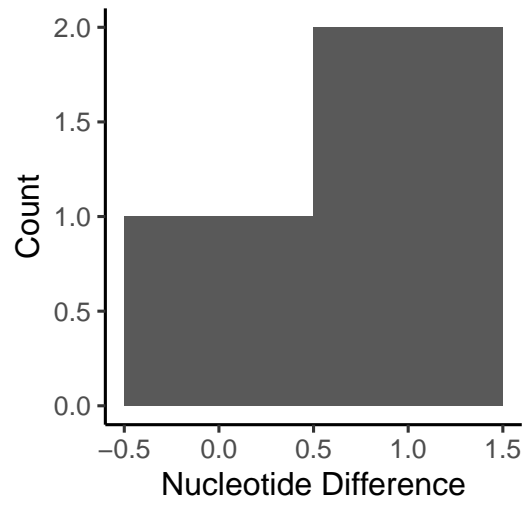


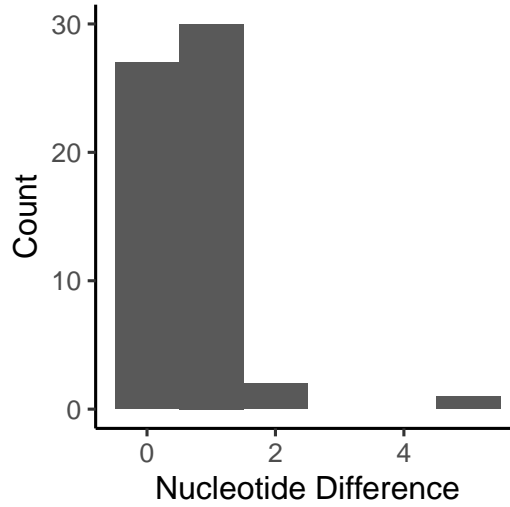
IGLV10-54*01

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



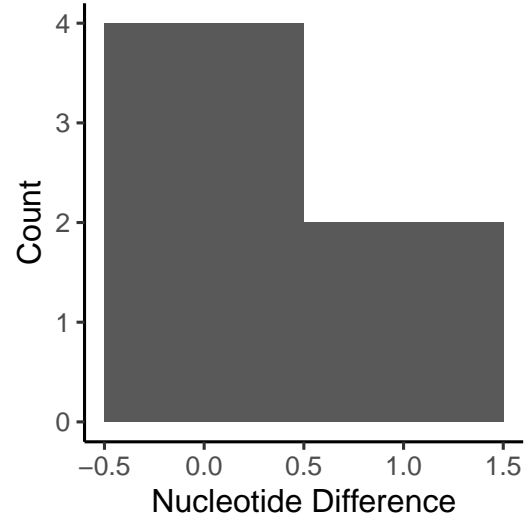
IGLV1-44*01

60 sequences assigned
27 (45%) exact matches, in which:
20 unique CDR3
3 unique J



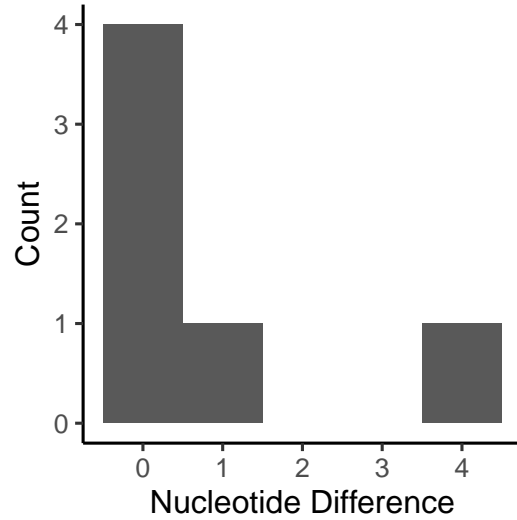
IGLV1-51*02

6 sequences assigned
4 (66.7%) exact matches, in which:
4 unique CDR3
1 unique J



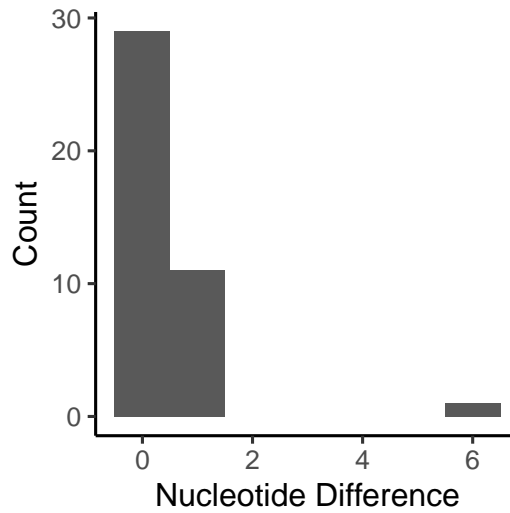
IGLV1-36*01

6 sequences assigned
4 (66.7%) exact matches, in which:
4 unique CDR3
2 unique J



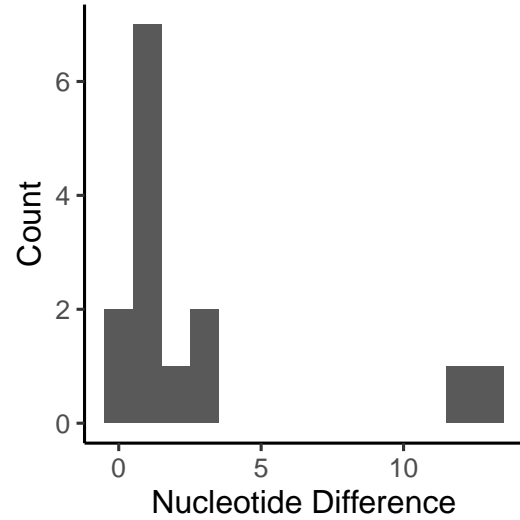
IGLV1-47*01

42 sequences assigned
29 (69%) exact matches, in which:
24 unique CDR3
2 unique J



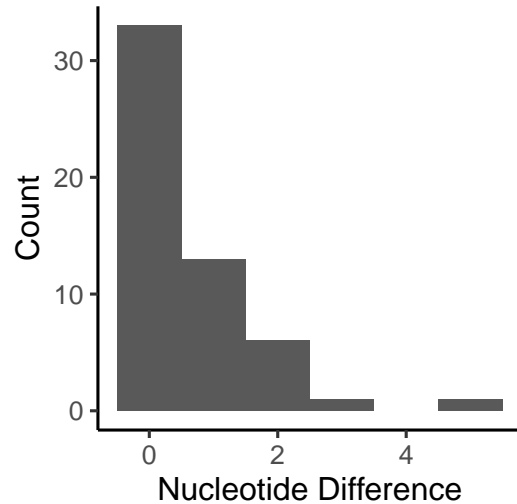
IGLV2-8*01

14 sequences assigned
2 (14.3%) exact matches, in which:
2 unique CDR3
2 unique J



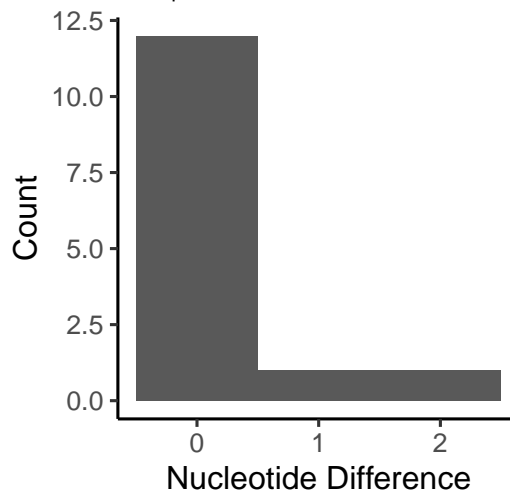
IGLV1-40*01

54 sequences assigned
33 (61.1%) exact matches, in which:
31 unique CDR3
3 unique J



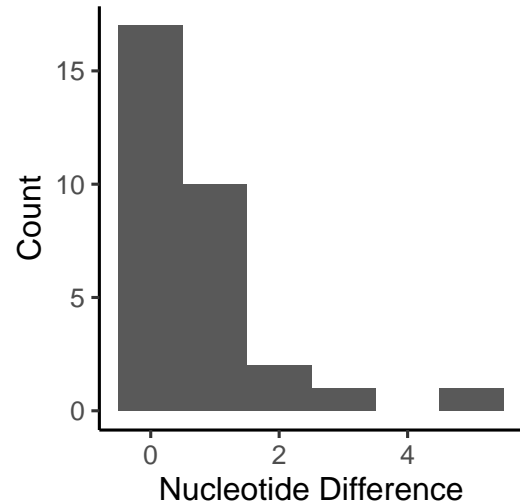
IGLV1-51*01

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



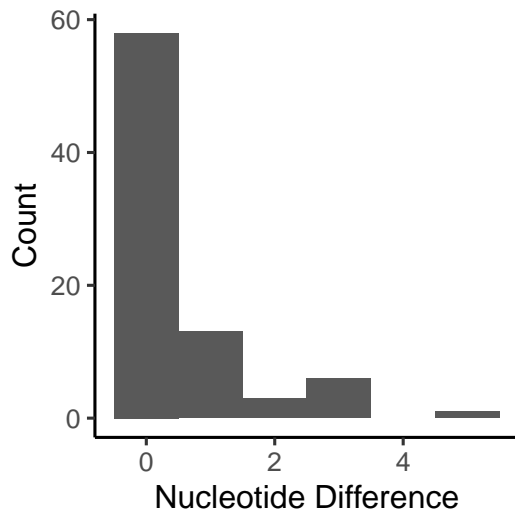
IGLV2-11*01

31 sequences assigned
17 (54.8%) exact matches, in which:
16 unique CDR3
2 unique J



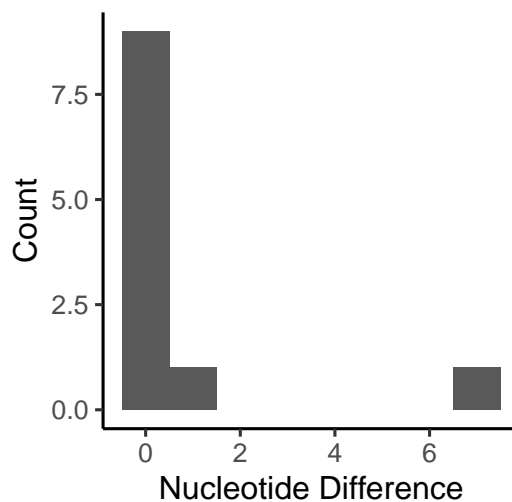
IGLV2-14*03EXT3

81 sequences assigned
58 (71.6%) exact matches, in which:
55 unique CDR3
2 unique J



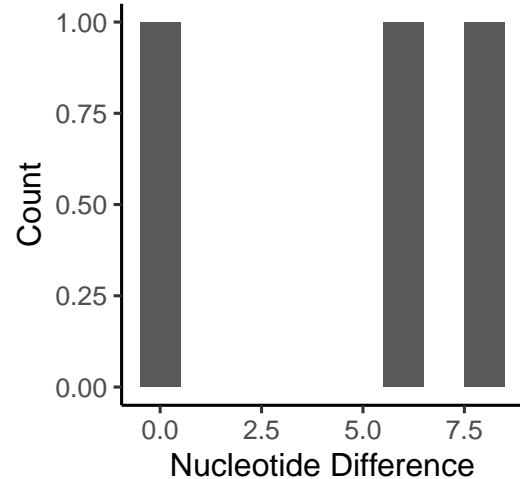
IGLV2-23*02EXT5

11 sequences assigned
9 (81.8%) exact matches, in which:
7 unique CDR3
2 unique J



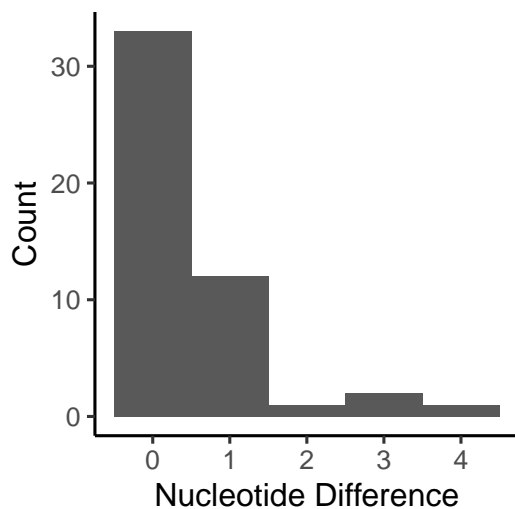
IGLV3-9*01EXT5

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



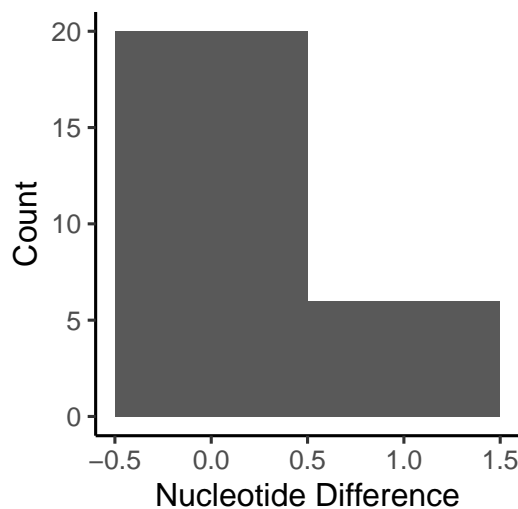
IGLV2-14*04EXT3

49 sequences assigned
33 (67.3%) exact matches, in which:
30 unique CDR3
2 unique J



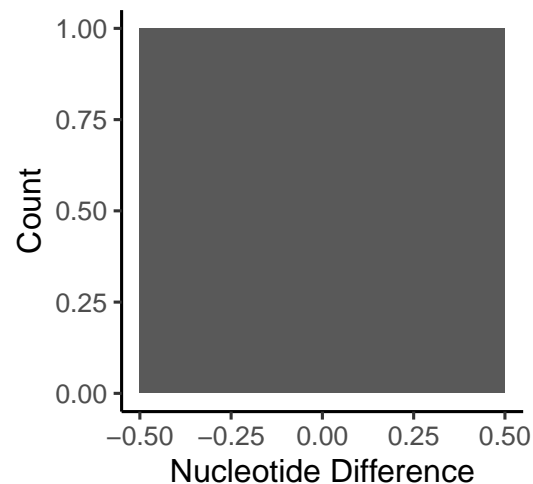
IGLV2-23*03EXT5

26 sequences assigned
20 (76.9%) exact matches, in which:
18 unique CDR3
2 unique J



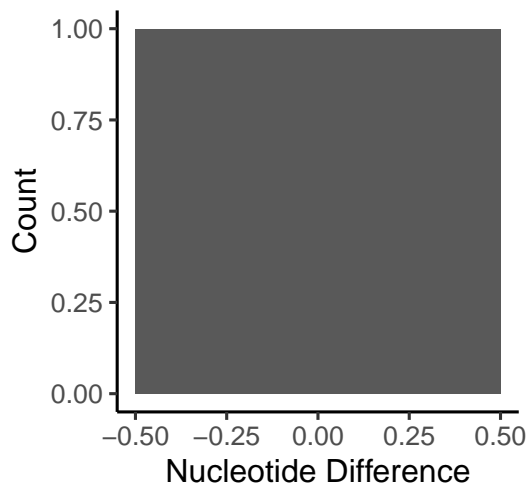
IGLV3-10*01

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



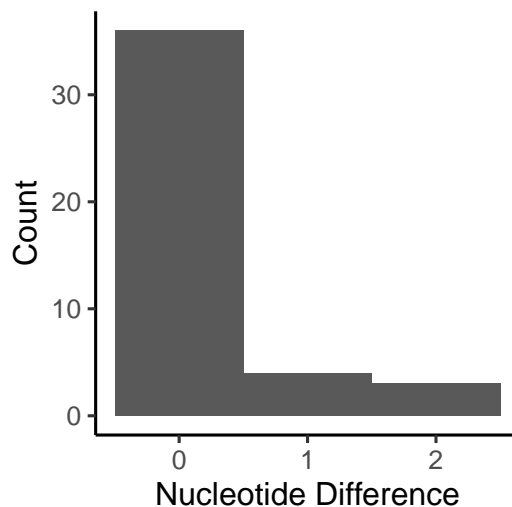
IGLV2-18*02

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



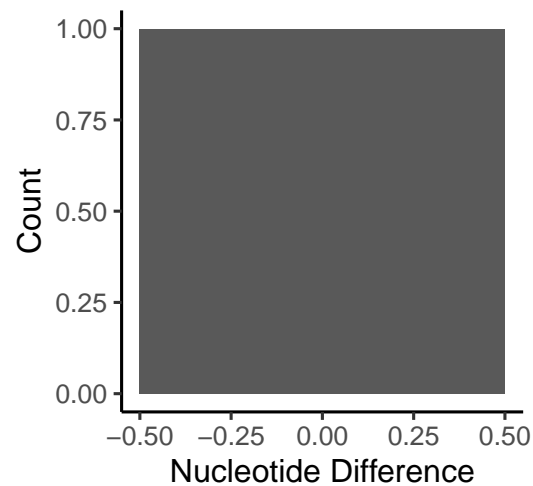
IGLV3-1*01

43 sequences assigned
36 (83.7%) exact matches, in which:
34 unique CDR3
2 unique J



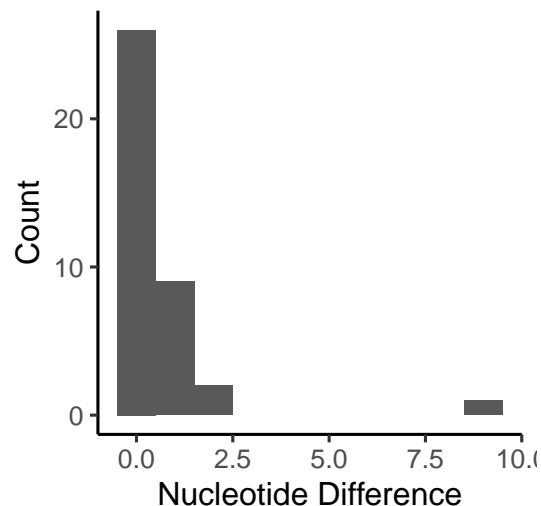
IGLV3-16*01

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



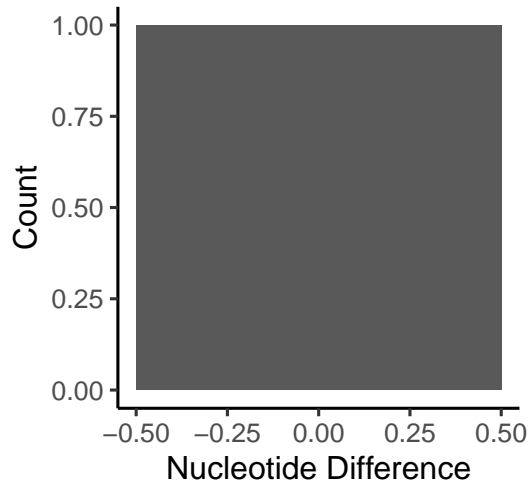
IGLV3-19*01

38 sequences assigned
26 (68.4%) exact matches, in which:
22 unique CDR3
2 unique J



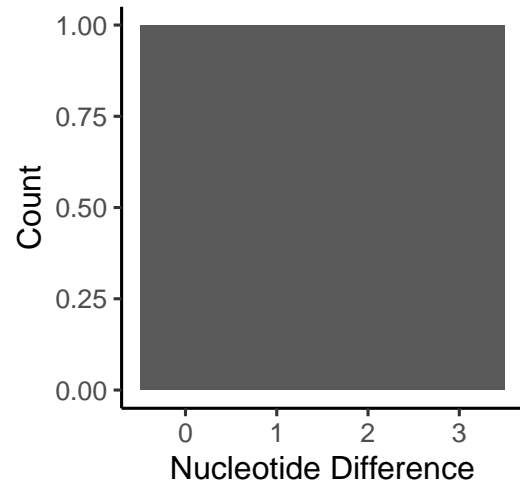
IGLV3-27*01

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



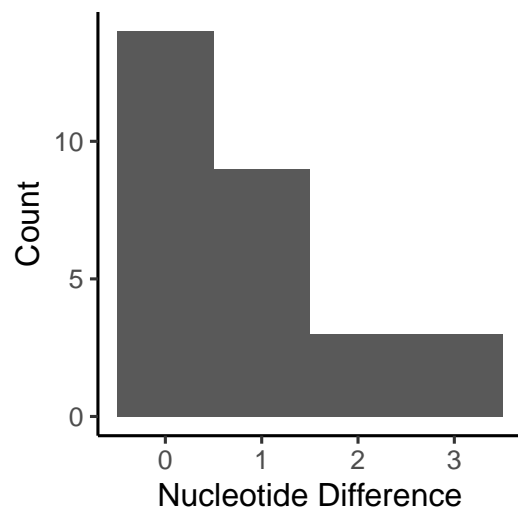
IGLV5-45*02

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



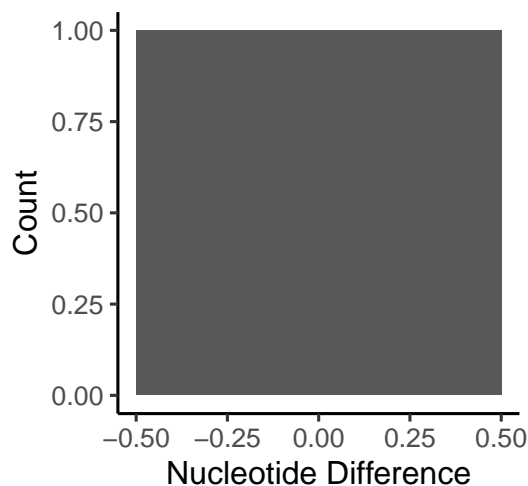
IGLV3-21*04

29 sequences assigned
14 (48.3%) exact matches, in which:
12 unique CDR3
2 unique J



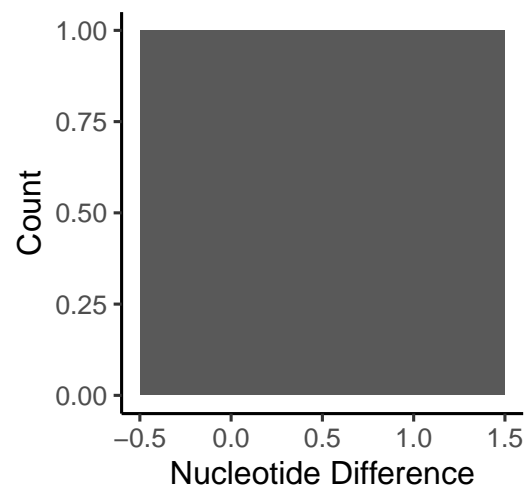
IGLV4-60*02

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



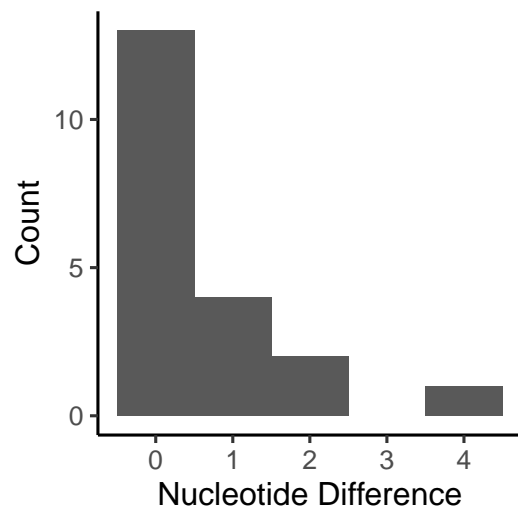
IGLV5-45*03

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



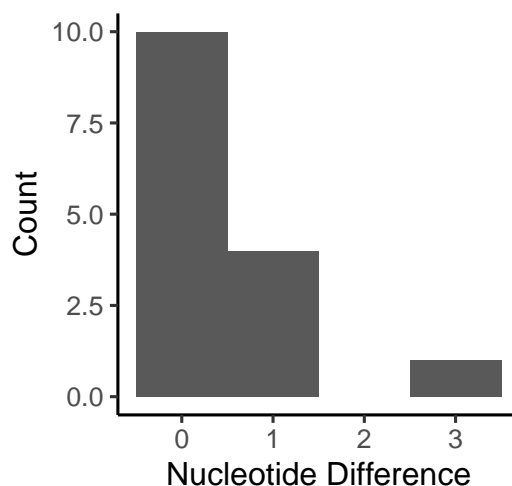
IGLV3-25*03

20 sequences assigned
13 (65%) exact matches, in which:
11 unique CDR3
2 unique J



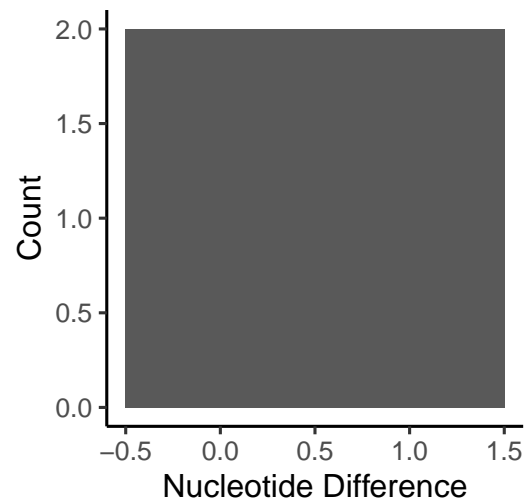
IGLV4-69*01

15 sequences assigned
10 (66.7%) exact matches, in which:
10 unique CDR3
1 unique J



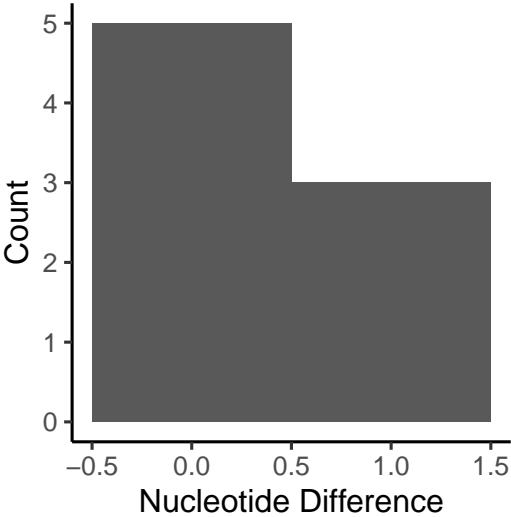
IGLV6-57*01

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



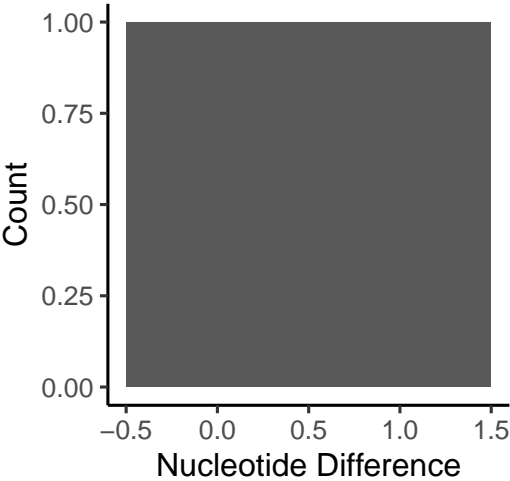
IGLV6-57*02

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



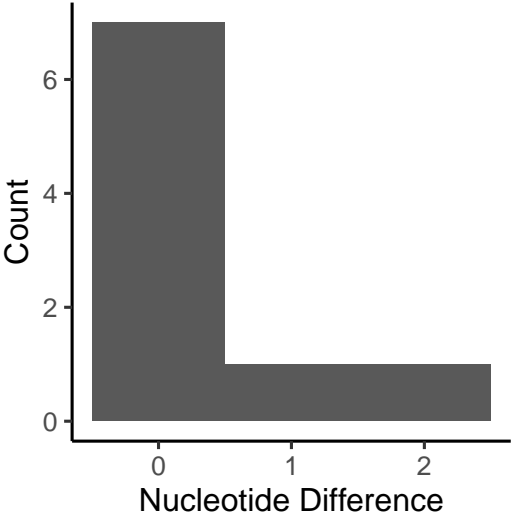
IGLV8-61*01

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



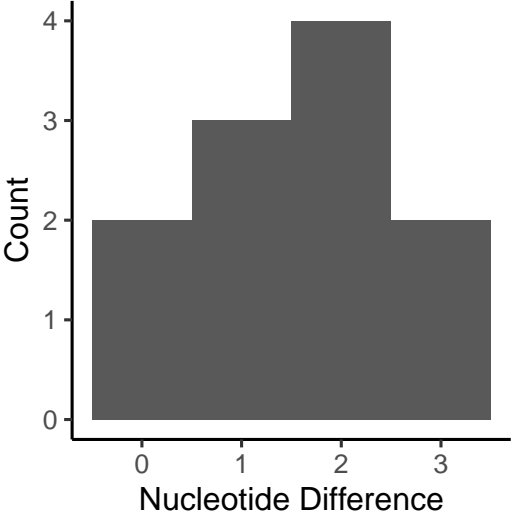
IGLV7-43*01

9 sequences assigned
7 (77.8%) exact matches, in which:
7 unique CDR3
2 unique J

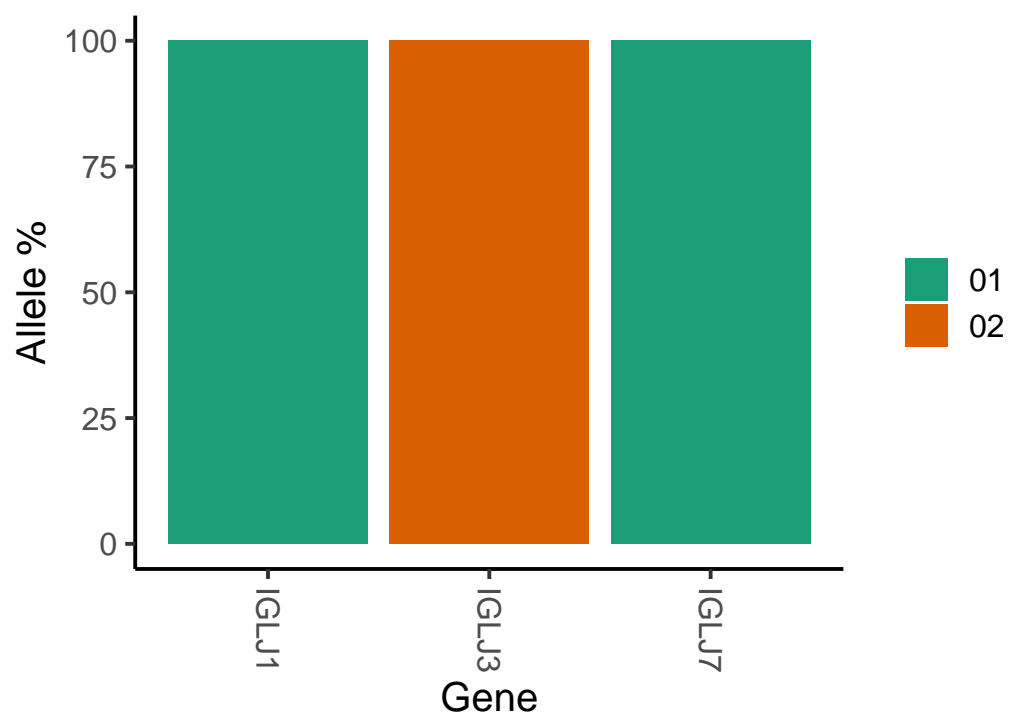


IGLV7-46*01

11 sequences assigned
2 (18.2%) exact matches, in which:
1 unique CDR3
1 unique J



Allele Usage



Warning – no inferred sequences found.