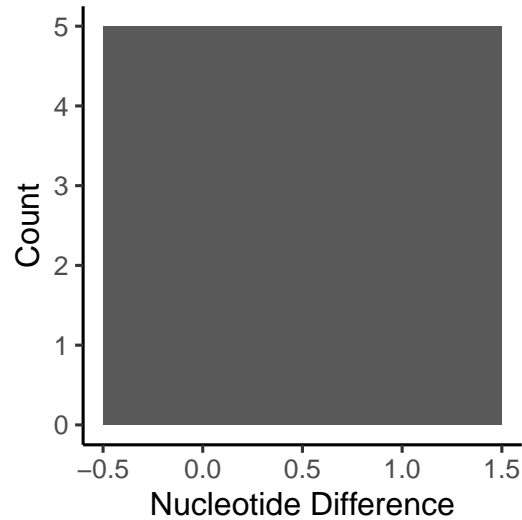


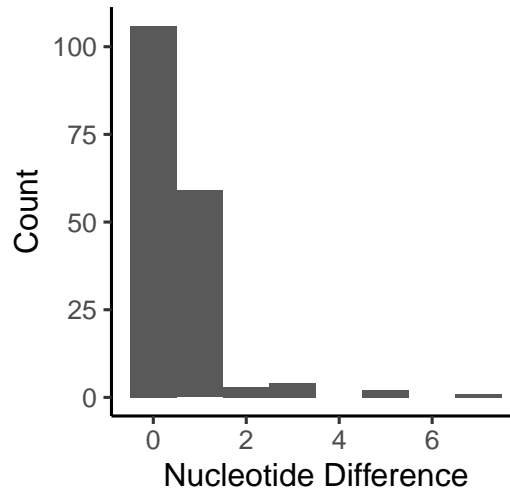
IGLV10-54*01

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



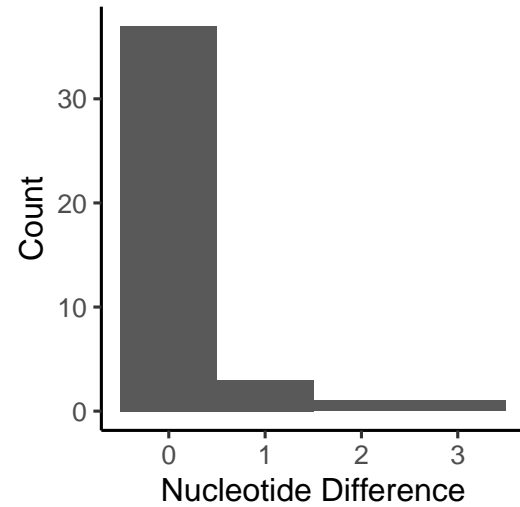
IGLV1-44*01

175 sequences assigned
106 (60.6%) exact matches, in which:
87 unique CDR3
3 unique J



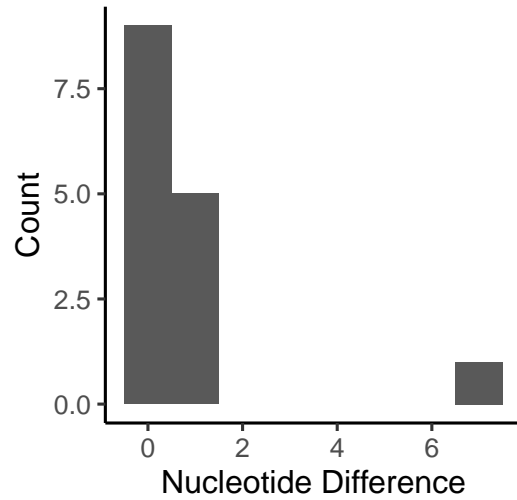
IGLV2-8*01

42 sequences assigned
37 (88.1%) exact matches, in which:
36 unique CDR3
2 unique J



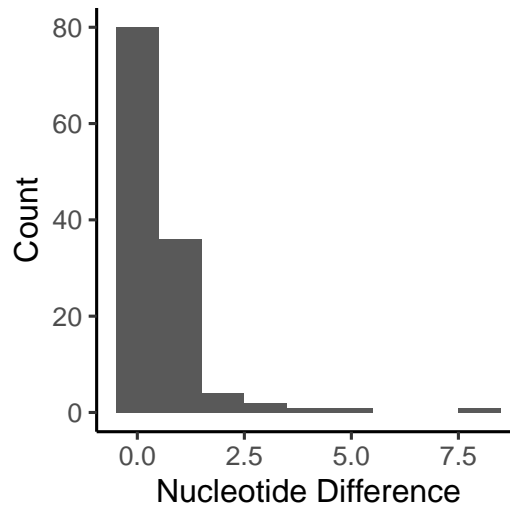
IGLV1-36*01

15 sequences assigned
9 (60%) exact matches, in which:
9 unique CDR3
2 unique J



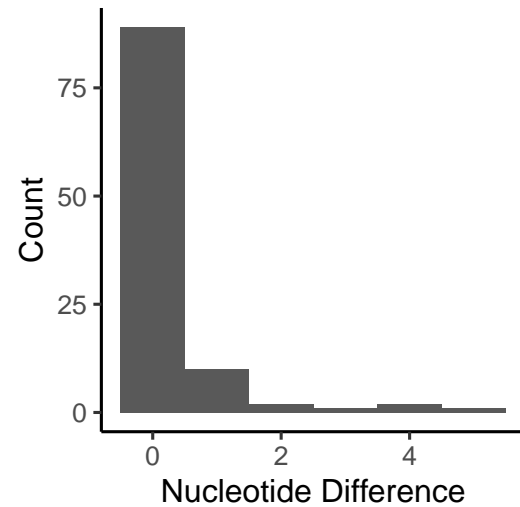
IGLV1-47*01

125 sequences assigned
80 (64%) exact matches, in which:
67 unique CDR3
2 unique J



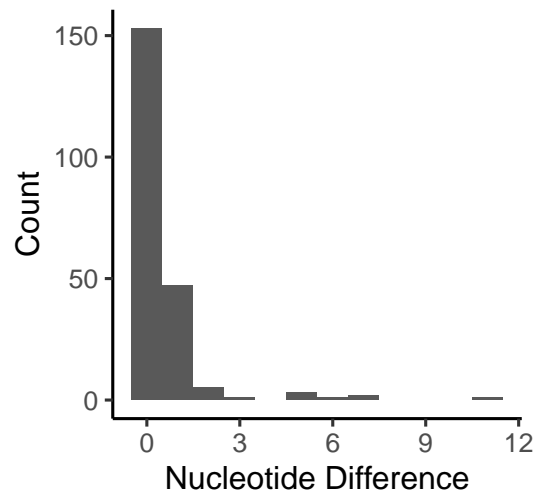
IGLV2-11*01

105 sequences assigned
89 (84.8%) exact matches, in which:
83 unique CDR3
2 unique J



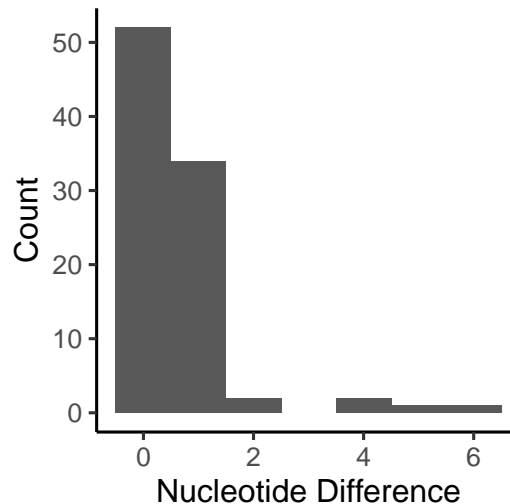
IGLV1-40*01

213 sequences assigned
153 (71.8%) exact matches, in which:
143 unique CDR3
3 unique J



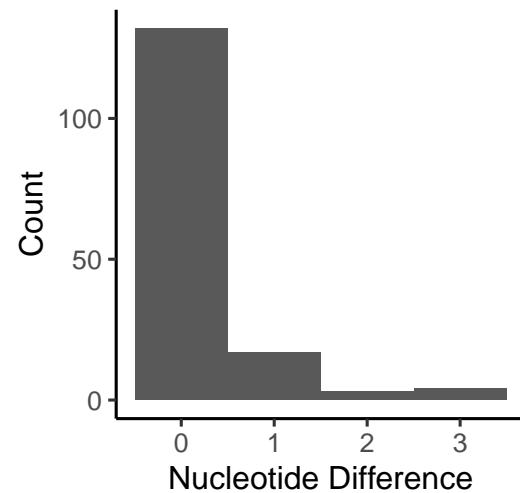
IGLV1-51*01

92 sequences assigned
52 (56.5%) exact matches, in which:
45 unique CDR3
2 unique J



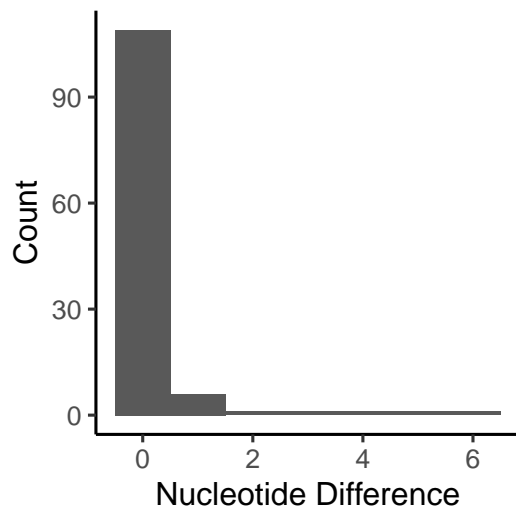
IGLV2-14*01

156 sequences assigned
132 (84.6%) exact matches, in which:
127 unique CDR3
2 unique J



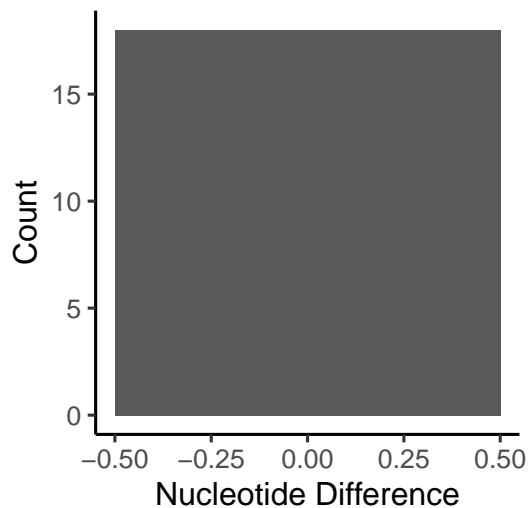
IGLV2-14*03EXT3

120 sequences assigned
109 (90.8%) exact matches, in which:
104 unique CDR3
2 unique J



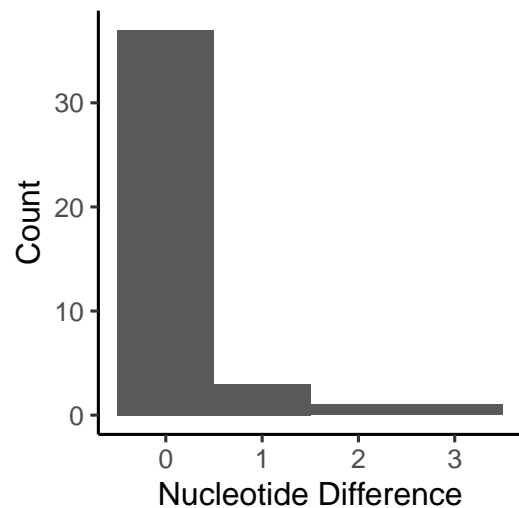
IGLV2-23*03EXT5

18 sequences assigned
18 (100%) exact matches, in which:
17 unique CDR3
2 unique J



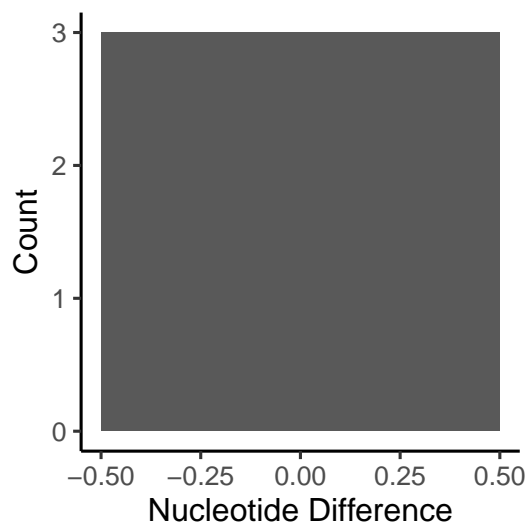
IGLV3-10*01

42 sequences assigned
37 (88.1%) exact matches, in which:
36 unique CDR3
3 unique J



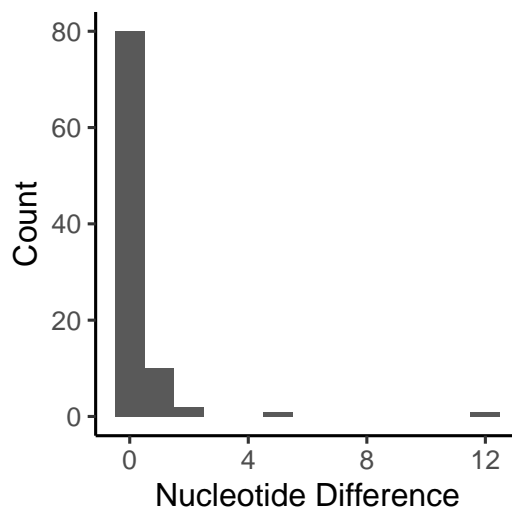
IGLV2-18*02

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



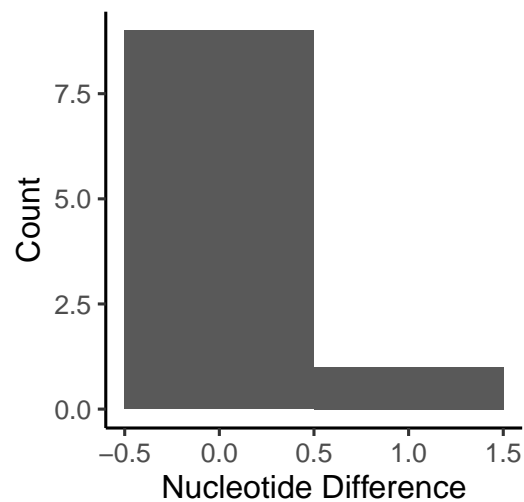
IGLV3-1*01

94 sequences assigned
80 (85.1%) exact matches, in which:
79 unique CDR3
2 unique J



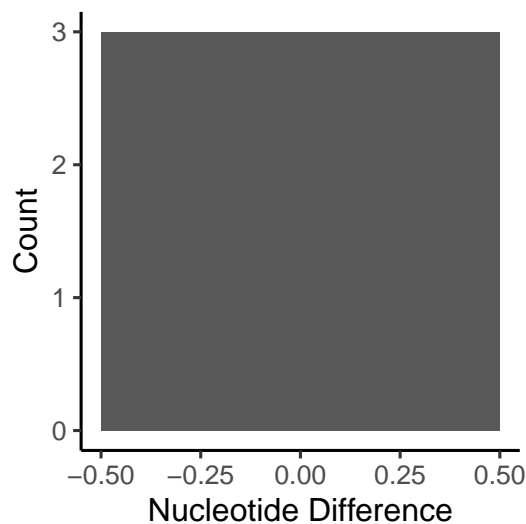
IGLV3-16*01

10 sequences assigned
9 (90%) exact matches, in which:
9 unique CDR3
1 unique J



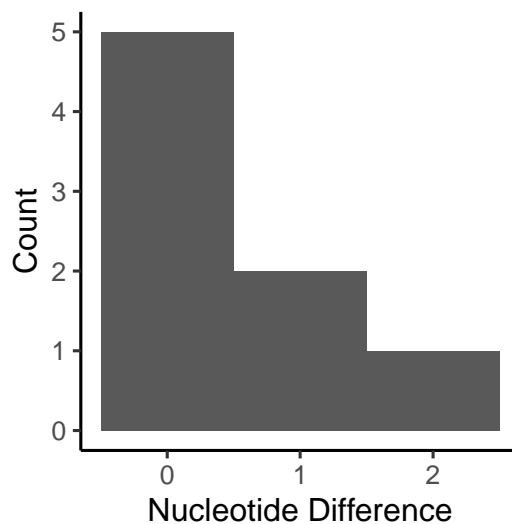
IGLV2-23*01

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



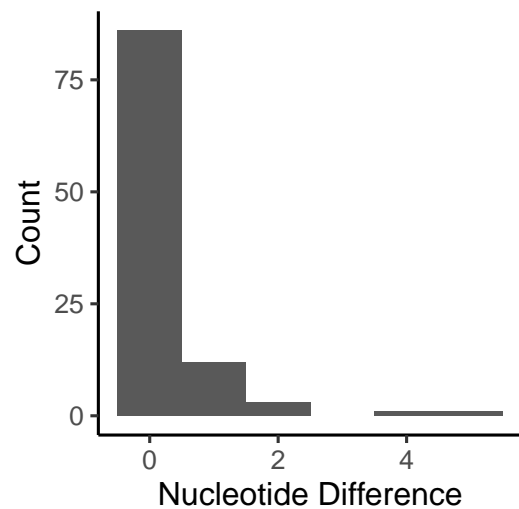
IGLV3-9*01EXT5

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



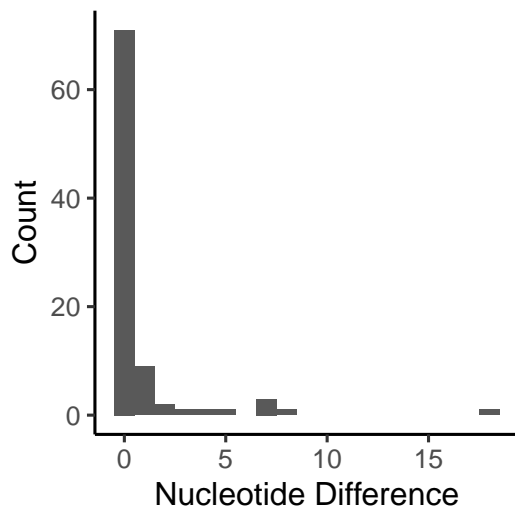
IGLV3-19*01

103 sequences assigned
86 (83.5%) exact matches, in which:
82 unique CDR3
3 unique J



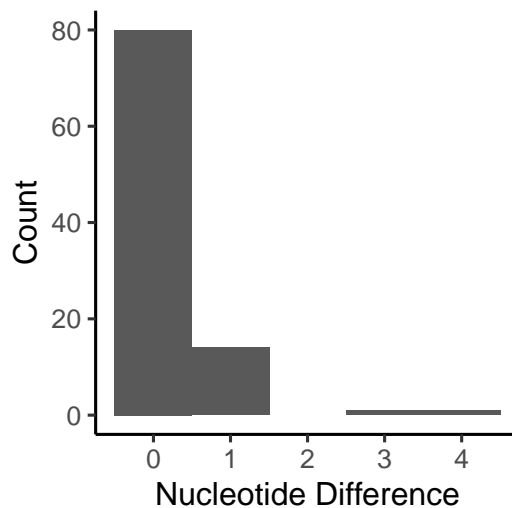
IGLV3-21*02

90 sequences assigned
71 (78.9%) exact matches, in which:
69 unique CDR3
2 unique J



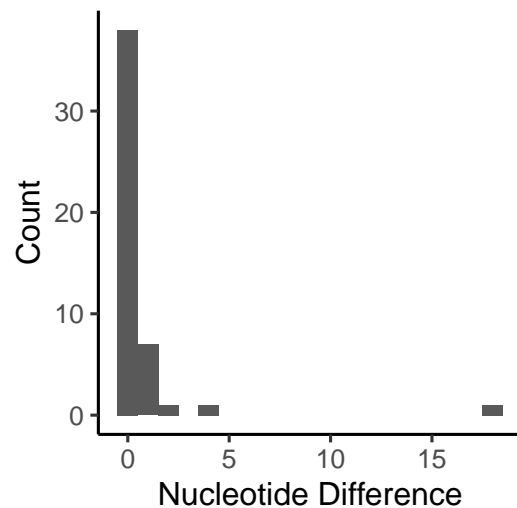
IGLV3-25*02

96 sequences assigned
80 (83.3%) exact matches, in which:
75 unique CDR3
2 unique J



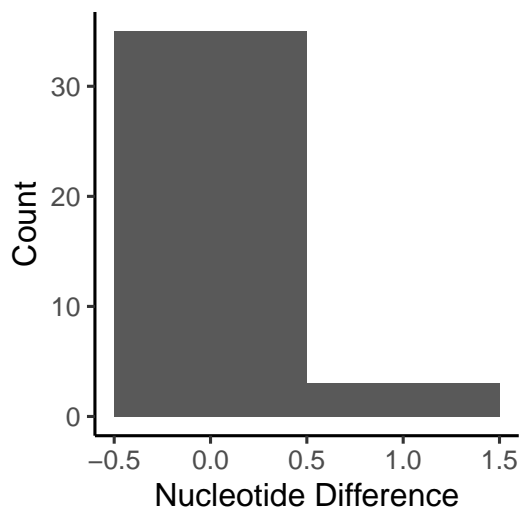
IGLV4-69*01

48 sequences assigned
38 (79.2%) exact matches, in which:
32 unique CDR3
2 unique J



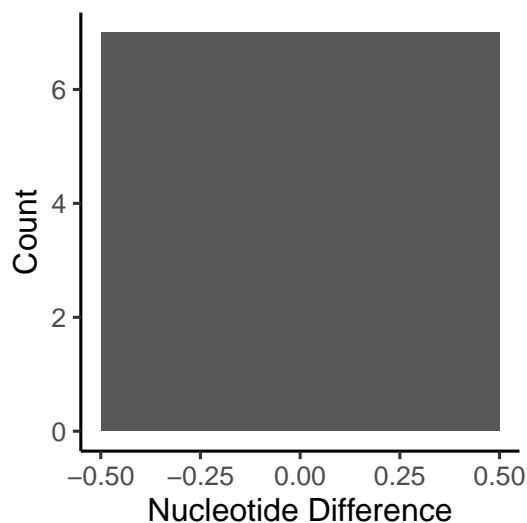
IGLV3-21*04

38 sequences assigned
35 (92.1%) exact matches, in which:
35 unique CDR3
2 unique J



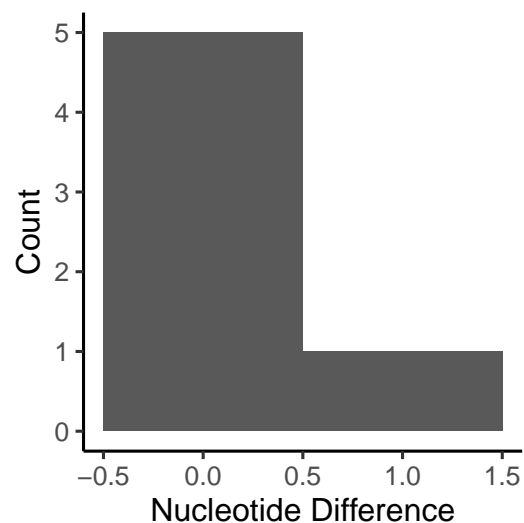
IGLV3-27*01

7 sequences assigned
7 (100%) exact matches, in which:
7 unique CDR3
1 unique J



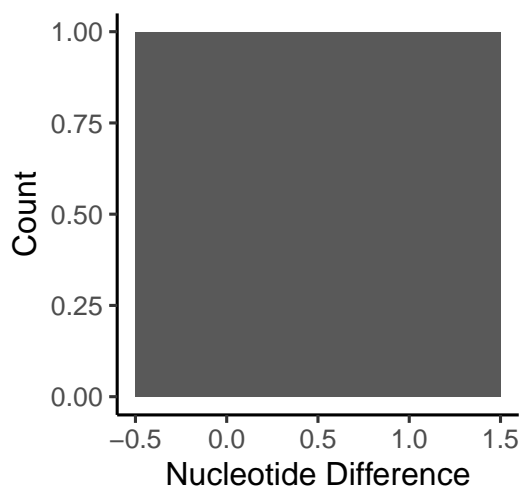
IGLV5-37*01

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



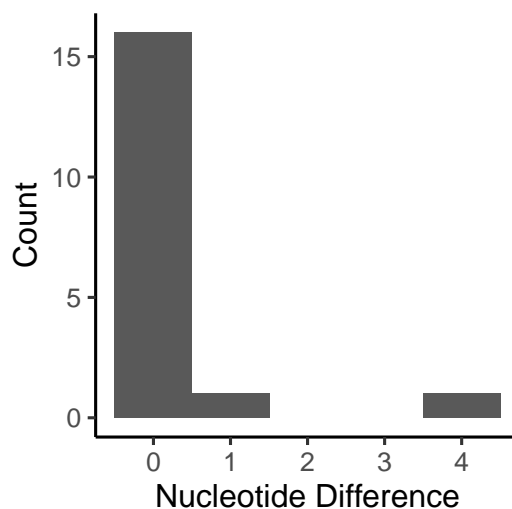
IGLV3-22*01

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



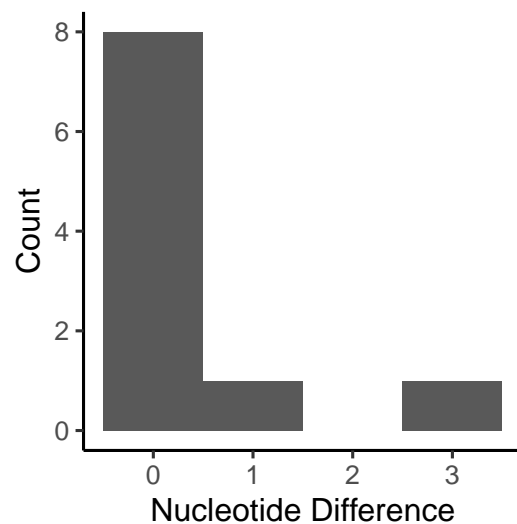
IGLV4-60*03EXT5

18 sequences assigned
16 (88.9%) exact matches, in which:
14 unique CDR3
2 unique J



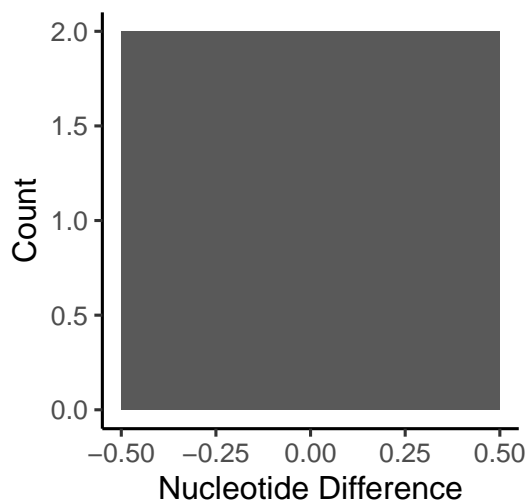
IGLV5-45*02

10 sequences assigned
8 (80%) exact matches, in which:
6 unique CDR3
3 unique J



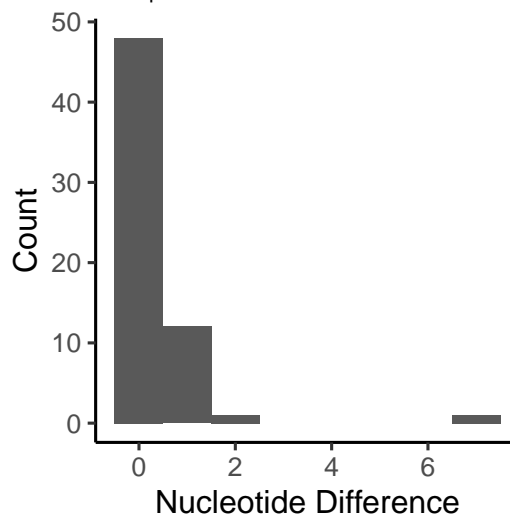
IGLV5-52*01

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



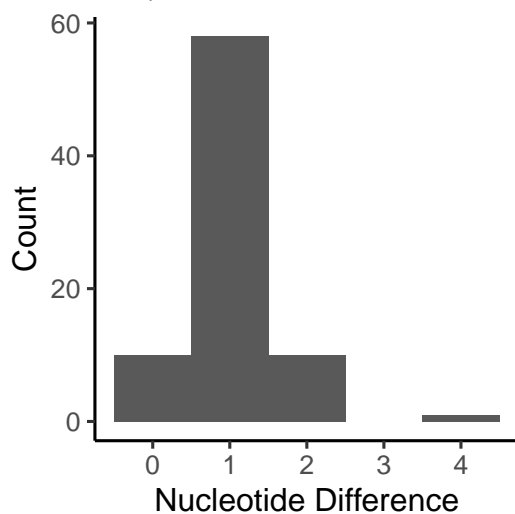
IGLV7-46*01

62 sequences assigned
48 (77.4%) exact matches, in which:
46 unique CDR3
2 unique J



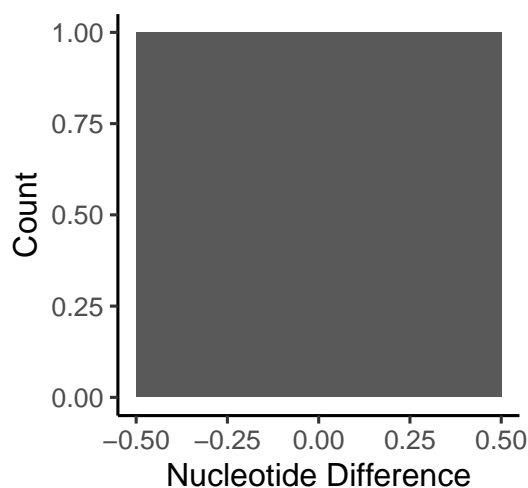
IGLV6-57*03

79 sequences assigned
10 (12.7%) exact matches, in which:
9 unique CDR3
1 unique J



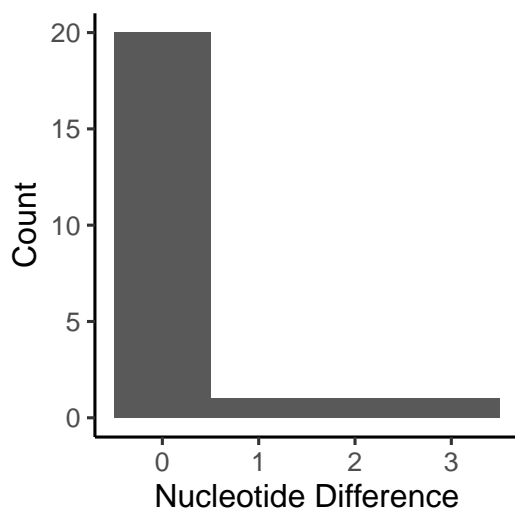
IGLV8-61*02

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



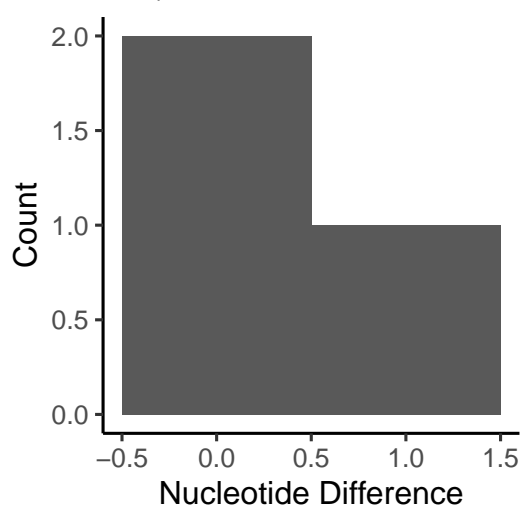
IGLV7-43*01

23 sequences assigned
20 (87%) exact matches, in which:
19 unique CDR3
2 unique J

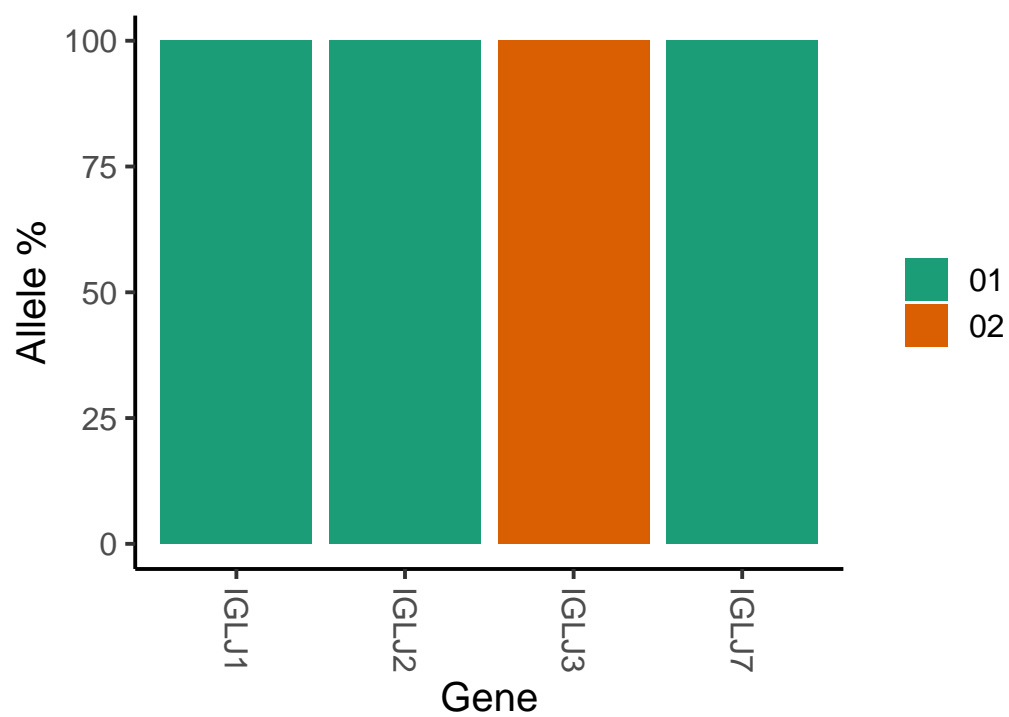


IGLV9-49*01

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



Allele Usage



Warning – no inferred sequences found.