

# Major Histocompatibility Complex

## PART-3

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# MHC molecule and Peptide interaction

- In human, various allelic form of class I & II MHC mol. have been identify.
- **Single Individual have only**
  - ✓ Up to 6 different class I MHC mol.
  - ✓ 12 or more different class II MHC mol.



**Limited number of MHC molecule able to present huge variants of antigenic peptide for T-Cell?**

**Yes**

**“MHC molecules can bind numerous variants of antigenic peptide”**

**Let us discuss how?**

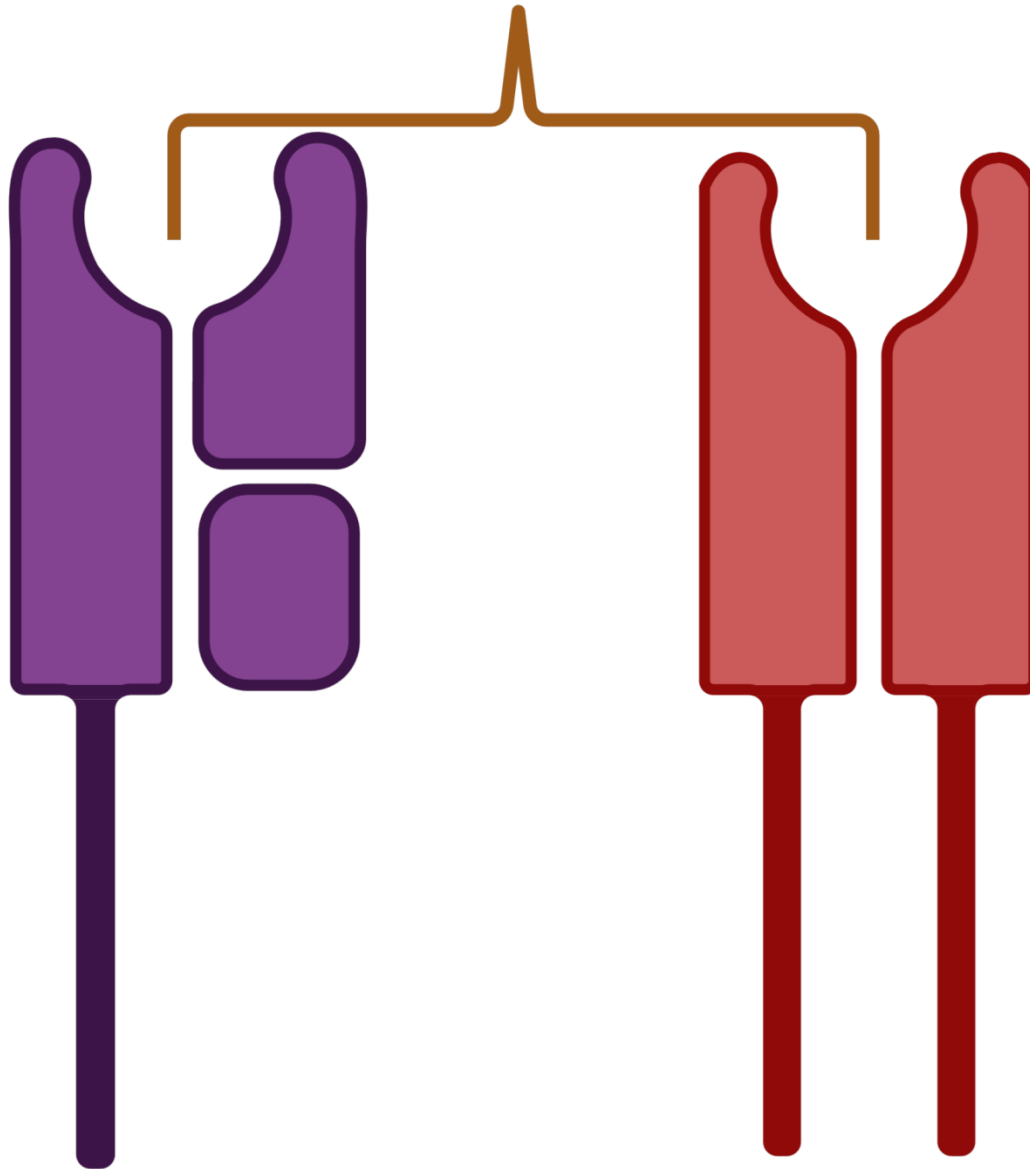
- **The broad specificity between antigenic peptides and MHC molecules is often referred to as “Promiscuous.”**
- **Association between antigenic peptide and MHC molecules are very stable under specific physiological conditions such as-**
  - **$K_d$  value range from approx.  $10^{-6}$  to  $10^{-10}$**

# Characteristic features of antigenic peptide-MHC molecule interaction

- Class I and Class II MHC molecules have a single antigenic peptide binding cleft.
- Generally accommodates one antigenic peptides at a time.
- Processed antigenic peptides must be compatible with MHC binding cleft.
- Compatible pair promote the interaction and activation of concerned immune cell.

- Peptide binding cleft is protected with a specific peptide (ERp57 in case of Class I MHC molecule) before encounter the antigenic peptide.
- Only few antigenic peptide-MHC molecules complex are capable to induce the immune response.
- MHC molecules express both self and non-self (antigenic) peptide on the surface of antigen presenting cells.
- T-cells capable to discriminate self and non-self (antigenic) peptide.
- After recognition, T-cell respond accordingly.

Single antigenic peptide binding cleft



Class I MHC Molecule

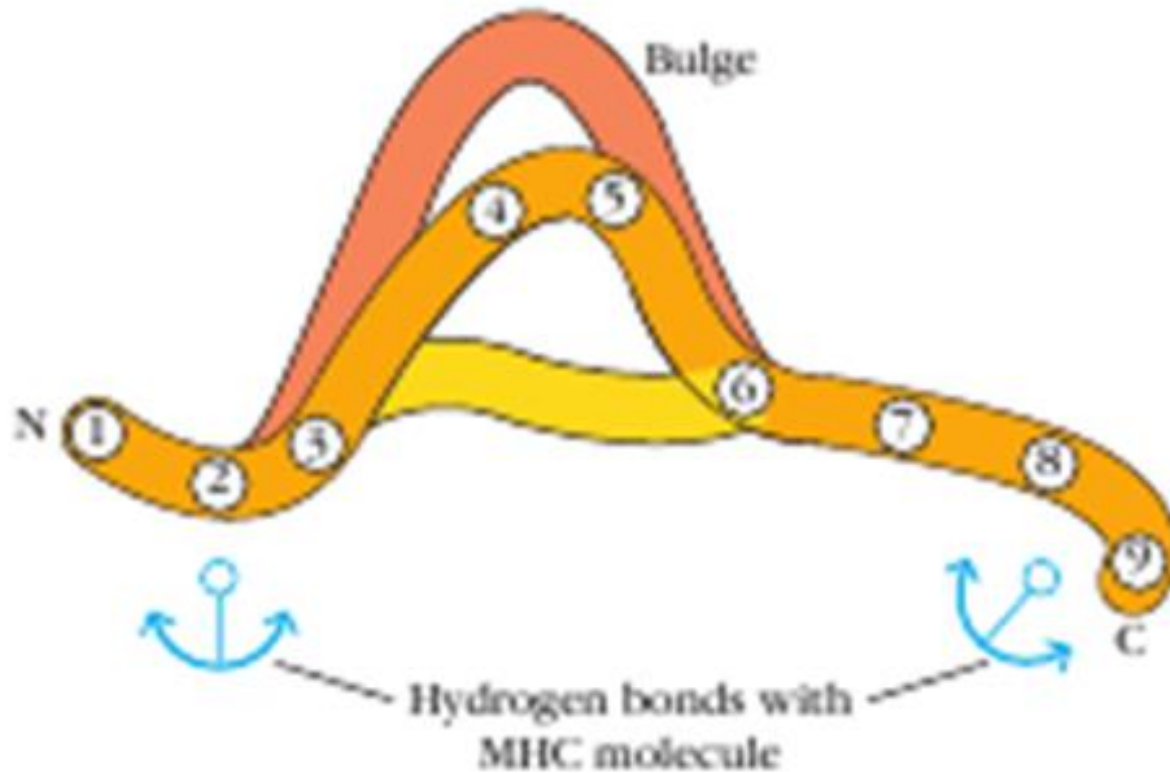
Class II MHC Molecule

# Class I MHC Molecule-Peptide Interaction

- Antigenic peptides are bind with class I molecule and expressed on the surface of APCs for CD8<sup>+</sup> T lymphocyte cell.
- Antigenic peptides are derived from endogenous altered antigenic proteins.
- Interaction between antigenic peptides and class I MHC molecules occur inside the Rough Endoplasmic Reticulum.
- This process referred as endogenous processing pathway.

- Every Class I MHC molecule have slightly different rule for peptide bindings
- All nucleated cells are able to express approx.  $10^{-5}$  copies of class I MHC molecule and each with its own promiscuity rules.
- Inherited Class I MHC allele will determine specificity of antigenic peptide with class I MHC molecule.
- Antigenic peptides are 8-10 AAs (9 AAs are common) long.





## Conformation of antigenic peptide of Class I MHC molecule

**Reference:** Kuby –Immunology; 7<sup>th</sup> Edition by Judith A. Owen, Jenni Punt, Sharon A. Stranford and Patricia P. Jones; Chapter-8: The Major Histocompatibility Complex and Antigen Presentation; Page: 267

- Specific amino acid residues are found at particular site of the antigenic peptide.
- Class I MHC molecule able to bind with thousands of different antigenic peptide due to presence of the similar amino acid residues at specific site.
- Specific sites amino acids (antigenic peptide) anchor the cleft of MHC molecule, they are called anchor residues.

- Allelic variants of class I MHC molecules have different amino acid residues in the antigenic peptide binding cleft.
- Class I MHC molecule contains-
  - ✓ Carboxyl-terminal anchor (at position 9) - contain hydrophobic residues such as leucine and isoleucine.
  - ✓ Amino-terminal anchor (at position 2<sup>nd</sup> or 2<sup>nd</sup> and 3<sup>rd</sup>).

- **Carboxyl-terminal anchor and Amino-terminal anchor firmly bind with peptide binding cleft.**
- **Amino acid residues between Carboxyl-terminal anchor and Amino-terminal anchor form an arch.**
- **Centre of arch (antigenic peptide) is bulge out from the cleft of MHC molecule .**
- **Centre of arch is interact with TCR.**

# Class II MHC Molecule-Peptide Interaction

- Antigenic peptides are bind with class II molecule and expressed on the surface of APCs for **CD4<sup>+</sup> T lymphocyte cell**.
- Class II MHC molecules also bind diverse antigenic peptide.
- Antigenic peptides are derived from exogenous protein (**either self or nonself**).
- Exogenous antigenic peptides are processed through **exogenous processing pathway**.

- **Antigenic peptides of class II MHC molecule derived from**
  - ✓ **Self membrane bound protein or**
  - ✓ **Foreign protein**
- **These protein uptake by cell through phagocytosis**
- **Uptake protein further processed through exogenous pathway.**
- **Antigenic peptides of class II MHC molecule contain 13 -18 amino acid residues.**

- Antigenic peptides binding cleft of class II MHC molecule **open at both end**.
- Due to open at both end, accommodate longer antigenic peptide than class I MHC molecule.
- Central core of 13 amino acid residues determine the ability of antigenic peptide to bind class II MHC molecules.
- Unlike class I antigenic peptide, Class II antigenic peptides have **internal conserved sequence motifs**.

- **Antigenic peptide of class II molecule bind inside the cleft with hydrogen bond.**
- **Hydrogen bond distributed throughout the binding site.**
- **Internal conserved sequence motifs contain 7-10 amino acids provide major contact point.**



- Major contact point contains
  - ✓ Amino terminus -aromatic or hydrophobic amino acid residues
  - ✓ Middle portion –Hydrophobic residues
  - ✓ Carboxyl terminus- Hydrophobic residues
- Many antigenic peptide have proline amino acid at second position and carboxyl terminal end.
- Proline amino acid contributes to peptide binding promiscuity.

**Thanks**