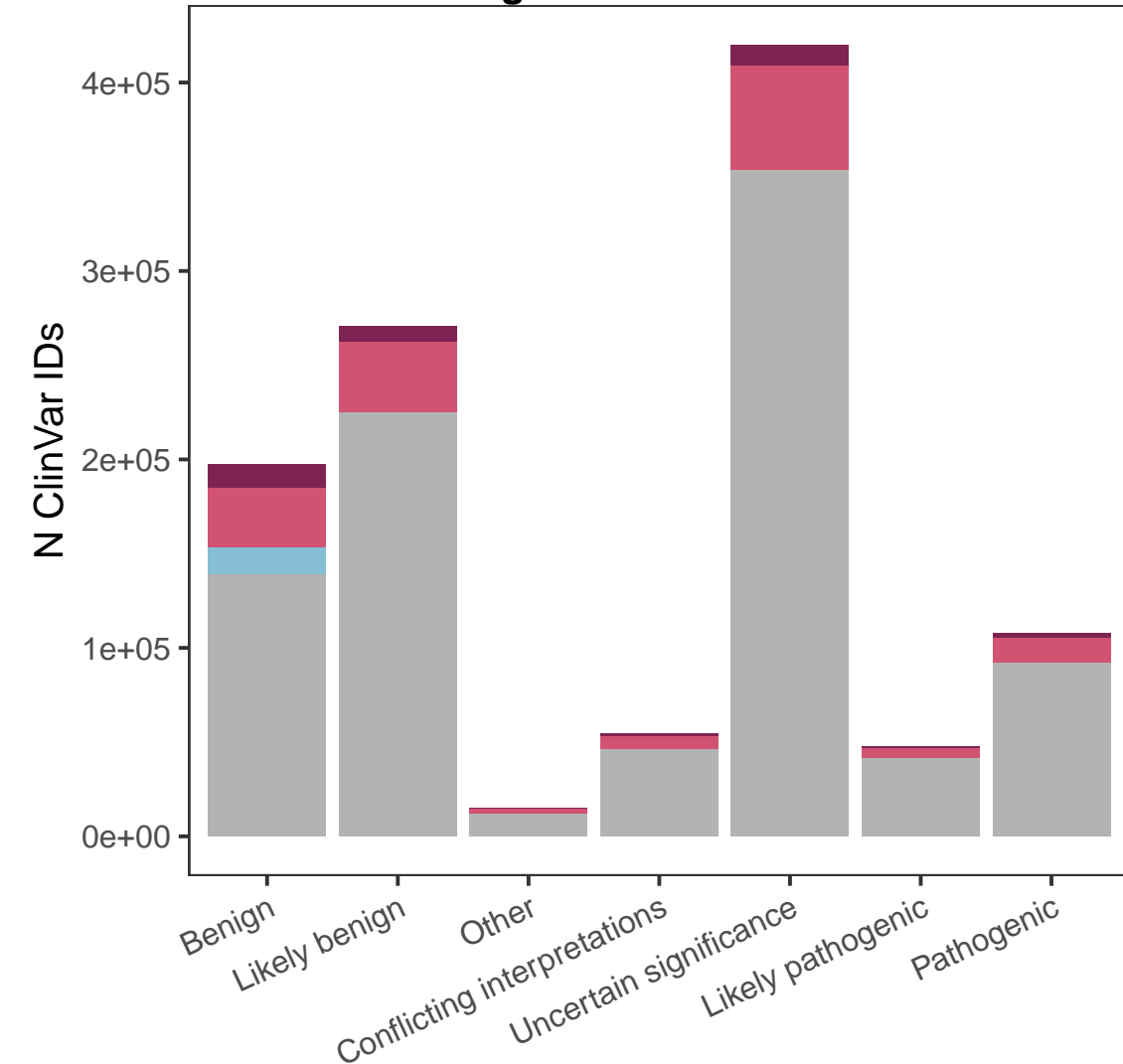
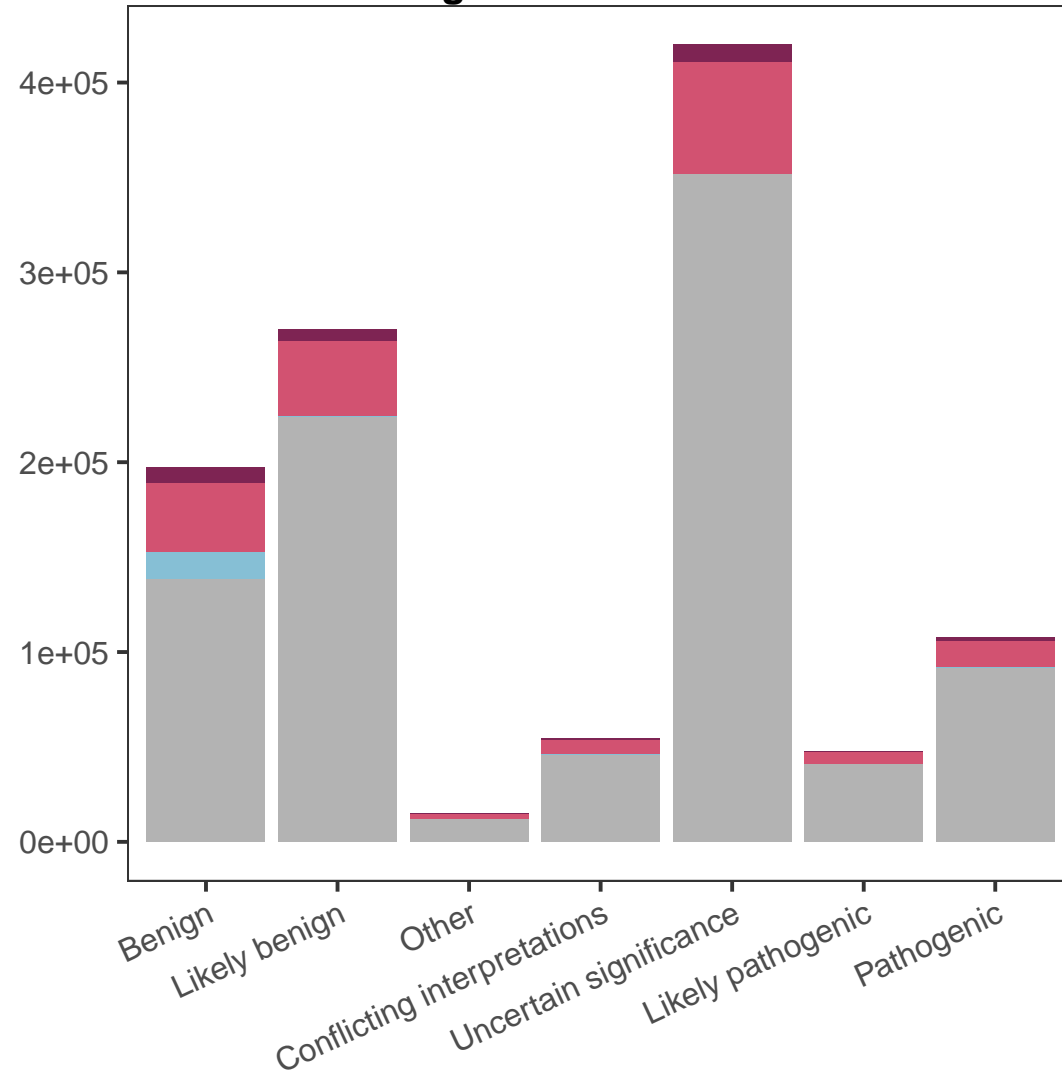


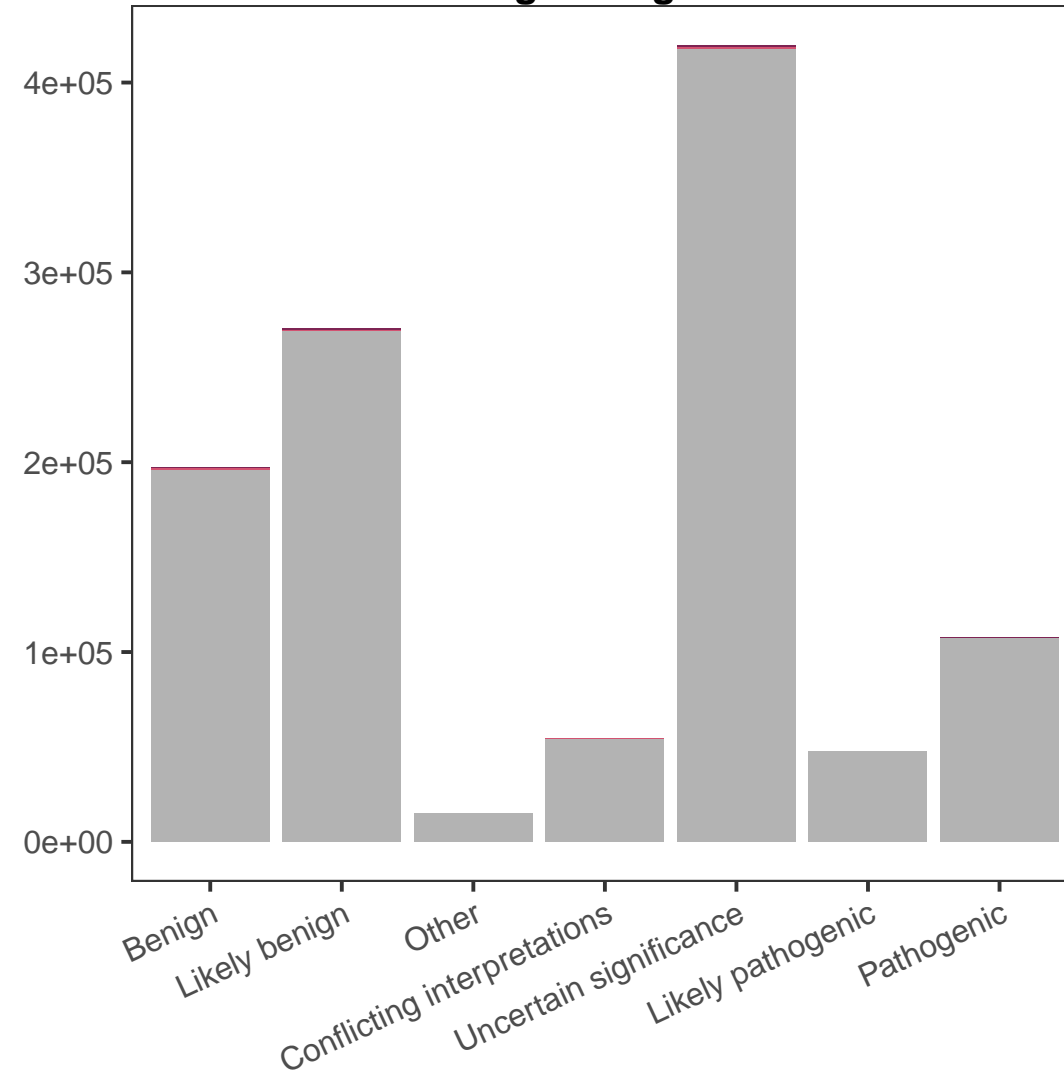
hg38vsT2T-CHM13



hg19vsT2T-CHM13



hg19vshg38



Sequence alignment

- > 5 mismatches | Indels
- [1, 5] mismatches
- Allele swapping
- Equal