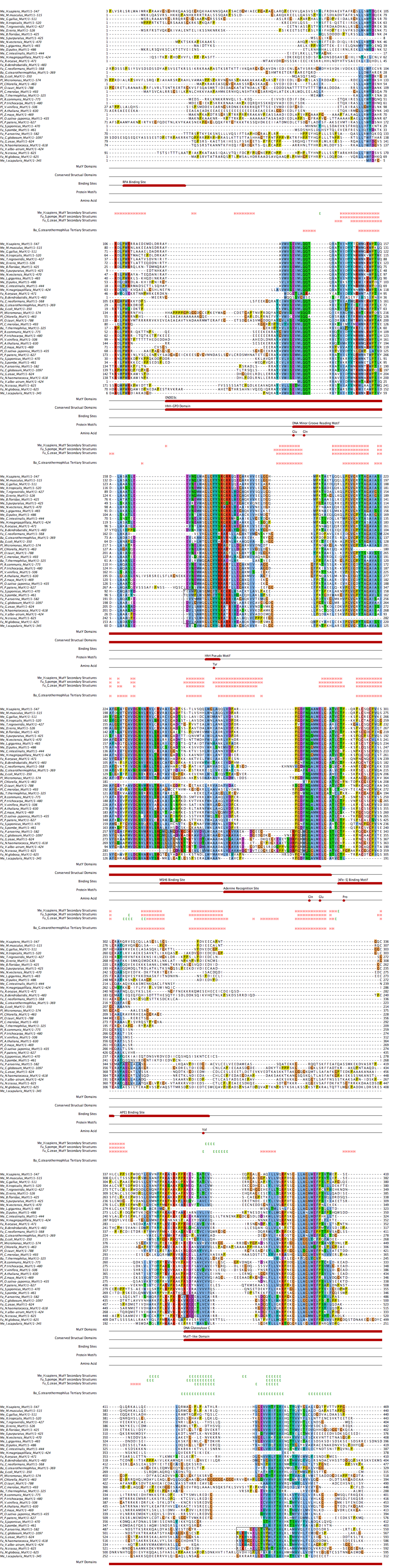
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Me_T.nigroviridis_MutY/1-427
Me_D.rerio_MutY/1-526
Me_B.floridae_MutY/1-425
Me_S.purpuratus_MutY/1-425
Me_N.vectensis_MutY/1-470
Me_L.gigantea_MutY/1-465
Me_D.pulex_MutY/1-486
Me_C.intestinalis_MutY/1-444
Me_H.magnipapilllata_MutY/1-424
Fu_R.oryzae_MutY/1-471
Fu_B.dendrobatidis_MutY/1-483
Fu_C.neoformans_MutY/1-568
Ba_G.stearothermophilus_MutY/1-369
Ba_E.coli_MutY/1-350
Pl_Micromonas_MutY/1-574
Pl_Chlorella_MutY/1-463
Pl_O.tauri_MutY/1-788
Pl_C.merolae_MutY/1-493
Ba_T.thermophilus_MutY/1-325
Pl_R.communis_MutY/1-775
Pl_P.trichocarpa_MutY/1-480
Pl_V.vinifera_MutY/1-506
Pl_A.thaliana_MutY/1-630
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Pl_O.sativa-japonica_MutY/1-435
Pl_P.patens_MutY/1-627
Fu_S.japonicus_MutY/1-470
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Fu_P.anserina_MutY/1-582
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Fu_N.haematococca_MutY/1-618
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Me_I.scapularis_MutY/1-345
MutY Domains
Conserved Structual Domains
Binding Sites
Protein Motifs
Amino Acid
Me_H.sapiens_MutY Secondary Structures
Fu_S.pompe_MutY secondary Structures
Fu_G.zeae_MutY Secondary Structures
Ba_G.stearothermophilus Tertiary Structures
RPA Binding Site
ENDO3c
HhH-GPD Domain
DNA Minor Groove Reading Motif
Glu
Gln
HhH Pseudo Motif
Tyr
MSH6 Binding Site
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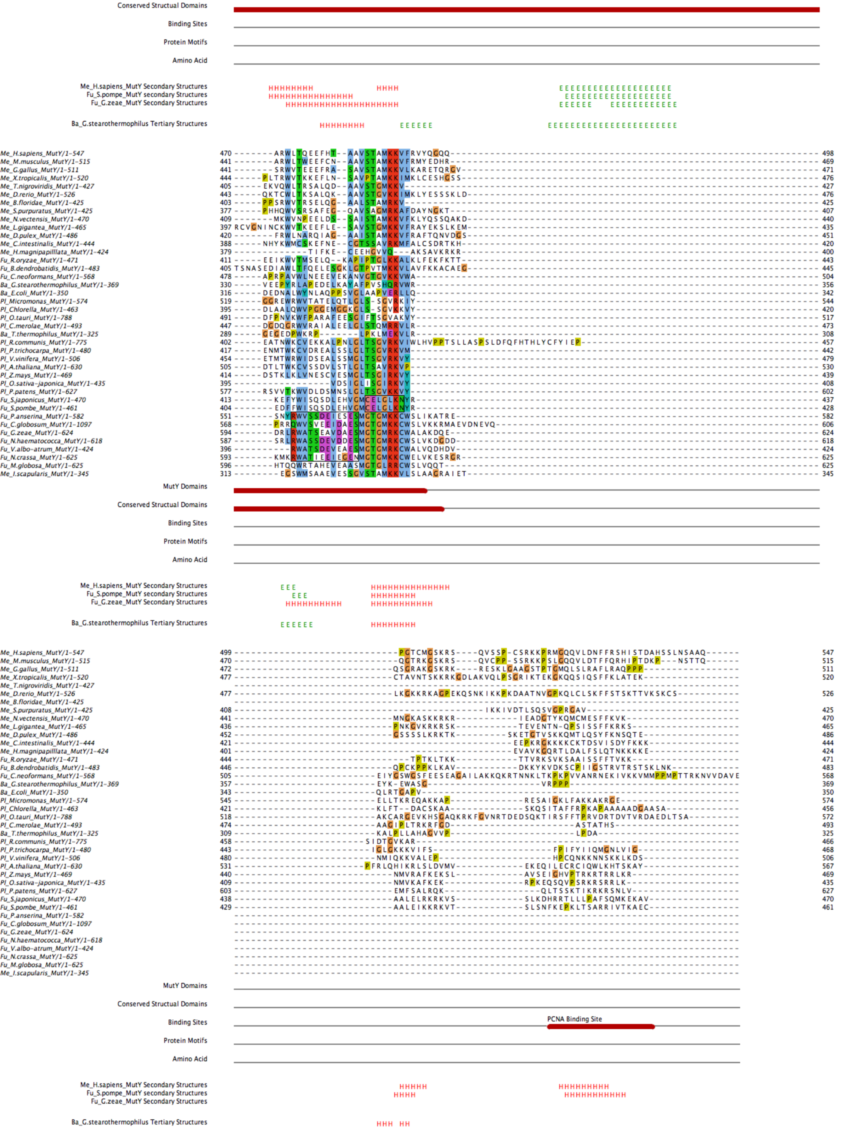
Adenine Recognition Site
Gln
Glu
Pro
APE1 Binding Site
Val
DNA Glycosylase C
MutT-like Domain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