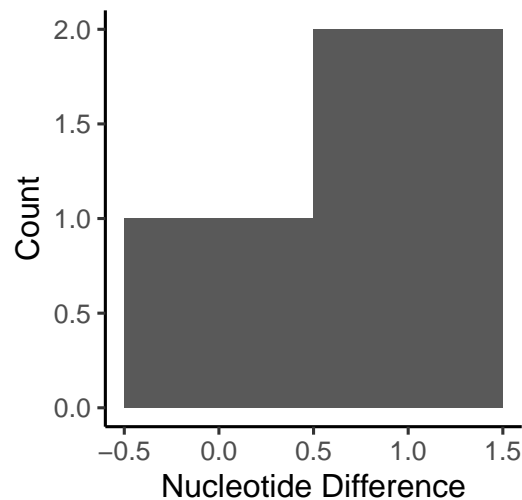


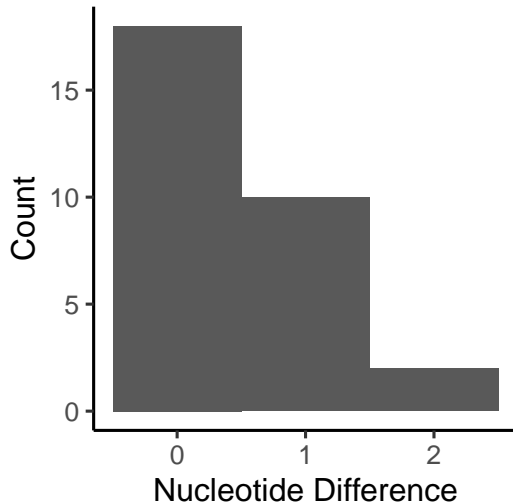
IGLV10–54*01

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



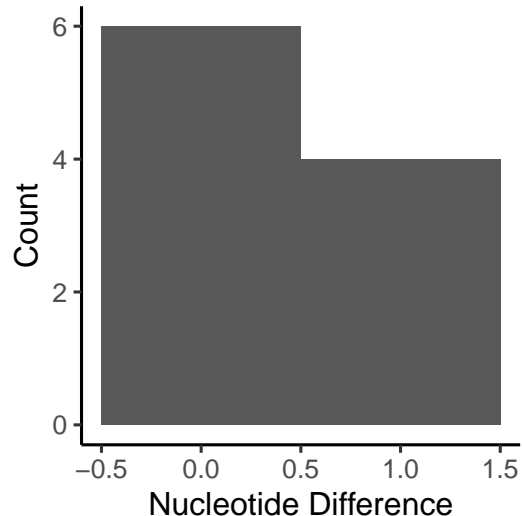
IGLV1–40*01

30 sequences assigned
18 (60%) exact matches, in which:
16 unique CDR3
3 unique J



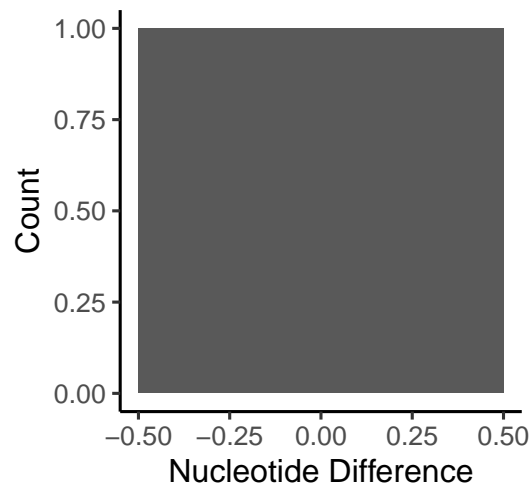
IGLV1–47*02

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



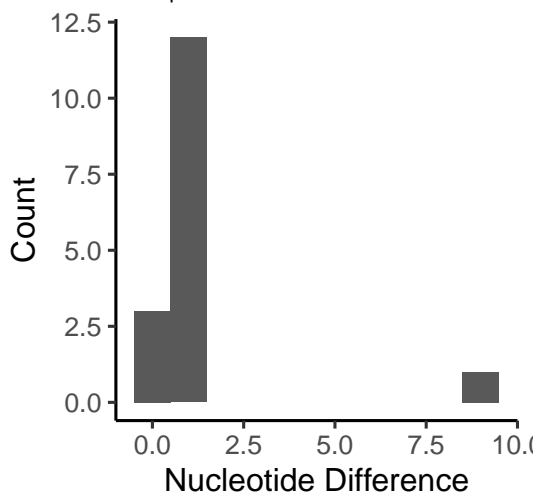
IGLV10–54*02

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



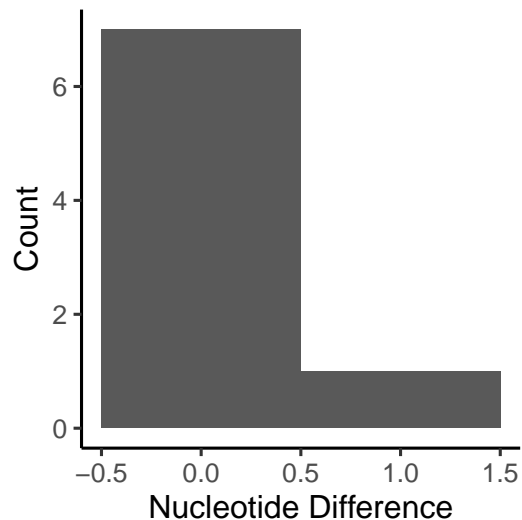
IGLV1–44*01

16 sequences assigned
3 (18.8%) exact matches, in which:
3 unique CDR3
2 unique J



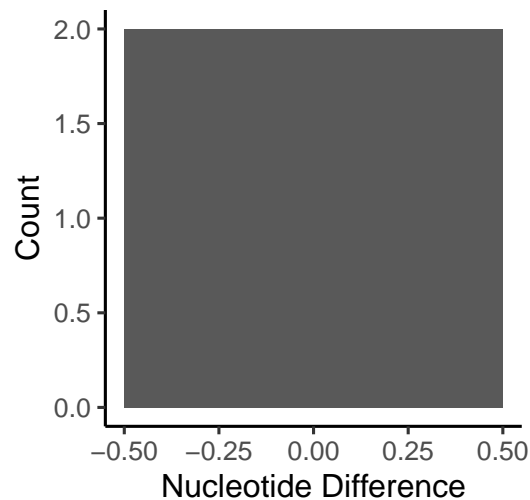
IGLV1–51*01

8 sequences assigned
7 (87.5%) exact matches, in which:
5 unique CDR3
3 unique J



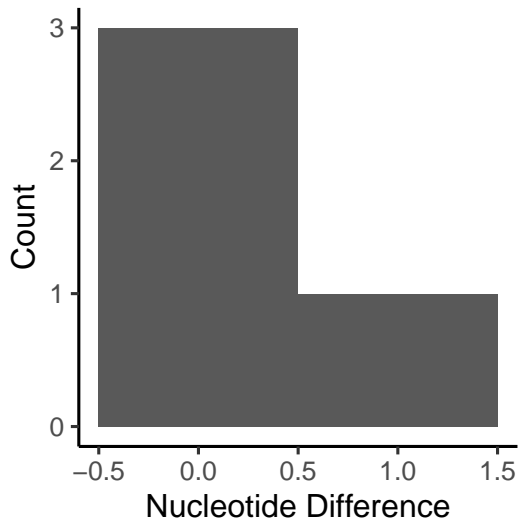
IGLV1–36*01

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



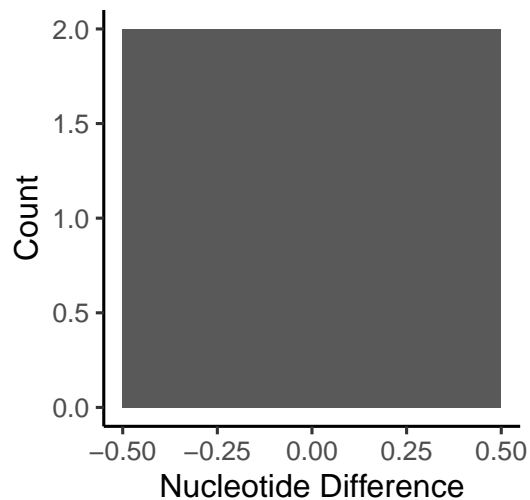
IGLV1–47*01

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



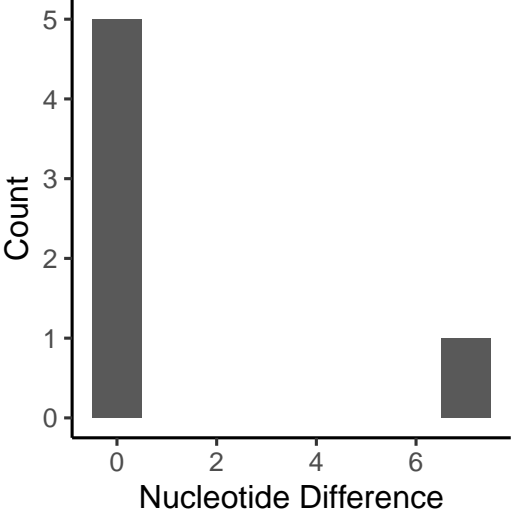
IGLV1–51*02

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



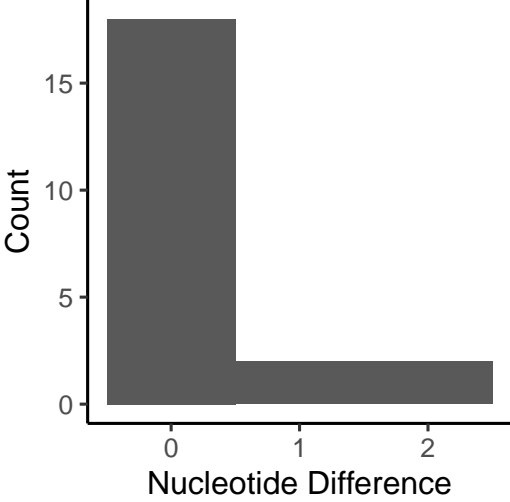
IGLV2-8*01

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



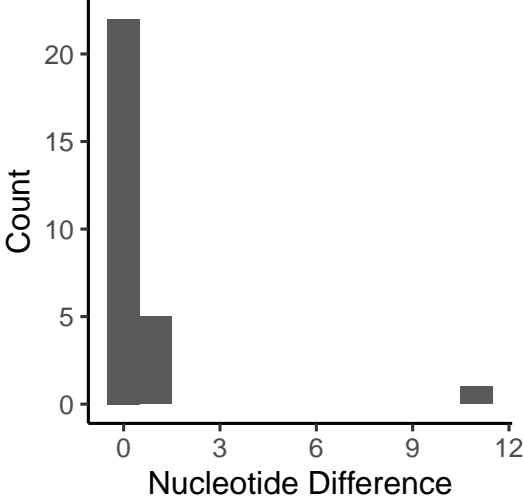
IGLV2-14*04EXT3

22 sequences assigned
18 (81.8%) exact matches, in which:
17 unique CDR3
2 unique J



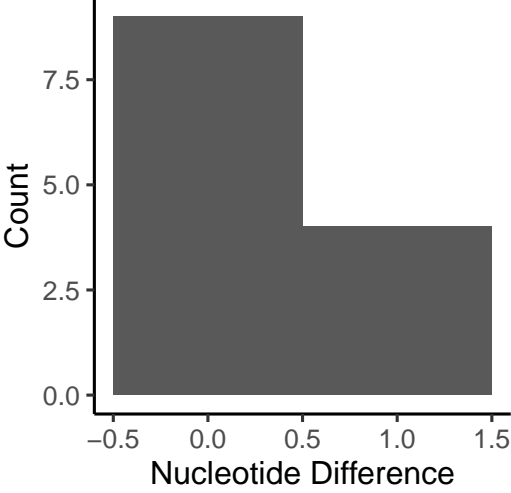
IGLV3-1*01

28 sequences assigned
22 (78.6%) exact matches, in which:
22 unique CDR3
3 unique J



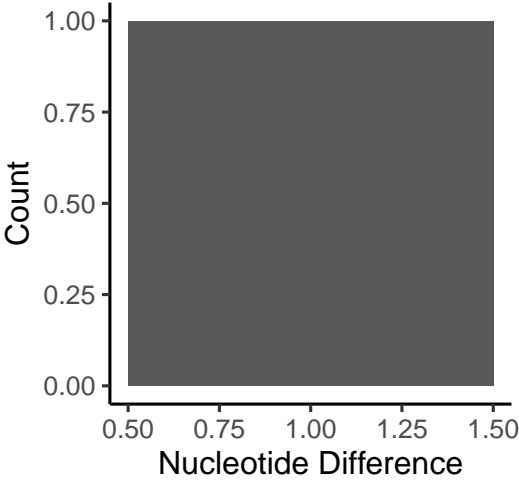
IGLV2-11*01

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



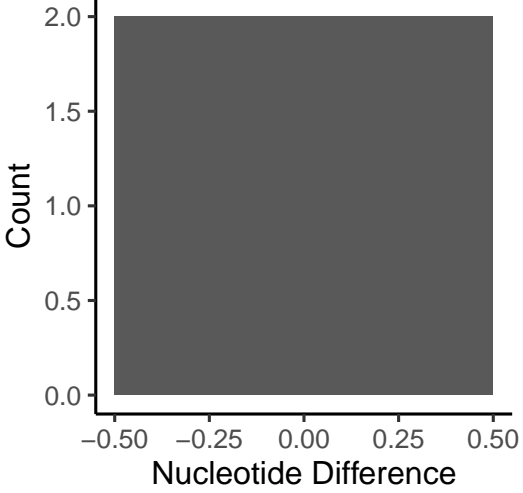
IGLV2-18*02

1 sequences assigned
No exact matches.



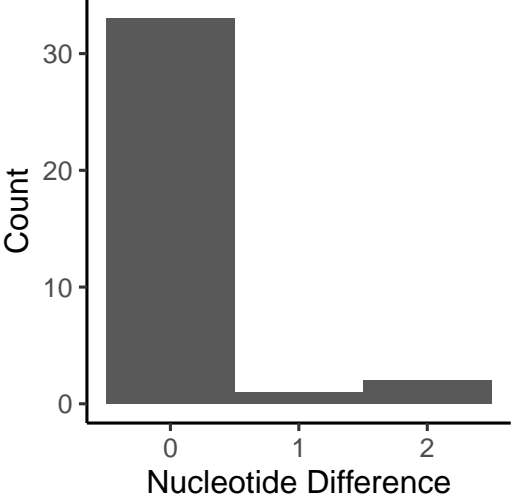
IGLV3-9*01EXT5

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



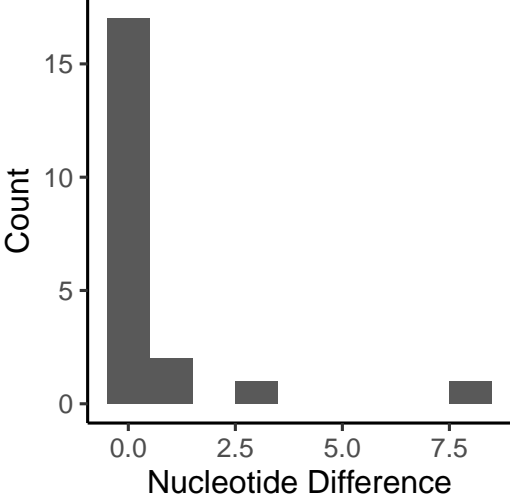
IGLV2-14*03EXT3

36 sequences assigned
33 (91.7%) exact matches, in which:
31 unique CDR3
2 unique J



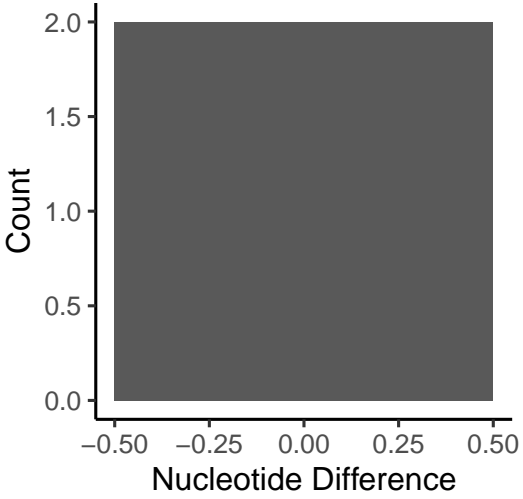
IGLV2-23*02EXT5

21 sequences assigned
17 (81%) exact matches, in which:
17 unique CDR3
2 unique J



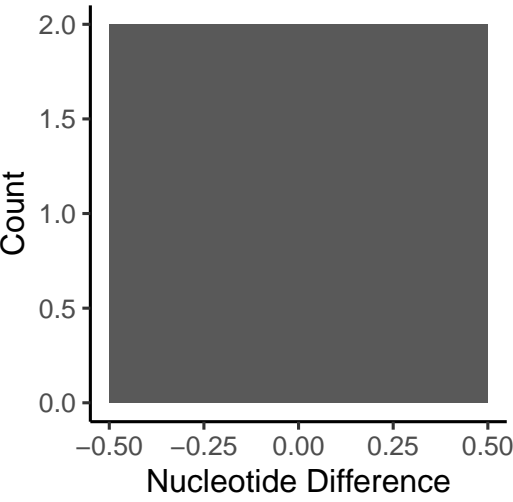
IGLV3-10*01

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



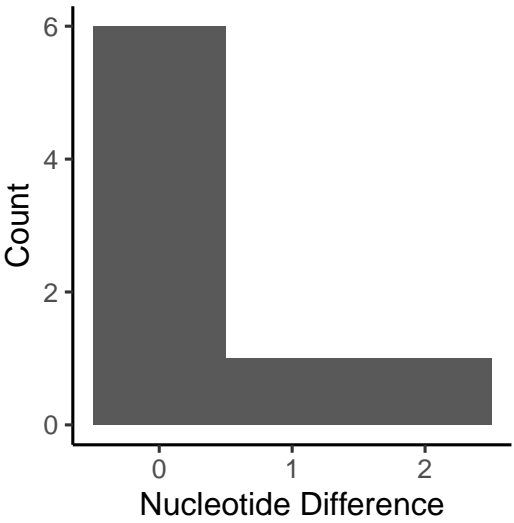
IGLV3-16*01

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



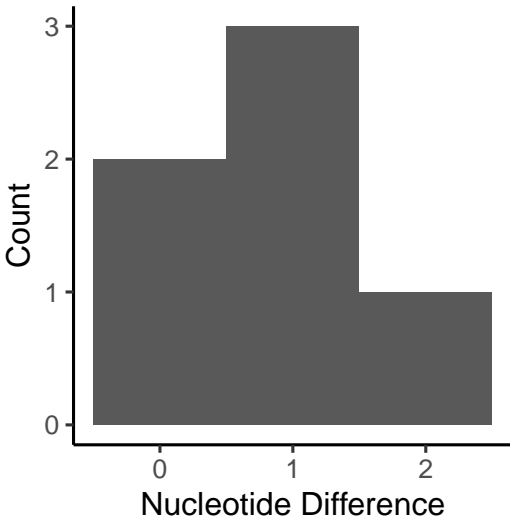
IGLV3-21*04

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



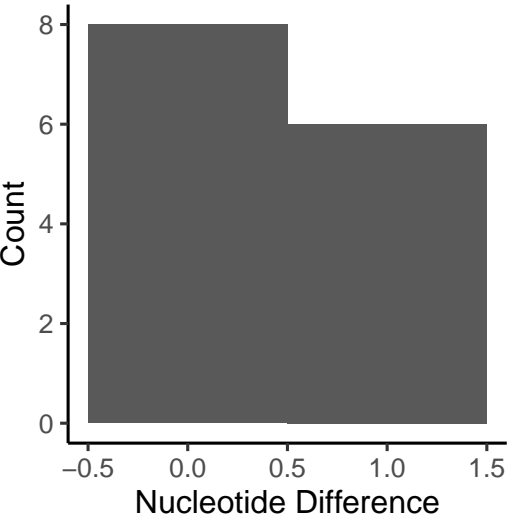
IGLV4-69*01

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



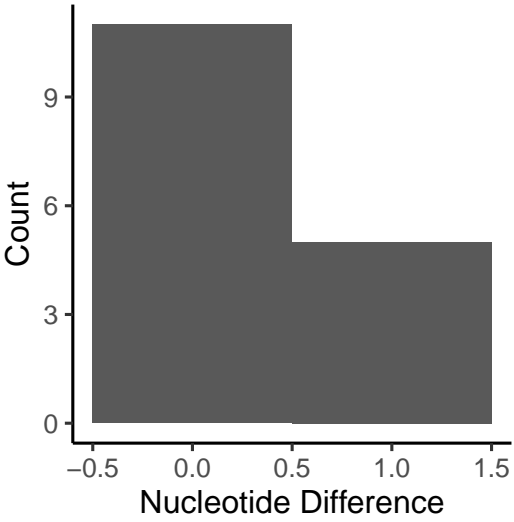
IGLV3-19*01

14 sequences assigned
8 (57.1%) exact matches, in which:
8 unique CDR3
2 unique J



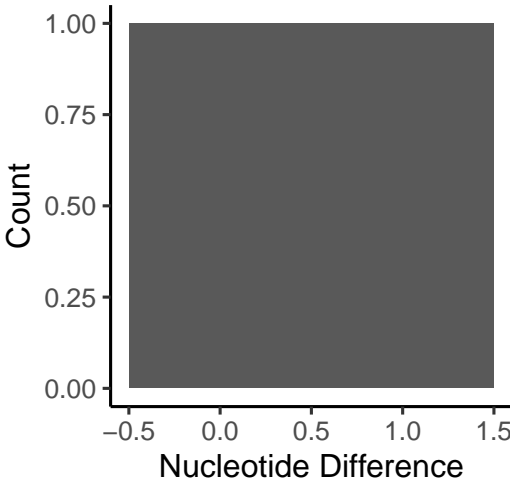
IGLV3-25*03

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



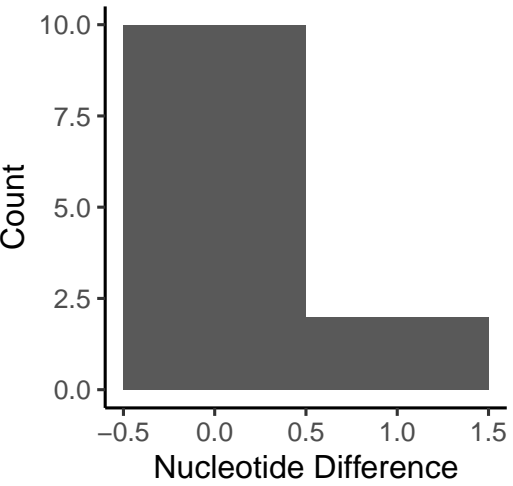
IGLV5-45*01

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



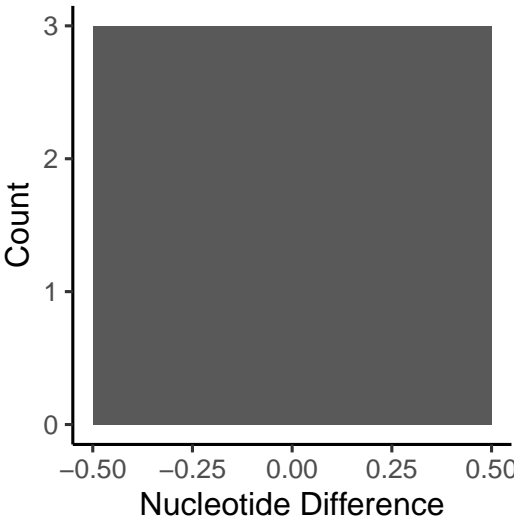
IGLV3-21*03

12 sequences assigned
10 (83.3%) exact matches, in which:
9 unique CDR3
1 unique J



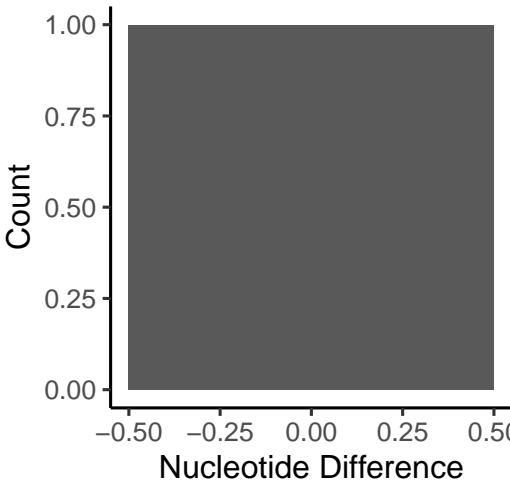
IGLV4-60*03EXT5

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



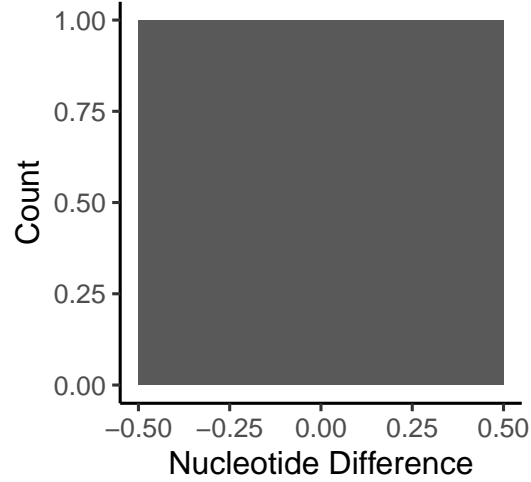
IGLV5-52*01

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



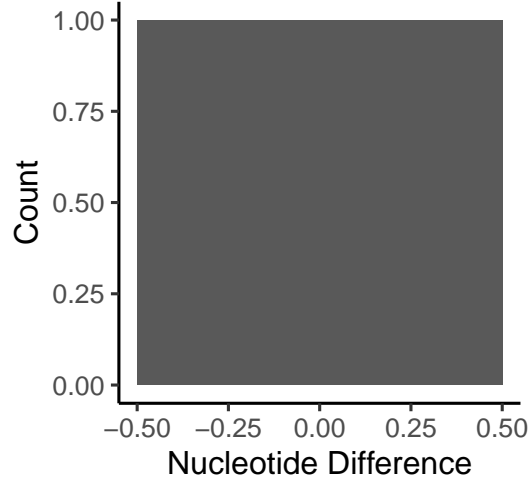
IGLV6-57*02

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



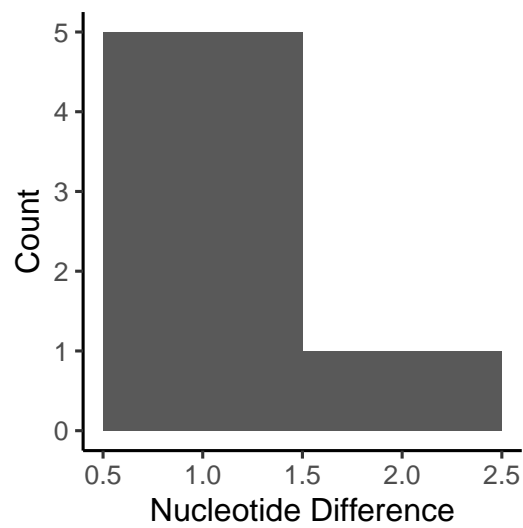
IGLV7-46*02

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



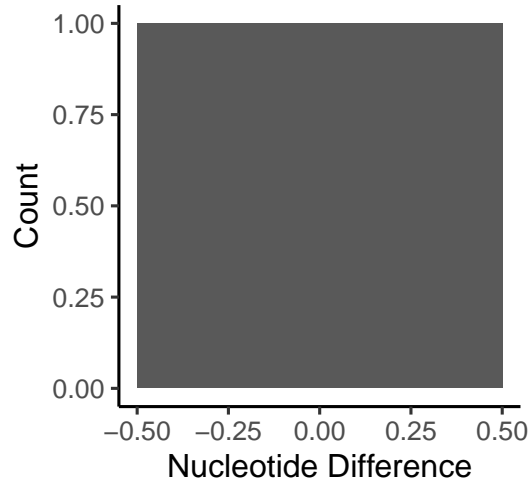
IGLV6-57*03

6 sequences assigned
No exact matches.



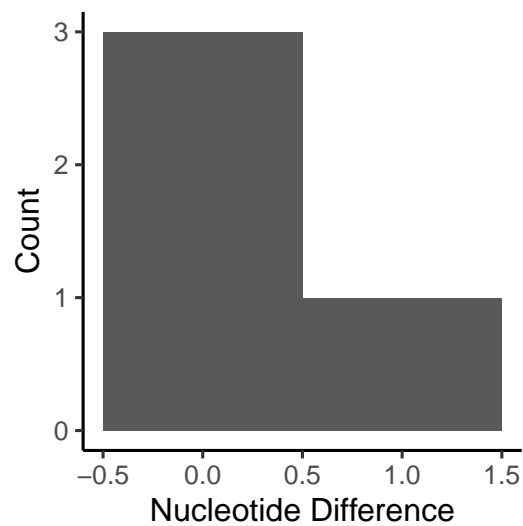
IGLV8-61*02

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



IGLV7-43*01

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



Allele Usage



Warning – no inferred sequences found.