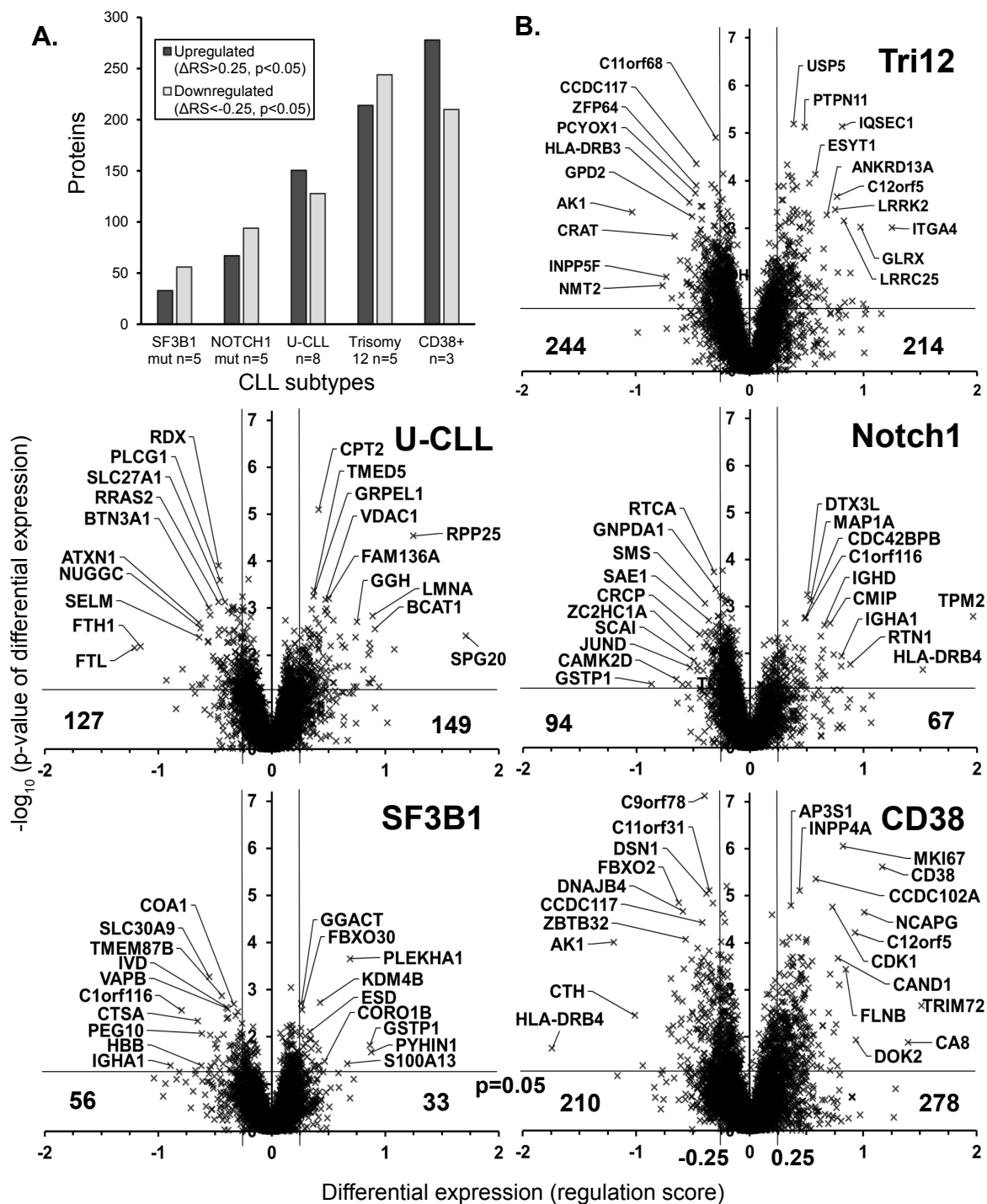
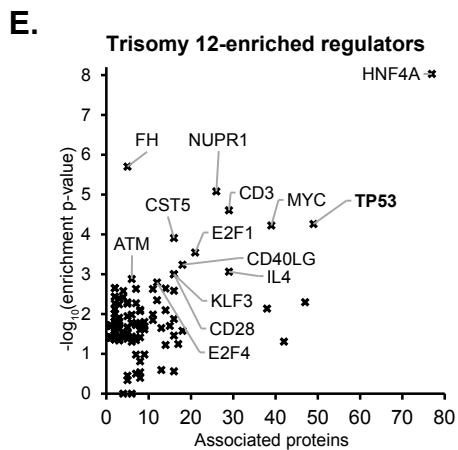
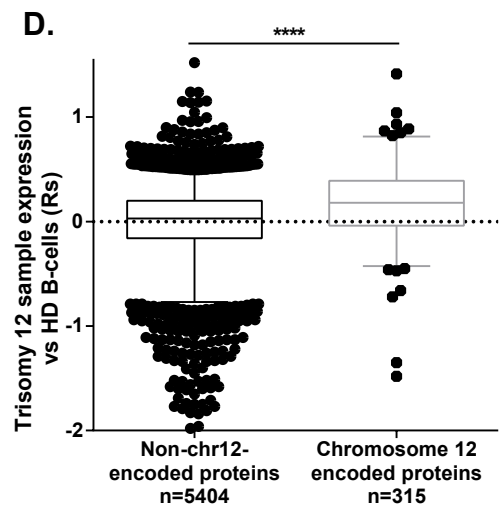
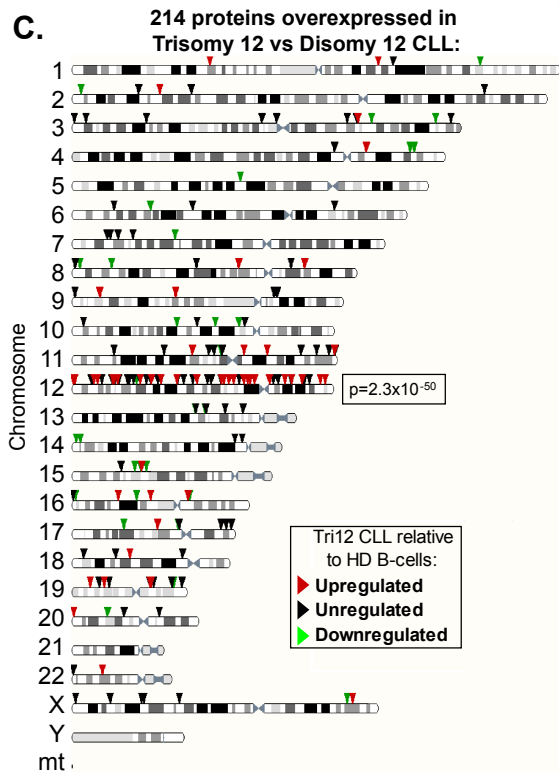


Figure S3. Analysis of emerging subtype-specific expression patterns across CLL.

A. Summary of subtype-specific protein expression determined by differential regulation score ($\Delta R_s = R_s \text{ subtype A} - R_s \text{ subtype B}$). **B.** Volcano plots highlighting examples of the most upregulated and downregulated proteins in each subtype. Statistical testing was applied as a means of highlighting an approximate signature. **C.** Chromosome enrichment analysis of the 214 trisomy 12-overexpressed proteins, additionally highlighting the expression of those proteins relative to B-cell controls. **D.** Chr12-encoded protein expression in trisomy 12 samples relative to HDB, compared to that of non-chr12 proteins. **E.** IPA upstream regulator analysis of the proteins regulated in trisomy 12 versus disomy 12 CLL. **F.** literature-derived associations between trisomy 12-upregulated chromosome 12 proteins and p53, highlighting the expression in individual samples. **G.** The location of each protein's gene on chromosome 12, of those described in F. **H.** Pathway describing those 10 proteins (from F.) suggested to interfere with effective p53 function.





F. Trisomy 12-overexpressed chr12-encoded p53-associated proteins

Protein	Evidence (Unique peptides, PSMs)	Log ₂ (ratios to HDB) (Disomy 12 vs Trisomy 12)	Tri12 ΔRs p-value	Effects on p53
Ubiquitin carboxyl-terminal hydrolase 5 (USP5)	26, 312		0.39 6.4E-6	Negatively regulates p53
Tyr phosphatase non-receptor type 11 (SHP2/PTPN11)	26, 197		0.48 7.4E-6	Signalling inhibits p53-induced senescence
Ubiquitin carboxyl-terminal hydrolase 15 (USP15)	26, 199		0.33 4.6E-5	Stabilises MDM2 expression
TP53-regulated inhibitor of apoptosis 1 (TRIA1/p53CSV)	3, 31		0.53 1.1E-4	Repressor of p21
Arf-GAP with GTPase, ANK/PH 2 (AGAP2/PIKE-A)	8, 51		0.42 1.3E-4	Knock-down stabilised p53, acts via AKT/MDM2
Proliferation-associated protein 2G4 (PA2G4/EBP1)	16, 421		0.25 1.3E-4	Enhances MDM2-p53 association
Thioredoxin reductase 1 (TXNRD1)	18, 141		0.38 1.5E-4	Inhibition increased p53-DNA binding
TP53-induced glycolysis + apoptosis regulator (TIGAR)	6, 45		0.77 2.1E-4	Inversely correlates with p53 expression
RAS oncogene family member Rab-35 (RAB35/1c)	7, 216		0.44 9.3E-4	Suppresses PRPK-induced p53 activation
Tyr phosphatase non-receptor type 6 (SHP1/PTPN6)	41, 1521		0.46 7.3E-3	Overexpression reduced p53 phosphorylation

G. Chromosome 12:



H. Tri12/Chr12 inhibitors of p53 activity

