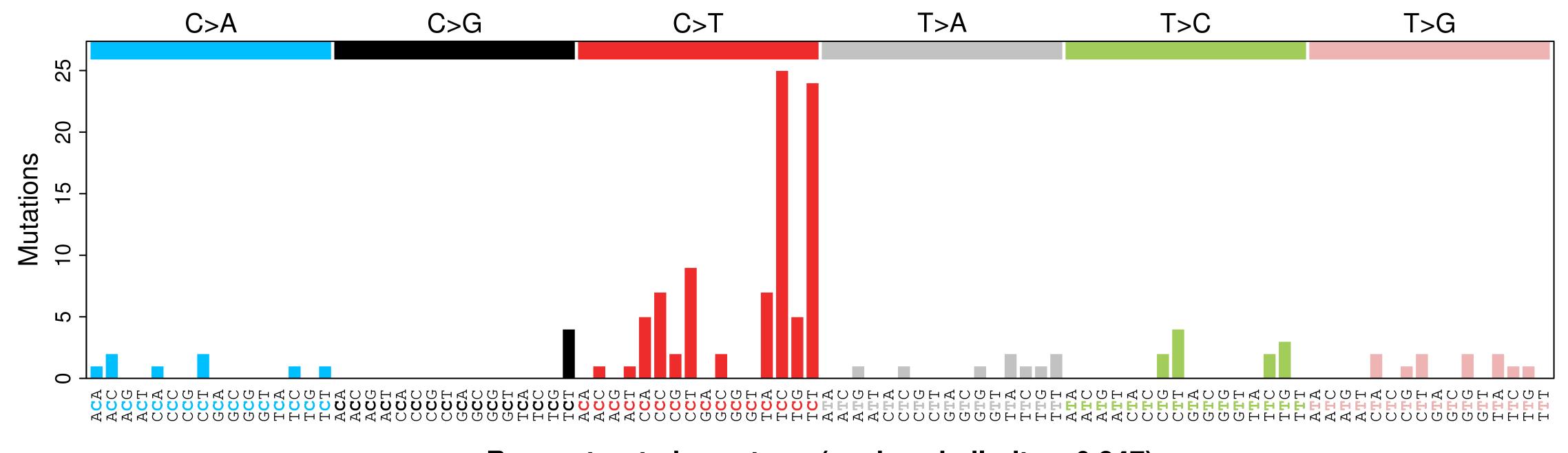
## CATD263a (131 mutations)



## Reconstructed spectrum (cosine similarity = 0.847)

