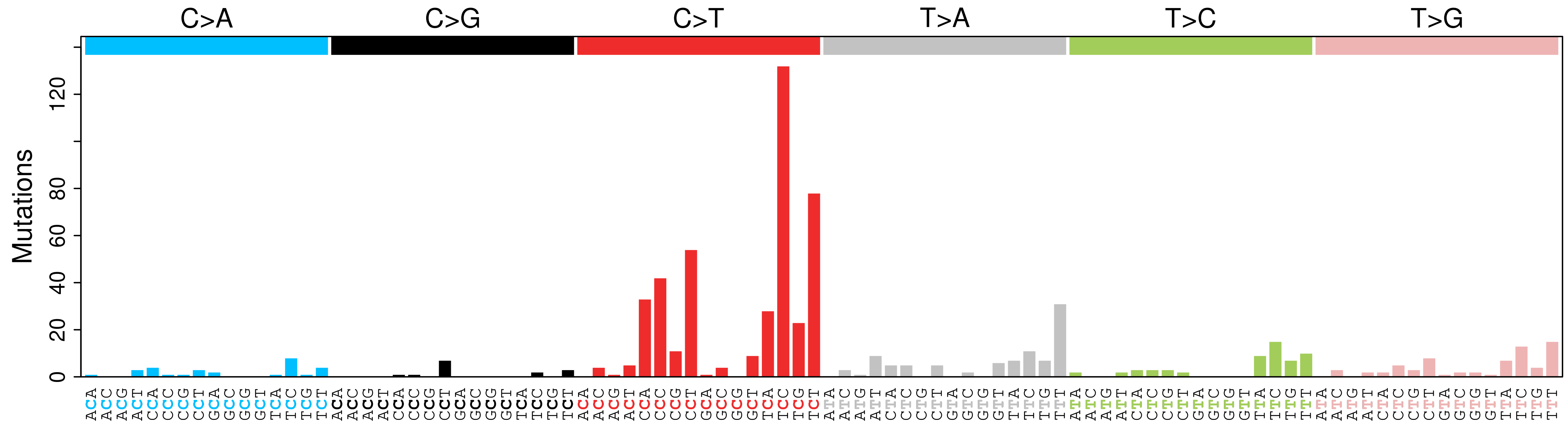


# CATD272a (684 mutations)



## Reconstructed spectrum (cosine similarity = 0.94)

