



IL-33 in asthma: From farming communities to animal models

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No Conflicts of Interest

Allergic asthma (Th2) is a chronic inflammatory disease of the airways

- Chronic inflammatory disease of the lung, affects over 300 million people ~7.5% of adult population (Braman Chest 2006, McCracken JAMA 2017)
- Symptoms: shortness of breath, chest tightness, and bouts of coughing or wheezing
- Pathology
 - Inflammation
 - Goblet cell mucus production
 - Smooth muscle thickening
 - Reversible airway constriction



Galli, et al. (2008) Nature and McCracken, et al. (2017) JAMA

Lung immunity to allergens



Genome Wide Association Studies (GWAS) to understand the genetic basis of asthma Low hanging fruit included IL33 and the IL33 receptor (IL1RL1/ST2)

 EVE Consortium: diverse ethnic backgrounds in US and Mexico, 2x10⁶ SNPs

• GABRIEL

Consortium:

European subjects, 582,892 SNPs, subjects from 23 studies





What is IL-33?



The NEW ENGLAND JOURNAL of MEDICINE

Efficacy and Safety of Itepekimab for Moderate-to-Severe Asthma



The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

Efficacy and Safety of Itepekimab in Patients with Moderate-to-Severe Asthma

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<u>Outline</u>

IL-33 is produced by dendritic cells and is involved in allergic sensitization in the lungs to multiple types of allergens

- IL-33 levels in Amish farm children are suppressed by their environment, but Hutterite children IL-33 levels are determined by their genetics
- IL-33 is produced by different lung cells in humans compared to mice.

Different stimuli promote development of various T helper responses



Signaling through FcγRIII diverts TLR-4 stimulated DCs toward DC^{Th2}



Bandukwala *et al,* (2007) JEM

Passive transfer of antigen specific IgG augments Th2 inflammation upon antigen challenge



Tjota, et al. JCI. 2013

What genes are regulating DC^{Th2} development?



Tjota, et, al. JCI. 2013 Williams, et al, Nature Comm 2013 Allergic responses can be triggered by a variety of structurally diverse allergens with varying biological functions

?

Immune

complexes

Type I (enzymatic):

- Dust Mite
- Fungal Spores
- Pollen

Type II (non-enzymatic):

- Animal Dander
- Latex
- Ovalbumin
- Wheat Flour



HDM extract targets multiple cellular pathways



Is HDM-mediated Th2 inflammation FcRy dependent?

HDM-mediated allergic airway inflammation is FcRy-dependent



Tjota et al, J. Allergy Clin Immun. 2014

HDM upregulates IL-33 in BMDCs in an FcRy-dependent manner



Tjota et al, J. Allergy Clin Immun. 2014

HDM-induced IL-33 upregulation in BMDCs is Dectin-2 dependent



Dectin 2^{-/-} bones provided by N. Barrett

Stimulation of FcRγ-containing receptors can lead to increased DC expression of IRF4 resulting in DC^{Th2} skewing



Bandukwala et al J. Exp Med. 2007 Tjota et al, J. Clin. Invest. 2013 Williams et al, Nature Comm 2013 Tjota et al, J. Allergy Clin Immun. 2014 Camacho, et al. JCI Insight, 2022

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The Farm Effect: Early childhood exposure protects against asthma and allergies



Erika von Mutius, JACI 2004

Amish farm children have a low asthma and atopy risk compared to Hutterites farm children

Previous Studies	Asthma	Allergic Sensitization (skin prick)
Amish children 6-12 (Holbreich et al. 2012)	5.2%	7.2%
Hutterite children 6-10 (Motika et al. 2011)	21.3%	33.3%

- Anabaptist founder populations from Europe
- Large families
- Long nursing time
- Germanic farming diet
- Consume raw milk
- Vaccinated
- Cats and dogs not allowed in home
- No television/radio exposure
- Children spend a lot of time outdoors
- No obesity in children
- *High level of genetic similarity between the two founder populations (by SNP analysis)*

Hutterite



Ree

Amish

mage credit: http://static.guim.co.uk/sysimages/Guardian/Pix/pictures/2012/8/1/134383410280 6/Amish-A-Secret-Life-008.jpg

Farm environment and exposures differ between Amish and Hutterite communities



Amish

- Family farms
- Traditional farming techniques
- Women and children help farm

Current Opinion in Immunology

Hutterites

- Communal farms
- Farming highly mechanized
- Women and children not involved in farming

Hypothesis

Amish children live in an environment that reduces risk of allergic sensitization through modulation of the immune response

Methods:

- Collect blood from 30 Amish and 30 Hutterite children
- Determine the specific changes in peripheral blood leukocyte (PBL) gene expression, cell phenotypes, and functions that could lead to reduced allergic sensitization.
- Tested the effect of house dust in a mouse model of asthma



Study Sample

Stein, Hrusch, Gozdz, et al. NEJM 2016

PBLs from Amish children produce less cytokine in response to LPS





IL-2, IL-5, IL-15, and IL-22 were trending down in the Amish



No difference observed in IL-10, IL-9, IL-13, IL-12p70, GM-CSF, MIP3A, IL-1β, IL-6, TNFα, IL-28A, IL-23, or IFNγ

Amish PBL have a globally dampened transcriptional response to LPS:



Genes significantly different in

Conclusions:

Amish kid's blood cells have a higher basal level of neutrophils and gene transcription of innate genes in neutrophils, but are hypo-responsive to innate stimulation.

Evidence of hyper-reactivity in a "bored" immune response in Hutterite children?

Tregs may be directly involved in the suppression of the innate immune cells in the Amish children.

Conclusions: Effects of Amish environment on circulating immune cells: Potential mechanisms for reduced asthma risk



Hrusch, Stein, et al. JACI 2019

PBLs from Amish children produce less IL-33 cytokine in response to LPS and at baseline



At baseline, IL-33 was the only cytokine that was different between the Amish and Hutterite children

Lung immunity to allergens



Genome Wide Association Studies (GWAS) have been performed to understand the genetic basis of asthma

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diverse ethnic backgrounds in US and Mexico, 2x10^6 SNPs

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European subjects, 582,892 SNPs, subjects from 23 studies





SNPs associated to increased asthma risk are located upstream of the IL33 gene



Bayesian fine-mapping quantifies uncertainty in variable selection



Risk allele of GWAS SNP rs1888909 is associated with IL33 mRNA and IL-33 protein levels



Protein levels (plasma) From Hutterite Children



No association was found in Amish children's plasma suggesting that the farming environment trumped their genetics



Aneas, I. Decker, D.Sperling, AI, Nobrega, M. Nature Comm 2021

The 5 kb asthma-associated region contain important regulatory activity



Aneas, Decker,.....Sperling, Nobrega. Nat Comm 2021

IL-33 is regulated by a causal SNP



Aneas, Decker,.....Sperling, Nobrega. Nat Comm 2021

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In the <u>lungs</u>, human *IL33*-reporter is expressed primarily in endothelial cells

BAC (+) Lung



Deletion of the 5kb insulator region eliminates expression in CD31+ MHC II+ microvascular endothelial cells



Deletion of the 5kb insulator region eliminates hIL-33reporter expression in tracheal basal epithelium and SMG





SNPs associated to increased asthma risk are located upstream of the IL33 gene



GFP (murine *II33* reporter) and Crimson (human *IL33* reporter) are expressed by distinct populations in the lungs



Crimso GFF Composite αCrimson α GFP

Lungs

Allergen sensitization and challenge *decreases* human-IL-33 reporter expression, but not murine IL-33 reporter expression



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IL-33 *decreases* human IL-33-reporter expression in endothelial cells suggesting a negative feedback loop



IL-33 *decreases IL33* mRNA expression in endothelial cells suggesting a negative feedback loop



Conclusions

- We have identified a 5kb region that confers tissue-specific expression of IL33
- The 5kb region has insulator activity and contains GWAS SNPs that affect its function
- Human IL-33 is primarily expressed in *endothelial* cells, not epithelial cells as in mice.
- IL-33 negatively regulates its own expression in endothelial cells, and this downregulation is dependent on the 5kb region

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Funded by NHLBI R01, NIAID U19 - and NIAID T32, F31, AHA postdoc Does rIL-33 downregulate *IL33* transcription by reducing chromatin accessibility at the *IL33* locus?



Domenick Kennedy

Extracellular IL-33 does not alter accessibility at the *IL33* locus



Does extracellular IL-33 increases *IL33* mRNA turnover?



Exogenous IL-33 treatment decreases nuclear *IL33* mRNA stability in human endothelial cells



Human IL33 mRNA in HUVECs

IL-33 *decrease* of human IL-33-reporter expression is dependent on the 5kb insulator region



DEL-BAC (+)

Acknowledgements

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Bayesian fine-mapping quantifies uncertainty in variable selection



- Heuristic LD approach deterministically chooses SNPs above certain LD threshold with the lead SNP
- Limitations: arbitrary threshold, doesn't quantify uncertainty



Credible set based on SNP PIPs

- Bayesian fine-mapping probabilistically evaluates each SNP's causal evidence
- **Posterior inclusion probability (PIP)**: the probability of a SNP being causal
- **95% credible set**: a set of SNPs that has 95% or greater probability of containing at least one causal SNP

Bayesian fine-mapping quantifies uncertainty in variable selection



SC-RNAseq demonstrate *IL33* is abundantly expressed in human lungs



Reyfman et al., 2017

Basal lung cells downregulate *IL33* mRNA as they differentiate in ALI epithelial cultures *in vitro*



Crimson (human *IL33* reporter) is expressed largely by non-hematopoietic cells in BAC (+) lungs



New BAC Tg mice that make hIL-33



Tissue	Hu CD4 (HuIL-33 reporter)	GFP (Mouse-IL33 reporter)	Double positive?
Epithelial cells	Neg	++++	None
Lym Endo	++++	+	+ (all GFP are also huCD4)
Vasc Endo	+	Neg	None
Fibroblast	Neg	++	None
Neutrophils	++++	Neg	None
Eosinophils	+/-	Neg	Noe
AM	+	Neg	None
Intersitial Macs	+	Neg	None
DC1	++++	Neg	None
DC2	++++	Neg	None
Ly6C+ Mono	+++	Neg	None

IL33-GFP FMO Neutrophils 10' 10⁶ 10⁵ mIL33+ 0.27 hCD4+ BAC(-) BAC(+) 0.76 104 107 107 mIL33+ mIL33+ 0.020 0.26 10⁶ 10⁶ -10 4 10⁵ 10⁵ 105 10⁶ - 10 hCD4+ hCD4+ hCD4-PE FMO 0.48 40.4 10⁴ 104 10 10⁶ mIL33-GFP 0 0 10⁵ mIL33+ 0.39 hCD4+ -10 ⁴ = -10⁴ -0.044 10⁴ 10⁵ 10⁵ 10⁶ 10⁶ -104 -104 104 104 0 0 hCD4-PE -10 4

Gated on live, CD45⁺, Lin (CD3/CD19/NK1.1)⁻ cells

0

-104

10⁵

104

106



Figure 4.1. *IL33* is expressed in the hematopoietic cells of human pulmonary tissue and varies widely between individuals. qPCR for *IL33* levels in the d0 RNA isolates from the human pulmonary hematopoietic cells of 50 lung donors.Values are



Figure 4.10. *IL33* is produced by a CD66b+ granulocytic population of lung cells. Distinct cell populations from human lungs were isolated by Flow Cytometric Sorting (FACS), from which cDNA was prepared (Continued on page 96).

Figure 4.10, continued. (A) R305 was sorted into four broad populations based on the following phenotypes: Red: CD45-, White: Granulocytes (CD45+CD66b+), Green: Dendritic Cells (CD45+CD11c+), and Blue: Other Cells (CD45+CD11c-CD66b-). *IL33* levels are shown relative to *ACTB*. (B) R317, R358, and R343 were sorted base on the following phenotypes: Red: CD45-, Grey: Monocytes and macrophages (CD45+CD14+), Orange: Lymphocytes (CD45+CD14-CD3/19+CD66b-), White: Granulocytes (CD45+CD14-CD3/19+CD66b-), White: Granulocytes (CD45+CD14-CD3/19-CD66b+), and Blue: Other Cells (CD45+CD14-CD3/19-CD66b+), *IL33* levels are relative to *GAPDH*. (C) R371 and R358 were sorted based on the following phenotypes: White: All granulocytes (CD45+CD14-CD3-CD19-CD66b+), Yellow: CD16hi granulocytes (CD45+CD14-CD3-CD19-CD66b+CD16hi), Cyan: CD16lo granulocytes (CD45+CD14-CD3-CD19-CD66b+CD16lo).