

HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc441

Average coverage per intergenic region

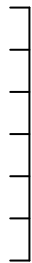
0.0005 0.0020 0.0035

-5kb

intergenic start

intergenic end

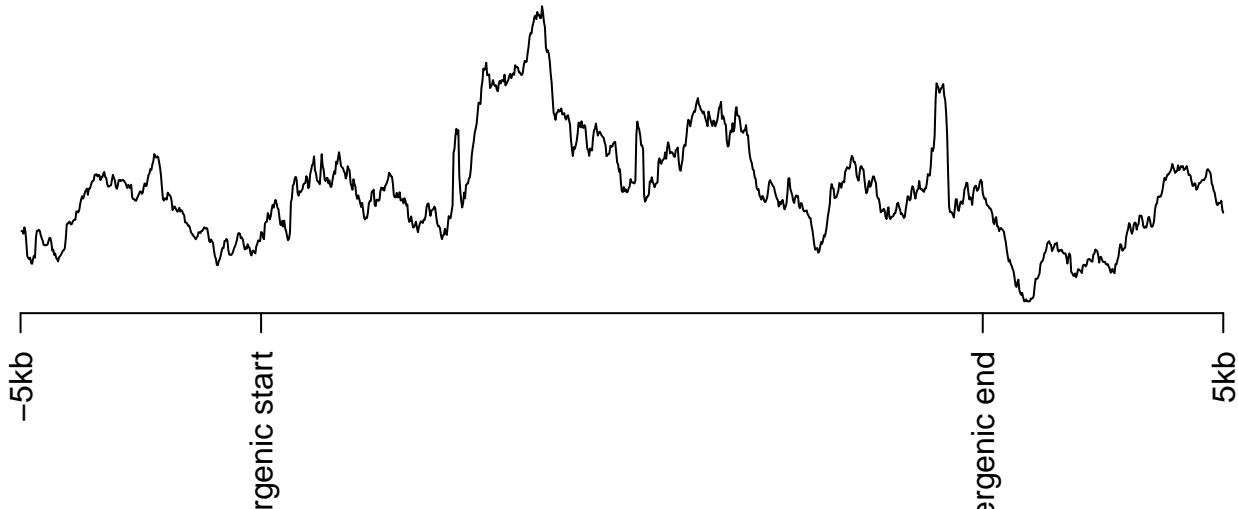
5kb



HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc442

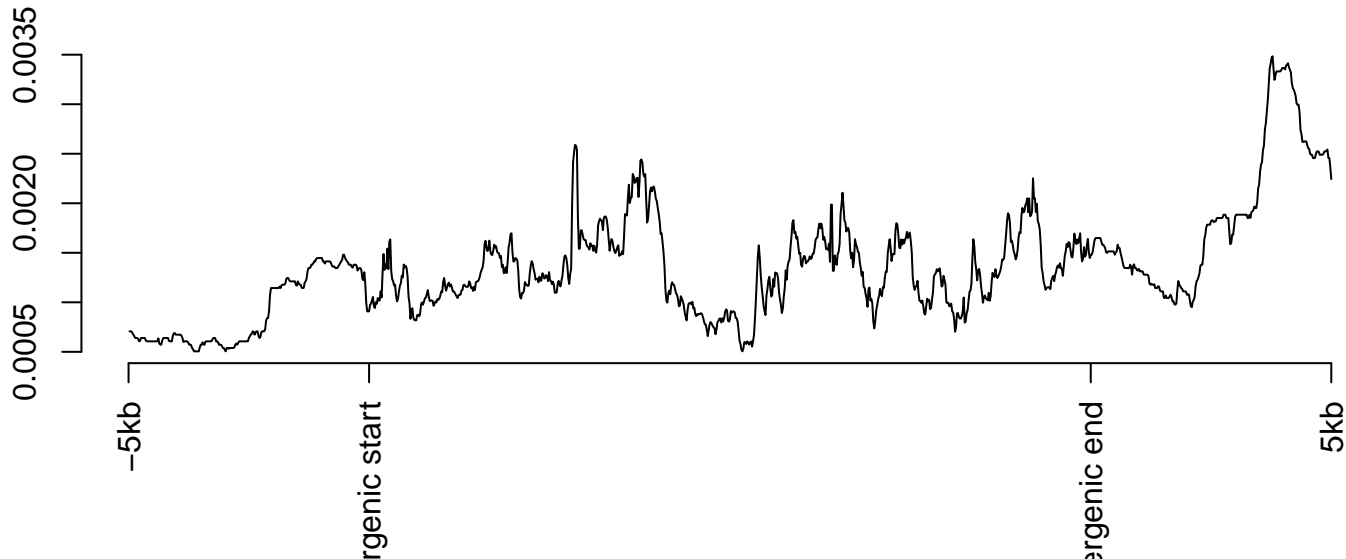
Average coverage per intergenic region

0.0025 0.0040 0.0055



Average coverage per intergenic region

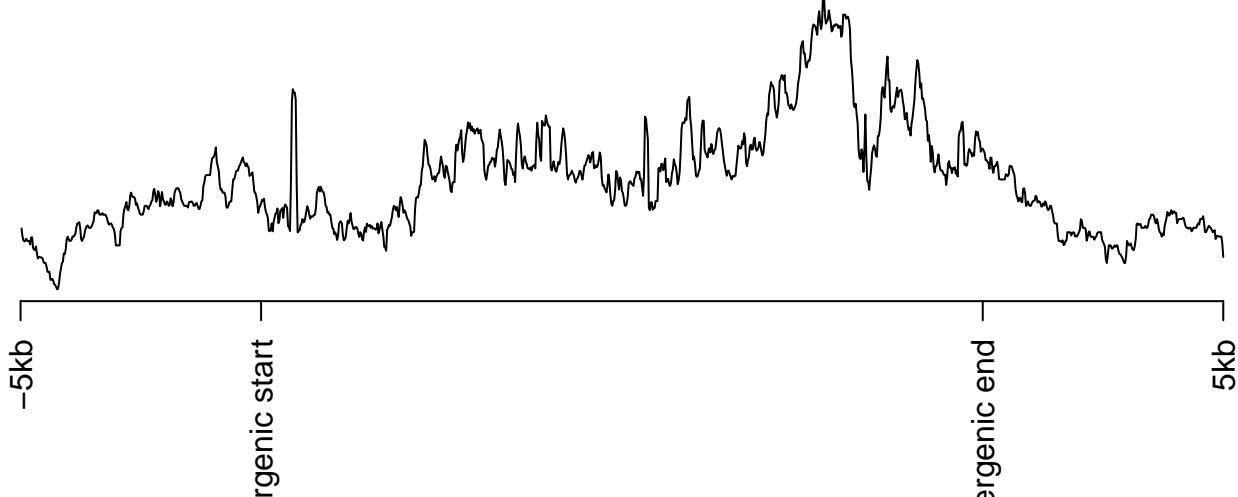
HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc443



Average coverage per intergenic region

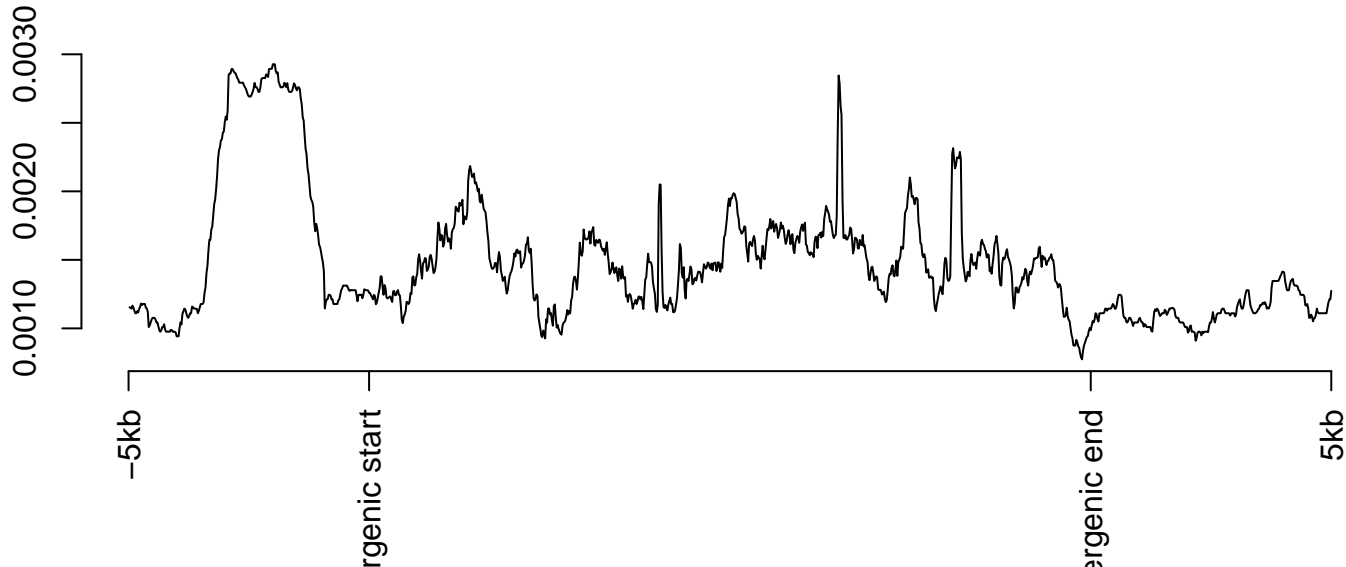
HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc444

0.0015 0.0025 0.0035



Average coverage per intergenic region

HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc445



HumanAstroCulture PositivePooled E.589 scCLTdegenNuc439

Average coverage per intergenic region

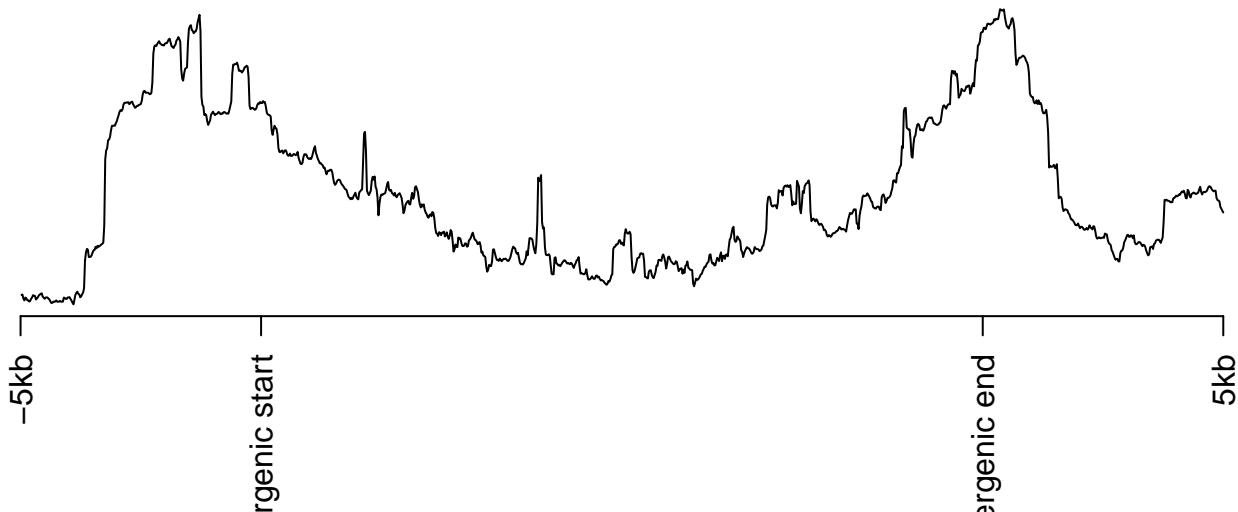
0.008
0.005
0.002

-5kb

intergenic start

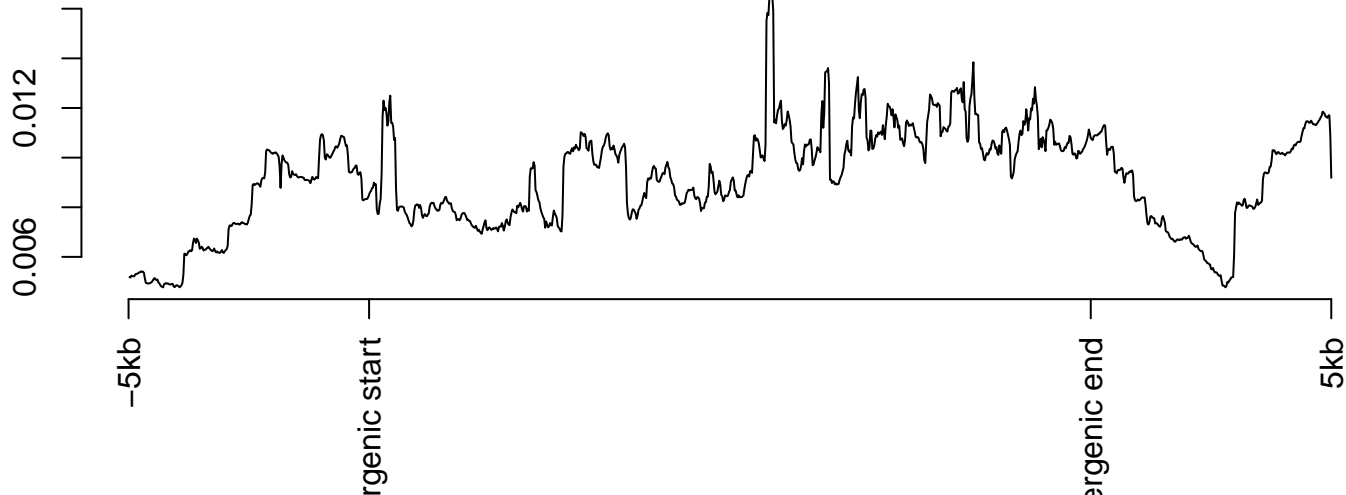
intergenic end

5kb



Average coverage per intergenic region

HumanAstroCulture PositivePooled E.589 scCLTdegenNuc440



Average coverage per intergenic region

HumanAstroCulture PositiveMerged E.– HumanAstroCulturePositiveMerged

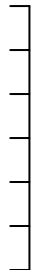
0.007 0.010 0.013

-5kb

intergenic start

intergenic end

5kb



Average coverage per intergenic region

HumanAstroCulture PositiveAll E.– HumanAstroCulturePositiveAll

