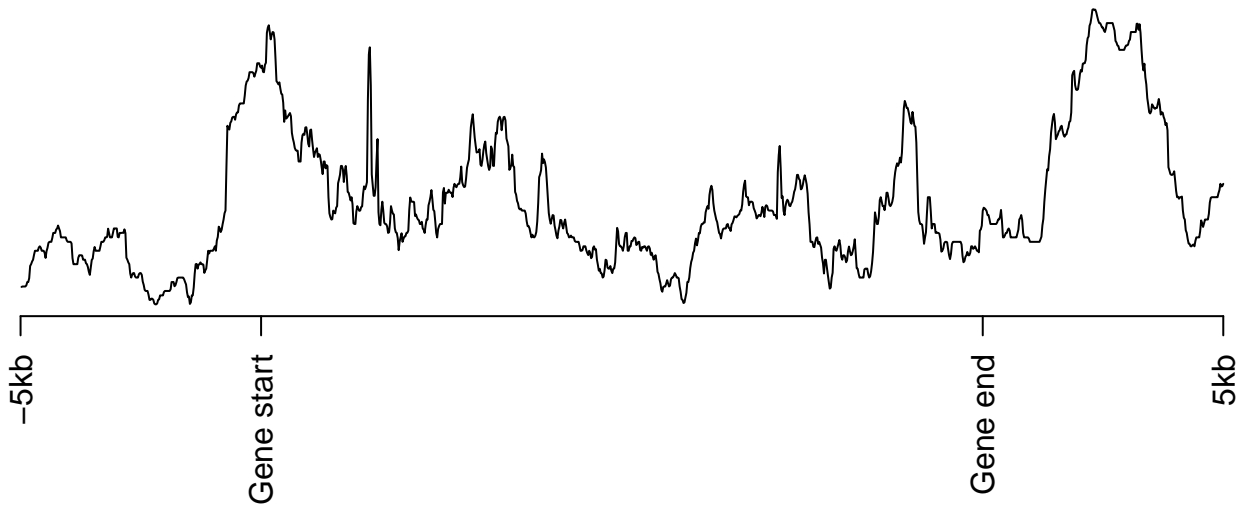


HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc441

Average coverage per gene

0.0005 0.0015



HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc442

Average coverage per gene

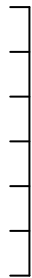
0.0060
0.0045
0.0030

-5kb

Gene start

Gene end

5kb



HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc443

Average coverage per gene

0.0005 0.0020

-5kb

Gene start

Gene end

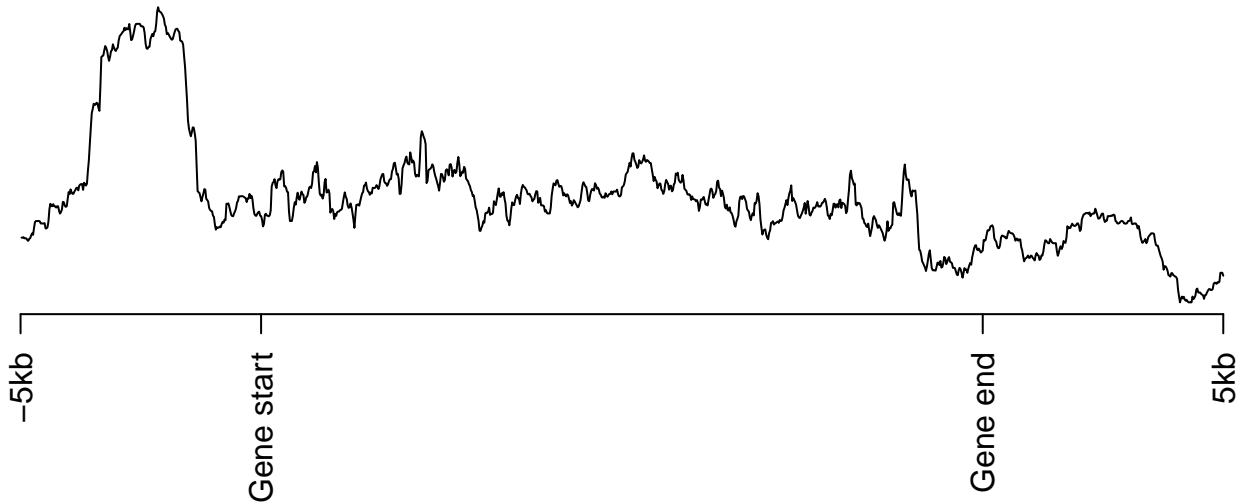
5kb



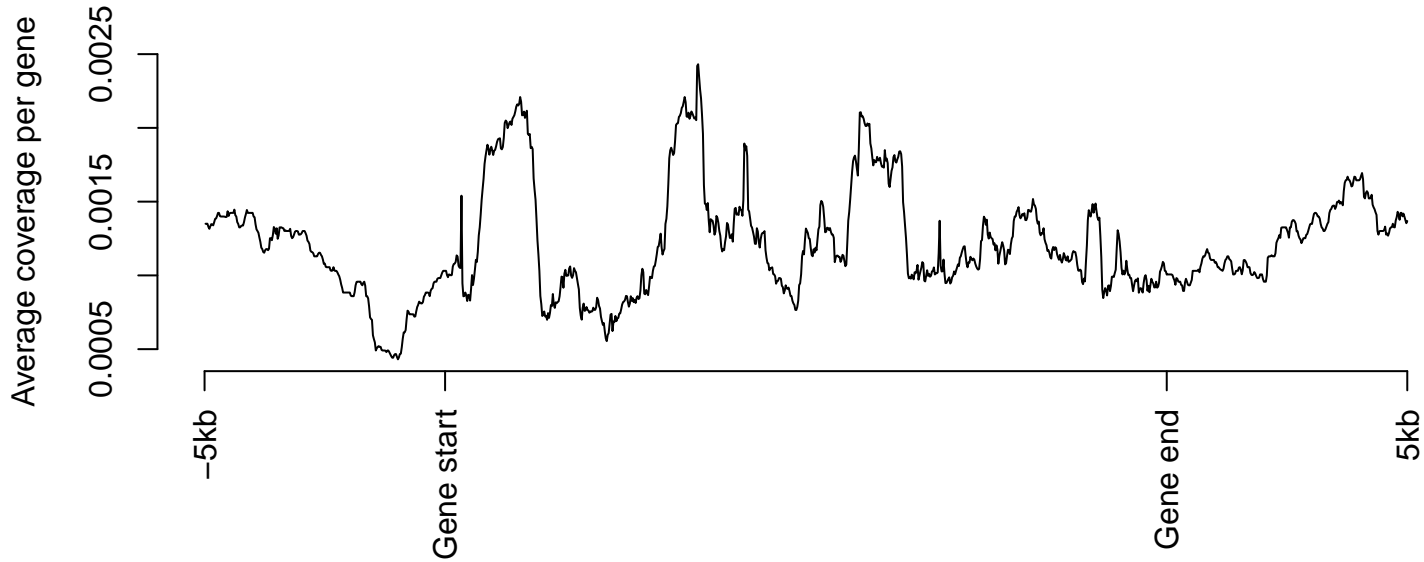
HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc444

Average coverage per gene

0.0015 0.0030



HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc445



HumanAstroCulture PositivePooled E.589 scCLTdegenNuc439

Average coverage per gene

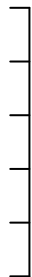
0.003 0.005 0.007

-5kb

Gene start

Gene end

5kb



HumanAstroCulture PositivePooled E.589 scCLTdegenNuc440

Average coverage per gene

0.006 0.010

-5kb

Gene start

Gene end

5kb



HumanAstroCulture PositiveMerged E.– HumanAstroCulturePositiveMerged

Average coverage per gene

0.012
0.008

-5kb

Gene start

Gene end

5kb



HumanAstroCulture PositiveAll E.– HumanAstroCulturePositiveAll

Average coverage per gene

0.018 0.024 0.030

-5kb

Gene start

Gene end

5kb

