Reddy analysis [1.96%; N = 969]

BRAF: NM_004333

R389H

R360Q

L597R

R671Q

Raf_RBD C1 PKinase Kinase 769 aa

G469A

D594... ...671N
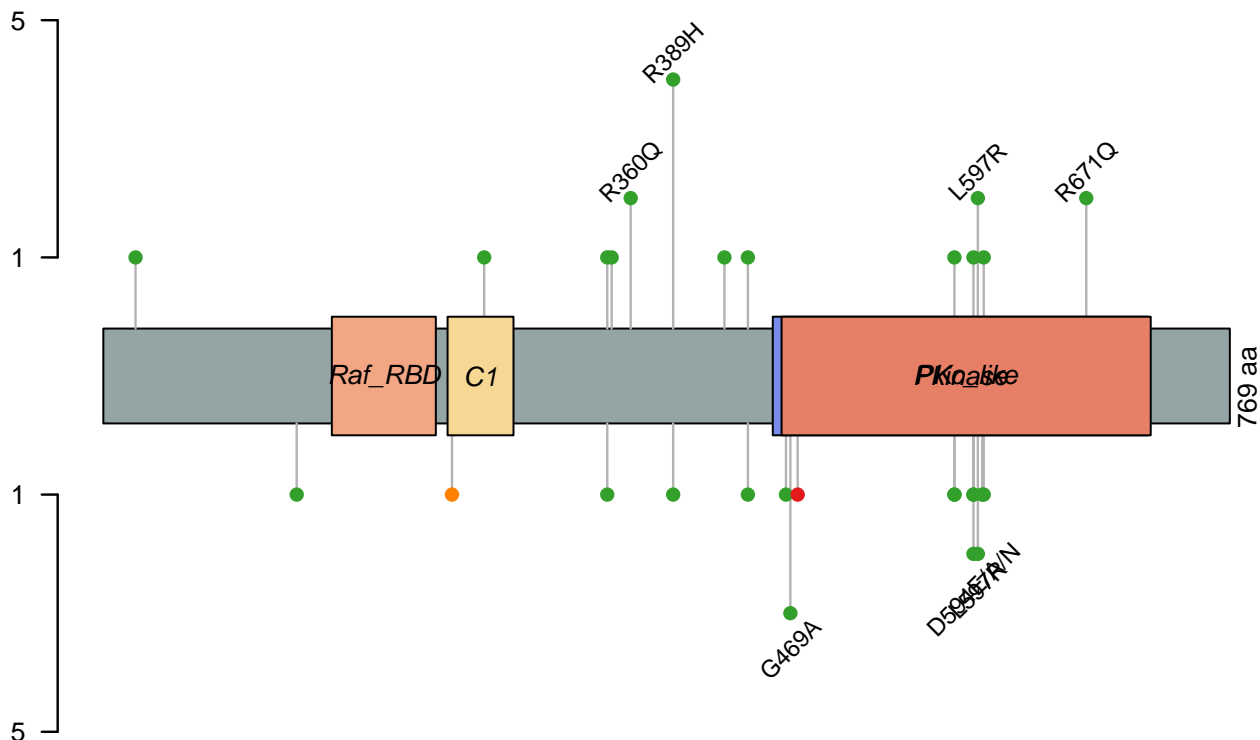
Reanalysis [2.09%; N = 958]

- ● Missense_Mutation
- ● Splice_Site
- ● Nonsense_Mutation