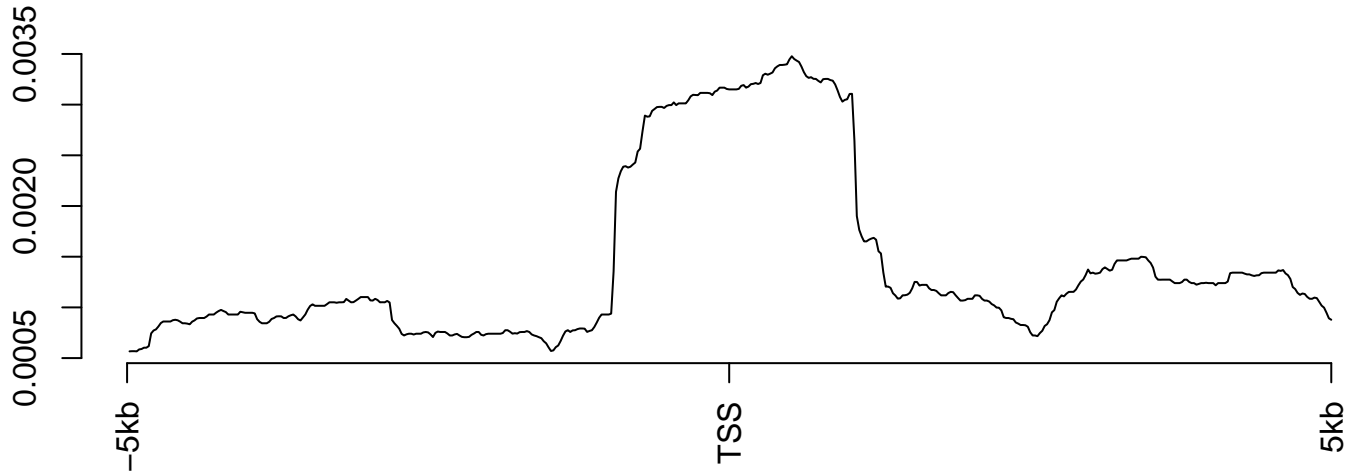
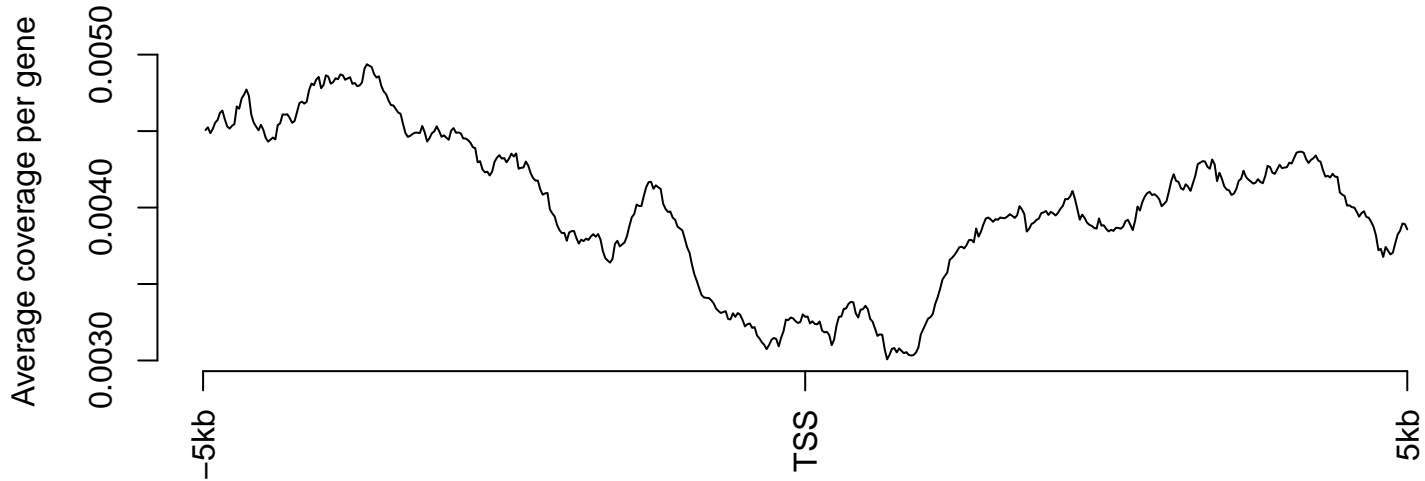


# HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc441

Average coverage per gene



# HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc442



# HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc443

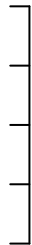
Average coverage per gene

0.0012  
0.0006

-5kb

TSS

5kb



# HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc444

Average coverage per gene

0.0020 0.0030

-5kb

TSS

5kb



# HumanAstroCulture PositiveSingle E.589 scCLTdegenNuc445

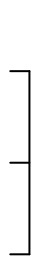
Average coverage per gene

0.0020  
0.0010

-5kb

TSS

5kb



# HumanAstroCulture PositivePooled E.589 scCLTdegenNuc439

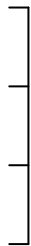
Average coverage per gene

0.007  
0.005

-5kb

TSS

5kb



# HumanAstroCulture PositivePooled E.589 scCLTdegenNuc440

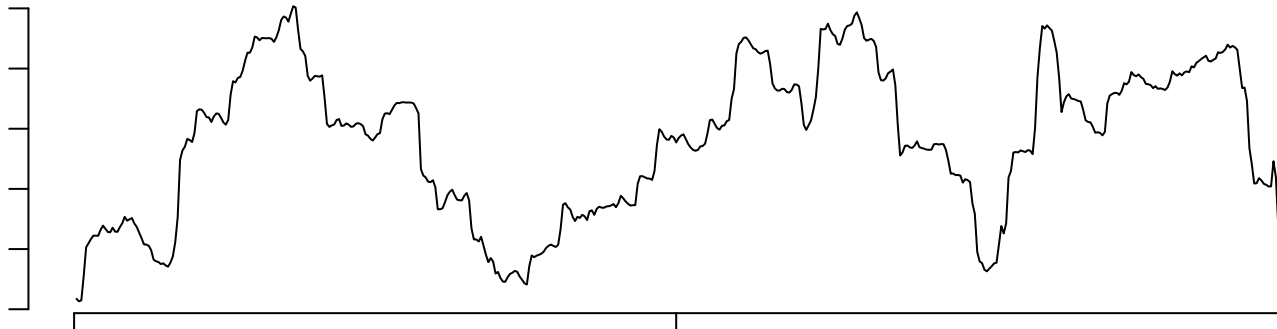
Average coverage per gene

0.007 0.009 0.011

-5kb

TSS

5kb



# HumanAstroCulture PositiveMerged E.– HumanAstroCulturePositiveMerged

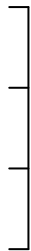
Average coverage per gene

0.011  
0.009

-5kb

TSS

5kb





# HumanAstroCulture PositiveAll E.– HumanAstroCulturePositiveAll

Average coverage per gene

0.022 0.026 0.030

-5kb

TSS

5kb

