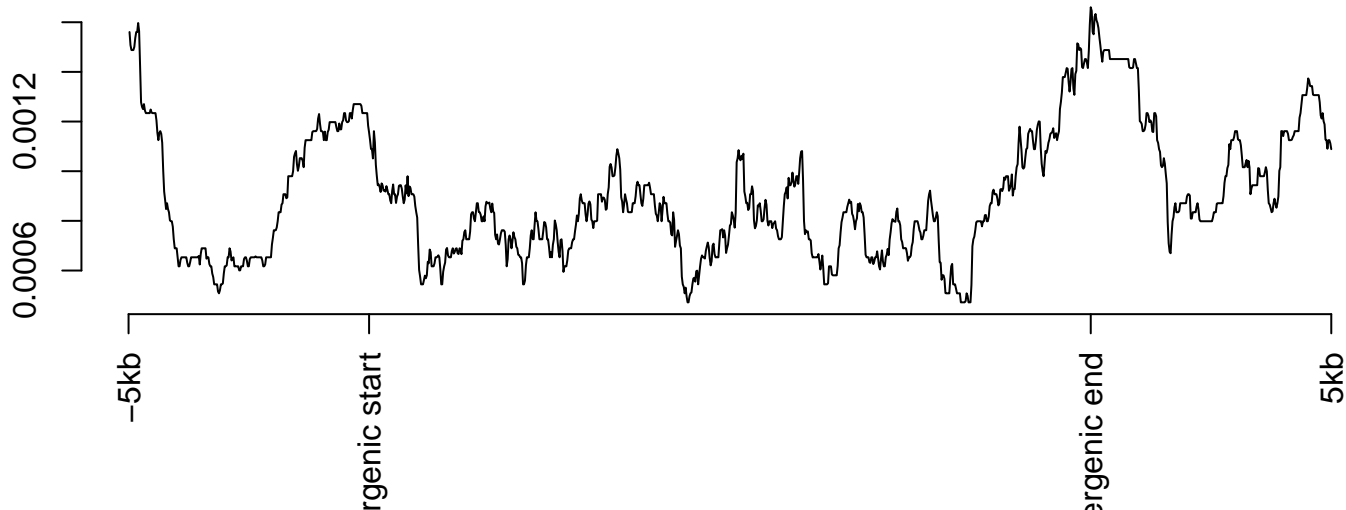


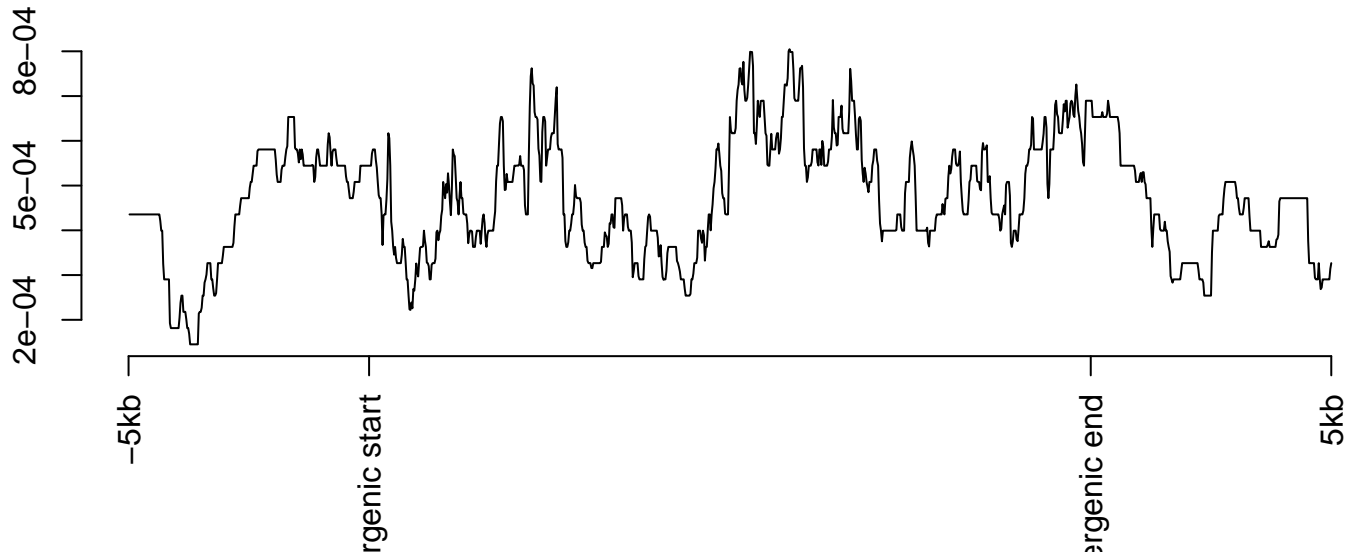
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc603



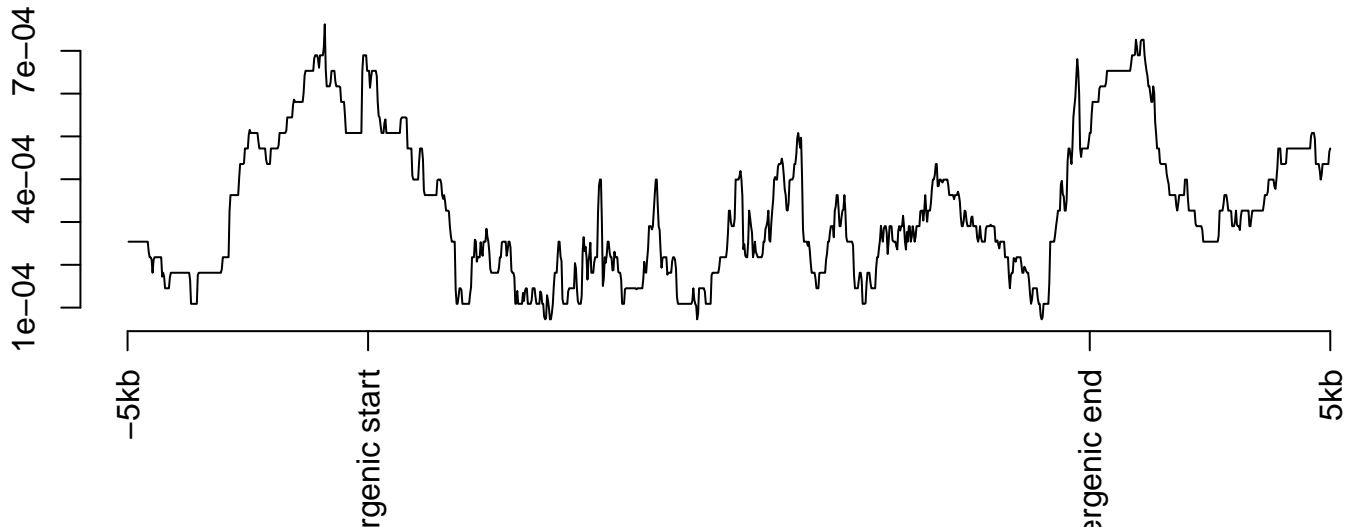
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc604



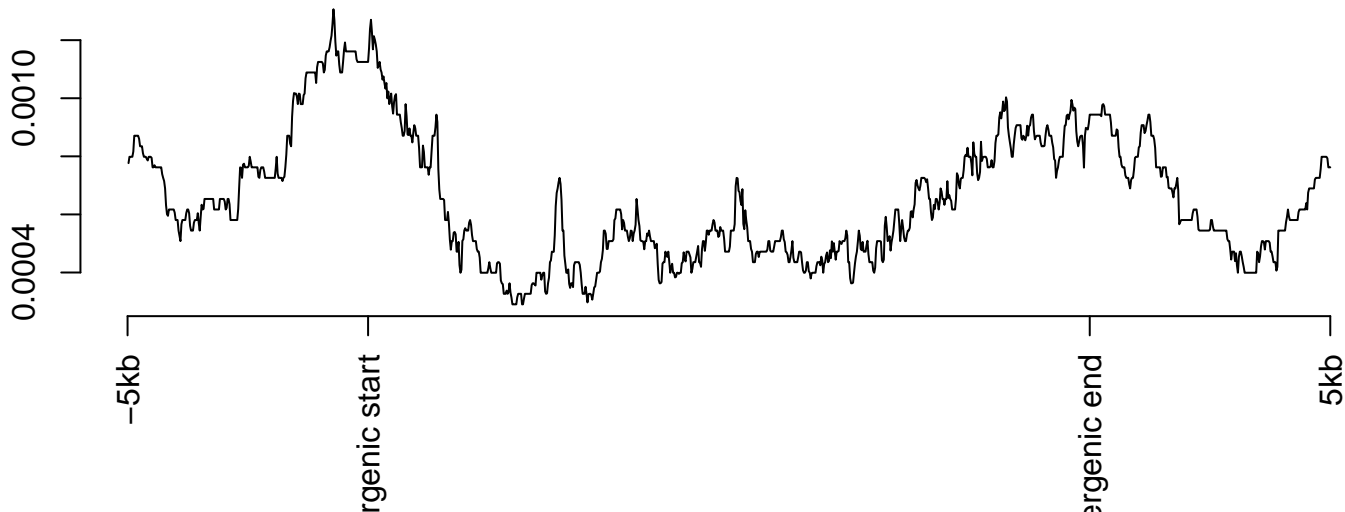
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc605



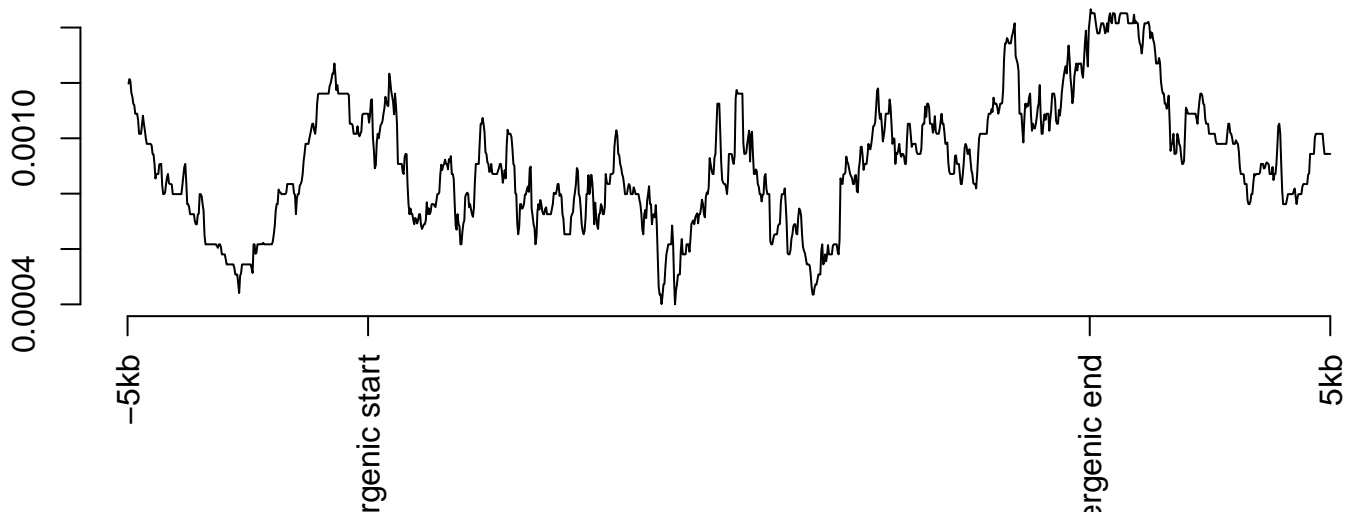
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc606



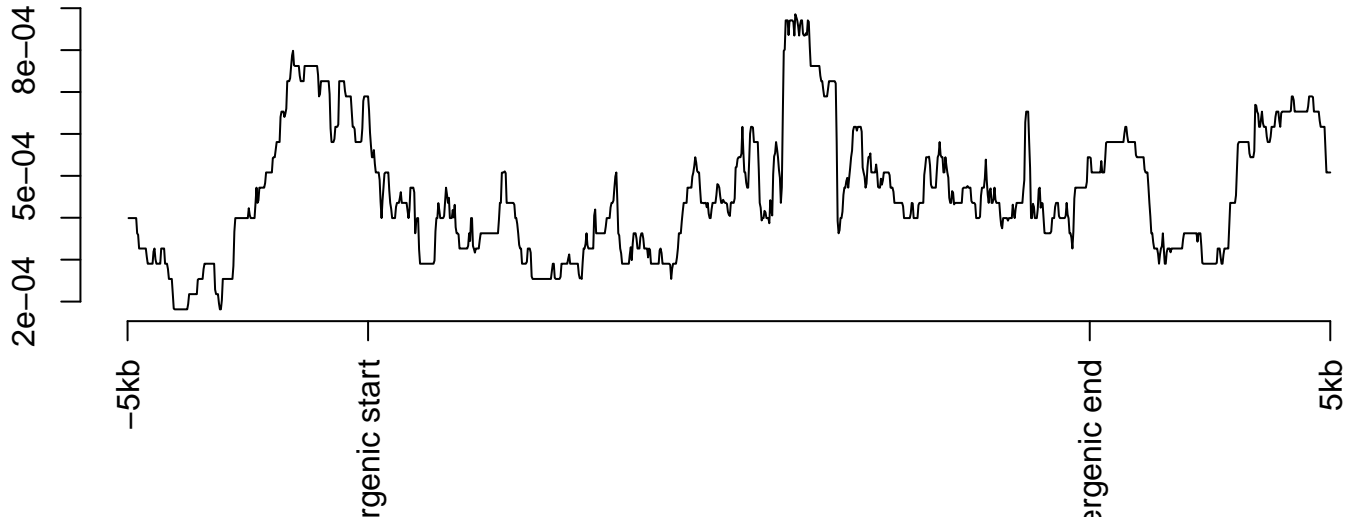
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.638 scCLTdegenNuc607



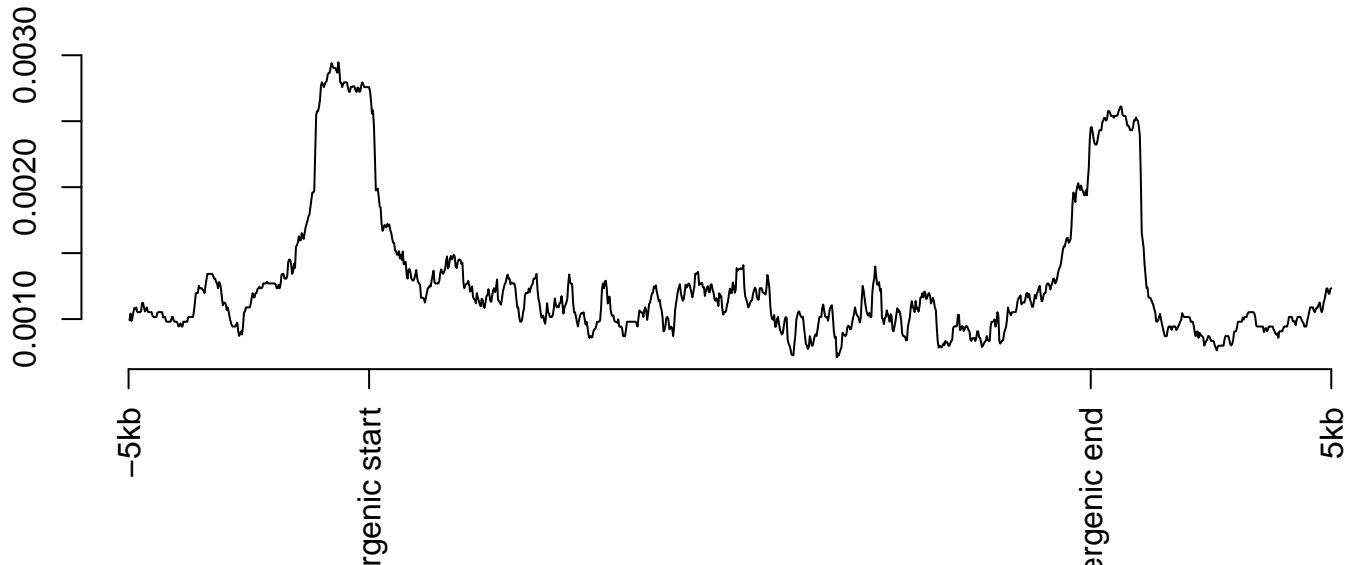
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.642 scCLTdegenNuc608b



Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc621



Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc622

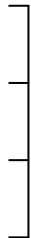
0.006
0.004

-5kb

intergenic start

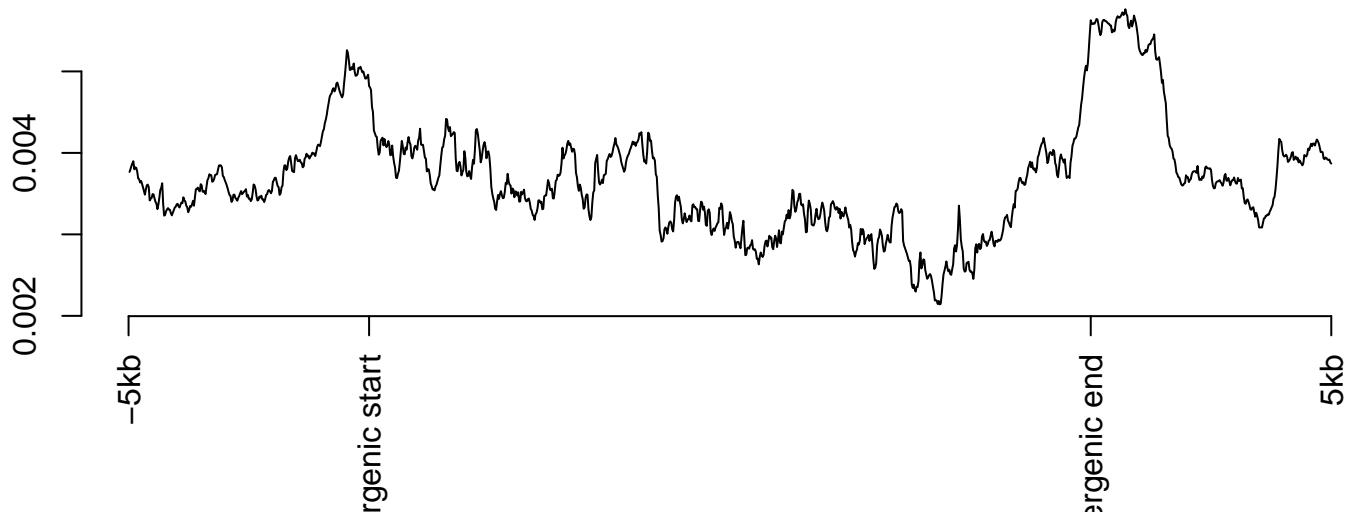
intergenic end

5kb



Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc623



Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc624

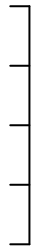
0.008 0.012 0.016

-5kb

intergenic start

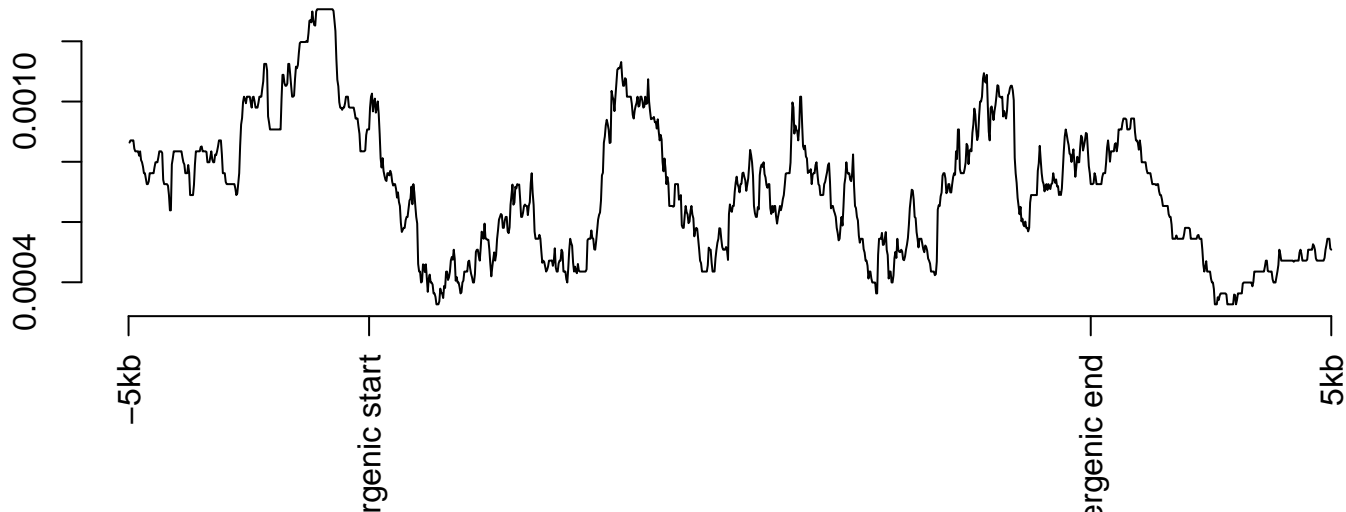
intergenic end

5kb



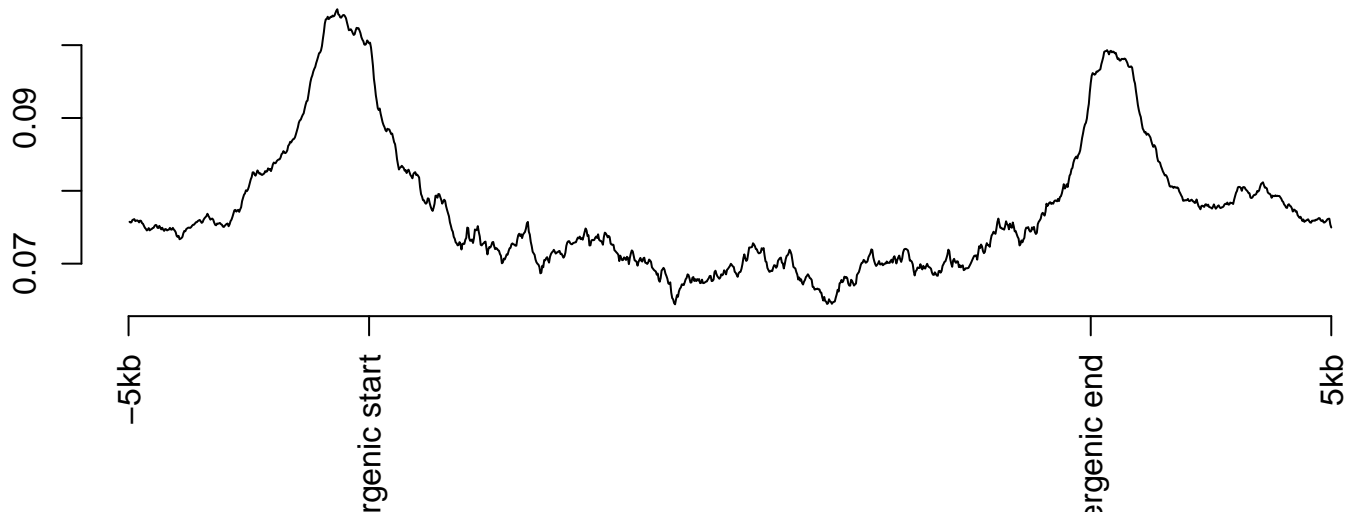
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc625



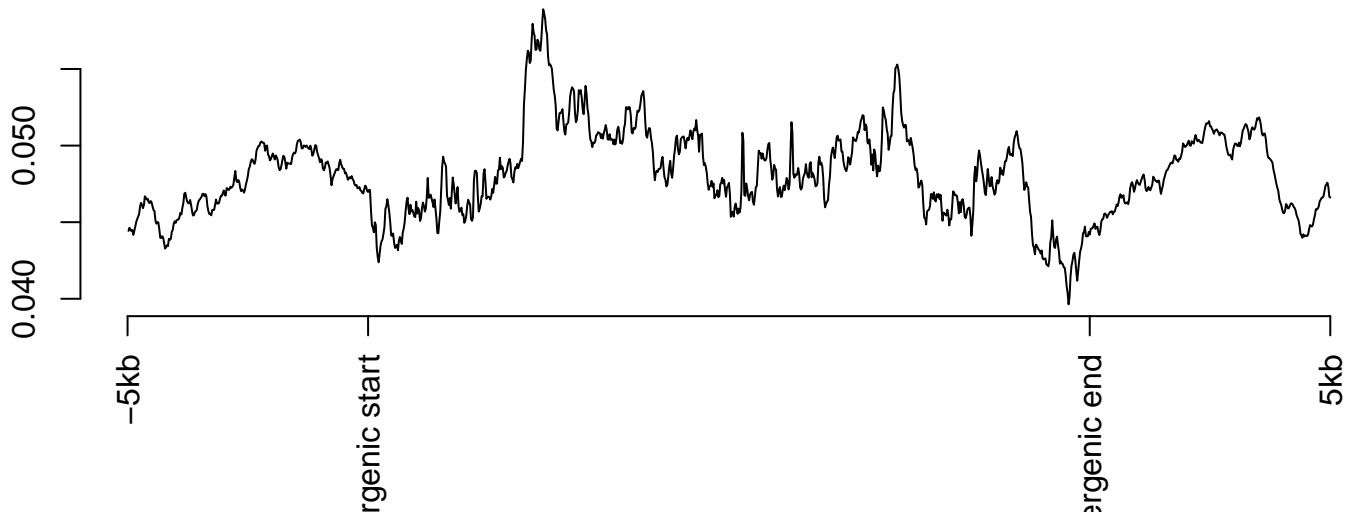
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc626



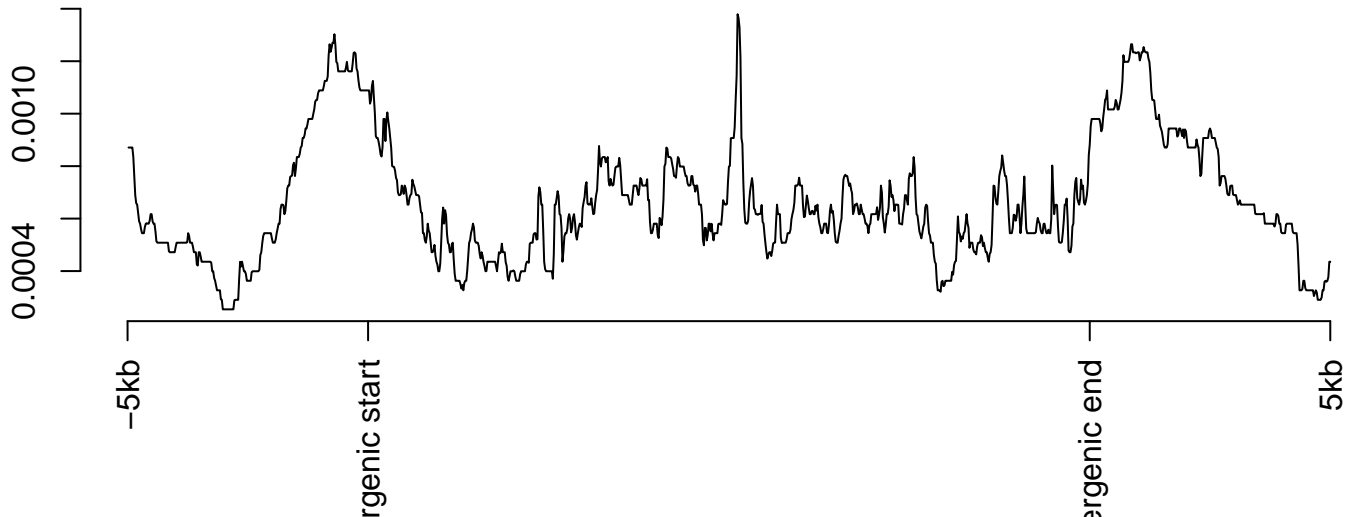
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.643 scCLTdegenNuc627



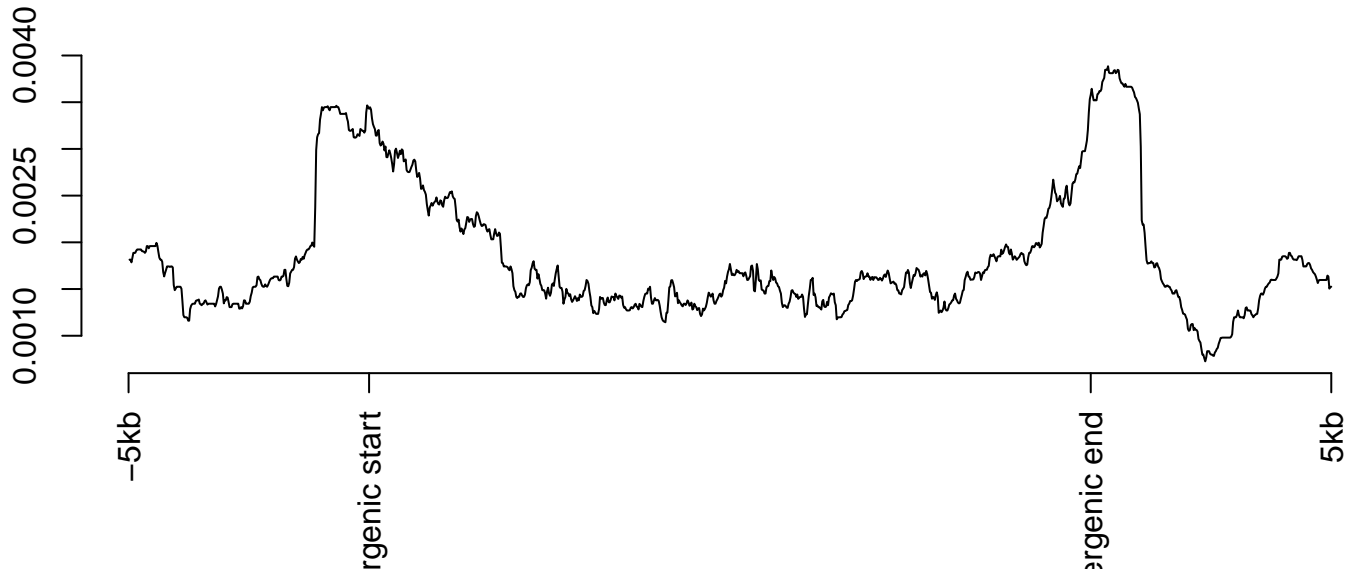
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.653 scCLTdegenNuc663



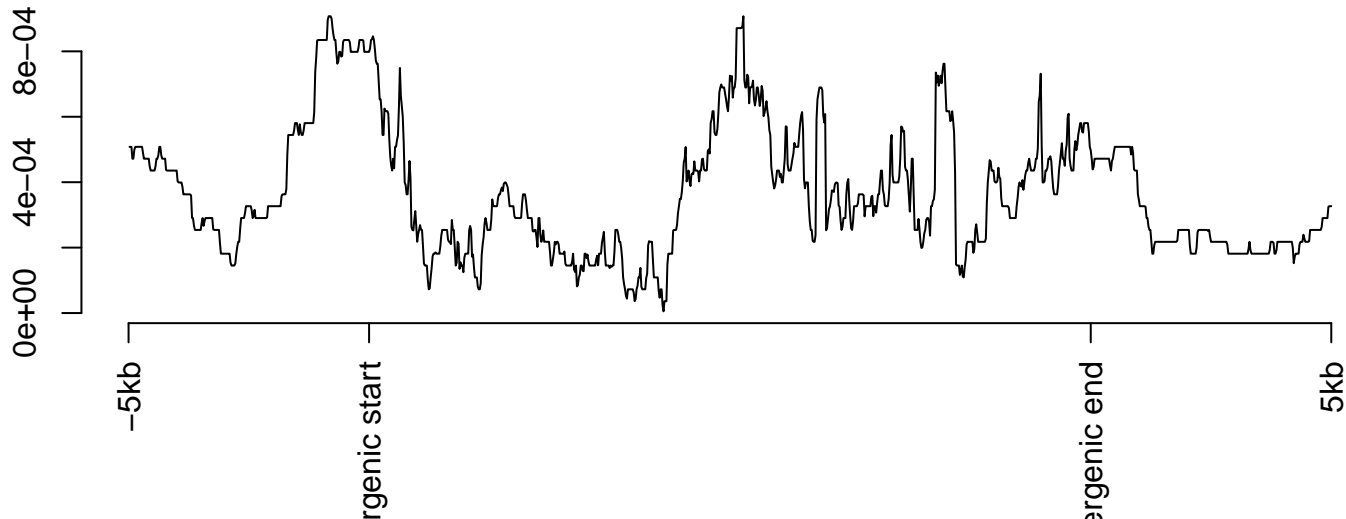
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.653 scCLTdegenNuc664



Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc697



Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc698

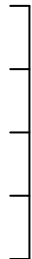
2e-04 6e-04 1e-03

-5kb

intergenic start

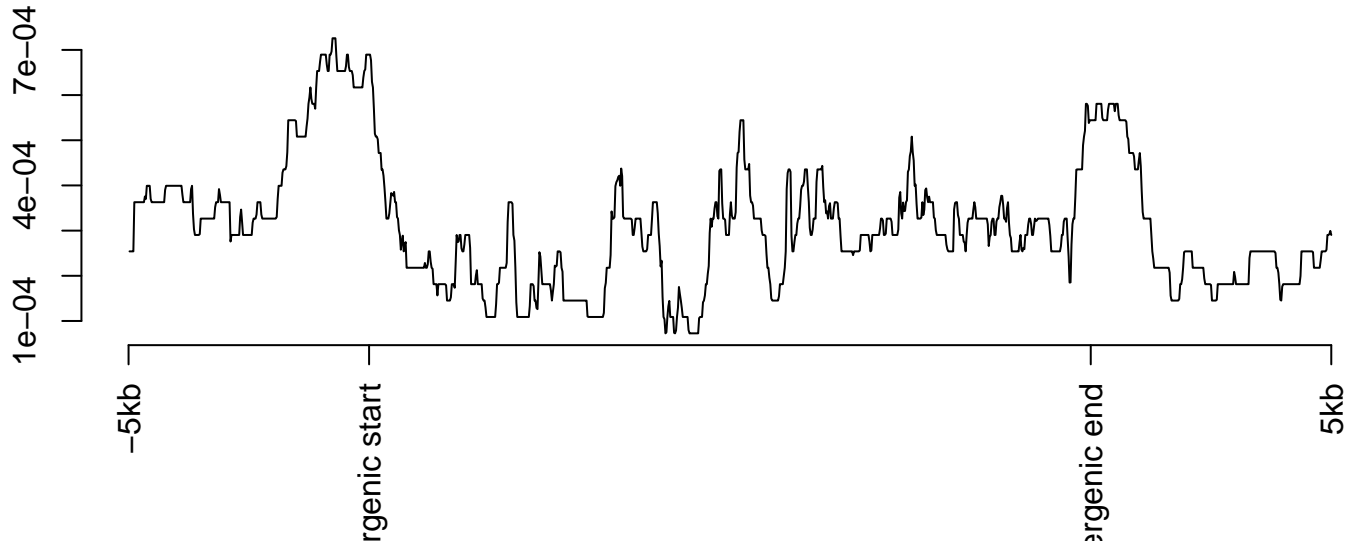
intergenic end

5kb



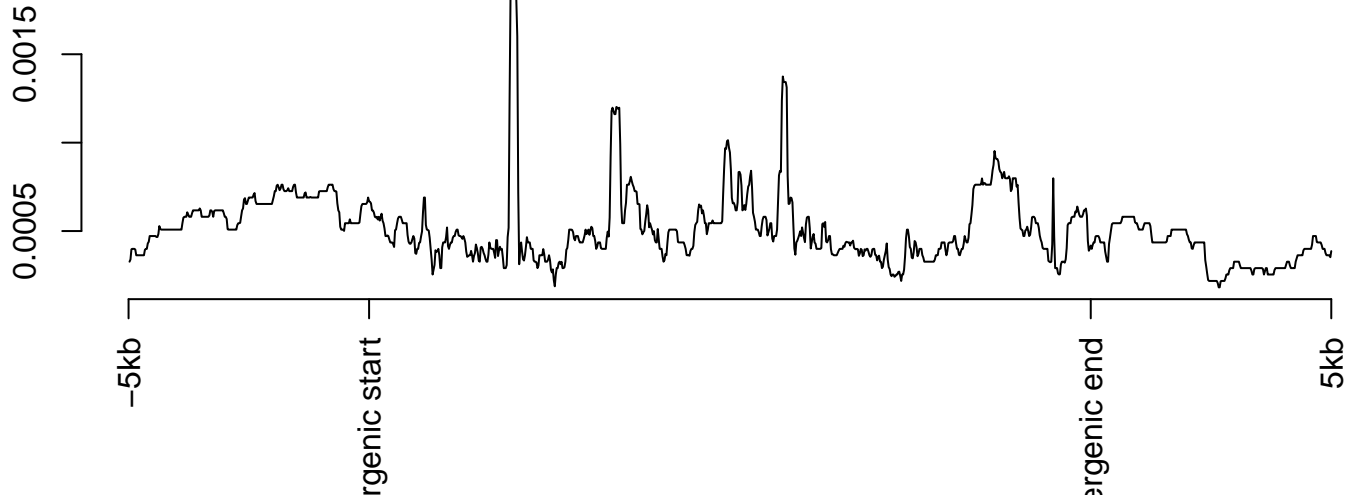
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.655 scCLTdegenNuc699



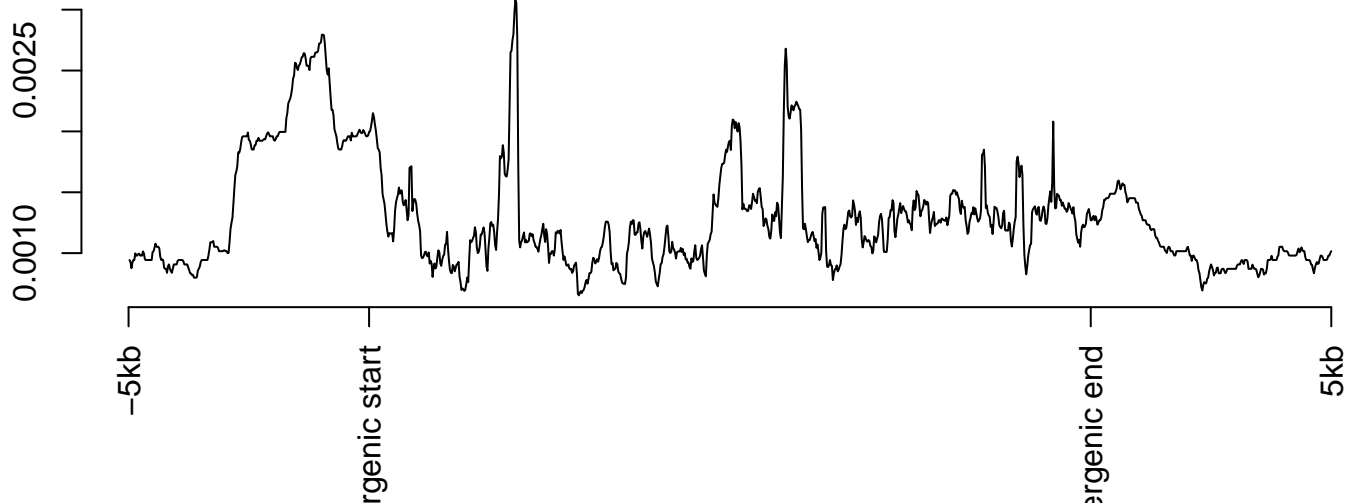
Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc776



Average coverage per intergenic region

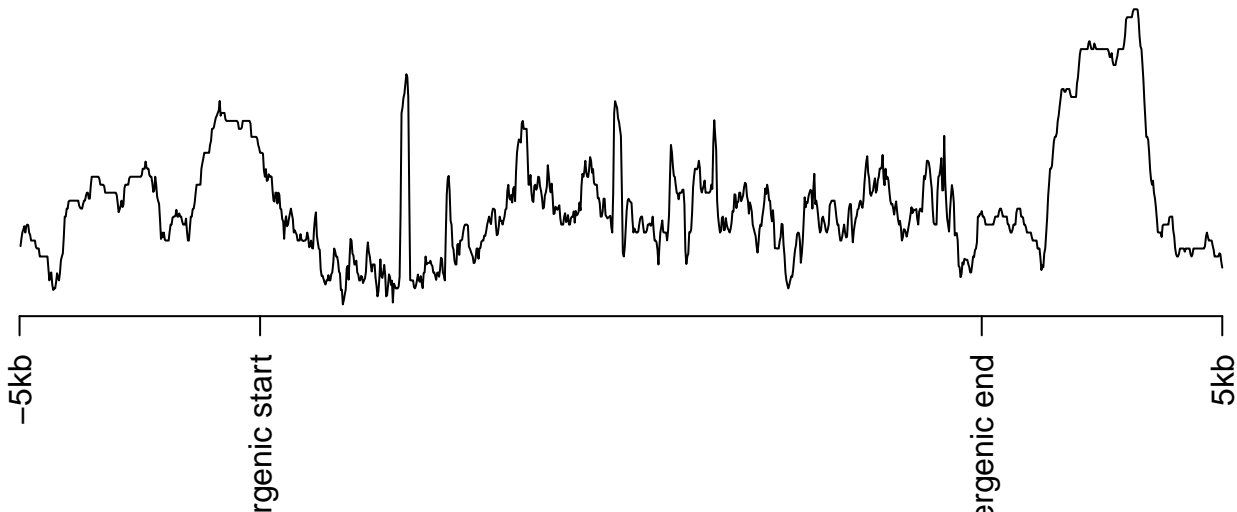
MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc777



Average coverage per intergenic region

MouseInterneuronSlice PositiveSingle E.666 scCLTdegenNuc778

0.0004 0.0010 0.0016



Average coverage per intergenic region

MouseInterneuronSlice PositiveMerged E.– MouseInterneuronSlicePositiveMerged

