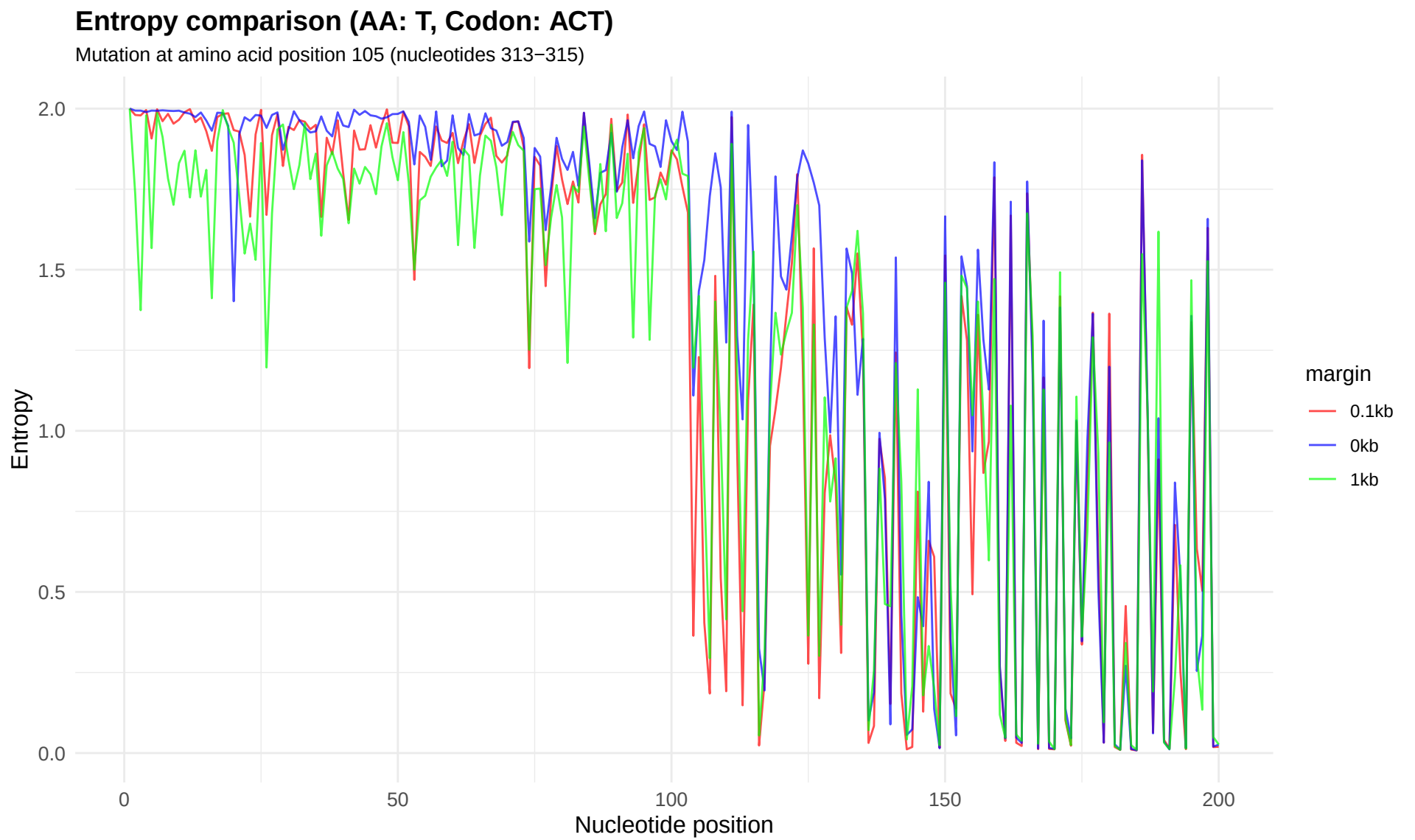
Entropy comparison (AA: T, Codon: ACT)
Mutation at amino acid position 105 (nucleotides 313−315)
Entropy
2.0
1.5
1.0
0.5
0.0
0
50
100
150
200
Nucleotide position
margin
0.1kb
0kb
1kb