

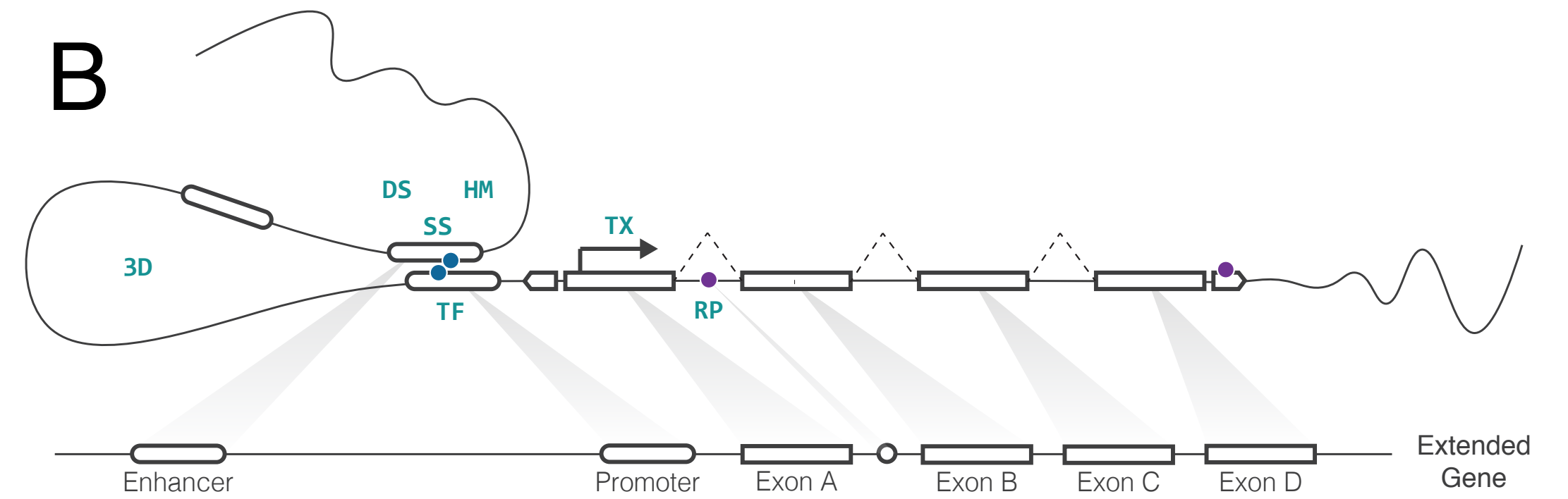
EC FigPackM

v170221

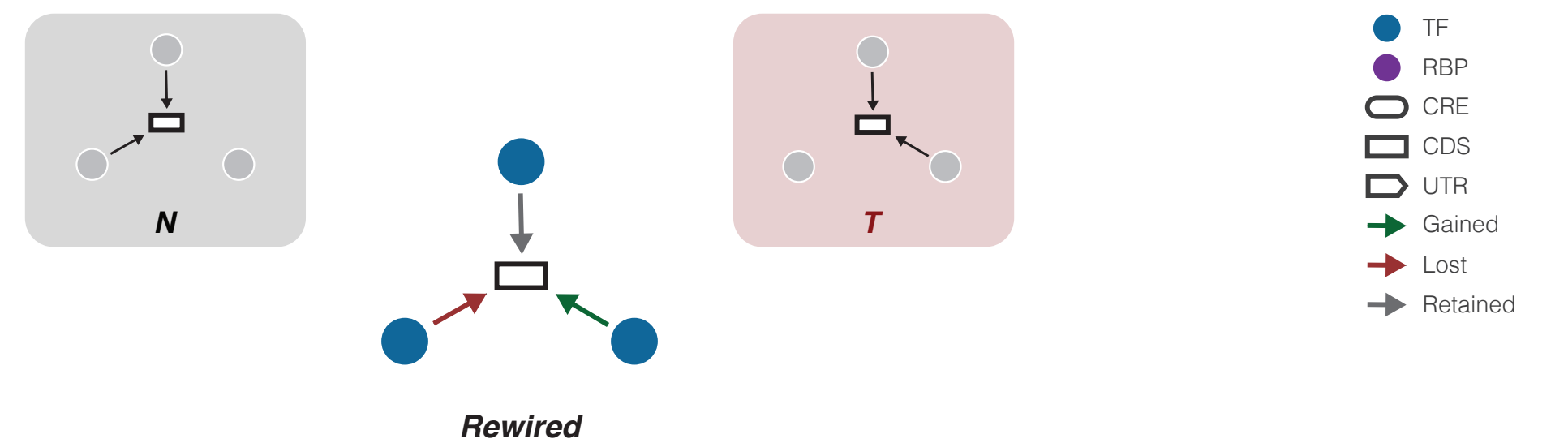
A

		K562	GM12878	HepG2	Liver	A549	IMR-90	MCF-7	MCF-10A	HeLa-S3
Transcription Factors TF	TF Total	207	99	95	7	31	9	52	4	59
	TFSS	129	71	71	6	23	6	37	3	39
	Chromatin Remodeller	17	6	6	1	2	1	2		6
	Cofactor	19	9	11		4		3		4
Histone Modifications HM	General (GTF)	15	7	6		4	1	3	1	9
	Histone	12	11	11	7	11	28	5	3	11
RNA-binding Proteins RP	eCLIP	89		70						
shRNA/siRNA Knockdown KD	TF	85		61				2		
	RBP	234		225						
3D Chromatin Structure 3D	ChIA-PET	3	3	1				4		2
	Hi-C	▲	▲			●	▲	▲	▲	
Chromatin Accessibility DS	DNase-seq	●	●	●		●	▲	●		●
	STARR-seq	●	●	▲				●		
Enhancers SS	WGBS	●	●	●	▲		▲	▲		
	RRBS	●	●	●	●	●	●	●	●	●
Methylation ME	RNA-seq	●	●	●	●	●	●	●	▲	●
	RAMPAGE	●	●							
Transcription TX	Repli-seq/chip	●	●	●			●	●		●
	SNV	▲	▲					▲		▲
Cell Line WGS WG	SV	▲	▲					▲		▲
	Cohort	150			82	197		116		
Patient Data PD	Expression	173		373		517		1100		546
		<i>T</i>	<i>N</i>	<i>T</i>	<i>N</i>	<i>T</i>	<i>N</i>	<i>T</i>	<i>N</i>	<i>T</i>
		Blood		Liver		Lung		Breast		Cervix

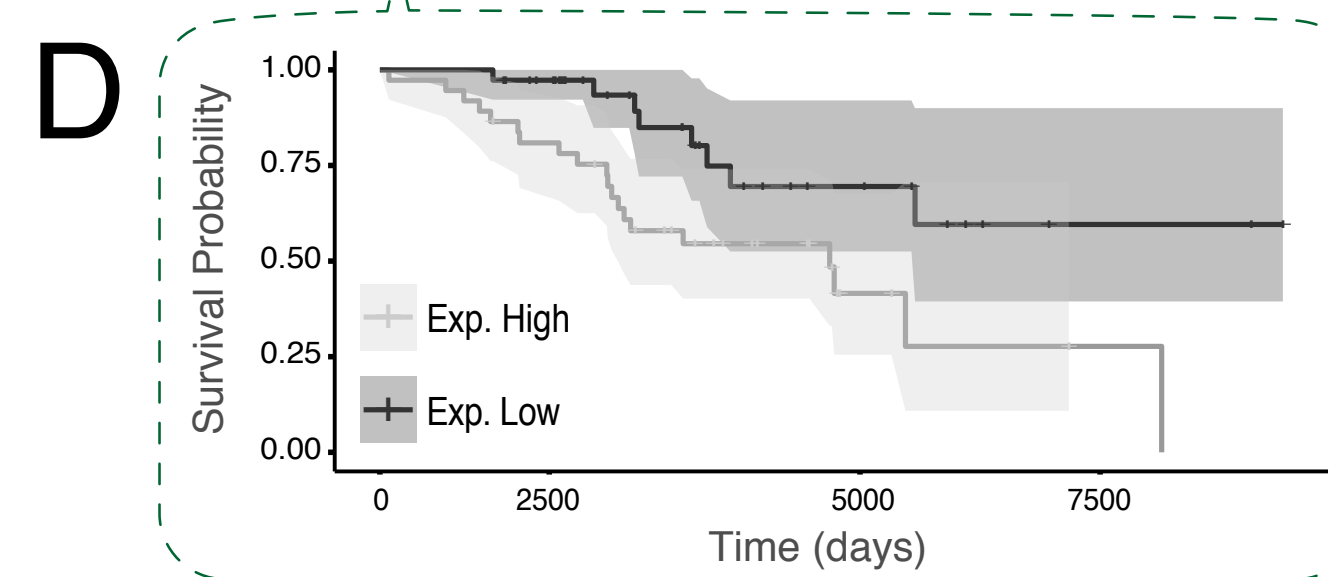
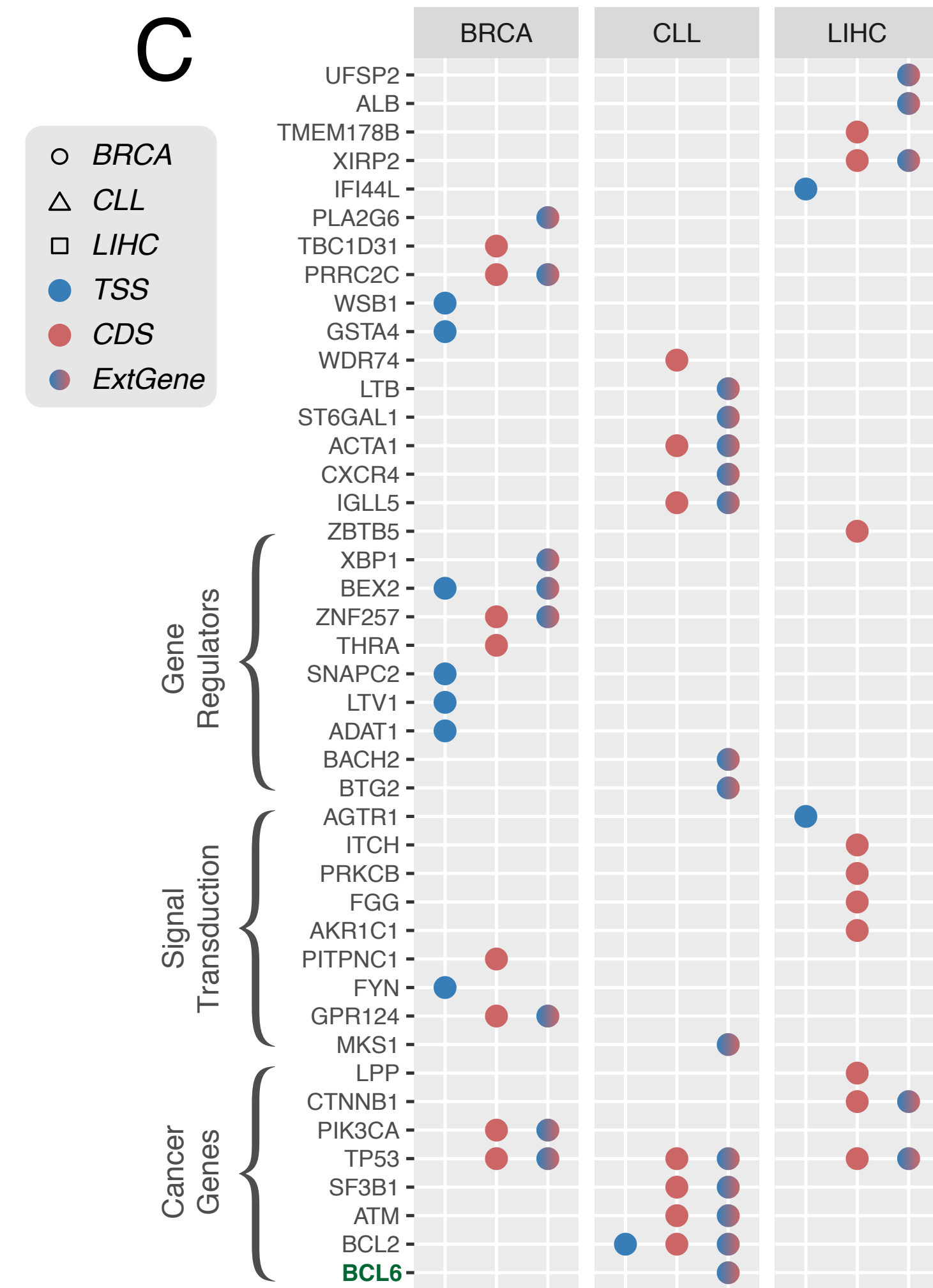
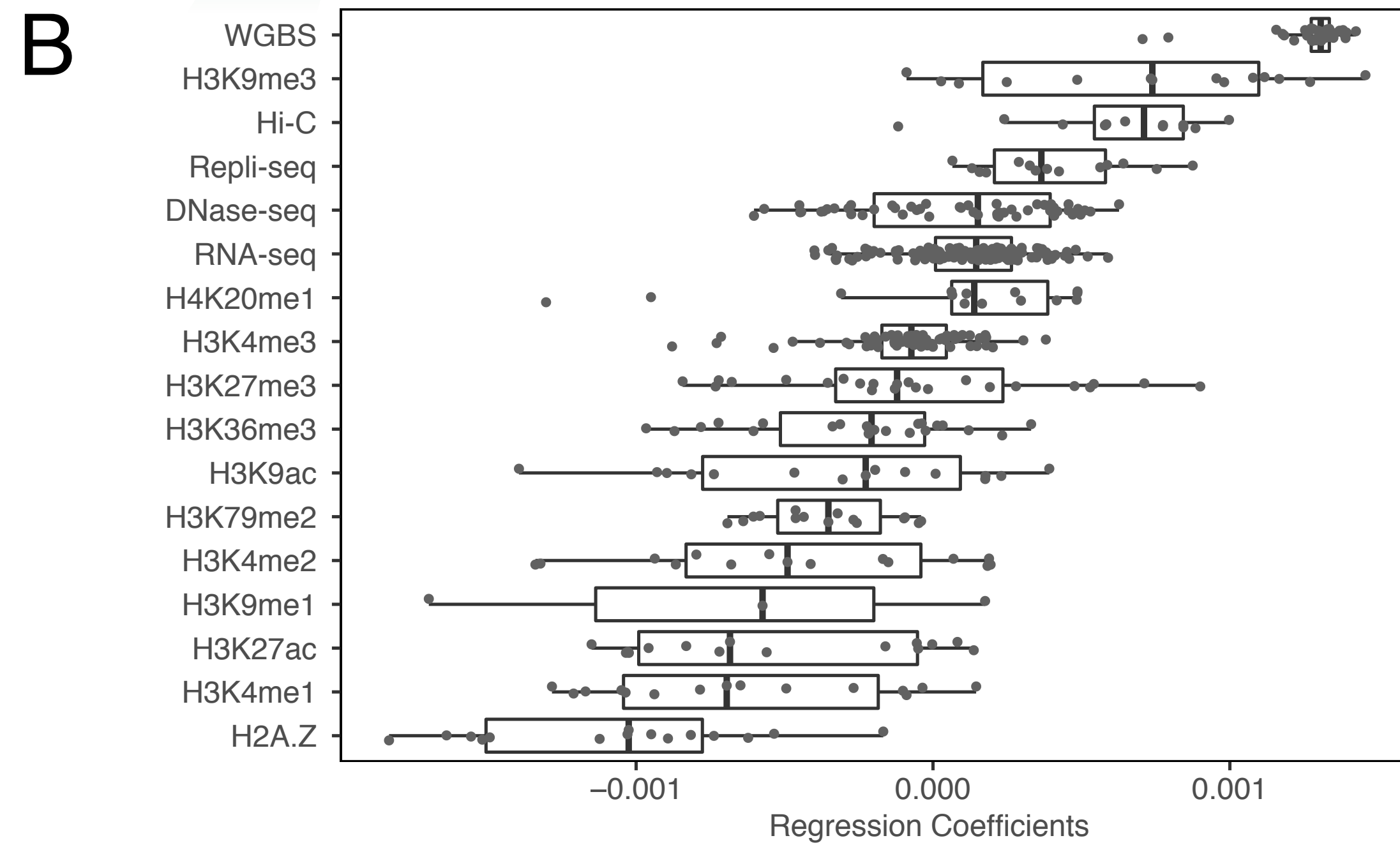
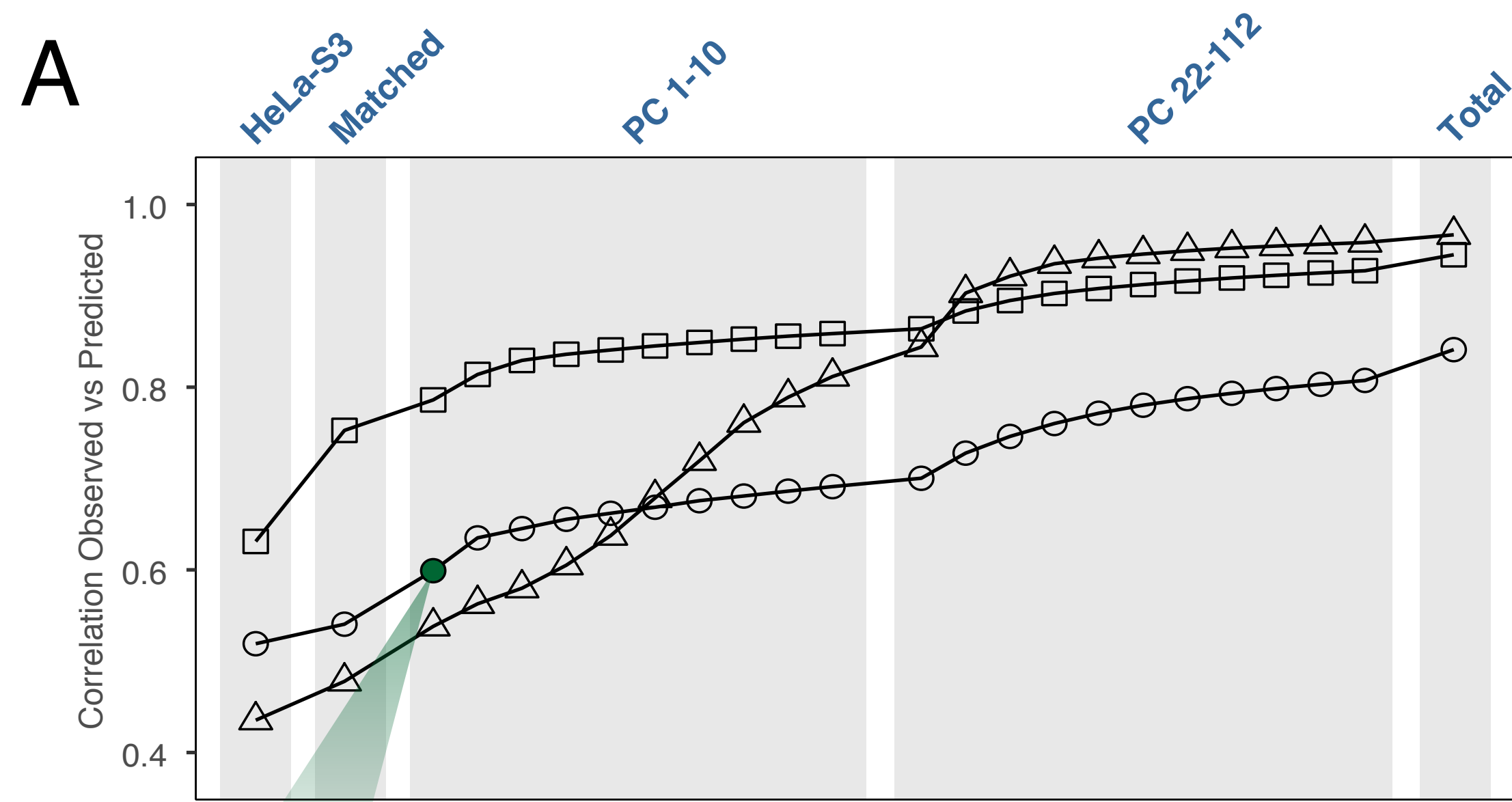
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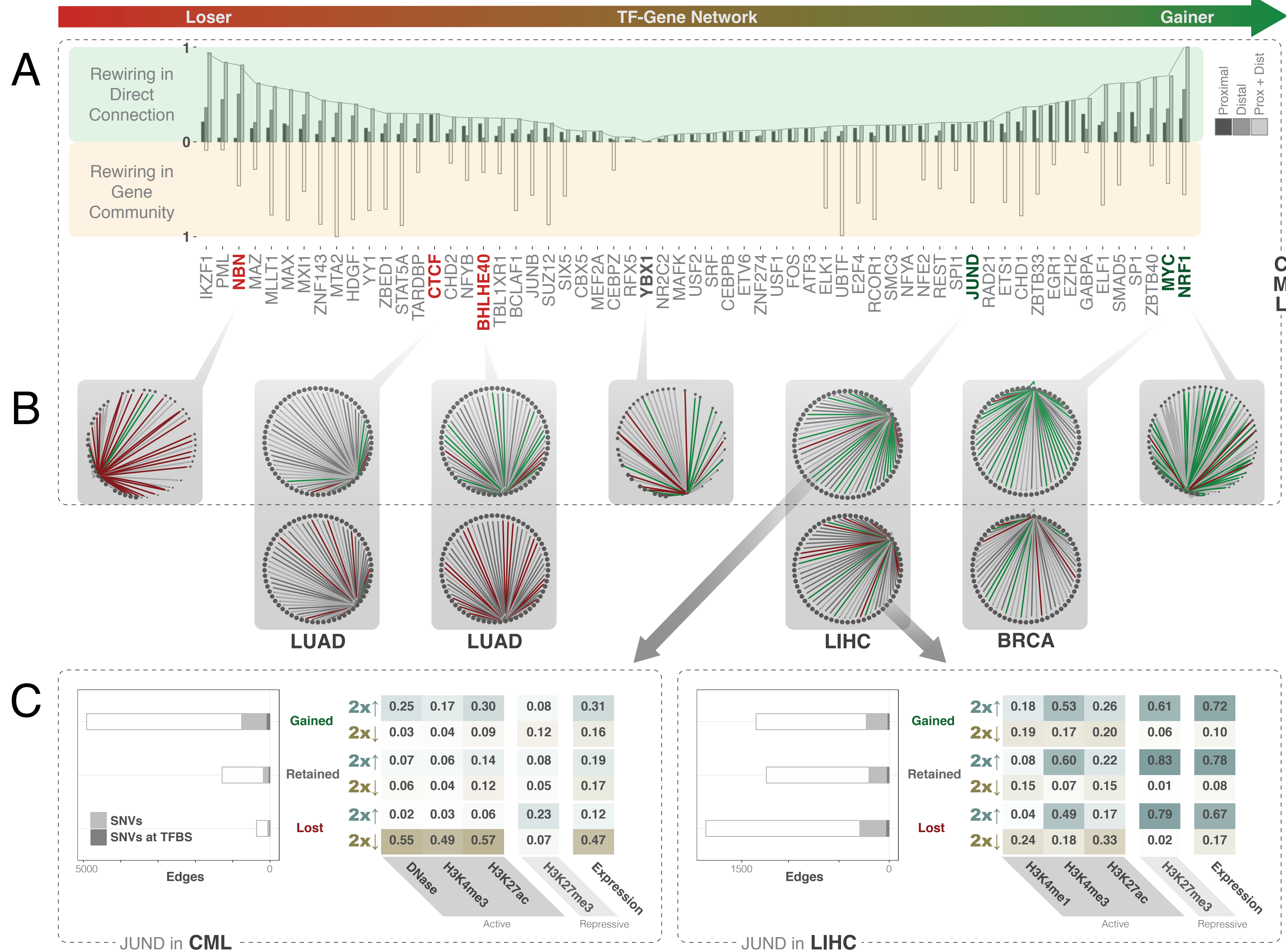


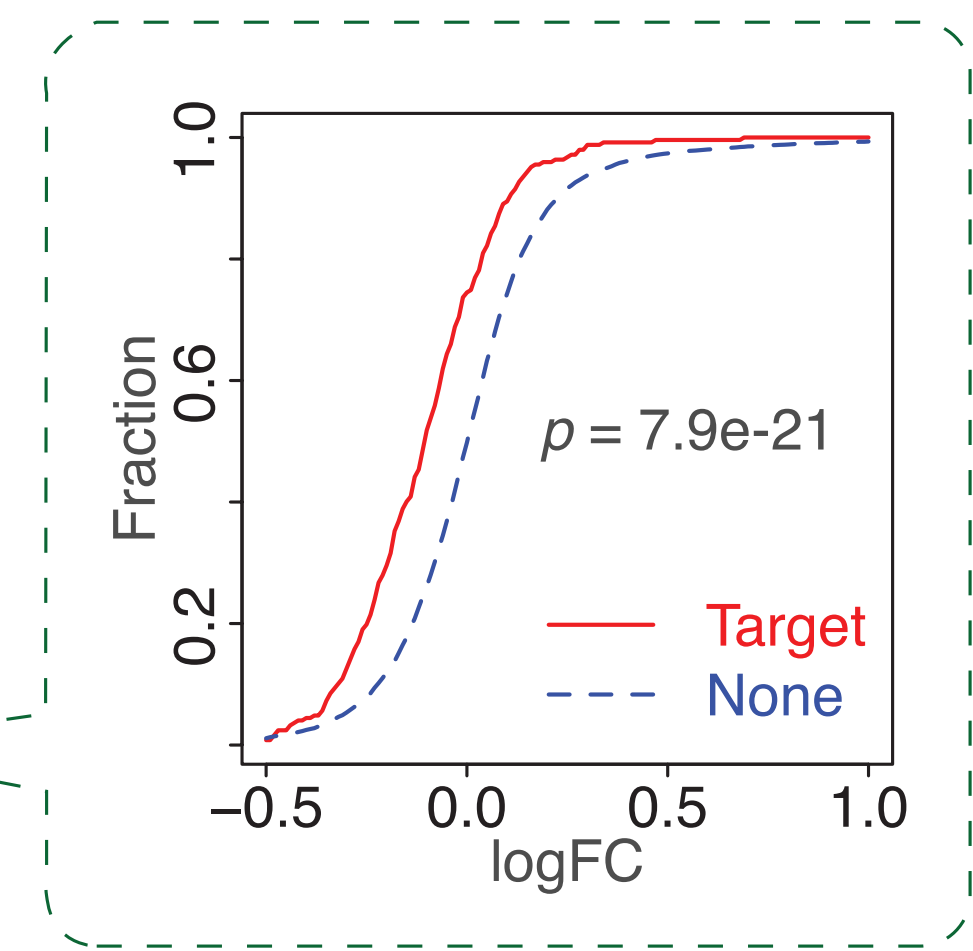
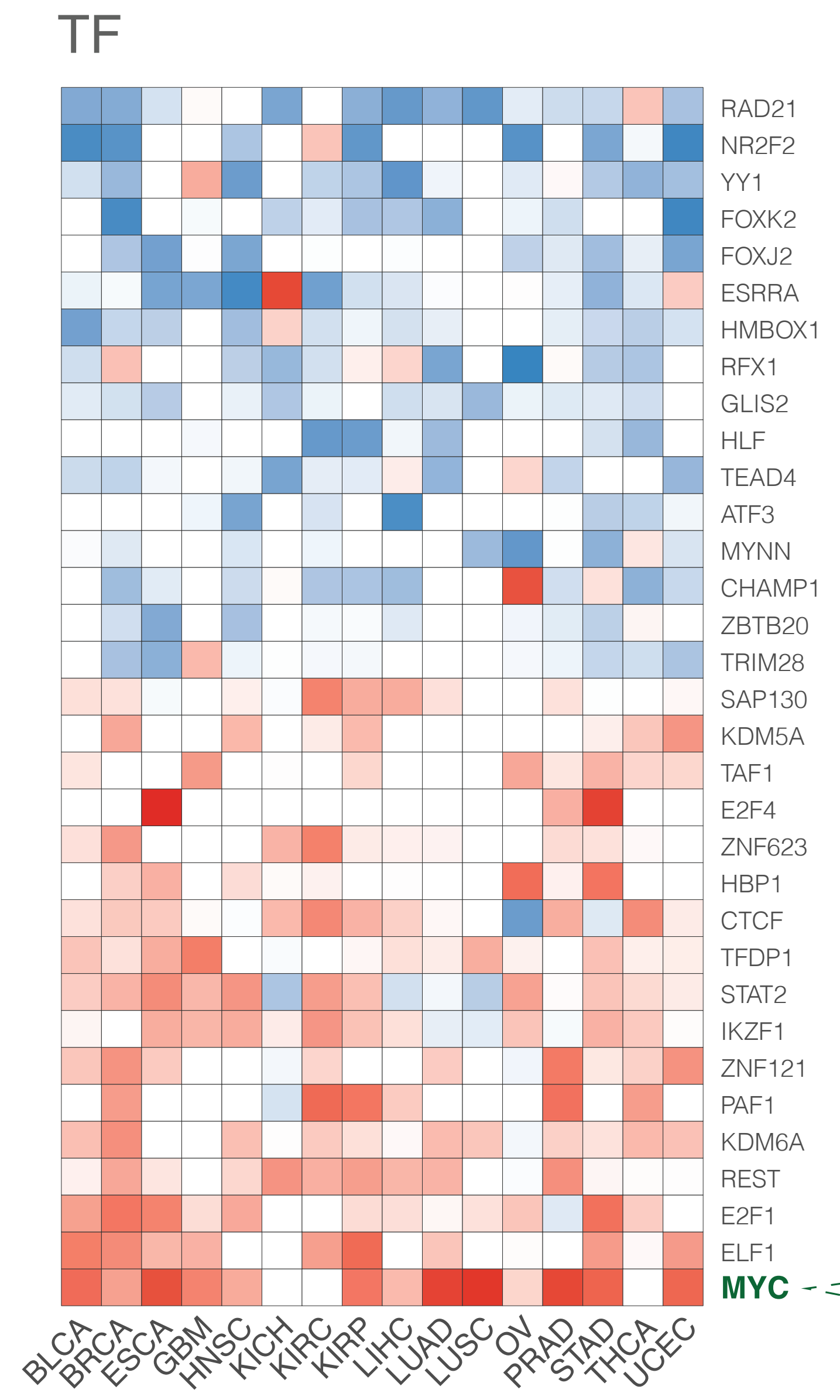
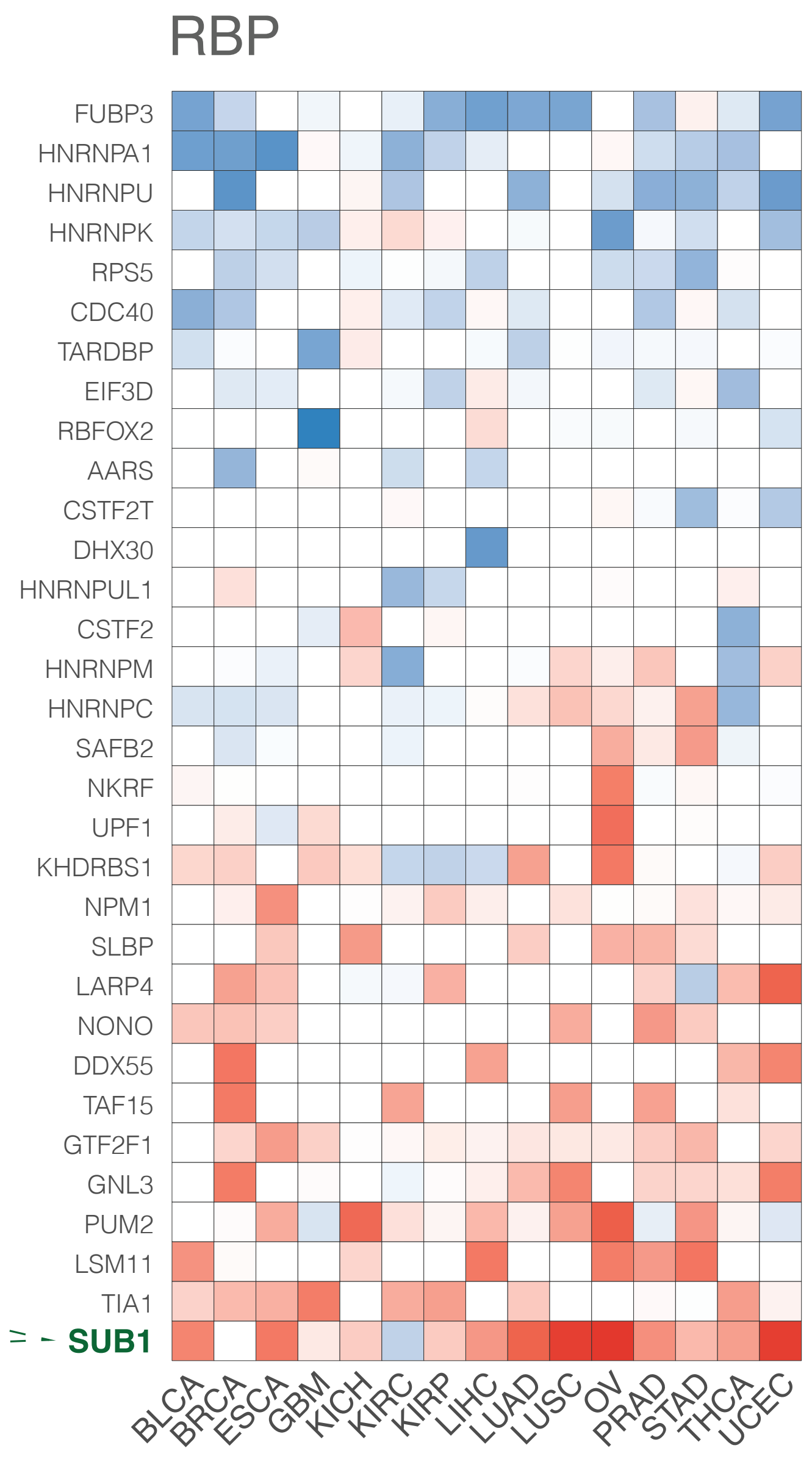
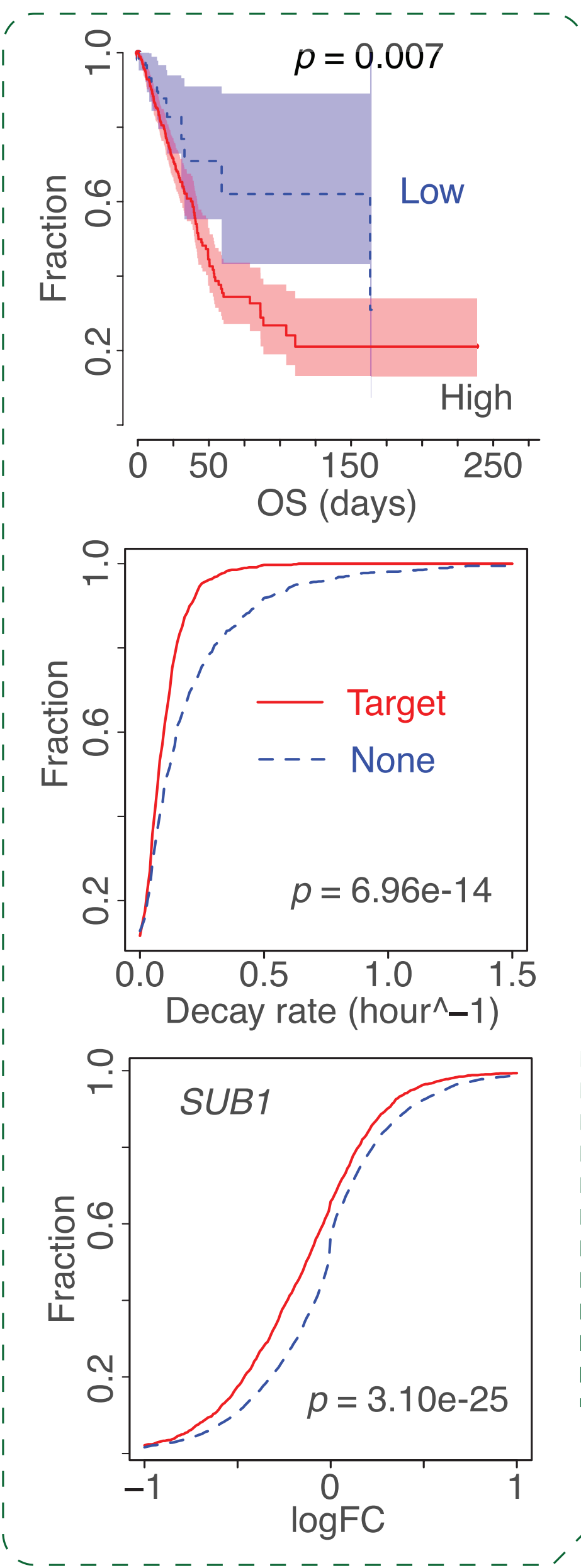
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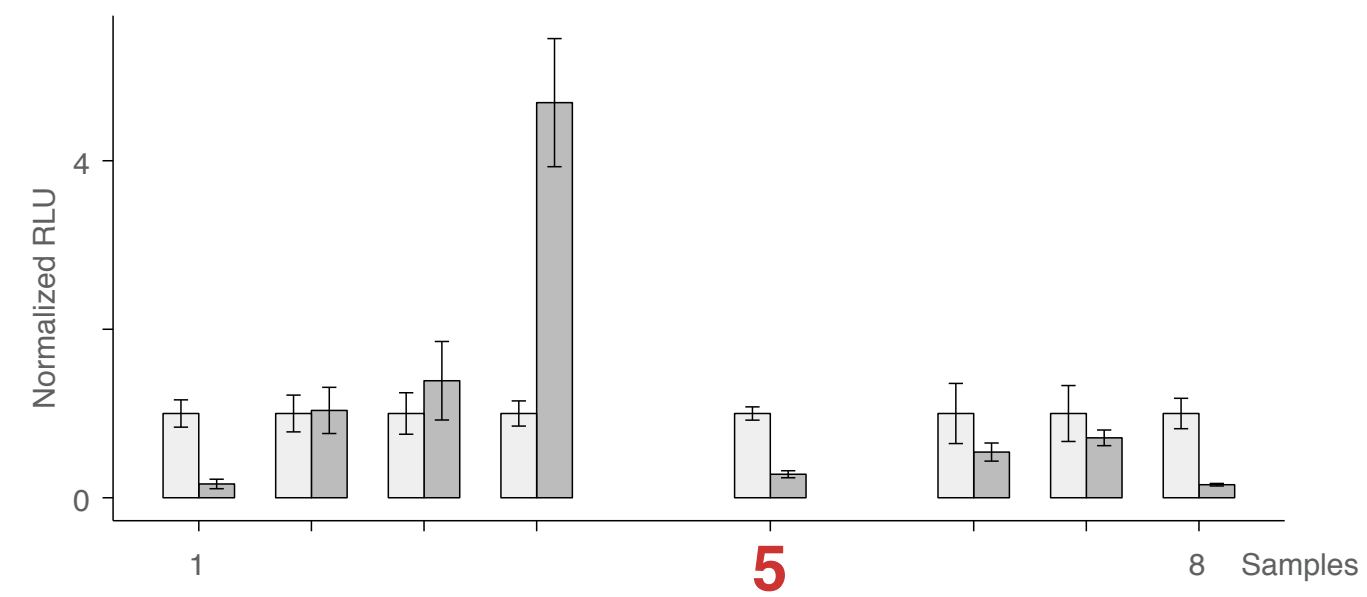
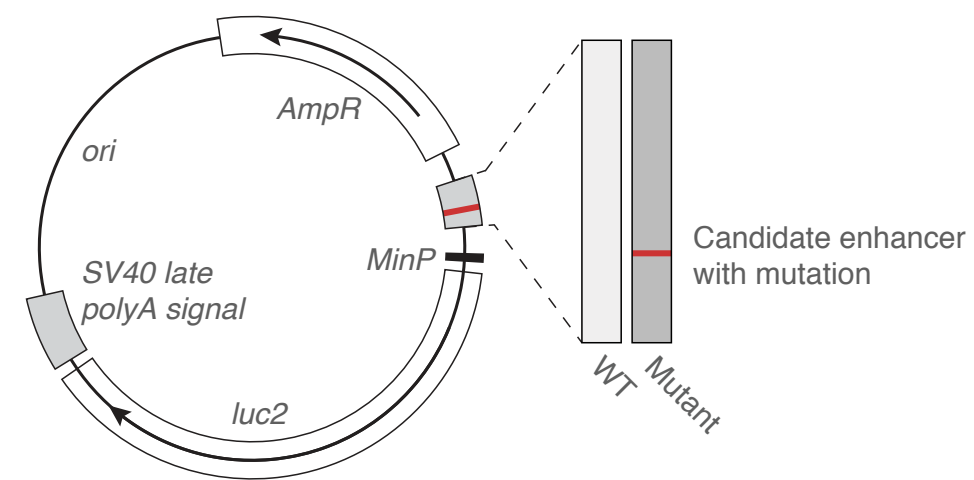
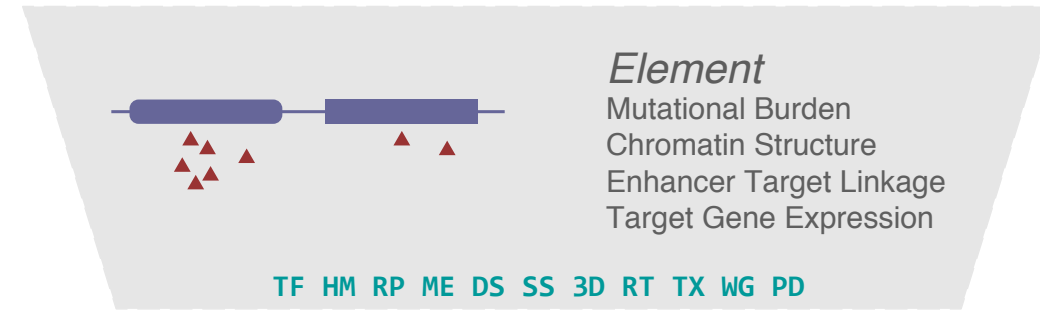
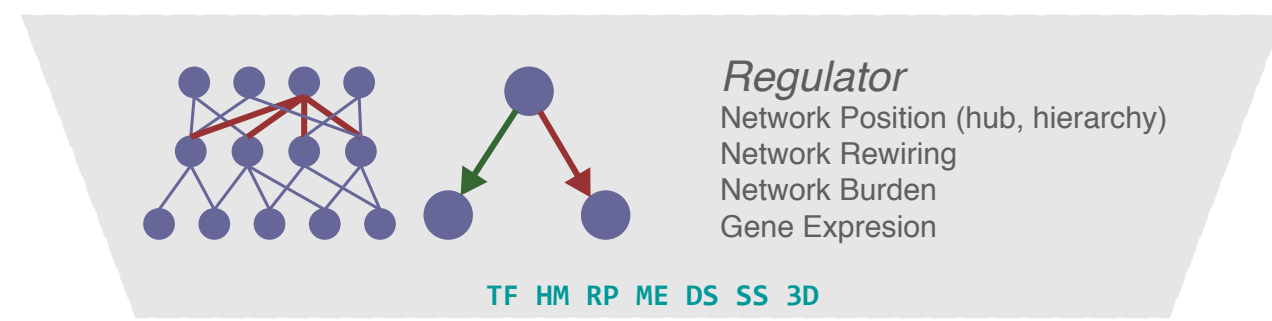


- TF
- RBP
- CRE
- CDS
- UTR
- Gained
- Lost
- Retained







A**B**