

Single-cell DN2 B Cell Atlas Reveals Novel Subtypes and Demographic Trends

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Motivation

- Double-negative (DN2) B cells are a novel type of B cells that **lack IgD and CD27**
- DN2 B cells are found abundantly within patients with **autoimmune diseases, elderly populations, and recently, in COVID-19 patients with severe symptoms**
- There is a lack of large-scale genomic research being done on DN2 B cells; establishing a cross-population atlas to further understand this novel immune cell is of high interest

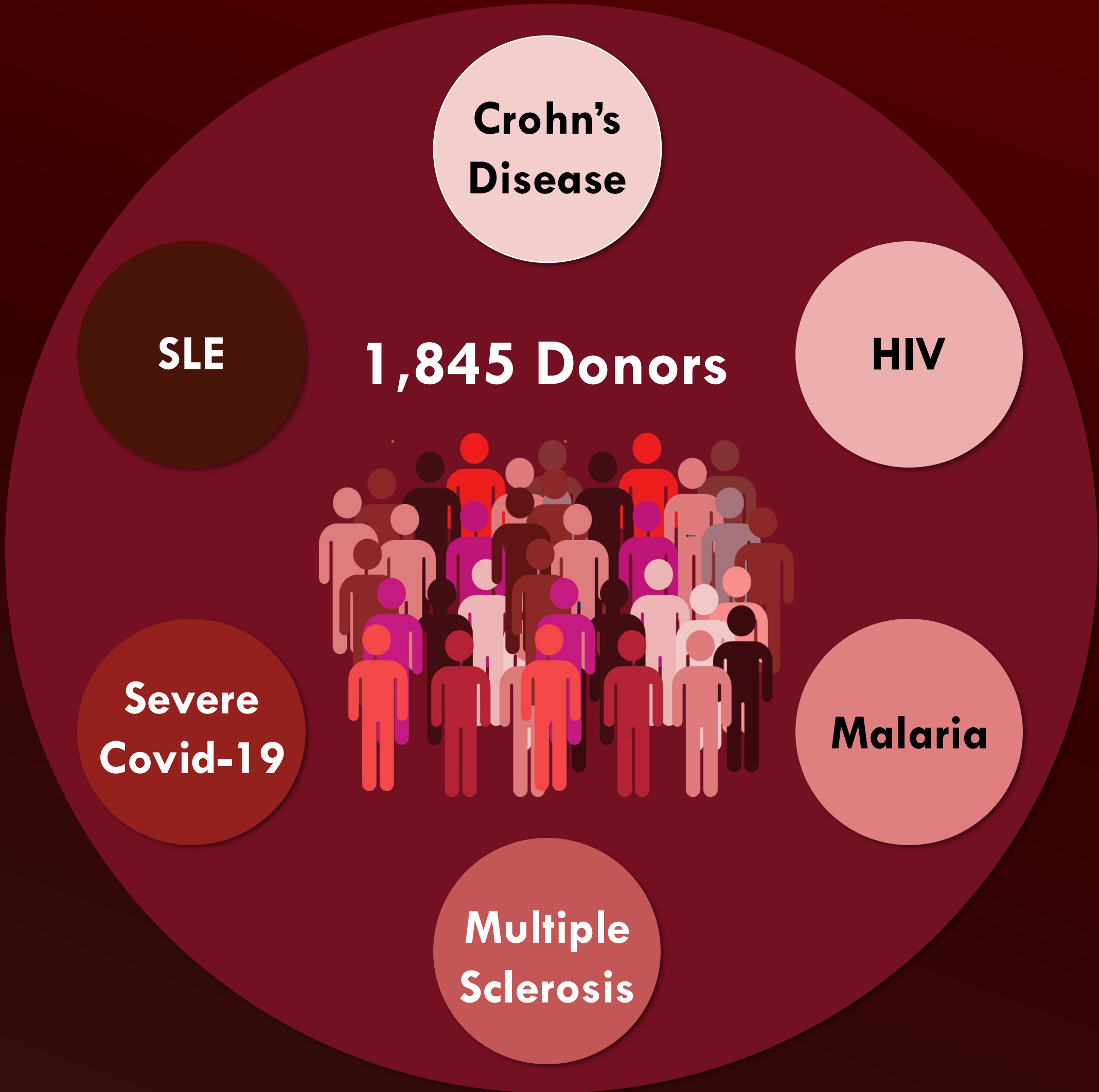
Aims:

- Establish** a healthy baseline for DN2 B cells by creating a DN2 atlas from 4 healthy PBMC datasets
- Incorporate** PBMC datasets from immune-related diseases into the atlas
- Assess** demographic variations and **Compare** DN2 characteristics in healthy and disease populations

METHODS

The Healthy and Disease DN2 cell Atlas, a harmonized, metadata-rich, cross-disease, single-cell omics resource, spanning over **30K DN2 B cells**.

Data acquisition:
Publicly available scRNA-seq data (Gene Expression Omnibus)



RESULTS

1 An Integrated B Cell Map of Health and Disease

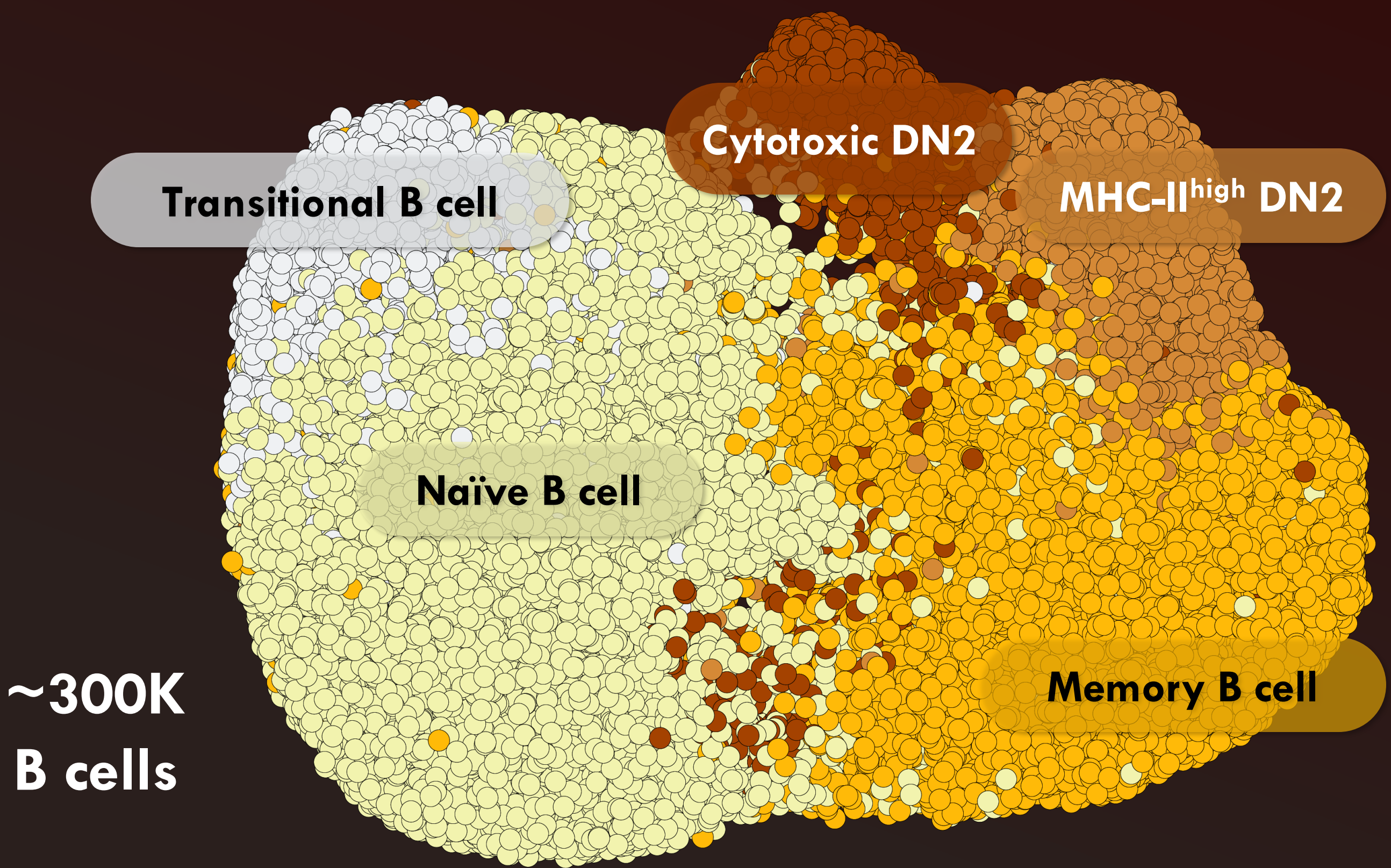


Fig. 1a | UMAP projection of healthy & diseased ~300K human peripheral blood mononuclear B cells using scRNA-seq and clustered into 5 subsets.

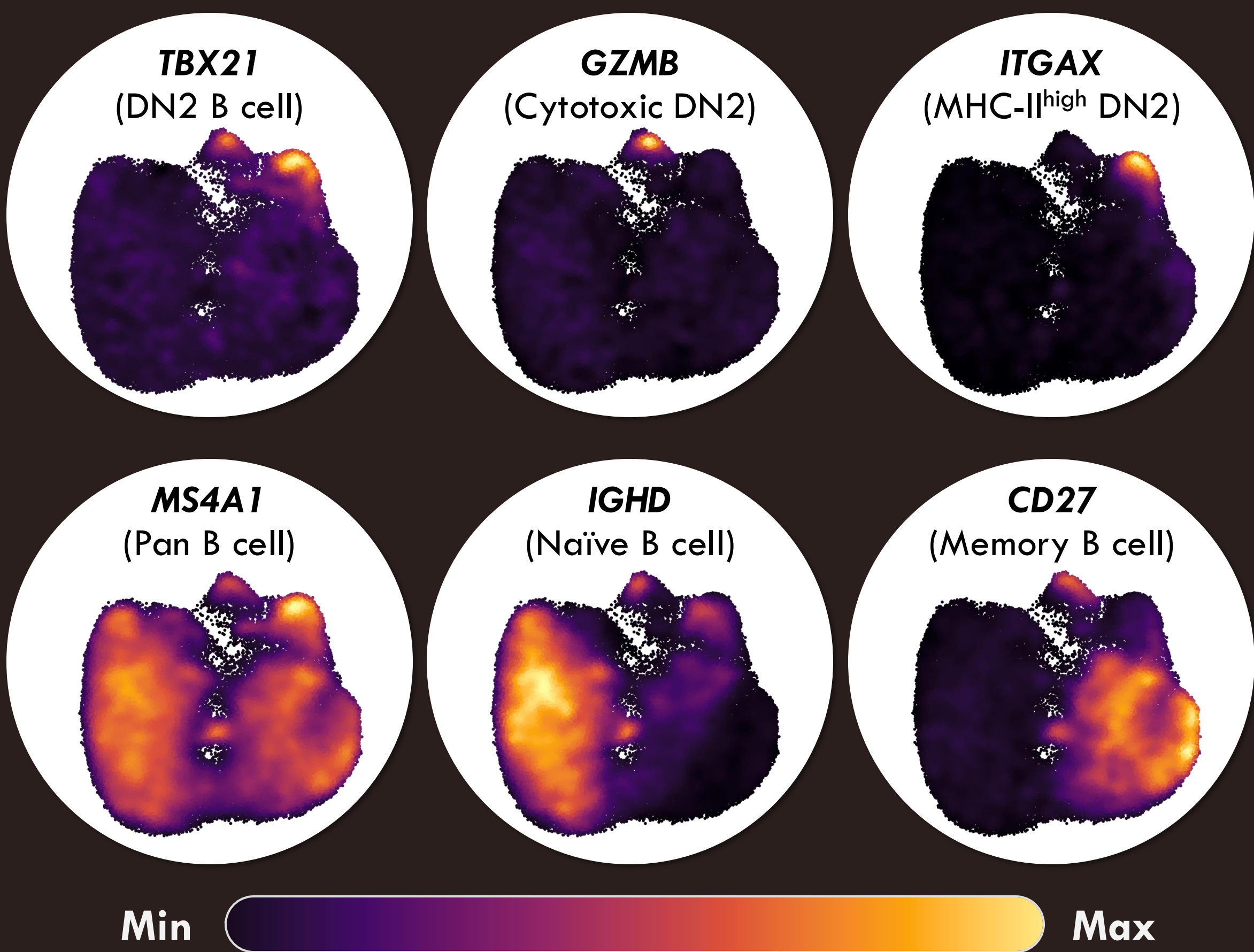


Fig. 1b | Density maps of above UMAP projection with characteristic B cell markers.

2 Proportion Analysis by Disease & Demographics

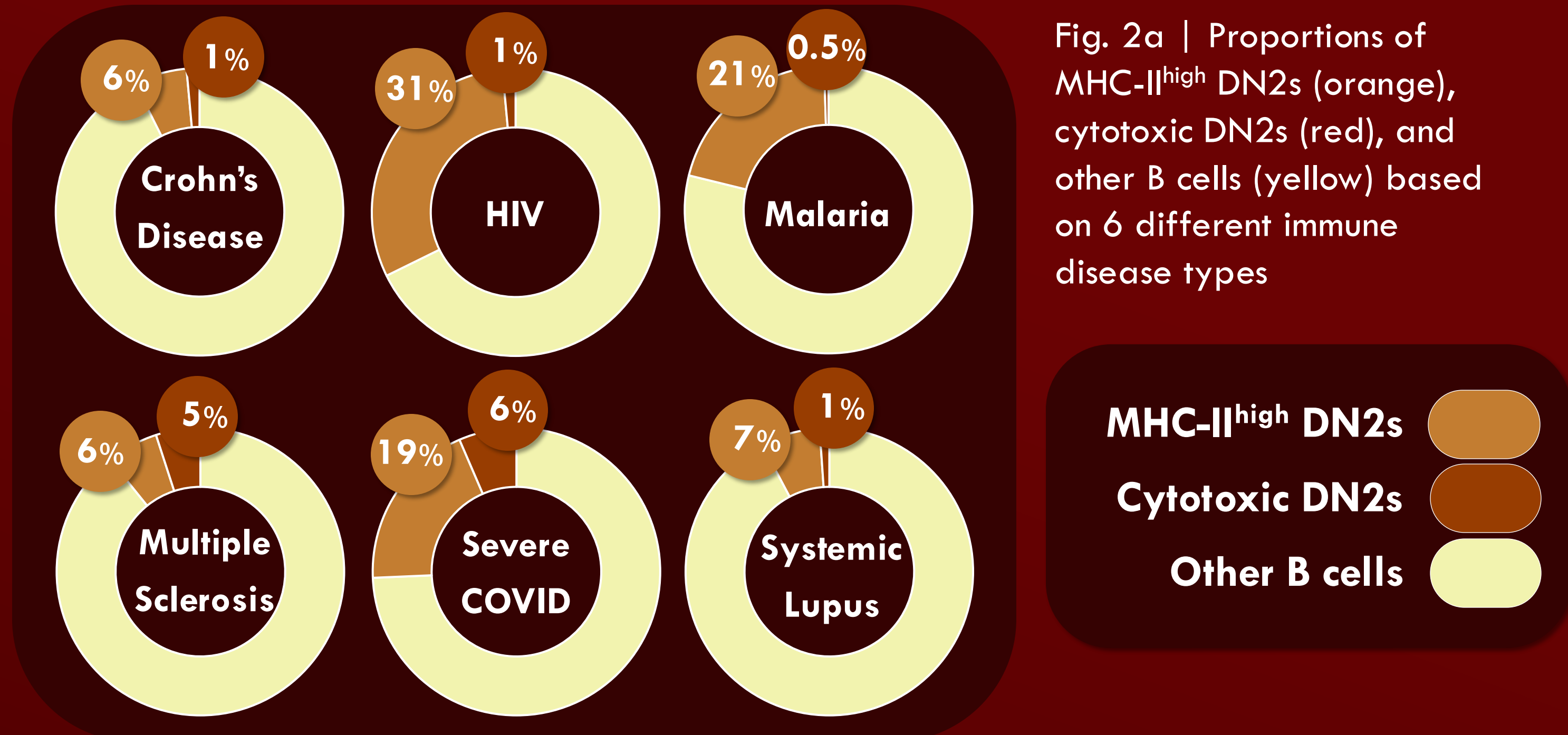


Fig. 2a | Proportions of MHC-II^{high} DN2s (orange), cytotoxic DN2s (red), and other B cells (yellow) based on 6 different immune disease types

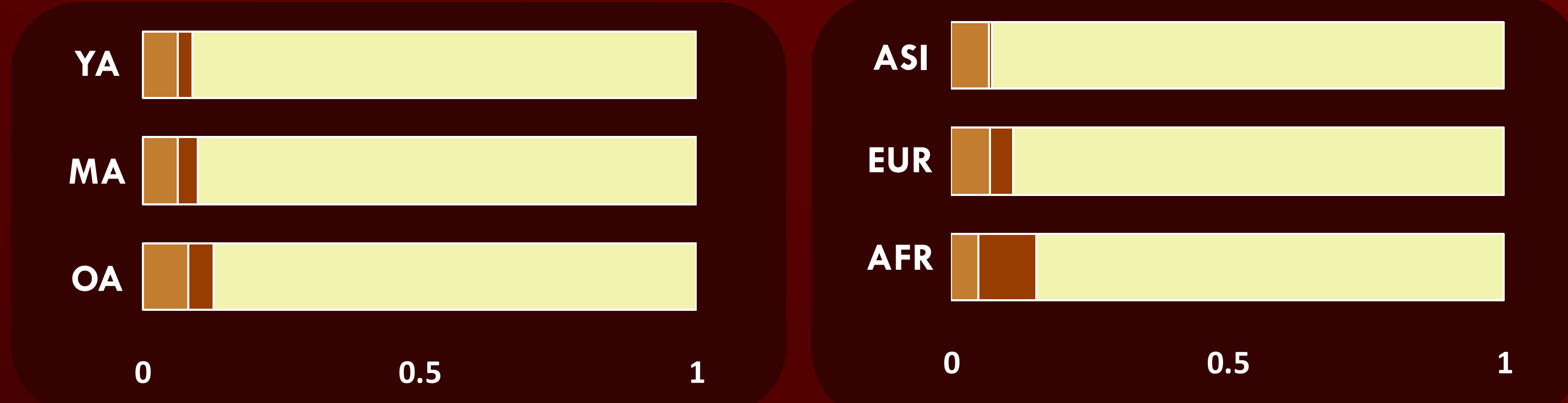


Fig. 2b | Proportions of classical DN2s (orange), cytotoxic DN2s (red), and other B cells (yellow) in healthy individuals based on age differences (Young Adult (YA) <45yrs, Middle-aged (MA) 46-65yrs, Older Adult (OA) >65 yrs)

Fig. 2c | Proportions of classical DN2s (orange), cytotoxic DN2s (red), and other B cells (yellow) in healthy individuals based on ethnicity differences (ASI= Asian, EUR= European, AFR= African American)

3 Signaling Pathways Associated with Disease

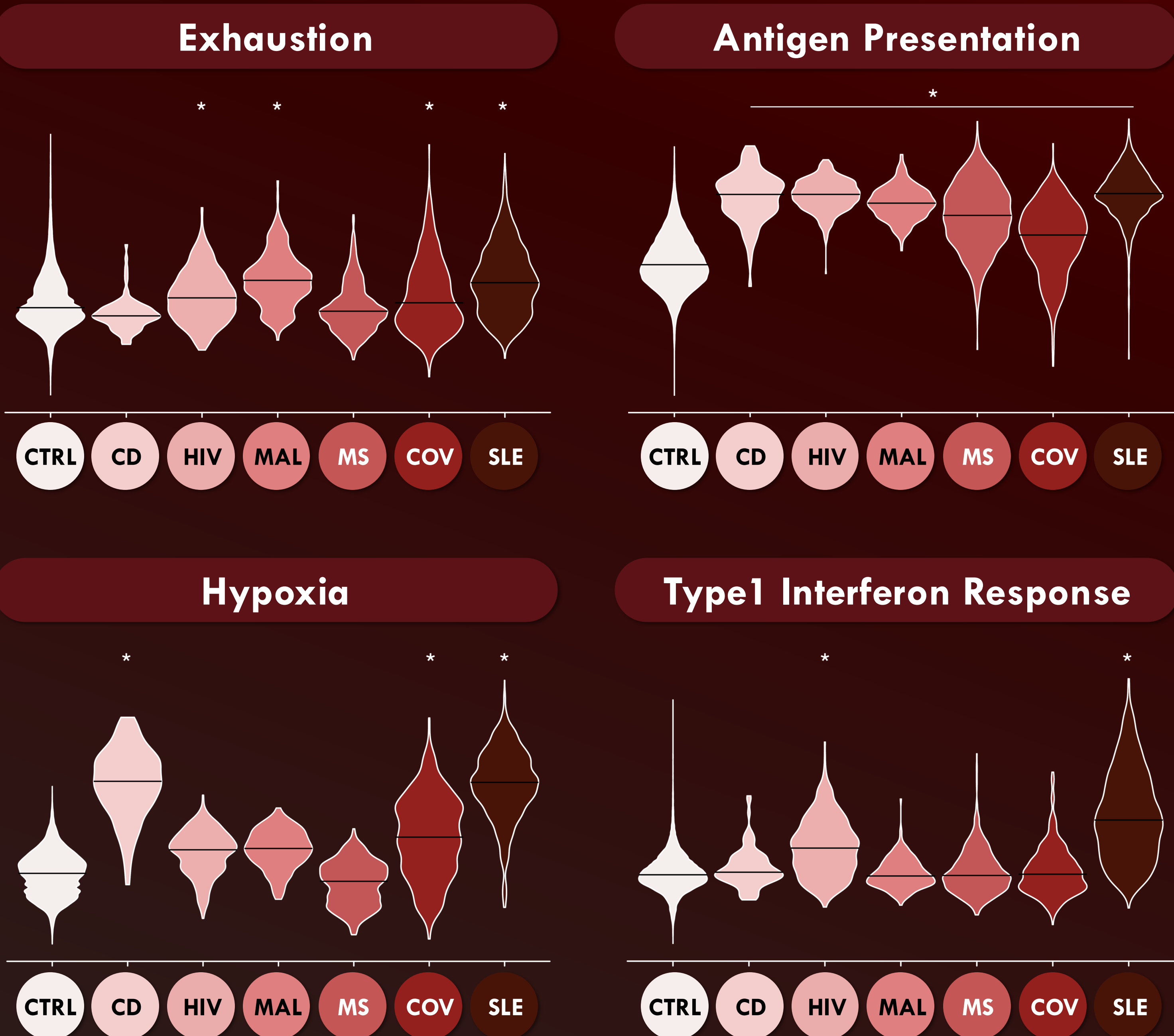


Fig. 3 | Violin plots depicting average expression of exhaustion, antigen presentation, hypoxia, and type 1 interferon-stimulated gene (ISG) signatures in DN2 B cells across individuals in cases and controls. Significant expression changes between cases vs controls are highlighted * $P_{\text{adjusted}} < 0.05$ (Wilcoxon). CTRL= Healthy; CD= Crohn's Disease; MAL= Malaria; MS= Multiple Sclerosis; COV= Severe Covid-19; SLE= Systemic Lupus Erythematosus

CONCLUSION

- scRNA-seq of PBMC reveals **distinct DN2 B cell subpopulations**
- DN2 subpopulations are **differentially expanded across chronic and autoimmune diseases**
- Ratio of DN2s to other B cells **increases with age**
- DN2 B cell subpopulations **vary across age and race**
- HIV and malaria** express a **significantly high proportion of MHC-II^{high} DN2 B cells**
- Increase of **cytotoxic DN2s** in **Severe COVID-19** and **multiple sclerosis**
- Disease DN2s express **distinct gene signatures** including genes encoding **hypoxia, antigen presentation, exhaustion, and type 1 IFN**