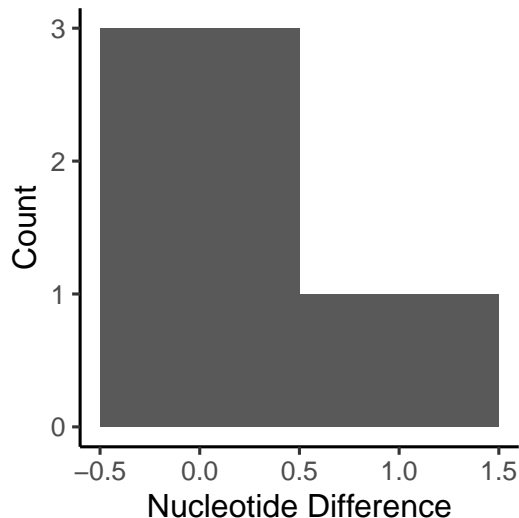


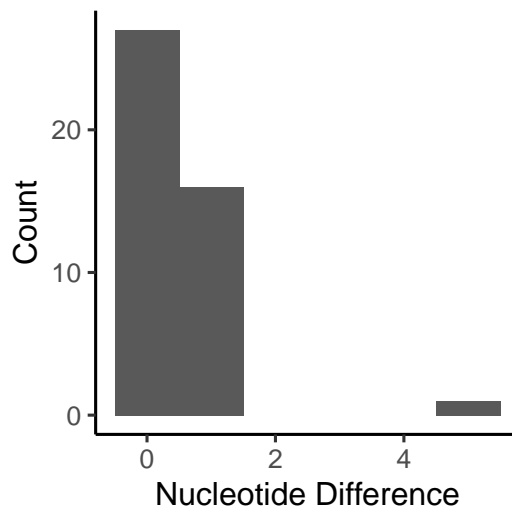
IGLV1-36*01

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



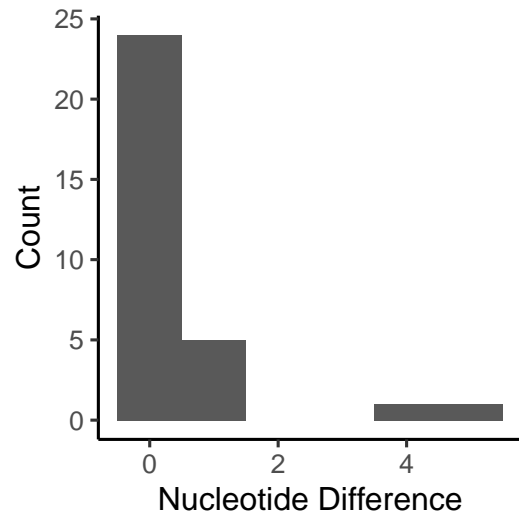
IGLV1-47*02

44 sequences assigned
27 (61.4%) exact matches, in which:
23 unique CDR3
2 unique J



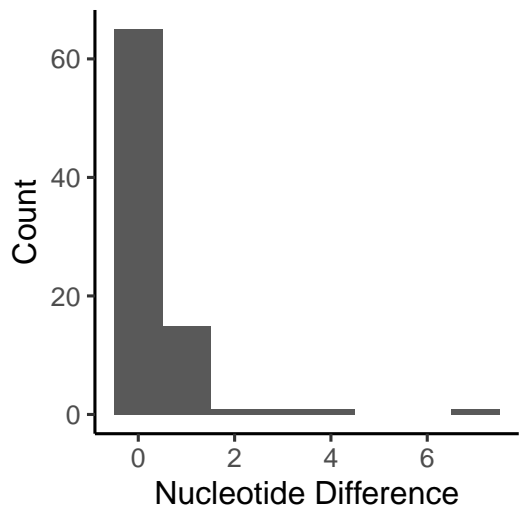
IGLV2-11*01

31 sequences assigned
24 (77.4%) exact matches, in which:
23 unique CDR3
2 unique J



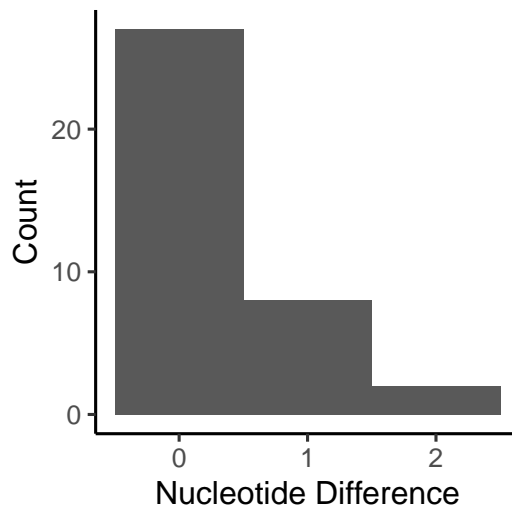
IGLV1-40*01

84 sequences assigned
65 (77.4%) exact matches, in which:
61 unique CDR3
3 unique J



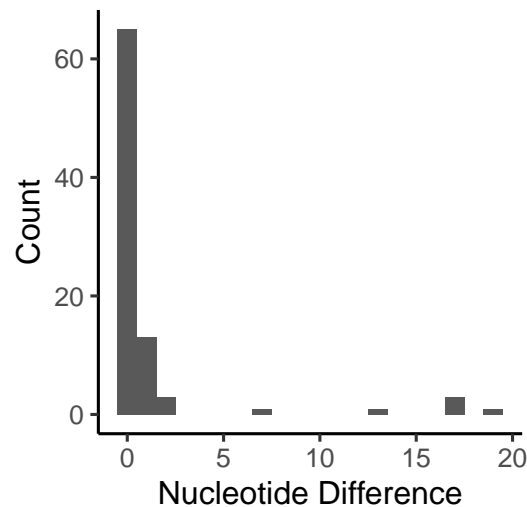
IGLV1-51*01

37 sequences assigned
27 (73%) exact matches, in which:
23 unique CDR3
2 unique J



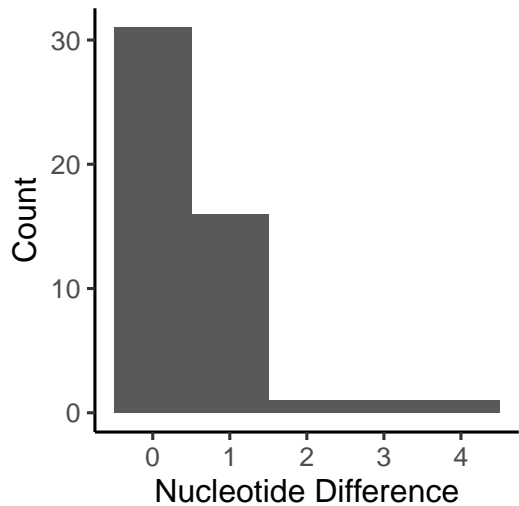
IGLV2-14*01

87 sequences assigned
65 (74.7%) exact matches, in which:
61 unique CDR3
2 unique J



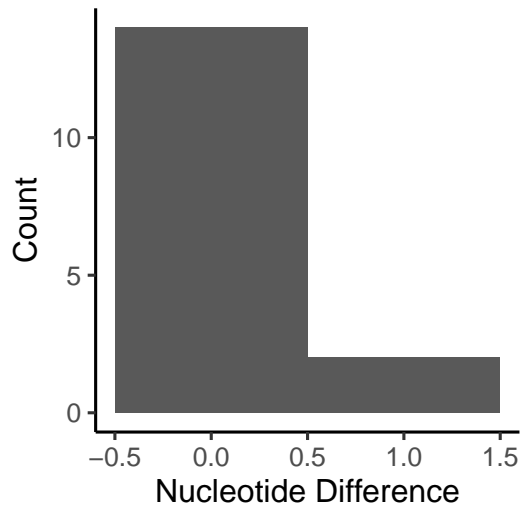
IGLV1-44*01

50 sequences assigned
31 (62%) exact matches, in which:
25 unique CDR3
2 unique J



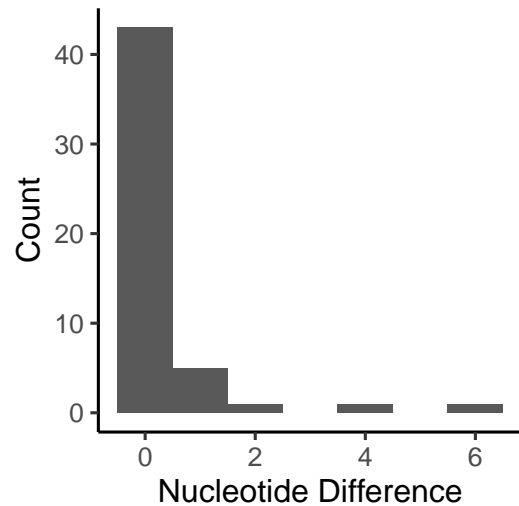
IGLV2-8*01

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



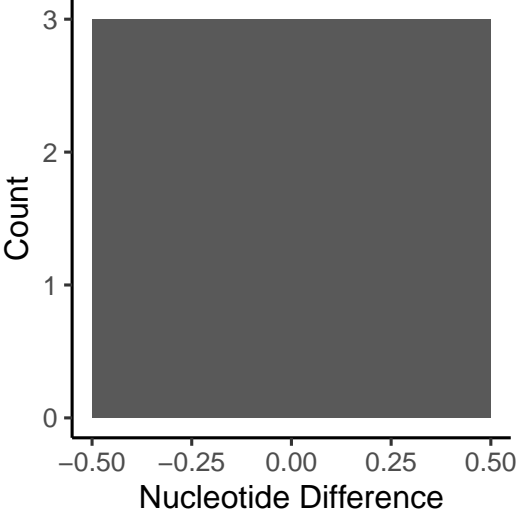
IGLV2-14*04EXT3

51 sequences assigned
43 (84.3%) exact matches, in which:
43 unique CDR3
2 unique J



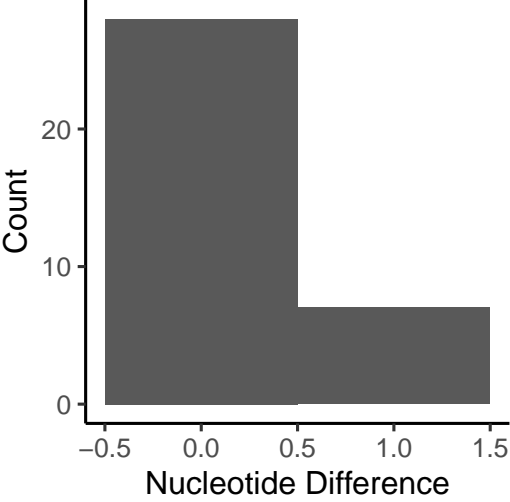
IGLV2-18*02

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



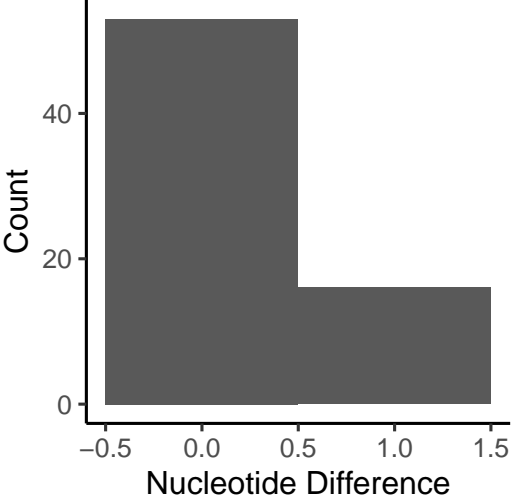
IGLV3-1*01

35 sequences assigned
28 (80%) exact matches, in which:
26 unique CDR3
2 unique J



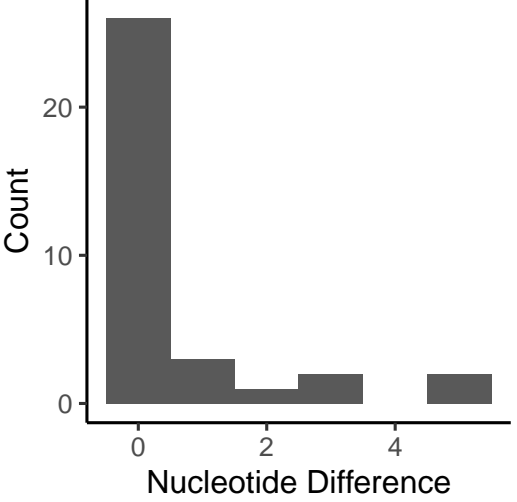
IGLV3-19*01

69 sequences assigned
53 (76.8%) exact matches, in which:
48 unique CDR3
3 unique J



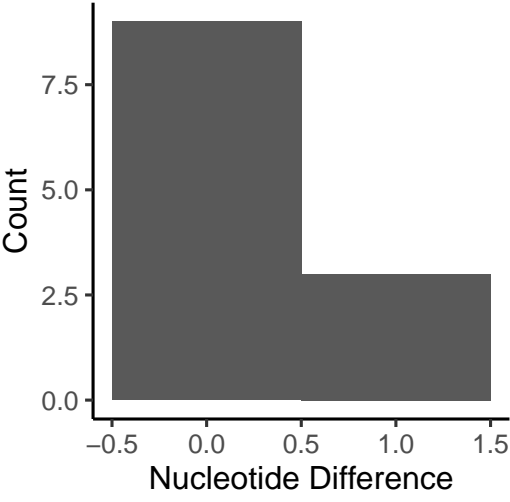
IGLV2-23*02EXT5

35 sequences assigned
26 (74.3%) exact matches, in which:
23 unique CDR3
4 unique J



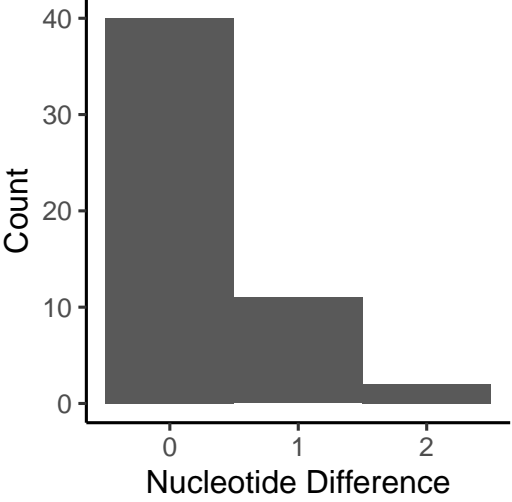
IGLV3-10*01

12 sequences assigned
9 (75%) exact matches, in which:
8 unique CDR3
2 unique J



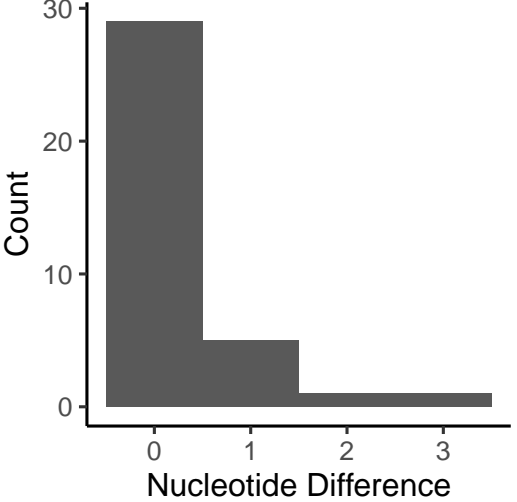
IGLV3-21*02

53 sequences assigned
40 (75.5%) exact matches, in which:
37 unique CDR3
2 unique J



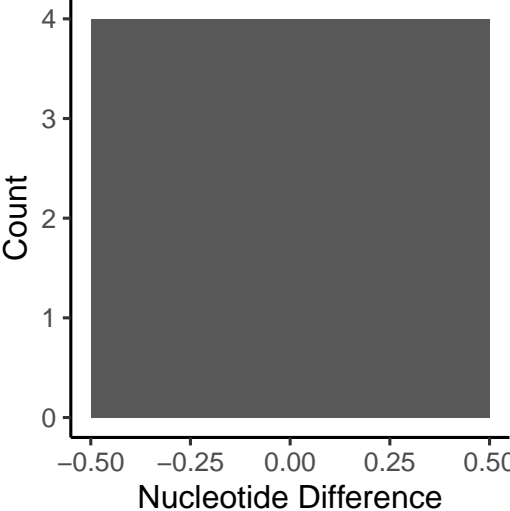
IGLV2-23*03EXT5

36 sequences assigned
29 (80.6%) exact matches, in which:
29 unique CDR3
3 unique J



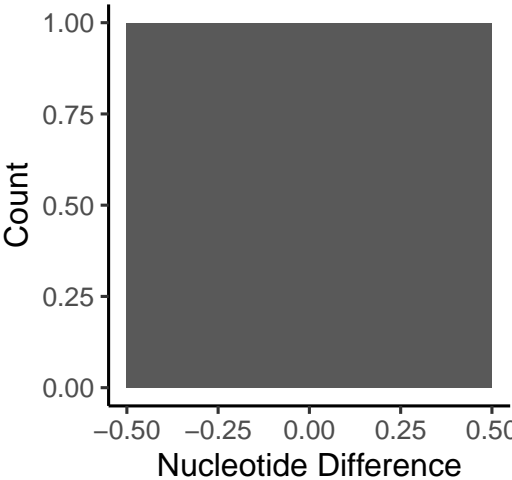
IGLV3-16*01

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



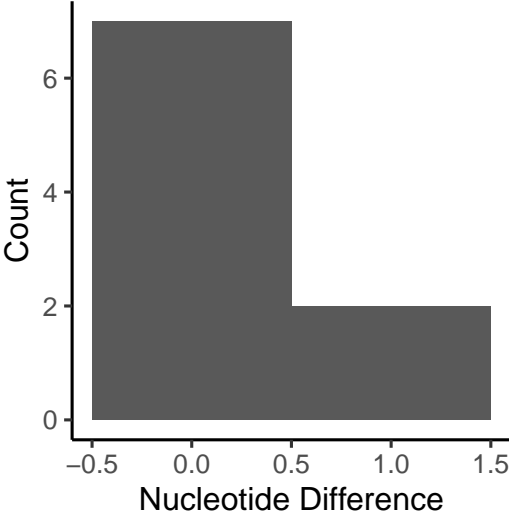
IGLV3-22*01

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



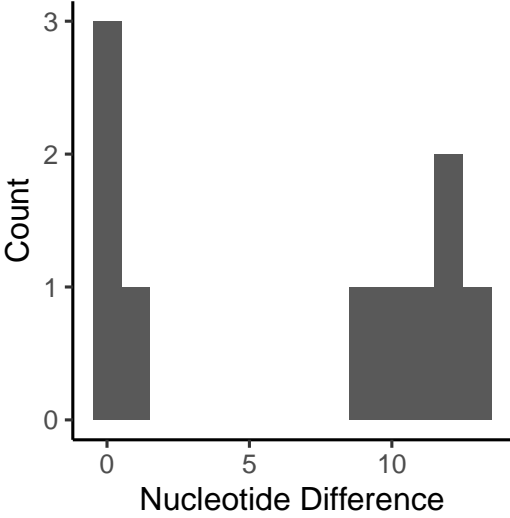
IGLV3–25*02

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



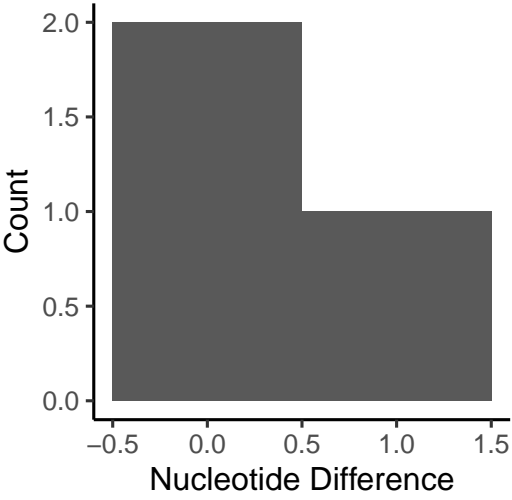
IGLV4–60*03EXT5

10 sequences assigned
3 (30%) exact matches, in which:
3 unique CDR3
2 unique J



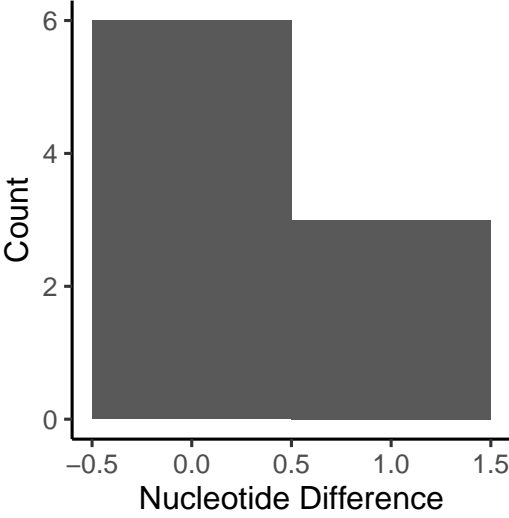
IGLV5–39*01

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



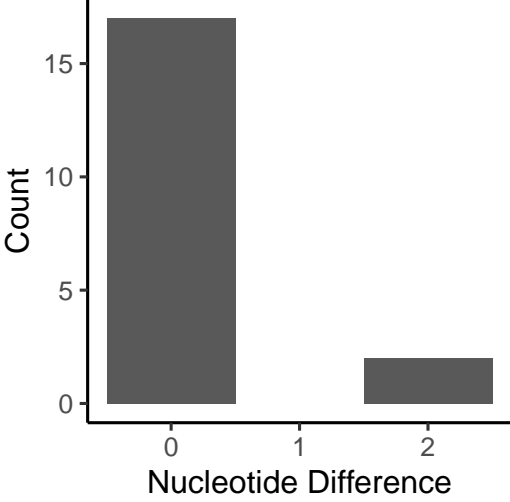
IGLV3–25*03

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



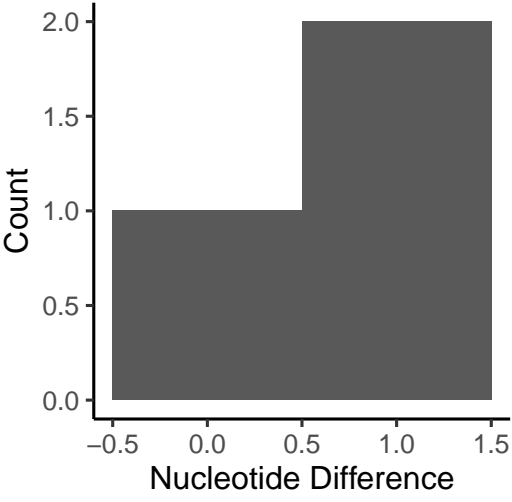
IGLV4–69*01

19 sequences assigned
17 (89.5%) exact matches, in which:
16 unique CDR3
2 unique J



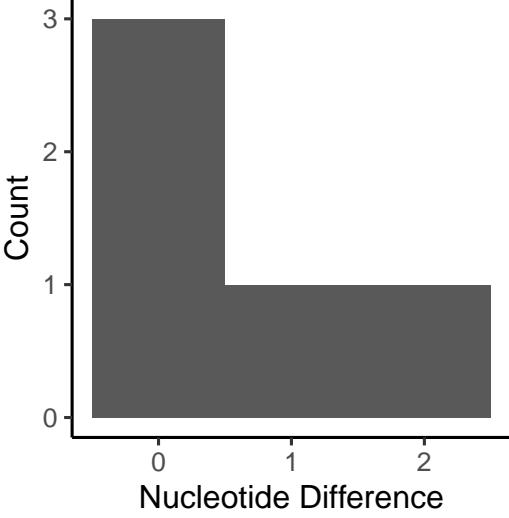
IGLV5–45*01

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



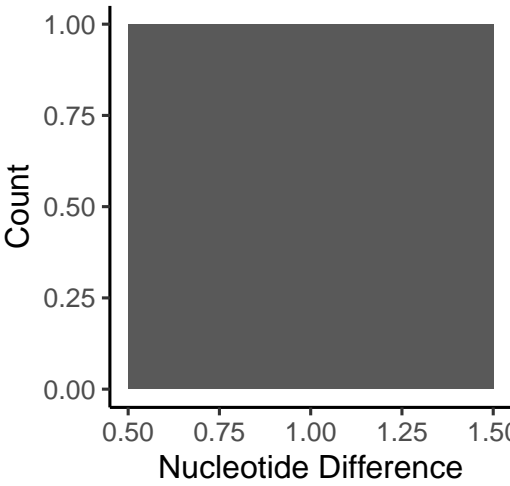
IGLV3–27*01

5 sequences assigned
3 (60%) exact matches, in which:
3 unique CDR3
1 unique J



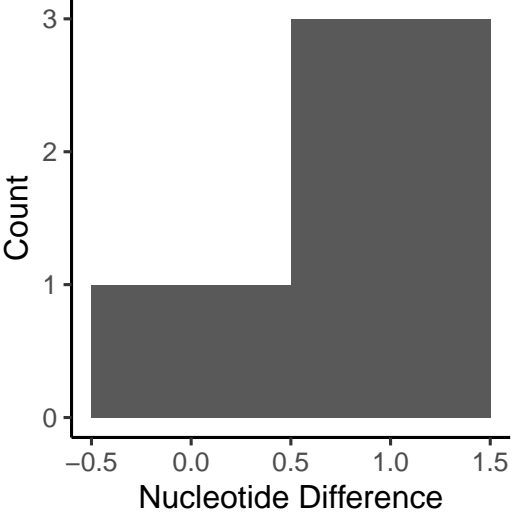
IGLV5–37*01

1 sequences assigned
No exact matches.



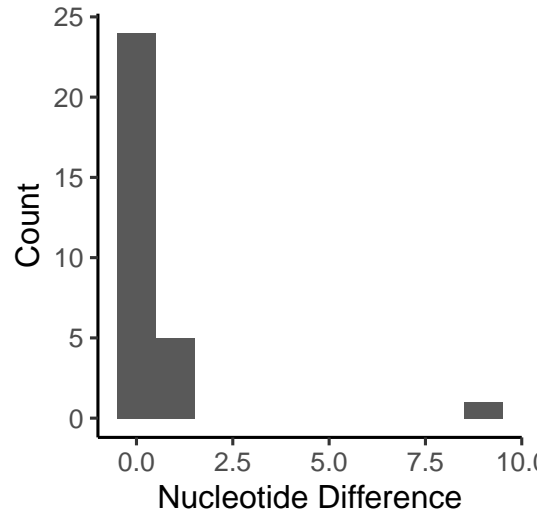
IGLV5–45*03

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



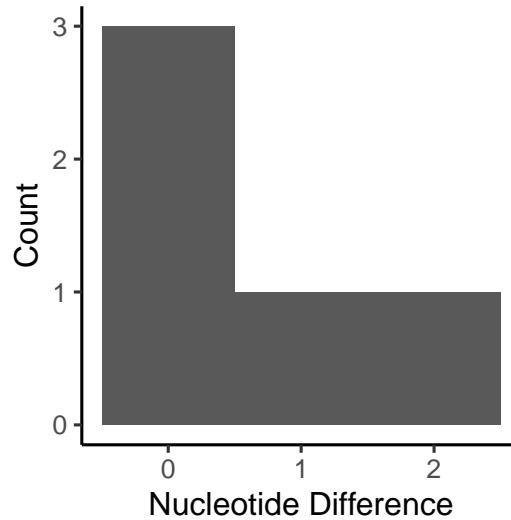
IGLV6-57*01

30 sequences assigned
24 (80%) exact matches, in which:
21 unique CDR3
2 unique J



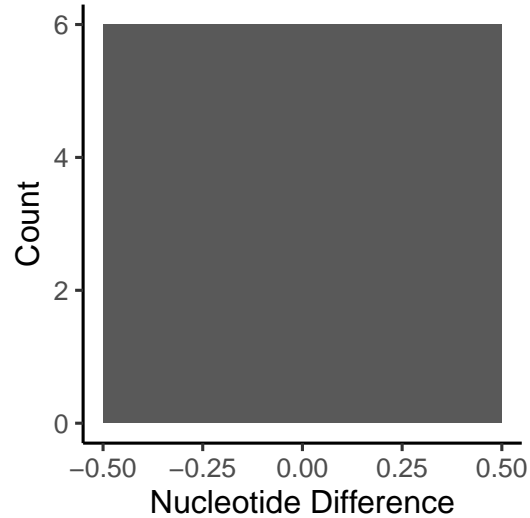
IGLV8-61*01

5 sequences assigned
3 (60%) exact matches, in which:
3 unique CDR3
2 unique J



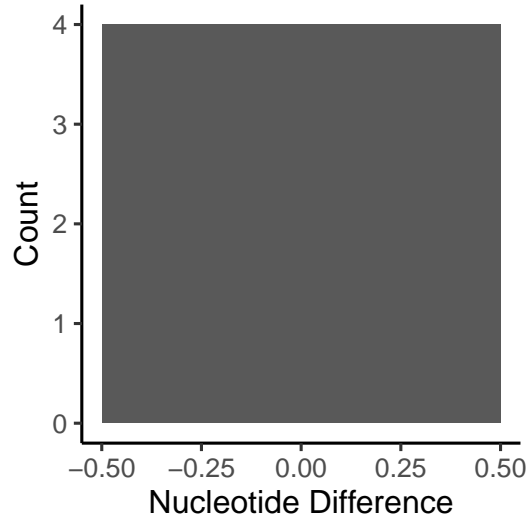
IGLV7-43*01

6 sequences assigned
6 (100%) exact matches, in which:
6 unique CDR3
2 unique J



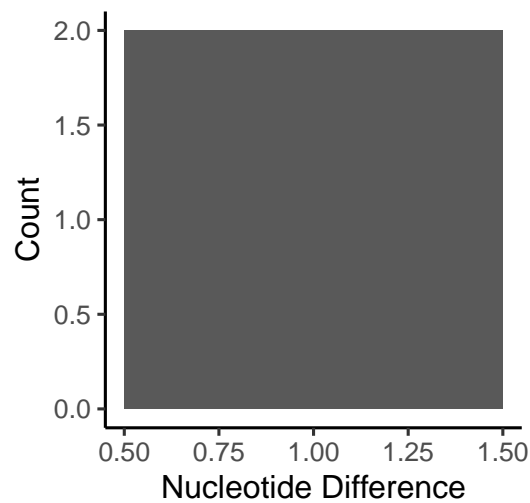
IGLV9-49*01

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

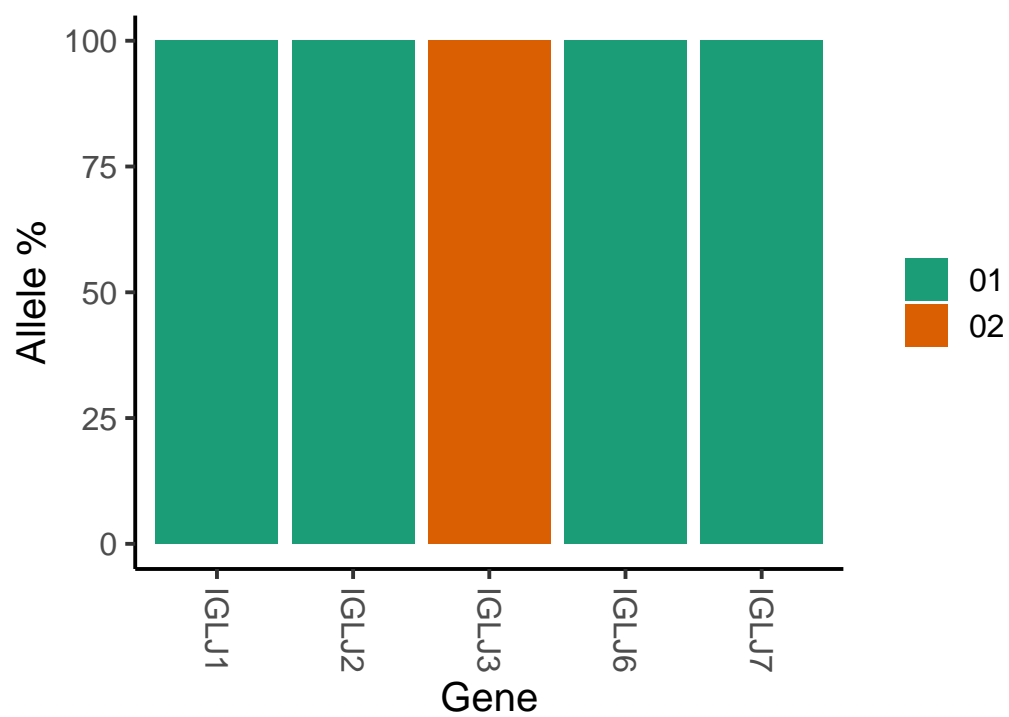


IGLV7-46*02

2 sequences assigned
No exact matches.



Allele Usage



Warning – no inferred sequences found.