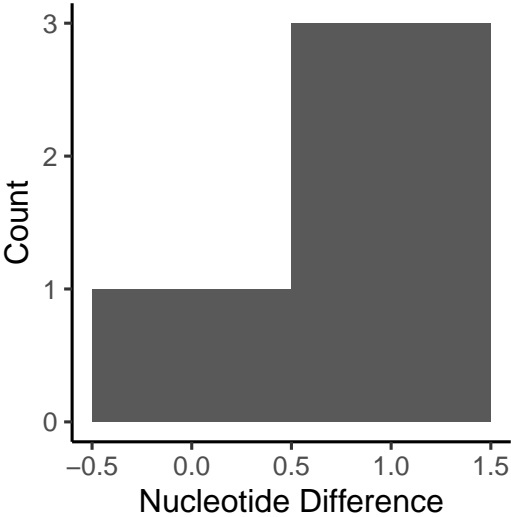


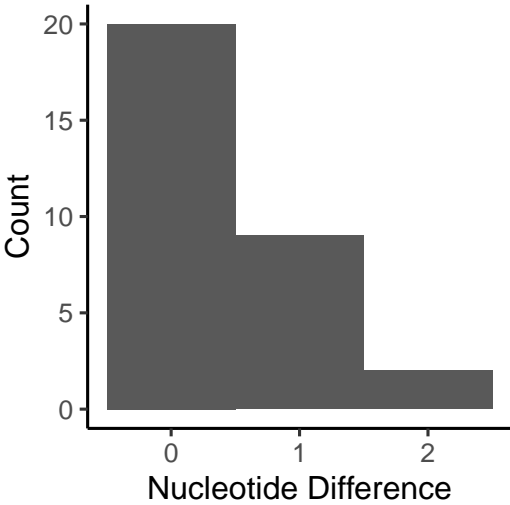
IGLV10–54*01

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



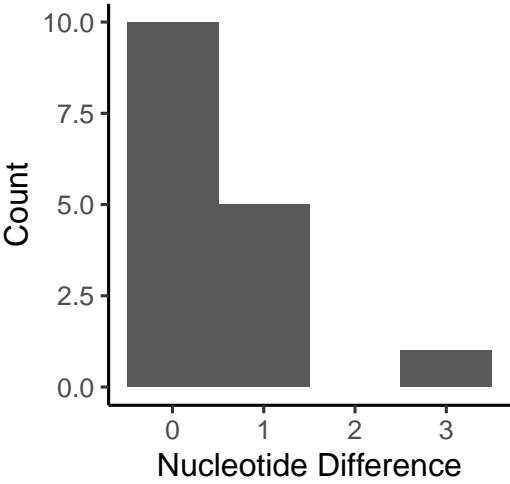
IGLV1–44*01

32 sequences assigned
20 (62.5%) exact matches, in which:
18 unique CDR3
3 unique J



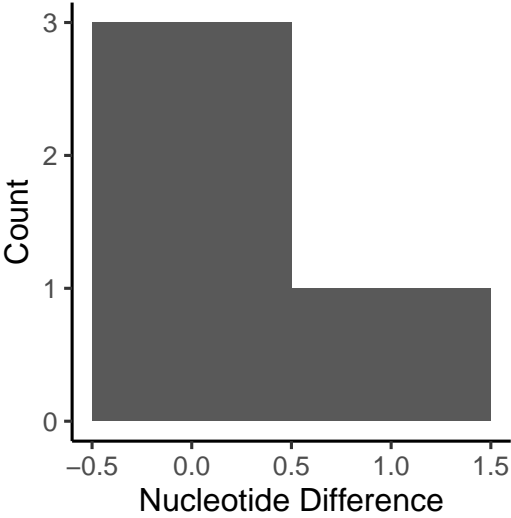
IGLV1–51*02

16 sequences assigned
10 (62.5%) exact matches, in which:
9 unique CDR3
3 unique J



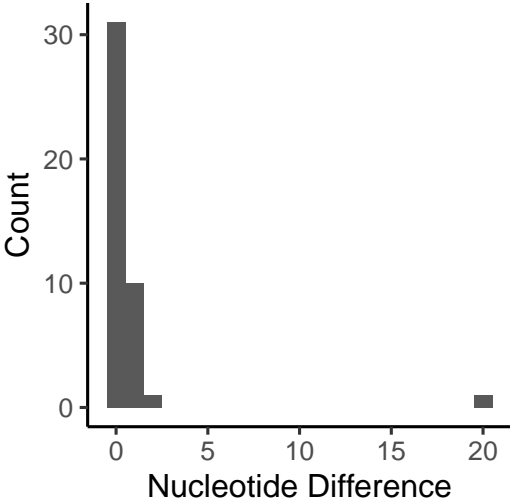
IGLV1–36*01

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



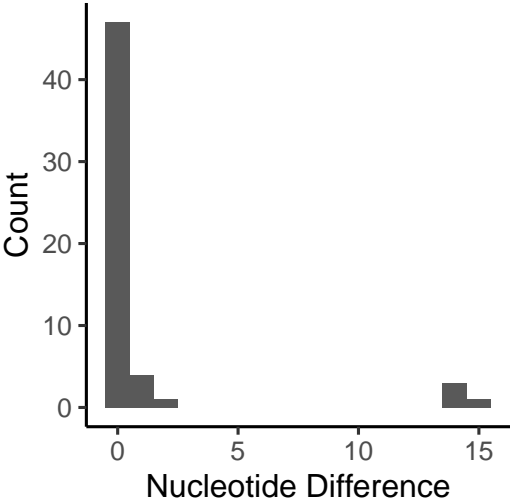
IGLV1–47*01

43 sequences assigned
31 (72.1%) exact matches, in which:
27 unique CDR3
4 unique J



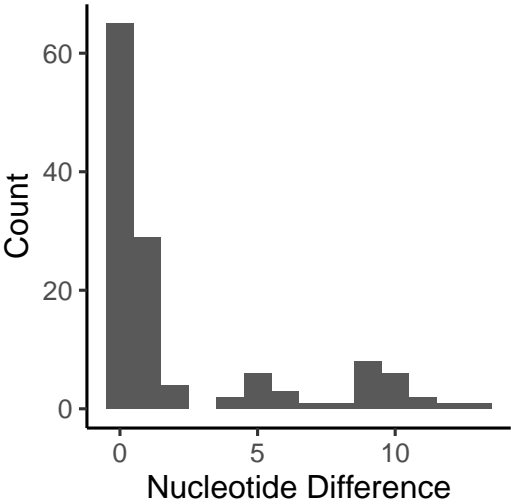
IGLV2–8*01

56 sequences assigned
47 (83.9%) exact matches, in which:
44 unique CDR3
3 unique J



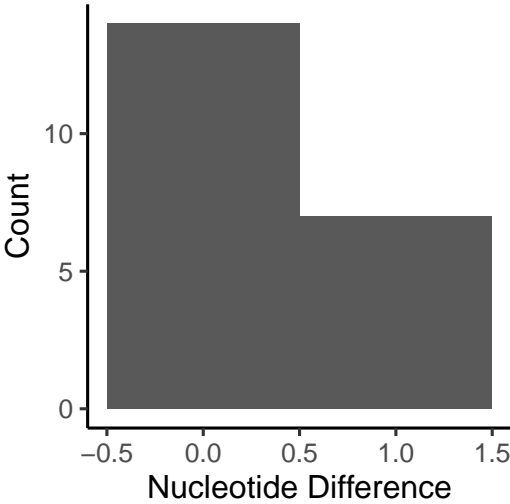
IGLV1–40*01

129 sequences assigned
65 (50.4%) exact matches, in which:
54 unique CDR3
3 unique J



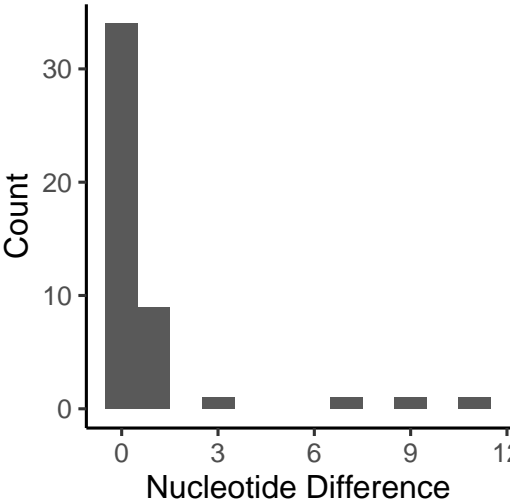
IGLV1–51*01

21 sequences assigned
14 (66.7%) exact matches, in which:
12 unique CDR3
2 unique J



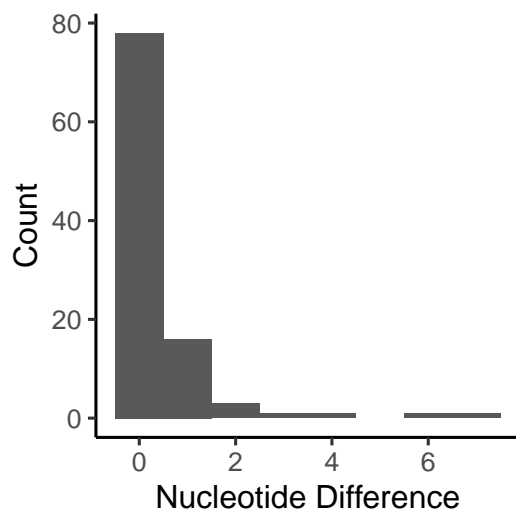
IGLV2–11*01

47 sequences assigned
34 (72.3%) exact matches, in which:
28 unique CDR3
3 unique J



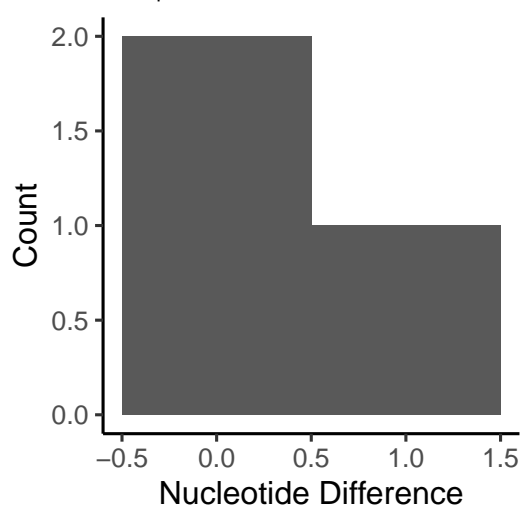
IGLV2-14*01

101 sequences assigned
78 (77.2%) exact matches, in which:
73 unique CDR3
3 unique J



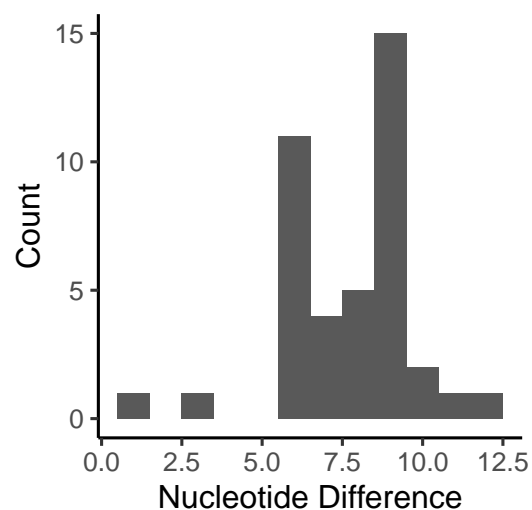
IGLV2-18*02

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



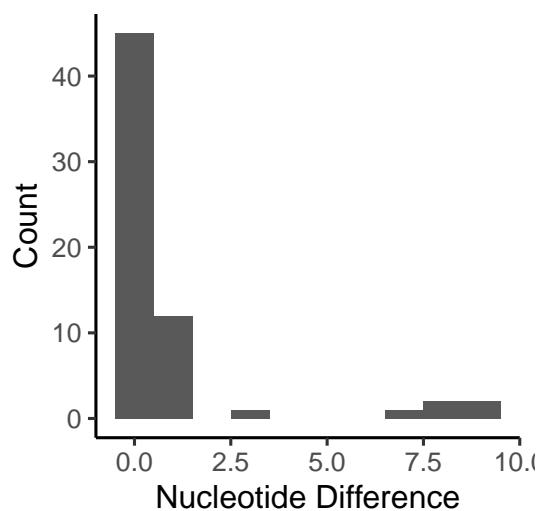
IGLV3-9*02

41 sequences assigned
No exact matches.



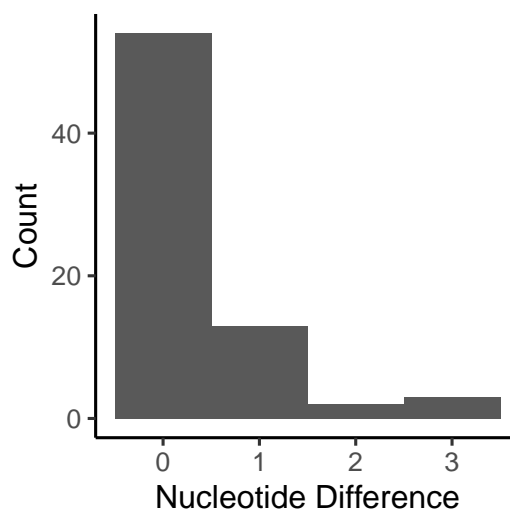
IGLV2-14*04EXT3

63 sequences assigned
45 (71.4%) exact matches, in which:
44 unique CDR3
3 unique J



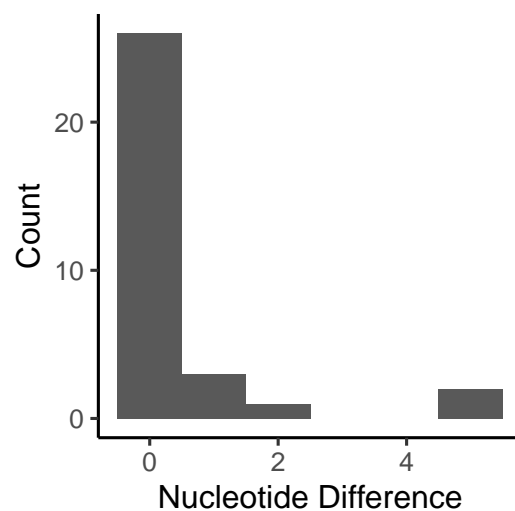
IGLV2-23*01

72 sequences assigned
54 (75%) exact matches, in which:
48 unique CDR3
4 unique J



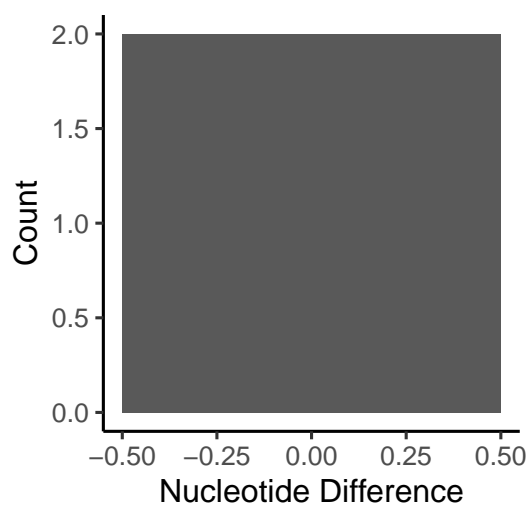
IGLV3-10*01

32 sequences assigned
26 (81.2%) exact matches, in which:
24 unique CDR3
3 unique J



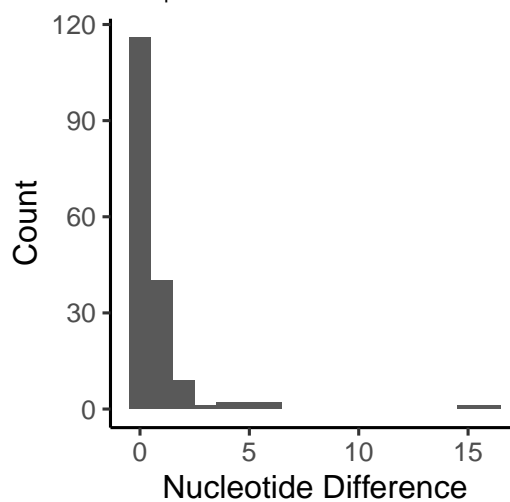
IGLV2-18*01

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



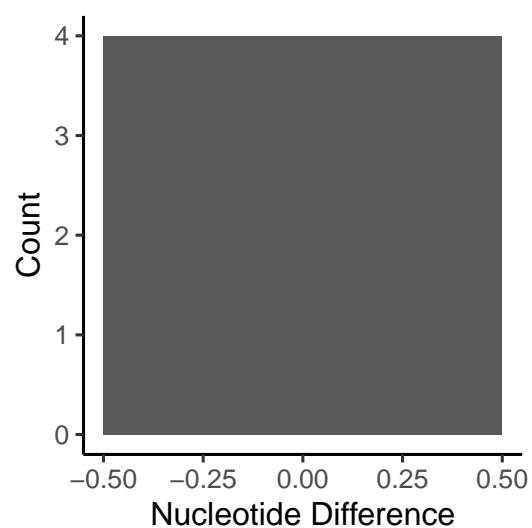
IGLV3-1*01

175 sequences assigned
116 (66.3%) exact matches, in which:
98 unique CDR3
3 unique J



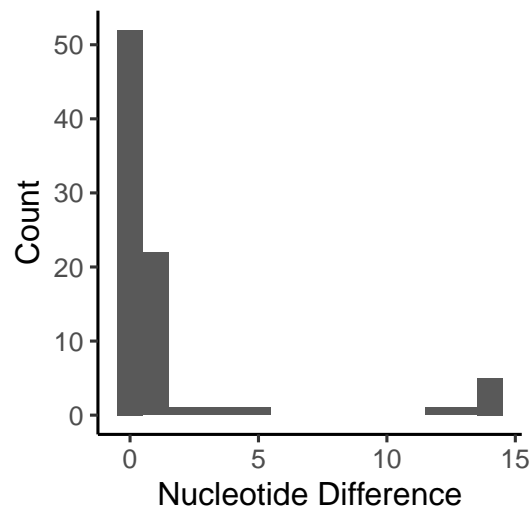
IGLV3-16*01

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



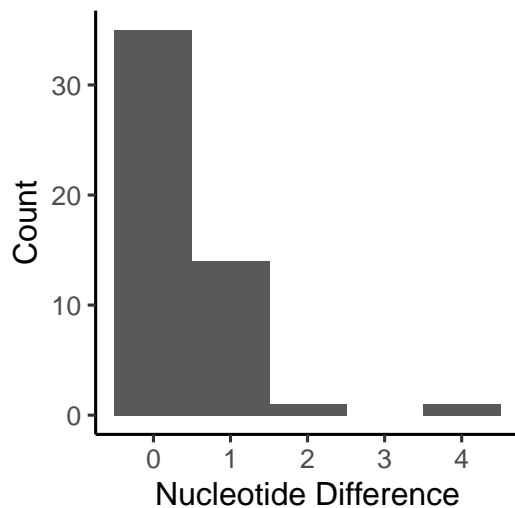
IGLV3-19*01

85 sequences assigned
52 (61.2%) exact matches, in which:
46 unique CDR3
3 unique J



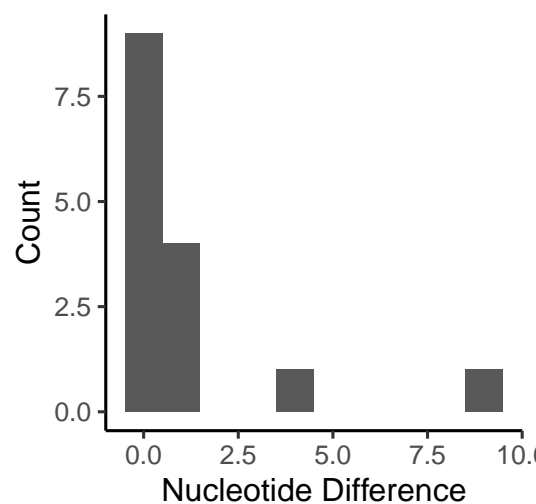
IGLV3-25*02

51 sequences assigned
35 (68.6%) exact matches, in which:
33 unique CDR3
3 unique J



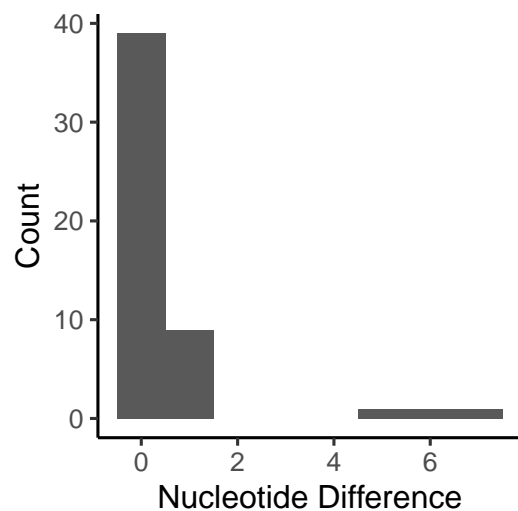
IGLV4-69*01

22 sequences assigned
9 (40.9%) exact matches, in which:
9 unique CDR3
2 unique J



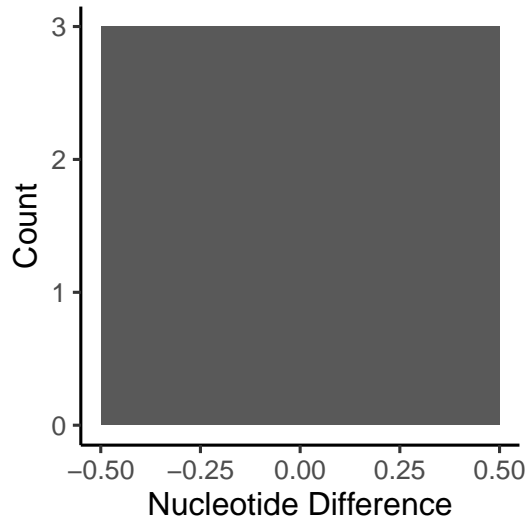
IGLV3-21*02

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



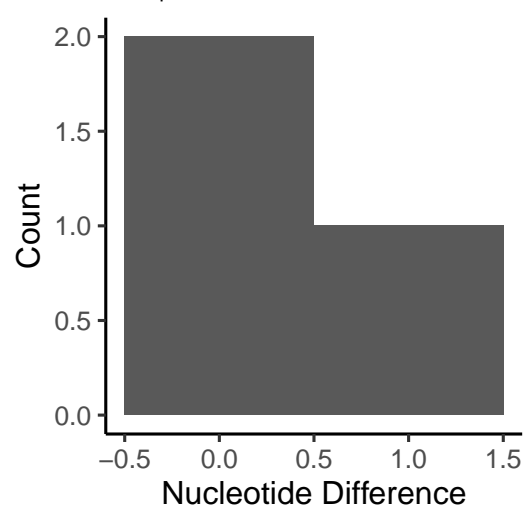
IGLV3-27*01

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



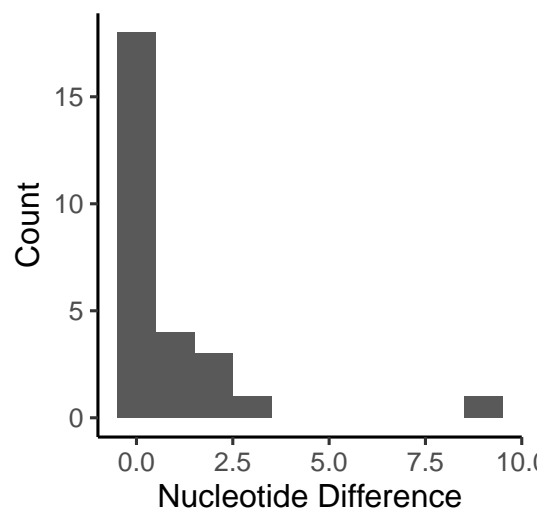
IGLV5-37*01

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



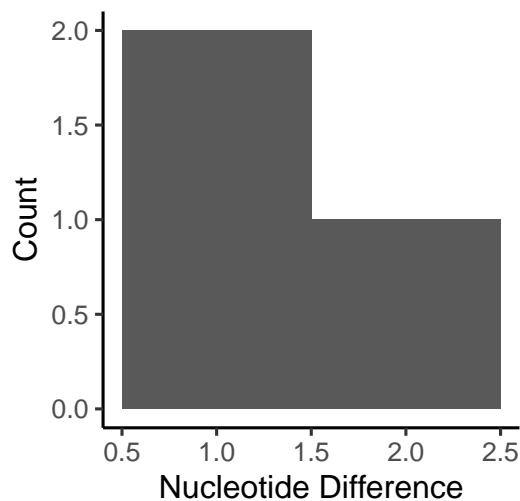
IGLV3-21*04

27 sequences assigned
18 (66.7%) exact matches, in which:
15 unique CDR3
4 unique J



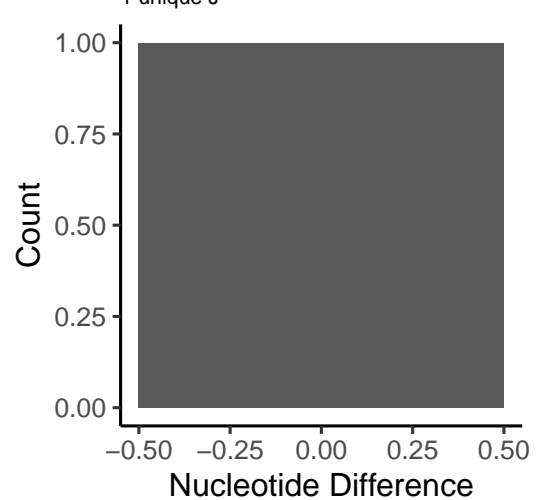
IGLV4-60*02

3 sequences assigned
No exact matches.



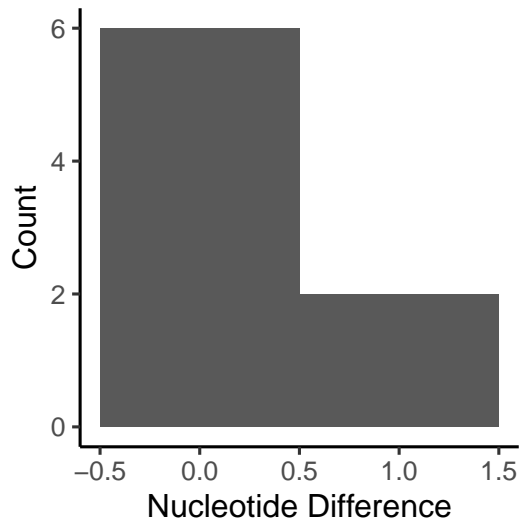
IGLV5-39*01

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



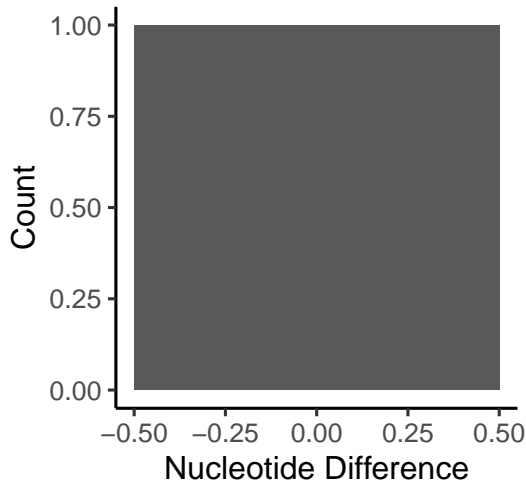
IGLV5-45*01

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



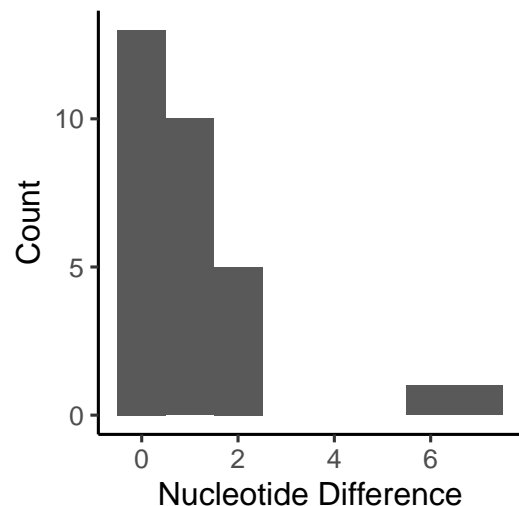
IGLV5-52*01

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



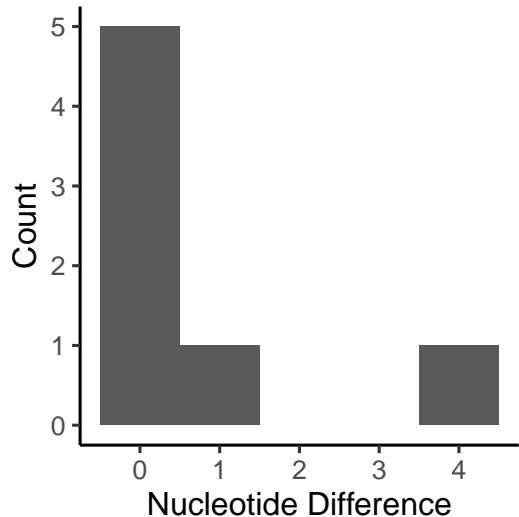
IGLV7-46*01

30 sequences assigned
13 (43.3%) exact matches, in which:
12 unique CDR3
4 unique J



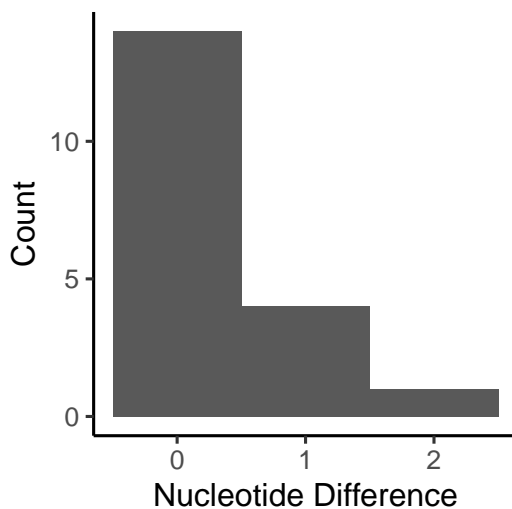
IGLV5-45*02

7 sequences assigned
5 (71.4%) exact matches, in which:
4 unique CDR3
2 unique J



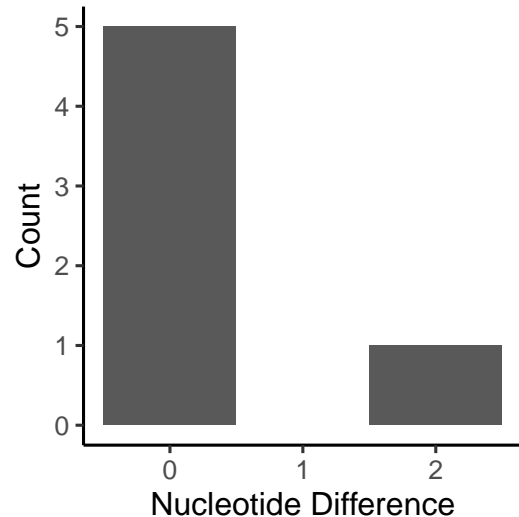
IGLV6-57*02

19 sequences assigned
14 (73.7%) exact matches, in which:
13 unique CDR3
3 unique J



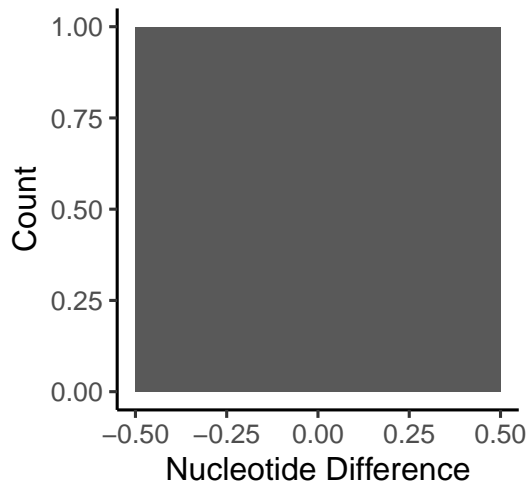
IGLV8-61*01

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



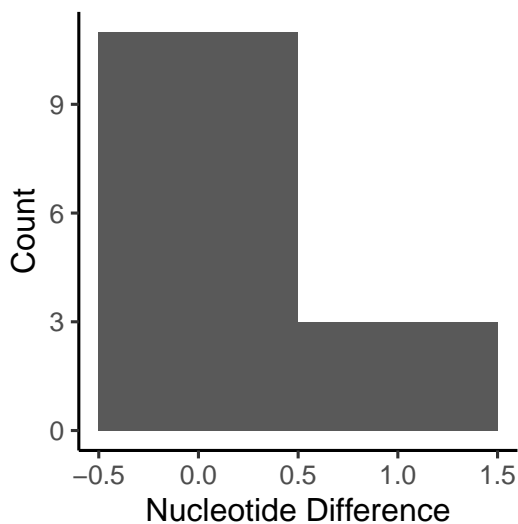
IGLV5-48*01

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



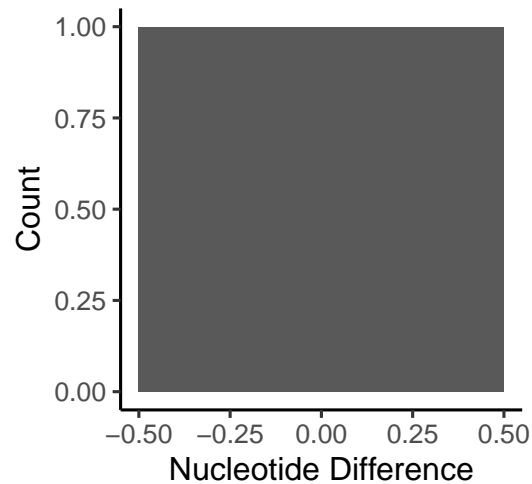
IGLV7-43*01

14 sequences assigned
11 (78.6%) exact matches, in which:
9 unique CDR3
3 unique J



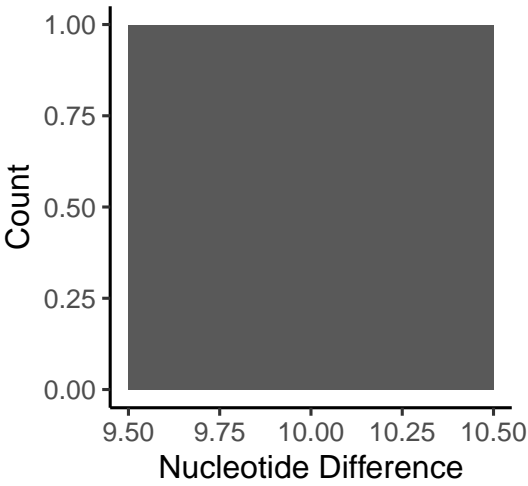
IGLV9-49*01

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

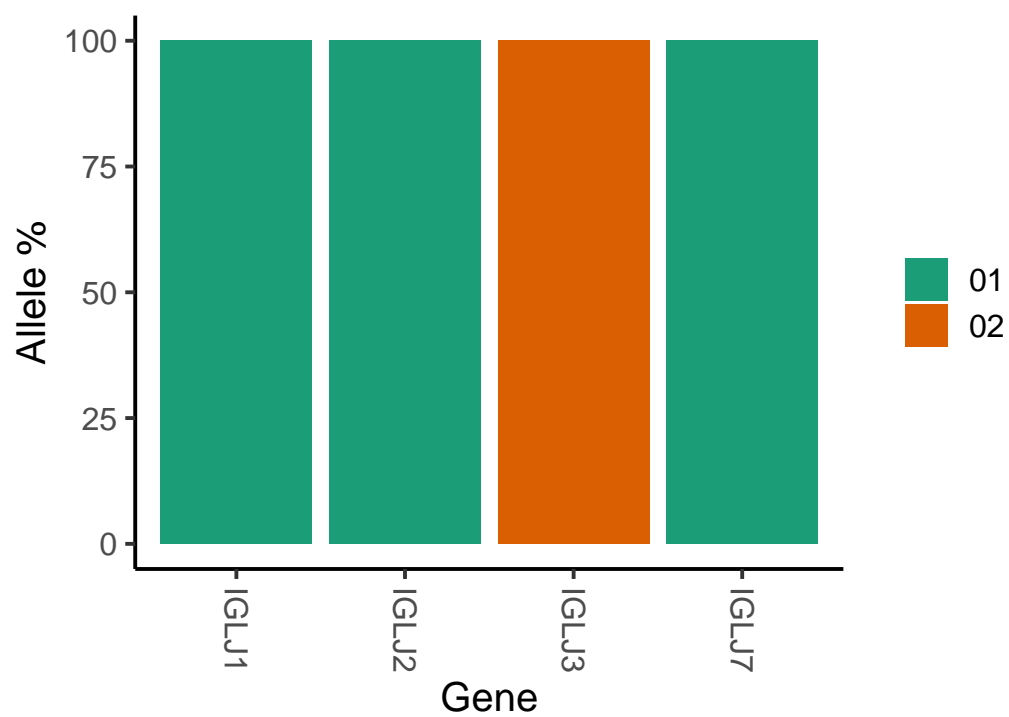


IGLV9-49*02

1 sequences assigned
No exact matches.



Allele Usage



Warning – no inferred sequences found.