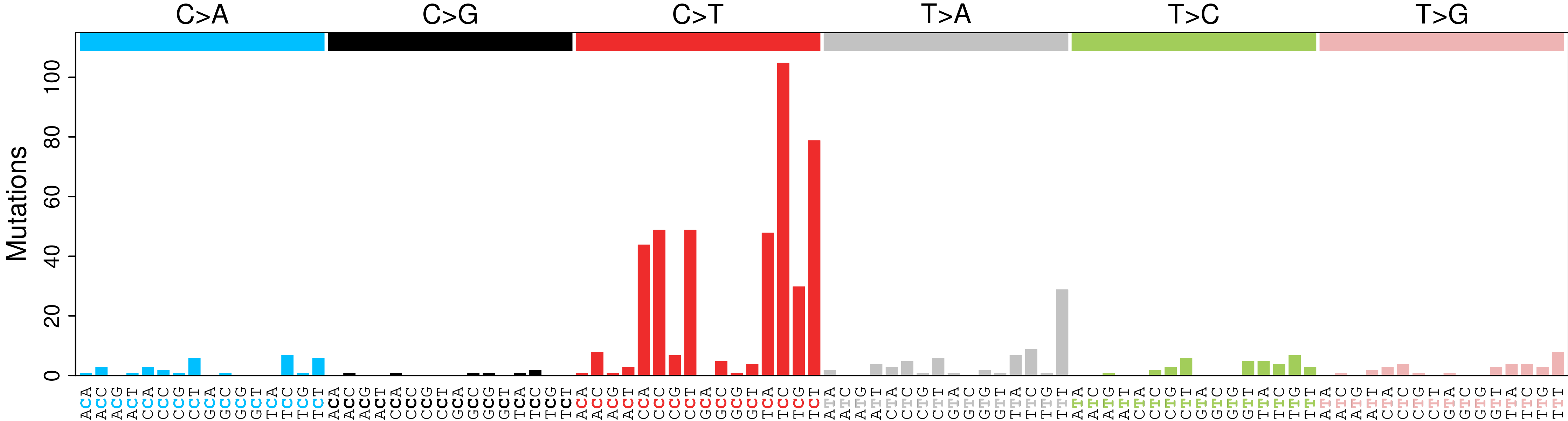


CATD0122a (614 mutations)



Reconstructed spectrum (cosine similarity = 0.971)

