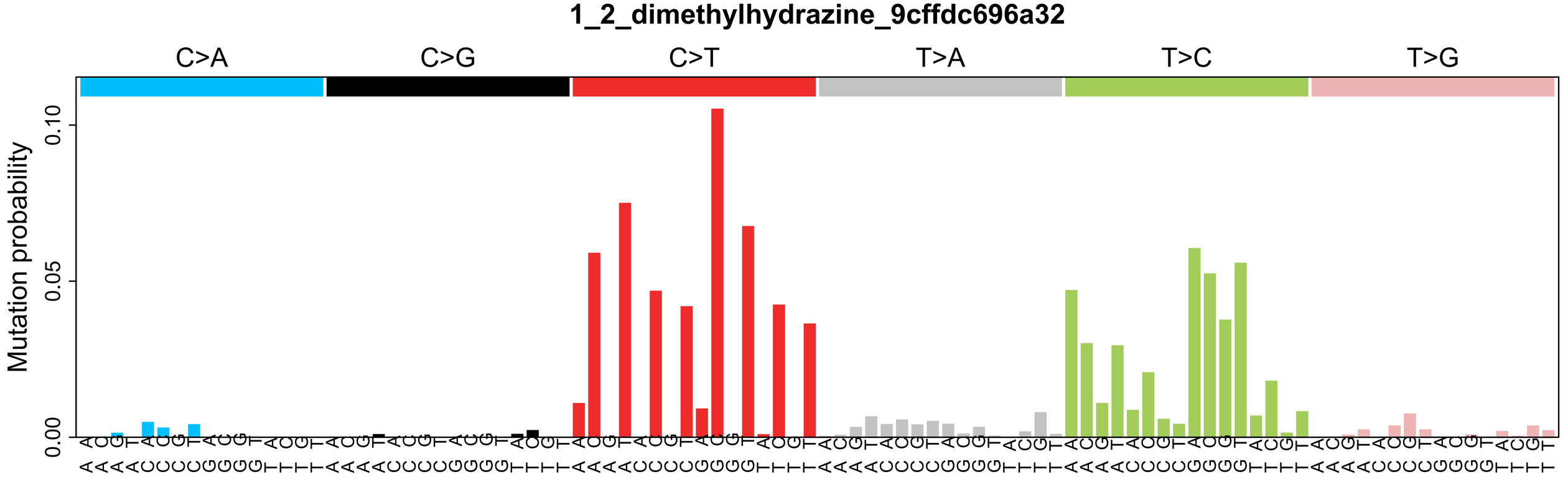
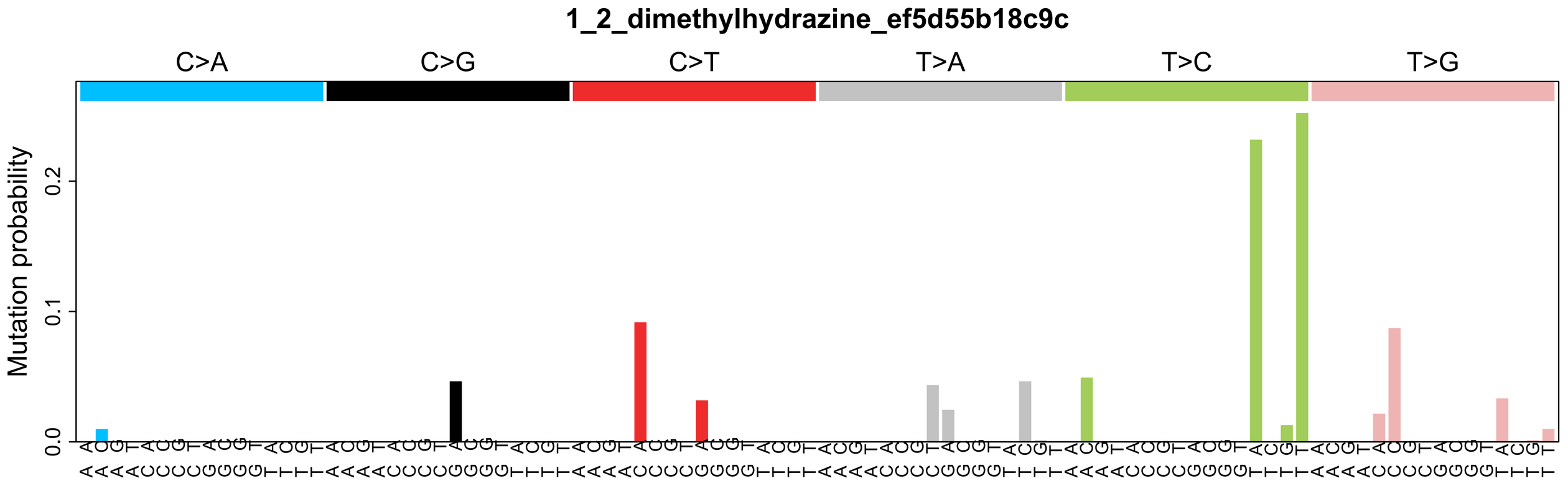
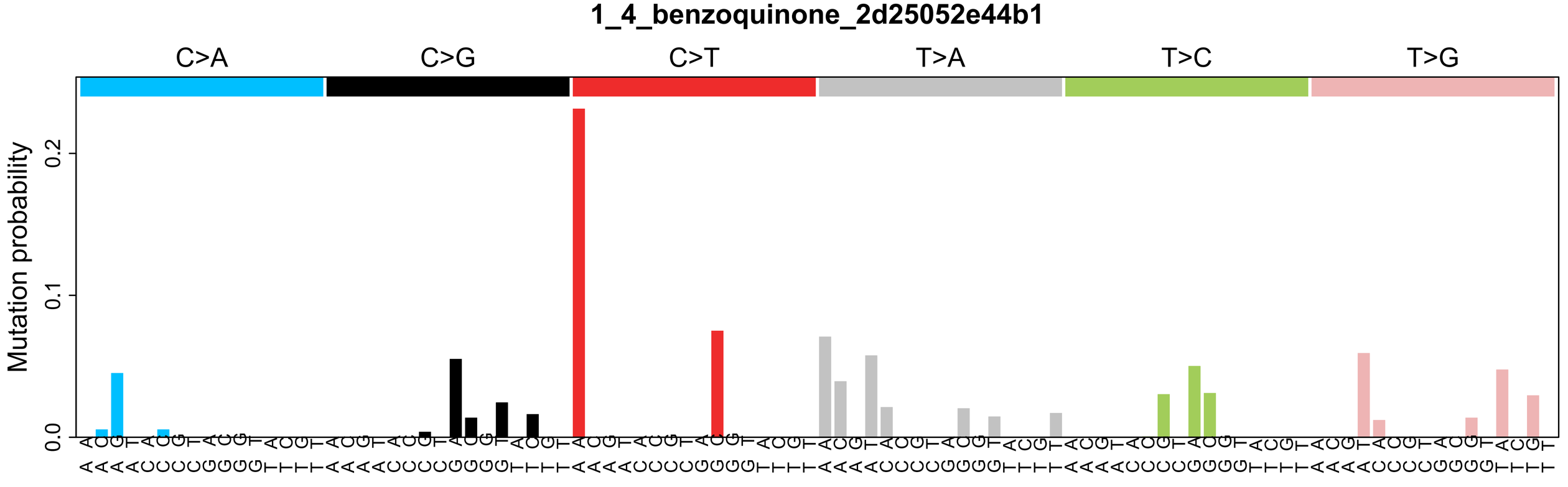
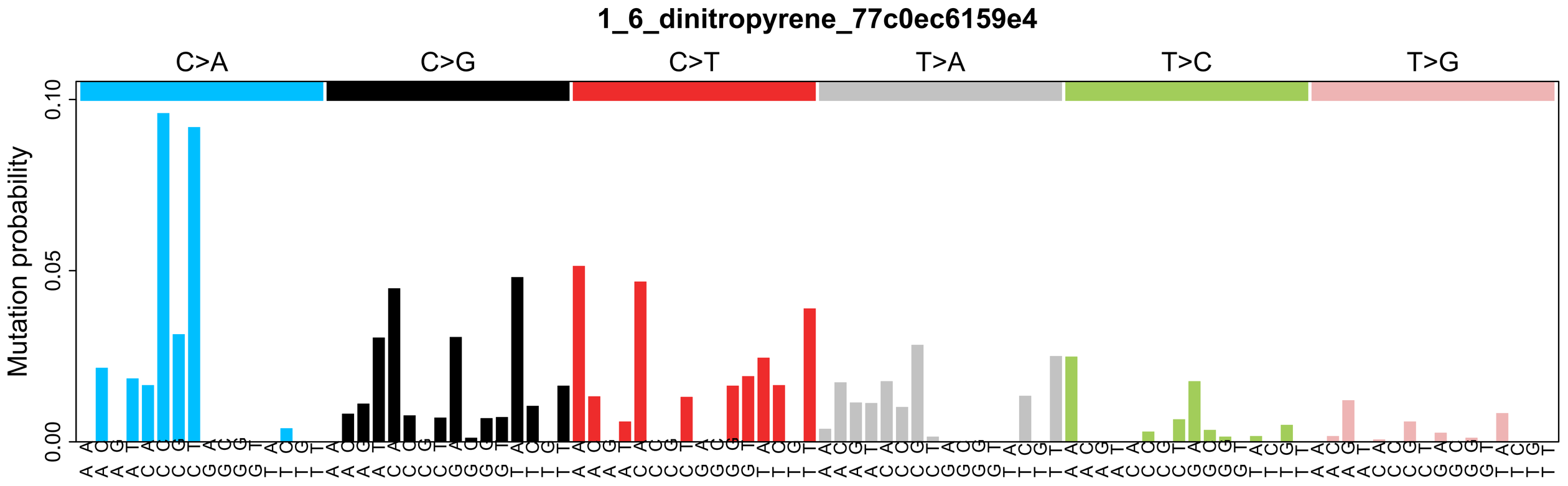
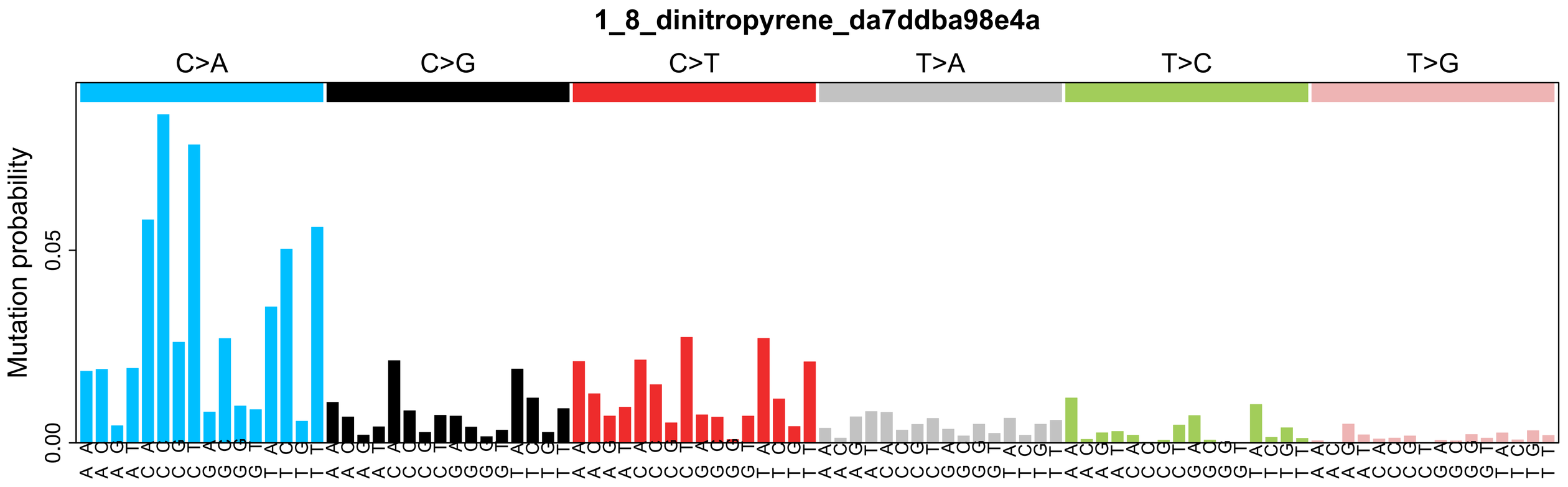
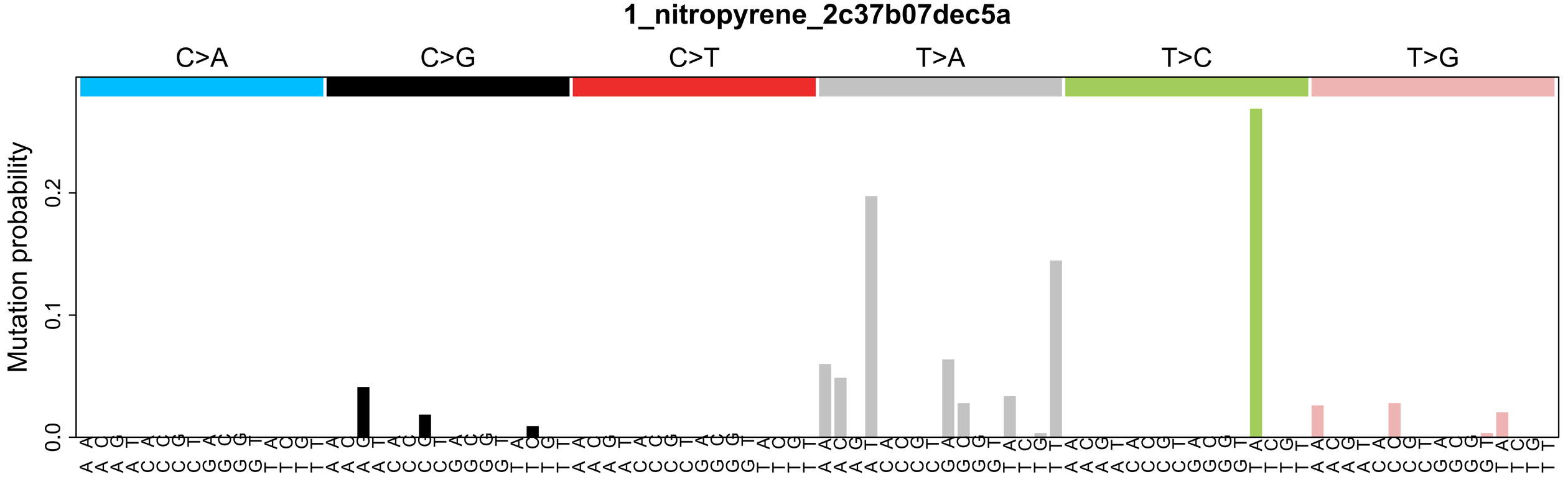

1_2_dimethylhydrazine_9cffdc696a32
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10

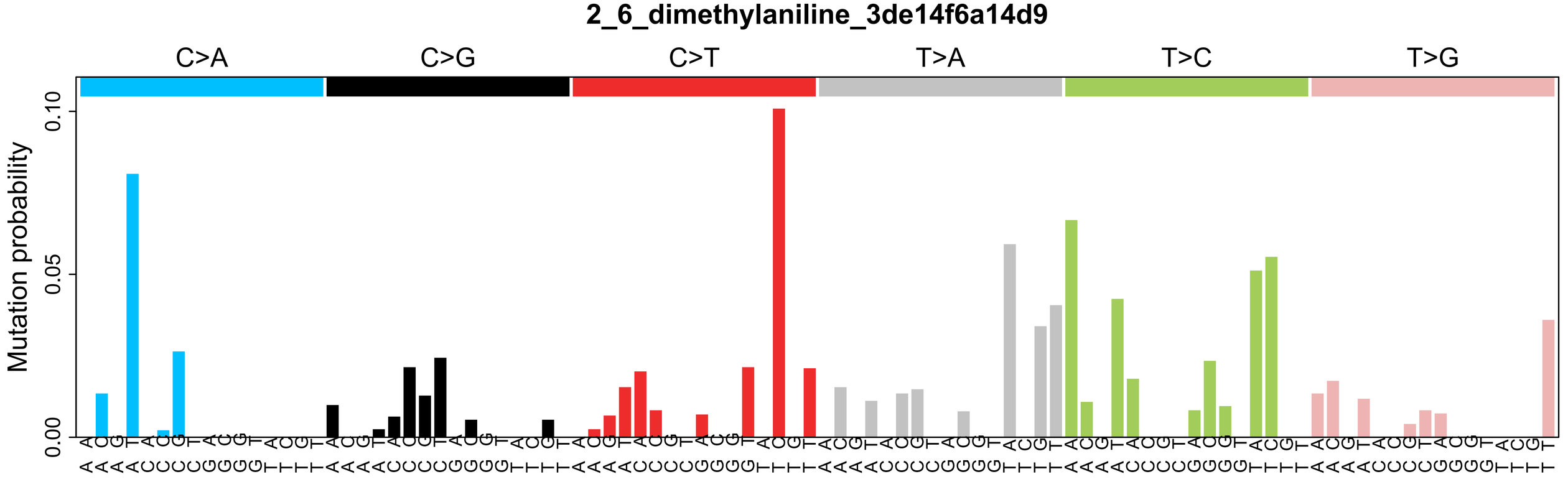

1_2_dimethylhydrazine_ef5d55b18c9c
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2

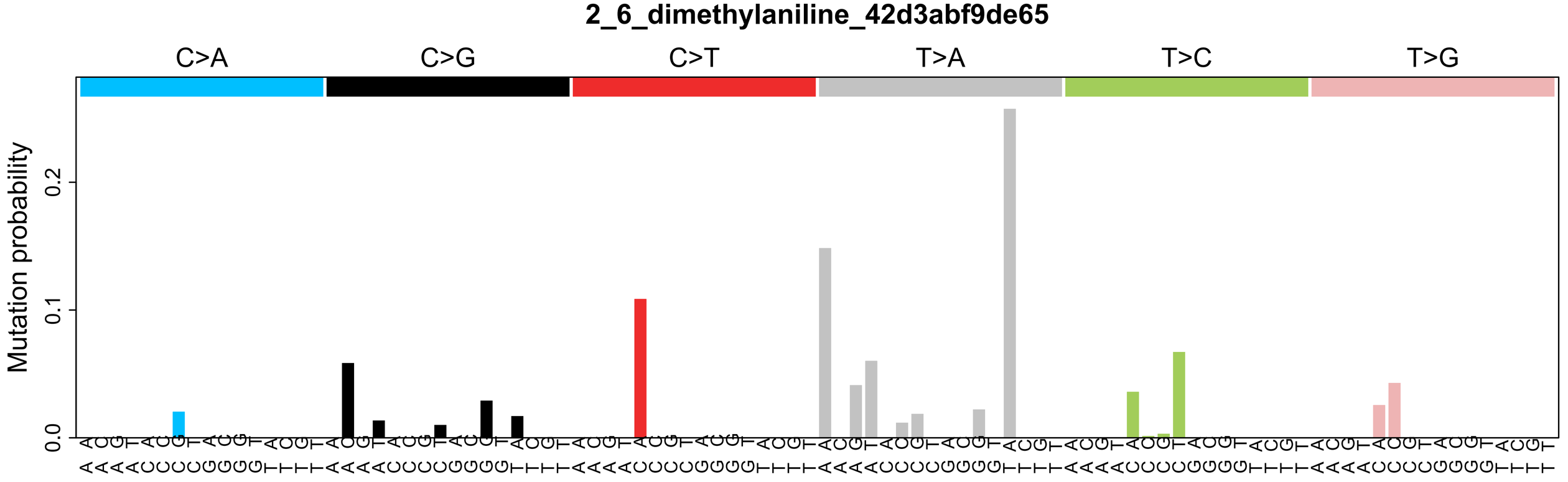

1_4_benzoquinone_2d25052e44b1
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2

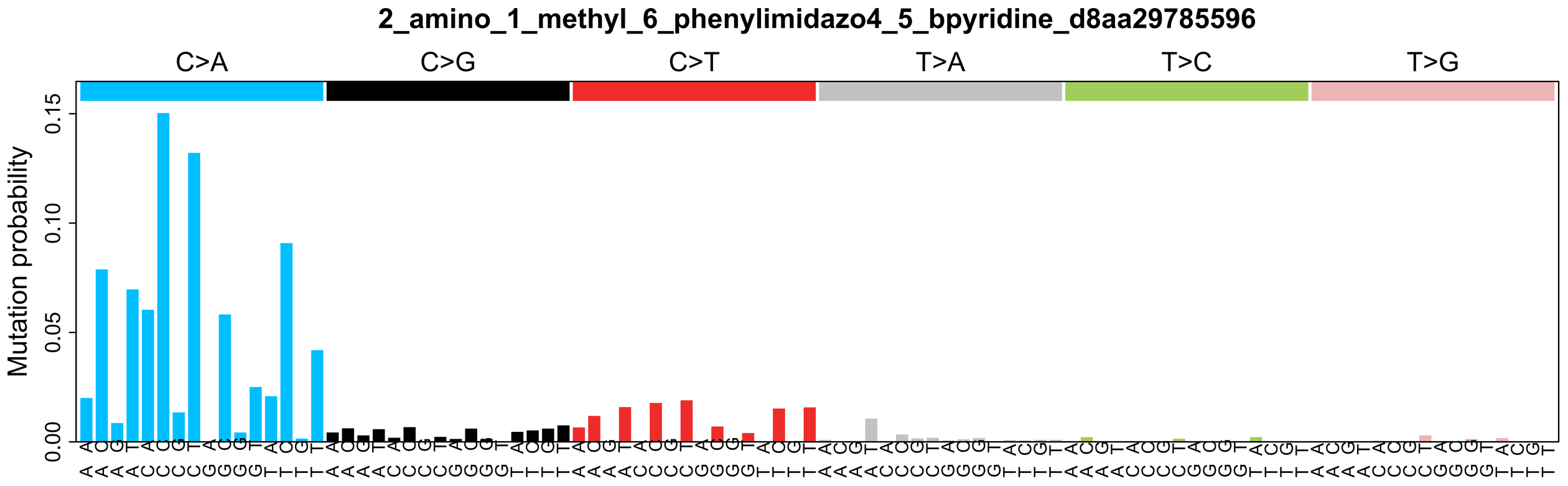

1_6_dinitropyrene_77c0ec6159e4
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00

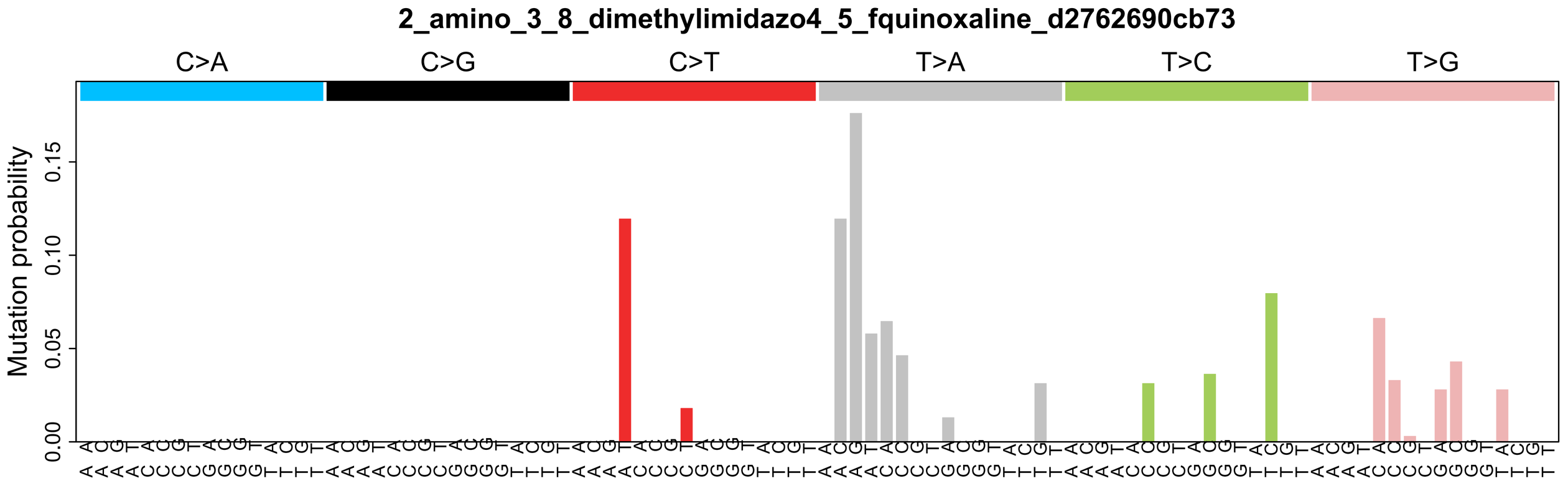

1_8_dinitropyrene_da7ddba98e4a
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.05
0.00

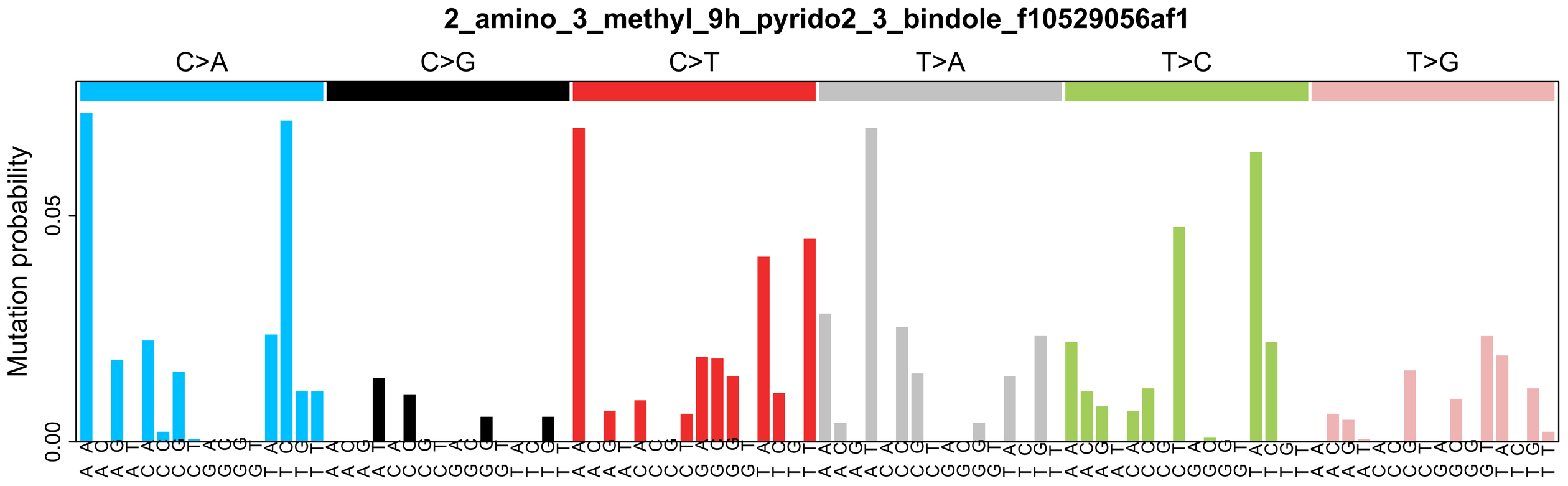

1_nitropyrene_2c37b07dec5a
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2

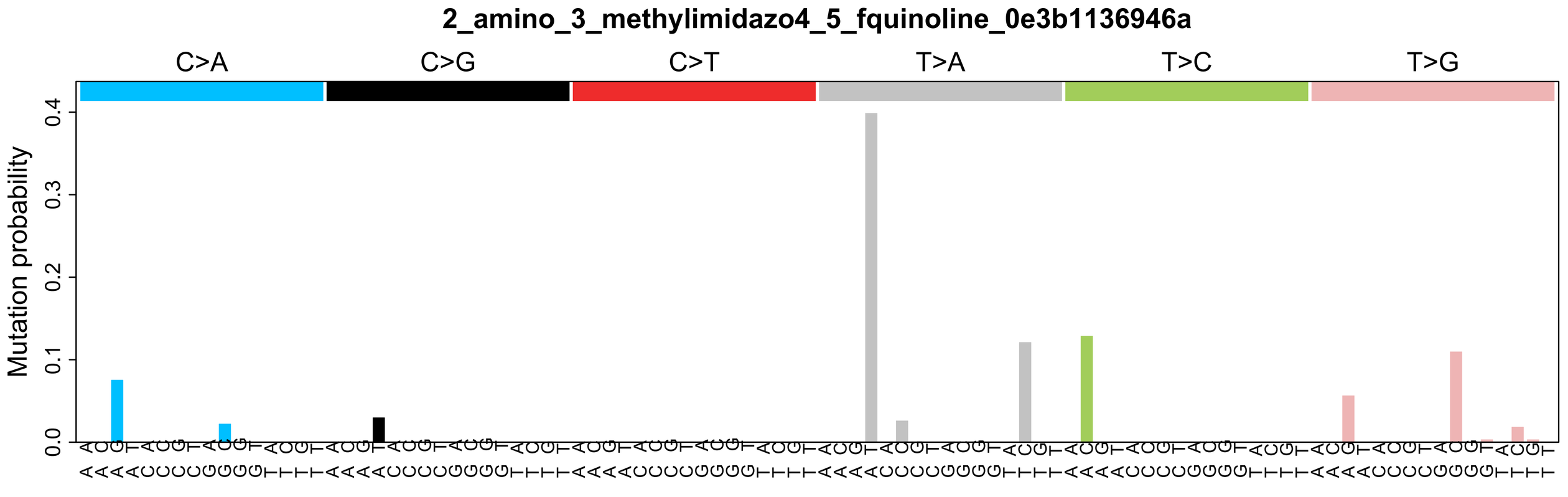

2_6_dimethylaniline_3de14f6a14d9
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10

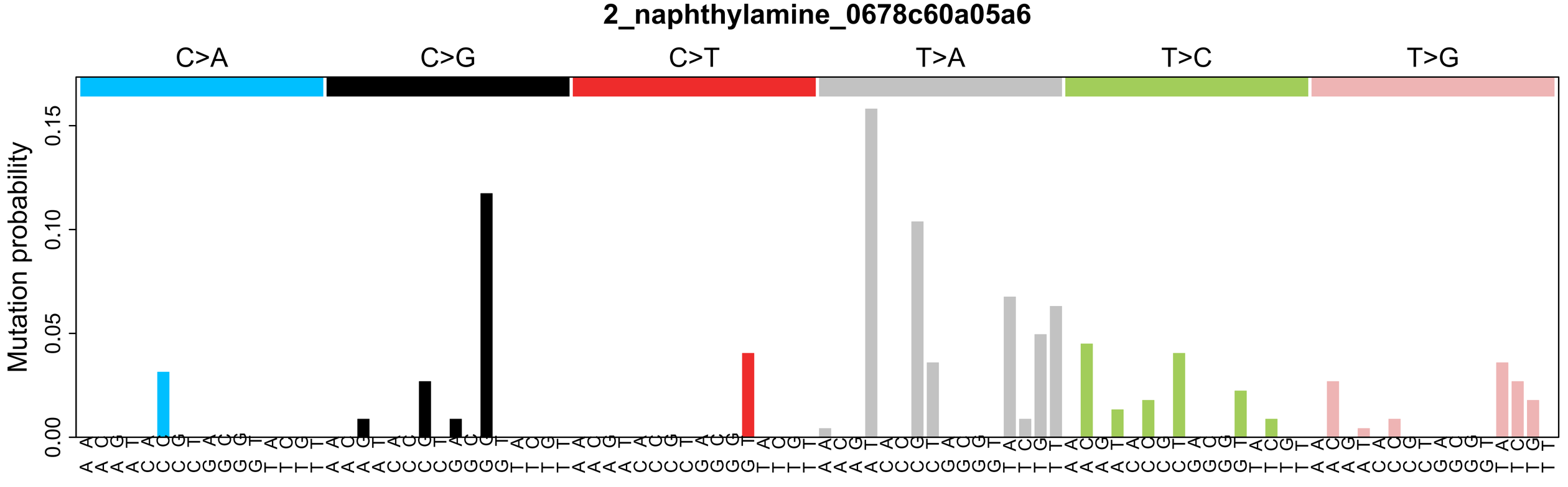

2_6_dimethylaniline_42d3abf9de65
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2

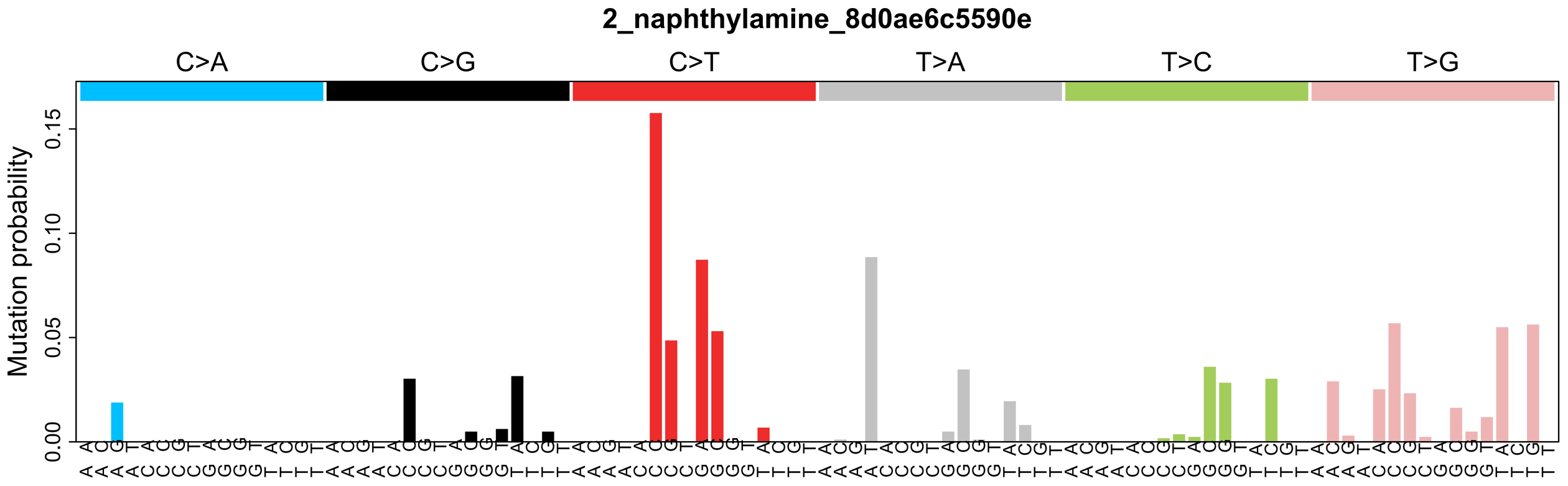

2_amino_1_methyl_6_phenylimidazo4_5_bpyridine_d8aa29785596
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15

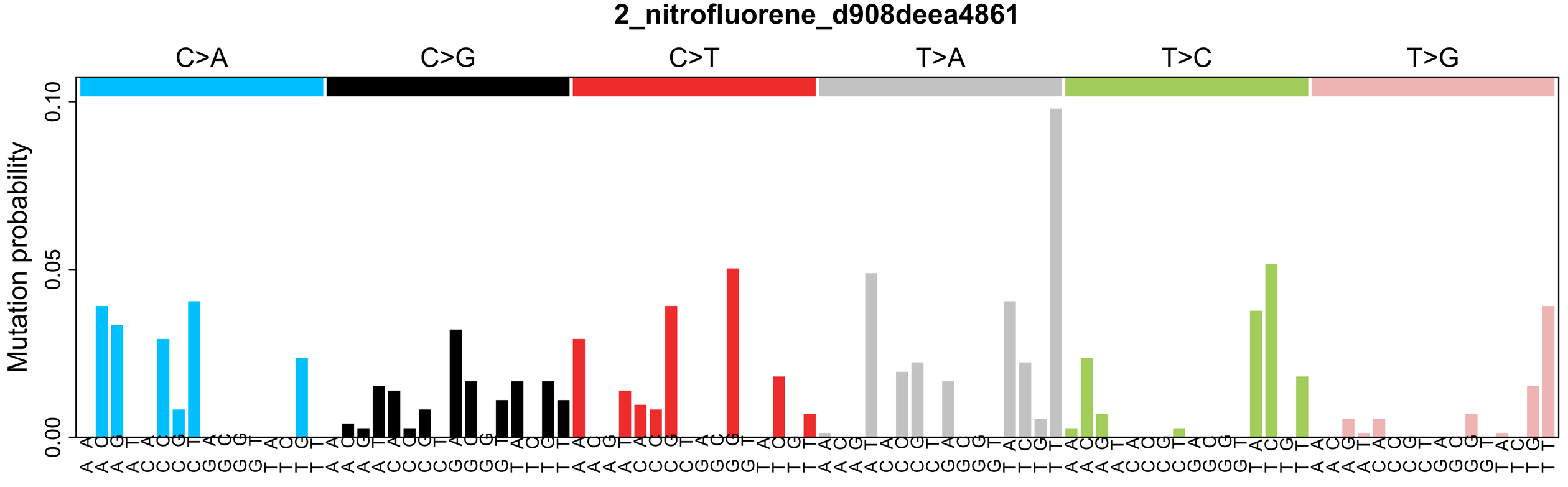

2_amino_3_8_dimethylimidazo4_5_fquinoxaline_d2762690cb73
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15

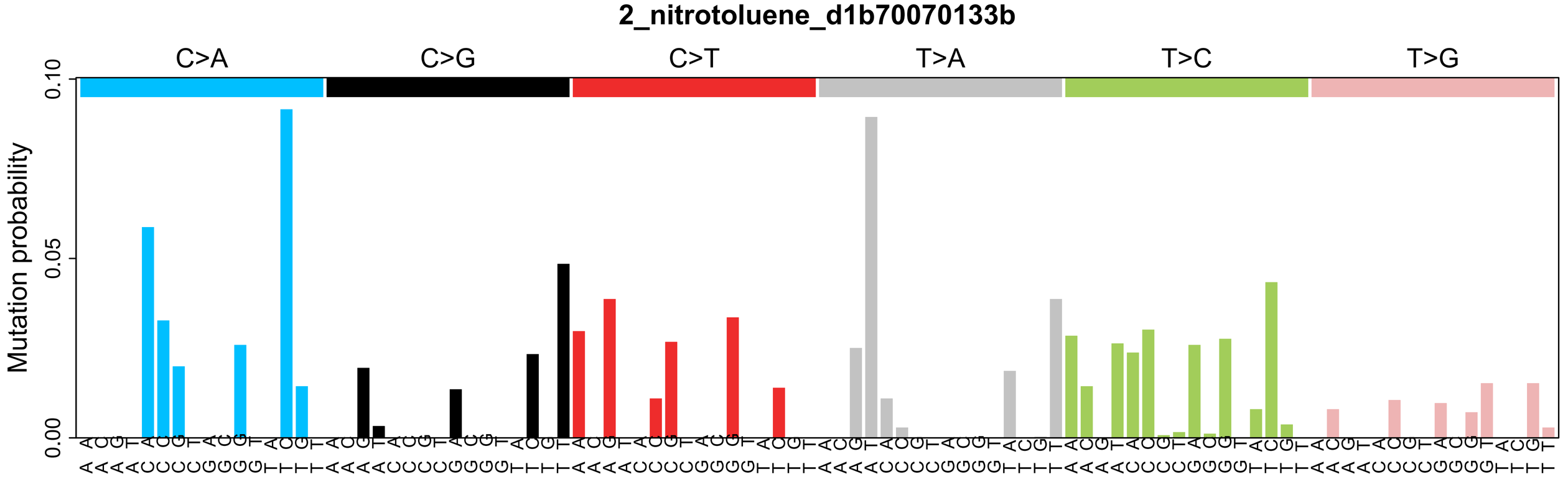

2_amino_3_methyl_9h_pyrido2_3_bindole_f10529056af1
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05

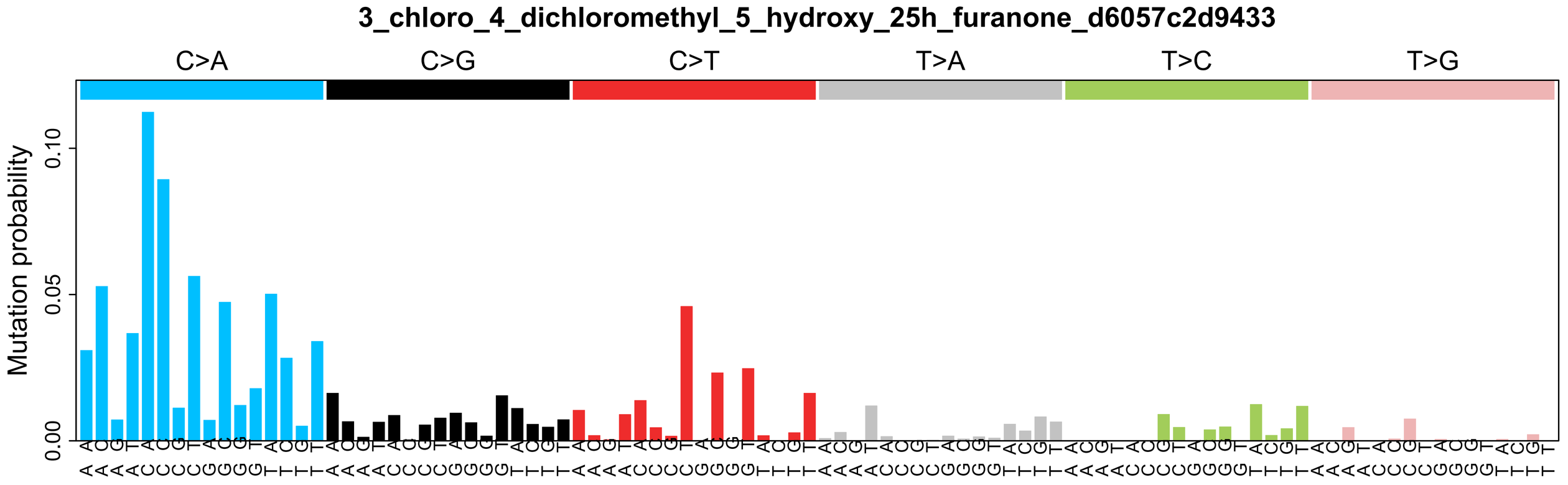

2_amino_3_methylimidazo4_5_fquinoline_0e3b1136946a
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2
0.3
0.4

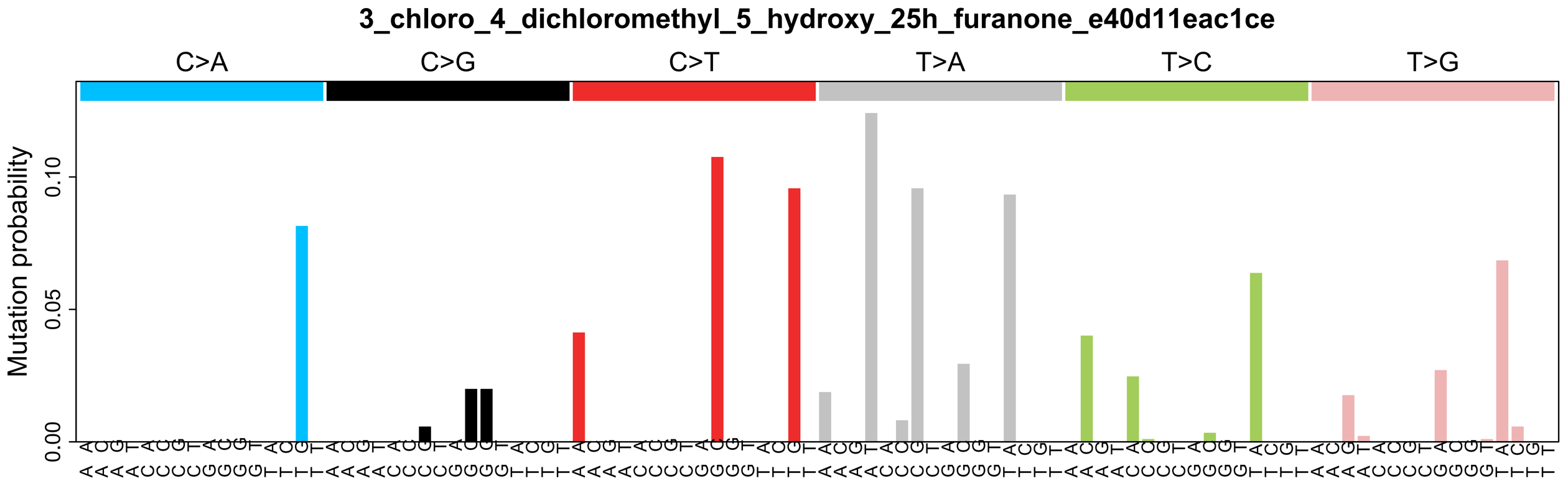

2_naphthylamine_0678c60a05a6
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15

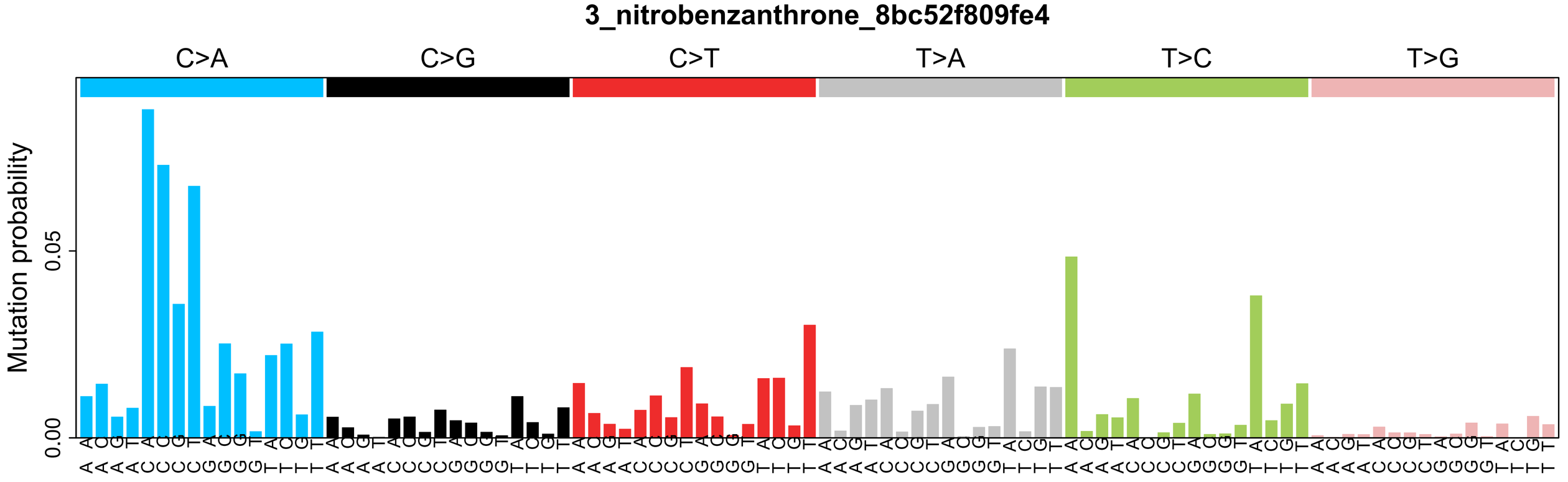

2_naphthylamine_8d0ae6c5590e
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15

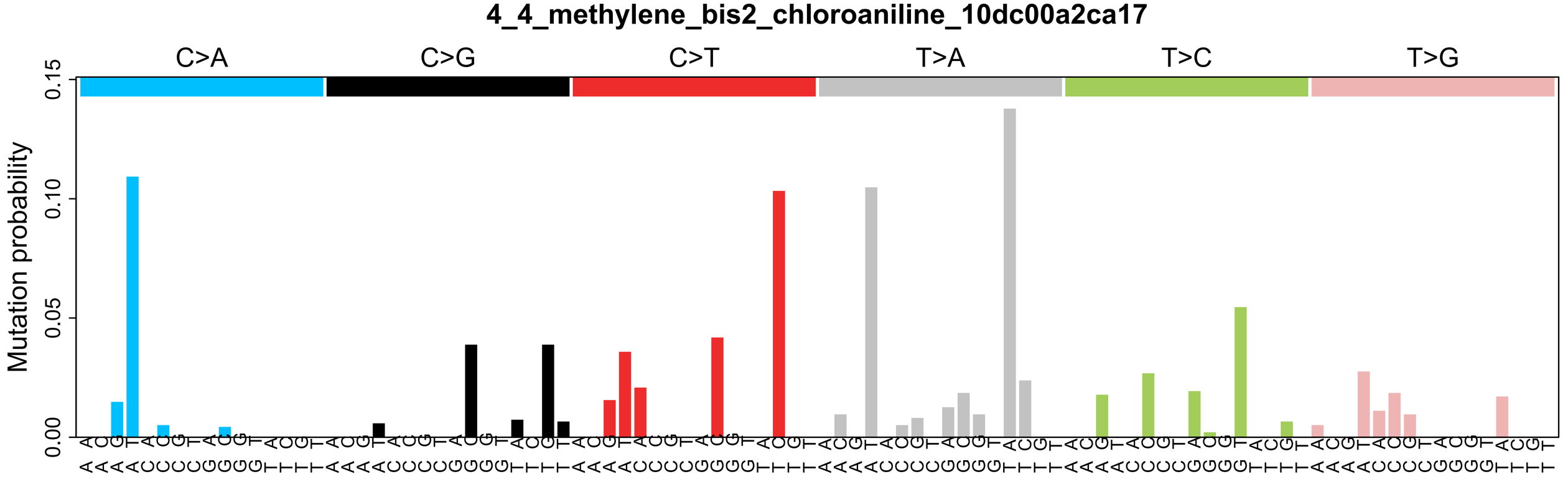

2_nitrofluorene_d908deea4861
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


2_nitrotoluene_d1b70070133b
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


3_chloro_4_dichloromethyl_5_hydroxy_25h_furanone_d6057c2d9433
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


3_chloro_4_dichloromethyl_5_hydroxy_25h_furanone_e40d11eac1ce
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


3_nitrobenzanthrone_8bc52f809fe4
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05


4_4_methylene_bis2_chloroaniline_10dc00a2ca17
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
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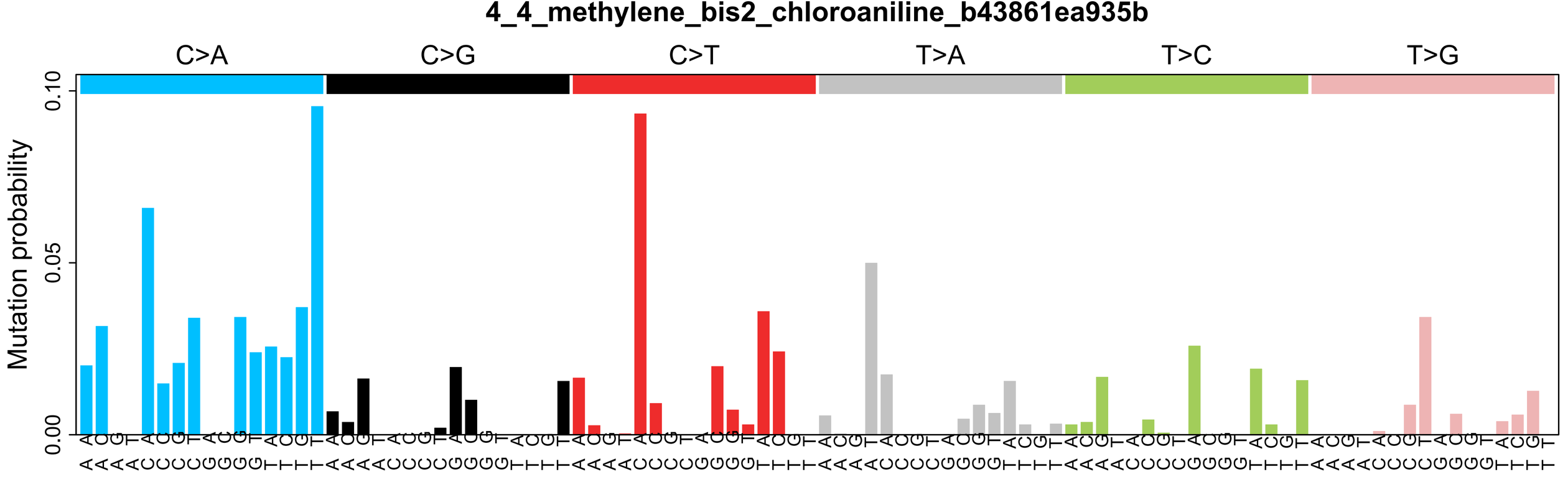
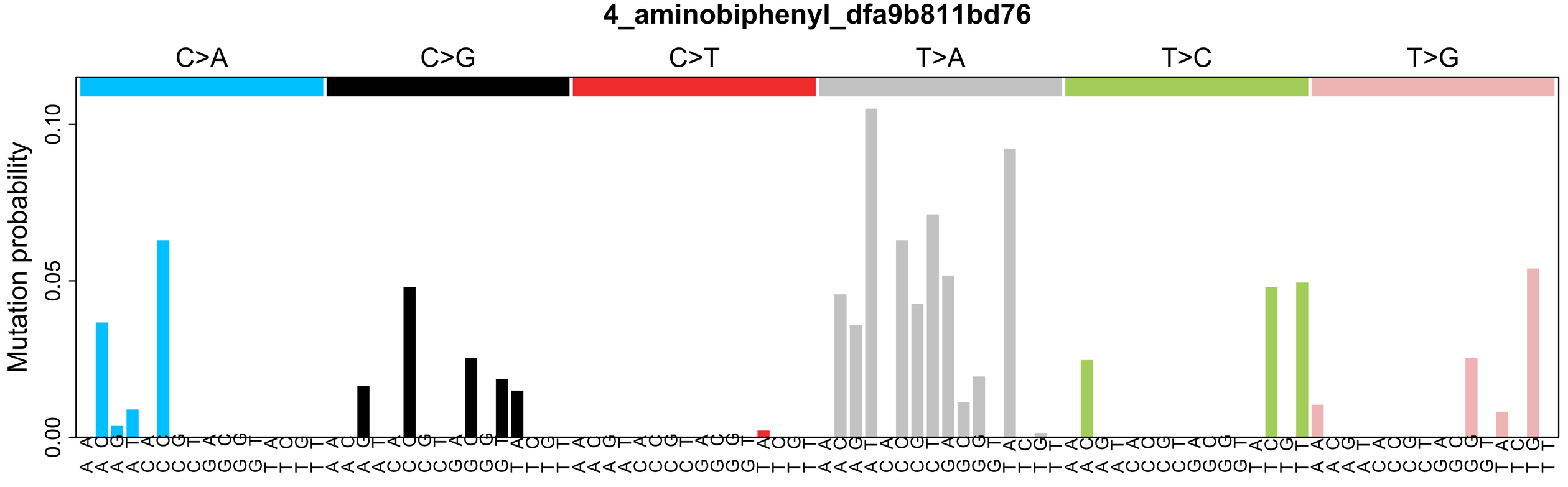
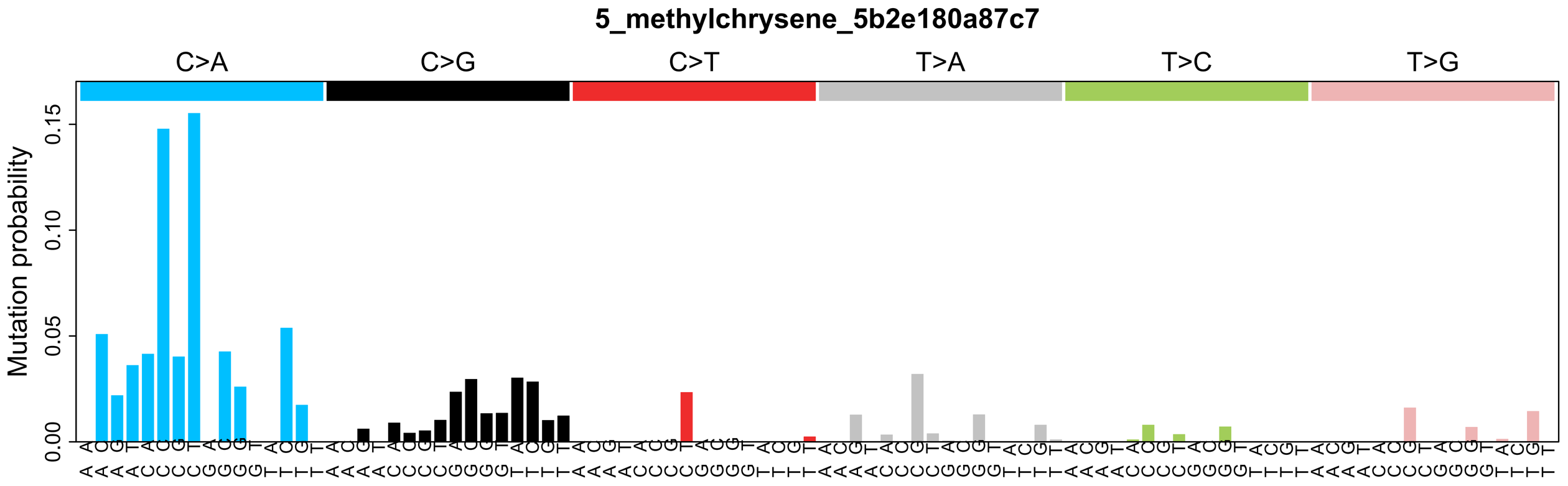
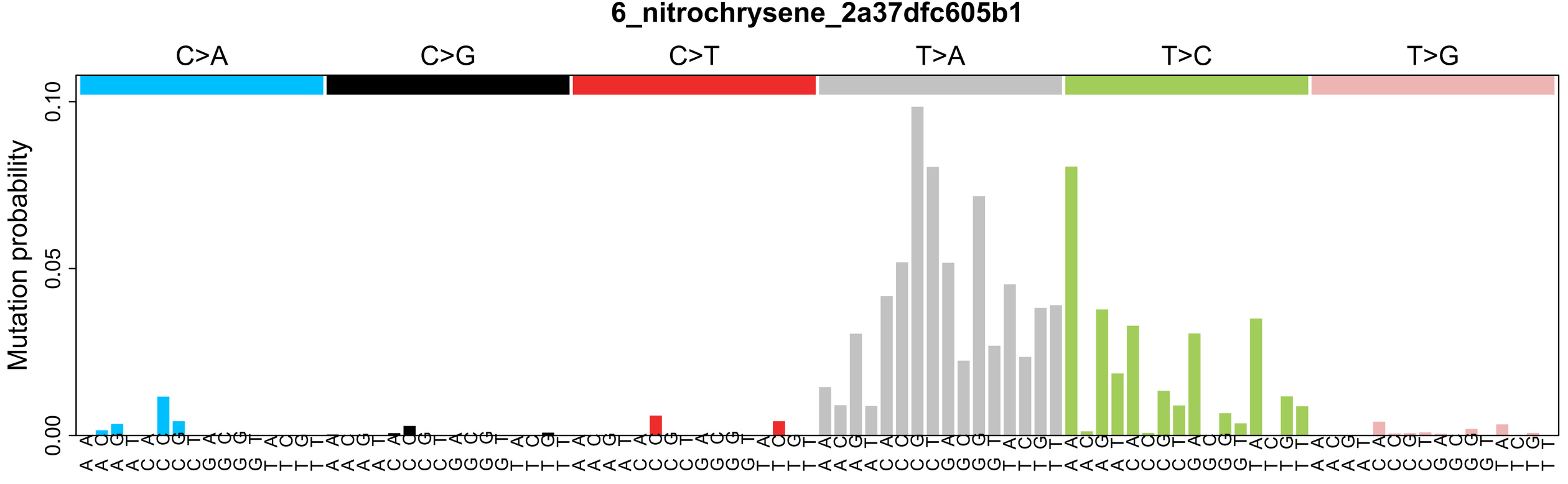
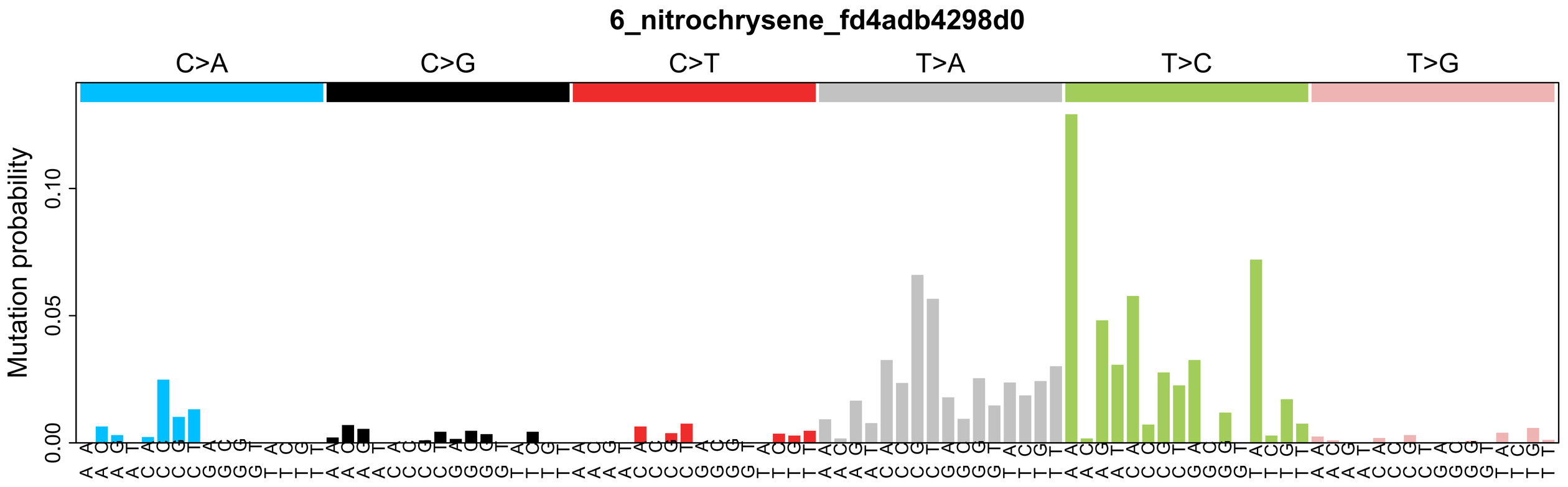
0.10
0.05
0.00


4_4_methylene_bis2_chloroaniline_b43861ea935b
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


4_aminobiphenyl_dfa9b811bd76
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


5_methylchrysene_5b2e180a87c7
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15


6_nitrochrysene_2a37dfc605b1
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10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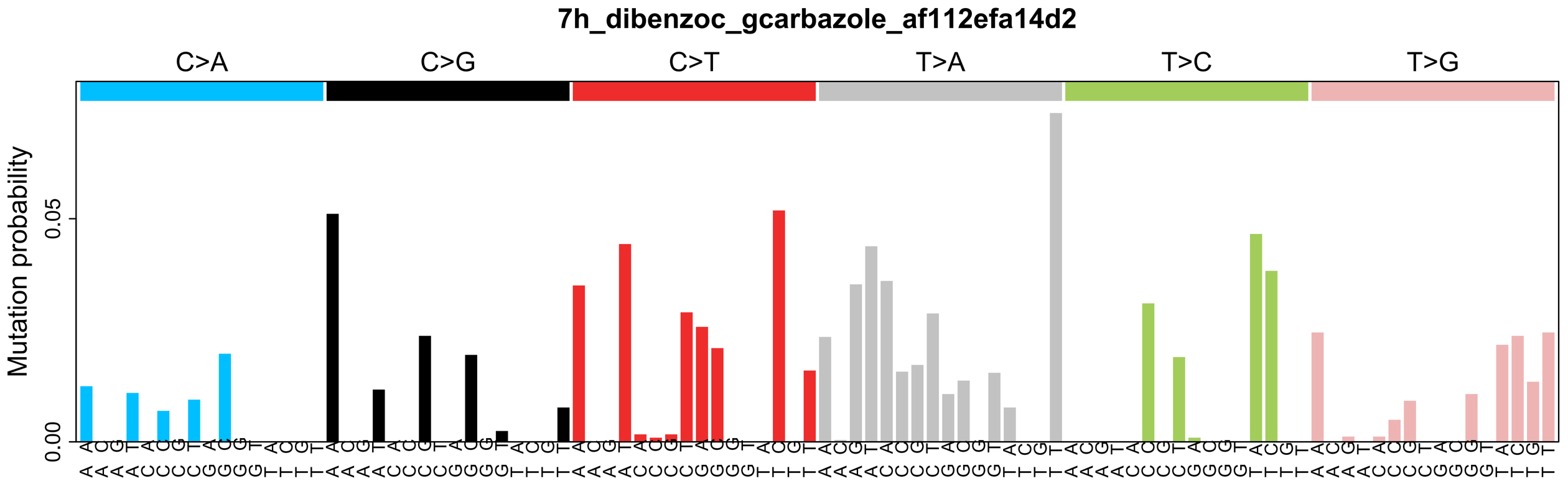
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


7h_dibenzoc_gcarbazole_af112efa14d2
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C>T
T>A
T>C
T>G
Mutation probability
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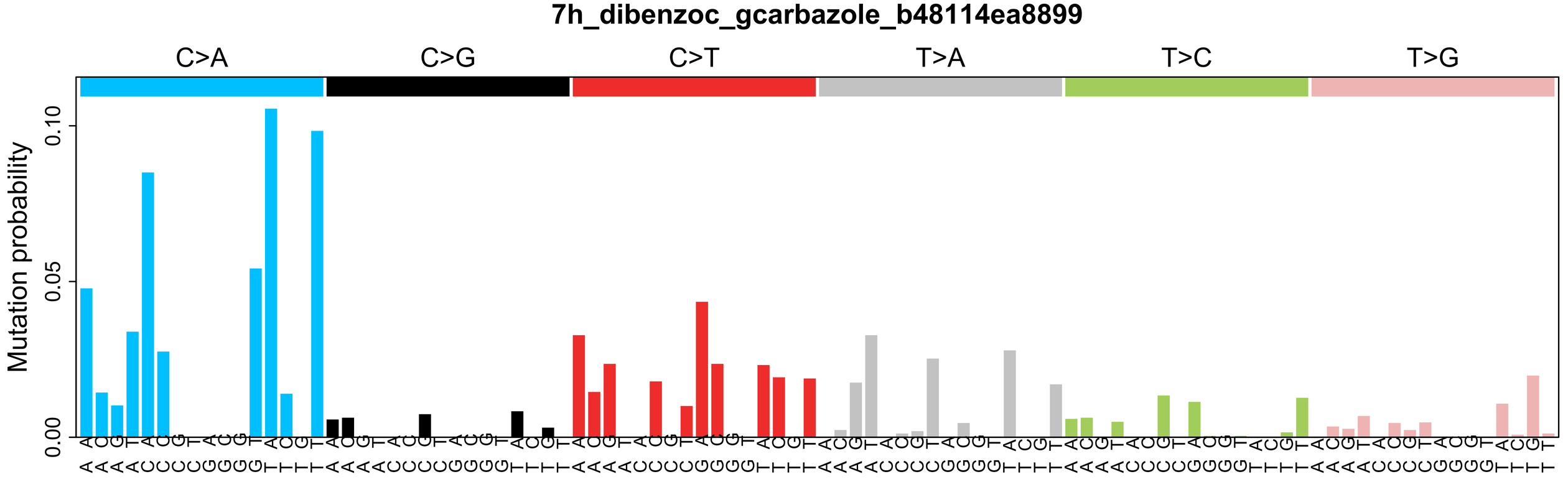
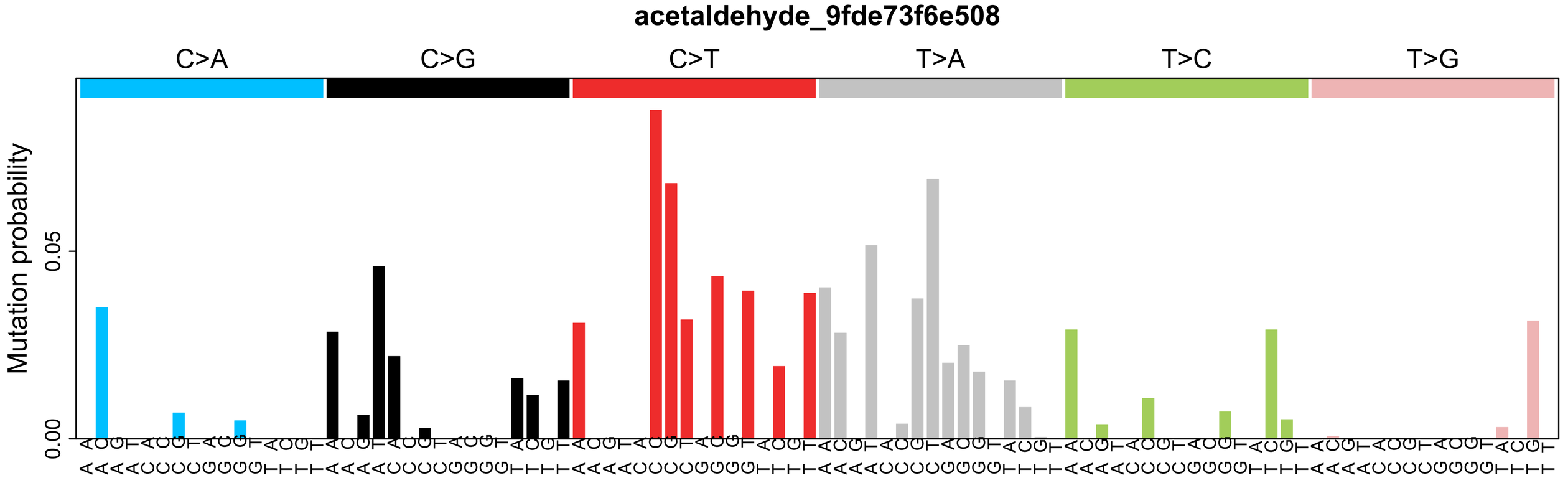
0.05


7h_dibenzoc_gcarbazole_b48114ea8899
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10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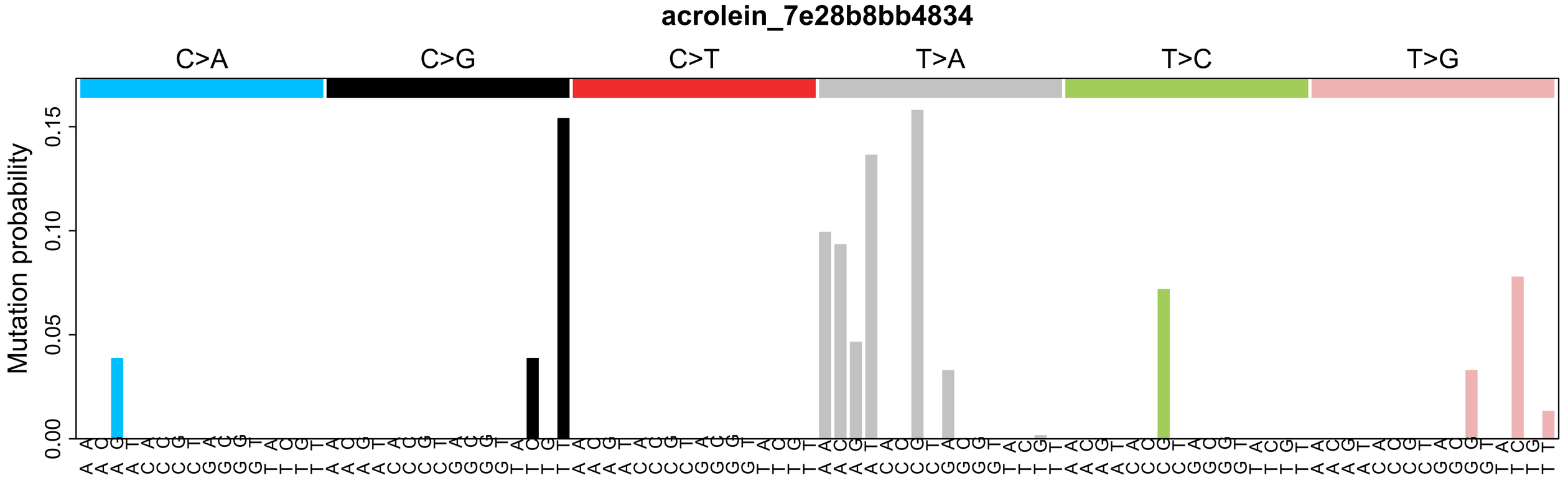
T>C
T>G
Mutation probability
0.00
0.05


acrolein_7e28b8bb4834
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C>G
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T>A
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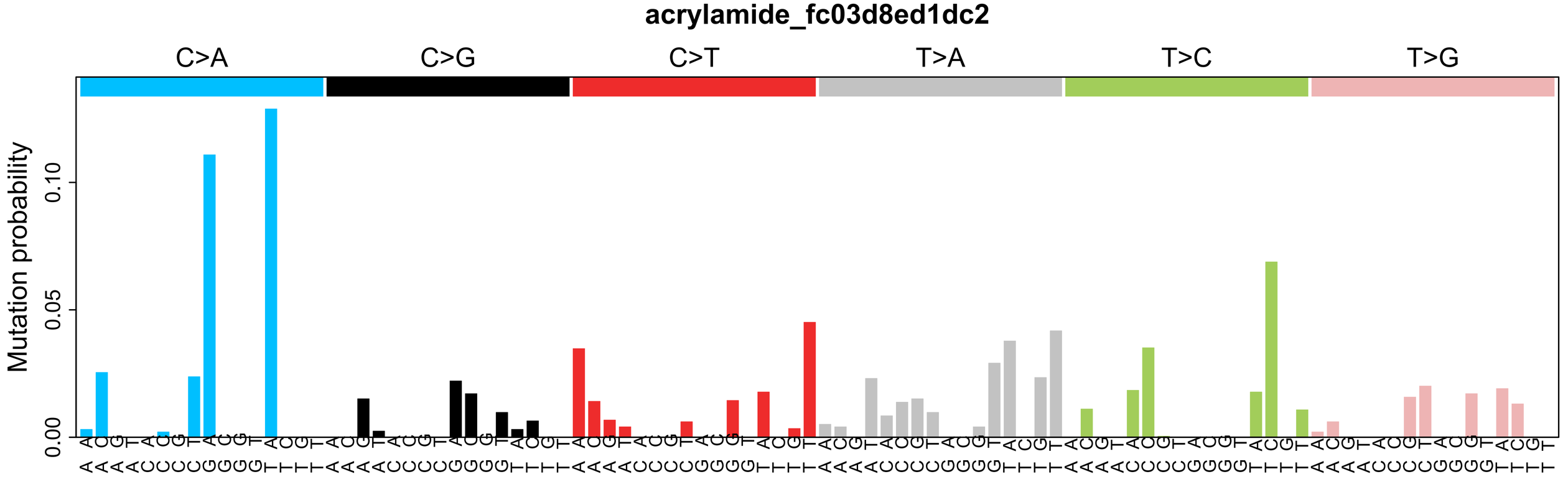
T>G
Mutation probability
0.00
0.05
0.10
0.15


acrylamide_fc03d8ed1dc2
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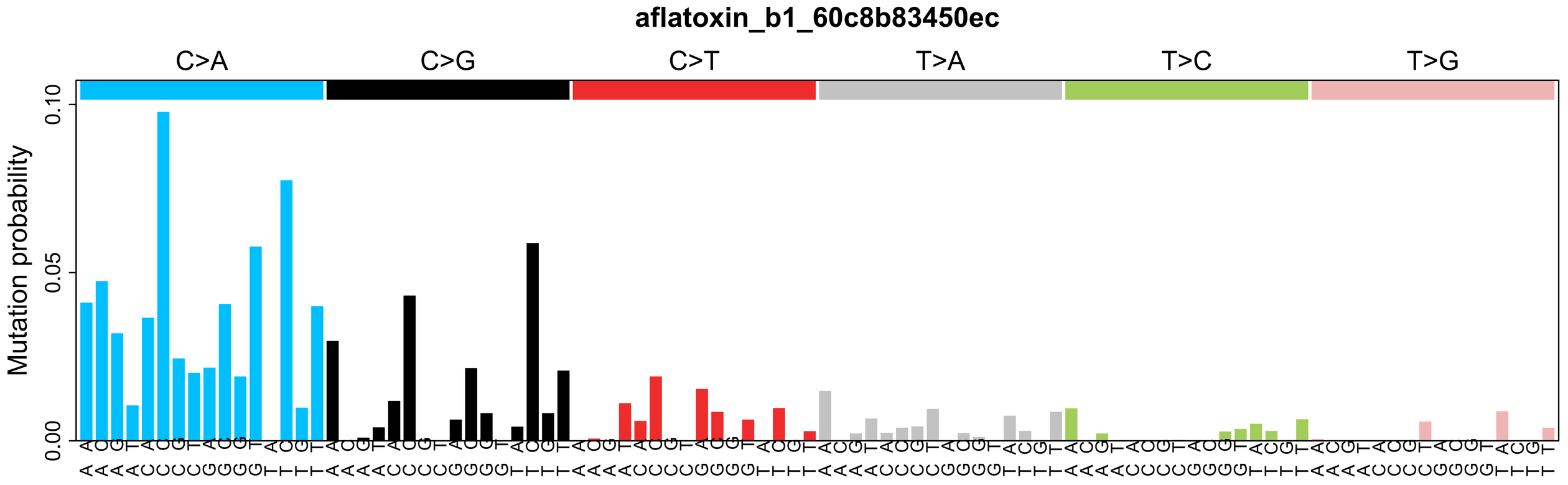
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


aflatoxin_b1_60c8b83450ec
C>A
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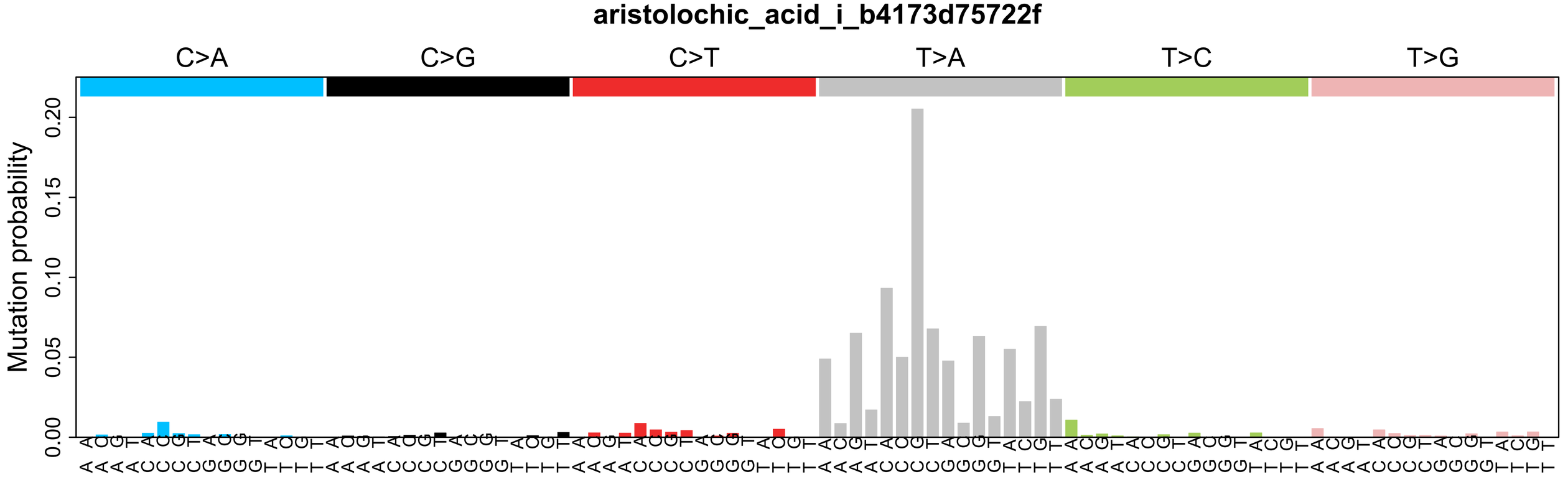
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


aristolochic_acid_i_b4173d75722f
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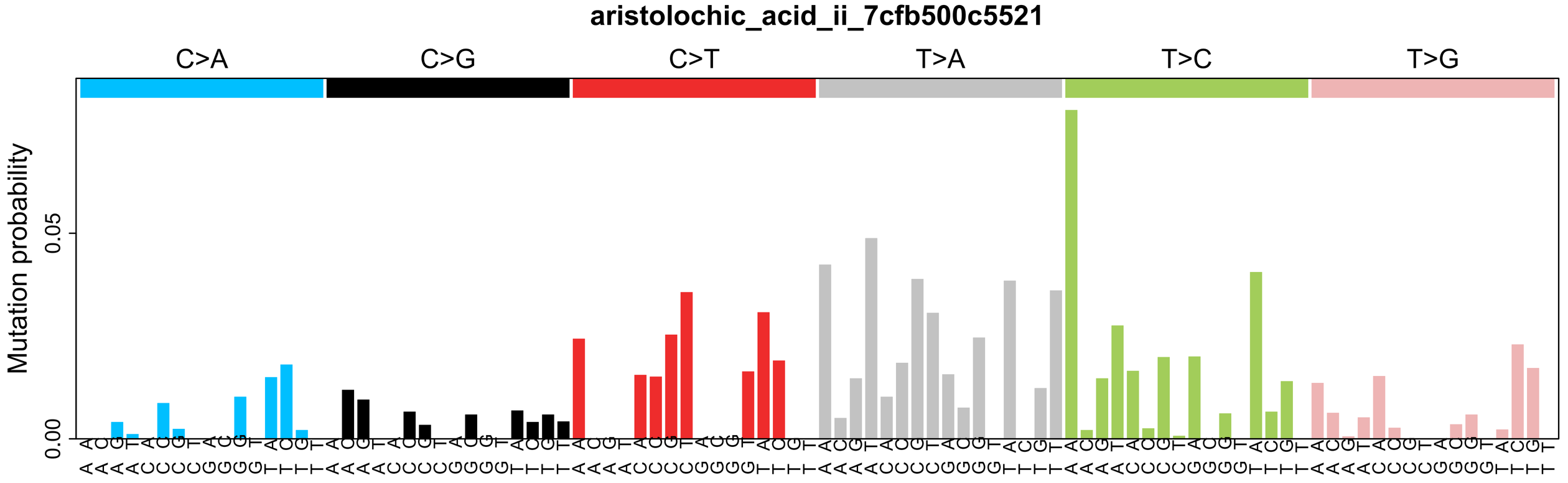
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15
0.20


aristolochic_acid_ii_7cfb500c5521
C>A
C>G
C>T
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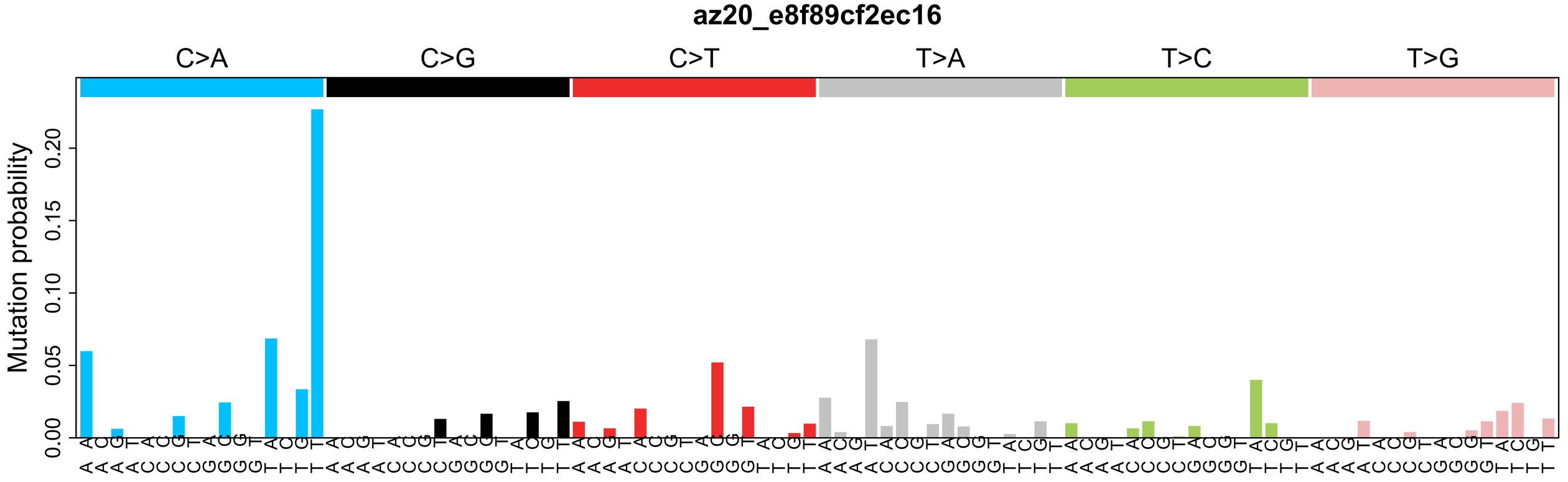
T>C
T>G
Mutation probability
0.00
0.05


az20_e8f89cf2ec16
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C>T
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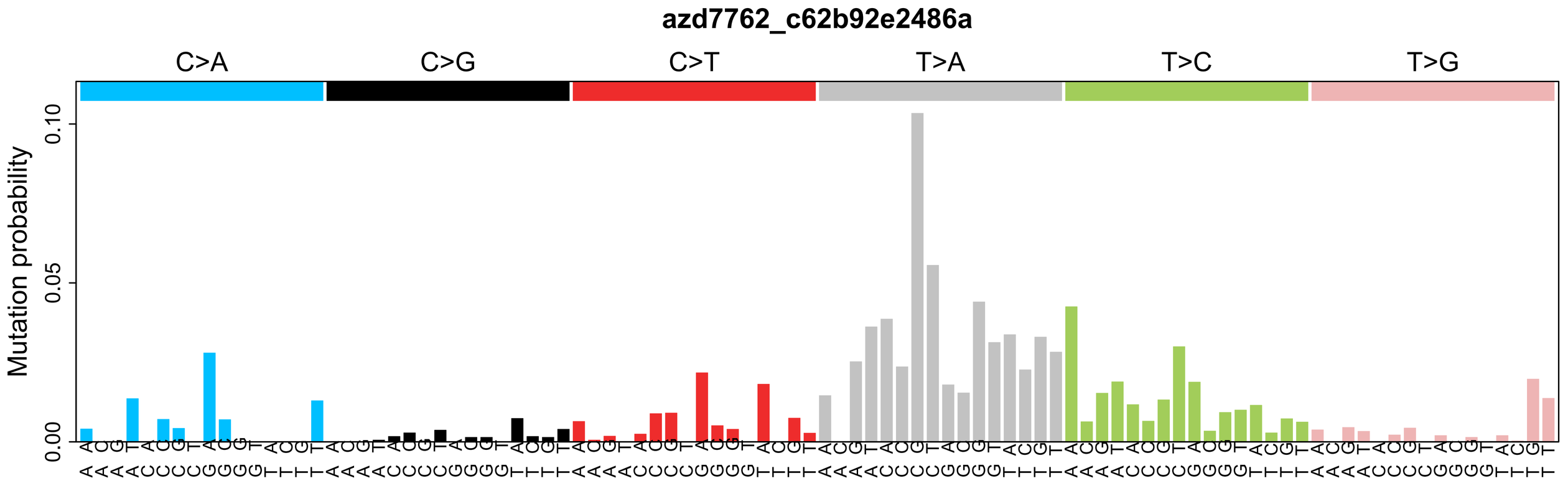
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15
0.20


azd7762_c62b92e2486a
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C>G
C>T
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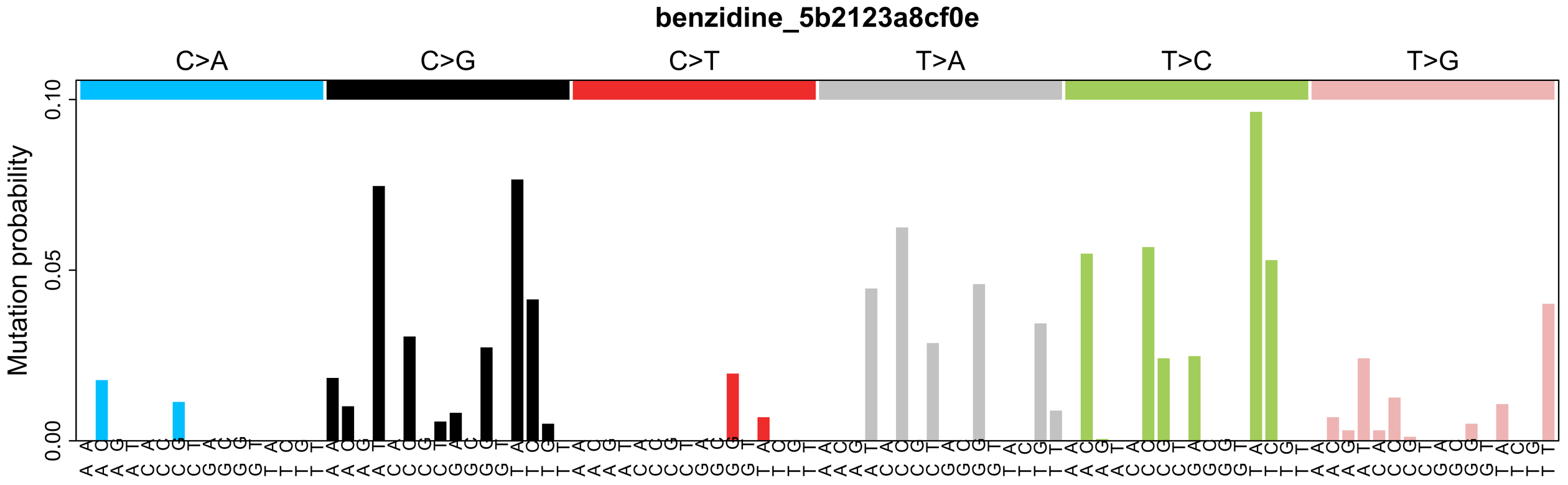
Mutation probability
0.00
0.05
0.10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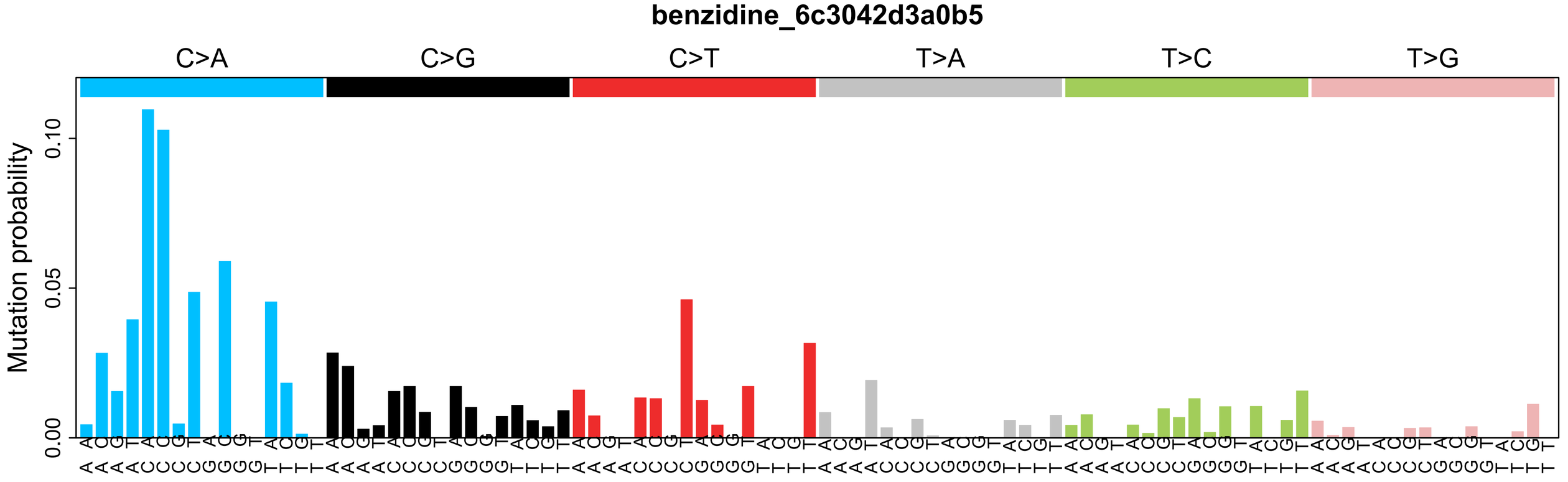
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


benzidine_6c3042d3a0b5
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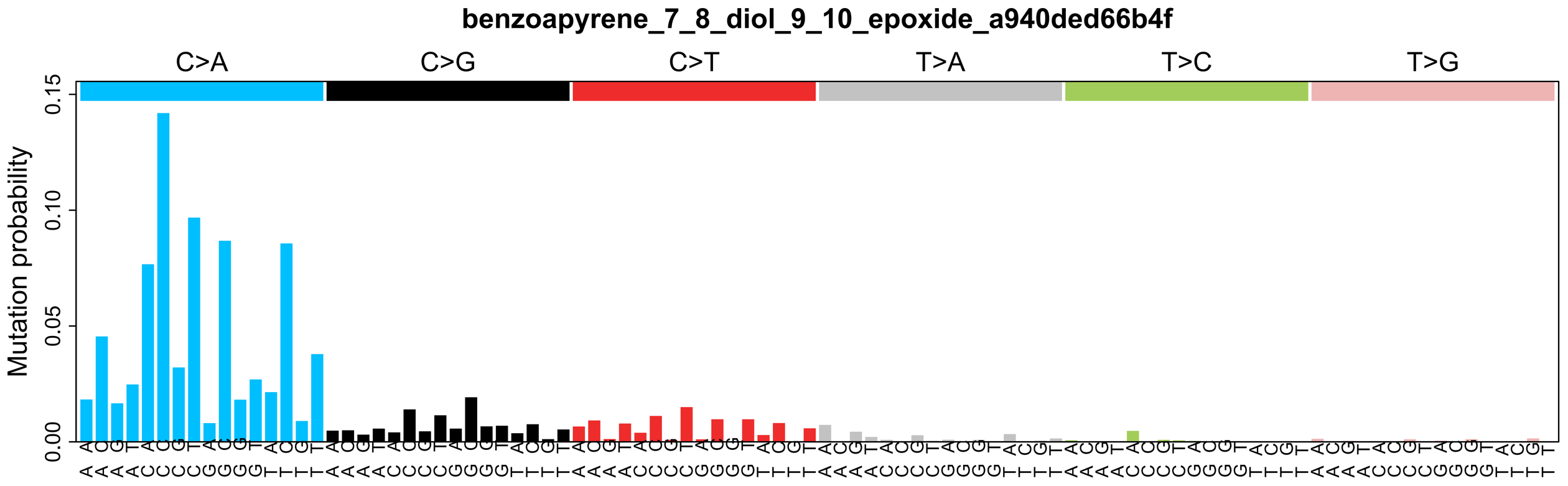
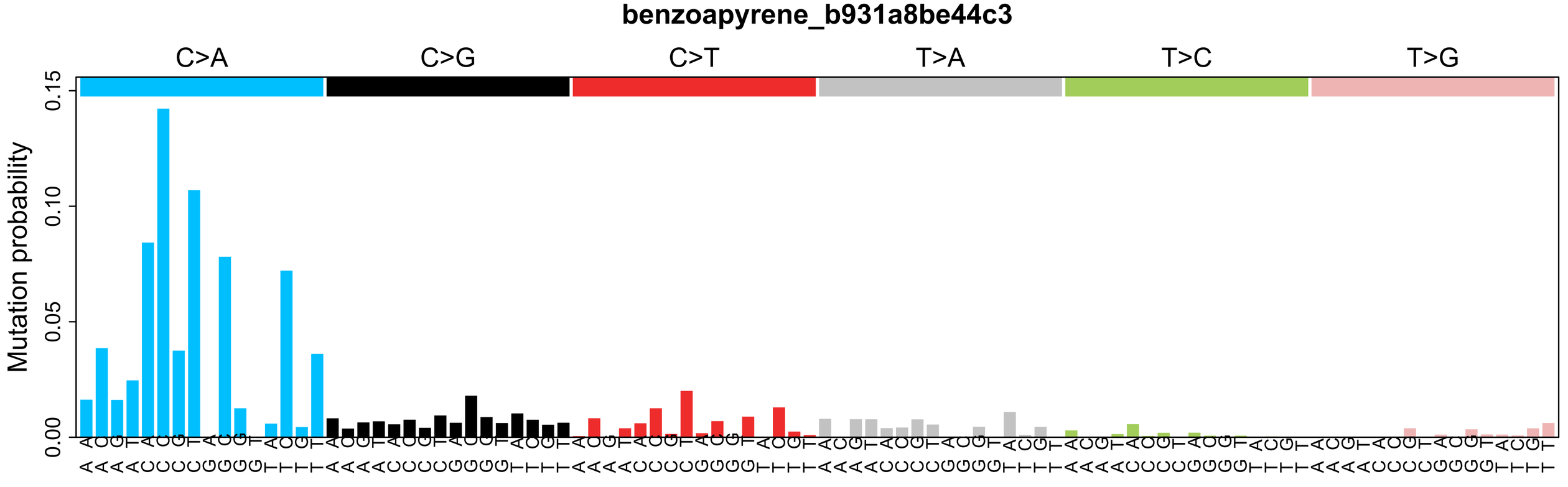
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


benzoapyrene_7_8_diol_9_10_epoxide_a940ded66b4f
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15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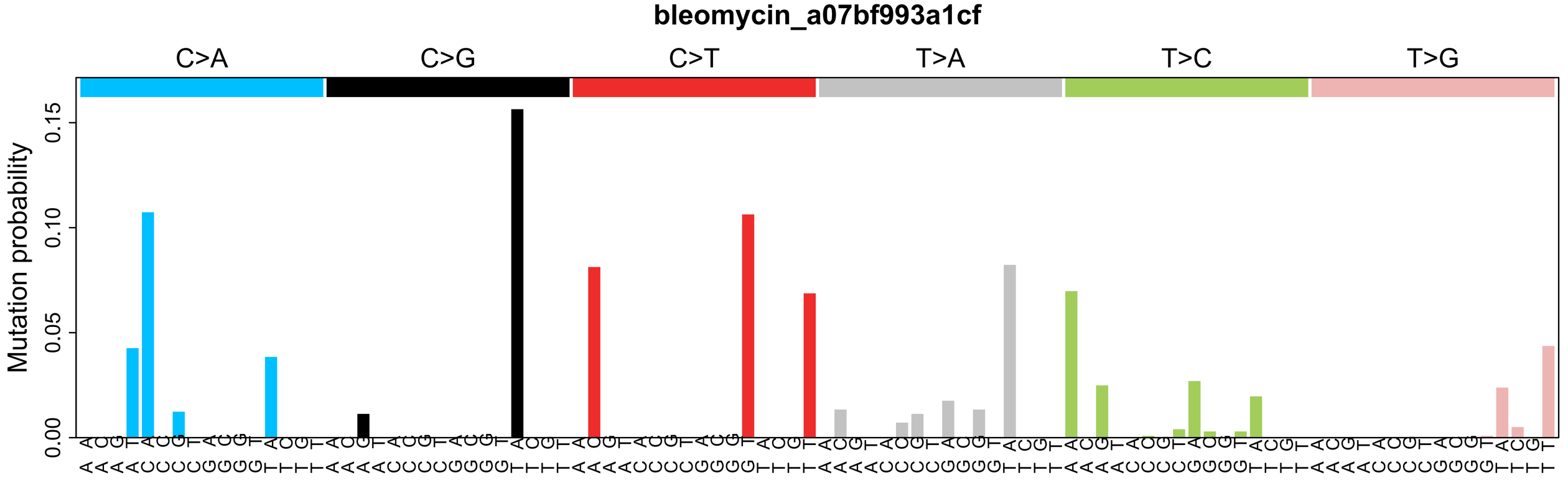
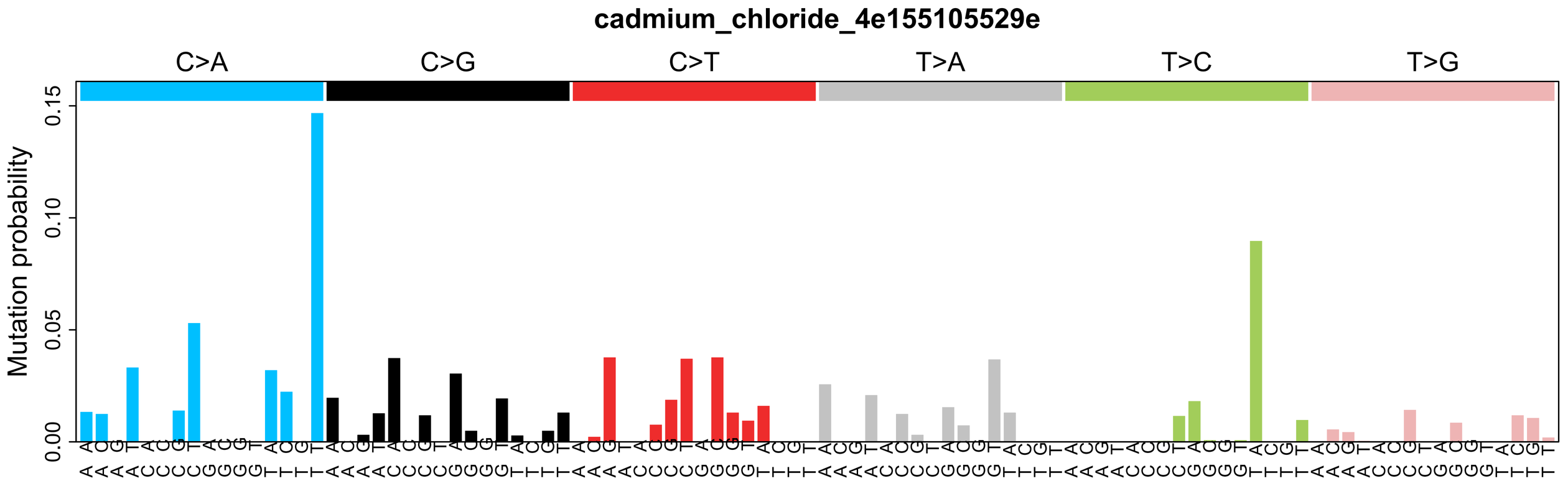
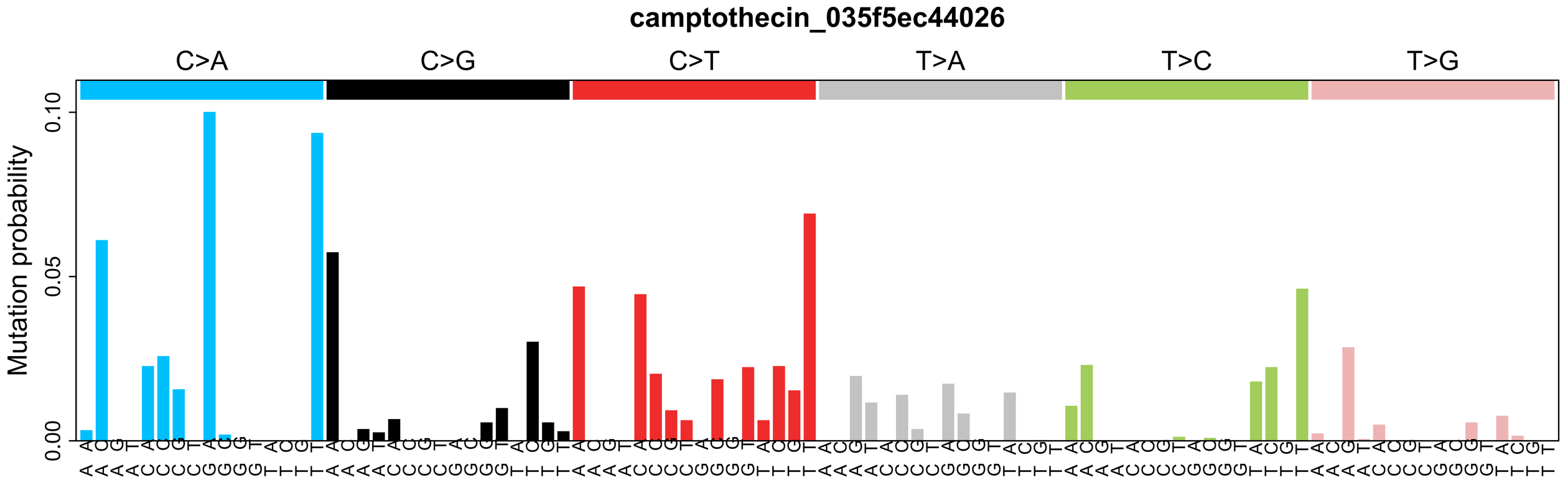
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15


bleomycin_a07bf993a1cf
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15


cadmium_chloride_4e155105529e
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15


camptothecin_035f5ec44026
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C>T
T>A
T>C
T>G
Mutation probability
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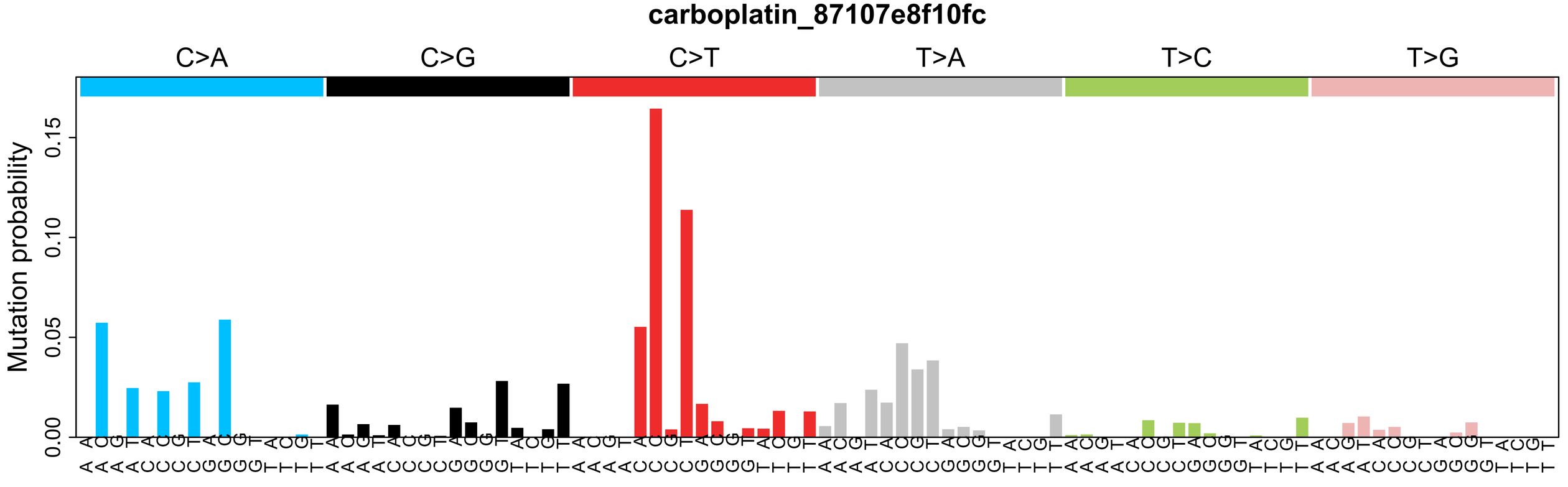
0.05
0.10


carboplatin_87107e8f10fc
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C>T
T>A
T>C
T>G
Mutation probability
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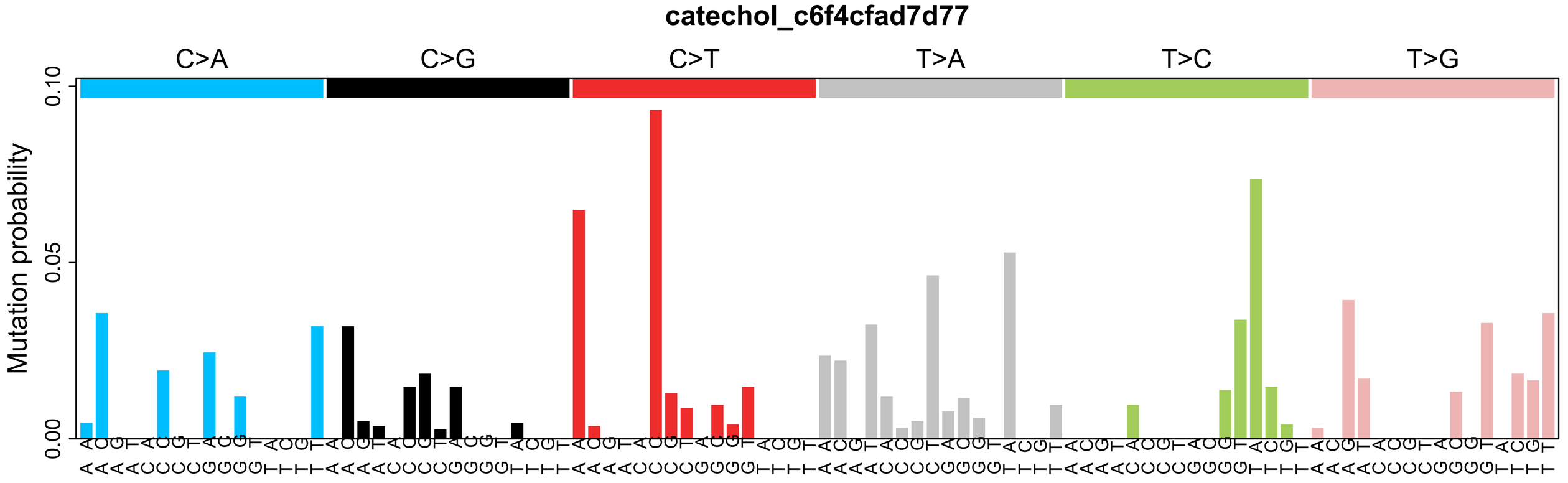
0.05
0.10
0.15


catechol_c6f4cfad7d77
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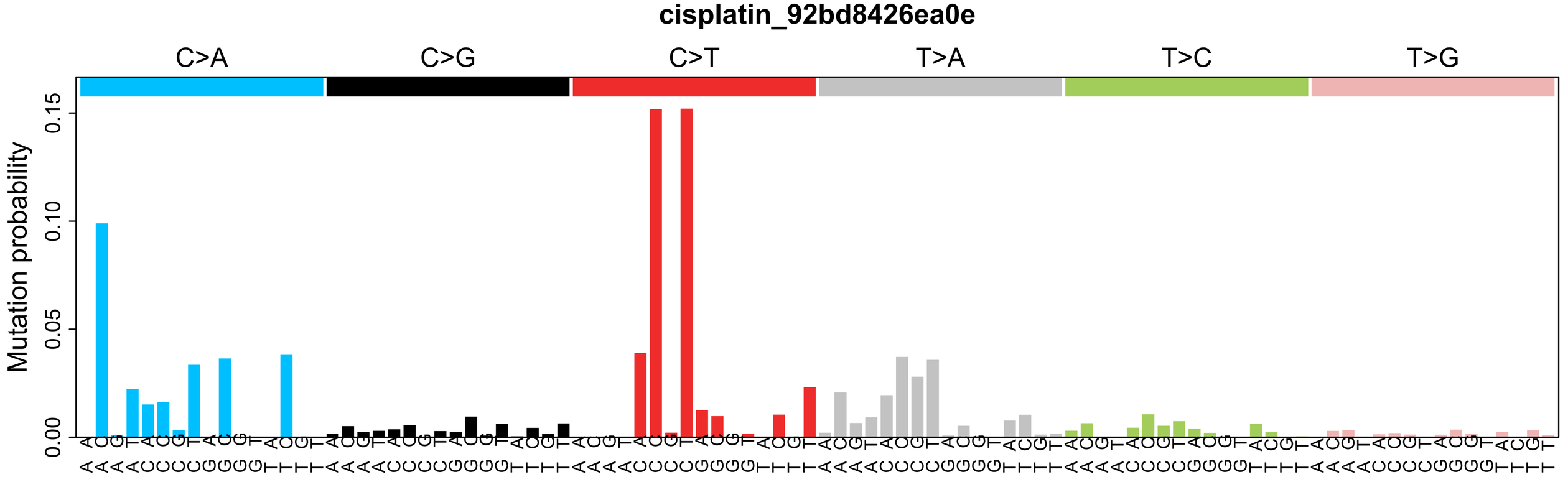
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00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T>G
Mutation probability
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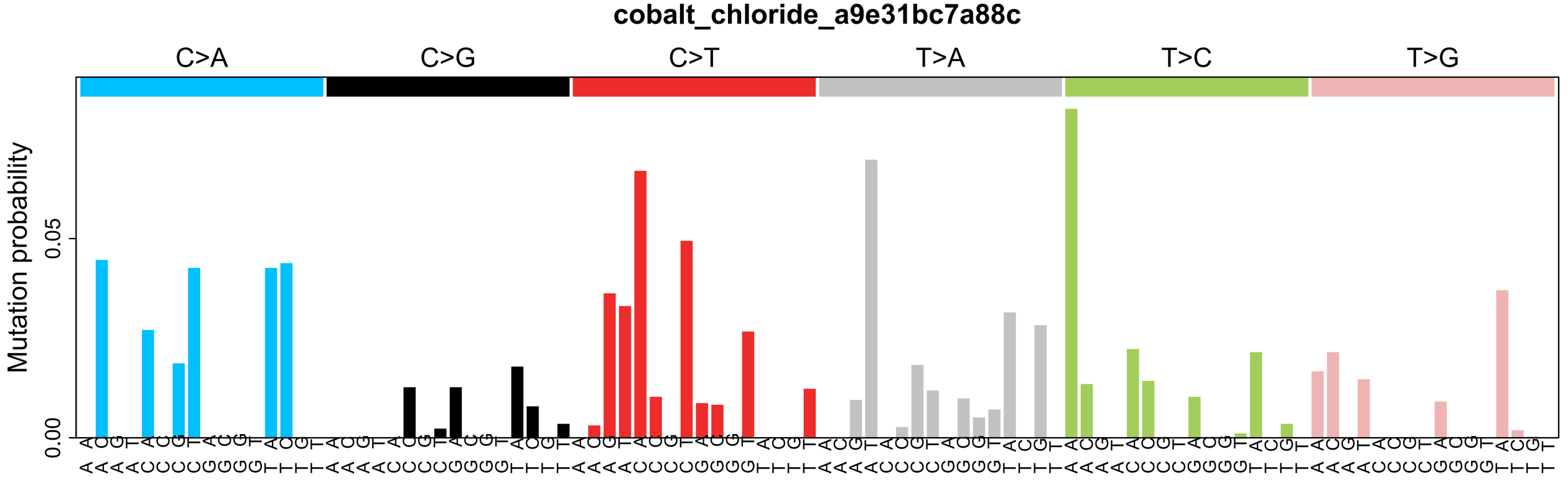
0.05
0.10
0.15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T>G
Mutation probability
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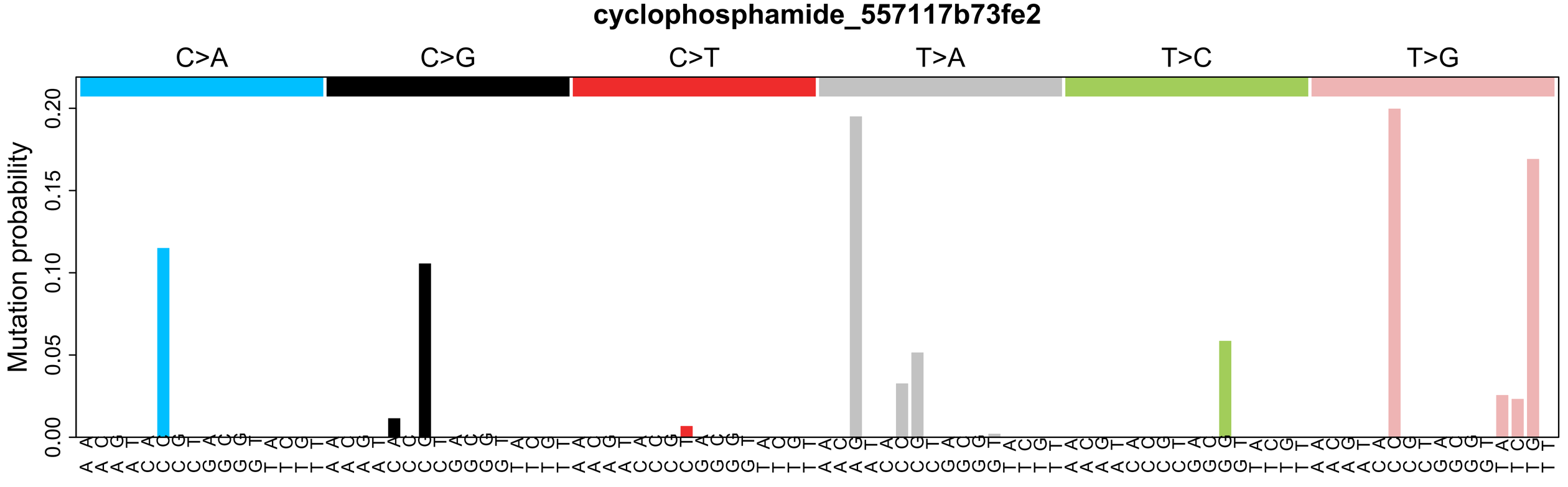
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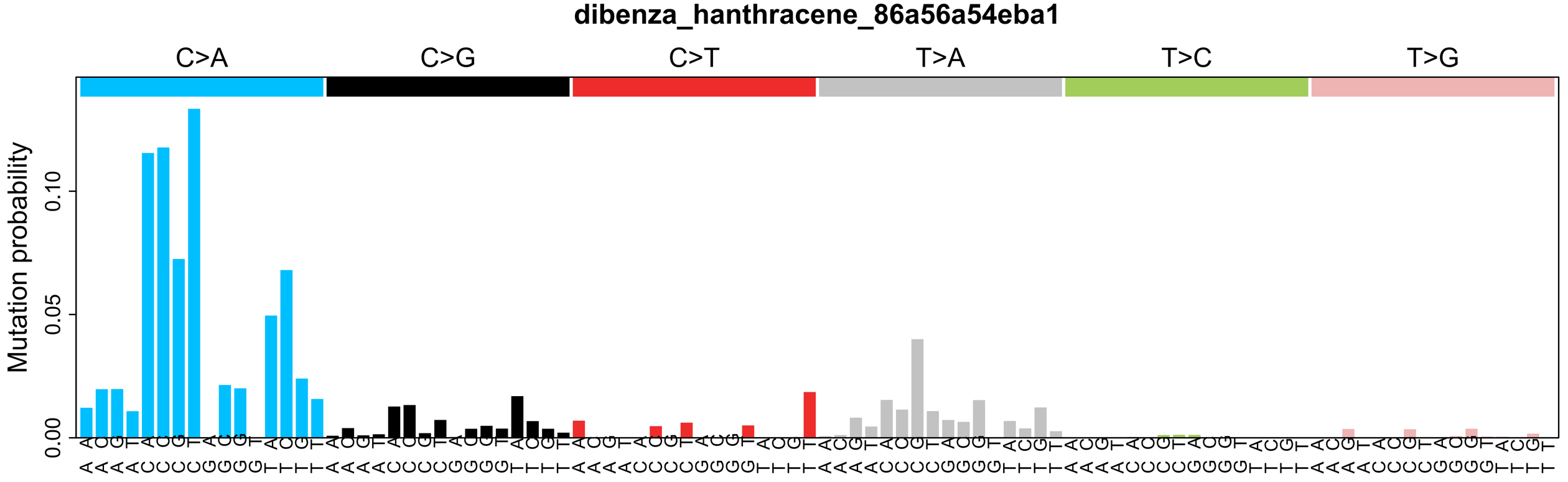
T>G
Mutation probability
0.00
0.05
0.10
0.15
0.20


dibenza_hanthracene_86a56a54eba1
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C>G
C>T
T>A
T>C
T>G
Mutation probability
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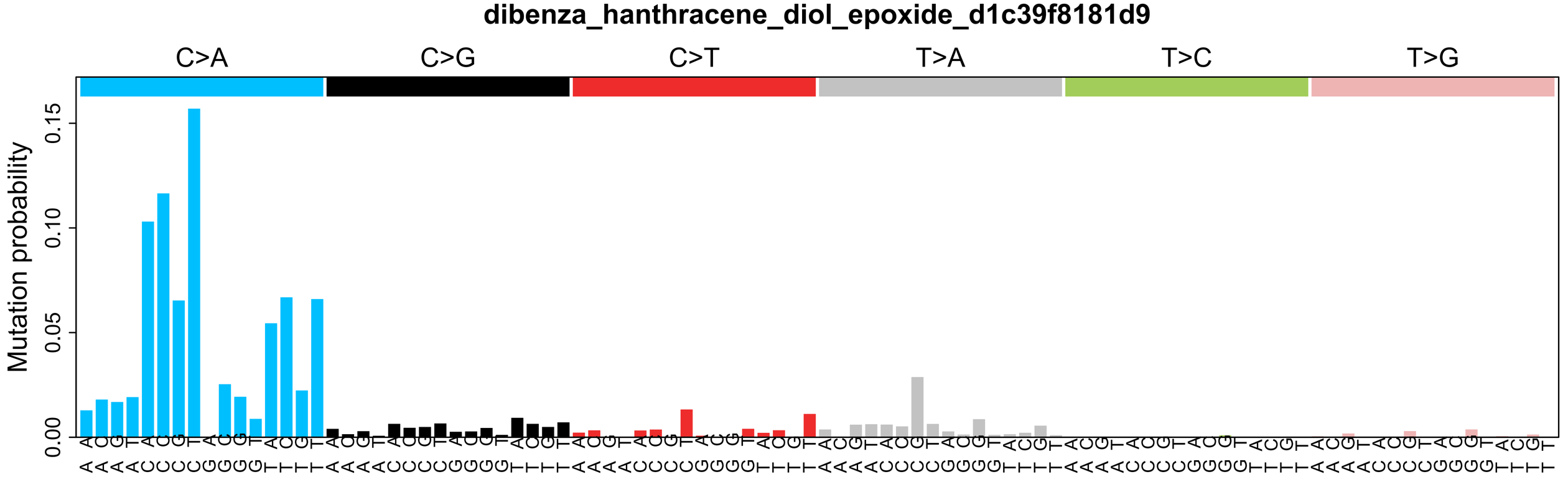
0.05
0.10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C>G
C>T
T>A
T>C
T>G
Mutation probability
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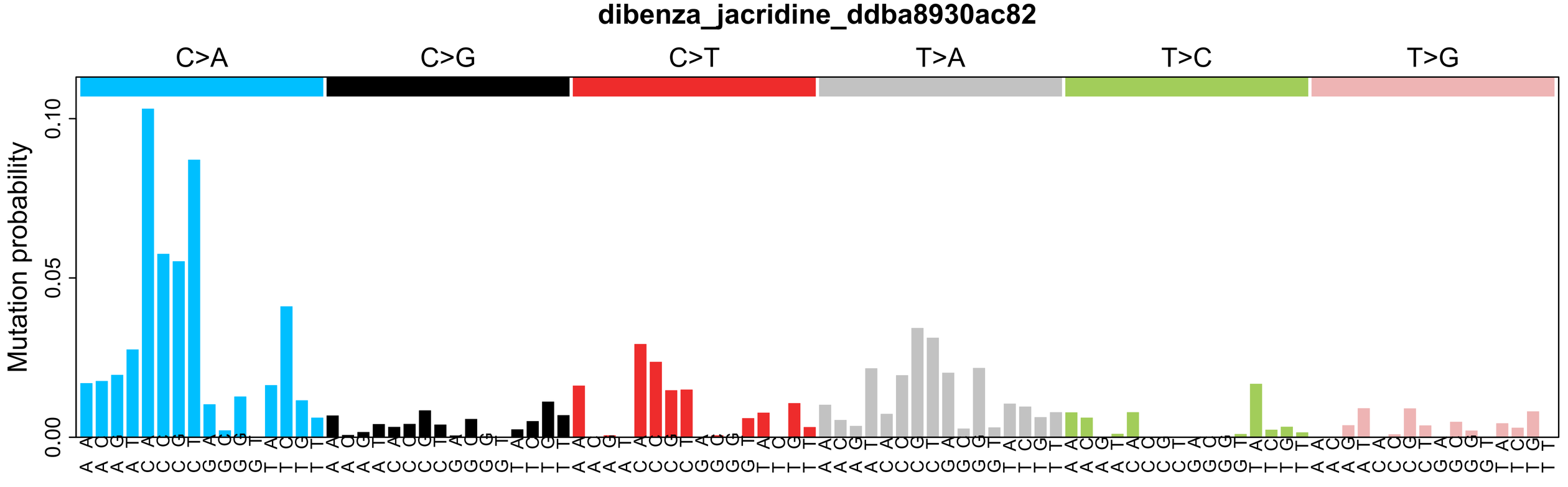
0.05
0.10
0.15


dibenza_jacridine_ddba8930ac82
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C>T
T>A
T>C
T>G
Mutation probability
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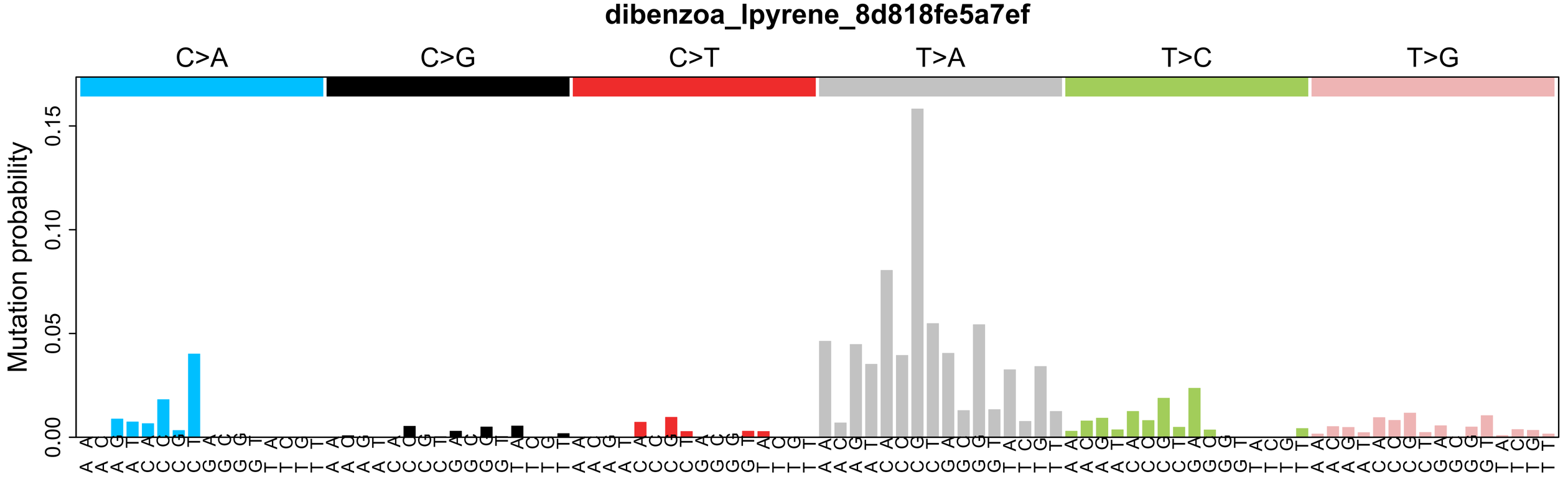
0.05
0.10


dibenzoa_lpyrene_8d818fe5a7ef
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C>T
T>A
T>C
T>G
Mutation probability
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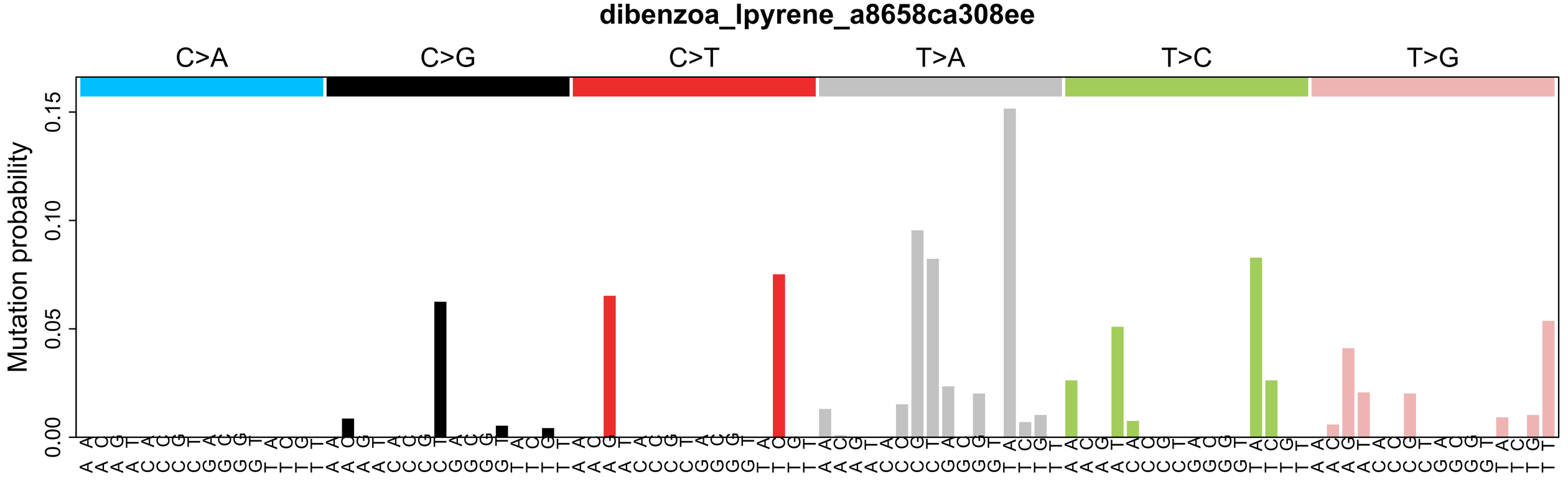
0.05
0.10
0.15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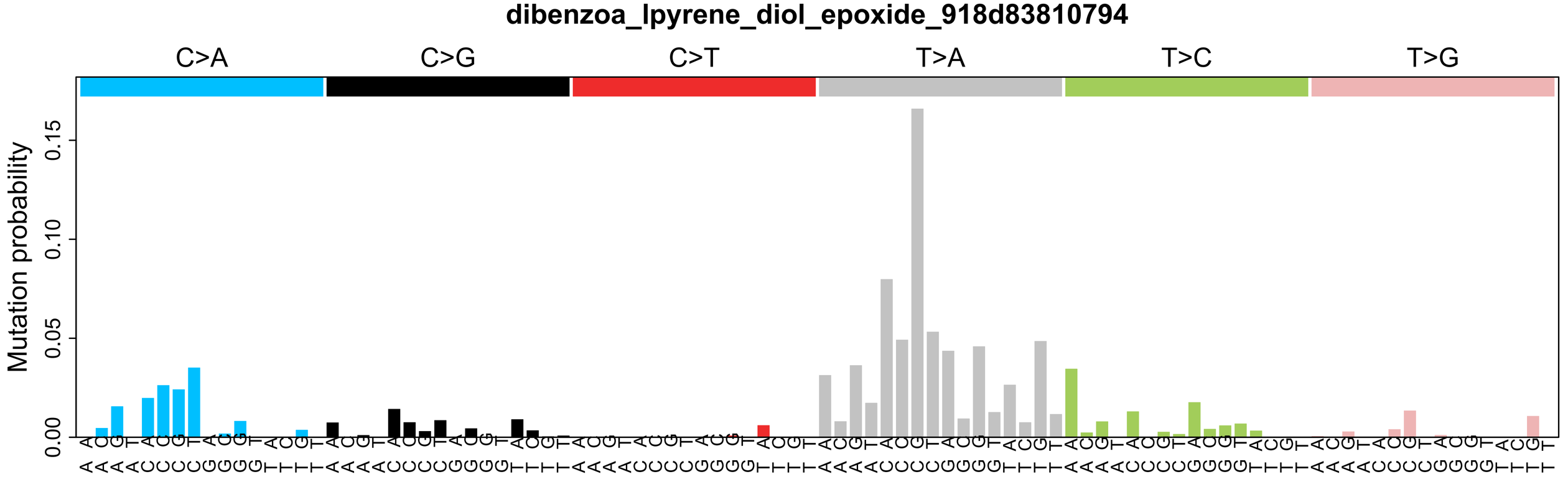
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15


dibenzoa_lpyrene_diol_epoxide_918d83810794
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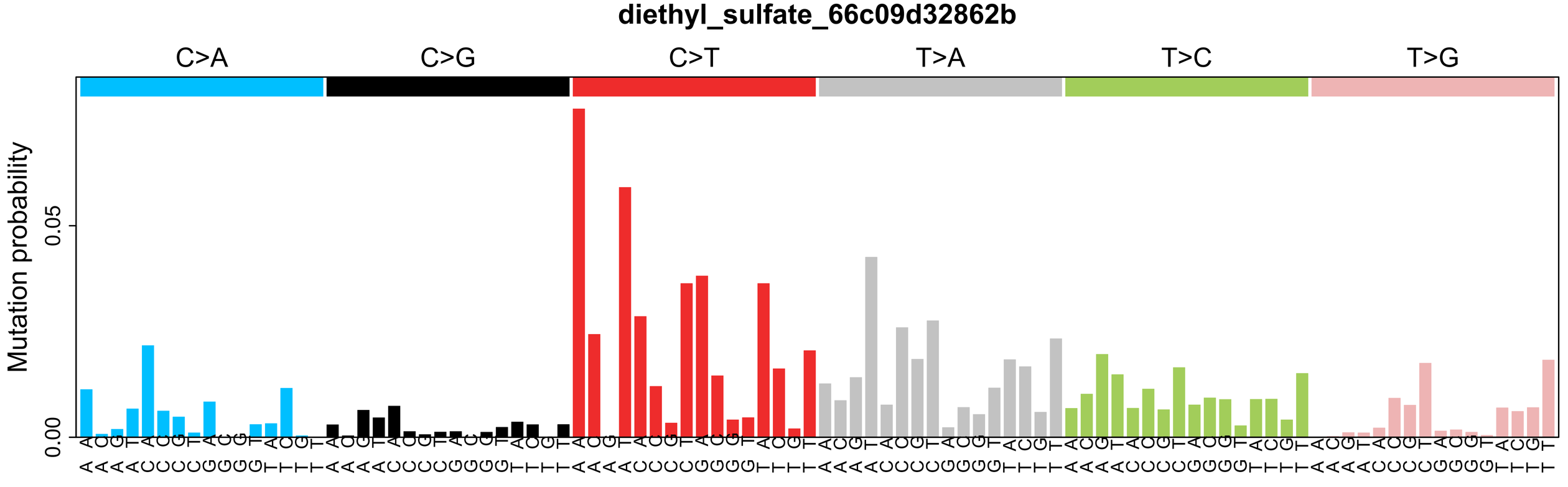
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15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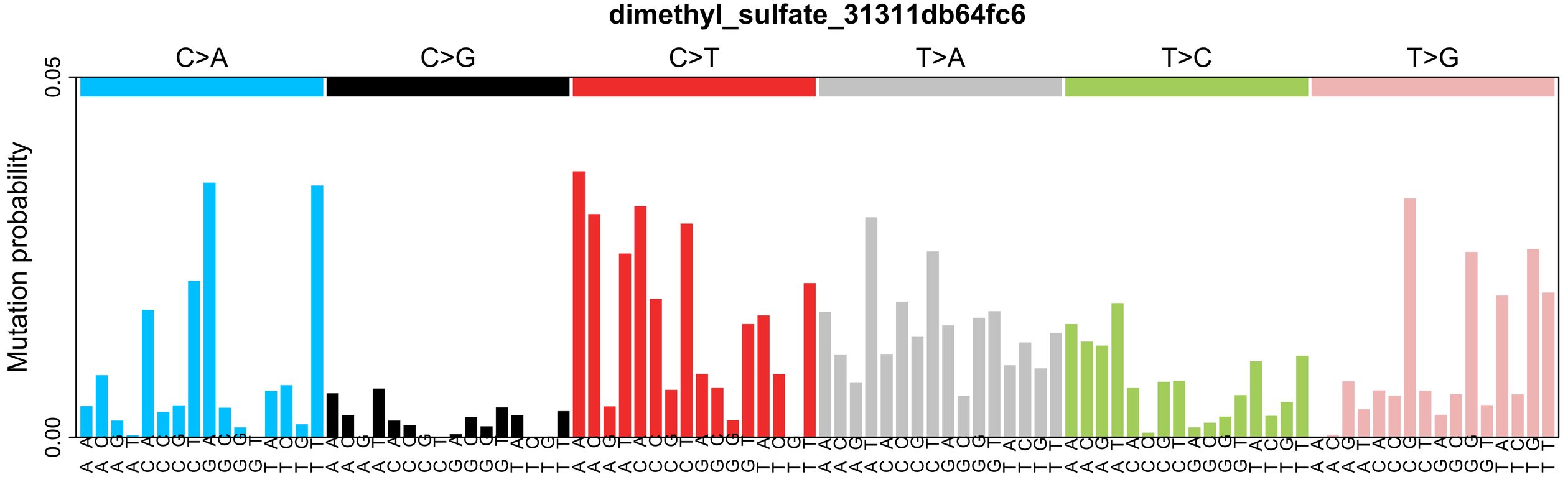
C>T
T>A
T>C
T>G
Mutation probability
0.05
0.00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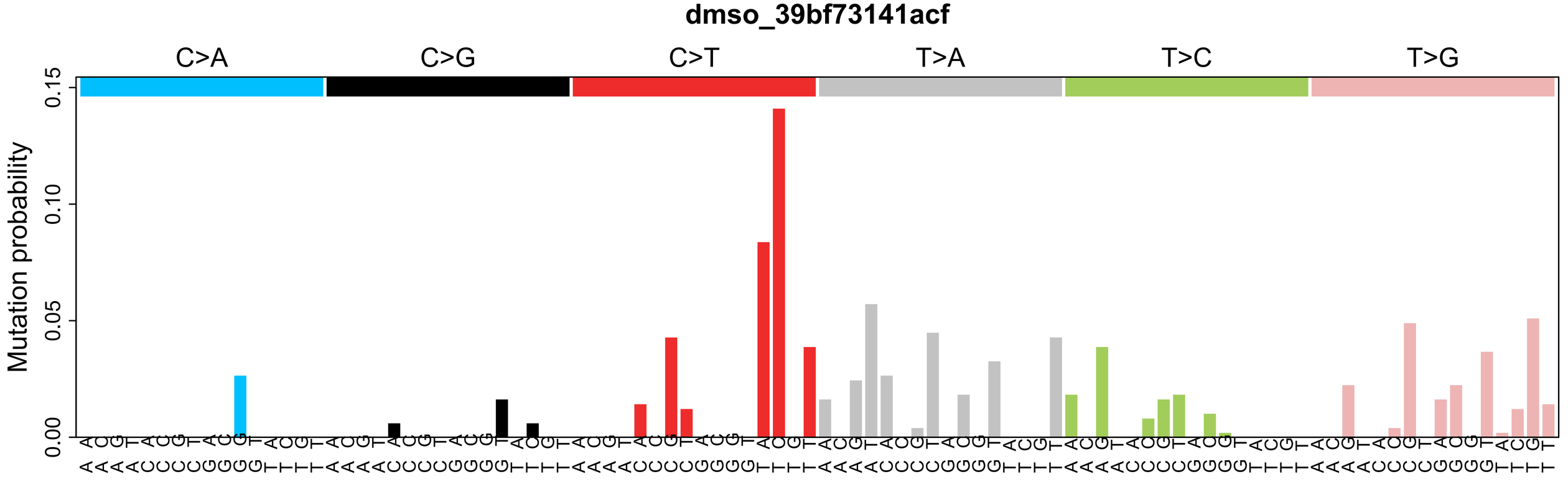
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.05
0.00


dmso_39bf73141acf
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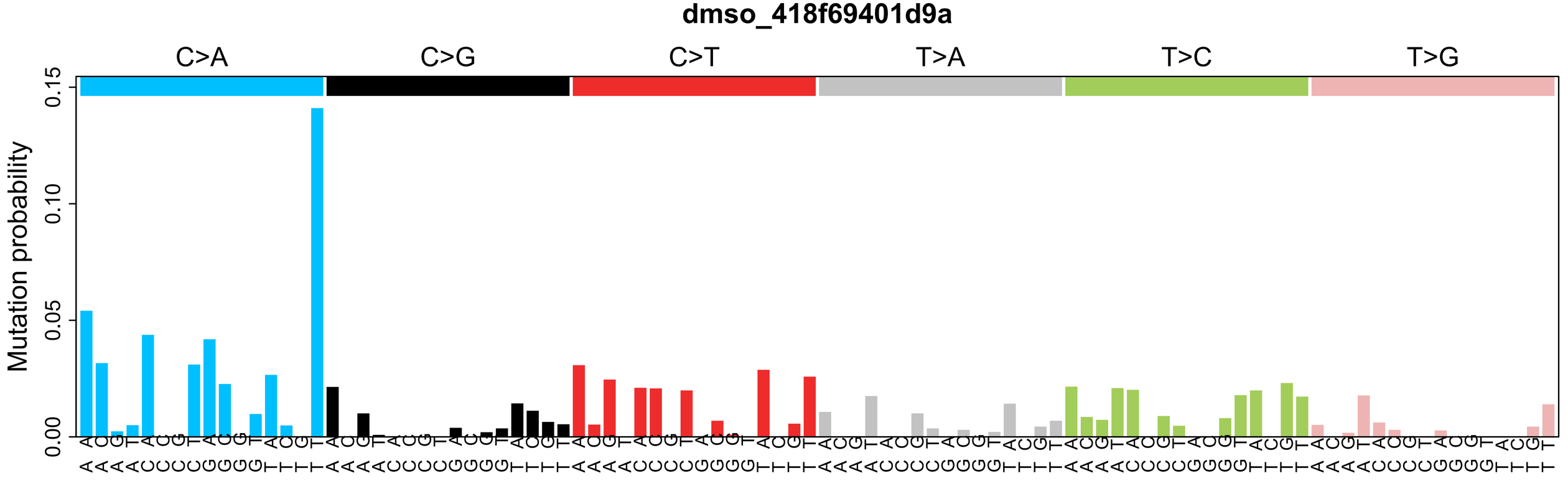
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15


dmso_418f69401d9a
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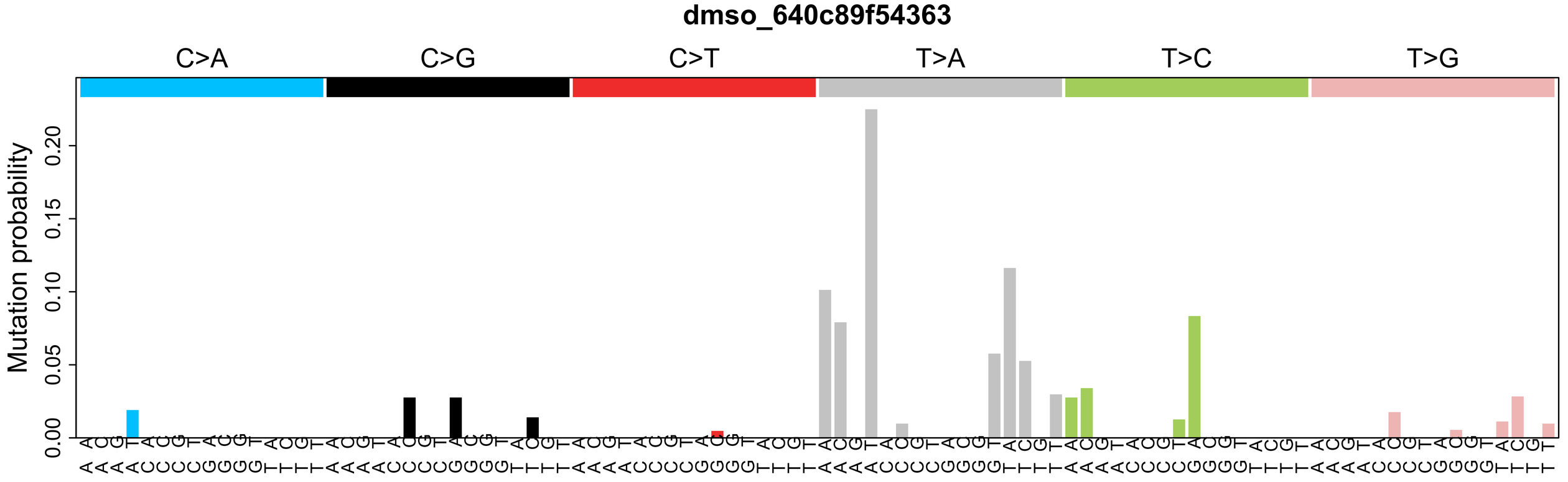
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15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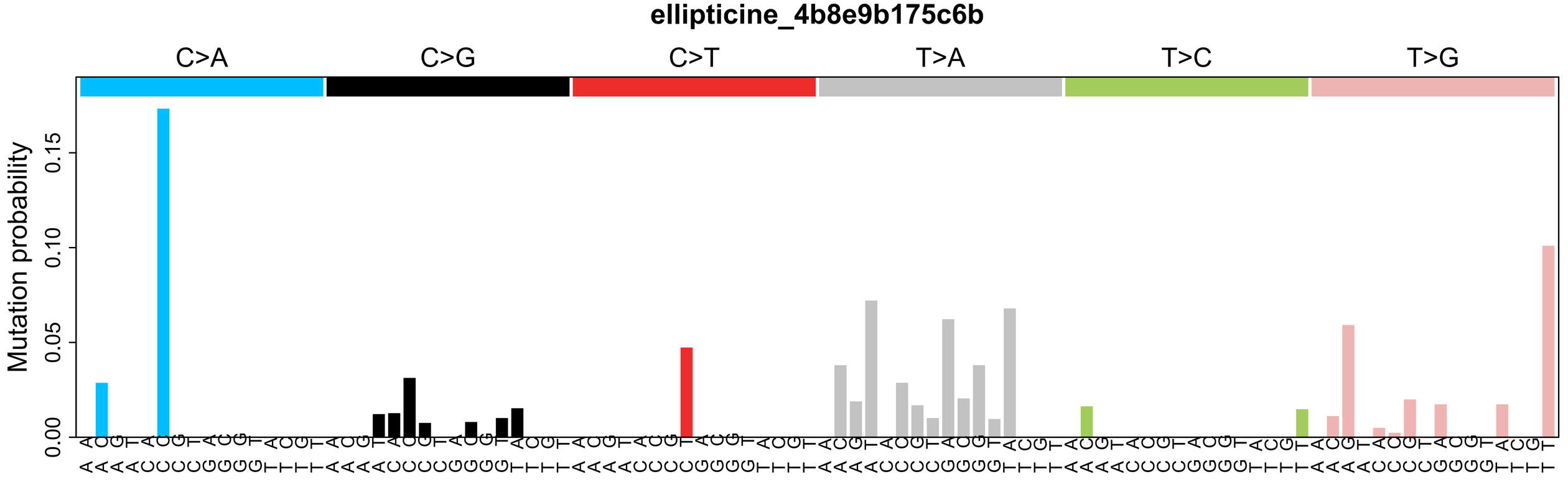
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15
0.20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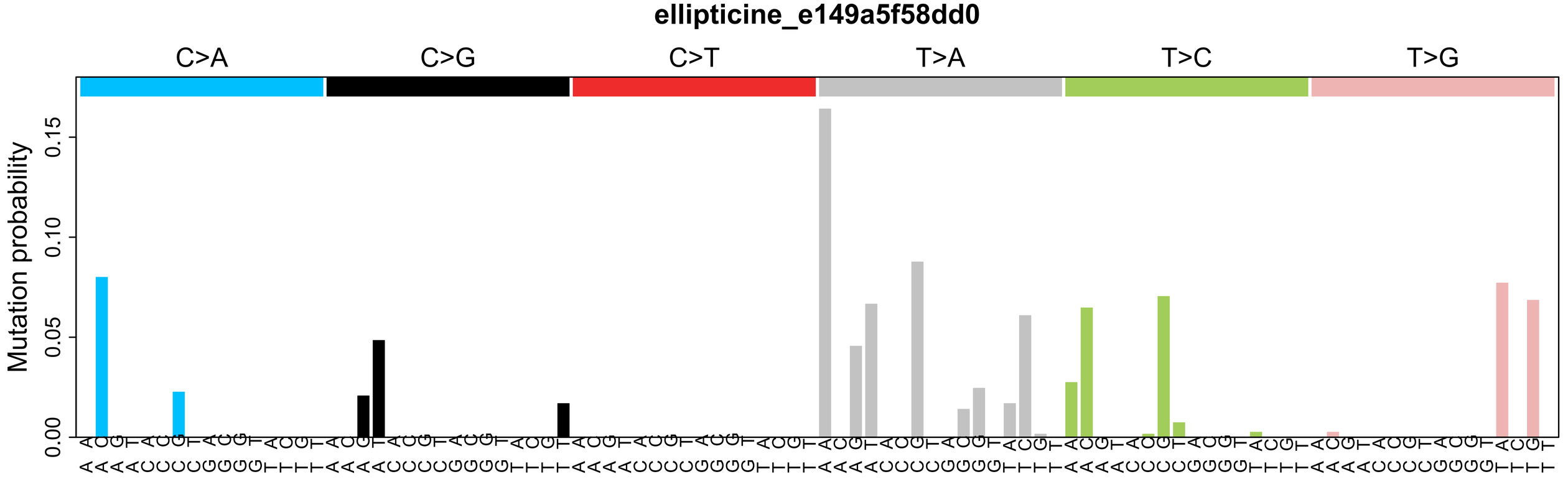
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15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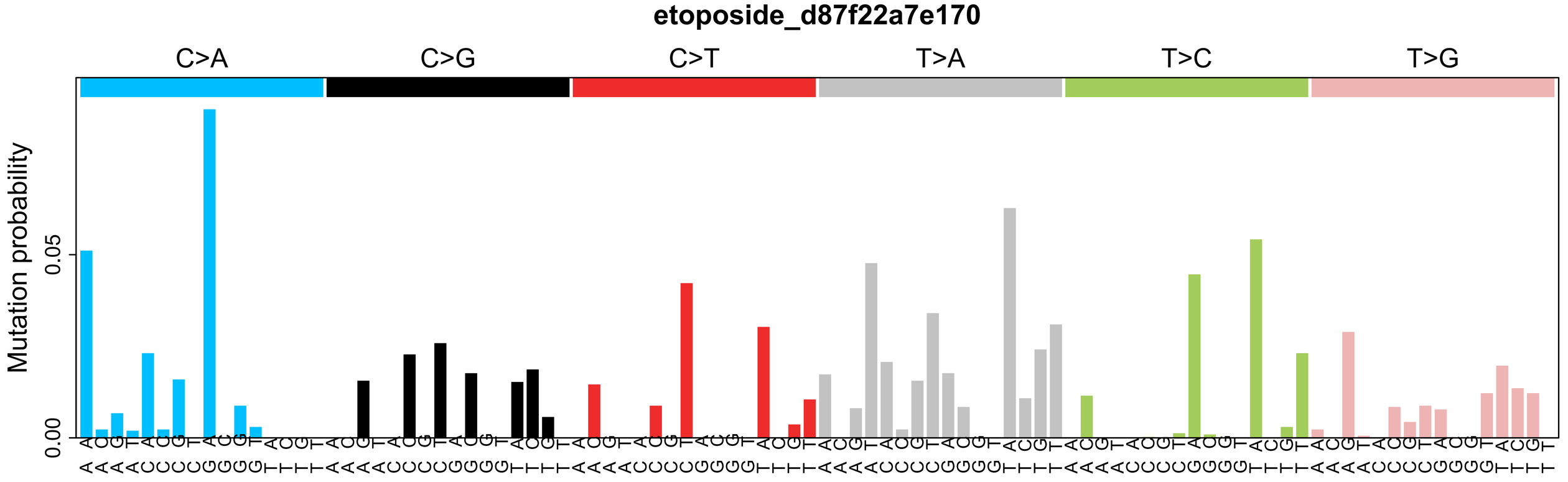
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15


etoposide_d87f22a7e170
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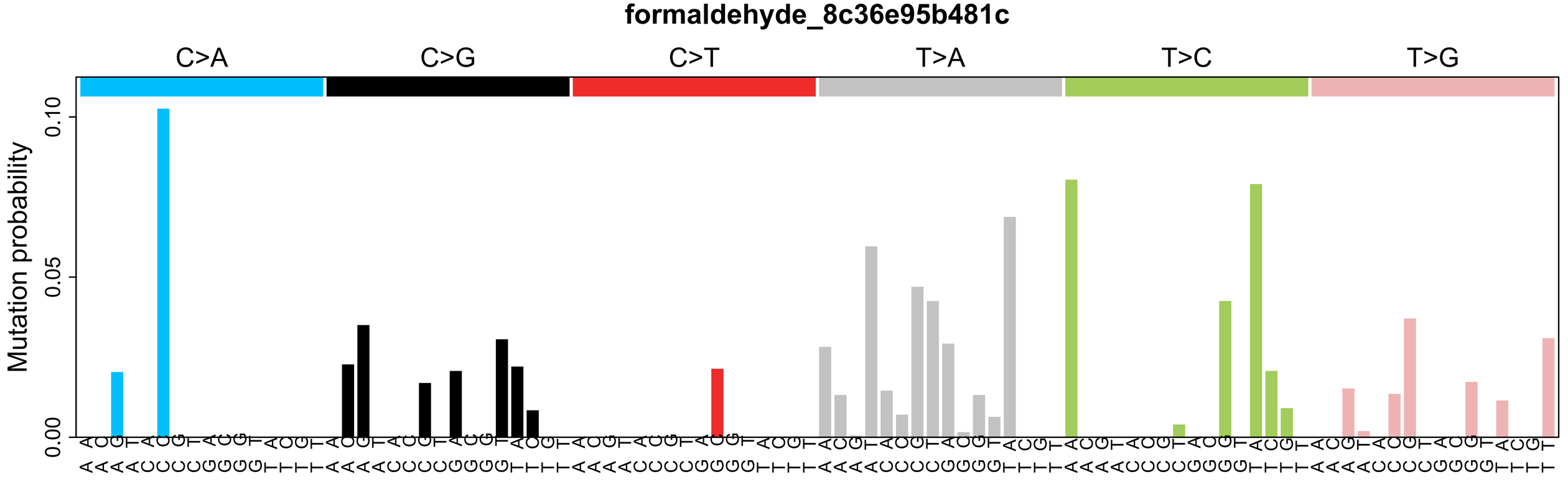
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05


formaldehyde_8c36e95b481c
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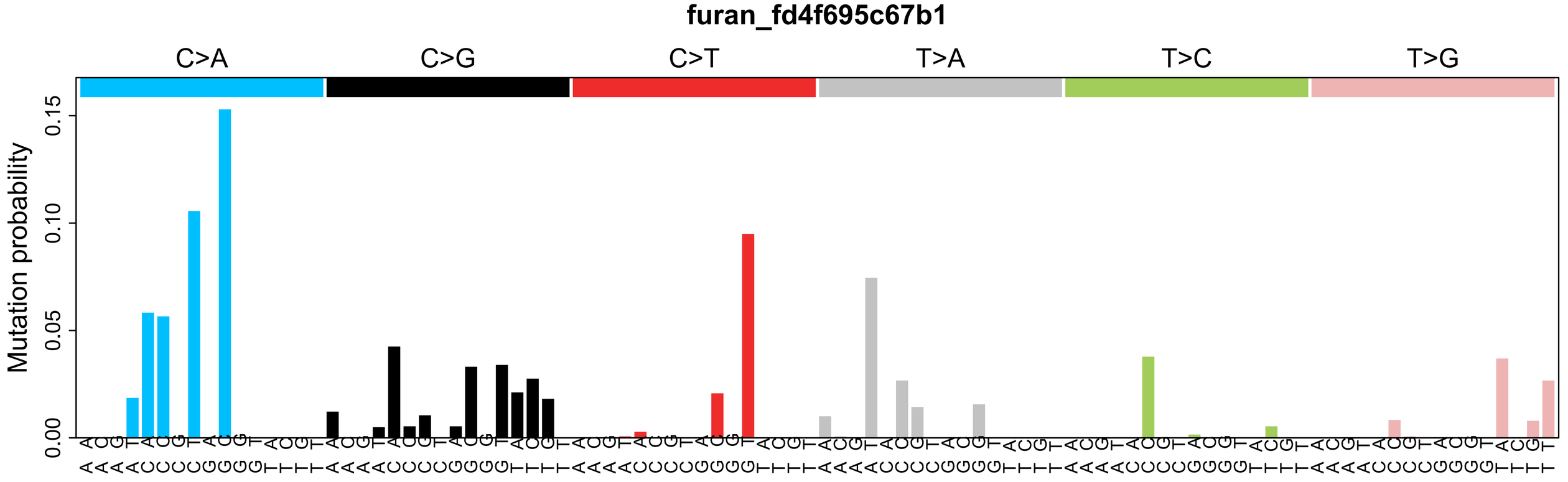
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


furan_fd4f695c67b1
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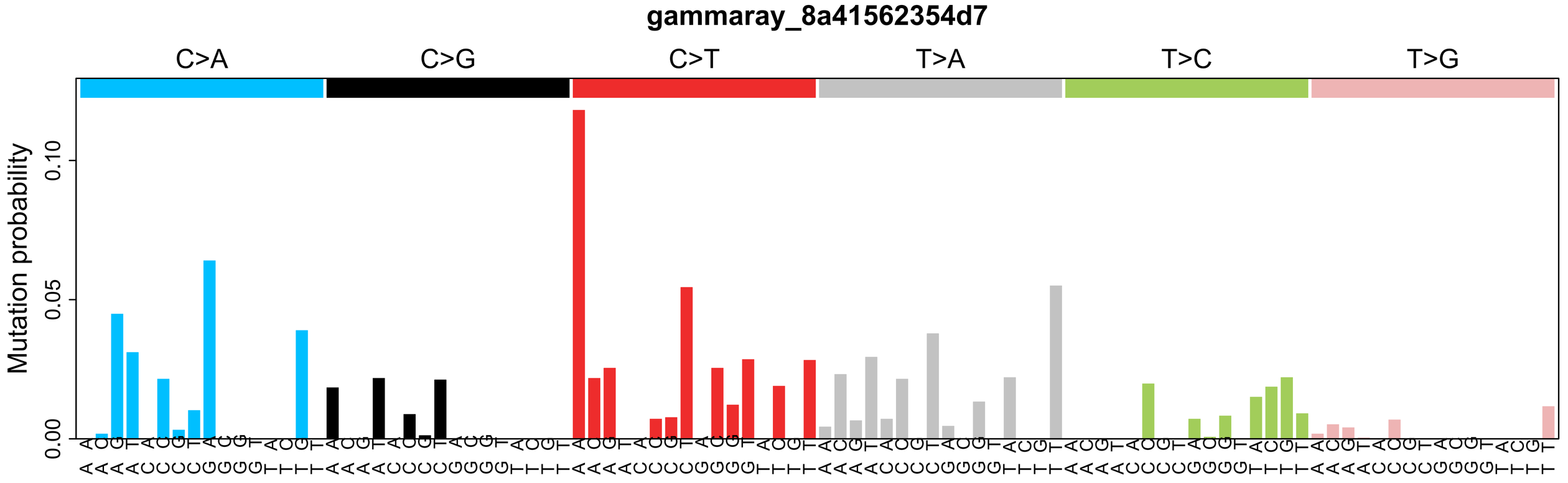
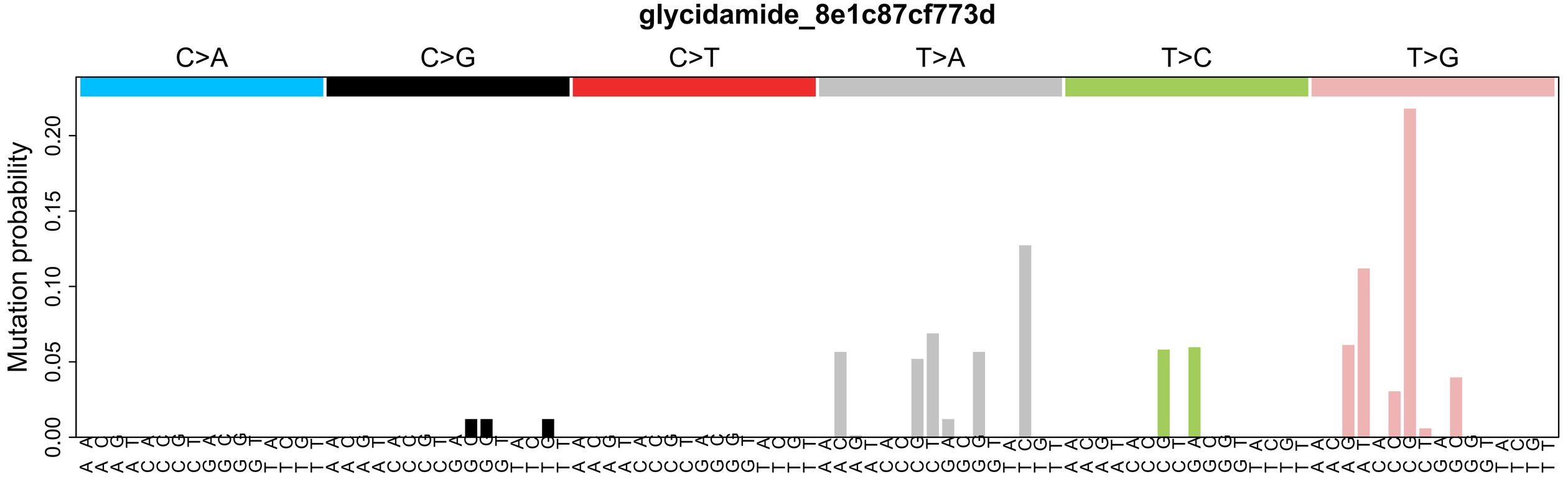
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15


gammaray_8a41562354d7
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


glycidamide_8e1c87cf773d
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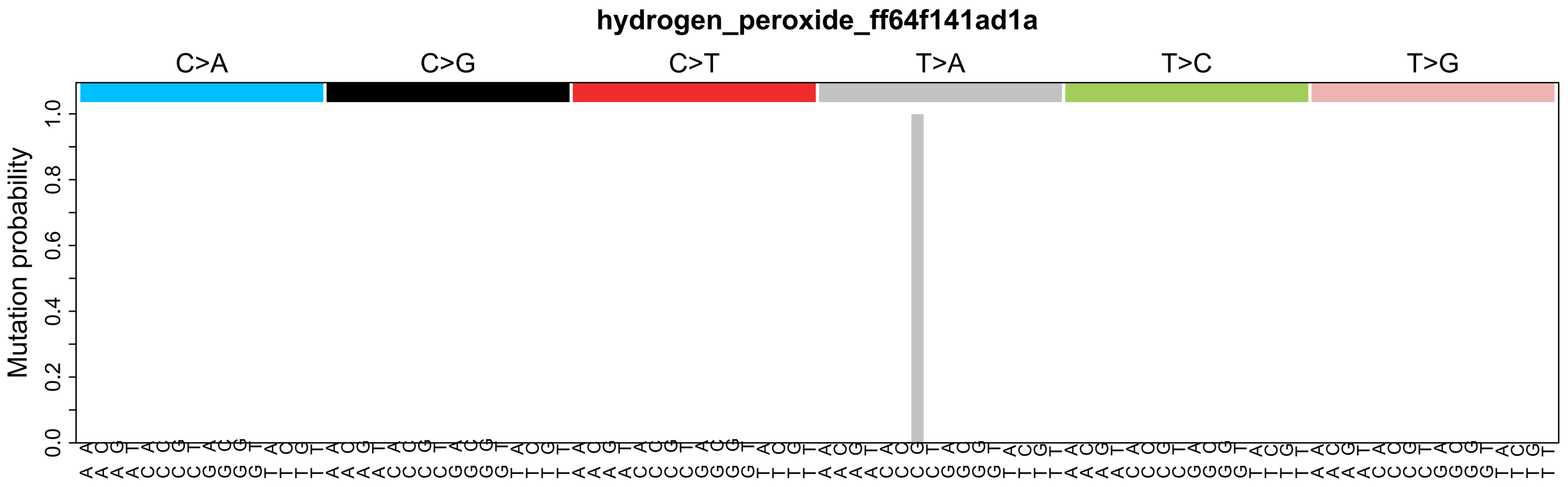
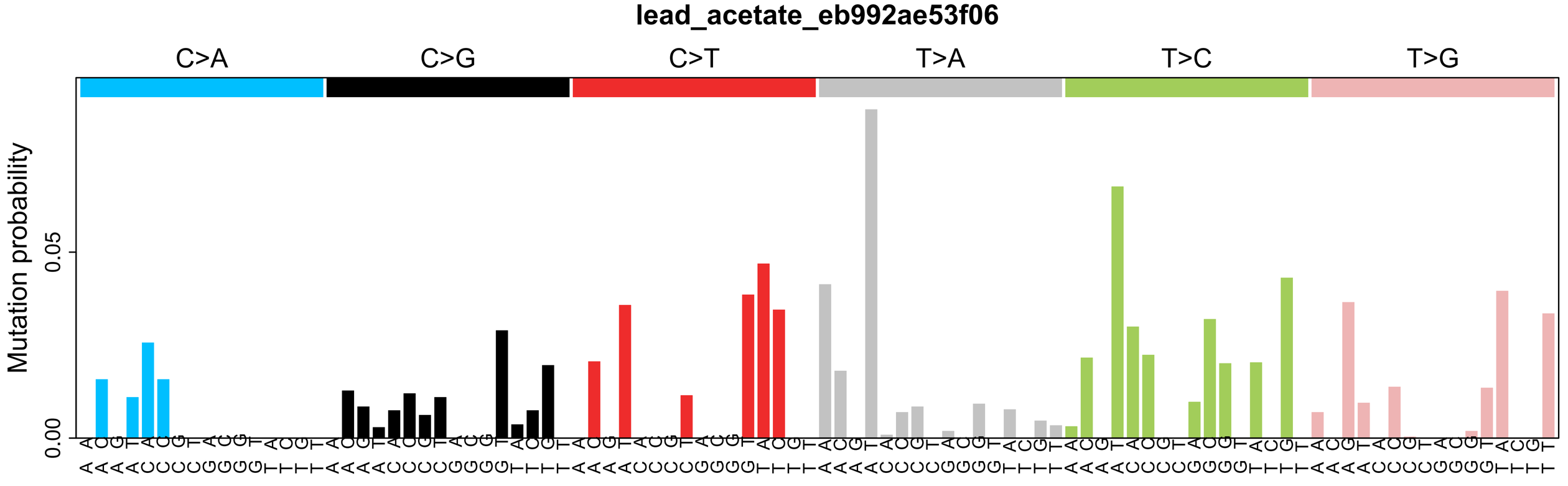
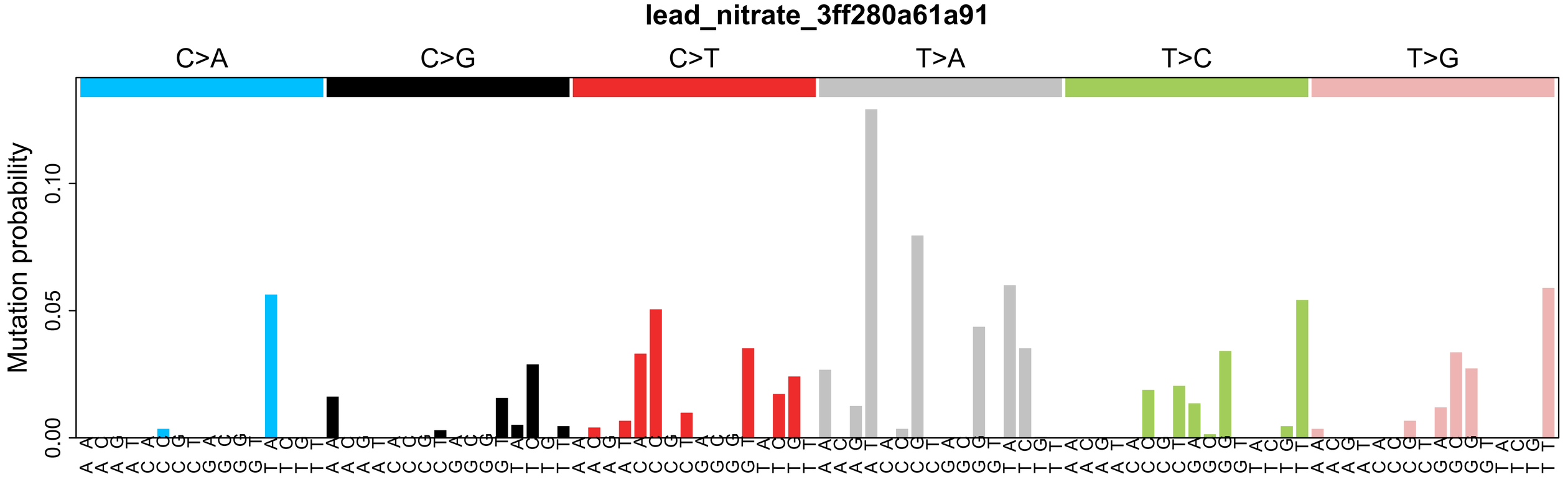
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15
0.20


hydrogen_peroxide_ff64f141ad1a
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.2
0.4
0.6
0.8
1.0


lead_acetate_eb992ae53f06
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05


lead_nitrate_3ff280a61a91
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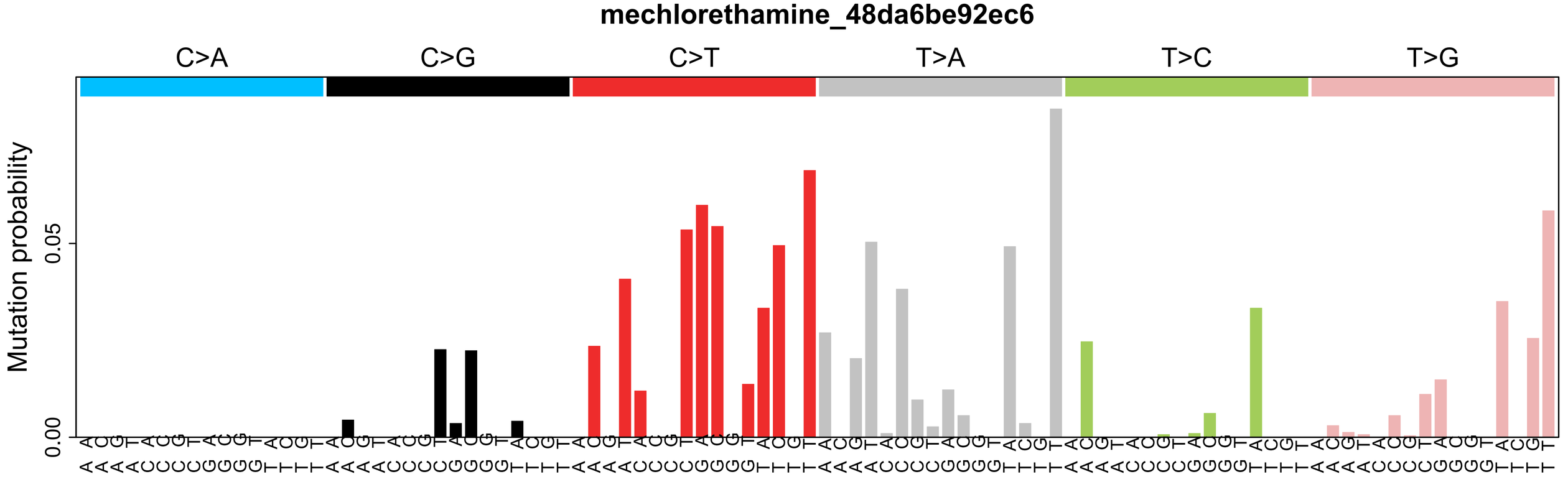
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


mechlorethamine_48da6be92ec6
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
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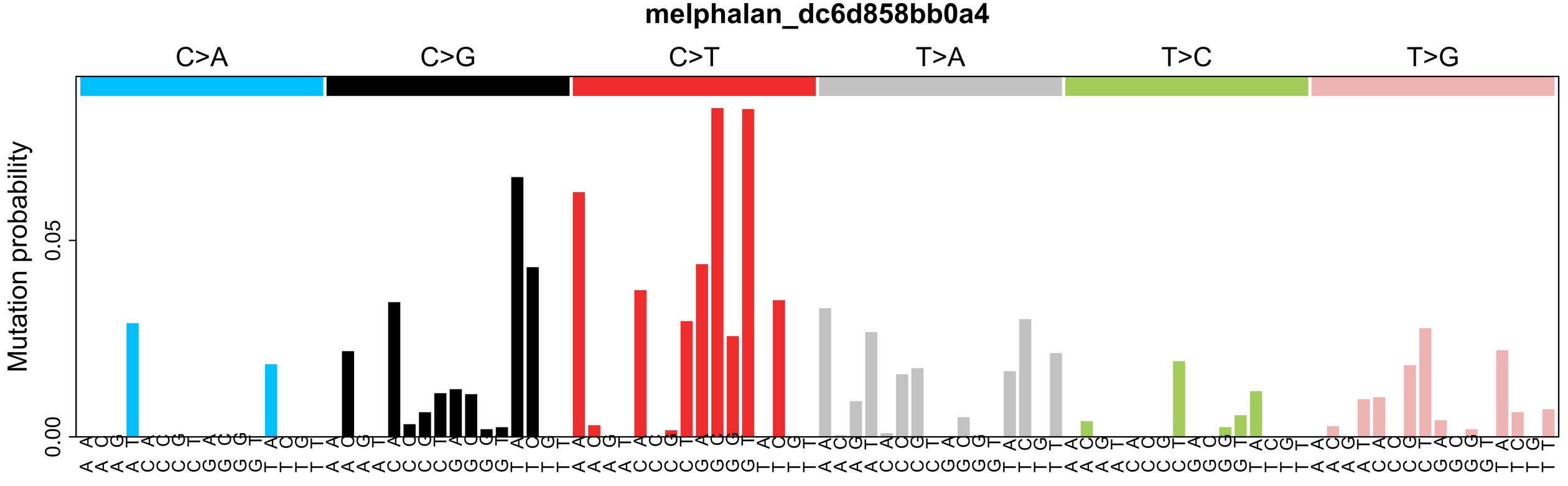
melphalan_dc6d858bb0a4
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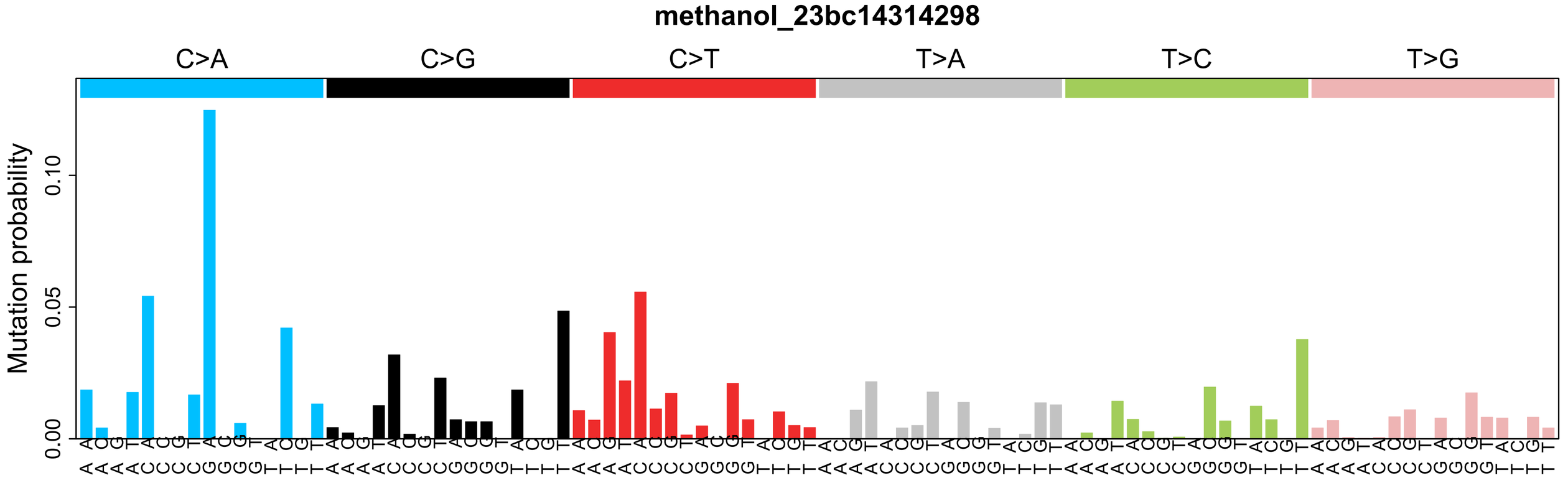
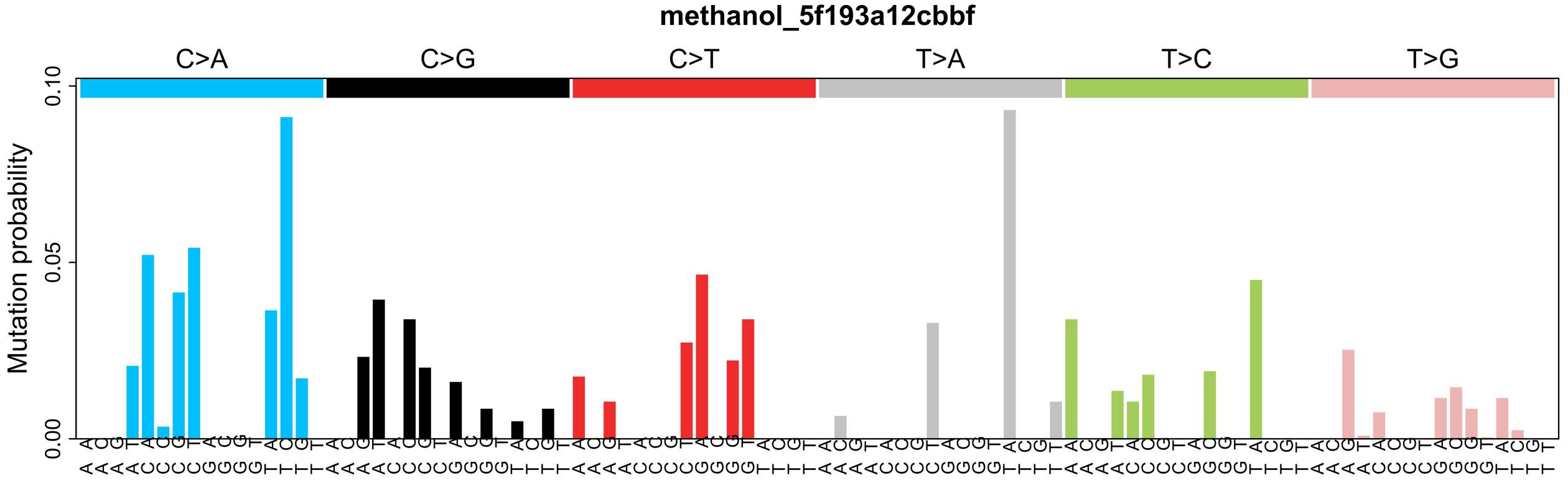
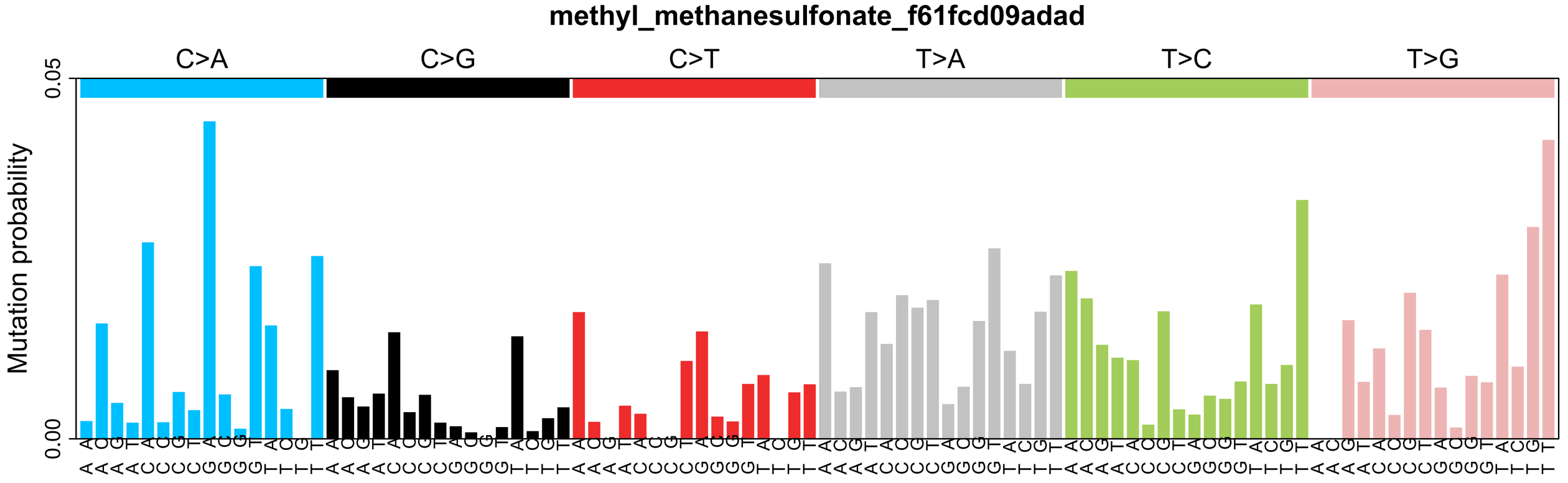
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05


methanol_23bc14314298
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


methanol_5f193a12cbbf
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


methyl_methanesulfonate_f61fcd09adad
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
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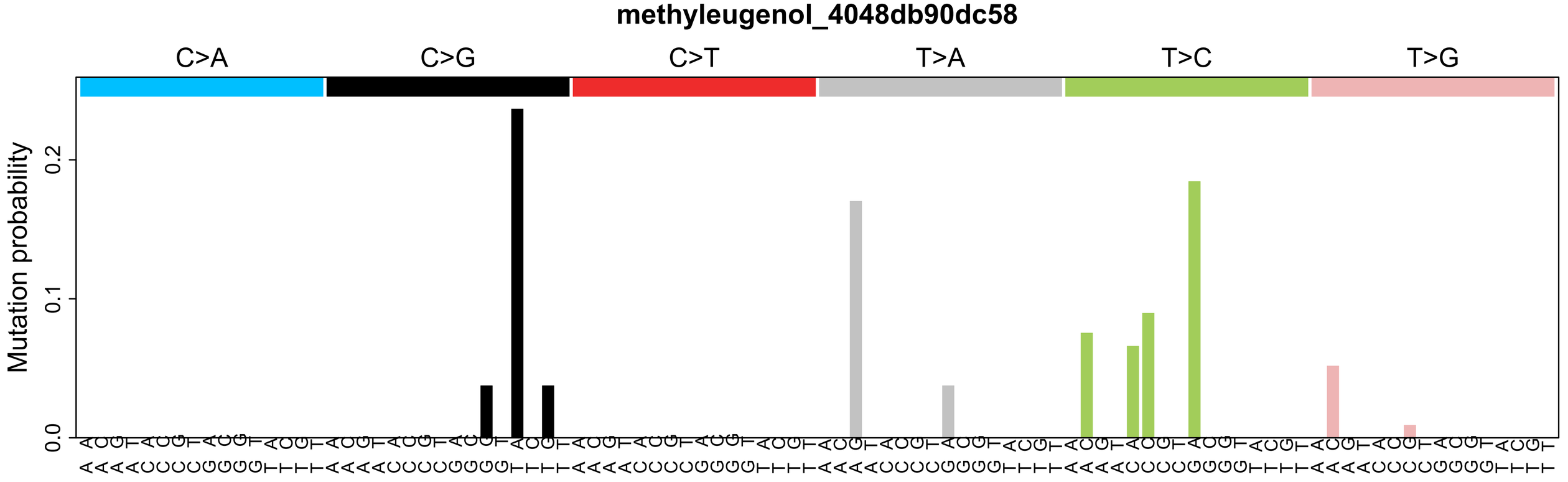
methyleugenol_4048db90dc58
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
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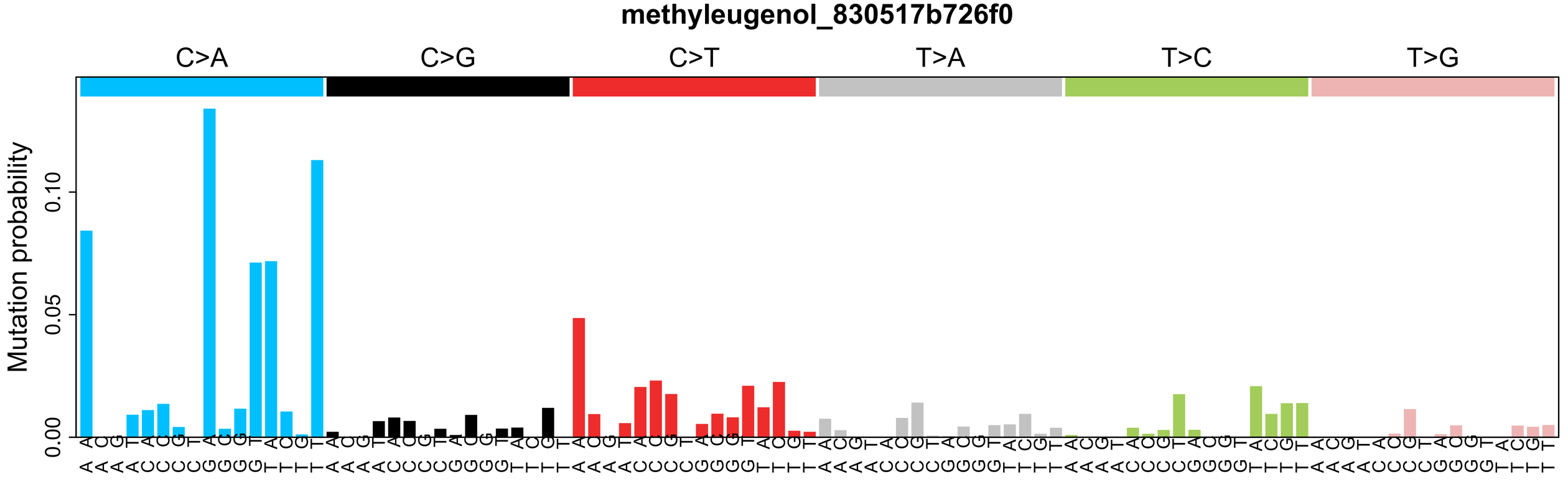
0.1
0.2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C>G
C>T
T>A
T>C
T>G
Mutation probability
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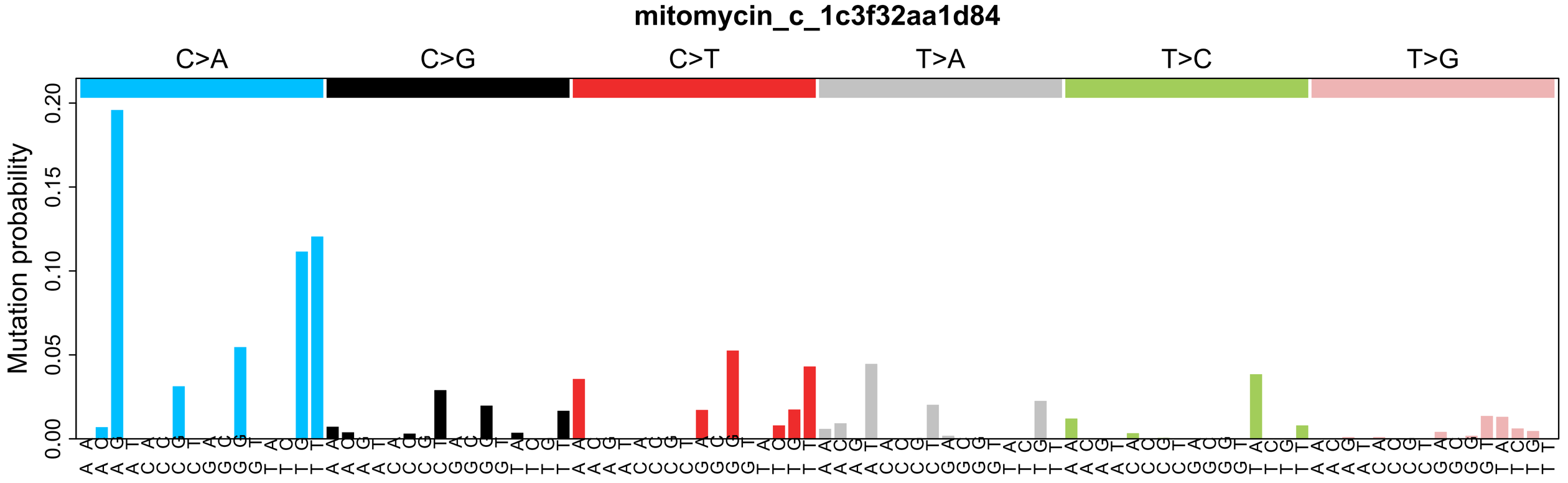
0.05
0.10


mitomycin_c_1c3f32aa1d84
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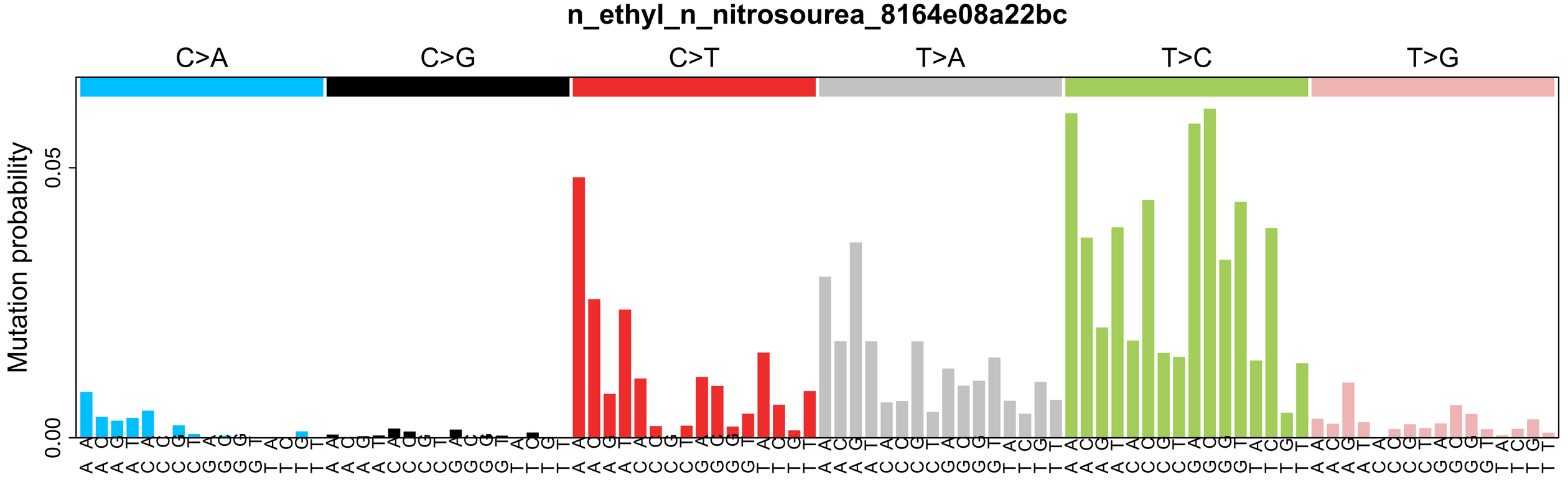
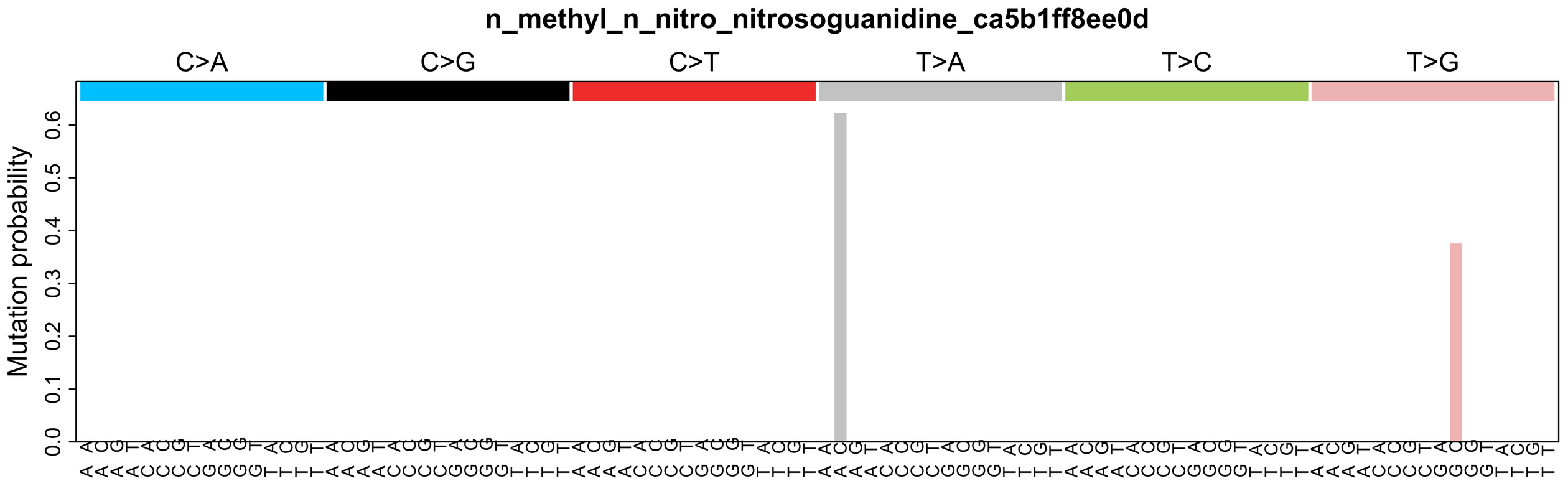
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15
0.20


n_ethyl_n_nitrosourea_8164e08a22bc
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.05
0.00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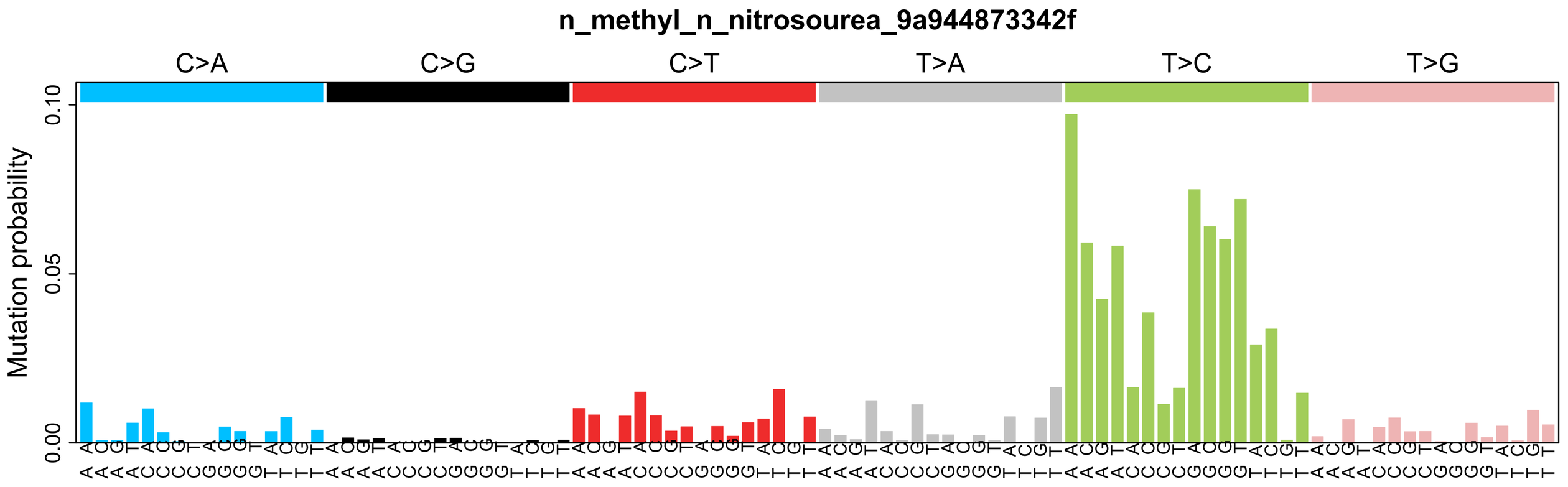
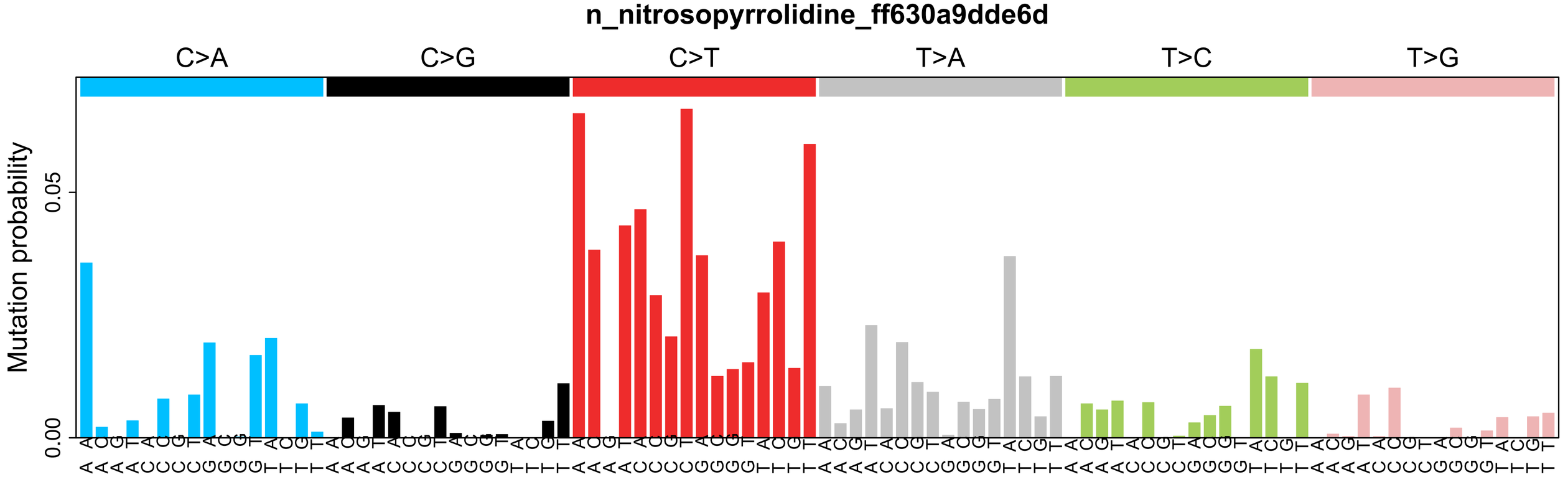
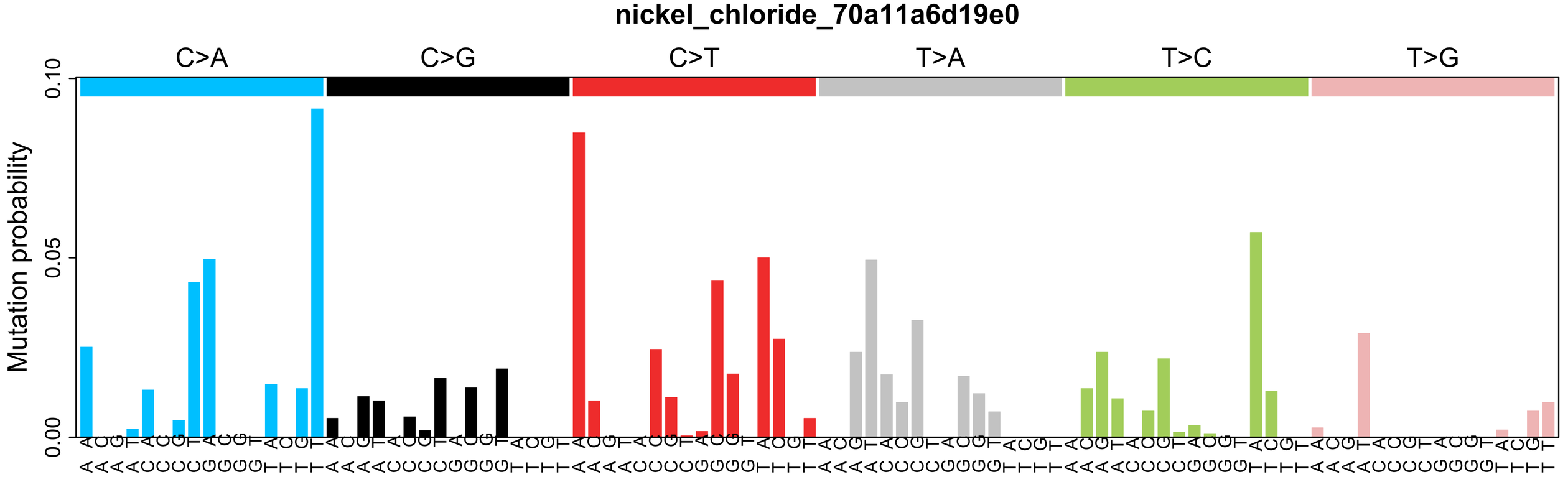
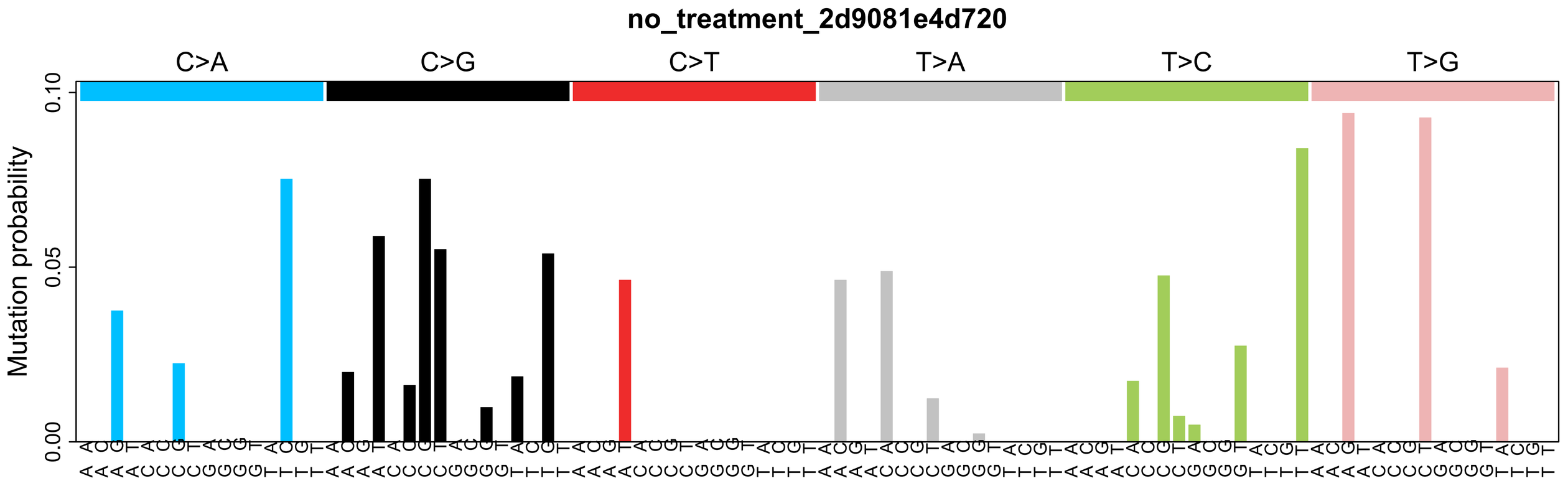
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2
0.3
0.4
0.5
0.6


n_methyl_n_nitrosourea_9a944873342f
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


n_nitrosopyrrolidine_ff630a9dde6d
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.05
0.00


nickel_chloride_70a11a6d19e0
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


no_treatment_2d9081e4d720
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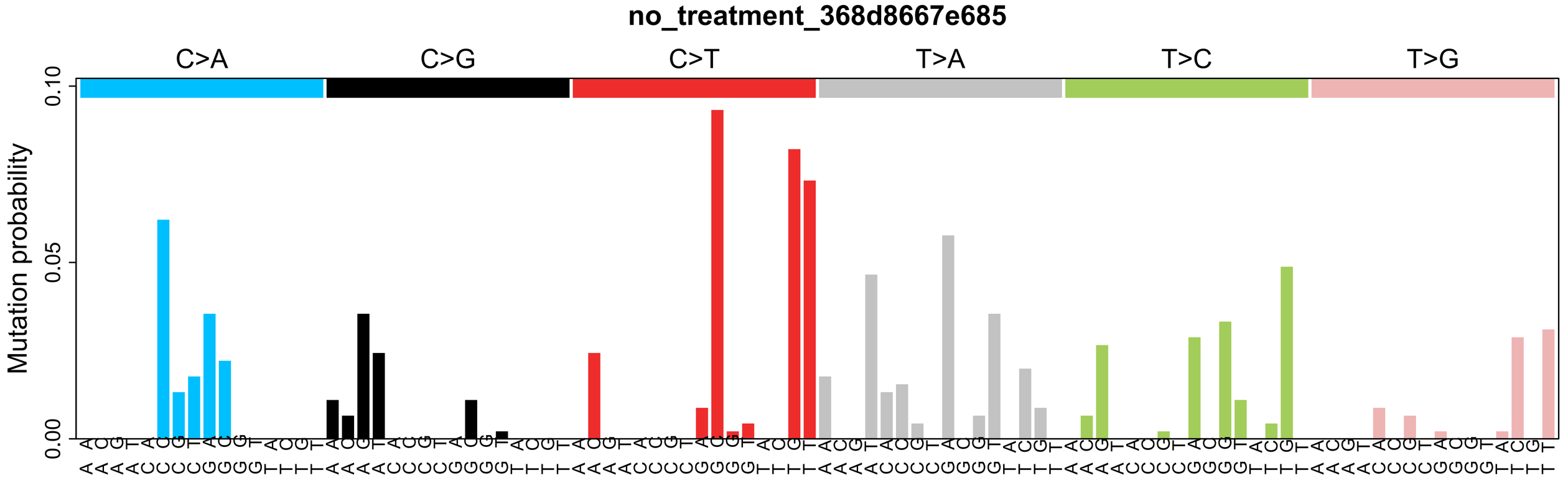
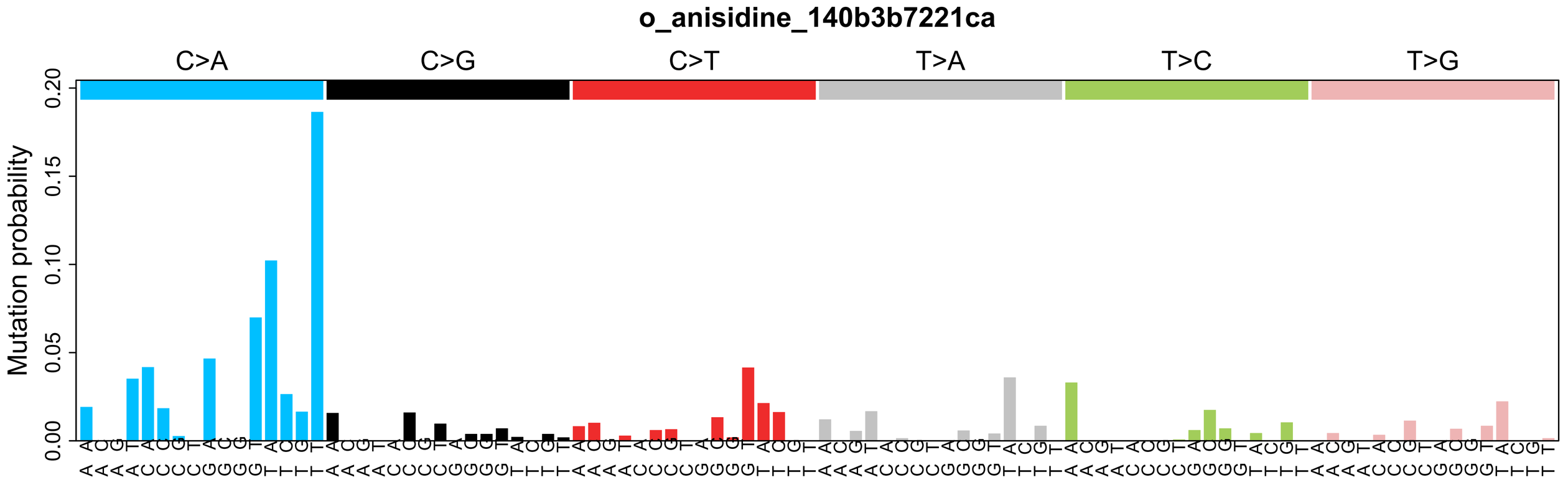
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


no_treatment_368d8667e685
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


o_anisidine_140b3b7221ca
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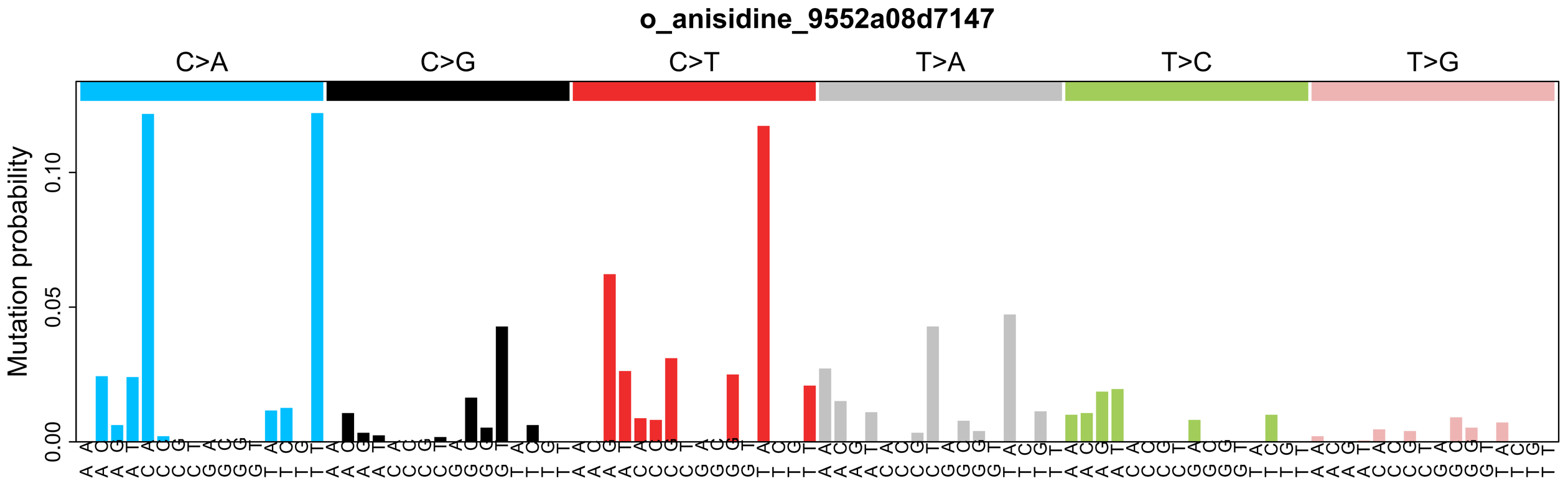
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15
0.20


o_anisidine_9552a08d7147
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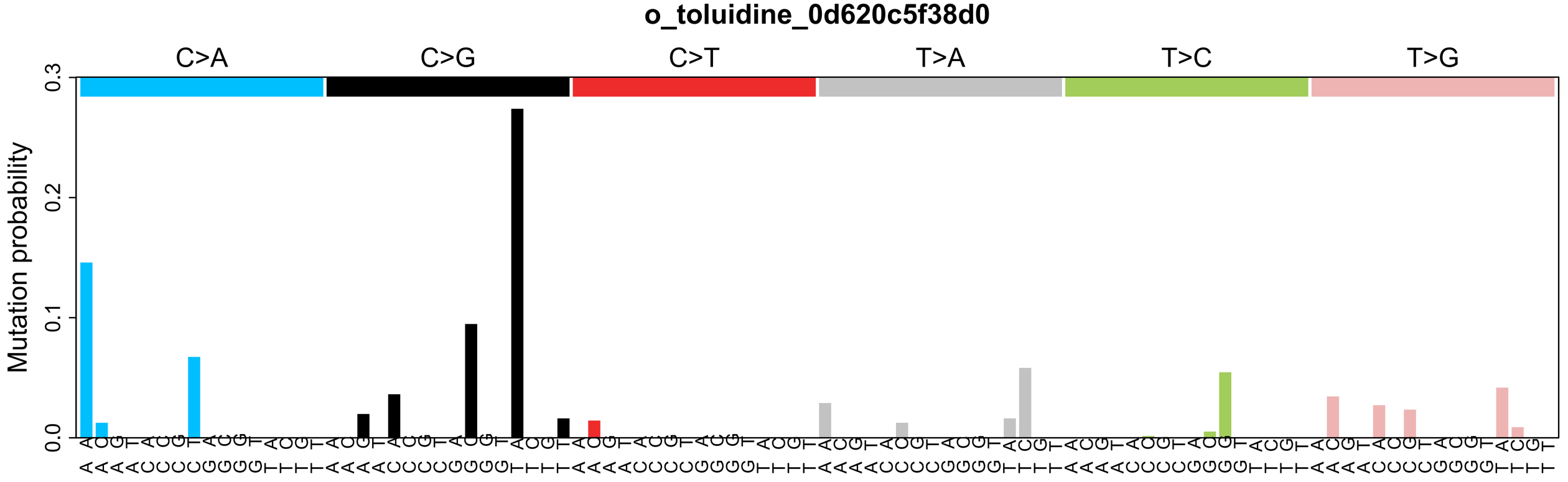
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


o_toluidine_0d620c5f38d0
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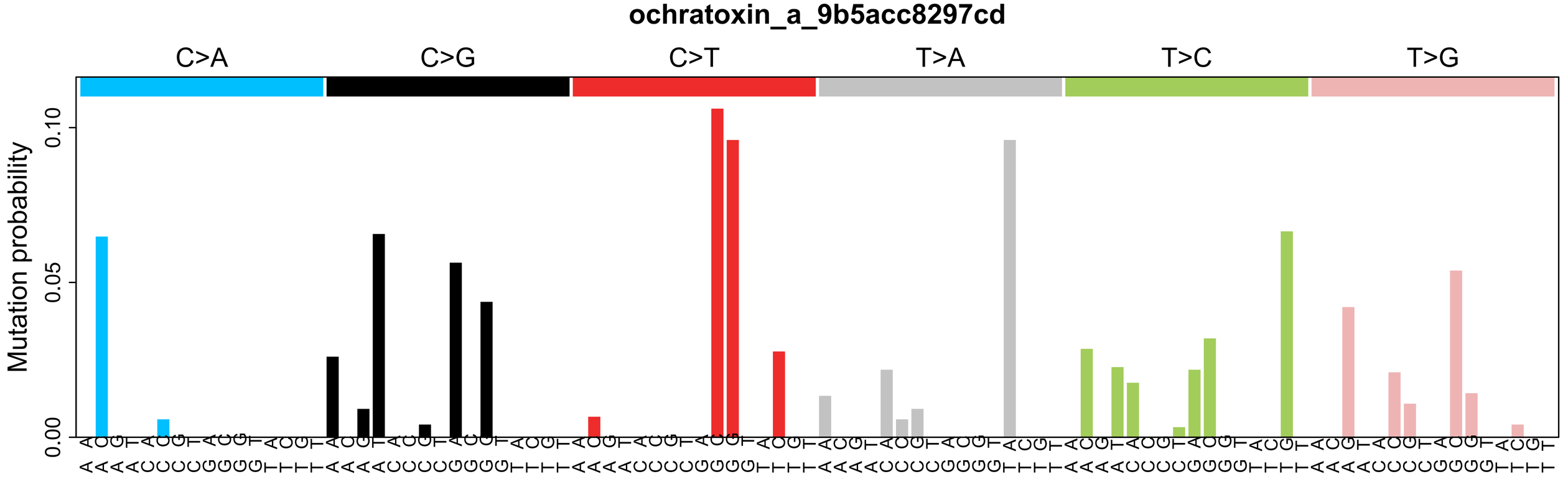
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2
0.3


ochratoxin_a_9b5acc8297cd
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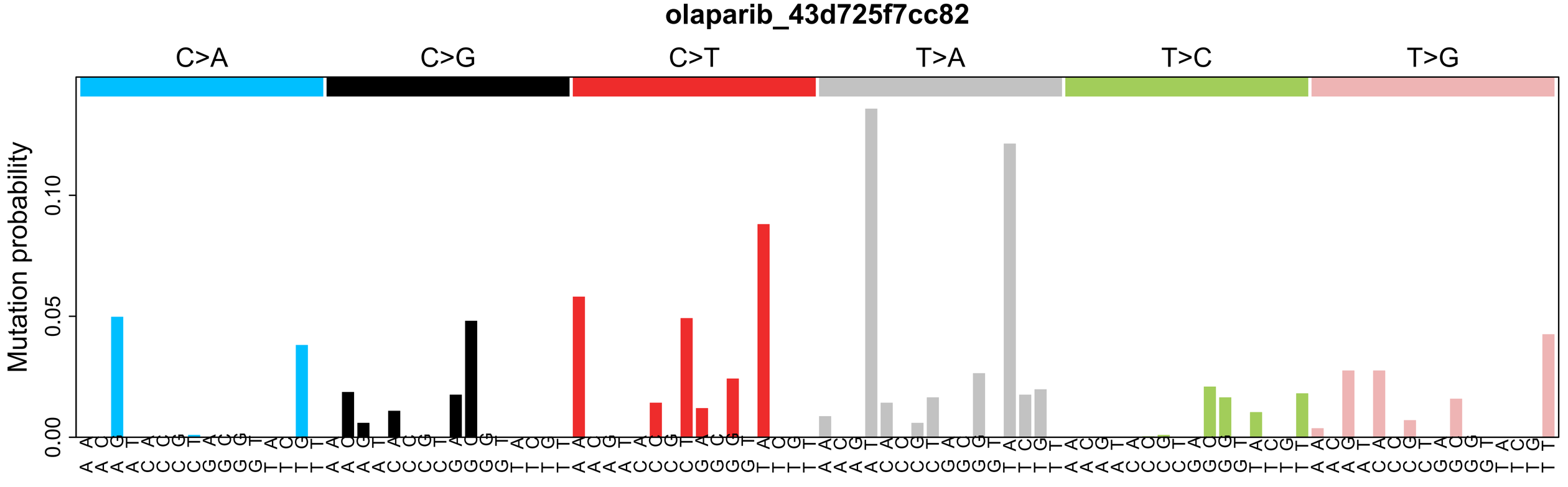
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


olaparib_43d725f7cc82
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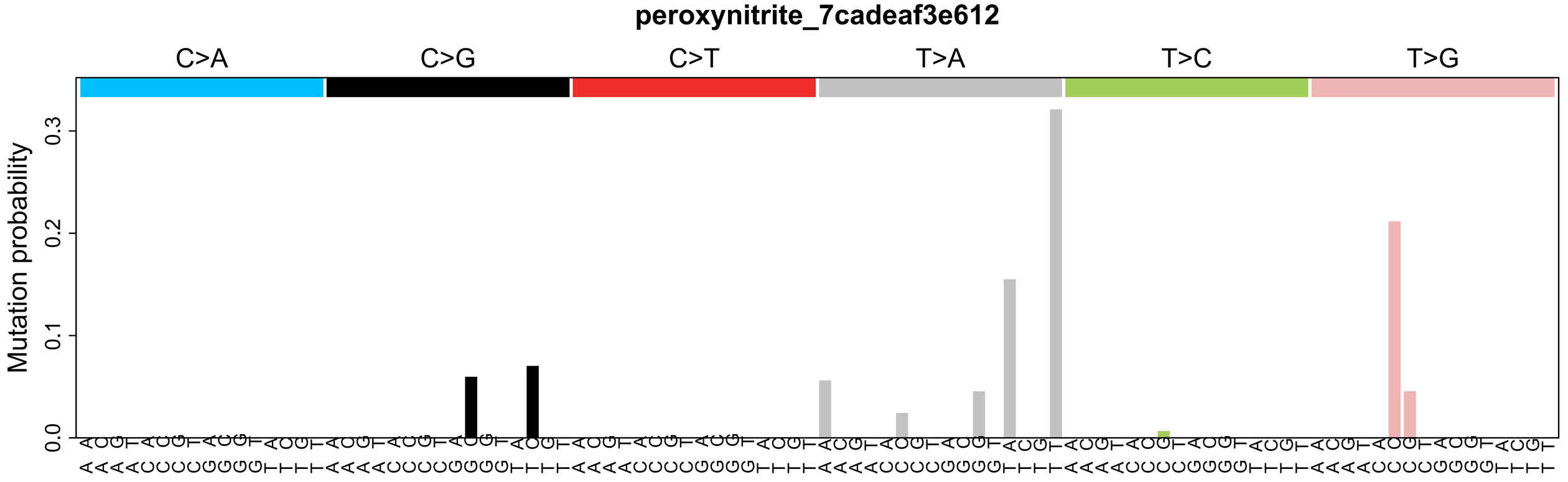
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


peroxynitrite_7cadeaf3e612
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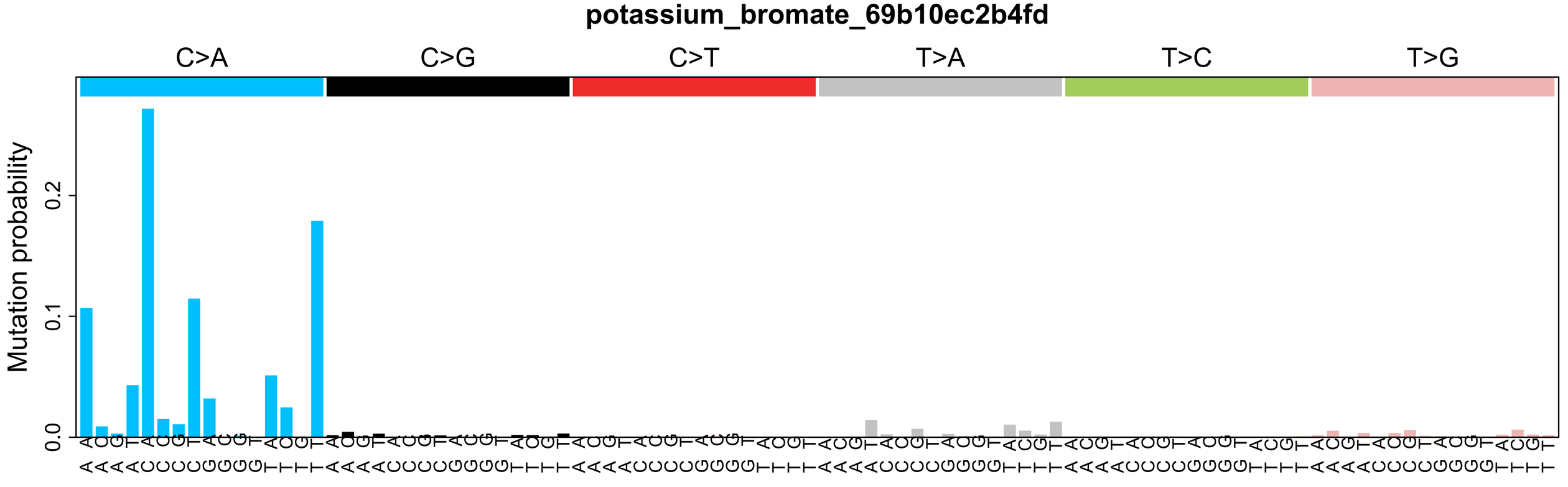
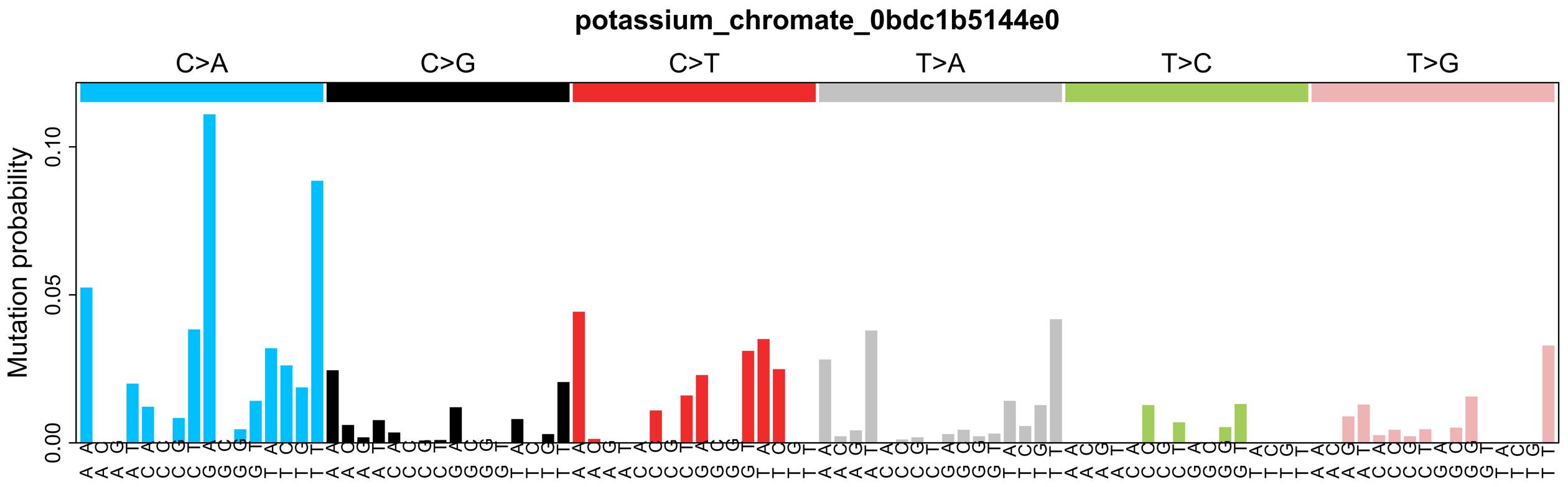
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2
0.3


potassium_bromate_69b10ec2b4fd
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2


potassium_chromate_0bdc1b5144e0
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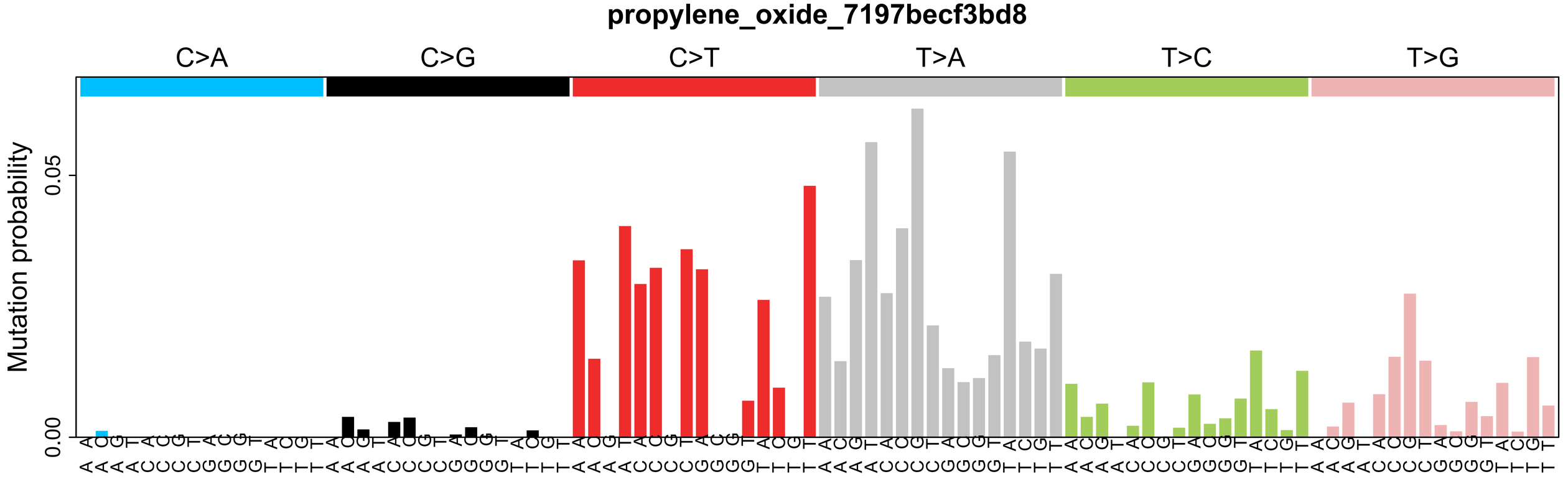
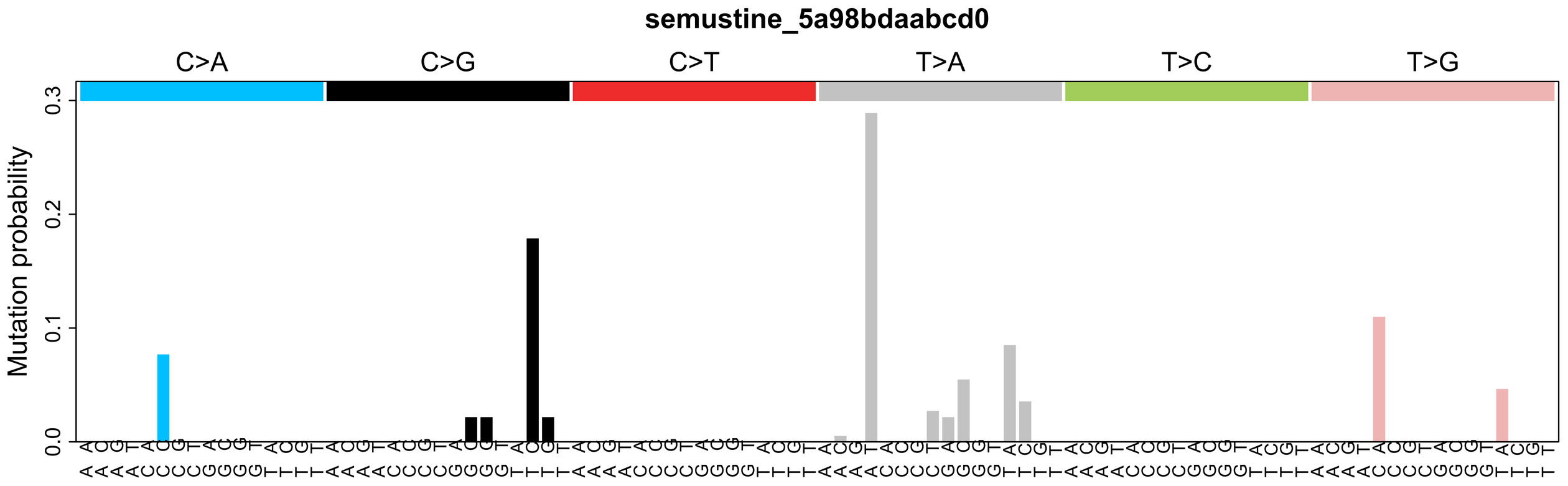
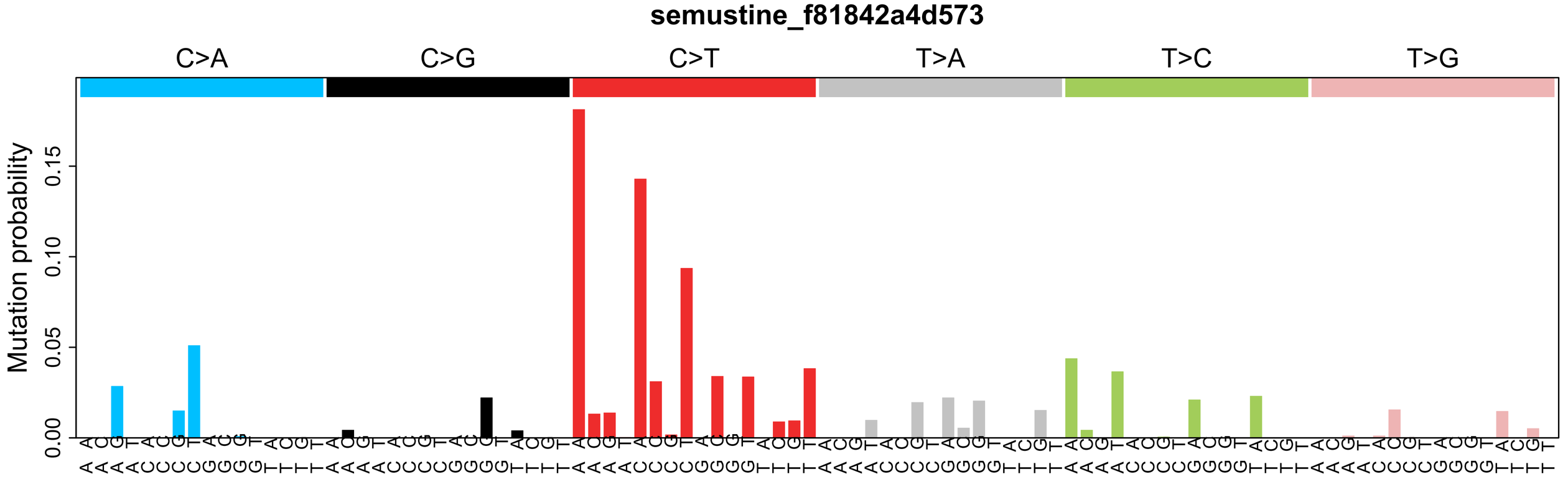
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


propylene_oxide_7197becf3bd8
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.05
0.00


semustine_5a98bdaabcd0
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2
0.3


semustine_f81842a4d573
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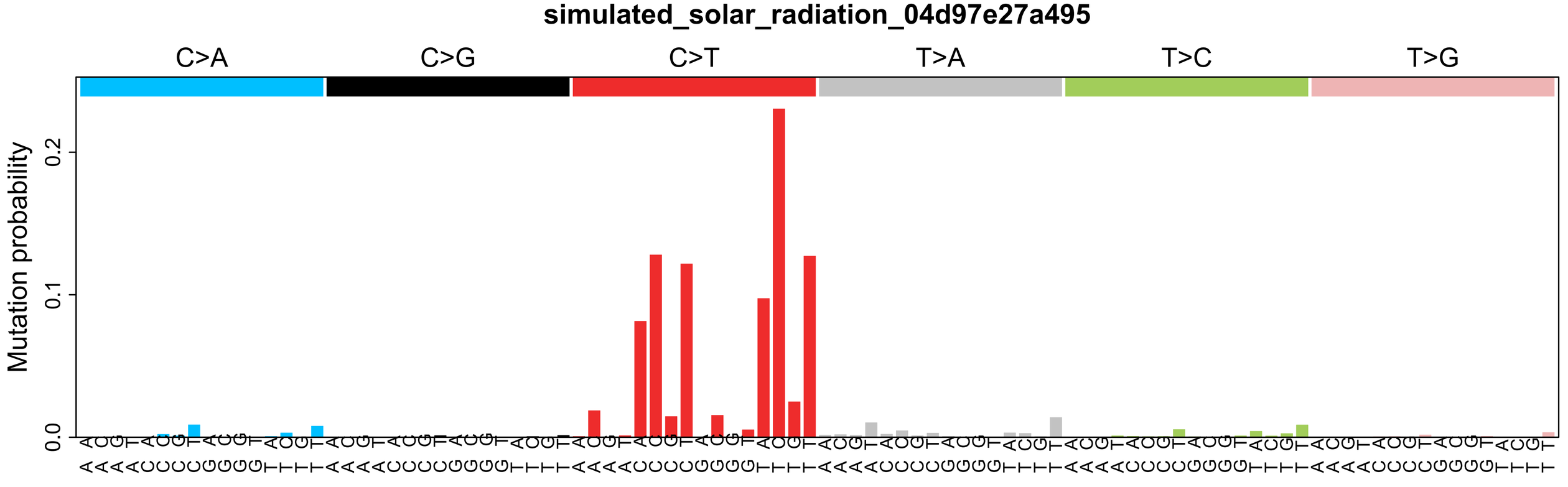
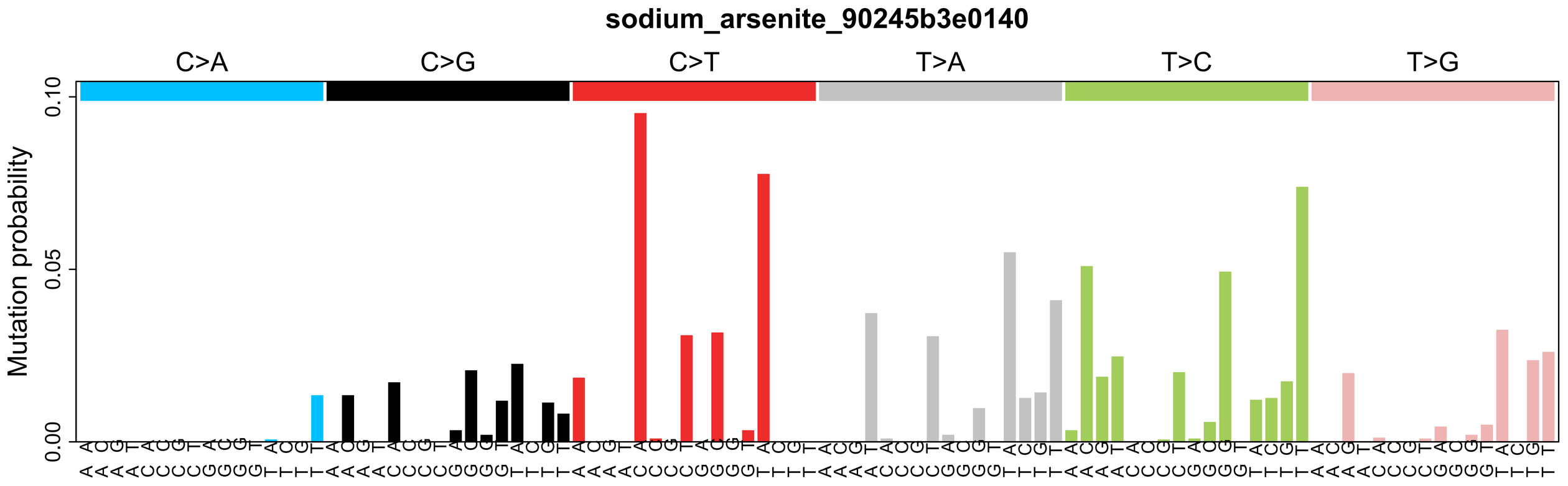
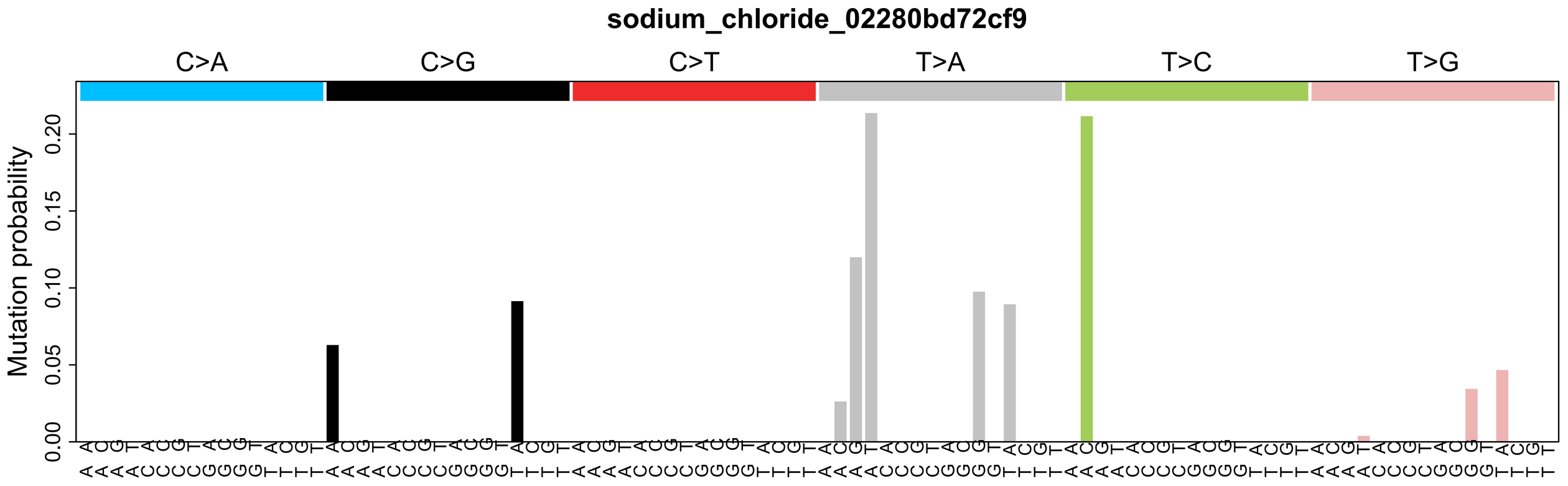
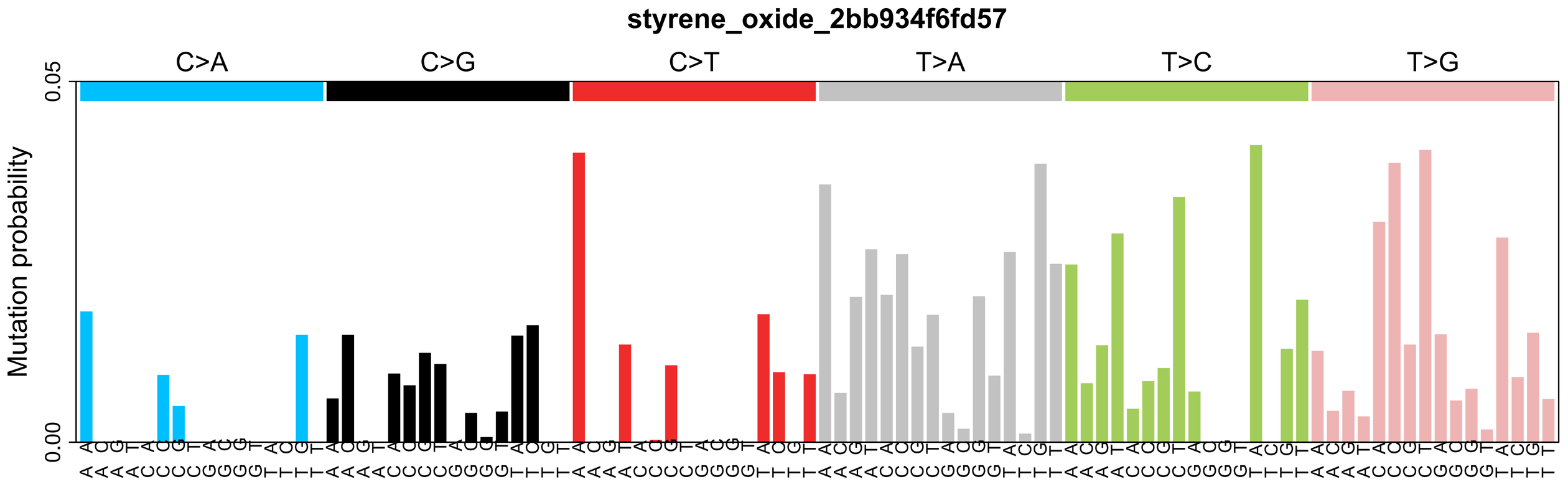
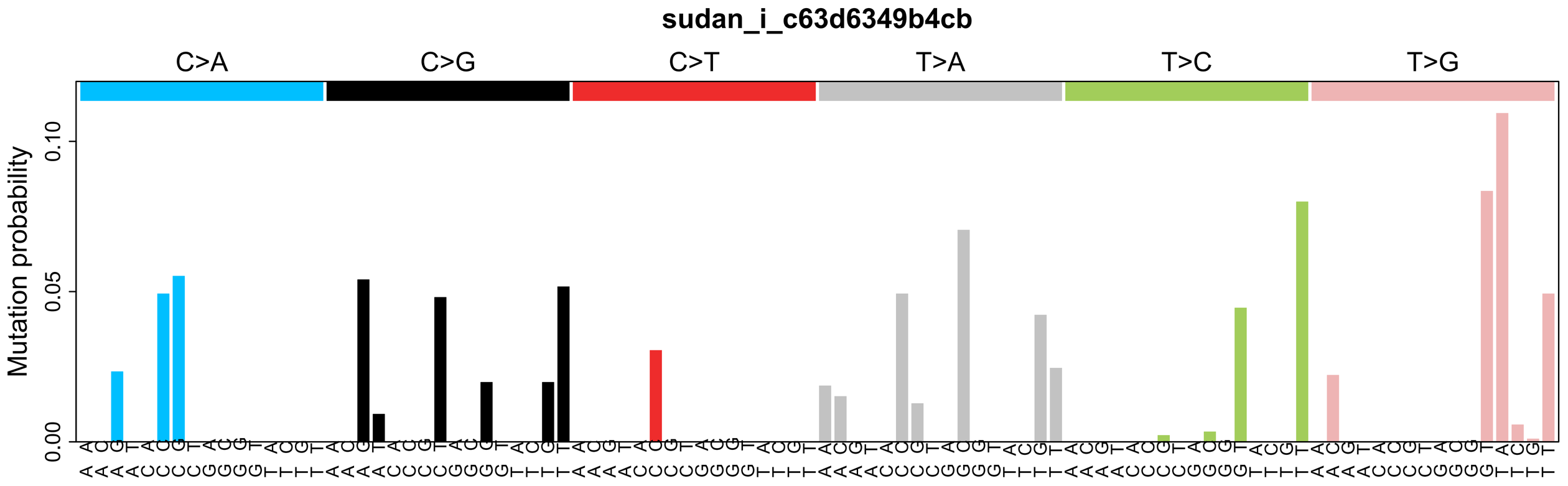
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15

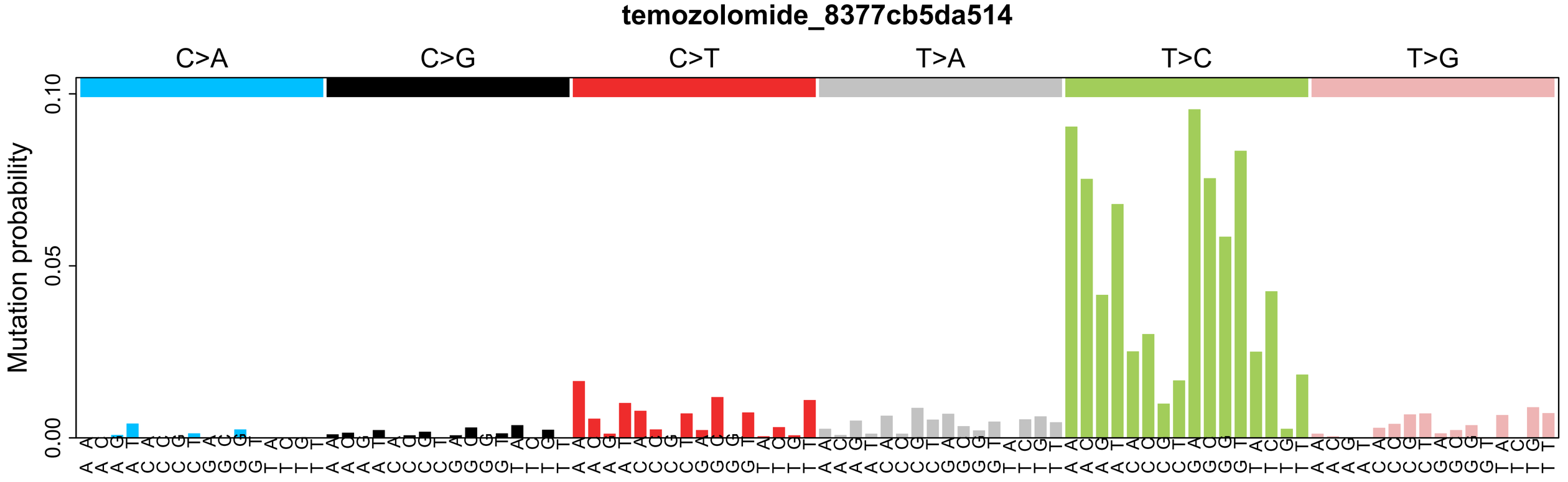

simulated_solar_radiation_04d97e27a495
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.0
0.1
0.2

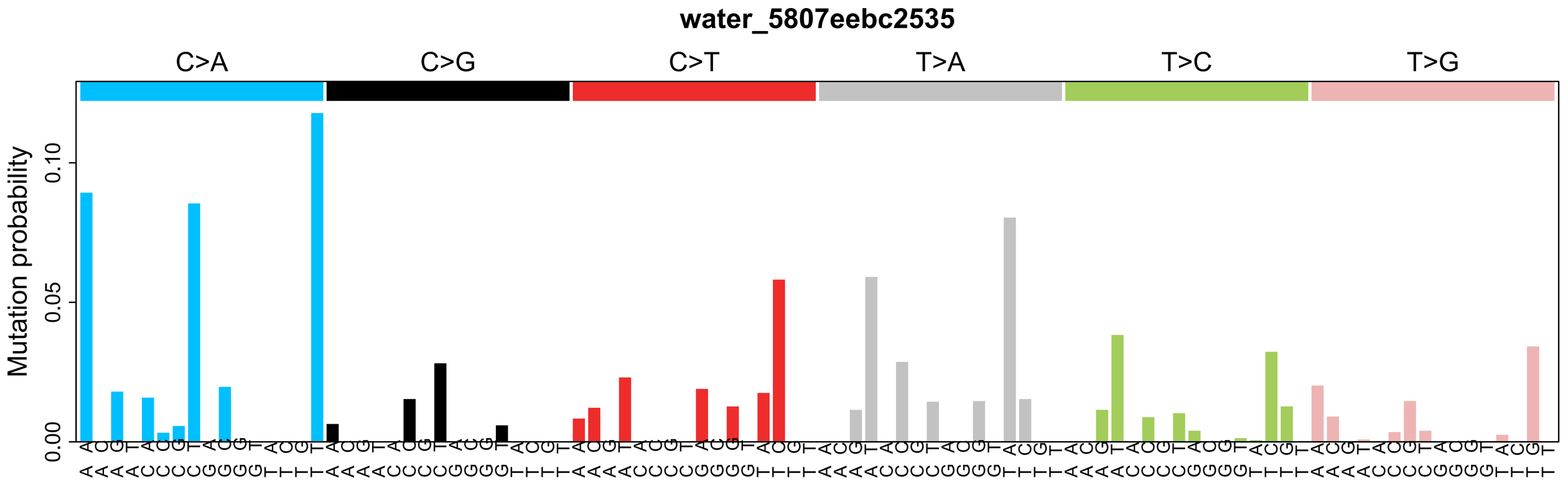

sodium_arsenite_90245b3e0140
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


sodium_chloride_02280bd72cf9
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10
0.15
0.20


styrene_oxide_2bb934f6fd57
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.05
0.00


sudan_i_c63d6349b4cb
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10


temozolomide_8377cb5da514
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.10
0.05
0.00


water_5807eebc2535
C>A
C>G
C>T
T>A
T>C
T>G
Mutation probability
0.00
0.05
0.10