Reddy analysis [1.65%; N = 969]

TCL1A: NM_001098725

5

R52H

R83Q

Y26F

1

1

TCL1_MTCP1

117 aa

1

Y26F/C/S

12

_40_splice

Reanalysis [2.51%; N = 958]

- Missense_Mutation
- Nonsense_Mutation
- Splice_Site