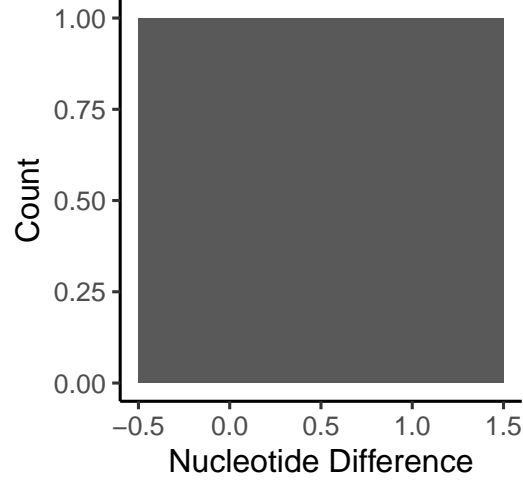


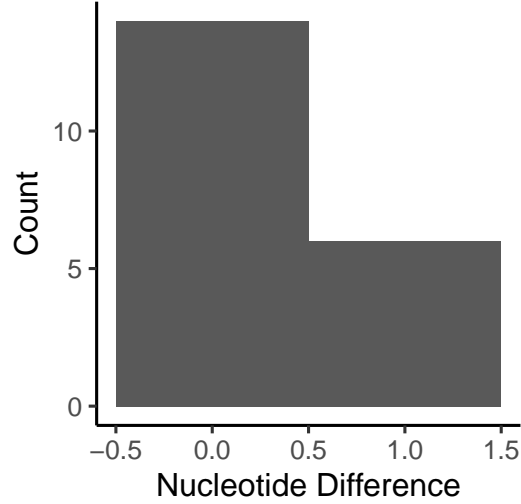
IGLV1-36*01

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



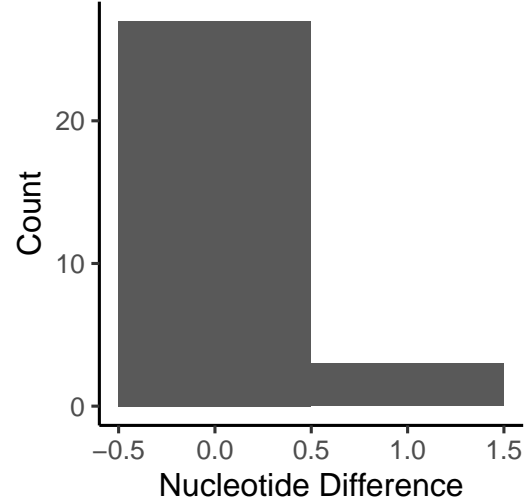
IGLV1-47*01

20 sequences assigned
14 (70%) exact matches, in which:
11 unique CDR3
3 unique J



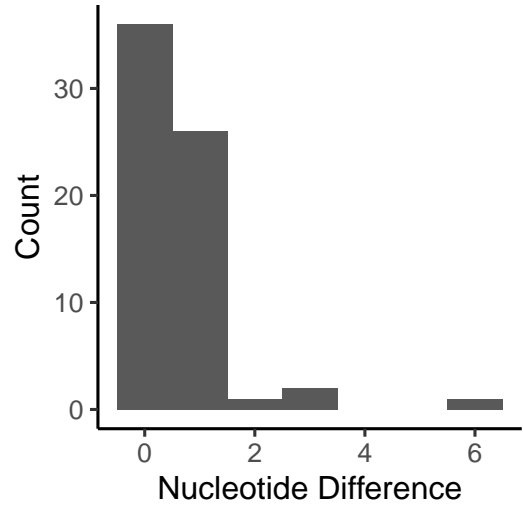
IGLV2-11*01

30 sequences assigned
27 (90%) exact matches, in which:
26 unique CDR3
3 unique J



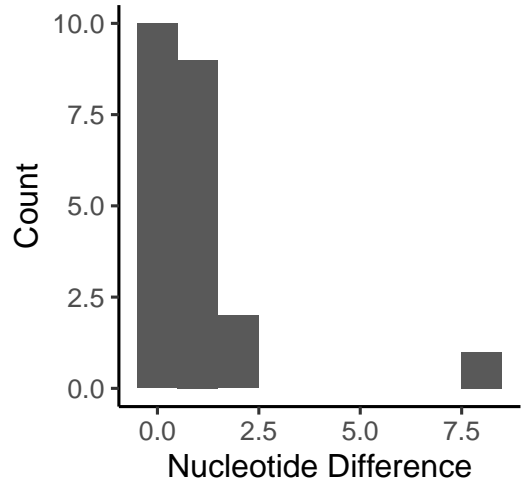
IGLV1-40*01

66 sequences assigned
36 (54.5%) exact matches, in which:
29 unique CDR3
4 unique J



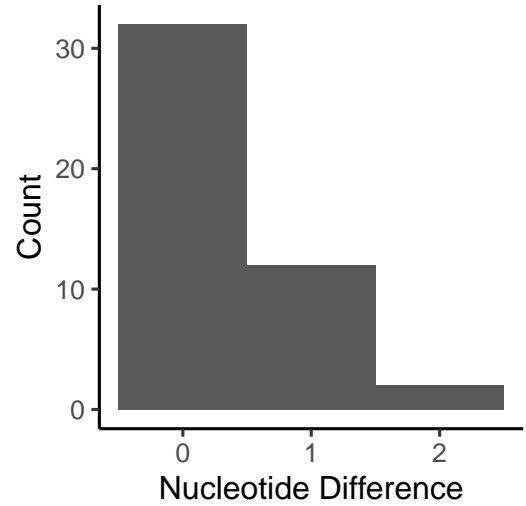
IGLV1-51*01

22 sequences assigned
10 (45.5%) exact matches, in which:
9 unique CDR3
2 unique J



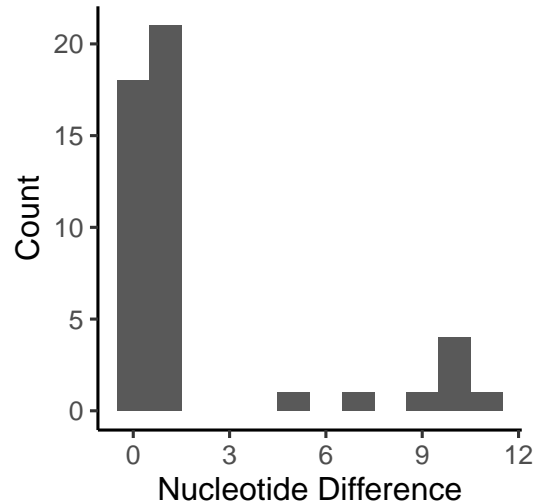
IGLV2-14*01

46 sequences assigned
32 (69.6%) exact matches, in which:
30 unique CDR3
3 unique J



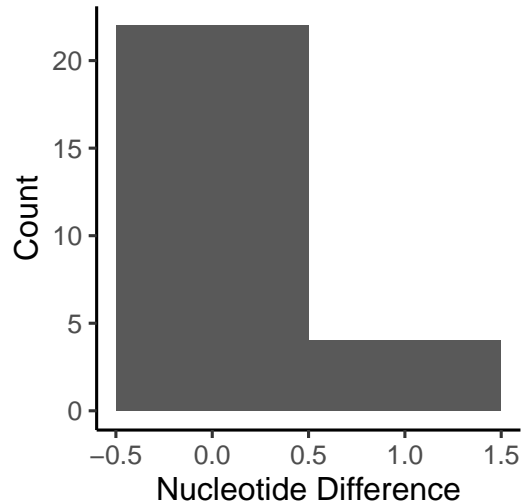
IGLV1-44*01

47 sequences assigned
18 (38.3%) exact matches, in which:
15 unique CDR3
3 unique J



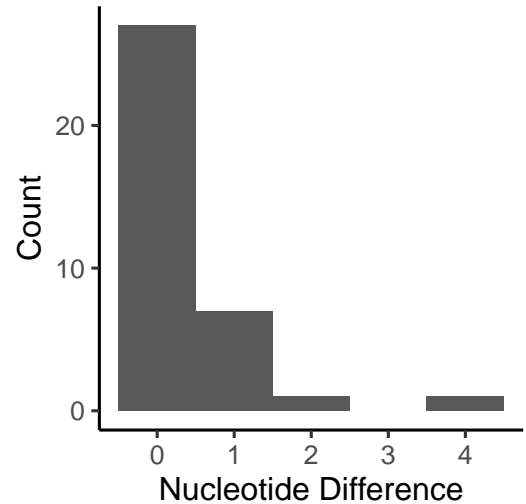
IGLV2-8*01

26 sequences assigned
22 (84.6%) exact matches, in which:
14 unique CDR3
3 unique J



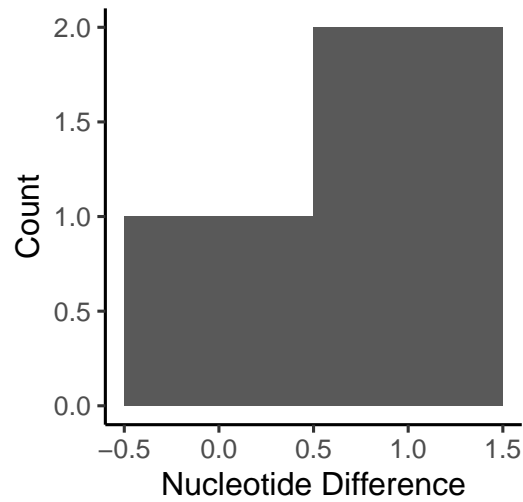
IGLV2-14*03EXT3

36 sequences assigned
27 (75%) exact matches, in which:
26 unique CDR3
4 unique J



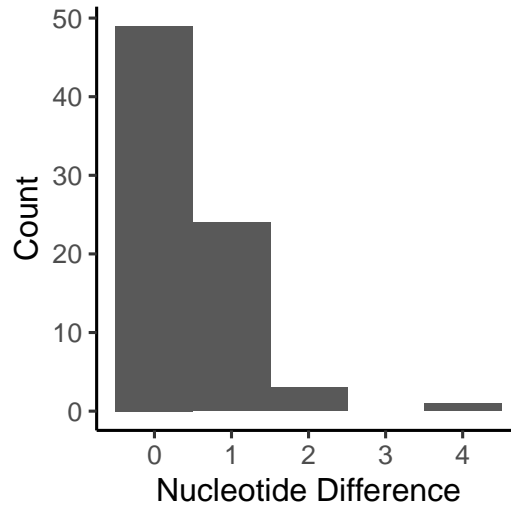
IGLV2-18*01

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



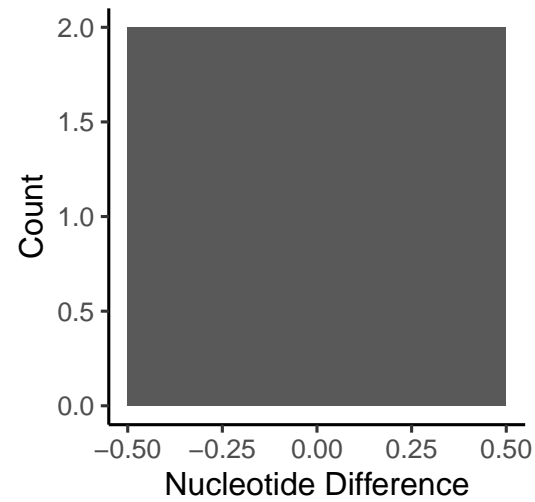
IGLV3-1*01

77 sequences assigned
49 (63.6%) exact matches, in which:
39 unique CDR3
4 unique J



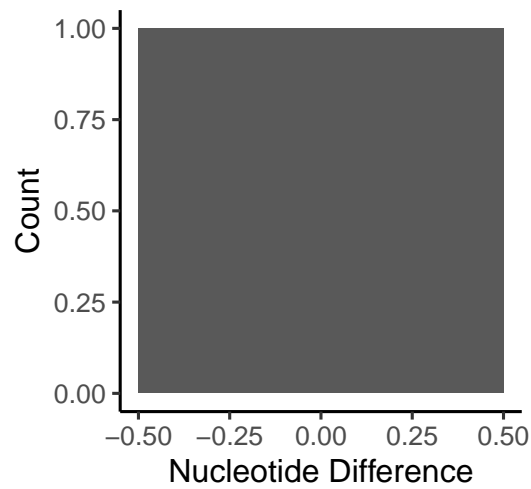
IGLV3-16*01

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



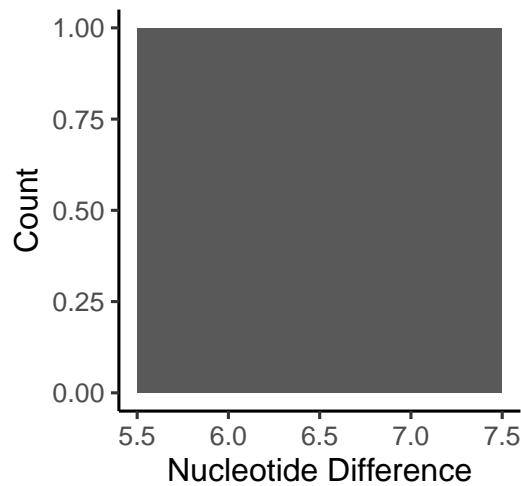
IGLV2-18*02

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



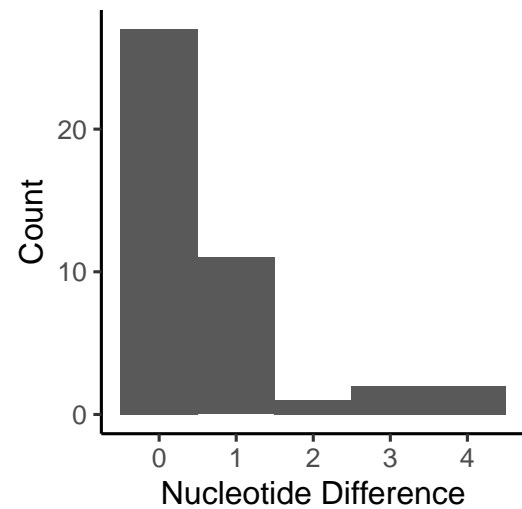
IGLV3-9*02

2 sequences assigned
No exact matches.



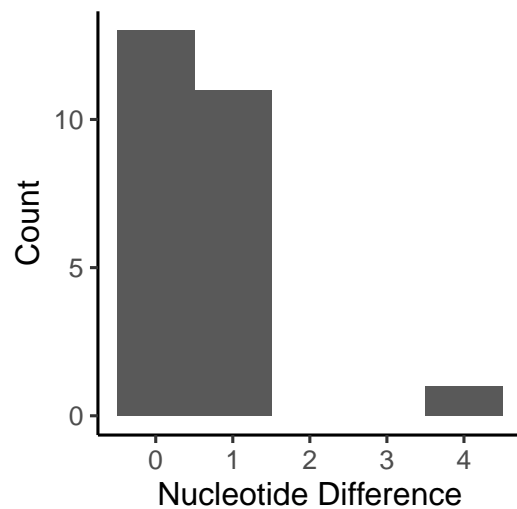
IGLV3-19*01

43 sequences assigned
27 (62.8%) exact matches, in which:
24 unique CDR3
3 unique J



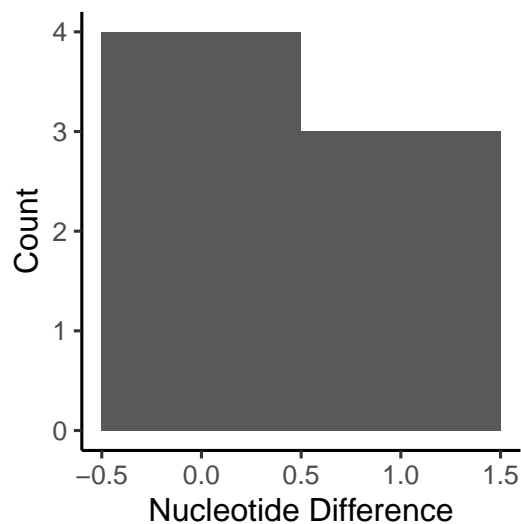
IGLV2-23*01

25 sequences assigned
13 (52%) exact matches, in which:
12 unique CDR3
3 unique J



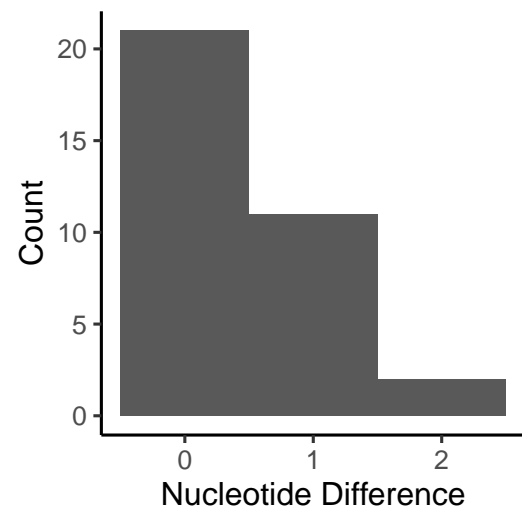
IGLV3-10*01

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



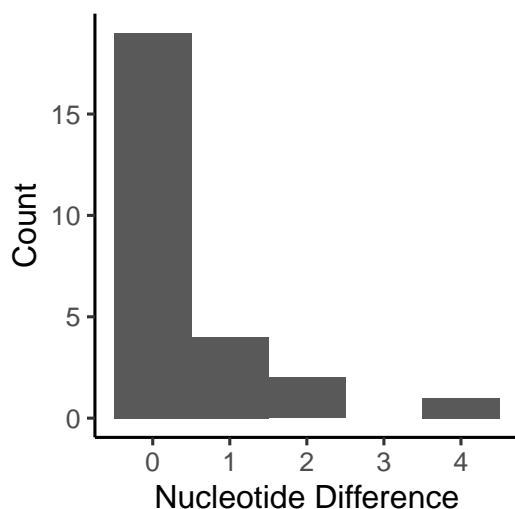
IGLV3-21*02

34 sequences assigned
21 (61.8%) exact matches, in which:
20 unique CDR3
3 unique J



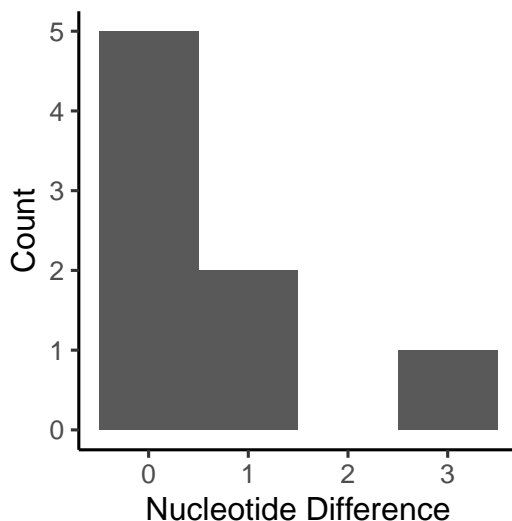
IGLV3–21*03

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



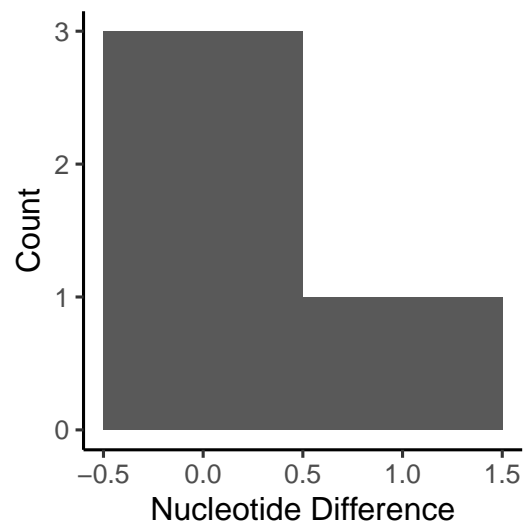
IGLV4–69*01

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



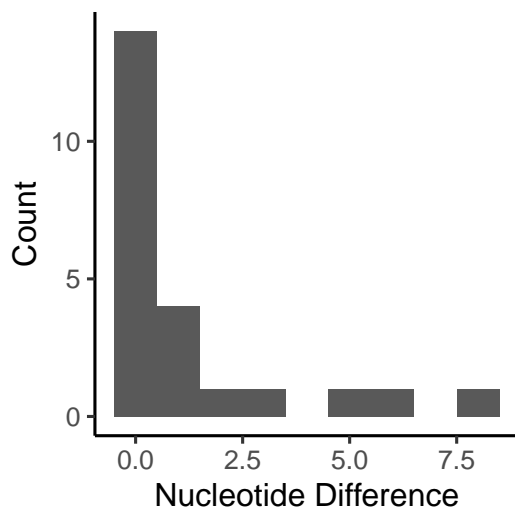
IGLV5–45*02

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



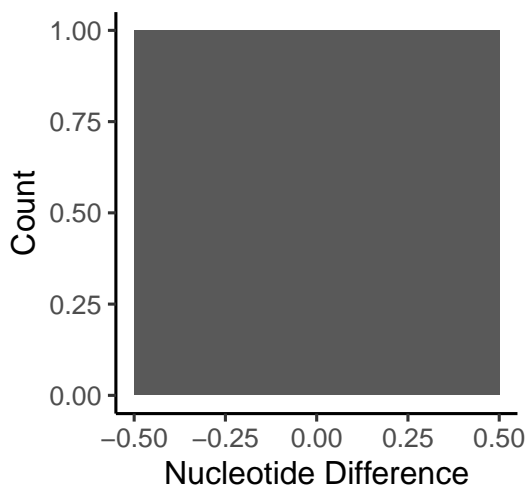
IGLV3–25*03

23 sequences assigned
14 (60.9%) exact matches, in which:
12 unique CDR3
3 unique J



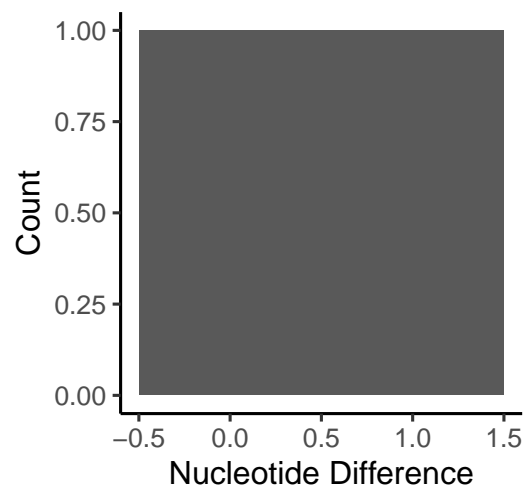
IGLV5–37*01

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



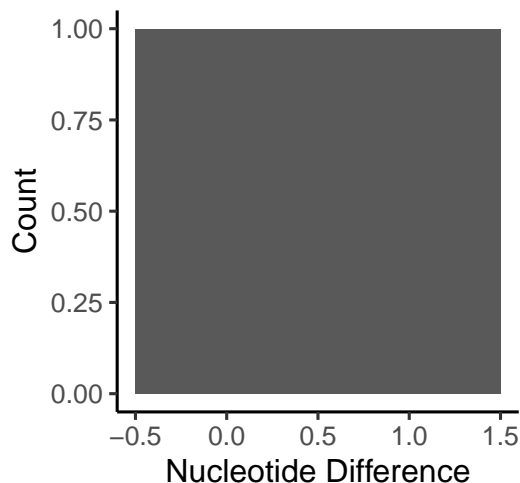
IGLV5–45*03

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



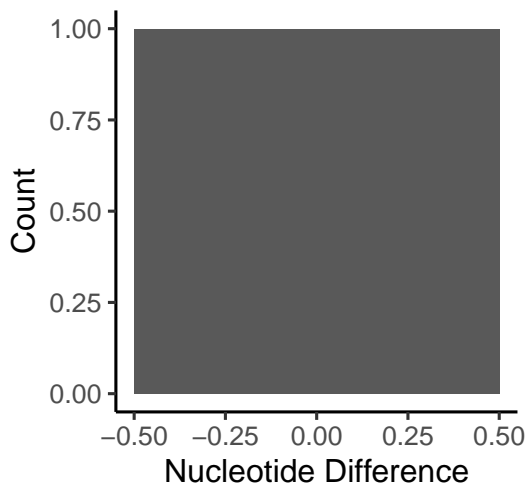
IGLV3–27*01

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



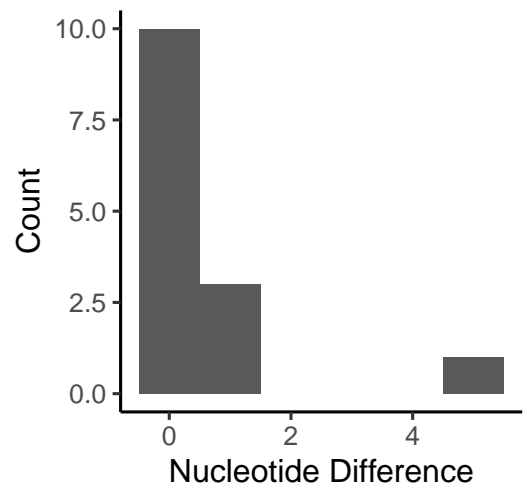
IGLV5–45*01

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



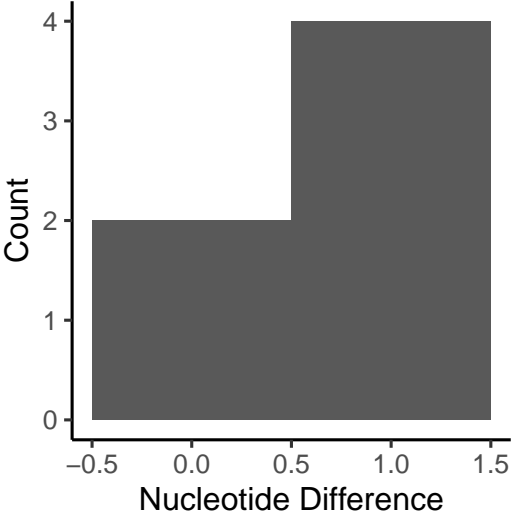
IGLV6–57*02

14 sequences assigned
10 (71.4%) exact matches, in which:
10 unique CDR3
3 unique J



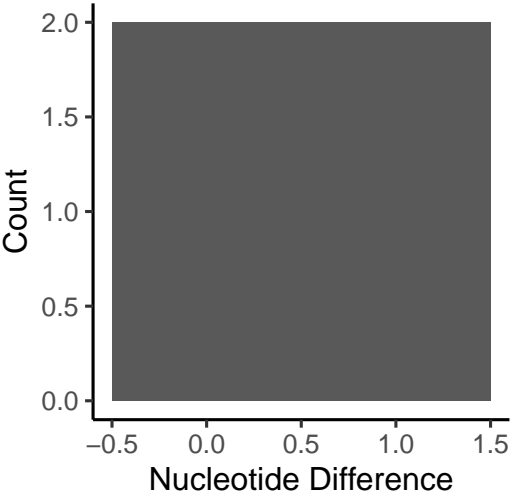
IGLV7-43*01

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



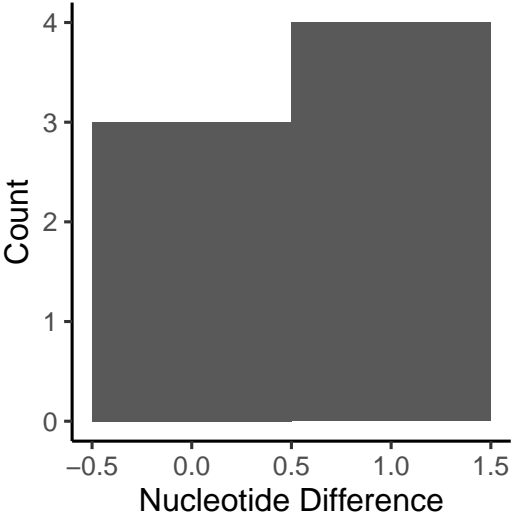
IGLV9-49*01

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



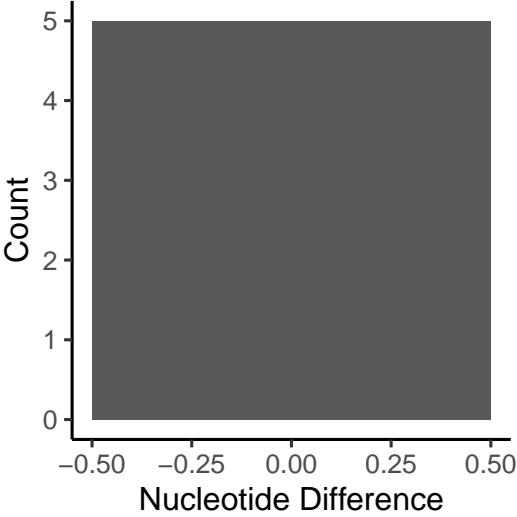
IGLV7-46*01

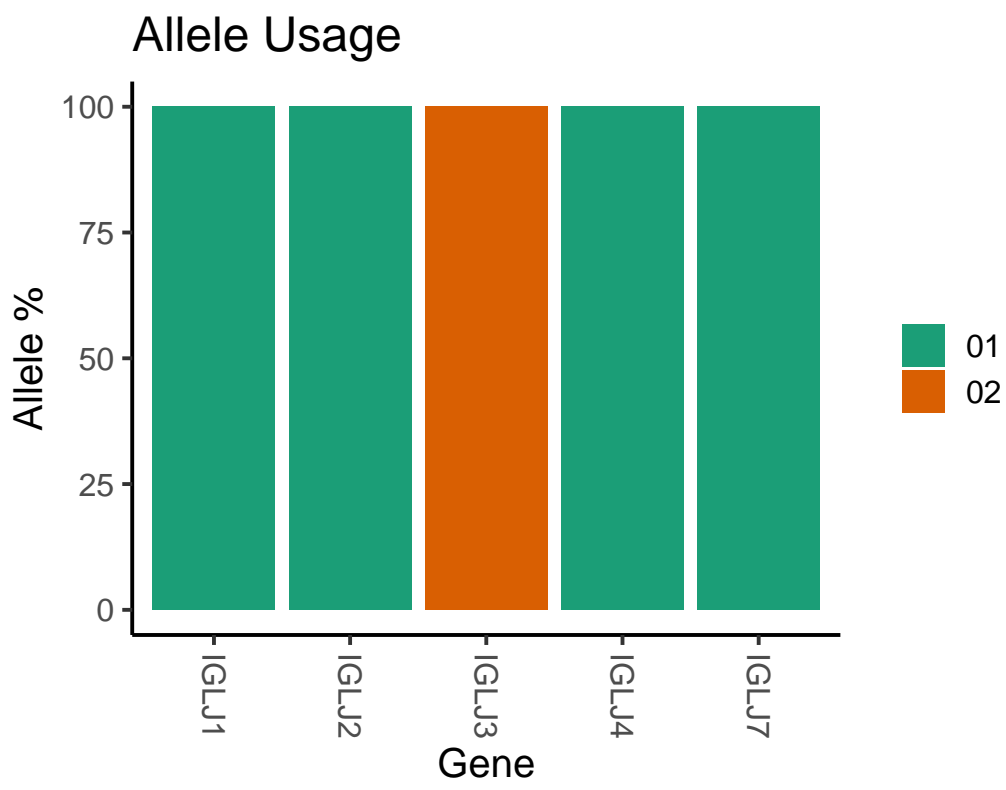
7 sequences assigned
3 (42.9%) exact matches, in which:
3 unique CDR3
2 unique J



IGLV8-61*01

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





Warning – no inferred sequences found.