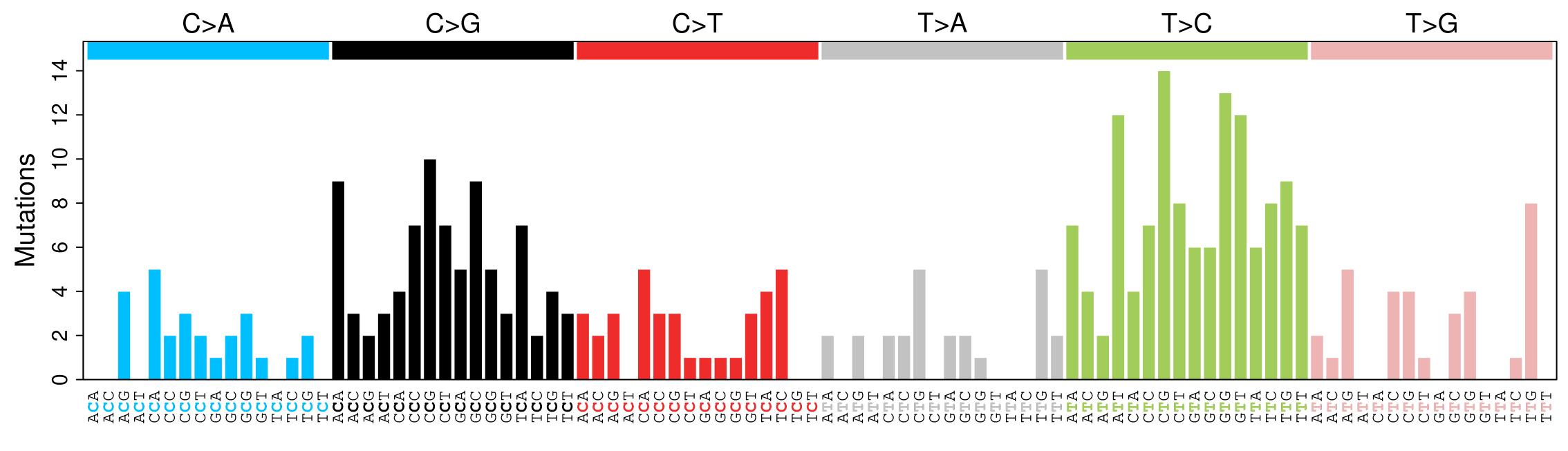
## CATD292a (327 mutations)



## **Reconstructed spectrum (cosine similarity = 0.747)**

