

K562TPA24hr PositiveSingle E.644 scCLTdegenNuc634

Average coverage per intergenic region

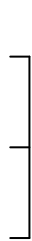
0.0015 0.0025

-5kb

intergenic start

intergenic end

5kb



Average coverage per intergenic region

K562TPA24hr PositiveSingle E.644 scCLTdegenNuc635

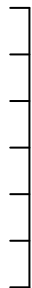
0.0016
0.0010
0.0004

-5kb

intergenic start

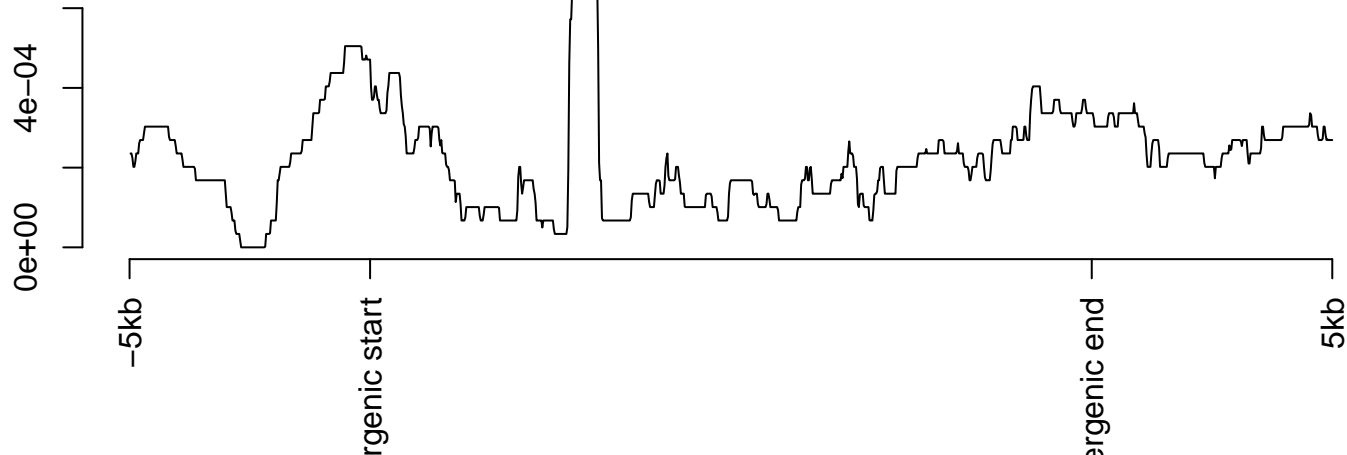
intergenic end

5kb



K562TPA24hr PositiveSingle E.651 scCLTdegenNuc636

Average coverage per intergenic region



K562TPA24hr PositiveSingle E.651 scCLTdegenNuc637

Average coverage per intergenic region

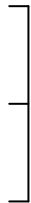
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-5kb

intergenic start

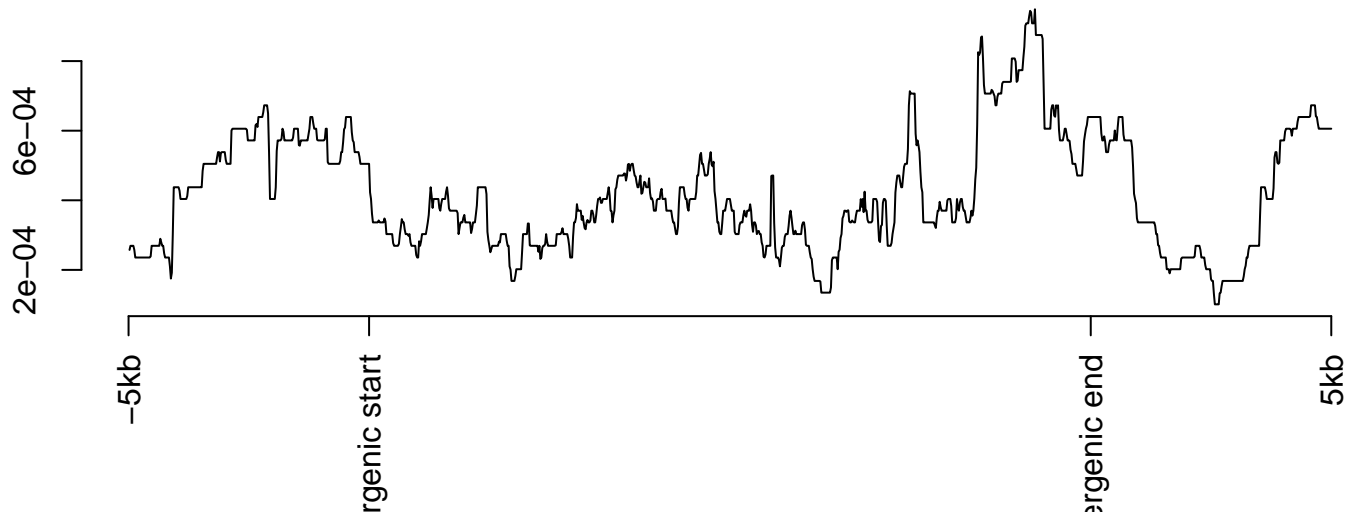
intergenic end

5kb



Average coverage per intergenic region

K562TPA24hr PositiveSingle E.651 scCLTdegenNuc638



Average coverage per intergenic region

K562TPA24hr PositiveMerged E.– K562TPA24hrPositiveMerged

