Evaluation of a Commercially Available Reverse Transcription-PCR Assay for Diagnosis of Enteroviral Infection in Archival and Prospectively Collected Cerebrospinal Fluid Specimens

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A commercially available reverse transcription (RT)-PCR method (AMPLICOR EV; Roche Diagnostic Systems, Inc., Branchburg, N.J.) was evaluated for detection of enteroviruses in cerebrospinal fluid from patients with neurological disease. This assay was compared with virus isolation in cell culture and an in-house RT-PCR method designed with a nonoverlapping region of the enteroviral genome. A panel of 200 cerebrospinal fluid specimens prospectively collected from patients with a wide variety of neurological symptoms, including 50 patients involved in three different outbreaks of acute aseptic meningitis, was assayed. A second panel of 97 archived cerebrospinal fluid specimens, stored for 2 to 5 years, from patients with aseptic meningitis associated with several enterovirus outbreaks was also studied. From the first panel, enteroviruses were detected in 13 of 50 specimens by cell culture (26%), in 43 of 50 specimens by AMPLICOR EV (86%), and in 46 of 50 specimens by the in-house assay (92%) from patients with aseptic meningitis associated with outbreak and 1 of 29, 3 of 29, and 4 of 29 specimens, respectively, from sporadic cases of aseptic meningitis. The remaining 121 cerebrospinal fluid specimens from patients with other neurological syndromes were negative by all tests. From the second panel, enteroviral RNA was detected by the AMPLICOR test (31 of 97 specimens, 32%) and the in-house assay (39 of 97 specimens, 40%). According to our results, patients with aseptic meningitis should be analyzed for enteroviral infection in cerebrospinal fluid by RT-PCR methods, and the AMPLICOR EV test is a suitable assay for diagnosis.

The enterovirus group includes 68 distinct serotypes of positive single-stranded RNA viruses which are human pathogens (poliovirus types 1 to 3, coxsackievirus groups A and B, echoviruses, and enteroviruses 68 to 71). Most enteroviral infections progress without clinical symptoms. However, enterovi-
um-high rates of enteroviral infection has not been validated (1, 20).

Several methods of enzymatic RNA reverse transcription (RT) followed by cDNA amplification (RT-PCR) have recently been introduced and used to obtain rapid diagnoses of enteroviral infection (3, 14, 15, 17, 22, 23). These new techniques showed greatly improved sensitivity compared to isolation in cell culture; however, typing of the virus strains, an important issue for epidemiological purposes, is not possible.

We have evaluated a commercially available RT-PCR test (AMPLICOR EV; Roche Diagnostic Systems, Branchburg, N.J.) for establishing the diagnosis of enteroviral infection of the CNS by comparison with isolation in cell culture and an in-house RT-PCR assay designed with a nonoverlapping region of the enteroviral genome. The latter methods were applied to CSF from patients with a variety of neurological symptoms for which lumbar puncture was routinely performed and to archival CSF samples from patients involved in several identified outbreaks of AM caused by enteroviruses.

MATERIALS AND METHODS

CSF specimens and patients. Two hundred consecutive CSF specimens, from the same number of patients with neurological symptoms for which lumbar puncture was routinely performed, were prospectively collected. After collection, 400 µl of each CSF specimen was immediately inoculated on appropriate cell lines for virus isolation. The remainder was subsequently aliquoted and frozen at −80°C for later detection of enteroviral RNA. For 106 specimens, the volume collected was not enough to perform the complete study. These specimens were diluted fourfold prior to aliquoting and culture.

Epidemiological data were highly suggestive of an acute enterovirus infection in 50 patients, because they were associated with three outbreaks of AM in Spain during the time of the study (November 1995 to May 1996). The remaining 150 patients presented with a wide variety of neurological symptoms, including 37 cases of neurological disorders associated with human immunodeficiency virus (HIV) infection and 7 cases associated with other causes of immunosuppression, 7 cases of suspected congenital infection, 29 cases of non-outbreak-associated or sporadic AM, and 29 cases of encephalitis; finally, a miscellaneous group of 41 patients presented with other neurological syndromes.

Ninety-seven archived CSF specimens, stored at −20°C for 2 to 5 years, from patients with AM who had been associated with several identified enterovirus outbreaks in Spain within a period of four years (1991 to 1994) were retrospectively selected and used. All of these specimens had previously been cultured, and the results of virus isolation tests were recovered from our records. Enteroviruses had been isolated in 22 of these CSF samples, including two echovirus type 4 (Echo-4), one Echo-7, three Echo-9, one Echo-11, one Echo-17, four Echo-18, one Coxsackievirus B6, and nine nonpoliovirus, untyped samples. The remaining 75 specimens were negative.

Cell culture and typing. Virus isolation was performed for each CSF sample (100 µl) on human embryo lung fibroblasts, human lung carcinoma cells (A549), THP-1 human monoblastic leukemia cells, and monkey kidney cells (LLC-MK2). Cells were grown at 37°C and observed daily for cytopathic effect during 15 days. If, after this time, no cytopathic effect had been observed, cultures were discarded. Enteroviruses isolated were typed by the standard method of virus neutralization (Lim-Banyesh-Melmuck immune serum pools).

AMPLICOR EV test. The AMPLICOR test was used according to the manufacturer’s instructions after a period of training and performance of a validation study with simulated specimens. In this way, the instruments and reagents involved in the PCR procedure, as well as the technicians performing the assay, were validated before processing of clinical specimens. The AMPLICOR EV test procedure has been described elsewhere (9). In short, viral RNA from specimens was extracted by mixing 100 µl of CSF with 400 µl of lysis solution, with incubation at room temperature for 10 min. RNA was precipitated by the addition of 500 µl of isopropanol and centrifugation at 16,000 × g for 10 min. The pellet was washed with 750 µl of 70% ethanol and resuspended in 200 µl of a guanidinium thiocyanate lysis buffer, which includes 100 molecules of a cold- and pure-formed genome fragment of enteroviruses DNA as an internal control for extraction and amplification procedures. As a first strand, DNA synthesis was performed by adding 1 µl of amplified product from the first reaction to 49 µl of a PCR mixture containing 60 mM Tris-HCl, 15 mM (NH4)2SO4, 2 mM MgCl2, 0.250 and 0.500, and negative if the OD450 was <0.250. However, in the commercially available kit, the equivocal range has been eliminated and the cutoff has been set at 0.350. PCR testing of each extracted sample was performed in duplicate, after amplification of the detection well per PCR (two PCR results were generated for each specimen). Both negative and positive control tubes were processed in each PCR run. The enzyme uracil-N-glycosylase (AmplErase; Roche), which recognizes and catalyzes the destruction of deoxuryridine-containing DNA, was included in the AMPLICOR master mix. This is a novel improvement designed to prevent false-positive amplification by carryover contamination (11).

The sensitivity of this test, as reported by Lina et al. (10) in a multicenter evaluation, ranged from 67 to 98% for viral titers of 1 to 10 50% tissue culture infective doses (TCID50)/0.1 ml but was only 16% for titers of 0.1 TCID50/0.1 ml. In-house RT-PCR assay. CSF specimens were also tested with the in-house RT-nested PCR assay, but only primers for enteroviruses and specific pseudorabies virus primers (as an internal control) were used in order to detect exclusively enteroviral RNA. Briefly, viral RNA from 50 µl of CSF specimen was extracted according to Casas et al. (4) by mixing it with 200 µl of a guanidinium thiocyanate lysis buffer, which includes 100 molecules of a cloned and purified genome fragment of pseudorabies virus DNA as an internal control for extraction and amplification procedures. Cells were incubated at room temperature for 5 min, and the sample was washed with 70% ethanol and disinfected with 70% ethanol and disinfected with 7% sterile double-distilled water. RT and the first PCR amplification of the target RNA were performed in a single reaction tube by using the Access RT-PCR System (Promega, Madison, Wis.). This system uses reverse transcriptase from avian myeloblastosis virus for first-strand DNA synthesis and the thermostable DNA polymerase from Thermus aquaticus (TF DNA polymerase) for DNA amplification. This simplifies the procedure and reduces the chance of contamination. After extraction, 5 µl of the dissolved pellet was added to 45 µl of an RT-PCR mixture composed of 2 mM MgSO4, 0.2 mM each deoxynucleoside triphosphate, 10 pmol of each downstream and upstream primer, 5 units of avian myeloblastosis virus reverse transcriptase, and 5 units of TF DNA polymerase; all of these reagents were used in a buffer compatible for both enzymes. Amplifications were carried out in a PTC-200 Peltier thermal cycle (MJ Research, Watertown, Mass.). Samples were subjected to an initial cycle of denaturation at 94°C for 1 min, annealing at 64°C for 1 min, and extension at 72°C for 1 min, followed by 45 cycles of 30 s at 94°C, 30 s at 55°C, and a final elongation at 72°C for 2 min. The amplified products were then performed by adding 1 µl of amplified product from the first reaction to 49 µl of a PCR mixture containing 60 mM Tris-HCl, 15 mM (NH4)2SO4, 2 mM MgCl2, 0.5 mM each deoxynucleoside triphosphate, 10 pmol of each sense and antisense primer, and 1.25 units of Taq polymerase (Perkin-Elmer). Amplification was carried out under the same conditions except for the annealing temperature (47°C), and only 30 cycles were performed. The nested amplification product was analyzed by electrophoresis through 2% agarose in a Tris-borate-EDTA gel stained with ethidium bromide (0.5 µg/ml).

The in-house PCR method has been shown to be highly sensitive (5), capable of detection of between 0.2 and 0.02 TCID50/0.1 ml with poliovirus type 1, coxsackievirus B1, coxsackievirus A16, Echo-4, Echo-9, and Echo-30.

Study design. All tests (cell culture, typing, AMPLICOR, and the in-house PCR) were performed at the Centro Nacional de Microbiologia laboratories, which receive clinical specimens from a high number of different hospitals all over Spain. The 200 prospectively collected CSF specimens were not specially selected for this study, but all CSF specimens sent to our laboratory for virological diagnosis were enrolled. Just after receipt, CSF specimens were cultured on appropriate cell lines and immediately aliquoted and frozen at −80°C for subsequent detection of enteroviral RNA. All AMPLICOR testing of each extracted sample was performed in duplicate; i.e., RNA from each extracted specimen was amplified twice and one detection per amplification was performed. In this way, each sample generated two PCR results. The in-house RT-PCR assay was performed in parallel on each CSF specimen. In order to avoid false-positive PCR results by carryover contamination, four distinct areas for reagent preparation, nucleic acid extraction and first amplification, nested PCR, and detection of products were established. Blinding of the study was guaranteed because cell culture, the AMPLICOR test, and the in-house assay were performed separately by different technicians without knowledge of the results obtained with the other assays.

Any discrepant result was resolved or reaffirmed by further PCR testing on a new frozen aliquot. Discrepant results included (i) discordant duplicate PCR results with the AMPLICOR test; (ii) concordant duplicate PCR results with the AMPLICOR test following between 0.250 and 0.500 absorbance unit (equivalent range); (iii) AMPLICOR PCR results for which the corresponding culture was discordant, i.e., PCR-negative and culture-positive or PCR-positive and culture-negative specimens; and (iv) specimens with discordant results by both PCR methods that were repeated by both PCR assays.

Statistical analysis. Comparisons between AMPLICOR test and cell culture isolation results were evaluated by McNemar’s test.
was positive for all tested specimens. Echo-9, Echo-11, Echo-30, coxsackievirus B6, and a nonpolio-terovirus types involved in these six specimens were Echo-4, specimens were also negative by the in-house assay. The en-
virus, untyped enterovirus. Three additional culture-positive
Echo-9, Echo-11, Echo-30 (four specimens), and a nonpolio-
failed to amplify with the AMPLICOR test. Enteroviruses iso-
positive. Moreover, six additional culture-positive specimens
Both RT-PCR methods yielded negative results for 51 speci-
39 of 97 specimens yielded amplification of enteroviral RNA
97 retrospectively collected specimens. Totals of 31 of 97 and
0.250 and 0.500 absorbance unit (equivocal range).
was positive in all tested specimens.
house PCR assay were discarded because the internal control
munocompromised patients. False-negative results with the in-
other than AM nor among the 44 specimens taken from im-
CSF specimens taken from patients with neurologic diseases
were found to be positive for enteroviral RNA by AMPLICOR
(10%) and four samples were found positive by the in-house
assay (14%). These four samples came from two infants, 1
month and 5 days old, and two adults, 33 and 49 years old.
All CSF specimens positive for enteroviruses by cell culture
were also positive by both RT-PCR tests. No positive results,
either by cell culture or RT-PCR, were obtained among the
CSF specimens taken from patients with neurologic diseases
other than AM nor among the 44 specimens taken from im-
munocompromised patients. False-negative results with the
in-house PCR assay were discarded because the internal control
was positive in all tested specimens.
There was a clear segregation between positive and negative
specimens with the AMPLICOR test, since none of the results
from the 200 prospective specimens analyzed fell between
0.250 and 0.500 absorbance unit (equivocal range).
Retrospective study. Table 2 shows the results obtained with
97 retrospectively collected specimens. Totals of 31 of 97 and
39 of 97 specimens yielded amplification of enteroviral RNA
by the AMPLICOR test and the in-house assay, respectively.
Both RT-PCR methods yielded negative results for 51 speci-
ments, including 3 specimens found previously by culture to be
positive. Moreover, six additional culture-positive specimens
failed to amplify with the AMPLICOR test. Enteroviruses iso-
lated from these nine specimens were typed as Echo-4, Echo-7,
Echo-9, Echo-11, Echo-30 (four specimens), and a nonpolio-
virus, untyped enterovirus. Three additional culture-positive
specimens were also negative by the in-house assay. The en-
terovirus types involved in these six specimens were Echo-4,
Echo-9, Echo-11, Echo-30, coxsackievirus B6, and a nonpolio-
virus, untyped enterovirus. False-negative results with the in-
house PCR assay were discarded, because the internal control
was positive for all tested specimens.
In order to emphasize the erratic nature of the results ob-
tained with archived CSF samples, Table 3 shows in detail the
results obtained for eight specimens with discordant duplicate
PCR results after the AMPLICOR test was performed in our
laboratory. These discordant results were resolved by repeated
AMPLICOR testing on new frozen aliquots sent to Roche
Molecular Systems (Somerville, N.J.).

RESULTS

Prospective study. The results obtained with the 156 CSF spec-
imens from immunocompetent patients included in the pro-
pective phase are summarized in Table 1. A total of 50 CSF
specimens from patients with AM associated with epidemic
outbreaks caused by enteroviruses were assayed. Thirteen of
them (13 of 50, 26%) gave positive results by cell culture and
both RT-PCR methods. Thirty additional cases were detected
by the AMPLICOR test; thus, a total of 43 positive specimens
(86%) were identified. All of these 43 positive samples were
also amplified, and therefore confirmed, by our in-house RT-
PCR, an alternative method which amplifies a nonoverlapping
region of the enteroviral genome. In addition, three further
cases were detected by the in-house assay (46 of 50, 92%). All
viral isolates were typed as Echo-30.
In only one of the 29 patients with sporadic AM was an en-
terovirus detected by cell culture and both RT-PCR methods;
it was typed as coxsackievirus B5. Two further cases were iden-
tified by both RT-PCR methods, and one additional case was
detected by the in-house assay. Thus, three CSF samples were
found to be positive for enteroviral RNA by AMPLICOR
(10%) and four samples were found positive by the in-house
assay (14%). These four samples came from two infants, 1
month and 5 days old, and two adults, 33 and 49 years old.
All CSF specimens positive for enteroviruses by cell culture
were also positive by both RT-PCR tests. No positive results,
either by cell culture or RT-PCR, were obtained among the
CSF specimens taken from patients with neurologic diseases
other than AM nor among the 44 specimens taken from im-
munocompromised patients. False-negative results with the
in-house PCR assay were discarded because the internal control
was positive in all tested specimens.

TABLE 1. Results of cell culture and RT-PCR tests for 156 CSF samples from immunocompetent patients included in the prospective study

<table>
<thead>
<tr>
<th>Neurologic syndrome</th>
<th>No. of CSF specimens</th>
<th>No. negative by all tests</th>
<th>Cell culture and both RT-PCR methods</th>
<th>Both RT-PCR methods</th>
<th>AMPLICOR RT-PCR</th>
<th>In-house RT-PCR</th>
</tr>
</thead>
<tbody>
<tr>
<td>Outbreak-associated AM</td>
<td>50</td>
<td>4</td>
<td>13</td>
<td>30</td>
<td>0</td>
<td>3</td>
</tr>
<tr>
<td>Sporadic AM</td>
<td>29</td>
<td>25</td>
<td>1</td>
<td>2</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>Other syndromes</td>
<td>77</td>
<td>77</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
</tr>
</tbody>
</table>

* All specimens which yielded positive results by culture were successfully amplified by both RT-PCR methods.
* Result(s) confirmed by testing of a new aliquot of CSF by both RT-PCR methods.

DISCUSSION

We have evaluated the AMPLICOR EV test to determine
the suitability of this method for detection of enteroviral RNA
in CSF specimens by comparing it with isolation in cell culture
and an in-house RT-PCR assay. Prospectively and retrospec-
tively collected clinical samples from a wide variety of syn-
dromes were assayed.

Prospective study. Despite testing of a broad spectrum of
neurological syndromes, including specimens from 29 patients
with encephalitis, 37 patients neurological disorders associated
with HIV infection and 7 patients with other causes of immu-
nosuppression, 7 patients with suspected congenital infection,
and 41 patients with other neurologic syndromes, enterovirus-
uses were detected only in specimens from 79 patients with AM.
Although some types of enteroviruses have been involved as
etiological agents in some cases of encephalitis (12), such cases
are uncommon and we did not find any positive results with
patients who presented with this disease. In addition, none of
the neurological disorders of the immunosuppressed patients
included in this study could be imputed to enteroviral infec-
tion; nevertheless, only one patient presented with gamma-
globulinemia, while the immunosuppression of the remaining
patients was caused by HIV infection or anticonvogenic treat-
ment. Several authors have suggested previously that enterov-
iral infection is probably an important cause of neurological
disease in patients with antibody deficiencies (18, 19).

Fortunately, we possessed 50 CSF specimens from patients
involved in enteroviral AM epidemics; therefore, a reliable
comparison between isolation in cell culture and RT-PCR
techniques could be done. Only 13 of 50 specimens (26%) yiel-
ded enteroviral growth in cell culture, while 43 of 50 (86%)

TABLE 2. Results of cell culture isolation and RT-PCR methods
for CSF specimens from the retrospective study

<table>
<thead>
<tr>
<th>AMPLICOR test result</th>
<th>Cell culture isolation</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>No. of positive specimens (n = 22)</td>
</tr>
<tr>
<td></td>
<td>with in-house RT-PCR</td>
</tr>
<tr>
<td>+</td>
<td>10</td>
</tr>
<tr>
<td>-</td>
<td>6</td>
</tr>
</tbody>
</table>
and 46 of 50 (92%) were positive for enteroviral RNA by the AMPLICOR test and the in-house RT-PCR assay, respectively. All of the culture-positive specimens were successfully amplified by both PCR methods, so no false-negative results were obtained. False-positive results with the AMPLICOR test were ruled out because all positive specimens were also amplified, and therefore confirmed, with the alternative RT-PCR assay, which amplifies a nonoverlapping region of the enteroviral genome. These data show that the AMPLICOR test is more sensitive by far than cell culture of CSF for diagnosis of enteroviral meningitis ($P = 0.22$ [McNemar’s test, Yates corrected]).

Three additional specimens that were positive by the in-house assay could not be amplified by the AMPLICOR test. These three samples were each assayed again with a new frozen aliquot. Although the initial volume of CSF used in the AMPLICOR test is double the volume used in the in-house assay, the amount of target RNA put into the amplification reaction mixture is the same for both PCR methods.

Possible reasons for the low rate of enteroviral detection by cell culture (13 of 50 specimens, 26%) compared with results obtained by RT-PCR methods could be either a low number of infectious particles in CSF specimens or numerous replication-defective or antibody-neutralized viruses which cannot be propagated in cell culture (13). Nevertheless, it is possible that previous dilution of some specimens decreased the sensitivity of cell culture. If we considered only the specimens from outbreak-associated AM cases that were tested undiluted (20 of 50), the percentage of positive specimens in cell culture would rise to 45% (9 of 20), which is still far below the 90% (18 of 20) positive specimens with the AMPLICOR test ($P = 0.55$ [Mc Nemar’s test, Yates corrected]) or 95% (19 of 20) with the in-house assay. Prior studies comparing cell culture and PCR assays for enteroviral detection in prospectively collected CSF samples from patients with AM have demonstrated that PCR is more sensitive than viral cultures. The rates of enterovirus detection by Yerly et al. for 38 patients whose specimens were obtained at intervals of enteroviruses (10, 1, or 0.1 TCID$_{50}$/0.1 ml).

Among our 29 patients with sporadic AM, the percentage of positive specimens detected by RT-PCR (3 of 29 [10%] with the AMPLICOR test and 4 of 29 [14%] with the in-house assay) was significantly lower than that found among the 50 outbreak-associated AM patients in this study or those reported in the studies cited above (23, 24). Nevertheless, neither of these two previous reports specified the origins of the CSF specimens as being from sporadic cases of AM or from cases involved in outbreaks, a well-established distinction in our study. Other epidemiologic factors, such as seasonal variations in the circulation of enteroviruses, year-to-year variations in the incidence of enteroviral infections, and cocirculation of other infectious agents causing AM at the time of the study, might also explain these differences. Note that based on the clear segregation of positive and negative results with the AMPLICOR test, in the commercially available kit the equivocal range has been eliminated and the cutoff has been set at 0.350.

**Retrospective study.** The analyses done on archived CSF specimens were less successful than those from prospectively collected samples. Only 31 of 97 specimens (32%) yielded a positive result by the AMPLICOR test; 39 of 97 (40%) did so by the in-house assay. In addition, both RT-PCR methods failed to detect three specimens which had been positive by culture, six specimens were AMPLICOR negative and culture positive, and three specimens were in-house negative and culture positive, for a total of 12 false-negative specimens. After typing, none of these 12 enteroviruses were found to be Echo-1, Echo-5, Echo-22, or Echo-23, four enteroviral types often missed by the AMPLICOR test (9). Moreover, the in-house PCR assay has previously been shown to be highly sensitive in detecting the enterovirus types involved in these false-negative results (5). Therefore, low sensitivity for detecting particular types of enteroviruses should be discarded as an explanation for these findings. The lack of reproducibility of the results obtained by the PCR assays in some of the archived samples (see Table 3) suggests that degradation of enteroviral RNA, caused by freezing-thawing and long-term storage of specimens at $-20^\circ C$ (8), likely accounts for the low sensitivity of the PCR tests in the retrospective study and indicates that archived CSF samples are not suitable for evaluation of the performance of such tests in diagnosis. Nevertheless, Rotbart et al. (16) reported high sensitivity (94.7%) and specificity (97.4%) for the AMPLICOR test in a study performed with archival CSF specimens stored at $-70^\circ C$.

In conclusion, the present study indicates that RT-PCR is a powerful tool for the diagnosis of AM syndromes due to enteroviral infection and shows that the AMPLICOR EV test is a reliable and standardized method for rapid and sensitive detection of enteroviruses in CSF. Long-term storage of enterovirus-containing CSF specimens is likely to lead to enteroviral RNA degradation that renders the specimen unsuitable for further testing. Archival CSF samples should not, therefore, be used for evaluation of PCR assays designed to detect enteroviral RNA in human CSF.

### ACKNOWLEDGMENTS

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### TABLE 3. Discordant duplicate PCR results by the AMPLICOR test with archived CSF specimens

<table>
<thead>
<tr>
<th>CSF specimen no.</th>
<th>AMPLICOR result(s) (OD$_{450}$)</th>
<th>In-house RT-PCR result</th>
<th>Cell culture result</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>1st assay</td>
<td>2nd assay</td>
<td>RMS$^a$</td>
</tr>
<tr>
<td>1</td>
<td>0.137, 0.414</td>
<td>0.198, 0.917</td>
<td>0.726</td>
</tr>
<tr>
<td>2</td>
<td>0.175, 0.798</td>
<td>0.213, 0.987</td>
<td>2.824</td>
</tr>
<tr>
<td>3</td>
<td>0.161, 0.332</td>
<td>0.158, 0.580</td>
<td>0.105</td>
</tr>
<tr>
<td>4</td>
<td>0.133, 0.135</td>
<td>0.155, 0.346</td>
<td>0.950</td>
</tr>
<tr>
<td>5</td>
<td>0.131, 0.544</td>
<td>0.151, 0.484</td>
<td>0.061</td>
</tr>
<tr>
<td>6</td>
<td>0.129, 0.518</td>
<td>0.159, 0.330</td>
<td>1.234</td>
</tr>
<tr>
<td>7</td>
<td>0.110, 0.127</td>
<td>0.184, 0.604</td>
<td>0.053</td>
</tr>
<tr>
<td>8</td>
<td>0.187, 0.521</td>
<td>0.150, 0.411</td>
<td>3.008</td>
</tr>
</tbody>
</table>

$^a$ Positive, >0.500; indeterminate, 0.250 to 0.500; negative, <0.250. Each specimen was assayed in duplicate (two microwell plate results) in the first and second assays with the AMPLICOR test. Results obtained by Roche Molecular Systems, Somerville, N.J.
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