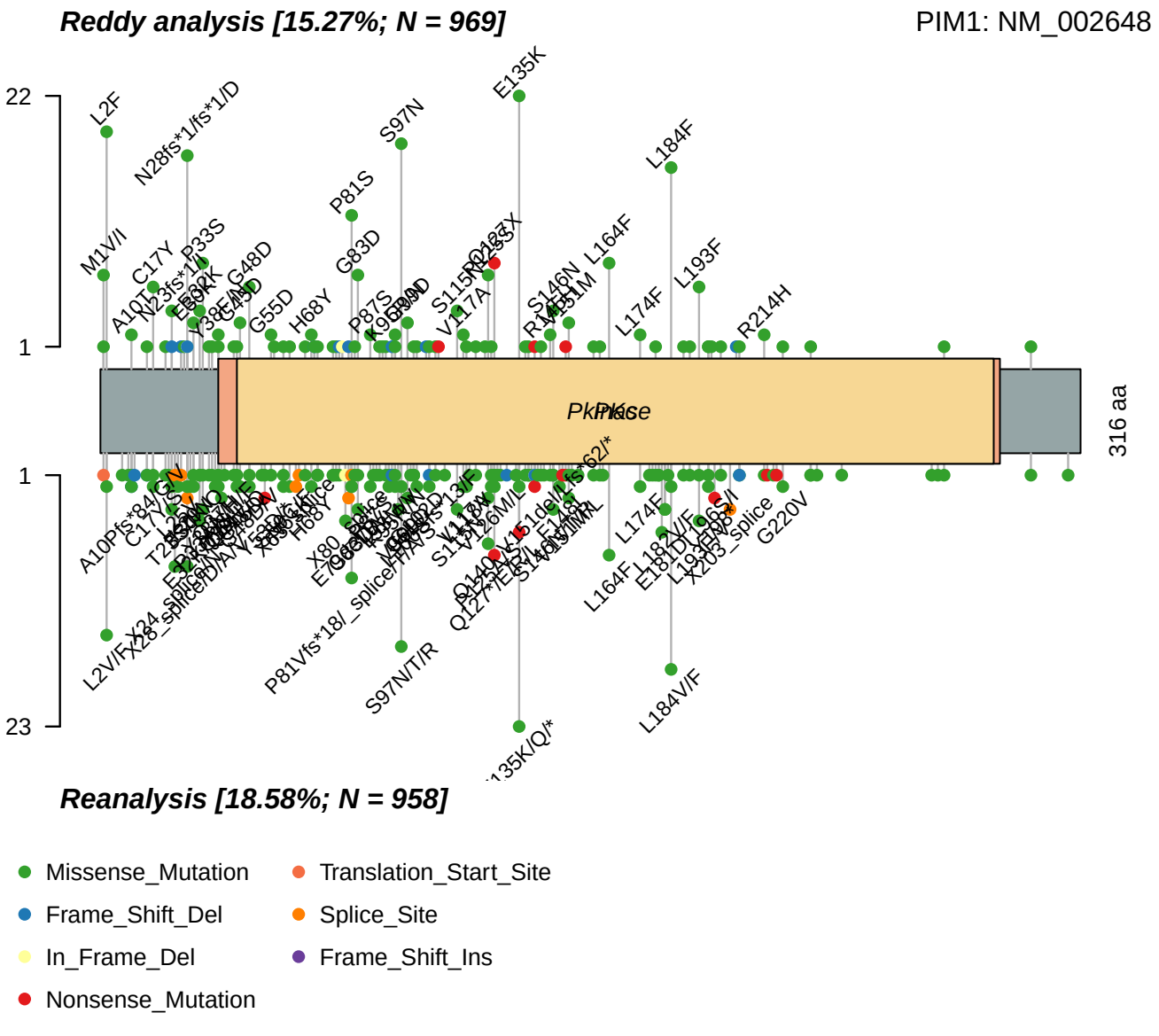
Reddy analysis [15.27%; N = 969]
PIM1: NM_002648
22
1
1
23
L2F
N28fs*1/fs*1/D
S97N
E135K
L184F
P81S
M1V/I
C17Y
P33S
G83D
G48D
G55D
H68Y
P87S
V117A
L164F
L193F
L174F
R214H
PKinase
316 aa
A10Pfs*84/G
L2V/F
P81Vfs*18/ _splice
S97N/T/R
L184V/F
G220V
X203_splice
Reanalysis [18.58%; N = 958]
Missense_Mutation
Frame_Shift_Del
In_Frame_Del
Nonsense_Mutation
Translation_Start_Site
Splice_Site
Frame_Shift_Ins