Advances in the biology, diagnosis and host–pathogen interactions of parvovirus B19

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Increased recognition of parvovirus B19 (B19), an erythrovirus, as a significant human pathogen that causes fetal loss and severe disease in immunocompromised patients has resulted in intensive efforts to understand the pathogenesis of B19-related disease, to improve diagnostic strategy that is deployed to detect B19 infection and blood-product contamination and, finally, to elucidate the nature of the cellular immune response that is elicited by the virus in diverse patient cohorts. It is becoming clear that at least three related erythrovirus strains (B19, A6/K71 and V9) are circulating in the general population and that viral entry into target cells is mediated by an expanding range of cellular receptors, including P antigen and \( \beta \)-integrins. Persistent infection by B19 is emerging as a contributory factor in autoimmune disease, a hypothesis that is constrained by the detection of B19 in the skin of apparently healthy individuals. B19 infection during pregnancy may account for thousands of incidences of fetal loss per annum in Europe, North America and beyond, yet there is currently only minimal screening of pregnant women to assess serological status, and thereby risk of infection, upon becoming pregnant. Whilst major advances in diagnosis of B19 infection have taken place, including standardization of serological and DNA-based detection methodologies, blood donations that are targeted at high-risk groups are only beginning to be screened for B19 IgG and DNA as a means of minimizing exposure of at-risk patients to the virus. It is now firmly established that a Th1-mediated cellular immune response is mounted in immunocompetent individuals, a finding that should contribute to the development of an effective vaccine to prevent B19 infection in selected high-risk groups, including sickle-cell anemics.

Parvovirus B19

In 1975, Yvonne Cossart discovered what was to become known as human parvovirus B19 (B19) (Cossart et al., 1975). B19 was first associated with disease in 1981, when it was linked to an aplastic crisis in a patient with sickle-cell disease. It has since been shown to cause erythema infectiosum (EI) (fifth disease of childhood), spontaneous abortion and some forms of acute arthritis (Anderson et al., 1983; Kinney et al., 1988; Woolf & Cohen, 1995).

B19 is a small, non-enveloped, ssDNA virus and, like all parvoviruses, the capsid proteins are arranged with icosahedral symmetry. B19 is 20–25 nm in diameter and has a genome of 5.6 kb (Clewley, 1984; Cotmore & Tattersall, 1984). The B19 capsid consists of an 83 kDa minor structural protein, VP1, and a 5 kDa major structural protein, VP2. VP2 makes up about 95% of the total capsid, with VP1 accounting for the remaining 5% (Ozawa et al., 1987). The sequences of the two proteins are collinear, with VP2 being identical to the carboxyl-terminus of VP1; however, VP1 comprises an additional 227 aa domain that is unique to the amino-terminal (Fig. 1). To the left of these sequences on the B19 genome is the ORF for a non-structural protein, NS1, which encodes a protein product of 77 kDa. NS1 is a phosphoprotein with important regulatory functions, including control of transcription (Momoeda et al., 1994a) and virus replication; it also plays a role in host-cell death (Ozawa et al., 1988). It possesses DNA-binding properties (Raab et al., 2002) and biochemical activities, such as ATPase, helicase and site-specific endonuclease activities, as well as nuclear localization signals (Li & Rhode, 1990; McCarty et al., 1992; Jindal et al., 1994; Brown & Young, 1997). B19 NS1 has also been shown to affect G1, but not G2, arrest in erythroid UT7/Epo-S1 cells (Morita et al., 2003). B19 uses a single promoter, p6, which is capable of expressing structural and non-structural genes differentially (Blundell et al., 1987;
Ozawa et al., 1987). It has been demonstrated that NS1 interacts directly with the p6 promoter and with cellular transcription factors Sp1/Sp3 to affect transcriptional regulation (Raab et al., 2002). Two other small polypeptides have been identified, one encoded by a region in the middle of the genome with a predicted Mr of 7–5 kDa and the other, which is encoded at the extreme right-hand end of the genome, with a predicted Mr of 11 kDa (St Amand et al., 1991). Spliced transcripts of these two polypeptides have been found in infected cells, but their function has been hitherto unknown (Luo & Astell, 1993).

As a result of increased awareness of and screening for B19, a number of novel genotypes have been identified. Servant et al. (2002) suggested that B19 should be classified as a genotype 1 erythrovirus, with newly identified strains A6 (Nguyen et al., 2002) and K71 (Hokynar et al., 2002) classified as genotype 2 and erythrovirus V9 (Nguyen et al., 1998) as the prototype of genotype 3. V9 has overall nucleotide sequence variation of about 12% from B19 isolates, with the majority of sequence divergence occurring in the 5′ portion of the VP1 unique region; however, differences in sequence are not restricted to this area, but are scattered throughout the genome (Nguyen et al., 1999; Heegaard et al., 2001). K71 persists in human skin and has a nucleotide divergence of 10.8% from B19 and 8.6% from V9 (Hokynar et al., 2002).

**Infecitivity, transmission and epidemiology**

The only known host for B19 is humans. The virus replicates in human erythroid progenitor cells (late erythroid cell precursors and burst-forming erythroid progenitors) of the bone marrow and blood, inhibiting erythropoiesis (Mortimer et al., 1983). Tropism of productive B19 infection is mainly due to the restrictive cellular distribution of the P blood group antigen globoside (Gb4) (Brown et al., 1993, 1994), which is found most commonly on cells of the erythroid lineage, but also on platelets, tissues from the heart, liver, lung, kidney and endothelium and on synovium (Cooling et al., 1995; Jordan & DeLoia, 1999). Individuals who lack erythrocyte P antigens are very rare (1 in 200 000) and apparently cannot be infected by B19 (Brown et al., 1994; Chipman et al., 1996).

The limited tropism of B19 is not fully understood, as reduced capsid expression has been observed in non-permissive cells; however, intracellular factors that are found only in erythroid cells are assumed to be essential for optimal transcription and virus replication (Ozawa et al., 1987; Kurpad et al., 1999; Gallinella et al., 2000). It has been shown that the level of P antigen expression on cells is not related directly to the efficiency of virus binding. In addition, some cell lines cannot be transduced by a B19 vector, despite P antigen expression and virus binding to the P antigen, thus indicating that a co-receptor is probably essential for virus entry into human cells (Weigel-Kelley et al., 2001). Thus, recent evidence suggests that the presence of the P antigen alone is not sufficient to gain entry into cells (Weigel-Kelley et al., 2001) and it has been suggested that multiple β-integrins may function as co-receptors for B19 cellular assimilation (Weigel-Kelley et al., 2003).

B19 studies have been hampered by difficulties in propagating the virus in vitro. Recent studies have indicated that infection under hypoxic conditions [1% (v/v) O2] causes upregulation of B19 expression, which is associated with increased virus replication and production of infectious virions (Pillet et al., 2002). Due to the cytotoxic nature of the non-structural protein NS1, no continuous cell line that propagates B19 has been established (Ozawa et al., 1987; Momoeda et al., 1994b).

B19 transmission occurs most commonly by personal contact via aerosol or respiratory secretions; however, contaminated blood products, such as clotting factor concentrates, are a source of iatrogenic transmission (Anderson et al., 1985; Lyon et al., 1989; Williams et al., 1990; Santagostino et al., 1994; Erdman et al., 1997). B19 can be transmitted transplacentally from an infected mother to her fetus, which may lead to non-immune fetal hydrops (NIHF), spontaneous abortion or intrauterine fetal death (IUFD) (Clewley et al., 1987; Miller et al., 1998; Skjoldbrand-Sparre et al., 2000). The P blood group antigen, which serves as a receptor for B19, has been detected on cells of the villous trophoblast of placental tissues in varying amounts during the course of pregnancy. In the first trimester, levels of the P antigen are very high; they begin to decline in the second trimester and become undetectable by the mid-stages of trimester 3 (Jordan & DeLoia, 1999). This high level of globoside receptor on placental cells in early pregnancy may act as a pathway for B19 to be transmitted from the mother to the fetus, whereby the virus can then infect erythroid progenitor cells for replication. In support of this hypothesis, Wegner & Jordan (2002) have shown conclusively that [125I]-labelled VP2 capsid interaction with villous cytotrophoblast cells is mediated by P antigen.

Infection with B19 is very common and cases of infection have been reported all over the world in all seasons. Seroprevalence increases with age and, by adulthood,
>70% of the adult population is seropositive (Kerr et al., 1999). Children are the main source of transmission and outbreaks can persist for months in schools and day-care centres, due to the relatively large number of seronegative children and close contact of children within this environment (Tuckerman et al., 1986; Grilli et al., 1989). The annual seroconversion rate among women of childbearing age has been estimated to be 1.5% during endemic periods and 13% during epidemics (Koch & Adler, 1989; Valeur-Jensen et al., 1999). Furthermore, infection by B19 during pregnancy can lead to spontaneous abortion or fetal anaemia. Consequently, the question must now be posed whether the B19 immune status of pregnant women should be determined routinely on initial presentation, to facilitate improved pregnancy outcome with respect to potential B19 infection.

**Manifestations and clinical symptoms**

B19 has been associated with an expanding range of clinical disorders since the discovery that it is the aetiopathological agent of EI. This is a mild childhood illness that is characterized by an erythematous rash that affects the face, trunk and limbs of the body. It is also associated with complications during pregnancy, acute arthropathy, disease in immunocompromised patients and transient aplastic crisis (TAC).

**B19 infection and pregnancy**

Exposure to and infection by B19 can lead to serious complications during pregnancy. Infection during pregnancy may result in fetal anaemia, spontaneous abortion and hydrops fetalis (Brown et al., 1984; Kinney et al., 1988; Heegaard & Hornsleth, 1995). About 30–40% of women are non-immune and do not possess neutralizing antibodies to B19 and, therefore, are susceptible to infection by this virus. A vertical transmission rate of 33% has been reported by the Public Health Laboratory Service in the UK (PHLS, 1990) and others have reported similar rates (Brown et al., 1984; Hall et al., 1990), although a recent study reported a transmission rate of 51% (Yaegashi, 2000). There are over four million births in Europe per annum (Eurostat, 1998) and, because 30% of pregnant women are B19-seronegative, over 1 200 000 European women are therefore susceptible to B19 infection during pregnancy. Assuming a combined rate of infection and fetal loss of 0.2% (Levy et al., 1997; Miller et al., 1998; Wattre et al., 1998), it can be estimated that approximately 3000 pregnancies per annum may be lost. Analogous birth rates in the USA and Canada imply that a similar incidence of fetal death due to B19 infection can be expected for these countries. These calculations are based solely on live births and, as the number of actual pregnancies is much higher, the above estimate is conservative. Pregnant women are most susceptible to B19 infection during epidemics and also when exposed to infected children in the home (Valeur-Jensen et al., 1999). During outbreaks, transmission rates of 25% in schools and 50% at home have been reported (Anderson et al., 1990). Most pregnant women are asymptomatic, but some do experience symptoms, such as exanthema and arthralgia (Komischke et al., 1997). As these symptoms are commonly associated with pregnancy, acute B19 infection can often be overlooked; however, routine screening for symptoms of B19 infection or seroconversion would overcome this problem.

Fetal death usually occurs 4–6 weeks post-infection, but has been reported up to 12 weeks after B19 symptomatic infection (Hedrick, 1996). A study of 427 pregnant women with B19 infection in the UK observed that fetal loss was confined to the first 20 weeks of gestation (Miller et al., 1998). This is supported by figures released in the UK and other studies, which reported that fetal loss as a consequence of intrauterine B19 infection is highest in, but not restricted to, the first 20 weeks of gestation (Hall et al., 1990; PHLS, 1990). The critical time of infection has since been narrowed down to the 16th week of gestation (Yaegashi et al., 1999). Most cases of fetal loss due to B19 infection have been reported in the second trimester (Enders & Biber, 1990; Torok, 1990; Wattre et al., 1998). This susceptibility could be attributed, at least in part, to the relative immaturity of the fetal immune response at this stage. More important, though, is the tropism that B19 has for erythroid progenitor cells (Yaegashi, 2000) and the fact that in the second trimester of pregnancy, the life-span of fetal red blood cells (RBCs) is shortened and RBC mass increases three- to fourfold during this period of gestation (Rodis et al., 1988). B19 replication within erythroid progenitor cells leads to apoptosis, which ultimately results in inhibition of erythropoiesis (Morey et al., 1993). Erythroblastopenia can then occur as a consequence of B19 replication, causing severe fetal anaemia.

Anaemia is an underlying factor in the development of hydrops. Fetal hydrops was first associated with B19 in 1984 (Brown et al., 1984). Since then, 10–20% of NIHF cases have been reported to be B19-associated (Yaegashi et al., 1994; Jordan, 1996) and, in a study of B19 infection in Japanese women during pregnancy, the risk of hydrops was determined to be about 10% (Yaegashi et al., 1999). NIHF usually occurs 2–4 weeks after maternal B19 infection (Komischke et al., 1997). Cases of IUFD that are associated with fetal hydrops and caused by B19 have been reported most commonly in the second trimester and, to a lesser extent (unquantified as yet), in the third trimester of pregnancy (Singh et al., 1997). When cases of IUFD that occurred during an 18 month period in the UK were examined, it was discovered that 11 deaths were caused by B19 in the second trimester and, of these, only three were hydropic (Wright et al., 1996). In a separate study over a 16 year period, ten cases of IUFD were reported, which presented in gestational weeks 15–29. Of those cases, 90% of the fetuses were hydropic, 30–40% had associated heart failure and three of the maternal infections were asymptomatic (Morey et al., 1993).

Until recently, third-trimester fetal loss or IUFD caused by acute B19 infection had not been widely reported. However,
of 93 IUFD cases that were examined, 7.5% had B19 DNA in placental tissue in the absence of fetal hydrops (Skjoldebrand-Sparre et al., 2000). Unusually, none of the infected pregnant women in this study showed any clinical symptoms of B19 infection. B19-associated IUFD in the final stages of gestation may have been overlooked previously, due to inadequate diagnostic procedures and the difference in clinical features of third-trimester B19 infection. The most striking observation in these IUFD cases was the lack of fetal hydrops and the fact that many of the cases had either delayed or absent B19 IgG responses. Histopathological examination of the fetuses revealed no major abnormalities. Similar reports of non-hydropic, third-trimester IUFD that was associated with B19 infection have been published (Tolfvenstam et al., 2001a). Here, it was revealed by PCR analysis of fetal or placental tissues that 15% of IUFD cases were attributable to B19 infection. This study also observed delayed B19-specific antibody responses, as the mothers involved had no serological evidence of an acute B19 infection. However, follow-up studies showed evidence of seroconversion within 6 months. Tissue samples showed no signs of virus inclusions and immunochemistry analysis revealed no evidence of B19 proteins (Tolfvenstam et al., 2001a). Although the concept of B19-induced, third-trimester fetal loss has proved somewhat controversial (Crowley et al., 2001; Sebire, 2001), it further illustrates the requirements for awareness of B19 pathogenesis and diagnostic B19 PCR screening during pregnancy. Furthermore, Nunoue et al. (2002) suggested strongly that prospective studies to evaluate the relationship between time of infection and IUFD, with and without signs of fetal hydrops, are necessary. In fact, B19 PCR may be the most sensitive way of diagnosing intrauterine B19 infection, especially as >50% of infected fetuses test negative for B19 IgM (Dieck et al., 1999).

Administration of high-titre intravenous immunoglobulin (IVIG) has been shown to be successful in treating fetal hydrops in some cases (Selbing et al., 1995; Alger, 1997). For cases of fetal infection, intrauterine blood transfusions may be beneficial (Schwarz et al., 1988a; Hansmann et al., 1989), especially in the case of hydrops, but this procedure does involve additional risks to the outcome of pregnancy (Berry et al., 1992; Cameron et al., 1997; Bousquet et al., 2000). A study by Wattre et al. (1998) reported two cases where intrauterine blood transfusions led to the cessation of symptoms and the birth of normal babies. In a separate study, 38 cases of B19-associated fetal hydrops were reported, 12 of which received intrauterine blood transfusion. Although three of these fetuses subsequently died, the probability of death among fetuses that did not receive a blood transfusion was significantly higher (Fairley et al., 1995). In addition, spontaneous resolution of hydrops without intervention has been reported, suggesting that treatment is not always necessary (Pryde et al., 1992).

B19 infection during pregnancy is not a significant cause of birth defects; however, at least one incidence of congenital cardiomyopathy has been linked to B19 infection (Barton et al., 1997).

**Arthopathy**

Like the rubella virus (Lee, 1960), B19 infection has also been linked to arthritis and arthralgias, most commonly in adults, but also in children (Reid et al., 1985). On average, 50% of adult cases of EI have associated joint manifestations that may persist for up to 1 month (Cassinotti et al., 1995). B19 arthritis is usually symmetrical, affecting mainly the small joints of the hands, wrists and knees (Reid et al., 1985). It is more common in females than males, with an estimated 60% of women with symptomatic disease that manifests in arthropathy (White et al., 1985; Woolf et al., 1989). Symptoms generally subside within 3 weeks without any damage to the joints (Woolf et al., 1991), but about 20% of affected women suffer persistent or recurring arthropathy. About 75% of these patients have an associated rash and <20% have the typical ‘slapped cheeks’ facial exanthem. B19 has been proposed as the causative agent of arthritic conditions that exhibit similar symptomologies to those found in rheumatoid arthritis (RA), juvenile RA and erosive polyarthritis, as recent B19 infection and high levels of B19 antibodies have been evident in many of these patients (White et al., 1985; Nocton et al., 1993; Mimori et al., 1994; Tyndall et al., 1994). It has been suggested that B19-associated arthritis is related to certain human leukocyte antigen (HLA) haplotypes of patients, with individuals of either HLA DR4 or B27 being most susceptible (Klouda et al., 1986; Jawad, 1993); however, it is unclear how B19 causes symptoms that are associated with arthritis. Analogous to the appearance of exanthema in EI, arthritis usually occurs after development of B19-specific antibodies. This suggests that symptoms may be due to formation of immune complexes. Despite the fact that the P antigen is expressed on synovium, it has been shown that synovial membrane cells are non-permissive to B19 (Miki & Chantler, 1992; Cooling et al., 1995). Normal human synovial fibroblasts have been shown to exhibit increased invasiveness following exposure to B19 viraeic serum, as judged by the acquired ability to degrade reconstituted cartilage matrix (Ray et al., 2001). B19 may gain entry to cells that possess the B19 receptor but are not actively dividing, resulting in the production of excessive cytotoxic NS1 (Ozawa et al., 1988). The B19 NS1 protein causes the secretion of proinflammatory cytokines, which could cause the inflammation and cell damage that are seen in patients with B19-associated arthritis and other inflammatory and autoimmune disorders that have been linked to B19 infections (Moffatt et al., 1996; Mitchell, 2002). In one study, antibodies that were specific for the non-structural protein NS1 were found in patients with persistent B19-associated arthropathy, but not in serum from individuals with evidence of past infection without complications (von Poblotzki et al., 1995a), thus suggesting an altered host response in the former cohort. However, others have disputed this, reporting similar NS1 antibody reactivity in
patients with chronic or acute B19-associated arthropathy (Mitchell et al., 2001) and recently infected healthy individuals (Searle et al., 1998; Ennis et al., 2001; Mitchell et al., 2001; Heegaard et al., 2002a). von Landenberg et al. (2003) further suggested that B19 may be involved directly in the induction of autoimmune reactions that are mediated, at least in part, by anti-phospholipid antibodies, because of the prevalence of these antibodies in persistently B19-infected individuals.

There is significant evidence of B19 DNA persistence in bone marrow, peripheral blood and synovial tissues of patients with chronic, B19-associated arthropathy (Foto et al., 1993; Musiani et al., 1995; Nikkari et al., 1995). However, it has also been shown that although B19 DNA persisted in the synovium tissue of 28% of children who presented with chronic arthritis, an even higher proportion (48%) of seropositive, immunocompetent volunteers had B19 DNA in their synovium tissues. These results imply that B19 DNA in synovium tissue may not be associated directly with symptoms of chronic arthropathy. None of the individuals tested had evidence of B19 DNA in their synovial fluid, bone marrow or blood and all were positive for B19 IgG antibodies (Söderlund et al., 1997). Nonetheless, a recent report further enhanced the correlation between B19 infection and rheumatic childhood disease (Lehmann et al., 2003). This work clearly elucidated a significant difference in serum and/or synovial fluid-derived B19 DNA (P < 0.0001) between control (9/124, 7%) and patient (26/74, 35%) specimens and concluded that the rate of persistent B19 infection in these patients is significantly higher than in age-matched controls.

The recent finding of B19 DNA in 64% (14/22) of control skin biopsies, compared to 50% (18/36) of chronic urticaria patients, confirms that caution should be exercised in drawing conclusions regarding B19 involvement in skin disorders and possibly in other B19-associated clinical disorders (Vuorinen et al., 2002).

**Chronic B19 infection in the immunocompromised host**

A host with a compromised immune system is particularly at risk of B19 infection, including people with AIDS, cancer patients who are receiving chemotherapy and transplant patients on immunosuppressive drugs (Young, 1996). Many are unable to produce neutralizing antibodies to clear the virus and this can lead to persistent infection, resulting in anaemia (Kurtzman et al., 1989a; Young, 1996). In one case study, an AIDS patient developed severe anaemia as a result of chronic pure red-cell aplasia that was caused by B19 infection (Koduri et al., 1997). Despite remission following IVIG transfusions, the patient suffered several recurrences of severe anaemia. In another report, sera obtained from transplant patients who were receiving bone-marrow grafts (n = 27) were analysed by PCR (Schleuning et al., 1999). Of the cohort tested, 15% were B19 DNA-positive and many of these patients also developed a reticular rash. Although development of a rash during B19 infection is thought to be mediated by formation of immune complexes, these patients did not exhibit any signs of a B19-specific antibody response. Therefore, it was hypothesized that the rash was a consequence of a direct virus effect on the skin; this finding is supported by the fact that B19 DNA has been found previously in a skin biopsy from a male patient with B19-associated fever, rash and polyarthritis (Nikkari et al., 1996). Persistent B19 infection results in chronic suppression of erythropoiesis with chronic anaemia. A report by Graeve et al. (1989) described how four children who were undergoing cancer chemotherapy treatment were infected by B19, resulting in chronic bone-marrow suppression. Schleuning et al. (1999) also reported that one of the transplant patients subsequently died from heart failure and B19 DNA was detected in the myocardium, but not in peripheral blood, indicating that heart failure was a consequence of B19 infection [heart failure has been recognized as a feature of B19 infection in the past (Chia & Jackson, 1996)]. Another of the transplant patients investigated was found to have developed hepatitis (Schleuning et al., 1999), which was also attributed to B19 infection (Yoto et al., 1996). As these transplant patients were subjected to strict decontamination procedures, including isolation in single rooms with positive airflow and decontaminated food, it is thus unlikely they contacted B19 via respiratory secretions. Platelet concentrates were screened for B19 prior to administration and were therefore not a likely source of transmission. However, as B19 DNA in clotting and immunoglobulin concentrates is known to cause infection, this may have been the route of transmission for the virus (Saldanha & Minor, 1996).

Many immunocompromised patients with chronic anaemia respond positively to IVIG therapy; however, individuals may suffer from recurrent relapses of aplasia (Koduri et al., 1997, 1999; Moudgil et al., 1997). In addition, administration of IVIG may not always be effective, as infection may persist despite treatment, particularly in transplant patients who are heavily immunosuppressed (Moudgil et al., 1997; Schleuning et al., 1999; Lui et al., 2001). To date, no data are available on the actual protective level of B19 IgG, although levels of >6 IU ml⁻¹ are thought to be protective (Searle et al., 1997). As patients fail to mount an antibody response, serological diagnosis is futile and detection of B19 infection is therefore usually achieved by B19 DNA detection via a PCR assay.

**Transient aplastic crisis (TAC)**

B19-associated TAC may occur in individuals who exhibit underlying chronic haemolytic disorders, such as hereditary spherocytosis (Beland et al., 1997). In 1981, it was discovered that B19 caused TAC in children with sickle-cell anaemia (Serjeant et al., 1981) and it is now clear from subsequent studies that almost 70% (118/177) of B19 infections in this cohort resulted in TAC (Serjeant et al., 2001).
Vaccine
No specific therapy is required for B19 infection in immunoocompetent individuals. Symptoms of arthropathy can be treated with non-steroidal, anti-inflammatory drugs.

Humoral immune response
B19 viraemia occurs 1 week after exposure and usually lasts about 5 days, with virus titres peaking on the first 2 days. B19-specific IgM antibodies are detected late in the viraemic stage (at about day 10 or 12) and can persist for up to 5 months (Anderson et al., 1985; Schwarz et al., 1988b; Yaegashi et al., 1989) but, in some patients, can last even longer (Musiani et al., 1995). Specific IgG antibodies are detectable about 15 days post-infection, remain high for several months and persist long-term. IgA antibodies are detectable for a short period following the onset of clinical symptoms (Erdman et al., 1991). Development of the antibody response corresponds to virus clearance and also, in the vast majority of cases of B19 infection in immunoocompetent individuals, protection from disease (Anderson et al., 1985). A study of children with sickle-cell disease showed that those who had had one episode of B19-associated TAC did not suffer a second episode (Serjeant et al., 1993).

Historically, the B19 VP1 protein, and in particular the VP1-unique region, was thought to be the immunodominant antigen and its incorporation into serological assays was thought to be essential (Rayment et al., 1990). However, it is now clear that this observation, which was based on the absence of antibodies to linear epitopes of VP2 when screened by Western blot, was somewhat erroneous. It has now been established conclusively that antibodies against capsid VP2 are maintained, even when B19 IgG directed against the VP1-unique region is lost (Kerr et al., 1999; Manaresi et al., 1999; Corcoran et al., 2000).

Specific anti-virus antibody is considered to be a significant mechanism of immune protection, based on the circumstantial evidence that high-dose immunoglobulin therapy is sometimes beneficial in infected patients (Kurtzman et al., 1989b; Schwarz et al., 1990). Antibodies against linear epitopes of VP2 and, to some extent, VP1 disappear abruptly after B19 infection, whereas IgG reactivity against conformational epitopes of both VP1 and VP2 persists (Söderlund et al., 1995; Kerr et al., 1999). Persistent infections that are associated with chronic anaemia where the immune response to B19 has failed to produce neutralizing antibodies or they have been at very low levels have been observed (Kurtzman et al., 1987, 1988; Coulombel et al., 1989).

Diagnosis of B19 infection – choice of antigen
Accurate laboratory diagnosis of recent B19 infection or past exposure relies on screening plasma specimens for either specific antibody reactivity against virus capsid proteins that are expressed in eukaryotic expression systems (e.g. the baculovirus expression system) or for B19 DNA by using PCR. Immunoassays that only incorporate Escherichia coli-expressed B19 antigens, which have undergone denaturation as part of the manufacturing process, will produce false-negative results, due to the absence of conformational epitopes (Jordan, 2000). A unique advantage of the eukaryotic baculovirus expression system is its ability to direct the post-translational protein folding that is necessary for the production of soluble, conformationally intact VP2 capsid proteins (Brown et al., 1990; Kerr et al., 1995a). Unlike B19 VP2, VP1 does not appear to form soluble capsid structures; however, VP1 has been produced as a ‘conformationally intact’ protein that retains conformational epitopes that are present in the native virion (Brown et al., 1990; Kerr et al., 1999).

A number of authors have stated that co-expression of VP1 and VP2 in eukaryotic expression systems results in the formation of empty capsids that are antigenically analogous to native B19 virions. Furthermore, it has been hypothesized that such co-capsids contain conformational epitopes that are essential for accurate detection of infection (Kajigaya et al., 1989, 1991; Franssila et al., 2001; Ballou et al., 2003). To date, there are no data to prove that co-capsids are actually present in such preparations and it cannot be excluded that separate, conformationally intact entities that are comprised of either VP1 or VP2 are in fact present.

More recent evidence from a number of authors now suggests that B19 NS1 IgG and IgM detection also plays a significant role in diagnosis of acute infection, thereby supplementing the role played by B19 capsid antigens as diagnostic antigens (Ennis et al., 2001; Heegaard et al., 2002a).

B19 IgM immunodetection
Acute B19 infections are confirmed by B19-specific IgM reactivity, whereas past infections are detected by IgG reactivity (Anderson et al., 1985). In most situations, IgM antibodies appear 7–10 days post-infection and are directed against linear and conformational epitopes of VP1 and VP2 (Palmer et al., 1996; Manaresi et al., 2001). It has been reported that IgM against conformational epitopes on VP1 and VP2 and against linear epitopes on VP1 appears at the same time post-infection and at the same frequency. However, it was also revealed that IgM reactivity against the minor capsid protein, VP1, may persist somewhat longer post-infection (Palmer et al., 1996; Manaresi et al., 2001). If IgM responses against conformational VP1 persist when other B19-specific IgM antibodies are absent, then diagnostic techniques that incorporate conformational VP1 may not be the most suitable markers of acute B19 infection. However, another study observed no difference in IgM reactivity against conformational epitopes of the capsid proteins in diagnosing B19 infection (Kerr et al., 1999). Furthermore, these authors observed no disparity in IgM reactivity against native (conformationally intact) and linearized antigens for both VP1 and VP2.

Presently, there is no International Standard preparation for B19 IgM and only one B19 IgM diagnostic test that has been
cleared by the US Food and Drug Administration (FDA) is available. It is a μγ-capture enzyme immunoassay (EIA) that utilizes B19 recombinant VP2 capsids for the detection of specific IgM in human serum or plasma. This immunoassay has 89.1% sensitivity and 99.4% specificity (Doyle et al., 2000) and is used widely for the diagnosis of recent B19 infection (Jordan, 2000; Mitchell et al., 2001; Vuorinen et al., 2002). Furthermore, validated alteration of the immunoassay cut-off, based on receiver operating characteristic analysis, facilitates improved immunoassay sensitivity, which may have a utility in detection of lower levels of B19-specific IgM in immunocompromised individuals and young children (Doyle et al., 2000). No evidence of cross-reactivity with other viral infections, such as rubella, mumps, varicella-zoster virus, cytomegalovirus, herpes simplex virus-1 (HSV-1) and HSV-2, is apparent when this immunoassay is used in clinical settings. Previous studies have reported cross-reactivity with rubella in several commercial B19 IgM assays (Sloots & Devine, 1996; Tolfvenstam et al., 1996) and as the symptoms of rubella infection are similar to those of B19 infection, this was a cause for concern, particularly in the diagnosis of infection in pregnant women. A false-positive rate of 5% was reported when specimens from healthy volunteers were analysed with a range of commercially available B19 IgM immunoassays, probably due to cross-reactivity and lack of specificity in these immunoassays (Tolfvenstam et al., 1996).

Detection of B19 NS1 IgM has received little attention as a marker of recent infection by B19. Ennis et al. (2001) observed that 27.5% (11/40) of specimens that were B19 VP2 IgM-positive also contained B19 NS1 IgM when tested by ELISA. Interestingly, when these samples were analysed by Western blot, there was no evidence of NS1 IgM reactivity, which indicates that conformational epitopes are important for detection.

### B19 IgG immunodetection

Development of B19 IgG antibodies coincides with a decline in the IgM response. IgG reactivity against conformational epitopes of VP1 and VP2 persists post-infection; however, for both capsid proteins, reactivity against linear epitopes declines post-infection [abruptly against VP2, but more slowly against VP1 (Söderlund et al., 1995; Kaikkonen et al., 1999; Kerr et al., 1999; Manaresi et al., 1999)]. Antibody reactivity against linear VP2 epitopes usually disappears within 6 months of B19 infection (Söderlund et al., 1995). This initial reactivity against linearized VP2 appears to be directed predominantly against a heptapeptide (amino acids 344–350) that was identified by analysis of acute-phase sera (Kaikkonen et al., 1999).

Although the antibody response wanes against linear epitopes on B19 capsid proteins, it persists against conformational epitopes of both capsid proteins. The only FDA-cleared B19 IgG immunoassay that is available as a marker of past infection is a microplate immunoassay that utilizes capsid VP2 to detect B19 and erythrovirus V9 IgG (Heegaard et al., 2002b; A. Garbarg-Chenon, personal communication). This baculovirus-based immunoassay has been compared to another commercially available, E. coli-based VP1 immunoassay for detection of B19 antibodies in the sera of pregnant women (Jordan, 2000). A number of equivocal results were obtained by using the B19 VP1 immunoassay. To verify results, samples were tested by using a commercially available VP1 immunofluorescent assay (IFA). This assay examines seroreactivity against conformationally intact VP1. Although this assay may be somewhat subjective, as it primarily measures the degree of specimen fluorescence, when used in parallel with the VP2 IgG immunoassay, it can confirm B19 reactivity. Results from the VP1 IFA were similar to those obtained by the baculovirus VP2 EIA and were in accordance with the fact that many of the samples that were found to be equivocal by the E. coli VP1 EIA had clinical histories of B19 exposure. Availability of a B19 IgG International Standard (2nd International Standard 2003; code 01/602; 77 IU per ampoule) should further assist in accurate confirmation of past B19 infection by standardizing B19 IgG determination from different laboratories that use a variety of test systems (Ferguson et al., 1997; Searle et al., 1997).

Recently, the importance of antibodies against the B19 non-structural protein NS1 has been investigated, with a view to improving diagnosis of B19 infection. The presence of B19 NS1 IgG was thought to be associated primarily with persistent B19 infection (von Poblotzki et al., 1995a, 1995b); however, several groups have subsequently found no significant difference between the level of NS1 IgG in control patients with past infection and those with chronic B19 infection (Searle et al., 1998; Venturoli et al., 1998; Jones et al., 1999). Mapping of B-cell epitopes on NS1 identified three antigenic regions (amino acids 191–206, 271–286 and 371–386) that were equally reactive with sera from healthy individuals with past B19 infection and patients who were infected persistently by B19 (Tolfvenstam et al., 2000). More recently, by using an E. coli expression system, NS1 IgG reactivity has been shown to be most prevalent in serum, following recent infection in pregnant women (61%) (Hemauer et al., 2000). These findings are supported by the work of Mitchell et al. (2001), who examined NS1 IgG reactivity in sera from individuals who were either infected by B19, had been exposed to B19 but were not infected, were suffering from a rash illness or chronic arthropathy, or were healthy controls. NS1 IgG reactivity was predominant in recently infected specimens and when follow-up samples from these individuals were analysed, the level of NS1-specific IgG reactivity had declined. In addition, there was no evidence of a connection between NS1 IgG antibodies and the development of arthropathy (Mitchell et al., 2001). The NS1-specific IgG response wanes post-infection as the virus is cleared from the body; therefore, NS1 IgG reactivity may have some value as a marker of recent infection, in conjunction with the detection of IgG against linear epitopes on VP2 (Ennis et al., 2001). This study demonstrated that 69% of children who had been infected recently by B19 were NS1
IgG-seropositive (Ennis et al., 2001). Heegaard et al. (2002a) also observed a seroprevalence of 60% B19 NS1 IgG in recently infected individuals (<6 weeks post-infection) and suggested that NS1 IgG detection may significantly improve immunoassay sensitivity.

It is now clear that B19 IgM and IgG detection is optimal in immunoassays that utilize VP2 capsids for antibody detection. Antibody (IgG/M) detection of B19 NS1 protein may assist in the confirmation of recent B19 infection, when used in combination with VP2 capsid-based immunoassays. Erythrovirus V9 antibody detection is also feasible, by using immunoassays that are based on B19 VP2 capsids.

**Cell-mediated immunity**

Cell-mediated immunity to B19 has not been studied extensively; this is due primarily to the fact that the humoral response was thought to be most important in combatting B19 infection. Indeed, initial attempts to demonstrate specific T-cell proliferative responses to B19 were unsuccessful (Kurtzman et al., 1989a) and, for some time, this work supported the prevailing theory that neutralizing antibody production was the major mechanism of immunity in B19. In 1996, *ex vivo* B19-specific CD4+ T-cell responses were first detected against *E. coli*-expressed VP1, VP2 and NS1 antigens (von Poblotzki et al., 1996). T-cell responses of 16 individuals were analysed (ten seropositive and six seronegative blood donors), none of whom had any evidence of acute infection. The majority (90%) of seropositive donors who were stimulated *ex vivo* by VP2 displayed specific T-cell responses, with 80% displaying VP1-specific responses. There was no significant difference in T-cell proliferation for NS1 between seropositive and seronegative individuals. Upon inclusion of mAbs that were specific for class I and class II HLA, it was found that HLA class II-specific antibodies inhibited T-cell proliferation, thus indicating that the effector T-cell population of B19 are CD4+ cells. Subsequent peripheral blood mononuclear cell (PBMC) depletion of either CD4+ or CD8+ T cells and stimulation of the remaining population confirmed this observation.

More recently, significant *ex vivo* T-cell reactivity was observed in PBMCs of recently and remotely infected individuals by using a B19 candidate vaccine (Franssila et al., 2001) and also the B19 recombinant proteins, VP1 and VP2 (Corcoran et al., 2000). T cells from recently infected individuals responded strongly to the B19 capsids, giving a mean T-cell stimulation index (SI) of 36 (Franssila et al., 2001). Blood donors with past infections gave comparable rates of T-cell stimulation. Seronegative individuals had SI values of about 3-3 and this study also showed that the responding population of T cells were CD4+. Although von Poblotzki et al. (1996) saw no difference in T-cell responses to NS1 in seronegative and seropositive individuals, significant responses to this antigen have been reported in recently infected individuals and patients who developed chronic arthralgia following B19 infection (Mitchell et al., 2001). T-cell responses to NS1 were not seen in healthy individuals with past B19 infection, except for two individuals who were also NS1 IgG-seropositive.

Cellular immune response to an epitope of NS1 that is recognized specifically by CD8+ T cells was investigated recently by using major histocompatibility complex tetrameric complex binding (Tolfvenstam et al., 2001b). The response of 21 individuals to this epitope was examined in healthy volunteers and human immunodeficiency virus (HIV)-1 infected adults and children. Sixteen of the volunteers were HLA-matched (HLA B35) and six were mismatched. Sixty-three per cent of matched individuals displayed specific CD8+ T-cell responses. Seventy-two per cent of matched individuals in the same cohort exhibited specific T-cell responses by using an interferon γ (IFN-γ) ELISpot assay. The level of B19-specific CD8+ T cells was similar among healthy and HIV-infected individuals. The results presented in this report showed the important cellular role of cytotoxic T cells in combatting B19 infection (Tolfvenstam et al., 2001b). B19-specific T-cell responses may now represent a novel method for confirming past B19 infection.

Recent evidence shows the importance of evaluating T-cell responses in understanding the nature of B19 infection. Chen et al. (2001) have identified an AIDS patient with persistent B19 infection who showed an initial remission of B19 infection. This remission was evident despite the lack of a specific antibody response, thus indicating a role for cellular immunity in combatting B19 infection. NS1-reactive lymphocytes have been detected in two B19-seronegative individuals who were exposed to the virus, indicating a possible subclinical B19 infection or perhaps a loss of antibodies against capsid proteins (Mitchell et al., 2001). The importance of cellular immunity in B19 was further emphasized in a report by Tolfvenstam et al. (2001b). Here, investigations of B19-specific CD8+ T-cell responses identified two healthy adults and two HIV-1-infected patients who were seronegative for B19, with specific T-cell responses against B19 by either IFN-γ ELISpot or tetramer binding studies, thus implying the presence of a cellular response in the absence of a humoral response.

Significant T-cell transcriptional activation has been reported in a patient with acute B19 infection, causing increased levels of interleukin (IL) 1β, IL6 and IFN-γ mRNA (Wagner et al., 1995). A subsequent study that analysed the sera of patients who were infected acutely by B19 showed that although IL1β, IL6, IFN-γ and tumour necrosis factor α (TNF-α) were secreted during the acute phase of infection, increased levels of both IFN-γ and TNF-α persisted and were detectable for 2–37 months later, during a follow-up study (Kerr et al., 2001). It has also been suggested that cytokine genetic polymorphisms may, in some way, affect the development of symptoms during B19 infection. To date, the transforming growth factor β (TGFβ) allele has been associated with skin rash at acute infection and the IFN-γ allele has been associated with NS1 antibody development (Kerr et al.,
In a study of recently infected children, it was shown that although strong T-cell proliferative responses were evident to both capsid proteins, production of the T helper (Th1) cytokine IFN-γ, but not of IL2, was impaired when compared to convalescent adults (Corcoran et al., 2000). In addition, *ex vivo* production of IFN-γ and IL2 that was observed in B19-seropositive pregnant women was lower than observed previously for healthy, non-pregnant individuals, suggesting a possible diminution of the maternal anti-virus immune response that may subsequently increase the risk of fetal B19 infection (Corcoran et al., 2003). Expression of the non-structural protein NS1 causes the production of increased levels of the inflammatory cytokine IL6 in a number of cell lines, including hematopoietic cell lines and human umbilical vein endothelial cells (Moffatt et al., 1996). IL6 is known to be involved in synovial cell proliferation and, in addition, high levels of IL6, along with other inflammatory cytokines, have been found in inflamed joints of patients with RA, which would suggest an association between IL6 production and the joint manifestations that are observed with B19 infection (Bataille et al., 1995). IL6 involvement in RA is supported by the fact that antibodies against IL6 cause inhibition of RA manifestations (Bataille et al., 1995). As well as increased IL6 production, high levels of IFN-γ, TNF-α and IL8 have been detected in the sera of infants with B19-associated acute myocarditis (Negro et al., 2000). IL2 production at the maternal–fetal interface in women who seroconverted to B19 during pregnancy is thought to determine the outcome of the pregnancy, with high levels of IL2 on the fetal side being associated with pregnancies that result in a poor outcome (Jordan et al., 2001).

**B19 vaccine**

Ballou et al. (2003) have shown recently that a recombinant vaccine (MEDI-491; Medimmune) that is comprised of B19 VP1 and VP2 capsid proteins could elicit neutralizing antibody titres in volunteer adults (*n* = 24). Sera from immunized individuals were also shown to be capable of inhibiting B19 replication *in vitro*. The efficacy of this formulation in preventing infection by B19 remains to be established; nonetheless, it is an encouraging and welcome advance in the fight against this ubiquitous pathogen.

**PCR detection of B19 DNA**

Many clinical laboratories now complement B19 antibody screening with diagnostic PCR and it is well-established that B19 PCR improves the sensitivity of detection of B19 infection (Skjøldebrand-Sparre et al., 2000; Manaresi et al., 2002). However, caution must be exercised with regard to the deployment of B19 PCR for a number of reasons: (i) the high viraemia that is associated with B19 infection, along with resultant amplicon load, may cause PCR false positivity – particularly when nested PCR is used for B19 detection; (ii) B19 DNA detection may not always be indicative of an acute infection; (iii) many PCR assays use in-house primer pairs of undefined sensitivity of detection; (iv) false negativity may be observed with respect to non-B19 strains (e.g. erythrovirus V9, K71 or A6), due to minor sequence differences (Hokynar et al., 2002; Nguyen et al., 2002; Servant et al., 2002); and, finally, (v) many extraction methods are suitable for DNA purification from serum or plasma only and not from solid tissue (e.g. placenta or fetal tissue). Notwithstanding these caveats, some of which are discussed in more detail below, B19 PCR is an important tool in the technologist’s armoury for detection of B19 infection.

During acute infection with B19, viral titres can reach ∼10¹² genome equivalents ml⁻¹ (Prowse et al., 1997). In the immunocompetent host, virus DNA is detectable for at least 1 month post-infection (Erdman et al., 1991). In chronic B19 infection, virus DNA can persist in the host without the presence of B19 IgM or IgG (Kurtzman et al., 1988; Frickhofen et al., 1990). However, it has also been shown that B19 DNA can persist in healthy, immunocompetent individuals at low levels for long periods (Cassinotti et al., 1993; Kerr et al., 1995b; Musiani et al., 1995; Cassinotti & Siegl, 2000). Thus, B19 DNA, detected by qualitative PCR analysis, is not always indicative of recent infection. Cassinotti & Siegl (2000) used quantitative PCR to follow the amount of specific B19 DNA in an immunocompetent patient, expressed as genome equivalents ml⁻¹, from the time of acute B19 infection until convalescence. A series of samples was taken over a 1 year period and was analysed by using a real-time fluorogenic PCR assay. During the viraemic stage of B19 infection, viral load reached levels of 8.8 × 10⁹ genome equivalents (ml blood)⁻¹. At this stage, the patient was positive for specific IgM and negative for IgG reactivity. At week 164, viral load had declined to 95 genome equivalents ml⁻¹, IgM reactivity was lost and conformational IgG reactivity was strong. Specimens taken after this time-point were undetectable for B19 DNA. Thus, whilst the actual amount of circulating B19 DNA that was present following B19 infection diminished dramatically after the first few weeks of infection, it persisted for some time before being cleared from the host, despite the development of circulating B19 IgG. This slow rate of B19 DNA clearance from an immunocompetent host could impact negatively on PCR as a diagnostic tool in differentiating between recent or chronic B19 infection in a situation where a qualitative PCR assay of unspecified sensitivity of detection was employed. However, with the introduction of the World Health Organization (WHO) International Standard for Parvovirus B19 DNA (NIBSC 99/800), PCR assay standardization has become possible (Saldanha et al., 2002). Using the WHO standard, a compatible PCR-ELISA that can detect levels as low as 1.6 × 10⁴ IU B19 DNA ml⁻¹ was established (Daly et al., 2002) and, by using real-time PCR technology, a sensitivity of detection of 15.4 IU ml⁻¹ (10 Baxter units ml⁻¹) (Aberham et al., 2001) was reached. Müller et al. (2002) and Thomas et al. (2003) have also described standardized B19 PCR assay systems. These standardized methods could be used not only in a diagnostic setting, but also for rapid screening of plasma mini-pools and blood products, thereby leading to determination of the amount of B19 DNA present and improved product safety.

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PCR may also be used in screening for the newly isolated erythrovirus V9 (Nguyen et al., 1998, 1999). A nested PCR assay that is capable of accurately detecting V9 and B19 DNA simultaneously, comprising a primary round of amplification by using a pair of consensus primers and a subsequent round of amplification by using separate primers for B19 and V9, has been developed (Heegaard et al., 2001). By using this PCR assay, clinical samples, including 100 B19 IgM-positive specimens and untreated plasma pools that represented 100 000 blood donor units from the Danish population, were screened for both V9 and B19 DNA. None of the specimens analysed were positive for V9 DNA, which may be due to the facts that this V9 isolate is an emerging virus and that this erythrovirus may actually be more divergent than thought previously (Heegaard et al., 2001). Thus, PCR could be used as a diagnostic tool for the identification of possible new erythrovirus isolates in cases where patients are negative for B19 DNA, but have displayed clinical symptoms of B19 infection.

B19 and blood-product safety

B19 can be transmitted through blood transfusions and plasma-derived products (Prowse et al., 1997; Santagostino et al., 1997). Screening of blood donations for the presence of B19 DNA is not routine (Blümel et al., 2002), despite the fact that this virus is highly resilient and, like the hepatitis A virus, can withstand denaturation, even at high temperatures (Santagostino et al., 1994). In fact, B19 can withstand processes that involve solvent/detergent treatment, lyophilization and temperatures of 100 °C for 30 min and, despite these harsh virucidal processes, still have the capacity to contaminate factor VIII and factor IX concentrates (Santagostino et al., 1997). B19 contamination of such purified blood products is particularly problematic as, in the absence of B19 IgG, the infectious potential of B19 may be enhanced (Blümel et al., 2002). The most recent determination of B19 prevalence is 1 in 625 blood donations (n = 16 859; range, 10^2 to >10^7 IU B19 DNA ml⁻¹) (Thomas et al., 2003). Previously, B19 had been estimated to be present in 1:16 000 transfusions, based on the mean incidence of B19 infection in a non-epidemic period (320 cases per 100 000 population) and the fact that viraemia lasts for about 7 days (Prowse et al., 1997). During epidemics, the incidence of viraemia in donations is greatly increased, with levels as high as 1:3790 reported in Ireland (O’Neill & Coyle, 1992) and 1:167 in Japan (Yoto et al., 1995).

The infectious level of B19 in blood products has yet to be established with certainty and is likely to depend on the level of B19 IgG that is co-present in the product, in addition to recipient immune status. As part of a phase IV study, a group of 100 healthy volunteers who were seronegative for B19 were given 1 unit of plasma that had been solvent/detergent-treated (Davenport et al., 2000). Of the volunteers who were screened subsequently for incidences of B19 infection, 18 % had seroconverted over the subsequent 3 months. Three of the ten batches of plasma that were used in the study were found retrospectively to contain high levels of B19 DNA (>10^7 genome equivalents ml⁻¹) and these batches coincided with the plasma that was administered to the volunteers who seroconverted. Interestingly, batches with low amounts of B19 (<10^4 genome equivalents ml⁻¹) did not cause B19 seroconversion. Upon discovery of these facts, the company that was involved in manufacturing this plasma (VITEX, Watertown, MA, USA) voluntarily recalled batches that were associated with B19 transmission and reviewed their screening process. Presently, plasma lots that contain high levels of B19 are eliminated from manufacturing batches of plasma. Thus, there is a level of virus, as yet undetermined, that will not cause B19 infection. Notably, Daly et al. (2002) undertook a retrospective study of plasma pools (n = 30) that were similar to those utilized in the study of Davenport et al. (2000) and found B19 IgG levels in the range of 64.7 ± 17.5 IU ml⁻¹. Thus, it is possible that this level of B19 IgG may be capable of preventing recipient B19 infection when transfused with plasma that is contaminated by low levels of B19 (<10^4 genome equivalents ml⁻¹). Blümel et al. (2002) have identified two incidences of B19 transmission by separate lots of clotting factor concentrates and have shown that B19 levels of 8.6 × 10^6 genome equivalents ml⁻¹ (volume, 180 ml) and 4 × 10^5 genome equivalents ml⁻¹ (volume, 966 ml) were responsible for seroconversion.

B19 seroprevalence is higher among haemophiliacs than the general population, presumably due to the fact that products that are derived from pooled plasma are more likely to contain infectious B19 DNA. This was shown conclusively when B19 seroprevalence in a haemophilic population was compared with that of normal healthy individuals and it was discovered that haemophilic children who were treated with virally inactivated clotting factor had an increased level of B19 seroprevalence (92 %) (Eis-Hübinger et al., 1996). Normally, in the general population, a continuous increase in B19 seroreactivity is observed with age, with a seroprevalence for adults over 60 that reaches about 72 %. Interestingly, the haemophilic population that was treated pre-1984 when transfused with plasma that is contaminated by low levels of B19 (<10^4 genome equivalents ml⁻¹) did not develop high B19 IgG levels, whereas the level of B19 seroprevalence in a haemophilic population was significantly increased (92 %) (Eis-Hübinger et al., 1996). Because of this, the Clotting Factor IX concentrates used in the study by Eis-Hübinger et al. (1996) were manufactured prior to 1984, and thus contained low levels of B19 DNA.

Despite the fact that B19 infection can be transmitted via contaminated blood products, there are presently no strict regulatory prerequisites governing B19 contamination of pooled plasma or blood products prior to product release. However, it should be acknowledged that many manufacturers now perform B19 PCR on plasma mini-pools, in order to eliminate high B19 viral load plasma (Aberham et al., 2001). PCR screening of blood products has been shown to facilitate removal of 23 B19 PCR-positive donations from a plasma pool of 6000, resulting in a 10–100-fold decrease in viral load (Prowse et al., 1997). However, the level of B19 DNA does not necessarily correlate to the rate of infectivity, as has been demonstrated previously for the canine parvovirus; in this case, heat treatment was found to reduce canine parvovirus infectivity by >100-fold, despite the fact that the PCR assay titre was unaltered (Hart et al., 1994).
Nonetheless, the issue of whether high-risk populations, such as pregnant women, immunocompromised patients and people with chronic anaemia, should undergo administration of any B19-containing products while the level of infectious B19 DNA is unknown and mini-pool screening is not mandatory must be addressed. The aforementioned availability of an International Standard preparation of B19 DNA (Saldanha et al., 2002), in addition to a number of compatible and quantitative B19 PCR detection systems (Aberham et al., 2001; Daly et al., 2002; Knöll et al., 2002; Müller et al., 2002; Thomas et al., 2003), should alleviate problems caused by ambiguity between results from laboratories that use various methods of measuring and expressing B19 DNA levels and help to determine the infectious dose for B19.

It is highly significant that, in the Netherlands, a recommendation has been made that individual donor screening for B19 IgG to identify individual donors with continually high antibody levels (at t = 0 and 6 months) should be initiated (Health Council for the Netherlands, 2002; Groeneveld & van der Noordaa, 2003). Selected individuals who maintain high B19 IgG levels will subsequently form a panel of plasma or blood-product donors for high-risk recipients, such as immunocompromised individuals, thereby minimizing the risk of B19 transmission from acutely infected, although asymptomatic, donors. It is our view that the introduction of such a screening algorithm sets the standard for blood-product safety in the future, specifically with respect to minimizing the risk of B19 transmission.

Assays that are based on exploitation of the P antigen receptor of B19, known as receptor-mediated haemagglutination, have been proposed as a cheap way to screen plasma or blood-product donors for high-risk recipients, such as immunocompromised individuals, thereby minimizing the risk of B19 transmission from acutely infected, although symptomatic, donors. It is our view that the introduction of such a screening algorithm sets the standard for blood-product safety in the future, specifically with respect to minimizing the risk of B19 transmission.

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References


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Fig. 2. Schematic depiction of T- and B-cell response to parvovirus B19 infection. Upon B19 infection, B cells divide to produce plasma cells and memory cells. Plasma cells secrete IgM antibodies that are specific for both conformational (N) and linear (D) epitopes of B19, which are detectable approximately 7 days post-infection. B19-specific IgG is detectable about 15 days post-infection and is directed initially against both linear and conformational epitopes of the capsid proteins (VP1 and VP2) and, to a lesser extent, against NS1, but declines against linear epitopes of the proteins in a time-dependent manner. Infection by B19 most likely confers lifelong protection on the host, due to the development of memory B cells that are specific for conformational regions of the B19 capsid and also linear regions of the VP1 protein (Corcoran et al., 2004). Antigen-presenting cells (APC) process the virus and display B19 peptides on their surface to Th cells. These Th cells then secrete cytokines that play a role in mediating anti-virus immunity and may also be associated with pathogenesis of B19 infection (e.g. IL6 is associated with RA).
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