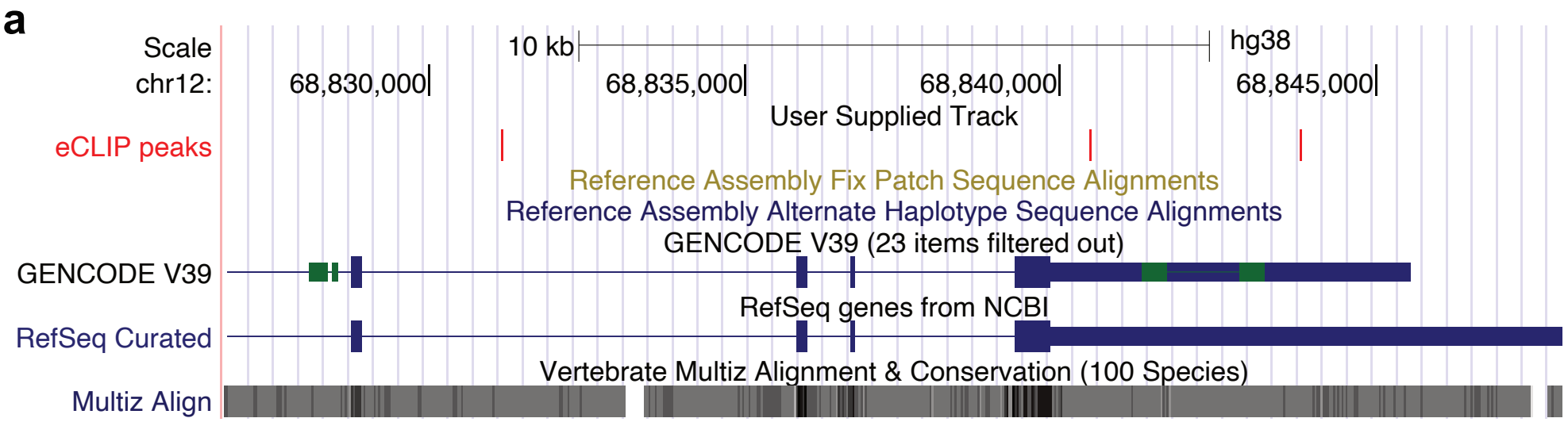
a
Scale
chr12:
10 kb
hg38
68,830,000
68,835,000
68,840,000
68,845,000
User Supplied Track
eCLIP peaks
Reference Assembly Fix Patch Sequence Alignments
Reference Assembly Alternate Haplotype Sequence Alignments
GENCODE V39 (23 items filtered out)
GENCODE V39
RefSeq genes from NCBI
RefSeq Curated
Vertebrate Multiz Alignment & Conservation (100 Species)
Multiz Align