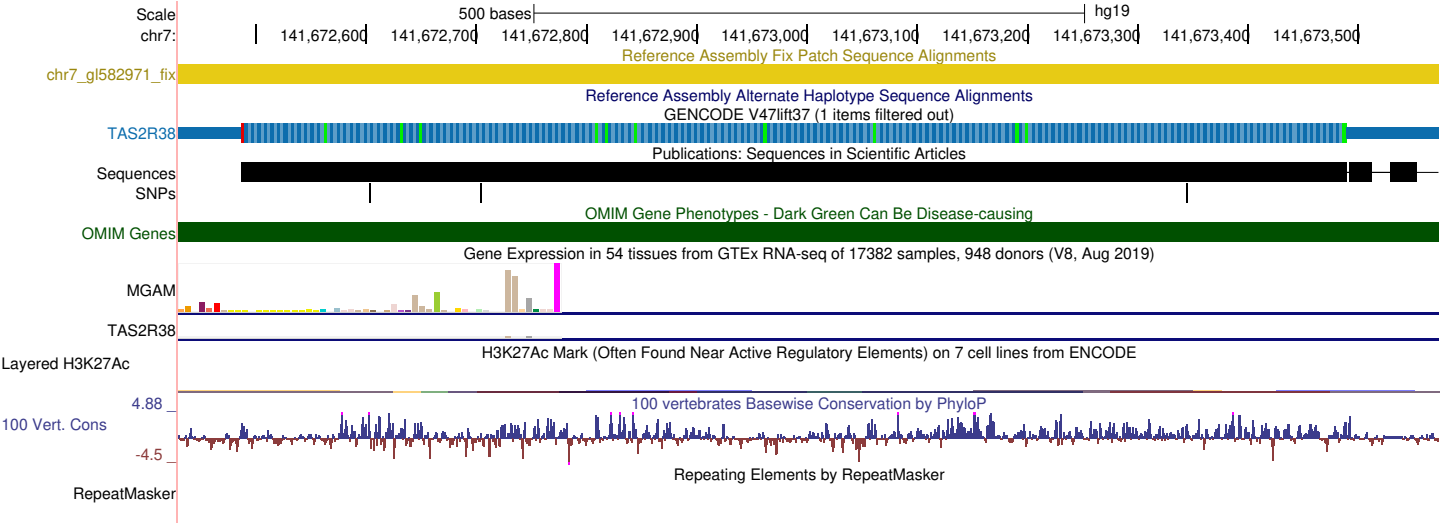
Scale
chr7:
500 bases
hg19
141,672,600
141,672,700
141,672,800
141,672,900
141,673,000
141,673,100
141,673,200
141,673,300
141,673,400
141,673,500
Reference Assembly Fix Patch Sequence Alignments
chr7_gl582971_fix
Reference Assembly Alternate Haplotype Sequence Alignments
GENCODE V47lift37 (1 items filtered out)
TAS2R38
Publications: Sequences in Scientific Articles
Sequences
SNPs
OMIM Gene Phenotypes - Dark Green Can Be Disease-causing
OMIM Genes
Gene Expression in 54 tissues from GTEx RNA-seq of 17382 samples, 948 donors (V8, Aug 2019)
MGAM
TAS2R38
Layered H3K27Ac
H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE
4.88
100 Vert. Cons
-4.5
100 vertebrates Basewise Conservation by PhyloP
Repeating Elements by RepeatMasker
RepeatMasker