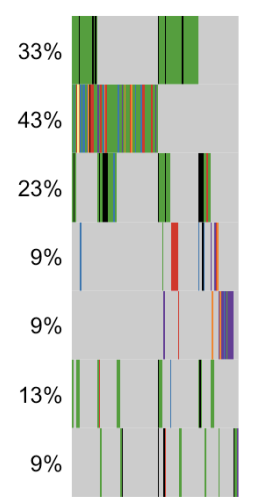
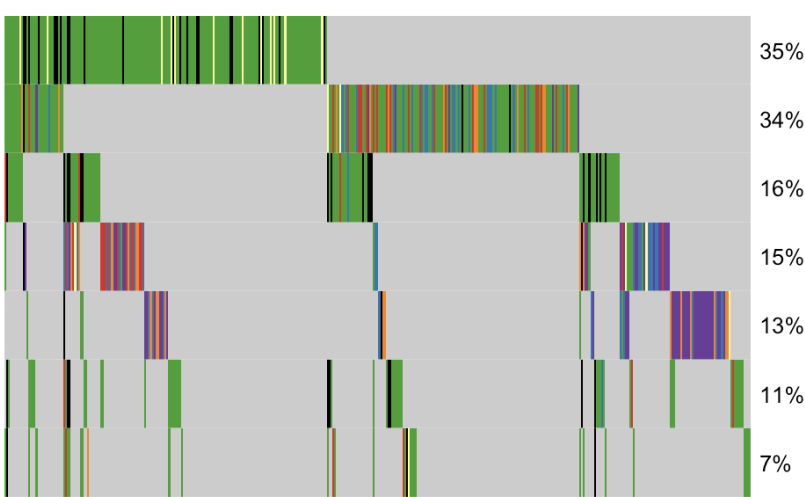


Positive Patients (N = 100)

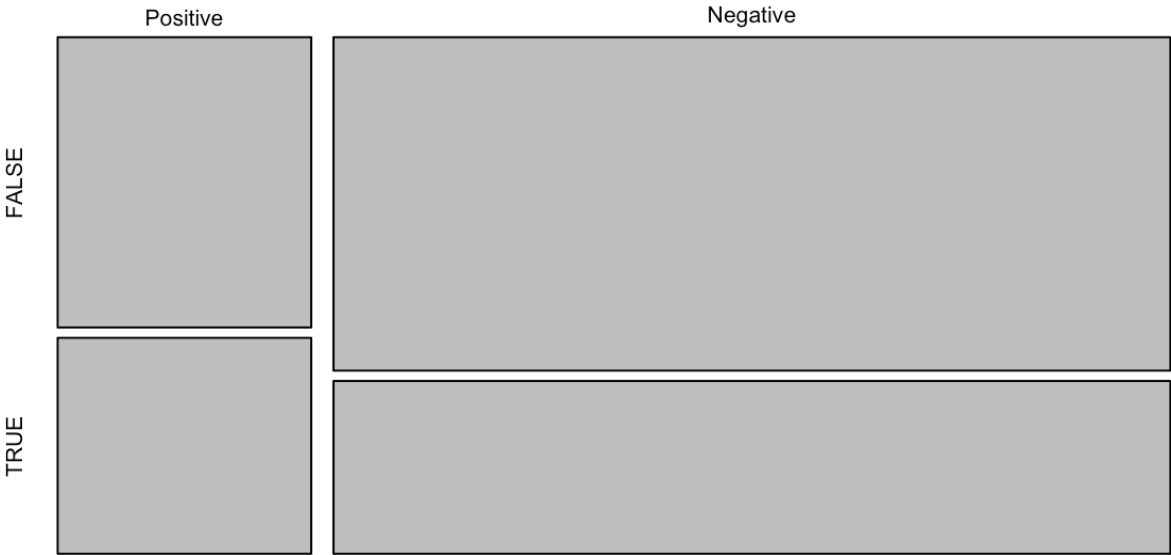


Negative Patients (N = 494)



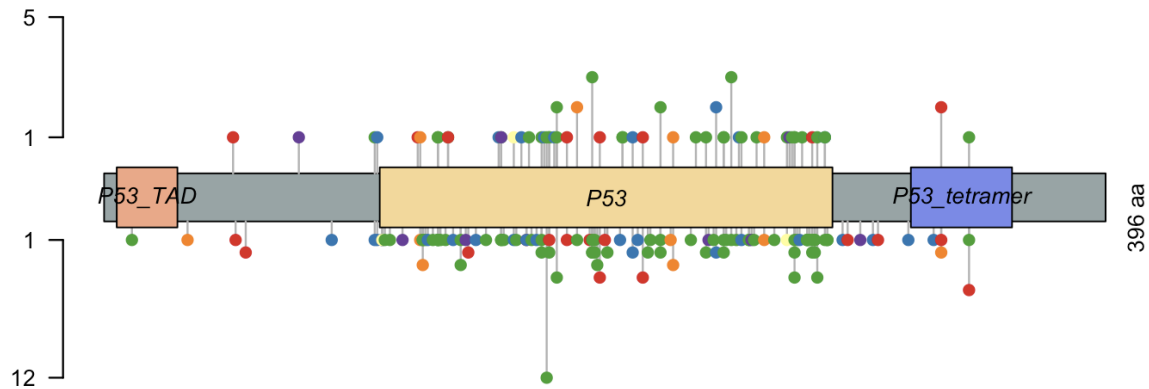
- Missense_Mutation
- Frame_Shift_Del
- Frame_Shift_Ins
- Splice_Site
- Nonsense_Mutation
- In_Frame_Del
- Multi_Hit
- In_Frame_Ins

contig



HER2 Positive [42.67%; N = 150]

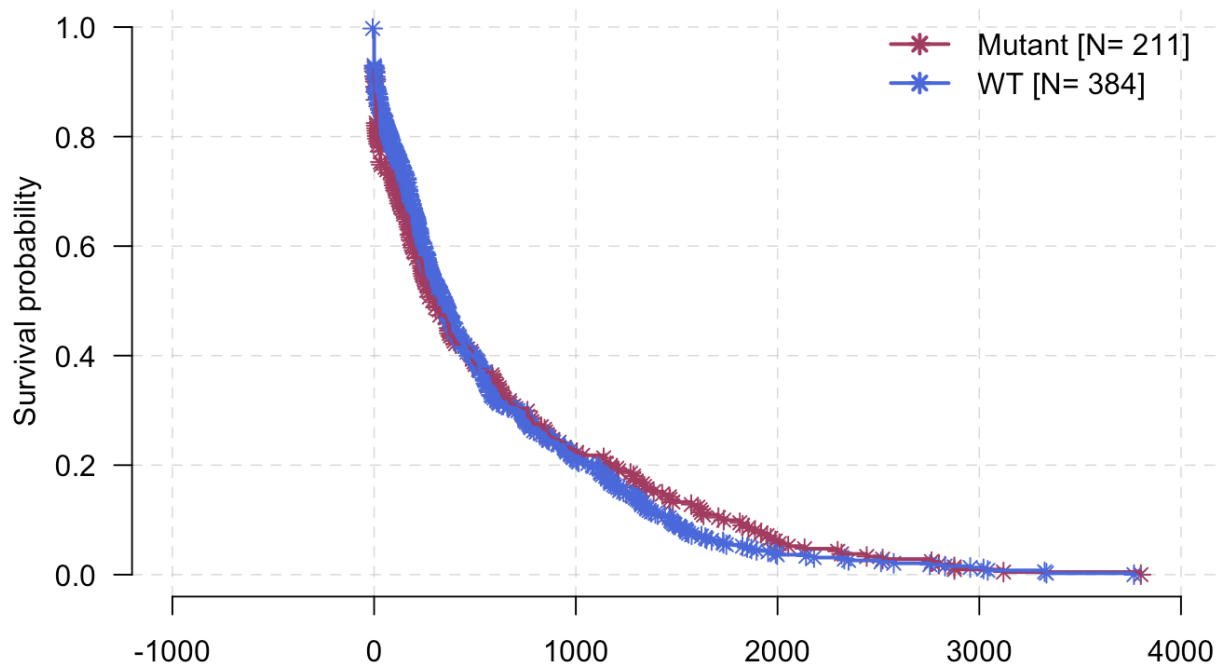
TP53: NM_000546



HER2 Negative [34.21%; N = 494]

- Nonsense_Mutation
- Frame_Shift_Del
- Frame_Shift_Ins
- Splice_Site
- Missense_Mutation
- In_Frame_Del

Mutant v/s WT



P-value: 0.632; HR: 0.96