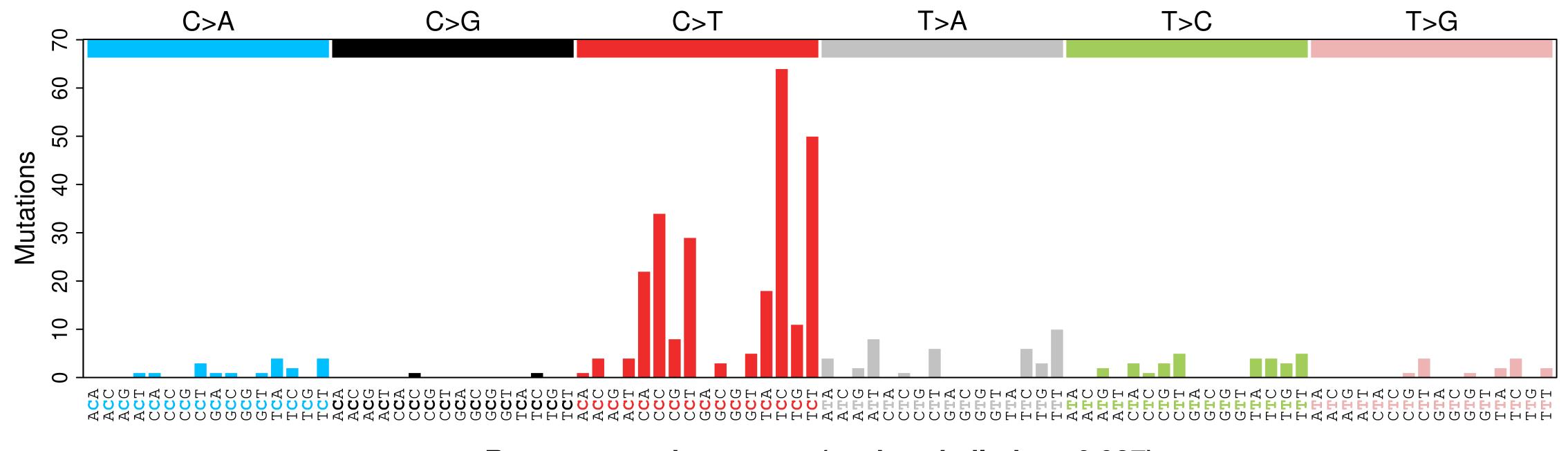
## CATD255a (357 mutations)



## **Reconstructed spectrum (cosine similarity = 0.937)**

