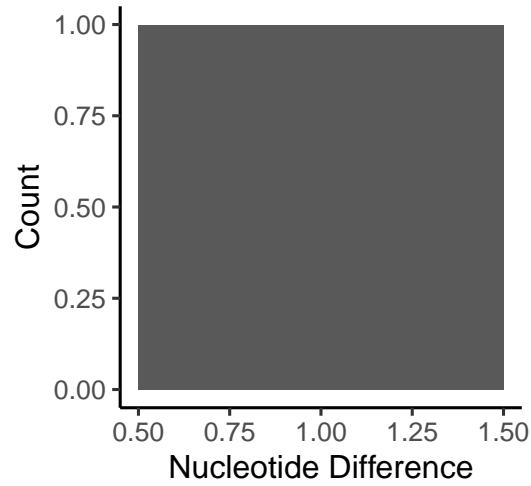


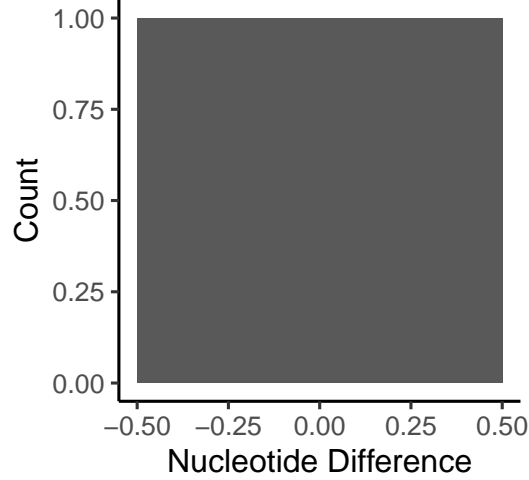
IGLV10-54*02

1 sequences assigned
No exact matches.



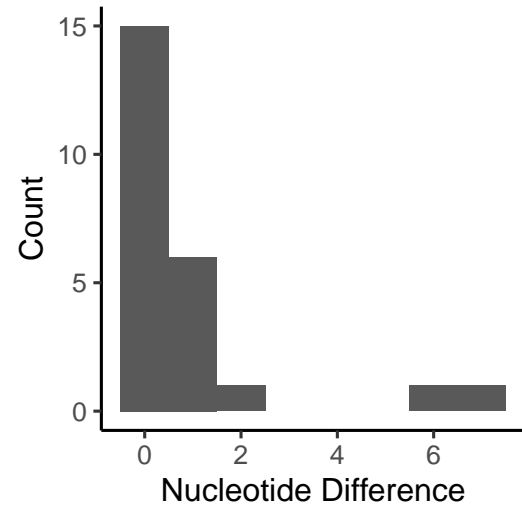
IGLV1-41*01

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



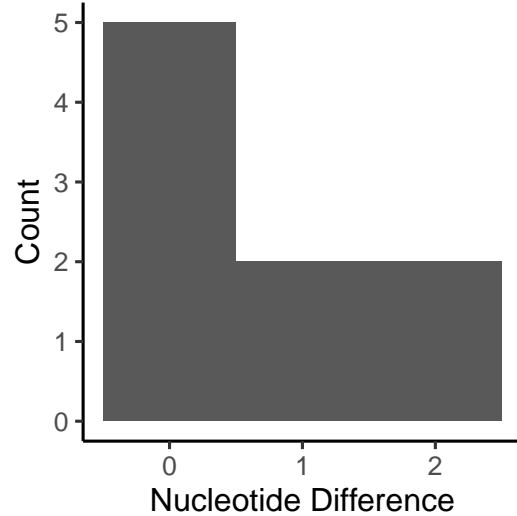
IGLV1-51*01

24 sequences assigned
15 (62.5%) exact matches, in which:
14 unique CDR3
3 unique J



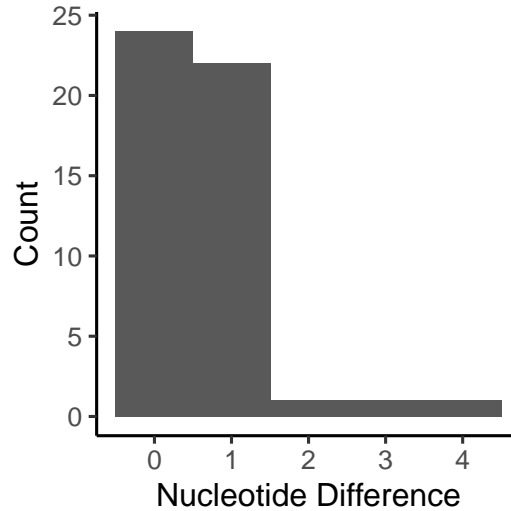
IGLV1-36*01

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



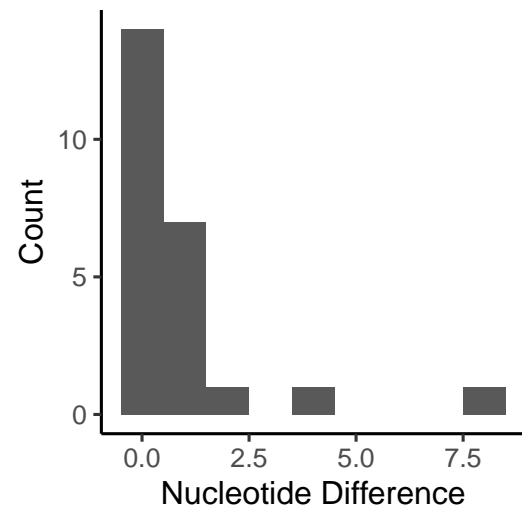
IGLV1-44*01

49 sequences assigned
24 (49%) exact matches, in which:
24 unique CDR3
4 unique J



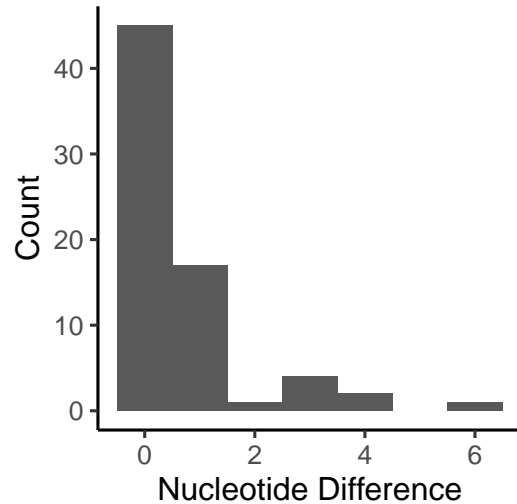
IGLV2-8*01

24 sequences assigned
14 (58.3%) exact matches, in which:
13 unique CDR3
3 unique J



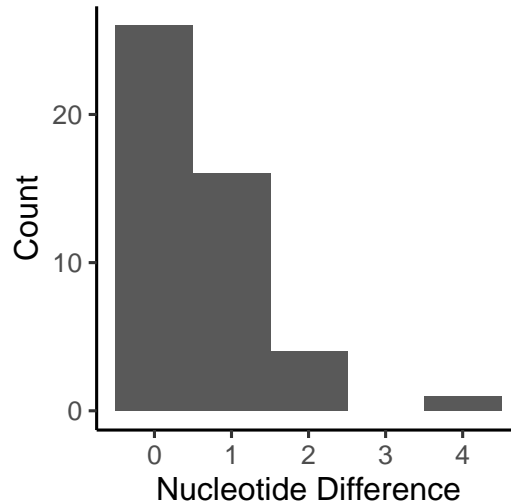
IGLV1-40*01

70 sequences assigned
45 (64.3%) exact matches, in which:
37 unique CDR3
3 unique J



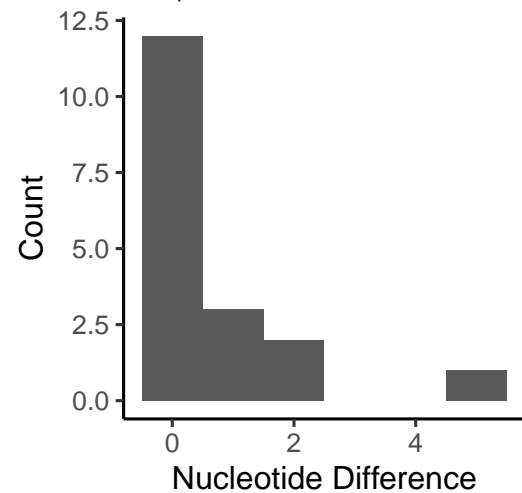
IGLV1-47*01

47 sequences assigned
26 (55.3%) exact matches, in which:
23 unique CDR3
4 unique J



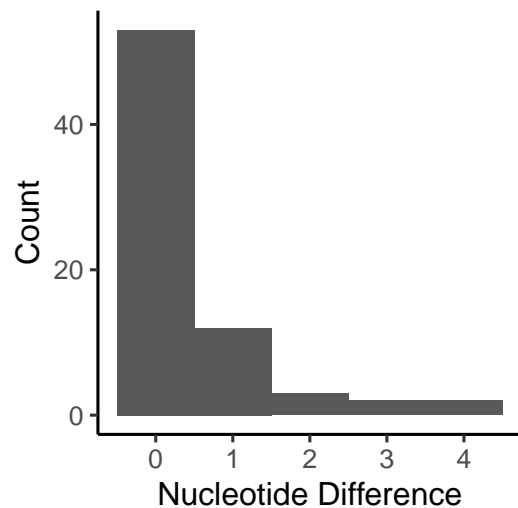
IGLV2-11*01

18 sequences assigned
12 (66.7%) exact matches, in which:
10 unique CDR3
3 unique J



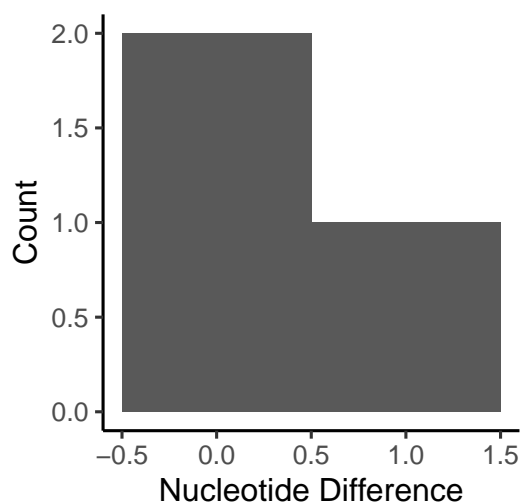
IGLV2-14*01

72 sequences assigned
53 (73.6%) exact matches, in which:
48 unique CDR3
4 unique J



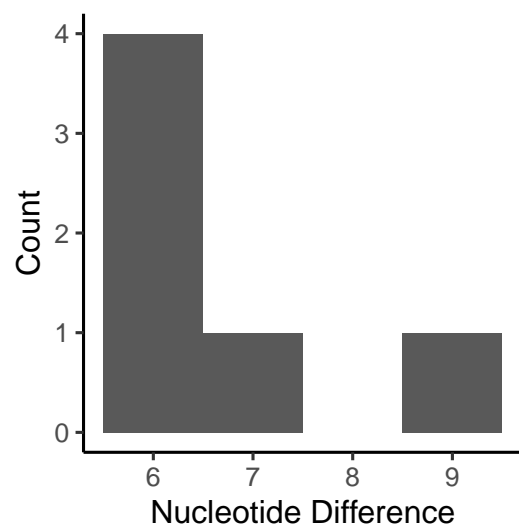
IGLV2-18*02

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



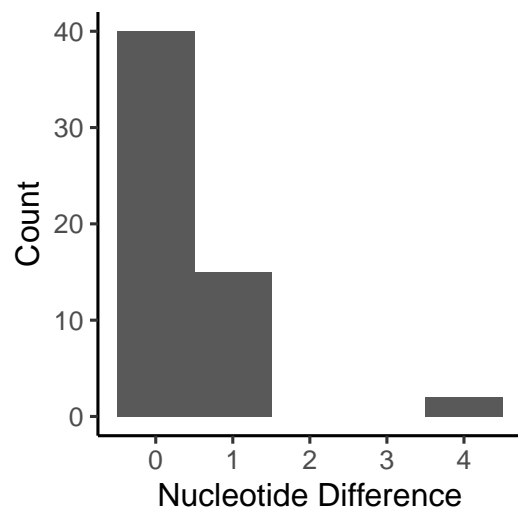
IGLV3-9*02

6 sequences assigned
No exact matches.



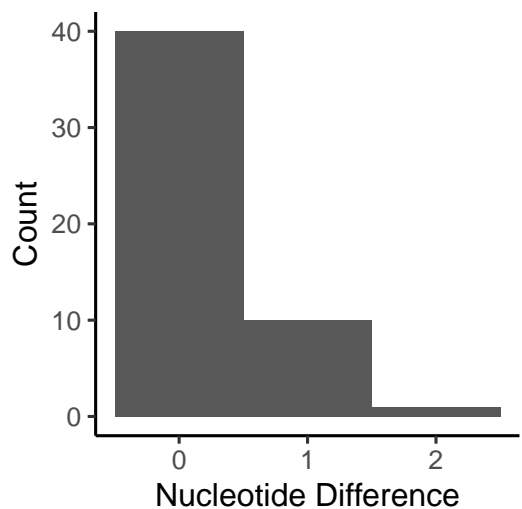
IGLV2-14*04EXT3

57 sequences assigned
40 (70.2%) exact matches, in which:
36 unique CDR3
3 unique J



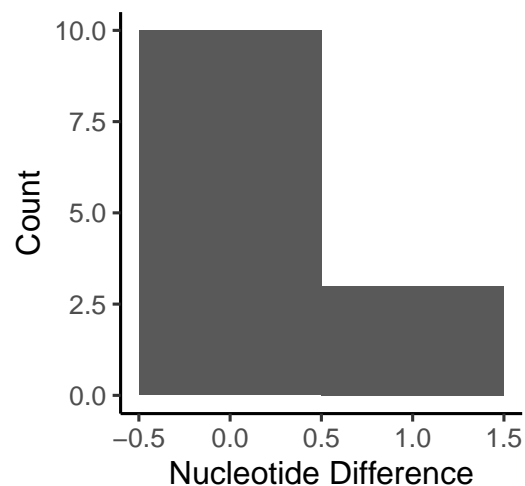
IGLV2-23*01_G170T

51 sequences assigned
40 (78.4%) exact matches, in which:
37 unique CDR3
5 unique J



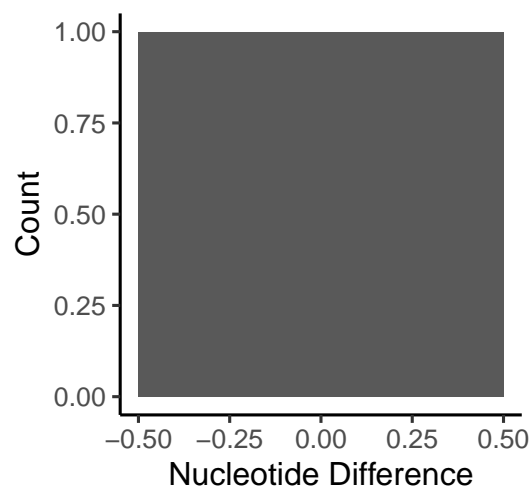
IGLV3-10*01

13 sequences assigned
10 (76.9%) exact matches, in which:
9 unique CDR3
2 unique J



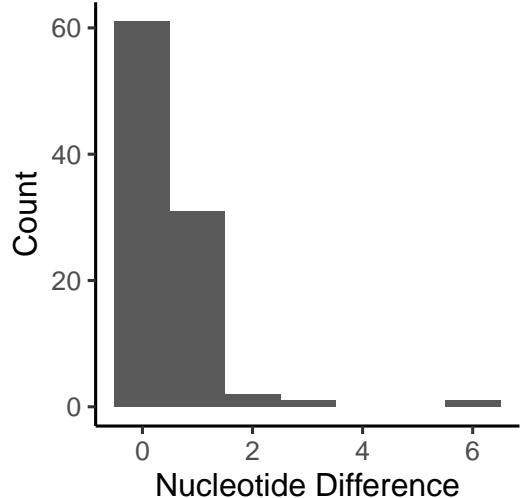
IGLV2-18*01

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



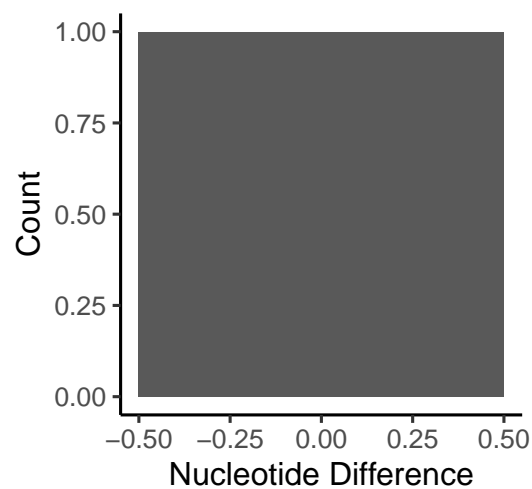
IGLV3-1*01

96 sequences assigned
61 (63.5%) exact matches, in which:
55 unique CDR3
3 unique J



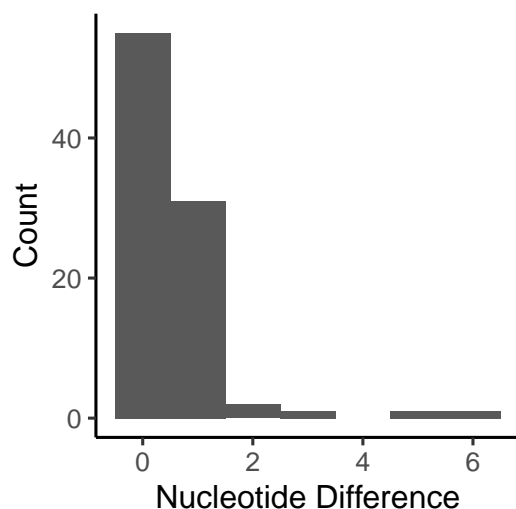
IGLV3-12*02

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



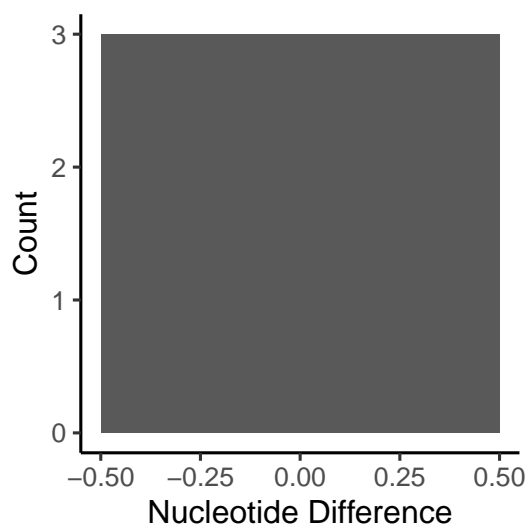
IGLV3–19*01

91 sequences assigned
55 (60.4%) exact matches, in which:
49 unique CDR3
3 unique J



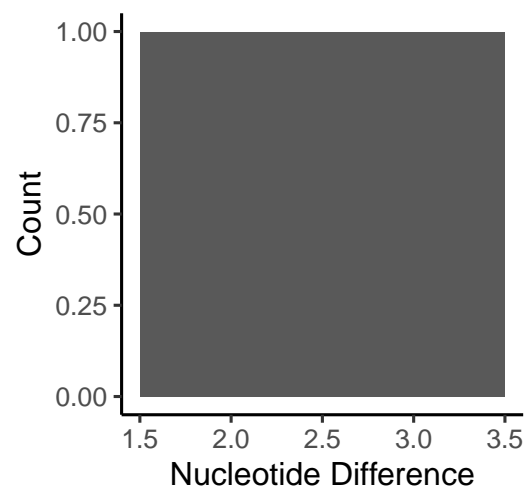
IGLV3–27*01

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



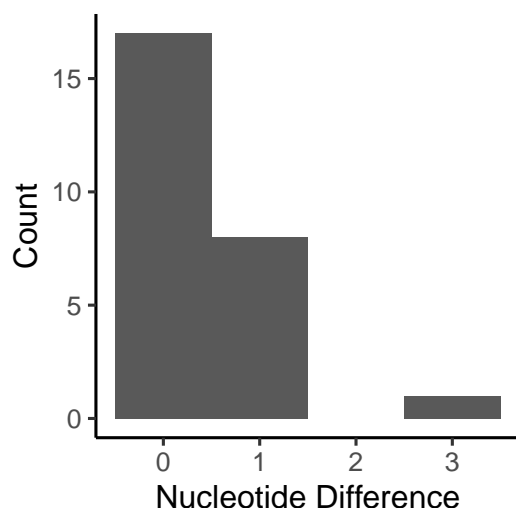
IGLV5–39*01

2 sequences assigned
No exact matches.



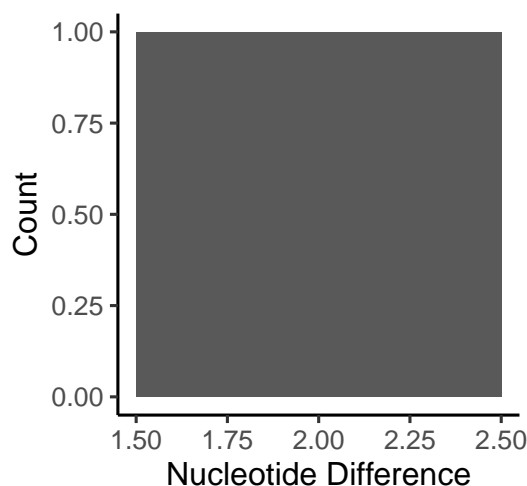
IGLV3–21*04

26 sequences assigned
17 (65.4%) exact matches, in which:
15 unique CDR3
3 unique J



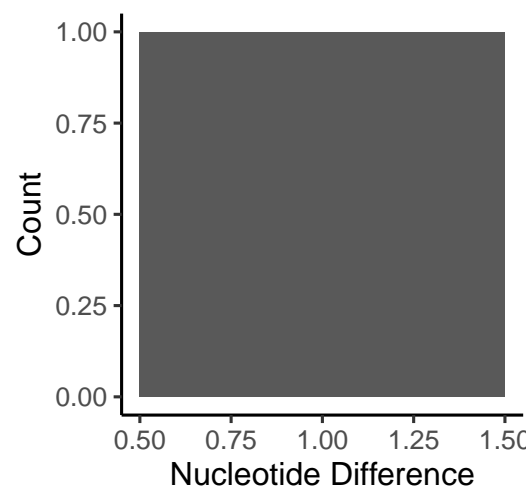
IGLV4–60*02

1 sequences assigned
No exact matches.



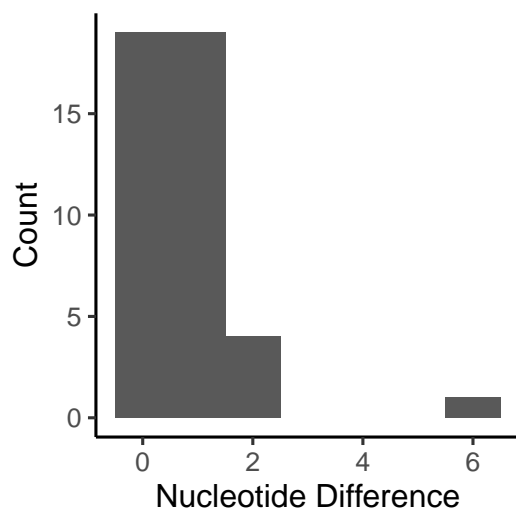
IGLV5–45*02

1 sequences assigned
No exact matches.



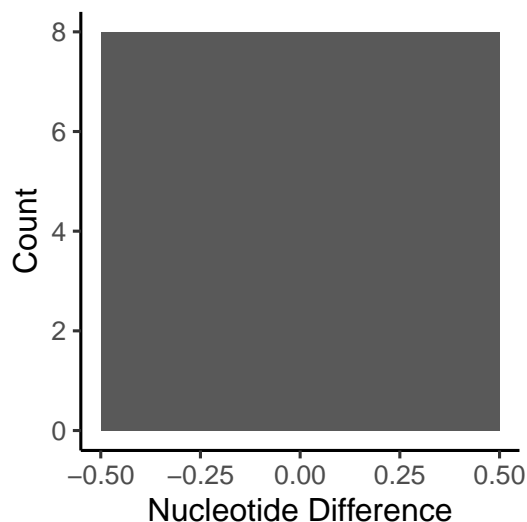
IGLV3–25*03

43 sequences assigned
19 (44.2%) exact matches, in which:
18 unique CDR3
4 unique J



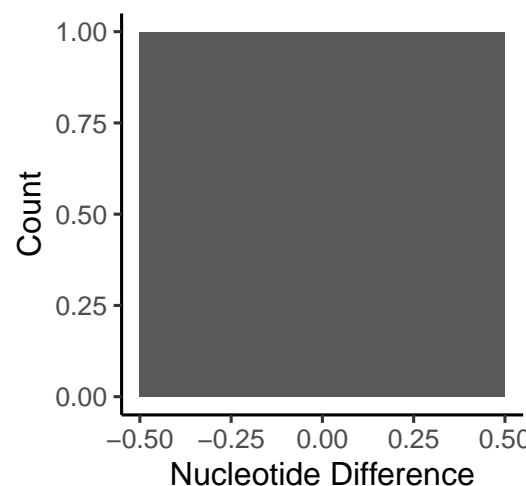
IGLV4–69*01

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



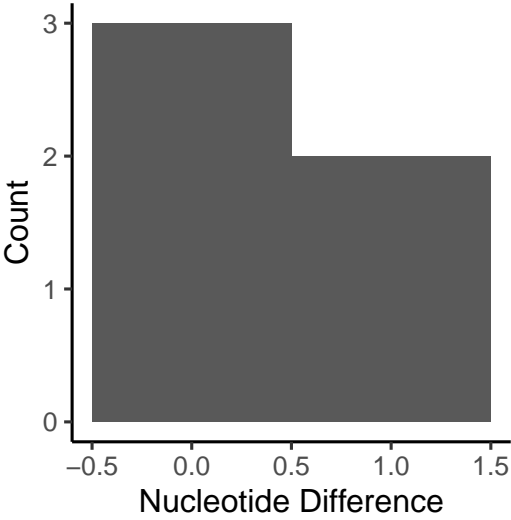
IGLV5–45*03

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



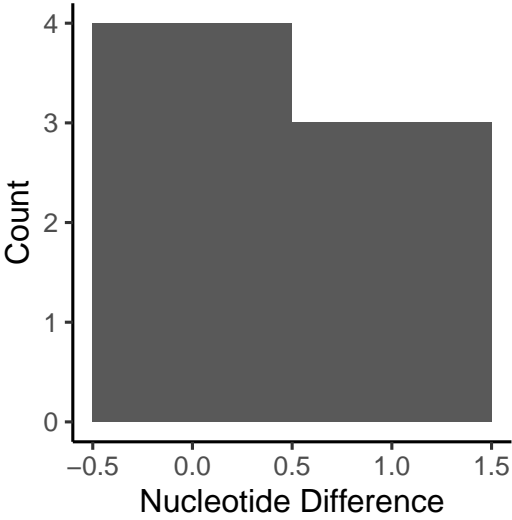
IGLV6–57*02

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



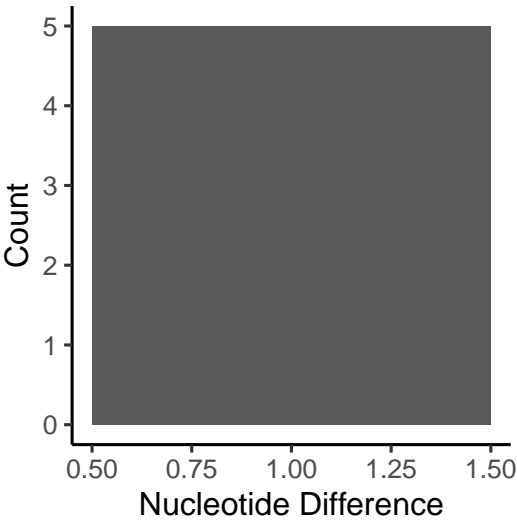
IGLV7–46*01

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



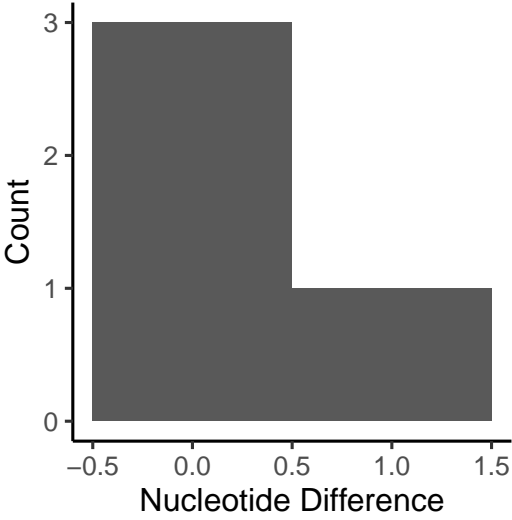
IGLV6–57*03

5 sequences assigned
No exact matches.



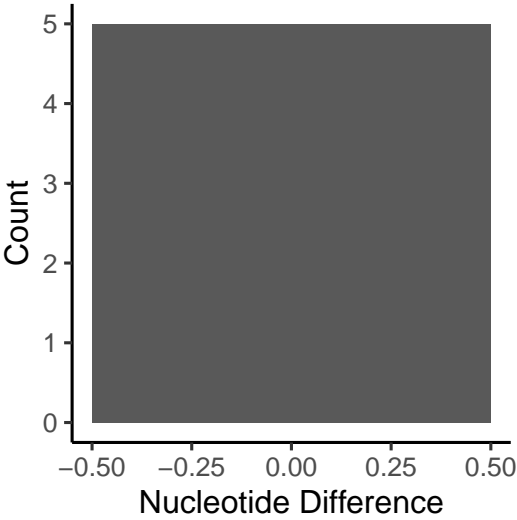
IGLV9–49*01

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

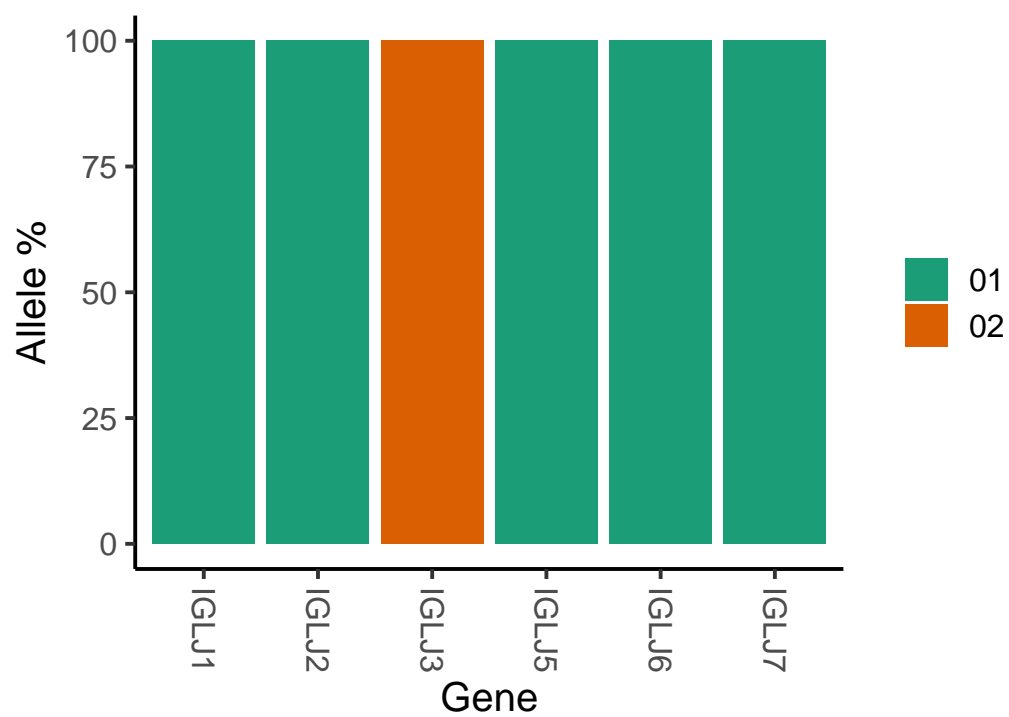


IGLV7–43*01

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



Allele Usage



Warning – no inferred sequences found.