

Disease-resistant genes: New approaches towards the control of parasitic diseases

Populations infected with various parasites have been estimated as 270 million for malaria, 200 million for schistosomiasis, 90 million for filariasis, 18 million for Chagas diseases and 12 million for leishmaniasis¹. Drug research is a top priority in many countries though drug use has many disadvantages. The various drugs and their toxic effects toward host system make drug use a threat to humans. Drug researchers are aware of the side effects and increasing drug resistance of the pathogens.

Recent studies have resulted in remarkable findings in identifying several disease-resistant loci in mice. These are identified as *Lsh*, *Ify* and *Bcg*, denoting resistance to *Salmonella typhimurium*, *Leishmania donovani*, *Mycobacterium bovis*²⁻⁵. These genes are located on the chromosome 1. Later studies have identified and selected several such genes in mouse⁶⁻¹⁰ for a variety of pathogenic infections (Table 1). How these genes confer resistance towards pathogenic infections is not known. Cloning of these

genes and their expression in other mammalian systems will give an understanding of how to combat parasitic diseases. Even though the hypothesis of one-gene: one-disease for hereditary diseases cannot be applicable in this context, one can think of polygenic localization of such disease-resistant factors. Recent progress in the isolation and characterization of candidate genes and new microsatellite-based markers for family linkage studies of the whole genome is at its frontiers. However, genome scanning approaches to gene mapping for infectious diseases are in preliminary stages in the case of parasitic diseases. This approach will be a fair play to overcome parasitic diseases and to tackle problems of drug use and its associated hazards.

Table 1. Various genes/loci identified in mammals have impact on pathogenesis resistance or susceptibility

| Gene/locus | Chromosome | Function encoded | Disease/pathogen |
|----------------------------|------------|--|--|
| MHC class II | 17 | Membrane glycoprotein presents antigen to CD4 ⁺ T cells | Malaria Leprosy |
| MHC class I | 17 | Membrane glycoprotein antigen to CD8 ⁺ T cell | <i>T. gondii</i> |
| MHC class I-b | 17 | Non classical MHC antigen presentation to CD8 ⁺ T cell | Listeria |
| TAP | 17 | Peptide transporters located in ER | Intracellular infections |
| LMP | 17 | Proteasomes: degradation of protein, antigen representation | Intracellular infections |
| Bcg/Ity/Lsh | 1 | Enhancement of APC function | <i>Mycobacteria</i> , <i>Leishmania</i> <i>Salmonella</i> |
| Vβ5 | 6 | Response to superantigen | <i>T. gondii</i> |
| Vβ6 | 6 | Response to viral superantigen | Polyoma virus |
| Vβ8 | 6 | Response to superantigen | Staphylococcus, <i>T. gondii</i> |
| VI(γ)δ subset | 13 | Response to Mycobacterial super-antigen | <i>Mycobacteria</i> , <i>Leishmania</i> |
| TNF-α | 17 | Glycoprotein | <i>T. gondii</i> , Malaria |
| IL-4 | 11 | Prototypic Th2 cytokine | <i>Leishmania</i> , Leprosy |
| INF-γ | 10 | Th1 cytokine | <i>Leishmania</i> , <i>Mycobacteria</i> |
| Cryp | 8 | Cell defense | <i>T. gondii</i> , <i>Salmonella</i> |
| NOS2 | 11 | No synthase; control pathogen multiplication | <i>Mycobacteria</i> , <i>Salmonella</i> <i>Leishmania</i> |
| Mx 1 | 16 | 75 kDa Mx 1 protein | Influenza |
| Anth 2 | 2 | Control pathogenesis | <i>Bacillus anthracis</i> |
| Bcga | 12 | Anergy to DTH | <i>Mycobacterium bovis</i> |
| HC | 2 | Pulmonary clearance of bacteria | <i>Staphylococcus aureus</i> |
| Lsr 1 | 2 | Bacterial multiplication | <i>Listeria monocytogenes</i> |
| Rict (Era 1, Api 1, Flu 1) | 5 | Control pathogenesis | <i>Rickettsia tsutsugamushi</i> |
| Ld | 17 | Brain parasite burden | <i>T. gondii</i> |
| Sc 11 | 8 | Resolution of cutaneous lesions | <i>Leishmania tropica major</i> |
| Sc 12 | 4 | Resolution of cutaneous lesions | <i>Leishmania maxicana maxicana</i> |
| Cmv 1 | 6 | Virus replication in spleen | Cytomegaloma virus |
| Fv 1 | 4 | Splenomegaly | Friend virus |
| Fv 2 | 9 | Splenomegaly | Friend virus |
| Hv 2 | 7 | Infection of macrophages and neurons | Mouse hepatitis virus 4 |
| Rmc 1 | 1 | Infectively retardation | Mink cell focus forming virus |
| Rmc f | 5 | Infectively retardation | Mink cell focus forming I |
| Rmp 2 | 2 | Survival | Mouse pox |
| Tmevd 1 | 6 | CNS demyelination | Theiler's encephelomyelitis virus |
| Tmevd 2 | 3 | CNS demyelination | Encephelomyelitis |
| Rmv 1, 2 | 1 | Viremia | Malaney virus I |

Mouse genes and their location on chromosomes have been described, however, human homologous genes have also been identified and many more will be identified within coming decades. CNS, central nervous system; ER, endoplasmic reticulum; T, *Toxoplasma*.

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