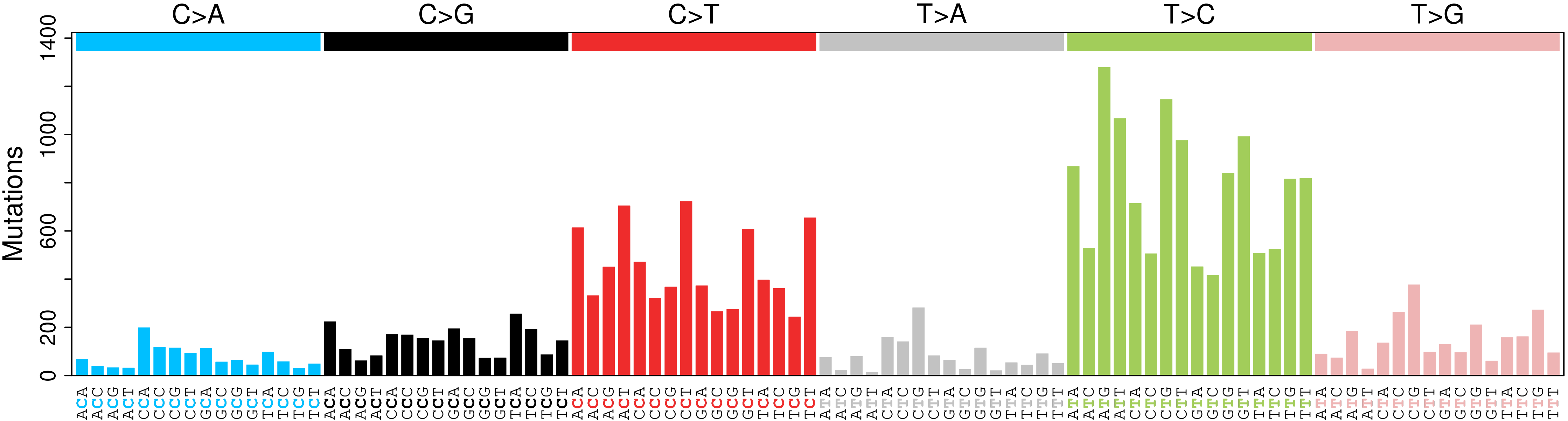
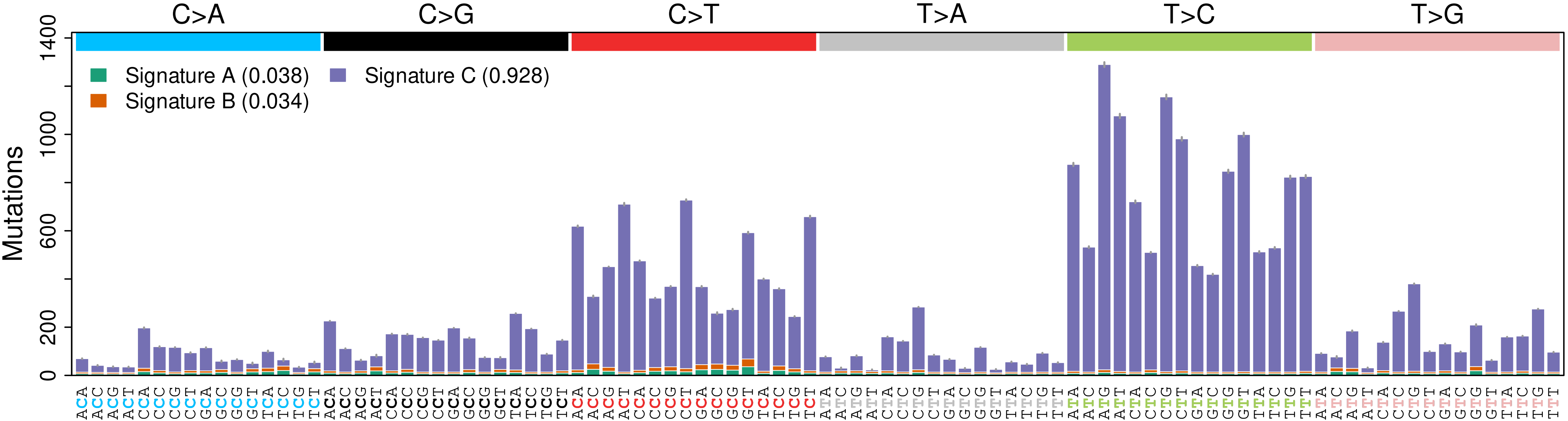


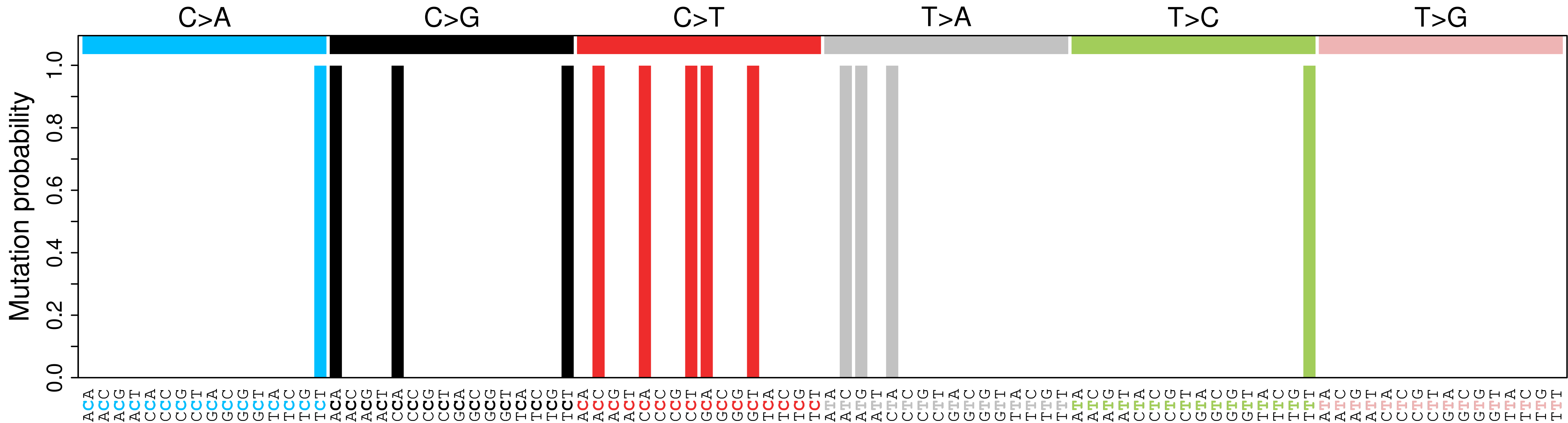
CATD309a (27,083 mutations)



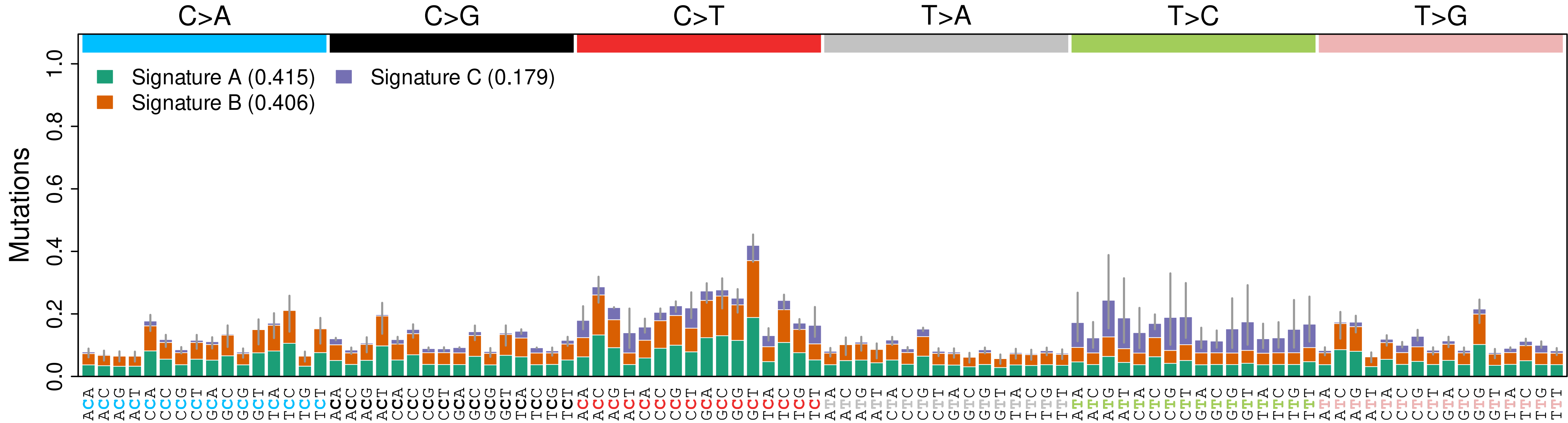
Reconstructed spectrum (cosine similarity = 1)



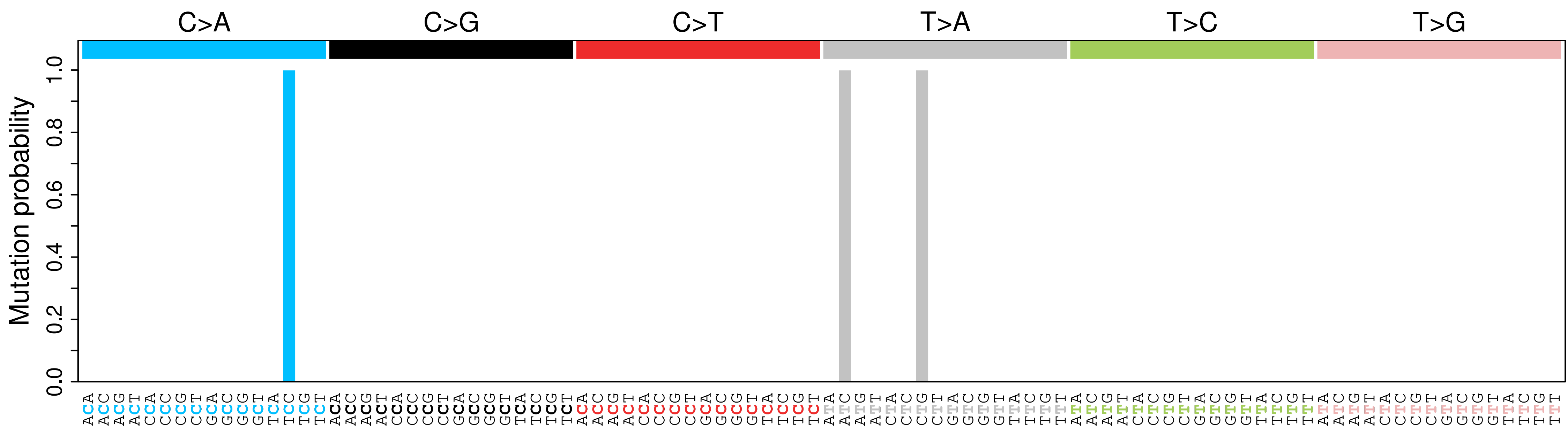
CATD319a



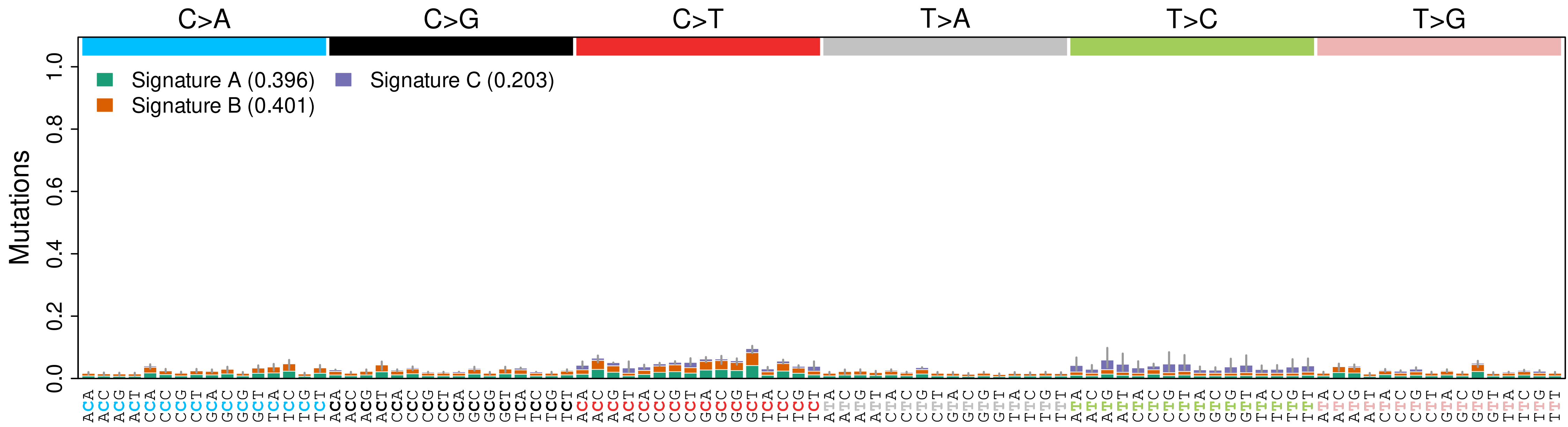
Reconstructed spectrum (cosine similarity = 0.449)



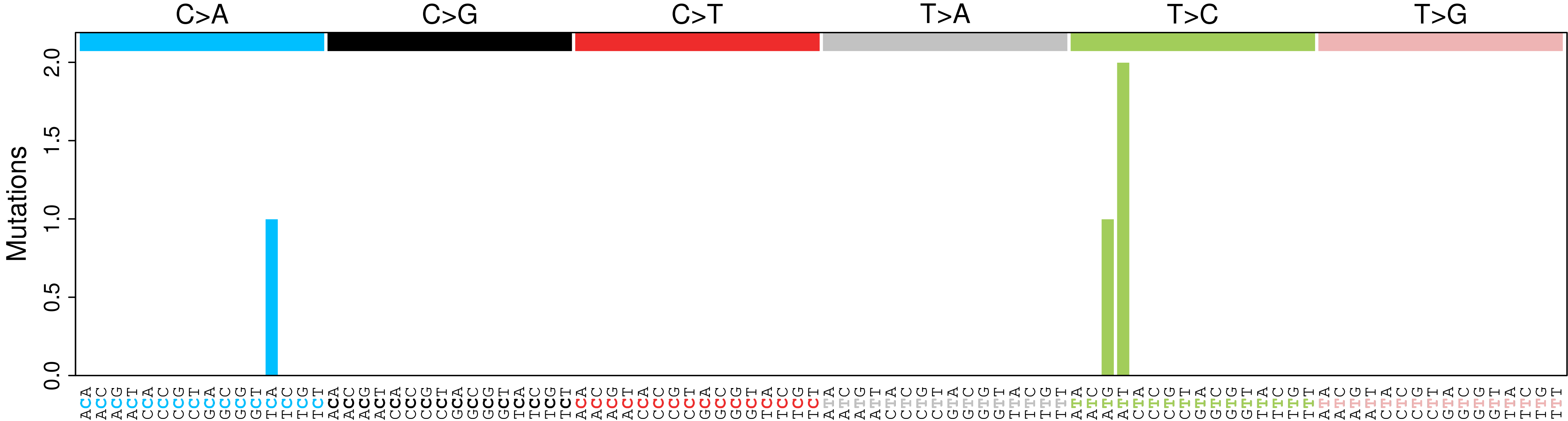
CATD310a



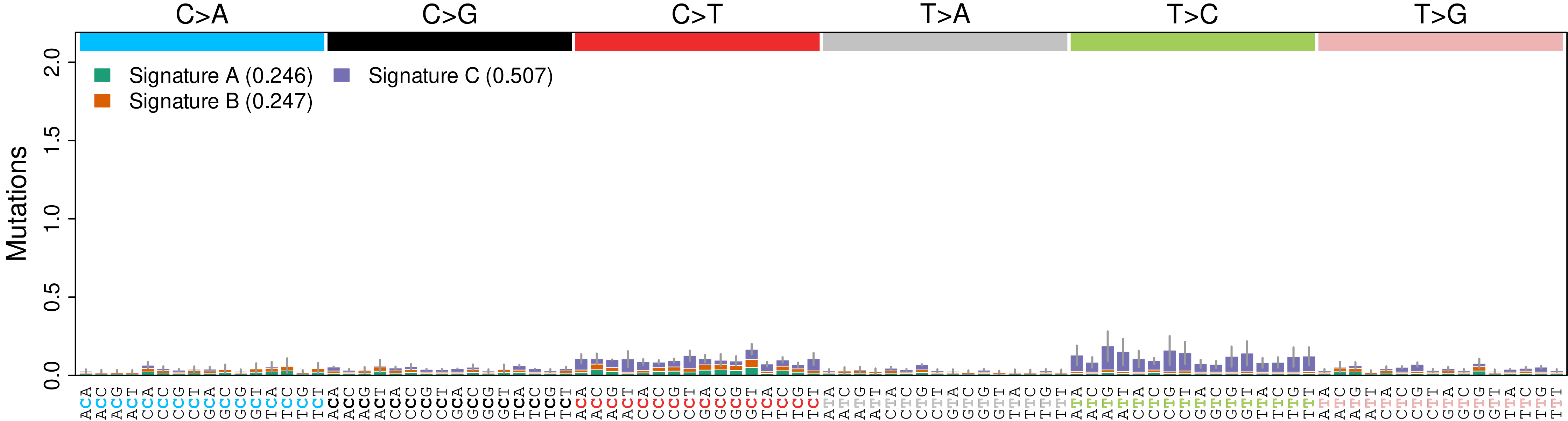
Reconstructed spectrum (cosine similarity = 0.181)



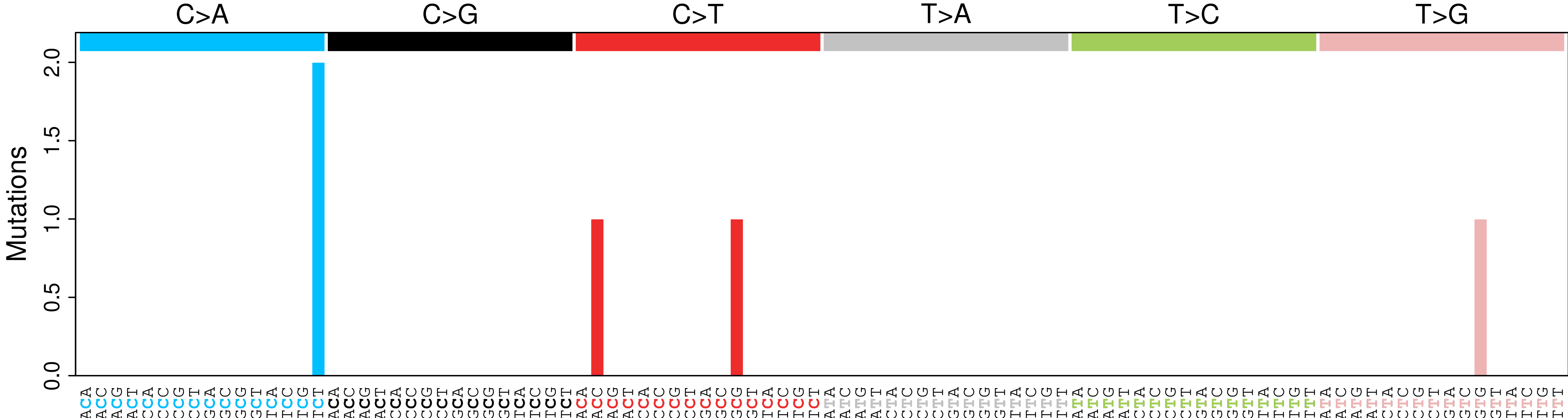
CATD297a (4 mutations)



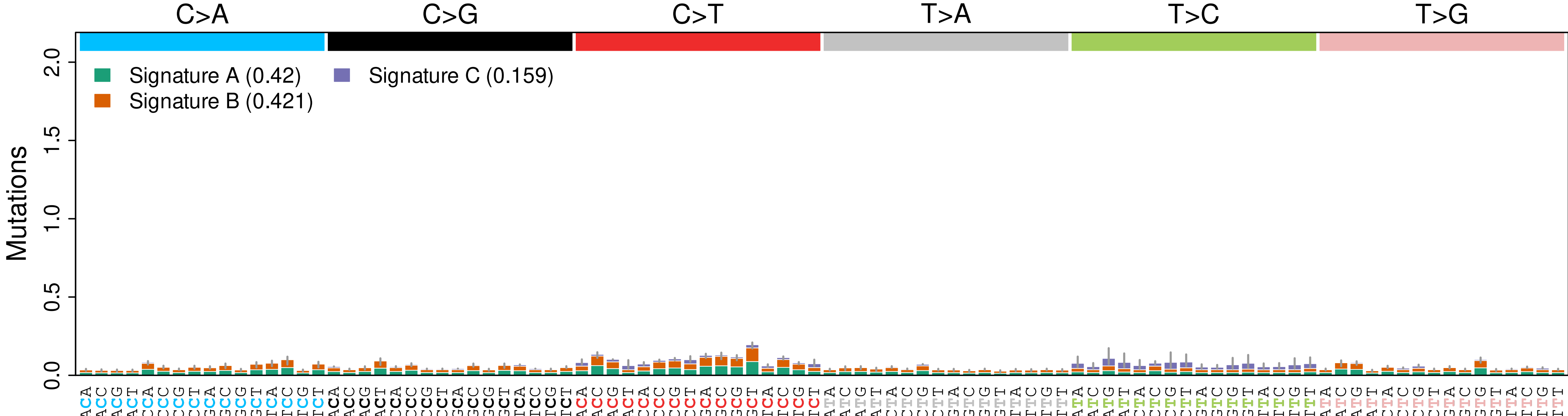
Reconstructed spectrum (cosine similarity = 0.31)



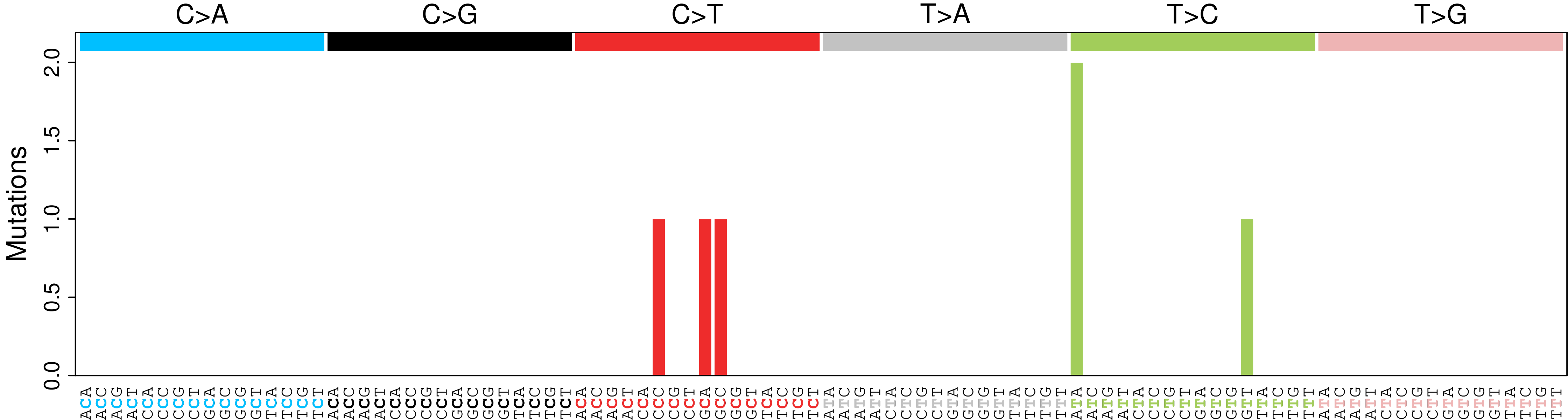
CATD323a (5 mutations)



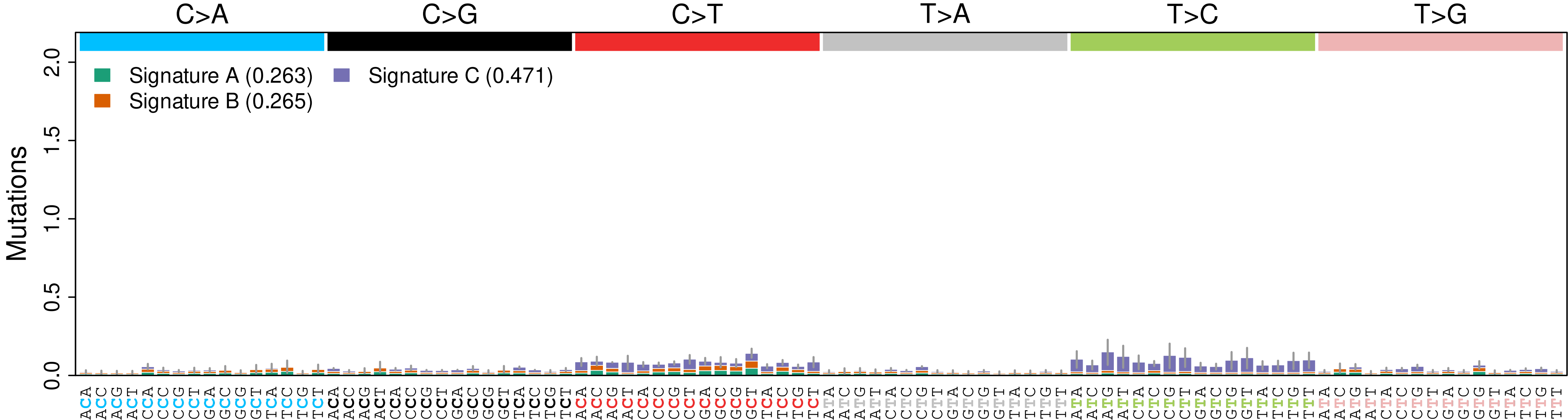
Reconstructed spectrum (cosine similarity = 0.28)



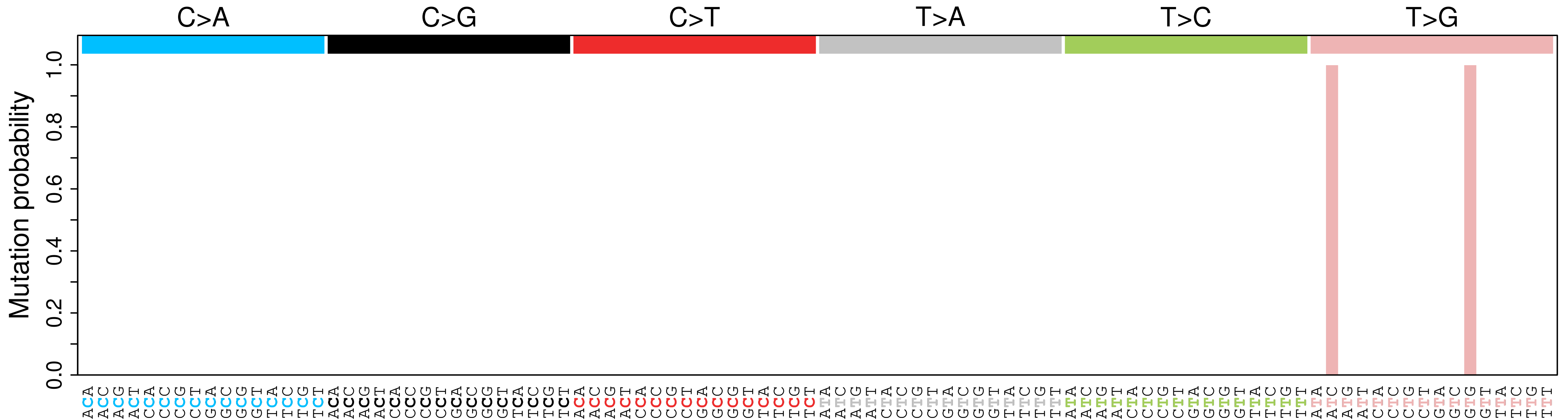
CATD314a (6 mutations)



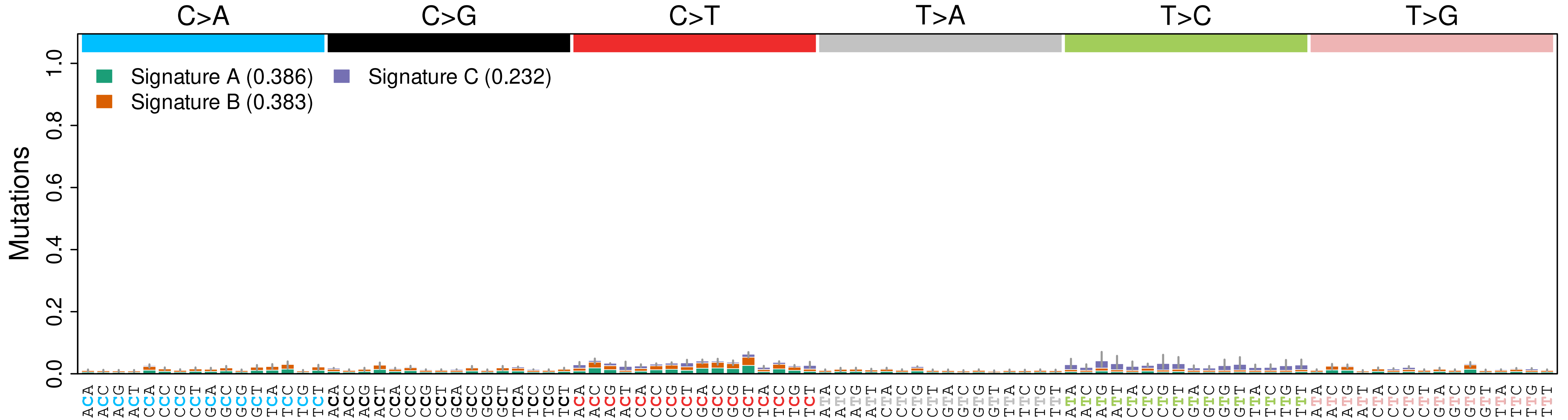
Reconstructed spectrum (cosine similarity = 0.337)



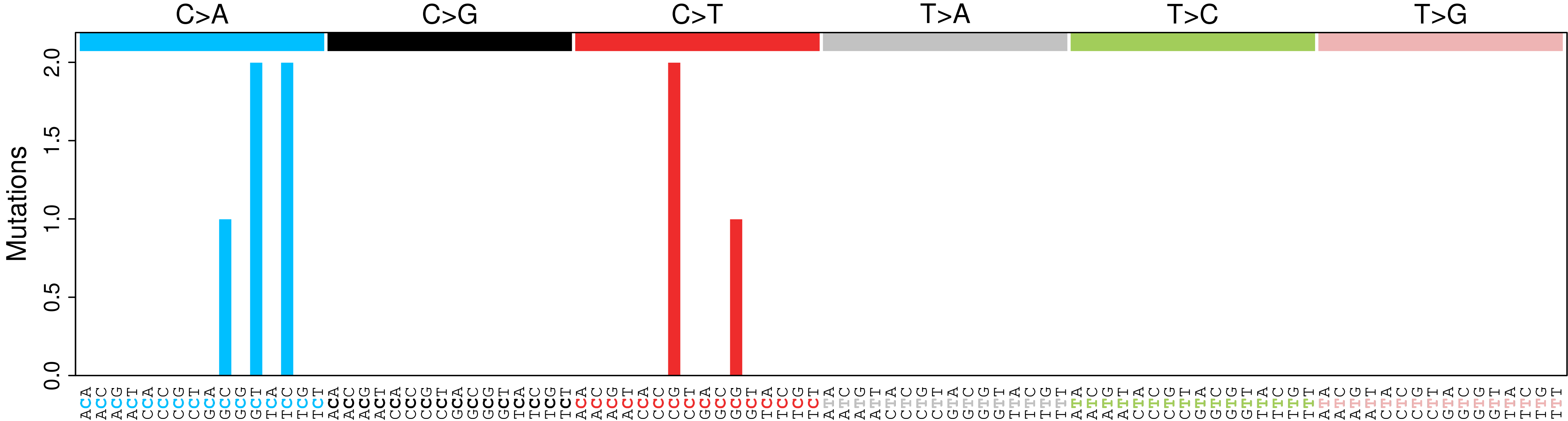
CATD311a



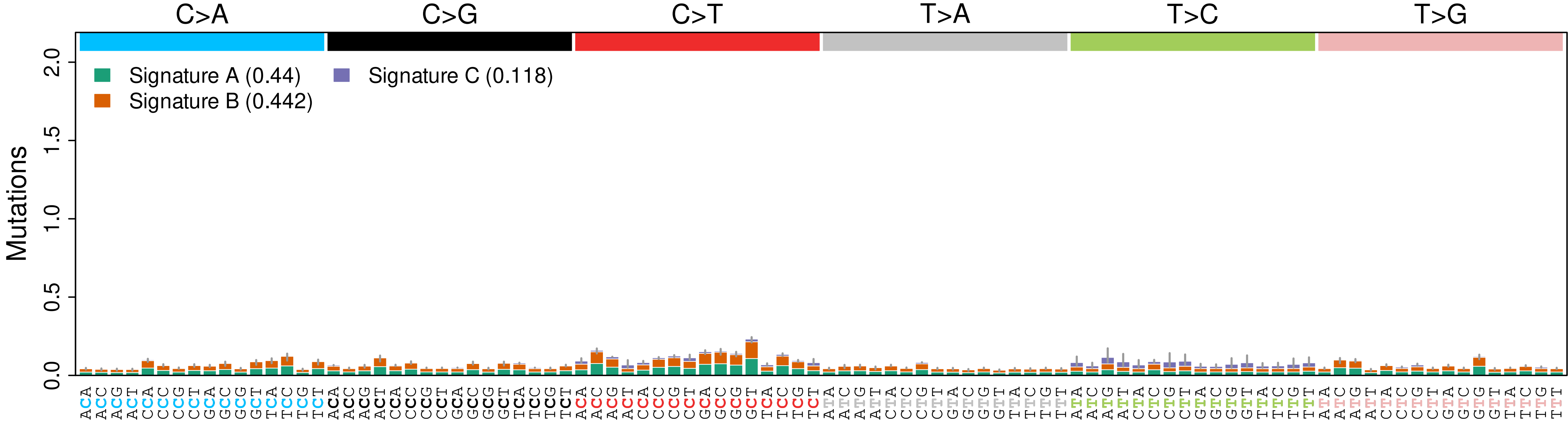
Reconstructed spectrum (cosine similarity = 0.179)



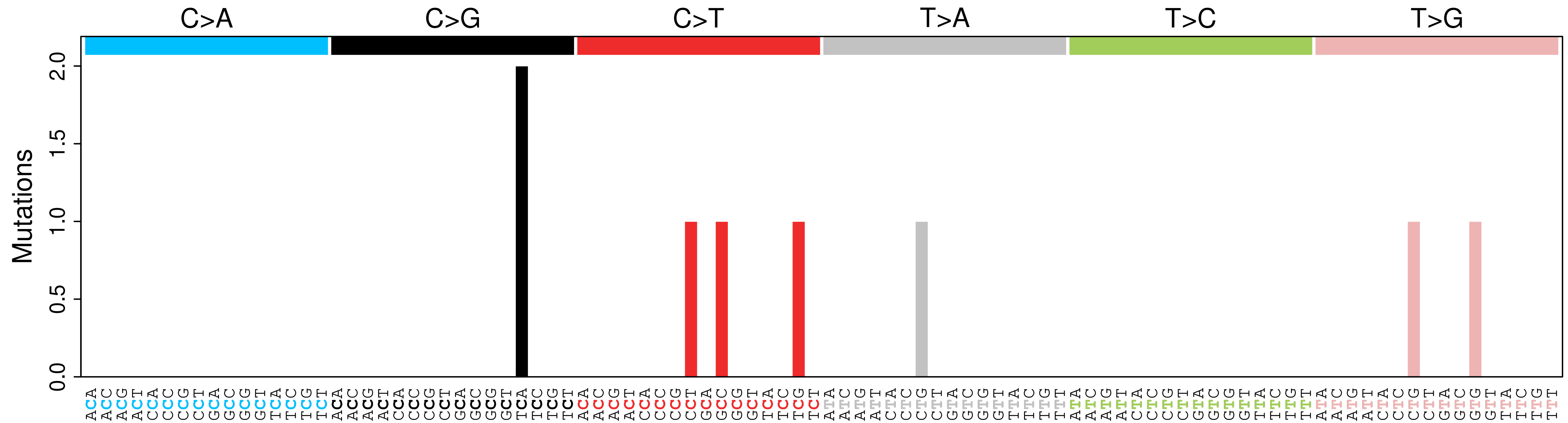
CATD300a (8 mutations)



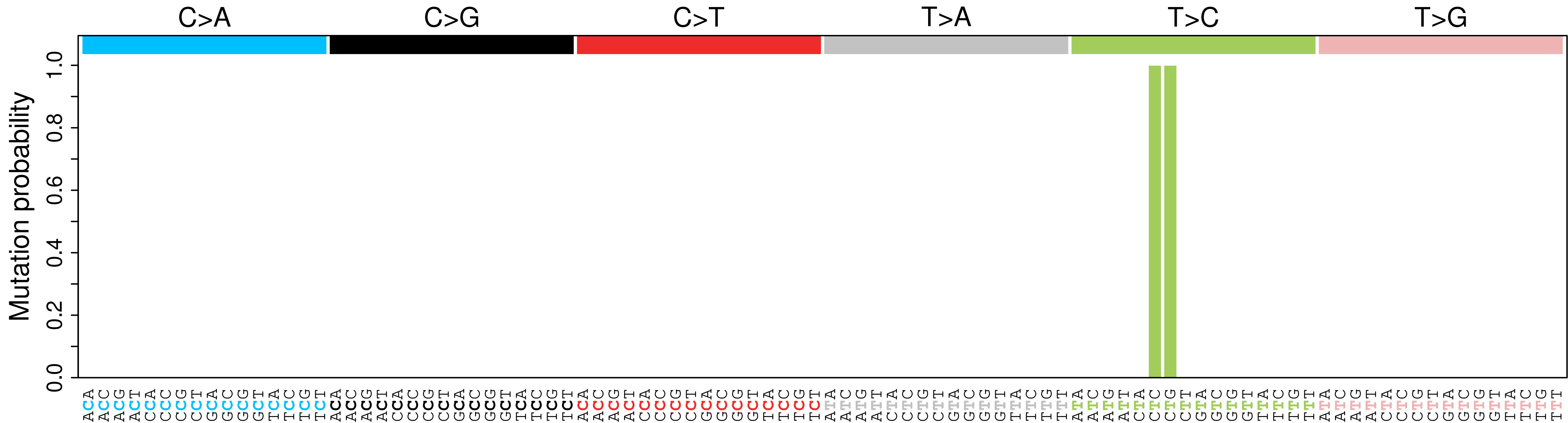
Reconstructed spectrum (cosine similarity = 0.301)



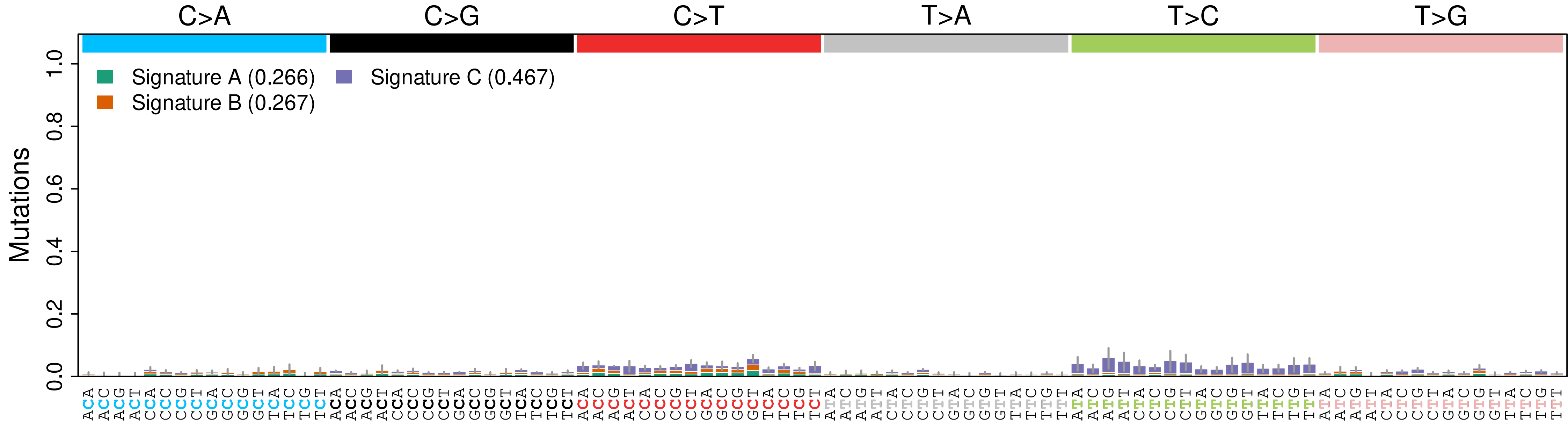
CATD328a (8 mutations)



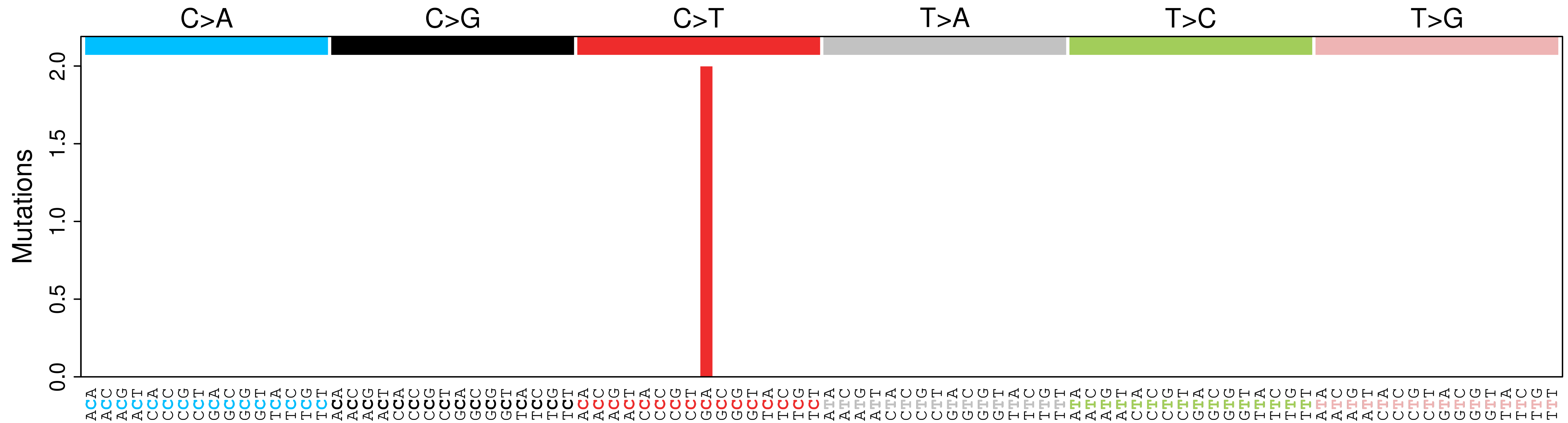
CATD0674a



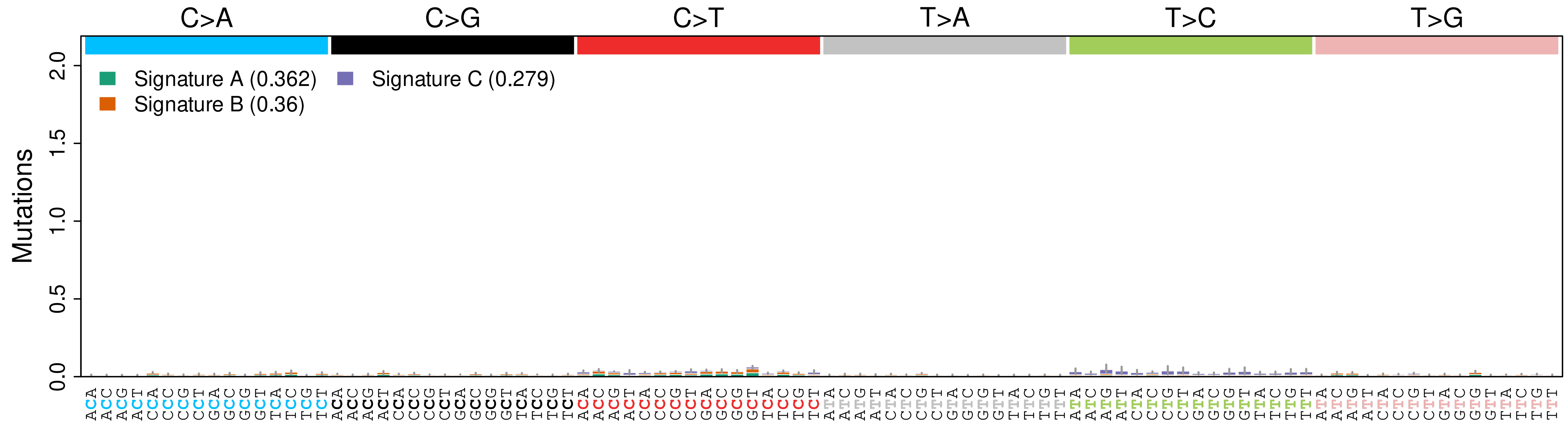
Reconstructed spectrum (cosine similarity = 0.242)



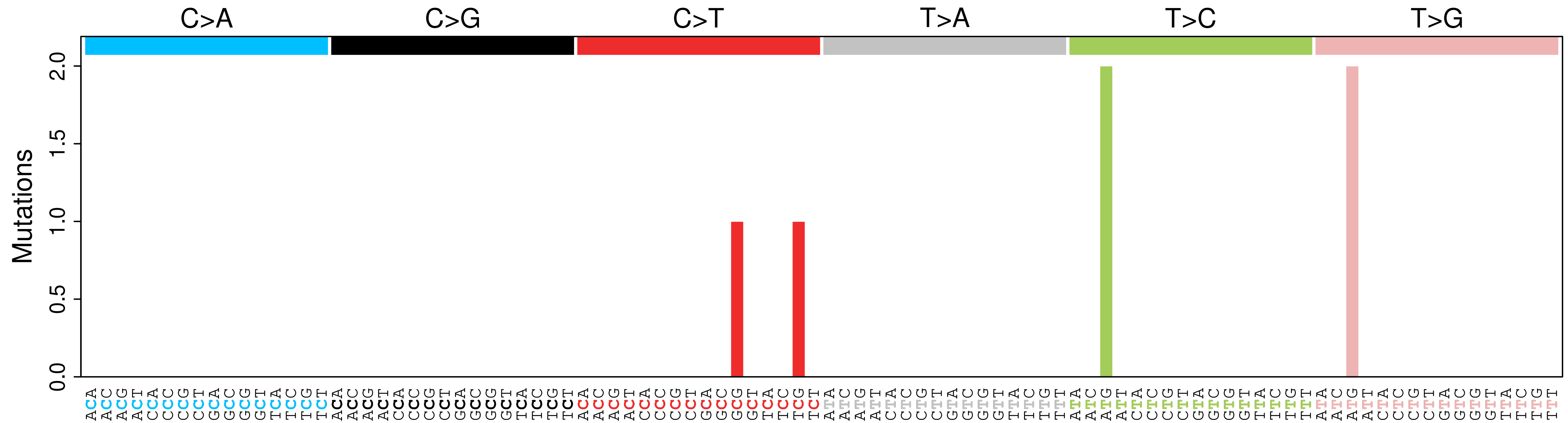
CATD318a (2 mutations)



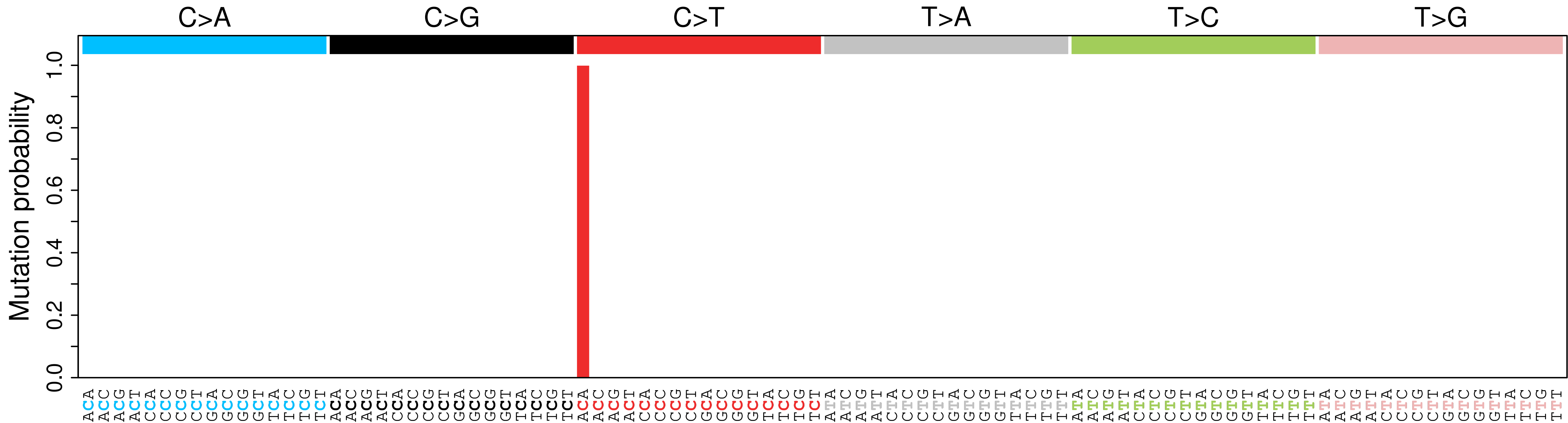
Reconstructed spectrum (cosine similarity = 0.177)



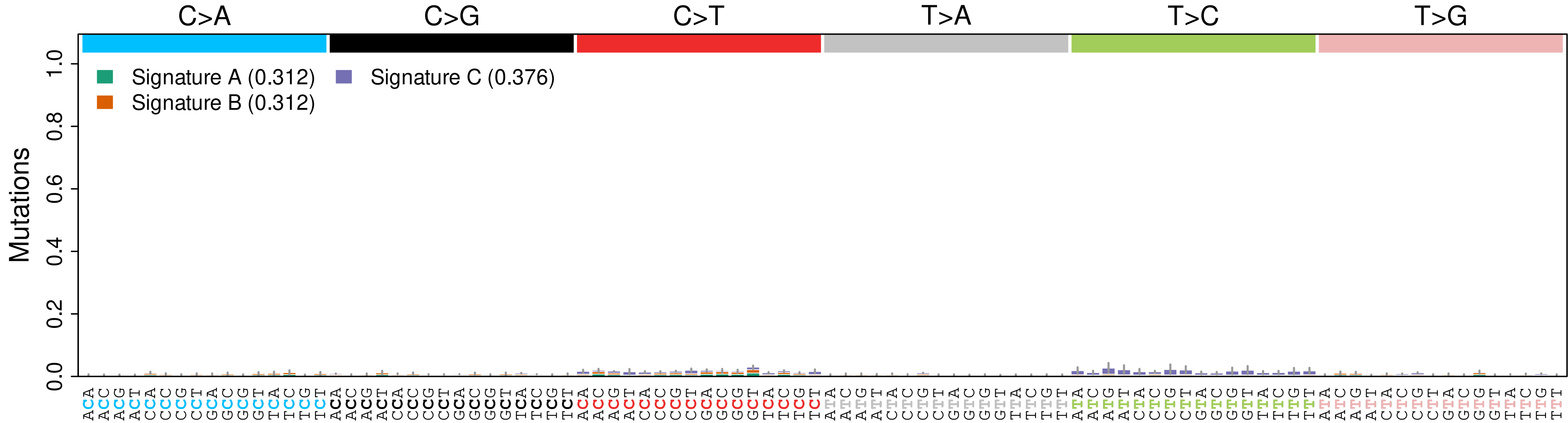
CATD301a (6 mutations)

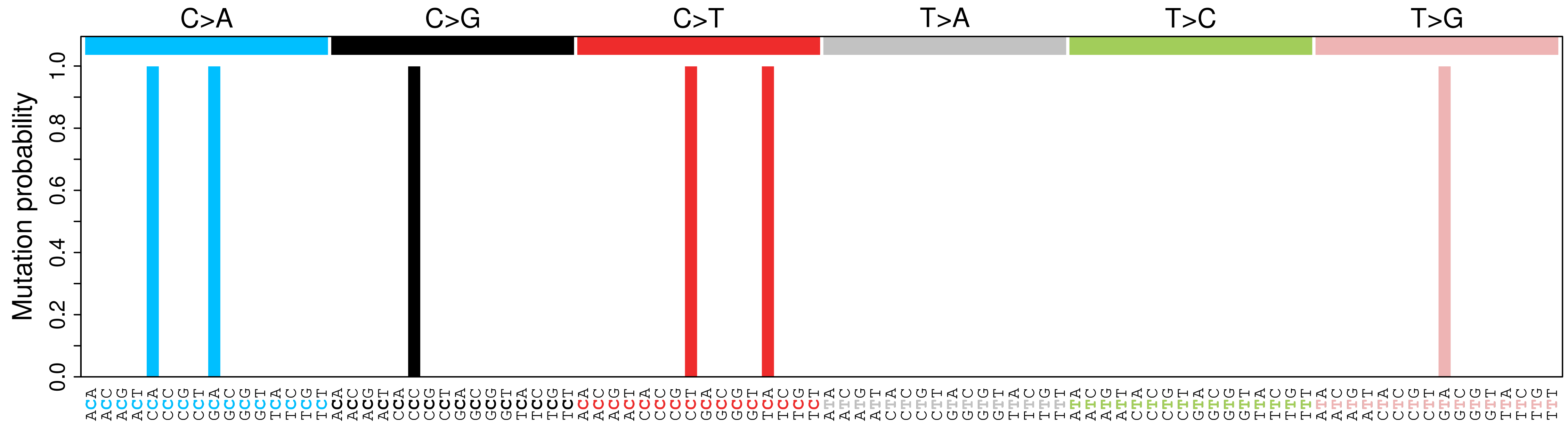


CATD308a

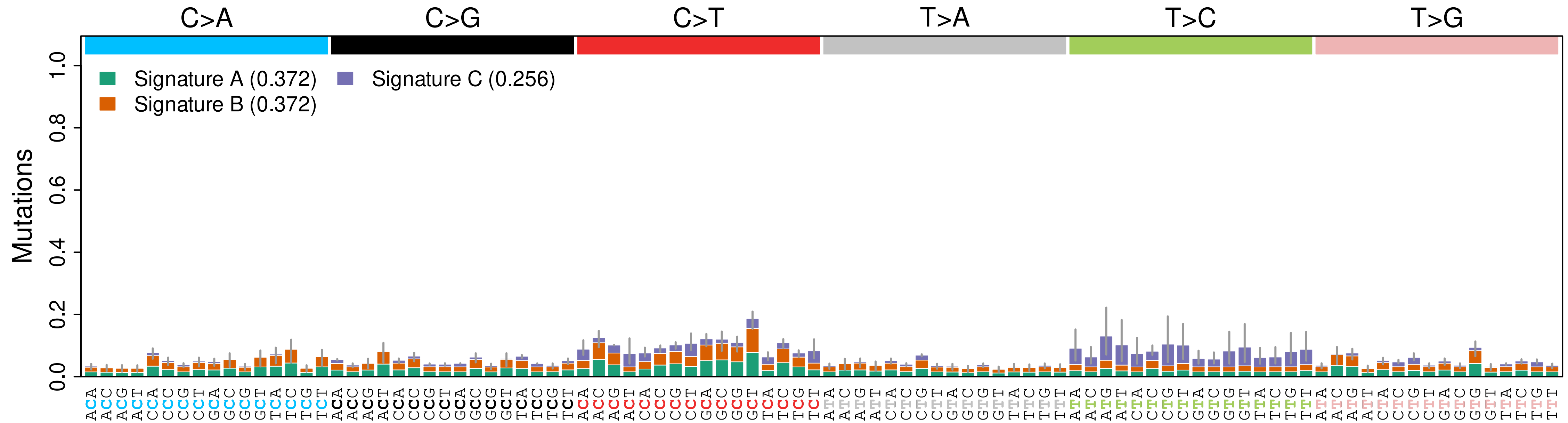


Reconstructed spectrum (cosine similarity = 0.14)

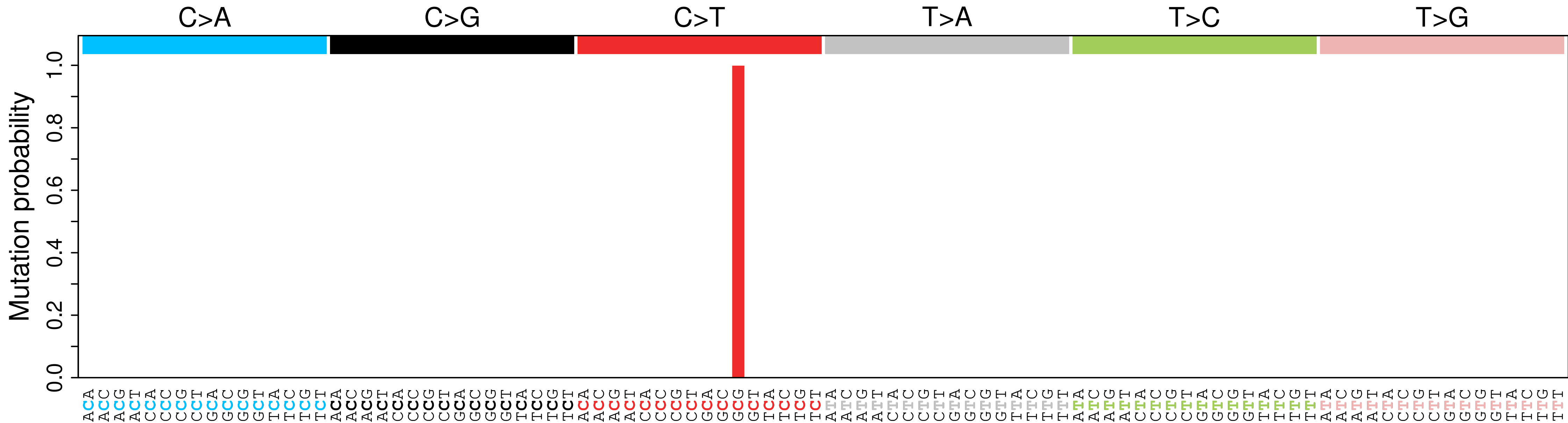


CATD296a

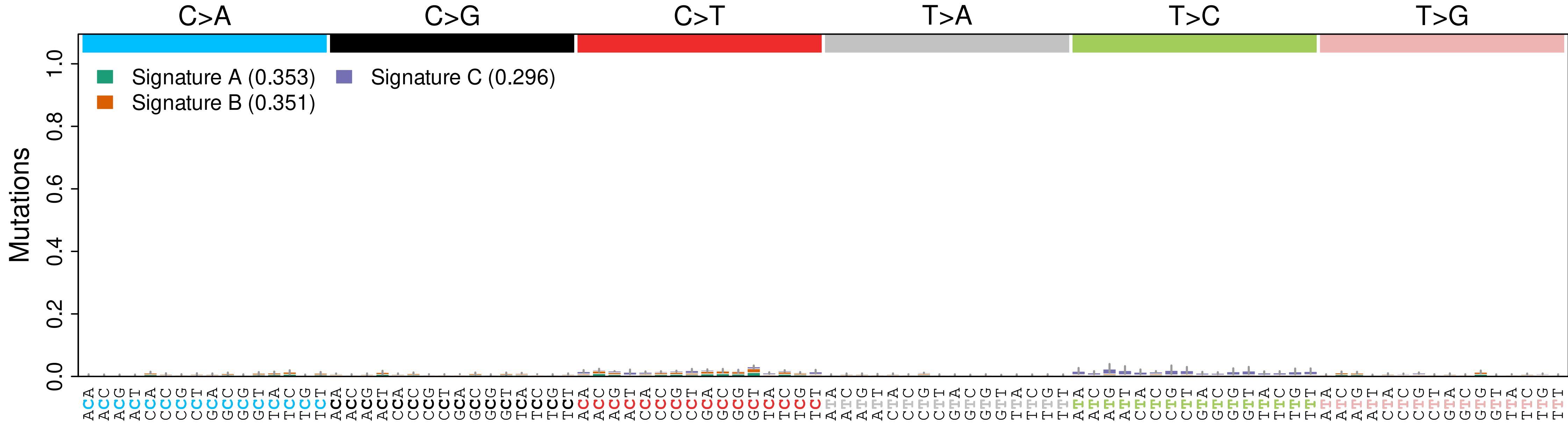
Reconstructed spectrum (cosine similarity = 0.249)



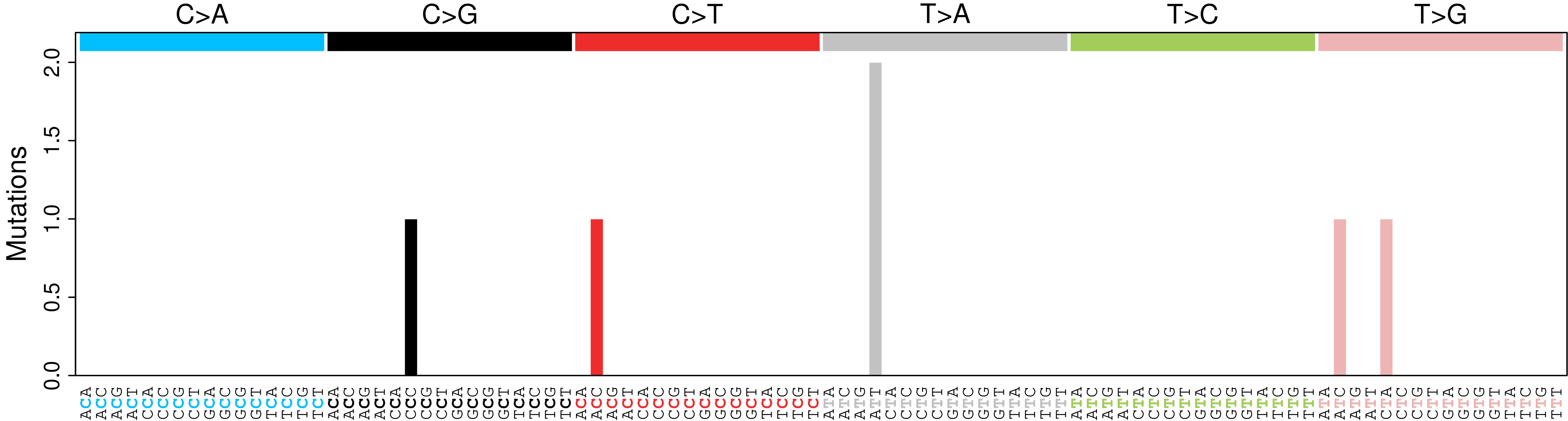
CATD0675a



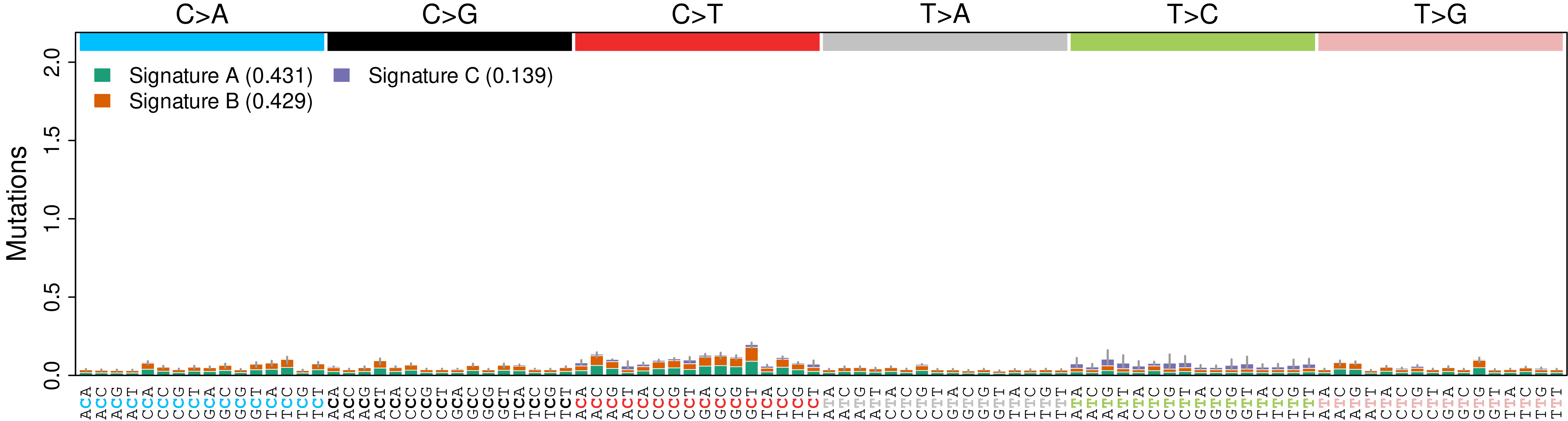
Reconstructed spectrum (cosine similarity = 0.157)



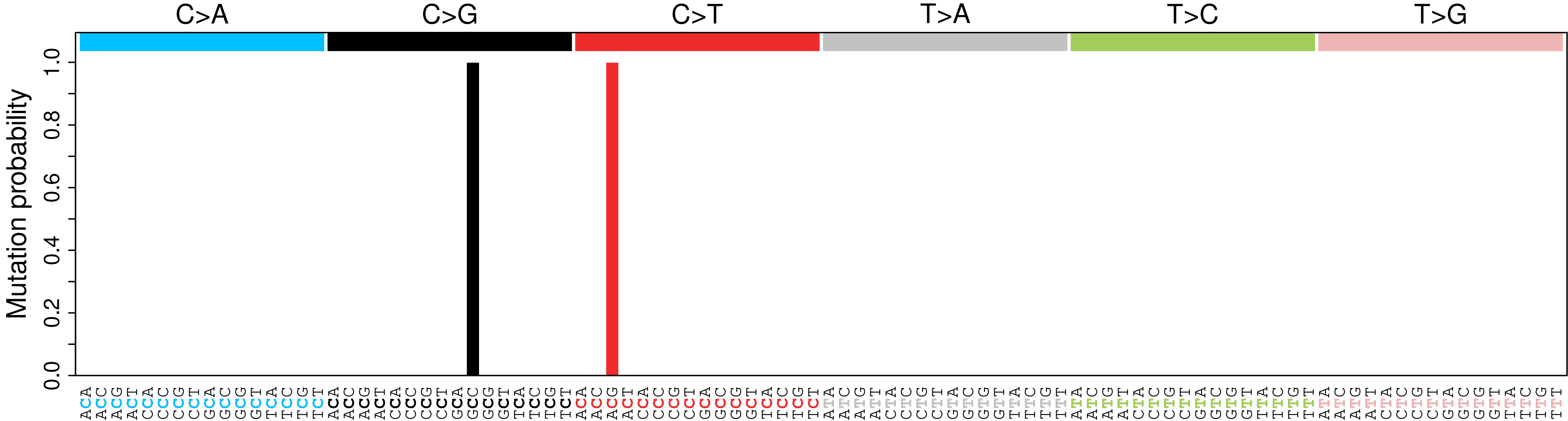
CATD326a (6 mutations)



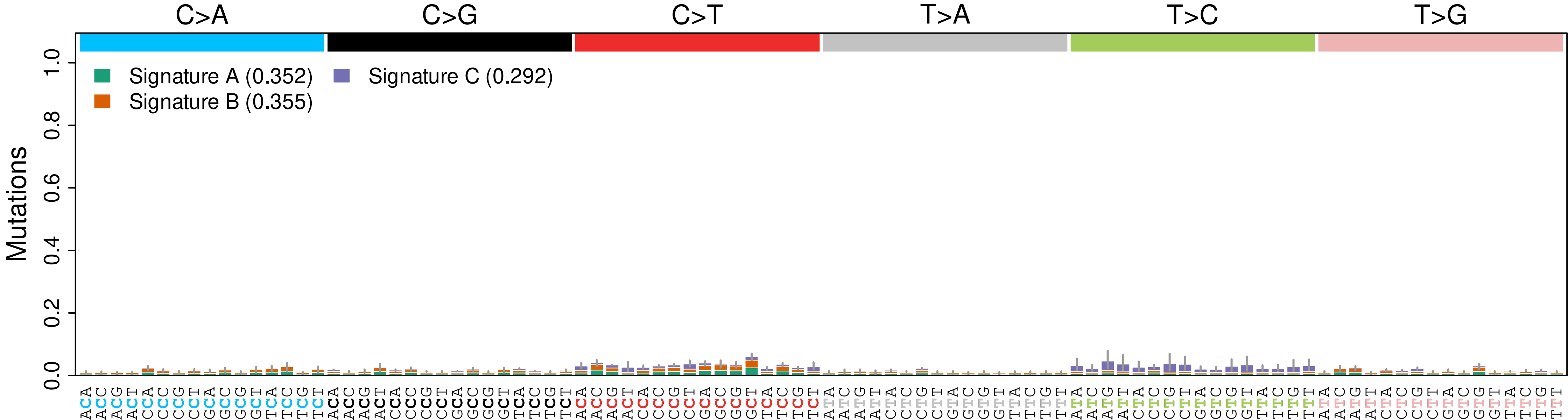
Reconstructed spectrum (cosine similarity = 0.226)



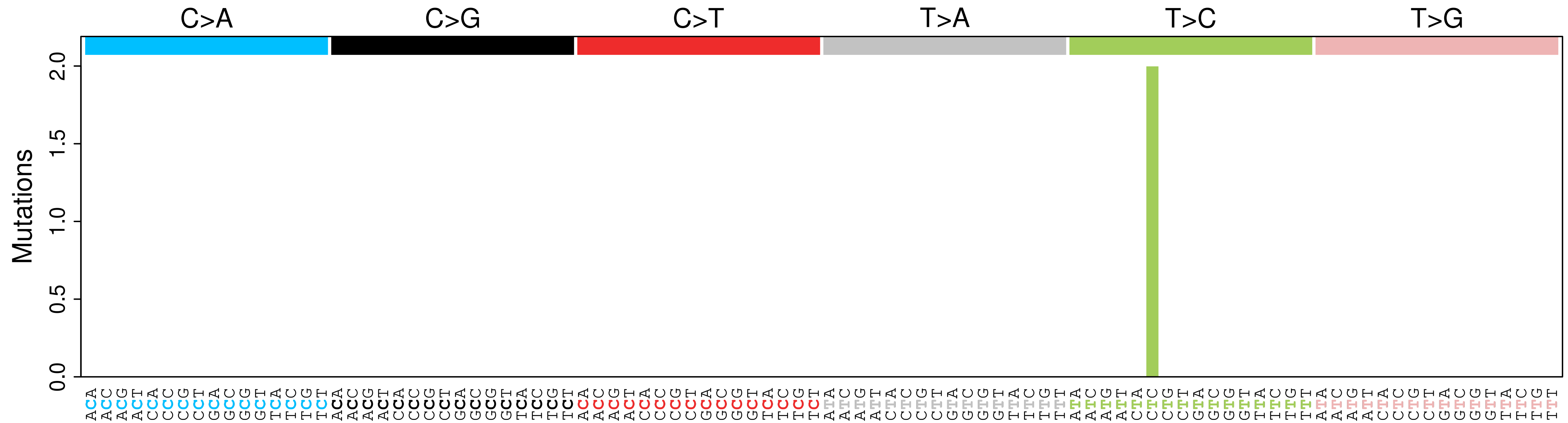
CATD0677a



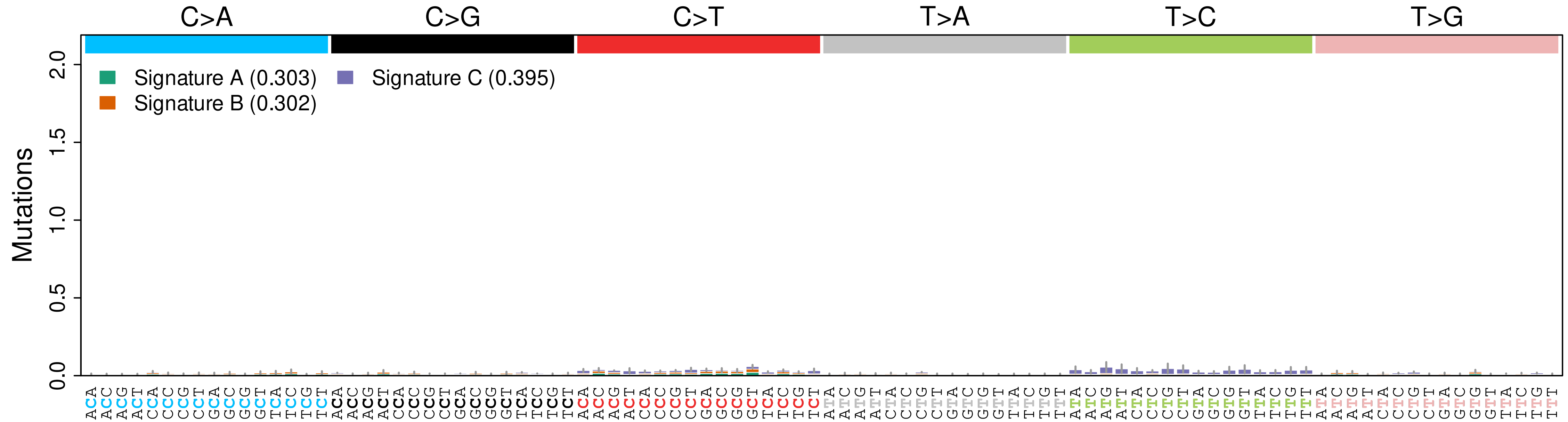
Reconstructed spectrum (cosine similarity = 0.169)



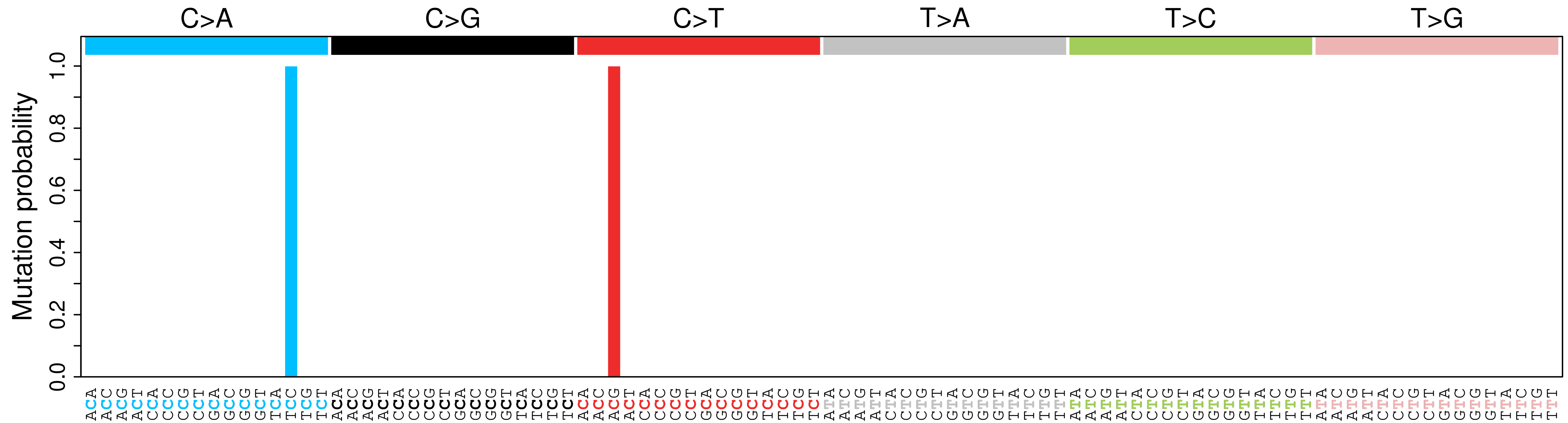
CATD304a (2 mutations)



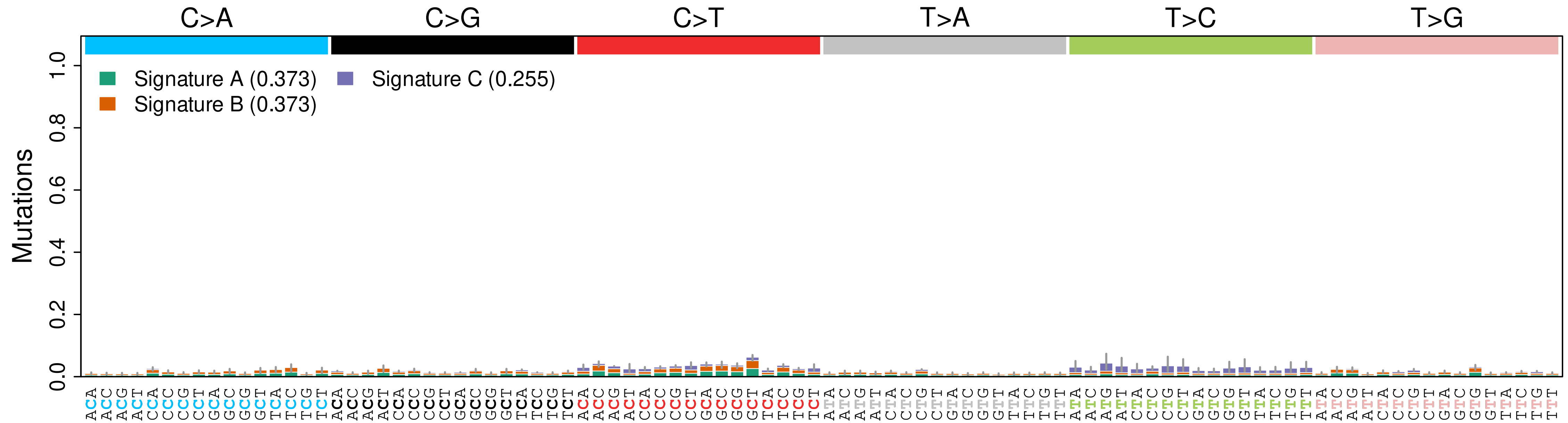
Reconstructed spectrum (cosine similarity = 0.127)



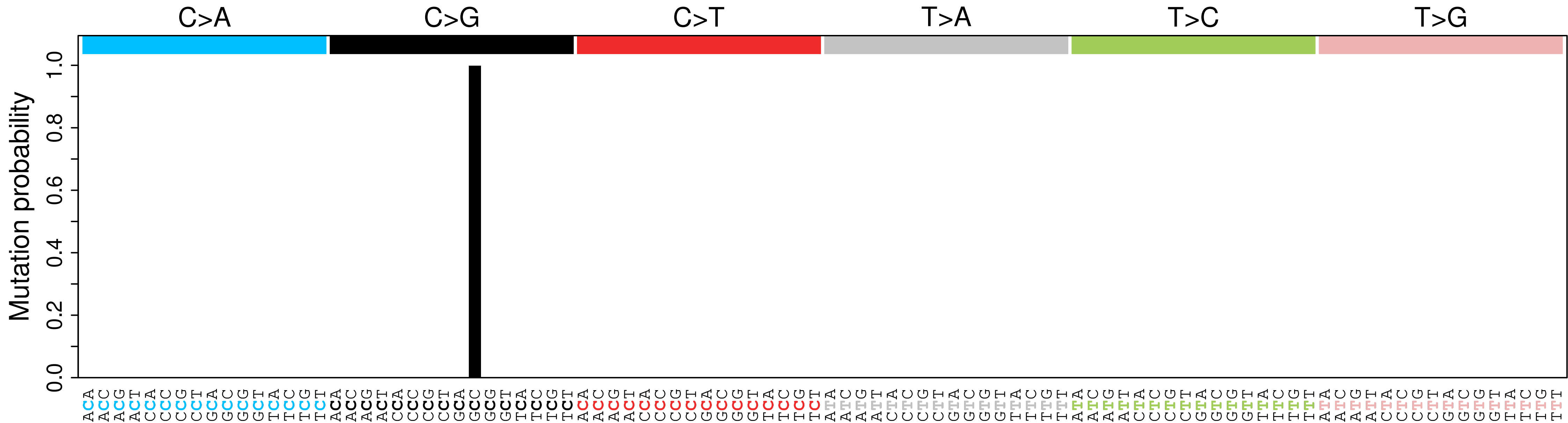
CATD299a



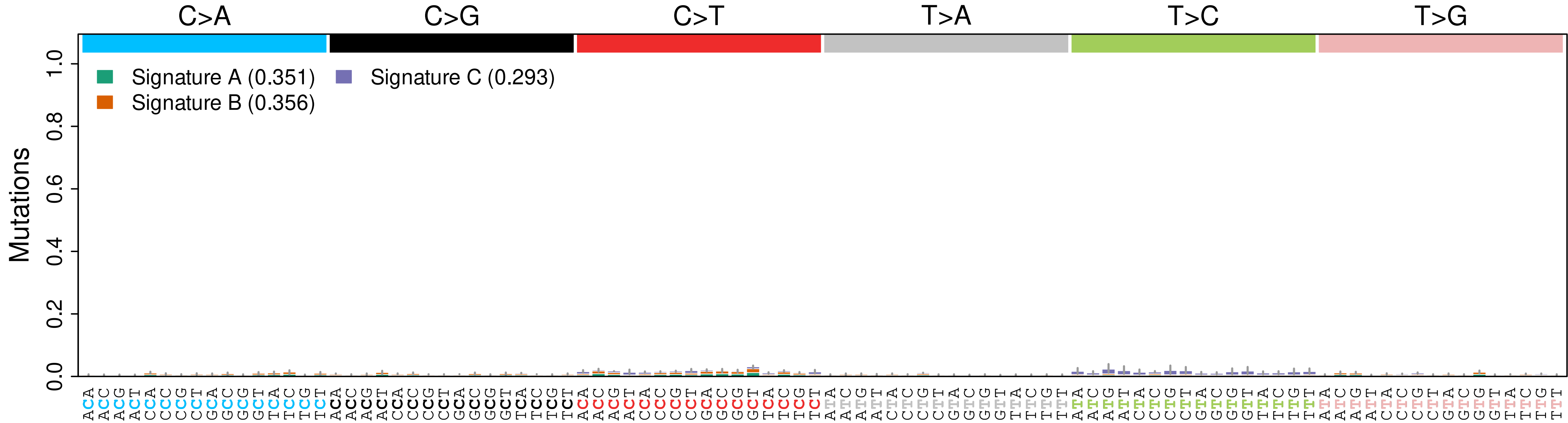
Reconstructed spectrum (cosine similarity = 0.2)



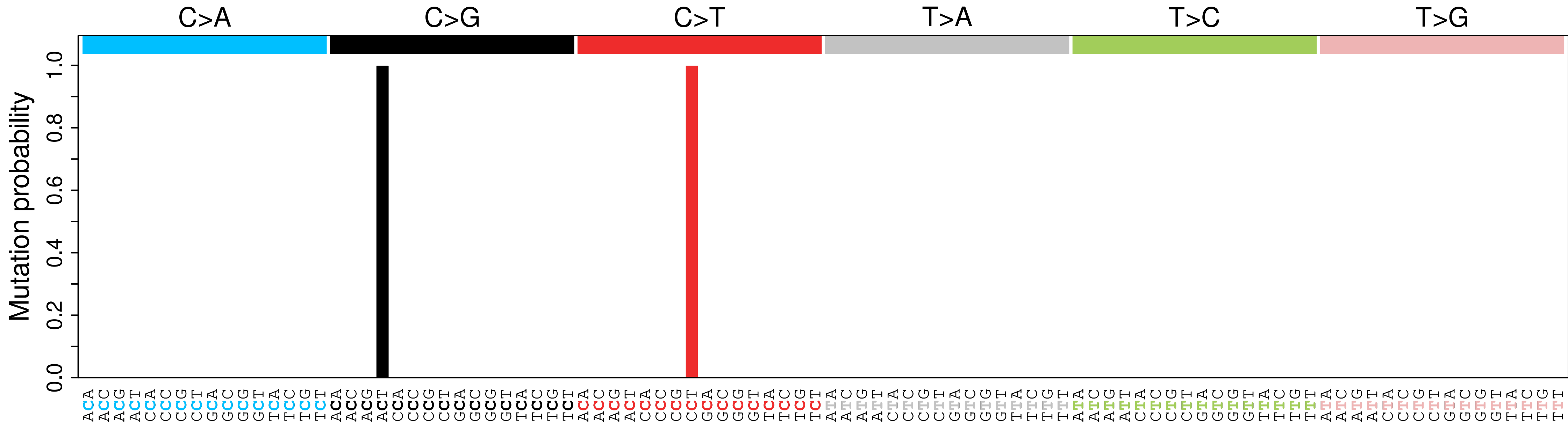
CATD317a



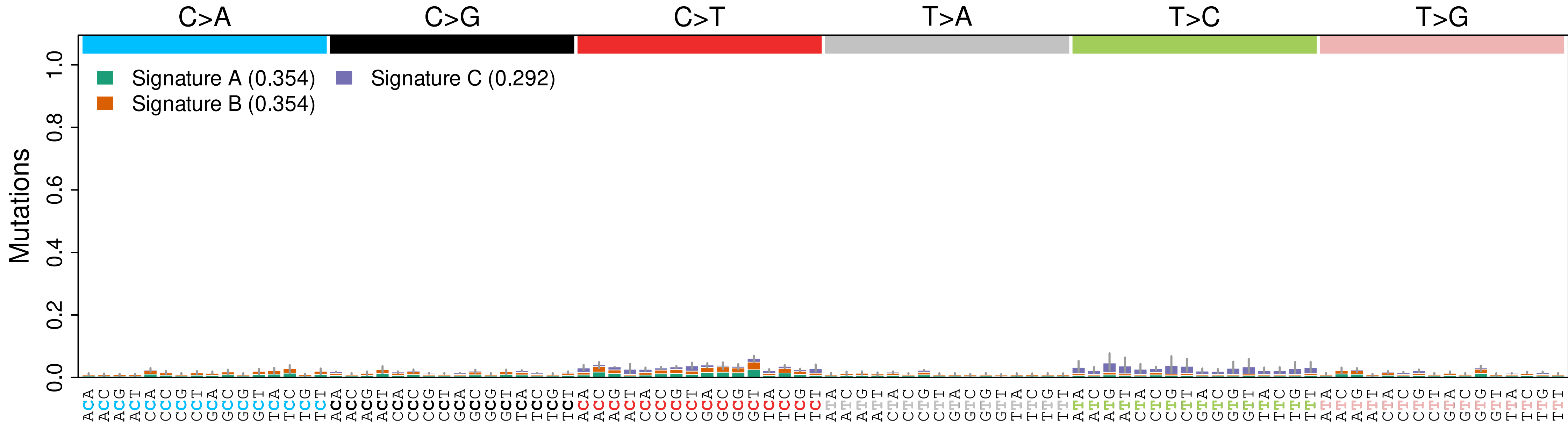
Reconstructed spectrum (cosine similarity = 0.09)



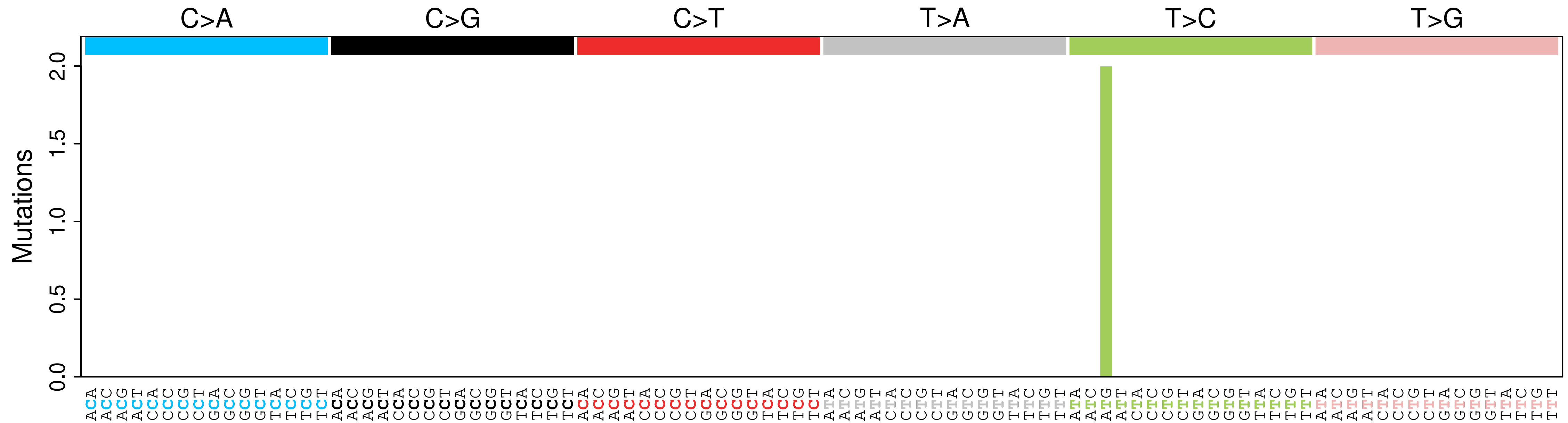
CATD307a



Reconstructed spectrum (cosine similarity = 0.198)



CATD303a (2 mutations)



Reconstructed spectrum (cosine similarity = 0.26)

