



Dietary gluten permanently reconfigures gastrointestinal tissue-resident immunity in celiac disease

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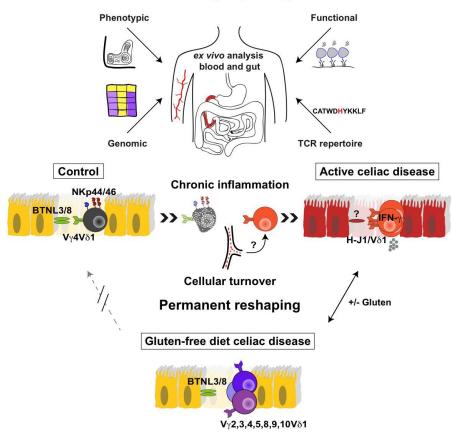
Vaccine Research Center, NIH, USA

The video

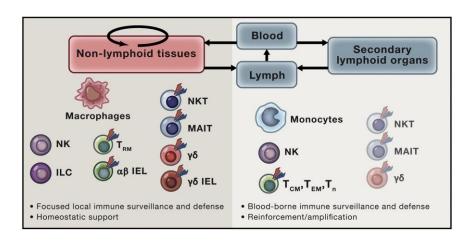
Chronically inflamed then never the same...

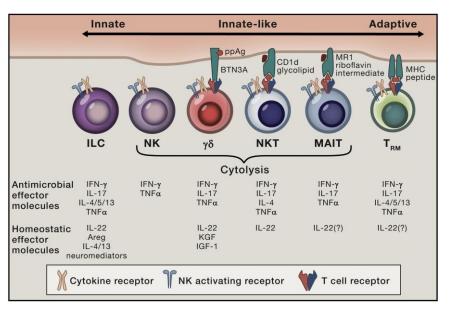
https://www.youtube.com/watch?v=nSkrQgugrc8

Tissue-resident lymphocyte turnover



Tissue-resident lymphocytes





Tissue-resident lymphocytes come in many different flavours (innate through adaptive)

Tissue-resident lymphocytes provide local immune surveillance and homeostatic functions

T_{RM} cells lack the capacity for tissue egress (CD62L⁻ CD69⁺ CD103⁺ S1PR1⁻)

T_{RM} cells proliferate *in situ* to generate tissue-specific immunological memory

T_{RM} cells are "new"

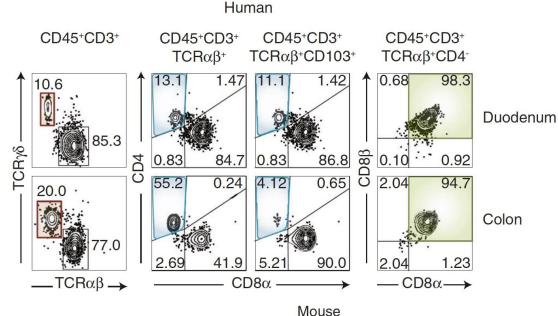
Intraepithelial lymphocytes (IELs) are "old"

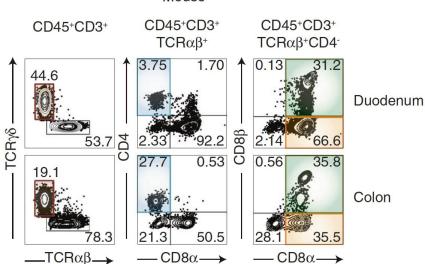
Gastrointestinal IELs

- TCRγδ+T IEL (% of CD3+ cells)
- TCR α β+ CD8 α β+ IEL (% of TCR α β+ CD4- IEL)
- TCR α β+ CD8 α α+ IEL (% of TCR α β+ CD4- IEL)
- TCR α β⁺ CD4⁺ IEL (% of TCR α β⁺ IEL)

The TCR $\gamma\delta$ IEL compartment is established early in life, independently of dietary/microbial insults, and is self-sustaining/stable (in mice)

The TCR $\gamma\delta$ IEL compartment does NOT form in the absence of BTNL molecules (in mice)





TCRγδ IELs in CeD



Active disease is characterized histologically by villous atrophy and immunologically by expanded populations of IELs



Adherence to a GFD leads to resolution of the villous abnormalities and decreased frequencies of CD4+ and CD8+ TCR $\alpha\beta$ IELs, but NOT decreased frequencies of TCR $\gamma\delta$ IELs



Jabri and Sollid, *Nat. Rev. Immunol.* **11**, 1281-1289 (2009) Kutlu *et al.*, *Gut* **34**, 208-214 (1993)



"The persistent increase in $TCR\gamma\delta$ IELs in coeliac disease patients who have recovered a normal mucosa suggests that $TCR\gamma\delta$ IEL do not induce directly epithelial damage."

Kutlu et al., Gut **34**, 208-214 (1993)

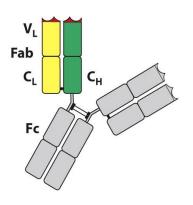


"By whatever means, the end-result of type B IEL activation may again be to suppress tissue infiltration by systemic T cells, as peak type B cell representation seems to correlate inversely with disease symptoms."

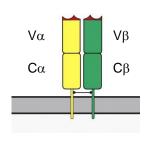


Hayday et al., Nat. Immunol. 2, 997-1003 (2001)

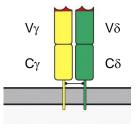
All pure speculation in the absence of defined antigens!!



B cells

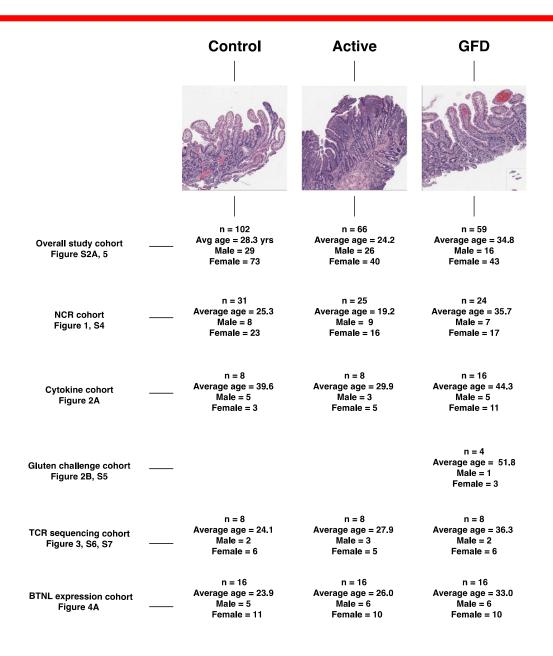


 $\alpha\beta$ T cells

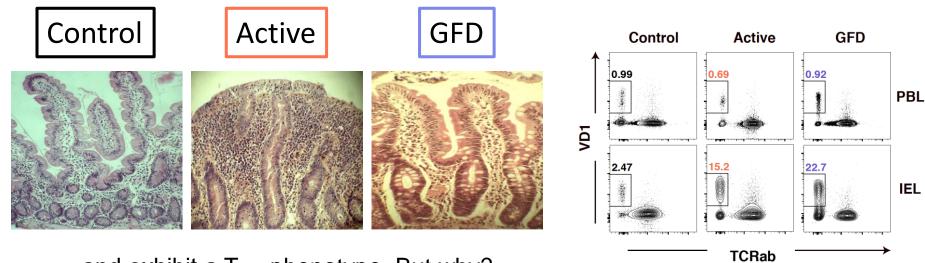


 $\gamma\delta$ T cells

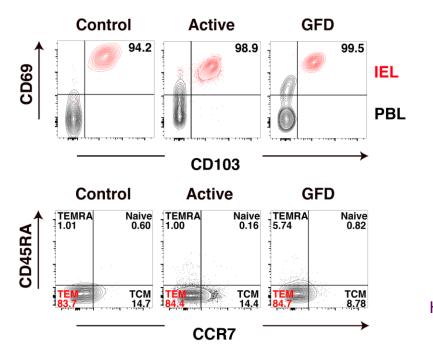
The cohort

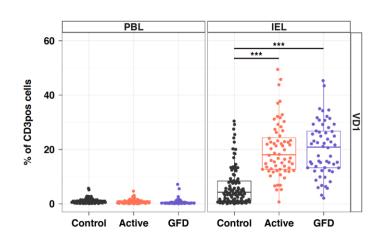


Vδ1+ IELs are expanded in active and GFD-treated CeD



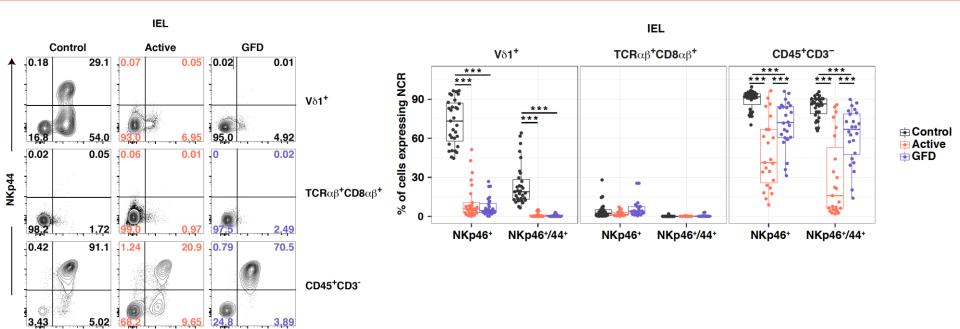
...and exhibit a T_{RM} phenotype. But why?





Halstensen et al., Scand. J. Immunol. 30, 665-672 (1989)

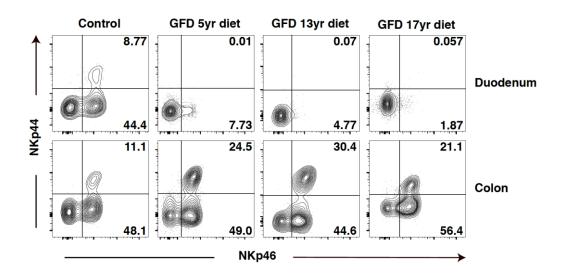
V δ 1+ IELs express activating NCRs in the healthy state but not in CeD



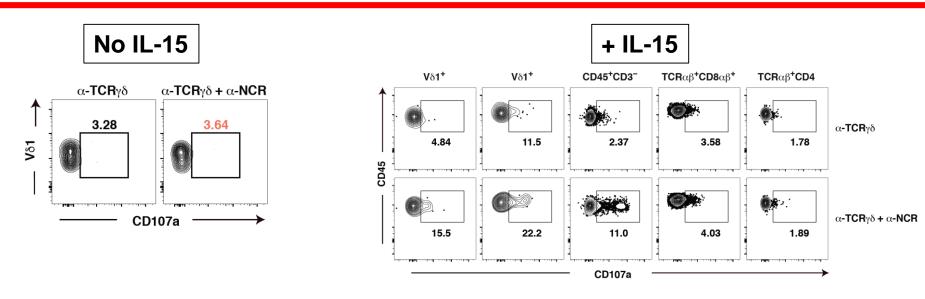
Natural cytotoxicity receptors (NCRs) bind heparan sulphate (HS) and "innately" trigger the cytolytic functions of NK cells

NKp46

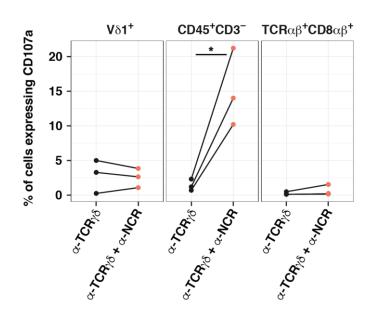
HS has been implicated in the biology of several cancers...

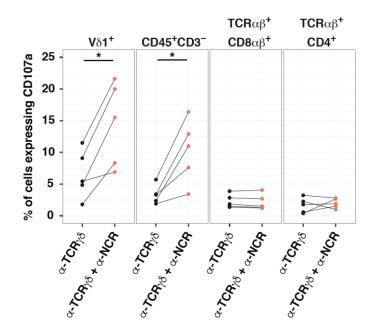


Vδ1+ IELs exhibit weak cytolytic activity in CeD

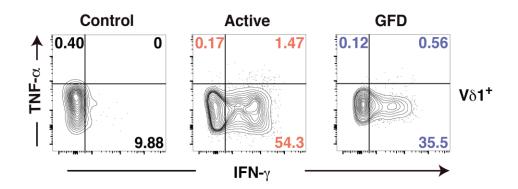


...as measured by surface mobilization of CD107a (LAMP). Enhanced by IL-15 and NCRs.

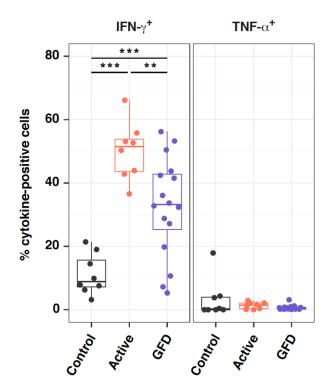




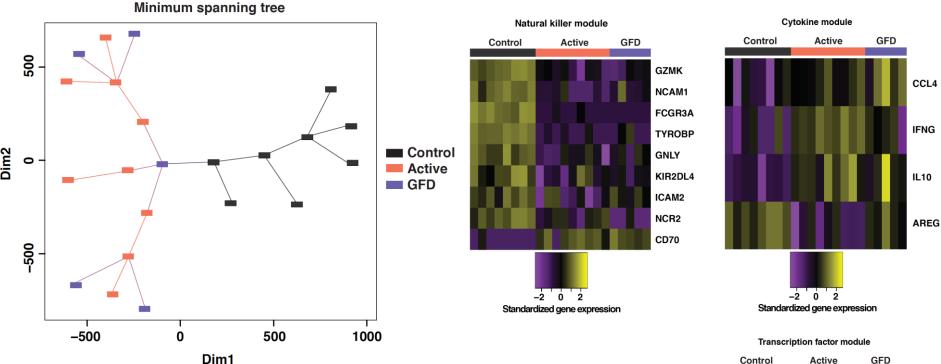
Vδ1+ IELs produce IFN-γ in active and GFD-treated CeD



...in response to PMA/IONO. I.e. $V\delta1^+$ IELs acquire a proinflammatory phenotype in CeD.



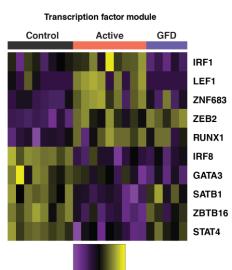
Vδ1+ IELs display a permanently altered transcriptional profile in CeD



NK module – control V δ 1⁺ IELs preferentially express archetypal NK receptor and cytolytic molecules

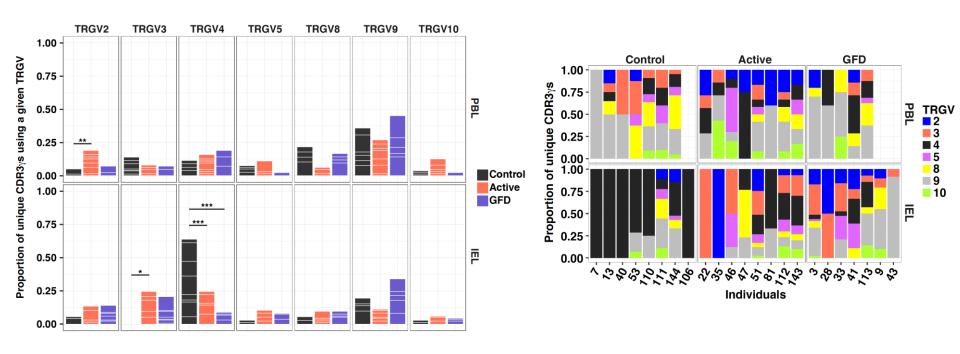
Cytokine module – control V δ 1⁺ IELs preferentially express genes associated with tissue, whereas V δ 1⁺ IELs from patients with CeD preferentially express proinflammatory cytokines

Transcription factor module - it's complicated but consistent!!



Standardized gene expression

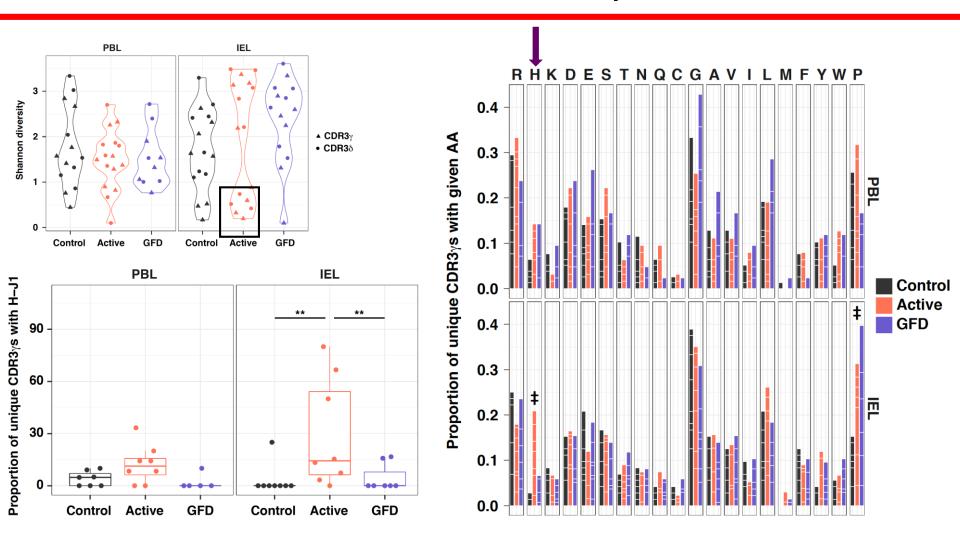
V δ 1+ IELs permanently lose the V γ 4+ "gut signature" in CeD



Vδ1+ IELs preferentially and in some cases exclusively pair with Vγ4+ chains in healthy controls, suggesting germline-driven anatomical localization to the gut

This Vγ4+ "gut signature" is permanently lost in patients with active or GFD-treated CeD, suggesting a coincident loss of an associated tissue-specific ligand for this component of the TCR

$V\delta 1^+$ IELs show evidence of clonal expansions in CeD



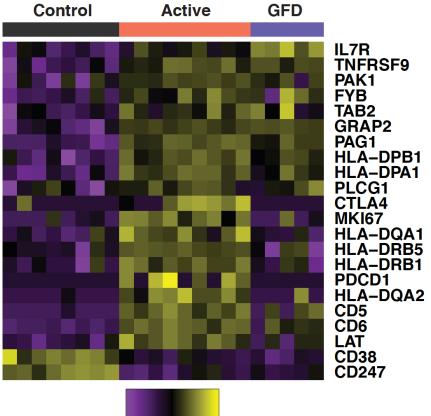
The Vδ1+ IEL repertoire is less diverse in a subset of patients with active CeD

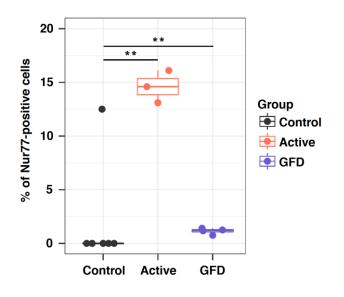
The somatically rearranged CDR3γ loop is enriched for H at position J−1 in patients with active or GFD-treated CeD

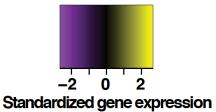
Patterns typically seen in adaptive TCRαβ responses…is there a specific "driver" antigen?

Vδ1+ IELs display hallmarks of TCR-mediated signaling in active CeD

TCR signaling/activation module

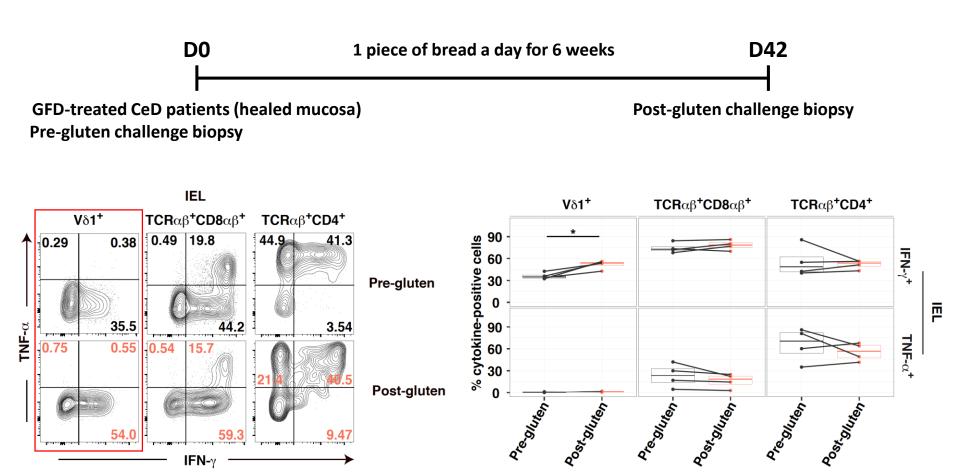






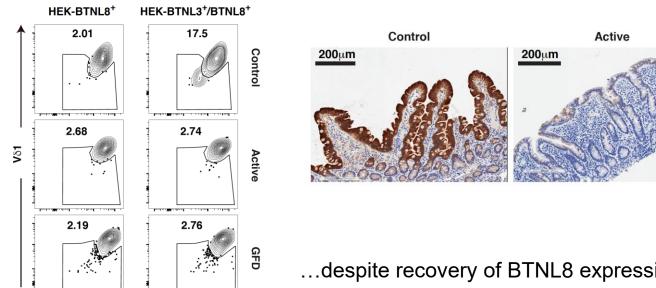
...again consistent with cognate recognition of antigen. But what?

Gluten challenge enhances IFN-γ production by Vδ1+ IELs in GFD-treated CeD



...ligand X? I.e. Specific recognition of a gluten-derived antigen or a self-molecule upregulated in the context of chronic inflammation triggered by gluten exposure?

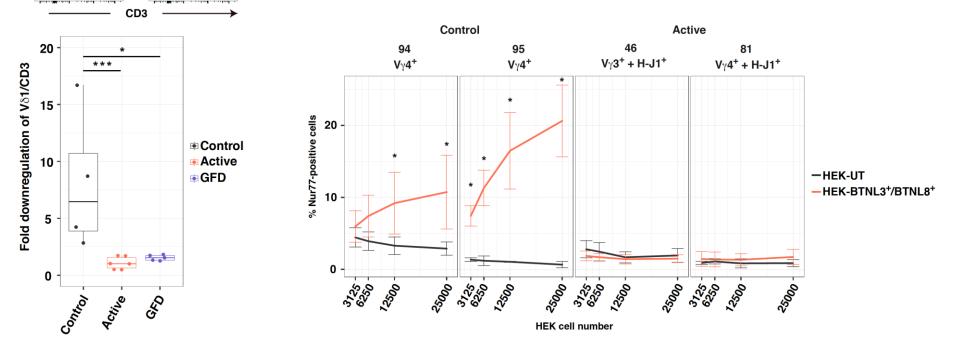
BTNL3/8-reactive V δ 1+ IELs are permanently lost in CeD



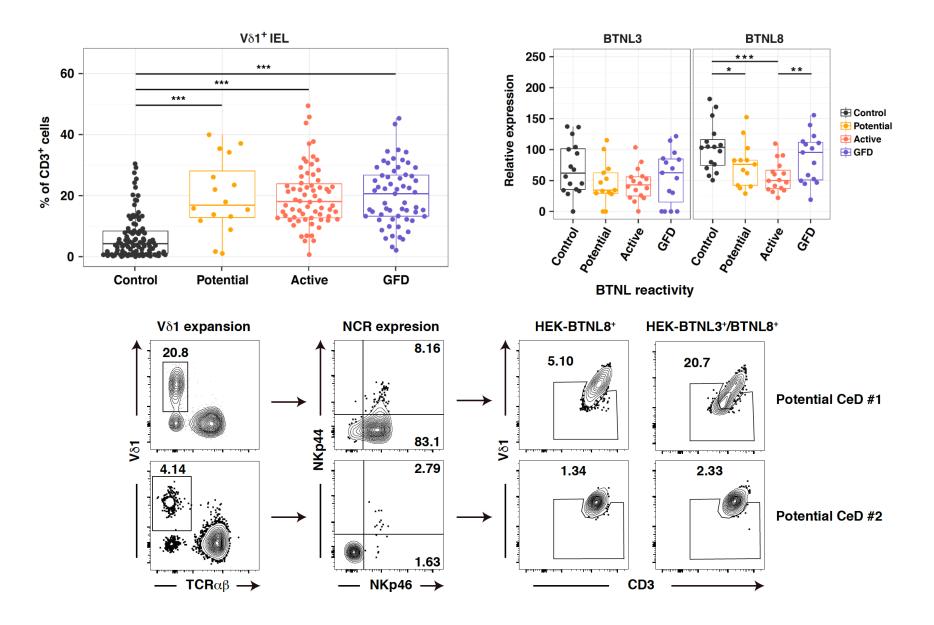


GFD

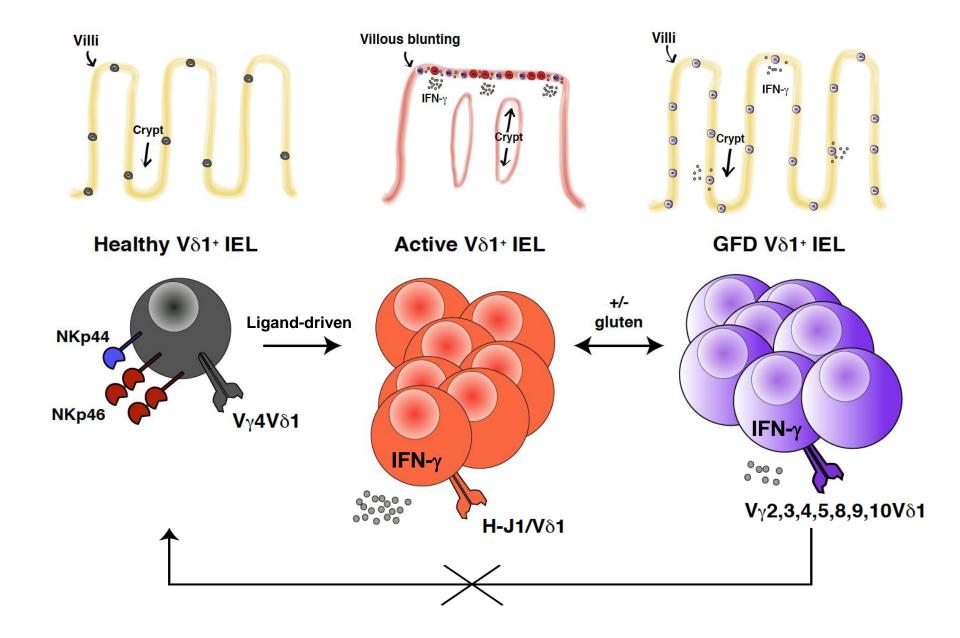
200μm



Dynamic remodeling of the V δ 1+ IEL compartment precedes tissue damage in CeD



Summary



Acknowledgements

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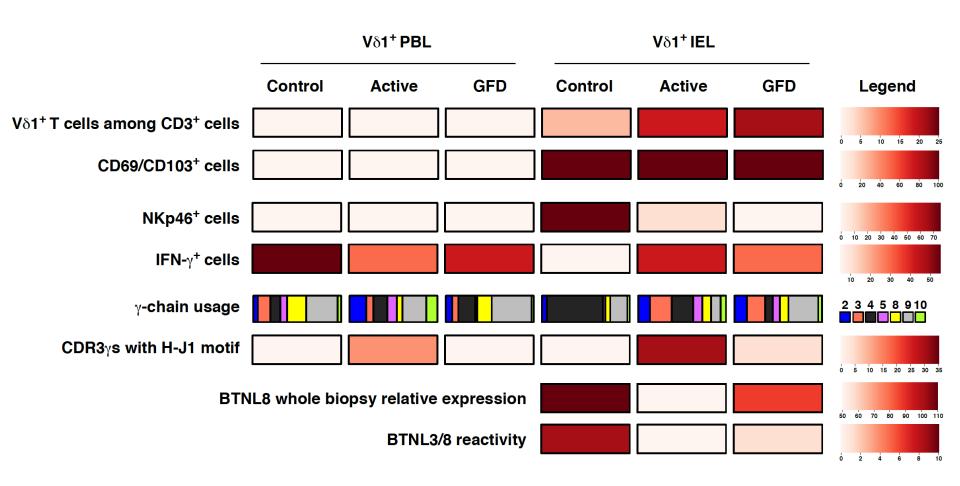
Vinod Kumar Cisca Wijmenga

Leiden * Netherlands

Vincent van Unen

...and Bob!!

Questions?



Above all, I must beware of all ultimate questions - they are too maddeningly unanswerable.

The Journal of a Disappointed Man (W.N.P. Barbellion)