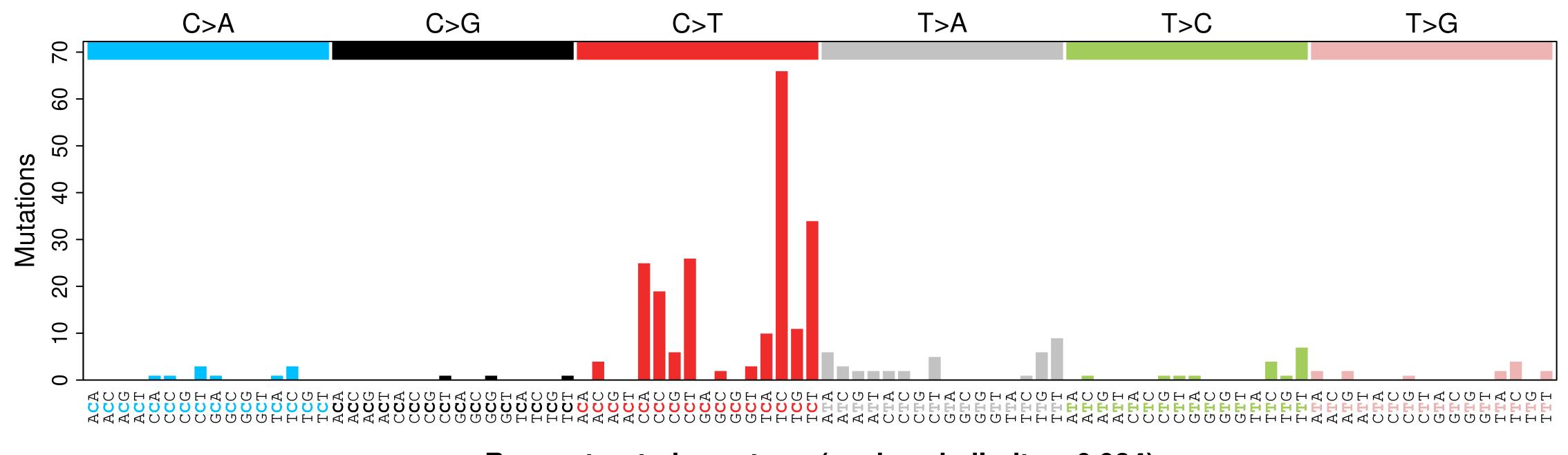
## CATD274a (286 mutations)



## Reconstructed spectrum (cosine similarity = 0.934)

