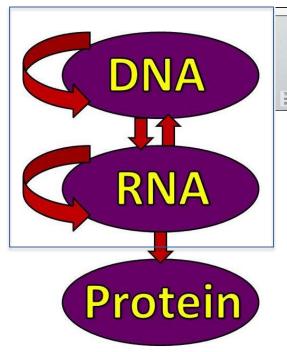
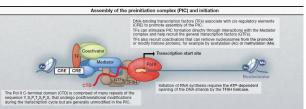
Chromosome Dynamics in T cells

Lark Kyun Kim, Ph.D.

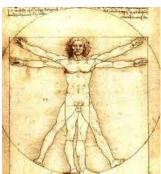
Assistant Professor
Severance Biomedical Science Institute
Yonsei University College of Medicine

Our Ultimate Goal









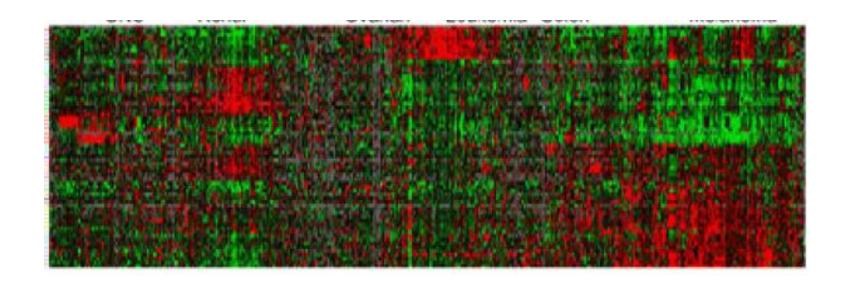
Mechanism



Biological phenomenon (in vivo and physiologically)

target genes biomarker

Genome-wide studies



Genome-wide sequencing

- RNA seq
- Exome seq
- ChIP seq
- Gro-seq
- Mi-seq
- LincRNA-seq

Bioinformatics

Top 20 (or whatever number) Up- or down- regulated gens

Question (I)

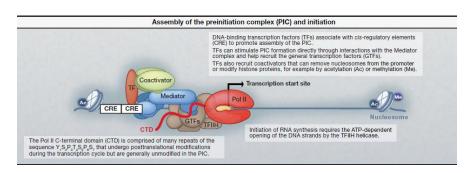
How many genes are being transcribed simultaneously in a given time point?

Things to think about

- 1. The number of transcription machinery such as RNA polymerase II and TBP is unlimited?
- 2. Transcription can undergo anywhere in the nucleus?

Question (II)

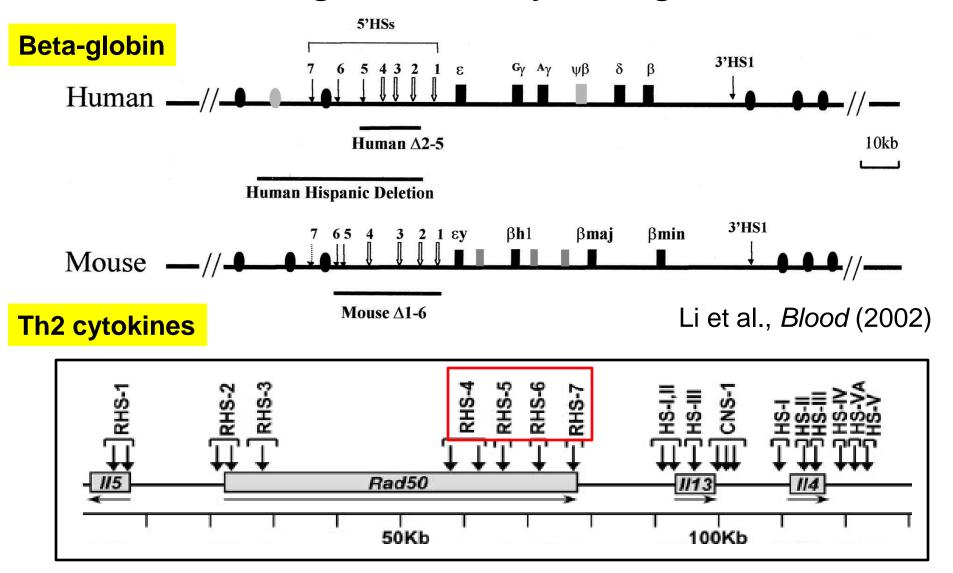
Each gene is regulated individually only via its own promoter and/or enhancer or a kind of control tower oversees the transcription globally?



Fromm et al., Cell (2013)

Locus control region (LCR)

- Beta-globin & Th2 cytokine genes-



Spilianakis et al., Nat Immunol (2004)

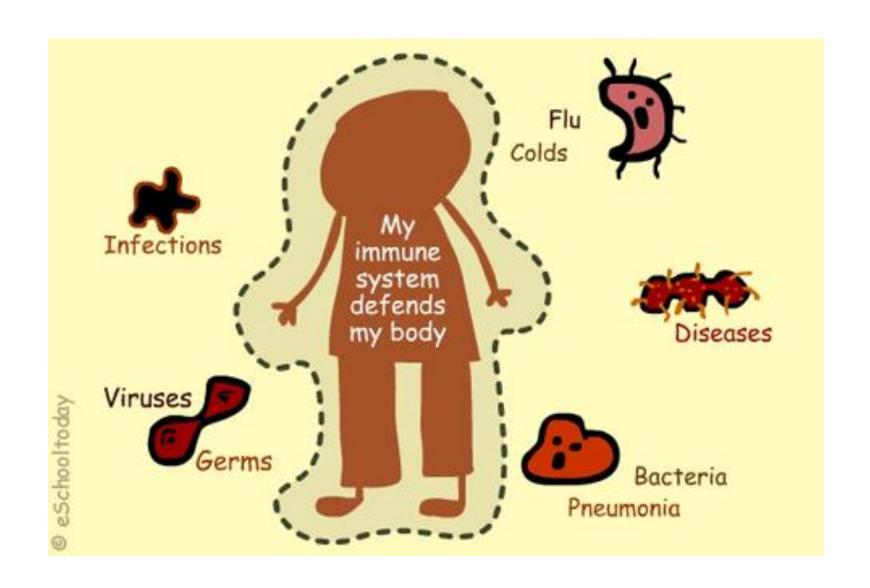
Question (III)

If there is a control tower which regulates a series of genes located closely at the same time,

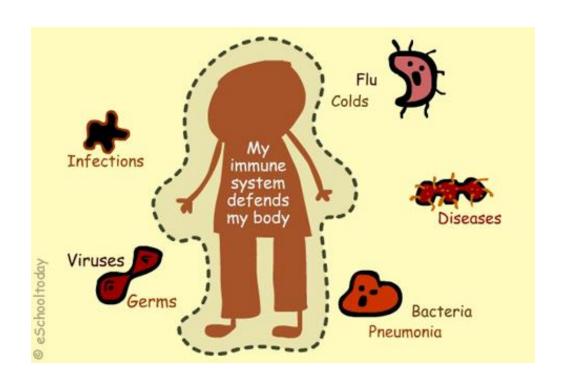
what happens?
in case of the relevant genes are located on
the different chromosomes

and moreover, one gene should be selected and the other should not

Immune System



Pathogens can be classified into 3 groups



Intracellular pathogens

Virus
Listeria
Salmonella
Legionella
Mycobacterium

Allergy

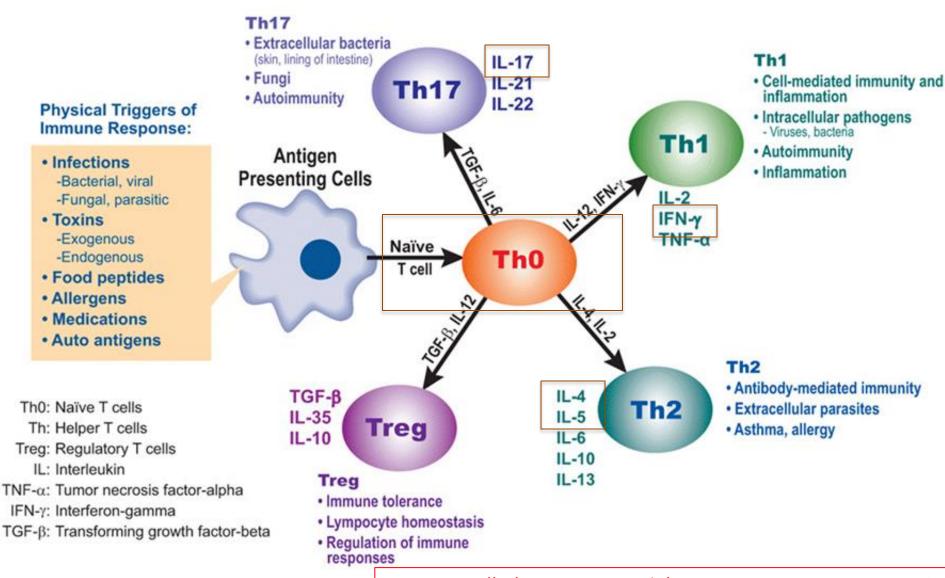
Food allergy
Atopic dermatitis
Asthma

Helminth

Extracellular pathogens

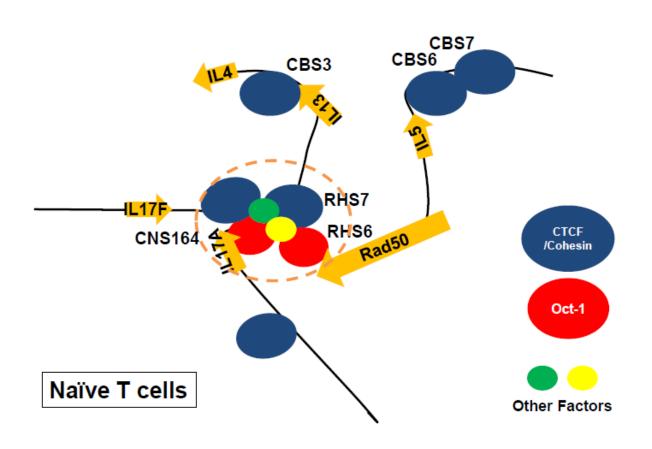
Bacillus
Mycoplasma
Pseudomonas
Vibrio
Streptococcus

T cells are key regulators of immune systems

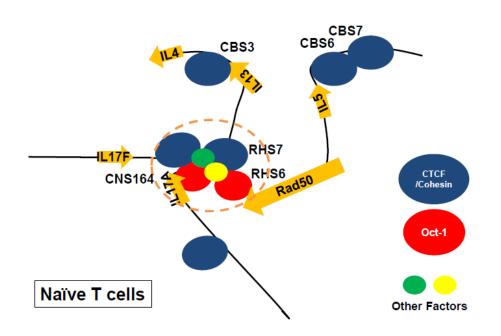


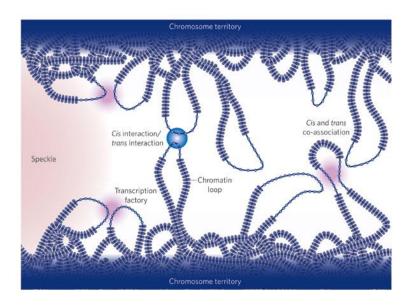
Naïve T cells have a potential to express IFNg, IL4 or IL17 These genes are located on different chromosomes

Interchromosomal association of each cytokine loci in naïve T cells



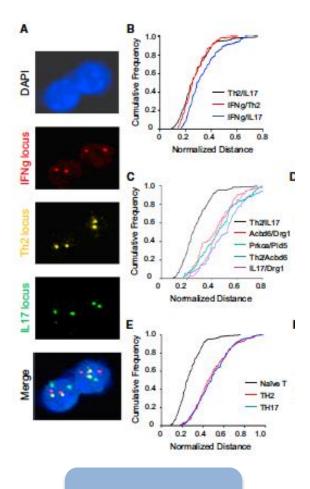
Interchromosomal association of each cytokine loci in naïve T cells

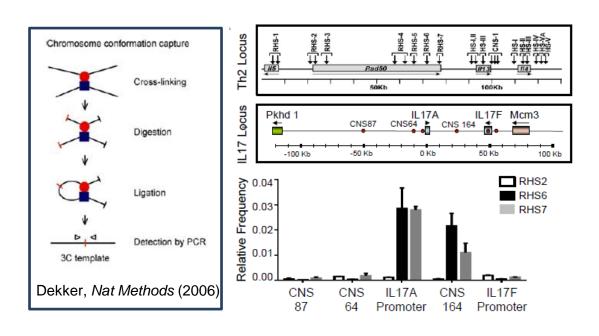




Fraser & Bickmore, *Nature* (2007)

Interchromosomal association of each cytokine loci in naïve T cells



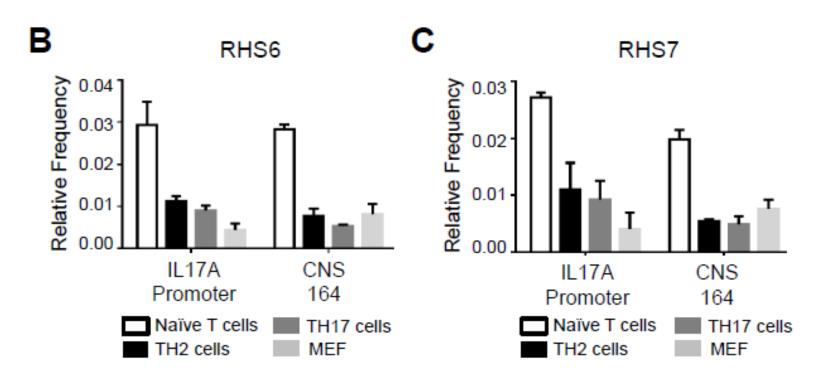


3C /4C Hi-C

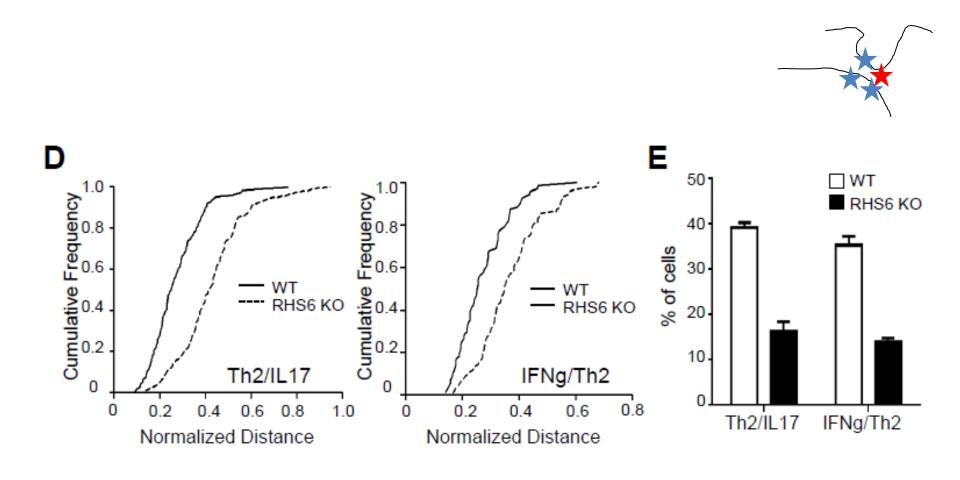
DNA FISH

3C qPCR b/w Th2 locus and IL17 locus





Interchromosomal association of cytokine loci in the absence of RHS6 in naïve T cells

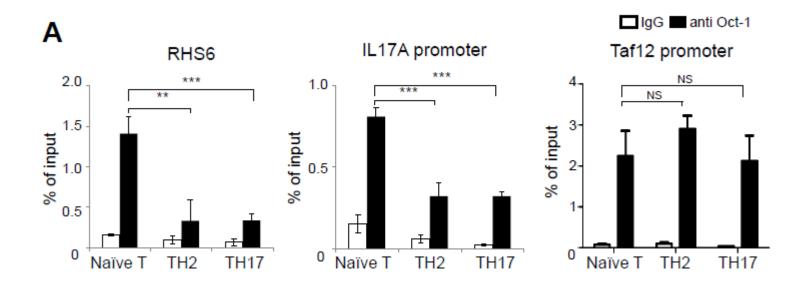


Binding proteins on interest regions

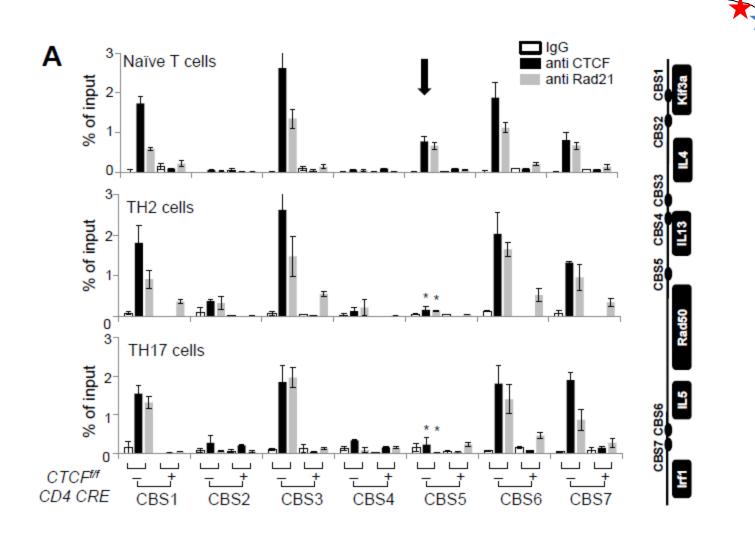
RHS6 of Th2 locus and promoter of IL-17 gene: Oct-1

RHS7 of Th2 locus and CNS164 of IL-17 locus: CTCF

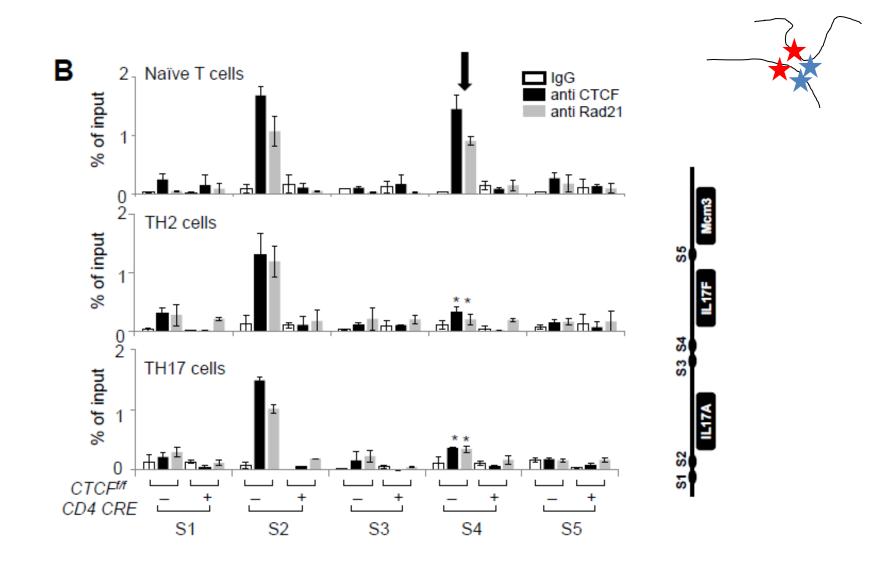
The binding of Oct-1 at RHS6 and IL-17 promoter in naïve T cell is reduced after differentiation



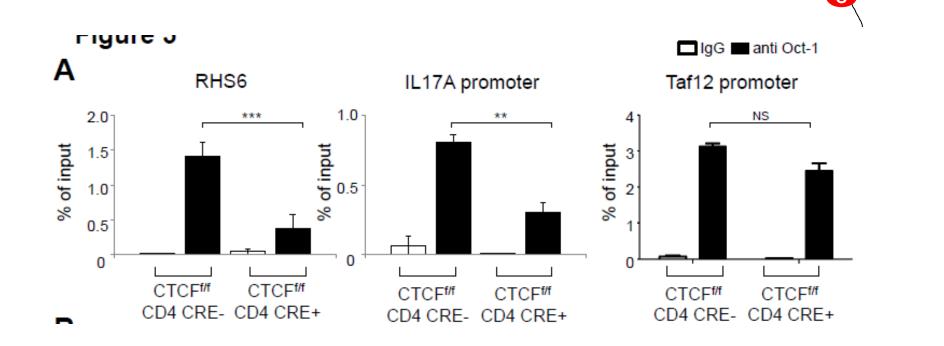
The binding of CTCF in the Th2 locus during T cell differentiation



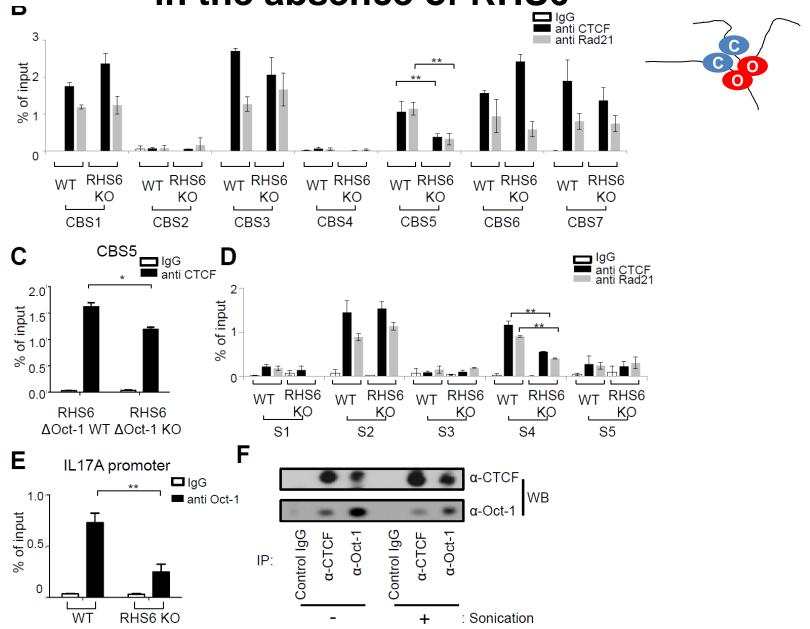
The binding of CTCF in the IL-17 locus during T cell differentiation



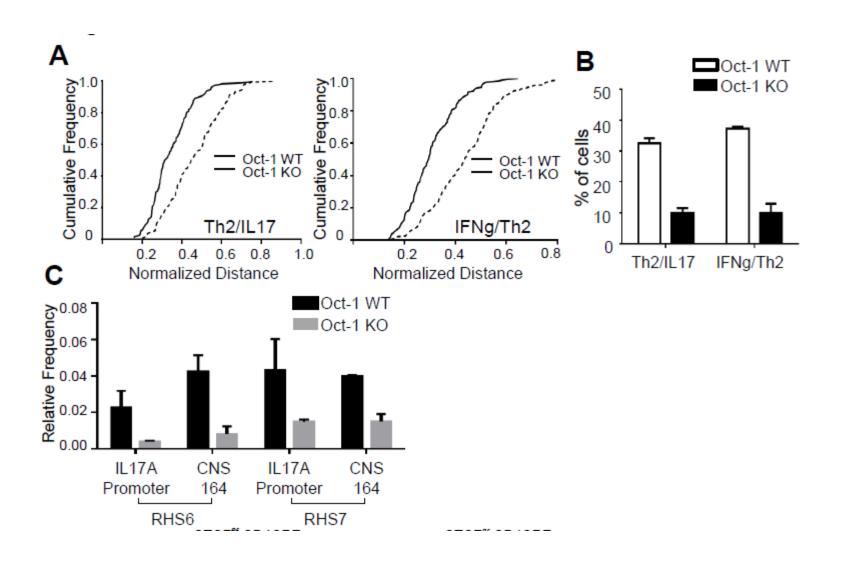
The binding of Oct-1 at RHS6 and IL-17 promoter in the absence of CTCF



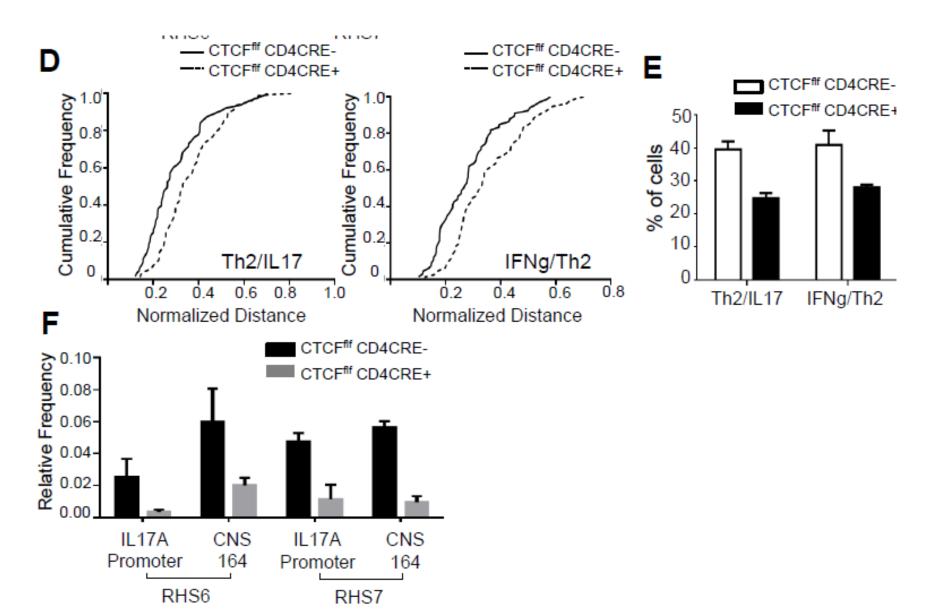
The binding of CTCF at RHS7 and CNS164 in the absence of RHS6



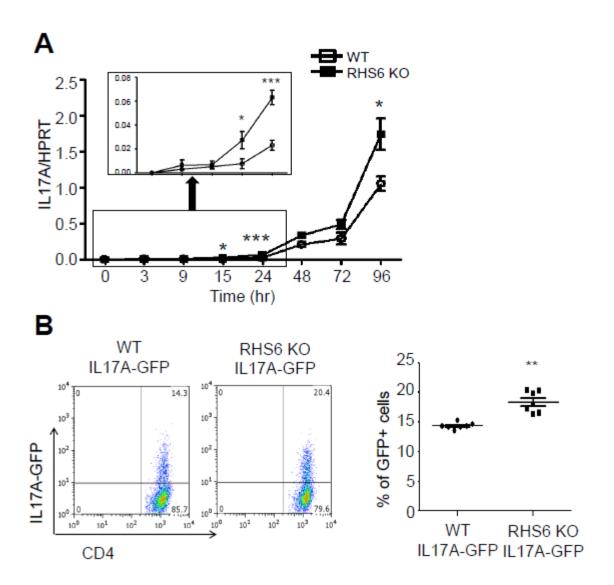
Interchromosomal association of cytokine loci in the absence of Oct-1



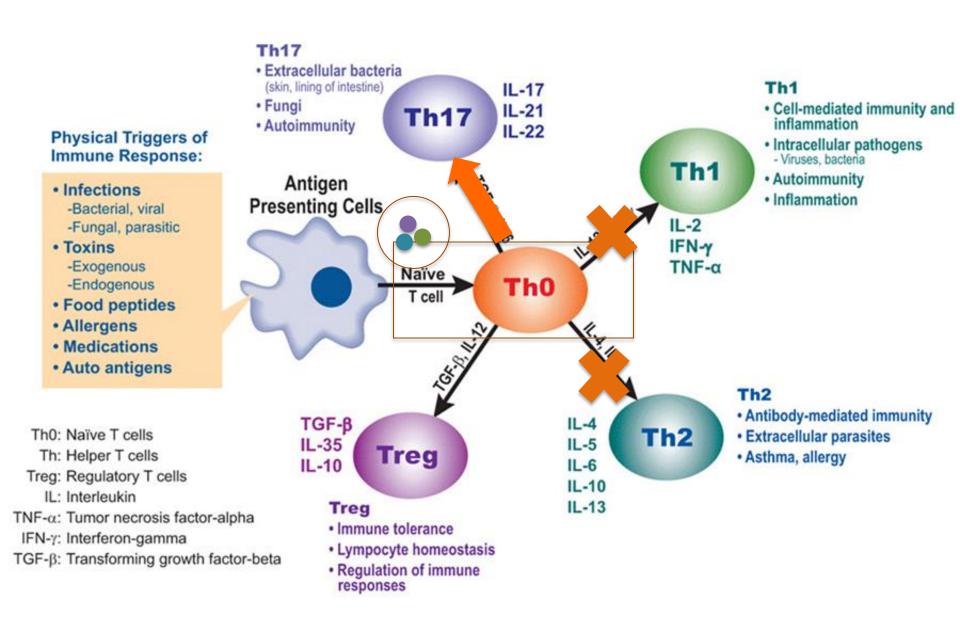
Interchromosomal association of cytokine loci in the absence of CTCF



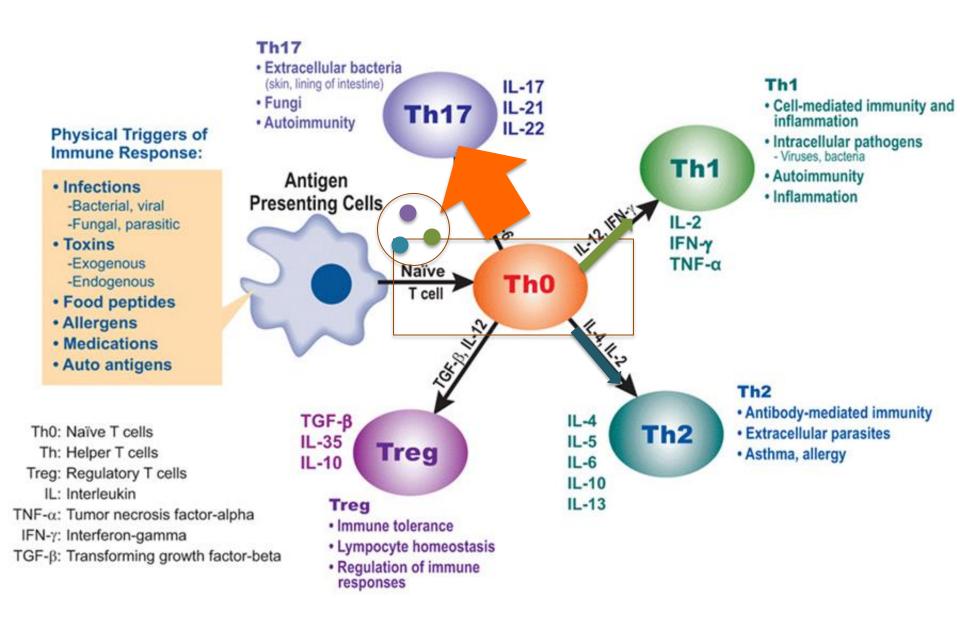
More Th17 cells generation in the absence of interchromosomal association



Aberrant production of effector T cells in the absence of interchromosomal association

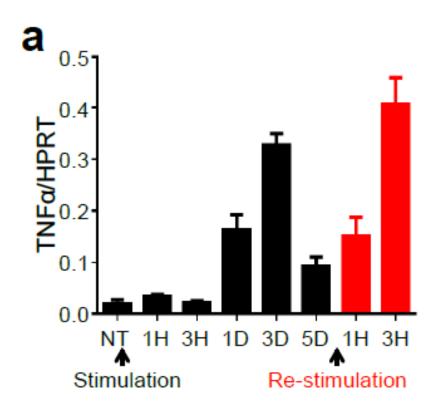


Aberrant production of effector T cells in the absence of interchromosomal association



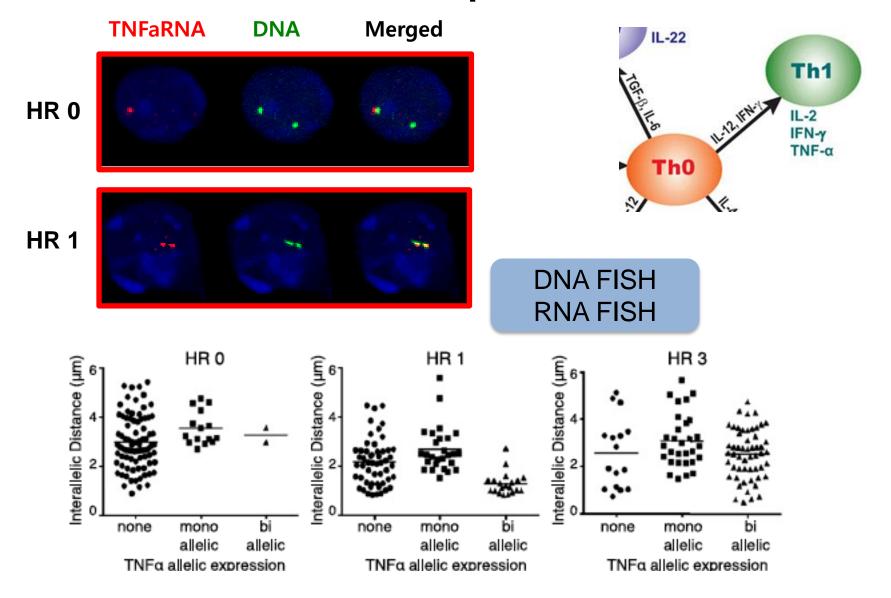
Another study on spatiotemporal regulation of transcription

Immunological memory is a key feature of immunity but molecular basis of this remains unclear

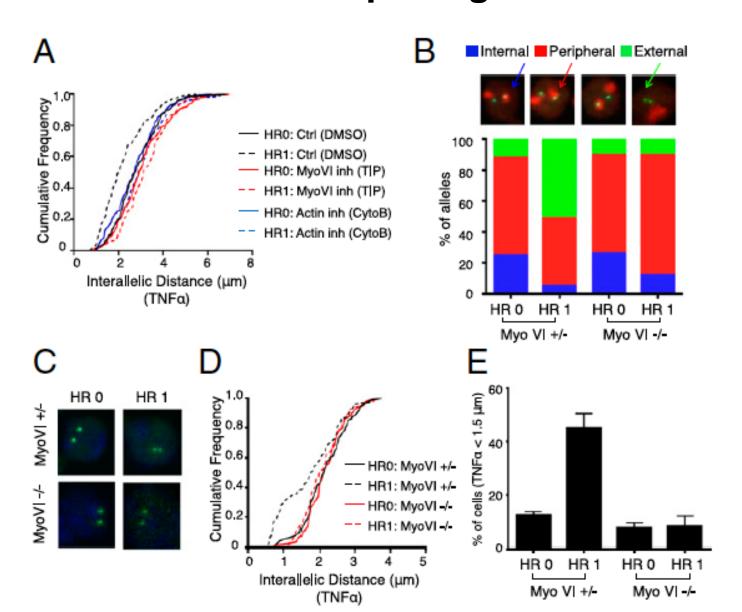


The resting Th1 cells can express TNF alpha rapidly and more efficiently upon previously encountered challenge

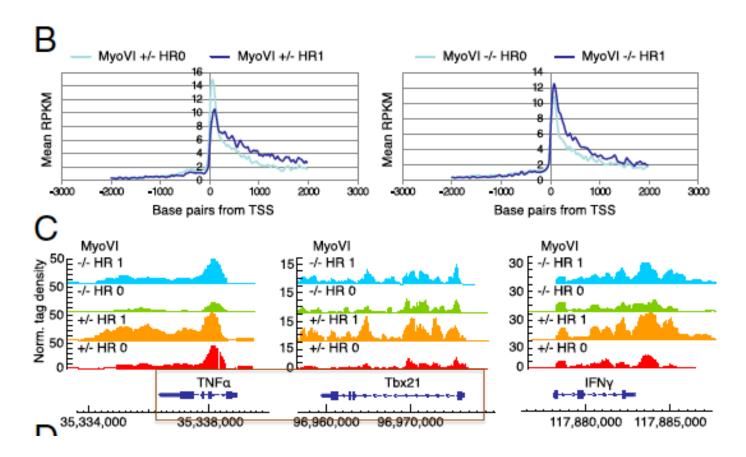
Allelic pairing of TNF loci is required for biallelic expression



Myosin VI is a motor protein that drives allelic pairing

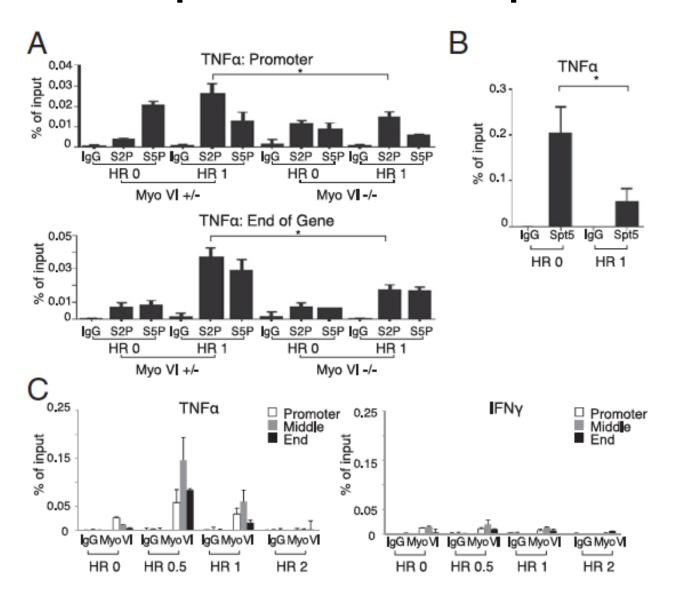


Pol II is paused at the promoter and waits for re-challenge

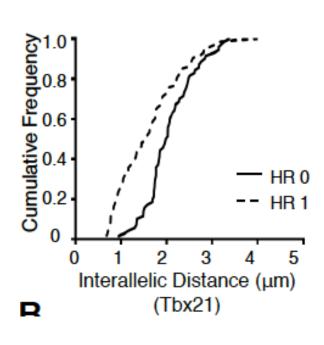


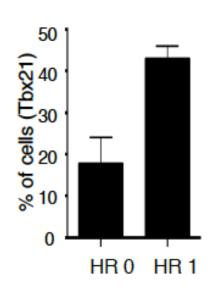
Gro-seq (Global run-on sequencing)

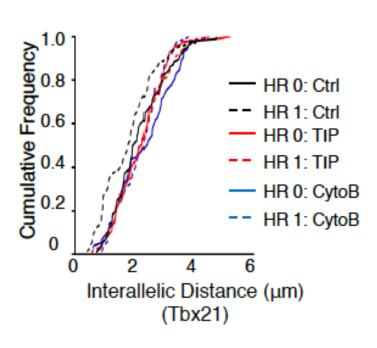
Initiation Pol II is already bound on the promoter of TNF alpha



T-bet also shows a homologous pairing in Th1 cells after re-stimulation







Current knowledge in the field

Nucleus is a well-organized structure.

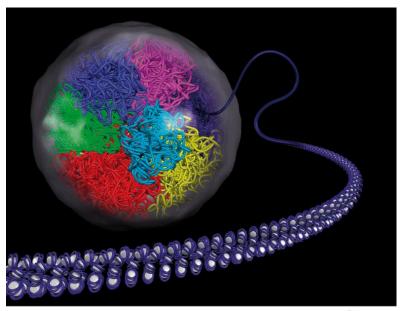
Chromosome territories

There has been a big advance of spatiotemporal regulation of transcription recently including us but most studies are done using cell lines and artificial system unlike ours.

PRINT ISSN: 0305-1048 ONLINE ISSN: 1362-4962

Nucleic Acids Research

VOLUME 37 ISSUE 19 2009 www.nar.oxfordjournals.org



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Fraser & Bickmore, Nature (2007)

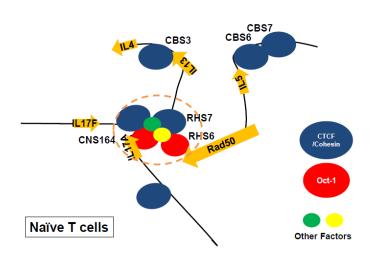
First demonstration of the relevance between interchromosomal associations and gene expression in primary cells

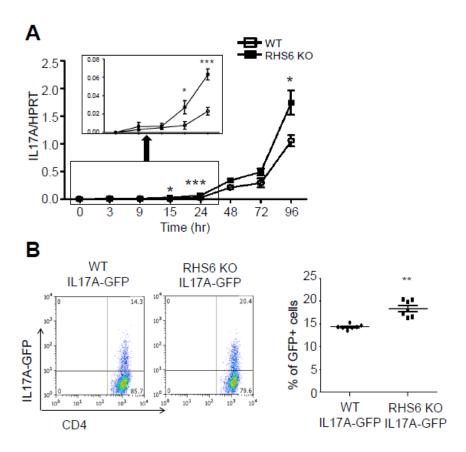


Molecular Cell
Article

Oct-1 Regulates IL-17 Expression by Directing Interchromosomal Associations in Conjunction with CTCF in T Cells

Lark Kyun Kim, 1,2 Enric Esplugues, 1,6 Cornelia E. Zorca, 1,3 Fabio Parisi, 4 Yuval Kluger, 4 Tae Hoon Kim, 3 Niels J. Galjart, 5 and Richard A. Flavell 1,2,4





Weakness → significant but not dramatic changes!!

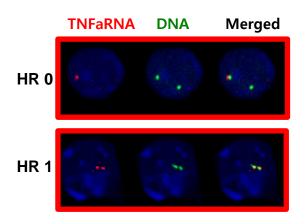
Current knowledge in the field

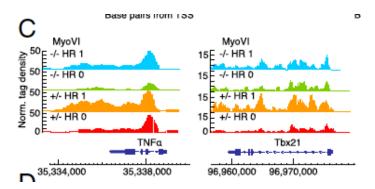
Homolog pairing by Myosin VI and biallelic expression : novel and hot

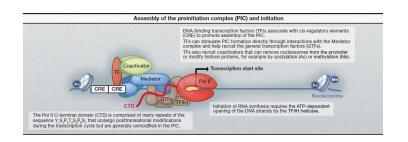
TNF: first coming cytokines

Tbet: master regulator of Th1 cells

Pol II pausing (rapid induced and developmentally Significant genes)







Current challenges

- 1. Direct evidence
- 2. More mechanism
- 3. Another examples

Acknowledgements

Yale University

Richard Flavell

David Schatz Tian Chi Tae Hoon Kim



Cornelia Zorca YA

All other lab members

Brown University

Min Jong Kang Chun Geun Lee Jack Elias

FUNDING



HHMI

Korea Research Foundation

