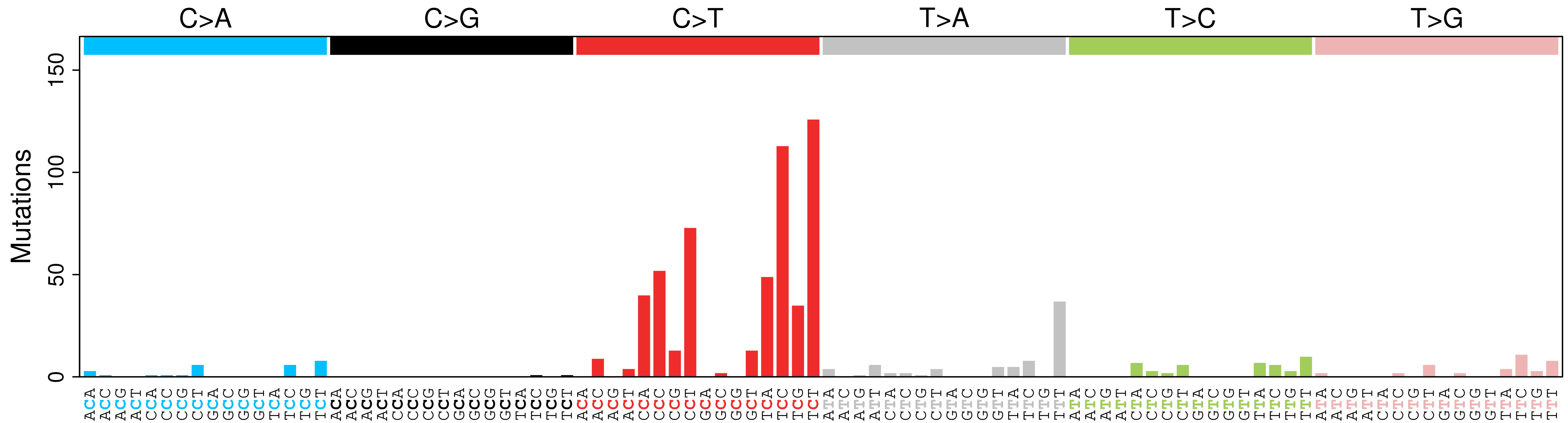


# CATD268a (715 mutations)



## Reconstructed spectrum (cosine similarity = 0.943)

