chr17:31,155,955-31,156,018

p13.2 p13.1 p12 p11.2 p11.1 q11.2 q12 q21.1 q21.31 q21.32 q22 q23.1 q23.3 q24.2 q24.3 q25.1 q25.3

65 bp

31,155,960 bp | 31,155,970 bp | 31,155,980 bp | 31,155,990 bp | 31,156,000 bp | 31,156,010 bp

KU10K-05709...combined.bw

[0 - 98]

KU10K-05709.5mc.hg38.aligned.w
hmod_hg38.bam Coverage

[0 - 28]

2

KU10K-05709.5mc.hg38.aligned.w
hmod_hg38.bam

Sequence

A C G T G t t t t t t t t t t c t t t t t t t t t C A G C T T C C A A T A A A A A C A G G A C A G C A G A A C A C A C A T A C C

L P I K T G Q Q N T H T

Refseq Genes

NF1