

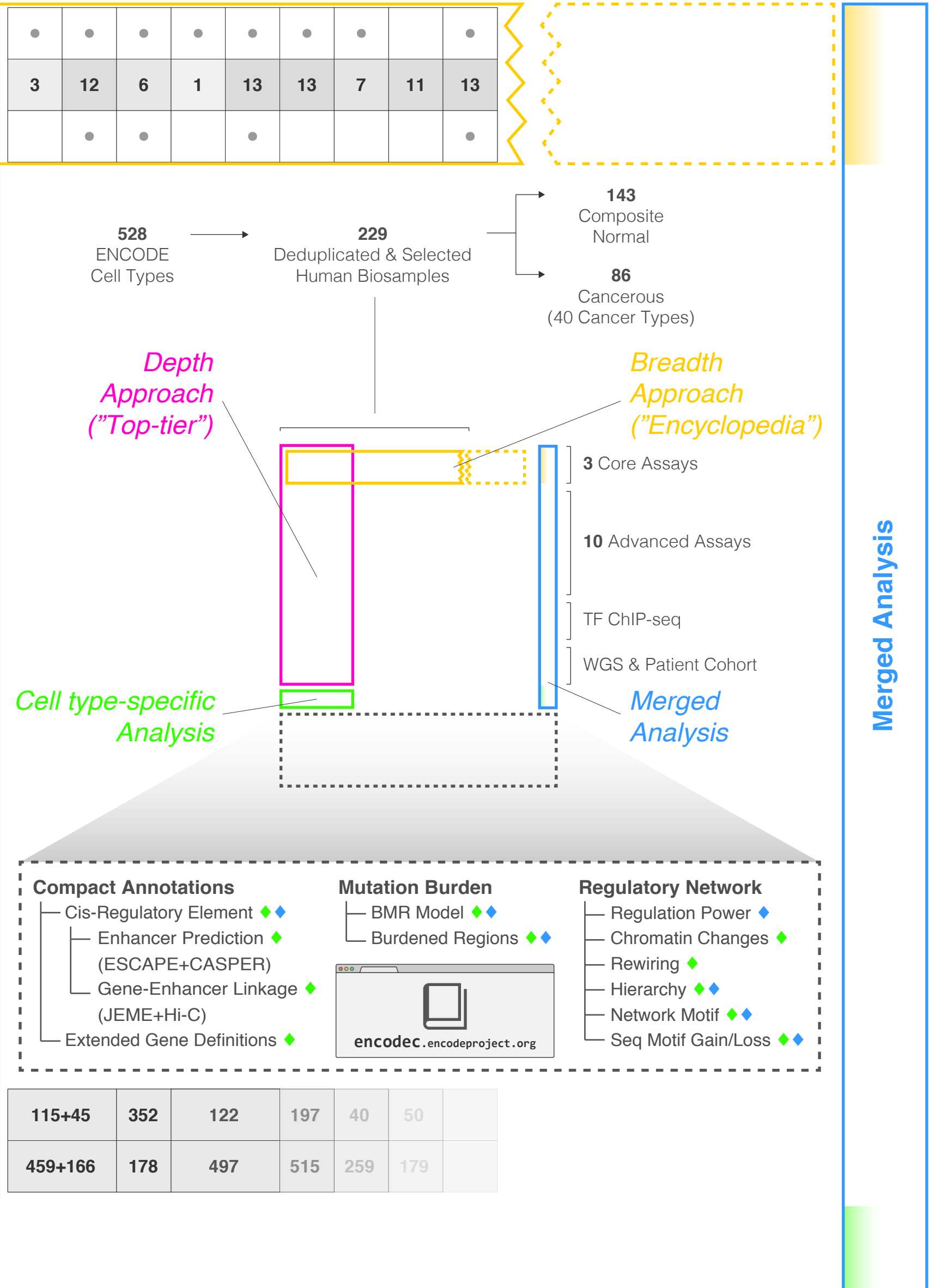
BIOSAMPLE →

ASSAY ↓

EN-CODEC

| | K562 | HepG2 | A549 | MCF-7 | HeLa-S3 | H1-hESC | Caco-2 | HCT116 | Panc1 | LNCaP | PC-3 | PC-9 | SK-N-MC | DND-41 | SK-N-SH | ... |
|--|------|-------|------|-------|---------|---------|-----------|--------|-------|-------|------|------|---------|--------|---------|-----|
| | CML | LIHC | LUAD | BRCA | Cervix | ESC | COAD+READ | PAAD | PRAD | LUAD | SARC | LAML | NB | | | |

| | | | | | | | | | | | | | | | | |
|----------------------------|--------------------------|-----|-----|-----|------|-----|----|---|----|---|---|----|----|---|----|----|
| Chromatin Accessibility DS | DNase-seq | • | • | • | • | • | • | • | • | • | • | • | • | • | • | • |
| | Histone Modifications HM | 12 | 11 | 11 | 5 | 11 | 11 | 3 | 12 | 6 | 1 | 13 | 13 | 7 | 11 | 13 |
| Transcription TX | RNA-seq | • | • | • | • | • | • | • | • | • | • | • | • | • | • | • |
| | RAMPAGE | • | | | | | | | | | | | | | | |
| RNA-binding Proteins RP | eCLIP | 89 | 70 | | | | | | | | | | | | | |
| | TF KD | 85 | 61 | | 2 | | | | | | | | | | | |
| shRNA/siRNA Knockdown KD | RBP KD | 234 | 225 | | | | | | | | | | | | | |
| | ChIA-PET | 3 | 1 | | 4 | 2 | | | | | | | | | | |
| 3D Chromatin Structure 3D | Hi-C | ▲ | | • | ▲ | | ▲ | | | | | | | | | |
| | STARR-seq | • | ▲ | | • | | | | | | | | | | | |
| Enhancers SS | WGBS | • | • | | ▲ | | • | | | | | | | | | |
| | RRBS | • | • | • | • | • | • | | | | | | | | | |
| Methylation ME | Repli-seq/chip | • | • | • | • | • | • | | | | | | | | | |
| | TF Total | 207 | 95 | 31 | 52 | 59 | 49 | | | | | | | | | |
| Transcription Factors TF | TFSS | 125 | 69 | 23 | 34 | 34 | 32 | | | | | | | | | |
| | Chromatin Remodeller | 31 | 13 | 3 | 7 | 9 | 9 | | | | | | | | | |
| | Cofactor | 20 | 7 | 3 | 4 | 6 | 3 | | | | | | | | | |
| | General (GTF) | 17 | 4 | 2 | 2 | 10 | 5 | | | | | | | | | |
| | Other | 14 | 2 | | 5 | | | | | | | | | | | |
| | | | | | | | | | | | | | | | | |
| Cell Line WGS WG | SNV | ▲ | | | ▲ | ▲ | | | | | | | | | | |
| | SV | ▲ | | | ▲ | ▲ | | | | | | | | | | |
| Patient Cohort PC | Mutation | 150 | 82 | 197 | 116 | | | | | | | | | | | |
| | Expression | 173 | 373 | 515 | 1100 | 546 | | | | | | | | | | |



Available from
 ● ENCODE
 ▲ External Resource
 ◆ EN-CODEC

Cell type-spec. Analysis

Merged Analysis

Fig 1

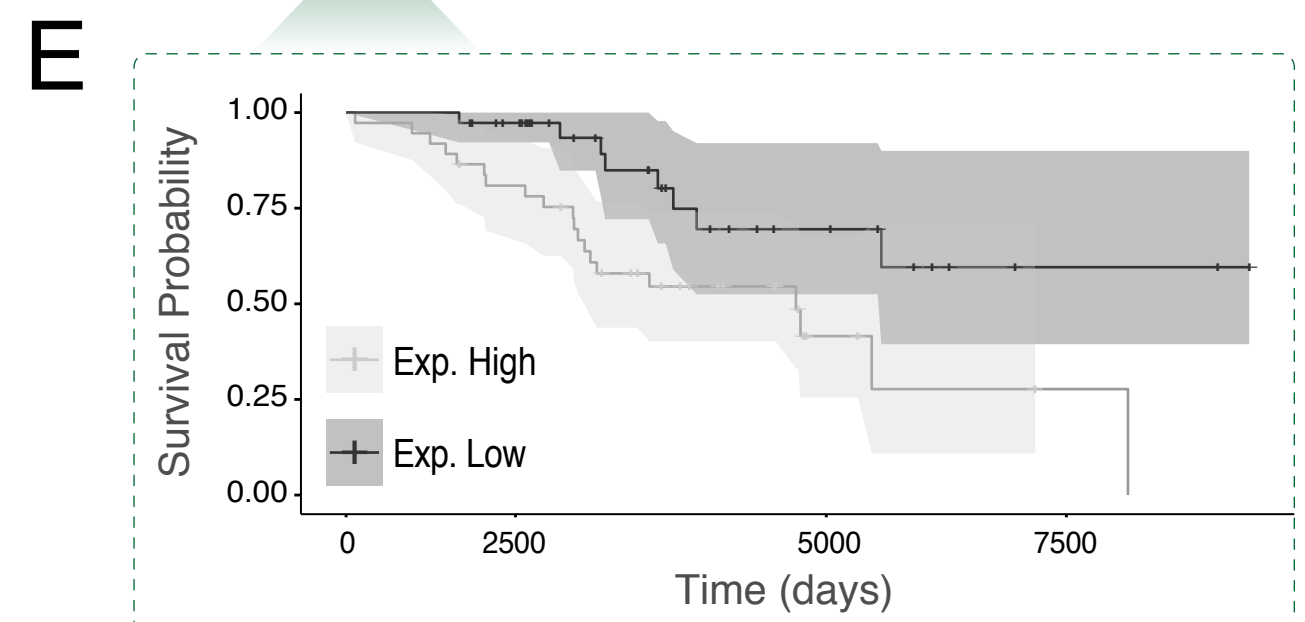
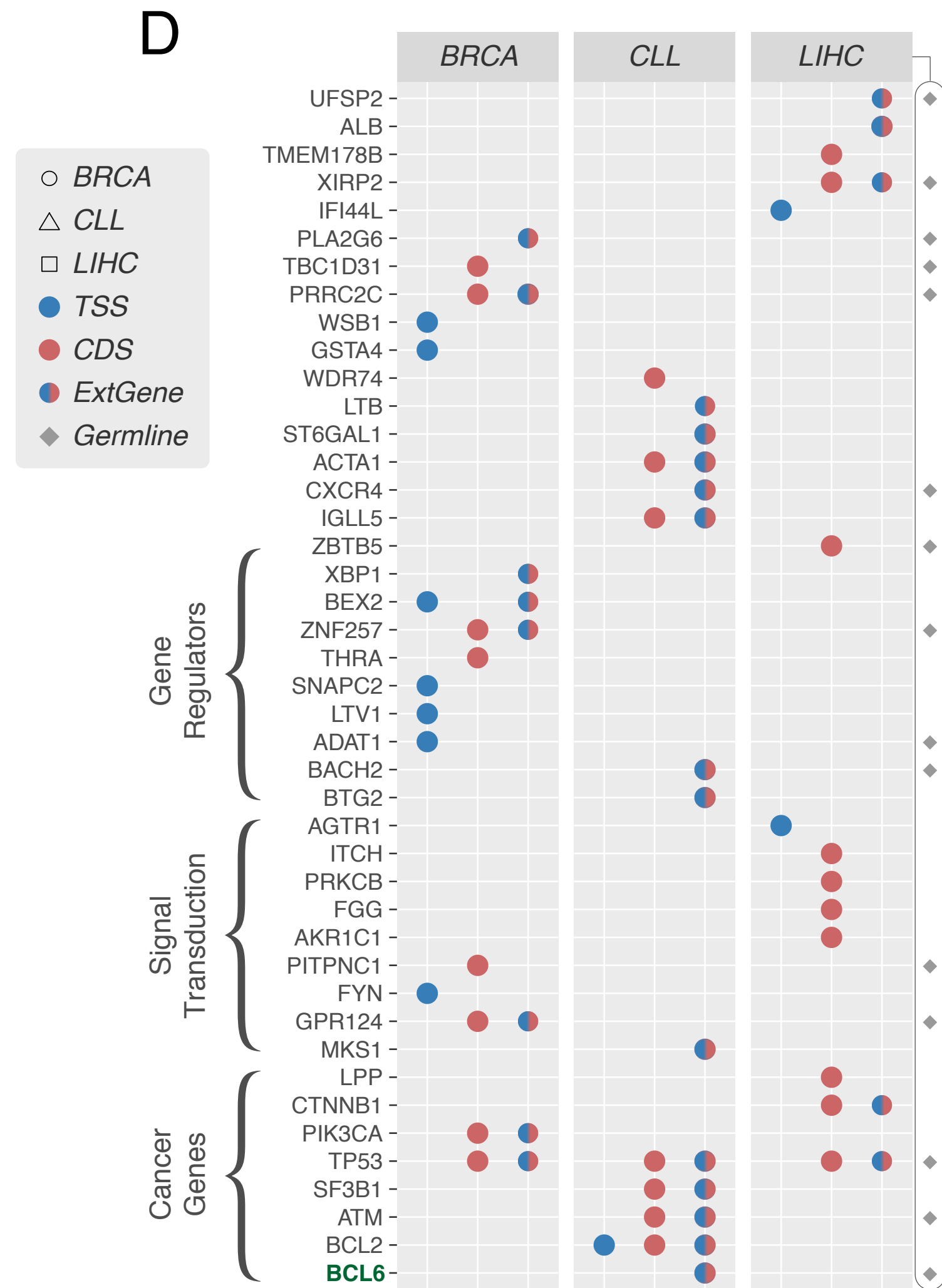
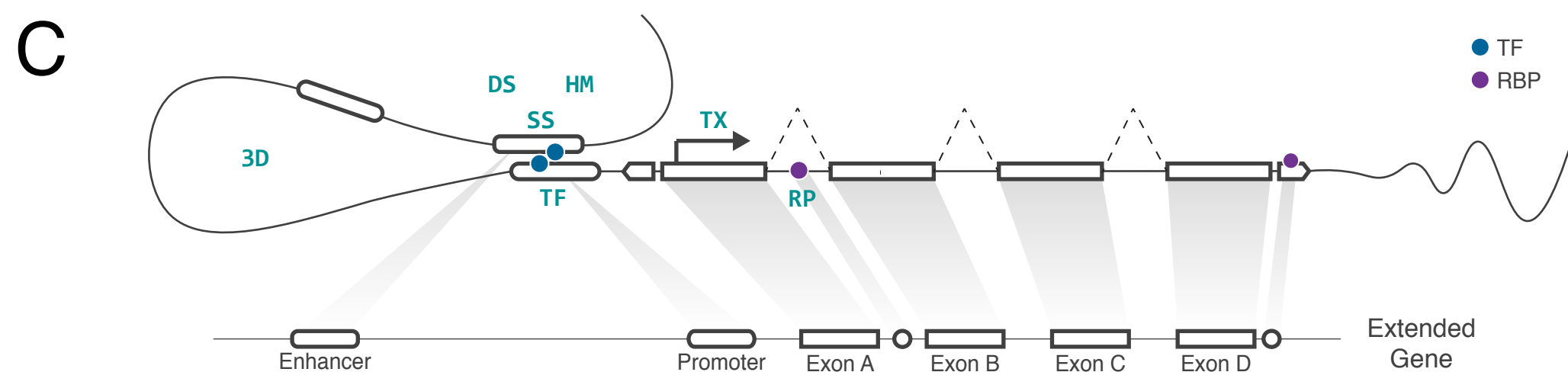
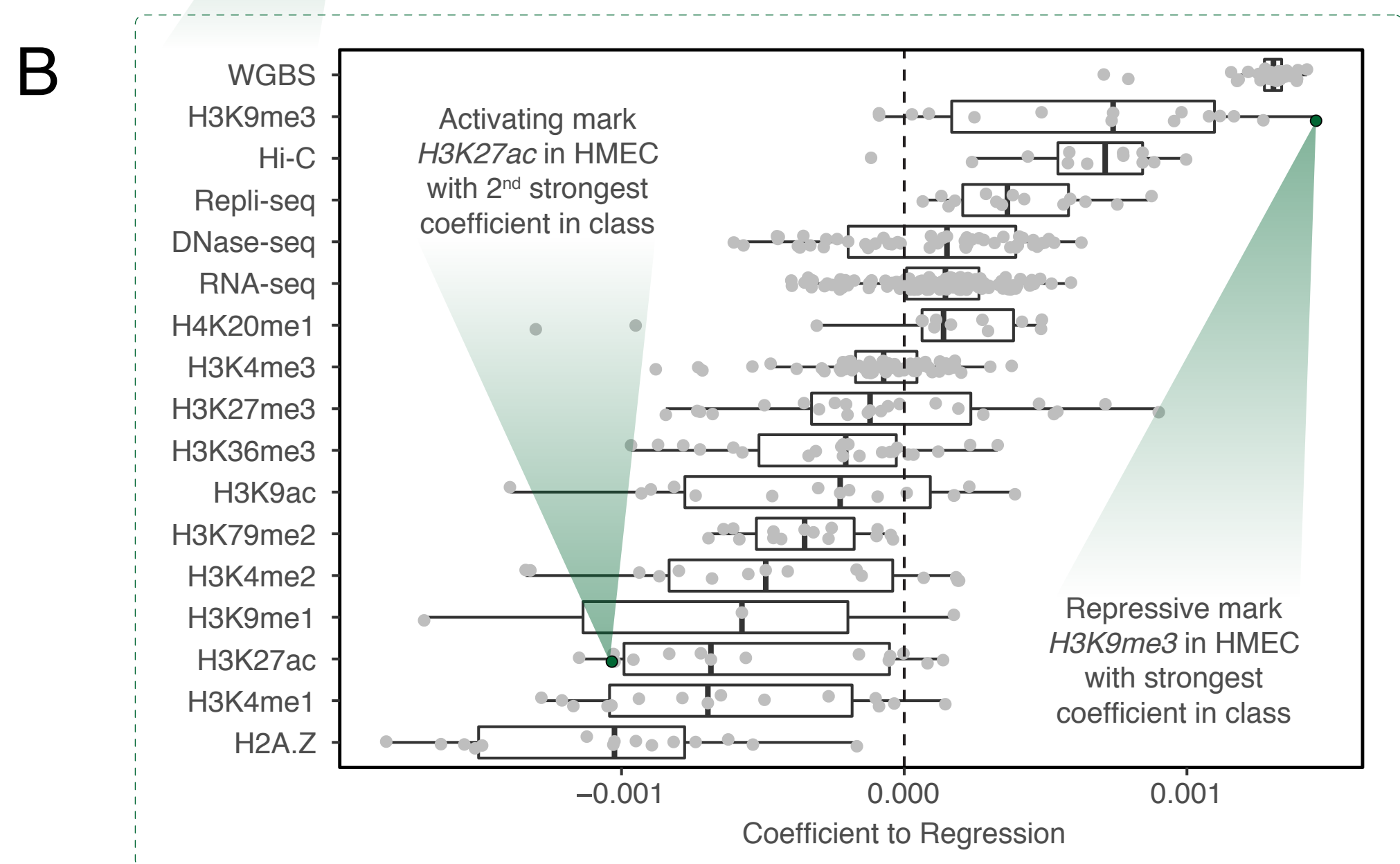
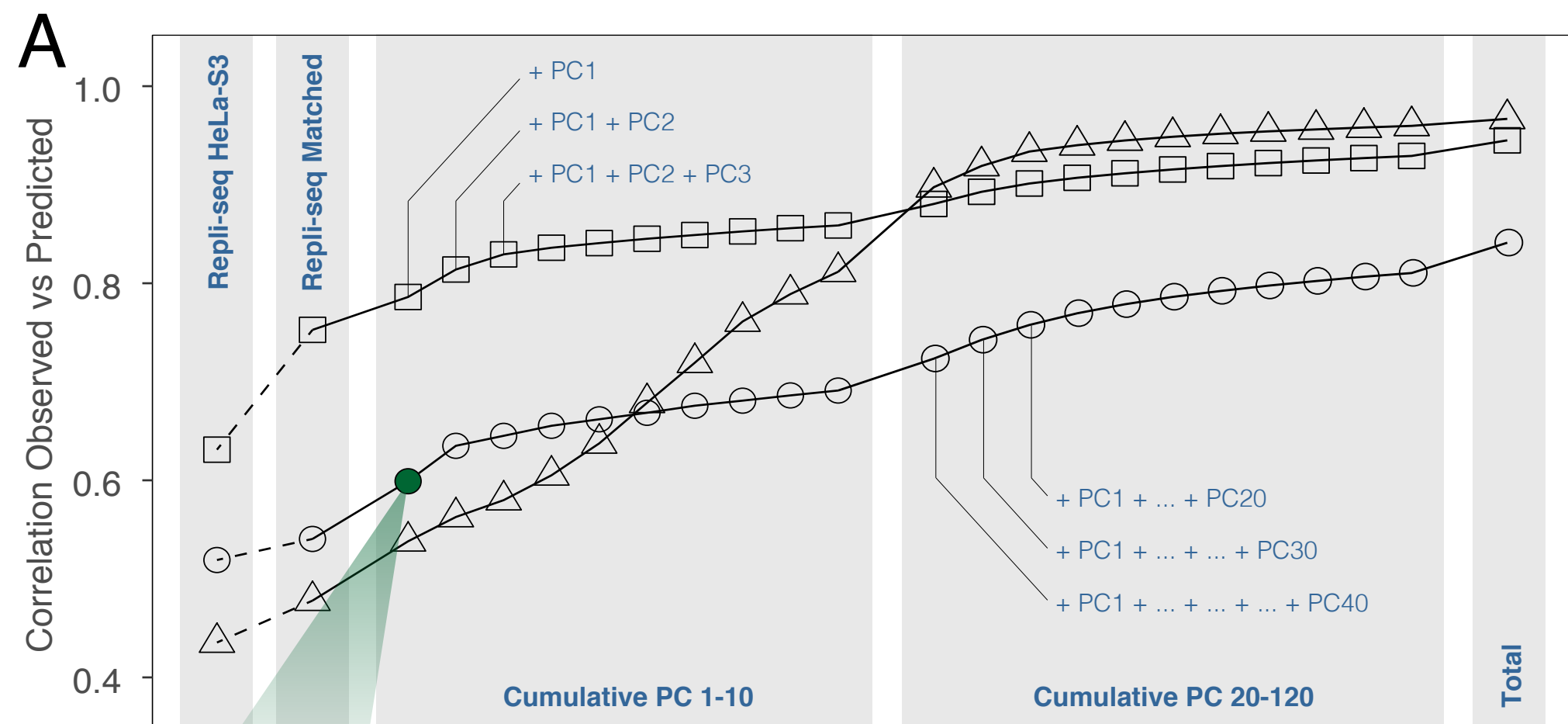


Fig 2

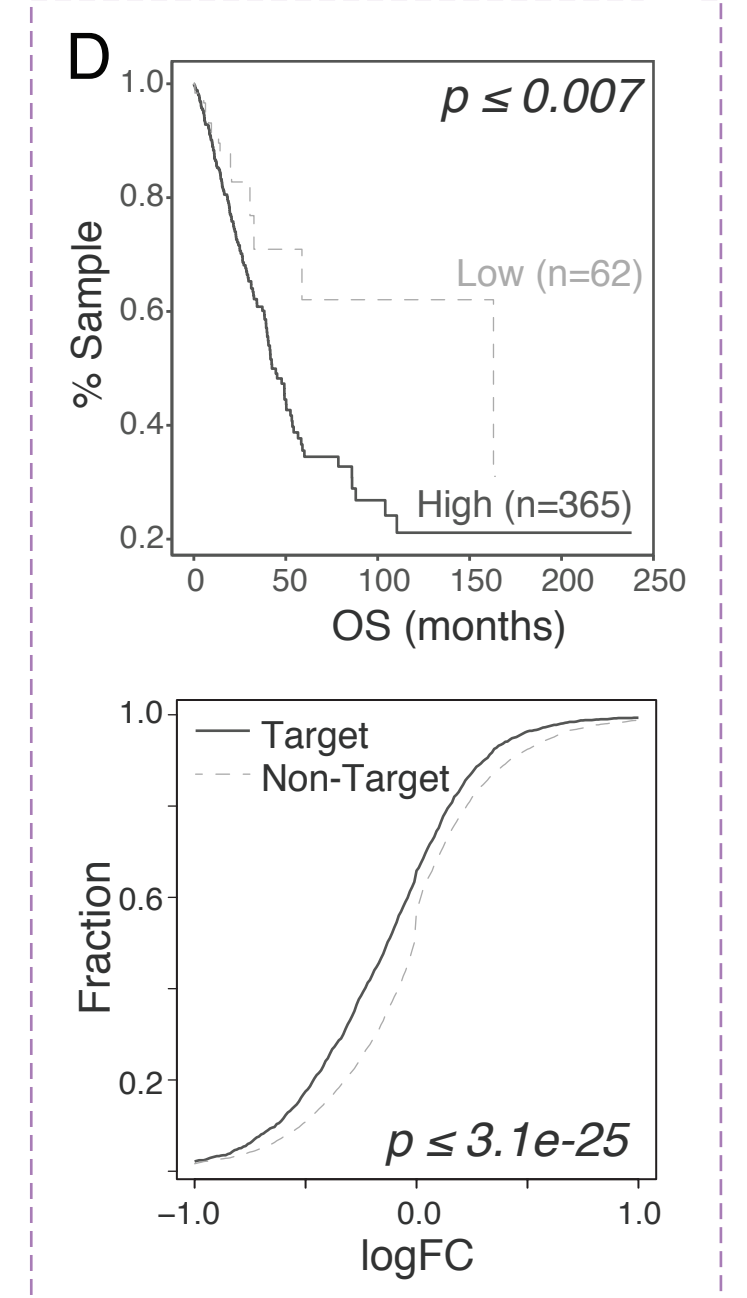
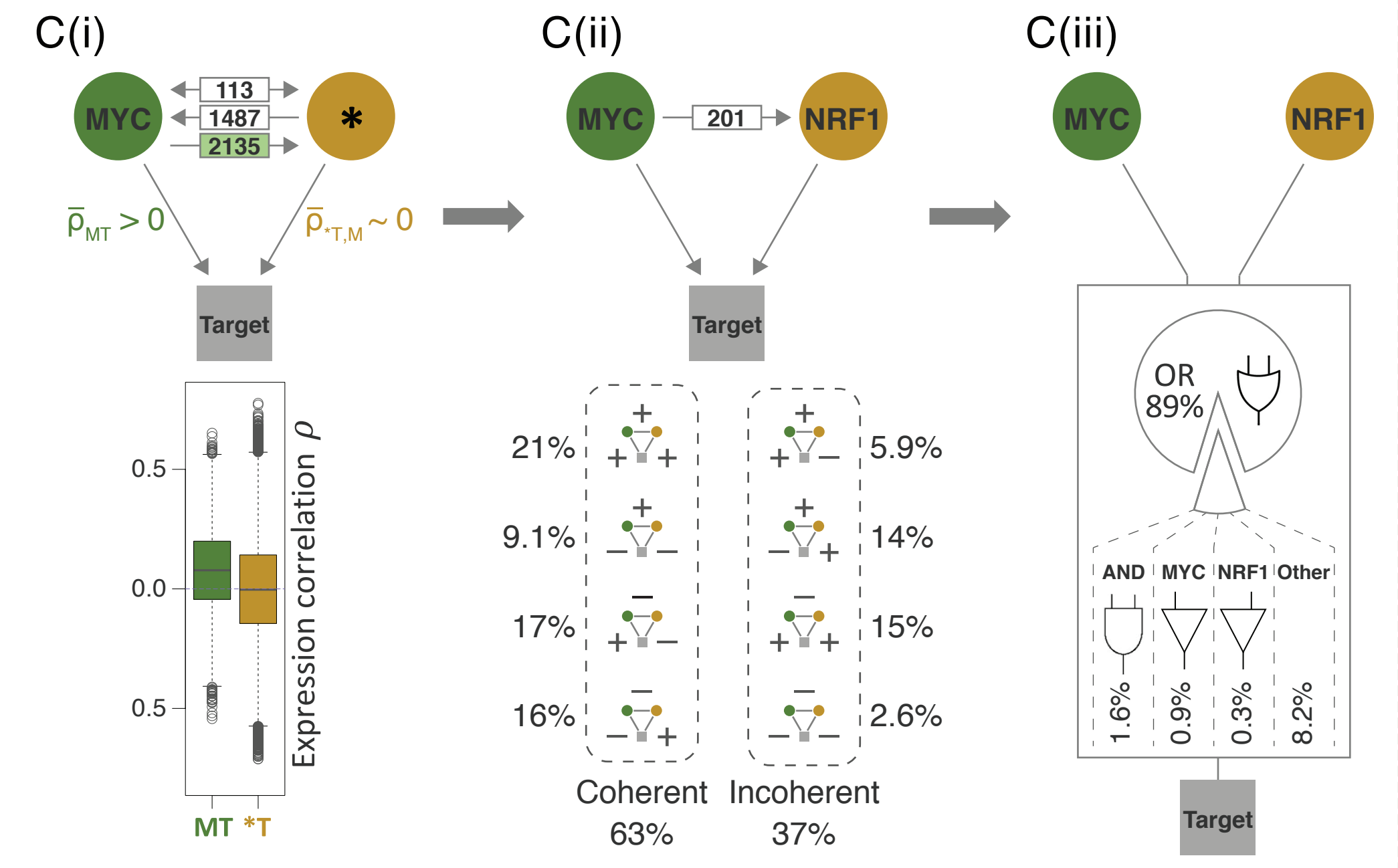
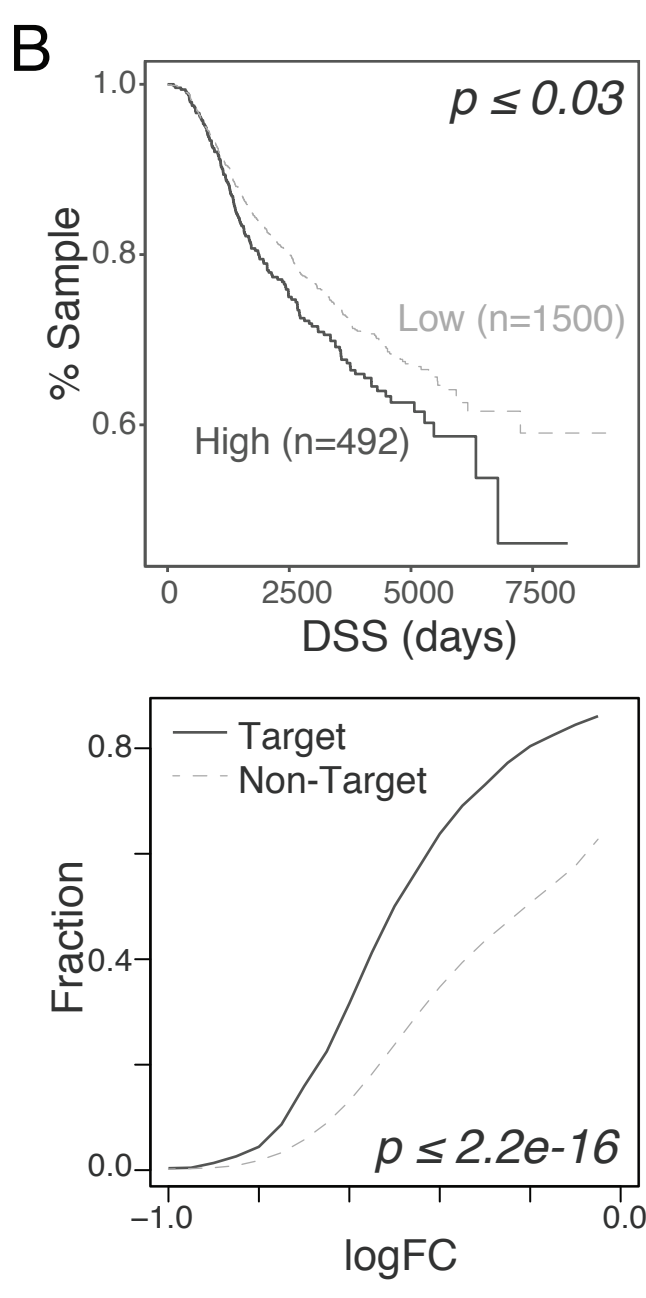
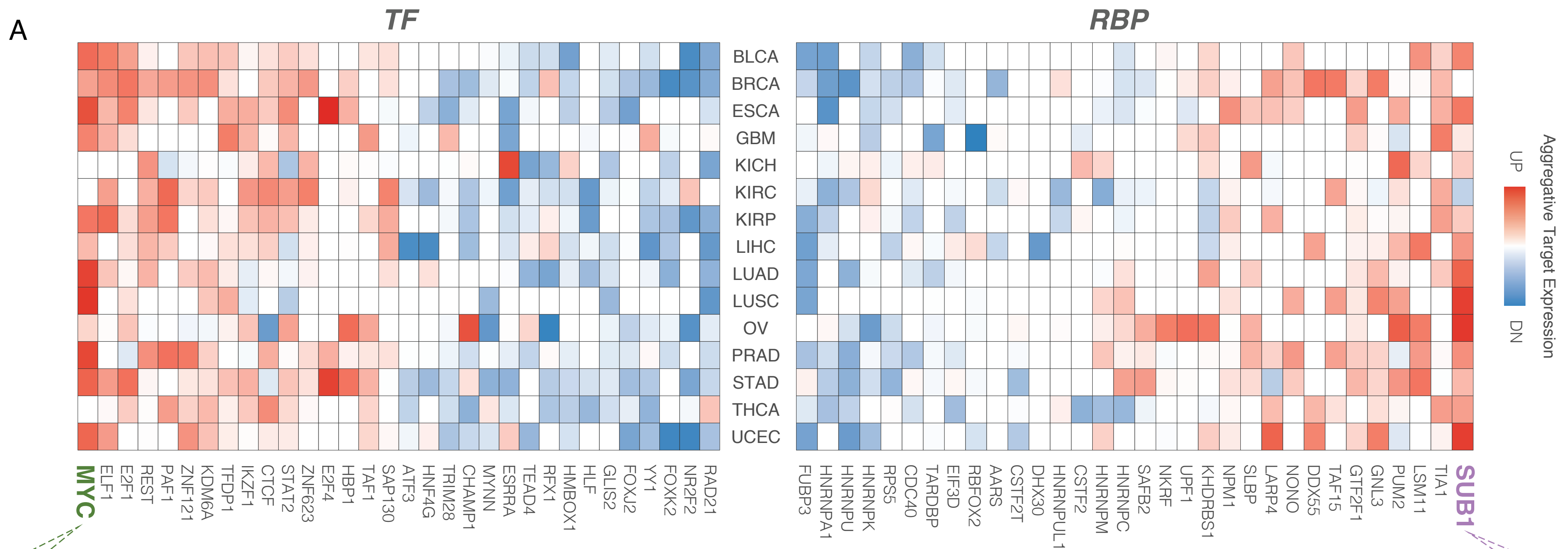
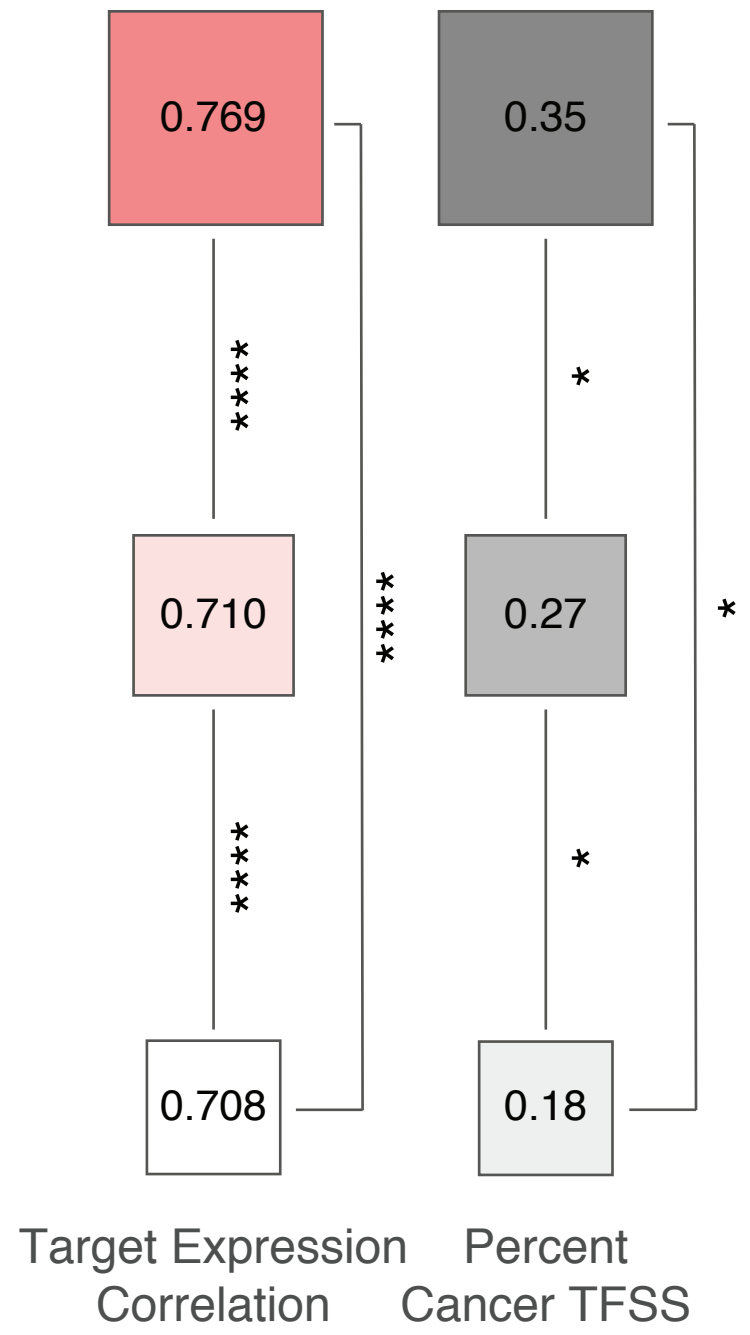
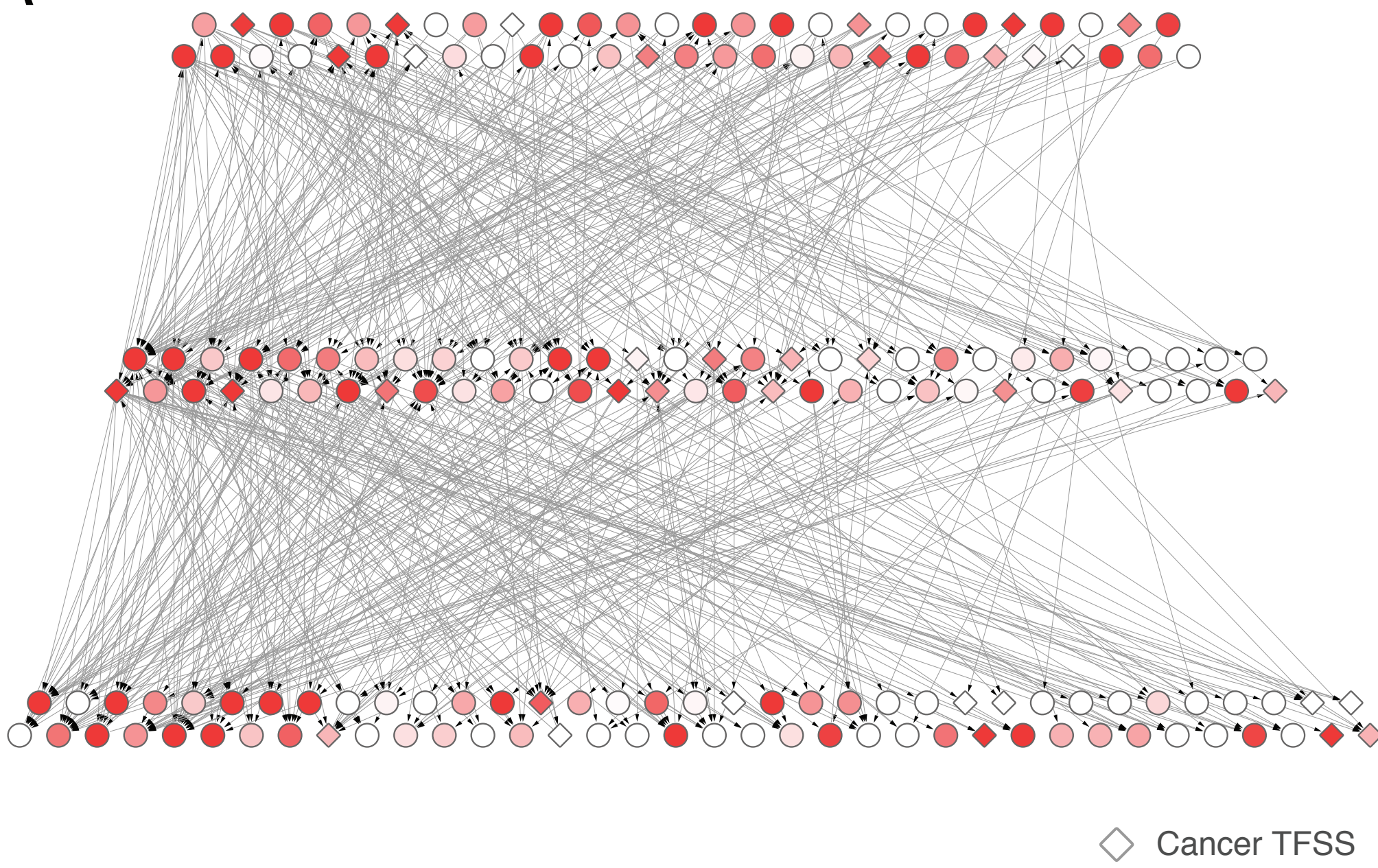


Fig 3

A



B

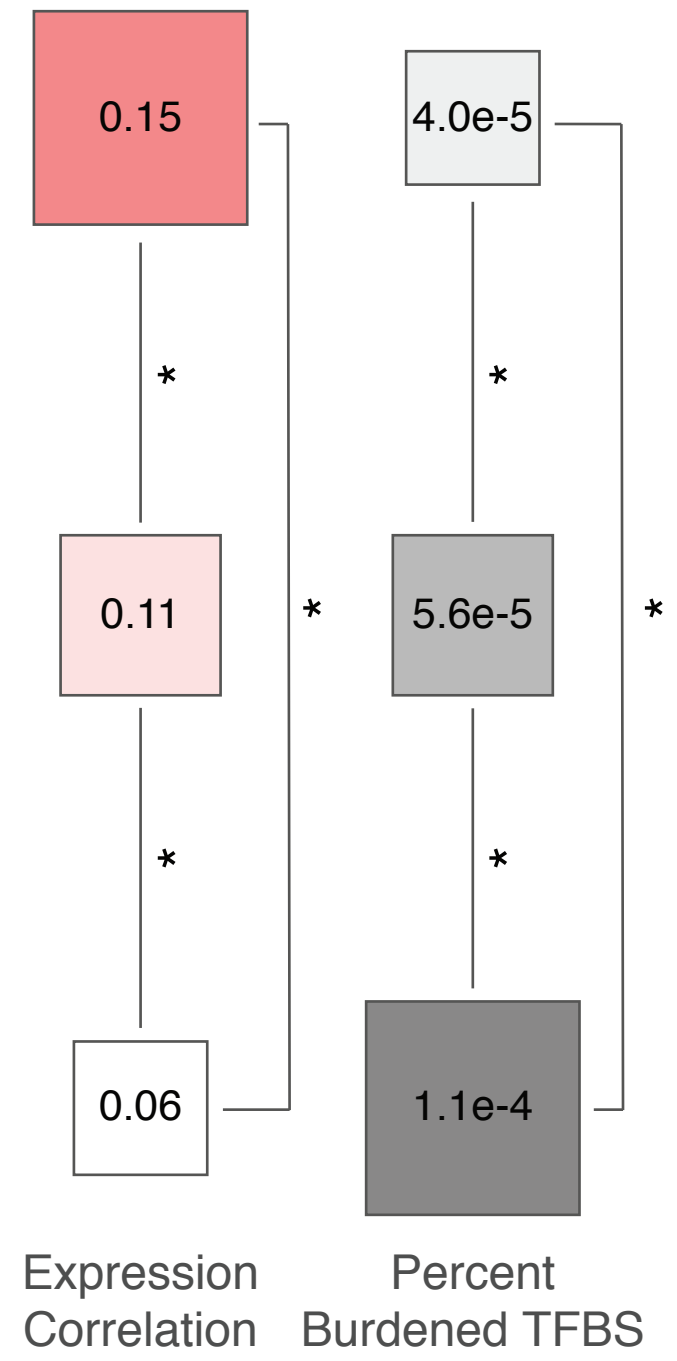
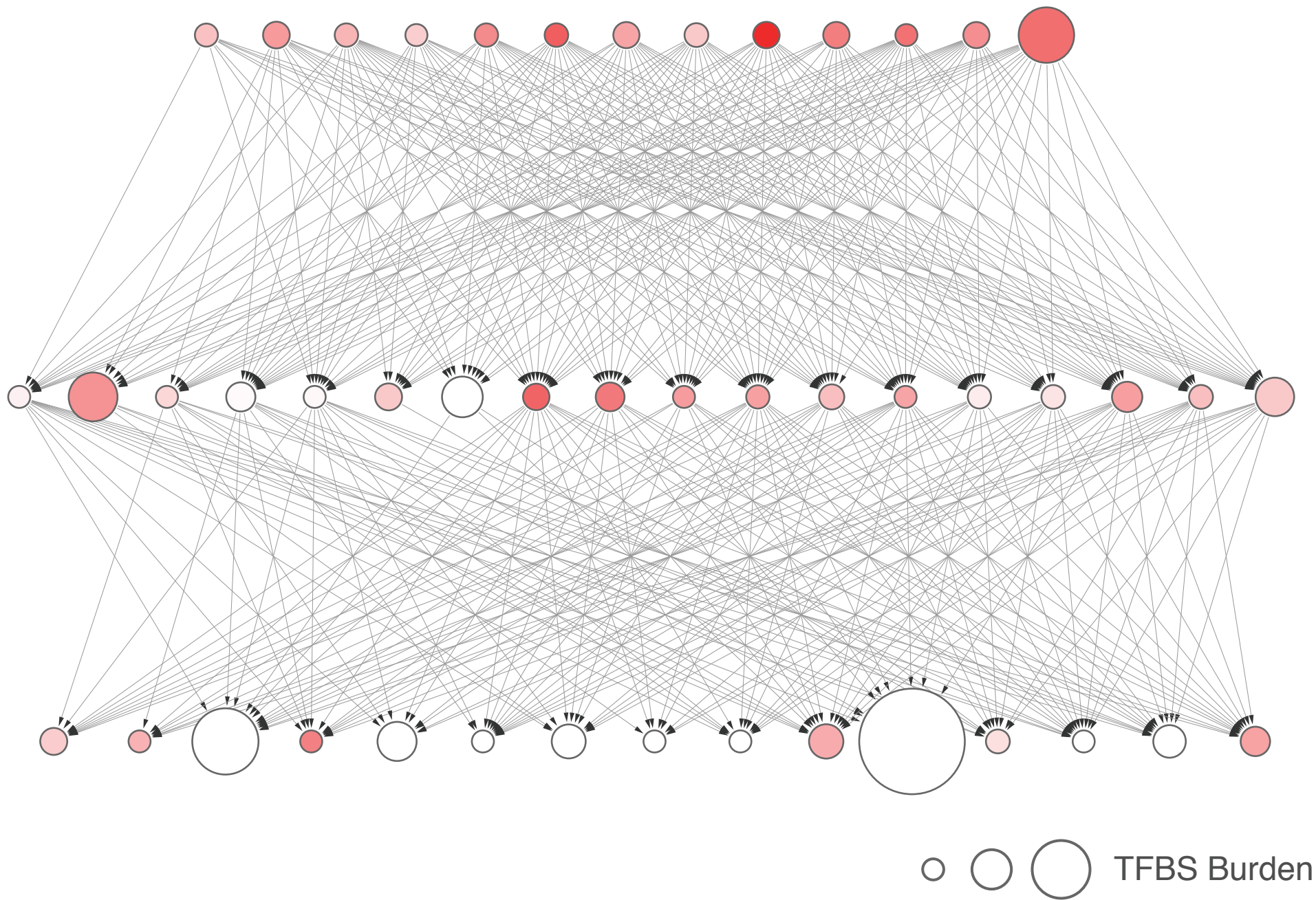


Fig 4

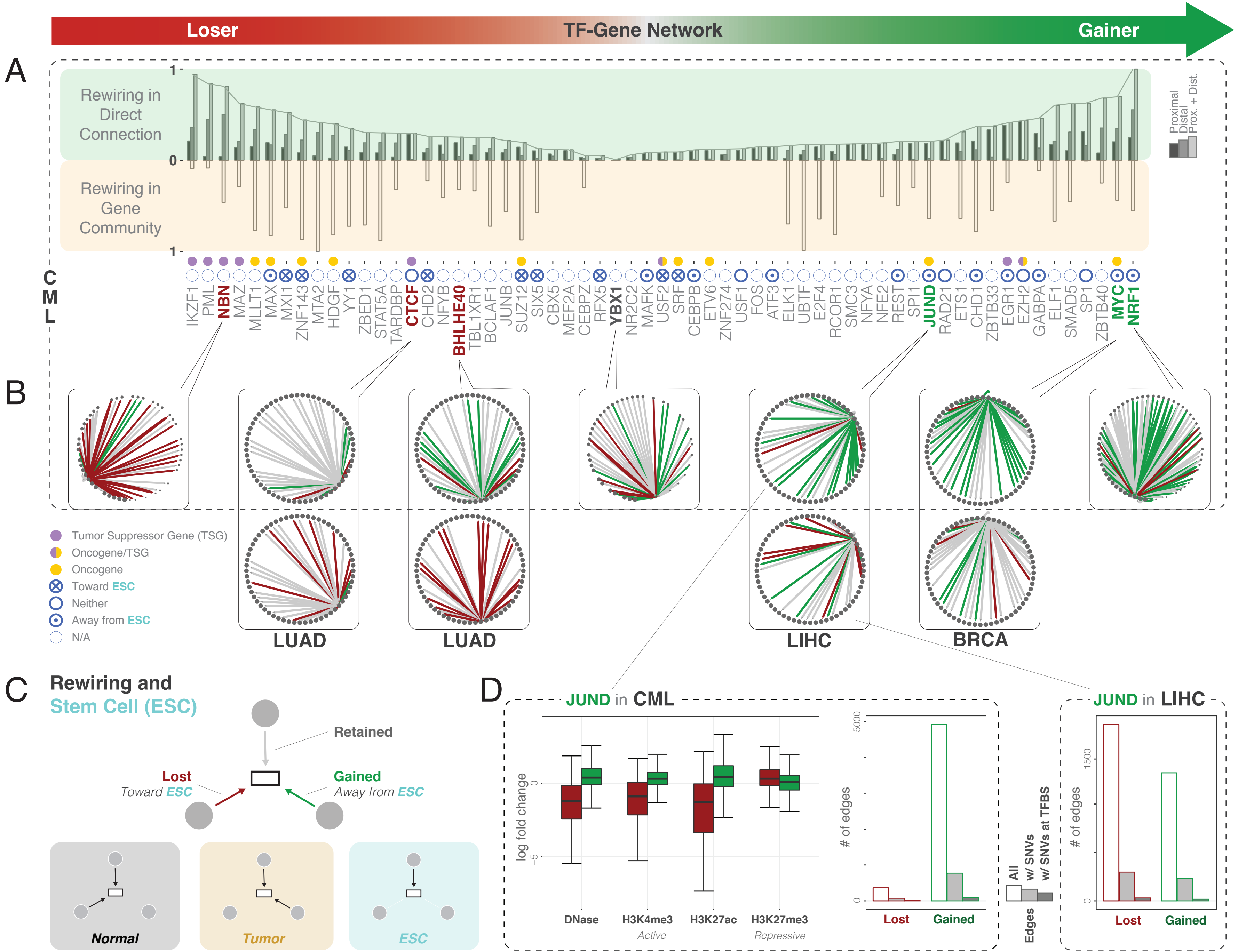


Fig 5

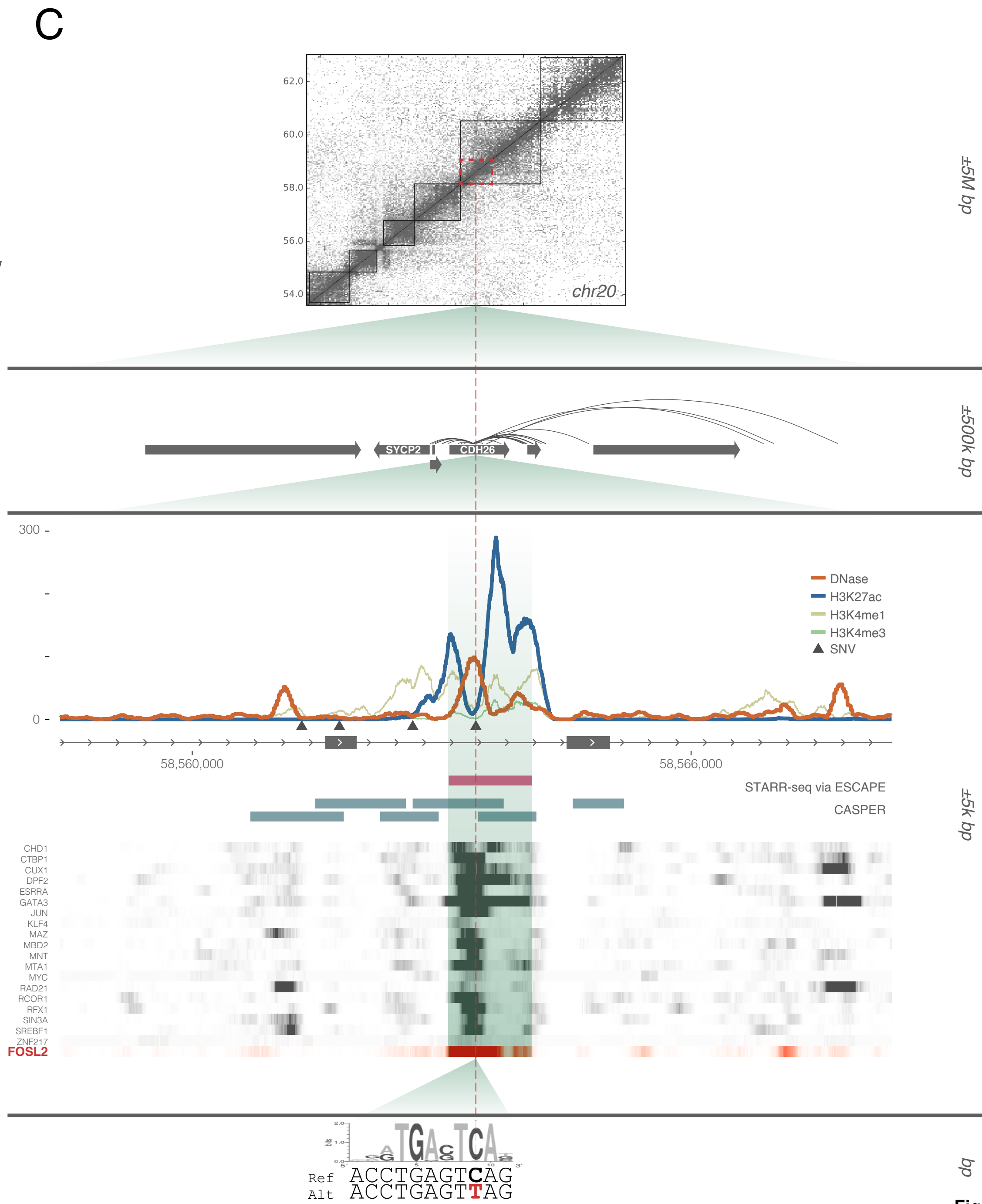
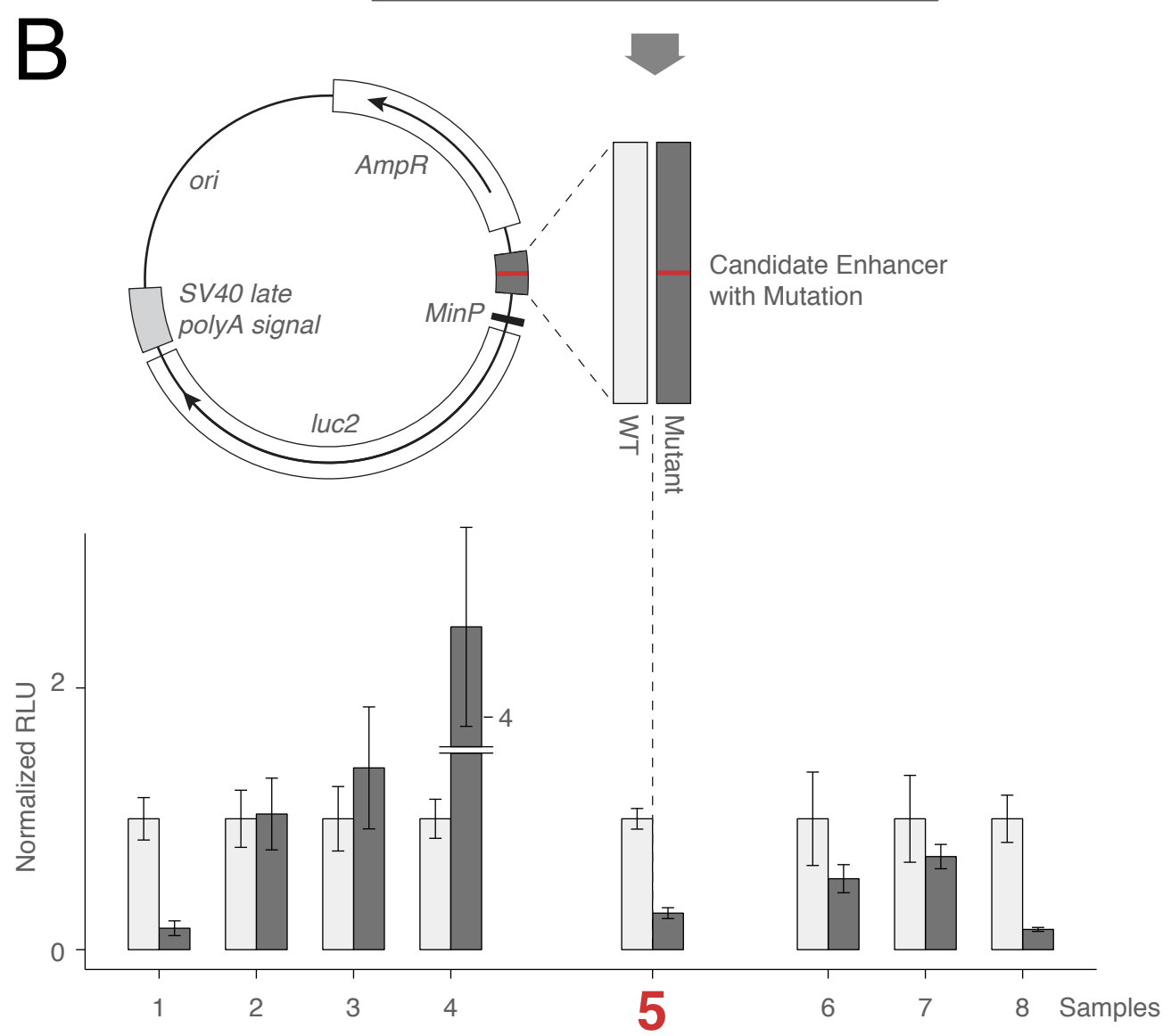
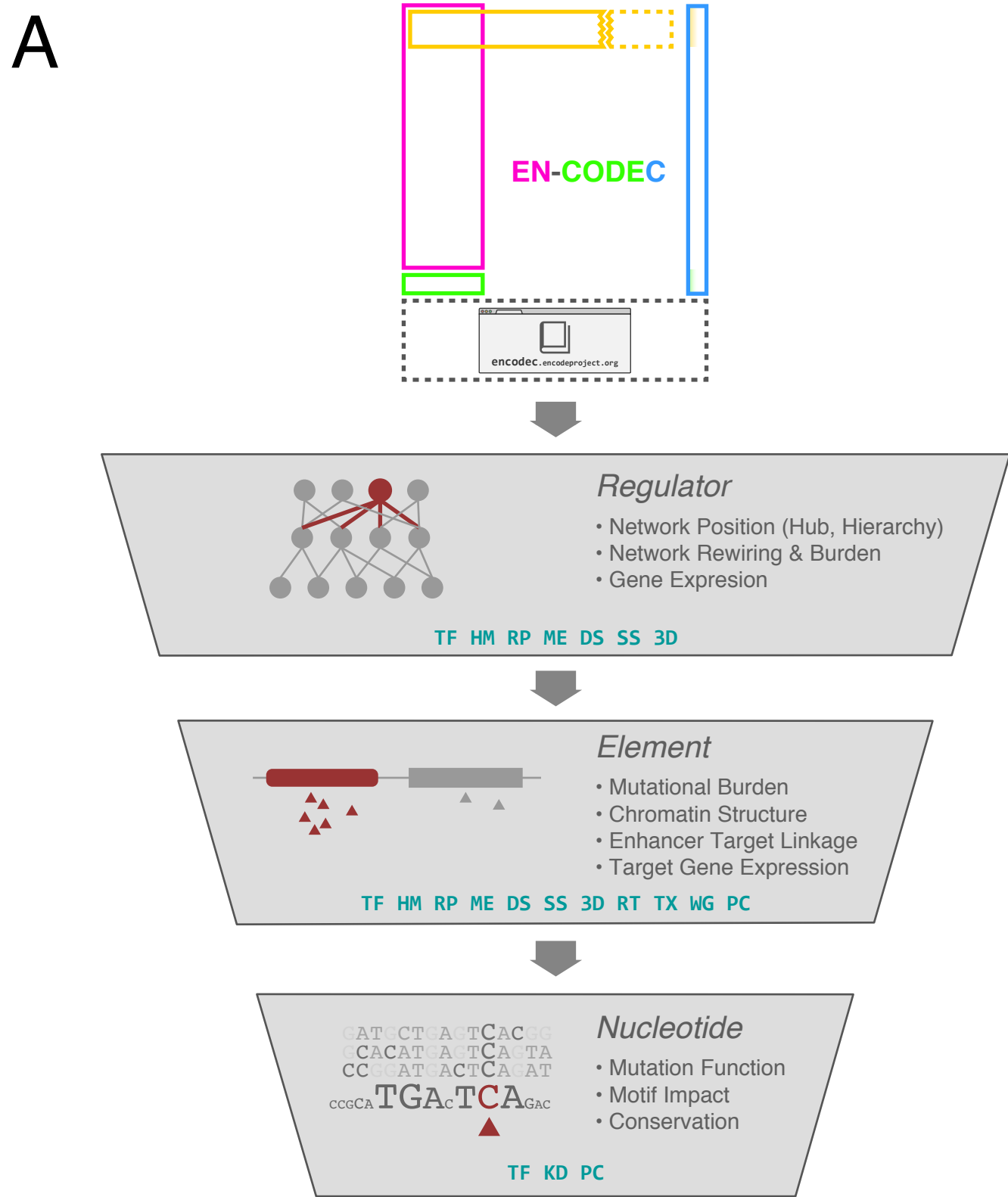


Fig 6