

Intravital Diagnosis of Human Rabies by PCR Using Saliva and Cerebrospinal Fluid

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An optimized reverse transcription (RT)-PCR protocol for the intravital detection of rabies virus genomic RNA was tested with clinical samples obtained from 28 patients suspected of having rabies, 9 of whom were confirmed to have had rabies by postmortem examination. RT-PCR using saliva combined with an immunofluorescence assay performed with skin biopsy samples allowed detection of rabies in the nine patients.

According to results of global surveillance by the World Health Organization, about 50,000 cases of human rabies occur each year (45), the majority of them in developing countries (63). Human rabies can be prevented through the combination of stringent animal vaccination, quarantine programs, and the availability of expensive vaccines and specific immunoglobulins (32). In spite of the fact that most developed countries have good public health measures in place to prevent human rabies, these countries are dealing with an increasing number of human rabies cases. This is clearly the case in the United States, where bat rabies virus variants as well as some imported cases of human rabies have recently achieved an increased public health significance (41, 51, 53, 56). This is also the case in France, where seven human cases were recorded from 1988 to 1997; only eight cases had been recorded in the preceding 20 years. All of the human rabies cases in France were imported. A similar situation has also been observed throughout the whole European Union. Of the 15 cases reported in the last 10 years, 12 were imported from Asia, Africa, and Latin America (46; unpublished data from France). These cases underscore the importance of alerting travellers of the risk of rabies contamination and of the prophylactic methods to prevent the disease (42). They also indicate the importance of maintaining a good system of surveillance, of keeping medical staffs well informed on clinical presentation of a disease which rarely occurs in developed countries, and of the necessity to develop diagnostic tools for the identification of rabies in these patients.

The clinical diagnosis of rabies is sometimes suggested by epidemiological (history of exposure) and clinical (e.g., paresthesia, hydrophobia) findings (36). However, the disease is often mistaken for other disorders (30). Differentiation from other neurologic diseases may require extensive investigations. Therefore, diagnosis is often confirmed late in the course of the disease or postmortem (31). Delays in diagnosis greatly increase the number of contacts that require postexposure prophylaxis. The average number of contacts (hospital personnel, family) receiving postexposure treatments (PET) is approximately 50 per case ($n = 19$) in France and between 41 and 55 per case in the United States (29, 35). In the United States, one case resulted in 209 PET (50), and 290 PET for one case

were reported recently in France. The early diagnosis of rabies is also essential to eliminate the expense and discomfort of unnecessary diagnostic tests and inappropriate therapy.

A wide variety of viruses, bacteria, and parasites, all of which are capable of causing aseptic meningitis and encephalitis, have been detected by PCR (39, 49, 52, 58). The objective of the present study was to establish a reverse transcription (RT)-PCR protocol for use in evaluating diagnostic specimens, including saliva and cerebrospinal fluid (CSF). CSF samples were centrifuged at $11,000 \times g$ for 20 min at 4°C. Total RNA was extracted from specimens of saliva and pellets of CSF by four different techniques, including the following: (i) proteinase K (34), (ii) guanidinium thiocyanate together with silica particles (5, 6), (iii) cationic surfactant (Catrimox-14; Iowa Biotechnology Corporation) (43), and (iv) chelating resin (Chelex 100; Bio-Rad) (60). The proteinase K method was performed as follows. Briefly, 200 μ l of biological fluid (saliva or CSF including the pellet) was incubated for 2 h at 37°C with 400 μ l of proteinase K buffer containing 40 μ g of proteinase K (Gibco BRL). The RNA was then purified by a phenol-chloroform extraction and precipitated in absolute ethanol. The pellets were resuspended in 100 μ l of pyrolyzed water. Considering that the N gene is the most conserved in the lyssa viruses (except some domains of the L protein gene) and that the sequence data concerning this gene are the most exhaustive, we used primers in the N gene that were shown to allow amplification of a wide range of genetically diverse lyssa viruses (7, 40). One microliter of primer N12 (5' GTAACACCTCTA CAATGG 3', positions 57 to 74; all the positions of the primers are given based on the PV strain sequence [59]) (100 ng/ μ l) was incubated with 2 μ l of RNA (1 μ g) at 65°C for 3 min and chilled on ice. Each tube was then incubated at 37°C for 90 min with 4 μ l of a solution containing each nucleotide triphosphate (10 mM), 0.6 μ l of RNasin (15 U), 1 μ l of dithiothreitol, 2 μ l of Super Script reverse transcriptase buffer (Gibco BRL), and 0.5 μ l of Super Script reverse transcriptase (100 U; Gibco BRL). The RNA-cDNA hybrids were diluted 10 times in Tris-EDTA buffer. Five microliters of diluted cDNA was mixed with 50 μ l containing a 1 μ M concentration of N12 and N40 (5' GCTTGATGATTGGAAGT 3', positions 1368 to 1349), 20 mM each nucleotide triphosphate, 10 mM Tris-HCl (pH 8.3), 50 mM KCl, 3 mM MgCl₂, and 2 U of *Taq* polymerase (Gibco BRL). PCR was performed with a GeneAmp PCR System 9600 (Perkin-Elmer) by using the following program: 1 cycle of denaturation at 94°C for 60 s, annealing at 50°C for 90 s, and elongation at 72°C for 90 s; 30 cycles of denaturation for 50 s;

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TABLE 1. Patients and specimens analyzed

Patient	No. of specimens from:			Total no. of specimens	Diagnosis for rabies ^a		
	CSF	Saliva	Bronchoalveolar washings		Postmortem	Intravital by IF assay of skin biopsy sample	Intravital by RT-PCR using saliva sample
P1	1			1	-	-	-
P2	1			1	-	-	-
P3	1			1	-	-	-
P4		1		1	+	-	+
P5	2	2		4	+	-	+
P6		1		1	-	-	-
P7	9	11	1	21	+	-	+
P8	2			2	-	-	-
P9	1			1	-	-	-
P10	1			1	-	-	-
P11	2	1		3	-	-	-
P12		1		1	-	-	-
P13	1	1		2	+	+	-
P14		1		1	-	-	-
P15	1			1	-	-	-
P16	1			1	-	-	-
P17		2		2	-	-	-
P18	1	1		2	+	+	-
P19	2	3		5	+	+	-
P20	3	4		7	+	+	-
P21	1	1		2	+	-	+
P22	1			1	-	-	-
P23	1			1	-	-	-
P24	1			1	-	-	-
P25	2			2	-	-	-
P26	1			1	-	-	-
P27	2	1		3	-	-	-
P28	3	13		16	+	+	+
Total	41	44	1	86			

^a +, positive; -, negative.

and a final cycle of elongation for 5 min. The optimum temperature for annealing and magnesium ion concentration for the PCR was determined by standard titration experiments of a cDNA of the N gene of strain 9147FRA. Five microliters of PCR products was diluted into 95 μ l of pyrolyzed water, denatured for 3 min at 95°C, and chilled on ice. The samples were then distributed onto a Hybond nylon transfer membrane (Amersham) with a Bio-Dot microfiltration unit (Bio-Rad) and fixed by irradiation on a UV transilluminator. Primer set N3 (5' GTCTCTTTGAAGCCTGAG 3', positions 113 to 130)-N23 (5' GGTCTCTCGTCAGTTCCAT 3', positions 464 to 446) and primer set N17 (5' TTCTTCCACAAGAAGCTTTG 3', positions 848 to 866)-N2 (5' CCCATATAGCATCCTAC 3', positions 1030 to 1013) were used to generate two digoxigenin-labelled probes by PCR. Hybridization with the two complementary digoxigenin-labelled probes and detection by chemiluminescence were performed by using a nonradioactive DNA labelling and detection kit (Boehringer).

The four different methods of RNA extraction were tested in parallel, each on two independent series of experimentally infected samples, to determine the threshold of detection of the RT-PCR. Briefly, negative specimens of saliva and CSF were pooled and infected with serial dilutions of a fox rabies isolate (isolate 9147FRA) (40). This virus was produced on BHK-21 cells infected at a multiplicity of infection of 0.1 and harvested 48 h after infection to limit the number of defective viral particles. The Catrimox-14 and Chelex 100 techniques produced negative results. In our hands, poor results were achieved when guanidinium thiocyanate was used together with silica particles (threshold of detection, around 10⁴ focus-

forming units [FFU]/ml). However, the results obtained with the proteinase K method were reproducible with a threshold of detection of 10² FFU/ml in the saliva and 10 FFU/ml in the CSF. These results are similar to the theoretical viral infectivity obtained by rapid tissue culture infection test (RTCIT) (20 FFU/ml) (9). The difference in the results for saliva and CSF can be explained by the presence of RNase in the saliva.

Forty-one specimens of CSF and 44 specimens of saliva collected daily by aspiration or by use of cotton swabs were obtained from 28 patients with manifestations of aseptic meningitis, encephalitis, and suspected rabies infection (Table 1). Rabies was confirmed in the brain tissue of nine of these patients by postmortem examination (8, 28). One patient each was infected in the Middle East, Latin America (38), and India, and six patients were infected in Africa (18). One of these patients (patient P13) was known to be infected with human immunodeficiency virus (1). The day of onset of illness was defined as the day when the first symptom attributed to rabies could be noted. These data are missing for two patients. The average duration of illness was 12.4 days for the seven remaining rabies-infected patients. The specificity of RT-PCR was one for the samples of saliva and CSF. Eleven of the 37 putative positive specimens of saliva (sensitivity, 0.30) and only 2 of 22 specimens of CSF (sensitivity, 0.09) were confirmed positive (Table 2). The bronchoalveolar washing was negative. Intravital diagnosis performed by RT-PCR identified the presence of rabies virus in the saliva of five patients (sensitivity for the patient, 0.56) and in the CSF of only two patients (sensitivity for the patient, 0.22). RT-PCR performed with saliva sam-

TABLE 2. Frequency of intravital diagnosis of rabies in nine patients in France (1986–1997)

Specimen type	Method(s) ^a	Frequency of rabies diagnosis ^b					Sensitivity ^d	
		0–4 days ^c	5–8 days	9–12 days	13–16 days	>16 days		
Serum	Serology (ELISA, RFFIT)	0/4	1/2	0/2		1/1	0/3	0.17 (<i>n</i> = 12)
CSF	Serology (ELISA, RFFIT)	0/4	0/1	0/3		1/1	0/3	0.08 (<i>n</i> = 12)
	RT-PCR	0/5	1/3	0/2	0/1	0/2	1/9	0.09 (<i>n</i> = 22)
Saliva	RREID	0/1	0/5	0/4	0/4	1/4	0/1	0.05 (<i>n</i> = 19)
	RT-PCR	1/5	3/8	1/5	0/4	3/6	3/9	0.30 (<i>n</i> = 37)
Skin biopsy	IF	3/3	2/2			0/1	1/1	0.86 (<i>n</i> = 7)

^a RFFIT, rapid fluorescent focus inhibition test; RREID, rapid rabies enzyme immunodiagnosis for rabies antigen detection.

^b The number of patients diagnosed with rabies relative to the total number is given.

^c Days after onset of symptoms.

^d *n* = number of samples.

ples is of particular help in early diagnosis of the disease (Table 2).

Patient P7, who experienced a long clinical period, was confirmed positive for rabies at the time of death. Specimens of saliva from patient P7 were confirmed positive consecutively on days 21 and 22 of the disease, became negative on day 23, and became positive again on day 24. To investigate the relationship between the intermittence of rabies virus excretion and the time of day of sample collection, 11 specimens of saliva were collected every 2 or 4 h from one patient (patient P28) during a 5-day period midway through the course of the disease. This period starts 2 days after the last collection of saliva which was found positive, 9 days after the onset of the symptoms, and 12 days before death. None of these specimens were confirmed positive. The question of titer and duration of virus shedding is of particular importance for rabies diagnosis because saliva remains the major source of exposure in contacts for whom rabies prophylaxis is recommended (35). The signal obtained after RT-PCR using some saliva samples indicated that the level of rabies virus in the saliva could reach more than 10³ FFU/ml (data not shown). This evidence confirms that any contact with saliva from a patient suspected of having rabies should be taken seriously.

Most of the conventional techniques used for postmortem analysis of the brain are of limited value to support the intravital diagnosis of rabies (9, 37, 61). We reviewed records of 39 cases of intravital diagnosis of rabies in the United States during the period 1960 to 1996 (3, 10–27) and of 16 cases of intravital diagnosis performed in France during the period 1970 to 1997 (Table 3). This confirms that the corneal smear first developed with mice by Schneider (54) was too insensitive for accurate clinical diagnosis (3, 4, 44, 62). The only test that has demonstrated reliable results is the immunofluorescence (IF) test on skin biopsy samples (62). In our study, the IF test (33), performed on frozen sections of the skin biopsy samples,

exhibited the highest sensitivity (sensitivity, 0.86; *n* = 7). It detected the presence of rabies virus very early in the course of the disease and could be considered one of the most important tests for intravital diagnosis. In routine laboratory testing, we noticed that the examination of a minimum of 20 sections was needed to ensure the observation of several hair follicles. Typical rabies nucleocapsid inclusions can be observed in the nerve endings around the base of some of these hair follicles (data not shown). RT-PCR assay of saliva and IF assay of skin sections together allowed a positive intravital diagnosis in all of the nine laboratory-confirmed cases of human rabies presented in this study. The average delay between the onset of clinical symptoms and the collection of specimens that confirmed the presence of rabies virus was 6.7 days (*n* = 7). The sensitivities of detection of rabies antibodies by enzyme-linked immunosorbent assay (ELISA) (48) and by seroneutralization on cell culture (57) were very low in serum as well as in CSF (results obtained from the sera and CSF of patients P13 and P28, who had received a PET protocol different from that recommended by the WHO Expert Committee on Rabies [64], were excluded) (Table 2). This confirms that serological testing is of limited value because seroconversion occurs late in the course of the disease (55) (Table 3). The detection of rabies antigen in the saliva by ELISA (47) also gave poor results.

On the basis of these results, we propose a simple testing protocol for intravital diagnosis of rabies. Two different specimens should be sent to the laboratory in the early phase of the disease, a skin biopsy specimen and a saliva specimen. Skin biopsy samples should be analyzed by direct IF and saliva samples should be analyzed by the RT-PCR method described above. Then, after 1 week of illness, serum samples should also be examined for the presence of rabies-specific antibodies. RT-PCR has also been proven to be a valuable adjunct to an epidemiological investigation because sequences can be ob-

TABLE 3. Results of laboratory confirmation of rabies attempted in 55 patients during life^a

Determination	Frequency of rabies diagnosis ^b					Sensitivity ^d
	0–4 days ^c	5–8 days	9–12 days	13–16 days	>16 days	
Antigen in skin biopsies	5/6	6/10	5/5	1/2	1/3	0.69 (<i>n</i> = 26)
Antibodies in serum	1/15	5/28	8/25	14/20	12/14	0.39 (<i>n</i> = 102)
Antibodies in CSF	0/7	0/12	1/10	3/8	6/10	0.21 (<i>n</i> = 47)
Antigen in corneal smears	0/4	1/16	3/16	2/7	2/16	0.14 (<i>n</i> = 59)

^a These results were obtained from the analysis of 39 cases of rabies in the United States (3, 10–27) and 16 cases in France (1970–May 1997) (unpublished results).

^b The number of patients diagnosed with rabies relative to the total number is given.

^c Days after onset of symptoms.

^d *n* = number of samples.

tained rapidly from PCR products, thus permitting rapid identification of the virus (2, 40).

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