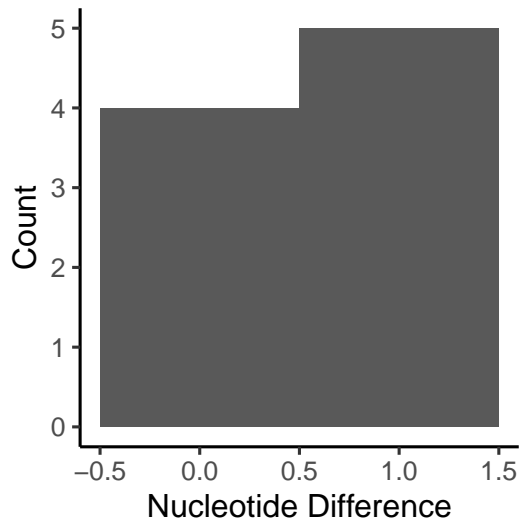


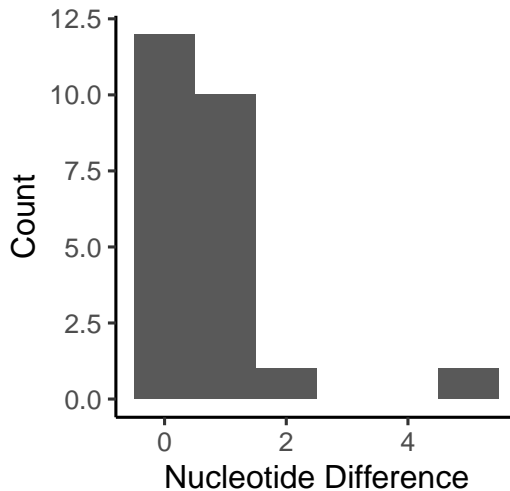
IGLV1-36*01

9 sequences assigned
4 (44.4%) exact matches, in which:
3 unique CDR3
2 unique J



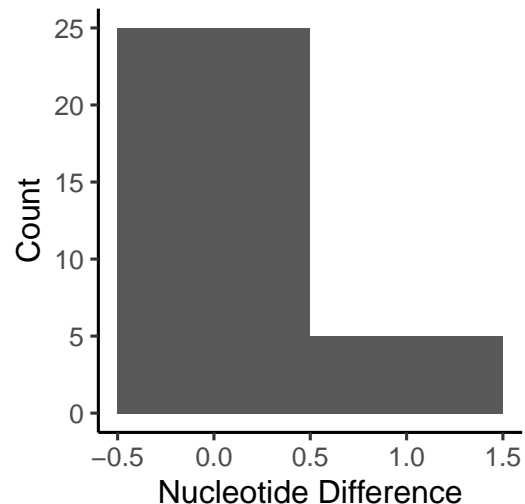
IGLV1-47*01

24 sequences assigned
12 (50%) exact matches, in which:
11 unique CDR3
2 unique J



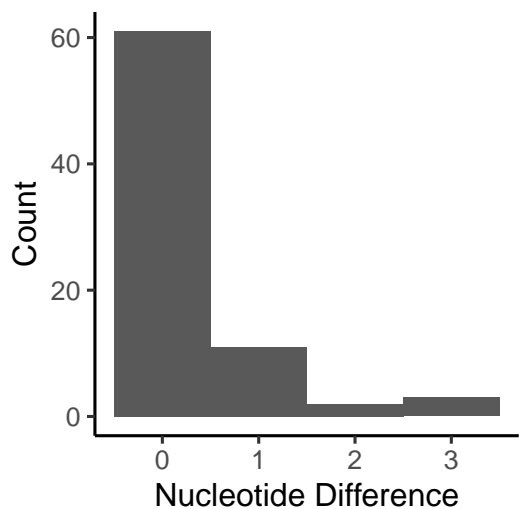
IGLV2-8*01

30 sequences assigned
25 (83.3%) exact matches, in which:
25 unique CDR3
2 unique J



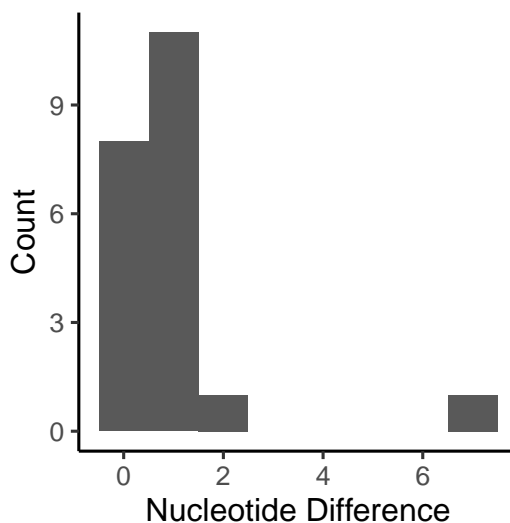
IGLV1-40*01

77 sequences assigned
61 (79.2%) exact matches, in which:
59 unique CDR3
3 unique J



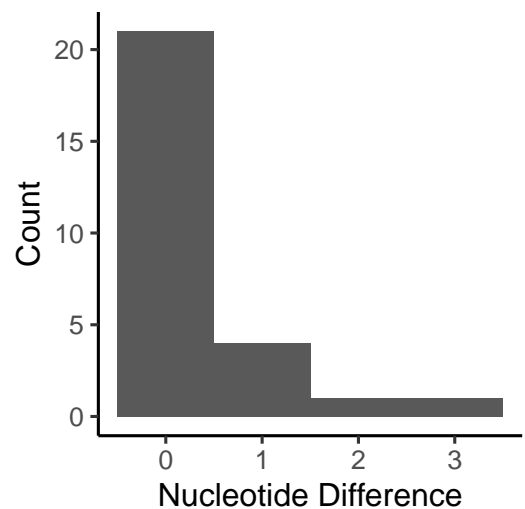
IGLV1-47*02

21 sequences assigned
8 (38.1%) exact matches, in which:
7 unique CDR3
2 unique J



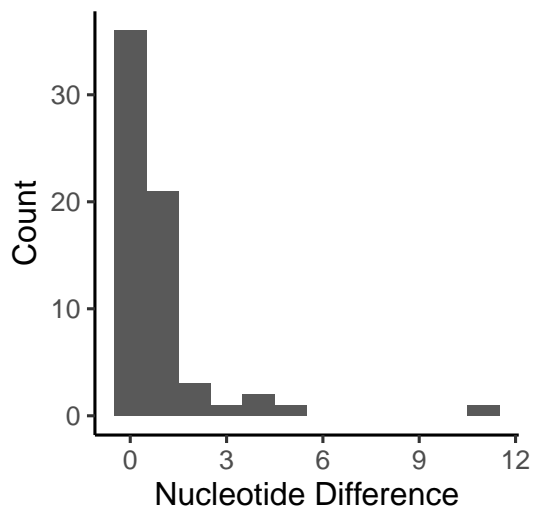
IGLV2-11*01

27 sequences assigned
21 (77.8%) exact matches, in which:
20 unique CDR3
2 unique J



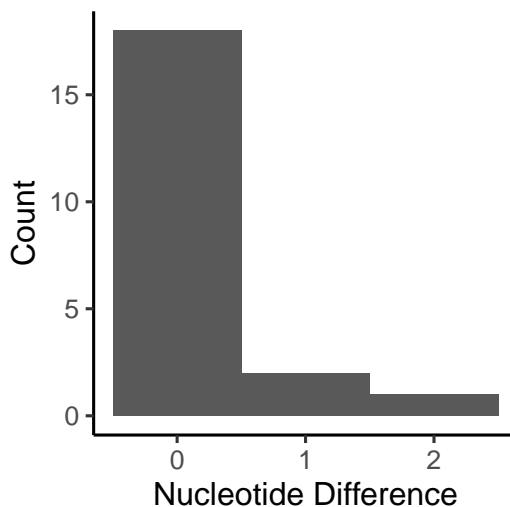
IGLV1-44*01

65 sequences assigned
36 (55.4%) exact matches, in which:
30 unique CDR3
3 unique J



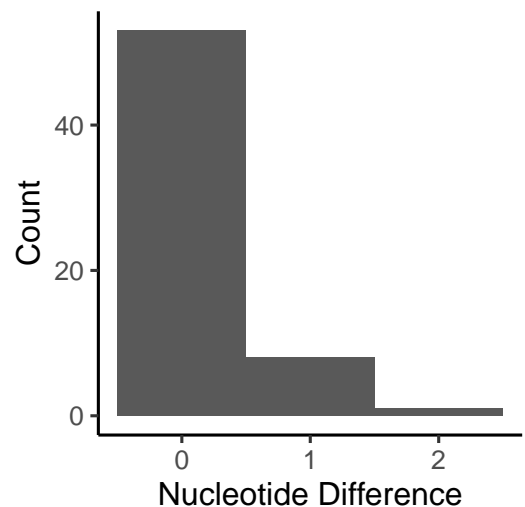
IGLV1-51*01

21 sequences assigned
18 (85.7%) exact matches, in which:
17 unique CDR3
3 unique J



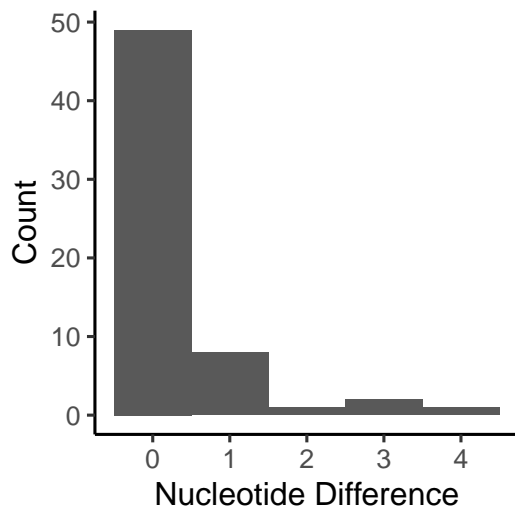
IGLV2-14*01

62 sequences assigned
53 (85.5%) exact matches, in which:
52 unique CDR3
2 unique J



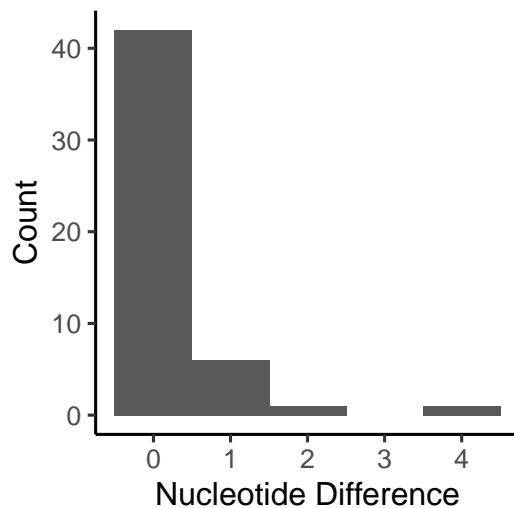
IGLV2-14*04EXT3

61 sequences assigned
49 (80.3%) exact matches, in which:
47 unique CDR3
2 unique J



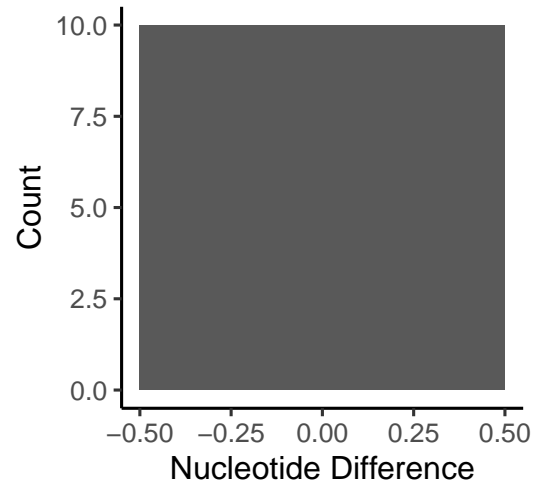
IGLV3-1*01

50 sequences assigned
42 (84%) exact matches, in which:
40 unique CDR3
3 unique J



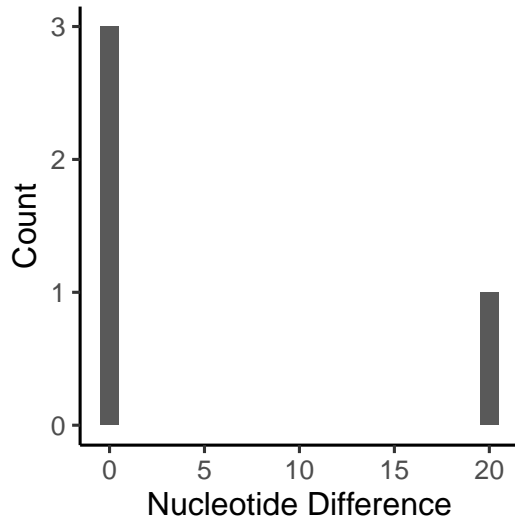
IGLV3-16*01

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



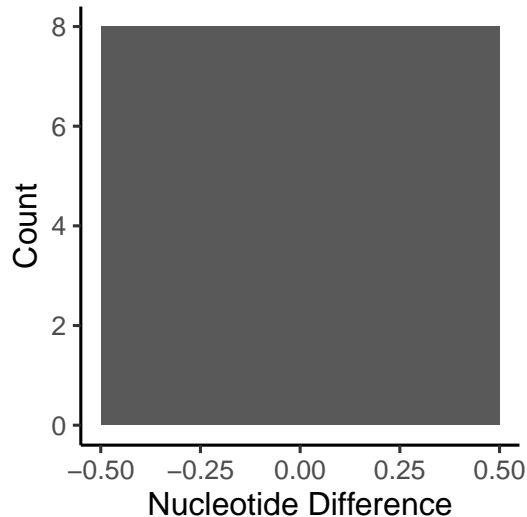
IGLV2-18*02

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



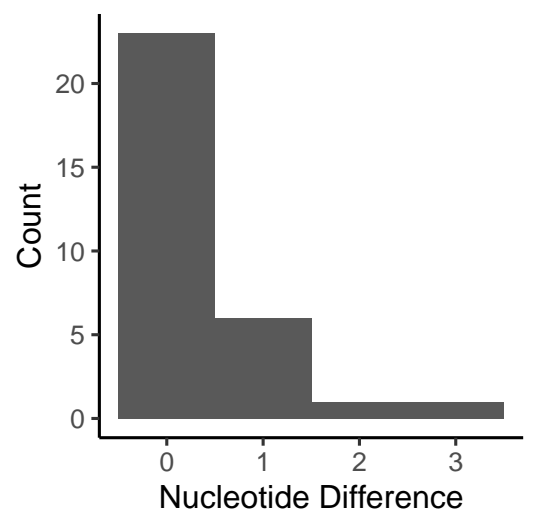
IGLV3-9*01EXT5

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



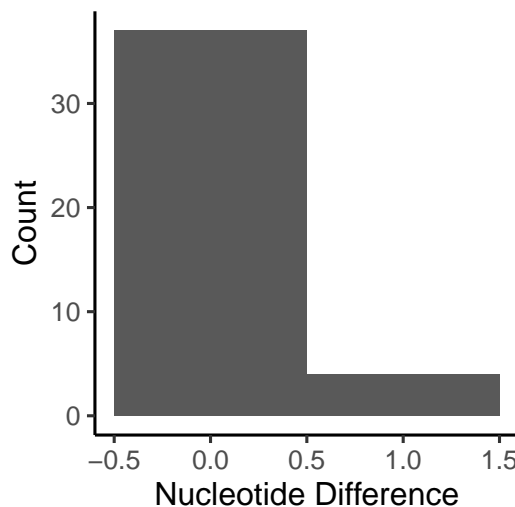
IGLV3-19*01

31 sequences assigned
23 (74.2%) exact matches, in which:
22 unique CDR3
2 unique J



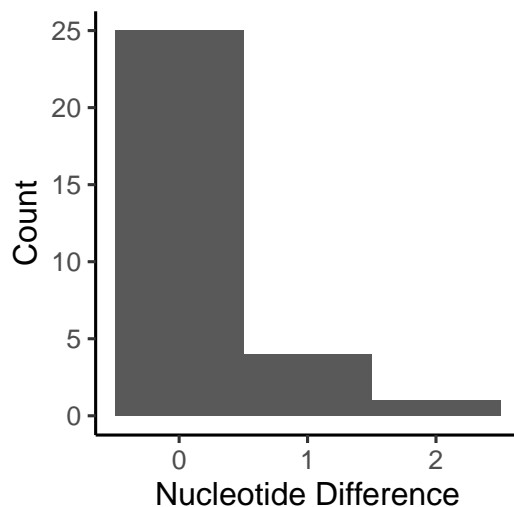
IGLV2-23*02EXT5

41 sequences assigned
37 (90.2%) exact matches, in which:
36 unique CDR3
3 unique J



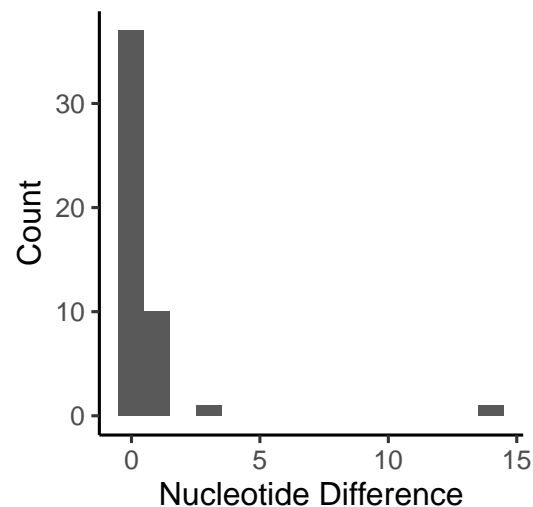
IGLV3-10*01

30 sequences assigned
25 (83.3%) exact matches, in which:
25 unique CDR3
2 unique J



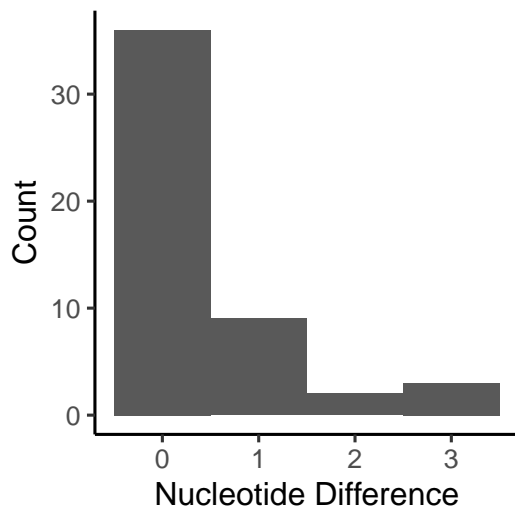
IGLV3-21*02

49 sequences assigned
37 (75.5%) exact matches, in which:
34 unique CDR3
2 unique J



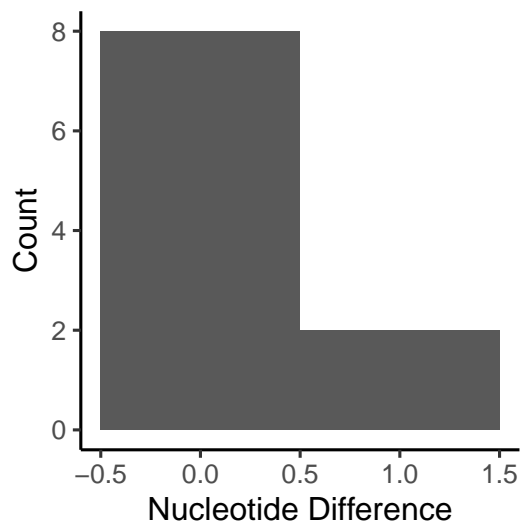
IGLV3–25*03

50 sequences assigned
36 (72%) exact matches, in which:
36 unique CDR3
2 unique J



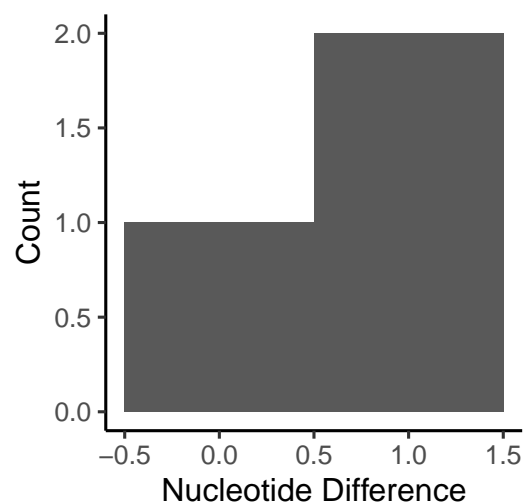
IGLV4–69*01

10 sequences assigned
8 (80%) exact matches, in which:
8 unique CDR3
1 unique J



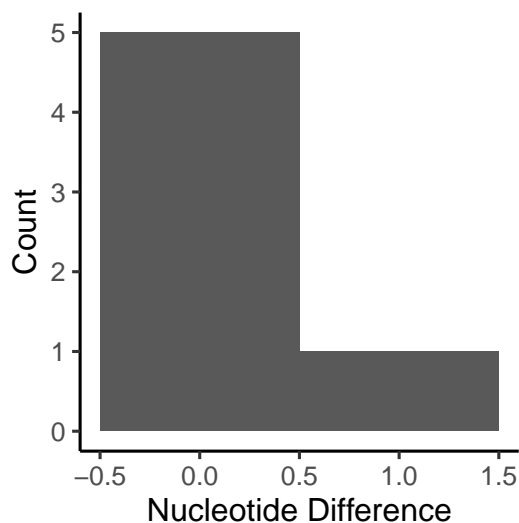
IGLV5–45*02

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



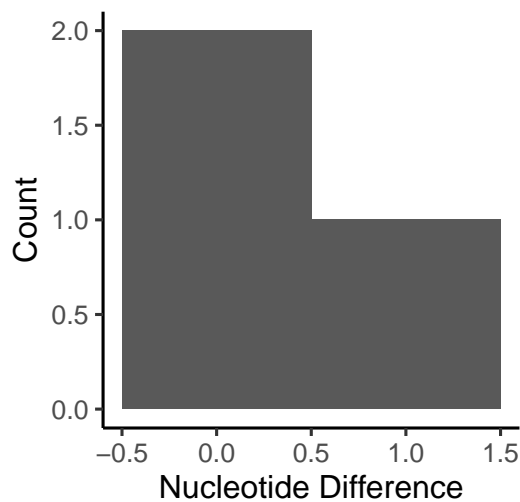
IGLV3–27*01

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



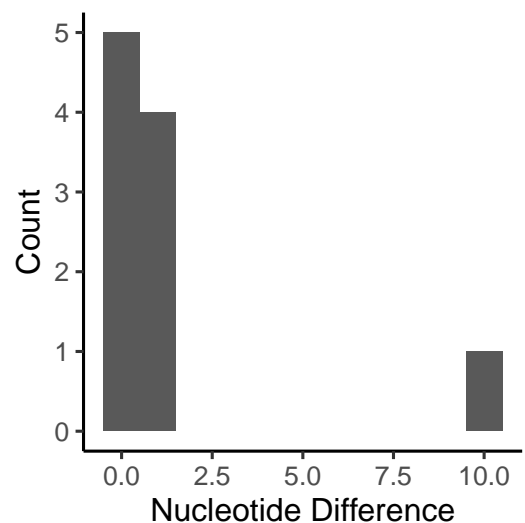
IGLV5–37*01

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



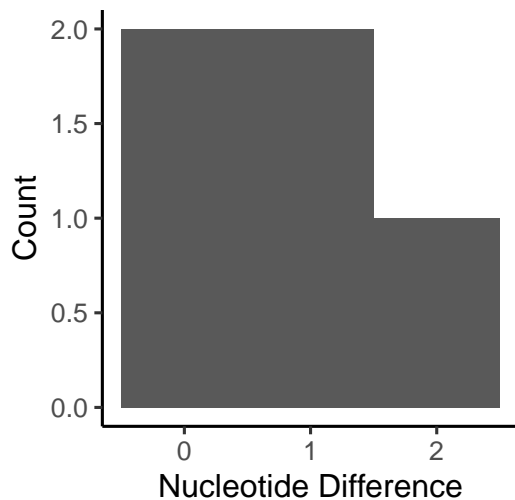
IGLV5–45*03

10 sequences assigned
5 (50%) exact matches, in which:
4 unique CDR3
2 unique J



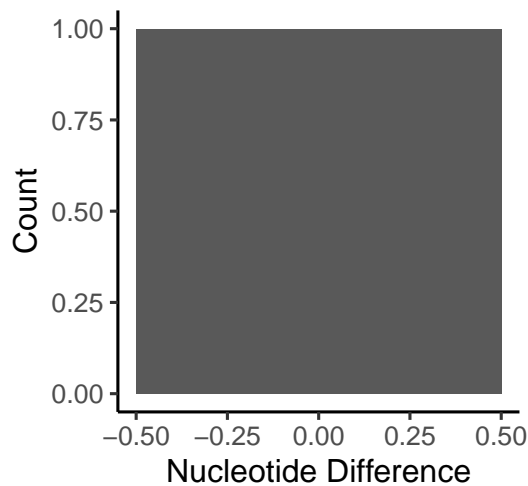
IGLV4–60*03EXT5

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



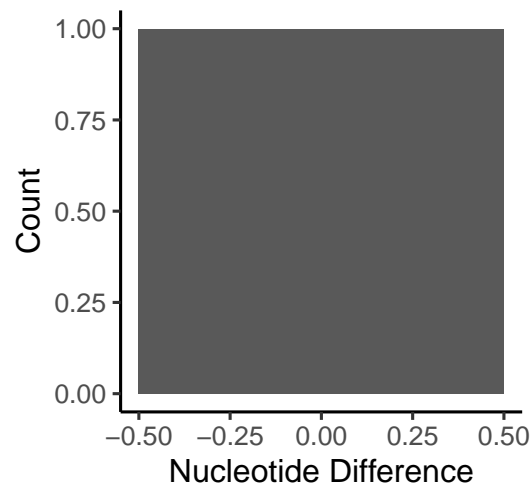
IGLV5–39*01

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



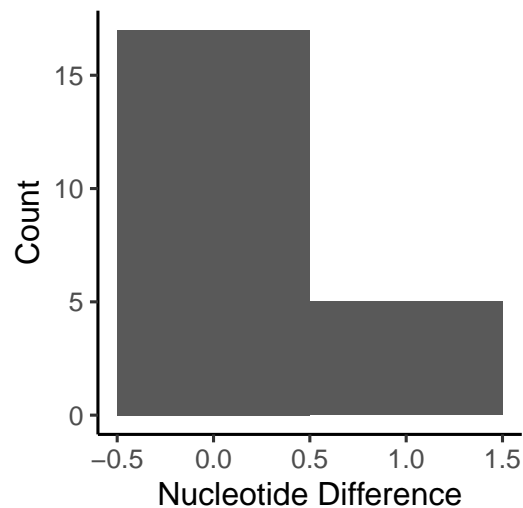
IGLV5–52*01

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



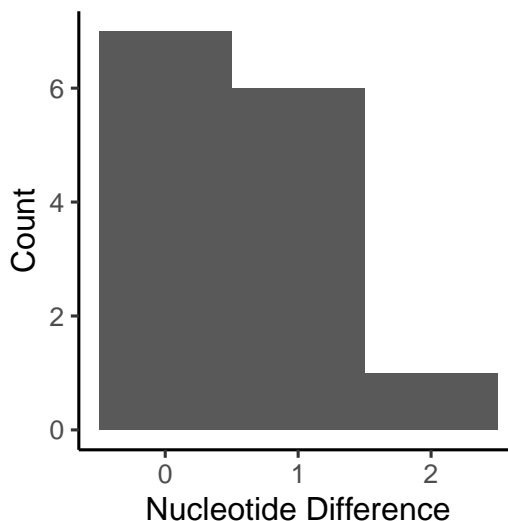
IGLV6-57*01

22 sequences assigned
17 (77.3%) exact matches, in which:
15 unique CDR3
4 unique J



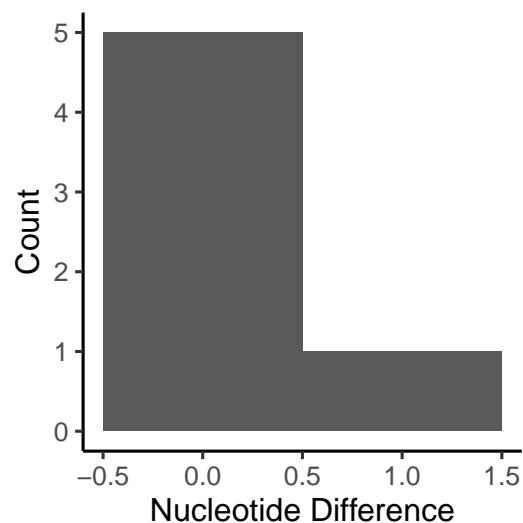
IGLV7-46*01

14 sequences assigned
7 (50%) exact matches, in which:
7 unique CDR3
1 unique J



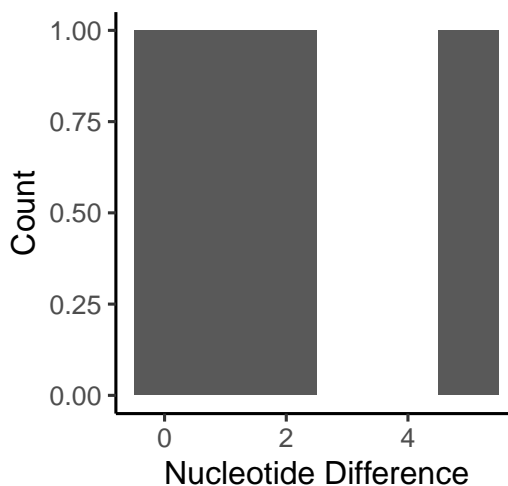
IGLV6-57*02

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



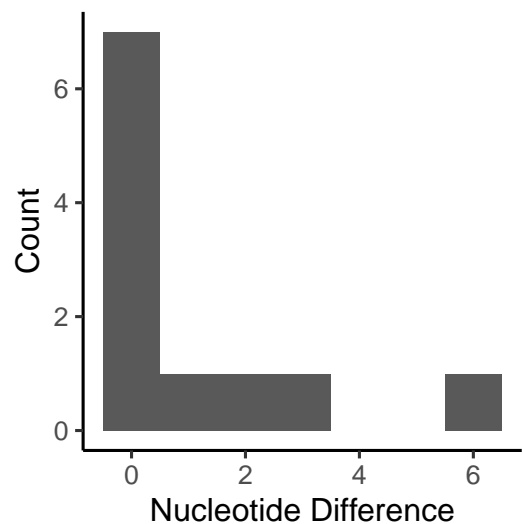
IGLV8-61*01

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



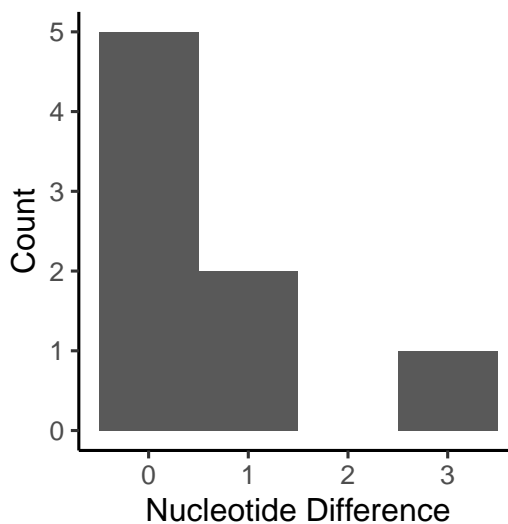
IGLV7-43*01

11 sequences assigned
7 (63.6%) exact matches, in which:
7 unique CDR3
2 unique J

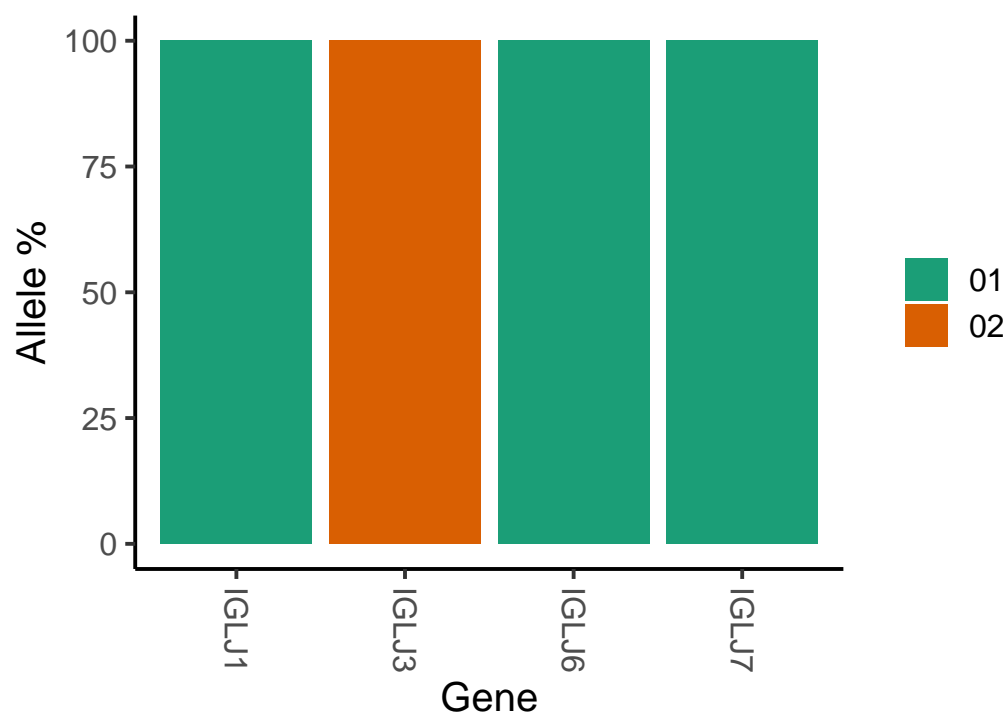


IGLV9-49*01

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



Allele Usage



Warning – no inferred sequences found.