

# **Microbiology 204**

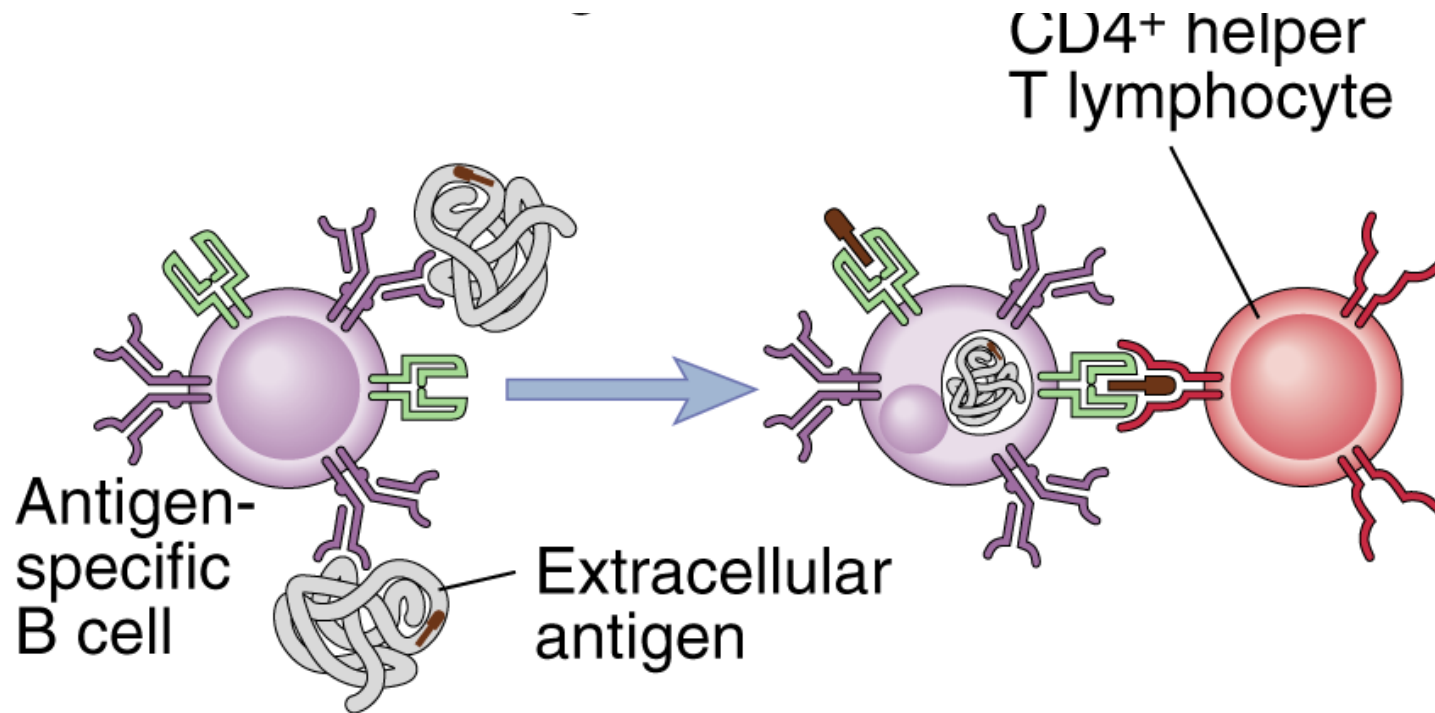
## **TCR Structure**

**Art Weiss**

**October 10, 2018**

**What do T cells see with their antigen receptors?**

# T cells and B cells use distinct antigen receptors to recognize fundamentally different forms of antigen

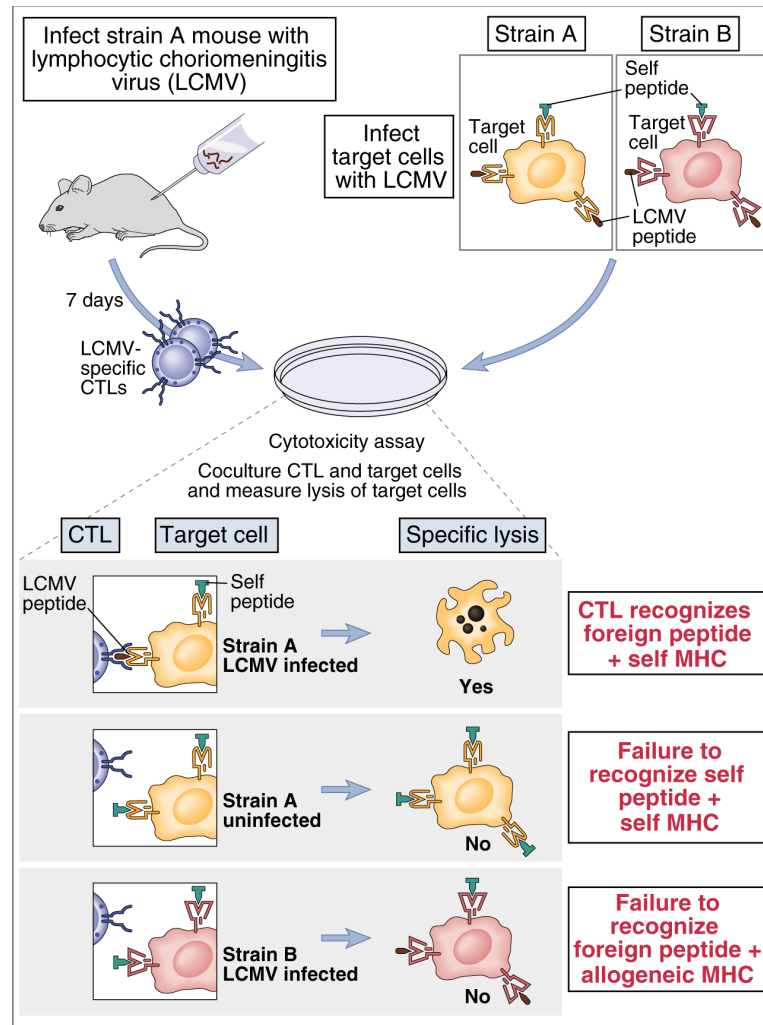


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B cells can recognize either **linear or conformational epitopes** of proteins, of carbohydrates or of lipids. The B cell antigen receptor is a form of membrane Ig.

T cells generally recognize “only” **linear peptide fragments** that are bound to MHC class I or class II molecules.

# MHC Restricted Recognition of Antigen



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Zinkernagel and Dougherty

Bevan

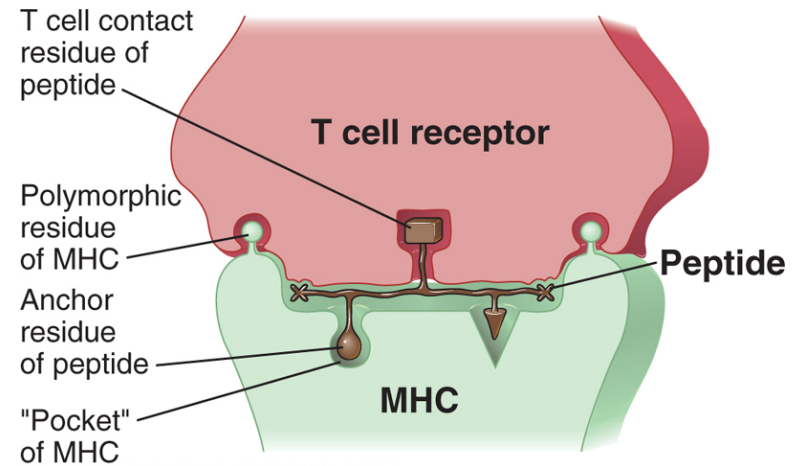
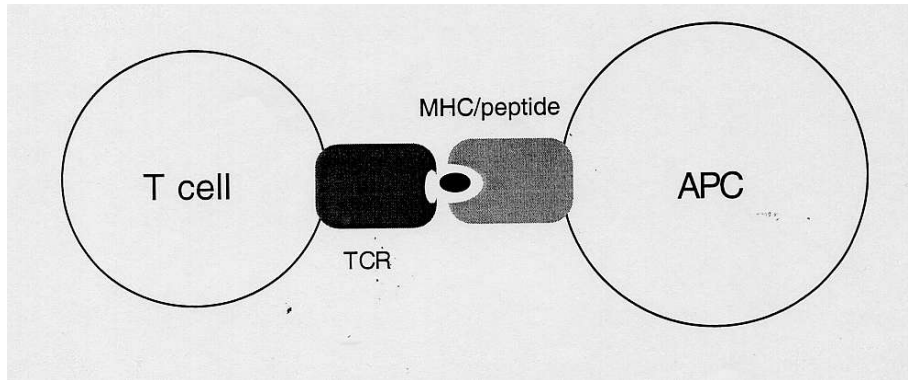
Mid -1970' s

***T cells only recognize specific peptide antigen in the context of self: MHC restriction. Specificity for self recognition is encoded in the MHC (Major Histocompatibility Complex).***



# MHC Restriction: How does the TCR simultaneously recognize MHC specificity and antigen specificity?

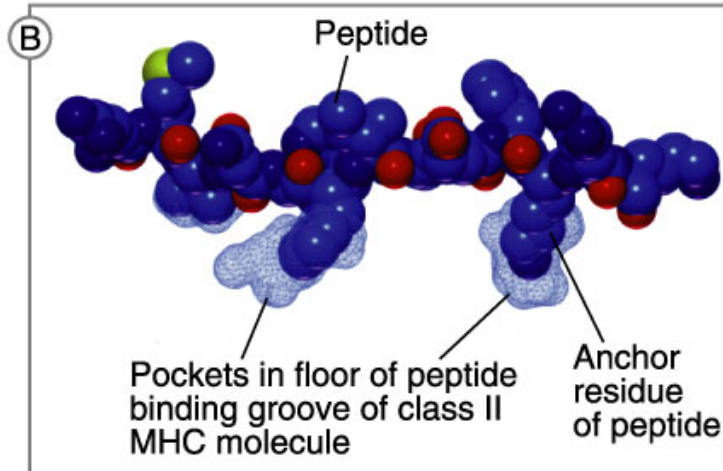
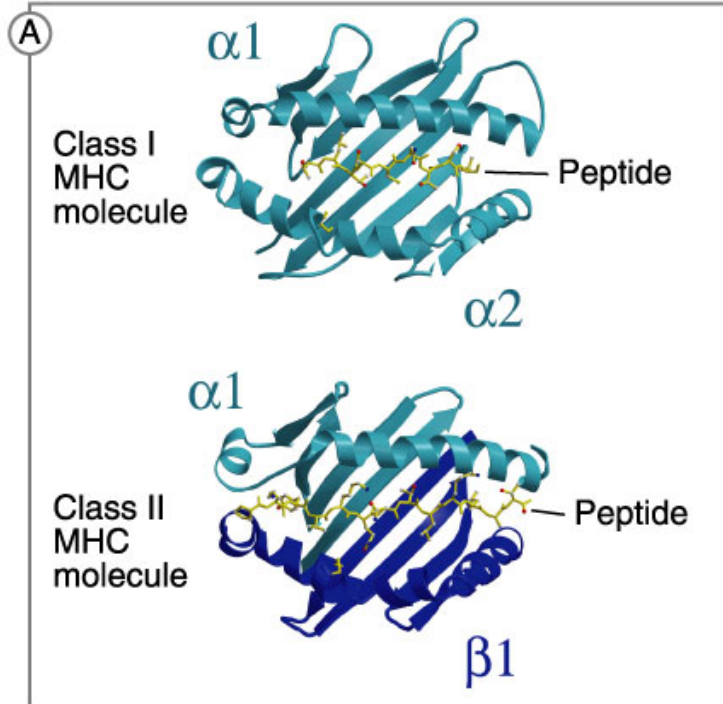
- One receptor or two receptors?
- Structure of the MHC provided the insight
- MHC molecules are designed to present peptides.



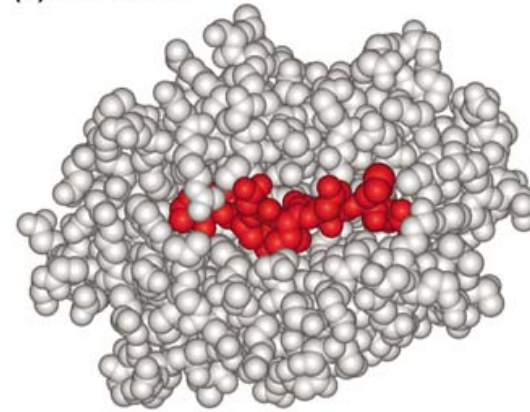
Abbas et al: Cellular and Molecular Immunology, 7e.  
Copyright © 2012, 2007, 2005, 2003, 2000, 1997, 1994, 1991 by Saunders, an imprint of Elsevier Inc.

*So, T cells simultaneously recognize a single peptide and MHC molecular complex!*

# Binding of Class I and Class II MHC Molecules to Peptide Ags

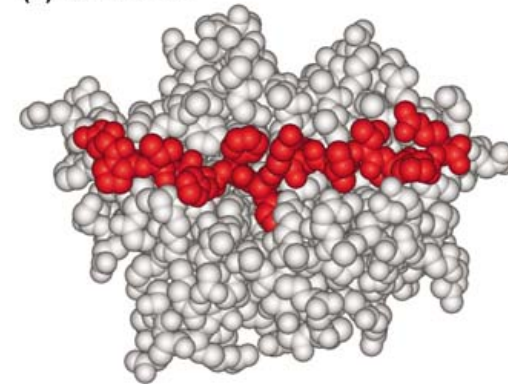


(a) MHC class I



© 1999–2007 New Science Press

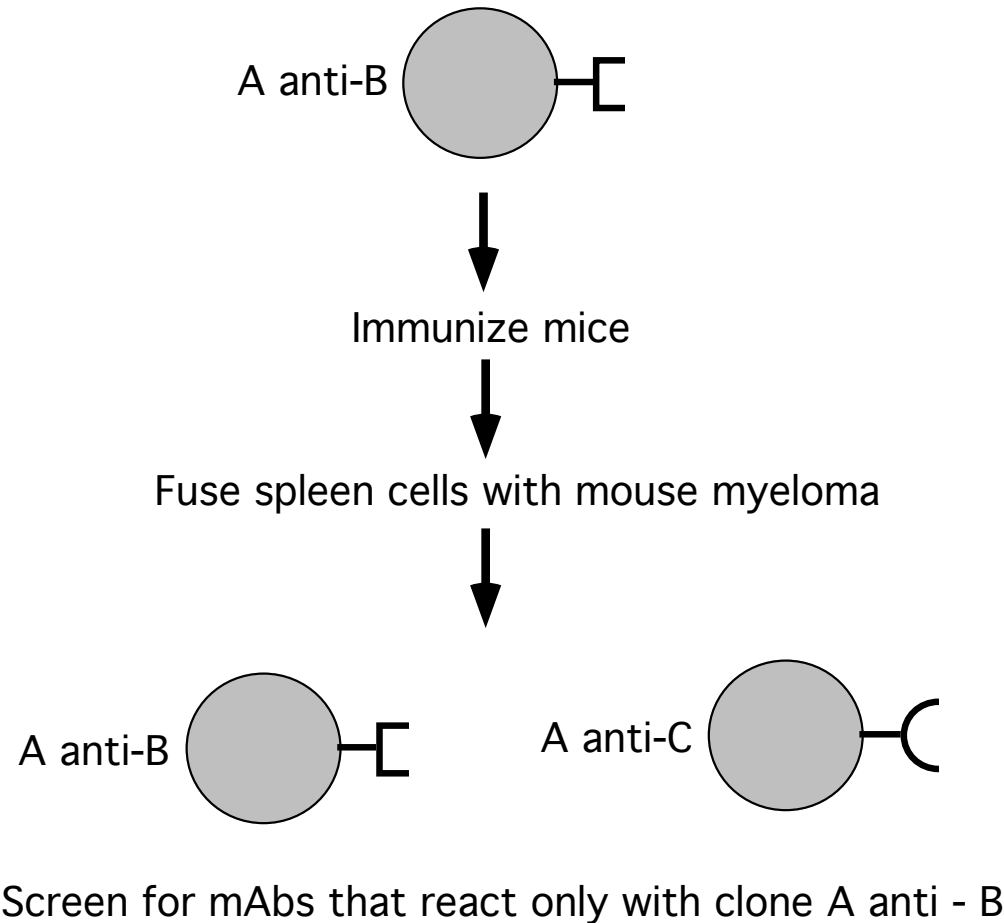
(b) MHC class II



# **Discovery and characterization of the TCR protein and genes**

# Identification of the TCR Protein

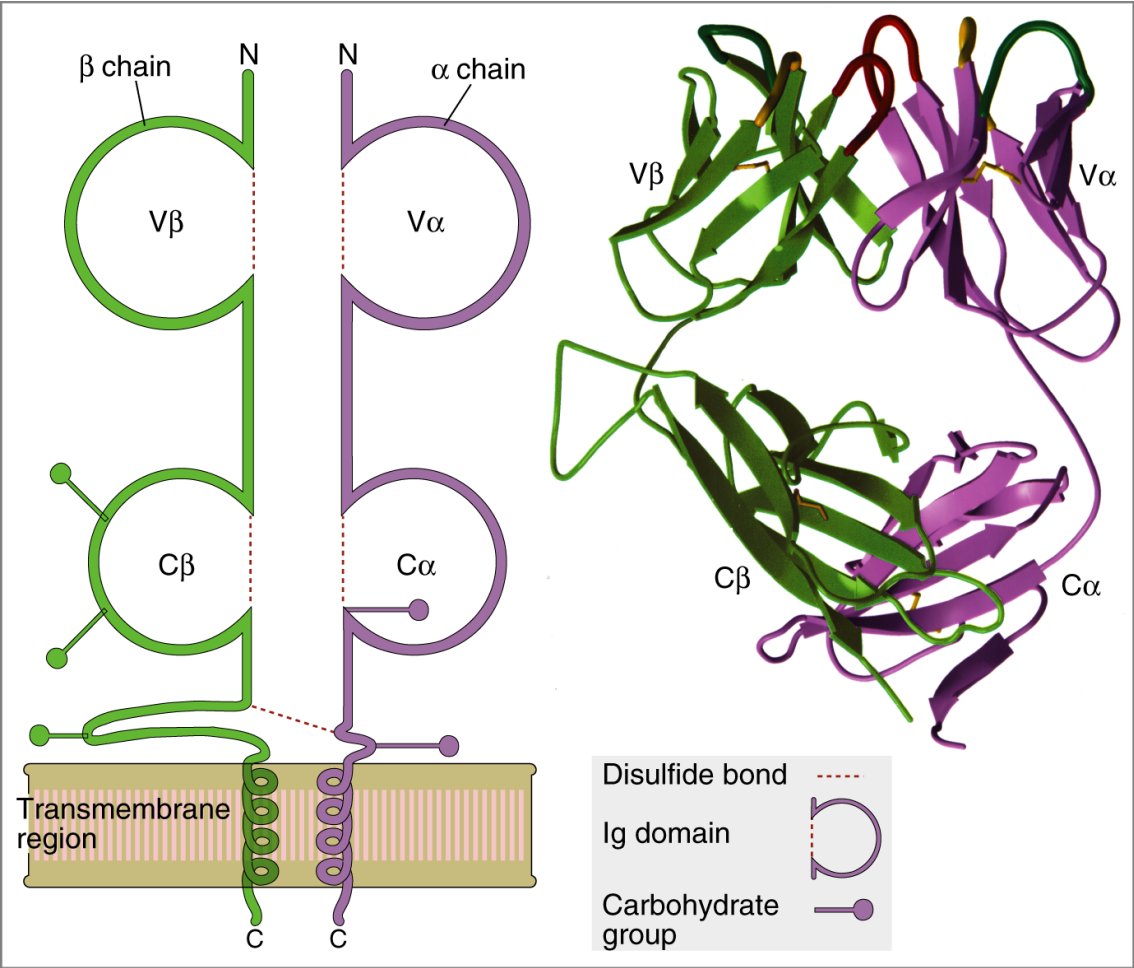
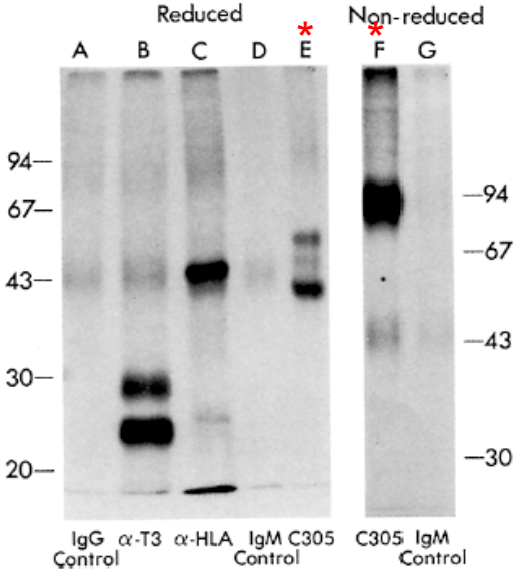
Generation of T cell *clone-specific* monoclonal antibodies (Allison, Reinherz, Kappler and Marrack, '82-' 83)



# Biochemical characterization of the TCR

Biochemical Characterization:

- 1. Disulfide-linked heterodimer
- 2. Transmembrane protein
- 3. Constant and variable regions
- 4. Both chains are glycoproteins



# Cloning the TCR $\beta$ -chain cDNA

Hedrick and Davis, 1984

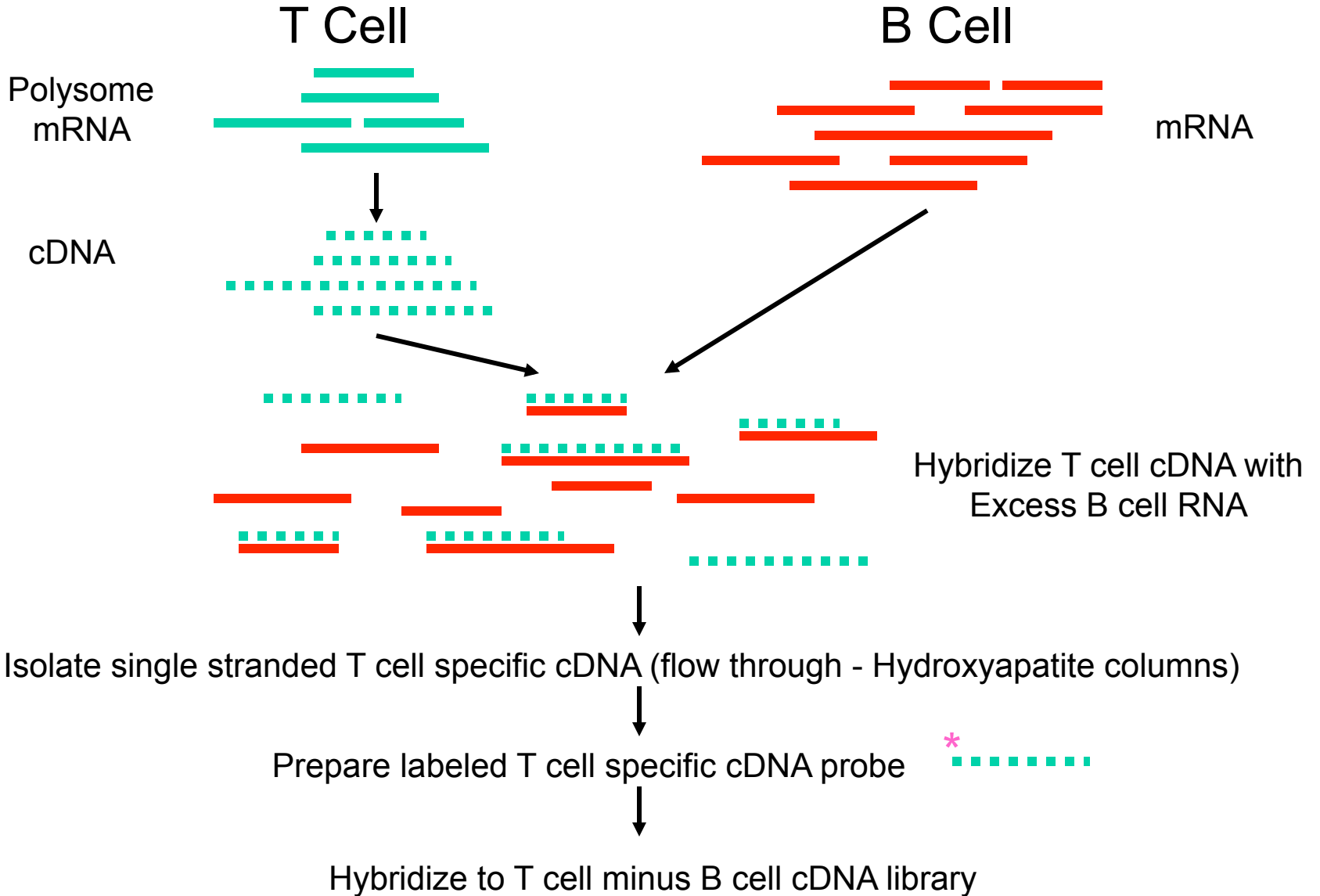
Yanagi and Mak, 1984

## Predictions:

1. T cell specific
2. Transmembrane protein
3. Genes should be rearranged in T cell but not in non-T cells
4. cDNA should encode **C**onstant and **V**ariable domains

# Isolation of TCR $\beta$ -chain cDNA

(Hedrick and Davis, 1984)



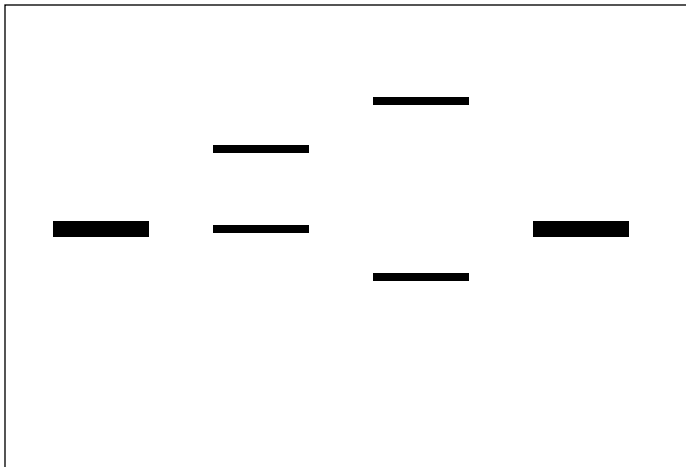
# TCR $\beta$ -chain cDNA



## Southern Blots: evidence for rearrangement (J-region probe)

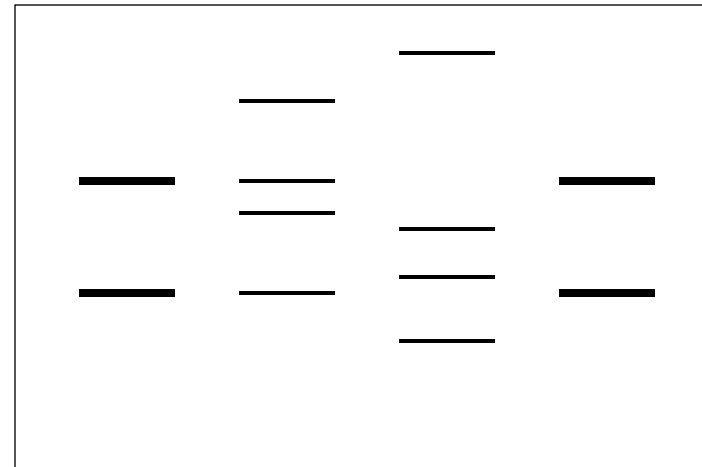
EcoR1 Digest

Liver    T cell clone 1    T cell clone 2    Kidney



BamH1 Digest

Liver    T cell clone 1    T cell clone 2    Kidney



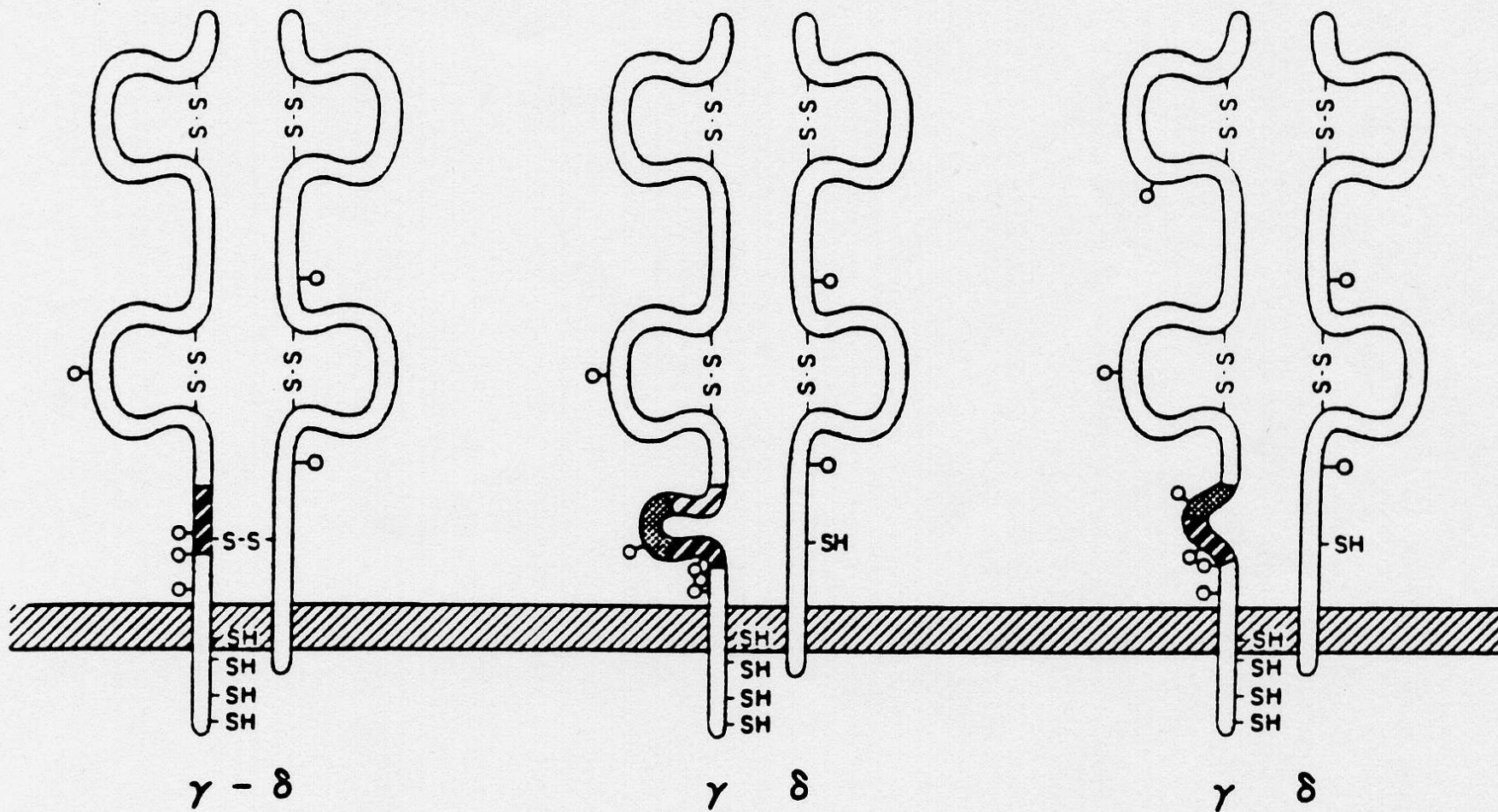


# THE $\gamma\delta$ T CELL RECEPTOR

FORM 1

FORM 2abc

FORM 2bc



PBL C1 ( $C_{\gamma 1}$ )

IDP2 ( $C_{\gamma 2}$ )

MOLT-13 ( $C_{\gamma 2}$ )

# $\gamma\delta$ T Cells

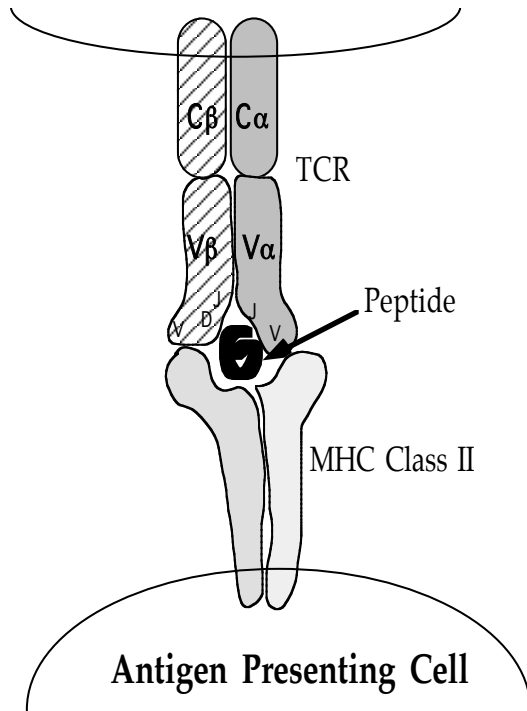
- Express  $\gamma\delta$  TCR heterodimer instead of the  $\alpha\beta$  TCR heterodimer
- Distinct lineage of T cells
- Most resting  $\gamma\delta$  T cells lack CD4 and CD8 coreceptors
- Activated  $\gamma\delta$  T cells can express CD8
- Minor subset in mouse and man (2-5%). Epithelial localization predominates.
- Expressed early in ontogeny
- MHC Restriction/recognition – little good evidence for “MHC restriction”, reactivity to some non-classical MHC molecule is well-documented, but there is no evidence for requirement
- Function:
  - Secrete lymphokines and mediate cytotoxicity
  - Role in bacterial infections (mycobacterial, and others)
  - Respond to non-peptidic ligands  
i.e. bacterial phospholipids, alkylamines,  
heat shock proteins,

# Both chains of the $\alpha\beta$ TCR heterodimer are involved in antigen and MHC recognition

$\beta$ -Chain	L	V	D	J	C
$\alpha$ -chain	L	V	J	C	

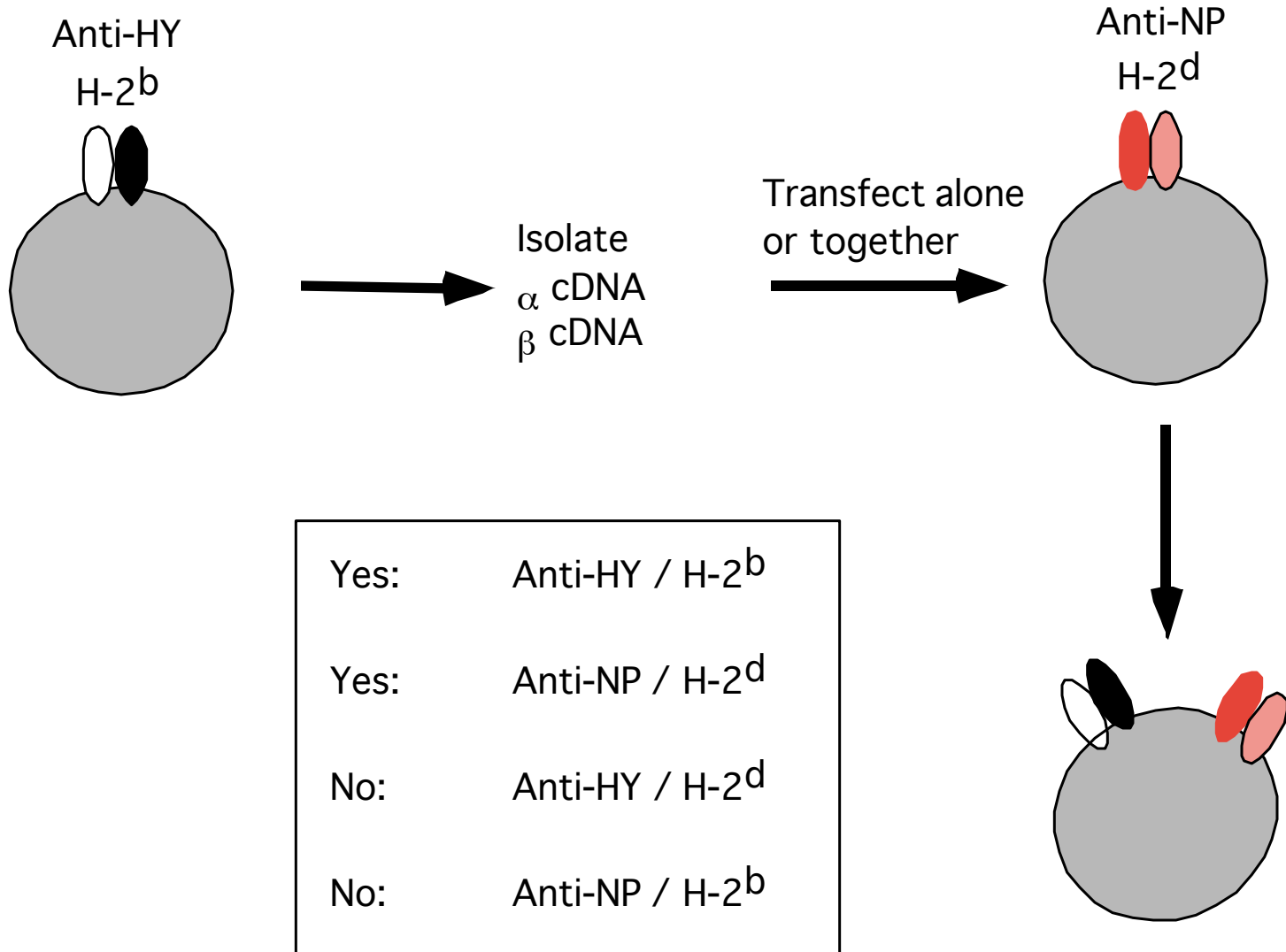
Expressed in T cells only  
 Clonally rearranged  
 Encodes a transmembrane glycoprotein  
 Homologous to light chains (Immunoglobulin fold)

T Cell

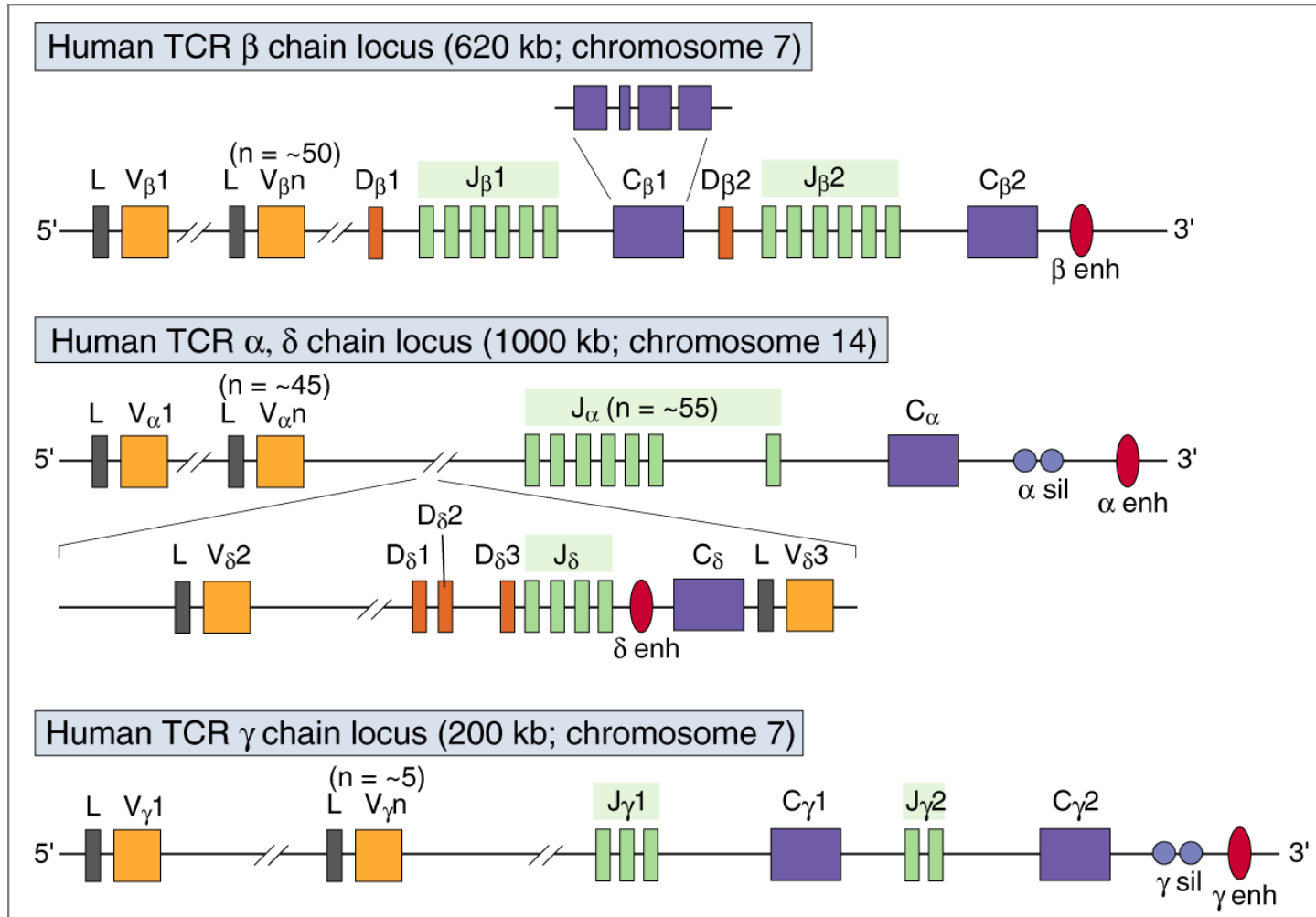


**$\alpha$  and  $\beta$  chains encode all of the information necessary for peptide and antigen specificity**  
 Transfection studies in T cell lines and hybridomas  
 TCR transgenic mice  
 **$\alpha$  and  $\beta$  chains both contribute to peptide and MHC molecule recognition**  
 Fusion of hybridomas

# $\alpha$ and $\beta$ chains of the TCR do not separately encode MHC or antigen specificity



# Human TCR gene loci



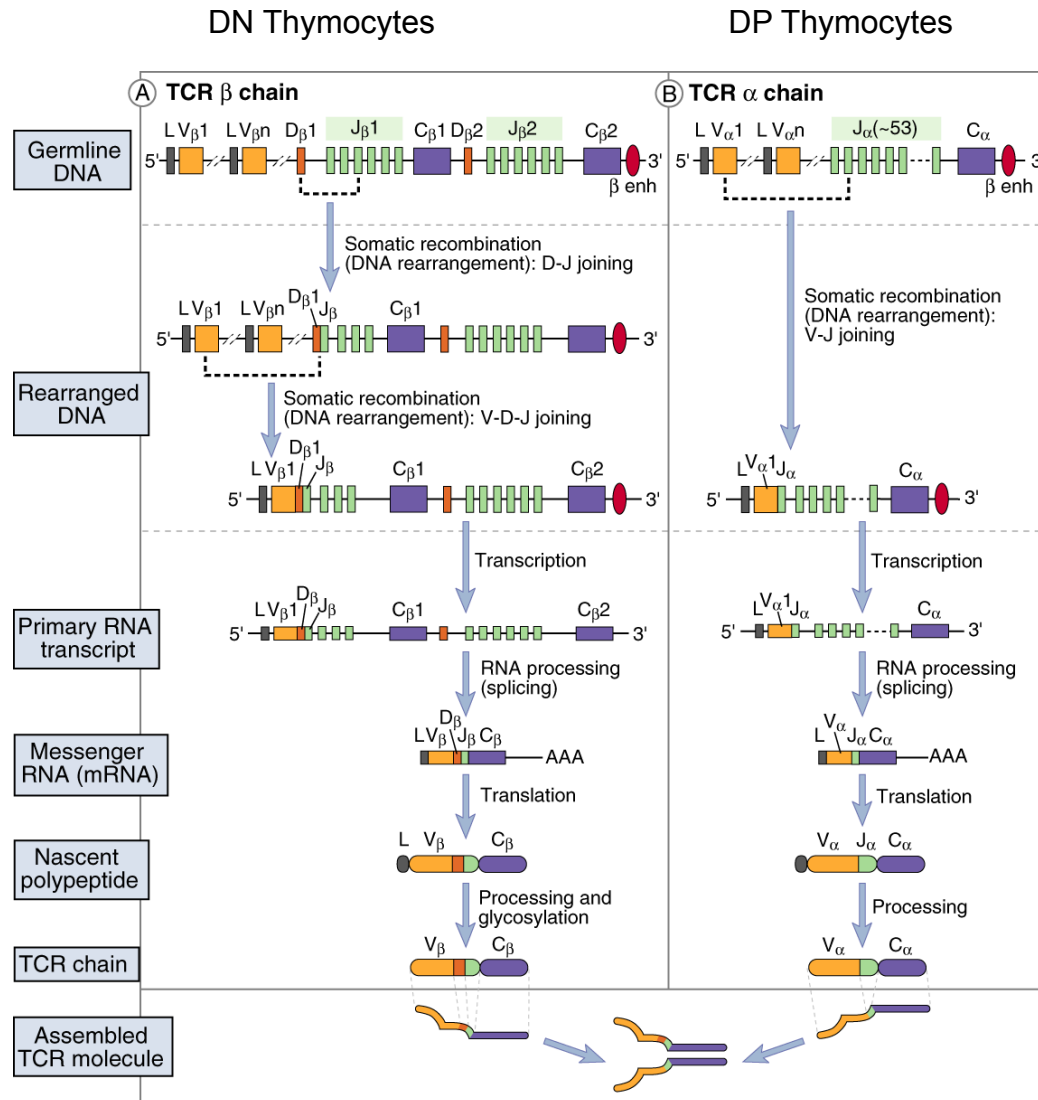
V segments 2 exons 80-250 nucleotides CDR1 and CDR2

J segments 1 exon 47-76 nucleotides

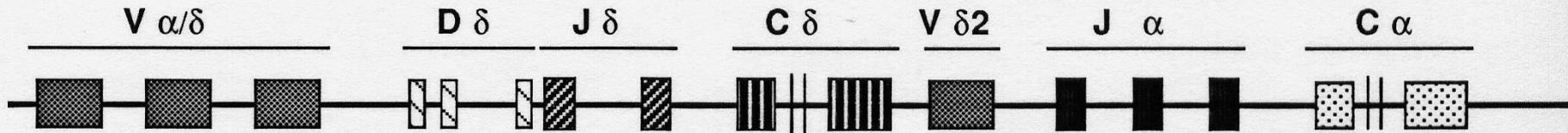
CDR3

D segments 1 exon 9-16 nucleotides

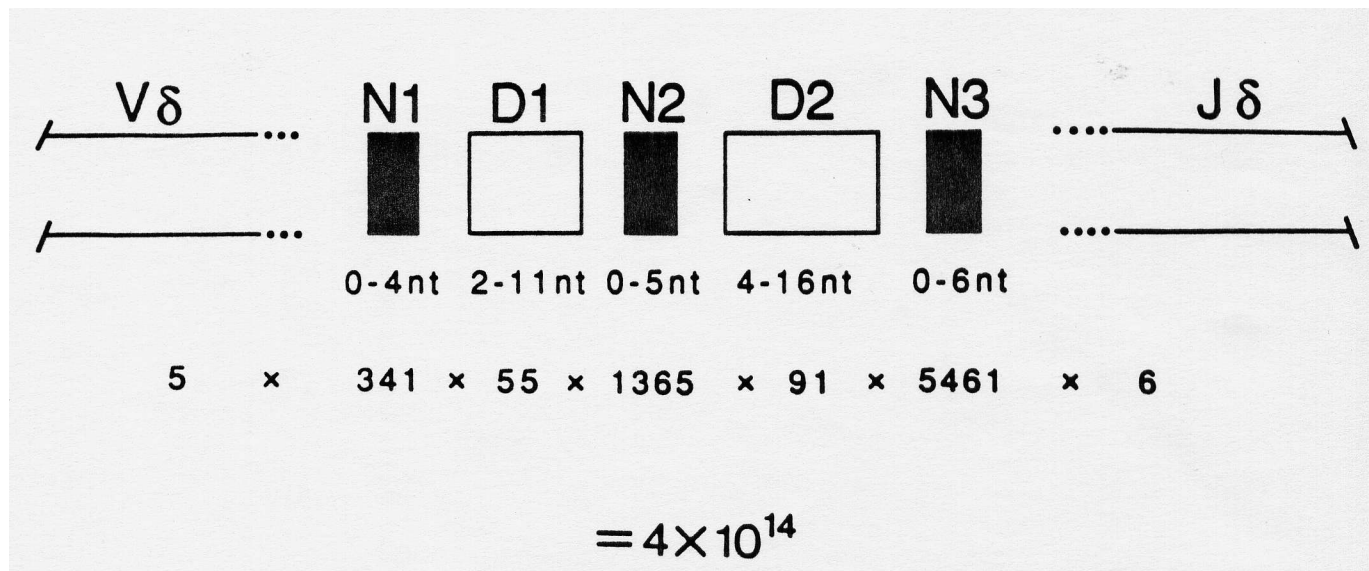
# TCR gene rearrangement occurs sequentially during T cell ontogeny



# Unusual organization of TCR gamma/delta genes



## Enormous potential of diversity in delta rearrangements



# Generating a diverse TCR repertoire

1. Recombination of different gene segments (V, D and J segments)
2. Recombination of different numbers of gene segments ( $\delta$  locus)
3. Imprecise joining of gene segments
4. “P” and “N” nucleotide addition (TdT)
5. Assembly of different combinations of rearranged  $\alpha$  and  $\beta$  chains

However, unlike immunoglobulin genes, **somatic mutation** of TCR genes does not take place.



# Comparison of diversity generated in TCR and BCR assembly

	Ig		TCR $\alpha\beta$		TCR $\gamma\delta$	
	H	L	$\alpha$	$\beta$	$\gamma$	$\delta$
Variable (V) segments	45	35	45	50	5	2
Diversity (D) segments	23	0	0	2	0	3
D' s in all frames	rarely	-	-	often	-	often
N-region addition	V-D, D-J	None	V-J	V-D, D-J	V-J	V-D1, D1-D2, D1-J
Joining segments	6	5	55	12	5	4
<b>Total potential diversity</b>	<b><math>\sim 10^{11}</math></b>		<b><math>\sim 10^{16}</math></b>		<b><math>\sim 10^{18}</math></b>	

# **Characteristics of TCR recognition of pMHC**

# Unusual features of TCR recognition of pMHC molecule complex

Simultaneous recognition of MHC specificity and peptide specificity

TCR affinity for peptide and MHC is very weak compared to antibodies:

**Kd of  $10^{-5}$  to  $10^{-7}$  M for TCR – dwell time of seconds for agonists**

Kd of  $10^{-7}$  to  $10^{-11}$  M for Ig

(Based on solution binding of monomers – flawed analysis)

Main determinant is off rate

TCR discrimination of self-pMHC vs agonist-pMHC is based on small differences in affinities (examples of only 10-fold)

Cell-Cell interaction context (avidity issues, coreceptors, particles/diffusion)

Tetramers of MHC/peptide can bind with higher *avidity*

Exquisite specificity despite low affinity:

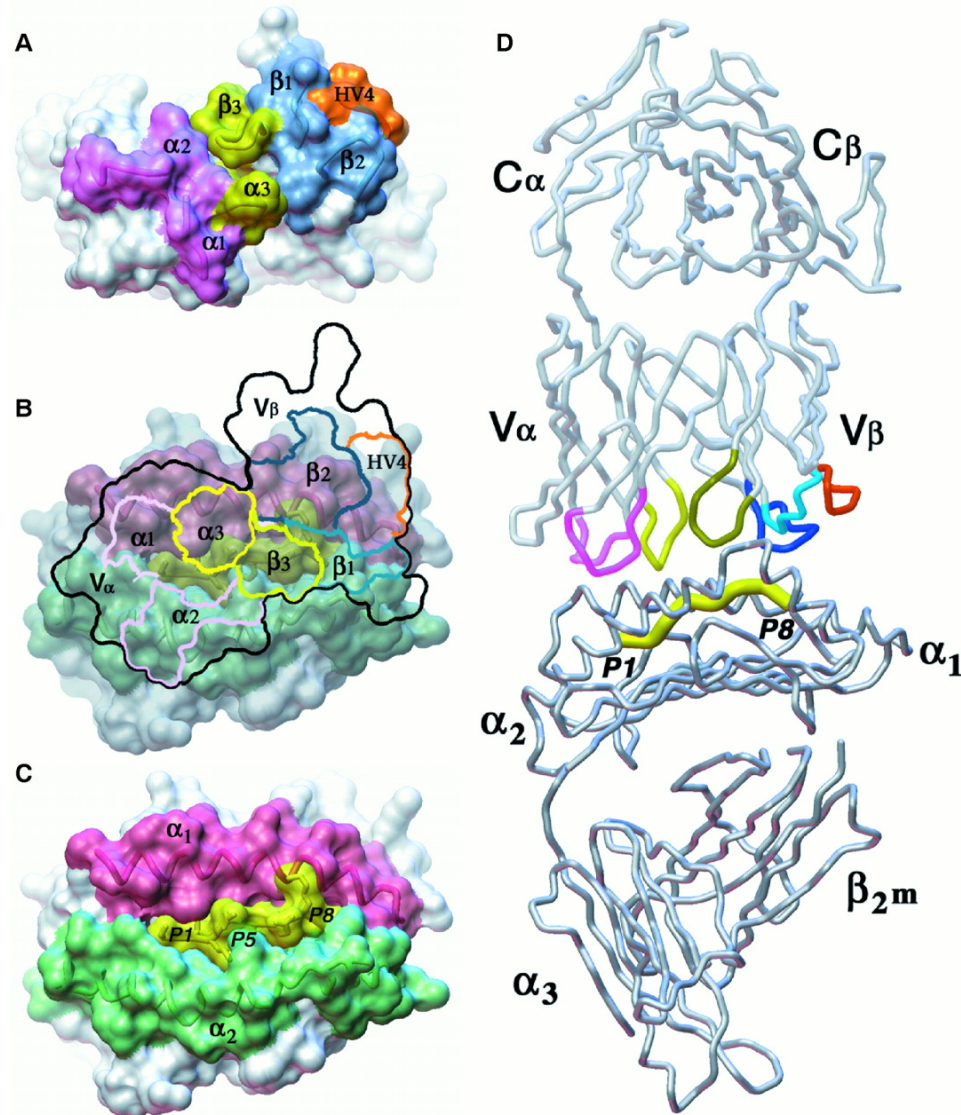
agonist peptides

altered peptide ligands

antagonist peptides

# Crystal structure of an $\alpha\beta$ TCR - Class I MHC/peptide

Garcia, et al., Science, 274:176, 1996



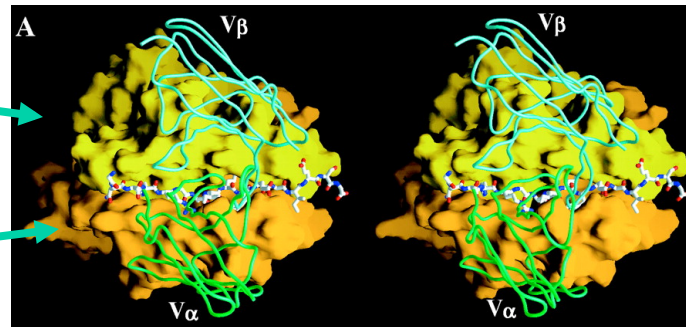
# Is the TCR and Class II MHC/peptide interaction oriented differently?

Reinherz, et al., Science, 286:1867, 1999

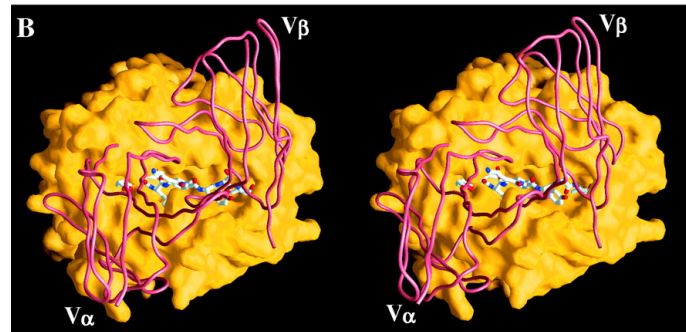
I-A alpha chain



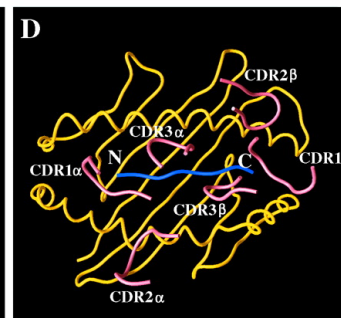
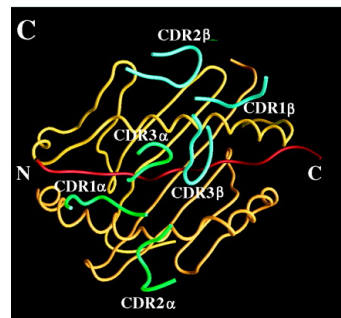
I-A beta chain



TCR-Class II peptide/MHC complex

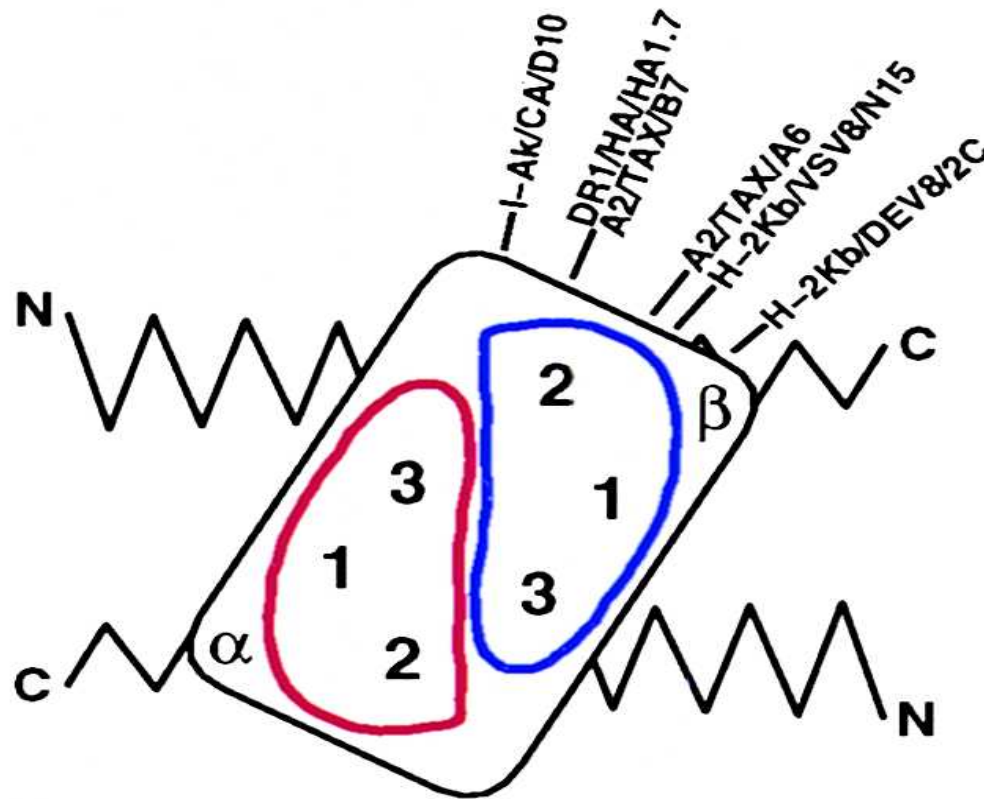


TCR-Class I peptide/MHC complex



# Distinct orientations of different TCR/MHC-peptide complexes

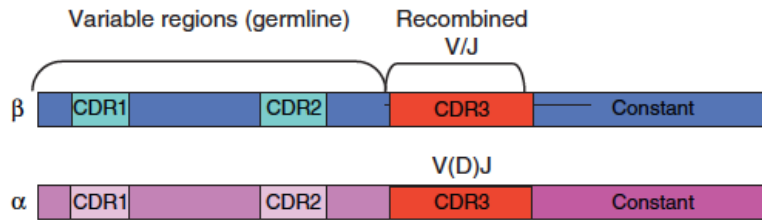
Hennecke and Wiley, Cell, 104:1, 2001



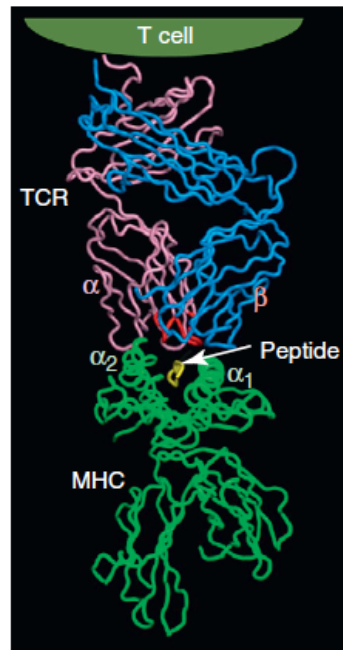
# CDR loops are involved in distinct recognition functions

Garcia, Trends Immunol., 2012

(a)

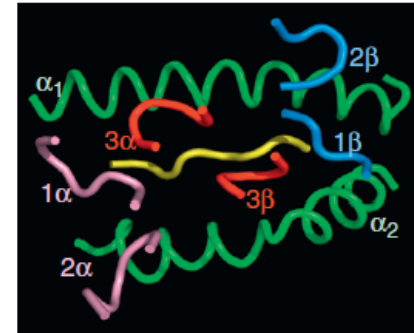


(b)



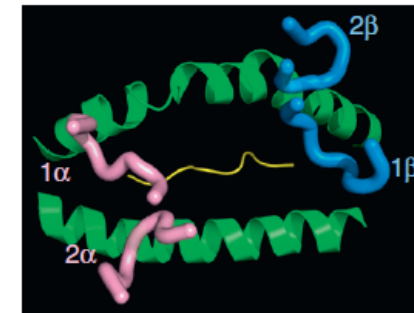
(c)

Canonical docking topology



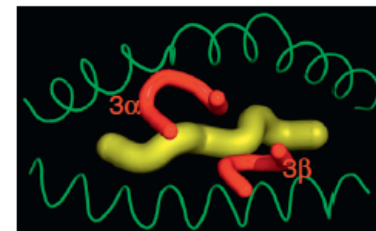
(d)

Germline-encoded recognition



(e)

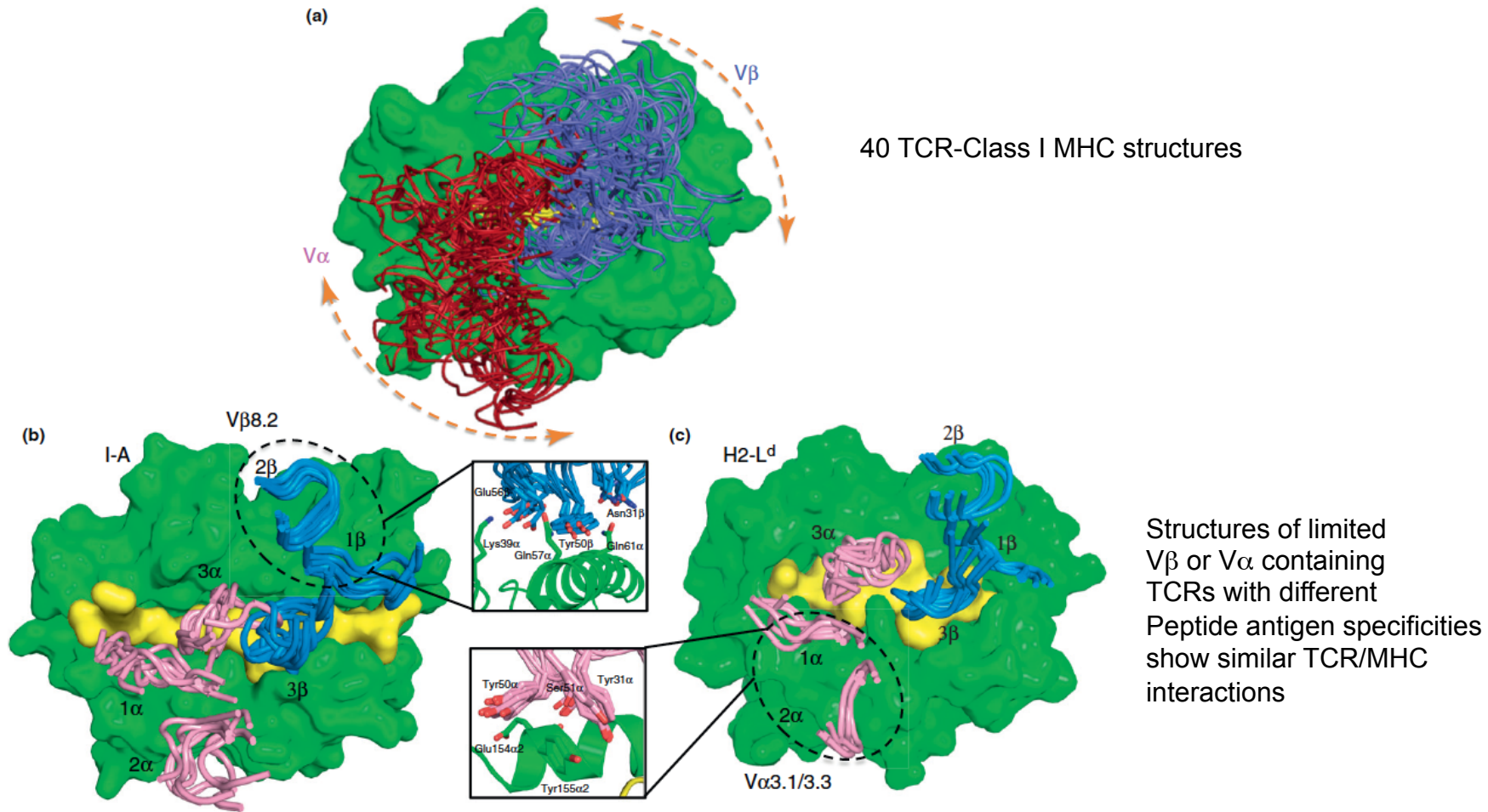
Imposed by selection/co-receptor





# $\alpha\beta$ TCR germline bias for MHC recognition

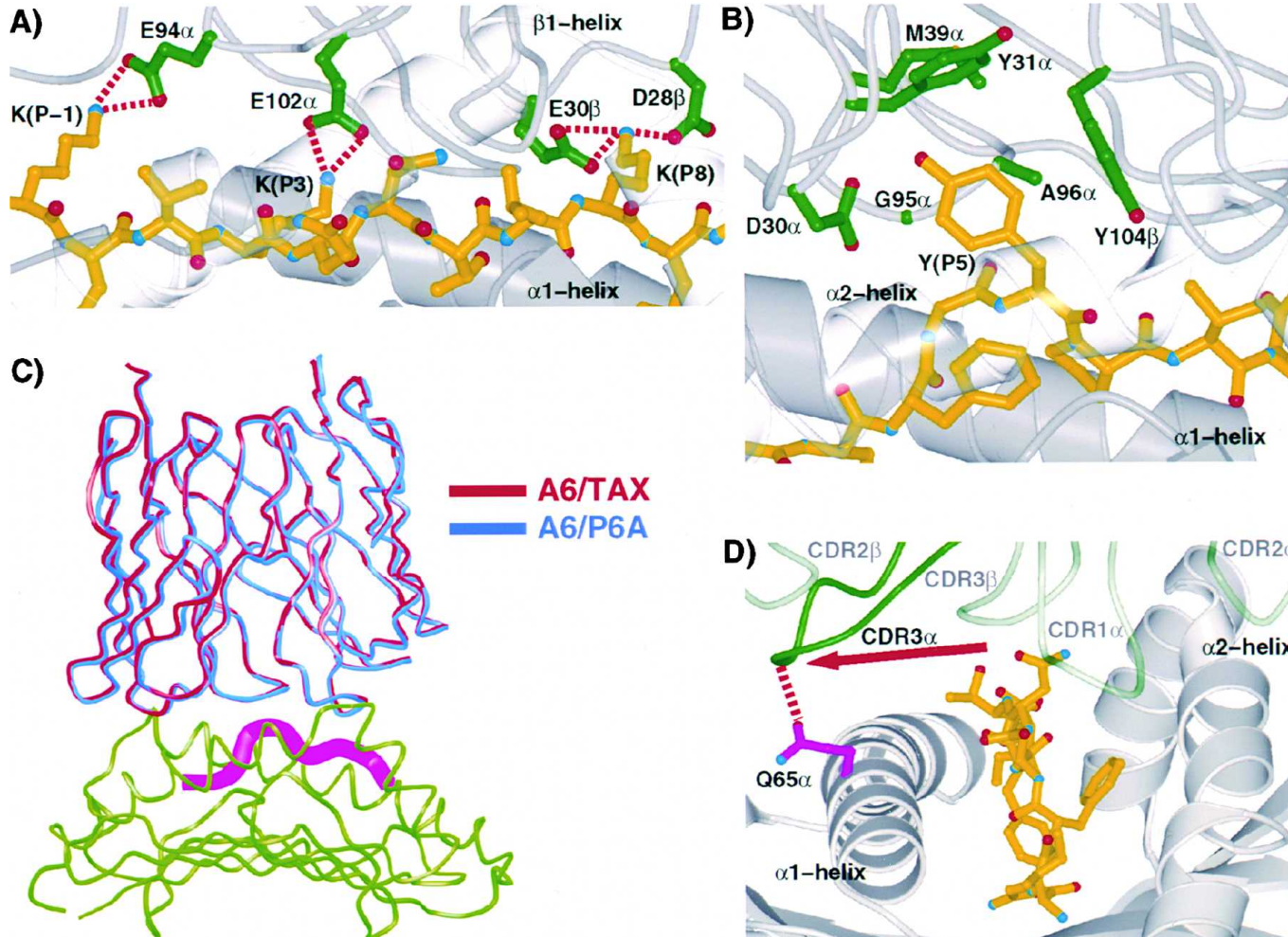
Garcia, Trends Immunol., 2012





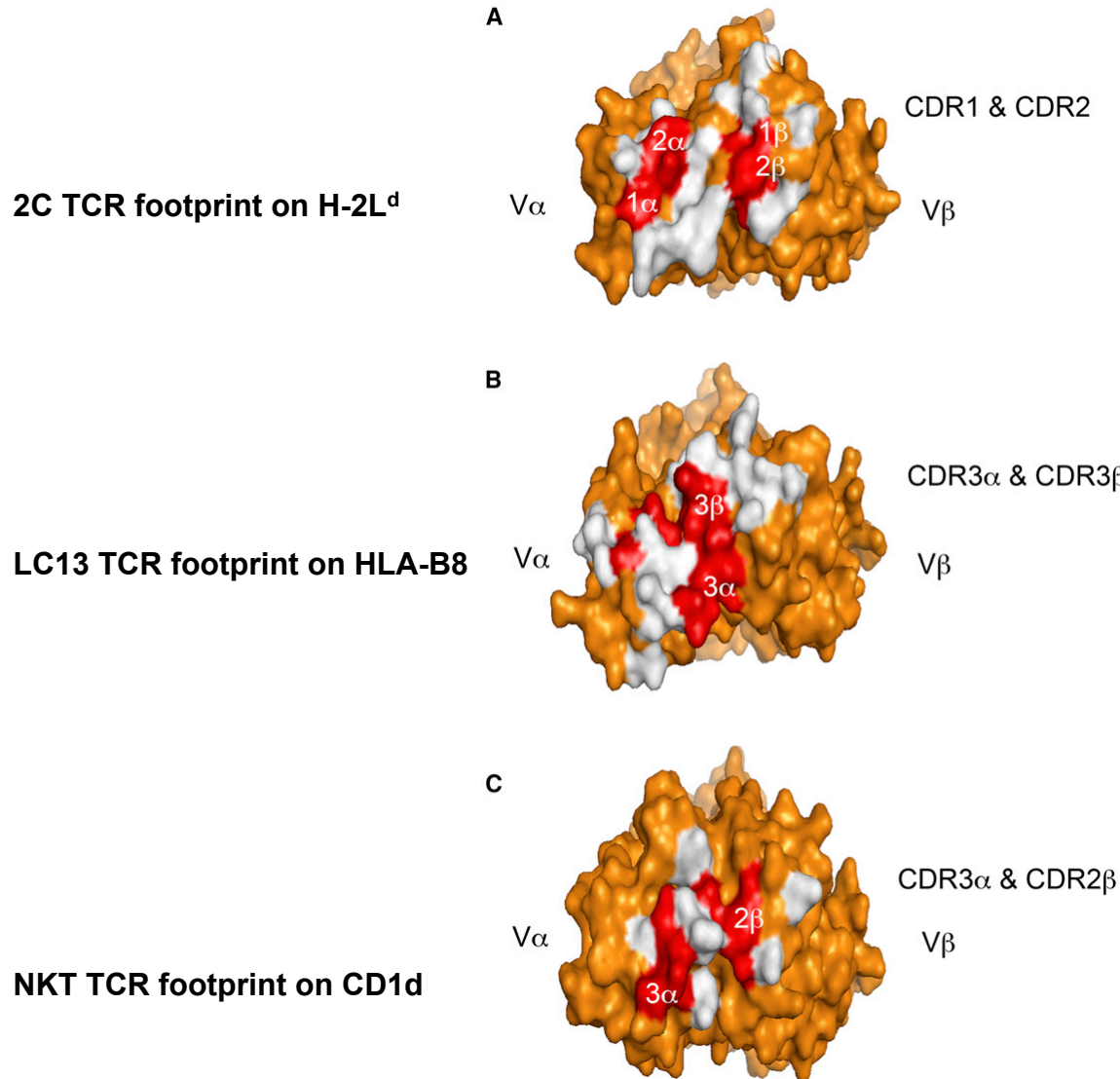
# The TCR can interact with MHC/peptide complexes via many different biochemical interactions

Hennecke and Wiley, Cell, 104:1, 2001

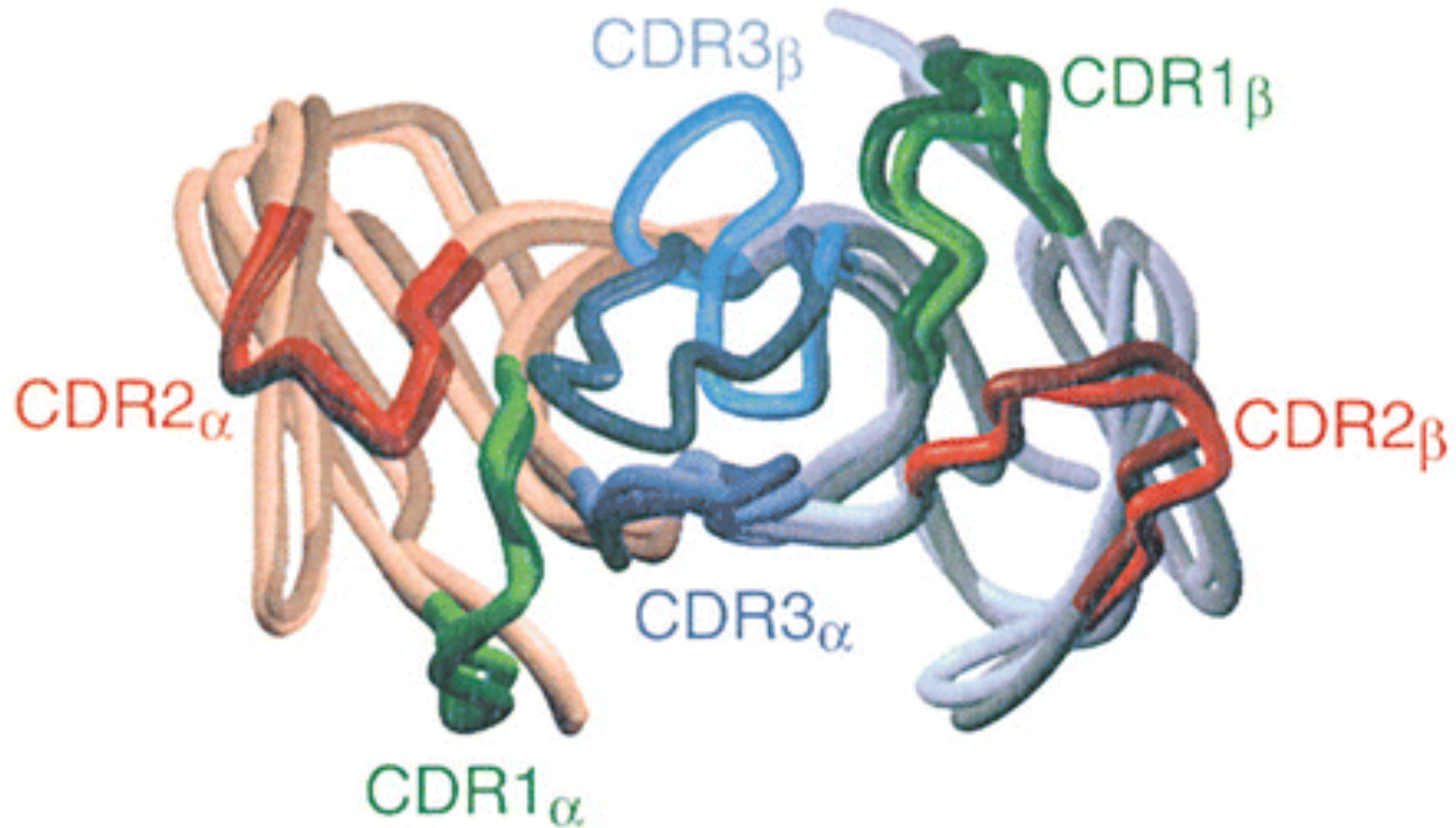


# Distinct structural and energetic ways that the TCR uses for antigen recognition

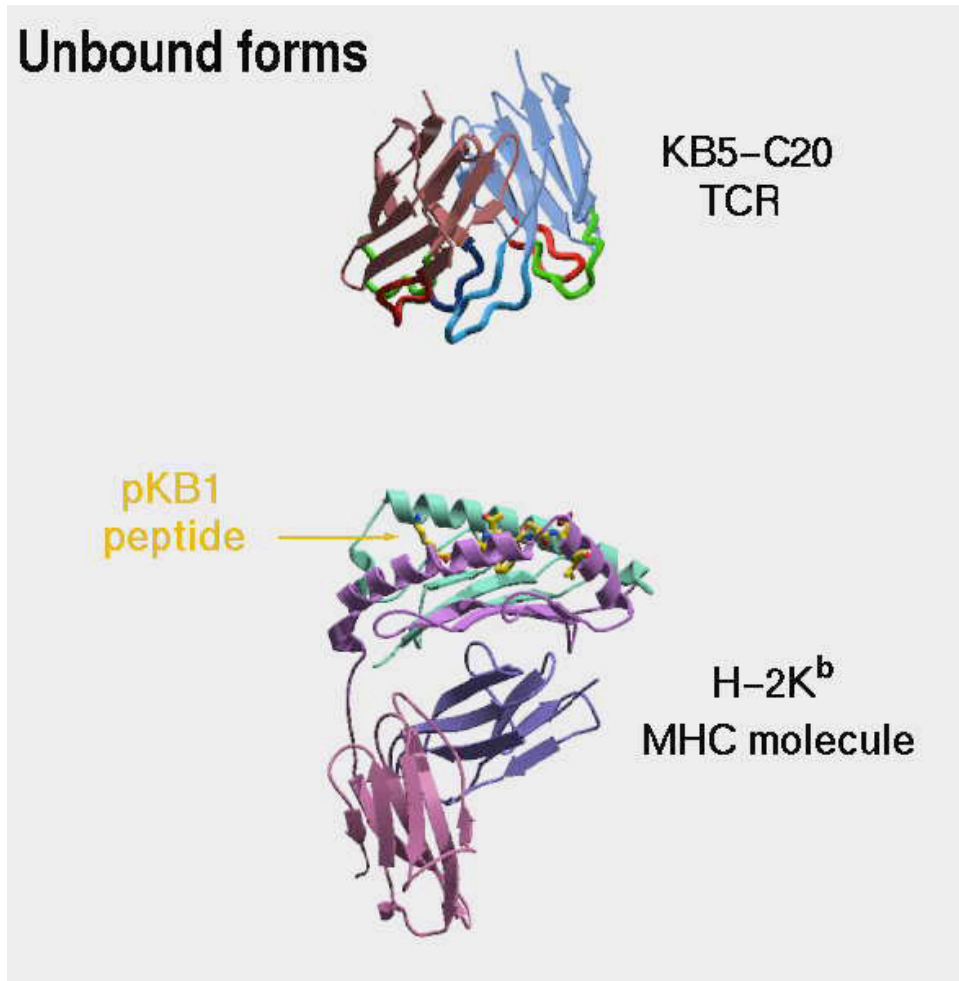
Godfrey, et al, Immunity 2008



# Distinct conformations of the TCR CDR3 loops in the ligand-unbound and bound states



# Model of a high degree flexibility in CDR3 during peptide/MHC docking



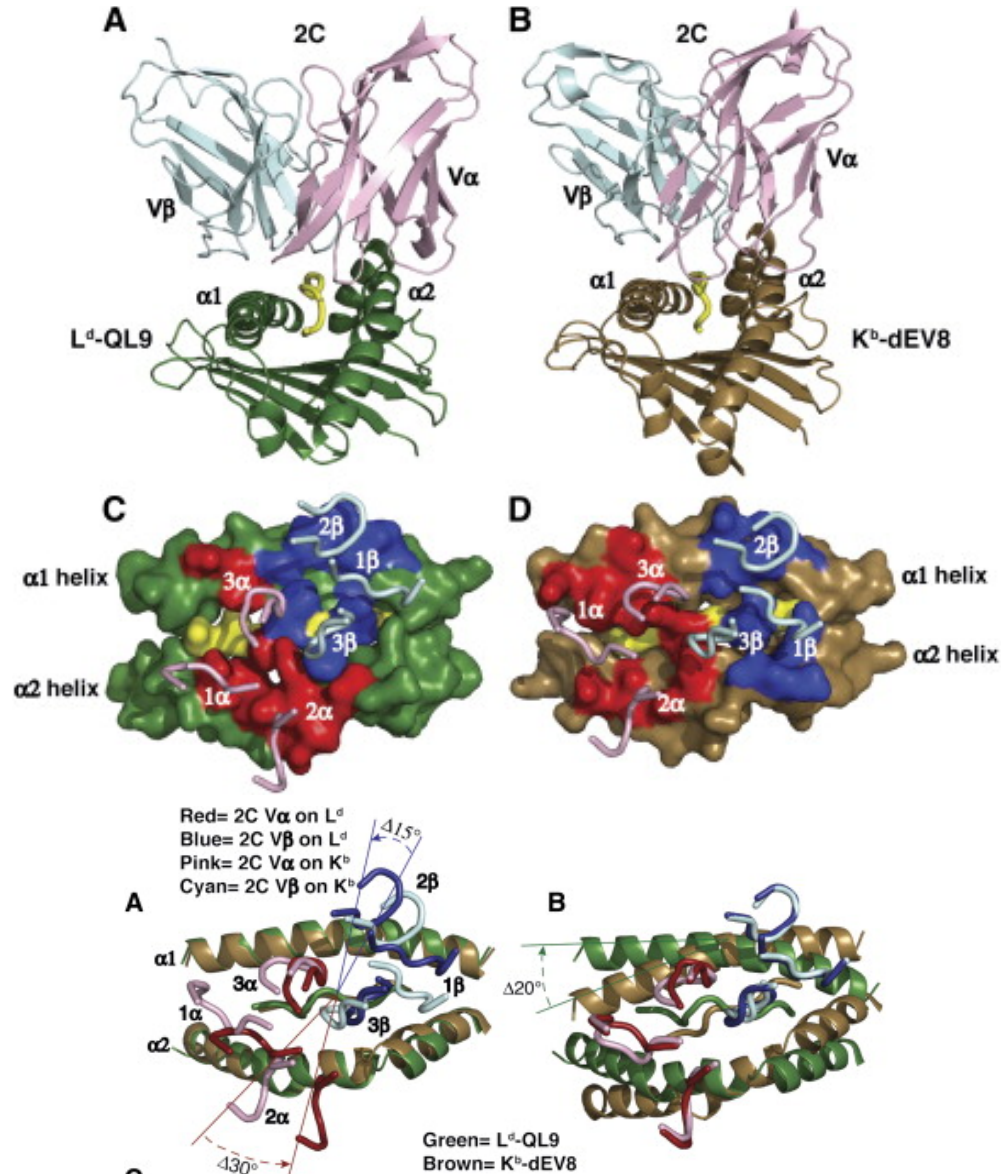


# The same TCR can adopt distinct conformations to be polyspecific

Colf, et al., Cell, 2007

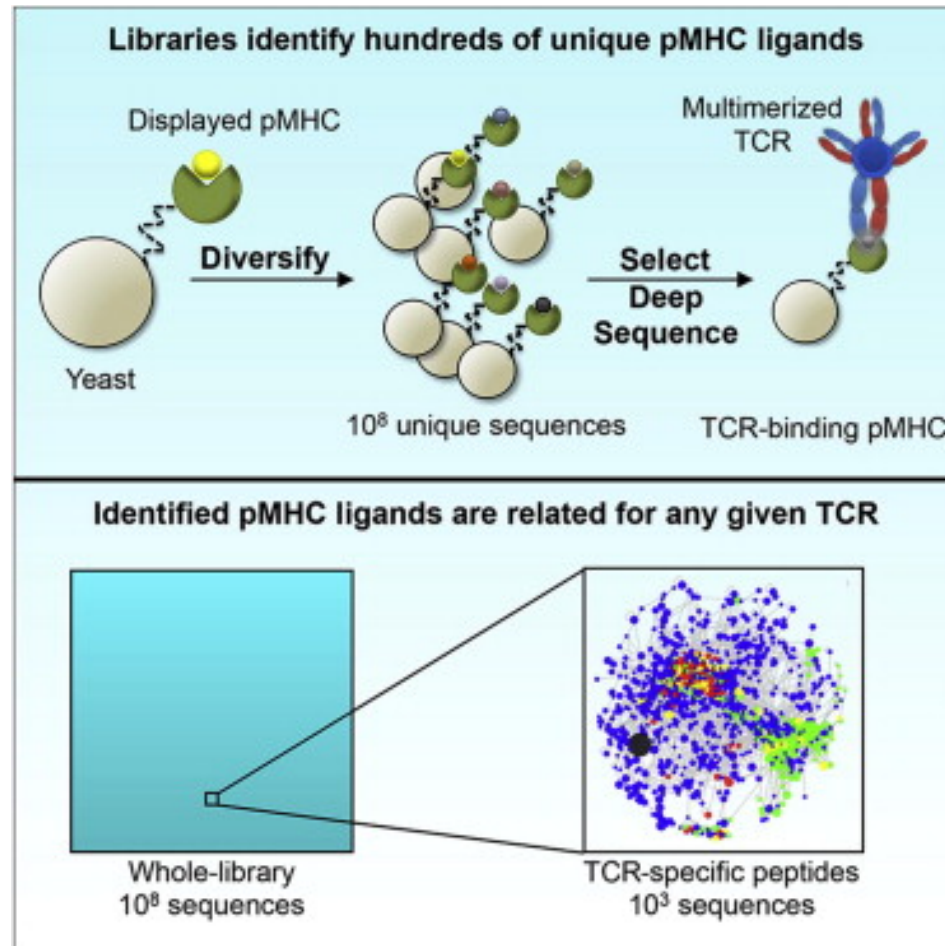
Allo-MHC  
plus peptide

Self-MHC  
plus peptide



# A Single TCR can Recognize Hundreds of Different Peptides Which Share Some Common Features

Michael E. Birnbaum, et al., Cell, 2014



**The TCR interacts with superantigens in  
a distinct manner**

# Superantigens

Bacterial enterotoxins

Staphylococcal, Streptococcal and Mycobacterial

Minor lymphocyte stimulating (Mls) antigen

Endogenous mouse retroviral products

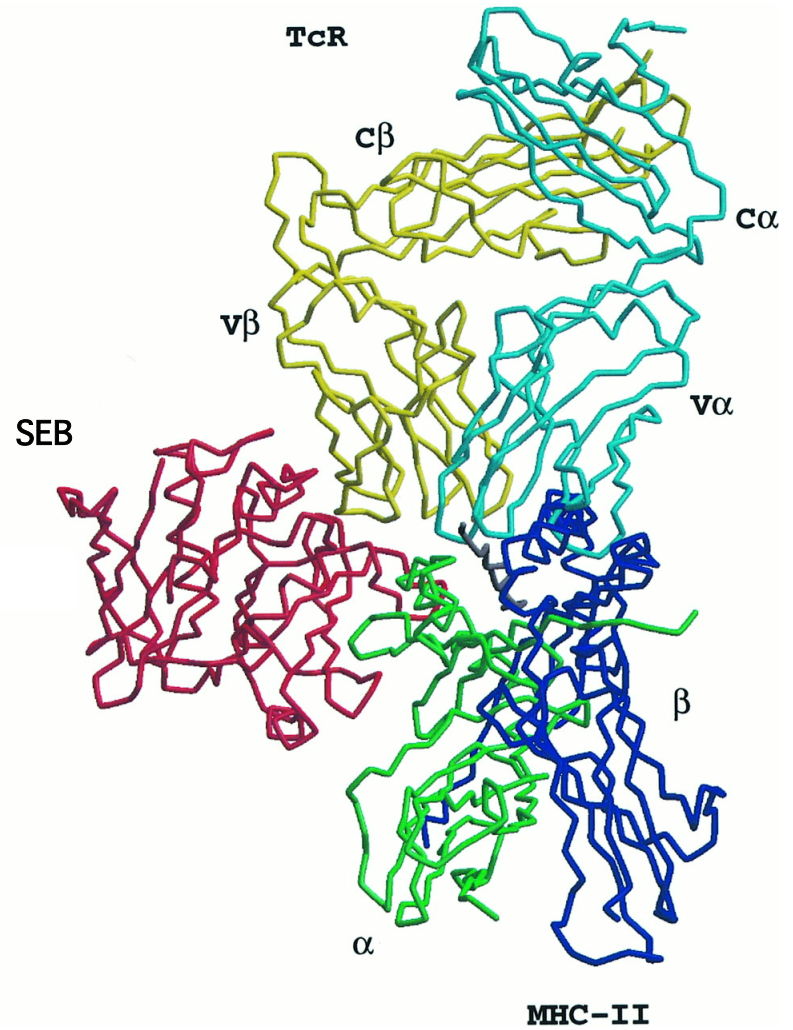
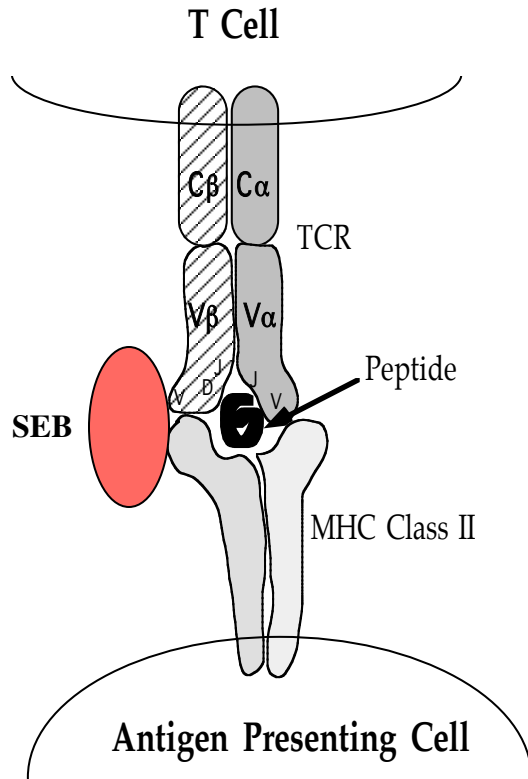
Unidentified endogenous antigens



# Comparison of Superantigens and Conventional Peptide Antigens

	<u>Conventional Antigens</u>	<u>Superantigens</u>
<b>Frequency of responsive T cells</b>	<b>1 in <math>10^4</math> to <math>10^5</math></b>	<b>1 in 4 to 20</b>
Interaction with the TCR	+	+
Interaction with MHC	+	+
MHC restricted recognition	+	-
<b>Requirement for processing</b>	<b>+</b>	<b>-</b>
<b>Binding to peptide groove in MHC</b>	<b>+</b>	<b>-</b>

# SEB/TCR/MHC structural model



# Superantigens have relative specificity for V $\beta$ segments

Toxin	V $\beta$ specificity	
	Human	Mouse
SEA	?	1, 3, 10, 11, 17
SEE	5.1, 6.1-3, 8, 18	11, 15, 17
SED	5, 12, ?	3, 7, 8.1-3, 11, 17
SEB	3, 12, 14, 15, 17, 20	3, 7, 8.1-3, 17
TSST1	2	3, 15, 17
ExFT	2	3, 10, 11, 15, 17
MAM	?	6, 8.1-3

Adapted from Marrack and Kappler, Science, 248:705, 1990

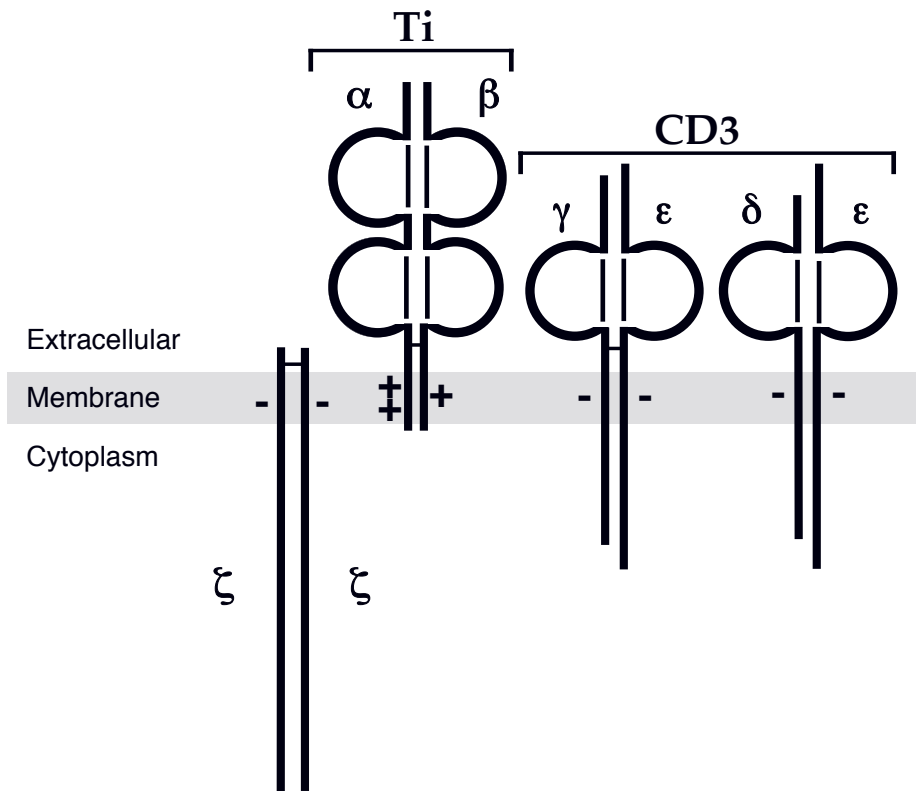
# Diseases Caused by Superantigens

Toxin	Organism	Disease
Staphylococcal enterotoxins (SE) A, B, C1, C2, C3, D and E	<i>S. aureus</i>	Food poisoning, Shock
Toxic Shock Syndrome Toxin (TSST1)	<i>S. aureus</i>	Toxic Shock Syndrome
Exfoliating Toxins A and B	<i>S. aureus</i>	Scalded Skin Syndrome
Pyrogenic exotoxins A, B, C	<i>S. pyogenes</i>	Fever, Rash, shock
<i>M. arthritides</i> mitogen	<i>M. arthritides</i>	Shock

Adapted from Marrack and Kappler, Science, 248:705, 1990

**The TCR is an oligomer**

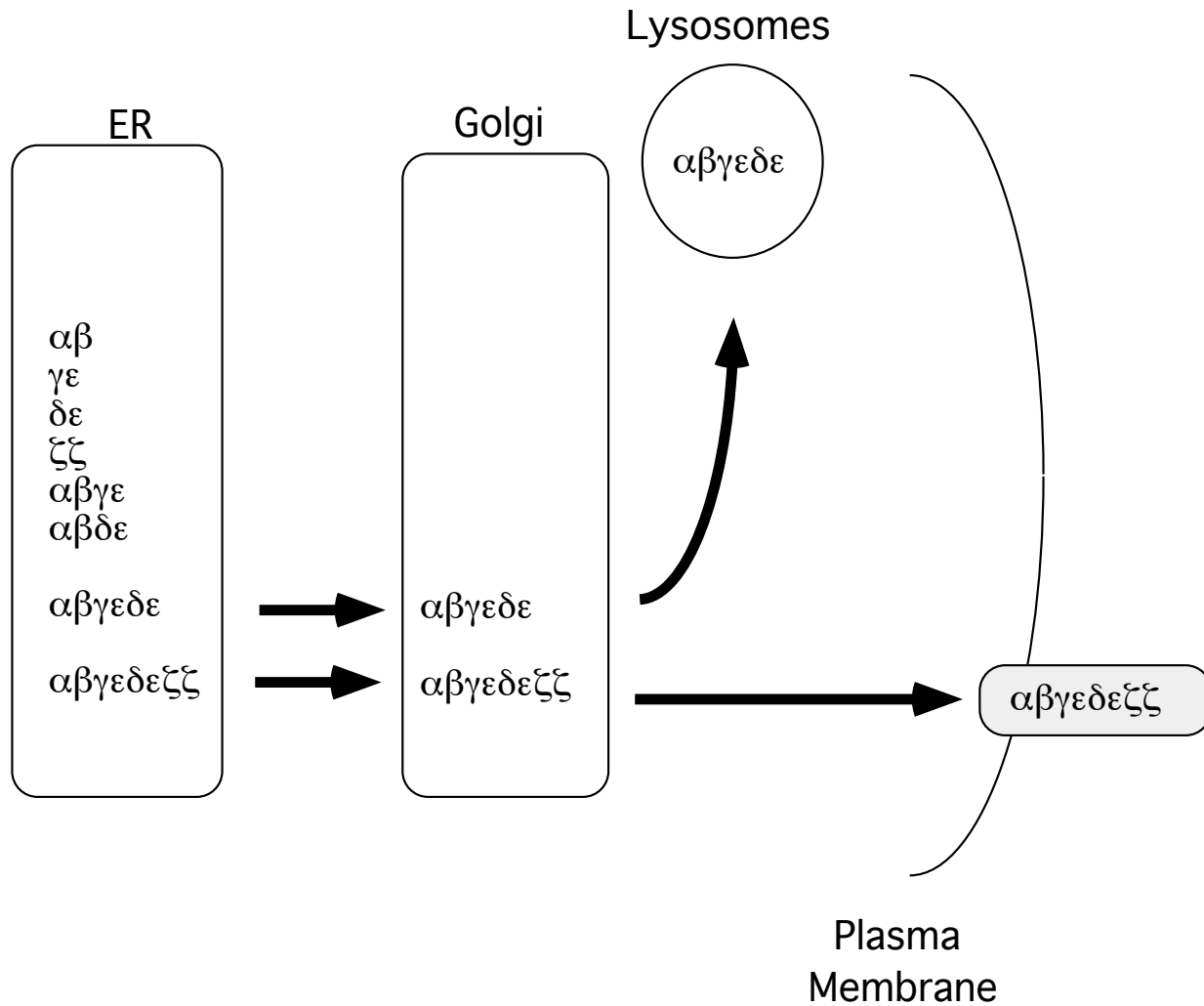
# The TCR is an oligomer



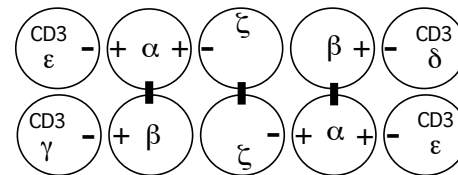
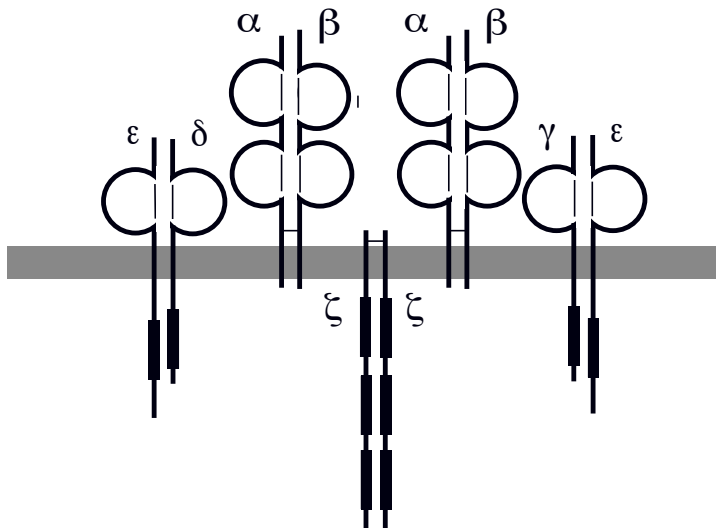
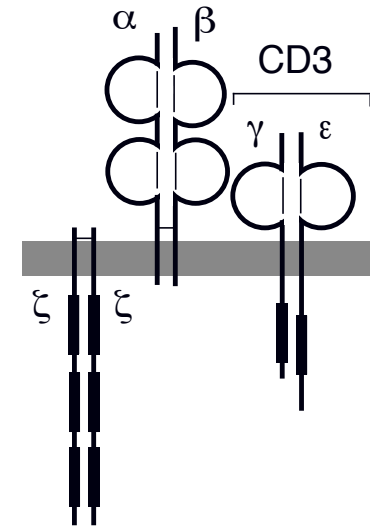
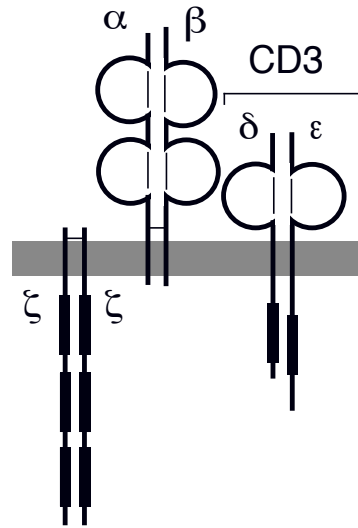
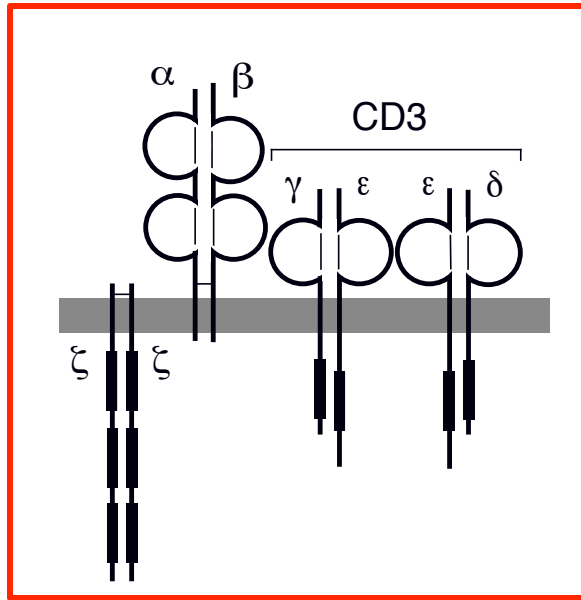
## Evidence:

1. Cointernalization of the CD3 and  $\alpha\beta$  heterodimer
2. Coimmunoprecipitation (very detergent dependent)
3. Chemical cross-linking ( $\beta$  and CD3  $\gamma$ )
4. Mutants (high CD3 expression requires  $\alpha\beta$ ,  $\gamma\delta$  or pre-TCR)
5. In vitro assembly studies

# TCR Assembly: Ordered Interactions and Quality Checkpoints

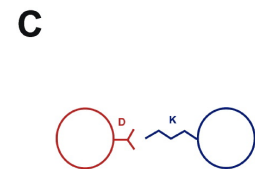
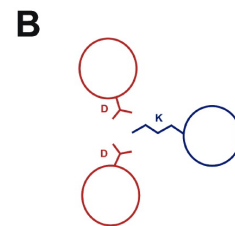
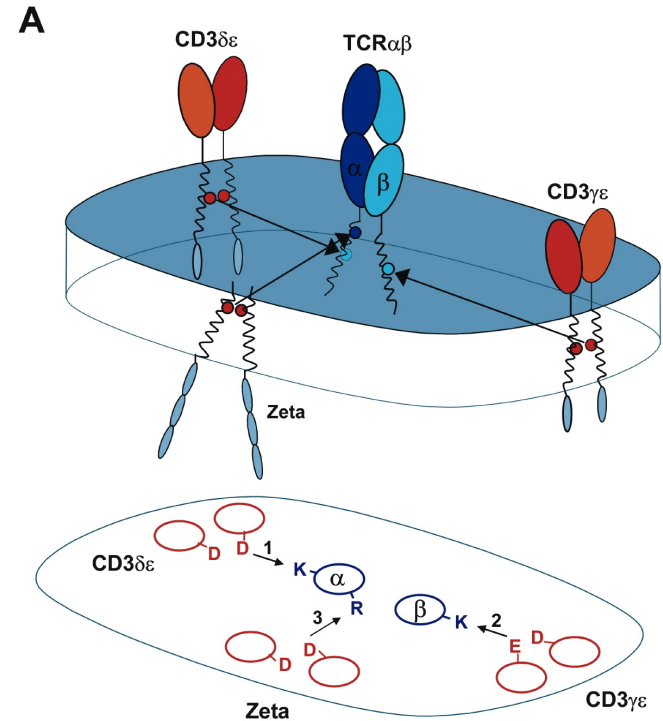
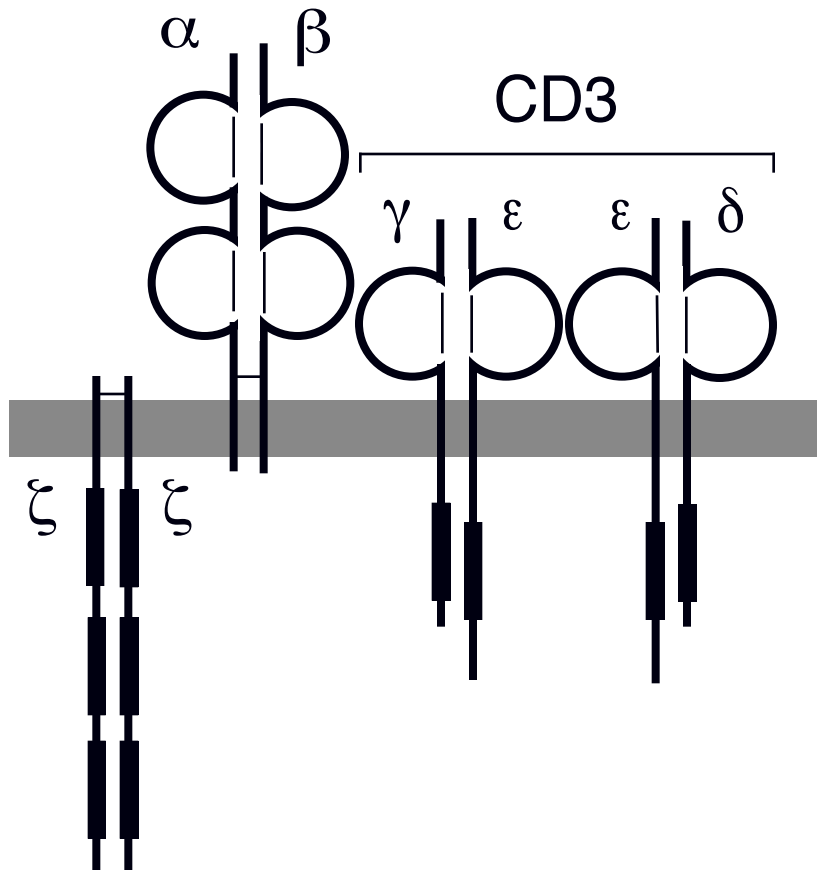


# TCR Stochiometry: Models



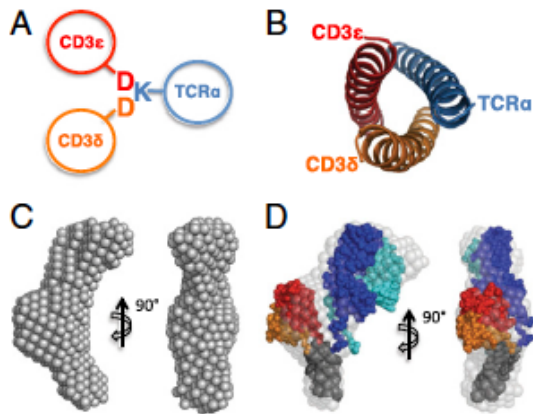


# In Vitro Assembly Favors a Single $\alpha\beta$ Heterodimer per TCR and Unusual Transmembrane Interactions

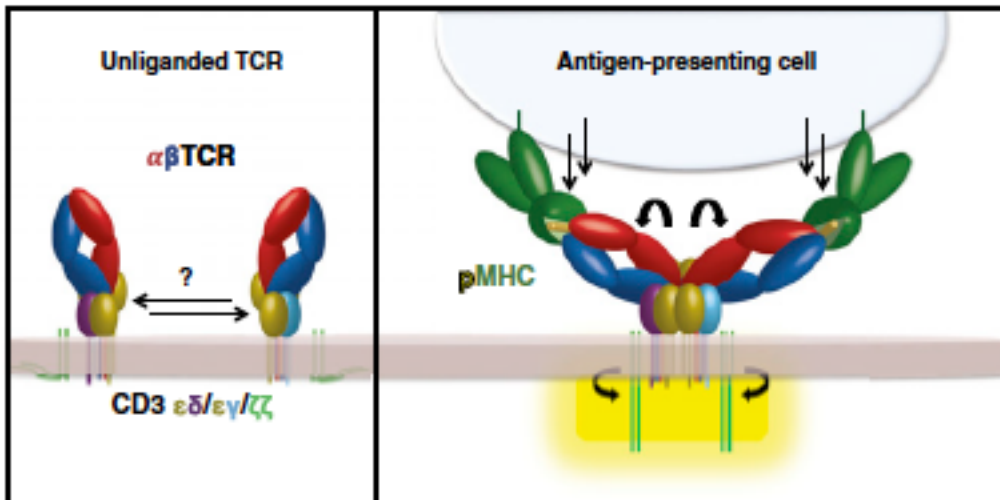
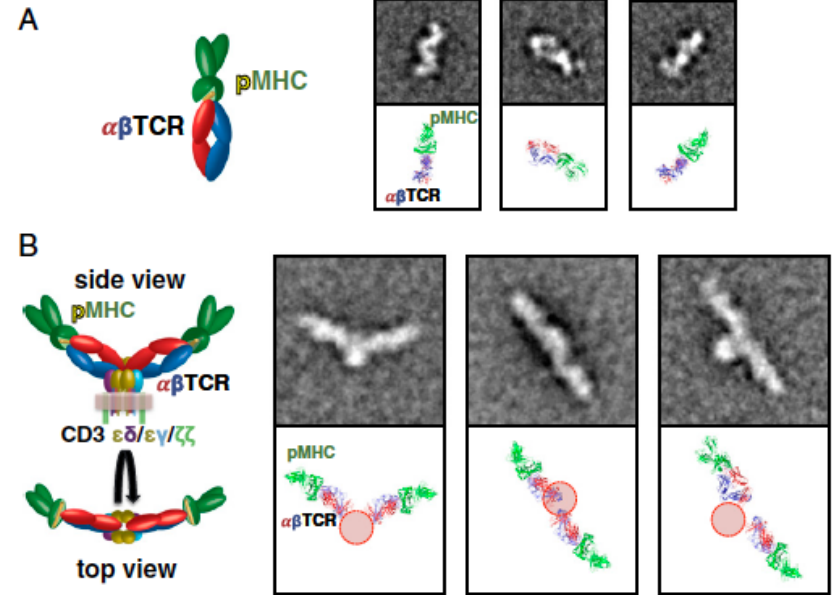


# Recent Biophysical Studies Suggest CD3 Chains Sit Below TCR $\alpha\beta$ Chains and the ligated TCR may Dimerize

Small angle X-ray scattering (SAX)



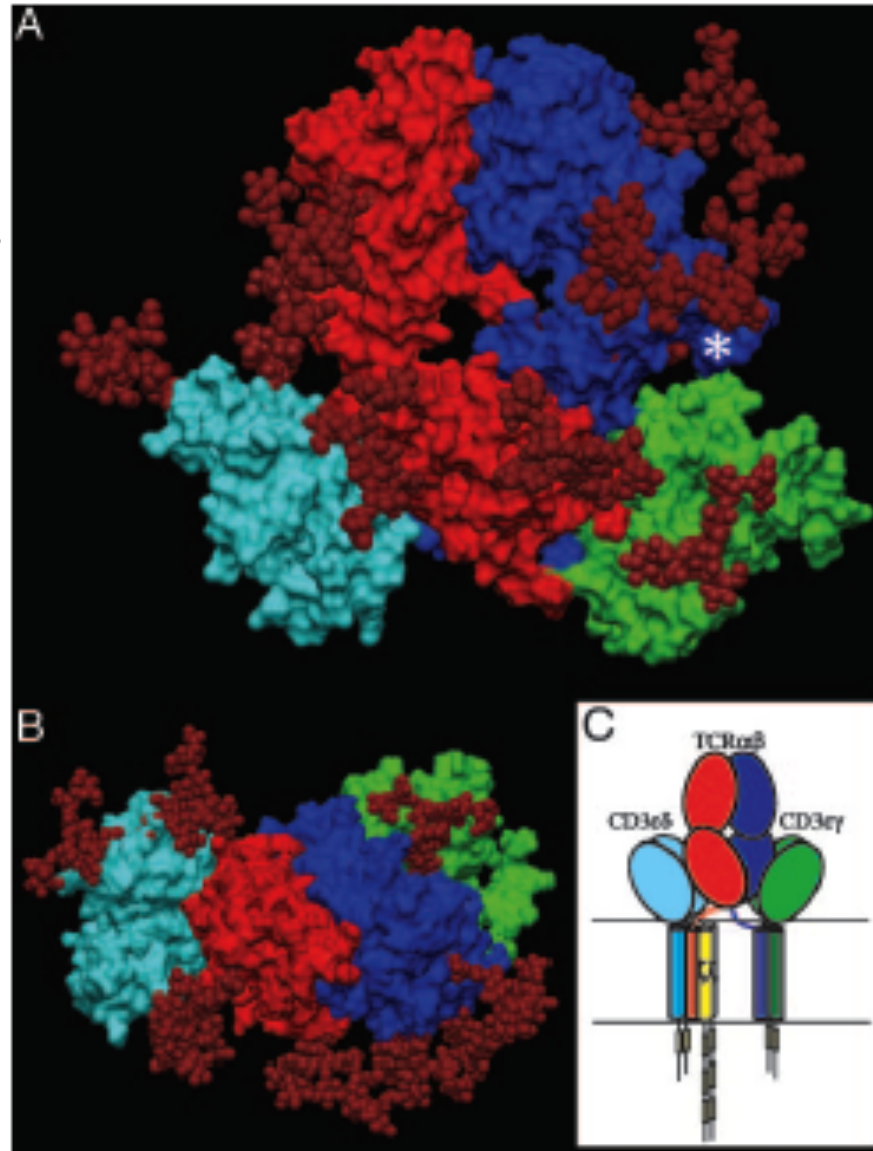
Negative Stain EM



# Model of TCR $\alpha\beta$ Heterodimer - CD3 complex

Sun, et al, PNAS, 2004

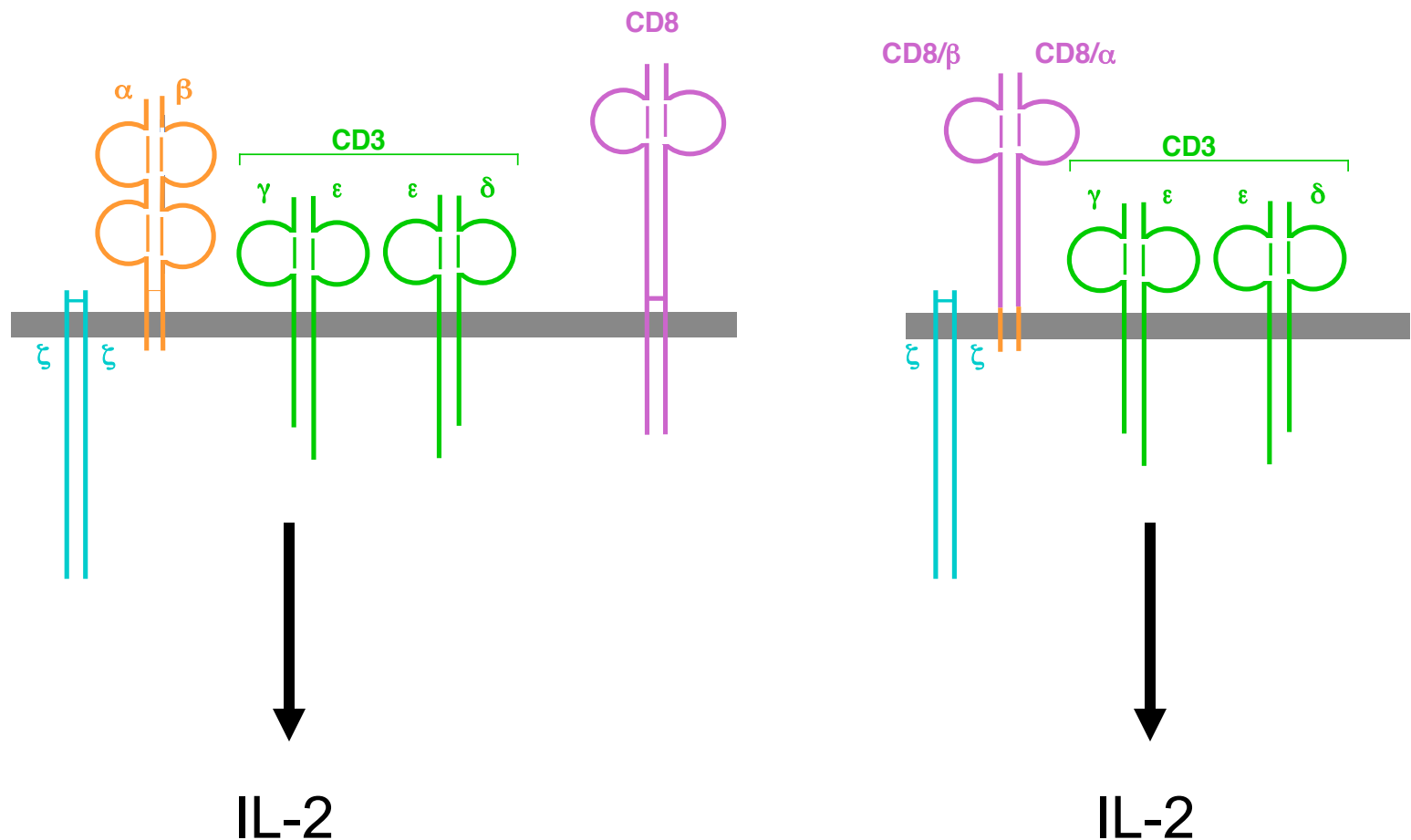
CD4 and CD8  
would be on this  
side - based on  
TCR and MHC  
interactions



**The CD3- and  $\zeta$ -chains are the signaling components of the oligomeric TCR**

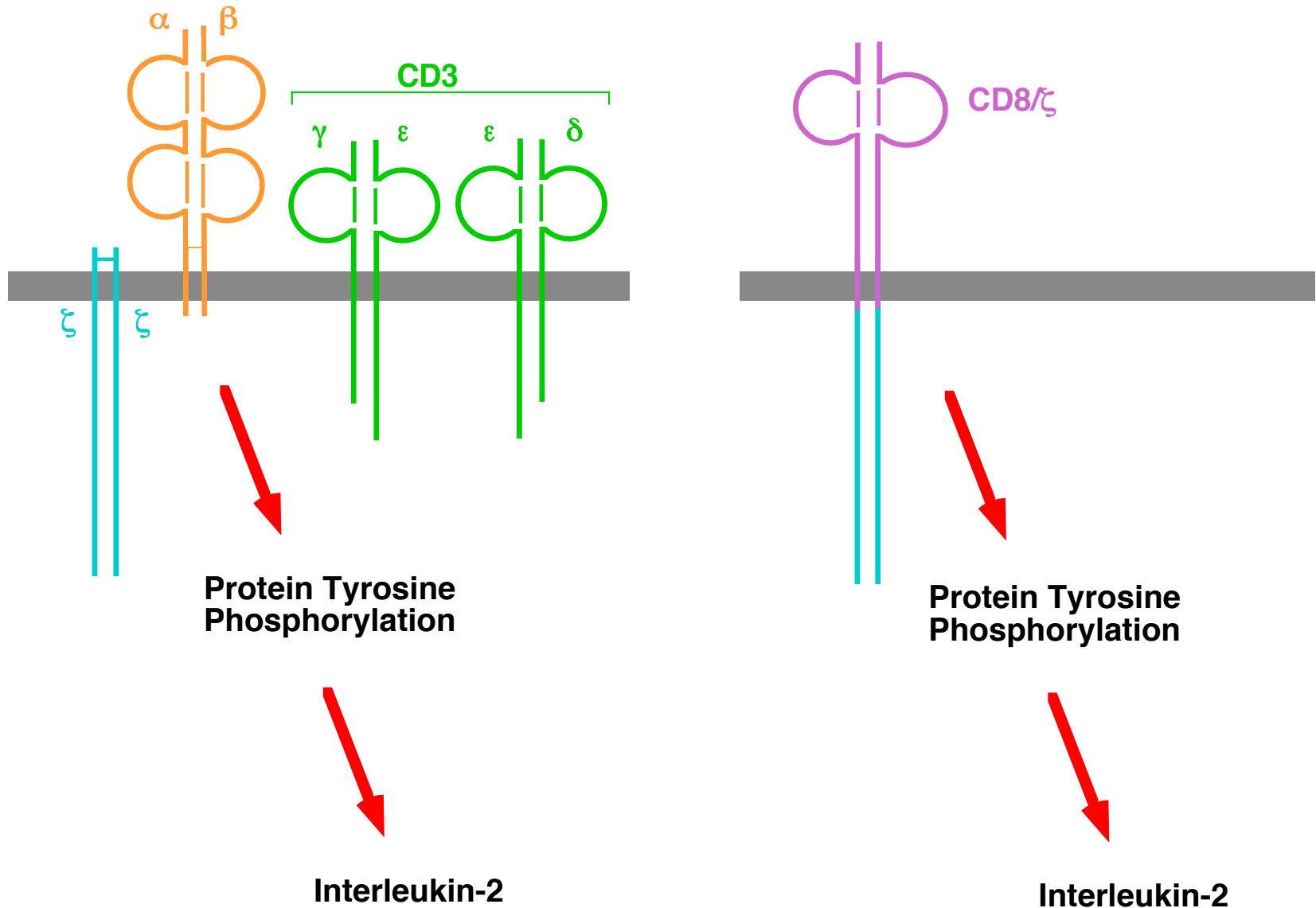
# Transmembrane domains allow structural and functional coupling of the $\alpha\beta$ heterodimer to CD3 chains

Tan and Weiss, J. Exp. Med, 1991



# TCR and CD8/ $\zeta$ Chimera Induce Similar Signals That Can Lead to T Cell Activation

Irving and Weiss, Cell, 1991

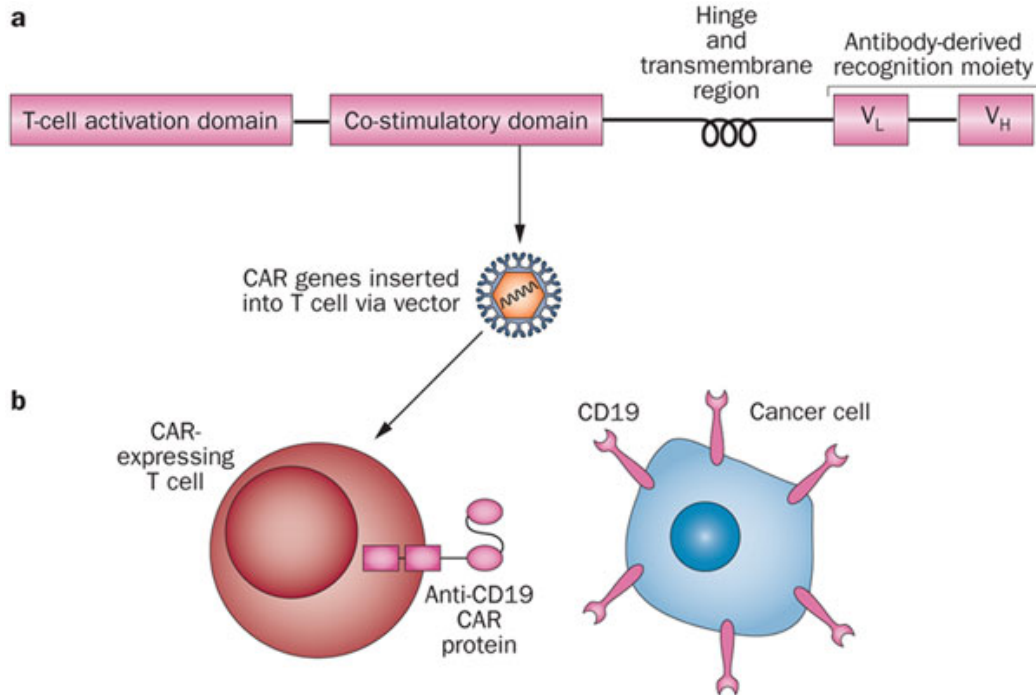


BRIEF REPORT

# Chimeric Antigen Receptor–Modified T Cells in Chronic Lymphoid Leukemia

David L. Porter, M.D., Bruce L. Levine, Ph.D., Michael Kalos, Ph.D., Adam Bagg, M.D., and Carl H. June, M.D.

*N Engl J Med* 2013;368:1509-18.  
DOI: 10.1056/NEJMoa1215134

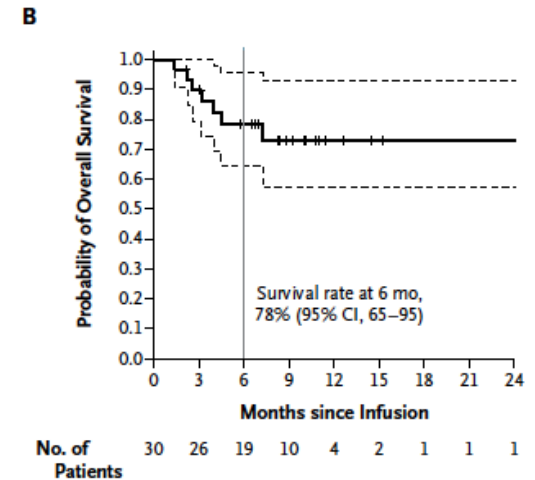


ORIGINAL ARTICLE

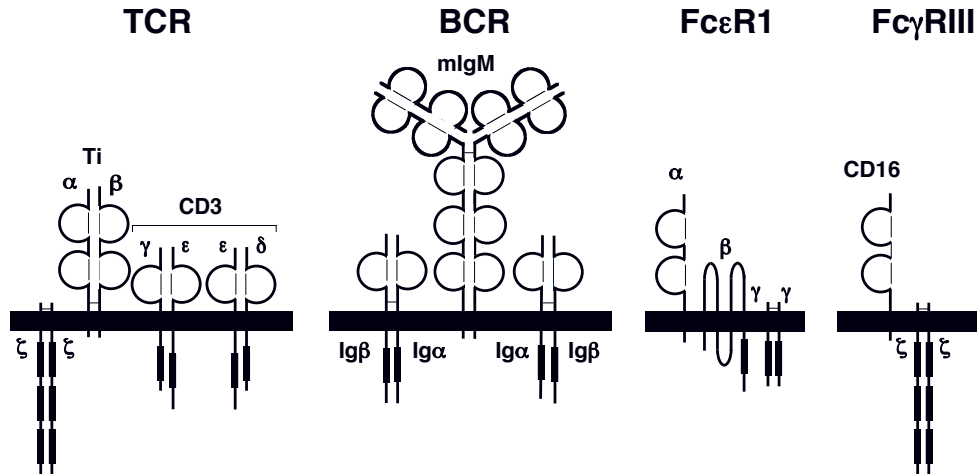
# Chimeric Antigen Receptor T Cells for Sustained Remissions in Leukemia

Shannon L. Maude, M.D., Ph.D., Noelle Frey, M.D., Pamela A. Shaw, Ph.D., Richard Aplenc, M.D., Ph.D., David M. Barrett, M.D., Ph.D., Nancy J. Bunin, M.D., Anne Chew, Ph.D., Vanessa E. Gonzalez, M.B.A., Zhaohui Zheng, M.S., Simon F. Lacey, Ph.D., Yolanda D. Mahnke, Ph.D., Jan J. Melenhorst, Ph.D., Susan R. Rheingold, M.D., Angela Shen, M.D., David T. Teachey, M.D., Bruce L. Levine, Ph.D., Carl H. June, M.D., David L. Porter, M.D., and Stephan A. Grupp, M.D., Ph.D.

*N Engl J Med* 2014;371:1507-17.  
DOI: 10.1056/NEJMoa1407222



# ITAM-containing receptors



## ITAMs (Immunoreceptor Tyrosine-Based Activation Motif)

hζ1	N Q L Y N E L N L G R R E E - Y D V L
hζ2	E G L Y N E L Q K D K M A E A Y S E I
hζ3	D G L Y Q G L S T A T K D T - Y D A L
hCD3γ	D Q L Y Q P L K D R E D D Q - Y S H L
hCD3ε	N P D Y E P I R K G Q R D L - Y S G L
hCD3δ	D Q V Y Q P L R D R D D A Q - Y S H L
hDAP12	E S P Y Q E L Q G Q R S D V - Y S D L
rIgE FcR γ	D A V Y T G L N T R N Q E T - Y E T L
rIgE FcR β	D R L Y E E L - H V Y S P I - Y S A L
mIg α	E N L Y E G L N L D D C S M - Y E D I
mIg β	D H T Y E G L N I D Q T A T - Y E D I
BLV gp30	D S D Y Q A L L P S A P E I - Y S H L
EBV LMP-2	H S D Y Q P L G T Q D Q S L - Y L G L
SIV Nef	G D L Y E R L L R A R G E T - Y G R L
KSHV	L Q D Y Y S L H D L C T E D - Y T Q P
Consensus	- - - Y - - L - - - - - - - Y - - L



# The ITAM as a Conserved Signal Transduction Module

ITAM can confer signal transduction function to heterologous receptors,  
17 aa are enough

ITAMs are encoded on 2 exons, evidence for evolutionary conservation

Tyrosines and Leucines (or Isoleucines) are critical, as is spacing between YXXL  
residues 7 and 8 aa spacer are OK, 6 is not

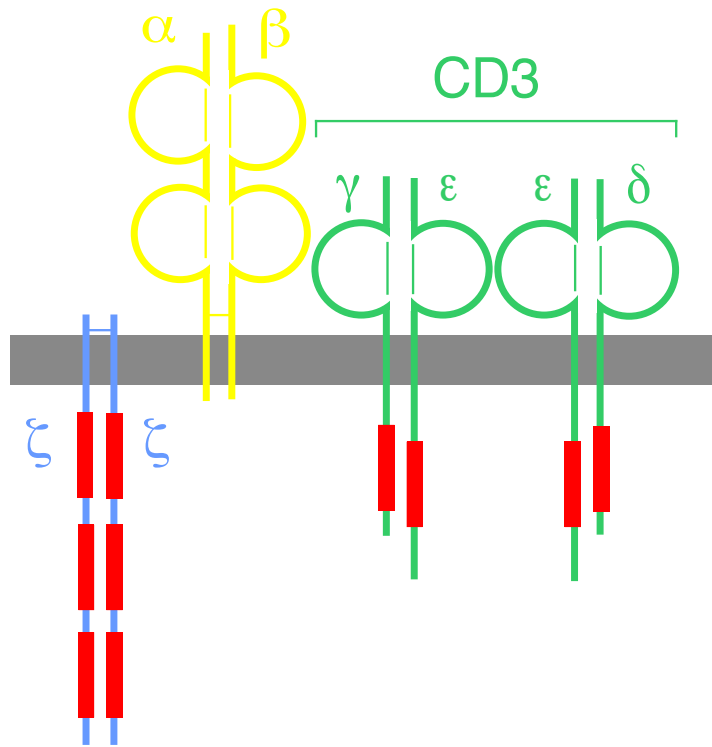
Function of redundancy: Signal Amplification vs Distinct Functions  
Multimers signal better  
Effector binding differences

Viruses usurp signaling function

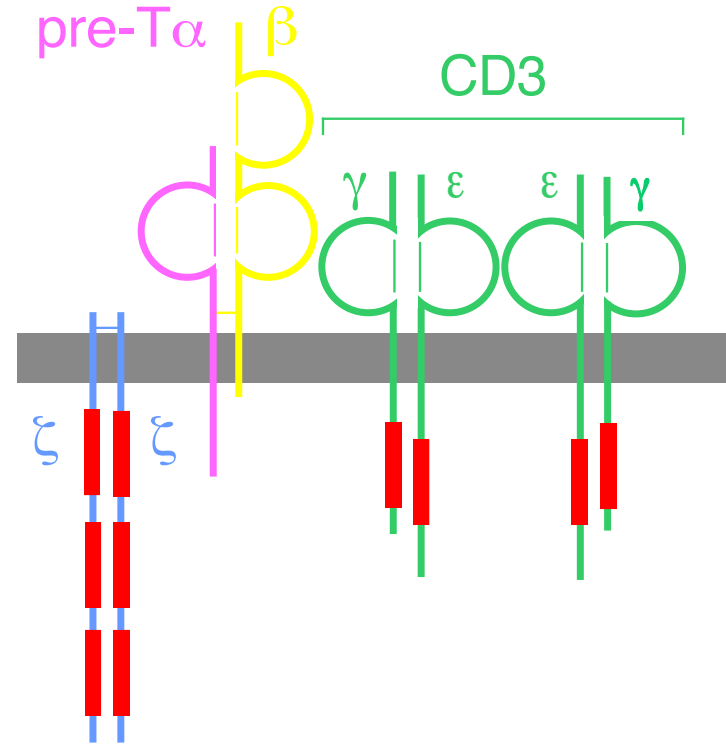
**The Pre-TCR, an immature receptor to  
assess functional  $\beta$ -chain rearrangement  
during thymus development**

# An immature form of the TCR has a surrogate for the $\alpha$ Chain, Pre-T $\alpha$

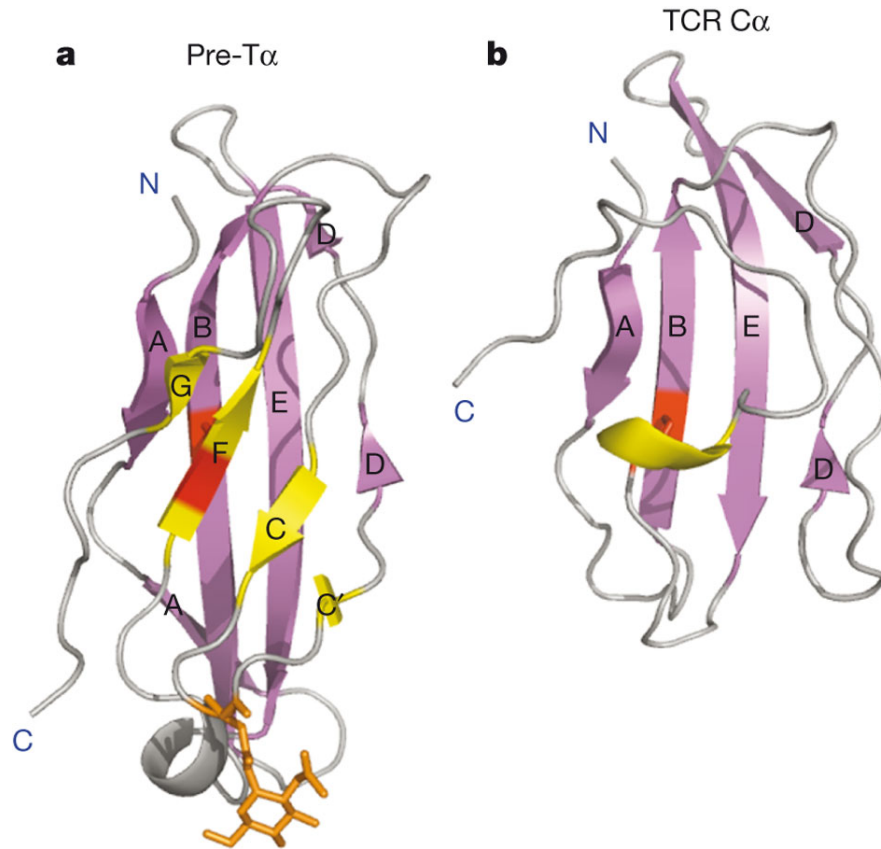
## TCR



## Pre-TCR

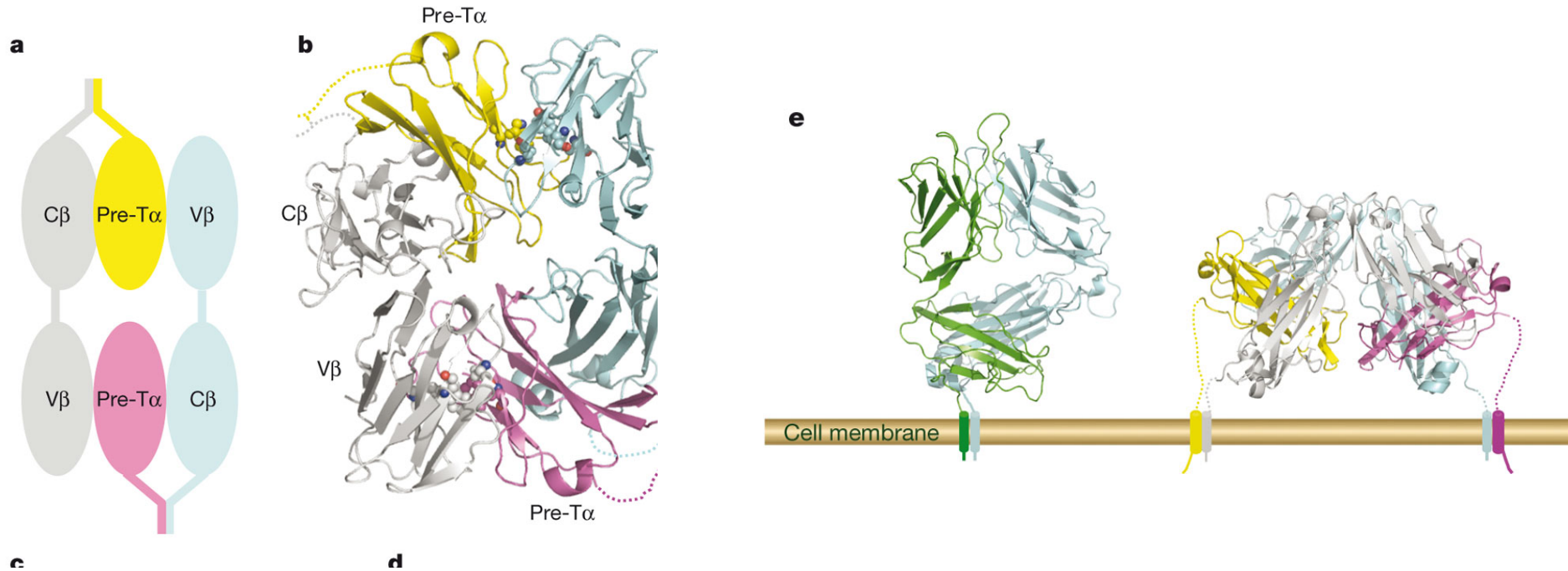


# The pre-T $\alpha$ structure



SS Pang *et al.* *Nature* **467**, 844-848 (2010) doi:10.1038/nature09448

# The pre-TCR dimer forms a constitutive dimer



SS Pang *et al. Nature* **467**, 844-848 (2010) doi:10.1038/nature09448

nature