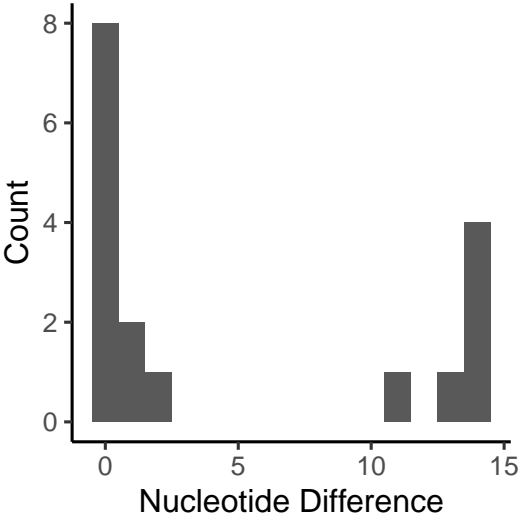


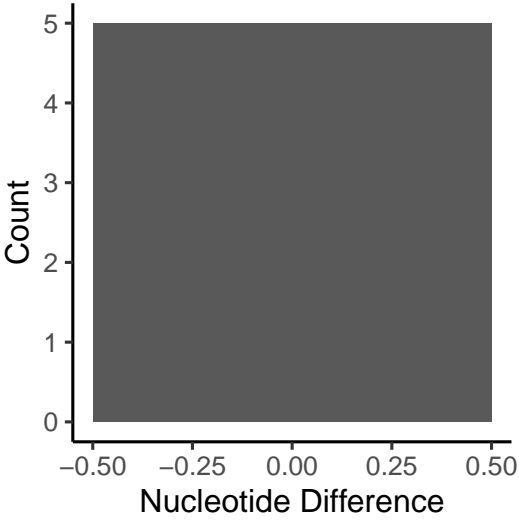
IGLV10–54*01

17 sequences assigned
8 (47.1%) exact matches, in which:
5 unique CDR3
1 unique J



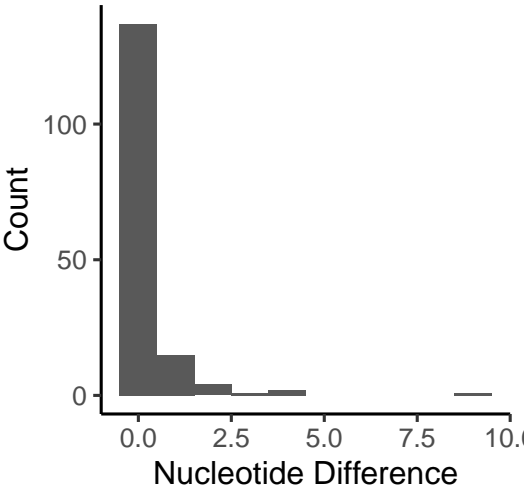
IGLV1–41*01

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



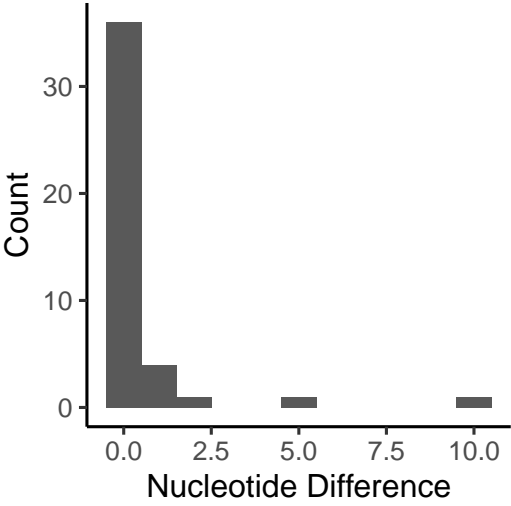
IGLV1–47*02

160 sequences assigned
137 (85.6%) exact matches, in which:
109 unique CDR3
3 unique J



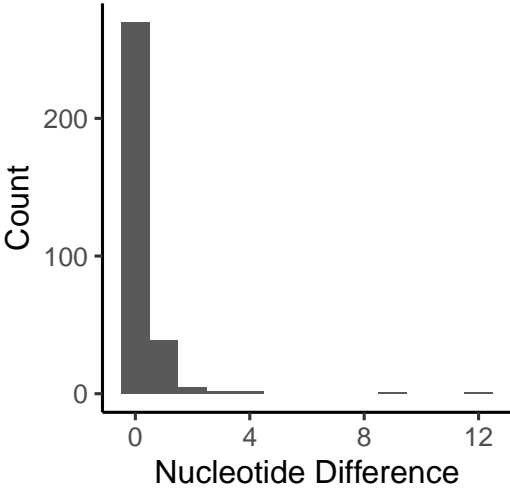
IGLV1–36*01

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



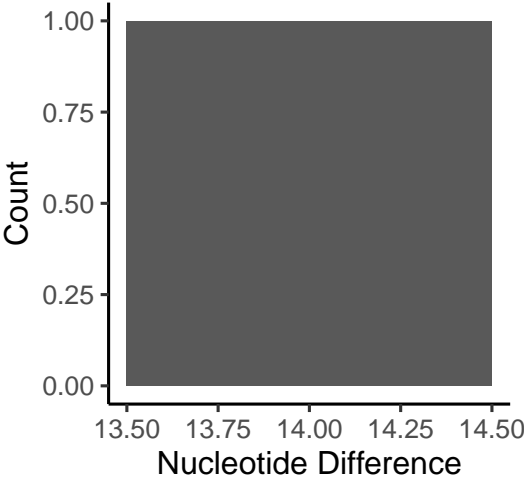
IGLV1–44*01

320 sequences assigned
270 (84.4%) exact matches, in which:
199 unique CDR3
4 unique J



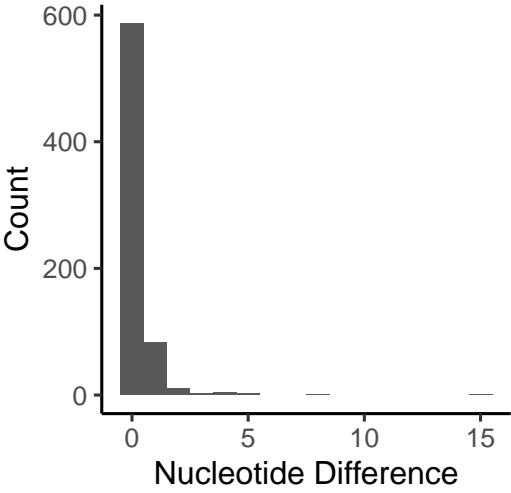
IGLV1–50*01

1 sequences assigned
No exact matches.



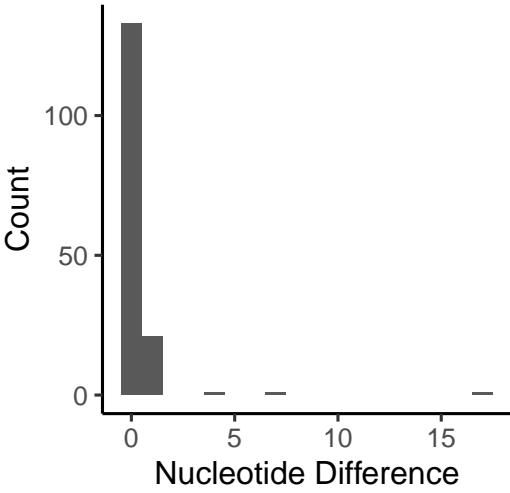
IGLV1–40*01

692 sequences assigned
587 (84.8%) exact matches, in which:
451 unique CDR3
5 unique J



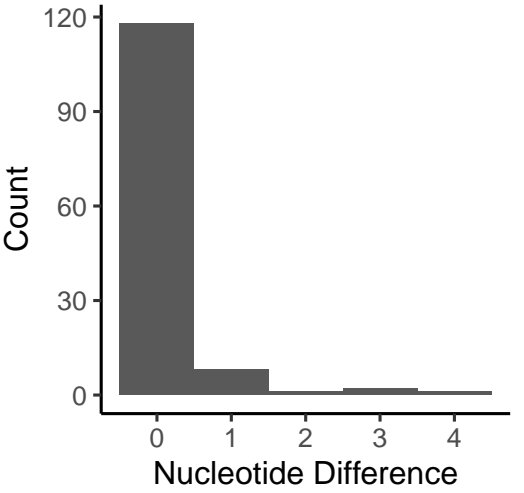
IGLV1–47*01

157 sequences assigned
133 (84.7%) exact matches, in which:
110 unique CDR3
4 unique J



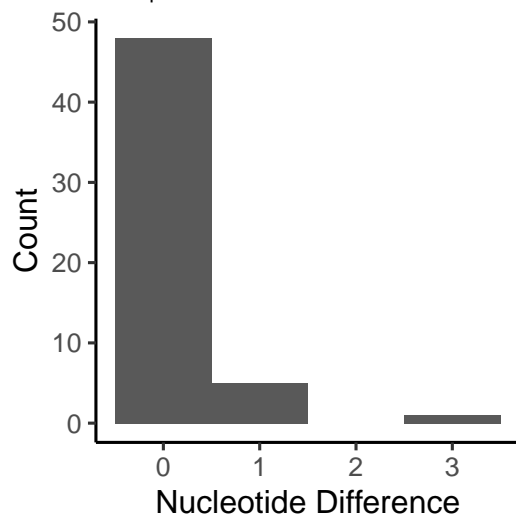
IGLV1–51*01

130 sequences assigned
118 (90.8%) exact matches, in which:
84 unique CDR3
3 unique J



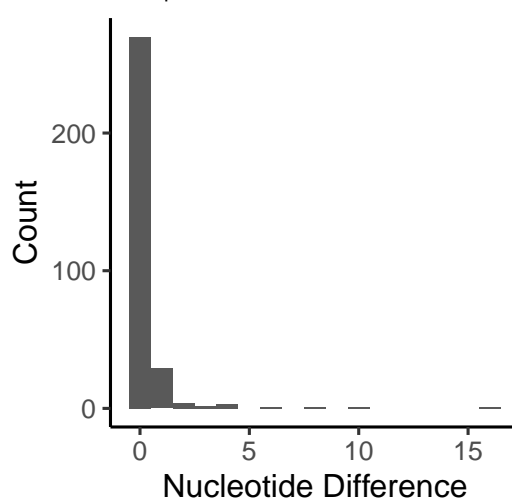
IGLV1-51*02

54 sequences assigned
48 (88.9%) exact matches, in which:
42 unique CDR3
3 unique J



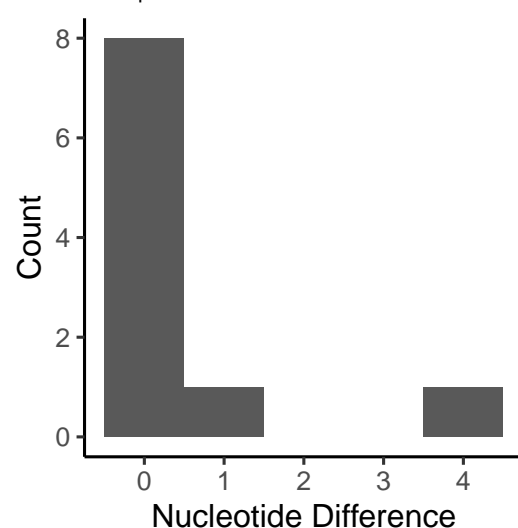
IGLV2-11*01

312 sequences assigned
270 (86.5%) exact matches, in which:
243 unique CDR3
3 unique J



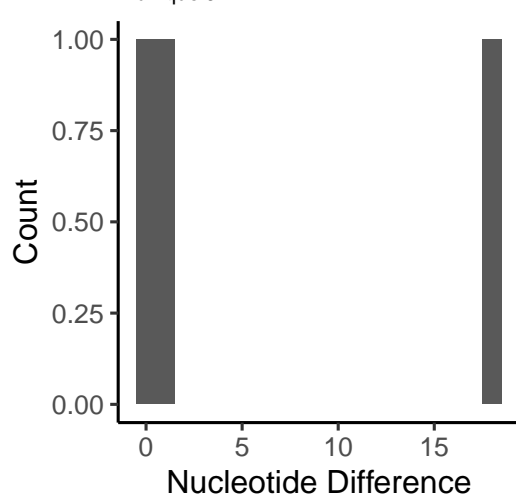
IGLV2-18*01

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



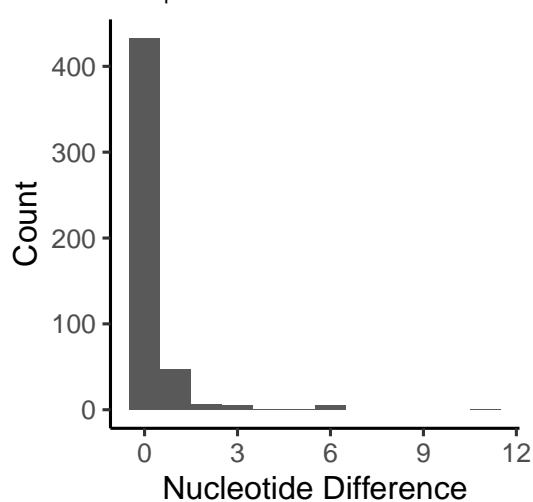
IGLV2-5*01

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



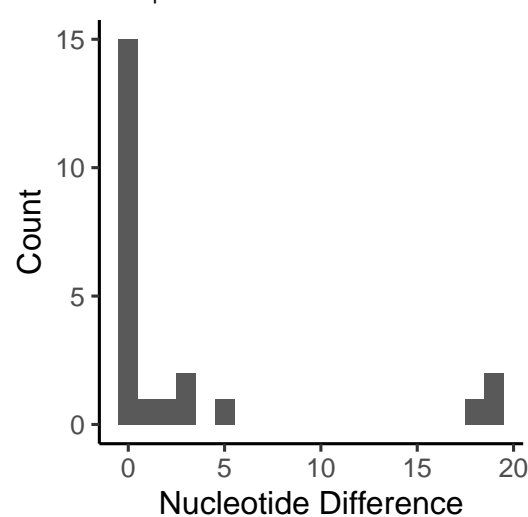
IGLV2-14*01

499 sequences assigned
433 (86.8%) exact matches, in which:
392 unique CDR3
2 unique J



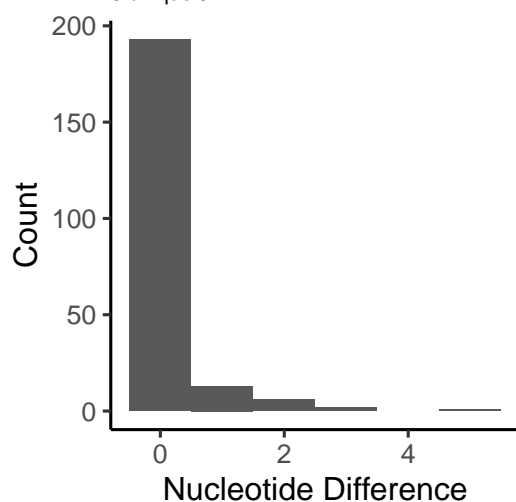
IGLV2-18*02

23 sequences assigned
15 (65.2%) exact matches, in which:
11 unique CDR3
2 unique J



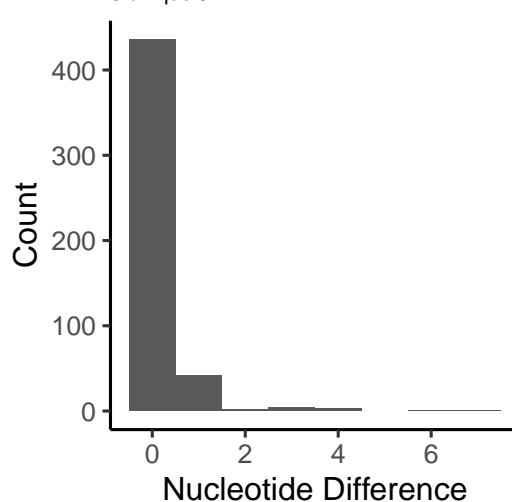
IGLV2-8*01

215 sequences assigned
193 (89.8%) exact matches, in which:
177 unique CDR3
3 unique J



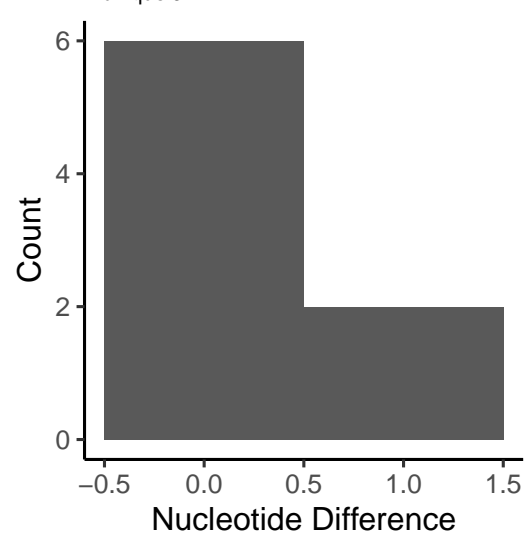
IGLV2-14*03EXT3

489 sequences assigned
436 (89.2%) exact matches, in which:
375 unique CDR3
3 unique J



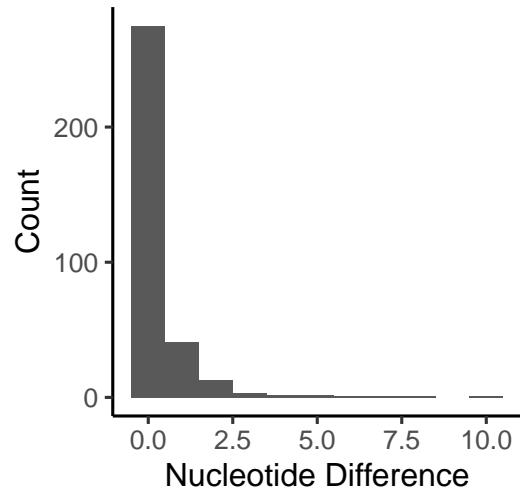
IGLV2-23*01

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



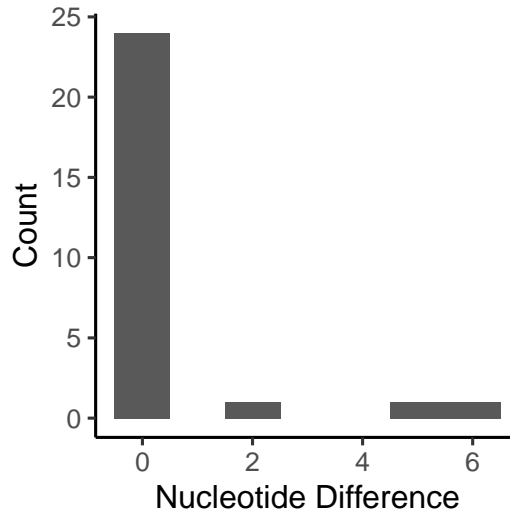
IGLV2-23*02EXT5

340 sequences assigned
275 (80.9%) exact matches, in which:
256 unique CDR3
4 unique J



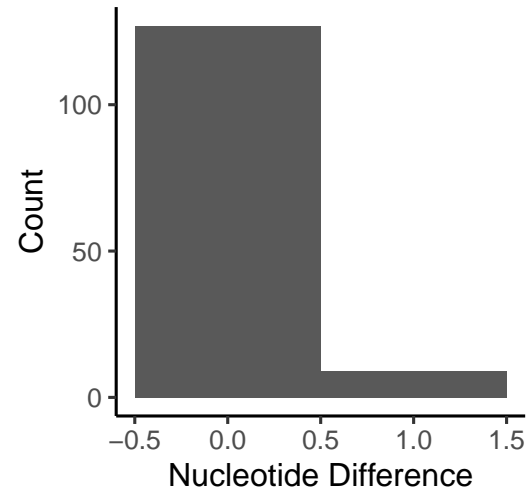
IGLV3-9*01EXT5

27 sequences assigned
24 (88.9%) exact matches, in which:
24 unique CDR3
2 unique J



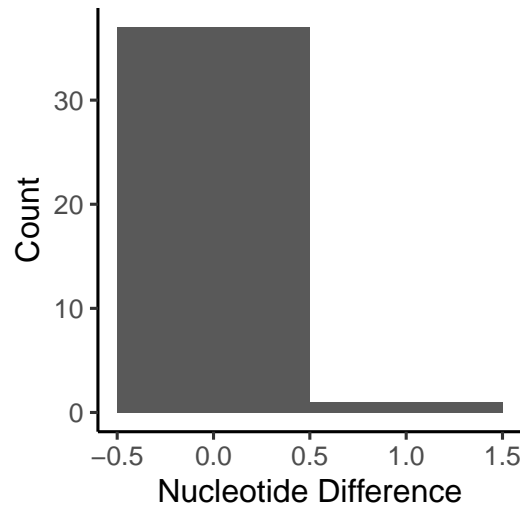
IGLV3-19*01

136 sequences assigned
127 (93.4%) exact matches, in which:
110 unique CDR3
4 unique J



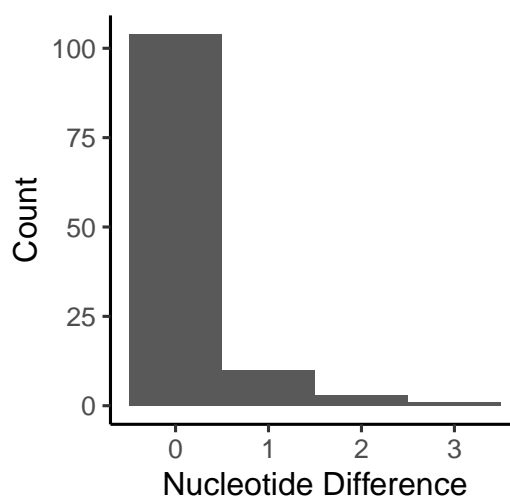
IGLV2-23*03EXT5

38 sequences assigned
37 (97.4%) exact matches, in which:
37 unique CDR3
2 unique J



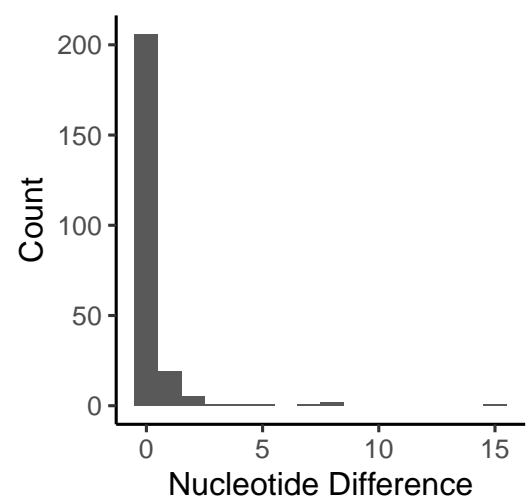
IGLV3-10*01

118 sequences assigned
104 (88.1%) exact matches, in which:
97 unique CDR3
3 unique J



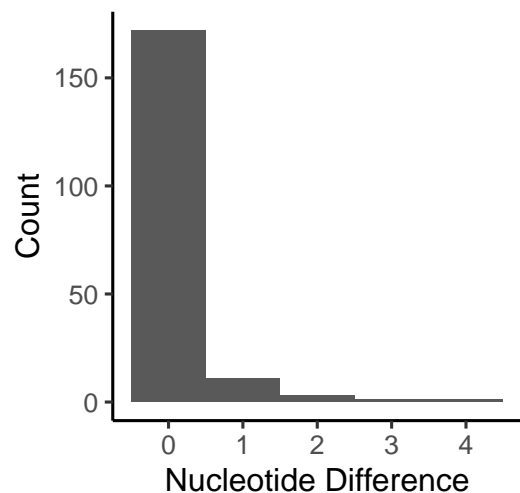
IGLV3-21*02

237 sequences assigned
206 (86.9%) exact matches, in which:
178 unique CDR3
3 unique J



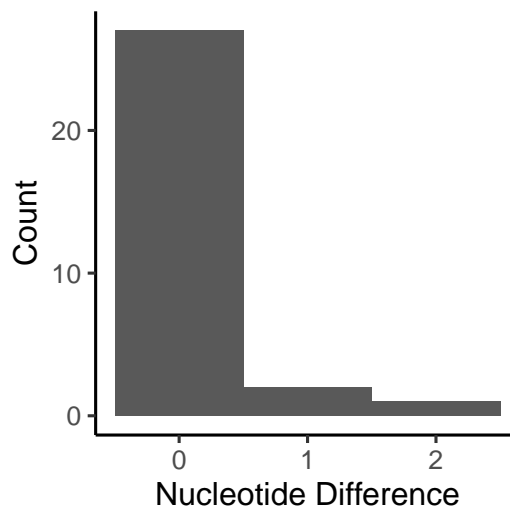
IGLV3-1*01

188 sequences assigned
172 (91.5%) exact matches, in which:
145 unique CDR3
2 unique J



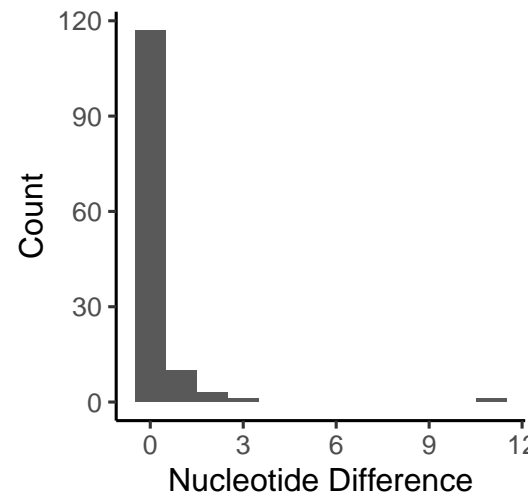
IGLV3-16*01

30 sequences assigned
27 (90%) exact matches, in which:
25 unique CDR3
3 unique J



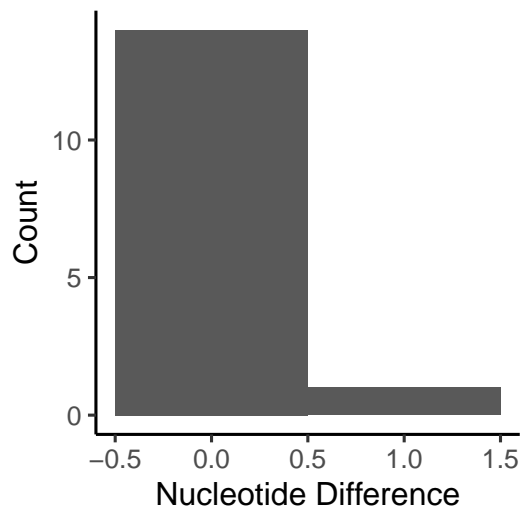
IGLV3-21*04

132 sequences assigned
117 (88.6%) exact matches, in which:
99 unique CDR3
3 unique J



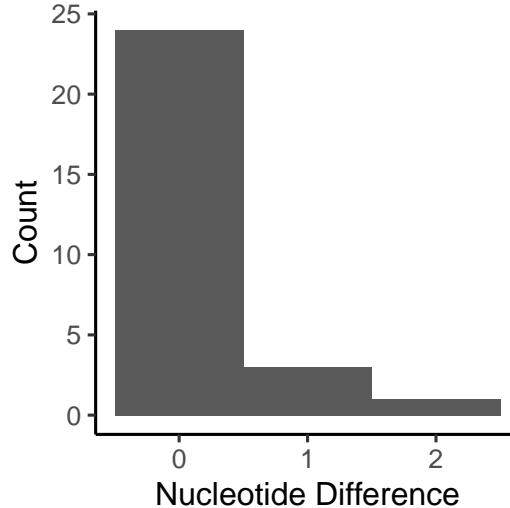
IGLV3-22*01

15 sequences assigned
14 (93.3%) exact matches, in which:
12 unique CDR3
1 unique J



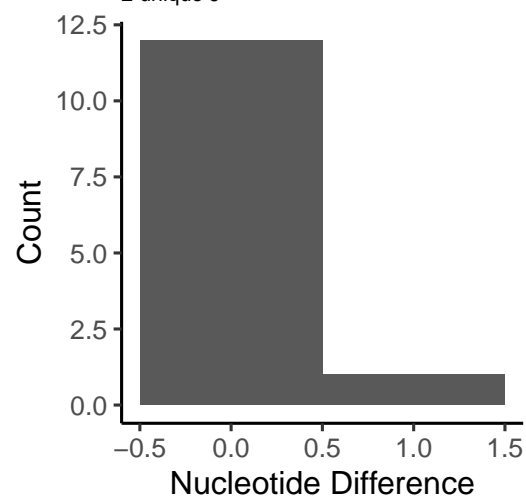
IGLV3-27*01

28 sequences assigned
24 (85.7%) exact matches, in which:
22 unique CDR3
3 unique J



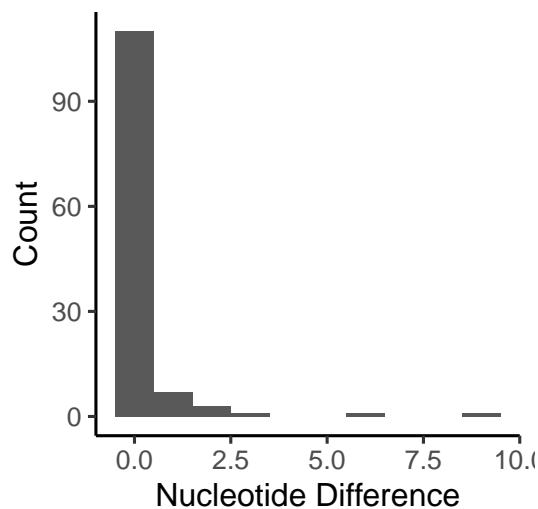
IGLV5-37*01

13 sequences assigned
12 (92.3%) exact matches, in which:
10 unique CDR3
2 unique J



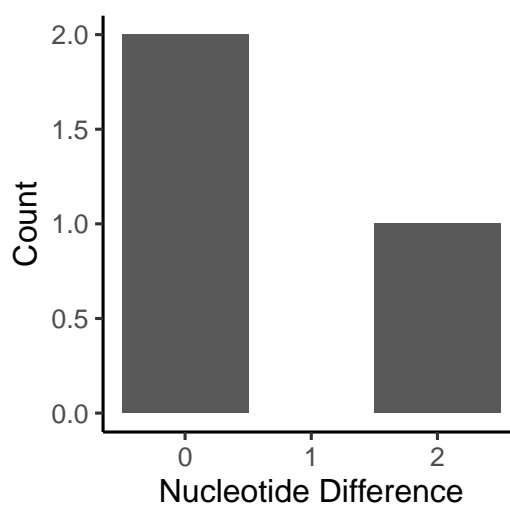
IGLV3-25*02

123 sequences assigned
110 (89.4%) exact matches, in which:
99 unique CDR3
2 unique J



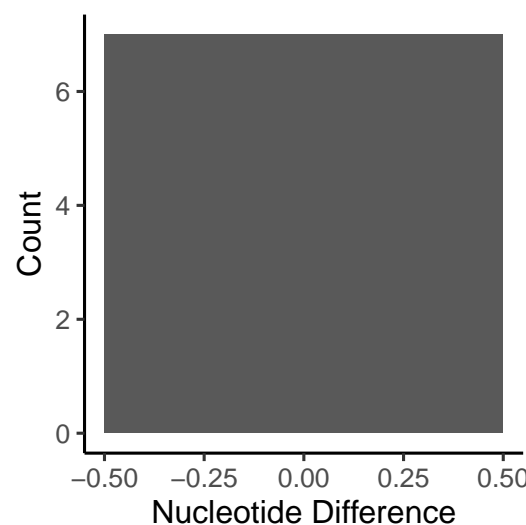
IGLV4-3*01

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



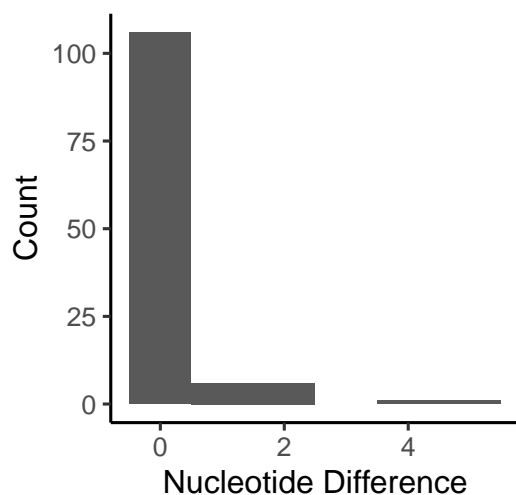
IGLV5-39*01

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



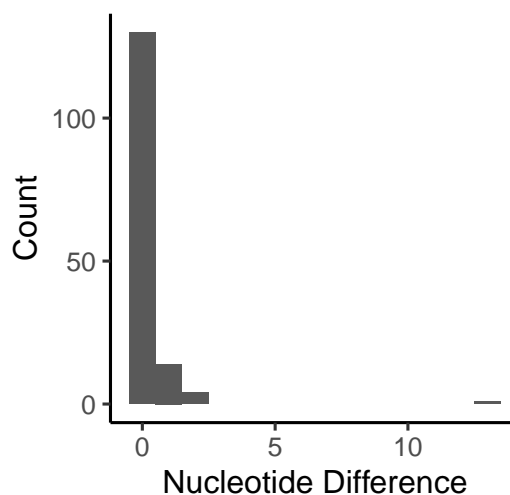
IGLV3-25*03

120 sequences assigned
106 (88.3%) exact matches, in which:
98 unique CDR3
3 unique J



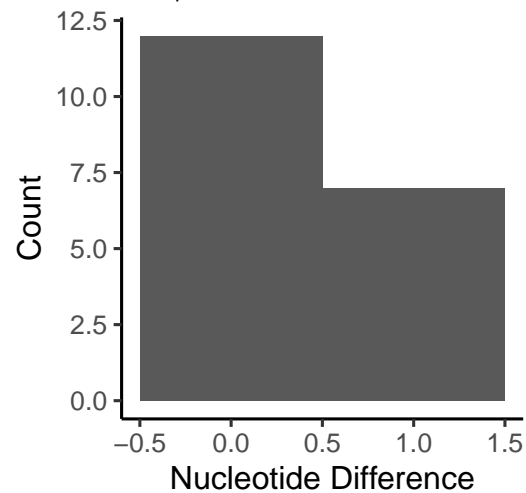
IGLV4-69*01

149 sequences assigned
130 (87.2%) exact matches, in which:
106 unique CDR3
3 unique J



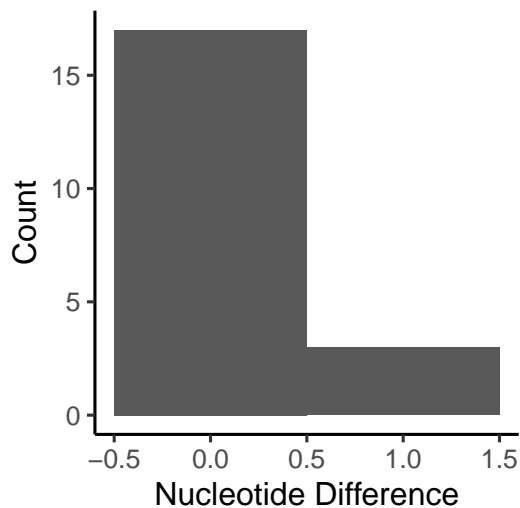
IGLV5-45*01

19 sequences assigned
12 (63.2%) exact matches, in which:
11 unique CDR3
2 unique J



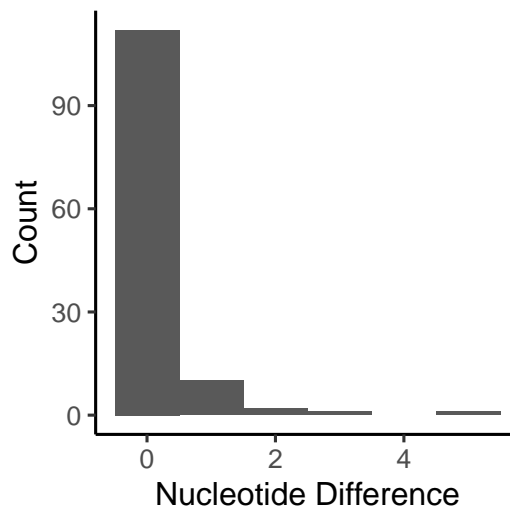
IGLV5-45*03

20 sequences assigned
17 (85%) exact matches, in which:
17 unique CDR3
5 unique J



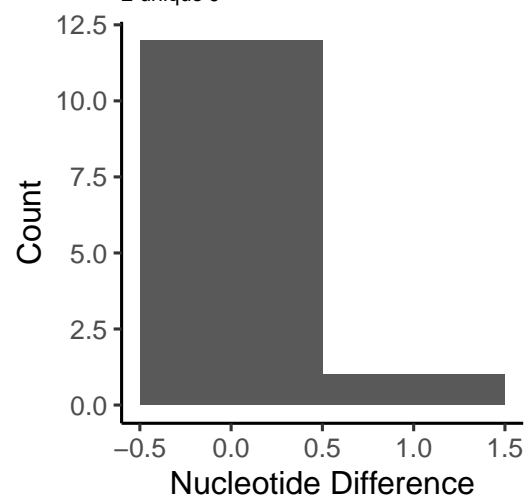
IGLV6-57*02

126 sequences assigned
112 (88.9%) exact matches, in which:
95 unique CDR3
4 unique J



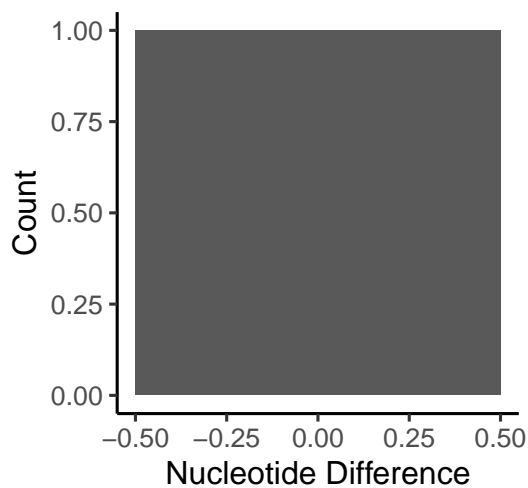
IGLV7-46*02

13 sequences assigned
12 (92.3%) exact matches, in which:
10 unique CDR3
2 unique J



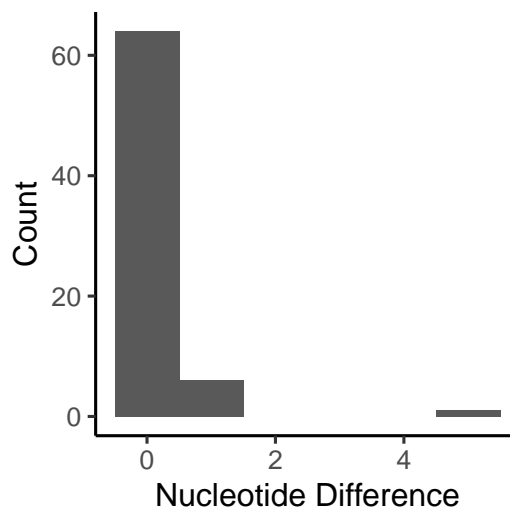
IGLV5-48*01

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



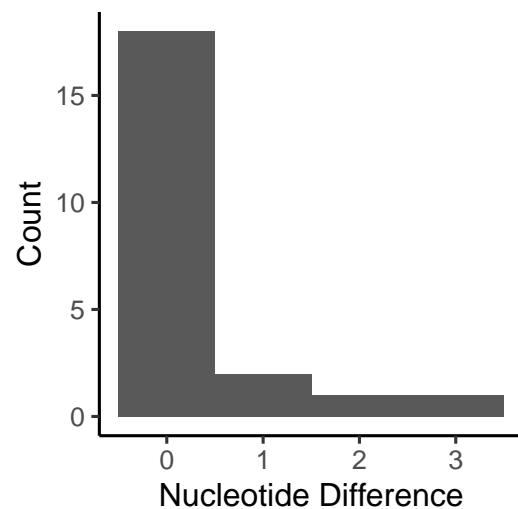
IGLV7-43*01

71 sequences assigned
64 (90.1%) exact matches, in which:
55 unique CDR3
2 unique J



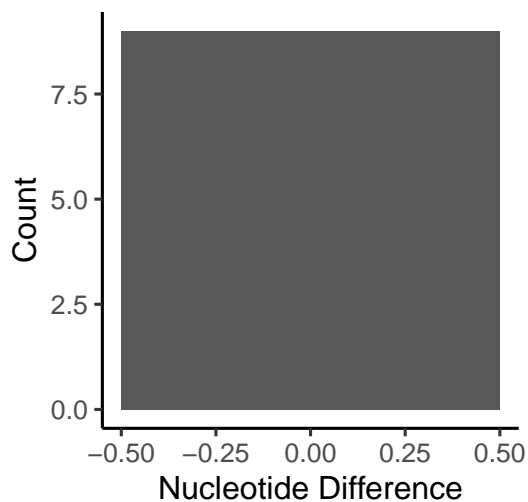
IGLV8-61*01

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



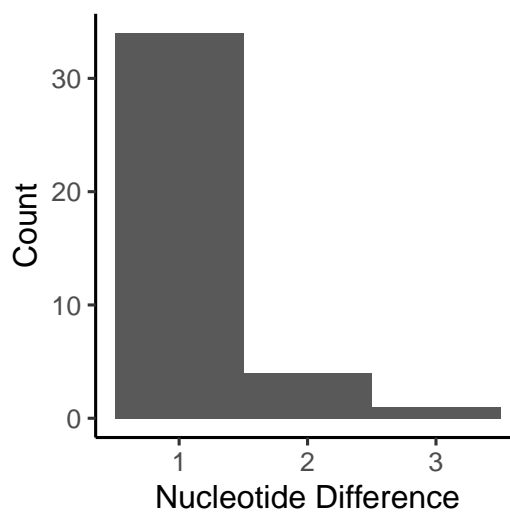
IGLV5-52*01

9 sequences assigned
9 (100%) exact matches, in which:
8 unique CDR3
2 unique J



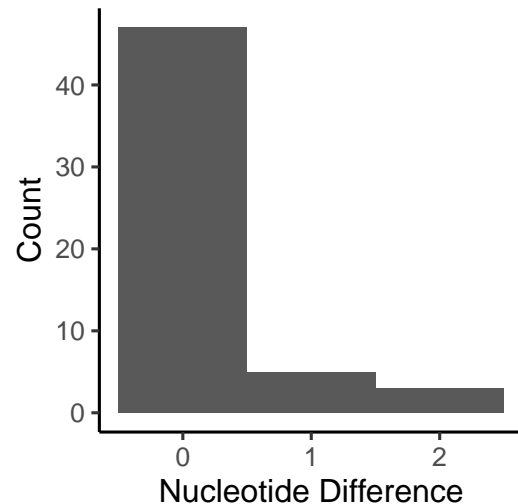
IGLV7-46*01

39 sequences assigned
No exact matches.

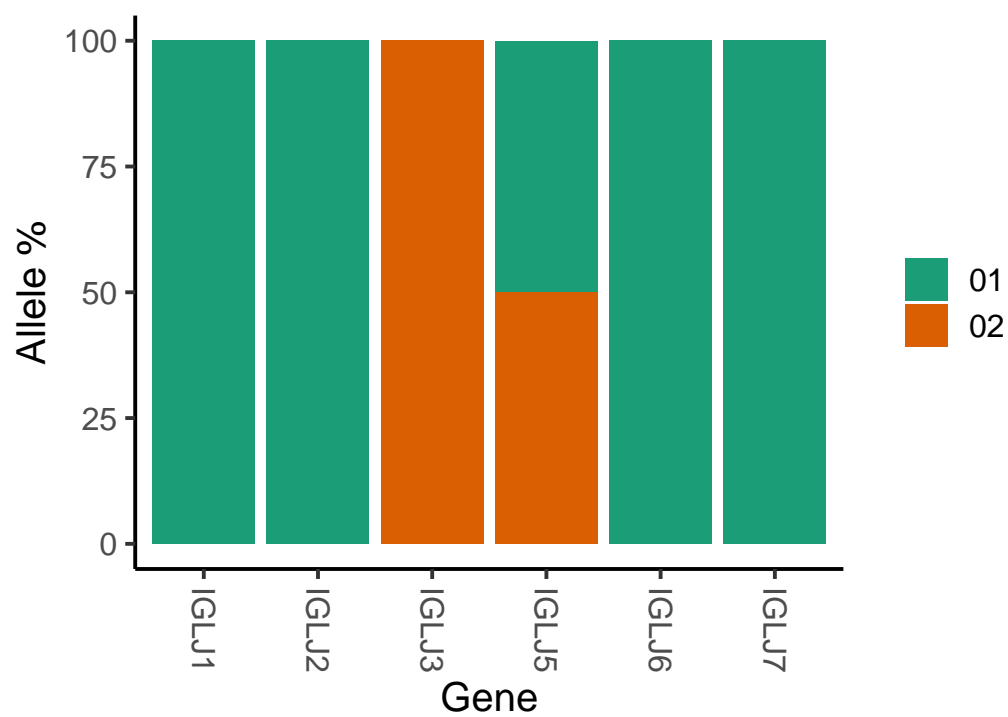


IGLV9-49*01

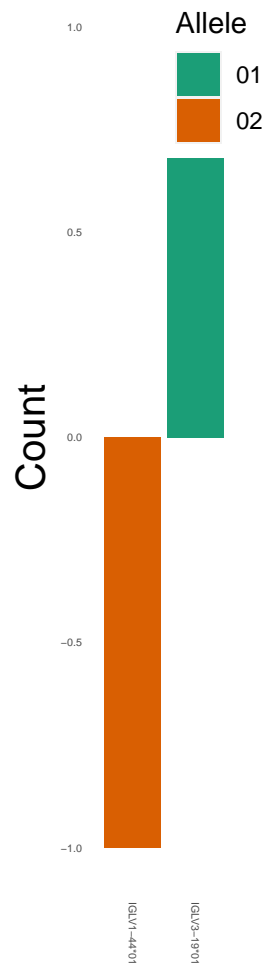
55 sequences assigned
47 (85.5%) exact matches, in which:
46 unique CDR3
3 unique J



Allele Usage



Sequence Count by IGLJ5 allele usage



Warning – no inferred sequences found.