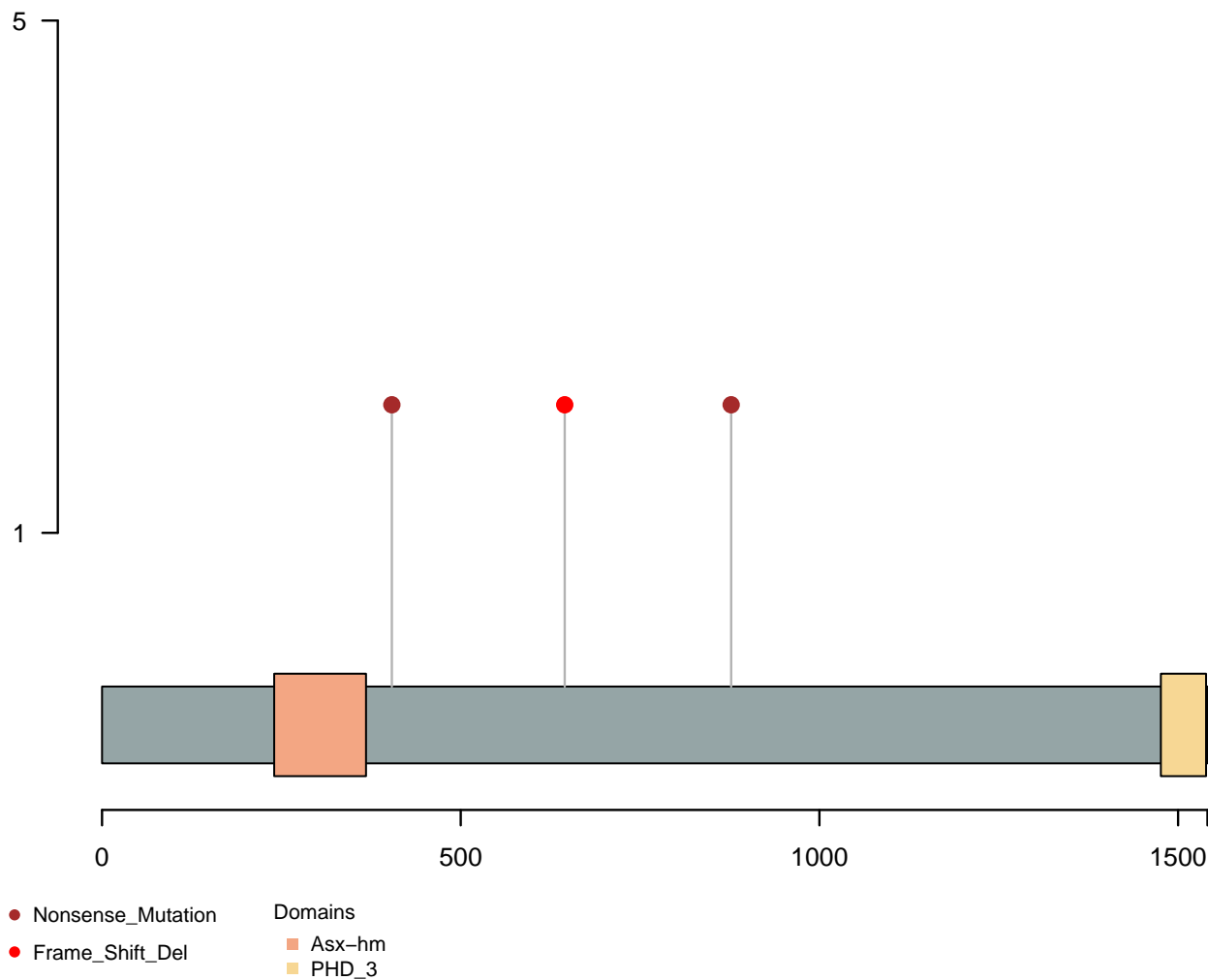


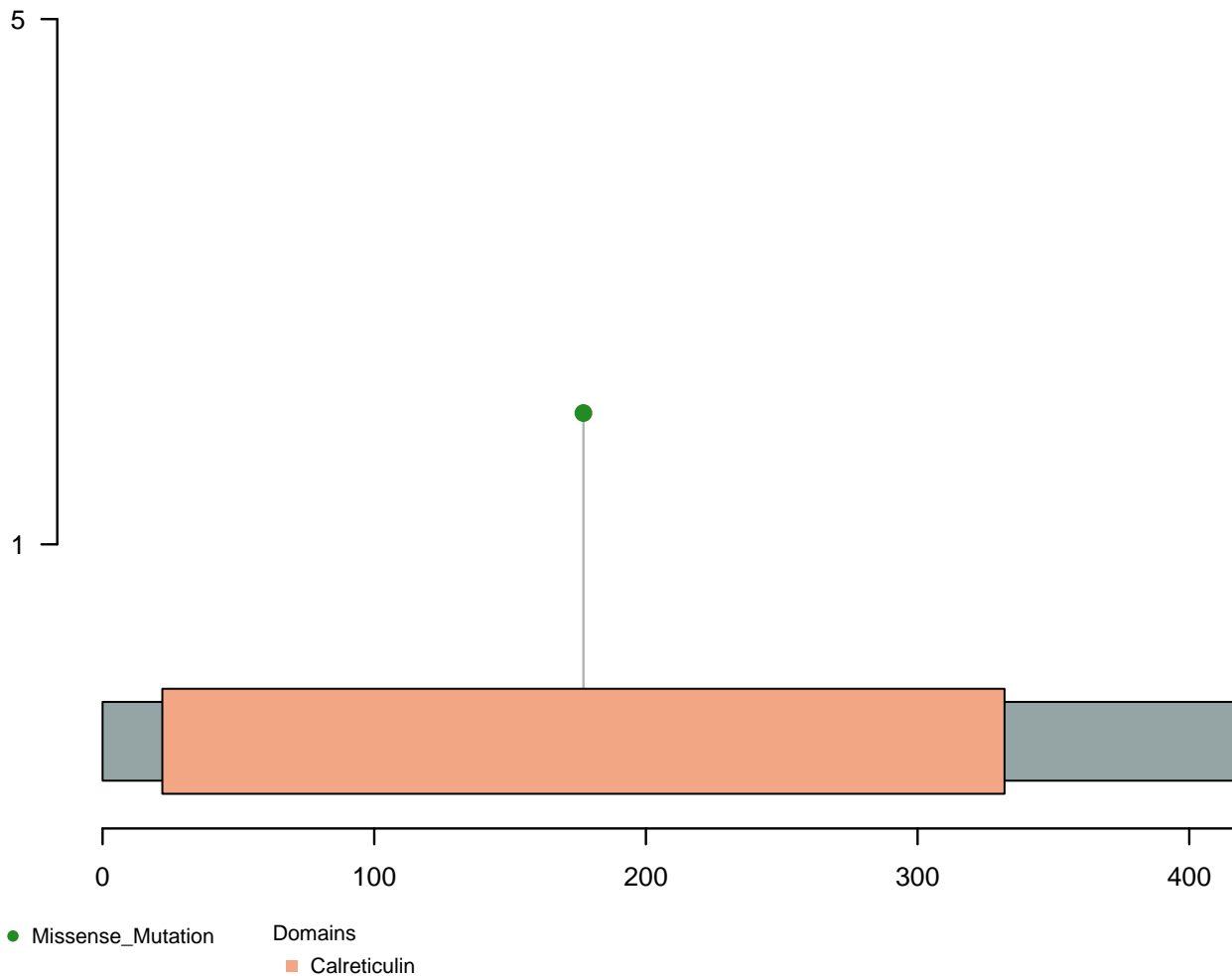
# ASXL1 : [Somatic Mutation Rate: 42.86%]

NM\_015338

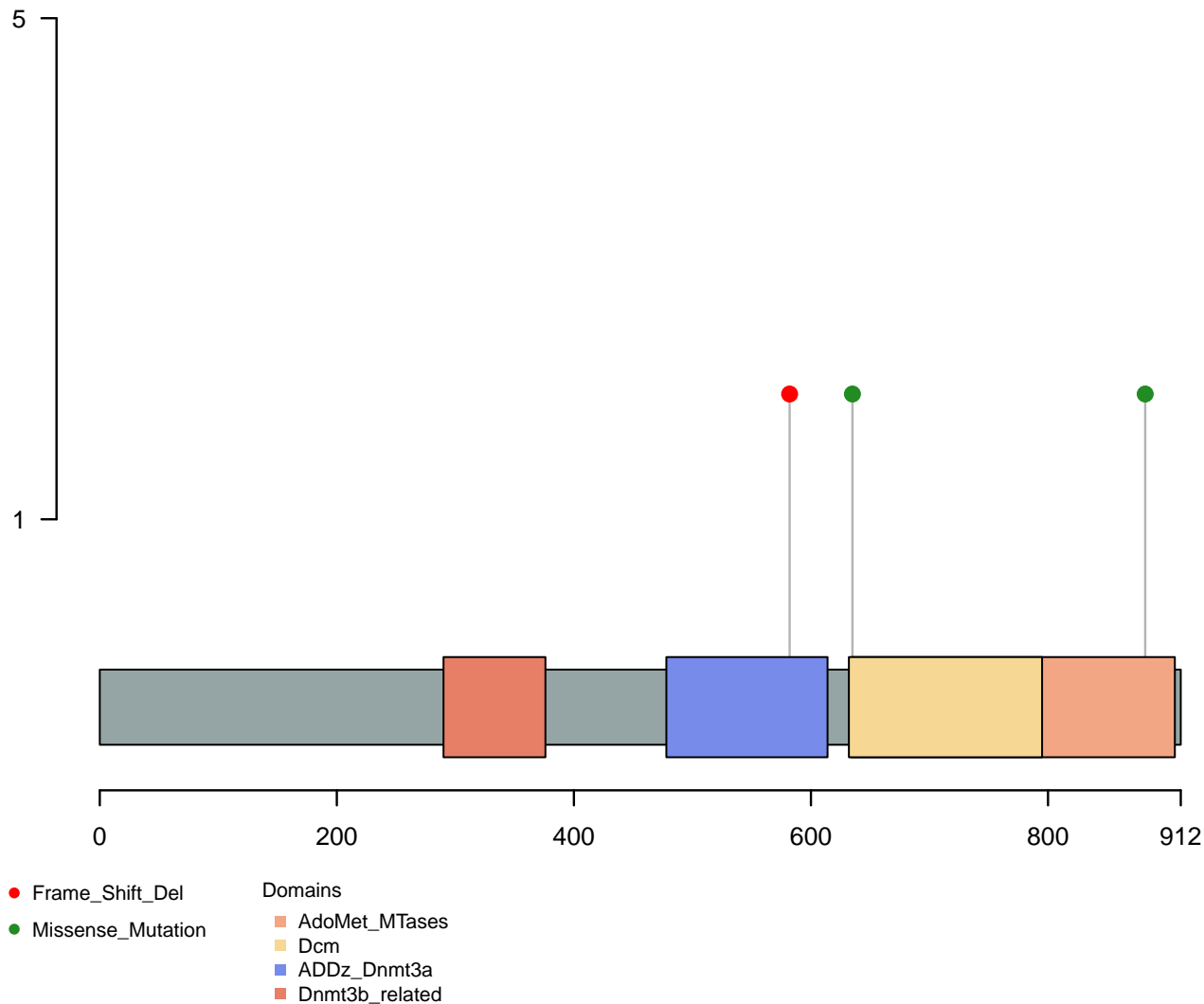


*CALR* : [Somatic Mutation Rate: 14.29%]

NM\_004343

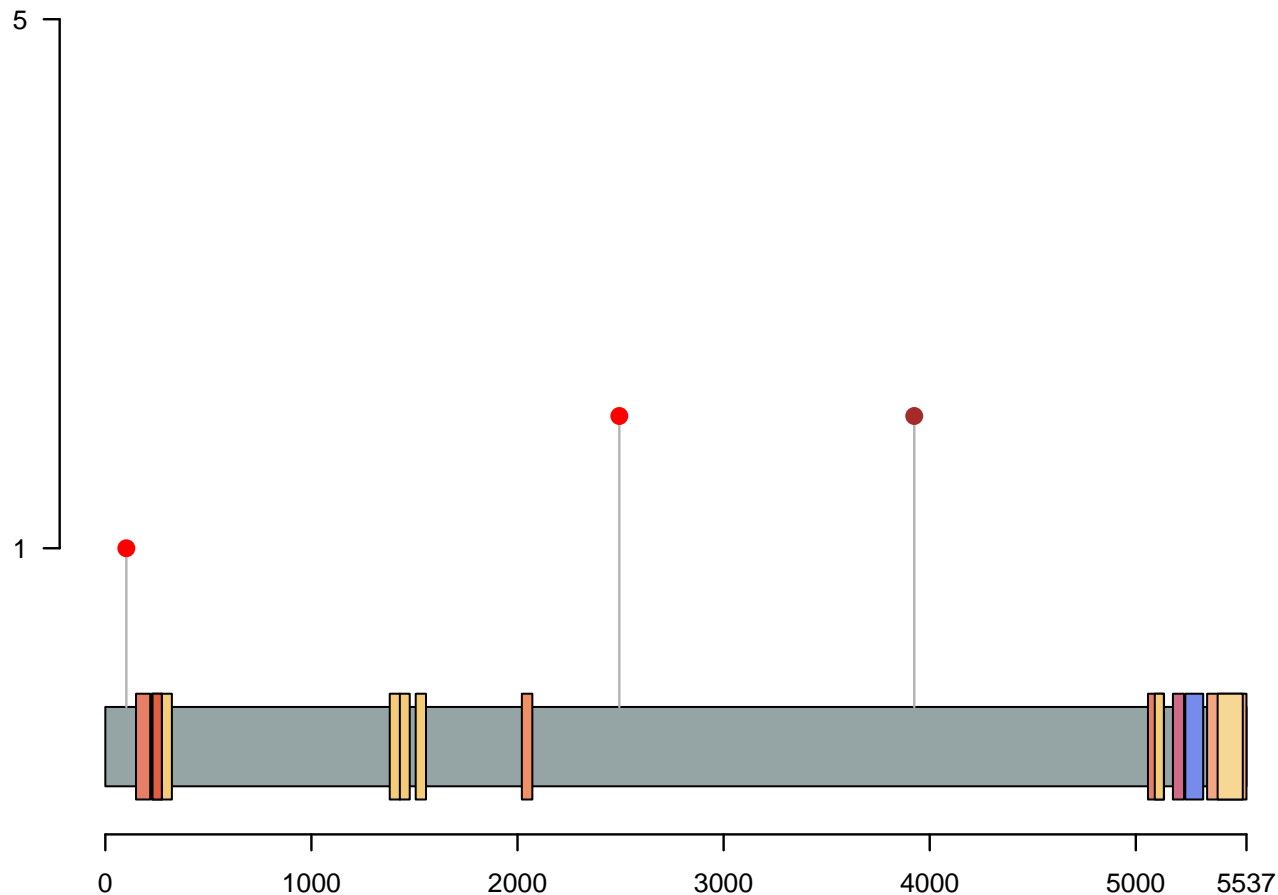


*DNMT3A* : [Somatic Mutation Rate: 42.86%]  
NM\_022552



*KMT2D* : [Somatic Mutation Rate: 35.71%]

NM\_003482



• Frame\_Shift\_Del

• Nonsense\_Mutation

Domains

COG2940

SET

FYRC

zf-HC5HC2H

FYRN

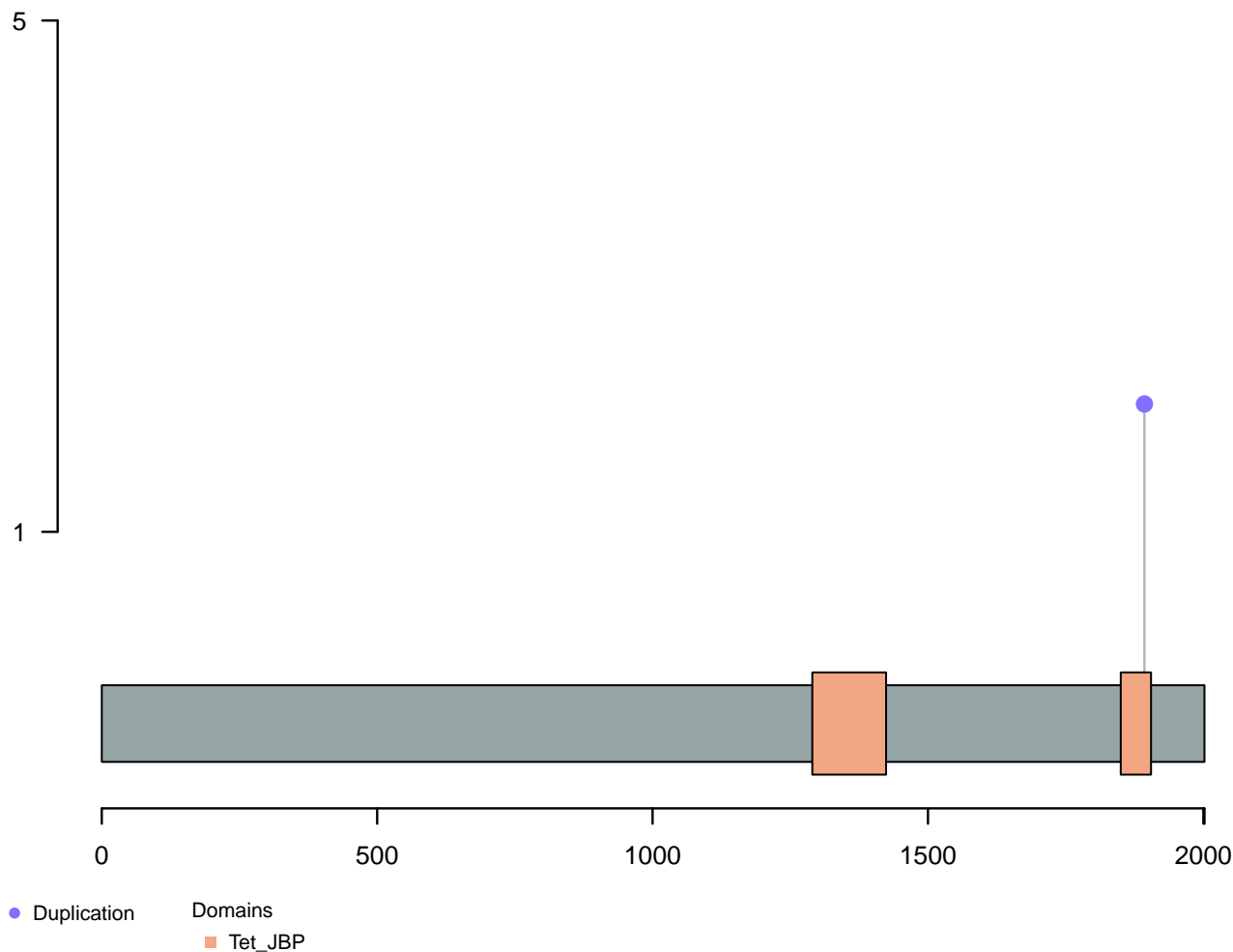
HMG-box

PHD

zf-C3HC4\_2

RING

*TET2* : [Somatic Mutation Rate: 14.29%]  
NM\_001127208



TP53 : [Somatic Mutation Rate: 14.29%]

NM\_000546

