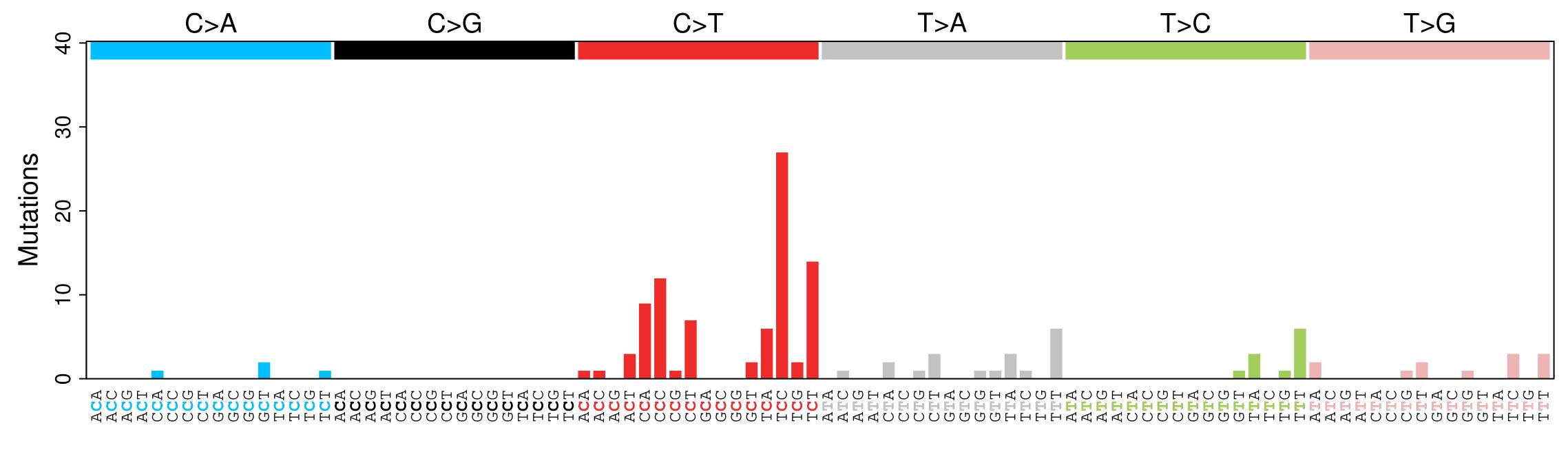
## CATD282a (131 mutations)



## Reconstructed spectrum (cosine similarity = 0.924)

