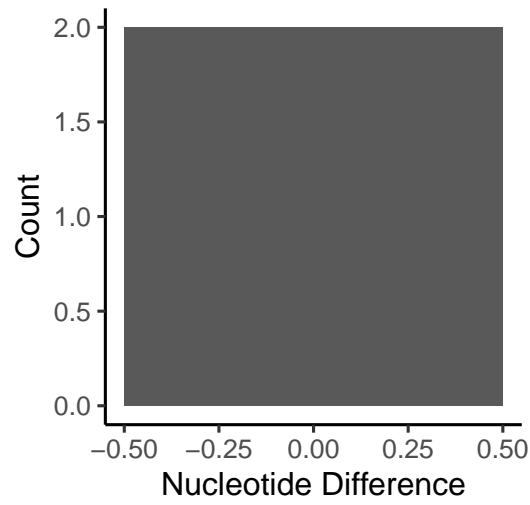


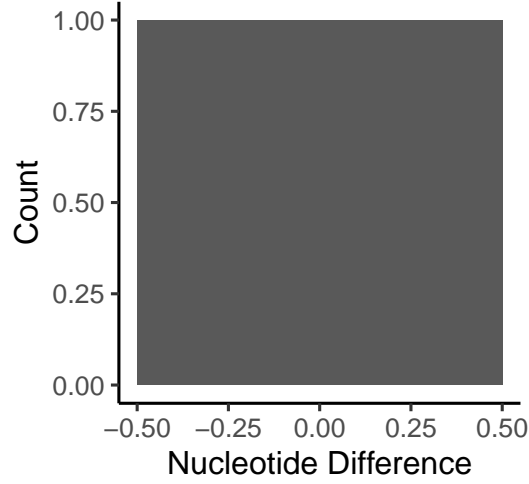
IGLV10-54*02

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



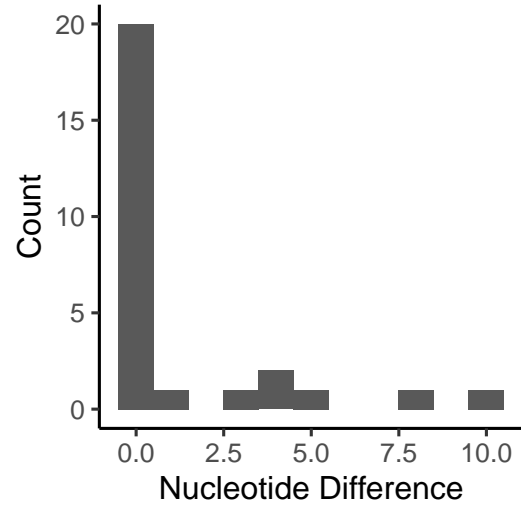
IGLV1-41*01

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



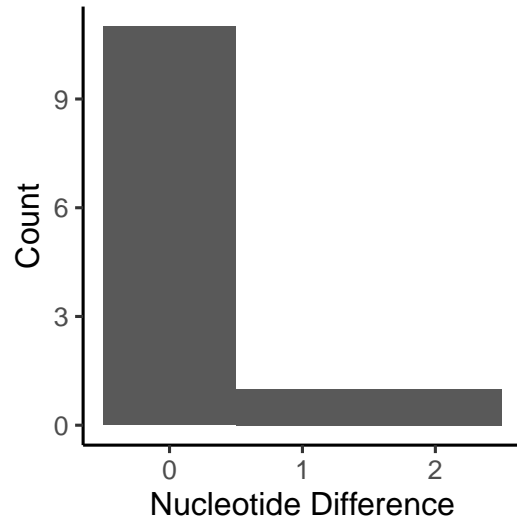
IGLV1-51*01

27 sequences assigned
20 (74.1%) exact matches, in which:
16 unique CDR3
2 unique J



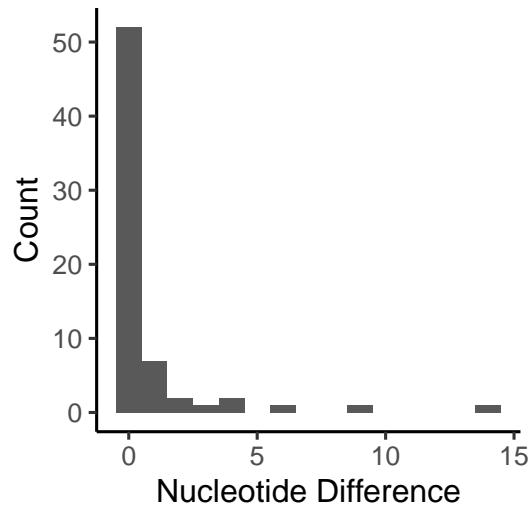
IGLV1-36*01

13 sequences assigned
11 (84.6%) exact matches, in which:
8 unique CDR3
3 unique J



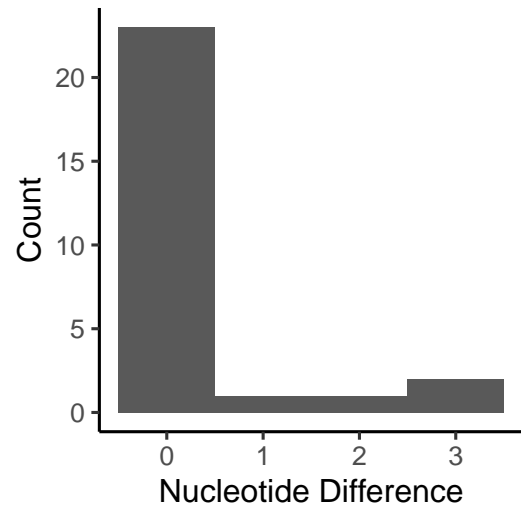
IGLV1-44*01

67 sequences assigned
52 (77.6%) exact matches, in which:
37 unique CDR3
3 unique J



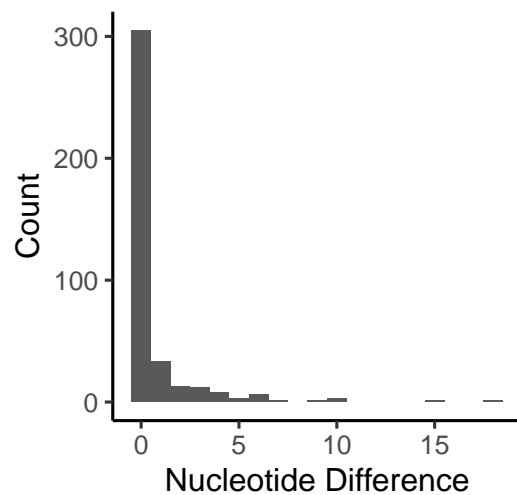
IGLV1-51*02

27 sequences assigned
23 (85.2%) exact matches, in which:
20 unique CDR3
2 unique J



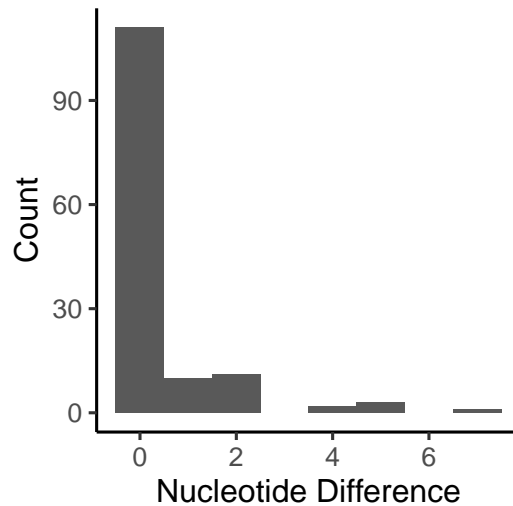
IGLV1-40*01

387 sequences assigned
305 (78.8%) exact matches, in which:
223 unique CDR3
3 unique J



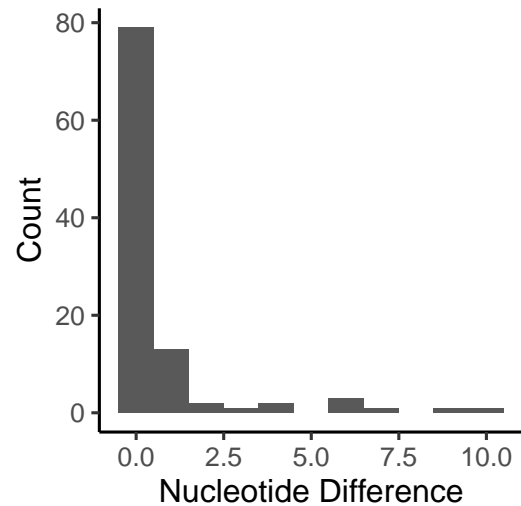
IGLV1-47*01

138 sequences assigned
111 (80.4%) exact matches, in which:
78 unique CDR3
3 unique J



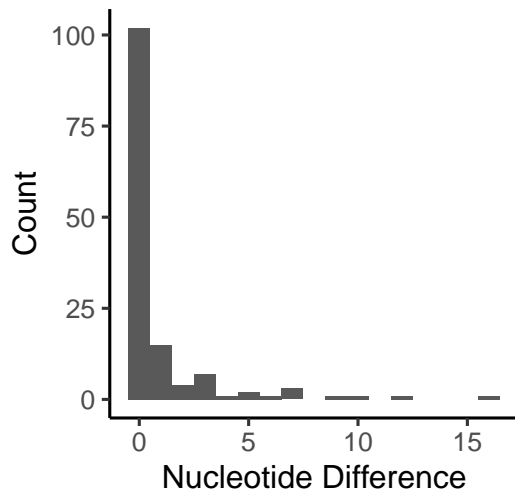
IGLV2-8*01

103 sequences assigned
79 (76.7%) exact matches, in which:
66 unique CDR3
2 unique J



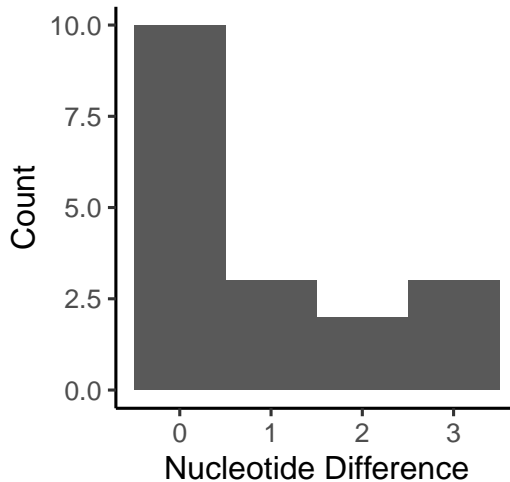
IGLV2-11*01

139 sequences assigned
102 (73.4%) exact matches, in which:
80 unique CDR3
2 unique J



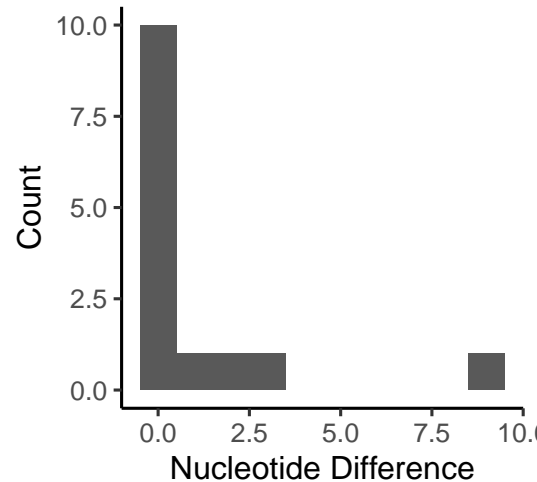
IGLV2-18*02

18 sequences assigned
10 (55.6%) exact matches, in which:
5 unique CDR3
2 unique J



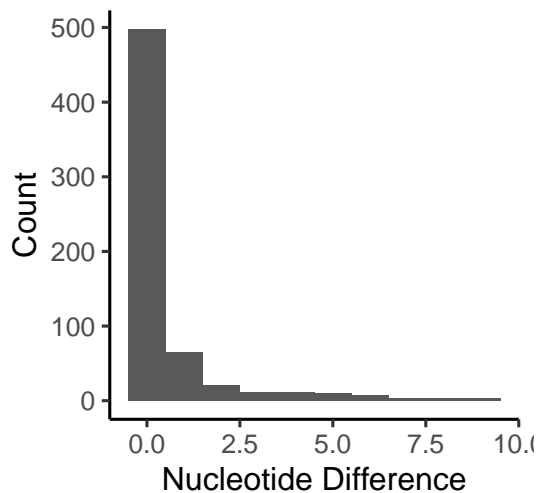
IGLV3-9*01EXT5

14 sequences assigned
10 (71.4%) exact matches, in which:
10 unique CDR3
3 unique J



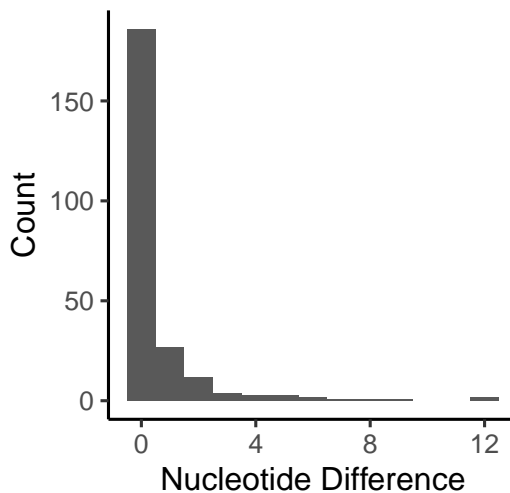
IGLV2-14*01

636 sequences assigned
498 (78.3%) exact matches, in which:
376 unique CDR3
5 unique J



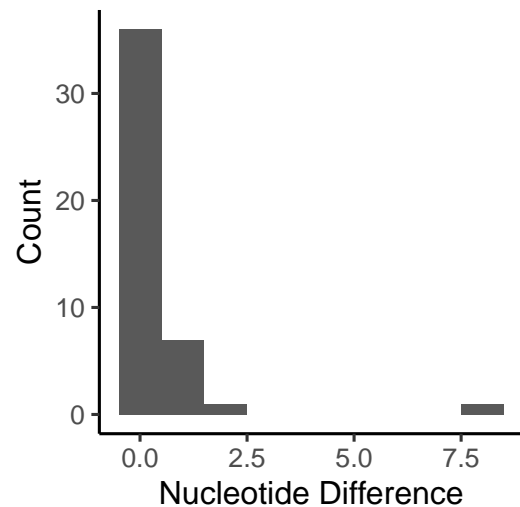
IGLV2-23*03EXT5

243 sequences assigned
186 (76.5%) exact matches, in which:
148 unique CDR3
3 unique J



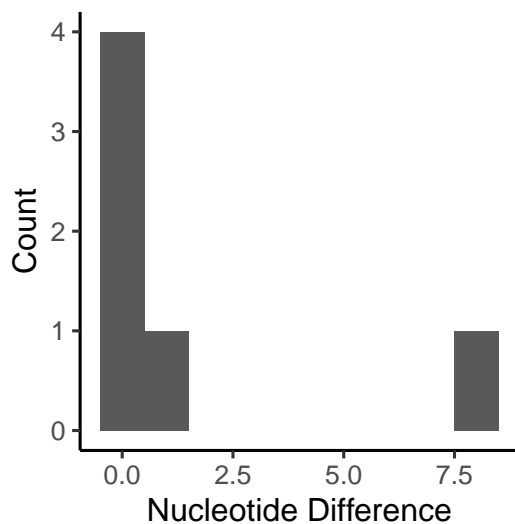
IGLV3-10*01

45 sequences assigned
36 (80%) exact matches, in which:
35 unique CDR3
2 unique J



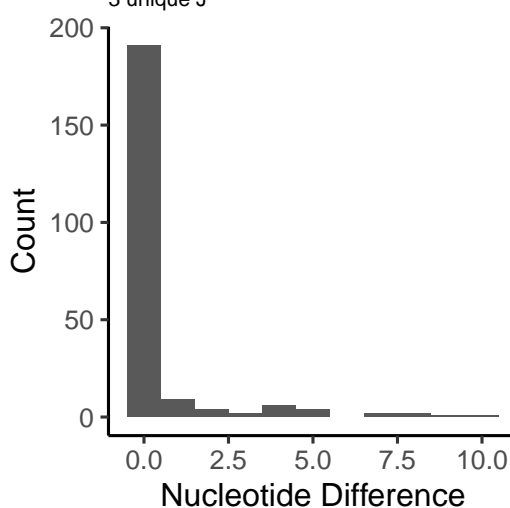
IGLV2-18*01

6 sequences assigned
4 (66.7%) exact matches, in which:
3 unique CDR3
2 unique J



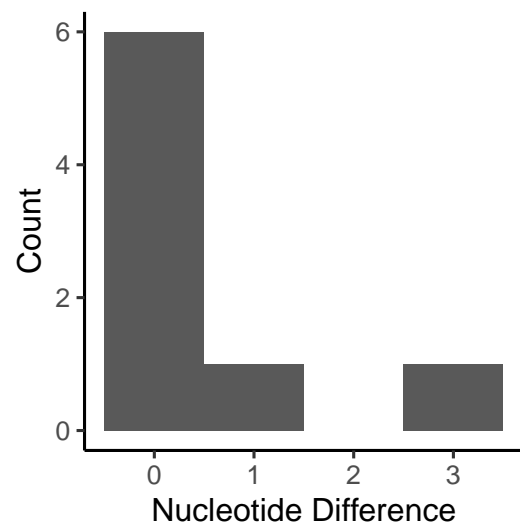
IGLV3-1*01

222 sequences assigned
191 (86%) exact matches, in which:
142 unique CDR3
3 unique J



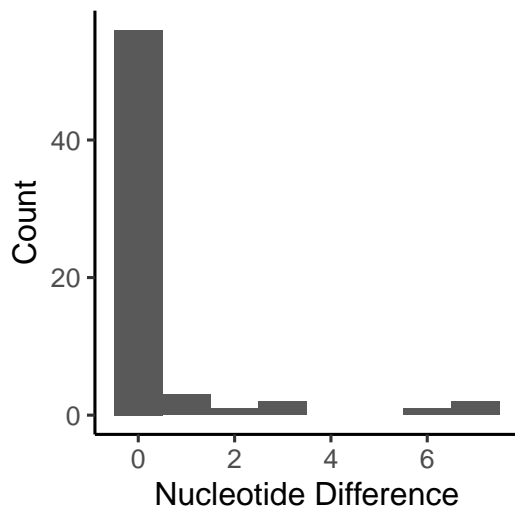
IGLV3-16*01

8 sequences assigned
6 (75%) exact matches, in which:
6 unique CDR3
2 unique J



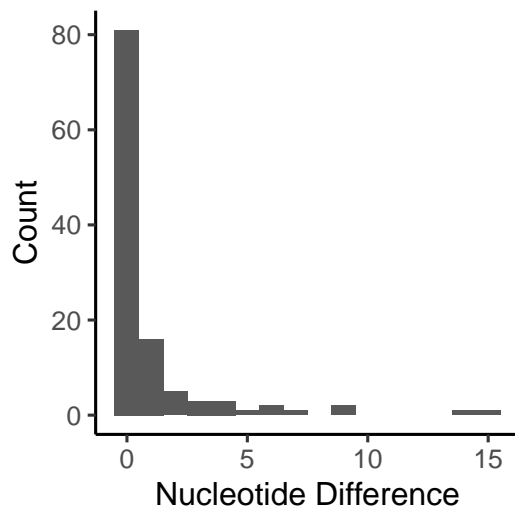
IGLV3-19*01

65 sequences assigned
56 (86.2%) exact matches, in which:
42 unique CDR3
2 unique J



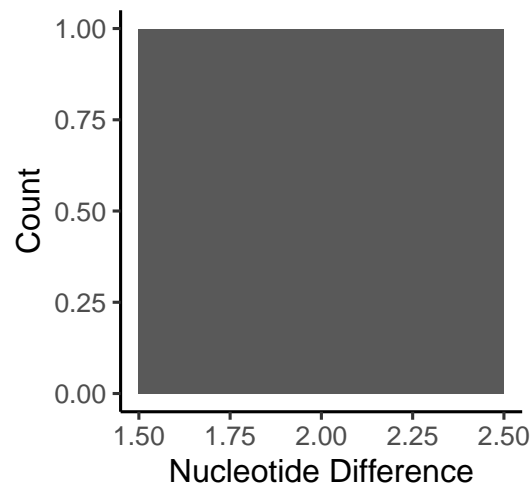
IGLV3-25*02

117 sequences assigned
81 (69.2%) exact matches, in which:
65 unique CDR3
2 unique J



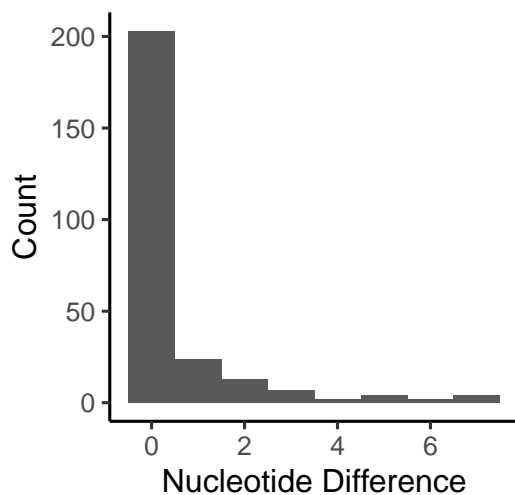
IGLV4-60*03EXT5

1 sequences assigned
No exact matches.



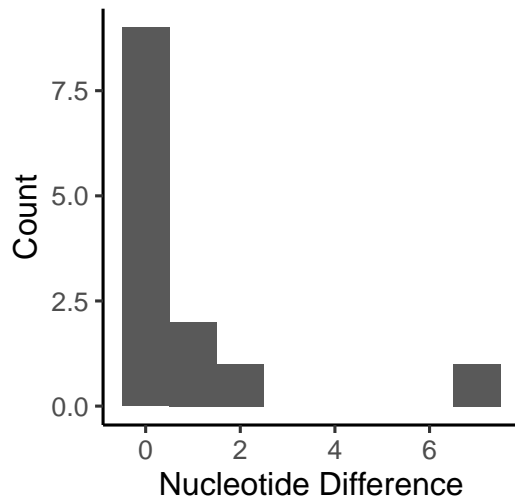
IGLV3-21*02

259 sequences assigned
203 (78.4%) exact matches, in which:
141 unique CDR3
4 unique J



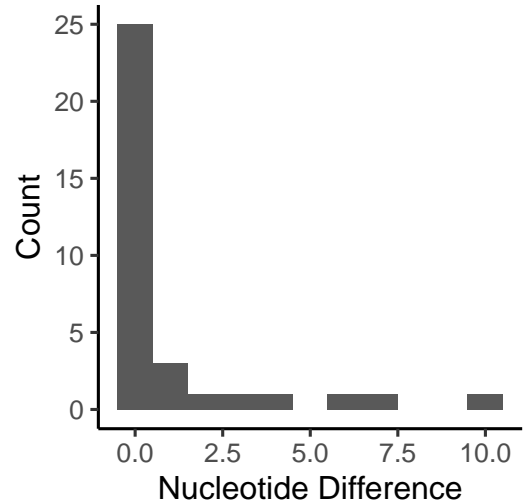
IGLV3-27*01

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



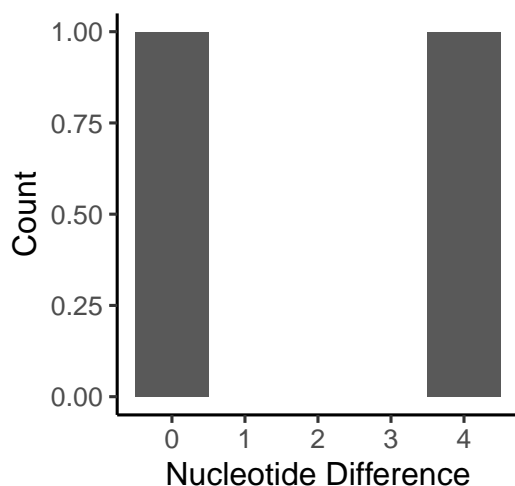
IGLV4-69*01

34 sequences assigned
25 (73.5%) exact matches, in which:
24 unique CDR3
4 unique J



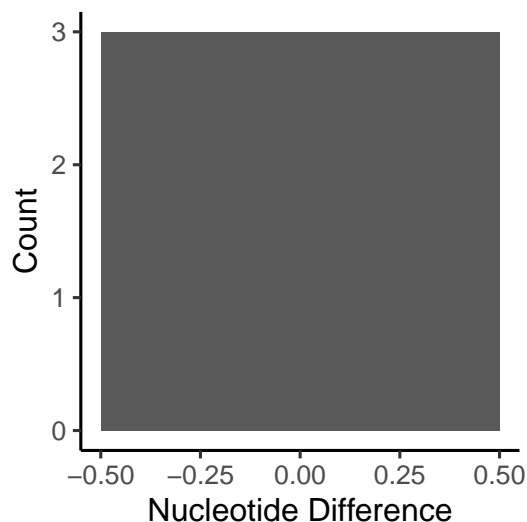
IGLV3-22*01

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



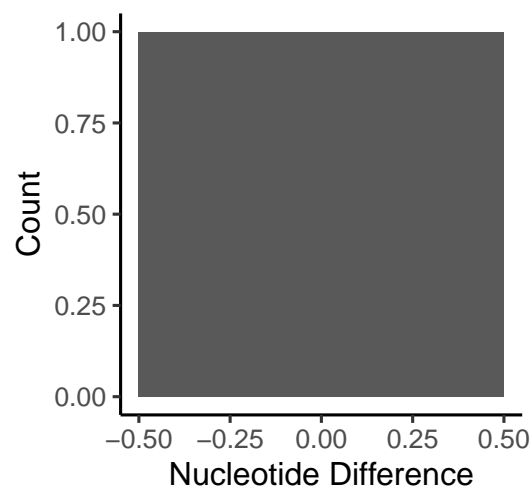
IGLV4-3*01

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



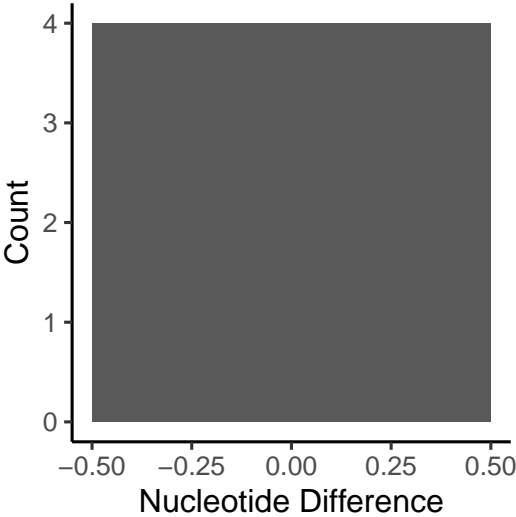
IGLV5-37*01

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



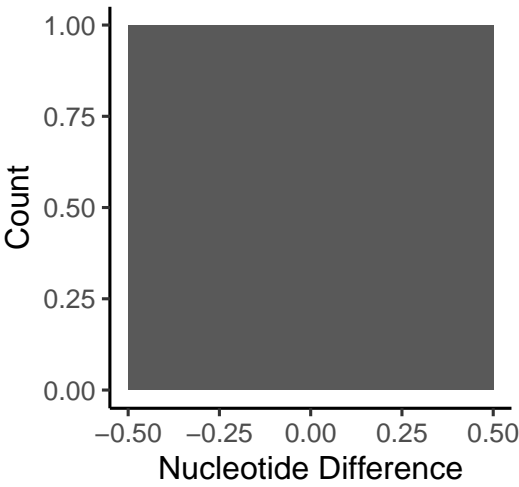
IGLV5–39*01

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



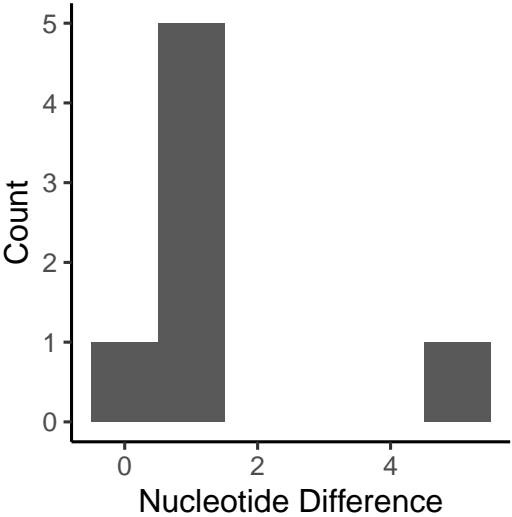
IGLV5–48*01

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



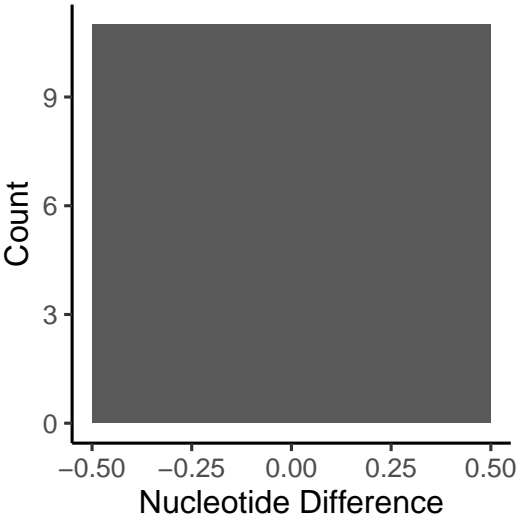
IGLV6–57*03

7 sequences assigned
1 (14.3%) exact matches, in which:
1 unique CDR3
1 unique J



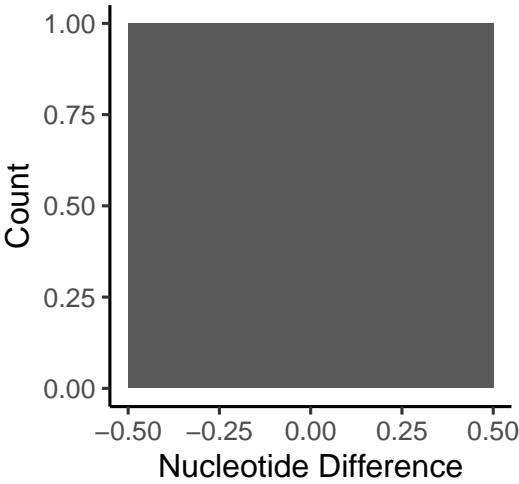
IGLV5–45*01

11 sequences assigned
11 (100%) exact matches, in which:
8 unique CDR3
3 unique J



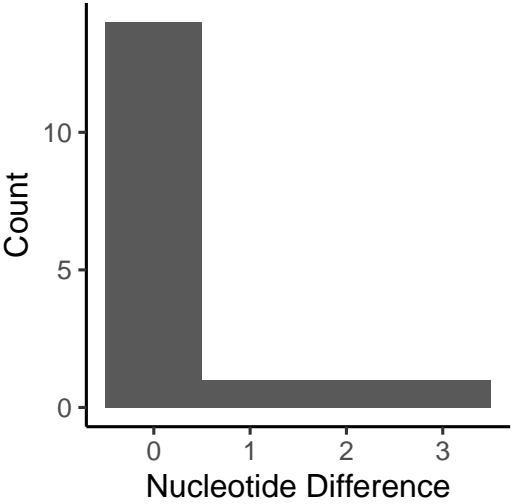
IGLV5–52*01

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



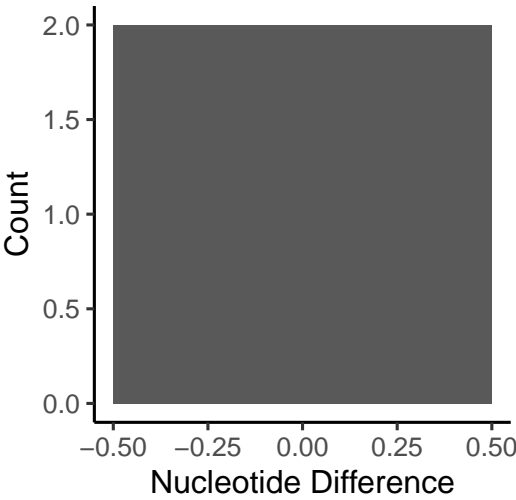
IGLV7–43*01

17 sequences assigned
14 (82.4%) exact matches, in which:
14 unique CDR3
2 unique J



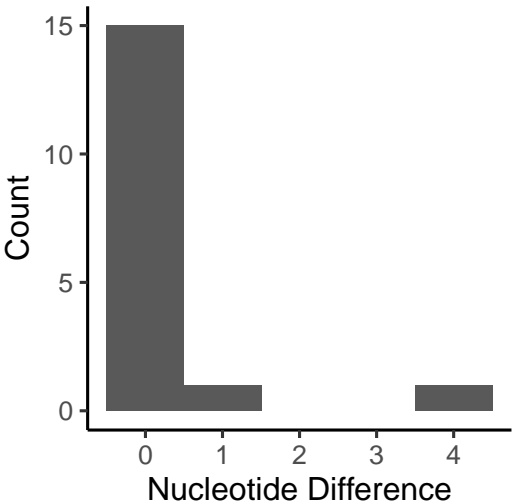
IGLV5–45*02

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



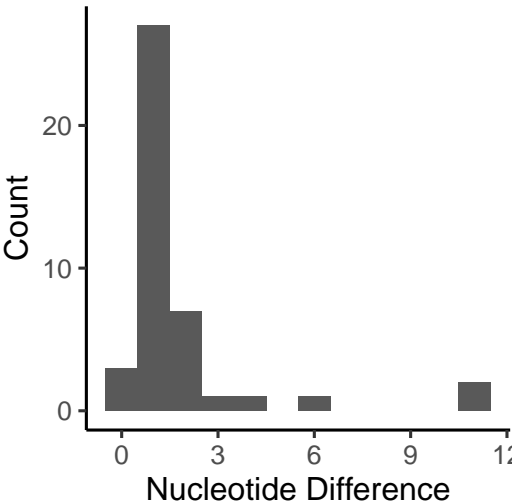
IGLV6–57*02

17 sequences assigned
15 (88.2%) exact matches, in which:
14 unique CDR3
3 unique J



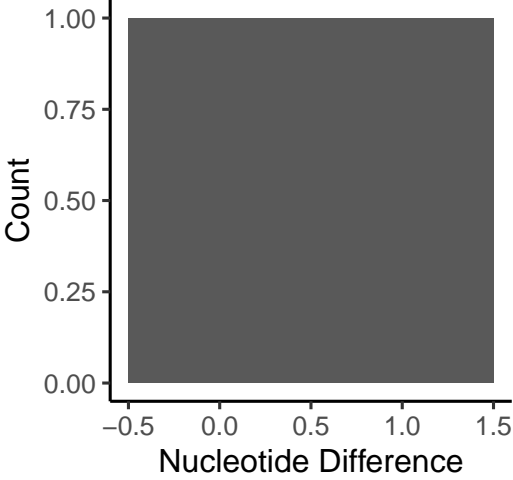
IGLV7–46*01

42 sequences assigned
3 (7.1%) exact matches, in which:
3 unique CDR3
1 unique J



IGLV9-49*01

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





Warning – no inferred sequences found.