responses is by the production of the cytokine IL-10. In this regard, children with a deficiency in the expression of IL-10 or the IL-10 receptor develop inflammatory bowel disease that mimics Crohn's disease and that can be cured by allogeneic stem cell transplantation. Finally, recent data indicate that B cells may also exert regulatory function, largely through the production of IL-10. Deficiency of IL-10-producing regulatory B cells can prolong the course of multiple sclerosis in an animal model, and such cells are thought to be functionally diminished in human SLE.

It should be apparent that no single mechanism can explain all the varied manifestations of autoimmunity or autoimmune disease. Furthermore, genetic evaluation has shown that convergence of a number of abnormalities is often required for the induction of an autoimmune disease. Additional factors that appear to be important determinants in the induction of autoimmunity include age, sex (many autoimmune diseases are far more common in women), exposure to infectious agents, and environmental contacts. How all of these disparate factors affect the capacity to develop self-reactivity is currently being investigated intensively.

GENETIC CONSIDERATIONS

Evidence in humans that there are susceptibility genes for autoimmunity comes from family studies and especially from studies of twins. Studies in type 1 diabetes mellitus, rheumatoid arthritis, multiple sclerosis, and SLE have shown that ~15–30% of pairs of monozygotic twins show disease concordance, whereas the figure is <5% for dizygotic twins. The occurrence of different autoimmune diseases within the same family has suggested that certain susceptibility genes may predispose to a variety of autoimmune diseases. Genome-wide association studies have begun to identify polymorphisms in individual genes that are associated with specific autoimmune diseases. More than 50 genetic polymorphisms associated with one or more autoimmune diseases have been identified to date. It is notable that some genes are associated with multiple autoimmune diseases, whereas others are specifically associated with only one autoimmune condition. Moreover, recent genetic evidence suggests that clusters of genetic risk factors can commonly be found in groups of autoimmune diseases. Four general clusters have been identified: one group of 6 genetic polymorphisms most frequently associated with Crohn's disease, psoriasis, and multiple sclerosis; a second cluster of 8 polymorphisms most strongly associated with celiac disease, rheumatoid arthritis, and SLE; a third cluster of 7 polymorphisms most strongly associated with type 1 diabetes, multiple sclerosis, and rheumatoid arthritis; and a fourth cluster of more than 12 polymorphisms most strongly associated with type 1 diabetes, rheumatoid arthritis, celiac disease, Crohn's disease, and SLE. These results imply that autoimmune diseases with widely different clinical presentations and patterns of organ involvement could involve similar immunopathogenic pathways. For example, the same allele of the gene encoding PTPN22 is associated with multiple autoimmune diseases. Its product is a phosphatase expressed by a variety of hematopoietic cells that downregulates antigen receptor–mediated stimulation of T and B cells. The risk allele is associated with type 1 diabetes mellitus, rheumatoid arthritis, and SLE in some populations. The explanation for the association of this polymorphism with autoimmune disease is uncertain, but it is likely that it diminishes antigen receptor signaling during lymphocyte development, permitting escape of autoreactive clones or decreased activation-induced apoptosis of autoantigen-reactive lymphocytes in the periphery. In recent years, genome-wide association studies have demonstrated a variety of other genes that are involved in human autoimmune diseases. Most genes individually confer a relatively low risk for autoimmune diseases and are found in normal individuals. No gene has been identified that is essential for autoimmune diseases. In addition to this evidence from humans, certain inbred mouse strains reproducibly develop specific spontaneous or experimentally induced autoimmune diseases, whereas others do not. These findings have led to an extensive search for genes that determine susceptibility to autoimmune disease and for genes that might be protective.

The strongest consistent association for susceptibility to autoimmune disease is with particular MHC alleles. It has been suggested that the association of MHC genotype with autoimmune disease relates to differences in the ability of different allelic variations of MHC molecules to present autoantigenic peptides to autoreactive T cells. An alternative hypothesis involves the role of MHC alleles in shaping the T cell receptor repertoire during T cell ontogeny in the thymus. In addition, specific MHC gene products may themselves be the source of peptides that can be recognized by T cells. Cross-reactivity between such MHC peptides and peptides derived from proteins produced by common microbes may trigger autoimmunity by molecular mimicry. However, MHC genotype alone does not determine the development of autoimmunity. Identical twins are far more likely to develop the same autoimmune disease than MHC-identical nontwin siblings; this observation suggests that genetic factors other than the MHC affect disease susceptibility. Studies of the genetics of type 1 diabetes mellitus, SLE, rheumatoid arthritis, and multiple sclerosis in humans and mice have identified several independently segregating disease susceptibility loci in addition to the MHC. Genes that encode molecules of the innate immune response are also involved in autoimmunity. In humans, inherited homozygous deficiency of the early proteins of the classic pathway of complement (C1q, C4, or C2) as well as genes involved in the type 1 interferon pathway are very strongly associated with the development of SLE.

IMMUNOPATHOGENIC MECHANISMS IN AUTOIMMUNE DISEASES

The mechanisms of tissue injury in autoimmune diseases can be divided into antibody-mediated and cell-mediated processes. Representative examples are listed in Table 377e-3.

TABLE 377e-3 MECHANISMS OF TISSUE DAMAGE IN AUTOIMMUNE DISEASE

Effector	Mechanism	Target	Disease
Autoantibody	Blocking or inactivation	α Chain of the nicotinic acetylcholine receptor	Myasthenia gravis
		Phospholipid–β_2-glycoprotein I complex	Antiphospholipid syndrome
		Insulin receptor	Insulin-resistant diabetes mellitus
		Intrinsic factor	Pernicious anemia
	Stimulation	TSH receptor (LATS)	Graves' disease
		Proteinase-3 (ANCA)	Granulomatosis with polyangiitis
		Epidermal cadherin	Pemphigus vulgaris
		Desmoglein 3	
	Complement activation	α_3 Chain of collagen IV	Goodpasture's syndrome
	Immune complex formation	Double-stranded DNA	Systemic lupus erythematosus
		Immunoglobulin	Rheumatoid arthritis
	Opsonization	Platelet GpIIb:IIIa	Autoimmune thrombocytopenic purpura
		Rh antigens, I antigen	Autoimmune hemolytic anemia
	Antibody-dependent cellular cytotoxicity	Thyroid peroxidase, thyroglobulin	Hashimoto's thyroiditis
T cells	Cytokine production	?	Rheumatoid arthritis, multiple sclerosis, type 1 diabetes mellitus
	Cellular cytotoxicity	?	Type 1 diabetes mellitus

Abbreviations: ANCA, antineutrophil cytoplasmic antibody; LATS, long-acting thyroid stimulator; TSH, thyroid-stimulating hormone.