

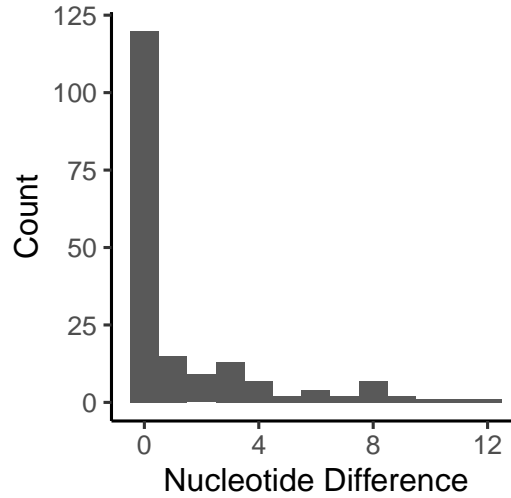
IGLV10-54*02

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



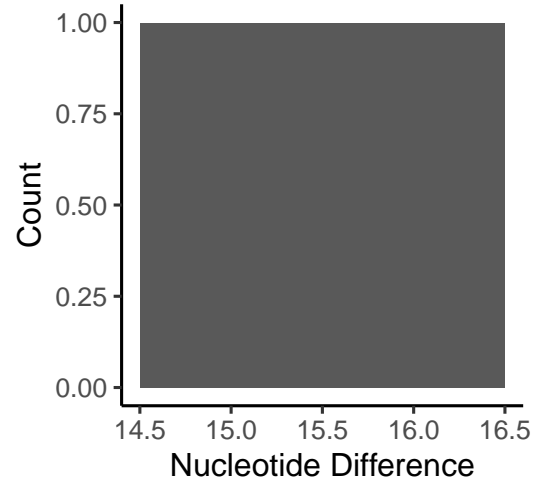
IGLV1-44*01

218 sequences assigned
120 (55%) exact matches, in which:
63 unique CDR3
3 unique J



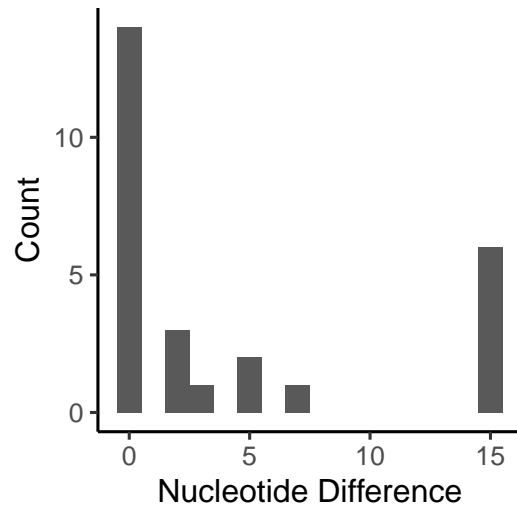
IGLV1-50*01

3 sequences assigned
No exact matches.



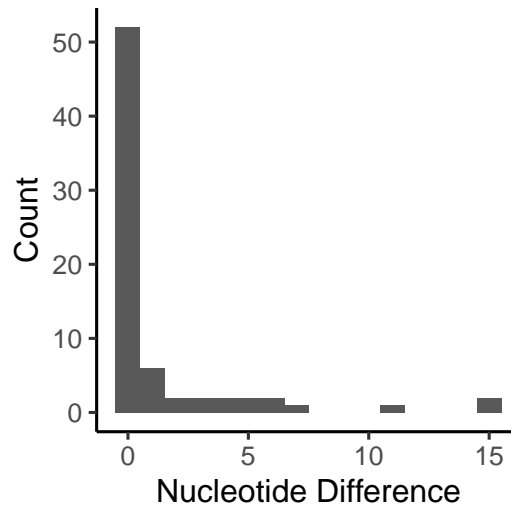
IGLV1-36*01

31 sequences assigned
14 (45.2%) exact matches, in which:
7 unique CDR3
2 unique J



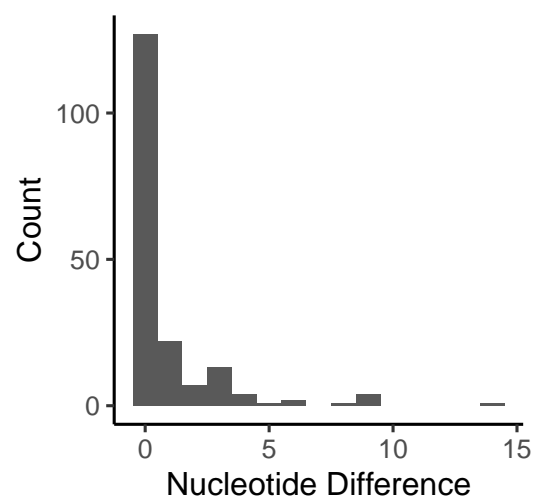
IGLV1-47*01

72 sequences assigned
52 (72.2%) exact matches, in which:
33 unique CDR3
2 unique J



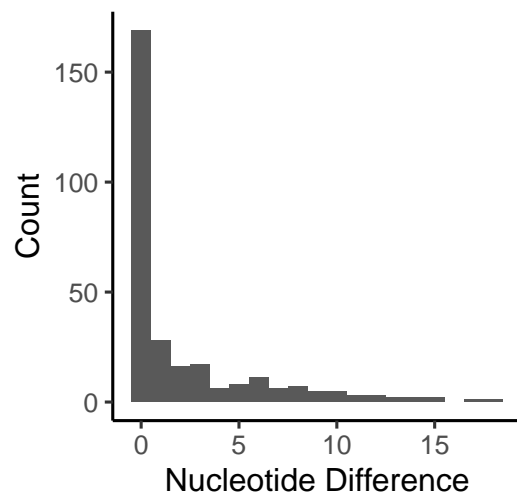
IGLV1-51*01

182 sequences assigned
127 (69.8%) exact matches, in which:
69 unique CDR3
3 unique J



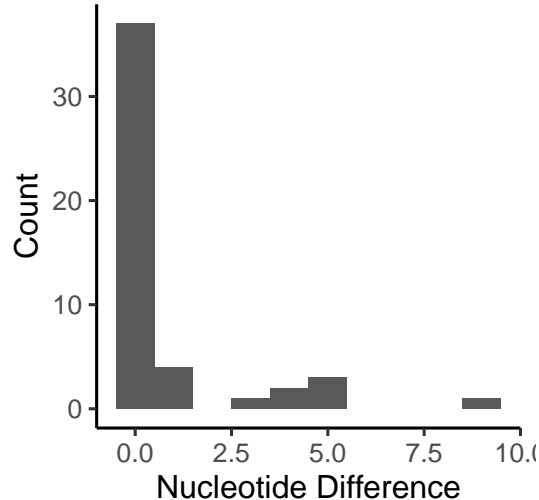
IGLV1-40*01

294 sequences assigned
169 (57.5%) exact matches, in which:
96 unique CDR3
3 unique J



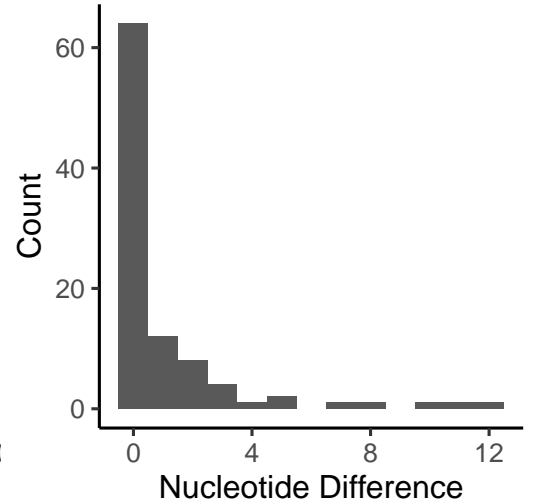
IGLV1-47*02

48 sequences assigned
37 (77.1%) exact matches, in which:
26 unique CDR3
2 unique J



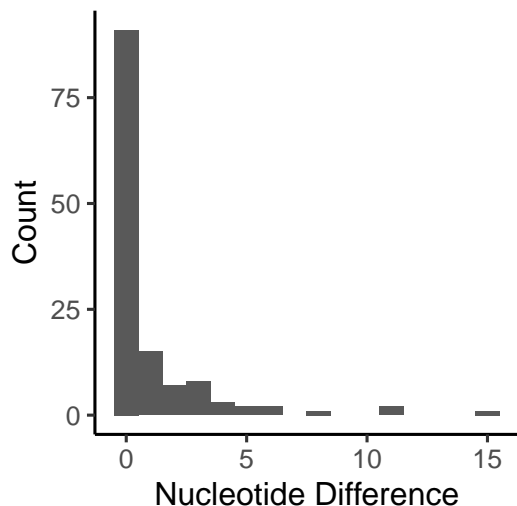
IGLV2-8*01

96 sequences assigned
64 (66.7%) exact matches, in which:
44 unique CDR3
2 unique J



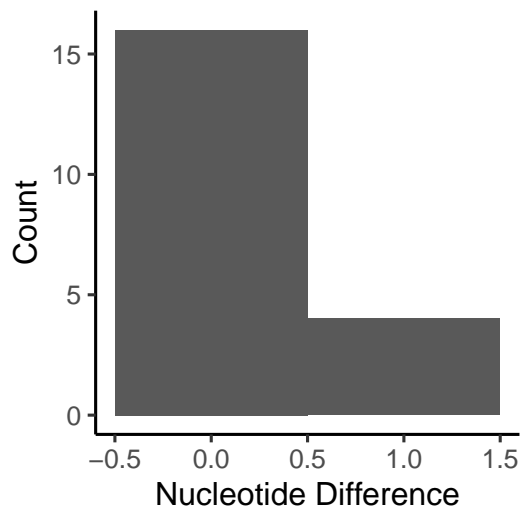
IGLV2-11*01

133 sequences assigned
91 (68.4%) exact matches, in which:
61 unique CDR3
2 unique J



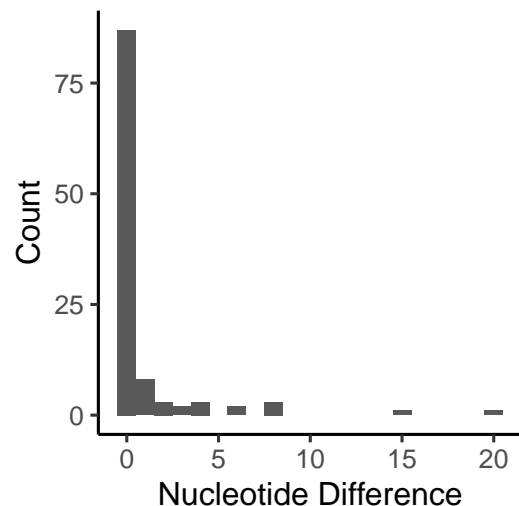
IGLV2-18*02

20 sequences assigned
16 (80%) exact matches, in which:
8 unique CDR3
2 unique J



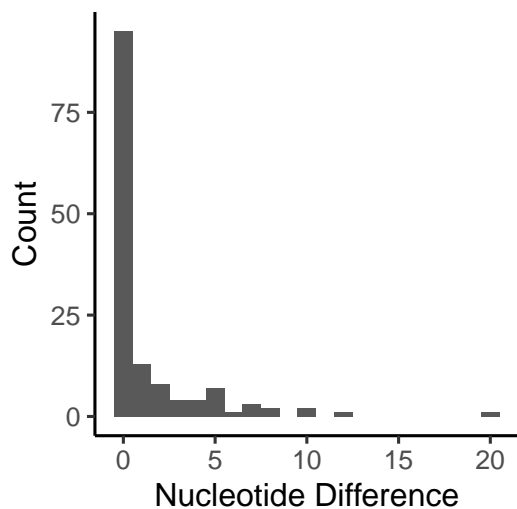
IGLV3-1*01

111 sequences assigned
87 (78.4%) exact matches, in which:
59 unique CDR3
2 unique J



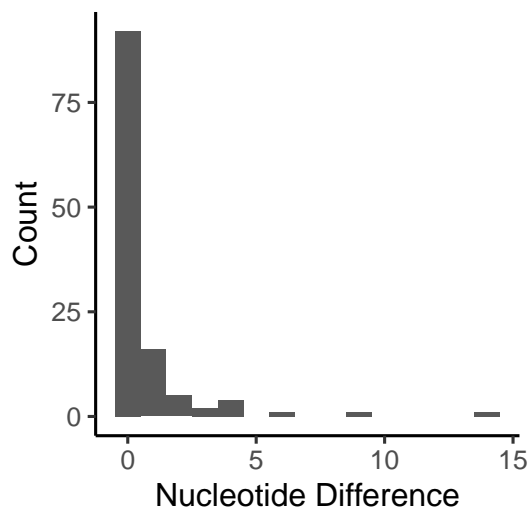
IGLV2-14*03EXT3

141 sequences assigned
95 (67.4%) exact matches, in which:
68 unique CDR3
2 unique J



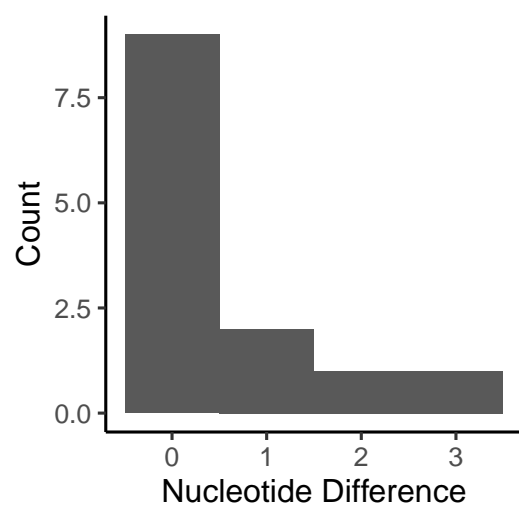
IGLV2-23*02EXT5

122 sequences assigned
92 (75.4%) exact matches, in which:
71 unique CDR3
2 unique J



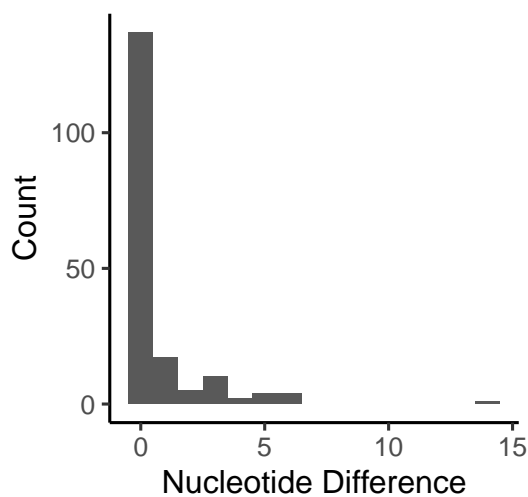
IGLV3-9*01EXT5

13 sequences assigned
9 (69.2%) exact matches, in which:
6 unique CDR3
2 unique J



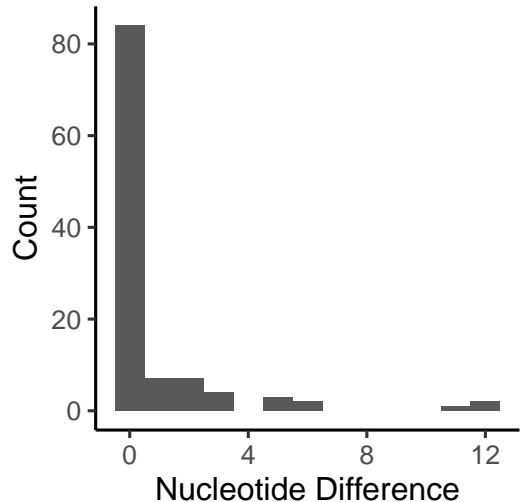
IGLV2-14*04EXT3

180 sequences assigned
137 (76.1%) exact matches, in which:
108 unique CDR3
3 unique J



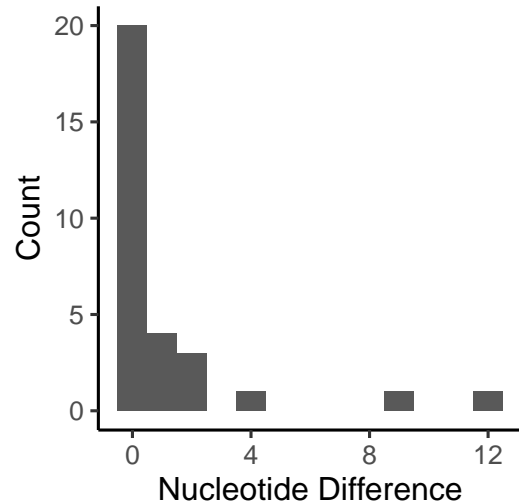
IGLV2-23*03EXT5

111 sequences assigned
84 (75.7%) exact matches, in which:
61 unique CDR3
4 unique J



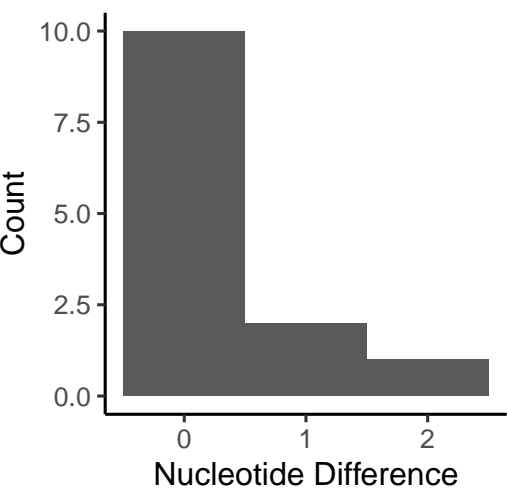
IGLV3-10*01

30 sequences assigned
20 (66.7%) exact matches, in which:
16 unique CDR3
2 unique J



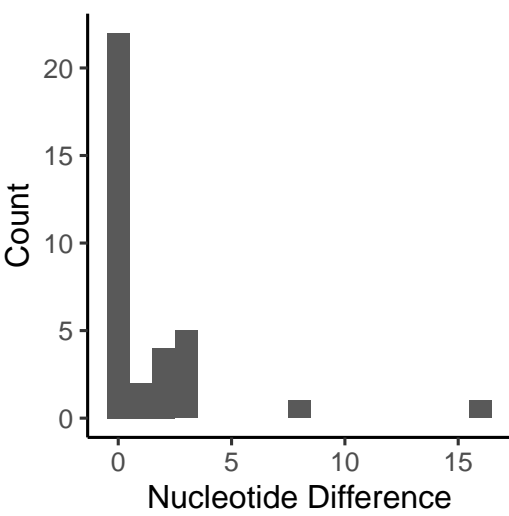
IGLV3-16*01

13 sequences assigned
10 (76.9%) exact matches, in which:
7 unique CDR3
2 unique J



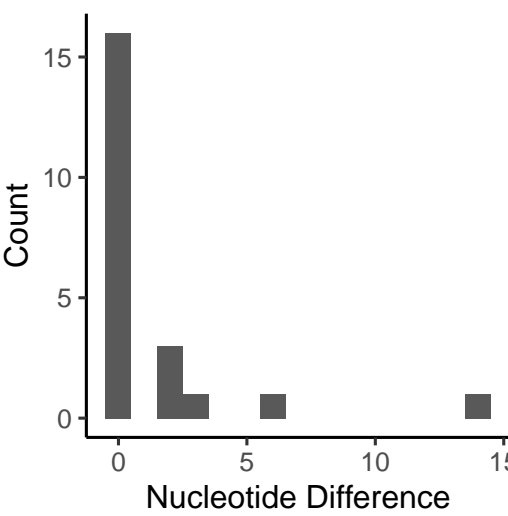
IGLV3-21*04

35 sequences assigned
22 (62.9%) exact matches, in which:
15 unique CDR3
2 unique J



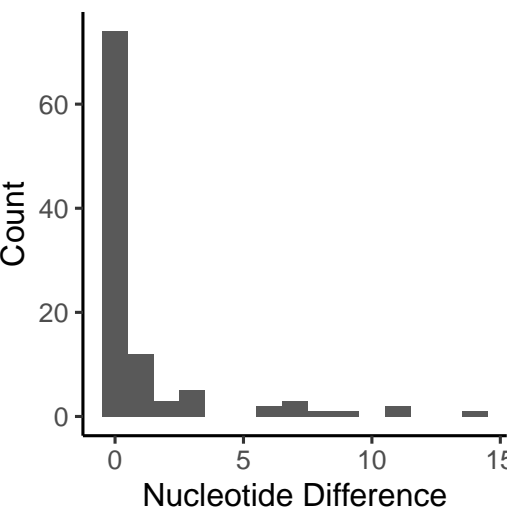
IGLV3-27*01

22 sequences assigned
16 (72.7%) exact matches, in which:
16 unique CDR3
2 unique J



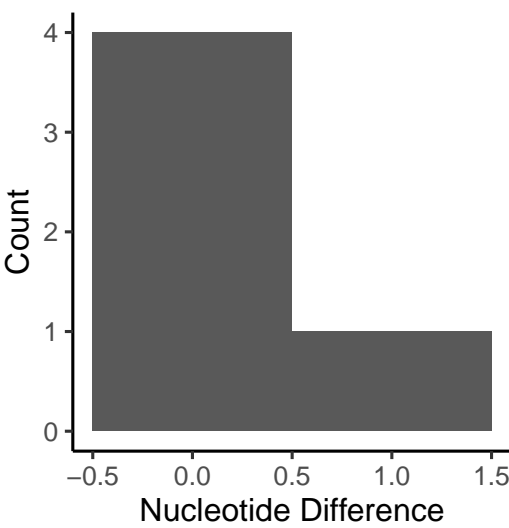
IGLV3-19*01

104 sequences assigned
74 (71.2%) exact matches, in which:
48 unique CDR3
2 unique J



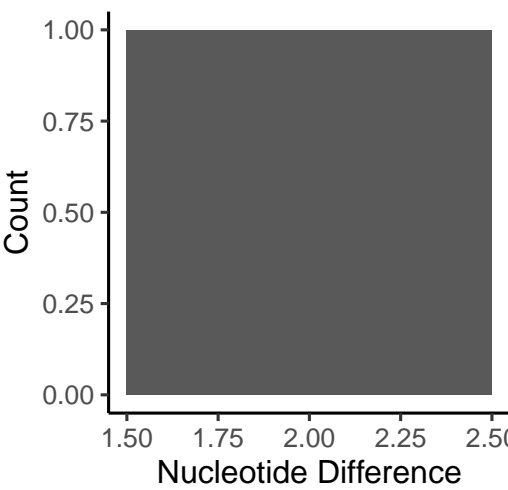
IGLV3-22*01

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



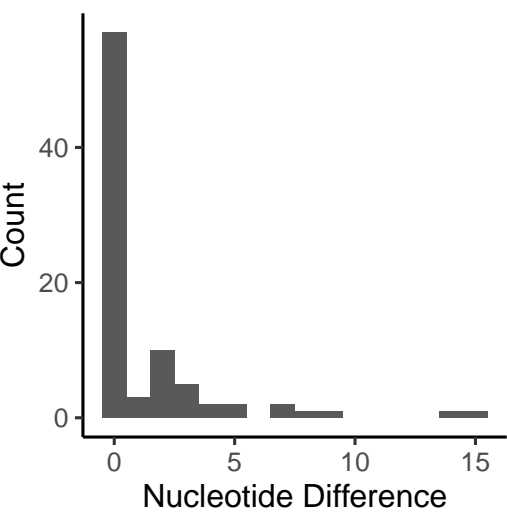
IGLV4-3*01

1 sequences assigned
No exact matches.



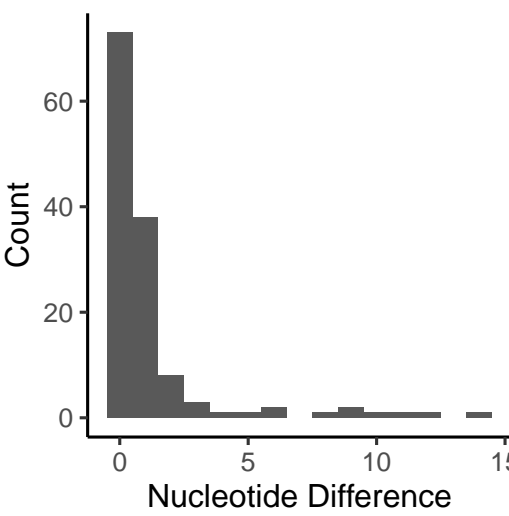
IGLV3-21*03

86 sequences assigned
57 (66.3%) exact matches, in which:
43 unique CDR3
2 unique J



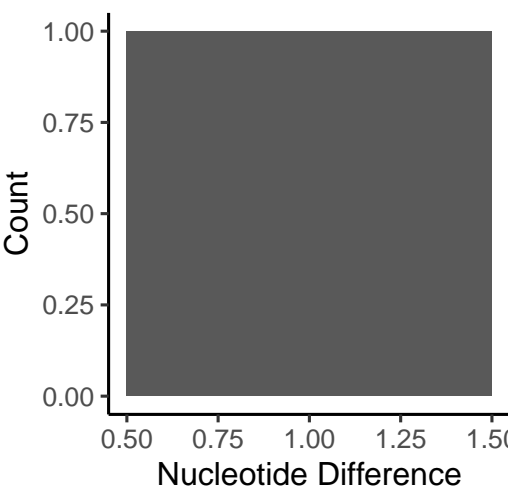
IGLV3-25*03

133 sequences assigned
73 (54.9%) exact matches, in which:
40 unique CDR3
2 unique J



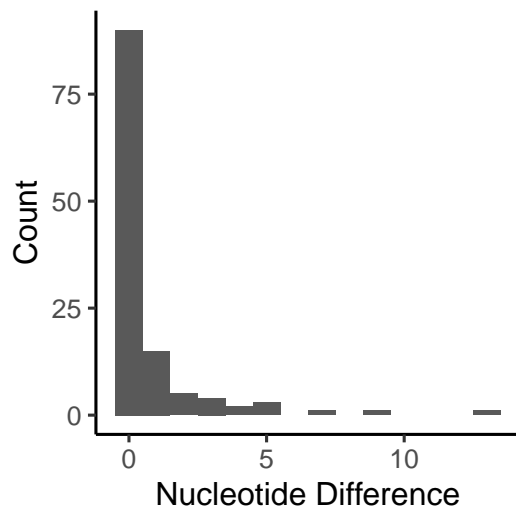
IGLV4-60*03EXT5

1 sequences assigned
No exact matches.



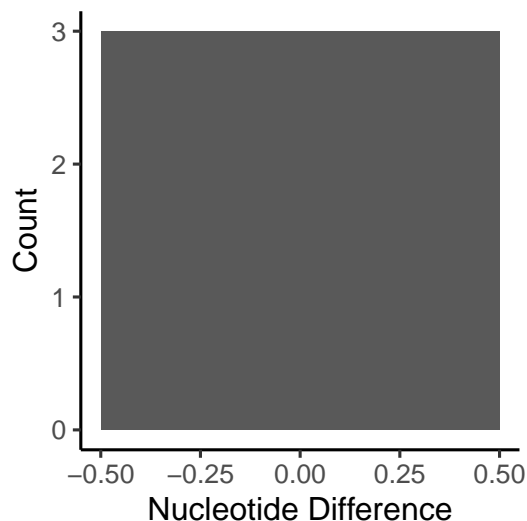
IGLV4-69*01

124 sequences assigned
90 (72.6%) exact matches, in which:
55 unique CDR3
2 unique J



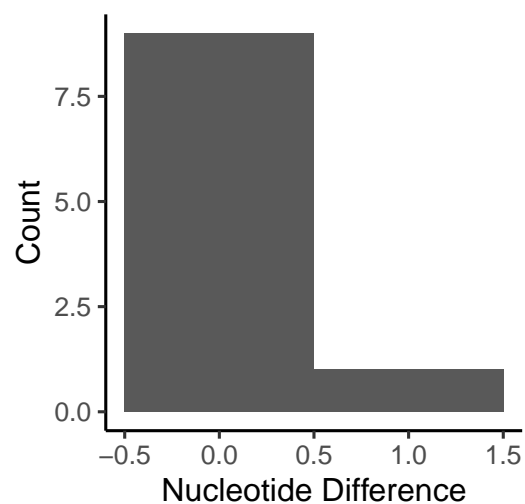
IGLV5-45*02

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



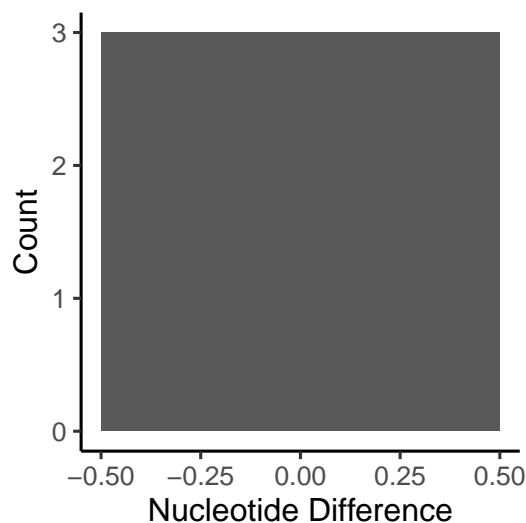
IGLV5-52*01

10 sequences assigned
9 (90%) exact matches, in which:
7 unique CDR3
1 unique J



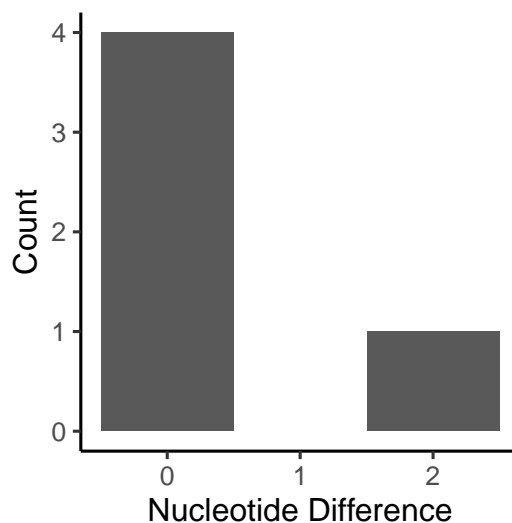
IGLV5-37*01

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



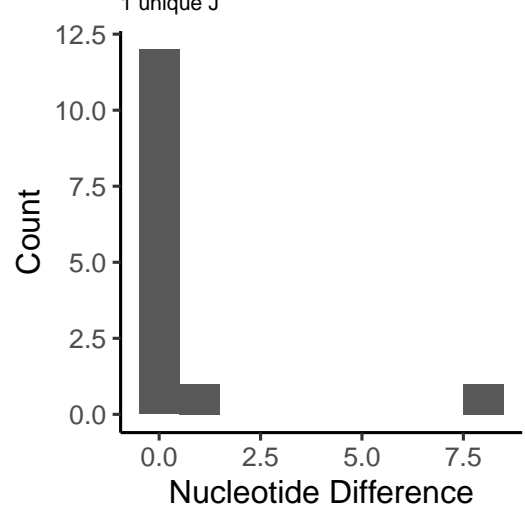
IGLV5-45*03

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



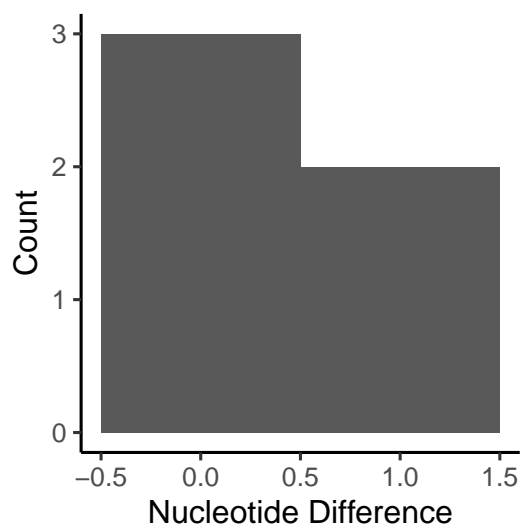
IGLV6-57*02

14 sequences assigned
12 (85.7%) exact matches, in which:
11 unique CDR3
1 unique J



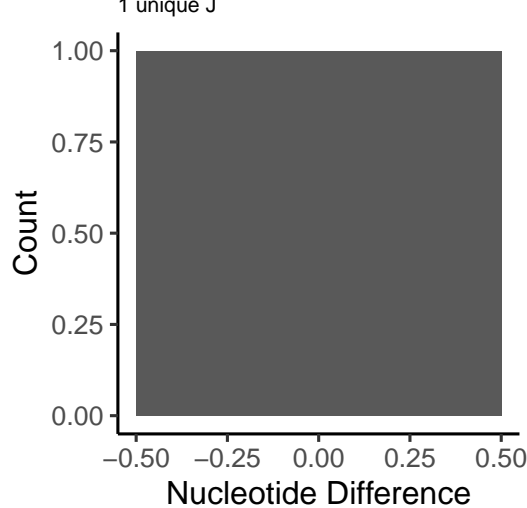
IGLV5-39*01

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



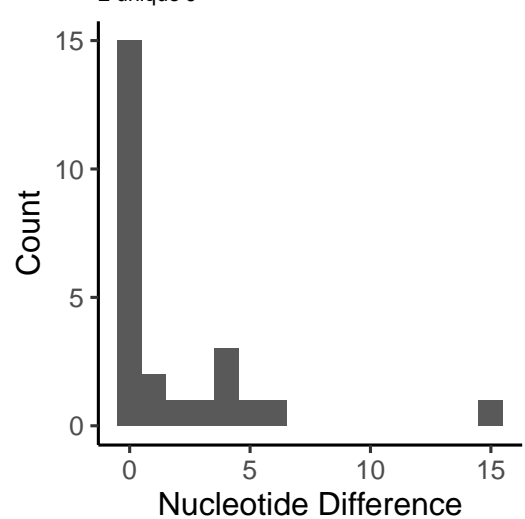
IGLV5-48*01

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



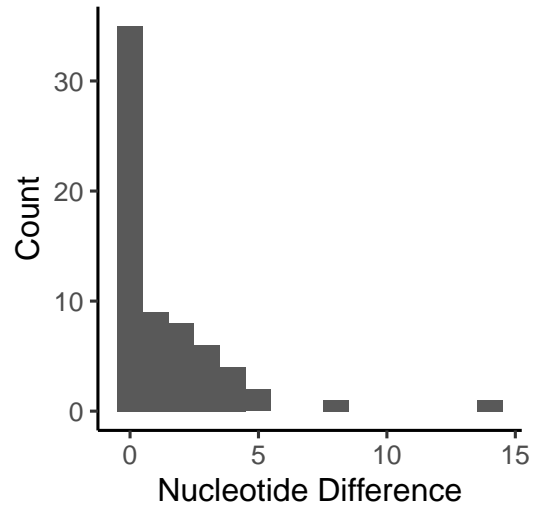
IGLV6-57*03

25 sequences assigned
15 (60%) exact matches, in which:
14 unique CDR3
2 unique J



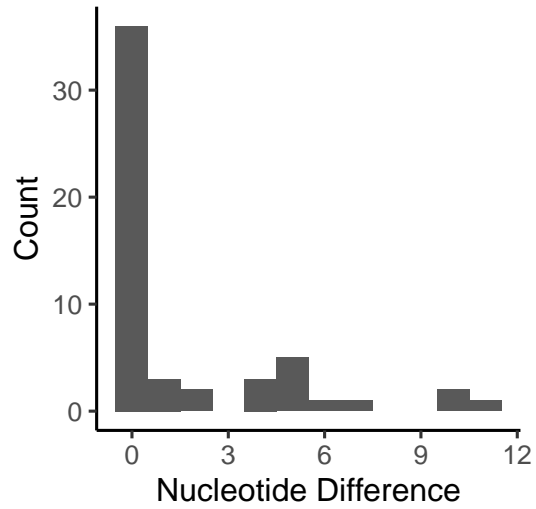
IGLV7-43*01

66 sequences assigned
35 (53%) exact matches, in which:
24 unique CDR3
2 unique J



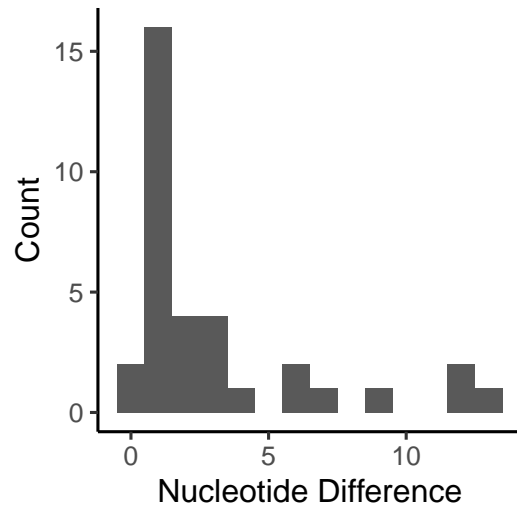
IGLV9-49*01

54 sequences assigned
36 (66.7%) exact matches, in which:
23 unique CDR3
2 unique J



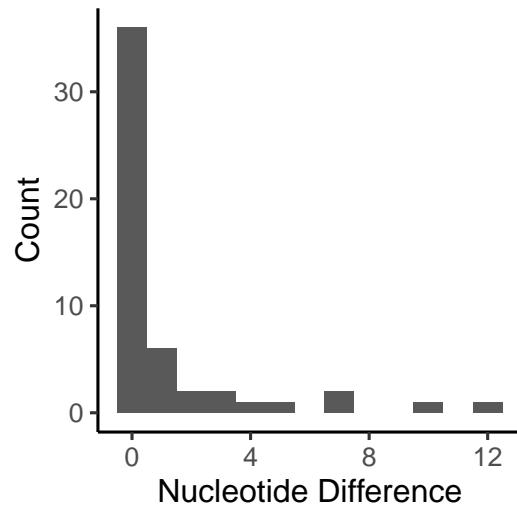
IGLV7-46*01

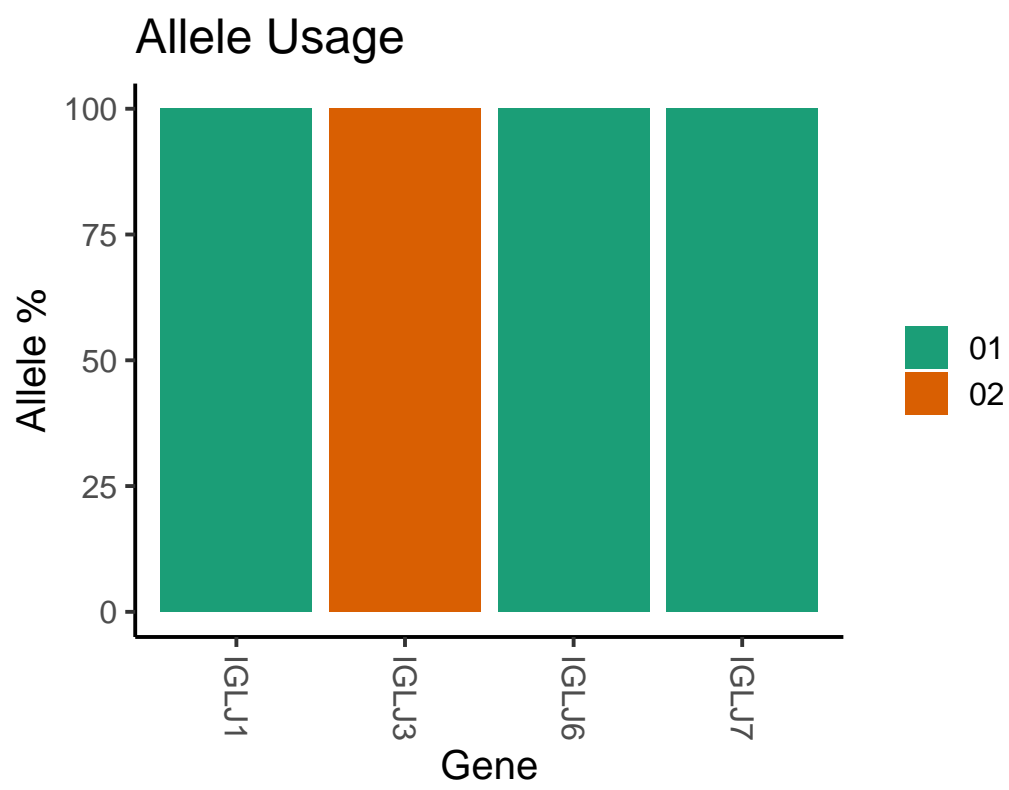
34 sequences assigned
2 (5.9%) exact matches, in which:
2 unique CDR3
2 unique J



IGLV8-61*01

52 sequences assigned
36 (69.2%) exact matches, in which:
22 unique CDR3
2 unique J





Warning – no inferred sequences found.