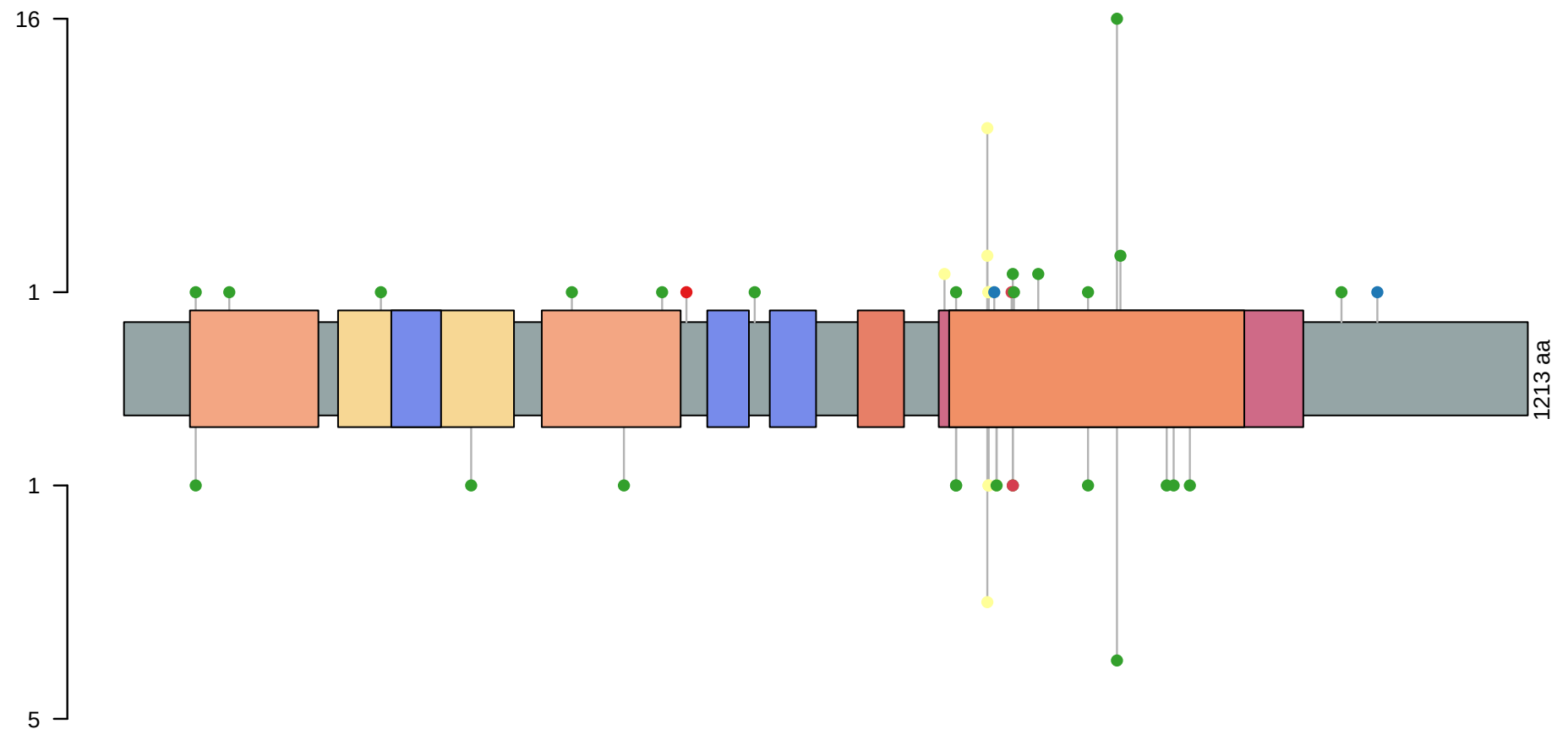


High [17.31%; N = 260]

EGFR: NM_005228



Low [8.21%; N = 207]

- Missense_Mutation
- Frame_Shift_Del
- Nonsense_Mutation
- In_Frame_Ins
- In_Frame_Del

Domains

- Recept_L_domain
- Furin-like
- FU
- TM_ErbB1
- PTKc_EGFR
- TyrKc