

医薬品
医薬部外品 研究報告 調査報告書
化粧品

識別番号・報告回数		報告日		第一報入手日 2008年8月1日	新医薬品等の区分 該当なし	厚生労働省処理欄
一般的名称	①乾燥抗 HBs 人免疫グロブリン ②ポリエチレングリコール処理抗 HBs 人免疫グロブリン	研究報告の 公表状況	TRANSFUSION 2008; 48 (7): 1333-1341	公表国 フランス		
販売名 (企業名)	①ヘブスプリン (ベネシス) ②静注用ヘブスプリン-IH (ベネシス)					
研究報告の概要	<p><背景> 2005年から2007年の間、チクングニヤウイルス (CHIKV) が、2006年2月に症例数の最大ピークとするレユニオン島での大流行を引き起こした。レユニオン島での供血は、2006年1月に中断された。</p> <p><研究デザインおよび方法> レユニオン島でウイルス血症の供血がされる平均リスクの推定を異なる流行期について計算した。計算には、定点観測の動向調査 (sentinel surveillance)、ウイルス血症の期間、および無症候感染の頻度から割り出した CHIKV 予想発現値を用いた。最後のこれら2つのパラメーターのデータは、最初は仮定に基づき、次いでアウトブレイクの期間に実施された検討をもとに出した。この予想リスクを、血小板ドネーションのスクリーニングのために実施した CHIKV 核酸増幅検査の結果と比較した。</p> <p><結果> アウトブレイクの期間中、リスクの平均値は、ドネーション 100,000 当たり 132 と予想された。このリスクは、2006年2月のアウトブレイクの最大期にピークに達し、ドネーション 100,000 当たり 1,500 であった。もし採血が中断されていなかったら、全体で 47 の供血がウイルス血症であったであろう。この期間、757,000 人住民のうちの 312,500 人が蚊を媒介にして感染していたと予想される。2006年1月から5月まで、予想リスク平均値 (0.7%) と血小板供血で観察されたリスク (0.4%) は同じ大きさであった。</p> <p><結論> この大きなアウトブレイクの間、ウイルス血症の供血の予想リスクは高かったが、蚊媒介の CHIKV 感染のリスクに比べ低かった。この予想リスクは、観察されたリスクの結果と一致したことによって裏付けられた。</p>					<p>使用上の注意記載状況・ その他参考事項等</p> <p>代表として静注用ヘブスプリン-IH の記載を示す。</p> <p>2. 重要な基本的注意 (1) 本剤の原材料となる血液については、HBs 抗原、抗HCV抗体、抗HIV-1抗体、抗HIV-2抗体陰性で、かつALT (GPT) 値でスクリーニングを実施している。更に、プールした試験血漿については、HIV-1、HBV及びHCVについて核酸増幅検査 (NAT) を実施し、適合した血漿を本剤の製造に使用しているが、当該NATの検出限界以下のウイルスが混入している可能性が常に存在する。本剤は、以上の検査に適合した高力価の抗HBs抗体を含有する血漿を原料として、Cohnの低温エタノール分画で得た画分からポリエチレングリコール4000処理、DEAEセファデックス処理等により抗HBs人免疫グロブリンを濃縮・精製した製剤であり、ウイルス不活化・除去を目的として、製造工程において60℃、10時間の液状加熱処理及び過膜処理 (ナノフィルトレーション) を施しているが、投与に際しては、次の点に十分注意すること。</p>
	報告企業の意見				今後の対応	
	レユニオン島におけるチクングニヤウイルス (CHIKV) の流行時の CHIKV 血症献血リスクに関する報告である。血漿分画製剤からのチクングニヤウイルス伝播の事例は報告されていない。また、万一原料血漿にチクングニヤウイルスが混入したとしても、BVDをモデルウイルスとしたウイルスバリデーション試験成績から、本剤の製造工程において十分に不活化・除去されると考えている。				本報告は本剤の安全性に影響を与えないと考えるので、特段の措置はとらない。	

147



TRANSFUSION COMPLICATIONS

Estimated risk of Chikungunya viremic blood donation during an epidemic on Reunion Island in the Indian Ocean, 2005 to 2007

Cécile Brouard, Pascale Bernillon, Isabelle Quatresous, Josiane Pillonel, Azzedine Assal, Henriette De Valk, and Jean-Claude Desenclos for the workgroup "Quantitative Estimation of the Risk of Blood Donation Contamination by Infectious Agents"

BACKGROUND: Between 2005 and 2007, Chikungunya virus (CHIKV) caused a massive epidemic on Reunion Island with a major peak in the number of cases in February 2006. Blood donation was interrupted on the island in January 2006.

STUDY DESIGN AND METHODS: Estimates of the mean risk of viremic blood donation on Reunion Island were computed for different phases of the epidemic. Calculations used CHIKV incidence estimates derived from sentinel surveillance, duration of viremia, and frequency of asymptomatic infection. Data on these two last parameters were initially based on hypotheses and subsequently obtained from studies carried out during the outbreak. The estimated risk was compared to the results of CHIKV nucleic acid testing (NAT) implemented for platelet (PLT) donations screening.

RESULTS: Over the course of the outbreak, the mean risk was estimated at 132 per 100,000 donations. The risk peaked at 1500 per 100,000 donations at the height of the outbreak in February 2006. In total, 47 blood donations would have been potentially viremic if blood collection had not been interrupted. During this period, an estimated 312,500 of 757,000 inhabitants had been infected by mosquito-borne transmission. From January to May 2006, the estimated mean risk (0.7%) and observed risk on PLT donations (0.4%) were of the same order of magnitude.

CONCLUSION: During this large outbreak, the estimated risk of viremic blood donation was high, but low compared to the risk of mosquito-borne CHIKV transmission. The estimated risk was corroborated by the concordant results with the observed risk.

Chikungunya virus (CHIKV) is an alphavirus that belongs to the *Togoviridae* family, transmitted by *Aedes* mosquitoes. It was first identified in 1952 during an outbreak in Tanzania.^{1,2} Afterward, it caused many outbreaks in Africa³⁻⁷ and in Asia.^{3,8-11} In Africa, a sylvatic transmission cycle between wild primates and mosquitoes is thought to maintain the virus, whereas in Asia, it is transmitted from human to human through an urban transmission cycle.³ CHIKV infection is mainly characterized by sudden onset of fever, arthralgia, myalgia, headache, and edemas.^{1,3,8,12,13} Other symptoms like rash, epistaxis, gingivorrhagia, nausea, vomiting, flushed face, or photophobia have also been described. The most typical clinical sign is polyarthralgia that is generally very painful, as suggested by its name Chikungunya meaning in the language of the Tanzanian Makonde plateau "that which bends up" in reference to the stooping posture adopted by patients because of the severity of the joint pains. The symptoms usually resolve within a few days, but in some severe cases, arthralgia may persist for months or years.^{3,13} Serosurveys implemented during prior outbreaks have demonstrated that Chikungunya infection can also be asymptomatic.⁹

In early 2005, CHIKV emerged for the first time in the southwest Indian Ocean region (Comoros, Reunion,

ABBREVIATIONS: CHIKV = Chikungunya virus; WNV = West Nile virus.

From the Institut de Veille Sanitaire (InVS) (French Institute of Public Health Surveillance), Saint-Maurice, France; Etablissement Français du Sang (EFS) (French Blood Services), Tours, France.

Address reprint requests to: Cécile Brouard, Institut de Veille Sanitaire, Département des Maladies Infectieuses, 12 Rue du Val d'Osne, 94415 Saint-Maurice, Cedex, France; e-mail: c.brouard@invs.sante.fr.

Received for publication October 1, 2007; revision received November 13, 2007, and accepted November 13, 2007.

doi: 10.1111/j.1537-2995.2008.01646.x

TRANSFUSION 2008;48:1333-1341.

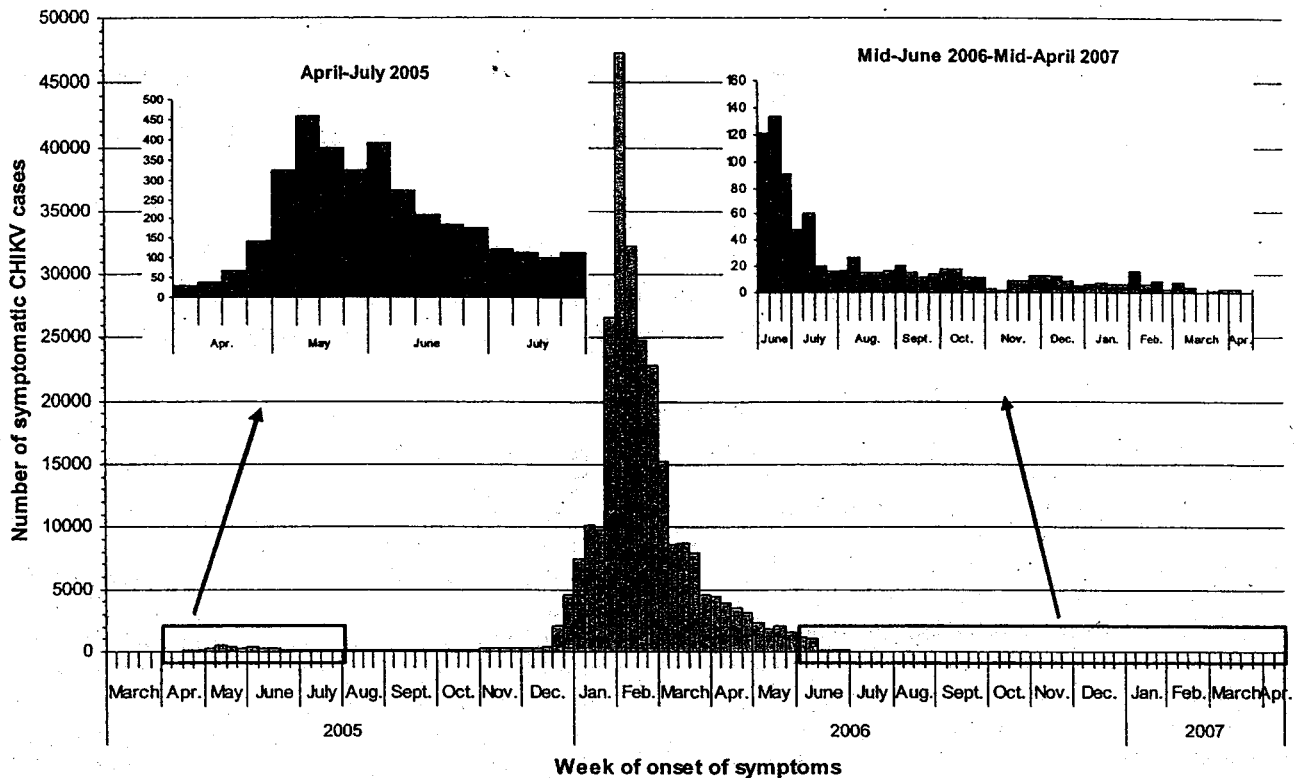


Fig. 1. Distribution of symptomatic cases of CHIKV infection per week of onset of symptoms, Reunion Island, March 28, 2005, through April 15, 2007.

Mayotte, Seychelles, Mauritius, and Madagascar Islands). On Reunion Island, the first cases were identified at the end of April 2005. After a first epidemic peak in May through June 2005 with a maximum of 450 cases during the second week of May, the number of cases decreased during the southern hemisphere winter season. At mid-December, an exponential increase in cases occurred, with almost 10,000 estimated cases at mid-January 2006 (Fig. 1). Because of concerns about the possible transmission of CHIKV by blood transfusion, the French Blood Services (EFS) interrupted blood donations on the island from January 23, 2006, except donations for platelets (PLTs) for which systematic screening for CHIKV genome by nucleic acid amplification testing (NAT) was set up.

At that moment, we estimated the risk of CHIKV viremic blood donation. Afterward, we updated these estimates since more accurate data were available on the incidence of infection and on the frequency of asymptomatic infections. We compared the estimated risk of viremic blood donation to the observed proportion of viremic PLT donations determined by CHIKV NAT screening.

MATERIALS AND METHODS

The estimates were performed by the French Institute of Public Health Surveillance (InVS) in the setting of a work-

group including the French Agency for the Safety of Health Products (Afssaps), the French Blood Services (EFS), and the National Institute for Blood Transfusion (INTS). In early 2005, this group initiated a project with the aim of obtaining a priori quantitative risk estimates of contamination of blood donations by infectious agents for various scenarios in terms of incidence and time-space distribution.¹⁴

General approach

The first estimates performed in January 2006 ("preliminary estimates") concerned the two following periods: Period A, from the detection of the first cases in April 2005 to mid-December 2005 when a large increase of cases occurred (March 28-December 18, 2005; 266 days); and Period B, from mid-December until the interruption of blood collection (December 19, 2005-January 22, 2006; 35 days; Fig. 2).

These estimates were later refined with consolidated incidence data, corrected for delayed care-seeking and delayed reporting and more precise estimates of the proportion of asymptomatic infections obtained through a seroepidemiologic survey carried out at the final phase of the outbreak ("retrospective estimates"). We also estimated the risk of viremic blood donation for five different

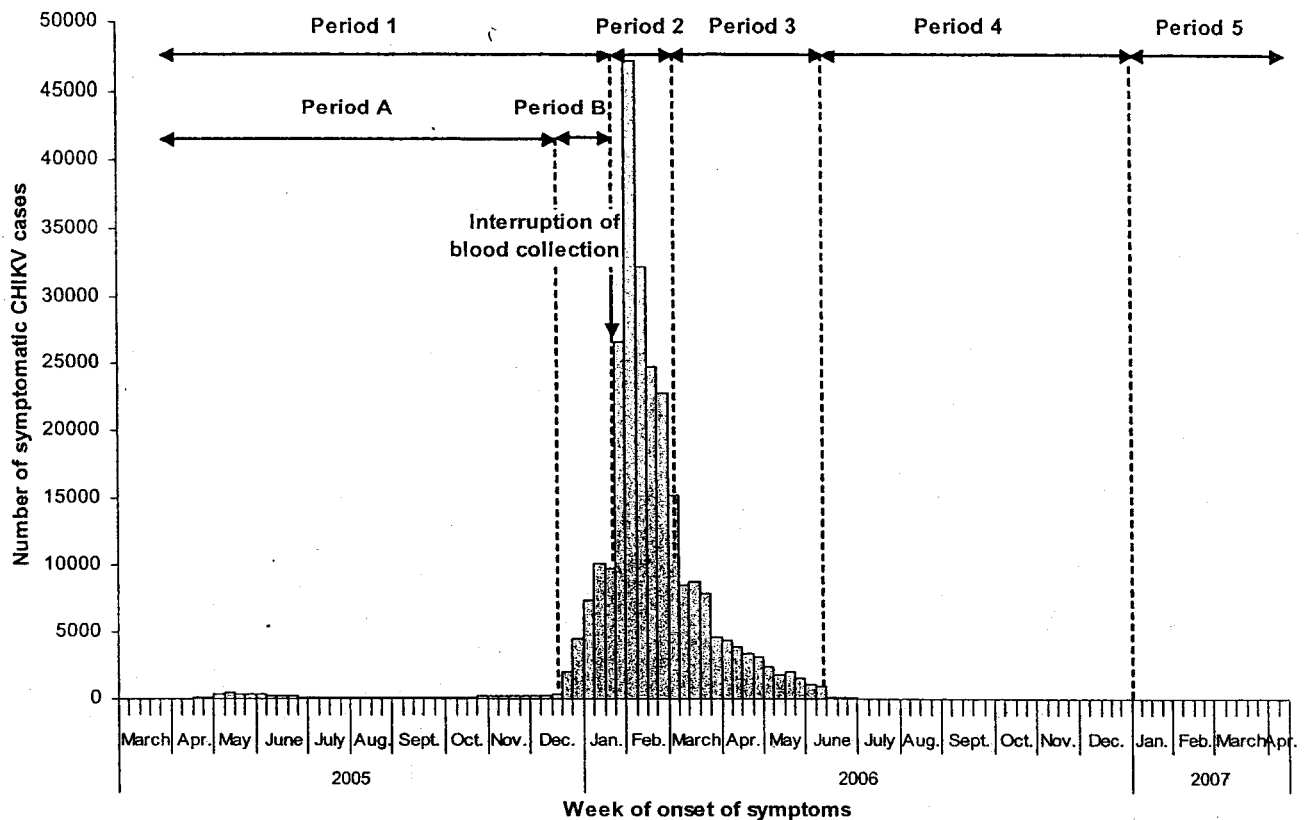


Fig. 2. Periods for risk estimates and distribution of symptomatic cases of CHIKV infection per week of onset of symptoms, Reunion Island, March 28, 2005, through April 15, 2007.

periods of the outbreak with these updated data (Fig. 2).

By use of the quarterly numbers of blood donations collected on Reunion Island in 2005 (unpublished data from EFS), we could then estimate the number of blood donations that would have been collected in 2006 if blood donations had not been interrupted.

To assess the validity of our risk estimates, we compared the estimated risk of viremic blood donation ("estimated risk") to the observed proportion of viremic PLT donations collected and screened for CHIKV genome ("observed risk") over the same period.

Statistical approach

An approximating formula developed by Biggerstaff and Petersen¹⁵ in 2002 for West Nile virus (WNV) was used to estimate the mean risk of viremic blood donation by CHIKV. This formula combines the proportion of asymptomatic (P_a) and symptomatic (P_s) infections with the duration of viremia among asymptomatic infected individuals (V_a) and the duration between onset of viremia and onset of symptoms in symptomatic patients (V_s). This provides the mean time an infected individual is viremic and asymptomatic. Dividing this mean duration of

viremia by the length of the outbreak period (L) then provides an estimate of the probability that an individual donates blood during viremia, assuming that a person with symptoms would self-defer or be excluded from donation by the predonation medical examination. Combined with the incidence (I) of the infection (including both symptomatic and asymptomatic infection), it gives an estimate of the mean risk of viremic blood donation:

$$\text{Mean risk} = \frac{(P_a \times V_a) + (P_s \times V_s)}{L} \times I.$$

As suggested by Biggerstaff and Petersen,¹⁵ risk confidence bounds were obtained by multiplying the confidence bounds of I by $[(P_a \times V_a) + (P_s \times V_s)]/L$. Confidence intervals (CIs) of I were calculated with Fleiss quadratic method.¹⁶

Data on duration of viremia

In January 2006, few data were available on the duration of CHIKV viremia. In 1964, Sarkar and coworkers¹⁷ described, from virologic studies of hemorrhagic fever in Calcutta, that CHIKV was most frequently isolated from blood within 48 hours after the onset of symptoms, but that it had been isolated as late as 6 days after the onset of illness.

The duration of viremia has been more extensively documented for dengue viruses: 1 or 2 days before the onset of symptoms and between 4 and 6 days and as late as 12 days after the first symptoms.¹⁸⁻²⁰ We thus used the following parameters for CHIKV: 1.5 days for the mean duration between onset of viremia and onset of symptoms among symptomatic patients (V_s) and $1.5 + 6 = 7.5$ days for the mean duration of viremia among asymptomatic infected individuals (V_a) assuming that the whole duration of viremia is similar in symptomatic and asymptomatic infections.

The same estimates of duration of viremia were used for the retrospective estimates since consistent observations were reported during the outbreak on Reunion Island. Thus, during this epidemic, CHIKV has been isolated from blood mostly within 5 days and as late as 12 days after the onset of symptoms. In some cases, CHIKV viremia might have persisted over 12 days since viral loads at 12 days were high.²¹

Data on the proportion of asymptomatic infections

For the preliminary estimates in January 2006, in the absence of data on the proportion of asymptomatic CHIKV infection, two hypotheses were formulated based on the proportion of asymptomatic infections reported during outbreaks of dengue:^{22,23} a minimal proportion of asymptomatic infection of 30 percent and a maximal proportion of 70 percent.

Between August and October 2006, a seroprevalence study was conducted among the general population of Reunion Island.²⁴ This survey showed that 38 percent of the inhabitants of Reunion Island had been infected by CHIKV. The preliminary results indicated that 6 percent of the study population had a positive CHIKV serology without having reported CHIKV symptoms. This suggests that approximately 15 percent of infected individuals during this outbreak may have had an asymptomatic infection. Therefore, this proportion of 15 percent was used for P_a for the retrospective estimates.

Incidence of CHIKV infection

We used the incidence data in the general population for the risk estimations assuming that potential blood donors had the same risk of CHIKV infection as the general population. The population of interest was the inhabitants of Reunion Island estimated at 756,745 by a population census conducted in 2004 by the National Institute for Statistics and Economics Studies (INSEE). CHIKV incidence data, by week of onset of symptoms, were obtained from the Reunion-Mayotte Interregional Epidemiology Unit, which had started surveillance for CHIKV infection as soon as the first cases were reported in April 2005. A suspect case of CHIKV infection was defined as a patient

with an abrupt onset of fever over 38.5°C associated with incapacitating arthralgia in the absence of any other potential cause of infection. From April to December 2005, surveillance relied on vector control teams, which conducted active and retrospective case-finding around the cases reported by a sentinel physician network, medical laboratories, private practitioners, and patients themselves. The number of cases took into account the symptomatic patients responding to the case definition whether or not they had consulted a general practitioner. During this period, approximately 67 suspect CHIKV cases were identified by active case-finding for every suspect case identified by the sentinel network physicians. From mid-December onward, the number of cases exceeded the capacity of the active surveillance system, and surveillance was then entirely based on the sentinel network. To estimate the total number of cases from the sentinel network data, the multiplier of 67, derived during the phase of active case finding, was used.²⁵

For the estimations of the risk of viremic donations, we calculated the estimated incidence of symptomatic and asymptomatic CHIKV infection by multiplying the estimated incidence of suspect cases by 100/(proportion of symptomatic infections).

RESULTS

Preliminary estimates

When the preliminary estimates were performed at the end of January 2006, the number of CHIKV suspect cases was 6500 for Period A and 25,000 for Period B. For Period A, the estimated mean risk of viremic blood donation was 15.2 per 100,000 donations, under the minimal hypothesis of 30 percent asymptomatic infections, and 61.3 per 100,000 donations, under the maximal hypothesis of 70 percent asymptomatic infections (Table 1). For Period B, the mean risk reached 445 per 100,000 donations, under the minimal hypothesis and 1,793 per 100,000 donations, under the maximal hypothesis.

Retrospective estimates

The retrospective estimates used the results of the seroprevalence survey that estimated the proportion of asymptomatic CHIKV infections during this outbreak at 15 percent. The updated estimate of the number of symptomatic cases was 6,864 for Period A and 34,002 for Period B (Table 2). Risk of viremic blood donation was then estimated at 9.6 and 362.5 per 100,000 donations for Periods A and B, respectively. The risk estimates for the five periods of the outbreak are shown in Table 3. Between the identification of the first CHIKV cases and the interruption of blood donations (Period 1), 7 of 14,450 blood donations collected could have been viremic. During

TABLE 1. Preliminary risk estimates of viremic blood donation, Reunion Island, March 28, 2005, through January 22, 2006

	Period A, Mar 28-Dec 18, 2005		Period B, Dec 19, 2005-Jan 22, 2006	
	Minimal hypothesis	Maximal hypothesis	Minimal hypothesis	Maximal hypothesis
Estimated number of symptomatic cases	6,500	6,500	25,000	25,000
Proportion of asymptomatic infections (%)	30	70	30	70
Estimated number of infected cases	9,286	21,667	35,714	83,333
Period length (days)	266	266	35	35
Estimated incidence of CHIKV infection per 100,000	1,227	2,863	4,720	11,012
Estimated risk of viremic blood donation				
Per 100,000 blood donations (95% CI)	15.2 (14.9-15.5)	61.3 (60.6-62.2)	445.0 (440.5-449.5)	1,793.4 (1,781.9-1,804.9)
Per estimated number of blood donations (95% CI)	2.0/12,800 (1.9-2.0)	7.9/12,800 (7.8-8.0)	7.1/1,600 (7.0-7.2)	28.7/1,600 (28.5-28.9)

TABLE 2. Retrospective risk estimates of viremic blood donation, Reunion Island, March 28, 2005, through January 22, 2006

	Period A, Mar 28- Dec 18, 2005	Period B, Dec 19, 2005- Jan 22, 2006
Estimated number of symptomatic cases	6,864	34,002
Proportion of asymptomatic infections (%)	15	15
Estimated number of infected cases	8,075	40,002
Period length (days)	266	35
Estimated incidence of CHIKV infection per 100,000	1,067	5,286
Estimated risk of viremic blood donation		
Per 100,000 blood donations (95% CI)	9.6 (9.4-9.8)	362.5 (359.0-366.0)
Per estimated number of blood donations (95% CI)	1.2/12,800 (1.2-1.3)	5.8/1,600 (5.7-5.9)

Period 2, at the height of the epidemic, the estimated risk of viremic blood donation was 1,500 per 100,000, that is, 29 potentially viremic donations if blood collection had continued. The estimated risk then decreased due to diminishing CHIKV transmission: 210 per 100,000 between March and June 2006 (Period 3), 1.4 per 100,000 for the second semester of 2006 (Period 4), and 0.27 per 100,000 for the first months of 2007 (Period 5), that is, 1 potentially viremic blood donation every 21 years on the basis of 17,500 blood donations collected each year. Finally, over the course of the outbreak, a total of 47 of 35,750 blood donations might have been viremic if blood collection had continued. Simultaneously, an estimated 312,500 of 757,000 inhabitants have been infected by mosquito-borne transmission.

Comparison between estimated risk and observed risk

Between January 23 and May 7, 2006, 2 of the 500 PLT donations screened for CHIKV RNA were positive (0.4%). One donor developed CHIKV symptoms on the day after the blood donation, the other remained asymptomatic. The risk of viremic blood donation over this period was estimated at 720 per 100,000 blood donations, that is, 0.72 percent.

Although an estimated 7 viremic donors had donated blood before the collection was interrupted, no case of transfusion-transmitted CHIKV infection has been identified during this period.

DISCUSSION

During this first and massive epidemic of CHIKV infection on Reunion Island, we computed estimates of the risk of CHIKV viremic blood donation, in real time during the ascending phase of the major epidemic peak, and afterward, we refined these estimates with newly available data. Although we underestimated the incidence of CHIKV infection in our preliminary calculations, we overestimated the proportion of asymptomatic infections. Consequently, the preliminary estimates were 1.2- to 6.4-fold greater than the retrospective calculations. The preliminary estimates, however, provided a right order of magnitude of the risk in real time in an emergency context. The retrospective calculations indicate a mean risk over the course of the outbreak, between April 2005 and April 2007, of 132 per 100,000 donations. The mean risk peaked at approximately 1,500 per 100,000 donations at the height of the outbreak in February 2006. In total, potentially 47 of 35,750 blood donations might have been viremic between April 2005 and April 2007 if blood collection had not been interrupted. We also estimated that 7 blood donations were viremic before the interruption of blood donations on the island. Therefore, this measure enabled the avoidance of 40 potentially viremic donations. By way of comparison, during the outbreak, the total number of individuals infected through mosquito-borne CHIKV transmission is estimated at 312,538 individuals.

This approach has several limitations. The estimates provided relate to a mean risk, which supposes that the risk is constant over the studied period and for the

TABLE 3. Retrospective risk estimates of viremic blood donation, Reunion Island, March 28, 2005, through April 15, 2007

	Period 1, Mar 28, 2005- Jan 22, 2006	Period 2, Jan 23, 2006- Mar 5, 2006	Period 3, Mar 6, 2006- Jun 11, 2006	Period 4, Jun 12, 2006- Dec 31, 2006	Period 5, Jan 1, 2007- Apr 15, 2007	Periods 1-5, Mar 28, 2005- Apr 15, 2007
Estimated number of symptomatic cases	40,866	169,008	54,936	772	75	265,657
Period length (days)	301	42	98	203	105	749
Proportion of asymptomatic infections (%)	15	15	15	15	15	15
Estimated number of infected cases	48,078	198,833	64,631	908	88	312,538
Estimated incidence of CHIKV infection per 100,000	6,353	26,275	8,541	120	12	41,300
Estimated risk of viremic blood donation Per 100,000 blood donations (95% CI)	50.7 (50.2-51.1)	1,501.4 (1,495.8-1,507.1)	209.2 (207.6-210.7)	1.4 (1.3-1.5)	0.27 (0.2-0.3)	132.3 (132.0-132.7)
Per-estimated number of blood donations (95% CI)	7.3/14,450 (7.3-7.4)	29.1/1,940 (29.0-29.2)	9.9/4,710 (9.8-9.9)	0.14/9,760 (0.13-0.15)	0.01/4,890 (0.01-0.02)	47.3/35,750 (47.2-47.4)

geographic area. Although estimates were performed for several periods selected according to the level of incidence, the number of cases and consequently the risk might have been highly variable during the studied period. In addition, the risk of infection varied by geographic area as later demonstrated by the seroprevalence survey that showed that 29.6 percent of the inhabitants of the North have been infected whereas in the East, this proportion reached 48 percent.²⁴ Consequently, the mean risk underestimates the maximal risk, corresponding to the peak of the outbreak and to the area where CHIKV transmission was maximal. This maximum risk, however, is highly time and space limited.

To obtain a more dynamic sight of the risk over the course of the epidemic and estimates of the maximal risk, it would have been necessary to develop an approach similar to the one proposed by Biggerstaff and Petersen^{15,26} for the WNV epidemic in 2002 in the United States. The latter is a statistical approach based on imputation and resampling techniques providing daily estimates of the risk of blood contamination in an epidemic setting. Conducting such an analysis in the context of this large and long-standing outbreak would have been computationally cumbersome. In our opinion, such a refinement was not essential in regard to the main objectives of the study, that is, providing a right order of magnitude of the risk as an aid for risk management. We considered that providing an approximation of the mean risk over five periods was a suitable alternative. To compute these mean risks, we therefore used the approximating formula proposed by Biggerstaff and Petersen.¹⁵ In 2003, Biggerstaff and Petersen demonstrated for the WNV epidemic in 2002 in the United States that the approximating formula provides a reasonable approximation to the mean risk of transfusion.¹⁵ The same work of comparison of the mean risks estimated by this method and by statistical resampling was carried out, in the setting of our workgroup, for an outbreak of acute hepatitis A in France that occurred in 1996 through 1997.^{14,27} It also concluded to a good concordance of the results of both methods. Note that the CIs presented with our mean risk estimates do not take into account the uncertainty on the duration of viremia, the proportion of asymptomatic infections, nor the coefficient of 67, used to estimate incidence of symptomatic infections from the sentinel network data. Even though this limitation led to artificially narrow CIs, point estimates of mean risk should not be affected.

Our incidence data were derived from a sentinel surveillance system. Because a clinical case definition was used, it is possible that other febrile illnesses, not due to CHIKV, were included in the case count. The positive predictive value of a clinical case definition, however, greatly improves if incidence is high. Therefore, the inclusion of noncases in the case count, leading to

overestimation of the incidence and hence the risk of viremic donation, is more likely to occur outside an outbreak setting. The validity of the extrapolated data derived from a sentinel surveillance system estimating the total number of cases in the community should also be questioned. The serosurvey estimated that 38 percent of the inhabitants had been infected and that 32 percent had suffered from symptomatic infections. These data are consistent with the 35 percent of the inhabitants having suffered from symptomatic illness, estimated by the sentinel surveillance system and corroborate our incidence estimates.

We assumed that potential blood donors had the same risk of CHIKV infection as the general population. This assumption was supported by the findings of the serosurvey that showed similar antibody prevalences among adults of both sexes.²⁴ In addition, when we applied age-specific CHIKV antibody prevalence rates of the serosurvey to the donor population of Reunion Island, the overall seroprevalence among donors was estimated at 37.2 percent, similar to the overall antibody prevalence in the general population (38%).

One major limitation of the validity of our estimates relates to lack of a precise knowledge on the distribution of the duration of asymptomatic viremia in individuals with apparent and inapparent infection. To refine the estimates, further studies are necessary to document the kinetics of CHIKV viremia. This approach also hypothesizes that symptomatic individuals would self-defer or be excluded by the predonation examination. In real life, this may not always be the case. In the United States, among the first 14 identified donors associated with transfusion-related WNV transmission to recipients, 3 were shown to have been symptomatic at the moment of the donation.¹⁵ Nevertheless, for CHIKV infection which is characterized by sudden onset of symptoms, this assumption is more plausible than for WNV which frequently causes paucisymptomatic infection.

Lack of data on the frequency of asymptomatic infection was the most important limiting factor for the preliminary estimates. This variable has a preponderant role in the risk estimate since it contributes both in the computation of the weighted mean of the duration of asymptomatic viremia and in the estimate of the incidence of infection. Valid data were available, however, for the retrospective calculations from the seroprevalence survey. This survey provided an estimate of the proportion of asymptomatic infections obtained directly among the studied population and for the epidemic CHIKV strain circulating.

In spite of the above limitations, the retrospective estimates are likely to give a good approximation of the real risk, as suggested by the observed risk of viremic PLT donations. From January to May 2006, this observed risk was 400 per 100,000 donations, of the same order of mag-

nitude as the risk of 720 per 100,000 donations estimated over the same period.

Up to date, CHIKV infections from transfusion of blood or blood components have not been reported in the literature. On Reunion Island, no case of transfusion-transmitted CHIKV infection has been identified in spite of the estimated seven viremic donations collected before donations were interrupted. Despite the lack of data about transfusion-transmitted CHIKV infection, the high viral load during the acute phase of the infection,^{21,28} the fact that several cases of CHIKV transmission have occurred among laboratory personnel handling infected blood,²⁹ and the fact that CHIKV has been transmitted to a health care worker drawing blood from an infected patient²⁸ provide evidence that transfusion-related transmission of CHIKV is highly plausible. It is possible that transfusion-related infections have not been recognized or have not been distinguished from infection from mosquito vectors. Also, the true transmission rate from viremic donors to recipients is not known. Several issues may influence the possibility of transmission of CHIKV through transfusion, such as the stability of the virus during storage of blood and the efficiency of virus elimination of blood processing methods, as viral inactivation. Also, the presence of IgM or IgG antibodies in donor blood may neutralize infectivity, as demonstrated for other viruses such as parvovirus³⁰ and suggested for WNV.³¹ In addition, the assessment of the risk of CHIKV transmission from a viremic donor to a recipient would need to take into account the recipient's immune status.

In conclusion, despite the absence of documented cases, blood transfusion-related CHIKV transmission is plausible and the risk of viremic donation can be substantial in an outbreak setting. During this large outbreak, the estimated risk of viremic blood donation was high, but low compared to the risk of mosquito-borne CHIKV transmission. Despite its limitations, this work provided a right order of magnitude of the risk of viremic blood donation in real time during the ascending phase of the epidemic peak. At this moment, the decision of interrupting blood collection relied on the precautionary principle. The low risk estimated for early 2007 was, however, useful to contribute to the decision making process to start again the collection of blood donations on the island from June 14, 2007. This illustrates how this approach may contribute to guiding prevention measures.

ACKNOWLEDGMENTS

The authors thank Reunion-Mayotte Interregional Epidemiology Unit (Reunion-Mayotte Cire) and the Reunion Regional Observatory of Health plus sentinel general practitioners for providing the case onset data and for their important work during this massive and long outbreak.

The workgroup "Quantitative Estimation of the Risk of Blood Donation Contamination by Infectious Agents" consists of (in alphabetical order):

Agence Française de Sécurité Sanitaire des Produits de Santé (Afssaps, French Agency for the Safety of Health Products): J.F. Legras, M. Martin, E. Pouchol, I. Sainte-Marie
 Etablissement Français du Sang (EFS; French Blood Services): A. Assal, P. Biagini, M.H. Elghouzzi, P. Gallian, P. Morel
 Institut National de Transfusion Sanguine (INTS): S. Laperche
 Institut de Veille Sanitaire (InVS) (French Institute of Public Health Surveillance): P. Bernillon, F. Biton, C. Brouard, I. Capek, E. Delarocque-Astagneau, H. de Valk, D. Jeannel, A. Mailles, J. Pilonel, I. Quatresous, L. Sanchez, N. Schwarz, V. Vaillant

REFERENCES

- Robinson MC. An epidemic of virus disease in Southern Province, Tanganyika Territory, in 1952-53. I. Clinical features. *Trans R Soc Trop Med Hyg* 1955;49:28-32.
- Lumsden WH. An epidemic of virus disease in Southern Province, Tanganyika Territory, in 1952-53. II. General description and epidemiology. *Trans R Soc Trop Med Hyg* 1955;49:33-57.
- Jupp PG, McIntosh BM. Chikungunya virus disease. In: Monath TP, editor. *The arboviruses: epidemiology and ecology*. Boca Raton (FL): CRC Press; 1988. p. 137-57.
- Pastorino B, Muyembe-Tamfum JJ, Bessaud M, Tock F, Tolou H, Durand JP, Peyrefitte CN. Epidemic resurgence of Chikungunya virus in democratic Republic of the Congo: identification of a new central African strain. *J Med Virol* 2004;74:277-82.
- Thonnon J, Spiegel A, Diallo M, Diallo A, Fontenille D. Chikungunya virus outbreak in Senegal in 1996 and 1997. *Bull Soc Pathol Exot* 1999;92:79-82.
- Lanciotti RS, Ludwig ML, Rwaguma EB, Lutwama JJ, Kram TM, Karabatsos N, Cropp BC, Miller BR. Emergence of epidemic O'nyong-nyong fever in Uganda after a 35-year absence: genetic characterization of the virus. *Virology* 1998;252:258-68.
- Saluzzo JF, Gonzalez JP, Herve JP, Georges AJ. Epidemiological study of arboviruses in the Central African Republic: demonstration of Chikungunya virus during 1978 and 1979. *Bull Soc Pathol Exot Filiales* 1980;73:390-9.
- Carey DE, Myers RM, DeRanitz CM, Jadhav M, Reuben R. The 1964 chikungunya epidemic at Vellore, South India, including observations on concurrent dengue. *Trans R Soc Trop Med Hyg* 1969;63:434-45.
- Porter KR, Tan R, Istary Y, Suharyono W, Sutaryo Widjaja S, Ma'Roef C, Listiyangingsih E, Kosasih H, Hueston L, McArdle J, Juffrie M. A serological study of Chikungunya virus transmission in Yogyakarta, Indonesia: evidence for the first outbreak since 1982. *Southeast Asian J Trop Med Pub Health* 2004;35:408-15.
- Nimmannitya S, Halstead SB, Cohen SN, Margiotta MR. Dengue and chikungunya virus infection in man in Thailand, 1962-1964. I. Observations on hospitalized patients with hemorrhagic fever. *Am J Trop Med Hyg* 1969;18:954-71.
- Thein S, La Linn M, Aaskov J, Aung MM, Aye M, Zaw A, Myint A. Development of a simple indirect enzyme-linked immunosorbent assay for the detection of immunoglobulin M antibody in serum from patients following an outbreak of chikungunya virus infection in Yangon, Myanmar. *Trans R Soc Trop Med Hyg* 1992;86:438-42.
- Borgherini G, Poubeau P, Staikowsky F, Lory M, Le Moullec N, Becquart JP, Wengling C, Michault A, Paganin F. Outbreak of chikungunya on Reunion Island: early clinical and laboratory features in 157 adult patients. *Clin Infect Dis* 2007;44:1401-7.
- Lambert J, Couturier E, Vaillant V. Infection à chikungunya. Etude descriptive des cas importés en France métropolitaine, 2005-2006. Saint-Maurice: Institut de Veille Sanitaire; 2007.
- Brouard C, De Valk H, Pilonel J; Groupe "Estimation Quantitative du Risque de Contamination d'un Don de Sang par des Agents Infectieux." Estimation quantitative du risque de contamination d'un don de sang par des agents infectieux. Saint-Maurice: Institut de Veille Sanitaire; 2007.
- Biggerstaff BJ, Petersen LR. Estimated risk of transmission of the West Nile virus through blood transfusion in the US, 2002. *Transfusion* 2003;43:1007-17.
- Fleiss JL. *Statistical methods for rates and proportions*. 2nd ed. New York: Wiley; 1981.
- Sarkar JK, Pavri KM, Chatterjee SN, Chakravarty SK, Anderson CR. Virological and serological studies of cases of haemorrhagic fever in Calcutta. Material collected by the Calcutta school of tropical medicine. *Indian J Med Res* 1964;52:684-91.
- Vaughn DW, Green S, Kalayanarooj S, Innis BL, Nimmannitya S, Suntayakorn S, Rothman AL, Ennis FA, Nisalak A. Dengue in the early febrile phase: viremia and antibody responses. *J Infect Dis* 1997;176:322-30.
- Gubler DJ, Suharyono W, Tan R, Abidin M, Sie A. Viraemia in patients with naturally acquired dengue infection. *Bull World Health Organ* 1981;59:623-30.
- Guzman MG, Kouri G. Dengue: an update. *Lancet Infect Dis* 2002;2:33-42.
- Laurent P, Le Roux K, Grivard P, Bertil G, Naze F, Picard M, Staikowsky F, Barau G, Schuffenecker I, Michault A. Development of a sensitive real-time reverse transcriptase PCR assay with an internal control to detect and quantify chikungunya virus. *Clin Chem* 2007;53:1408-14.
- Waterman SH, Novak RJ, Sather GE, Bailey RE, Rios I, Gubler DJ. Dengue transmission in two Puerto Rican communities in 1982. *Am J Trop Med Hyg* 1985;34:625-32.
- McBride WJ, Mullner H, LaBrooy JT, Wronski I. The 1993 dengue 2 epidemic in North Queensland: a serosurvey and comparison of hemagglutination inhibition with an ELISA. *Am J Trop Med Hyg* 1998;59:457-61.

24. Perau J, Catteau C, Michault A, Parain C, Favier F. Fin 2006, 300 000 personnes avaient été atteintes par le chikungunya. *Economie de la Réunion* 2007;129:16-7.
25. Renault P, Solet JL, Sissoko D, Balleydier E, Larrieu S, Filleul L, Lassalle C, Thiria J, Rachou E, De Valk H, Ilef D, Ledrans M, Quatresous I, Quenel P, Pierre V. A major epidemic of chikungunya virus infection on Reunion Island, France, 2005-2006. *Am J Trop Med Hyg* 2007;77:727-31.
26. Biggerstaff BJ, Petersen LR. Estimated risk of West Nile virus transmission through blood transfusion during an epidemic in Queens, New York City. *Transfusion* 2002;42:1019-26.
27. Sanchez-Garrido L, Bernillon P, Delarocque-Astagneau E, Brouard C, Pillonel J, Santa-Olalla P, De Valk H, Desenclos JC. Modélisation du risque de contamination des dons de sang par le virus de l'hépatite A. *Journées de veille sanitaire* 2005, 29-30 November; Paris.
28. Parola P, De Lamballerie X, Jourdan J, Rovey C, Vaillant V, Minodier P, Brouqui P, Flahault A, Raoult D, Charrel RN. Novel chikungunya virus variant in travelers returning from Indian Ocean islands. *Emerg Infect Dis* 2006;12:1493-9.
29. Centers for Disease Control and Prevention and National Institute of Health. *Biosafety in Microbiological and Biomedical Laboratories (BMBL)*. 5th ed. Washington (DC): US Government; 2007.
30. Parsyan A, Candotti D. Human erythrovirus B19 and blood transfusion—an update. *Transfus Med* 2007;17:263-78.
31. Busch MP, Caglioti S, Robertson EF, McAuley JD, Tobler LH, Kamel H, Linnen JM, Shyamala V, Tomasulo P, Kleinman SH. Screening the blood supply for West Nile virus RNA by nucleic acid amplification testing. *N Engl J Med* 2005;353:460-7. ■

