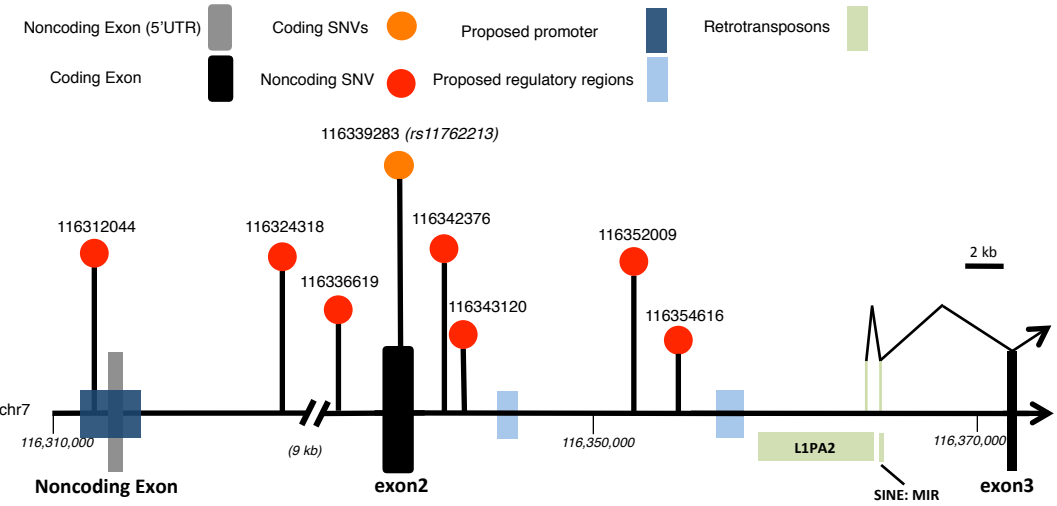
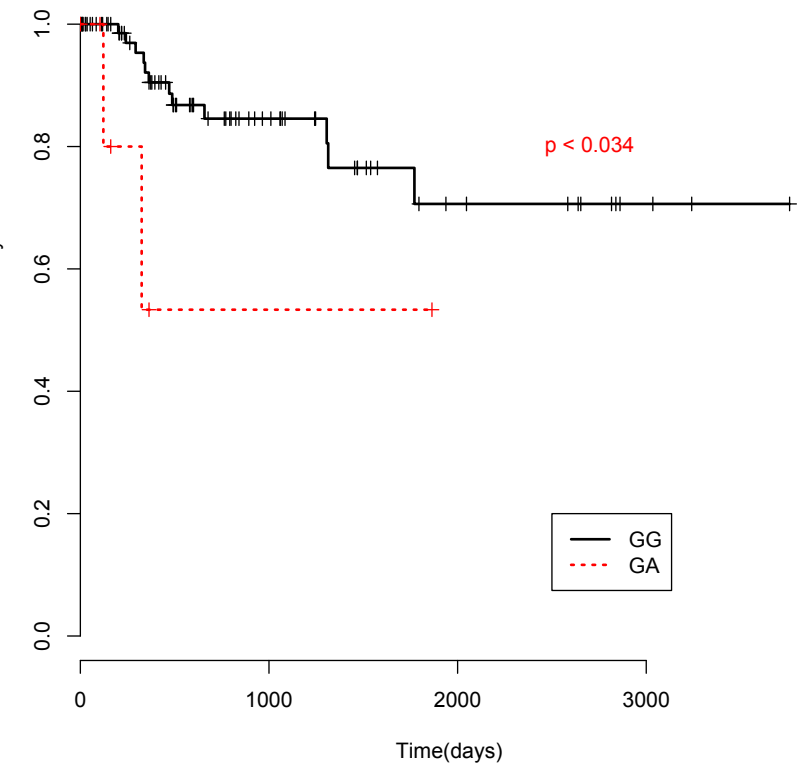


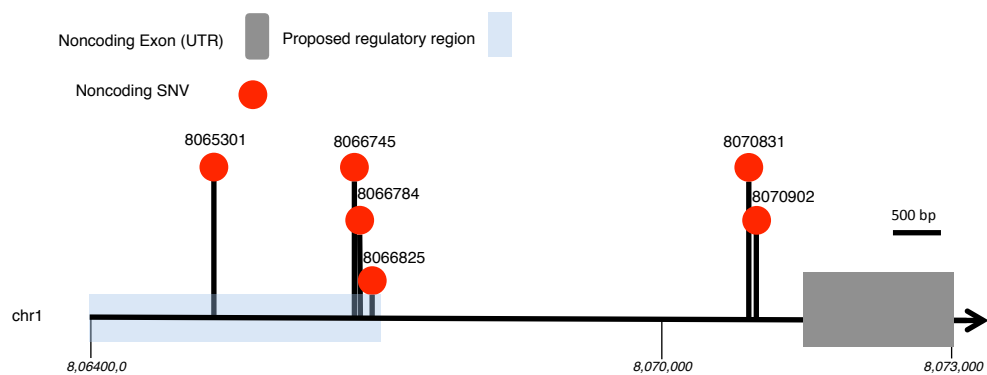
A MET



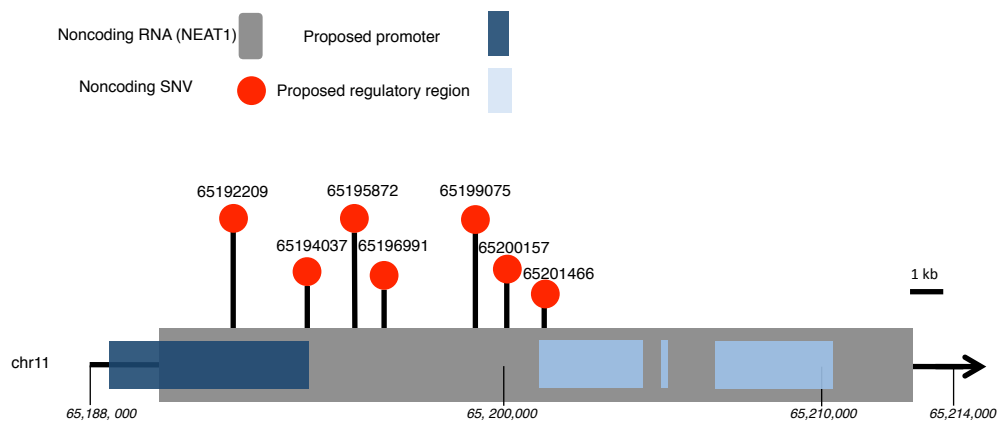
B



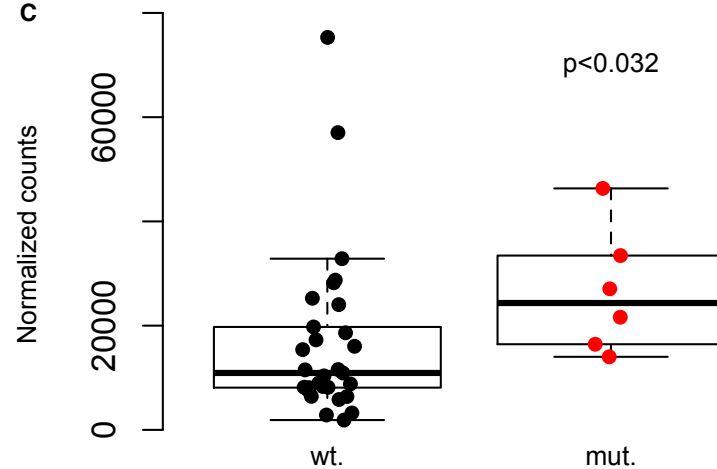
A ERRFI1



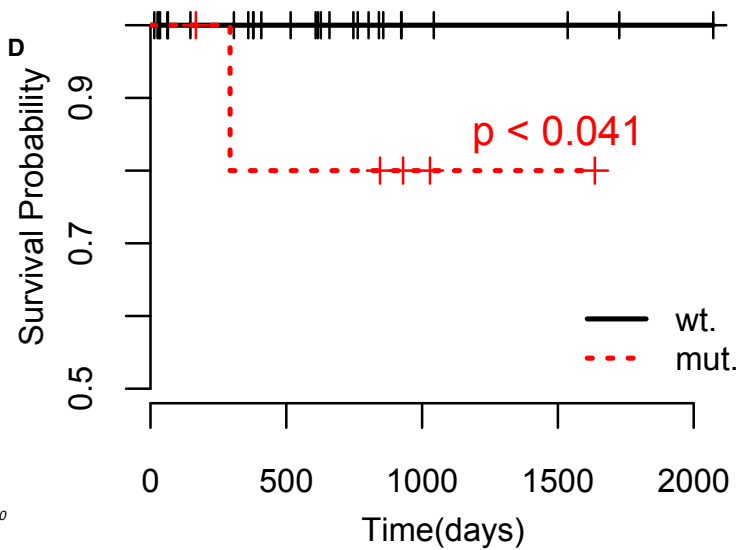
B NEAT1

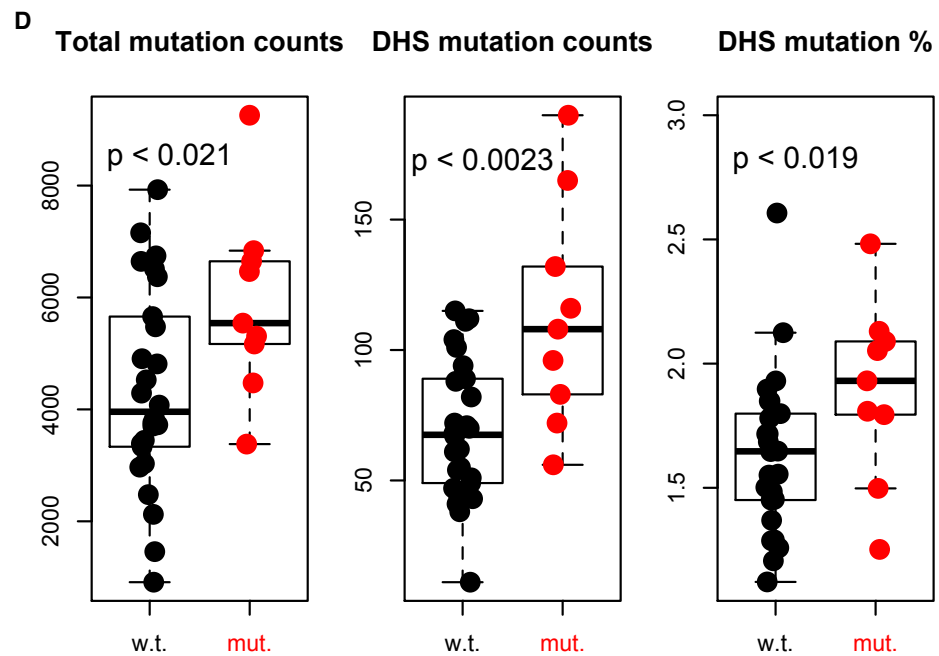
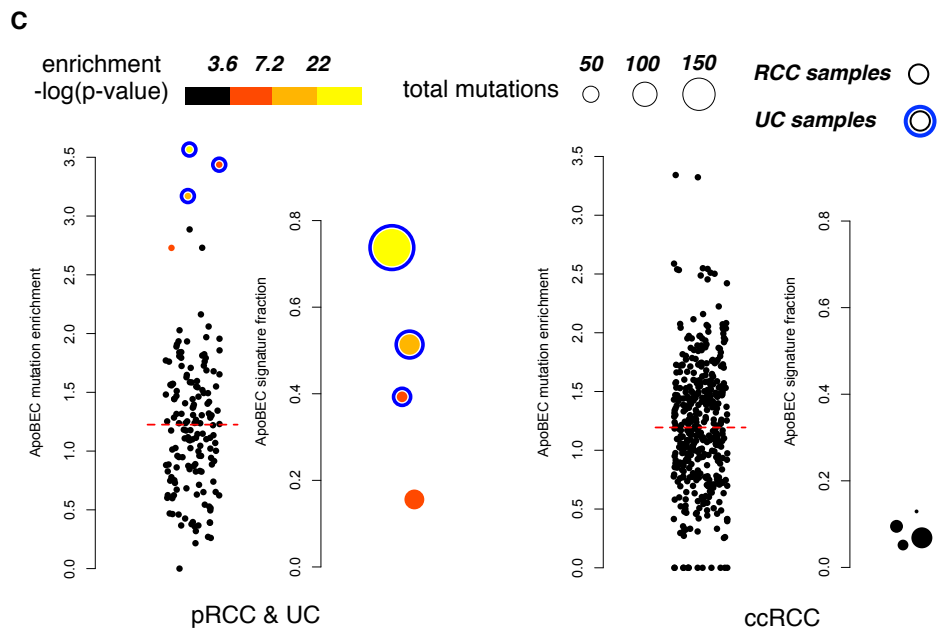
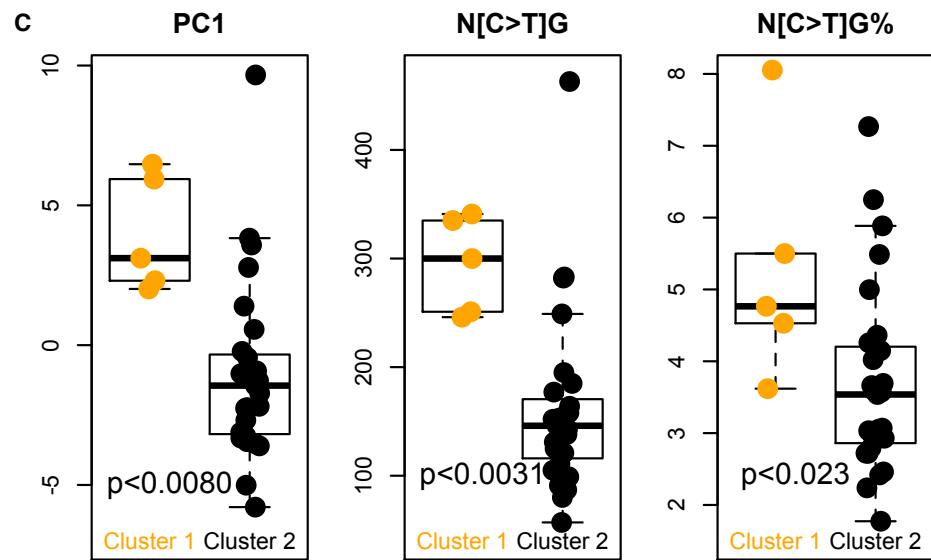
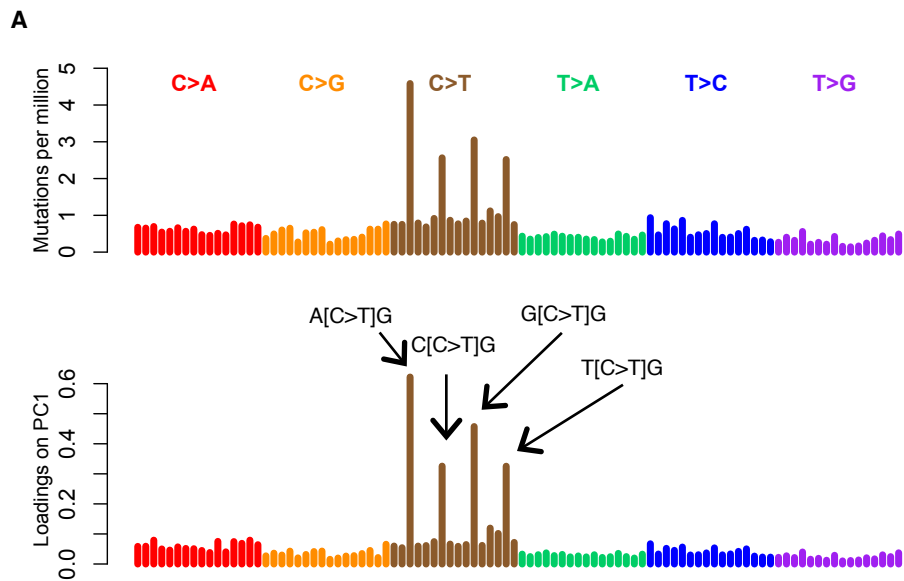


C

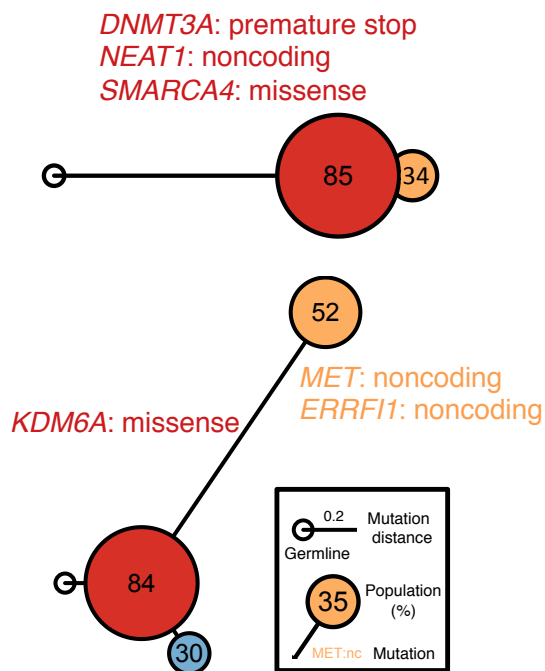


D

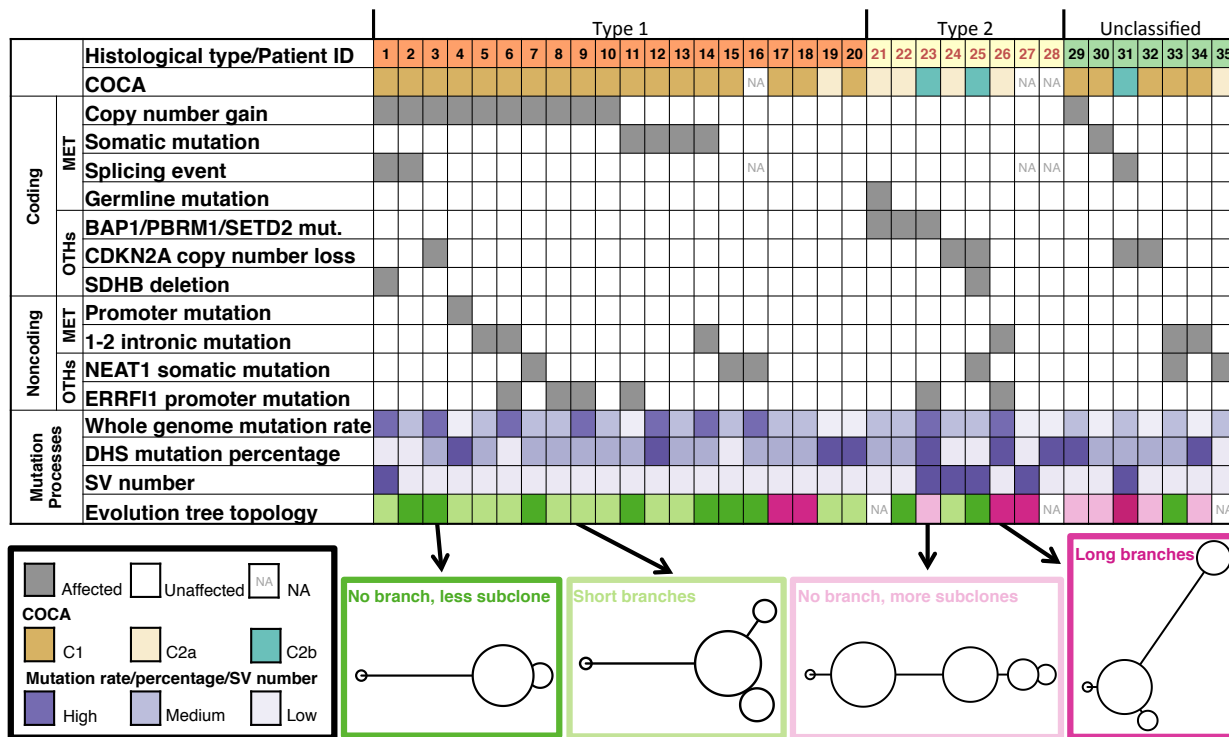


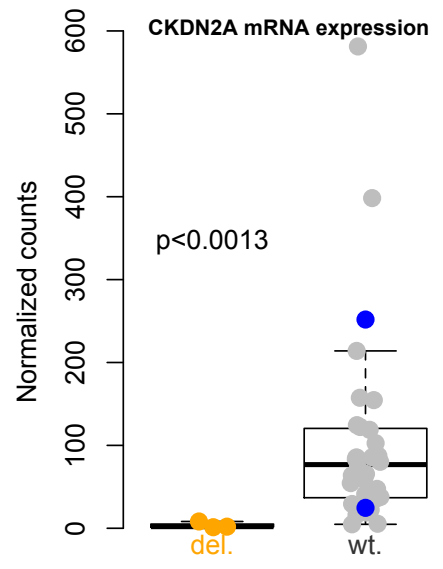
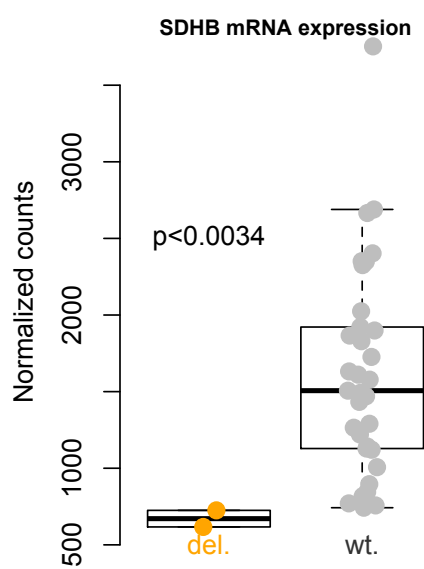
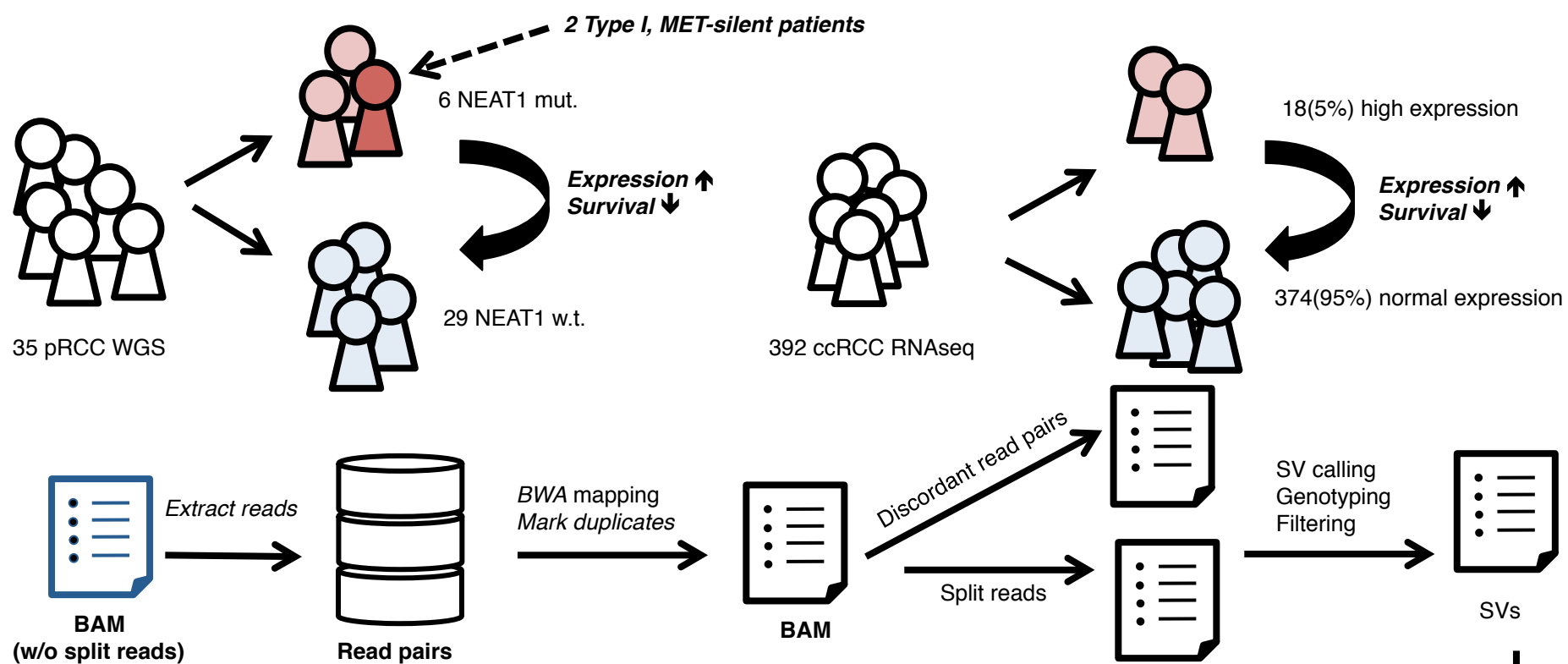


A



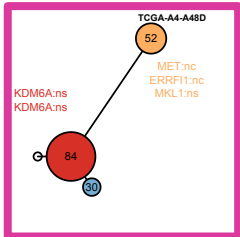
B



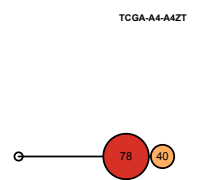


- 3 *CDKN2A* deletions
 2 *SDHB* deletions
 1 *STAG2* deletion
 1 *DNMT3A* deletion
 1 *EGFR* duplication
 1 *HIF1A* duplication
 ...

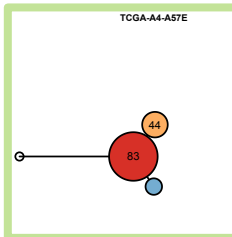
Prioritize



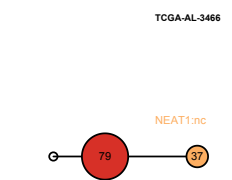
TCGA-AL-3473



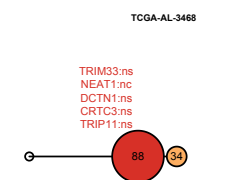
TCGA-AL-45DJ



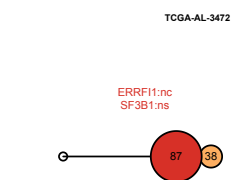
TCGA-B1-A47M



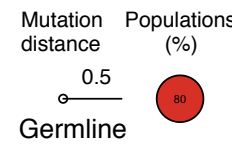
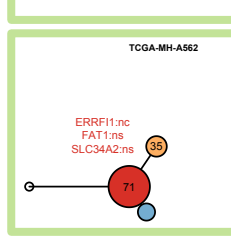
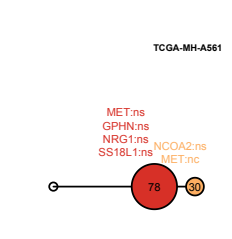
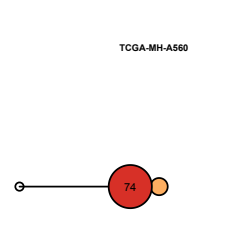
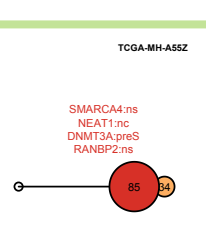
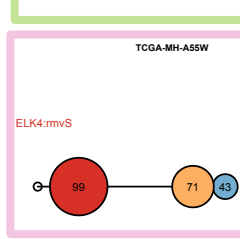
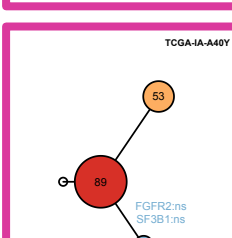
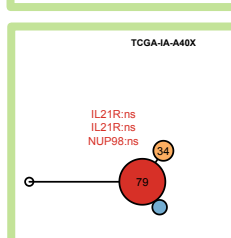
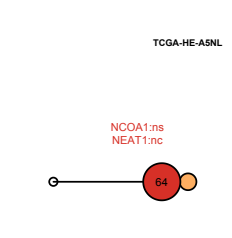
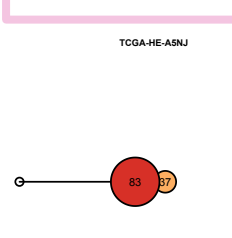
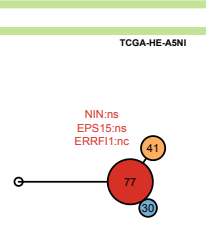
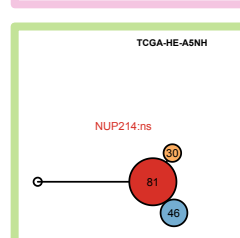
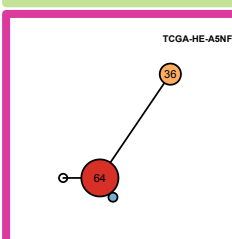
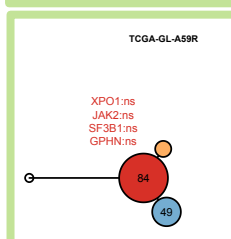
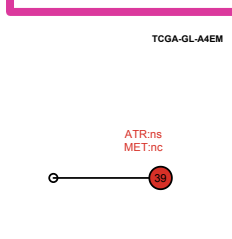
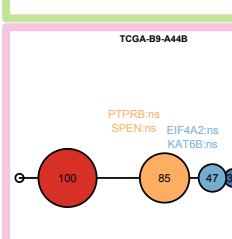
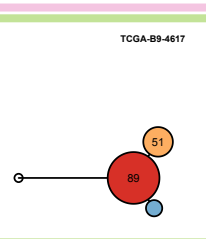
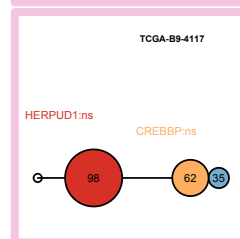
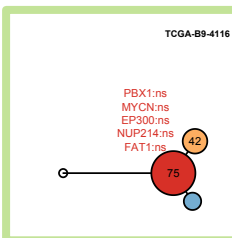
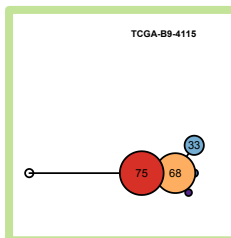
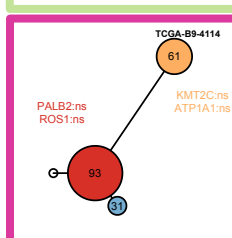
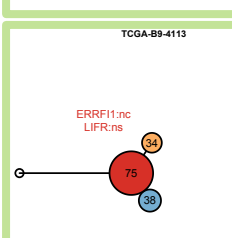
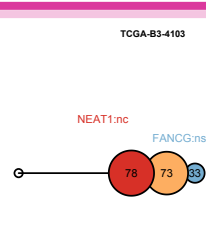
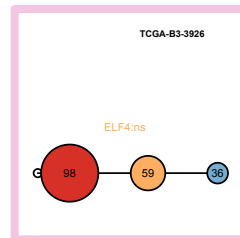
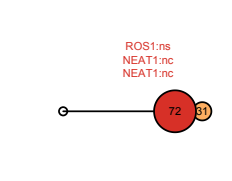
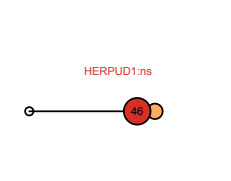
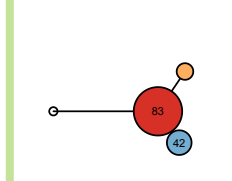
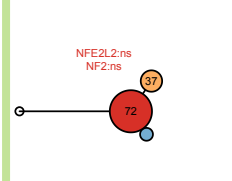
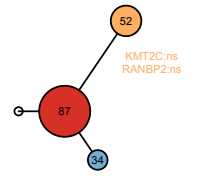
TCGA-B1-A47N



TCGA-B1-A47O



TCGA-B3-3925



Mutation% in early replicated regions

