

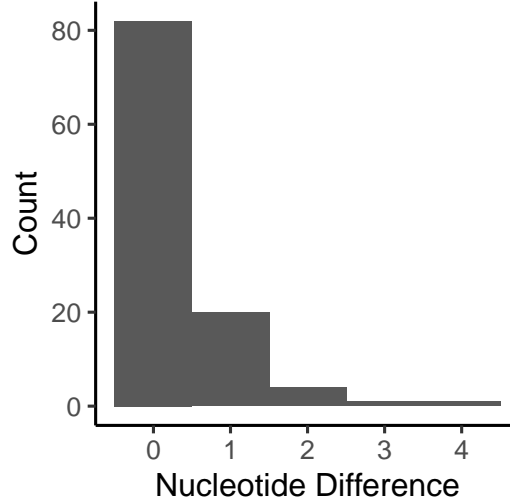
IGLV10-54*01

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



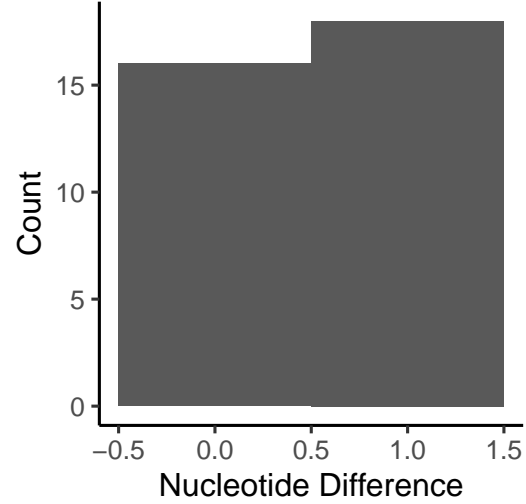
IGLV1-40*01

108 sequences assigned
82 (75.9%) exact matches, in which:
77 unique CDR3
3 unique J



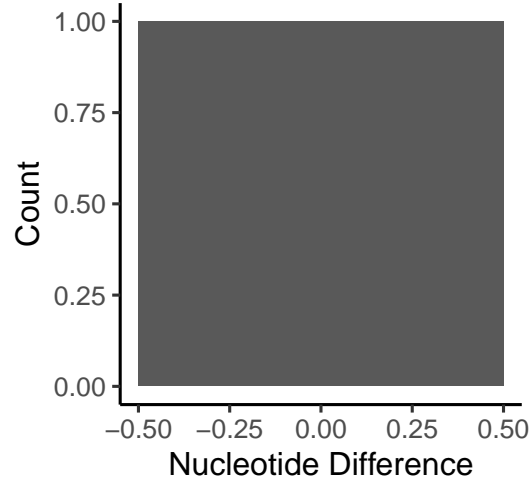
IGLV1-47*01

34 sequences assigned
16 (47.1%) exact matches, in which:
16 unique CDR3
2 unique J



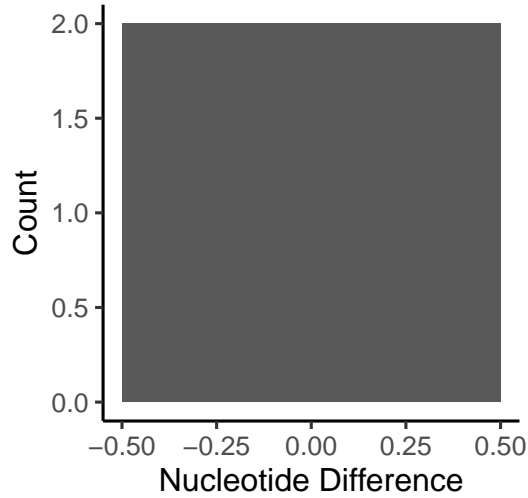
IGLV10-54*02

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



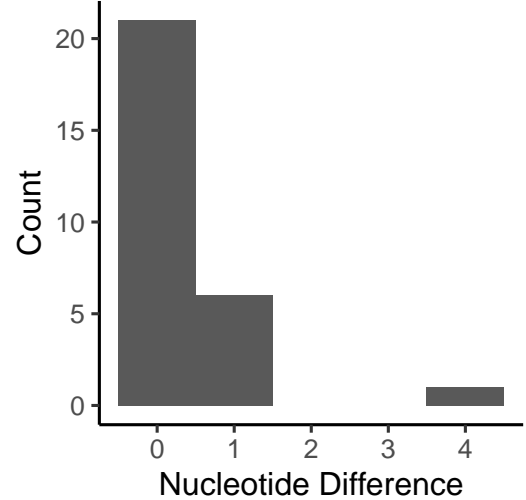
IGLV1-41*01

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



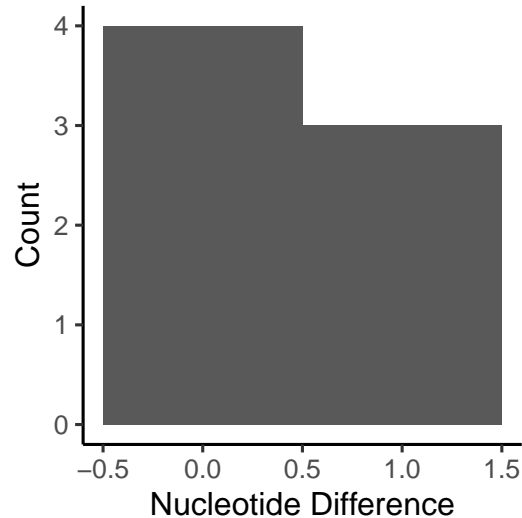
IGLV1-51*01

28 sequences assigned
21 (75%) exact matches, in which:
18 unique CDR3
3 unique J



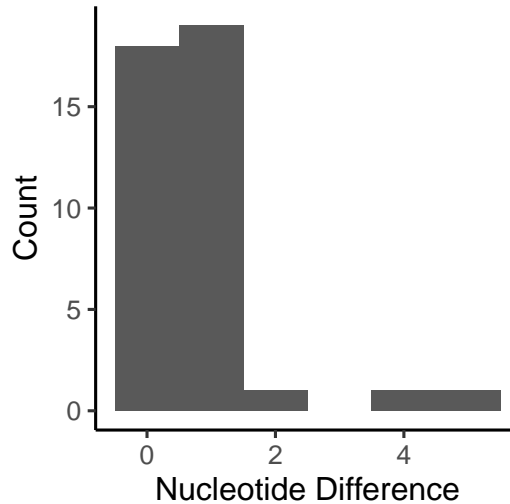
IGLV1-36*01

7 sequences assigned
4 (57.1%) exact matches, in which:
3 unique CDR3
2 unique J



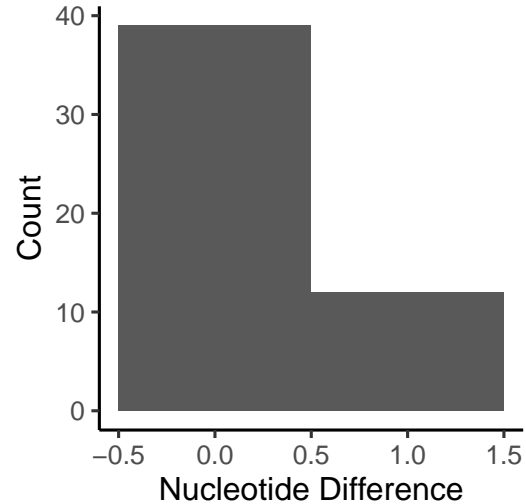
IGLV1-44*01

40 sequences assigned
18 (45%) exact matches, in which:
16 unique CDR3
2 unique J



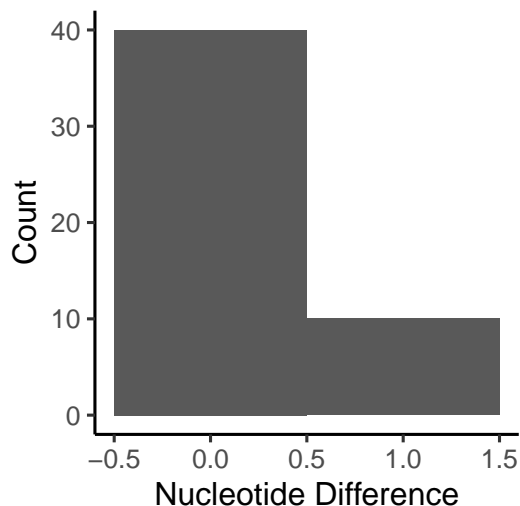
IGLV2-8*01

51 sequences assigned
39 (76.5%) exact matches, in which:
36 unique CDR3
2 unique J



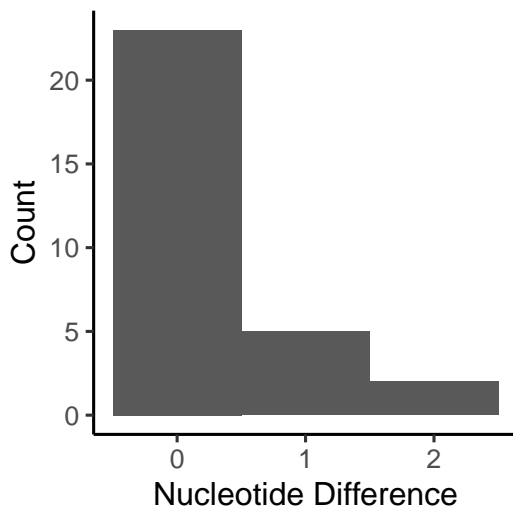
IGLV2-11*01

50 sequences assigned
40 (80%) exact matches, in which:
39 unique CDR3
2 unique J



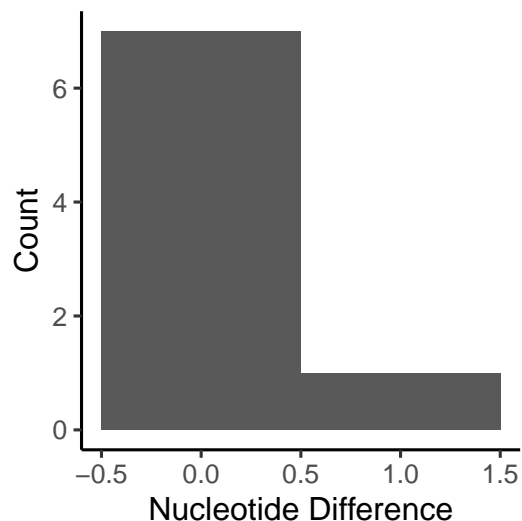
IGLV2-23*02EXT5

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



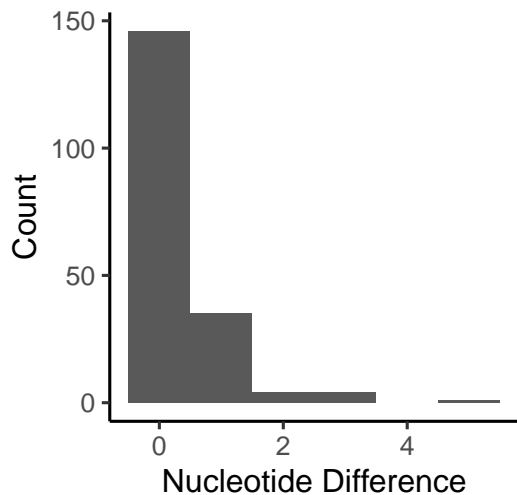
IGLV3-9*01EXT5

8 sequences assigned
7 (87.5%) exact matches, in which:
6 unique CDR3
2 unique J



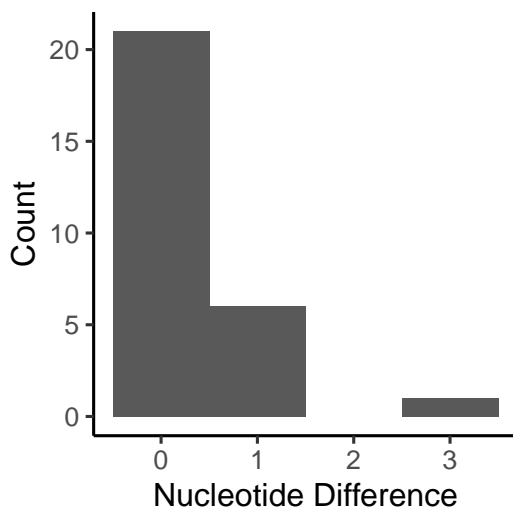
IGLV2-14*03EXT3

190 sequences assigned
146 (76.8%) exact matches, in which:
134 unique CDR3
2 unique J



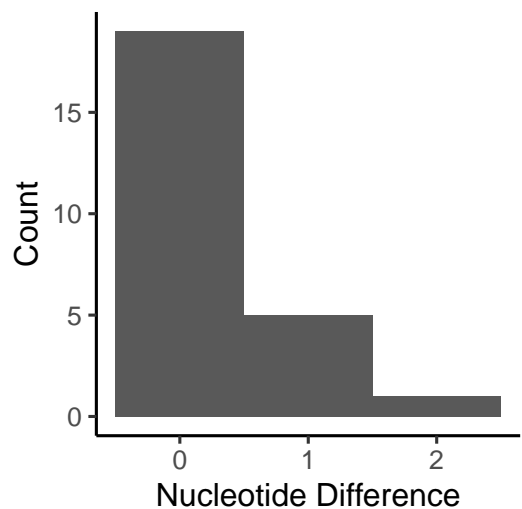
IGLV2-23*03EXT5

28 sequences assigned
21 (75%) exact matches, in which:
19 unique CDR3
2 unique J



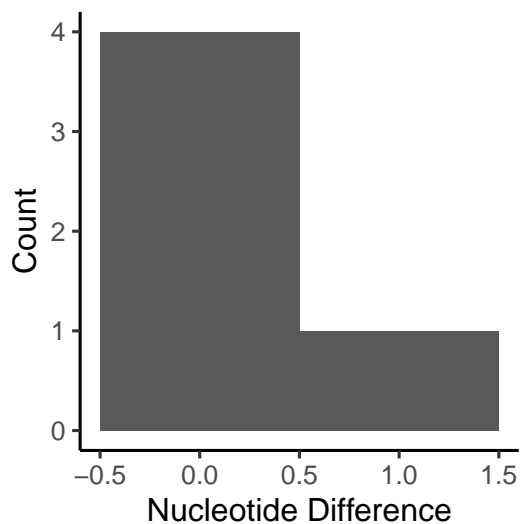
IGLV3-10*01

25 sequences assigned
19 (76%) exact matches, in which:
19 unique CDR3
3 unique J



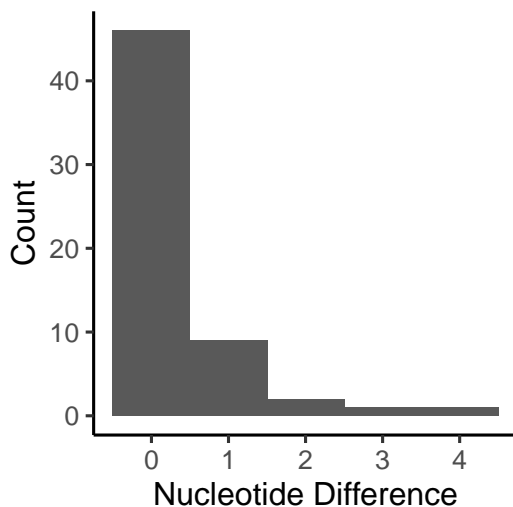
IGLV2-18*02

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



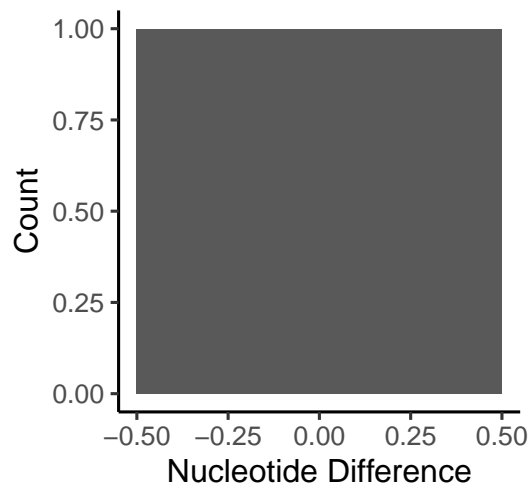
IGLV3-1*01

60 sequences assigned
46 (76.7%) exact matches, in which:
43 unique CDR3
2 unique J



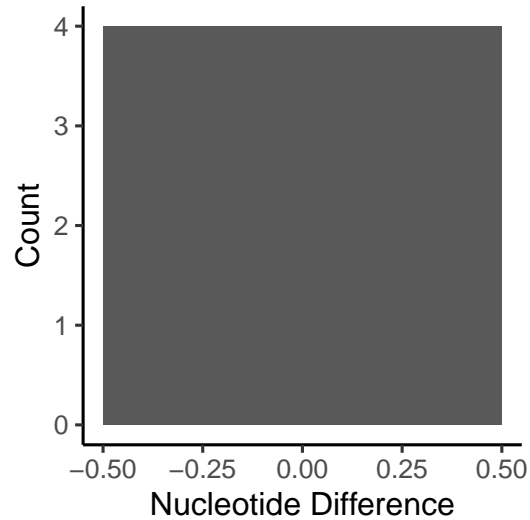
IGLV3-12*01

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



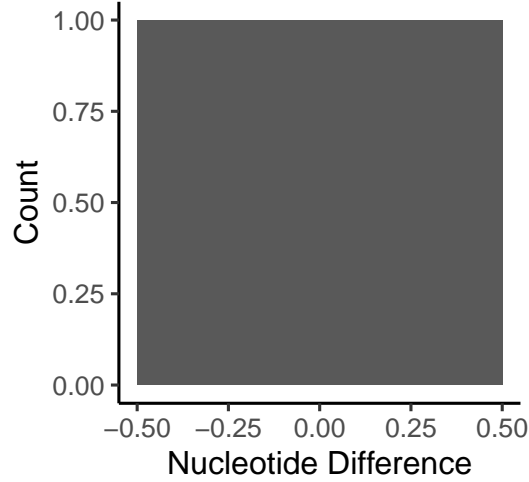
IGLV3–16*01

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



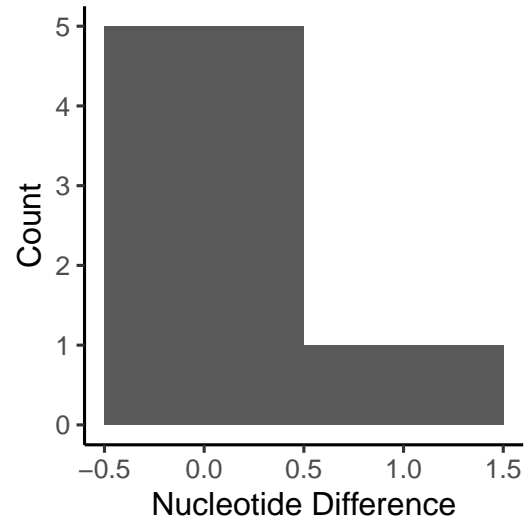
IGLV3–22*01

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



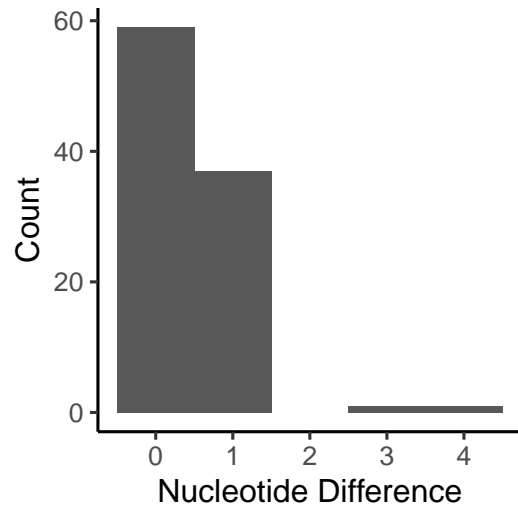
IGLV4–60*03EXT5

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



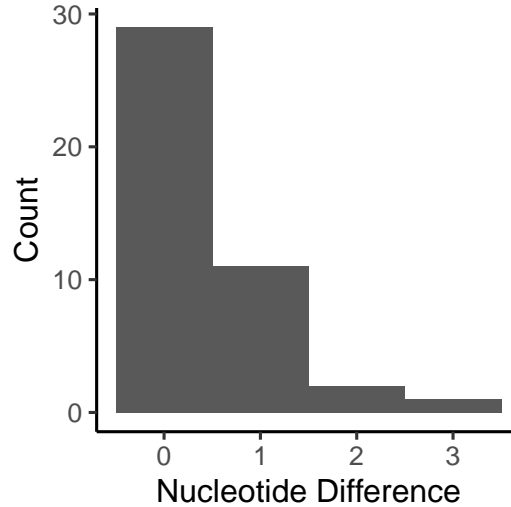
IGLV3–19*01

98 sequences assigned
59 (60.2%) exact matches, in which:
50 unique CDR3
3 unique J



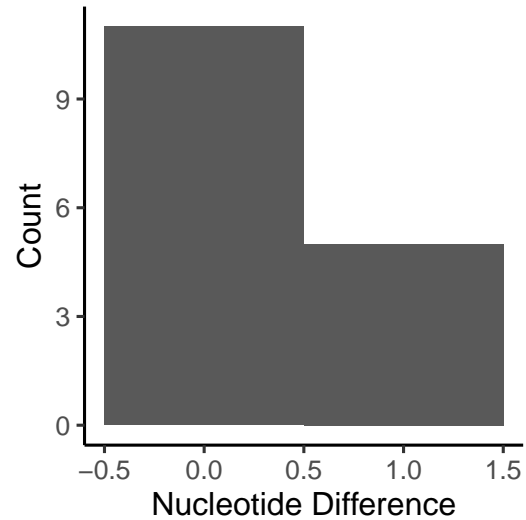
IGLV3–25*03

43 sequences assigned
29 (67.4%) exact matches, in which:
26 unique CDR3
3 unique J



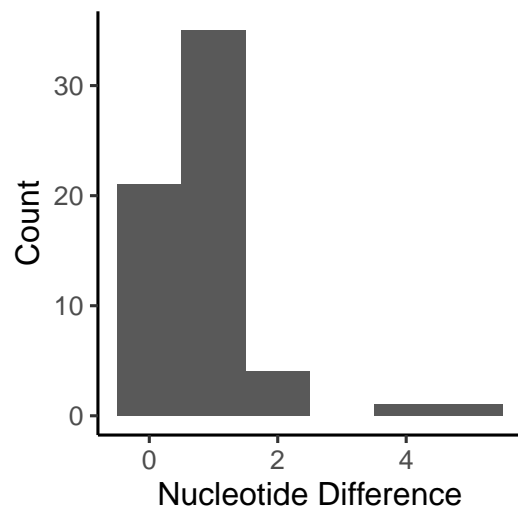
IGLV4–69*01

16 sequences assigned
11 (68.8%) exact matches, in which:
11 unique CDR3
3 unique J



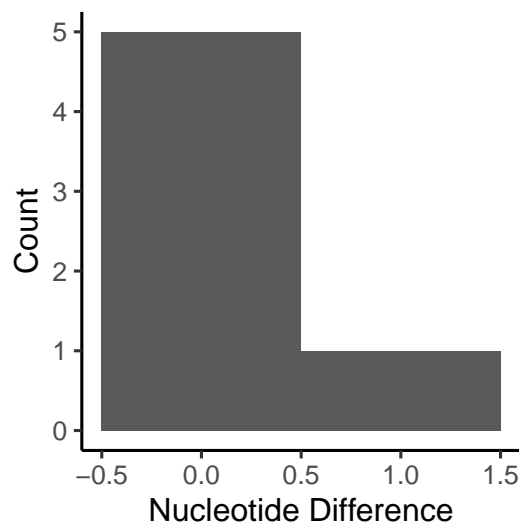
IGLV3–21*04

62 sequences assigned
21 (33.9%) exact matches, in which:
20 unique CDR3
2 unique J



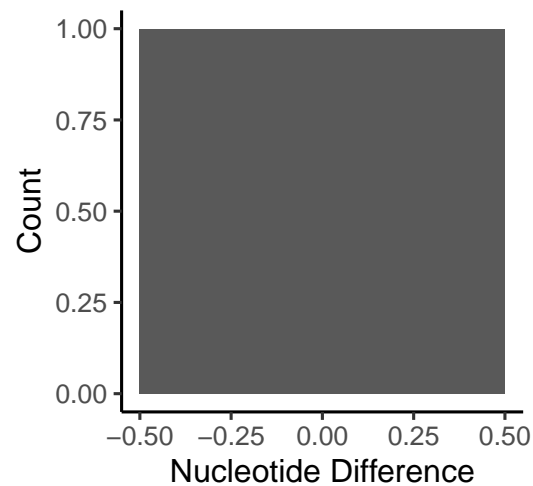
IGLV3–27*01

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



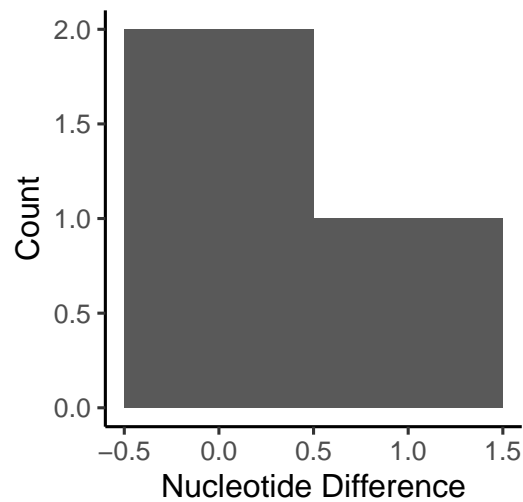
IGLV5–39*01

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



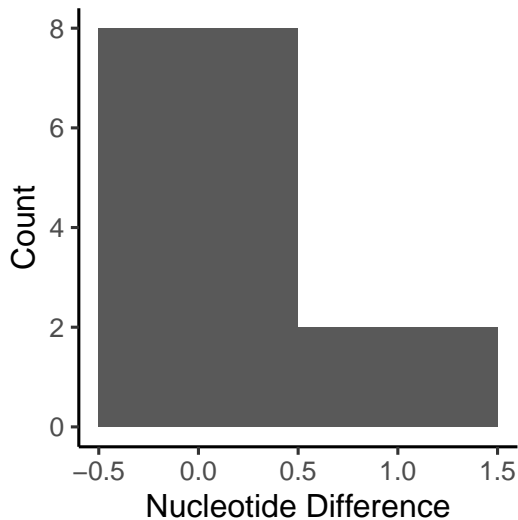
IGLV5-45*01

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



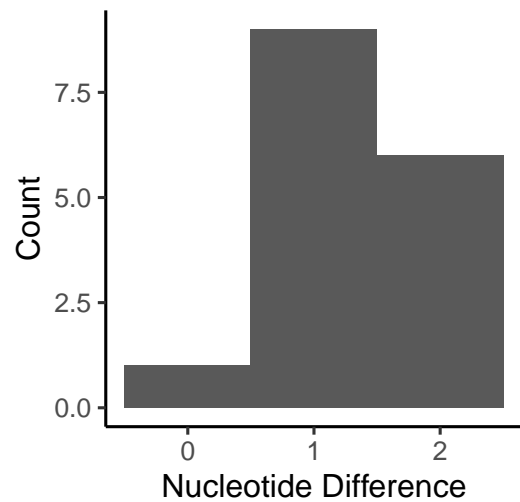
IGLV6-57*02

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



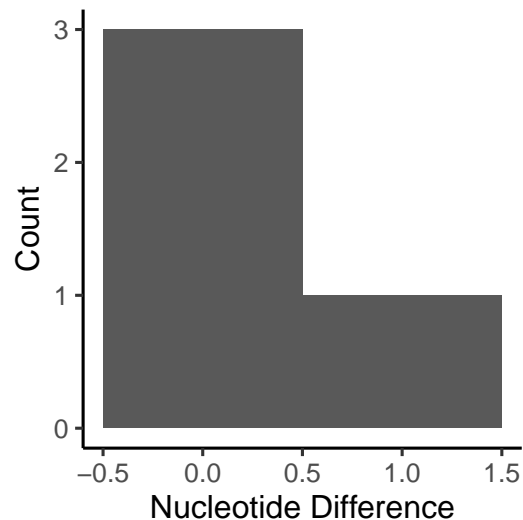
IGLV7-46*01

16 sequences assigned
1 (6.2%) exact matches, in which:
1 unique CDR3
1 unique J



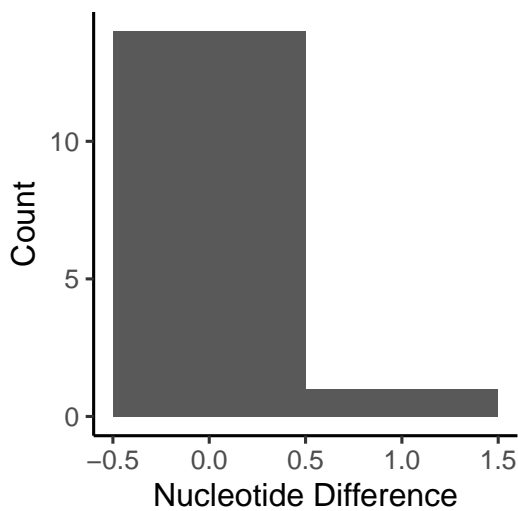
IGLV5-45*02

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



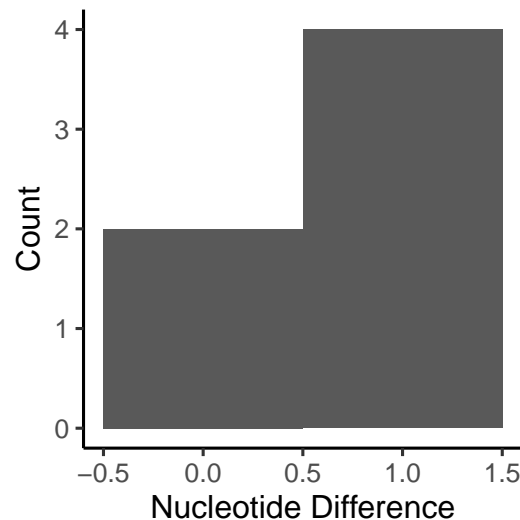
IGLV6-57*03

15 sequences assigned
14 (93.3%) exact matches, in which:
14 unique CDR3
2 unique J



IGLV8-61*01

6 sequences assigned
2 (33.3%) exact matches, in which:
2 unique CDR3
1 unique J



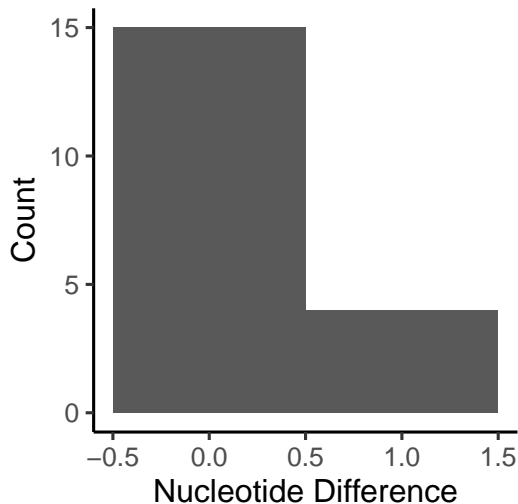
IGLV5-48*01

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



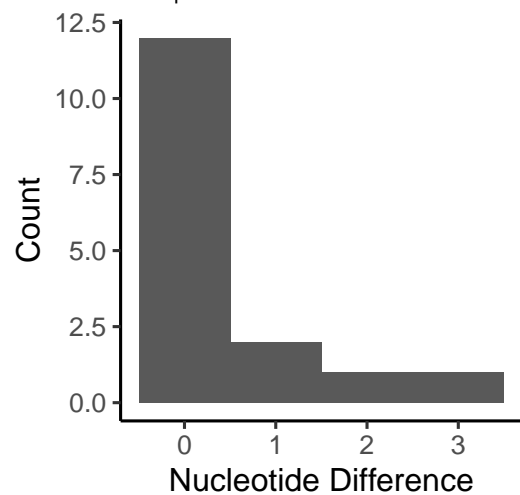
IGLV7-43*01

19 sequences assigned
15 (78.9%) exact matches, in which:
15 unique CDR3
2 unique J



IGLV9-49*01

16 sequences assigned
12 (75%) exact matches, in which:
11 unique CDR3
3 unique J



Allele Usage



Warning – no inferred sequences found.