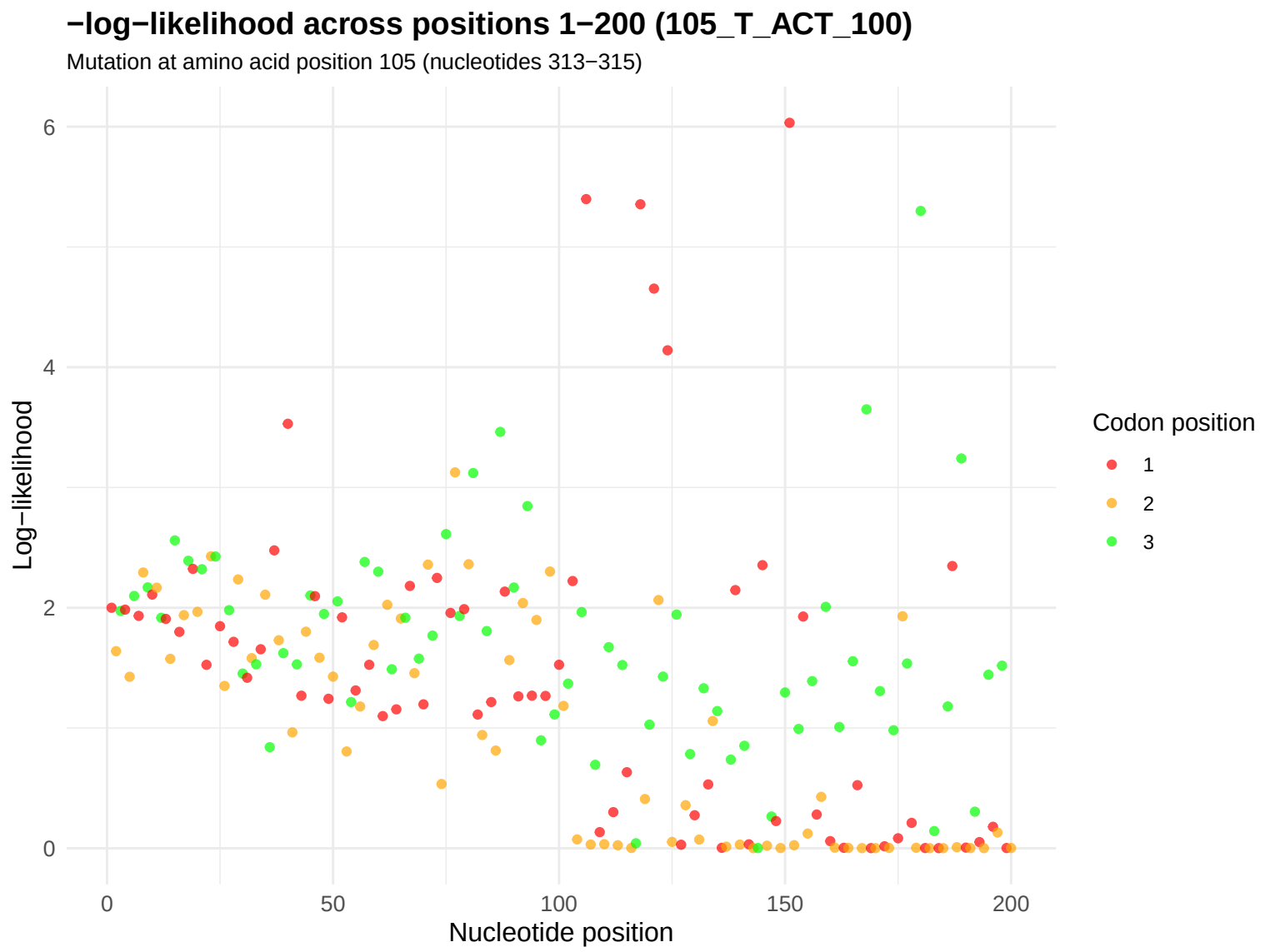
−log−likelihood across positions 1−200 (105_T_ACT_100)
Mutation at amino acid position 105 (nucleotides 313−315)
Log−likelihood
Nucleotide position
0
2
4
6
0
50
100
150
200
Codon position
1
2
3