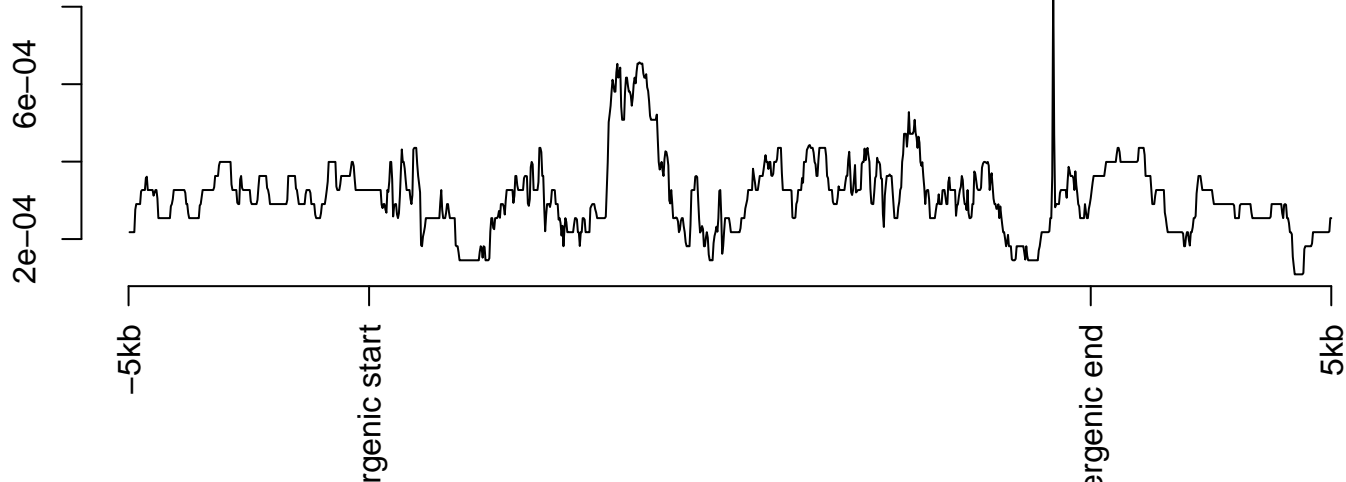
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.670 scCLTdegenNuc840



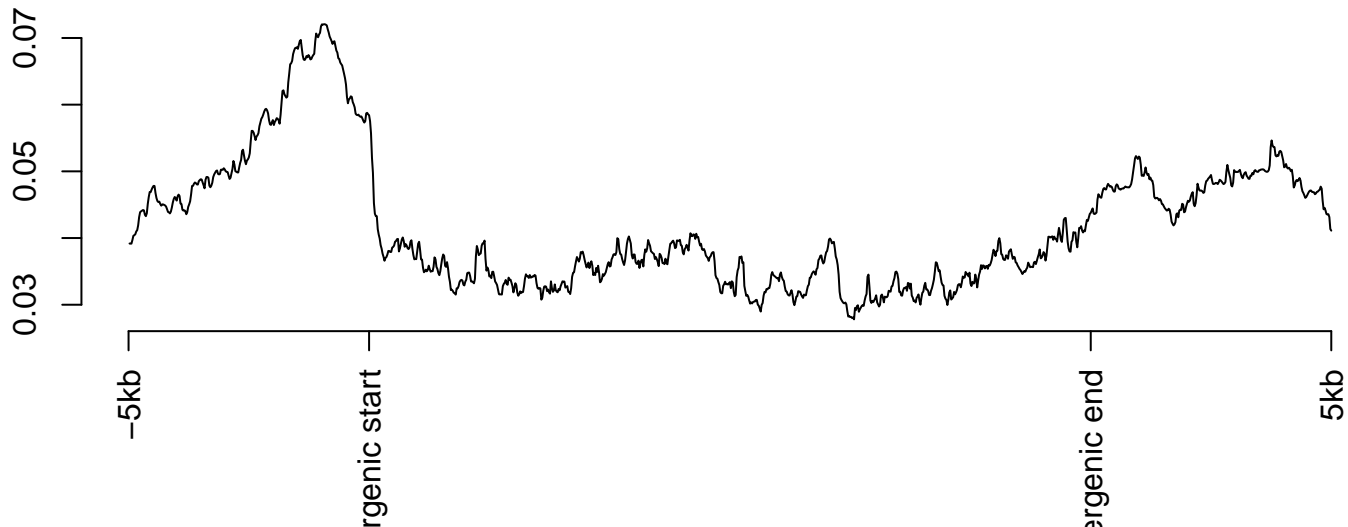
Average coverage per intergenic region

MouseNeuronSlice PositiveSingle E.670 scCLTdegenNuc843



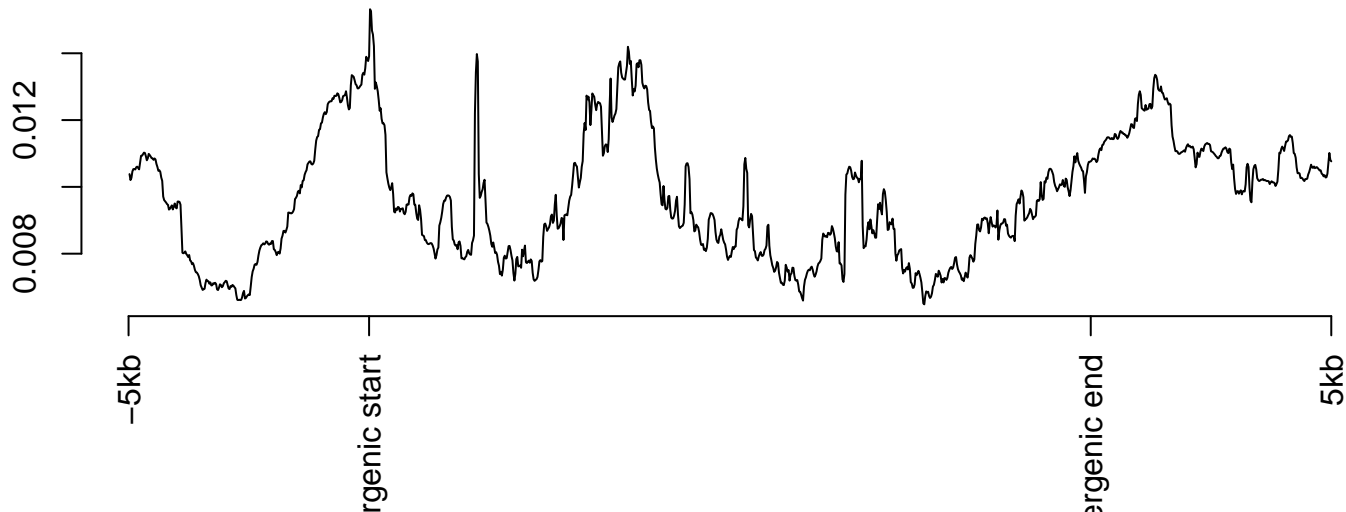
Average coverage per intergenic region

MouseNeuronSlice PositivePooled E.584 scCLTdegenNuc413



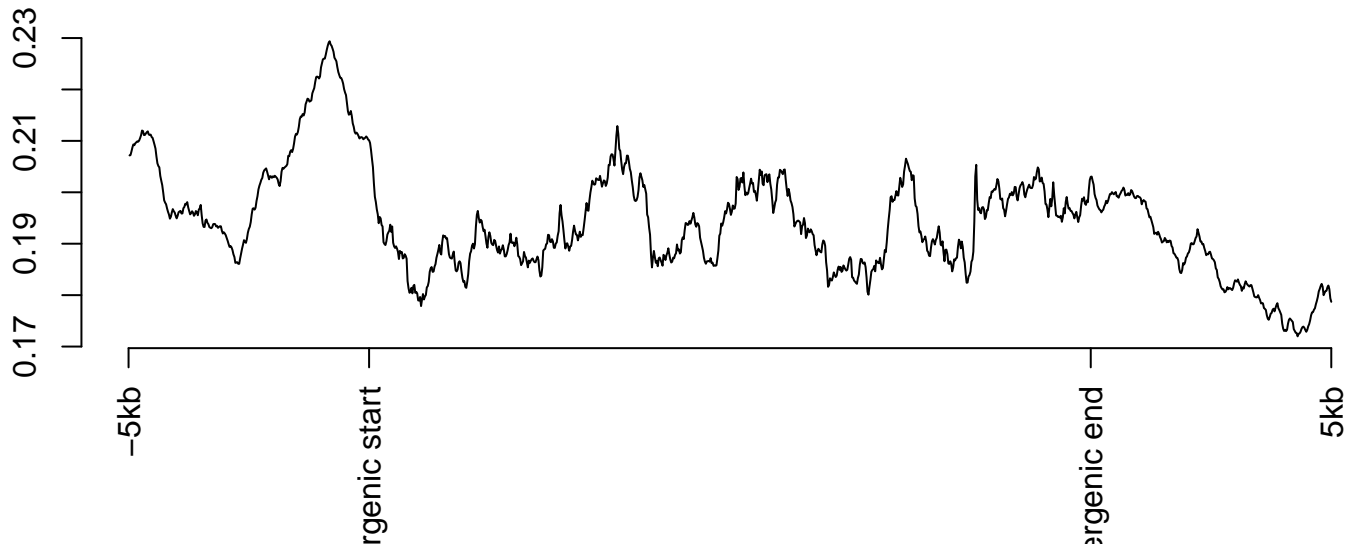
Average coverage per intergenic region

MouseNeuronSlice PositivePooled E.589 scCLTdegenNuc462



Average coverage per intergenic region

MouseNeuronSlice PositiveMerged E.– MouseNeuronSlicePositiveMerged



Average coverage per intergenic region

MouseNeuronSlice PositiveAll E.– MouseNeuronSlicePositiveAll

