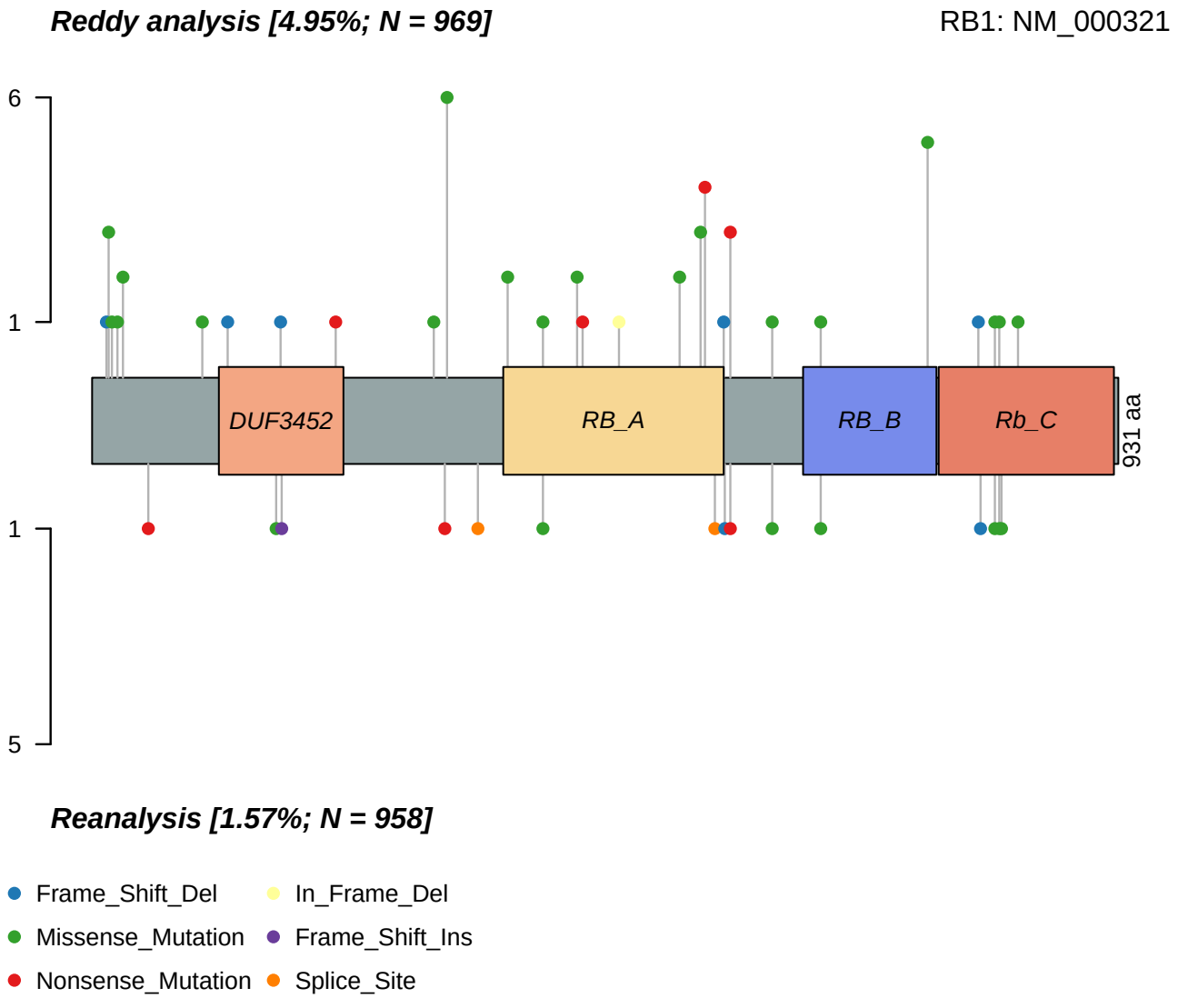
Reddy analysis [4.95%; N = 969]
RB1: NM_000321
6
1
1
5
DUF3452
RB_A
RB_B
Rb_C
931 aa
Reanalysis [1.57%; N = 958]
Frame_Shift_Del
In_Frame_Del
Missense_Mutation
Frame_Shift_Ins
Nonsense_Mutation
Splice_Site