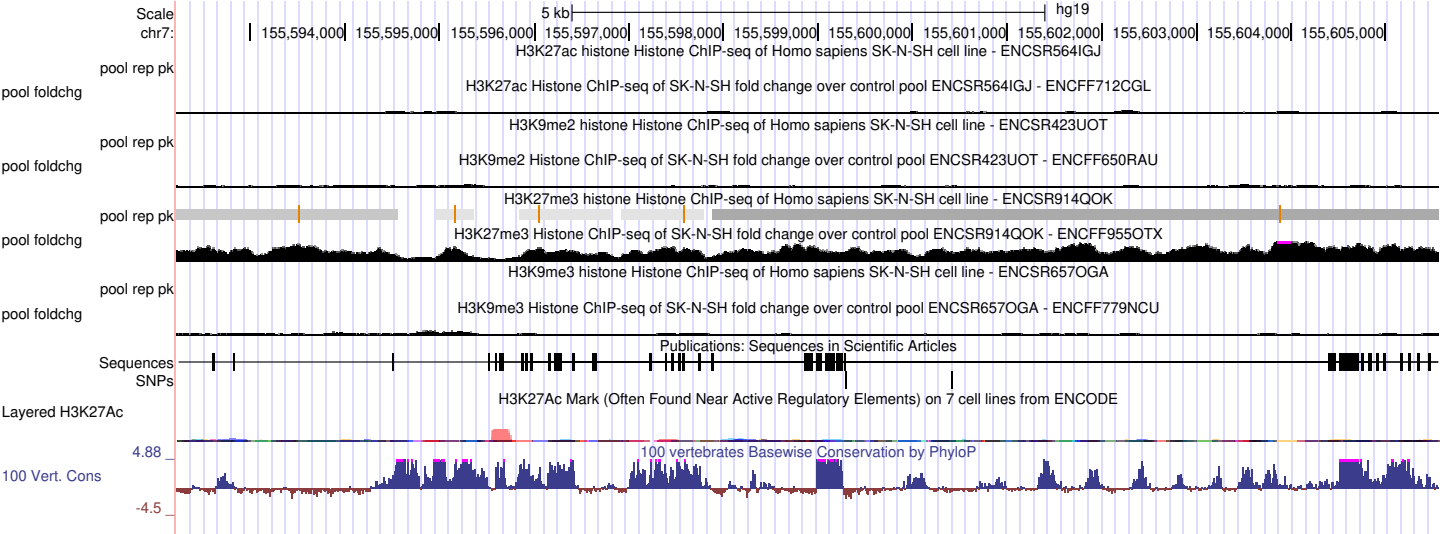
Scale
chr7:
5 kb
hg19
155,594,000
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H3K27ac histone Histone ChIP-seq of Homo sapiens SK-N-SH cell line - ENCSR564IGJ
pool rep pk
pool foldchg
H3K27ac Histone ChIP-seq of SK-N-SH fold change over control pool ENCSR564IGJ - ENCFF712CGL
H3K9me2 histone Histone ChIP-seq of Homo sapiens SK-N-SH cell line - ENCSR423UOT
pool rep pk
pool foldchg
H3K9me2 Histone ChIP-seq of SK-N-SH fold change over control pool ENCSR423UOT - ENCFF650RAU
H3K27me3 histone Histone ChIP-seq of Homo sapiens SK-N-SH cell line - ENCSR914QOK
pool rep pk
pool foldchg
H3K27me3 Histone ChIP-seq of SK-N-SH fold change over control pool ENCSR914QOK - ENCFF955OTX
H3K9me3 histone Histone ChIP-seq of Homo sapiens SK-N-SH cell line - ENCSR657OGA
pool rep pk
pool foldchg
H3K9me3 Histone ChIP-seq of SK-N-SH fold change over control pool ENCSR657OGA - ENCFF779NCU
Publications: Sequences in Scientific Articles
Sequences
SNPs
H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE
Layered H3K27Ac
4.88
100 vertebrates Basewise Conservation by PhyloP
100 Vert. Cons
-4.5