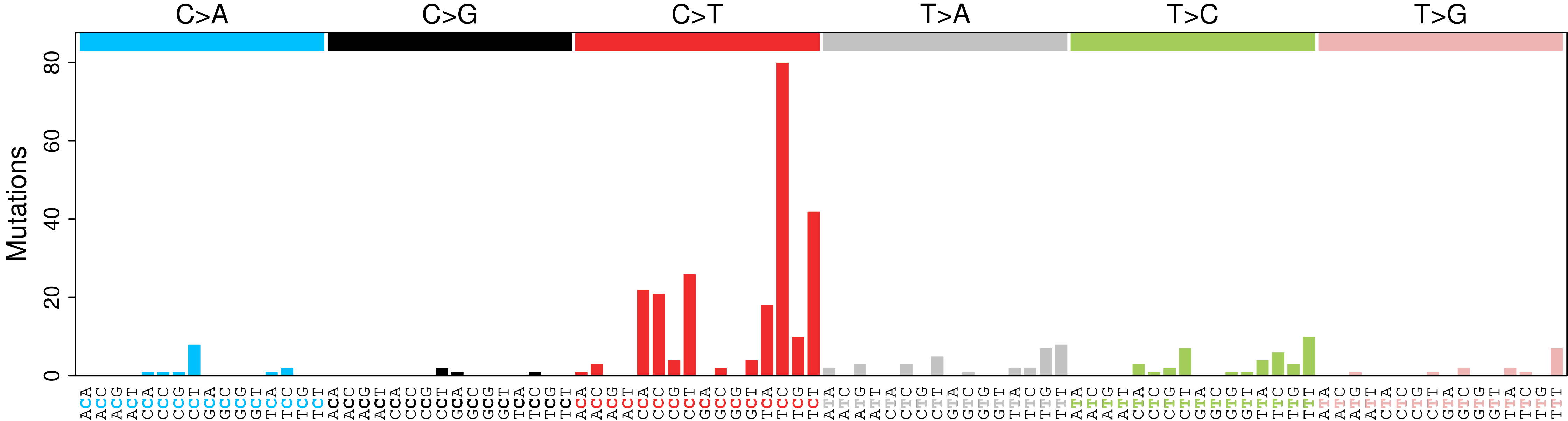


CATD250a (336 mutations)



Reconstructed spectrum (cosine similarity = 0.936)

