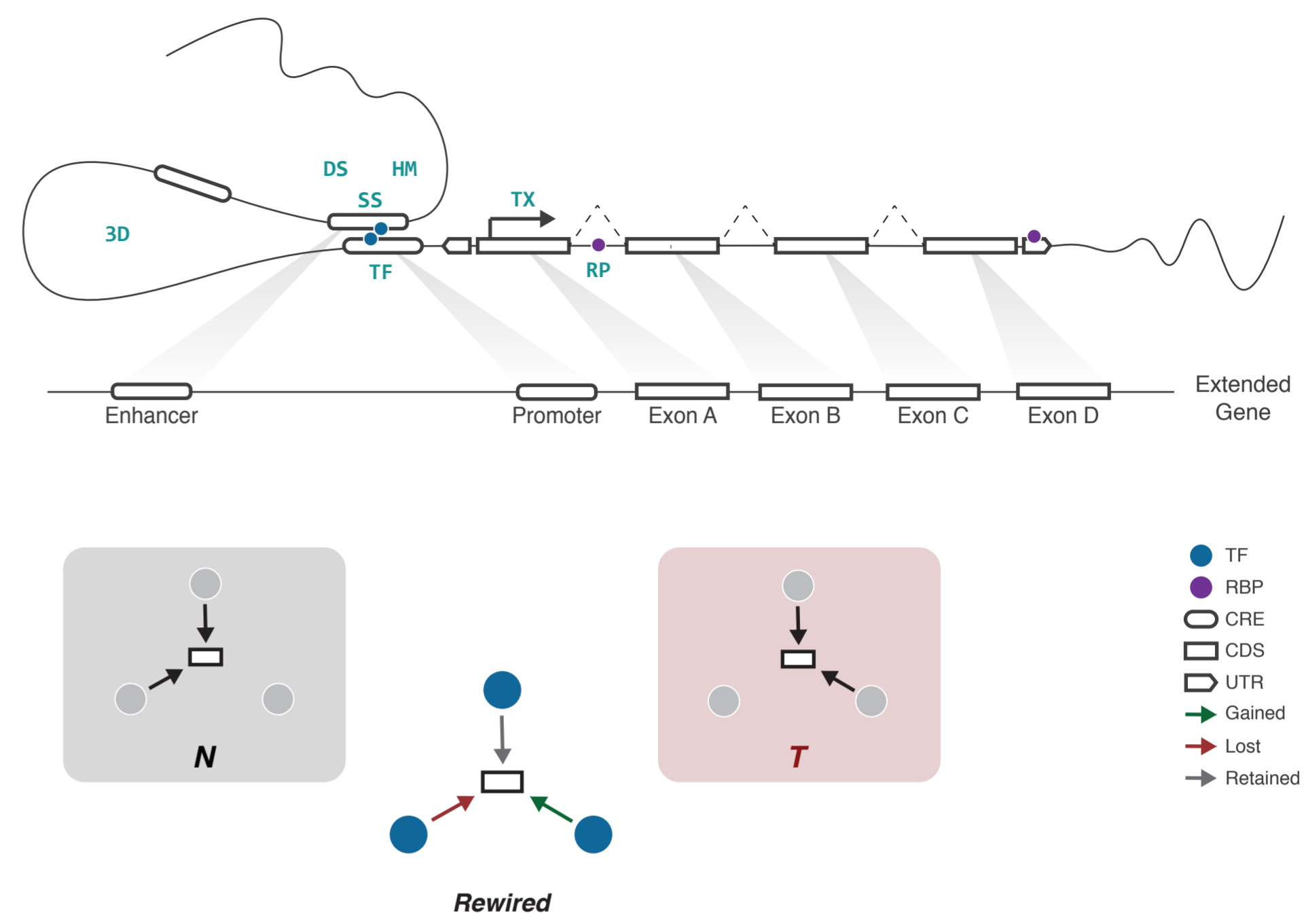
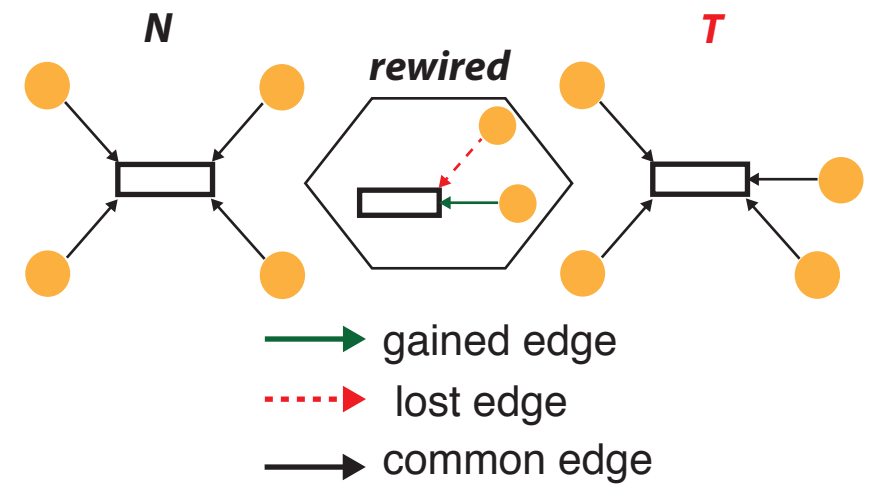
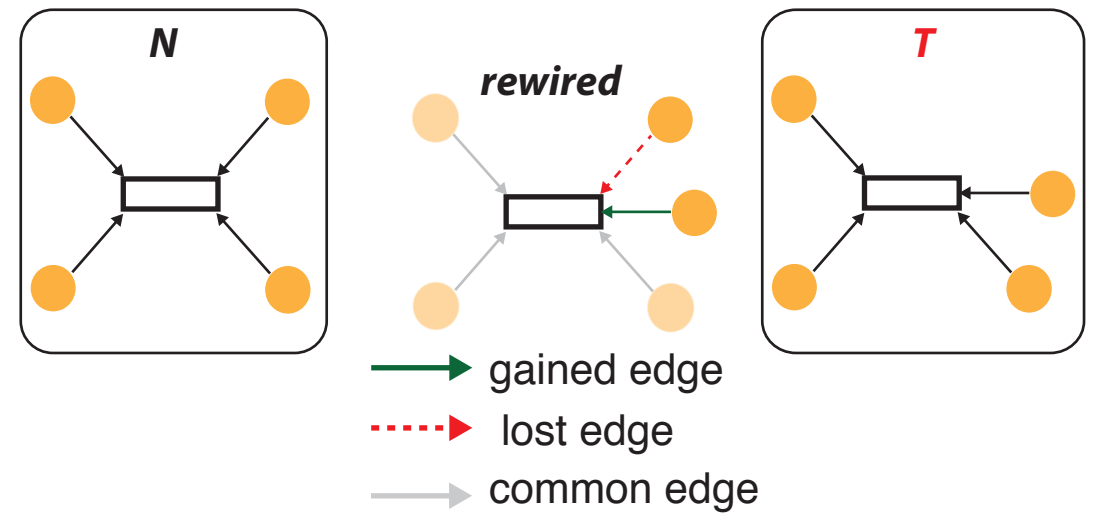
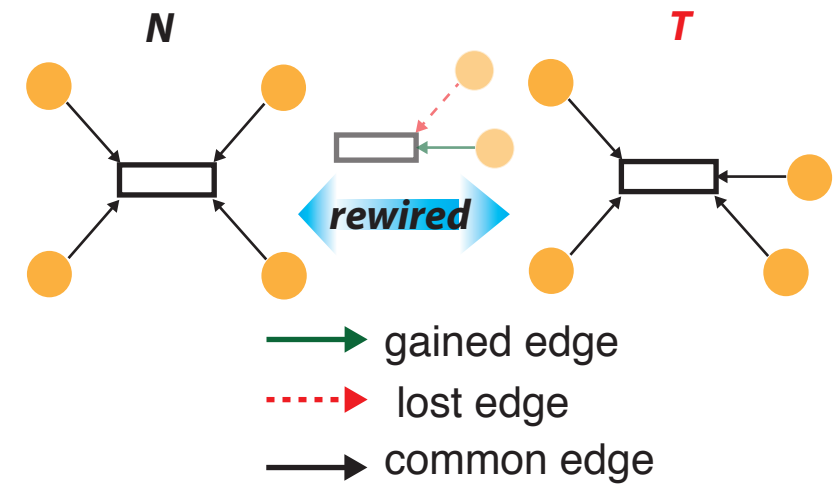


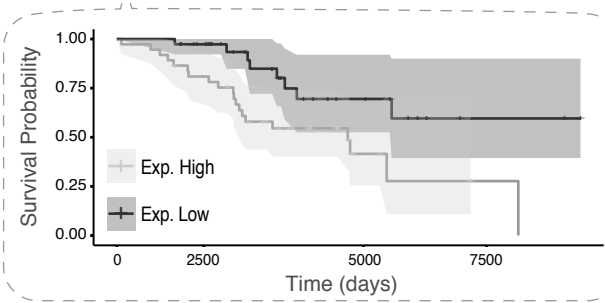
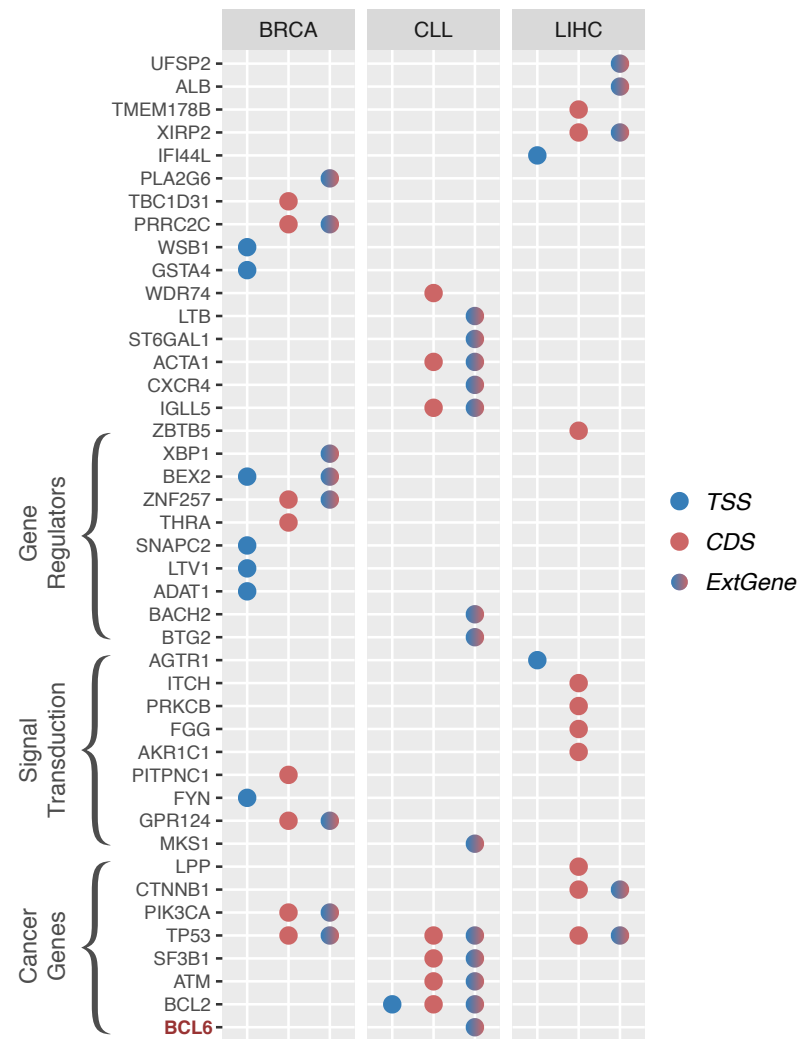
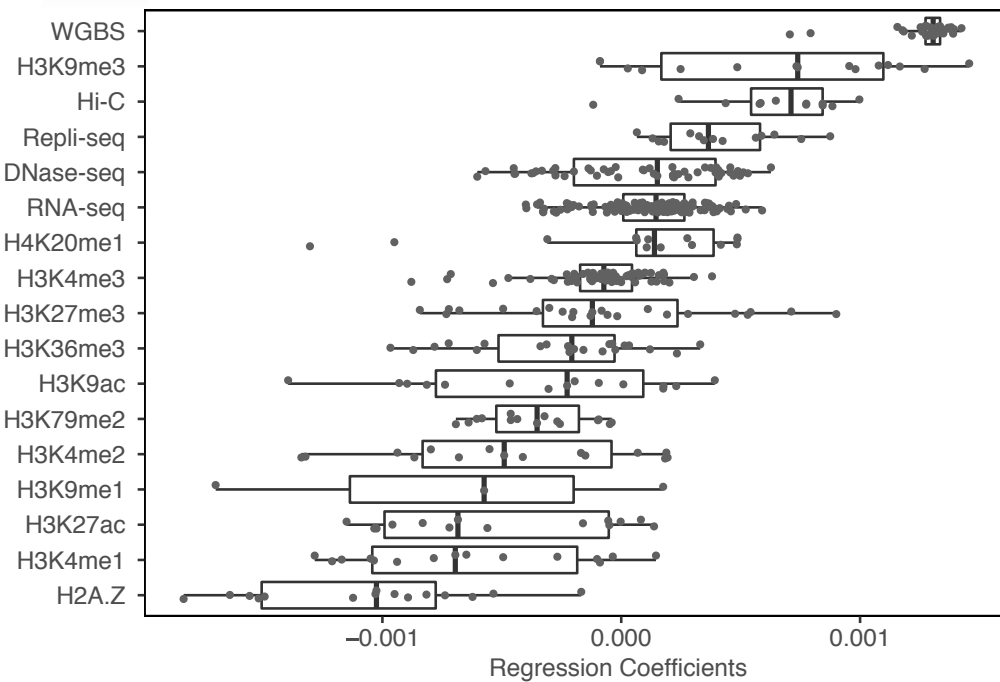
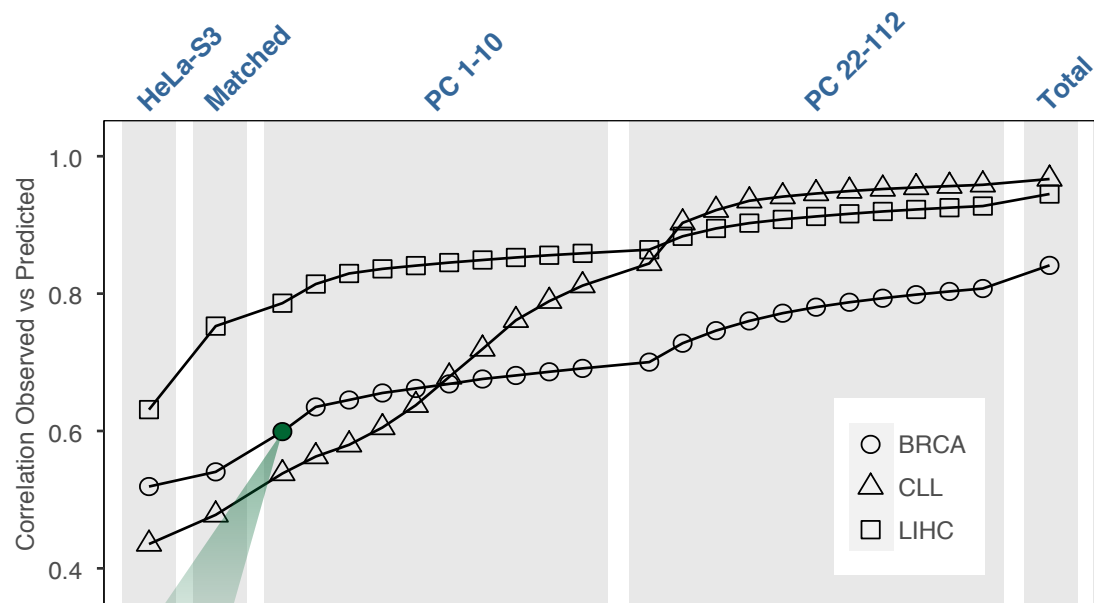
# MAIN FIGURES

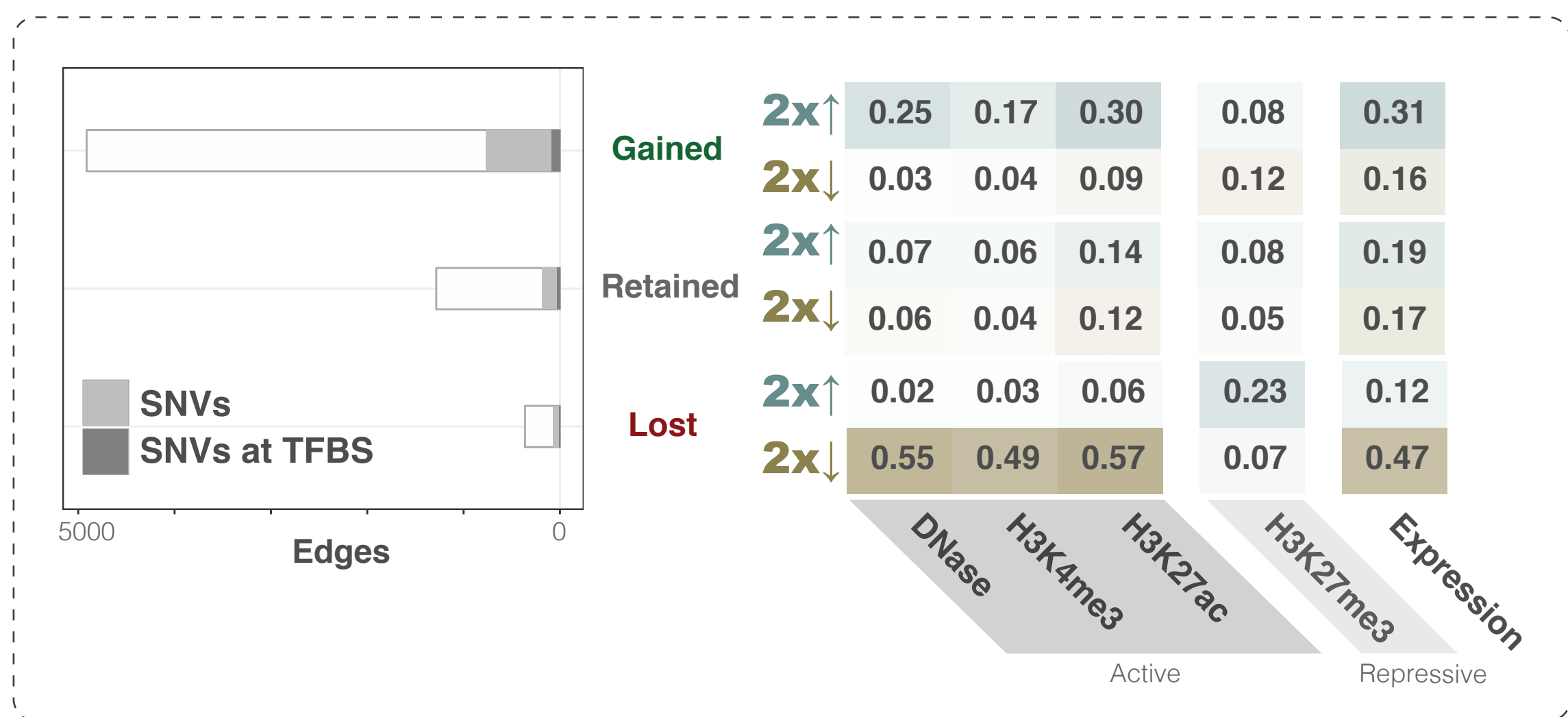
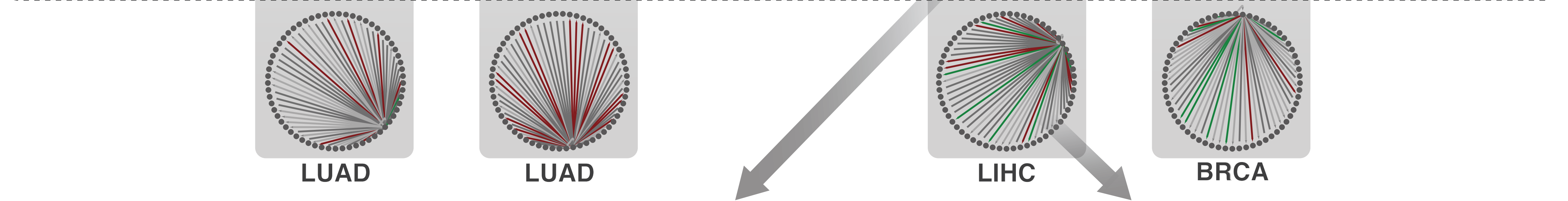
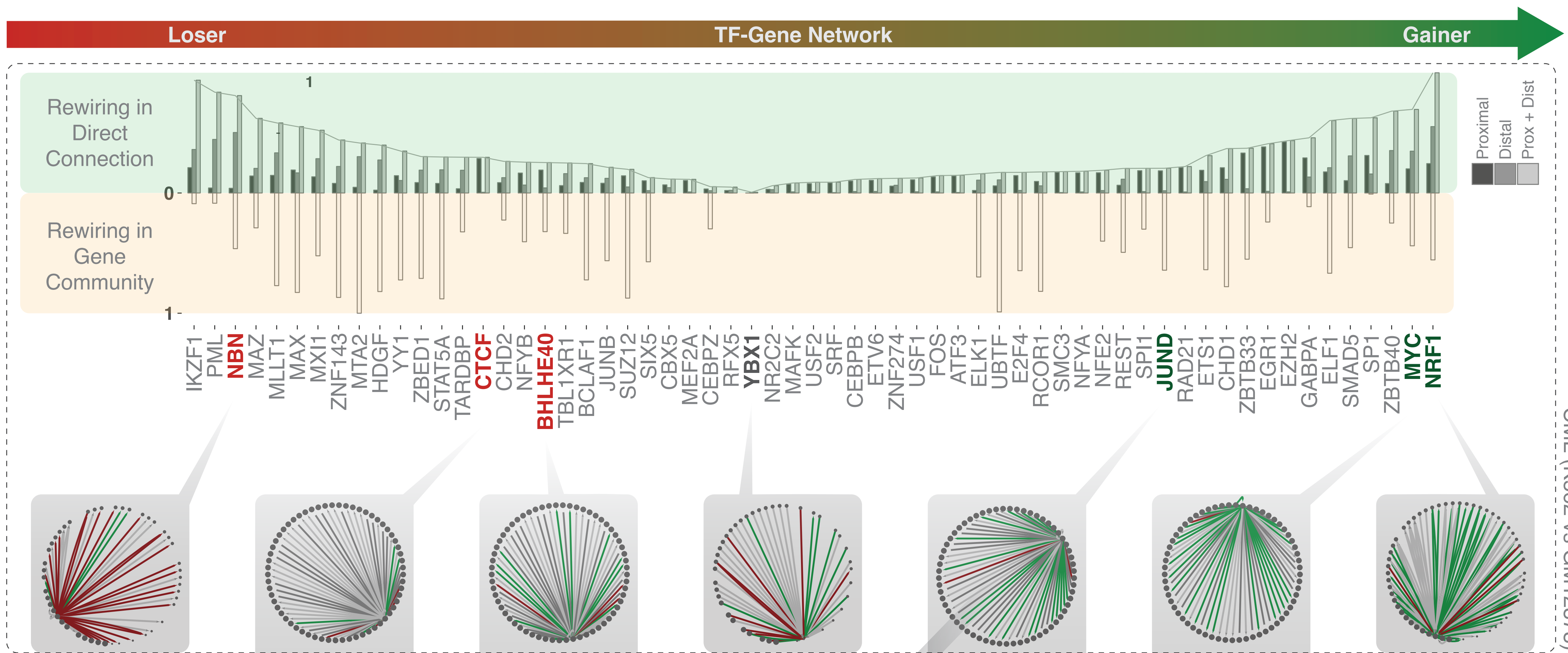
		<i>K562</i>	<i>GM12878</i>	<i>HepG2</i>	<i>Liver</i>	<i>A549</i>	<i>IMR-90</i>	<i>MCF-7</i>	<i>MCF-10A</i>	<i>HeLa-S3</i>
Transcription Factors <b>TF</b>	<b>TF Total</b>	207	99	95	7	31	9	52	4	59
	TFSS	129	71	71	6	23	6	37	3	39
	Chromatin Remodeller	17	6	6	1	2	1	2		6
	Cofactor	19	9	11		4		3		4
	General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

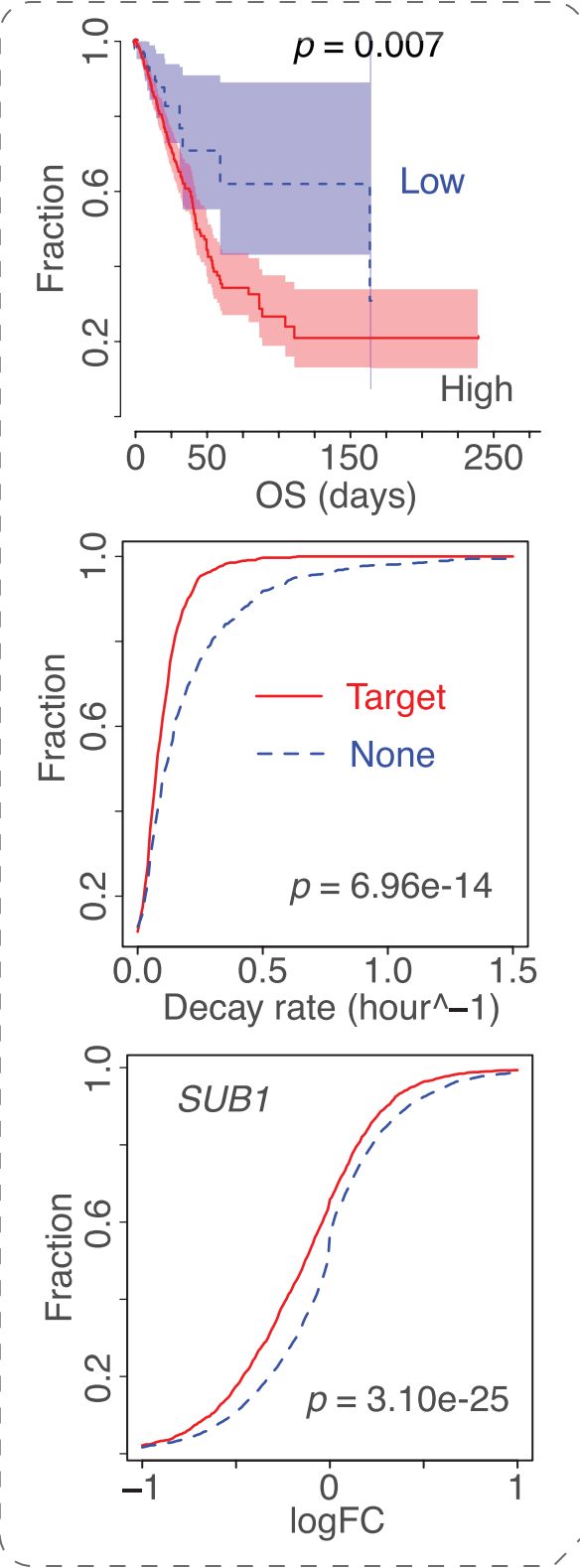


- TF
- RBP
- CRE
- CDS
- ▭ UTR
- Gained
- Lost
- Retained

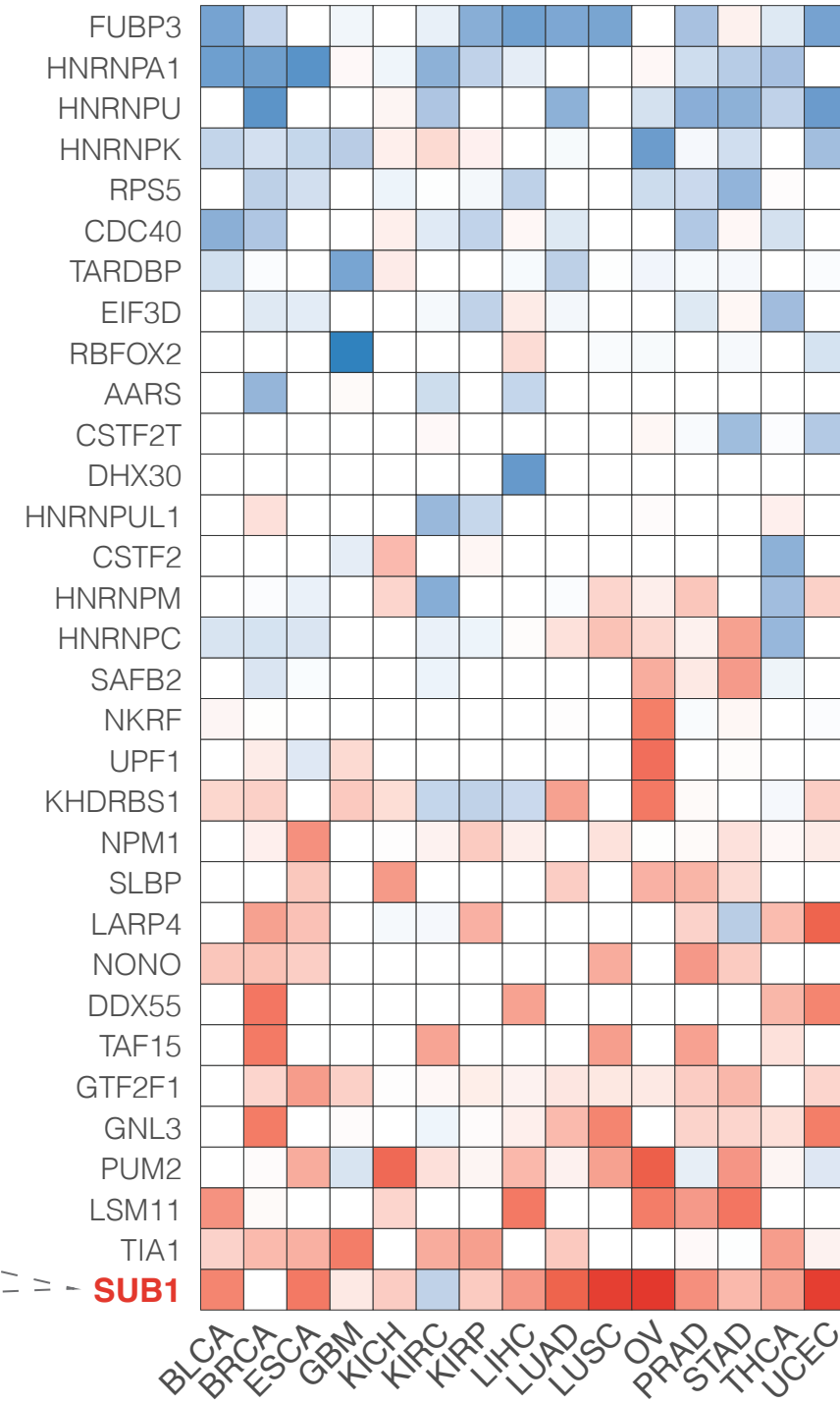








### RNA-Binding Proteins



### Transcription Factors

