

16-18 June 2023



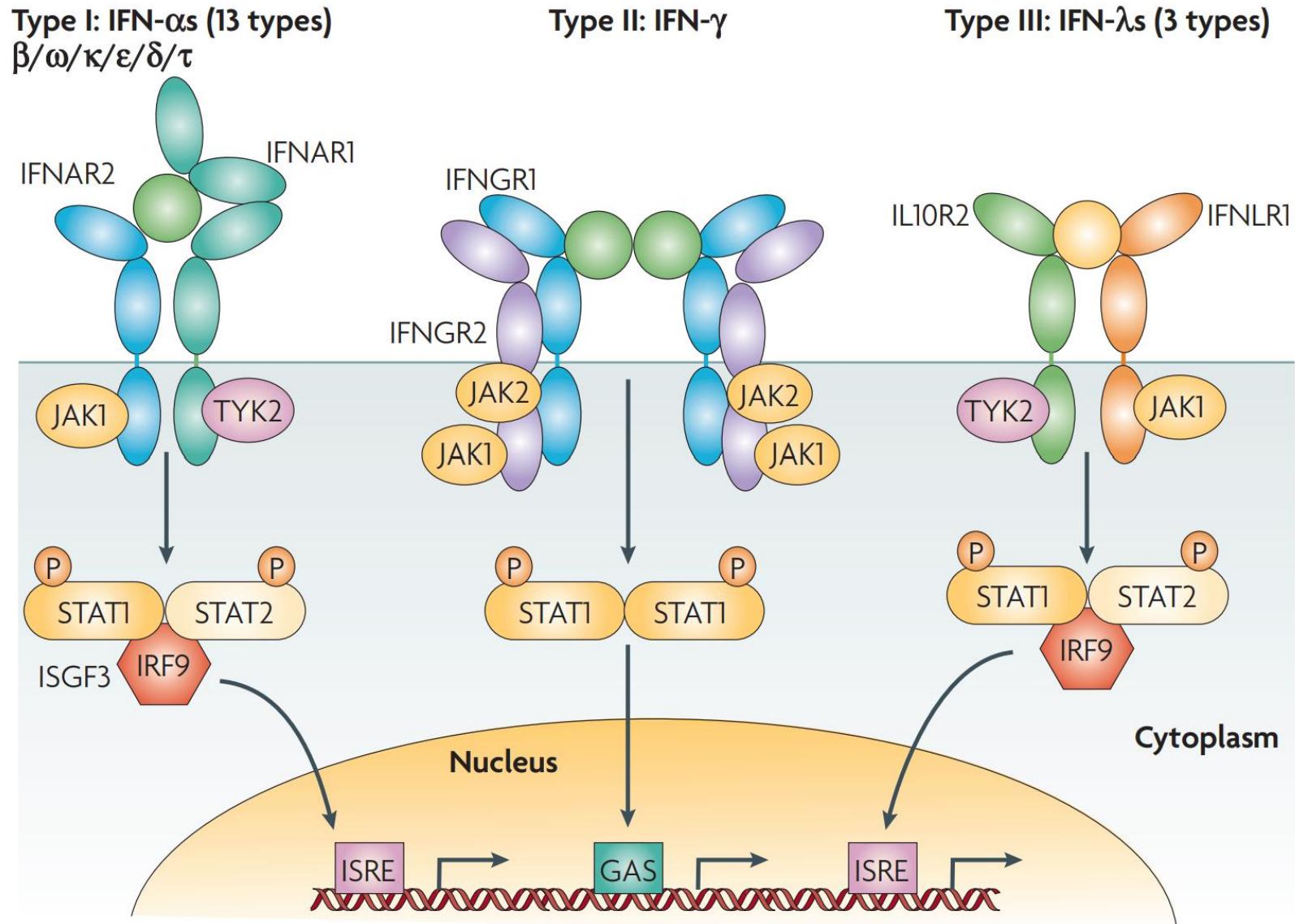
Current Immunological trends in clinical topics:

Cells and pathways producing interferon in health and disease

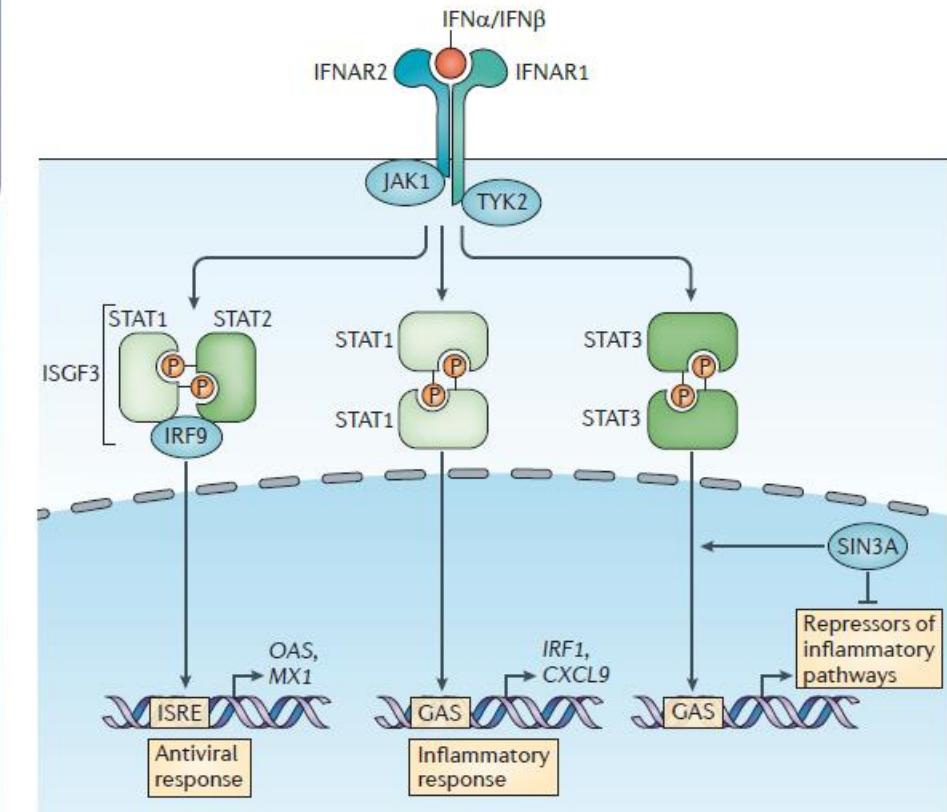
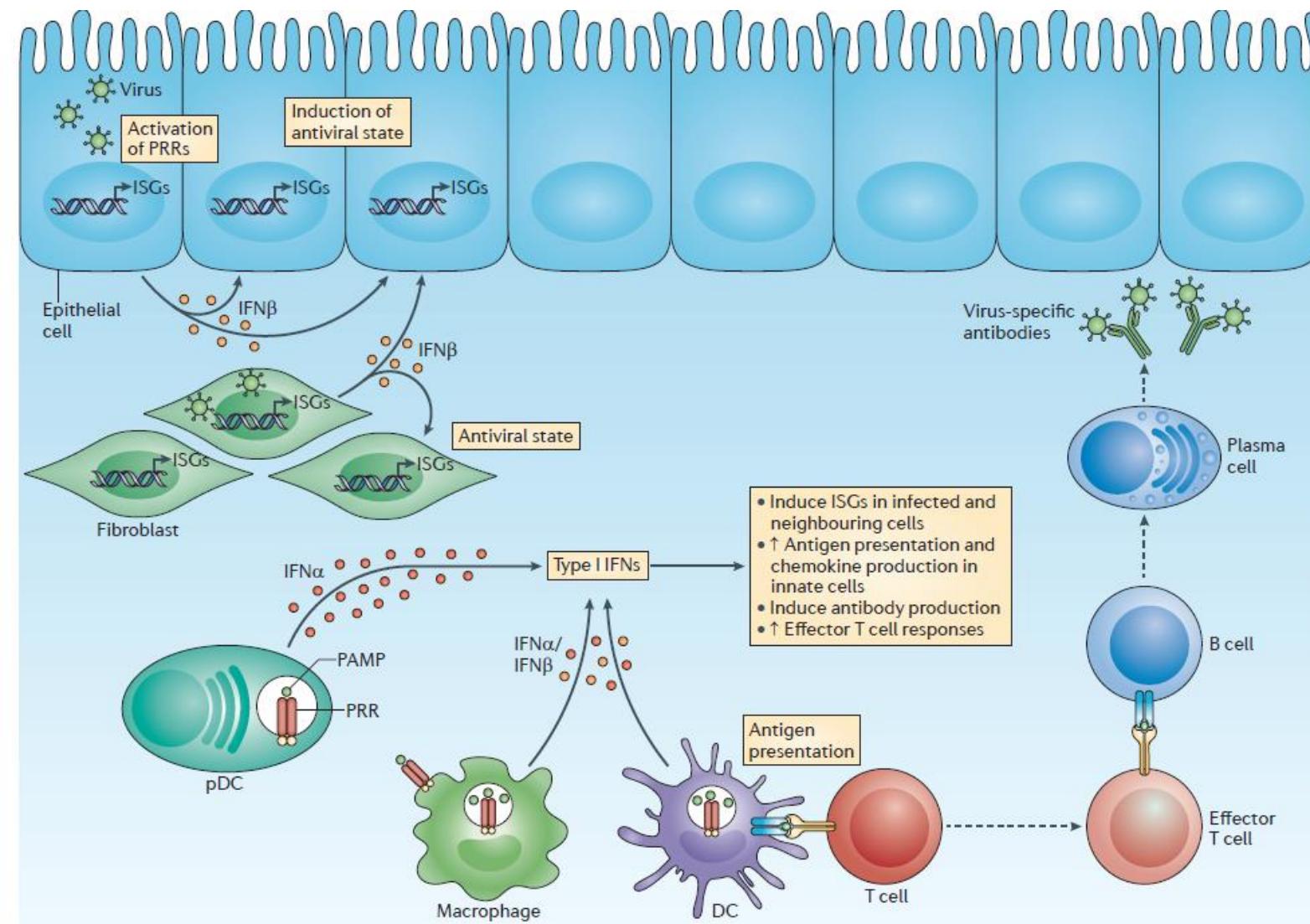


Aggelos Banos,
Rheumatology Resident, Asklepeion Voulas
PostDoctoral Fellow, BRFAA

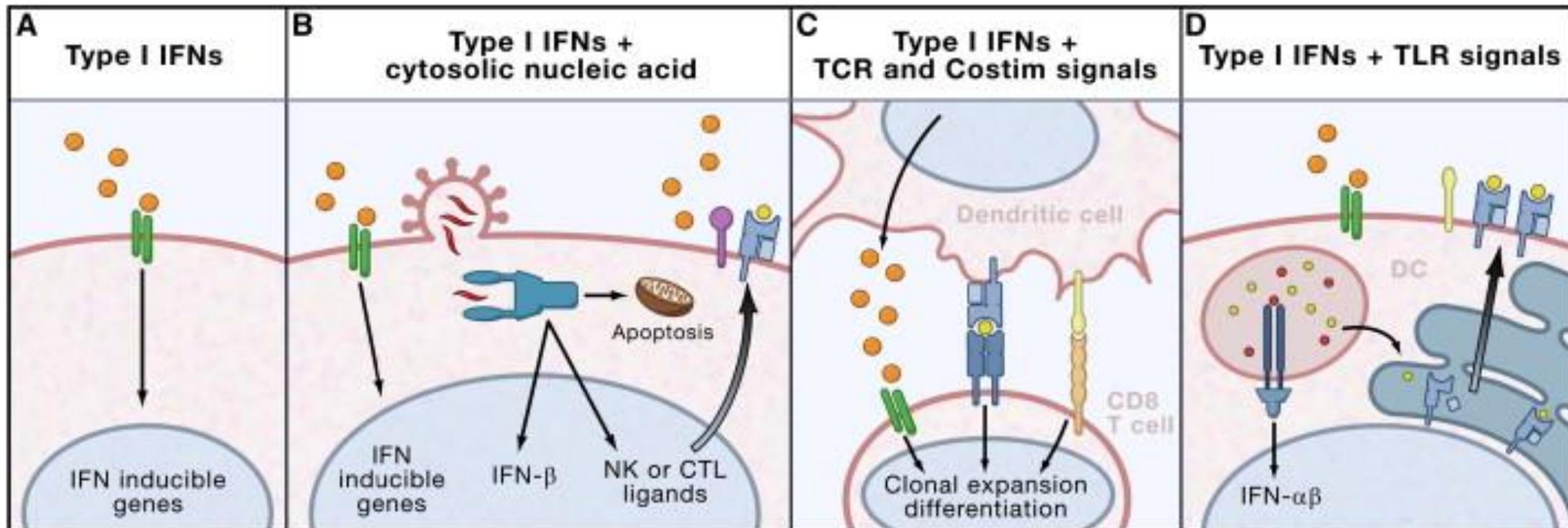
Type I, II and III interferons



Regulation of type I interferon responses

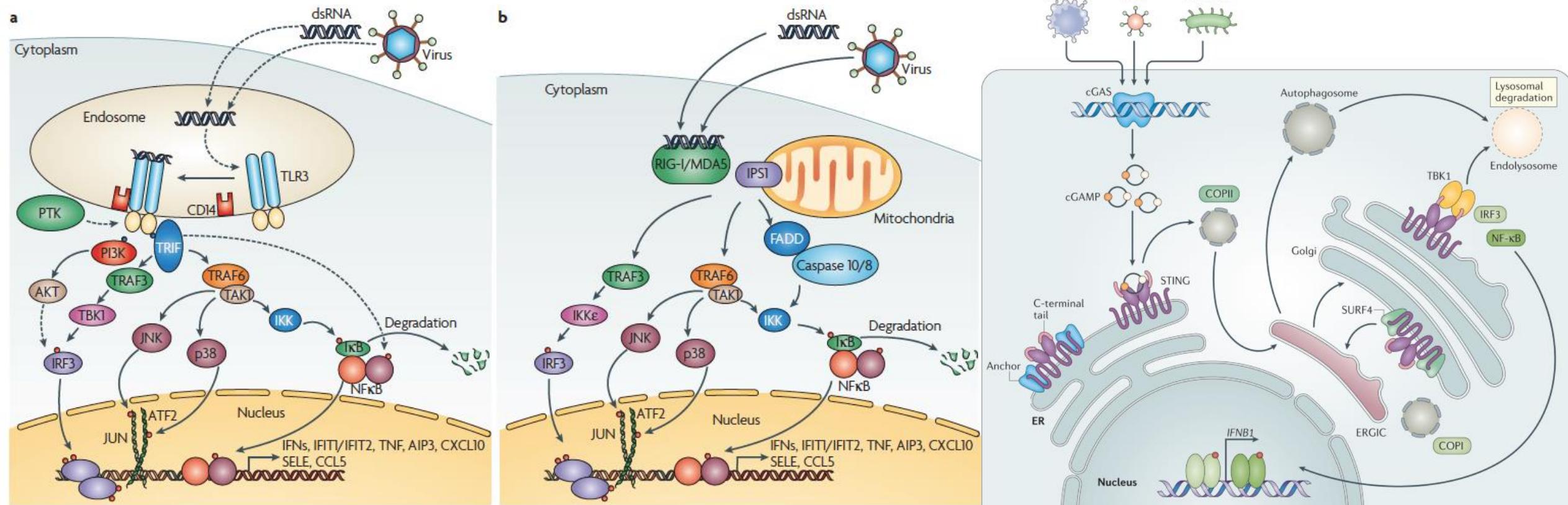


Type I IFN in host defense

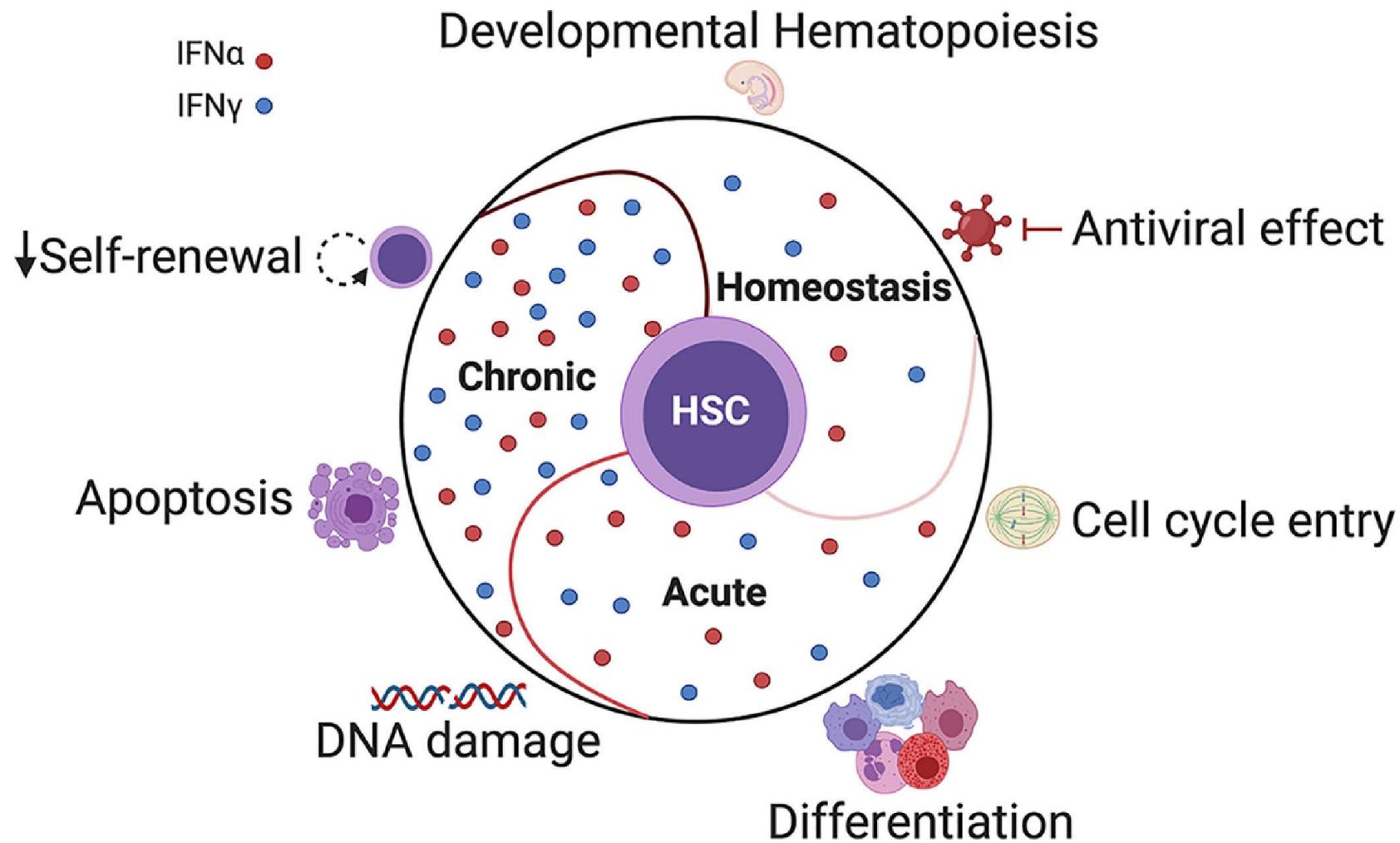


- Base line induction of IFN signaling in uninfected cells
- Infection: sense of cytosolic nucleic acid → IFN beta → paracrine activation
- Clonal expansion and differentitaion of CD8+ cells through IFN signaling
- DCs sense IFNs and cross-present antigens

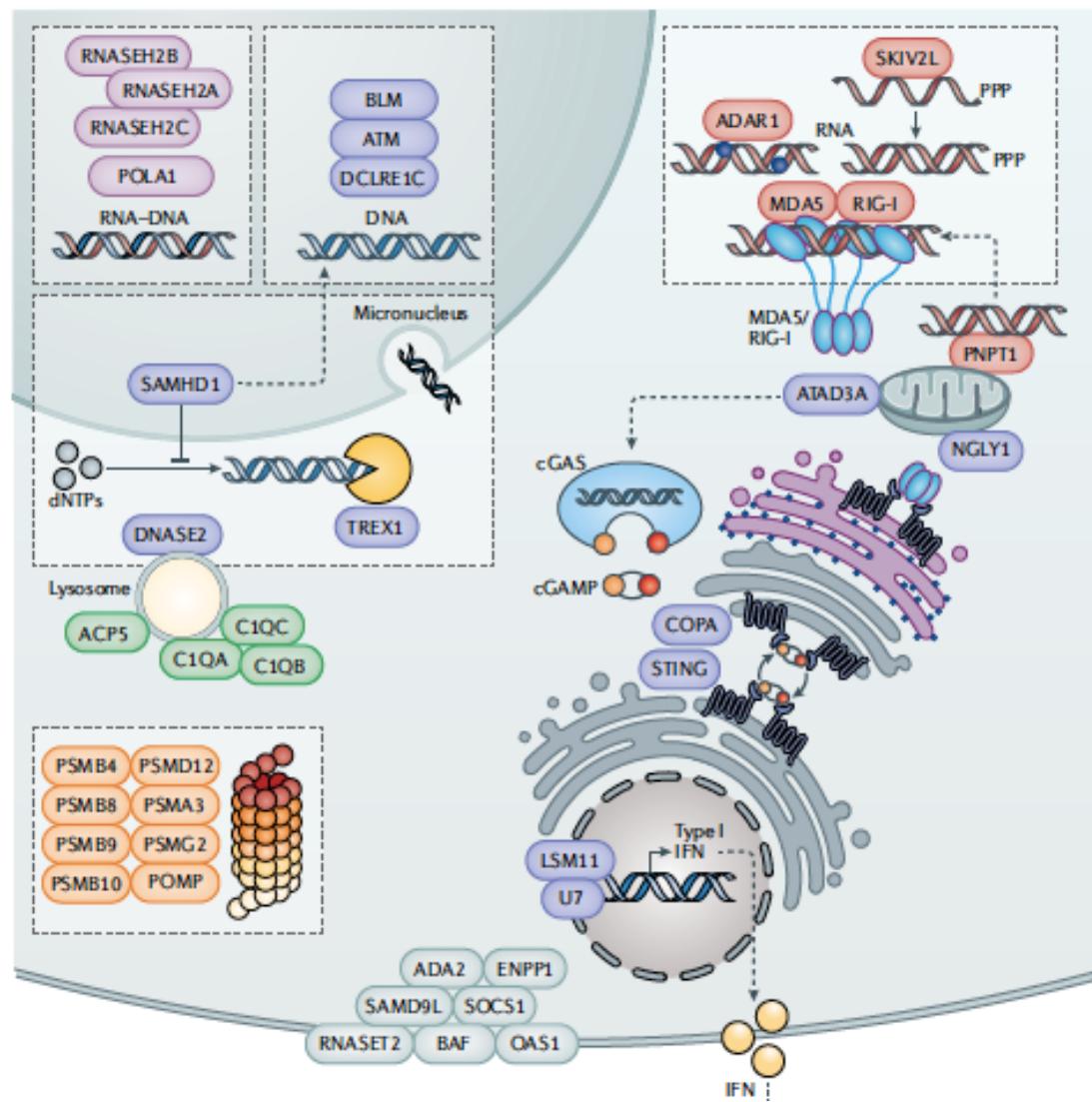
Type I IFN signaling



Effect of interferons in hematopoiesis

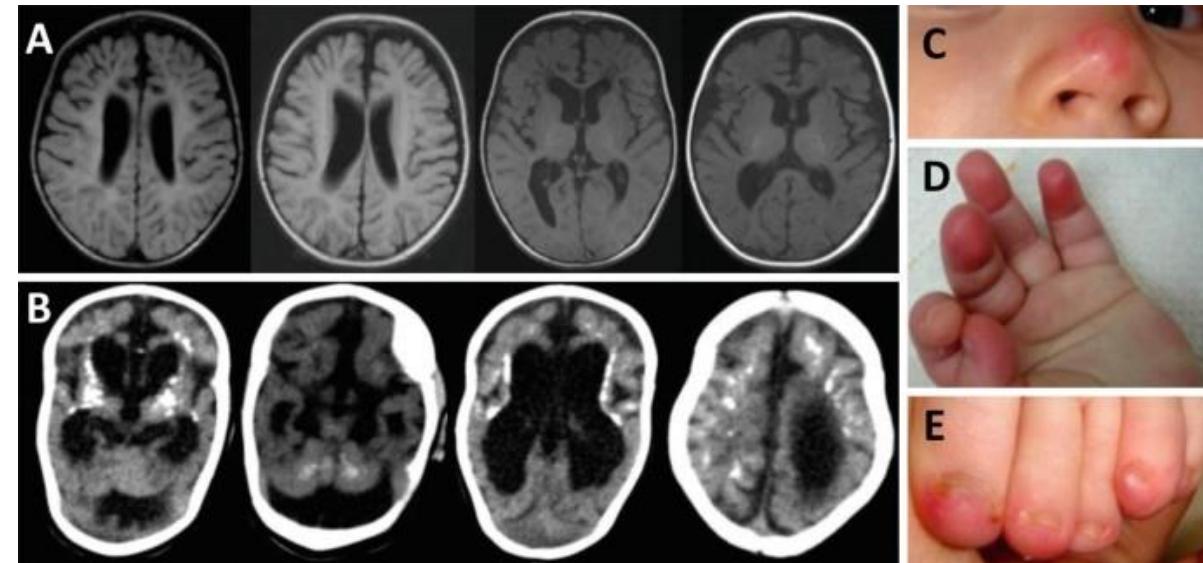


Genetic Interferonopathies



Crow and Stetson, Nat Rev Immun 2022
 Tungler et al, Enc of Med Imm. 2019
 Liu et al NEJM 2014
 Volpi et al, Clinical Immunology 2018

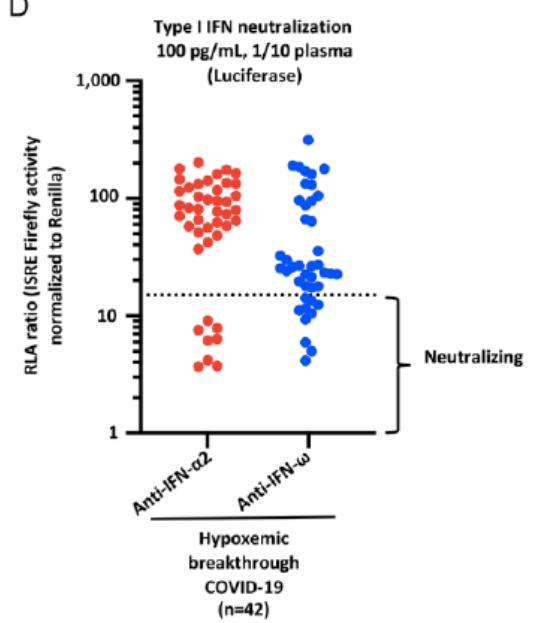
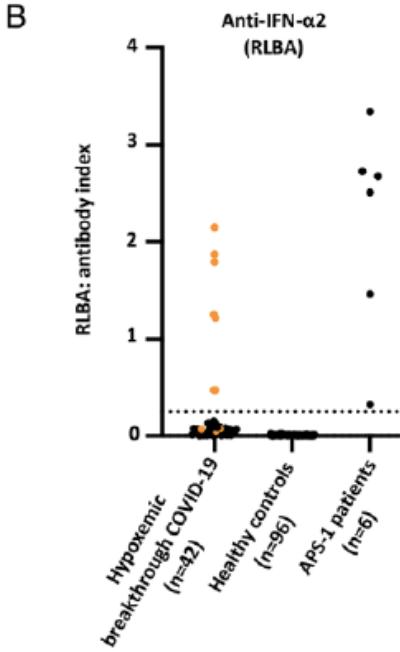
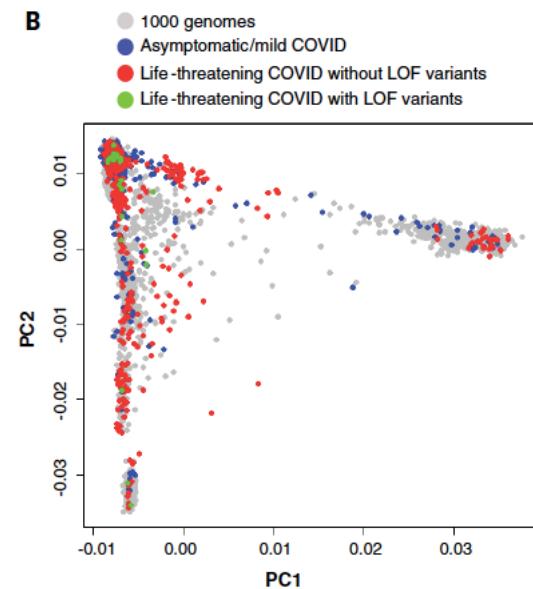
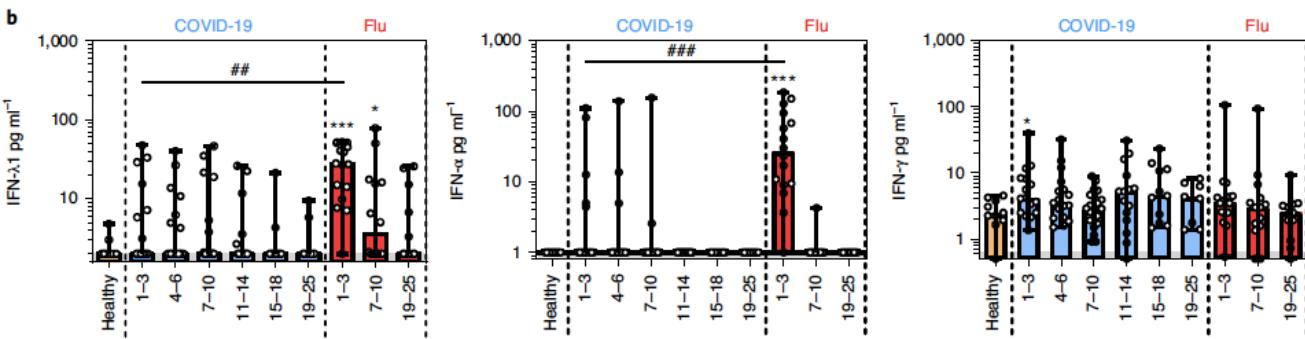
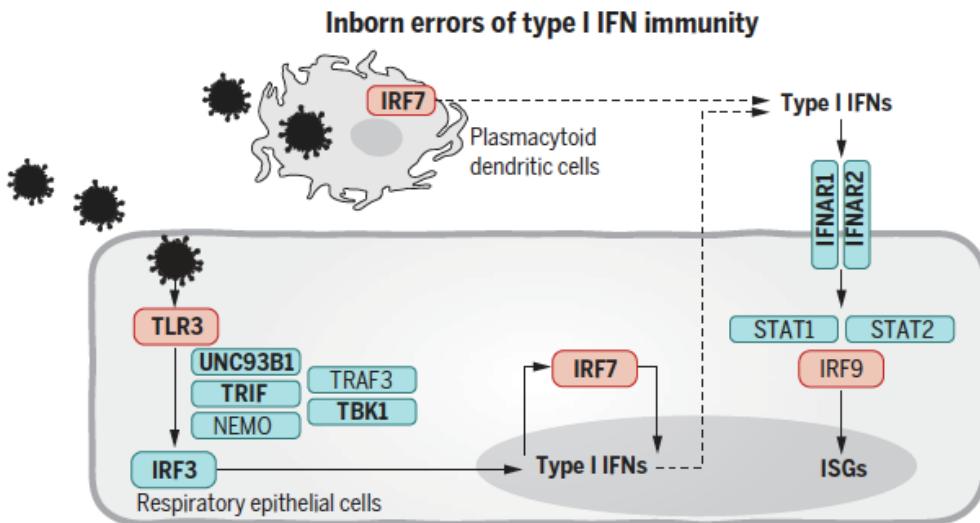
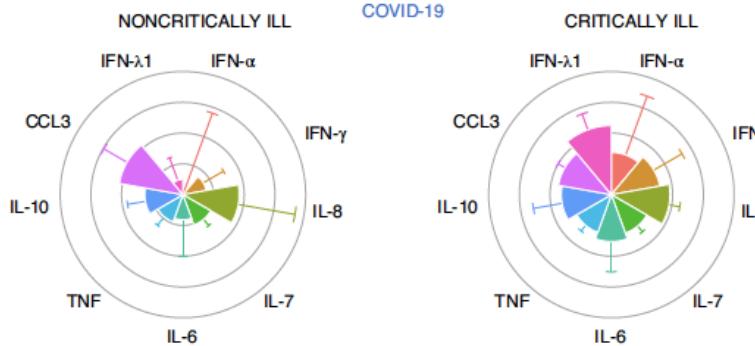
Aicardi-Gutierrez Syndrome



SAVI Syndrome

COPA syndrome

IFNs in COVID19

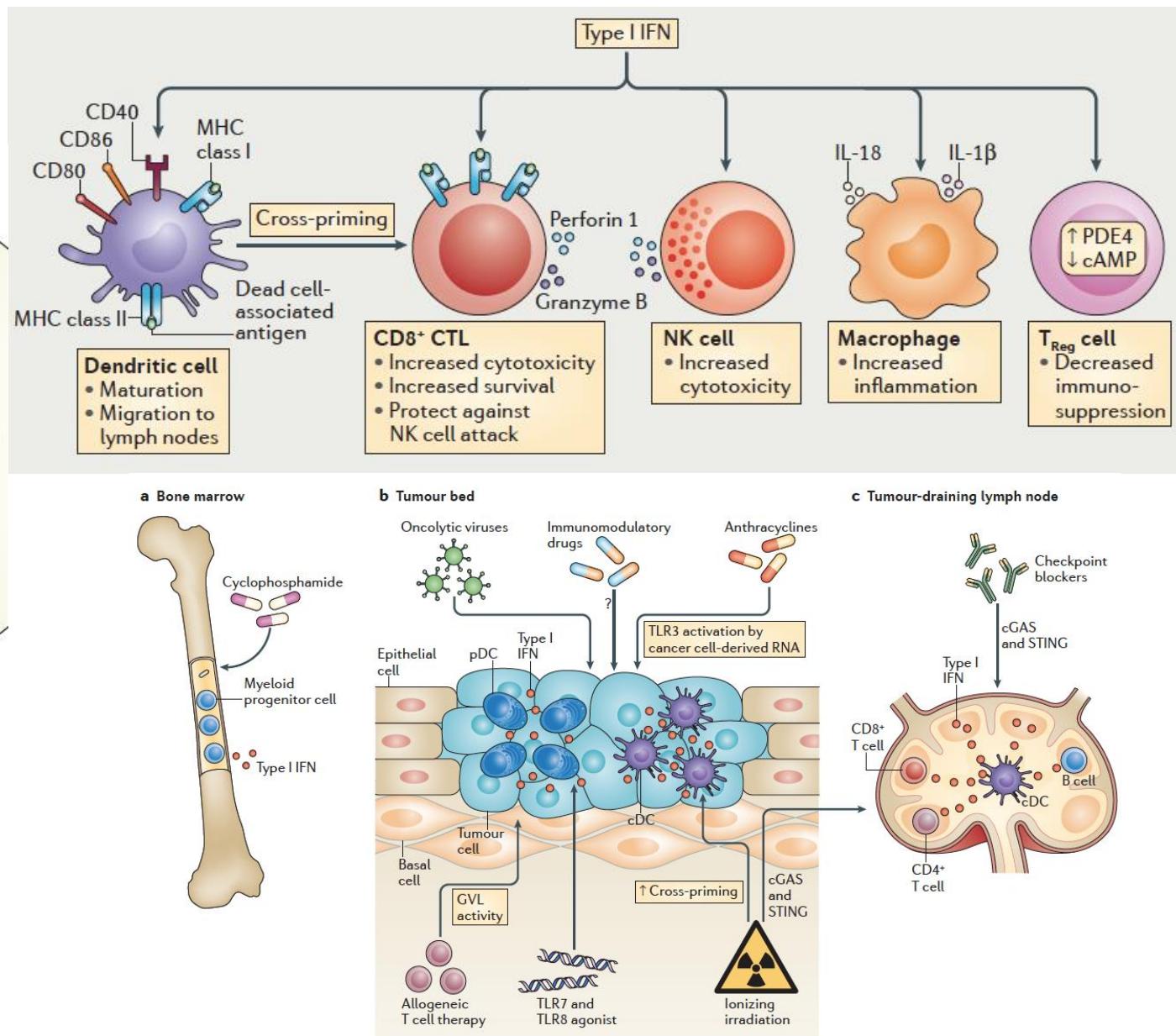
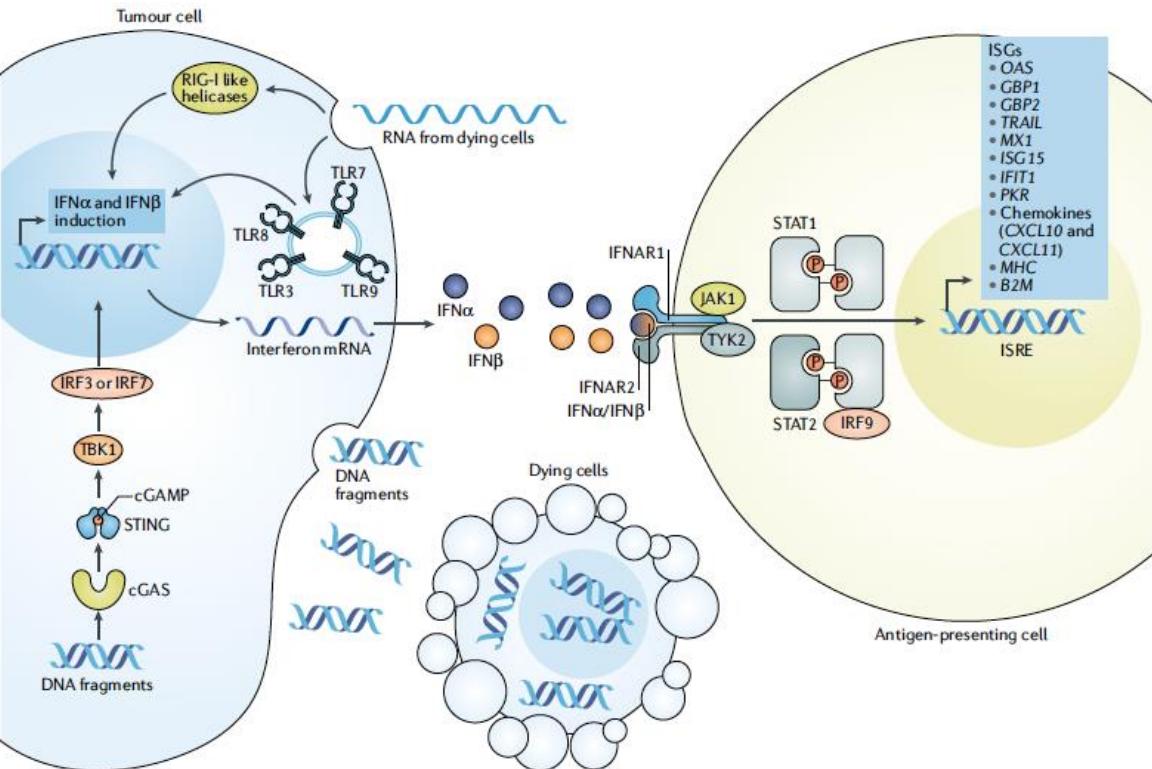


Galani I. et al, Nat Immun 2021

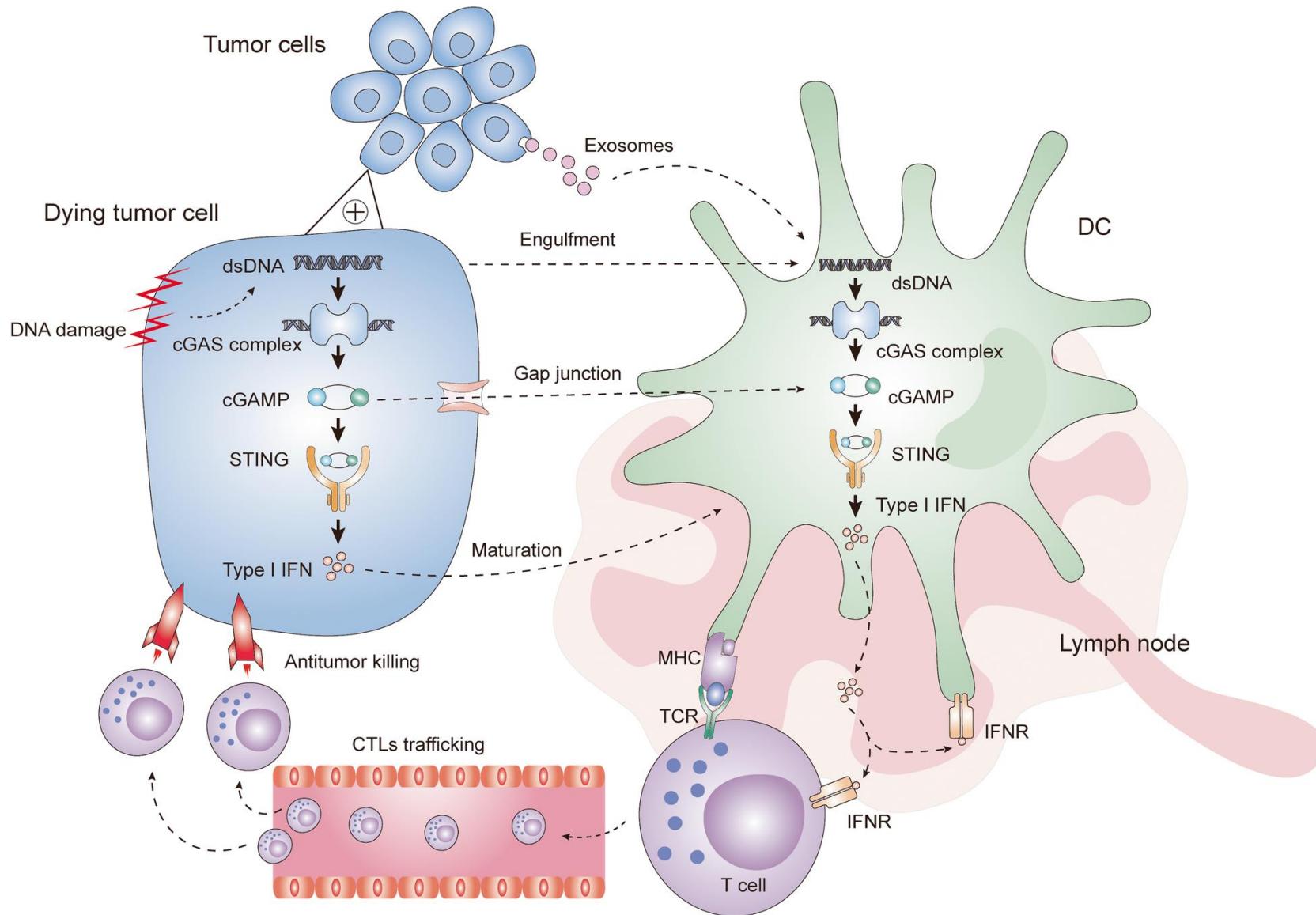
Bastard P. et al, Science Immunology 2022

Zhang Q, et al Science 2020

Interferon impact on cancer biology

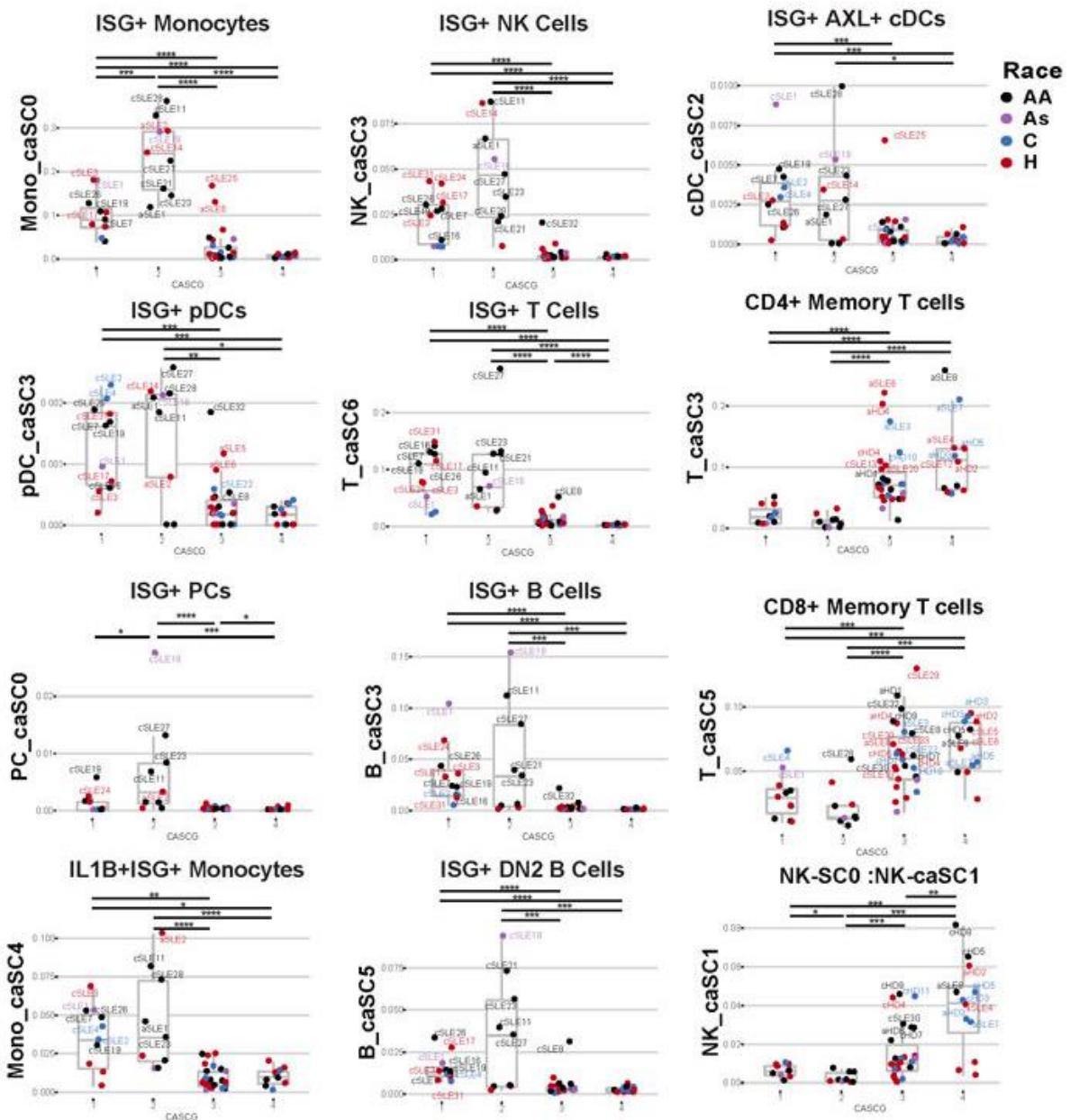
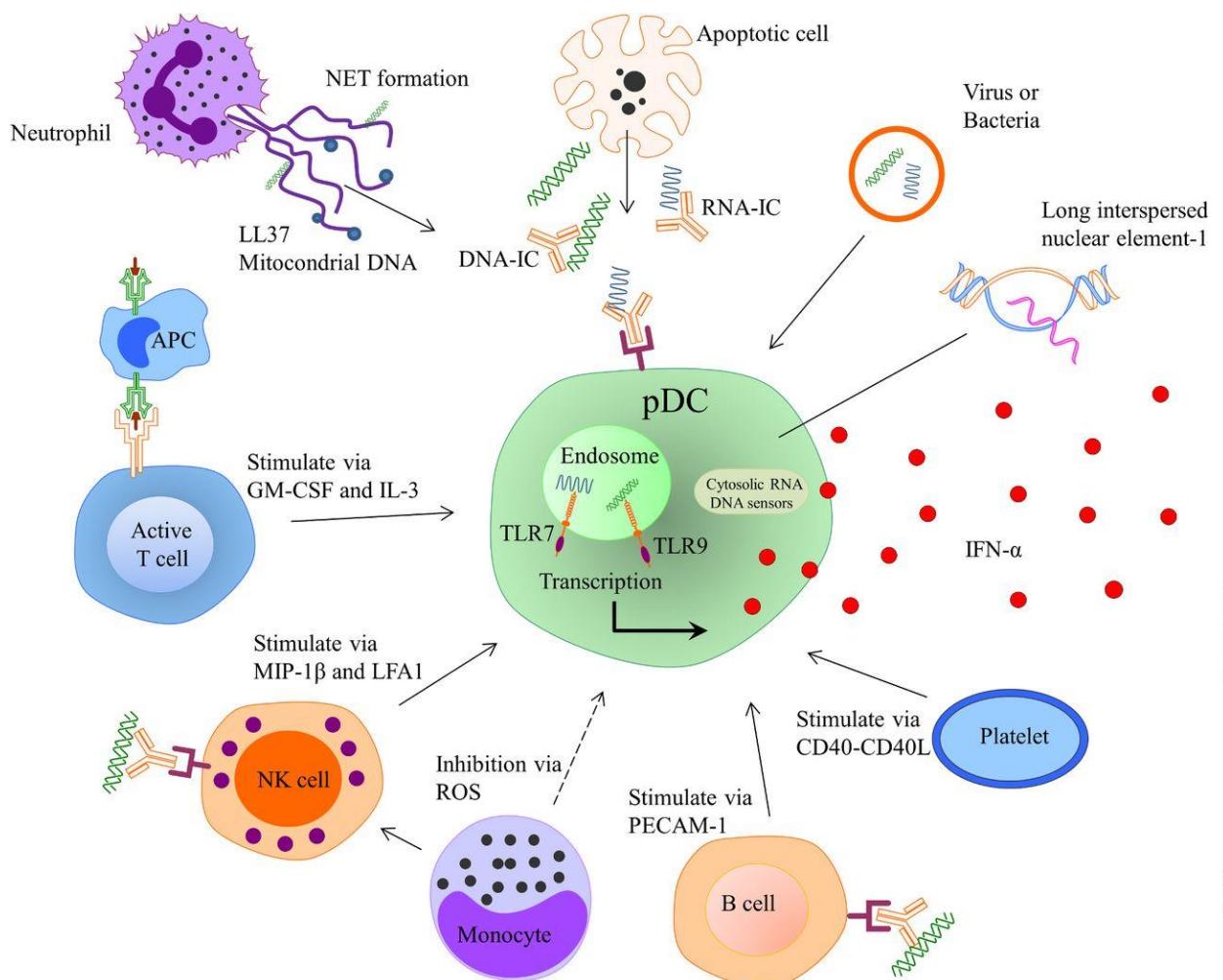


cGAS/STING pathway and Type I IFN in tumor immunity

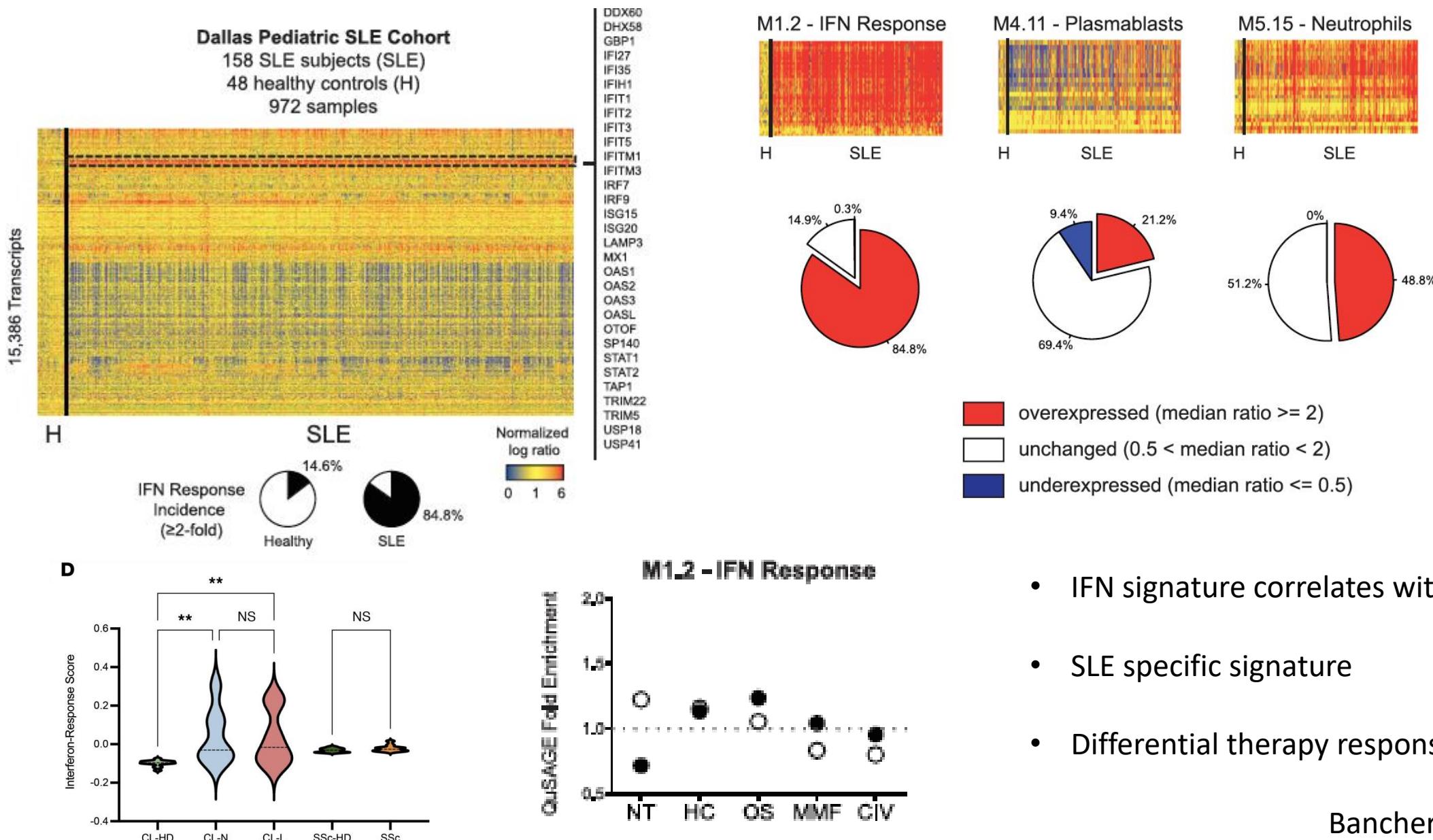


IFN pathway in SLE

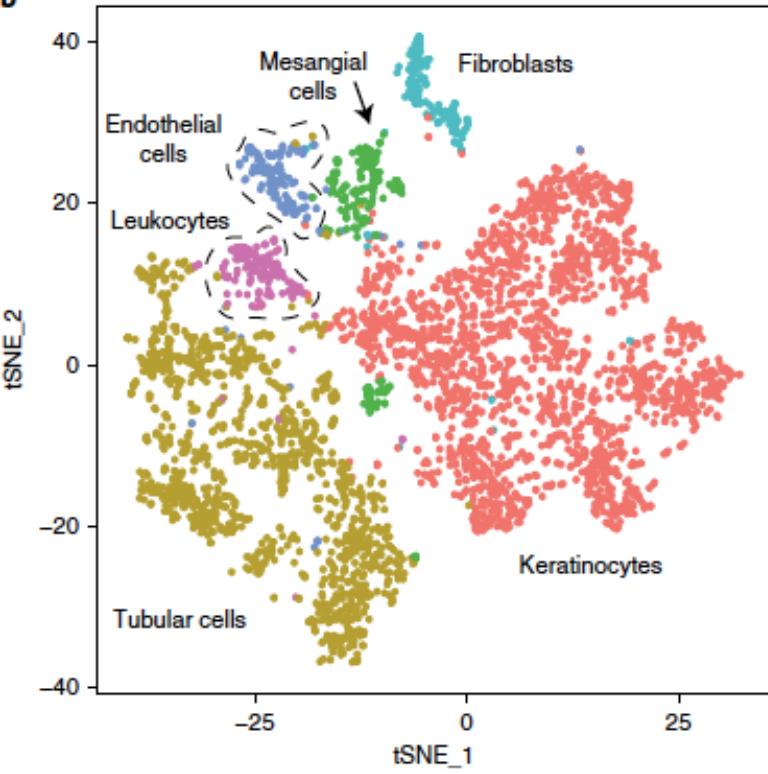
Cell type-specific profile



Molecular Stratification of Lupus Patients

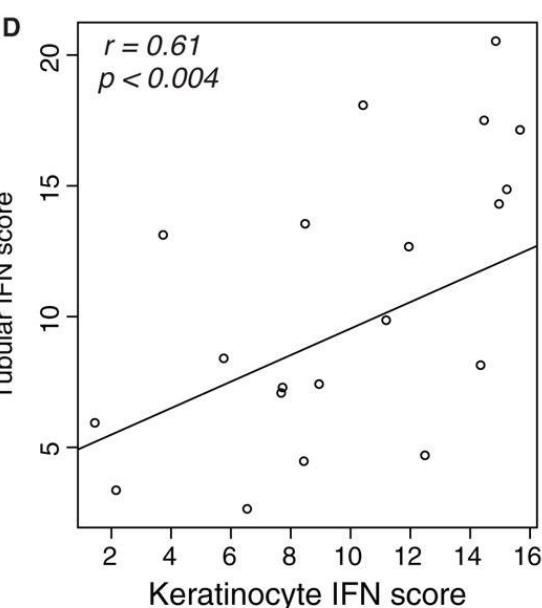
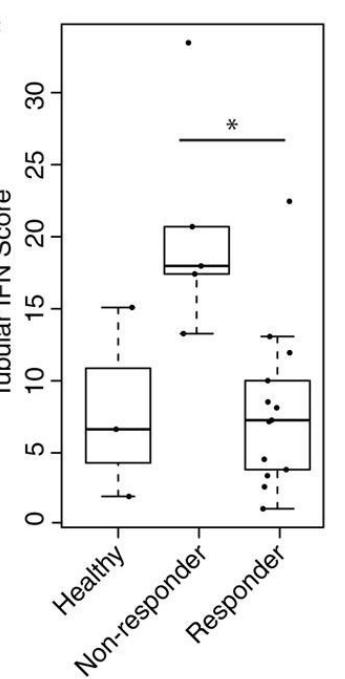
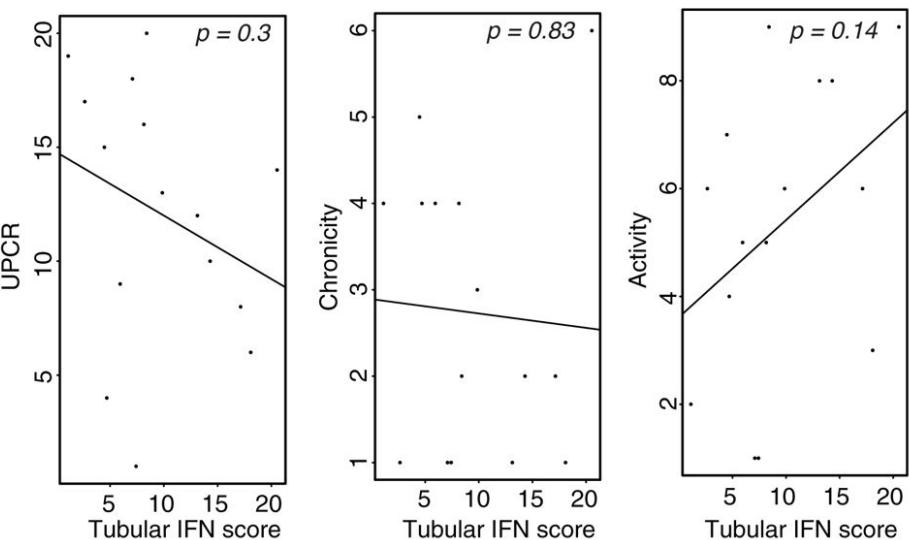
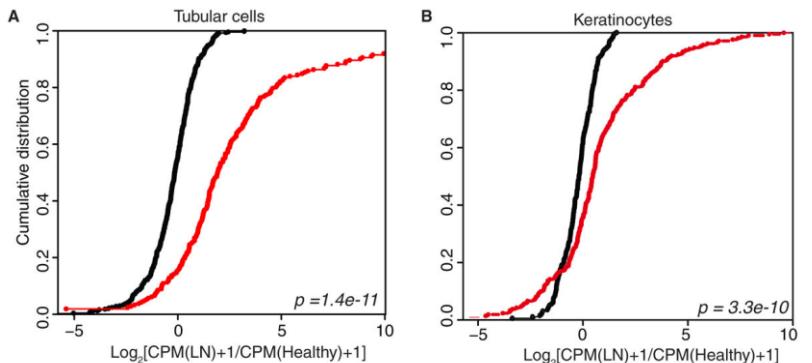


Tubular cell and keratinocyte single-cell transcriptomics applied to lupus nephritis reveal type I IFN and fibrosis relevant pathways

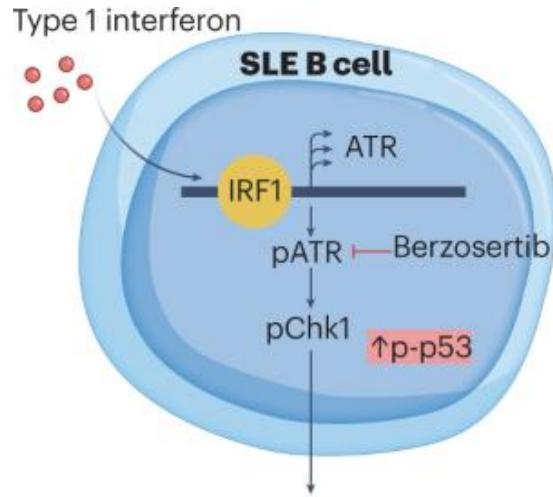
b

Single-cell analysis of the same individual: Skin vs Kidney

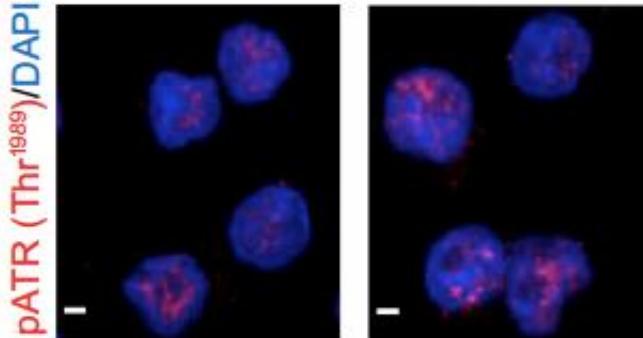
- Higher tubular IFN score in non-responders
- High correlation of keratinocyte vs tubular IFN score
- High correlation of LN damage indices to IFN score



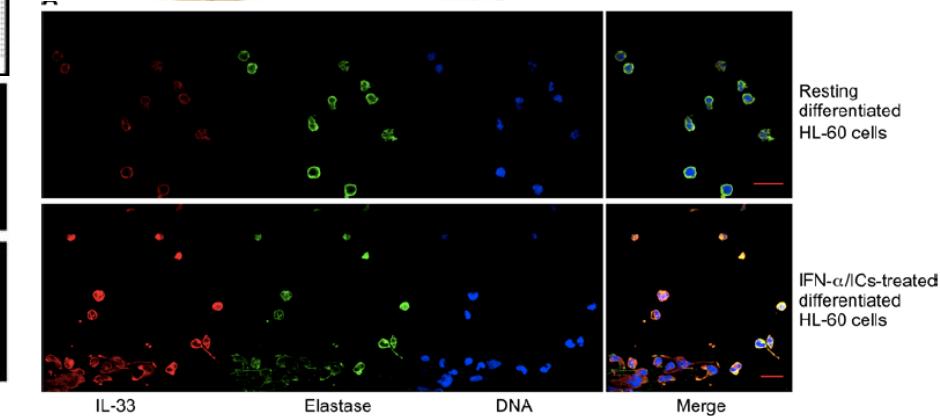
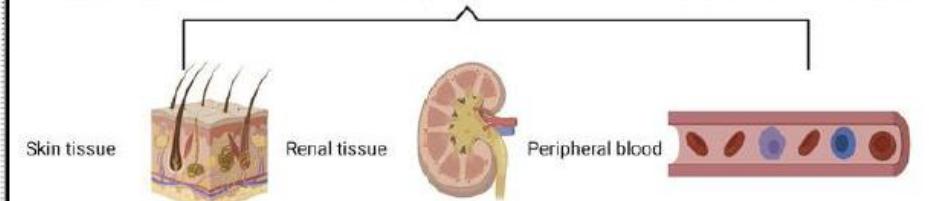
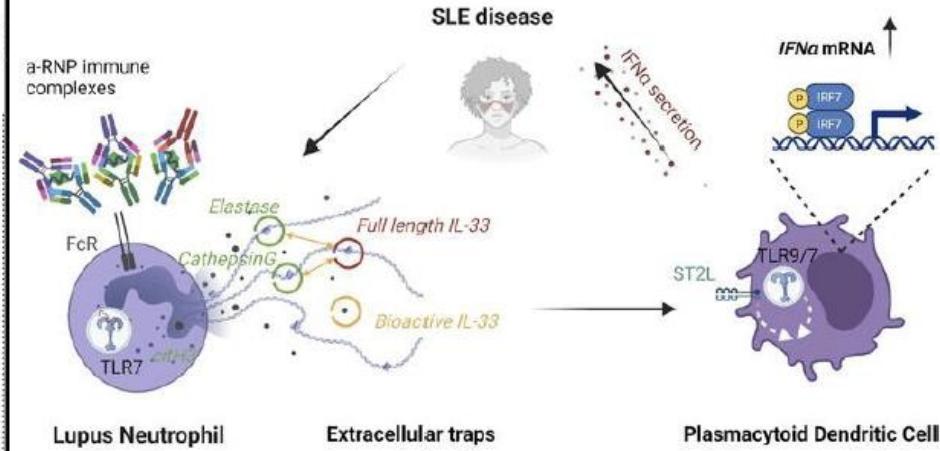
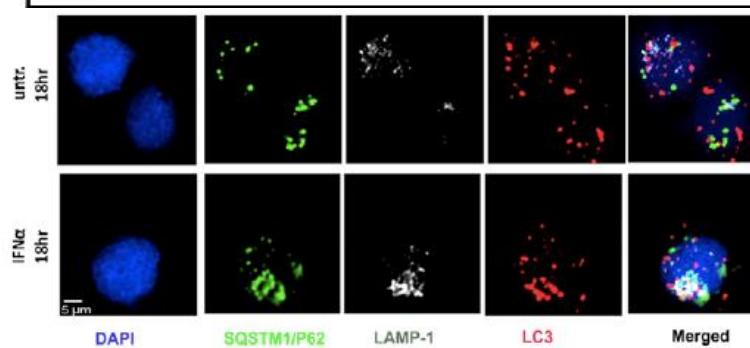
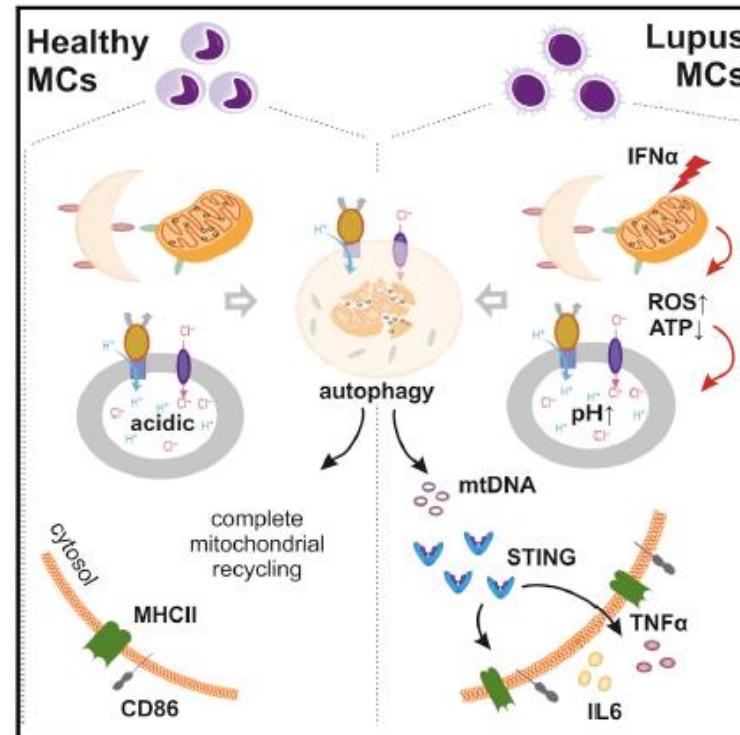
Different mechanisms promote autoimmunity through IFN signaling



IFN- α



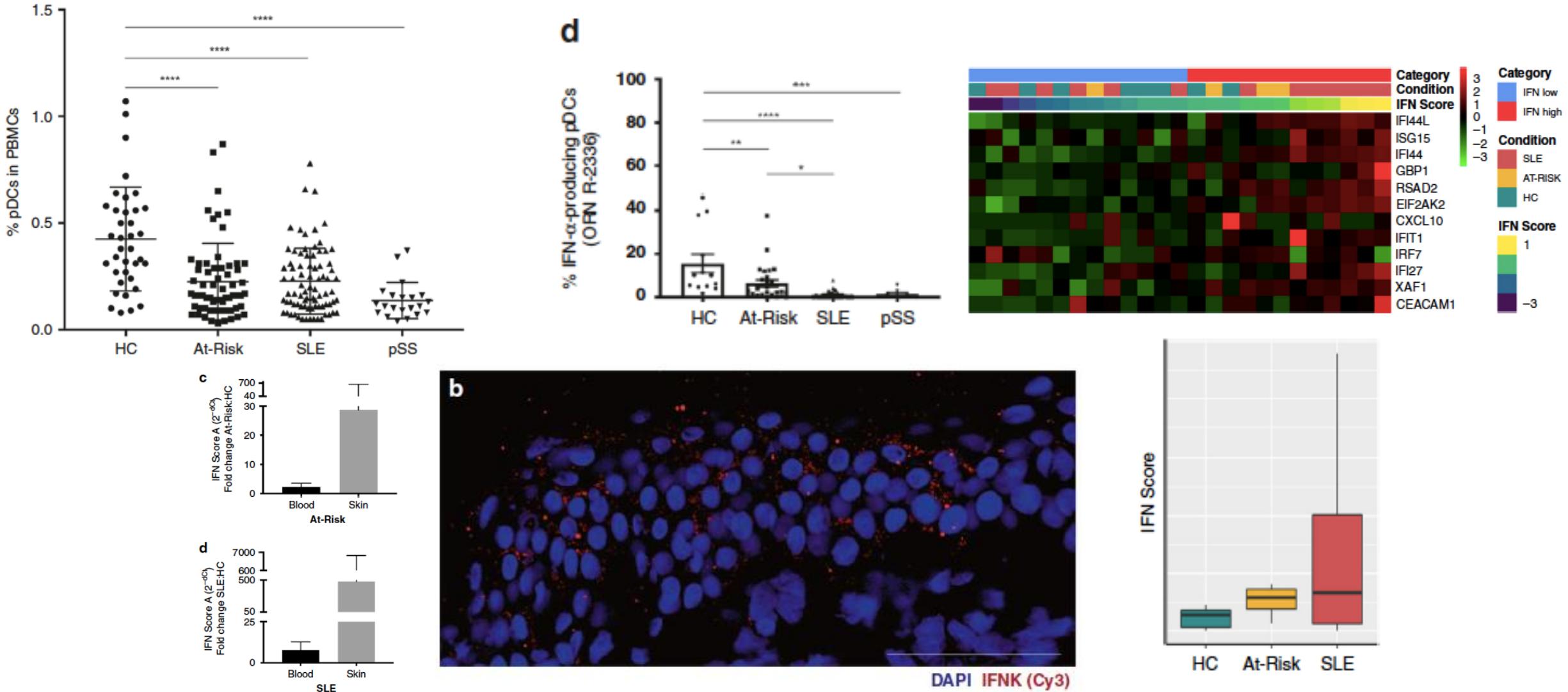
pATR (Thr¹⁸⁸⁹)/DAPI



Functionally impaired plasmacytoid dendritic cells and non-haematopoietic sources of type I interferon characterize human autoimmunity

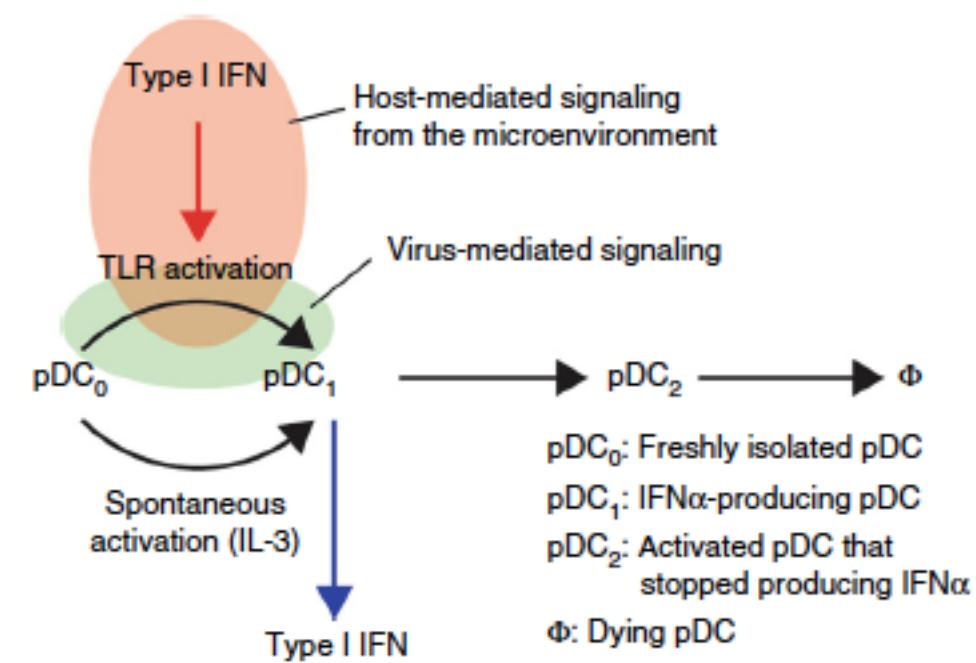
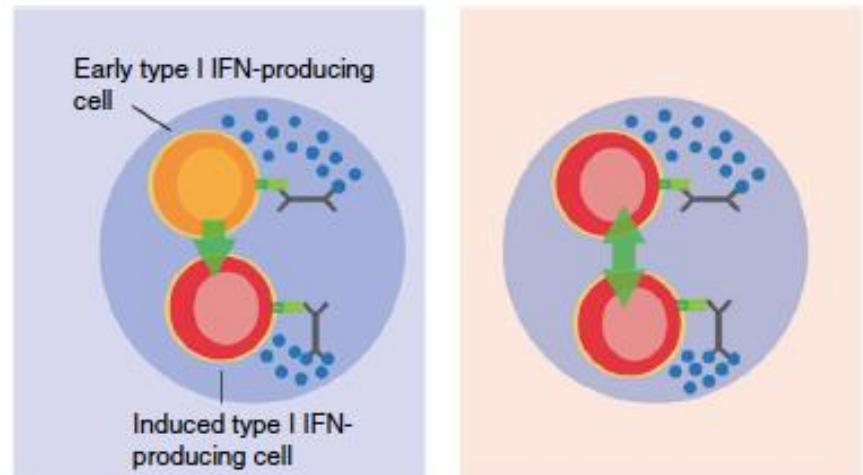
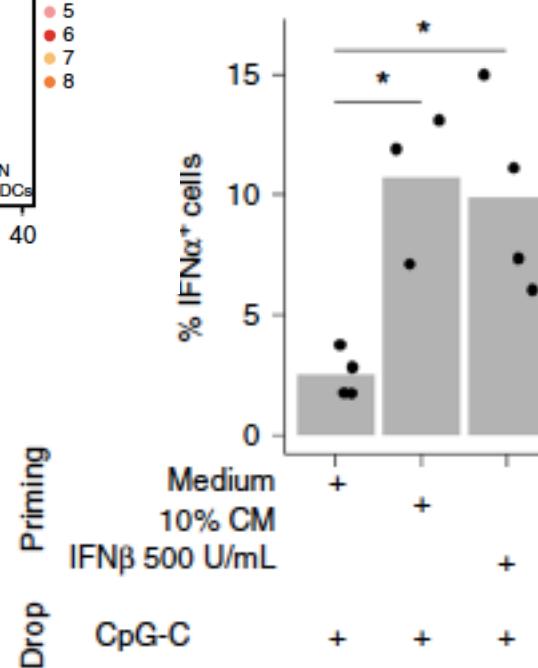
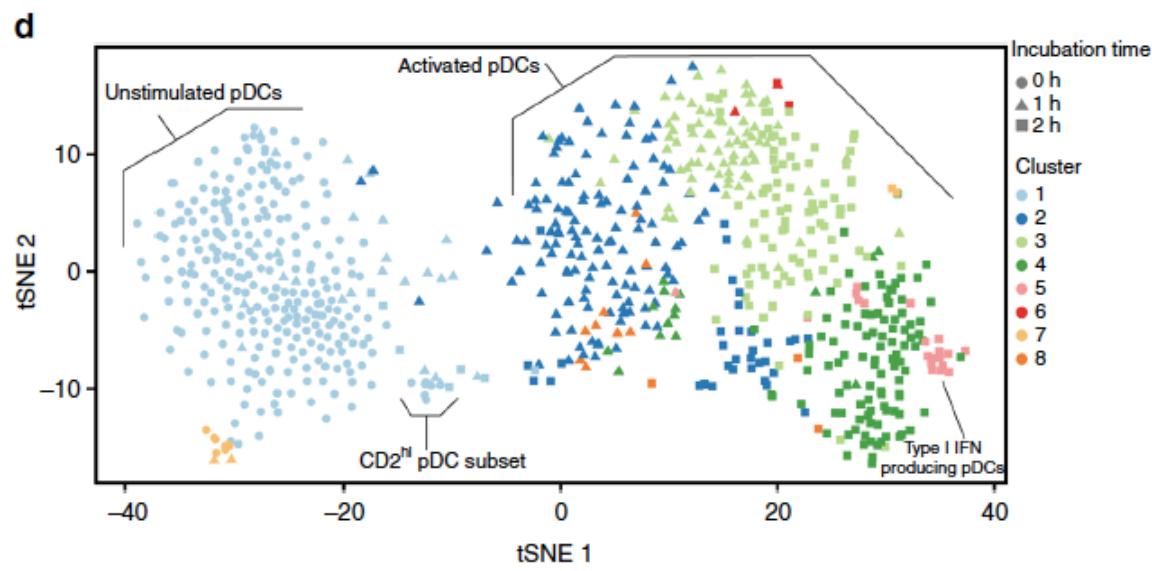
Antonios Psarras^{1,2,3}, Adewonuola Alase¹, Agne Antanaviciute⁴, Ian M. Carr⁴, Md Yuziaful Md Yusof^{1,2}, Miriam Wittmann^{1,2}, Paul Emery^{1,2}, George C. Tsokos^{1,2} & Edward M. Vital^{1,2}✉

Current theories of the cell source of IFNs

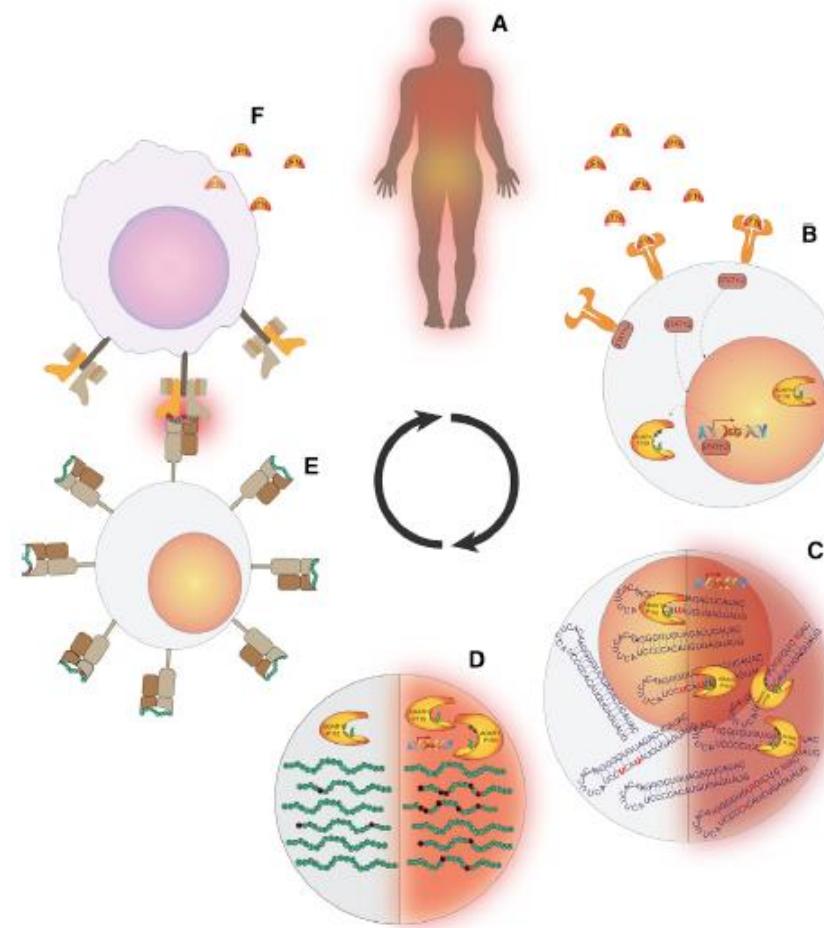
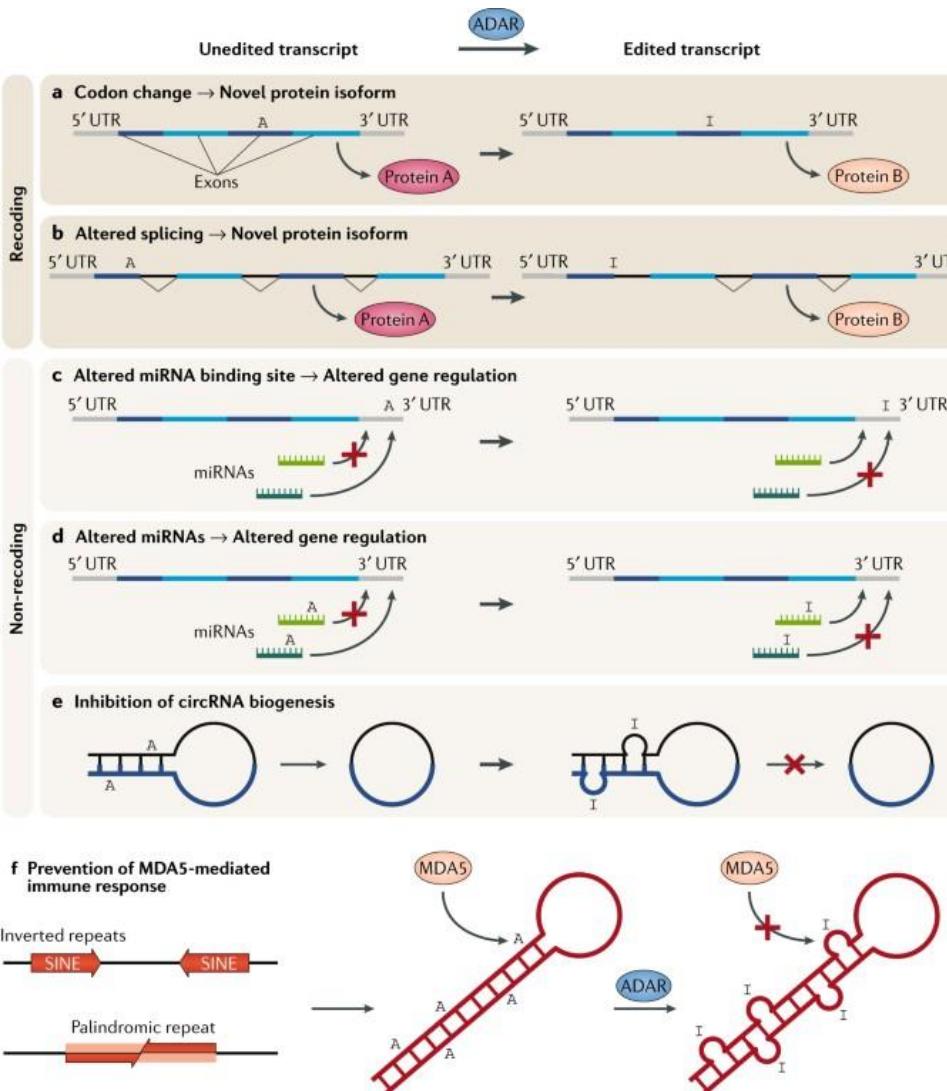


Single-cell analysis reveals that stochasticity and paracrine signaling control interferon-alpha production by plasmacytoid dendritic cells

Florian Wimmers^{1,2}, Nikita Subedi^{3,4}, Nicole van Buuringen¹, Daan Heister¹, Judith Vivié⁵, Inge Beeren-Reinieren¹, Rob Woestenenk⁶, Harry Dolstra⁶, Aigars Piruska⁷, Joannes F.M. Jacobs⁸, Alexander van Oudenaarden⁵, Carl G. Figdor¹, Wilhelm T.S. Huck⁷, I. Jolanda M. de Vries¹ & Jurjen Tel^{1,3,4}



RNA editing (A-to-I) as mechanism of inflammatory response

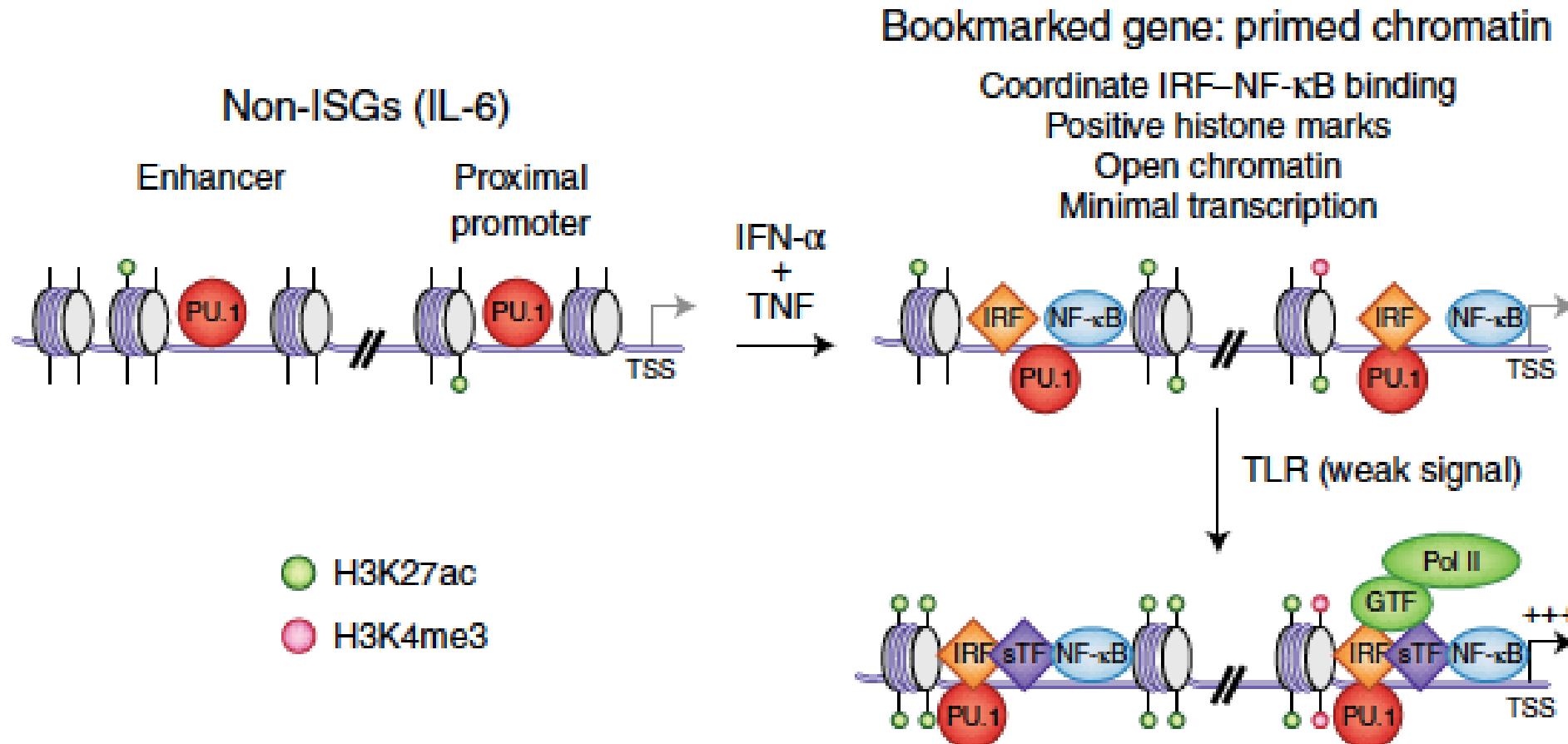


RNA editing in SLE

- Interferon uptake in cells
- JAK-STAT pathway
- ADAR1 upregulation
- Higher editing rates
- More recoded proteins
- Degradation-epitope formation
- Antigen presentation
- Trigger of immune response
- Loop

Eisenberg et al., *Nature Rev Genetics*, 2018
 Roth SH et al., *Cell Reports*, 2018

Epigenetic regulation through type I IFN



2022 EULAR points to consider for the measurement, reporting and application of IFN-I pathway activation assays in clinical research and practice

Javier Rodríguez-Carrio ^{1,2}, Agata Burska, ² Philip G Conaghan ^{1,2}, Willem A Dik, ³
 Robert Biesen ^{1,2}, Maija-Leena Eloranta, ⁵ Giulio Cavalli, ⁶ Marianne Visser, ⁷
 Dimitrios T Boumpas ⁸, George Bertsias, ⁹ Marie Wahren-Herlenius ^{10,11},
 Jan Rehwinkel, ¹² Marie-Louise Frémont ¹³, Mary K Crow ¹⁴,
 Lars Rönnblom ¹⁵, Marjan A Versnel ¹⁵, Edward M Vital ²

Fundamental/basic unmet needs

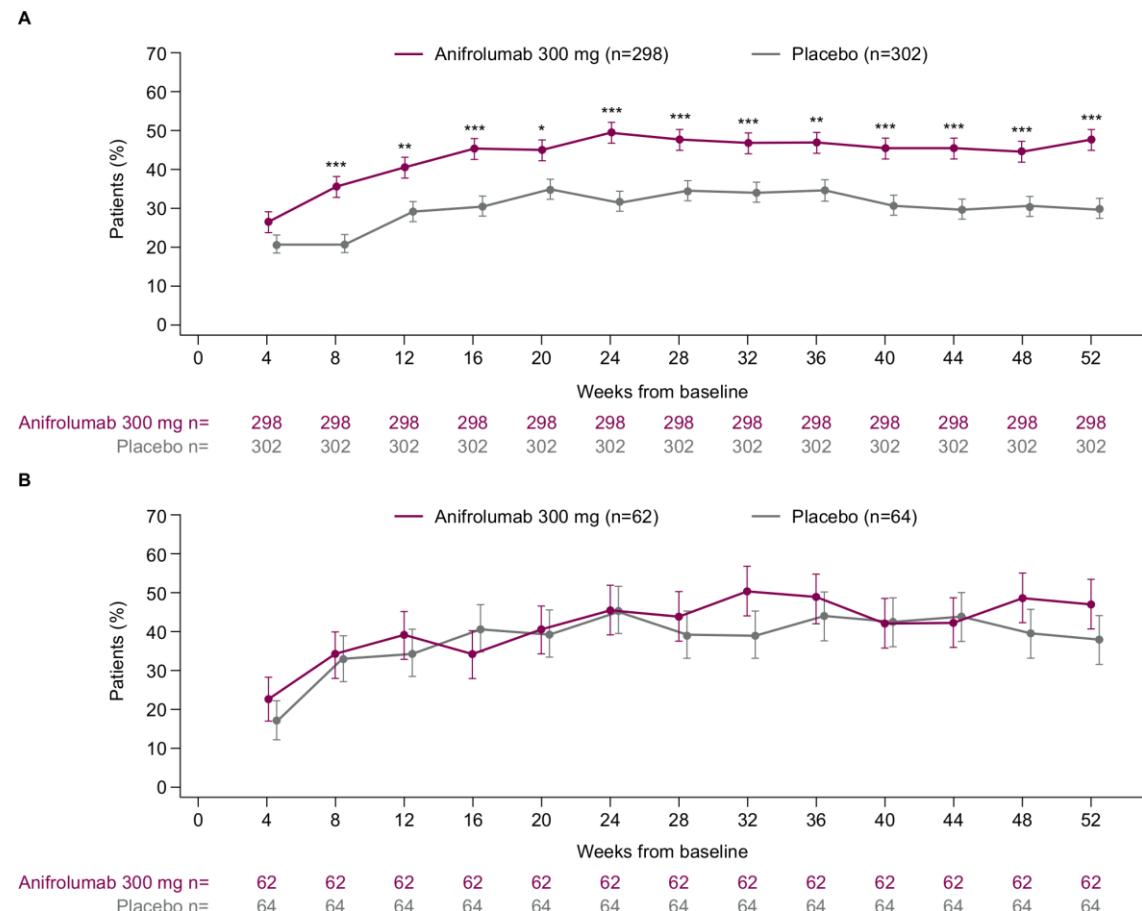
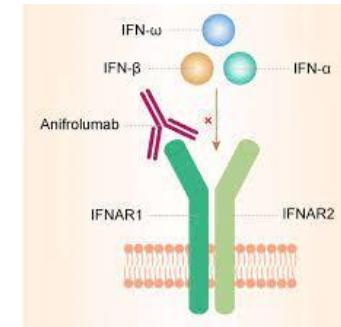
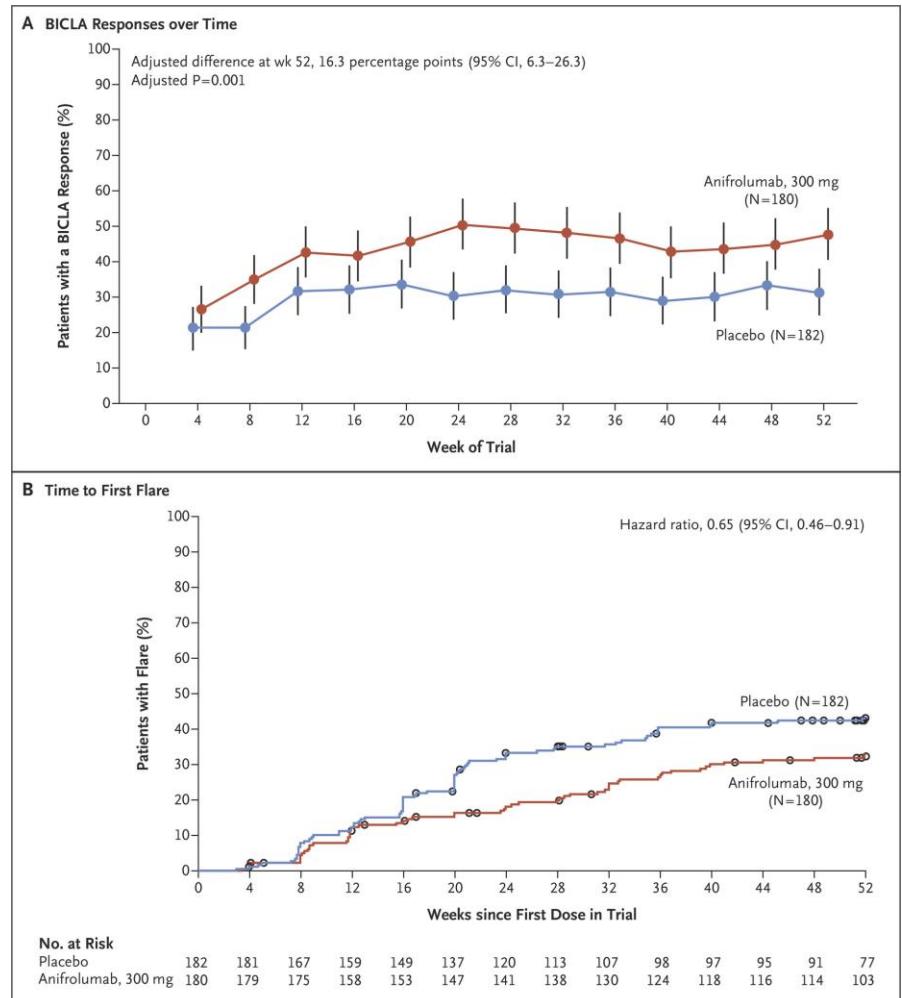
- ⇒ A better understanding of whether different type I interferons (IFN-Is), in particular IFN α s, have unique and/or redundant functions may help in the development of more precise tools for clinical use.
- ⇒ For IFN-stimulated genes:
 - ⇒ Identify the sets of ISGs induced by different IFNs in relevant primary cell types.
 - ⇒ Characterise differences in cell sensitivity to IFN-Is and tissue and cell-specific ISGs profiles.
 - ⇒ Characterise molecular, cellular and biochemical functions of ISGs.
 - ⇒ Identify which of the hundreds of ISGs typically induced actually mediate pathology in rheumatic and musculoskeletal diseases (RMDs).
 - ⇒ Investigate IFN-repressed factors.
- ⇒ Development of assays that directly, sensitively and specifically measure subtypes of IFN-I.

Table 1 Overarching principles and points to consider for the measurement and reporting of IFN-I pathway assays in clinical research and practice

	Level of evidence	Level of agreement (mean±SD, n (%)) scorings ≥8/10
Overarching principles		
A.The IFN pathway is a complex system with multiple subtypes of IFNs and diverse downstream effects on gene and protein expression.	N/A	9.76±0.66 17 (100)
B. IFN-I pathway activation is a common hallmark in many RMDs. Although IFN-I pathway activation is associated with some clinical manifestations, the utility of IFN-I pathway assays in clinical practice requires further validation for most contexts.	N/A	9.29±0.98 16 (94.1)
Points to consider		
1. Task force consensus terminology should be considered for reporting IFN assays measurement.	5	9.58±0.79 17 (100)
2. Existing assays measure different aspects of the IFN-I pathway; they do not reflect the entirety of the pathway and some are not specific for IFN-I. The most appropriate assay will depend on the research or clinical question and should be justified.	4	9.76±0.56 17 (100)
3. Publications on novel IFN-I pathway assays should report whether they specifically reflect IFN-I, and to the extent possible, which IFN-I is measured.	5	9.58±0.61 17 (100)
4. For assays that evaluate pathways downstream of the IFN-I receptor (eg, IFN-stimulated gene expression or protein scores) the choice of components needs to be justified. For gene expression scores, the known subsets of IFN-stimulated genes should be described separately.	5	9.41±0.87 16 (94.1)
5. IFN-I pathway is consistently activated in several RMDs, but assays measuring IFN-I pathway activation cannot be currently recommended for diagnostic purposes.	2b/3b	8.58±1.83 12 (70.5)
6. IFN-I pathway assays define more severe subgroups within many RMDs, so they should be considered in stratification studies.	2b/3b	8.70±1.31 12 (70.5)
7. IFN-I pathway activation is associated with disease activity in some RMDs, especially SLE and myositis, but its added value in clinical decision making is uncertain.	2b/3b	8.82±1.18 14 (82.3)
8. IFN-I pathway assays can predict disease exacerbations, in particular flare occurrence in patients with SLE, but further work should be performed to determine to what extent they outperform current instruments.	2b	9.00±1.00 16 (94.1)
9. IFN-I pathway assays might predict progression from preclinical autoimmunity to clinical disease.	2b	8.00±1.69 11 (64.7)
10. In SLE, IFN-I pathway assays may be useful in predicting response to IFN-I targeting therapies.	2b	8.76±1.20 14 (82.3)
11. IFN-I pathway assay results may be affected by some treatments (eg, IFN-targeted therapies and high-dose glucocorticoids), and timing of sample collection should be taken into account and reported.	2b/3b	9.70±0.46 17 (100)

IFN-I, type I interferon; RMD, rheumatic and musculoskeletal disease; SLE, systematic literature review.

Anifrolumab: IFN targeted therapy in SLE



Vital et al. ARD 2021
Morand et al NEJM 2020

Interferon gene expression signature in rheumatoid arthritis neutrophils correlates with a good response to TNFi therapy

Helen L. Wright¹, Huw B. Thomas¹, Robert J. Moots² and Steven W. Edwards¹

