

Major Histocompatibility Complex

PART- 4

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Pathways for processing antigen

Two pathways are proposed-

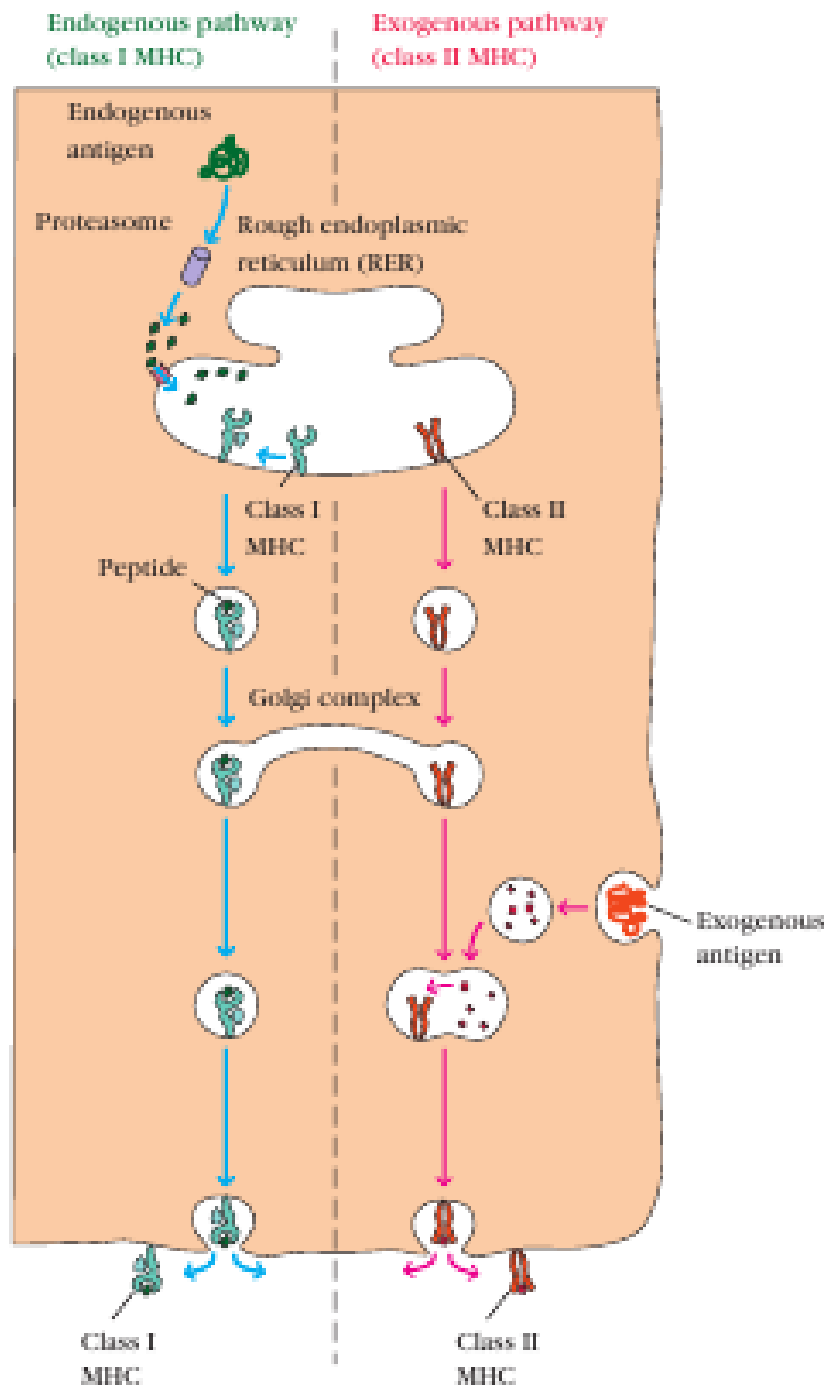
- 1. Cytosolic or Endogenous pathway**
- 2. Exogenous pathway**

Cytosolic or Endogenous Pathway

- Antigenic peptides for class I MHC molecule are processed through cytosolic or endogenous pathway.
- For this pathway, antigen are generated within cell.
- **Proteasome** (A proteolytic system): degraded irregular/ antigenic intracellular proteins into short peptide.

Overview of endogenous and exogenous pathways for the antigenic peptides

Reference: Kuby – Immunology; 7th Edition by Judith A. Owen, Jenni Punt, Sharon A. Stranford and Patricia P. Jones; Chapter-8: The Major Histocompatibility Complex and Antigen Presentation; Page: 284



General pathway of protein degradation

PROTEASOME

- Found in all eukaryotic cell
- Degrade proteins into short peptides
- Size- 20S
- Composed of 14 subunits
- Structure: barrel like symmetrical ring
- Target protein degrade in proteasome when ubiquitin (small protein) attached.

PROTEASOME CONT.....

- **Proteasome complex consists of**
 - **20S base**
 - **19S regulatory component**
- **Central hollow of proteasome degrade ubiquitin-protein complex.**
- **Cleavage of peptide bond- ATP dependent.**
- **Break down of protein through proteasome complex is also utilize by the immune system.**
- **Degraded antigenic peptides are displayed by APCs through class I MHC molecules.**

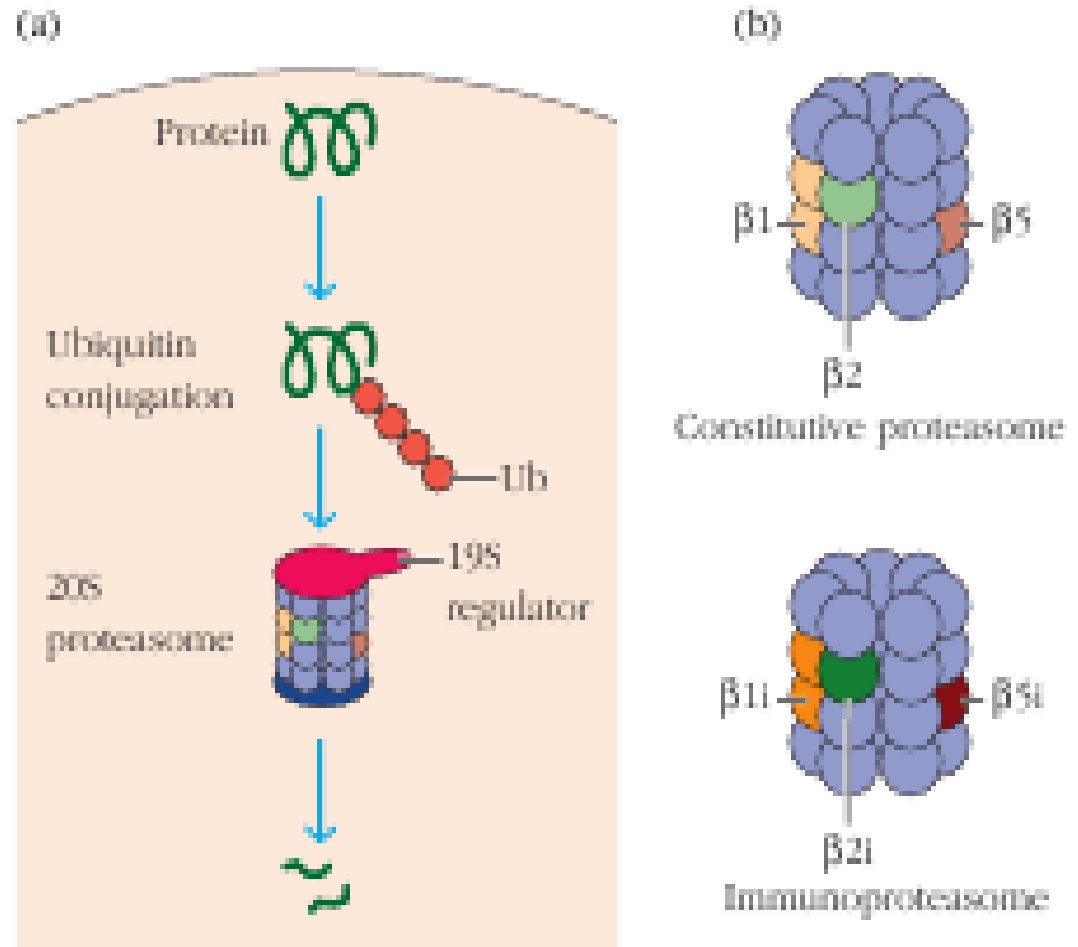
Specific pathway of protein degradation

IMMUNOPROTEASOME

- Found in infected APCs
- Size equal to proteasome
- Its unique components are induced by
 - Interferon- γ and
 - TNF- α
- Interferon- γ and TNF- α act as signal molecule for LMP2 and LMP7 genes
- Product of LMP2 and LMP7 genes (replacement catalytic protein subunit) convert standard proteasome into immunoproteasomes.
- Immunoproteasomes work more efficiently than standard proteasome.

PROTEASOME &

IMMUNOPROTEASOME



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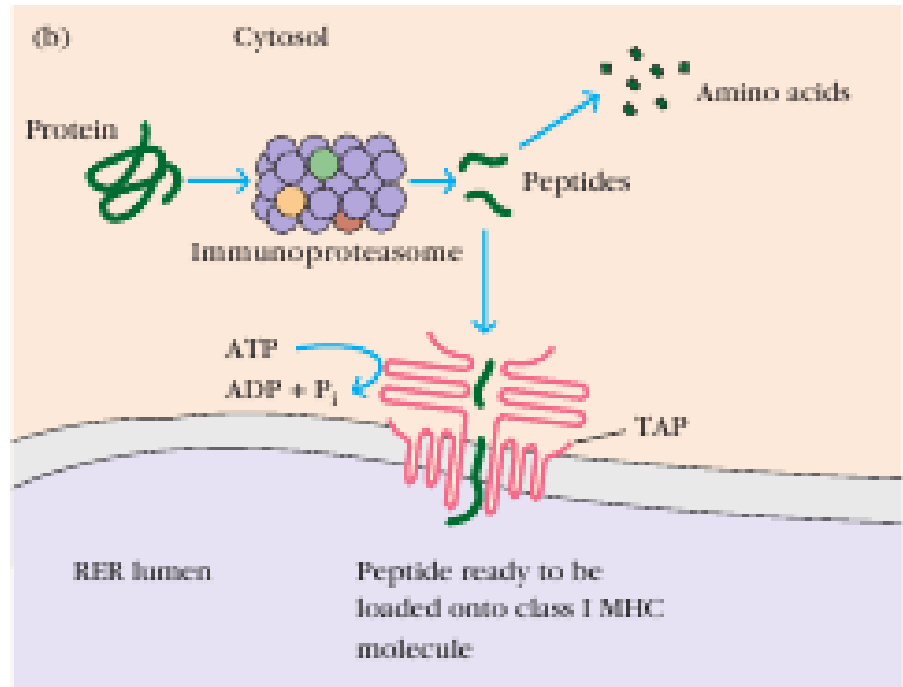
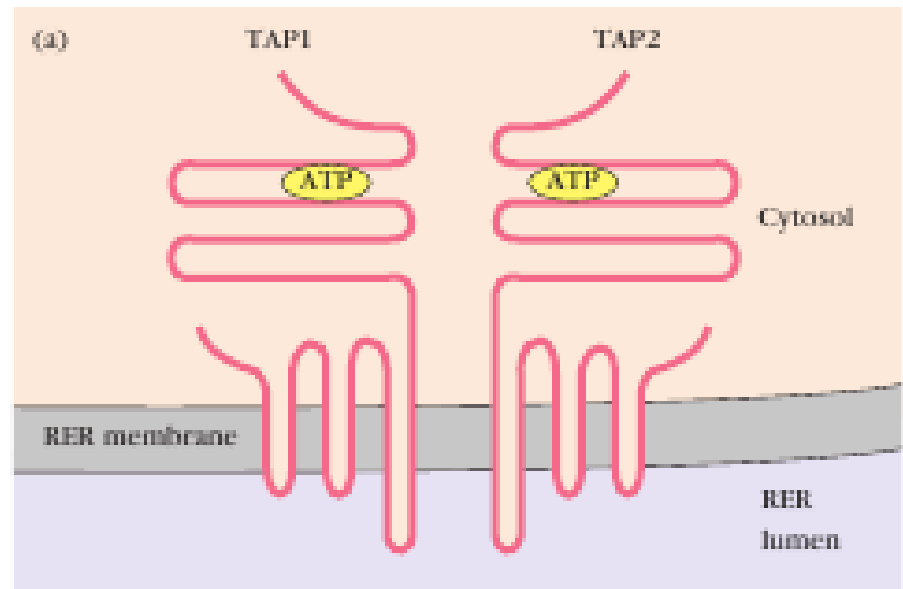
Transporter Protein- TAP

- TAP-membrane spanning heterodimer
- Consists of two protein
 - TAP1 and
 - TAP2
- Have two domain
 - RER lumen domain
 - Cytosolic domain-ATP binding
- Antigenic peptides formed by proteasome are translocated by TAP into RER lumen.
- Hydrolysis of ATP required during transportation.

- TAP show affinity for antigenic peptide (8-16 AAs).
- ERAP (Endoplasmic reticulum aminopeptidase)
 - Found in ER lumen
 - Trimmed longer peptide
- TAP1 and TAP2 protein synthesized by
 - TAP1 and TAP2 gene
 - Mapped within class II MHC region
 - Allelic form exist in population
- Deficiency of TAP leads to disease syndrome.

Transporter Protein- TAP and processing of antigenic peptides

Reference: Kuby –Immunology; 7th Edition by Judith A. Owen, Jenni Punt, Sharon A. Stranford and Patricia P. Jones; Chapter-8: The Major Histocompatibility Complex and Antigen Presentation; Page: 286





Deficiencies in TAP Can Lead to Bare Lymphocyte Syndrome

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Role of molecular chaperone in assembly of Class I MHC Molecule

- Molecular Chaperones are-
- Calnexin
 - found in ER membrane
 - Help in folding of class I α chain with ERp57 protein (enzymatic activity)
- Calnexin release after assembling of α chain and β microglobulin.

- **Molecular chaperone-**

- Calreticulin and
 - Tapasin
- } immediately associated with class I MHC molecule after release calnexin

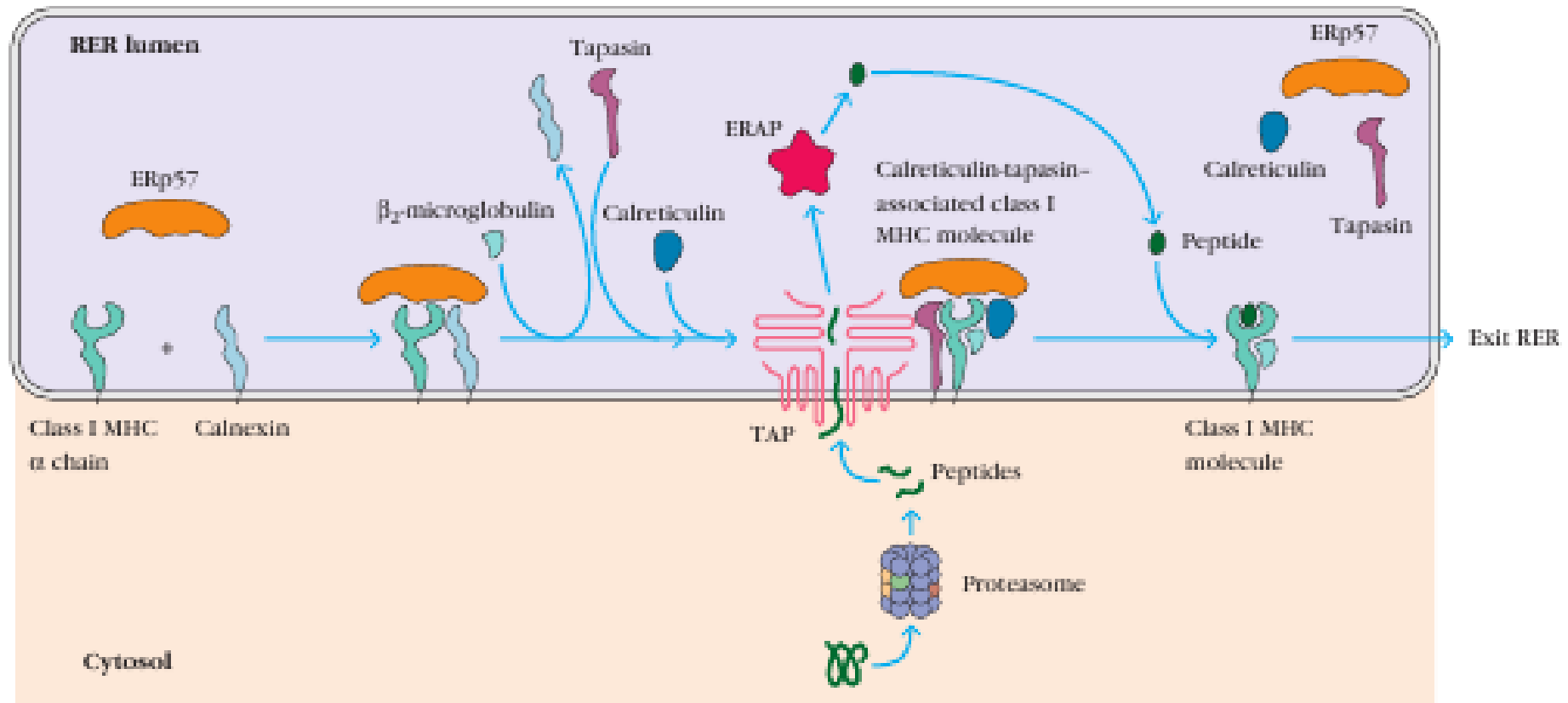
- **Tapasin (TAP associated protein)-**

- ✓ Bring transporter in proximity with class I MHC molecule
 - ✓ Allow acquiring of antigenic peptide with class I MHC molecule.

- **ERAP1**

- ✓ Help in removing residue from amino-terminal of peptide to enhance the binding capability with class I molecule.

Assembly and stabilization of class I MHC molecules



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Thanks