

# K562 NegCtrlSingle E.571 scCLTdegenNuc383

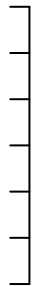
Average coverage per gene

0.00025  
0.00005

-5kb

TSS

5kb



# K562 NegCtrlSingle E.571 scCLTdegenNuc384

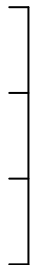
Average coverage per gene

0.00000 0.00010

-5kb

TSS

5kb



# K562 NegCtrlSingle E.571 scCLTdegenNuc386

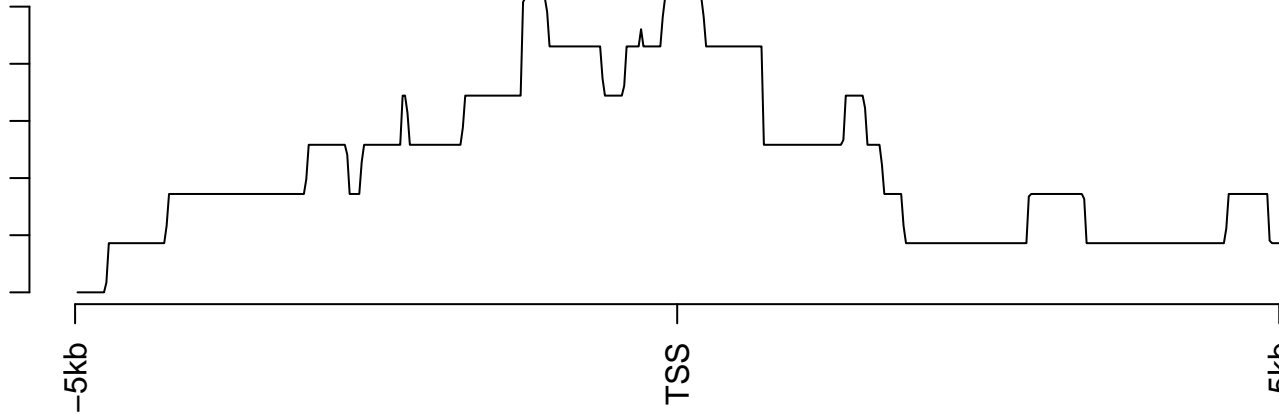
Average coverage per gene

0e+00 6e-05

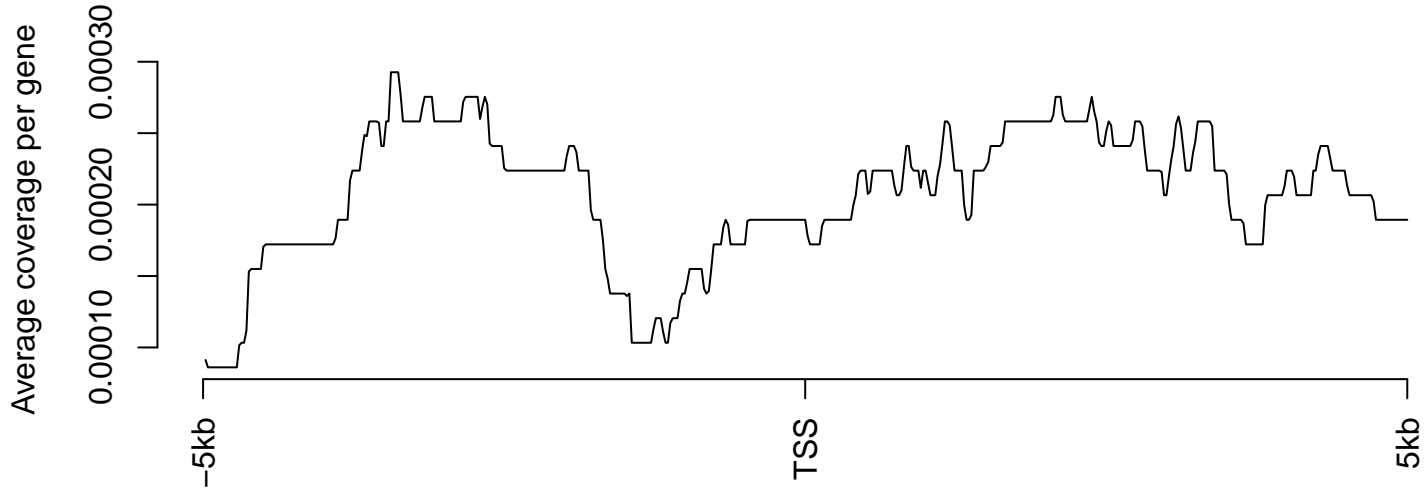
-5kb

TSS

5kb



# K562 NegCtrlSingle E.579 scCLTdegenNuc417



# K562 NegCtrlSingle E.579 scCLTdegenNuc418

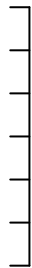
Average coverage per gene

0.00030 0.00050

-5kb

TSS

5kb



# K562 NegCtrlSingle E.579 scCLTdegenNuc420

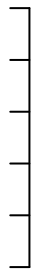
Average coverage per gene

0.00045  
0.00030

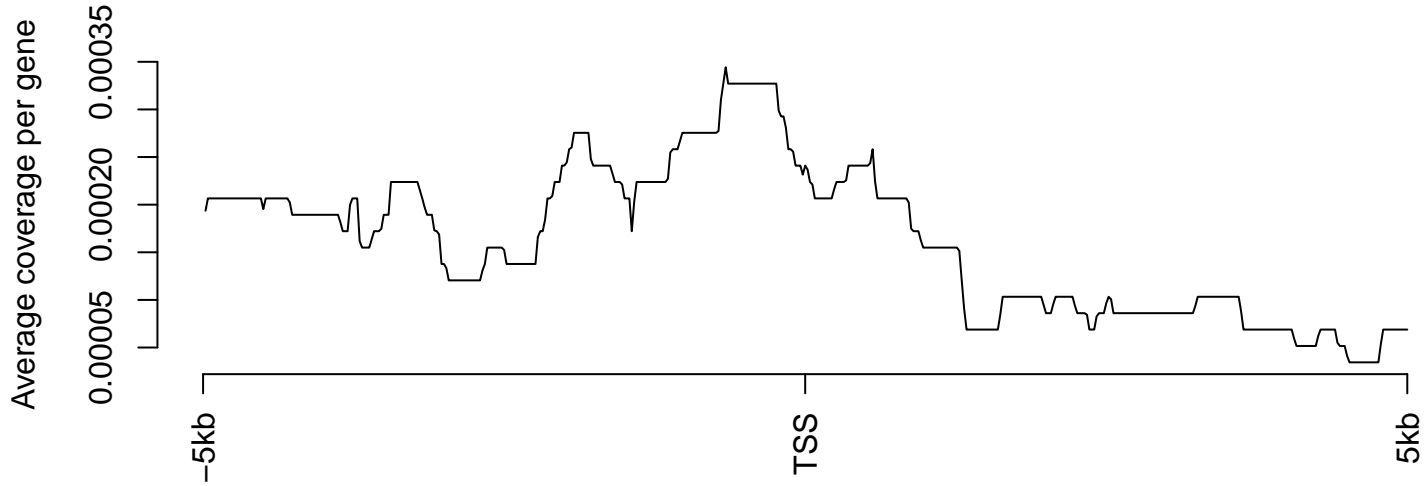
-5kb

TSS

5kb



# K562 NegCtrlSingle E.579 scCLTdegenNuc422



# K562 NegCtrlSingle E.579 scCLTdegenNuc423

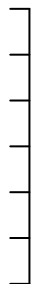
Average coverage per gene

4e-04 7e-04 1e-03

-5kb

TSS

5kb





# K562 NegCtrlSingle E.579 scCLTdegenNuc424

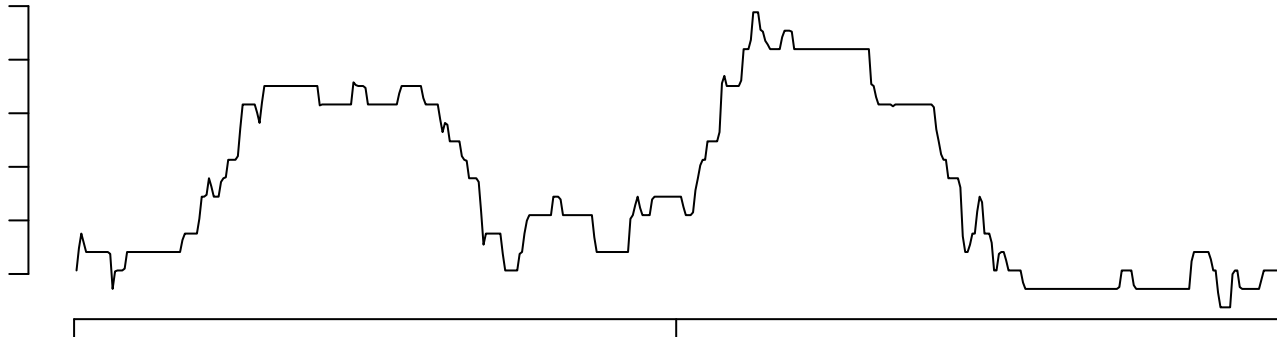
Average coverage per gene

0.00010 0.00025

-5kb

TSS

5kb



# K562 NegCtrlSingle E.579 scCLTdegenNuc425

Average coverage per gene

0.00010 0.00020

-5kb

TSS

5kb



# K562 NegCtrlSingle E.671 scCLTdegenNuc897

Average coverage per gene

0.0020 0.0030

-5kb

TSS

5kb



# K562 NegCtrlSingle E.671 scCLTdegenNuc898

Average coverage per gene

0.0004 0.0007 0.0010

-5kb

TSS

5kb



# K562 NegCtrlSingle E.671 scCLTdegenNuc900

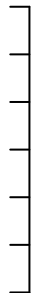
Average coverage per gene

3e-04  
6e-04  
9e-04

-5kb

TSS

5kb



# K562 NegCtrlSingle E.671 scCLTdegenNuc902

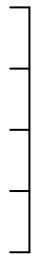
Average coverage per gene

1e-04 3e-04 5e-04

-5kb

TSS

5kb



# K562 NegCtrlSingle E.671 scCLTdegenNuc903

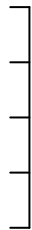
Average coverage per gene

0.0018  
0.0012

-5kb

TSS

5kb



# K562 NegCtrlSingle E.676 scCLTdegenNuc950

Average coverage per gene

0.0004 0.0008 0.0012

-5kb

TSS

5kb





# K562 NegCtrlSingle E.678 scCLTdegenNuc957

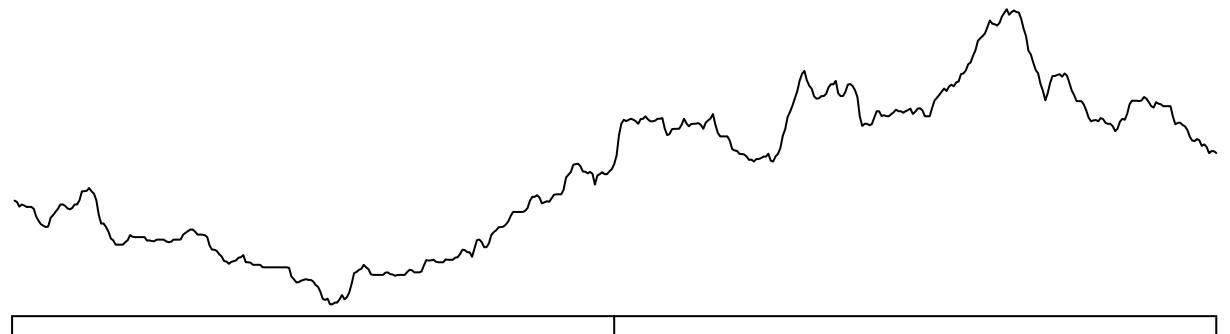
Average coverage per gene

0.0010 0.0020

-5kb

TSS

5kb



# K562 NegCtrlPooled E.591 scCLTdegenNuc463

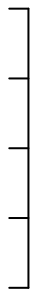
Average coverage per gene

0.010  
0.008  
0.006

-5kb

TSS

5kb



# K562 NegCtrlPooled E.591 scCLTdegenNuc464

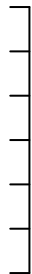
Average coverage per gene

0.003 0.006 0.009

-5kb

TSS

5kb



# K562 NegCtrlMerged E.- K562NegCtrlMerged

Average coverage per gene

0.009 0.011

-5kb

TSS

5kb



# K562 NegCtrlAll E.- K562NegCtrlAll

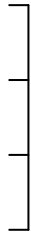
Average coverage per gene

0.022 0.026

-5kb

TSS

5kb



# K562 PositiveSingle E.571 scCLTdegenNuc404

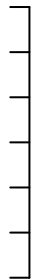
Average coverage per gene

0.00025 0.00045

-5kb

TSS

5kb



# K562 PositiveSingle E.571 scCLTdegenNuc406

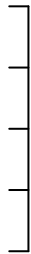
Average coverage per gene

0.00030  
0.00015

-5kb

TSS

5kb



# K562 PositiveSingle E.571 scCLTdegenNuc408

Average coverage per gene

7e-04

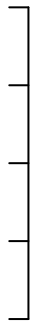
5e-04

3e-04

-5kb

TSS

5kb





# K562 PositiveSingle E.571 scCLTdegenNuc409

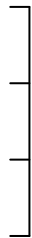
Average coverage per gene

0.0040 0.0050

-5kb

TSS

5kb



# K562 PositiveSingle E.571 scCLTdegenNuc410

Average coverage per gene

0.0014  
0.0008

-5kb

TSS

5kb



# K562 PositiveSingle E.571 scCLTdegenNuc411

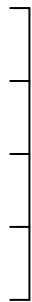
Average coverage per gene

0.0004 0.0008 0.0012

-5kb

TSS

5kb



# K562 PositiveSingle E.574 scCLTdegenNuc334

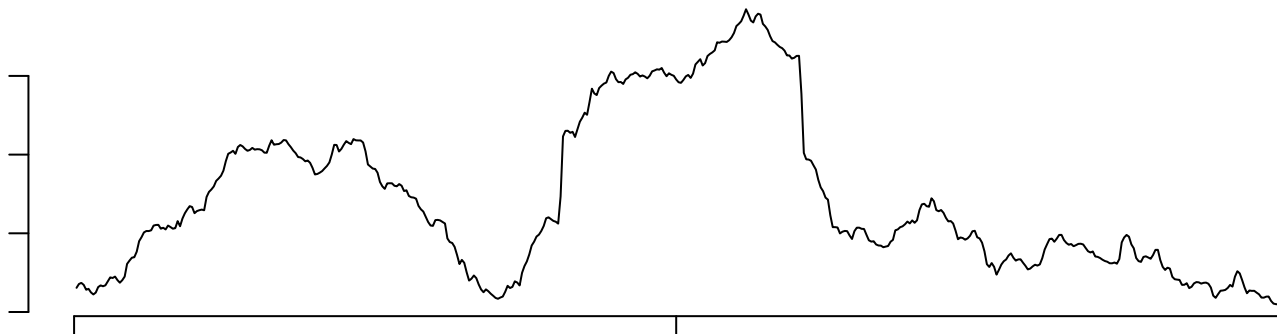
Average coverage per gene

0.004 0.006

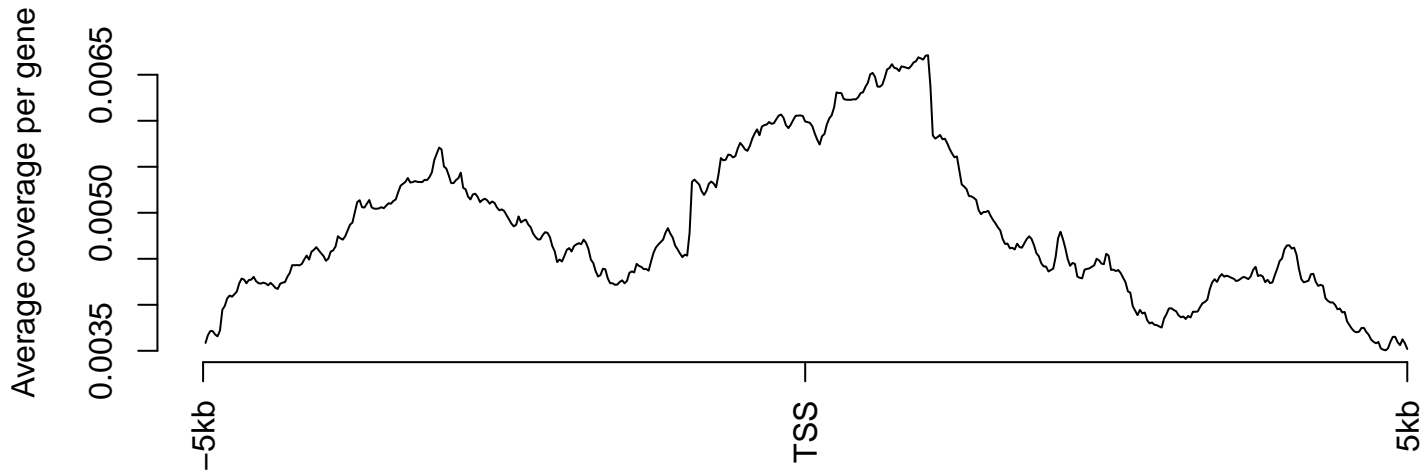
-5kb

TSS

5kb



# K562 PositiveSingle E.574 scCLTdegenNuc335



# K562 PositiveSingle E.574 scCLTdegenNuc336

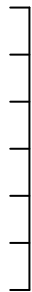
Average coverage per gene

0.0080  
0.0065  
0.0050

-5kb

TSS

5kb



# K562 PositiveSingle E.579 scCLTdegenNuc337

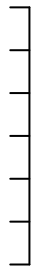
Average coverage per gene

0.0020 0.0026 0.0032

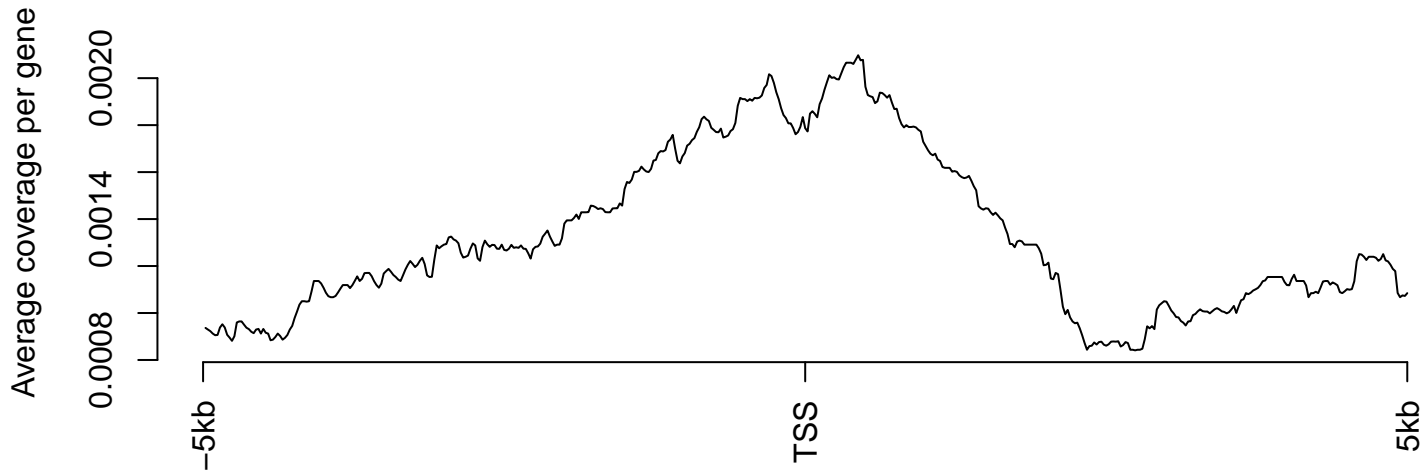
-5kb

TSS

5kb



# K562 PositiveSingle E.579 scCLTdegenNuc338





# K562 PositiveSingle E.579 scCLTdegenNuc339

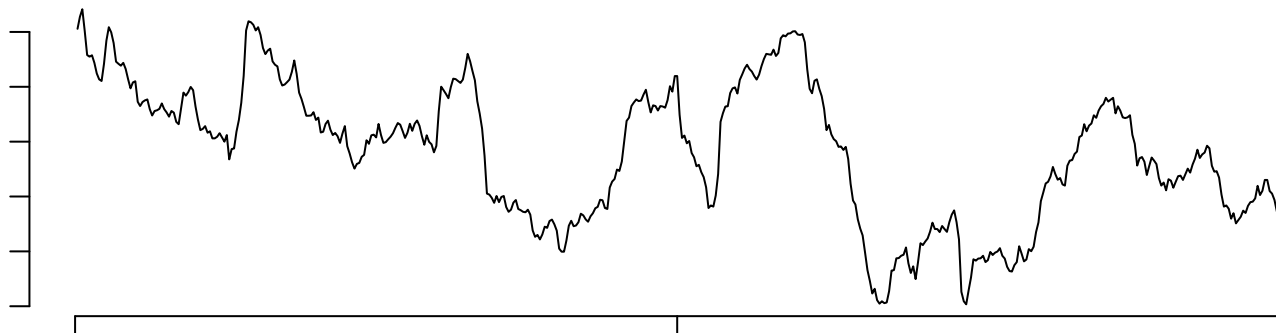
Average coverage per gene

0.0034  
0.0028

-5kb

TSS

5kb



# K562 PositiveSingle E.579 scCLTdegenNuc340

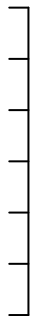
Average coverage per gene

0.0030 0.0045 0.0060

-5kb

TSS

5kb



# K562 PositiveSingle E.579 scCLTdegenNuc341

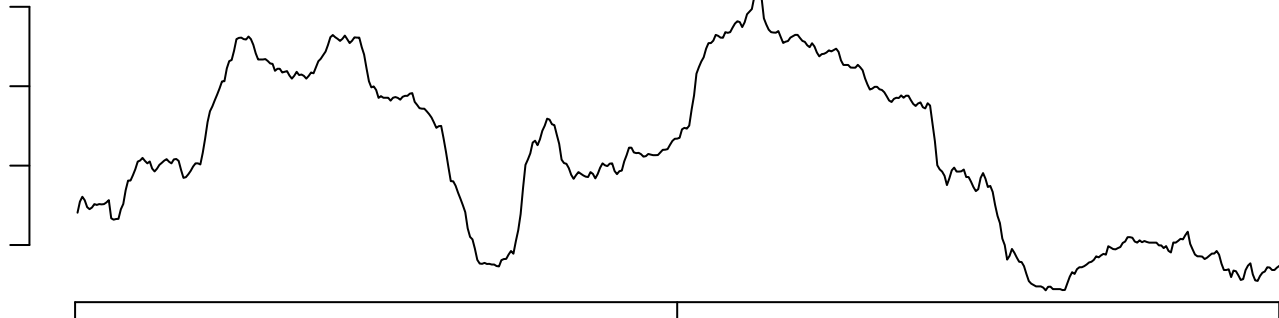
Average coverage per gene

0.0015  
0.0025

-5kb

TSS

5kb



# K562 PositiveSingle E.580 scCLTdegenNuc342

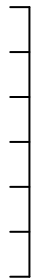
Average coverage per gene

0.0022  
0.0016  
0.0010

-5kb

TSS

5kb



# K562 PositiveSingle E.580 scCLTdegenNuc343

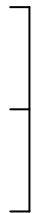
Average coverage per gene

0.0015  
0.0025

-5kb

TSS

5kb



# K562 PositiveSingle E.580 scCLTdegenNuc345

Average coverage per gene

0.0016 0.0022

-5kb

TSS

5kb



# K562 PositiveSingle E.580 scCLTdegenNuc428

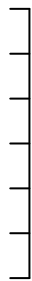
Average coverage per gene

0.017  
0.014  
0.011

-5kb

TSS

5kb



# K562 PositiveSingle E.588 scCLTdegenNuc323

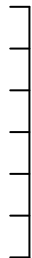
Average coverage per gene

0.0017  
0.0014  
0.0011

-5kb

TSS

5kb





# K562 PositiveSingle E.588 scCLTdegenNuc324

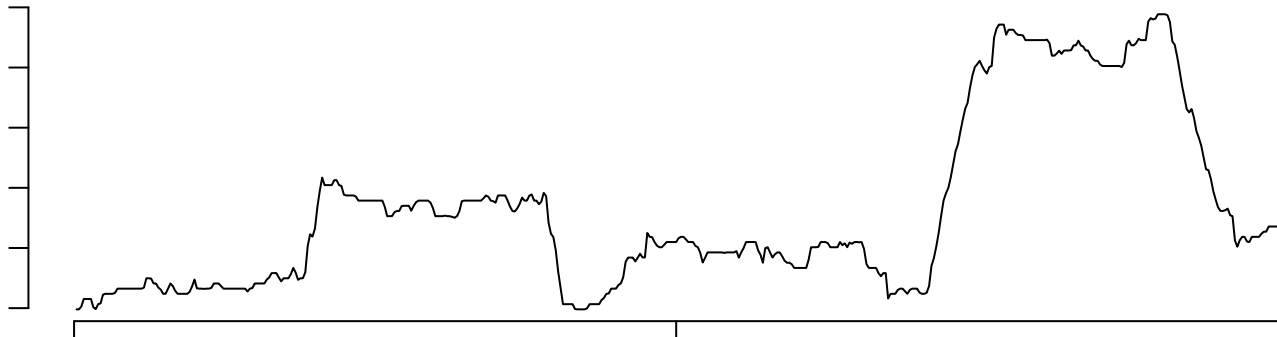
Average coverage per gene

0.0010  
0.0004

-5kb

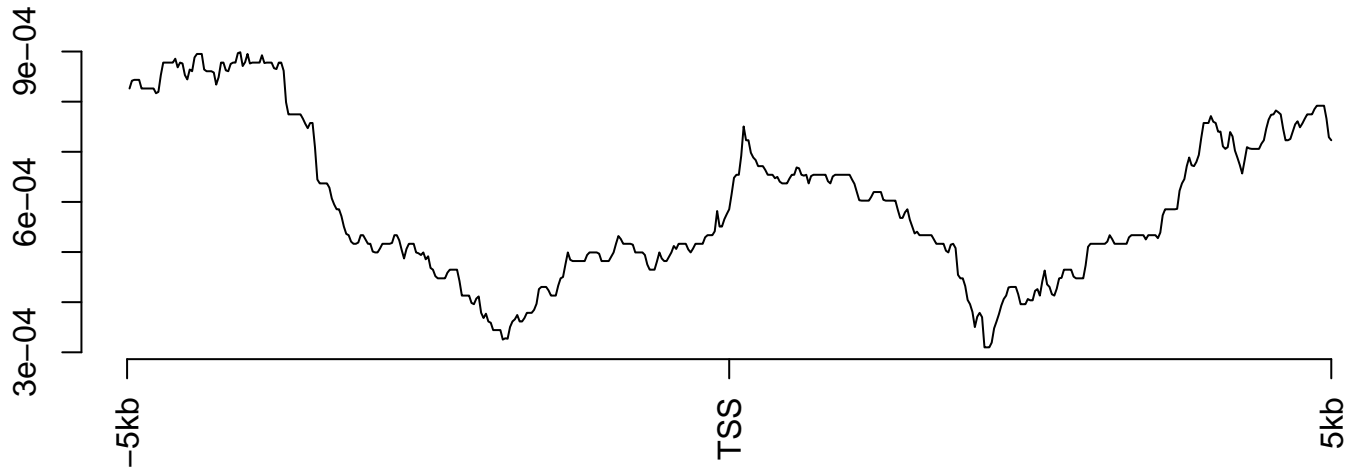
TSS

5kb



# K562 PositiveSingle E.588 scCLTdegenNuc325

Average coverage per gene



# K562 PositiveSingle E.588 scCLTdegenNuc446

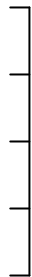
Average coverage per gene

0.0025 0.0035 0.0045

-5kb

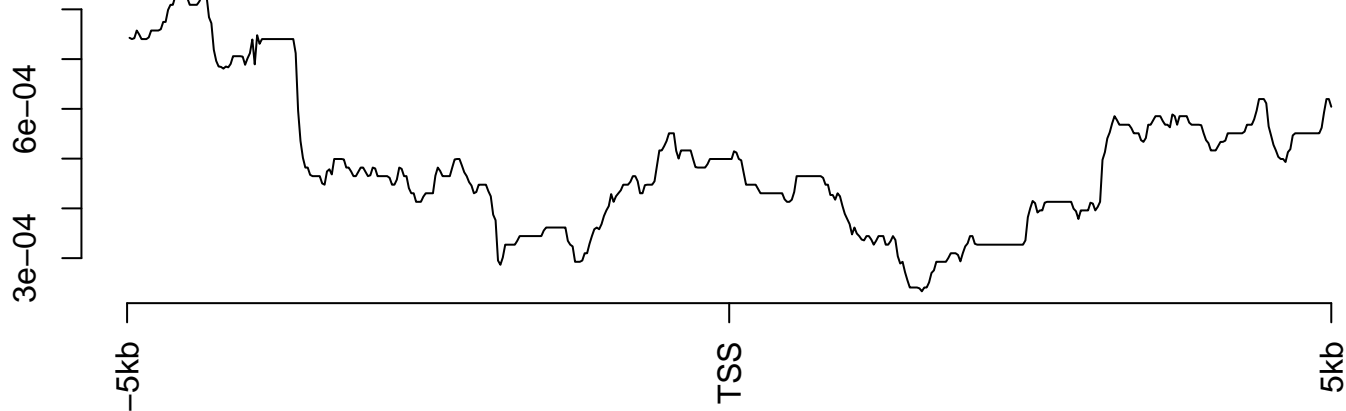
TSS

5kb



# K562 PositiveSingle E.588 scCLTdegenNuc447

Average coverage per gene



# K562 PositiveSingle E.588 scCLTdegenNuc448

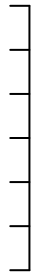
Average coverage per gene

1e-04 4e-04 7e-04

-5kb

TSS

5kb



# K562 PositiveSingle E.588 scCLTdegenNuc452

Average coverage per gene

0.0035  
0.0025

-5kb

TSS

5kb



# K562 PositiveSingle E.588 scCLTdegenNuc454

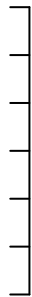
Average coverage per gene

0.0015  
0.0012  
0.0009

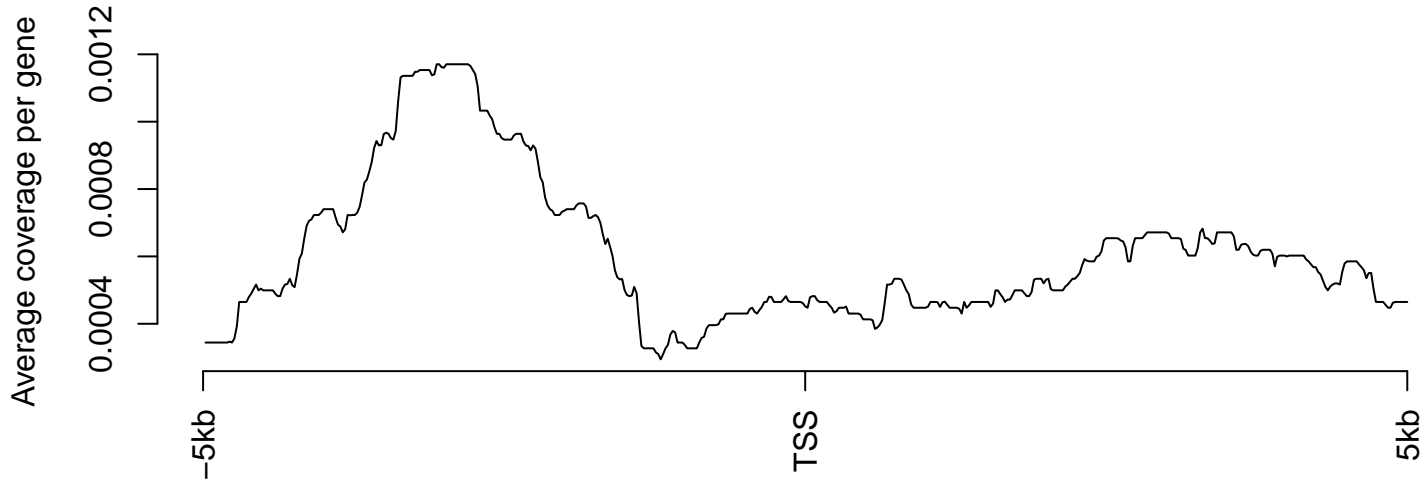
-5kb

TSS

5kb



# K562 PositiveSingle E.588 scCLTdegenNuc455





# K562 PositiveSingle E.588 scCLTdegenNuc456

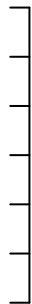
Average coverage per gene

0.0030  
0.0024  
0.0018

-5kb

TSS

5kb



# K562 PositiveSingle E.589 scCLTdegenNuc326

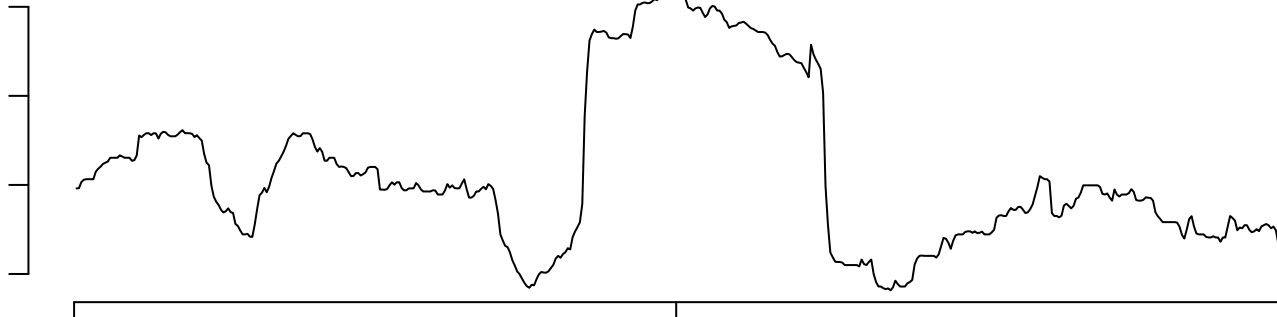
Average coverage per gene

0.0020  
0.0010

-5kb

TSS

5kb



# K562 PositiveSingle E.589 scCLTdegenNuc348

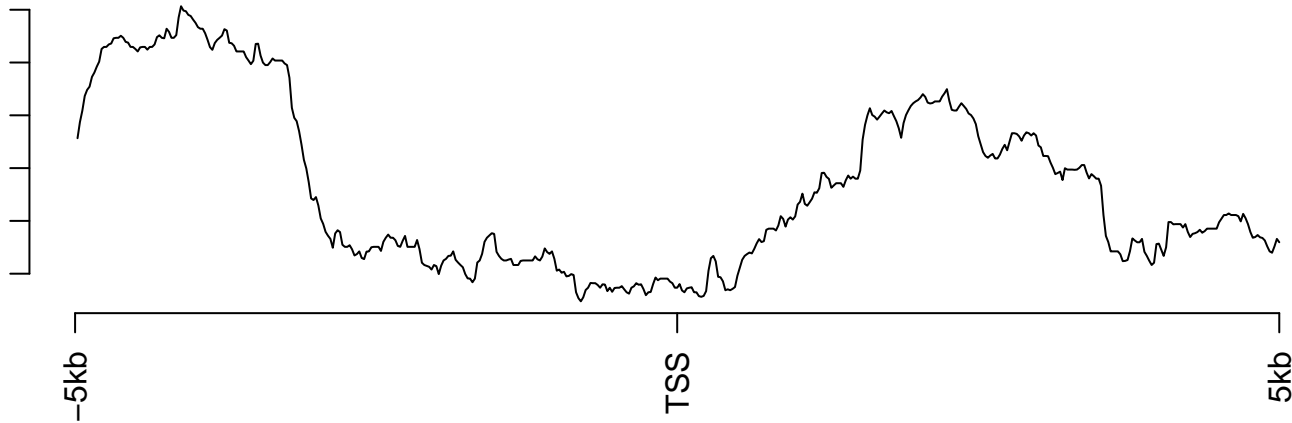
Average coverage per gene

0.0016  
0.0010

-5kb

TSS

5kb



# K562 PositiveSingle E.665 scCLTdegenNuc766

Average coverage per gene

0.0035  
0.0025

-5kb

TSS

5kb



# K562 PositiveSingle E.665 scCLTdegenNuc767

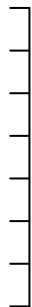
Average coverage per gene

0.0055  
0.0040  
0.0025

-5kb

TSS

5kb



# K562 PositiveSingle E.665 scCLTdegenNuc768

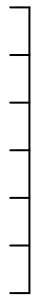
Average coverage per gene

0.0002 0.0008 0.0014

-5kb

TSS

5kb



# K562 PositiveSingle E.665 scCLTdegenNuc769

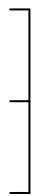
Average coverage per gene

0.0025  
0.0015

-5kb

TSS

5kb



# K562 PositiveSingle E.665 scCLTdegenNuc770

Average coverage per gene

0.0010 0.0013

-5kb

TSS

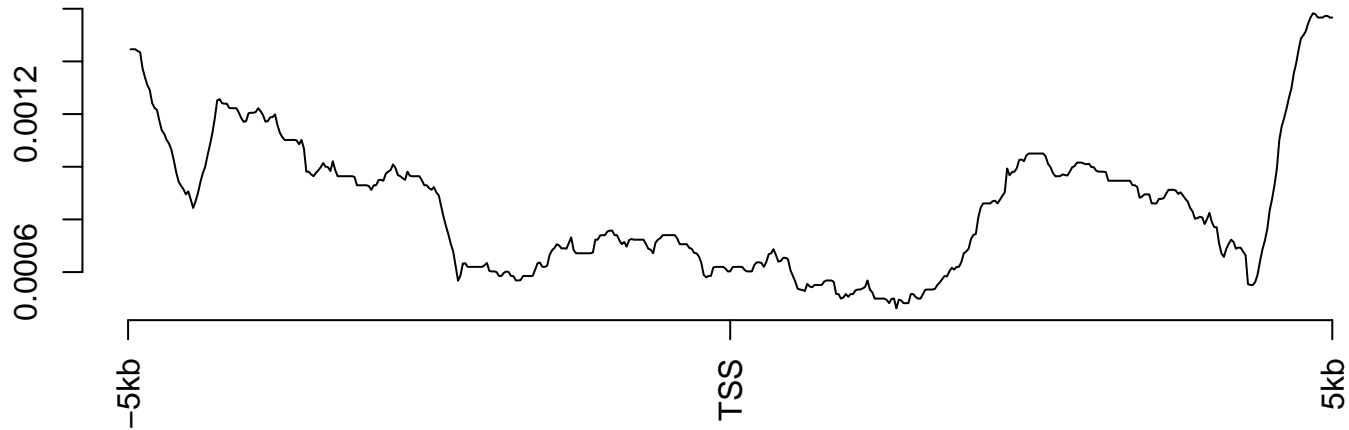
5kb





# K562 PositiveSingle E.665 scCLTdegenNuc771

Average coverage per gene



# K562 PositiveSingle E.665 scCLTdegenNuc772

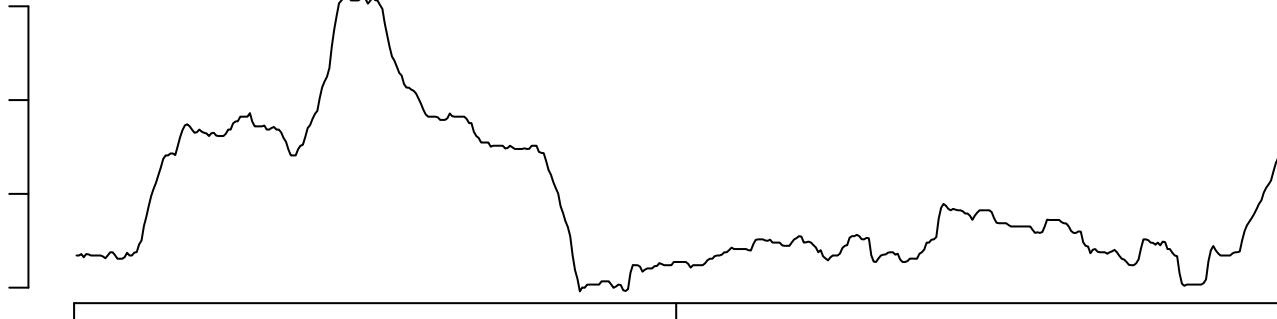
Average coverage per gene

0.0015  
0.0005

-5kb

TSS

5kb



# K562 PositiveSingle E.665 scCLTdegenNuc773

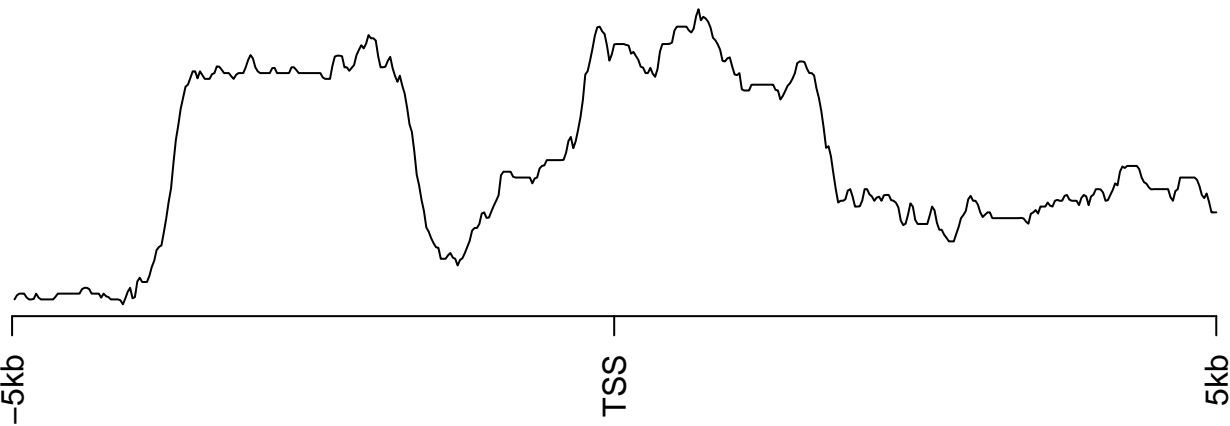
Average coverage per gene

0.0006 0.0010

-5kb

TSS

5kb



# K562 PositiveSingle E.665 scCLTdegenNuc774

Average coverage per gene

0.0002 0.0008

-5kb

TSS

5kb



# K562 PositiveSingle E.665 scCLTdegenNuc775

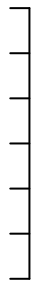
Average coverage per gene

0.00015 0.00035

-5kb

TSS

5kb



# K562 PositiveSingle E.673 scCLTdegenNuc906\_b

Average coverage per gene

2e-04  
5e-04

-5kb

TSS

5kb



# K562 PositiveSingle E.673 scCLTdegenNuc907\_b

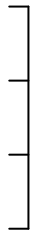
Average coverage per gene

2e-04  
4e-04

-5kb

TSS

5kb



# K562 PositiveSingle E.673 scCLTdegenNuc908\_b

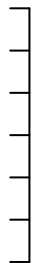
Average coverage per gene

0.0006 0.0012 0.0018

-5kb

TSS

5kb





# K562 PositiveSingle E.675 scCLTdegenNuc933

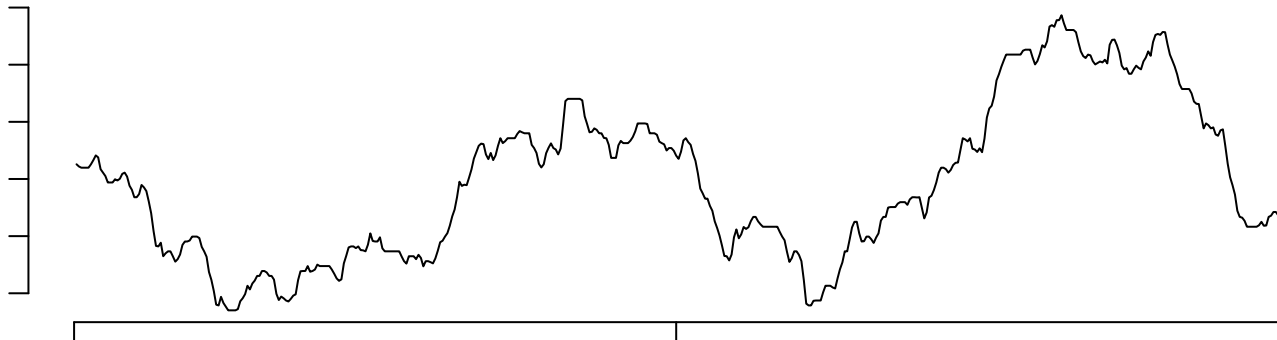
Average coverage per gene

0.0014  
0.0008

-5kb

TSS

5kb



# K562 PositiveSingle E.675 scCLTdegenNuc935

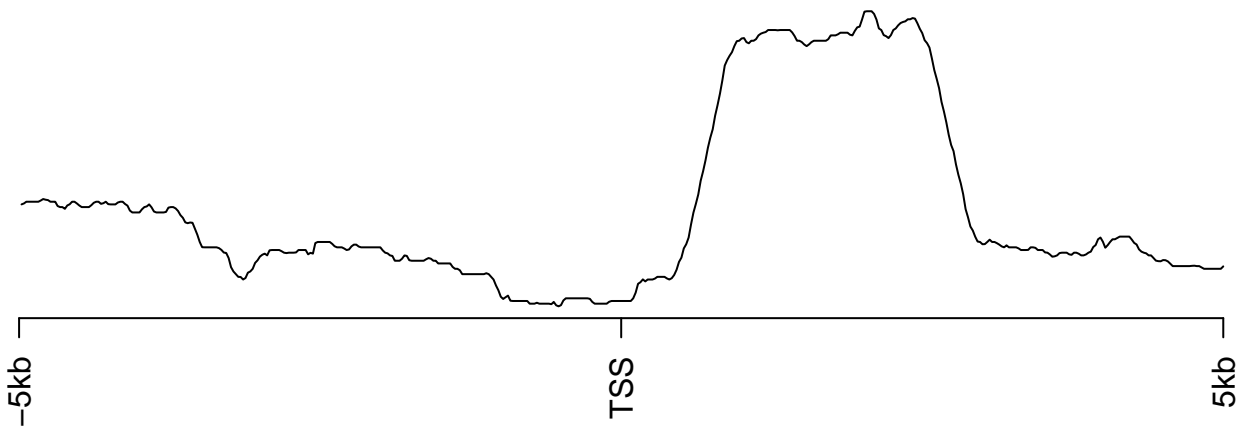
Average coverage per gene

0.0015  
0.0005

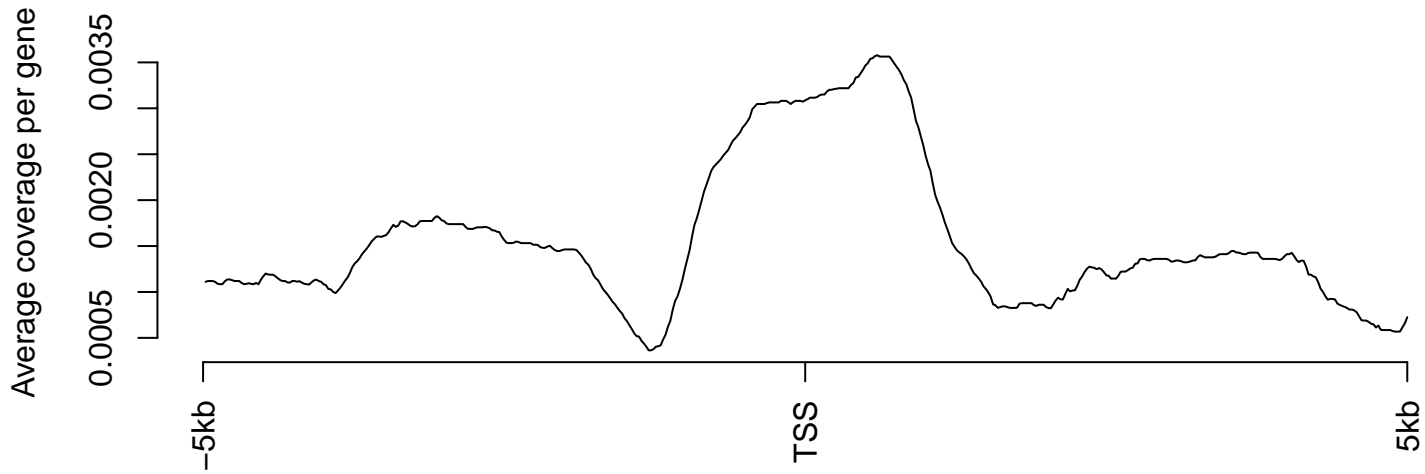
-5kb

TSS

5kb



# K562 PositiveSingle E.675 scCLTdegenNuc936



# K562 PositiveSingle E.675 scCLTdegenNuc938

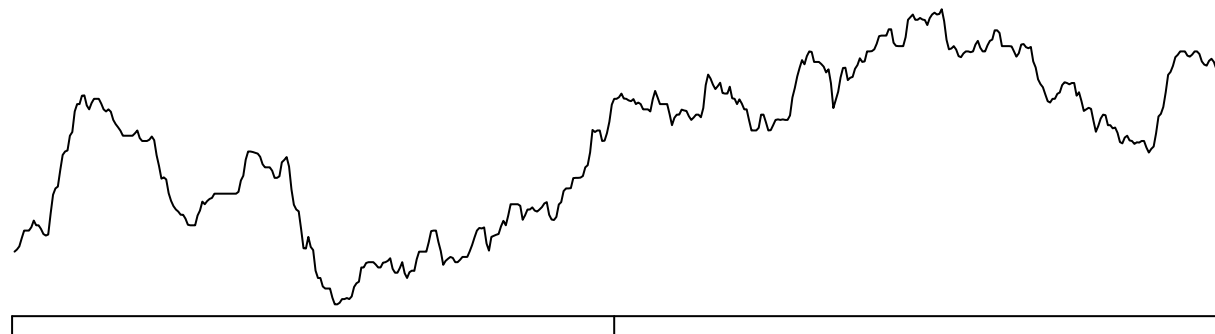
Average coverage per gene

0.0014  
0.0008

-5kb

TSS

5kb



# K562 PositiveSingle E.675 scCLTdegenNuc942

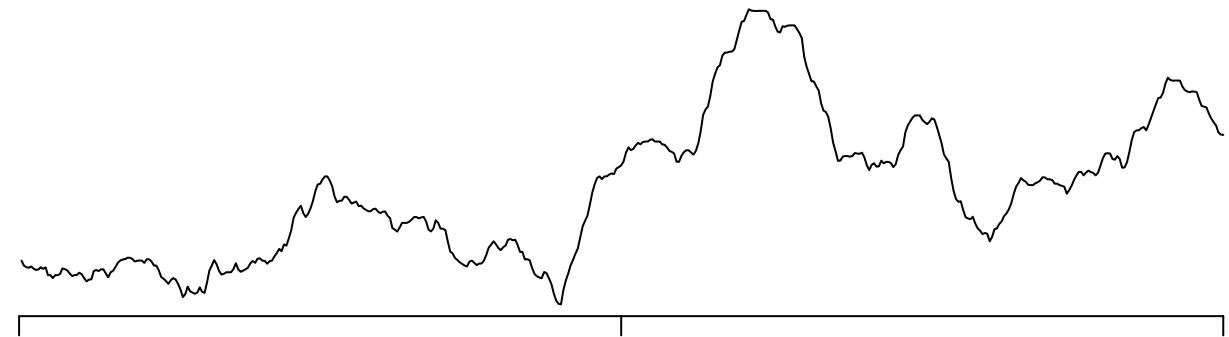
Average coverage per gene

0.0015 0.0025

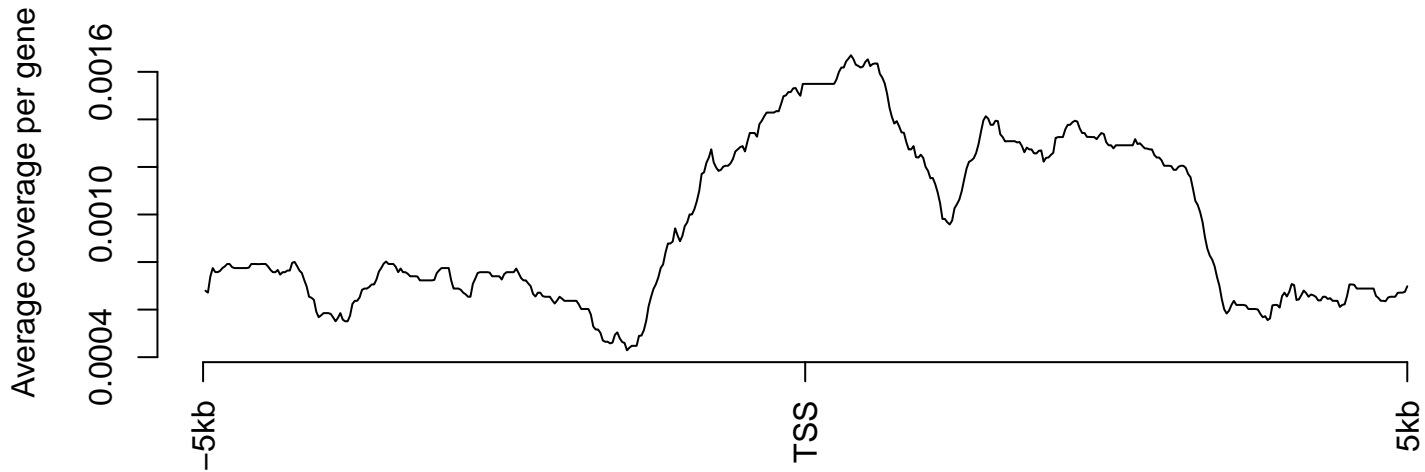
-5kb

TSS

5kb



# K562 PositiveSingle E.675 scCLTdegenNuc943



# K562 PositiveSingle E.676 scCLTdegenNuc944

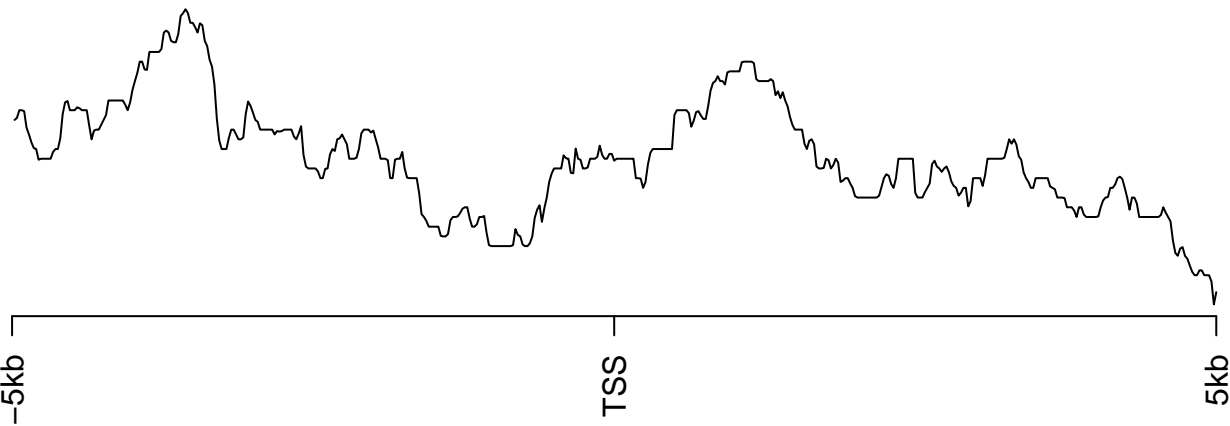
Average coverage per gene

4e-04  
7e-04

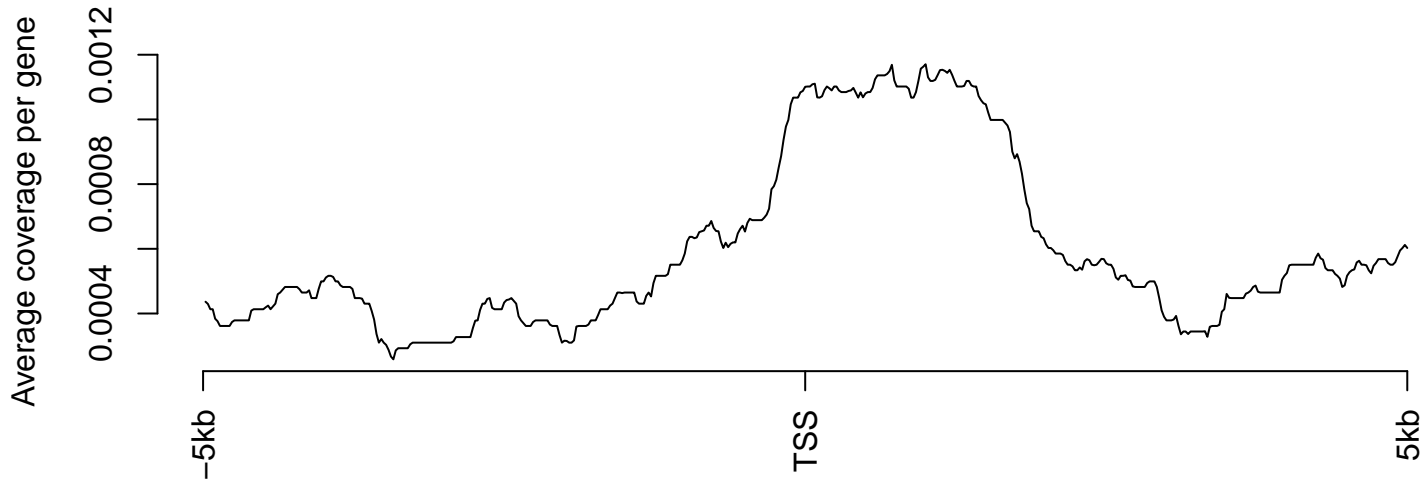
-5kb

TSS

5kb



# K562 PositiveSingle E.676 scCLTdegenNuc945





# K562 PositiveSingle E.676 scCLTdegenNuc946

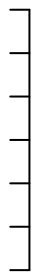
Average coverage per gene

0.0010 0.0016 0.0022

-5kb

TSS

5kb



# K562 PositiveSingle E.676 scCLTdegenNuc947

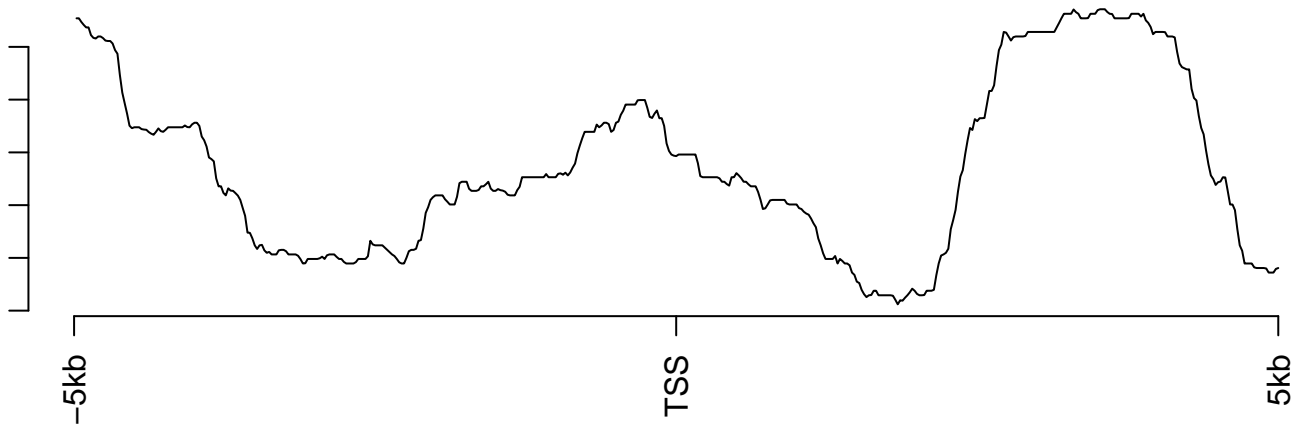
Average coverage per gene

0.0002 0.0008

-5kb

TSS

5kb



# K562 PositiveSingle E.676 scCLTdegenNuc948

Average coverage per gene

2e-04  
6e-04

-5kb

TSS

5kb



# K562 PositiveSingle E.676 scCLTdegenNuc949

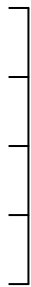
Average coverage per gene

0.006  
0.004  
0.002

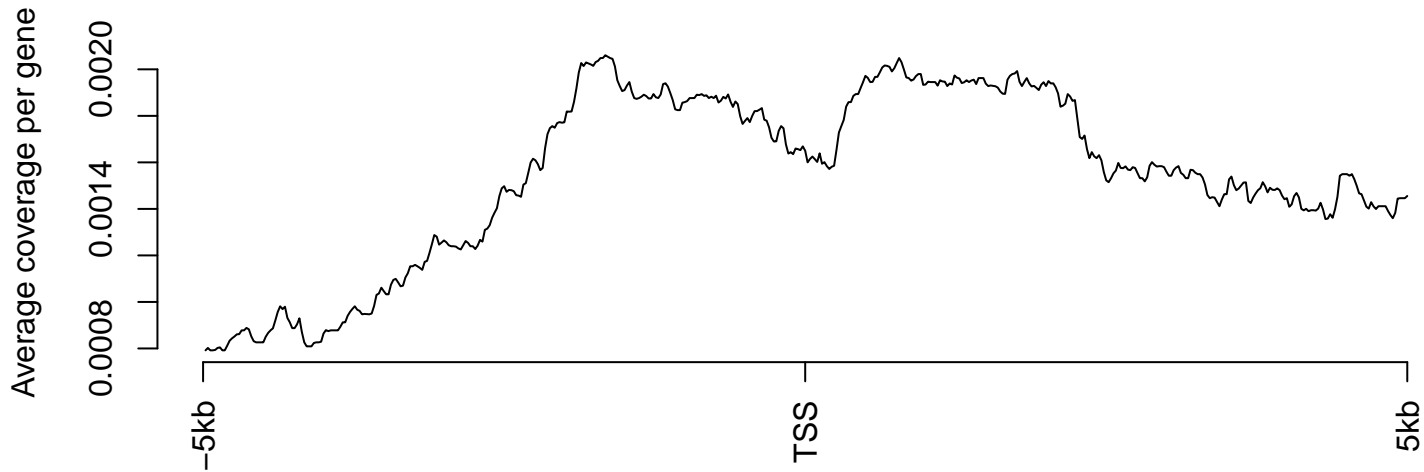
-5kb

TSS

5kb



# K562 PositiveSingle E.678 scCLTdegenNuc951



# K562 PositiveSingle E.678 scCLTdegenNuc952

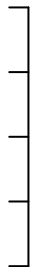
Average coverage per gene

0.0006 0.0010 0.0014

-5kb

TSS

5kb



# K562 PositiveSingle E.678 scCLTdegenNuc953

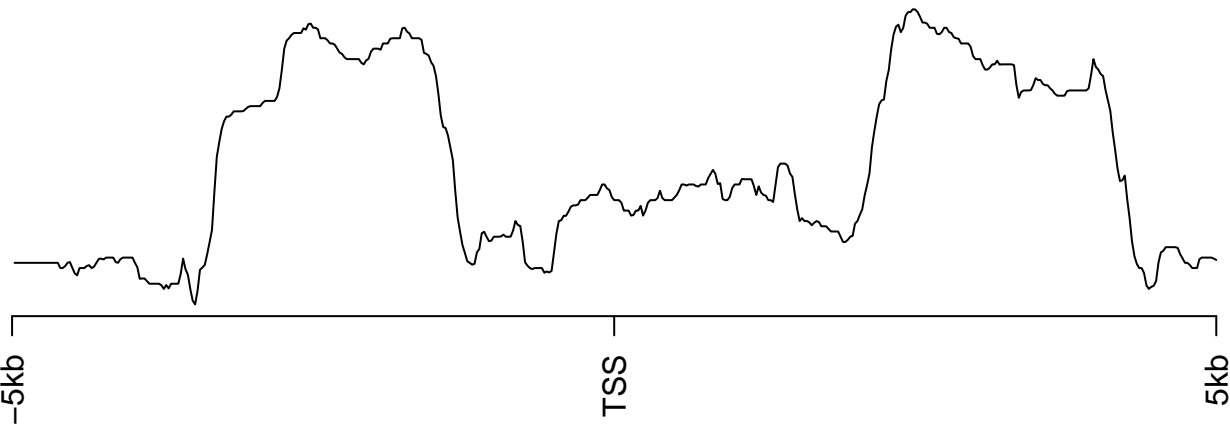
Average coverage per gene

0.0012  
0.0006

-5kb

TSS

5kb



# K562 PositiveSingle E.678 scCLTdegenNuc954

Average coverage per gene

0.0006 0.0012

-5kb

TSS

5kb





# K562 PositiveSingle E.678 scCLTdegenNuc955

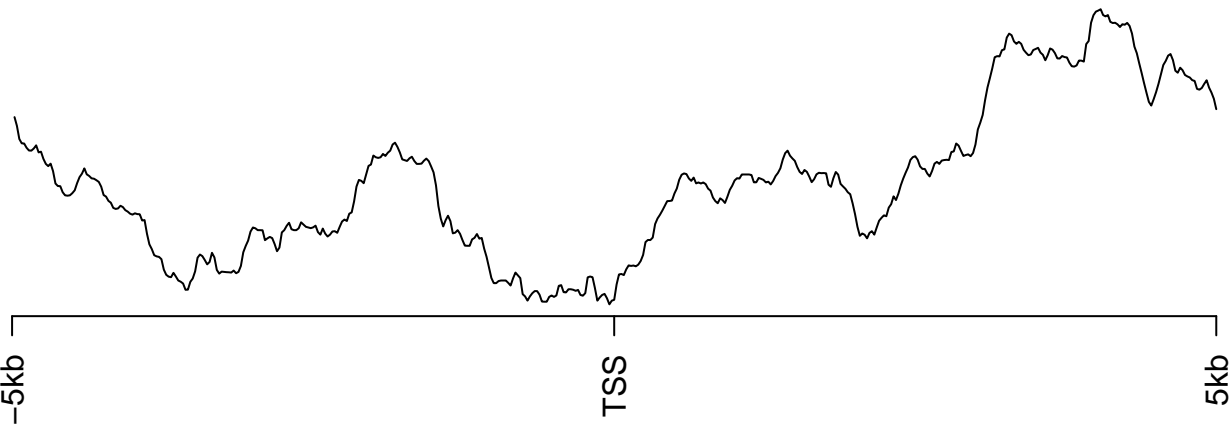
Average coverage per gene

0.0025 0.0035

-5kb

TSS

5kb



# K562 PositiveSingle E.678 scCLTdegenNuc956

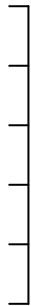
Average coverage per gene

0.0006 0.0012

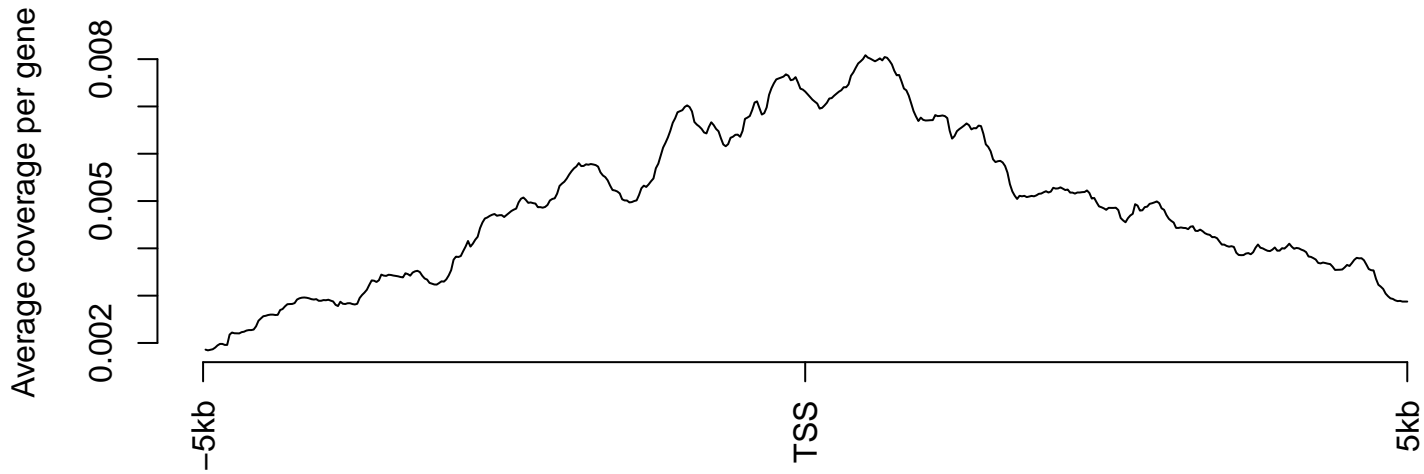
-5kb

TSS

5kb



# K562 PositiveSingle E.683 scCLTdegenNuc1004



# K562 PositiveSingle E.683 scCLTdegenNuc1005

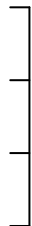
Average coverage per gene

0.0025 0.0035

-5kb

TSS

5kb



# K562 PositiveSingle E.683 scCLTdegenNuc1006

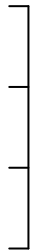
Average coverage per gene

0.0025 0.0035

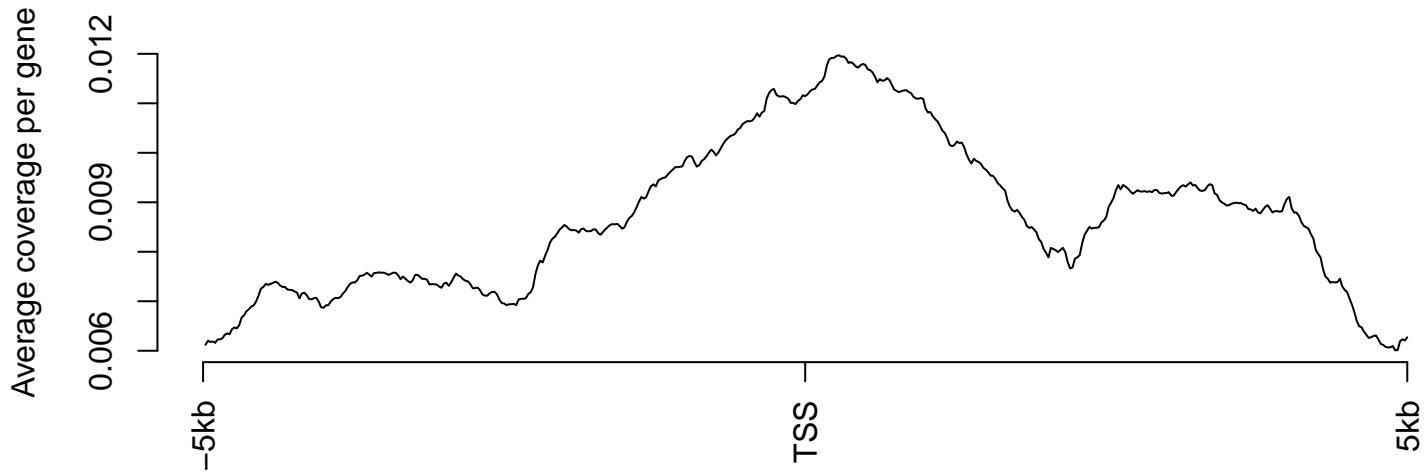
-5kb

TSS

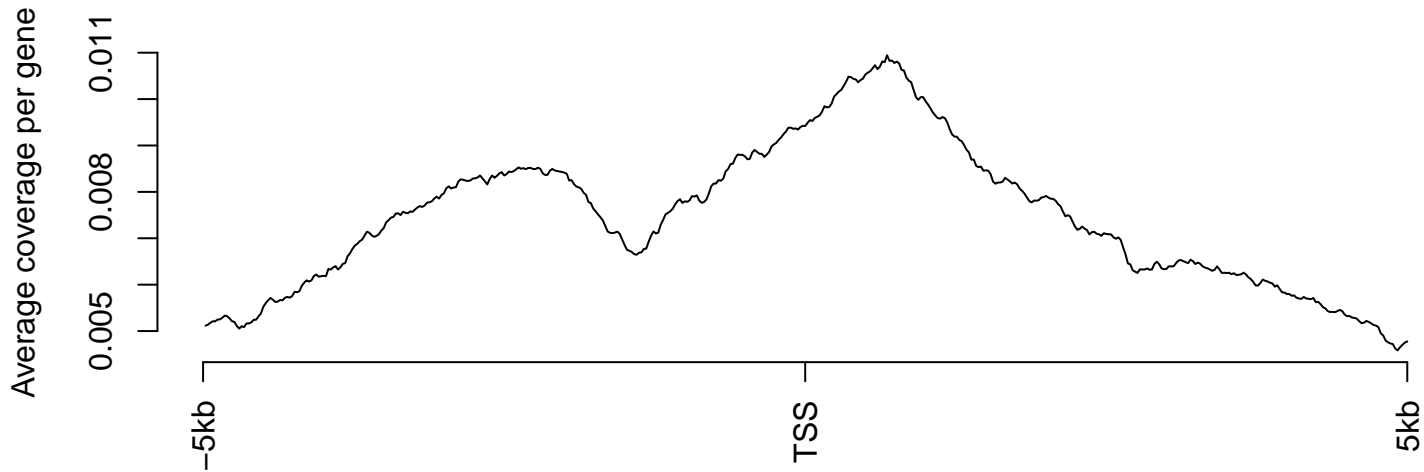
5kb



# K562 PositiveSingle E.683 scCLTdegenNuc1007



# K562 PositiveSingle E.683 scCLTdegenNuc1008



# K562 PositiveSingle E.683 scCLTdegenNuc1009

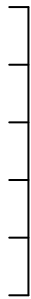
Average coverage per gene

0.0025 0.0040

-5kb

TSS

5kb





# K562 PositiveSingle E.683 scCLTdegenNuc1010

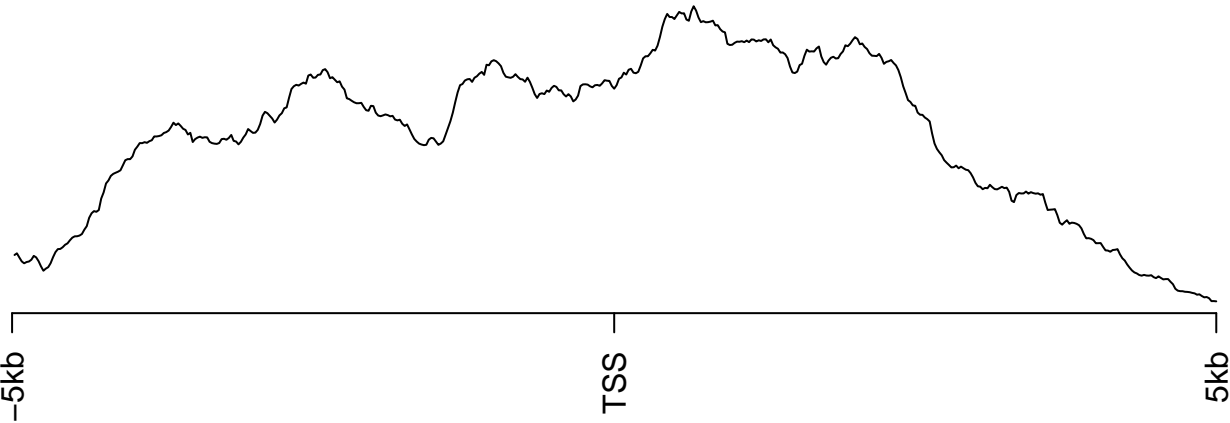
Average coverage per gene

0.004 0.007

-5kb

TSS

5kb



# K562 PositiveSingle E.683 scCLTdegenNuc1012

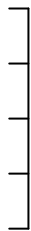
Average coverage per gene

0.004 0.006 0.008

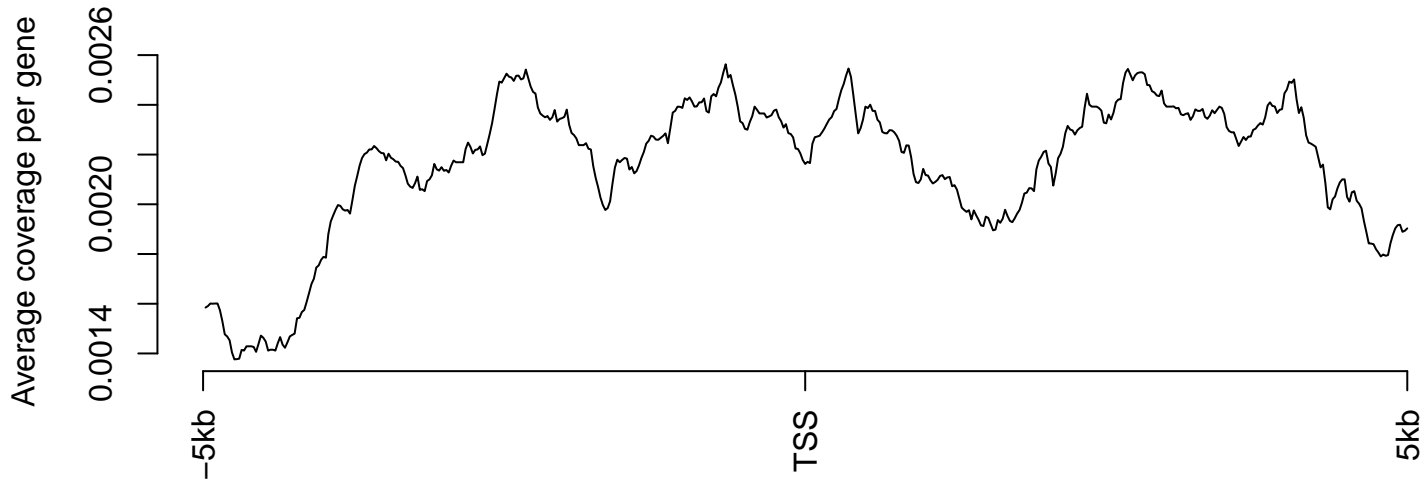
-5kb

TSS

5kb



# K562 PositiveSingle E.683 scCLTdegenNuc1013



# K562 PositiveSingle E.683 scCLTdegenNuc979

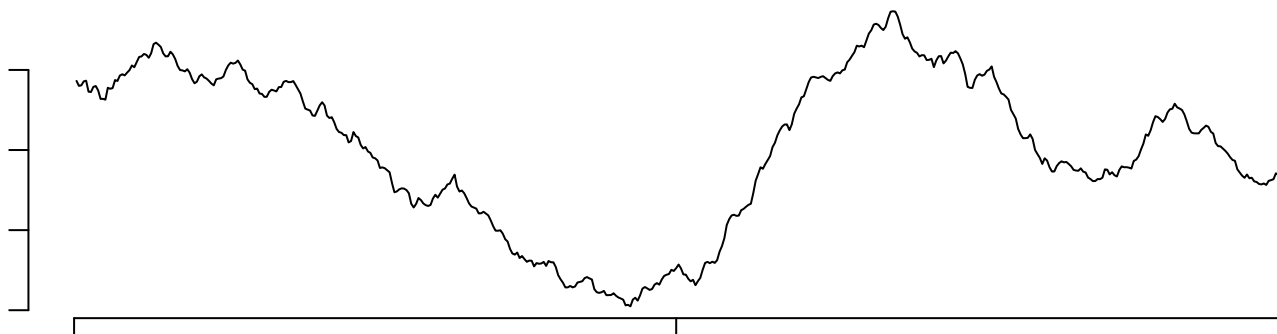
Average coverage per gene

0.007  
0.005

-5kb

TSS

5kb



# K562 PositiveSingle E.694 scCLTdegenNuc1077

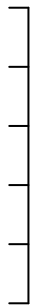
Average coverage per gene

1e-04  
3e-04  
5e-04

-5kb

TSS

5kb



# K562 PositiveSingle E.694 scCLTdegenNuc1078

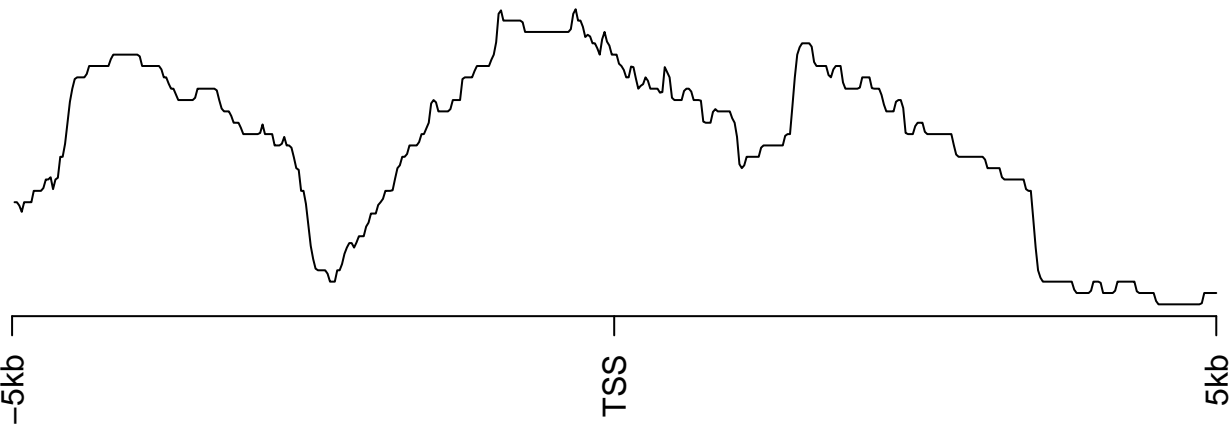
Average coverage per gene

1e-04 3e-04

-5kb

TSS

5kb



# K562 PositiveSingle E.694 scCLTdegenNuc1079

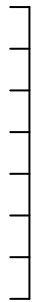
Average coverage per gene

0.0006 0.0012 0.0018

-5kb

TSS

5kb



# K562 PositiveSingle E.694 scCLTdegenNuc1080

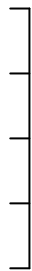
Average coverage per gene

0.0015 0.0025 0.0035

-5kb

TSS

5kb





# K562 PositivePooled E.574 scCLTdegenNuc333

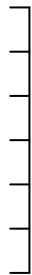
Average coverage per gene

0.018 0.024 0.030

-5kb

TSS

5kb



# K562 PositivePooled E.588 scCLTdegenNuc451

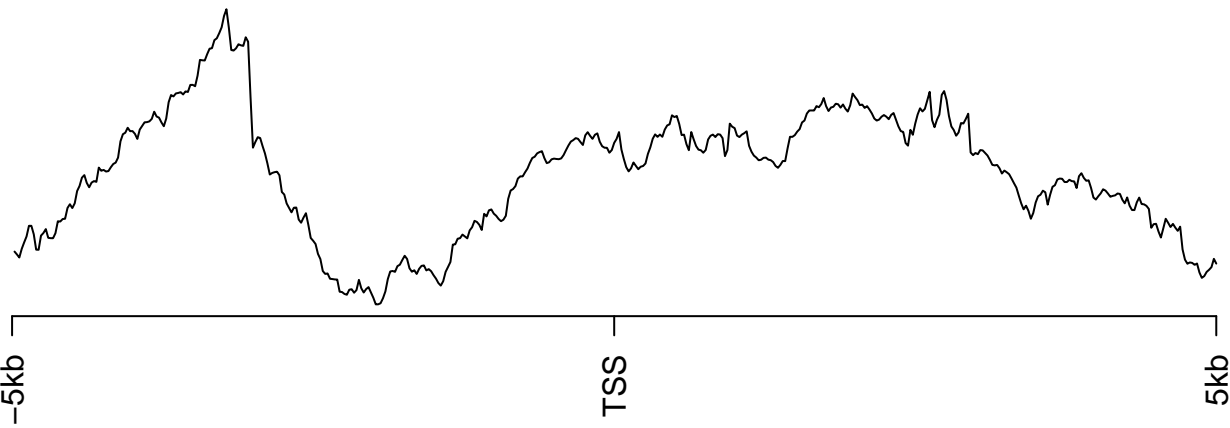
Average coverage per gene

0.018 0.021

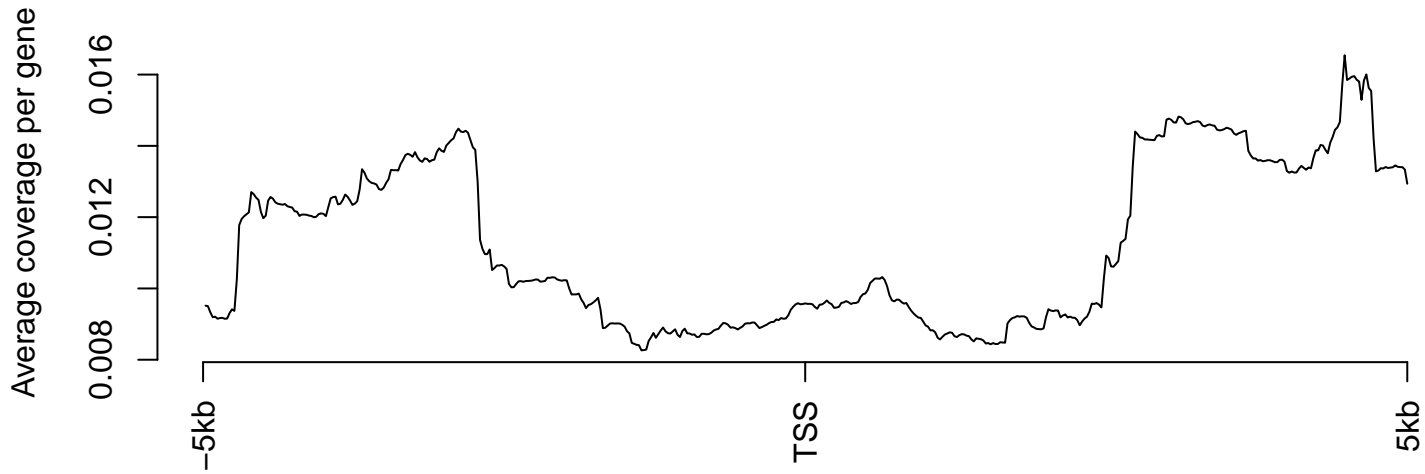
-5kb

TSS

5kb



# K562 PositivePooled E.588 scCLTdegenNuc457



# K562 PositivePooled E.652 scCLTdegenNuc656

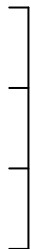
Average coverage per gene

0.06  
0.04

-5kb

TSS

5kb



# K562 PositivePooled E.652 scCLTdegenNuc657

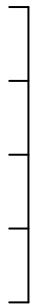
Average coverage per gene

0.09  
0.07  
0.05

-5kb

TSS

5kb



# K562 PositivePooled E.683 scCLTdegenNuc1011

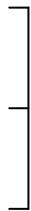
Average coverage per gene

0.10 0.15 0.20

-5kb

TSS

5kb



# K562 PositiveMerged E.– K562PositiveMerged

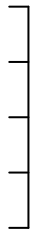
Average coverage per gene

0.15 0.17 0.19

-5kb

TSS

5kb



# K562 PositiveAll E.- K562PositiveAll

Average coverage per gene

0.35 0.45 0.55

-5kb

TSS

5kb

