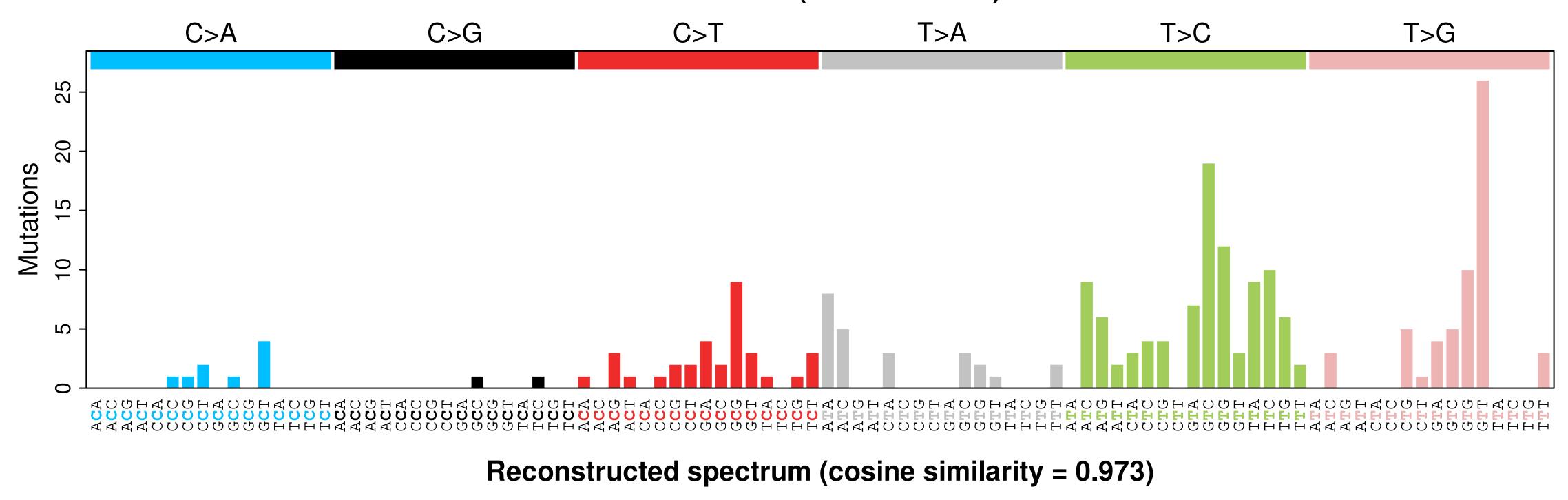
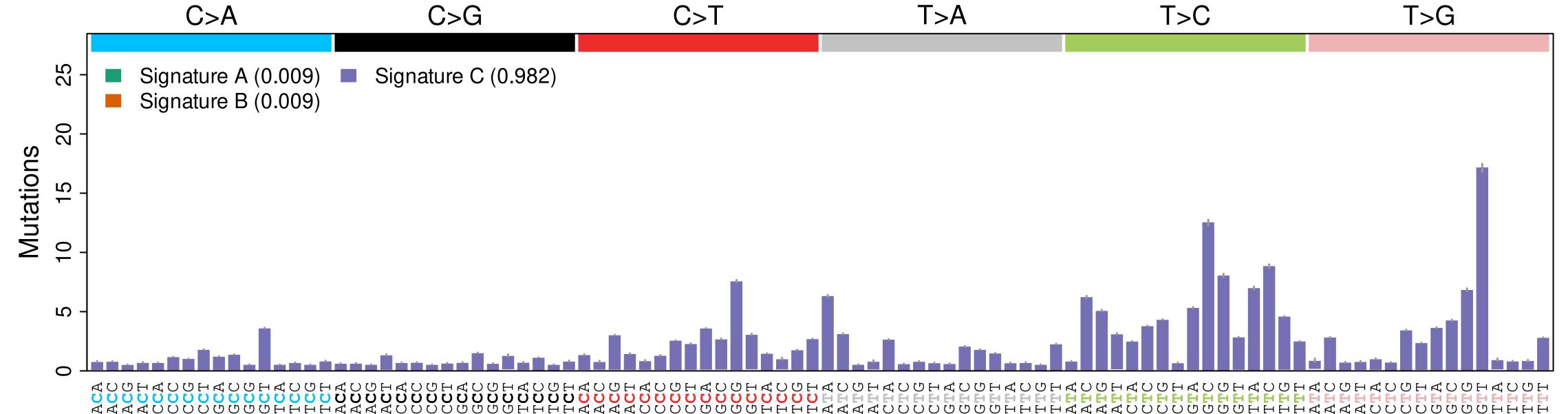
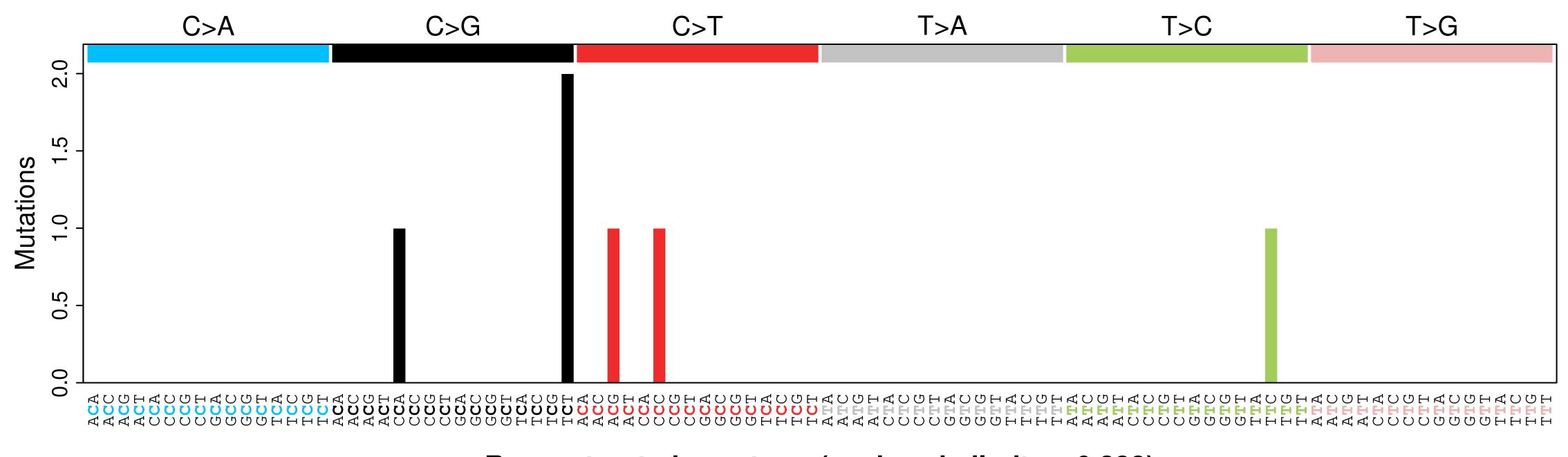
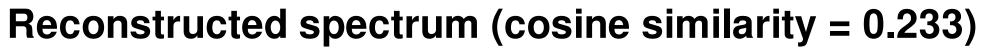
CATD0735a (221 mutations)

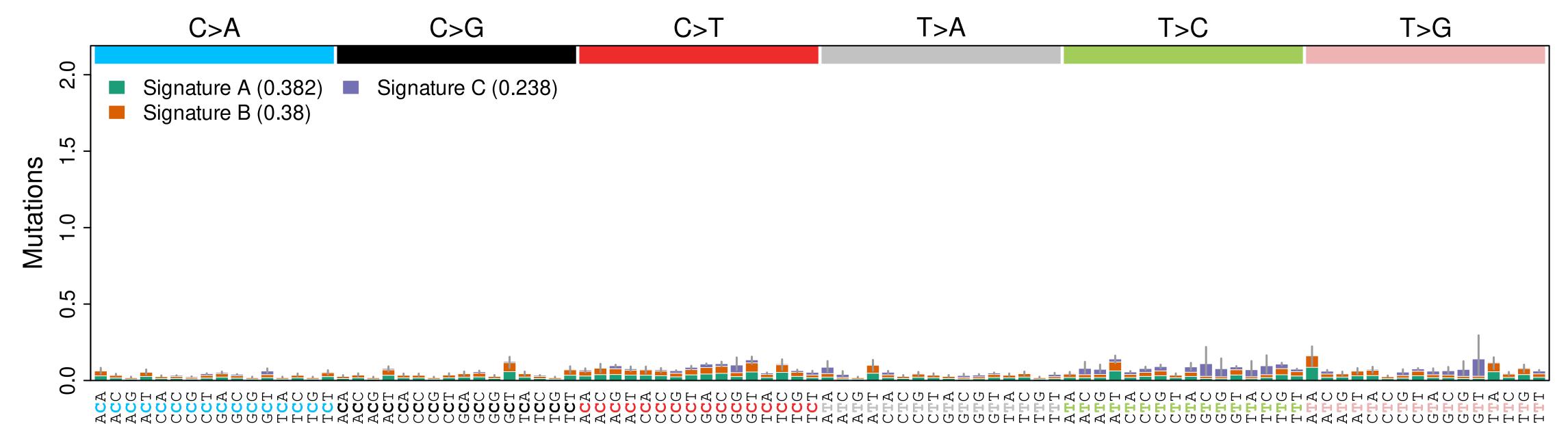




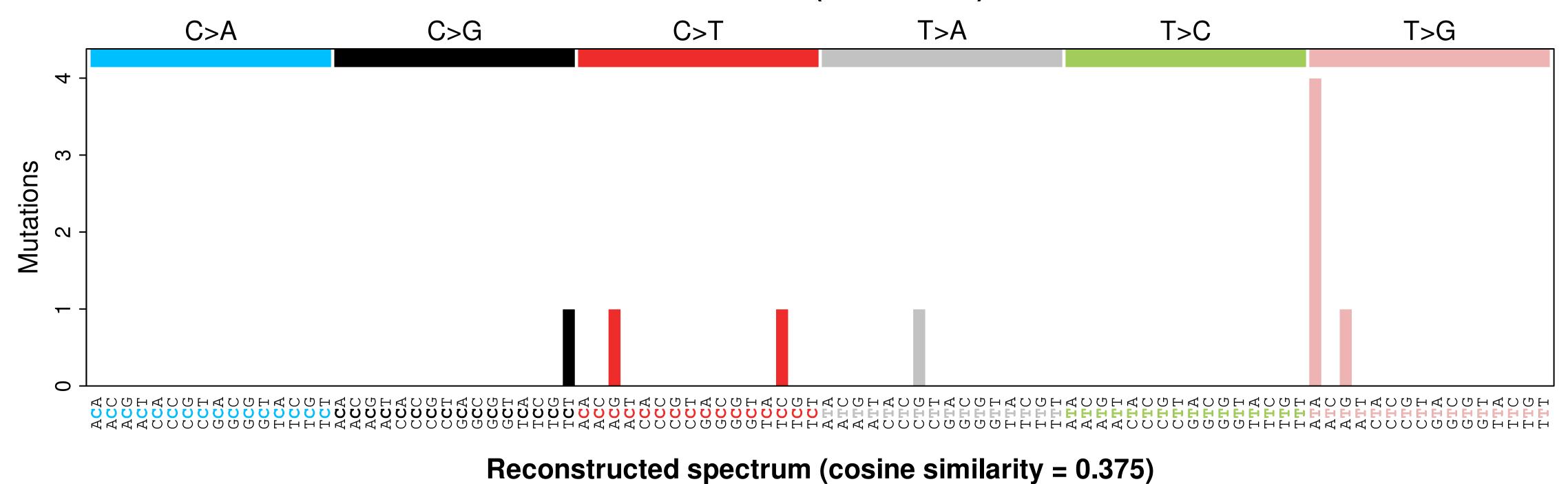
CATD0575a (6 mutations)



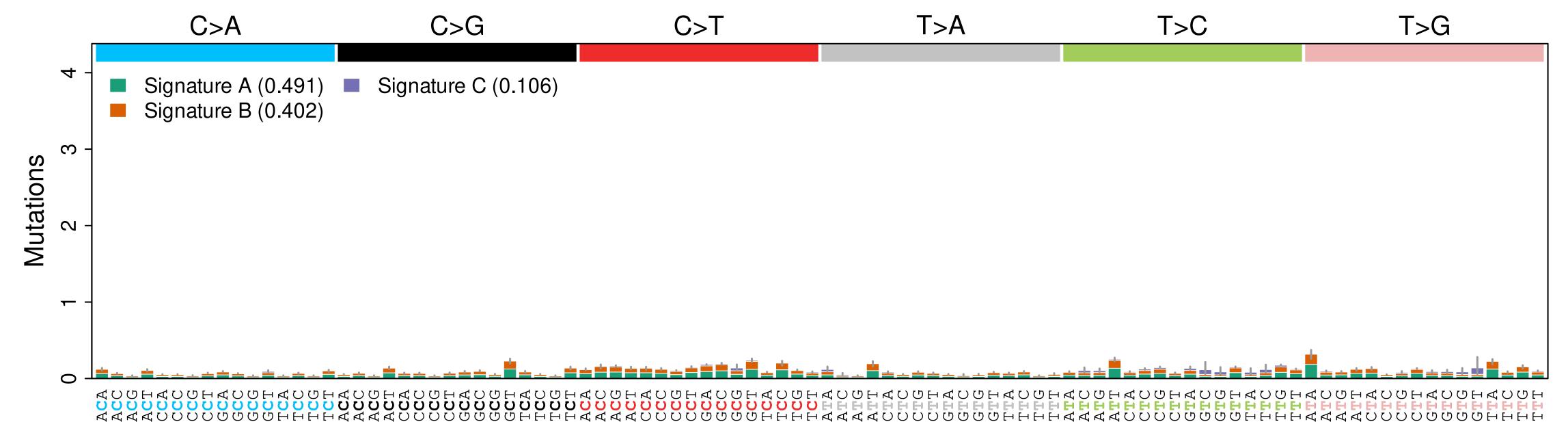




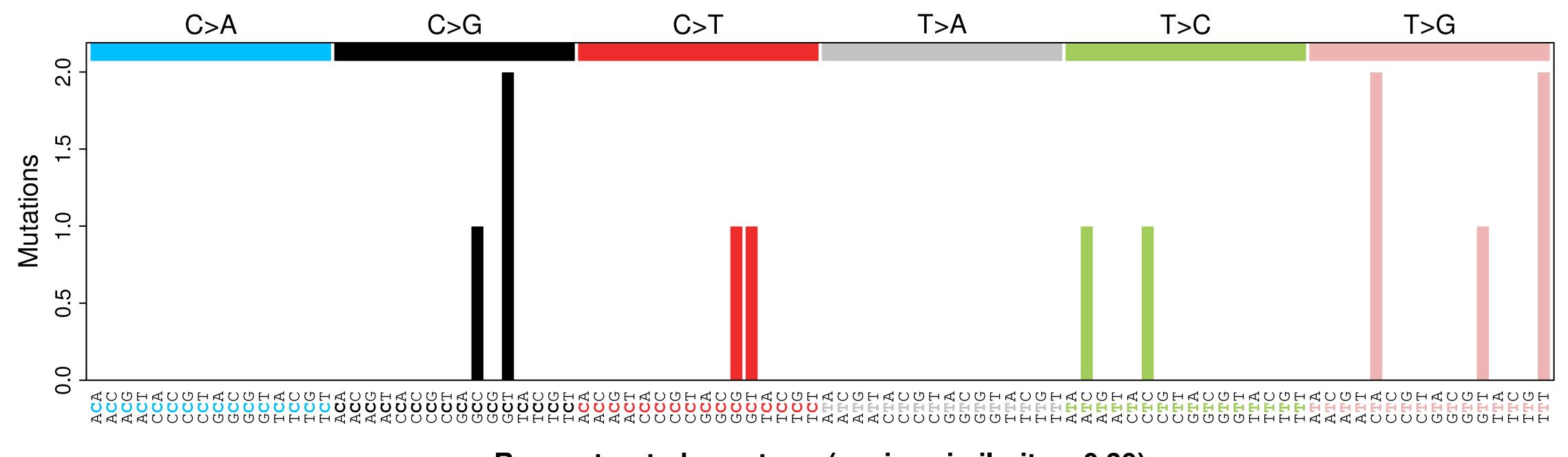
CATD0584a (9 mutations)



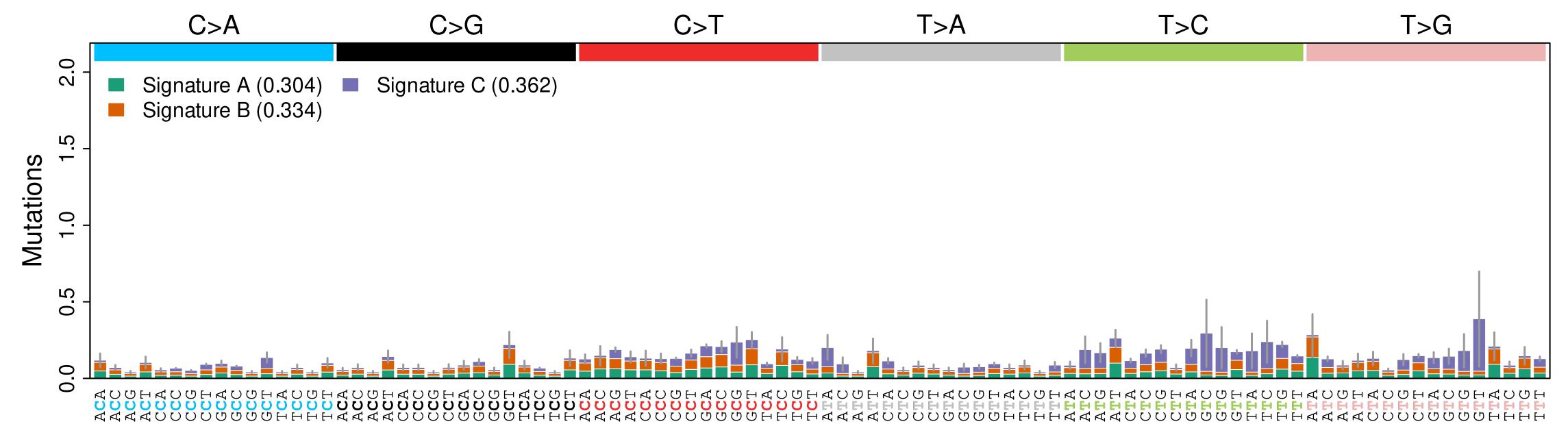




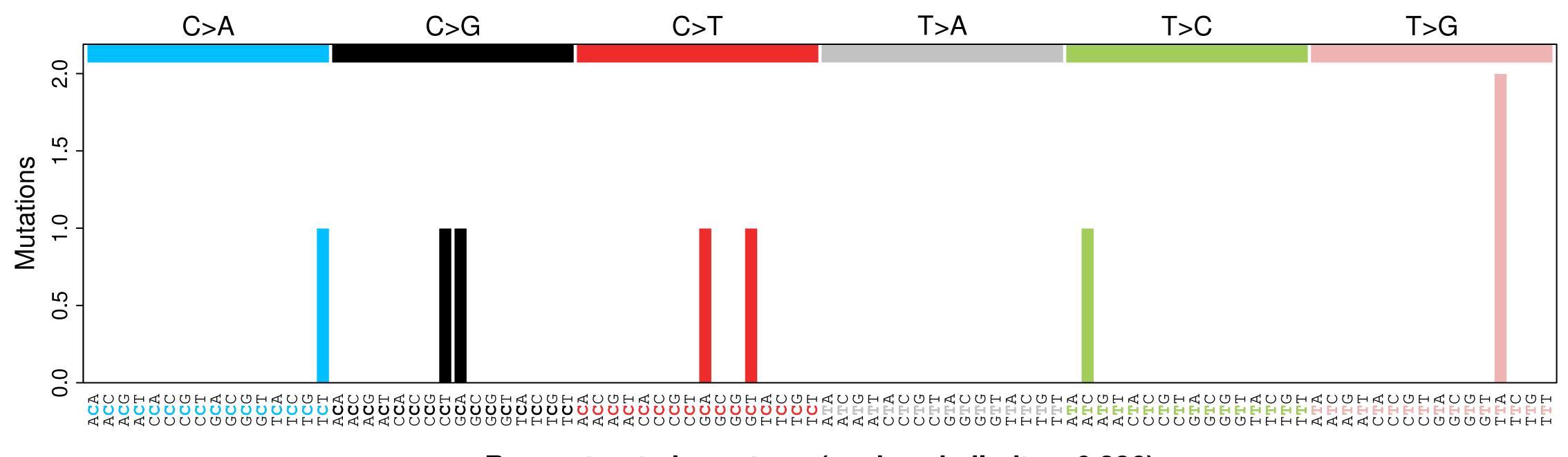
CATD0600a (12 mutations)

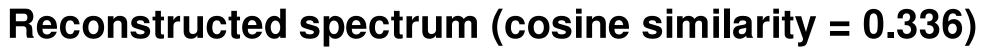


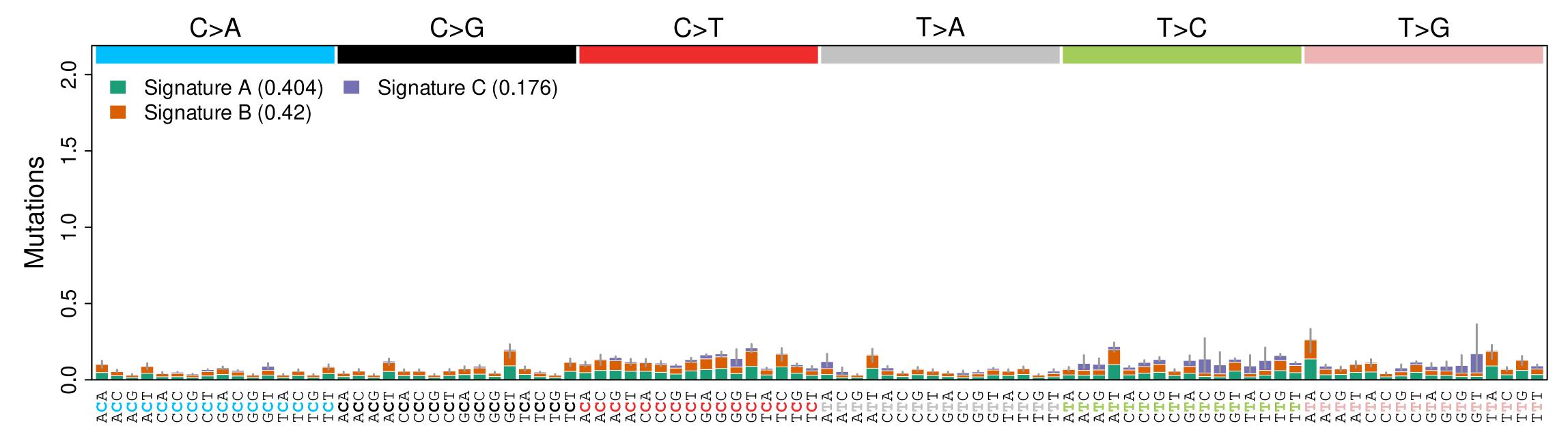
Reconstructed spectrum (cosine similarity = 0.39)



CATD0576a (8 mutations)

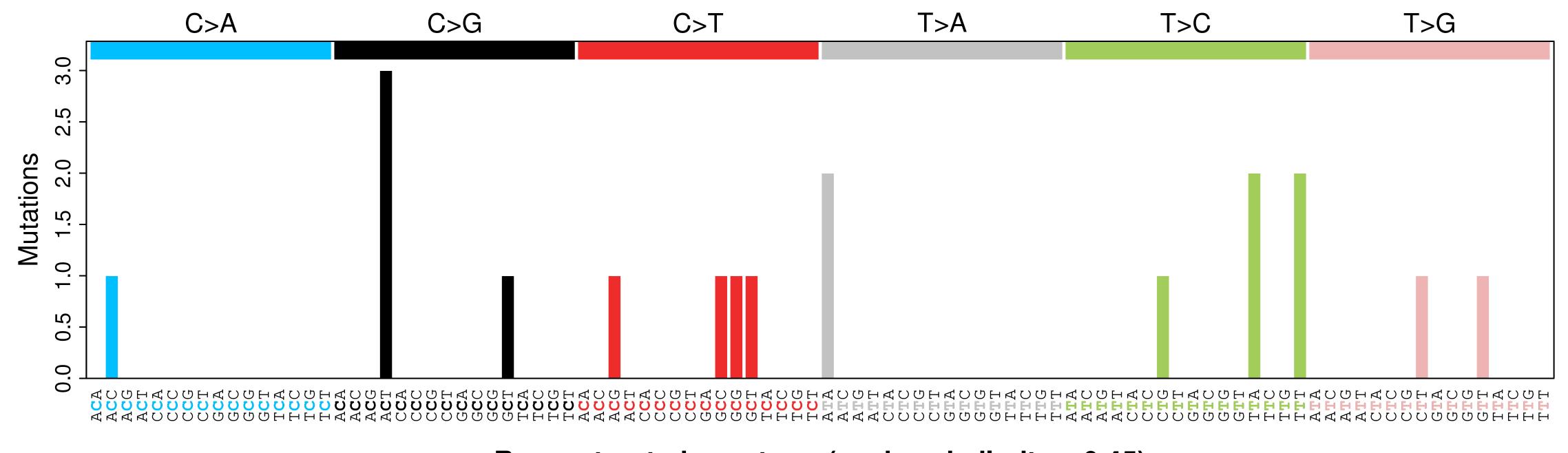




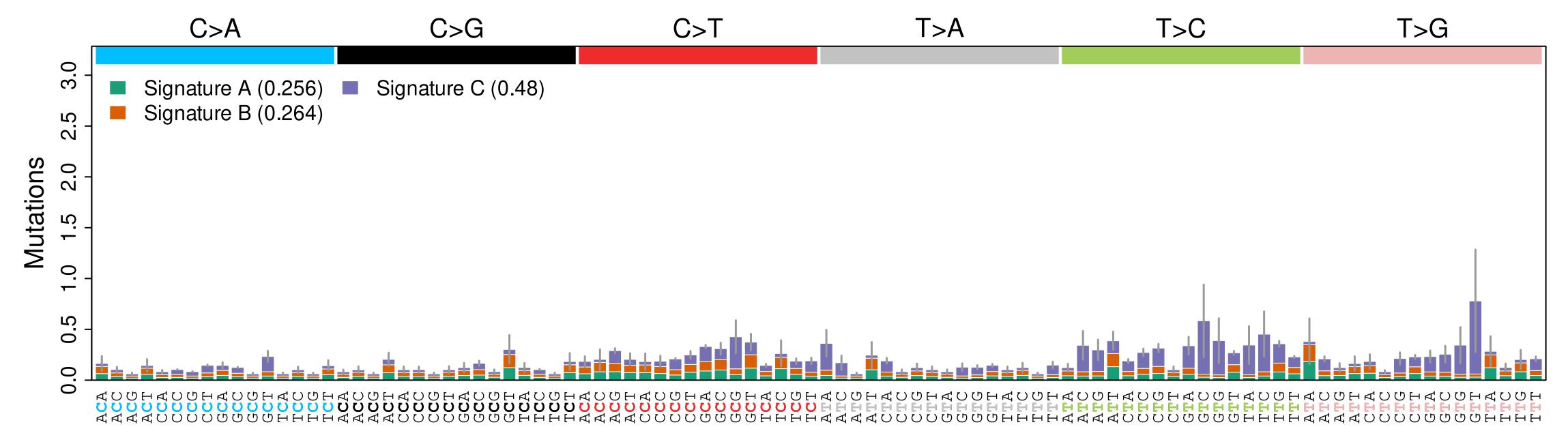


CATD0679a (6 mutations) T>G C>A C>G C>T T>A T>C 2.0 1.5 Mutations 0.5 0.0 **Reconstructed spectrum (cosine similarity = 0.317)** C>A C>G T>A T>C T>G C>T 2.0 Signature A (0.377) Signature C (0.231) Signature B (0.392) 5 Mutations 2 0

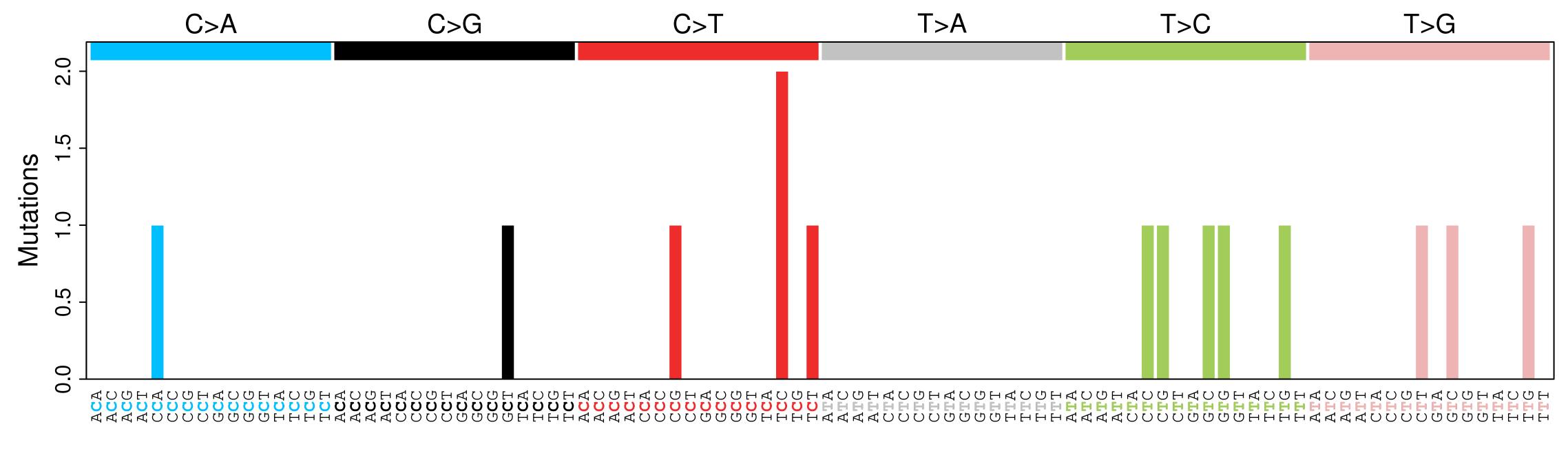
CATD0591a (18 mutations)



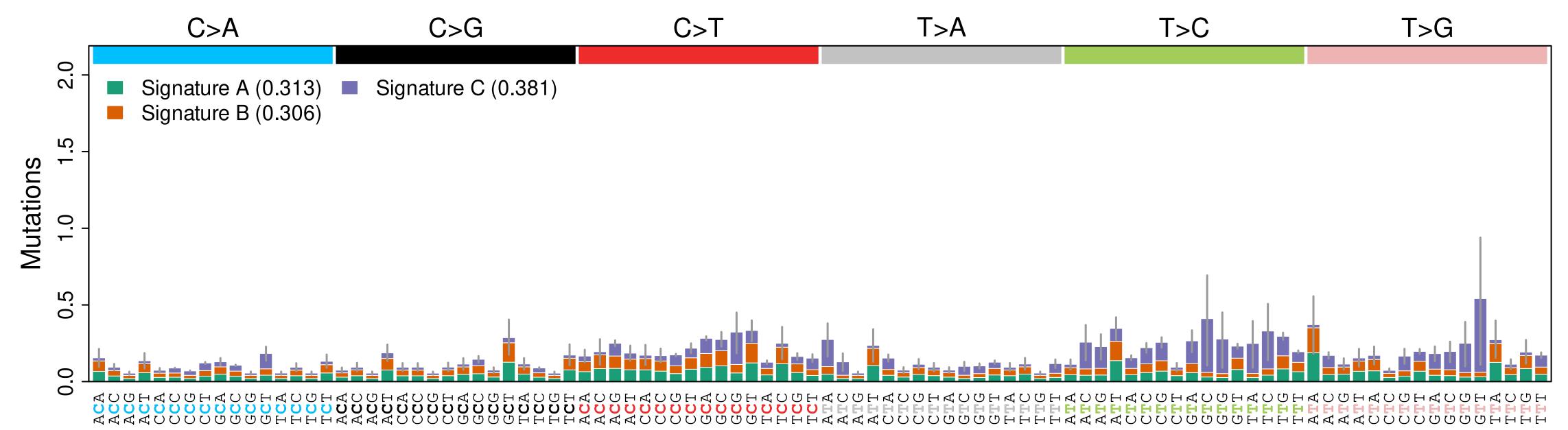
Reconstructed spectrum (cosine similarity = 0.45)



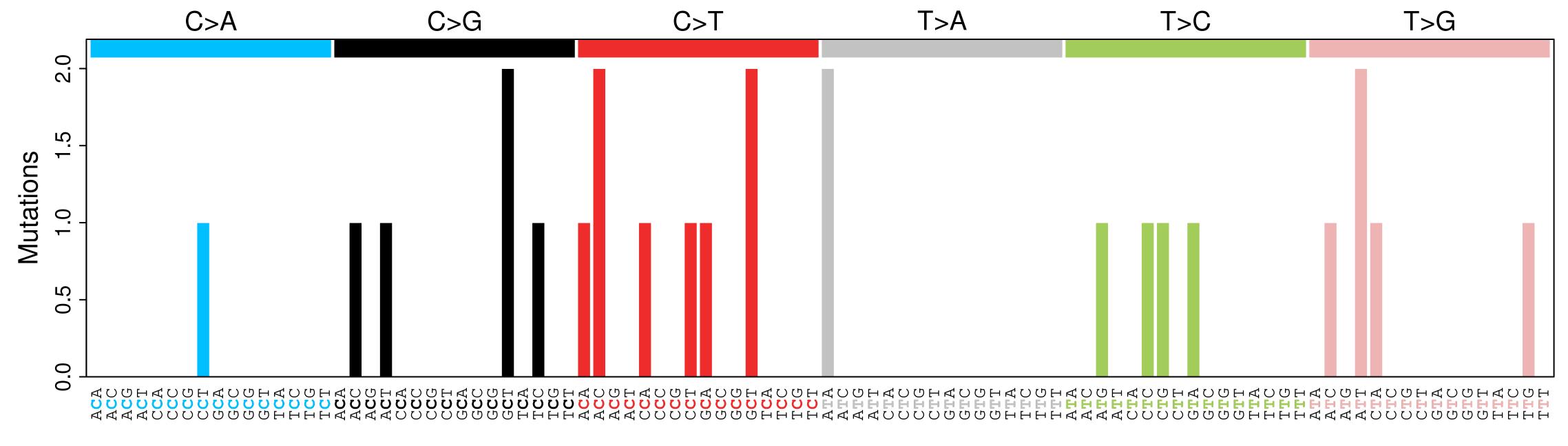
CATD0686a (14 mutations)



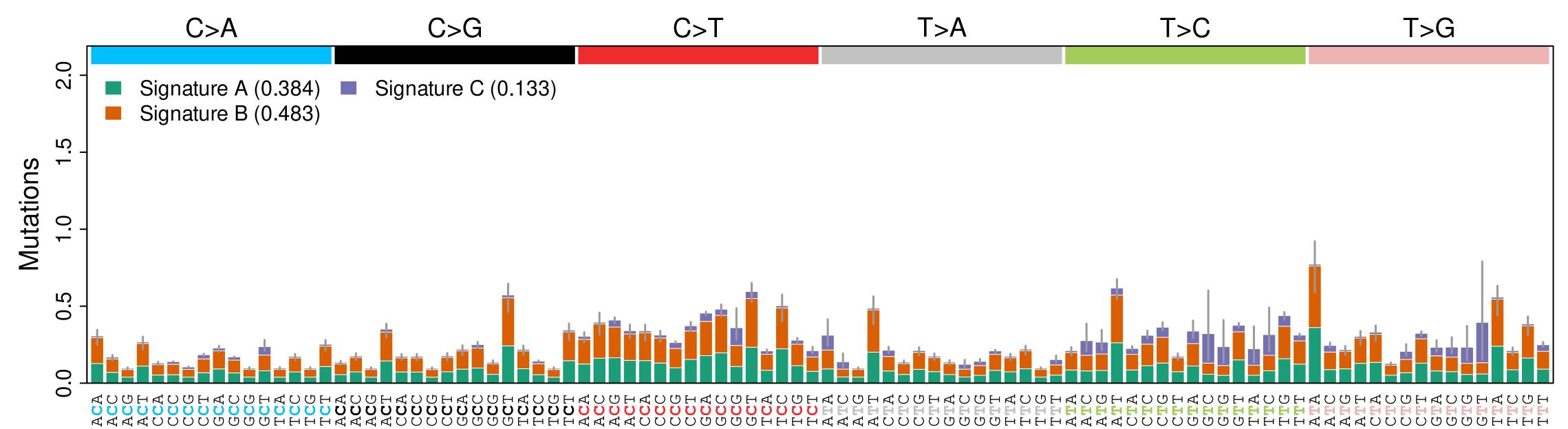
Reconstructed spectrum (cosine similarity = 0.434)



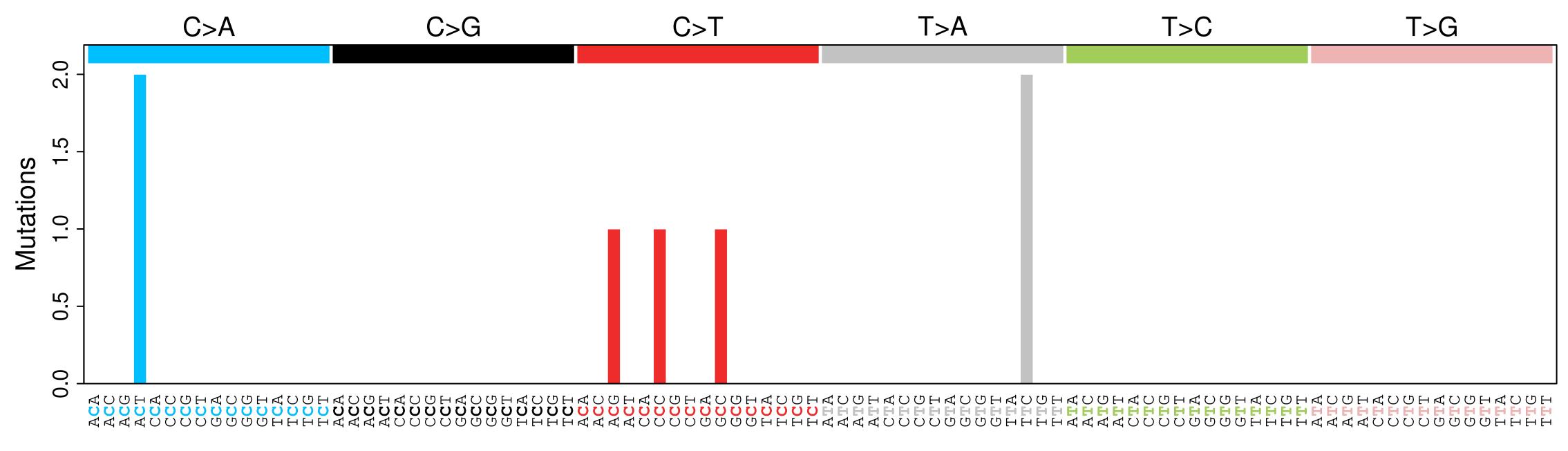
CATD0734a (25 mutations)



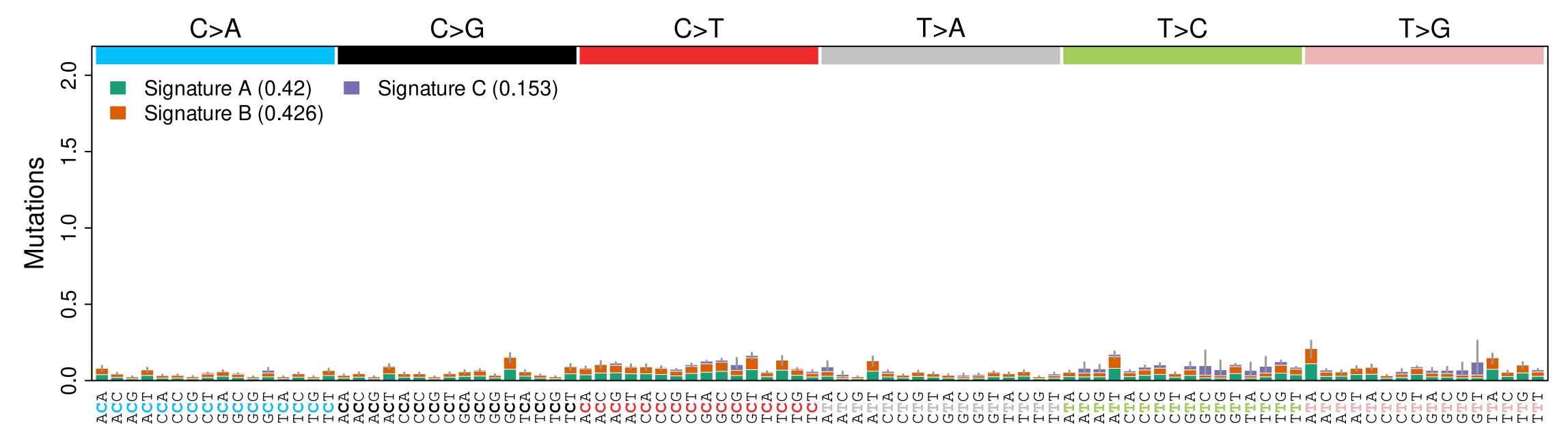
Reconstructed spectrum (cosine similarity = 0.527)



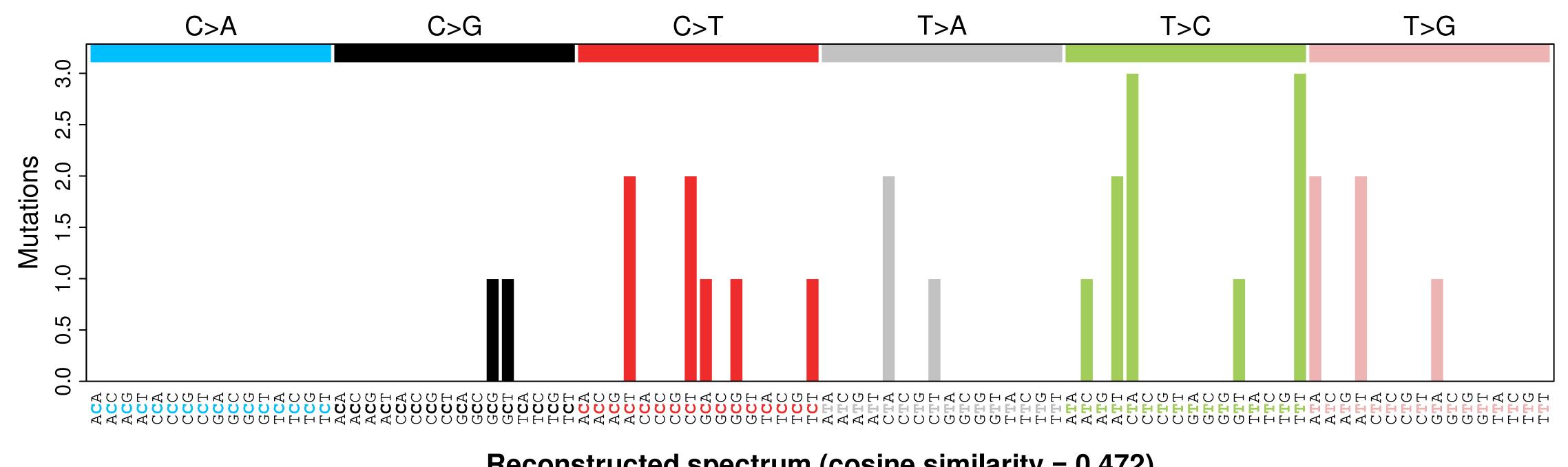
CATD0678a (7 mutations)



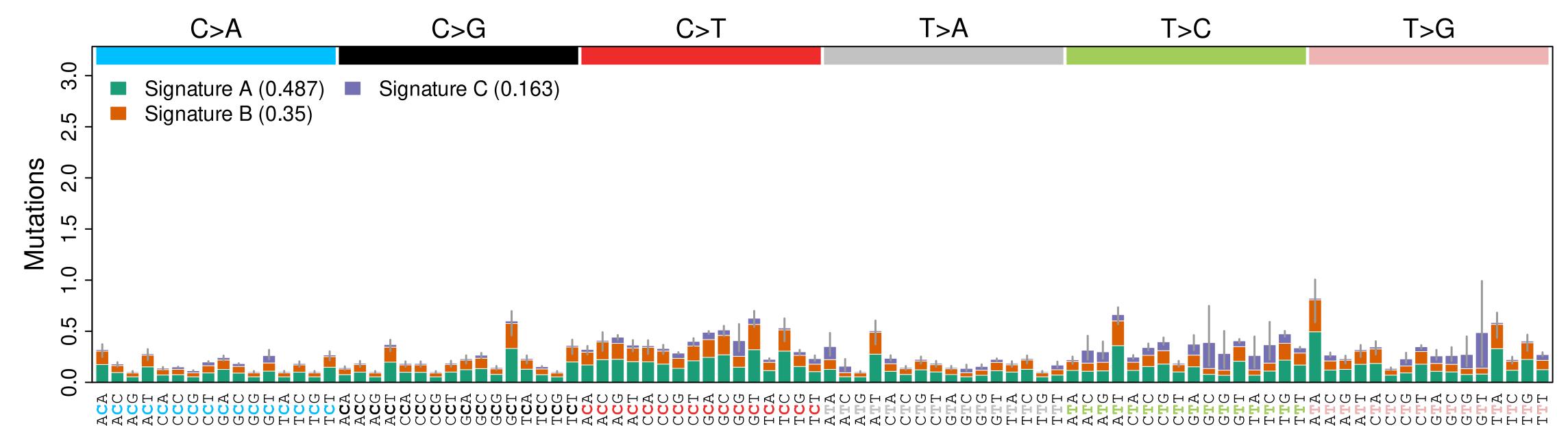
Reconstructed spectrum (cosine similarity = 0.227)



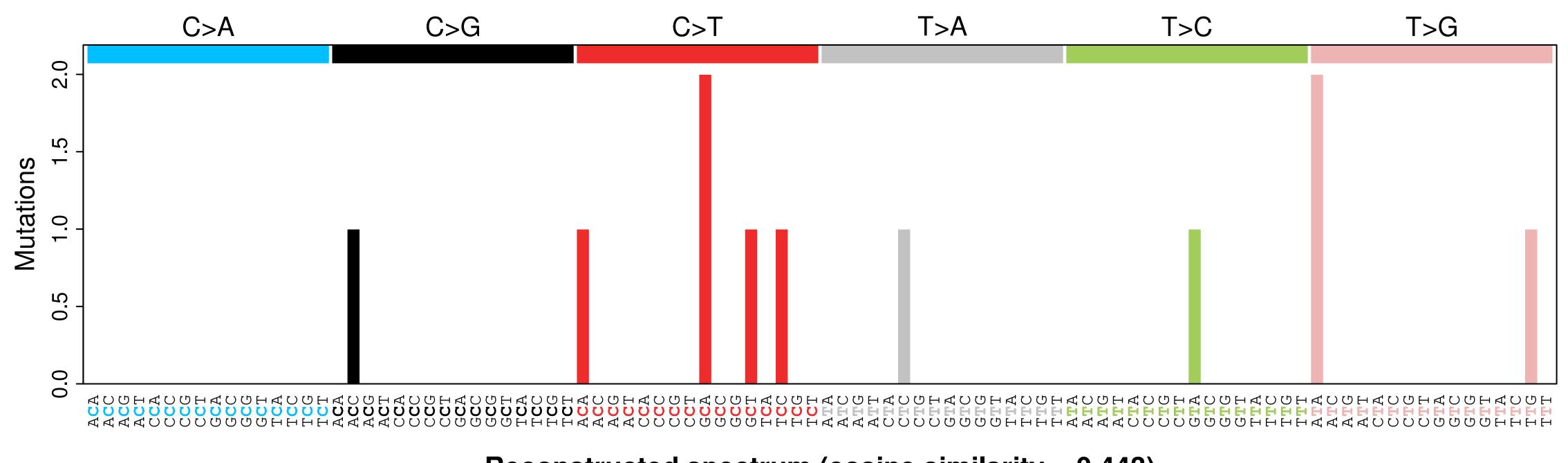
CATD0729a (27 mutations)



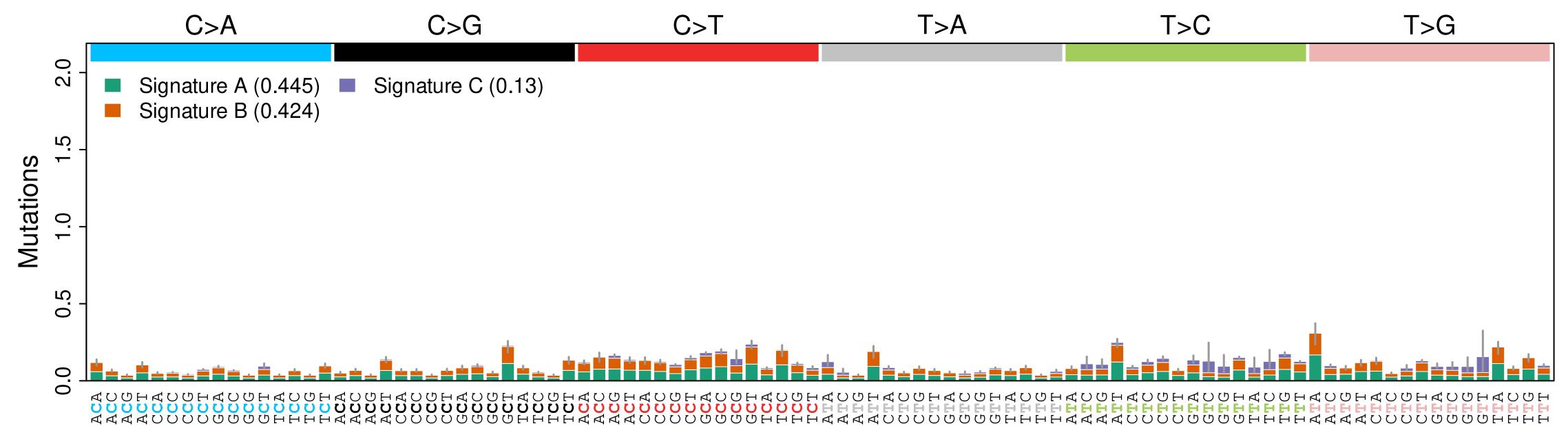




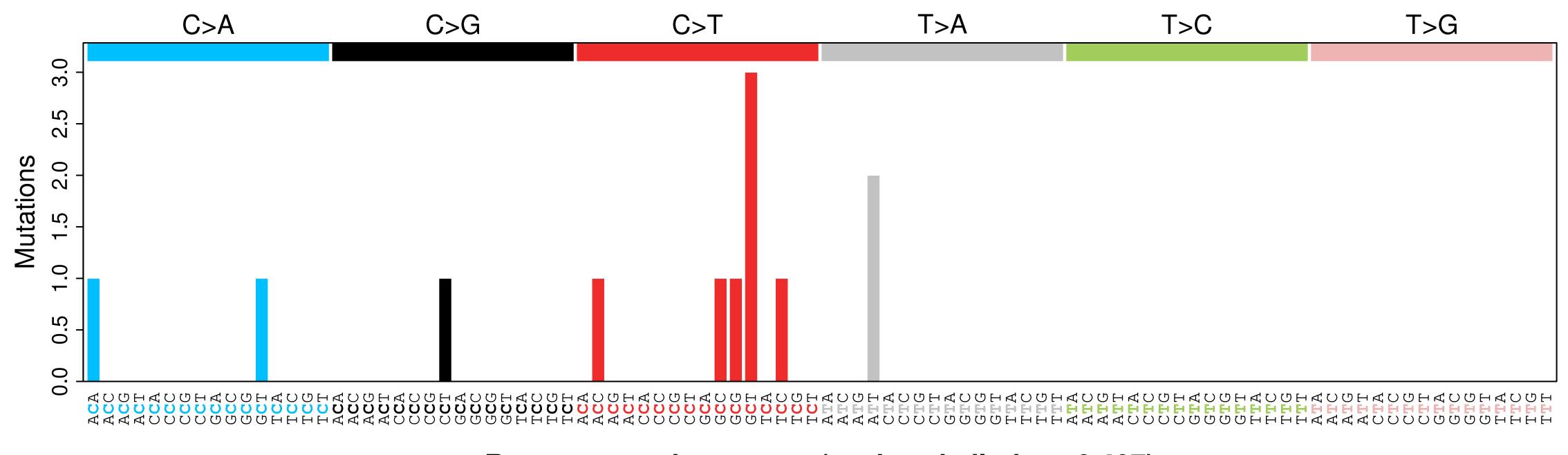
CATD0609a (11 mutations)



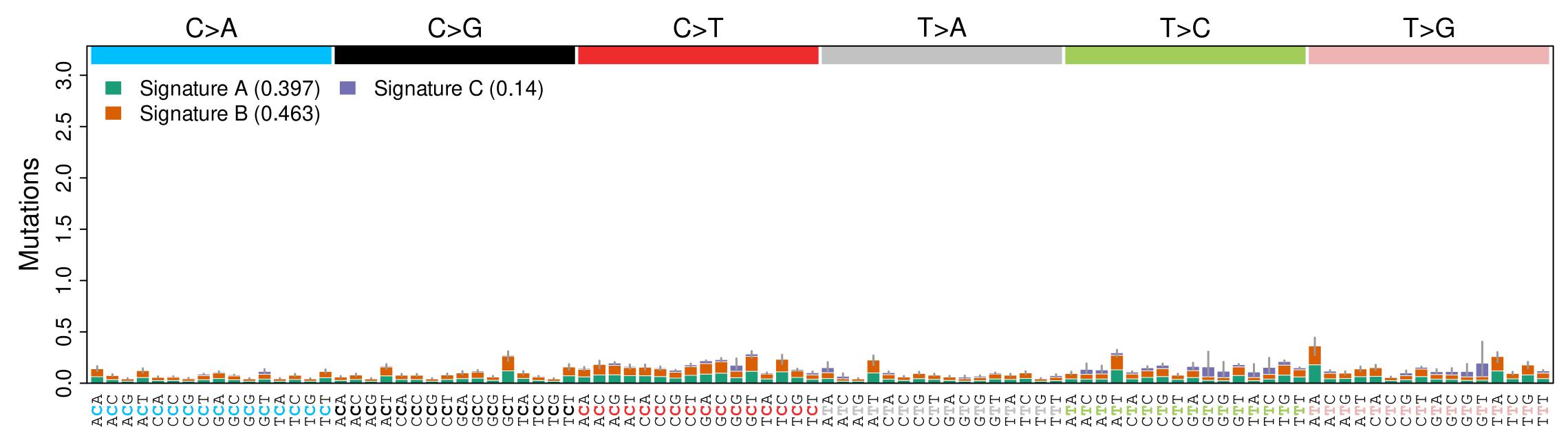
Reconstructed spectrum (cosine similarity = 0.443)



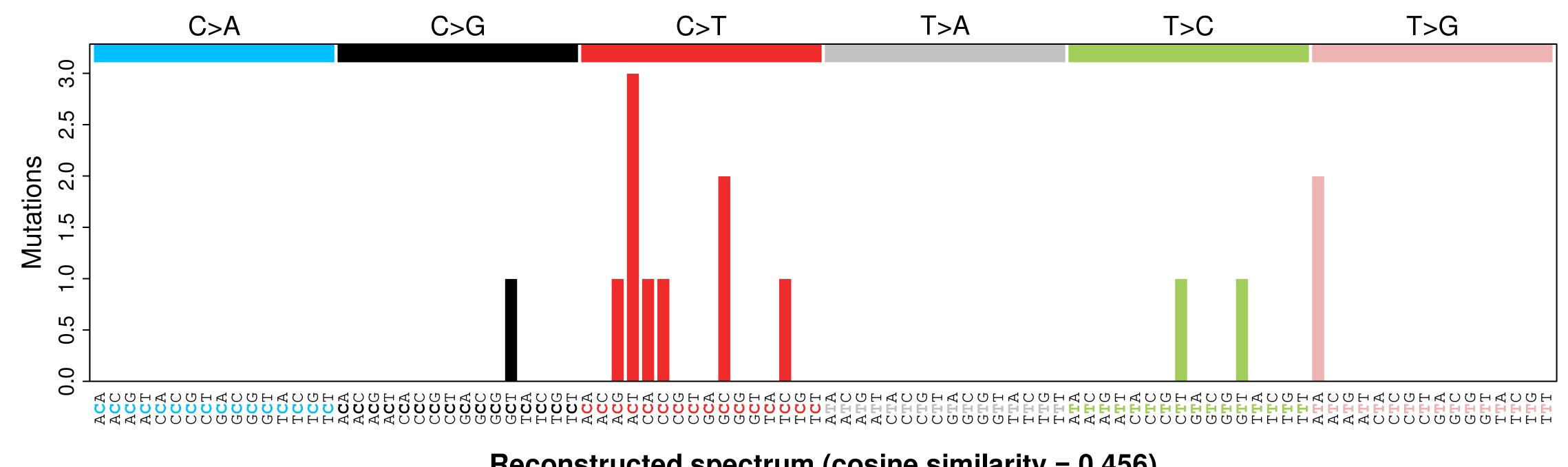
CATD0601a (12 mutations)



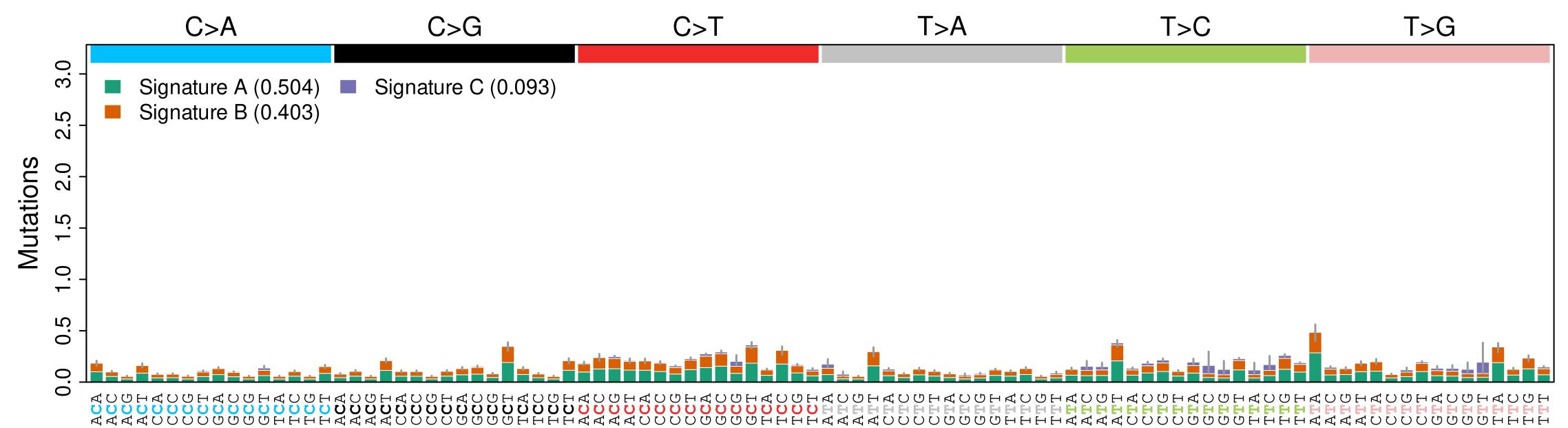




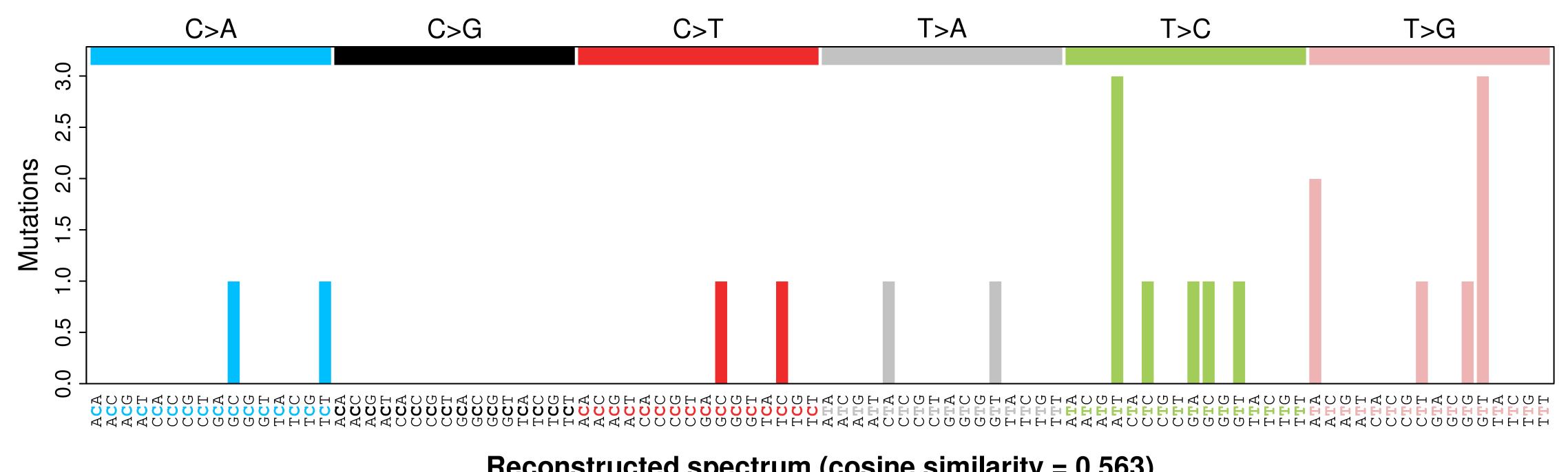
CATD0731a (14 mutations)

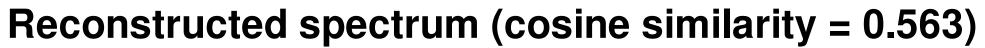


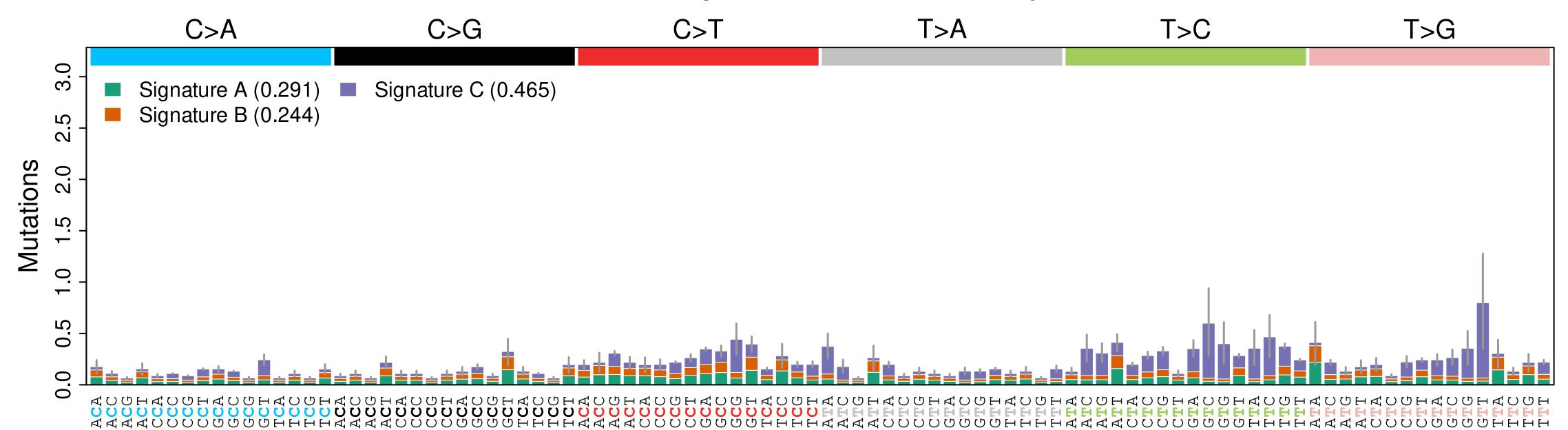
Reconstructed spectrum (cosine similarity = 0.456)



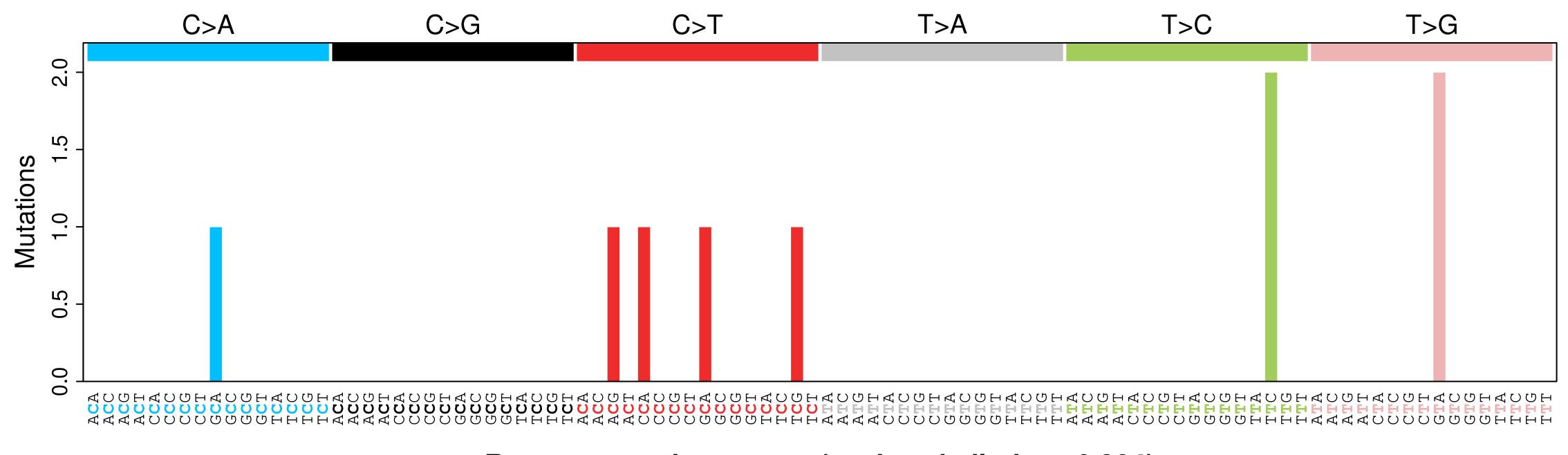
CATD0723a (20 mutations)

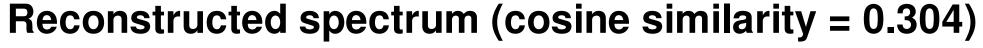


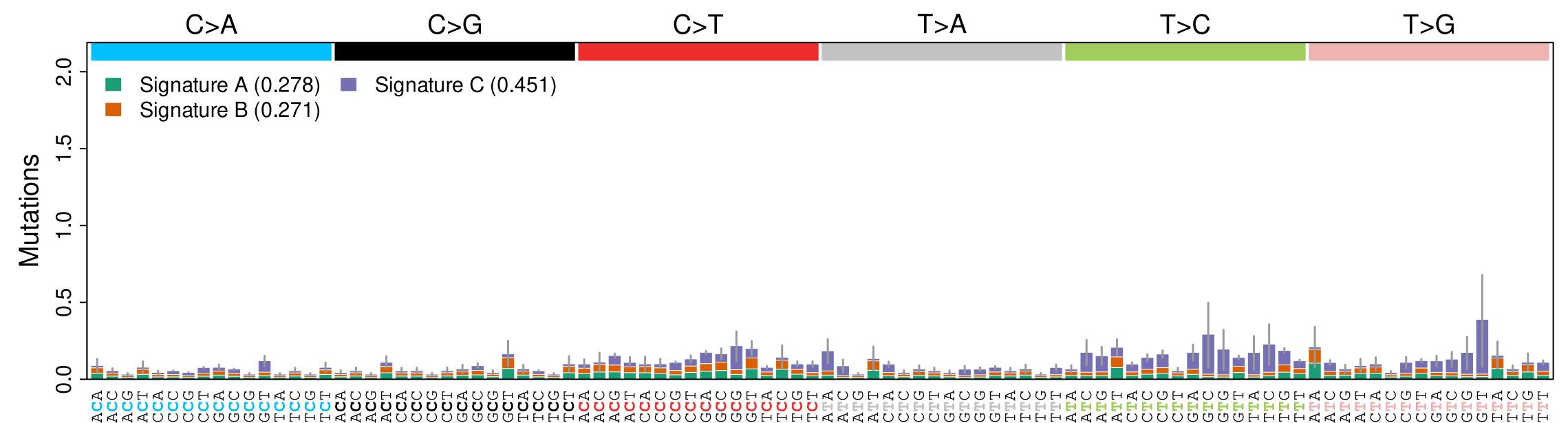




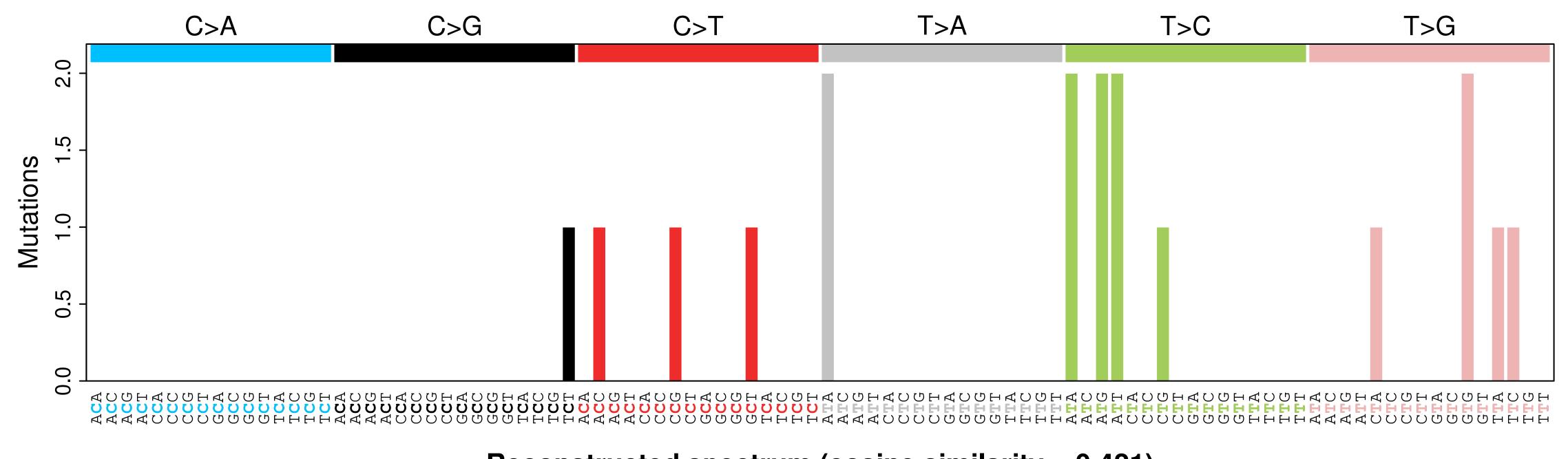
CATD0730a (9 mutations)



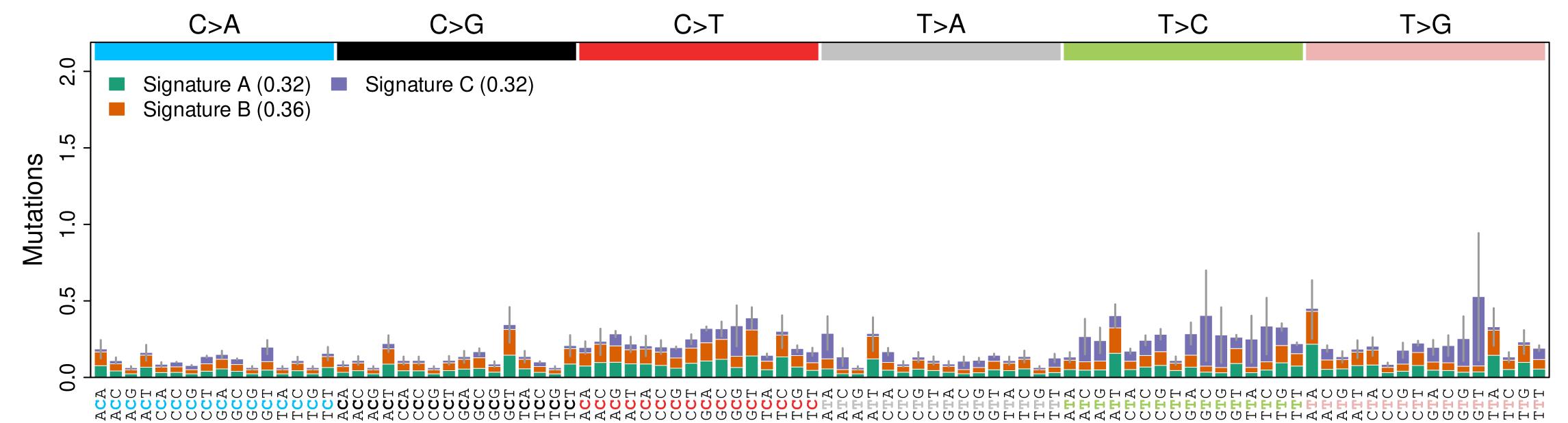




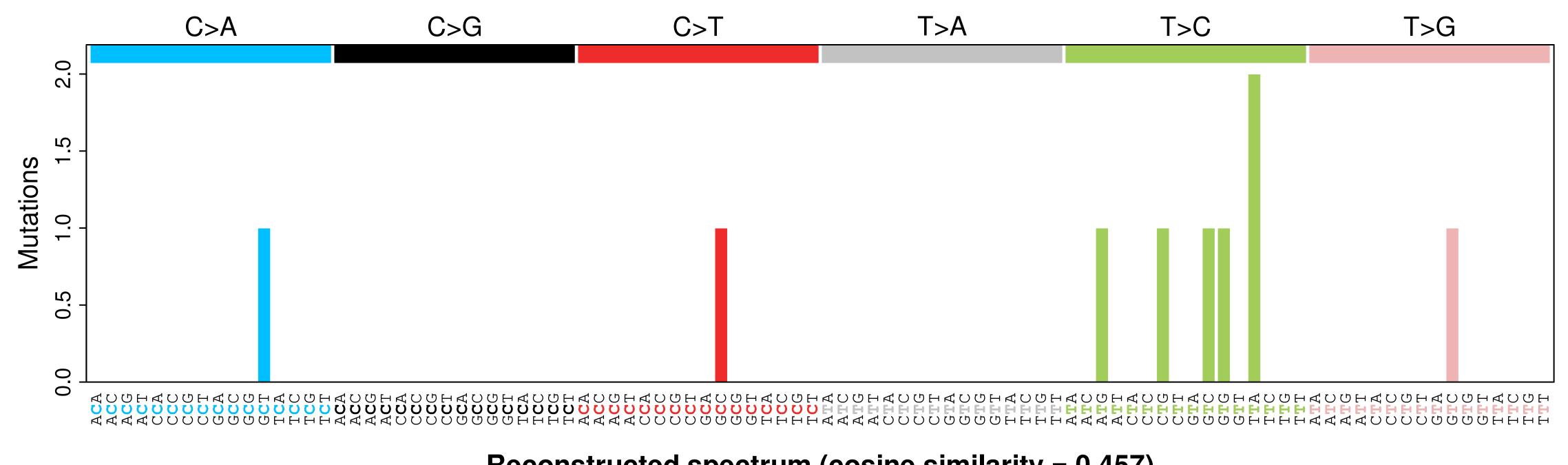
CATD0586a (18 mutations)

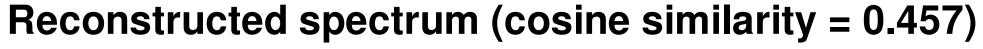


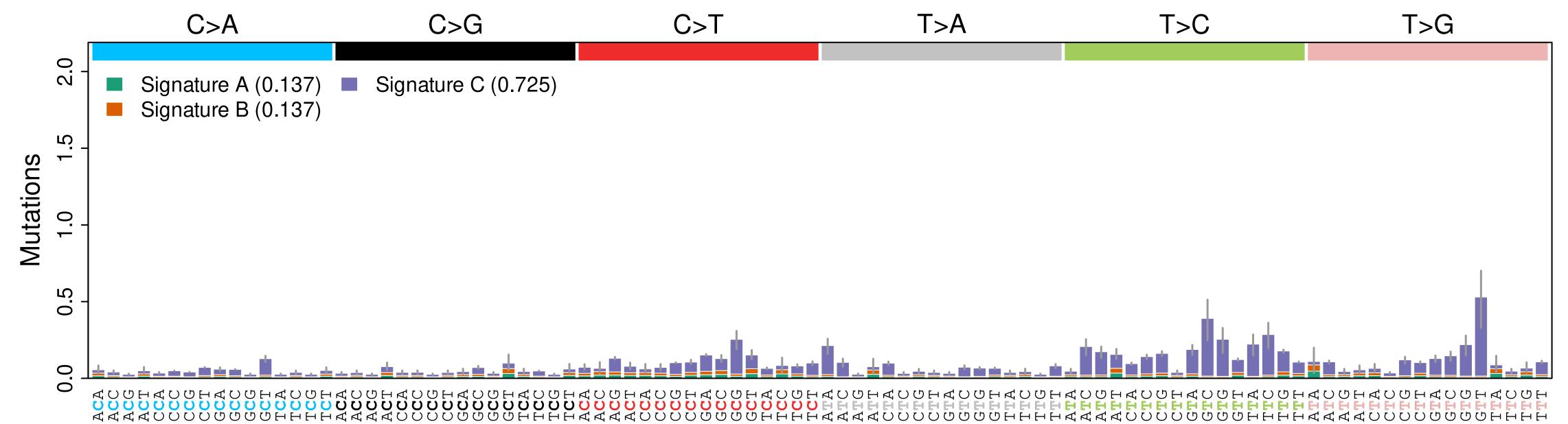
Reconstructed spectrum (cosine similarity = 0.421)



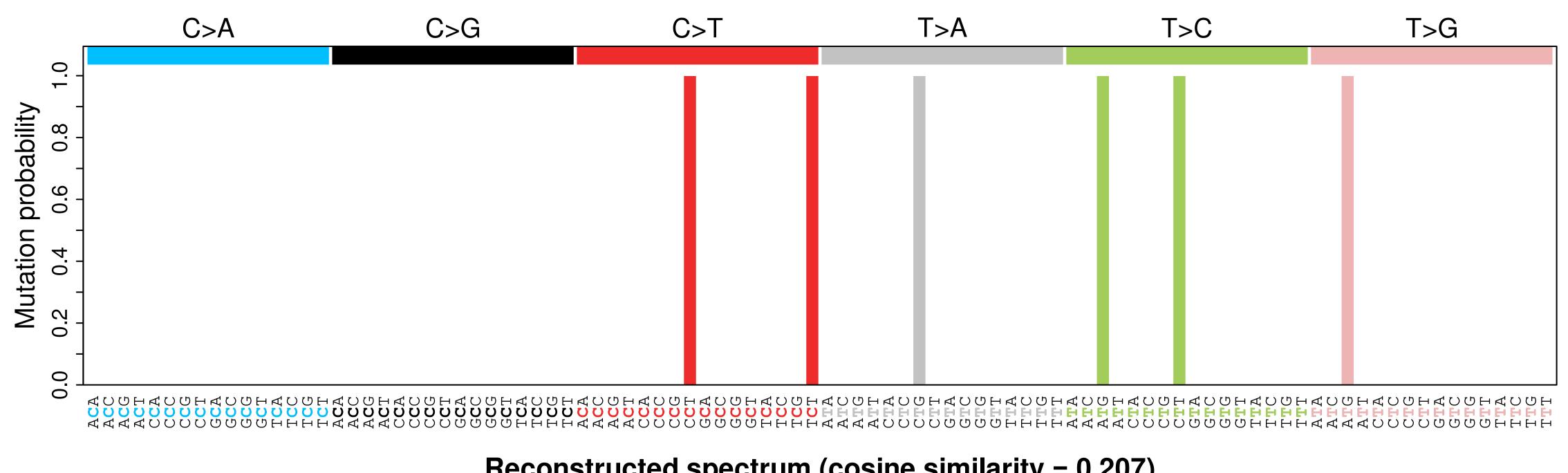
CATD0733a (9 mutations)



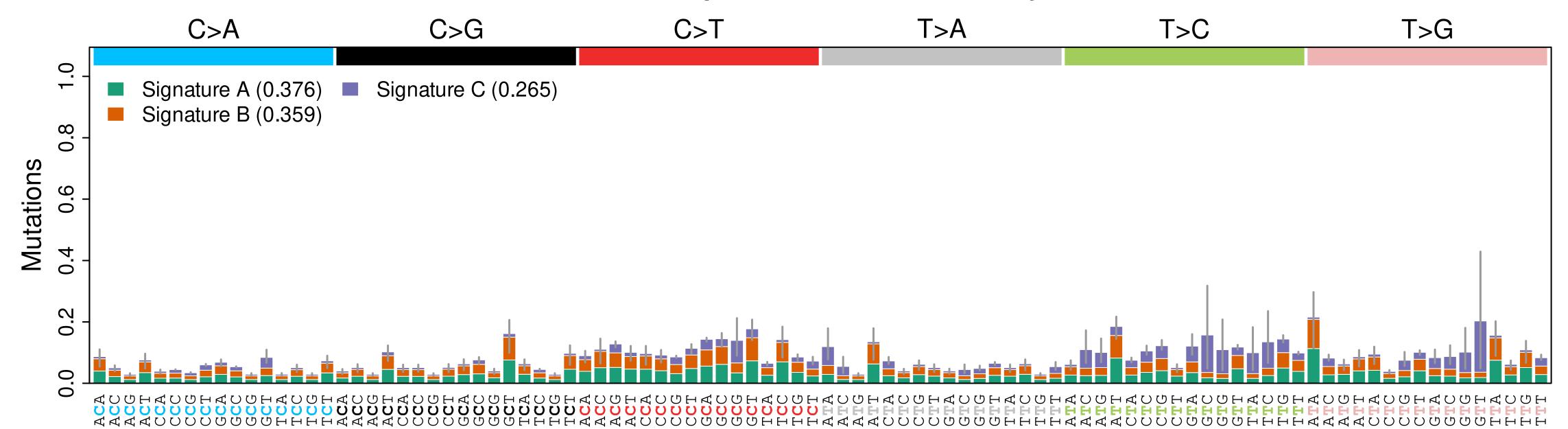




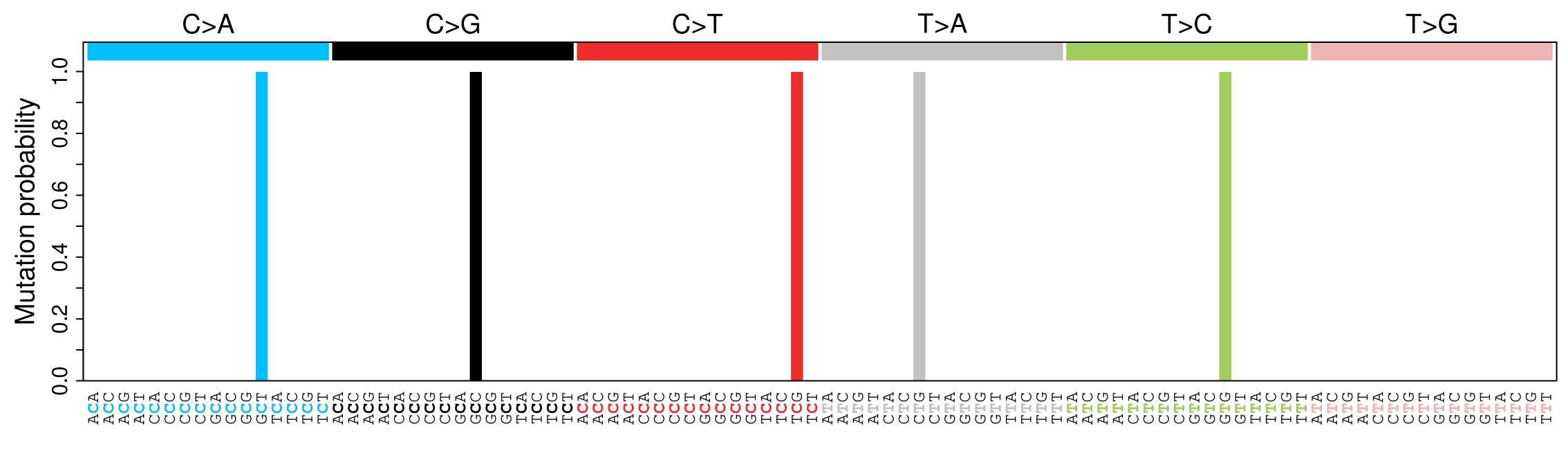
CATD0687a

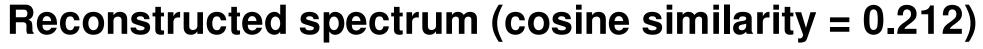


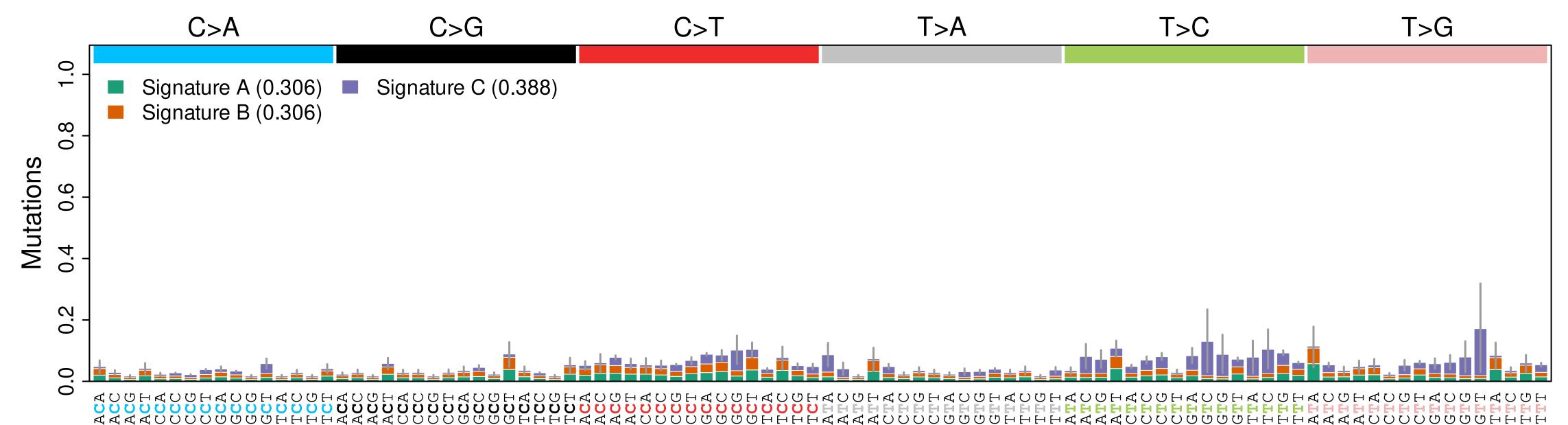




CATD0578a

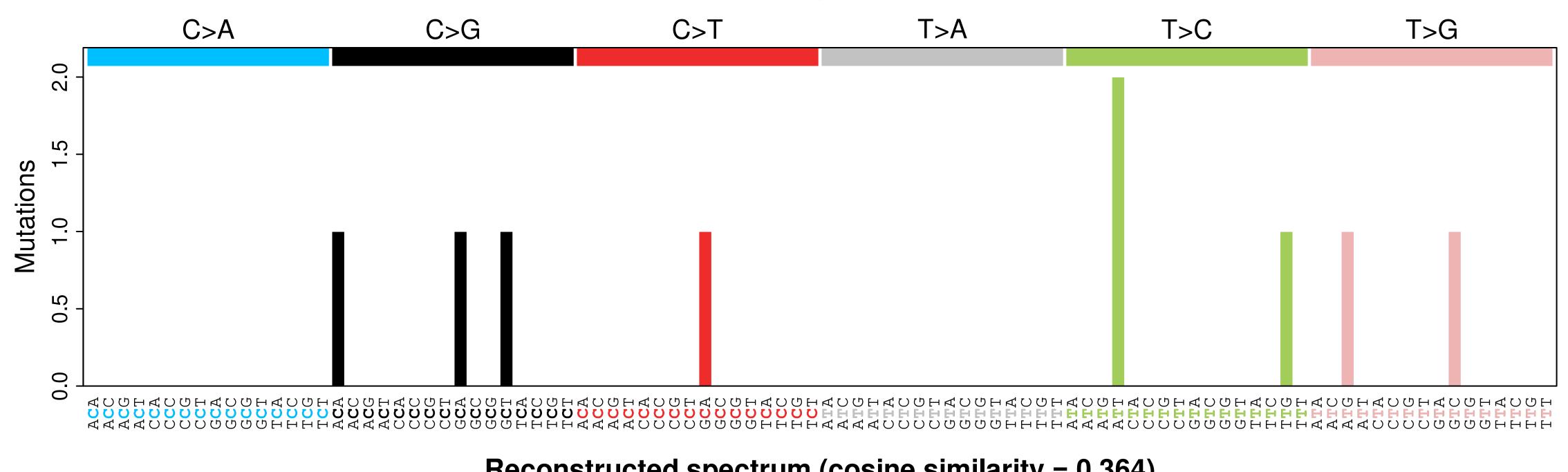


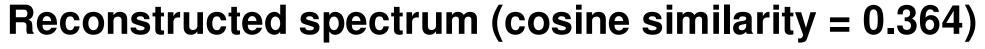


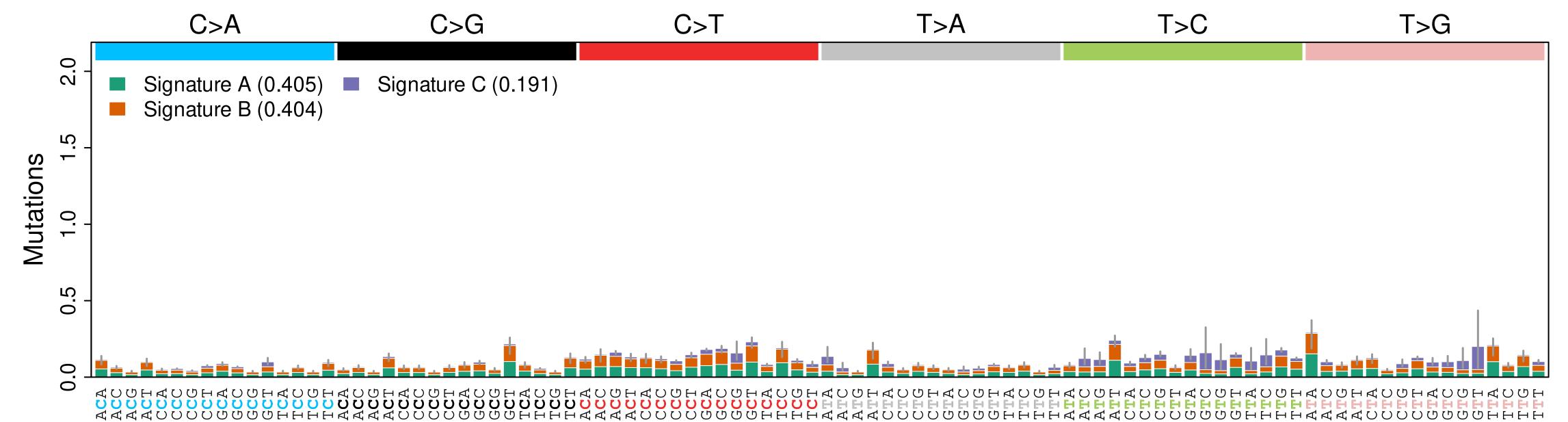


CATD0587a (4 mutations) T>G C>A C>G C>T T>A T>C 2.0 5 Mutations 0.5 0.0 **Reconstructed spectrum (cosine similarity = 0.161)** C>A C>G T>A T>C T>G C>T 2.0 Signature A (0.328) Signature C (0.349) Signature B (0.323) 5 Mutations 2 0

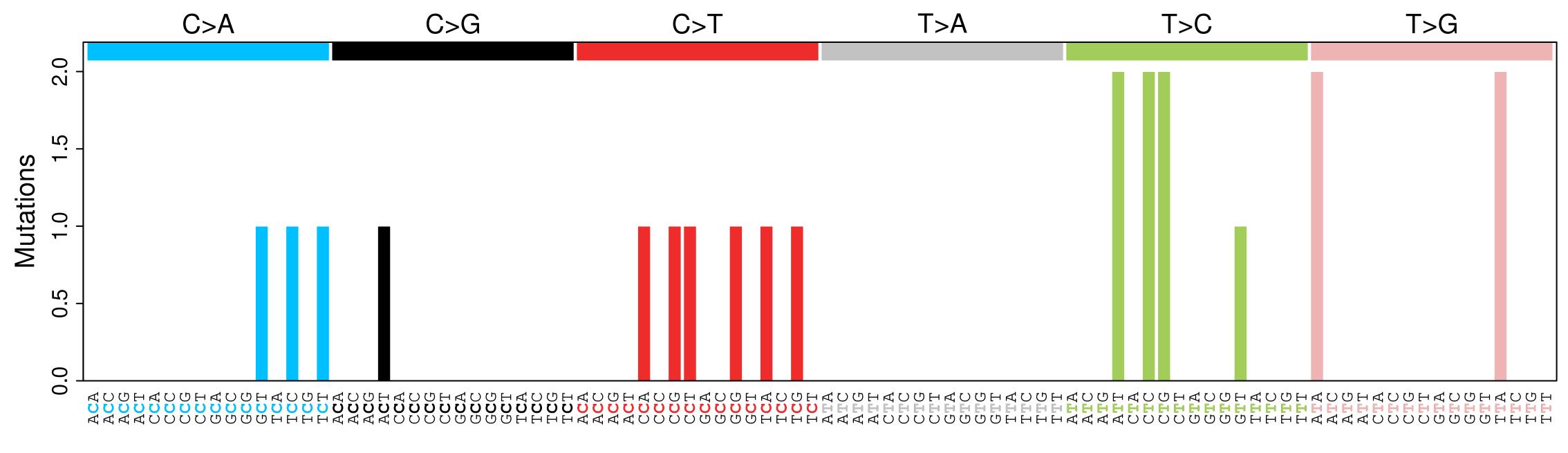
CATD0589a (9 mutations)

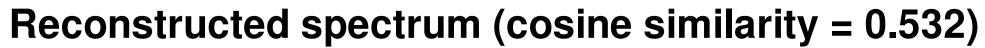


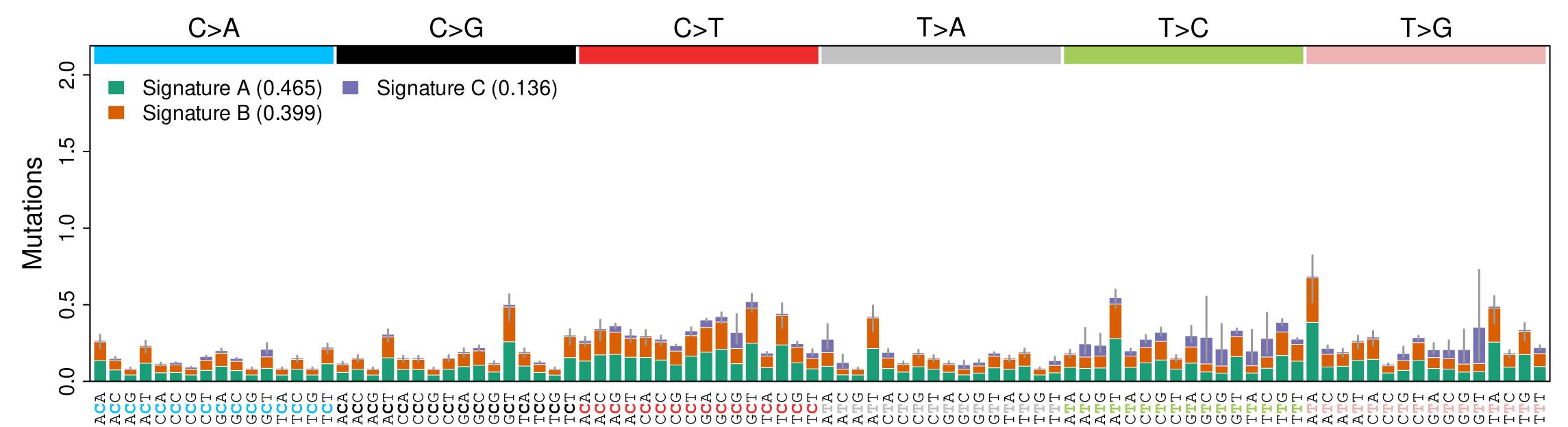




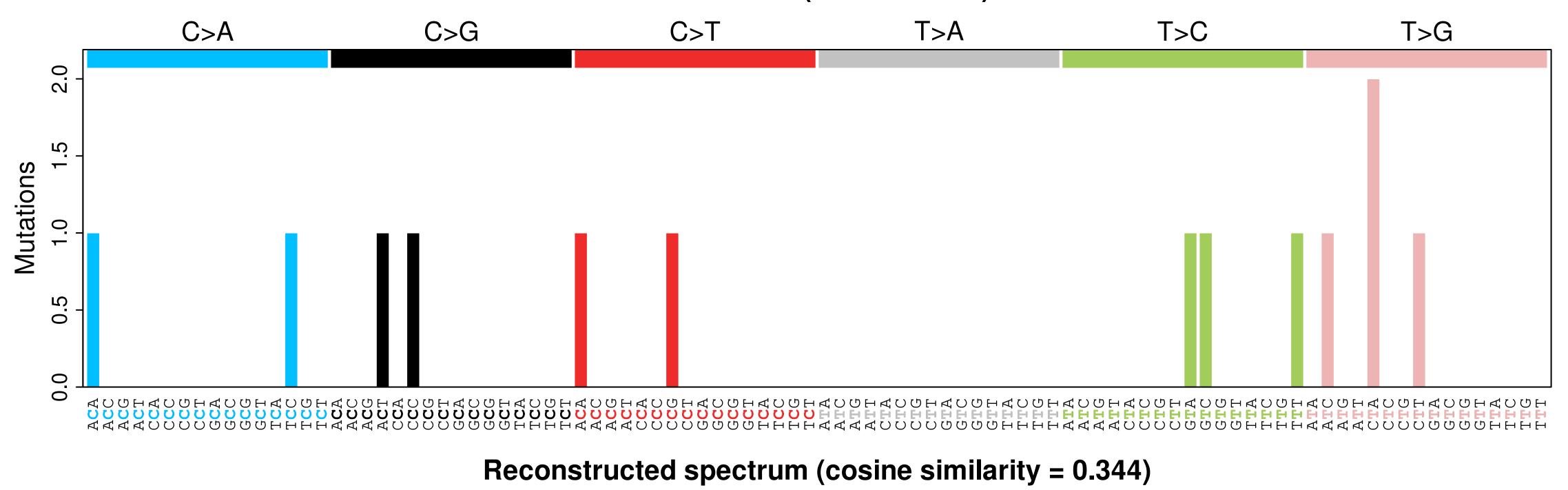
CATD0590a (21 mutations)

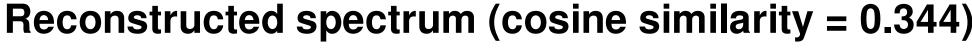


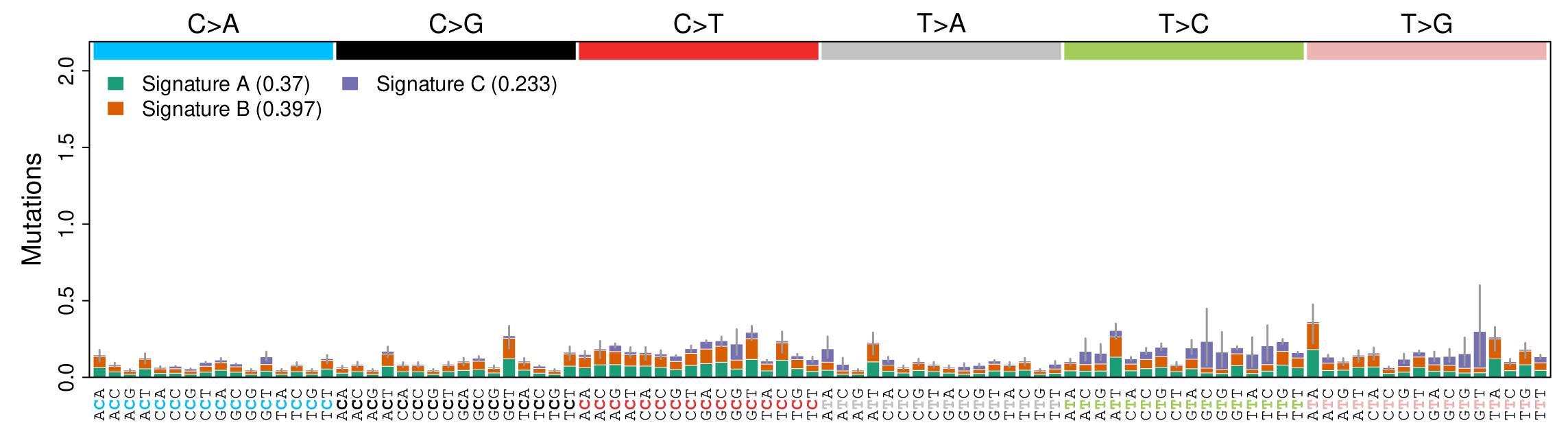




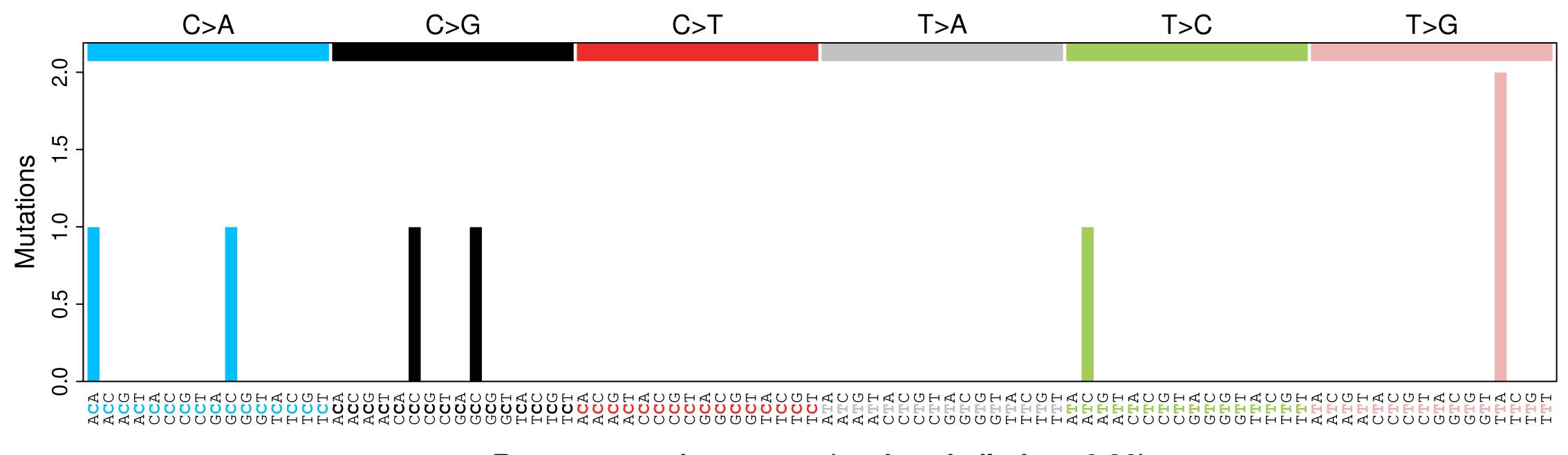
CATD0727a (13 mutations)



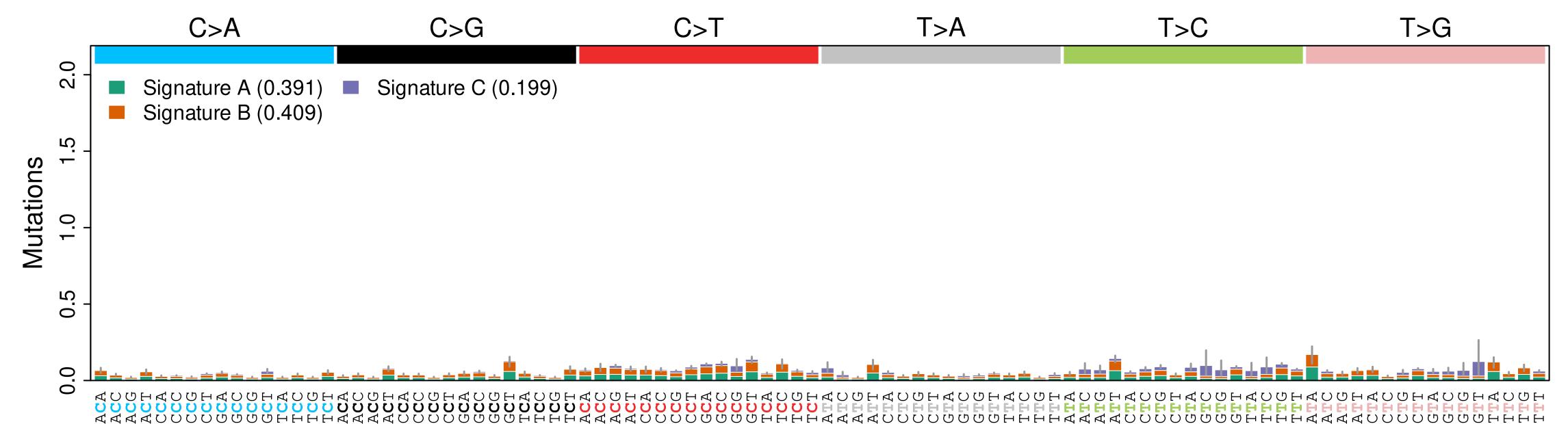




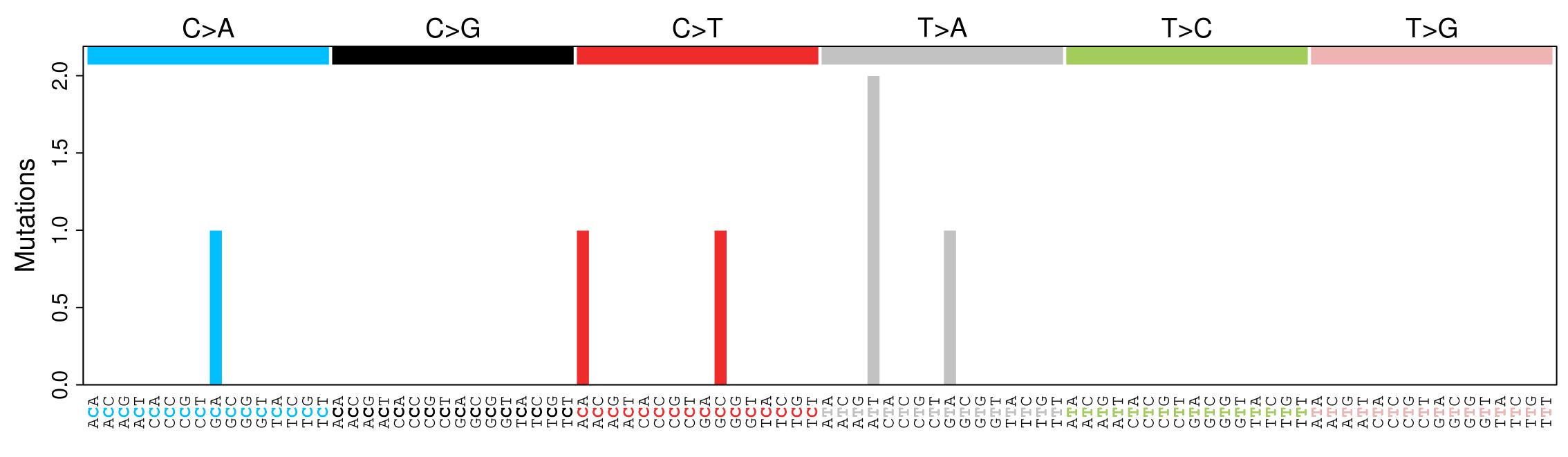
CATD0583a (7 mutations)



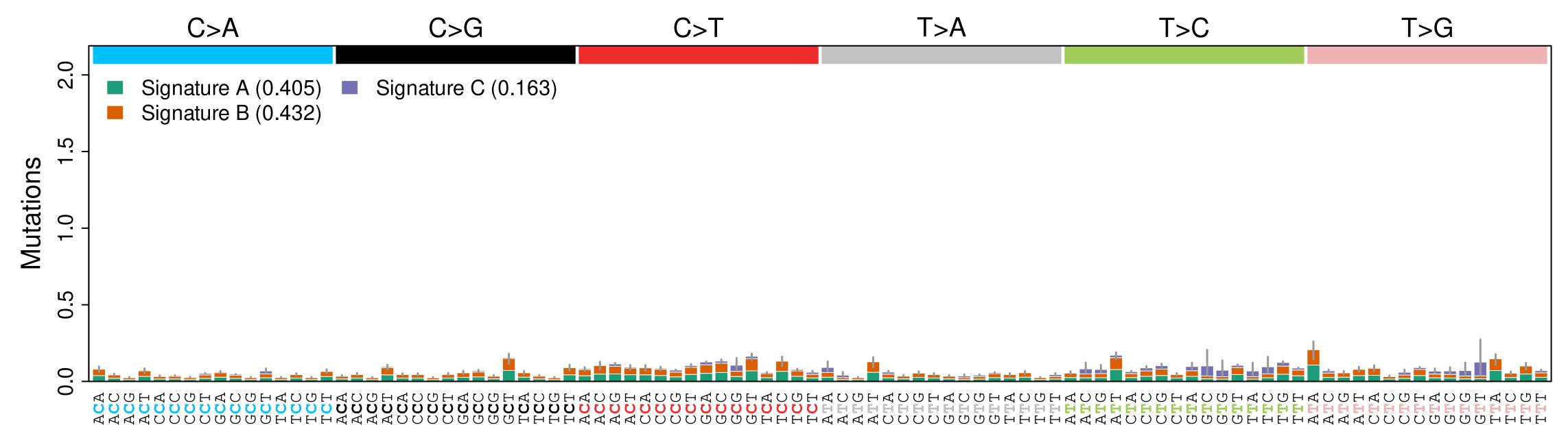
Reconstructed spectrum (cosine similarity = 0.26)



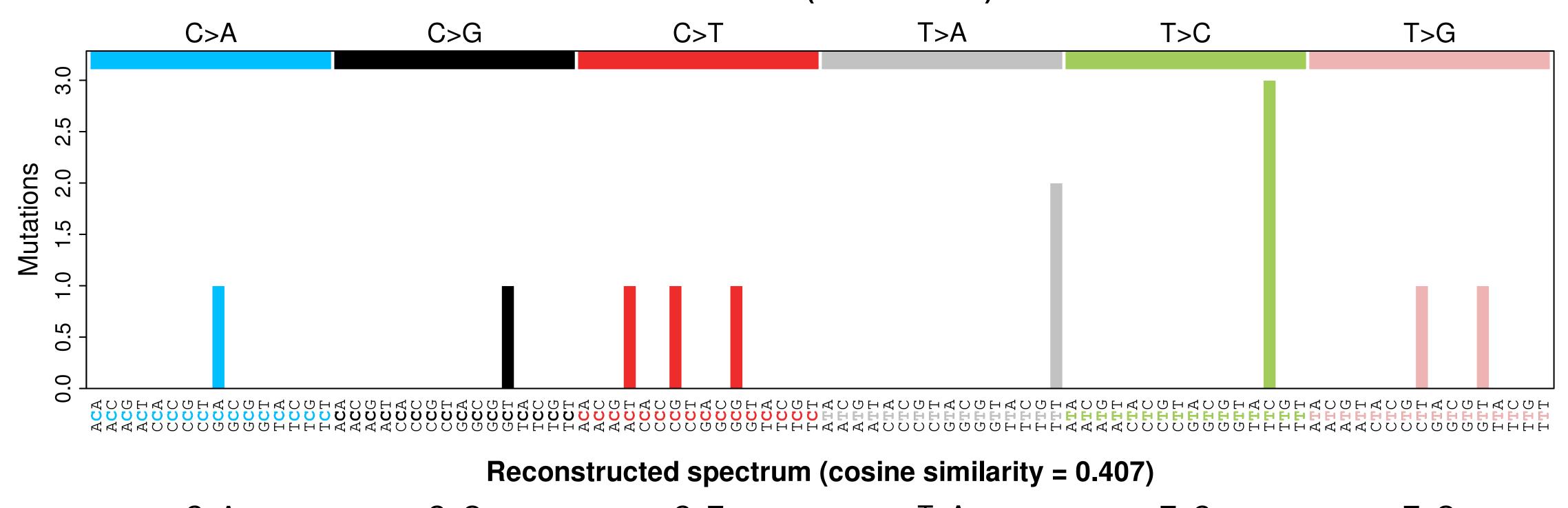
CATD0732a (6 mutations)

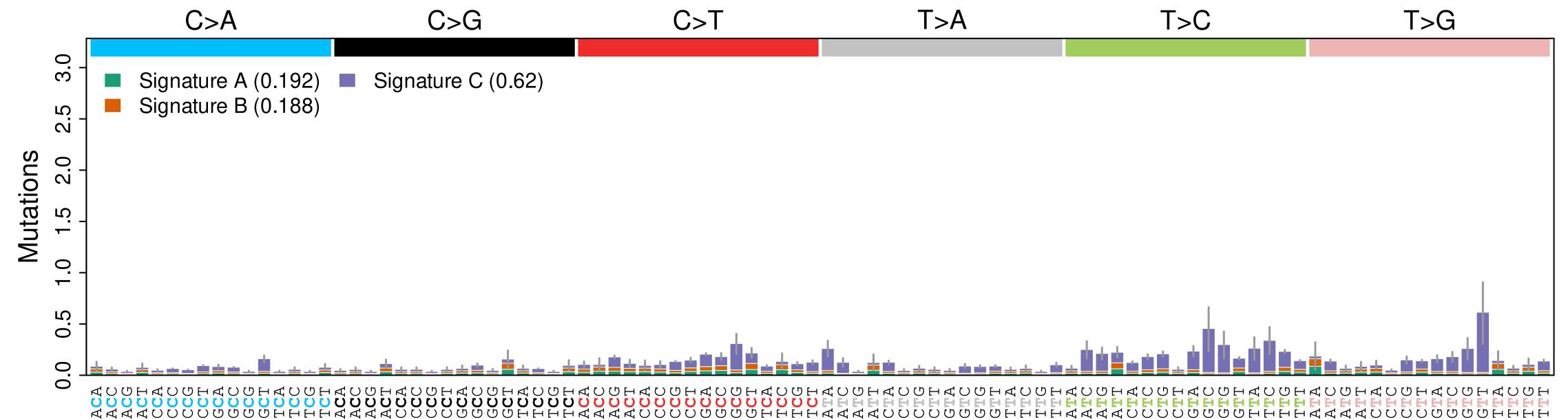


Reconstructed spectrum (cosine similarity = 0.257)

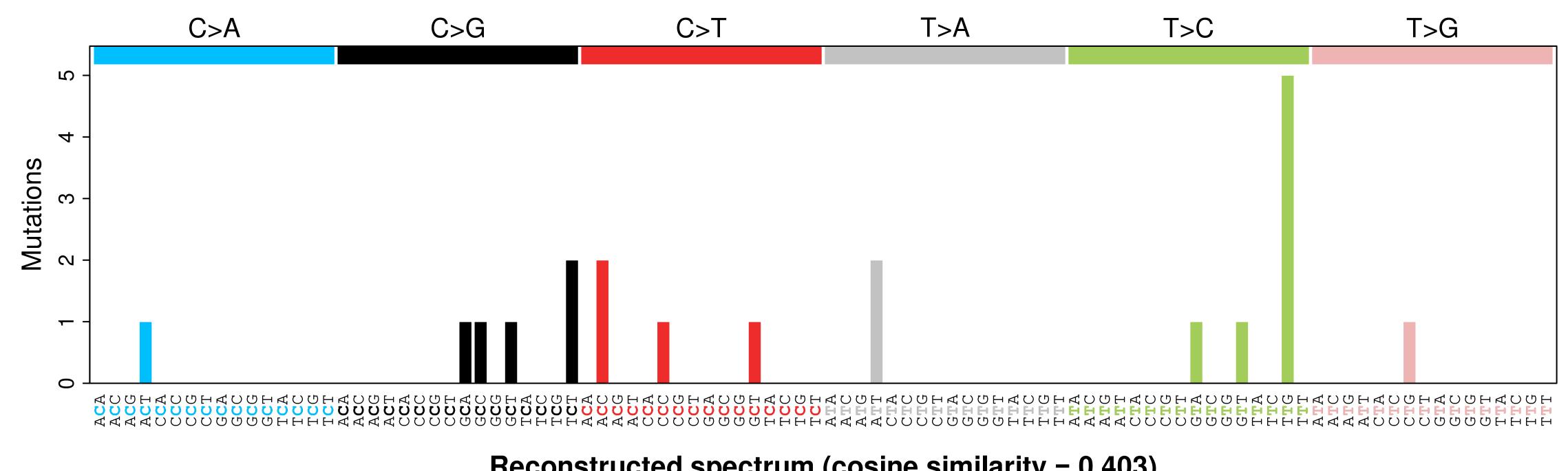


CATD0725a (12 mutations)

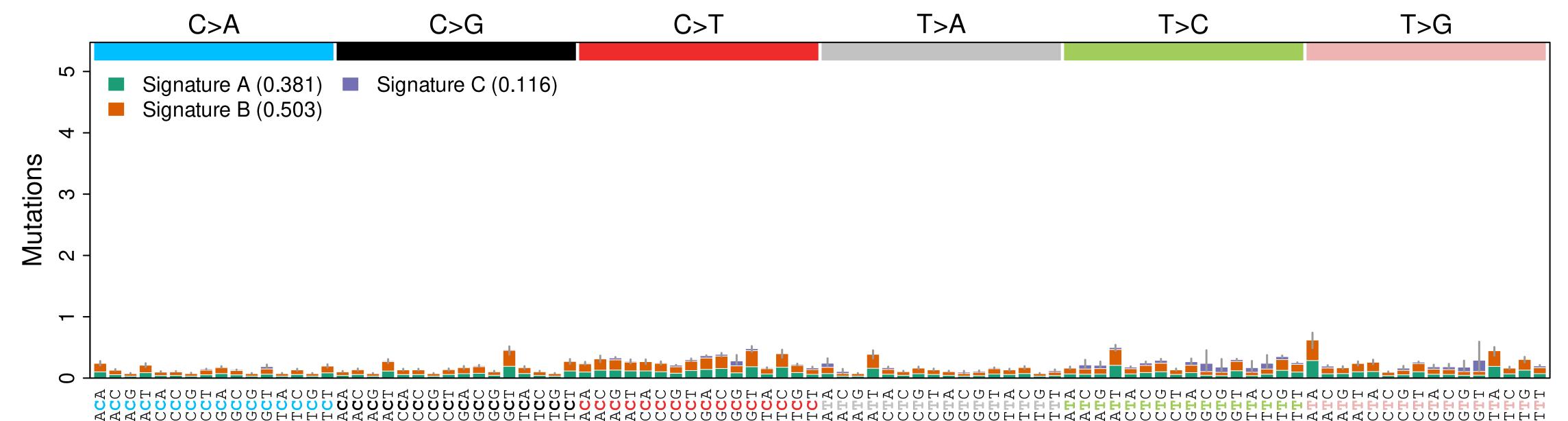




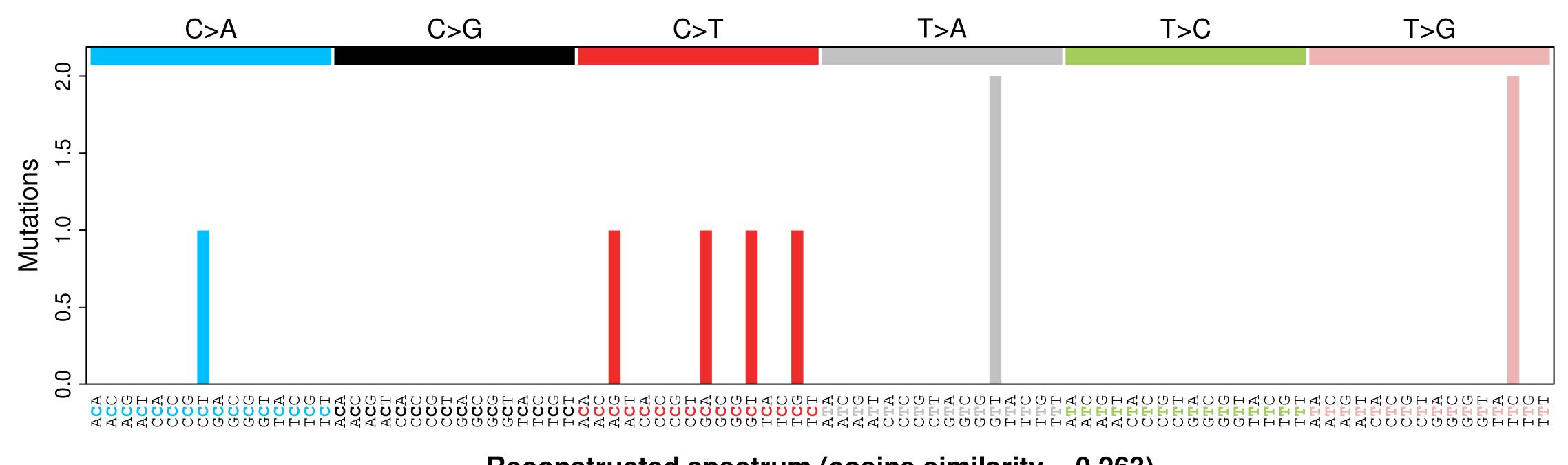
CATD0577a (20 mutations)

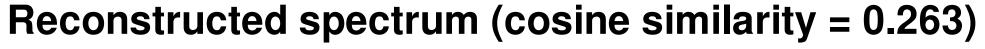


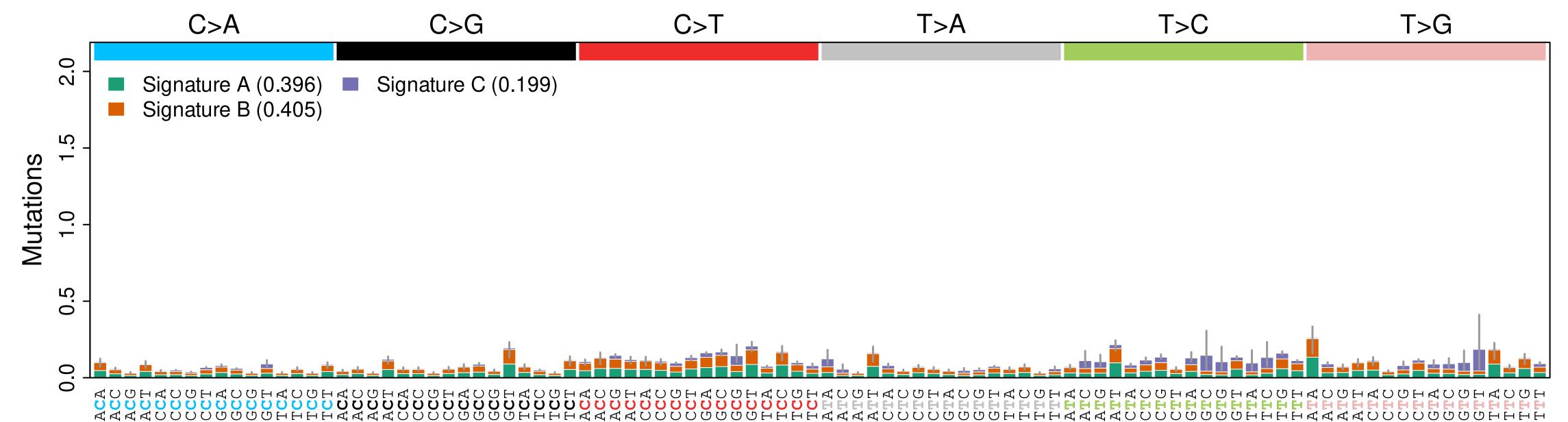
Reconstructed spectrum (cosine similarity = 0.403)



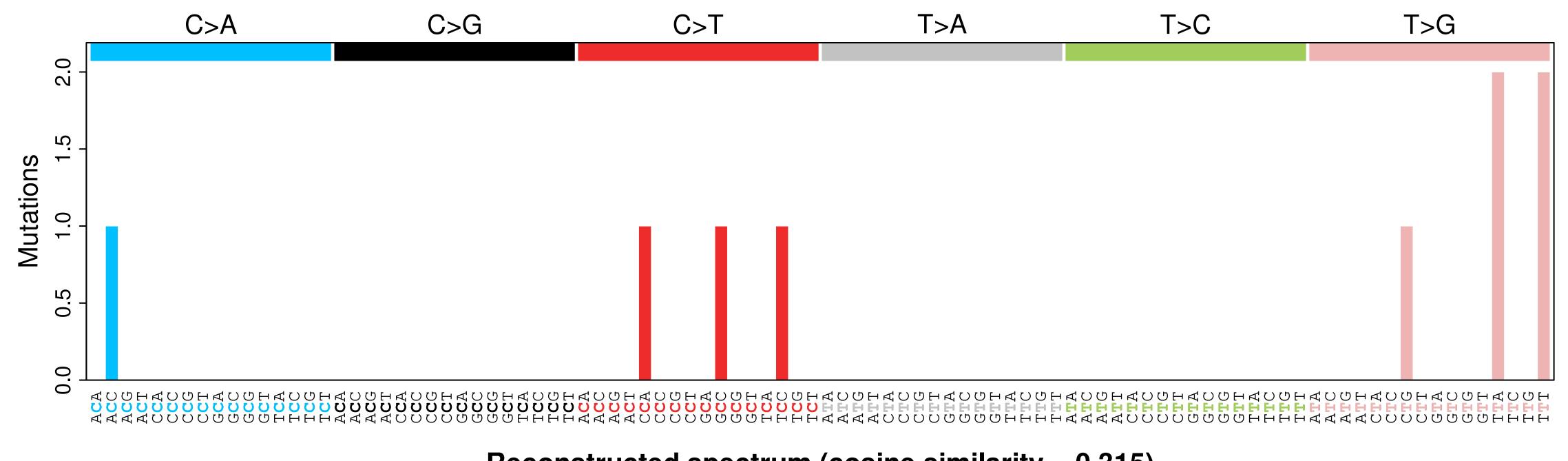
CATD0606a (9 mutations)

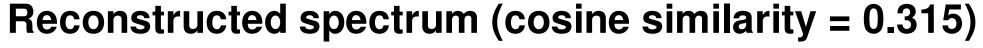


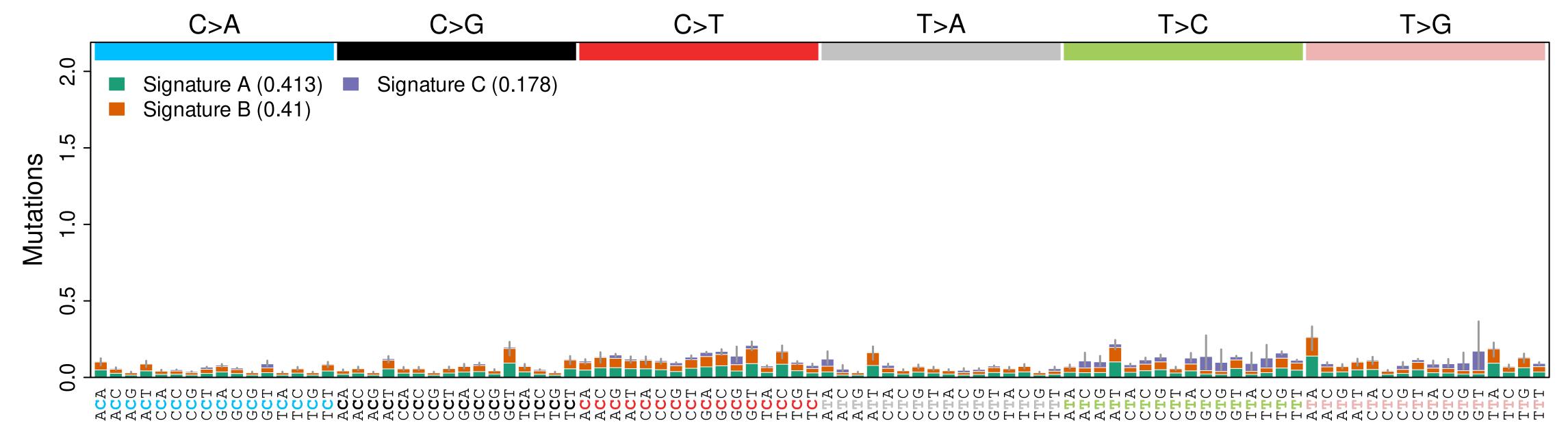




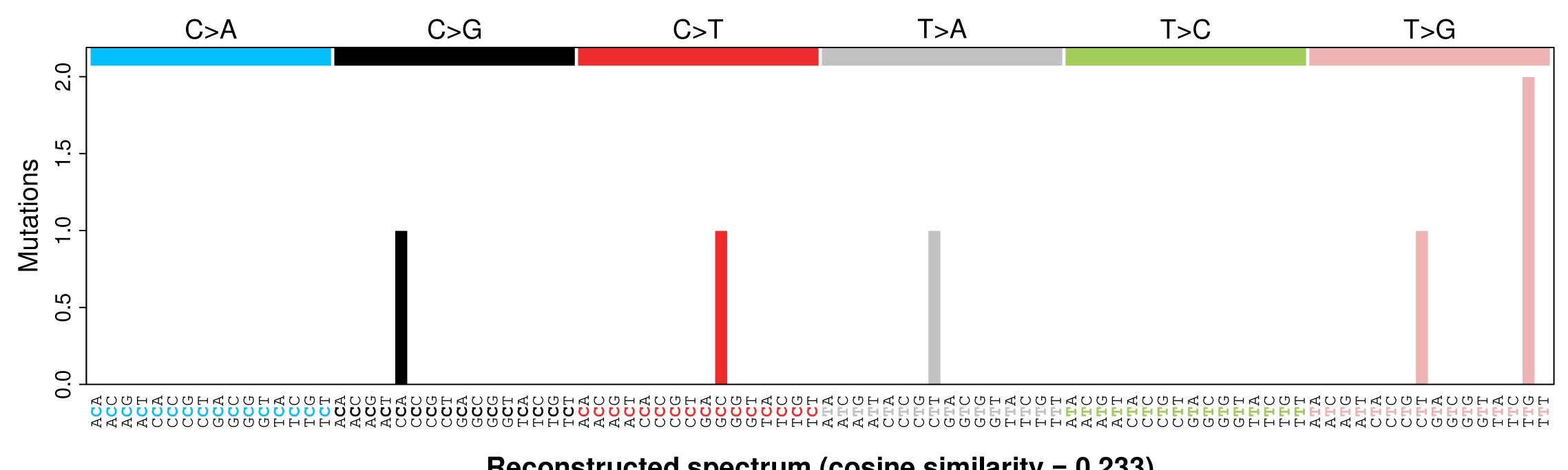
CATD0588a (9 mutations)



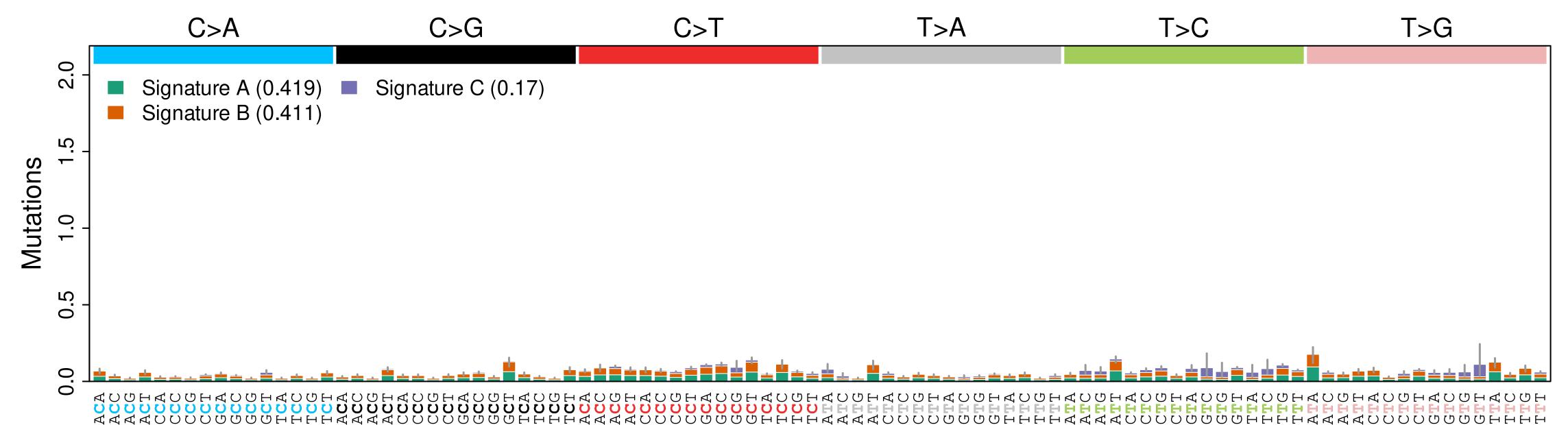




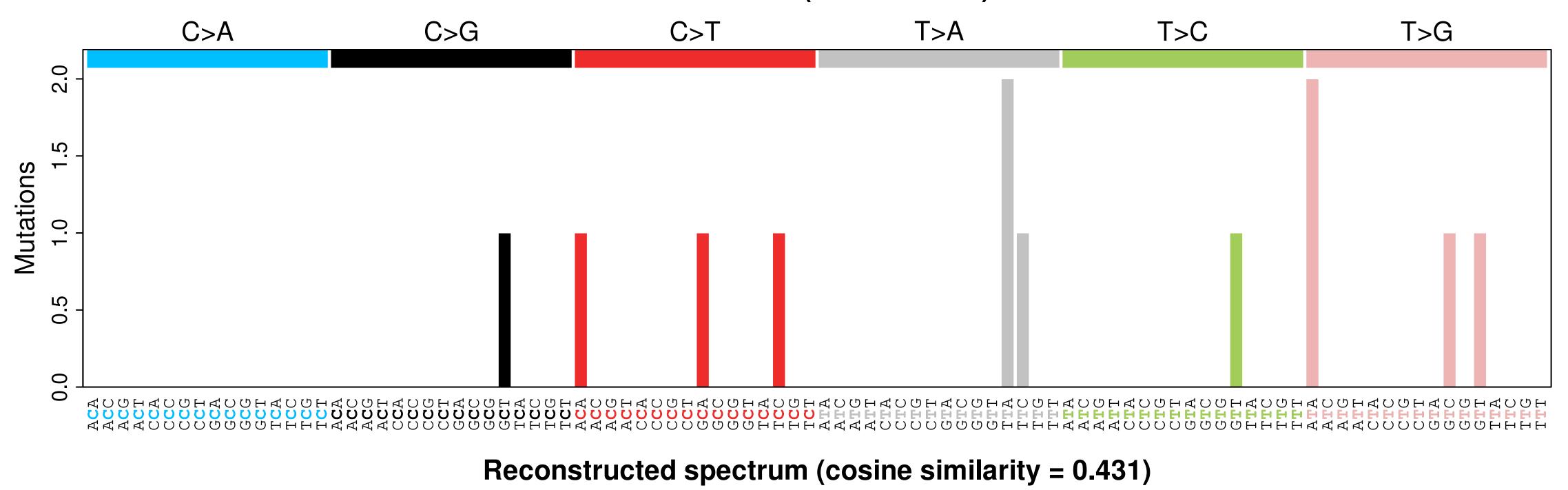
CATD0572a (6 mutations)

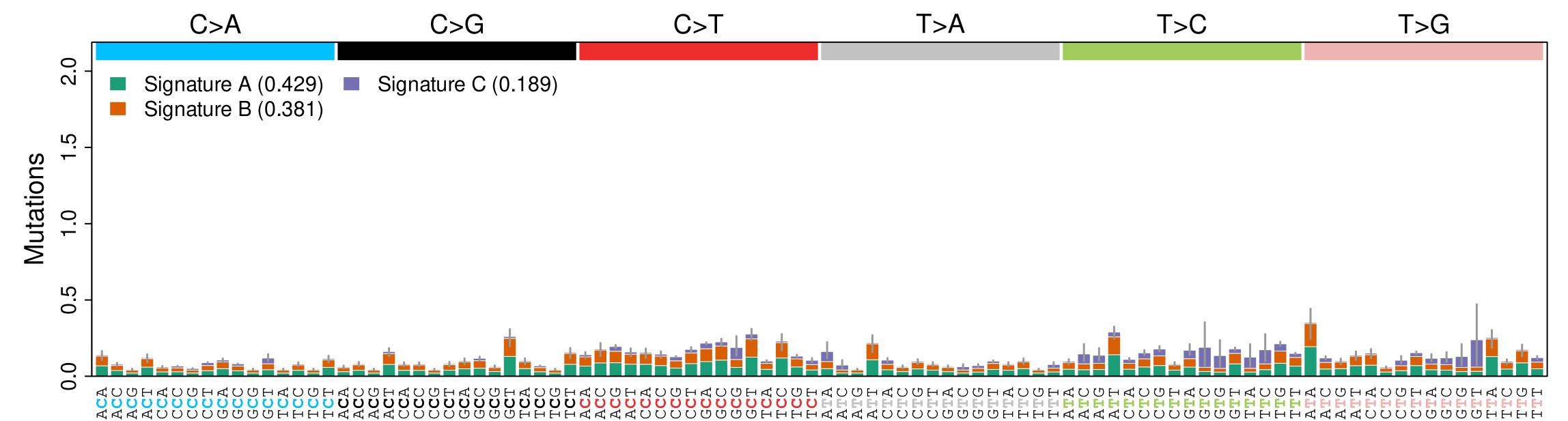


Reconstructed spectrum (cosine similarity = 0.233)

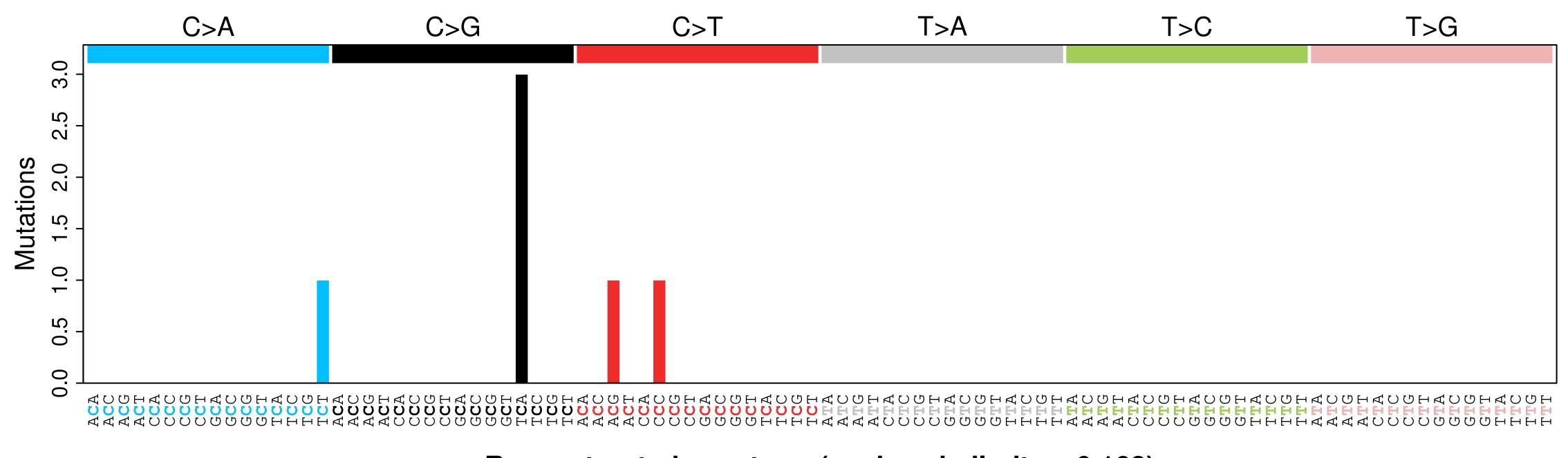


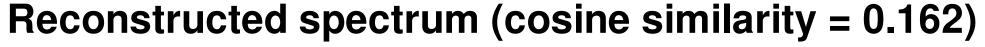
CATD0573a (12 mutations)

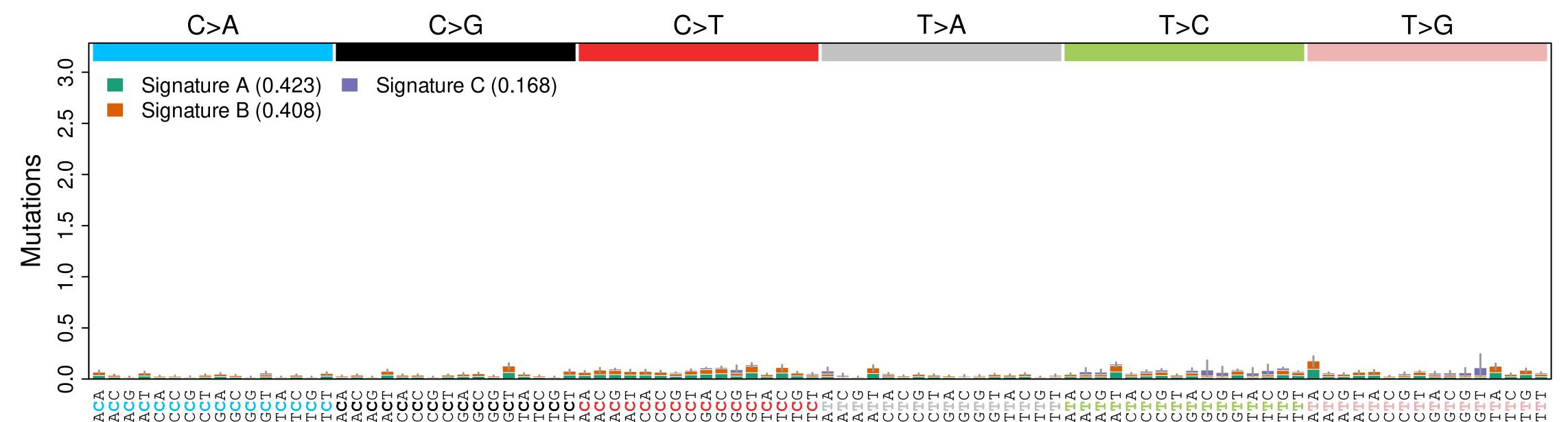




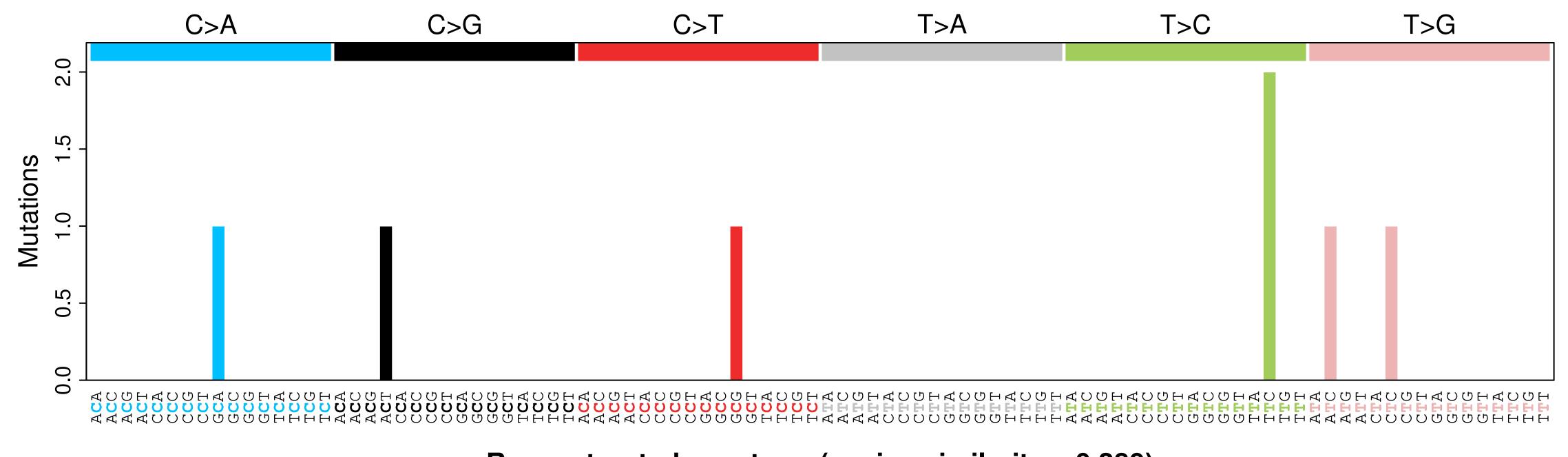
CATD0592a (6 mutations)

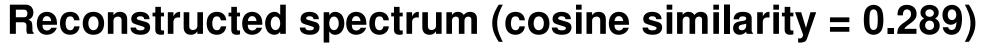


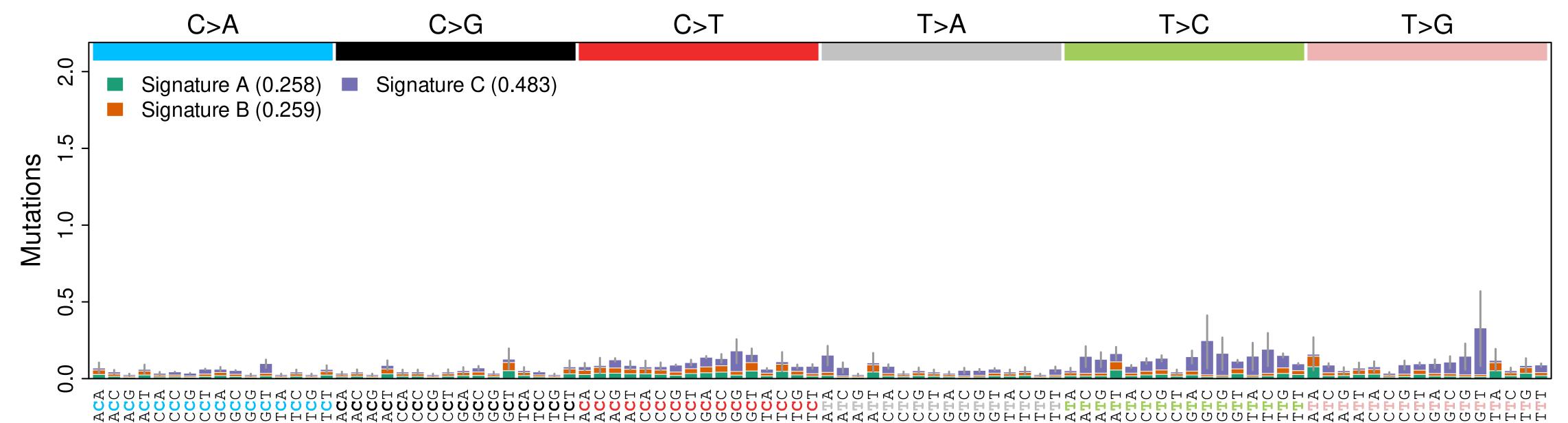




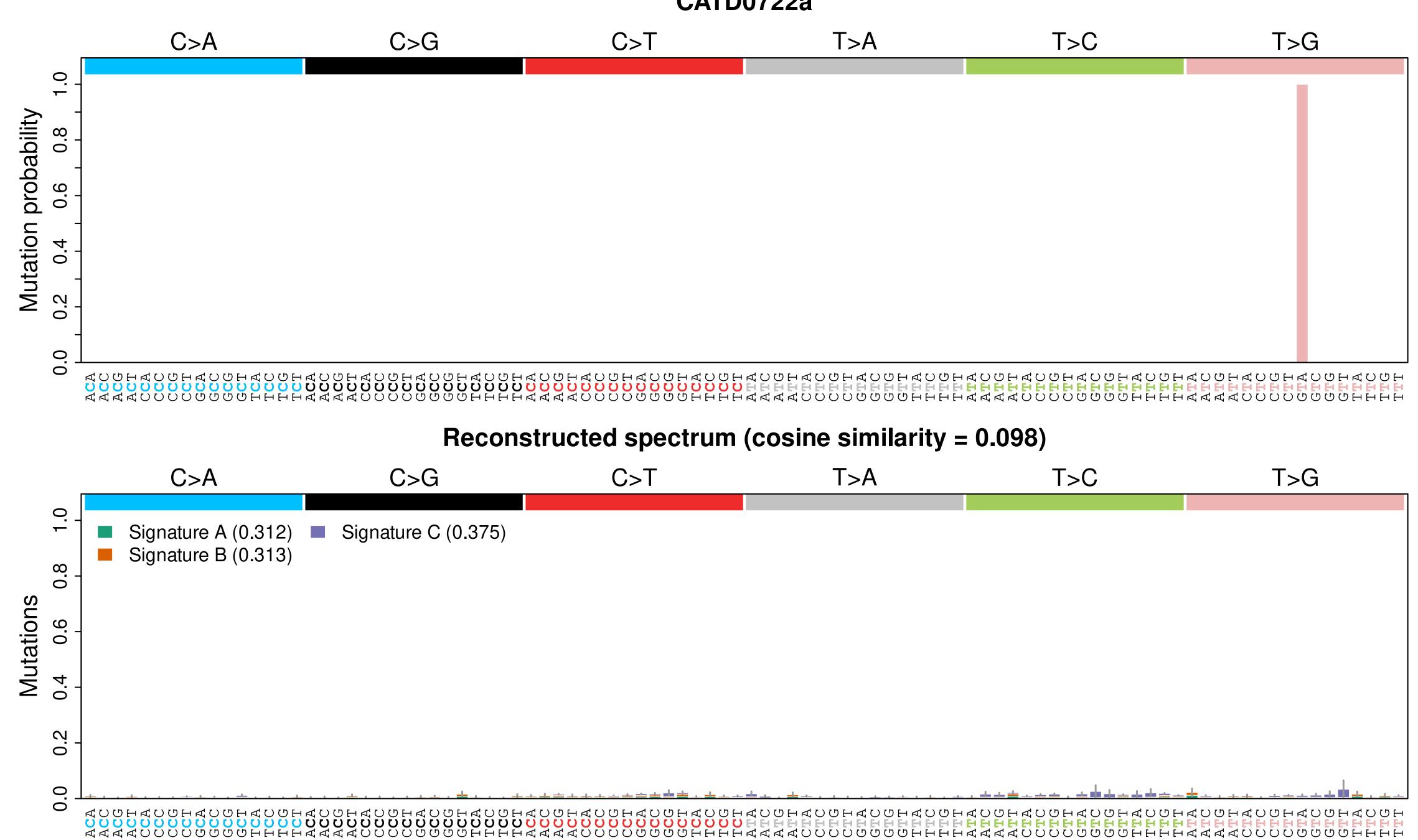
CATD0603a (7 mutations)



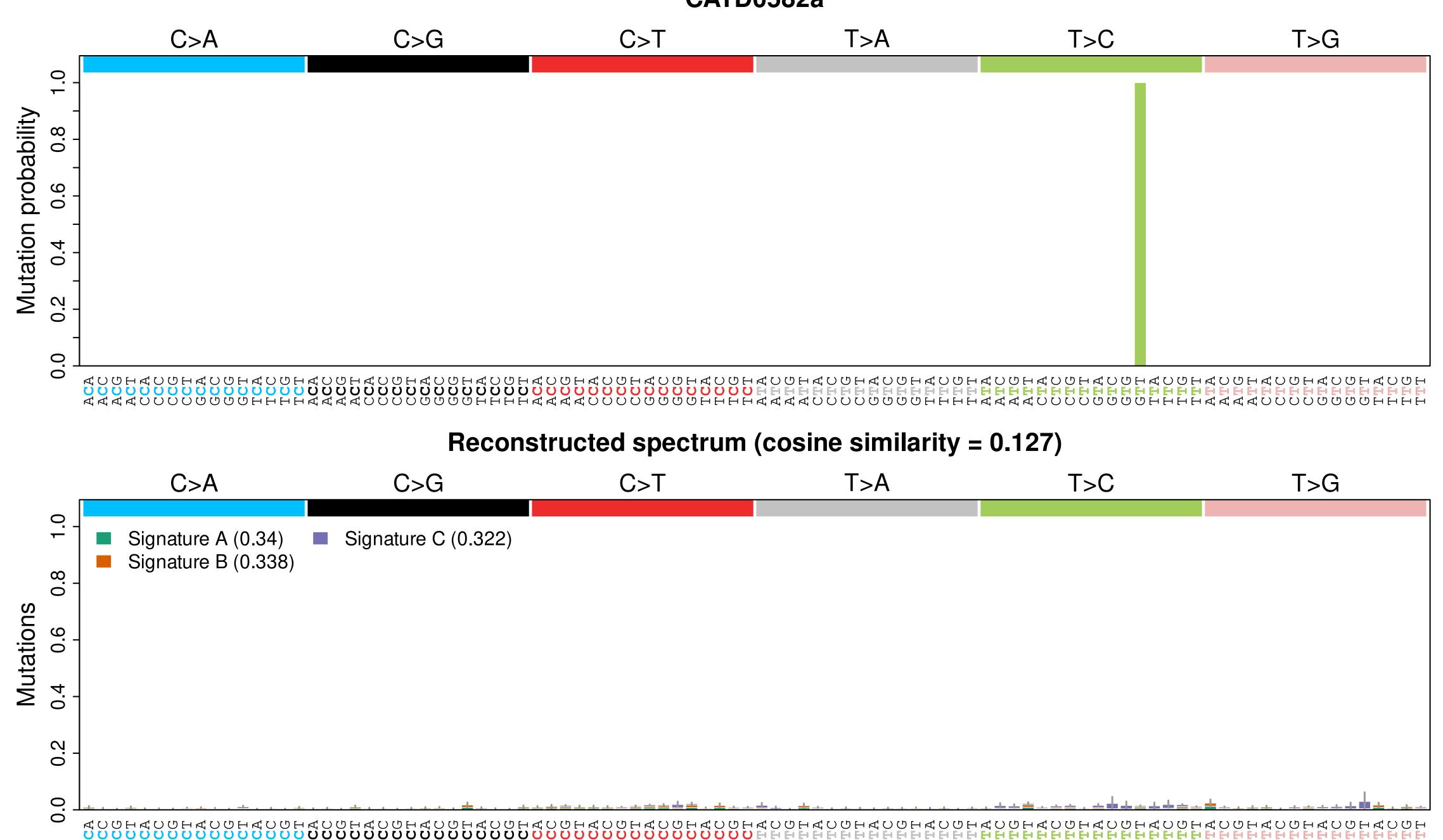




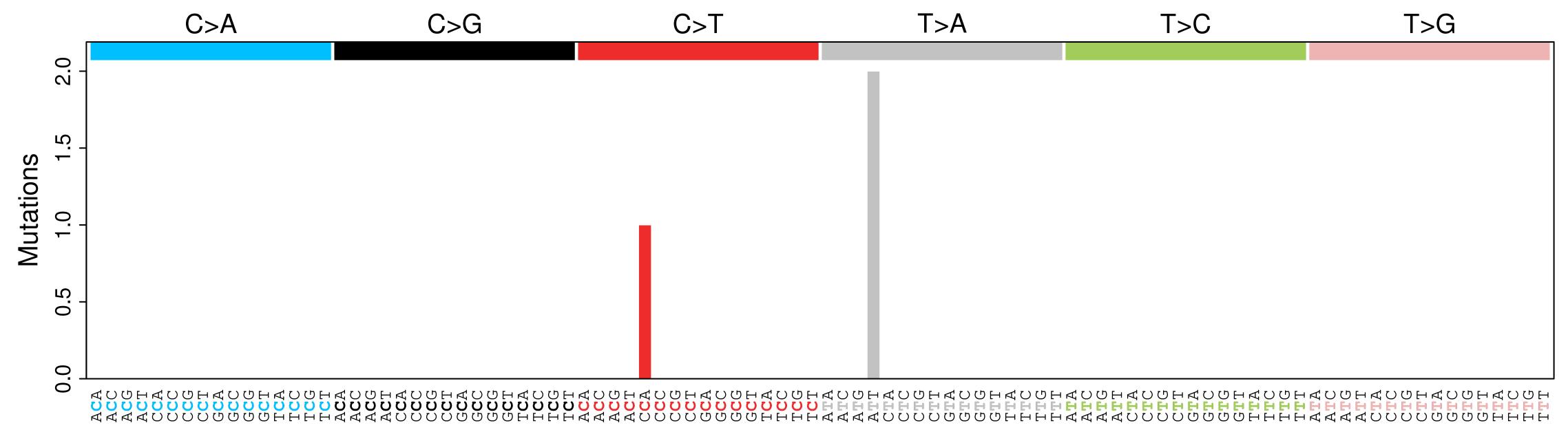




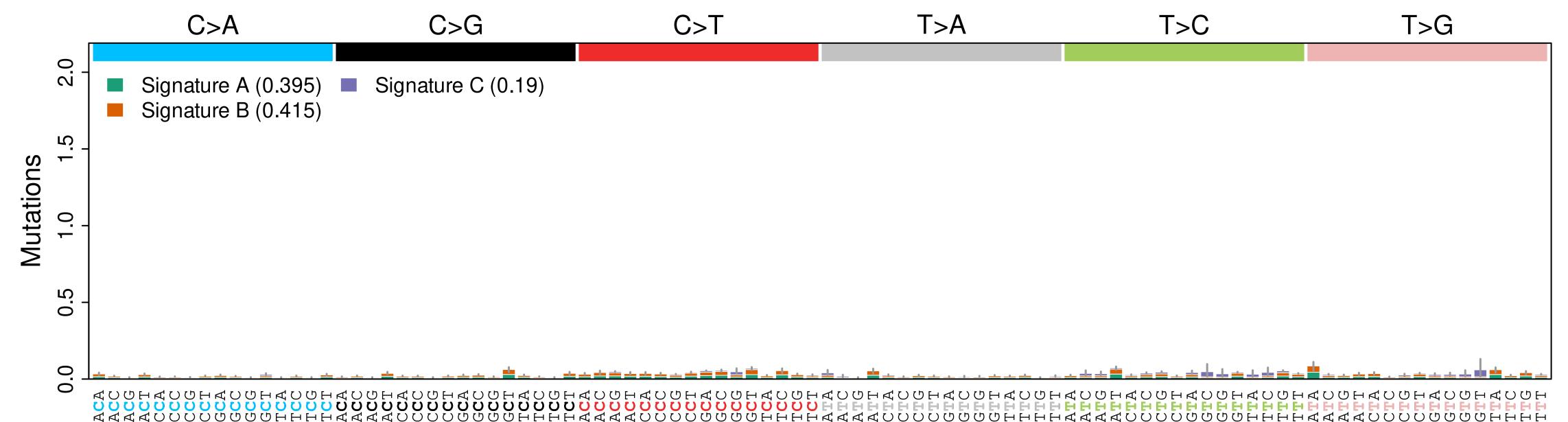




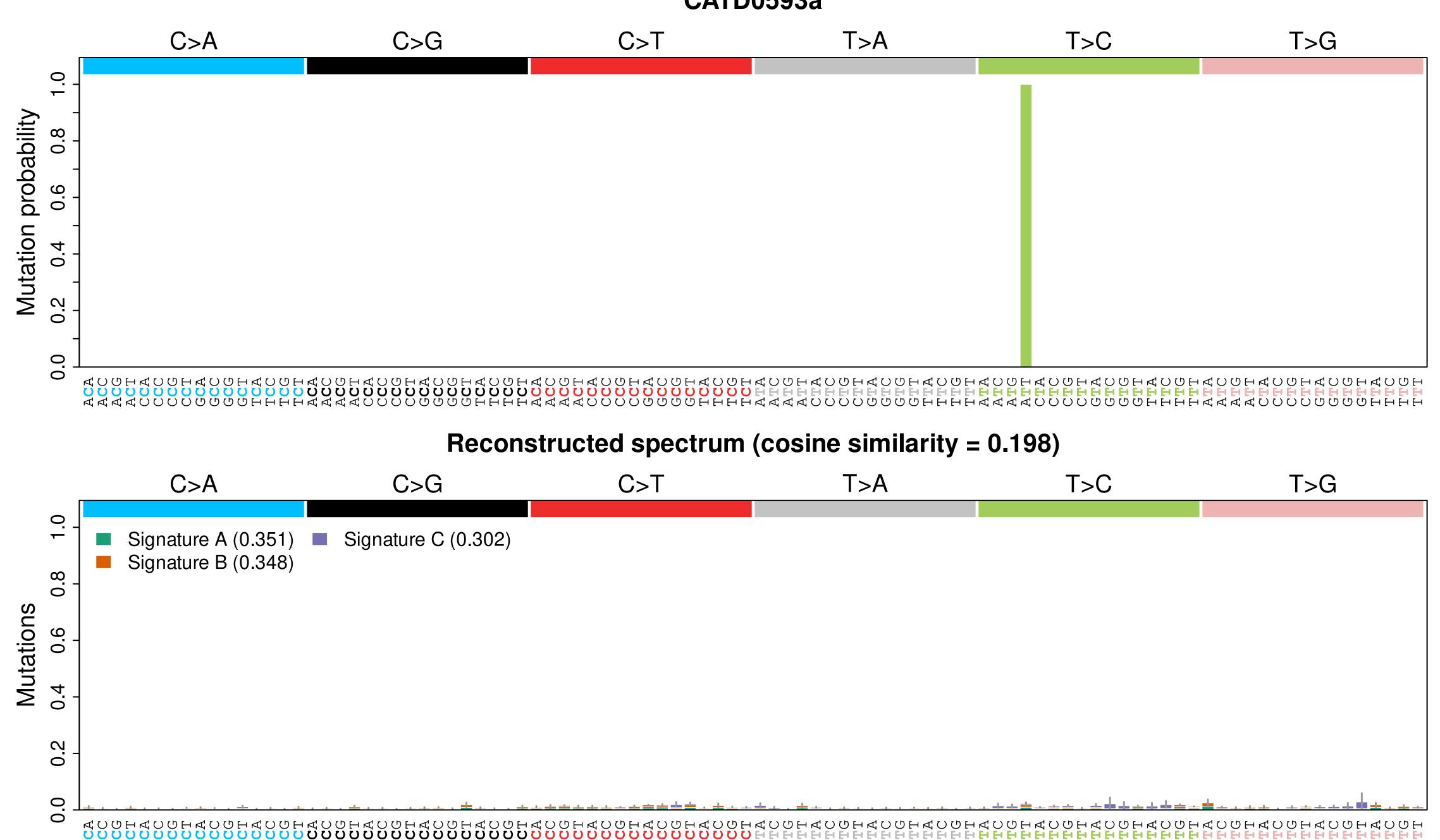
CATD0585a (3 mutations)



Reconstructed spectrum (cosine similarity = 0.195)



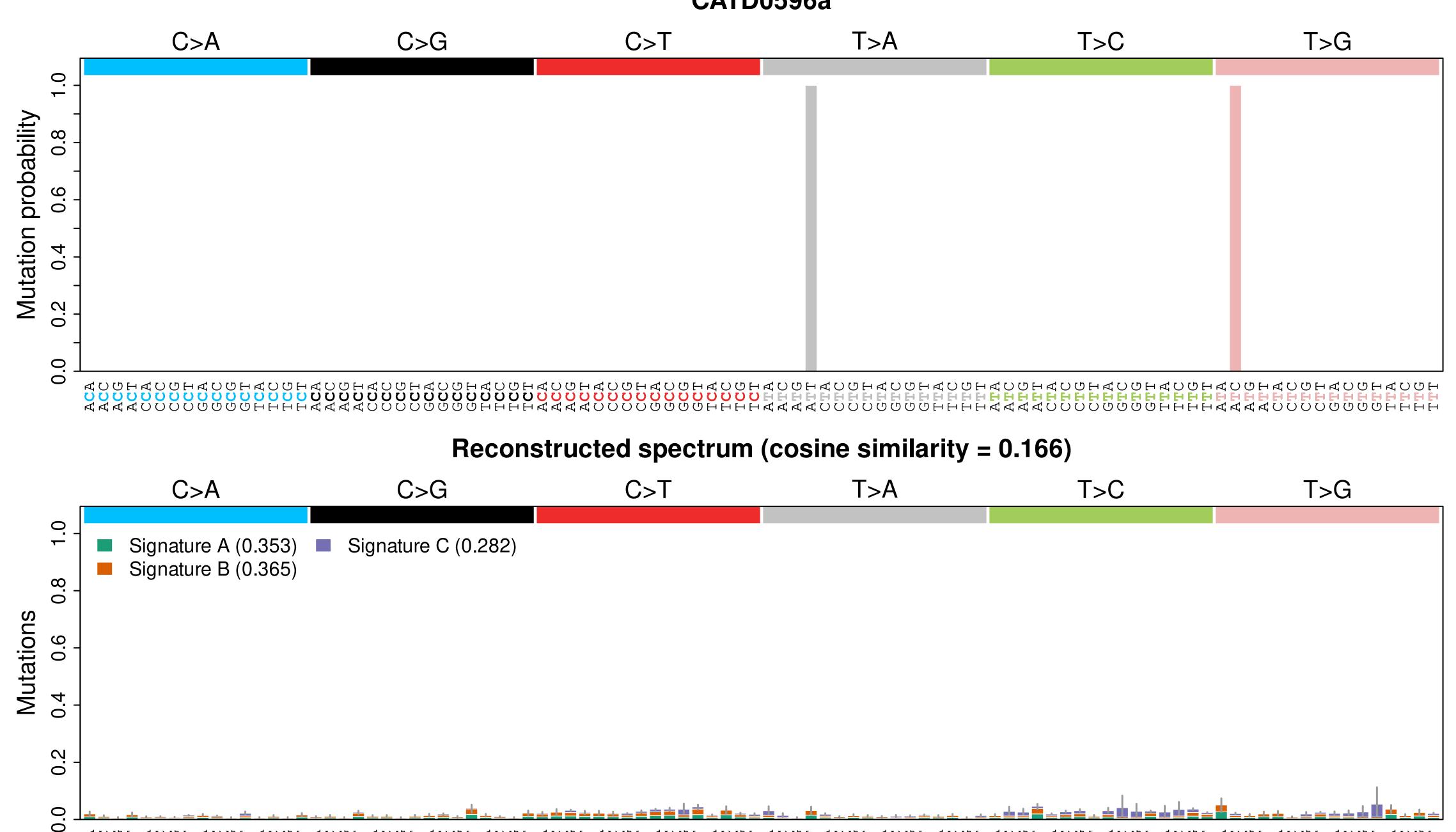




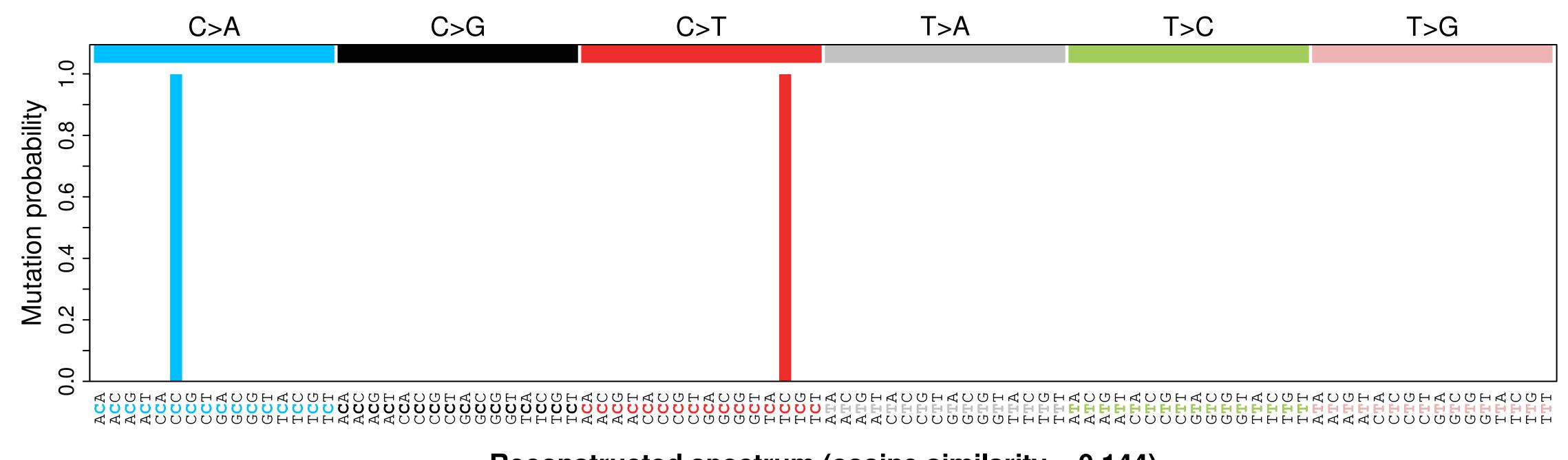
CATD0581a (3 mutations) T>G C>A C>G C>T T>A T>C 2.0 5 Mutations 0.0 **Reconstructed spectrum (cosine similarity = 0.149)** C>A C>G T>A T>C T>G C>T 2.0 Signature A (0.389) Signature C (0.198) Signature B (0.413) 5 Mutations 5 0 0

CATD0736a (4 mutations) T>G C>A C>G C>T T>A T>C 2.0 1.5 Mutations 0.5 0.0 **Reconstructed spectrum (cosine similarity = 0.285)** C>A C>G T>A T>C T>G C>T 2.0 Signature A (0.316) Signature C (0.379) Signature B (0.304) 5 Mutations 2 0

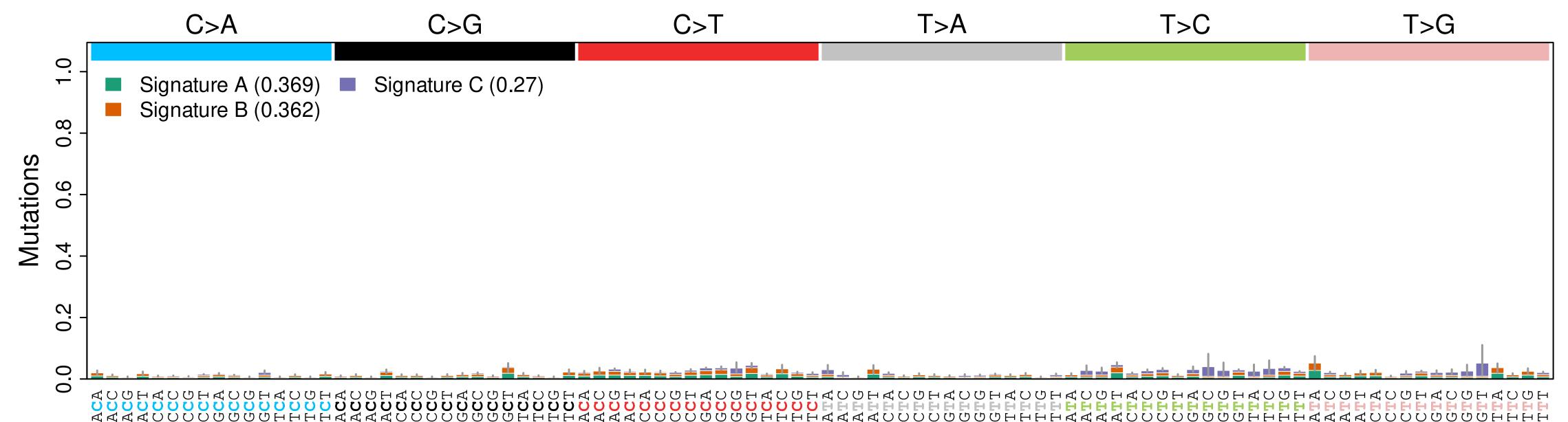




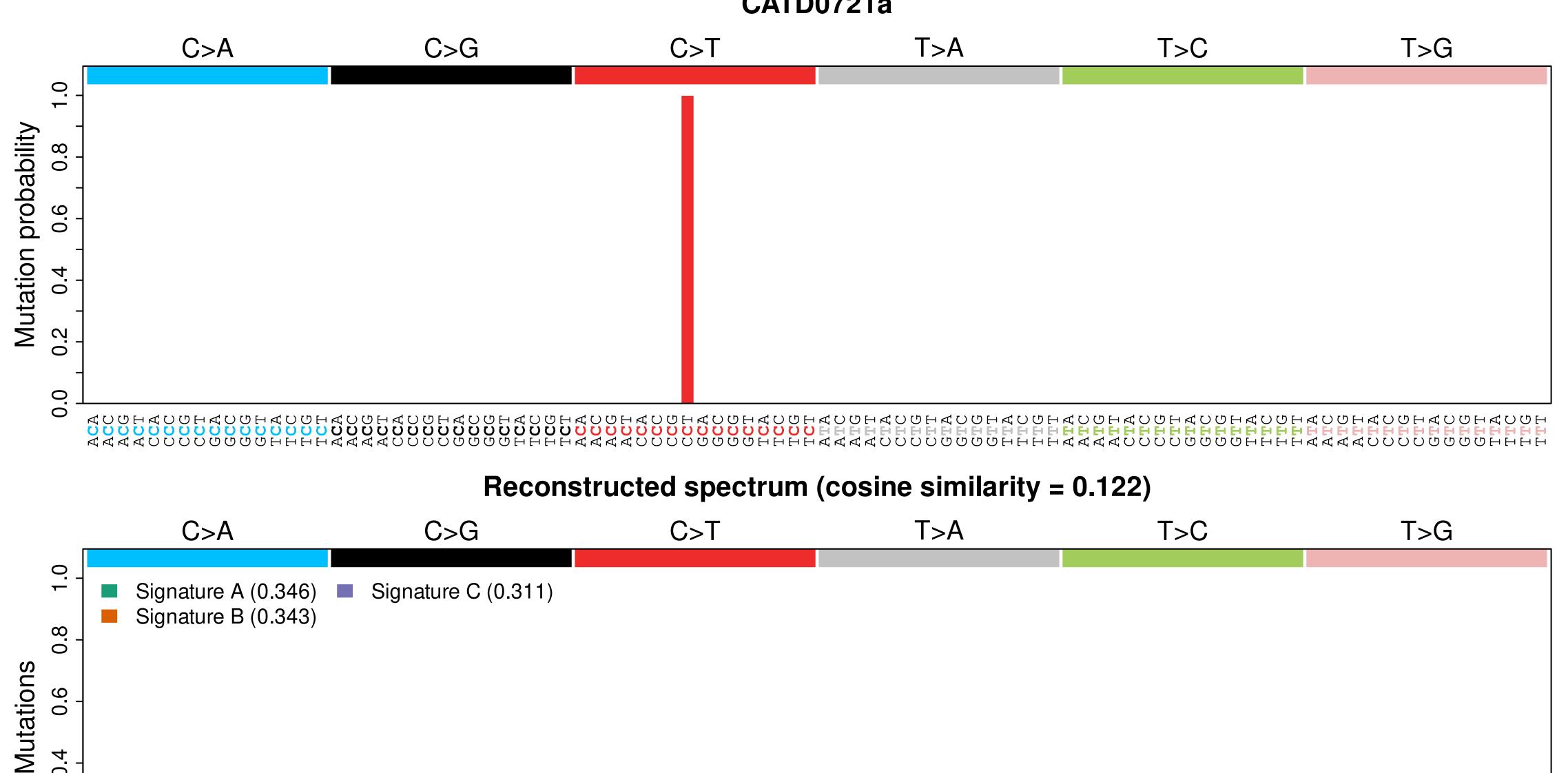




Reconstructed spectrum (cosine similarity = 0.144)

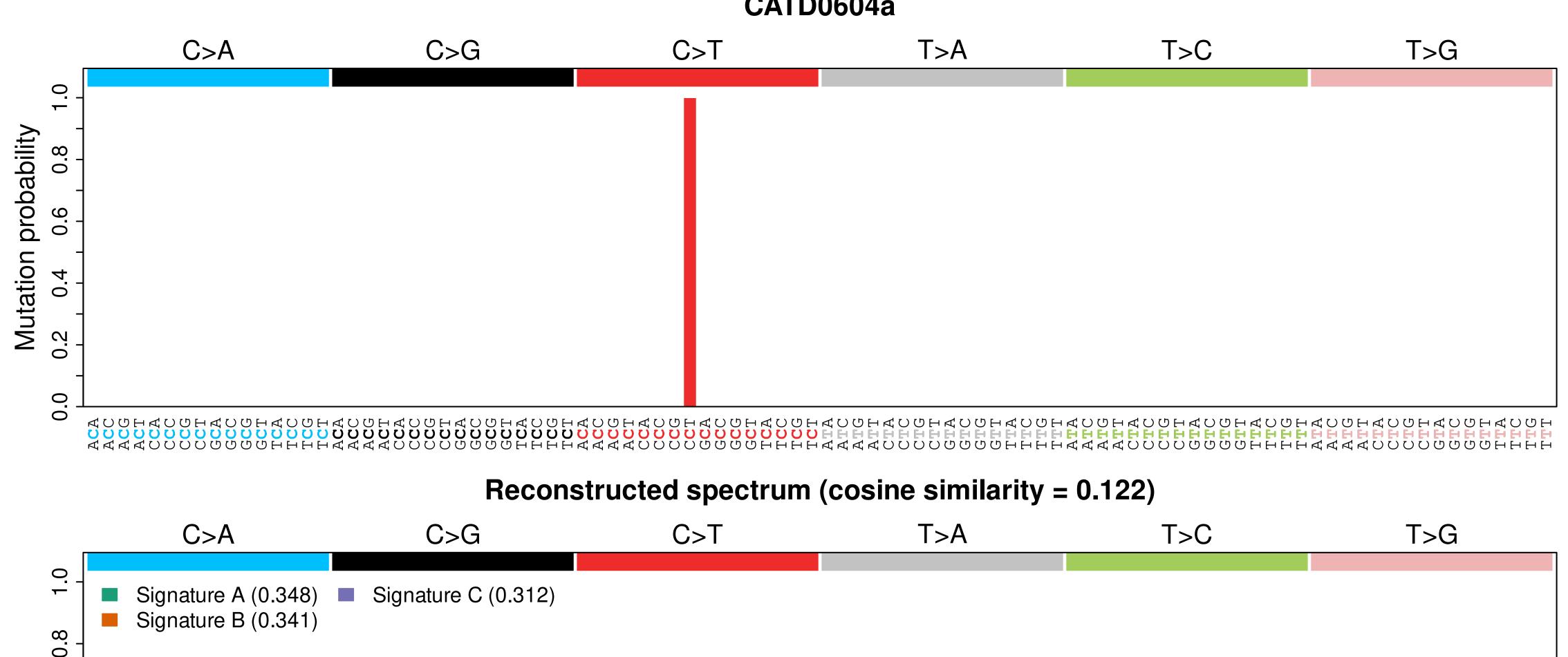


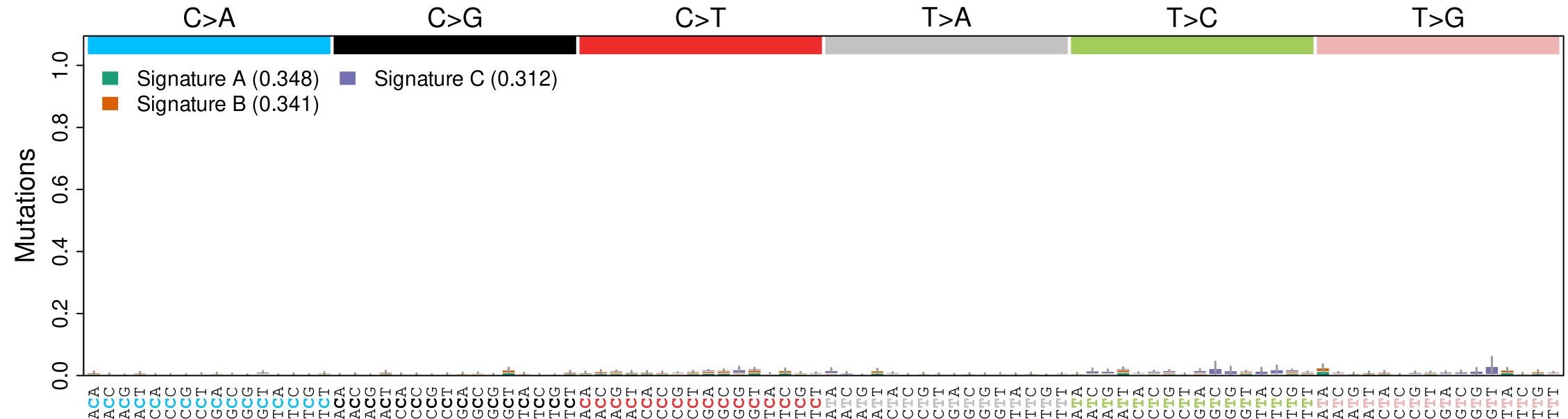




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CATD0720a

