

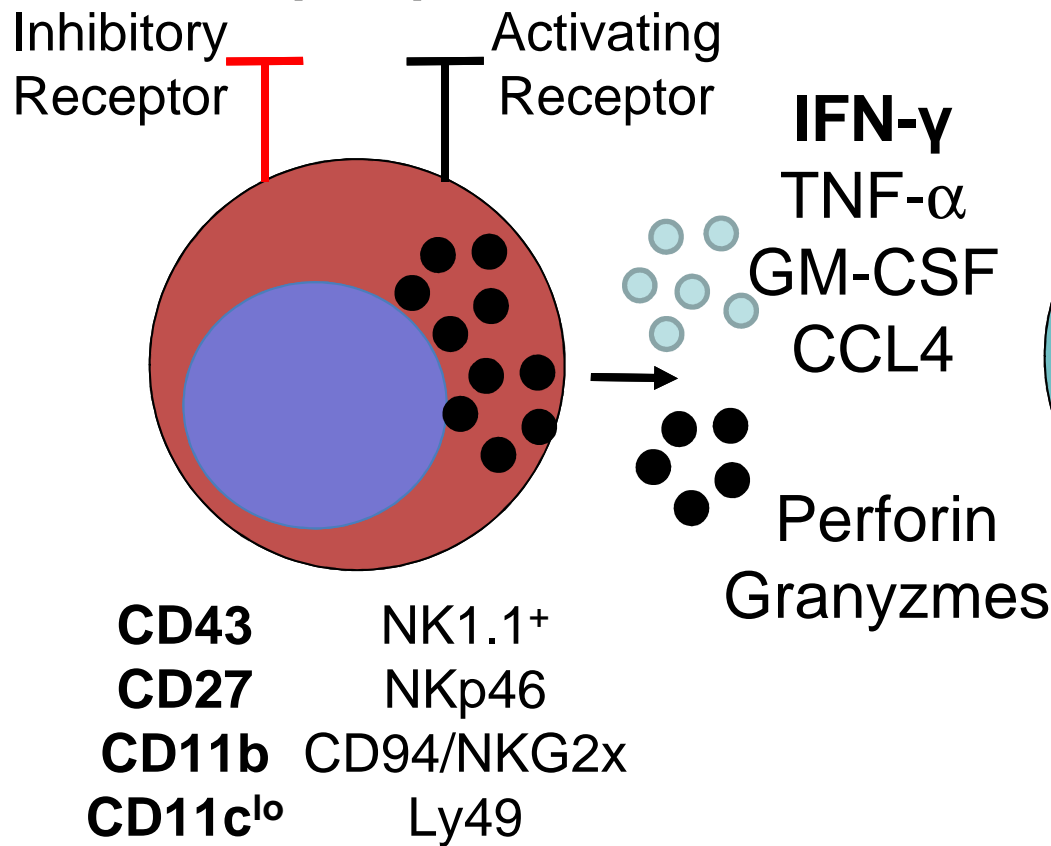


Innate lymphoid cells (ILC) in immunity

**Marco Colonna
Washington University in St Louis**

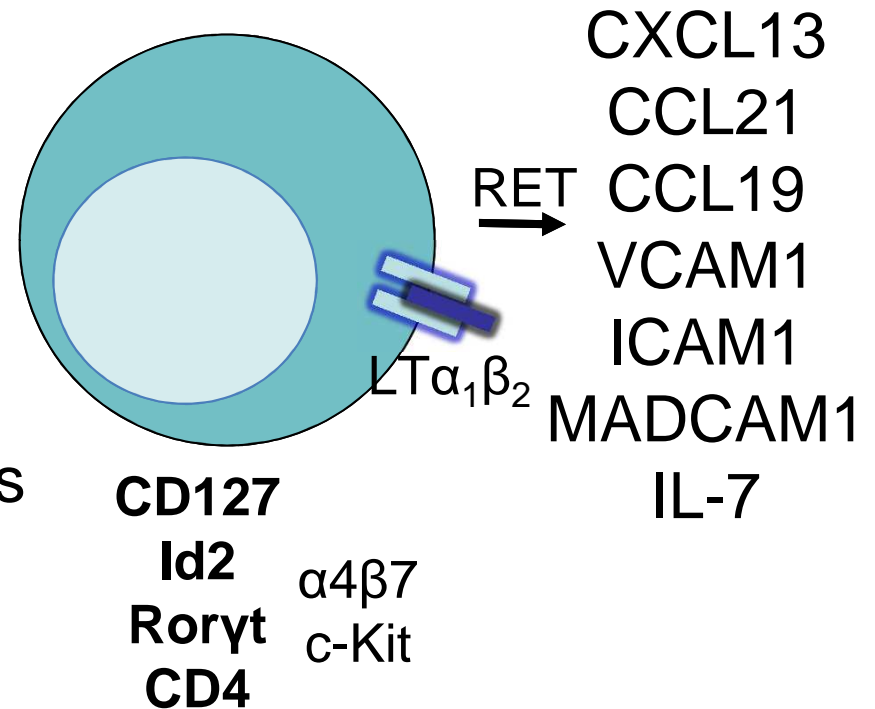
Innate Lymphoid Cells (ILC) before 2009:

Natural killer cell (NK)



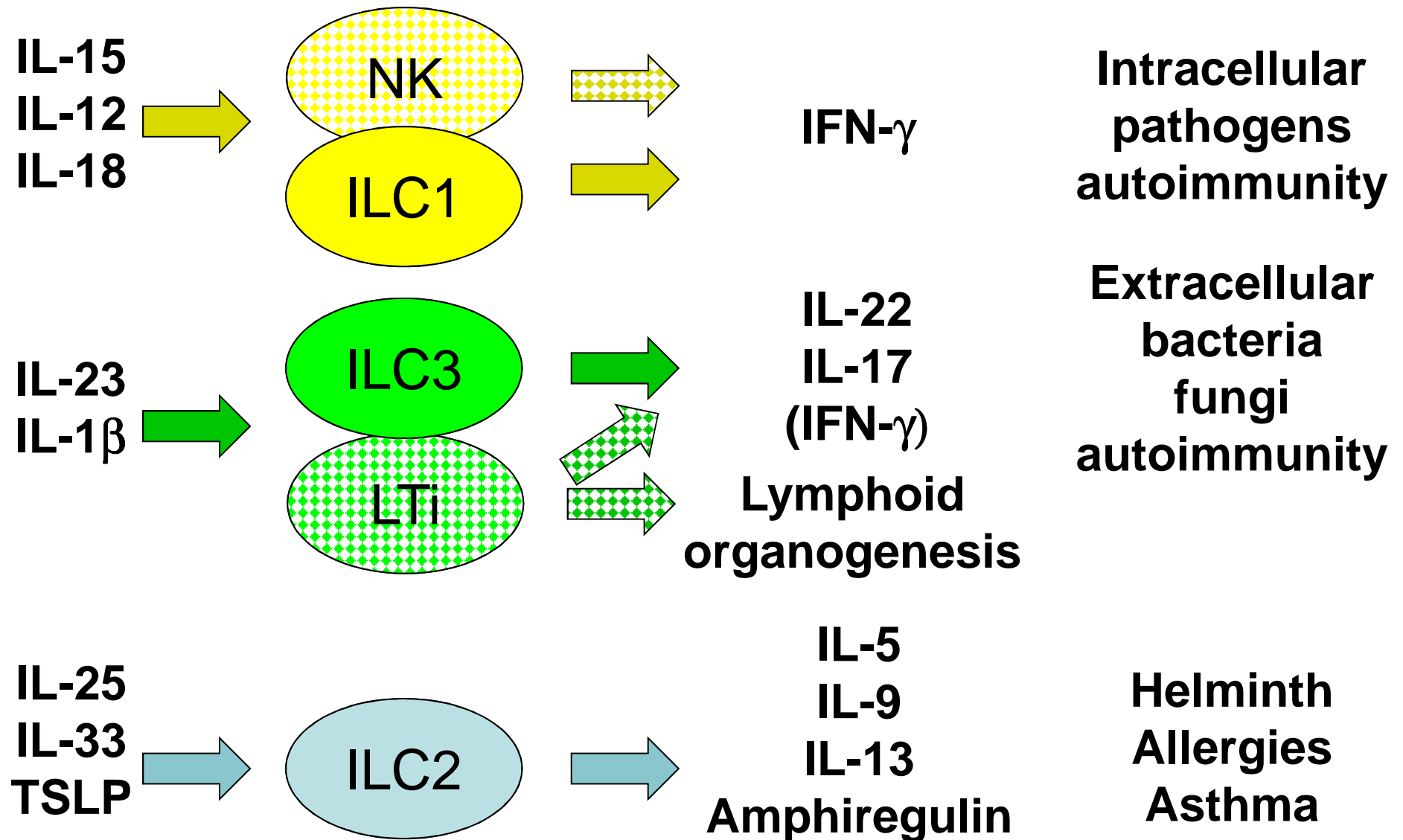
Cytotoxicity

Lymphoid tissue inducer cell (LTi)

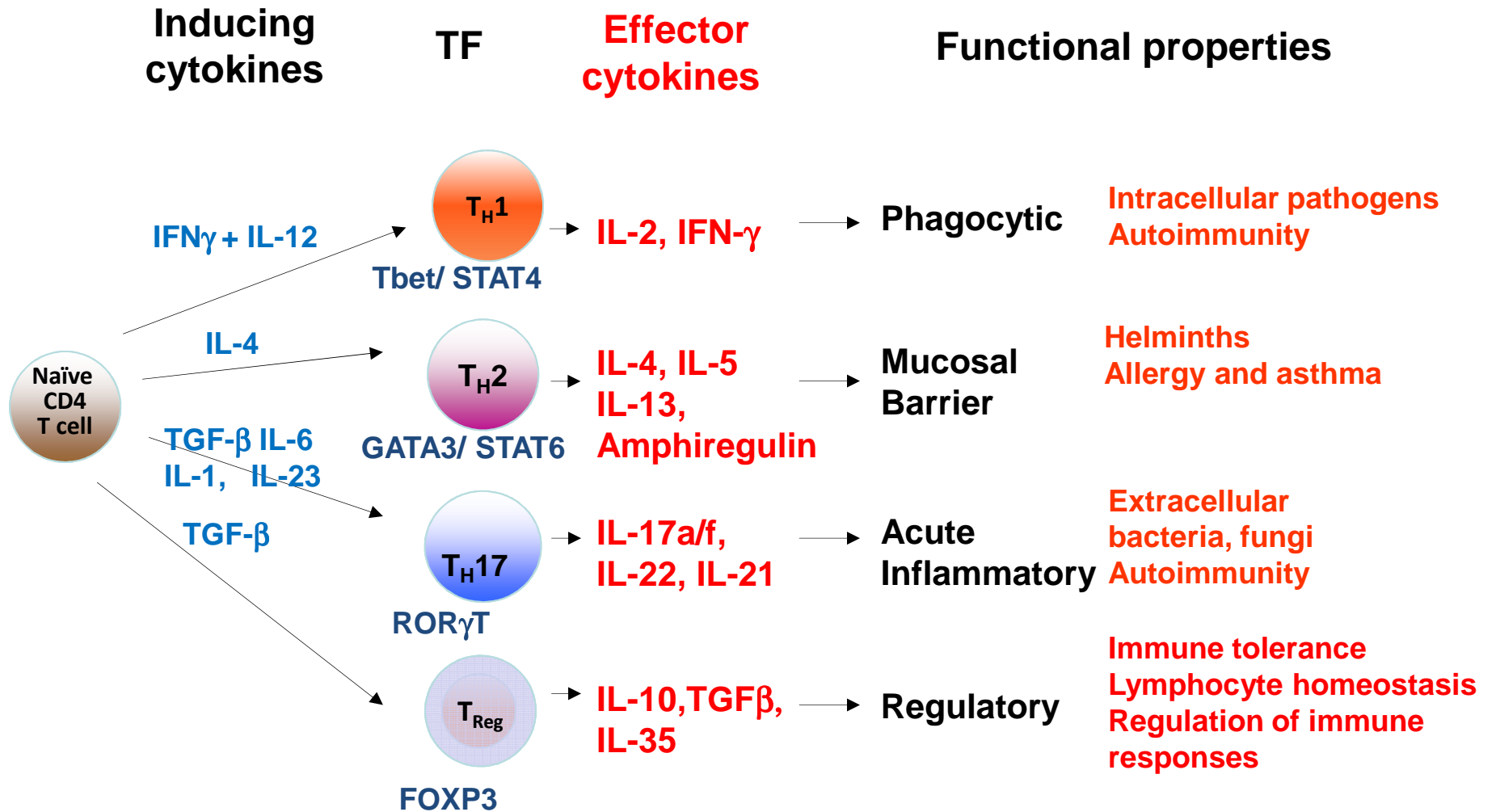


Lymphoid organogenesis

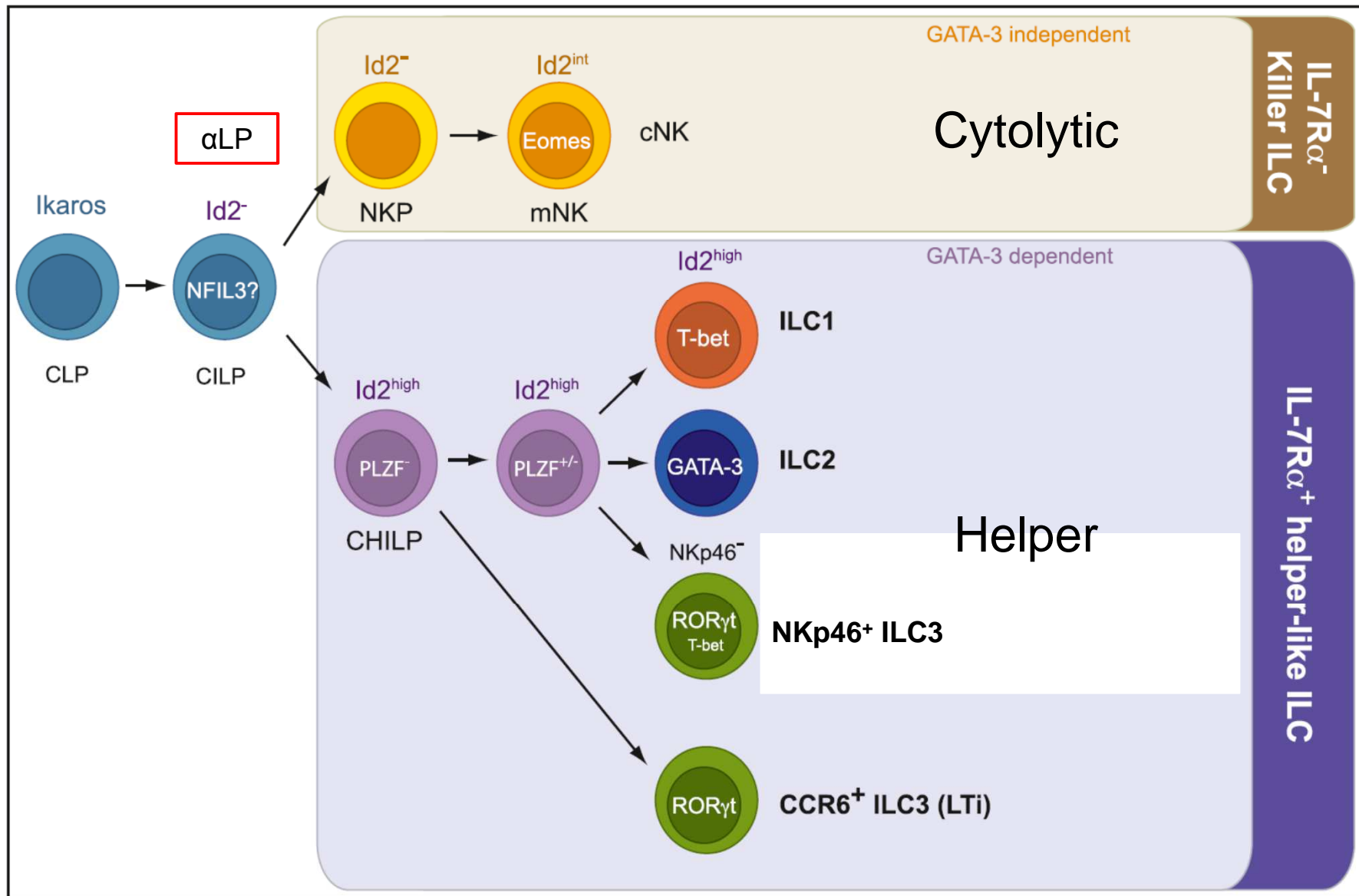
Innate Lymphoid Cells (ILC) after 2009



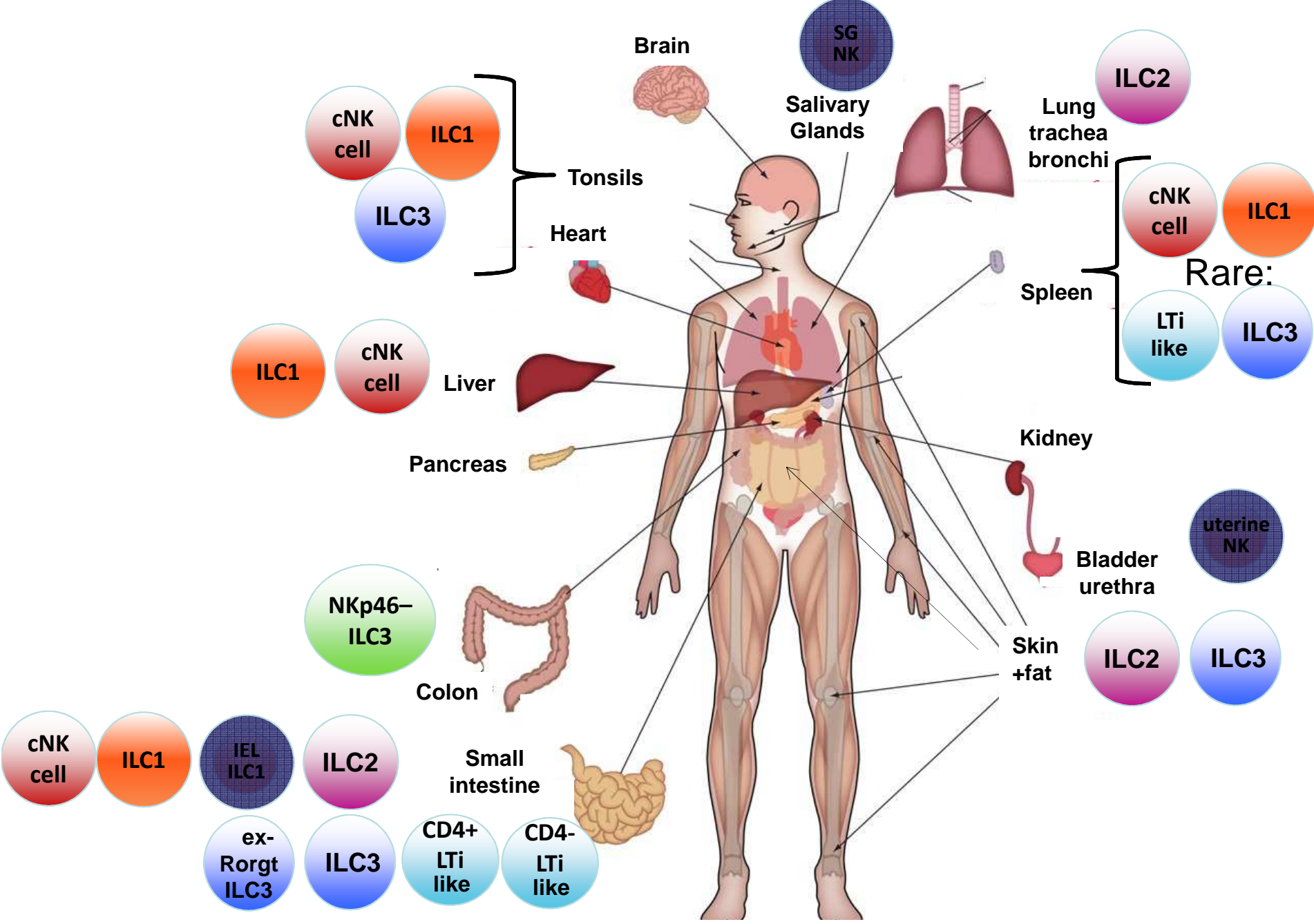
Functional modules of CD4 Th cells



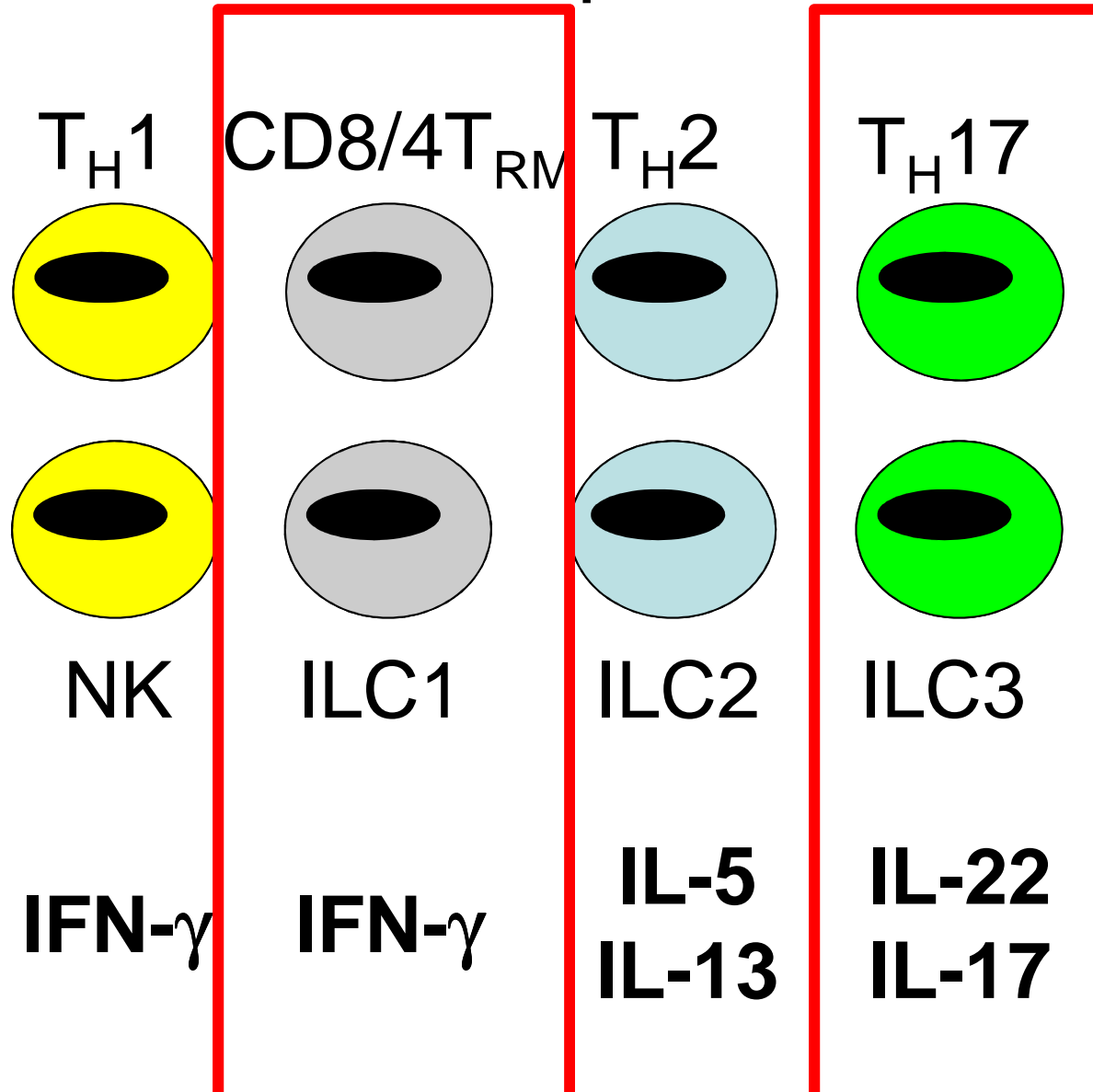
Transcription factors and cytokines in ILC development



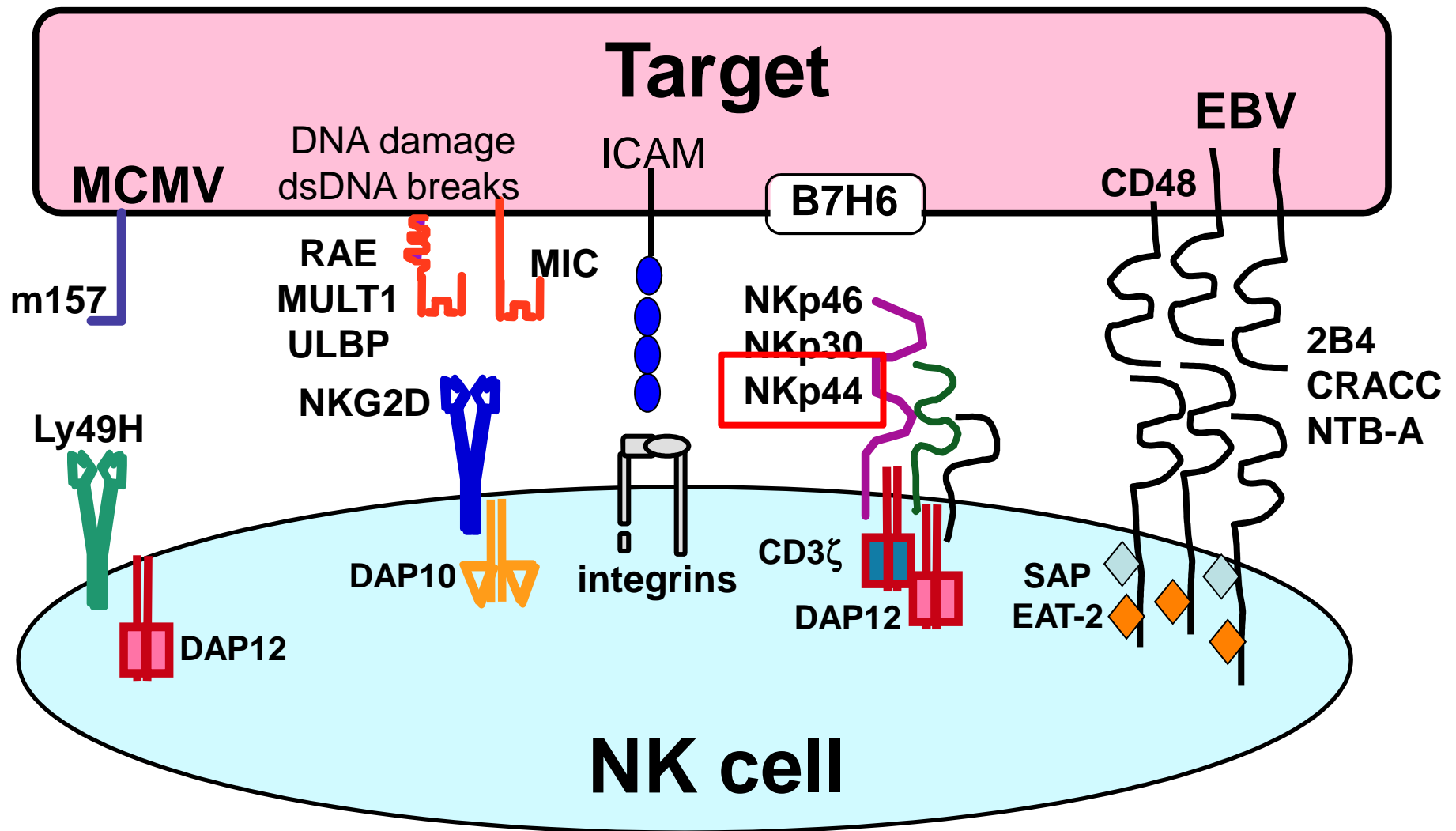
ILCs are tissue resident cells



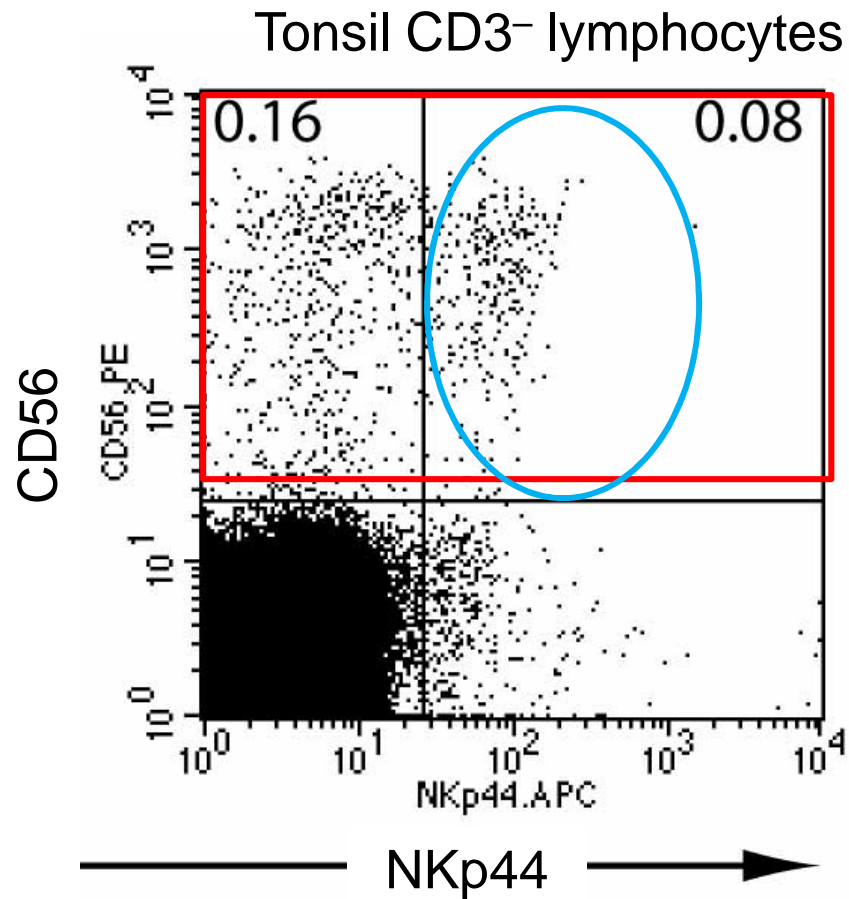
Similarity of functional modules in innate and adaptive immunity



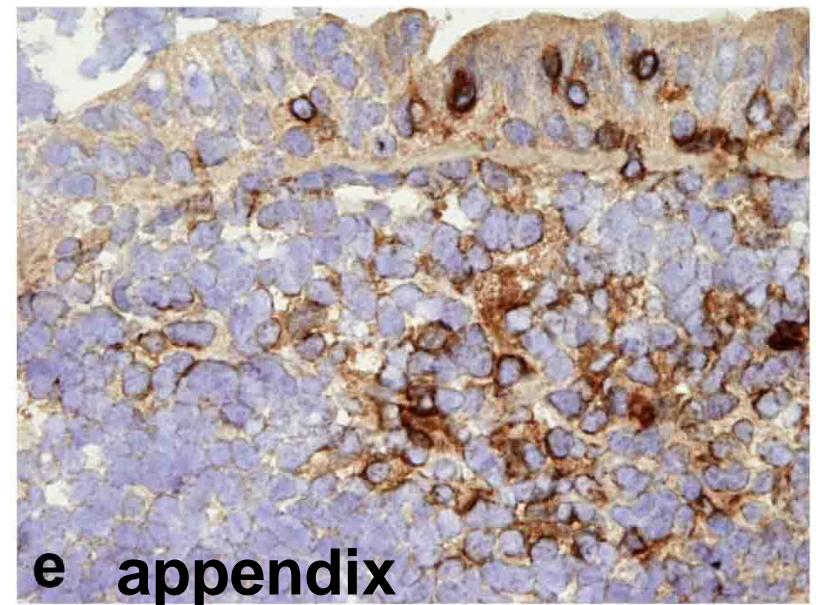
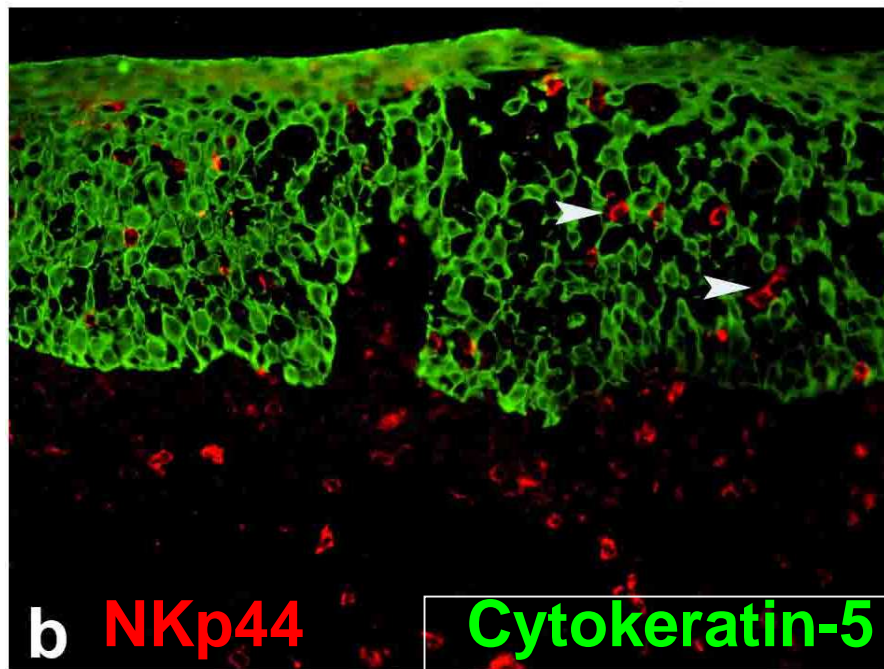
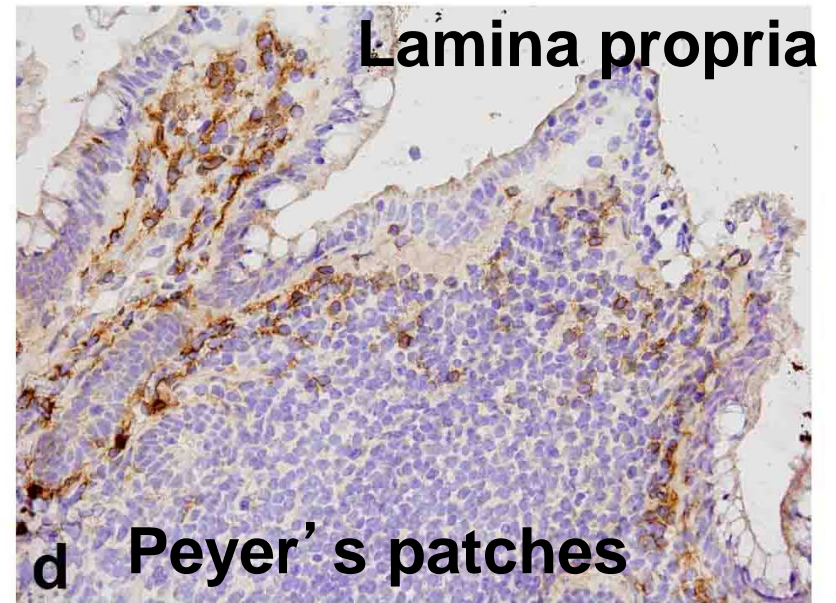
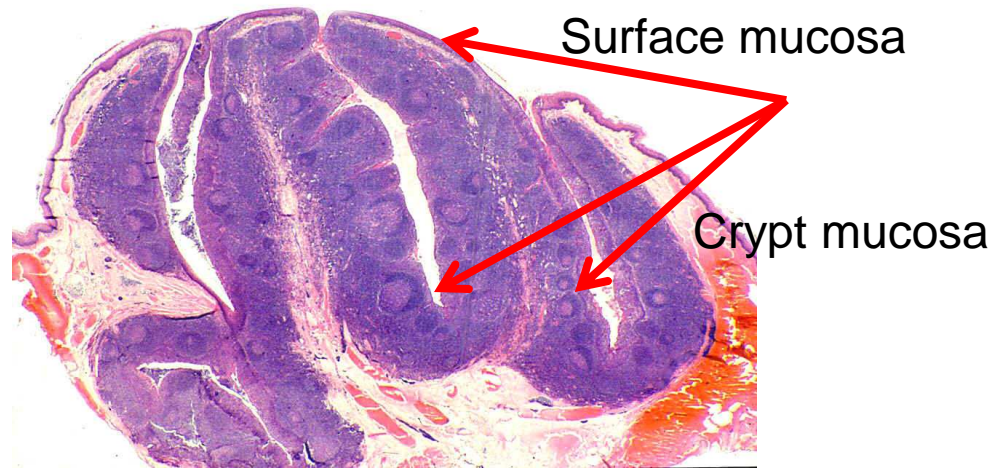
NK cells have multiple receptors



NKp44 is expressed in the human tonsil

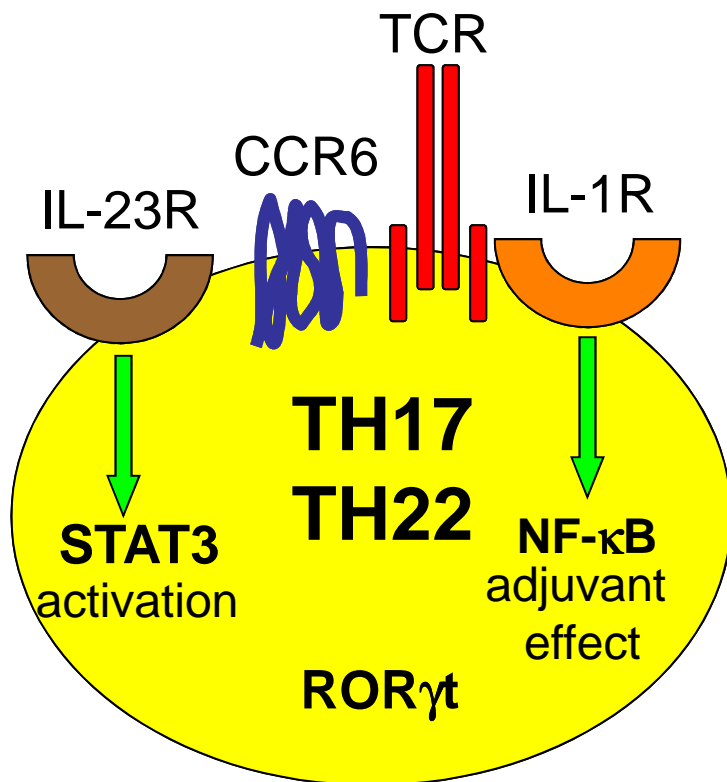


CD56+ NKp44+ cells are found in human oral and gastrointestinal mucosae



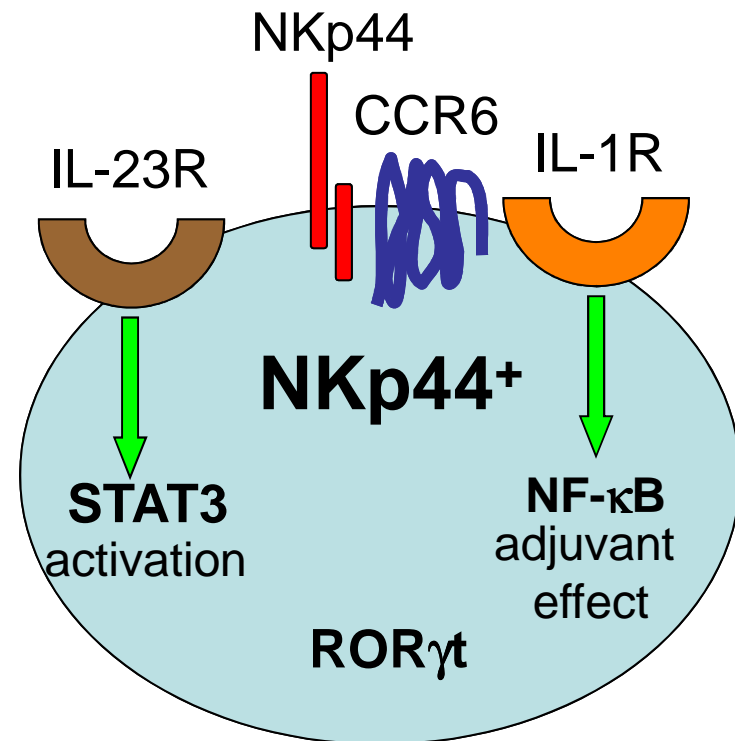
Similarity between human NKp44+ cells and TH17/TH22

Type IL-17/22
CD4 T cells



IL-17 and/or IL-22

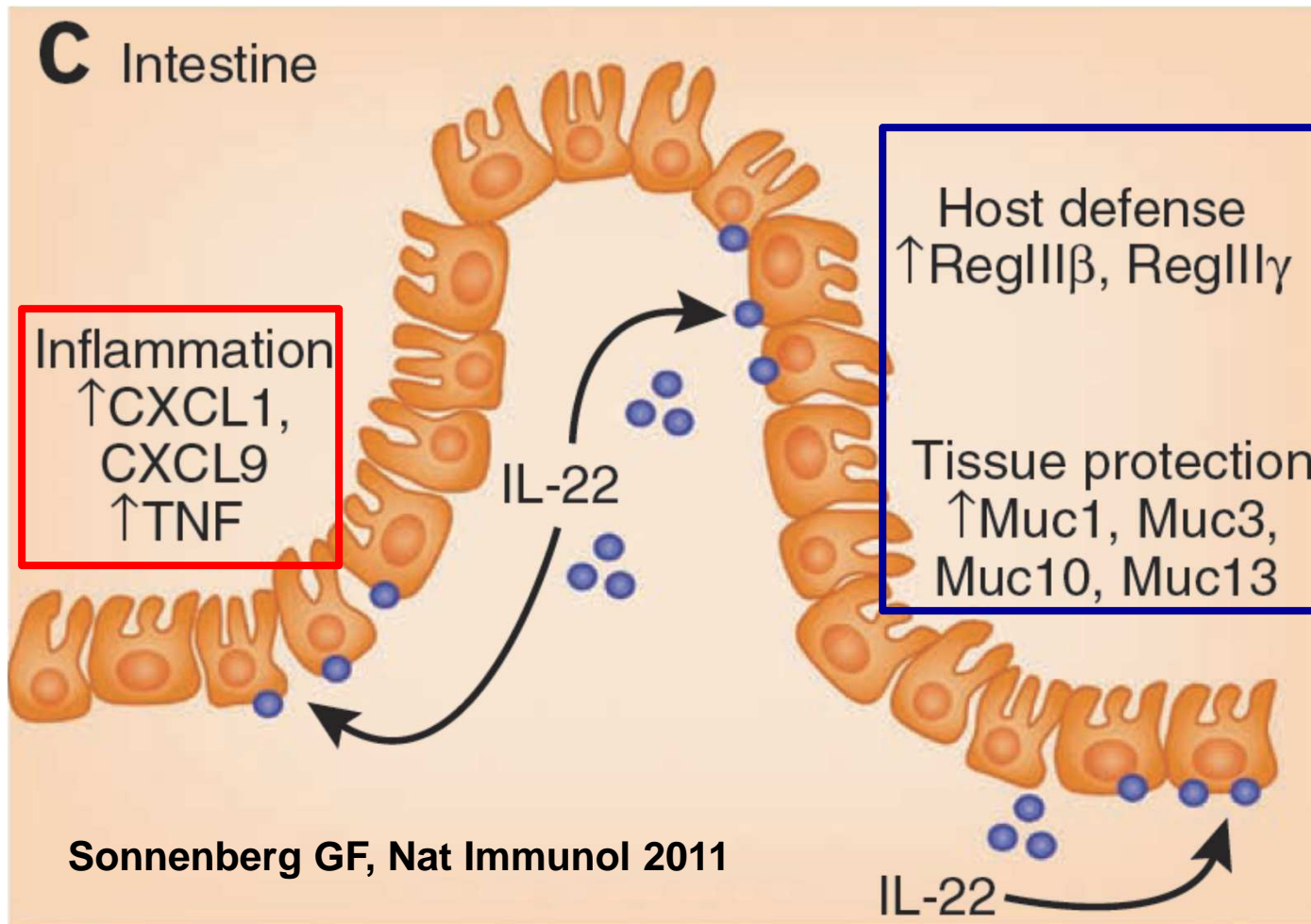
NKp44+ cells
Innate lymphoid cells type 3
(ILC3)



IL-22

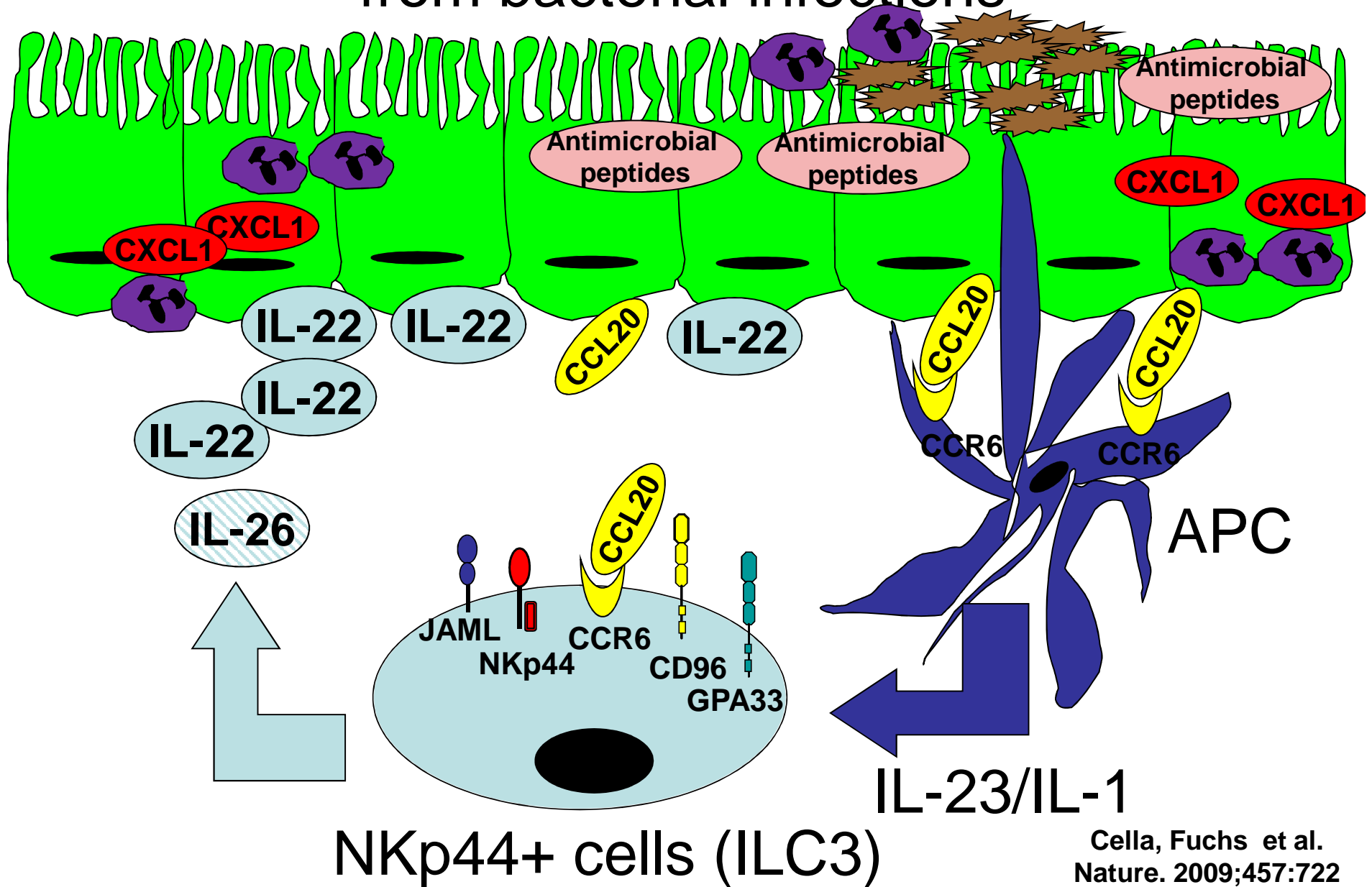
Cella, Fuchs et al.
Nature. 2009;457:722

IL-22 triggers epithelial cell secretion of antimicrobial peptides and chemokines

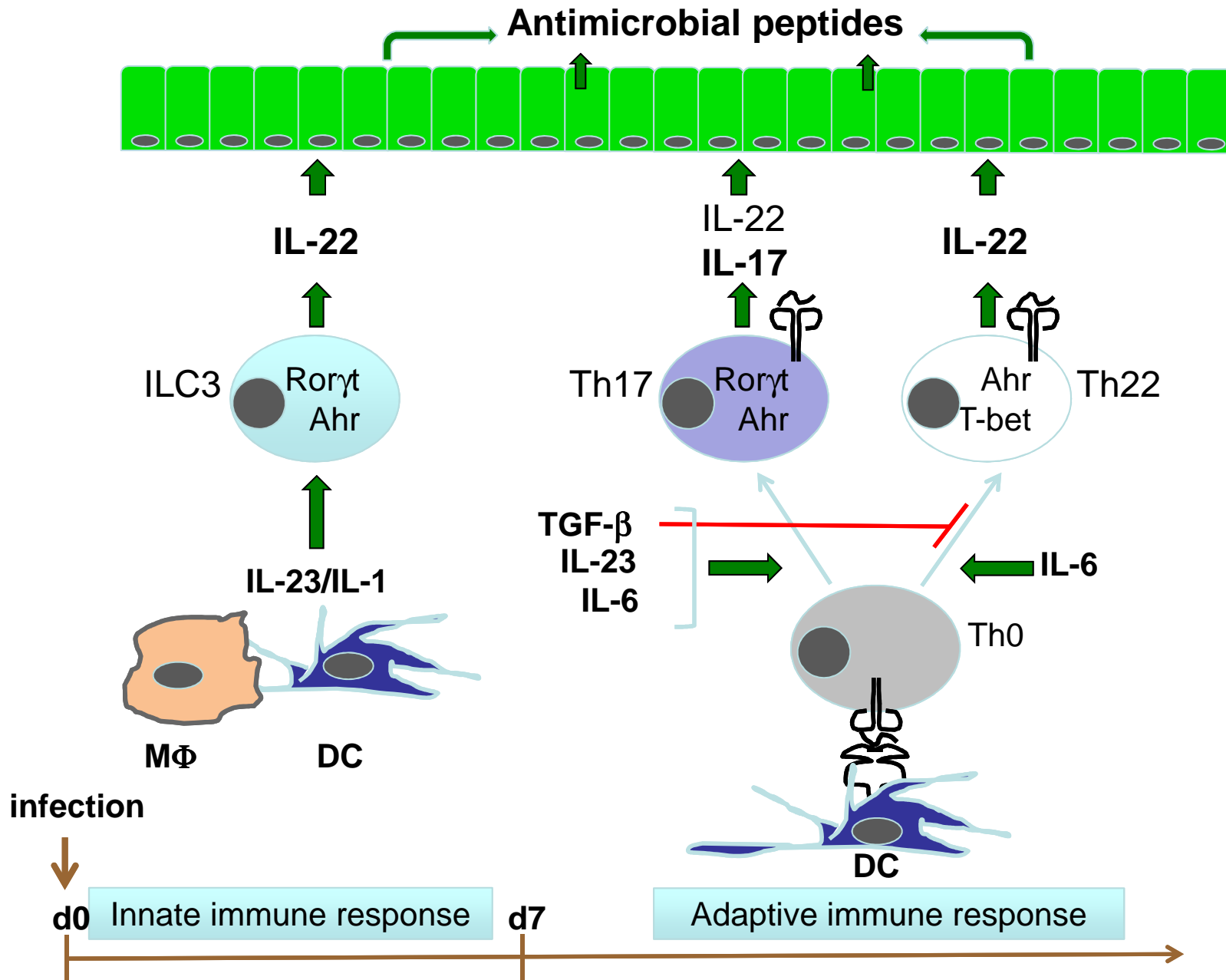


IL-22 protects from bacterial infections but excessive secretion can cause inflammation (psoriasis)

Hypothesis: ILC3 protect the intestinal mucosa from bacterial infections



Cooperation between ILC3 and T cells



Mouse models for understanding the function of ILC3

- **Genetic models:** *Rag*^{-/-} x *γc*^{-/-}
- **Antibody depletion in RAG^{-/-} mice**
 - anti-NK1.1 in *Rag*^{-/-} mice
 - anti-CD90 (anti-Thy1) in *Rag*^{-/-} mice

Use of antibodies in *Rag*^{-/-} mice or *Rag*^{-/-} x *γc*^{-/-} will deplete the cell population in question but will also eliminate other populations as well

Need of lineage-specific deletion mutants

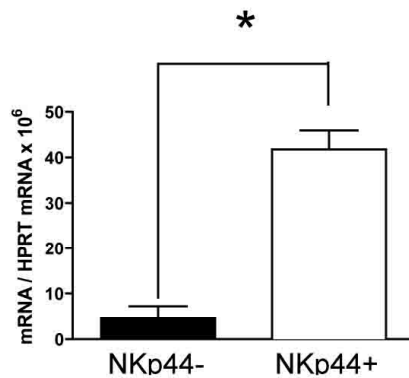
Which transcription factors are required for ILC3 development?

Jacob Lee

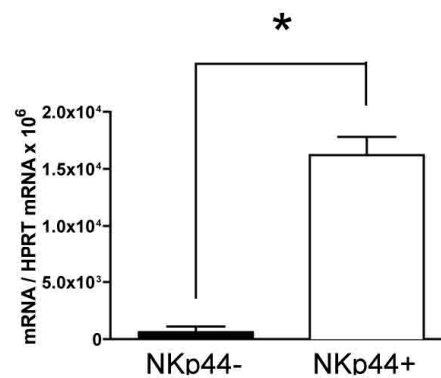
NKp44+ cells express the transcription factor AHR

Retinoic acid-Related Orphan Receptors

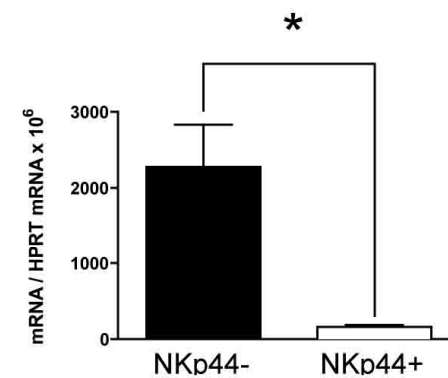
Rora



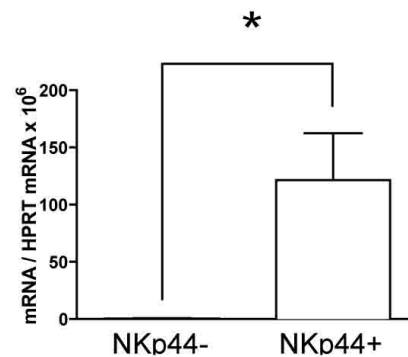
Rorc



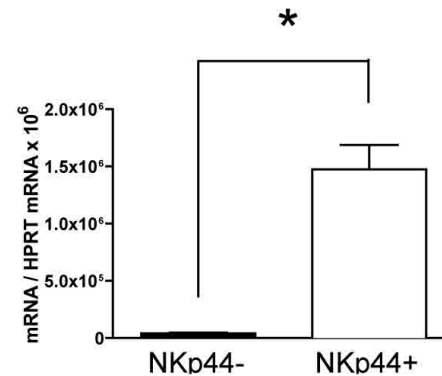
Foxp3



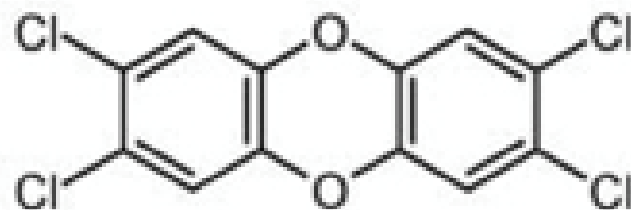
Irf4



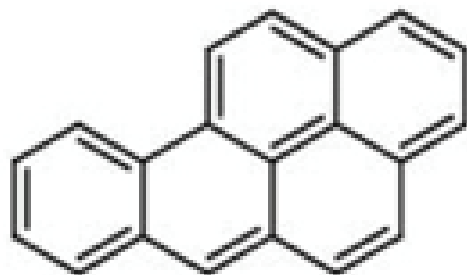
Aryl Hydrocarbon Receptor
Ahr



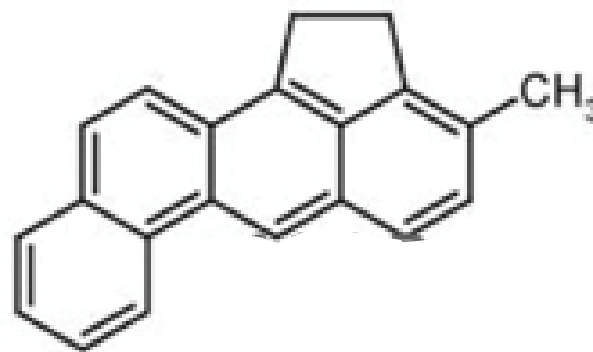
Aryl Hydrocarbon Receptor (AHR): a sensor of polycyclic aromatic hydrocarbons



2,3,7,8-tetrachlorodibenzo-p-dioxin

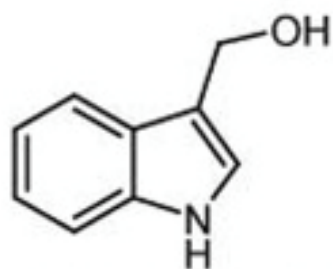


Benzo[a]pyrene



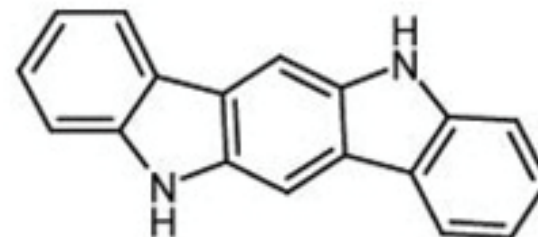
3-methylcholanthrene

AHR ligands include nutritional components



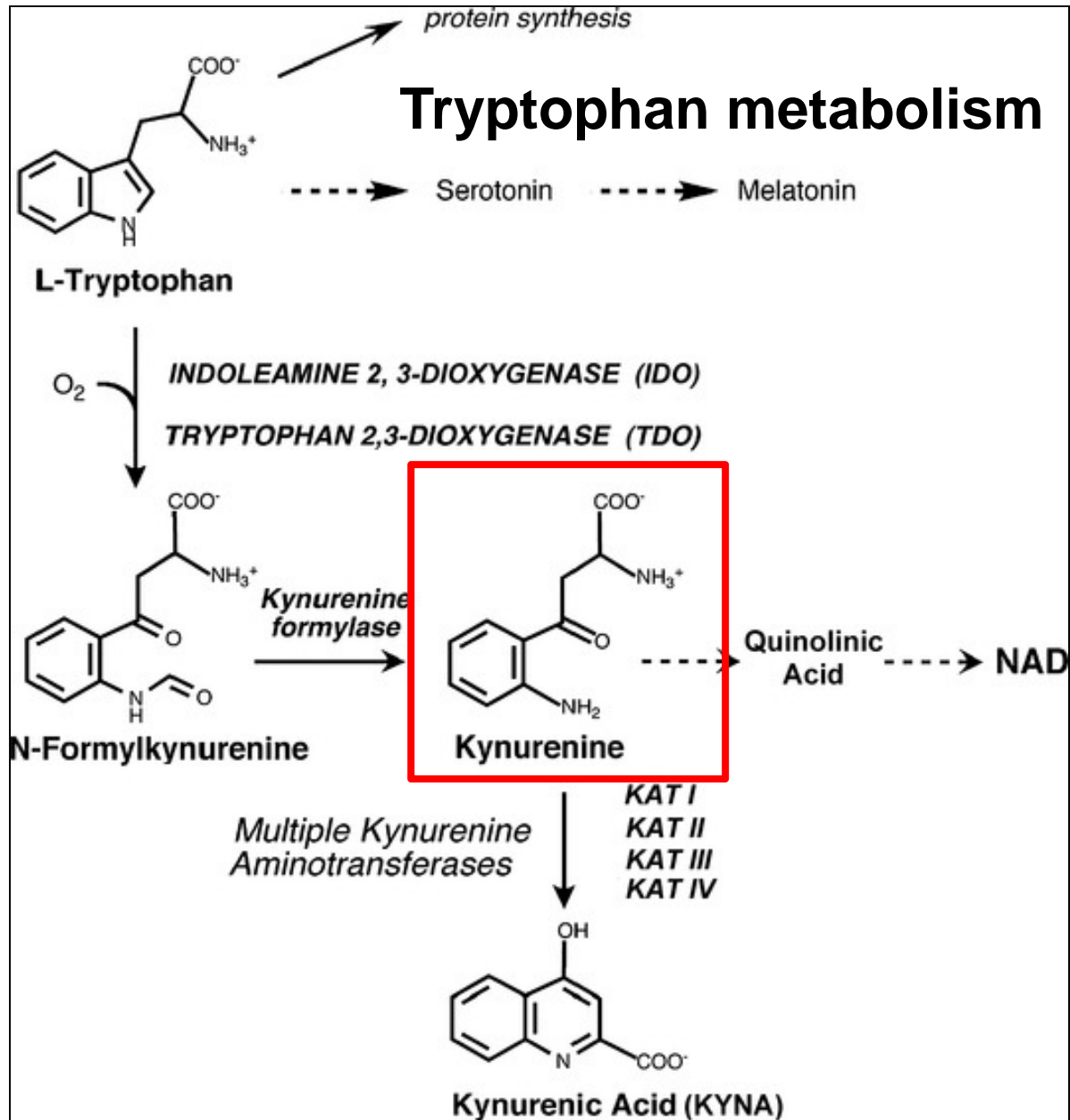
Indole-3-carbinol (I3C)

Acidic pH



Indolo[3,2-*b*]carbazole (ICZ)

Endogenous AHR ligands

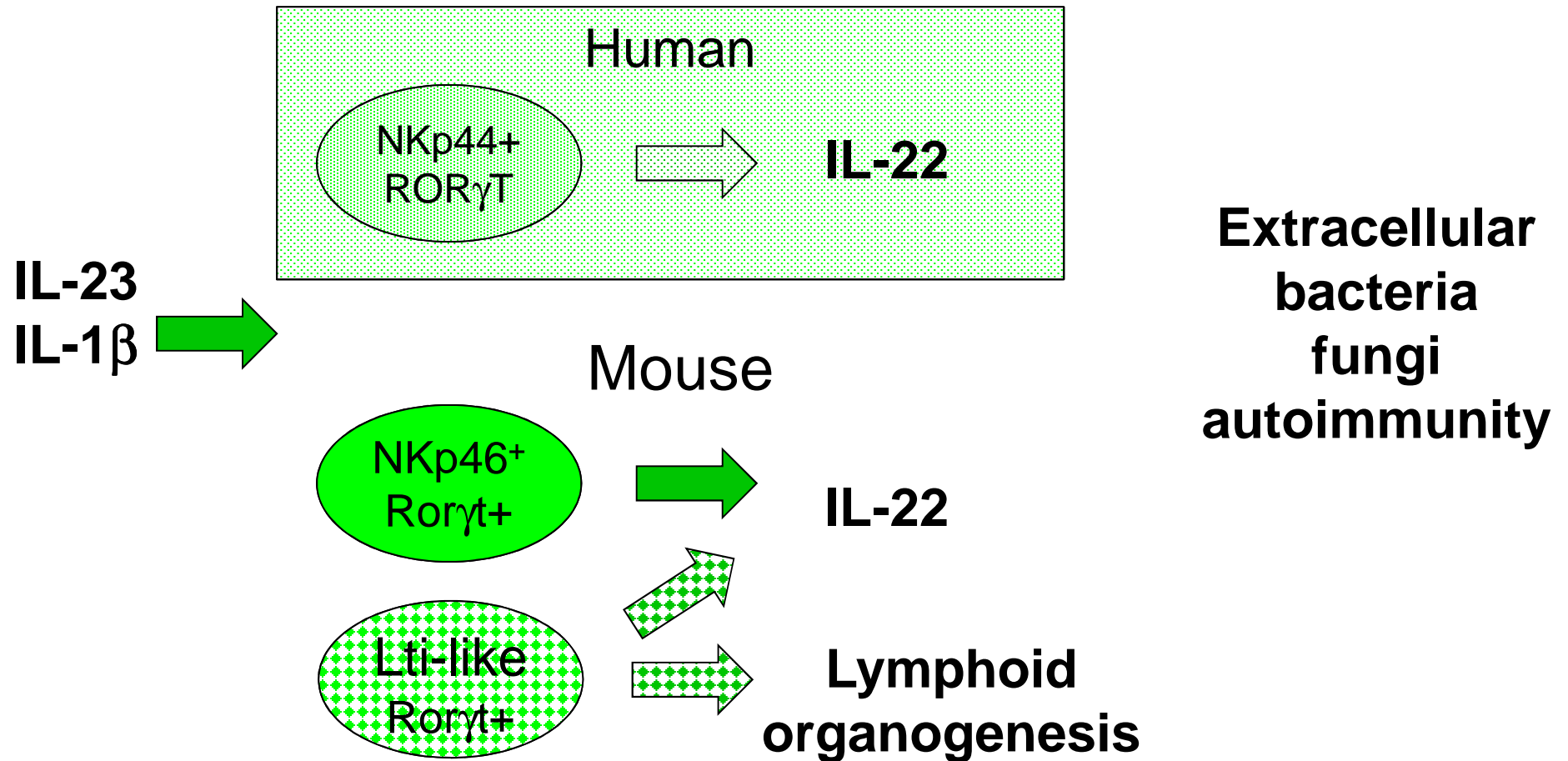


Can we target AHR to block the development of ILC3?

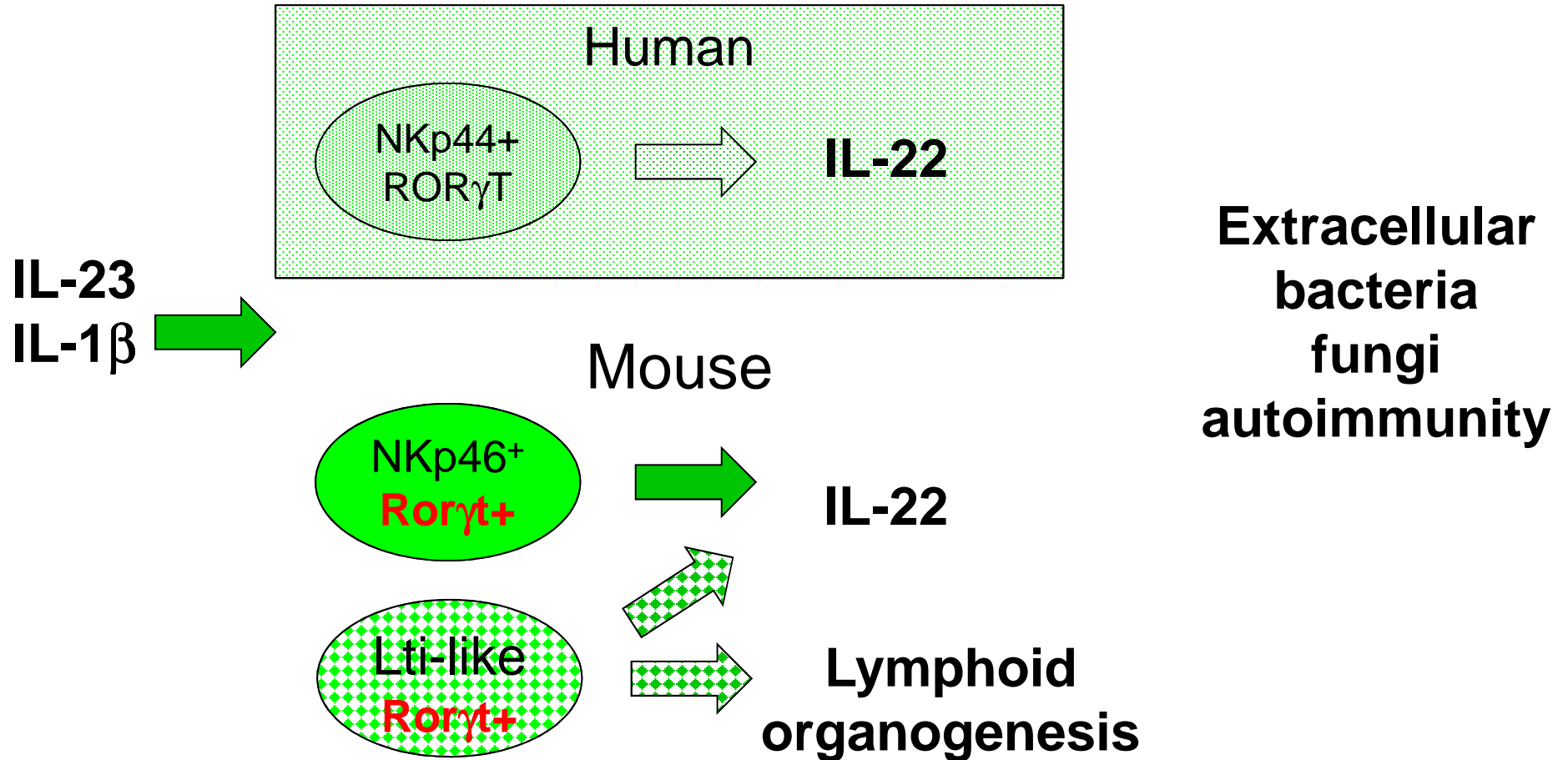
Analysis of conditional *Ahr*^{-/-} mice

Chris Bradfield

ILC3 subsets in mouse small intestine

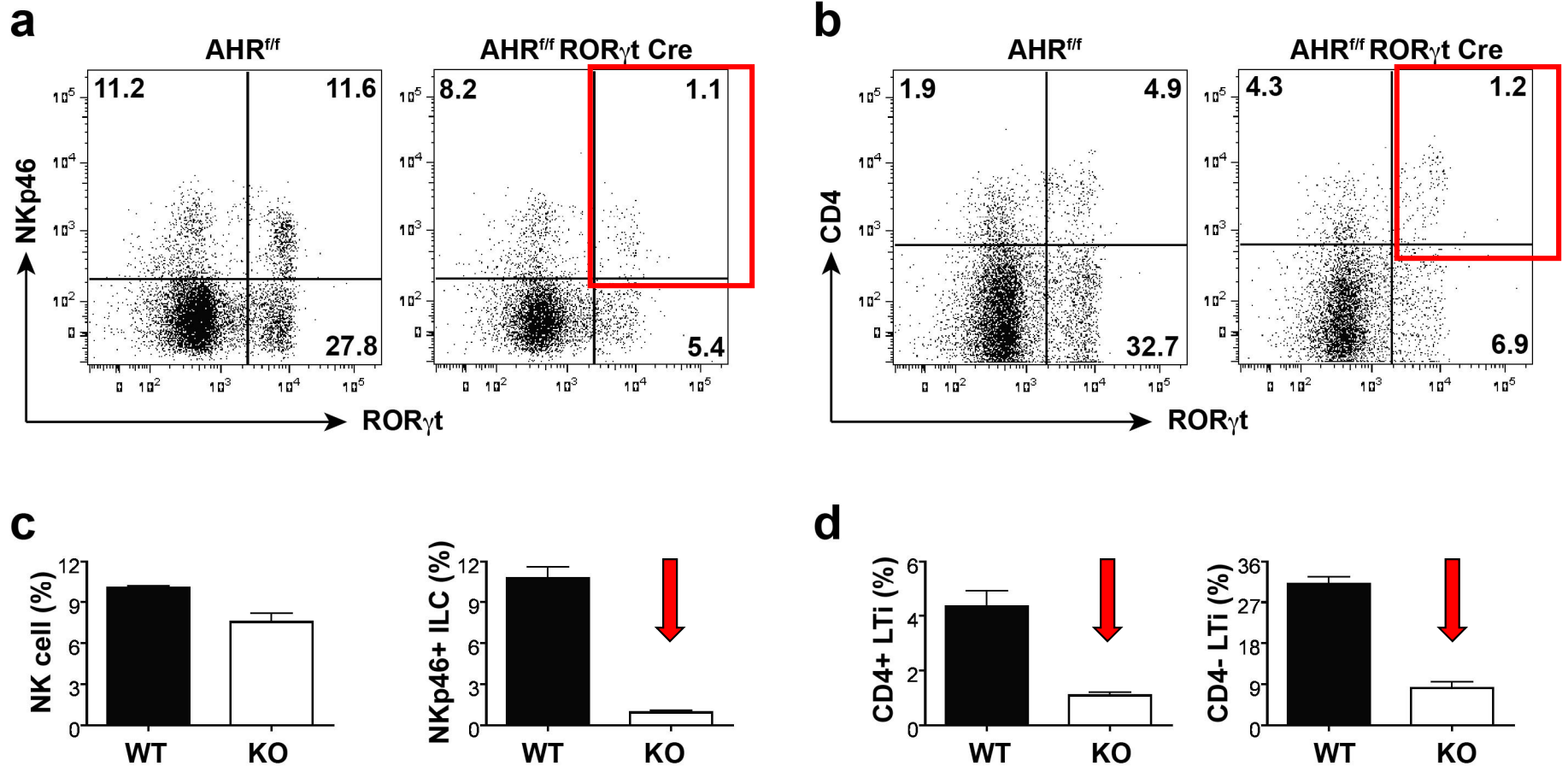


ILC3 subsets in mouse small intestine

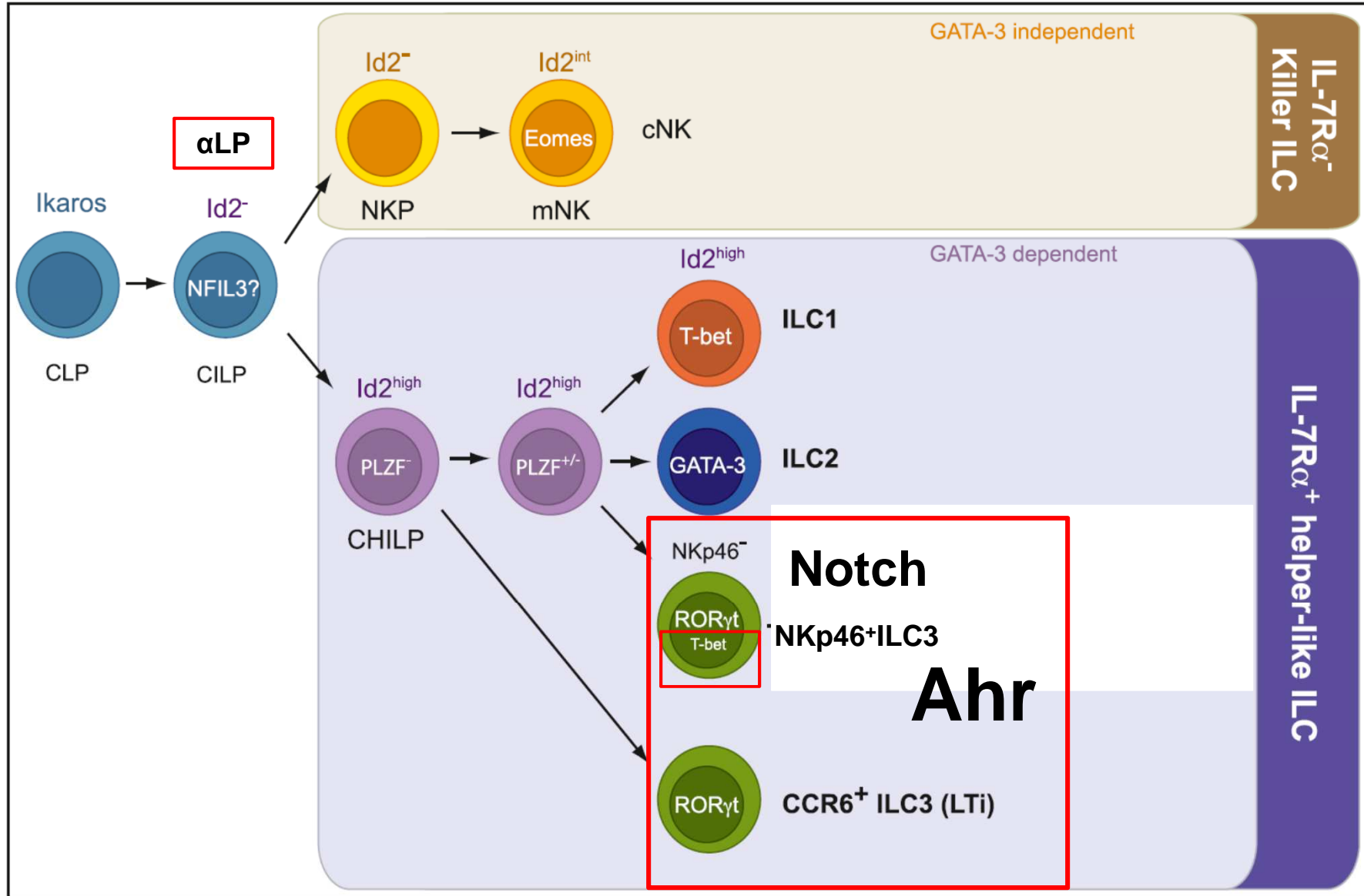


Conditional *Ahr*-deficient mice

Ahr^{fl/fl} x *Royt-cre*



AHR is required for ILC3 development



ILC3-deficient mice

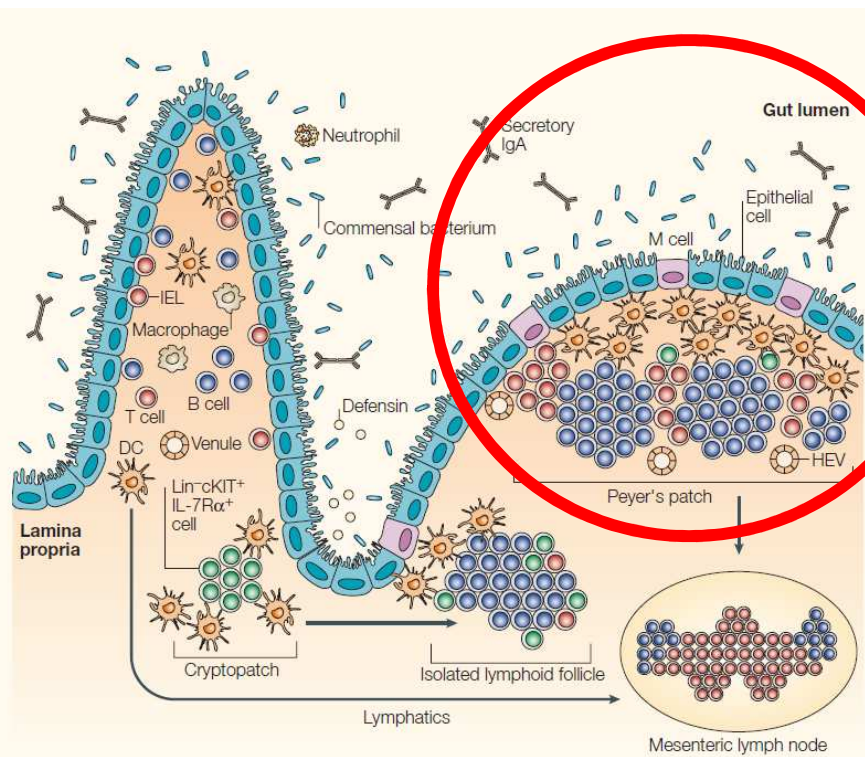
Mice	Affected cells
AHR ^{f/f} x ROR γ t-Cre	ILC3
AHR ^{f/f} x ROR γ t-Cre x TCR $\beta\delta$ ^{-/-}	ILC3, T cells

**Are ILC3 important for
intestinal homeostasis?**

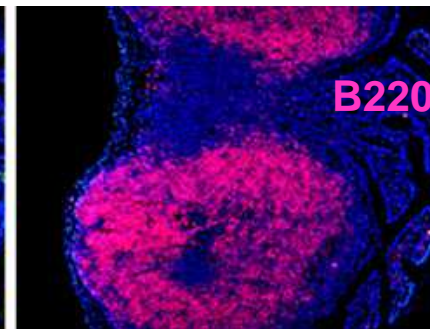
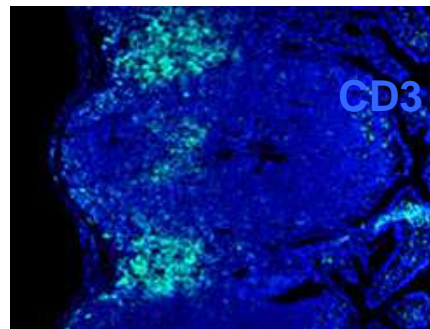
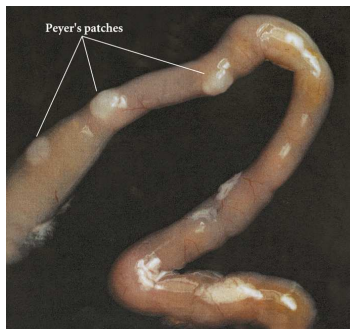
Intestinal lymphoid tissues

Peyer's Patches

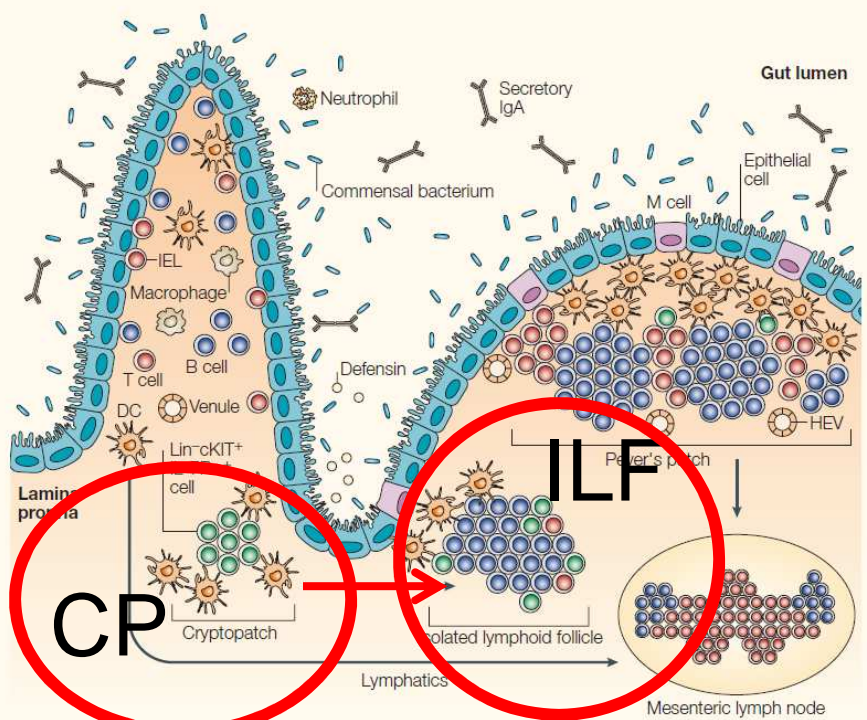
Develop pre-natally



Eberl, Nat Rev Immunol 2005

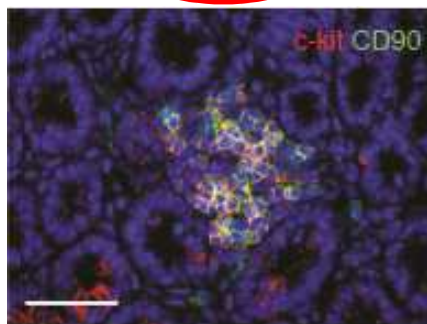


Intestinal lymphoid tissues



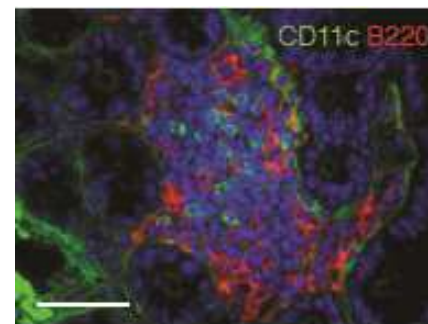
Develop post-natally

Eberl, Nat Rev Immunol 2005



cryptopatches

CD90+(Thy1+) c-Kit+ Lti-ILC

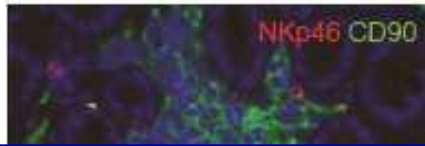


Isolated lymphoid follicles

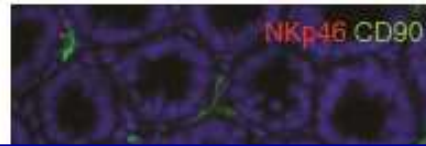
B220+/CD11c+

ILC3-deficient mice lack cryptopatches and isolated lymphoid follicles

WT

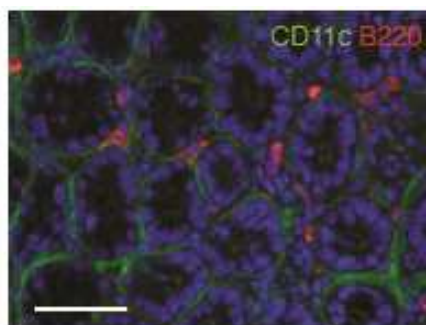
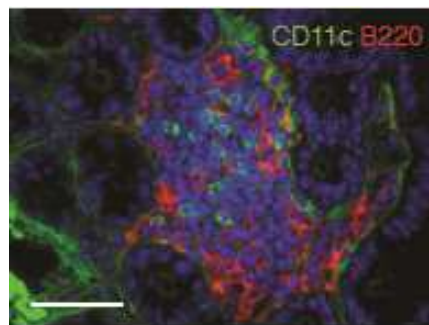


ILC3-deficient



- CD90+ (Thy1+) clusters identifying CP are absent in ILC3-deficient mice

Prenatal lymphoid tissues (Peyer's Patches) are conserved



- CD11c+B220+ clusters identifying ILFs are absent in Ahr-deficient mice

Rodney Newberry
Keely McDonald

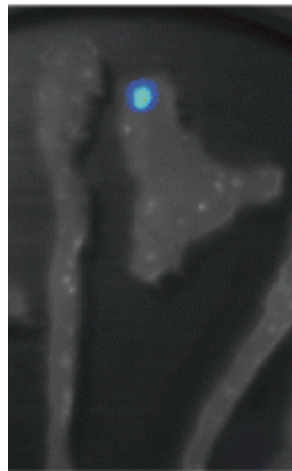
Are ILC3 important in host response to pathogenic bacteria?

Christina Song

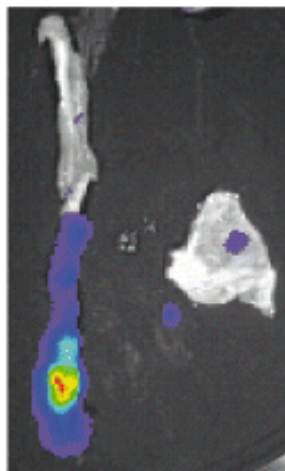
Citrobacter rodentium infection

- *C. rodentium* is a natural mouse pathogen related to enteropathogenic *E. coli*
- The infection is restricted to the lumen of the intestine
- In B6 mice, *C. rodentium* infection initiates on the surface of caecal patches, peaks at d7-14 pi and is cleared by d21-28 pi.

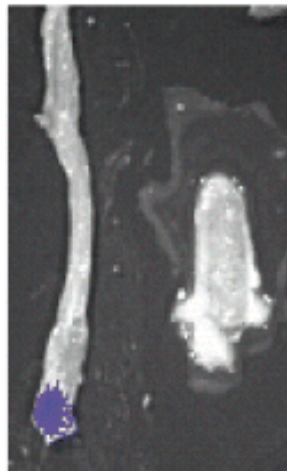
Oral gavage of LB grown *C. rodentium*



Day 3



Day 10

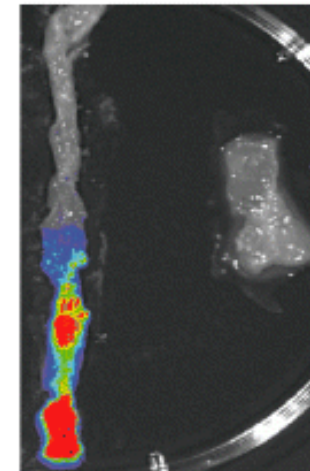


Day 17



Day 21

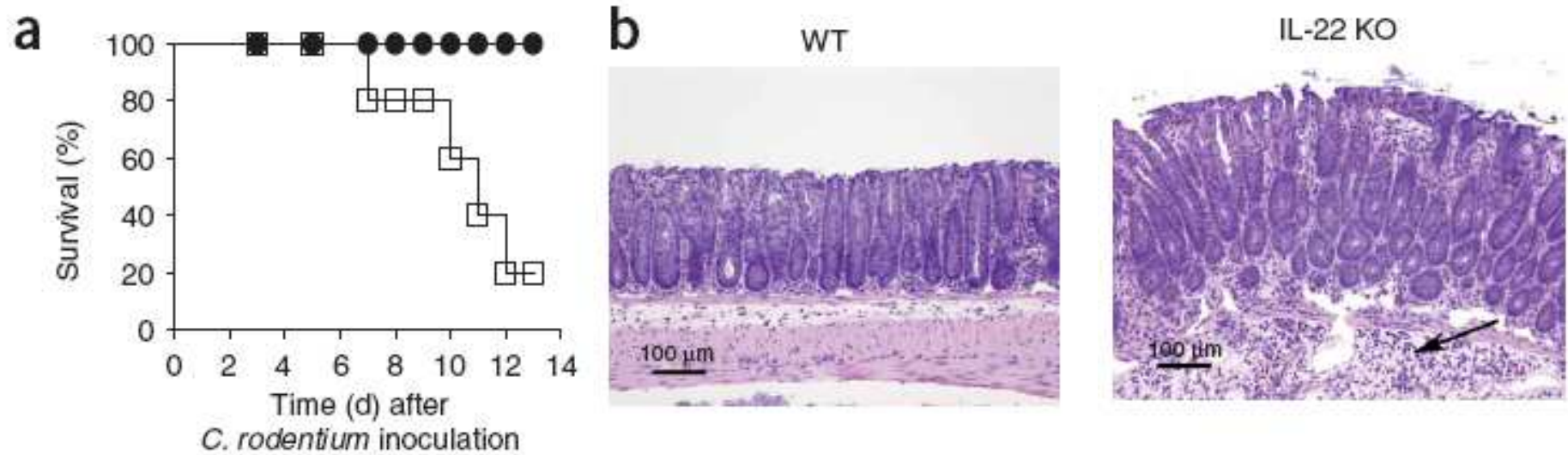
Natural transmission



Day 3

Mundy R et al. Cell Microbiol. 2005;7:1697

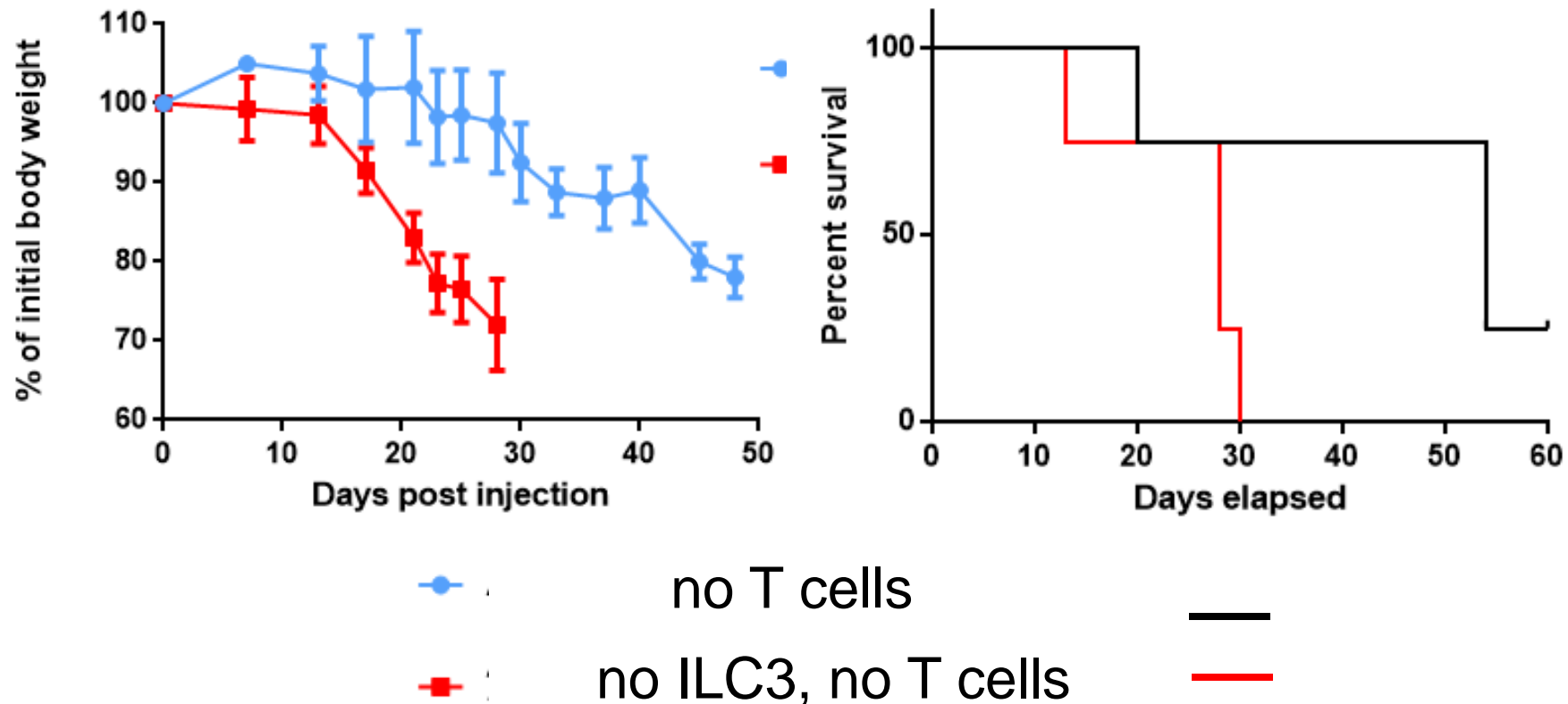
Host defense against *C. rodentium* infection depends on IL-22



Zheng Y, et al. Nat Med. 2008;14:282.

C. rodentium infection in the absence of ILC3 and T cells

AHR^{f/f} x ROR γ t-cre x TCR β / δ ko (no ILC3, no T cells) vs AHR^{f/f} x TCR β / δ -/- (no T cells)



C. rodentium infection in mice lacking ILC3 only

AHR^{f/f} x ROR γ t-cre (no ILC3) vs
AHR^{f/f}

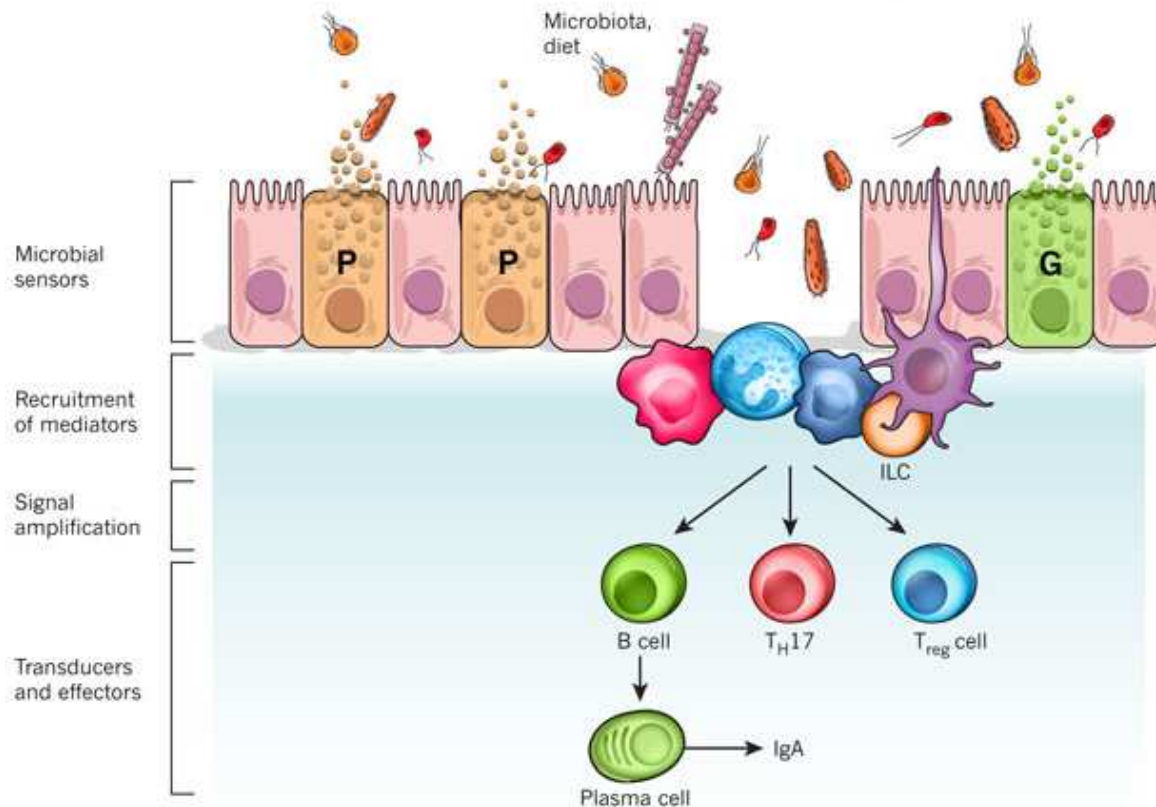
ILC3 protect the gastrointestinal tract
from bacterial infections
Partial redundancy with T cells

60
0 10 20 30 0 10 20 30
Days post infection Days elapsed

● ctr —

■ no ILC3 —

Do ILC3 contribute to IBD?



IBD-related processes

Epithelial barrier

*GNA12**, *HNFA4*, *CDH1*, *ERRFI1*, *MUC19*, *ITLN1**

Restitution

REL, *PTGER4*, *NKX2-3*, *STAT3*, *ERRFI1*, *HNFA4*, *PLA2G7E*

Solute transport

SLC9A4, *SLC22A5*, *SLC22A4**, *AQP12A/B*, *SLC9A3*, *SLC26A3*

Paneth cells

*ITLN1**, *NOD2**, *ATG16L1**, *XBP1**

Innate mucosal defence

*NOD2**, *ITLN1**, *CARD9**, *REL*, *SLC11A1*, *FCGR2A*/B*

Immune cell recruitment

CCL11/CCL2/CCL7/CCL8, *CCR6*, *IL8RA/IL8RB*, *MST1**

Antigen presentation

*ERAP2**, *LNPEP*, *DENND1B*

IL-23/TH17

*IL23R**, *JAK2*, *TYK2**, *STAT3*, *ICOSLG*, *IL21*, *TNFSF15*

T-cell regulation

NDIFP1, *TNFSF8*, *TAGAP*, *IL2*, *IL2RA*, *TNFRSF9*, *PIM3*, *IL7R**, *IL12B*, *IL23R**, *PRDM1*, *ICOSLG*, *TNFSF8*, *IFNG*, *IL21*

B-cell regulation

IL5, *IKZF1*, *BACH2*, *IL7R**, *IRF5*

Immune tolerance

IL10, *IL27**, *SBNO2*, *CREM*, *IL1R1/IL1R2*, *NOD2**

IL-23R
STAT3

Cellular responses

Autophagy

*ATG16L1**, *IRGM*, *NOD2**, *LRRK2*, *CUL2*, *PARK7*, *DAP*

ER stress

CPEB4, *ORMDL3*, *SERINC3*, *XBP1**

Intracellular logistics

VAMP3, *KIF21B*, *TTL8*, *FGFR10P*, *CEP72*, *TPPP*

Cell migration

ARPC2, *LSP1*, *AAMP*

Apoptosis/necroptosis

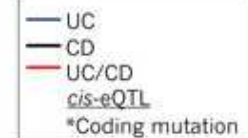
FASLG, *THADA**, *DAP*, *PUS10*, *MST1**

Carbohydrate metabolism

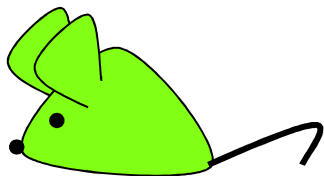
*GCKR**, *SLC2A4RG*

Oxidative stress

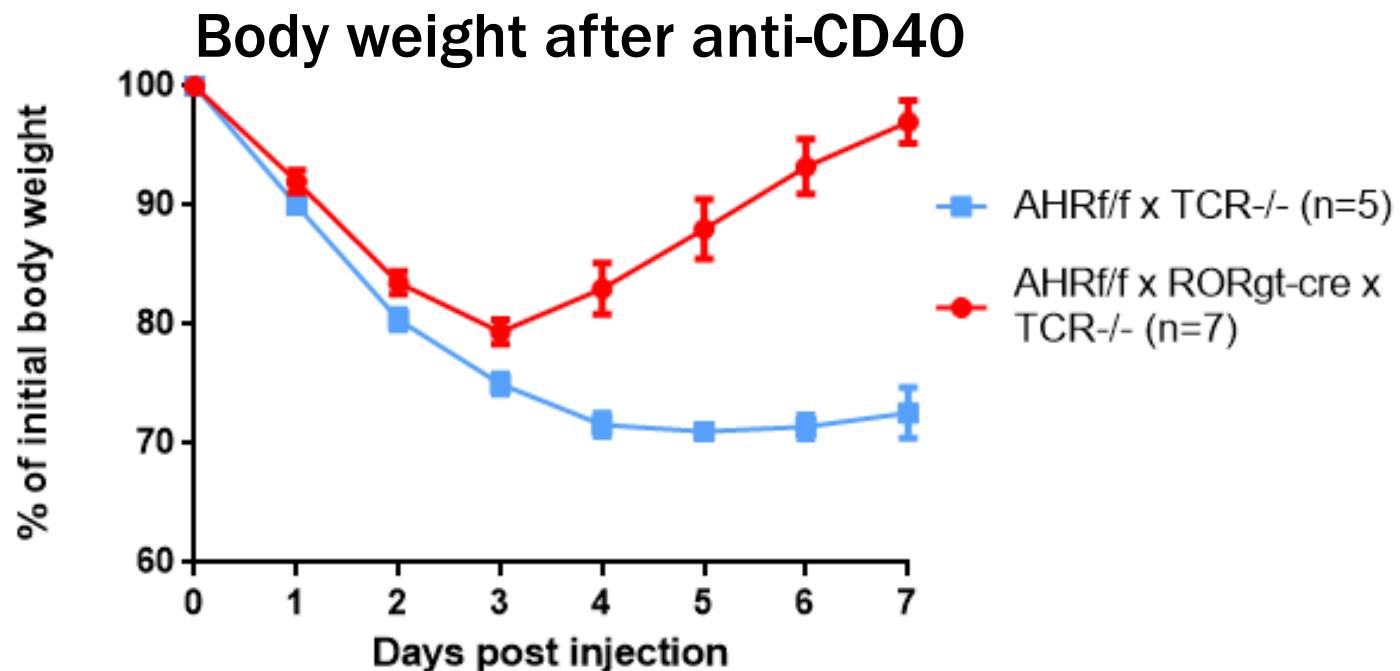
PRDX5, *BACH2*, *ADO*, *GPX4*, *GPX1**, *SLC22A4*, *LRRK2*, *NOD2**, *CARD9**, *HSPA6*, *DLD*, *PARK7*, *UTS2**, *PEX13*



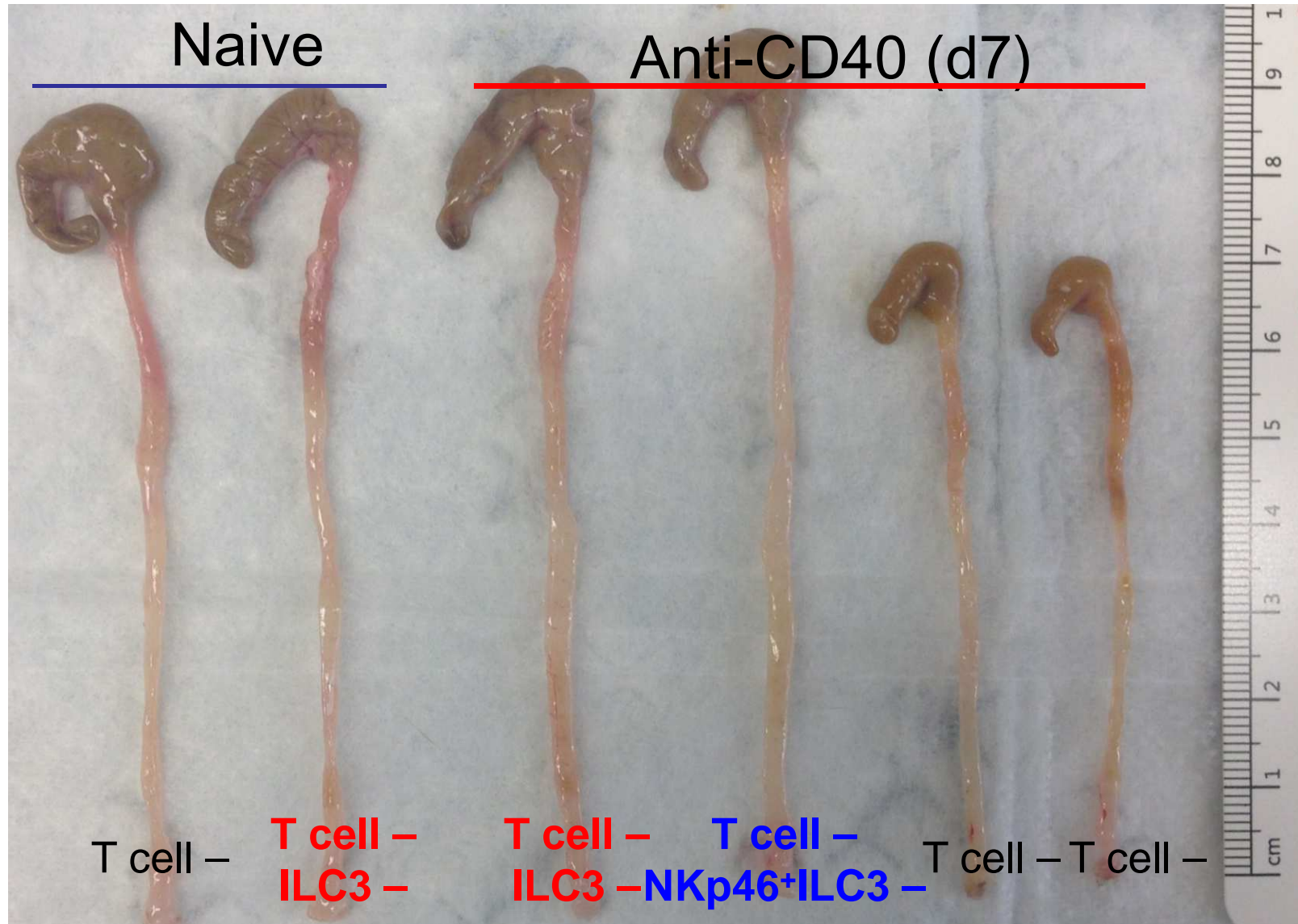
ILC3 worsen anti-CD40-induced colitis



AHR^{f/f} x ROR γ t-cre x TCR β/δ ko (**no ILC3, no T cells**) vs
AHR^{f/f} x TCR β/δ -/- (**no T cells**)



Colon length 7 days after anti-CD40



T cell – **T cell –** **T cell –** **T cell –** T cell – T cell –
 ILC3 – **ILC3 –** **NKp46+ILC3 –**

Ctrl
(Ahrf/f x
TCRbd/-)

ROR γ t-cre x
Ahrf/f x
TCRbd/-

ROR γ t-cre x
Ahrf/f x
TCRbd/-

ROR γ tf/f x
NKp46-cre x
TCRbd/-

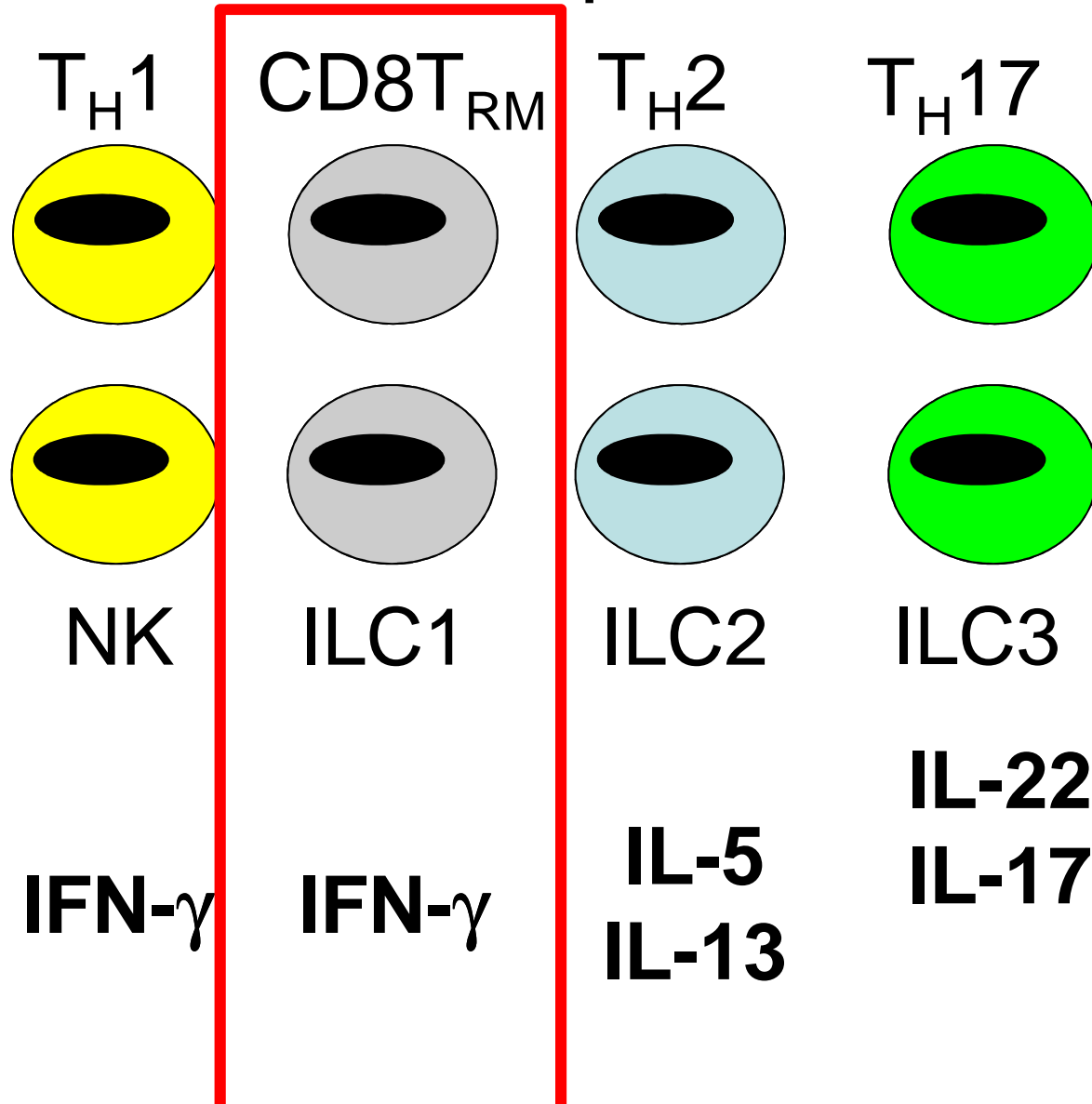
Ctrl
(Ahrf/f x
TCRbd/-)

Cre Ctrl
(ROR γ t-cre+ x
TCRbd/-)

ILC3: two-edged sword

- Induce the development of ILF but have no obvious role in containment of commensal microbiota
- Protect from *C. rodentium* infections together with T cells
- Promote inflammation in a model of IBD

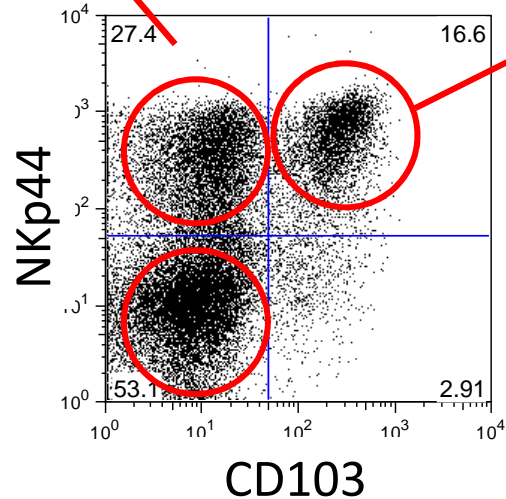
Similarity of functional modules in innate and adaptive immunity



Human tonsils harbor 2 major subsets of NKp44⁺ ILCs

Produce IL-22 upon stimulation with IL-23

NKp44⁺ ILC3



Localized within tonsil surface and crypt epithelium

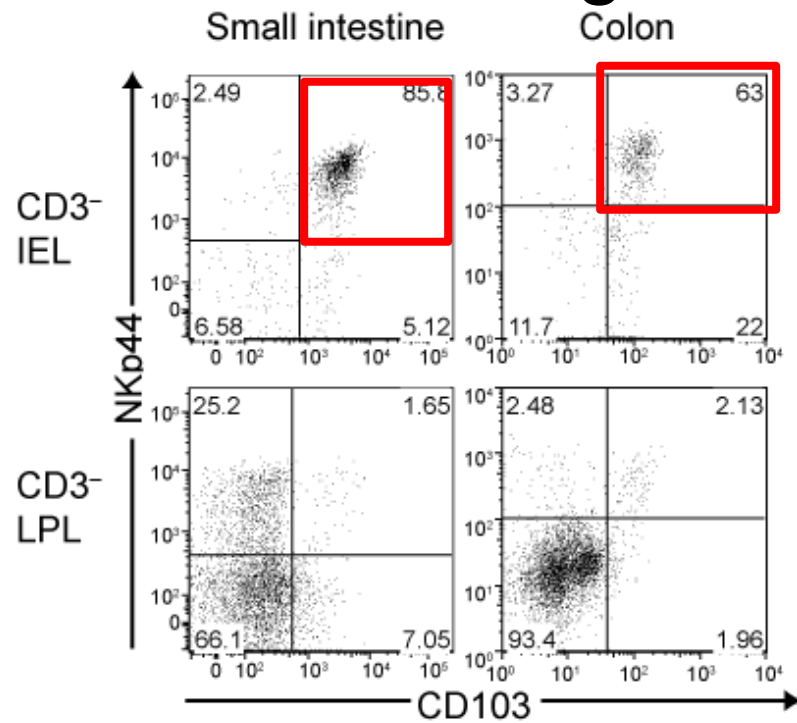
Secrete IFN- γ upon stimulation with IL-12 + IL-15

Limited cytolytic capacity

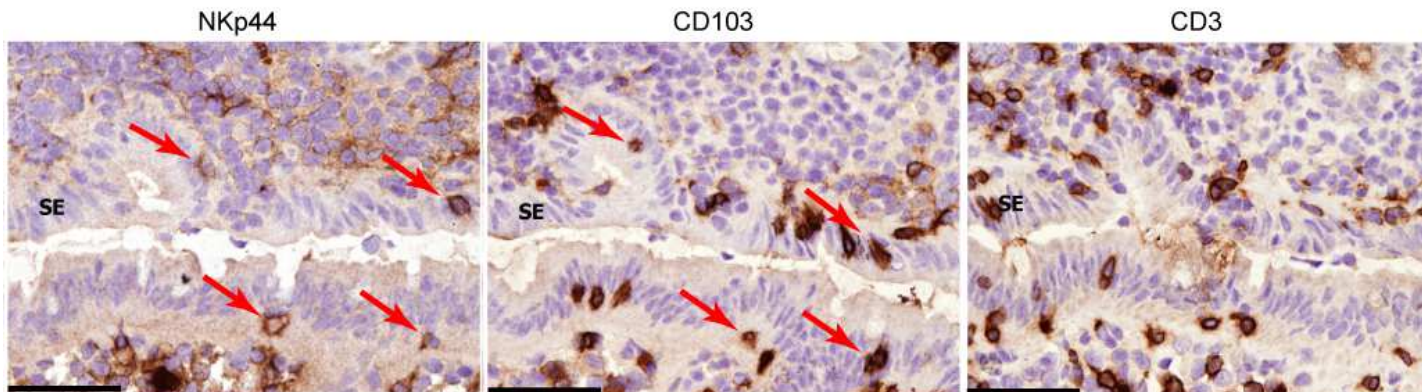
Gated on CD56+

Anja Fuchs

Intraepithelial CD103⁺ ILC1 are present in human gastrointestinal tract

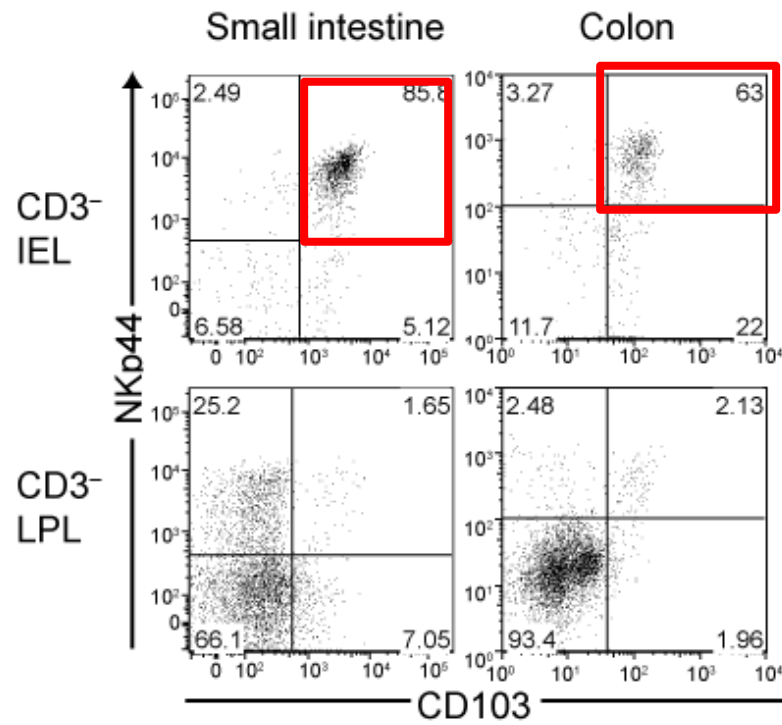


Gated on CD45⁺CD3⁻ lymphocytes

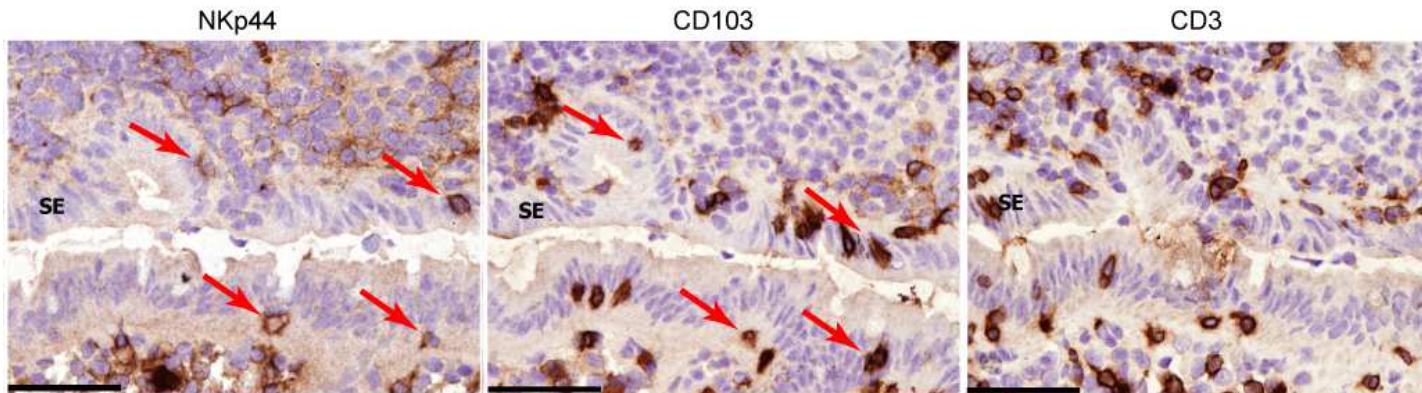
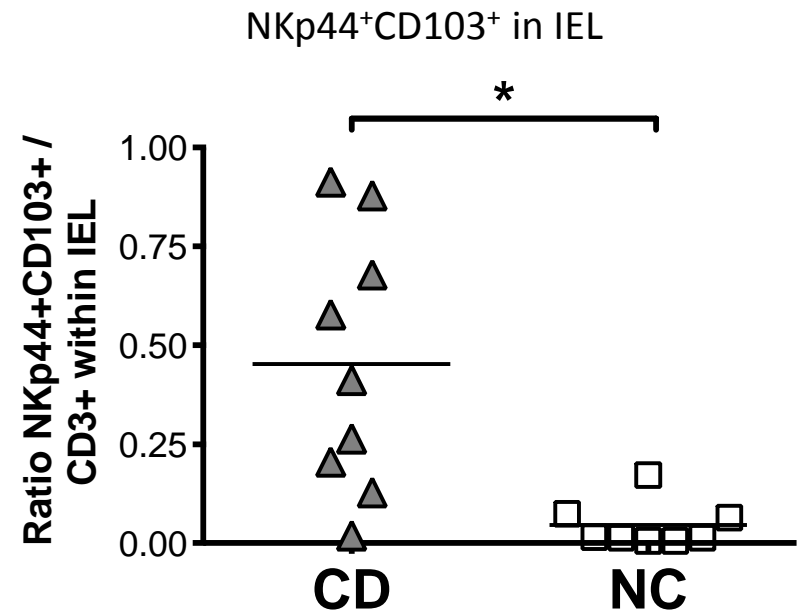


Human appendix

Increased CD103⁺ ILC1 frequencies in the ileum of individuals with Crohn's disease

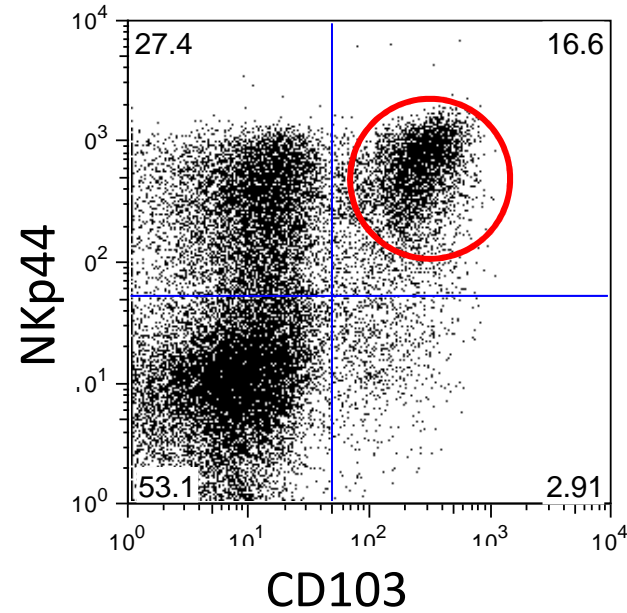


Gated on CD45⁺CD3⁻ lymphocytes

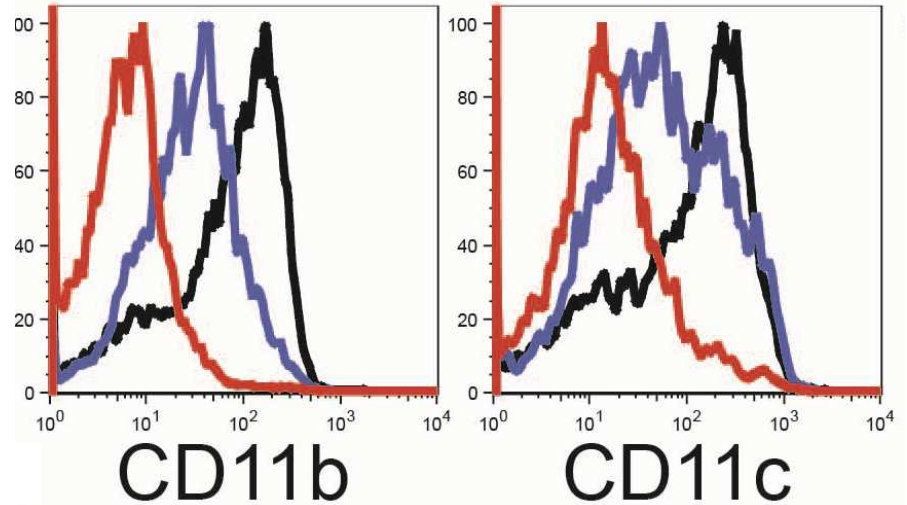
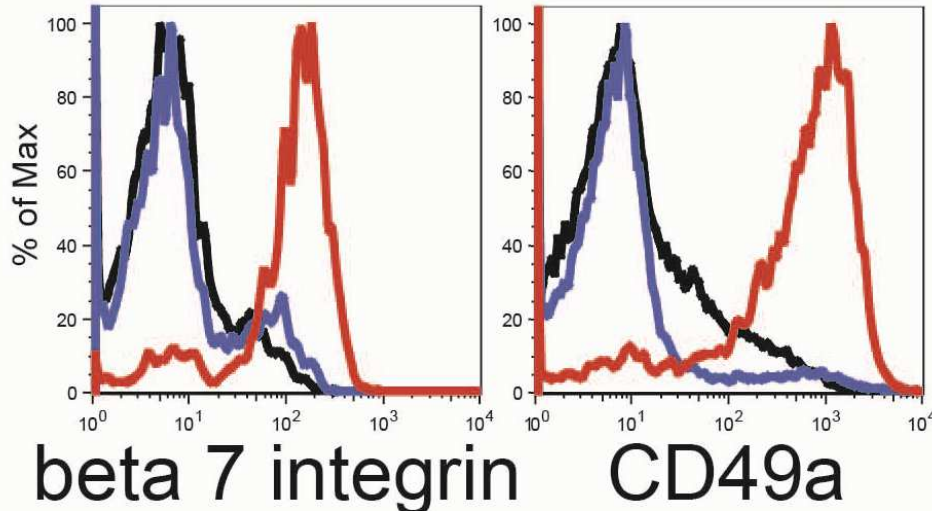





Human appendix

NK cells or distinct ILC subset?

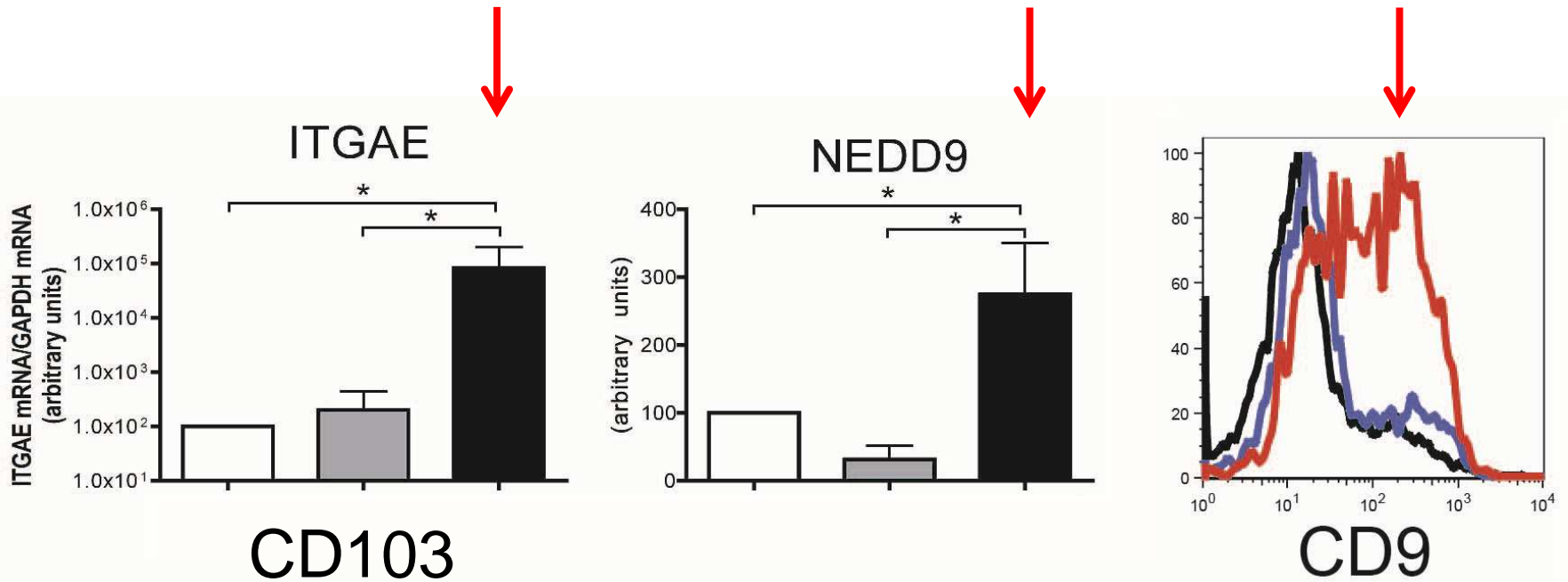


Unique integrin pattern (VLA1+, CD103/Beta7+)



-  NKp44+CD103+
-  ILC3
-  NKp44- (conventional NK)

Unique features: TGF-beta imprinting



- NKp44+CD103+
- ILC3
- NKp44- (conventional NK)

Similarity with Tissue Resident Memory CD8 T cells (T_{RM})

Memory



RECIRCULATE

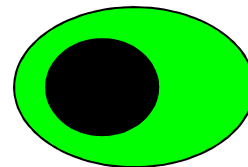
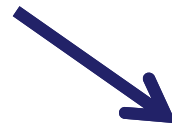
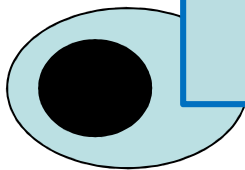
T_{EM}

Peripheral

Naive

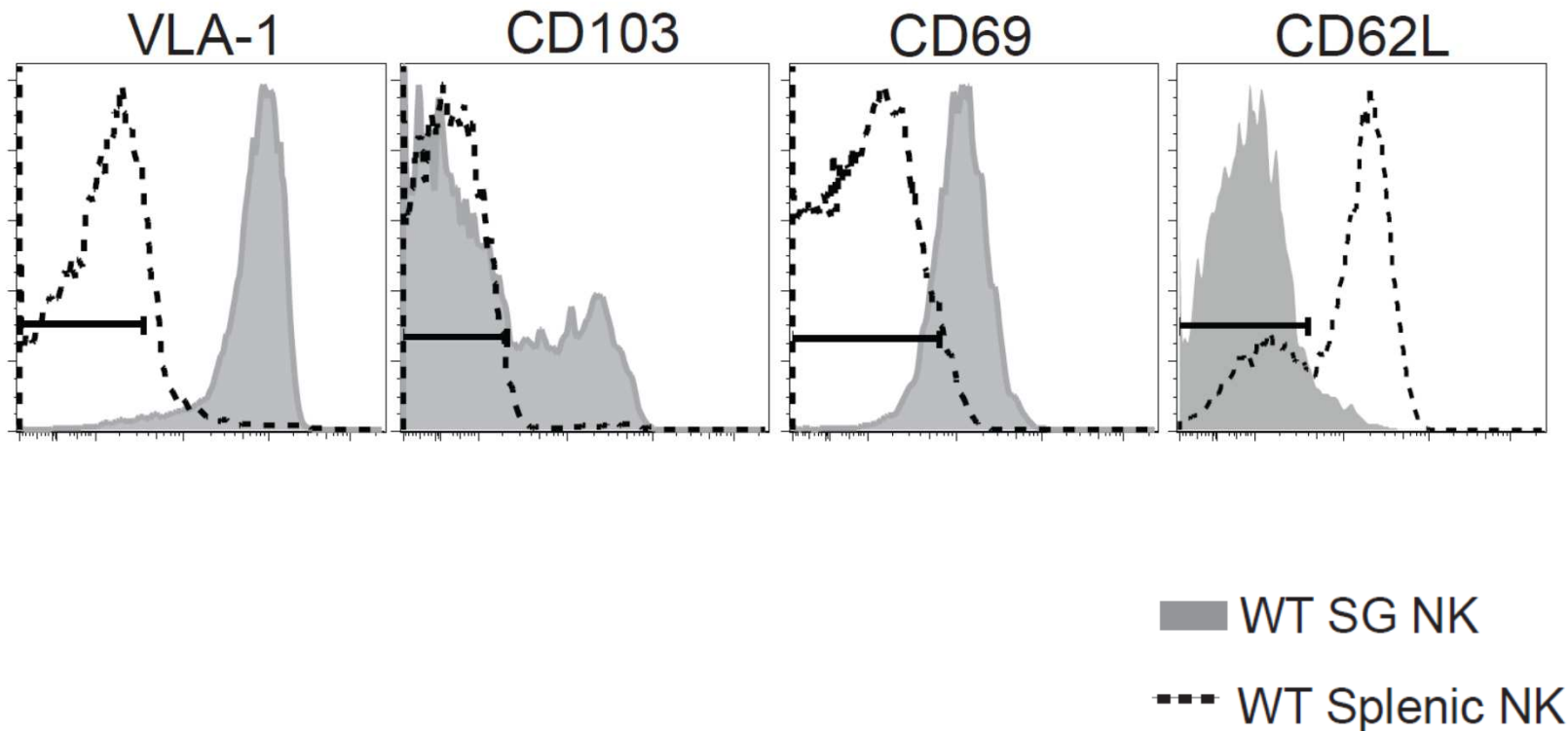
Intraepithelial NKp44+CD103+ are
tissue-resident ILC1

Secondary
Lymph
Nodes



T_{RM}
**Retained in
peripheral tissues**

Salivary gland NK cells are very similar to intestinal intraepithelial ILC1



Victor Cortez

NK cells vs ILC1

Recirculating

Tissue resident



NK

BM, spleen, LN,
siLP



IEL
ILC1

Intestinal
epithelium



uterus

also SG

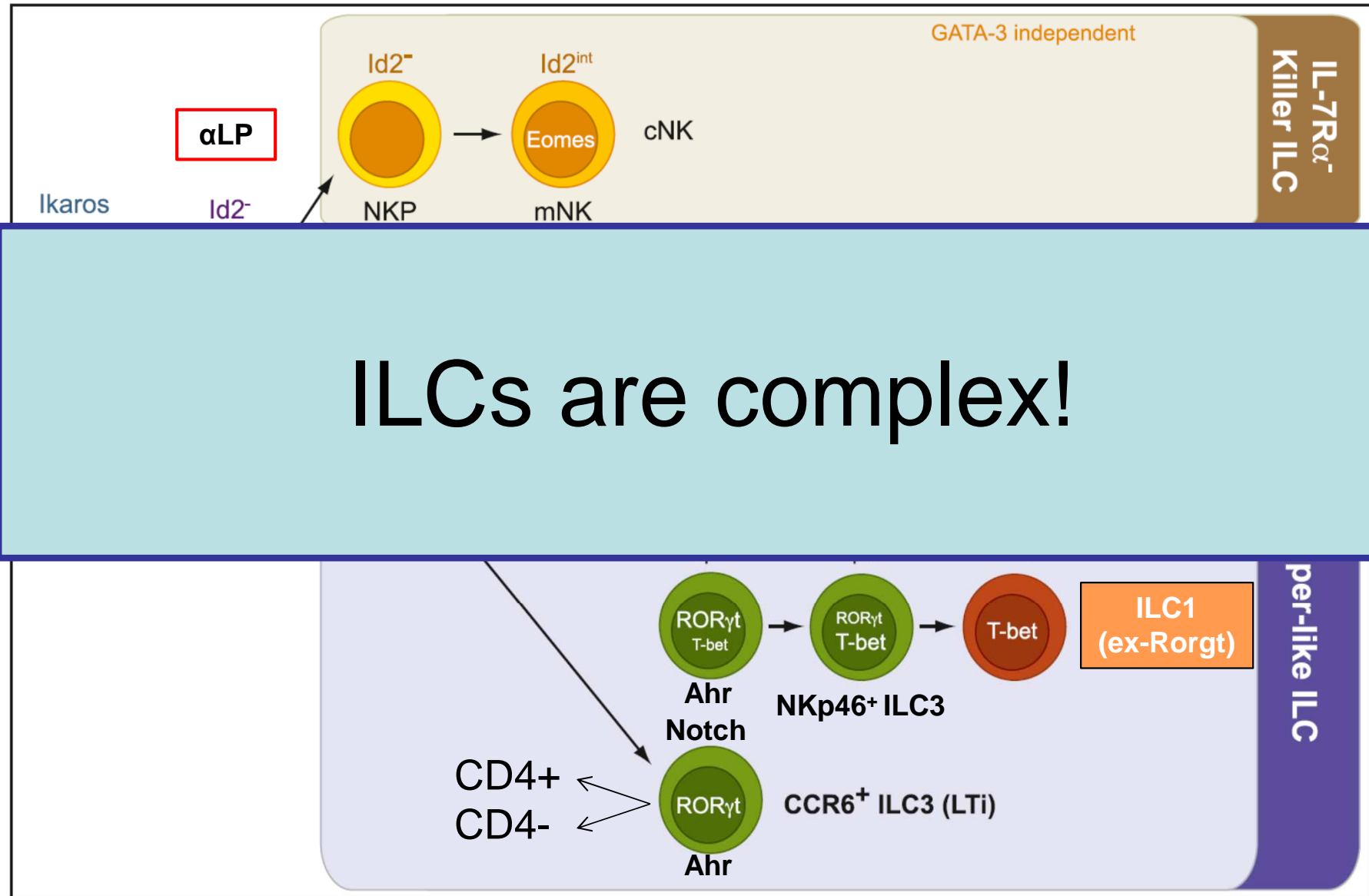


Liver
ILC1

also siLP
and skin

VLA2	positive	negative	positive*	negative
VLA1	negative	positive	positive	positive
TRAIL	negative	positive	positive	positive
CD127	negative	dim	negative	positive*
IL-15	Dependent	Partially independent	Dependent	Dependent
Tbet	Dependent	Dependent	Dependent	Dependent
Eomes	Dependent	Independent	Independent	Independent
Nfil3	Partially independent	Dependent	Independent	Dependent

Heterogeneity of ILC



Understanding ILC diversity
Unique and shared transcriptional
programs of mouse ILCs

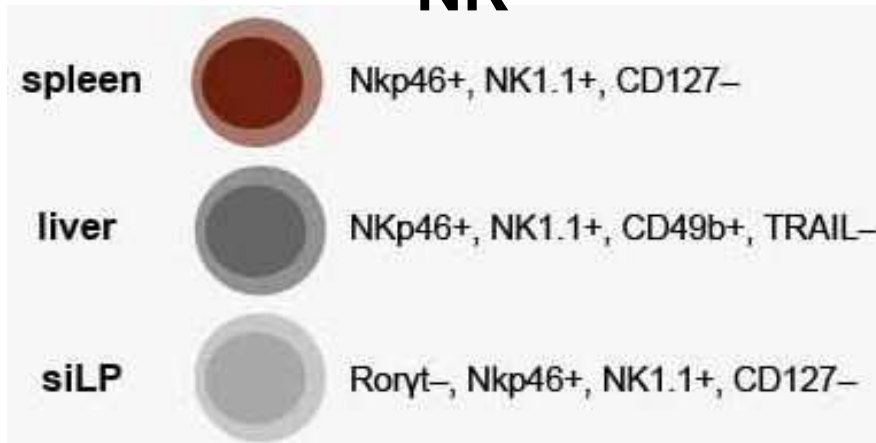
In collaboration with the Immgen
Consortium

<http://www.immgen.org/>

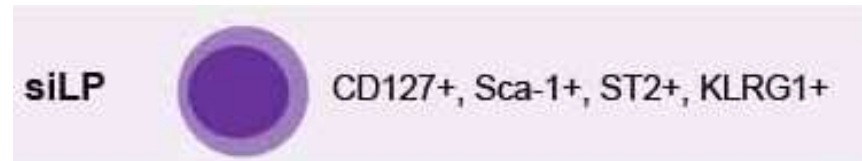
Michelle Robinette

ILC classes and subsets

NK



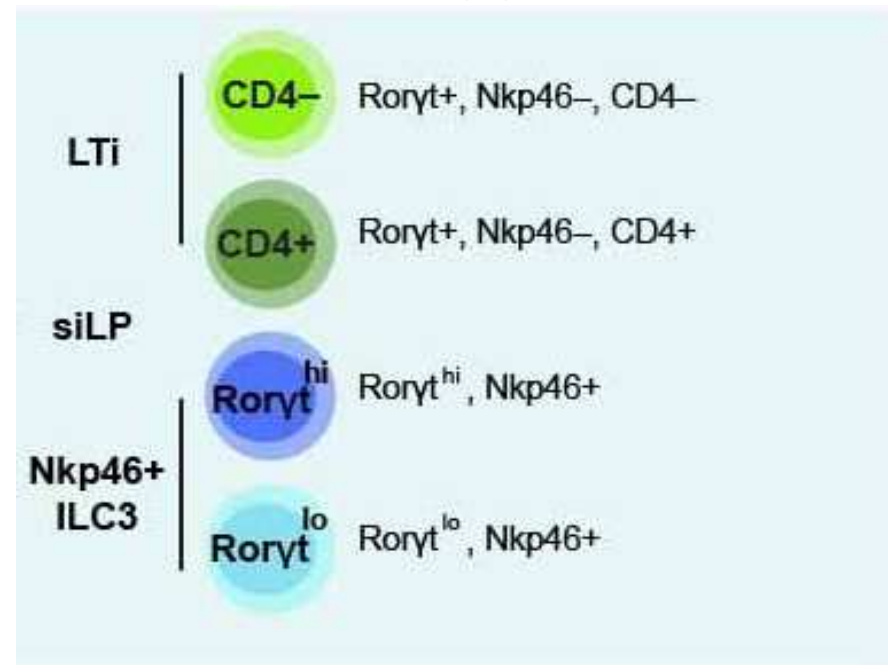
ILC2



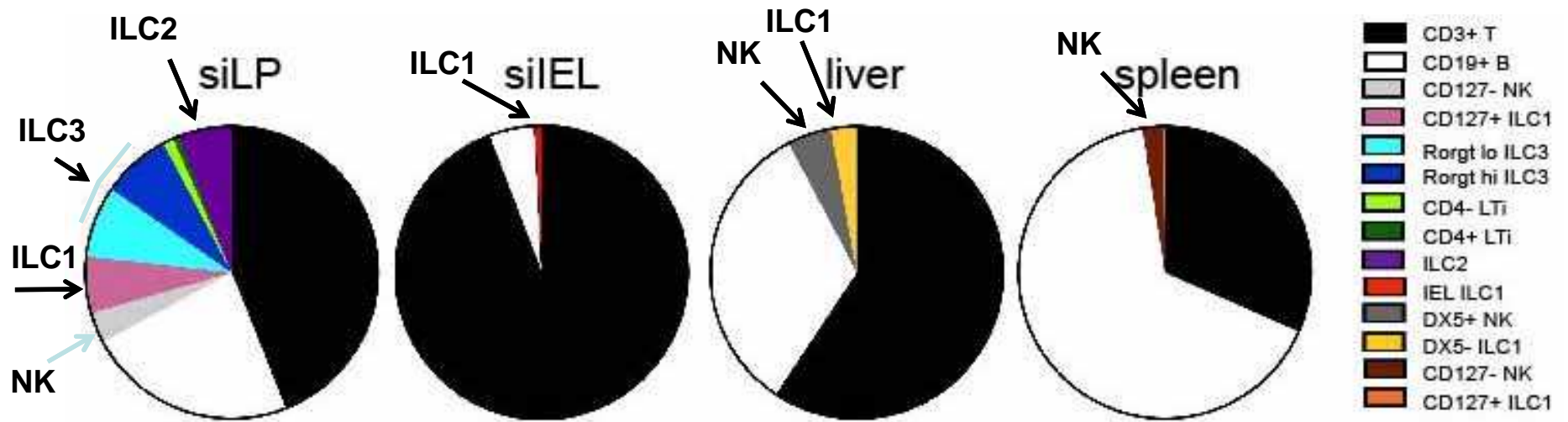
ILC1



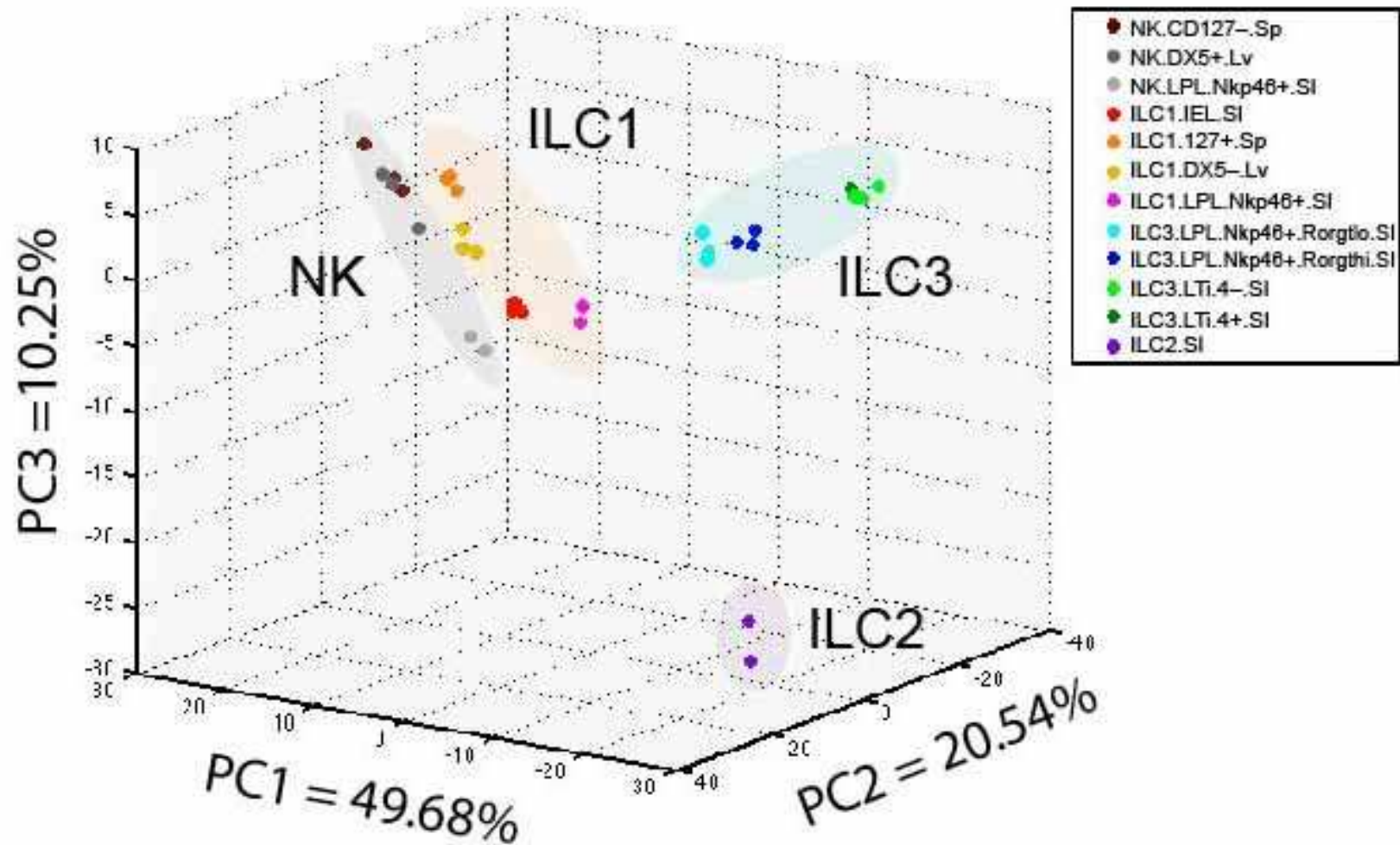
ILC3



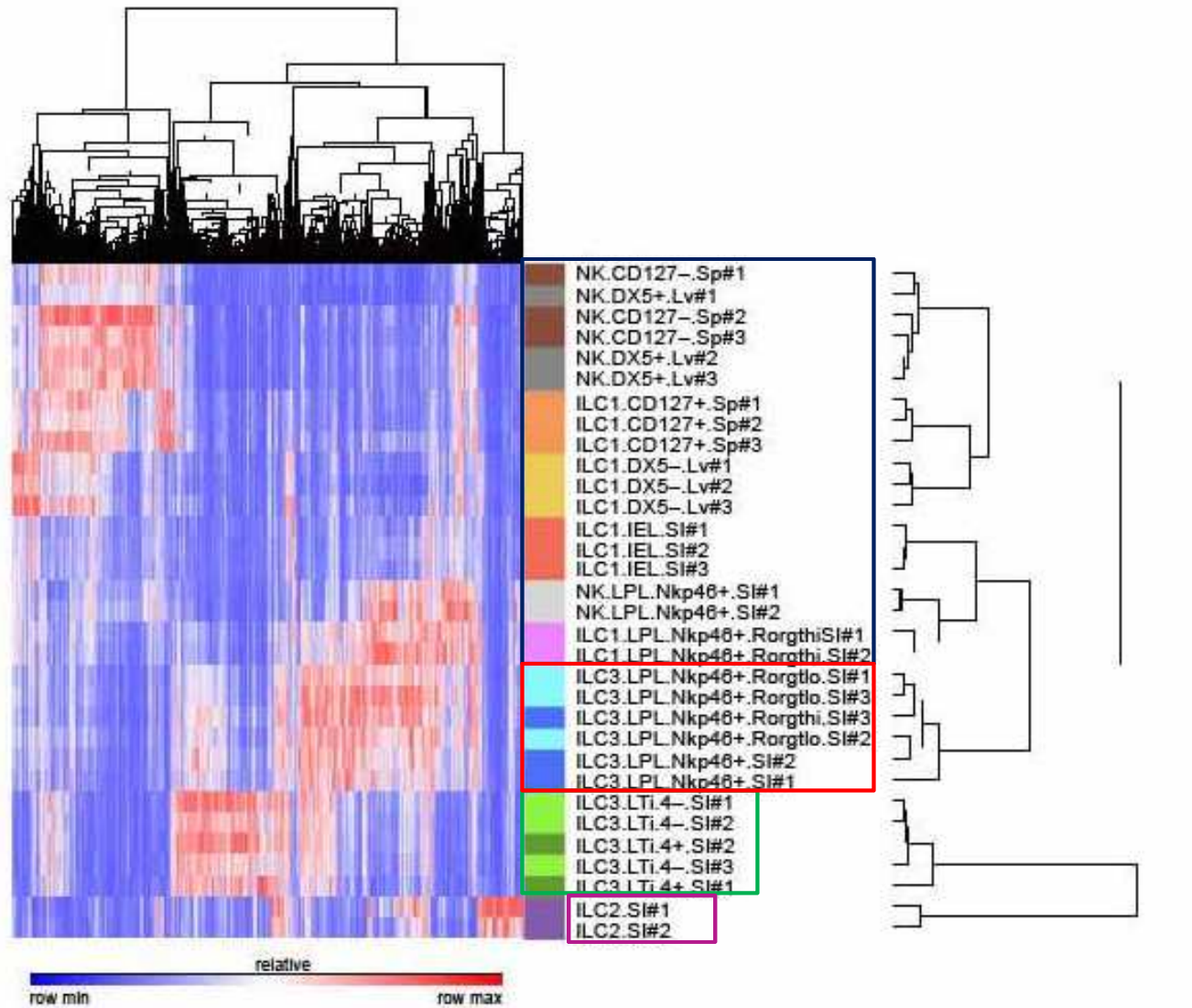
ILC subsets frequency



Principal component analysis of NK cell and ILC diversity



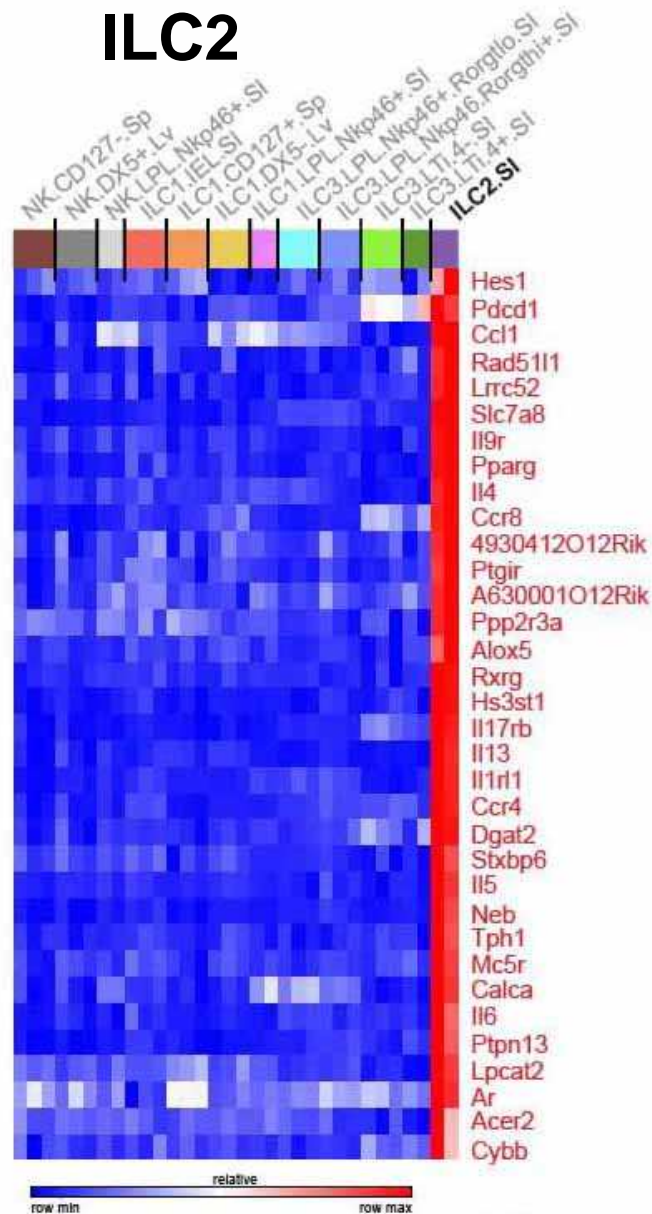
ILC1 and NK cells are intermixed



Hierarchical clustering of the top 10% differentially expressed genes

Heatmap of ILC2 unique transcripts

ILC2

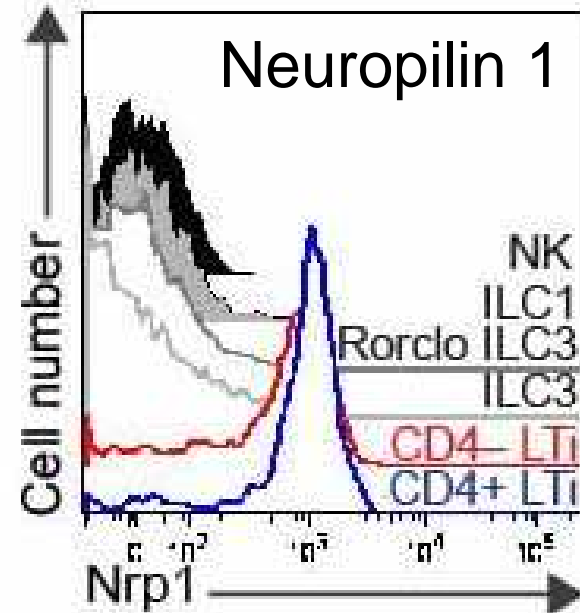
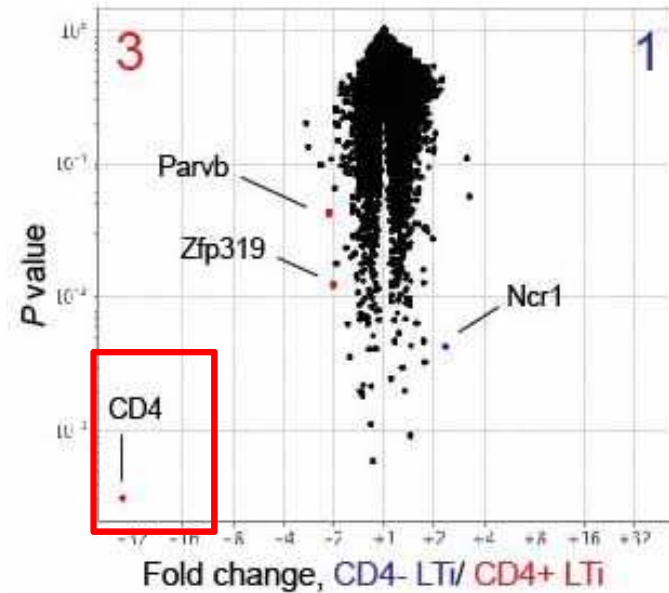
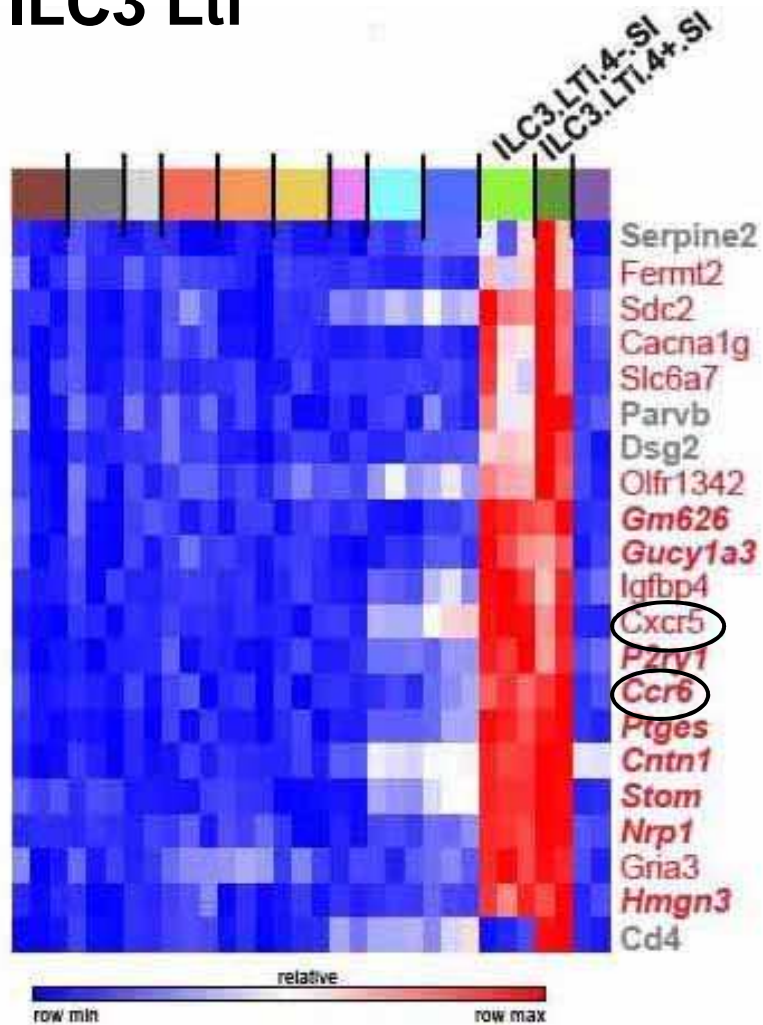


IL4
 IL5
 IL13
 IL9
 IL9R
 IL33R

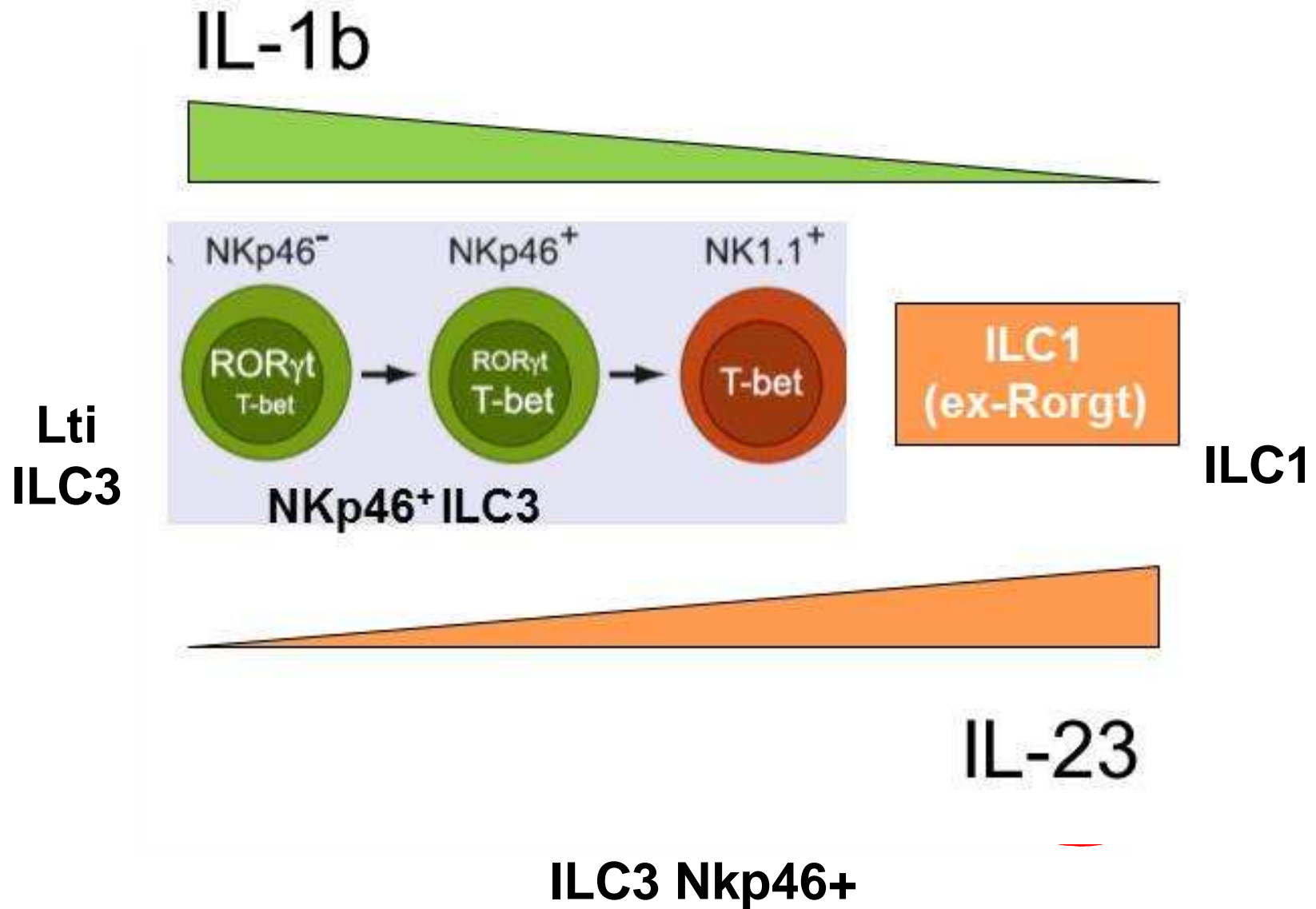
M2-like polarization
 Arginase 1
 PPAR γ
 Dgat2
 Lpcat2

Heatmap of ILC3 (Lti-like) unique transcripts

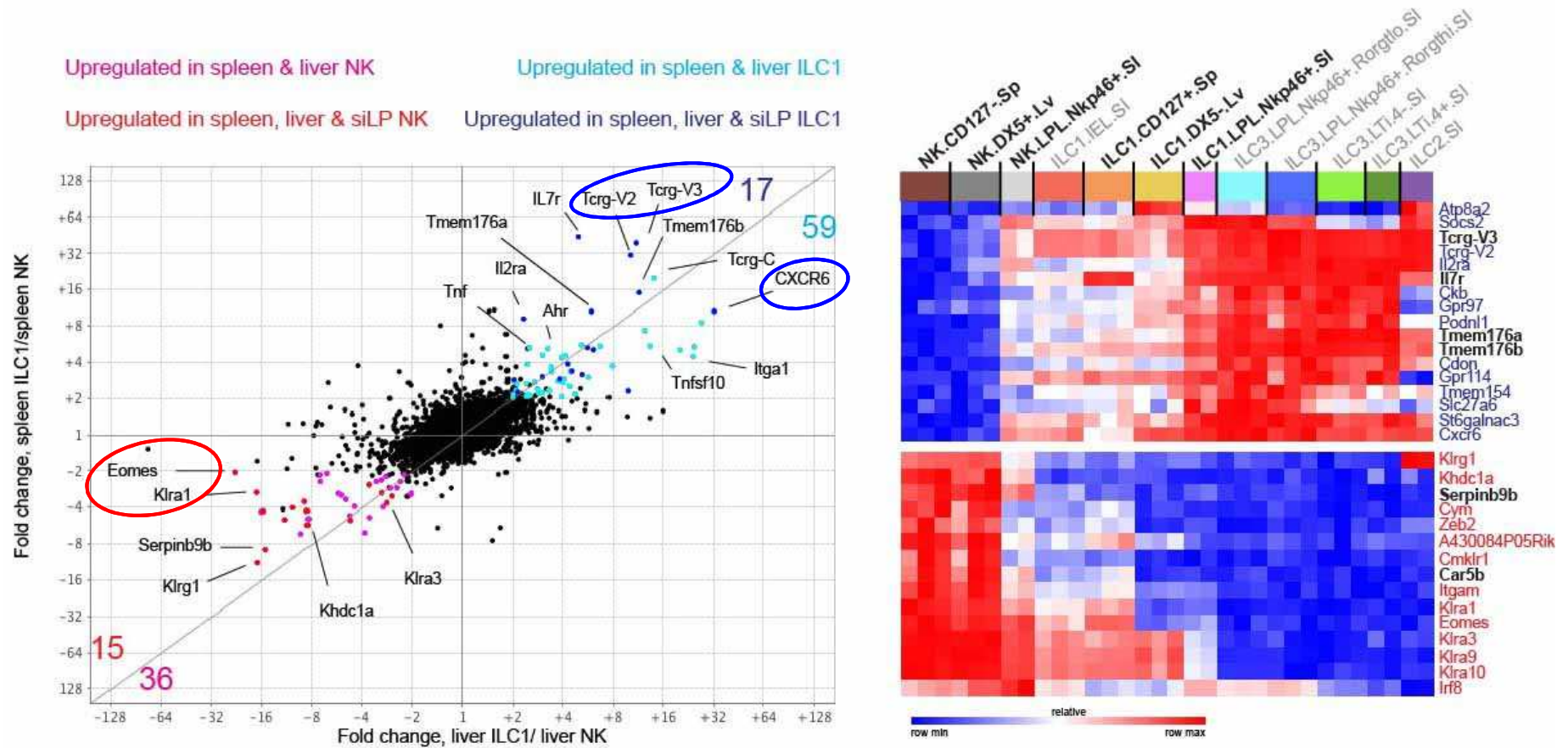
ILC3 Lti



NKp46+ILC3 share transcripts with Lti ILC3 but also ILC1

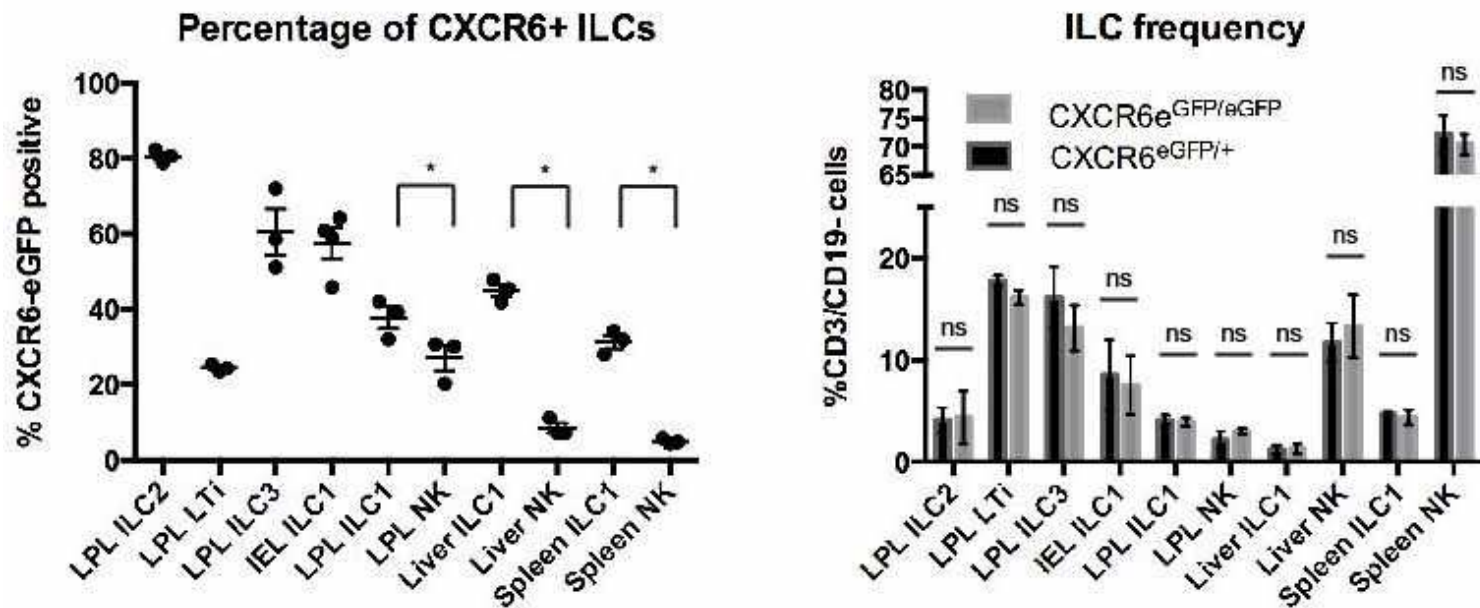


ILC1 transcriptional profile is intermediate between NK cells and ILCs



Core ILC signature

ILC core signature: CXCR6



CXCR6 deficiency has no impact on ILC frequencies but influences cytokine responses

Conclusions

- **ILC2** express transcripts indicative of M2 polarization
- Neuropilin1 is defining marker of Lti **ILC3**
- NKp46+ **ILC3** may have functional plasticity and convert into ILC1
- **Intestinal ILC and NK** share an “intestinal” gene signature probably reflecting environmental exposure
- There is a major functional overlap between **ILC1** and **NK** cells
- A core signature for ILC distinct from NK:
 - CXCR6, IL-7R, GLT of TCR γ , TMEM176a/b

Lab 2014



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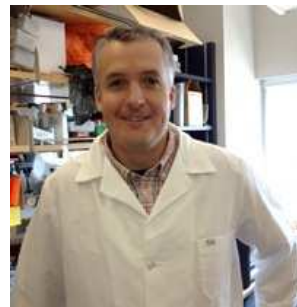
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