

K562 NegCtrlSingle E.571 scCLTdegenNuc383

Average coverage per gene

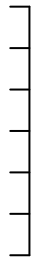
0.00005 0.00025

-5kb

Gene start

Gene end

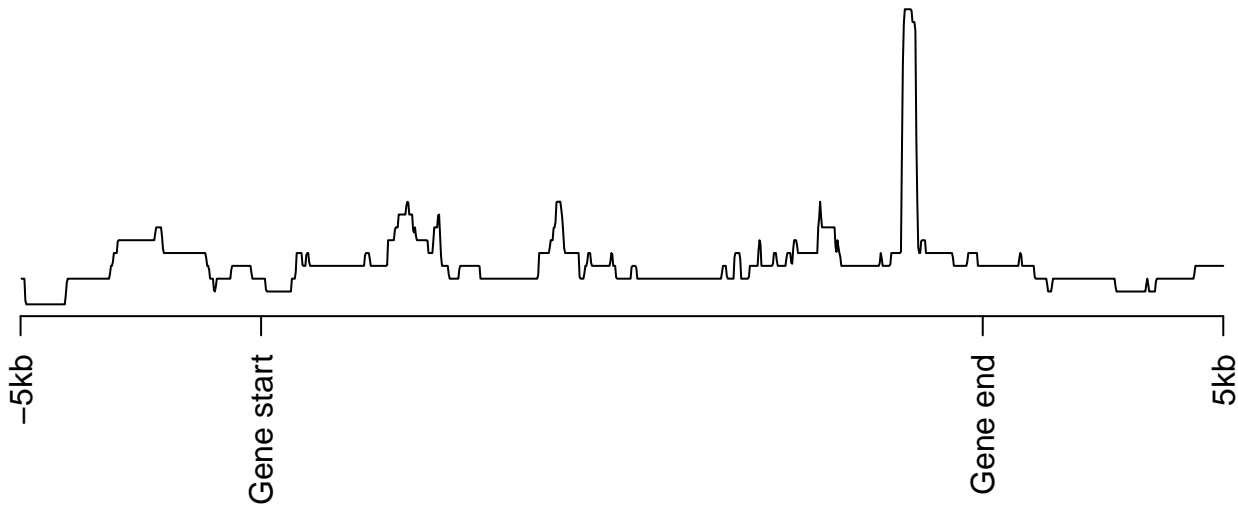
5kb



K562 NegCtrlSingle E.571 scCLTdegenNuc384

Average coverage per gene

0e+00 3e-04



K562 NegCtrlSingle E.571 scCLTdegenNuc386

Average coverage per gene

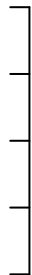
0e+00 2e-04 4e-04

-5kb

Gene start

Gene end

5kb



K562 NegCtrlSingle E.579 scCLTdegenNuc417

Average coverage per gene

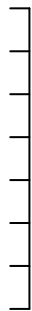
0e+00 3e-04 6e-04

-5kb

Gene start

Gene end

5kb



K562 NegCtrlSingle E.579 scCLTdegenNuc418

Average coverage per gene

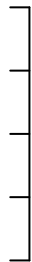
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

5kb



K562 NegCtrlSingle E.579 scCLTdegenNuc420

Average coverage per gene

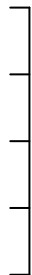
2e-04 6e-04 1e-03

-5kb

Gene start

Gene end

5kb



K562 NegCtrlSingle E.579 scCLTdegenNuc422

Average coverage per gene

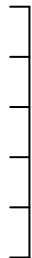
1e-04
4e-04

-5kb

Gene start

Gene end

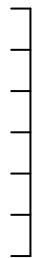
5kb



K562 NegCtrlSingle E.579 scCLTdegenNuc423

Average coverage per gene

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

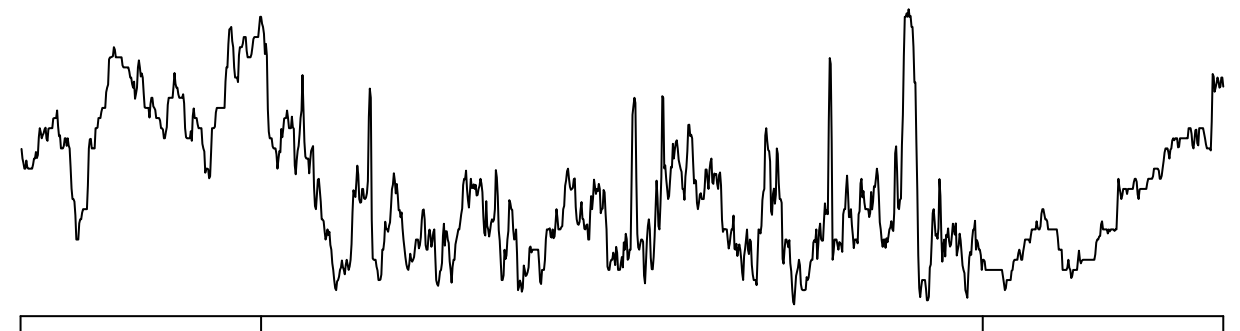


-5kb

Gene start

Gene end

5kb



K562 NegCtrlSingle E.579 scCLTdegenNuc424

Average coverage per gene

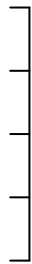
1e-04 3e-04 5e-04

-5kb

Gene start

Gene end

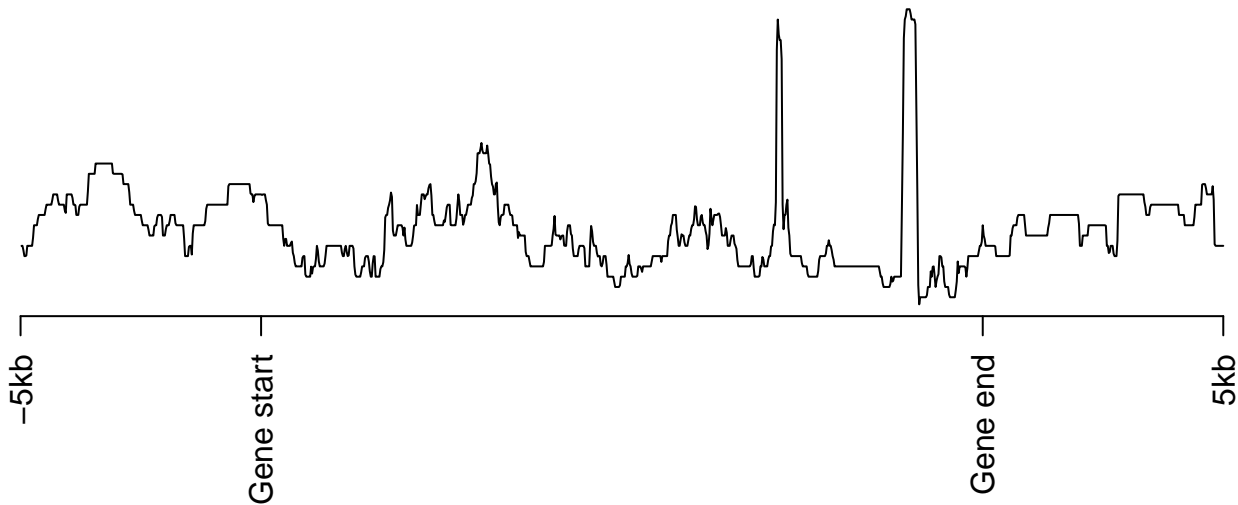
5kb



K562 NegCtrlSingle E.579 scCLTdegenNuc425

Average coverage per gene

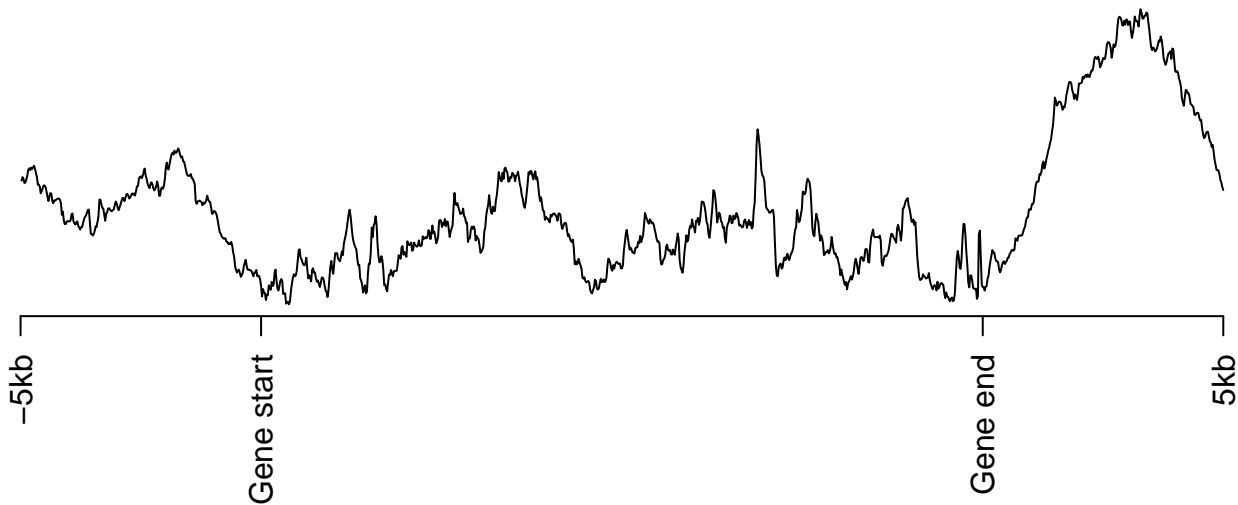
0e+00 3e-04 6e-04



K562 NegCtrlSingle E.671 scCLTdegenNuc897

Average coverage per gene

0.0035
0.0020



K562 NegCtrlSingle E.671 scCLTdegenNuc898

Average coverage per gene

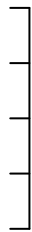
0.0020
0.0005

-5kb

Gene start

Gene end

5kb



K562 NegCtrlSingle E.671 scCLTdegenNuc900

Average coverage per gene

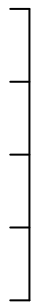
0e+00 4e-04 8e-04

-5kb

Gene start

Gene end

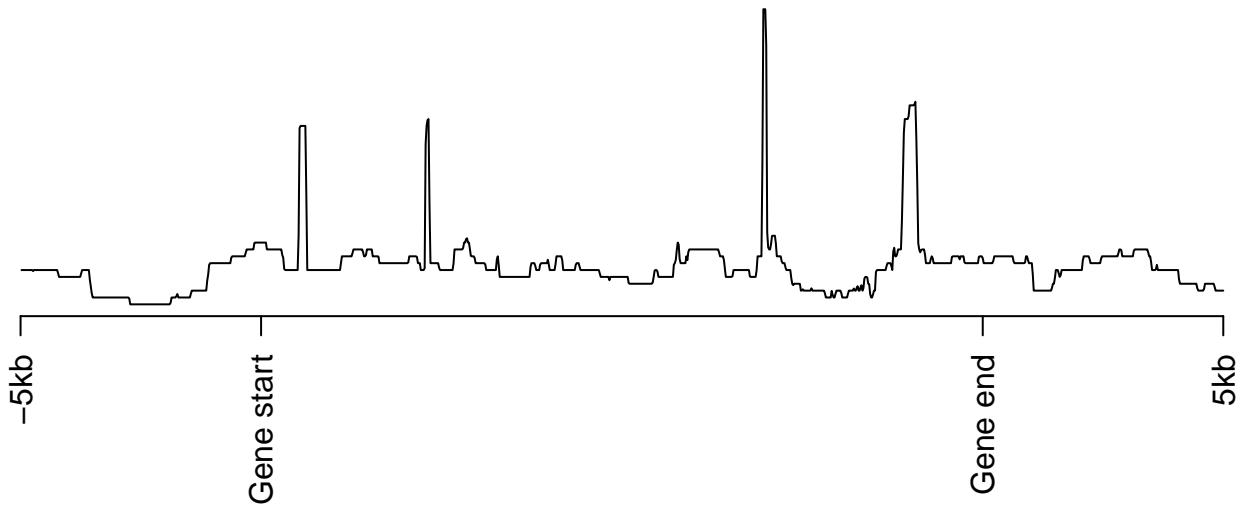
5kb



K562 NegCtrlSingle E.671 scCLTdegenNuc902

Average coverage per gene

0e+00 6e-04



K562 NegCtrlSingle E.671 scCLTdegenNuc903

Average coverage per gene

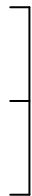
0.0020
0.0010

-5kb

Gene start

Gene end

5kb



K562 NegCtrlSingle E.676 scCLTdegenNuc950

Average coverage per gene

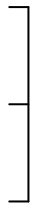
0.0015
0.0005

-5kb

Gene start

Gene end

5kb



K562 NegCtrlSingle E.678 scCLTdegenNuc957

Average coverage per gene

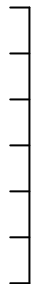
0.0005 0.0020 0.0035

-5kb

Gene start

Gene end

5kb



K562 NegCtrlPooled E.591 scCLTdegenNuc463

Average coverage per gene

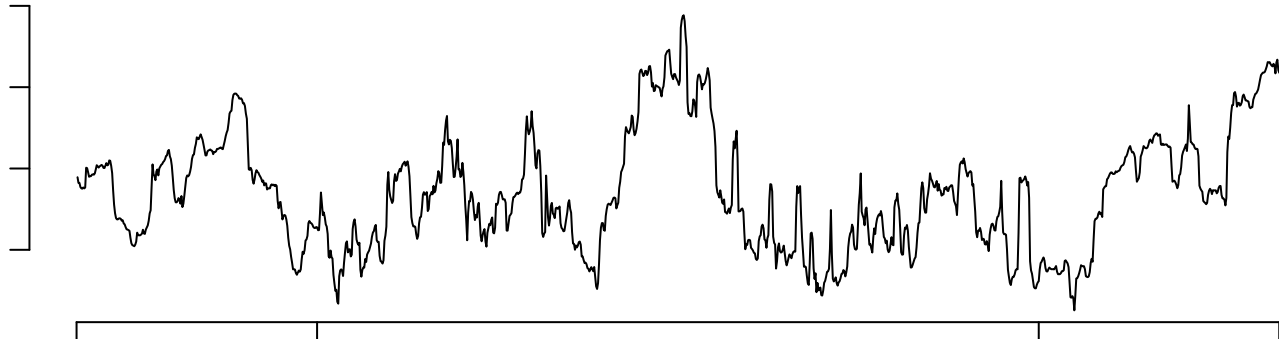
0.010
0.006

-5kb

Gene start

Gene end

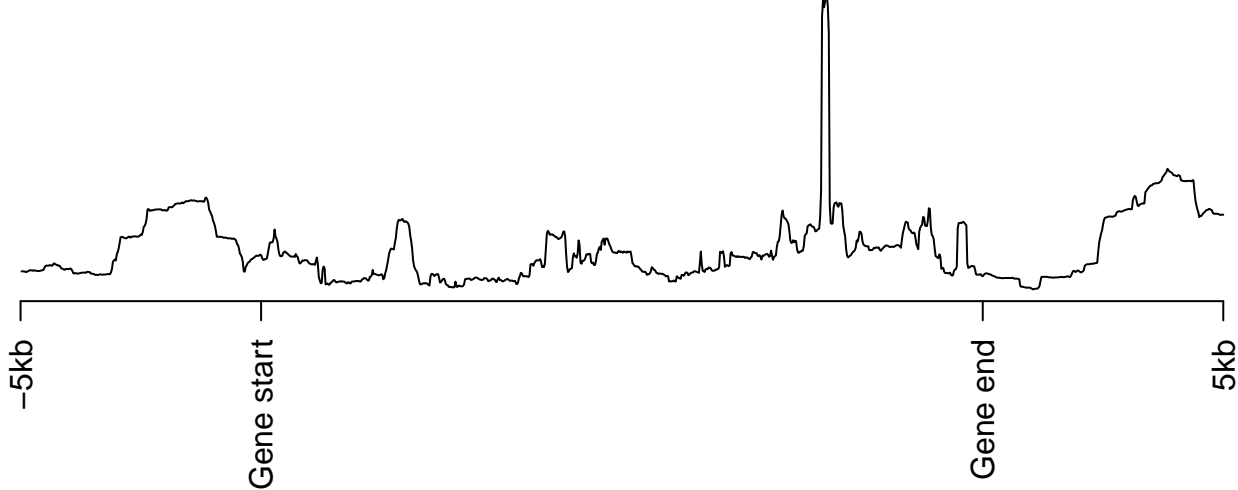
5kb



K562 NegCtrlPooled E.591 scCLTdegenNuc464

Average coverage per gene

0.005 0.015 0.025



K562 NegCtrlMerged E.- K562NegCtrlMerged

Average coverage per gene

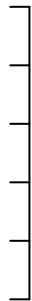
0.006 0.010 0.014

-5kb

Gene start

Gene end

5kb



K562 NegCtrlAll E.- K562NegCtrlAll

Average coverage per gene

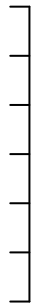
0.015 0.030 0.045

-5kb

Gene start

Gene end

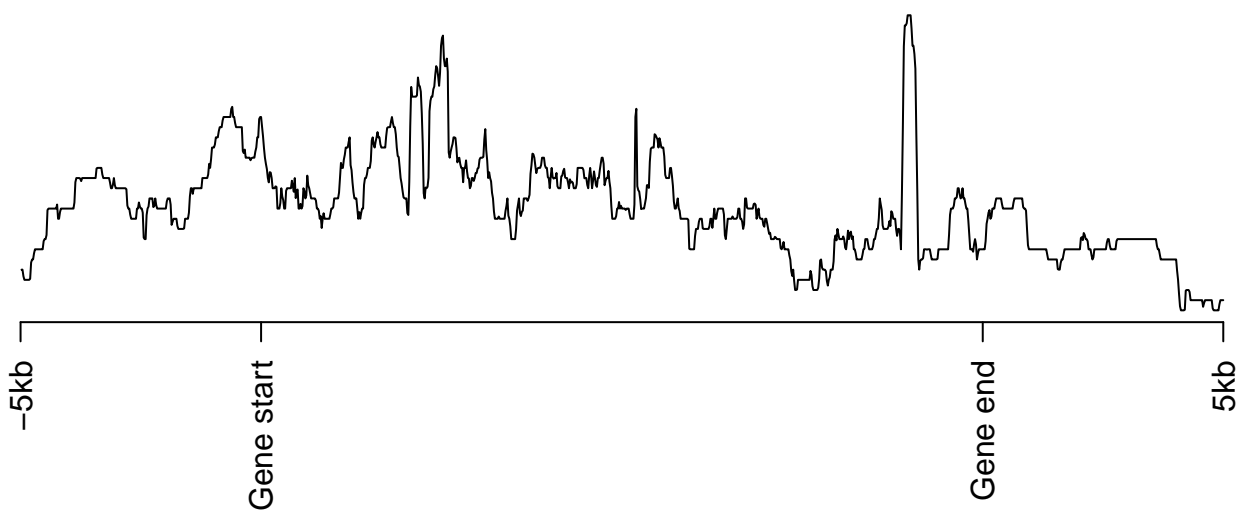
5kb



K562 PositiveSingle E.571 scCLTdegenNuc404

Average coverage per gene

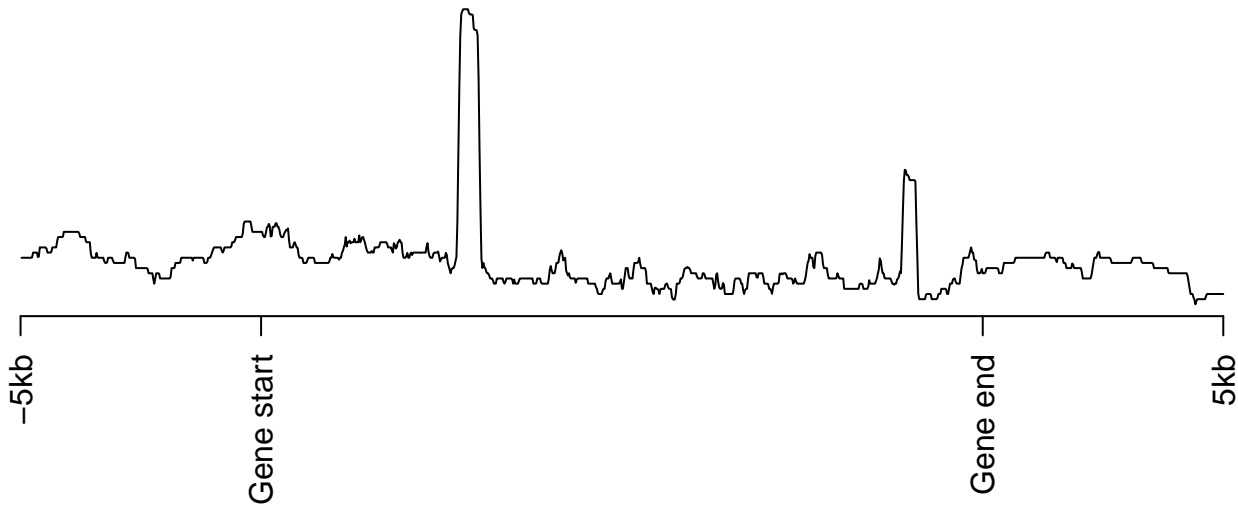
1e-04
4e-04
7e-04



K562 PositiveSingle E.571 scCLTdegenNuc406

Average coverage per gene

0.0000 0.0006 0.0012



K562 PositiveSingle E.571 scCLTdegenNuc408

Average coverage per gene

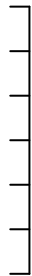
2e-04 5e-04 8e-04

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.571 scCLTdegenNuc409

Average coverage per gene

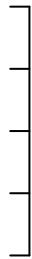
0.0045
0.0030

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.571 scCLTdegenNuc410

Average coverage per gene

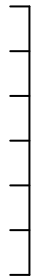
0.0006 0.0012 0.0018

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.571 scCLTdegenNuc411

Average coverage per gene

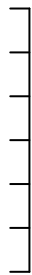
0.0002 0.0008 0.0014

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.574 scCLTdegenNuc334

Average coverage per gene

0.0035 0.0045 0.0055

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.574 scCLTdegenNuc335

Average coverage per gene

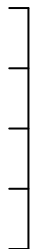
0.0030
0.0045

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.574 scCLTdegenNuc336

Average coverage per gene

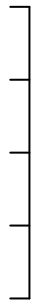
0.008
0.006
0.004

-5kb

Gene start

Gene end

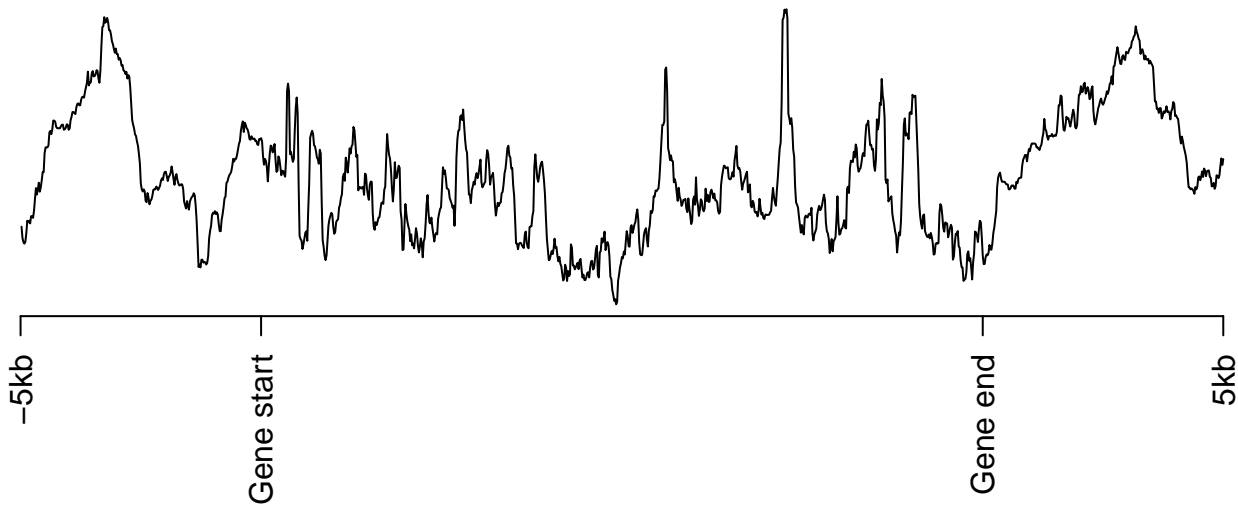
5kb



K562 PositiveSingle E.579 scCLTdegenNuc337

Average coverage per gene

0.0030
0.0020



K562 PositiveSingle E.579 scCLTdegenNuc338

Average coverage per gene

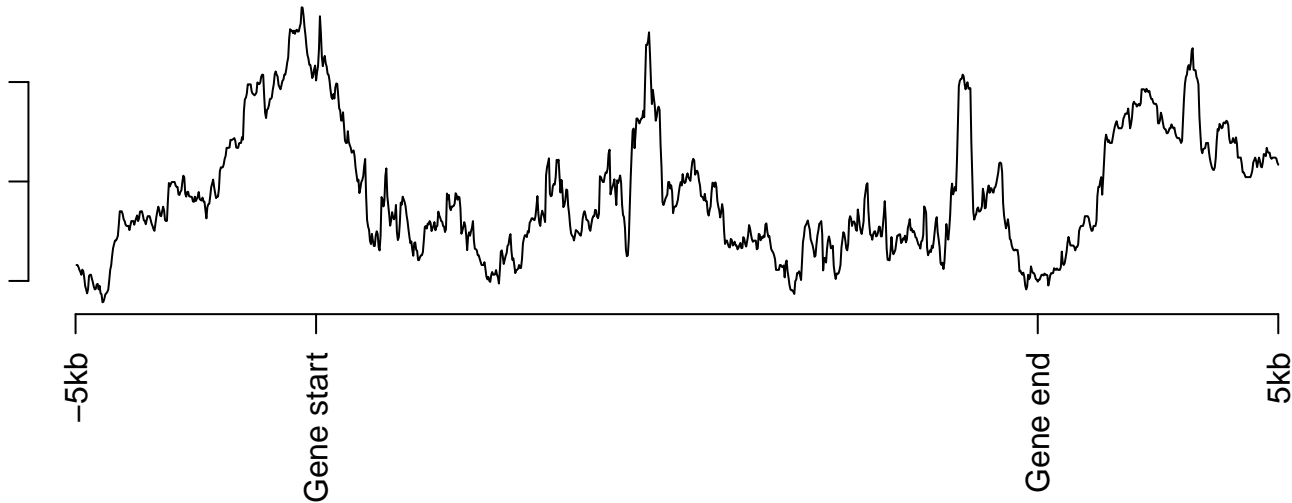
0.0020
0.0010

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.579 scCLTdegenNuc339

Average coverage per gene

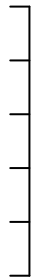
0.0040
0.0025

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.579 scCLTdegenNuc340

Average coverage per gene

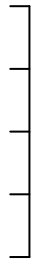
0.0025 0.0035 0.0045

-5kb

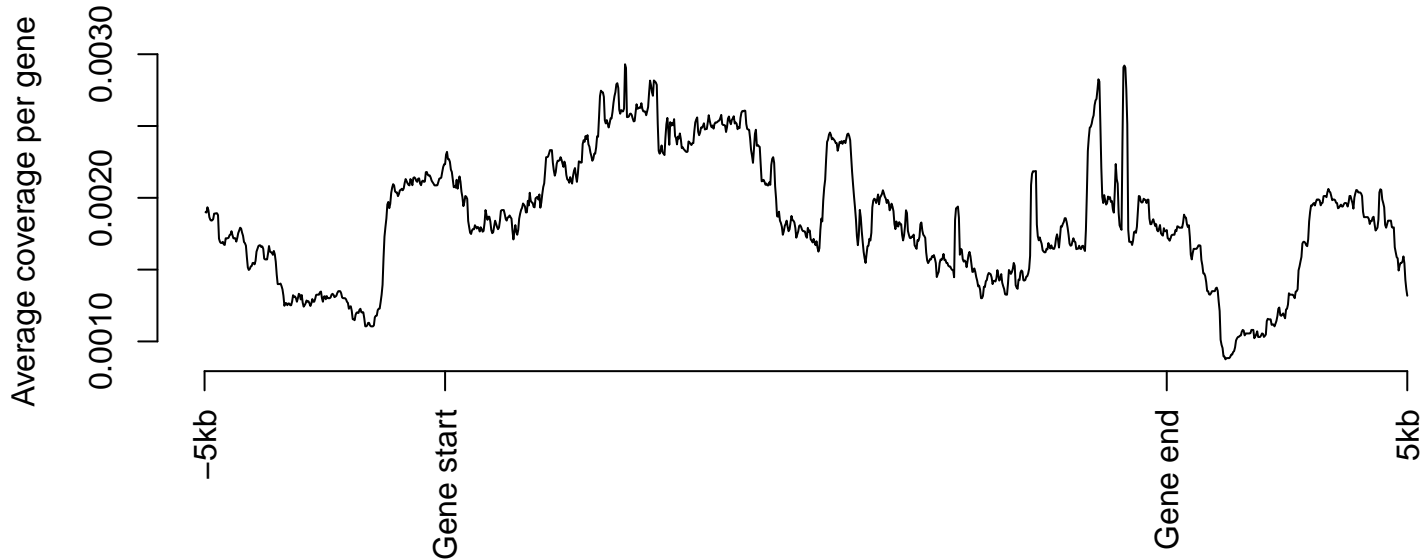
Gene start

Gene end

5kb



K562 PositiveSingle E.579 scCLTdegenNuc341



K562 PositiveSingle E.580 scCLTdegenNuc342

Average coverage per gene

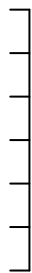
0.0020
0.0014
0.0008

-5kb

Gene start

Gene end

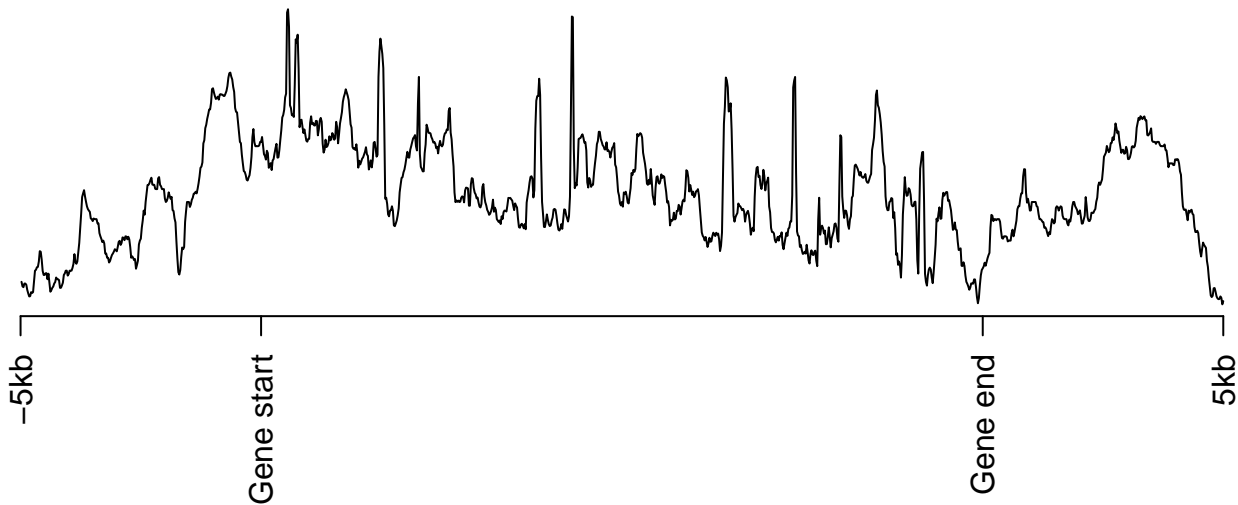
5kb



K562 PositiveSingle E.580 scCLTdegenNuc343

Average coverage per gene

0.0015 0.0025



K562 PositiveSingle E.580 scCLTdegenNuc345

Average coverage per gene

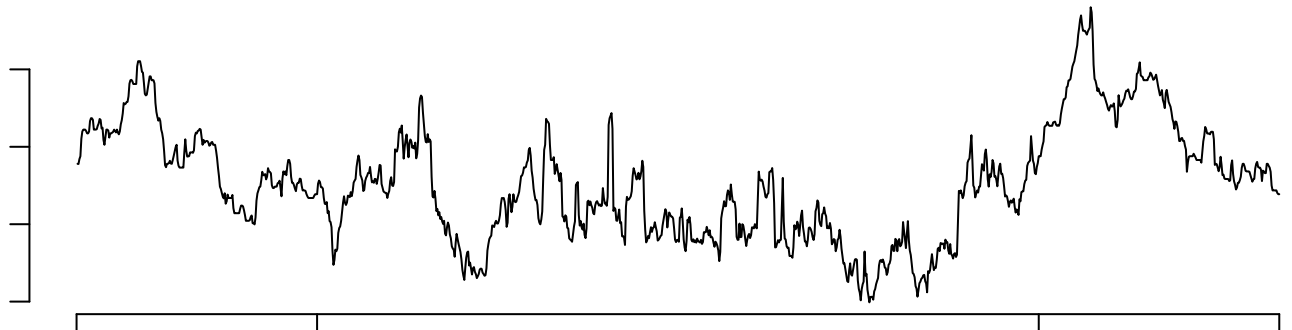
0.0010 0.0020

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.580 scCLTdegenNuc428

Average coverage per gene

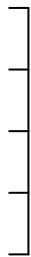
0.018
0.014
0.010

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.588 scCLTdegenNuc323

Average coverage per gene

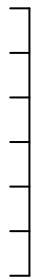
0.0018
0.0012
0.0006

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.588 scCLTdegenNuc324

Average coverage per gene

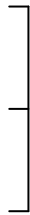
0.0015
0.0005

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.588 scCLTdegenNuc325

Average coverage per gene

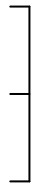
0.0015
0.0005

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.588 scCLTdegenNuc446

Average coverage per gene

0.004
0.002

-5kb

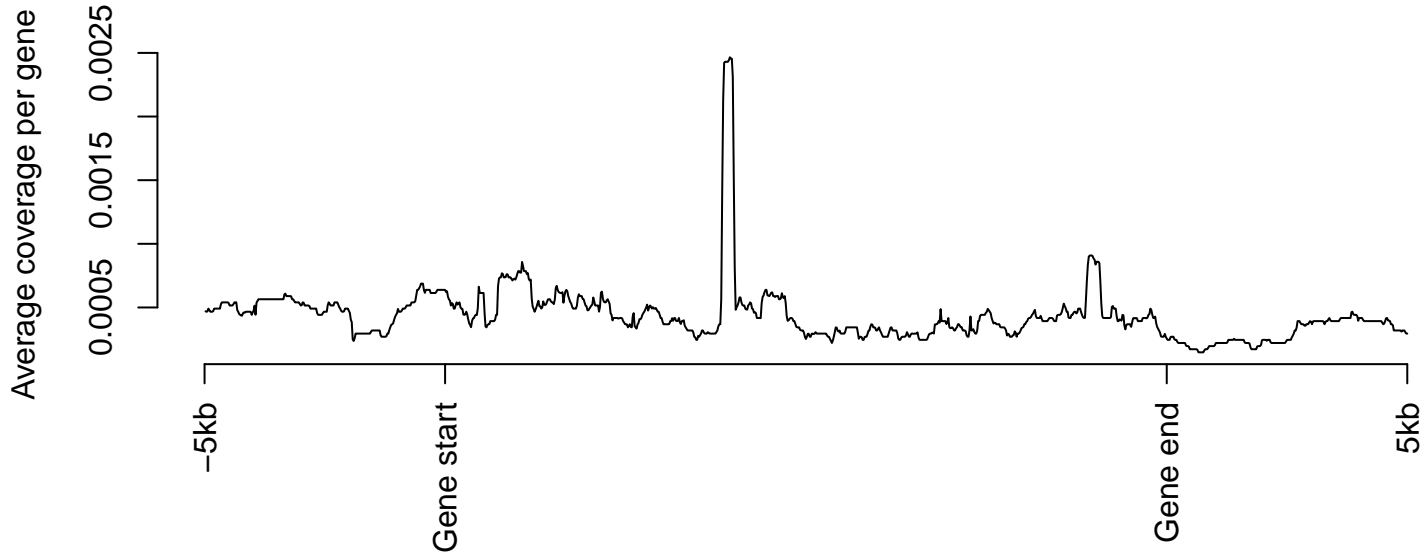
Gene start

Gene end

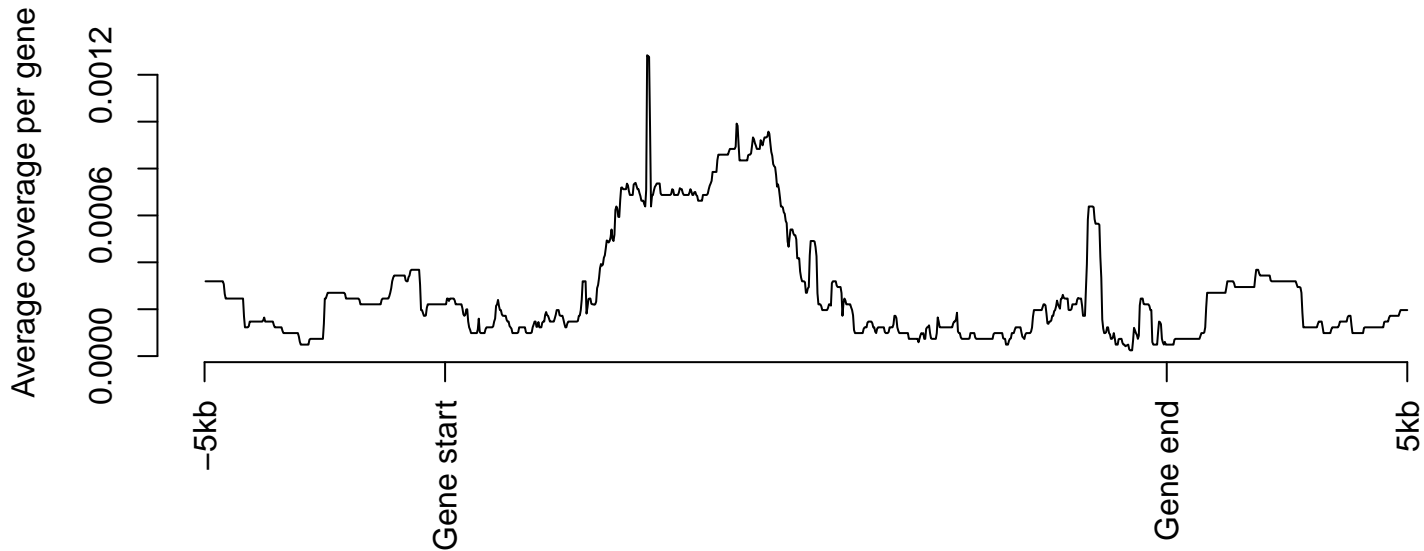
5kb



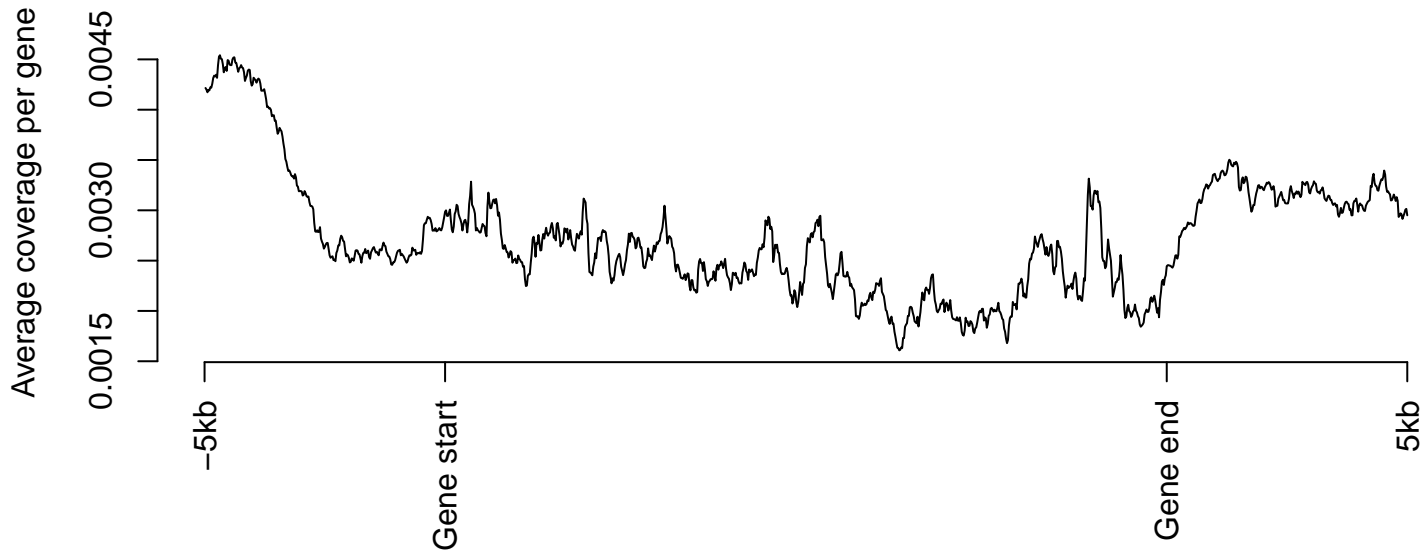
K562 PositiveSingle E.588 scCLTdegenNuc447



K562 PositiveSingle E.588 scCLTdegenNuc448



K562 PositiveSingle E.588 scCLTdegenNuc452



K562 PositiveSingle E.588 scCLTdegenNuc454

Average coverage per gene

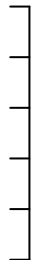
0.0006 0.0012

-5kb

Gene start

Gene end

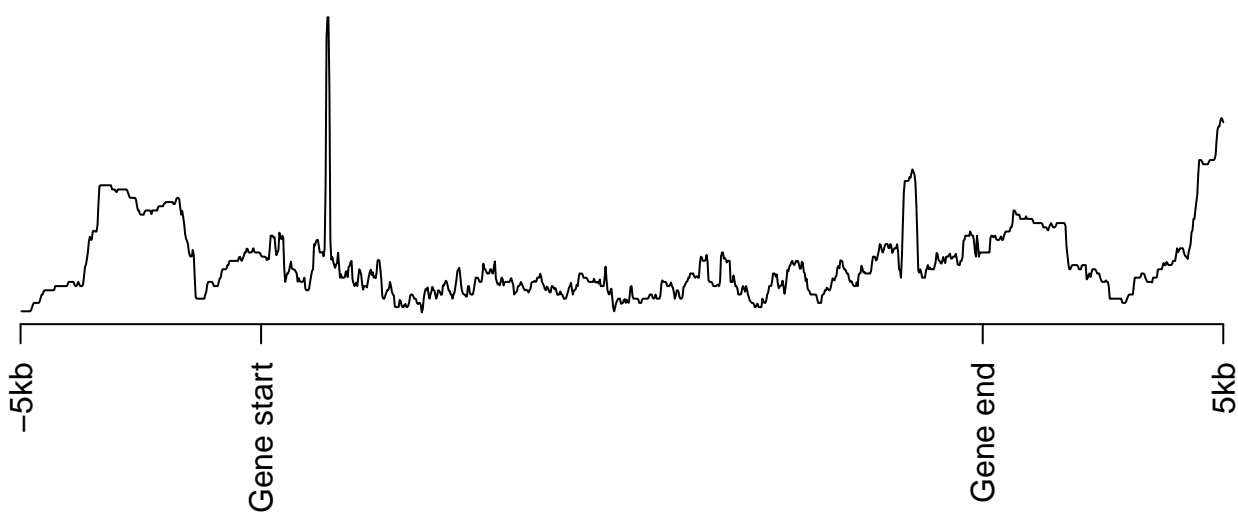
5kb



K562 PositiveSingle E.588 scCLTdegenNuc455

Average coverage per gene

0.0005
0.0015



K562 PositiveSingle E.588 scCLTdegenNuc456

Average coverage per gene

0.0015 0.0025

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.589 scCLTdegenNuc326

Average coverage per gene

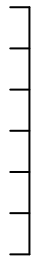
0.0018
0.0010

-5kb

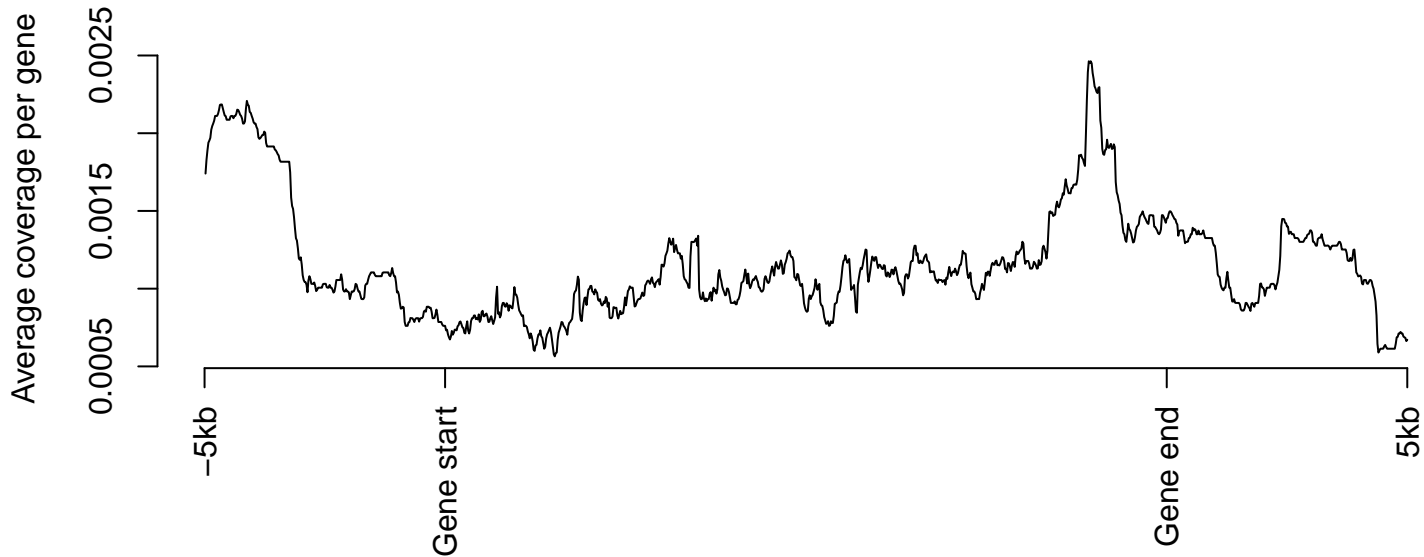
Gene start

Gene end

5kb



K562 PositiveSingle E.589 scCLTdegenNuc348



K562 PositiveSingle E.665 scCLTdegenNuc766

Average coverage per gene

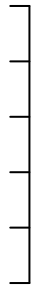
0.0035
0.0020

-5kb

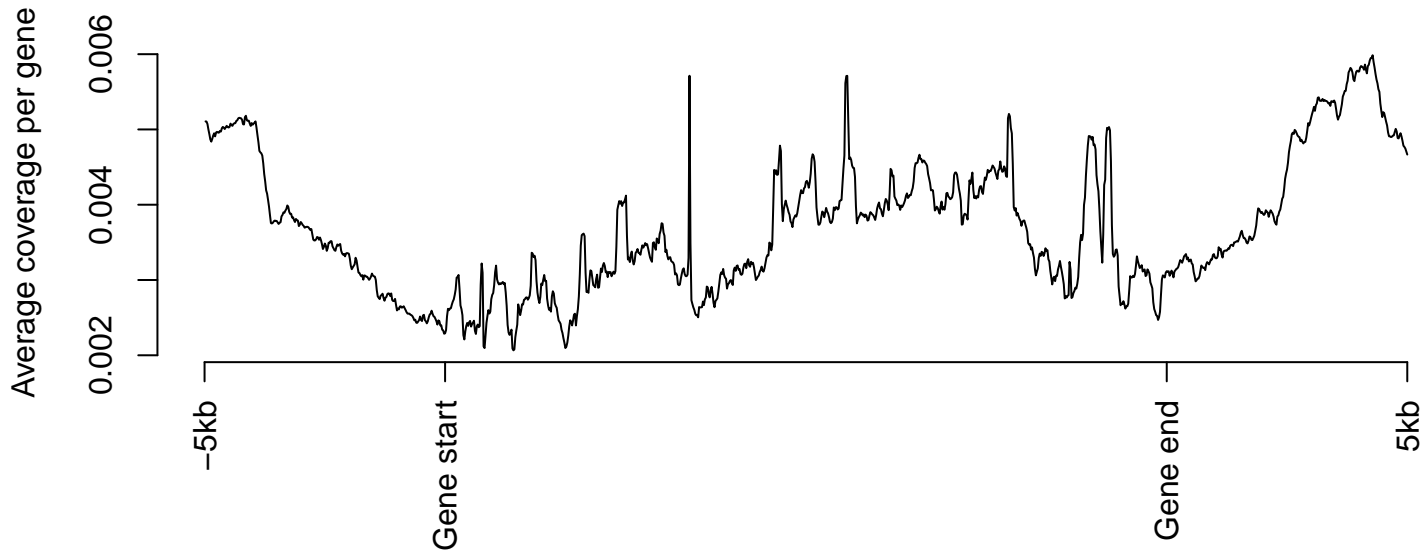
Gene start

Gene end

5kb



K562 PositiveSingle E.665 scCLTdegenNuc767



K562 PositiveSingle E.665 scCLTdegenNuc768

Average coverage per gene

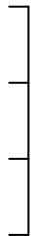
0.0015
0.0005

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.665 scCLTdegenNuc769

Average coverage per gene

0.0020
0.0010

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.665 scCLTdegenNuc770

Average coverage per gene

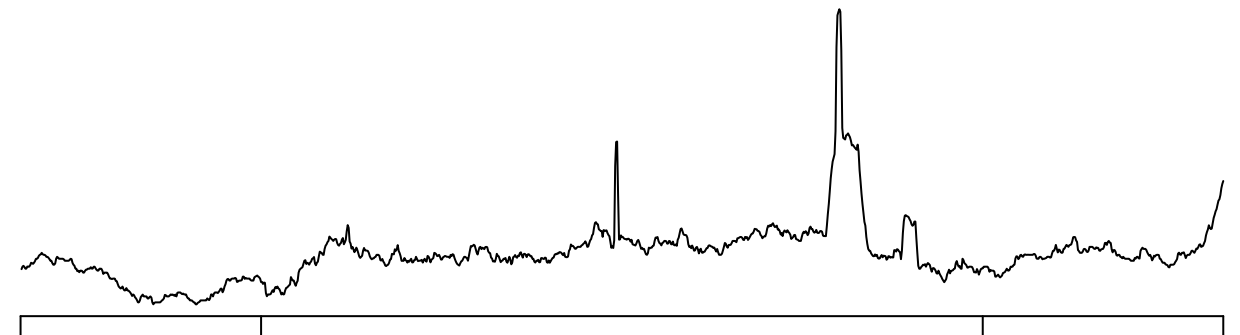
0.003
0.001

-5kb

Gene start

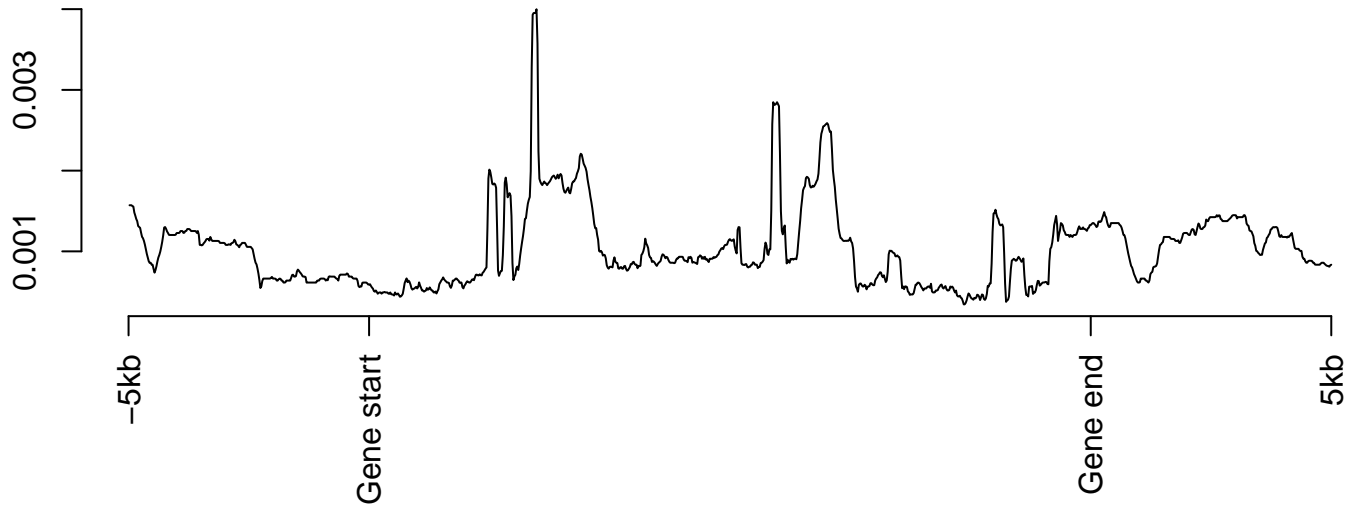
Gene end

5kb



K562 PositiveSingle E.665 scCLTdegenNuc771

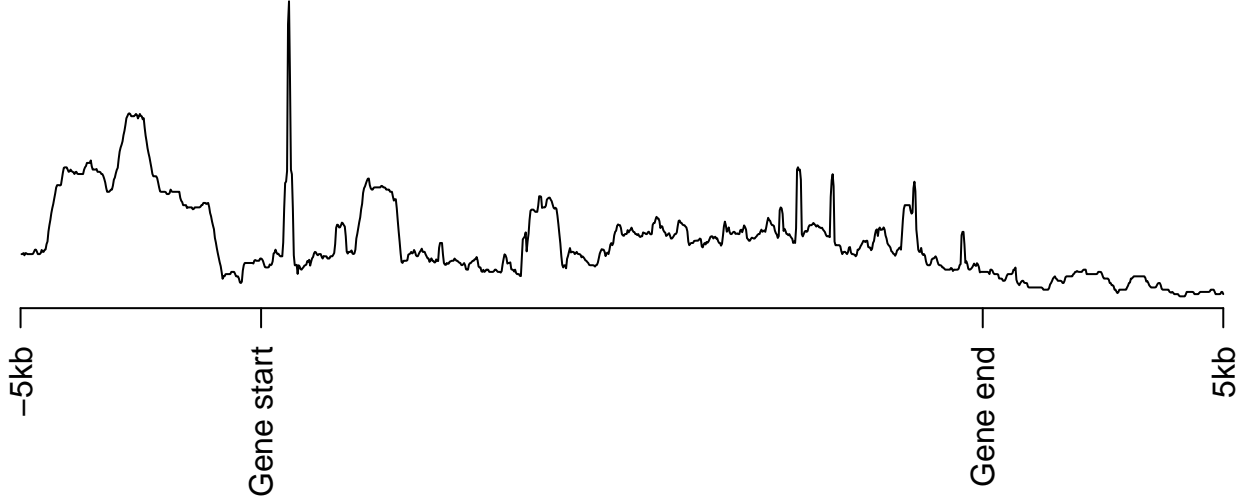
Average coverage per gene



K562 PositiveSingle E.665 scCLTdegenNuc772

Average coverage per gene

0.0035
0.0020
0.0005



K562 PositiveSingle E.665 scCLTdegenNuc773

Average coverage per gene

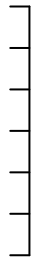
0.0010
0.0002

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.665 scCLTdegenNuc774

Average coverage per gene

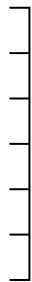
0.0002 0.0008 0.0014

-5kb

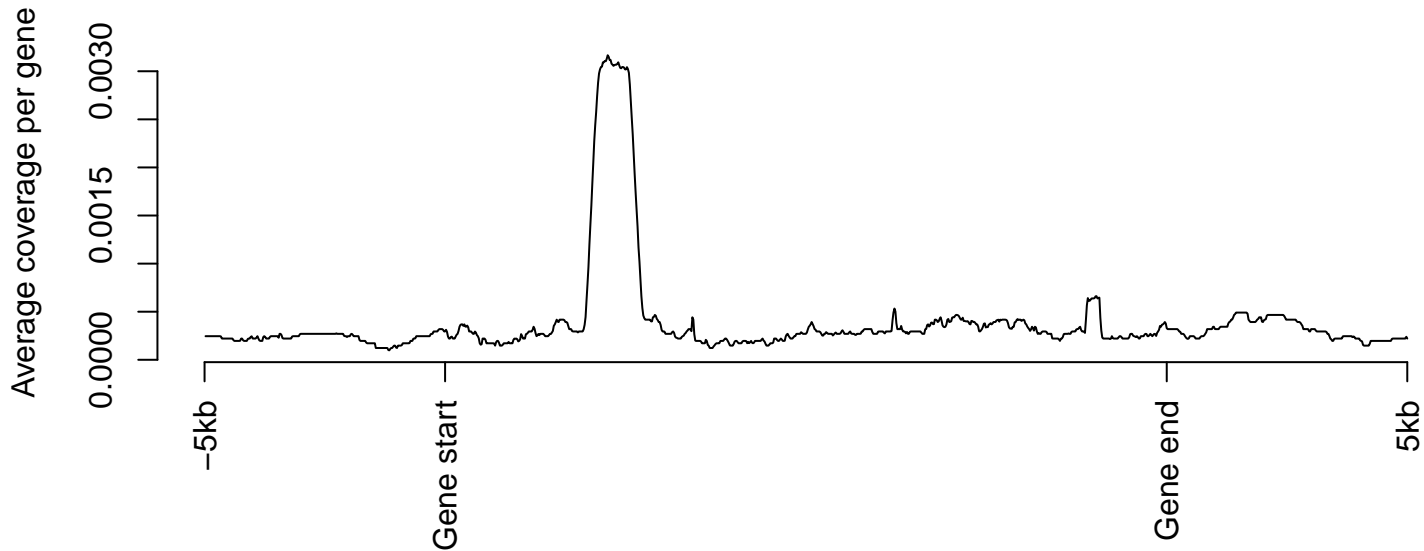
Gene start

Gene end

5kb



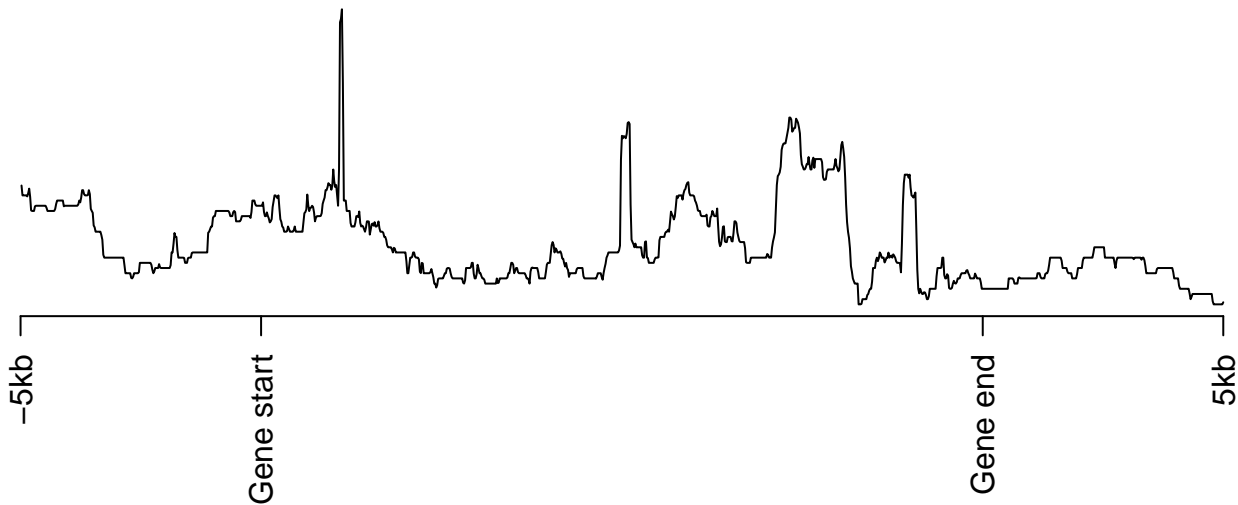
K562 PositiveSingle E.665 scCLTdegenNuc775



K562 PositiveSingle E.673 scCLTdegenNuc906_b

Average coverage per gene

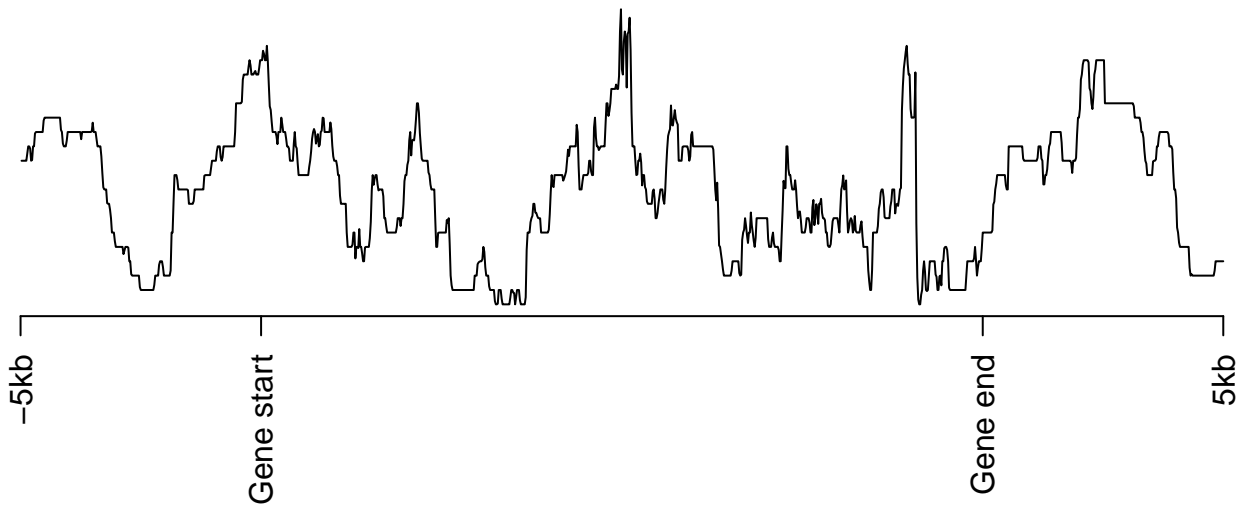
0.0014
0.0008
0.0002



K562 PositiveSingle E.673 scCLTdegenNuc907_b

Average coverage per gene

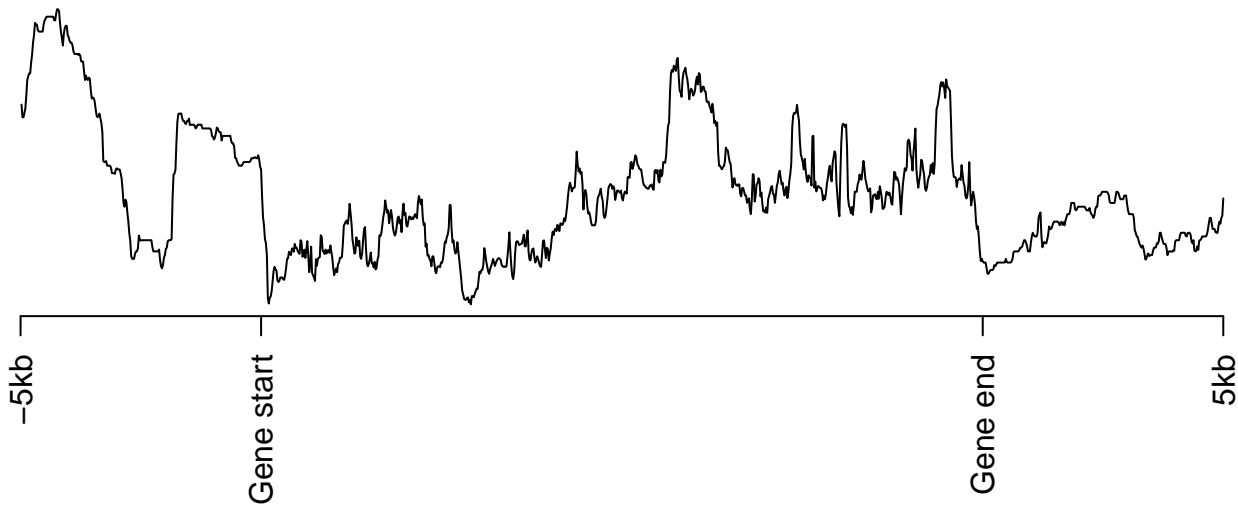
1e-04 3e-04 5e-04



K562 PositiveSingle E.673 scCLTdegenNuc908_b

Average coverage per gene

0.0005 0.0015



K562 PositiveSingle E.675 scCLTdegenNuc933

Average coverage per gene

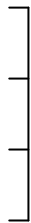
0.0010 0.0020

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.675 scCLTdegenNuc935

Average coverage per gene

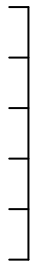
0.0005 0.0020

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.675 scCLTdegenNuc936

Average coverage per gene

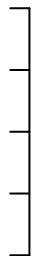
0.001 0.003 0.005

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.675 scCLTdegenNuc938

Average coverage per gene

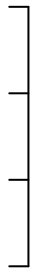
0.0005
0.0015

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.675 scCLTdegenNuc942

Average coverage per gene

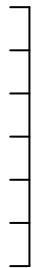
0.0010 0.0025 0.0040

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.675 scCLTdegenNuc943

Average coverage per gene

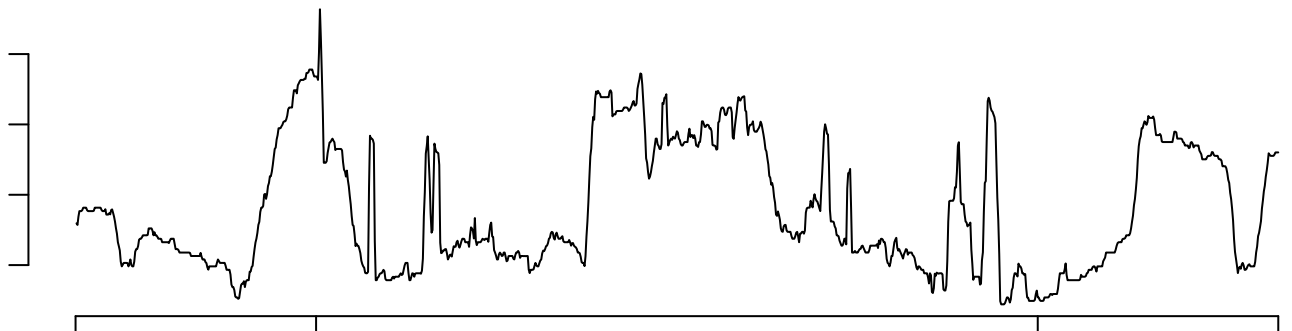
0.0005 0.0015

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.676 scCLTdegenNuc944

Average coverage per gene

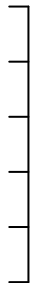
0.0008
0.0002

-5kb

Gene start

Gene end

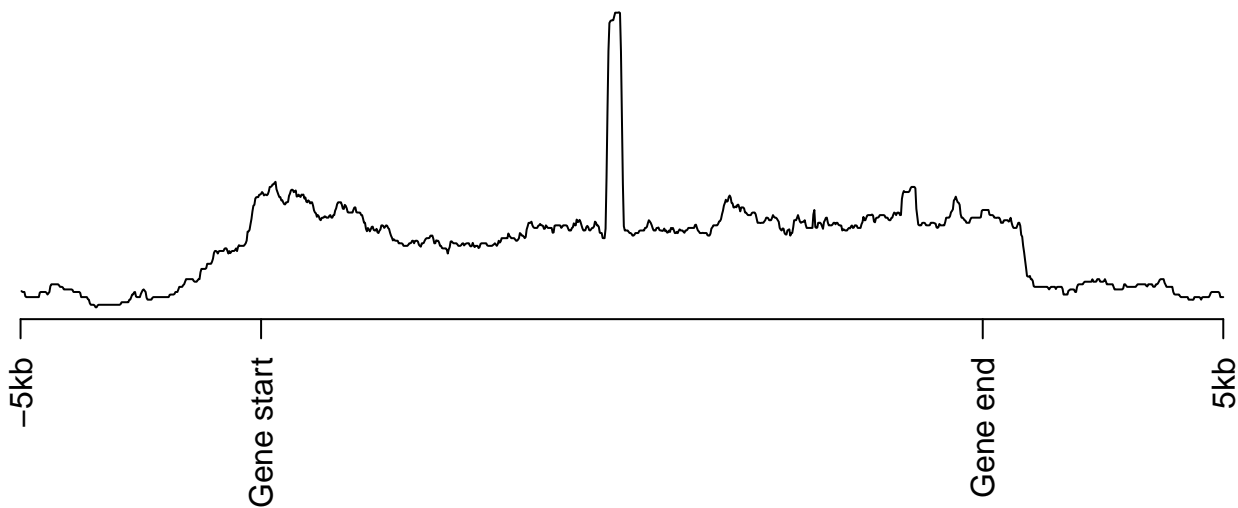
5kb



K562 PositiveSingle E.676 scCLTdegenNuc945

Average coverage per gene

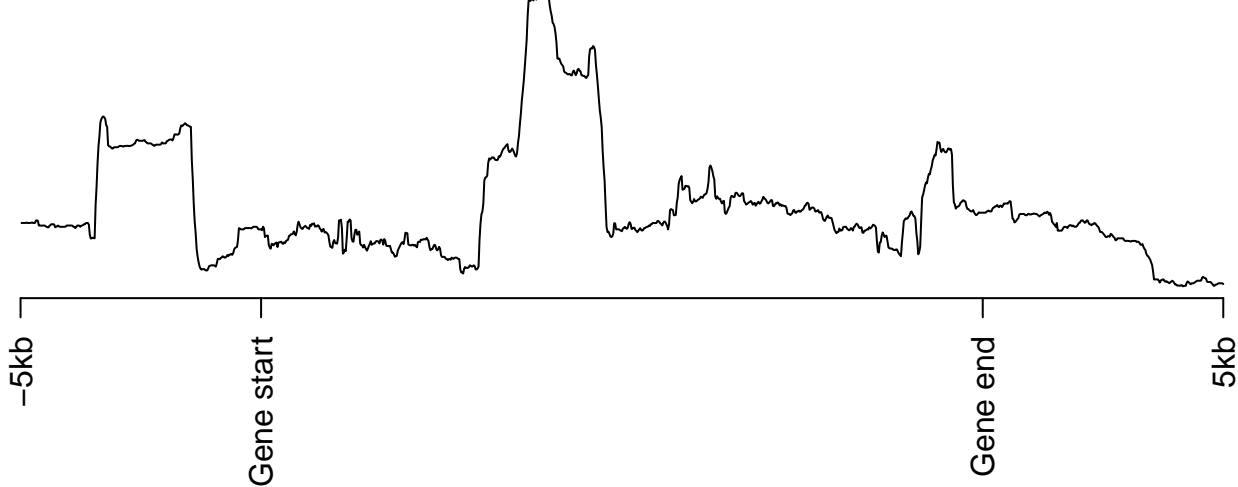
0.0020
0.0005



K562 PositiveSingle E.676 scCLTdegenNuc946

Average coverage per gene

0.001
0.004



K562 PositiveSingle E.676 scCLTdegenNuc947

Average coverage per gene

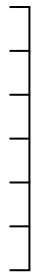
0.0016
0.0010
0.0004

-5kb

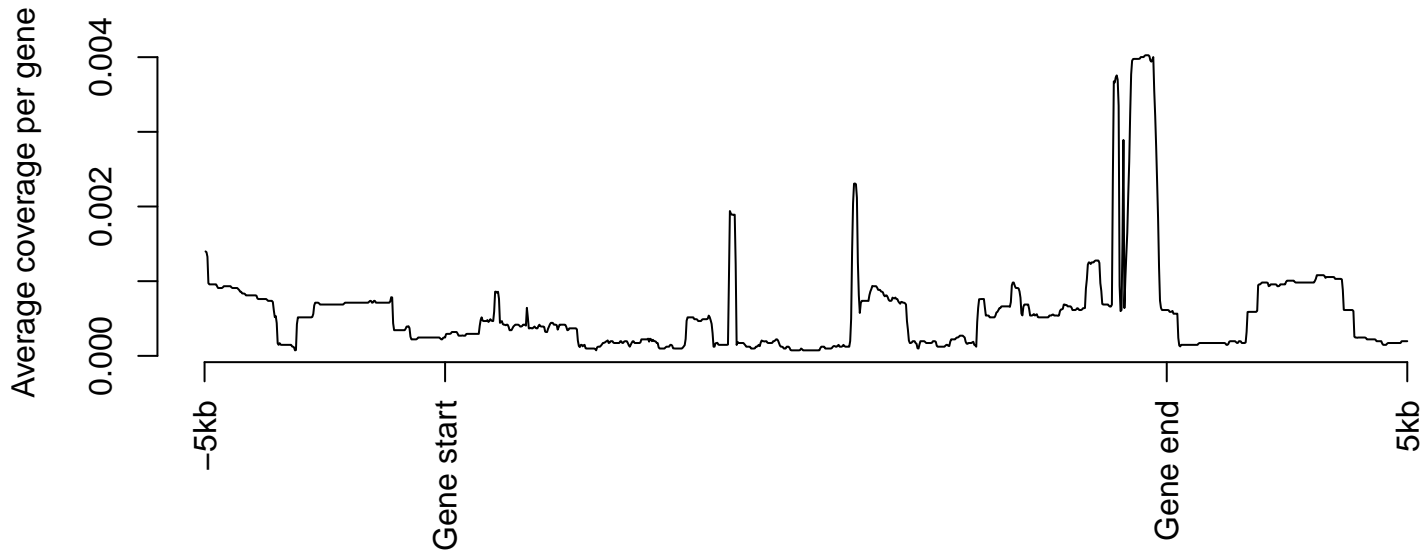
Gene start

Gene end

5kb



K562 PositiveSingle E.676 scCLTdegenNuc948



K562 PositiveSingle E.676 scCLTdegenNuc949

Average coverage per gene

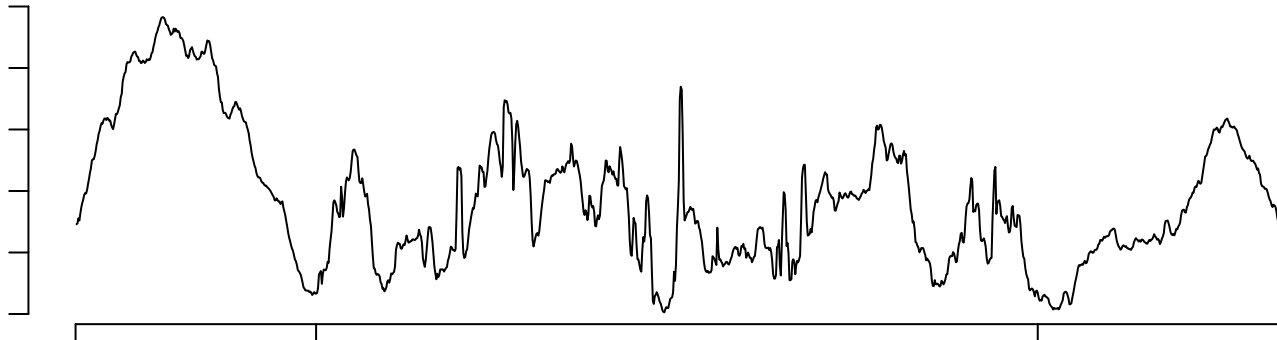
0.006
0.004
0.002

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.678 scCLTdegenNuc951

Average coverage per gene

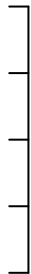
0.0005 0.0015 0.0025

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.678 scCLTdegenNuc952

Average coverage per gene

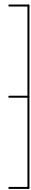
0.0015
0.0005

-5kb

Gene start

Gene end

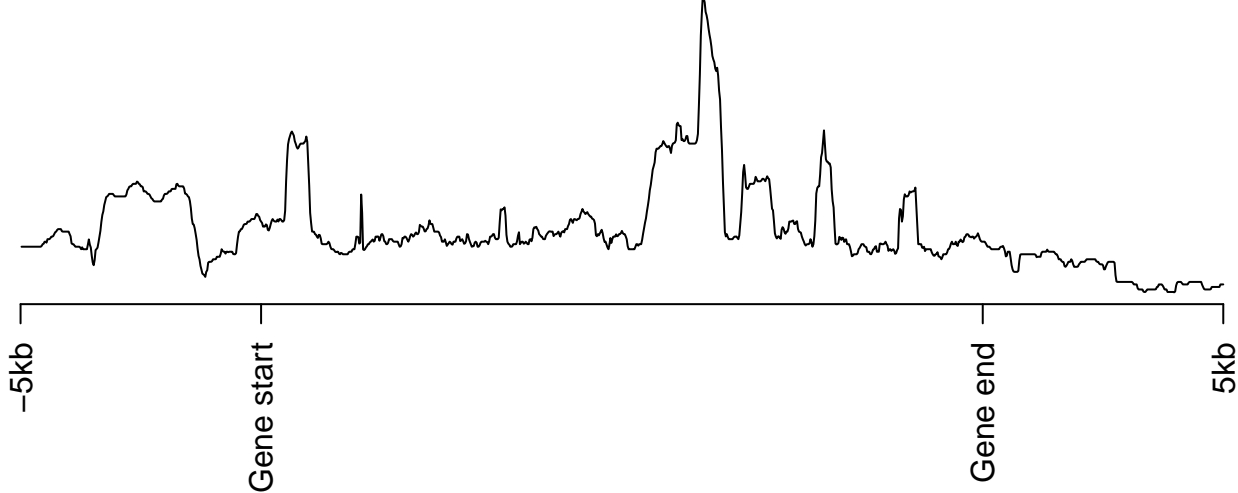
5kb



K562 PositiveSingle E.678 scCLTdegenNuc953

Average coverage per gene

0.0005 0.0020



K562 PositiveSingle E.678 scCLTdegenNuc954

Average coverage per gene

0.003
0.001

-5kb

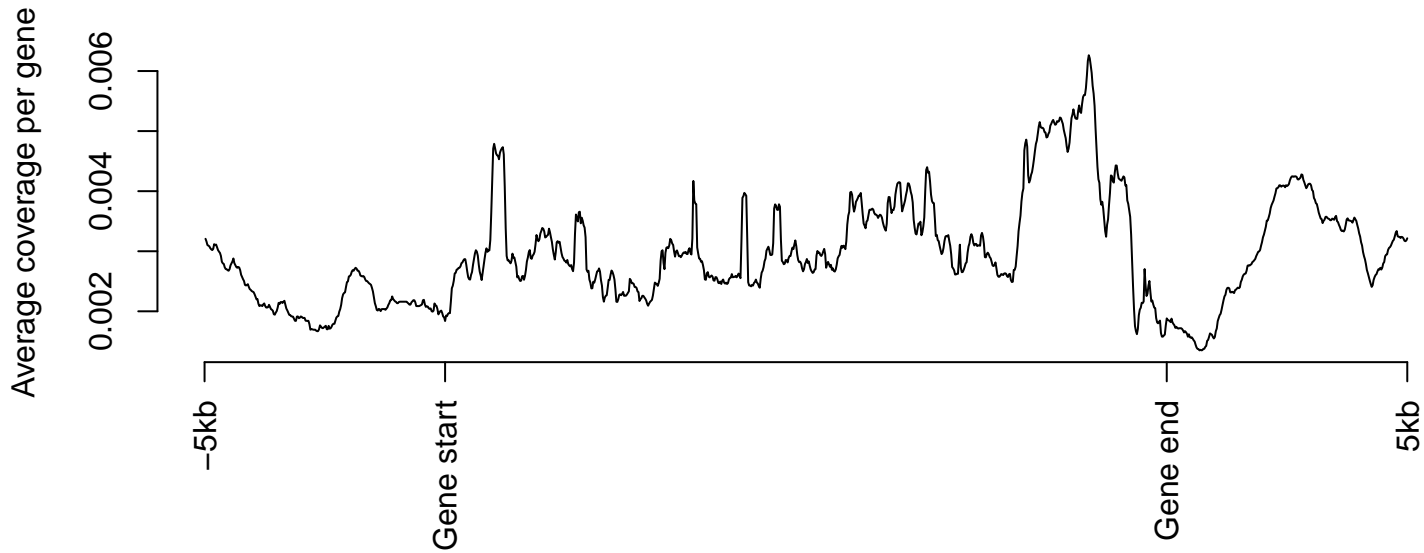
Gene start

Gene end

5kb



K562 PositiveSingle E.678 scCLTdegenNuc955



K562 PositiveSingle E.678 scCLTdegenNuc956

Average coverage per gene

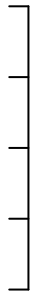
0.0025
0.0015
0.0005

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.683 scCLTdegenNuc1004

Average coverage per gene

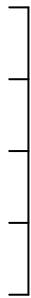
0.005
0.003
0.001

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.683 scCLTdegenNuc1005

Average coverage per gene

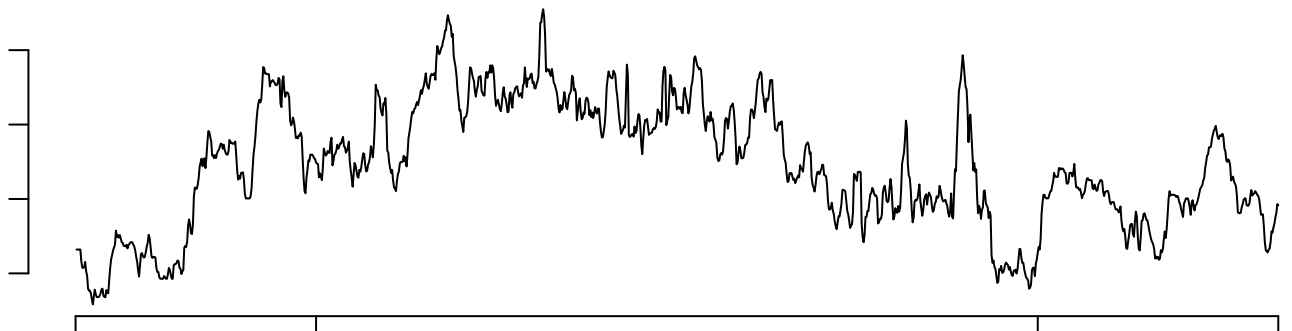
0.0020 0.0030

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.683 scCLTdegenNuc1006

Average coverage per gene

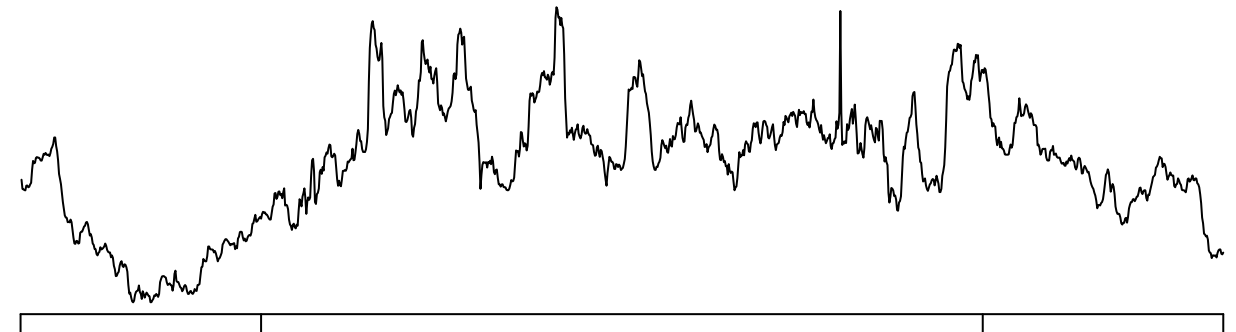
0.0035
0.0020

-5kb

Gene start

Gene end

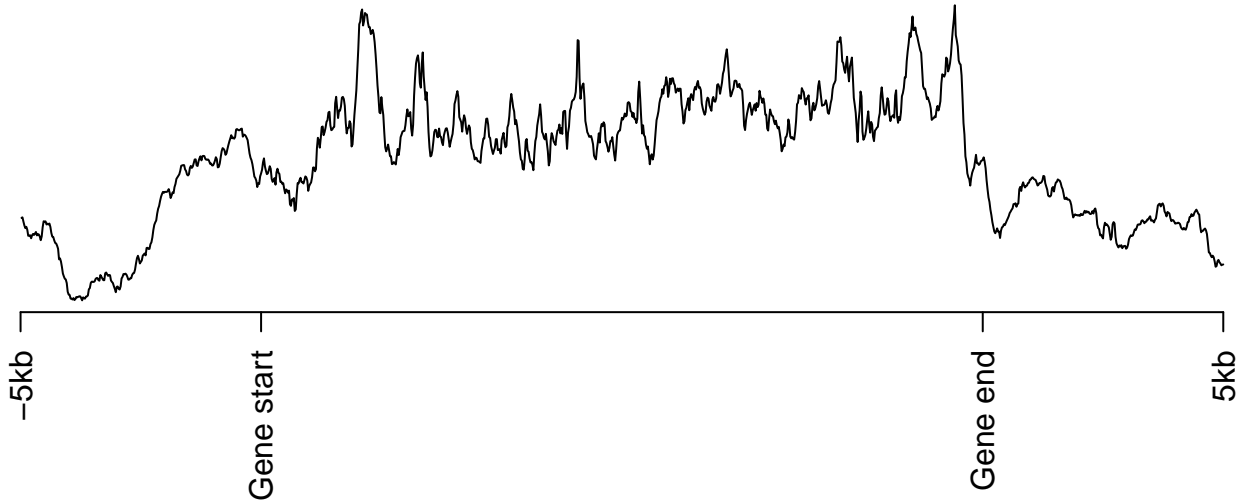
5kb



K562 PositiveSingle E.683 scCLTdegenNuc1007

Average coverage per gene

0.005 0.007 0.009



K562 PositiveSingle E.683 scCLTdegenNuc1008

Average coverage per gene

0.003 0.005 0.007

-5kb

Gene start

Gene end

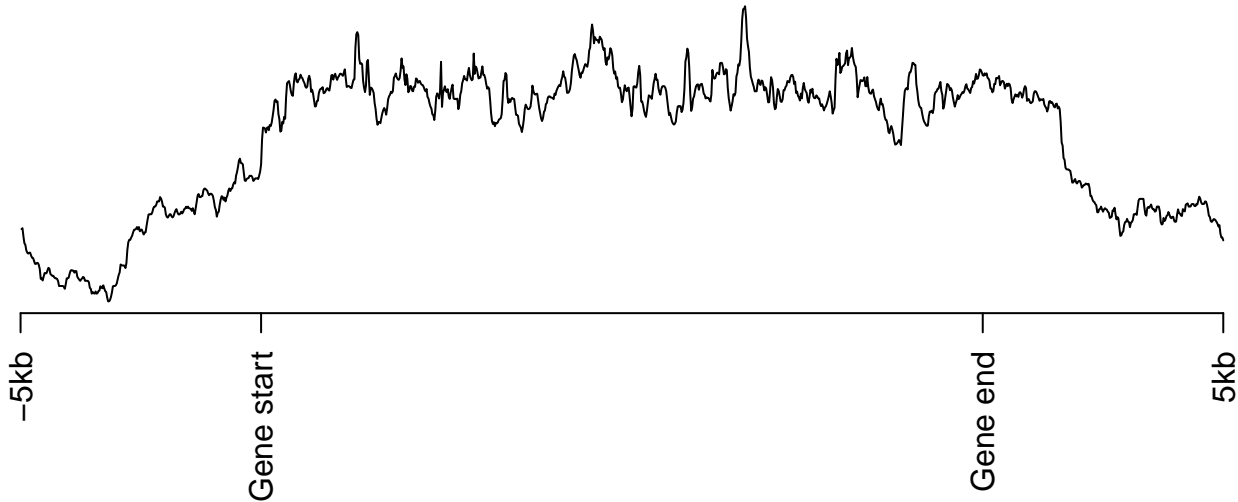
5kb



K562 PositiveSingle E.683 scCLTdegenNuc1009

Average coverage per gene

0.0015 0.0030



K562 PositiveSingle E.683 scCLTdegenNuc1010

Average coverage per gene

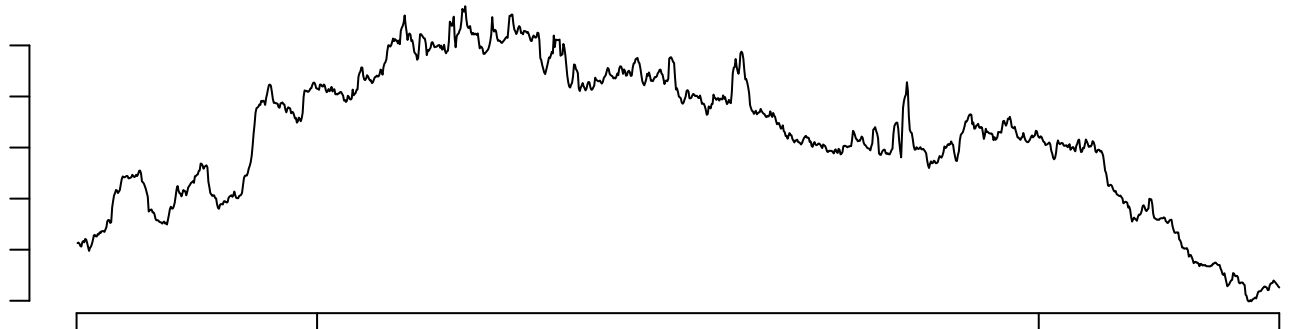
0.002 0.005

-5kb

Gene start

Gene end

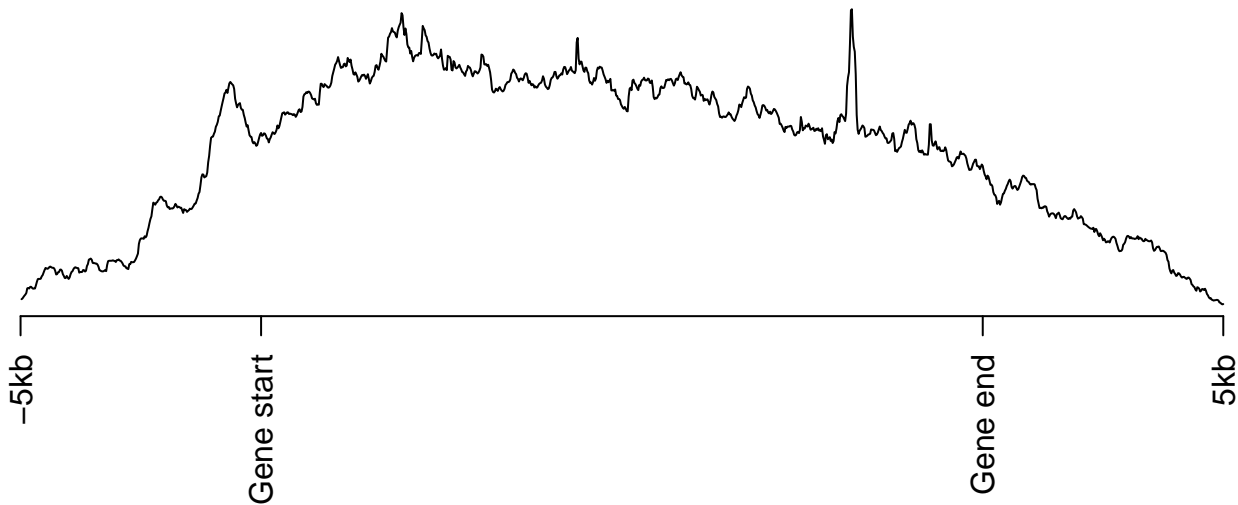
5kb



K562 PositiveSingle E.683 scCLTdegenNuc1012

Average coverage per gene

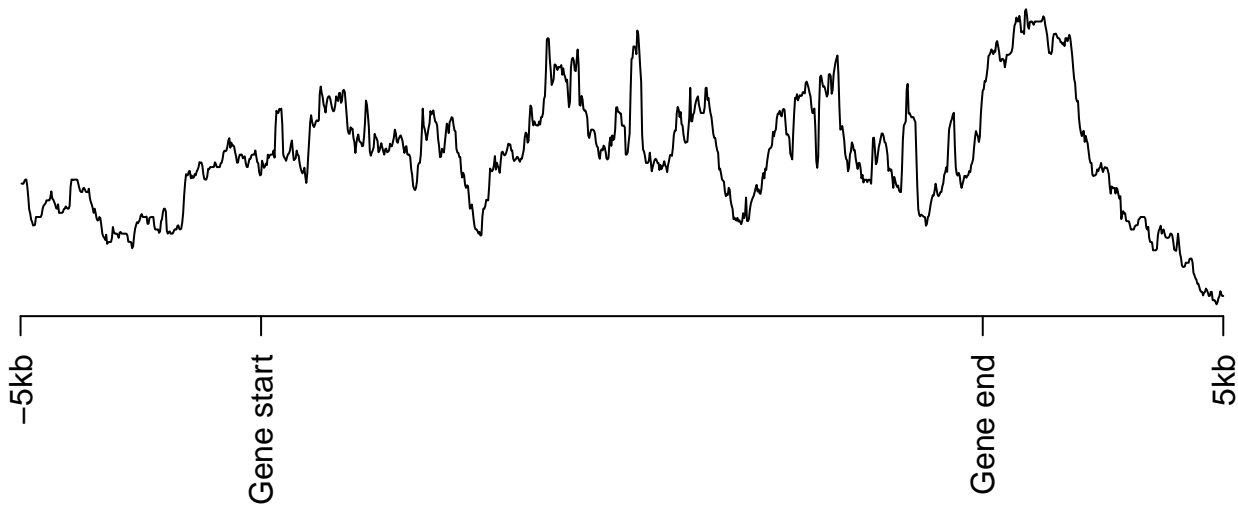
0.002 0.004 0.006



K562 PositiveSingle E.683 scCLTdegenNuc1013

Average coverage per gene

0.0010 0.0020



K562 PositiveSingle E.683 scCLTdegenNuc979

Average coverage per gene

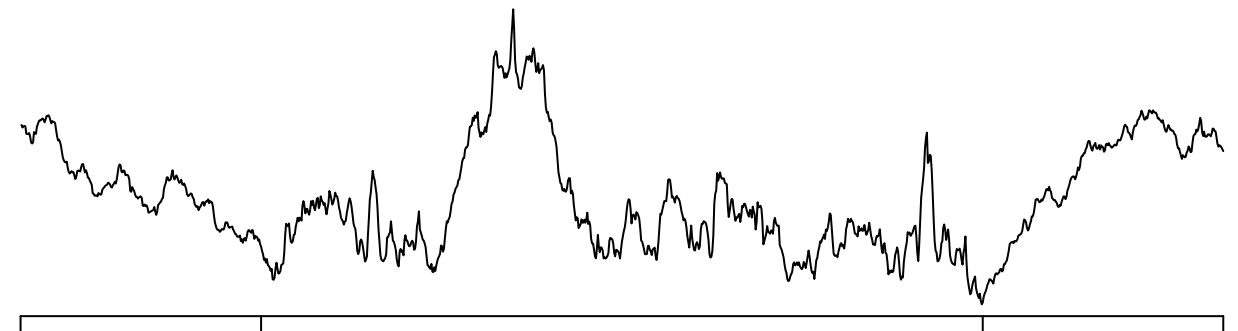
0.004 0.006 0.008

-5kb

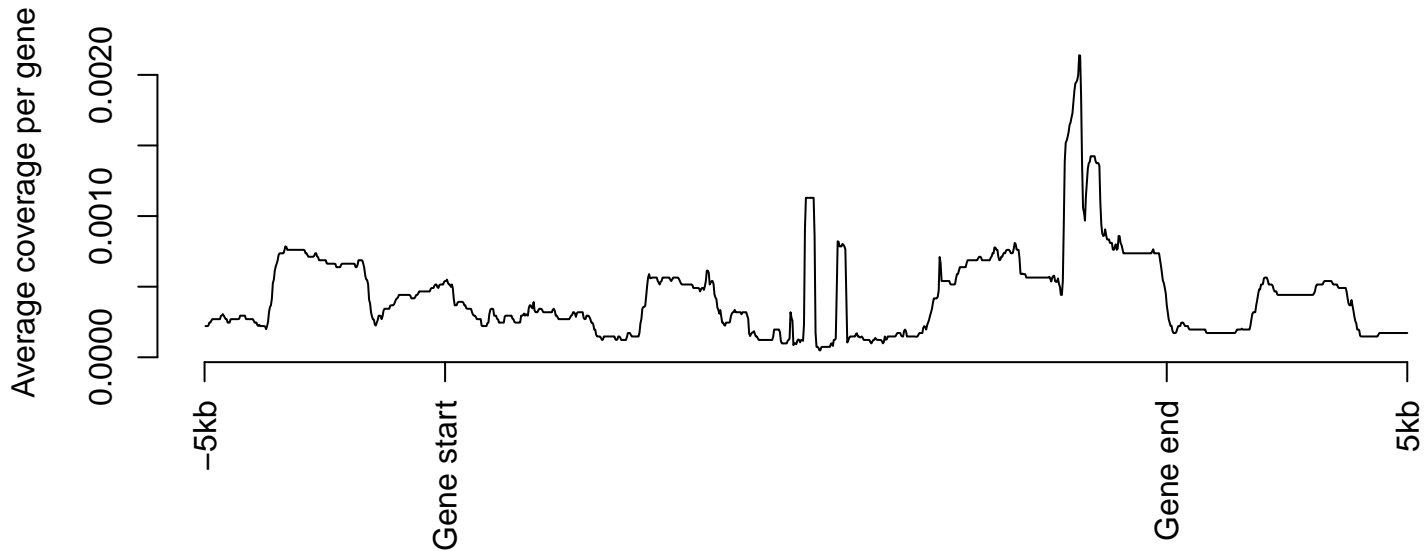
Gene start

Gene end

5kb

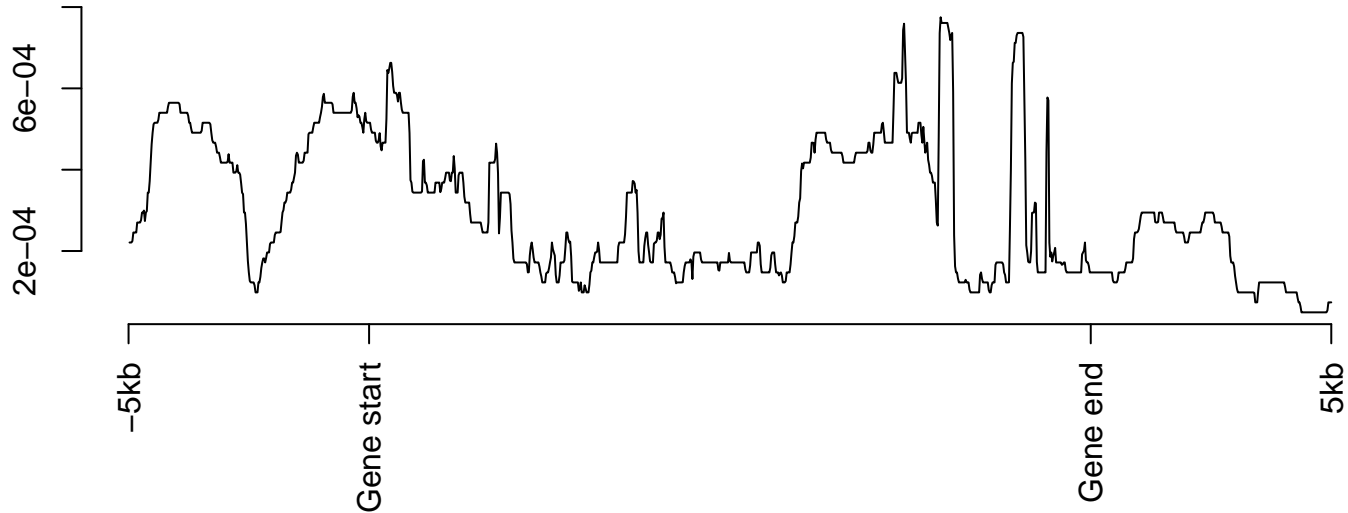


K562 PositiveSingle E.694 scCLTdegenNuc1077



K562 PositiveSingle E.694 scCLTdegenNuc1078

Average coverage per gene



K562 PositiveSingle E.694 scCLTdegenNuc1079

Average coverage per gene

0.0005 0.0015

-5kb

Gene start

Gene end

5kb



K562 PositiveSingle E.694 scCLTdegenNuc1080

Average coverage per gene

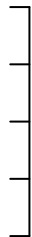
0.0030
0.0015

-5kb

Gene start

Gene end

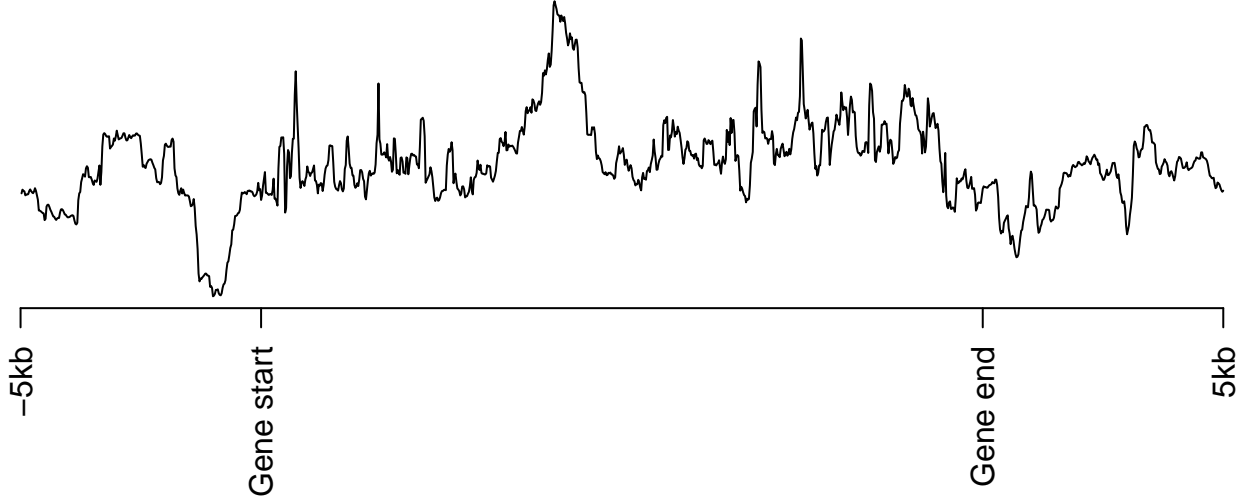
5kb



K562 PositivePooled E.574 scCLTdegenNuc333

Average coverage per gene

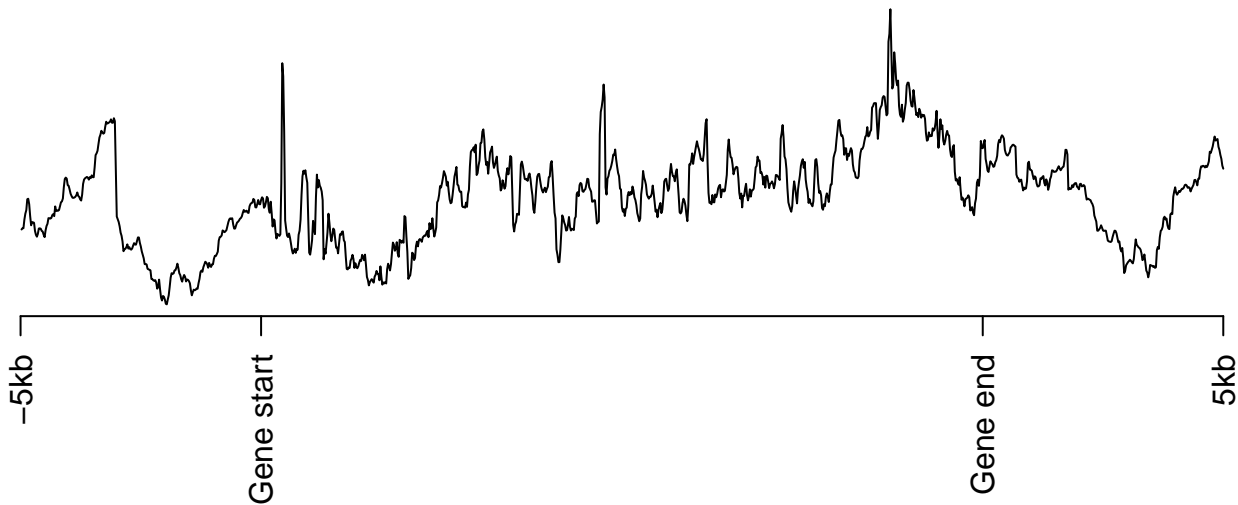
0.012 0.018 0.024



K562 PositivePooled E.588 scCLTdegenNuc451

Average coverage per gene

0.018 0.022 0.026



K562 PositivePooled E.588 scCLTdegenNuc457

Average coverage per gene

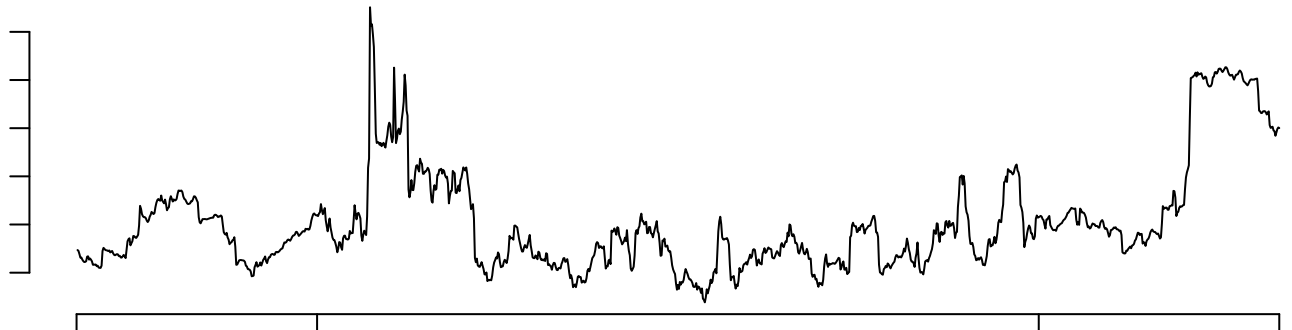
0.014
0.008

-5kb

Gene start

Gene end

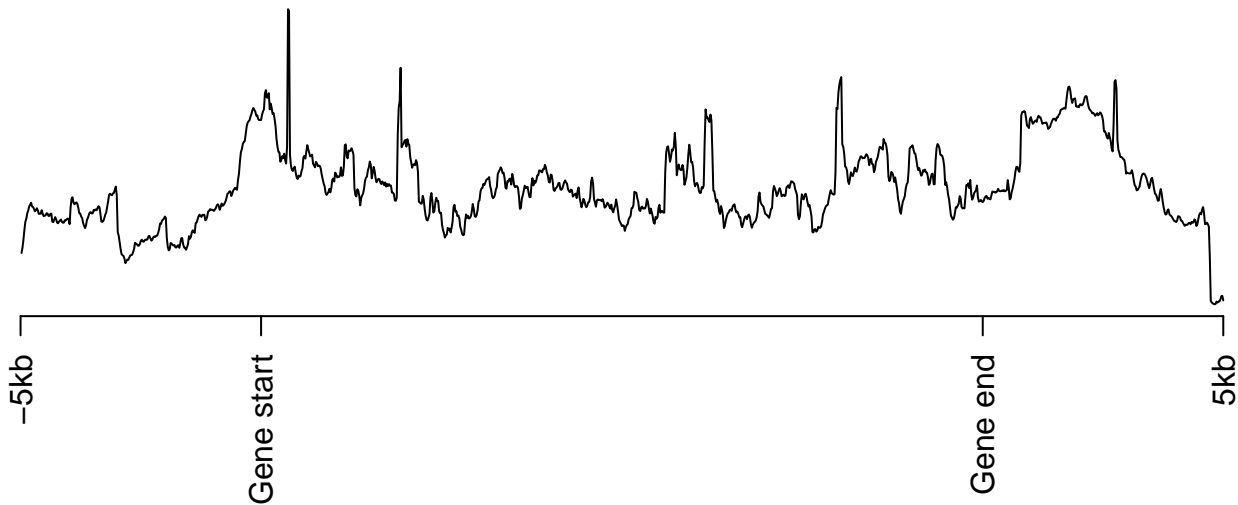
5kb



K562 PositivePooled E.652 scCLTdegenNuc656

Average coverage per gene

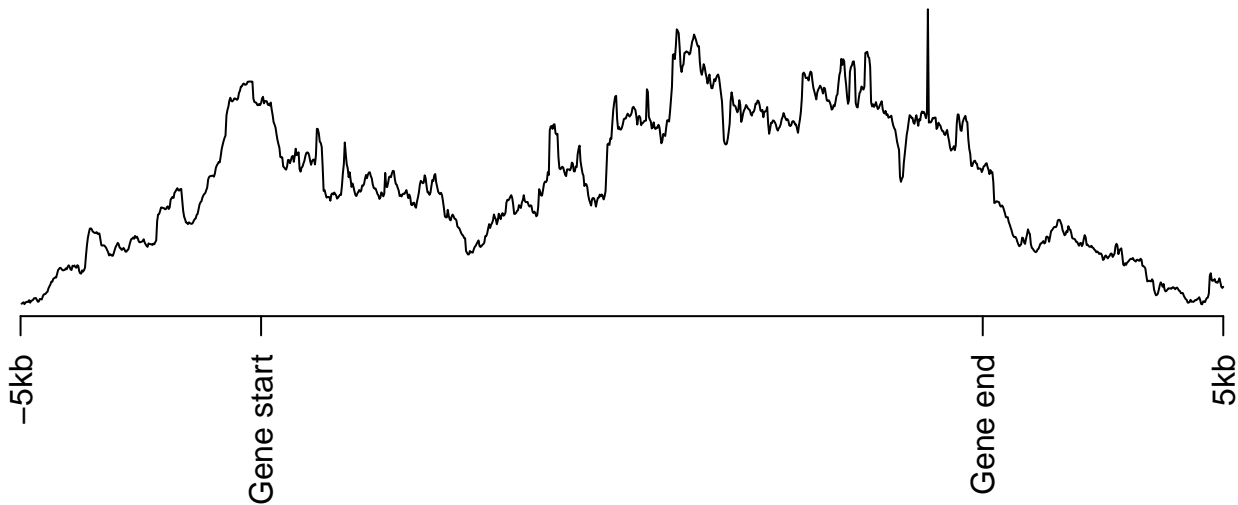
0.030 0.040 0.050



K562 PositivePooled E.652 scCLTdegenNuc657

Average coverage per gene

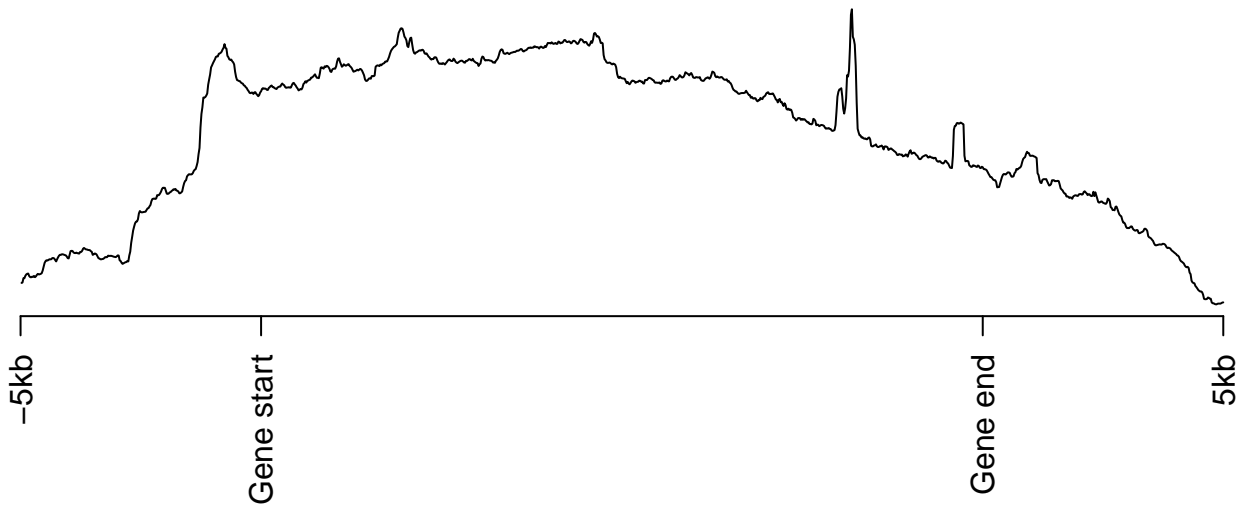
0.07
0.05



K562 PositivePooled E.683 scCLTdegenNuc1011

Average coverage per gene

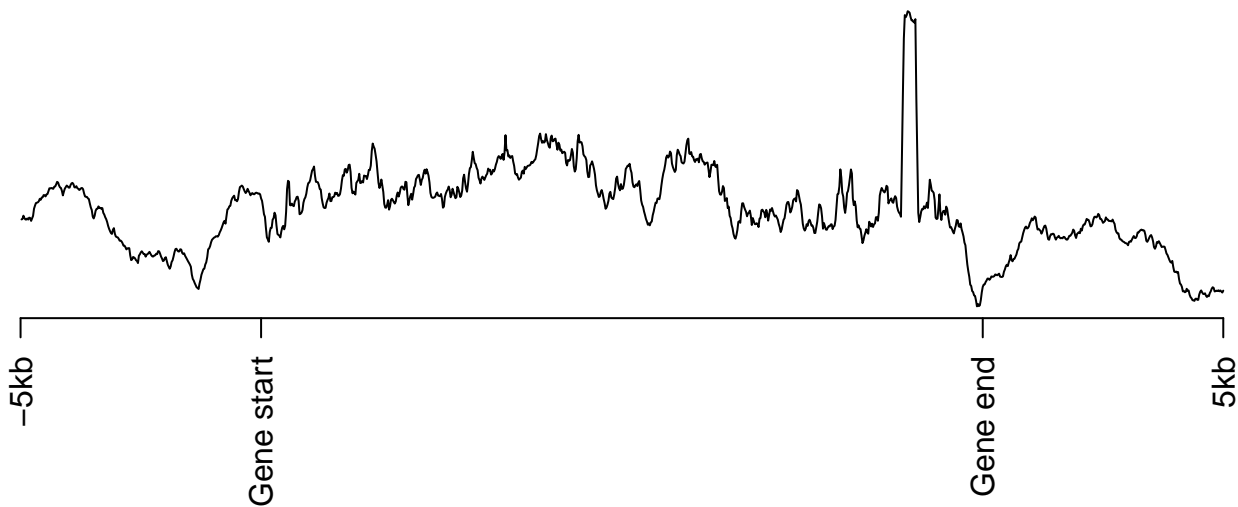
0.04 0.08 0.12



K562 PositiveMerged E.- K562PositiveMerged

Average coverage per gene

0.14 0.16 0.18



K562 PositiveAll E.- K562PositiveAll

Average coverage per gene

0.30
0.40

-5kb

Gene start

Gene end

5kb

