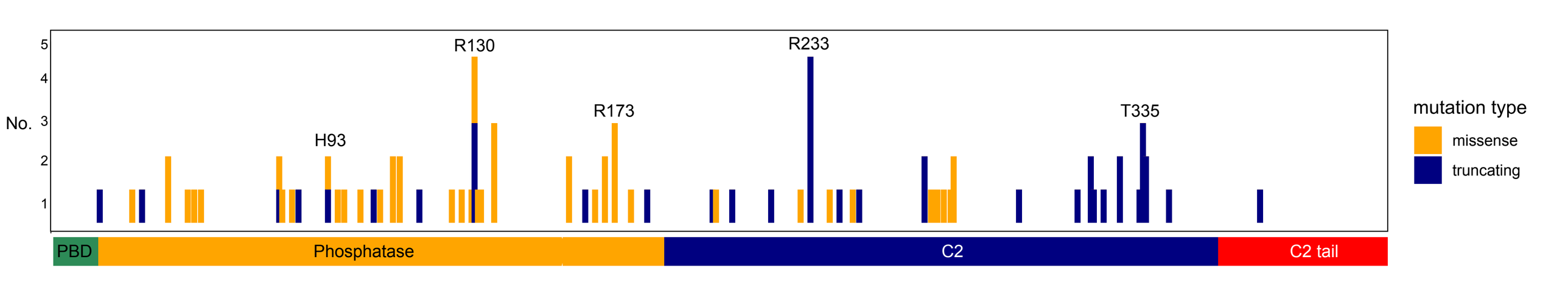
No.
5
4
3
2
1
R130
R233
R173
T335
H93
mutation type
missense
truncating
PBD
Phosphatase
C2
C2 tail