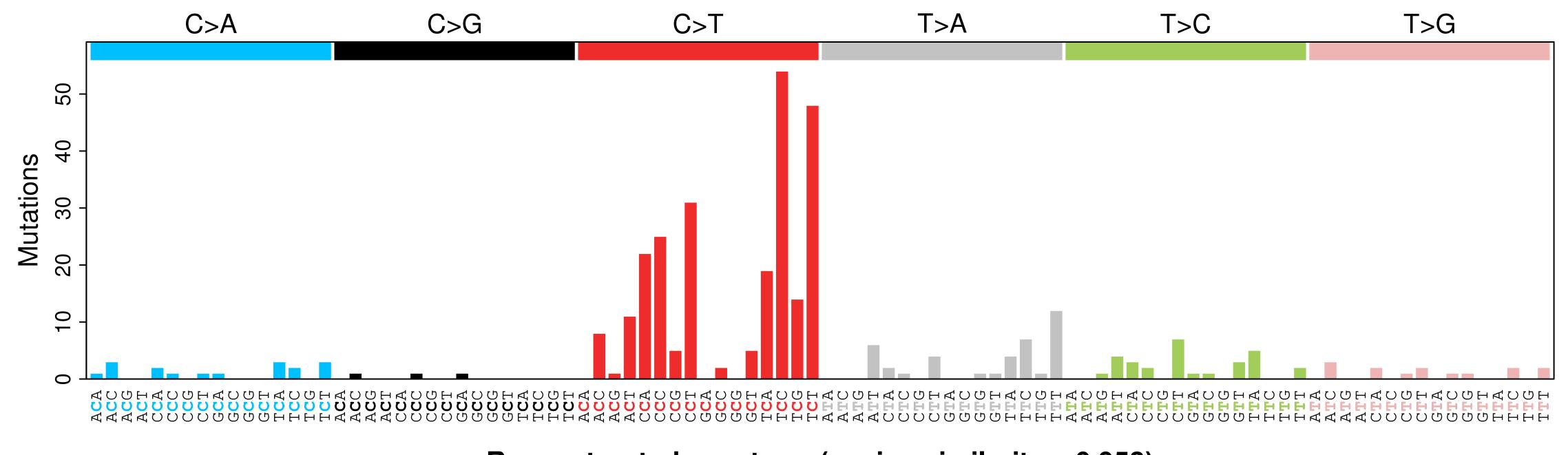
## CATD276a (347 mutations)



## Reconstructed spectrum (cosine similarity = 0.953)

