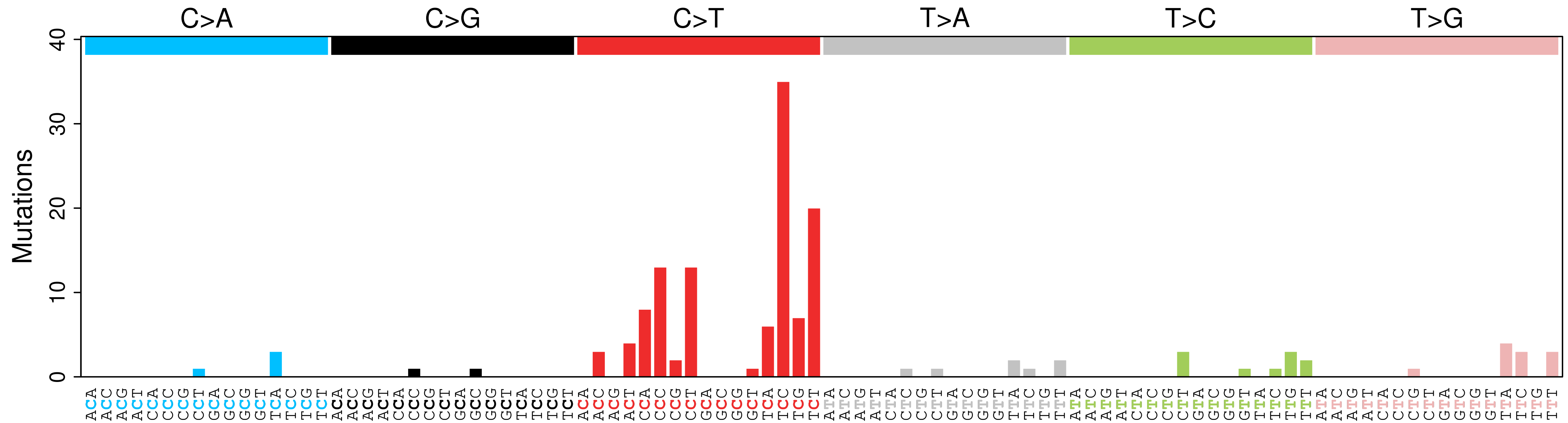


## CATD265a (146 mutations)



### Reconstructed spectrum (cosine similarity = 0.95)

