

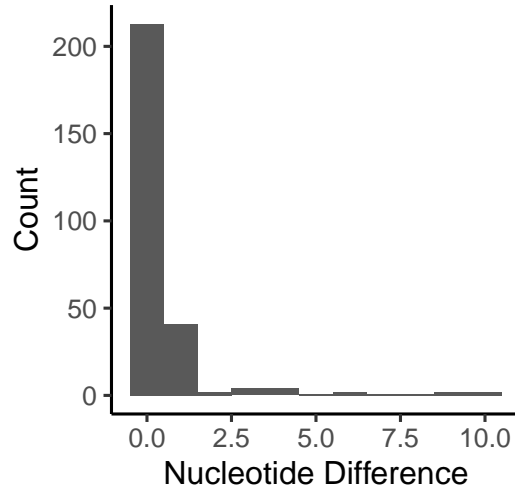
IGLV10-54*01

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



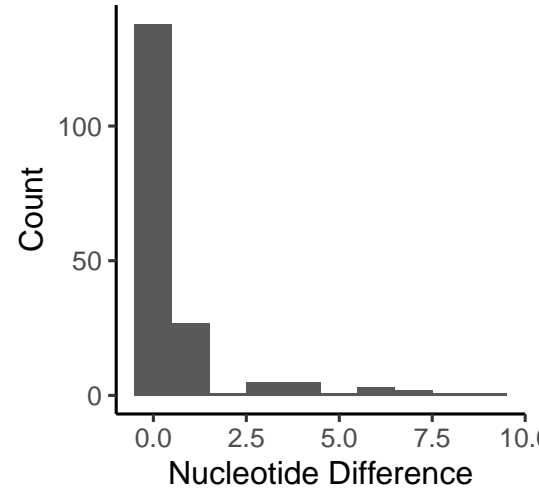
IGLV1-40*01

273 sequences assigned
213 (78%) exact matches, in which:
195 unique CDR3
3 unique J



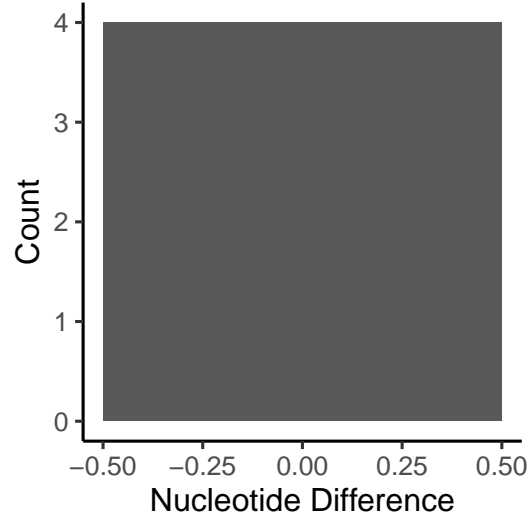
IGLV1-47*01

184 sequences assigned
138 (75%) exact matches, in which:
113 unique CDR3
3 unique J



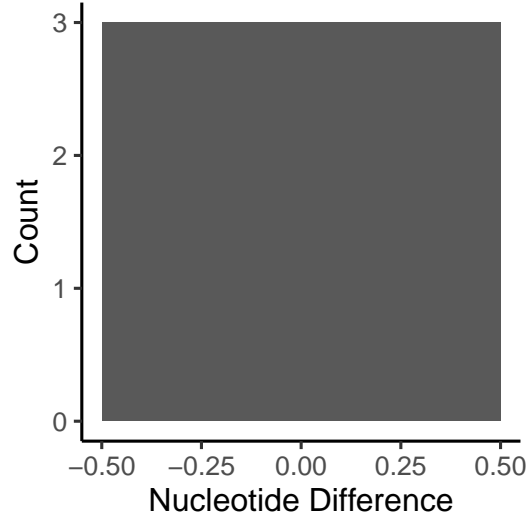
IGLV10-54*02

4 sequences assigned
4 (100%) exact matches, in which:
4 unique CDR3
1 unique J



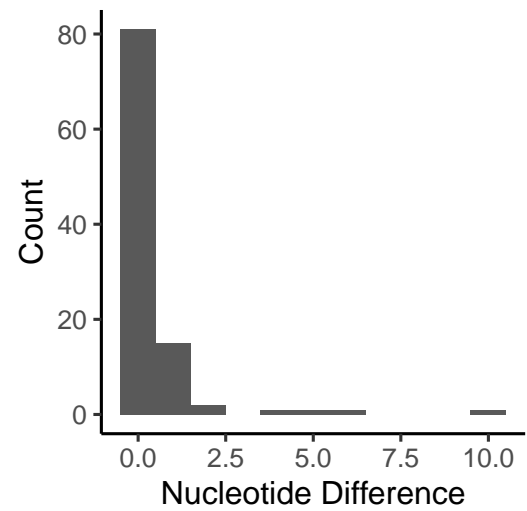
IGLV1-41*01

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



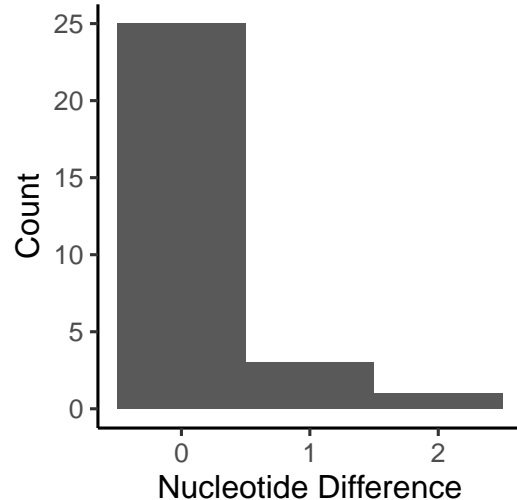
IGLV1-51*01

102 sequences assigned
81 (79.4%) exact matches, in which:
75 unique CDR3
3 unique J



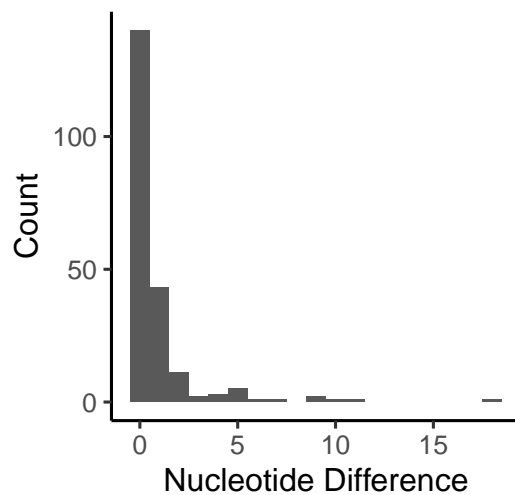
IGLV1-36*01

29 sequences assigned
25 (86.2%) exact matches, in which:
23 unique CDR3
2 unique J



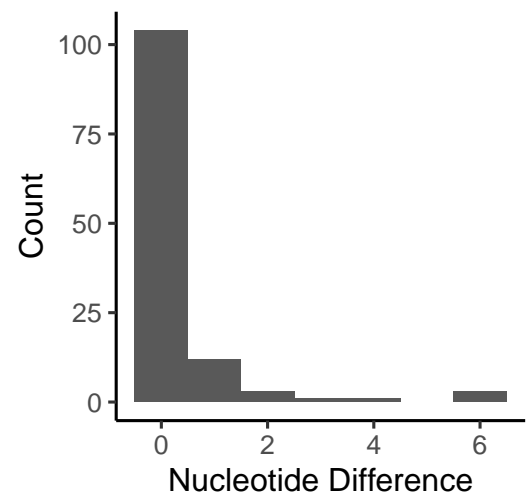
IGLV1-44*01

211 sequences assigned
140 (66.4%) exact matches, in which:
118 unique CDR3
3 unique J



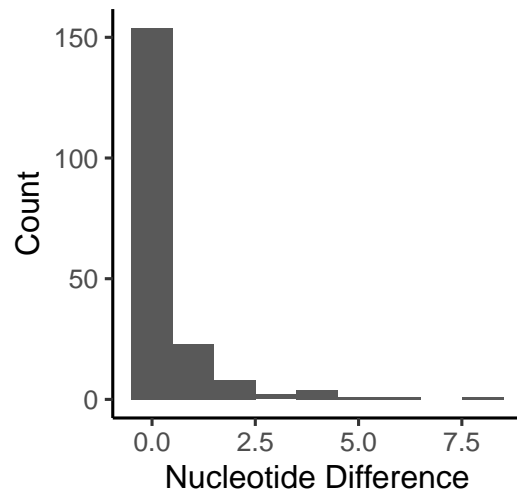
IGLV2-8*01

124 sequences assigned
104 (83.9%) exact matches, in which:
97 unique CDR3
3 unique J



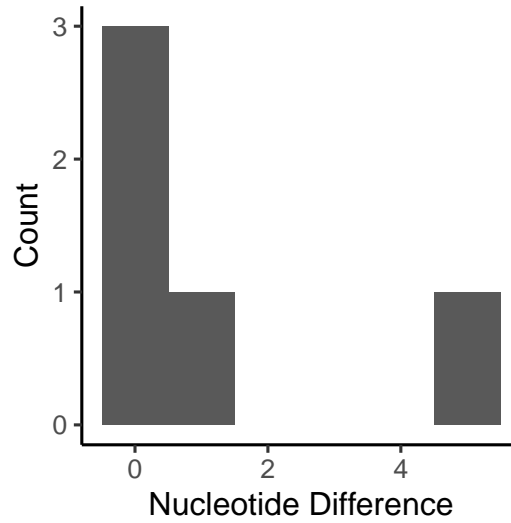
IGLV2-11*01

194 sequences assigned
154 (79.4%) exact matches, in which:
143 unique CDR3
2 unique J



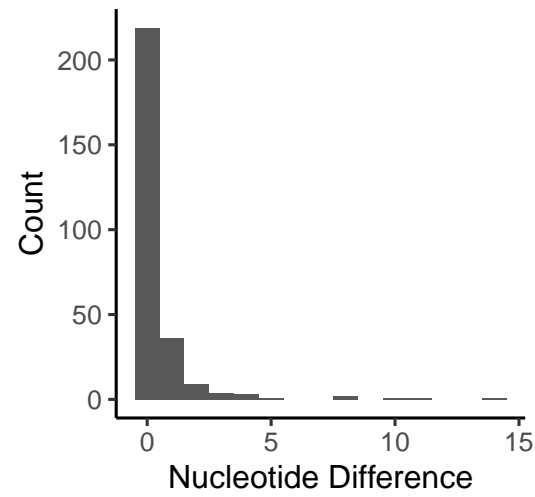
IGLV2-18*02

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



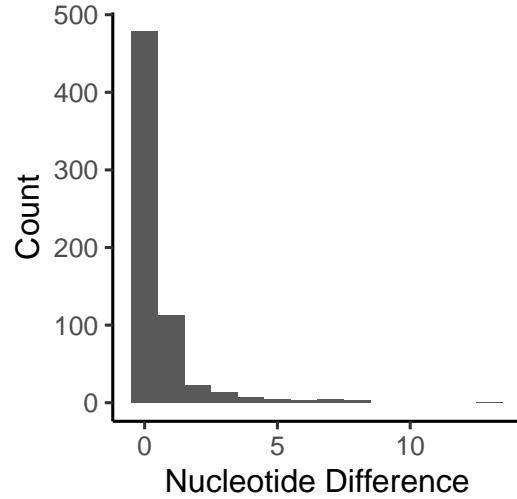
IGLV3-1*01

277 sequences assigned
219 (79.1%) exact matches, in which:
192 unique CDR3
3 unique J



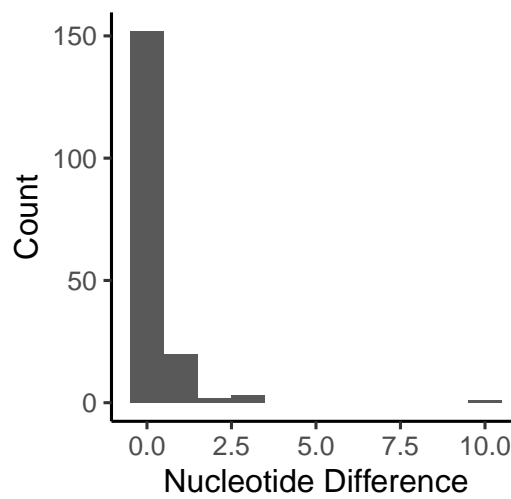
IGLV2-14*01

651 sequences assigned
479 (73.6%) exact matches, in which:
419 unique CDR3
3 unique J



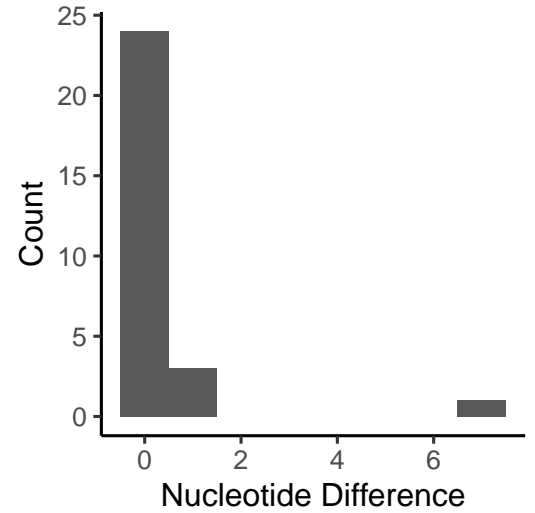
IGLV2-23*02EXT5

178 sequences assigned
152 (85.4%) exact matches, in which:
142 unique CDR3
2 unique J



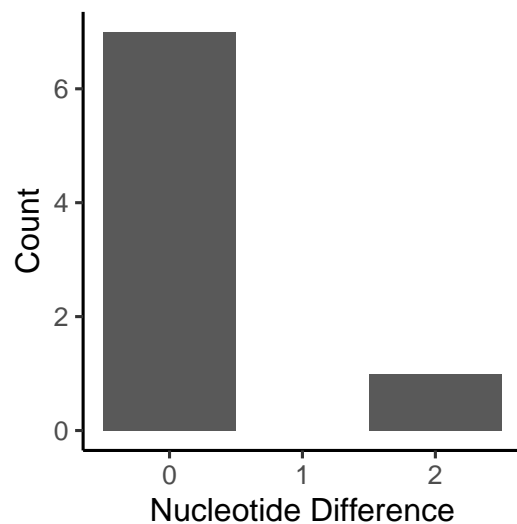
IGLV3-9*01EXT5

28 sequences assigned
24 (85.7%) exact matches, in which:
22 unique CDR3
2 unique J



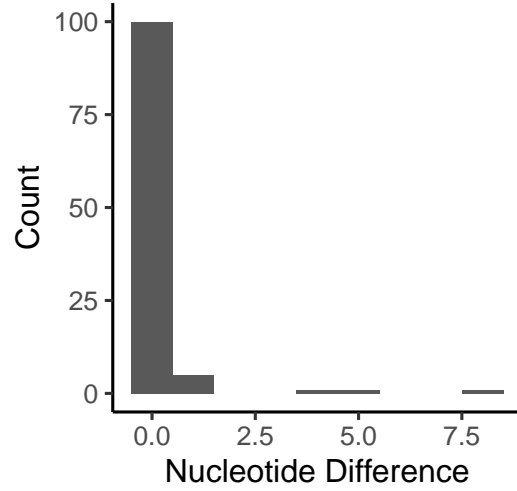
IGLV2-18*01

8 sequences assigned
7 (87.5%) exact matches, in which:
6 unique CDR3
2 unique J



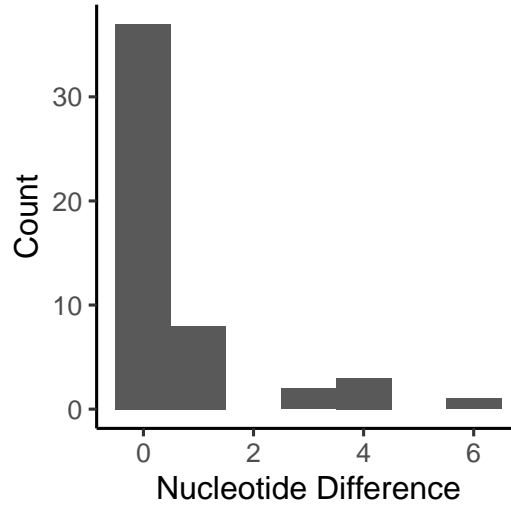
IGLV3-19*01

108 sequences assigned
100 (92.6%) exact matches, in which:
93 unique CDR3
2 unique J



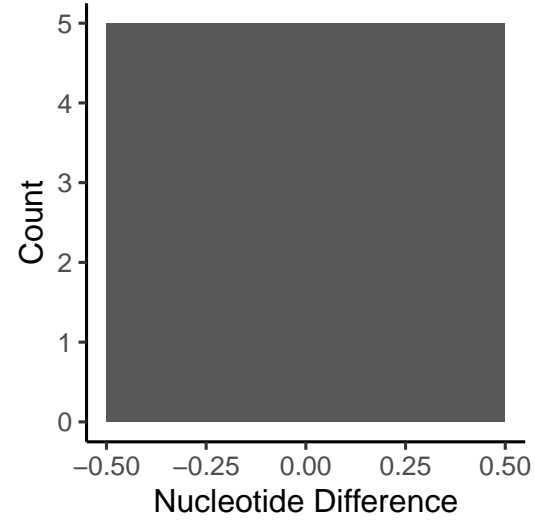
IGLV3-25*02

51 sequences assigned
37 (72.5%) exact matches, in which:
33 unique CDR3
2 unique J



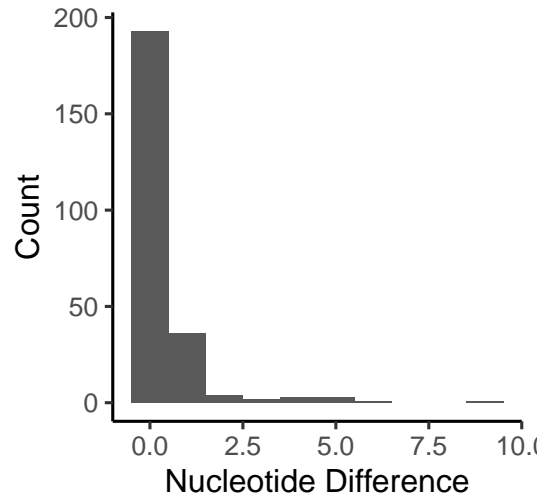
IGLV4-3*01

5 sequences assigned
5 (100%) exact matches, in which:
5 unique CDR3
2 unique J



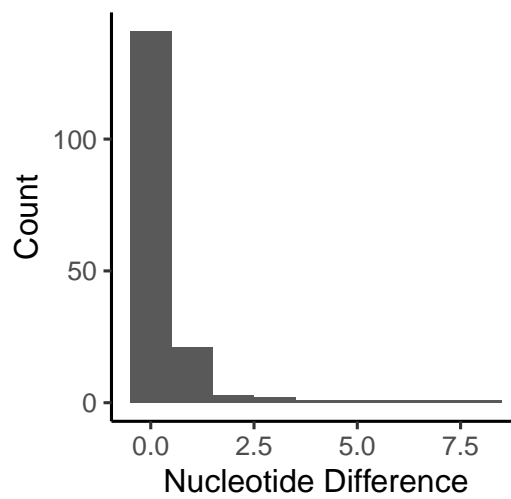
IGLV3-21*02

243 sequences assigned
193 (79.4%) exact matches, in which:
170 unique CDR3
4 unique J



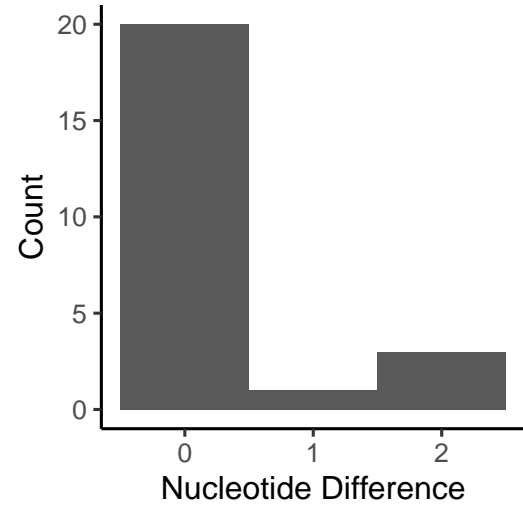
IGLV3-25*03

172 sequences assigned
141 (82%) exact matches, in which:
135 unique CDR3
3 unique J



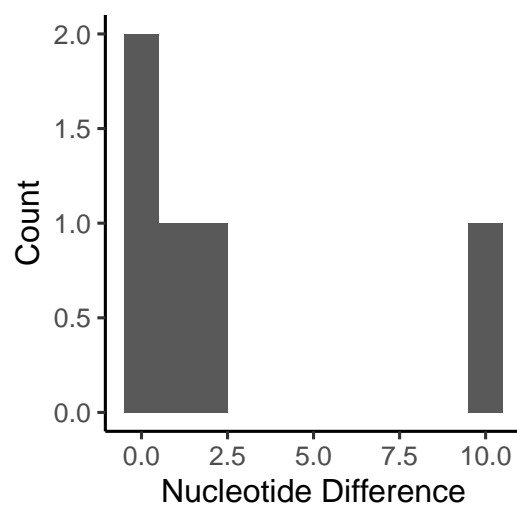
IGLV4-60*02

24 sequences assigned
20 (83.3%) exact matches, in which:
17 unique CDR3
3 unique J



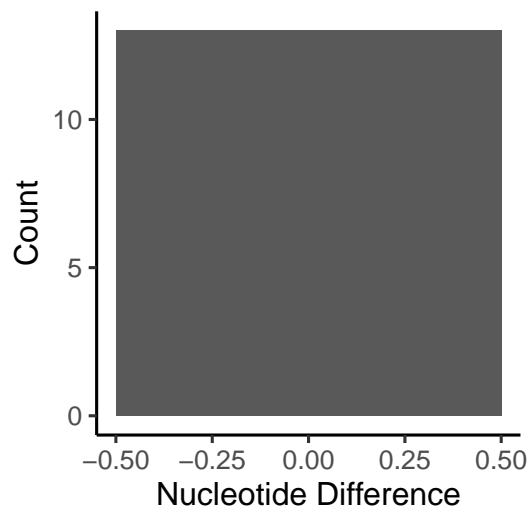
IGLV3-22*01

5 sequences assigned
2 (40%) exact matches, in which:
2 unique CDR3
1 unique J



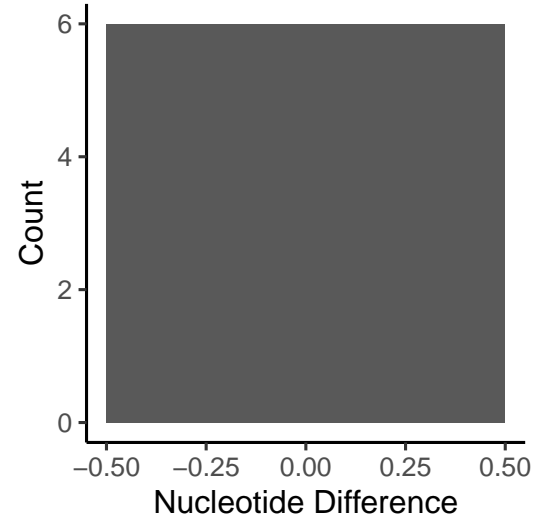
IGLV3-27*01

13 sequences assigned
13 (100%) exact matches, in which:
13 unique CDR3
3 unique J



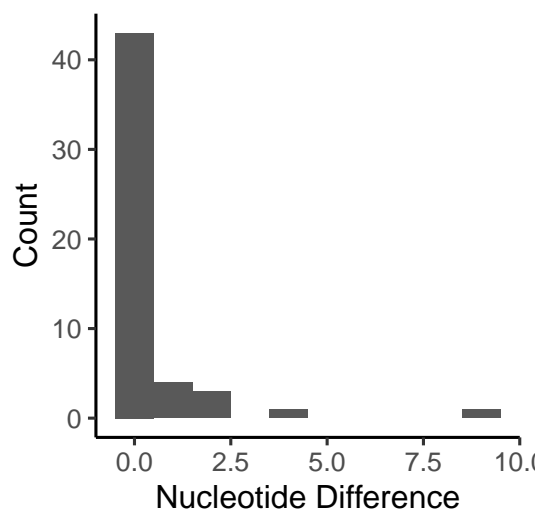
IGLV4-60*03EXT5

6 sequences assigned
6 (100%) exact matches, in which:
6 unique CDR3
2 unique J



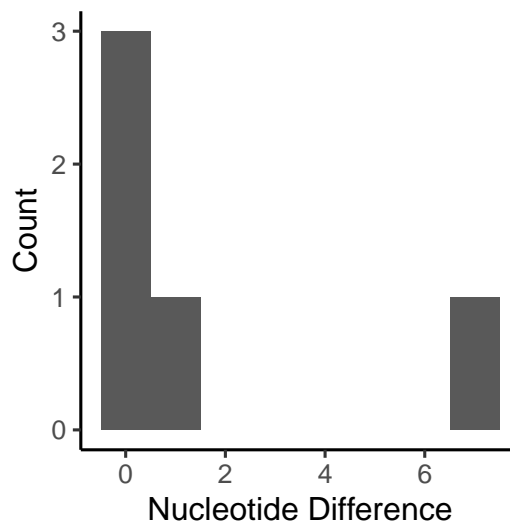
IGLV4-69*01

52 sequences assigned
43 (82.7%) exact matches, in which:
39 unique CDR3
3 unique J



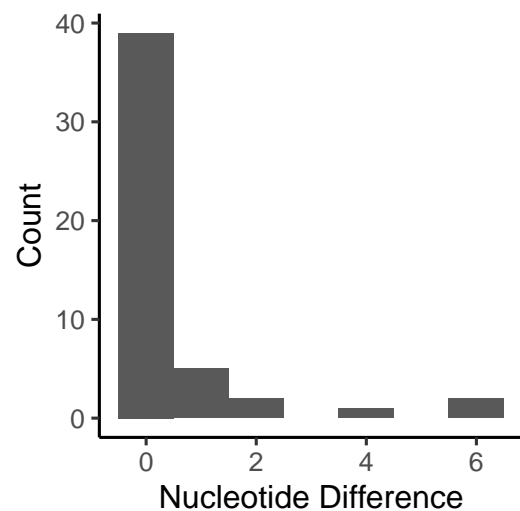
IGLV5-45*02

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



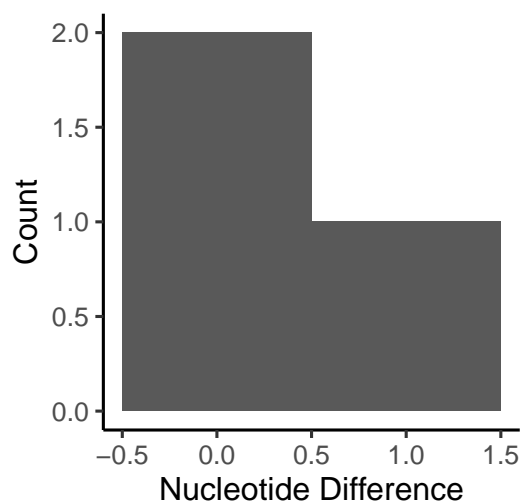
IGLV7-43*01

49 sequences assigned
39 (79.6%) exact matches, in which:
36 unique CDR3
2 unique J



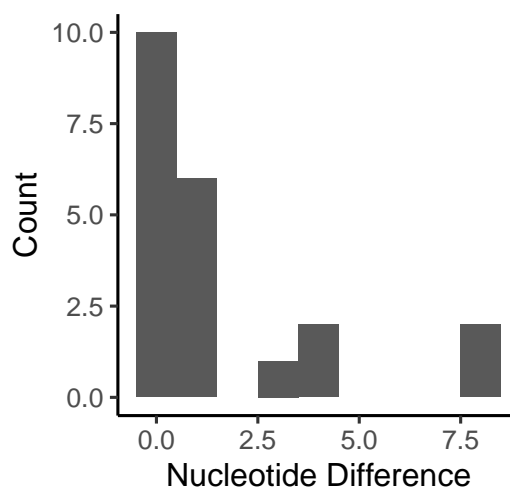
IGLV5-37*01

3 sequences assigned
2 (66.7%) exact matches, in which:
1 unique CDR3
1 unique J



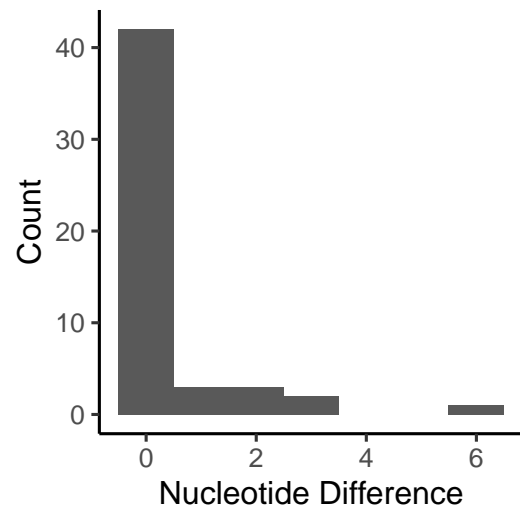
IGLV5-45*03

21 sequences assigned
10 (47.6%) exact matches, in which:
9 unique CDR3
3 unique J



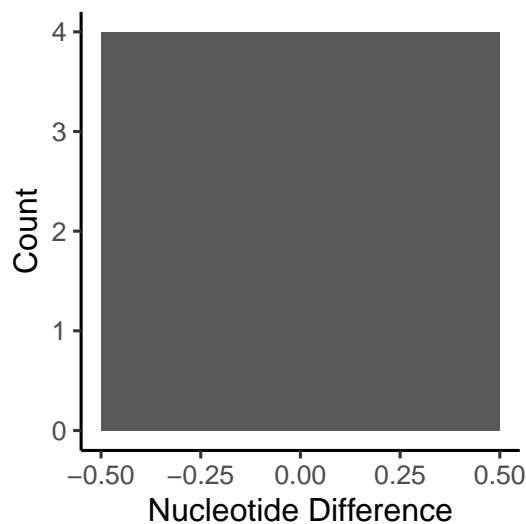
IGLV7-46*01

51 sequences assigned
42 (82.4%) exact matches, in which:
38 unique CDR3
3 unique J



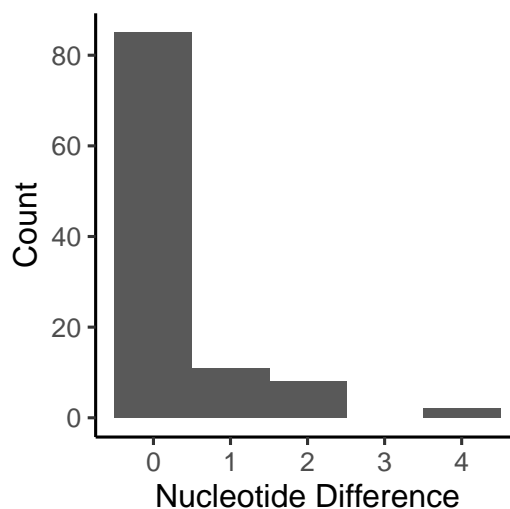
IGLV5-39*01

4 sequences assigned
4 (100%) exact matches, in which:
4 unique CDR3
1 unique J



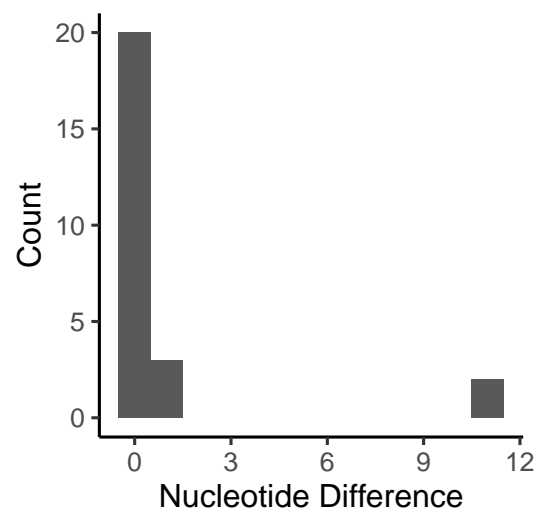
IGLV6-57*01

106 sequences assigned
85 (80.2%) exact matches, in which:
77 unique CDR3
4 unique J



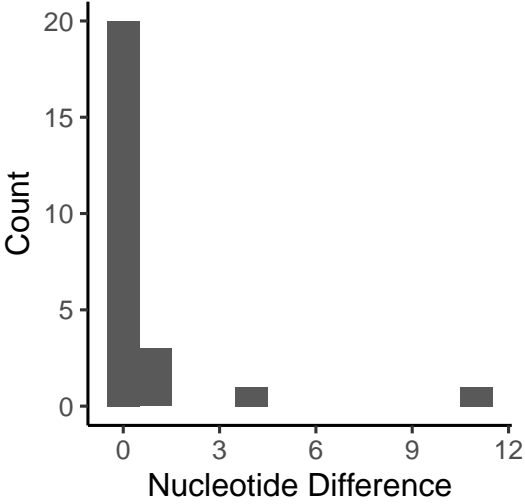
IGLV8-61*01

25 sequences assigned
20 (80%) exact matches, in which:
18 unique CDR3
4 unique J



IGLV9-49*01

25 sequences assigned
20 (80%) exact matches, in which:
19 unique CDR3
3 unique J



Allele Usage



Warning – no inferred sequences found.