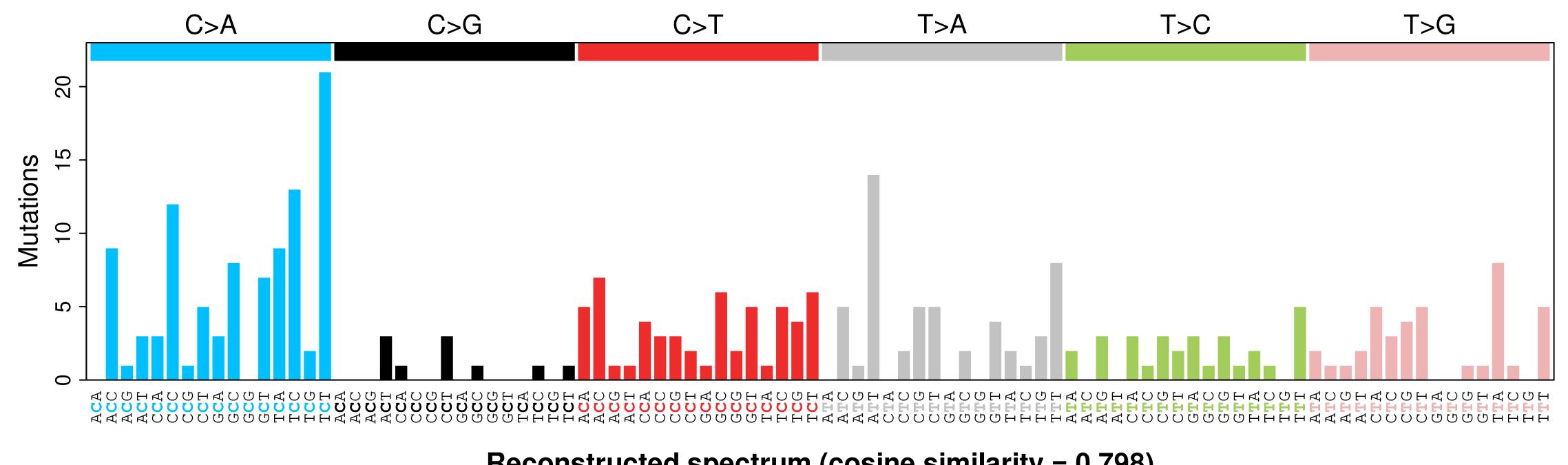
## CATD186a (284 mutations)



## Reconstructed spectrum (cosine similarity = 0.798)

