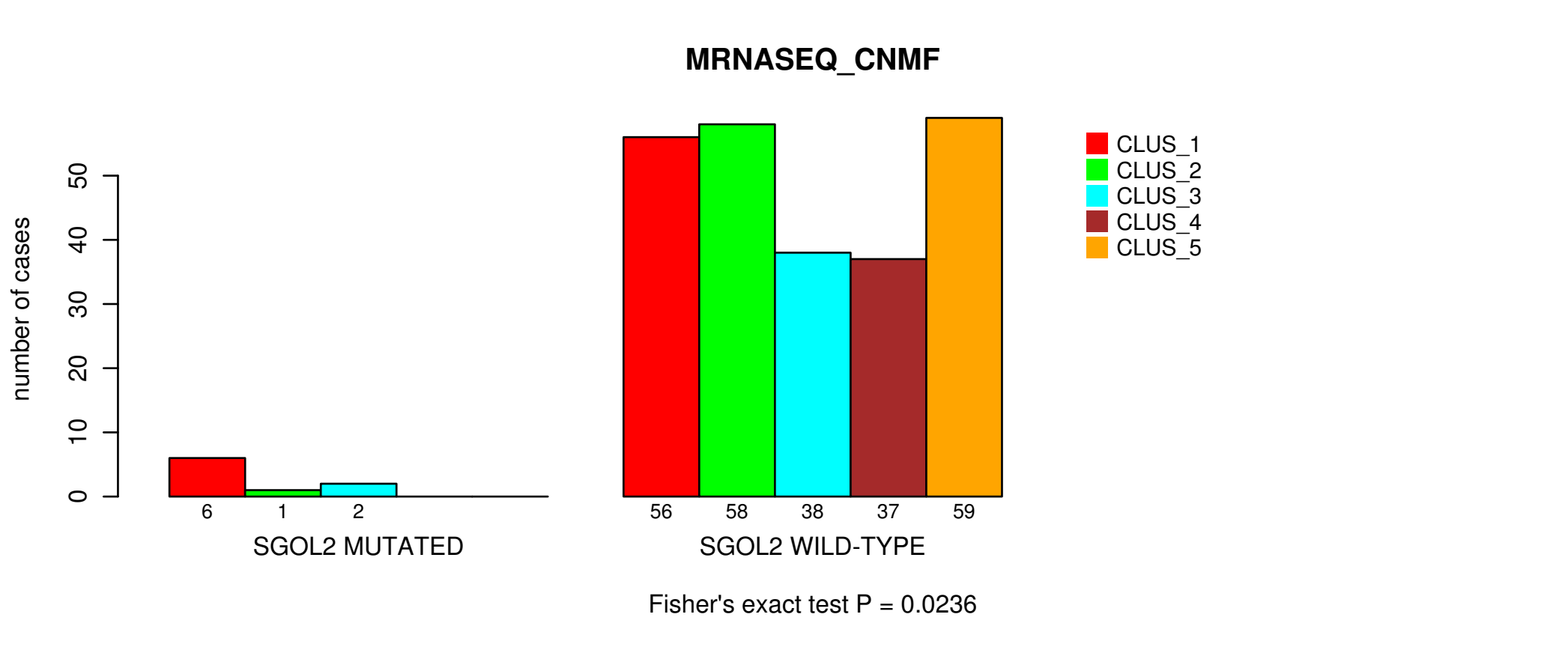
MRNASEQ_CNMF
number of cases
0
10
20
30
40
50
6
1
2
56
58
38
37
59
SGOL2 MUTATED
SGOL2 WILD-TYPE
CLUS_1
CLUS_2
CLUS_3
CLUS_4
CLUS_5
Fisher's exact test P = 0.0236