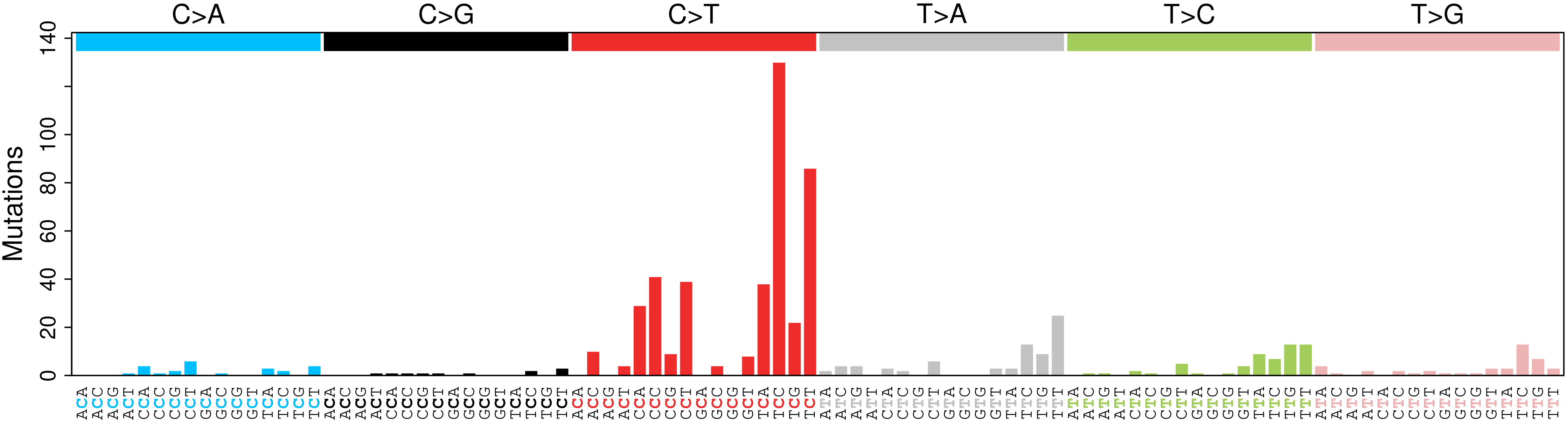


CATD264a (631 mutations)



Reconstructed spectrum (cosine similarity = 0.949)

