Reddy analysis [4.93%; N = 973]
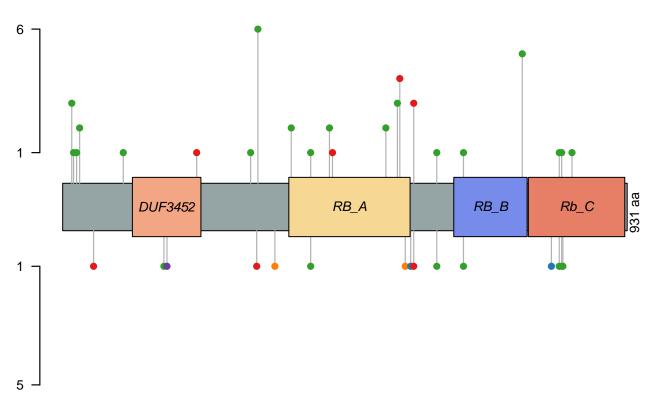
RB1: NM_000321

931 aa

DUF3452 • RB_A • RB_B • Rb_C

Reanalysis [1.57%; N = 958]

- Missense_Mutation
- Splice_Site
- Nonsense_Mutation
- Frame_Shift_Del
- Frame_Shift_Ins