**S4 Fig. B cell subset and cluster analysis.** (A-C) Healthy and convalescent donor cell frequencies for (A) CD11c+, (B) T-bet+, and (C) DN subsets among viable CD19+ lymphocytes. Bars represent mean +/- SD. Statistical analysis between each donor subgroup was done with non-parametric Kruskal-Wallis test with Dunn's correction for multiple comparisons. Adjusted \( p \) value was used to determine family-wise significance at alpha = 0.05. Healthy control and total convalescent groups also compared by Mann-Whitney test with two-tailed \( p \) value, alpha = 0.05. Healthy control \( n = 24 \), conv. total \( n = 40 \), asympt. \( n = 5 \), conv. early \( n = 12 \), conv. mid \( n = 6 \), conv. late \( n = 17 \), except for T-bet (B), where \( n = 4, 9, 6, \) and 16, respectively, due to technical limitations. (D) tSNE representation of flow cytometry data sub-gated to highlight key clusters of viable CD19+ lymphocytes from a representative convalescent donor with an enriched memory B cell subset. Heatmap overlay shows median expression for the target below each plot.