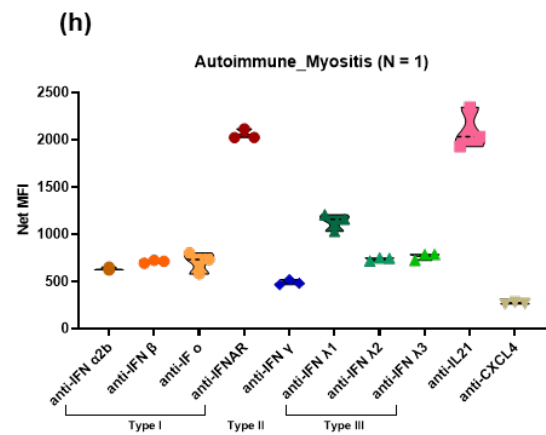
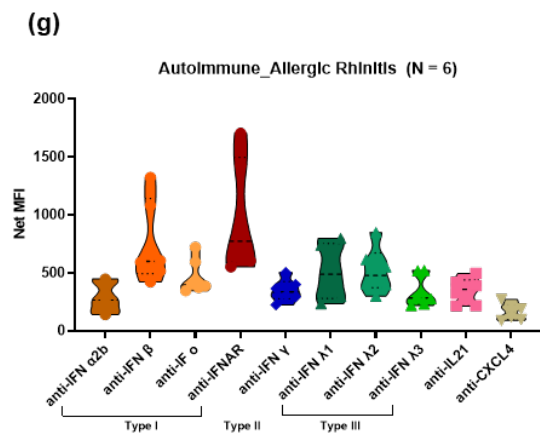
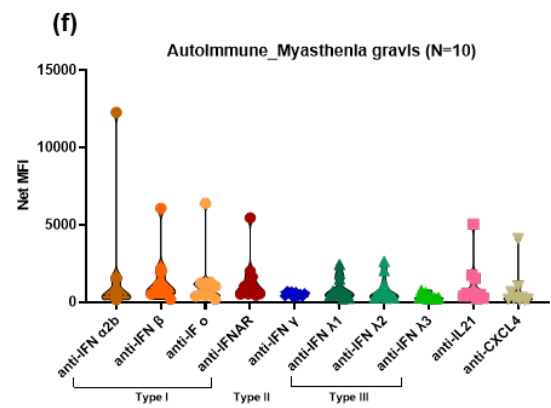
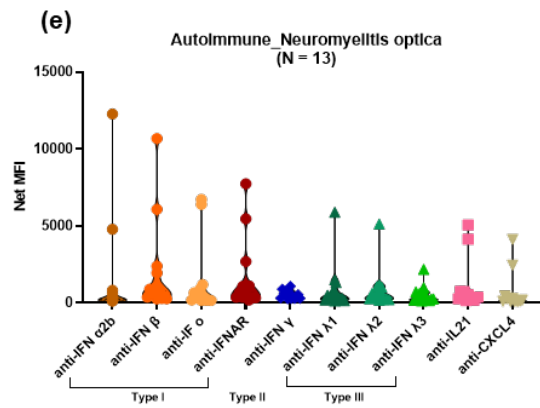
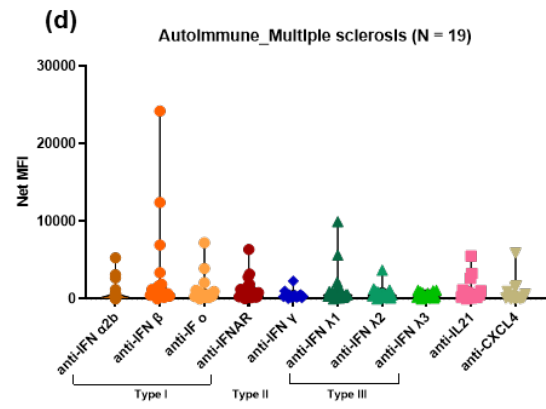
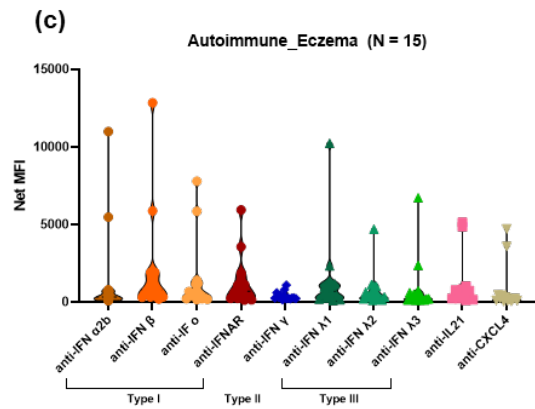
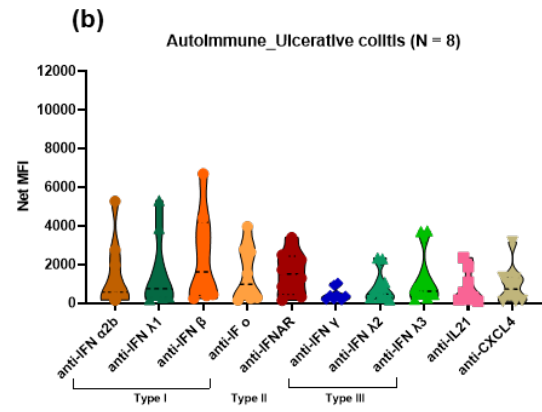
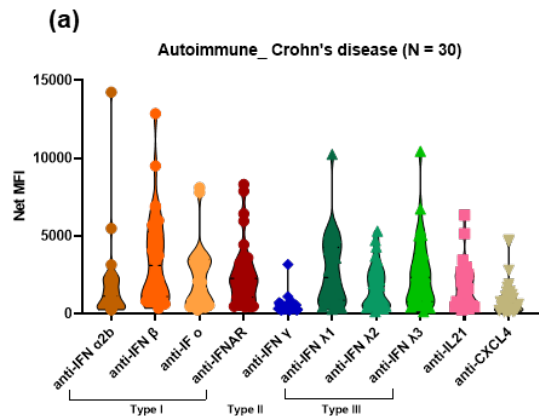
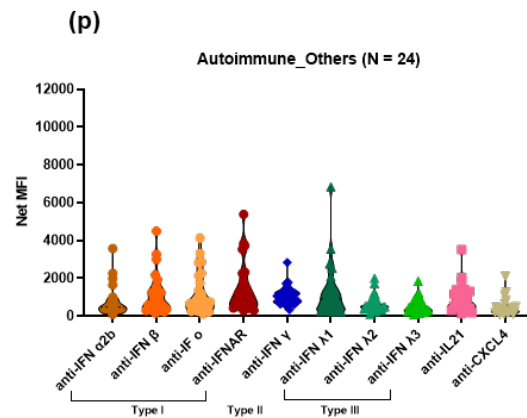
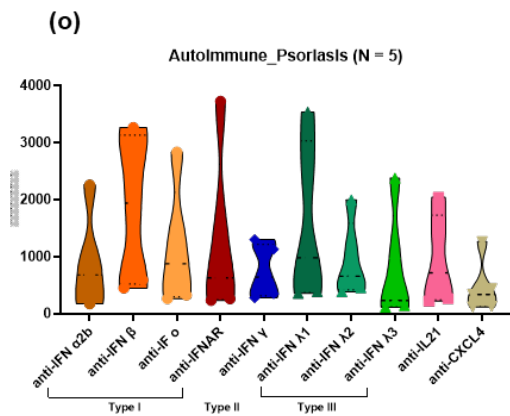
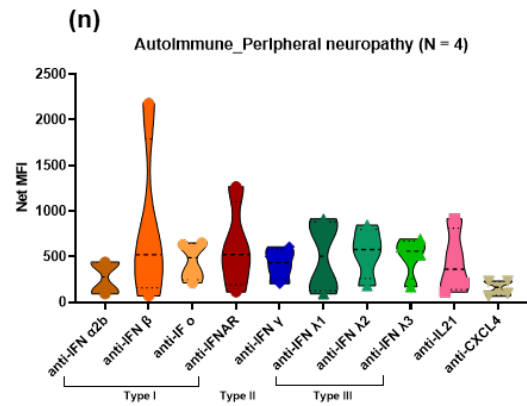
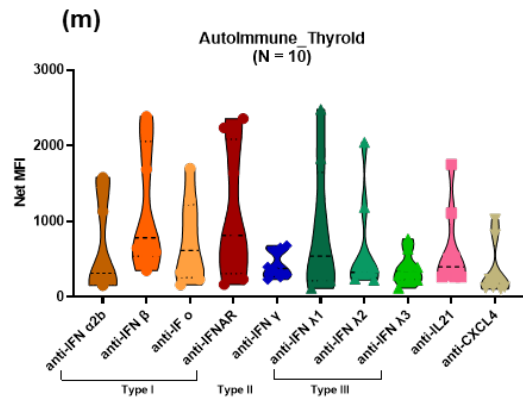
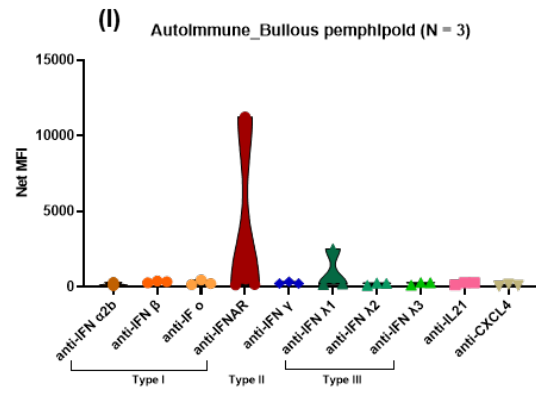
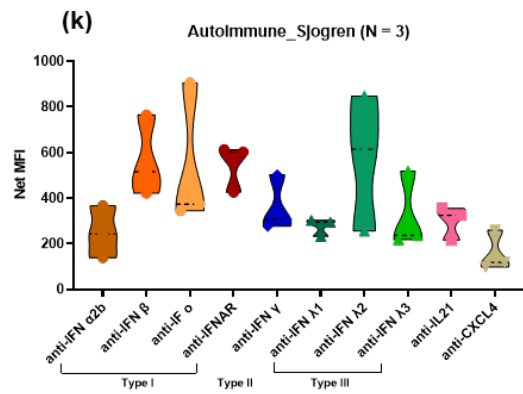
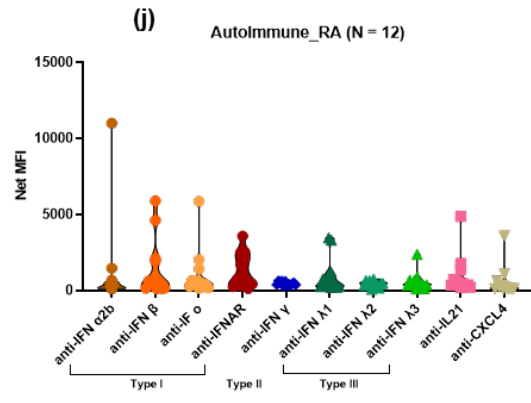
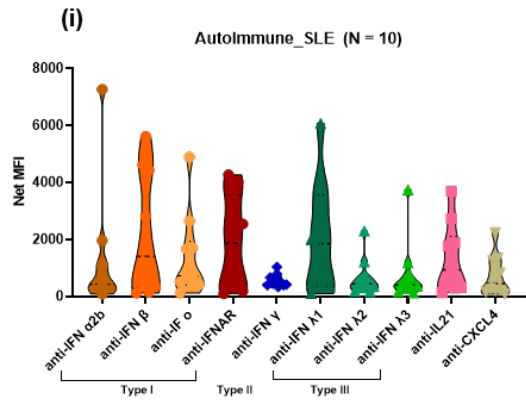
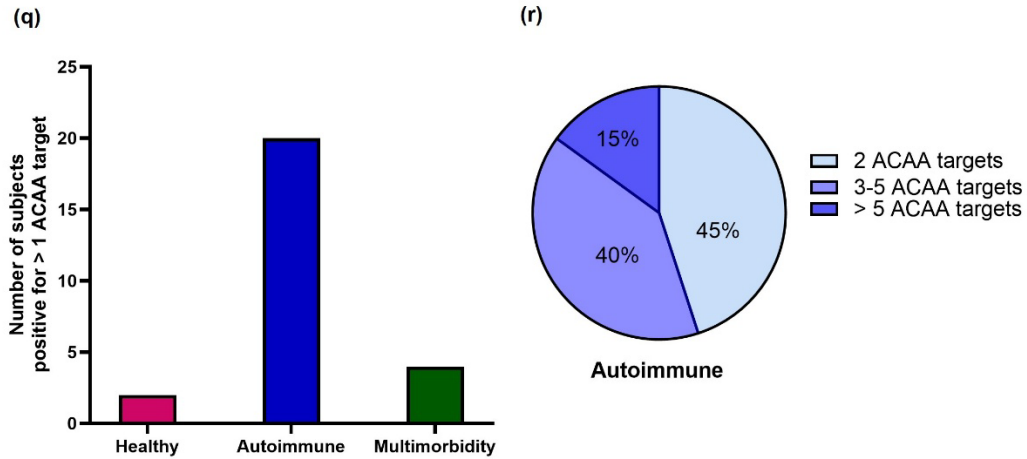
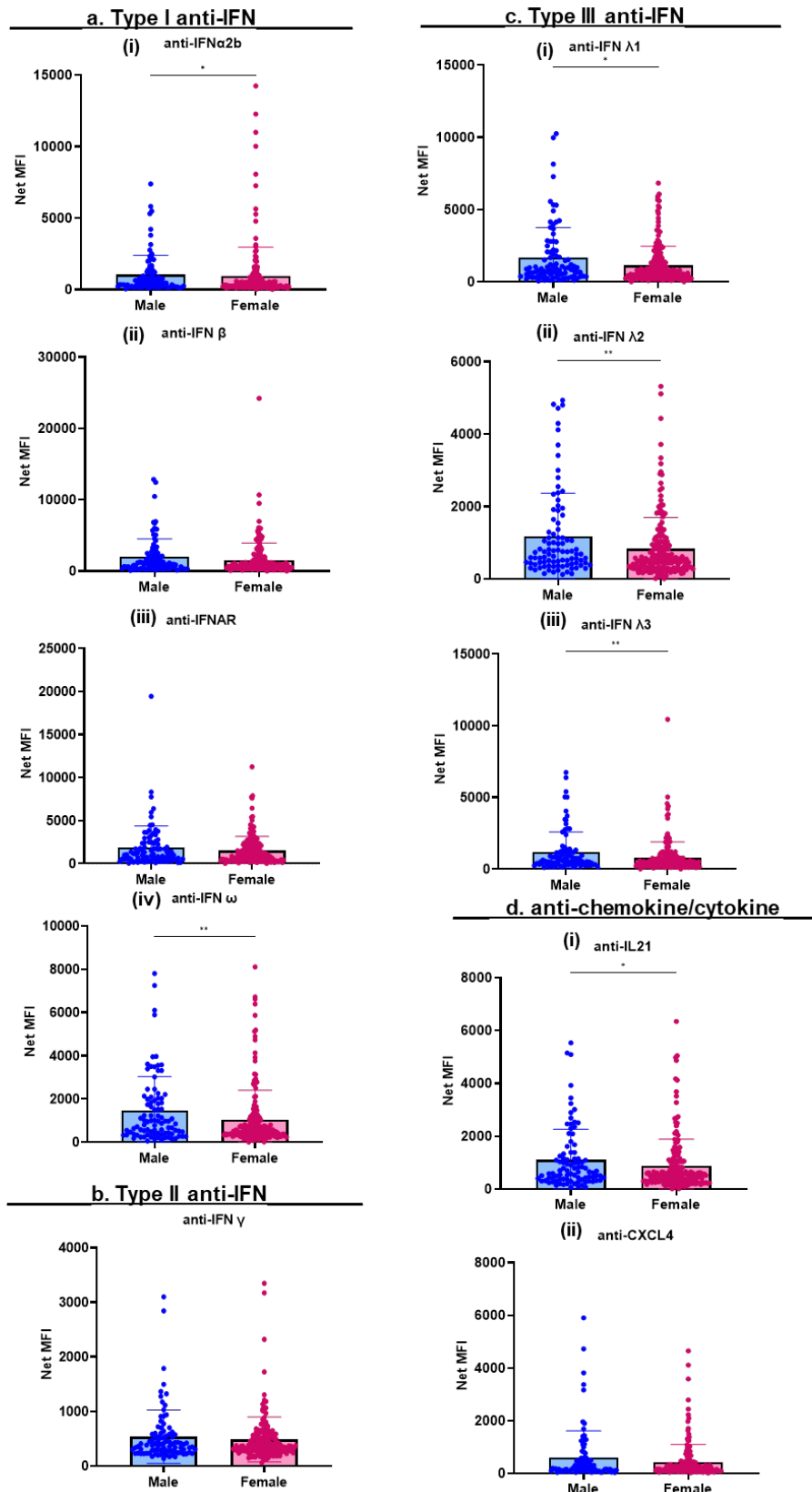








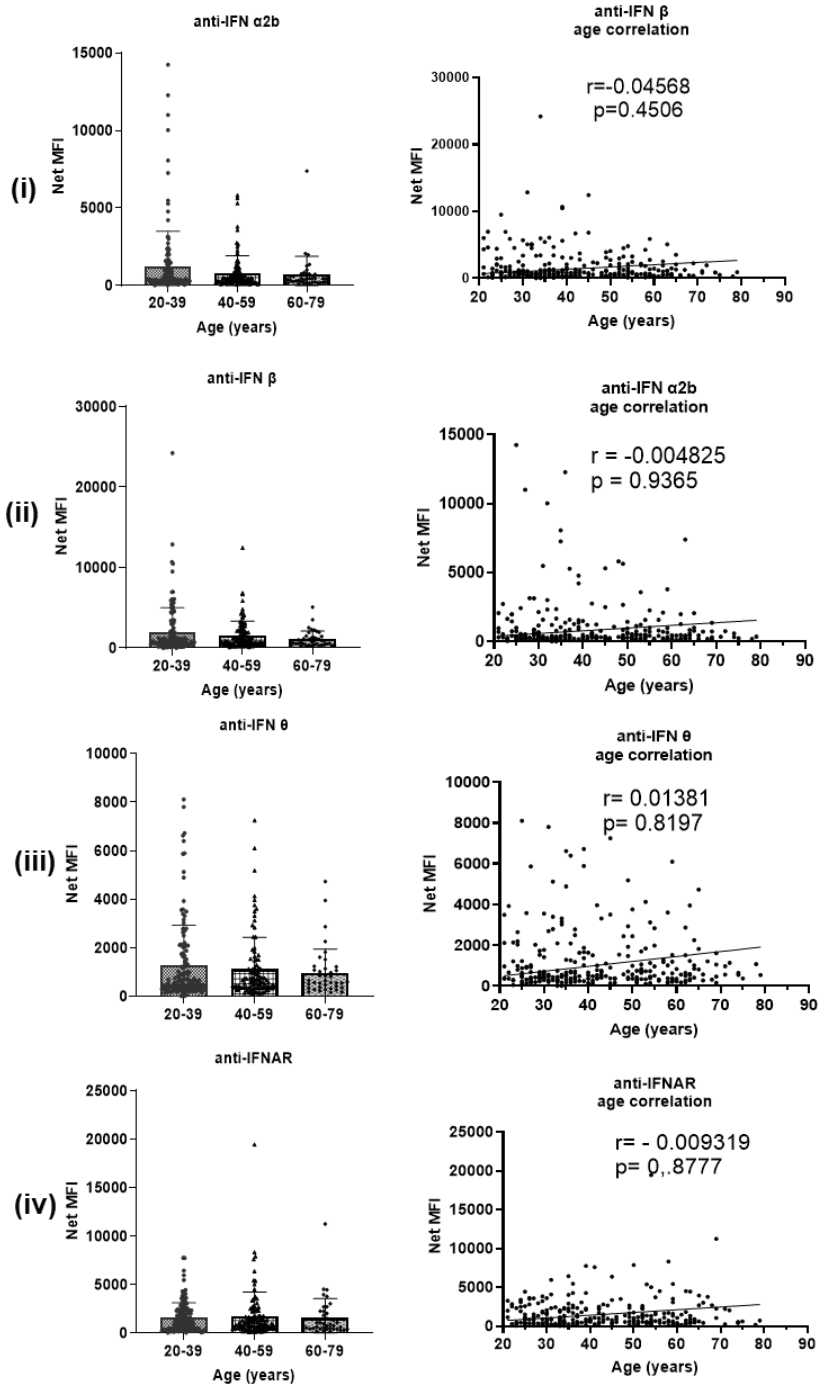
Supplementary figure 1: An autoimmune disease-specific representation of the anti-cytokine autoantibody levels. The autoimmune group was stratified based on the various conditions presented by the individuals. Truncated box-violin plots were employed to display the net MFI autoantibody levels across the different targets evaluated in the magnetic-bead based Luminex assay. The plots are represented as follows: (a-p) Each graph represents one specific anti-cytokine autoantibody target. All data is represented as mean $\pm$ S.E.M for 3 technical replicate experiments. (q)- The bar graphs represent the number of subjects across the 3 different subsets, that display net MFI values above the defined threshold values for more than 1 ACAA target. (q-r)The pie chart focuses on the autoimmune group and is divided into percentage of subjects which display net MFI values above the defined threshold values for 2, 3-5 and >5 ACAA targets respectively. The percentage values for each group is reflected on the pie chart.



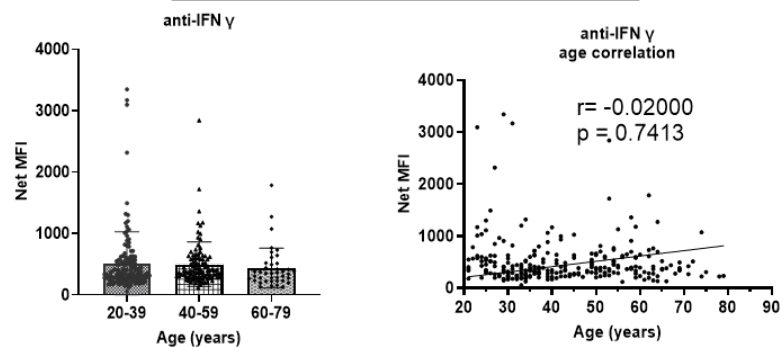
Supplementary figure 2: Assessment of sex as a demographic factor influencing autoantibody levels:

The measured net MFI autoantibody levels for the cohort was distributed based on males vs females for each target on the anti-cytokine autoantibody panel. Dotplots were used to represent the comparison of net MFI levels between the 2 groups. a (i-iv)- Type I anti- IFNs, b- Type II anti-IFN, c (i-iii)-Type III anti-IFNs and d (i and ii)-anti-cytokine/anti-chemokine. Each graph evaluates one specific anti-cytokine autoantibody. Each dot represents the net MFI value of one individual. The statistical evaluation was performed using the Mann-Whitney. All data is represented as mean $\pm$ S.E.M for 3 technical replicate experiments.

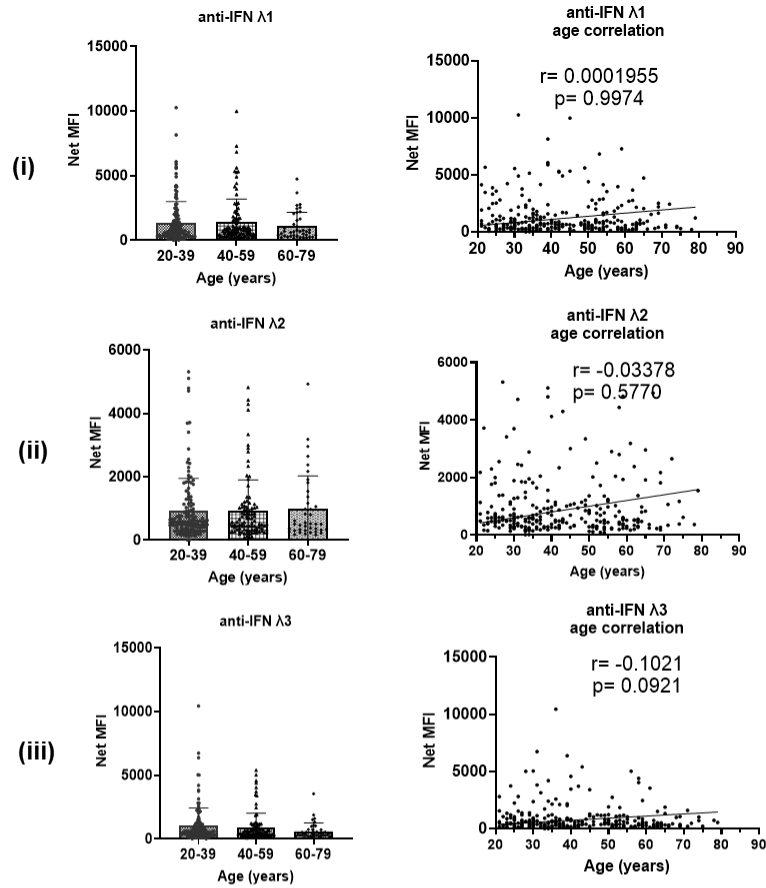
a. Type I anti-IFN



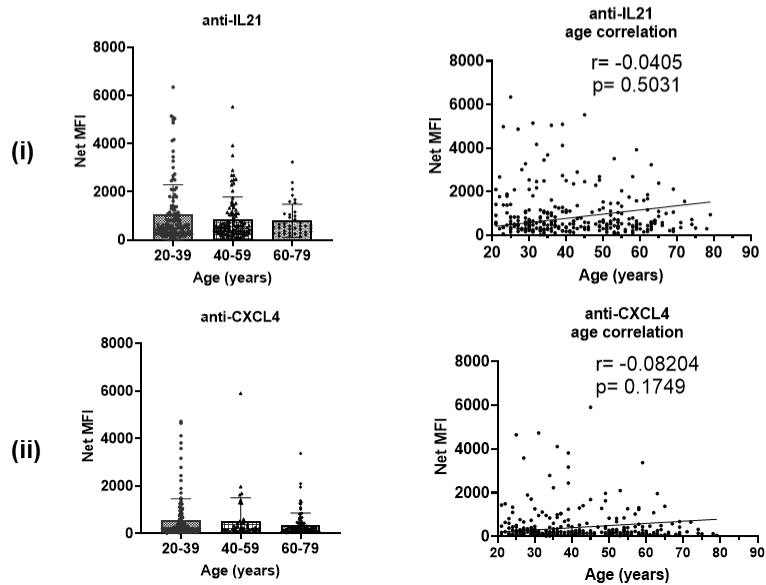
b. Type II anti-IFN



c. Type III anti-IFN

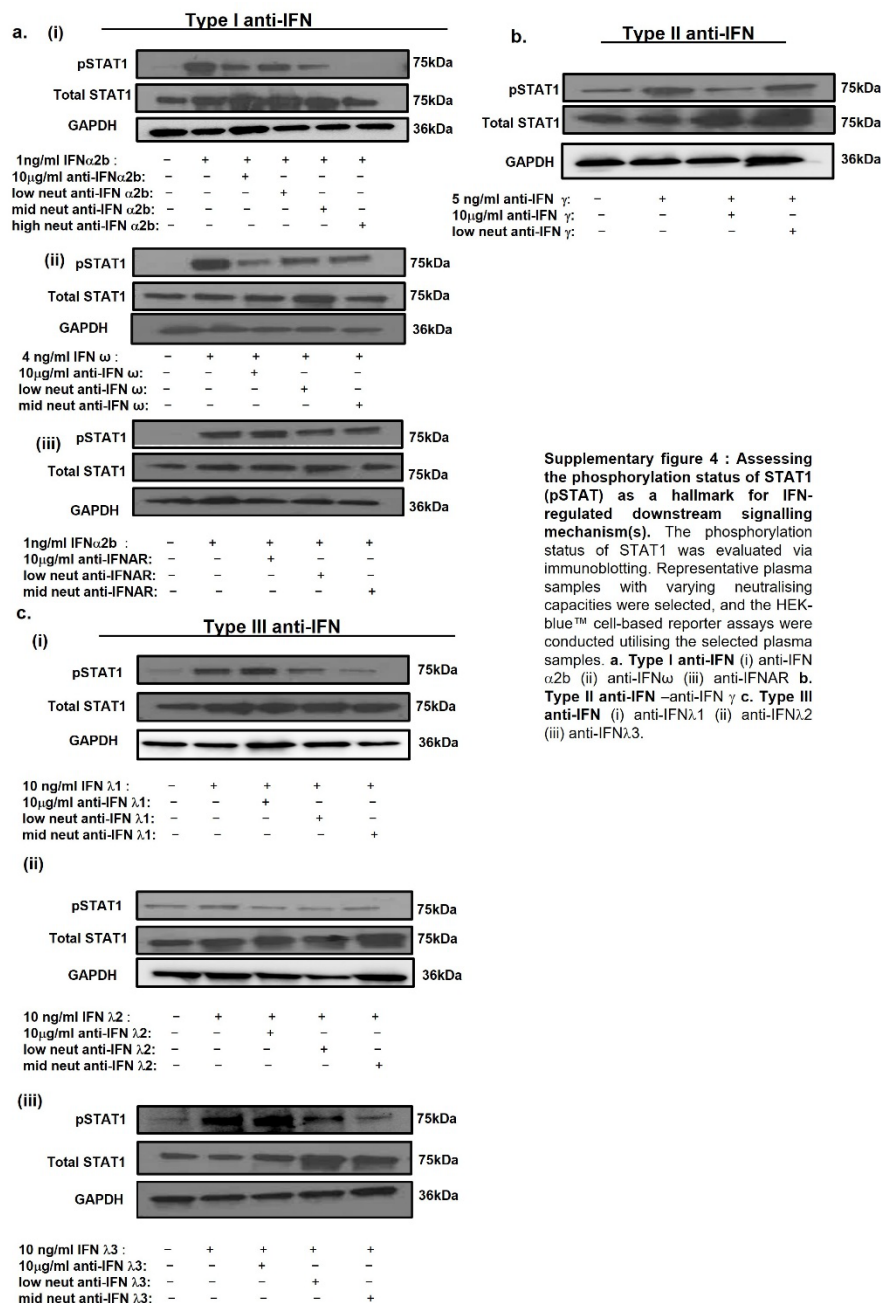


d. anti-chemokine/cytokine



Supplementary figure 3: Assessment of age as a demographic factor influencing autoantibody levels:

The measured net MFI autoantibody levels for the cohort was distributed based on 3 major age groups; 20-39, 40-59 and 60-79, for each target on the anti-cytokine autoantibody panel. Dotplots were used to represent the comparison of net MFI levels within the age groups. a (i-iv)- Type I anti- IFNs, b- Type II anti-IFN, c (i-iii)-Type III anti-IFNs and d (i and ii)- anti-cytokine/anti-chemokine. Each graph evaluates one specific anti-cytokine autoantibody. Each dot represents the net MFI value of one individual. The statistical evaluation was performed using the Kruskal-Wallis test followed by Dunn's test. Simple linear regression analyses were conducted, followed by a non-parametric spearman correlation analysis. Each graph evaluates one specific anti-cytokine autoantibody. Each dot represents the net MFI value of one individual. The spearman correlation r is indicated on each linear regression graph. All data is represented as mean $\pm$ S.E.M for 3 technical replicate experiments.



Supplementary figure 4 : Assessing the phosphorylation status of STAT1 (pSTAT) as a hallmark for IFN-regulated downstream signalling mechanism(s). The phosphorylation status of STAT1 was evaluated via immunoblotting. Representative plasma samples with varying neutralising capacities were selected, and the HEK-blue™ cell-based reporter assays were conducted utilising the selected plasma samples. **a. Type I anti-IFN** (i) anti-IFN α 2b (ii) anti-IFN ω (iii) anti-IFNAR **b. Type II anti-IFN** –anti-IFN γ **c. Type III anti-IFN** (i) anti-IFN λ 1 (ii) anti-IFN λ 2 (iii) anti-IFN λ 3.

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	Net MFI threshold value
anti-IFN α 2b	4720.46
anti-IFN β	5710.07
anti-IF γ	4118.30
anti-IFNAR	6056.31
anti-IFN λ 1	1869.98
anti-IFN λ 2	4541.81
anti-IFN λ 3	3505.28
anti-IL21	4358.37
anti-IL21	3209.20
anti-CXCL4	1847.49

Supplementary table 1: The average net MFI levels for each specific autoantibody target were calculated for all 3 groups representing the study cohort, and tabulated as shown above.

	Average Net MFI values		
	Healthy	Autoimmune	Multimorbidity
anti-IFN α 2b	682.28	1151.80	1024.22
anti-IFN β	1205.67	2096.70	1441.52
anti-IF γ	820.42	1350.31	1267.82
anti-IFNAR	1402.26	1714.70	1775.80
anti-IFN λ 1	433.44	588.86	364.38
anti-IFN λ 2	918.76	1526.00	1371.89
anti-IFN λ 3	793.41	926.21	1195.55
anti-IL21	933.15	1004.77	629.83
anti-IL21	661.24	1080.84	1061.70
anti-CXCL4	345.92	646.80	333.78

Supplementary table 2: Net MFI threshold values for each autoantibody target was established by assigning the healthy individuals as a reference group and applying the following formula: (Net MFI threshold= $\text{Mean}_{\text{MFI healthy}} \pm 3 * \text{Av. Std.dev}_{\text{MFI healthy}}$). The respective threshold values were then tabulated as shown above.

(Autoimmune)

Biomarker regulated	Type of ACAA displaying significant association					
IL-12B	anti-IFN β	anti-IFN λ 2	anti-IFN λ 3			
IL17A	anti-IFN α 2b	anti-IFN ω	anti-IFNAR	anti-IFN λ 1	anti-IL21	
MCP3	anti-IFN β	anti-IFN λ 1	anti-CXCL4			
CASP8	anti-IFN α 2b	anti-IFN ω	anti-IFNAR	anti-IFN λ 1	anti-IL21	
GDNF	anti-IFN λ 1					
TNF	anti-IFNAR					

Supplementary table 3: Significant associations between target biomarker and ACAA levels in the autoimmune group.

(Multimorbidity)

Biomarker regulated	Type of ACAA displaying significant association					
CCL20	anti-IFN α 2b	anti-IFN ω				
CCL19	anti-IFN α 2b	anti-IFN β	anti-IFN ω	anti-IFN λ 1	anti-IL21	
TNFRSF9	anti-IFN ω	anti-IFN γ	anti-CXCL4			
FGF23	anti-IL21					
MCP3	anti-IFN β					
SLAMF1	anti-IFN ω	anti-IL21				
FGF5	anti-IFNAR	anti-IFN λ 1				
IL15RA	anti-IFN λ 3					
GDNF	anti-IFN λ 2	IL-21				
CDCP1	anti-IFN ω					
CXCL10	anti-CXCL4					
CCL25	anti-IFN ω	anti-CXCL4				

Supplementary table 4: Significant associations between target biomarker and ACAA levels in the multimorbidity group