

## MAJOR HISTOCOMPATIBILITY COMPLEX

**Q.What is major histocompatibility complex?**

- i)Every mammal's species possesses a tightly linked cluster of genes, the major histocompatibility complex (MHC), whose products play roles in intercellular recognition and in discrimination between self and nonself.
- ii)The MHC participates in the development of both humoral and cell mediated immune responses.
- iii)While antibodies may react with antigens alone, most T cells recognize antigen only when it is combined with an MHC molecule.
- iv)The particular set of MHC molecules expressed by an individual influences the repertoire of antigens to which that individual's  $T_H$  and  $T_C$  cells can respond.
- v)The major histocompatibility complex is a collection of genes arrayed within a long continuous stretch of DNA on chromosome 6 in humans and on chromosome 17 in mice.
- vi)The MHC is referred to as the HLA complex in humans and as the H-2 complex in mice.



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**Q.State the role of MHC genes./ Q.How MHC I differs from MHC II?**

- i)Class I MHC genes encode glycoproteins expressed on the surface of nearly all nucleated cells.
  - ii)The major function of the class I gene products is presentation of peptide antigens to  $T_C$  cells.
  - iii)Class II MHC genes encode glycoproteins expressed primarily on antigen-presenting cells (**macrophages, dendritic cells, and B cells**), where they present processed antigenic peptides to  $T_H$  cells.
  - iv)Class III MHC genes encode, in addition to other products, *various secreted proteins that have immune functions, including components of the complement system and molecules involved in inflammation.*
- The loci constituting the MHC are highly polymorphic, that is, many alternative forms of the gene, or alleles, exist at each locus among the population.

Class I and class II MHC molecules are membrane-bound glycoproteins that are closely related in both structure and function. Both types of membrane glycoproteins function as highly specialized antigen-presenting molecules that form unusually stable complexes with antigenic peptides, displaying them on the cell surface for recognition by T cells. In contrast, class III MHC molecules are a group of unrelated proteins that do not share structural similarity and common function with class I and II molecules.

**Q.Give a comparative analysis between MHC I and MHC II./ Q.Give characteristic features of MHC I/ Q.Give characteristic features of MHC II.**

- i)Structural analyses have revealed that the  $\alpha$  chain of class I MHC molecules is organized into three external domains ( $\alpha_1$ ,  $\alpha_2$ , and  $\alpha_3$ ), each containing approximately **90 amino acids**.
- ii)A transmembrane domain of about 25 hydrophobic amino acids followed by a short stretch of charged (hydrophilic) amino acids and a cytoplasmic anchor segment of 30 amino acids.
- iii)The  $\beta$ -microglobulin is similar in size and organization to the  $\alpha_3$  domain; it does not contain a transmembrane region and is noncovalently bound to the class I glycoprotein.
- iv)Sequence data reveal homology between the  $\alpha_3$  domain,  $\beta_2$ -microglobulin, and the constant-region domains in immunoglobulins.
- v)The  $\alpha_1$  and  $\alpha_2$  domains interact to form a platform of eight antiparallel  $\beta$ -strands spanned by two long  $\alpha$  helical regions.
- vi)The structure forms a deep groove, or cleft with the long  $\alpha$  helices as sides and the  $\beta$  strands of the  $\beta$  sheet as the bottom (**Figure 1**). This peptide-binding cleft is located on the top surface of the *class I MHC molecule*, and it is large enough to bind a peptide of 8–10 amino acids.
- vii)Processed antigen and self-peptides bound to the  $\alpha_1$  and  $\alpha_2$  domains in this deep groove. The  $\alpha_3$  domain and  $\beta_2$ -microglobulin are organized into two  $\beta$  pleated sheets each formed by antiparallel  $\beta$  strands of amino acids.
- viii)The  $\alpha_3$  domain appears to be highly conserved among class I MHC molecules and contains a sequence that interacts with the **CD8 membrane molecule** present on  $T_C$  cells.