Surveillance for Unexplained Deaths and Critical Illnesses Due to Possibly Infectious Causes, United States, 1995–1998


Population-based surveillance for unexplained death and critical illness possibly due to infectious causes (UNEX) was conducted in four U.S. Emerging Infections Program sites (population 7.7 million) from May 1, 1995, to December 31, 1998, to define the incidence, epidemiologic features, and etiology of this syndrome. A case was defined as death or critical illness in a hospitalized, previously healthy person, 1 to 49 years of age, with infection hallmarks but no cause identified after routine testing. A total of 137 cases were identified (incidence rate 0.5 per 100,000 per year). Patients’ median age was 20 years, 72 (53%) were female, 112 (82%) were white, and 41 (30%) died. The most common clinical presentations were neurologic (29%), respiratory (27%), and cardiac (21%). Infectious causes were identified for 34 cases (28% of the 122 cases with clinical specimens); 23 (68%) were diagnosed by reference serologic tests, and 11 (32%) by polymerase chain reaction-based methods. The UNEX network model would improve U.S. diagnostic capacities and preparedness for emerging infections.

The 1992 Institute of Medicine report—Emerging Infections, Microbial Threats to Health in the United States (1)—highlighted the need for a more effective means to detect emerging infectious diseases. In response to this report and as part of the Emerging Infections Program (EIP) (2), the Centers for Disease Control and Prevention (CDC) collaborated with state health departments and academic institutions to develop a pilot surveillance strategy for early detection of new and unrecognized infectious diseases in the United States. This project—Surveillance for Unexplained Deaths and Critical Illnesses Due to Possibly Infectious Causes—was developed on the basis of two observations. The first was the realization that supposedly new infectious diseases identified in the United States in recent decades had been occurring long before they were recognized and identified. The second was important progress in molecular diagnostic methods, which in some instances has allowed new infectious agents to be identified and characterized with molecular probes, making in vitro cultivation unnecessary.

In 1995, we initiated population-based surveillance for unexplained deaths and critical illnesses due to possibly infectious etiologies (UNEX) at four U.S. sites. The objectives of this effort were to define the incidence, epidemiologic features, and possible causes of these deaths and illnesses; create a bank of clinical specimens for future testing as new pathogens and methods are identified; and assist in building U.S. capacity for detecting and responding to uncommon and previously unrecognized pathogens. This report describes the methods we developed to reach these goals and the results of the first 3.5 years of surveillance.

Methods

Surveillance Sites

Population-based surveillance for UNEX was initiated on May 1, 1995, among persons 1 to 49 years of age residing in the San Francisco Bay area (Alameda, Contra Costa, and San Francisco Counties) of California (n=2,168,810); in New Haven County, Connecticut (n=556,592); in the entire state of Minnesota (n=3,419,760); and among persons 1 to 39 years of age residing in Oregon (n=1,544,466).2 All these sites were participants in the EIP, and the total population targeted for


2Oregon used a different age cut-off because of limited resources.
surveillance was 7.7 million. We report the results of surveil-
ance for cumulative cases through December 31, 1998.

Case Definition
An UNEX case was defined as illness in a previously
healthy resident of a surveillance area who was 1 to 49 years
old (1 to 39 years old in Oregon) and who died or was hospi-
talized with a life-threatening illness with hallmarks of an
infectious disease for which no cause was identified through
routine testing initiated by health-care providers. A previously
healthy person was defined as a patient without a preexisting
known systemic, chronic medical illness diagnosed before the
acute onset of the UNEX. Such preexisting conditions
included malignancy; HIV infection; chronic cardiac, pulmo-

dary, renal, hepatic, or rheumatologic disease; or diabetes mel-

litus. Patients were also excluded from the study if they had
received any immunosuppressive therapy, had evidence of
toxic ingestion or exposure, had trauma before their illness, or
acquired their illness ≥48 hours after hospital admission
(Appendix I).

A life-threatening illness was defined as any illness requir-
ing admission to an intensive-care unit (ICU). Hallmarks of an
infectious disease were defined as the following: fever or his-
tory of fever, leukocytosis, histopathologic evidence of an
acute infectious process, or a physician-diagnosed syndrome
consistent with an infectious etiology, including encephalitis
or meningitis, fulminant hepatitis or hepatic failure, myocarditis,
adult respiratory distress syndrome, respiratory failure, or sepsis.

Case Finding and Ascertainment
Patients meeting the case definition were sought at surveil-
ance sites through various mechanisms. Practicing clinicians
in all surveillance sites were informed about the project
through letters and bulletins and presentations at local and
regional professional society meetings. Personnel at some sur-
veillance sites attempted to identify cases more actively
through regular communications with persons working in
ICUs and local medical examiners or through routine review
of ICU admission records. Physicians and other health profes-
sionals were asked to report suspected cases by telephone to
local surveillance site personnel. When a case was reported, a
screening form was completed to determine if the patient met
the case definition. This surveillance system was not designed
to provide timely reporting or testing.

Surveillance Audit
To evaluate the sensitivity of the surveillance system, per-
sonnel at all surveillance sites conducted a retrospective
review of death records from their surveillance areas, and three
sites (California, Connecticut, and Oregon) reviewed all hospi-
tal discharge data in their areas for a period of at least 1 year.
All death certificates for the age groups included in the sur-
veillance were reviewed for the presence of specific Interna-
tional Classification of Disease codes (ICD-9), selected for
their potential to identify unexplained deaths due to possibly
infectious causes (3). Persons whose death records included
ICD-9 codes indicating a disqualifying underlying medical
condition were excluded. Once potential cases were identified,
the patients’ medical records were reviewed. If the records
were not available, the primary physician was contacted to
determine if the patient met the surveillance case definition.
The sensitivity of the surveillance system for detecting deaths
(SD) was calculated by dividing the number of deaths (D1)
detected through surveillance alone by the total number of
deaths (D1+D2) detected through both surveillance and death
record review (D2): SD=D1/D1+D2. The sensitivity of the sur-
veillance system (SC) for detecting critical illness cases was
calculated by dividing the number of such cases (C1) detected
through surveillance alone by the total number of cases
(C1+C2) found through both surveillance and hospital dis-
charge review (C2): SC=C1/C1+C2.

Collection of Clinical Information and Specimens
For patients meeting the case definition, surveillance site
personnel completed a case report form that included demo-
graphic, epidemiologic, and clinical information. This inform-
ation was collected through interview of physicians caring
for the patient, review of the medical record, and contact with
the patient or the patient’s family. Cases were assigned a clin-
ical syndrome depending on the predominant system involved,
the basis of information provided by the physician. These
syndromes included neurologic (encephalitis, meningitis), car-
diac (myocarditis, pericarditis, endocarditis), respiratory
(pneumonitis), and hepatic (hepatitis). Syndromes such as sep-
sis, in which no predominant organ system was involved, were
classified as “other.” The hospital laboratories were requested
to save all remaining clinical specimens obtained as part of
routine clinical management, including biopsies and autopsies.

Laboratory Testing
For the first 2 years of the study, the project investigators
selected diagnostic tests individually for each case. Decisions
were made on the basis of clinical, epidemiologic, and histo-
logic data; previous laboratory testing ordered by the health-
care providers; and availability, timing, quality, and quantity of
clinical specimens. In the third year of the project, based on
information gained to date, a set of standardized syndrome-
specific laboratory testing protocols was developed for respir-
atory, neurologic, cardiac, and hepatic syndromes (Appendix
II: available online at URL: http://www.cdc.gov/ncidod/EID/ vol8no2/pdf/01-0165-app2.pdf). These protocols prioritized
testing based on available clinical and epidemiologic informa-
tion and a differential diagnosis; they guided a first round of
laboratory testing which, if negative, prompted a customized
second round of testing. Cases that did not fit any of these four
syndromes were discussed by the project investigators on an
individual basis.
Histopathologic Testing

Whenever possible, in addition to initial examination by local pathologists, tissue specimens were examined by CDC pathologists to help guide further laboratory testing decisions. CDC pathologists have available a unique set of antibodies and probes for immunohistochemistry (IHC) and in-situ hybridization (ISH); these and other special studies, such as chemical stains, were selected based on all available case information. IHC tests were performed by a two-step indirect immunoperoxidase technique with various antibodies (4). ISH tests used digoxigenin-labeled probes with an immunoperoxidase staining protocol (5). Positive and negative controls were run in parallel with case specimens.

Testing for Viral Pathogens

The California Department of Health Services (CDHS) Viral and Rickettsial Diseases Laboratory was the primary testing site for viral pathogens other than the hepatitis viruses. Serologic tests were available for immunoglobulin (Ig) G directed against 19 viral pathogens and for IgM directed against 14 of these. When only a single serum specimen was available, the presence of both IgM and IgG was assessed, either by enzyme immunosorbent assay (EIA), indirect immunofluorescence assay (IFA), or both (6). Paired sera were tested by EIA or IFA for increase in IgG titer. Additional testing included nucleic acid amplification by polymerase chain reaction (PCR) for selected viral pathogens if adequate specimens were available (7-9). An increase in IgG titer by EIA was interpreted as evidence of recent or current infection if the ratio of convalescent- to acute-phase indices was ≥1.5; an index is determined by the equation (optical density [OD]-positive antigen - OD-negative antigen)/predetermined positive threshold OD (usually 0.1). The CDHS diagnostic assays for IgM to B19V, Cytomegalovirus, Hantavirus (SNV), herpes simplex virus, MeV, MuV, RUBV, SLEV, VZV, and WEEV have varying minimum positive values, with indices from 1.0 and 2.0. For the enterovirus IgM assay, which detects the presence of enterovirus group antibody in serum, a ratio of OD-negative antigen to OD-positive antigen ≥2.0. For the enterovirus IgM assay, which detects the presence of enterovirus group antibody in serum, a ratio of OD-negative antigen to OD-positive antigen ≥2.0. For the enterovirus IgM assay, which detects the presence of enterovirus group antibody in serum, a ratio of OD-negative antigen to OD-positive antigen ≥2.0. For the enterovirus IgM assay, which detects the presence of enterovirus group antibody in serum, a ratio of OD-negative antigen to OD-positive antigen ≥2.0.

Bacterial Broad-Range Ribosomal DNA (rDNA) PCR

DNA extraction from clinical specimens was performed as described (10,11). All clinical specimens tested with the broad-range bacterial rDNA PCR were analyzed by using at least one of the following primer pairs: fD1mod (positions 8-27 in Escherichia coli 16S rRNA gene) (12) and 16S1RR-B (575-556) (13); 8F2 (8-27) and 806R (806-787); and 515F (515-533) and 13R (1390-1371). PCR products were characterized by direct sequencing or cloning and sequencing, followed by comparison with rDNA sequences available in GenBank (11).

Criteria for Causation

Cases were defined as having definite, probable, possible, or no microbial etiology (Table 1). These levels of certainty for the causal role of an infectious agent reflected the integration of several factors, including the relationship of anatomic site of detection to site of disease, reliability of the method, and whether the putative agent was a known cause of the clinical syndrome under investigation. Cases were classified as explained if results showed a definite or probable disease cause and as unexplained if results indicated a possible infectious cause or none at all.

Statistical Analysis

Analysis was performed by SAS 6.12 (SAS Institute, Cary, NC). Denominators for the population under surveillance, obtained from the 1992 intercensuses (14), included all persons in the age groups under surveillance at the various sites; denominators including only previously healthy persons are not available, and no attempt was made to estimate this frac-

3IC was available at CDC for the following pathogens: Acanthamoeba culbertsoni; adenovirus; Bacillus anthracis; Balamuthia spp.; Bartonella henselae, Bartonella quintana; Brucella spp.; Chlamydia spp.; Coccioidiodes spp; Coxiella burnetii; Crimerror-Congo hemorrhagic fever virus; Cryptococcus spp.; Cytomegalovirus; Dengue virus; Eastern equine encephalitis virus; Ebola virus; Ehrlichia chaffeensis; Enterovirus (Pan-Enterovirus); human enterovirus 71; Flavivirus; Japanese encephalitis serocomplex group (West Nile virus, St. Louis encephalitis virus [SLEV], Japanese encephalitis virus [JEV], Francisella tularensis; Group A streptococci; Guanarito virus (Venezuelan hemorrhagic fever virus); Hantavirus; Helicobacter pylori; Hendra virus; herpes simplex viruses 1 and 2; Histoplasma spp.; human granulocytic ehrlichiosis; Human hanta- pesivirus 6; HIV-1; HIV-2; B19 virus (B19V); Influenza A virus (FLUA); Influenza B virus (FLUB); Junin virus (Argentine hemorrhagic fever); La Crosse virus; Lassa virus; Legionella pneumophila serogroups 1, 5, 6; Leptospira spp.; Listeria monocytogenes; Lymphocytic choriomeningitis virus (LCMV); Machupo virus (Bolivian hemorrhagic fever); Marburg virus; measles (Edmonston) virus (MeV); Mycobacterium spp.; Myo- plasma pneumoniae; Naegleria fowleri; Neisseria meningitidis C; Nipah virus; Human parainfluenza virus types 1 and 3 (HPIV 1,3); Rabies virus (RABV); Human respiratory syncytial virus (HRSV); Rickettsia spp.; Orientia group; Rickettsia spp. spotted fever group; Rickettsia spp. Typhus group; Rift Valley fever virus; Rotavirus; Streptococcus pneumoniae; Toxoplasma gondii; Treponema pallidum; Trypanosoma cruzi; varicella-zoster virus (VZV); Venezuelan equine encephalitis virus; Western equine encephalomyelitis virus (WEEV); Yellow fever virus; and Yersinia pestis.

4The following viral tests were used at the CDHS Viral and Rickettsial Diseases Laboratory: IgG was detected by both EIA and IFA for adenovirus, HHV-6, HHV-8, herpes simplex virus, FLUA, FLUB, MeV, Mumps virus (MuV), HPIV-1,4, HRSV, Rubella virus (RUBV), VZV, SLEV, and WEEV. IgG was detected by EIA only for Hantavirus (Sin Nombre virus [SNV]) and B-19. IgG was detected by IFA only for Epstein-Barr virus (viral capsid antigen), LCV, and RABV. IgM was detected by both EIA and IFA for HHV-6, herpes simplex virus, MeV, MuV, HPIV-1,4, HRSV, RUBV, and VZV. IgM was detected by EIA only for enterovirus, hantavirus (SNV), and B19V. IgM was detected by IFA only for Epstein-Barr virus. PCR tests performed were Herpesvirus consensus PCR (6); enterovirus PCR, modified from (7) [Antisense primer (1R): 5’-ATT GTC ACC ATA AGC AGC CA; sense primer (1L): 5’-CCT CGG GCC CCT GAA TGG GAA TAA T]; and adenovirus PCR (8).
Table 1. Classification of laboratory test results and cases,\(^a\) surveillance for unexplained death and critical illness possibly due to infectious causes (UNEX), 1995–1998

<table>
<thead>
<tr>
<th>A</th>
<th>B</th>
<th>C</th>
</tr>
</thead>
<tbody>
<tr>
<td>1. Detection of organism by culture from involved site(^b)</td>
<td>1. Detection of organism by culture, IF, IHC, IEM, ISH, or PCR(^a) in blood or clinically relevant site(^c)</td>
<td>1. Detection of organism by culture, IF, IHC, IEM, ISH, or PCR from uninvolved, but nonmucosal, noncutaneous site</td>
</tr>
<tr>
<td>2. Detection of organism by direct immunologic staining (i.e., IF, IHC, IEM) at involved site</td>
<td>2. Positive serology: 24-fold change in IgG/IgA titer or significantly elevated IgM titer</td>
<td></td>
</tr>
<tr>
<td>3. Detection of organism by DNA/RNA ISH at involved site</td>
<td>3. Detection of organism by EM(^a) at involved site</td>
<td></td>
</tr>
<tr>
<td>4. Detection of organism by PCR(^a) at involved site</td>
<td>4. Detection of other specific microbial antigen at characteristic site (e.g., urine, CSF)</td>
<td></td>
</tr>
</tbody>
</table>

\(^a\)Case classification: A case was considered to have a definite explanation if the organism was a well-recognized cause of the syndrome and there was one test from column A, or 2 from column B. A case was considered to have a probable explanation if the organism was a well-recognized cause of the syndrome and there was one test from column B, or if the organism was not a well-recognized cause of the syndrome and there was one test from column A, or 2 from column B. A case was considered to have a possible explanation if the organism was not a well-recognized cause of the syndrome and there was one test from column B, or if there was one test from column C, regardless of whether organism is known to cause the syndrome. 

\(^b\) Involved\(^f\) refers to the presence of typical pathology.

\(^c\) IF = immunofluorescence, IHC = immunohistochemistry, IEM = immuno-electron microscopy, ISH = in situ hybridization, PCR = polymerase chain reaction, Ig = immunoglobulin, EM = electron microscopy, CSF = cerebrospinal fluid.

\(^d\) For example, bronchoalveolar lavage in respiratory syndrome.

\(^e\) EM is often nonspecific and may not permit reliable microbial identification without further characterization (e.g., IEM).

\(^f\) Specific or broad range PCR/reverse transcriptase-PCR; product must be characterized beyond size determination (e.g., sequencing, single-strand conformation polymorphism, restriction fragment-length polymorphism, or probe hybridization).

Results

Epidemiology

From May 1, 1995, to December 31, 1998, 525 possible cases were reported to UNEX personnel; 388 of these reports were excluded. The three most common reasons for exclusion were the presence of a preexisting medical condition (33%), residence outside the surveillance area (17%), and cause identified by local health-care providers on further testing (26%). Among cases excluded for the last reason, 72% had an infectious cause identified.

A total of 137 cases met the case definition, for a minimum overall annual rate of 0.5 per 100,000 population. After data were adjusted for age and race, this rate translates into 920 cases in the United States each year. The overall annual incidence rates remained stable over time, but varied among the different sites from 0.3 to 2.3 per 100,000 per year. The highest rate was in Connecticut, where active surveillance was conducted in a well-defined population of approximately 500,000 persons. Forty-one (30%) of the case-patients died, of whom 30 (73%) had autopsies performed, reflecting a rate much higher than the national autopsy rate of <11% (15). Cases were reported a median of 6 days from time of admission to the hospital (0 to 289 days).

The median age of case-patients was 20 years; 20 (15%) were 1 to 4 years of age, 53% were female, and 82% were white. The incidence rates varied by age group (Figure 1) but did not differ by sex and race. No differences were observed in the seasonal distribution of cases, nor was there clustering of cases by time or place. As for exposures, 54% of all cases were reported to have pets, which is similar to national rates of pet ownership: 54% to 64% (American Veterinary Medical Association U.S. Pet Ownership and Demographics Sourcebook); 8% had traveled outside the United States in the year before hospitalization, and 4% had received transfusions at least once in their lifetime.

Clinical Features

Table 2 summarizes the distribution of cases and the proportion explained by surveillance, as well as the syndrome-specific case-death ratios. The largest proportion of cases presented as a neurologic syndrome, followed closely by respiratory syndrome. The highest syndrome-specific case-death ratio was seen among cases with cardiac syndrome (46%) and the lowest among cases with neurologic syndrome (18%). An example of a case is described in Appendix III.

Surveillance Audit

Table 3 summarizes the results of the surveillance audits. The site-specific sensitivity (\(S_{P}\)) of our prospective surveillance for detecting unexplained deaths ranged from 38% in California to 100% in both Connecticut and Minnesota. Retrospective death record review identified 25% to 100% of all deaths detected through surveillance. Cases detected through surveillance but not by death record review were missed by the...
latter because the death certificates did not have the specified ICD-9 codes. The review of hospital discharge data focused on one tertiary-care referral hospital under surveillance in California and the one in Oregon, but included the entire surveillance area in Connecticut (16). Of potential cases identified by the selected ICD-9 codes, 90% to 96% were excluded, indicating the lack of specificity of these codes. The sensitivity of our prospective surveillance to detect only critical illnesses due to infectious causes in the United States. To our knowledge, this is the first public health attempt to describe the features of this problem, in spite of its clinical complexities. This project established the infrastructure needed to detect UNEX cases, attempt to identify their etiology, and ultimately identify new infectious agents. However, since this project was a pilot study, it was difficult to standardize many of its aspects. Many lessons were learned during this project, whether related to the best surveillance methods to use or the laboratory testing process. In

<table>
<thead>
<tr>
<th>Syndrome</th>
<th>No. (%)</th>
<th>No. of deaths (%)</th>
<th>No. of explained /cases with specimens (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Neurologic</td>
<td>39 (29)</td>
<td>7 (18)</td>
<td>15/37 (41)</td>
</tr>
<tr>
<td>Respiratory</td>
<td>36 (26)</td>
<td>11 (31)</td>
<td>13/33 (39)</td>
</tr>
<tr>
<td>Cardiac</td>
<td>28 (20)</td>
<td>13 (46)</td>
<td>3/22 (14)</td>
</tr>
<tr>
<td>Multisystem</td>
<td>18 (13)</td>
<td>4 (22)</td>
<td>3/15 (20)</td>
</tr>
<tr>
<td>Hepatic</td>
<td>9 (7)</td>
<td>4 (44)</td>
<td>0/8 (0)</td>
</tr>
<tr>
<td>Other</td>
<td>7 (5)</td>
<td>2 (29)</td>
<td>0/7 (0)</td>
</tr>
<tr>
<td>All cases</td>
<td>137</td>
<td>41 (30)</td>
<td>34/122 (28)</td>
</tr>
</tbody>
</table>

**Search for Etiologic Agents**

Of the 137 UNEX case-patients, 122 had specimens available for testing; 10 of these had tissue specimens only. Of the 122 cases, 34 (28%) could be attributed to a specific infectious agent; these agents were classified as definite or probable causes of the illness, based on our criteria (Table 1). Specific infectious causes and the laboratory methods used for diagnosis are listed in Table 4. Table 5 lists additional infectious causes for possible cases that did not meet our criteria for definite or probable causation. All the infectious agents identified in this study were previously recognized bacterial and viral pathogens. One patient, admitted because of syncope, was found to have a complete heart block and had evidence of simultaneous infection with *Borrelia burgdorferi* and *Ehrlichia chaffeensis*, which has been previously reported (17). A number of cases met the clinical definition for various infectious diseases syndromes, including toxic shock syndrome (five cases), but did not meet our definition for an explained case. In addition, four cases had evidence of polyclonal serologic response to multiple infectious agents and therefore could not be attributed to a specific etiology. The proportion of explained cases was largest among those with neurologic syndromes, followed by those with respiratory syndromes; it was higher among surviving patients (29%) than among patients who died (15%), although this difference was not statistically significant (p=0.2) (Figure 2). Explained cases were similar to unexplained cases in terms of patient age, sex, and race, but were reported sooner after admission than unexplained cases (median 4 vs. 7.5 days, respectively; p=0.1). The proportion of explained cases during 1998 (7 [17%] of 41), when laboratory testing protocols were used routinely for first-round testing, did not differ significantly from the same proportion for cases enrolled during 1995-1997 (27 [28%] of 96) when no such protocol was used (p>0.05).

Clinical specimens from each enrolled patient underwent an average of 28 laboratory tests (up to 103 tests). The mean number of tests performed did not differ substantially for explained and unexplained cases (30 vs. 27, respectively). None of the cases with only histologic specimens available had an infectious cause identified. Of the 34 explained cases, 23 (68%) were explained by using serologic tests, 7 (21%) by specific primer PCR, and 4 (12%) by 16S rDNA PCR. Among the 122 cases with specimens, serologic testing provided the highest yield in identifying infectious causes (23 [22%] of 104), followed by specific primer PCR (7 [10%] of 70) and 16S rDNA PCR (4 [8%] of 48). An infectious etiology was more likely to be identified in cases with paired serum specimens (14 [23%] of 62) than in those with single serum specimens (2 [5%] of 42) (p=0.05).

**Discussion**

This study is the first to measure the population burden of unexplained deaths and critical illness from possibly infectious causes in the United States. To our knowledge, this is the first public health attempt to describe the features of this problem, in spite of its clinical complexities. This project established the infrastructure needed to detect UNEX cases, attempt to identify their etiology, and ultimately identify new infectious agents. However, since this project was a pilot study, it was difficult to standardize many of its aspects. Many lessons were learned during this project, whether related to the best surveillance methods to use or the laboratory testing process. In
addition, data obtained in the first 3.5 years of this project suggest that UNEX occur in previously healthy persons at rates similar to those of other conditions of clear public health concern and priority (18). Of obvious concern is also the large proportion of these deaths and severe illnesses that remains unexplained after extensive laboratory testing. Our findings highlight the substantial limitations of available diagnostic tests for infectious diseases and the need for improved tests and novel approaches to identify infectious disease agents.

Our surveillance estimated the burden of disease only among previously healthy persons 1 to 49 years of age. Since a different age cut-off was used in Oregon, the final rates of disease were adjusted for age and race. The lower age limit was chosen to avoid confusion with congenital problems seen in infants but to include most children in day care, where infectious diseases are common and new infectious diseases might spread rapidly. The upper age limit was intended to exclude an expected increased proportion of unexplained deaths due to noninfectious causes among persons ≥50 years of age. Although immunocompromised patients are more susceptible to a variety of infectious diseases, available resources and a concern that the clinical relevance of novel microbial findings would be more difficult to interpret in immunocompromised persons compelled us to focus on previously healthy persons. In addition, many of the new infectious diseases first identified in these persons have subsequently been found to affect persons with normal immune systems (19,20).

The surveillance methods adopted during this project were customized to meet the objectives of this study, taking into consideration the limitations of local resources; therefore UNEX cannot be easily compared with other classical surveillance systems. The different methods of surveillance used at the four sites allowed us, through the surveillance audits and validation, to determine how these differences affected case-finding. For example, investigators in Connecticut were able to detect most UNEX cases largely because they conducted more active surveillance in a smaller population base; in this site, surveillance focused on all seven hospitals in New Haven County. At the academic tertiary-care hospital, EIP staff reviewed ICU admission logs and communicated with clinicians daily. At the other six hospitals, a stimulated passive surveillance system was used in which physicians and infection control practitioners were given reminders several times per year. The active prospective method captured a greater proportion of total cases (86% of cases at the single hospital) then did the passive methods (50% of total cases at the six remaining hospitals).

If this surveillance is to be expanded, different methods may be chosen, depending on availability of resources and overall objectives. Less resource-intensive passive surveillance may be used if the goal is to monitor trends in disease occurrence. For example, although analyzing all death certifi-

**Table 4. Infectious disease causes for explained cases, UNEX,1995–1998, California, Oregon, Connecticut, and Minnesota (n=34)**

<table>
<thead>
<tr>
<th>Syndrome</th>
<th>Etiology (n)</th>
<th>Tests (n)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Neurologic</td>
<td>Neisseria meningitidis (4)</td>
<td>16S rDNA PCR (2), PCR (1), EIA IgM (1)</td>
</tr>
<tr>
<td></td>
<td>Bartonella henselae (1)</td>
<td>PCR, IFA IgG</td>
</tr>
<tr>
<td></td>
<td>Bartonella spp. (2)</td>
<td>IFA IgG</td>
</tr>
<tr>
<td></td>
<td>Chlamydia pneumoniae (1)</td>
<td>MIF IgG</td>
</tr>
<tr>
<td></td>
<td>Mycoplasma pneumoniae (1)</td>
<td>EIA IgM/IgG</td>
</tr>
<tr>
<td></td>
<td>Cytomegalovirus (1)</td>
<td>EIA &amp; IFA IgG</td>
</tr>
<tr>
<td></td>
<td>Coxsackie B (1)</td>
<td>EIA IgM, viral culture</td>
</tr>
<tr>
<td></td>
<td>Enterovirus (1)</td>
<td>EIA IgM</td>
</tr>
<tr>
<td></td>
<td>Epstein-Barr virus b (1)</td>
<td>IFA IgG (VCA and EA)</td>
</tr>
<tr>
<td>Respiratory</td>
<td>Human herpes virus 6 (1)</td>
<td>IFA and EIA IgM (IgM and IgG)</td>
</tr>
<tr>
<td></td>
<td>Mumps virus (1)</td>
<td>IFA IgM, IFA and EIA IgG</td>
</tr>
<tr>
<td></td>
<td>Chlamydia pneumoniae (2)</td>
<td>MIF IgG (2), IFA IgM</td>
</tr>
<tr>
<td></td>
<td>Mycoplasma pneumoniae (4)</td>
<td>PCR (blood), EIA IgM/IgG</td>
</tr>
<tr>
<td></td>
<td>Streptococcus pneumoniae (2)</td>
<td>16S rDNA PCR (pleural fluid)</td>
</tr>
<tr>
<td></td>
<td>Legionella spp. (1)</td>
<td>PCR (from lung)</td>
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<tr>
<td></td>
<td>Adenovirus (1)</td>
<td>EIA and IFA IgG</td>
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<tr>
<td></td>
<td>Influenza B virus (1)</td>
<td>EIA and IFA IgG</td>
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<tr>
<td></td>
<td>Influenza A virus (1)</td>
<td>EIA and IFA IgG (IgM, IFA IgG)</td>
</tr>
<tr>
<td></td>
<td>Human parainfluenza virus (1)</td>
<td>EIA and IFA IgG</td>
</tr>
<tr>
<td></td>
<td>Borrelia burgdorferi/</td>
<td>EIA/IFA flagella IgG, Western blot (IgG IgM)</td>
</tr>
<tr>
<td></td>
<td>Ehrlichia chaffeensis (1)</td>
<td></td>
</tr>
<tr>
<td></td>
<td>Enterovirus (1)</td>
<td>EIA IgM</td>
</tr>
<tr>
<td></td>
<td>Legionella spp. (1)</td>
<td>PCR (heart)</td>
</tr>
<tr>
<td>Multisystem</td>
<td>Neisseria meningitidis (1)</td>
<td>PCR (CSF)</td>
</tr>
<tr>
<td></td>
<td>Adenovirus (1)</td>
<td>PCR (blood)</td>
</tr>
<tr>
<td></td>
<td>Enterovirus (1)</td>
<td>IgM EIA</td>
</tr>
</tbody>
</table>

*EIA = enzyme immunosorbent assay, IFA = indirect immunofluorescent assay, IG = immunoglobulin, MIF = microimmunofluorescence, PCR = polymerase chain reaction.

*See Appendix III for a detailed description of this case.

**Table 5. Infectious causes for “possibly” explained cases, UNEX, 1995–98, California, Oregon, Connecticut, and Minnesota (n=34)**

<table>
<thead>
<tr>
<th>Syndrome</th>
<th>Etiology (n)</th>
<th>Tests (n)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Neurologic</td>
<td>Mycoplasma pneumoniae</td>
<td>Remel EIA (IgM/IgG)</td>
</tr>
<tr>
<td></td>
<td>Influenza B virus (FLUBV)</td>
<td>Nasopharyngeal culture</td>
</tr>
<tr>
<td></td>
<td>Varicella-zoster virus (reactivation)</td>
<td>EIA/IFA IgG</td>
</tr>
<tr>
<td>Respiratory</td>
<td>Enterovirus</td>
<td>EIA IgM</td>
</tr>
<tr>
<td>Cardiac</td>
<td>Chlamydia pneumoniae</td>
<td>MIF IgG</td>
</tr>
<tr>
<td></td>
<td>Adenovirus</td>
<td>EIA IgM</td>
</tr>
<tr>
<td></td>
<td>FLUBV</td>
<td>IFA IgM</td>
</tr>
<tr>
<td>Other</td>
<td>Enterovirus (2)</td>
<td>IgM EIA</td>
</tr>
</tbody>
</table>

*EIA = enzyme immunosorbent assay, IFA = indirect immunofluorescent assay, IG = immunoglobulin, MIF = microimmunofluorescence, PCR = polymerase chain reaction.

*Other syndromes included one case with thrombotic thrombocytopenic purpura and one with hemolytic uremic syndrome.
cates for UNEX cases would be prohibitively time-consuming, electronically searching only the certificates in which the manner of death was recorded as natural, undetermined, or pending investigation could substantially decrease the workload. Under a passive system, maintaining good communication between study staff and clinical staff (clinicians, pathologists, and infection control practitioners) is critical and aided by the provision of diagnostic testing not locally available (such as serologic testing for hantavirus or toxin testing for botulism) and timely feedback of study results. Such collaboration may be critical to early diagnosis of diseases that produce characteristic clinical syndromes (e.g., potential bioterrorist agents such as botulism) or that are not readily confirmed by clinical laboratories.

Before initiating this project, we had reviewed multiple cause-of-death data for the United States to estimate the number of unexplained deaths from possibly infectious causes at these EIP sites (3). In 1992, the rate of unexplained deaths among healthy persons 1 to 49 years of age was 8.9 per 100,000 population. The discrepancy between this rate and that found in our study (0.5 per 100,000) is likely due to the low specificity of ICD-9 codes in excluding persons with previous health problems, as well as the problems related to retrospective analysis in general. For at least two reasons, we expect that the incidence of UNEX found in this study represents only a minimal estimate of the true burden of this problem. First, the denominator in our calculations included all persons in our designated groups, since we chose not to estimate the fraction of previously healthy persons in the surveillance populations at the four sites. Second, the differences in incidence rates between the four surveillance sites and results of the surveillance audits support the assumption that the overall rate detected was a minimal estimate of overall disease.

An important unresolved issue from our study is the large proportion of cases that remained unexplained, even after extensive laboratory testing. Although a standardized protocol for testing was used only during 1998, the proportion of explained cases before and after this protocol was used did not differ substantially. Some illnesses may have noninfectious causes, especially given the lack of specificity in our clinical criteria for case inclusion and in the features of infection in general. In cardiac syndromes, for example, myocarditis and myocardial infarction can have very similar presentations. Some cases may have been caused by microbial products such as toxins without the presence of the organism or substantial amounts of its nucleic acids. Laboratory methods for screening and detection of toxins remain inadequate. For some patients, specimens were not available from the primary site of disease, were severely limited in quantity, or were only available from late in the course of the disease; in many cases, multiple serum specimens were not available, autopsies were incomplete, and tissue specimens were obtained only from unaffected organs. Finally, the breadth of our testing methods may not have been adequate. Since broad-range PCR methods were applied only to bacteria and a limited range of viruses, many other potential agents may have been missed. Our approach to the detection of viral pathogens relied more heavily on serologic and immunohistochemical techniques, in part because of the difficulty in designing a comprehensive set of consensus PCR primers for all known viral families (21). In our study, viral testing was also constrained by limited experience with certain IgM assays. The development, testing, and application of comprehensive broad-range viral and fungal consensus primers for use in PCR assays may be helpful. Through this project, we created a population-based bank of clinical specimens that may prove valuable in the search for newly recognized etiologic agents, the development of diagnostic tests, and the standardization of nucleic acid-based techniques for identifying previously unknown etiologic agents.

This project represents an attempt to build capacity for early detection and response to emerging infectious diseases threats in the United States and elsewhere. The usefulness of this surveillance system for UNEX was recently illustrated during an outbreak of West Nile virus encephalitis in the northeastern United States (22) and an outbreak of unexplained illness among injecting drug users in Scotland and Ireland (23); initial reports of illness from both these investigations were received through the UNEX surveillance project, and initial testing was conducted through the infrastructure developed for this project. Future surveillance for UNEX may benefit from simplified case-finding methods, improved specimen quality, and more focused syndrome-specific surveillance. Once validated, surveillance methods may be adopted by the broader public health community. Such surveillance approaches will strengthen the collaboration between clinicians, laboratorians, and public health professionals, leading to improved detection of unexplained deaths and critical illnesses, including possible bioterrorism events, and better monitoring of emerging infectious diseases.

Acknowledgments

We thank the following scientists for their extensive help in discussion and laboratory testing of the various UNEX cases: CDC: Barbara Anderson, Larry Anderson, Mary Bajani, Bernie Beall, David Beall, Robert Benson, Cheryl Bopp, Sandra Bragg, George Carlone, Leon Carter, Roberta Challener, May Chu, Cheryl Elie, Dean Erdman,
Dr. Hajjeh is chief of the epidemiology unit of the mycotic diseases branch, Division of Bacterial and Mycotic Diseases, Centers for Disease Control and Prevention. Her research interests include improving surveillance for infectious diseases, epidemiology of fungal diseases, and evaluation of prevention strategies.

References

Address for correspondence: Rana A. Hajjeh, Division of Bacterial and Mycotic Diseases, CDC, Mailstop C09, 1600 Clifton Rd, NE, Atlanta, GA 30333, USA; fax: 404-639-3059; e-mail: rlh5@cdc.gov
Appendix I

Case Definition, Surveillance for Unexplained Deaths and Critical Illnesses Due to Possibly Infectious Causes, United States, 1995–1998

Previously Healthy

Patients are considered previously healthy who had no known preexisting chronic medical condition before the onset of the illness resulting in hospitalization or death, including malignancy; HIV infection; chronic cardiac; pulmonary, renal, hepatic, or rheumatologic disease; or diabetes mellitus. These patients have no history of immunosuppressive therapy, trauma thought to be related to illness, evidence of toxic ingestion or exposure, or nosocomial infection.

Reasons for Exclusion

1. A history of a malignancy other than nonmelanoma skin cancer
2. HIV infection identified during hospitalization, previously or after discharge
3. History of physician-confirmed myocardial infarction, angina with known coronary artery disease, or congestive heart failure
4. Any history of hospital admission for asthma or other pulmonary diseases except for uncomplicated pneumonia
5. History of dialysis or chronically elevated blood urea nitrogen and creatinine
6. Biopsy-proven liver disease of any kind or chronic coagulopathy or chronic Hepatitis B or C virus infection as a result of hepatic insufficiency
7. Physician-confirmed rheumatologic conditions requiring chronic or intermittent medical therapy with oral steroids or other immunosuppressive drugs
8. Any known physician-confirmed diabetes mellitus previously or during hospitalization
9. Development of hallmarks of infection >48 hours after hospital admission
10. Any mention of a history of excessive alcohol use, alcohol abuse, or alcoholism is a reason for exclusion (e.g., delirium tremens, withdrawal seizures, alcoholic neuropathy, persistent liver function test abnormalities, gastrointestinal bleeding, coagulopathy, or hypoalbuminemia).
11. Any mention of injecting drug use
12. Any history of neurologic disease, including seizures,
13. Obesity, defined as body mass index ≥30 or “obese” noted in medical chart
14. Physician-confirmed diagnosis of anorexia

Not Reasons for Exclusion

1. Hypertension or a history of hypertension
2. Any history of inhaler use
3. Pyelonephritis or nephrolithiasis or a history of either of these conditions in the absence of a chronically elevated blood urea nitrogen and creatinine
4. History of hepatitis
5. Pregnancy

Appendix II

Algorithm for Meningo-Encephalitis available online only at URL: http://www.cdc.gov/ncidod/EID/vol8no2/pdf/01-0165-app2.pdf

Appendix III

An Example of a Clinical Case Surveillance for Unexplained Deaths and Critical Illnesses Due to Possibly Infectious Causes, United States, 1995–1998

A 22-month-old boy from Oregon was healthy except for previous bouts of otitis media, for which tympanostomy tubes had been placed. Three days before admission, in May 1997, tactile fever was noted, and one day before admission, the patient had decreased activity and rhinorrhea. On the day of admission, he vomited twice. In the emergency room, he had a temperature of 39.2°C, was irritable and lethargic, and had nuchal rigidity. A complete blood count showed a total leukocyte count (WBC) of 18,300 (69% segmented cells, 8% bands). Cerebrospinal fluid (CSF) analysis showed a WBC of 54 (25% segmented cells, 75% monocytes), protein 38 mg/dL, and glucose 70 mg/dL. The patient was hospitalized and initially treated with ceftriaxone. On the next day, he became less responsive, and abnormal posturing developed in the left upper and lower extremities. A computed tomography scan of the head (without and with contrast) was normal. He was transferred to a tertiary-care center, where an electroencephalogram showed moderate generalized slowing and recurrent right hemispheric electrographic seizures. A magnetic resonance imaging scan done on the same day showed a diffusely increased white matter signal consistent with viral encephalitis or acute disseminated encephalomyelitis. The patient received acyclovir for 3 days. His responsiveness and clinical condition gradually improved, and he was transferred to a rehabilitation service 17 days after admission. Initial work-up at the hospital revealed negative blood cultures and negative bacterial and viral cultures of the CSF. PCR for Epstein-Barr virus in the blood and CSF was negative, as was PCR for herpes simplex virus in CSF.

The patient was enrolled in the UNEX project and evaluated. Specimens available for testing included acute- and convalescent-phase serum and CSF specimens. A variety of tests were conducted (see neurologic syndrome testing protocol in Appendix II). Because the quantities of specimens available were limited, testing was prioritized. First-round testing was negative for Cytomegalovirus, HHV-6, and arboviruses. However, testing for IgG antibodies (by IFA) for Epstein-Barr viral capsid antibodies showed a fourfold rise in titer between acute- and convalescent-phase serum specimens; testing for IgG antibodies (also by IFA) to Epstein-Barr early antigen revealed a fourfold decrease in titer between convalescent- and acute-phase serum specimens, indicating acute Epstein-Barr infection.
Appendix I

Case Definition, Surveillance for Unexplained Deaths and Critical Illnesses Due to Possibly Infectious Causes, United States, 1995-1998

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5. History of dialysis or chronically elevated blood urea nitrogen and creatinine
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7. Physician-confirmed rheumatologic conditions requiring chronic or intermittent medical therapy with oral steroids or other immunosuppressive drugs

8. Any known physician-confirmed diabetes mellitus previously or during hospitalization

9. Development of hallmarks of infection >48 hours after hospital admission

10. Any mention of a history of excessive alcohol use, alcohol abuse, or alcoholism is a reason for exclusion.(e.g., delirium tremens, withdrawal seizures, alcoholic neuropathy, persistent liver function test abnormalities, gastrointestinal bleeding, coagulopathy, or hypoalbuminemia).

11. Any mention of injecting drug use

12. Any history of neurologic disease, including seizures,

13. Obesity, defined as body mass index \( \geq 30 \) or “obese” noted in medical chart

14. Physician-confirmed diagnosis of anorexia

**Not Reasons for Exclusion**

1. Hypertension or a history of hypertension

2. Any history of inhaler use

3. Pyelonephritis or nephrolithiasis or a history of either of these conditions in the absence of a chronically elevated blood urea nitrogen and creatinine

4. History of hepatitis

5. Pregnancy
Algorithm for Meningo-Encephalitis

Specimens (brain, meninges, spinal cord, nerve) will be sent to Centers for Disease Control and Prevention (CDC) for testing by monoclonal antibody, hybridization, electron microscopy, or polymerase chain reaction (PCR). Before any testing, 250 µL (or other amount) from each specimen will be stored in a serobank. First-round tests will be performed on all specimens. Culture for bacterial and viral organisms are to be performed at primary site.

Date of symptom onset __/__/__
Date of hospital admission __/__/__

Available specimens (with dates)
- Serum
- Blood
- Cerebrospinal fluid (CSF)
- Blood culture
- Pleural fluid
- Pericardial fluid
- Peritoneal fluid
- Other

First-Round Testing

**Viral Panel**
Viral PCR preferable if acute-phase samples available (CSF, serum, whole blood, saliva [for rabies ≥1.0 mL])
- Adenovirus (100 µL serum, CSF)
- Cytomegalovirus
- *Enterovirus (coxsackie A and B, Echo, etc.)*
- Epstein-Barr Virus
- Herpes simplex virus (HSV) (150 µL serum, CSF)
- *Rabies virus (RABV)* (saliva)
- *Varicella-zoster virus (VZV)*
- *Human herpesvirus (HHV-6)* (for children <3 years of age)

**Arboviruses**
- California encephalitis group
- *Colorado tick fever virus (CTFV)*
- *Eastern equine encephalomyelitis virus (EEEV)*
- *St. Louis encephalitis virus (SLEV)*
- *Venezuelan equine encephalomyelitis virus (VEEV)*
- *Western equine encephalomyelitis virus (WEEV)*

Serologic testing on CSF and serum--(0.25 mL)
- Adenovirus immunoglobulin (Ig) M (enzyme immunoassay [EIA]-2.5 µL serum, immunofluorescent assay [IFA]-30 µL serum)
Adenovirus IgG (EIA-2.5 µL serum, IFA-50 µL serum)
Cytomegalovirus IgM (EIA-2.5 µL serum, IFA-30 µL serum)
Cytomegalovirus IgG (EIA-2.5 µL serum, IFA-50 µL serum)
Epstein-Barr virus IgM (IFA-30 µL serum)
Epstein-Barr virus IgG (IFA-50 µL serum)
Epstein-Barr nucleocapsid antigen (IFA-100 µL serum)
*Enterovirus* IgM (*Coxsackie, Echo*) (EIA 5 µL serum, 50 µL CSF)
HSV IgM (EIA-2.5 µL, IFA-30 µL serum)
HSV IgG (EIA-2.5 µL, IFA-50 µL serum)
*Influenza A virus* (FLUAV) IgM (EIA-2.5 µL serum, IFA-30 µL serum)
FLUAV IgG (EIA-2.5 µL serum, IFA-50 µL serum)
*Influenza B virus* (FLUBV) IgM (EIA-2.5 µL serum, IFA-30 µL serum)
FLUBV IgG (EIA-2.5 µL serum, IFA-50 µL serum)
*Measles (Edmonston) virus* (MeV) IgM (EIA-2.5 µL, IFA-30 µL serum)
MeV IgG (EIA-2.5 µL, IFA-50 µL serum)
*Human parainfluenza virus* (HPIV 1-4) IgM (complement fixation [CF]-serum, EIA-2.5 µL serum, IFA-30 µL serum)
HPIV 1-4 IgM (CF-serum, EIA-2.5 µL serum, IFA-50 µL serum)
RABV IgG (IIF-50 µL serum, CSF; NT-50 µL serum, CSF)
VZV IgM (EIA-2.5 µL, IFA-30 µL serum)
VZV IgG (EIA-2.5 µL, IFA-50 µL serum)

**Arboviruses**
California group IgG (IFA capture-50 µL serum, CSF)
EEEV IgM (enzyme-linked immunosorbent assay [ELISA] capture-2.5 µL serum, CSF)
EEEV IgG (ELISA capture-2.5 µL serum, CSF)
*Powassan virus* (POWV) IgM (ELISA capture-2.5 µL serum, CSF)
POWV IgG (ELISA capture-2.5 µL serum, CSF)
SLEV IgM (ELISA capture-2.5 µL serum, CSF)
SLEV IgG (ELISA capture-2.5 µL serum, CSF)
VEEV IgM (ELISA capture-2.5 µL serum, CSF)
VEEV IgG (ELISA capture-2.5 µL serum, CSF)
WEEV IgM (ELISA capture-2.5 µL serum, CSF)
WEEV IgG (ELISA capture-2.5 µL serum, CSF)
HHV-6 IgM (EIA-2.5 µL serum, IFA-30 µL serum) (for children < 3 years of age)
HHV-6 IgG (IFA-50 µL serum)

**Bacterial panel**
Criteria: CSF PMNs >400, total protein >100, glucose <40, sudden death, prior antibiotic use
CSF--(0.5 mL)
*Borrelia burgdorferi* IgM (ELISA-5 µL CSF)
*B. burgdorferi* IgG (ELISA-5 µL CSF)
*Haemophilus influenzae* antigen (Ag) (LA-50 µL CSF)
*Mycoplasma pneumoniae* IgM/IgG (Remel ELISA-50 µL serum)
*Neisseria meningitidis* PCR (100 µL)
N. meningitidis Ag (LA-A,C,Y,W-135-250 µL CSF)
Streptococcus pneumoniae Ag (LA-50 µL CSF)
Syphilis (VDRL-CSF)

Serologic testing--(0.5 mL)
B. burgdorferi IgM (ELISA-5 µL serum)
B. burgdorferi IgG (ELISA-5 µL serum)
H. influenzae Ag (LA 50 µL)
H. influenzae IgM (ELISA-30 µL serum)
H. influenzae IgG (ELISA-30 µL serum)
Leptospira spp. (IgM/IgG MAT- 60 µL,IgM ELISA-5 µL)
M. pneumoniae IgM/IgG (Remel EIA-50 µL serum)
M. pneumoniae IgG (Zeus EIA-50 µL serum)
N. meningitidis Ag (LA-A,C,Y,W-135-250 µL serum)
N. meningitidis (ELISA-100 µL serum)
N. meningitidis A IgM
N. meningitidis A IgG
N. meningitidis B IgM
N. meningitidis B IgG
N. meningitidis C IgM
N. meningitidis C IgG
N. meningitidis Y IgM
N. meningitidis Y IgG
N. meningitidis W-135 IgM
N. meningitidis W-135 IgG
Syphilis (RPR-serum, FTA-ABS)

Serologic testing for Bartonella spp.
Bartonella elizabethae IgM (IFA-30 µL blood)
B. elizabethae IgG (IFA-50 µL blood)
B. henselae IgM (IFA-30 µL blood)
B. henselae IgG (IFA-50 µL blood)
B. quintana IgM (IFA-serum)
B. quintana IgG (IFA-serum)

Fungal panel (0.5 mL)
Cryptococcis neoformans Ag (LA-350 µL urine, serum, CSF)
Endemic Fungal Panel (CF-75 µL- serum, CSF; ID-100 µL serum, CSF)
Coccidioides inmitis (CF, ID, ELISA-serum, CSF)
Blastomyces dermatitidis (CF, ID, ELISA-serum, CSF)
Histoplasma capsulatum (CF, ID-serum , CSF)
H. capsulatum Ag (RIA-urine, serum, CSF; molecular probe-serum)
Paracoccidioides brasiliensis (CF, ID-serum, CSF)

Rash: (0.5 mL)
Erythema
Toxic Shock Syndrome Toxin-1
Maculopapular
*Ehrlichia chaffeensis* IgM (IFA-30 µL serum)
*E. chaffeensis* IgG (IFA-50 µL serum)
*E. phagocytophilia* (HGE) IgM (IFA-30 µL serum)
*E. phagocytophilia* (HGE) IgG (IFA-50 µL serum)
*Rickettsia rickettsii* IgM (IFA-30 µL serum)
*R. rickettsii* IgG (IFA-50 µL serum)
*R. typhi* IgM (IFA-30 µL serum)
*R. typhi* IgG (IFA-50 µL serum)
*Rubella virus* (RUBV) IgM (EIA-2.5 µL, IFA-30 µL serum)
RUBV IgG (EIA-2.5 µL, IFA-50 µL serum)
Vesiculonodular
*Variola* IgM (EIA-2.5 µL, IFA-30 µL serum)
*Variola* IgG (EIA-2.5 µL, IFA-50 µL serum)

Eosinophilia >8% or eosinophils in CSF:
*Paragonimus westermanii* (IB-serum)
*Schistosoma* spp. IgM/IgG (screening EIA-serum, speciate IB-serum)
*Strongyloides stercoralis* (serum)
*Taenia* spp. (cysticercosis) (IB-serum, CSF)
*Toxocara* spp. IgM/IgG (EIA-serum)
*Trichinella spiralis* IgM/IgG (EIA, bentonite flocculation-serum)

Hemorrhage: (0.5 mL)
with history of travel to Africa or South America
Hemorrhagic fever panel (*Crimean Congo hemorrhagic fever virus* [C-CHFV], *Lassa virus* [LASV], *Machupo virus* [MACV], *Junin virus* [JUNV], *Rift Valley fever virus* [RVFV], *Marburg virus* [MBGV], Ebola virus [EBOV], *Omsk hemorrhagic fever virus* [OHFV])
Hemorrhagic fever Ag (ELISA 250 µL serum)
Hemorrhagic fever IgM (ELISA 100 µL serum)
Hemorrhagic fever IgG (ELISA 100 µL serum)

**High-Priority Second Round (if not previously performed)**
Bacterial PCR (CSF and serum)—(0.5 mL)
*Mycobacterium tuberculosis*
*Neisseria meningitidis* PCR (100 µL)
16S universal probe (serum, whole blood, CSF)—(100 µL)

Bacterial serologic testing
Bacterial tests—(1.0 mL)
*Borrelia recurrentis* IgM (ELISA-5 µL serum)
*B. recurrentis* IgG (ELISA-5 µL serum)
Brucella spp. (MAT-serum 20 µL)

*Chlamydia pneumoniae* IgM (MIF-10 µL serum, EIA-2.5 µL serum, IFA-30 µL serum)
*C. pneumoniae* IgG (MIF-10 µL serum, EIA-2.5 µL serum, IFA-50 µL serum)

*Chlamydia psittaci* IgM (MIF-10 µL serum, EIA-2.5 µL serum, IFA-30 µL serum)
*C. psittaci* IgG (MIF-10 µL serum, EIA-2.5 µL serum, IFA-50 µL serum)

*Legionella pneumophila* IgM (IFA-25 µL serum, ELISA-10 µL serum)
*L. pneumophila* IgG (IFA-25 µL serum, ELISA-10 µL serum)
*L. pneumophila* Ag (RIA-100 µL urine)

*Listeria* IgM (ELISA-30 µL serum)
*Listeria* IgG (ELISA-30 µL serum)

Rickettsial serologic testing—(0.5 mL)
*Coxiella burnetti* IgM (IFA-5 µL serum)
*C. burnetti* IgG (IFA-50 µL serum)
*Ehrlichia canis* IgM (IFA-30 µL serum)
*E. canis* IgG (IFA-50 µL serum)
*E. chaffeensis* IgM (IFA-30 µL serum)
*E. chaffeensis* IgG (IFA-50 µL serum)
*E. phagocytophilia* (HGE) IgM (IFA-30 µL serum)
*E. phagocytophilia* (HGE) IgG (IFA-50 µL serum)
*Rickettsia rickettsii* IgM (IFA-30 µL serum)
*R. rickettsii* IgG (IFA-50 µL serum)
*R. typhi* IgM (IFA-30 µL serum)
*R. typhi* IgG (IFA-50 µL serum)

Viral PCR—
Cache Valley virus (CVV)
Jamestown Canyon virus (JCV)
*Murray Valley encephalitis virus* (MVEV)
*Powassan virus* (POWV)
Snowshoe hare virus (SSHV)
Tickborne complex

Viral serologic testing (serum)—(1.0 mL)
*Lymphocytic choriomeningitis virus* (LCM) Ag (EIA-50 µL serum)
LCM IgM (IF-serum)
LCM IgG (IF-serum)
MeV IgM (EIA-2.5 µL, IFA-30 µL serum)
MeV IgG (EIA-2.5 µL, IFA-50 µL serum)
*Mumps virus* (MuV) IgM (EIA-2.5 µL, IFA-30 µL serum)
MuV IgG (EIA-2.5 µL, IFA-50 µL serum)
RUBV IgM (EIA-2.5 µL, IFA-30 µL serum)
RUBV IgG (EIA-2.5 µL, IFA-50 µL serum)
Parasitology serologic testing-(0.5 mL)

*Babesia microti* (IFA-serum)
*Babesia* WA-1 (IFA-serum)
Malaria IgG (IFA-serum)
*Strongyloides stercoralis* (serum)

**Other tests to consider**

Viral PCR--
HTLV-I
with travel to Africa
*Rift Valley fever virus* (RVFV)
*West Nile virus* (WNV)
with travel to Asia
*Chikungunya virus* (CHIV)
Hendra virus (HeV)
*Japanese encephalitis virus* (JEV)
*Kyanasur Forest disease virus* (KFDV)

Viral serologic testing--(0.5 mL)
CVV Ag (ELISA 50 µL serum)
CVV IgM (ELISA 5 µL serum)
CVV IgG (ELISA 5 µL serum)
CHIV IgM (ELISA 5 µL serum)
CHIV IgG (ELISA 5 µL serum)
*Dengue virus* (DENV) (1-4) IgM (IF-serum)
DENV (1-4) IgG (IF-serum)
JCV IgM (ELISA 5 µL serum)
JCV IgG (ELISA 5 µL serum)
JEEV IgM (ELISA 5 µL serum)
JEEV IgG (ELISA 5 µL serum)
KFDV IgM (ELISA 5 µL serum)
KFDV IgG (ELISA 5 µL serum)
RVFV IgM (ELISA 5 µL serum)
RVFV IgG (ELISA 5 µL serum)
Tickborne complex IgM (ELISA 5 µL serum)
Tickborne complex IgG (ELISA 5 µL serum)
WNV IgM (ELISA 5 µL serum)
WNV IgG (ELISA 5 µL serum)
Yellow fever virus (YFV) Ag (ELISA 50 µL serum)
YFV IgM (ELISA 5 µL serum)
YFV IgG (ELISA 5 µL serum)

HTLV-I IgM (ELISA 5 µL serum)
HTLV-I IgG (ELISA 5 µL serum)
*Hepatitis A virus* (HAV) IgM (EIA-10 µL serum)
Hepatitis B virus (HBV) surface Ag (EIA-150 µL serum)
HBV surface Ab (EIA-200 µL serum)
HBV core IgM (EIA-10 µL serum)
HBV core total (EIA-100 µL serum)
HBV eAg (EIA-200 µL serum)
HBV eAb (EIA-100 µL serum)
Hepatitis C virus (HCV) IgM/IgG (EIA-30 µL serum with confirmation)
Hepatitis delta virus (HDV) IgM (EIA-10 µL serum)
HDV total (EIA-100 µL serum)
HDV Ag (EIA-50 µL serum)
Hepatitis E virus (HEV) IgM (EIA-50 µL serum)
HEV IgG (EIA-10 µL serum)

Bacterial tests--(1.0 mL)
Afipia felis IgM (IFA-30 µL blood)
A. felis IgG (IFA-50 µL blood)
Tropheryma whippelii IgM (IFA-30 µL blood)
T. whippelii IgG (IFA-50 µL blood)

Bartonella tests
B. quintana PCR (EDTA blood)
B. elizabethae PCR (EDTA blood)
B. henselae PCR (EDTA blood)
with travel to Latin America
B. bacilliformis PCR (EDTA blood)
B. bacilliformis IgM (IFA-30 µL blood)
B. bacilliformis IgG (IFA-50 µL blood)

Rickettsial serologies--(0.5 mL)
Rickettsia akari IgM (IFA-30 µL blood)
R. akari IgG (IFA-50 µL blood)
R. prowazekii IgM (IFA-30 µL blood)
R. prowazekii IgG (IFA-50 µL blood)
with travel to Africa
R. africae IgM (IFA-30 µL blood)
R. africae IgG (IFA-50 µL blood)
R. conorii IgM (IFA-30 µL blood)
R. conorii IgG (IFA-50 µL blood)
with travel to SE Asia, SW Pacific, Australia
Orientia tsutsugamushi IgM (IFA-30 µL blood)
O. tsutsugamushi IgG (IFA-50 µL blood)

Rickettsial PCR--(0.5 mL)
Ehrlichia chaffeensis
E. phagocytophilia (Human granulocytic ehrlichiosis)
O. tsutsumagushi
R. africae
R. akari
R. conorii
R. prowazekii
R. rickettsii
R. typhi

Parasitic serologic testing--(0.5 mL)
Paragonimus westermani (IB-serum)
Schistosoma spp. IgM/IgG (screening EIA-serum, speciate IB-serum)
Taenia spp. (cysticercosis) (IB-serum, CSF)
Trichinella spiralis IgM/IgG (EIA, bentonite flocculation-serum)
Toxocara spp. IgM/IgG (EIA,-serum)
Appendix III

An Example of a Clinical Case Surveillance for Unexplained Deaths and Critical Illnesses Due to Possibly Infectious Causes, United States, 1995-1998

A 22-month-old boy from Oregon was healthy except for previous bouts of otitis media, for which tympanostomy tubes had been placed. Three days before admission, in May 1997, tactile fever was noted, and one day before admission, the patient had decreased activity and rhinorrhea. On the day of admission, he vomited twice. In the emergency room, he had a temperature of 39.2°C, was irritable and lethargic, and had nuchal rigidity. A complete blood count showed a total leukocyte count (WBC) of 18,300 (69% segmented cells, 8% bands). Cerebrospinal fluid (CSF) analysis showed a WBC of 54 (25% segmented cells, 75% monocytes), protein 38 mg/dL, and glucose 70 mg/dL. The patient was hospitalized and initially treated with ceftriaxone. On the next day, he became less responsive, and abnormal posturing developed in the left upper and lower extremities. A computed tomography scan of the head (without and with contrast) was normal. He was transferred to a tertiary-care center, where an electroencephalogram showed moderate generalized slowing and recurrent right hemispheric electrographic seizures. A magnetic resonance imaging scan done on the same day showed a diffusely increased white matter signal consistent with viral encephalitis or acute disseminated encephalomyelitis. The patient received acyclovir for 3 days. His responsiveness and clinical condition gradually improved, and he was transferred to a rehabilitation service 17 days after admission. Initial work-up at the hospital revealed negative blood cultures and negative bacterial and viral cultures of the CSF. PCR for Epstein-Barr virus in the blood and CSF was negative, as was PCR for Herpes simplex virus in CSF.

The patient was enrolled in the UNEX project and evaluated. Specimens available for testing included acute- and convalescent-phase serum and CSF specimens. A variety of tests were conducted (see neurologic syndrome testing protocol in Appendix II available online at http://www.cdc.gov/ncidod/eid/vol8no2/pdf/01-0165-app2.pdf). Because the quantities of
specimens available were limited, testing was prioritized. First-round testing was negative for 
Cytomegalovirus, HHV-6, and arboviruses. However, testing for IgG antibodies (by IFA) for 
Epstein-Barr viral capsid antibodies showed a fourfold rise in titer between acute- and 
convalescent-phase serum specimens; testing for IgG antibodies (also by IFA) to Epstein-Barr 
early antigen revealed a fourfold decrease in titer between convalescent- and acute-phase serum 
specimens, indicating acute Epstein-Barr infection.