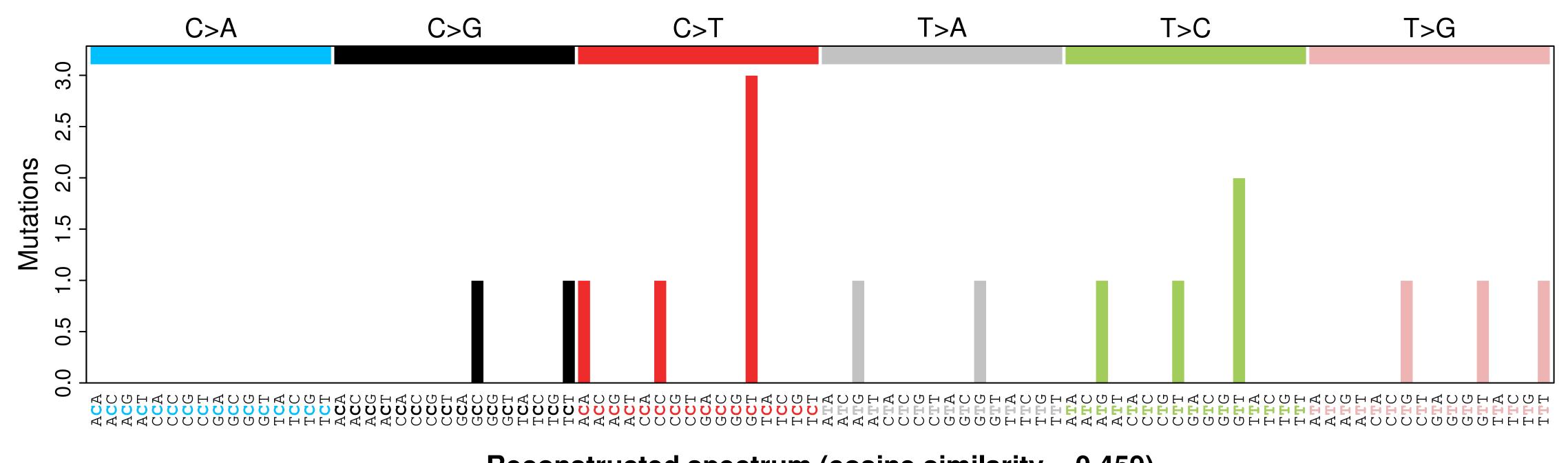
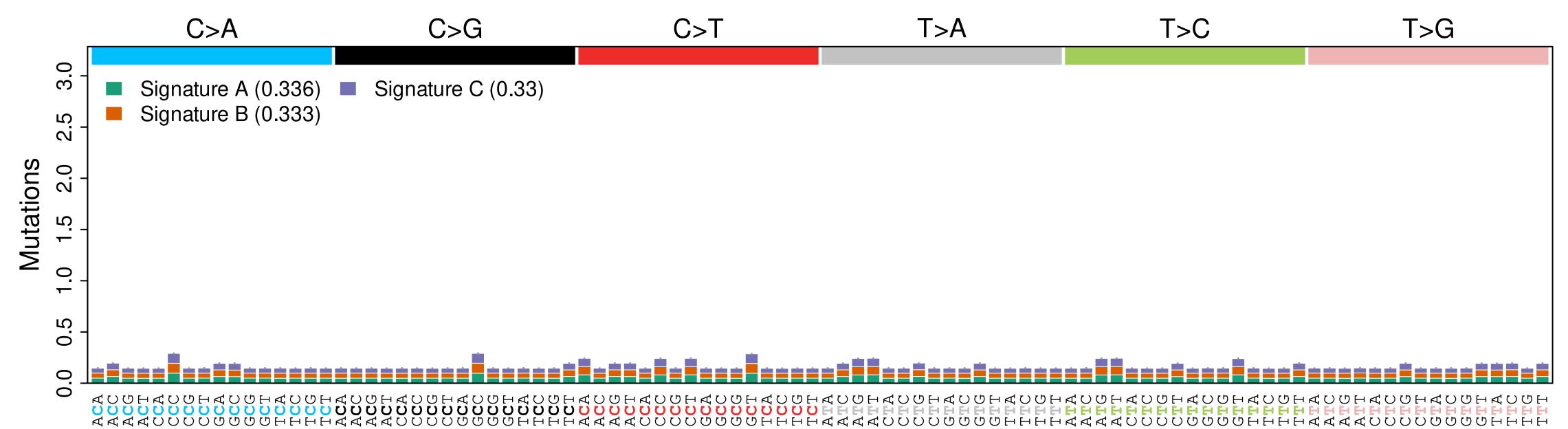
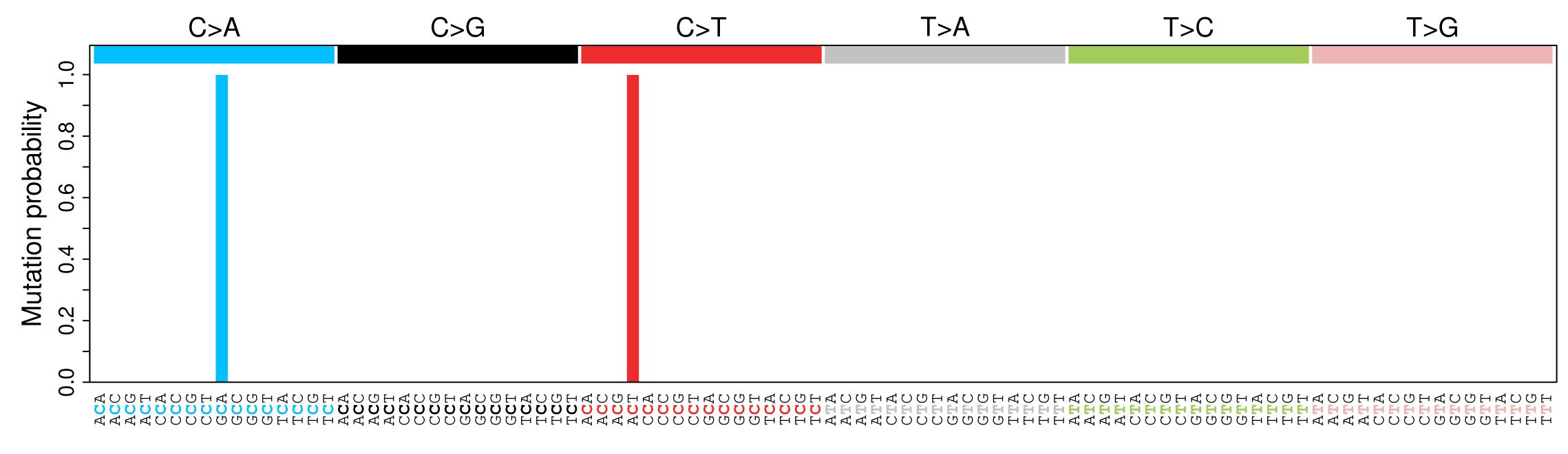
CATD0569a (16 mutations)



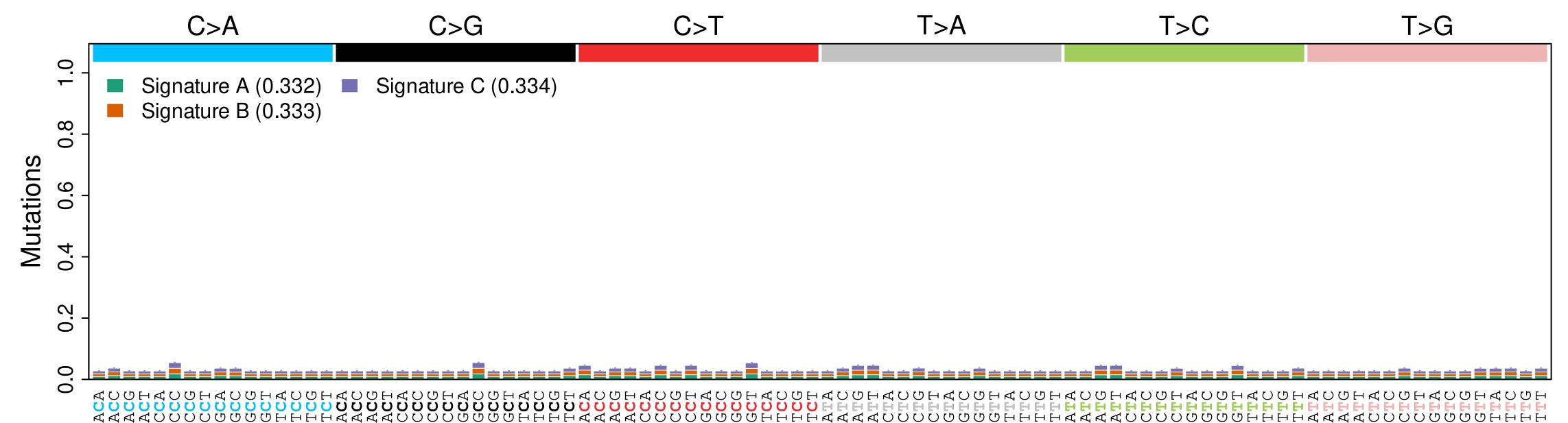
Reconstructed spectrum (cosine similarity = 0.459)



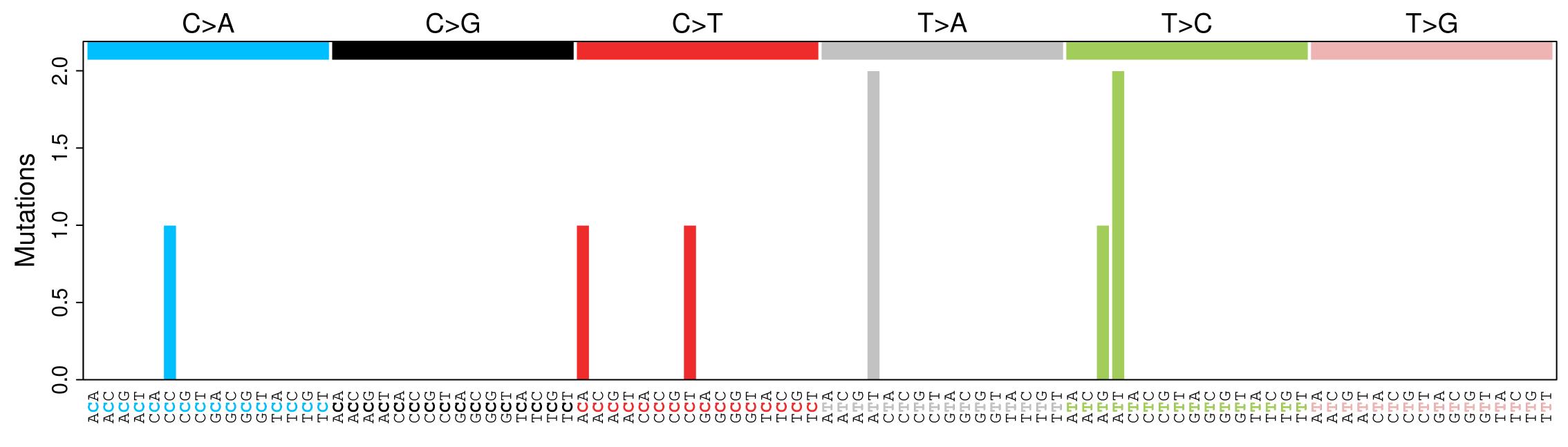




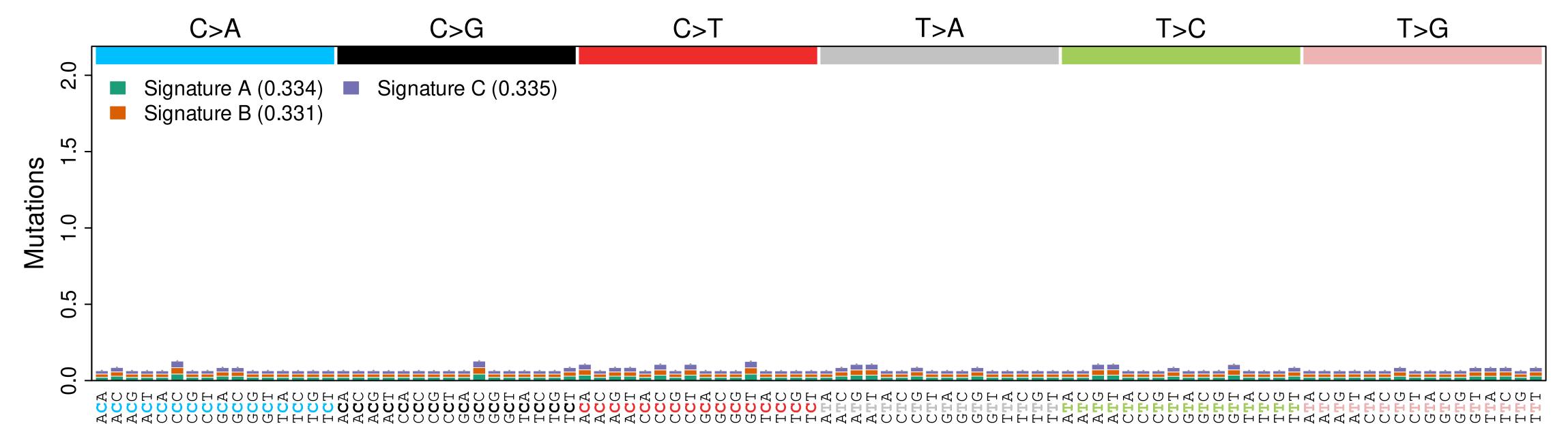
Reconstructed spectrum (cosine similarity = 0.165)



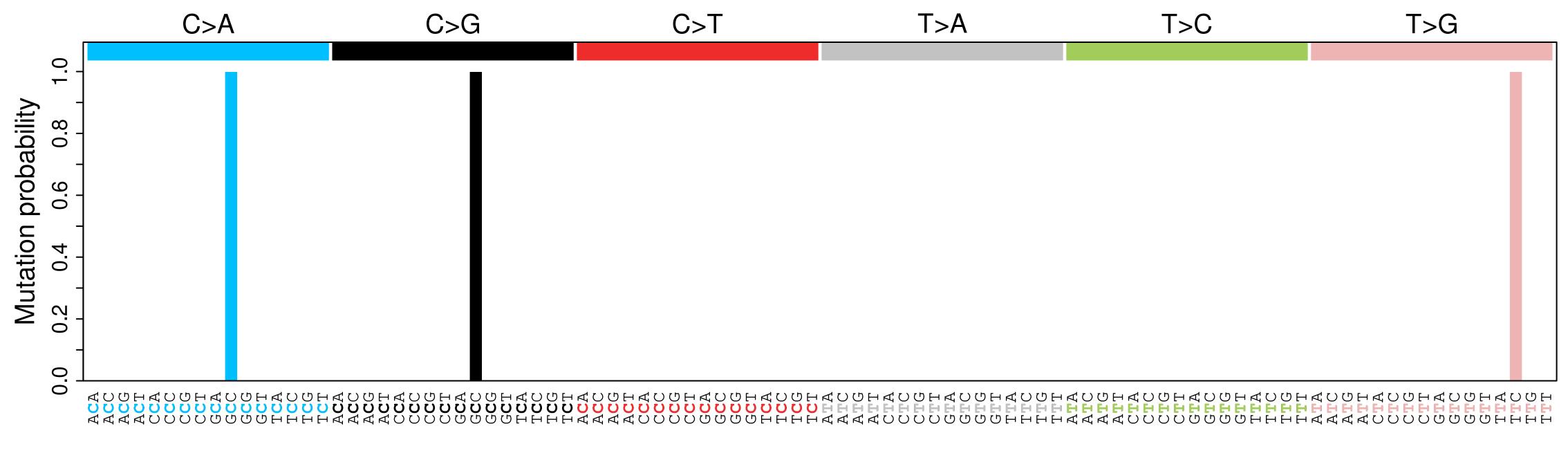
CATD0565a (8 mutations)

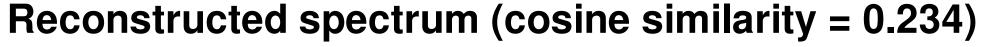


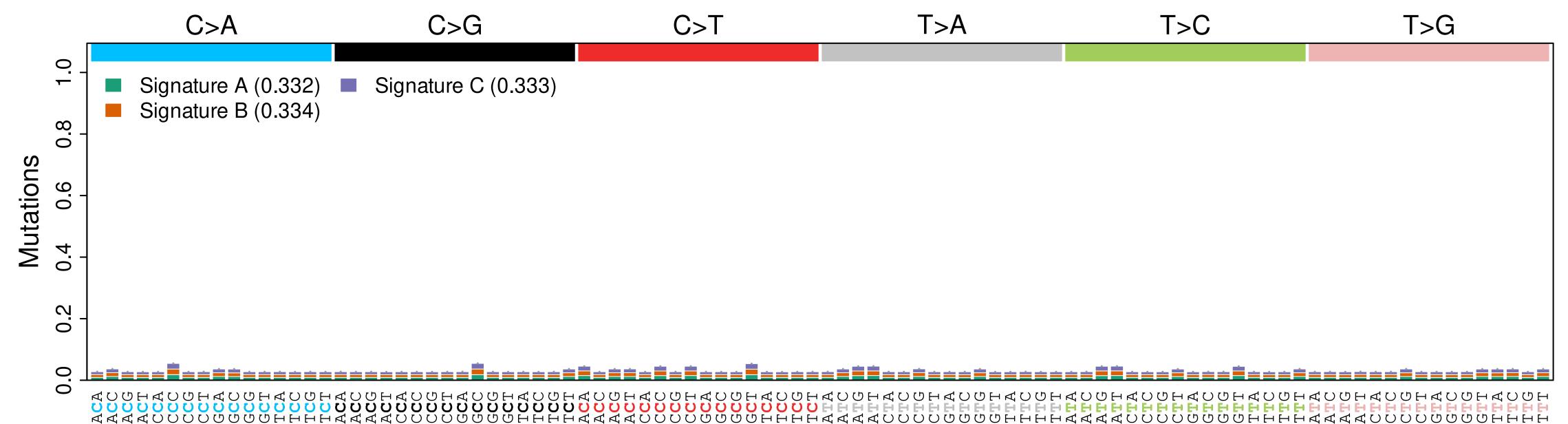
Reconstructed spectrum (cosine similarity = 0.343)



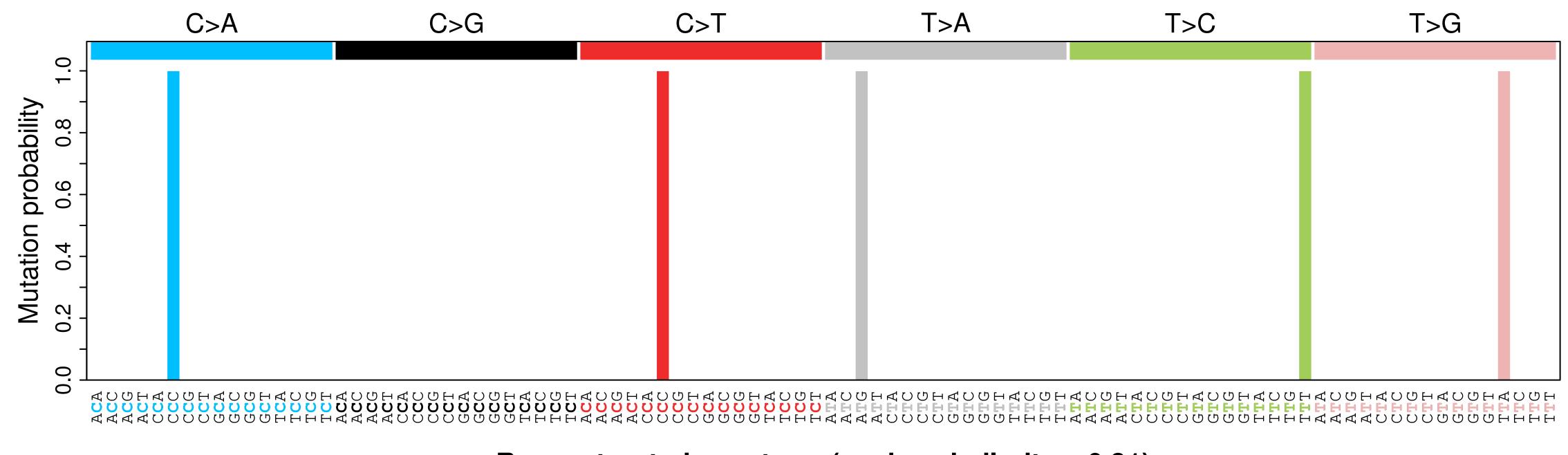


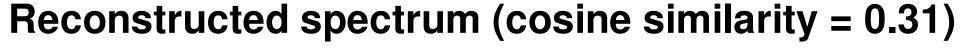


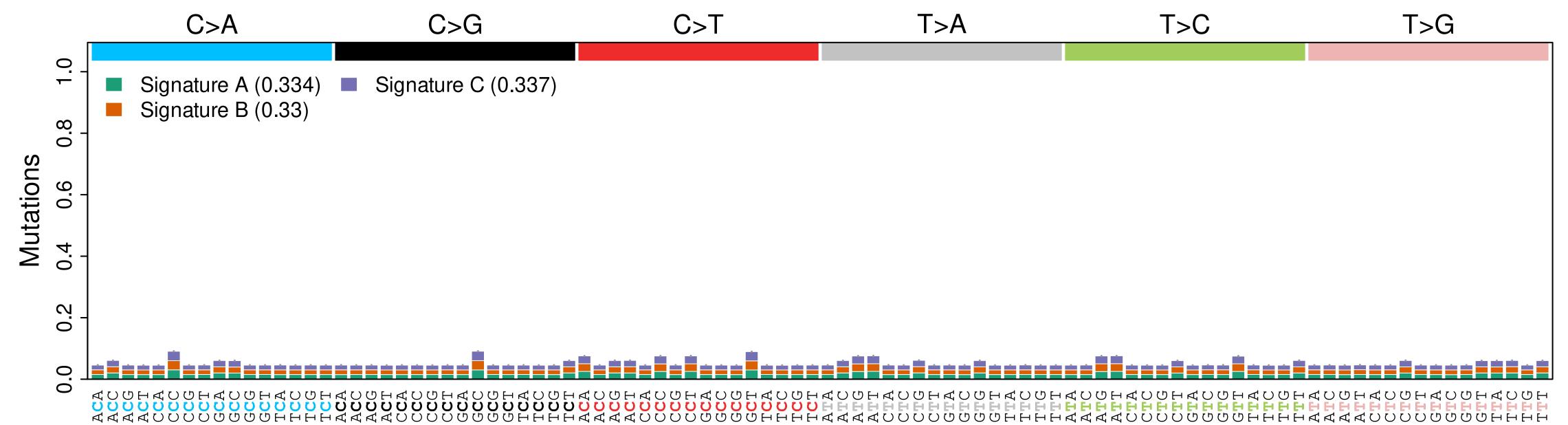




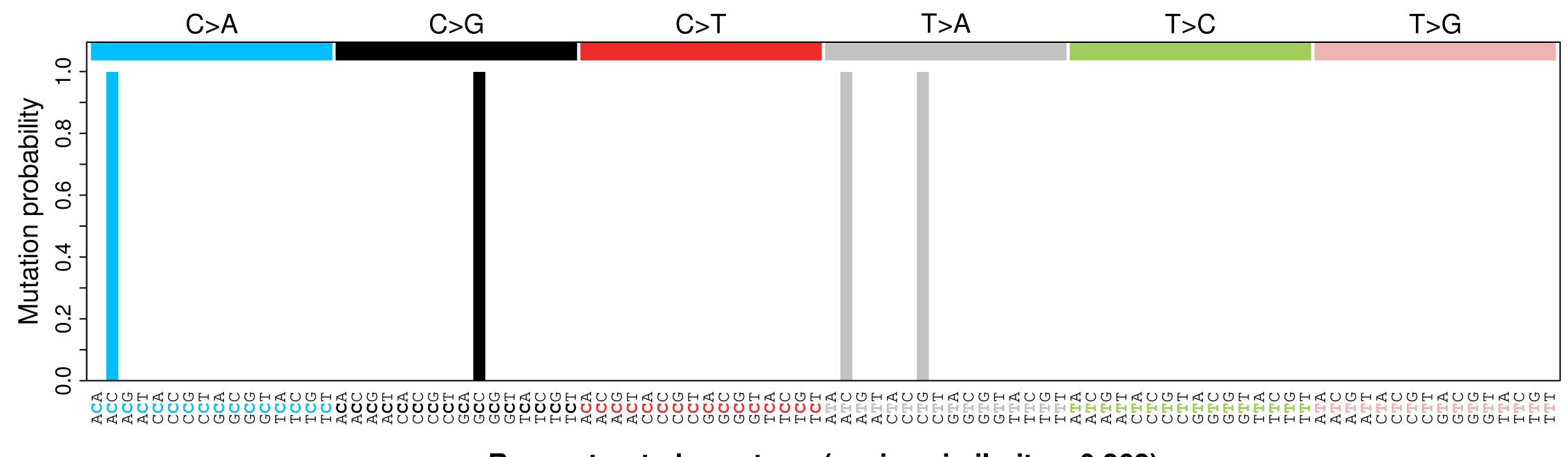












Reconstructed spectrum (cosine similarity = 0.262)

