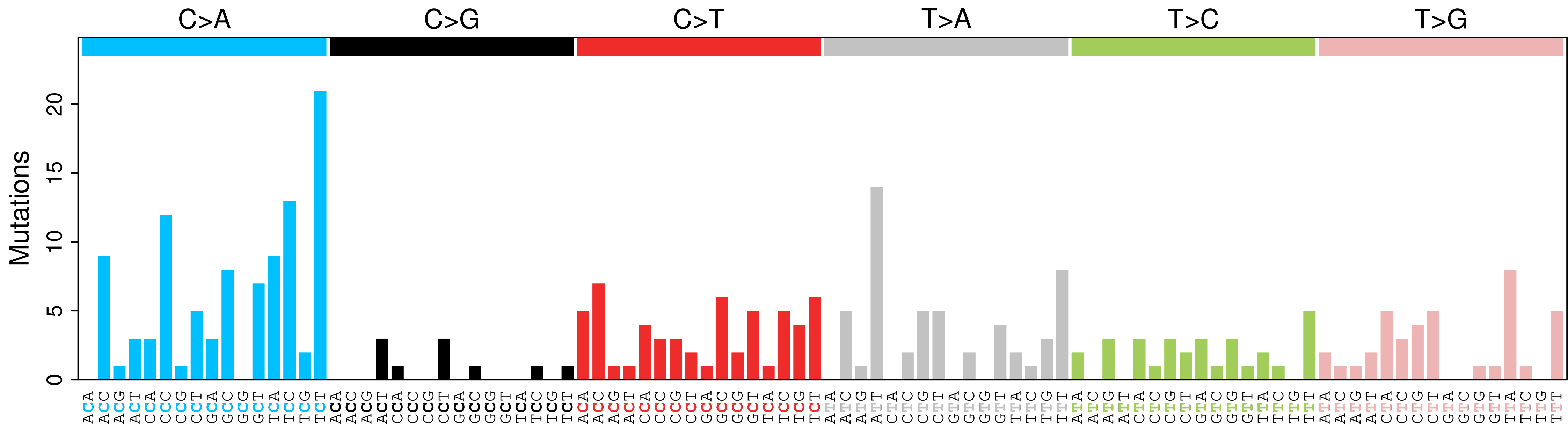CATD186a (284 mutations)

Reconstructed spectrum (cosine similarity = 0.852)

SBS1 (0.006)   SBS36 (0.018)
SBS2 (0.005)   SBS37 (0.007)
SBS3 (0.013)   SBS38 (0.007)
SBS4 (0.021)   SBS39 (0.008)
SBS5 (0.009)   SBS40a (0.011)
SBS6 (0.007)   SBS40b (0.014)
SBS7a (0.007)   SBS40c (0.011)
SBS7b (0.009)   SBS41 (0.012)
SBS7c (0.03)   SBS42 (0.026)
SBS7d (0.006)   SBS43 (0.005)
SBS8 (0.033)   SBS44 (0.012)
SBS9 (0.03)   SBS45 (0.015)