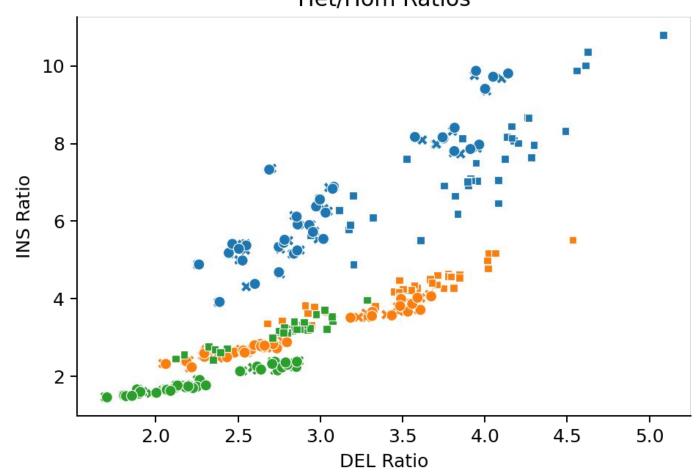


b)

Missing Variants Per-Locus

Het/Hom Ratios



merge

- exact
- strict
- loose reference
- hg19
- **s** grch38
- chm13

