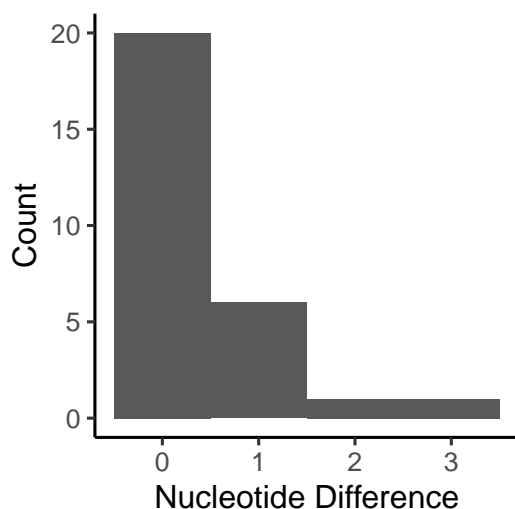


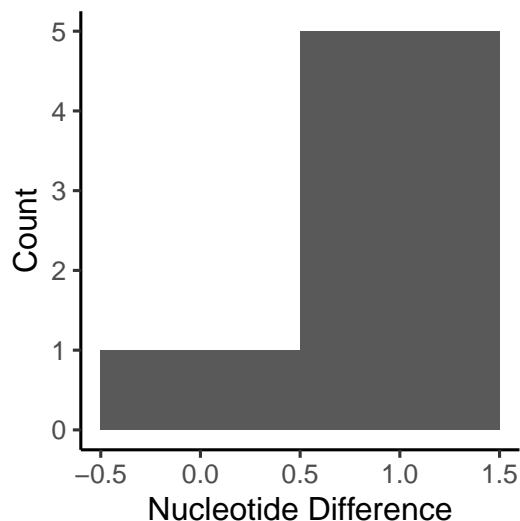
IGLV1-40*01

28 sequences assigned
20 (71.4%) exact matches, in which:
17 unique CDR3
3 unique J



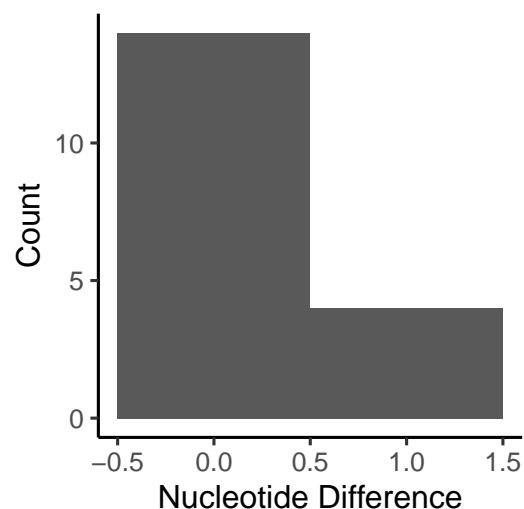
IGLV1-51*01

6 sequences assigned
1 (16.7%) exact matches, in which:
1 unique CDR3
1 unique J



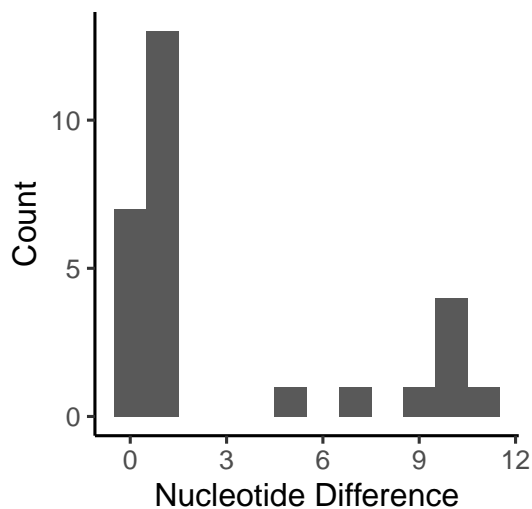
IGLV2-14*01

18 sequences assigned
14 (77.8%) exact matches, in which:
14 unique CDR3
2 unique J



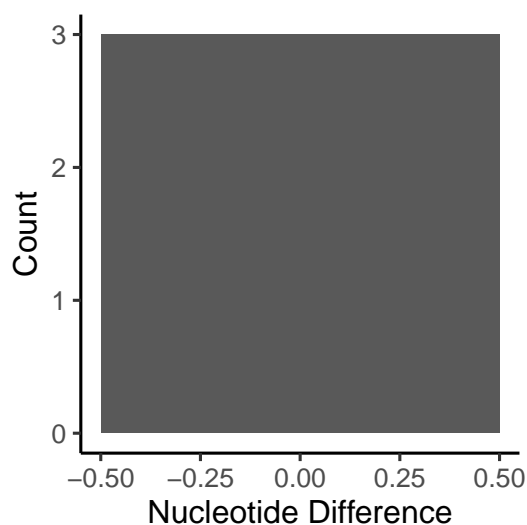
IGLV1-44*01

28 sequences assigned
7 (25%) exact matches, in which:
6 unique CDR3
2 unique J



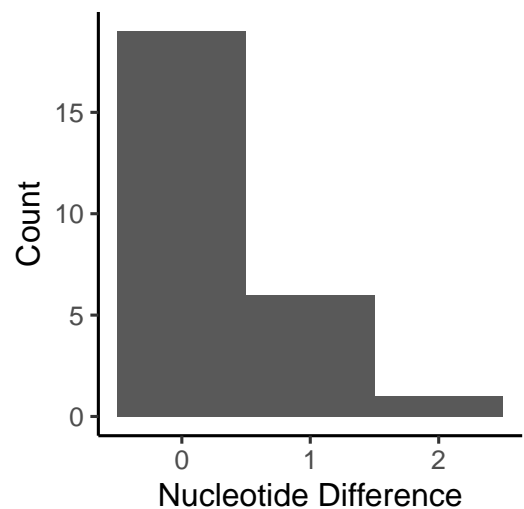
IGLV2-8*01

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



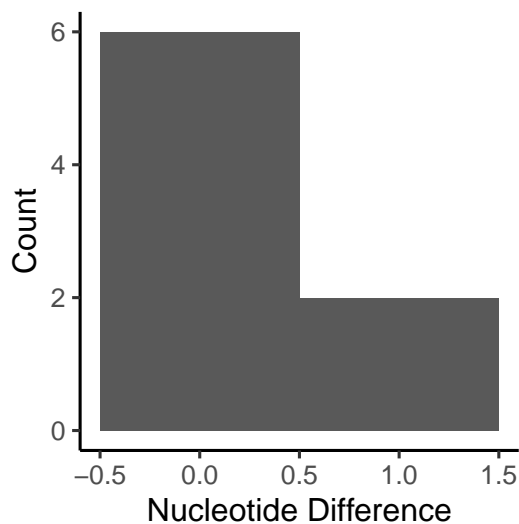
IGLV2-14*03EXT3

26 sequences assigned
19 (73.1%) exact matches, in which:
19 unique CDR3
3 unique J



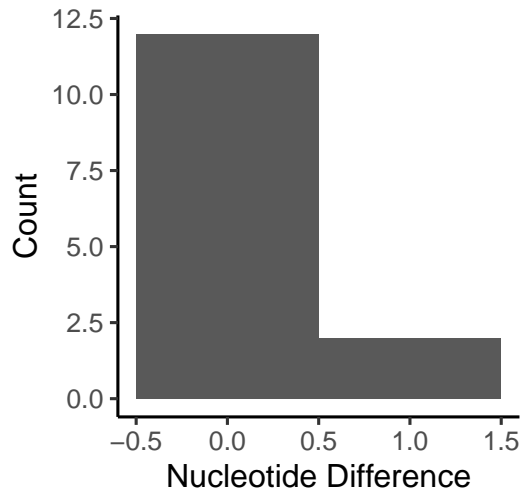
IGLV1-47*01

8 sequences assigned
6 (75%) exact matches, in which:
5 unique CDR3
2 unique J



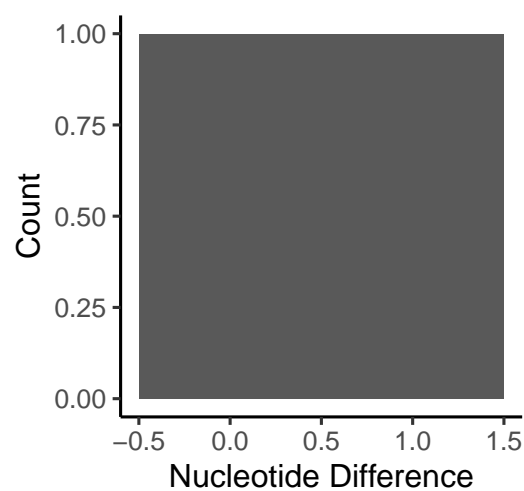
IGLV2-11*01

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



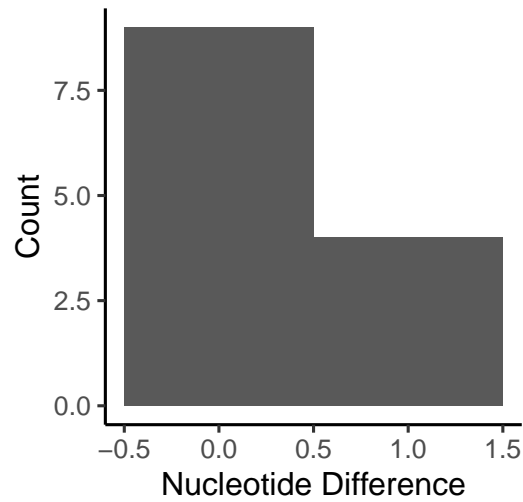
IGLV2-18*01

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



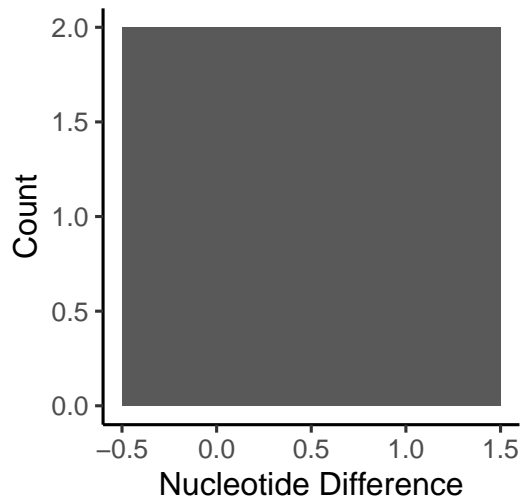
IGLV2–23*03EXT5

13 sequences assigned
9 (69.2%) exact matches, in which:
8 unique CDR3
2 unique J



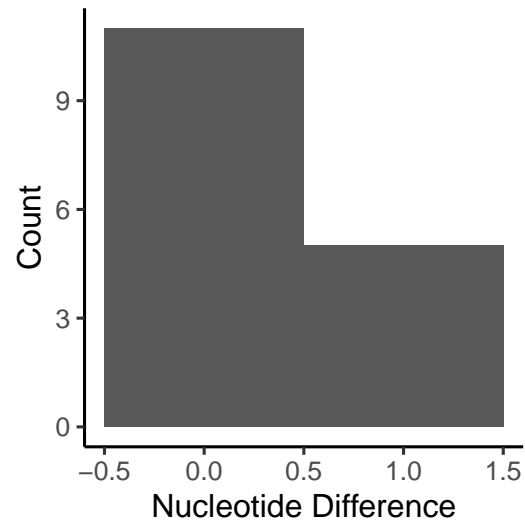
IGLV3–10*01

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



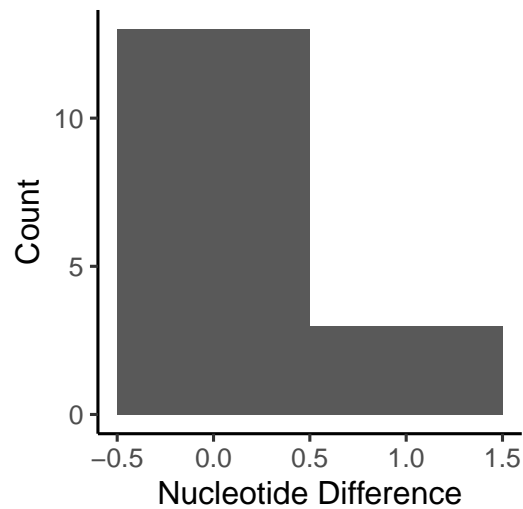
IGLV3–21*02

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



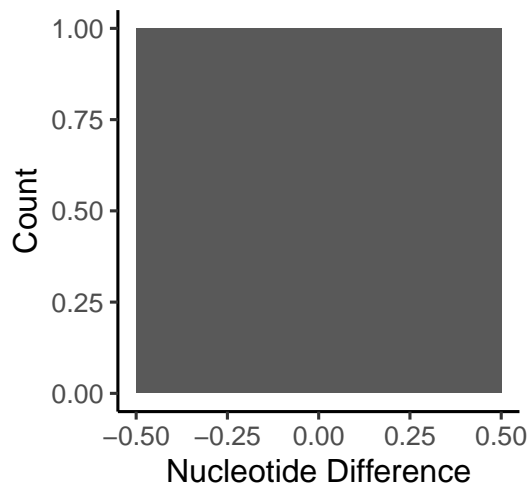
IGLV3–1*01

16 sequences assigned
13 (81.2%) exact matches, in which:
12 unique CDR3
3 unique J



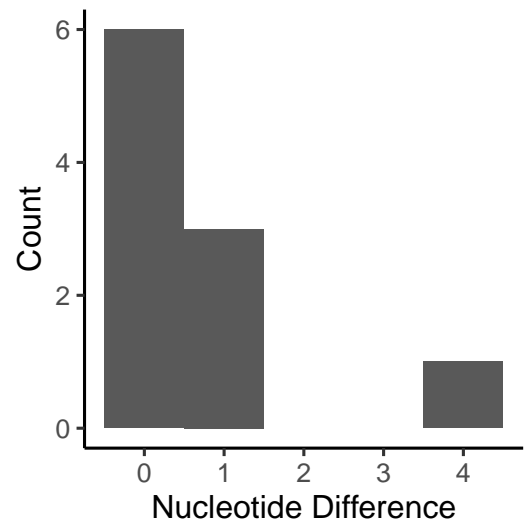
IGLV3–16*01

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



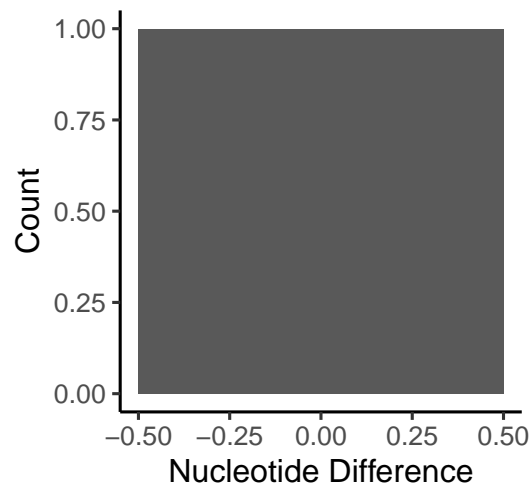
IGLV3–21*03

10 sequences assigned
6 (60%) exact matches, in which:
6 unique CDR3
3 unique J



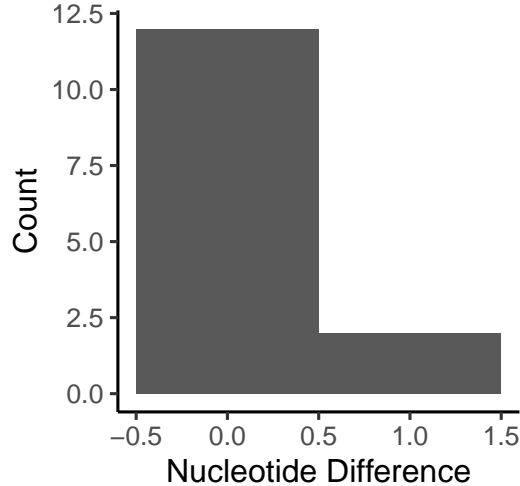
IGLV3–9*01EXT5

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



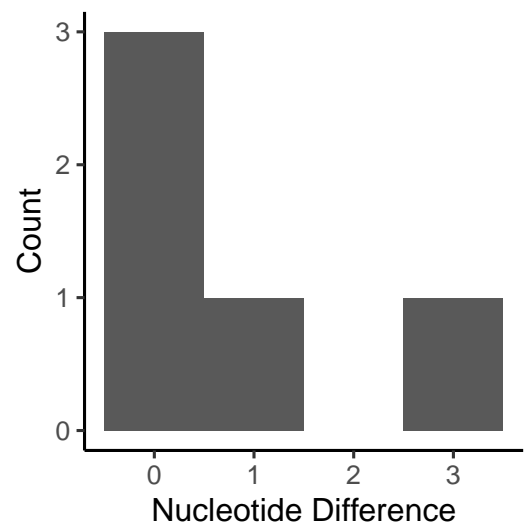
IGLV3–19*01

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



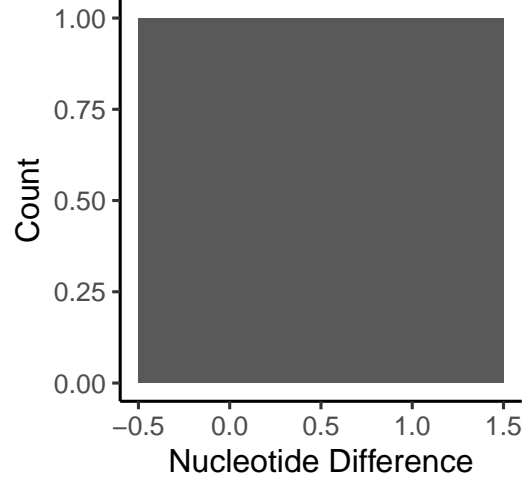
IGLV3–25*03

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



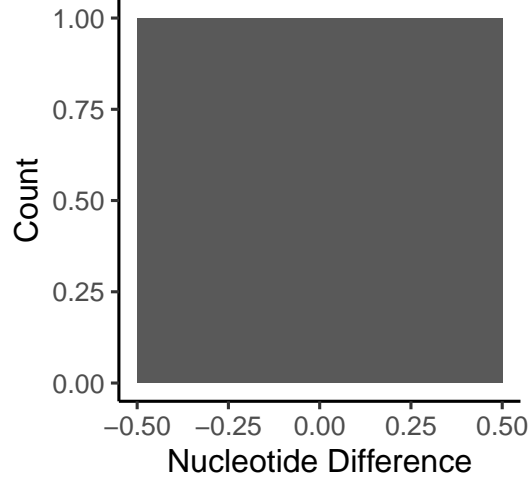
IGLV3–27*01

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



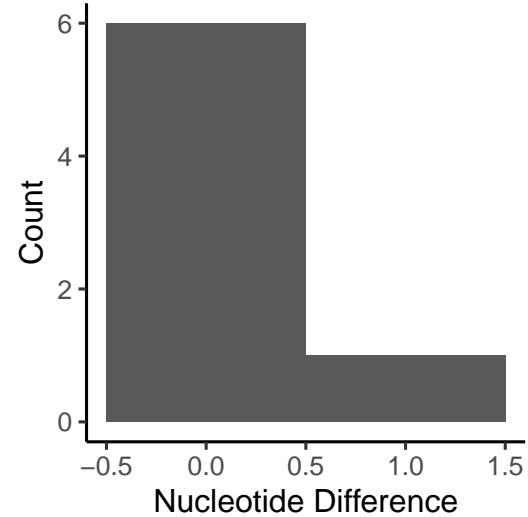
IGLV5–45*01

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



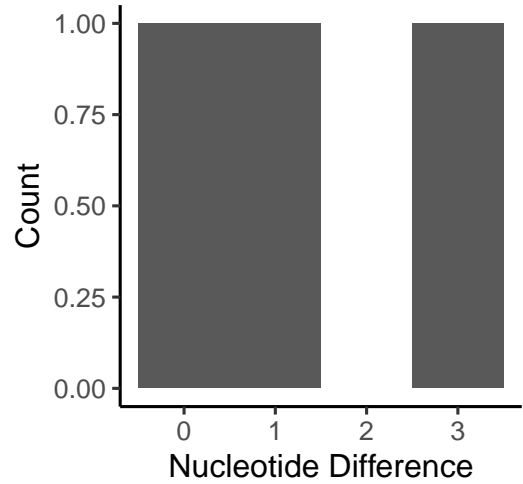
IGLV6–57*02

7 sequences assigned
6 (85.7%) exact matches, in which:
6 unique CDR3
2 unique J



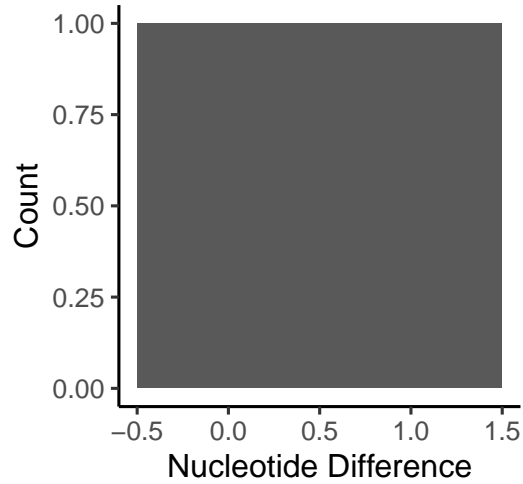
IGLV4–69*01

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



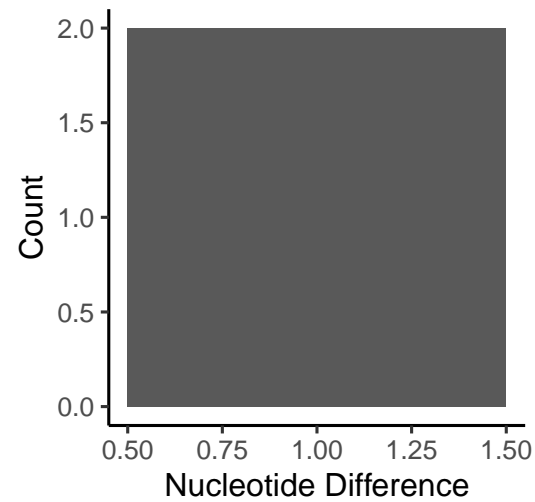
IGLV5–45*02

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



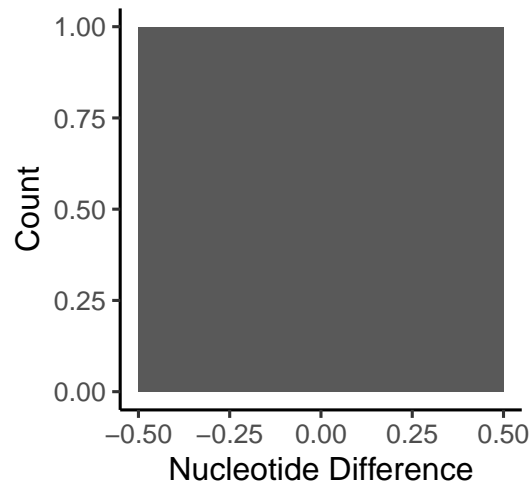
IGLV7–43*01

2 sequences assigned
No exact matches.



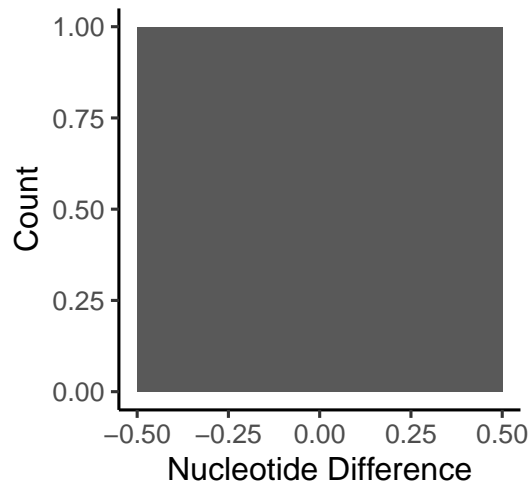
IGLV5–37*01

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



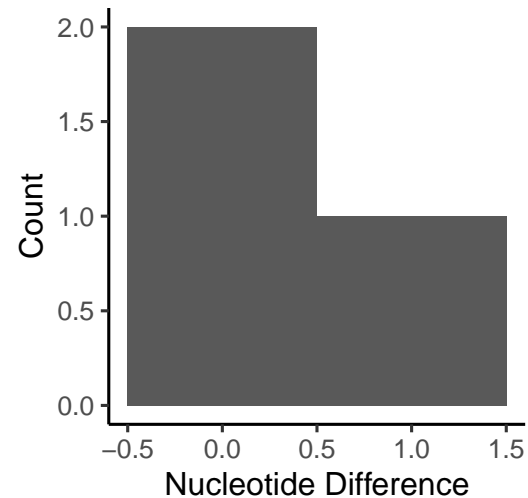
IGLV5–45*03

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



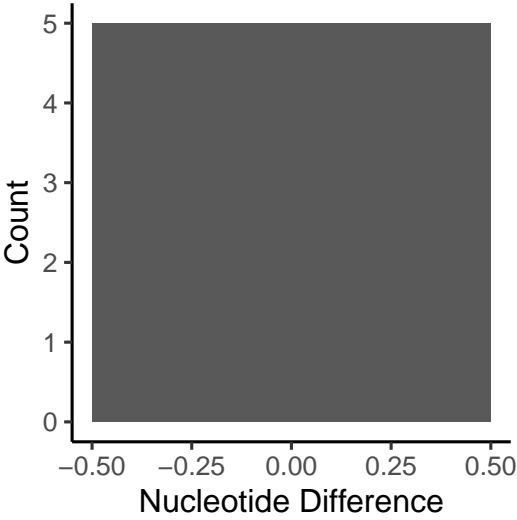
IGLV7–46*01

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



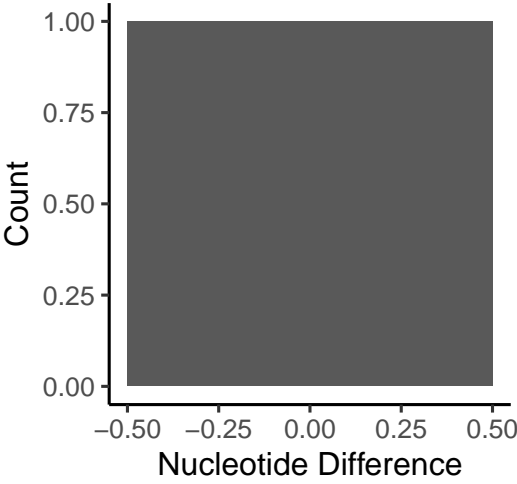
IGLV8-61*01

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

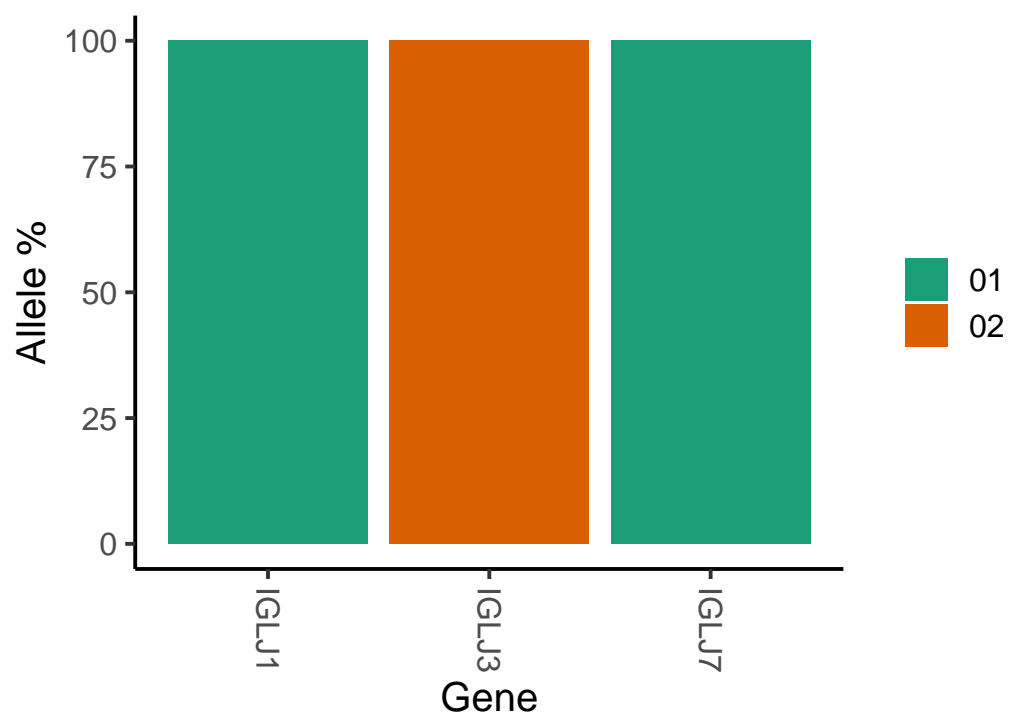


IGLV9-49*01

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



Allele Usage



Warning – no inferred sequences found.