

UCSC Genome Browser on Human Feb. 2009 (GRCh37/hg19) Assembly

move <<< << < > >> >>> zoom in 1.5x 3x 10x base zoom out 1.5x 3x 10x

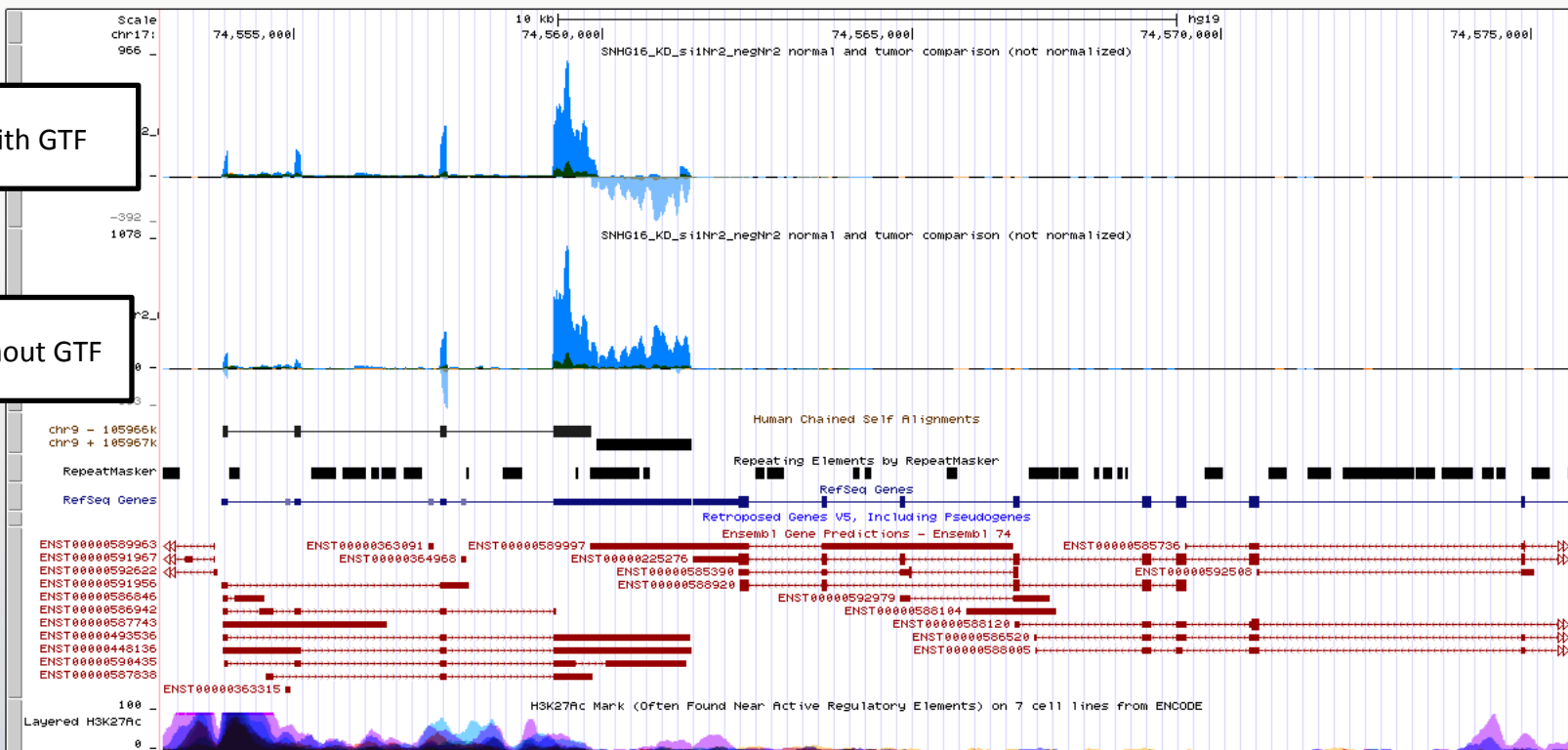
chr17:74,552,891-74,575,645 22,755 bp. enter position, gene symbol or search terms

go

chr17 (q25.1) p13.3 p13.2 17p13.1 17p12 17p11.2 17q11.2 17q12 21.2 q21.31 21.33 17q22 q23.2 q24.2 q24.3 q25.1 17q25.3

With GTF

Without GTF

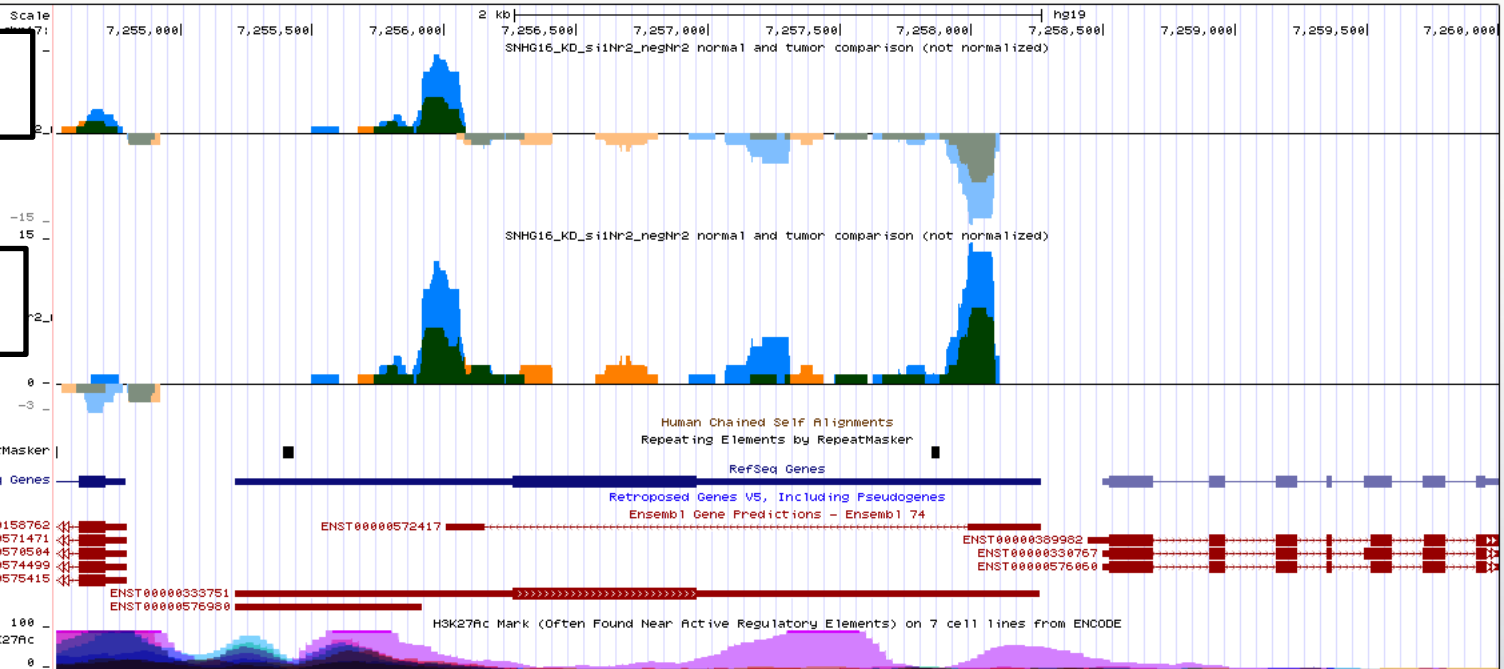


UCSC Genome Browser on Human Feb. 2009 (GRCh37/hg19) Assembly

move <<< << < > >> >>> zoom in 1.5x 3x 10x base zoom out 1.5x 3x 10x

chr17:7,254,529-7,260,001 5,473 bp. | go

chr17 (p13.1) p13.3 p13.2 p13.1 17p12 17p11.2 17q11.2 17q12 21.2 q21.31 21.33 17q22 q23.2 q24.2 q24.3 q25.1 17q25.3



With GTF

Without GTF

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move <<< << < > >> >>> zoom in 1.5x 3x 10x base zoom out 1.5x 3x 10x

chr17:7,207,461-7,216,998 9,538 bp.

go

chr17 (p13.1) p13.3 p13.2 p13.1 17p12 17p11.2 17q11.2 17q12 21.2 q21.31 21.33 17q22 q23.2 q24.2 q24.3 q25.1 17q25.3

Scale chr17: 7,208,000| 7,209,000| 7,210,000| 7,211,000| 7,212,000| 7,213,000| 7,214,000| 7,215,000| 7,216,000| hg19
1324
2 kb
SNHG16_KD_si1Nr2_negNr2 normal and tumor comparison (not normalized)

With GTF

SNHG16_KD_si1Nr2_negNr2 normal and tumor comparison (not normalized)

Without GTF

chr10 - 82006k
chr10 + 81272k
chr17 + 76153k
chr19 + 45631k
chr9 + 2875k

Human Chained Self Alignments

Repeating Elements by RepeatMasker

RefSeq Genes

Retroposed Genes V5, Including Pseudogenes

Ensembl Gene Predictions - Ensembl 74

ENST00000336452
ENST00000355058
ENST00000336458
ENST00000576930
ENST00000572815
ENST00000573542
ENST00000575001
ENST00000571955
ENST00000419711
ENST00000573714
ENST00000416015
ENST00000571569
ENST00000571098
ENST00000573053
ENST00000577040
ENST00000572700
ENST00000531950
ENST00000572172
ENST00000570780
ENST00000573607
ENST00000369167
ENST00000573684
ENST00000575474
ENST00000315601
ENST00000574458
ENST00000360728
ENST00000571697

H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE

100
Layered H3K27Ac
0