

Supplementary File 2:

Most intronic MAPs from microproteins modulated by TAK-981 are lowly expressed in healthy samples. Heatmap showing the mean RNA expression [$\log(\text{RPHM} + 1)$] of all MAPs from intron-derived microprotein coding regions across normal tissues (GTEx), AML patient samples (Leucegene and TCGA cohorts), pooled medullary thymic epithelial cells (mTECs), and AML cell lines treated with DMSO or TAK-981. Black frames highlight regions with mean expression >8.55 RPHM.

