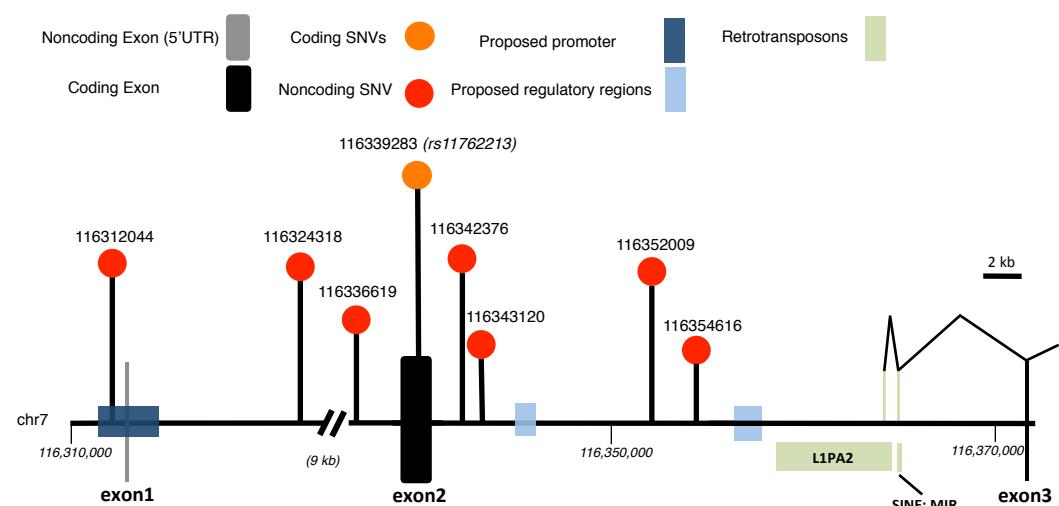
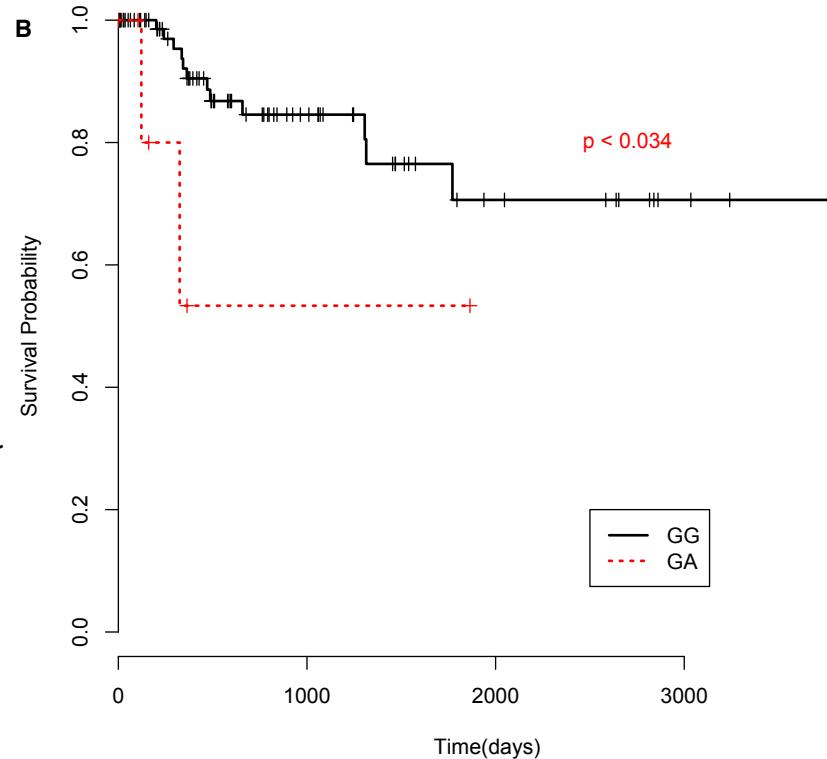


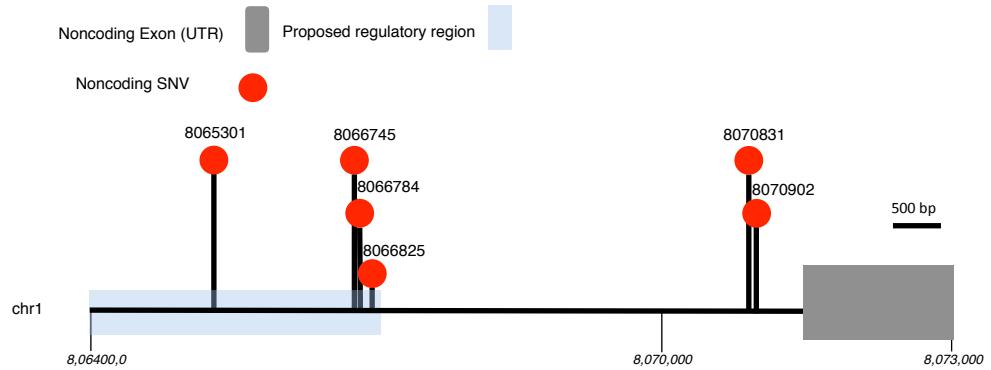
A MET



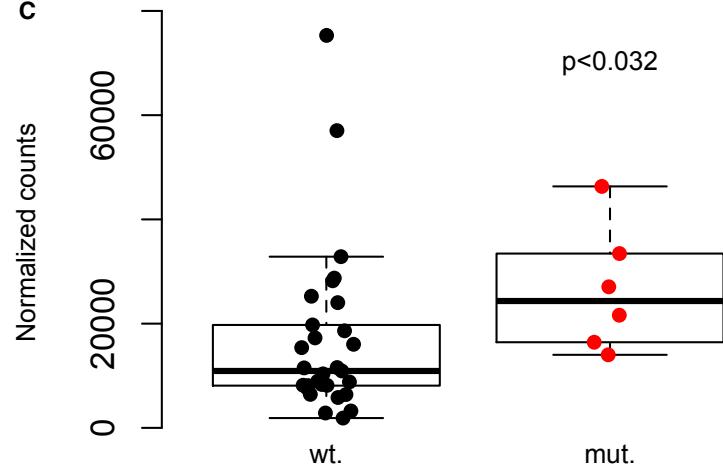
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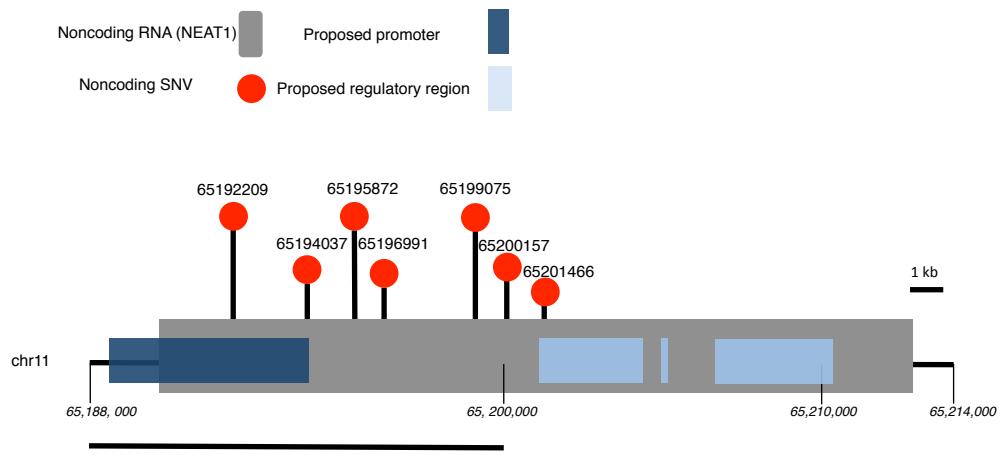
A ERF1



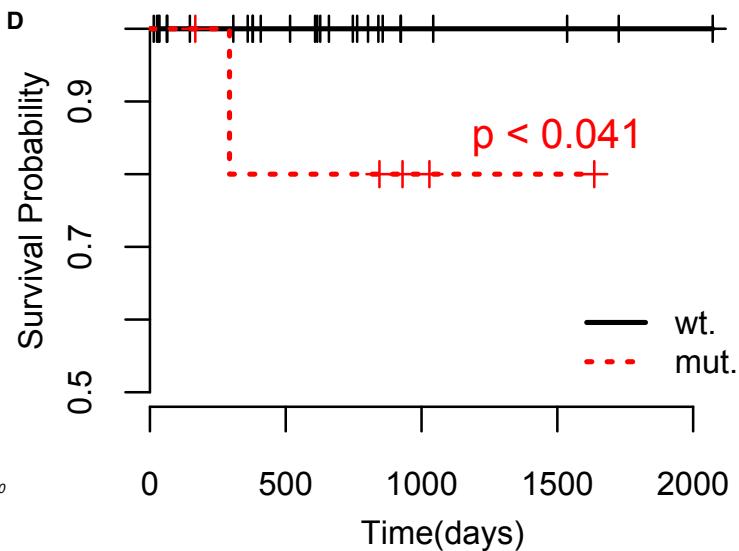
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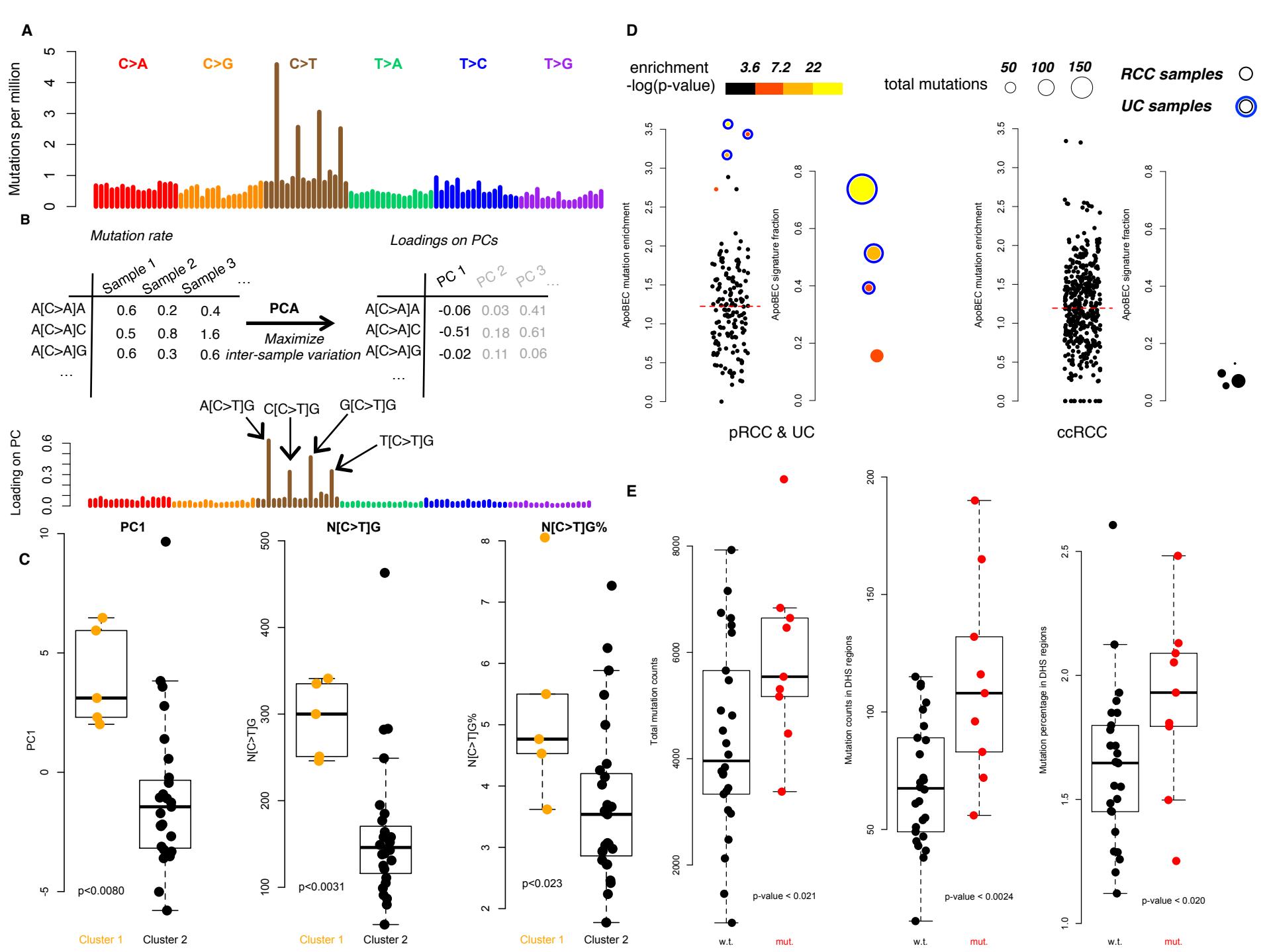


B NEAT1

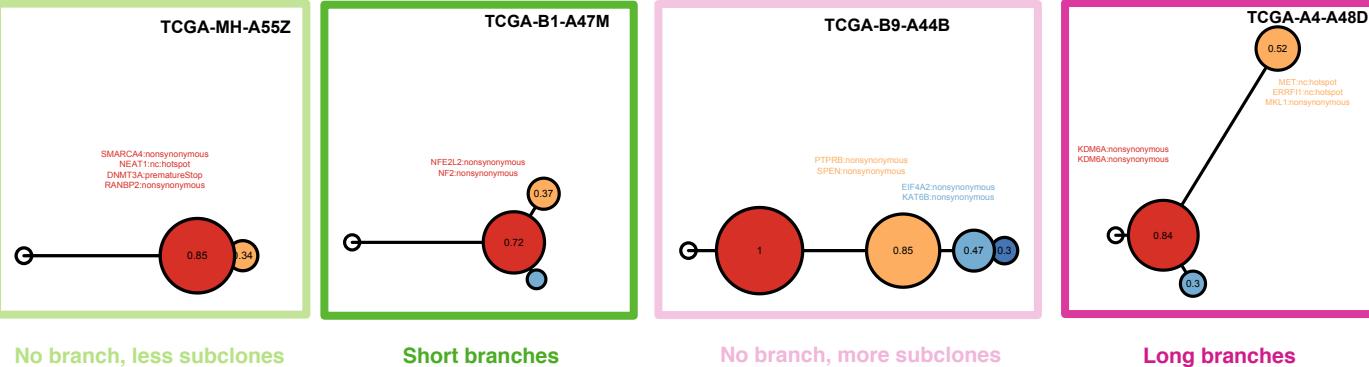
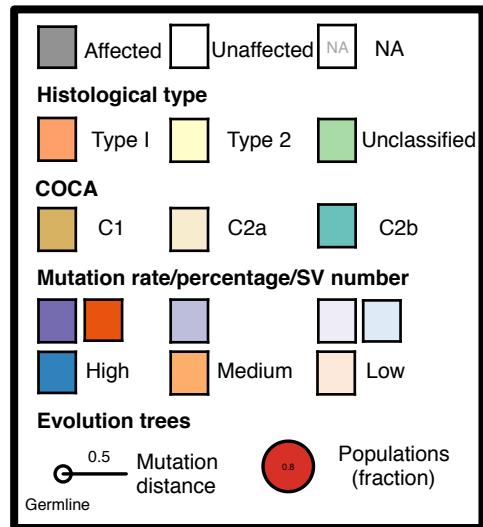


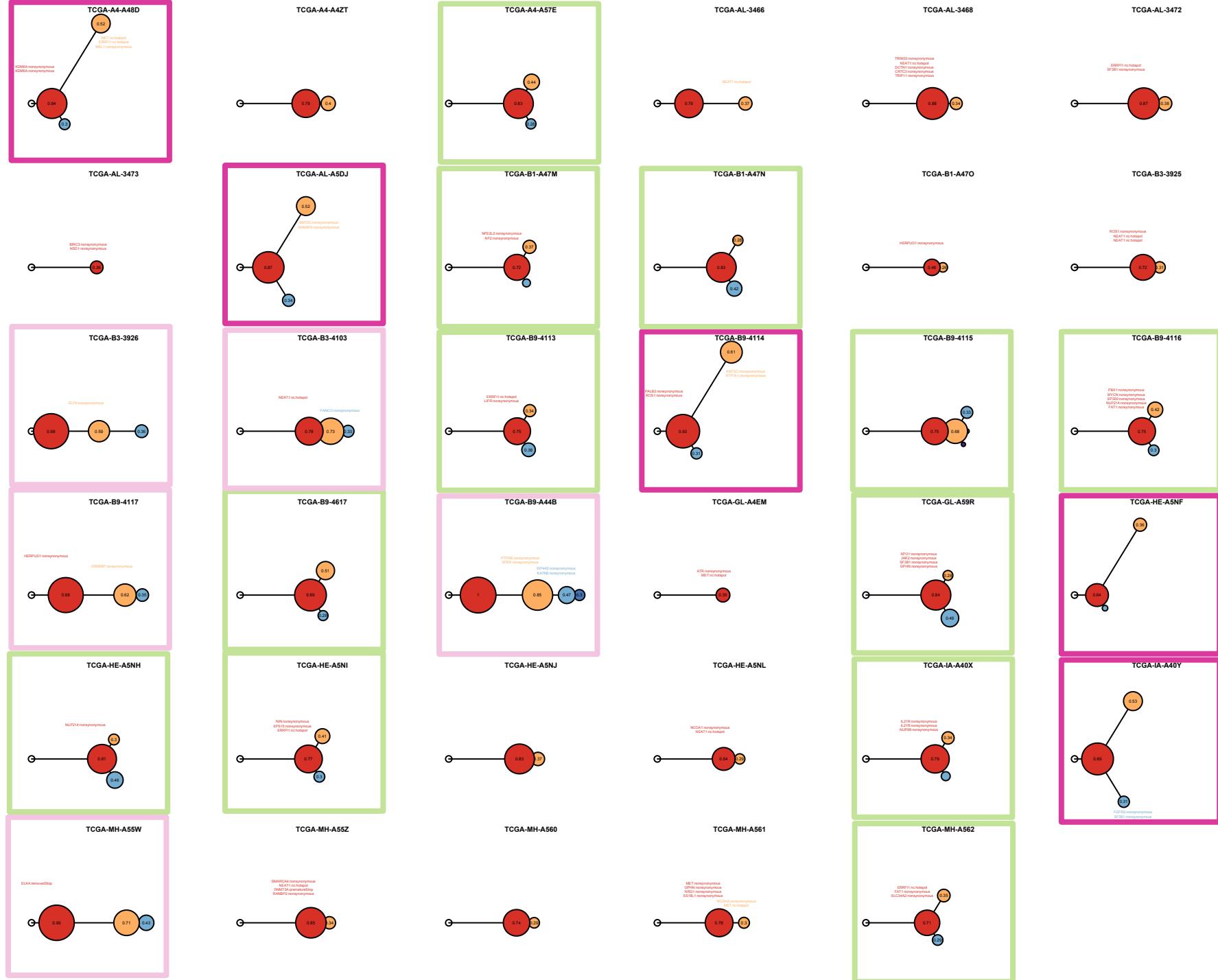
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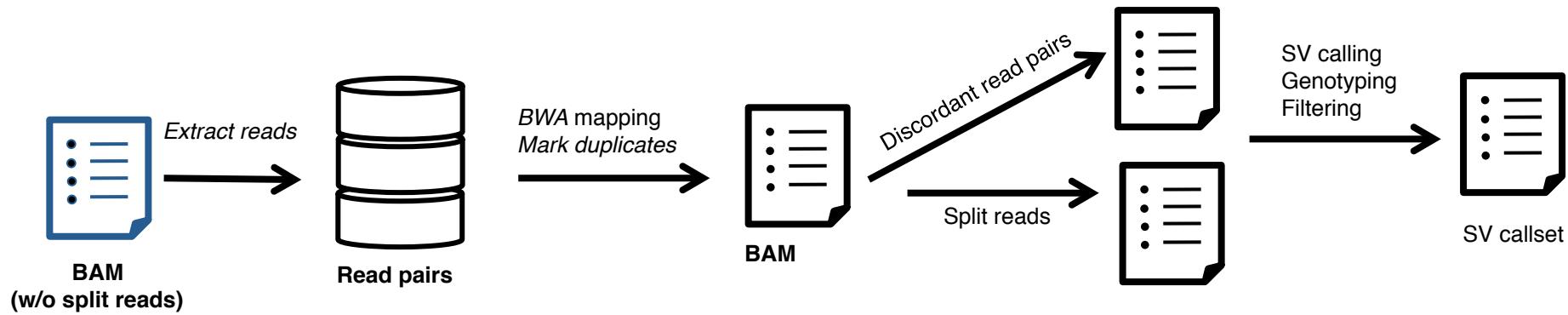
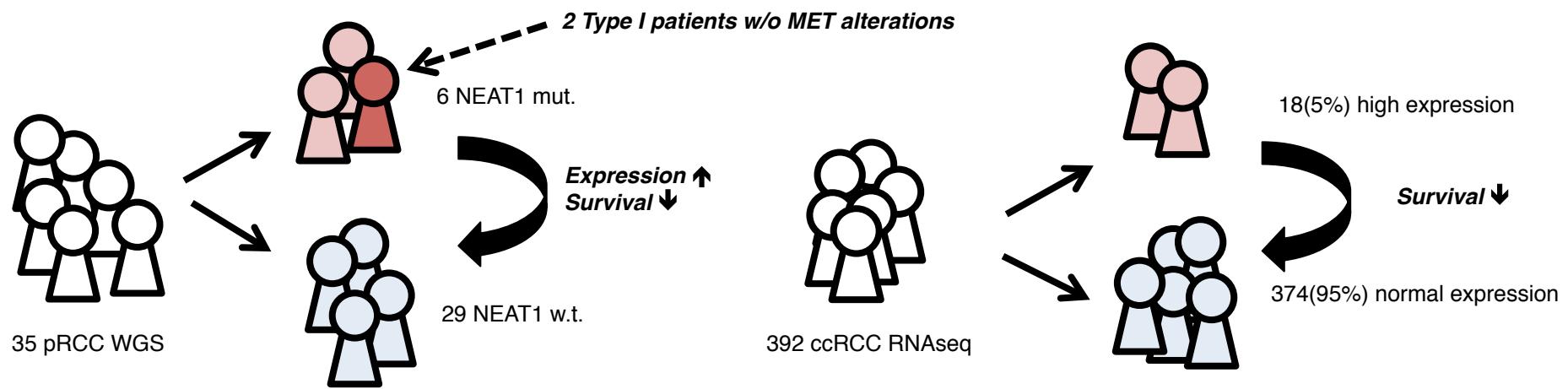




	Histological type/Patient ID	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35
	COCA															NA												NA	NA							
Coding	MET copy number gain																																			
	MET somatic mut.																																			
	MET splicing event																																			
	MET germline mut.																																			
	BAP1/PBRM1/SETD2 somatic mut.																																			
	CDKN2A copy number loss																																			
	SDHB deletion																																			
Noncoding	MET promoter mut.																																			
	MET 1-2 intronic mut.																																			
	NEAT1 somatic mut.																																			
	ERRFI1 promoter mut.																																			
Mutation Processes	Whole genome mutation rate																																			
	DHS mutation percentage																																			
	SV number																																			
	Evolution tree topology																																			







Worse OS

**Median OS (month):
36 v.s. 77**