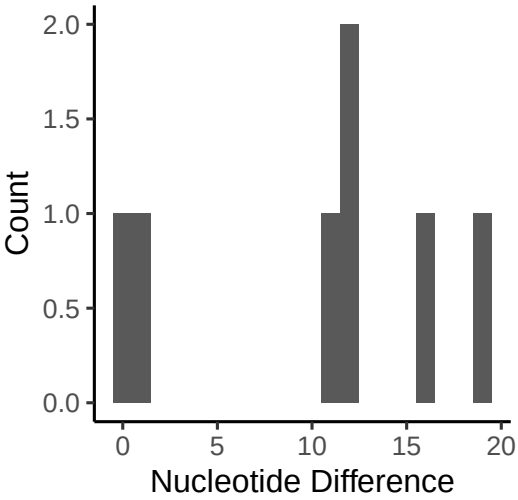
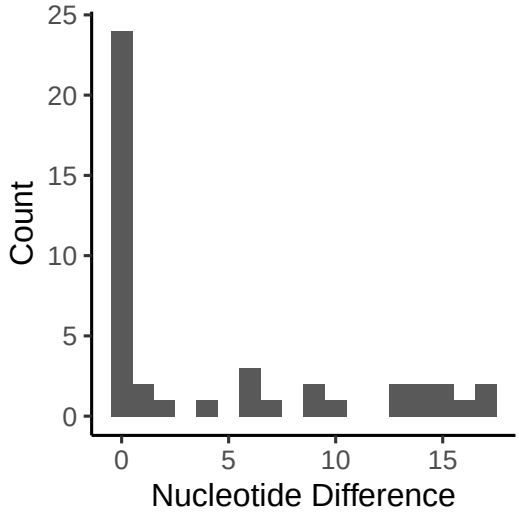
IGHV1−2*06

11 sequences assigned
1 (9.1%) exact matches, in which:
1 unique CDR3
1 unique J



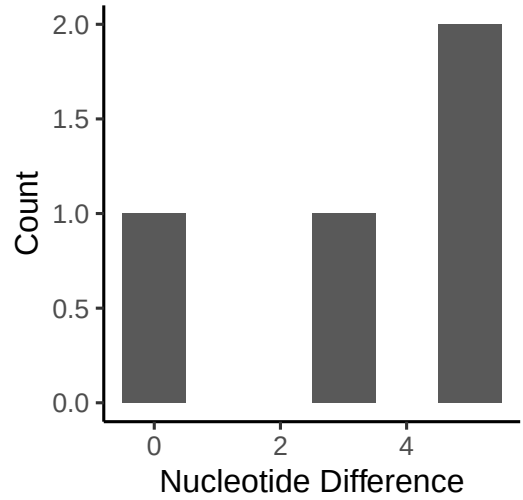
IGHV1−18*01

51 sequences assigned
24 (47.1%) exact matches, in which:
23 unique CDR3
6 unique J



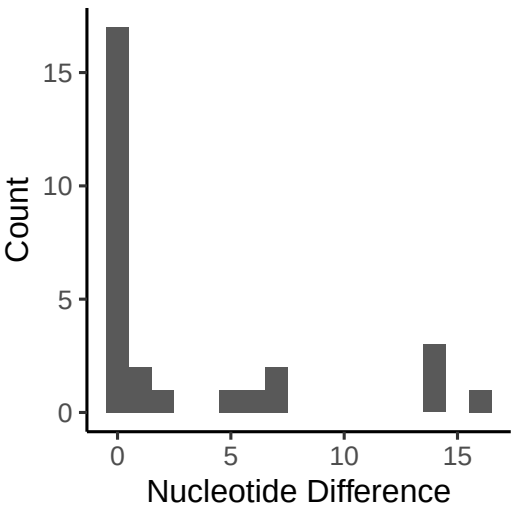
IGHV1−58*01

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



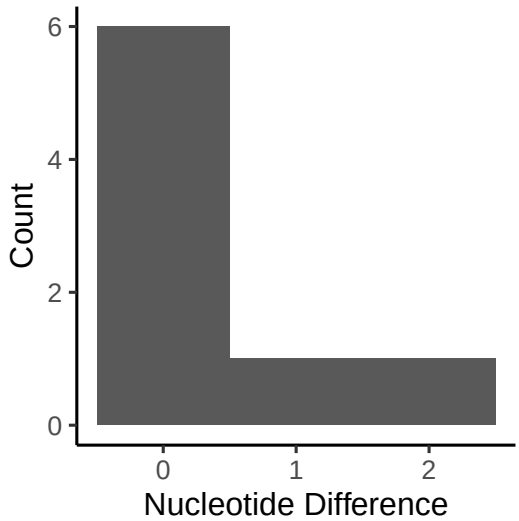
IGHV1−3*01

33 sequences assigned
17 (51.5%) exact matches, in which:
17 unique CDR3
6 unique J



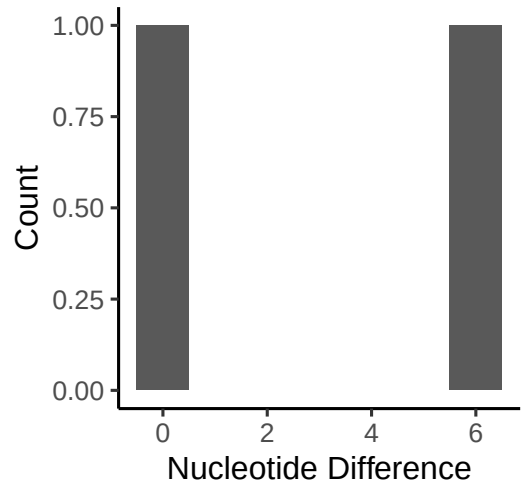
IGHV1−24*01

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



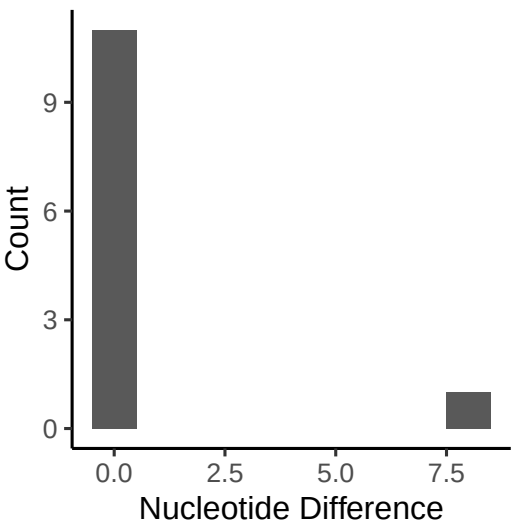
IGHV1−69−2*01

6 sequences assigned
1 (16.7%) exact matches, in which:
1 unique CDR3
1 unique J



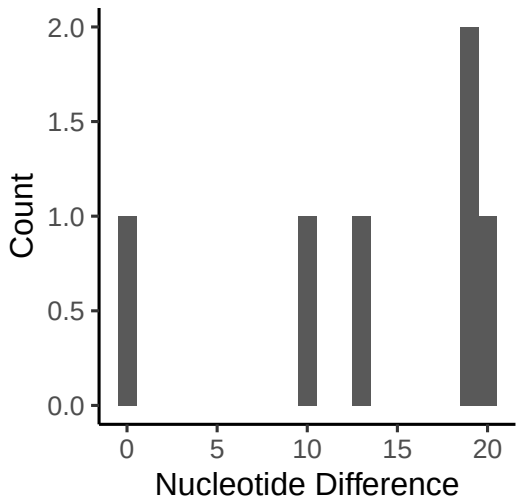
IGHV1−8*01

13 sequences assigned
11 (84.6%) exact matches, in which:
11 unique CDR3
4 unique J



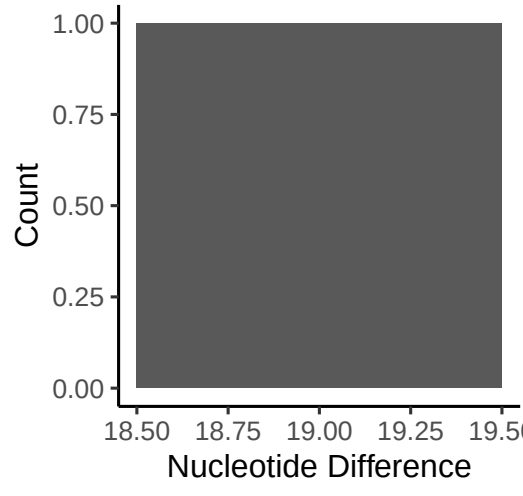
IGHV1−46*01

7 sequences assigned
1 (14.3%) exact matches, in which:
1 unique CDR3
1 unique J



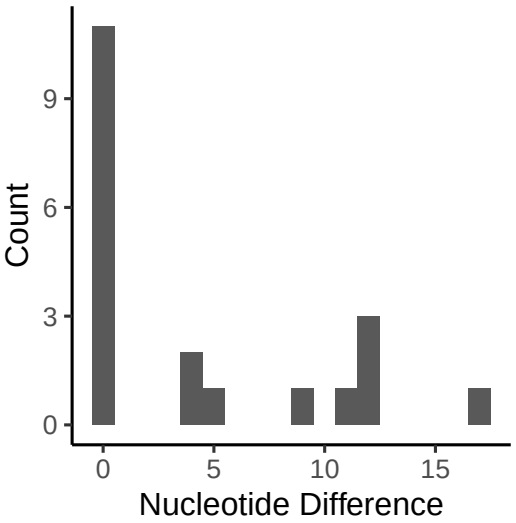
IGHV1−69*06

5 sequences assigned
No exact matches.



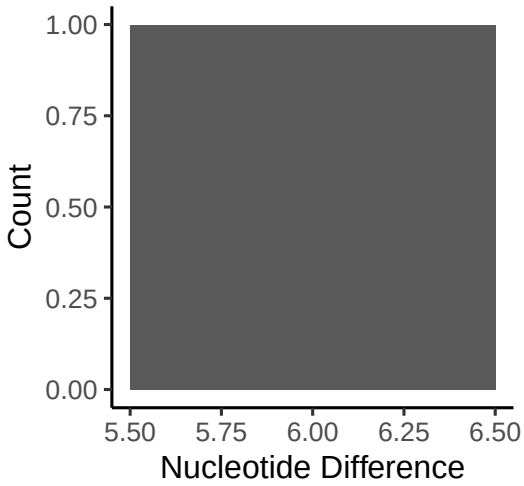
IGHV2−5*02

23 sequences assigned
11 (47.8%) exact matches, in which:
11 unique CDR3
3 unique J



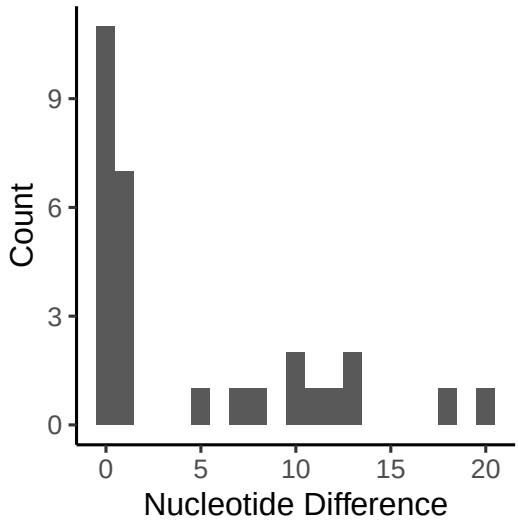
IGHV2−70*13

1 sequences assigned
No exact matches.



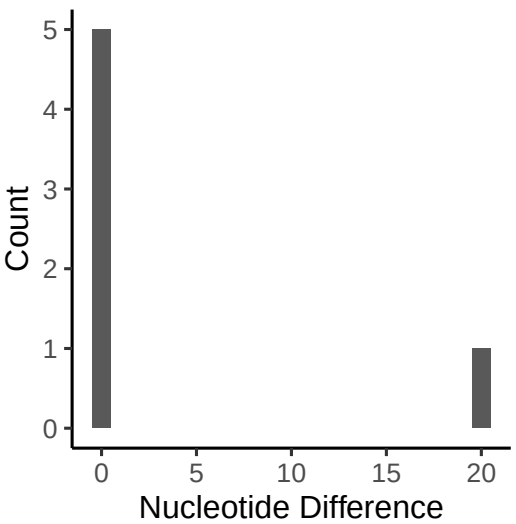
IGHV3−11*01

38 sequences assigned
11 (28.9%) exact matches, in which:
11 unique CDR3
5 unique J



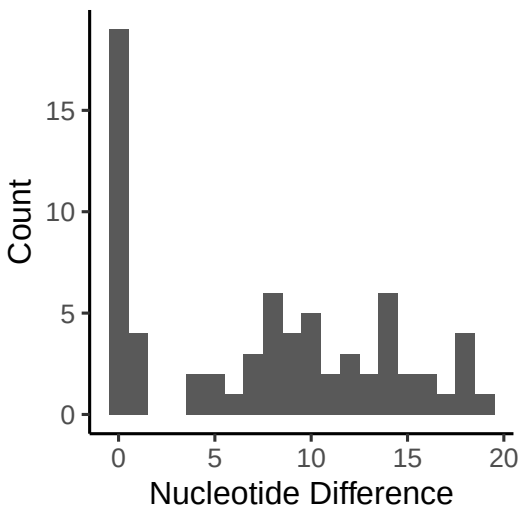
IGHV2−26*01

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



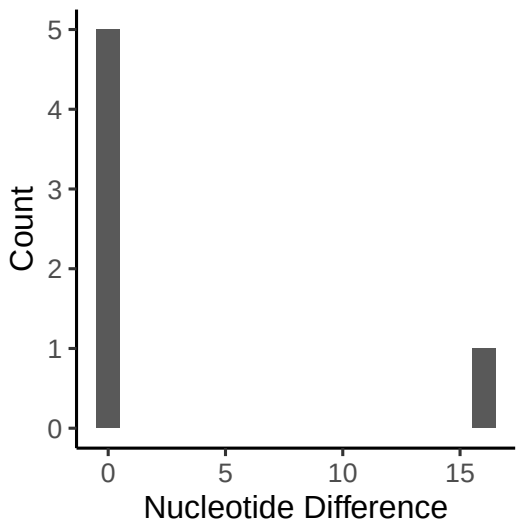
IGHV3−7*01

83 sequences assigned
19 (22.9%) exact matches, in which:
19 unique CDR3
5 unique J



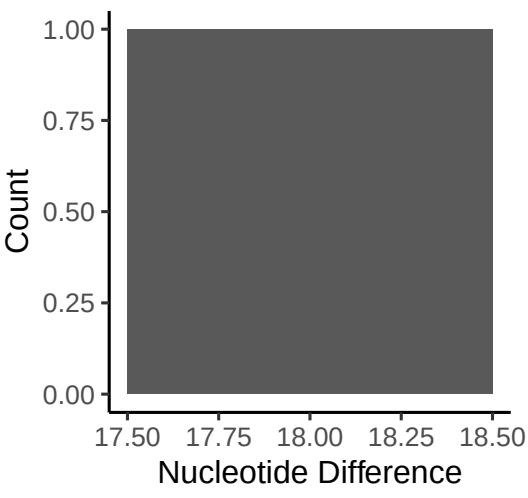
IGHV3−13*01

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



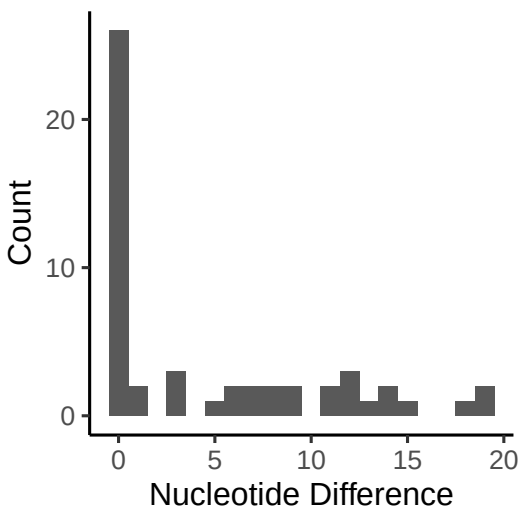
IGHV2−26*03

1 sequences assigned
No exact matches.



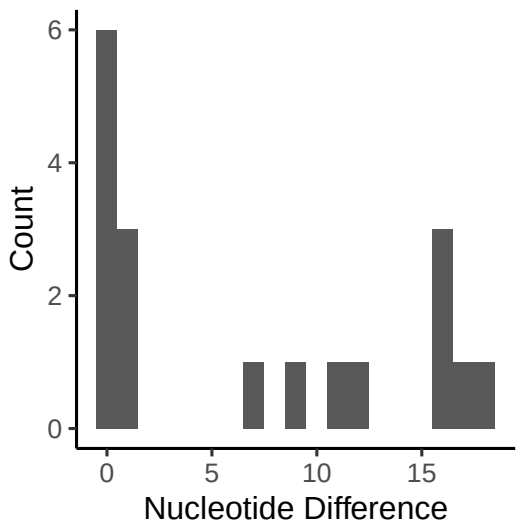
IGHV3−9*01

61 sequences assigned
26 (42.6%) exact matches, in which:
26 unique CDR3
5 unique J



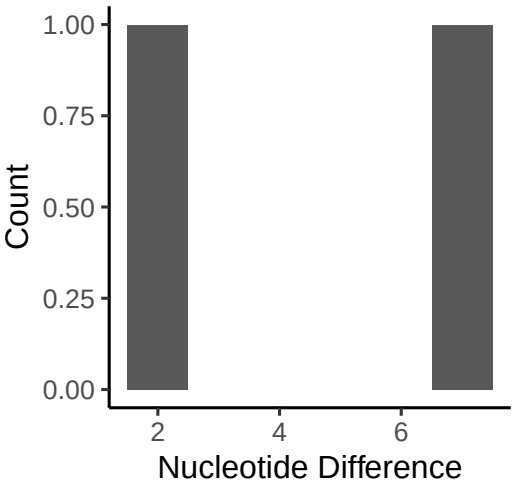
IGHV3−15*01

22 sequences assigned
6 (27.3%) exact matches, in which:
6 unique CDR3
3 unique J



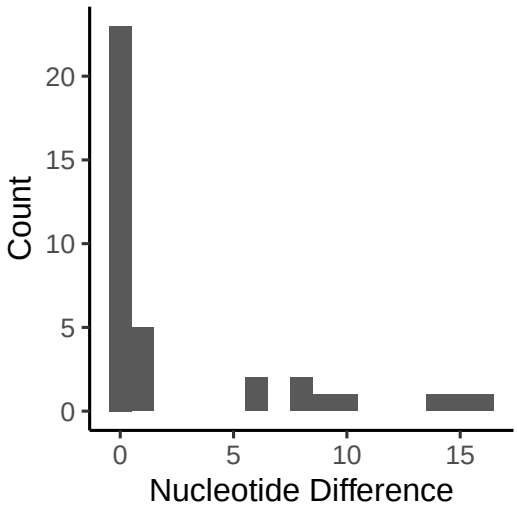
IGHV3−20*01

2 sequences assigned
No exact matches.



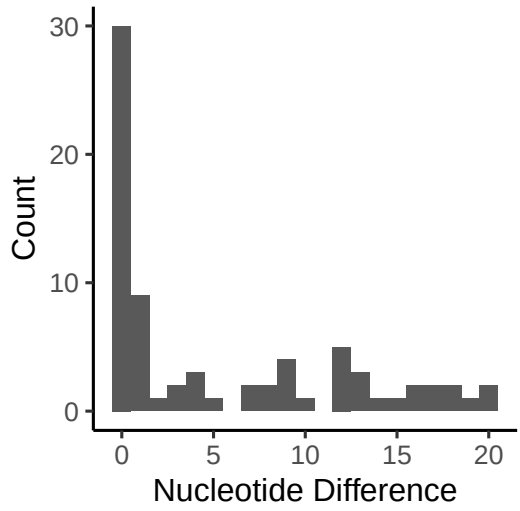
IGHV3−30−3*01

37 sequences assigned
23 (62.2%) exact matches, in which:
23 unique CDR3
5 unique J



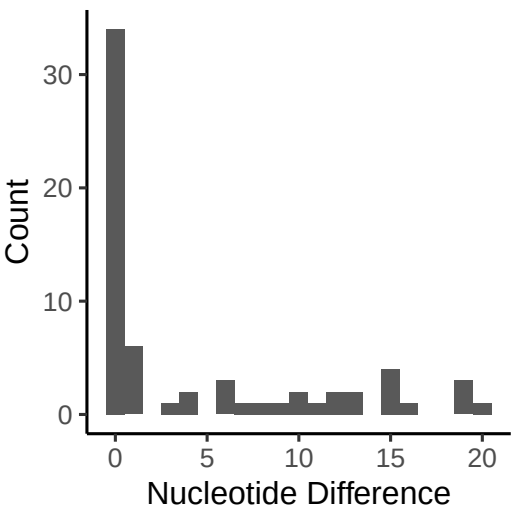
IGHV3−48*03

86 sequences assigned
30 (34.9%) exact matches, in which:
30 unique CDR3
5 unique J



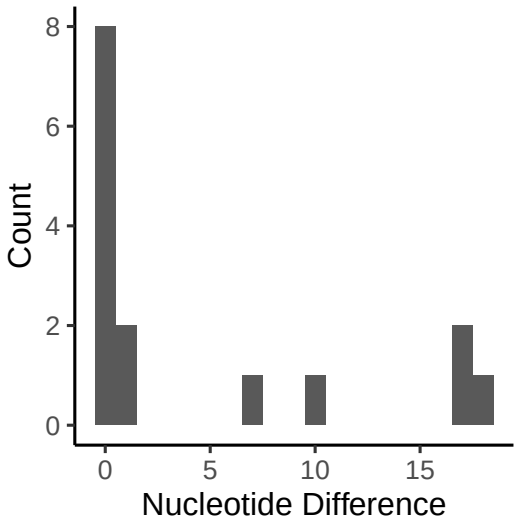
IGHV3−21*01

78 sequences assigned
34 (43.6%) exact matches, in which:
34 unique CDR3
5 unique J



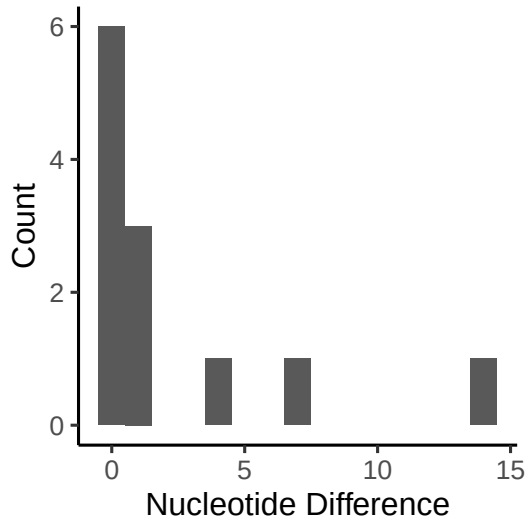
IGHV3−33*01

18 sequences assigned
8 (44.4%) exact matches, in which:
8 unique CDR3
4 unique J



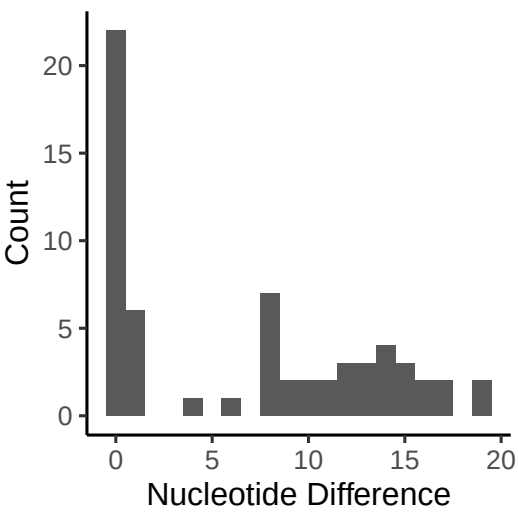
IGHV3−49*04

21 sequences assigned
6 (28.6%) exact matches, in which:
6 unique CDR3
2 unique J



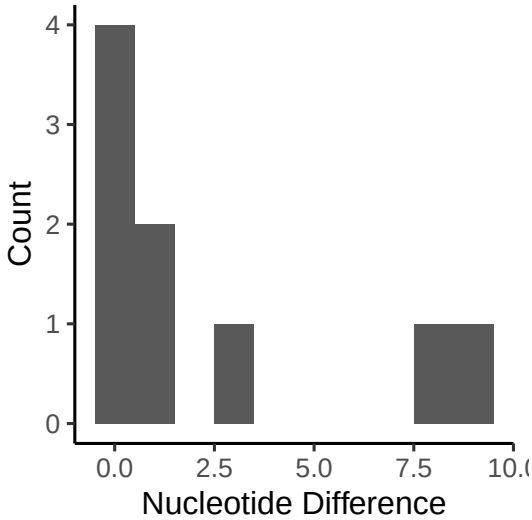
IGHV3−23*04

82 sequences assigned
22 (26.8%) exact matches, in which:
22 unique CDR3
3 unique J



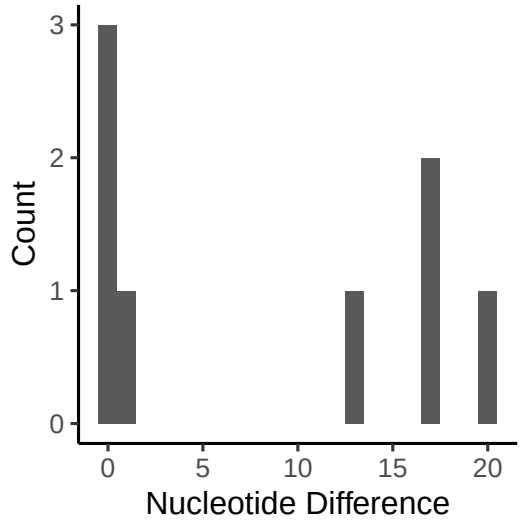
IGHV3−43*02

15 sequences assigned
4 (26.7%) exact matches, in which:
4 unique CDR3
2 unique J



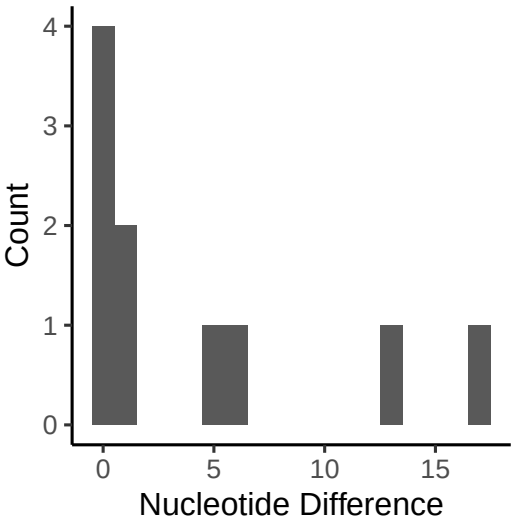
IGHV3−53*02

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



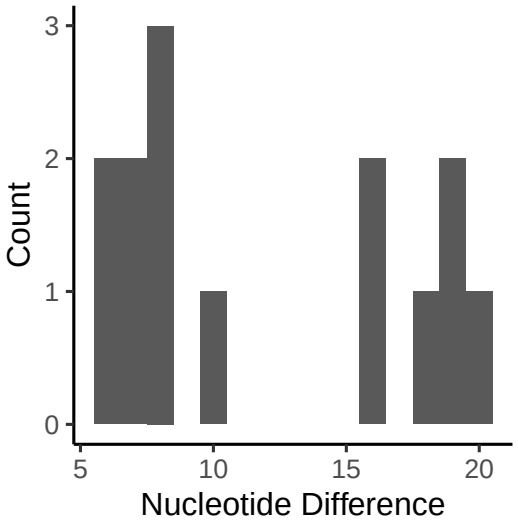
IGHV3−53*04

11 sequences assigned
4 (36.4%) exact matches, in which:
4 unique CDR3
3 unique J



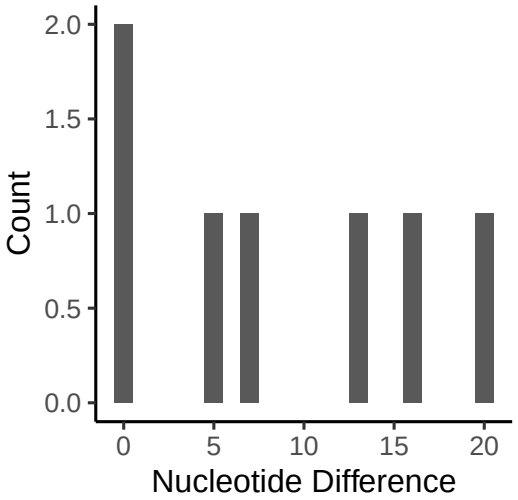
IGHV3−74*01

19 sequences assigned
No exact matches.



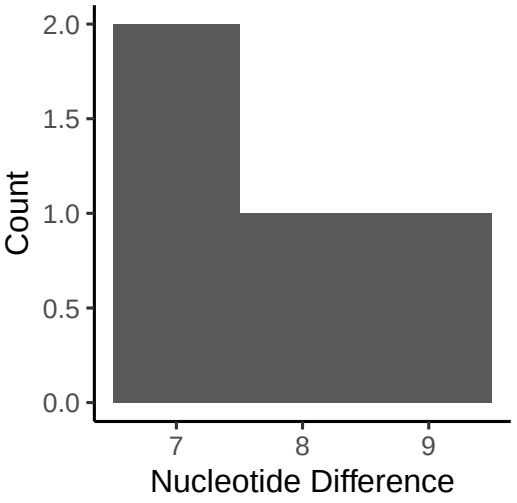
IGHV4−30−4*01

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



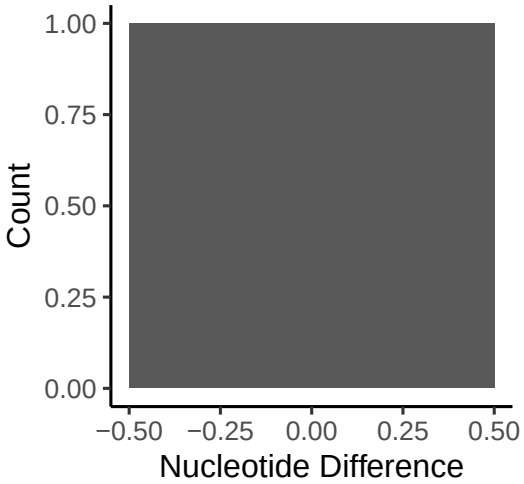
IGHV3−72*01

7 sequences assigned
No exact matches.



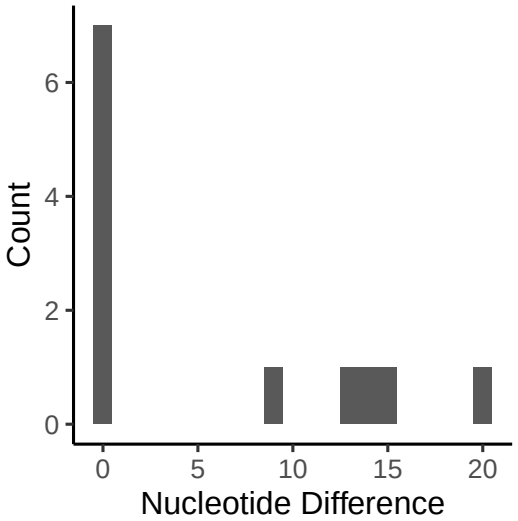
IGHV3−43D*04

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



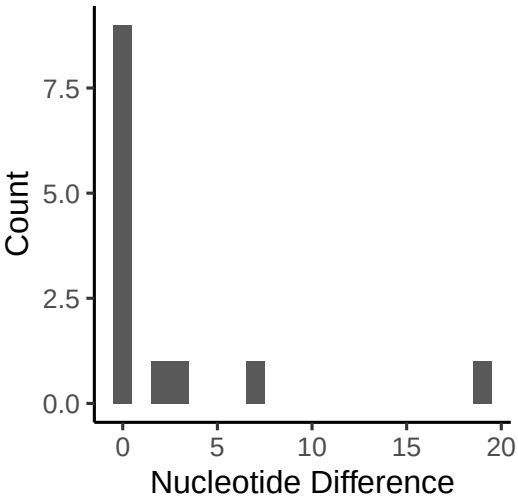
IGHV4−31*03

14 sequences assigned
7 (50%) exact matches, in which:
7 unique CDR3
4 unique J



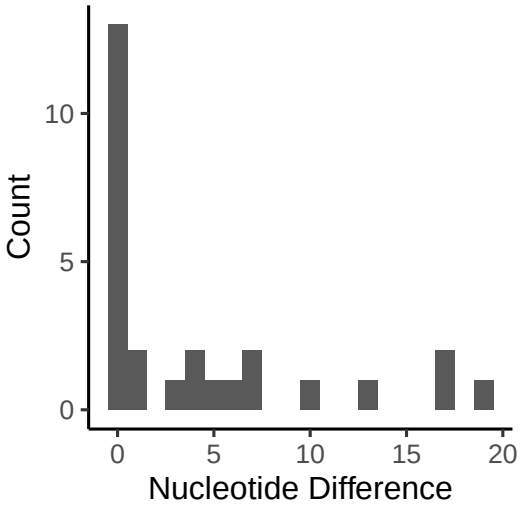
IGHV3−73*02

18 sequences assigned
9 (50%) exact matches, in which:
9 unique CDR3
4 unique J



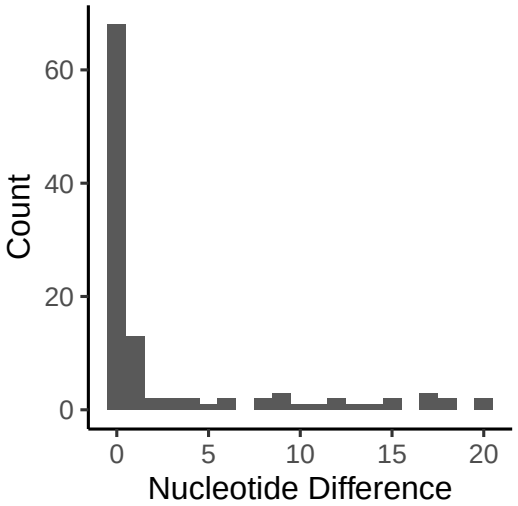
IGHV4−4*02

34 sequences assigned
13 (38.2%) exact matches, in which:
13 unique CDR3
3 unique J



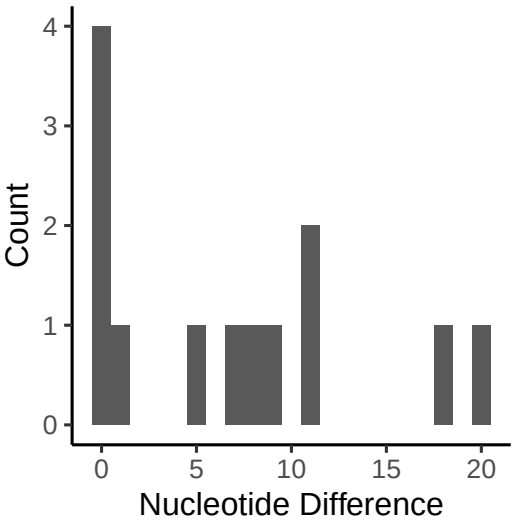
IGHV4−34*01

124 sequences assigned
68 (54.8%) exact matches, in which:
68 unique CDR3
7 unique J



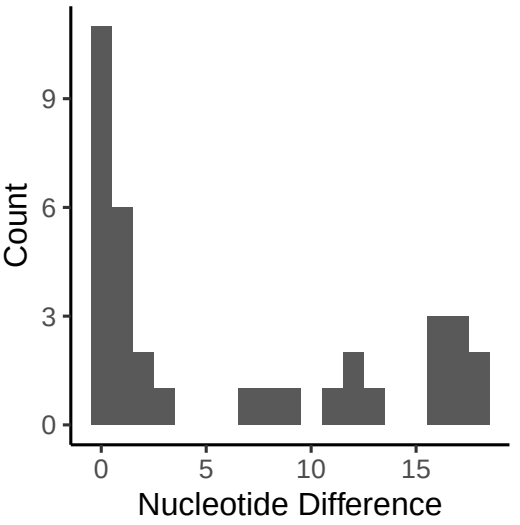
IGHV4−38−2*01

14 sequences assigned
4 (28.6%) exact matches, in which:
4 unique CDR3
2 unique J



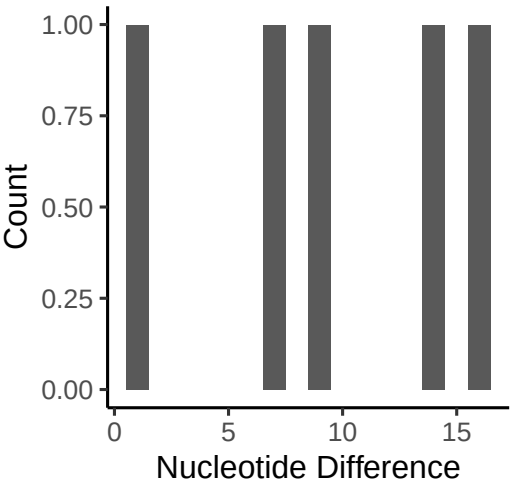
IGHV4−59*01

40 sequences assigned
11 (27.5%) exact matches, in which:
11 unique CDR3
4 unique J



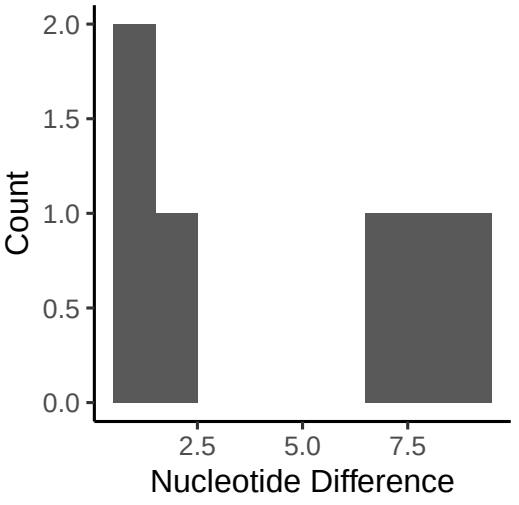
IGHV7−4−1*02

13 sequences assigned
No exact matches.



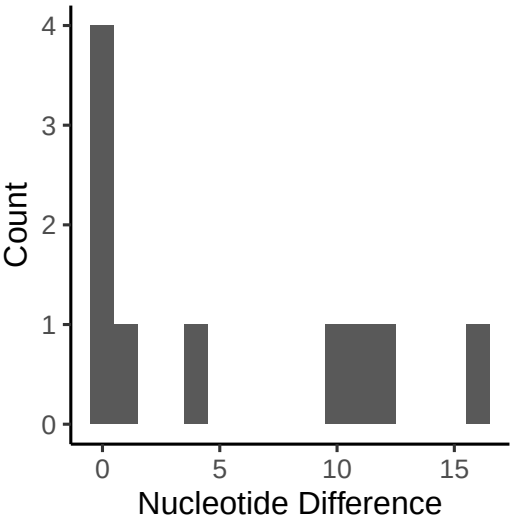
IGHV4−38−2*02

8 sequences assigned
No exact matches.



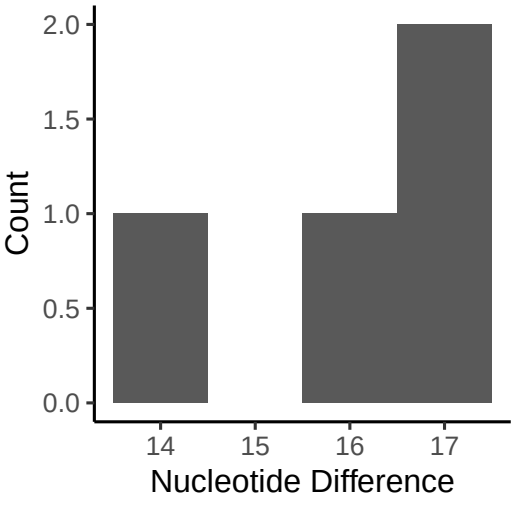
IGHV4−61*01

17 sequences assigned
4 (23.5%) exact matches, in which:
4 unique CDR3
3 unique J



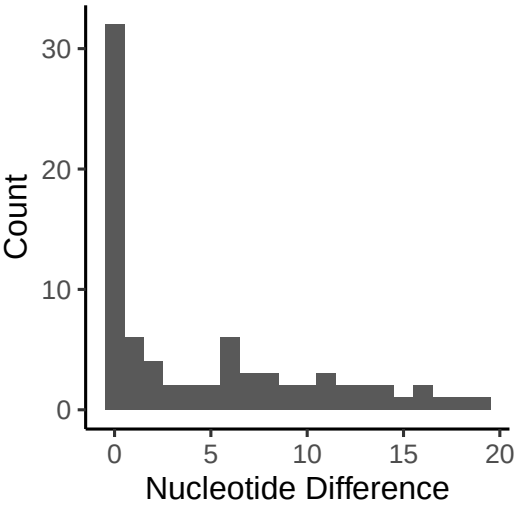
IGHV4−39*01

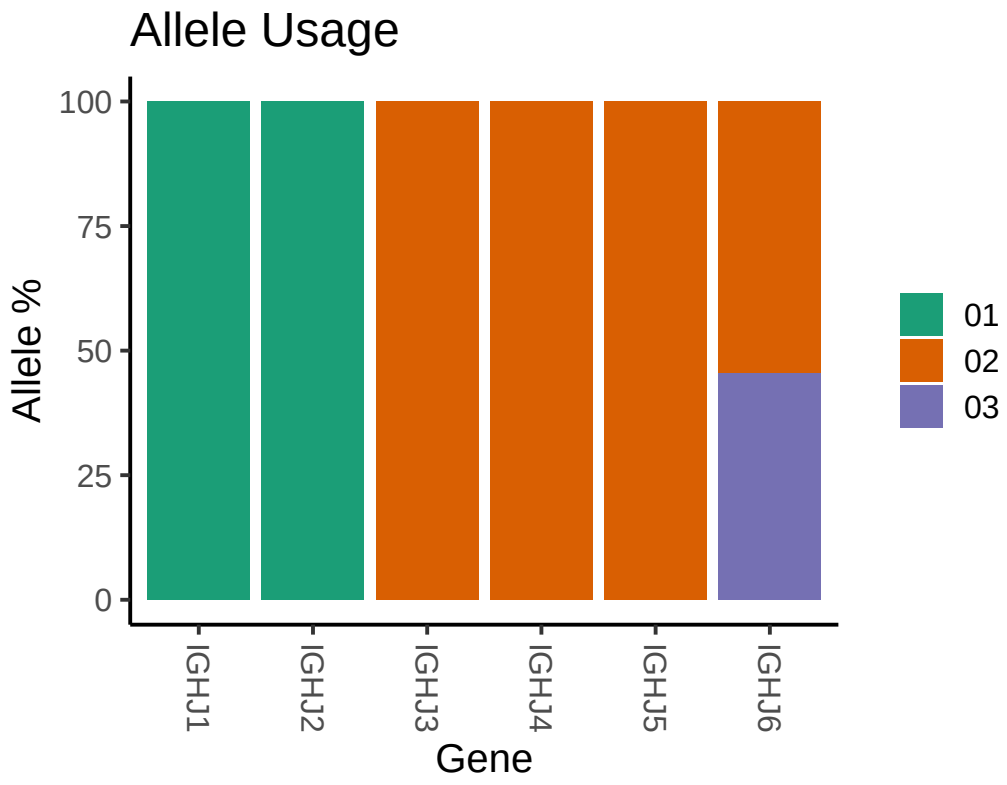
9 sequences assigned
No exact matches.



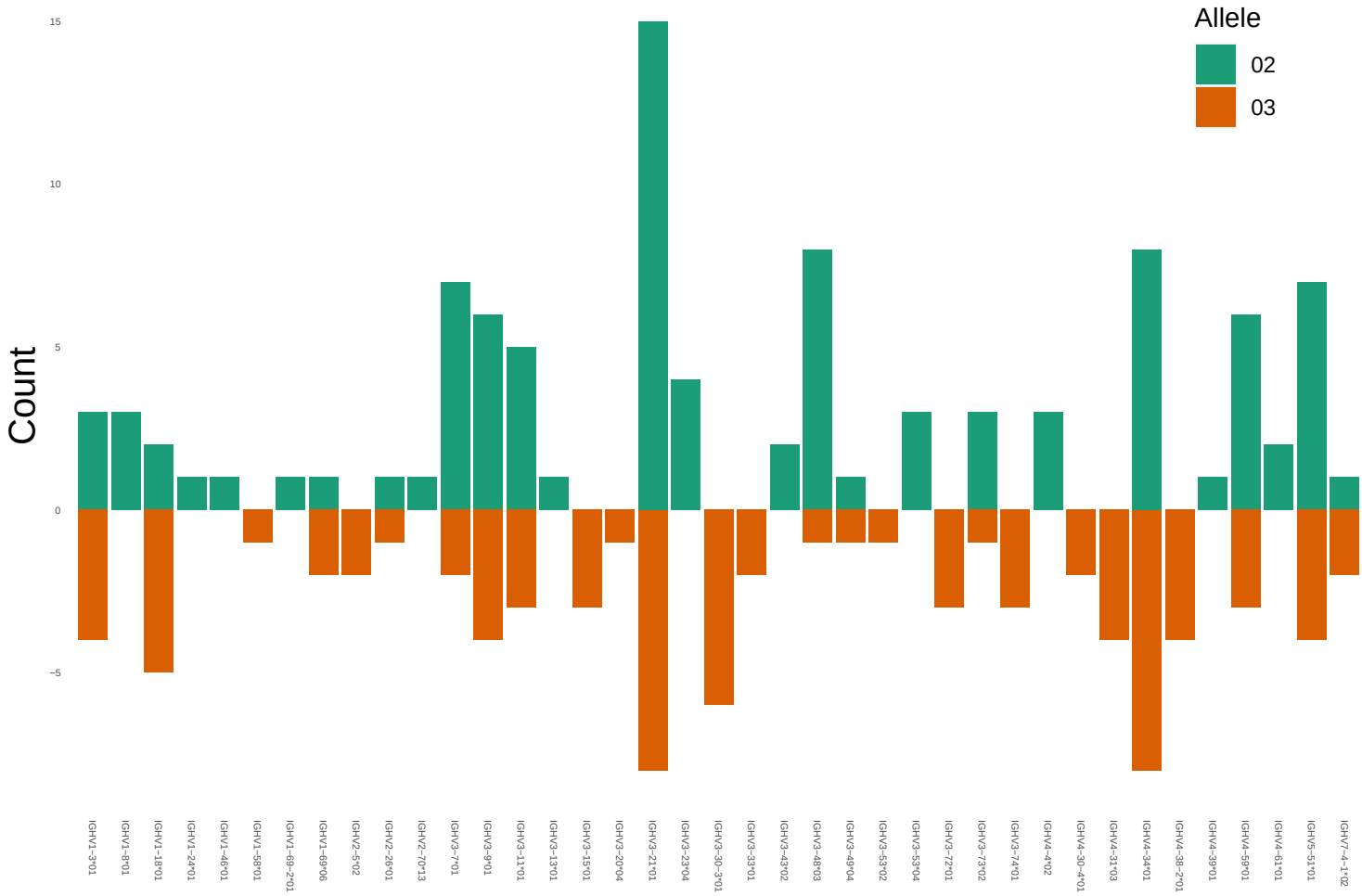
IGHV5−51*01

91 sequences assigned
32 (35.2%) exact matches, in which:
32 unique CDR3
6 unique J




Allele Usage
100
75
50
25
0
Allele %
IGHJ1
IGHJ2
IGHJ3
IGHJ4
IGHJ5
IGHJ6
Gene
01
02
03

Sequence Count by IGHJ6 allele usage



Warning − no inferred sequences found.

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Warning − no inferred sequences found.

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