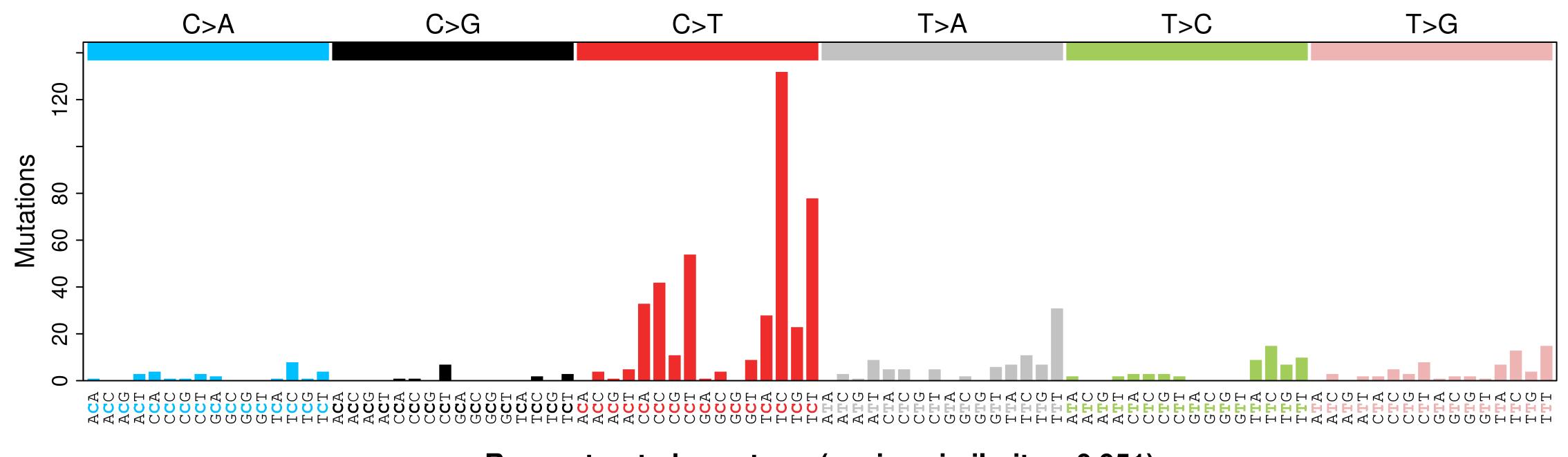
## CATD272a (684 mutations)



## **Reconstructed spectrum (cosine similarity = 0.951)**

