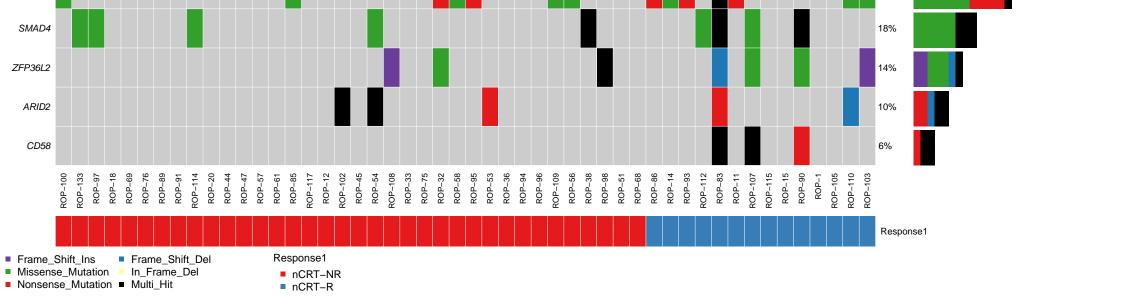
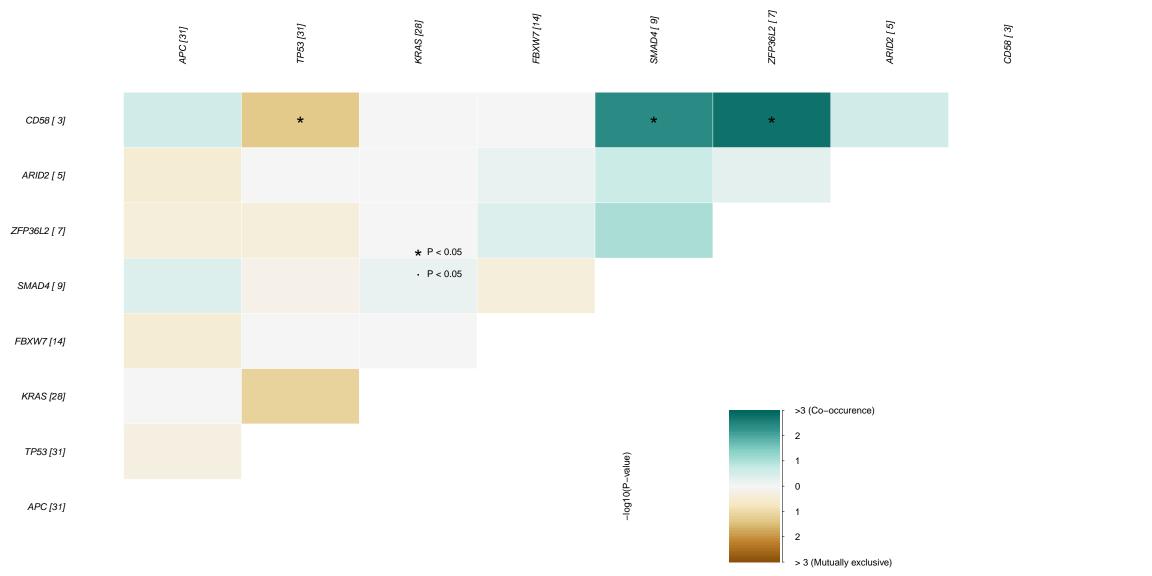
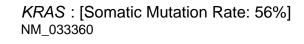
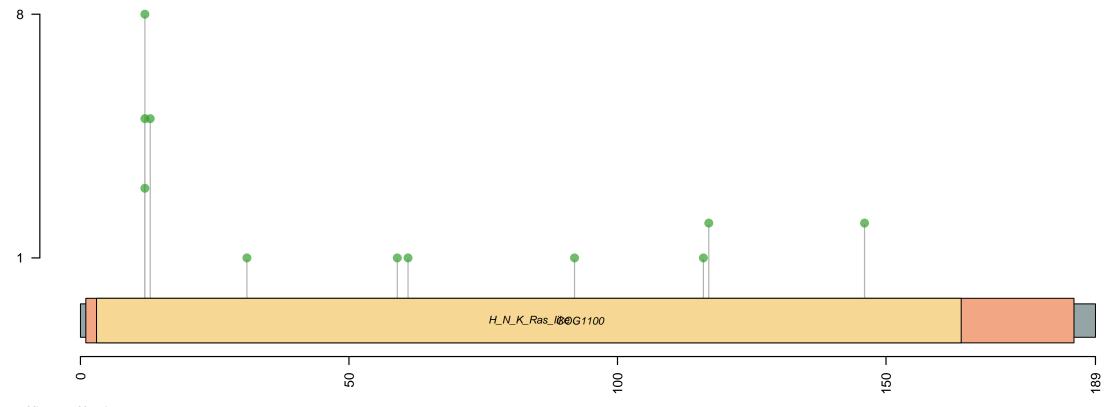
Altered in 50 (100%) of 50 samples. 16 TMB 31 No. of samples APC 62% TP53 62% KRAS 56% FBXW7 28% SMAD4 18% ZFP36L2 14% ARID2 CD58 6% ROP-97 ROP-18 ROP-69 ROP-76 ROP-12 ROP-45 ROP-54 ROP-108 ROP-33 ROP-95 ROP-53 ROP-94 ROP-96 ROP-109 ROP-98 ROP-51 ROP-68 ROP-93 ROP-112 ROP-11 ROP-107 ROP-90 ROP-105 ROP-110 ROP-103 ROP-20 ROP-47 ROP-57 ROP-61 ROP-85 ROP-117 ROP-75 ROP-32 ROP-58 ROP-56 ROP-38 ROP-86 ROP-14 ROP-15 ROP-44 ROP-1 Response1



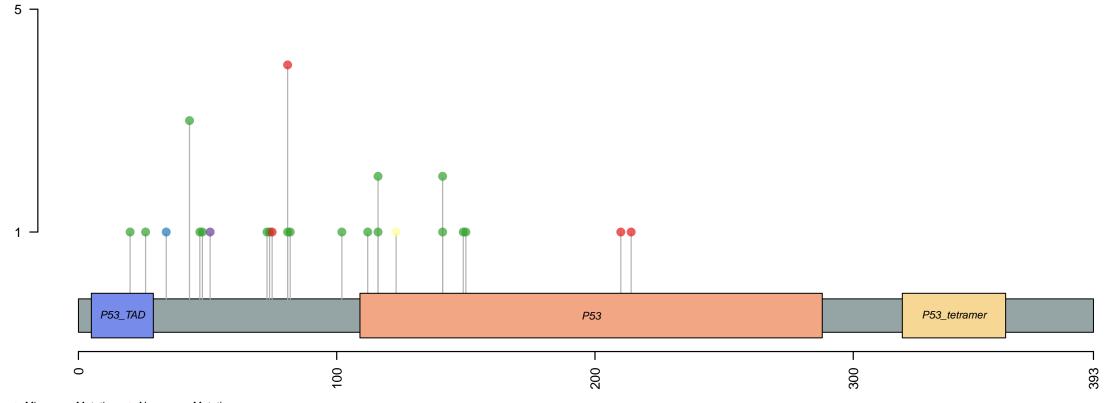






Missense_Mutation

TP53 : [Somatic Mutation Rate: 62%] NM_000546

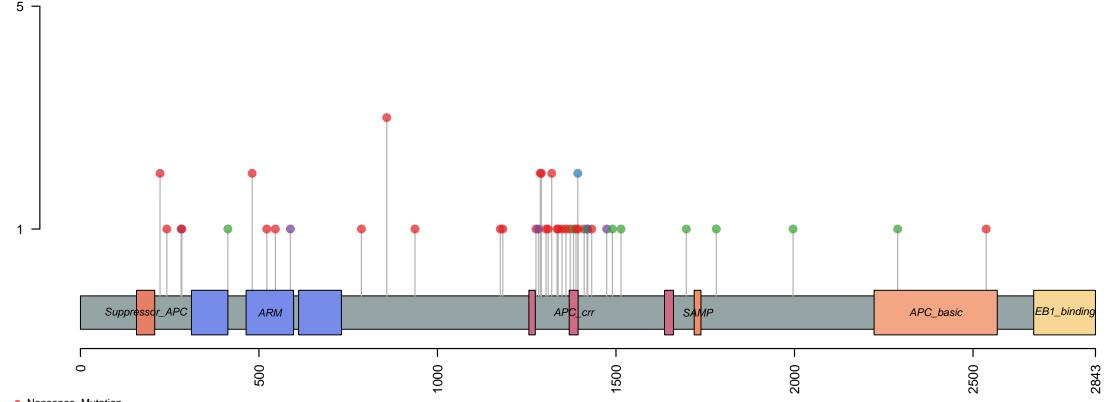


Missense_MutationNonsense_Mutation

Frame_Shift_DelIn_Frame_Del

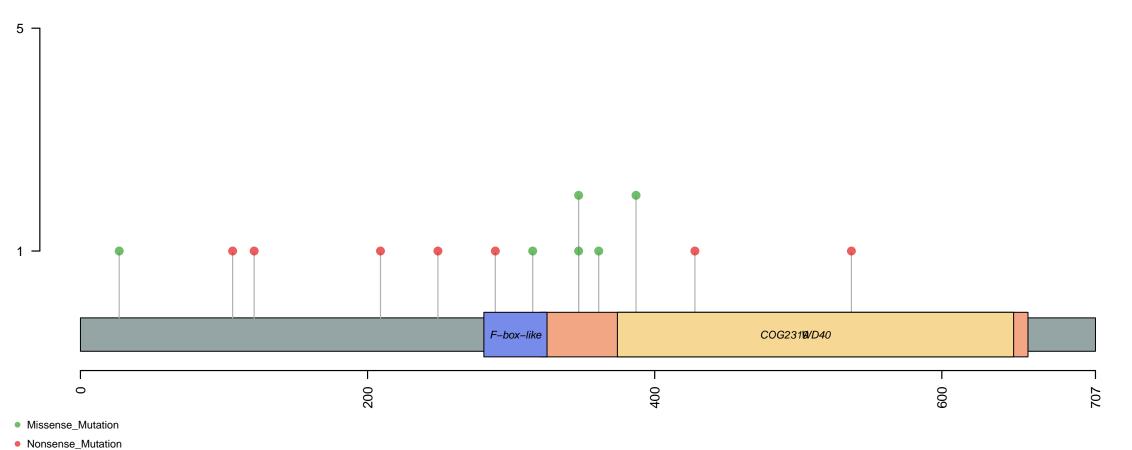
Frame_Shift_Ins

APC: [Somatic Mutation Rate: 62%] NM_000038

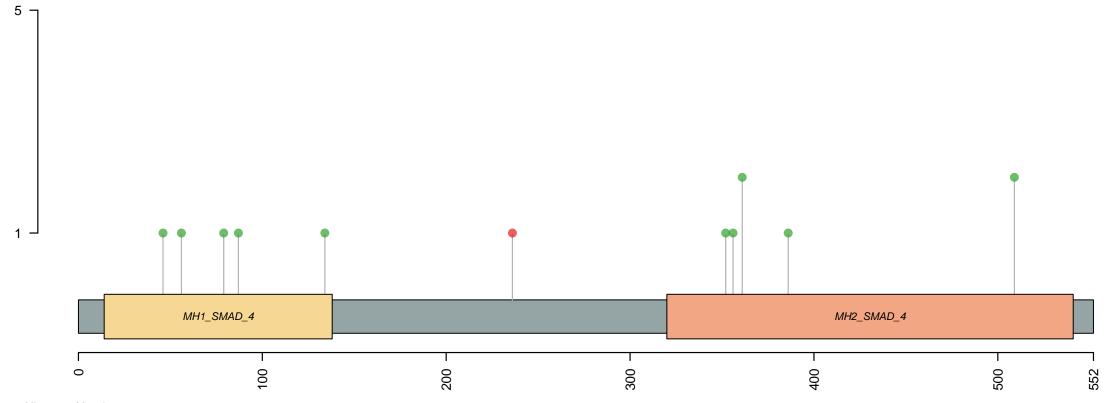


- Nonsense_Mutation
- Frame_Shift_Ins
- Missense_Mutation
- Frame_Shift_Del

FBXW7 : [Somatic Mutation Rate: 28%] NM_033632



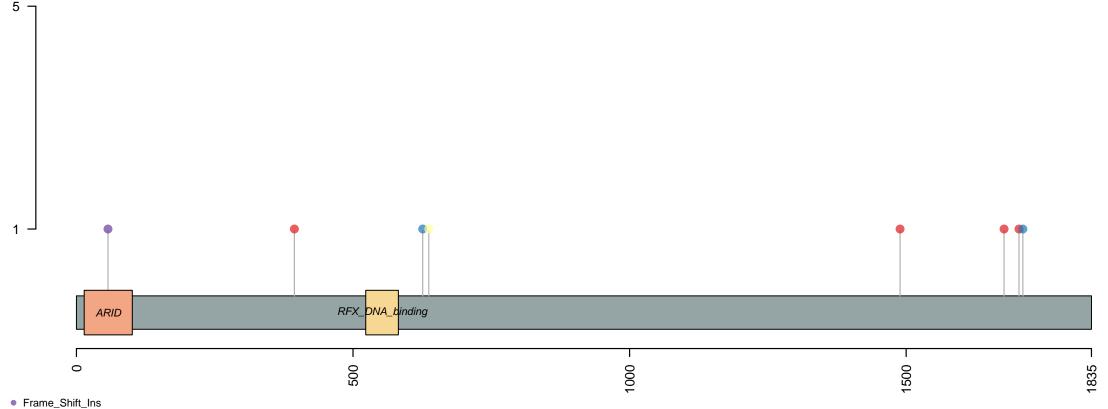
SMAD4 : [Somatic Mutation Rate: 18%] NM_005359



Missense_Mutation

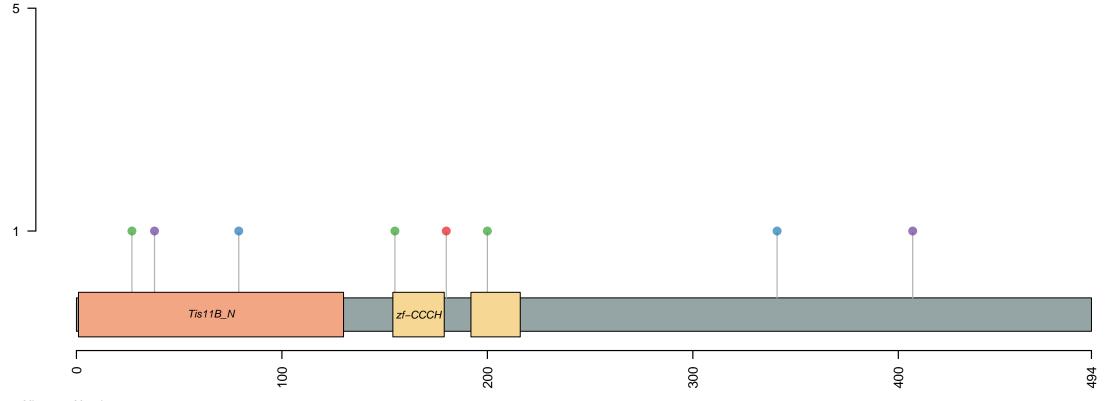
Nonsense_Mutation

ARID2: [Somatic Mutation Rate: 10%] NM_152641

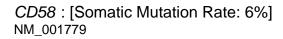


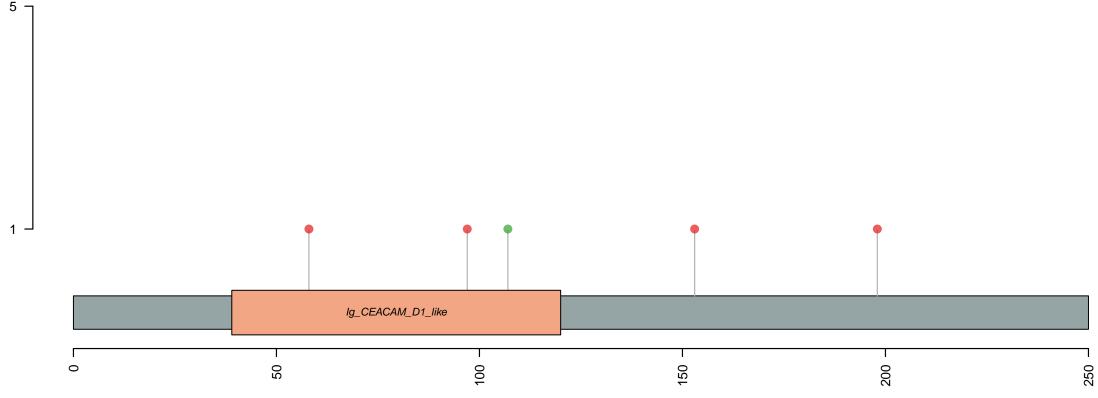
- Nonsense_Mutation
- Frame_Shift_Del
- In_Frame_Del

ZFP36L2 : [Somatic Mutation Rate: 14%] NM_006887



- Missense_Mutation
- Frame_Shift_Ins
- Frame_Shift_Del
- Nonsense_Mutation





Nonsense_Mutation

Missense_Mutation