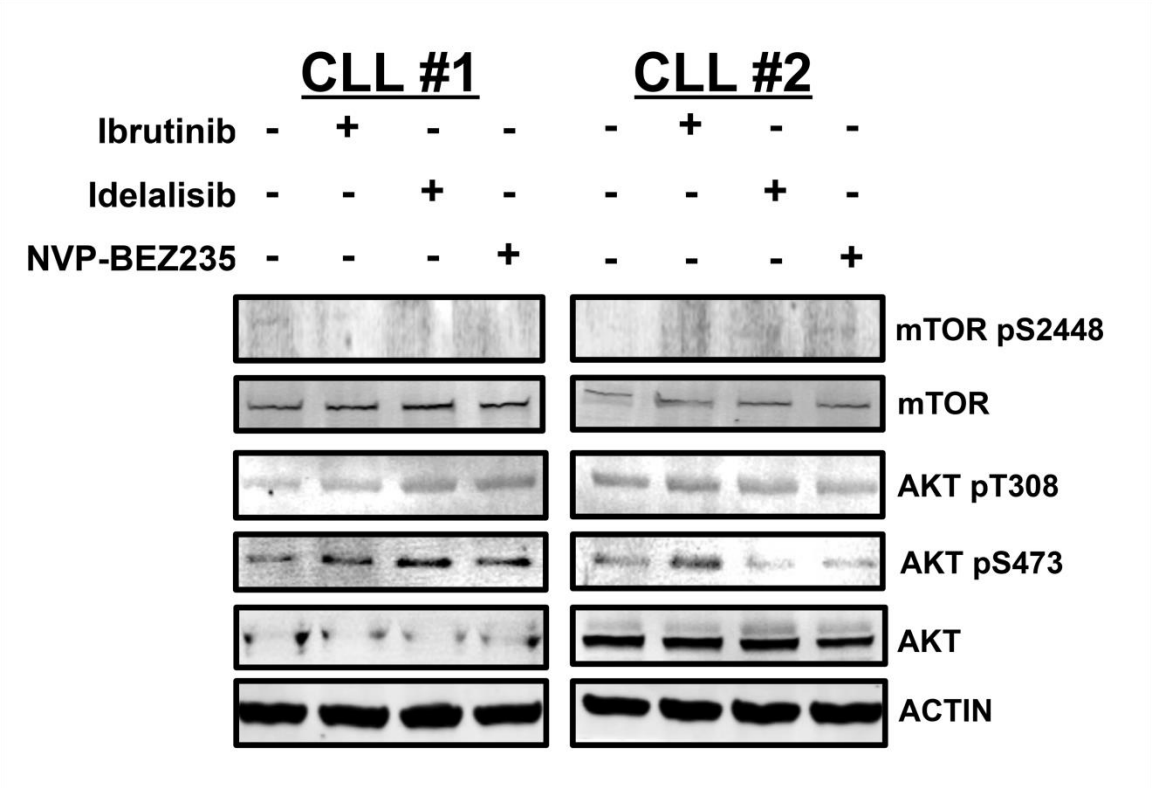
RPPA-based protein profiling reveals eIF4G overexpression and 4E-BP1 serine 65 phosphorylation as molecular events that correspond with a pro-survival phenotype in chronic lymphocytic leukemia

Supplementary Material

Supplementary File S1: Normalized RPPA intensities of each analyzed antibody probe from CLL and normal B-cell samples.

Supplementary File S2: Ingenuity pathway analysis of proteins significantly altered between CLL and normal B-cells.



Supplementary Figure S1: Comparison of AKT and mTOR phosphorylation in inhibitor treated CLL cells co-cultured with HS-5 stromal cells.