# Clinical features and laboratory findings of human parvovirus B19 in human immunodeficiency virus-infected patients

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Immunocompromised patients may develop severe chronic anaemia when infected by human parvovirus B19 (B19V). However, this is not the case in human immunodeficiency virus (HIV)-infected patients with good adherence to highly active antiretroviral treatment (HAART). In this study, we investigated the clinical evolution of five HIV-infected patients receiving HAART who had B19V infections confirmed by serum polymerase chain reaction. Four of the patients were infected with genotype 1a strains and the remaining patient was infected with a genotype 3b strain. Anaemia was detected in three of the patients, but all patients recovered without requiring immunoglobulin and/or blood transfusions. In all cases, the attending physicians did not suspect the B19V infections. There was no apparent relationship between the infecting genotype and the clinical course. In the HAART era, B19V infections in HIV-positive patients may be limited, subtle or unapparent.

Key words: HIV - human parvovirus B19 - seroconversion

Parvovirus B19 (B19V), a member of the genus *Ery-throvirus* within the Parvoviridae family, has been grouped into three distinct genotypes: (i) genotype 1, with subtypes 1a (the prototypic virus) and 1b, (ii) genotype 2 (A6 and LaLi strains) and (iii) genotype 3, with subtypes 3a (V9 strains) and 3b (D91.1 strains) (Nguyen et al. 1999, 2002, Servant et al. 2002, Fauquet et al. 2005).

B19V primarily infects erythroid cells, leading to transient inhibition of erythropoiesis. In immunocompetent individuals, it usually causes an acute and self-limited childhood disease known as erythema infectiosum (EI) ("slapped cheek" rash or 5th disease). Adults with EI (particularly females) frequently present with joint symptoms. However, most patients are asymptomatic (Woolf et al. 1989). Immunocompromised patients who are unable to produce neutralising antibodies may develop severe chronic anaemia. In human immunodeficiency virus (HIV)-infected patients with residual immunity, the clinical manifestations, if any, are those of fifth disease (Frickhofen & Young 1990). With the advent of highly active antiretroviral treatment (HAART), several studies, including one from our group (de Azevedo et al. 2012), have shown a decrease in cases of anaemia caused by B19V (Mylonakis et al. 1999, Ware & Moore 2001).

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To better understand the importance of B19V infection in the HAART era, a study to estimate the frequency of B19V seroconversion in a cohort of HIVinfected patients was conducted by our group during an eight-year period (2001-2008) at the Infectious Diseases Department of Antonio Pedro University Hospital (HUAP) at the Fluminense Federal University (Niterói, state of Rio de Janeiro, Brazil) (de Azevedo et al. 2009). Seroconversions were detected in approximately 30% (28 of 88) of our anti-B19 IgG-negative patients, a similar proportion to that found by others (Chernak et al. 1995, Mylonakis et al. 1999). Most seroconversions occurred during incidence peaks of a B19V infection in Niterói (Oliveira et al. 2005, de Azevedo et al. 2012). All sera from the 88 patients were tested by polymerase chain reactions (PCRs) and B19 DNA was detected in five of the patients, four of whom also exhibited seroconversion. This paper describes the clinical and laboratory findings of B19V infections in these five HIV-infected patients.

# PATIENTS, MATERIALS AND METHODS

Data, such as haemoglobin (Hb) concentration, CD4<sup>+</sup> T cell counts and B19V IgG and IgM serology, were retrieved from a previous study by our group (de Azevedo et al. 2012). ELISAs (Biotrin International<sup>TM</sup>, Dublin, Ireland) were performed, according to the manufacturer's instructions, to detect IgG and IgM antibodies to B19 in all sera. The following definitions were used in this study: (i) anaemia was defined according to the World Health Organization (WHO 2001) criteria as a Hb concentration below 13 g/dL in men and below 12 g/dL in women and (ii) severe anaemia was defined as a Hb concentration below 7 g/dL. PCRs were performed to detect and genotype B19V. Briefly, viral DNA was extracted from serum samples of 88 HIV-positive patients using a QIAamp DNA Blood Mini Kit (QIAGEN, Hilden, Germany) according to the manufacturer's instructions. For the screening of B19V DNA, PCR was performed using the primers E1905F (nt 1905-1923) and E1987R (nt 2007-1987), which amplify a 102-bp fragment of the NS1 gene as previously described by Nguyen et al. (2002), with some modifications (de Azevedo et al. 2012). This PCR can routinely detect as few as 20 copies of the B19V DNA (Nguyen et al. 2012).

To genotype the B19V strains detected in HIV-positive patients, a semi-nested PCR using the primer pairs P12/P16 (4127-4689) and P13/P16 (4214-4689) for partial amplification of the VP1/VP2 capsid gene was performed as described by Durigon et al. (1993). The 476bp amplicons were purified using GFX<sup>TM</sup> PCR DNA and the Gel Band Purification kit (GE, Healthcare, UK) and were then subjected to direct sequencing using the BigDye terminator v. 1.1 cycle sequencing kit (Applied Biosystems, CA, USA) (Otto et al. 2008).

Sequences were retrieved and analysed by the Bio-Edit sequence alignment editor, v. 7.0.1 (mbio.ncsu.edu/ BioEdit/bioedit.html) and they were compared with the following sequences available in GenBank (Hall 1999): genotype 1a: M24682-Wi (UK); genotype 1b: DQ35706 and DQ35705 (Ghana); genotype 2: AY06477-A6 (Denmark) and AY04416-LaLi; genotype 3a: NC0049-V9 (Italy); genotype 3b: Z68146 (UK), AF162273 (Italy), AY083234-D01.1 (France), DQ234778, DQ234779.2 (Ghana), EF089227, EF089229.1 (Brazil/PA) and AY647977.1 (Brazil/SP).

*Ethics* - Written informed consent was obtained from all volunteers and the project was approved by the Hospital Review Board (CMM/HUAP 134/05).

#### RESULTS

Case 1 - A male patient diagnosed with acquired immune deficiency syndrome (AIDS) in 1996 at 41 years of age, with poor adherence to HAART. The observation period started in December 2001 (CD4+ T cell count: 158 cells/mm3; anti B19V IgG: negative). There was a gradual decrease in CD4<sup>+</sup> T cell counts, which reached 3 cells/ mm<sup>3</sup> in August 2004. In February 2005, the patient was hospitalised because of fever, hepatosplenomegaly, severe anaemia (Hb: 5.9 g/dL) and multiple opportunistic infections (oro-oesophageal candidiasis, perianal condylomata and herpes). The patient underwent a bone marrow biopsy (February 2005) and was started on empirical treatment for atypical mycobacteriosis. Clinical improvement was observed and the patient did well afterward, which was attributed to a better adherence to HAART (lamivudine, stavudine, efavirenz and lopinavir/ritonavir). The biopsy report described "overall cellularity of 60%, with presence of the three series. The myeloid/ erythroid ratio was 2:1, with predominance of immature forms, numerous myeloblasts and moderate increase in the number of red blood cells with megaloblastic maturation. Megakarvocvtes were observed in the usual number, but with frequent atypical forms. There were rare

erythroblasts with nuclear viral inclusions consistent with B19V. Conclusion: myelodysplastic changes secondary to HIV infection, compatible with B19V infection". The patient received neither immunoglobulin therapy nor a blood transfusion. There was a gradual recovery from anaemia during 2005. In November 2005, stavudine was replaced by zidovudine to improve adherence. In December 2005, seroconversion (anti-B19V IgG-positive and anti-B19V IgM-positive) and the presence of B19V DNA (strain RJ 011/2005, genotype 1a) were both detected in association with an increase in the CD4<sup>+</sup> T cell count (34 cells/mm<sup>3</sup>) and resolution of anaemia (Hb: 13.9 g/dL) (Table I). A subsequent serum sample (June 2006) was negative for B19V DNA.

*Case 2* - A female patient diagnosed with AIDS in 1997 at 54 years of age, when she was started on HAART with good adherence to therapy and immune recovery. The observation period began in December 2001 (CD4<sup>+</sup> T cell count: 598 cells/mm<sup>3</sup>). Seroconversion to B19V (anti-B19 IgG) was observed in June 2005. There is no record of blood counts during the period of seroconversion (November 2004-June 2005). During this period, there were no complaints or changes in physical examination suggestive of anaemia. A serum sample collected in June 2005 (strain RJ 026/2005, genotype 1a) was B19V DNA-positive by PCR. The next available blood sample (September 2005) remained B19V DNA-positive (Table II), but the subsequent sample (January 2006) was negative for B19V DNA.

*Case 3* - A woman diagnosed with AIDS in 1996 at 36 years of age, when she was started on HAART, followed by immunological recovery. The observation period began in January 2002 (CD4<sup>+</sup> T count: 520 cells/mm<sup>3</sup>). B19V seroconversion (anti-B19 IgM and anti-B19 IgG) was observed in September 2005 (CD4<sup>+</sup> T count: 649 cells/mm<sup>3</sup>). At that time, there were no complaints or physical examination changes suggestive of anaemia. A serum sample collected in September 2005 (strain RJ 071/2005, genotype 3b) was B19V DNA-positive (Table III). Two subsequent samples collected in June 2006 and September 2006 were, respectively, positive and negative for B19V DNA.

*Case 4* - A man diagnosed with AIDS in November 2003 at 33 years of age. At that time, he presented with mild anaemia (Hb = 12.8 g/dL) attributed to HIV infection. He began HAART in September 2004. The observation period started in November 2003 (CD4<sup>+</sup> T cell count: 215 cells/mm<sup>3</sup>). B19V seroconversion (anti-B19 IgG) was observed in April 2007 (CD4<sup>+</sup> T cell count: 504 cells/mm<sup>3</sup>). Hb levels below 13.0 g/dL, suggestive of mild anaemia, were observed from November 2003-July 2006, before the period of seroconversion (anti-B19 IgG) (July 2006-April 2007). A sample collected in July 2006 (strain RJ 350/2006, genotype 1a) was B19V DNApositive (Table IV). A subsequent sample (April 2007) was negative for B19V DNA.

*Case 5* - A man diagnosed with AIDS in November 2001 at 35 years of age. The observation period began in January 2002 (CD4<sup>+</sup> T cell count: 80 cells/mm<sup>3</sup>),

					TABLF Laboratory data	E I a of Case 1					
Month/year	Dec/2001	Mar/2002	Aug/2002	Jul/2003	Feb/2004	Aug/2004	Feb/2005	Mar/2005	Jul/2005	Sep/2005	Dec/2005
IgG anti-B19V	NEG		NEG	NEG	NEG	NEG					POS
IgM anti-B19V	NEG	,	NEG	NEG	NEG	NEG	,	ı	ı	·	POS
PCR-B19	I	ı	I	I	I	NEG	ı	ı	ı	ı	POS
CD4 <sup>+</sup> (cells/mm <sup>3</sup> )	158	ı	118	45	6	6	ı	ı	1	ı	34
Hb (g/dL)	·	13.1		ı	·		5.9	6.9	10	10.5	13.9
B19V: parvovirus B	19; Hb: haemo	globin; NEG: 1	negative; PCR:	polymerase ch	nain reaction; P	OS: positive; -:	lack of data in	the medical rec	cords.		
					TABLE	П					
					Laboratory data	a of Case 2					
Month/year	Dec/2001	May/20(	0ct//	2002 0	)ct/2003	Jun/2003	Jul/2004	Nov/20	004 Ju	n/2005	Sep/2005
IgG anti-B19V	NEG	ı	IN	ĐE	NEG	NEG	NEG	NEC	5	POS	POS
IgM anti-B19V	NEG		IN	BG	NEG	NEG	NEG	NEC		NEG	NEG
PCR-B19			·		ı	ı	I	NEC	<b>ר</b> ז	POS	POS
CD4 <sup>+</sup> (cells/mm <sup>3</sup> )	589	'	24	41	474	702	712	769		613	702
Hb (g/dL)	·	12.6	13	3.5						ı	13
B19V: parvovirus B	19; Hb: haemo	globin; NEG: r	legative; PCR:	polymerase ch	ain reaction; P(	OS: positive; -: ]	ack of data in t	he medical rec	cords.		
					TARLF	III					
					Laboratory data	a of Case 3					
Month/year	Jan/2002	Oct/200	2 Dec/	2002	Apr/2003	Sep/2003	May/2004	Aug/2(	004 Fe	b/2005	Sep/2005

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B19V: parvovirus B19; Hb: haemoglobin; NEG: negative; PCR: polymerase chain reaction; POS: positive; -: lack of data in the medical records.

POS POS POS 649

NEG NEG NEG 610

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CD4<sup>+</sup> (cells/mm<sup>3</sup>)

Hb (g/dL)

NEG

NEG

lgG anti-B19V IgM anti-B19V PCR-B19

NEG

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when HAART (zidovudine, lamivudine and efavirenz) was started and it ended in October 2005 (CD4<sup>+</sup> T cell count: 244 cells/mm<sup>3</sup>). The patient had several opportunistic infections and anaemia during the follow-up and a decrease in Hb level (5.9 g/dL) was observed in June 2002. In addition, CD4<sup>+</sup> T cell counts less than 100 cells/mm<sup>3</sup> were observed until November 2002. As the first test for the presence of IgG antibodies was indeterminate or weakly positive, these samples were tested again and found to be negative. Nevertheless, B19 DNA was detected in one of the samples (November 2002, strain RJ 109/2002, genotype 1a). At this time, the Hb level was 9.14 g/dL and, thus, no longer compatible with severe anaemia (Table V). There is no record of a B19V outbreak during 2002/2003 (Oliveira et al. 2005).

According to sequence analysis of the VP1/VP2 capsid region, the strains RJ 011/2005, RJ 026/2005, RJ 109/2002 and RJ 350/2006 shared high nucleotide (nt) identity (100-99%) with 1a B19V isolates from other countries (EU78564, M2468, M13178, EU478541, EU14278 and JN211184). One strain (RJ 071/2005) displayed 99.3-97.2% nt identity with 3b B19V Brazilian isolates (EF089227 and EF089229). Based on their nt identity with other B19V isolates previously described, four of the B19V strains in this study were characterised as belonging to genotype 1a and one B19V strain was characterised as belonging to genotype 3b.

# DISCUSSION

This is the first study to analyse clinical manifestations in HIV-infected patients during an outbreak of B19V in Brazil. Persistent B19 infection has been reported to be associated with HIV infection, possibly due to inability to mount a protective B19-specific antibody response (Chernak et al. 1995, van Elsacker-Niele et al. 1996) and some reports have shown that complete remission of B19-associated pure red cell aplasia can be achieved just by treating patients with HAART (Mylonakis et al. 1999, Ware & Moore 2001).

Our findings show that the impact of restoring the immune response by either starting or continuing HAART may explain the reduction and even the elimination, of severe haematological changes caused by B19V infections, even before a detectable increase in the number of CD4<sup>+</sup> T cells in the blood occurs (Cases 1, 4 and 5). Although anaemia had been detected in three patients (2 of whom had severe anaemia), all of the patients recovered and none required immunoglobulin and/or blood transfusions, including the two patients for whom no data existed in the medical records regarding the occurrence of anaemia. Additionally, our results also show that B19 infection may be apparent only in retrospective analyses. as there are many causes of anaemia in HIV-infected patients, including co-infection by mycobacteria, fungus or cytomegalovirus, adverse drug effects, lymphoma or

	Laboratory data of Case 4										
Month/year	Nov/2003	Jan/2006	Jul/2006	Apr/2007	Jan/2008	Mar/2008	Jul/2008				
IgG anti-B19V	NEG	NEG	NEG	POS	POS	POS	POS				
IgM anti-B19V	NEG	NEG	NEG	NEG	NEG	NEG	NEG				
PCR-B19	-	NEG	POS	NEG	-	-	-				
CD4 <sup>+</sup> (cells/mm <sup>3</sup> )	215	364	327	504	539	285	528				
Hb (g/dL)	12.8	12.3	12.5	14.2	-	13.9	13.6				

TABLE IV

B19V: parvovirus B19; Hb: haemoglobin; NEG: negative; PCR: polymerase chain reaction; POS: positive; -: lack of data in the medical records.

TABLE V Laboratory data of Case 5

			-				
Month/year	Jan/2002	Jun/2002	Nov/2002	Jun/2003	Jul/2004	May/2005	Oct/2005
Test 1 IgG anti-B19V	IND	NEG	+/-a	IND	NEG	IND	IND
Test 2 IgG antiB19V	NEG						
Test 3 IgG anti-B19V	NEG						
IgM anti-B19V	NEG						
PCR-B19	-	NEG	POS	NEG	-	-	-
CD4 <sup>+</sup> (cells/mm <sup>3</sup> )	80	61	36	314	253	344	244
Hb (g/dL)	-	5.92	9.14	11.3	12.3	-	13.7

*a*: weakly positive; B19V: parvovirus B19; Hb: haemoglobin; IND: indetermined; NEG: negative; PCR: polymerase chain reaction; POS: positive; -: lack of data in the medical records.

the direct effect of HIV infection itself on the bone marrow (Abkowitz et al. 1997, Koduri 2000). As there are usually no clinical clues of a B19V infection in these patients, the presence of anaemia, especially during B19V epidemics, should alert the physician to the possibility of B19 infection.

One of our patients (Case 3) had viraemia persisting for at least nine months (from September 2005-June 2006). This propensity to develop protracted B19V viraemias in HIV-infected and other immunosuppressed individuals was described long ago (Kurtzman et al. 1989b, Weiland et al. 1989, Frickhofen et al. 1990, Gyllensten et al. 1994). Prolonged viraemias have also been described in healthy blood donors (Hitzler & Runkel 2002).

Dissociation between the serological and molecular markers was observed in Case 5, most likely due to the inability of this immunocompromised patient to produce neutralising antibodies. According to Kurtzman et al. (1989a), the immune response may be quantitatively and qualitatively altered by immunosuppression or during persistent infections. Consequently, in this clinical context, diagnostic genome detection has been advocated by some authors (Calvet et al. 1999).

It is well known that genotype 1 is the most common B19V genotype detected world-wide (Servant-Delmas et al. 2010). In this study, genotype 1 was detected in four of the five HIV-infected patients. This genotype was also detected in EI patients during outbreaks in Niterói (Pereira et al. 2010). Unexpectedly, a genotype 3b strain was detected in a sample from September 2005.

Ferry et al. (2009) found that the detection of B19V genotype 2 or 3 DNA was infrequent in the blood samples of a large cohort of immunocompromised, HIV-infected anaemic patients, despite the use of highly sensitive real-time PCR methods. All three genotypes have been identified in Brazil, although genotype 1 is the most common (Sanabani et al. 2006, Freitas et al. 2008, Keller et al. 2009, Garcia 2010). Genotype 3b was identified in a serum sample obtained from a patient with systemic lupus erythematosus in 1999 in Belém, state of Pará (Freitas et al. 2008); in a sample from a patient from São Paulo, state of São Paulo, with anaemia after a kidney transplantation in 2004 (Keller et al. 2009) and in children with exanthaematic disease (Freitas et al. 2008).

This study had the limitations inherent in retrospective studies involving the review of medical records. One such limitation was the long interval between blood collections. These long intervals made it difficult to precisely know the time of B19V seroconversion and they prevented complete analysis of the interactions between serological (especially IgM) and molecular markers. Another limitation was not performing quantitative PCR (qPCR), as it is known that the virus may persist for a long time and maintain persistent viraemias in HIV-positive patients (Frickhofen et al. 1990, Miao et al. 2012). However, the following evidence of primary infections in our patients was found: (i) the previous serum samples (in relation to the PCR-positive samples) were negative by PCR, (ii) the subsequent samples (in relation to the PCR-positive samples) were also negative by PCR, (iii) seroconversions were observed in four of the five cases and (iv) all but one case occurred during an outbreak of B19V infection in Niterói. If these positive PCRs corresponded to long persistent infections in HIV-positive patients who were recovering their immunity, a decrease in the intensity of the viraemia would be expected by the time of seroconversion, together with a switch from positive to negative PCR results. This outcome did not occur, as the PCR results changed from negative to positive at the time of seroconversion and then changed from positive to negative afterward. Considering the high sensitivity of qPCR and the long intervals between the blood collections, the use of this technique would be expected to detect low DNA levels in the blood samples despite the presence of circulating IgG antibodies. However, this finding would not change the interpretation of the obtained results.

In conclusion, our findings show that B19V infections in HIV-infected patients may be limited, subtle and apparent only in retrospective analyses due to the variety of causes of anaemia in these patients and the adequate use of HAART. The finding of four strains of genotype 1 and one strain of genotype 3 is consistent with their relative frequency in Brazil. There were no apparent relationships between the infecting genotype and the clinical course.

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## REFERENCES

- Abkowitz JL, Brown KE, Wood RW, Kovach NL, Green SW, Young NS 1997. Clinical relevance of parvovirus B19 as a cause of anaemia in patients with human immunodeficiency virus infection. *J Infect Dis 176*: 269-273.
- Calvet A, Pujol MO, Bertocchi M, Bastien O, Boissonnat P, Mornex JF 1999. Parvovirus B19 infection in thoracic organ transplant recipients. J Clin Virol 13: 37-42.
- Chernak E, Dubin G, Henry D, Naides SJ, Hodinka RL, MacGregor RR, Friedman HM 1995. Infection due to parvovirus B19 in patients infected with human immunodeficiency virus. *Clin Infect Dis* 20: 170e3.
- de Azevedo KML, Setúbal S, Camacho LAB, Garcia RCNC, Siqueira MM, Pereira RFA, de Oliveira SA 2012. Parvovirus B19 seroconversion in a cohort of human immunodeficiency virus-infected patients. *Mem Inst Oswaldo Cruz 107*: 356-361.
- de Azevedo KML, Setúbal S, Camacho LAB, Velarde LGC, de Oliveira SA 2009. Seroepidemiological study of human parvovirus B19 among human immunodeficiency virus-infected patients in a middle-sized city in Rio de Janeiro, Brazil. *Mem Inst Oswaldo Cruz 104*: 901-904.
- Durigon EL, Erdman DD, Gary WG, Pallansh MA, Torok TJ, Anderson LJ 1993. Multiple prime pairs for polymerase chain reaction (PCR) amplification of human parvovirus B19 DNA. J Virol Methods 44: 155-165.
- Fauquet CM, Mayo MA, Maniloff J, Desselberger U, Ball LA 2005. Virus taxonomy: VIIIth report of the ICTV, Elsevier/Academic Press, London, 1162 pp.

- Ferry T, Hirschel B, Dang T, Meylan P, Delhumeau C, Rauch A, Weber R, Elzi L, Bernasconi E, Schimid P, Calmy A 2009. Infrequent replication of parvovirus B19 and erythrovirus genotypes 2 and 3 among HIV-infected patients with chronic anemia. *Clin Inf Dis* 50: 115-118.
- Freitas RB, Melo FL, Oliveira DS, Romano CM, Freitas MRC, Macedo O, Linhares AC, Zanotto PM, Durigon EL 2008. Molecular characterization of human erythrovirus strains obtained from patients with several clinical presentations in the Amazon Region of Brazil. J Clin Virol 43: 60-65.
- Frickhofen N, Young NS 1990. Polymerase chain reaction for detection of parvovirus B19 in immunodeficient patients with anaemia. Behring *Inst Mitt 85*: 46-54.
- Frickhofen N, Abkowitz JL, Safford M, Berry JM, Antunez-de-Mayolo J, Astrow A, Cohen R, Halperin I, King L, Mintzer D, Cohen B, Young NS 1990. Persistent B19 parvovirus infection in patients infected with human immunodeficiency virus type 1 (HIV-1): a treatable cause of anemia in AIDS. *Ann Int Med 113*: 926-933.
- Garcia SO 2010. O significado das variantes de eritrovirus em pacientes com citopenias de origem desconhecida, MsC Thesis, Universidade de São Paulo, São Paulo, 169 pp.
- Gyllensten K, Sonnerborg A, Jorup-Ronstrom C, Halvarsson M, Yun Z 1994. Parvovirus B19 infection in HIV-1 infected patients with anemia. *Infection* 22: 356-358.
- Hall TA 1999. BioEdit: biological sequence alignment editor for Windows. Available from: mbio.ncsu. edu/BioEdit/bioedit.html.
- Hitzler WE, Runkel S 2002. Prevalence of human parvovirus B19 in blood donors as determined by a haemagglutination assay and verified by the polymerase chain reaction. *Vox Sang 82*: 18-23.
- Keller LW, Barbosa LM, Melo FL, Pereira LM, Neto ED, Lanhez LE, Durigon EL 2009. Phylogenetic analysis of a near-full-length sequence of an erythrovirus genotype 3 strain isolated in Brazil. *Arch Virol 154*: 1685-1687.
- Koduri 2000. Parvovirus B19-related anaemia in HIV-infected patients. AIDS Patient Care STDS 14: 7-11.
- Kurtzman GJ, Cohen BJ, Field AM, Oseas R, Blaese RM, Young NS 1989a. Immune response to B19 parvovirus and an antibody defect in persistent viral infection. J Clin Invest 84: 1114-1123.
- Kurtzman GJ, Frickhofen N, Kimball J, Jenkins DW, Nienhuis AW, Young NS 1989b. Pure red-cell aplasia of 10 years' duration due to persistent parvovirus B19 infection and its cure with immunoglobulin therapy. N Engl J Med 321: 519-523.
- Miao H, Zhu J, Yin H, Ke L, Gao L, Pan Z, Yang X, Li W 2012. Human immunodeficiency virus/human parvovirus B19 co-infection in blood donors and AIDS patients in Sichuan, China. *Blood Transfus 10*: 502-514.
- Mylonakis E, Dickinson BP, Mileno MD, Flanigan T, Schiffman FJ, Mega A, Rich JD 1999. Persistent parvovirus related anaemia of

seven years duration in a HIV infected patient: complete remission of highly active antiretroviral therapy. *Am J Hem 60*: 164-166.

- Nguyen QT, Sifer C, Schneider V, Allaume X, Servant A, Bernaudin F, Auguste V, Garbarg-Chenon A 1999. Novel human erythrovirus associated with transient aplastic anaemia. J Clin Microbiol 37: 2483-2487.
- Nguyen QT, Wong S, Heegaard ED, Brown KE 2002. Identification and a characterization of a second novel human erythrovirus variant A6. *Virology 301*: 374-380.
- Oliveira SA, Marinho TAS, Silva ASV, Rodrigues NB, Faria RA, Setúbal S 2005. Estudo da frequência da infecção pelo parvovírus B19 em casos de doenças exantemáticas no período de 1994 a 2004. Braz J Infect Dis 9 (Suppl.): 73.
- Otto T, Vasconcellos E, Gomes L, Moreira A, Degrave W, Mendonça-Lima L and Alves-Ferreira M 2008. Chormapipe: a pipeline for analysis, quality control and management for a DNA sequencing facility. *Gen Mol Res 7*: 861-871.
- Pereira RFA, Cubel Garcia RCN, Azevedo KLM, Setúbal S, Siqueira MAMT, Castro TX, Oliveira SA 2012. Molecular characterization of human parvovirus B19 strains circulating in Niterói/RJ - Brazil. Virus Reviews & Research 17 (Suppl.): 158-159.
- Sanabani S, Kleine Neto W, Pereira J, Sabino EC 2006. Sequence variability of human erythroviruses present in bone marrow of Brazilian patients with various B19-related hematological symptoms. J Clin Microbiol 44: 604-606.
- Servant A, Laperche S, Lallemand F, Marinho V, de SaintMaur G, Meritet JF, Garbarg-Chenon A 2002. Genetic diversity within human erythroviruses identification of three genotypes. *J Virol* 76: 9124-9134.
- Servant-Delmas A, Lefrere JJ, Morinet F, Pillet S 2010. Advances in human B19 virus biology. *J Virol 84*: 9658-9665.
- van Elsacker-Niele AMW, Kroon FP, van der Ende ME, Salimans MM, Spaan WJ, Kroes AC 1996. Prevalence of parvovirus B19 infection in patients infected with human immunodeficiency virus. *Clin Infect Dis* 23: 1255-1260.
- Ware AJ, Moore T 2001. Resolution of chronic parvovirus B19-induced anaemia by use of highly active antiretroviral therapy in a patient with acquired immunodeficiency syndrome. *Clin Infect Dis* 32: 122-123.
- Weiland HT, Salimans MM, Fibbe WE, Kluin PM, Cohen BJ 1989. Prolonged parvovirus B19 infection with severe anaemia in a bone marrow transplant patient. Br J Haematol 71: 300.
- WHO World Health Organization 2001. Iron deficiency anaemia. Assessment, prevention and control. A guide for programme managers. Available from: who.int/nutrition/publications/en/ ida assessment prevention control.pdf.
- Woolf AD, Campion GV, Chishick A, Wise S, Cohen BJ, Klouda PT, Caul O, Dieppe PA 1989. Clinical manifestations of human parvovirus B19 in adults. *Arch Intern Med 149*: 1153-1156.