• For MHC class II, the  $\beta$  polypeptide chain is far more variable than the  $\alpha$  chain.

MHC genes are co-dominantly expressed so that the variant of for example *HLA-A* inherited from both parents will both be expressed. Therefore, a given individual might express HLA-A2 and HLA-A74, HLA-B8 and HLA-B27 and HLA-C1 and HLA-C6, one set inherited from their mother and the other from their father. The same co-dominant expression is seen with the MHC class II molecules.

### **Structure of MHC molecules**

## MHC class I

- The MHC class I molecule is a heterodimer.
- It is comprised of a highly polymorphic  $\alpha$  polypeptide chain of 44kDa. This  $\alpha$  chain is organized into three globular domains ( $\alpha_1$ ,  $\alpha_2$  and  $\alpha_3$ ) that protrudes from the cell surface; a hydrophobic section anchors the molecule in the membrane and it also have a short cytoplasmic N-terminal tail.
- This  $\alpha$  chain is linked to the smaller, non-polymorphic, 12kDa polypeptide called  $\beta$ -microglobulin. This  $\beta$ -microglobulin ensures correct folding of the MHC class I molecule and is not involved in peptide binding.
- The  $\beta$ -microglobulin and  $\alpha_3$  regions of the MHC class I molecule for into what resemble an immunoglobulin domain.
- The  $\alpha_1$  and  $\alpha_2$ , which are most distal to the membrane memb
- The peptide-binding great of MHC class and ecules is closed at both ends, thereby allowing it to occupy smaller peptides.

# MITO USS

The MHC class II molecule is also a heterodimer.

- It is comprised of a 34kDa  $\alpha$  polypeptide organized into two globular domains ( $\alpha_1$  and  $\alpha_2$ ) and a 29kDa  $\beta$  polypeptide chain, also organized into two globular domains ( $\beta_1$  and  $\beta_2$ ).
- $\alpha_2$  and  $\beta_2$  (nearest to surface) fold into immunoglobulin domains.
- $\alpha_1$  and  $\beta_1$  (distal to surface) form two extended helices above a  $\beta$ -pleated sheet, similar to the  $\alpha_1$  and  $\alpha_2$  of MHC class I. Again, this forms the peptide-binding groove.
- The peptide-binding groove of MHC class II molecules is open at both ends, thereby allowing it to occupy larger peptides.

# **Peptide-binding to MHC**

#### To MHC class I

- MHC class I molecules present peptides of 8-10 amino acids in length.
- The peptides that bind to MHC class I are derived from endogenous antigens.
- The forces involved in peptide binding to MHC are non-covalent.