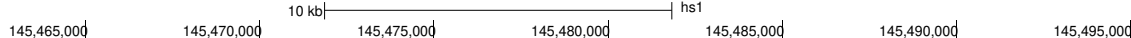


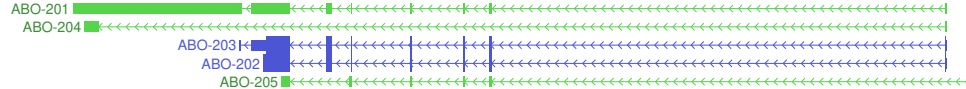
Scale
chr9:



Centromeric Satellite Annotation
CHM13 unique in comparison to GRCh38/hg38

Consensus rDNA models
Perfect Matches to Short Sequence (CAGCAGCAGCAGCAGCAG)

CAT + Liftoff Gene Annotations



NCBI RefSeq genes, predicted subset (XM_* or XR_*)

NCBI RefSeq other annotations (not NM_*, NR_*, XM_*, XR_*, NP_* or YP_*)