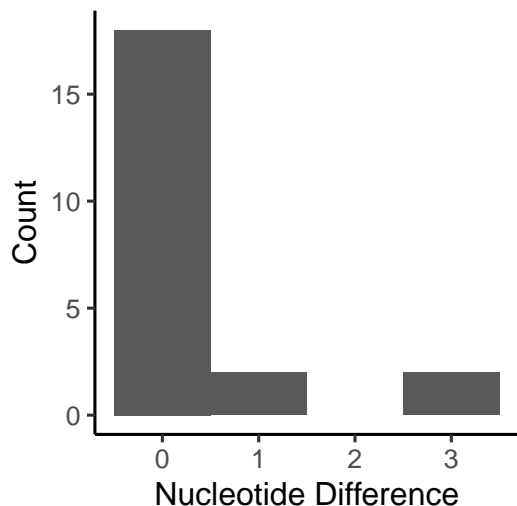


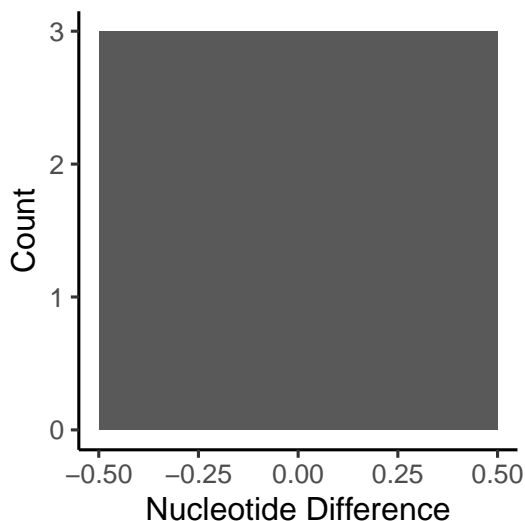
IGLV1-40*01

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



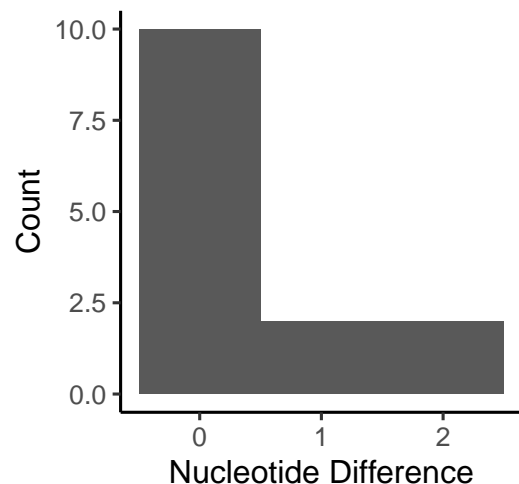
IGLV1-51*01

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



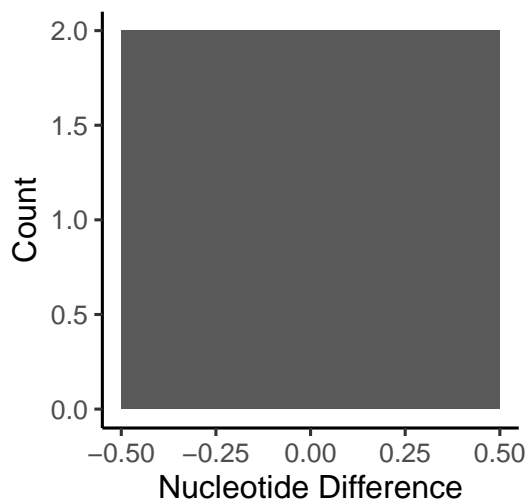
IGLV2-14*01

14 sequences assigned
10 (71.4%) exact matches, in which:
10 unique CDR3
3 unique J



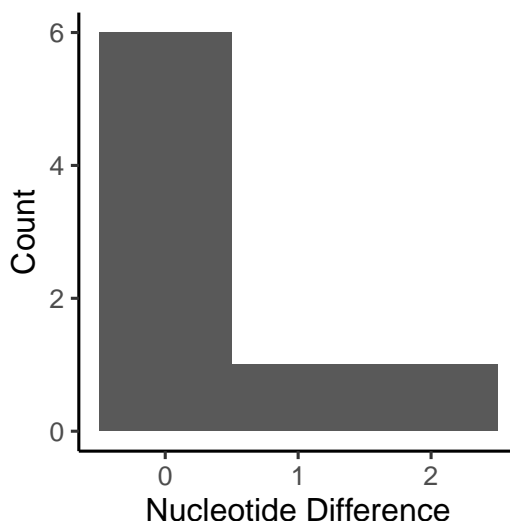
IGLV1-44*01

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



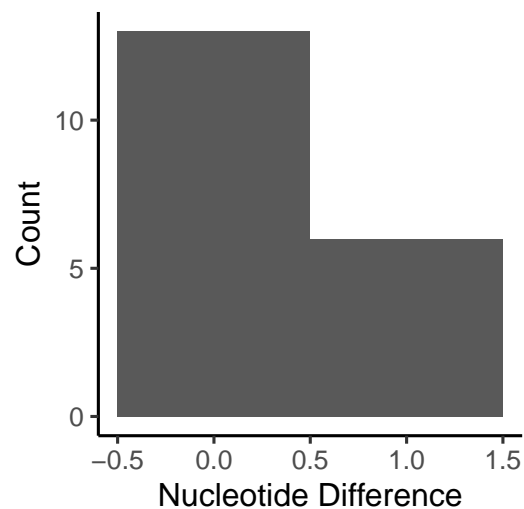
IGLV2-8*01

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



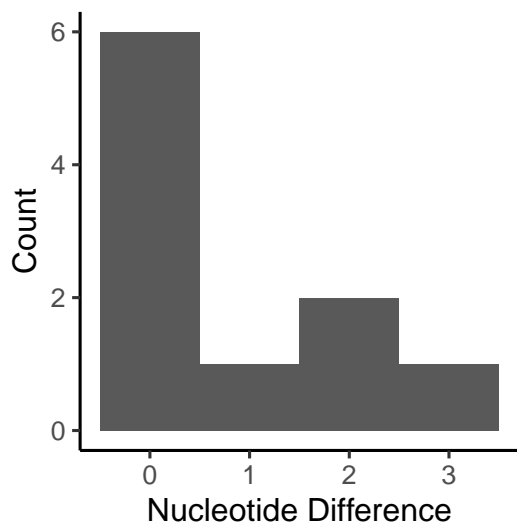
IGLV2-14*04EXT3

19 sequences assigned
13 (68.4%) exact matches, in which:
11 unique CDR3
3 unique J



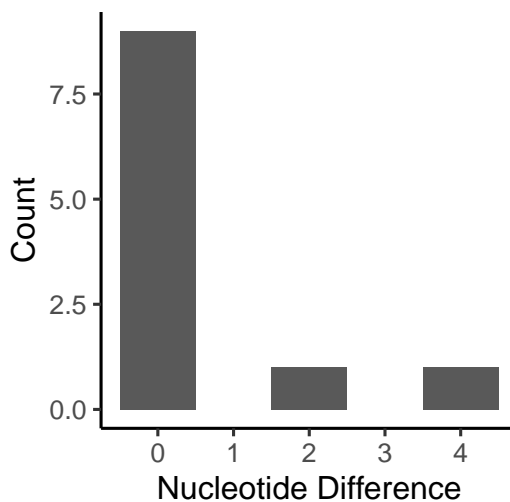
IGLV1-47*01

10 sequences assigned
6 (60%) exact matches, in which:
5 unique CDR3
2 unique J



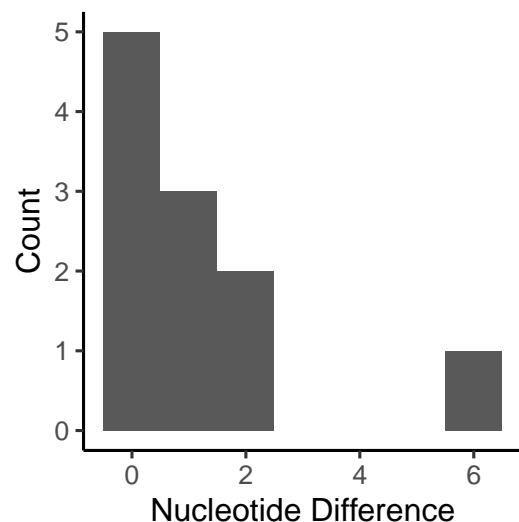
IGLV2-11*01

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



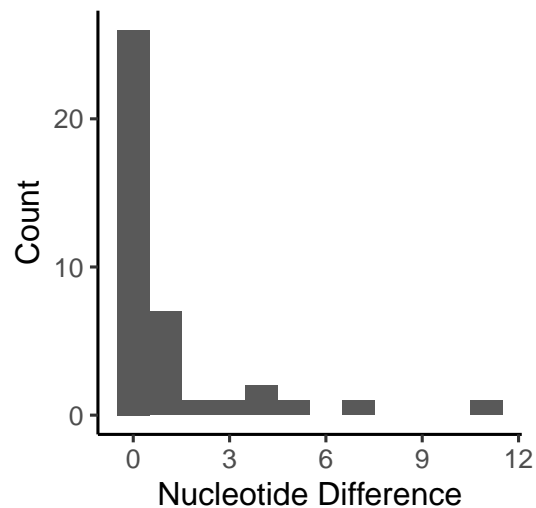
IGLV2-23*01

11 sequences assigned
5 (45.5%) exact matches, in which:
5 unique CDR3
3 unique J



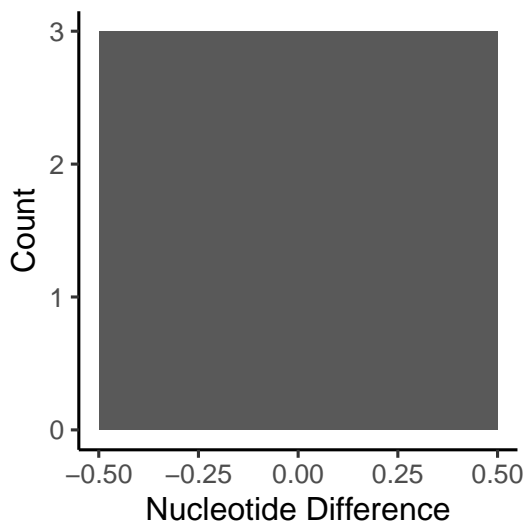
IGLV3-1*01

40 sequences assigned
26 (65%) exact matches, in which:
18 unique CDR3
2 unique J



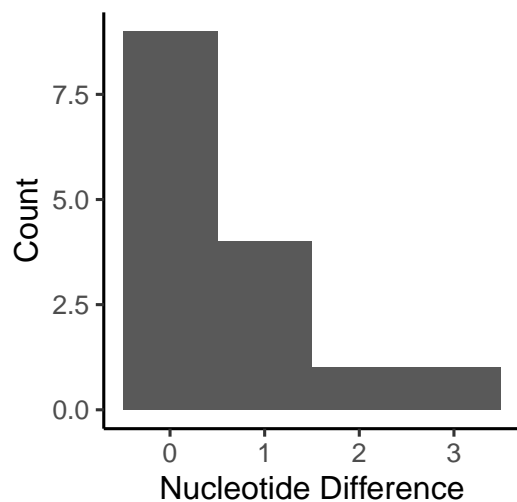
IGLV3-16*01

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



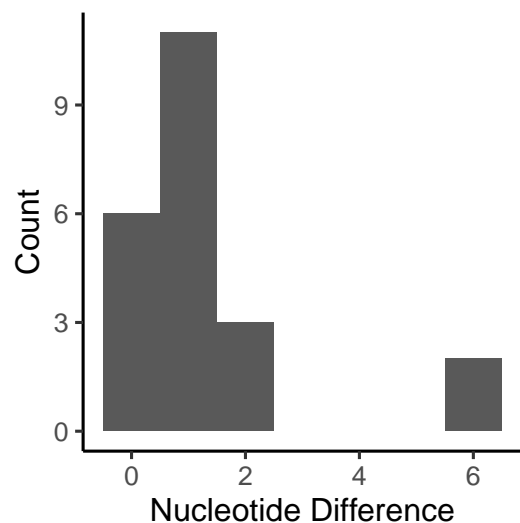
IGLV3-21*04

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



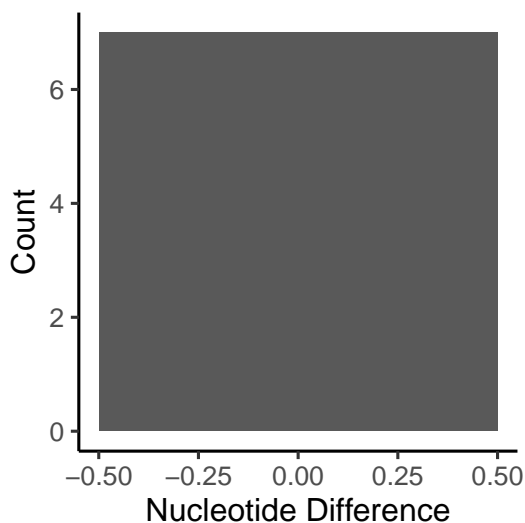
IGLV3-1*01_G153A_G194L

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



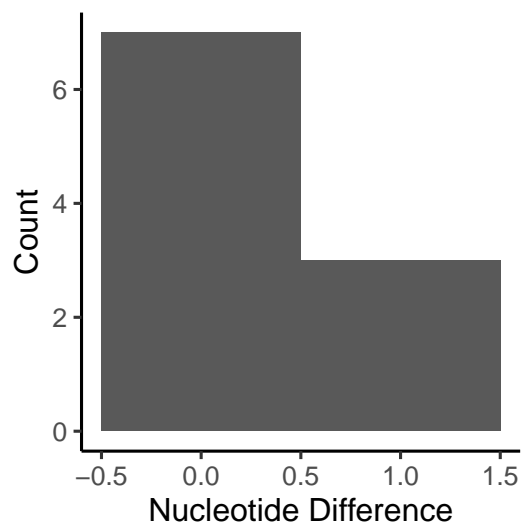
IGLV3-19*01

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



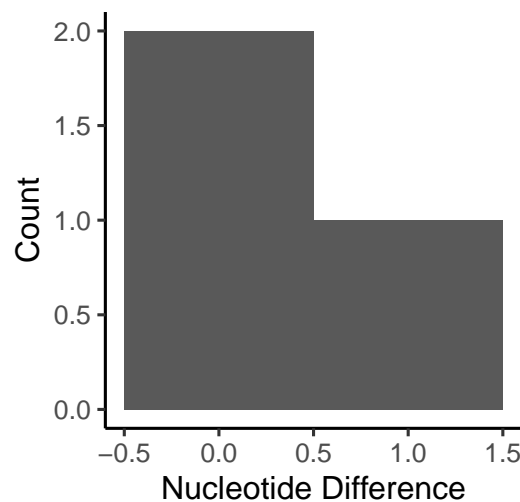
IGLV3-25*03

10 sequences assigned
7 (70%) exact matches, in which:
5 unique CDR3
3 unique J



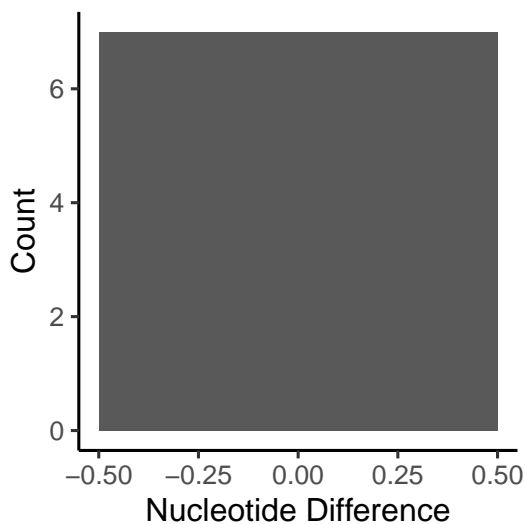
IGLV3-10*01

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



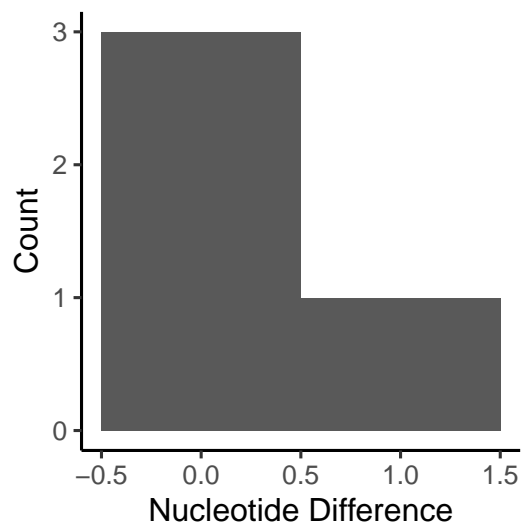
IGLV3-21*02

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



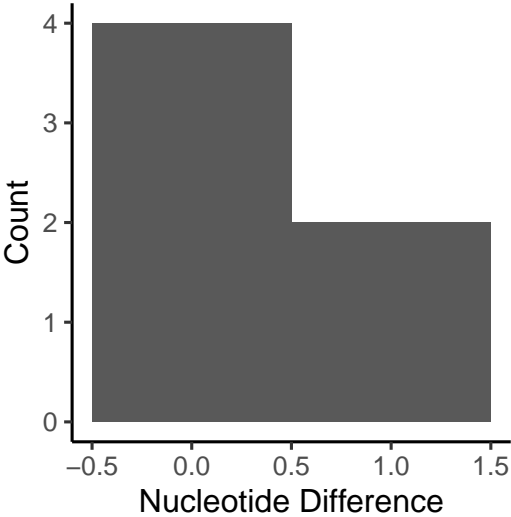
IGLV4-69*01

4 sequences assigned
3 (75%) exact matches, in which:
2 unique CDR3
2 unique J



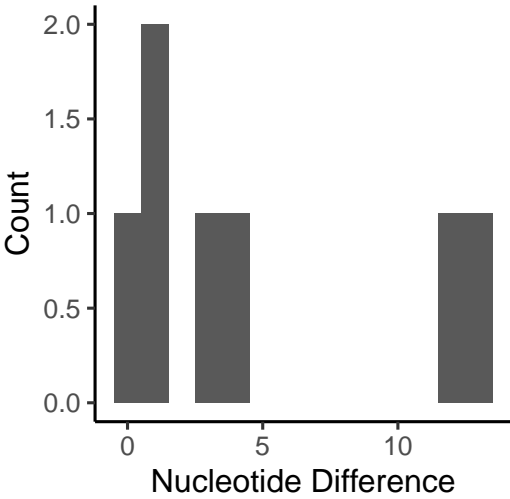
IGLV5-45*01

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



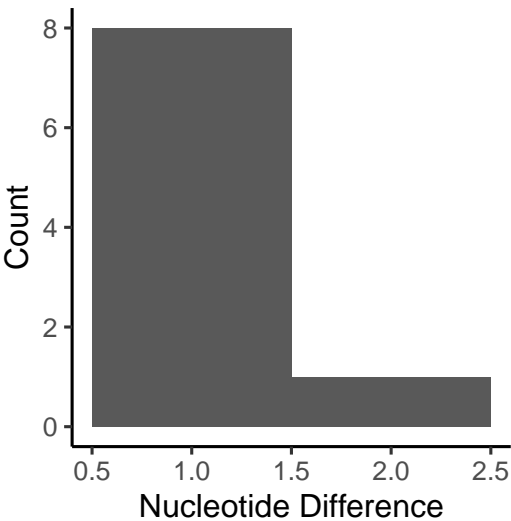
IGLV7-46*01

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



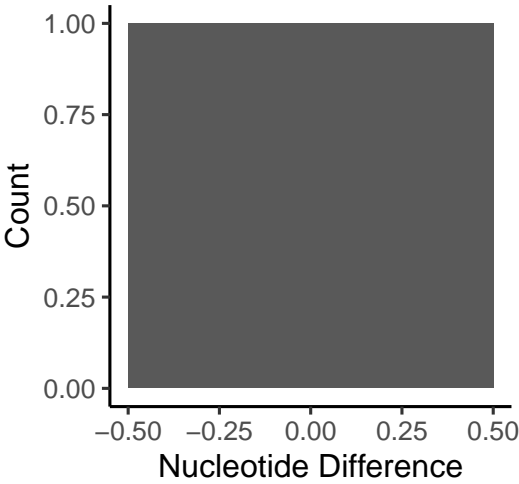
IGLV6-57*03

9 sequences assigned
No exact matches.



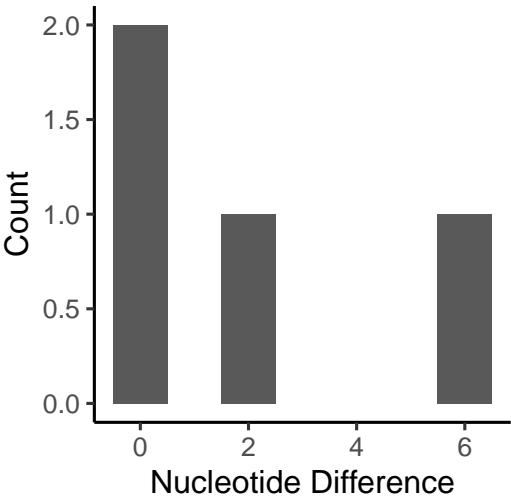
IGLV8-61*01

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



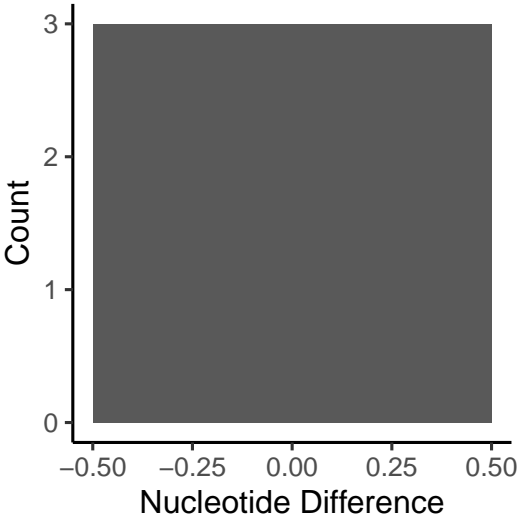
IGLV7-43*01

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

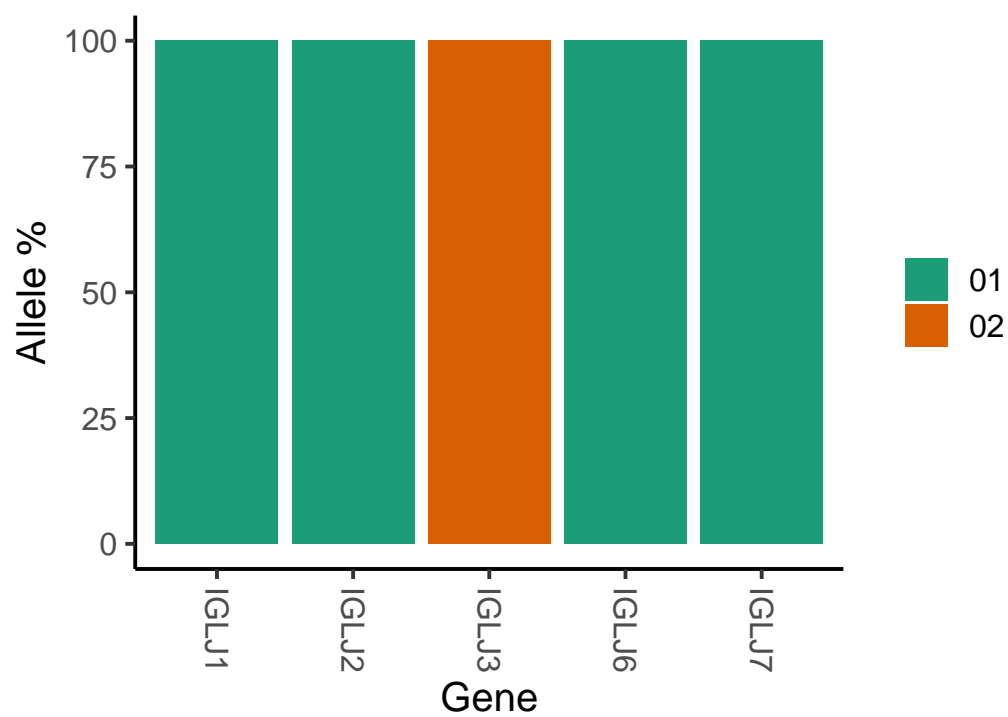


IGLV9-49*01

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



Allele Usage



Warning – no inferred sequences found.