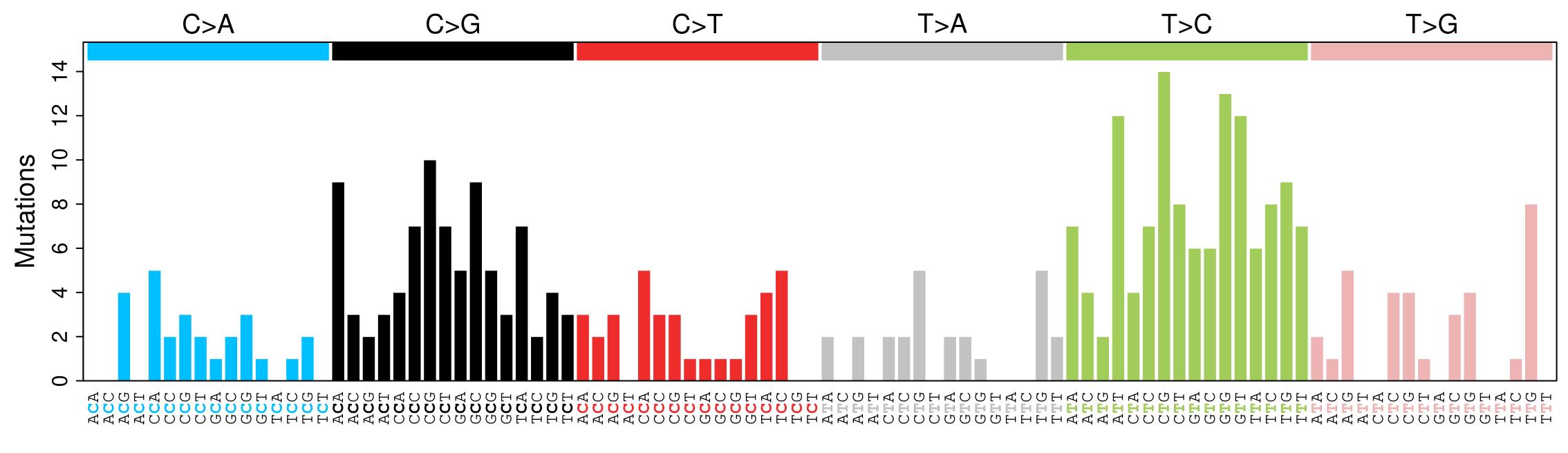
## CATD292a (327 mutations)



## Reconstructed spectrum (cosine similarity = 0.828)

