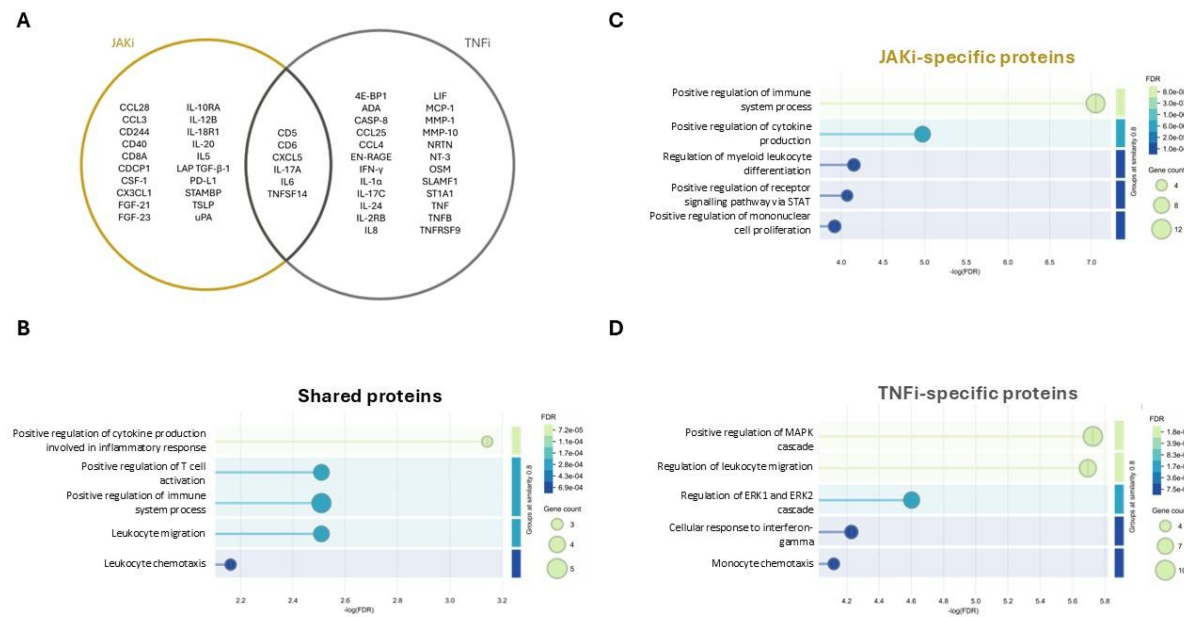




Supplementary Figure S3. Comparative analysis of *ex vivo* JAKi- and TNFi-modulated protein signatures reveals treatment-specific biological pathways.- (A) Venn diagram showing the overlap between proteins differentially secreted *ex vivo* by immune cells after JAKi or TNFi treatment. Shared proteins are shown in the overlapping area, while treatment-specific proteins appear in the corresponding sections. (B-D) Functional enrichment analysis based on Gene Ontology biological process terms for shared proteins (B), JAKi-specific proteins (C), and TNFi-specific proteins (D). Dot size represents the number of proteins associated with each term, and color indicates the level of statistical significance expressed as false discovery rate (FDR).