We investigated the extent of lymphocytic choriomeningitis virus (LCMV) infection in employees and rodents at 3 commercial breeding facilities. Of 97 employees tested, 31 (32%) had IgM and/or IgG to LCMV, and aseptic meningitis was diagnosed in 4 employees. Of 1,820 rodents tested in 1 facility, 382 (21%) mice (Mus musculus) had detectable IgG, and 13 (0.7%) were positive by reverse transcription PCR; LCMV was isolated from 8. Rats (Rattus norvegicus) were not found to be infected. S-segment RNA sequence was similar to strains previously isolated in North America. Contact by wild mice with colony mice was the likely source for LCMV, and shipments of infected mice among facilities spread the infection. The breeding colonies were depopulated to prevent further human infections. Future outbreaks can be prevented with monitoring and management, and employees should be made aware of LCMV risks and prevention.

Author affiliations: Centers for Disease Control and Prevention, Atlanta, Georgia, USA (B. Knust, U. Ströher, L. Edison, C.G. Albarino, L. Guerrero, S. Campbell, A. Gibbons, Z. Reed, D. Cannon, C. Manning, B. Petersen, S.T. Nichol, P.E. Rollin); Georgia Department of Public Health, Atlanta (L. Edison); Indiana Board of Animal Health, Indianapolis, Indiana, USA (J. Lovejoy, D. Metcalf, B. Marsh); Deaconess Hospital, Evansville, Indiana, USA (E. Armeanu); Indiana State Department of Health, Indianapolis (J. House); Vanderburgh County Health Department, Evansville (D. Cory); Green River District Health Department, Owensboro, Kentucky, USA (C. Horton, J. Austin); and Kentucky Department for Public Health, Frankfort, Kentucky, USA (K.L. Fowler, J. Poe, K.E. Humbaugh)

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from patient 1 were positive for LCMV IgM by ELISA, indicating recent LCMV infection. Reverse transcription PCR (RT-PCR) for LCMV was negative, indicating that viremia was no longer present.

The Vanderburgh (Indiana) County Health Department, in conjunction with the Indiana State Department of Health, Indiana Board of Animal Health, and CDC, initiated an outbreak investigation to determine the extent of LCMV infection in the staff and rodents in facility A (4). Trace-back investigations also identified a distributor (facility B) where live rats (Rattus norvegicus) and mice (M. musculus) from facility A were handled and packaged for sale as live and frozen animal food in 21 states (5). An additional mouse breeding facility in Kentucky (facility C) had shipped live mice to facility B, which redistributed them to facility A shortly before the outbreak. We describe the diagnostic and epidemiologic aspects of this investigation and the response taken to control the outbreak at these facilities.

Materials and Methods

Employee Serosurvey

As part of the investigation, we provided all current and former employees of facilities A, B, and C within the previous 6 months the opportunity to be interviewed and tested for LCMV as part of the investigation. County health officials administered a questionnaire that collected information about recent clinical illness and work habits at the facilities, including hygiene measures. A blood sample was collected from each interviewee and refrigerated for transport to CDC, where IgG and IgM ELISA were performed (online Technical Appendix, wwwnc.cdc.gov/eid/article/20/2/pdfs/13-0860.pdf). We considered a current or former employee to be recently infected (within the past 2–3 months) with LCMV if he or she had IgM with or without IgG. An employee for whom only IgG was detectable by ELISA was considered to have had a previous infection. If employee had evidence of recent LCMV infection and had sought medical care because of illness, we reviewed his or her medical records. All employees signed an informed consent form.

Rodent Testing

Adult breeding rodents (mice and rats) from facility A were sampled to determine LCMV infection status for each room (online Technical Appendix). In accordance with the facility’s standard operating procedures for processing feeder rodents, we euthanized mice and rats with carbon dioxide gas, and the carcasses were frozen and shipped to CDC on dry ice. Carcasses were thawed, and animals were dissected under Biosafety Level 3 conditions. Heart blood and small sections of kidney and spleen were collected. Rodent specimens were tested for LCMV RNA by RT-PCR and for LCMV IgG by ELISA. RT-PCR–positive specimens were inoculated onto cell culture to isolate virus, and viral RNA was sequenced and compared with other LCMV strains (online Technical Appendix).

Statistical Analysis

All data were collected by using a computerized spreadsheet (Microsoft Excel, Microsoft, Redmond, WA, USA) and analyzed by using statistical analysis software (SAS 9.3, SAS Institute, Cary, NC, USA). Rodent location information was combined with diagnostic test results to calculate an observed prevalence of IgG seropositivity and RT-PCR positivity per room sample. Ninety-five percent CIs for the prevalence of mice with LCMV antibodies and apparent viremia per room population were estimated by using a binomial distribution equation, and estimated ranges of antibody-positive and viremic rodents per room were calculated from these ranges.

Employee questionnaire data were combined with diagnostic test results, and a case–control comparison was performed. We defined an LCMV case-patient as an employee who had detectable LCMV IgM and/or IgG and controls as employees of the same facility who had negative test results. Statistics for each facility were examined separately. Univariate analysis was conducted, and p<0.05 was considered statistically significant.

Results

Human LCMV Investigation

All 52 employees of facility A were tested by ELISA. Fifteen (29%) had detectable anti-LCMV antibodies; 13 had IgM and IgG, indicating recent infection. Nine IgM/IgG-positive employees reported recent clinical illness, including 5 who had sought medical treatment. Symptoms described most frequently were headache, nausea and vomiting, subjective fever, decreased appetite, diarrhea, muscle ache, and stiff neck (Table 1). Aseptic meningitis was diagnosed in 4 employees from facility A after lumbar punctures demonstrated lymphocytic CSF (Table 2). Dates of onset for laboratory-confirmed aseptic meningitis ranged from April 7 through May 14, 2012. All case-patients recovered fully.

All 13 employees of facility B were tested; 1 (8%) had evidence of previous infection (LCMV IgG only). This 38-year-old woman did not recall any recent distinct clinical illness that fit the clinical profile for LCMV infection. She had not directly handled any live mice but had handled and labeled shipping packages of live mice from facilities A and C in the months before being tested.

Thirty-two of 36 facility C employees were tested; 15 (47%) had detectable LCMV antibodies, and 11 (34%) had evidence of recent LCMV infection. In facility C, 1 of the
tested employees was pregnant and negative for LCMV antibodies, and a 29-year-old man who was IgG/IgM positive had visited the emergency department in May 2012 because of chest pain and headaches. Electrocardiogram, serum chemistry, and complete blood count did not show any abnormalities. Altogether, 97 employees of facilities A, B, and C were tested, and 31 (32%) had LCMV antibodies. Job duties of employees at facilities A, B, and C ranged from administrative/managerial duties to direct handling of live and euthanized rodents to cleaning. Most employees conducted multiple duties in multiple buildings. No particular job duty was associated with LCMV infection (data not shown). Employees had worked at the facilities for 2 days–20 years, and 1 case-patient had worked at facility A for only 5 days before becoming ill. We found no association between length of time employed and previous infection. For employees had worked at the facilities for 2 days–20 years, and 1 case-patient had worked at facility A for only 5 days before becoming ill. We found no association between length of time employed and previous infection.

For facility A, working in building 2 and smoking were independently associated with having recent or previous LCMV infection (Table 3, Appendix). For facilities B and C, no specific factors evaluated were associated with LCMV infection in employees tested (Table 3, Appendix). At the time of the investigation, all 97 employees reported washing hands after handling the rodents, and 90%–100% of employees reported wearing masks and gloves, although many admitted having begun using these items only when the LCMV outbreak was suspected.

### Rodent LCMV Investigation

**Facility A**

Facility A, located in Indiana, bred and raised mice and rats exclusively for sale as feeder animals for reptiles and birds of prey. Most rodents were euthanized and frozen on-site for sale as frozen feeder rodents; however, live rodents also were shipped. Live and frozen rodents were transported to facility B, also in Indiana, for storage, sale, and distribution. Facility A had 4 buildings that housed breeding rodents; each building was subdivided into rooms by species (Table 3, Appendix). At the time of the investigation, facility A housed ≈155,000 adult mice and ≈14,000 adult rats. In accordance with standard procedures, baited rodent traps were set outside and inside buildings at regular intervals throughout the facility; any mice that were caught were promptly euthanized. Rodent feed was stored indoors on pallets.

In May 2012, a total of 1,820 mice and rats from facility A were tested for LCMV IgG by ELISA and for LCMV RNA by RT-PCR. Of 1,421 mice tested, 296 (20.8%) had detectable IgG, and 10 (0.7%) had detectable RNA; apparent prevalence varied by room (Table 4). No RT-PCR–positive mice had detectable LCMV IgG. Only 1 mouse room tested (building 3, room 2) contained no IgG- or RT-PCR–positive mice, indicating that LCMV infection was widespread. The estimated number of viremic mice in a

### Table 1. Symptoms reported by 9 employees of facility A who tested positive for LCMV IgM and reported recent illness, Indiana, USA, 2012

<table>
<thead>
<tr>
<th>Symptom or sign</th>
<th>Present no. (%)</th>
<th>Absent no. (%)</th>
<th>Unknown no.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Headache</td>
<td>9 (100)</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Nausea/vomiting</td>
<td>9 (100)</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Fever</td>
<td>8 (89)</td>
<td>1 (11)</td>
<td>0</td>
</tr>
<tr>
<td>Decreased appetite</td>
<td>8 (89)</td>
<td>1 (11)</td>
<td>0</td>
</tr>
<tr>
<td>Diarrhea</td>
<td>6 (67)</td>
<td>3 (33)</td>
<td>0</td>
</tr>
<tr>
<td>Muscle ache</td>
<td>6 (67)</td>
<td>3 (33)</td>
<td>0</td>
</tr>
<tr>
<td>Stiff neck</td>
<td>6 (67)</td>
<td>3 (33)</td>
<td>0</td>
</tr>
<tr>
<td>Joint pain</td>
<td>5 (56)</td>
<td>4 (44)</td>
<td>0</td>
</tr>
<tr>
<td>Malaise</td>
<td>5 (56)</td>
<td>2 (22)</td>
<td>2 (22)</td>
</tr>
<tr>
<td>Cough</td>
<td>4 (44)</td>
<td>5 (56)</td>
<td>0</td>
</tr>
<tr>
<td>Drowsiness</td>
<td>3 (33)</td>
<td>6 (67)</td>
<td>0</td>
</tr>
<tr>
<td>Sensory disturbance</td>
<td>2 (22)</td>
<td>6 (67)</td>
<td>1 (11)</td>
</tr>
<tr>
<td>Parotid pain</td>
<td>1 (11)</td>
<td>7 (78)</td>
<td>1 (11)</td>
</tr>
<tr>
<td>Confusion</td>
<td>1 (11)</td>
<td>8 (89)</td>
<td>0</td>
</tr>
<tr>
<td>Increased blood leukocytes</td>
<td>3 (33)</td>
<td>1 (11)</td>
<td>5 (56)</td>
</tr>
<tr>
<td>Increased CSF leukocytes</td>
<td>4 (44)</td>
<td>0</td>
<td>5 (56)</td>
</tr>
<tr>
<td>Increased CSF protein</td>
<td>4 (44)</td>
<td>0</td>
<td>5 (56)</td>
</tr>
</tbody>
</table>

*LCMV, lymphocytic choriomeningitis virus; CSF, cerebrospinal fluid.*

### Table 2. Demographic and clinical features of laboratory-confirmed illness in 5 facility A employees who sought health care after exposure to LCMV-infected mice, Indiana, USA, 2012

<table>
<thead>
<tr>
<th>Patient no.</th>
<th>Age, y/sex</th>
<th>Time employed</th>
<th>Onset date</th>
<th>Symptoms</th>
<th>Clinical findings†</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>27/F</td>
<td>2 y</td>
<td>Apr 20</td>
<td>Fever, muscle ache, nausea, vomiting, abdominal pain, diarrhea, malaise, headache, stiff neck</td>
<td>Lymphocytic CSF, leukocytosis (13.8), elevated protein in CSF, appendicitis, UTI</td>
</tr>
<tr>
<td>2</td>
<td>34/M</td>
<td>1.5 y</td>
<td>Apr 11</td>
<td>Fever, nausea, vomiting, malaise, headache, stiff neck</td>
<td>Lymphocytic CSF (1,311/μL), elevated protein in CSF</td>
</tr>
<tr>
<td>3</td>
<td>28/F</td>
<td>1.5 y</td>
<td>Apr 7</td>
<td>Fever, muscle ache, nausea, vomiting, cough, joint pain, photophobia, headache, diarrhea</td>
<td>UTI; no blood or CSF collected during medical evaluation</td>
</tr>
<tr>
<td>4</td>
<td>48/M</td>
<td>5 d</td>
<td>Apr 27</td>
<td>Fever, muscle ache, nausea, vomiting, abdominal pain, cough, headache, diarrhea</td>
<td>Lymphocytic CSF, elevated protein in CSF, mild anemia (34.9%), CT normal</td>
</tr>
<tr>
<td>5</td>
<td>38/M</td>
<td>2 mo</td>
<td>May 14</td>
<td>Fever, muscle ache, nausea, vomiting, abdominal pain, cough, headache, diarrhea</td>
<td>Lymphocytic CSF (320/μL), elevated protein in CSF (77), leukocytosis (15.1), anemia (Hgb 11.5), CT normal</td>
</tr>
</tbody>
</table>

*LCMV, lymphocytic choriomeningitis virus; CSF, cerebrospinal fluid; UTI, urinary tract infection; CT, computed tomographic scan; Hgb, hemoglobin.†Lymphocytic CSF is defined as >5 lymphocytes/μL; leukocytosis is >11,000 erythrocytes/μL; and elevated protein in CSF is >60 mg/dL.*

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strain 201202467 belongs to lineage I, a heavily popula
tLCMV. The phylogenetic analysis (Figure) shows that
generated GenBank, corresponding to representative strains of
the 34 sequences available
shown), suggesting a single origin of LCMV. The com-
PCR-positive mice. The sequence analysis of a 630-bp
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given room at the time of sampling varied from 0 to 472 (as
estimated by the number of RT-PCR–positive mice) (95% CI 0–1,180 viremic mice per room) (Table 4). None of 399 rats tested were positive by ELISA or RT-PCR.

LCMV was successfully isolated from 8 of 10 RT-
PCR–positive mice. The sequence analysis of a 630-bp amplicon from the S-segment from the 8 isolates showed a high degree of similarity (98.6%–99.7% identity; data not shown), suggesting a single origin of LCMV. The complete S-segment sequence of the prototype outbreak strain (201202467 Indiana, GenBank accession no. KF732824) was aligned and compared with 34 sequences available in GenBank, corresponding to representative strains of LCMV. The phylogenetic analysis (Figure) shows that strain 201202467 belongs to lineage I, a heavily populat-
ed LCMV clade with numerous isolates from Europe and North America (6).

After evidence of LCMV infection was detected, the Indiana Board of Animal Health (Indianapolis, IN, USA) issued quarantines and stop-movement orders on all live and frozen rodents from facility A. After animal testing was done, regulatory restrictions were lifted on rats but were retained on mice. More than 400,000 mice were euthanized and buried on site. All rodent feed and bedding was incinerated, and the facility was disinfected by using a 0.1% bleach solution. Employees were provided N95 respirators, gloves, boots, and water-resistant coveralls to wear when handling the mice and possibly contaminated equipment, and they were instructed about how to properly use the equipment and disinfect appropriately to limit their risk for infection. Once these measures were fully imple-
mentation, no additional employees became ill throughout the depopulation process, which ended on July 11, 2012.

All frozen rodents from facility A that were remaining in storage at facility B were destroyed. Live rodents originating from facility A were sold to >500 different purchas-
erS in 21 different states (5).

### Facility C

Facility C exclusively produced frozen feeder mice that were euthanized on site in Kentucky and then transported to facility B for storage and distribution. In March 2012, ≈90,000 live mice were shipped from facility C to facility A via facility B to replace breeding stock. Other than this instance, live shipments were not common. Management was similar to that of facility A. After the LCMV outbreak among facility A employees, the owners of facility C had 50 mice tested at a commercial laboratory (IDEXX RADIL, Columbia, MO, USA); 33 (66%) were antibody-positive by immunofluorescent antibody testing. No mice were tested by CDC. The owners of facility C reported this finding to the Indiana Board of Animal Health (Indianapolis, IN, USA); 33 (66%) were antibody-positive by immunofluorescent antibody testing. No mice were tested by CDC. The owners of facility C reported this finding to

### Table 4. Test results for LCMV from mice and rats in facility A, Indiana, USA, 2012*

<table>
<thead>
<tr>
<th>Building/room</th>
<th>Species</th>
<th>No. adult rodents in room</th>
<th>Sample size</th>
<th>IgG positive</th>
<th>Estimated no. per room (95% CI)†</th>
<th>RT-PCR positive</th>
<th>Estimated no. per room (95% CI) †</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Rat</td>
<td>1,512</td>
<td>110</td>
<td>0</td>
<td>0 (0–41)</td>
<td>0</td>
<td>0 (0–41)</td>
</tr>
<tr>
<td>2</td>
<td>Rat</td>
<td>2,058</td>
<td>83</td>
<td>0</td>
<td>0 (0–72)</td>
<td>0</td>
<td>0 (0–72)</td>
</tr>
<tr>
<td>3</td>
<td>Mouse</td>
<td>13,104</td>
<td>110</td>
<td>28 (25)</td>
<td>3,342 (2,306–4,534)</td>
<td>4 (4)</td>
<td>472 (131–1,179)</td>
</tr>
<tr>
<td>4</td>
<td>Mouse</td>
<td>10,368</td>
<td>110</td>
<td>25 (23)</td>
<td>2,354 (1,586–3,287)</td>
<td>2 (2)</td>
<td>187 (21–664)</td>
</tr>
<tr>
<td>5</td>
<td>Mouse</td>
<td>13,104</td>
<td>102</td>
<td>36 (35)</td>
<td>4,626 (3,420–5,949)</td>
<td>0</td>
<td>0 (0–380)</td>
</tr>
<tr>
<td>6</td>
<td>Mouse</td>
<td>10,368</td>
<td>110</td>
<td>42 (38)</td>
<td>3,961 (3,017–4,966)</td>
<td>1 (1)</td>
<td>93 (2–508)</td>
</tr>
</tbody>
</table>

*LCMV, lymphocytic choriomeningitis virus; RT-PCR, reverse transcription PCR.
†Binomial estimators were used to calculate 95% CIs for the entire adult population for each room, based on the number of test-positive animals detected with the samples obtained.
‡All RT-PCR–positive animals were LCMV IgG-negative as measured by ELISA.
RESEARCH

with black eyes (colony mice are albino), indicating that wild mice had interbred with the colony mice.

Discussion

LCMV is endemic in wild *M. musculus* populations across the United States and throughout the world. Sporadic human LCMV cases occur after contact with infected wild house mice, but the virus has the potential to cause large epidemics when it enters high-density rodent populations, as in the outbreak of this report. Previous outbreaks of human disease have been linked to contact with pet hamsters and laboratory animals (7–11). In the current outbreak investigation, we found that nearly one third of rodent facility employees tested had LCMV antibodies. This overall attack rate is consistent with previous outbreaks of LCMV in hamsters and nude mice at research facilities (10,12,13).

Employees of rodent breeding facilities of all kinds should be aware of the risks posed by exposure to rodents infected with LCMV, and monitoring programs should be in place to detect and control infections in rodents. Commercial laboratory rodent breeding colonies have developed management practices to avoid contact between wild mice and colony animals; the US Department of Health and Human Services and the Federation of European Laboratory Animal Science Associations recommend routine virologic and serologic monitoring to detect pathogens, including LCMV (2,14). Facilities producing rodents for the pet and feeder-rodent industries should adapt these

Figure. Phylogenetic tree comparing S RNA genomes of most representative lymphocytic choriomeningitis virus (LCMV) strains. Evolutionary analysis was conducted in MEGA5 (online Technical Appendix, wwwnc.cdc.gov/eid/article/20/2/pdfs/13-0860.pdf) by using the neighbor-joining method. Bootstrap values listed at the nodes provide statistical support for 1,000 replicates. Branches corresponding to partitions reproduced in <50% bootstrap replicates are collapsed. Scale bar indicates substitutions per site. The main LCMV lineages are indicated with roman numerals (I–IV). After the strain denomination, the location and date it was isolated are noted. A color version of this figure is available online (www.cdc.gov/EID/content/20/2/130860-F.htm)
practices to avoid such outbreaks. When LCMV or LCMV antibodies are detected in rodents or employees of a rodent breeding operation, all animal-handling personnel should wear protective equipment, including a respirator. After personal protective gear and training in its proper use were provided to facility employees, LCMV infection was not laboratory confirmed in any additional employees of facility A throughout the depopulation process, which implies that use of such equipment can reduce the risk for infection.

In facility A, many mice had LCMV antibodies, and several were viremic at the time of the investigation. The rodent testing results fit with the human testing results; that is, working in building 2, which had the highest prevalence of LCMV antibody-positive and RT-PCR–positive mice, was significantly associated with employee infections. Of the ≈13,000 adult mice in room 1 of building 2, ∼131–1,179 were viremic. In such a high-density environment, the virus can be present in aerosol form (15), which explains why many employees of facilities A and C had detectable antibodies. In facility B, which only transiently held live mice for further distribution, only 1 (8%) of 13 employees had detectable LCMV antibodies, suggesting that less intensive exposure to mice put employees at lower risk for infection. Smoking, a risk factor for many bacterial and viral infections, including tuberculosis, pneumococcal pneumonia, and influenza (16), was associated with LCMV infection in facility A employees. The structural changes to the respiratory epithelium and modulation of immune function hypothetically impair the smoker’s immune response. The physical act of smoking also might facilitate transfer of pathogens from the hands to the mouth.

The proportion of infected employees in whom clinical illness developed varied by facility. In facility A, 69% of employees who had recent infection reported illness, and 33% had symptoms severe enough to cause them to seek medical care. Conversely, all facility C employees appeared to have asymptomatic infections, except for 1 with nonspecific illness. Previous LCMV outbreak investigations have found various rates of clinical illness (9,10,13), with asymptomatic infections of 25%–55%. Because employee age and sex did not differ among facilities, the reason for the difference in disease manifestation remains unclear. The longevity of LCMV IgM detectable in peripheral blood is not well-established for humans; thus, some employees with IgM who did not develop symptoms might have been infected previously.

In immunocompetent adults, the neurologic form of LCMV infection classically has biphasic features consisting of a nonspecific initial phase, with fever, myalgia, and headache most commonly observed, and can include nausea and/or vomiting and retroorbital pain (8,17). Symptoms may subside after several days to be followed by a neurologic phase comprising meningeal symptoms with fever, headache, nuchal rigidity, vomiting, and light sensitivity. In LCMV patients in whom aseptic meningitis is diagnosed, CSF characteristically shows a lymphocytic pleocytosis, and elevated protein and decreased glucose also might be present (1,3); in this report, all 4 case-patients on whom lumbar punctures were performed had lymphocytic CSF with elevated protein. Symptoms frequently reported by case-patients in this outbreak included headache; fever; and abdominal symptoms, such as nausea, vomiting, and diarrhea. Such nonspecific symptoms obscure the clinical diagnosis. A thorough clinical history covering relevant animal contacts remains vital to determining the source and appropriate etiologies to test for in work-ups of aseptic meningitis.

Although all case-patients in this LCMV outbreak recovered from their illness, the specter of more severe disease manifestations remains a cause for concern. Infections during pregnancy can result in spontaneous abortion or characteristic congenital defects, such as chorioretinitis, microencephaly or macroencephaly, and hydrocephalus (18). Mental retardation, vision deficits, cerebral palsy, and epilepsy are potential lifelong manifestations (19). Since 2005, 5 outbreaks of LCMV have occurred after organ transplantation, resulting in the death of 14 (82%) of 17 organ recipients (20–23). Concern about these severe forms of LCMV infection led the outbreak response team to recommend depopulating the mouse breeding facilities and to conduct a trace-forward investigation of live mice that were sold and shipped from facility B (5).

LCMV is transmitted horizontally and vertically in affected rodents (17,24). Horizontal infections, acquired through direct contact with infected rodents or indirect contact with contaminated fomites, can lead rodents to shed infectious virus for a few weeks to a few months. When mice are exposed in utero to LCMV, they become persistently infected and shed the virus throughout their lives, including to all offspring, which also will be persistently infected (2,25,26). Infections in rodents are inapparent. As a result, when LCMV is introduced to a high-density environment, such as the breeding colonies of this outbreak, the number of infected rodents can silently reach very high numbers and thus pose a risk to the humans coming in contact with them.

The primary goal of the rodent sampling scheme was to detect LCMV infection in the colony; only adult mice were sampled because we assumed that they would have the greatest likelihood of having detectable antibodies. If younger mice had been tested, the seroprevalence is likely to have been lower and the proportion of viremic animals higher. In this investigation, 21% of adult mice tested had antibodies; this prevalence is higher than that found in wild mouse populations not associated with human infections (2%–9%) (27–29).
These molecular investigations demonstrated a unique strain of LCMV, 201202467, in facility A mice, which suggests a single introduction and transmission event throughout the breeding facility. Although we were unable to sample mice from facility C, it is likely that the outbreak strain was the same as in facility A and was introduced during the wild mouse infestation that had occurred. Molecular analysis also shows the close relationship of strain 201202467 with another LCMV strain isolated from a mouse found in an infested house in Michigan in 2005 (30). The geographic distribution of this LCMV strain throughout North America is not known (7).

As in previous outbreaks, interactions between wild mice and colony rodents frequently introduce LCMV into breeding colonies (7). In the absence of control measures and monitoring, movements of live mice among breeder mice can contribute to the spread of potentially infected mice. No human or rodent LCMV vaccine is available, and no treatment exists. Therefore, prevention measures are necessary and must rely on wild rodent exclusion, infection control, and microbiological monitoring (2). When LCMV antibodies are detected in colony mice, transmission must be assumed to be ongoing, and all possibly infected or exposed rodents should be removed from the colony by euthanasia and disposal to mitigate human risk (24,31).

In conclusion, laboratory and epidemiologic investigations effectively identified a large outbreak of LCMV in 3 commercial mouse breeding facilities and associated infections in several employees. The presumptive source of virus introduction was contact between wild mice and colony mice, and the outbreak spread among facilities when mice were transported for use as breeding stock. The breeding colonies were depopulated to prevent further human infections. Future outbreaks can be prevented with strict biosecurity and microbiological monitoring, and employees should be made aware of the symptoms of LCMV infection and prevention measures.

Acknowledgments

We gratefully acknowledge Gayle Galland, Adam Langer, and Christian Witkovskie for their assistance in accessing and inspecting facility C.

Dr Knust is an epidemiologist and veterinarian in CDC’s Viral Special Pathogens Branch. Her primary research interests include epidemiology and prevention of zoonotic diseases.

References


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Technical Appendix

Detailed Animal Sampling and Laboratory Methods

Facility A Rodent Sampling

In facility A, numbers of adult breeding rodents in each room were estimated. Each room was treated as a discrete population. Hypergeometric probability equations were used to determine sample size, similar to a previous lymphocytic choriomeningitis virus (LCMV) investigation (1). Given the estimated range of adult rodents per room (1,500–15,000), a sample size of 110 animals was estimated to have a 96.5%–96.9% probability of detecting at least 1 antibody-positive animal if the overall seroprevalence was ≥3%. Systematic sampling was used to select at least 1 adult animal from each rack in the room, with varying pan locations within the rack. Juvenile animals were not sampled.

Diagnostic Assays

ELISA

Human serum was tested for LCMV IgM and IgG by using in-house ELISA as described (2). Mouse and rat serum was tested with a modified LCMV IgG ELISA by using goat antimouse conjugate (Cat 31446, Pierce, Rockford, IL, USA) or antirat conjugate (Cat 14-16-06, Kirkegaard & Perry Laboratories, Gaithersburg, MD, USA).

RT-PCR and Virus Genome RNA Sequencing

Tissue specimens of mouse liver and spleen (≈100 mg) were collected in 500 μL of lysis binding solution (Ambion Life Technologies, Austin, TX, USA) and homogenized in a high-throughput tissue grinder (Geno/Grinder 2000, SPEX SamplePrep, Metuchen, NJ, USA). Total RNA was extracted with MagMax-96 total RNA isolation kit (SuperScrip III One-Step RT-PCR System with Platinum Taq High Fidelity, Invitrogen Life Technologies, Grand Island, NY, USA),
following the manufacturer’s instructions. A 655-bp product corresponding to the nucleoprotein gene was amplified by using a generic primer set capable of amplifying all known strains of LCMV. The primers used in these reactions were as follows: LCMV1748F (AIATIATRCARTCCATRAGIGCRCA) and LCM2377R (TCIGGIGARGGITGGCCITAYAT). The RT-PCR was done as described (1) with SuperScript III One-Step RT-PCR System with Platinum Taq High Fidelity (Invitrogen Life Technologies), according to the manufacturer instructions. The amplified nucleoprotein gene fragments from 8 positive samples were sequenced by using standard dideoxy-sequencing techniques (Invitrogen Life Technologies). To further characterize the LCMV strain, the complete S RNA (∼3,411 nt) from 1 representative sample (201202467) was amplified by RT-PCR, as described (3), by using the arenavirus standard primer Are19c (CGCACAGTGGATCCTAGGC). Multiple sequence alignments and phylogenetic analysis were done using MEGA4 (4).

Virus Isolation

For all mice found to have virus RNA detected by RT-PCR, 100 mg of mixed spleen and liver tissues were ground manually in Hank’s Balanced Salt Solution and centrifuged. 100 μL of the resulting tissue homogenate suspension was inoculated on confluent Vero E6 cells. Flasks were incubated at 37°C and monitored for cytopathic effects. On day 7, culture medium was changed and cells checked by immunofluorescence assay (IFA) for the presence of LCMV. Negative cultures were kept until day 14 for a final IFA before being discarded. Positive cultures were harvested and stored.

References

