

# K562TPAnone PositiveSingle E.644 scCLTdegenNuc628

Average coverage per intergenic region

0.0005 0.0015

-5kb

intergenic start

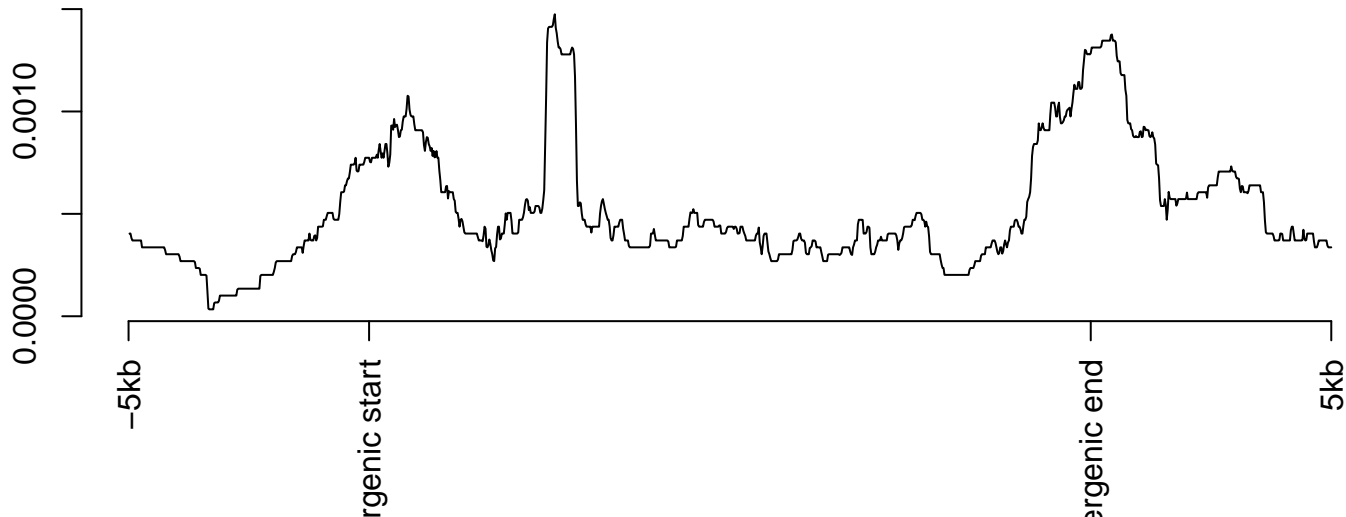
intergenic end

5kb



Average coverage per intergenic region

K562TPAnone PositiveSingle E.644 scCLTdegenNuc629



Average coverage per intergenic region

K562TPAnone PositiveSingle E.644 scCLTdegenNuc630

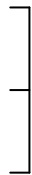
0.0020  
0.0010

-5kb

intergenic start

intergenic end

5kb



Average coverage per intergenic region

K562TPAnone PositiveMerged E.– K562TPAnonePositiveMerged

