

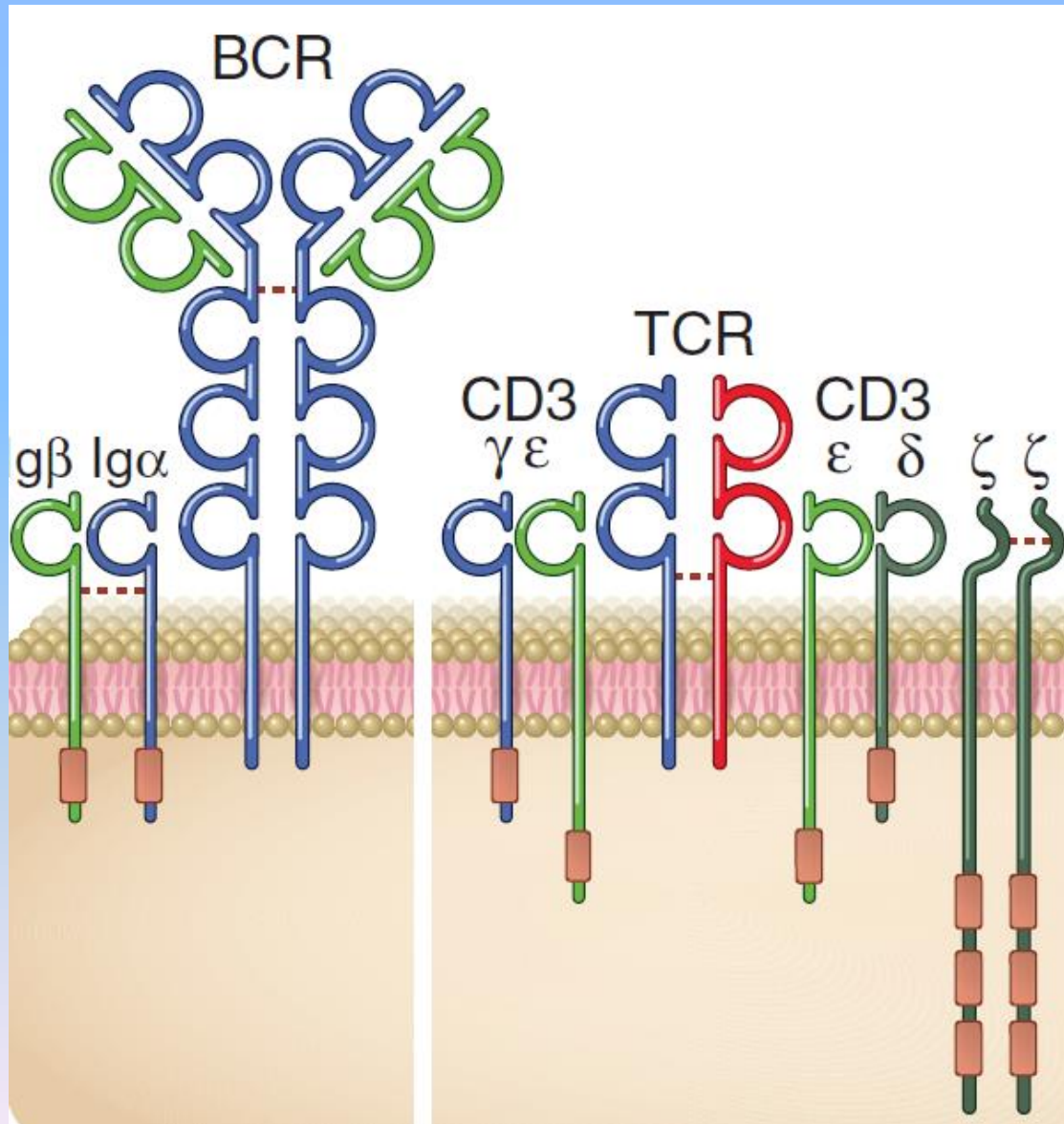
Graduate Program in Immunology
BMI5905 - Effector Mechanisms of Immune Response

Antigen Presentation and Peripheral Tolerance

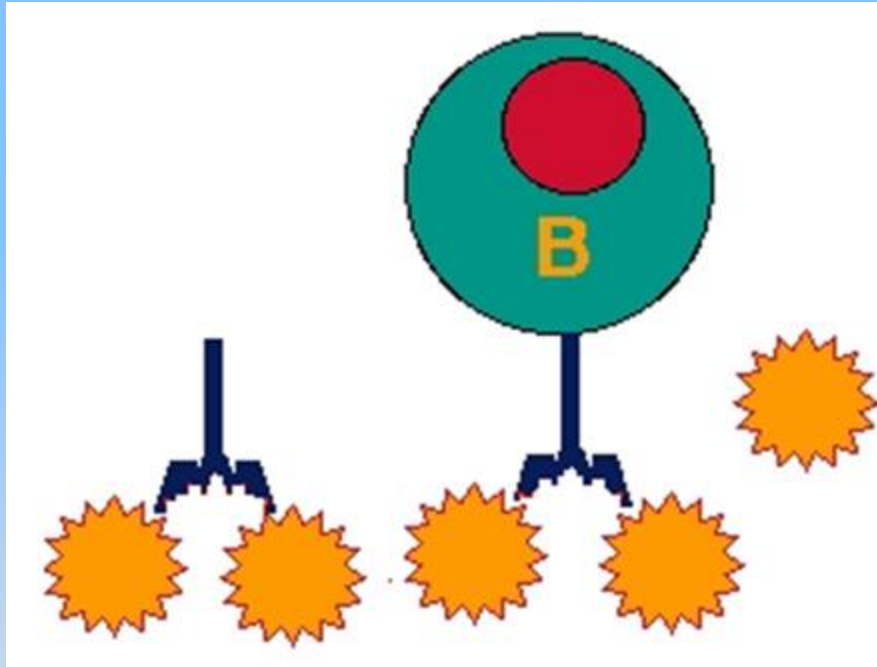
Prof. Anderson Sá-Nunes

Department of Immunology
Institute of Biomedical Sciences
University of Sao Paulo

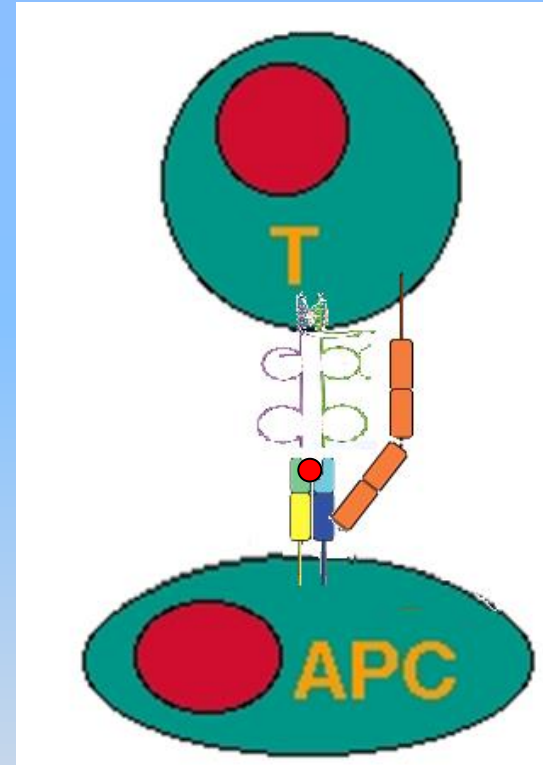
Antigen Receptors of Acquired Immunity



Antigen Recognition by Lymphocyte Receptors

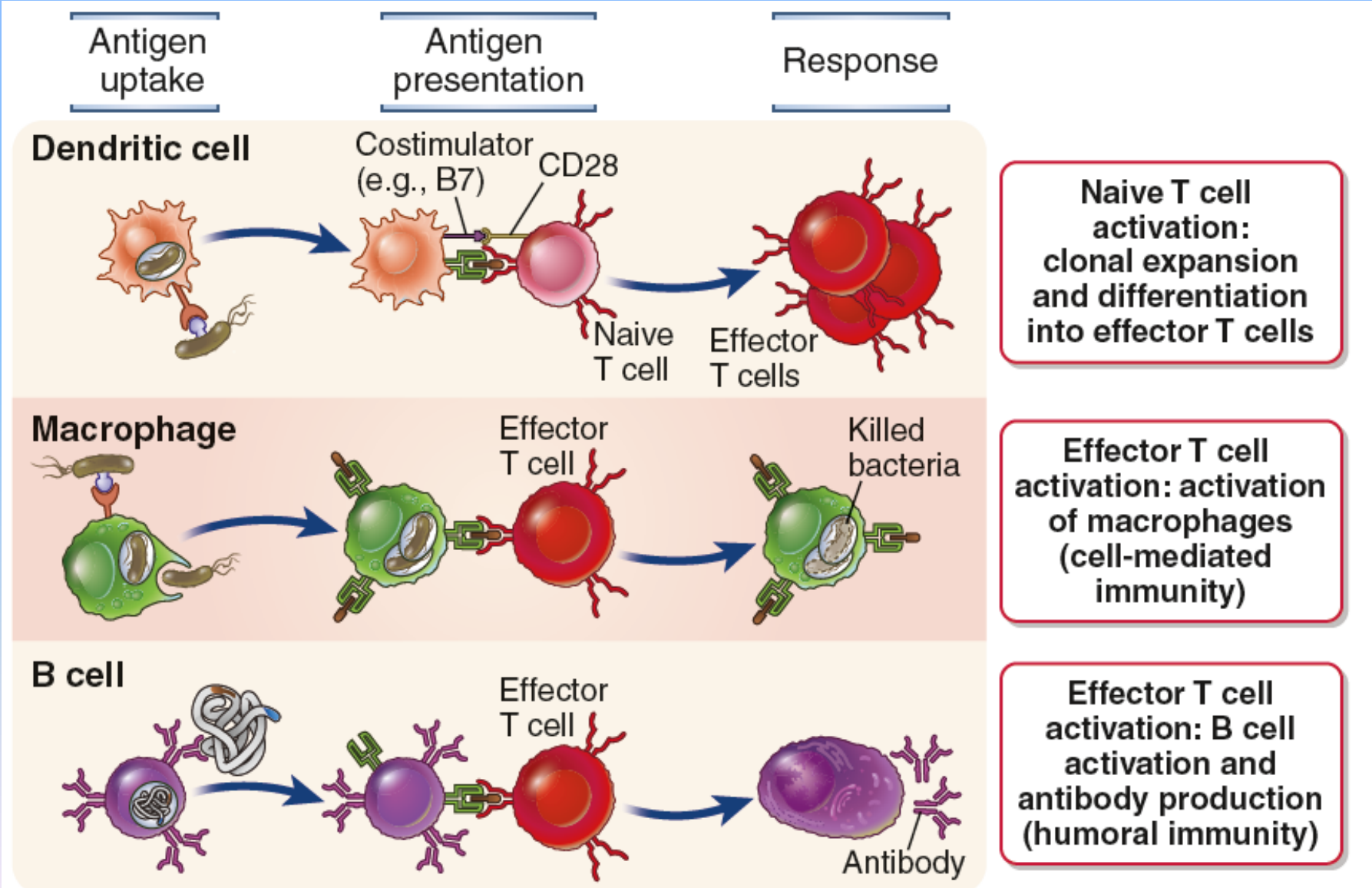


BCRs recognize antigens on both native or denatured conformation

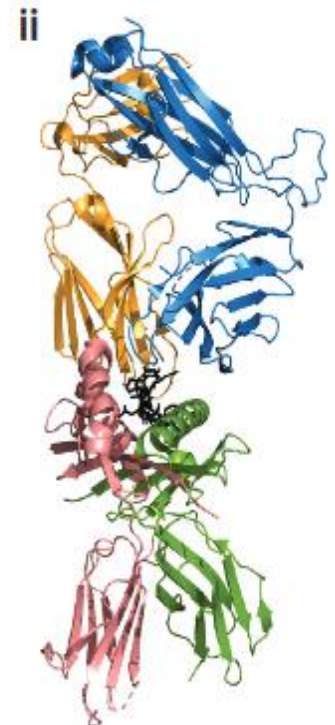
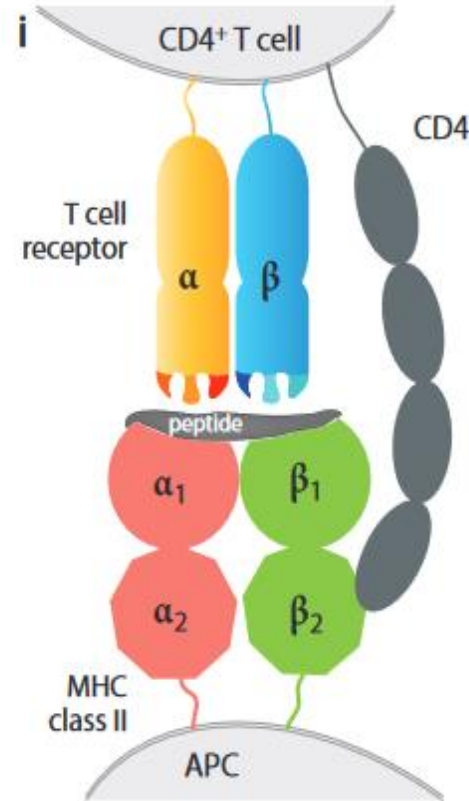
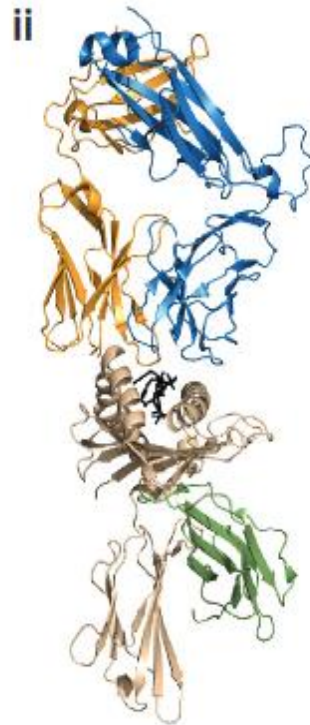
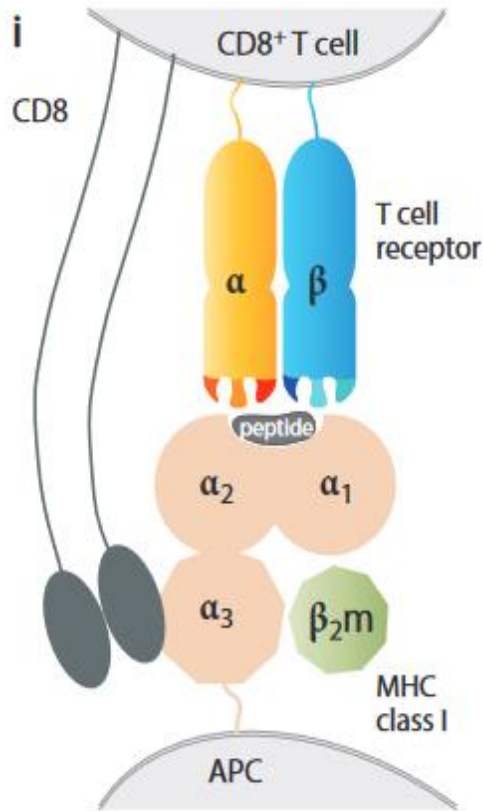


TCRs do not recognize antigens directly. Small parts of protein antigens (peptides) need to be bound to MHC molecules and displayed by antigen presenting cells (APC)

Antigen Presenting Cells



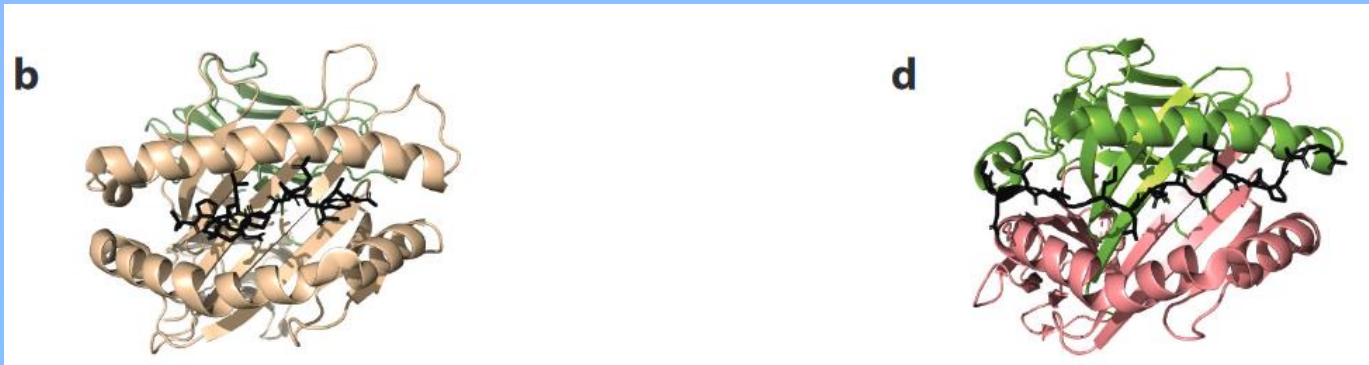
MHC-peptide/TCR interactions



HLA-A
HLA-B
HLA-C

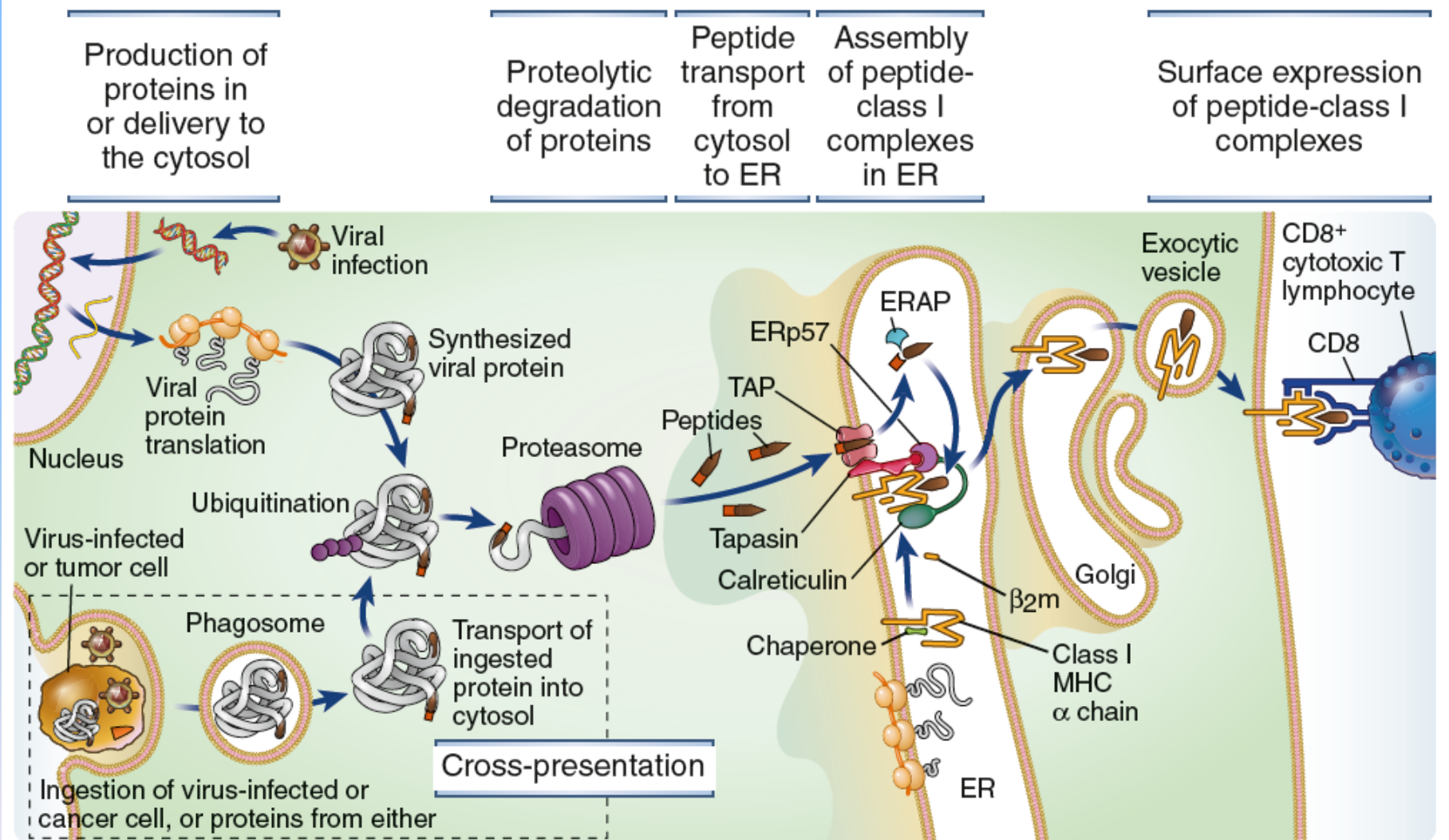
HLA-DP
HLA-DQ
HLA-DR

Peptide binding to MHC molecules and allotypes

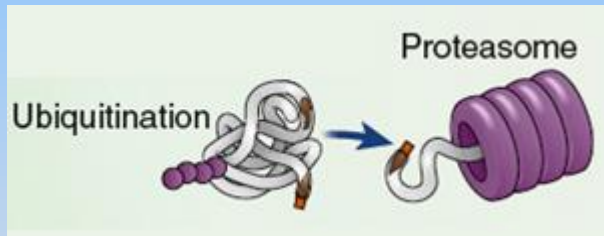


HLA allotype	Number of HLA allotypes
HLA-A	3,629
HLA-B	4,572
HLA-C	3,447
HLA-E	12
HLA-F	6
HLA-G	19
HLA class I	11,685
HLA-DP	3,160
HLA-DQ	1,278
HLA-DR	1,055
HLA-DM	11
HLA-DO	8
HLA class II	5,512

MHC Class I Pathway



Ubiquitin/Proteasome System



Ubiquitin

- Identified in 1974, sequenced one year later
- Present in all living organisms
- 76 aminoacids (8.5 kDa)
- Extremely conserved:
Human X Yeast = 96% identity

BIOCHEMISTRY, VOL. 14, NO. 10, 2214-2218, 1975

The Complete Amino Acid Sequence of Ubiquitin, an Adenylate Cyclase Stimulating Polypeptide Probably Universal in Living Cells[†]

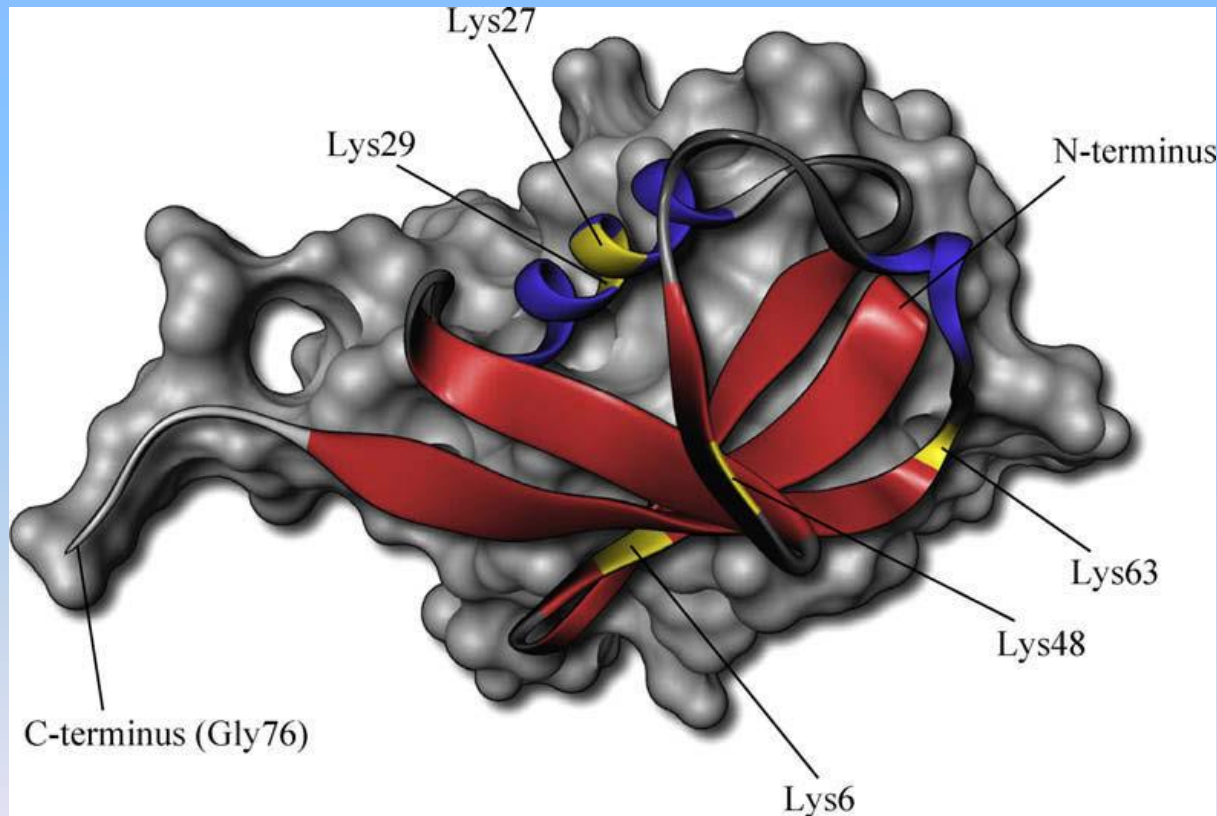
David H. Schlesinger,* Gideon Goldstein, and Hugh D. Niall[‡]

MQIFV**K**TLTG**K**TITLEVEPSDTIENV**KAK**IQD**K**EGIPPDQQRLIFAG**K**QLEDGRTLSDYNIQ**K**ESTLHLVLRRLGG

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MQIFV**K**TLTG**K**TITLEVEPSDTIENV**KAK**IQD**K**EGIPPDQQRLLIFAG**K**QLEDGRTLSDYNIQ**K**ESTLHLVLRRLRGG

Elucidation of basic functions of ubiquitin and protein ubiquitination pathways



The Nobel Prize in Chemistry 2004
Aaron Ciechanover, Avram Hershko, Irwin Rose

The Nobel Prize in Chemistry 2004 was awarded jointly to Aaron Ciechanover, Avram Hershko and Irwin Rose "for the discovery of ubiquitin-mediated protein degradation".

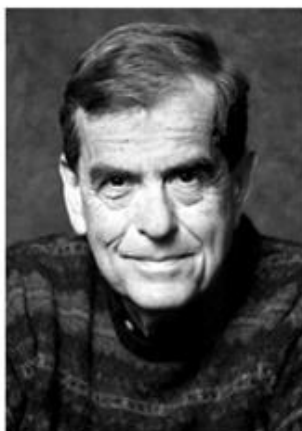


Photo: D. Porges

Aaron Ciechanover

Aaron Ciechanover

Born: 1 October 1947, Haifa, British Protectorate of Palestine (now Israel)

Affiliation at the time of the award:
Technion – Israel Institute of Technology, Haifa, Israel

Prize motivation: "for the discovery of ubiquitin-mediated protein degradation"

Field: Biochemistry



Photo: D. Porges

Avram Hershko

Avram Hershko

Born: 31 December 1937, Karcag, Hungary

Affiliation at the time of the award:
Technion – Israel Institute of Technology, Haifa, Israel

Prize motivation: "for the discovery of ubiquitin-mediated protein degradation"

Field: Biochemistry



Irwin Rose

Irwin Rose

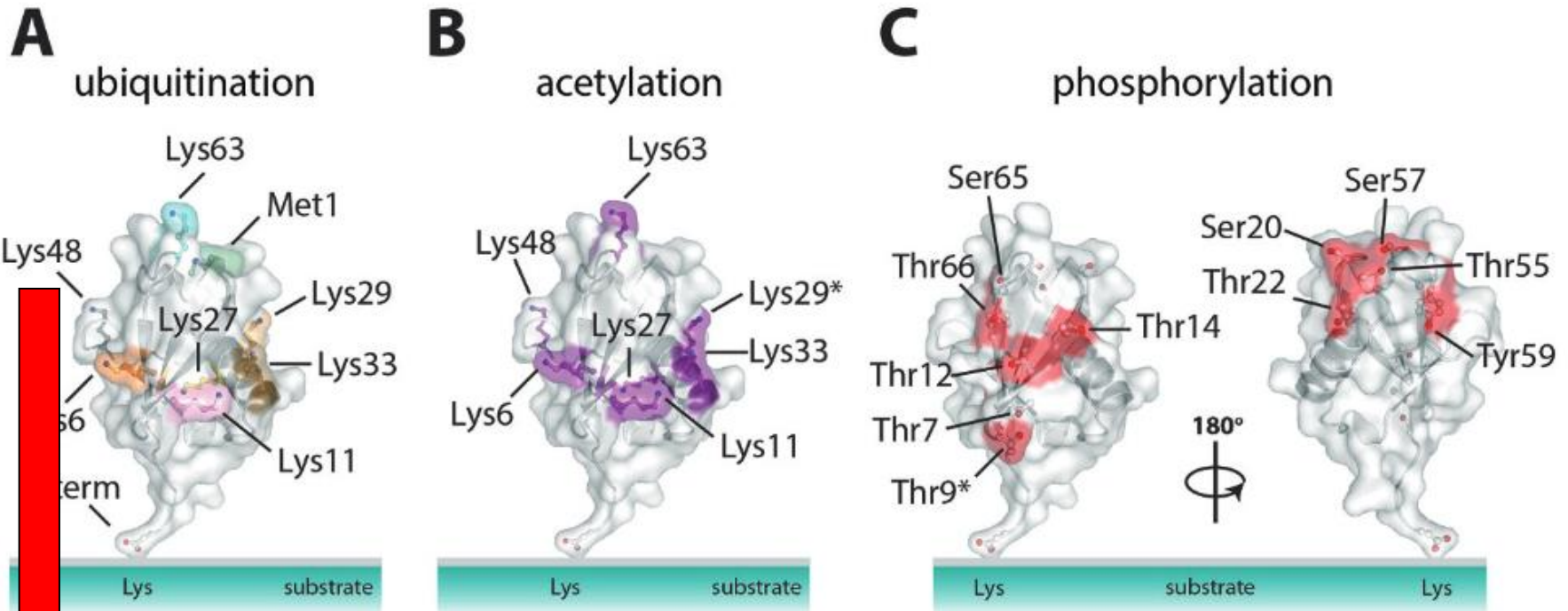
Born: 16 July 1926, Brooklyn, NY, USA

Affiliation at the time of the award:
University of California, Irvine, CA, USA

Prize motivation: "for the discovery of ubiquitin-mediated protein degradation"

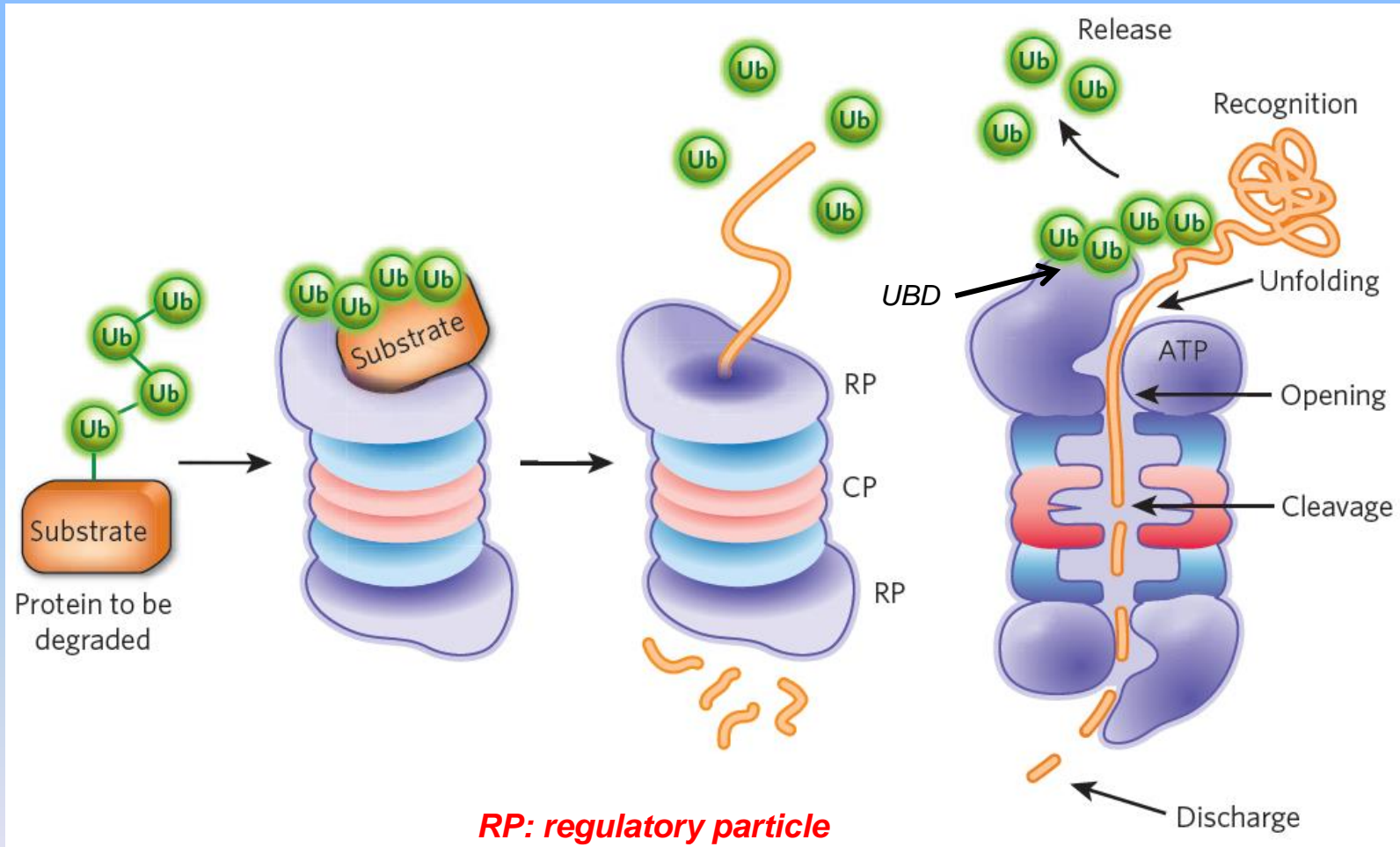
Field: Biochemistry

Ubiquitin Modifications



***~50% of all modifications
Target proteins to the proteasome***

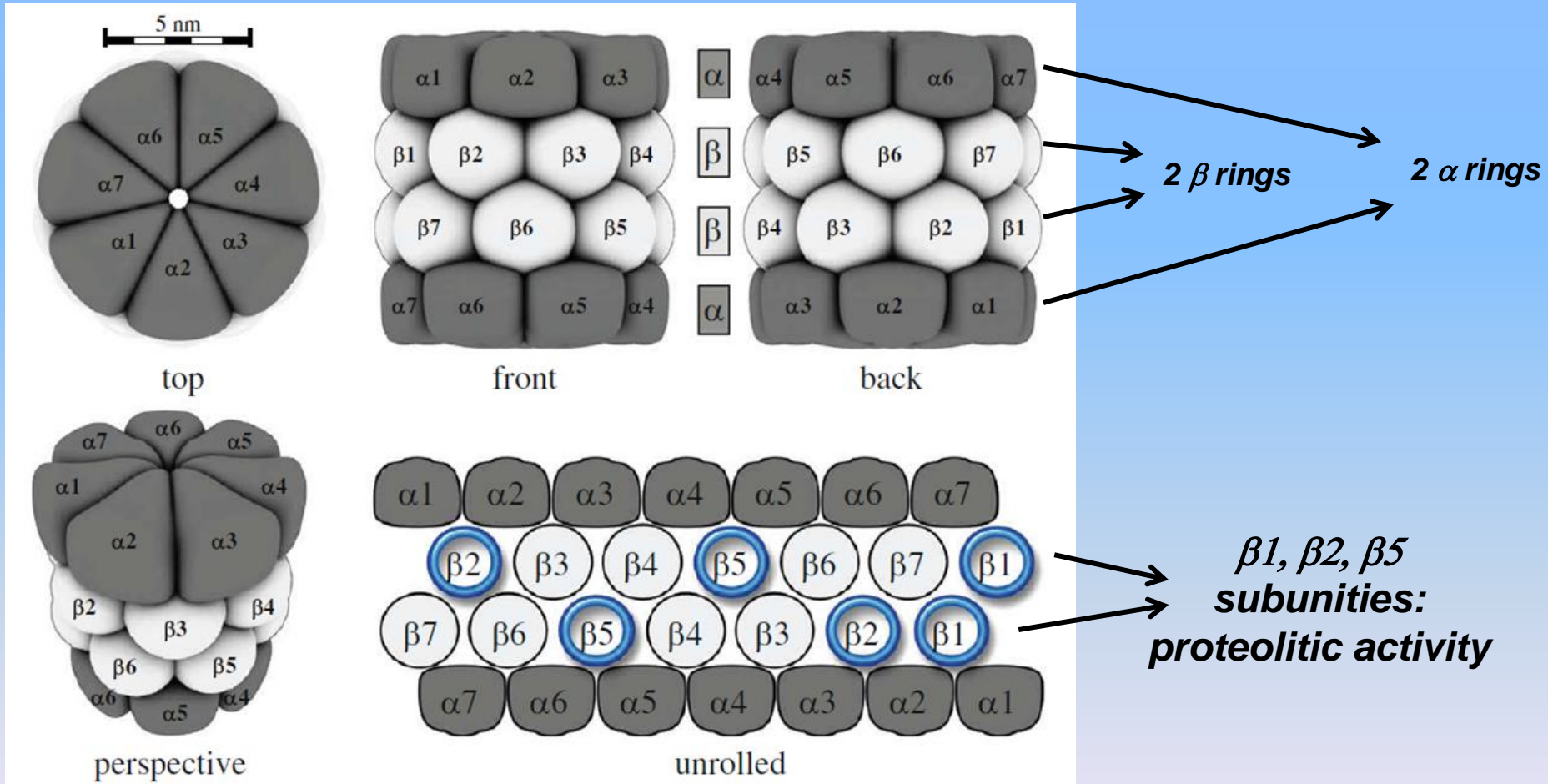
Lys48 Poliubiquitination: Proteasome Degradation



RP: regulatory particle
CP: core particle (20S)
UBD: ubiquitin-binding domain

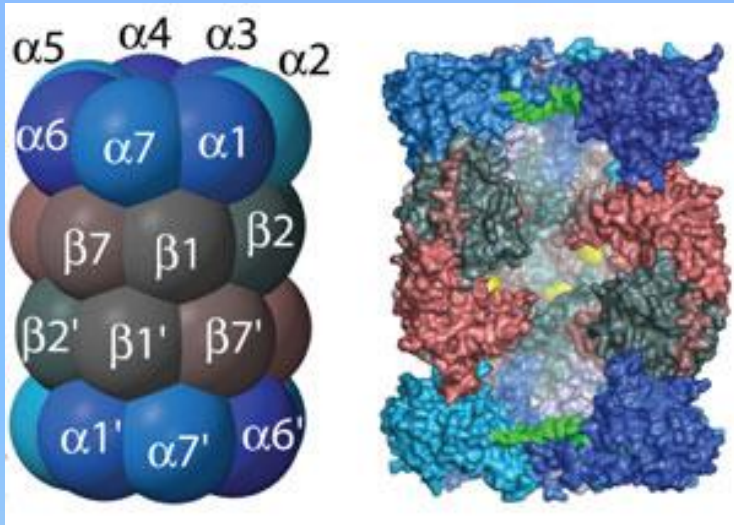
Proteasome 20S (20S Core)

Present in all eukariotic and some prokariotic organisms



Types of Proteasomes

<http://picsdigger.com/image/6d0c4b8c/>



Cylindrin

Macroxyproteinase

Multicatalytic proteinase complex

Prosome

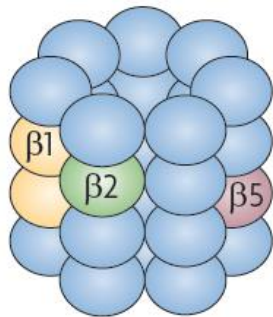
2-35 aa peptides

Peaks of:

2-3 aa

8-10 aa

20-30 aa



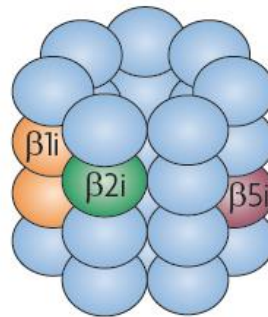
Constitutive proteasome

$\beta 1$ (PSMB6, Y, δ)

$\beta 2$ (PSMB7, Z, MC14)

$\beta 5$ (PSMB5, X, MB1, ϵ)

Optimized for 8-10 aa



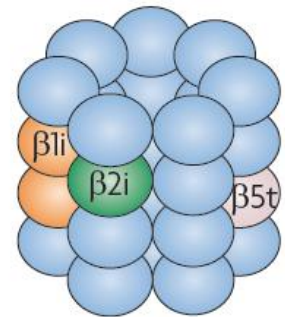
Immunoproteasome

$\beta 1i$ (PSMB9, LMP2)

$\beta 2i$ (PSMB10, LMP10, MECL1)

$\beta 5i$ (PSMB8, LMP7)

$\beta 5t$ KO: less CD8+ T cells



Thymoproteasome

$\beta 1i$ (PSMB9, LMP2)

$\beta 2i$ (PSMB10, LMP10, MECL1)

$\beta 5t$ (PSMB11)

Proteasome Regulators

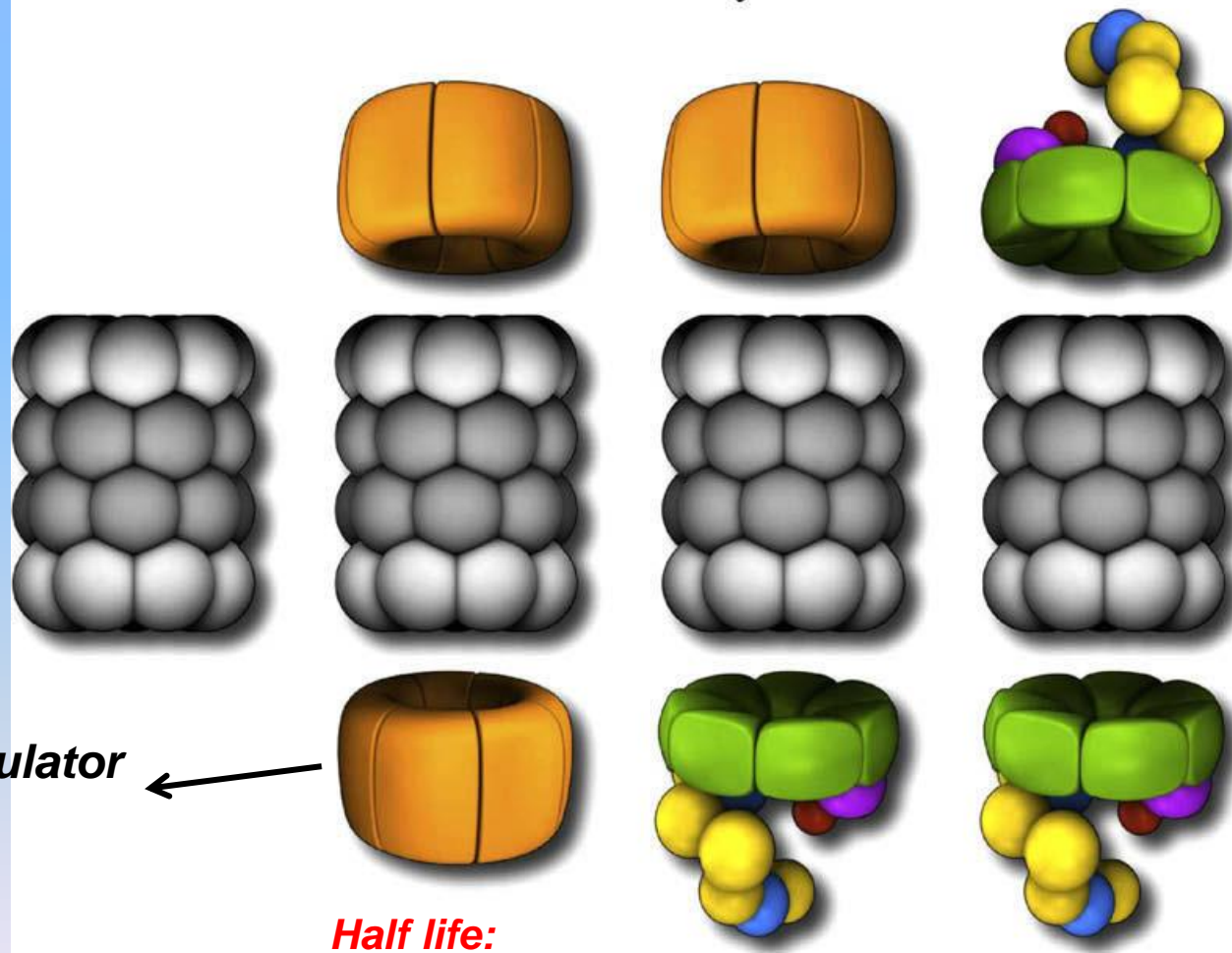
HeLa cells 41±5%
20S

20±9%
immuno

24±9%
hybrid

15±3%
26S

MW: 2 MDa



**Ubiquitin
binding
region**

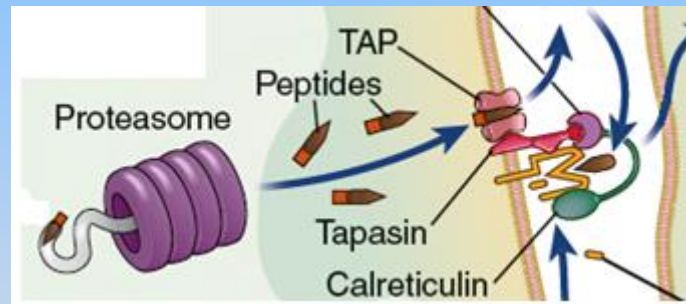
**11S regulator
(PA28)**

**19S
regulator**

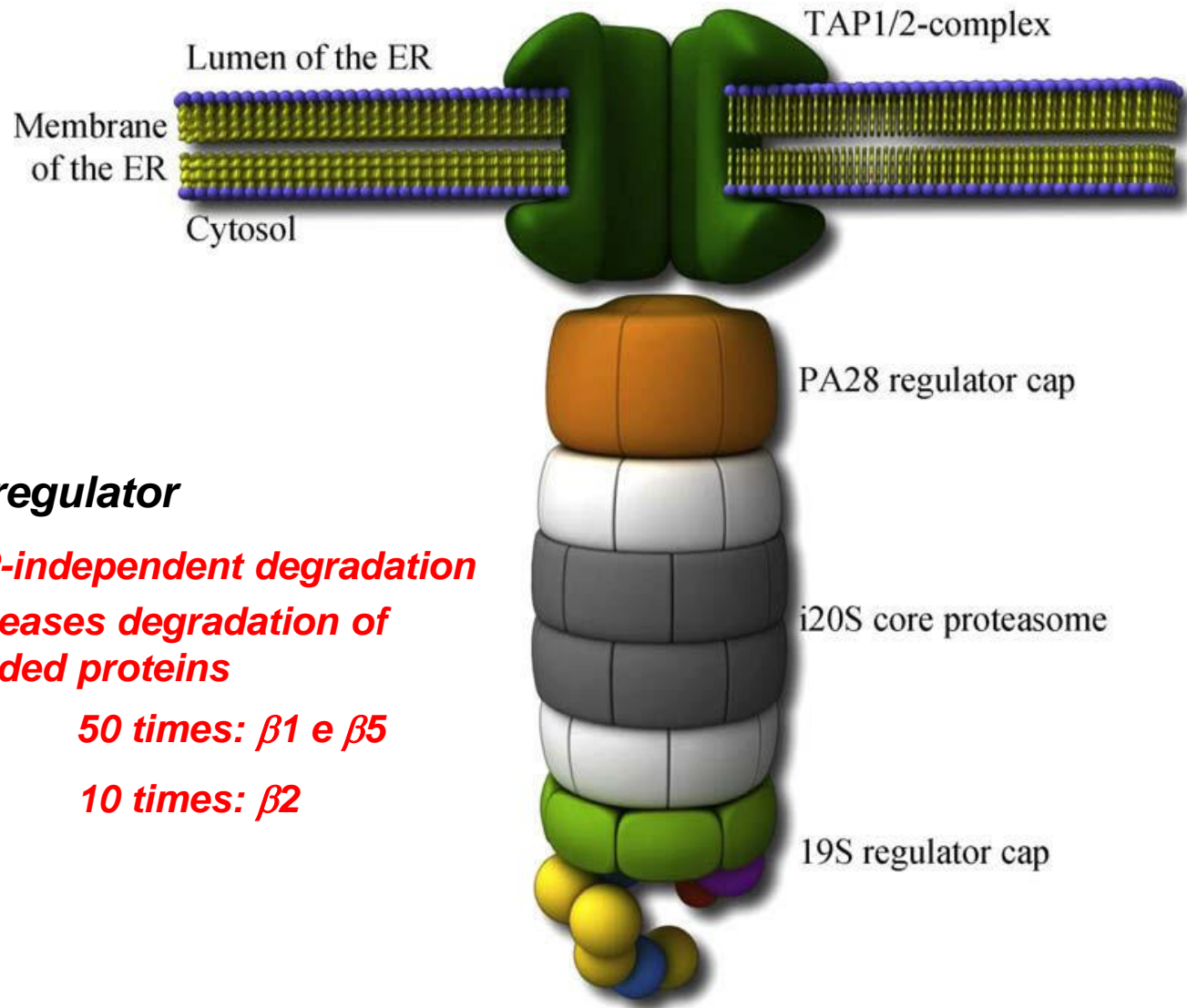
**Half life:
27 hours**

**Half life:
8-12 days**

MHC Class I Pathway



Molecular Coupling Proteasome-TAP 1/2



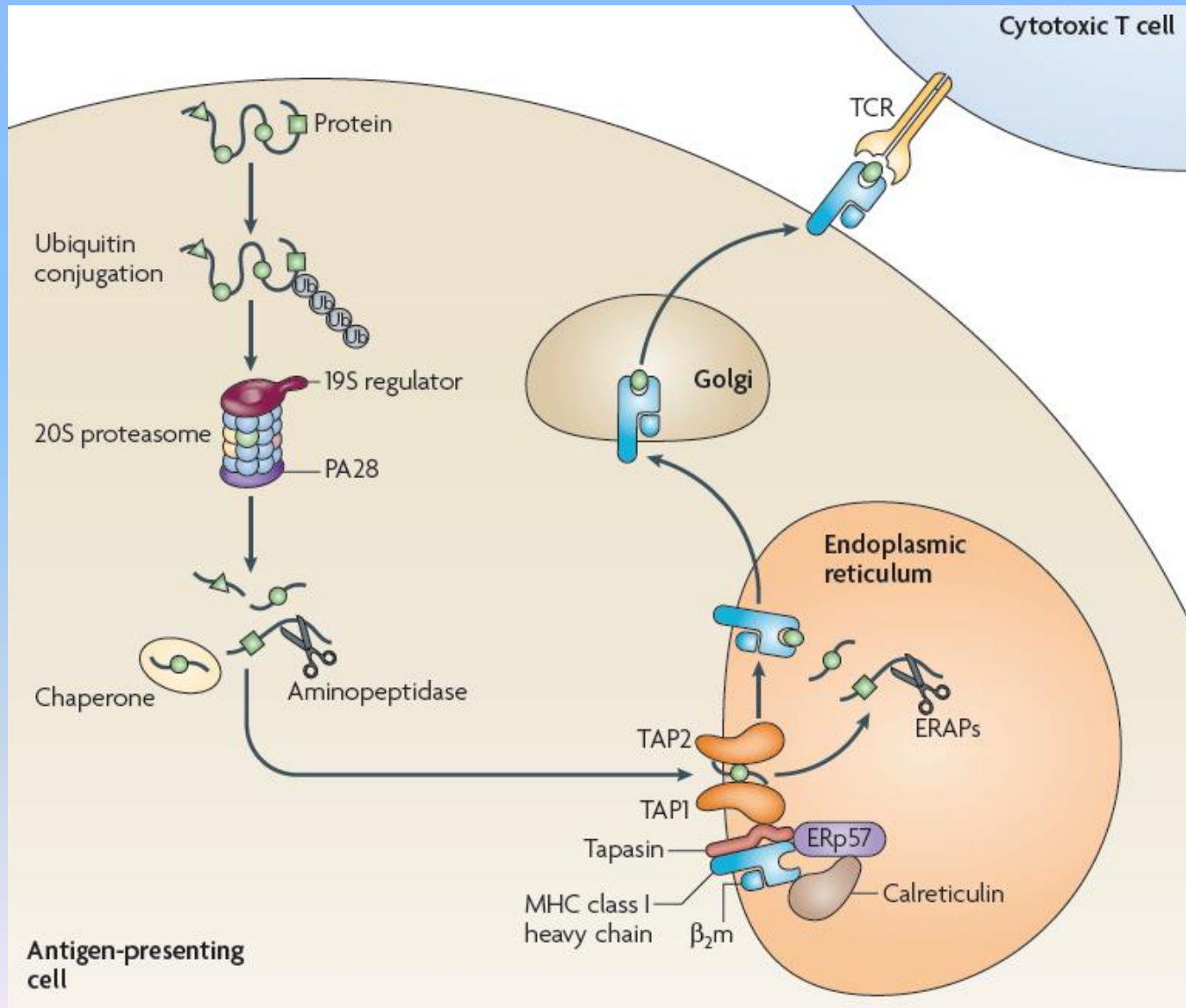
11S regulator

- **ATP-independent degradation**
- **Increases degradation of unfolded proteins**

50 times: $\beta 1$ e $\beta 5$

10 times: $\beta 2$

Processing of "Endogenous" Peptides



MHC Class II Pathway

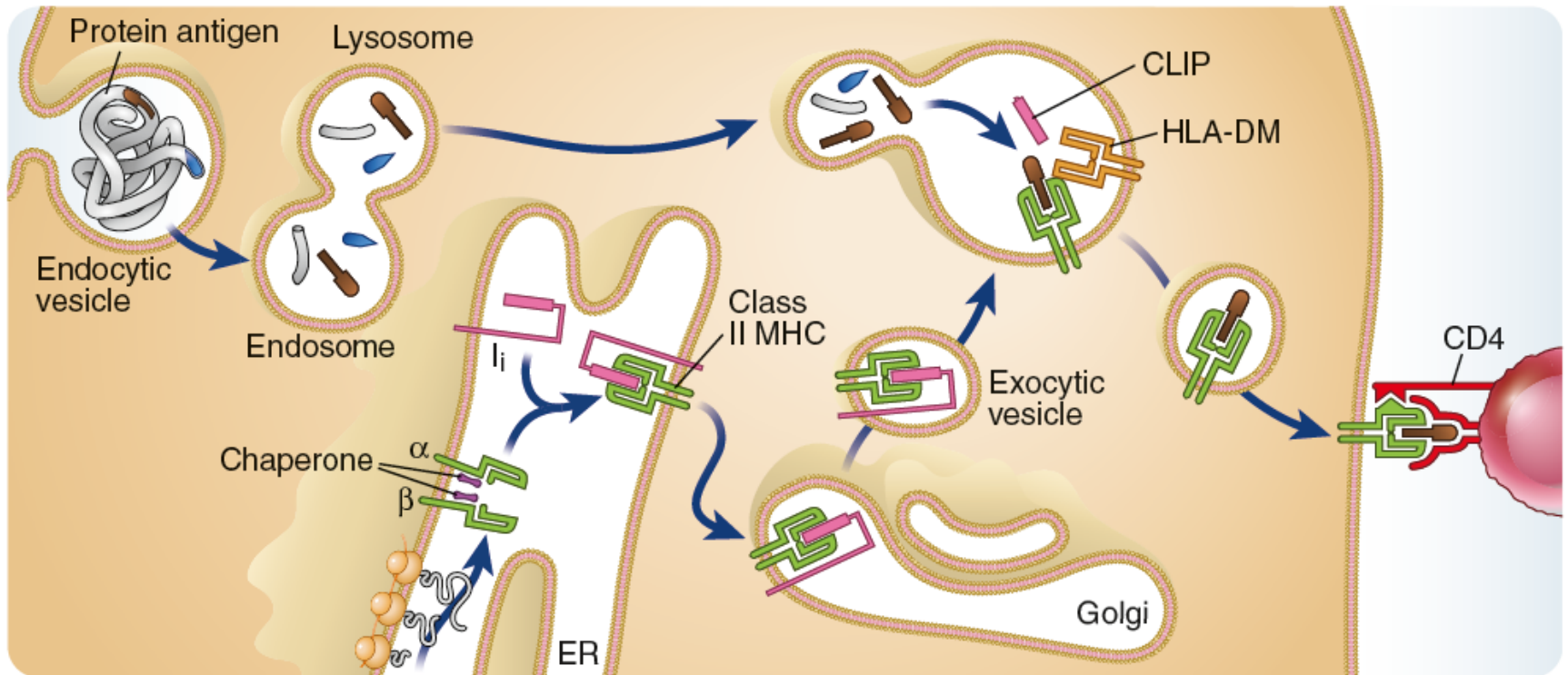
Uptake of extracellular proteins into vesicular compartments of APC

Processing of internalized proteins in endosomal/lysosomal vesicles

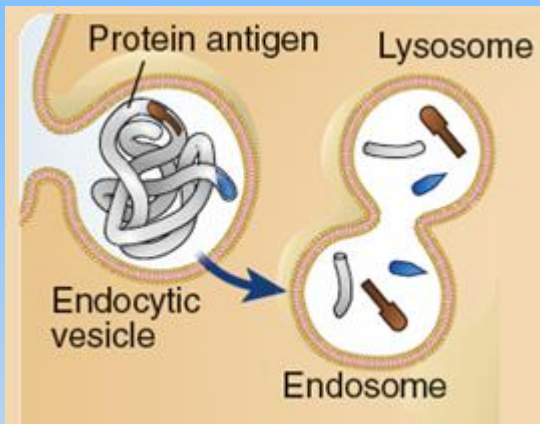
Biosynthesis and transport of class II MHC molecules to endosomes

Association of processed peptides with class II MHC molecules in lysosomes

Expression of peptide-MHC complexes on cell surface



MHC Class II Pathway



History of Lysosomal Proteases

Lysosomotropic Agents (Biochem Pharmacol 23:2495-2531, 1974)

Proc. Natl. Acad. Sci. USA
Vol. 79, pp. 175-178, January 1982
Immunology

Decrease in macrophage antigen catabolism caused by ammonia and chloroquine is associated with inhibition of antigen presentation to T cells

(macrophage-lymphocyte interactions/T-cell immunity/lysosomotropic agents/antibacterial immunity)

H. KIRK ZIEGLER* AND EMIL R. UNANUE†

Table 1. Inhibition of antigen presentation with NH₄Cl and chloroquine

Assay	Control, %	10 mM NH ₄ Cl		0.1 mM chloroquine	
		Observed, %	Δ,%	Observed, %	Δ,%
Antigen uptake*	15 ± 1	13 ± 2	13	15 ± 2	0
Antigen ingestion†	66 ± 2	63 ± 2	5	67 ± 6	-2
Antigen catabolism‡	29 ± 4	13 ± 3	55	14 ± 6	52
I-A expression§	54 ± 4	59 ± 2	-9	56 ± 4	-4
T cell-macrophage binding¶					
Before antigen handling	70 ± 7	26 ± 8	63	30 ± 8	57
After antigen handling	84 ± 8	70 ± 11	17	64 ± 10	24

Peritoneal macrophages
¹²⁵I-Listeria monocytogenes

Drug exposure (1 h):

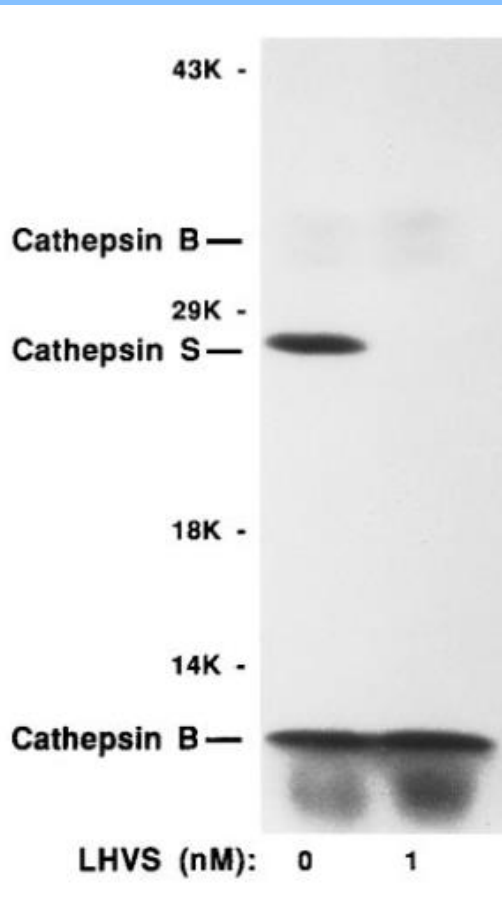
- Before handling: starting 30 min before Listeria uptake
- After handling: 30 min after Listeria uptake

Cathepsin S: essential to antigen presentation

Immunity, 1996, 4: 357-366.

Essential Role for Cathepsin S in MHC Class II-Associated Invariant Chain Processing and Peptide Loading

Richard J. Riese,* Paula R. Wolf,† Dieter Brömme,‡ Lisa R. Natkin,* José A. Villadangos,† Hidde L. Ploegh,† and Harold A. Chapman*



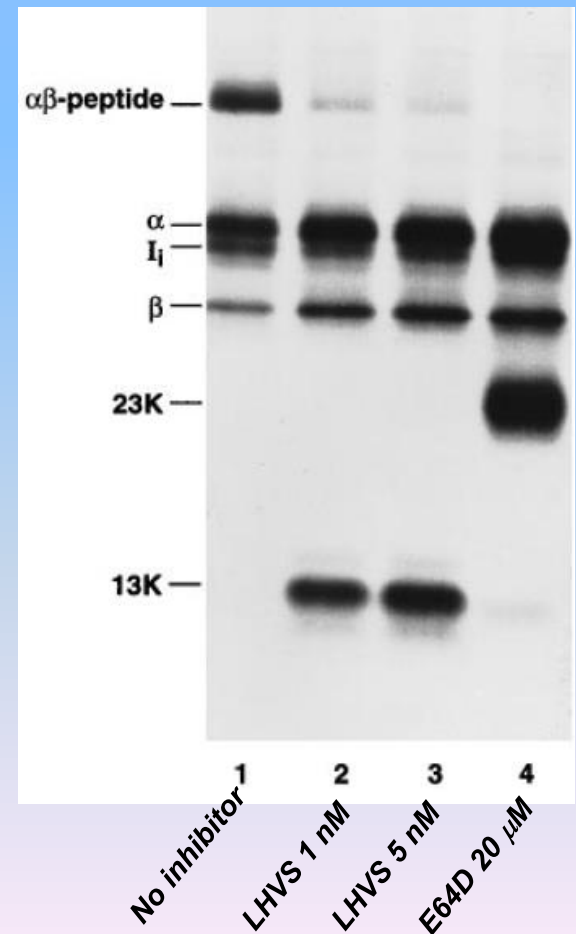
³⁵[S]-methionine/cysteine

Treatments:

- LHVS (Cathepsin S inhibitor)

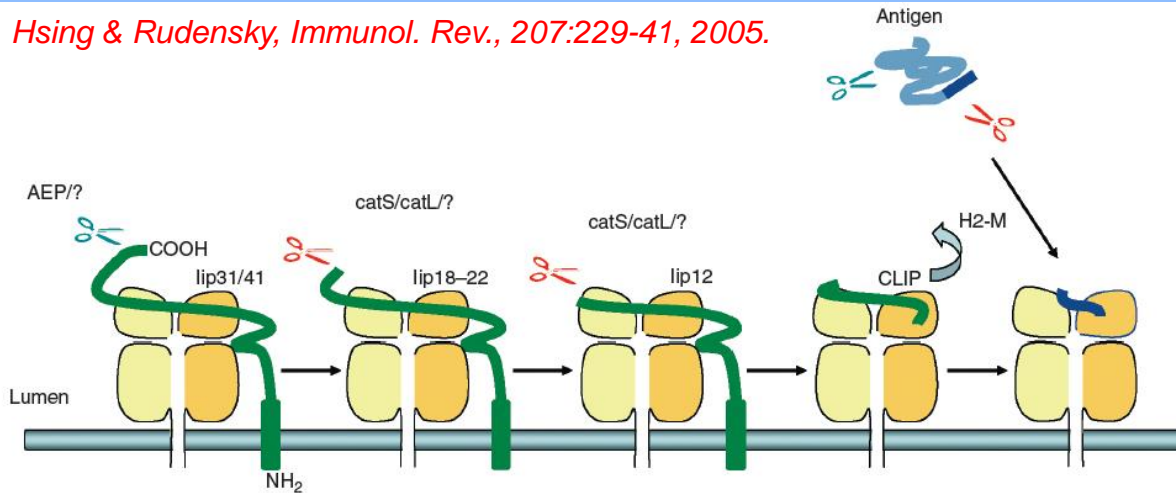
- E64D (broad spectrum cystein protease inhibitor)

Immunoprecipitation:
anti-HLA-DR1





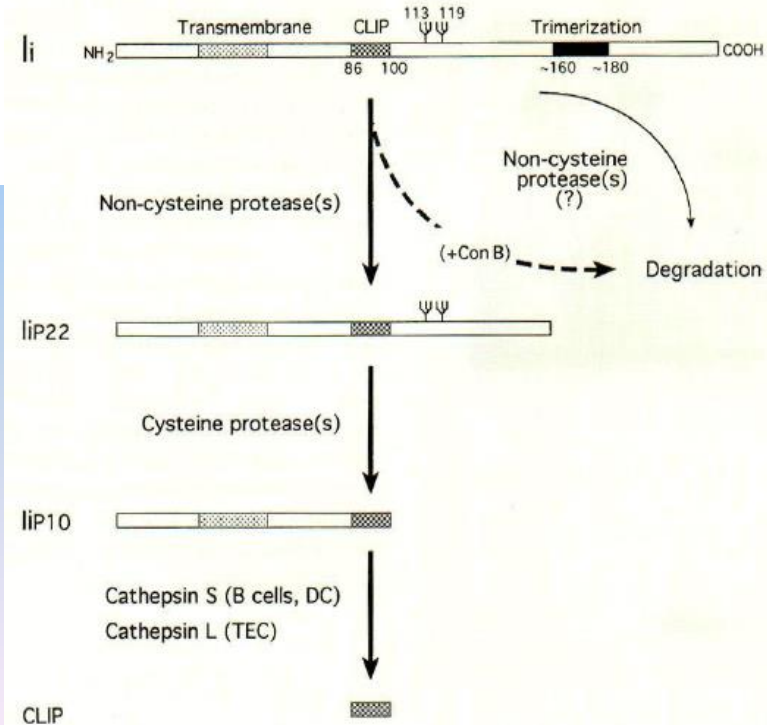
Invariant Chain Degradation

Hsing & Rudensky, *Immunol. Rev.*, 207:229-41, 2005.



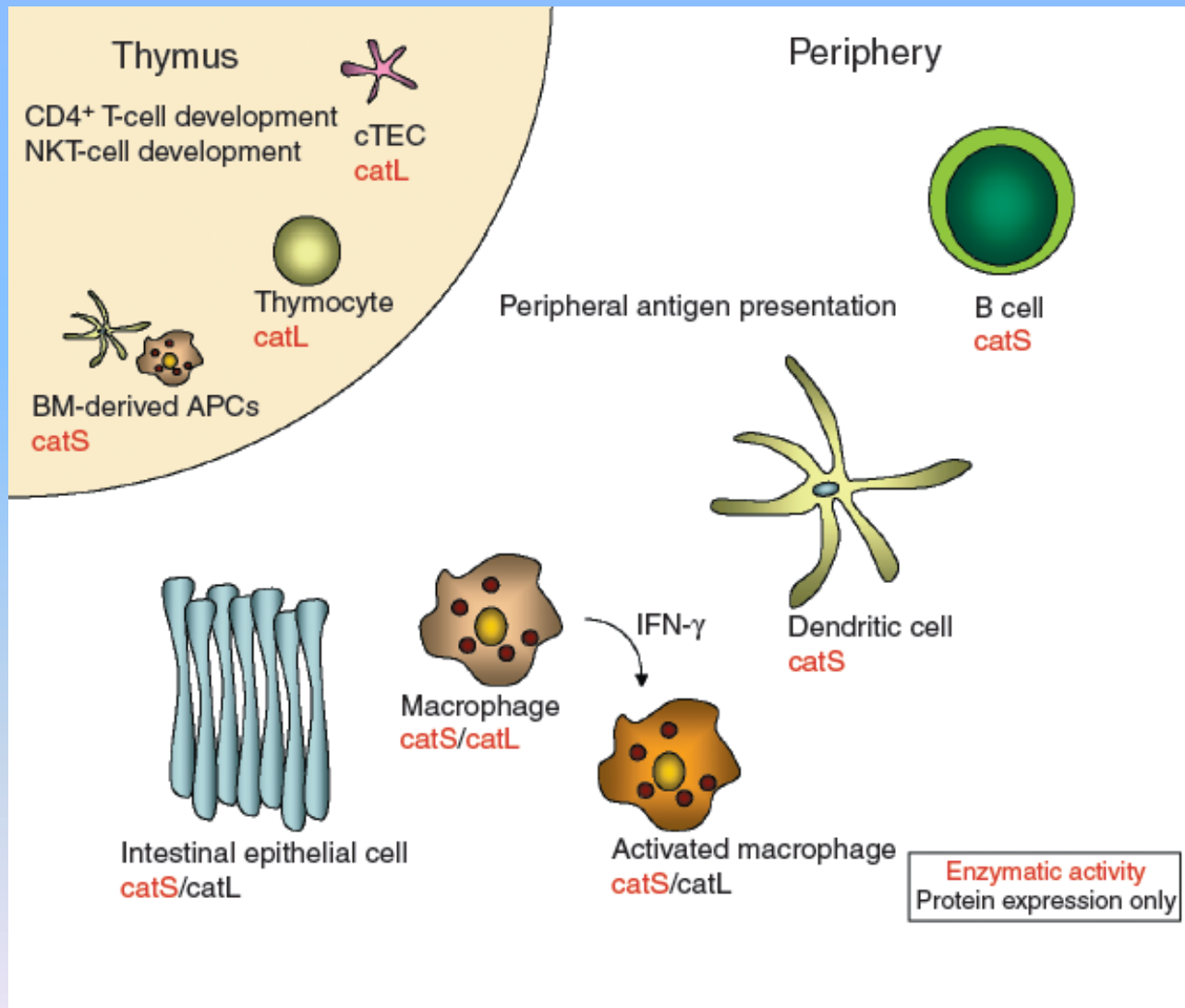
Cytosol

-  Leupeptin-insensitive cysteine or aspartic proteases
-  Leupeptin-sensitive cysteine proteases

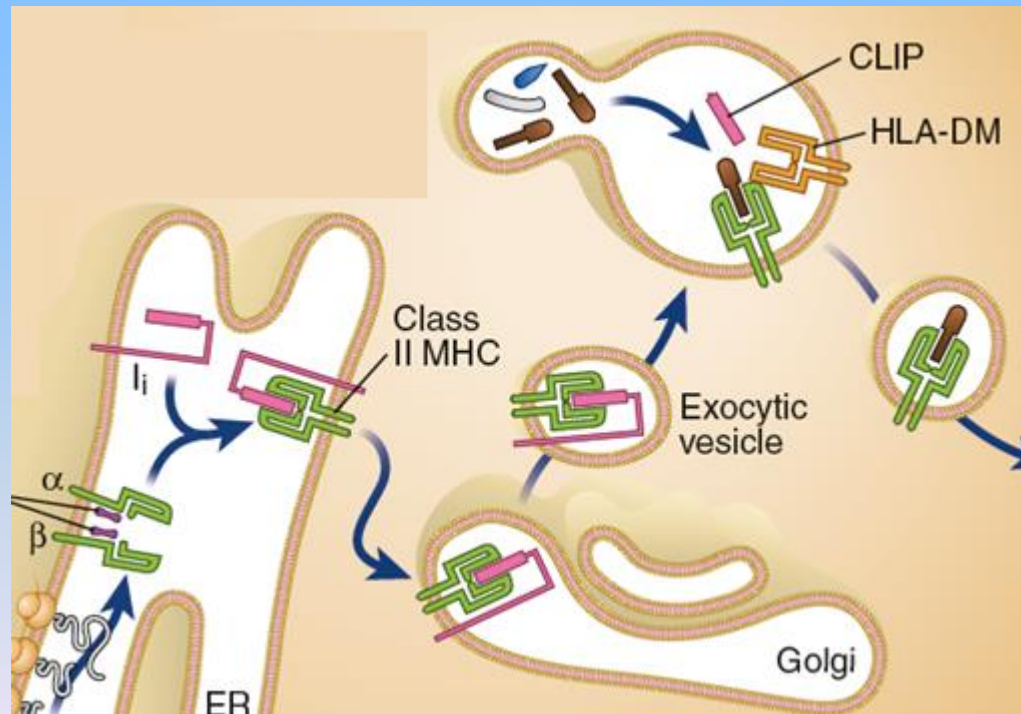


Villadangos et al., *Immunological Reviews* 172:109-20, 1999.

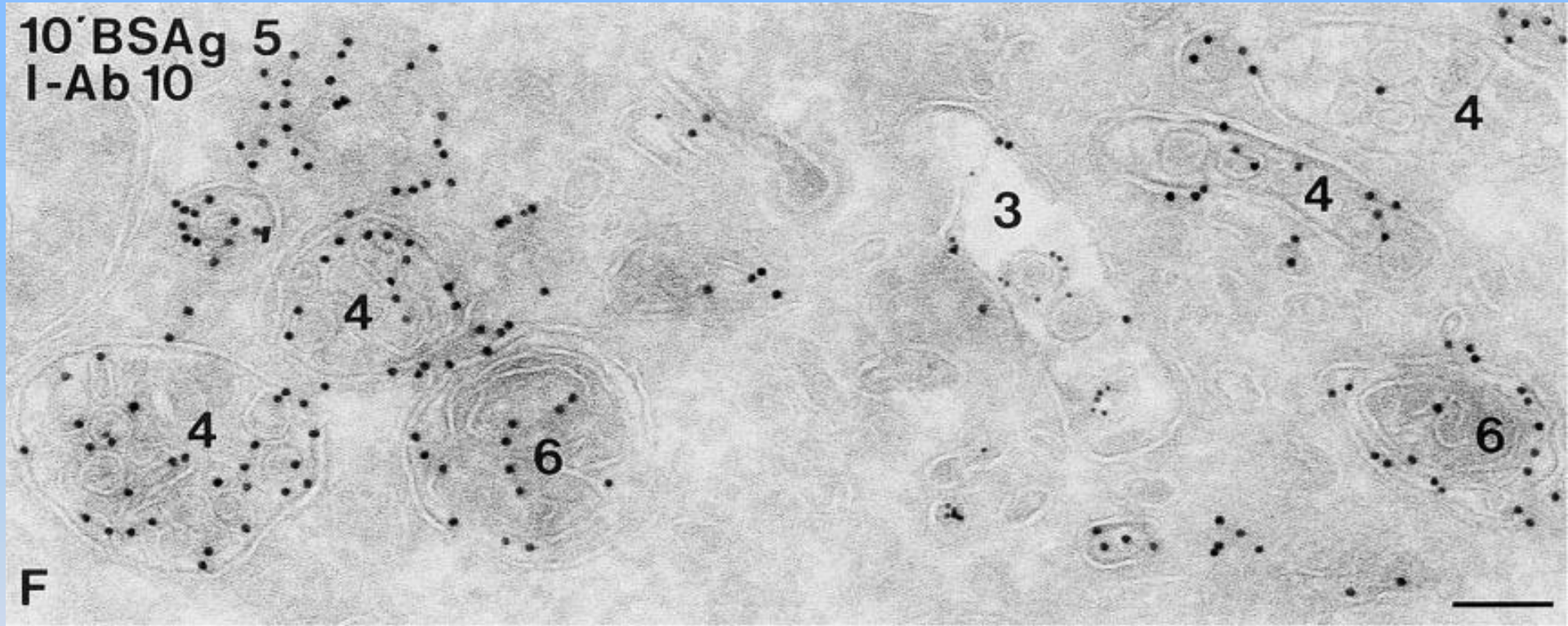
Cathepsin S and L Expression by APCs



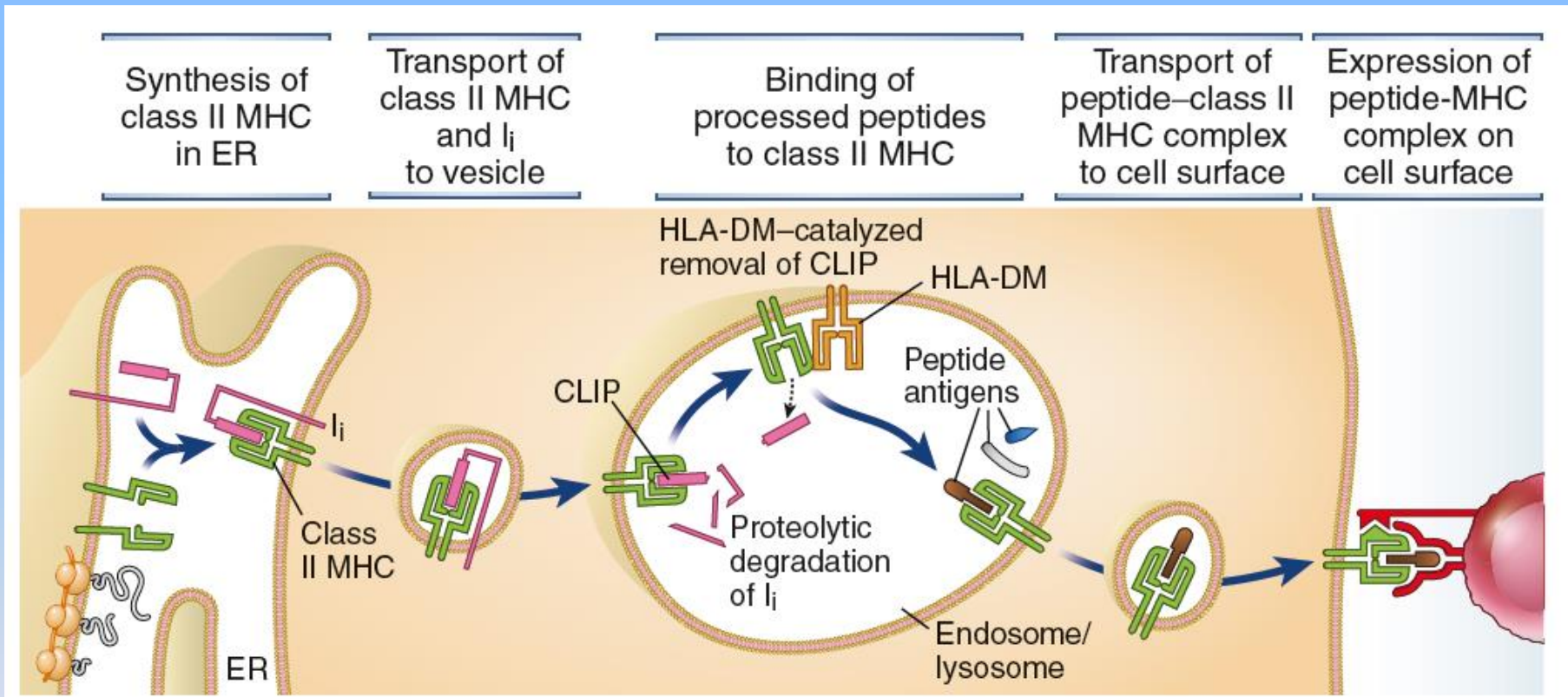
MHC Class II Pathway



Class II Compartment (MIIC)



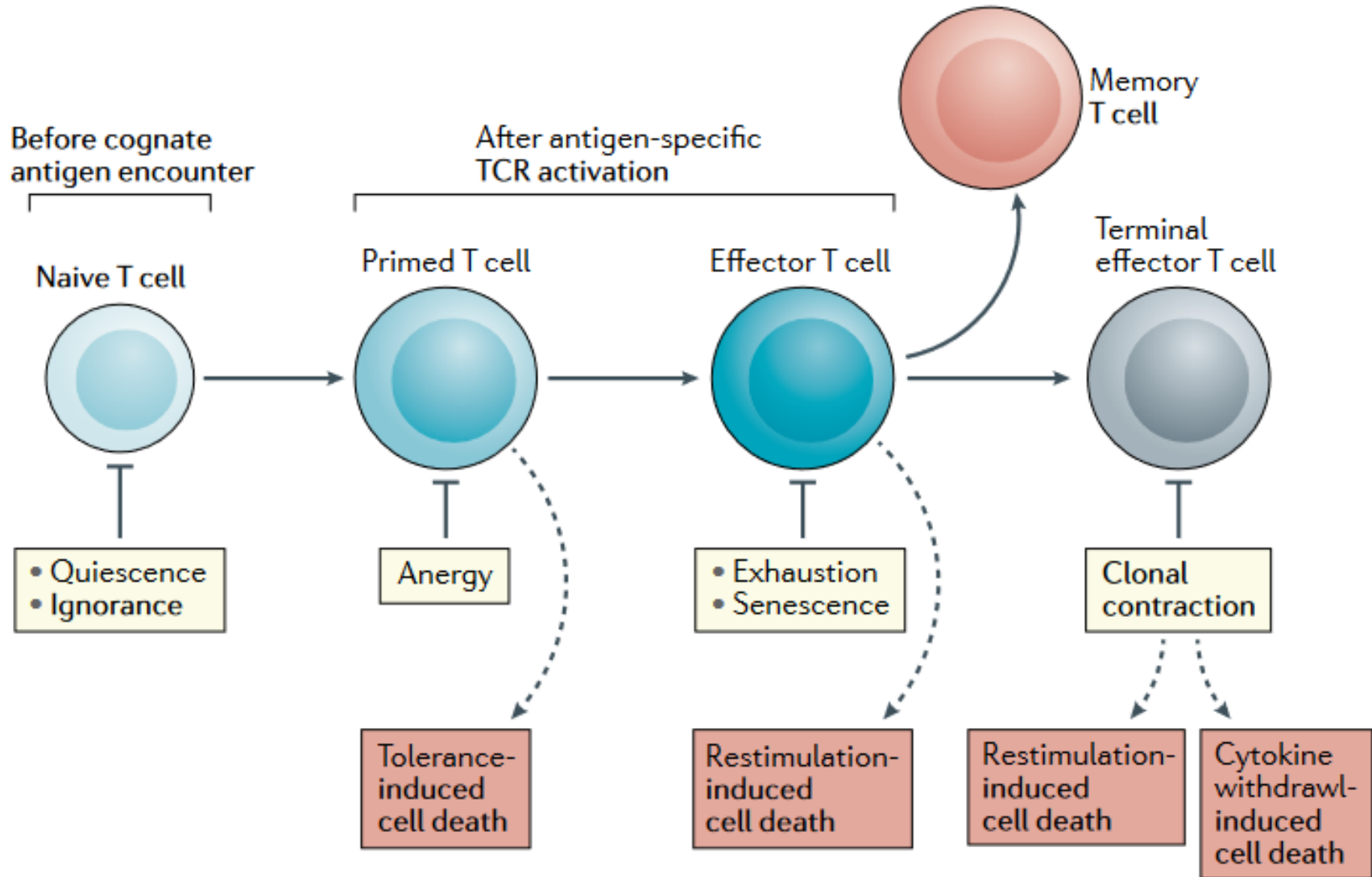
Final Steps of MHC Class II Pathway



Peripheral T cell Tolerance: Facts

- ***Thymic deletion of self-reactive T cells is only ~60–70% efficient***
- ***Peripheral naive T cell repertoire contains a significant portion of low-avidity, self-reactive T cells***
- ***Peripheral tolerance checkpoints include:***
 - 1) ***Mechanisms that act directly on the responding T lymphocyte (T cell-intrinsic mechanisms)***
 - 2) ***Mechanisms that depend on other cell subsets, such as regulatory T cells (Treg cells) and dendritic cells (T cell-extrinsic mechanisms).***

Checkpoints for Peripheral T cell Tolerance



Checkpoints for Peripheral T cell Tolerance

- ***Quiescence: sleeping?***
- ***Ignorance: it is there, but they just do not know.***
- ***Anergy: insufficient signals? The starting motor does not work...***
- ***Exhaustion: tired, they are done...***
- ***Senescence: just getting old!***
- ***Deletion: shot and killed!***

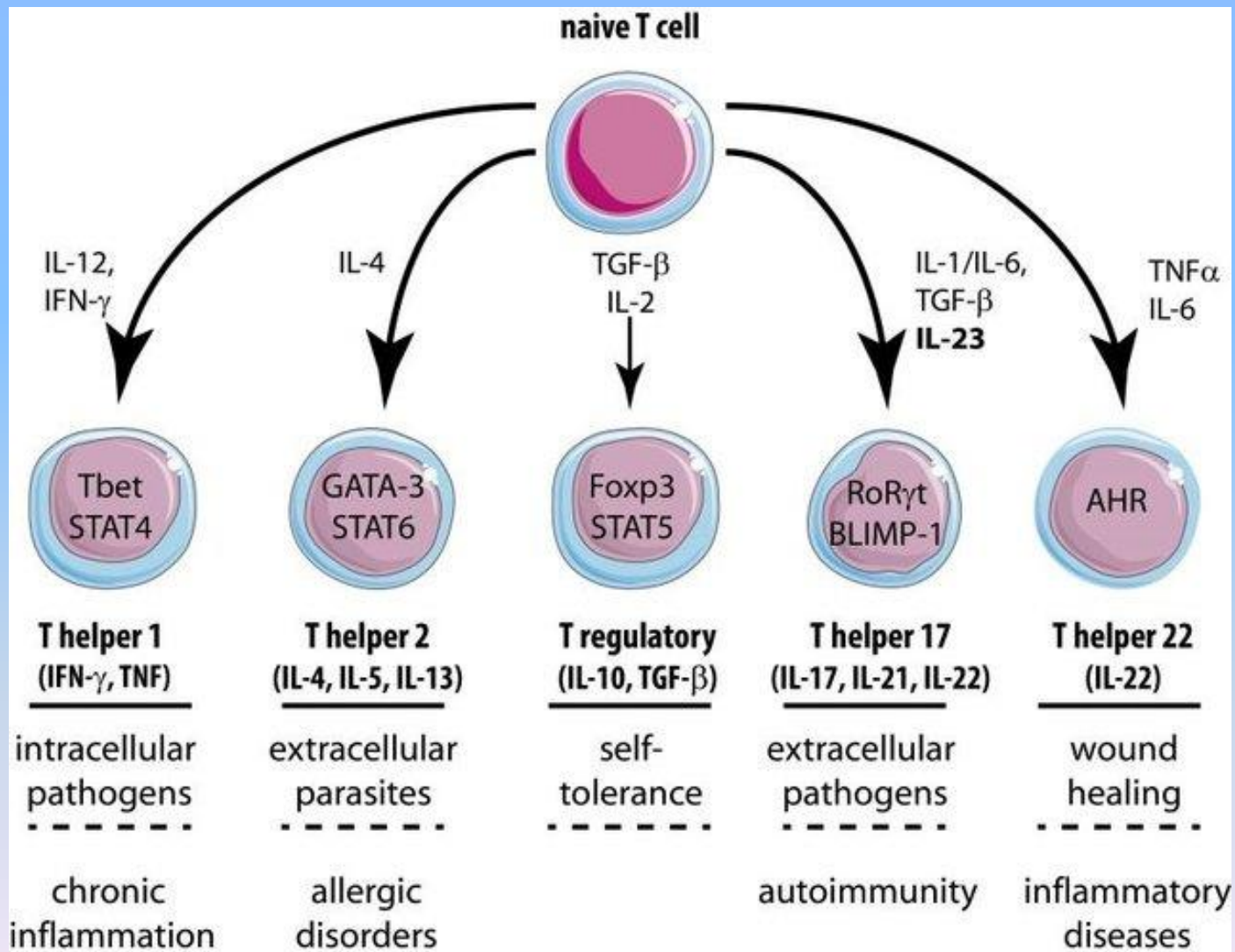
Checkpoints for Peripheral T cell Tolerance

Table 1 | Summary of the main regulators and markers of each tolerance checkpoint in T cells

Factor	Quiescence	Ignorance	Anergy	Exhaustion	Senescence	Deletional tolerance
Surface receptors	TGFβR1 (REF. ¹⁵)	Unknown	CD73 (REFS ^{78,79})	PD1 (REF. ¹⁷⁷)	NKG2D ¹⁹³	FAS (also known as CD95) ^{160,161}
	VISTA ¹⁷		FR4 (REFS ^{78,79})	TIGIT ¹⁷⁸	IFNα/IFNAR ¹¹⁹	TNFR1(REFS ^{164,181})
			LAG3 (REF. ⁸⁵)	LAG3 (REF. ¹⁷⁹)		TRAILR1 and TRAILR2 (REFS ^{182,183})
			NRP1 (REF. ⁷⁹)	TIM3 (REF. ¹⁸⁰)		
Signalling molecules	BTG1/BTG2 (REF. ¹⁸)	Unknown	DGKα ⁶²	SHP1 (REF. ¹⁸⁷)	TAB1 (REF. ¹²⁹)	CASP8 (REF. ¹⁶¹)
	TSC1/TSC2 (REFS ^{51,52})		CBLB ⁶⁷	SHP2 (REFS ^{187,188})	Sestrin 2 (REF. ¹⁵⁵)	BID ¹⁹⁰
			GRAIL ¹⁶⁴	PTPN2 (REF. ¹⁸⁹)		BIM (also known as BCL-2L1) ^{145,146}
			ITCH ^{185,186}			
Transcription factors	KLF2 (REFS ^{20,22})	Unknown	NFAT1 (REF. ⁶⁴)	IRF4 (REF. ¹⁹³)	–	–
	FOXO1 (REFS ^{26–28})		EGR2 (REFS ^{66–68})	NR4A1 (REF. ¹⁰⁵)		
	RUNX1 (REFS ^{29,30})		EGR3 (REF. ⁶⁷)	GATA3 (REF. ¹⁹⁴)		
	TOB1 (REFS ^{14,15})		NR4A1 (REF. ⁷²)	TOX ^{105–106}		
	FOXP1 (REFS ^{191,192})		TOB1 (REF. ¹⁵)	BATF ¹⁰²		
			BLIMP1 (REF. ¹⁹⁵)			
			EOMES ¹⁹⁶			

DGKα, diacylglycerol kinase-α; IFNα, interferon-α.

Development of T cell subsets



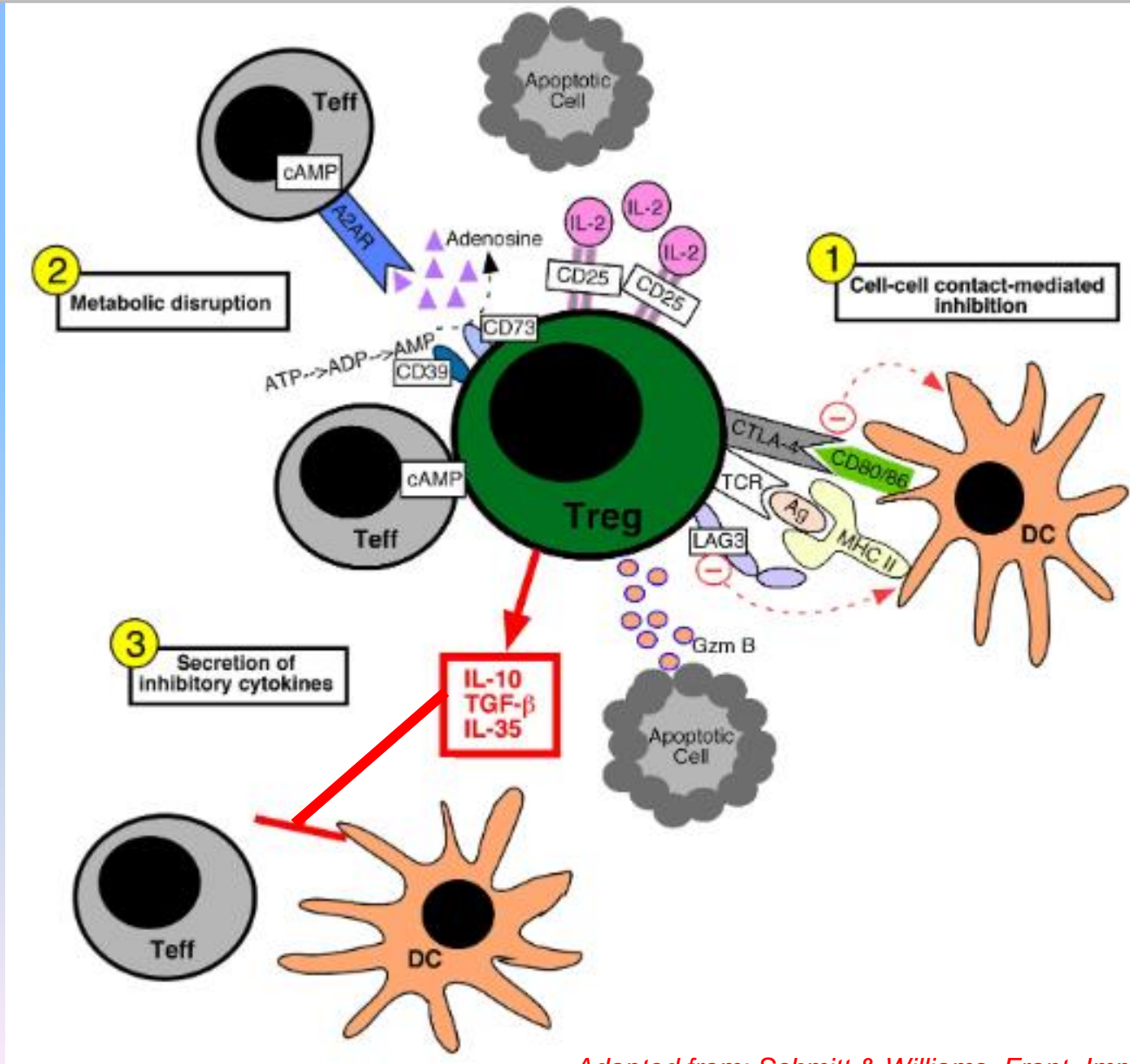
Regulatory T cell subsets

TABLE 1 | Different Treg subsets identified in human and/or mouse.

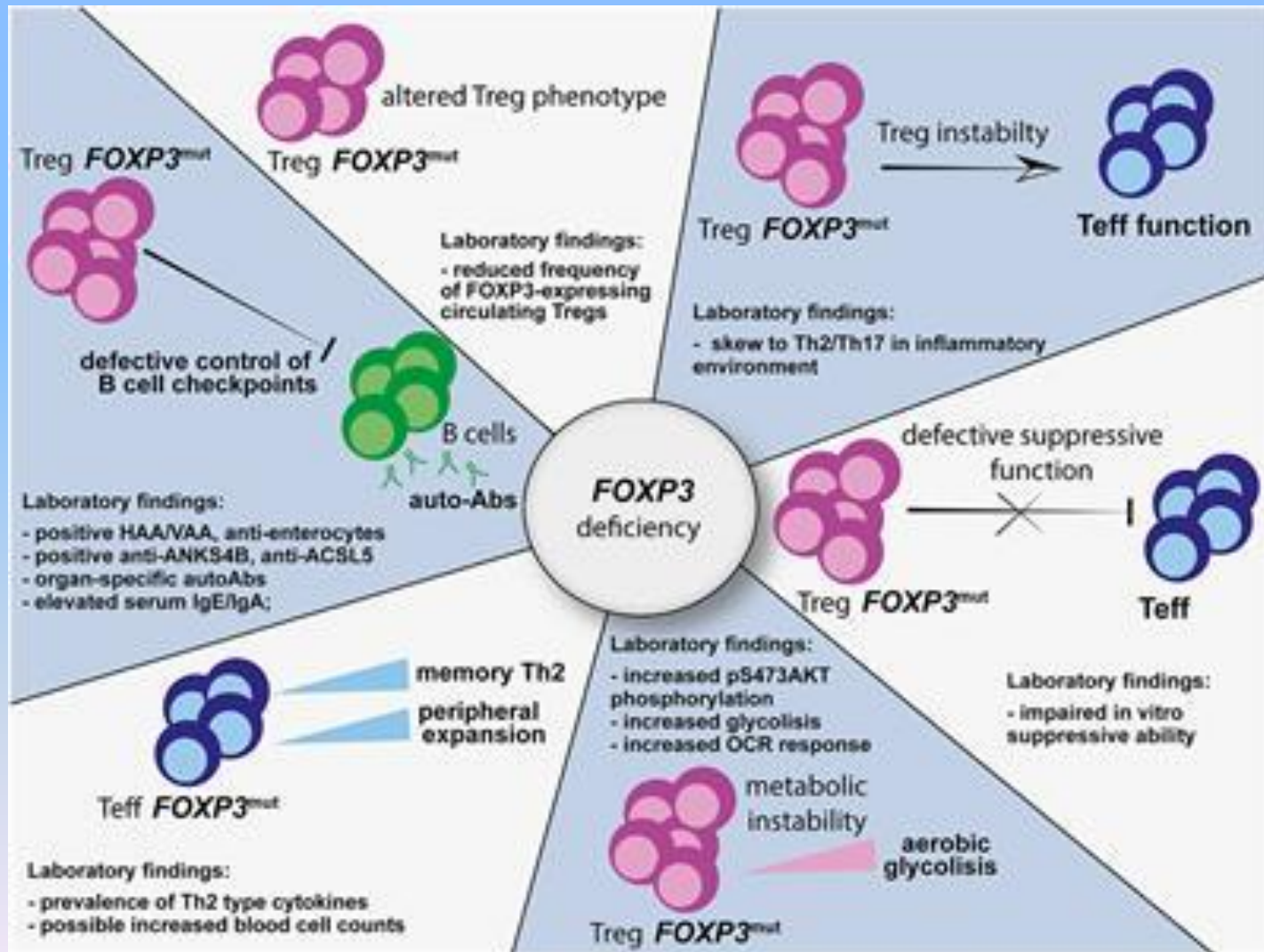
Cell population	Human	Mouse	Reference
CD4 ⁺ CD25 ⁺ Foxp3 ⁺	✓	✓	(5, 6)
CD4 ⁺ Foxp3 ^{low} CD45RA ⁺	✓	✗	(12)
CD4 ⁺ Foxp3 ^{high} CD45RA ⁻	✓	✗	(12)
CD4 ⁺ Foxp3 ⁻ IL10 ⁺ (Tr1)	✓	✓	(37)
CD4 ⁺ Foxp3 ⁻ TGFβ ⁺ (Th3)	✓	✓	(38)
CD4 ⁺ Foxp3 ⁻ IL35 ⁺ (iTr35)	✓	✓	(39)
CD4 ⁺ Foxp3 ⁻ IL10 ⁺ TGFβ ⁺ (Treg of B cells)	✗	✓	(40)
CD8 ⁺ Foxp3 ⁺ and/or CD28 ⁻ and/or CD25 ⁺	✓	✗	(41–43)
CD8 ⁺ CD45RA ⁺ CCR7 ⁺ Foxp3 ⁺	✓	✗	(44)
CD8 ⁺ CD45RC ^{low/-}	✓	✓	(45)
CD8 ⁺ CD122 ⁺	✗	✓	(46)
CD8 ⁺ HLA-DR ⁺	✓	✗	(47)
CD8 ⁺ HLA-E ⁺	✓	✗	(48)
CD8 ⁺ Qa-1 ⁺	✗	✓	(48)
γδ T cells	✓	✓	(49)
NKT	✓	✓	(50)

Treg cells are categorized based on CD4 or CD8 surface markers expression.

Mechanisms of Suppression by Regulatory T cells




Immune dysregulation, polyendocrinopathy, enteropathy, X-linked (IPEX) syndrome




Immune dysregulation, polyendocrinopathy, enteropathy, X-linked (IPEX) syndrome

Classical triad of symptoms

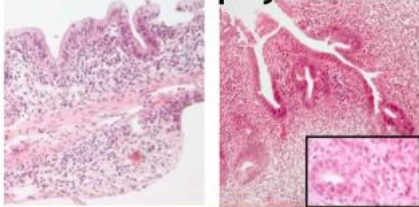
Dermatitis




Type 1 diabetes




Enteropathy with villous atrophy




Alopecia




Autoimmune thyroiditis



Recurrent infections

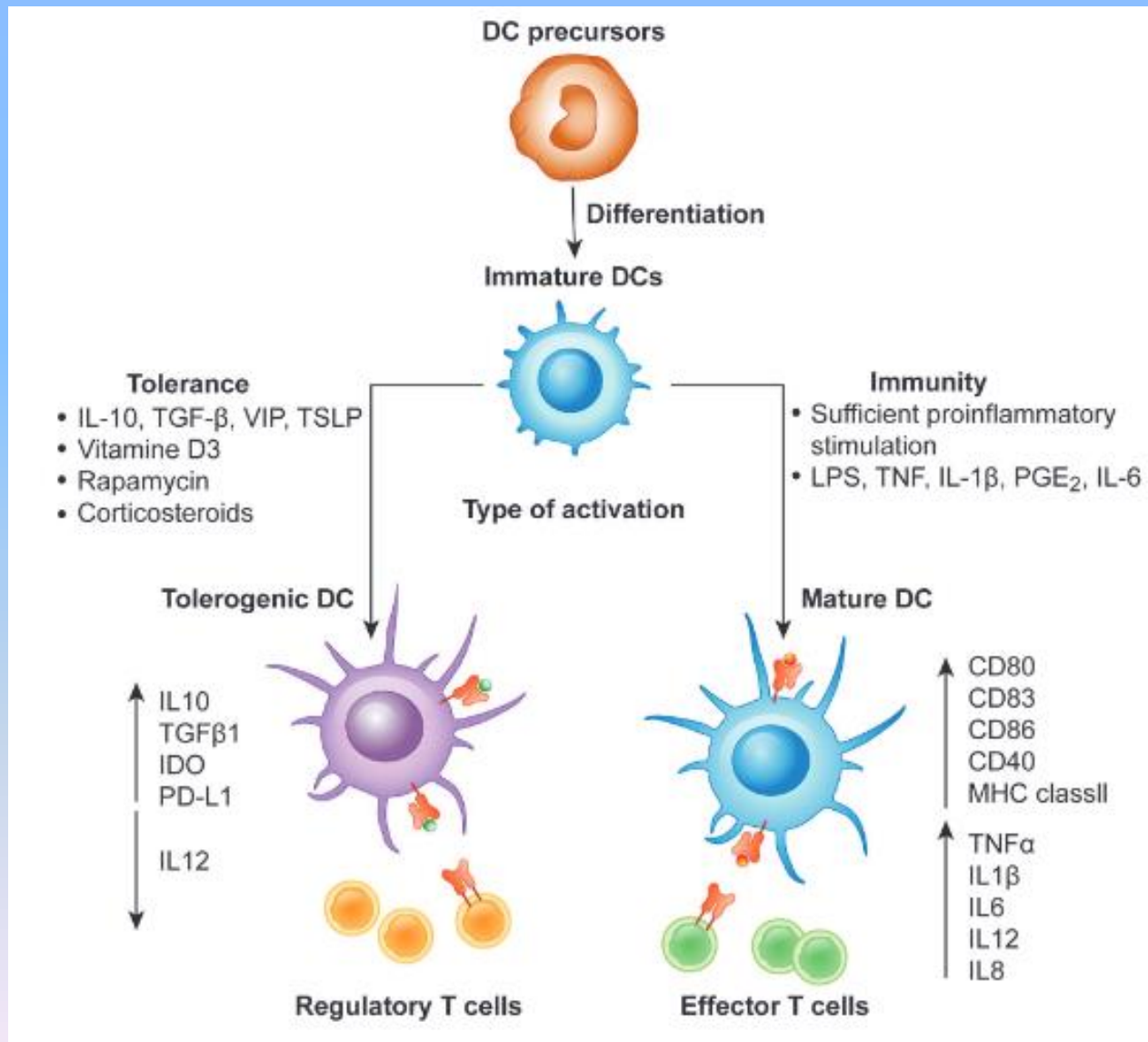


Autoimmune hepatitis

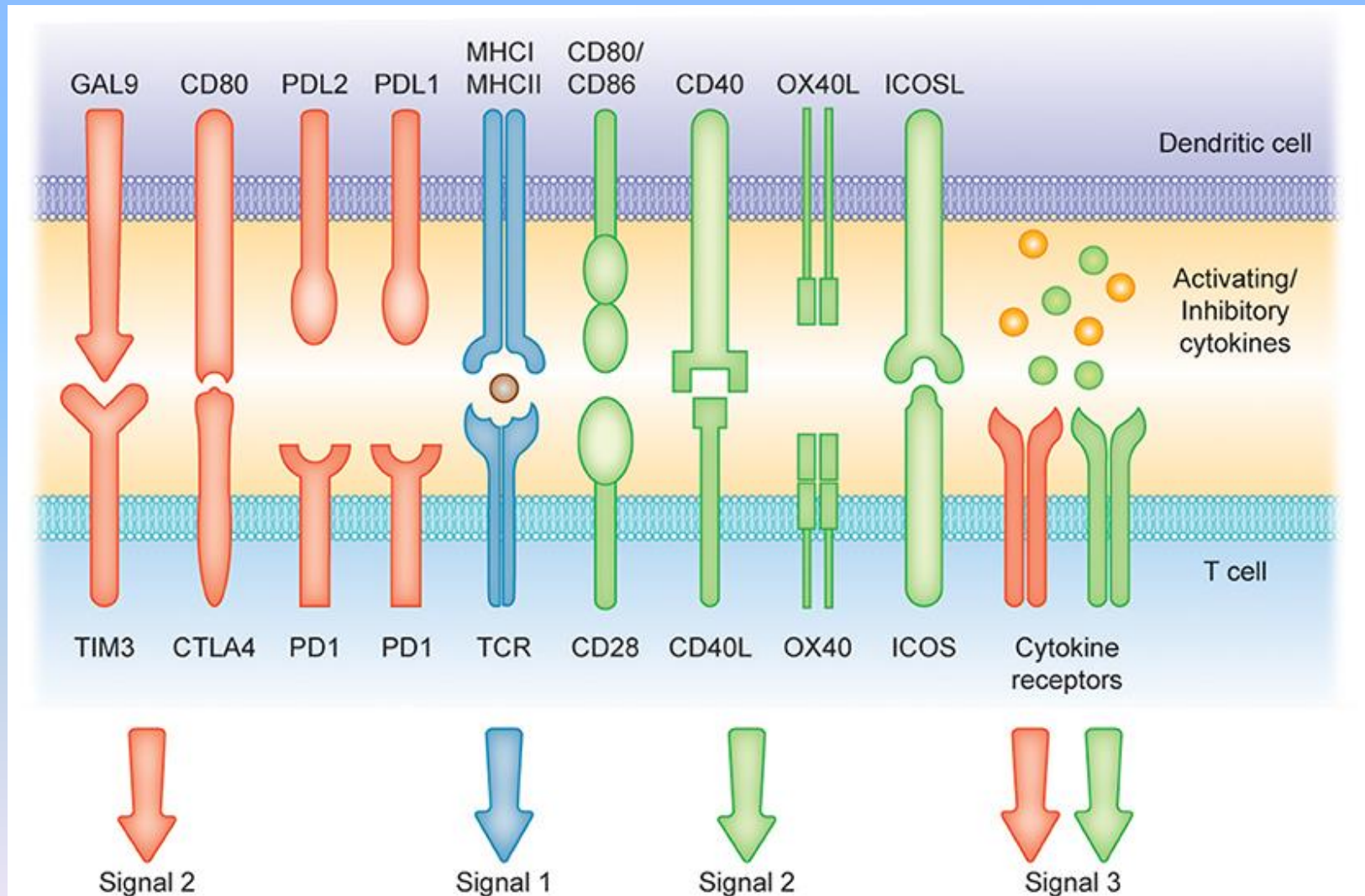


Additional symptoms

Dendritic cells: tolerance versus immunity



Tolerance or immunity depend on the molecular Interaction with T cells



Extrathymic Role for AIRE in peripheral DCs

Cell
PRESS

Immunity
Article

Extrathymic *Aire*-Expressing Cells Are a Distinct Bone Marrow-Derived Population that Induce Functional Inactivation of CD4⁺ T Cells

James M. Gardner,^{1,2,7} Todd C. Metzger,^{1,7} Eileen J. McMahon,^{1,3} Byron B. Au-Yeung,⁴ Anna K. Krawisz,¹ Wen Lu,¹ Jeffrey D. Price,⁵ Kellsey P. Johannes,¹ Ansuman T. Satpathy,⁶ Kenneth M. Murphy,⁶ Kristin V. Tarbell,⁵ Arthur Weiss,⁴ and Mark S. Anderson^{1,*}

¹Diabetes Center, University of California, San Francisco, San Francisco, CA 94143-0540, USA

²Department of Surgery, University of California, San Francisco, San Francisco, CA 94143-0540, USA

³Department of Biology, Westmont College, Santa Barbara, CA 93108, USA

⁴Howard Hughes Medical Institute, Rosalind Russell Medical Research Center for Arthritis, Department of Medicine, Department of Microbiology and Immunology, University of California, San Francisco, San Francisco, CA 94143-0540, USA

⁵Immune Tolerance Section, Diabetes Branch, National Institute of Diabetes and Digestive and Kidney Diseases, National Institutes of Health, Bethesda, MD 20892, USA

⁶Howard Hughes Medical Institute, Department of Pathology and Immunology, School of Medicine, Washington University in St. Louis, St. Louis, MO 63110, USA

⁷These authors contributed equally to this work

*Correspondence: manderson@diabetes.ucsf.edu

<http://dx.doi.org/10.1016/j.immuni.2013.08.005>