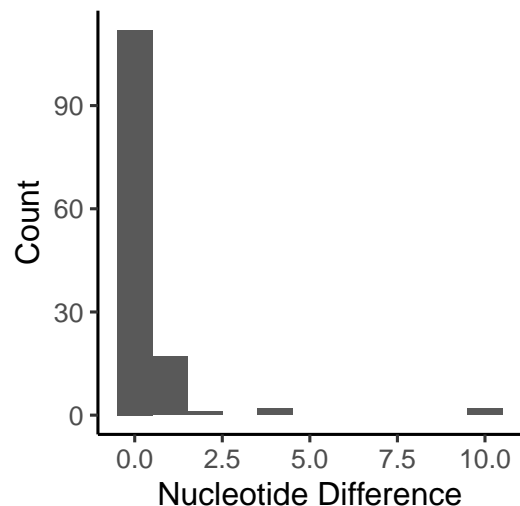


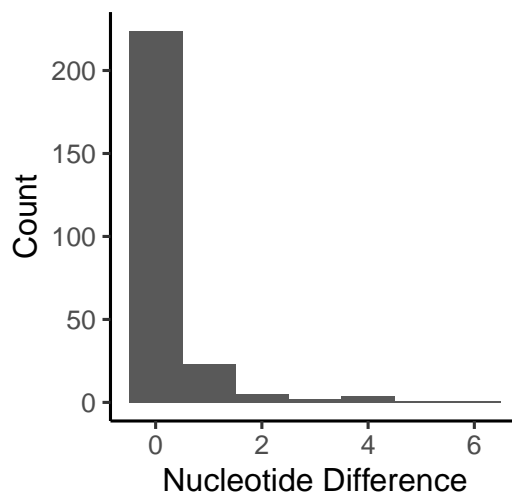
IGHV1-2*04

134 sequences assigned
112 (83.6%) exact matches, in which:
109 unique CDR3
5 unique J



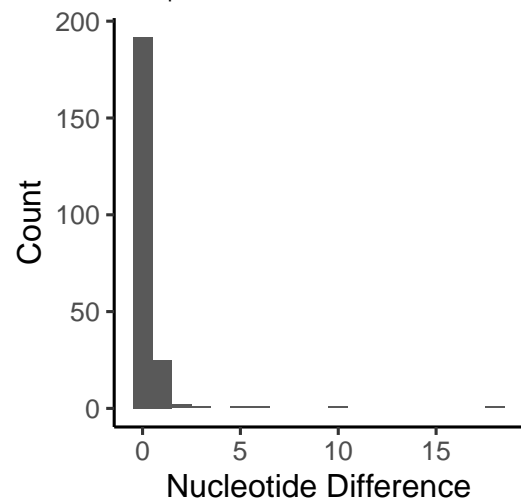
IGHV1-8*01

260 sequences assigned
224 (86.2%) exact matches, in which:
219 unique CDR3
6 unique J



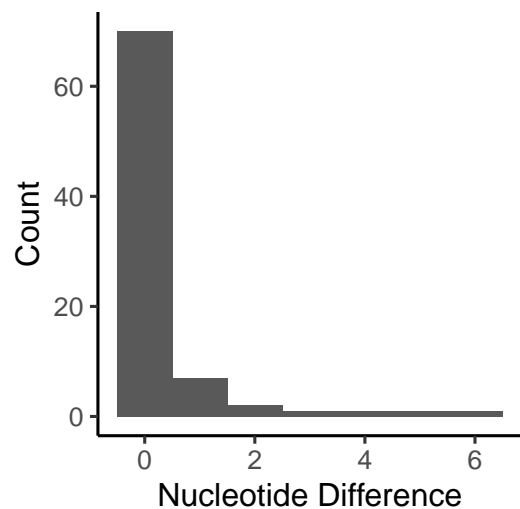
IGHV1-46*01

225 sequences assigned
192 (85.3%) exact matches, in which:
190 unique CDR3
6 unique J



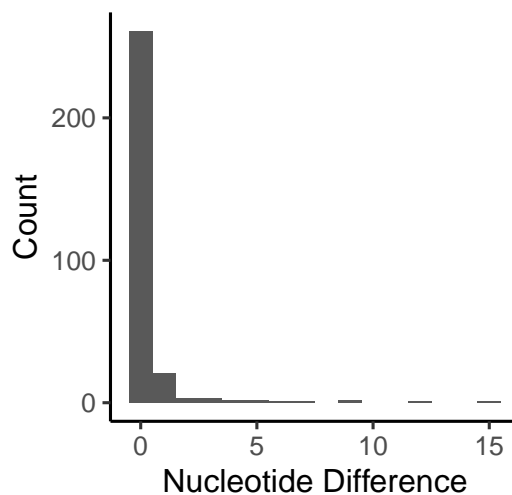
IGHV1-3*01

83 sequences assigned
70 (84.3%) exact matches, in which:
69 unique CDR3
5 unique J



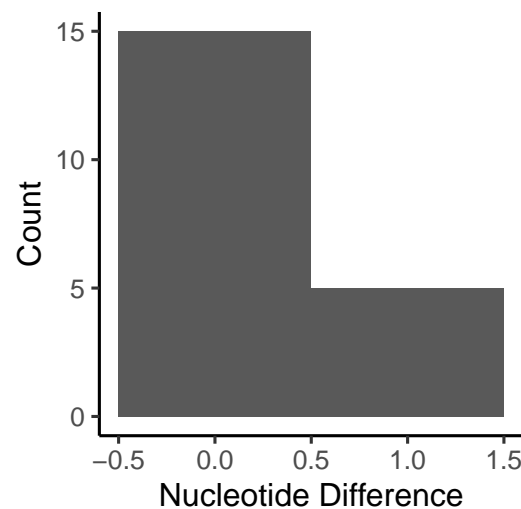
IGHV1-18*01

298 sequences assigned
261 (87.6%) exact matches, in which:
256 unique CDR3
6 unique J



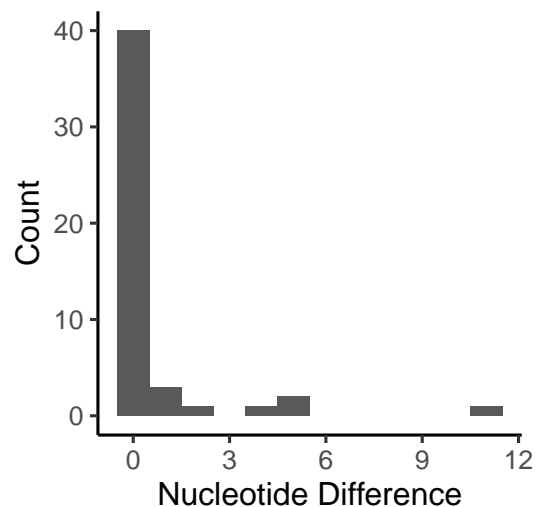
IGHV1-58*02

20 sequences assigned
15 (75%) exact matches, in which:
15 unique CDR3
5 unique J



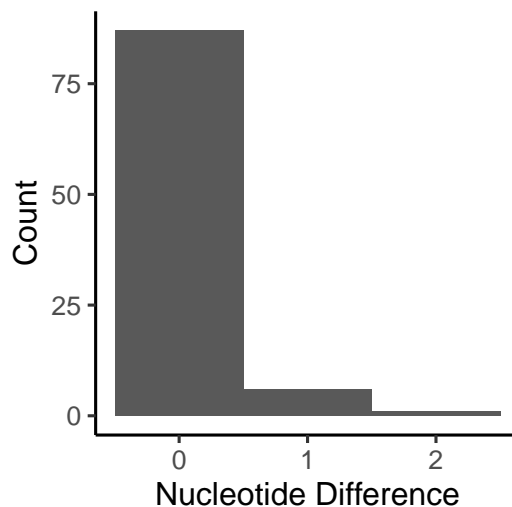
IGHV1-3*05

48 sequences assigned
40 (83.3%) exact matches, in which:
40 unique CDR3
6 unique J



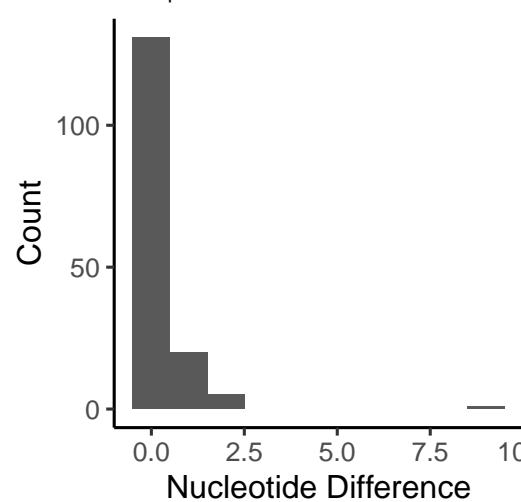
IGHV1-24*01

94 sequences assigned
87 (92.6%) exact matches, in which:
87 unique CDR3
5 unique J



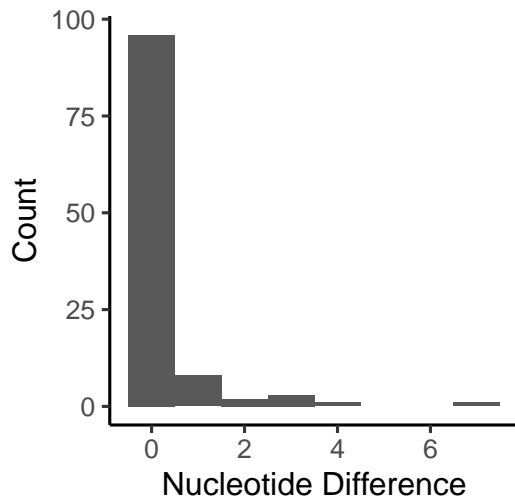
IGHV1-69*04

157 sequences assigned
131 (83.4%) exact matches, in which:
127 unique CDR3
6 unique J



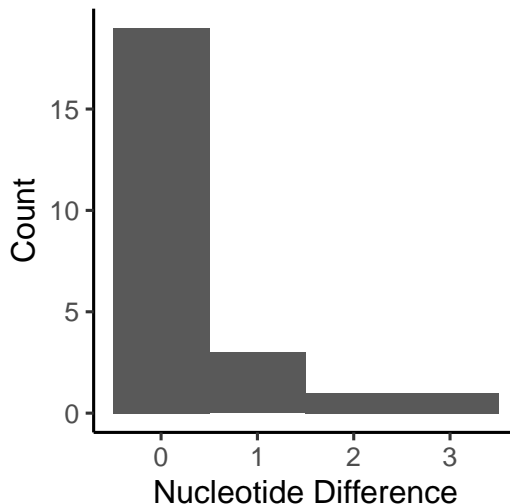
IGHV1–69*08_C191T

111 sequences assigned
96 (86.5%) exact matches, in which:
94 unique CDR3
6 unique J



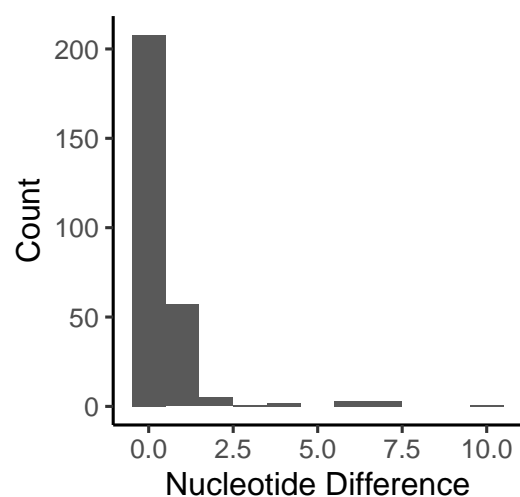
IGHV2–70*15

24 sequences assigned
19 (79.2%) exact matches, in which:
19 unique CDR3
6 unique J



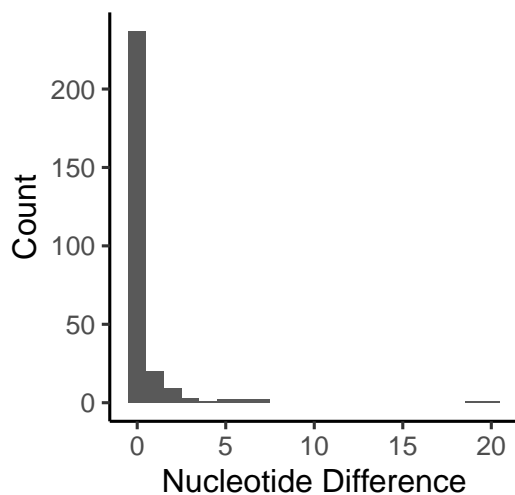
IGHV3–11*01

280 sequences assigned
208 (74.3%) exact matches, in which:
202 unique CDR3
6 unique J



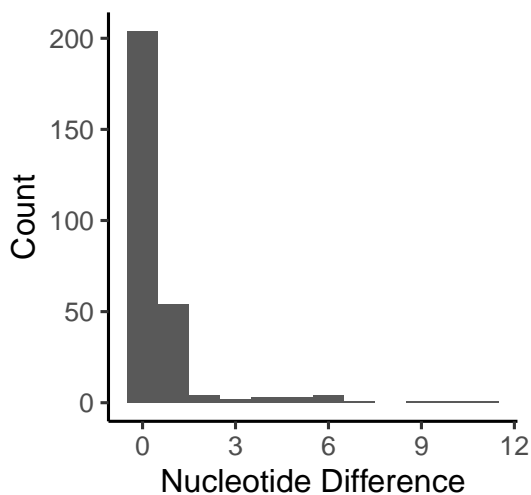
IGHV2–5*02

278 sequences assigned
237 (85.3%) exact matches, in which:
236 unique CDR3
6 unique J



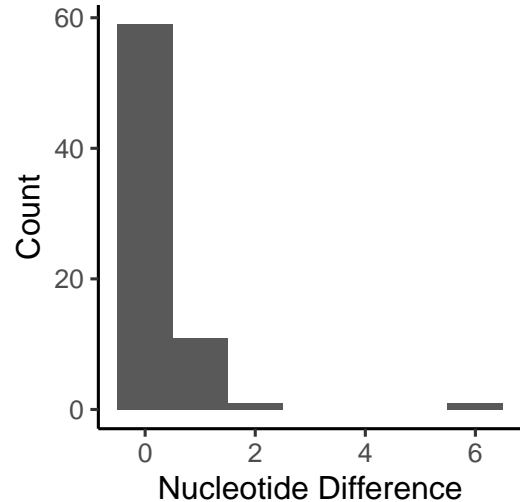
IGHV3–7*01

286 sequences assigned
204 (71.3%) exact matches, in which:
201 unique CDR3
6 unique J



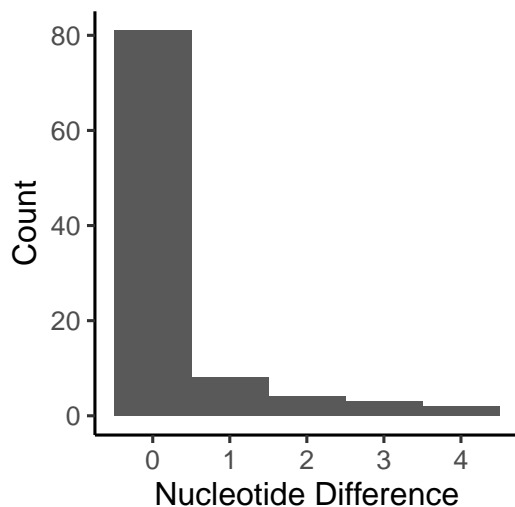
IGHV3–13*01

72 sequences assigned
59 (81.9%) exact matches, in which:
58 unique CDR3
5 unique J



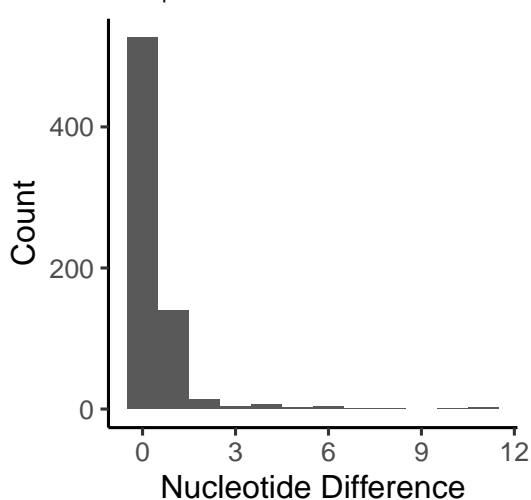
IGHV2–26*01

98 sequences assigned
81 (82.7%) exact matches, in which:
80 unique CDR3
5 unique J



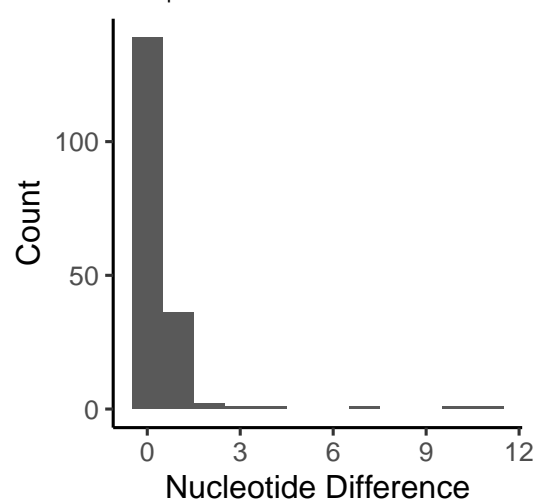
IGHV3–9*01

699 sequences assigned
527 (75.4%) exact matches, in which:
507 unique CDR3
6 unique J



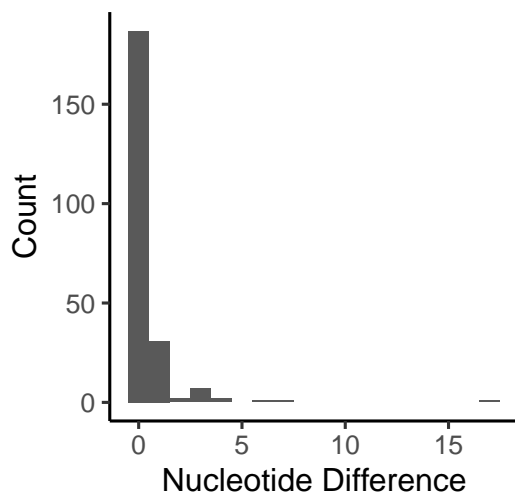
IGHV3–15*01

182 sequences assigned
139 (76.4%) exact matches, in which:
136 unique CDR3
5 unique J



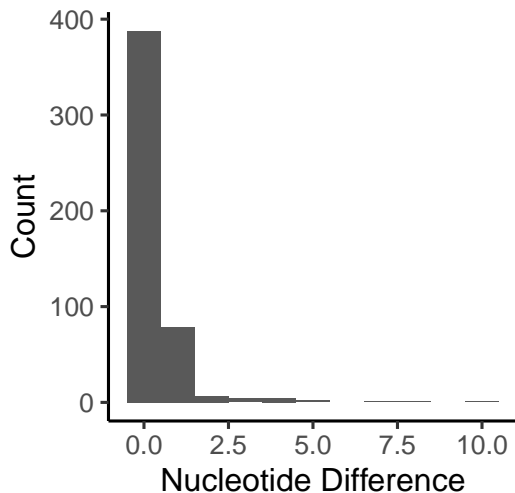
IGHV3-15*07

232 sequences assigned
187 (80.6%) exact matches, in which:
184 unique CDR3
6 unique J



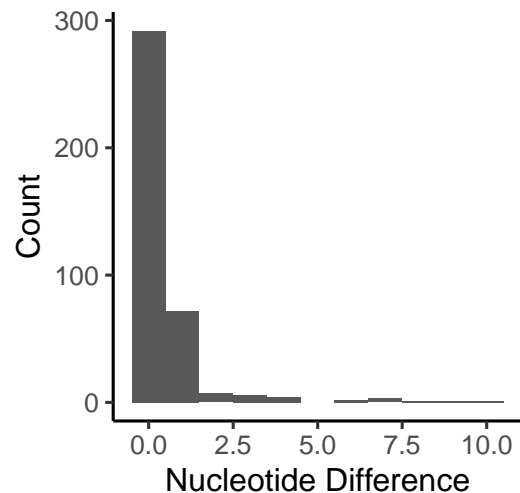
IGHV3-21*01

489 sequences assigned
388 (79.3%) exact matches, in which:
379 unique CDR3
6 unique J



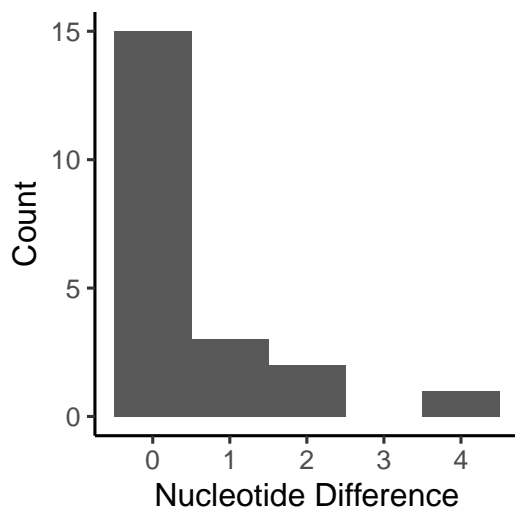
IGHV3-33*01

389 sequences assigned
292 (75.1%) exact matches, in which:
282 unique CDR3
6 unique J



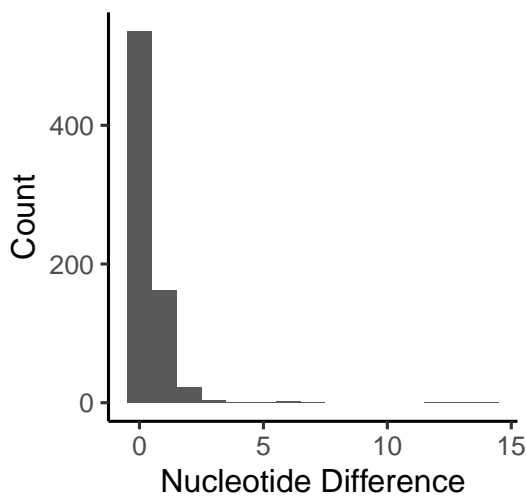
IGHV3-20*01

21 sequences assigned
15 (71.4%) exact matches, in which:
15 unique CDR3
5 unique J



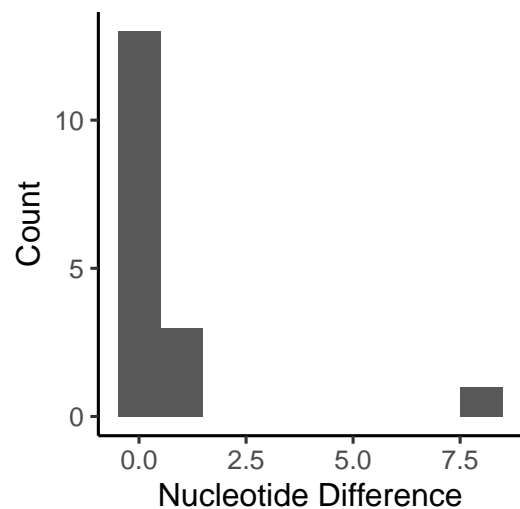
IGHV3-30-3*01

736 sequences assigned
536 (72.8%) exact matches, in which:
521 unique CDR3
6 unique J



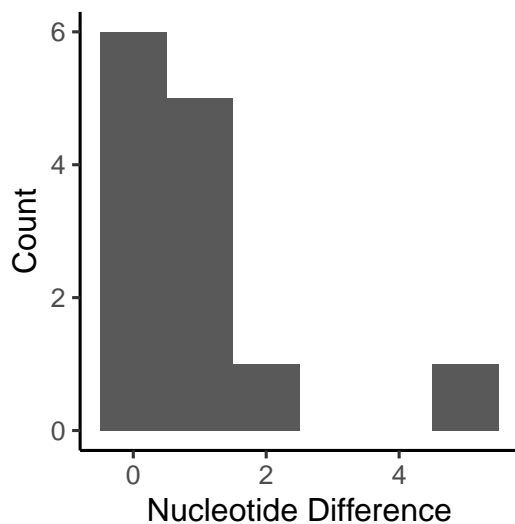
IGHV3-43*01

17 sequences assigned
13 (76.5%) exact matches, in which:
13 unique CDR3
3 unique J



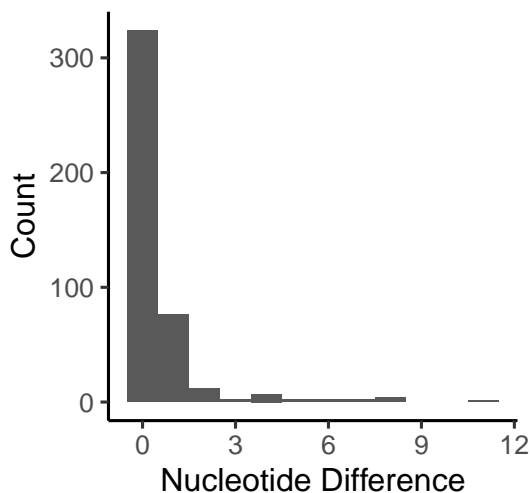
IGHV3-20*04

13 sequences assigned
6 (46.2%) exact matches, in which:
6 unique CDR3
3 unique J



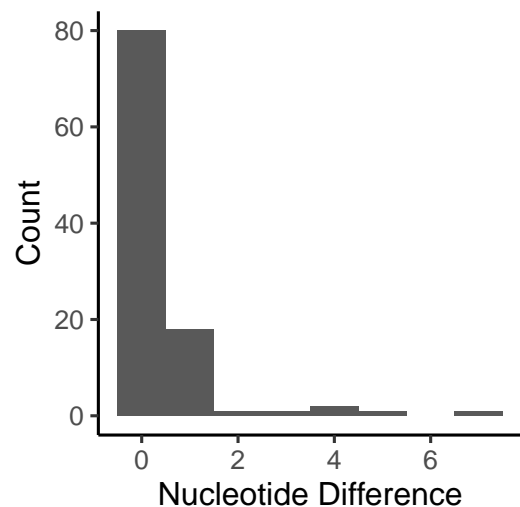
IGHV3-30*03

432 sequences assigned
324 (75%) exact matches, in which:
313 unique CDR3
6 unique J



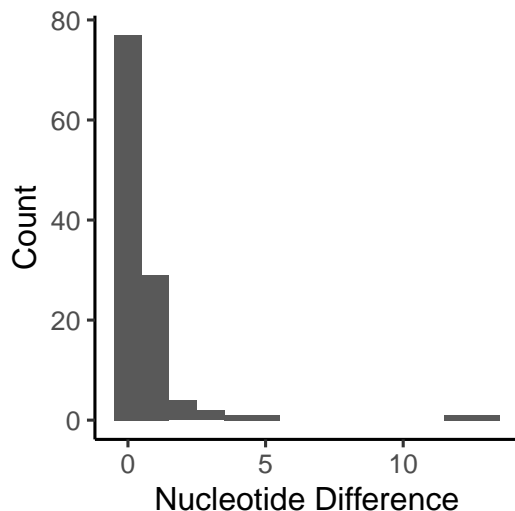
IGHV3-43*02

104 sequences assigned
80 (76.9%) exact matches, in which:
74 unique CDR3
6 unique J



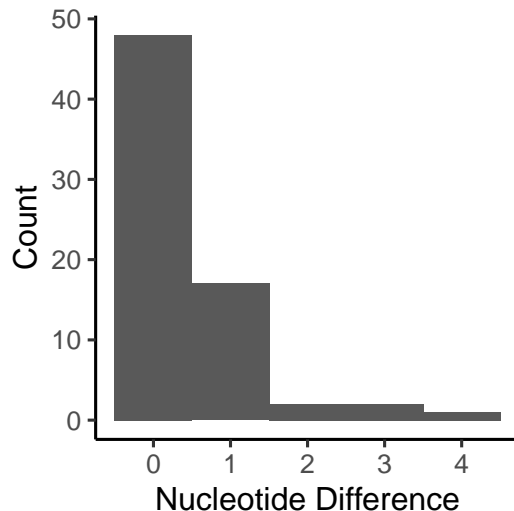
IGHV3-48*02

117 sequences assigned
77 (65.8%) exact matches, in which:
77 unique CDR3
6 unique J



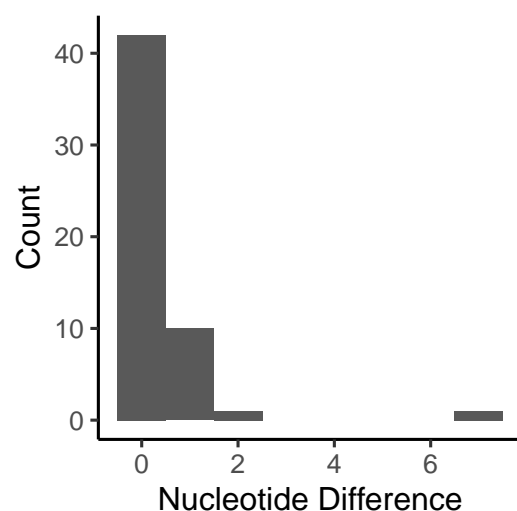
IGHV3-49*05

70 sequences assigned
48 (68.6%) exact matches, in which:
46 unique CDR3
4 unique J



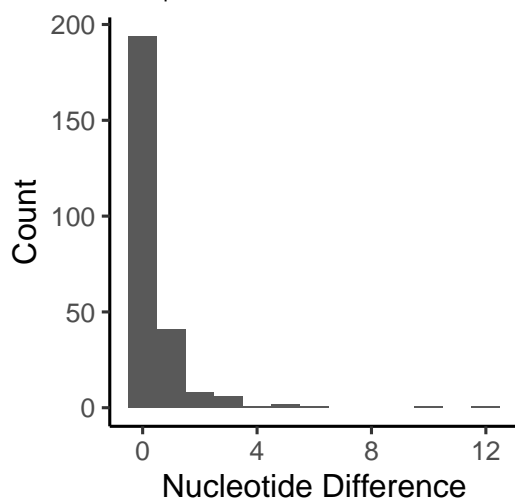
IGHV3-64*01

54 sequences assigned
42 (77.8%) exact matches, in which:
39 unique CDR3
4 unique J



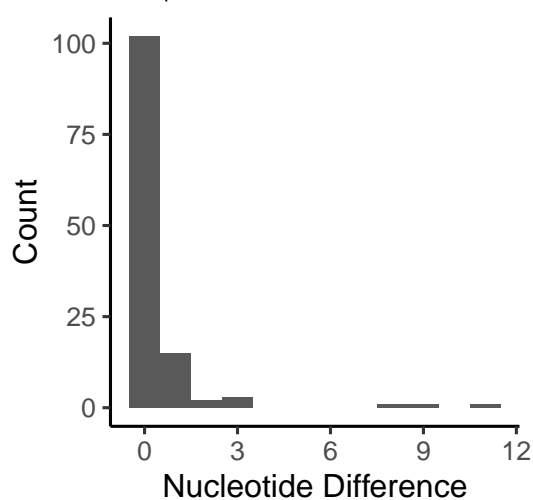
IGHV3-48*03

256 sequences assigned
194 (75.8%) exact matches, in which:
190 unique CDR3
6 unique J



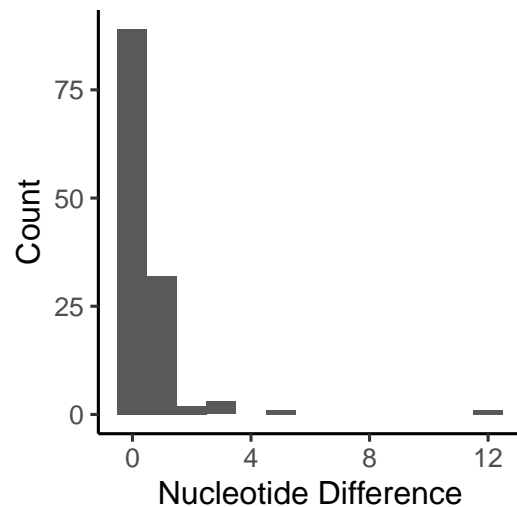
IGHV3-53*01

125 sequences assigned
102 (81.6%) exact matches, in which:
98 unique CDR3
6 unique J



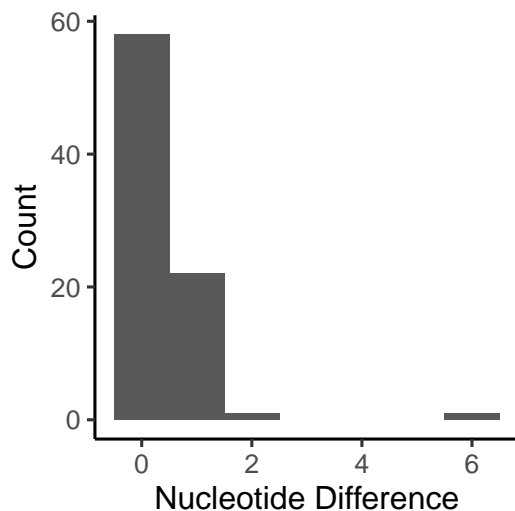
IGHV3-66*01

128 sequences assigned
89 (69.5%) exact matches, in which:
88 unique CDR3
6 unique J



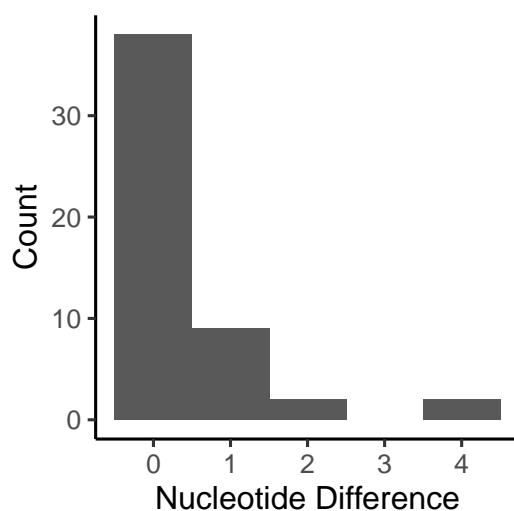
IGHV3-49*04

82 sequences assigned
58 (70.7%) exact matches, in which:
58 unique CDR3
6 unique J



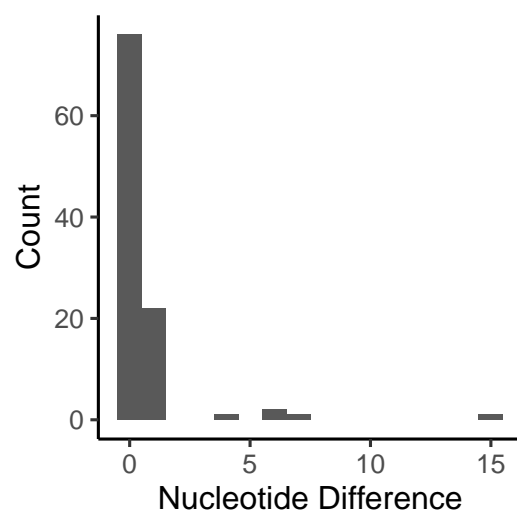
IGHV3-53*04

51 sequences assigned
38 (74.5%) exact matches, in which:
38 unique CDR3
5 unique J



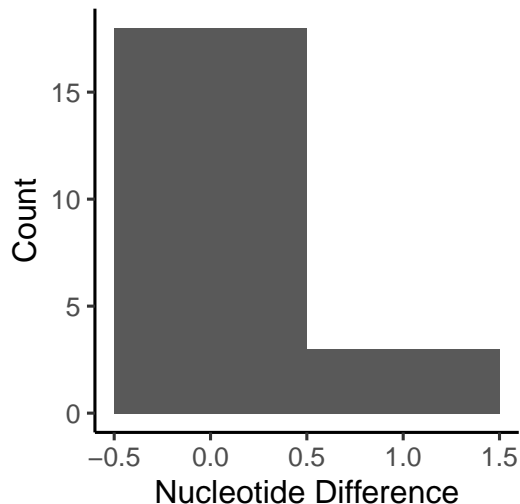
IGHV3-66*02

103 sequences assigned
76 (73.8%) exact matches, in which:
76 unique CDR3
5 unique J



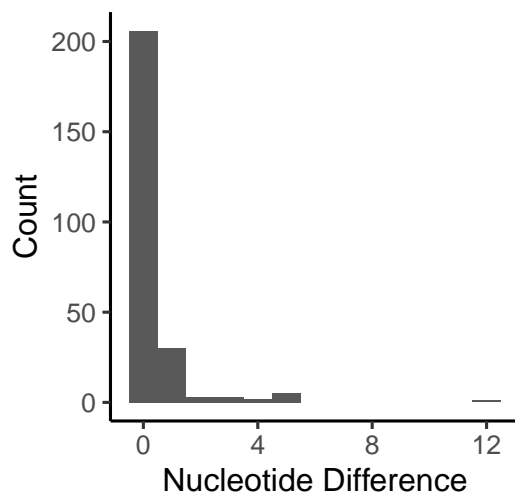
IGHV3-72*01

21 sequences assigned
18 (85.7%) exact matches, in which:
17 unique CDR3
5 unique J



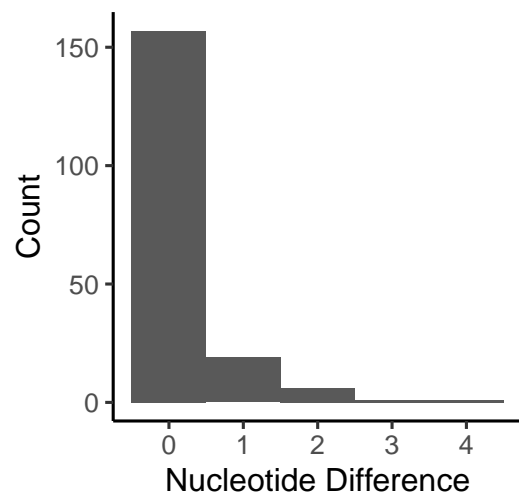
IGHV4-4*02

250 sequences assigned
206 (82.4%) exact matches, in which:
203 unique CDR3
6 unique J



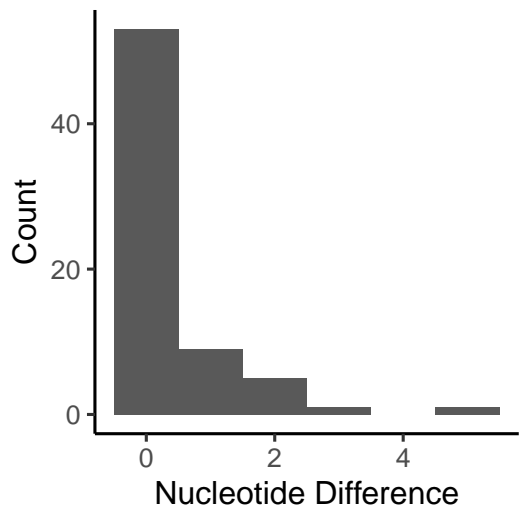
IGHV4-30-4*01

184 sequences assigned
157 (85.3%) exact matches, in which:
152 unique CDR3
6 unique J



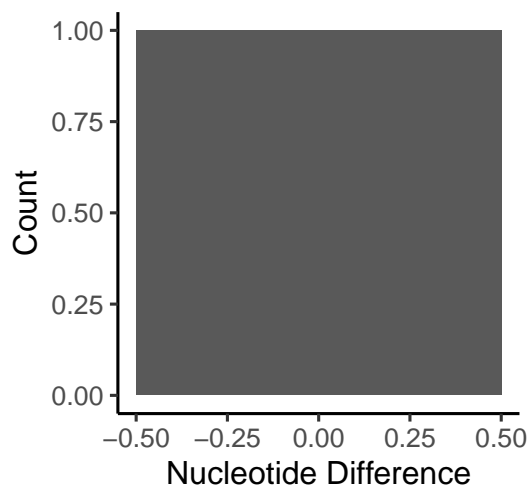
IGHV3-73*01

69 sequences assigned
53 (76.8%) exact matches, in which:
52 unique CDR3
5 unique J



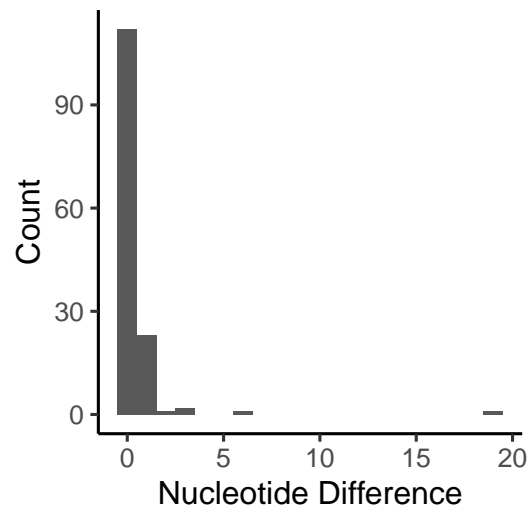
IGHV4-28*01

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



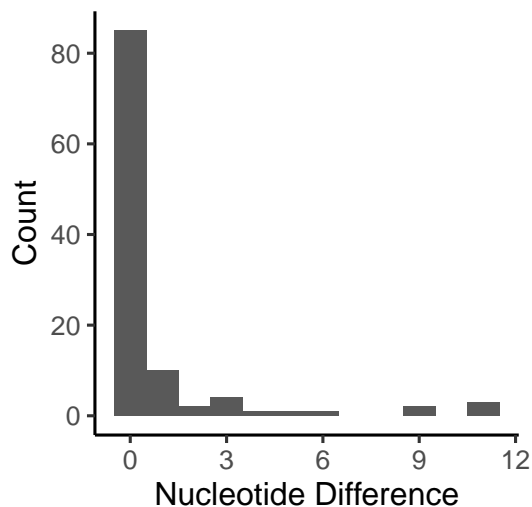
IGHV4-30-2*01

140 sequences assigned
112 (80%) exact matches, in which:
109 unique CDR3
6 unique J



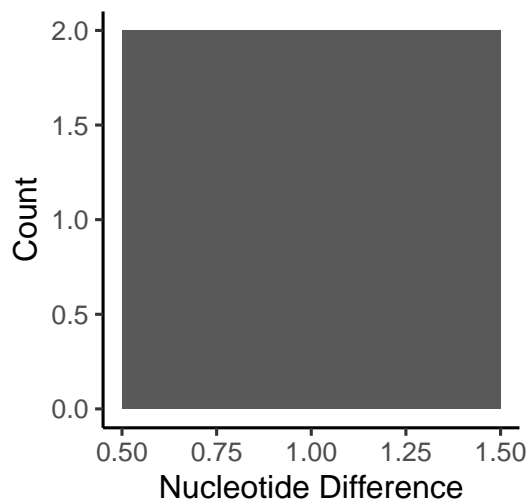
IGHV3-74*01

109 sequences assigned
85 (78%) exact matches, in which:
81 unique CDR3
6 unique J



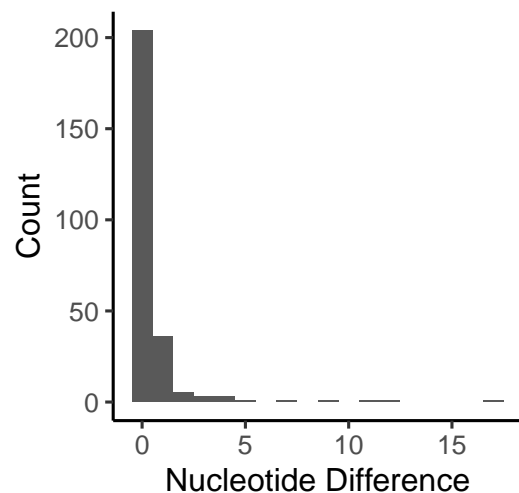
IGHV4-28*04

2 sequences assigned
No exact matches.



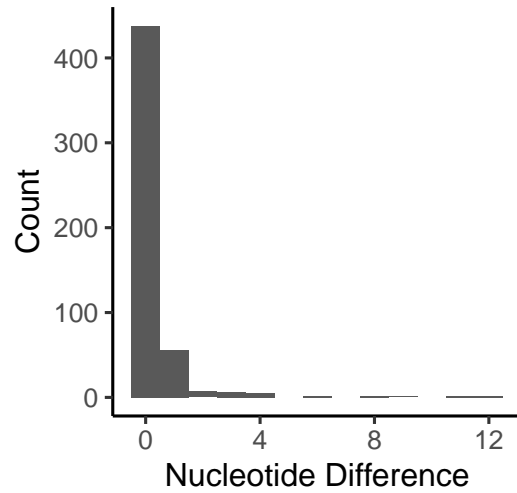
IGHV4-31*03

257 sequences assigned
204 (79.4%) exact matches, in which:
200 unique CDR3
6 unique J



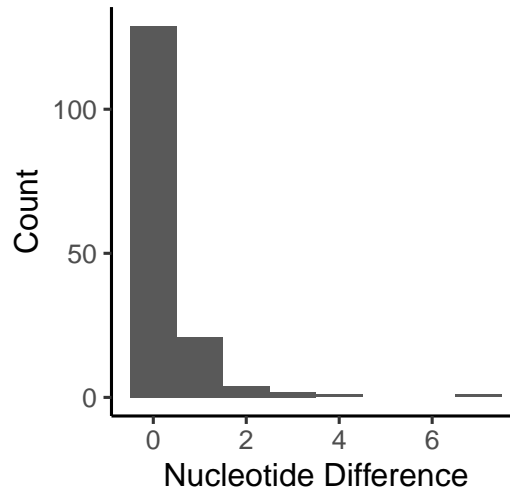
IGHV4-34*01

522 sequences assigned
438 (83.9%) exact matches, in which:
427 unique CDR3
6 unique J



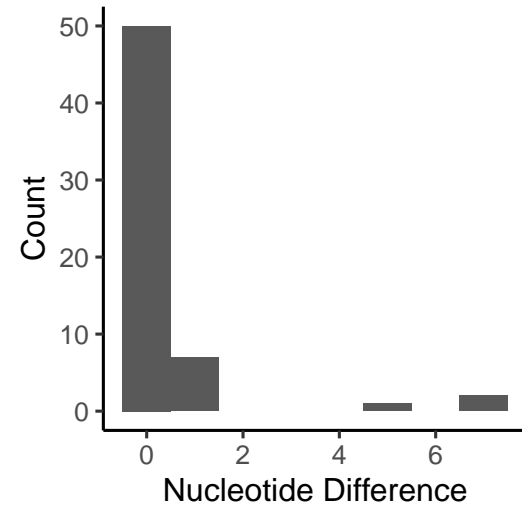
IGHV4-59*08

158 sequences assigned
129 (81.6%) exact matches, in which:
127 unique CDR3
6 unique J



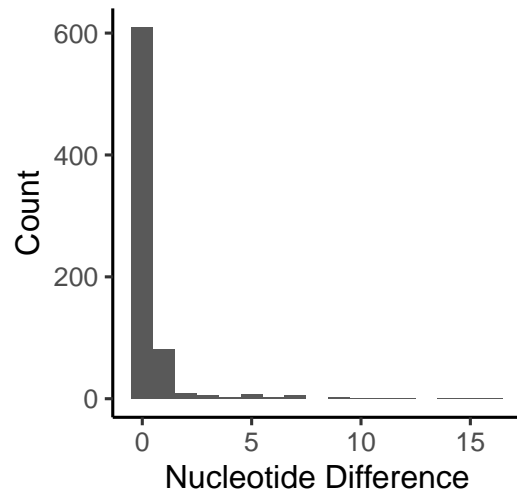
IGHV6-1*01

60 sequences assigned
50 (83.3%) exact matches, in which:
48 unique CDR3
5 unique J



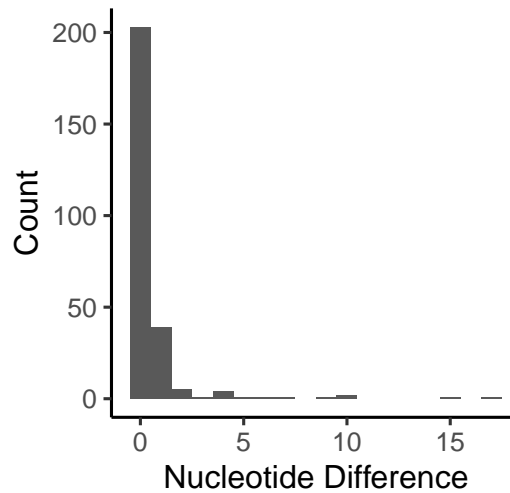
IGHV4-39*01

732 sequences assigned
610 (83.3%) exact matches, in which:
587 unique CDR3
6 unique J



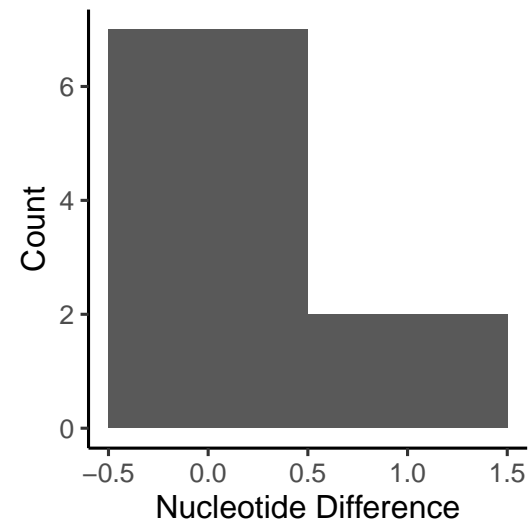
IGHV4-61*02

260 sequences assigned
203 (78.1%) exact matches, in which:
203 unique CDR3
6 unique J



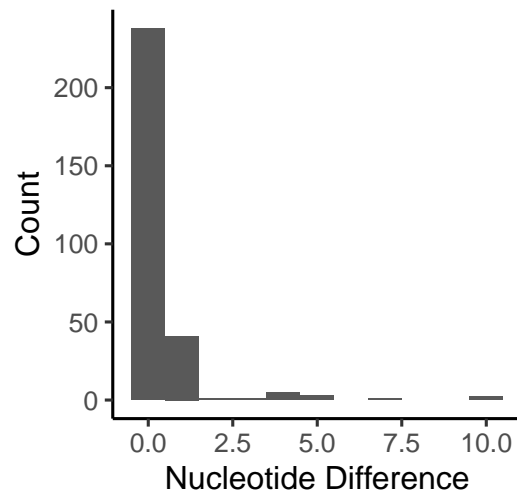
IGHV7-4-1*01

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



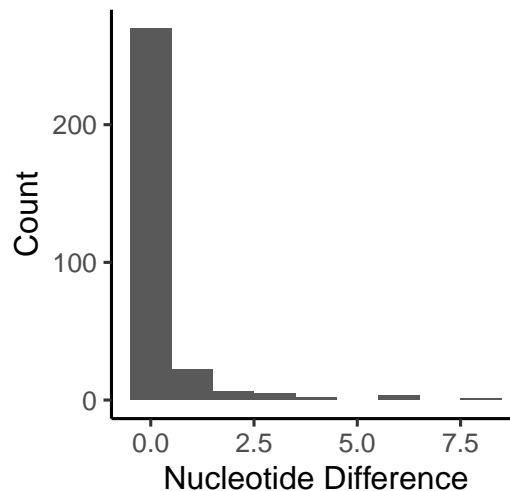
IGHV4-59*01

292 sequences assigned
238 (81.5%) exact matches, in which:
238 unique CDR3
6 unique J



IGHV5-51*01

309 sequences assigned
270 (87.4%) exact matches, in which:
264 unique CDR3
6 unique J





Warning – no inferred sequences found.