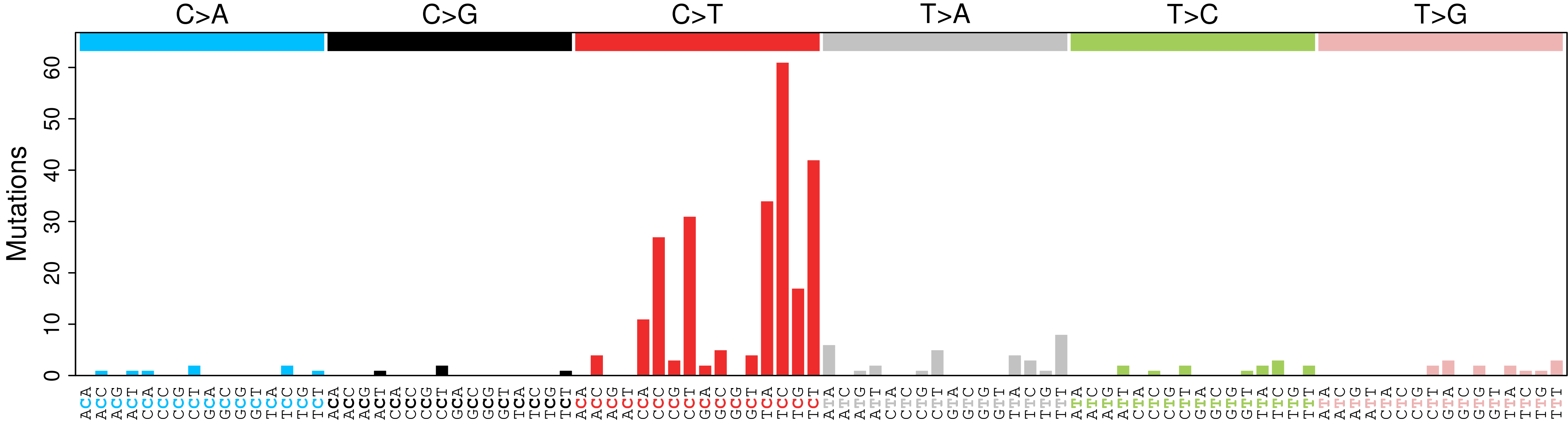**CATD0123a (311 mutations)**

Reconstructed spectrum (cosine similarity = 0.966)

SBS1 (0.003)
SBS2 (0.073)
SBS3 (0.005)
SBS4 (0.004)
SBS5 (0.005)
SBS6 (0.004)
SBS7a (0.259)
SBS7b (0.184)
SBS7c (0.027)
SBS7d (0.005)
SBS8 (0.005)
SBS9 (0.006)
SBS36 (0.004)
SBS37 (0.005)
SBS38 (0.004)
SBS39 (0.004)
SBS40a (0.006)
SBS40b (0.006)
SBS40c (0.005)
SBS41 (0.007)
SBS42 (0.006)
SBS43 (0.006)
SBS44 (0.004)
SBS45 (0.004)