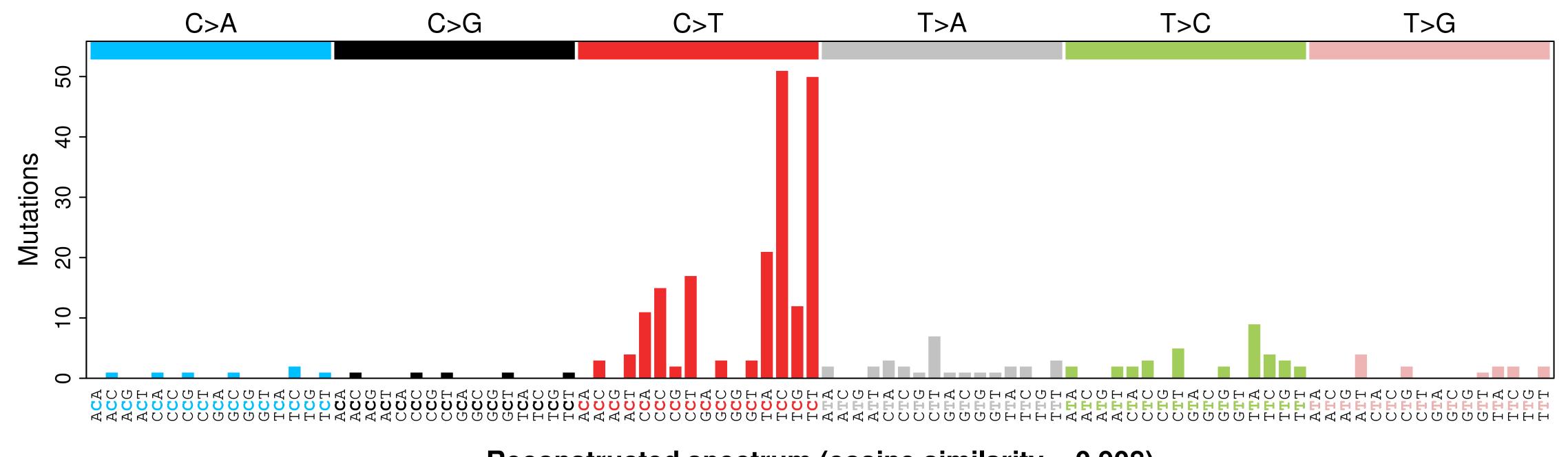
## CATD256a (279 mutations)



## Reconstructed spectrum (cosine similarity = 0.908)

