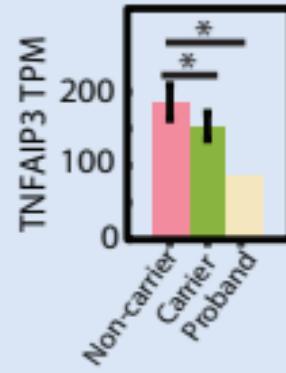
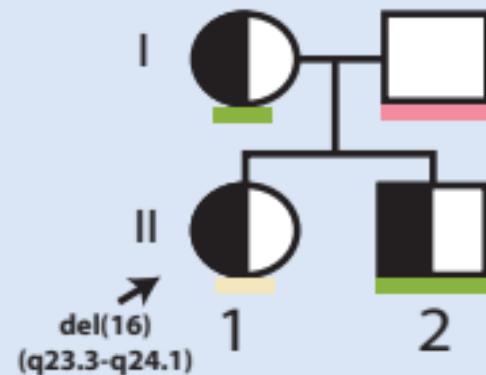


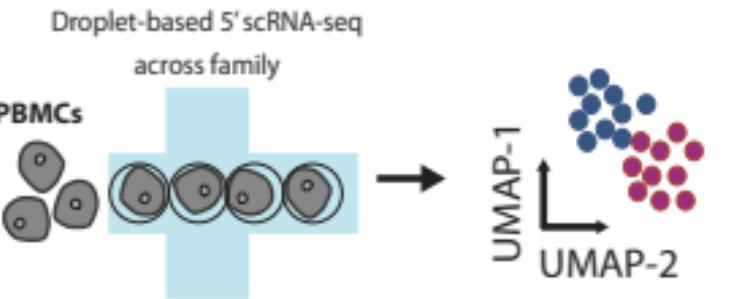
Evan's Syndrome Driven by a Familial Novel TNFAIP3 Splice Variant and De-Novo MBTPS1 Deletion Genotype

c. -15.-2A A>G, heterozygous



Reduced A20 expression in splice variant carriers & more significant in proband

Single-Cell RNA-seq Reveals Skewed Cytotoxic Immune Skewing and Activation in Proband Phenotype



Enrichment of cytotoxic CD8+ T cells, Natural Killer cells, and non-classical monocytes

Flow Cytometry and scRNA-seq suggests lineage specific NF-κB Dysegregation Mechanism

TFNAIP3 (A20) ↓
Impaired NF-κB negative feedback

MBTPS1 (S1P) ↓
Reduced ATF6 UPR buffering
Altered SREBP processing

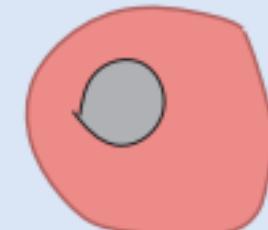
Monocyte

NF-κB ↑
UPR ↑
Inflammation ↑



CD8 T-cell

NF-κB ↓
Exhaustion ↑
Cytotoxic IFN ↑



Chronic Inflammatory Loop = *Evan's Syndrome (AIHA and ITP)*