

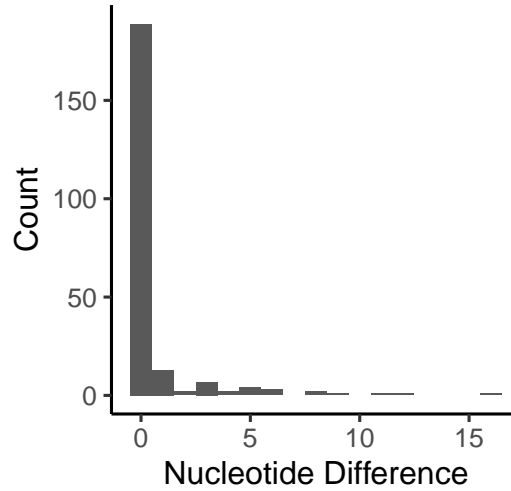
IGLV10-54*01

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



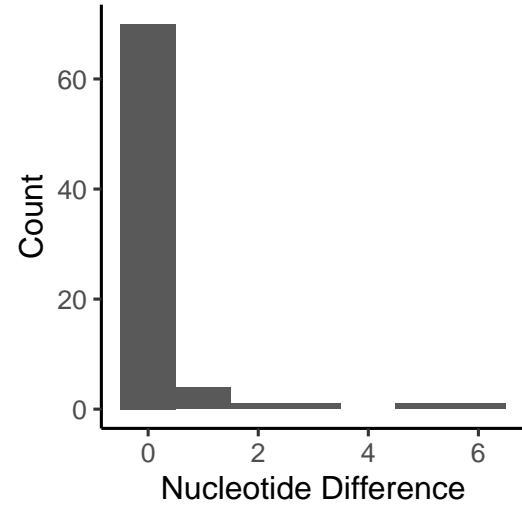
IGLV1-40*01

227 sequences assigned
189 (83.3%) exact matches, in which:
173 unique CDR3
4 unique J



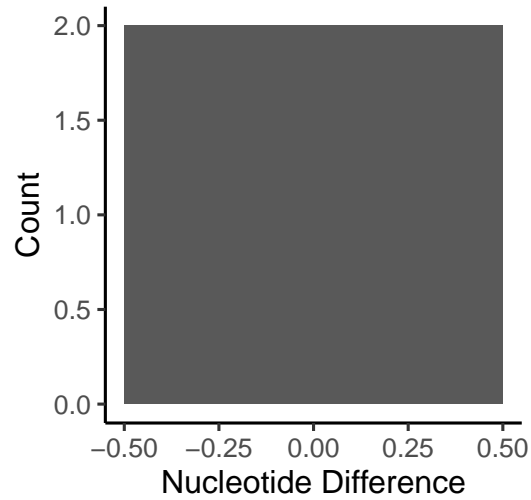
IGLV1-47*01

78 sequences assigned
70 (89.7%) exact matches, in which:
66 unique CDR3
4 unique J



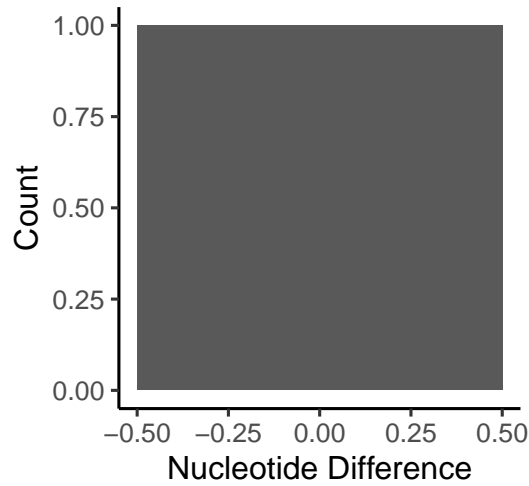
IGLV10-54*02

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



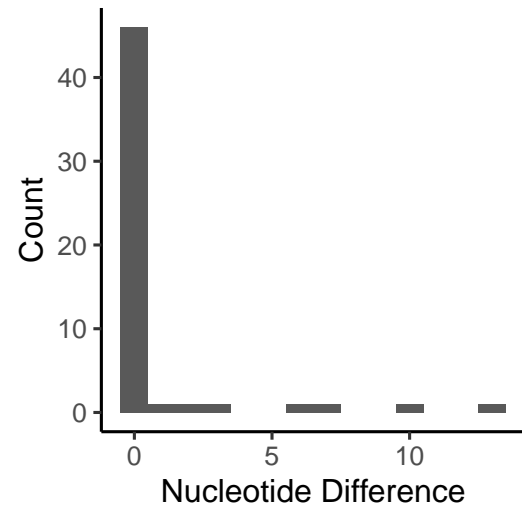
IGLV1-41*01

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



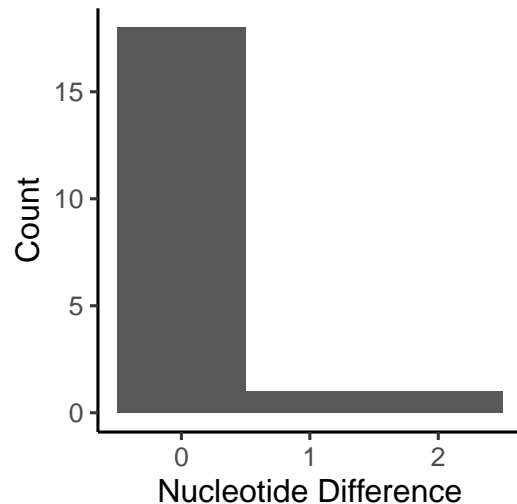
IGLV1-47*02

53 sequences assigned
46 (86.8%) exact matches, in which:
42 unique CDR3
3 unique J



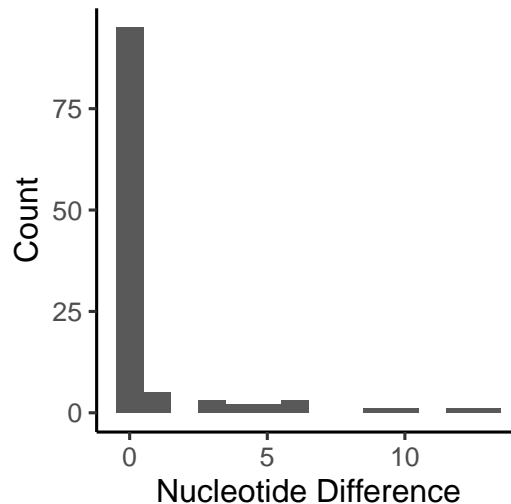
IGLV1-36*01

20 sequences assigned
18 (90%) exact matches, in which:
18 unique CDR3
3 unique J



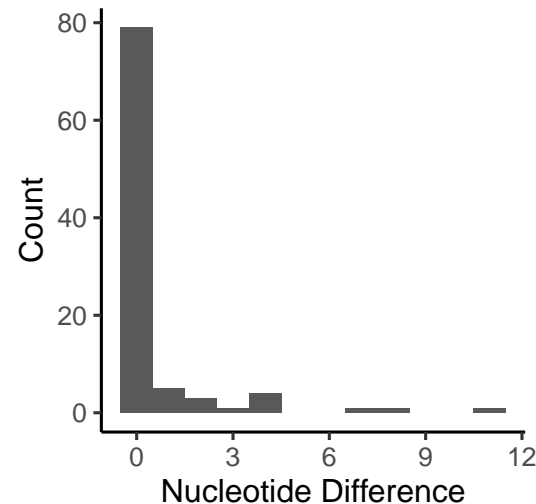
IGLV1-44*01

115 sequences assigned
95 (82.6%) exact matches, in which:
85 unique CDR3
4 unique J



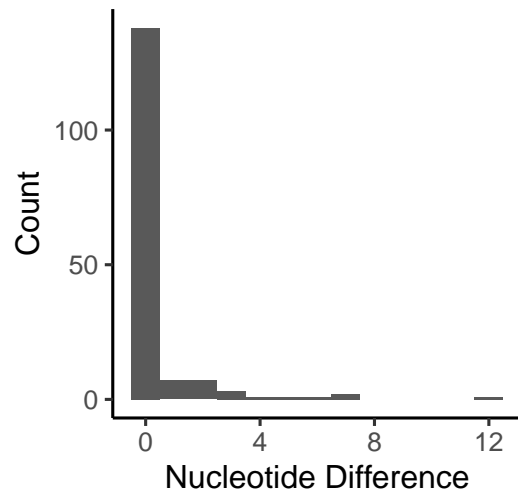
IGLV1-51*01

96 sequences assigned
79 (82.3%) exact matches, in which:
60 unique CDR3
4 unique J



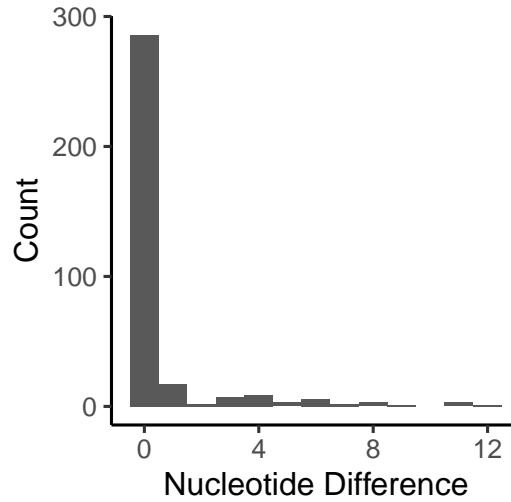
IGLV2-8*01

161 sequences assigned
138 (85.7%) exact matches, in which:
126 unique CDR3
4 unique J



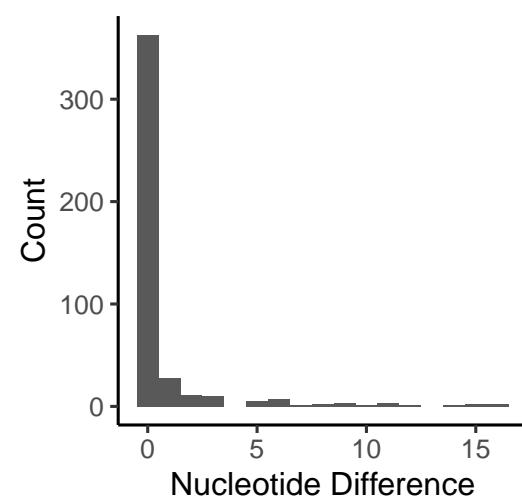
IGLV2-14*03EXT3

341 sequences assigned
286 (83.9%) exact matches, in which:
258 unique CDR3
3 unique J



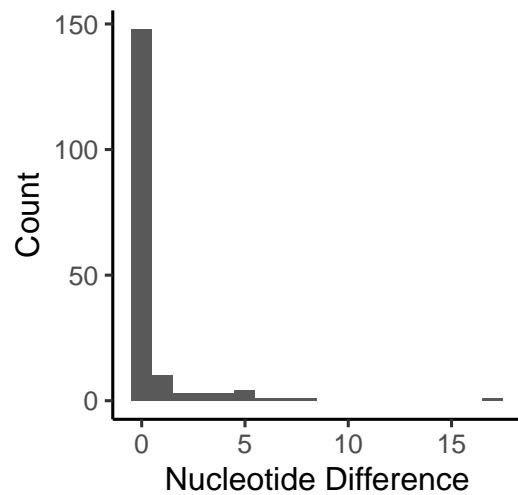
IGLV2-23*01

440 sequences assigned
363 (82.5%) exact matches, in which:
334 unique CDR3
4 unique J



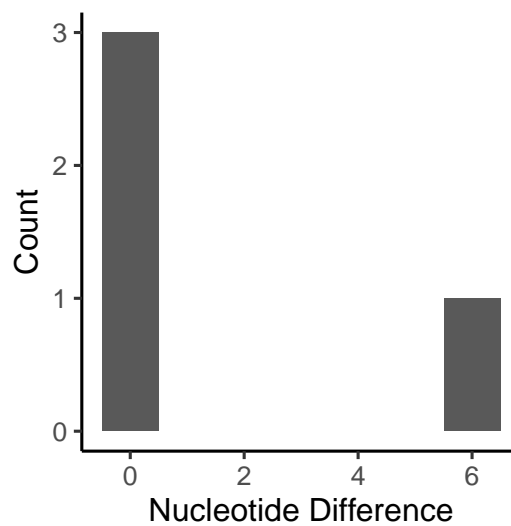
IGLV2-11*01

175 sequences assigned
148 (84.6%) exact matches, in which:
141 unique CDR3
3 unique J



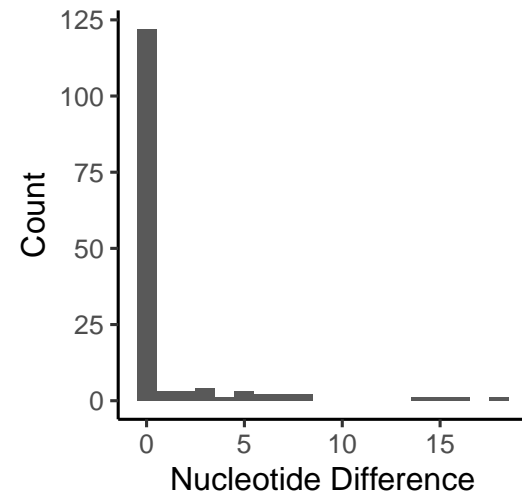
IGLV2-18*01

4 sequences assigned
3 (75%) exact matches, in which:
3 unique CDR3
2 unique J



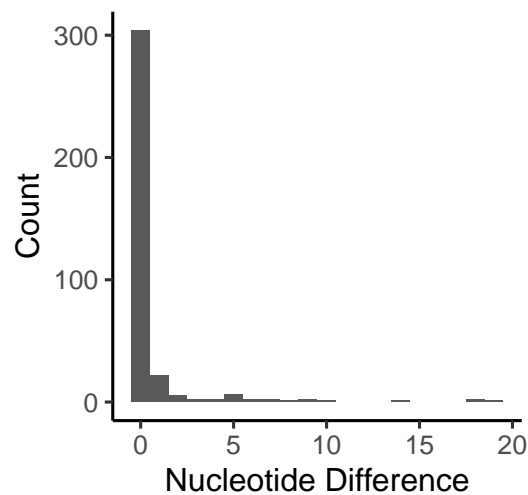
IGLV3-1*01

148 sequences assigned
122 (82.4%) exact matches, in which:
113 unique CDR3
4 unique J



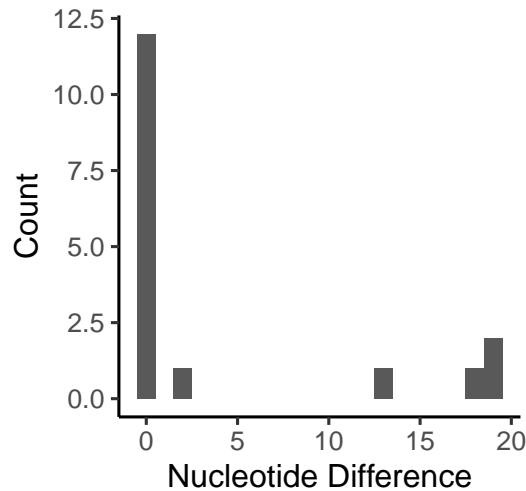
IGLV2-14*01

353 sequences assigned
304 (86.1%) exact matches, in which:
276 unique CDR3
3 unique J



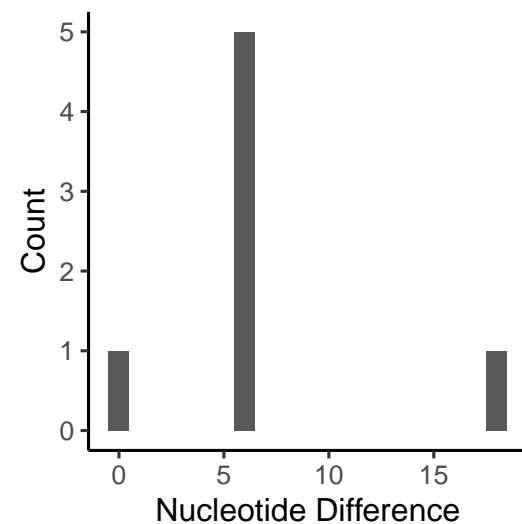
IGLV2-18*02

17 sequences assigned
12 (70.6%) exact matches, in which:
11 unique CDR3
3 unique J



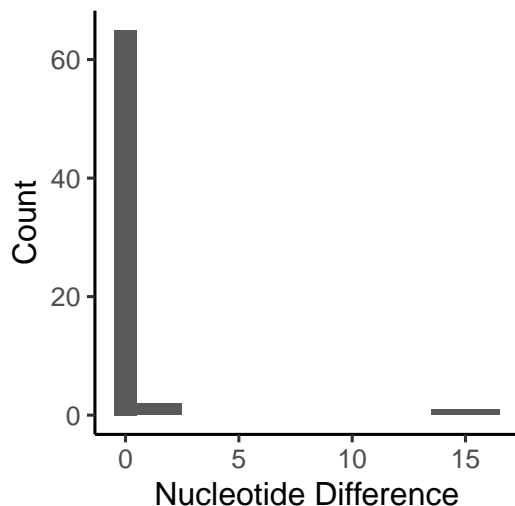
IGLV3-9*02

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



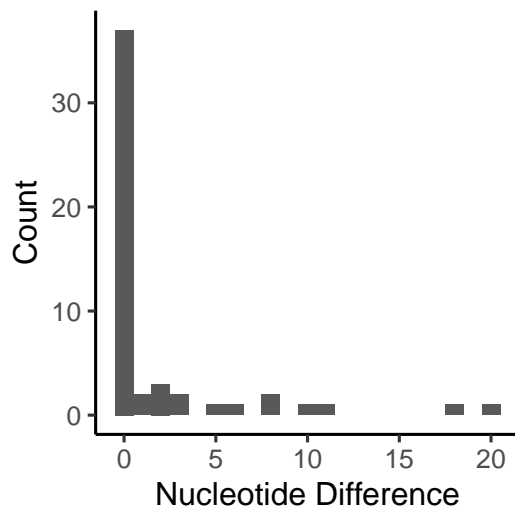
IGLV3-10*01

72 sequences assigned
65 (90.3%) exact matches, in which:
64 unique CDR3
3 unique J



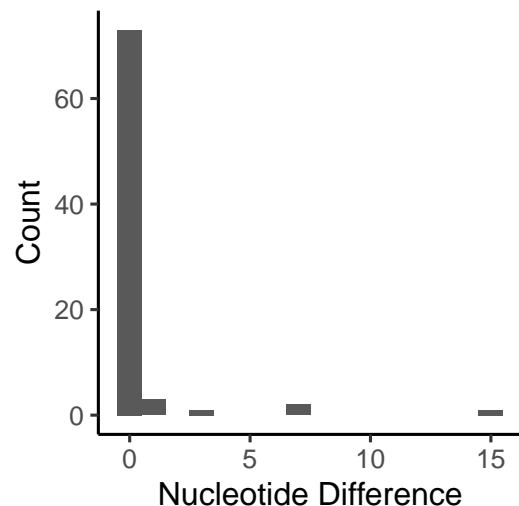
IGLV3-21*02

52 sequences assigned
37 (71.2%) exact matches, in which:
35 unique CDR3
4 unique J



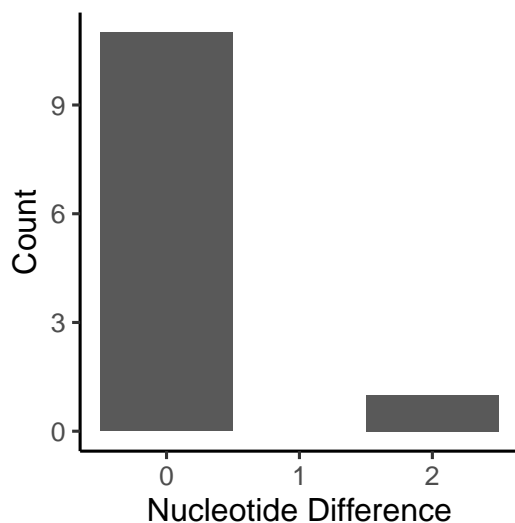
IGLV3-25*02

80 sequences assigned
73 (91.2%) exact matches, in which:
71 unique CDR3
3 unique J



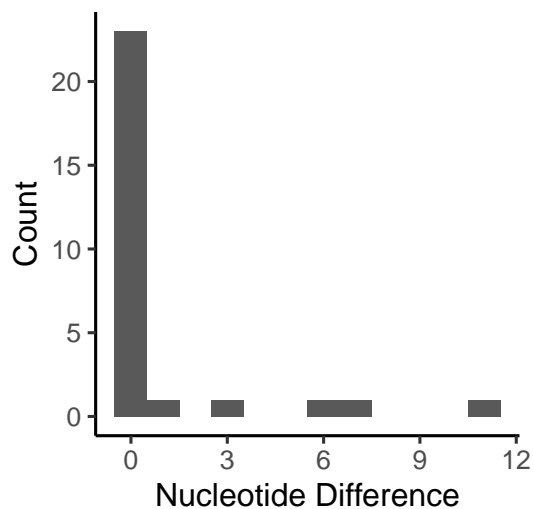
IGLV3-16*01

12 sequences assigned
11 (91.7%) exact matches, in which:
11 unique CDR3
4 unique J



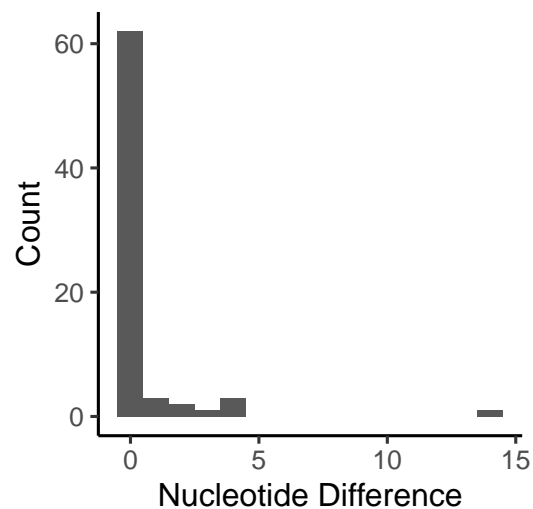
IGLV3-21*04

30 sequences assigned
23 (76.7%) exact matches, in which:
21 unique CDR3
3 unique J



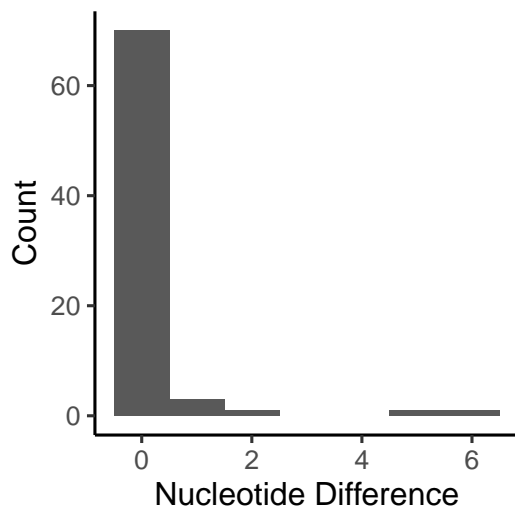
IGLV3-25*03

72 sequences assigned
62 (86.1%) exact matches, in which:
59 unique CDR3
5 unique J



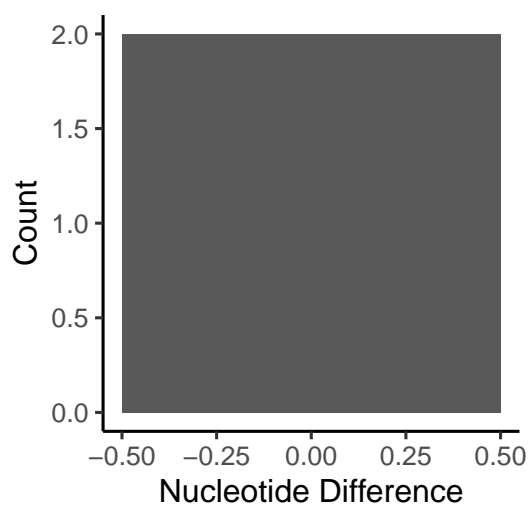
IGLV3-19*01

76 sequences assigned
70 (92.1%) exact matches, in which:
62 unique CDR3
3 unique J



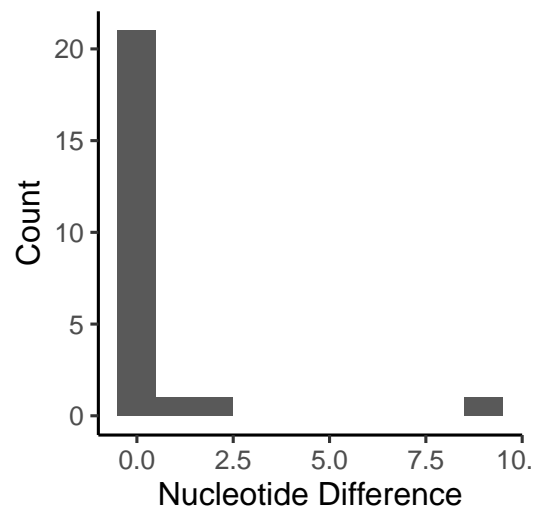
IGLV3-22*01

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



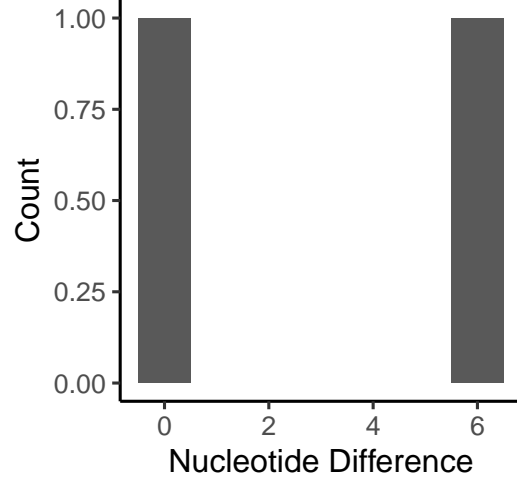
IGLV3-27*01

25 sequences assigned
21 (84%) exact matches, in which:
20 unique CDR3
2 unique J



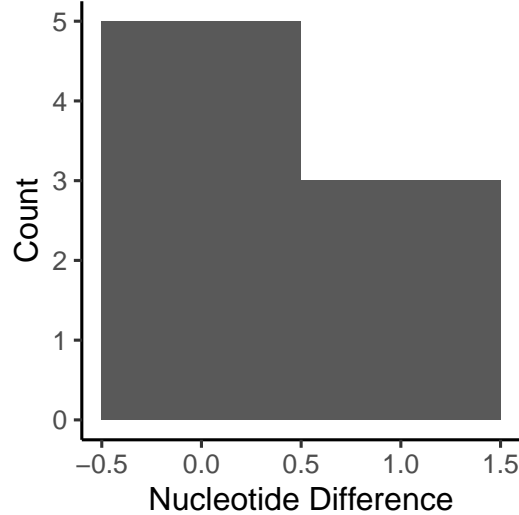
IGLV4-3*01

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



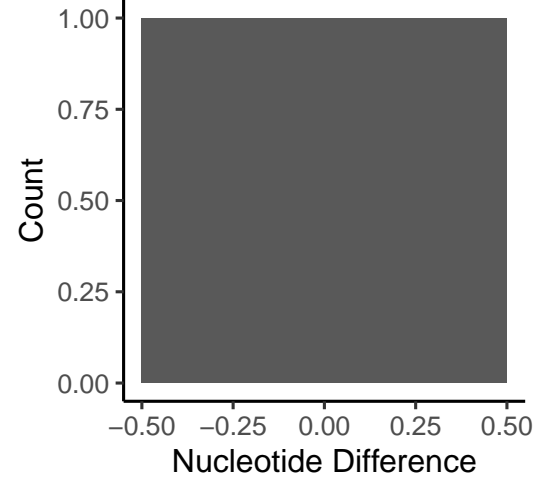
IGLV5-37*01

8 sequences assigned
5 (62.5%) exact matches, in which:
5 unique CDR3
2 unique J



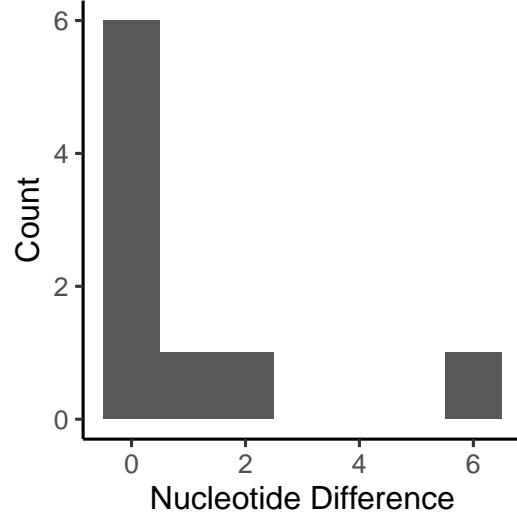
IGLV5-48*01

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



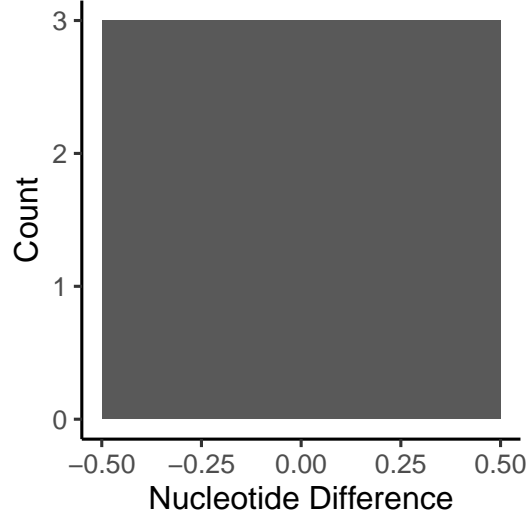
IGLV4-60*02

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



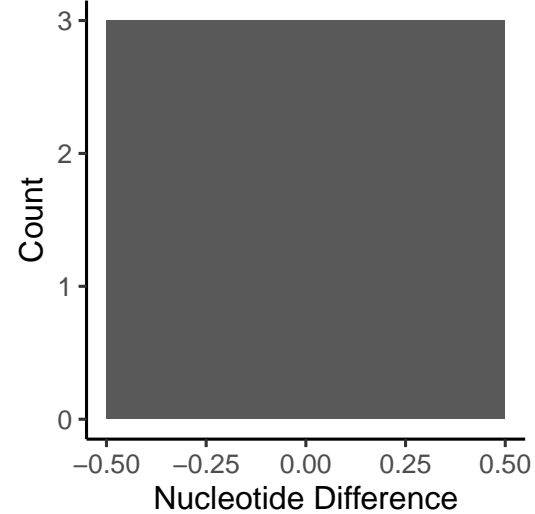
IGLV5-39*01

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



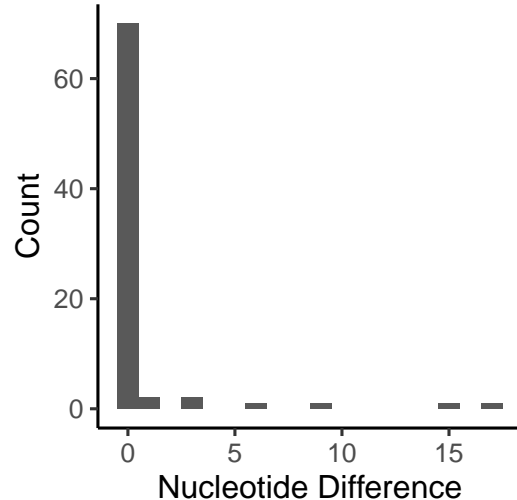
IGLV5-52*01

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



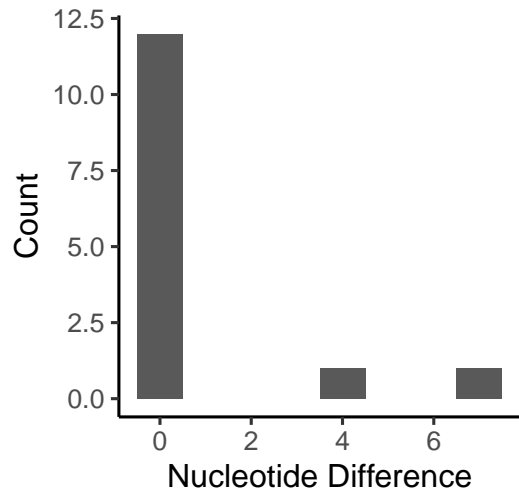
IGLV4-69*01

78 sequences assigned
70 (89.7%) exact matches, in which:
59 unique CDR3
4 unique J



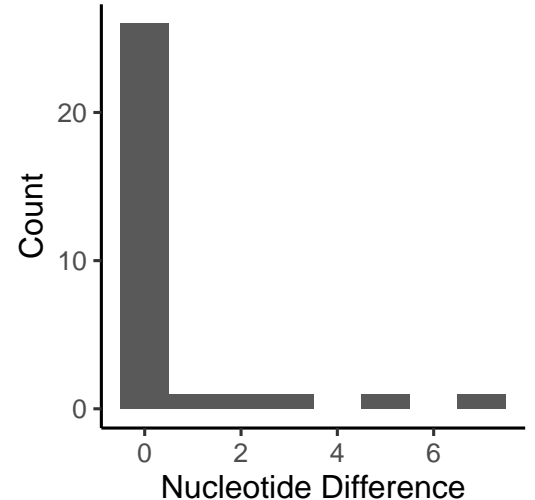
IGLV5-45*03

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



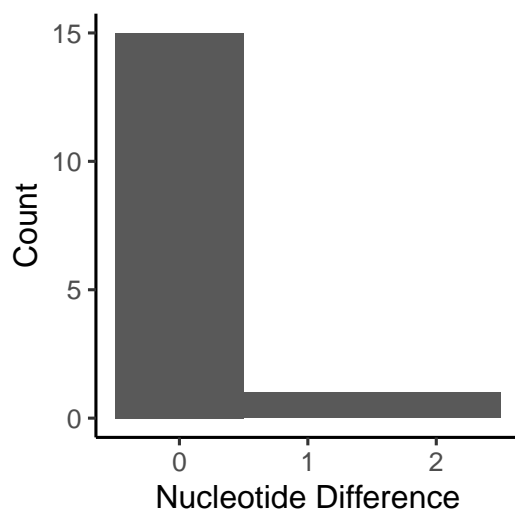
IGLV6-57*01

31 sequences assigned
26 (83.9%) exact matches, in which:
25 unique CDR3
2 unique J



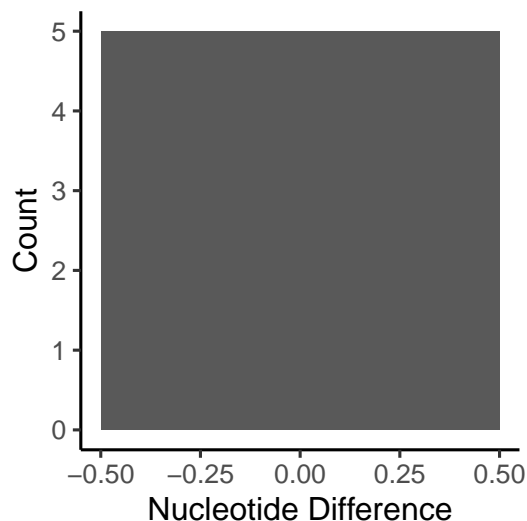
IGLV6-57*02

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



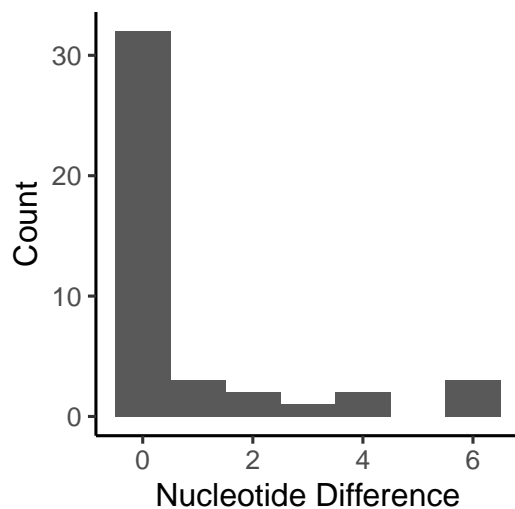
IGLV7-46*02

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



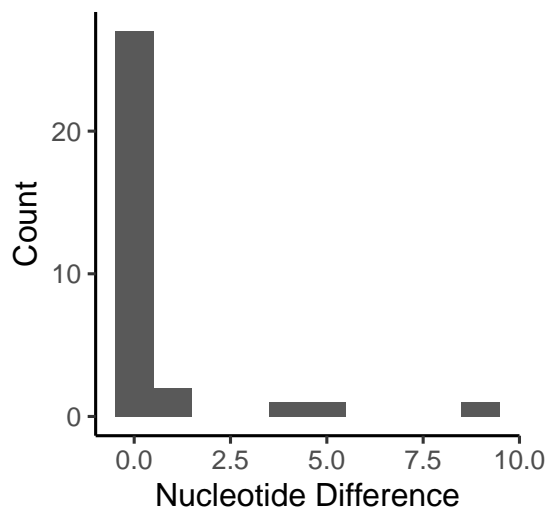
IGLV7-43*01

43 sequences assigned
32 (74.4%) exact matches, in which:
29 unique CDR3
3 unique J



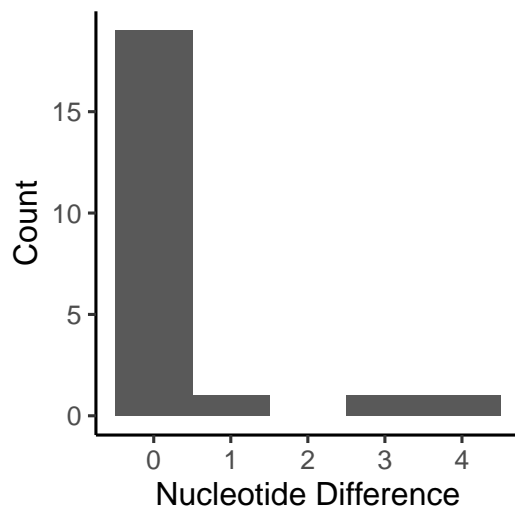
IGLV8-61*01

32 sequences assigned
27 (84.4%) exact matches, in which:
25 unique CDR3
4 unique J



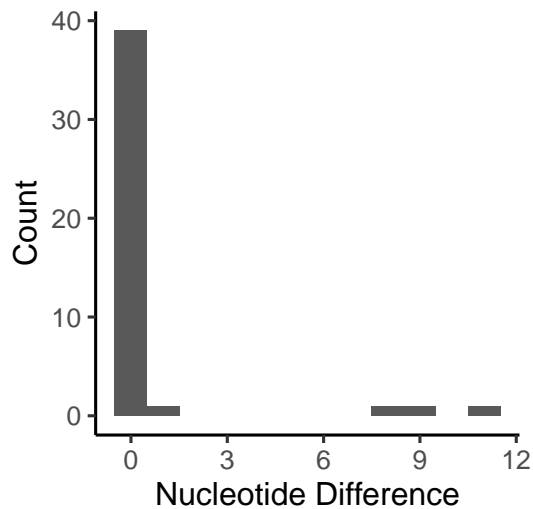
IGLV7-46*01

22 sequences assigned
19 (86.4%) exact matches, in which:
18 unique CDR3
3 unique J



IGLV9-49*01

44 sequences assigned
39 (88.6%) exact matches, in which:
36 unique CDR3
3 unique J



Allele Usage



Warning – no inferred sequences found.