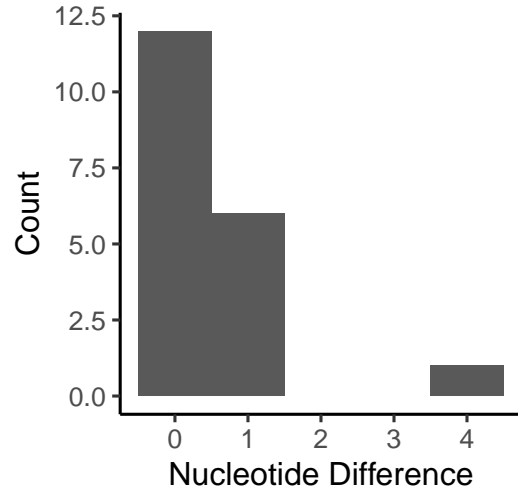


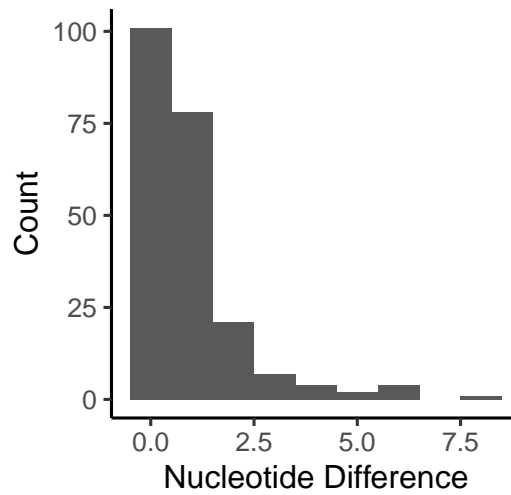
IGLV10-54*01

19 sequences assigned
12 (63.2%) exact matches, in which:
11 unique CDR3
1 unique J



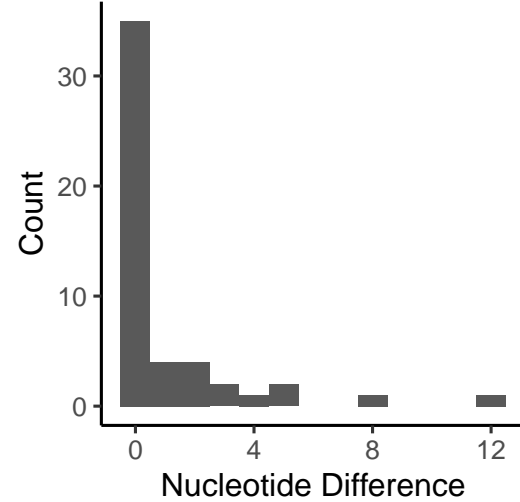
IGLV1-44*01

218 sequences assigned
101 (46.3%) exact matches, in which:
85 unique CDR3
3 unique J



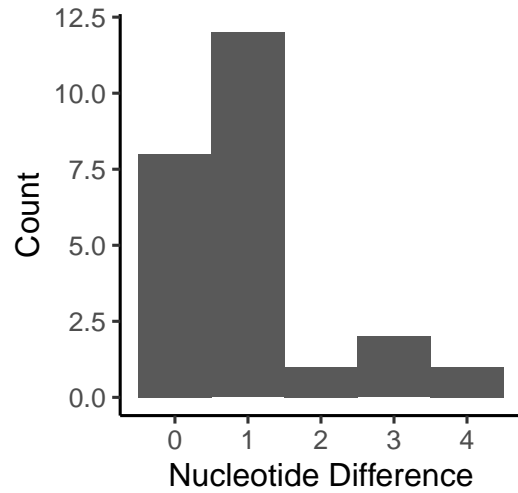
IGLV2-8*01

50 sequences assigned
35 (70%) exact matches, in which:
30 unique CDR3
3 unique J



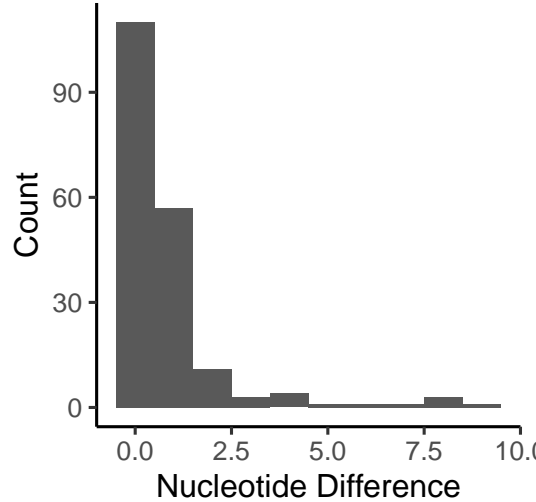
IGLV1-36*01

24 sequences assigned
8 (33.3%) exact matches, in which:
8 unique CDR3
2 unique J



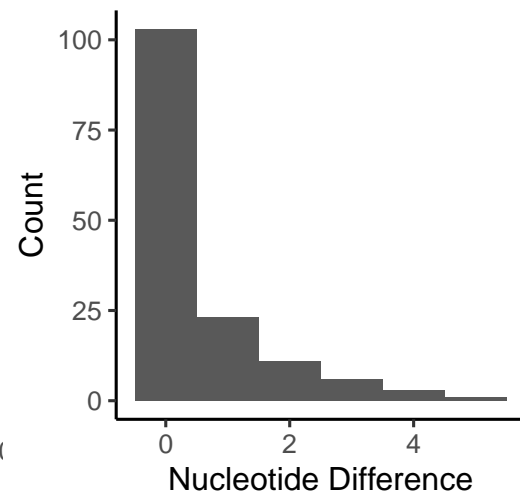
IGLV1-47*01

192 sequences assigned
110 (57.3%) exact matches, in which:
101 unique CDR3
3 unique J



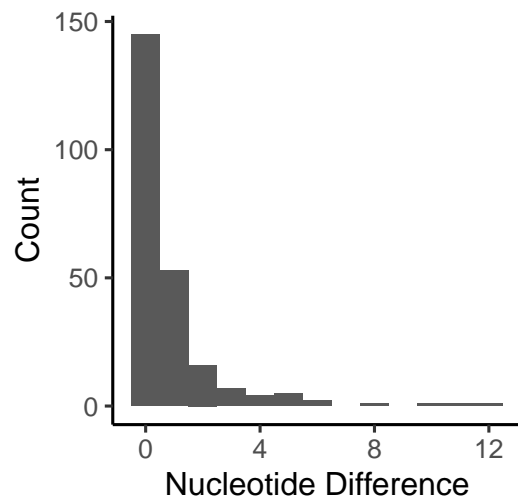
IGLV2-11*01

147 sequences assigned
103 (70.1%) exact matches, in which:
93 unique CDR3
3 unique J



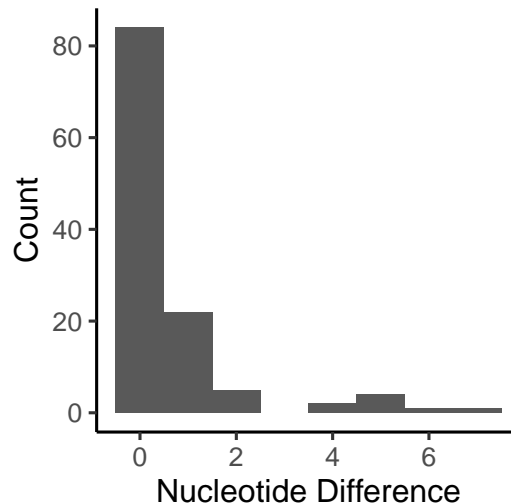
IGLV1-40*01

237 sequences assigned
145 (61.2%) exact matches, in which:
123 unique CDR3
2 unique J



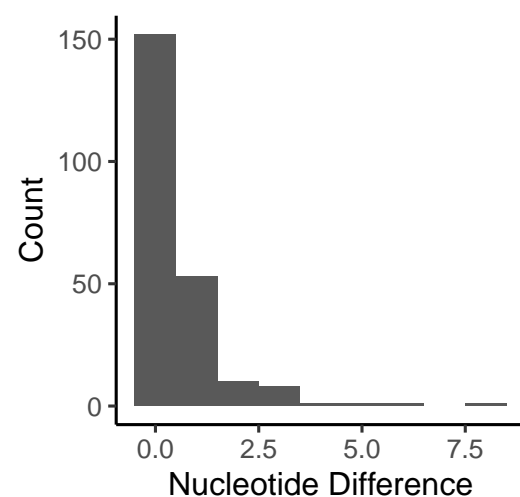
IGLV1-51*01

119 sequences assigned
84 (70.6%) exact matches, in which:
65 unique CDR3
3 unique J



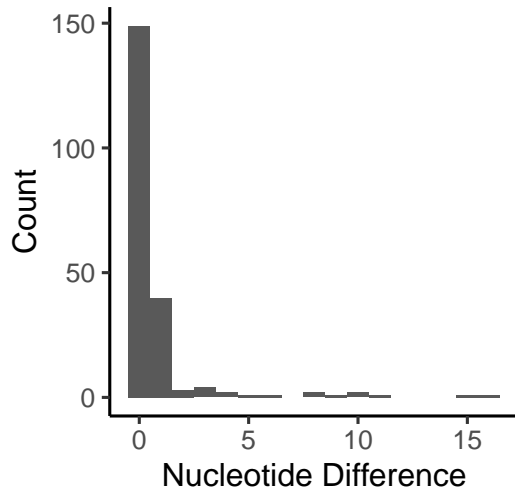
IGLV2-14*01

227 sequences assigned
152 (67%) exact matches, in which:
137 unique CDR3
2 unique J



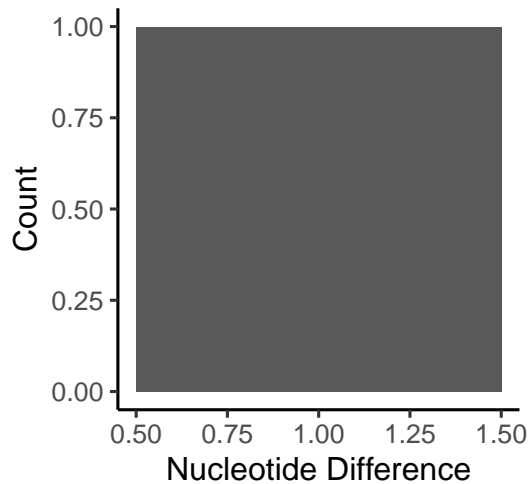
IGLV2-14*03EXT3

208 sequences assigned
149 (71.6%) exact matches, in which:
143 unique CDR3
3 unique J



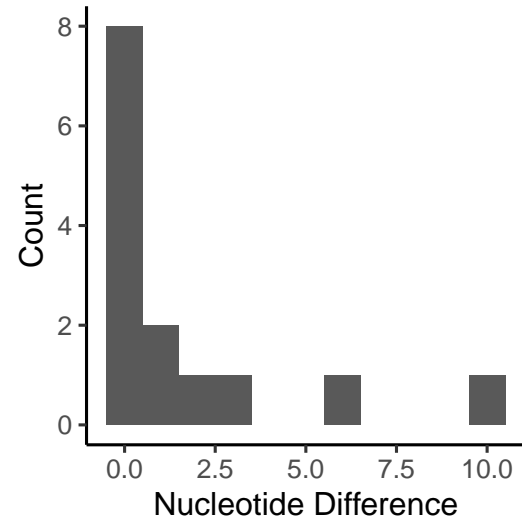
IGLV2-18*03

1 sequences assigned
No exact matches.



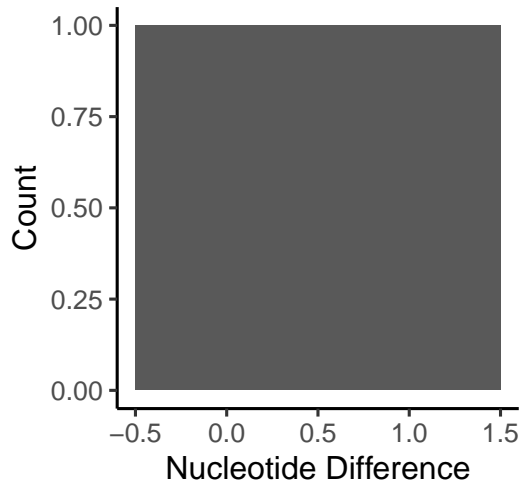
IGLV3-9*01EXT5

14 sequences assigned
8 (57.1%) exact matches, in which:
7 unique CDR3
2 unique J



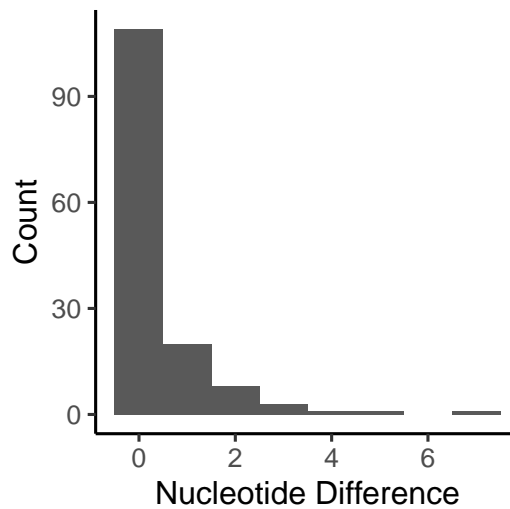
IGLV2-18*01

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



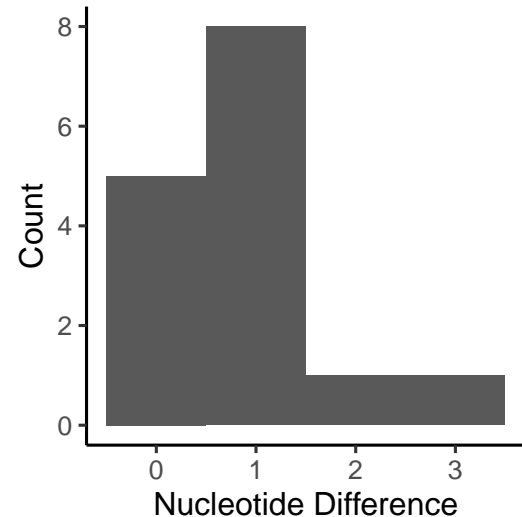
IGLV2-23*02EXT5

143 sequences assigned
109 (76.2%) exact matches, in which:
100 unique CDR3
3 unique J



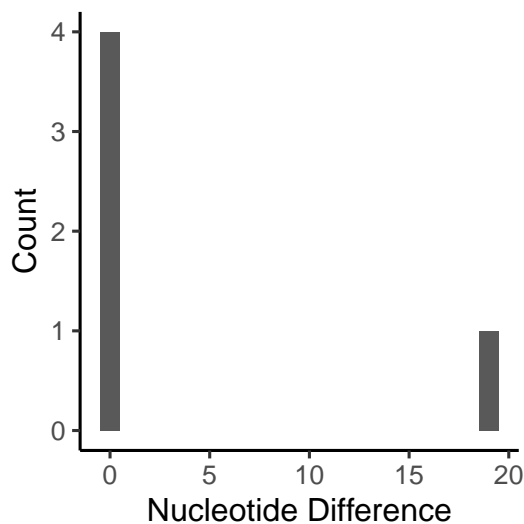
IGLV3-10*01

15 sequences assigned
5 (33.3%) exact matches, in which:
5 unique CDR3
2 unique J



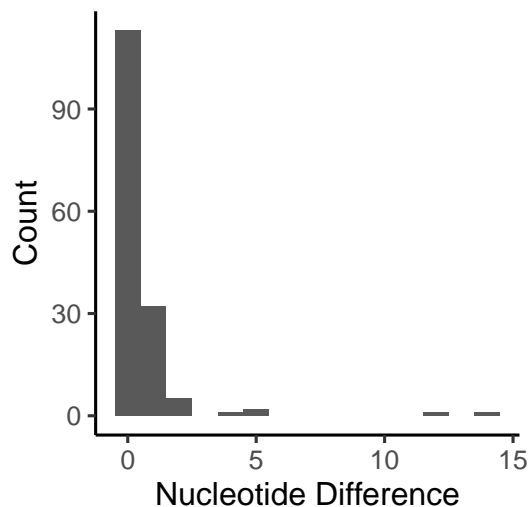
IGLV2-18*02

5 sequences assigned
4 (80%) exact matches, in which:
2 unique CDR3
2 unique J



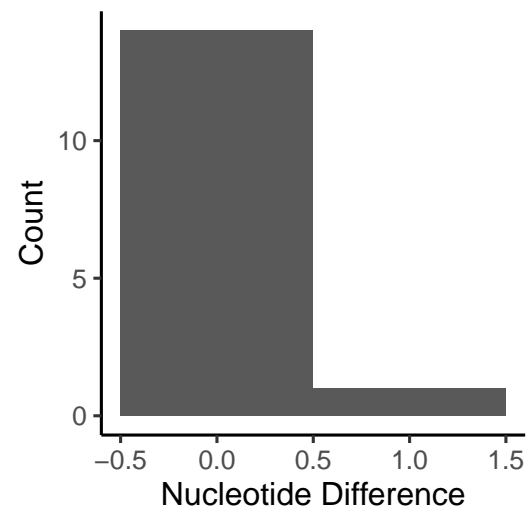
IGLV3-1*01

155 sequences assigned
113 (72.9%) exact matches, in which:
105 unique CDR3
2 unique J



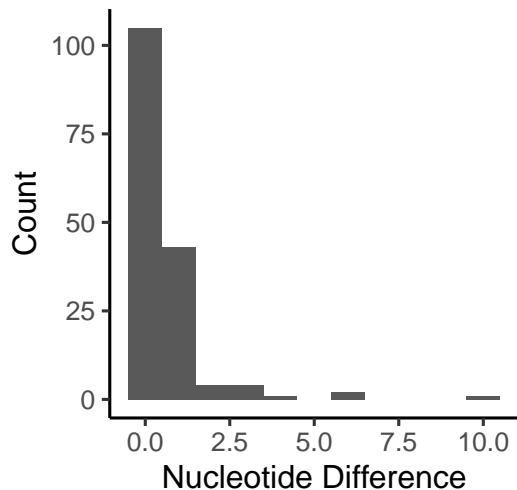
IGLV3-16*01

15 sequences assigned
14 (93.3%) exact matches, in which:
13 unique CDR3
3 unique J



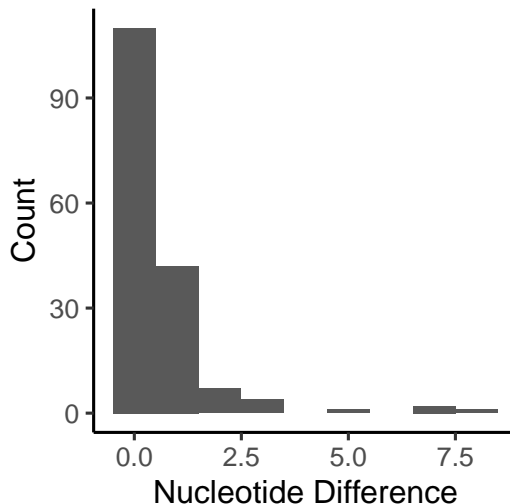
IGLV3-19*01

160 sequences assigned
105 (65.6%) exact matches, in which:
90 unique CDR3
2 unique J



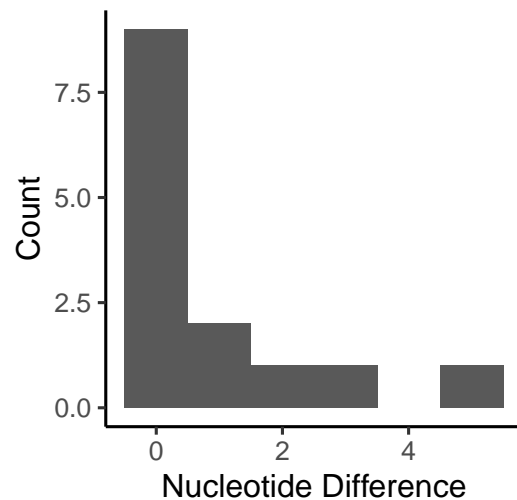
IGLV3-25*03

167 sequences assigned
110 (65.9%) exact matches, in which:
94 unique CDR3
3 unique J



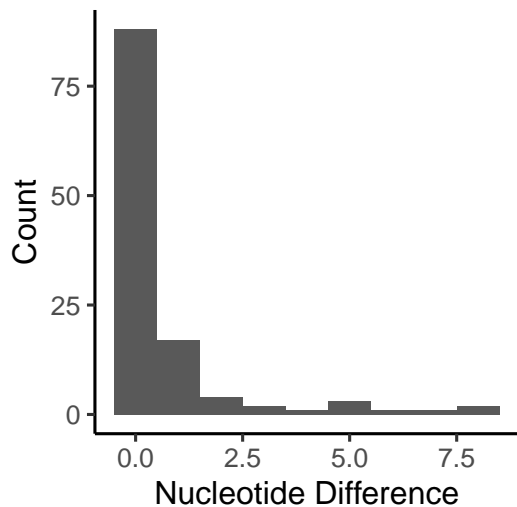
IGLV4-60*03EXT5

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



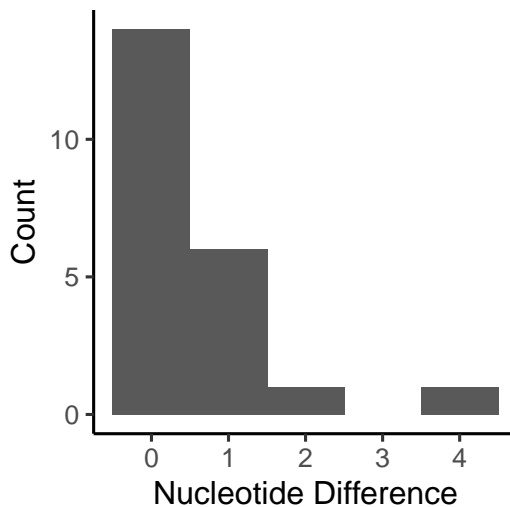
IGLV3-21*02

119 sequences assigned
88 (73.9%) exact matches, in which:
73 unique CDR3
3 unique J



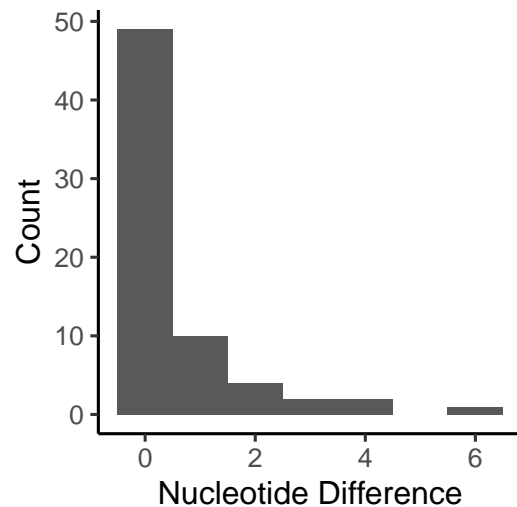
IGLV3-27*01

22 sequences assigned
14 (63.6%) exact matches, in which:
14 unique CDR3
1 unique J



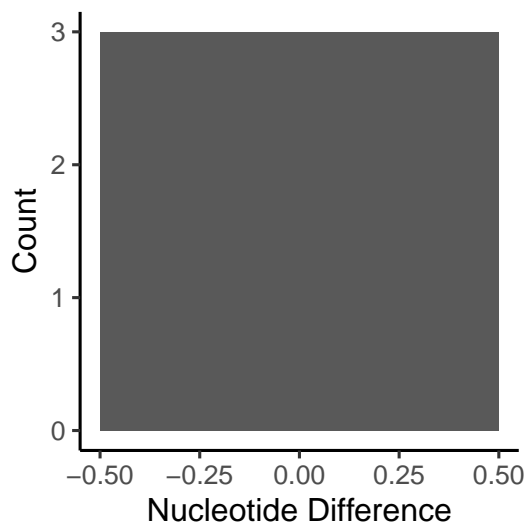
IGLV4-69*01

68 sequences assigned
49 (72.1%) exact matches, in which:
41 unique CDR3
2 unique J



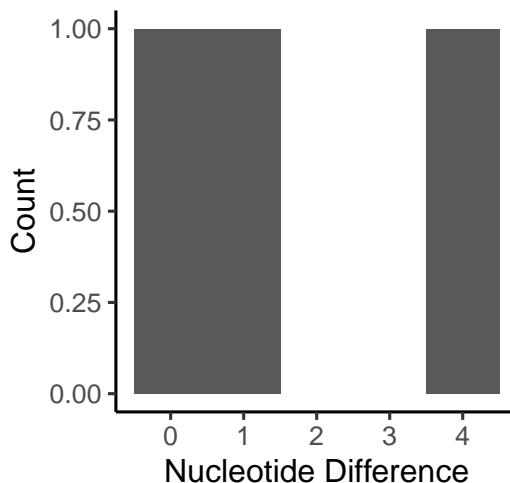
IGLV3-22*01

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



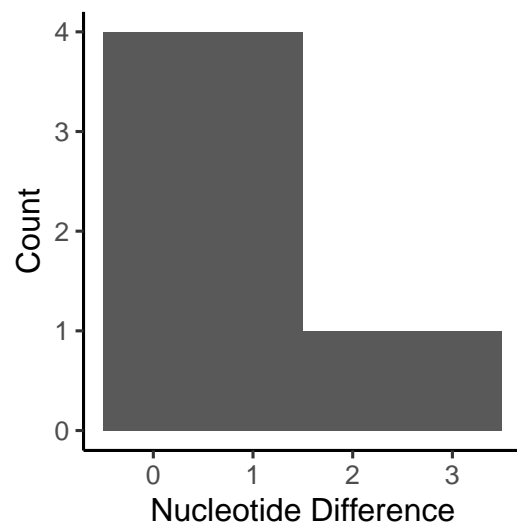
IGLV4-3*01

3 sequences assigned
1 (33.3%) exact matches, in which:
1 unique CDR3
1 unique J



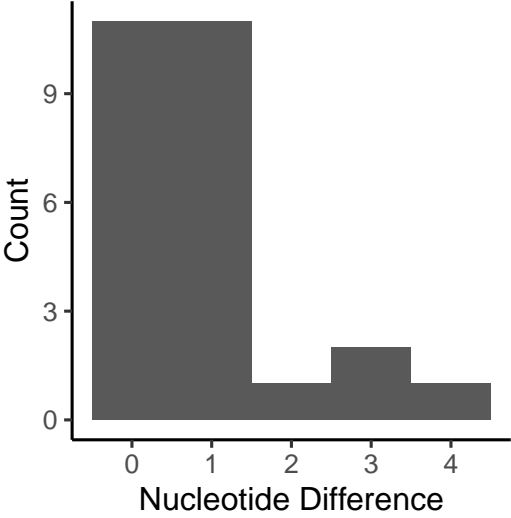
IGLV5-37*01

10 sequences assigned
4 (40%) exact matches, in which:
4 unique CDR3
2 unique J



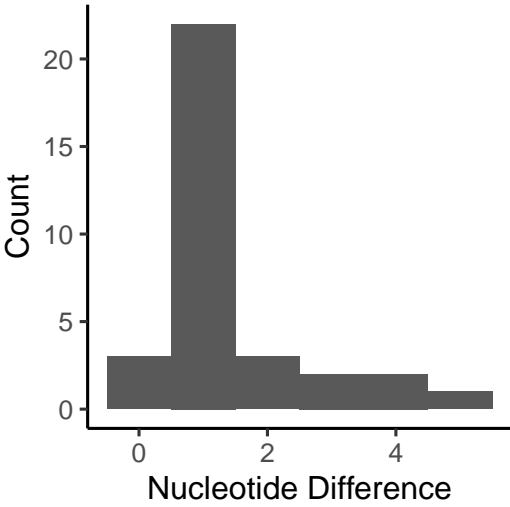
IGLV5-45*02

26 sequences assigned
11 (42.3%) exact matches, in which:
6 unique CDR3
1 unique J



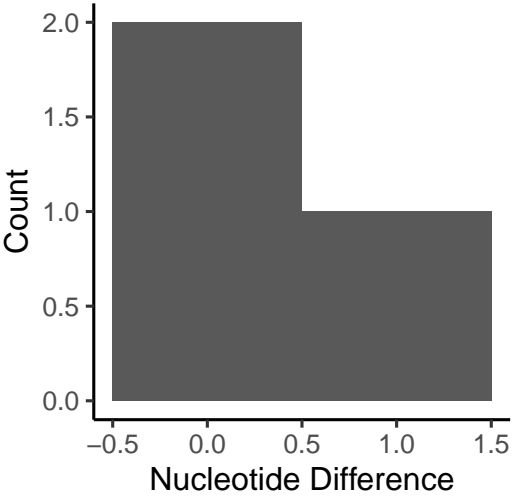
IGLV6-57*03

33 sequences assigned
3 (9.1%) exact matches, in which:
3 unique CDR3
1 unique J



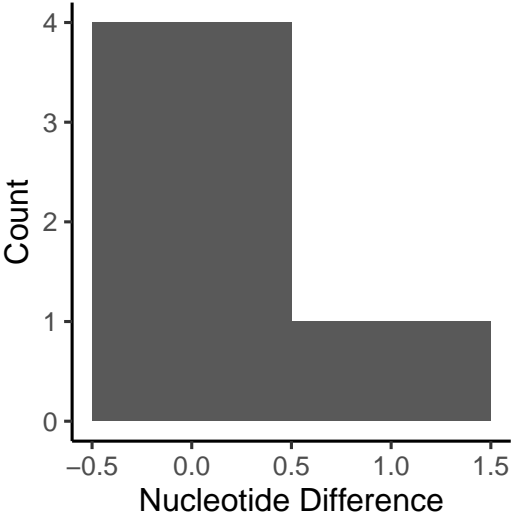
IGLV8-61*02

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



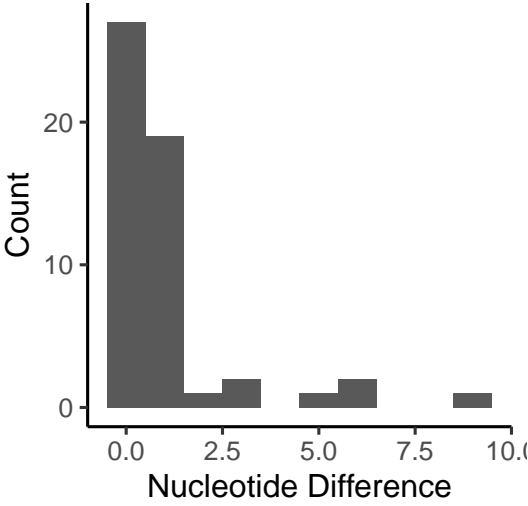
IGLV5-52*01

5 sequences assigned
4 (80%) exact matches, in which:
4 unique CDR3
2 unique J



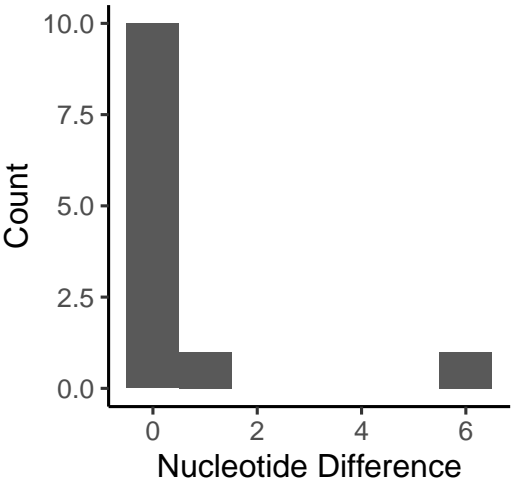
IGLV7-43*01

53 sequences assigned
27 (50.9%) exact matches, in which:
25 unique CDR3
2 unique J



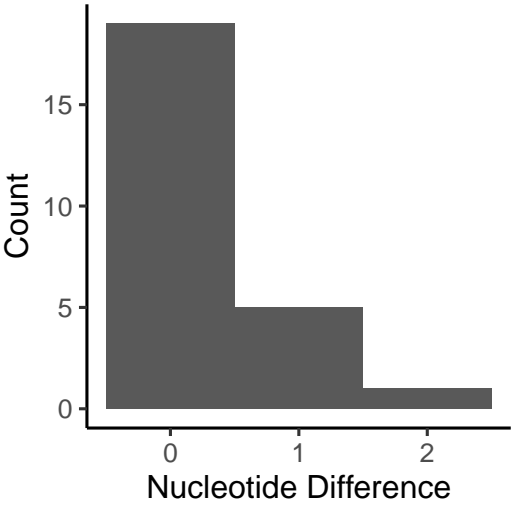
IGLV9-49*01

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



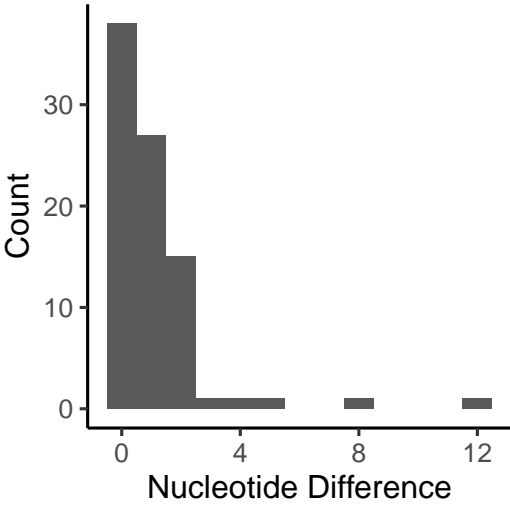
IGLV6-57*02

25 sequences assigned
19 (76%) exact matches, in which:
18 unique CDR3
1 unique J

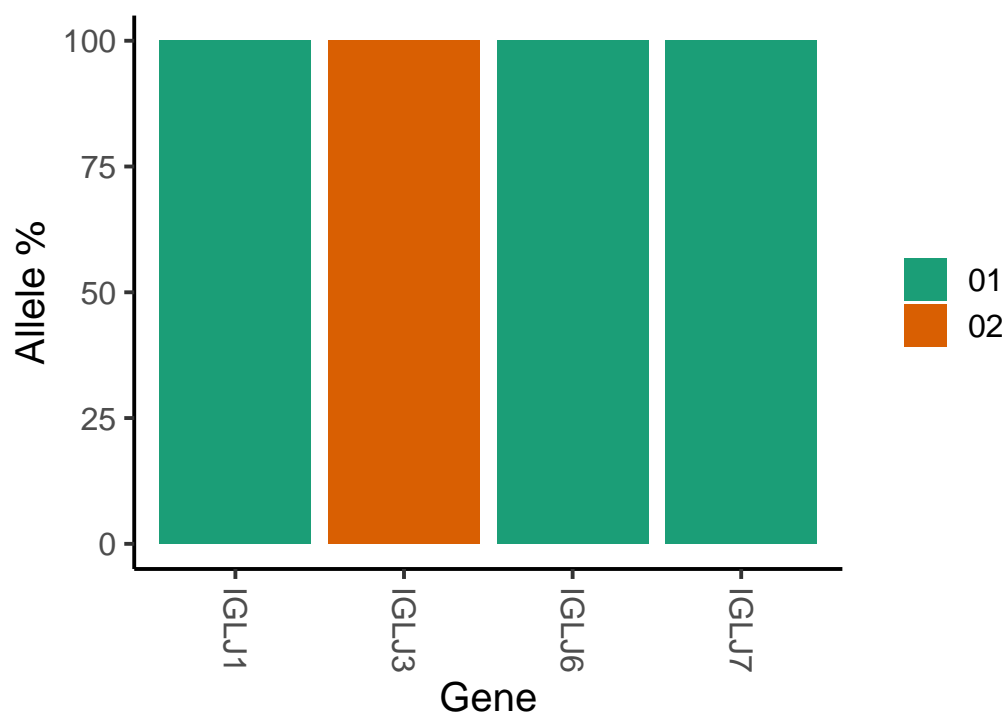


IGLV7-46*01

85 sequences assigned
38 (44.7%) exact matches, in which:
33 unique CDR3
2 unique J



Allele Usage



Warning – no inferred sequences found.