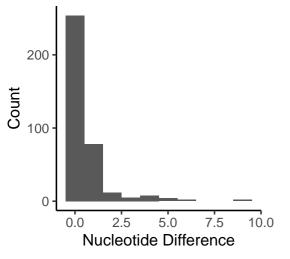


IGKV1E-39*01

365 sequences assigned 254 (69.6%) exact matches, in which: 216 unique CDR3 5 unique J



IGKV2E-40*01

5 sequences assigned 3 (60%) exact matches, in which: 3 unique CDR3 3 unique J

