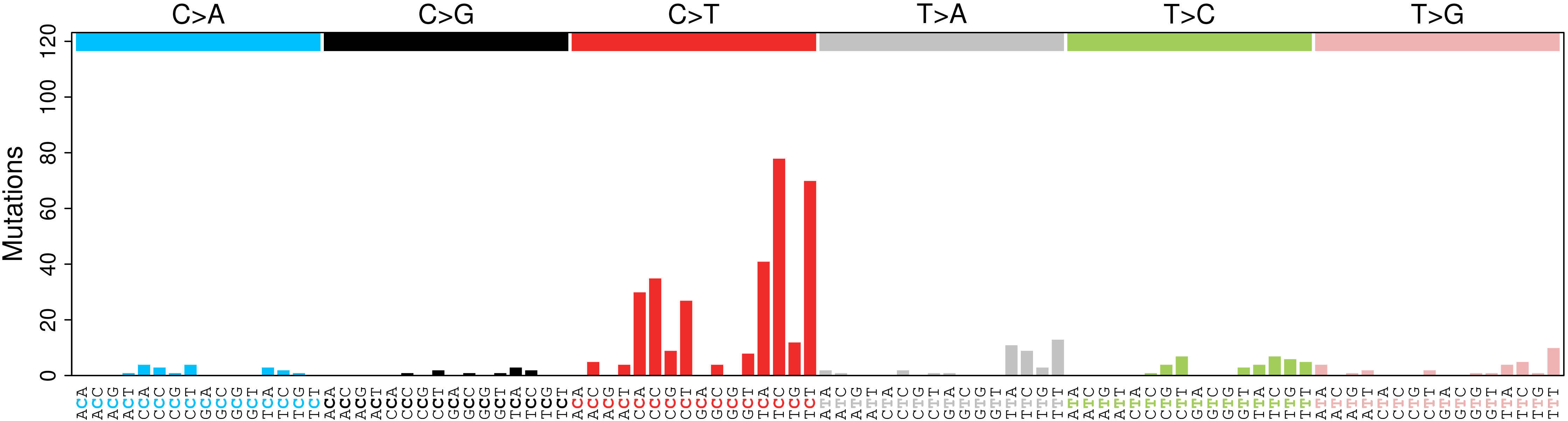CATD248a (463 mutations)

Reconstructed spectrum (cosine similarity = 0.946)

SBS2 (0.11)
SBS7a (0.198)
SBS7b (0.692)