

Genetics of celiac disease

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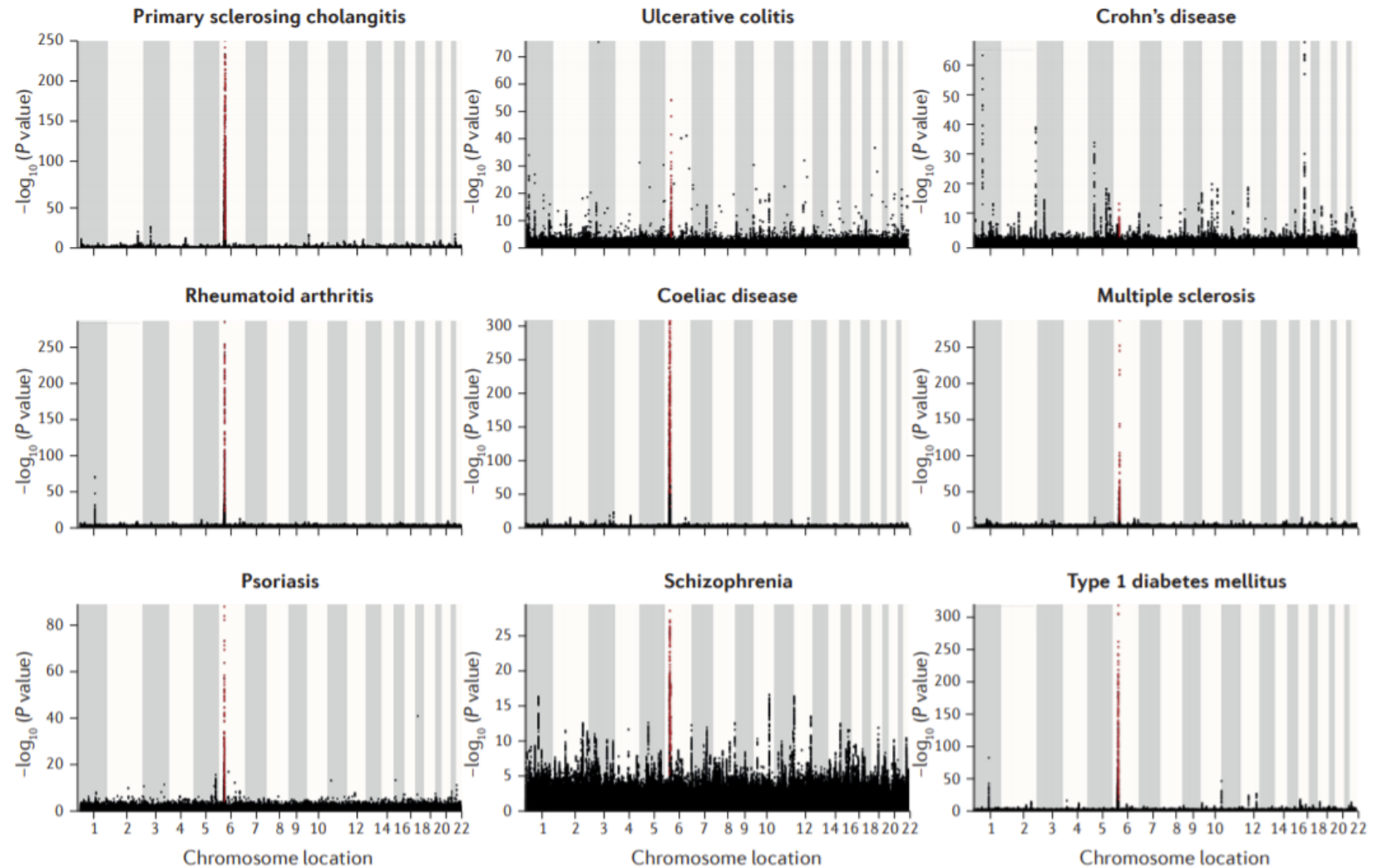
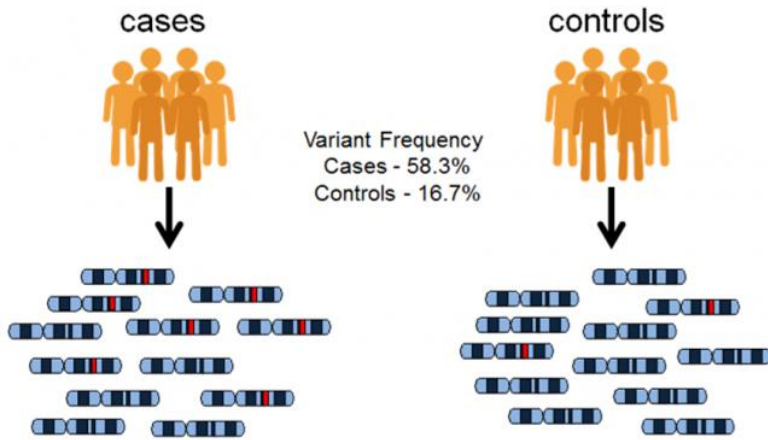
Celiac disease



- Inflammation of the small intestine in genetically susceptible individuals.
- Caused by inappropriate immune response against gluten proteins in food.
- Leads to killing of intestinal epithelial cells, villous atrophy, malabsorption.

HLA and autoimmunity

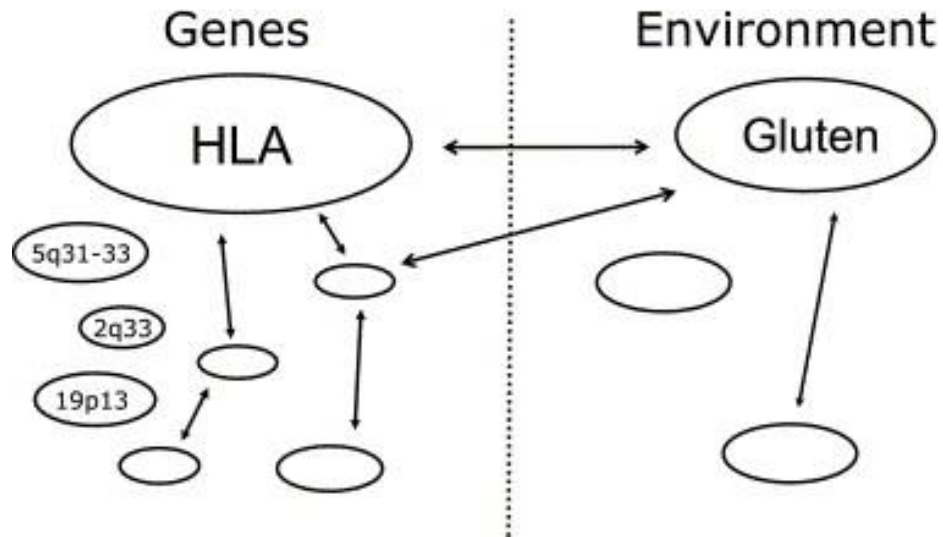
Genome-wide
association studies
(GWAS)



● MHC ● Other

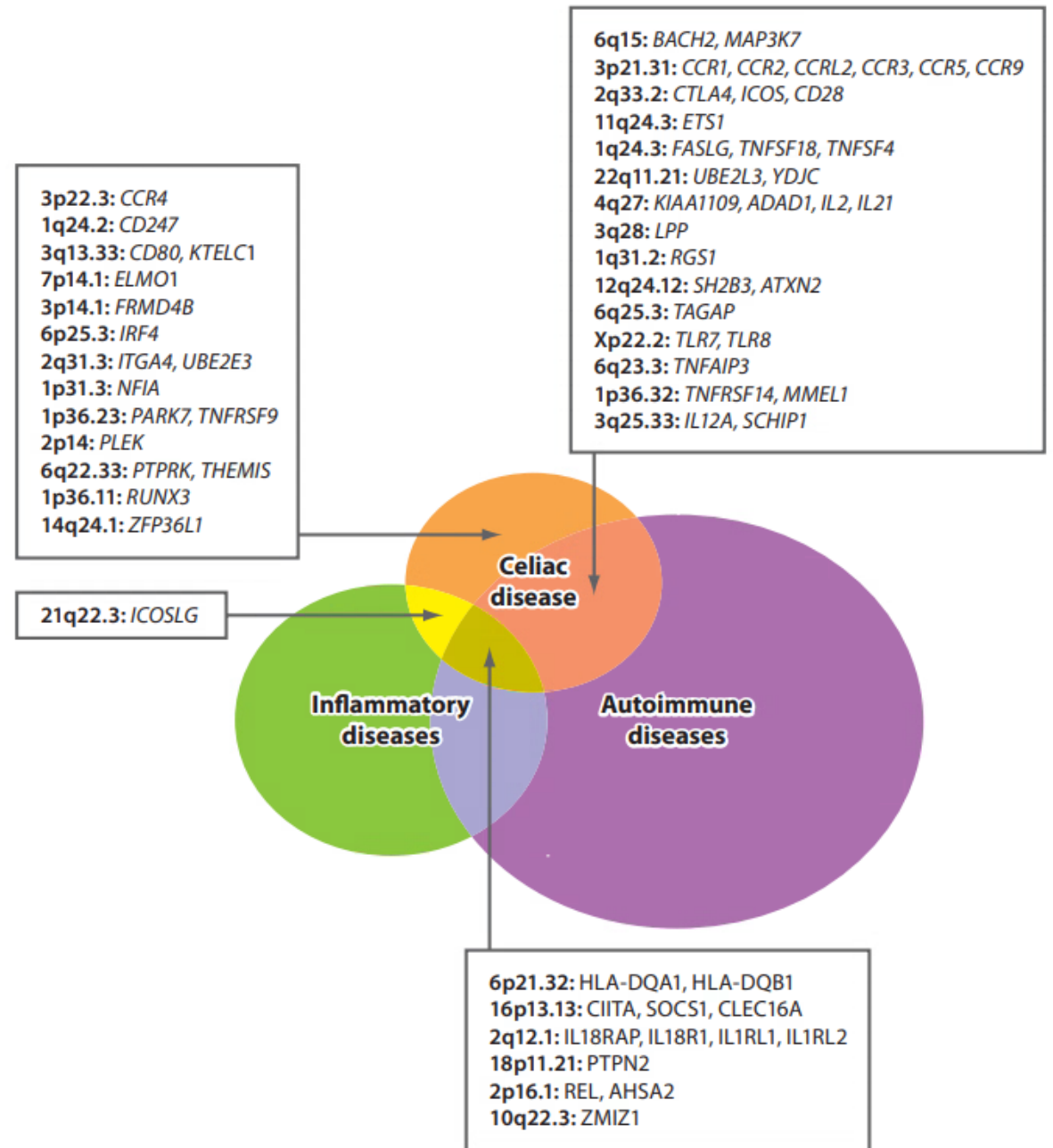
Genetics

- High concordance rate in monozygotic twins, 50-80%.
- Both HLA and non-HLA genes (>40).
- Effect of HLA dominates.
- Missing heritability (~50%).

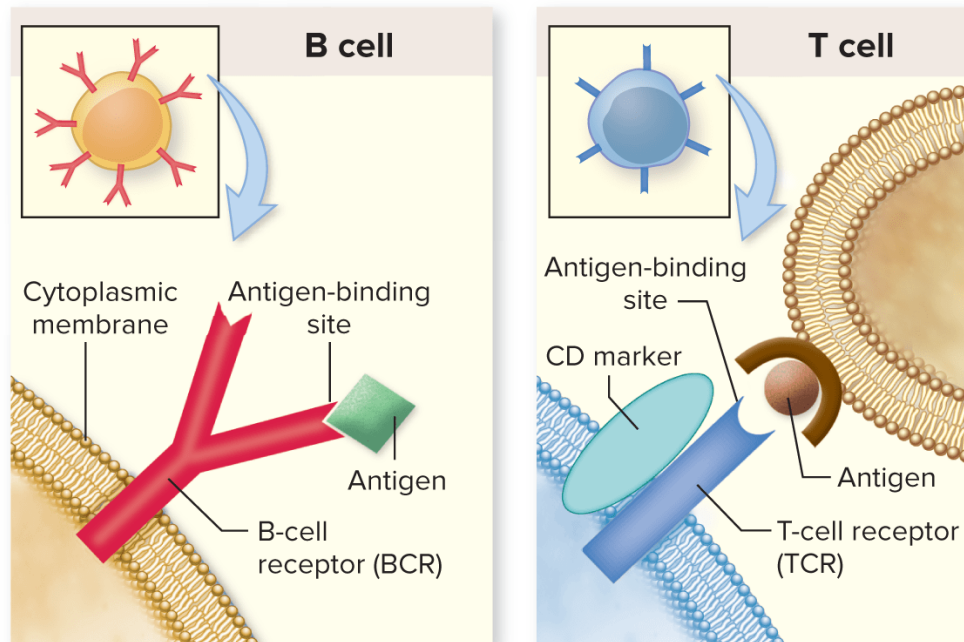


Non-HLA genes

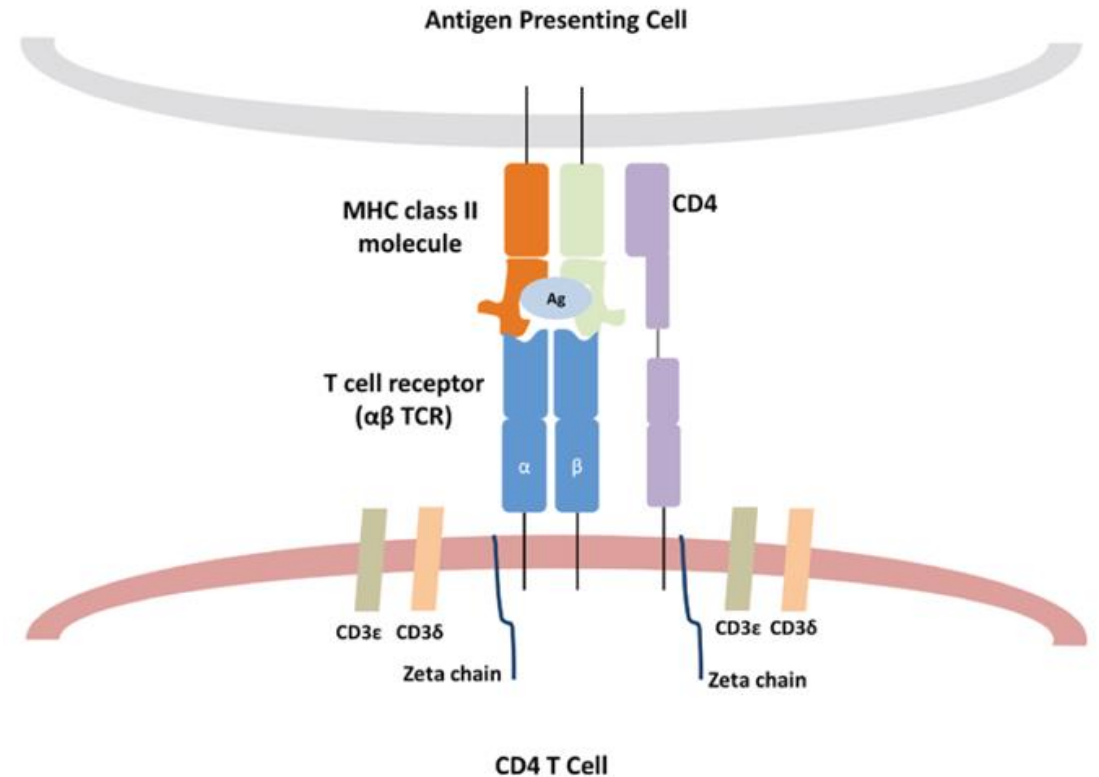
- 95% of SNPs in non-coding regions.
- Genes in associated regions implicated in lymphocyte activation and trafficking.
- Points to an important role of CD4+ T cell activation.



Antigen receptors on B and T cells



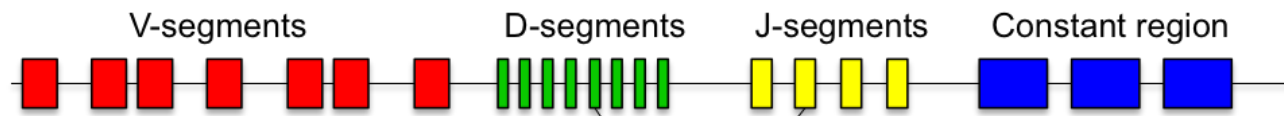
Activation, proliferation



Generation of antigen receptors: V(D)J recombination

- Same mechanism in B and T cells
- Each new cell gets a unique antigen receptor → collection = repertoire
- Theoretically $>10^{15}$ unique receptors

Germline configuration:



(1) D to J recombination



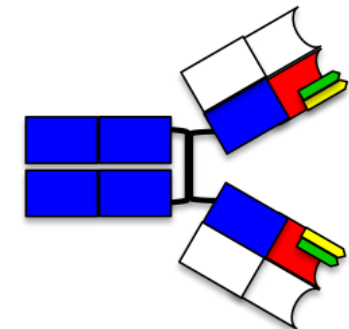
(2) V to DJ recombination



(3) Transcription & splicing

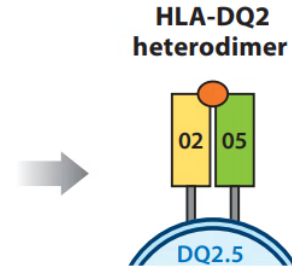
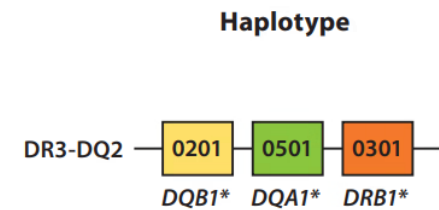


(4) Translation & assembly



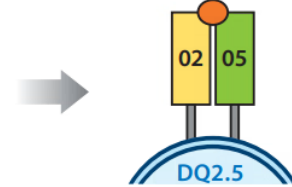
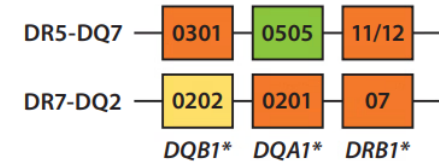
HLA molecules associated with celiac disease

HLA-DQ2

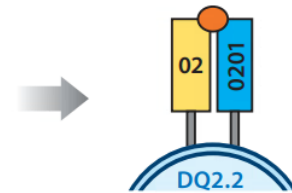


Predisposition for celiac disease

Very high

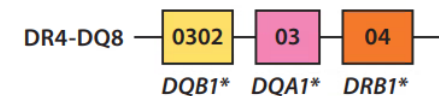


Very high



Low

HLA-DQ8



High

Epitope*

Previous names

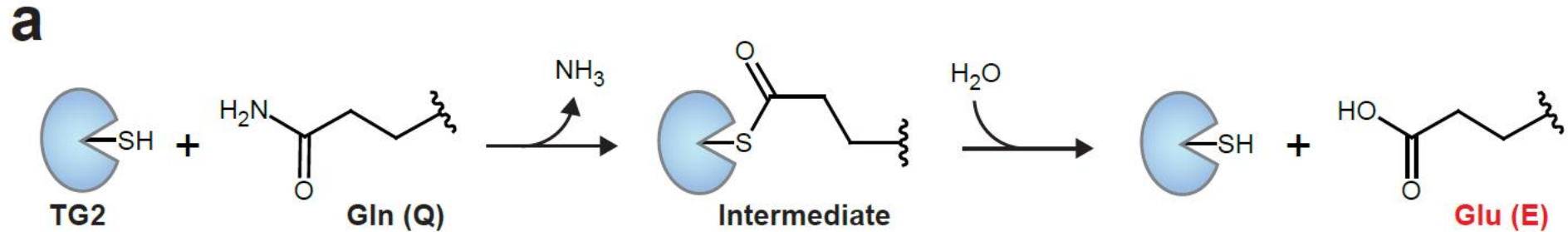
Peptide-binding register†

DQ2.5 restricted epitopes

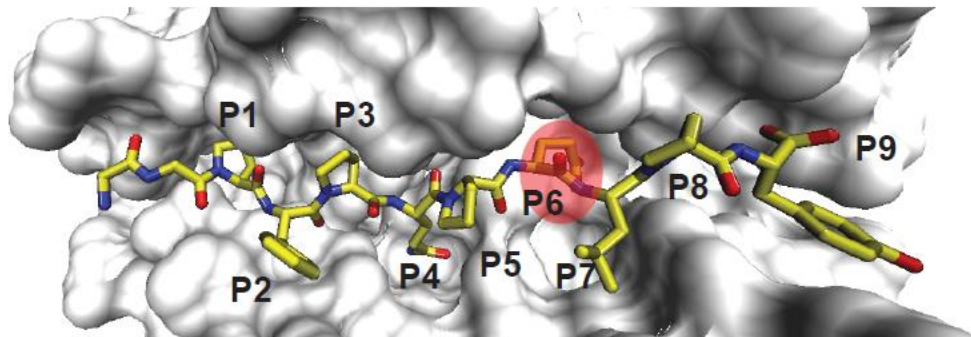
Epitope*	Previous names	Peptide-binding register†								
		1	2	3	4	5	6	7	8	9
DQ2.5-glia- α 1a	DQ2- α -I, α 9	P	F	P	Q	P	E	L	P	Y ^{††}
DQ2.5-glia- α 1b	DQ2- α -III	P	Y	P	Q	P	E	L	P	Y
DQ2.5-glia- α 2	DQ2- α -II, α 2	P	Q	P	E	L	P	Y	P	Q
DQ2.5-glia- α 3	glia- α 20	F	R	P	E	Q	P	Y	P	Q
DQ2.5-glia- γ 1	DQ2- γ -I	P	Q	Q	S	F	P	E	Q	Q
DQ2.5-glia- γ 2	DQ2- γ -II, γ 30	I	Q	P	E	Q	P	A	Q	L
DQ2.5-glia- γ 3	DQ2- γ -III	Q	Q	P	E	Q	P	Y	P	Q
DQ2.5-glia- γ 4a	DQ2- γ -IV	S	Q	P	E	Q	E	F	P	Q
DQ2.5-glia- γ 4b	DQ2- γ -VIIC	P	Q	P	E	Q	E	F	P	Q
DQ2.5-glia- γ 4c	DQ2- γ -VIIa	Q	Q	P	E	Q	P	F	P	Q
DQ2.5-glia- γ 4d	DQ2- γ -VIIB	P	Q	P	E	Q	P	F	C	Q
DQ2.5-glia- γ 4e [§]		L	Q	P	E	Q	P	F	P	Q
DQ2.5-glia- γ 5	DQ2- γ -VI	Q	Q	P	F	P	E	Q	P	Q
DQ2.5-glia- ω 1	DQ2- ω -I	P	F	P	Q	P	E	Q	P	F
DQ2.5-glia- ω 2	DQ2- ω -II	P	Q	P	E	Q	P	F	P	W

Molecular basis for HLA association

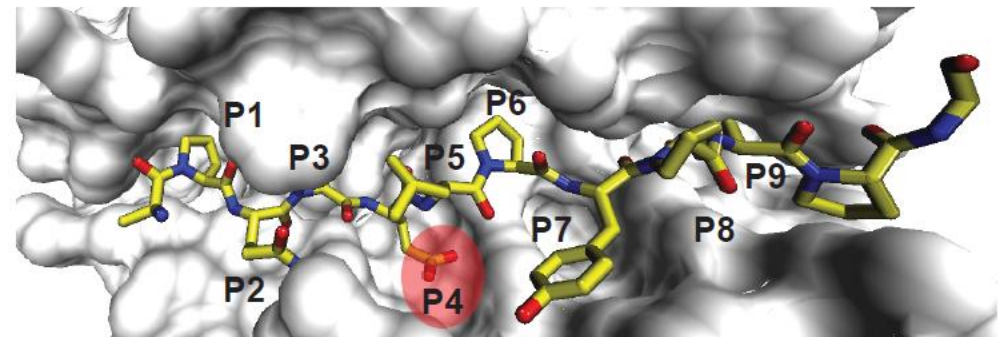
- Preference for negatively charged peptides.
- Gluten deamidation mediated by transglutaminase 2 (TG2).



b DQ2.5-glia- α 1a: P-F-P-Q-P-E-L-P-Y



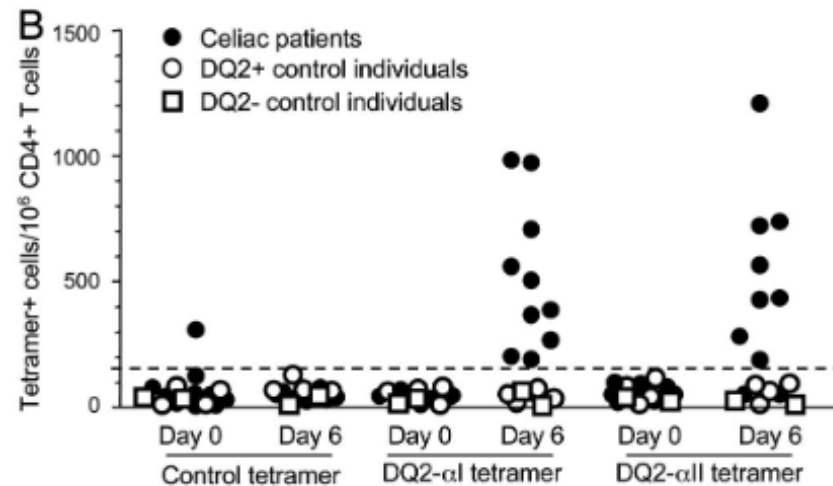
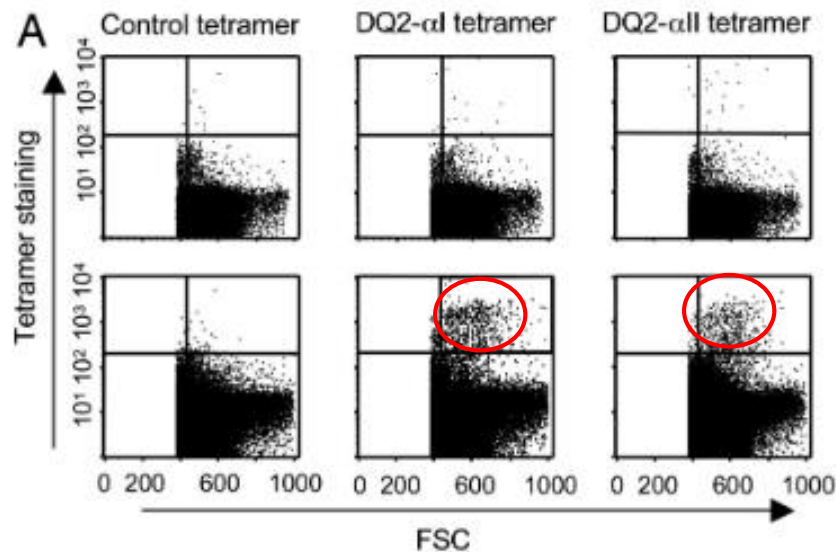
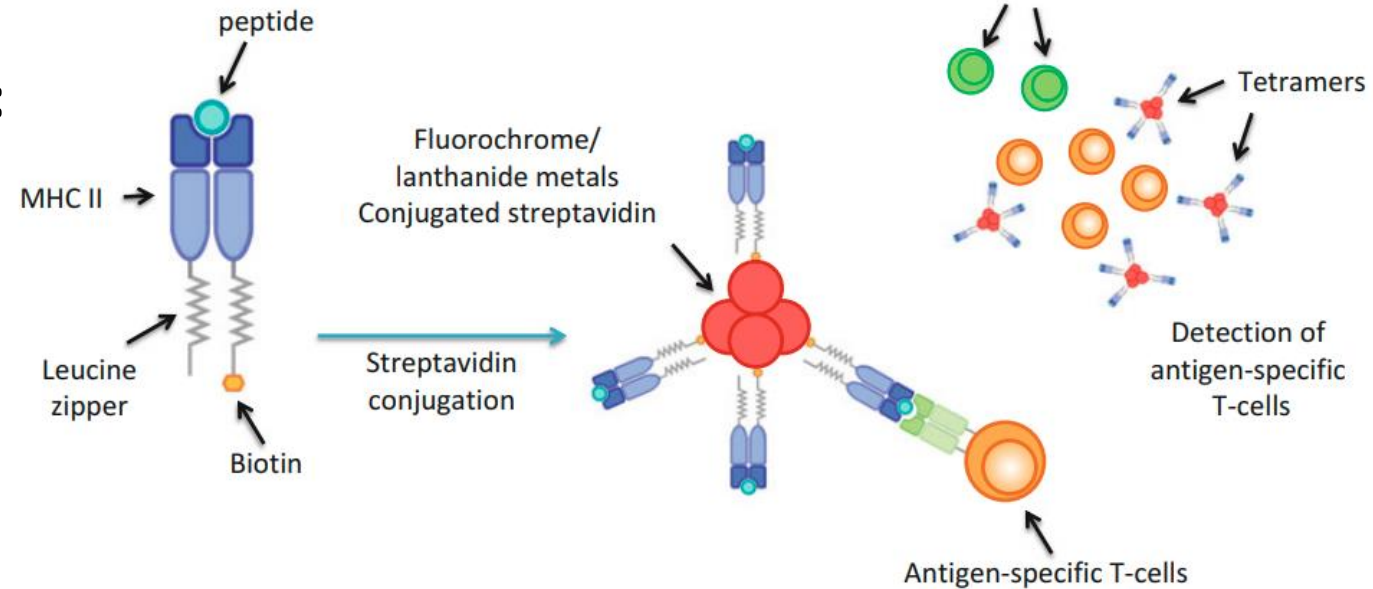
DQ2.5-glia- α 2: P-Q-P-E-L-P-Y-P-Q



Detection of gluten-specific T cells

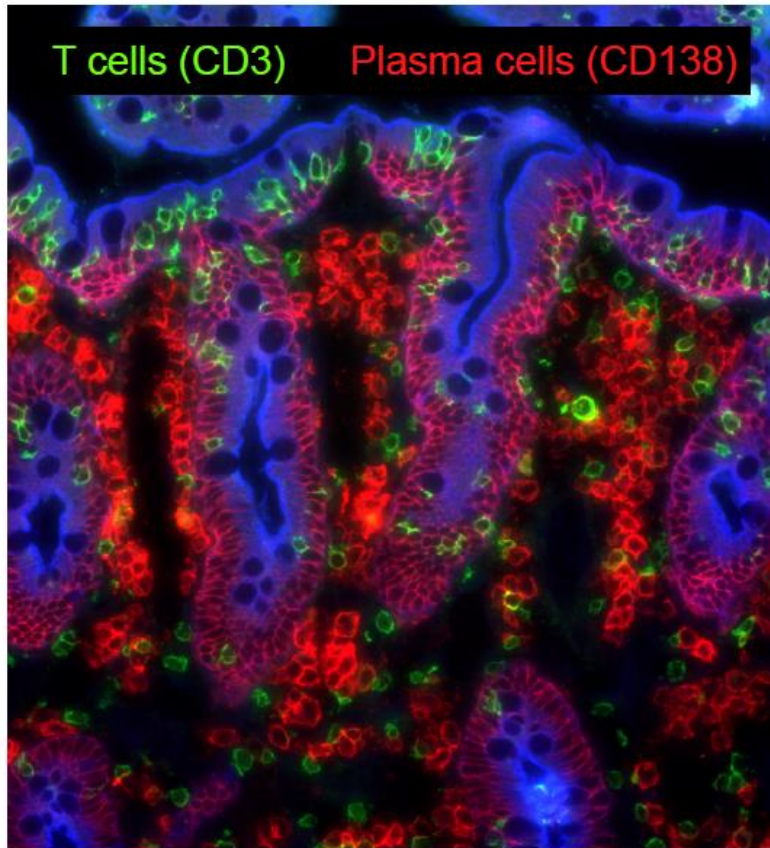
Advantages of studying celiac disease:

- Well-defined antigens.
- Good access to the disease lesion.
- Response can be turned on/off.

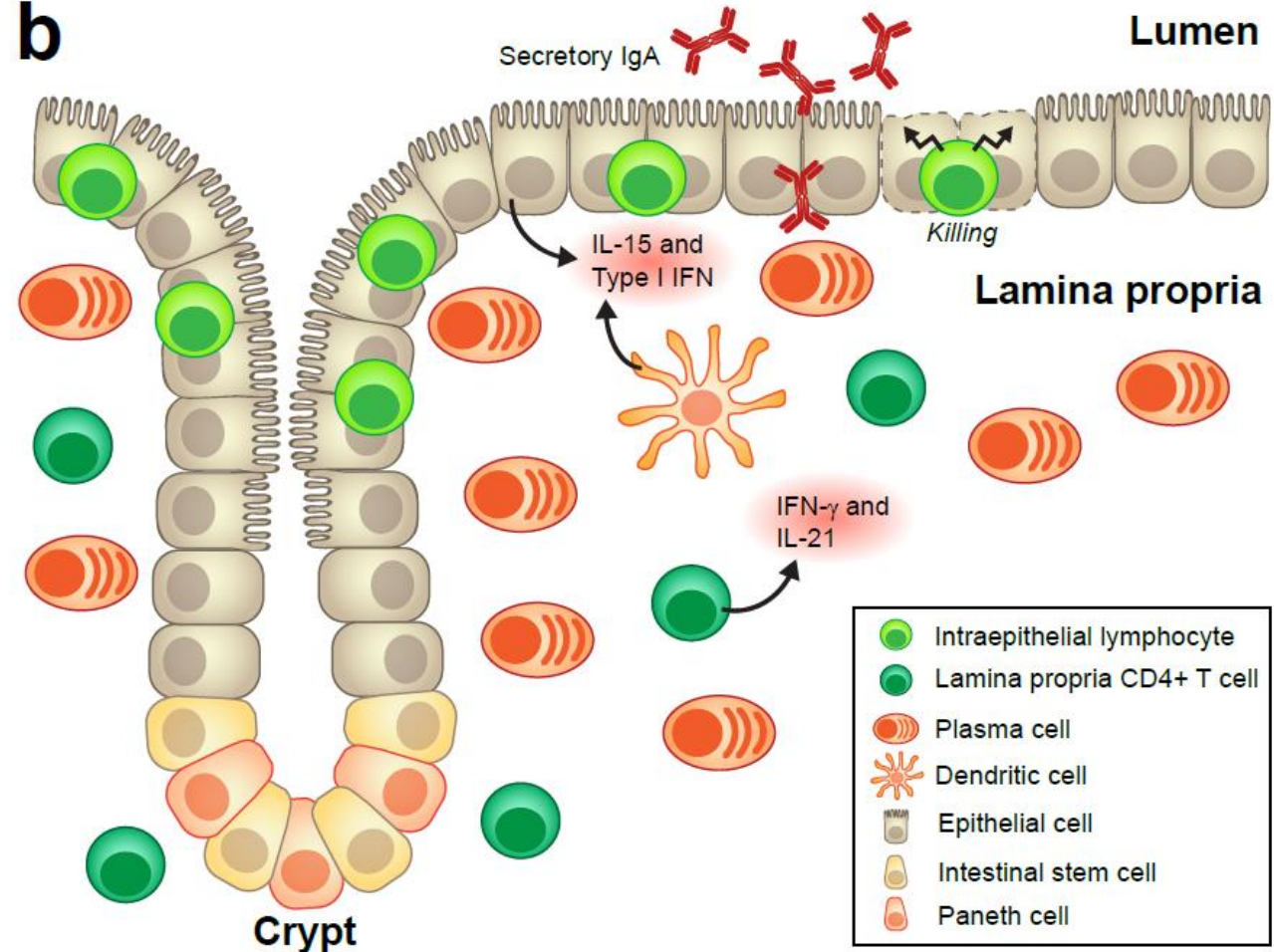


Immune cell infiltration in the gut

a

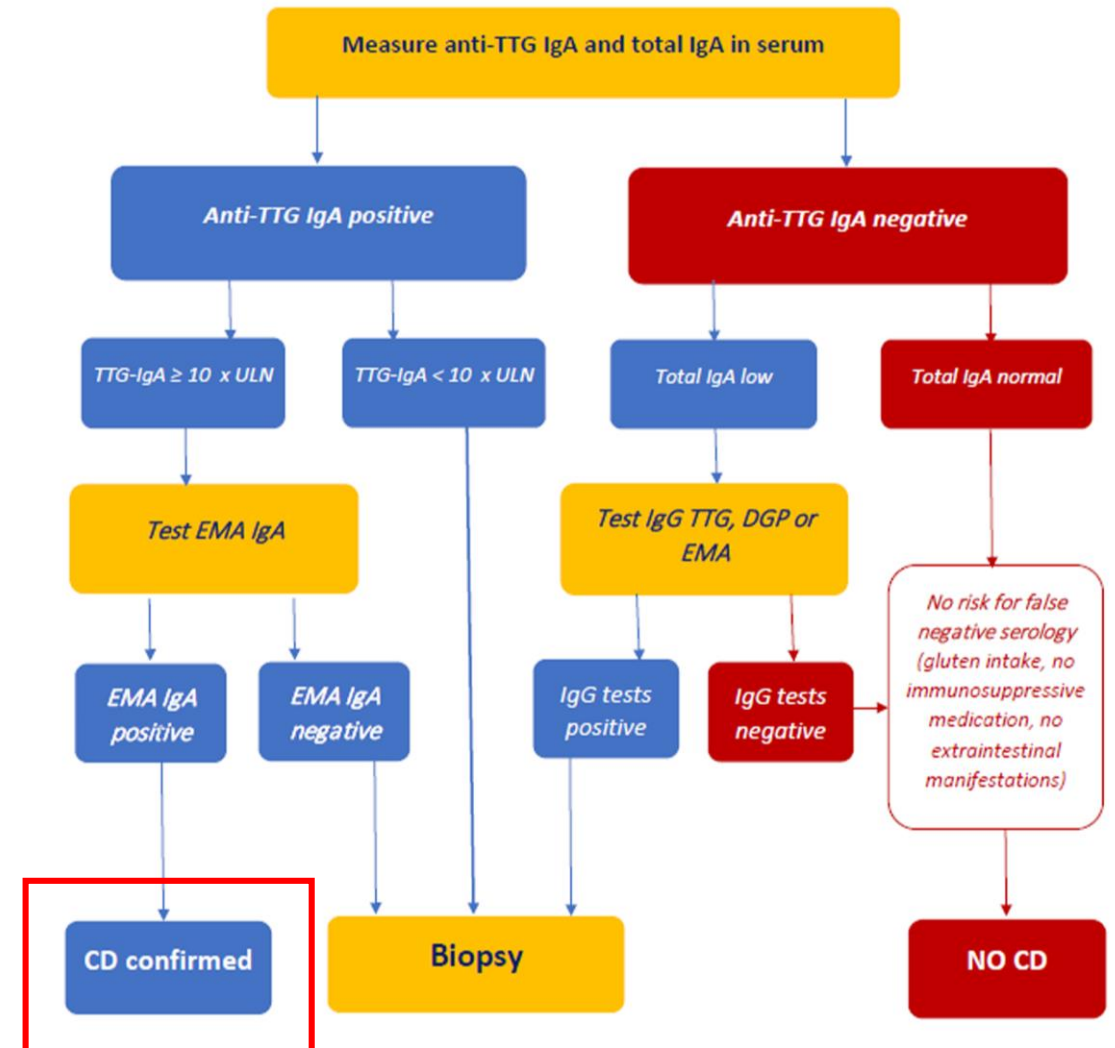


b



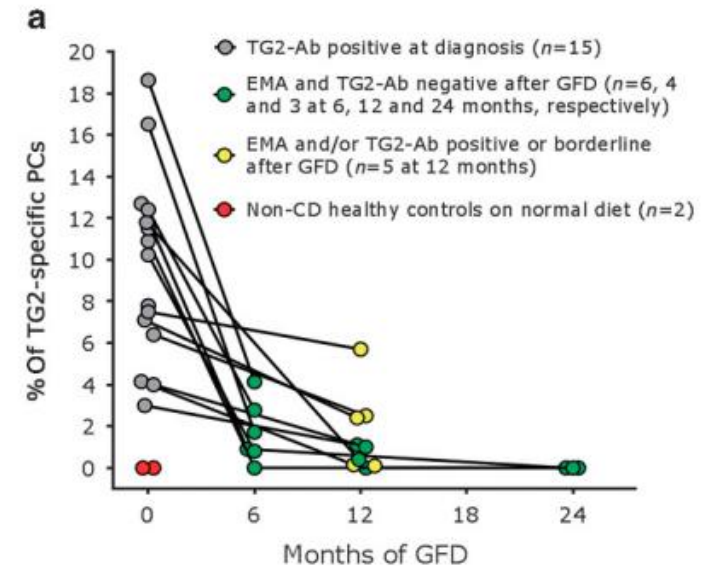
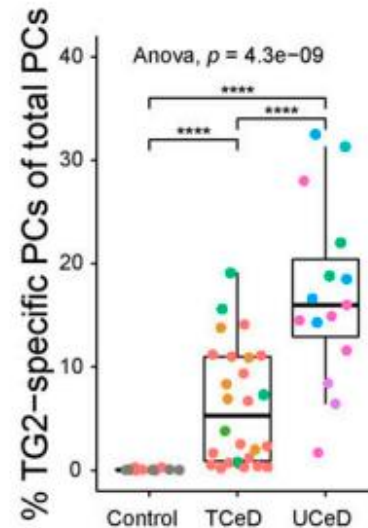
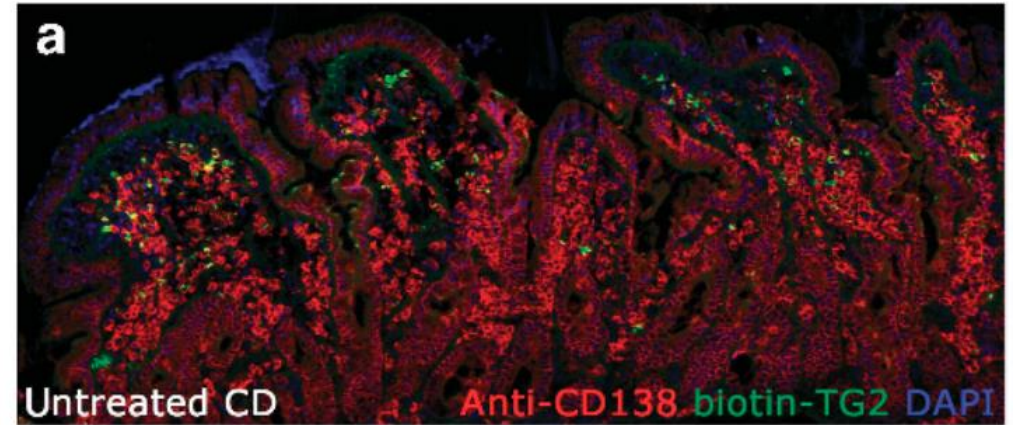
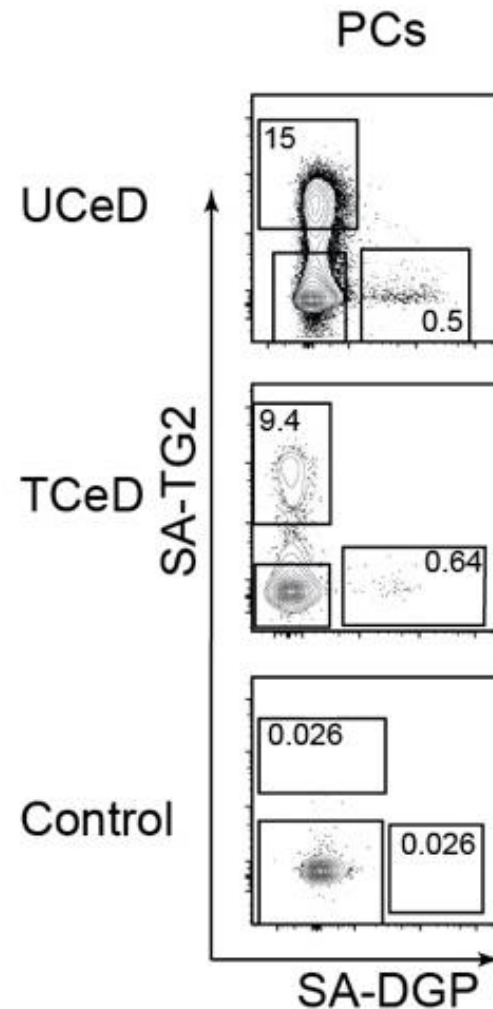
Serum antibodies in celiac disease

- Anti-deamidated gluten peptide (DGP) antibodies (IgG > IgA)
- Anti-tissue transglutaminase (TTG or TG2) antibodies (IgA > IgG)
- Detection with recombinant TG2 or tissue sections – endomysial antibodies (EMA)



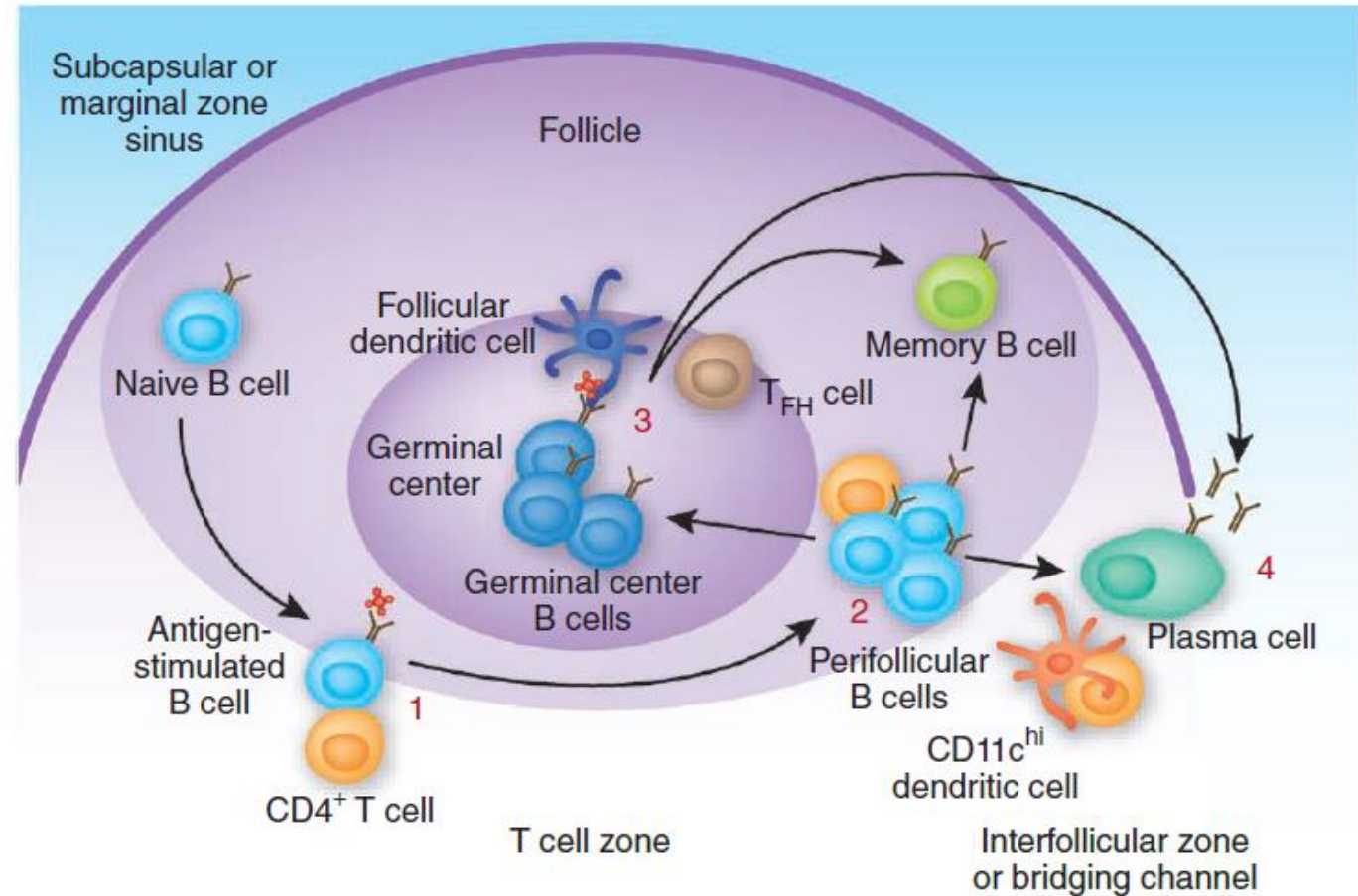
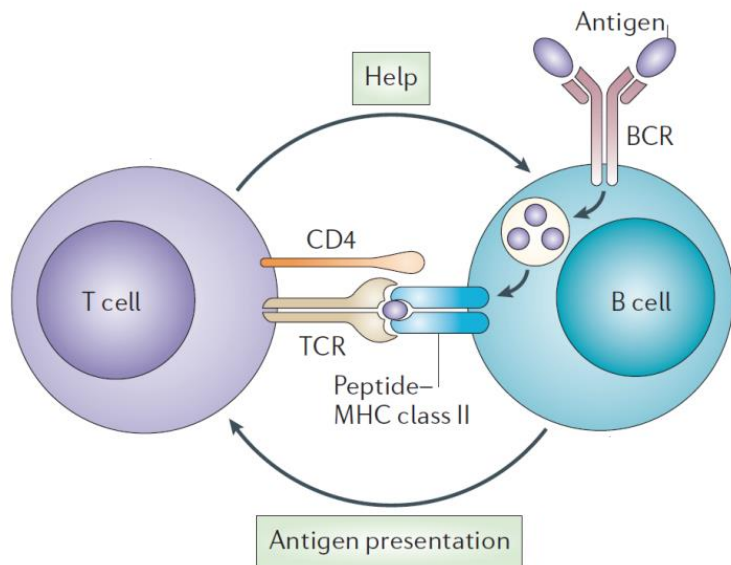
Antibody-secreting plasma cells in the gut

- 10-20% TG2-specific.
- <1% DGP-specific.
- Dependent on gluten in the diet.



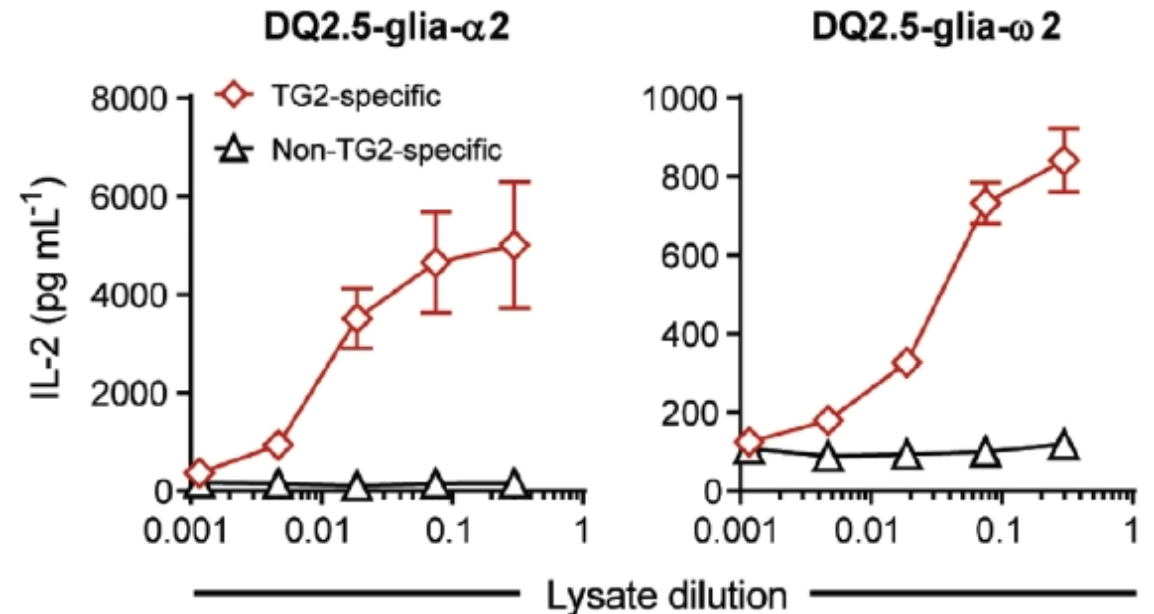
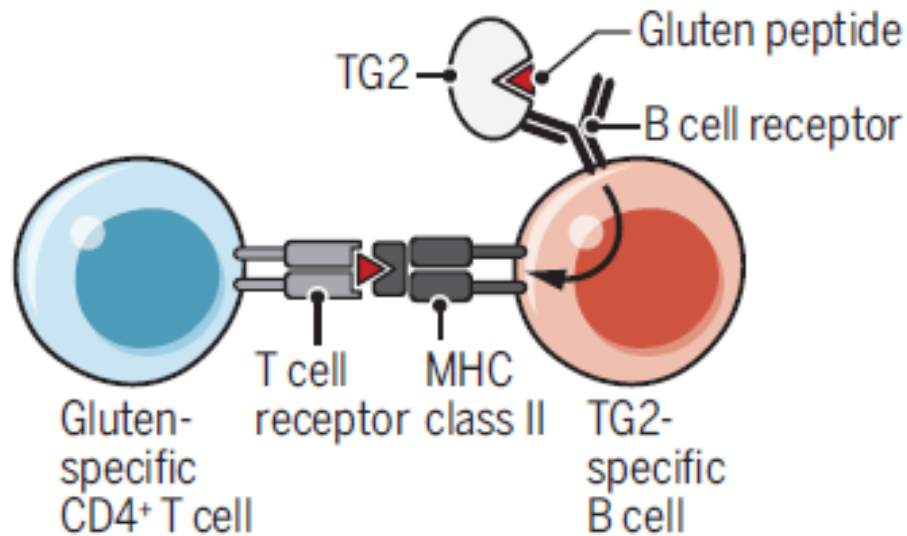
B-cell activation

- Antigen binding → BCR signaling.
- Antigen uptake → interaction with cognate CD4⁺ T cells.
- Proliferation, differentiation to memory cells and plasma cells (antibody production).

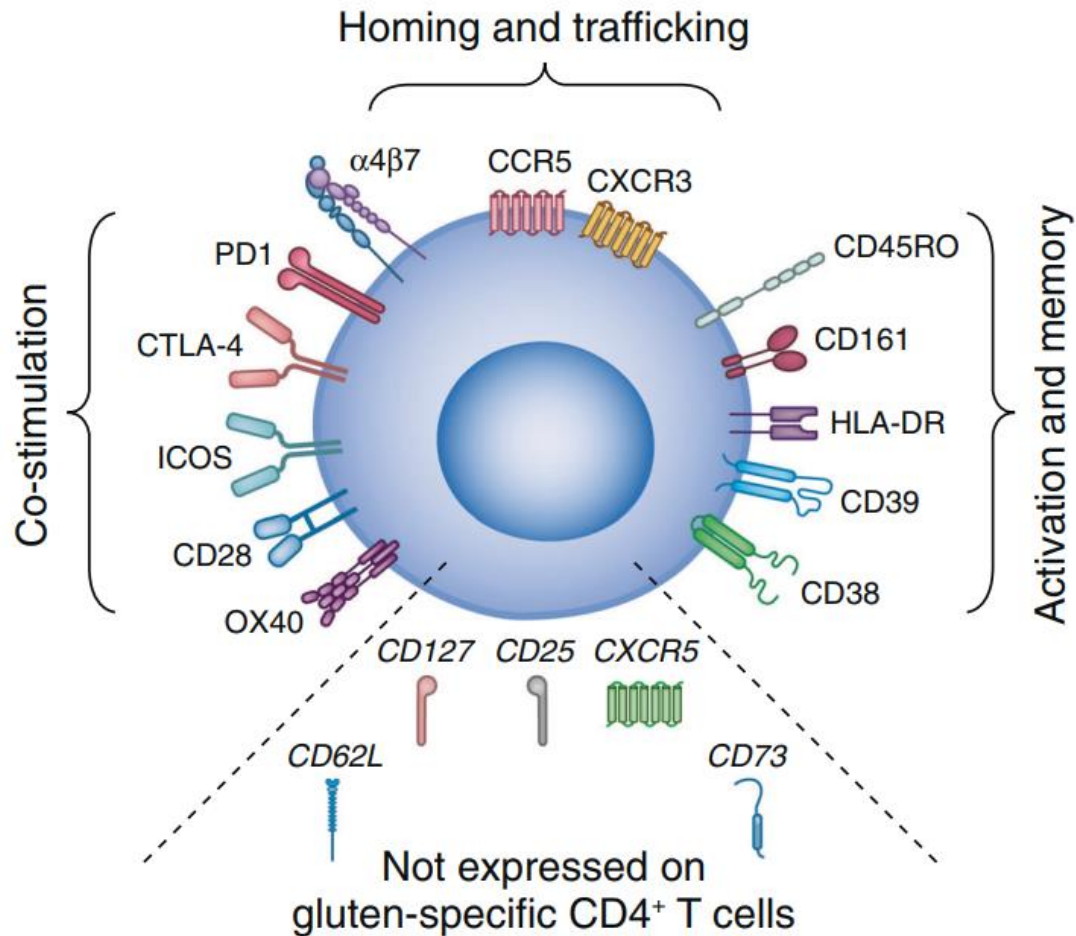


What is the role of TG2-specific B cells / antibodies?

- Soluble TG2-specific antibodies do not have a clear pathogenic effect.
- B cells with TG2-specific BCR can present antigen to gluten-specific T cells.
- TG2-specific B cells main antigen-presenting cell for gluten-specific CD4+ T cells.

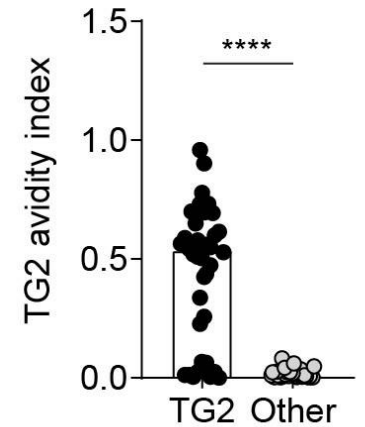
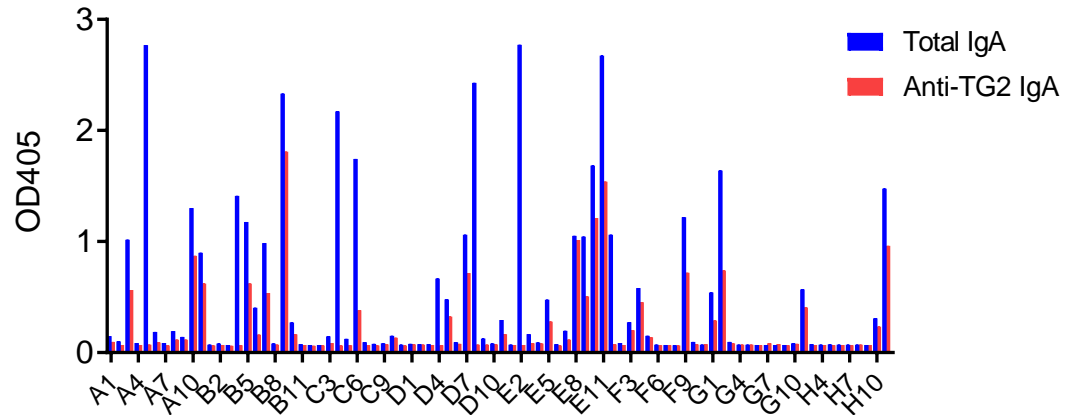
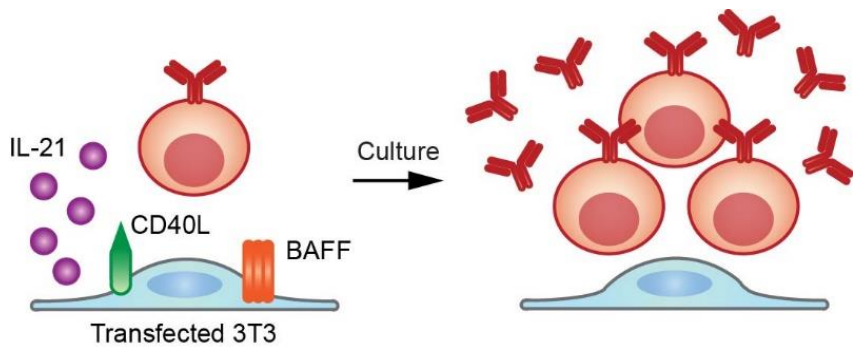
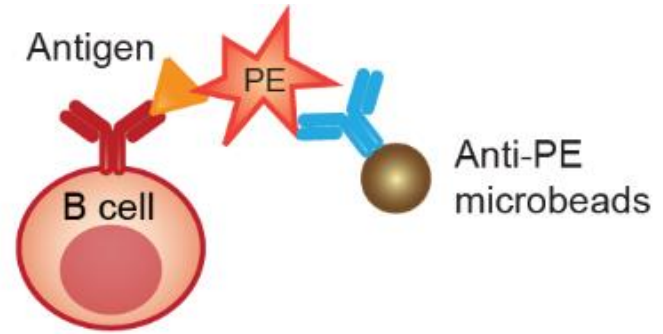
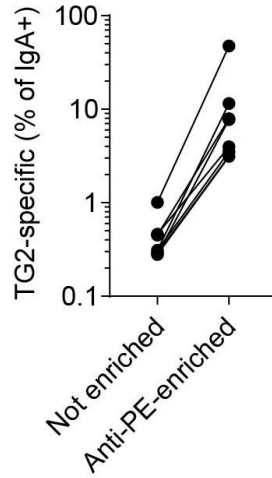
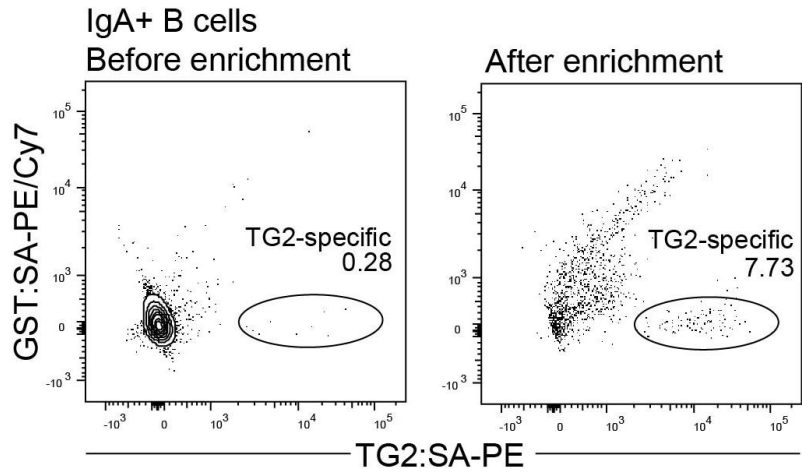


T-cell phenotype



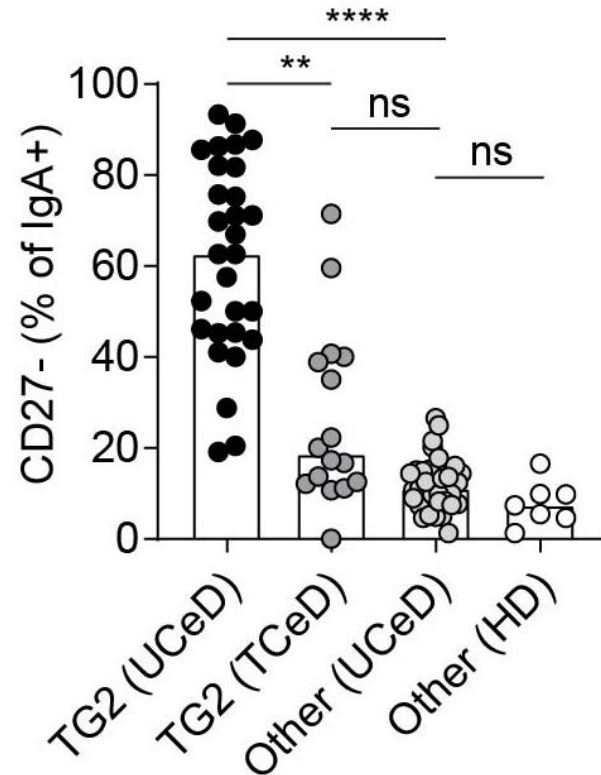
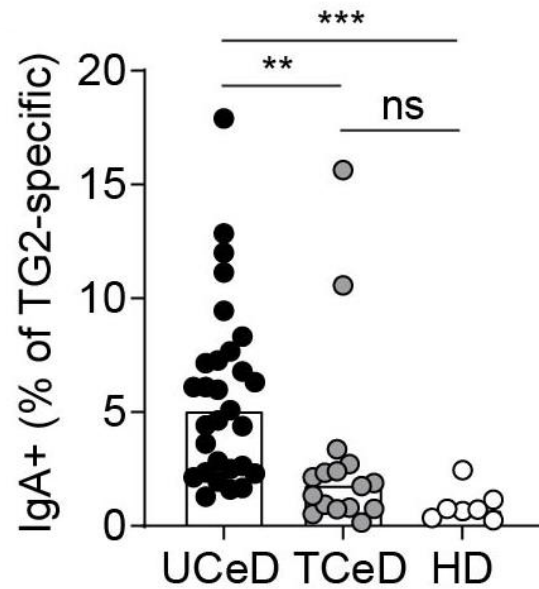
- Characterization by mass cytometry and single-cell RNA sequencing.
- Surface markers indicating activation and gut-homing.
- Secretion of B-helper cytokines IL-21 and CXCL13.
- Similar to previously described “T peripheral helper cells” (Tph).

Identifying TG2-specific B cells in blood



TG2-specific IgA+ B cells

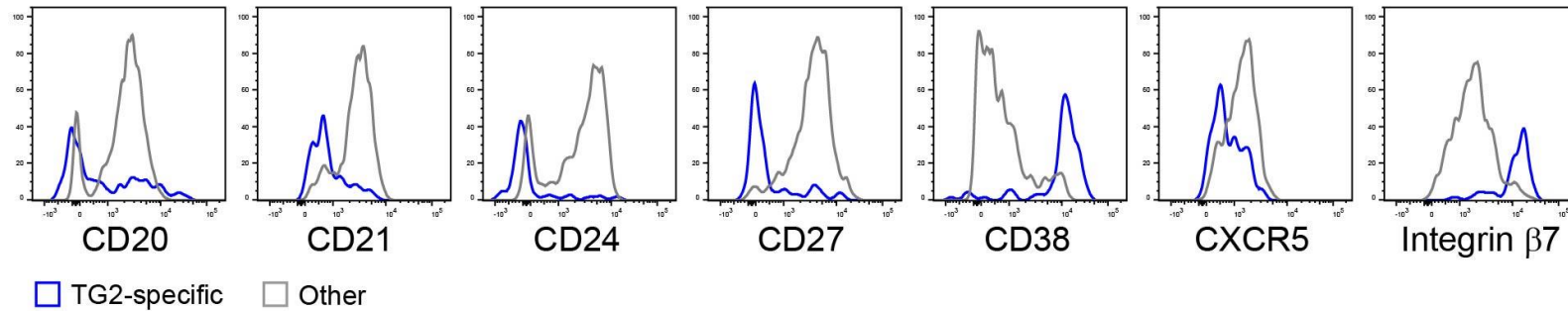
- Present in untreated patients.
- Negative for the classical memory marker CD27.



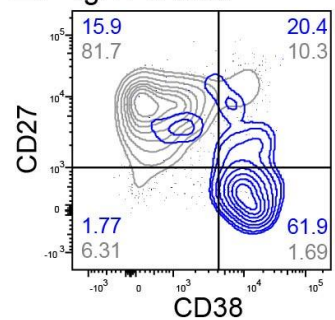
B-cell phenotype – flow cytometry

- Surface markers indicating activated, gut homing memory B cells / plasmablasts.
- Activated cells disappear rapidly on gluten-free diet.

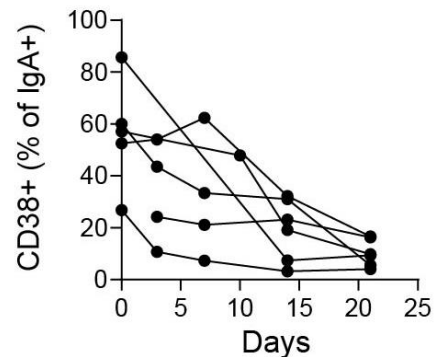
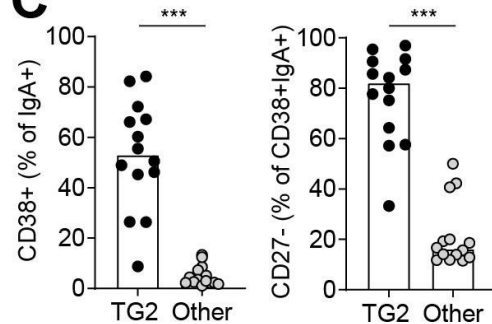
A IgA+ B cells



B IgA+ B cells

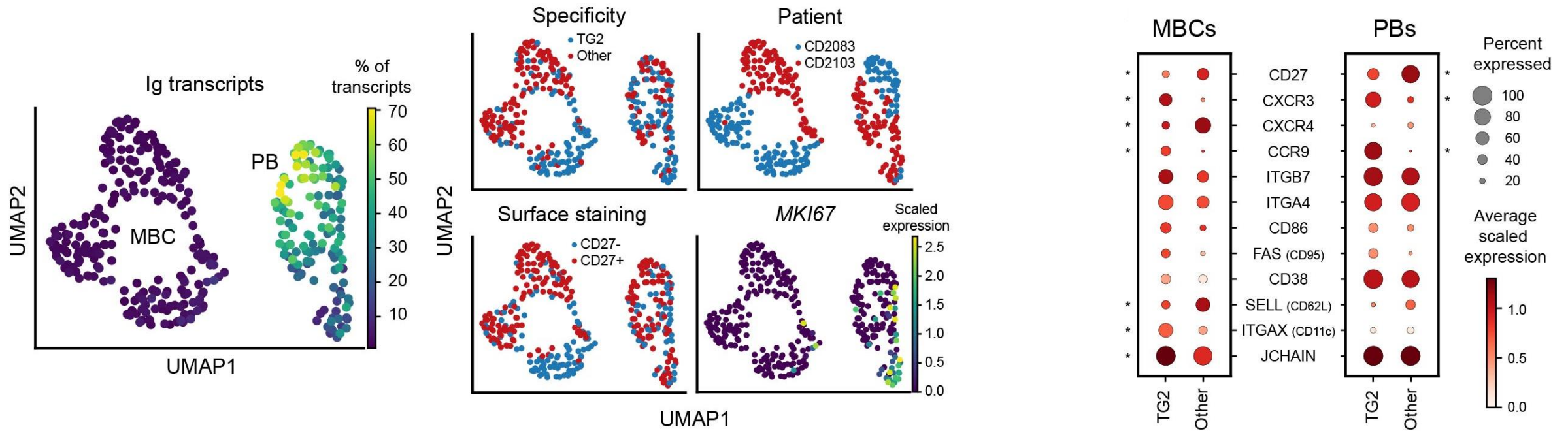


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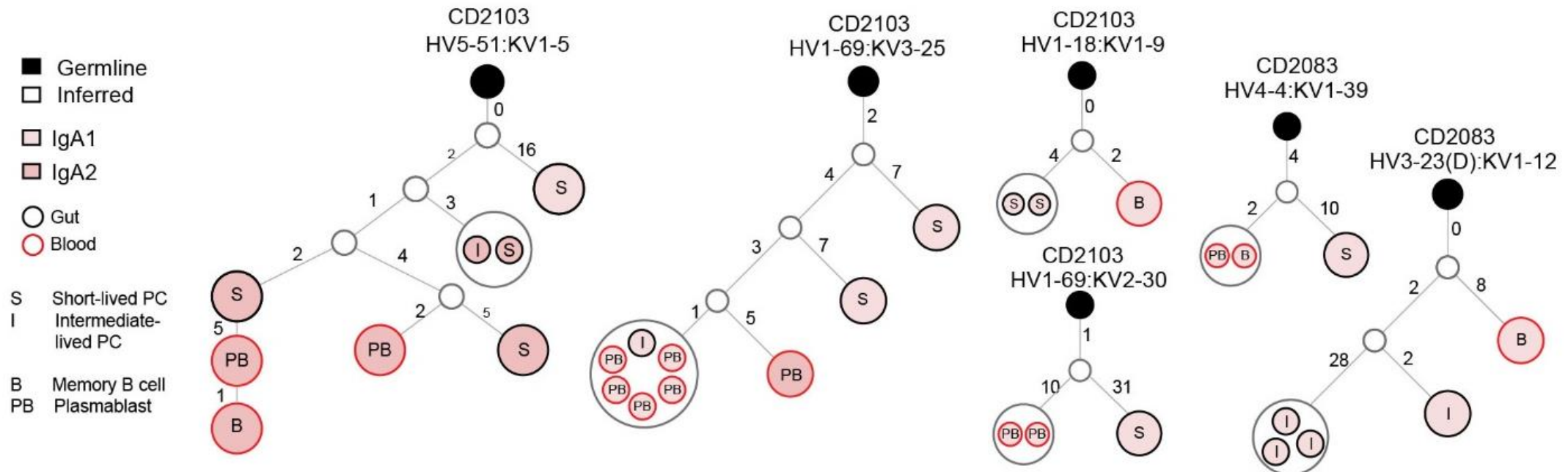
Single-cell RNAseq

- Two main clusters: memory B cells and plasmablasts.
- CD27 negative, activation, gut-homing.

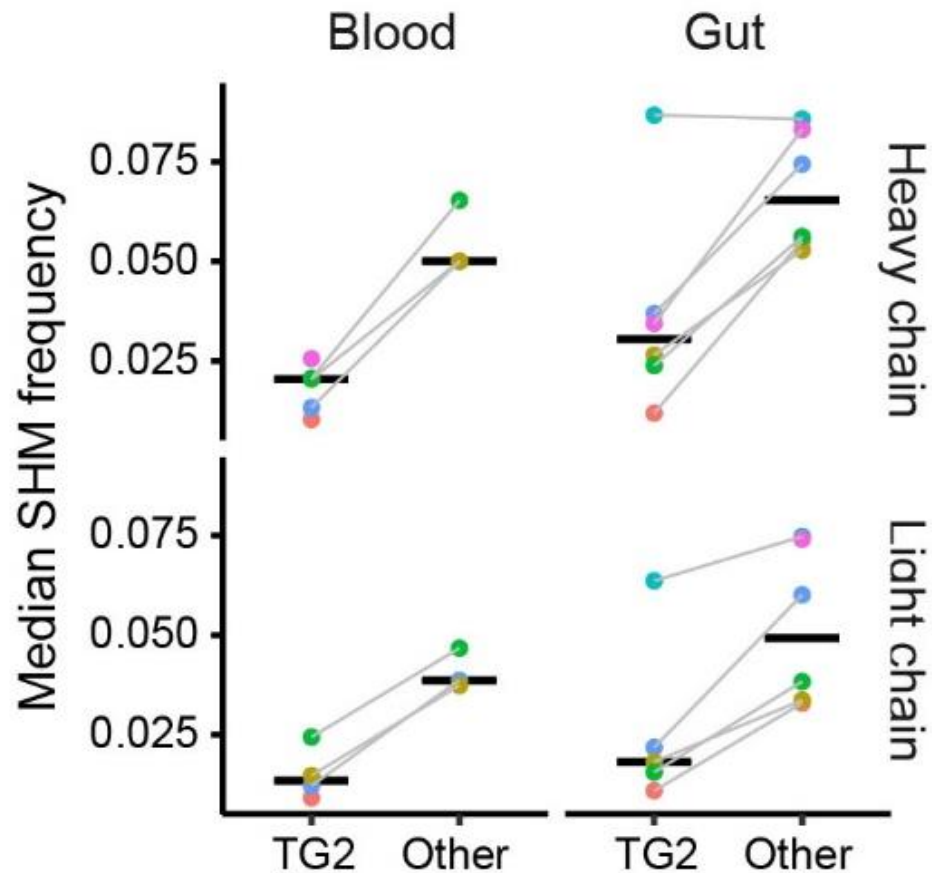


V(D)J sequences

- Clonal overlap between TG2-specific cells in blood and gut (same ancestral B cells).
- Cells in blood are precursors of gut plasma cells.

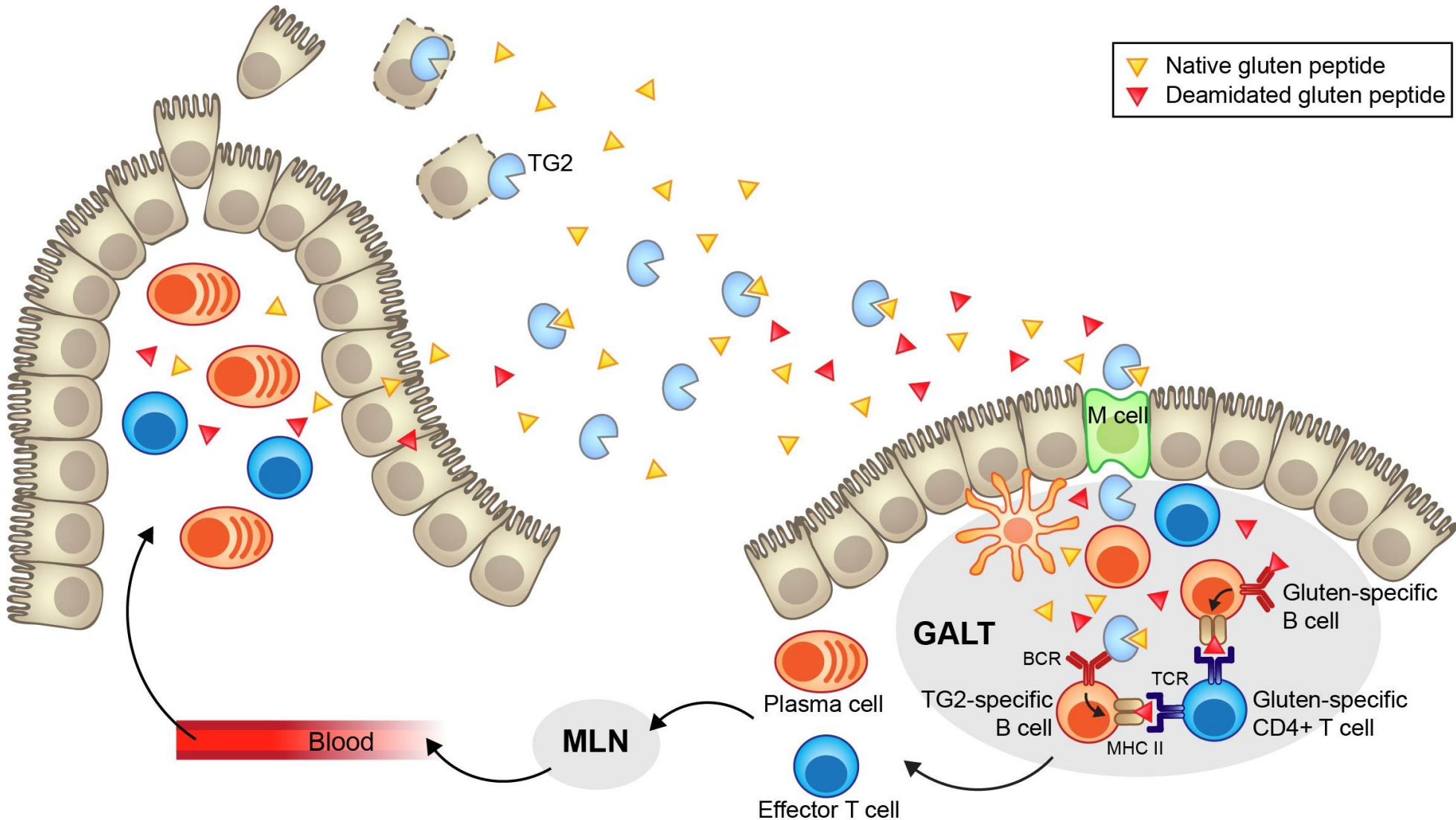


BCR mutations



- TG2-specific cells have fewer mutations than other B cells/plasma cells.
- SHM mainly happens in germinal centers.
- CD27 is expressed in germinal centers.
- Lack of CD27 and SHM could indicate extrafollicular origin.
- Supported by phenotype of gluten-specific T cells (Tph).

A model for celiac disease pathogenesis



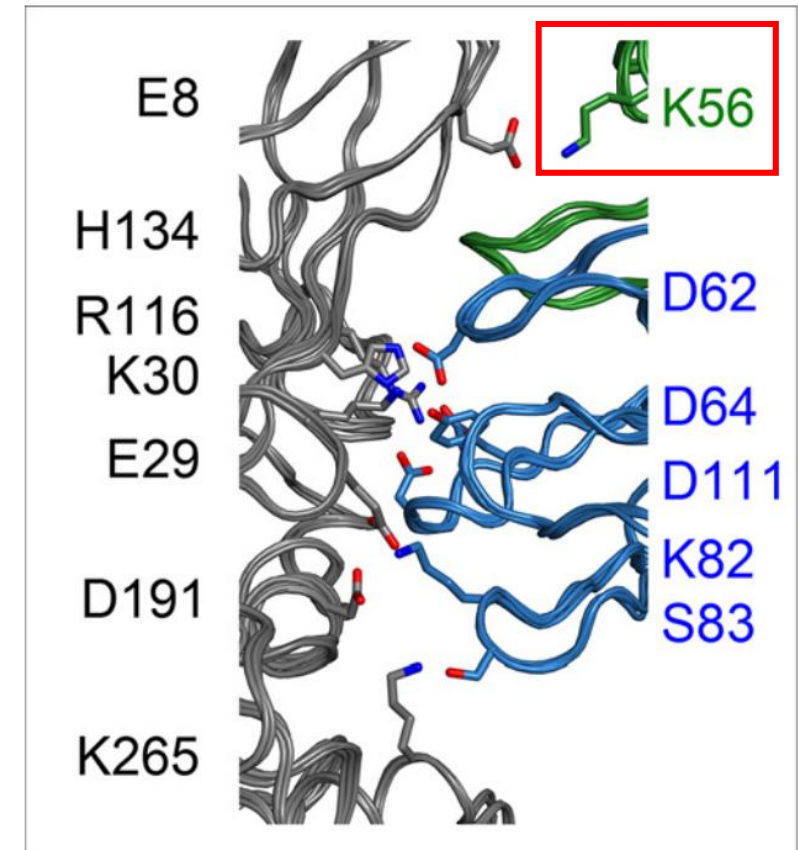
Could BCR and TCR be disease susceptibility genes?

Preference for certain V-genes observed for:

- TG2-specific B cells/plasma cells
- DGP-specific plasma cells
- Gluten-specific T cells

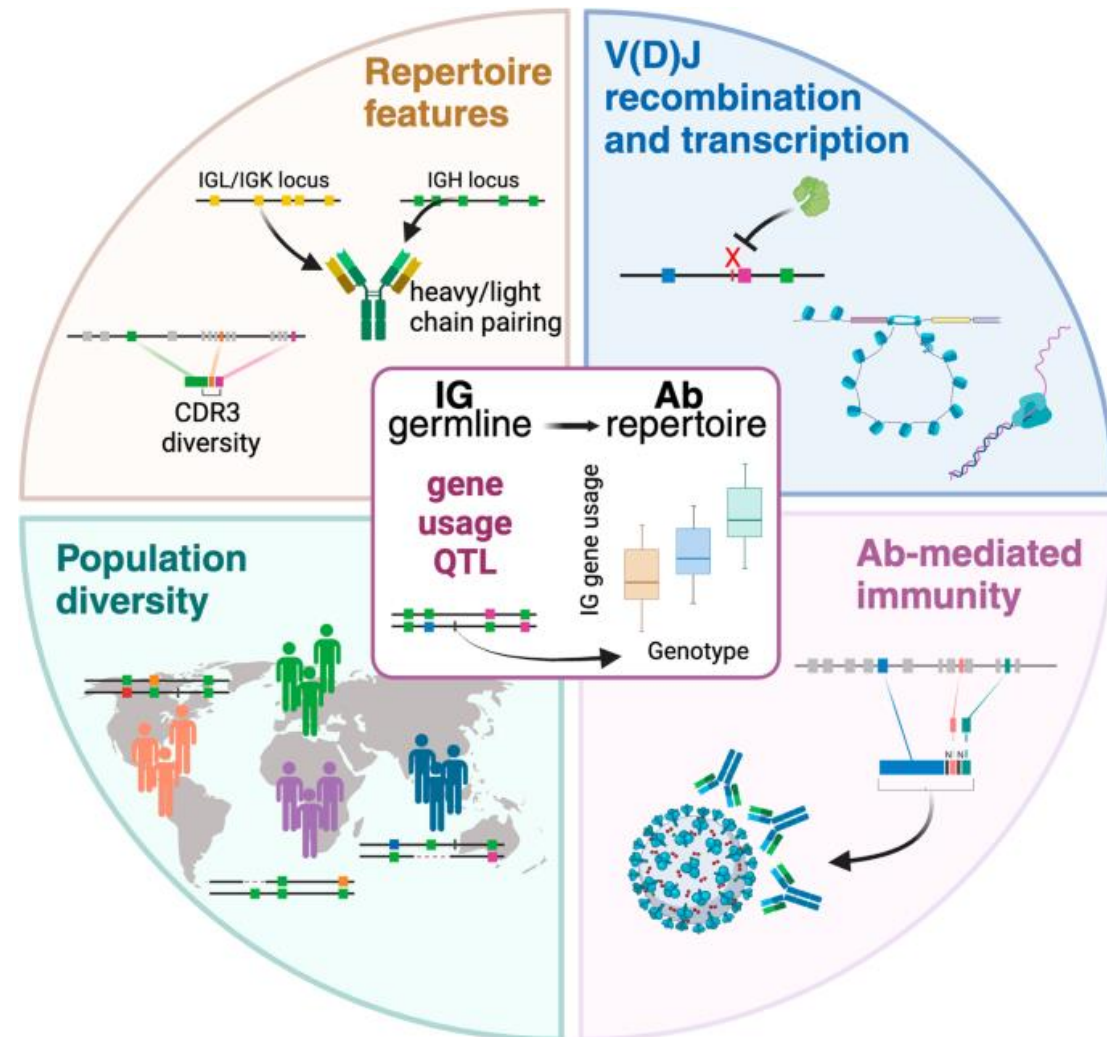
Most common heavy/light chain combination for TG2-specific cells: *IGHV5-51:IGKV1-5*

IGKV1-5 mutation K56D
→ 200 fold reduction in affinity



Genetic diversity in BCR and TCR genes

- Bigger diversity than previously appreciated
- Both in coding and non-coding regions
- Can affect repertoire composition
- Implications for precursor frequencies
- BCR and TCR regions not covered in GWAS



Conclusions

- HLA dominates genetic predisposition for celiac disease
- Both HLA and non-HLA genes point to T-cell activation as key pathogenic event
- B cells likely act as main antigen-presenting cells for gluten-specific T cells
- TG2-specific B cells internalize enzyme-substrate complexes of TG2 and gluten
- Directly links antigen uptake and deamidation
- Phenotype of both T cells and B cells suggests extrafollicular interactions at induction sites
- Importance of TCR and BCR specificity implicates them as potential susceptibility genes

Acknowledgements

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- Filipa Vaz
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- Louise Risnes

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- Knut Lundin

Akershus University Hospital

- Jørgen Jahnsen

And all the patients!!!

HELSE  SØR-ØST



UiO  **University of Oslo**