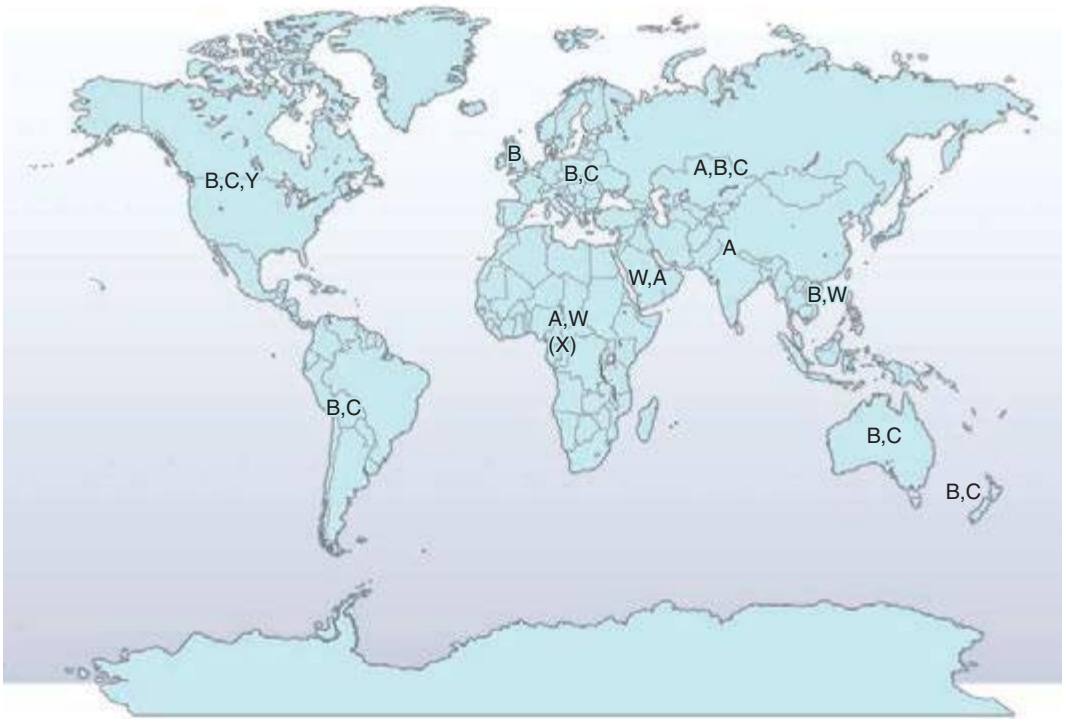

FIGURE 180-4 **Global distribution of meningococcal serogroups, 1999–2009.**

1990s in North America, the median ages for patients with disease due to serogroups B, C, Y, and W were 6, 17, 24, and 33 years, respectively. After early childhood, a second peak of disease occurs among adolescents and young adults (15–25 years of age) in Europe and North America. It is thought that this peak relates to social behaviors and environmental exposures in this age group, as discussed below. Most cases of infection with *N. meningitidis* in developed countries today are sporadic, and the rarity of the disease suggests that individual susceptibility may be important. A number of factors probably contribute to individual susceptibility, including the host's genetic constitution, environment, and contact with a carrier or a case.

The best-documented genetic association with meningococcal disease is complement deficiency, chiefly of the terminal complement components (C5–9), properdin, or factor D; such a deficiency increases the risk of disease by up to 600-fold and may result in recurrent attacks. Complement components are believed to be important for the bactericidal activity of serum, which is considered the principal mechanism of immunity against invasive meningococcal disease. However, when investigated, complement deficiency is found in only a very small proportion of individuals with meningococcal disease (0.3%). Conversely, 7–20% of persons whose disease is caused by the less common serogroups (W, X, Y, Z, 29E) have a complement deficiency. Complement deficiency appears to be associated with serogroup B disease only rarely. Individuals with recurrences of meningococcal disease, particularly those caused by non-B serogroups, should be assessed for complement deficiency by measurement of total hemolytic complement activity. There is also limited evidence that hyposplenism (through reduction in phagocytic capacity) and hypogammaglobulinemia (through absence of specific antibody) increase the risk of meningococcal disease. Genetic studies have revealed various associations with disease susceptibility, including complement and mannose-binding lectin deficiency, single-nucleotide polymorphisms in Toll-like receptor (TLR) 4 and complement factor H, and variants of Fc gamma receptors.

Factors that increase the chance of a susceptible individual's acquiring *N. meningitidis* via the respiratory route also increase the risk of meningococcal disease. Acquisition occurs through close contact with carriers as a result of overcrowding (e.g., in poor socioeconomic settings, in refugee camps, during the Hajj pilgrimage to Mecca, and during freshman-year residence in college dormitories) and certain social behaviors (e.g., attendance at bars and nightclubs, kissing). Secondary cases may occur in close contacts of an index case (e.g., household members and persons kissing the infected individual); the risk to these contacts may be as high as 1000 times the background rate in the population. Factors that damage the nasopharyngeal epithelium also increase the risk of both colonization with *N. meningitidis* and invasive disease. The most important of these factors are cigarette smoking (odds ratio, 4.1) and passive exposure to cigarette smoke. In addition, recent viral respiratory tract infection, infection with *Mycoplasma* species, and winter or the dry season have been associated with meningococcal disease; all of these factors presumably either increase the expression of adhesion molecules in the nasopharynx, thus enhancing meningococcal adhesion, or facilitate meningococcal invasion of the bloodstream.

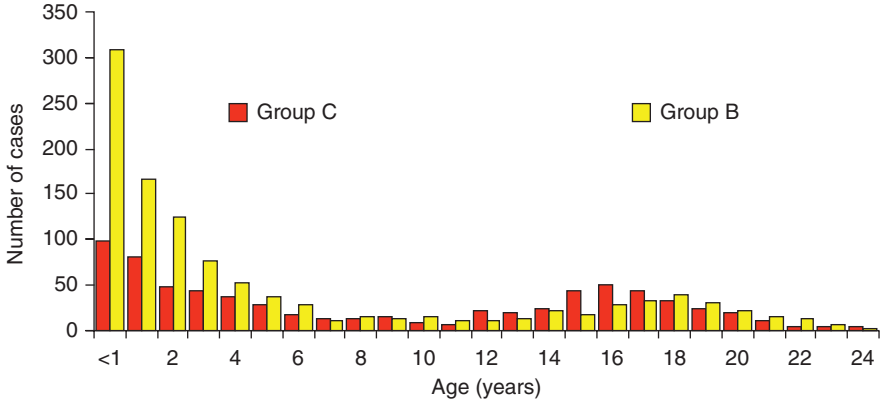

FIGURE 180-5 **Age distribution of serogroups B and C meningococcal disease in England and Wales, 1998–1999.** *(Health Protection Agency, UK; www.hpa.org.uk.)*