

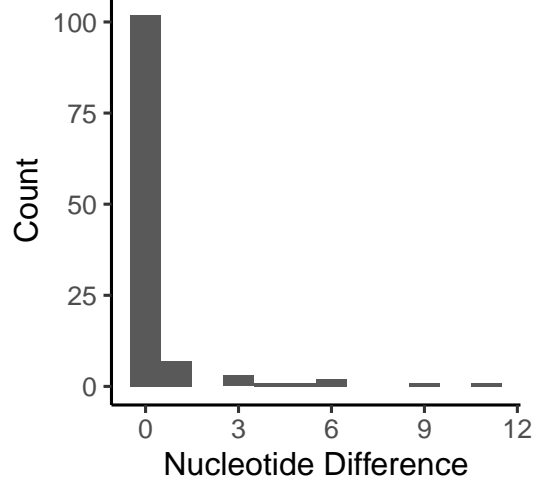
IGLV10-54*01

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



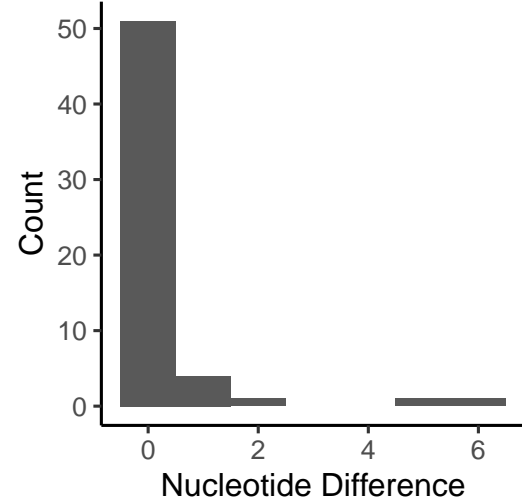
IGLV1-40*01

118 sequences assigned
102 (86.4%) exact matches, in which:
93 unique CDR3
3 unique J



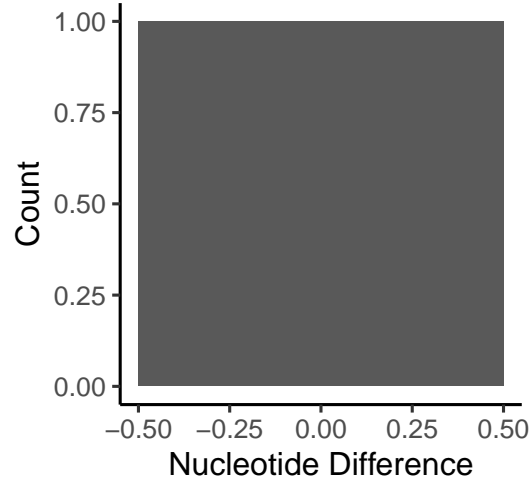
IGLV1-47*01

58 sequences assigned
51 (87.9%) exact matches, in which:
49 unique CDR3
3 unique J



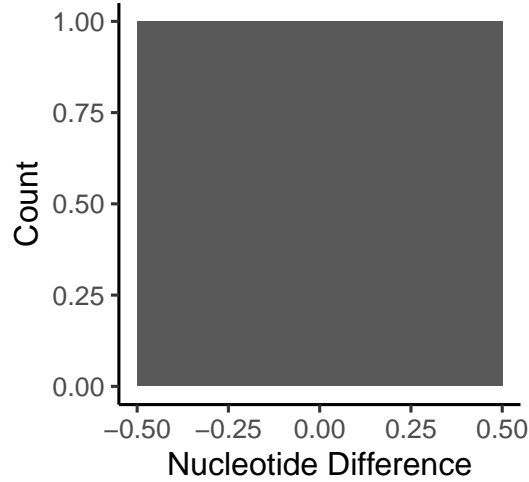
IGLV10-54*02

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



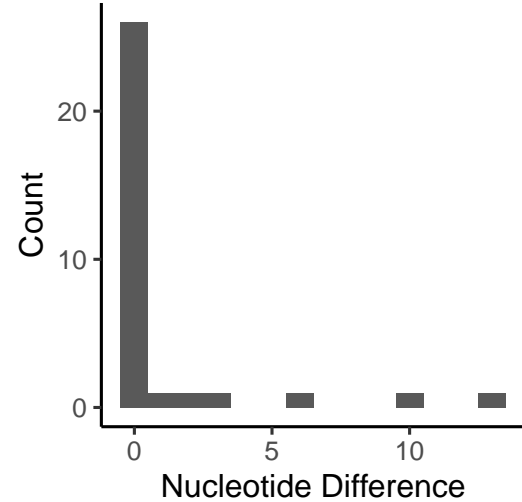
IGLV1-41*01

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



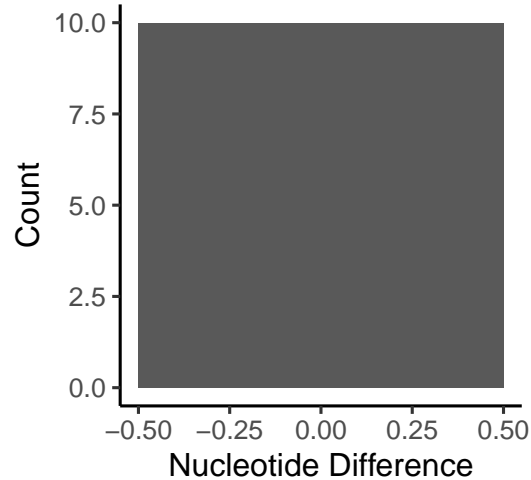
IGLV1-47*02

32 sequences assigned
26 (81.2%) exact matches, in which:
25 unique CDR3
2 unique J



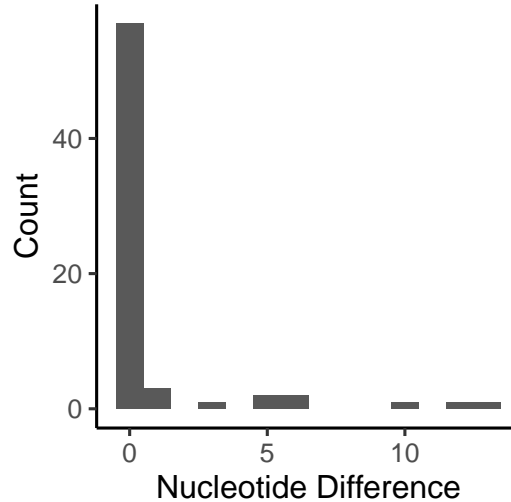
IGLV1-36*01

10 sequences assigned
10 (100%) exact matches, in which:
10 unique CDR3
2 unique J



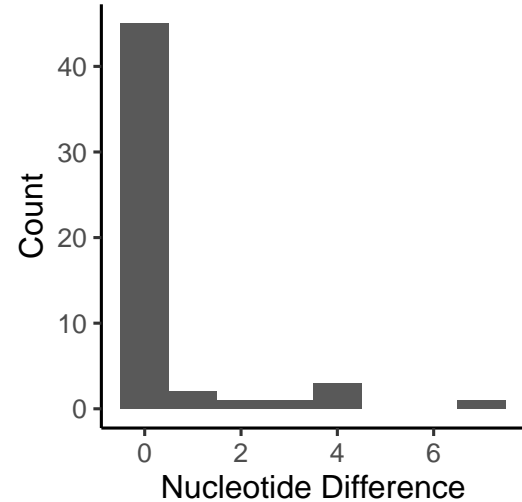
IGLV1-44*01

69 sequences assigned
57 (82.6%) exact matches, in which:
52 unique CDR3
3 unique J



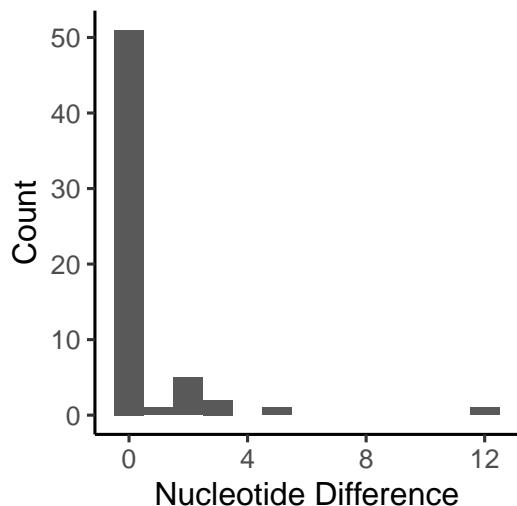
IGLV1-51*01

53 sequences assigned
45 (84.9%) exact matches, in which:
37 unique CDR3
3 unique J



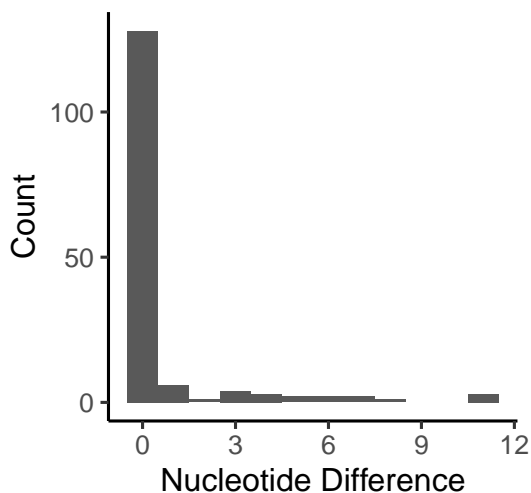
IGLV2-8*01

61 sequences assigned
51 (83.6%) exact matches, in which:
50 unique CDR3
3 unique J



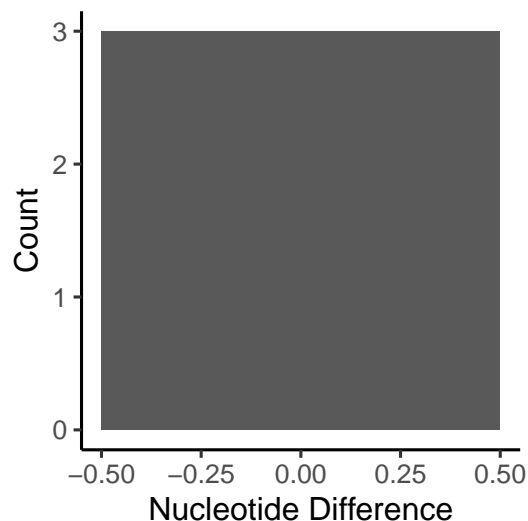
IGLV2-14*03EXT3

153 sequences assigned
128 (83.7%) exact matches, in which:
127 unique CDR3
3 unique J



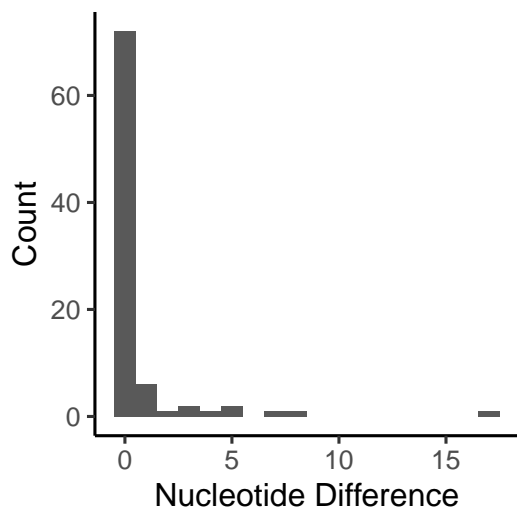
IGLV2-23*01

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



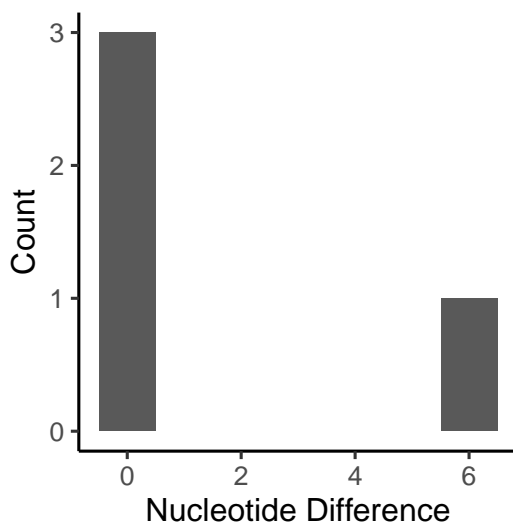
IGLV2-11*01

87 sequences assigned
72 (82.8%) exact matches, in which:
69 unique CDR3
2 unique J



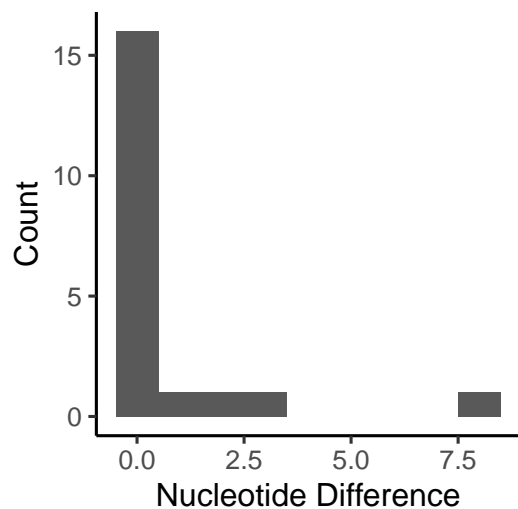
IGLV2-18*01

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



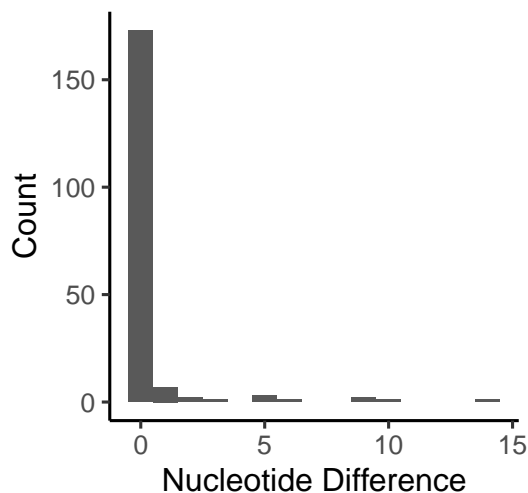
IGLV2-23*03EXT5

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



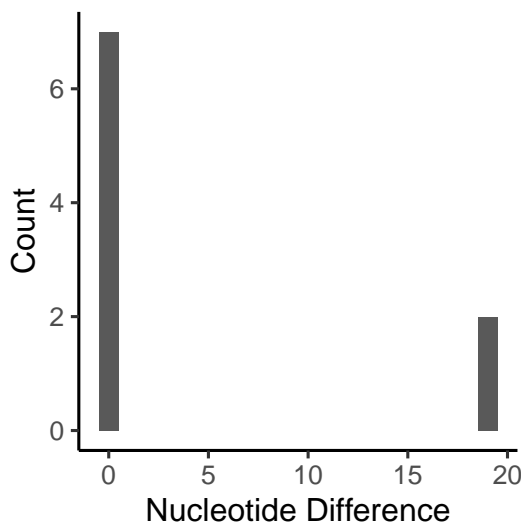
IGLV2-14*01

191 sequences assigned
173 (90.6%) exact matches, in which:
167 unique CDR3
2 unique J



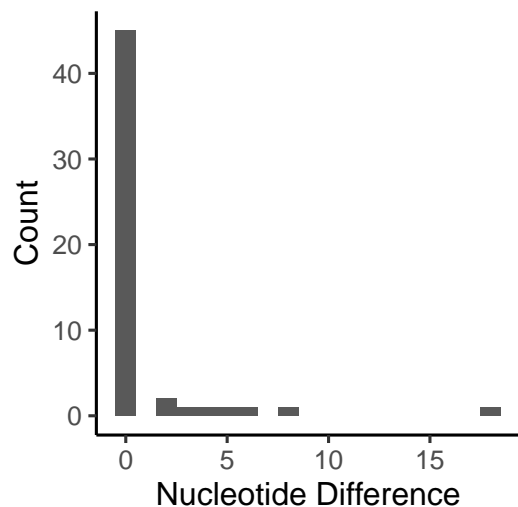
IGLV2-18*02

9 sequences assigned
7 (77.8%) exact matches, in which:
6 unique CDR3
2 unique J



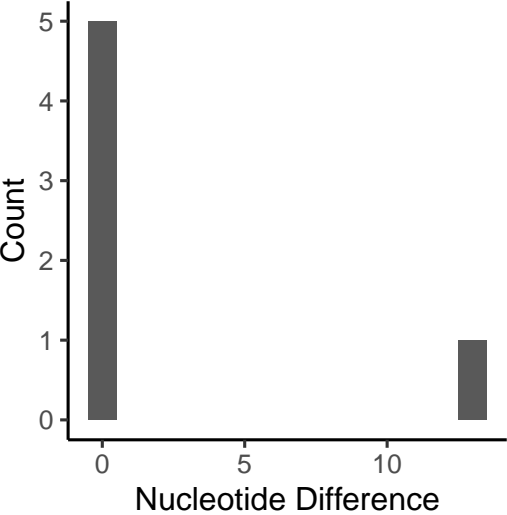
IGLV3-1*01

53 sequences assigned
45 (84.9%) exact matches, in which:
45 unique CDR3
3 unique J



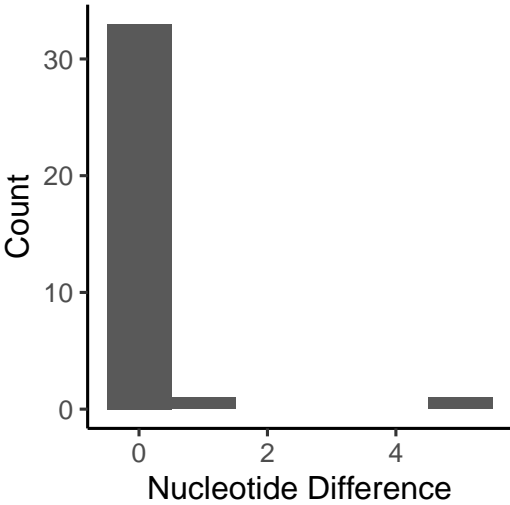
IGLV3-9*01EXT5

6 sequences assigned
5 (83.3%) exact matches, in which:
5 unique CDR3
2 unique J



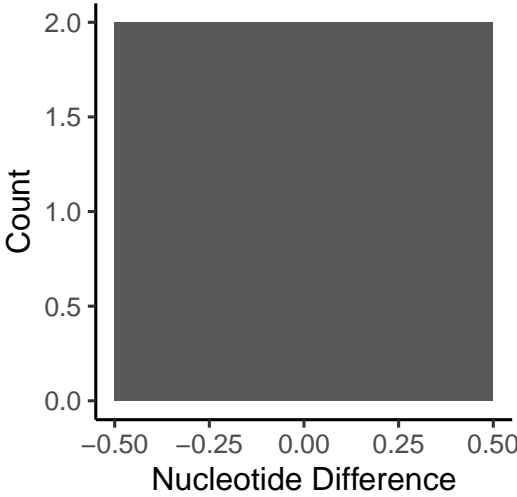
IGLV3-19*01

35 sequences assigned
33 (94.3%) exact matches, in which:
29 unique CDR3
2 unique J



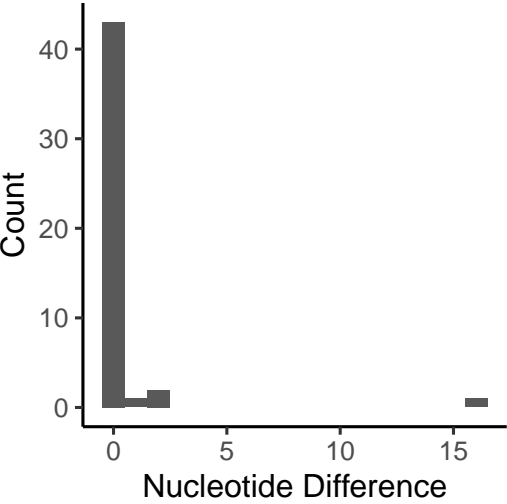
IGLV3-22*01

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



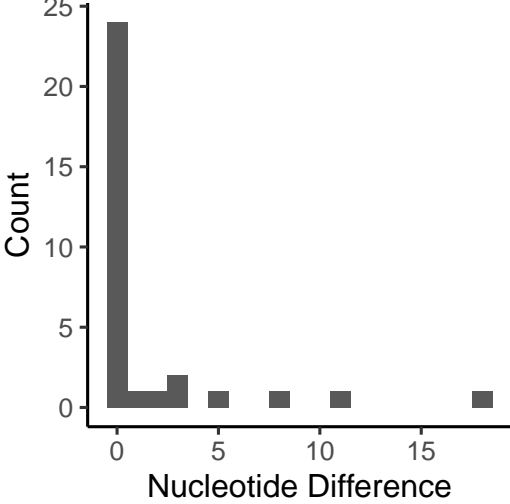
IGLV3-10*01

47 sequences assigned
43 (91.5%) exact matches, in which:
42 unique CDR3
2 unique J



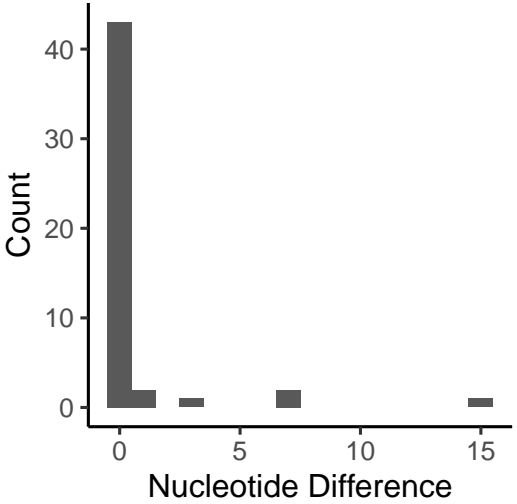
IGLV3-21*02

32 sequences assigned
24 (75%) exact matches, in which:
22 unique CDR3
3 unique J



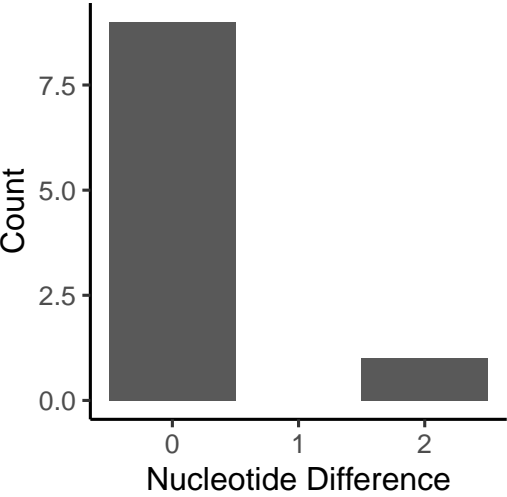
IGLV3-25*02

49 sequences assigned
43 (87.8%) exact matches, in which:
41 unique CDR3
2 unique J



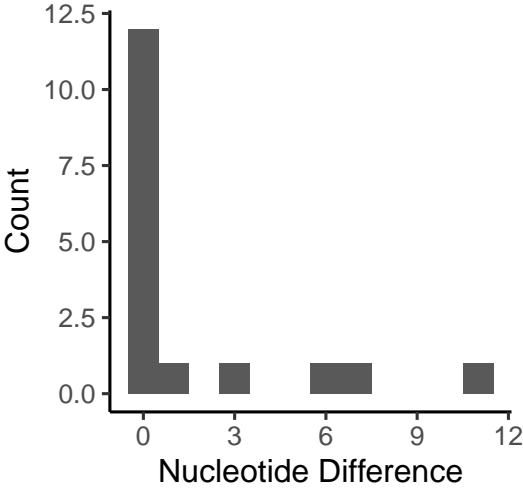
IGLV3-16*01

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



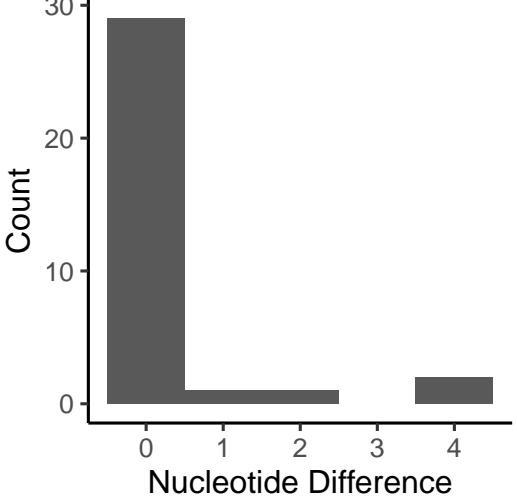
IGLV3-21*04

18 sequences assigned
12 (66.7%) exact matches, in which:
12 unique CDR3
2 unique J



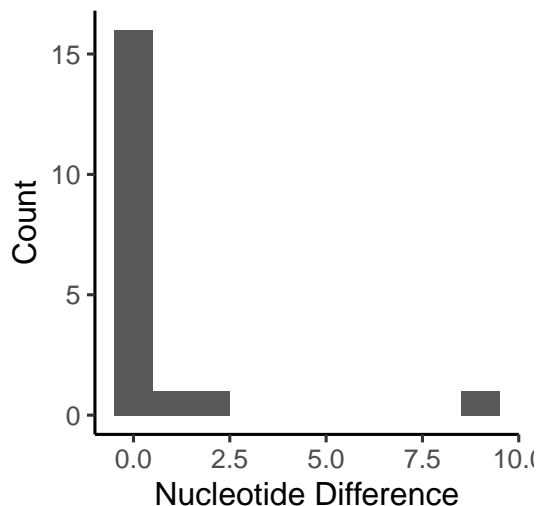
IGLV3-25*03

33 sequences assigned
29 (87.9%) exact matches, in which:
29 unique CDR3
4 unique J



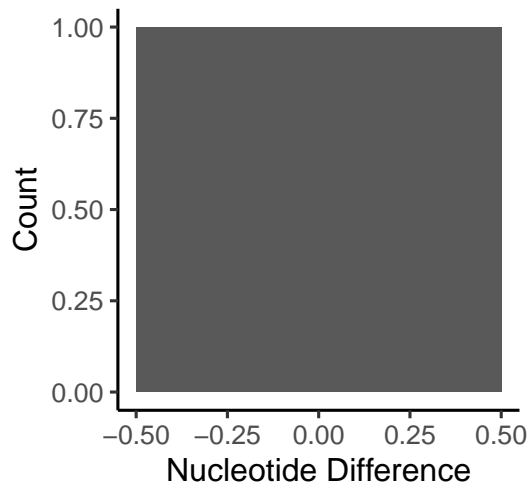
IGLV3-27*01

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



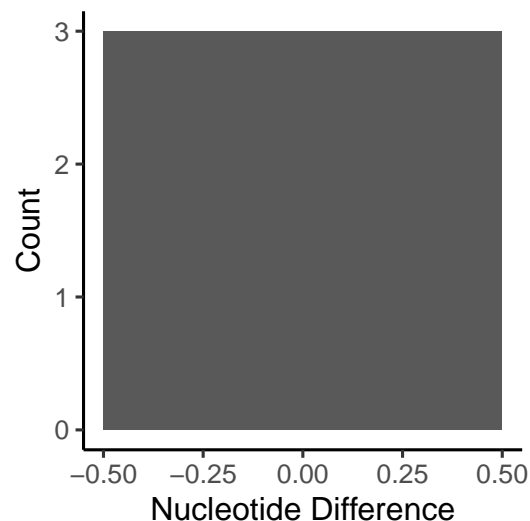
IGLV4-60*03EXT5

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



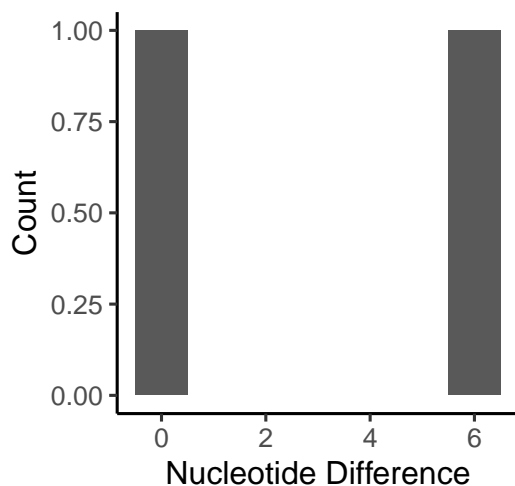
IGLV5-39*01

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



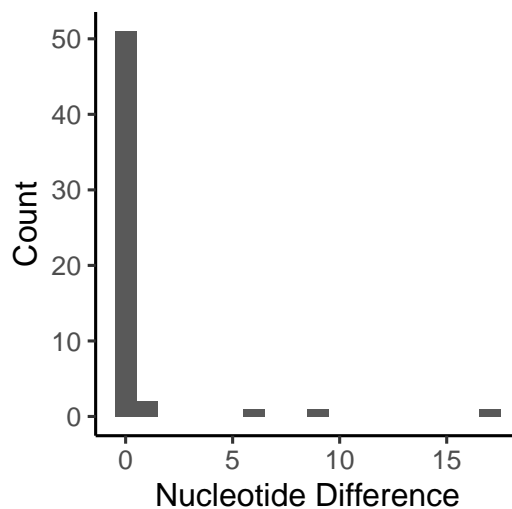
IGLV4-3*01

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



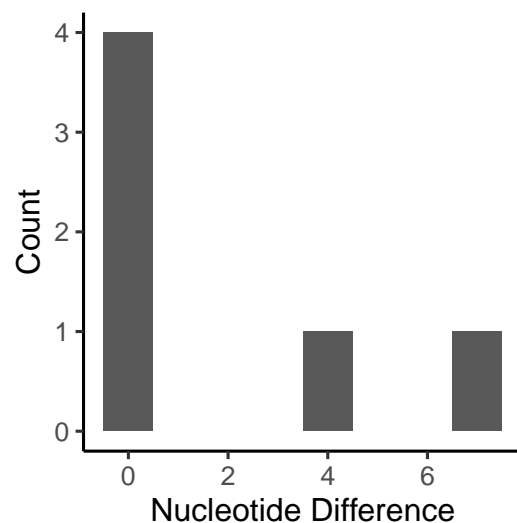
IGLV4-69*01

56 sequences assigned
51 (91.1%) exact matches, in which:
43 unique CDR3
3 unique J



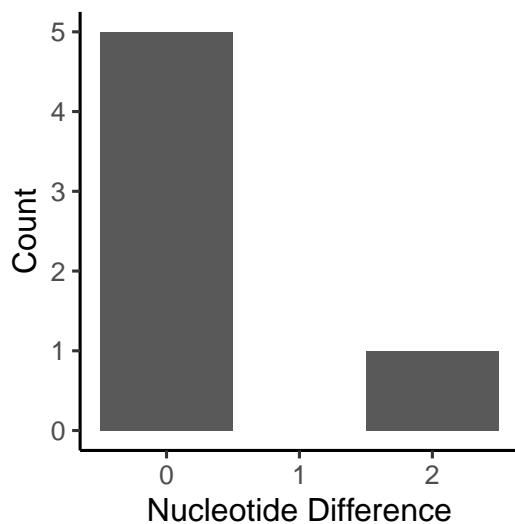
IGLV5-45*03

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



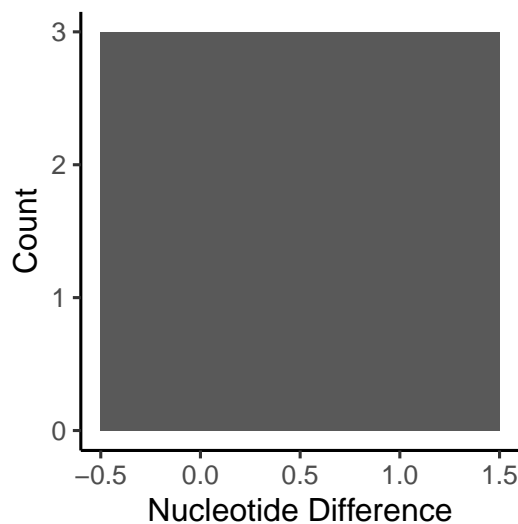
IGLV4-60*02

6 sequences assigned
5 (83.3%) exact matches, in which:
5 unique CDR3
1 unique J



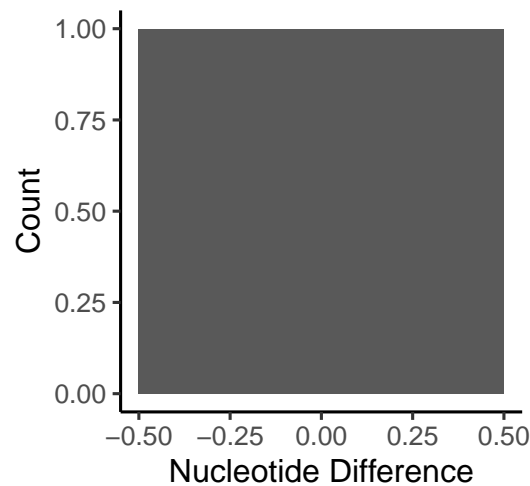
IGLV5-37*01

6 sequences assigned
3 (50%) exact matches, in which:
3 unique CDR3
1 unique J



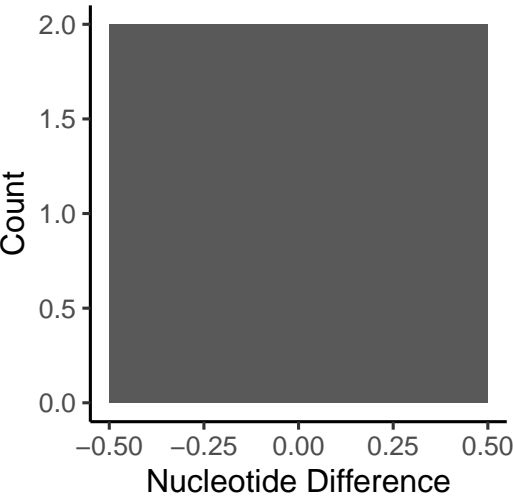
IGLV5-48*01

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



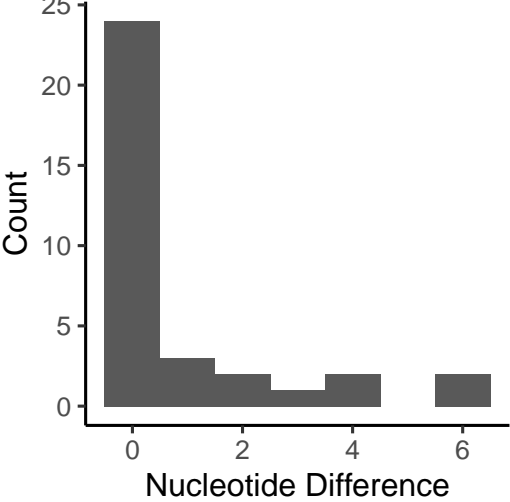
IGLV5-52*01

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



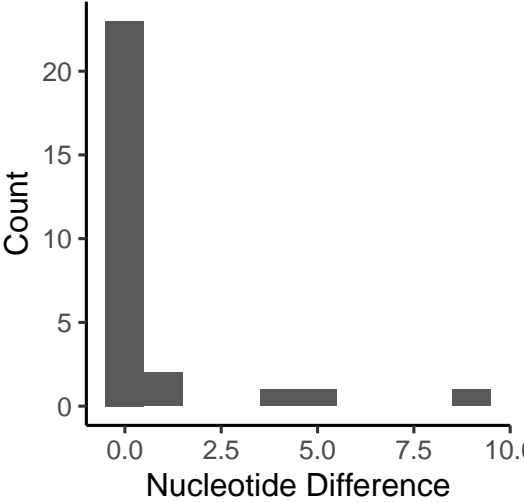
IGLV7-43*01

34 sequences assigned
24 (70.6%) exact matches, in which:
21 unique CDR3
2 unique J



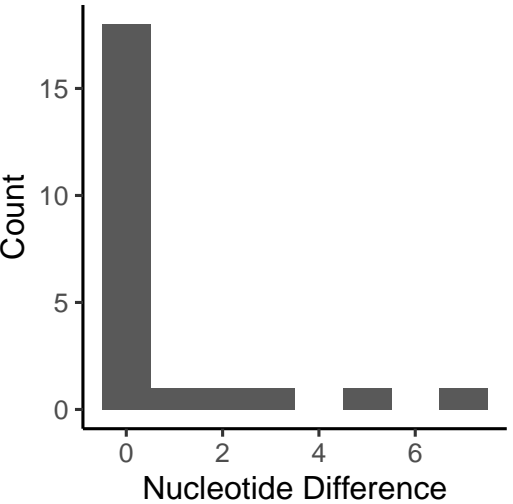
IGLV8-61*01

28 sequences assigned
23 (82.1%) exact matches, in which:
21 unique CDR3
3 unique J



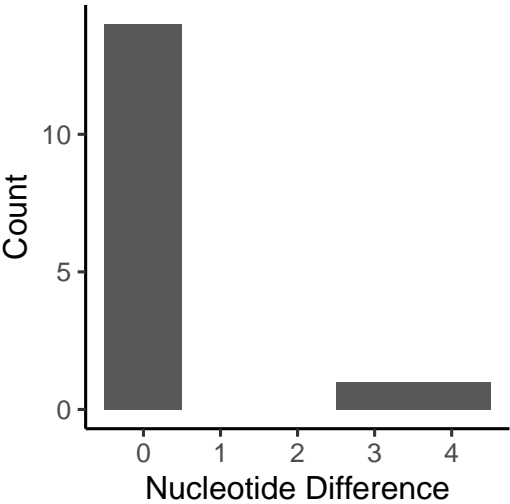
IGLV6-57*01

23 sequences assigned
18 (78.3%) exact matches, in which:
18 unique CDR3
1 unique J



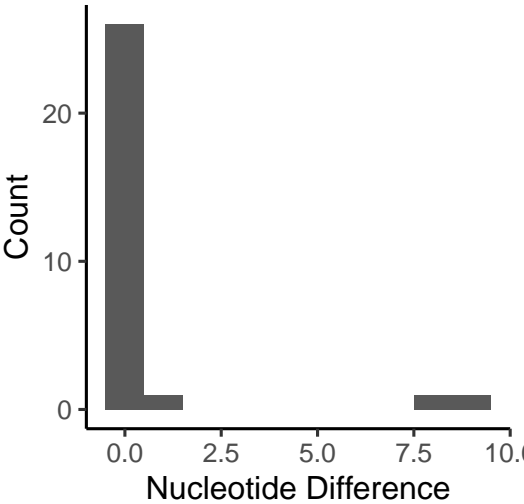
IGLV7-46*01

16 sequences assigned
14 (87.5%) exact matches, in which:
13 unique CDR3
2 unique J



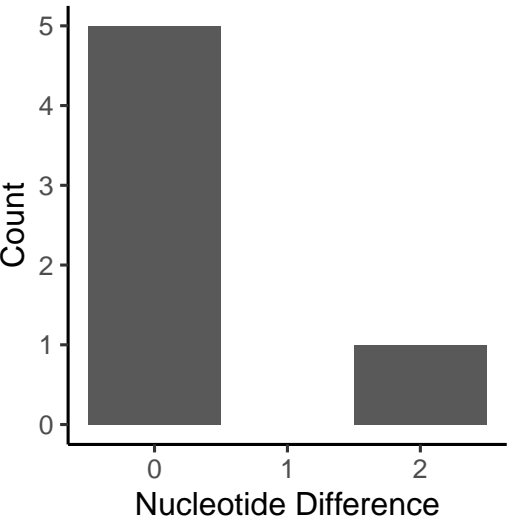
IGLV9-49*01

29 sequences assigned
26 (89.7%) exact matches, in which:
23 unique CDR3
2 unique J



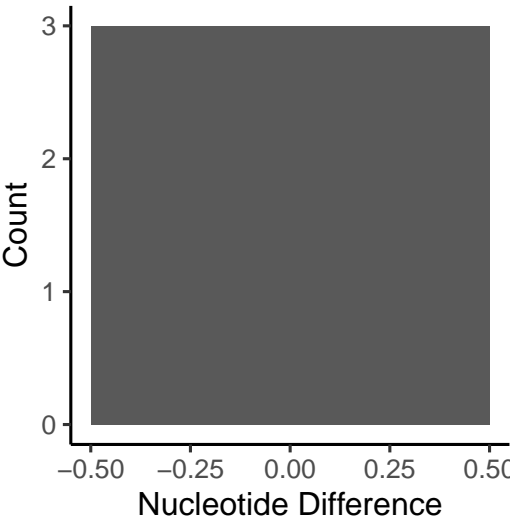
IGLV6-57*02

6 sequences assigned
5 (83.3%) exact matches, in which:
5 unique CDR3
1 unique J



IGLV7-46*02

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



Allele Usage



Warning – no inferred sequences found.