requiring simultaneous anti-infective therapy, nutritional support, and immunosuppression. HSCT provides a curative approach.

Functional T Cell Defects (Fig. 374-2) A subset of T cell PIDs with autosomal inheritance is characterized by partially preserved T cell differentiation but defective activation resulting in abnormal effector function. There are many causes of these defects, but all lead to susceptibility to viral and opportunistic infections, chronic diarrhea, and failure to thrive, with onset during childhood. Careful phenotyping and in vitro functional assays are required to identify these diseases, the best characterized of which are the following.

ZETA-ASSOCIATED PROTEIN 70 (ZAP70) DEFICIENCY Zeta-associated protein 70 (ZAP70) is recruited to the TCR following antigen recognition. A ZAP70 deficiency leads typically to an almost complete absence of CD8+ T cells; CD4+ T cells are present but cannot be activated in vitro by TCR stimulation.

CALCIUM SIGNALING DEFECTS A small number of patients have been reported who exhibit a profound defect in in vitro T and B cell activation as a result of defective antigen receptor-mediated Ca^{2+} influx. This defect is caused by a mutation in the calcium channel gene (*ORAI-*) or its activator (*STIM-1*). It is noteworthy that these patients are also prone to autoimmune manifestations (blood cytopenias) and exhibit a nonprogressive muscle disease.

HUMAN LEUKOCYTE ANTIGEN (HLA) CLASS II DEFICIENCY Defective expression of HLA class II molecules is the hallmark of a group of four recessive genetic defects all of which affect molecules (RFX5, RFXAP, RFXANK, and CIITA) involved in the transactivation of the genes coding for HLA class II. As a result, low but variable CD4+ T cell counts are observed in addition to defective antigen-specific T and B cell responses. These patients are particularly susceptible to herpesvirus, adenovirus, and enterovirus infections and chronic gut/liver *Cryptosporidium* infections.

HLA CLASS I DEFICIENCY Defective expression of molecules involved in antigen presentation by HLA class I molecules (i.e., TAP-1, TAP-2, and Tapasin) leads to reduced CD8+ T cell counts, loss of HLA class I antigen expression, and a particular phenotype consisting of chronic obstructive pulmonary disease and severe vasculitis.

OTHER DEFECTS A variety of other T cell PIDs have been described, some of which are associated with a precise molecular defect (e.g., IL-2-inducible T cell kinase [ITK] deficiency, IL-21 receptor deficiency, CARD11 deficiency). These conditions are also characterized by profound vulnerability to infections, such as severe Epstein-Barr virus (EBV)–induced B cell proliferation and autoimmune disorders in ITK deficiency. Milder phenotypes are associated with CD8 and CD3γ deficiencies.

HSCT is indicated for most of these diseases, although the prognosis is worse than in SCID because many patients are chronically infected at the time of diagnosis. Fairly aggressive immunosuppression and myeloablation may be necessary to achieve engraftment of allogeneic stem cells.

T Cell Primary Immunodeficiencies with DNA Repair Defects This is a group of PIDs characterized by a combination of T and B cell defects of variable intensity, together with a number of nonimmunologic features resulting from DNA fragility. The autosomal recessive disorder *ataxia-telangiectasia* (AT) is the most frequently encountered condition in this group. It has an incidence of 1:40,000 live births and causes B cell defects (low IgA, IgG2 deficiency, and low antibody production), which often require immunoglobulin replacement. AT is associated with a progressive T cell immunodeficiency. As the name suggests, the hallmark features of AT are telangiectasia and cerebellar ataxia. The latter manifestations may not be detectable before the age of 3–4 years, so that AT should be considered in young children with IgA deficiency and recurrent and problematic infections. Diagnosis is based on a cytogenetic analysis showing excessive chromosomal rearrangements (mostly affecting chromosomes 7 and 14) in lymphocytes. AT is caused by a mutation in the gene encoding the ATM protein—a kinase that plays an important role in the detection and repair of DNA lesions (or cell death if the lesions are too numerous) by triggering several different pathways. Overall, AT is a progressive disease that carries a very high risk of lymphoma, leukemia, and (during adulthood) carcinomas. A variant of AT ("AT-like disease") is caused by mutation in the *MRE11* gene.

Nijmegen breakage syndrome (NBS) is a less common condition that also results from chromosome instability (with the same cytogenetic abnormalities as in AT). NBS is characterized by a severe T and B cell combined immune deficiency with autosomal recessive inheritance. Individuals with NBS exhibit microcephaly and a bird-like face, but have neither ataxia nor telangiectasia. The risk of malignancies is very high. NBS results from a deficiency in nibrin (NBSI, a protein associated with MRE11 and Rad50 that is involved in checking DNA lesions) caused by hypomorphic mutations.

Severe forms of *dyskeratosis congenita* (also known as Hoyeraal-Hreidarsson syndrome) combine a progressive immunodeficiency that can also include an absence of B and NK lymphocytes, progressive bone marrow failure, microcephaly, in utero growth retardation, and gastrointestinal disease. The disease can be X-linked or, more rarely, autosomal recessive. It is caused by the mutation of genes encoding telomere maintenance proteins, including dyskerin (DKC1).

Finally, *immunodeficiency with centromeric and facial anomalies* (ICF) is a complex syndrome of autosomal recessive inheritance that variably combines a mild T cell immune deficiency with a more severe B cell immune deficiency, coarse face, digestive disease, and mild mental retardation. A diagnostic feature is the detection by cytogenetic analysis of multiradial aspects in multiple chromosomes (most frequently 1, 9, and 16) corresponding to an abnormal DNA structure secondary to defective DNA methylation. It is the consequence of a deficiency in the DNA methyltransferase DNMT3B, or ZBTB24.

T Cell Primary Immunodeficiencies with Hyper-IgE Several T cell PIDs are associated with elevated serum IgE levels (as in Omenn syndrome). A condition sometimes referred to as *autosomal recessive hyper-IgE syndrome* is notably characterized by recurrent bacterial infections in the skin and respiratory tract and severe skin and mucosal infections by pox viruses and human papillomaviruses, together with severe allergic manifestations. T and B lymphocyte counts are low. Mutations in the *DOCK8* gene have been found in most of these patients. This condition is an indication for HSCT.

A very rare, related condition with autosomal recessive inheritance that causes a similar susceptibility to infection with various microbes (see above), including mycobacteria, reportedly results from a deficiency in Tyk-2, a JAK family kinase involved in the signaling of many different cytokine receptors.

Autosomal Dominant Hyper-IgE Syndrome This unique condition, the *autosomal dominant hyper-IgE syndrome*, is usually diagnosed by the combination of recurrent skin and lung infections that can be complicated by pneumatoceles. Infections are caused by pyogenic bacteria and fungi. Several other manifestations characterize hyper-IgE syndrome, including facial dysmorphy, defective loss of primary teeth, hyperextensibility, scoliosis, and osteoporosis. Elevated serum IgE levels are typical of this syndrome. Defective T_H17 effector responses have been shown to account at least in part for the specific patterns of susceptibility to particular microbes. This condition is caused by a heterozygous (dominant) mutation in the gene encoding the transcription factor STAT3 that is required in a number of signaling pathways following binding of cytokine to cytokine receptors (such as that of IL-6 and the IL-6 receptor). It also results in partially defective antibody production because of defective IL-21R signaling. Hence, immunoglobulin substitution can be considered as prophylaxis of bacterial infections.

Cartilage Hair Hypoplasia The autosomal recessive *cartilage hair hypoplasia* (CHH) disease is characterized by short-limb dwarfism, metaphyseal dysostosis, and sparse hair, together with a combined T and B cell PID of extremely variable intensity (ranging from quasi-SCID to no clinically significant immune defects). The condition can predispose to erythroblastopenia, autoimmunity, and tumors. It is caused by mutations in the *RMRP* gene for a noncoding ribosome-associated RNA.