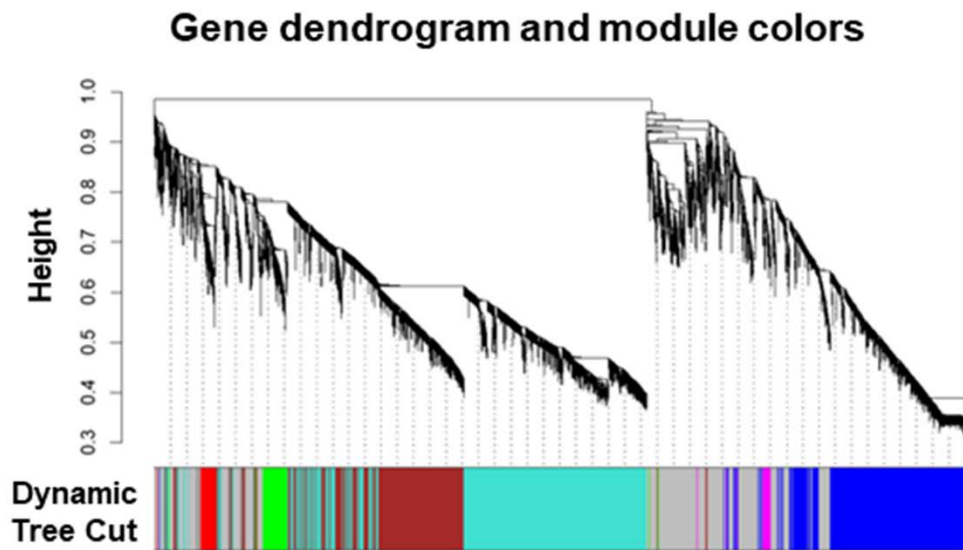**Figure S1: WGCNA-identified modules in SpiD3-treated OSU-CLL cells**

Figure S1. WGCNA-identified modules in SpiD3-treated OSU-CLL cells.
OSU-CLL cells treated with vehicle or SpiD3 (1, 2 µM) for 4 h were subject to RNA-sequencing analysis and weighted gene co-expression network analysis (WGCNA). Gene dendrogram and colors of WGCNA-identified modules for the top 500 differentially expressed genes ($P = 0.05$) in SpiD3-treated cells are shown.