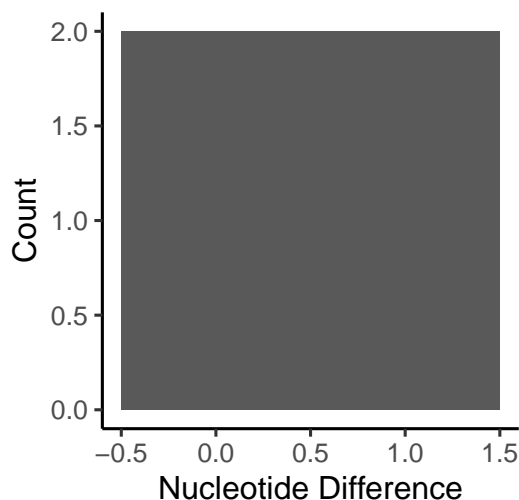


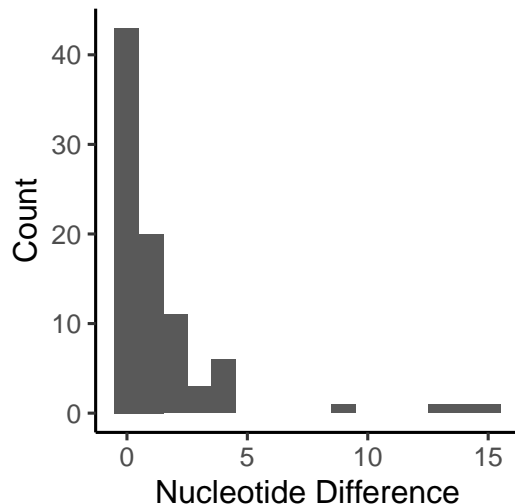
IGLV10-54*01

4 sequences assigned
2 (50%) exact matches, in which:
2 unique CDR3
1 unique J



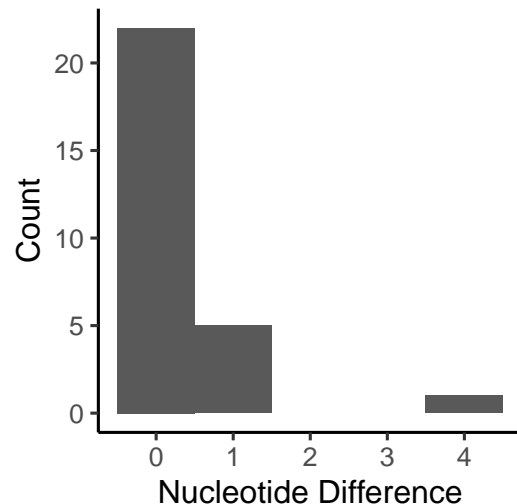
IGLV1-44*01

87 sequences assigned
43 (49.4%) exact matches, in which:
39 unique CDR3
3 unique J



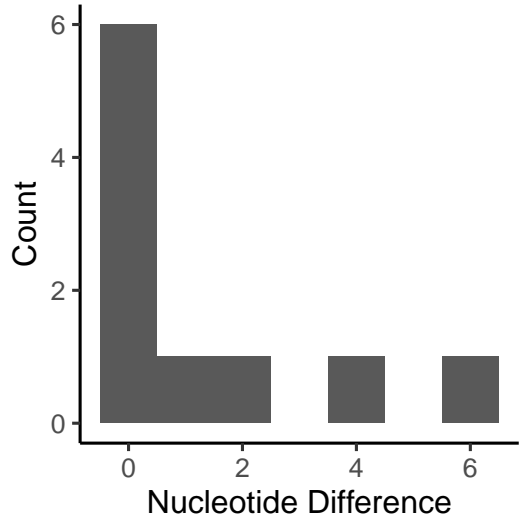
IGLV1-51*02

28 sequences assigned
22 (78.6%) exact matches, in which:
20 unique CDR3
2 unique J



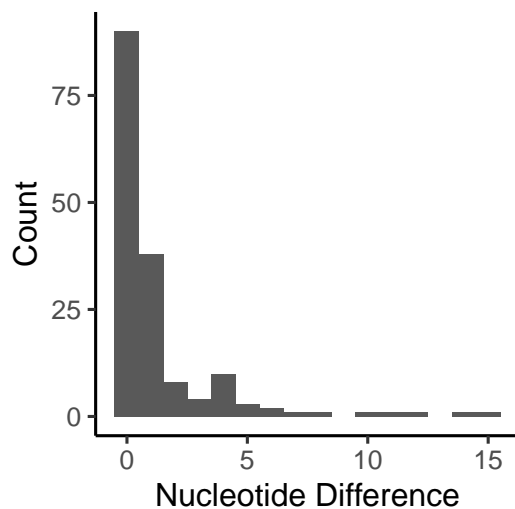
IGLV1-36*01

10 sequences assigned
6 (60%) exact matches, in which:
6 unique CDR3
2 unique J



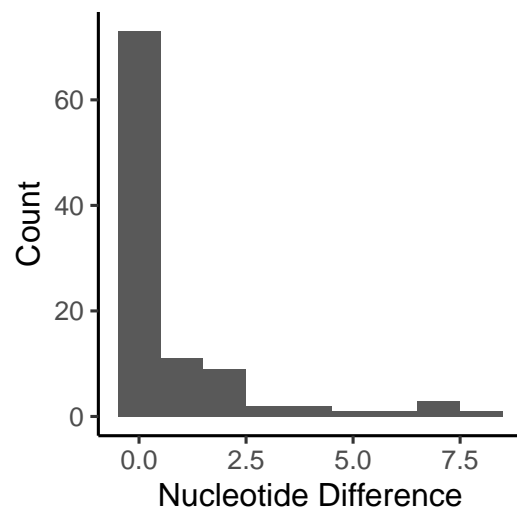
IGLV1-47*01

162 sequences assigned
90 (55.6%) exact matches, in which:
83 unique CDR3
3 unique J



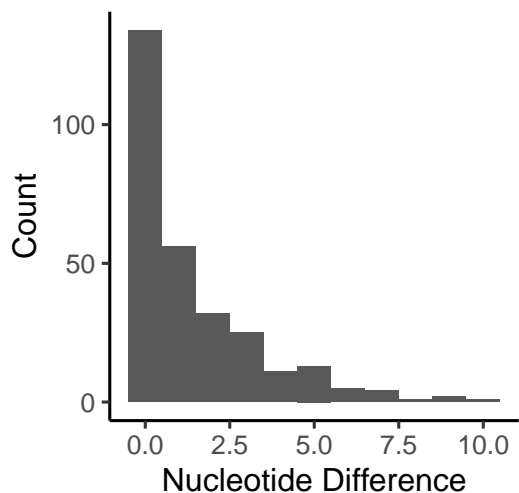
IGLV2-8*01

104 sequences assigned
73 (70.2%) exact matches, in which:
71 unique CDR3
3 unique J



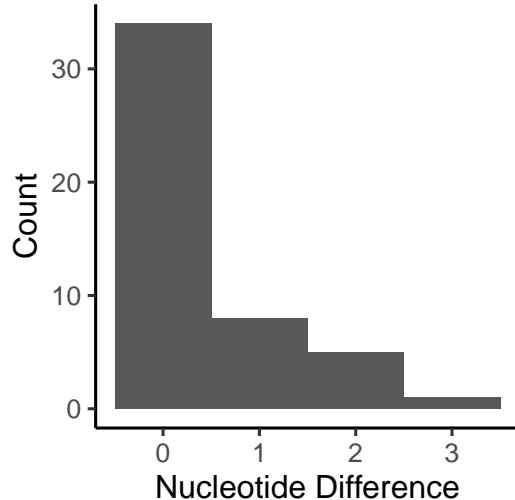
IGLV1-40*01

284 sequences assigned
134 (47.2%) exact matches, in which:
125 unique CDR3
3 unique J



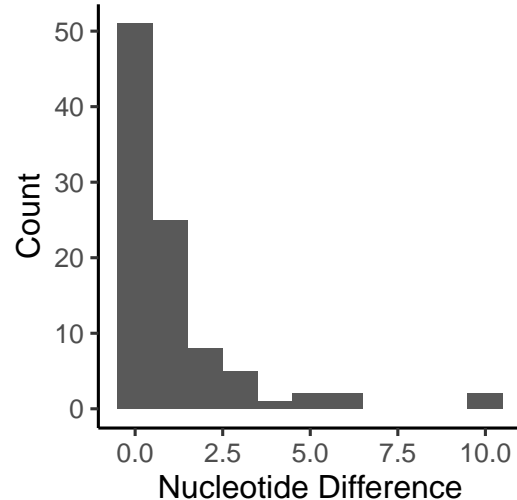
IGLV1-51*01

48 sequences assigned
34 (70.8%) exact matches, in which:
31 unique CDR3
3 unique J



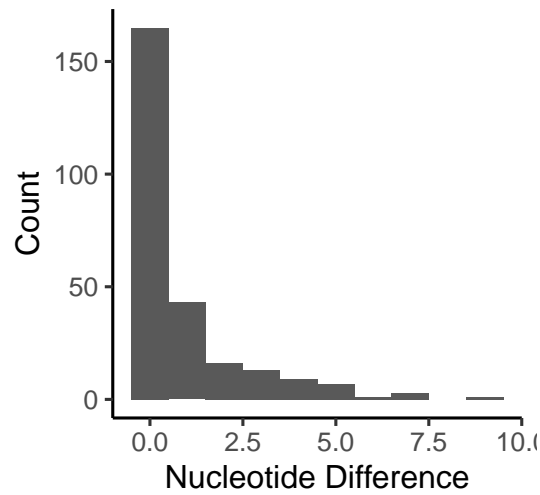
IGLV2-11*01

96 sequences assigned
51 (53.1%) exact matches, in which:
45 unique CDR3
2 unique J



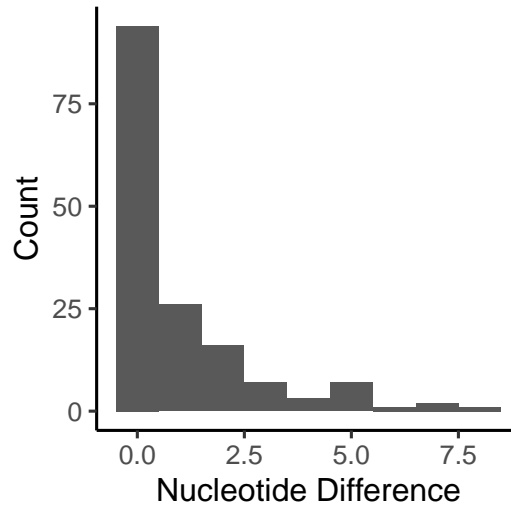
IGLV2-14*01

258 sequences assigned
165 (64%) exact matches, in which:
151 unique CDR3
3 unique J



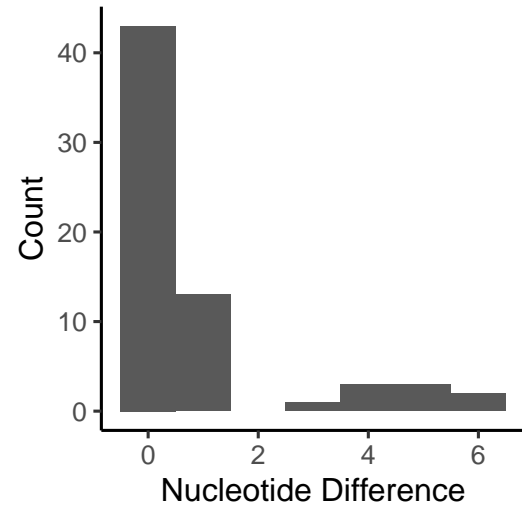
IGLV2-23*03EXT5

157 sequences assigned
94 (59.9%) exact matches, in which:
91 unique CDR3
4 unique J



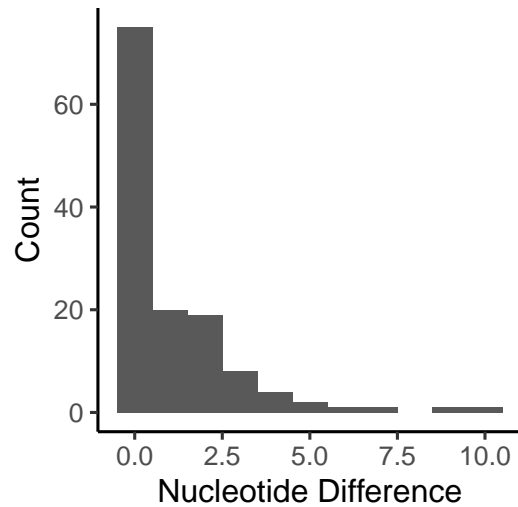
IGLV3-10*01

65 sequences assigned
43 (66.2%) exact matches, in which:
43 unique CDR3
4 unique J



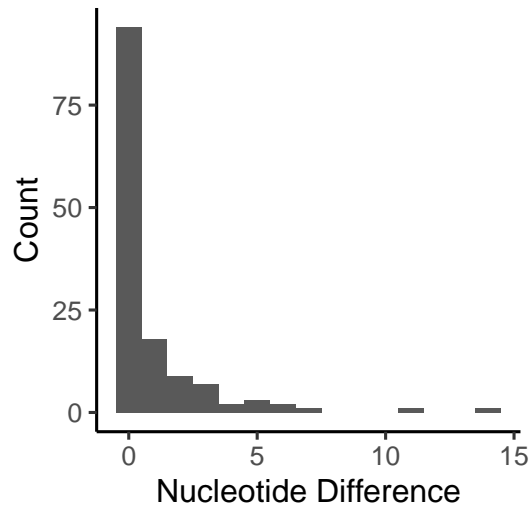
IGLV2-14*04EXT3

132 sequences assigned
75 (56.8%) exact matches, in which:
71 unique CDR3
2 unique J



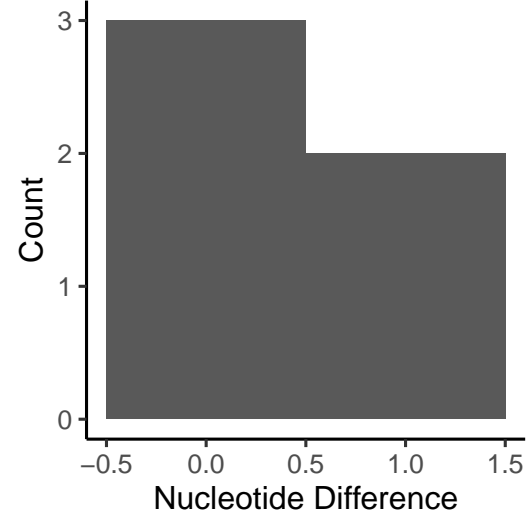
IGLV3-1*01

138 sequences assigned
94 (68.1%) exact matches, in which:
92 unique CDR3
2 unique J



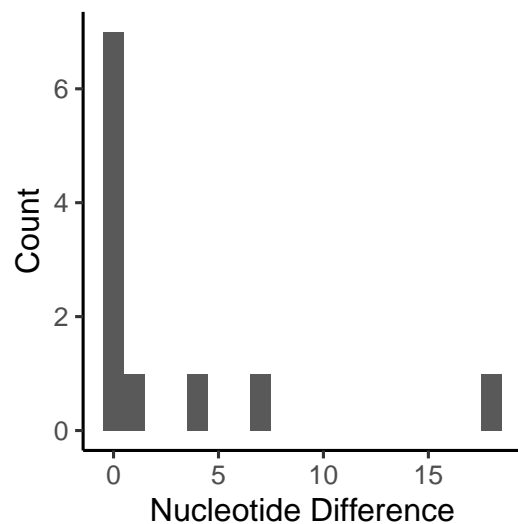
IGLV3-16*01

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



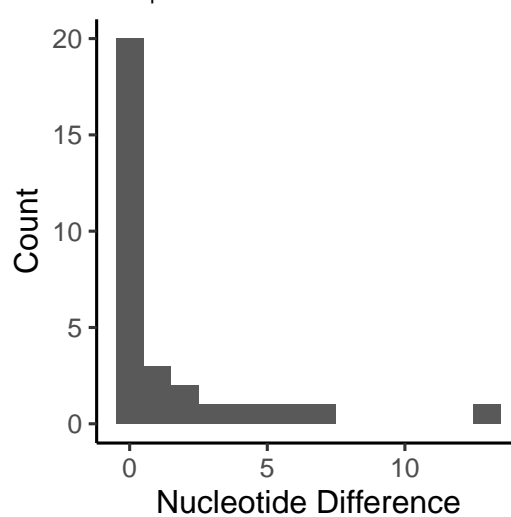
IGLV2-18*02

11 sequences assigned
7 (63.6%) exact matches, in which:
6 unique CDR3
2 unique J



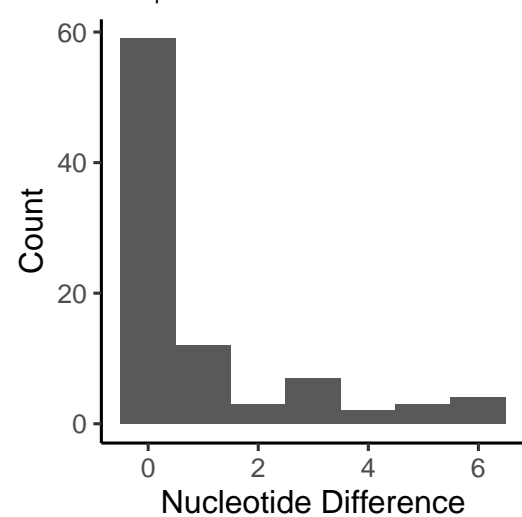
IGLV3-9*01EXT5

31 sequences assigned
20 (64.5%) exact matches, in which:
19 unique CDR3
2 unique J



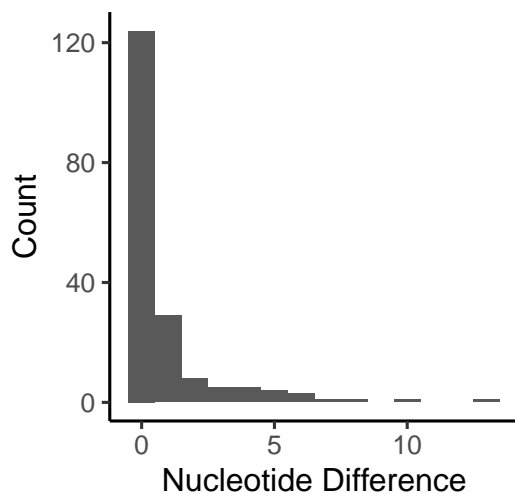
IGLV3-19*01

90 sequences assigned
59 (65.6%) exact matches, in which:
56 unique CDR3
3 unique J



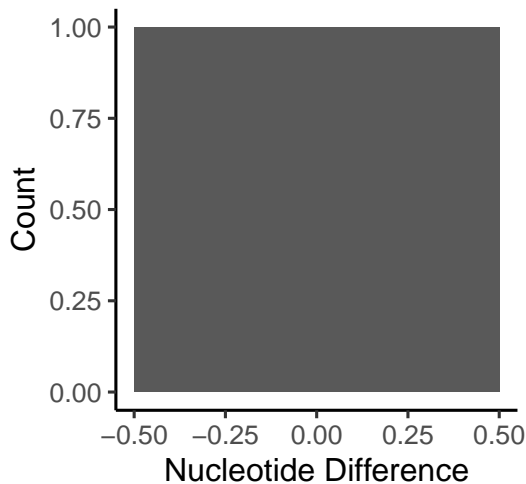
IGLV3-21*02

182 sequences assigned
124 (68.1%) exact matches, in which:
111 unique CDR3
4 unique J



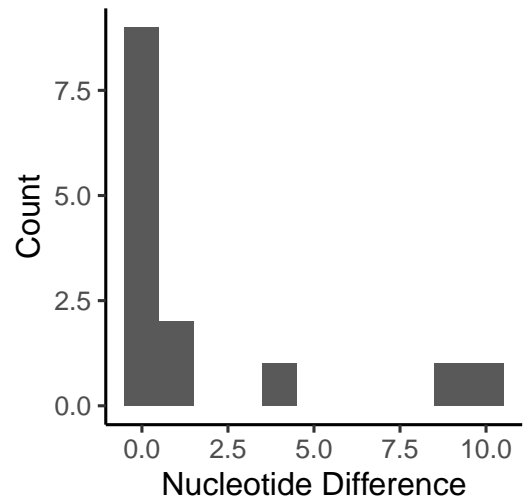
IGLV4-3*01

1 sequences assigned
1 (100%) exact matches, in which:
1 unique CDR3
1 unique J



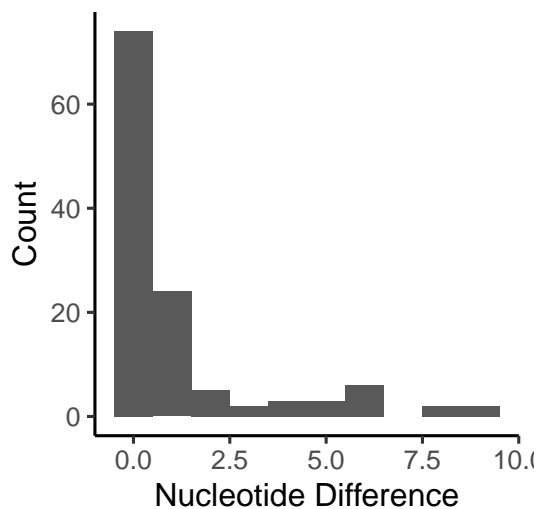
IGLV5-45*01

14 sequences assigned
9 (64.3%) exact matches, in which:
6 unique CDR3
2 unique J



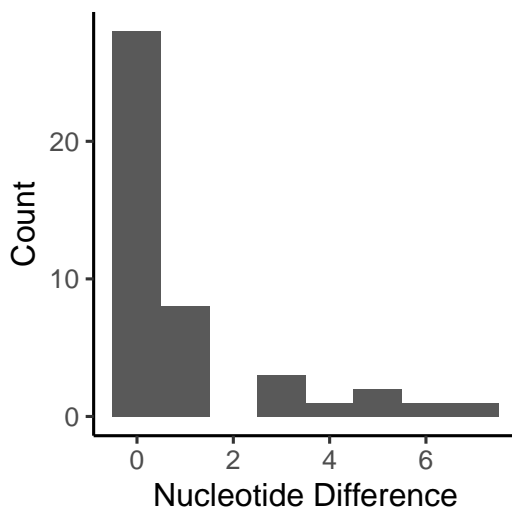
IGLV3-25*03

121 sequences assigned
74 (61.2%) exact matches, in which:
70 unique CDR3
3 unique J



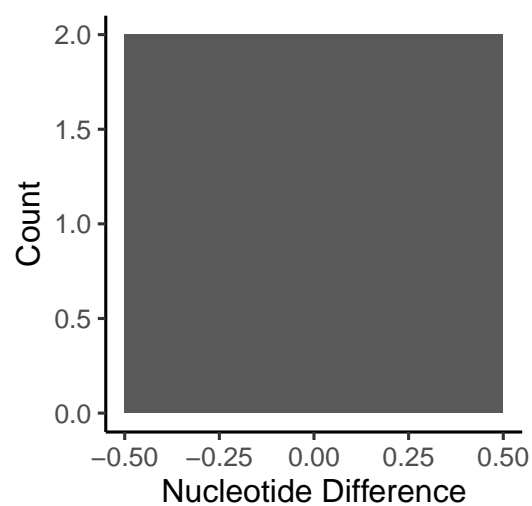
IGLV4-69*01

44 sequences assigned
28 (63.6%) exact matches, in which:
27 unique CDR3
3 unique J



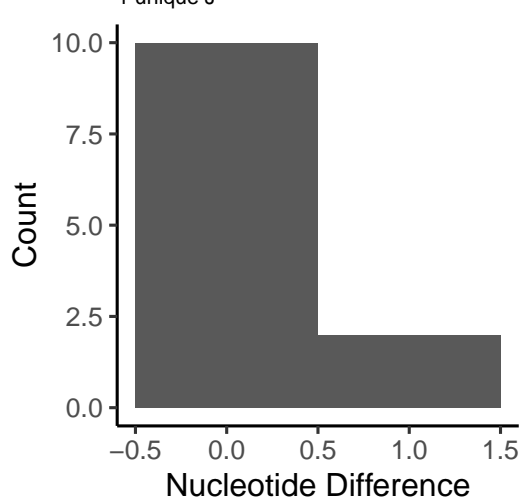
IGLV5-48*01

2 sequences assigned
2 (100%) exact matches, in which:
2 unique CDR3
2 unique J



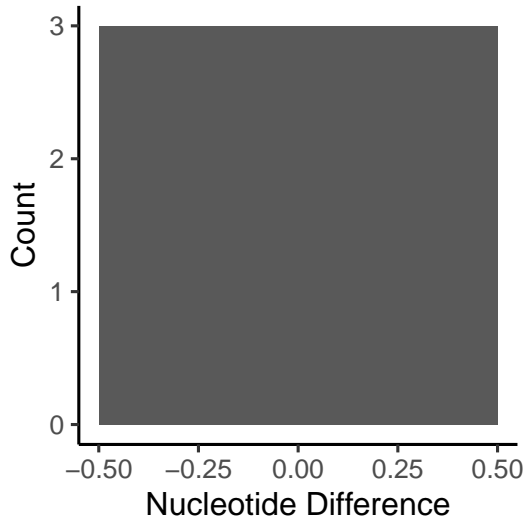
IGLV3-27*01

12 sequences assigned
10 (83.3%) exact matches, in which:
10 unique CDR3
1 unique J



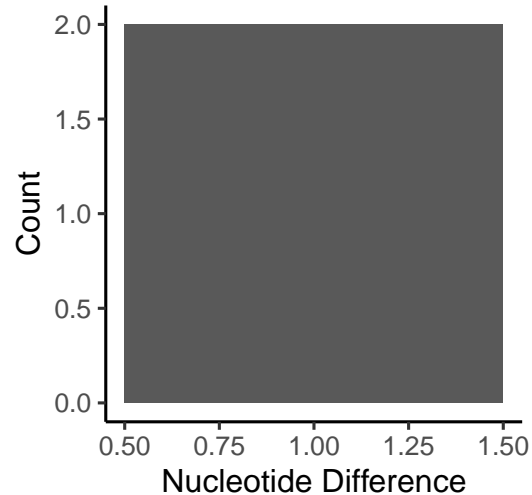
IGLV5-37*01

3 sequences assigned
3 (100%) exact matches, in which:
3 unique CDR3
2 unique J



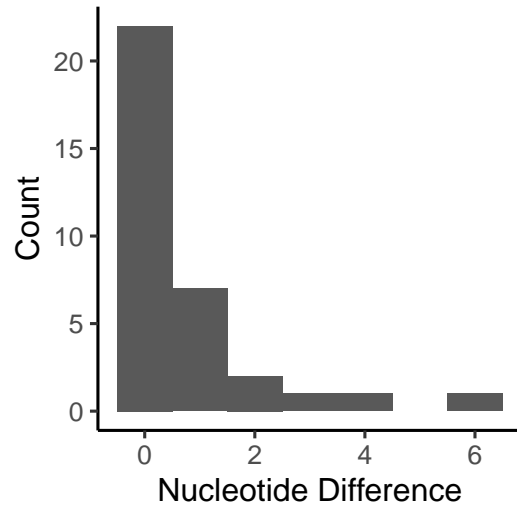
IGLV5-52*01

2 sequences assigned
No exact matches.



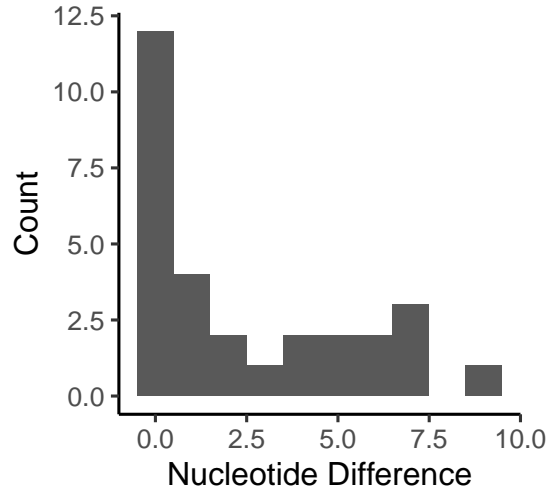
IGLV6-57*02

34 sequences assigned
22 (64.7%) exact matches, in which:
20 unique CDR3
3 unique J



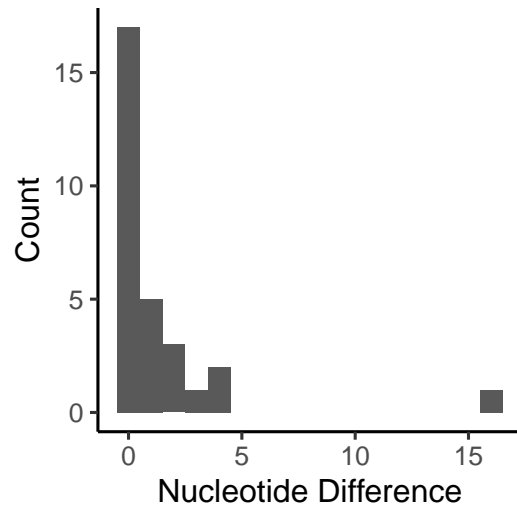
IGLV8-61*01

29 sequences assigned
12 (41.4%) exact matches, in which:
11 unique CDR3
2 unique J



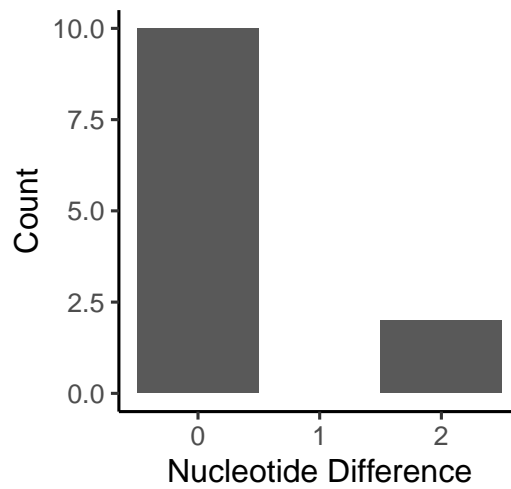
IGLV7-43*01

29 sequences assigned
17 (58.6%) exact matches, in which:
15 unique CDR3
2 unique J



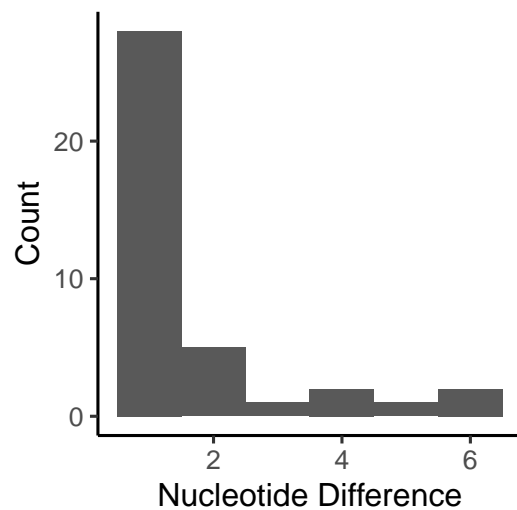
IGLV9-49*01

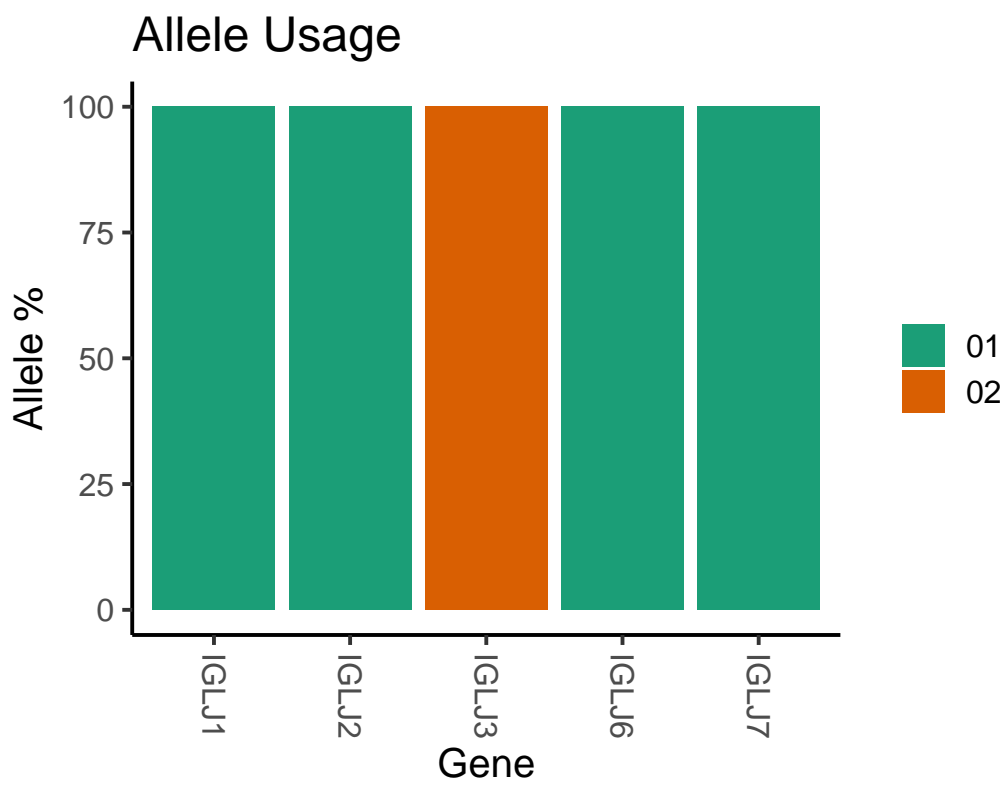
12 sequences assigned
10 (83.3%) exact matches, in which:
10 unique CDR3
2 unique J



IGLV7-46*01

39 sequences assigned
No exact matches.





Warning – no inferred sequences found.