

**Programa de Pós-graduação em Imunologia ICB/USP**

**Disciplina BMI 5904**

**Reconhecimento no Sistema Imune**



## **Aula 7 - MHC/HLA**

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**Lab. Imunogenetica/Dep.Imunologia/ICB/USP**

# Reconhecimento

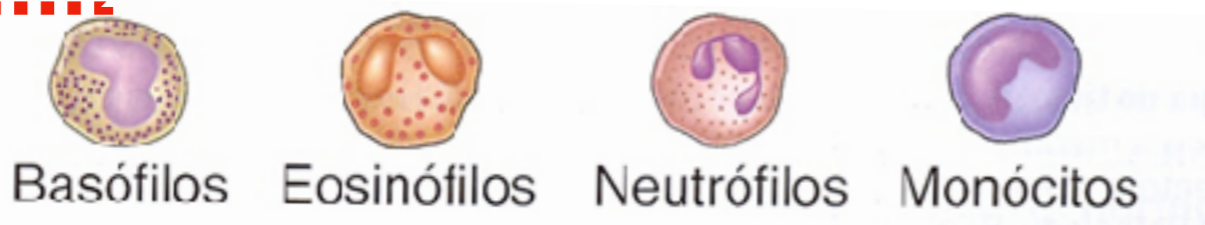
## SISTEMA IMUNE



**PADRÕES MOLECULARES  
(MAMPs, PAMPs, NAMPs,  
DAMPs, HAMPs, VAMPs....)**



**PRRs**



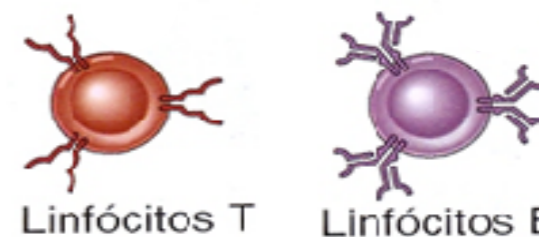
**&...todas as outras células**

*20-40 genes para reconhecer um limitado  
numero de pattern “mais prevalentes e  
conservados”*

**ANTIGENOS  
ESPECIFICOS**



**TCR, BCR, Ig**



*Rec com potencial de  
reconhecer um amplo leque  
de antígenos (recombinação  
somática randômica)*

# MHC/HLA

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- ◇ Glico-proteínas (super família das Ig) da superfície celular
- ◇ Codificadas pelos genes MHC (humanos: HLA no chr 6)
- ◇ Tipo: **MHC-I e MHC-II, e MHC-I non-classica, MHC-I like**

MHC-I, MHC-II

elevado polimorfismo

MHC-I non classicas

baixo polimorfismo

MHC-I like

monomorfico



# MHC/HLA

Glico-proteínas (super familia das Ig)

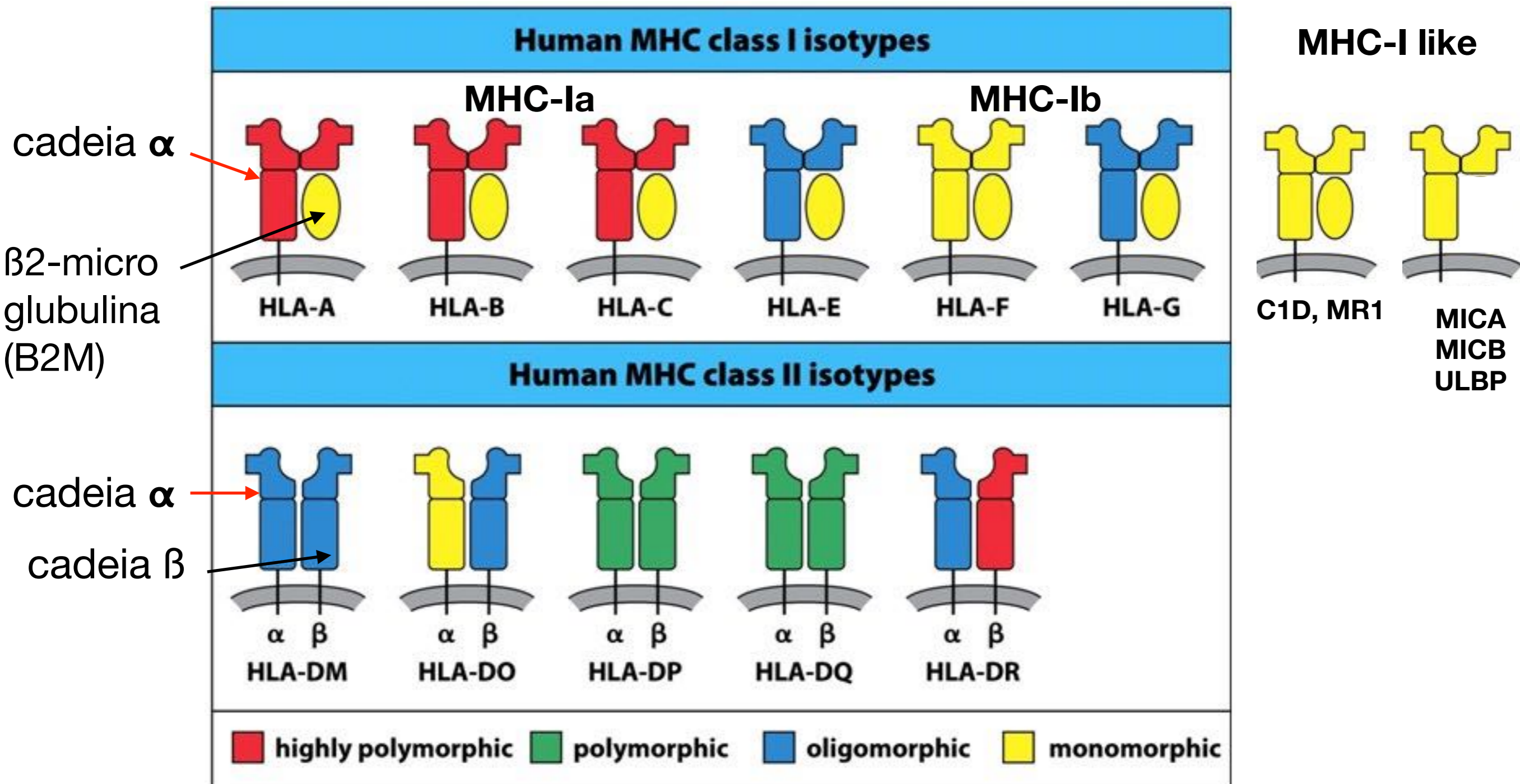


Figure 5.24 The Immune System, 3ed. (© Garland Science 2009)

# MHC/HLA

<i>MHC I</i>	<i>Immune function</i>	<i>ligand binding between the <math>\alpha 1</math> and <math>\alpha 2</math> domains</i>	<i>TCR binding</i>	<i>NKR binding</i>	<i><math>\beta_2m</math>-associated</i>
<i>Classical MHC I molecules</i>					
HLA-A	Yes	Peptide	Yes	LCR/ LIR, KIR <sup>a</sup>	Yes
HLA-B	Yes	Peptide	Yes	LIR, KIR <sup>a</sup>	Yes
HLA-C	Yes	Peptide	Yes	LIR, KIR	Yes
<i>Non-classical MHC I molecules</i>					
HLA-E	Yes	Peptide	Yes	CD94/NKG2A/C, LIR	Yes
HLA-G	Yes	Peptide	Yes	LIR, KIR2DL4 <sup>165</sup>	Yes
HLA-F	Yes <sup>b</sup>	No <sup>c</sup>		LIR	Yes
CD1	Yes	Lipids	Yes		Yes
MICA	Yes	No	Yes ( $\gamma\delta$ lineage <sup>167</sup> )	NKG2D	No
MICB	Yes	No	Yes ( $\gamma\delta$ lineage <sup>167</sup> )	NKG2D	No
ULBPs	Yes	No	ULBP4 ( $\gamma\delta$ lineage <sup>168</sup> )	NKG2D	No
FcRn <sup>169</sup>	Yes	No	No		Yes
HFE <sup>170</sup>	No	No	No		Yes
MR1 <sup>171</sup>	Yes	Vitamin B metabolites	Yes (MAIT cells)		Yes
EPCR <sup>172</sup>	Yes	Phospholipid	Yes ( $\gamma\delta$ lineage <sup>173</sup> )		No
ZAG <sup>174</sup>	No	Fatty acids			No
<i>HCMV-encoded MHC I-like molecules</i>					
UL18	Yes	Peptide	No	LIR1	Yes
UL142	Yes				
UL37					

Chr6

Chr6

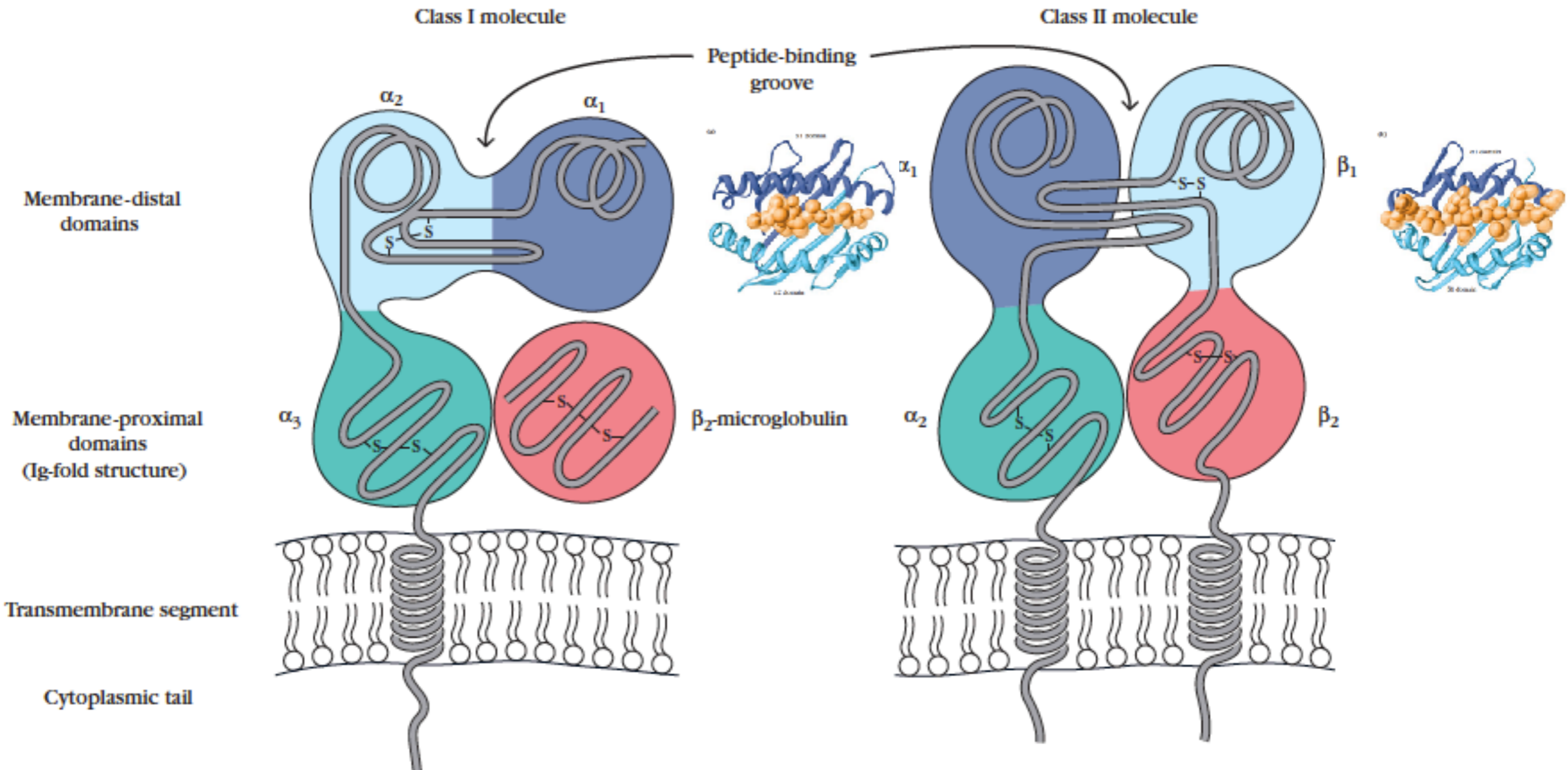
# MHC/HLA

## MHC-I

- 1 cadeia variável ( $\alpha$ )
- 1 cadeia invariável ( $\beta_2$ -microglobulina)
- Sítio de ligação do Ag ( $\alpha_1 + \alpha_2$ ): **8-10 AA**

## MHC-II

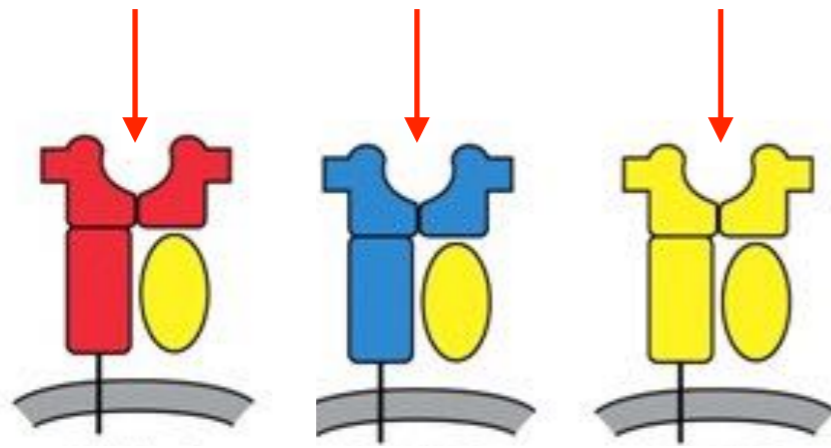
- 1 cadeia variável ( $\alpha$ )
- 1 cadeia variável ( $\beta$ )
- Sítio de ligação do Ag ( $\alpha_1 + \beta_1$ ): **13-18 AA**



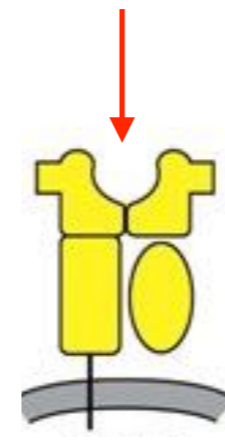


# Classical & Non-classical MHC-I

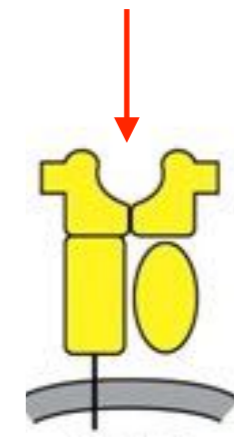
Broadly similar structure and binding groove



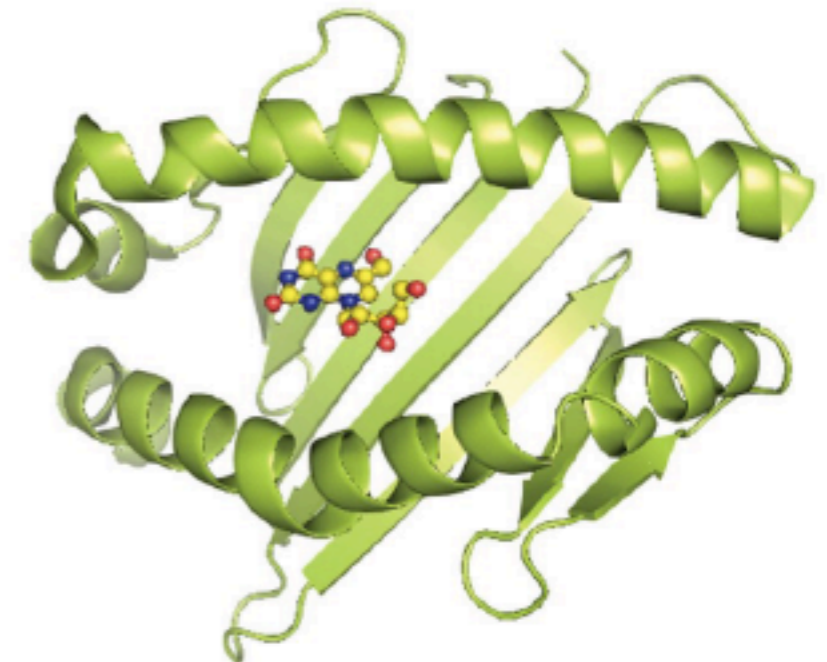
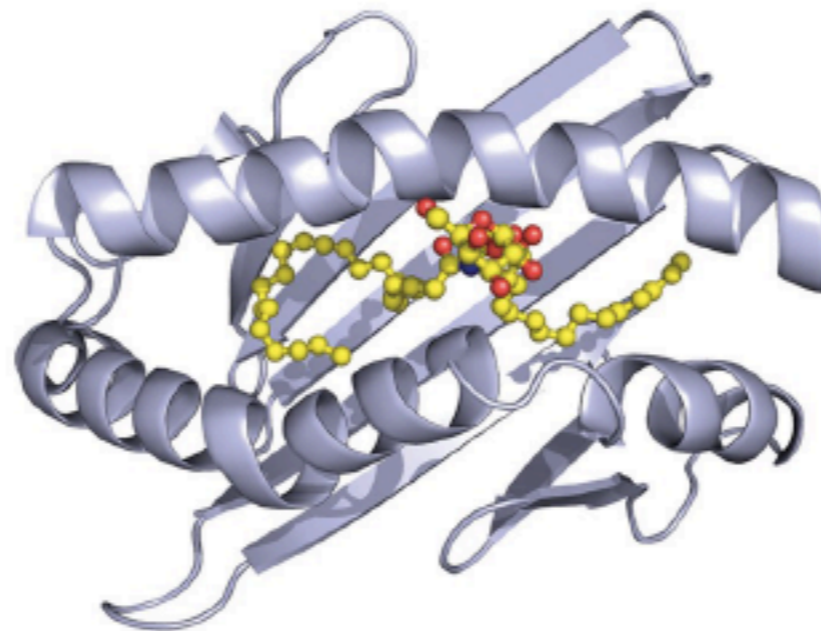
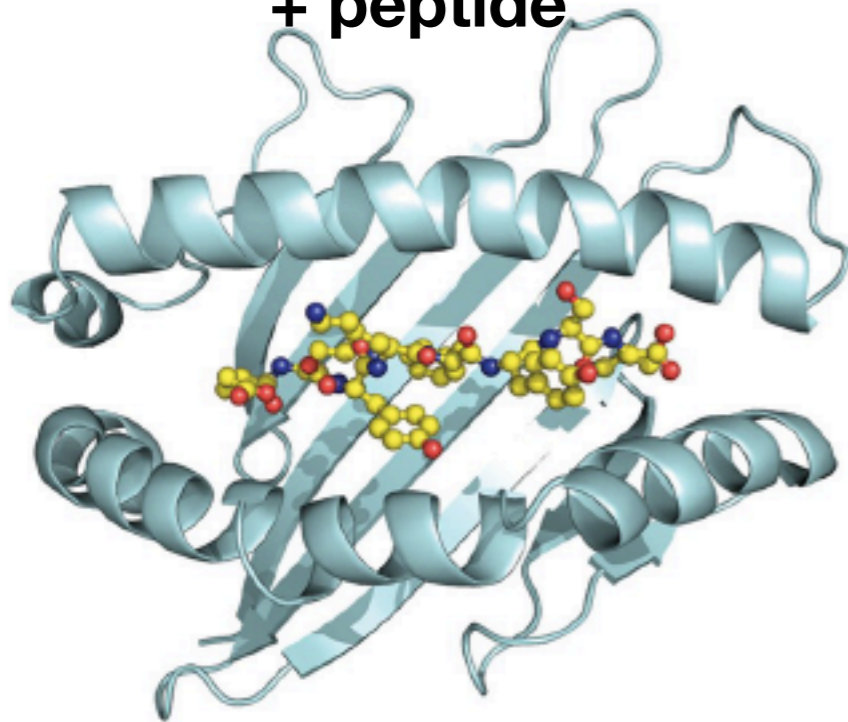
**MHC-I**  
+ peptide



**CD1**  
+ lipid



**MR1**  
+ small metabolite



**Endogenous molecules**

# Apresentação de Ag via MHC-I

**MHC presente em todas as células!**

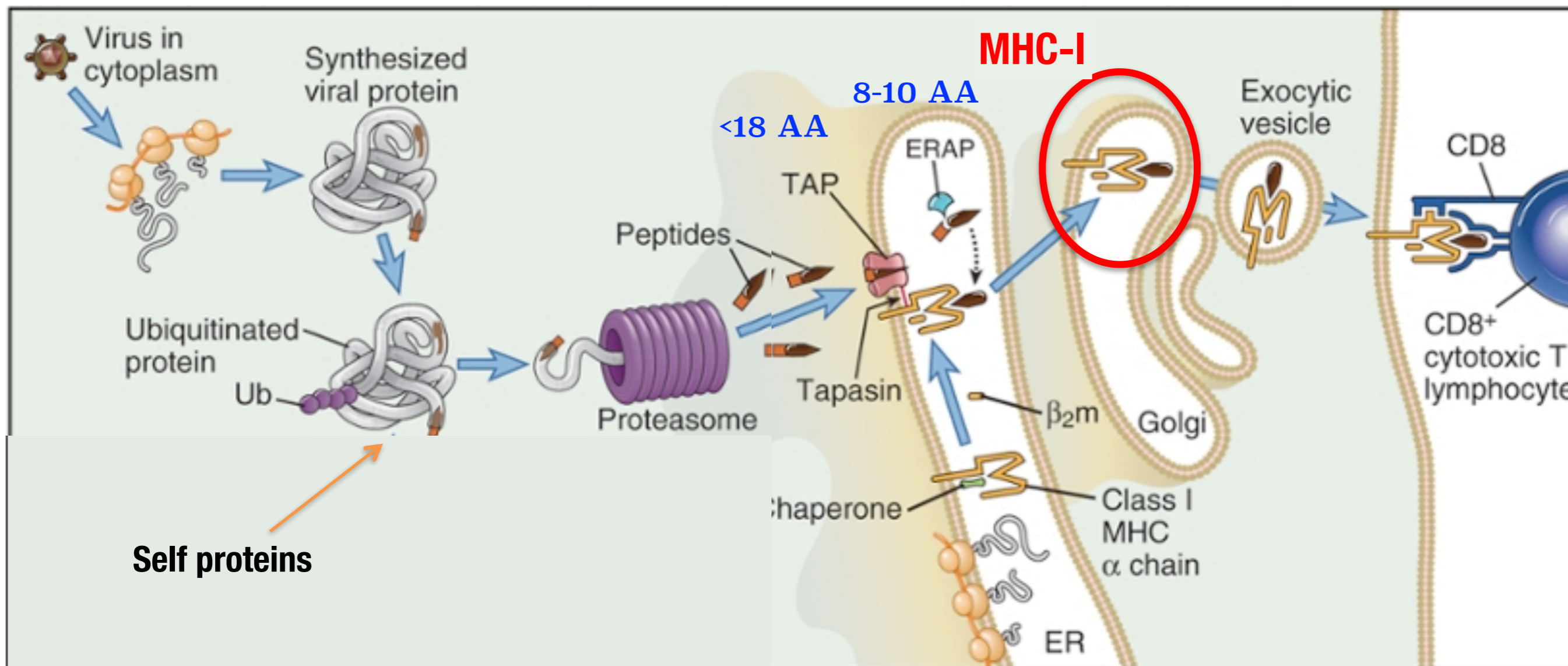
no citosol de qualquer célula  
turn over de proteínas endógenas  
infecção viral  
→ síntese de proteínas virais

Digestão no  
**PROTEASOMA**  
→ peptídeos de  
até 18 AA

Peptídeos  
transportados  
no ER (TAP)  
ERAP1/2: 8-10 AA

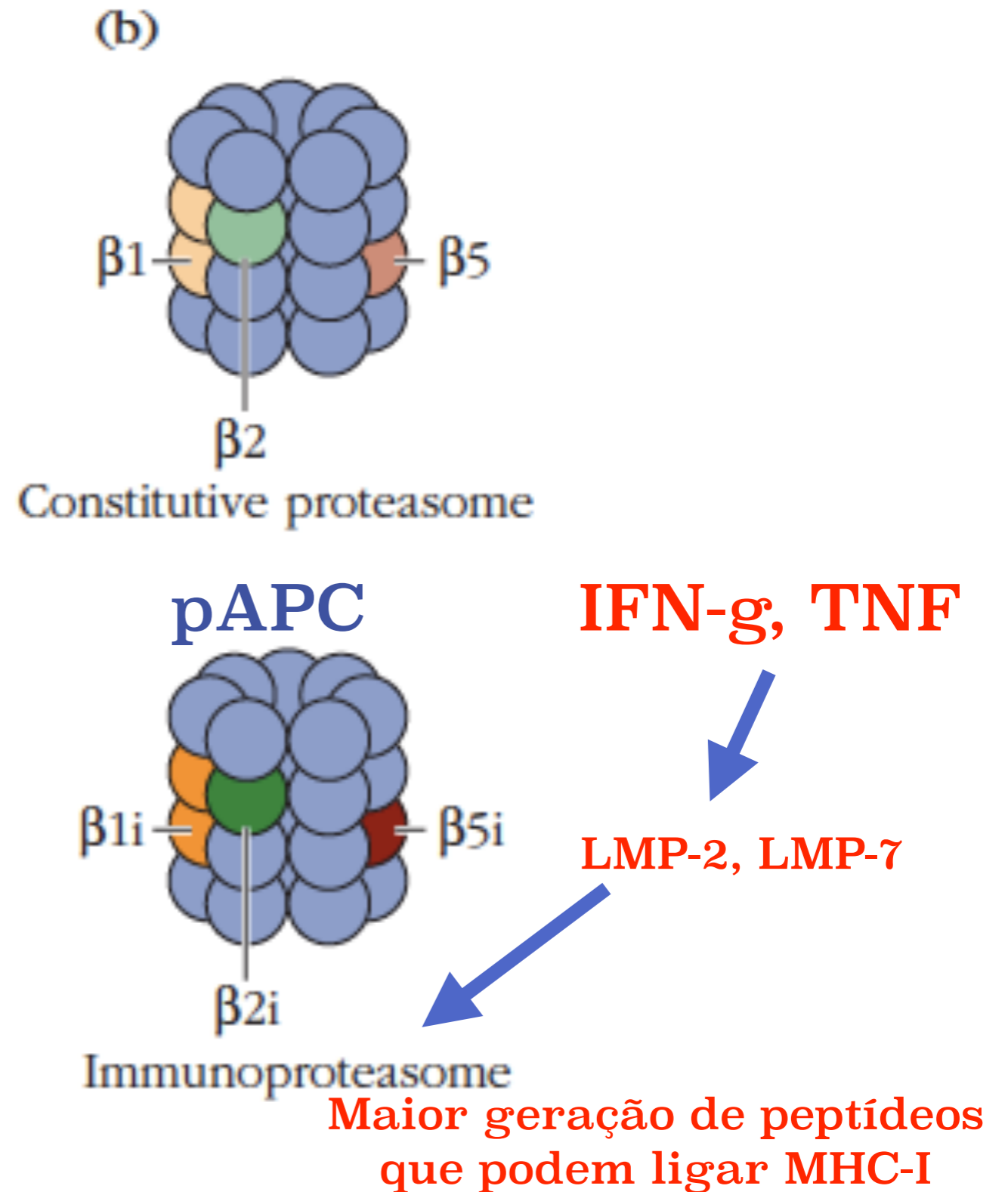
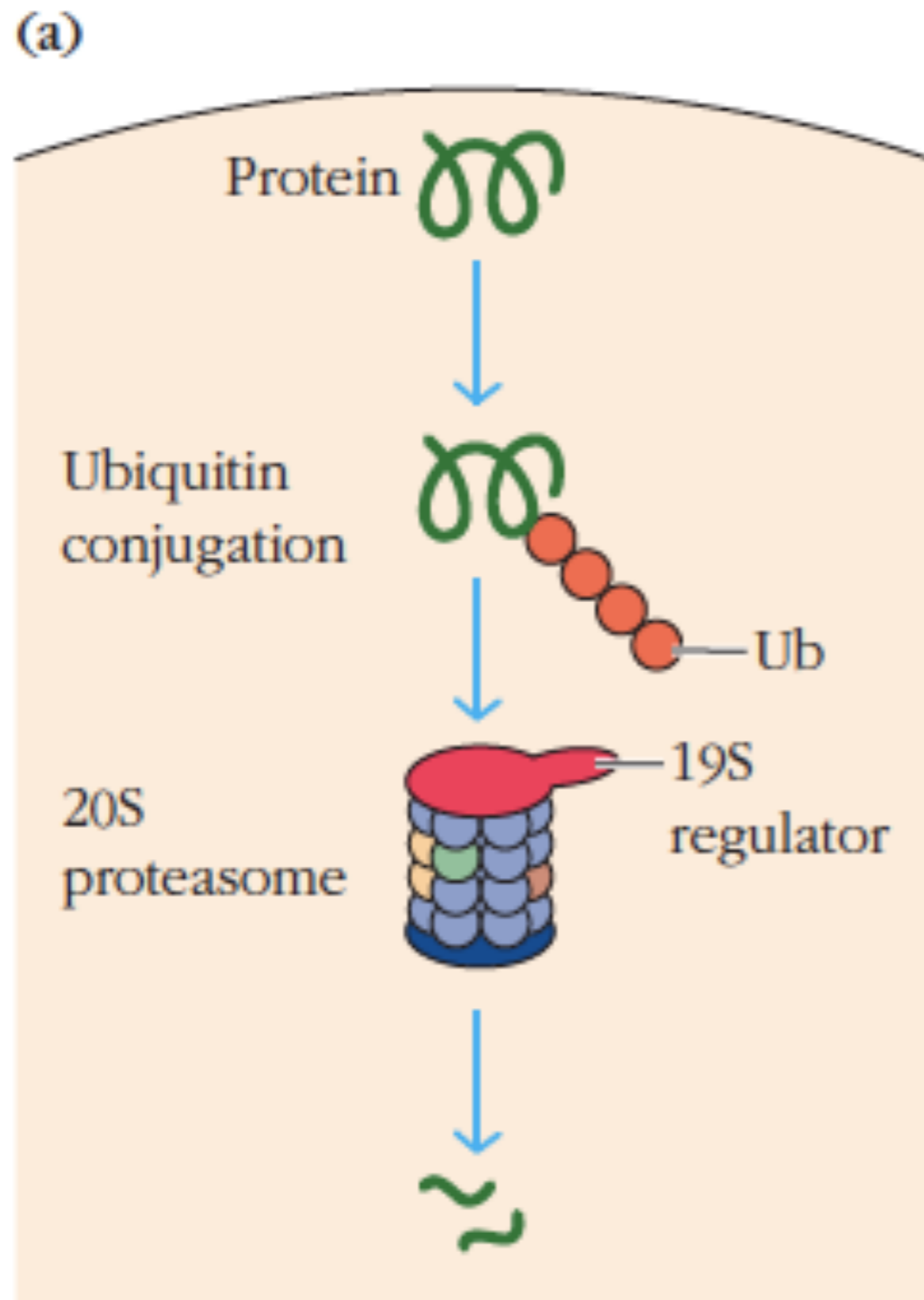
Peptídeos  
encaixam  
no MHC-I

Complexos  
MHC-I/peptídeo  
apresentados na S  
(via Golgi)

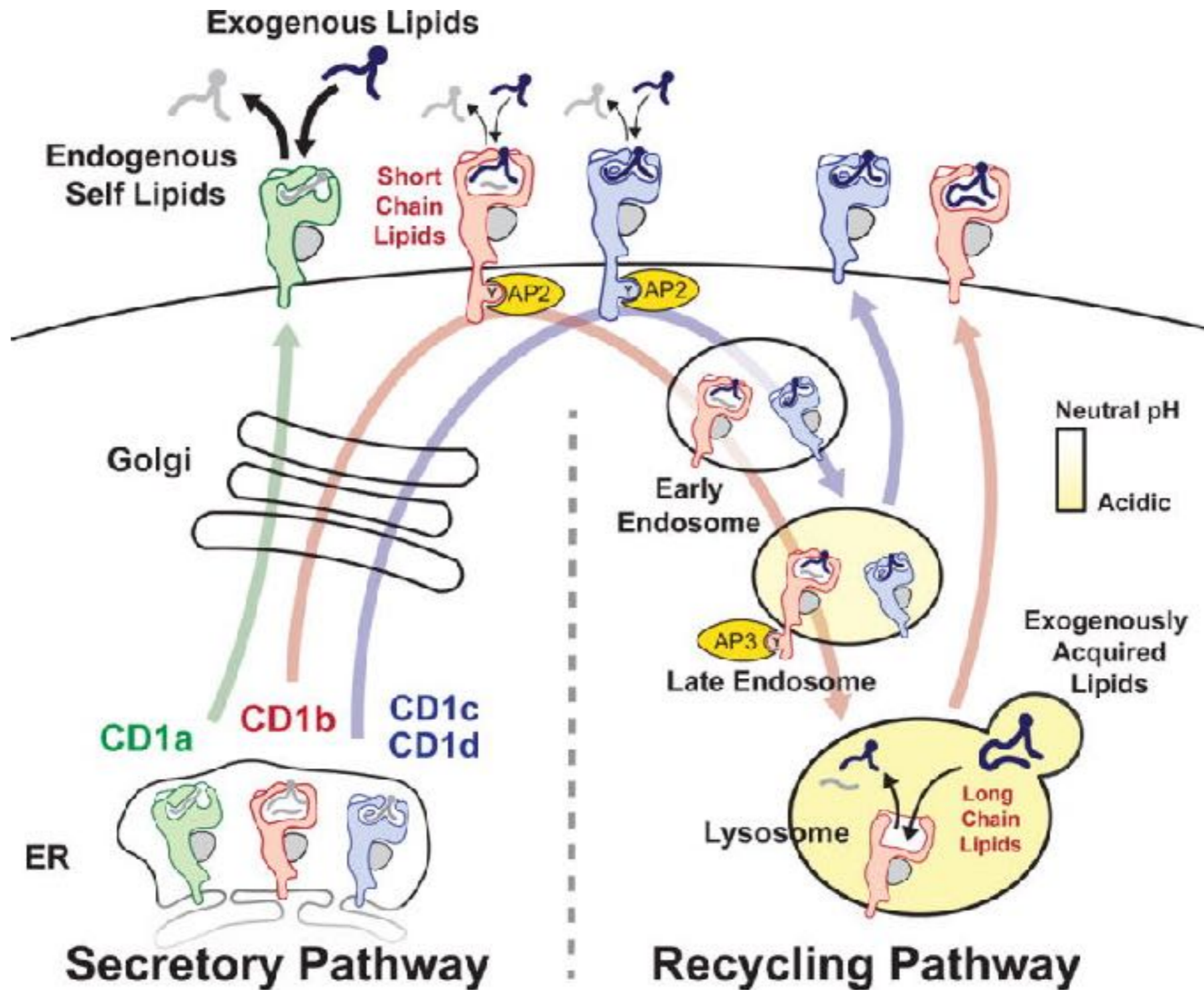




# Proteassoma e Immunoproteassoma

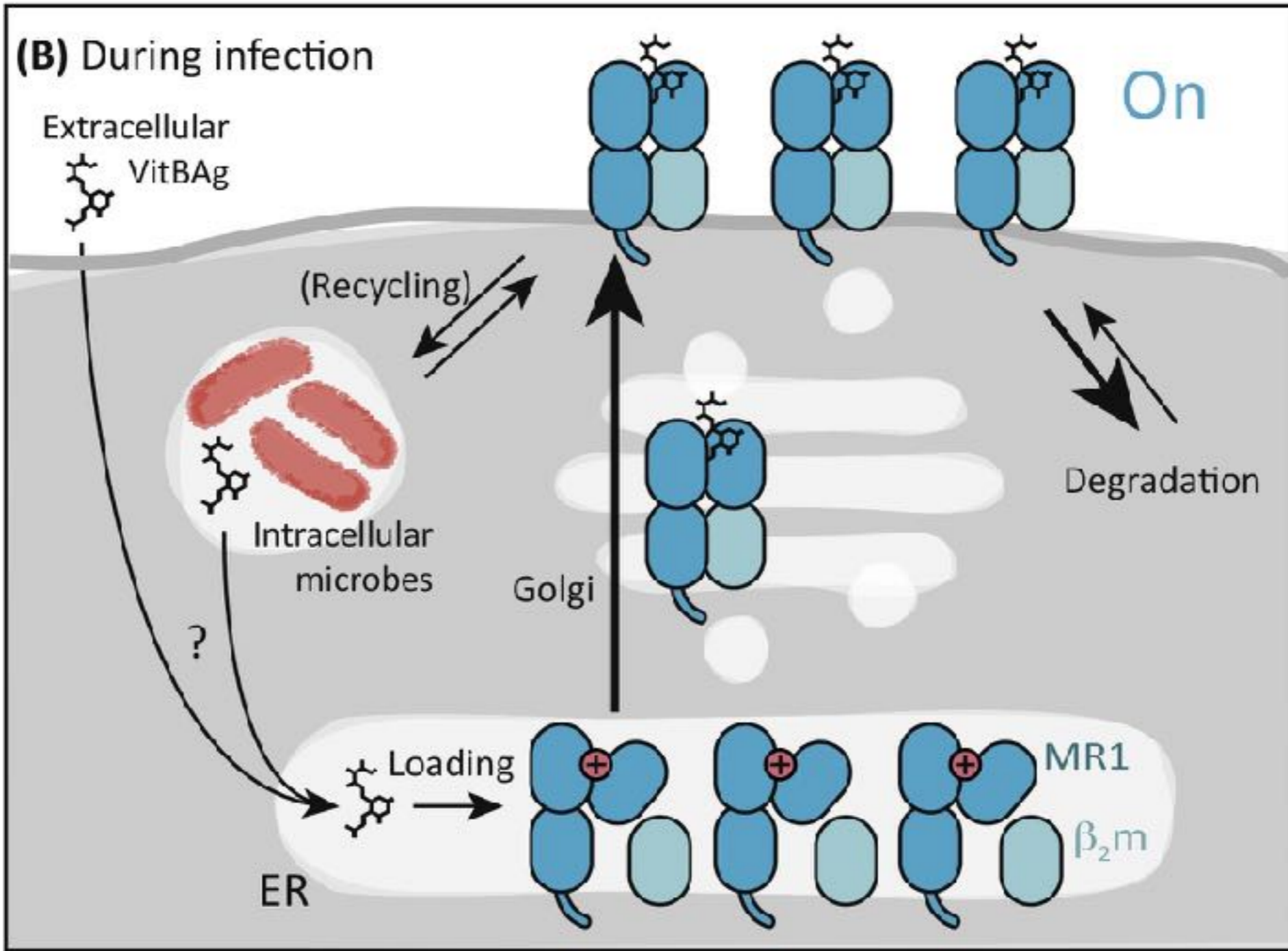


# Apresentação de lipídios via CD1



# Apresentação de metabólitos via MR1

metabolites derived from the synthesis of vitamin B2 (riboflavin)





# Apresentação de Ag via MHC-II

## MHC presente nas APC profissionais

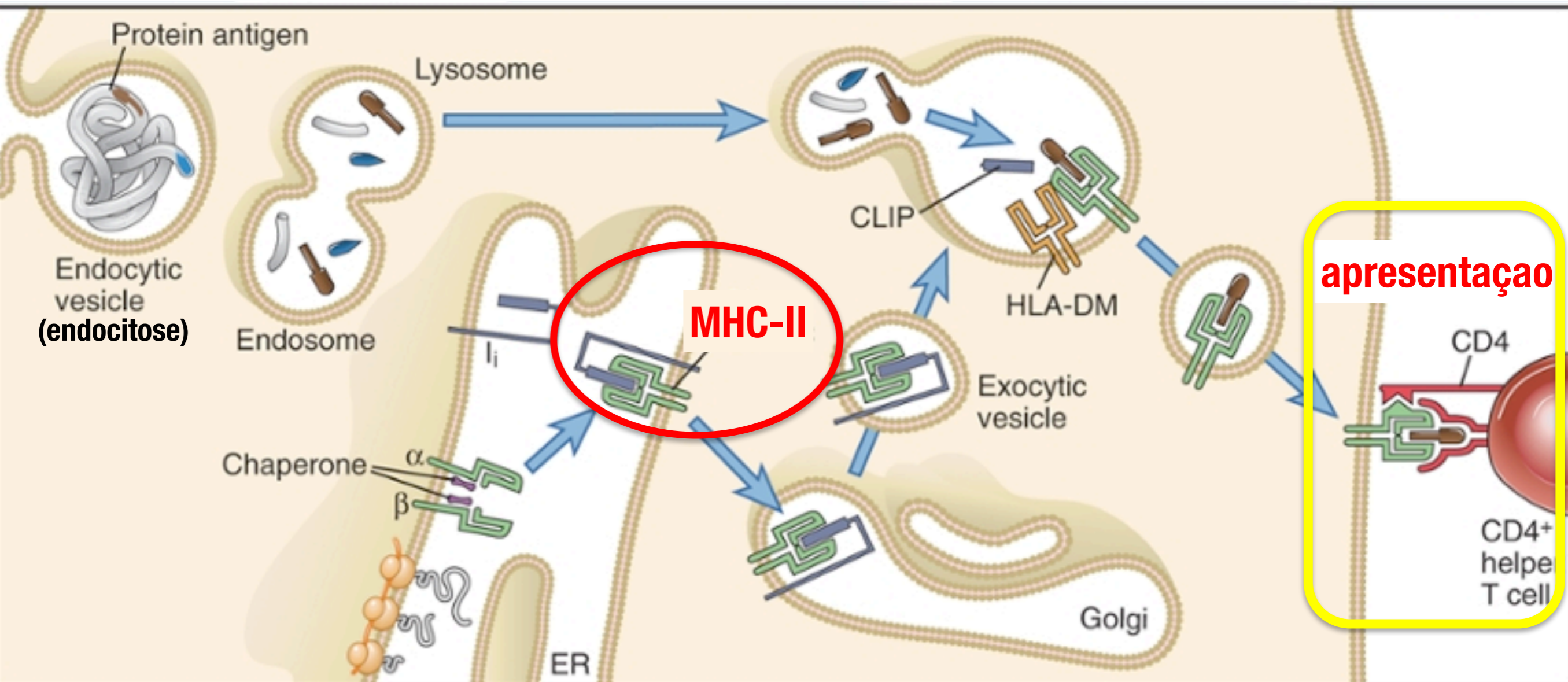
Captação de proteínas extracelulares em compartimentos vesiculares

Digestão das proteínas

Biosíntese e transporte do MHC-II aos endosomas (via Golgi)

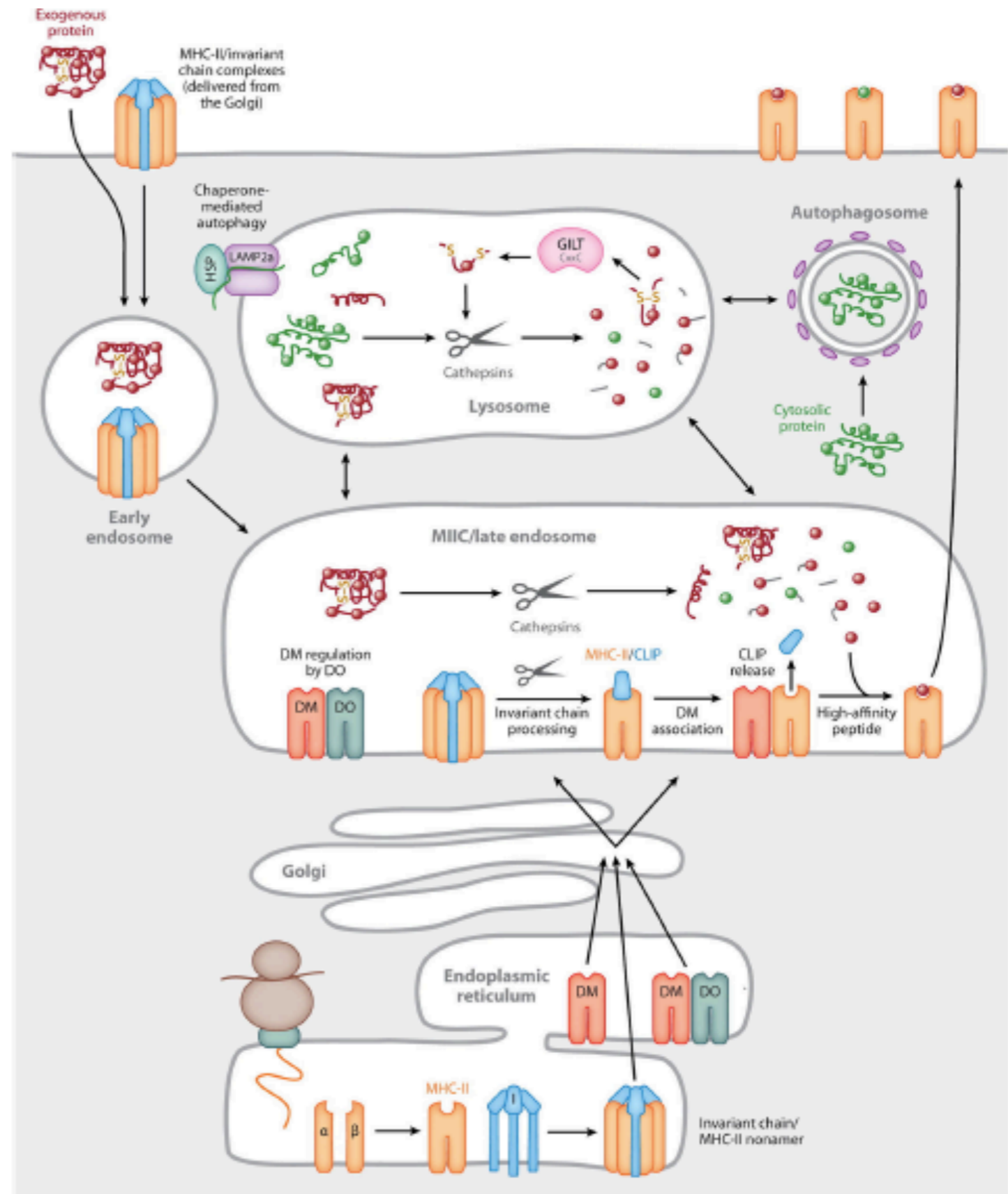
Associação dos peptídeos originados com MHC-II

Expressão dos complexos peptídeo-MHC na superfície da APC



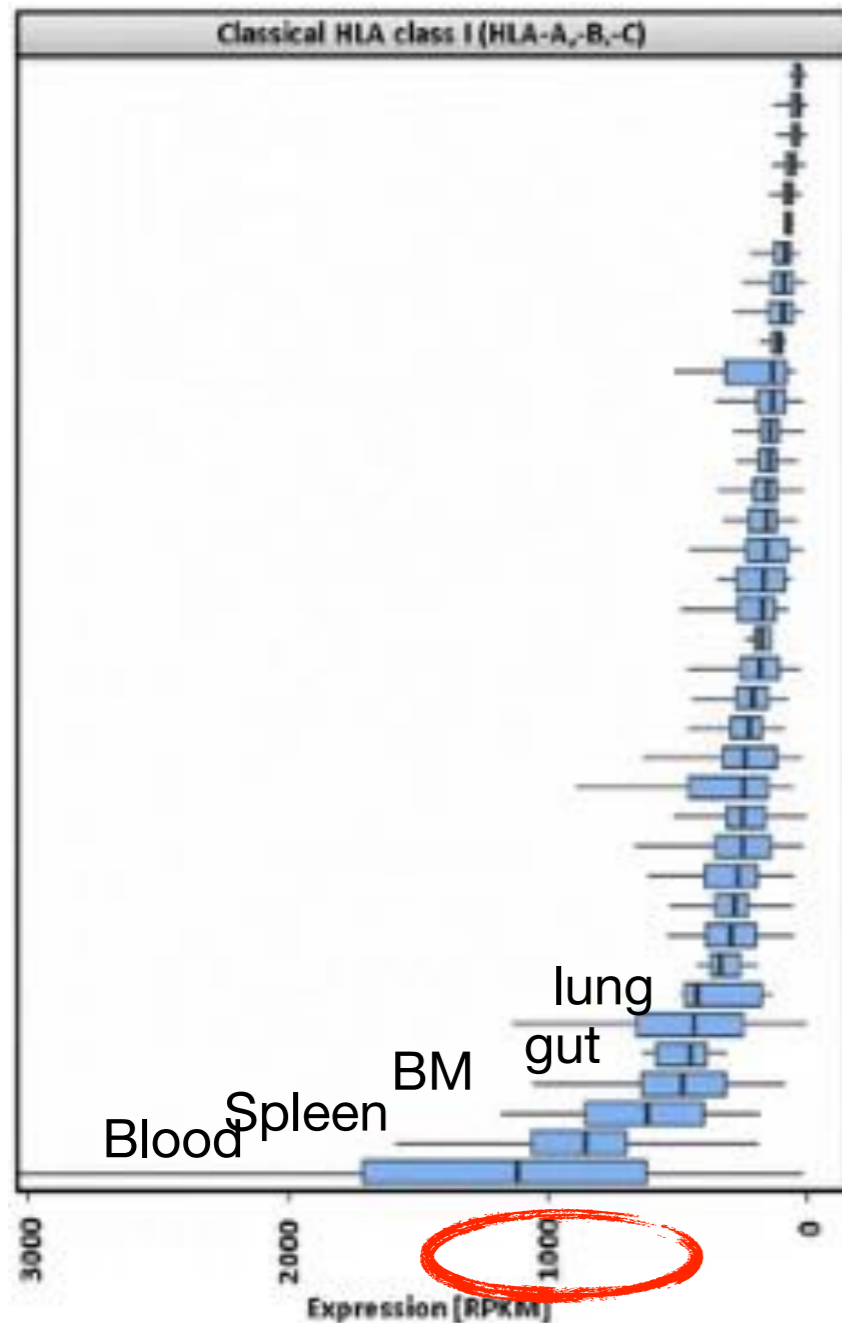
# Apresentação de Ag via MHC-II

Not only phagocytes or endocytosed molecules but also “self” proteins (autophagy)

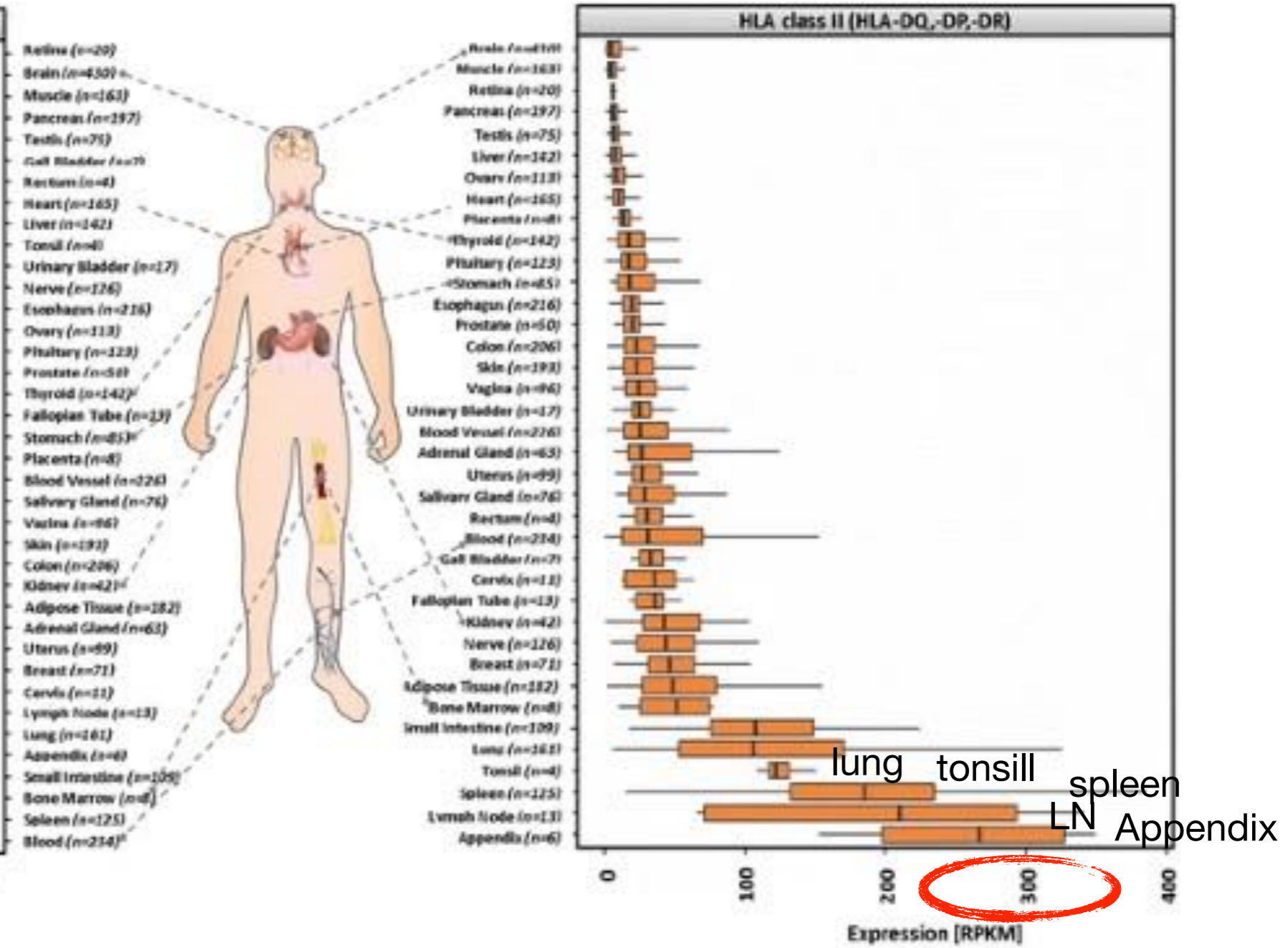


# Classical MHC genes expression

## MHC-I (ubiquitous)



## MHC-II (mainly APCs)





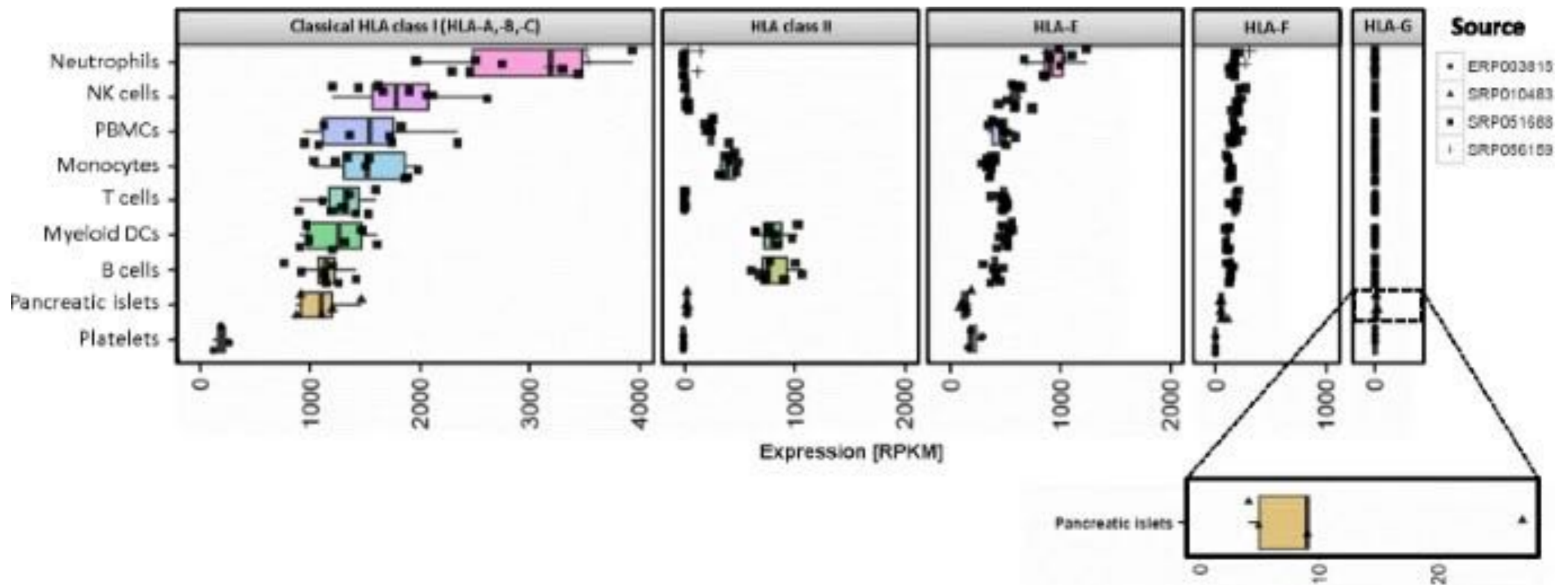
# MHC genes expression

**MHC-I**  
(ubiquitous)

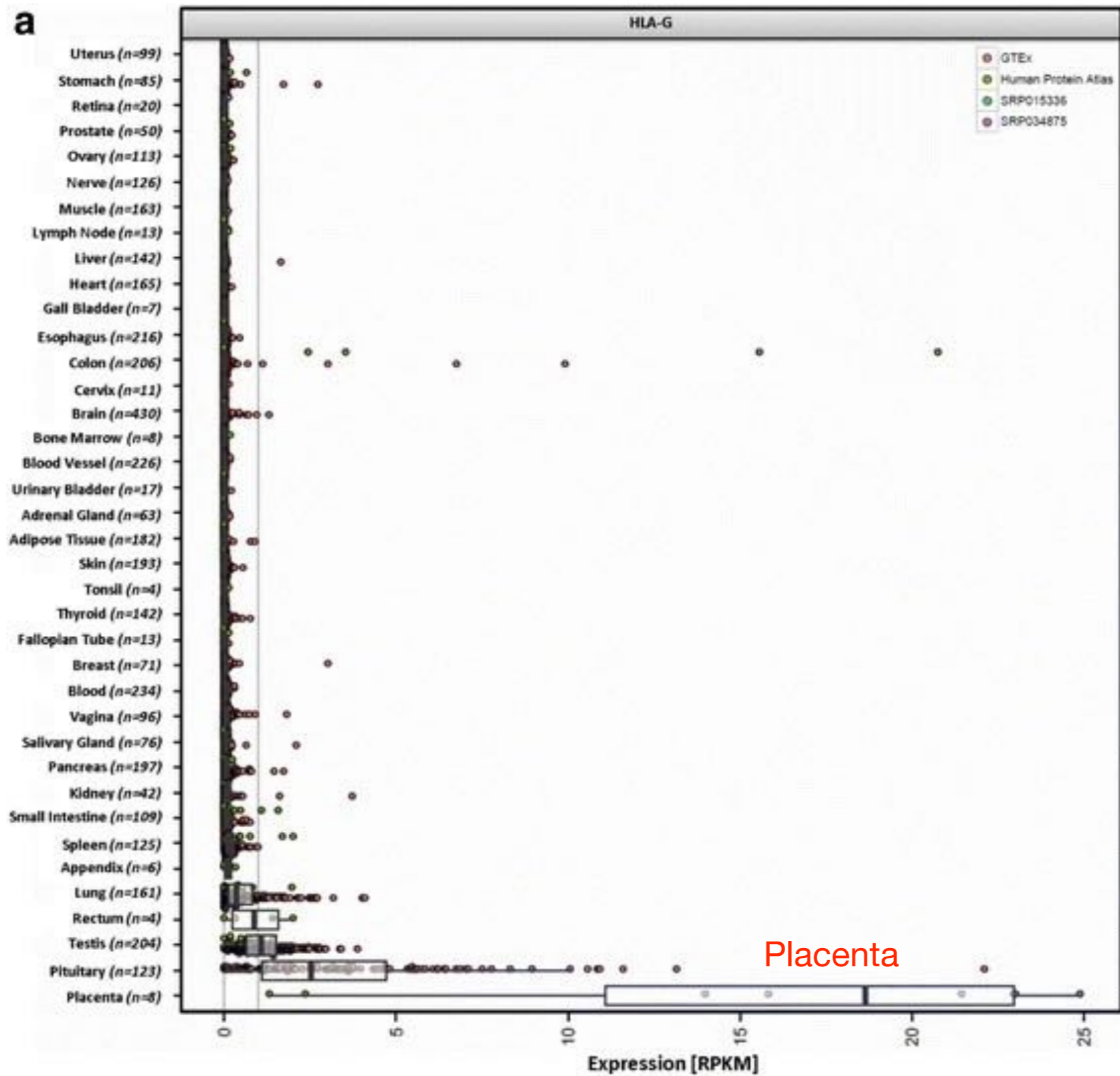
**MHC-II**  
(mainly APCs)

**HLA-E**  
(ubiquitous)

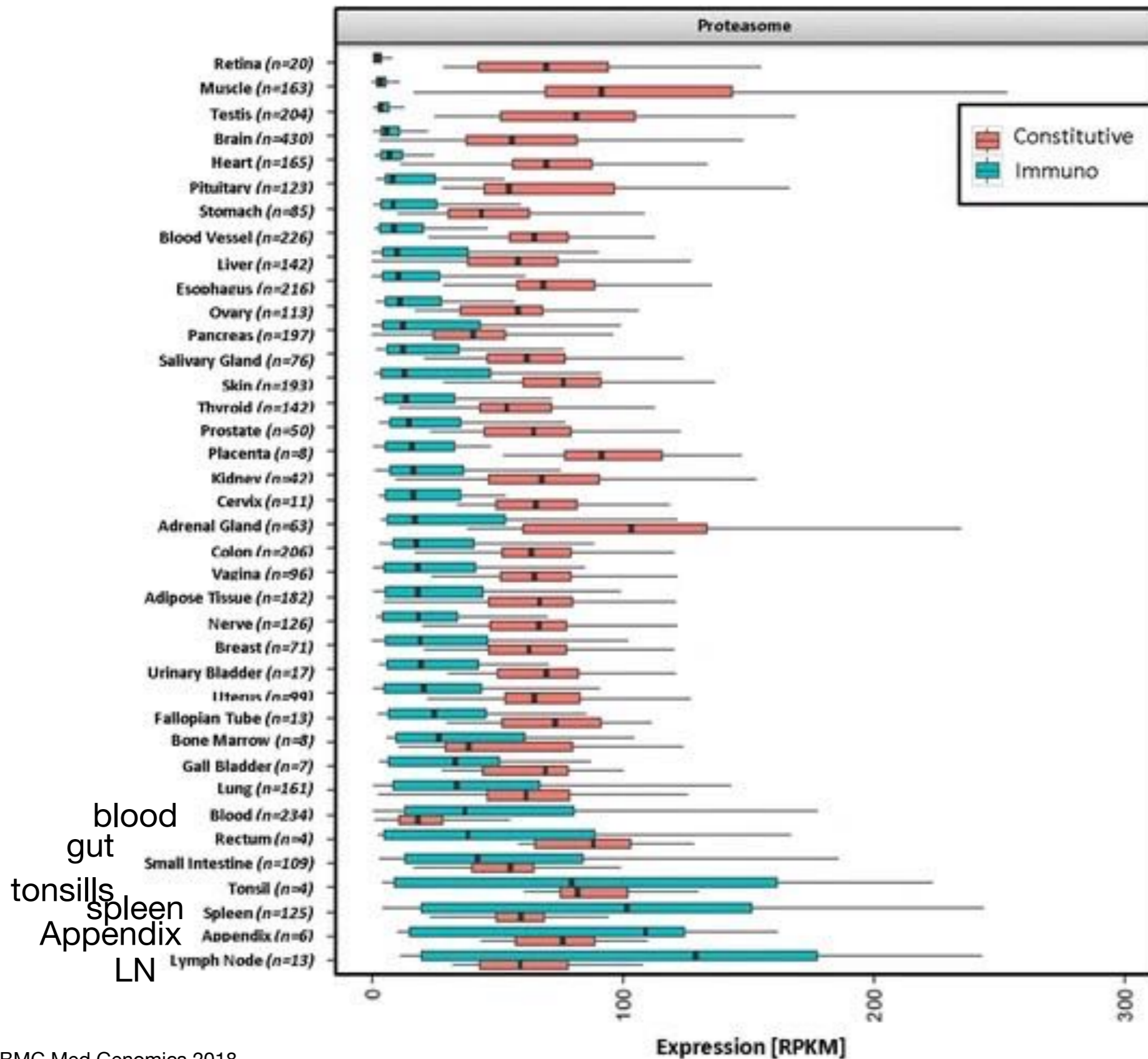
**HLA-G**  
(restricted)



# HLA-G expression



# Proteasome expression





# Funções dos MHC

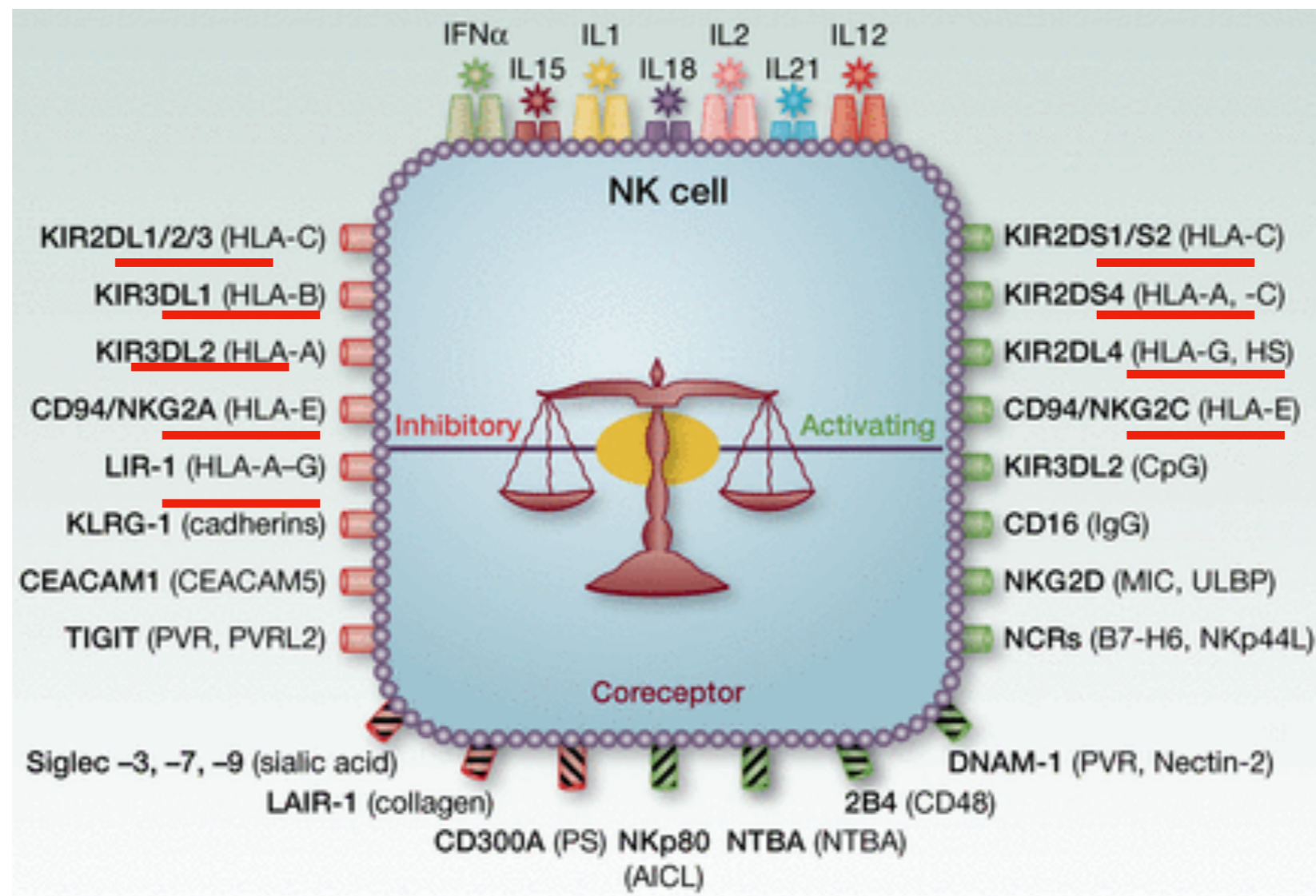
- ligandos de Receptores NK cells (imunovigilância)

ex: HLA-E + NKG2A/CD94 = inibição

HLA-E + NKG2C/CD94 = ativação

HLA-A + KIR3DL2 = inibição

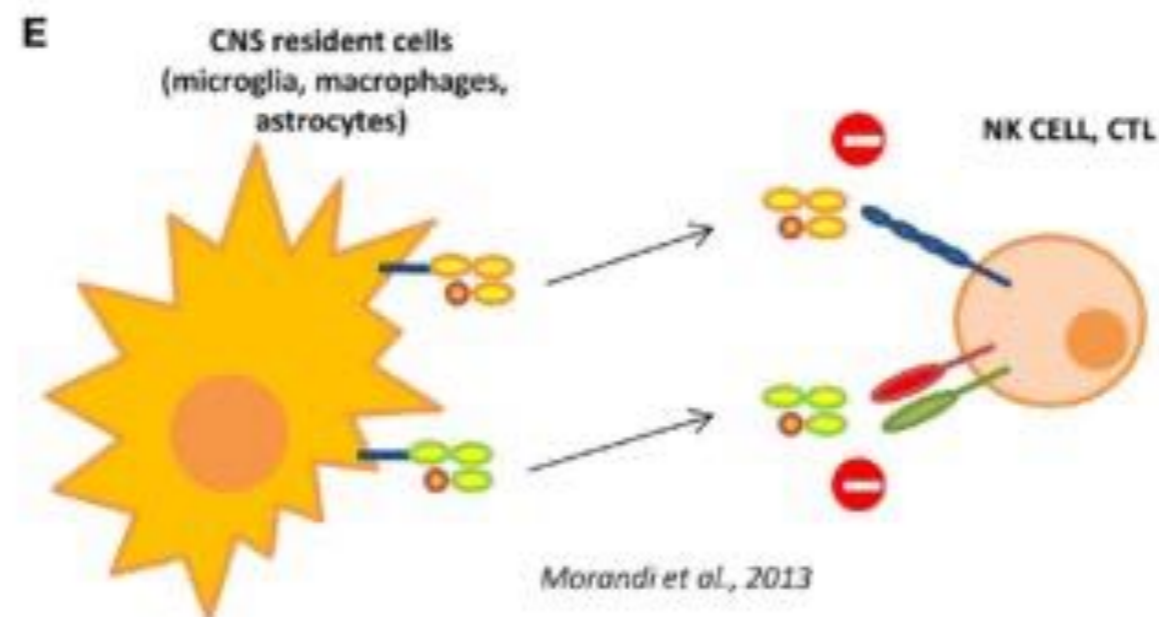
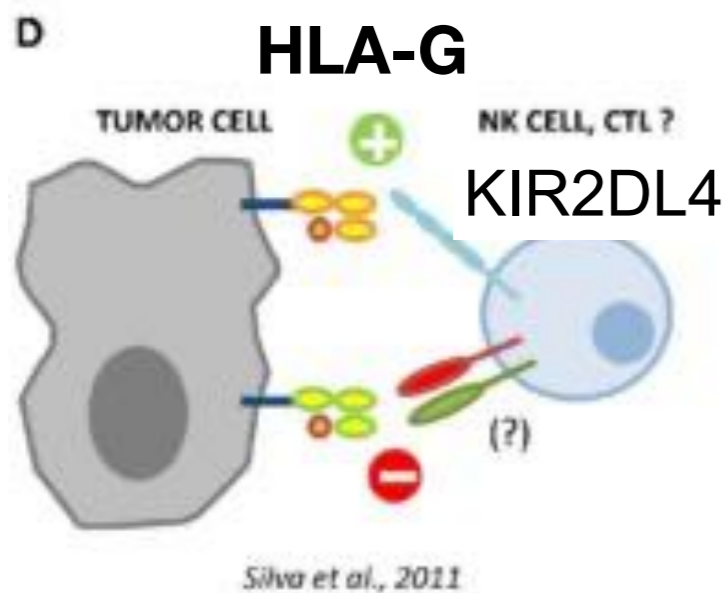
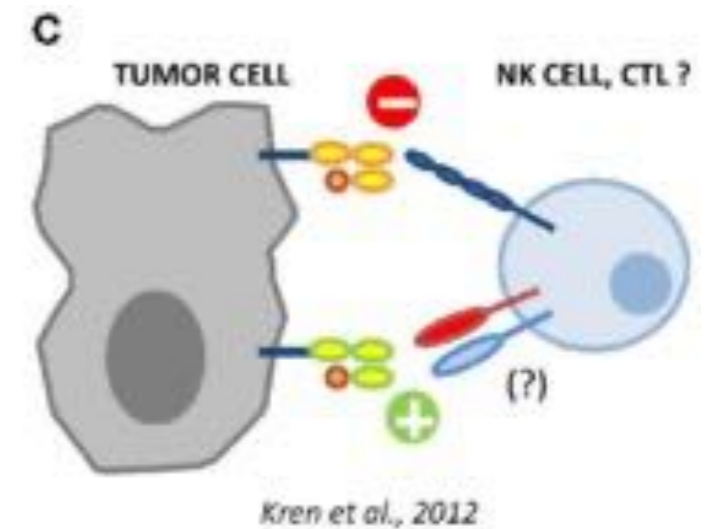
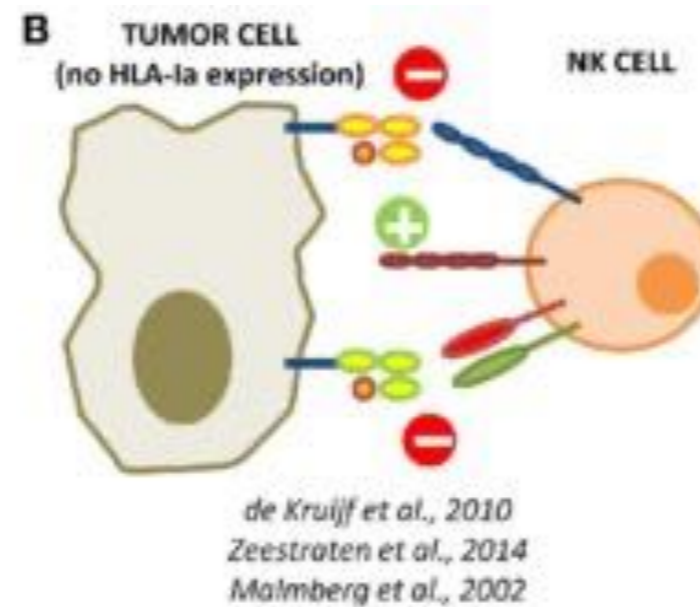
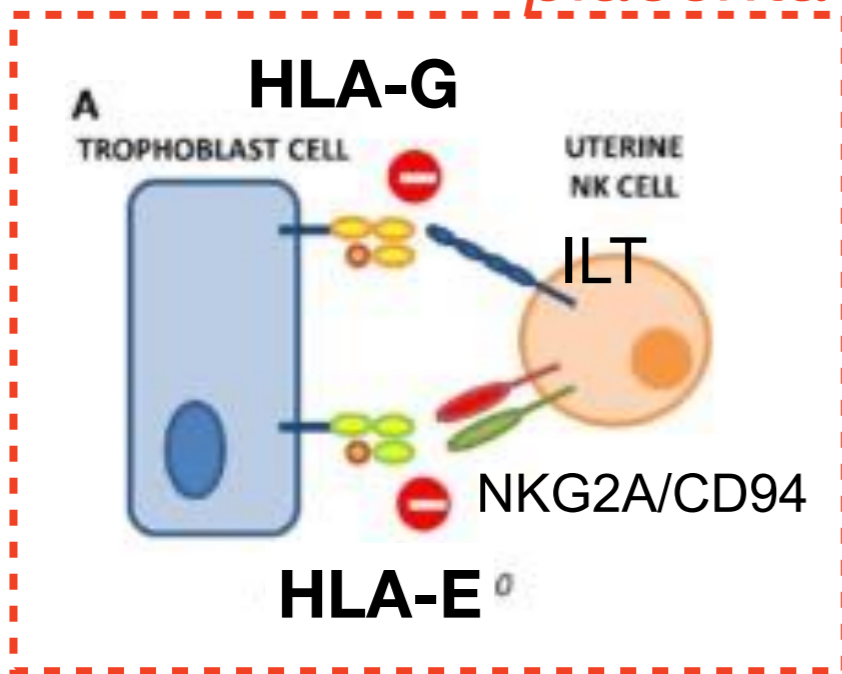
HLA-A + KIR2DS4 = ativação



# Funções dos MHC

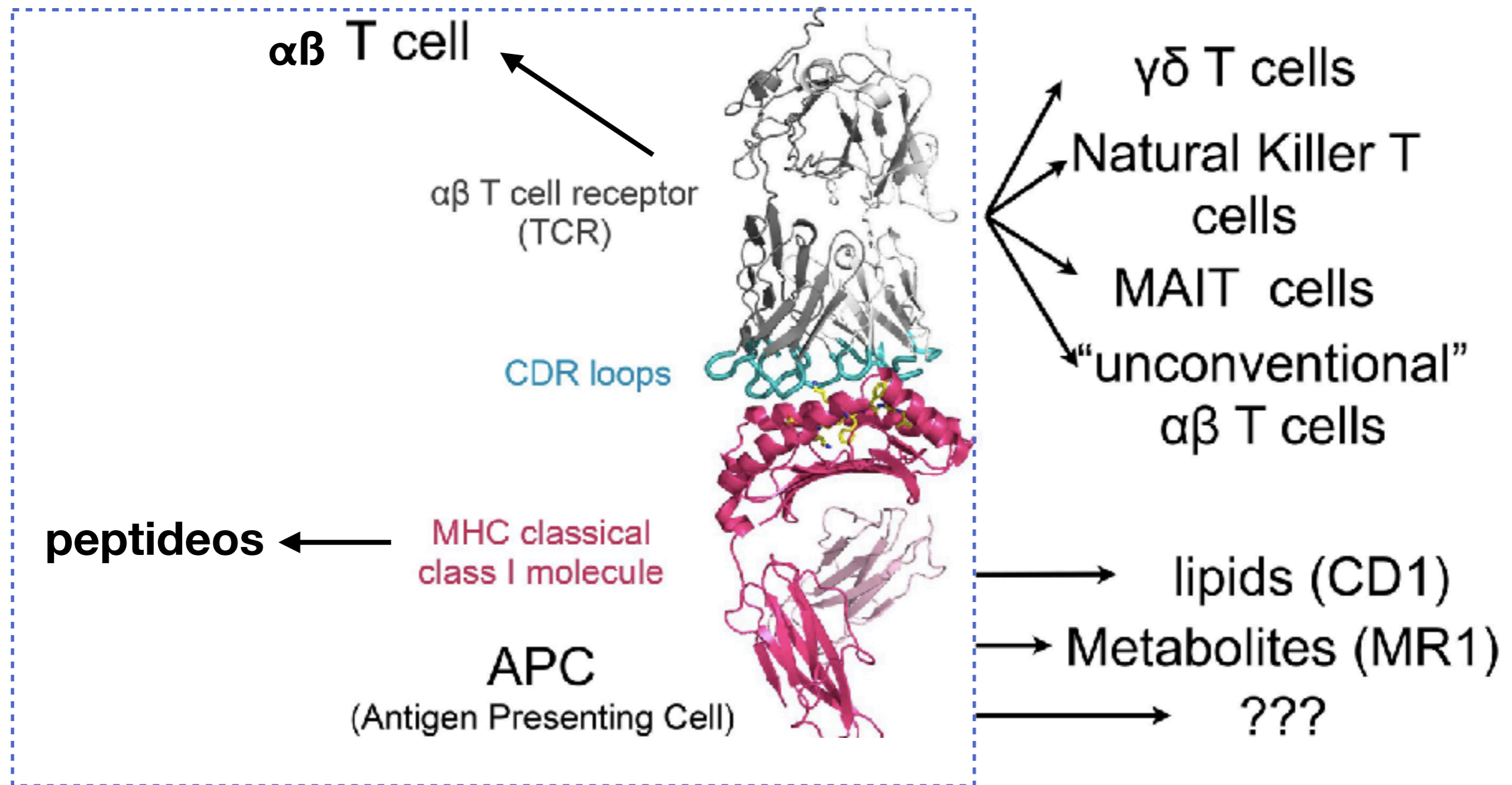
- imunoregulação

*placenta*



# Funções dos MHC

- apresentação de antígenos aos linfócitos T (**ligando do TCR**; ativação imunidade adquirida)



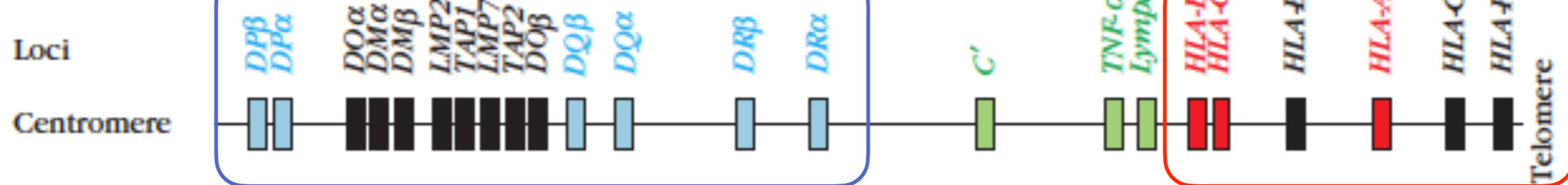


# Organização do locus MHC

HUMAN CHROMOSOME 6



Chr6



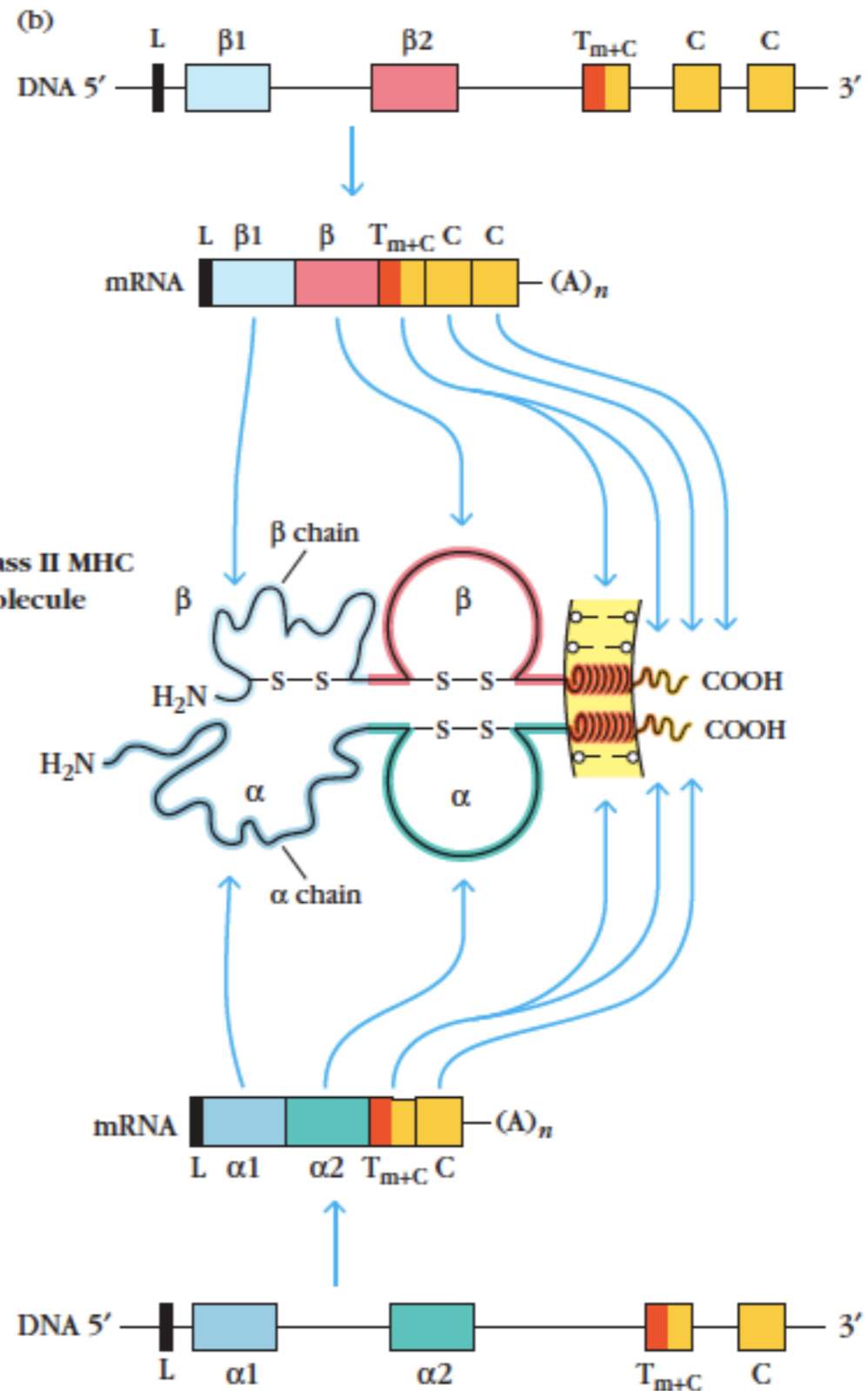
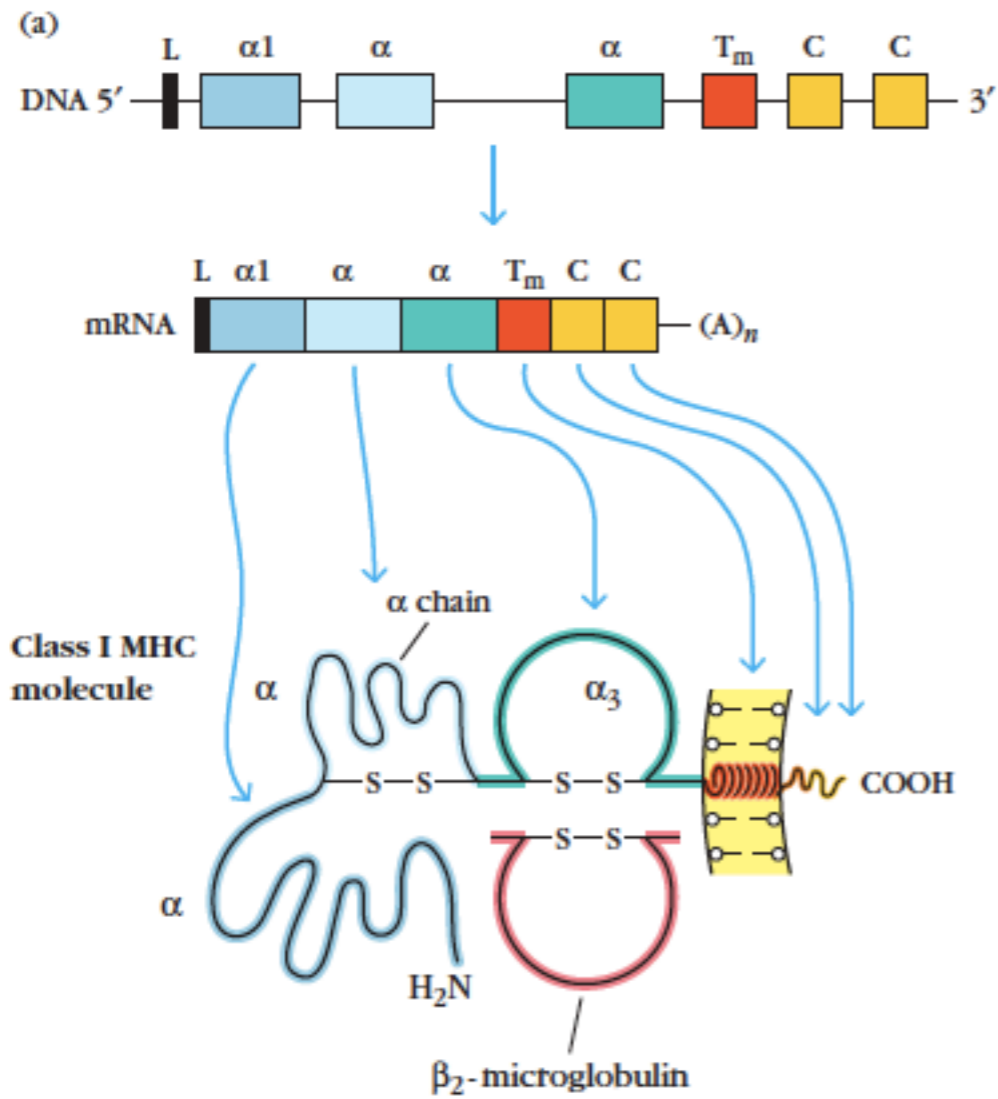
Human HLA complex

Complex	HLA									
MHC class	II			III		I				
Region	DP	DQ	DR	C4, C2, BF		B	C	A		
Gene products	DP αβ	DQ αβ	DR αβ	C' proteins		TNF-α	Lymphotoxin-α	HLA-B	HLA-C	HLA-A

# MHC/HLA

## MHC-I

## MHC-II



# Variabilidade do MHC

Several hundred different allelic variants of class I and II MHC molecules

## Regioes variaveis no sitio de ligacao para o peptideo

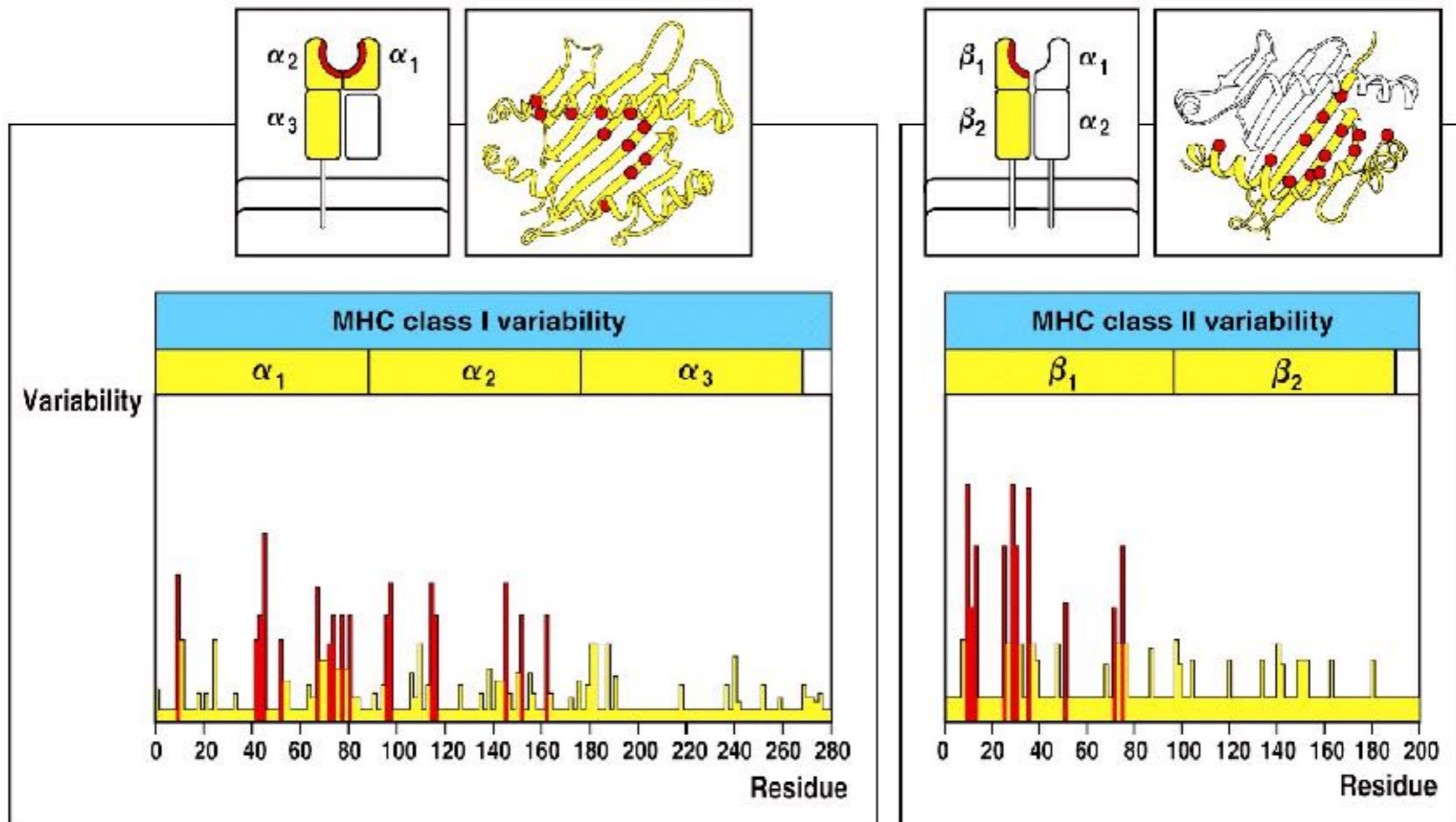


Figure 5-16 Immunobiology, 6/e. (© Garland Science 2005)



# Variabilidade do MHC

- Genes MHC são muito polimórficos e muitos alelos existem para cada gene
- Muito raro 2 indivíduos compartilhar os 2 sets de genes HLA (problemas transplante!)

**TABLE 8-3** Genetic diversity of MHC loci in the human population

MHC CLASS I	
HLA locus	Number of allotypes (proteins)
A	1448
B	1988
C	1119
E	3
F	4
G	16
MHC CLASS II	
HLA locus	Number of allotypes (proteins)
DMA	4
DMB	7
DOA	3
DOB	5
DPA1	17
DPB1	134
DQA1	47
DQB1	126
DRA	2
DRB1	860
DRB3	46
DRB4	8
DRB5	17

Source: Data obtained from <http://hla.alleles.org>, a Web site maintained by the HLA Informatics Group based at the Anthony Nolan Trust in the United Kingdom, with up-to-date information on the numbers of HLA alleles and proteins.

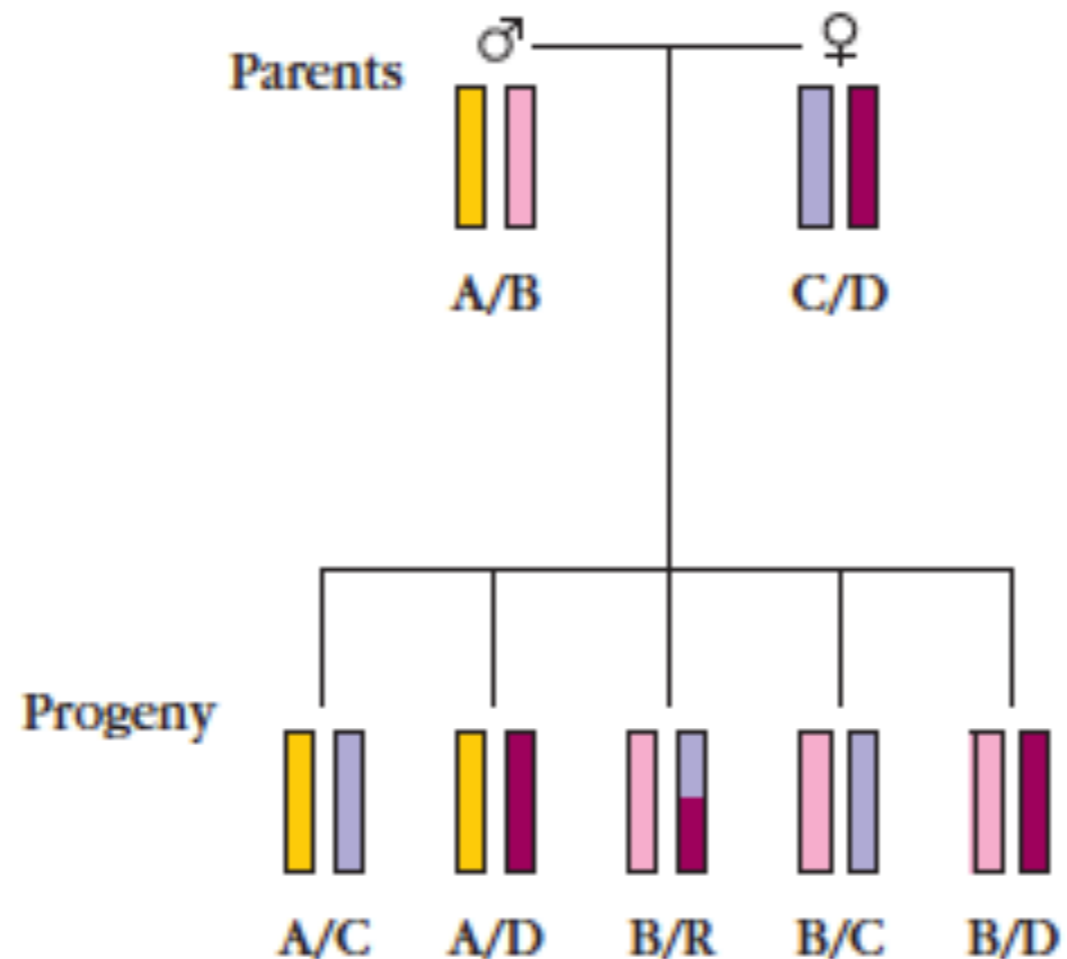
DQ Serotype	DR-DQ haplotype	DR B1	DQ A1	DQ B1	Freq % <sup>[5]</sup>	
DQ2	DR17-DQ2	0301	0501	0201	13.1	
	DR7-DQ2	0701	0201	0202	11.1	
DQ4	DR8-DQ4	0801	0401	0402	2.2	
		0101	0101	0501	9.1	
DQ5	DR1-DQ5	0102	0101	0501	1.4	
		0103	0101	0501	0.5	
		1001	0104	0501	0.7	
		1601	0102	0502	1.0	
		1401	0104	0503	2.0	
DQ6	DR15-DQ6	1502	0103	0601	0.7	
		1501	0102	0602	14.2	
		1301	0103	0603	5.6	
		DR13-DQ6	1302	0102	0604	0.7
		1302	0102	0609	3.4	
DQ7	DR11-DQ7	1101	0505	0301	5.6	
		1104	0505	0301	2.7	
		DR12-DQ7	1201	0505	0301	1.1
		DR13-DQ7	1303	0505	0301	0.7
DQ8	DR4-DQ7	0401	0303	0301	5.3	
		0407	0303	0301	0.9	
		0402	0301	0302	1.0	
DQ9	DR4-DQ8	0404	0301	0302	4.2	
		0401	0302	0302	0.7	
DQ9	DR7-DQ9	0701	0201	0303	3.7	
		0901	0302	0303	0.8	

# Genetica do MHC

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- Cada indivíduo é geralmente heterozigoto para cada locus (materno/paterno=)
- Diferentes genes vizinhos (haplotipos; LD)
- Apesar da baixa taxa de recombinação, essa contribui pela diversidade dos loci

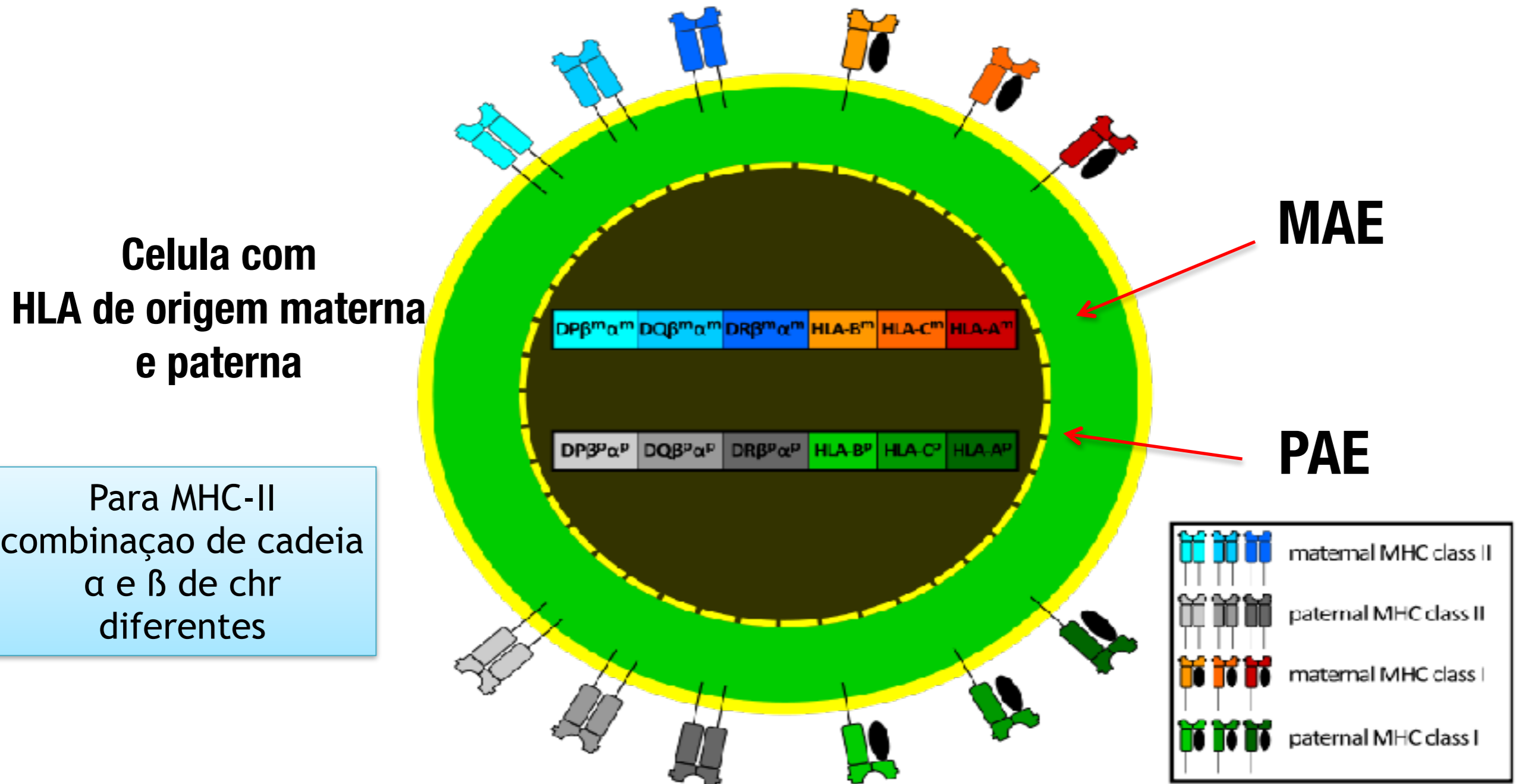
(c) Inheritance of HLA haplotypes in a typical human family



- *Genitor e filho compartilha apenas um haplotipo*
- *Ha uma chance de 25% que 2 irmaos herdem haplotipos compatíveis de HLA*

# Genetica do MHC

➤ Cada célula expressa diferentes genes MHC de modo co-dominante

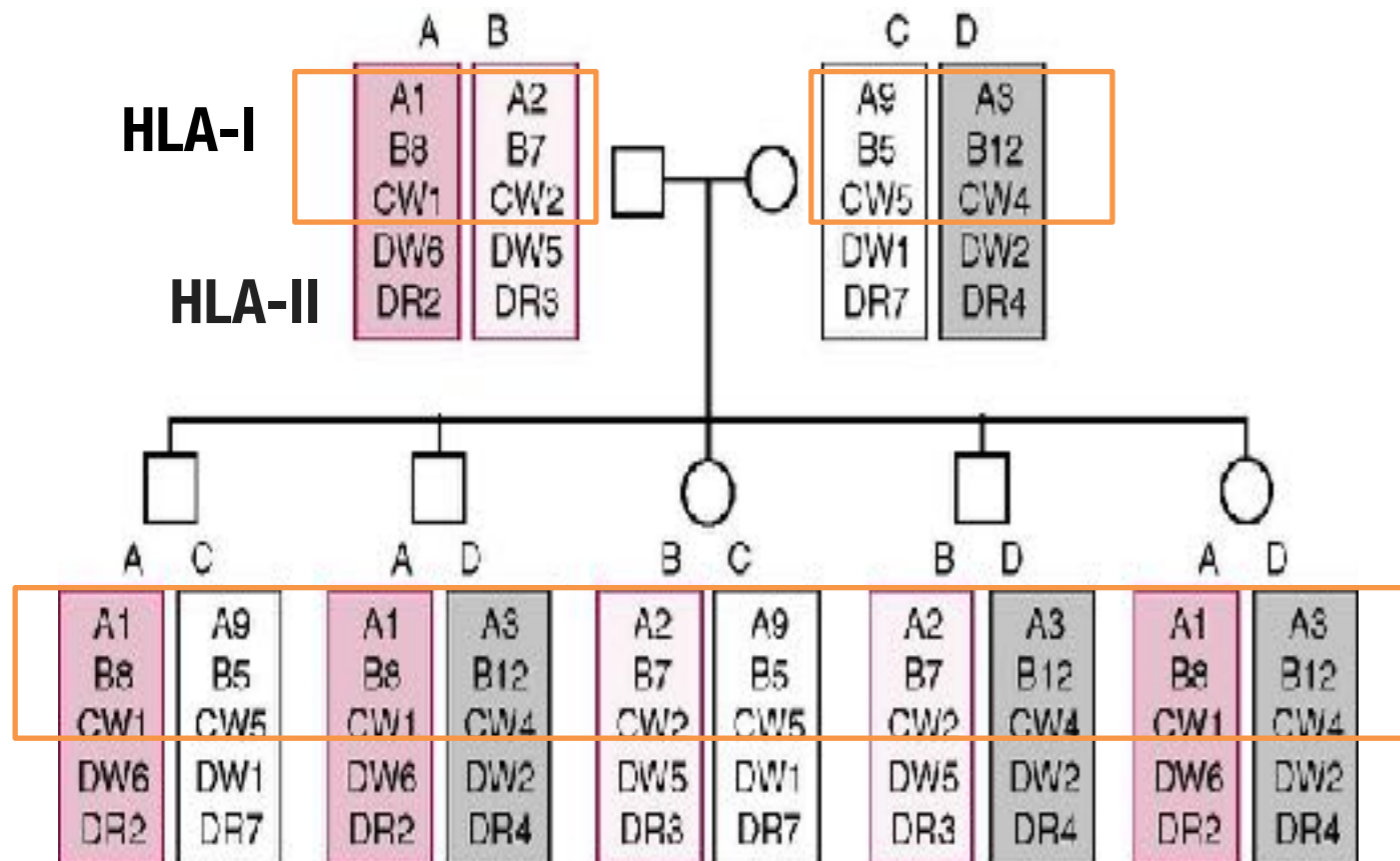




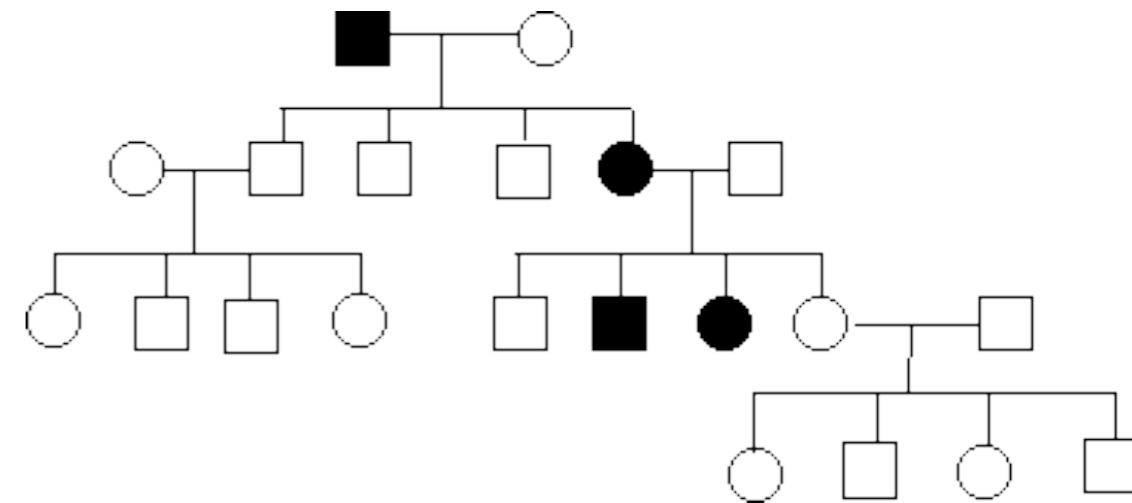
# Genetica do MHC

muitos genes codominantes x polimorfismo  
alelico x possibilidade de recombinação

MUITA VARIABILIDADE



*Herança de tipo mendeliano*



# Genetica do MHC

- ◇ possibilidade de ter diferentes combinações alelicas para apresentar de modo ótimo um maior número possível de Ag aos vários linfócitos T → **Vantagem seletiva**
- ◇ A maioria dos alelos são presentes com frequência diferente nas populações
- ◇ Isso indica que as mutações não são “neutras” mas que aconteceu alguma “seleção” favorecendo alguns alelos

**HLA-A\*0201** representa:

- 95% dos alelos A2 na população caucasiana
- 25% dos alelos A2 na população chinesa Han
- 3% dos alelos A2 população indiana

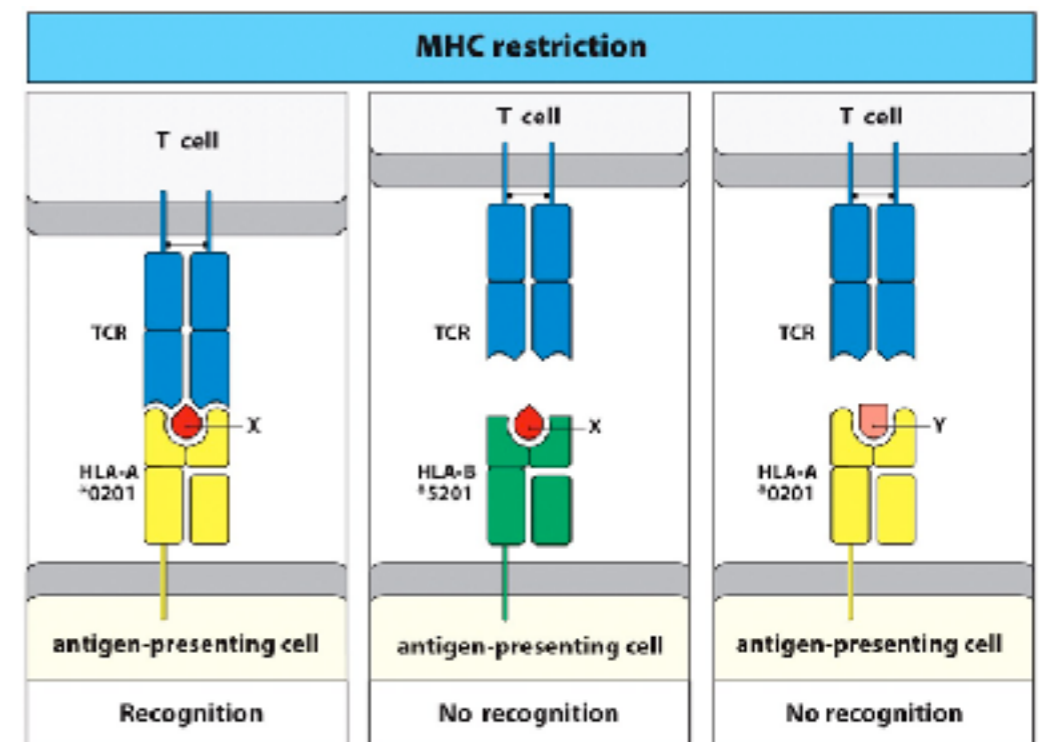
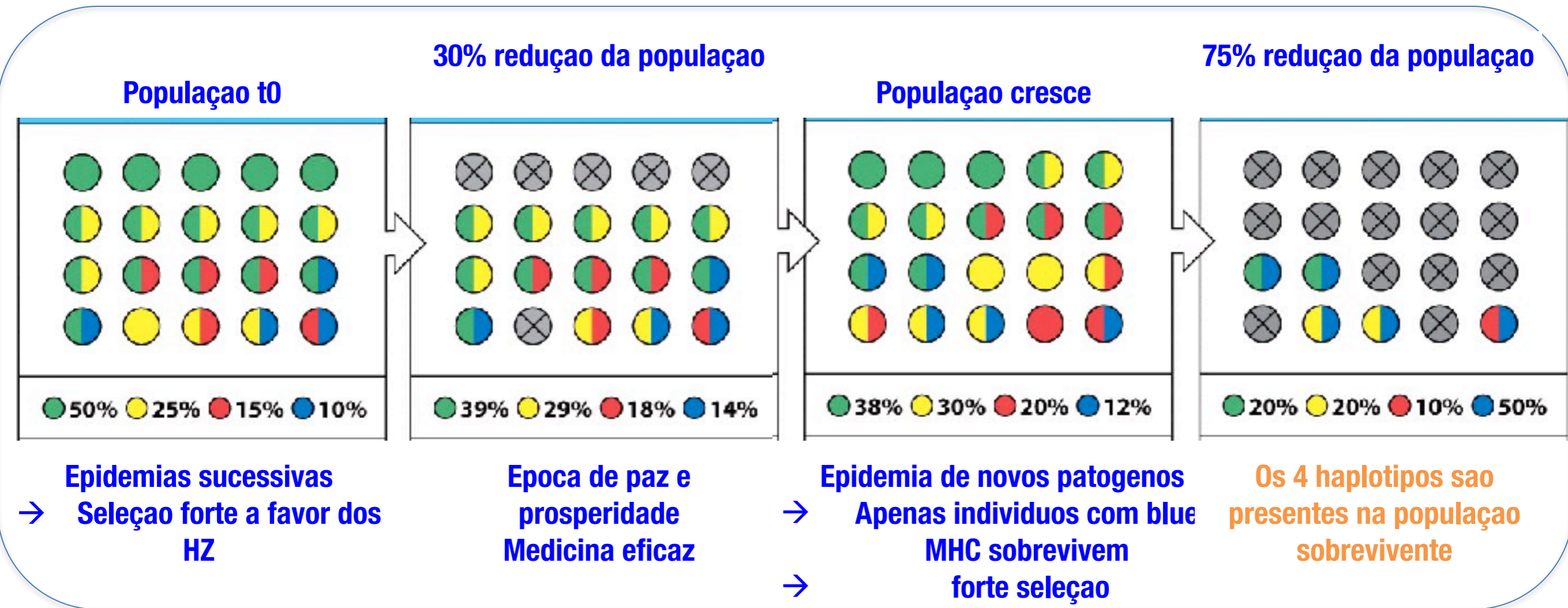


Figure 5.21 The Immune System, Dec. 10 Garland Science 2009

# Genetica do MHC





# MHC

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Several hundred different allelic variants of class I and II MHC molecules

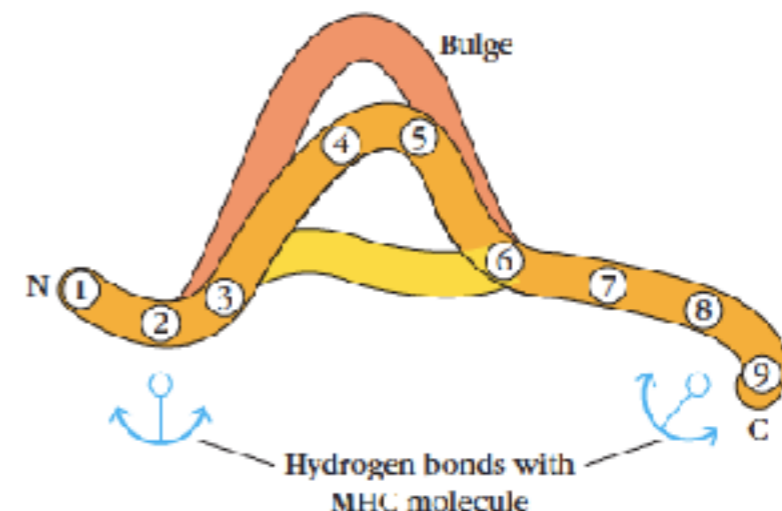
Any one individual express only a small number of these molecules

- up to 6 different class I molecules
- 12 or more different class II molecules

**Such a limited number of molecules for an enormous array of antigen peptides**

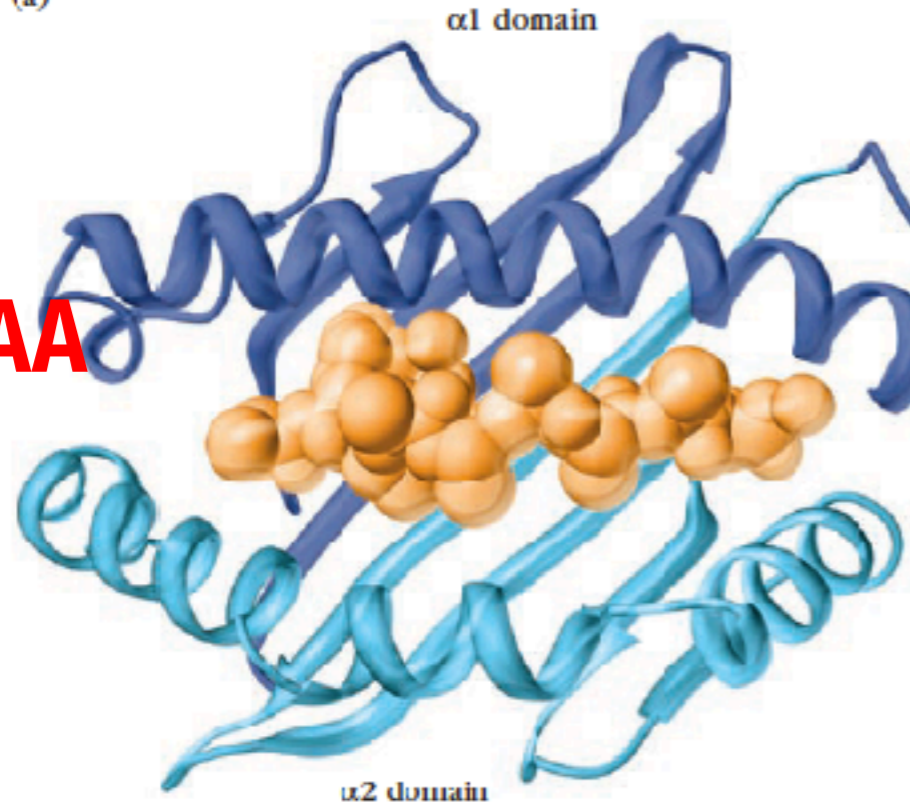
The peptide binding of MHC-I and -II molecules does not exhibit the specificity characteristic of antigen-binding by antibodies and TCR

A given MHC molecule can bind numerous different peptides, and some peptides can bind to several different MHC (promiscuous binding)

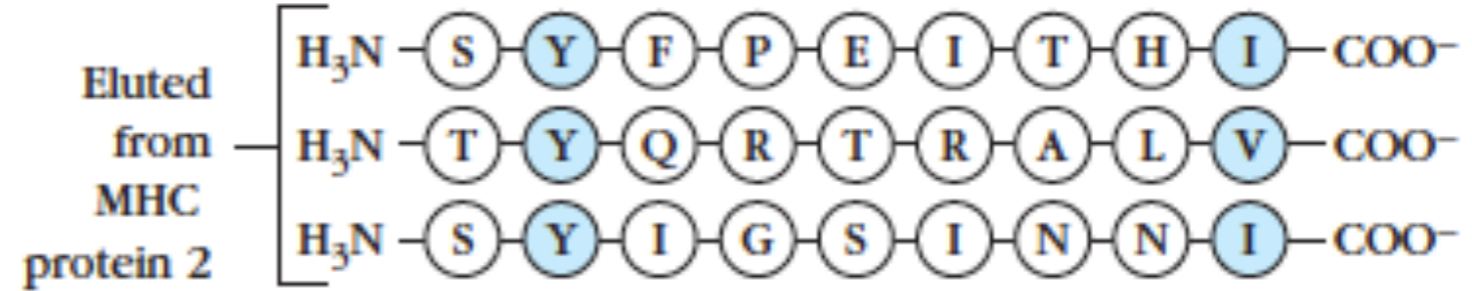


# MHC

(a)

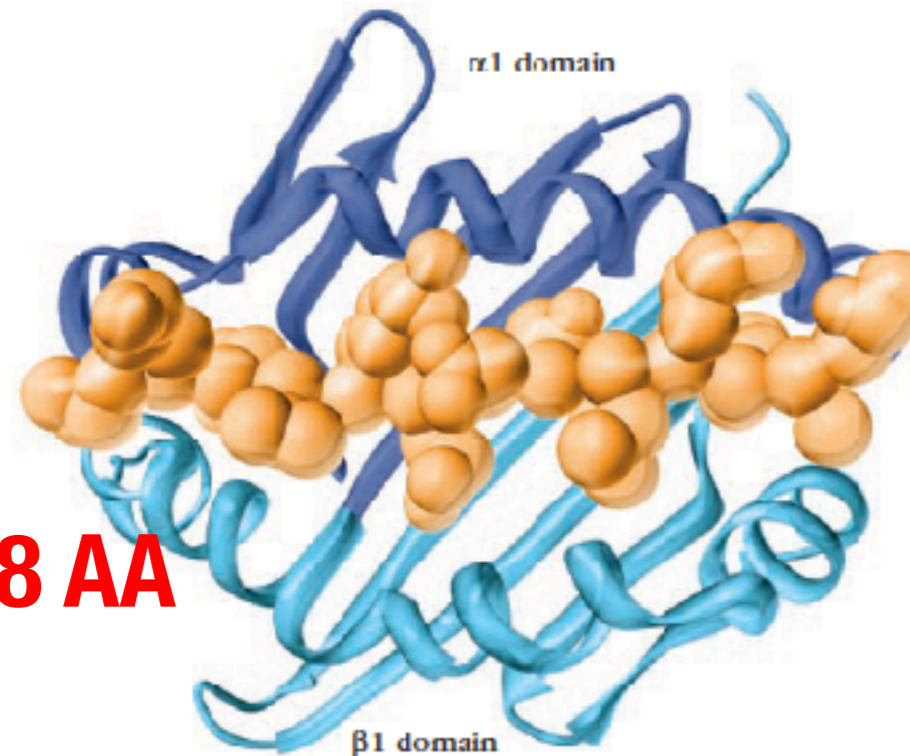


**Residuos ancora**



A = alanine	K = lysine	R = arginine
E = glutamic acid	L = leucine	S = serine
F = phenylalanine	N = asparagine	T = threonine
G = glycine	P = proline	V = valine
H = histidine	Q = glutamine	Y = tyrosine
I = isoleucine		

(b)



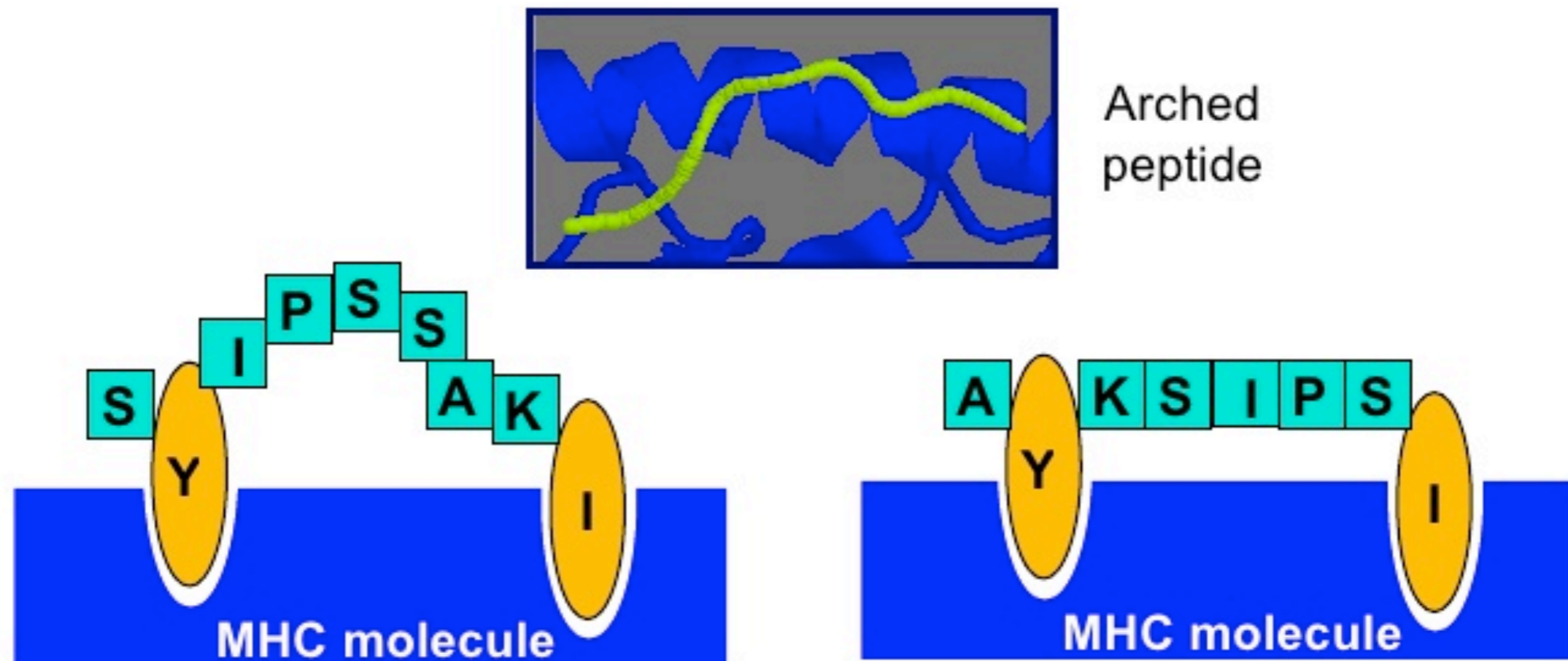
**13-18 AA**

**8-10 AA**

# MHC

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**MHC molecules can bind peptides of different length**



Complementary anchor residues & pockets provide the broad specificity of a particular type of MHC molecule for peptides

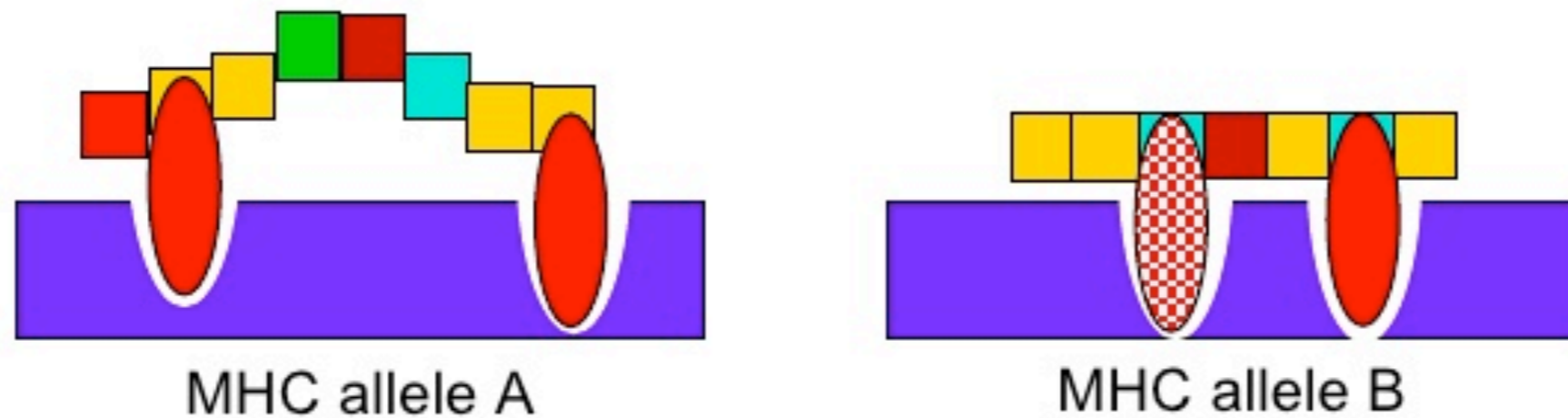
Peptide sequence between anchors can vary  
Number of amino acids between anchors can vary



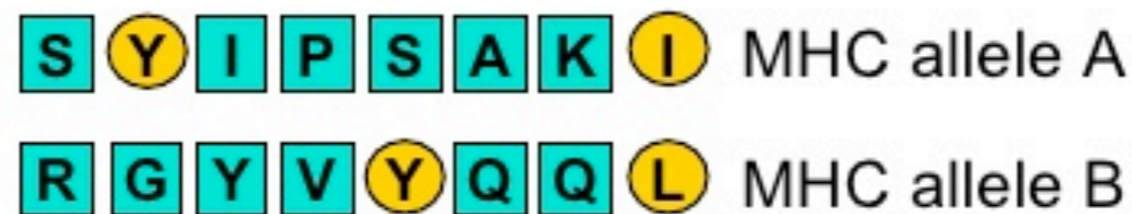
# MHC

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## Polymorphism in the MHC affects peptide antigen binding



Changes in the pockets, walls and floor of the peptide binding cleft alter peptide MHC interactions and determine which peptides bind.



Products of different MHC alleles bind a different repertoire of peptides

