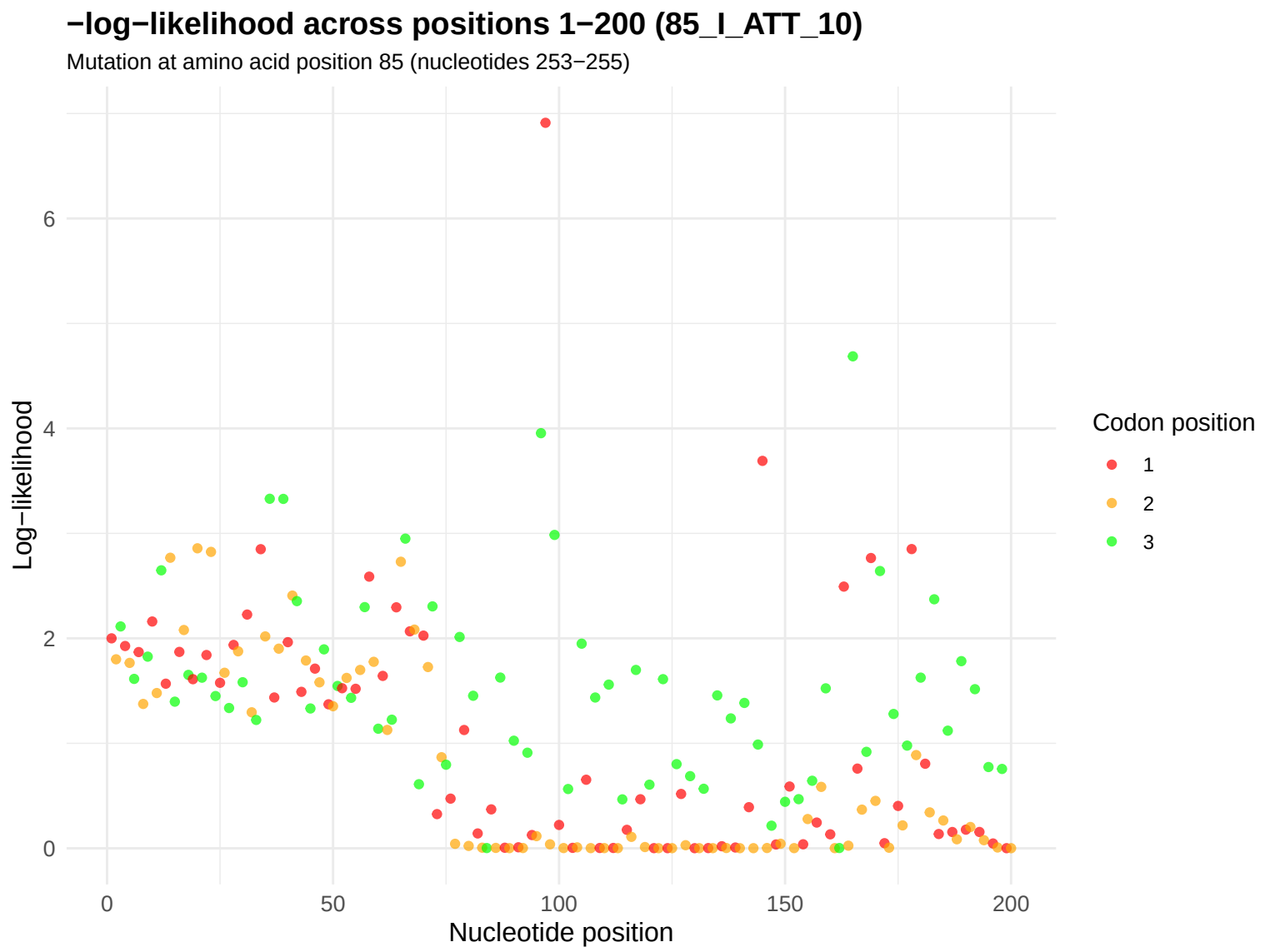
−log−likelihood across positions 1−200 (85_I_ATT_10)
Mutation at amino acid position 85 (nucleotides 253−255)
Log−likelihood
0
2
4
6
0
50
100
150
200
Nucleotide position
Codon position
1
2
3