In this study, AHF lot B was derived solely from plasma tested by a high-titer B19 NAT screening procedure and had no detectable B19 DNA. In contrast, AHF lot A was mostly derived from unscreened plasma. The transmission case might not have occurred had B19 NAT screening been performed. That is, if donations with high levels of B19 DNA had been identified, the high-titer plasma pool for the implicated lot, 10⁷ geq per mL, would not have existed. A B19 transmission by a similar S/D-treated, immunoaffinity-purified, AHF product to a seronegative child with mild hemophilia A, who had not been previously infused with any blood product, has been documented. As in most reported cases, however, sequencing analysis was not performed and the amount of B19 DNA infused was unknown.

Little is known regarding the correlation between a product's infectivity and its B19 DNA content. The B19 infectious dose in susceptible individuals, that is, presumably seronegative persons, would be expected to vary depending on whether the product contained anti-B19 igG antibodies. For example, pooled plasma, S/D-treated, had levels of anti-B19 IgG^{11,29} approximately 40 IU per mL in every product lot because each pool of plasma represented up to 2500 plasma donations. Only those seronegative volunteers infused with a 200-mL dose of product lots containing greater than 10⁷ geq per mL B19 DNA were infected, whereas those infused with an equal volume of lots containing less than 10⁴ geq per mL did not seroconvert.^{29,30,32}

In a separate transmission case, a seronegative child was infected by infusing a dry heat–treated FVIII concentrate, which contained 4×10^3 geq per mL B19 DNA, over a period of 52 days. ¹⁷ The total infectious dose for this case was equivalent to 4×10^6 geq of B19 DNA from a product whose anti-B19 content, if any, was unknown. In our study, the seronegative recipient was infected by receiving a total of 2×10^4 geq of B19 DNA from a product that contained no detectable B19 IgG.

In conclusion, we have confirmed B19 transmission in a recipient of a S/D-treated high-purity AHF product derived from mostly B19 NAT unscreened plasma. The seronegative recipient became infected after receiving 2×10^4 geq (or IU) of B19 DNA present in the product.

Therefore, to safeguard the viral safety with respect to B19, minipool screening by B19 NAT should be implemented to reduce the level of potentially infectious B19 virus in the resulting products, especially those without the presence of anti-B19.

ACKNOWLEDGMENTS

We thank J.S. Finlayson, PhD, for critical review of the manuscript and Donald Baker, PhD, of Baxter BioScience for providing not only the information regarding NAT screening status relevant to implicated AHF lots but also relevant plasma pools for analysis. In addition, we appreciate the staff of the Dartmouth Hitchcock Hemophilia and Thrombosis Center for assistance with collecting data for this project and Mary Hitchcock Memorial Hospital for identification of the specific implicated lots of AHP.

REFERENCES

- Young NS, Brown KE. Mechanisms of disease: parvovirus B19. N Engl J Med 2004;350:586-97.
- Anderson MJ, Tsou C, Parker RA, et al. Detection of antibodies and antigen of human parvovirus B19 by enzyme-linked immunosorbent assay. J Clin Microbiol 1986;24:522-6.
- Cohen BJ, Buckley MM. The prevalence of antibody to human parvovirus B19 in England and Wales. J Med Microbiol 1988;25:151-3.
- Jordan J, Tiangco B, Kiss J, Koch W. Human parvovirus B19: prevalence of viral DNA in volunteer blood donors and clinical outcomes of transfusion recipients. Vox Sang 1998;75:97-102.
- Aubin JT, Defer C, Vidaud M, Maniez MM, Flan B. Largescale screening for human parvovirus B19 DNA by PCR. application to the quality control of plasma for fractionation. Vox Sang 2000;78:7-12.
- Weimer T, Streichert S, Watson C, Gröner A. High-titer screening PCR: a successful strategy for reducing the parvovirus B19 load in plasma pools for fractionation. Transfusion 2001;41:1500-4.
- Siegl G, Cassinotti P. Presence and significance of parvovirus B19 in blood and blood products. Biologicals 1998;26:89-94.
- Luban NLC. Human parvoviruses: implications for transfusion medicine. Transfusion 1994;34:821-7.
- Saldanha J, Minor P. Detection of human parvovirus B19
 DNA in plasma pools and blood products derived from these
 pools: implications for efficiency and consistency of removal
 of B19 DNA during manufacture, Br J Haematol 1996;93:714
- Eis-Hübinger AM, Sasowski U, Brackmann HH. Parvovirus B19 DNA contamination in coagulation factor VIII products. Thromb Haemost 1999;81:476-7.
- Schmidt I, Blürnel J, Seitz H, Willkommen H, Löwer J. Parvovirus B19 DNA in plasma pools and plasma derivatives. Vox Sang, 2001;81:228-35.
- Azzi A, Morfini M, Mannucci PM. The transfusionassociated transmission of parvovirus B19. Transfus Med Rev 1999;13:194-204.
- Azzi A, Ciappi S, Zakvrzewska K, et al. Human parvovirus B19 infection in hemophiliacs first infused with two highpurity, virally attenuated factor VIII concentrates. Am J Hematol 1992;39:228-30.
- Yee TT, Cohen BJ, Pasi KJ, Lee CA. Transmission of symptomatic parvovirus B19 infection by clotting factor concentrate. Br J Haematol 1996;93:457-9.
- Santagostino E, Mannucci PM, Gringeri A, et al.
 Transmission of parvovirus B19 by coagulation factor

- concentrates exposed to 100°C heat after lyophilization. Transfusion 1997;37:517-22.
- 16. Matsui H, Sugimoto M, Tsuji S, et al. Transient hypoplastic anemia caused by primary human parvovirus B19 infection in a previously untreated patient with hemophilia transfused with a plasma-derived, monoclonal antibodypurified factor VIII concentrate [case report]. J Pediatr Hematol Oncol 1999;21:74-6.
- Blümel J, Schmidt J, Effenberger W, et al. Parvovirus B19 transmission by heat-treated clotting factor concentrates. Transfusion 2002;42:1473-81.
- Schosser R, Keller-Stanislawski B, Nübling M, Löwer J. Causality assessment of suspected virus transmission by human plasma products. Transfusion 2001;41:1020-9.
- Saldanha J, Lelie N, Yu MW, Heath A: B19 Collaborative Study Group. Establishment of the first World Health Organization International Standard for human parvovirus B19 DNA nucleic acid amplification techniques. Vox Sang 2002:82:24-31.
- Blood safety monitoring among persons with bleeding disorders—United States, May 1998-June 2002. MMWR Morb Mortal Wkly Rep 2003;51:1152-4.
- Shade RO, Blundell MC, Cotmore SF, Tattersall P, Astell CR. Nucleotide sequence and genome organization of human parvovirus B19 isolated from the serum of a child during aplastic crisis. J Virol 1986;58:921-36.
- Blundell MC, Beard C, Astell CR. In vitro identification of a B19 parvovirus promoter. Virology 1987;157:534-8.
- Bhattacharyya SP, Tan D, Guo ZP, e tal. Presence of human parvovirus B19 DNA in factor VIII concentrates: effects of viral clearance and product purification procedures. Haemophilia 2000;6:353.
- Blümel J, Schmidt I, Willkommen H, Löwer J. Inactivation of parvovirus B19 during pasteurization of human serum albumin. Transfusion 2002;42:1011-8.
- 25. Yunoki M, Tsujikawa M, Urayama T, et al. Heat sensitivity of human parvovirus B19. Vox Sang 2003;84:164-9.

- Burnouf T, Radosevich M. Nanofiltration of plasma-derived biopharmaceutical products. Haemophilia 2003;9:24-37.
- Servant A, Laperche S, Lallemand F, et al. Genetic diversity within human erythroviruses: identification of three genotypes. J Virol 2002;76:9124-34.
- 28. Dorsch S, Kaufmann B, Schaible U, et al. The VP1-unique region of parvovirus B19: amino acid variability and antigenic stability. J Gen Virol 2001;82:191-9.
- Blood Products Advisory Committee. Nucleic acid testing of blood donors for human parvovirus B19 [Internet]. Rockville (MD): U.S. Food and Drug Administration; 1999, Available from: http://www.fda.gov/ohrms/dockets/ac/99/ transcript/3548tl.rtf
- Davenport R, Geohas G, Cohen S, et al. Phase IV study of PLAS+ SD: hepatitis A (HAV) and parvovirus B19 safety results. Blood 2000;96:451a.
- Koenigbauer U, Eastlund T, Day JW. Clinical illness due to parvovirus B19 infection after infusion of solvent/detergenttreated pooled plasma. Transfusion 2000;40:1203-6.
- Brown KE, Young NS, Alving BM, LH.Barbosa. Parvovirus B19: implications for transfusion medicine, summary of a workshop. Transfusion 2001;41:130-5.
- Tabor E, Yu MW, Hewlett I, Epstein JS. Summary of a workshop on the implementation of NAT to screen donors of blood and plasma for viruses. Transfusion 2000;40: 1273-5.
- Hewlett IK, Yu MW, Epstein JS. Implementation of donor screening for infectious agents transmitted by blood by nucleic acid technology. The International Forum. Vox Sang 2002;82:87-111.
- Tabor E, Epstein JS. NAT screening of blood and plasma donations: evolution of technology and regulatory policy. Transfusion 2002;42:1230-7.
- 36. Aberham C, Pendl C, Gross P, Zerlauth G, Gessner M. A quantitative, internally controlled real-time PCR assay for the detection of parvovirus B19 DNA. J Virol Methods 2001;92:183-91.

医薬品 研究報告 調査報告書

戦別番号・報告回数	第1報		1	第一報入手日 新医薬品等 2005年3月10日 該当なし		の区分	厚生労働省処理欄
一般的名称乾燥濃縮人アンチトロンビンIII販売名(企業名)アンスロビン P-ベーリング (ZLB ベーリング株式会社)		·Ш		International J of Hem			
		研究報告の公表状況	80(2004) 301-305 Current safety of the blood supply in the United States		公表国 米国		
NAT などの名られなどの名られないでは、 いべかりりる。ア間ガスがマシーでT. cruzi抗患がでいる。 WNVのNATA	ている血液製剤の安全対策に関するロイルスのテストを実施した。新興感染症であるシャーガス、ダニによる感染である。米国で年間 1-2 例の輸血感染があったためである。 は、ラテンアメリカでは一般的体のテストはない。献血時にシャでT. cruziが感染した例はない。 算入により、約1000件のWNV-R り場合、検出できず感染する恐れ	た結果、HIV と HC ス病、バベシア症、 国で輸血によるバベ とがない。輸血感染 がある。献血時の問 りな寄生虫(T. cruzi) ャーガス病である。 NA陽性の献血ドナ	で 感染は献血 1500 万代マラリア、WNV と vC マラリア、WNV と vC でかけない かっぱい かっぱい かっぱい かっぱい かっぱい 大国で4代か 問診されるだけである マーを見つけ、感染の伝	ついて1件に減少して1Dに注意しなければな けの報告がある。日本はないが、高齢者、免疫低いドナーは排除している。カナダで2件の確定しる。臨床研究によると、	いる。 らない。 (1 件の報告が K下患者、無脾 いるが、最近の いた感染報告が 輸血を受けた。	ある。献血時の 症患者は危険性 感染例はこの問 ある。現在米国 総計120,000人の	
報告企業の意見		今後の対応			· 		
シャーガス病は、T. あり、本剤を含む弊 過(0.22 ミクロン等) 染報告は存在するが 53℃、30 分の加熱で マラリアも同様に、 ある。 シャーガス病、マラ	血液供給を脅かす情報を入手し cruziによる感染症で、大きされ 社の血漿分画製剤における製造 で十分除去できるものである。 が、血漿分画製剤からの感染の で死滅する報告がある。 血液製剤の製造過程で寄生虫は リア、バベシア症に関して、弊 安全である。	が約 20 ミクロンで 生工程、特に滅菌ろ また輸血からの感 報告はない。また は除去される報告が		興感染症に関する情報	収集に努める所	所存である。	





International Journal of HEMATOLOGY

Current Safety of the Blood Supply in the United States

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Received August 16, 2004; received in revised form May 24, 2004; accepted September 3, 2004

Abstract

In common with other developed countries, the United States has placed a great deal of emphasis on blood safety. As a result of careful donor selection and the use of advanced tests, including nucleic acid testing (NAT), the risk of transmission of human immunodeficiency virus and hepatitis C virus has been reduced to about 1 in 1.5 million donations. NAT for hepatitis B virus has not been introduced, but nevertheless the risk is low. Attention recently has been focused on emerging infections. NAT for West Nile virus was implemented within 6 to 8 months of recognition of the need to prevent transfusion transmission of this newly introduced virus. Approximately 1000 potentially infectious donations were identified and removed from the blood supply during the 2003 season. Other emerging infections attracting attention include Chagas' disease, babesiosis, malaria, and variant Creutzfeldt-Jakob disease.

Int J Hematol. 2004;80:301-305. doi: 10.1532/IJH97.04123 ©2004 The Japanese Society of Hematology

Key words: Blood transfusion; HIV; Hepatitis C; Hepatitis B; Emerging infections

1. Introduction

In the United States, blood safety depends on selection of voluntary donors, extensive use of screening questions, laboratory testing, and maintenance of deferral registries. These processes are highly regulated and are managed under voluntary quality systems such as the standards of the American Association of Blood Banks (AABB). Over the years, there has been a process of continuing improvement, particularly in testing. This process has resulted in a very low frequency of residual infectivity from the blood supply, at least for hepatitis and retroviral infections. The recent introduction of nucleic acid testing (NAT) has had a major impact on safety [1-3]. At the same time, a number of new threats to blood safety have appeared and necessitated additional donor deferral and/or testing measures [4]. Notable among these new infections have been West Nile virus (WNV) and variant Creutzfeldt-Jakob disease (vCJD).

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2. Current Risk of Hepatitis Viruses and Retroviruses

The original approach to controlling transfusion-transmitted hepatitis and, later, acquired immunodeficiency syndrome (AIDS) involved careful questioning of donors about their medical history and risk behaviors. The majority of these questions are still in place, despite the use of tests of increasing sensitivity. Overall, however, very few donors are deferred as a result of these questions, but there is good evidence that almost 2% of donors may fail to report deferrable risk behaviors during the donation process [5]. Nevertheless, both the prevalence and incidence of human immunodeficiency virus (HIV), hepatitis B virus (HBV), and hepatitis C virus (HCV) infection among donors is much lower than that attributable to the general population (Table 1) [6]. It seems likely that the majority of infected and at-risk individuals do not donate as a result of the use of a voluntary donor population along with broad public education.

In the United States, the following tests are performed on all blood donations: antibodies to hepatitis B core antigen (anti-HBc), hepatitis B surface antigen (HBsAg), antibodies to hepatitis C virus (anti-HCV), antibodies to HIV-1 and HIV-2 (anti-HIV-1/2), antibodies to human T-lymphotrophic virus I (HTLV-I) and HTLV-II (anti-HTLV-I/II), serologic test for syphilis, and minipool NAT for HIV and HCV RNA. In addition, all donations are tested by investigational NAT

next 2 years, a similar number of cases were seen, but surveillance studies revealed that the virus was spreading to a larger area each year. The major amplifying hosts were a number of bird species, and significant avian mortality occurred. At that stage in the epidemic, there was little concern about the risk of transfusion transmission of WNV, although a risk estimate was published for the initial outbreak [12]. In 2002, however, there was an enormous outbreak of human cases, totaling 4156 with 284 deaths and affecting the majority of the continental United States. Of most concern, 61 potential cases of transfusion-transmitted WNV infections were reported, and of these 23 were confirmed [13]. In all cases in which samples of the implicated donations were available, it was found that readily detectable levels of WNV RNA were present.

These observations led to rapid development and implementation of NAT for WNV. The decision to move toward such testing was made in September 2002, and tests were fully in place before the start of the 2003 WNV season in July. During 2003 there were 9388 human cases of WNV disease with 246 deaths. The cases occurred over an even broader geographical area than that affected in 2002. Blood collectors identified approximately 1000 WNV RNA-positive donations (a rate of about 1 in 5000), preventing many potential infections among blood recipients. In some localities, the frequency of finding RNA-positive donors was extremely high (for example, 1 in 47 in parts of Nebraska) [14]. As a result of concerns that NAT in minipools did not detect all RNApositive donations and the eventual demonstration of recipient infections attributable to such low-titer samples, limited single-donation testing was implemented in some areas of highest incidence of WNV infection. It was clear that this approach did detect some otherwise undetectable, potentially infectious samples. The practice of performing resource-limited single-donation testing of this type continued into the WNV season in 2004. During 2003, 6 confirmed cases of transfusion-transmitted WNV were reported. All seemed to be attributable to donations with very low titers of WNV, below the levels detectable by minipool testing. As of this writing, however, there has been no authenticated case in which transmission has been attributable to a blood unit with detectable levels of WNV immunoglobulin M (IgM), although it is known that IgM and WNV RNA may coexist for a time. Thus the unexpected emergence of WNV and the finding of its transmissibility by transfusion posed a significant challenge in 2002 [15]. Rapid development and implementation of NAT clearly had a significant impact on the problem, although it has not proven possible to entirely eliminate the risk.

4. Other Infections

4.1. Syphilis

All donations are tested for syphilis with treponemal tests, nontreponemal tests, or both. There has been no reported case of transfusion syphilis in the United States for well over 40 years. It is possible that this outcome is a result of continuing testing, and it has not proven possible to eliminate the requirement for such testing [16]. In recent studies, however,

Orton and her colleagues did not find treponemal DNA and/or RNA in 169 blood donor samples with confirmed positive serological test results for syphilis [17]. Thus the potential for detection of an infectious sample appears to be low.

4.2. Malaria

Malaria is probably the infection most frequently transmitted by transfusion. However, such transmission is a rarity in the United States with only 1 or 2 cases annually [18]. Approximately 1000 cases of imported malaria are diagnosed each year in the United States. This number is small compared with the numbers in, for example, Western Europe. There is a comprehensive effort to exclude at-risk donors by careful questioning about their travel history. Many of the recent cases of transfusion transmission of malaria appear to be attributable to failures in the questioning process. Although endemic malaria has been eliminated from the United States, there is concern about the occurrence of epidemiologically unexplainable cases, most recently in Virginia and Florida. At least some of such cases are attributable to mosquito-borne transmission from migrant workers or travelers, but it is clear that a questioning strategy would be ineffective in identifying such secondary cases if the individuals were to present to give blood. This is a situation that deserves future scrutiny.

4.3. Chagas' Disease

It is well-established that Chagas' disease (caused by the protozoan parasite Trypanosoma cruzi) is transmitted by blood transfusion. In Latin America, where human infection is endemic, it is estimated that a recipient of parasitemic blood has a 12% to 50% chance of being infected. Because infection is often lifelong, population movements from endemic areas lead to the presence of infected and potentially infectious individuals in nonendemic areas such as the United States. There have been a total of 6 wellauthenticated transfusion transmitted cases of T cruzi infection in the United States (4 cases) and Canada (2 cases) [4]. These cases are thought to be a substantial minority of the cases that might occur, because the disease is not readily diagnosed, nor is it often suspected. One of the recognized cases was identified only as a result of careful follow-up of a patient inadvertently given a transfusion of seropositive platelets [19]. Essentially all cases were traced to donors who had been infected early in life in areas of endemicity. There is currently no testing for T cruzi antibodies in the United States, and donors are asked only if they have had Chagas' disease. This measure is very insensitive [20]. Seroprevalence studies have shown that in areas with a high proportion of migrants from Latin America, as many as approximately 1 in 7500 donors may be in the seropositive state, and approximately 60% of these donors actually have parasitemia, as demonstrated by polymerase chain reaction analysis and or parasite culture [4]. It is thought that the national seroprevalence rate may be between 1 in 40,000 and 1 in 25,000, suggesting a potential for a few hundred infections each year. Lookback studies, however, did not identify any infected recipients within a group of 19 patients who received blood been shown that a rapid response to a newly emerging, transfusion-transmissible agent is possible, as in the case of WNV.

References

- Dodd RY, Notari EP, Stramer SL. Current prevalence and incidence of infectious disease markers and estimated window-period risk in the American Red Cross blood donor population. Transfusion. 2002;42:975-979.
- Stramer SL, Glynn SA, Kleinman SH, et al. Detection of HIV-1 and HCV infections among antibody-negative blood donors by nucleic acid-amplification testing. N Engl J Med. 2004;351:760-768.
- Goodman JL. The safety and availability of blood and tissues: progress and challenges. N Engl J Med. 2004;351:819-822.
- Dodd RY, Leiby DA. Emerging infectious threats to the blood supply. Annu Rev Med. 2004;55:191-207.
- Williams AE, Thomson RA, Schreiber GB, et al. Estimates of infectious disease risk factors in US blood donors. JAMA. 1997;277: 967-972.
- Zou S, Dodd RY, Stramer SL, Strong DM. Probability of viremia with HBV, HCV, HIV, and HTLV among tissue donors in the United States. N Engl J Med. 2004;351:751-759.
- Delwart EL, Kalmin ND, Jones TS, et al. First report of human immunodeficiency virus transmission via an RNA-screened blood donation. Vox Sang. 2004;86:171-177.
- Phelps R, Robbins K, Liberti T, et al. Window-period human immunodeficiency virus transmission to two recipients by an adolescent blood donor. *Transfusion*. 2004;44:929-933.
- Schüttler CG, Caspari G, Jursch CA, et al. Hepatitis C virus transmission by a blood donation negative in nucleic acid amplification tests for viral RNA. Lancet. 2000;355:41-42.
- Busch MP. Closing the windows on viral transmission by blood transfusion. In: Stramer SL, ed. Blood Safety in the New Millenium. Bethesda, MD: American Association of Blood Banks; 2001:33-54.
- Janssen RS, Satten GA, Stramer SL, et al. New testing strategy to detect early HIV-1 infection for use in incidence estimates and for clinical and prevention purposes. JAMA. 1998;280:42-48.
- Biggerstaff BJ, Petersen LR. Estimated risk of West Nile virus transmission through blood transfusion during an epidemic in Queens, New York City. Transfusion. 2002;42:1019-1026.
- Pealer LN, Marfin AA, Petersen LR, et al. Transmission of West Nile virus through blood transfusion in the United States in 2002. N Engl J Med. 2003;349:1236-1245.

- Dodd RY. Emerging infections, transfusion safety, and epidemiology. N Engl J Med. 2003;349:1205-1206.
- Biggerstaff BJ, Petersen LR. Estimated risk of transmission of the West Nile virus through blood transfusion in the US, 2002. Transfusion. 2003;43:1007-1017.
- Orton S. Syphilis and blood donors: what we know, what we do not know, and what we need to know. Transfus Med Rev. 2001;15: 282-291.
- Orton SL, Liu H, Dodd RY, et al. Prevalence of circulating *Tre-ponema pallidum* DNA and RNA in blood donors with confirmed-positive syphilis tests. *Transfusion*. 2002;42:94-99.
- Mungai M, Tegtmeier G, Chamberland M, et al. Transfusiontransmitted malaria in the United States from 1963 through 1999. N Engl J Med. 2001;344:1973-1978.
- Leiby DA, Lenes BA, Tibbals MA, et al. Prospective evaluation of a patient with *Trypanosoma cruzi* infection transmitted by transfusion. N Engl J Med. 1999;341:1237-1239.
- Leiby DA, Read EJ, Lenes BA, et al. Seroepidemiology of Trypanosoma cruzi, etiologic agent of Chagas' disease, in US blood donors. J Infect Dis. 1997;176:1047-1052.
- Leiby DA, Rentas FJ, Nelson KE, et al. Evidence of Trypanosoma cruzi infection (Chagas' disease) among patients undergoing cardiac surgery. Circulation. 2000;102:2978-2982.
- Matsui T, Inoue R, Kajimoto K, et al. First documentation of transfusion-associated babesiosis in Japan [in Japanese]. Rinsho Ketsueki. 2000:41:628-634.
- Prusiner SB. Prion diseases and the BSE crisis. Science. 1997;278: 245-251.
- Collinge J. Prion diseases of humans and animals: their causes and molecular basis. Annu Rev Neurosci. 2001;24:519-550.
- Collinge J. Variant Creutzfeldt-Jakob disease. Lancet. 1999;354: 317-323.
- Peden AH, Head MW, Ritchie DL, et al. Preclinical vCJD after blood transfusion in a PRNP codon 129 heterozygous patient. Lancet. 2004;364:527-529.
- Llewelyn CA, Hewitt PE, Knight RS, et al. Possible transmission of variant Creutzfeldt-Jakob disease by blood transfusion. *Lancet*. 2004;363:417-421.
- Kuehnert MJ, Roth VR, Haley NR, et al. Transfusion-transmitted bacterial infection in the United States, 1998 through 2000. Transfusion. 2001;41:1493-1499.
- Ness P, Braine H, King K, et al. Single-donor platelets reduce the risk of septic platelet transfusion reactions. *Transfusion*. 2001;41: 857-861.