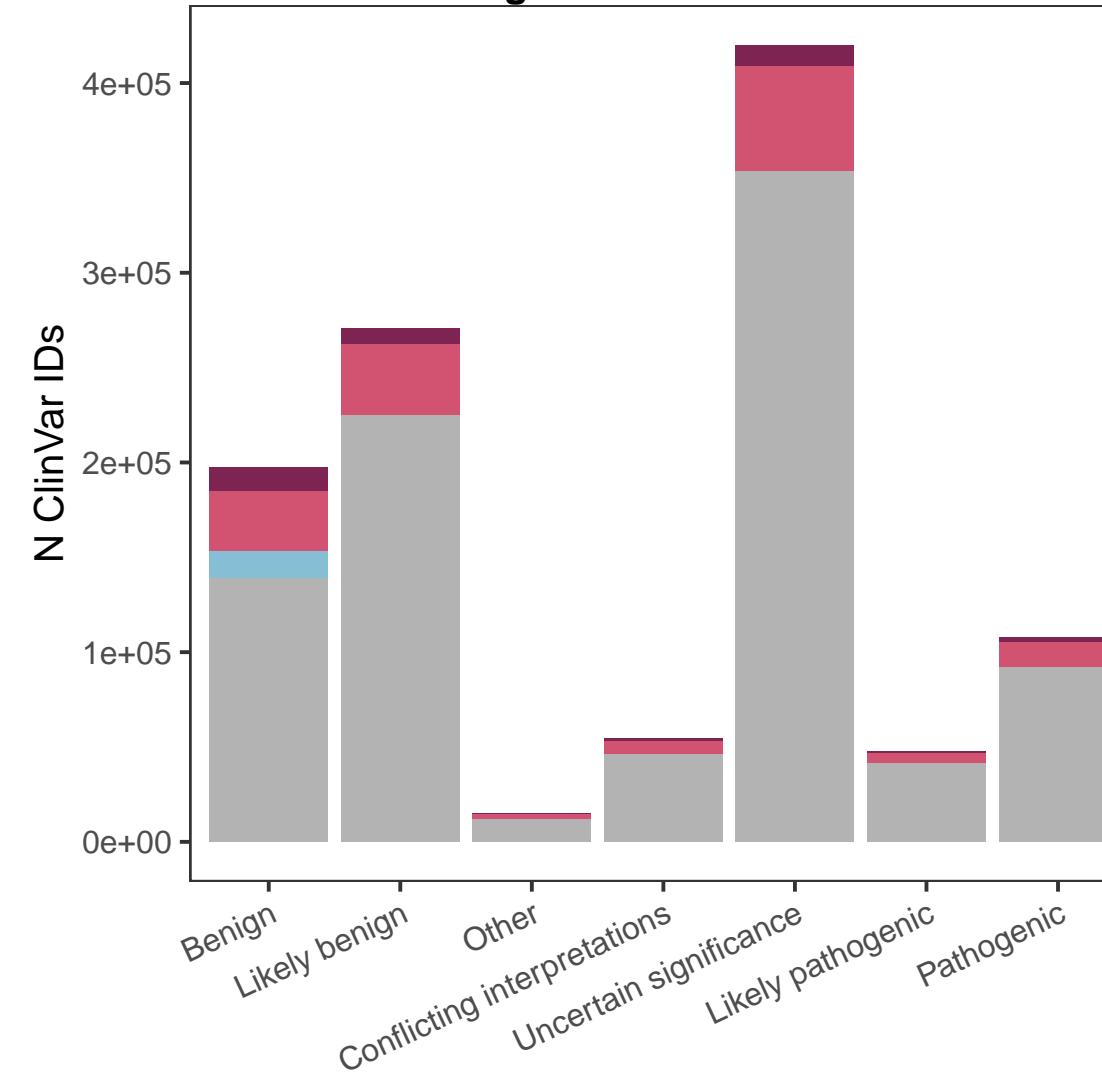
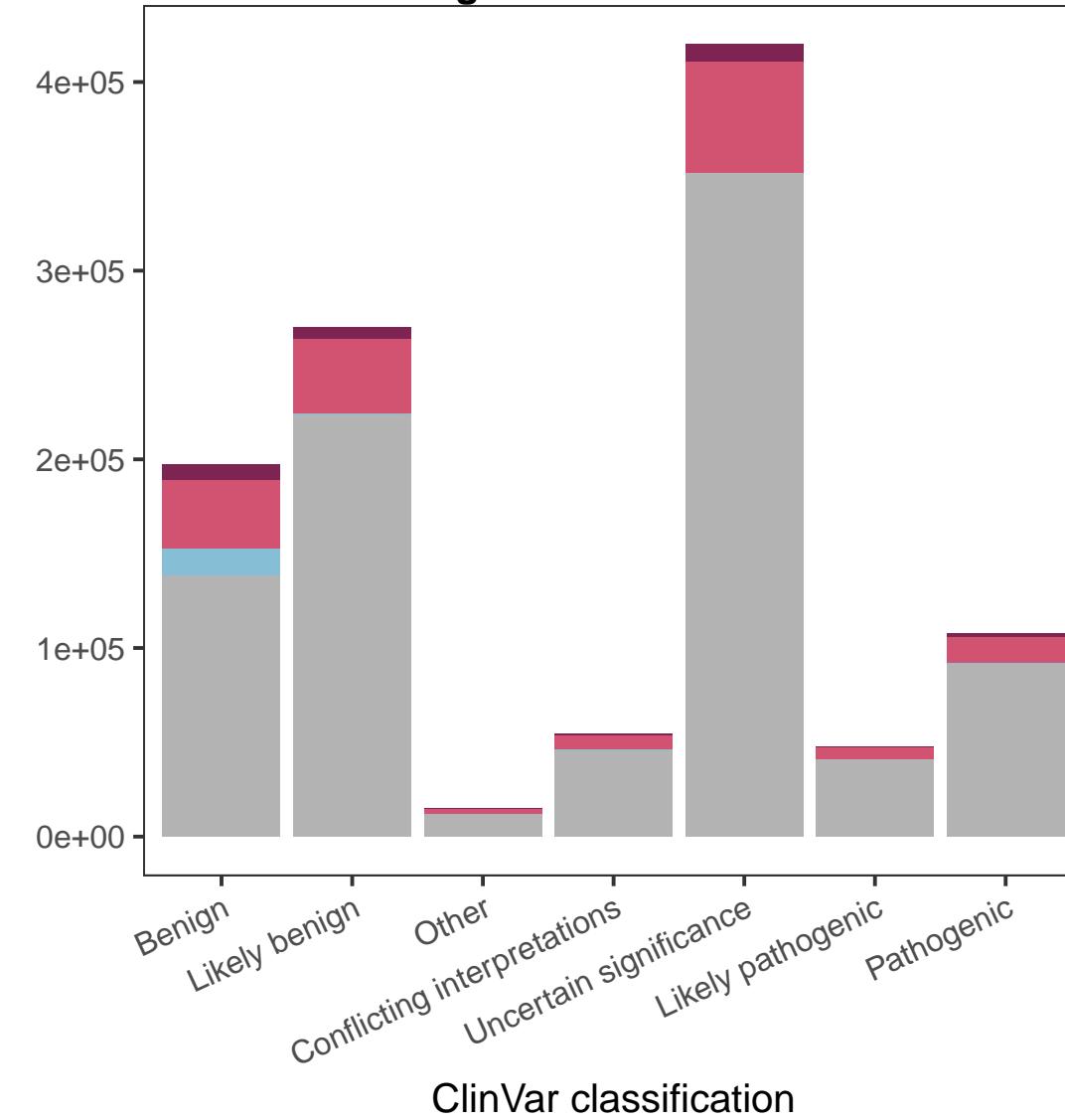


**hg38vsT2T-CHM13****hg19vsT2T-CHM13**

ClinVar classification

Sequence alignment

&gt; 5 mismatches | Indels

[1, 5] mismatches

Allele swapping

Equal

**hg19vshg38**