living conditions. Half the population included arrived in Spain after 2000, a fact that illustrates the increasing immigration rates from Latin America observed over the past years. Another interesting result from the questionnaire was that the information obtained about living conditions in the Chagas disease endemic area (rural area, adobe house) did not correlate with the presence or absence of antibodies to T. cruzi. People born in endemic regions (7 of 11 positive donors) generally declared that they had never lived in a rural environment or an adobe house (Table 2), as is commonly assumed. Hence, this question is not useful for differentiation purposes. Interestingly, the same conclusion was drawn from the Berlin study, in which 95 of 100 immigrants declared that they came from an urban area, including the 5 cases of confirmed Chagas disease.21

The two serologic assays used in this study were chosen because at the beginning of the study they were commercially available and EC-marketed. Both are based on recombinant antigens, whereas the third conventional in-house ELISA is based on whole parasite lysate. All samples confirmed as positive had been initially reactive with both recombinant antigens assays, and all samples initially reactive with only one assay presented a nonreactive result in the in-house ELISA and were considered false-positive samples. It is worth noting that many discrepant results observed between both assays corresponded to low 0.9 to 1 signal-to-cutoff rates for bioelisa Chagas (Biokit) or doubtful reactions with ID-PaGIA (DiaMed), which were all considered as initially reactive in this study. Additionally, it should be mentioned that the T. cruzi ELISA test system performed on all initially reactive samples (with one or two tests) confirmed the results obtained with the conventional in-house ELISA. The high rate of inconclusive or false-positive results obtained when one diagnostic test is used underscores the need to confirm all initially positive results with a second serologic technique. In any case, there is still a need for a real confirmatory test to overcome the issues of discrepancies and false results (positive or negative). The ID-PaGIA assay allows testing of a small number of samples at a time. Although this system has the drawback of rather subjective reading, it could be useful in blood centers with a small volume of donations and is now even more reliable since a third antigen has been recently added to increase the sensitivity of the test. The ELISA format, which allows for automation and objective reading, should be indicated in other blood centers. An even more appropriate strategy would be the use of two screening tests, one based on recombinant antigens and the other on crude antigens.20

In summary, this study reports a seroprevalence of *T. cruzi* infection of 0.62 percent among at-risk donors in Catalonia and emphasizes the need to include individuals who have resided in, but were not necessarily born in

endemic areas as at-risk donors. The difficulty of this type of selective screening is proper identification of the risk population, which essentially depends on the predonation interview. Latin Americans accounted for more than 1 percent of the total of donors in our study, and this substantial contribution underscores the need to accept them as donors.

In the future, techniques to inactivate or reduce the parasite load, which are currently under development or evaluation, ^{29,30} might be applicable to blood components. At this time, however, detection of *T. cruzi* infection is the only preventive measure available to accept at-risk blood donors.

ACKNOWLEDGMENTS

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医薬品 研究報告 調査報告書

应采印·则九秋口·阙里取口官										
識別番号·報告回数			報告日	第一報入手日 2008. 10. 15	新医薬品等の区分 該当なし	総合機構処理欄				
一般的名称 人全血液			BuaNews online, Moi	n 13 Oct 公表国						
販売名(企業名)		赤」(日本赤十字社) 日赤」(日本赤十字社))	研究報告の公表状況							
研究報告の 南アフリカ、ヨシンと ウイルスリカで国際にイアフリカでの 南で、カラックので、カラックの で、カラックののが、カーのでの 1人目間のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人目のと 1人日のと 1日	特定された。ウイルス 染症研究所(NICD) 措置が必要である」と うか、ならびに当該ら に存在することはま は、9月中旬に重篤 の患者の搬送に同行 がした他の3名の患 がインフルエンザ様症	と者を出したウイルスは感染マウスの排泄と保健省は共同で、との声明を発表した。 との声明を発表した。 とイルスの分布についだ示されていないといるななでずンビアカテムを対象ですができる。 者は退院したことができまれる。 を発症した教急を	は、暫定的に西アフリカのは物を介し、人間の食物を このウイルスが体液を介し 3名の死因を確定するにいて検討を行う必要があるいICDは述べた。 いら搬送され、Morningside 形亡し、間もなく看護師が確認されているが、依然と 数命士であり、もう1名は20、現在は安定している。	ハウスダストを汚染ー してヒトからヒトに感染 は更なる検査が必要 。ヒトに疾患を引き起 Medi-Clinicに入院 死亡した。 して2名が厳重な監	する可能性がある。 まするため、「患者の看護 である。新たなアレナウ とすアレナウイルスが南 し、2日後に死亡した。約 現下に置かれている。1	使用上の注意記載状況・その他参考事項等 人全血液-LR「日赤」 照射人全血液-LR「日赤」 血液を介するウイルス、 細菌、原虫等の感染 vCJD等の伝播のリスク				
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Compiled by the Government Communication and Information System Date: 13 Oct 2008

Title: Unknown illness identified as Arenavirus

By Luyanda Makapela

Johannesburg - The virus which has caused the death of three people has been provisionally identified as the rodentborne Arenavirus.

The Arenavirus, related to the Lassa Fever Virus of West Africa, causes chronic infections in multimammate mice. Infected mice's excretion contains the virus which can contaminate human food or house dust.

A joint statement by the National Institute for Communicable Disease (NICD) and the Department of Health explained that the Arenavirus is a disease spread from human to human through the contact of body fluids.

"Special precautions are required in nursing patients," a statement said.

The finding follows blood samples being sent to Atlanta, in the United States to determine the cause of the deaths of three people who had been suspected of contracting Viral Haemorrhagic Fever.

The virus is similar to Lassa Fever, the department said. It has previously been found in rodents elsewhere in Africa, but has not been found to cause disease in humans other than in West Africa.

Further tests are needed to confirm the diagnosis by growing the virus in culture.

"It needs to be determined whether it is a previously unrecognised member of the Areaviruses, and what its distribution is. There is no indication as yet that Arenaviruses which cause disease in humans are present in South African rodents," the NICD said.

The first victim, who had to be flown in from Zambia in a critical condition, was admitted to the Morningside Medi-Clinic in mid September. She died two days later.

About two weeks later, the paramedic who had flown in with the first victim, was admitted at the same clinic presenting the same symptoms.

A nurse, Gladys Mthembu died shortly afterwards. According to certain reports Ms Mthembu's family has been given a go-ahead to continue with the funeral arrangements as her bedroom had been cordoned off by health officials

Maria Mokubung, a cleaner at the Morningside Medi-Clinic, who also died last weekend has since been ruled out as a possible victim of the virus

Meanwhile the Gauteng Health Department has confirmed that the three other patients, including nurse's female supervisor, who had been under observation for showing symptoms of the virus have been discharged.

They had been in contact with the nurse who died.

However, departmental spokesperson Phumelele Kaunda said there were two contacts that were still under active surveillance after being admitted for observation.

The one patient is a paramedic who had contact with the first patient and developed fever and flu-like symptoms. He was admitted initially in Flora Clinic and then transferred to Morningside Medi-Clinic with a diagnosis of kidney stones.

The other patient is a nurse who attended to the second patient and developed signs and symptoms similar to the first three patients. She is being treated in isolation and received the anti-viral medication, ribavirin. The patient is presently stable.

Gauteng Health MEC Brian Hlongwa meanwhile has sent condolences to the families of those that were killed by the viral infection, particularly families of health professionals who died in the line of duty.

"This illustrates the dedication of our health professionals and the need to society to respect and honour the work that they do," said MEC Hlongwa.

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RSS...RSS....RSS..... What is RSS feed? Click here to find out He also thanked the NICD, the National Health Laboratory Service, Centre for Disease Control in Atlanta and the World Health Organisation for ensuring that the results were made available soon. - BuaNews

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①人血清アルブミン、②人血清アルブミン、③人血清アルブミン*、④人免役グロブリン、⑤乾燥ペプシン処理人免疫グロブリン、⑥乾 燥スルホ化人免疫グロブリン、⑦乾燥スルホ化人免疫グロブリン*、⑧乾燥濃縮人活性化プロテインC、⑨乾燥濃縮人血液凝固第WI因子、 般 的 名 称|⑩乾燥濃縮人血液凝固第IX因子、⑪乾燥抗破傷風人免疫グロブリン、⑫抗 HBs 人免疫グロブリン、⑬トロンビン、⑭フィブリノゲン加 第XⅢ因子、⑩乾燥濃縮人アンチトロンビンⅢ、⑩ヒスタミン加人免疫グロブリン製剤、⑪人血清アルブミン*、⑱人血清アルブミン*、 ①献血アルブミン 20 "化血研"、②献血アルブミン 25 "化血研"、③人血清アルブミン "化血研"*、④ "化血研"ガンマーグロブリン、 ⑤献血静注グロブリン"化血研"、⑥献血ベニロン- I、⑦ベニロン*、⑧注射用アナクトC2,500 単位、⑨コンファクトF、⑩ノバクト 版 売 名(企 業 名) M、⑪テタノセーラ、⑫へパトセーラ、⑬トロンビン"化血研"、⑭ボルヒール、⑯アンスロビンP、⑯ヒスタグロビン、⑪アルブミン 20%化血研*、®アルブミン 5%化血研*、⑩静注グロブリン*、⑳ノバクトF*、㉑アンスロビン P 1500 注射用 アレナウイルス属は、エンベロープをもつ 1 本鎖 RNA(-)ウイルスである。齧歯類に寄生し、慢性腎臓感染をおこす。齧歯類の尿中 は高ウイルス価であり、ヒトの食品やハウスダストを汚染する。曝露したヒトは偶発的宿主となる。このウイルスの原型はリンパ球性脈 絡膜髄膜炎ウイルス(LCMV)であり、ヒトに感染するとインフルエンザ様症状、無菌性髄膜炎もしくは重症髄膜脳炎を発症する。出血 熱症候群の原因となる Arenaviruses は南米(New World arenaviruses)から数多く報告されている。いわゆる Old World arenaviruses は世界中に分布する LCMV と、西アフリカのナイジェリア、シエラレオネ、リベリア、ギニアを中心に1年間に最大50万人が感染し、 実際にはさらに広い地域に分布すると見られているラッサ熱ウイルスである。ラッサ熱ウイルス感染の臨床症状としては、不顕性、軽症 発熱性疾患から劇症出血性疾患まで様々であり、致死率は一般的な社会環境における1~2%から、入院患者では20%、院内感染では40% 以上に及ぶこともある。西アフリカ一帯に生息する野ネズミの一種であるマストミス(Mastomys natalensis)は、ラッサ熱ウイルスの 最重要宿主であり、その分布は、西アフリカから東アフリカ一帯と、南アフリカ北東端まで南に広がっている。他の Mastomys 種とも 分布域が重複し、アレナウイルスは過去にはアフリカ南部の齧歯類でも確認されている。 報告企業の意見 (http://www.forth.go.jp/cgi-bin/promed/search.cgi?title_link=20081029-0050&button_detail=on) 弊所の血漿分画製剤の製造工程には、冷エタノール分画工程、ウイルス除去膜ろ過工程あるいは加熱工程等の原理の異なるウイルス除 去及び不活化工程が存在しているので、ウイルスクリアランスが期待される。 - 各製造工程のウイルス除去・不活化効果は、「血漿分画製剤のウイルスに対する安全性確保に関するガイドライン(医薬発第 1047 号、 平成 11 年 8 月 30 日)」に従い、ウシウイルス性下痢ウイルス (BVDV)、仮性狂犬病ウイルス (PRV)、ブタパルボウイルス (PPV)、A 型肝炎ウイルス (HAV) または脳心筋炎ウイルス (EMCV) をモデルウイルスとして、ウイルスプロセスバリデーションを実施し、評 価を行っている。今回報告したアレナウイルス属は、エンベロープの有無、核酸の種類等からモデルウイルスとしては BVDV が該当す ると考えられるが、上記バリデーションの結果から、BVDVの除去・不活化効果を有することを確認している。 また、これまでに当該製剤によるアレナウイルス感染の報告例は無い。

以上の点から、当該製剤はアレナウイルスに対する安全性を確保していると考える。



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Subject PRO/AH/EDR> Undiagnosed fatalities - S. Africa ex Zambia (10); arenavirus

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Arena virus outbreak, South Africa - Update

This updates all previous reports and includes available data as of 24 Oct 2008. An outbreak of infection due to an arenavirus was identified in South Africa in early October 2008. A total of 5 cases has been reported for the period 12 Sep to 24 Oct 2008.

The primary case (case 1) had onset of illness on 2 Sep 2008. An additional 3 secondary cases (case 2, 3 and 4) and 1 tertiary case (case 5) have been confirmed to have an arenavirus infection by laboratory testing. The primary case and 3 secondary cases have died. The tertiary case is currently hospitalized. Ages of cases ranged from 33 to 47 years. 4 cases were female and 1 male. The source of infection is, as yet, unknown for the primary case. The other 4 cases all had potential exposure to blood and/or body fluids of a primary or secondary case in the health—care setting.

The primary case was a safari booking agent resident in Zambia. The patient was flown to South Africa for medical care in a critically ill condition on 12 Sep 2008, and died on 14 Sep 2008. Case 2 was a paramedic who cared for case 1 during the transfer from Zambia on 12 Sep 2008 and case 3 was a nurse who cared for case 1 in the intensive care unit from 12-14 Sep 2008. Case 2 was admitted on 27 Sep 2008 and died on 2 Oct 2008 and case 3 was admitted on 30 Sep 2008 and died on 5 Oct 2008. On 14 Sep 2008, case 4 performed terminal cleaning of the room in which case 1 was hospitalized. The 5th patient is a nurse who . cared for case 2 from 27 Sep 2008 to 2 Oct 2008. She became ill on 9 Oct 2008 and is currently critical but stable. Ribavirin has been used for treatment in this case based on good evidence of efficacy in patients with Lassa fever (an arenavirus infection). The estimated incubation period (interval from exposure to symptom onset) in secondary and tertiary cases ranges from 7 to 13 days, In 4 patients who died, the interval from onset of illness to death ranged from 9 to 12 days (Figure 1).

Only limited clinical data are currently available for case 4; who presented late in the course of illness with bleeding and confusion and died soon thereafter. Clinical features of the remaining 4 cases, for which more clinical data were available, are presented. All patients presented initially with a non-specific flu-like illness with symptoms of fever headache and myalgia. The illness increased in severity over 7 days with all 4 patients developing diarrhoea and pharyngitis during the course of illness. A morbiliform rash on the face and trunk was reported in 4 cases on day 6 – 8 of illness. Facial swelling occurred in 3 patients. There appeared to be an initial clinical improvement after hospital admission in 3 patients, followed by clinical deterioration. Sudden and rapid deterioration

with respiratory distress, neurological signs and circulatory -collapse were terminal features in all patients who died. Bleeding was not a prominent feature. However, one patient had a petechial rash and another had oozing of blood from venepuncture sites. Chest pain was reported in case 1.

At the time of admission all patients had thrombocytopenia (range: 42-104 X109/L). Liver transaminases (AST and ALT) were available for 4 of 5 cases and were variable at the time of admission, however all 4 patients had raised AST and ALT during the course of their illness. Leucopenia was present on admission in 2 patients and 3 patients had a normal white blood cell count on admission. 4 patients subsequently developed leucocytosis during the course of hospitalisation. All contacts (family members, friends and healthcare staff) are being monitored with twice daily temperature measurements for a period of 21 days after the last exposure to a known case. In addition, safe burial of the deceased has been supervised by environmental health officers. Full personal protective equipment (PPE) and isolation precautions as per VHF protocols have been instituted.

The causative agent in this outbreak was initially identified as an Old World arenavirus by immunohistochemical tests performed at the Infectious Diseases Pathology Branch of the Centers for Disease Control and Prevention in Atlanta, USA, and on autopsy liver and skin samples taken with biopsy needles and skin punches in the Special Pathogens Unit of the National Institute for Communicable Diseases. National Health Laboratory Service, Sandringham (SPU-NICD/ NHLS), South Africa, from cases 2 and 3 on 9 Oct 2008 under biosafety level 4 laboratory conditions. Subsequently, infection with an Old World arenavirus has been confirmed in all 5 cases by positive PCR results and virus isolation by SPUNICD/ NHLS and CDC. Analysis of sequencing data generated at SPU-NICD/NHLS, Columbia University, New York, and CDC, Atlanta appears to indicate that the current outbreak is caused by a unique Old World arenavirus.

There are currently no additional suspected cases. The outbreak appears to be contained and has been confined to individuals with very close contact in a health-care setting. Monitoring of contacts, active case finding and investigation and management of suspected cases will continue as needed. Further characterization of the causative agent is under way and investigation into the source of infection in the primary case is required. Additional studies to determine whether mild/asymptomatic infection occurred amongst close contacts and other exposed individuals would be essential in better characterizing the extent of this outbreak and clinical spectrum of disease:

Arenaviruses are a family of enveloped negative sense single-stranded RNA viruses. Members of the family are parasites of rodents, in which they establish chronic renal infection. High titres of virus are present in rodent urine, which can contaminate human food or house dust. Exposed humans may become infected as accidental hosts. The prototype of the family is lymphocytic choriomeningitis (LCM) virus and infection of humans with this virus may present as an influenza-like illness, aseptic meningitis or severe meningo-encephalomyelitis. Arenaviruses which cause a haemorrhagic fever syndrome are well documented in South America (New World arenaviruses, including Junin, Machupo, Sabia and Guanarito viruses). The so-called Old World arenaviruses include LCM which in fact has a worldwide distribution, and Lassa fever virus which affects up to 500 000 people annually in West Africa, specifically in Nigeria, Sierra Leone, Liberia and Guinea, but the virus is suspected to be more widely distributed in that region.

The clinical spectrum of Lassa fever virus infection ranges from inapparent, through mild febrile illness to fulminant haemorrhagic disease, and mortality rates vary from 1-2 percent among cases in the community at large, through 20 percent among hospitalized patients, to >40 percent in nosocomial outbreaks. The multimammate mouse (Mastomys natalensis), which is the most important host of Lassa fever virus, has a distribution extending from West Africa across to East Africa and from there southwards to the northeasterncomer of South Africa. Its distribution overlaps with that of other Mastomys species, and arenaviruses have been found in southern African rodents in the past, but there has been no previous association of these viruses with human disease despite sustained monitoring. Preliminary

testing indicates that the virus associated with the present nosocomial disease outbreak is a distinct new member of the family.

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[This update provides a definitive account of the recent outbreak of arenavirus—associated disease in South Africa. A primary case (case 1) had onset of illness on 2 Sep 2008. An additional 3 secondary cases (case 2, 3 and 4) and 1 tertiary case (case 5) have been confirmed to have an arenavirus infection by laboratory testing. Case 5 (not previously reported) is a nurse who cared for case 2 from 27 Sep 2008 to 2 Oct 2008. She became ill on 9 Oct 2008 and is currently critical but stable. Cases 1, 2, 3 and 4 did not survive infection.

Infection with an Old World arenavirus has been confirmed in all 5 cases by positive PCR results and virus isolation by SPUNICD/ NHLS and CDC. Analysis of sequencing data generated at SPU-NICD/NHLS, Columbia University, New York, and CDC, Atlanta, appears to indicate that the current outbreak is caused by a unique Old World arenavirus.

There are currently no additional suspected cases. The outbreak appears to be contained and has been confined to individuals with very close contact in a health-care setting. Monitoring of contacts, active case finding and investigation and management of suspected cases are continuing. Further characterization of the causative agent is under way, as is investigation into the source of infection in the primary case.

- Mod.CP]

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[see also:

Undiagnosed fatalities – S. Africa ex Zambia (09): arenavirus 20081018.3300
Undiagnosed fatalities – S. Africa ex Zambia (08): arenavirus 20081013.3241
Undiagnosed fatalities – S. Africa ex Zambia (07): arenavirus 20081012.3234
Undiagnosed fatalities – South Africa ex Zambia (06): WHO 20081010.3211
Undiagnosed fatalities – South Africa ex Zambia (05) 20081008.3192
Undiagnosed fatalities – South Africa ex Zambia (04) 20081008.3188
Undiagnosed fatalities – South Africa ex Zambia (03) 20081007.3178
Undiagnosed fatalities – South Africa ex Zambia (02) 20081006.3157
Undiagnosed fatalities – South Africa ex Zambia: RFI 20081005.3139
Undiagnosed fatalities – South Africa ex Zambia: RFI 20081005.3139

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