

Micro 204

TCR Signal Transduction

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October 15, 2018

Signal Transduction: What is it?

Definition: *Signal transduction is the transmission of information from outside of the cell or within the cell to convey information that could induce or affect a change in the cellular status.*

How is it accomplished? (a few examples)

- Changes in protein or lipid phosphorylation

- Other protein modifications (i.e., ubiquitylation, sulfenylation glycosylation, etc.)

- Changes in protein localization

- Changes in protein:protein interactions

- Changes in protein:lipid interactions

- Assembly of macromolecular complexes

- Disassembly of protein complexes

- Protein degradation

- Changes in lipid turnover

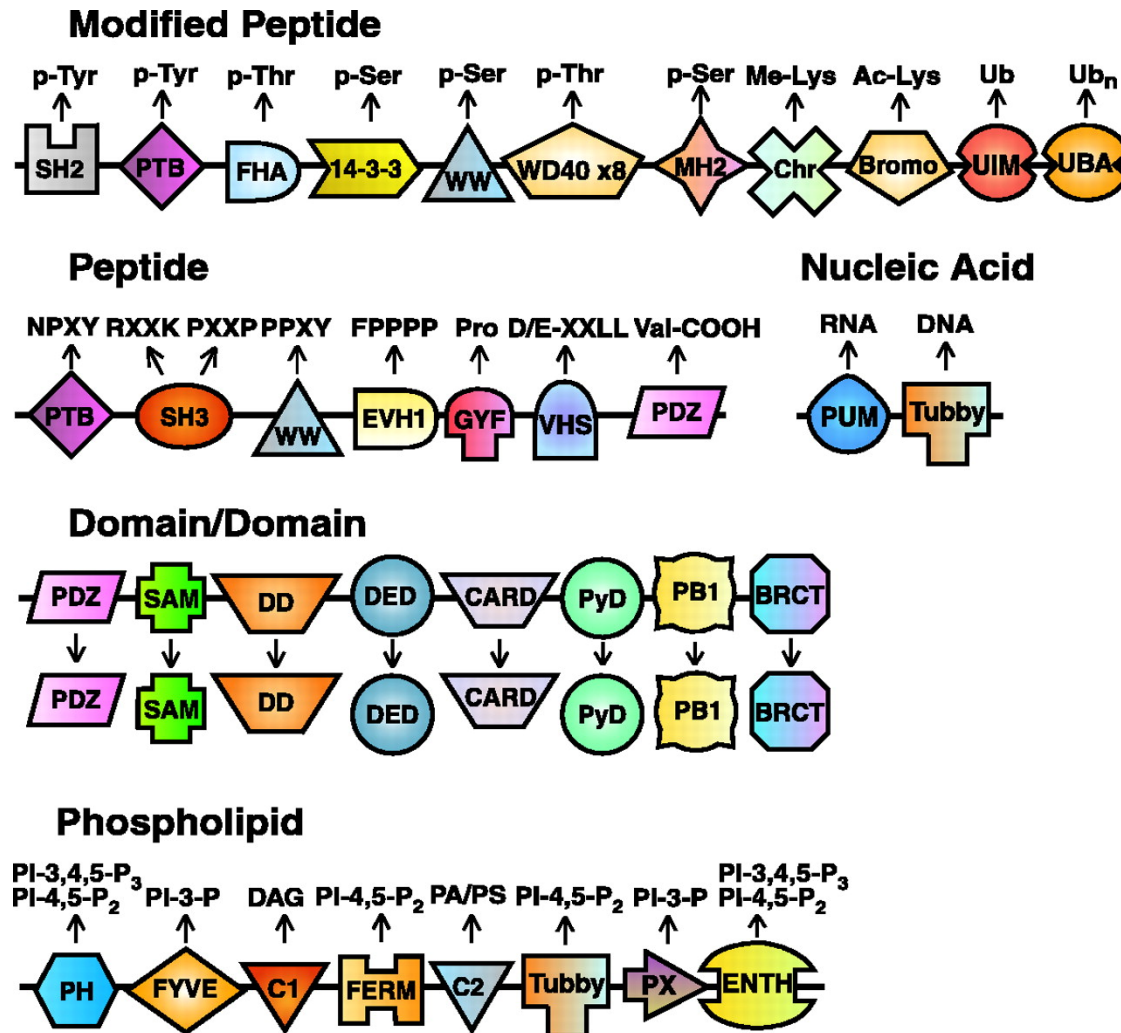
- Generation of second messengers

- Ionic fluxes

- Signal dampening or amplification

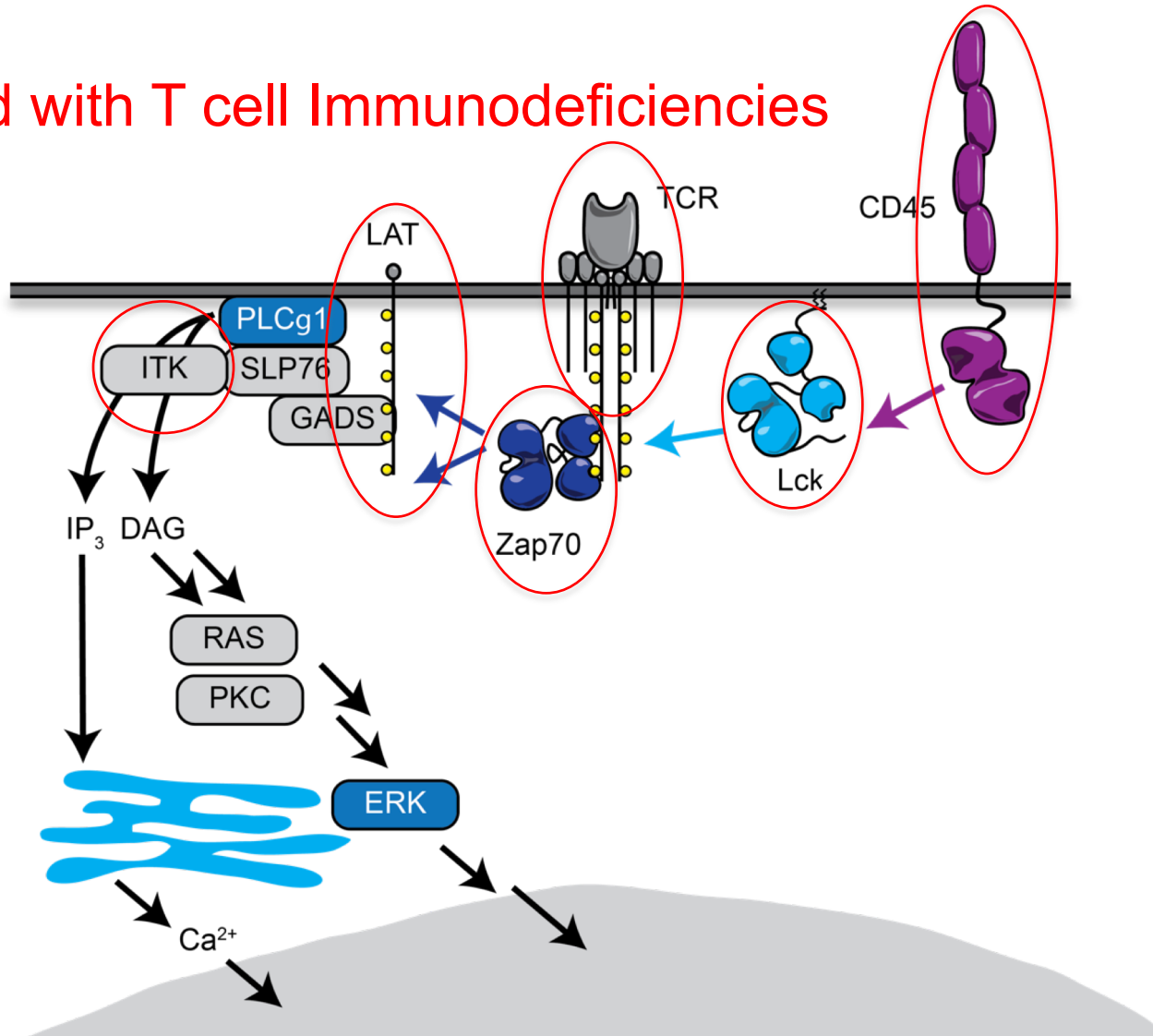
- Feedback loops, positive or negative

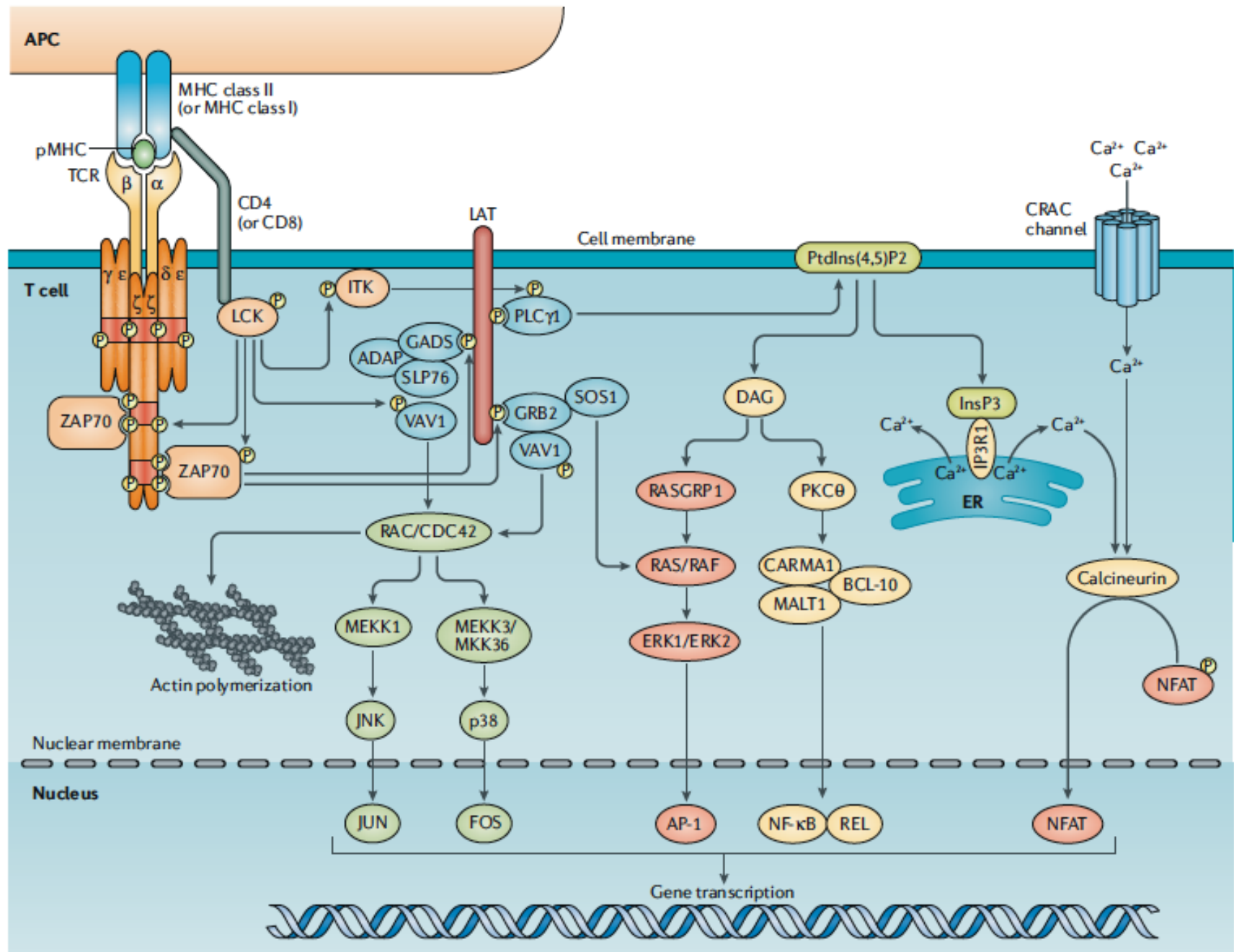
Modular Protein Interaction Domains Are Involved in Signaling Pathways



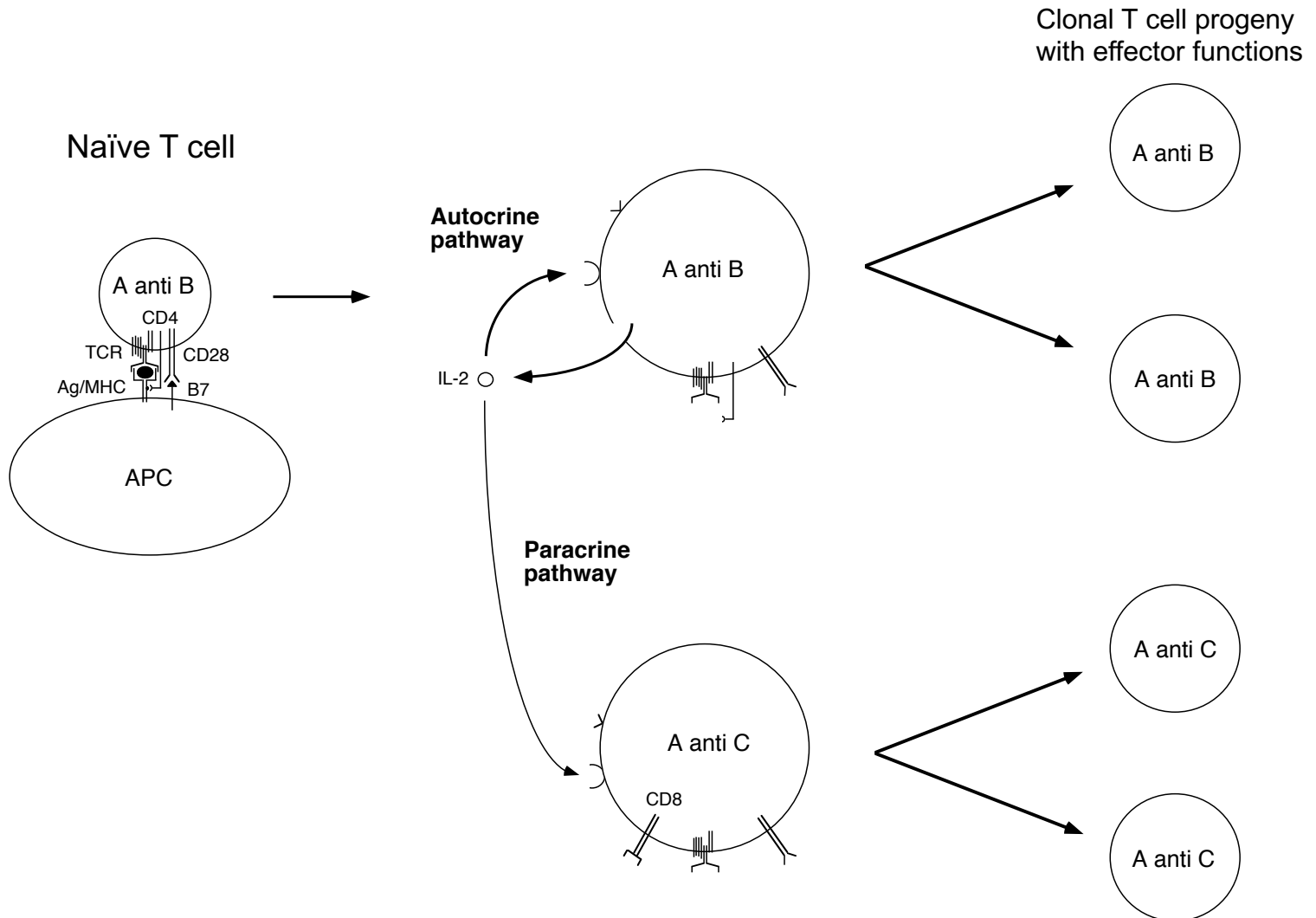
Early TCR signaling events

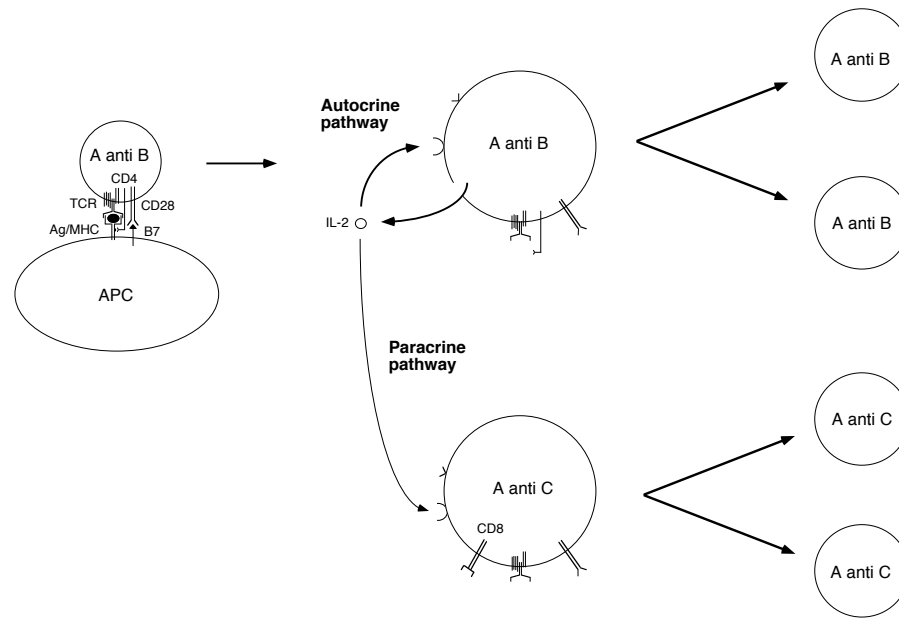
Associated with T cell Immunodeficiencies





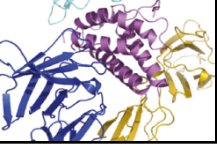
TCR Signaling is Required for the Activation, Clonal Proliferation and Differentiation of Naïve T Cells



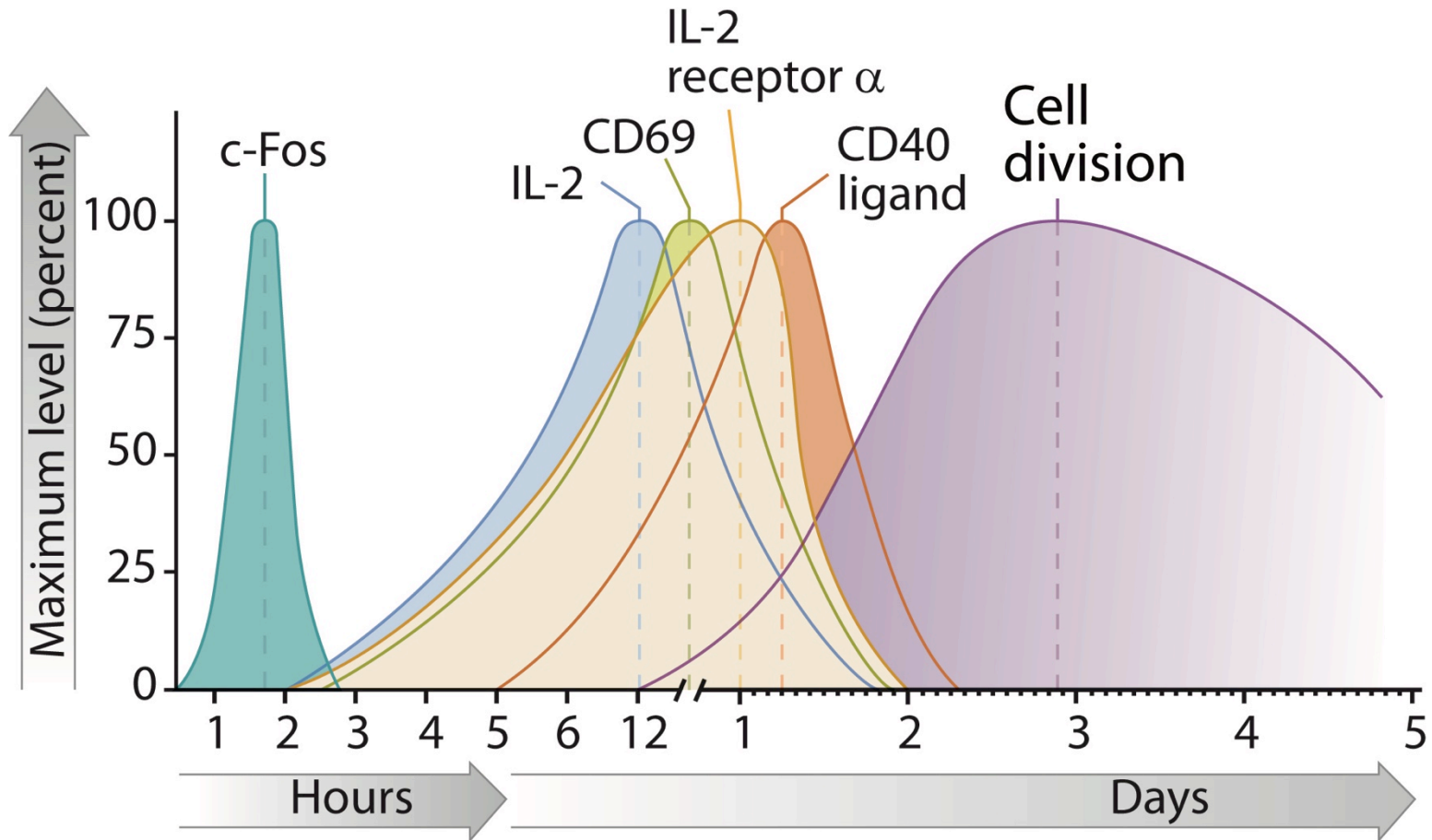


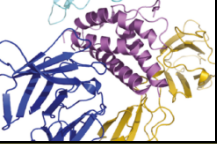
Key Points:

- 1. T cell activation occurs during a complex cell-cell interaction**
- 2. Stimulation of a quiescent G_0 T cell results in cell cycle progression as well as enormous metabolic and synthetic changes and cell enlargement**
- 3. Activation is initiated by the TCR, but also requires multiple other receptors involved in intercellular interactions (i.e., costimulation via CD28)**
- 4. Key event is the production of lymphokines and the cellular response to them**
- 5. End result is the clonal expansion and acquisition of effector functions**

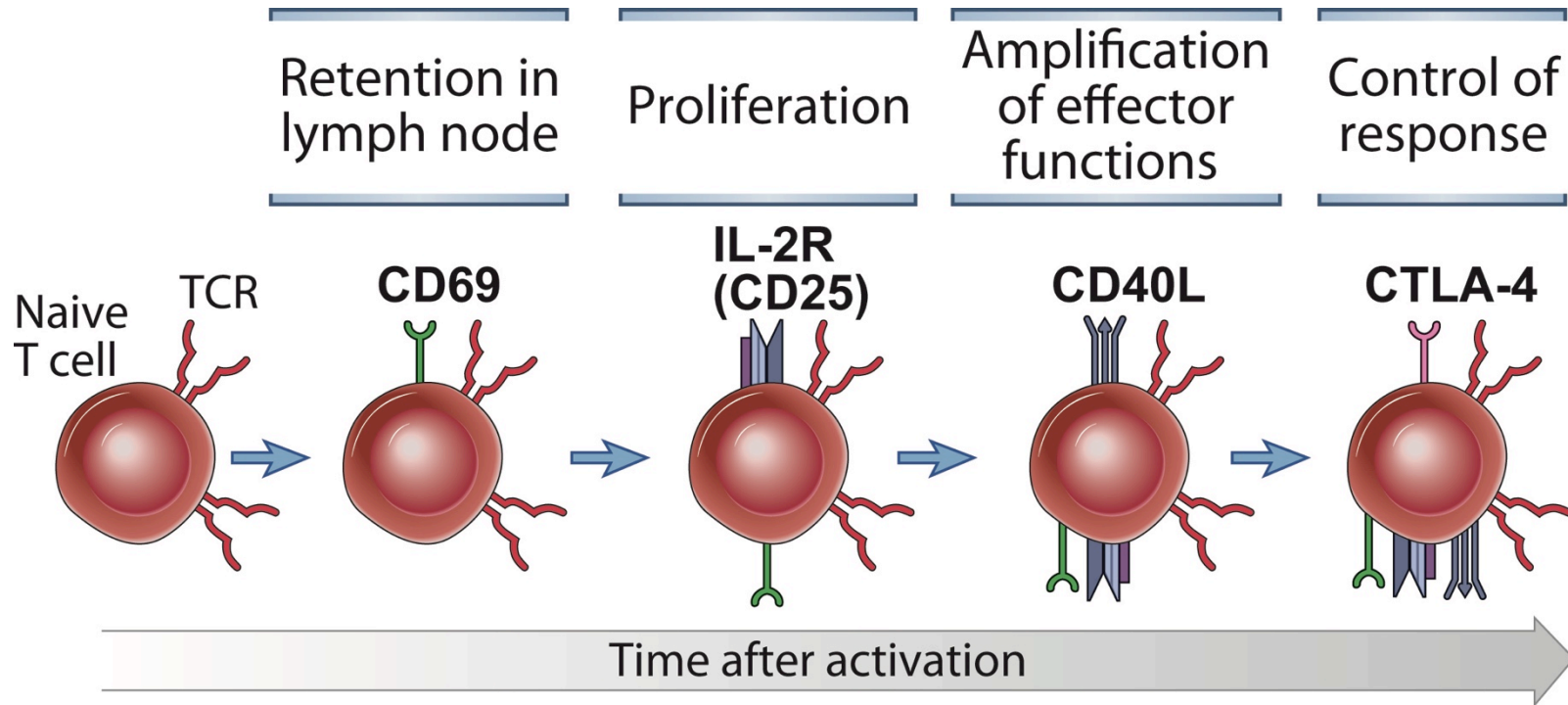


Genes Expressed After T Cell Activation: Time Scale





Molecules Expressed After T Cell Activation



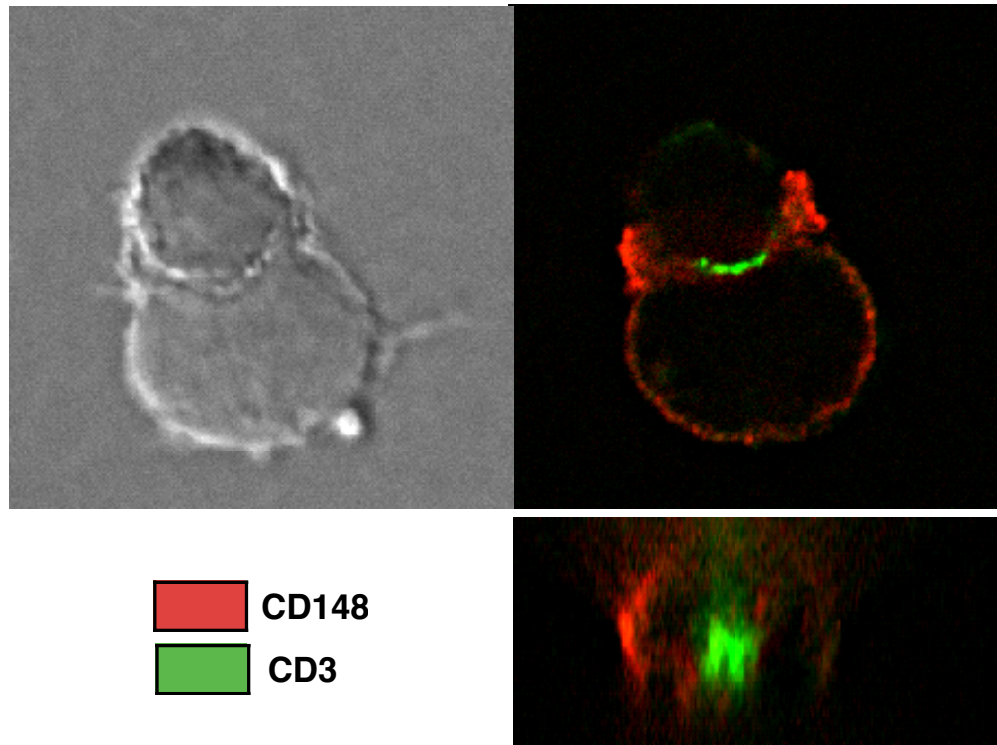
Consequences of Antigen Receptor Signaling

- Biochemical changes
 - Protein phosphorylation and other modifications
 - Phospholipid turnover
 - Ionic changes
 - Protein relocalization
 - Proteolysis
- Cytoskeletal Reorganization/Cell Polarization
 - Directed cytotoxicity
 - Directed cytokine secretion
- Gene Transcription
 - Cytokines and their receptors
 - Cell surface molecules
 - Antibody production
- Effector Functions
 - Cytokine secretion
 - Cytotoxicity
 - Antibody class switching
- Cell proliferation
- Apoptosis

Immunologic Synapse:

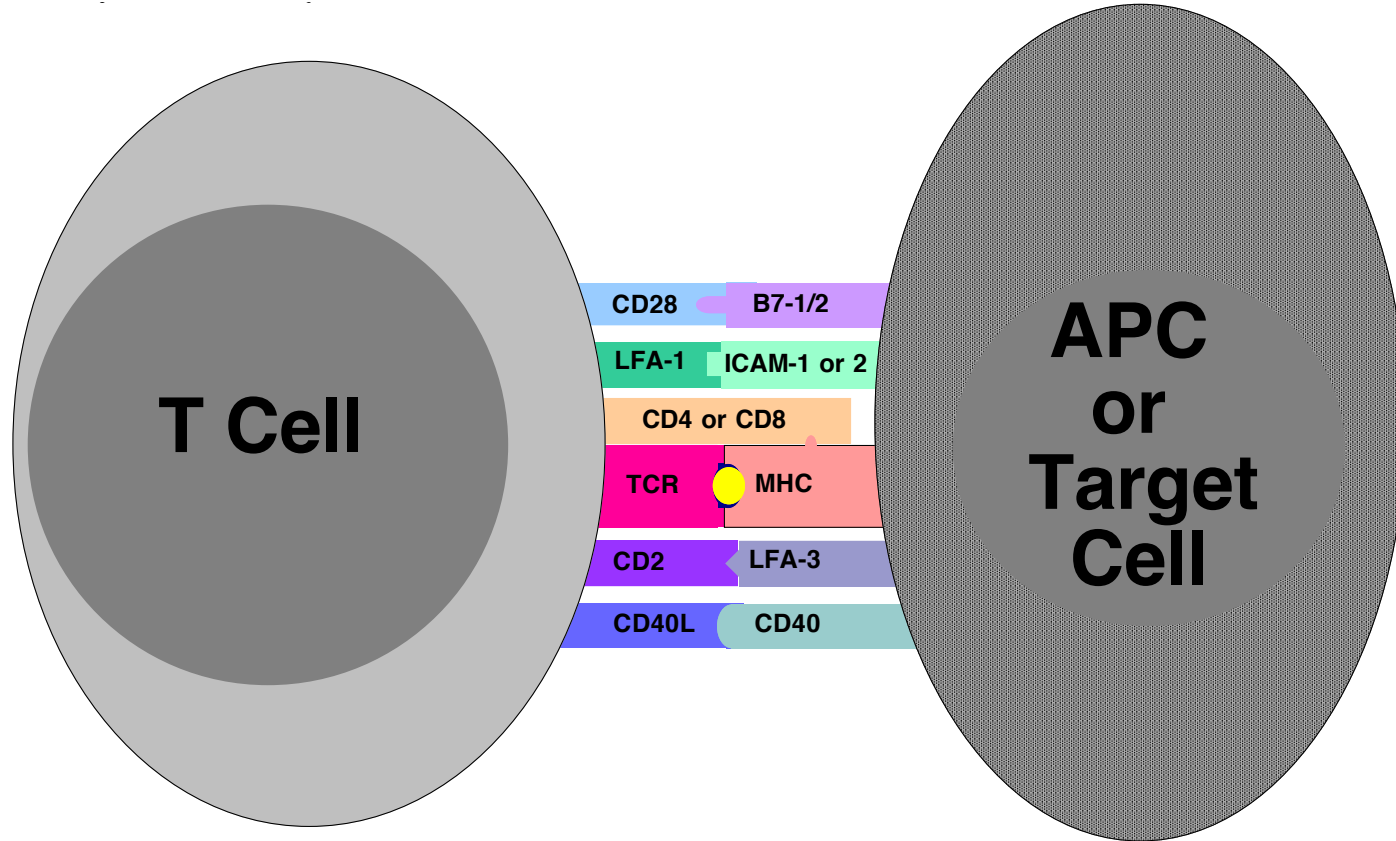
A spatial and temporal segregation of cell surface molecules

Jurkat:APC conjugates with SAg



Lin and Weiss, J. Cell Biol., 2003

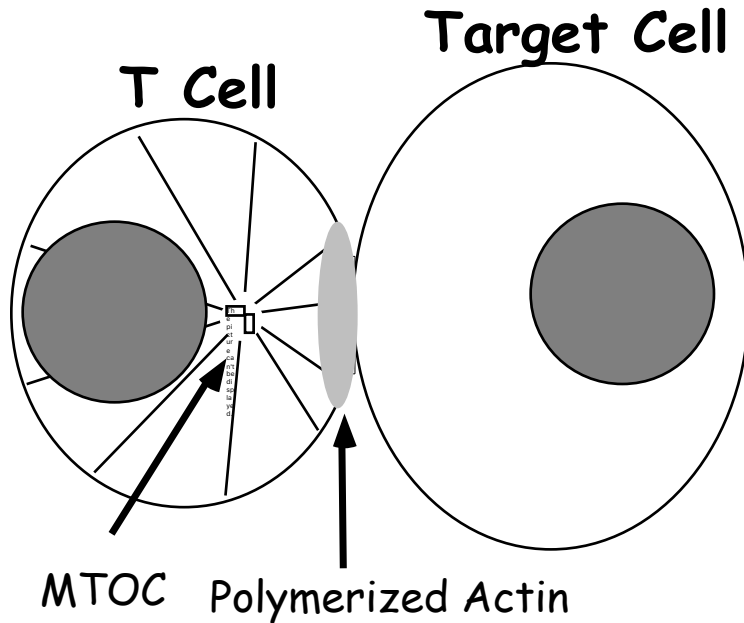
Complexity of the T Cell : Antigen Presenting Cell Interaction



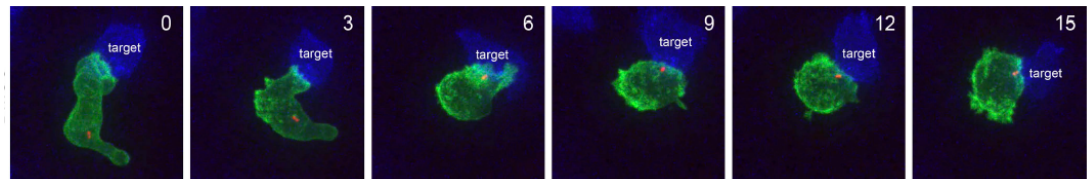
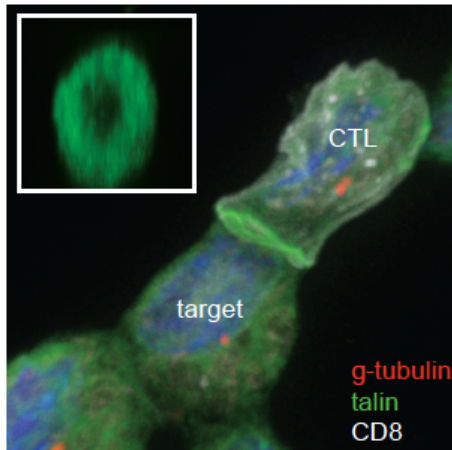
Intermolecular Interactions

| | |
|----------------------|---------------|
| Antigen recognition: | TCR – pMHC |
| Coreceptor function: | CD4/CD8 - MHC |
| Adhesion: | LFA-1 – ICAM1 |
| | CD2 – LFA-3 |
| | CD40L – CD40 |
| Costimulation: | CD28 – B7-1/2 |
| Differentiation: | Cytokines |

T Cells Polarize During Antigen-Specific Recognition

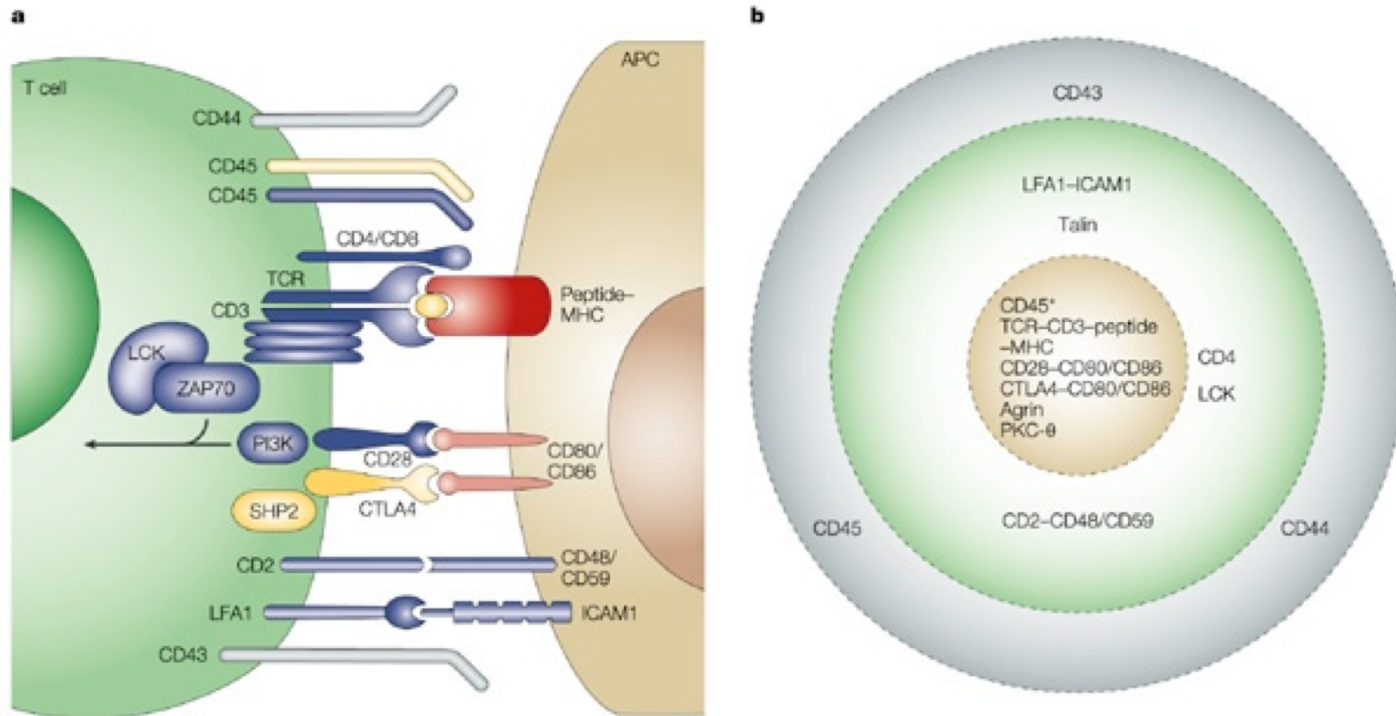


1. T cell cytoskeleton polarizes; APC does not
2. MTOC and microtubules polarize
3. Actin cytoskeleton polymerizes and polarizes
4. Requires TCR signals: Rho and Rac GTPases
5. Polarization of cytoskeletal elements contribute to polarized secretion of cytolytic granules or lymphokines



Immunologic Synapse:

A spatial and temporal segregation of cell surface molecules

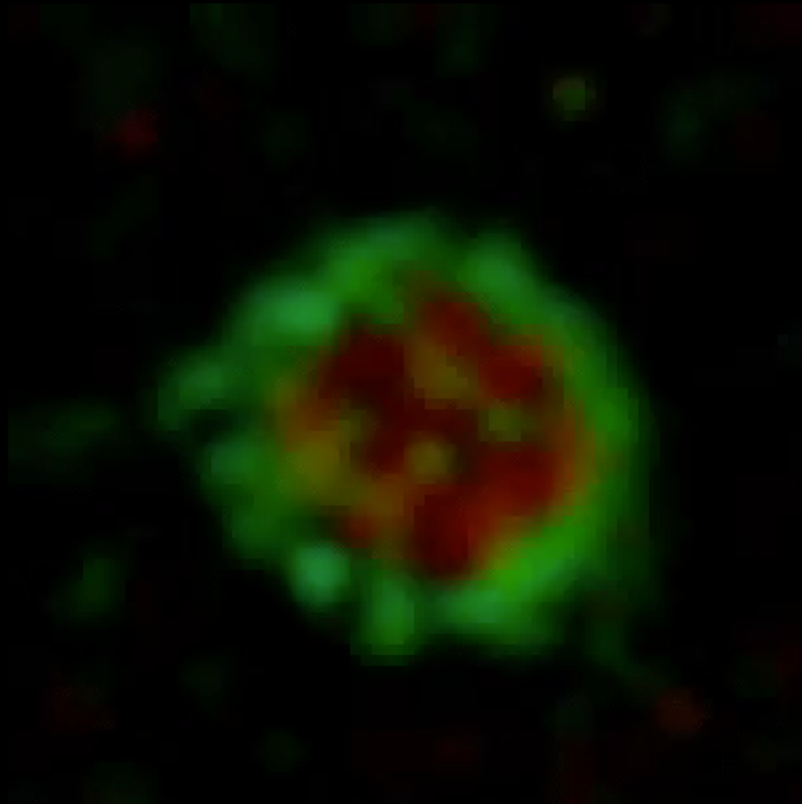


Nature Reviews | Immunology

Avi Kupfer - deconvolution IF microscopy of T cell/APC interactions - organized, bulls-eye type of structure

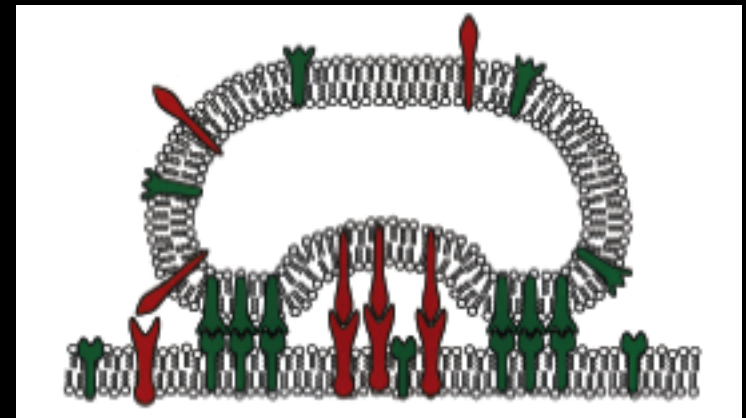
Non-activating, altered, peptides do not support the formation of these structures

Formation of an Immunological Synapse Involves Coordinated Spatial Organization



Red= ICAM-1

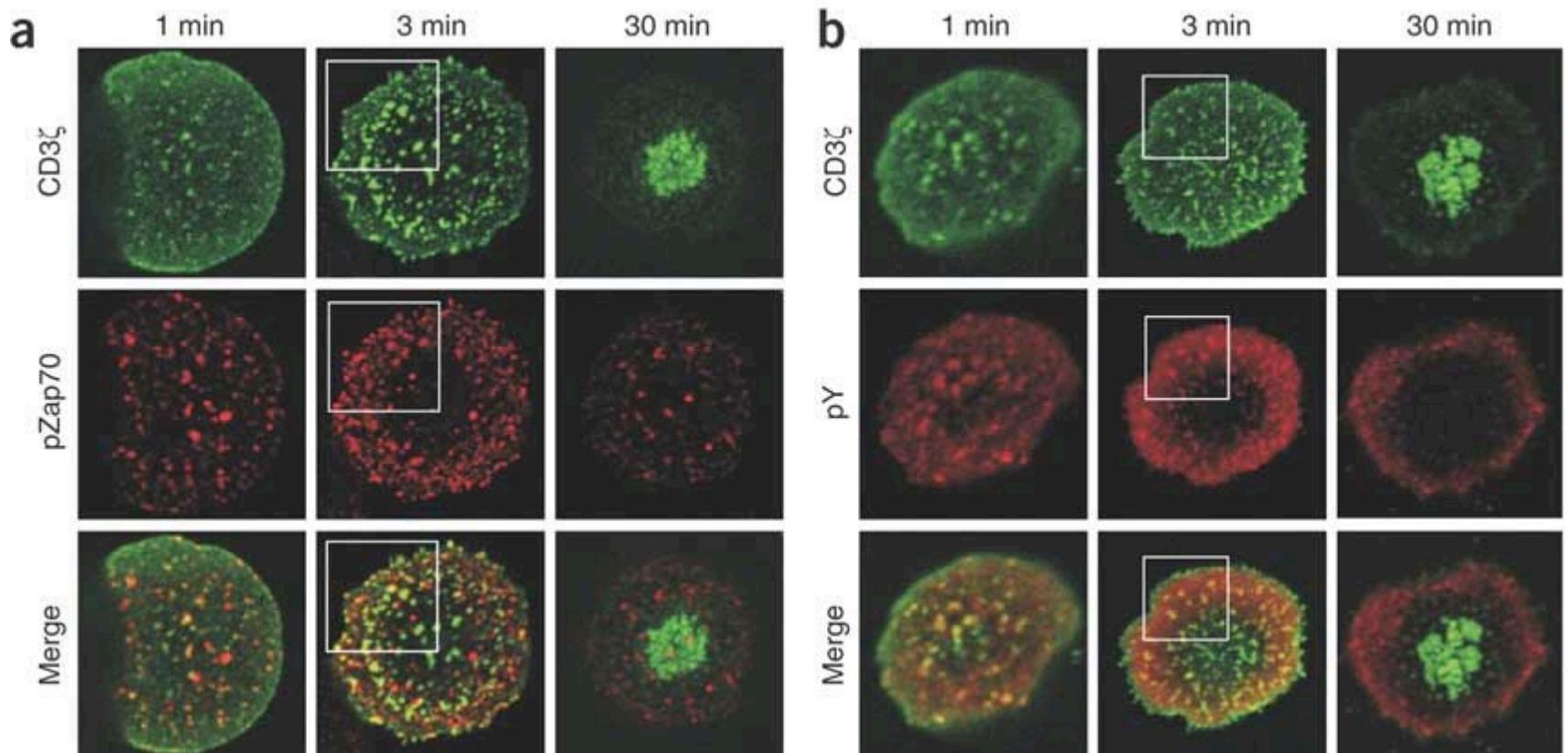
Green=MHC/pep



Grakoui et al. (1999) *Science* 285:221.

Signaling Occurs in TCR Microclusters at the Periphery of the Synapse

Yokosuka et al., Nature Immunol. 6, 1253, 2005



Function(s) of the Immunological Synapse?

1. Signal Initiation: *probably not*

P-tyr and Calcium responses precede mature synapse formation

2. Signal enhancement: *perhaps*

But no biochemical correlate and mature synapse does not appear to be required (CD2-AP KO signals better than wt but does not form synapse)

Peripheral microclusters appear to be most active signaling complexes

3. Directed secretion: *fits well*

Directed secretion by CTL and Th cells makes teological sense and has been well documented

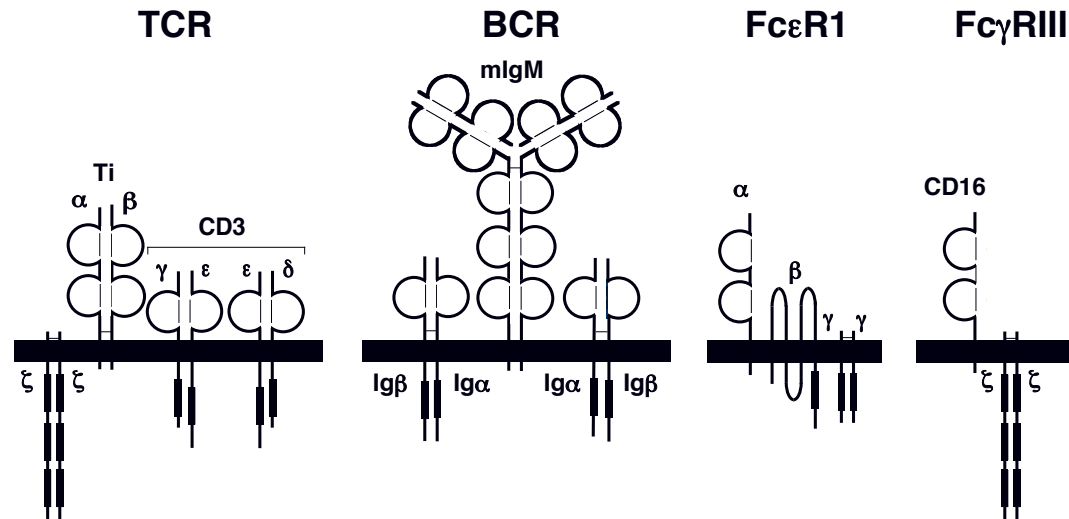
4. Receptor downmodulation: *likely*

Modeling and CD2-AP KO data supports this

New data on TCR microclusters also supportive

Some of the central TCR clusters are in internal structures and may even be in extracellular vesicles

How does ligand binding induce TCR signaling?



ITAMs (Immunoreceptor Tyrosine-Based Activation Motif)

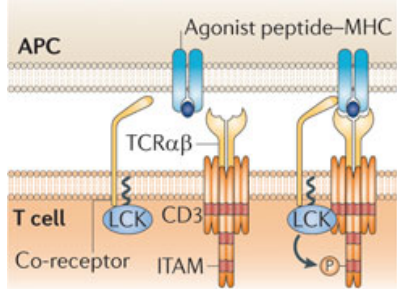
| | |
|------------|---------------------------------------|
| hζ1 | N Q L Y N E L N L G R R E E - Y D V L |
| hζ2 | E G L Y N E L Q K D K M A E A Y S E I |
| hζ3 | D G L Y Q G L S T A T K D T - Y D A L |
| hCD3γ | D Q L Y Q P L K D R E D D Q - Y S H L |
| hCD3ε | N P D Y E P I R K G Q R D L - Y S G 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 |

1. *How do TCRs transmit their ligand occupancy state across the plasma membrane?*
2. *Do TCRs undergo conformational changes when they bind ligand?*

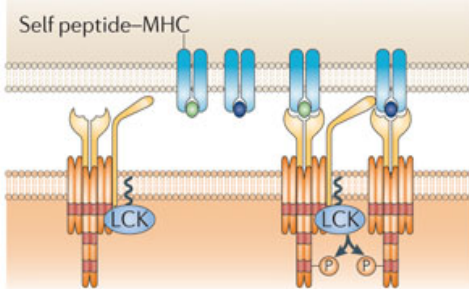
Models of TCR triggering

Aggregation

a Co-receptor heterodimerization

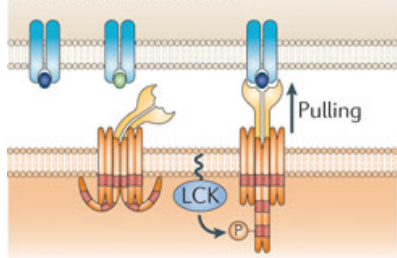


b Pseudodimer

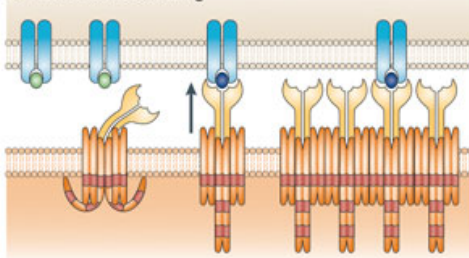


Conformational change

c Piston-like movement

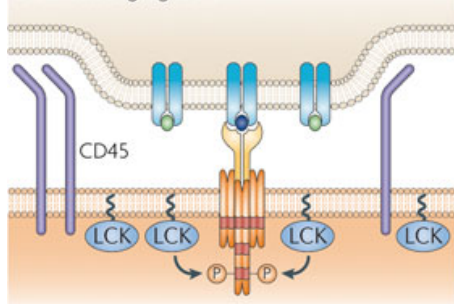


d Induced clustering

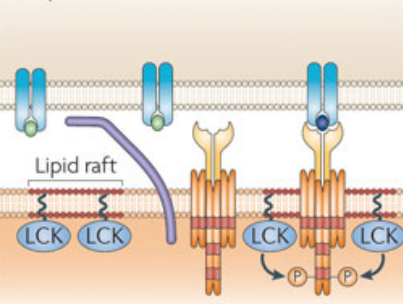


Segregation or redistribution

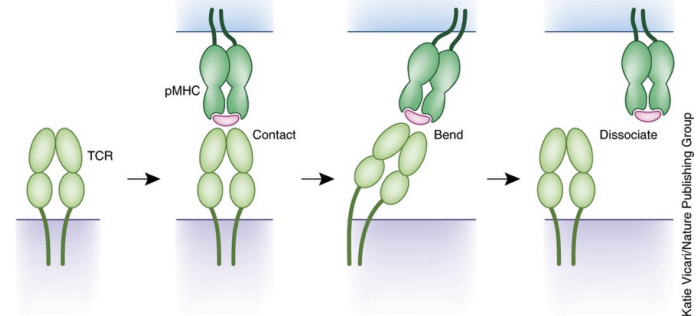
e Kinetic segregation



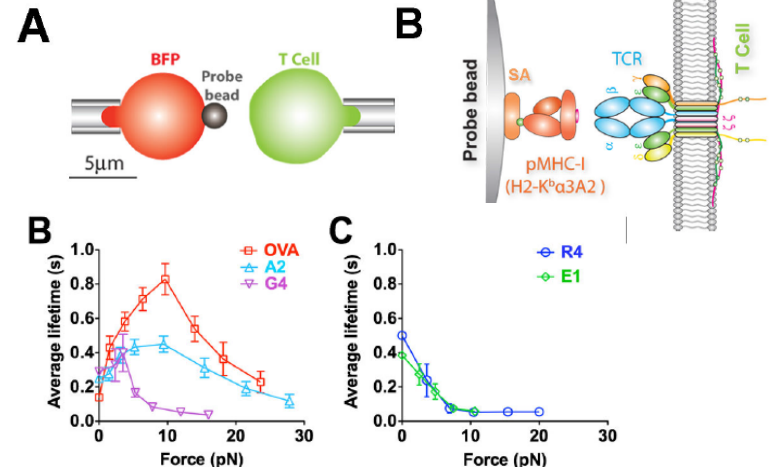
f Lipid raft



Mechano-sensory model



Catch-Bond Model

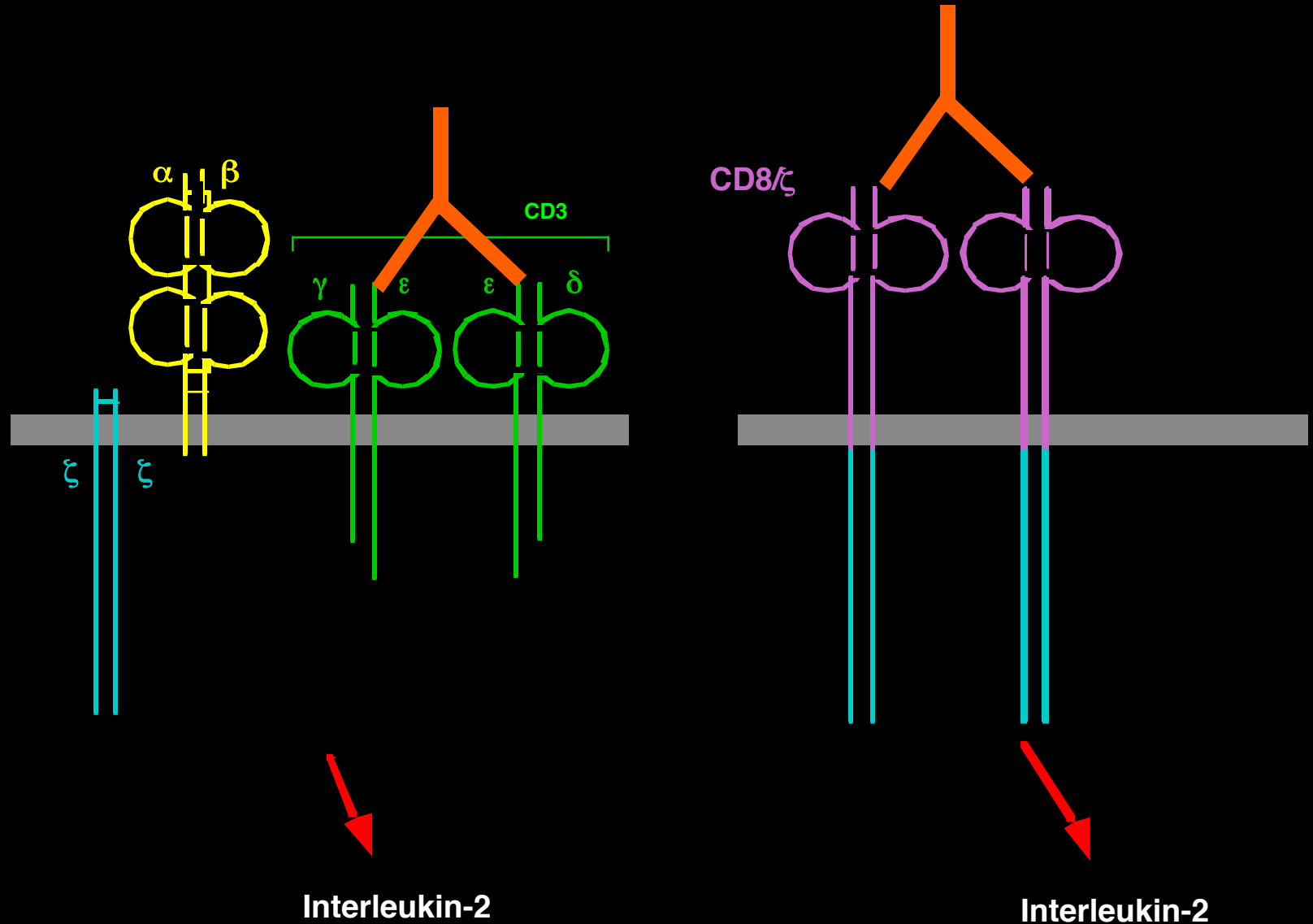


Van der Merwe & Dushek, Nature Rev. Immunol
Chakraborty and Weiss, Nature Immunol., 2014;
Liu, et al., Cell, 2015

Support for the Oligomerization Model

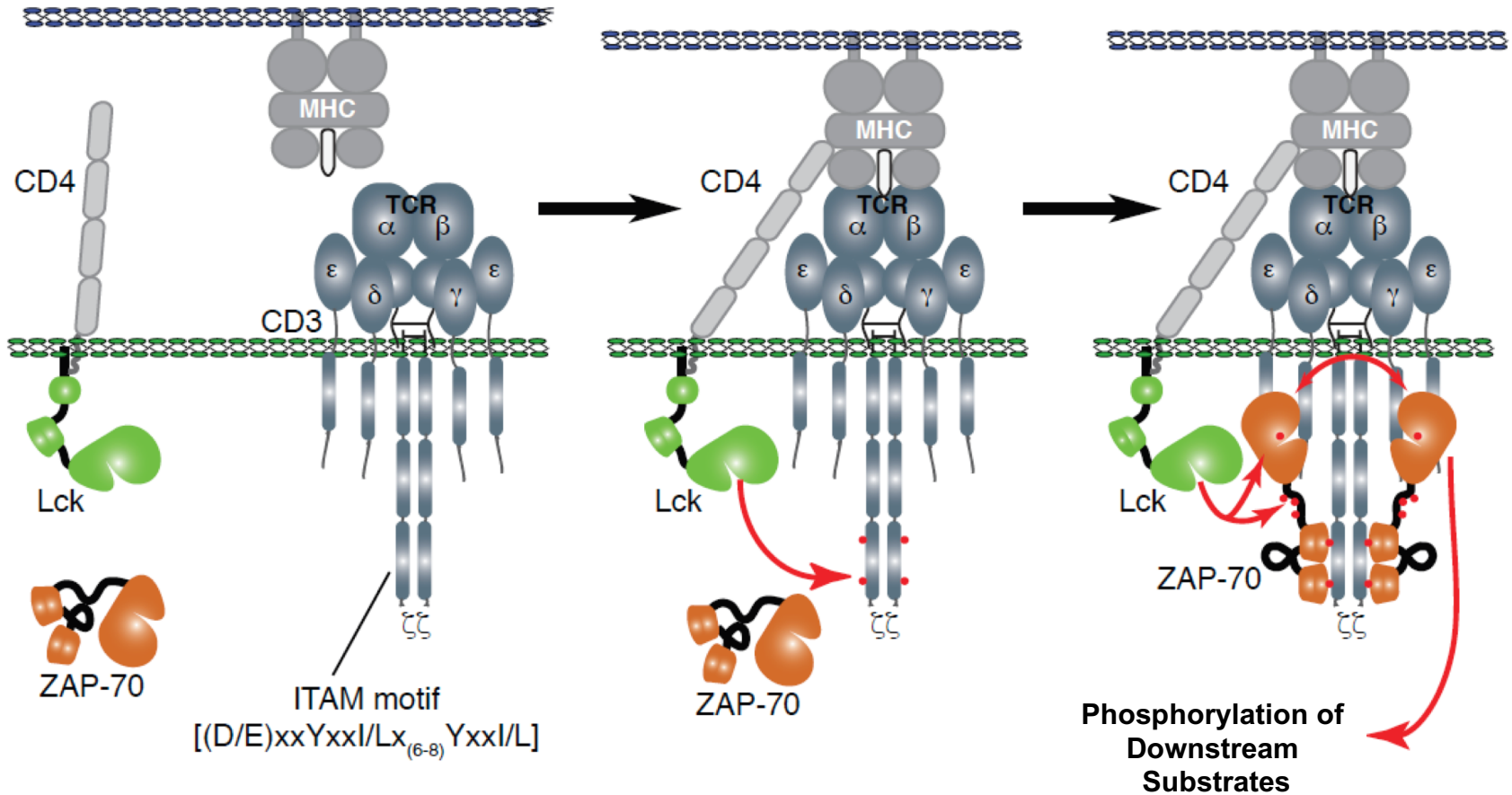
- Bivalent (but usually not univalent Fab) mAbs can induce T cell responses
- Many different anti-TCR mAbs can induce TCR signaling
- The more multivalent the mAb or pMHC (i.e., tetramer), the better
- Chimeric receptors (ITAM containing chimeras) signal
- Hetero-oligomerization with coreceptors (ie. CD4) increases sensitivity to peptide/MHC molecule

mAbs Against the TCR/CD3 Complex or Chimeric Receptors can Induce Similar Signals and Responses



A Hetero-Oligomerization Model:

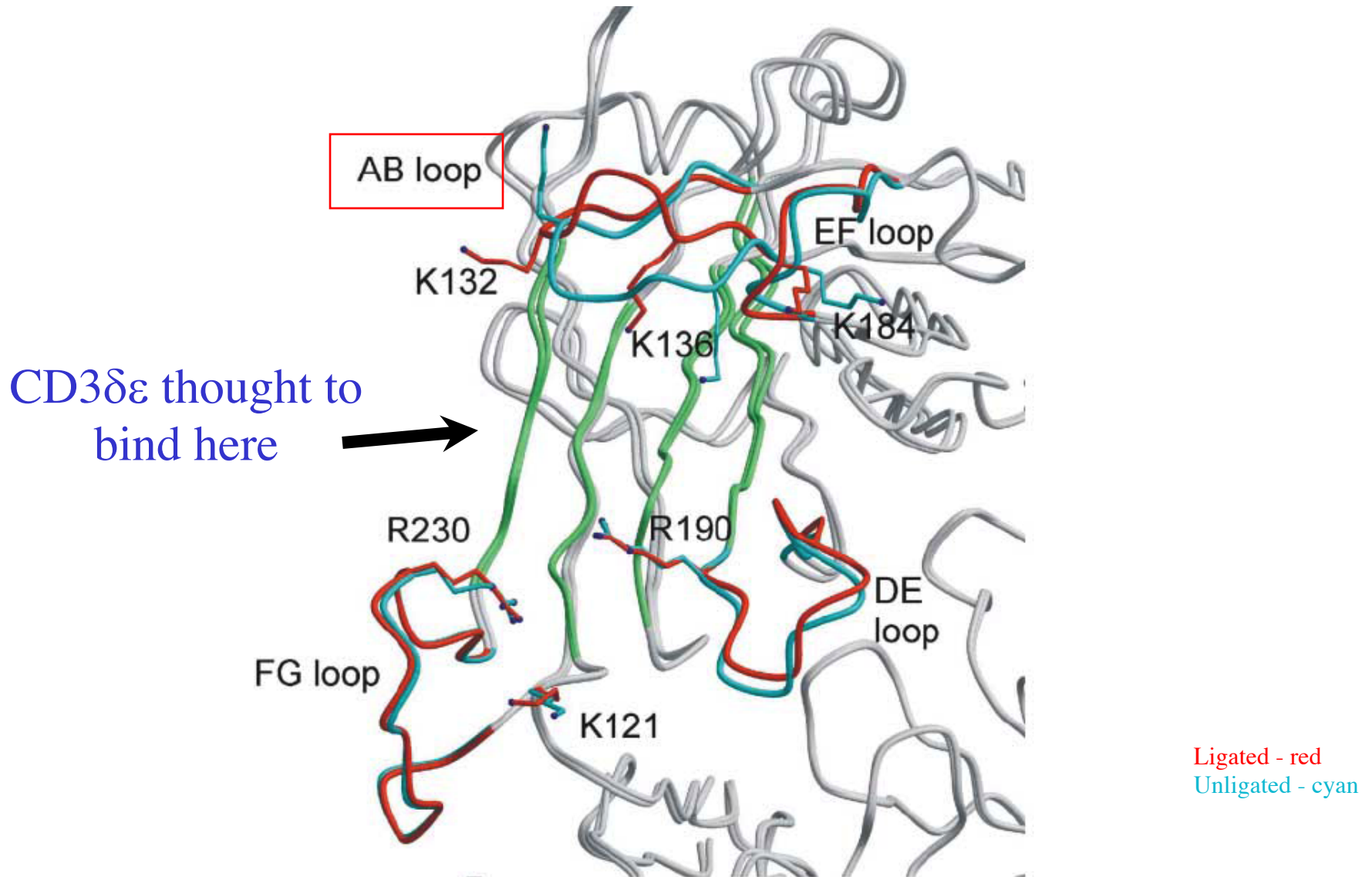
CD4 and TCR are Brought Together During Antigen Recognition



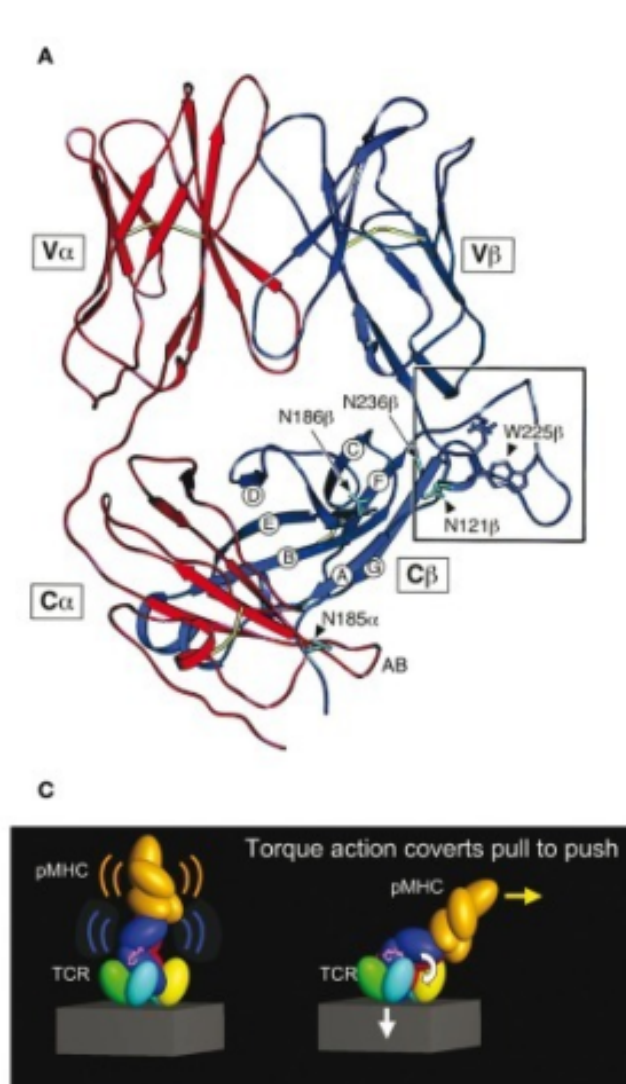
Some data support a conformational change

Conformational Change in the TCR α Constant Region A-B Loop in the LC13 TCR

Kjer-Nielsen, et al., Immunity, 2003

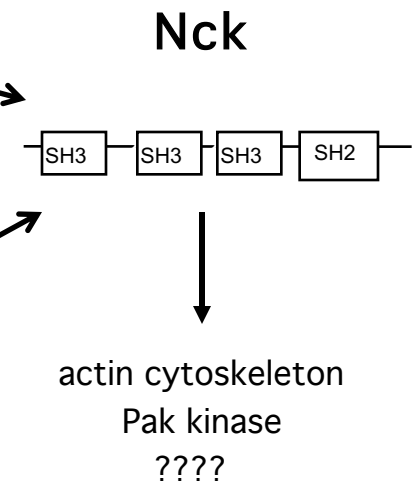
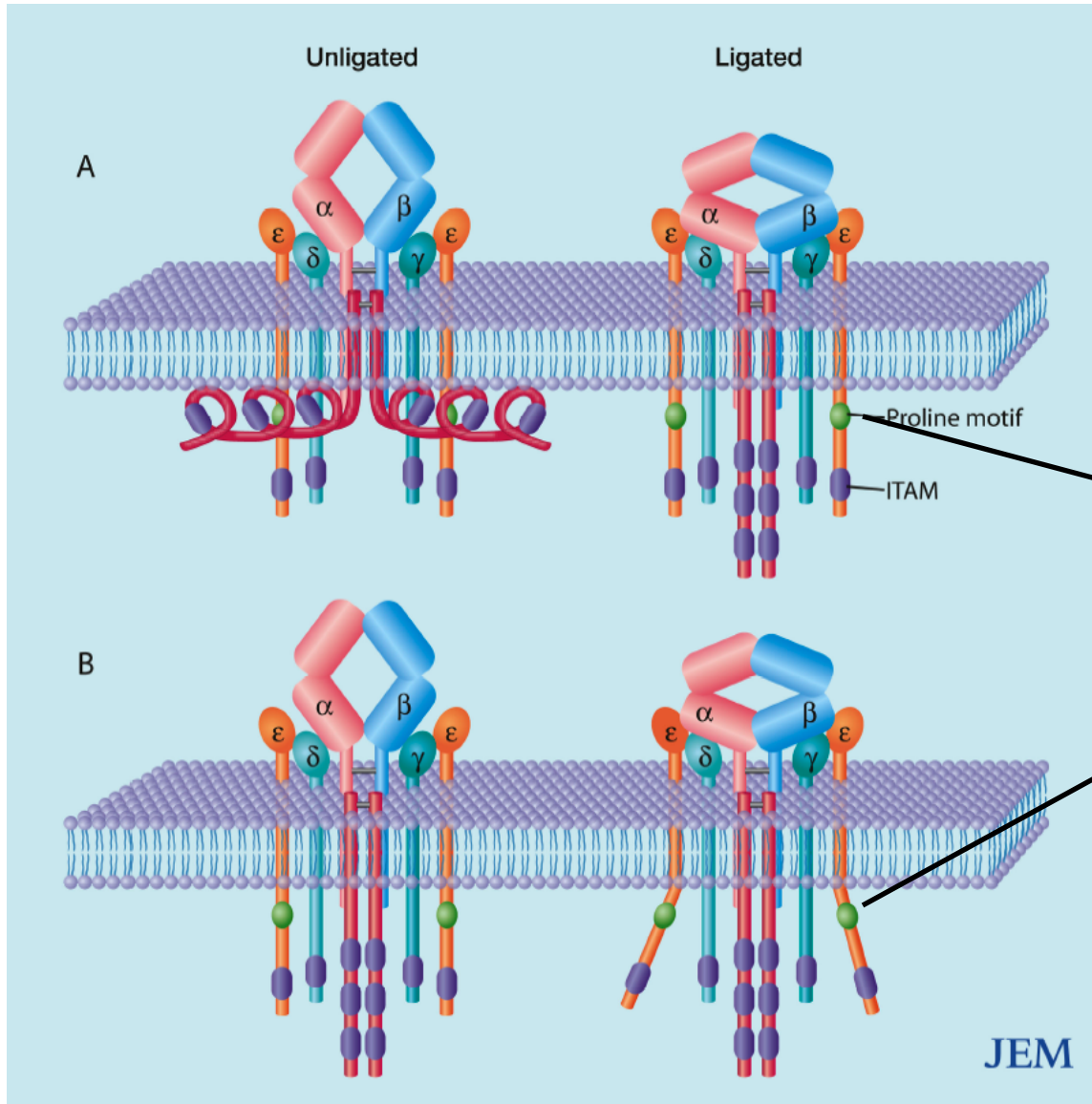


The TCR as a Mechanosensor



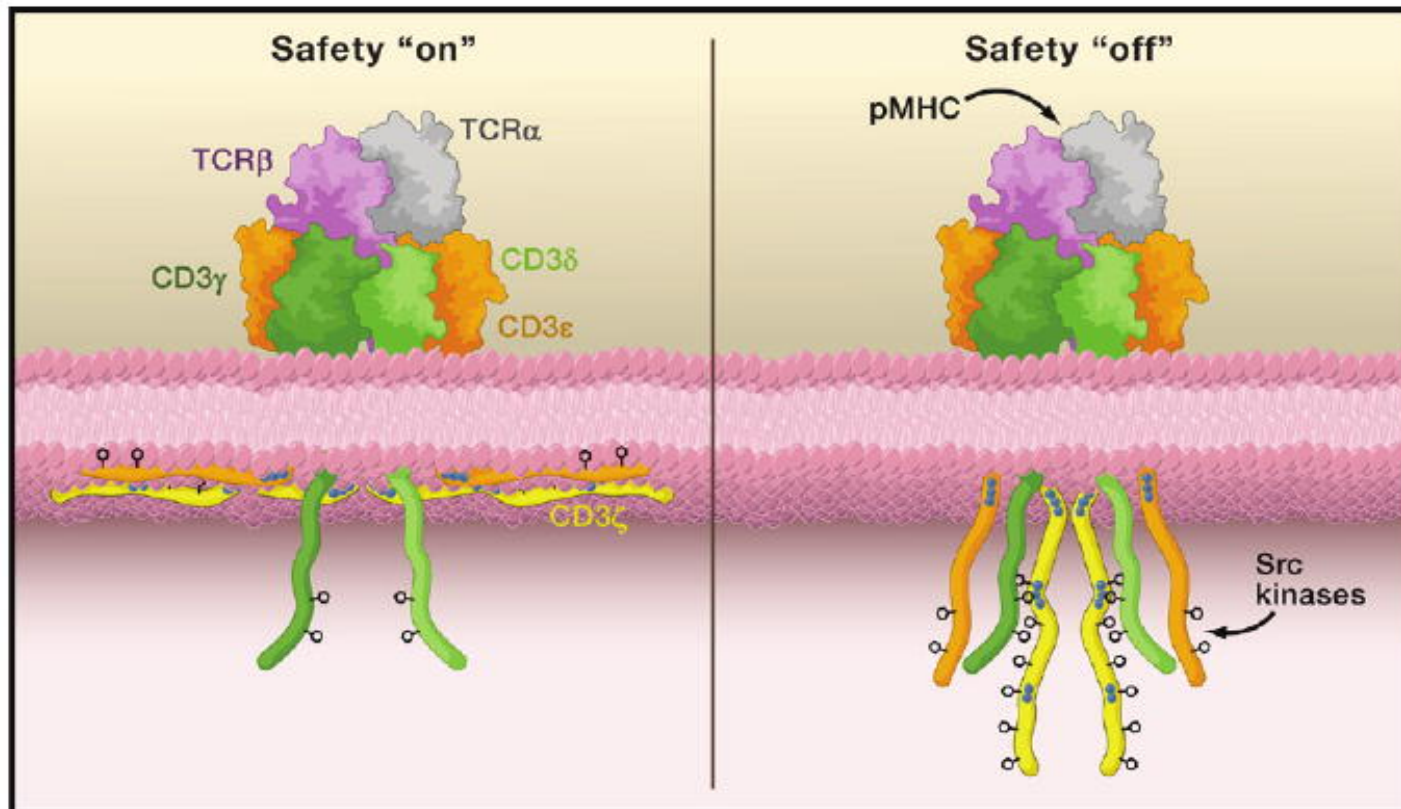
A CD3 ϵ Proline Motif May Become Accessible During a Putative TCR Conformational Change: *Models*

(Alarcon, et al, *Cell*, 2002;
Gil, et al, *J. Exp. Med.*, 2005;
Levin and Weiss, *J. Exp. Med.*, 2005)



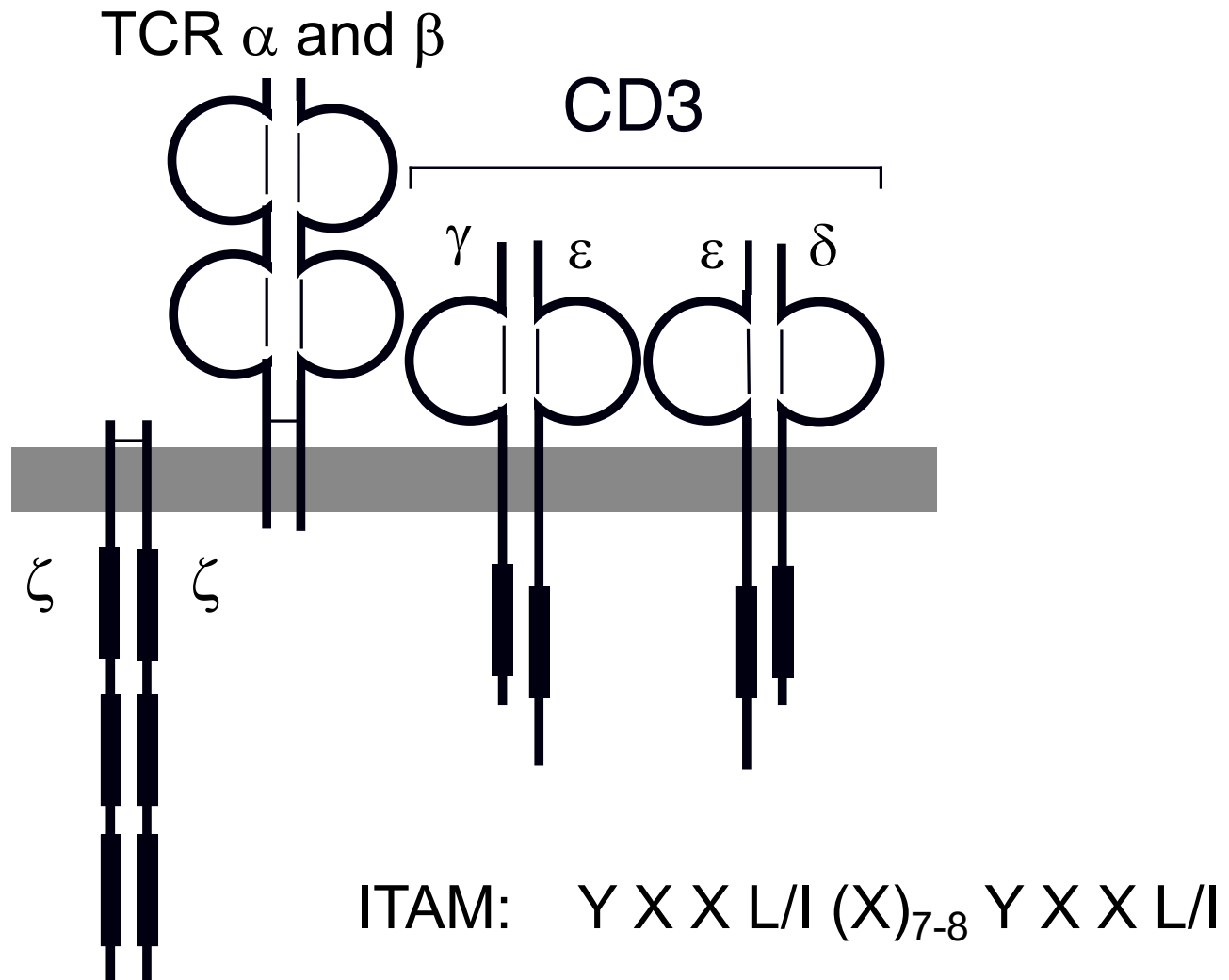
Non-Phosphorylated Tyrosines in ITAMs May be Sequestered by Hydrophobic Interactions with Lipids in Plasma Membrane

Liu, et al., Cell, 2008

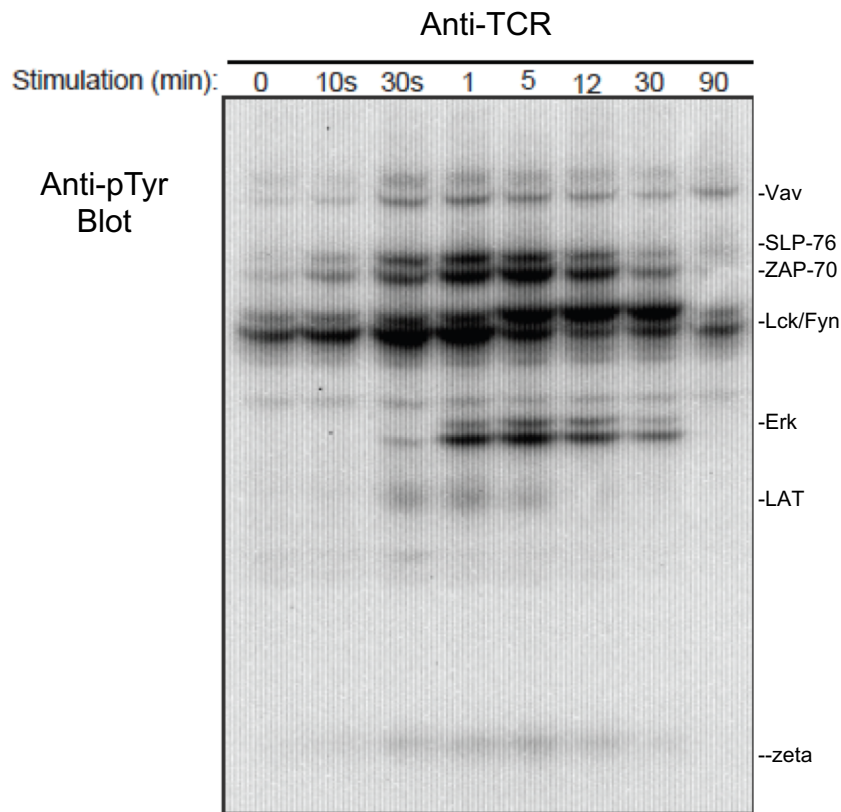


Preview by Kuhns and Davis, Cell, 2008

The TCR is an Oligomer and Utilizes ITAMs to Initiate Signal Transduction



TCR Cross-linking Induces Rapid Protein Tyrosine Phosphorylation

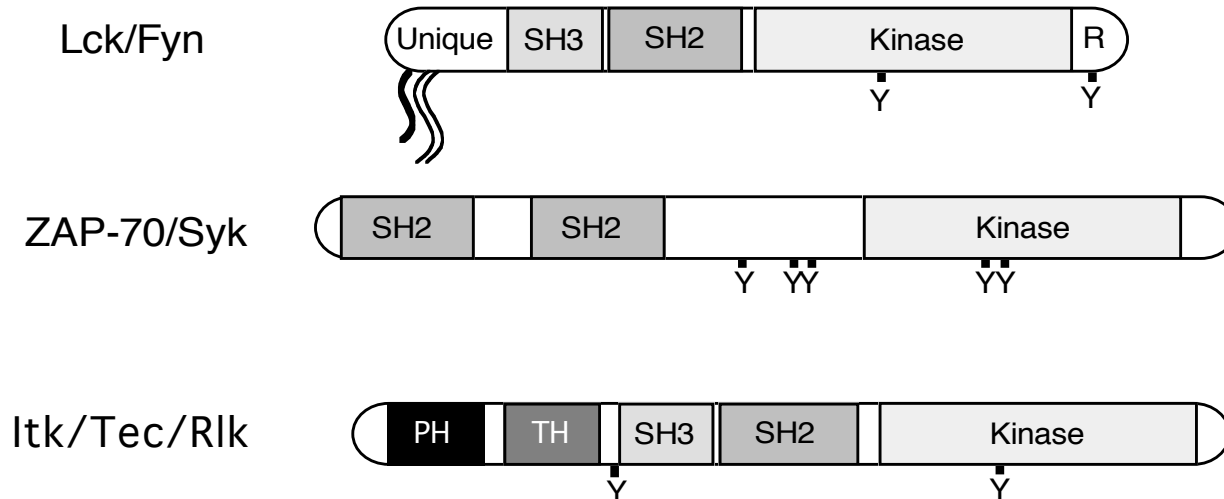


TCR signaling is initiated by protein tyrosine phosphorylation which can:

- Induce conformational changes
- Promote recruitment to the membrane
- Promote protein-protein interactions
- Activate enzymes (kinases, phosphatases, PI3-kinase, PLC- γ 1)
- Inactivate enzymes
- Promote ubiquitination

Jurkat T cell Leukemic Line
stimulated with anti-TCR mAb

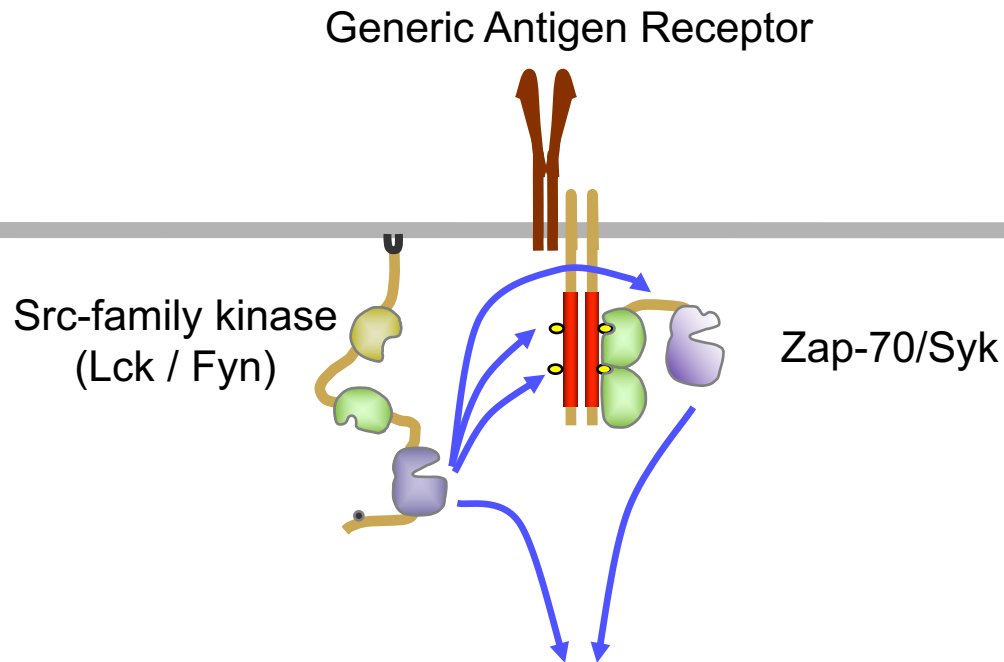
Three Families of Cytoplasmic Protein Tyrosine Kinases Are Involved in TCR Signal Transduction



Protein Interaction modules

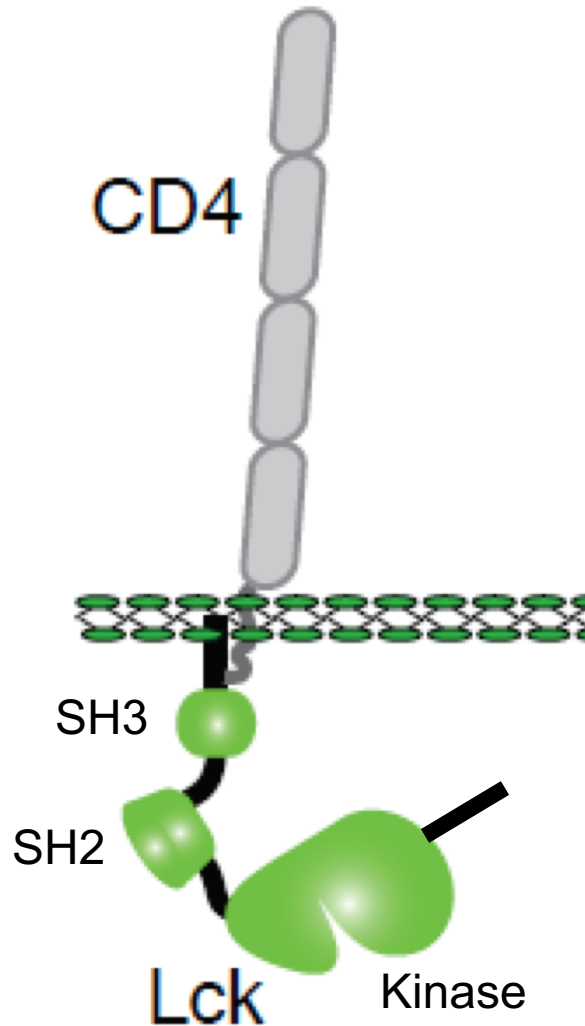
- SH2 Domains** - Interact with tyrosine phosphorylated sequences
- SH3 Domains** - Interact with proline-rich sequences
- PH Domains** - Interact with membrane phospholipids
- TH Domains** - Unique to Tec family kinases - proline rich

SFKs Are Critical Initiators of Antigen Receptor Signaling, So their Regulation is Important



Tyrosine phosphorylation, MAPK activation (P-Erk)
Intra-cellular calcium increase...

Lck, the Kinase that Initiates TCR signaling



Associates non-covalently
with **CD4** and **CD8** via
C-C rich motifs

Src Kinase Family Member

Peripheral Membrane Protein

Palmytoylated
Myristoylated
Enriched in lipid rafts

Lipid rafts, GEMs, DIGs

Cholesterol-rich microdomains in the plasma membrane
(25 to 70 nm in diameter)

Enriched in GPI-linked receptors and glycosphingolipids

Enriched in many signaling molecules, including:
PIP2 and PIP3, several GTPases including Ras,
Src kinases, some heterotrimeric G-proteins

Operationally defined based on detergent solubility

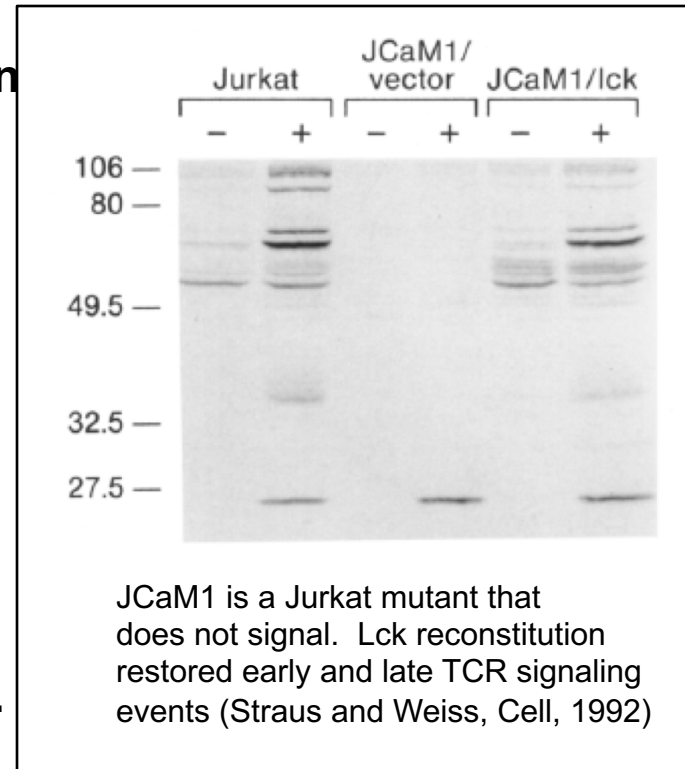
Some of Lck (50% or less) partitions into lipid rafts
(reversibility and dynamic changes?)

May be a focal point of signal initiation or sequestration

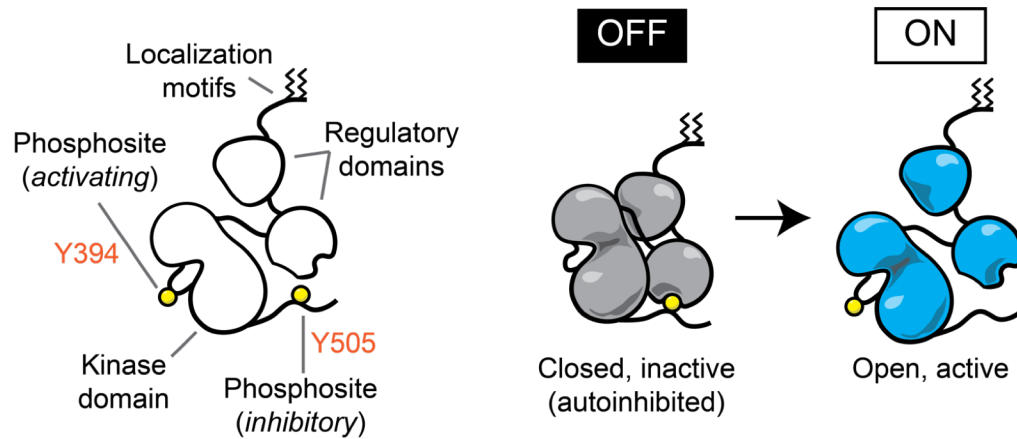
- A topic of considerable controversy!

Lck is Involved in TCR Signaling

- A pool of active Lck exists in resting T cells and may not increase following TCR stimulation
- Repositioning Lck may be most critical.
- KO in mice results in a relative block in thymocyte development.
- Lck-deficient T cells have a TCR signaling defect (ITAMs are not phosphorylated, nor are distal events induced).
- CD4-Lck or CD8-Lck associations are required for optimal antigen responsiveness.

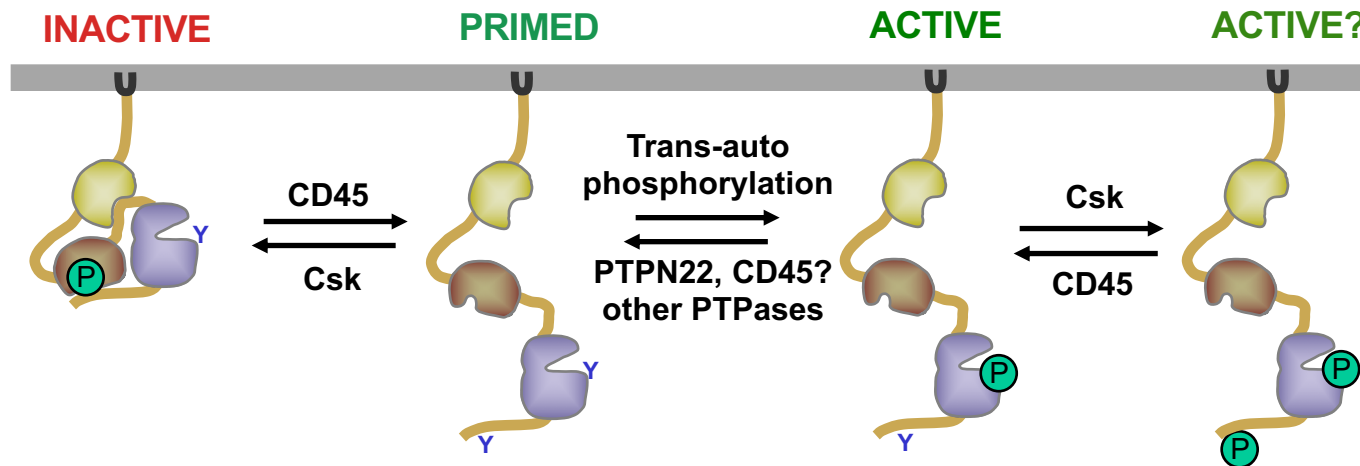


Regulating Src Family Kinases

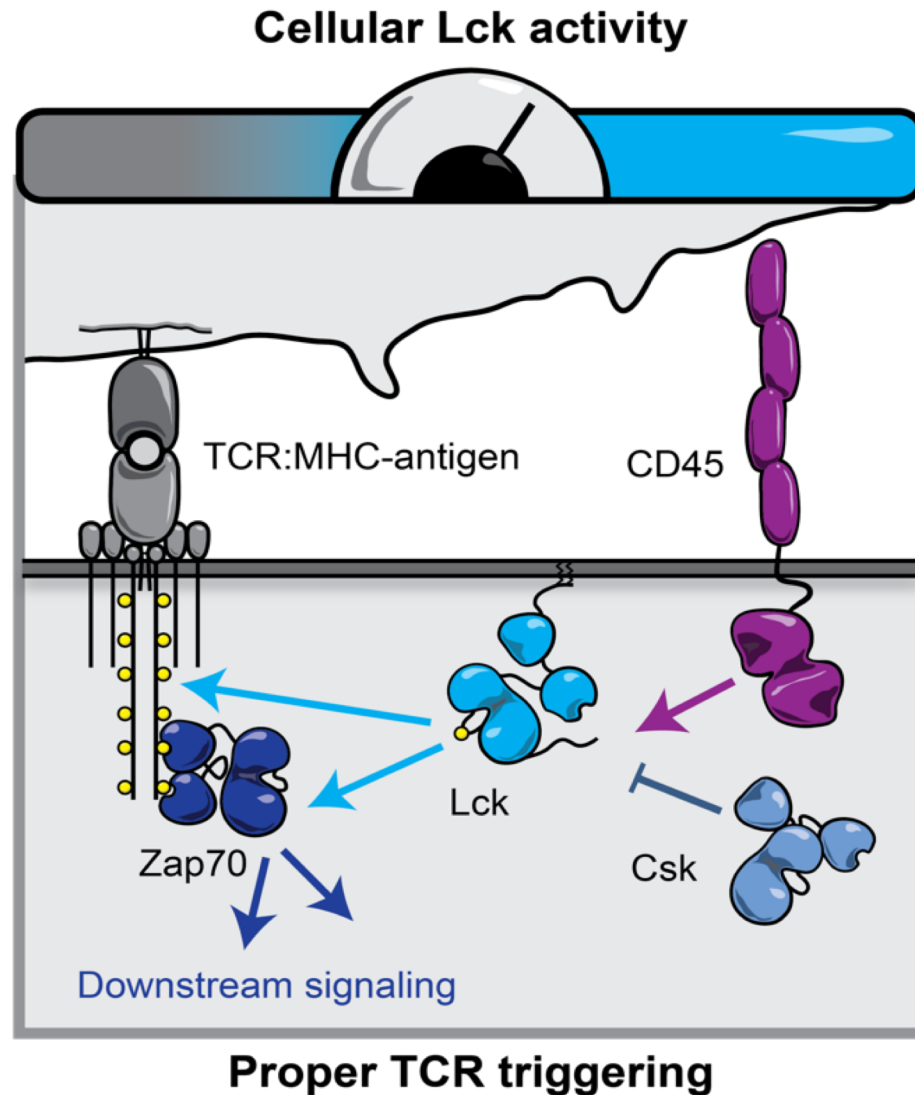


- Lck and Fyn are Src family kinases found in T cells
- Conformation determines activity
- Phosphorylation regulates conformation & therefore activity

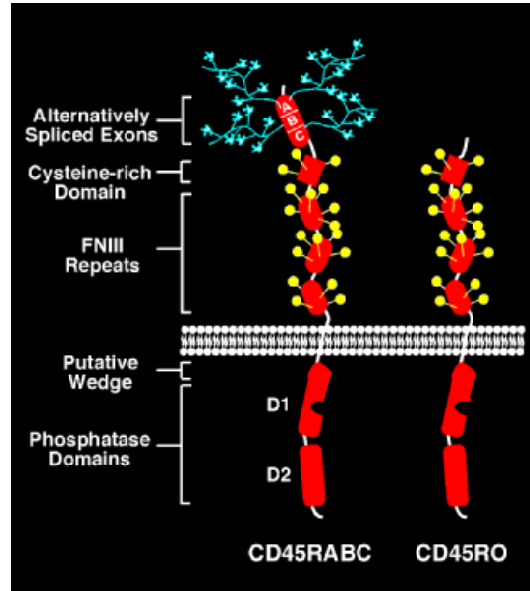
Various activation states of SFKs in T cells



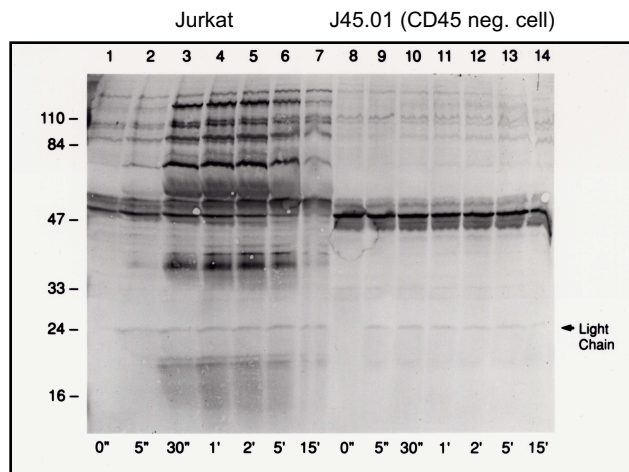
A dynamic interplay of CD45 and Csk is important in setting the level of Lck activity



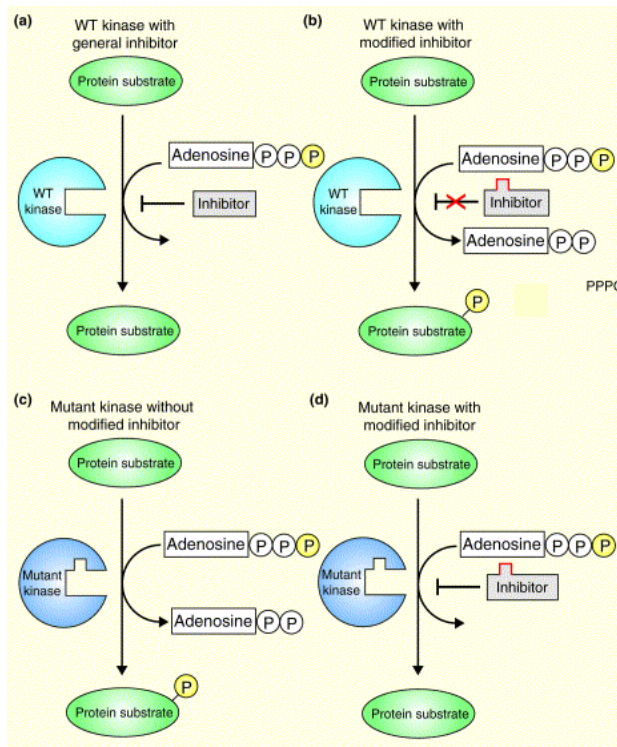
CD45, a Tyrosine Phosphatase Plays a Positive and a Negative Regulatory Role for TCR Induced Tyrosine Phosphorylation



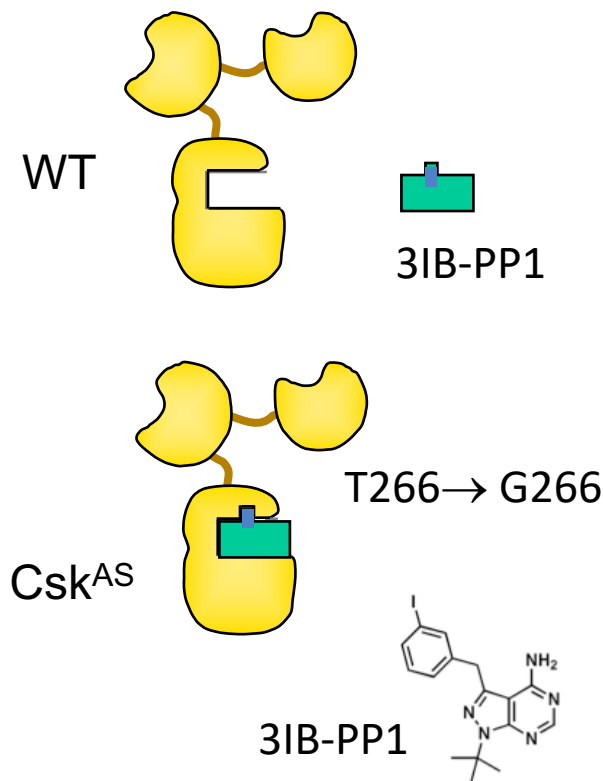
- Receptor-like protein tyrosine phosphatase expressed on all nucleated hematopoietic cells, expressed at very high levels (25 μ M)
- Distinct isoforms expressed in a cell-type and developmental-stage specific manner
- Required for T cell and B cell antigen receptor signaling and for normal development - dephosphorylates negative regulatory tyrosine in Src kinases.
- Mutations associated with SCID and autoimmunity in humans and mice
- Jurkat and T cell clone deficient in CD45 does not signal
- Contains tandemly duplicated phosphatase domains, but only domain 1 is functionally active.



A unique analog-sensitive allele of Csk can be rapidly and specifically inhibited

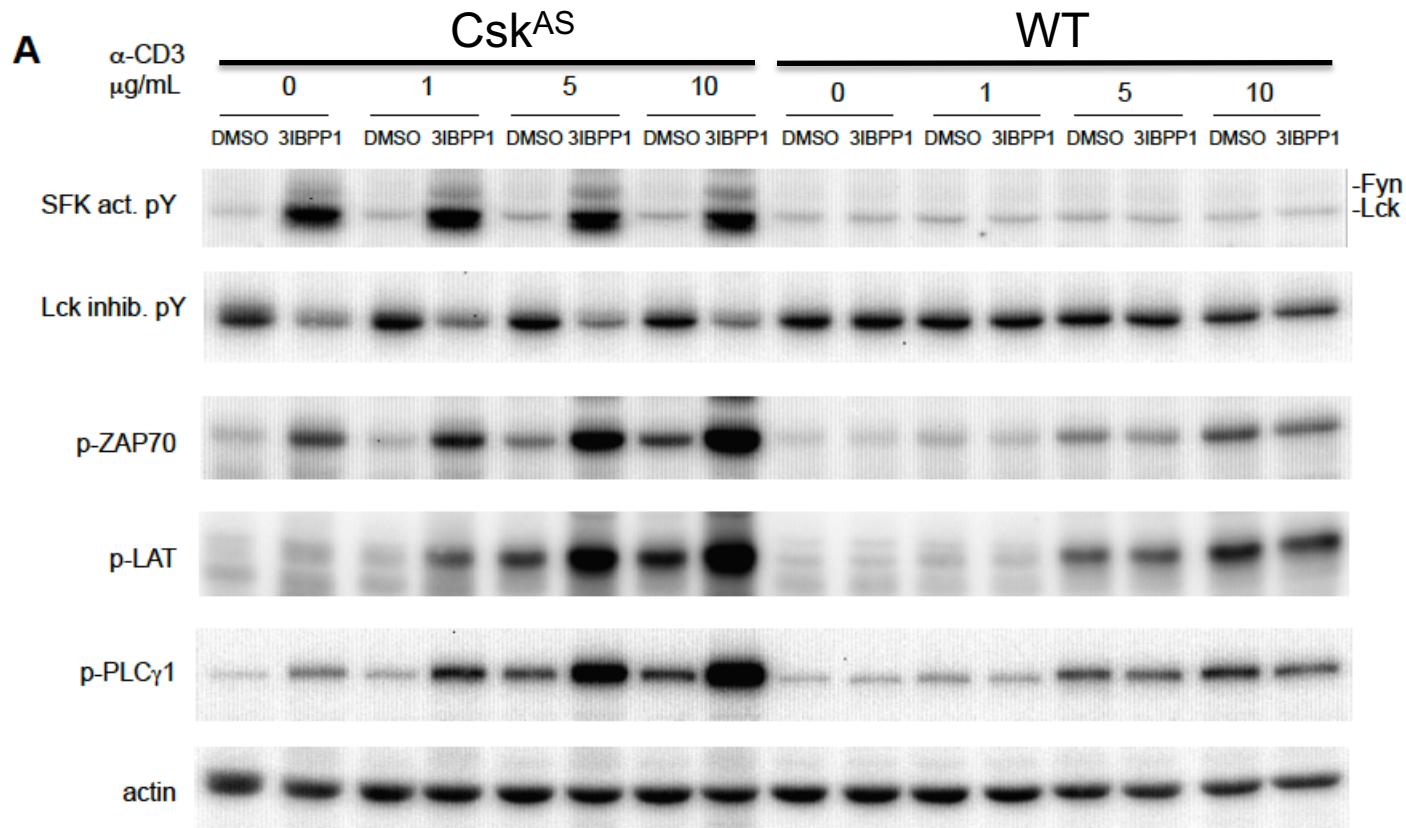


Alaimo et al, Curr Op Chem Bio, 2001

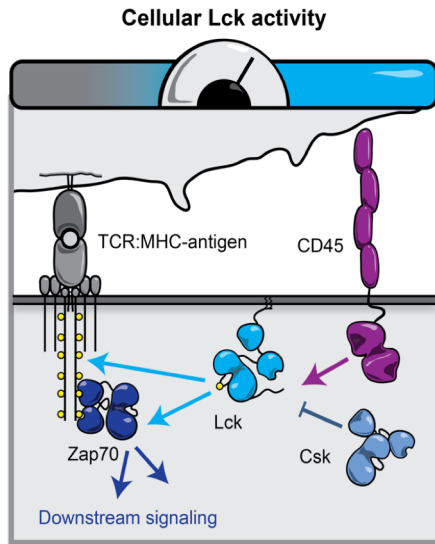


Synergy of Csk inhibition and TCR stimulation occurs downstream of Lck phosphorylation

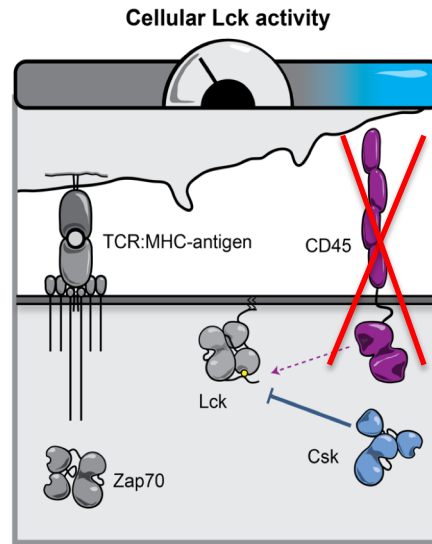
Mouse CD4 T cells



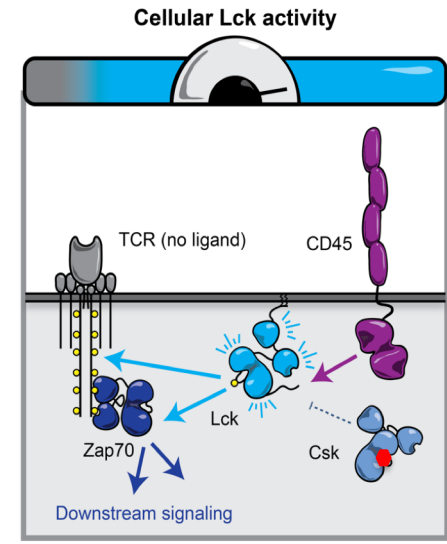
Lck activity is required and sets a threshold for T cell activation



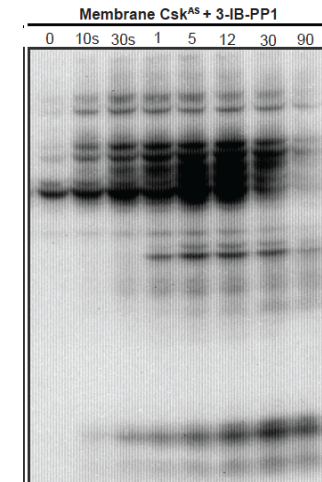
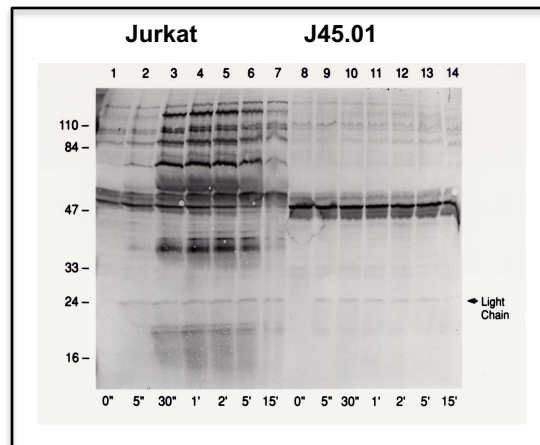
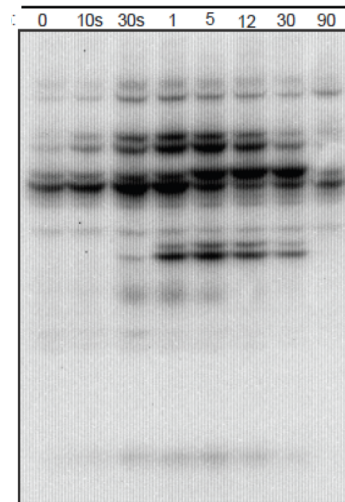
Proper TCR triggering



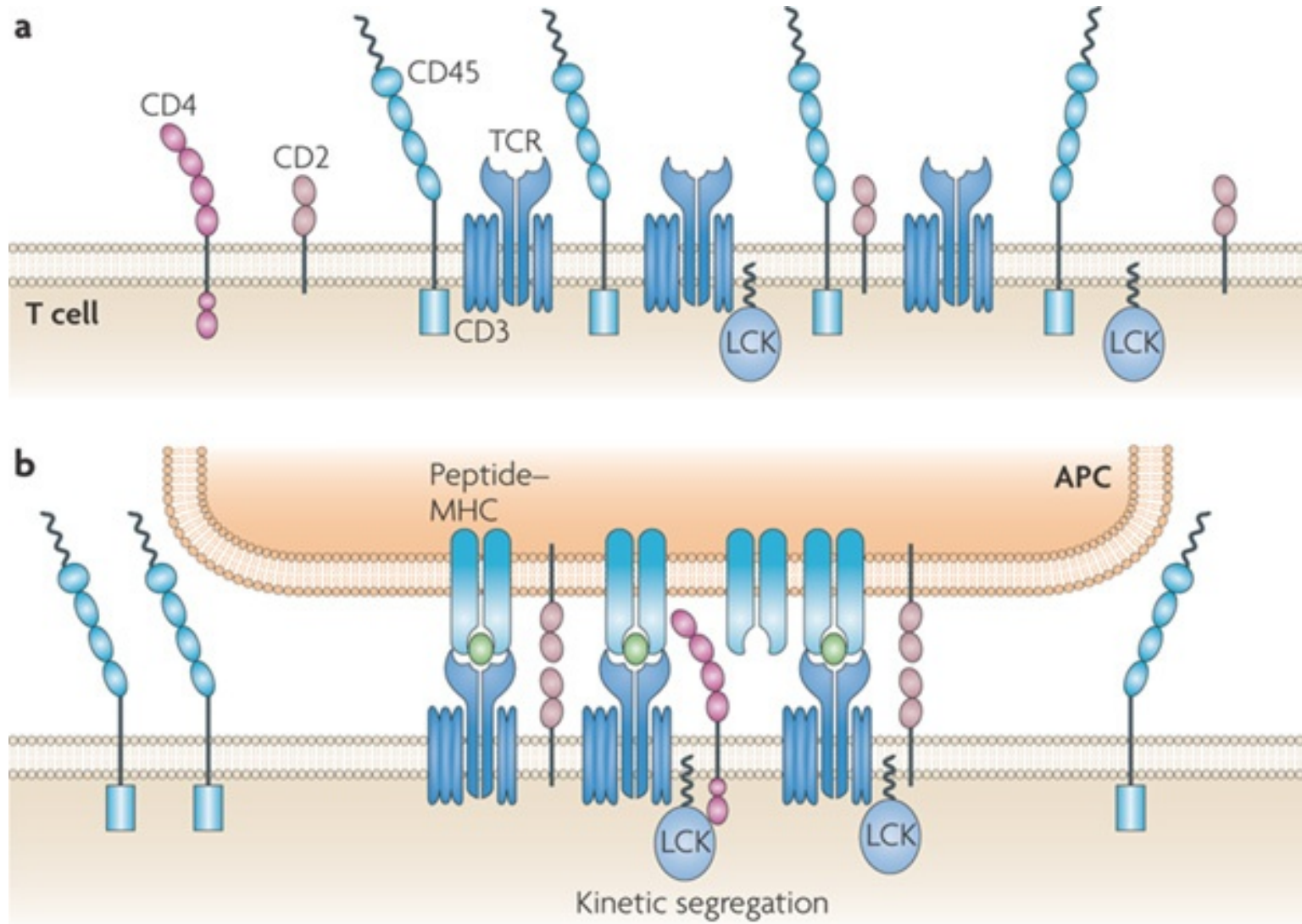
Decreased Lck activity - Ignorance



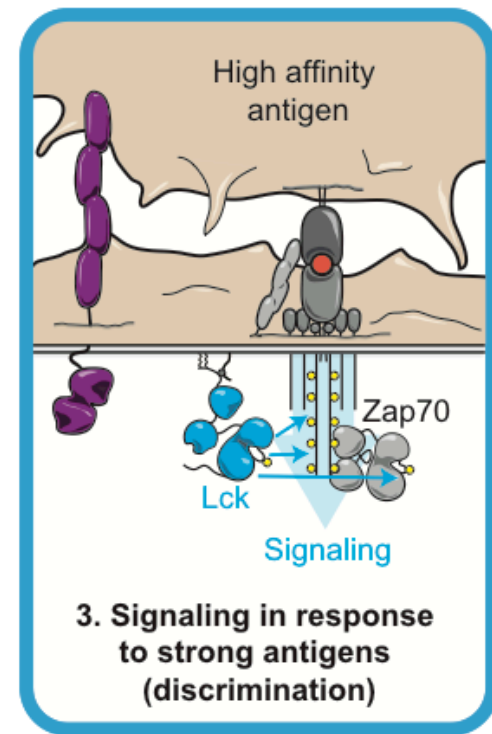
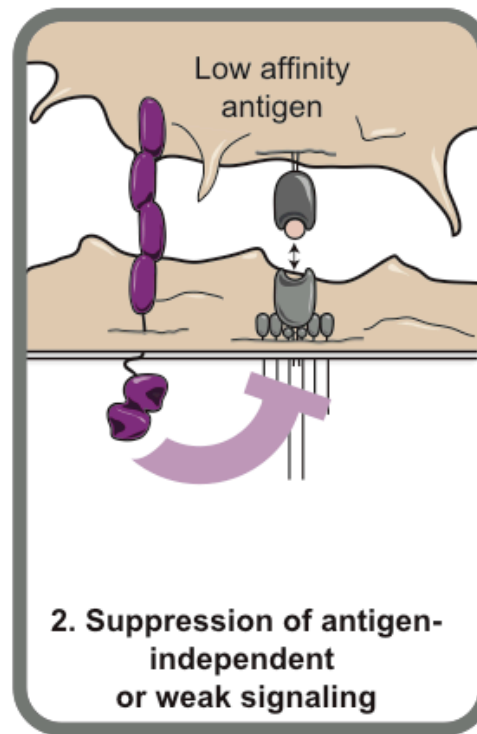
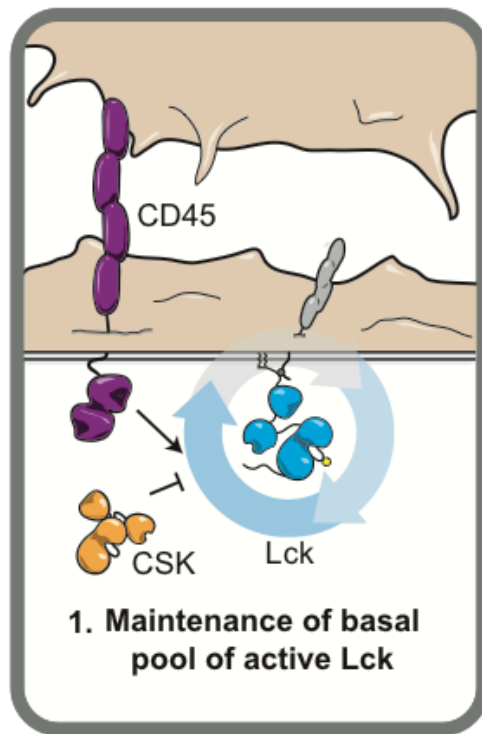
Increased Lck activity - Increased triggering



Kinetic Segregation Model: Separation of CD45 PTP via Large EC Domain Segregates PTP activity from pTyr Events

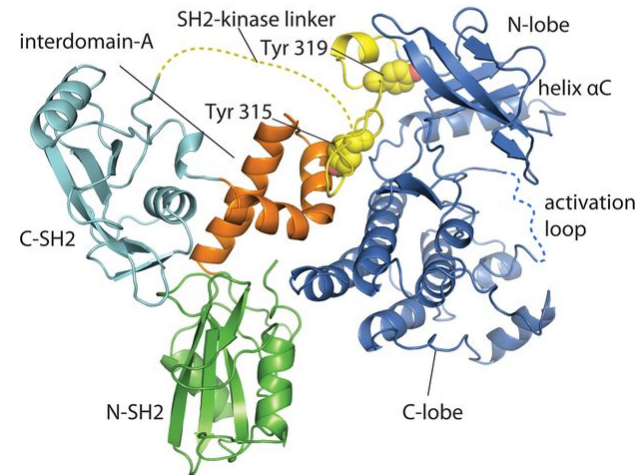
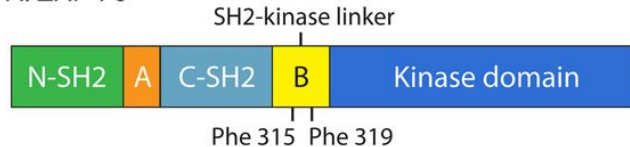


TCR ligand discrimination can be influenced by the relative positive and negative regulators of Lck and by ligand potency



ZAP-70 in TCR Signaling

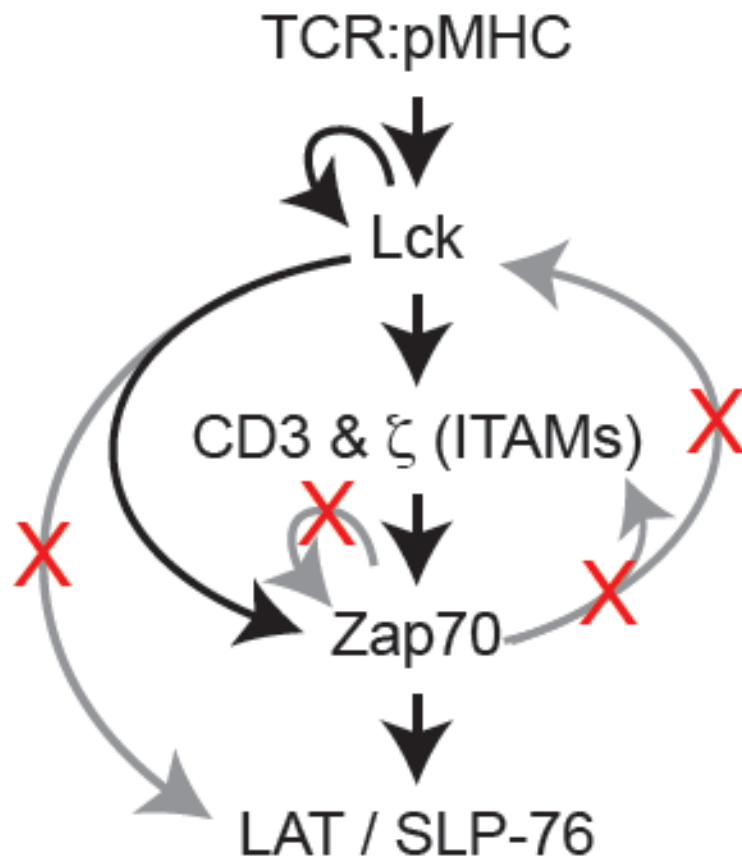
A. ZAP-70



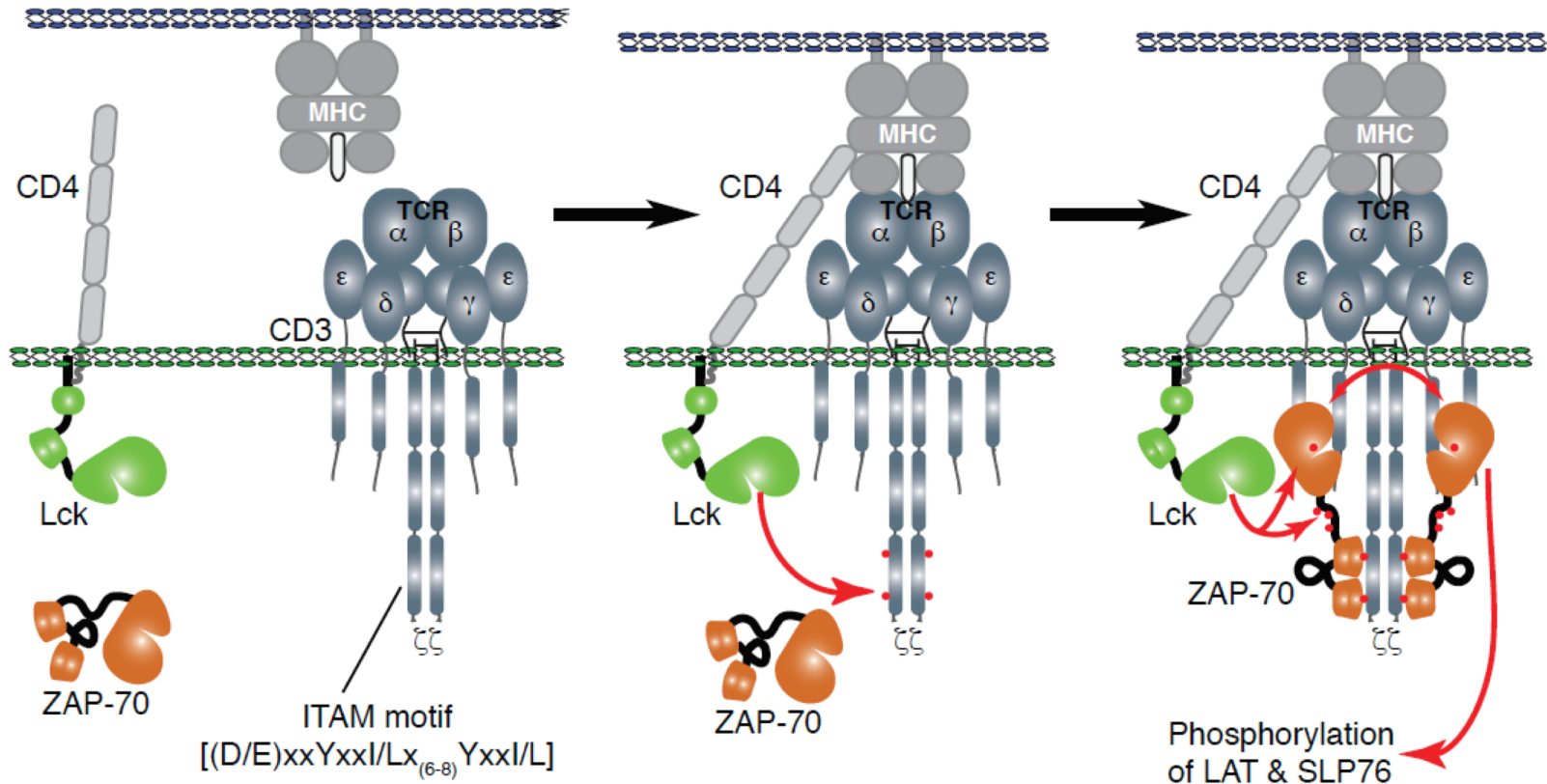
- Expressed only in T cells and NK cells
- Cooperative binding of tandem SH2 domains to doubly phosphorylated ITAMs
- Substrates: LAT, SIp-76
- ZAP-70 LOF mutations:
 - Cell line - loss of TCR signaling
 - Mouse KO - developmental arrest (DP → SP)
 - Human SCID - No CD8 T cells
 - CD4 T cells with TCR signaling defect

**Why did Zap70 evolve?
Why isn't Syk good enough?**

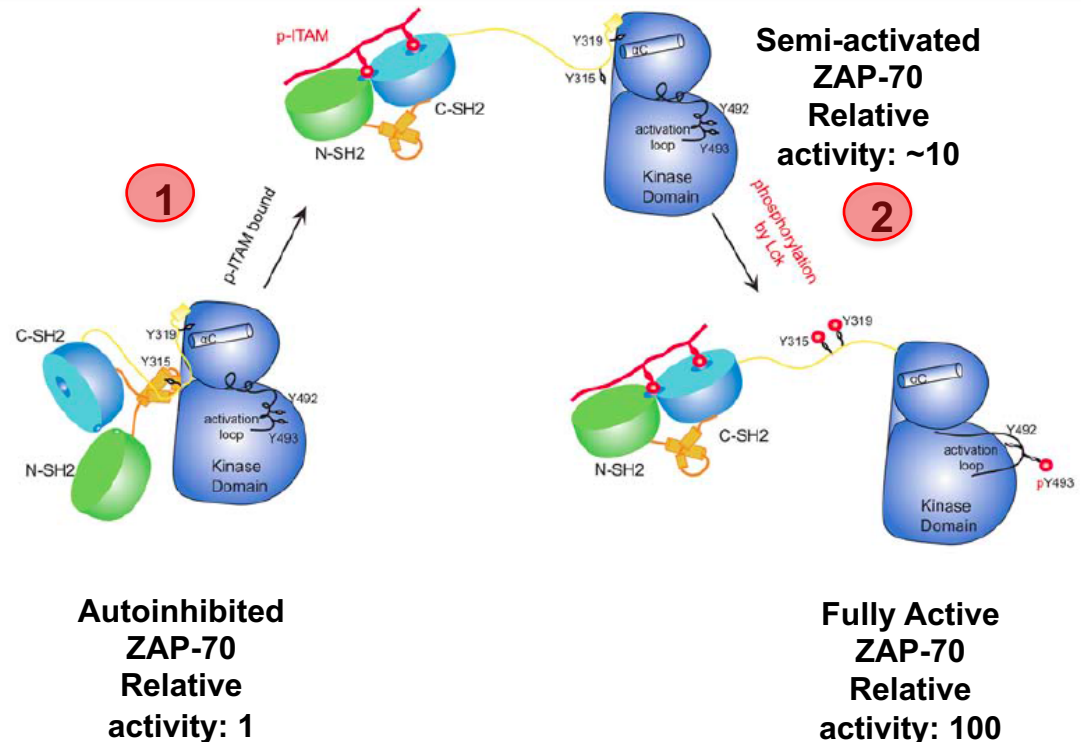
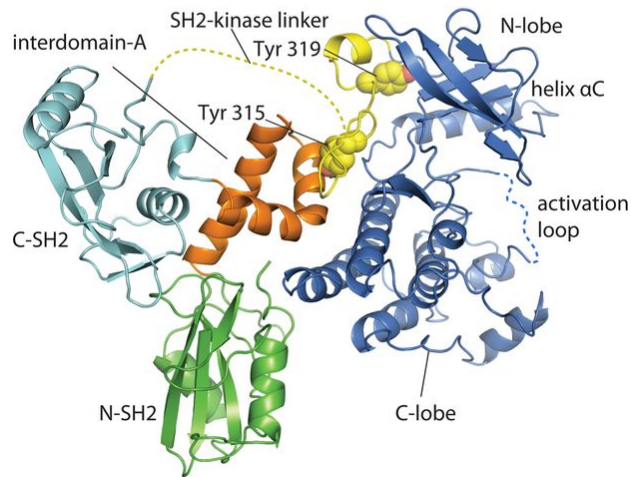
TCR signaling hierarchy



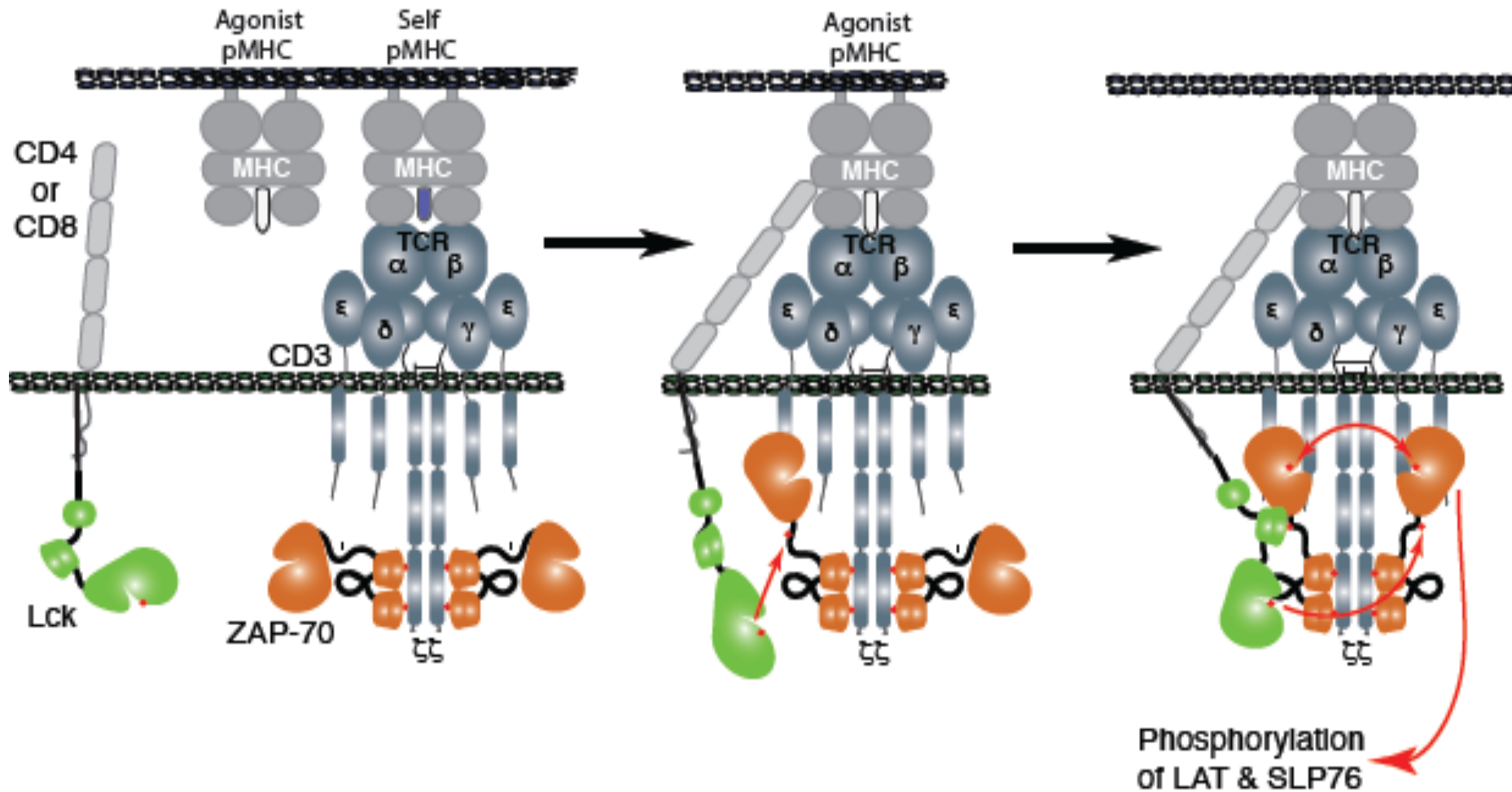
Model requiring TCR and coreceptor (CD4 or CD8) function: Zap70 and Lck evolved to enforce a requirement for MHC recognition by the TCR



Two Allosteric Changes Involving ITAM Binding and Lck-Mediated Phosphorylation are Necessary for ZAP-70 activation



Lck phosphorylation of and its binding to ZAP-70 that is already bound to the TCR- ζ chain may be Critical for Activation of the TCR

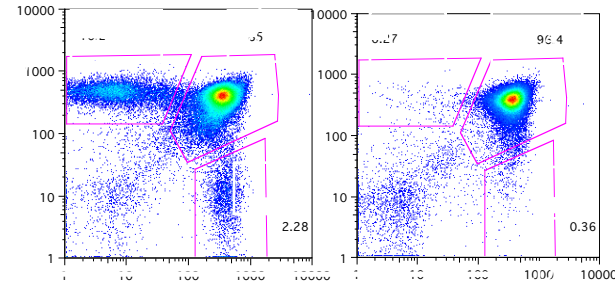


ZAP-70 in Disease

ZAP-70 deficiency

Human (CD8 T cell-deficient SCID)

Mouse (T cell-deficient SCID)



Spontaneous mouse model of arthritis

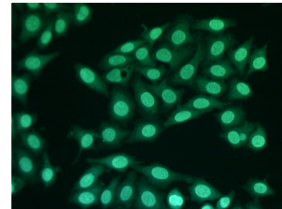
Hypomorphic *skg* allele – alters development

(Sakaguchi, et al., Nature, 2005)



Catalytic domain ENU mouse mutants - hyper-IgE and autoimmunity

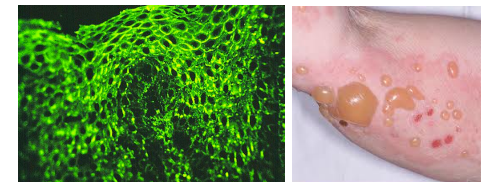
Level of TCR signaling alters development (Siggs, et al, Immunity, 2007)



SH2 + catalytic domain mutants – human autoimmunity

Bullous pemphigoid, anti-Factor VIII antibodies,

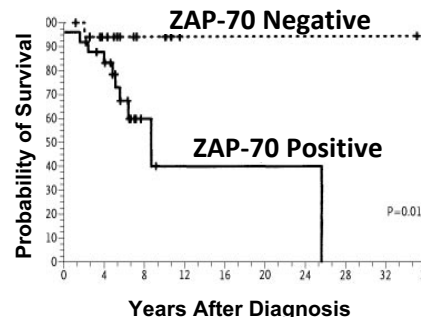
Glomerulonephritis, IBD (Chan, et al., J Exp Med, in press)



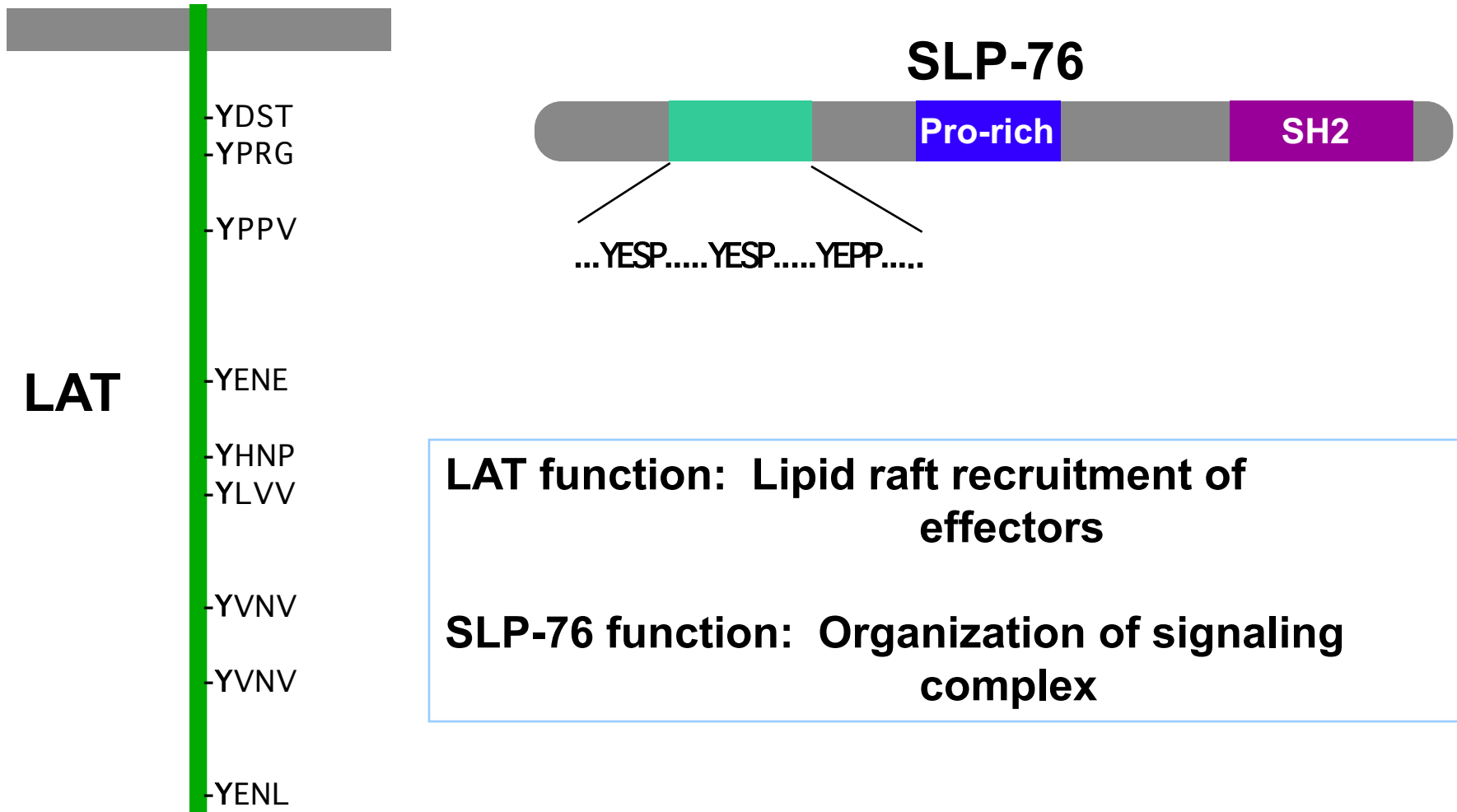
Aberrant expression in B cell CLL

ZAP-70+ CLL: Poor prognosis

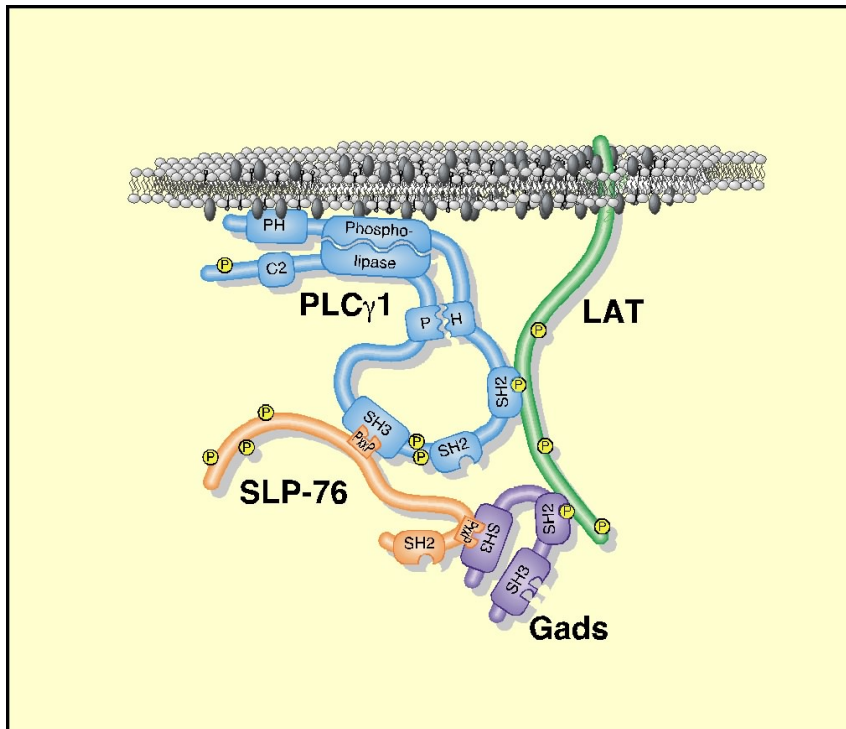
(Rassenti, et al., NEJM 2004)



Two Adapters Are ZAP-70 Substrates: Critically Required for TCR Downstream Signaling



Assembly of LAT and SLP-76 into a Signalosome in Lipid Rafts Is required for Multiple Downstream TCR Signaling Pathways

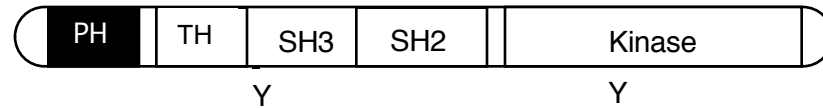


Events requiring LAT & SLP-76

PLC γ 1 activation
Ras activation
Rac activation
HPK1 activation

Tec Kinases in T cells

Itk/Tec/Rlk

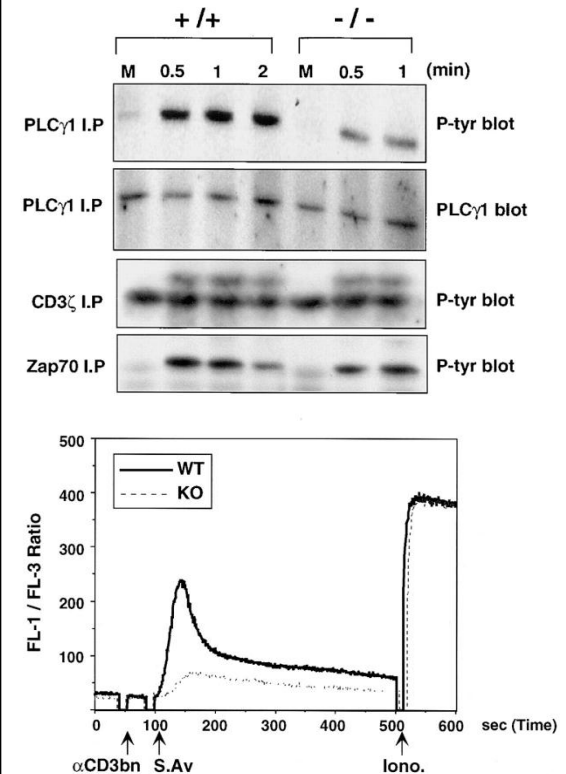


Itk

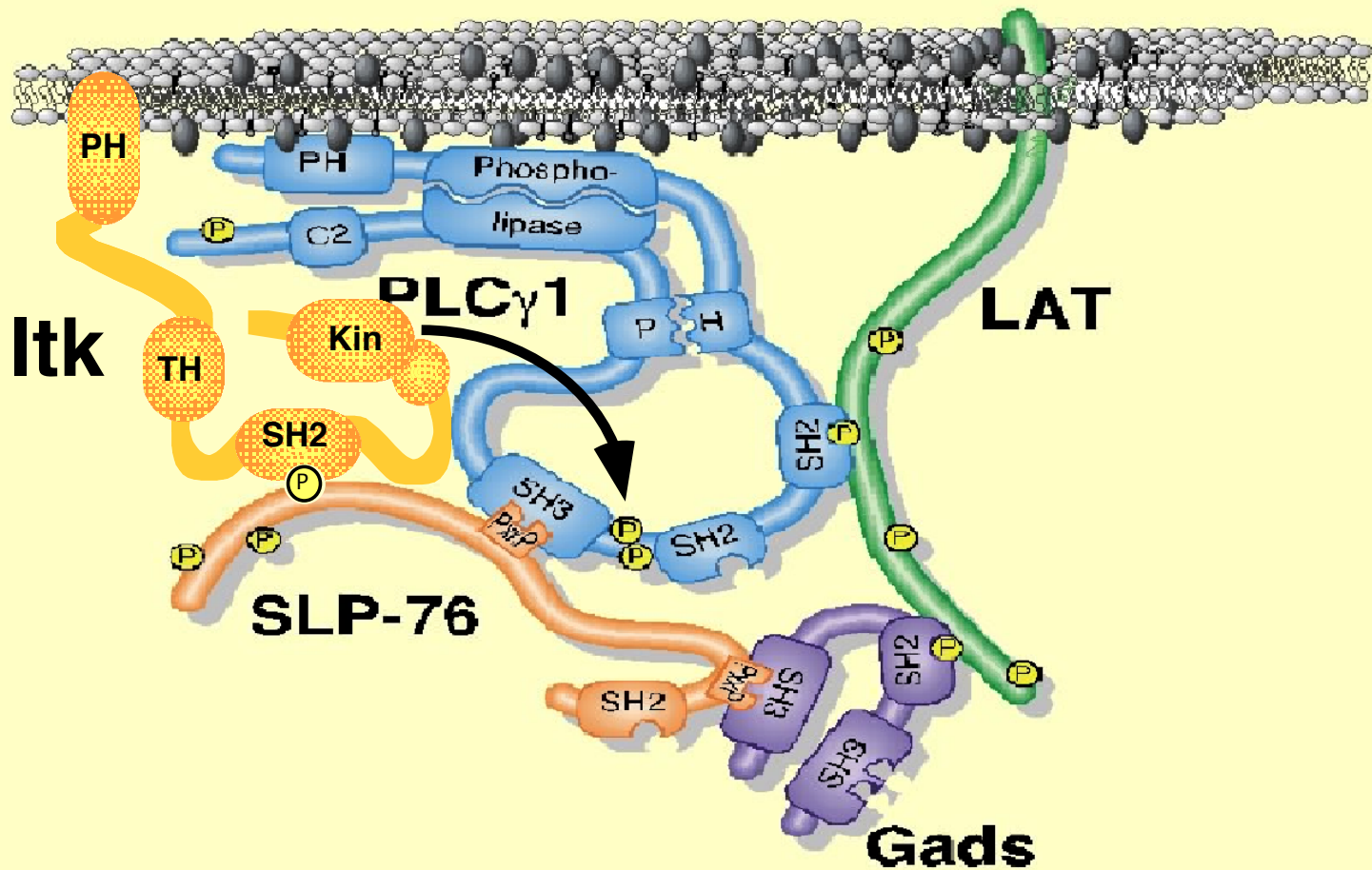
- Expressed predominantly in T cells
- PH domain binds to PIP3 which recruits Itk to membrane
- Kinase domain is phosphorylated and activated (by Lck)
- TH and SH3 domains interact in basal state to inhibit kinase
- Autophosphorylation of Y180 adjacent to SH3 domain allows Itk to unfold, rendering SH2 and SH3 domains accessible
- Likely to phosphorylate and activate PLC γ 1
- KO in mice results in mild defect in PLC activation and Ca²⁺ increase (redundant roles of other family members, Rlk and Tec, in T cells)

Defective PLC γ 1 phosphorylation and Ca²⁺ increase in Itk null T cells

Liu, et al., JEM, 1998



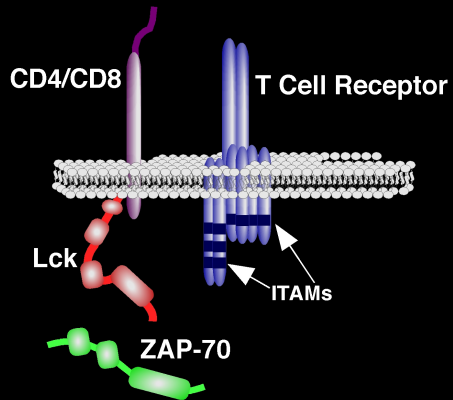
Itk is recruited to the Signalosome and Phosphorylates PLC γ 1



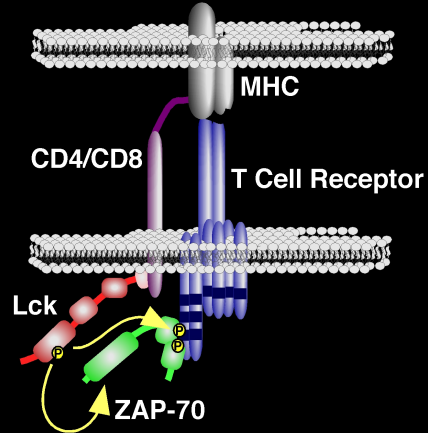
Sequential Events in TCR Signal Transduction

A

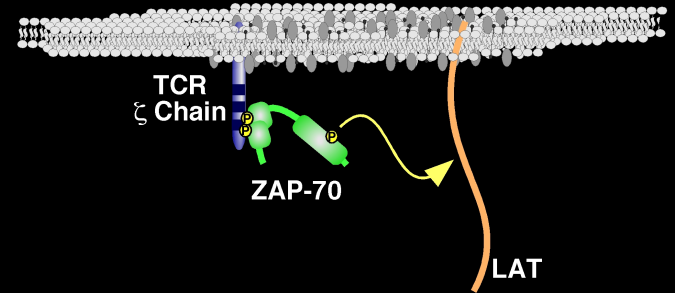
Resting



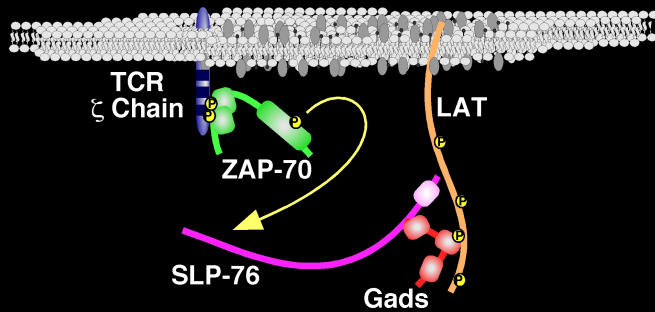
Stimulated



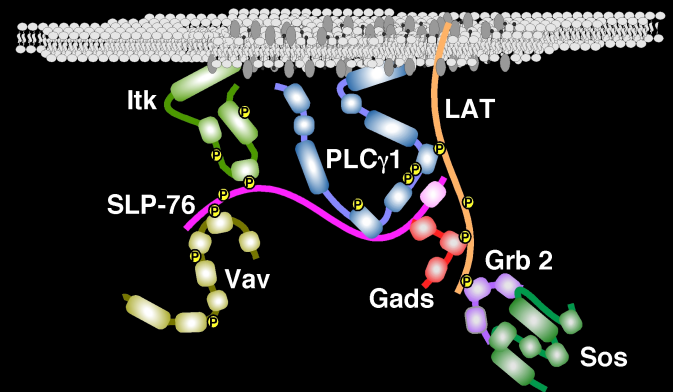
B



C



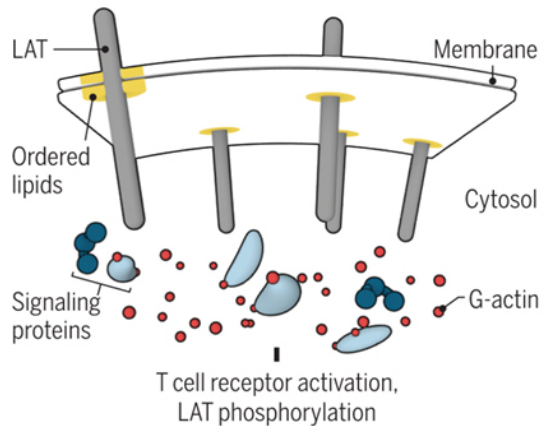
D



Dynamic In Vitro Assembly of Signaling Components

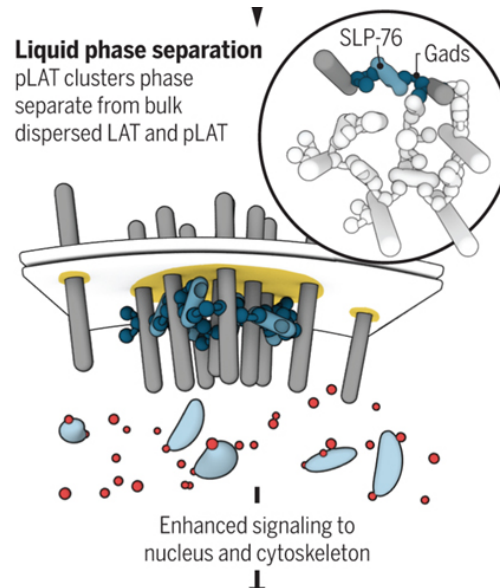
Dispersed bulk phase

Unphosphorylated LAT proteins span the membrane but favor ordered membrane regions (yellow)



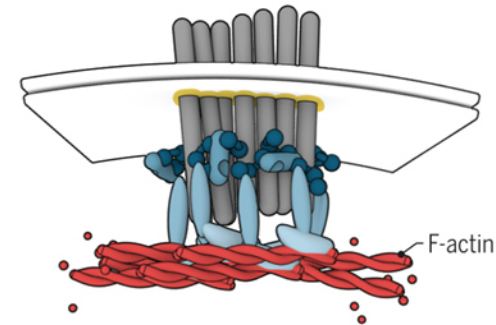
Liquid phase separation

pLAT clusters phase separate from bulk dispersed LAT and pLAT



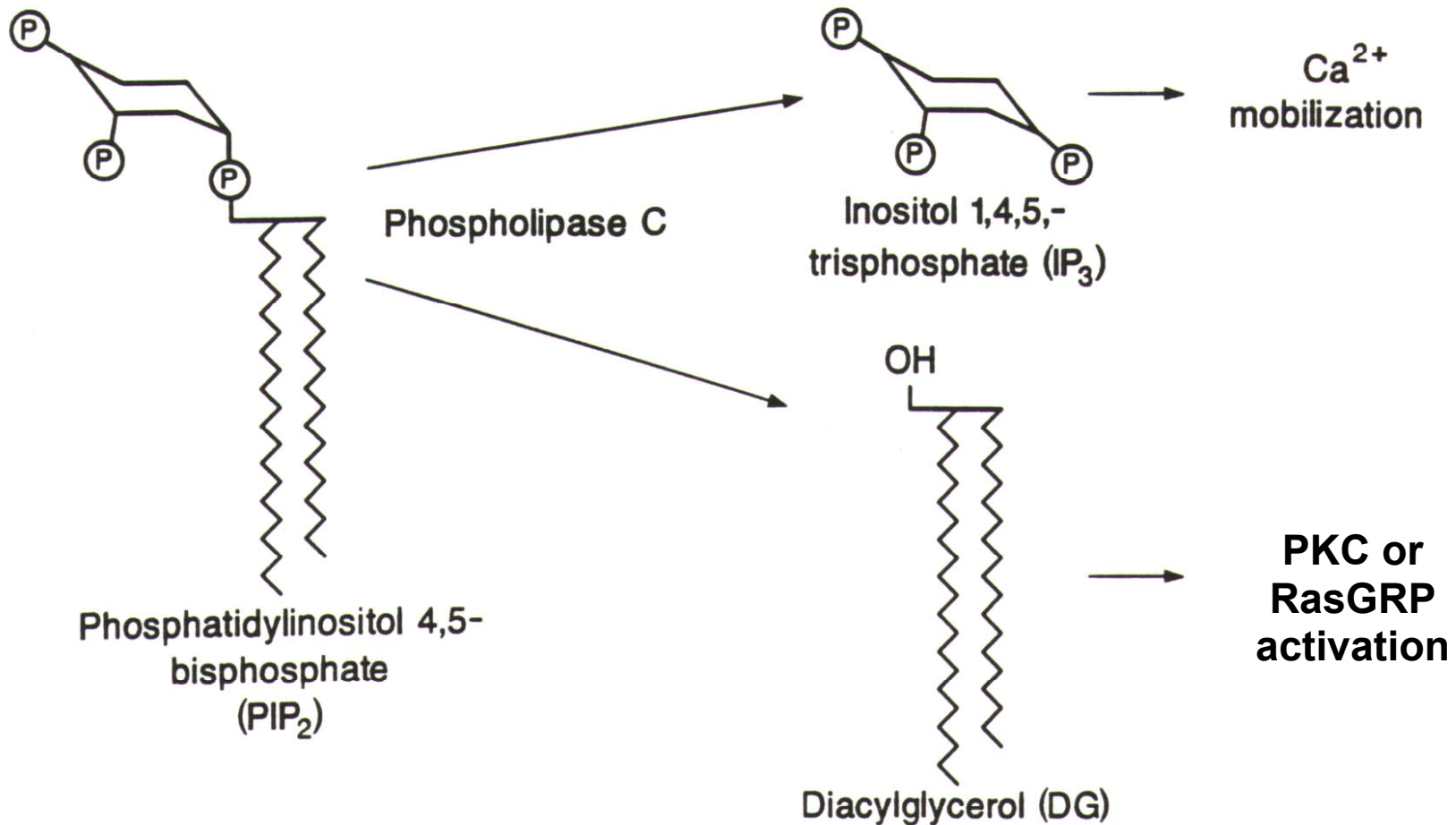
Solid phase scaffold

The pLAT cluster triggers, then takes the shape of, an F-actin gel



Dustin and Muller, Science, 2016
Su, et al., Science, 2016

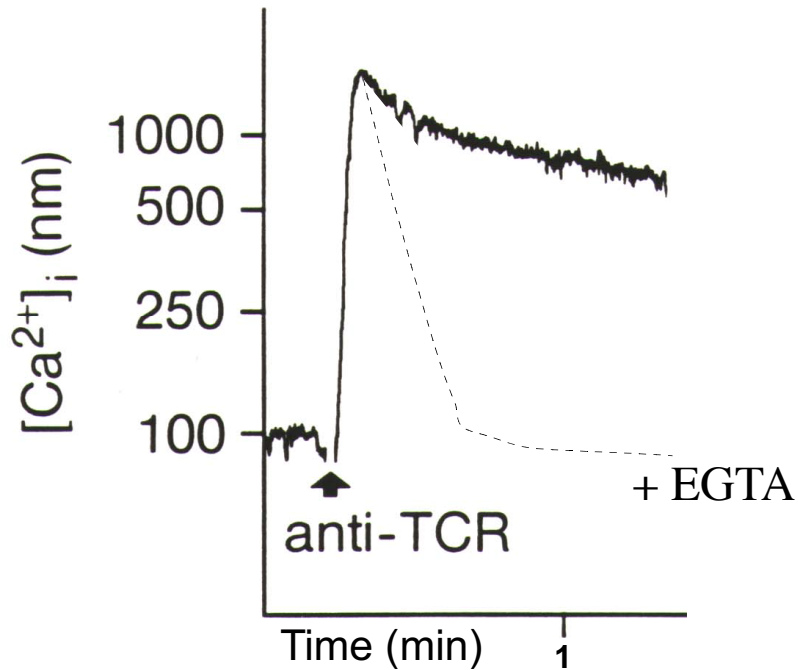
PLC Activation Leads to Second Messenger Generation and Mobilization of Calcium as well as PKC and RasGRP Activation



Evidence for the Importance of the Phosphatidylinositol Pathway in T cells

1. Activating and synergistic effects of calcium ionophores and phorbol esters
2. Inhibitory effects of calcium chelators and PKC inhibitors
3. Positive effects of activating alleles of calcineurin and PKC or Ras)
4. Ability of heterologous G-protein coupled receptors that activate PLC beta but not Src or Syk PTKs to induce IL-2 gene activation
5. T cell activation defects in PKC theta or RasGRP1 mice

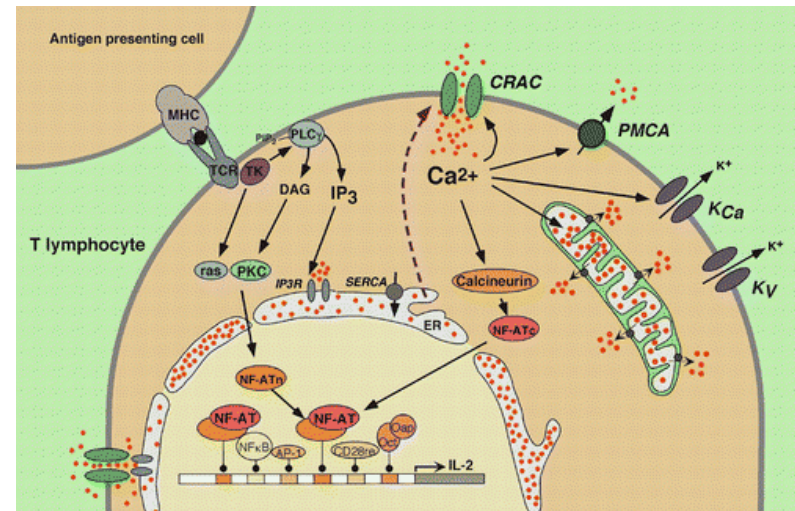
The Phosphatidylinositol Pathway: Calcium Mobilization after TCR Stimulation



Initial release from Intracellular stores
(results from IP₃ interacting with receptors)

Depletion of intracellular stores results in a
transmembrane flux of calcium, I_{crac}

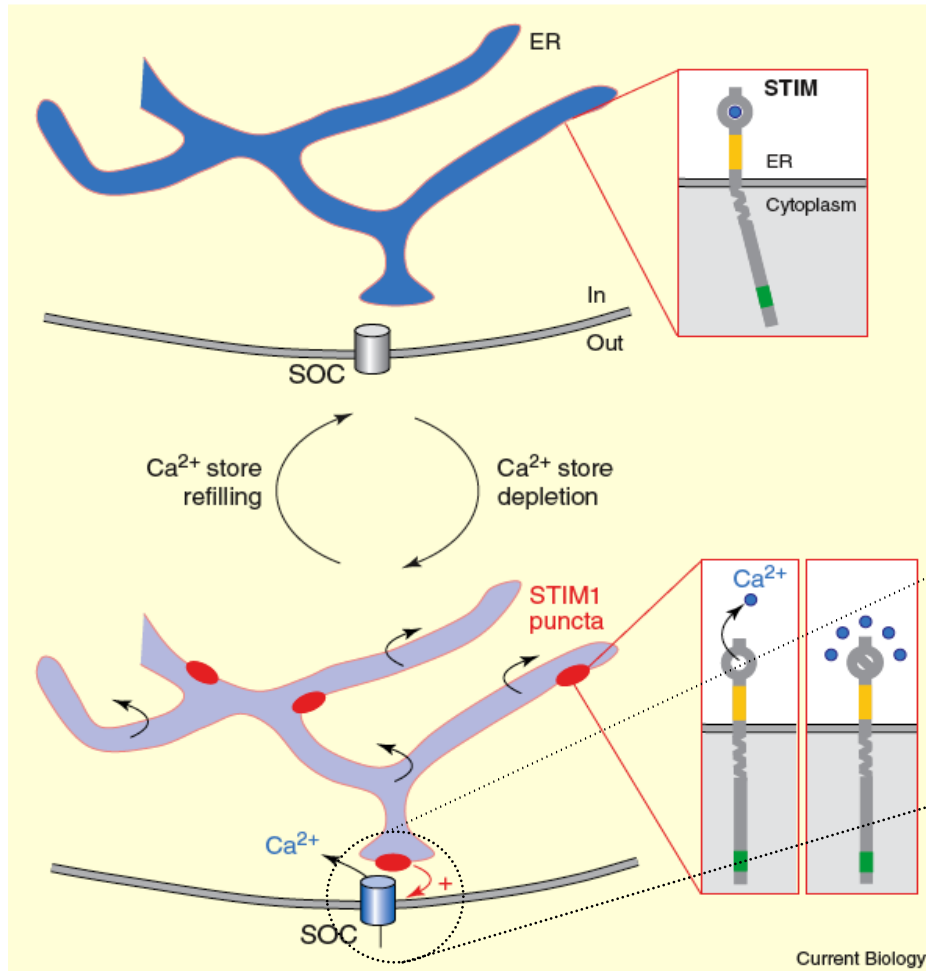
Lewis Annu. Rev. Immuno. 2001



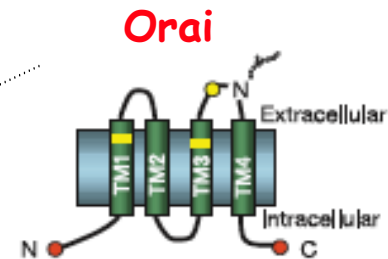
Downstream Consequences:

- Activation of Calcineurin (Ca/CaM-dependent serine/threonine phosphatase), the target of Cyclosporin A and FK506
- Activation of Ca/CaM kinase

STIM is the ER Calcium Sensor and Orai is a Component of Icrac



Depletion of intracellular stores results in altered function and distribution of ER resident protein STIM. STIM migrates in “puncta” towards plasma membrane where it regulates the recently identified Icrac channel, Orai, in the plasma membrane to allow a transmembrane flux of calcium



Orai initially identified as a consequence of tracking down the basis of unusual calcium pathway-defective SCID

The Phosphatidylinositol Pathway: PKC activation after TCR Stimulation

PKC - a large family of serine/threonine kinases

Multiple isozymes in T cells

DAG-regulated, phorbol ester responsive

Some are calcium responsive

PKC theta localizes to cSMAC, and KO mice have selective defect in NF-kB activation in T cells

Function:

TCR downregulation

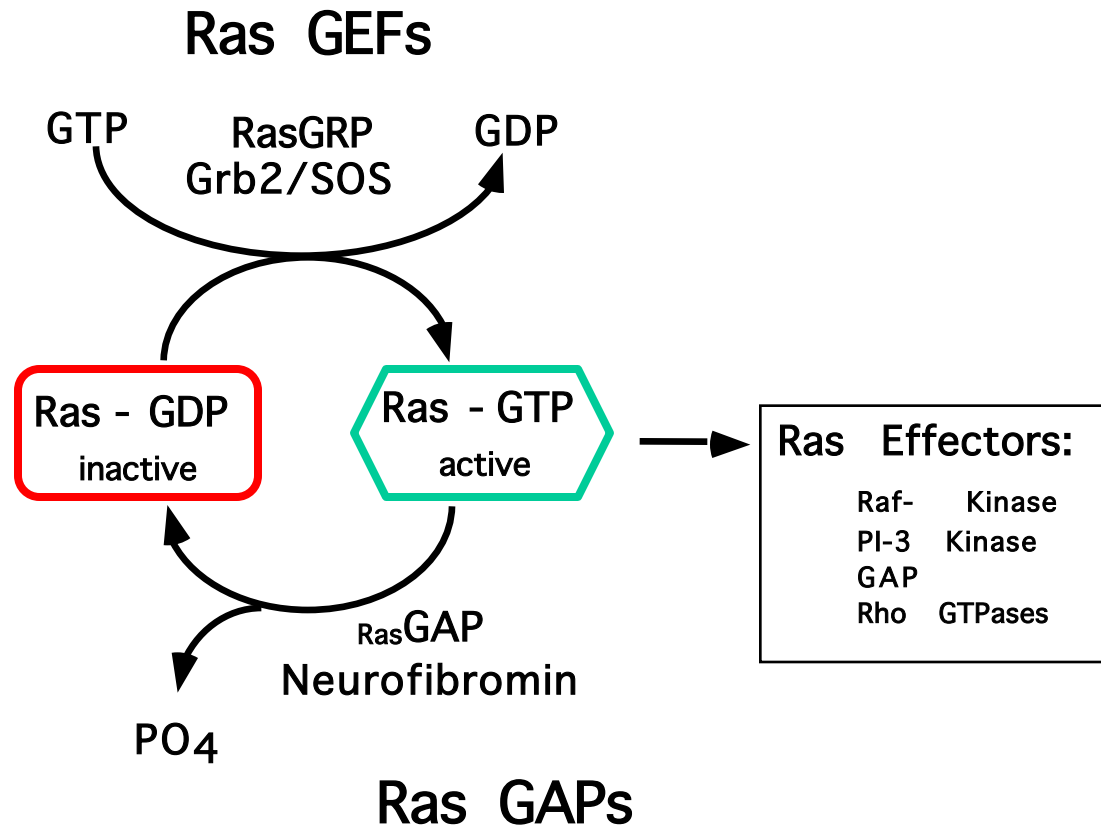
NFkB activation

Activation of RasGRP

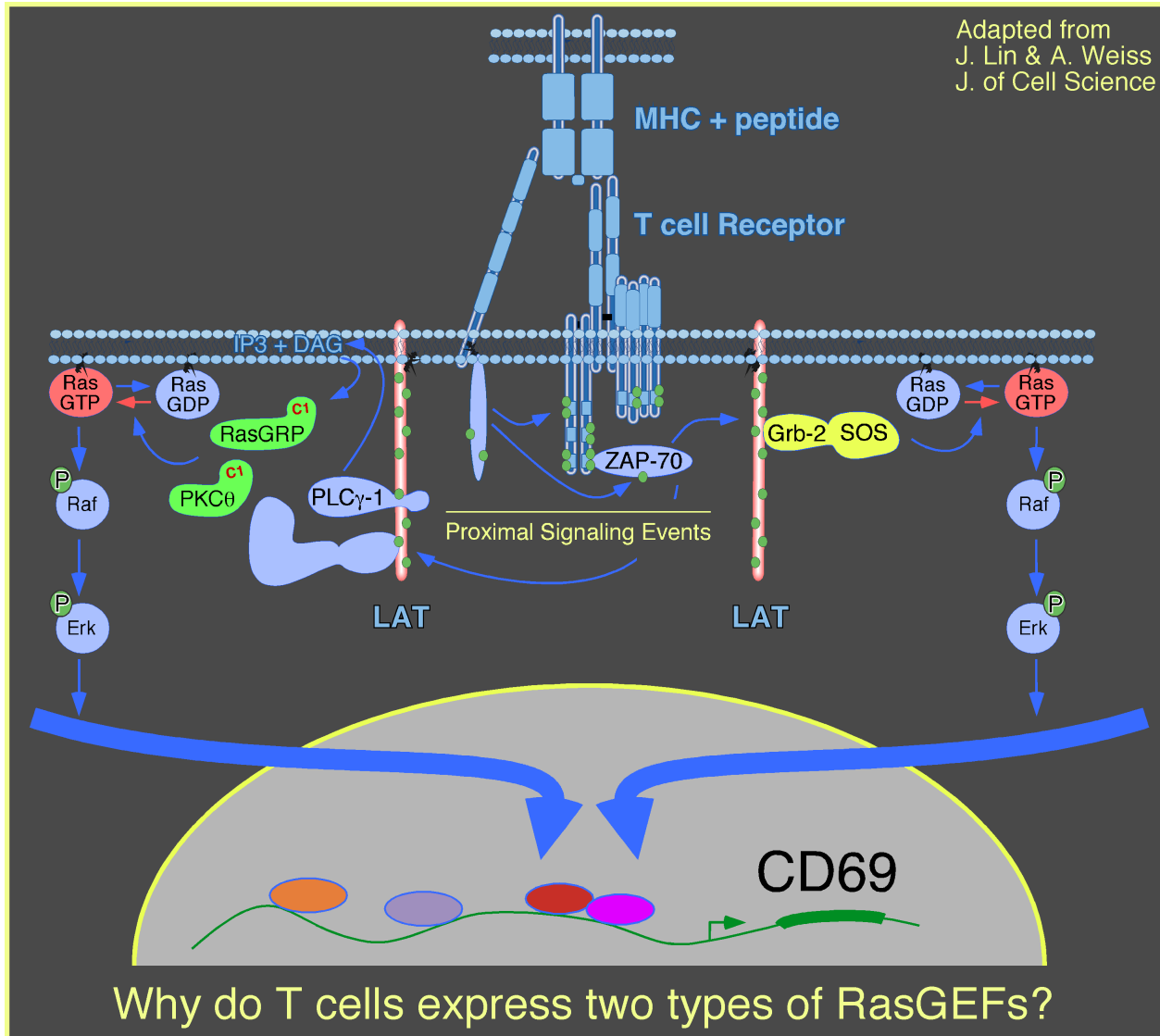
The diagram illustrates the TCR signaling pathway. The TCR (blue) is associated with the LcK (yellow) and ZAP-70 (yellow) kinases. The LAT (green) protein is phosphorylated (P) and recruits PLCγ (orange) and SLP76 (orange). SLP76 recruits Vav (orange), which activates PKCθ (pink) and PDK1 (pink). PI3K (pink) is also activated and phosphorylates PDK1. PKCθ and PDK1 phosphorylate CARMA1 (blue), BCL10 (blue), MALT1 (blue), and TRAF6 (green). This complex recruits IKKα (purple), IKKβ (purple), and NEMO (purple). IKKα and IKKβ are ubiquitinated (orange) and phosphorylated (P). The phosphorylated IKK complex then phosphorylates IκB (yellow), leading to the release of NF-κB (yellow) for nuclear translocation. A legend indicates that 'P' represents phosphorylation and orange ovals represent ubiquitination.

van Oers and Chen, Science, 2005

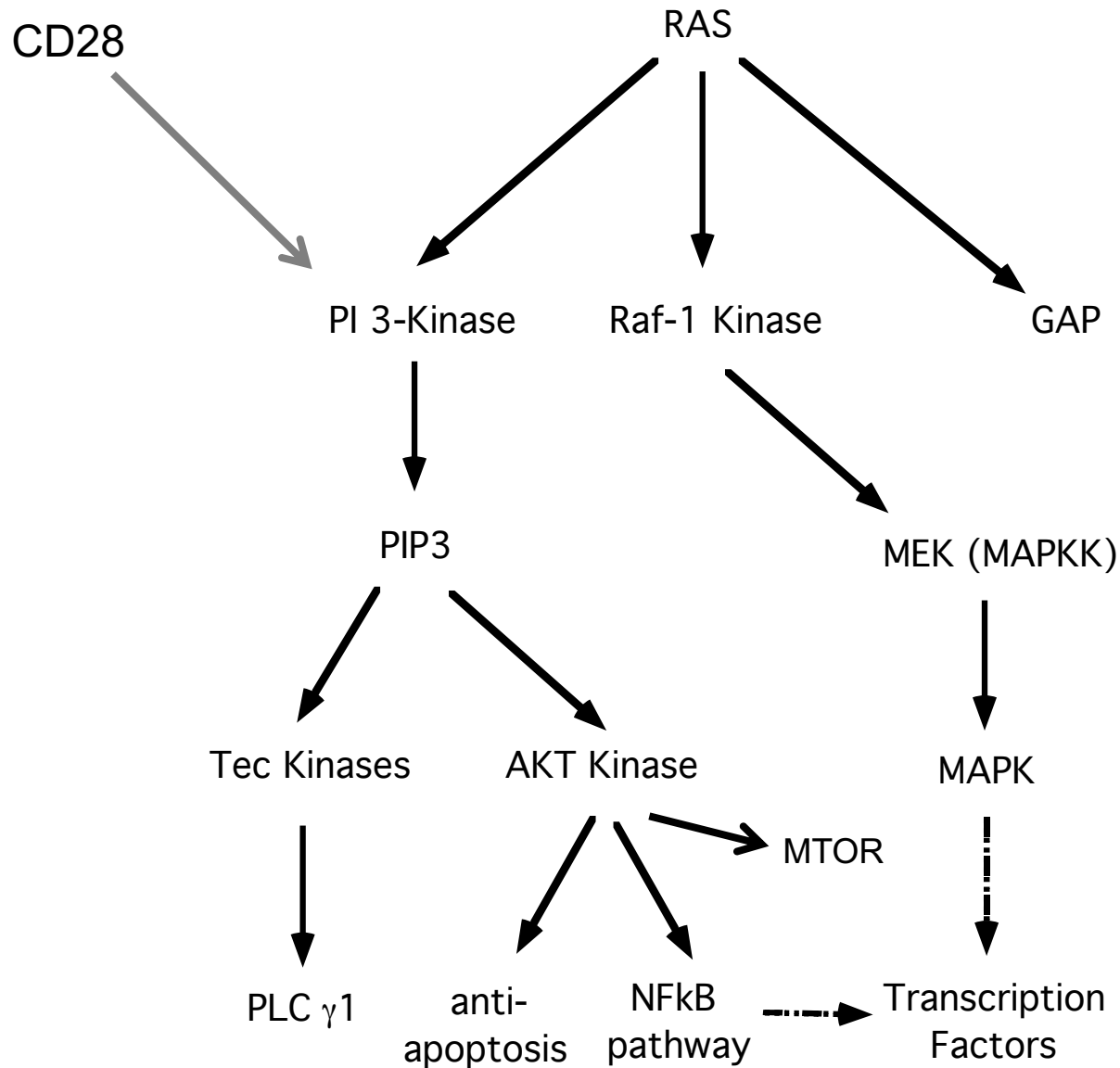
TCR Stimulation Activates Ras Via Guanine Nucleotide Exchange Factors (GEFs)



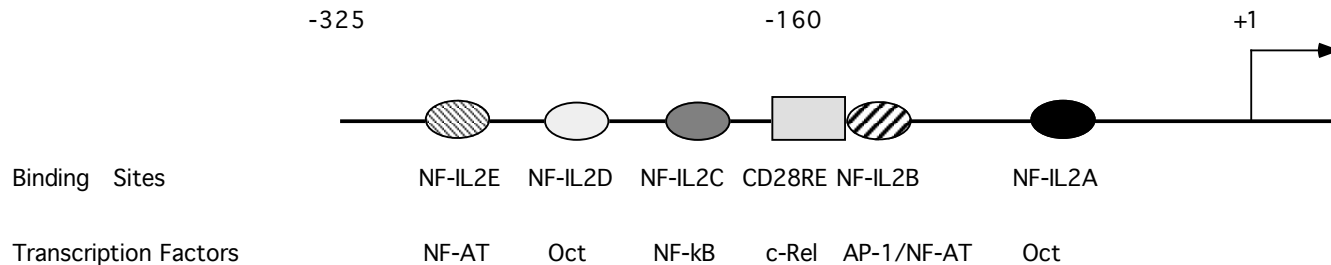
The TCR Uses Two GEFs to Activate Ras



Ras Activates Multiple Signaling Pathways



IL-2 Gene Regulation



Transcriptional Regulation

IL-2 gene is inactive in naive T cells

Site of for signal integration (TCR and costimulatory signals)

Transcriptional Activation

Cascade of Events:

Immediate-early genes are involved (i.e., Jun and Fos)

- delayed temporal expression of IL-2 transcripts
- requirement for protein synthesis

Nuclear translocation of preformed-inactive factor (NFAT)

- requires calcium flux
- requires calcineurin activation
- dephosphorylation of inactive cytoplasmic NFAT

Post-transcriptional Regulation:

mRNA stabilization via AU-rich 3' untranslated region

Activation of Transcription Factors in T Cells

Phosphorylation, release, and degradation of I κ B

Dephosphorylation of cytoplasmic NFAT

MAP kinase, SAP kinase pathways

