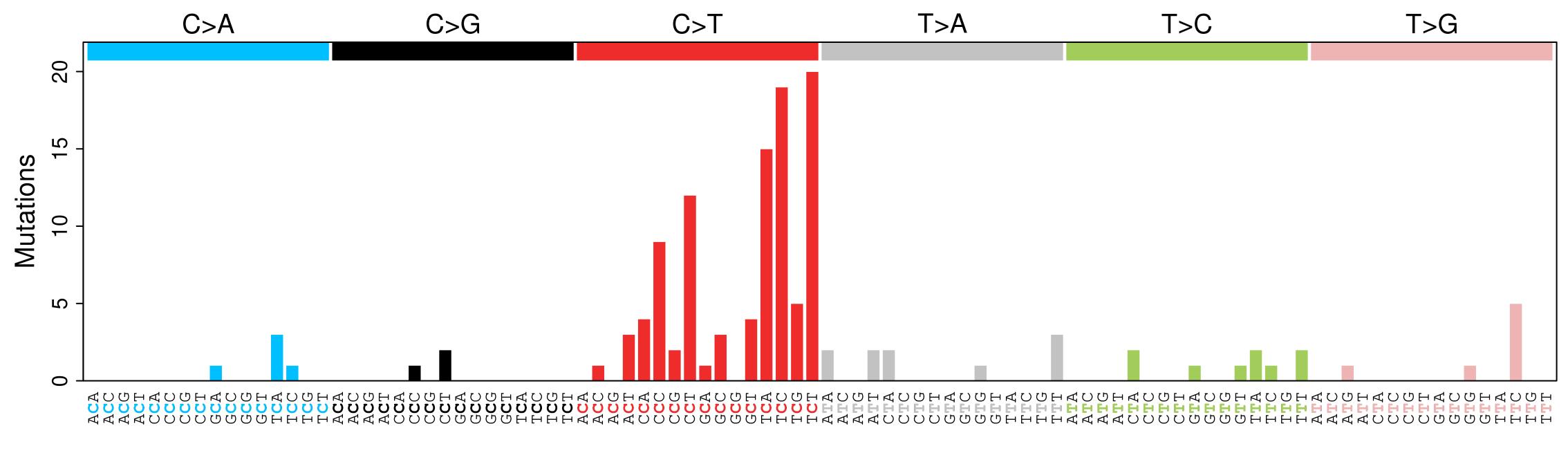CATD0653a (132 mutations)

Reconstructed spectrum (cosine similarity = 0.909)

SBS1 (0.006)
SBS2 (0.098)
SBS3 (0.008)
SBS4 (0.006)
SBS5 (0.009)
SBS6 (0.008)
SBS7a (0.117)
SBS7b (0.075)
SBS7c (0.015)
SBS7d (0.01)
SBS8 (0.007)
SBS9 (0.007)
SBS36 (0.007)
SBS37 (0.009)
SBS38 (0.005)
SBS39 (0.008)
SBS40a (0.01)
SBS40b (0.011)
SBS40c (0.009)
SBS41 (0.009)
SBS42 (0.01)
SBS43 (0.006)
SBS44 (0.007)
SBS45 (0.005)