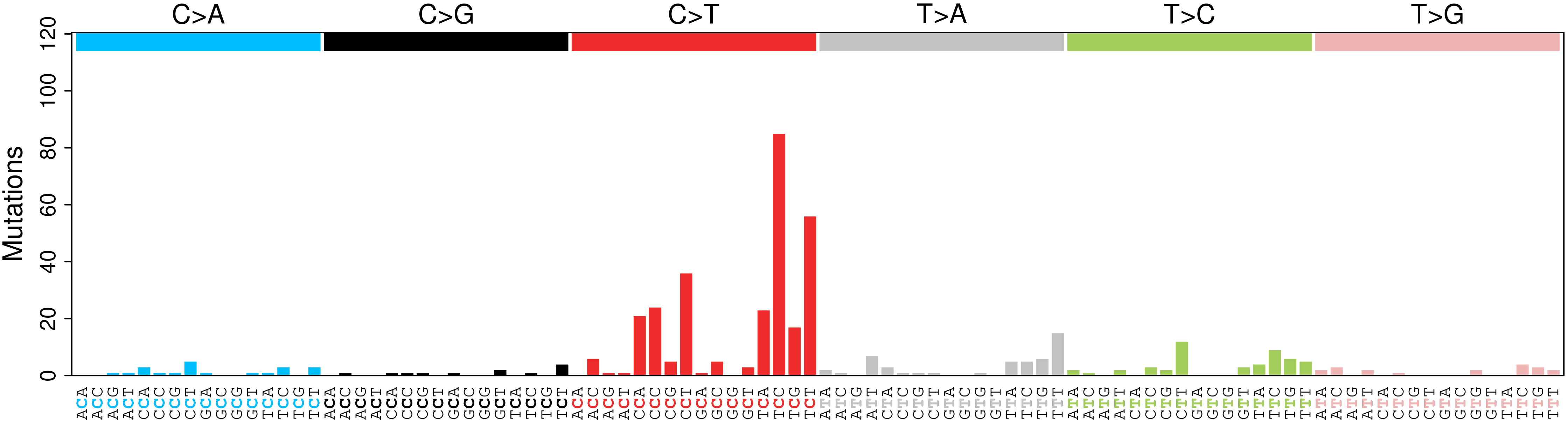


CATD0128a (433 mutations)



Reconstructed spectrum (cosine similarity = 0.932)

