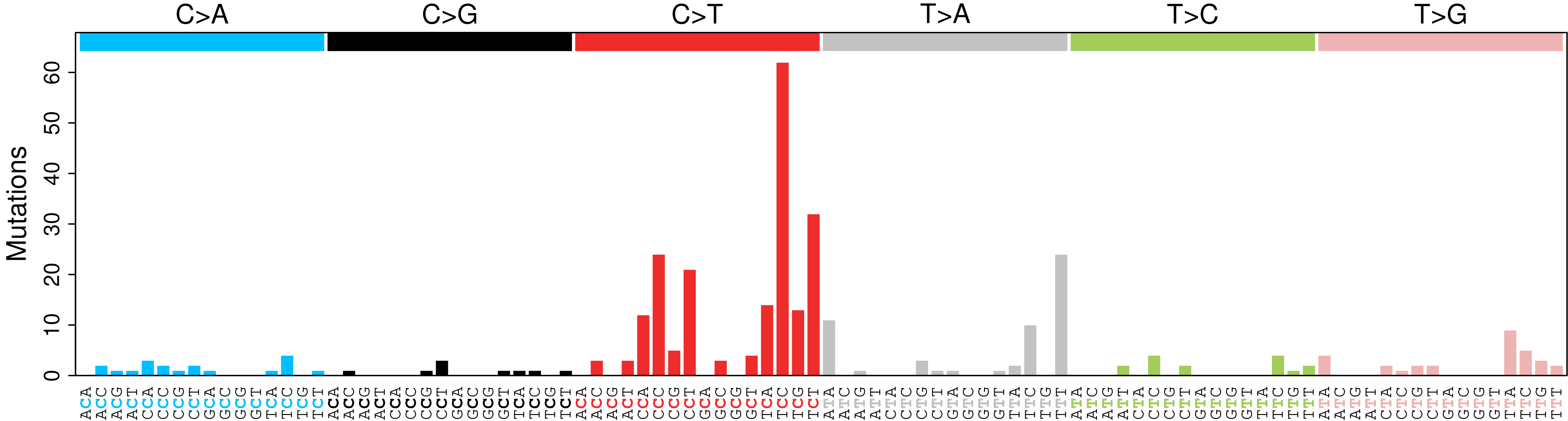


CATD262a (323 mutations)



Reconstructed spectrum (cosine similarity = 0.926)

