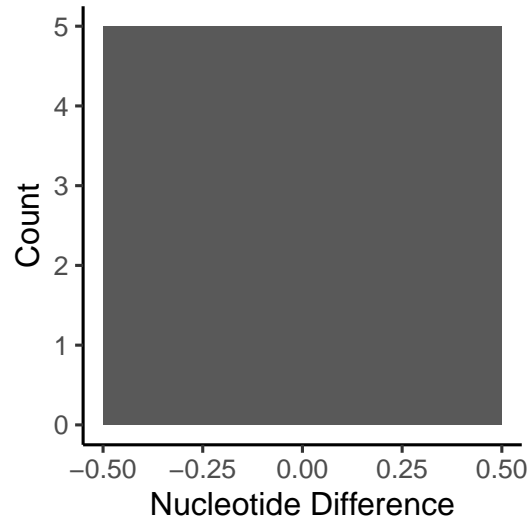


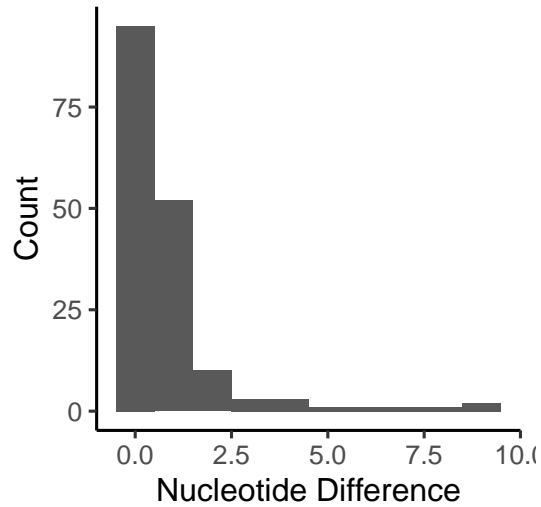
IGLV10-54*01

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



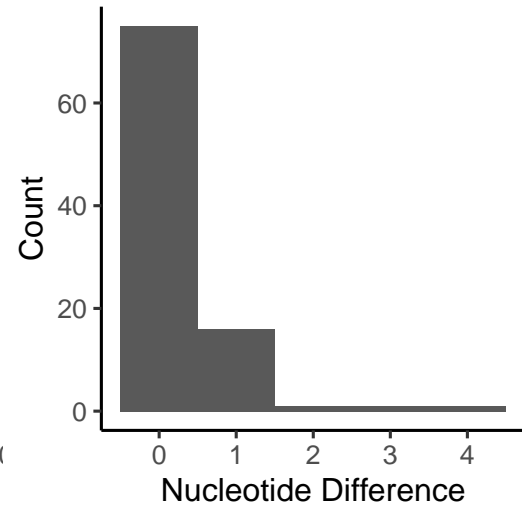
IGLV1-44*01

169 sequences assigned
95 (56.2%) exact matches, in which:
86 unique CDR3
3 unique J



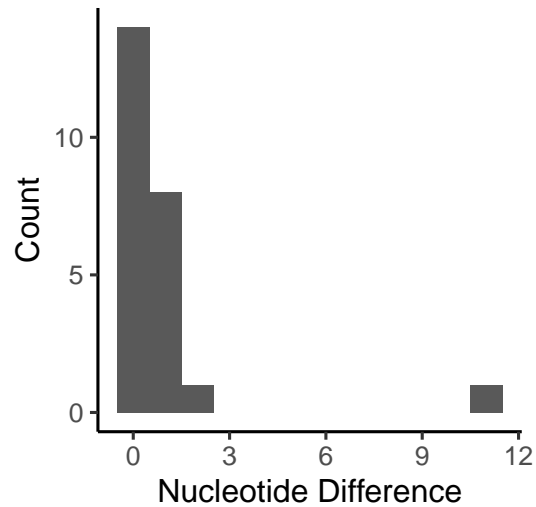
IGLV1-51*01

94 sequences assigned
75 (79.8%) exact matches, in which:
59 unique CDR3
4 unique J



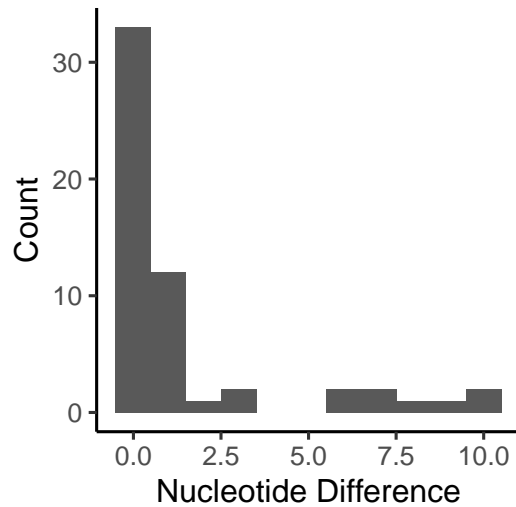
IGLV1-36*01

24 sequences assigned
14 (58.3%) exact matches, in which:
11 unique CDR3
3 unique J



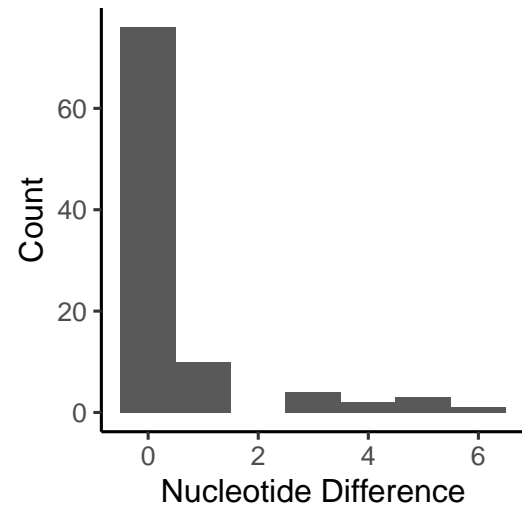
IGLV1-47*01

56 sequences assigned
33 (58.9%) exact matches, in which:
28 unique CDR3
4 unique J



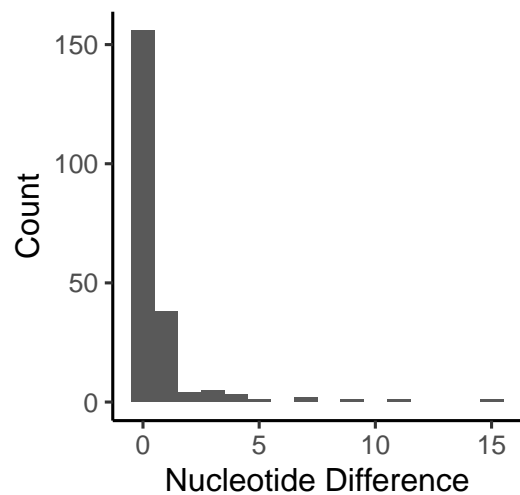
IGLV2-8*01

96 sequences assigned
76 (79.2%) exact matches, in which:
69 unique CDR3
3 unique J



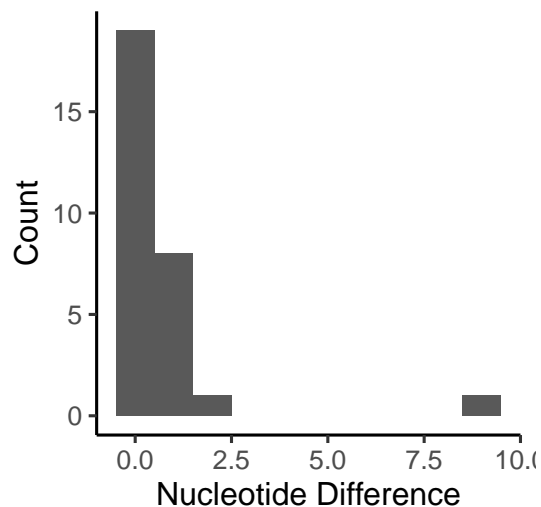
IGLV1-40*01

212 sequences assigned
156 (73.6%) exact matches, in which:
145 unique CDR3
4 unique J



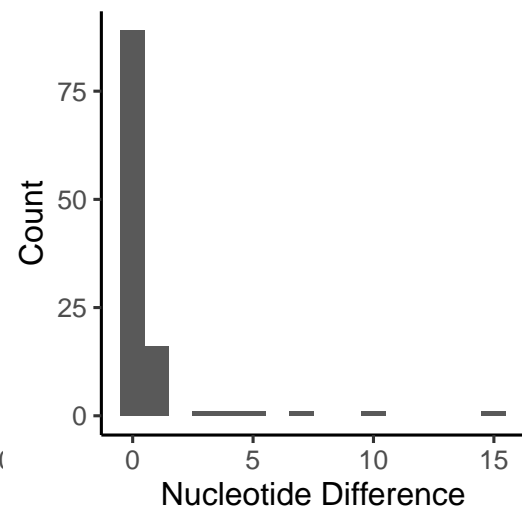
IGLV1-47*02

29 sequences assigned
19 (65.5%) exact matches, in which:
17 unique CDR3
2 unique J



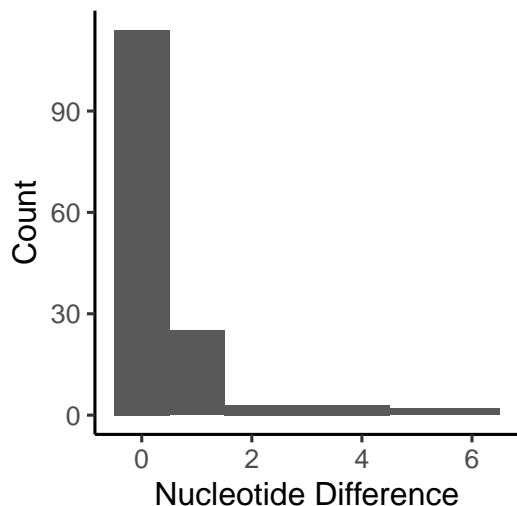
IGLV2-11*01

111 sequences assigned
89 (80.2%) exact matches, in which:
87 unique CDR3
3 unique J



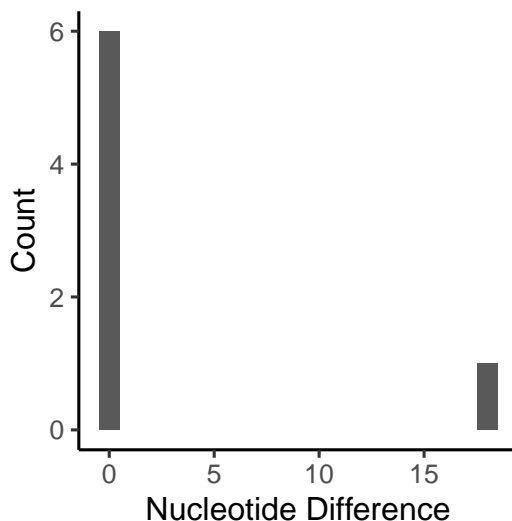
IGLV2-14*01

152 sequences assigned
114 (75%) exact matches, in which:
109 unique CDR3
3 unique J



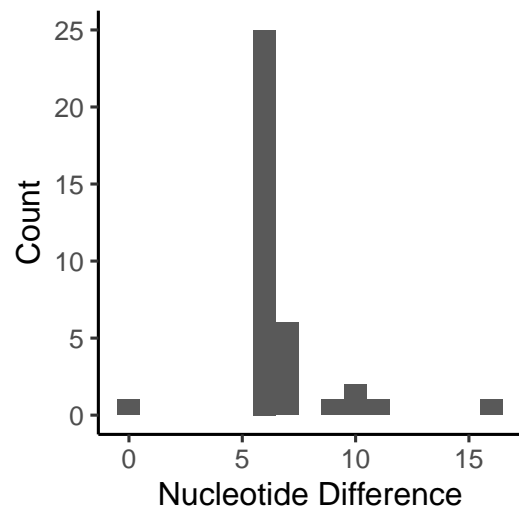
IGLV2-18*02

7 sequences assigned
6 (85.7%) exact matches, in which:
6 unique CDR3
2 unique J



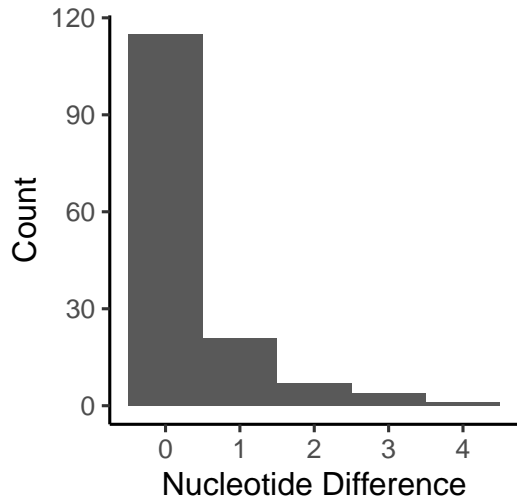
IGLV3-9*02

37 sequences assigned
1 (2.7%) exact matches, in which:
1 unique CDR3
1 unique J



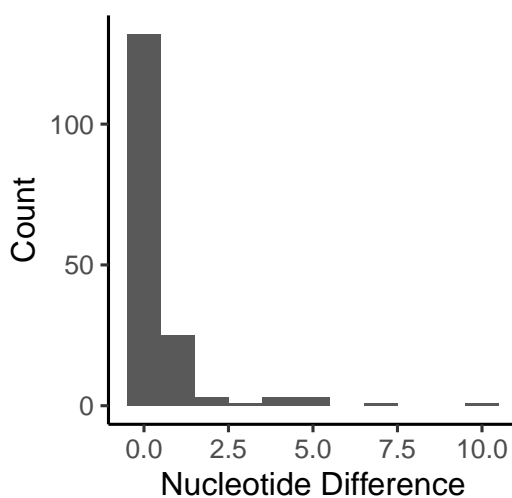
IGLV2-14*04EXT3

148 sequences assigned
115 (77.7%) exact matches, in which:
104 unique CDR3
2 unique J



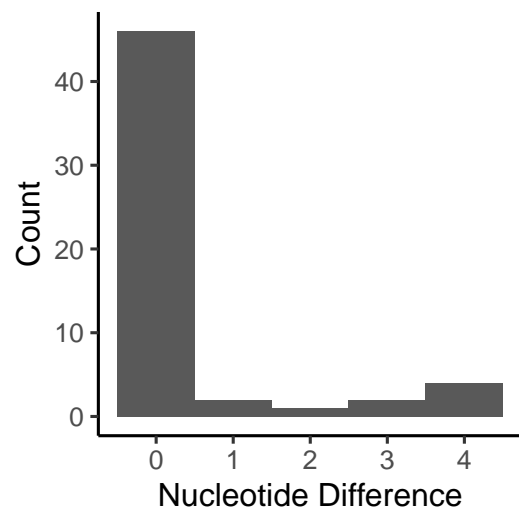
IGLV2-23*01

169 sequences assigned
132 (78.1%) exact matches, in which:
122 unique CDR3
4 unique J



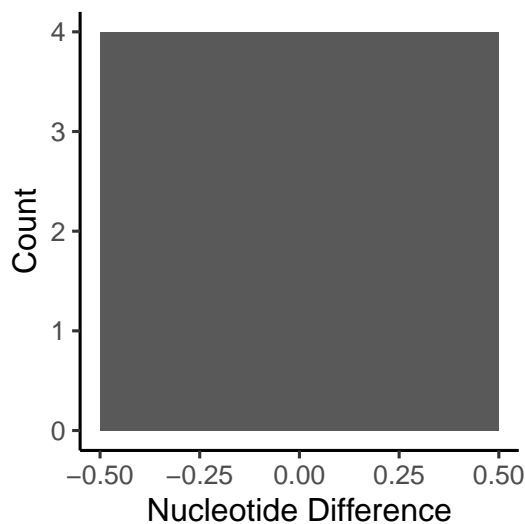
IGLV3-10*01

55 sequences assigned
46 (83.6%) exact matches, in which:
41 unique CDR3
3 unique J



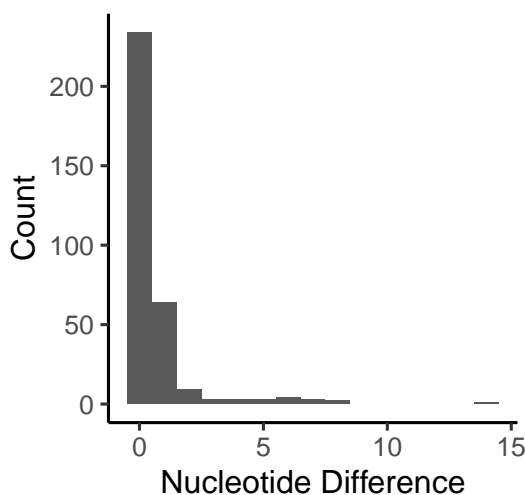
IGLV2-18*01

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



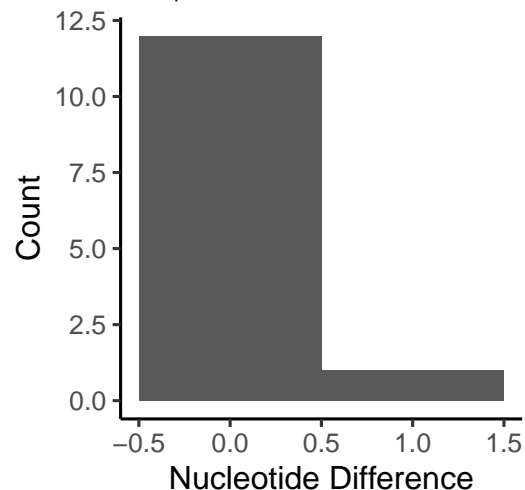
IGLV3-1*01

326 sequences assigned
234 (71.8%) exact matches, in which:
203 unique CDR3
3 unique J



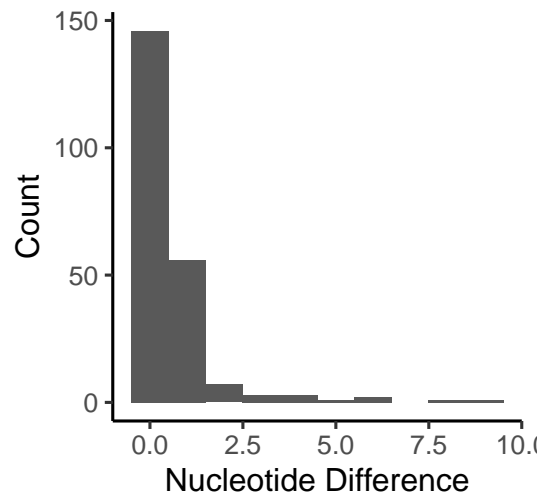
IGLV3-16*01

13 sequences assigned
12 (92.3%) exact matches, in which:
12 unique CDR3
4 unique J



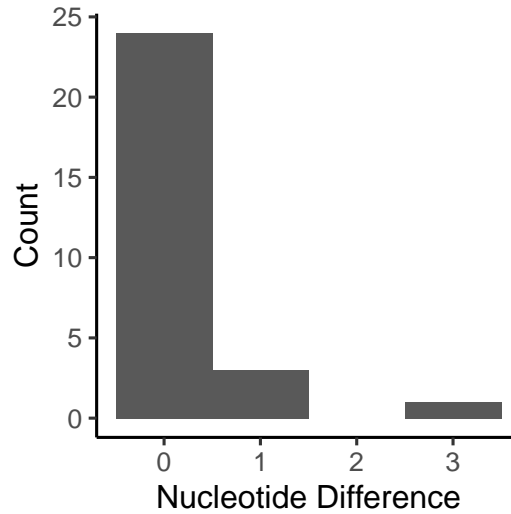
IGLV3–19*01

220 sequences assigned
146 (66.4%) exact matches, in which:
124 unique CDR3
4 unique J



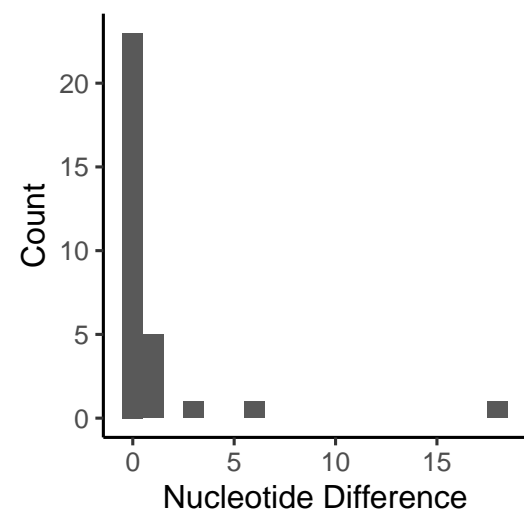
IGLV3–27*01

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



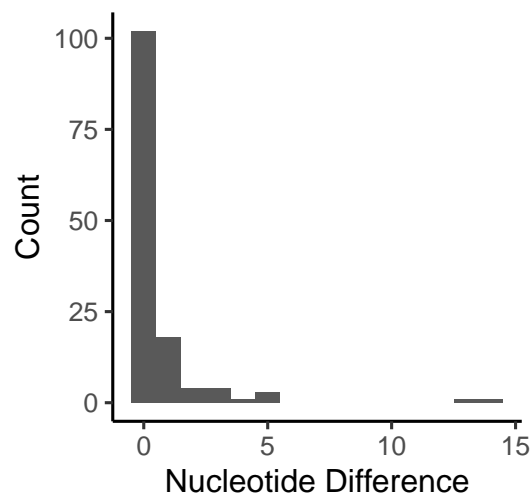
IGLV4–69*01

31 sequences assigned
23 (74.2%) exact matches, in which:
20 unique CDR3
2 unique J



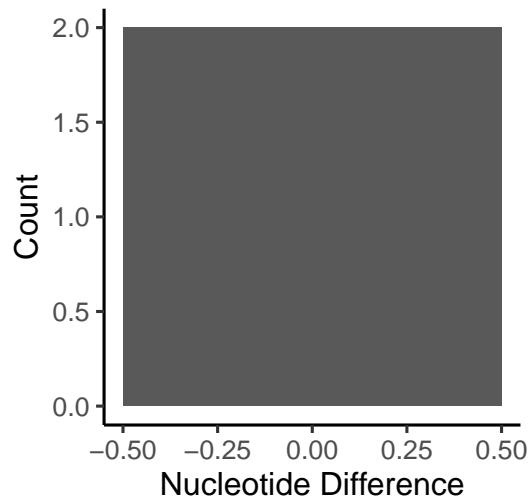
IGLV3–21*04

134 sequences assigned
102 (76.1%) exact matches, in which:
91 unique CDR3
4 unique J



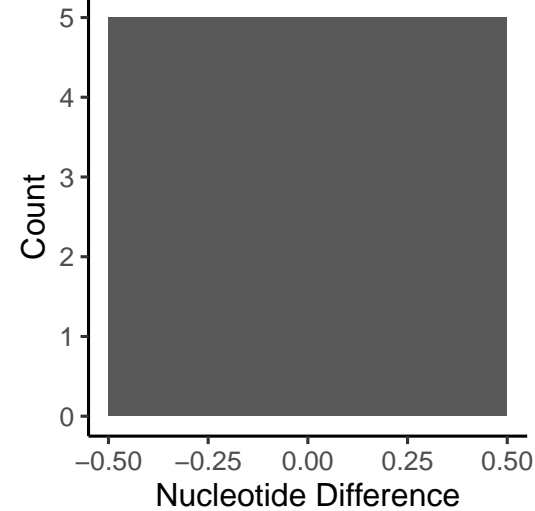
IGLV4–3*01

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



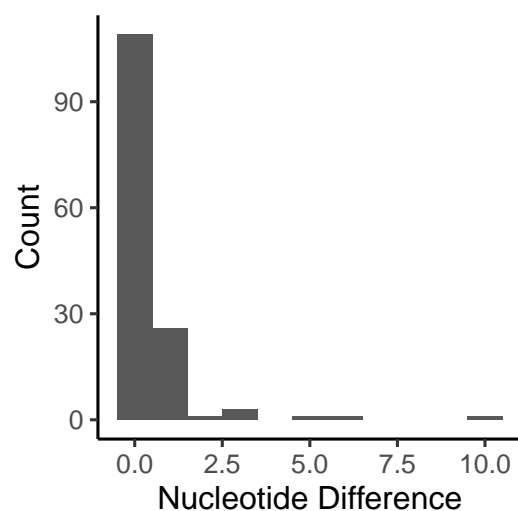
IGLV5–37*01

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



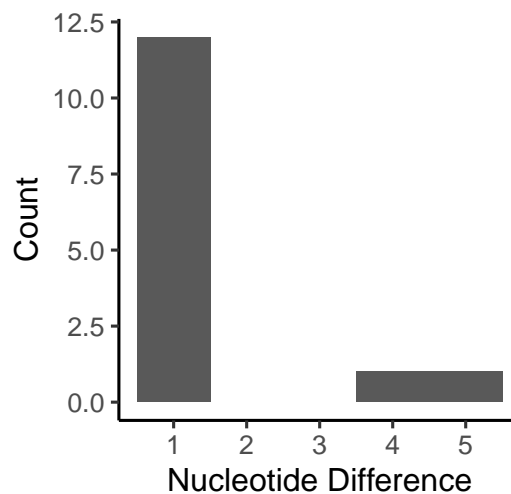
IGLV3–25*03

142 sequences assigned
109 (76.8%) exact matches, in which:
100 unique CDR3
4 unique J



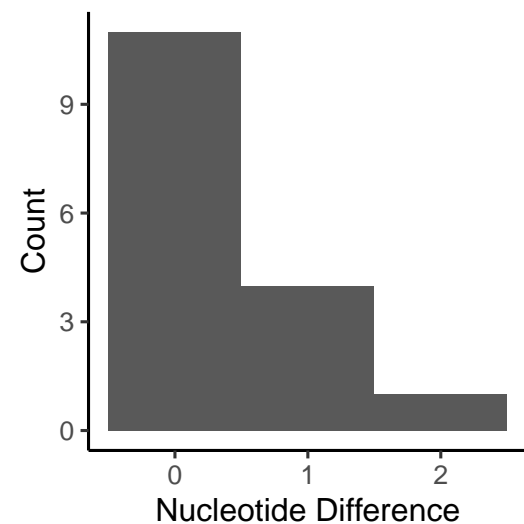
IGLV4–60*02

14 sequences assigned
No exact matches.



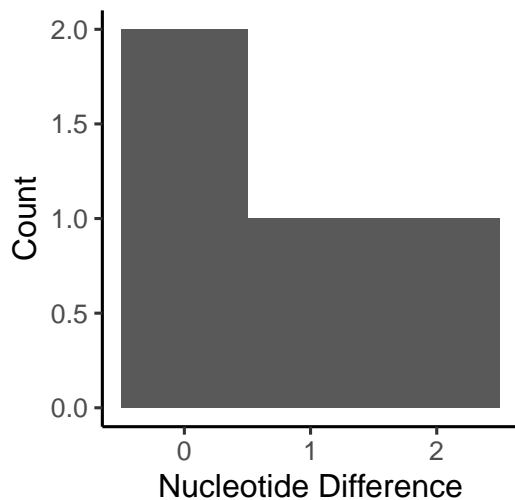
IGLV5–45*02

16 sequences assigned
11 (68.8%) exact matches, in which:
9 unique CDR3
2 unique J



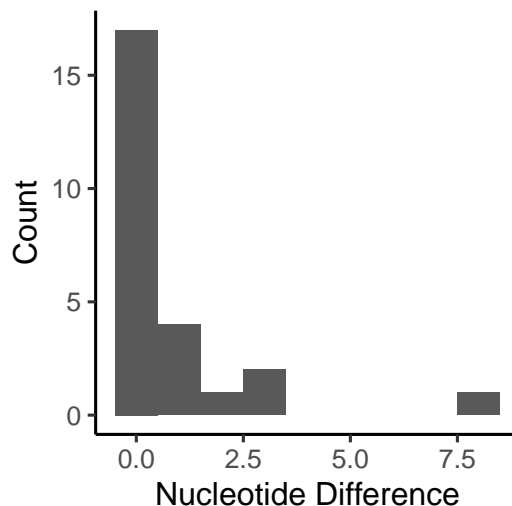
IGLV5-52*01

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



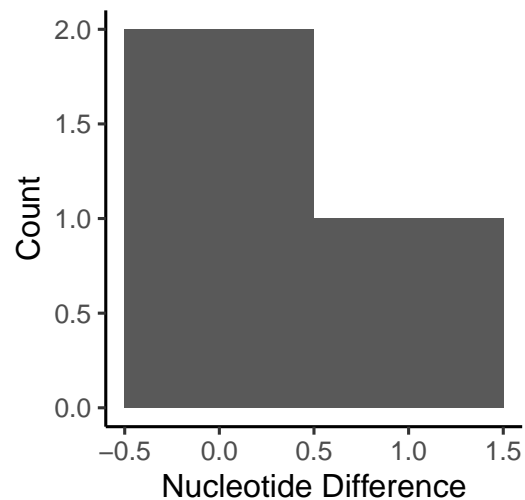
IGLV7-43*01

25 sequences assigned
17 (68%) exact matches, in which:
15 unique CDR3
3 unique J



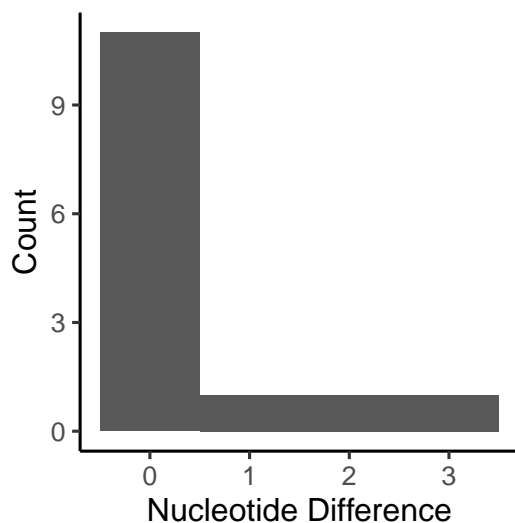
IGLV9-49*01

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



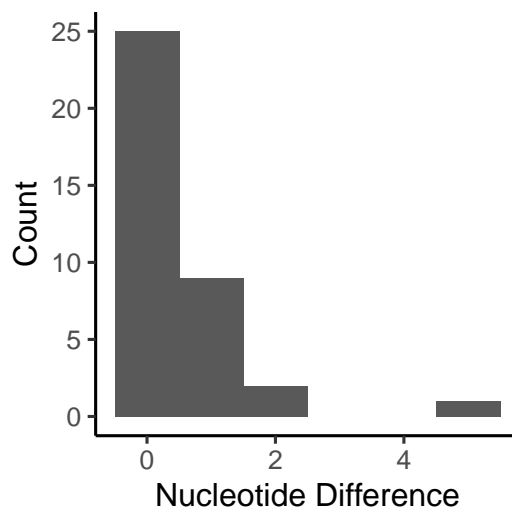
IGLV6-57*02

14 sequences assigned
11 (78.6%) exact matches, in which:
11 unique CDR3
2 unique J



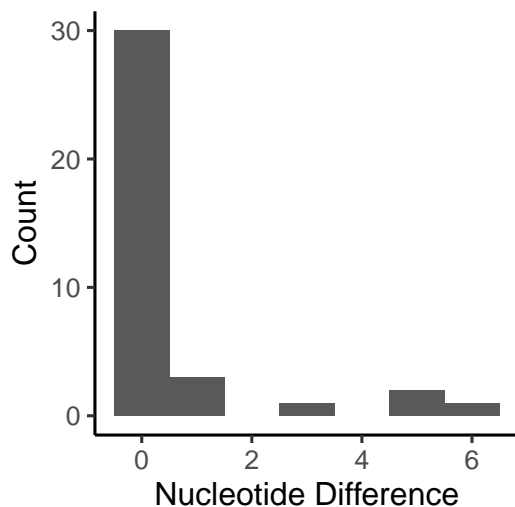
IGLV7-46*01

37 sequences assigned
25 (67.6%) exact matches, in which:
25 unique CDR3
3 unique J



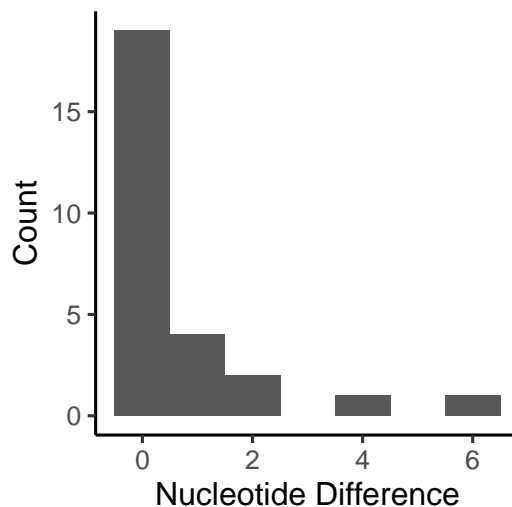
IGLV6-57*03

37 sequences assigned
30 (81.1%) exact matches, in which:
27 unique CDR3
3 unique J

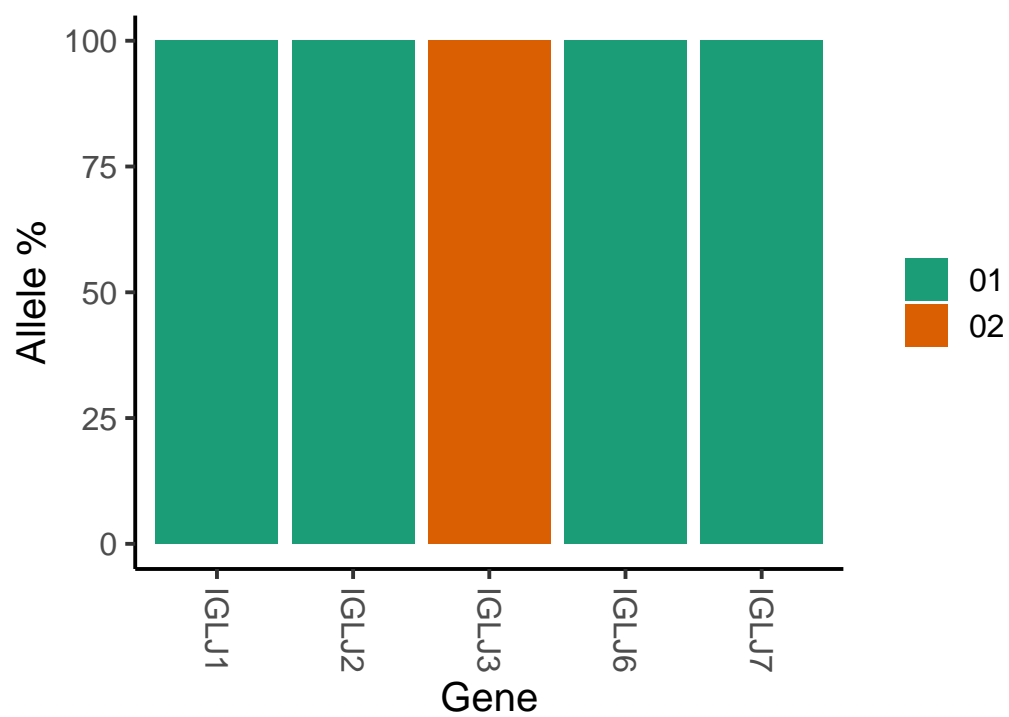


IGLV8-61*01

28 sequences assigned
19 (67.9%) exact matches, in which:
18 unique CDR3
4 unique J



Allele Usage



Warning – no inferred sequences found.