Enterovirus and herpesviridae family as etiologic agents of lymphomonocytary meningitis, Southern Brazil

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ABSTRACT

Viral meningitis is a common infectious disease of the central nervous system (CNS) that occurs worldwide. The aim of this study was to identify the etiologic agent of lymphomonocytary meningitis in Curitiba, PR, Brazil. During the period of July 2005 to December 2006, 460 cerebrospinal fluid (CSF) samples with lymphomonocytary meningitis were analyzed by PCR methodologies. Fifty nine (12.8%) samples were positive. Enteroviruses was present in 49 (83%) samples and herpes virus family in 10 (17%), of these 6 (10%) herpes simplex virus, 1 (2%) Epstein Barr virus, 2 (3%) human herpes virus type 6 and 1 (2%) mixed infection of enterovirus and Epstein Barr virus. As conclusion enterovirus was the most frequent virus, with circulation during summer and was observed with higher frequency between 4 to 17 years of age. PCR methodology is an important method for rapid detection of RNA enterovirus and DNA herpesvirus in CSF.

Key words: aseptic meningitis, lymphocytic meningitis, lymphomonocytary meningitis, cerebrospinal fluid, central nervous system, enterovirus, herpesvirus.

Enterovirus e família herpesviridae como agentes etiológicos de meningites linfomonocitárias, no sul do Brasil

RESUMO

A meningite viral é uma síndrome infecciosa comum do sistema nervoso central (SNC), que ocorre no mundo inteiro. O objetivo deste estudo foi identificar o agente etiológico de meningite linfomonocitária em Curitiba, PR, Brasil. Durante o período de julho de 2005 a dezembro de 2006, 460 amostras com meningite linfomonocitária foram analisadas por metodologias de PCR. Cinquenta e nove (12,8%) amostras foram positivas. Enterovirus estava presente em 49 (83%) amostras e herpes vírus em 10 (17%), destas 6 (10%) HSV, 1 (2%) EBV, 2 (3%) HHV- 6 e 1 (2%) infecção mista de enterovirus e EBV. Conclui-se que o enterovirus foi o vírus mais frequente, com a circulação durante o verão. Houve maior número de amostras positivas entre 4 a 17 anos. A metodologia de PCR é um importante método para a detecção rápida de RNA de enterovirus e DNA do herpesvirus no LCR. Palavras-chave: meningite asséptica, meningite linfocítica, meningite linfomonocitária, líquido cefalorraquidiano, sistema nervoso central, enterovirus, herpesvirus.

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Viral meningitis is a common infectious disease of the central nervous system (CNS) that occurs worldwide, especially during the summer. Almost 90% of cases are caused by enterovirus such as, coxsackievirus and echovirus which contain several serotypes¹. They are easily transmitted by direct contact with respiratory secretions that can cause epidemic outbreaks with high financial impact and morbidity rates. These viruses are typically geographically confined and some serotypes may arise in endemic regions, with gradual changes in the predominant serotypes present year to year 1,2. Enterovirus meningitis is the most common agent identified in most viral meningitis cases, followed by herpesviridae family viruses³. Viral meningitis is responsible for approximately 50% of inpatients with meningitis in the city of Curitiba, Southern Brazil⁴.

The number of notified cases of meningitis from all etiologies (bacterial, viral and not otherwise specified) in Curitiba from 1995 to 2003 was from 612 to 896. It is estimated that 40% of the meningitis cases were caused by viruses and the coefficient for the incidence of viral meningitis was around 29 cases/100,000 inhabitants⁴. There is little information on the etiology of viral meningitis in Brazil especially in the Southern region. In this study we identified the etiologic agent of viral meningitis that occurred in Curitiba, the capital of Paraná State, in Southern Brazil, in the July 2005 to December 2006 period using biology molecular approaches.

METHOD

The city of Curitiba is located in the southern Brazil at an altitude of 934.6 m, and has subtropical climate. It is the seventh largest city in Brazil with a population of around 1,850.000 inhabitants and a population density of 4,202.83 hab/km. In the period of July 2005 to December 2006 a total of 671 cerebrospinal fluid (CSF) samples were collected from patients with clinical suspicion of lymphomonocytary meningitis seen at three different reference hospitals in Curitiba (Hospital Universitário Evangélico, Hospital Pequeno Príncipe and Hospital de Clínicas-UFPR). CSF samples were stored in a RNA/ DNAse free and sterile tube and sent to the virology laboratory for less than 24 hours, on ice. There was a label on each sample containing epidemiological and clinical data the samples were stored at -70°C until molecular analysis.

Inclusion criteria – CSF samples were included in the study based on the clinical criteria of meningitis: acute onset (<3 weeks), presence of fever (>38°C), meningo-radicular signs and headache. CSF biochemical and cytological pattern of viral meningitis are: WBCs ≥5 cells/mm³ with predominance of lymphocytes, CSF glucose normal (>45 mg/dL), CSF lactate <3.5 mmol/L⁵.

Exclusion criteria – [1] Clinical diagnosis of encephalitis, defined as acute onset (<3 weeks), the presence of fever (>38°C) and signs or symptoms that suggested brain parenchyma involvement (consciousness and/or personality alterations, seizures or focal neurological signs)⁶; [2] Samples with CSF biochemical and cytological characteristics of bacterial meningitis: WBCs >5 cells/mm³ with predominance of neutrophils, low CSF glucose normal (<45 mg/dL), CSF lactate >3.5 mmol/L; [3] Samples improperly stored, non-refrigerated samples or sent improperly to the virology section; [4] Identification in CSF of an etiologic agent other than virus.

CSF total protein was quantified by the sulphosalicylic acid turbidimetric method and CSF glucose by the enzymatic method. CSF total cell count (WBCs) was assessed using a Fuchs Rosenthal chamber. For differential cell count CSF samples were concentrated in a cytospin and the slides stained by the May Grünwald-Giemsa technique.

The study was approved by the Ethical and Research Committee on Humans Beings from the Hospital de Clínicas, UFPR.

Extraction of viral RNA/DNA

The extraction of viral RNA or DNA was followed by a previously described protocol⁷. Briefly, 200 μL of lysis buffer (Guanidine Isothyocianate (GuSCN), Invitrogen (USA) 4 M, 0.5% of N-Lauroylsarcosine salt solution, Fluka (USA), 1 mM dithiotreitol (DTT), Invitrogen (USA), 25 mM sodium citrate, Sigma (USA), 20 μg/tube of glycogen, Sigma (USA), 100 copies of plasmid with pseudo rabies virus (PRV), used as internal control were added to $50 \mu L$ of CSF. After, vortexed tube was incubated room temperature for 10 minutes. 250 µL of isopropyl alcohol (-20°C) was added, vortexed and centrifuged for 10 minutes, 17,000 rpm at 4°C. Pellet was washed with 500 µL of ethanol 70, centrifuged as described. Supernatant was discarded and the open tube incubated at 56°C for 10 minutes. The dried pellet was resuspended in ultrapure water and stored at -70° C.

Reverse transcription

Viral RNA was transcribed to obtain cDNA as described elsewhere 8 . The master mix was prepared in a volume of 17.5 $\mu L/tube$. The mix contained dideoxynucleotides triphosphates 2.5 mM (dNTPs), 5X buffer (Invitrogen, USA), dithiothreitol (DTT) 0.1 M (invitrogen, USA), 0.25 μL of RNAse Out (Invitrogen, USA) and transcriptase enzyme Superscript II (Invitrogen, USA) 0.25 μL of the enterovirus antisense primer was added to 10 μL of the extracted product. After 2 minutes at 94°C ice cooling, 17.5 μL of the mix was added to the tube, which was stored at 45°C for 1 hour.

Amplification - PCR

Enterovirus – PCR for enterovirus was based on the amplification of the 5'-UTR region of the gene, which is a highly conserved region in most enterovirus serotypes according to Casas et al.9. For the first PCR the primer sequences were: EV1- Reverse 5'GAAACACG-GACACCAAAGTAGTCG3' and EV1 - Forward 5'CG-GTACCTTTGTRCGCCTGTTTTA3'. The amplification consisted of two rounds. For the first PCR the master mix contained 10 mM Tris-HCL (pH 8.8), 3.5 mM MgCl₂, 2.5 mM KCl, 1.5 U of Taq polymerase and 0.5 μL of each external primer in a concentration of 10 pmol, the final volume was adjusted with ultra pure water to $22.5~\mu L$ and $2.5~\mu L$ of cDNA. The amplification was performed with an Eppendorf thermocycler in one cycle of 94°C for 2 minutes, followed by 40 cycles: 94°C for 30 seconds, 48°C for 30 seconds and 72°C for 2 minutes, after extension on 72°C for 10 minutes. For the second PCR the primer sequences were as follows: EV2 - Reverse 5'GGATTAGCCGCATTCAGGG3'; EV2 - Forward 5'CAAGCACTTCTGTTTCCCCG3'. Master mix for the second PCR was described above and the volume of Tag polymerase was 0.25 µL. 0.5 µL from the first amplification product and 24.5 µL from the mix were used. The second amplification was performed with an Eppendorf thermocycler in one cycle of 94°C for 2 minutes, followed by 30 cycles of 94°C for 30 seconds, 52°C for 30 seconds, 72°C for 1 minute, after an extension on 72°C for 10 minutes. Amplification products were analyzed in 1% agarosis gel. The expected product for enterovirus was of 306-316 pb and for the internal control, PRV 147 pb.

Herpesviridae family – The herpesviruses were identified by the methodology described by Johnson et al.10, which was standardized to detect eight species of human herpesviruses. Primers were designed for a well conserved region of the DNA polymerase gene. The first pair of primers (PAN1) was used to amplify herpes simplex virus type 1 (HSV-1), herpes simplex virus type 2 (HSV-2), Epstein Barr virus (EBV), Cytomegalovirus (CMV), and human herpes virus type 8 (HHV-8) and had the following sequences: HSV P1 - 5'-GTGGTG-GACTTTGCCAGCCTGTACCC-3' and HSV-P2 -5'-TAAACATGGAGTCCGTGTCGCCGTAGATGA-3'. The primer pair 2 (PAN2) was used to amplify varicella zoster virus (VZV), human herpes virus type 6 (HHV-6) (variant A or B), and HHV-7 presented the following sequence VZV-P1 - 5'-TCGTGTTTTGATTTTCAAAGTT-TATATCC-3' and VZV-P2 5'-ATAAACACACAATCC-GTATCACCATAAATAACCT-3'. Polymerase Chain Reaction was performed in a total volume of 25 µL. The mix contained 2.5 µL of 10X buffer (Invitrogen, USA), 0.75 µL magnesium chloride (MgCl₂) 50 mM (Invitrogen), 8 μL dNTPs 1.25 mM (Invitrogen), 1.25 μL of dimethyl sulfoxide (DMSO) (Merck), 1.75 µL of each primer in a concentration of 37.5 pmol, 0.35 µL de Taq polymerase (Invitrogen), ultra pure water and 2,5 μL from DNA. The amplification was performed with an Eppendorf cycler. For the first pair of primers the cycle was as follows: 1 cycle 95°C for 12 minutes, 3 cycles of 95°C for 1 minute, 60°C for 1 minute, 72°C for 1 minute, 37 cycles of 95°C for 1 minute, 55°C for 45 seconds, 72°C for 1 minute, 1 cycle 72°C for 3 minutes. For the second pair of primers the cycles were: 1 cycle 95°C for 12 minutes, 40 cycles of 95°C for 1 minute, 47°C for 1 minute, 72°C for 1 minute, 1 cycle 72°C for 3 minutes. The amplification products were analyzed by electrophoresis using 1% agarosis gel. The expected product for PAN 1 was 532 bp and for PAN 2 536 bp. For each reaction positive and negative samples as well as water were used as controls.

Restriction enzyme digestion

Samples with products of approximately 532 or 536 bp were selected for enzyme digestion. Each reaction mixture in which amplicons were detected was subjected to digestion with the restriction enzymes BamHI and BstUI (New England Biolabs, USA). The digestion mixture consisted of 10 μ L of PCR mixture, 1.5 μ L of the appropriate enzyme buffer, 1 μ L of enzyme, and 2.5 μ L of double-distilled water for a total volume of 15 μ L. The reaction mixtures were incubated for 1 h at 37°C (BamHI) or 60°C (BstUI) 10 .

Statistical analysis

Variables were compared using Student T test, Chi Square and Mann Whitney tests, as appropriated. Results are shown as mean and standard deviation or frequencies and percentage. A p value <0.05 was considered significant.

RESULTS

During the July 2005 to December 2006 period a total 671 CSF samples were evaluated, with 460 samples (460 patients) fulfilling the inclusion criteria for the study. A total 59/460 (12.8%) samples showed positive results for enterovirus (n=48), herpes simplex virus (n=6), Epstein Barr virus (n=1), human herpes virus type 6 (n=2) and a mixed infection with both enterovirus and Epstein Barr virus (n=1). Enteroviruses represented 83% of positive samples herpes virus family 17%. Samples of herpesvirus samples were digested by restriction enzymes and identified according to the pattern of the signals in the agarosis gel being 6 (10%) for herpes simplex virus, 1 (2%) for Epstein Barr virus and 2 (3%) for human herpes virus type 6 and 1 (2%) co infection of enterovirus and Epstein Barr virus (Fig 1). CMV, VZV, HHV-7, HHV-8 were not

Nogative camples	Positive samples
Negative samples	Positive samples

	Negative samples IQR		Positive samples IQR			-	
	Mean±SD	25%-75%	Median	Mean±SD	25%-75%	Median	р
RBC cell/mm³	580±2890	1-48	5	36.9±183	0.6-13.5	2	0.0071*
WBC cell/mm³	163±271	24-181	80	170.4±279	16-183.5	98	0.7987
Neutrophils (%)	6±17	0-1	1	34.3 ± 33.4	8.5-58.5	28	0.3401
Lymphocytes (%)	80.6±90	51-99	96	63 ± 30.50	39-91.5	61.5	0.0348*
Monocytes-macrophages (%)	17.3 ± 33.2	1-9	1	11.09 ± 23.20	1-8.5	1	0.8212
Total protein (mg/dl)	51.5±31.72	32-62	44	53.5±43.6	28-56.5	43.8	0.5564
Glucose (mg/dl)	68.4±55	57-70	62	70.8±33	59-75	64.5	0.2344

^{*}p<0.05 Mann Whitney Test. RBC: red blood cells; WBC: white blood cells; IQR: interquartiles reference; SD: standard deviation.

found. All negative samples for the tested viruses were positive for the PRV internal control assay indicating efficient nucleic acid extraction.

Biochemical and cellular CSF characteristics of positive and negative samples are shown in Table 1. CSF samples with negative results had RBC mean of 580±2890 cell/mm³ compared to the positive samples where the mean was 36.9 ± 183 cell/mm³ (p=0.0071).

There were higher numbers of positive samples during summer months with a median temperature of 21.5°C (Fig 2), mainly for enteroviruses. Herpesviruses were distributed along the year but its circulation was not observed during the winter months.

There were a high number of positive samples for the 4-6 age group (34%) and 7-17 age group (31%). The same distribution was observed for the cases with suspicion of acute lymphomonocytary meningitis with PCR negative results, being 234 (59%) male and 161 (41%) female patients (p=0.6677). The same pattern was observed for positive samples, 33 (57%) male and 25 (43%) female. The mean age of patients for positive enteroviruses was 6±5 years old and for herpesviruses 11±14 years old. Table 2 shows the age intervals for positive and negative samples. The percentage for positive and negative samples, was great for 4-17 age group (296/453) (p=0.0012). It is noteworthy, enterovirus positive samples (48/59) the mean age was 6±5 years and for herpesviruses (10/58)

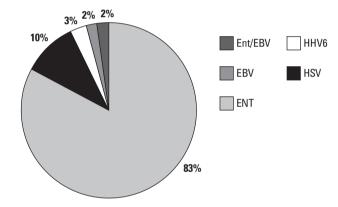


Fig 1. Percentage for positive samples in the period of the study for enterovirus (ENT), herpes simplex virus (HSV) human herpes virus type 6 (HHV6) Epstein Barr virus (EBV) and mixed infection enterovirus and Epstein Barr virus (ENT/EBV).

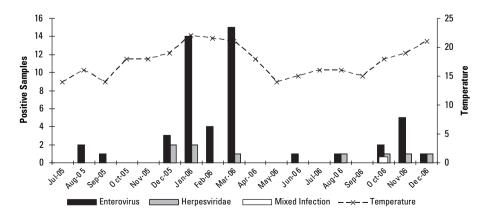
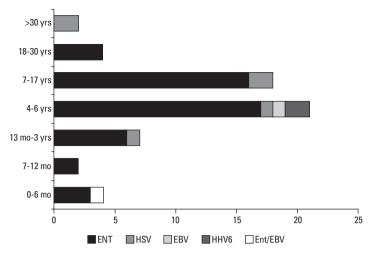


Fig 2. Montly distribution of positive samples and average temperature.

ENT: enterovirus; HSV: herpes simplex virus; EBV: Epstein Barr virus; HHV6: human herpes virus type 6; ENT/EBV: mixed infection; Temp: average temperature.



ENT: enterovirus; HSV: herpes simplex virus; EBV: Epstein Barr virus; HHV6: human herpes virus type 6; ENT/EBV: mixed infection.

Fig 3. Distribution of positive CSF samples by age.

were 11 ± 14 years old (p=0.7645). The mean age for negative patients was 7 ± 7.5 years old. Fig 3 shows the distribution of positive samples according to age. Enterovirus was present in all ages under 30 years. However, positive samples were more prevalent in the ages of 4-17 age group (66.8%). On the other hand, positivity for *Herpesviridae* family was lower with proportional distribution in all ages. The mixed infection with enterovirus and Epstein Barr virus was observed in a 3 month old child.

DISCUSSION

There is little information on viral infections of the CNS in Southern Brazil. The findings of the present study have contributed to a better understanding of the epidemiology of these viruses in this region. We showed here that enterovirus is the main etiologic agent of viral meningitis in Curitiba and probably in the south Brazil, which is in accordance with different reports for other geographic regions^{11,12}. The circulation of the enterovirus occurred mainly in the summer and autumn similarly to that described for the northern hemisphere. In temperate climates the EVs appear markedly during summer/autumm seasonality. However, a high yearround incidence occurs in tropical and subtropical areas. It is believed that fecal-oral spread of these agents is facilitated, particularly among children, during periods of warm weather. In addition to direct person to person transmission, EVs may be recovered from houseflies, wastewater and sewage¹³. Our data is in agreement with those different authors who affirm the importance of establishing a diagnostic tool for patient management as well as carrying out epidemiological analysis to monitor the reappearance of new strains and epidemics by

Table 2. CSF negatives and positives samples from patients. epidemiological characteristics.

epideimological enaracteristics.							
	Enterovirus (n=48)	Family Herpesviridae (n=10)	Negative (n=395)				
Gender							
Μ	26 (54%)	7 (70%)	234 (59%)				
F	22 (46%)	3 (30%)	161 (41%)				
Distribution by age							
0-6 mo	3 (6.25%)	0 (10%)	15 (3.8%)				
7-12 mo	2 (4.16%)	2 (20%)	37 (9.4%)				
13 mo-3 yrs	6 (12.5%)	0	67 (17%)				
4-6 yrs	17 (35.4%)	3 (30%)	132 (33.4%)				
7-17 yrs	16 (33.3%)	2 (20%)	131 (33.1%)				
18-30 yrs	4 (8.39%)	1 (10%)	6 (1.5%)				
>30 yrs	0	2 (20%)	7 (1.8%)				

Kruskal Wallis: p=0.0012.

enteroviruses^{14,15}. Santos et al. described the frequency of enterovirus meningitis analyzing CSF samples collected in 11 states in Brazil¹⁶. Enterovirus was detected in 15.8% of them with a predominance of Echovirus type 30, and five meningitis outbreaks were identified during the study period from 1998 to 2003. The authors pointed out the importance of evaluating epidemiological features responsible for the frequency of viral meningitis and phylogenetic relationships between serotypes in Brazil¹⁶. Mendonza et al. described virus involvement in SNC infections in Brazil¹⁷. This analysis was performed in 200 CSF samples and viral genome was detected in 23.34%, which includes CMV (6%), HSV1 (5%), VZV and EBV (0.5%) and enterovirus (11.34%)¹⁷. Another study in northern Brazil found 13.2% of positive CSF samples for enterovirus using cellular culture, immunofluorescence and neutralization test. The authors reported the importance of determining a national picture of viral meningitis in Brazil since it is still incomplete¹⁸. Despite the use of different methodologies, the proportion of positive samples was similar to that observed in this study. The incidence of the different viruses varies globally according to different populations, methodologies and objectives of the study. Enteroviruses detected by molecular biology RT-PCR or Real Time PCR or cellular culture analysis show an incidence varying from 10 to 46%^{9,12,15,17,19}. Herpesviruses, HHV6, VZV, and a mixed infection by HHV6/EV, were detected in a lower proportion²⁰⁻²². In the present study in addition to enterovirus, positive samples for HSV (10%), EBV (3%) and HHV6 (3%) were observed. Several studies have already shown mixed infection²³. The involvement of herpesviruses as etiologic agents of meningitis or encephalitis has been extensively described in immunocompetent as well as immunossupressed patients²³⁻²⁵. Some authors report the difficulty to distinguish aseptic from bacterial meningitis due to the overlap of clinical symptoms and laboratory findings^{26,27}. In this context, CSF examination is an essential part of the diagnostic analysis of patients with suspected CNS viral infection²⁸. The classical pattern of viral meningitis is an increase of CSF WBCs, lymphomonocytary with normal glucose and elevated lactic acid. However, in the very acute stages of viral meningitis, chiefly in enteroviral meningitis, polymorphonuclear granulocytes may predominate and shifts to a mononuclear pleocytosis within 6 hours^{27,29}. These patients also have benefic with the virus PCR analysis of CSF, as if the reaction is positive the patient is not submitted to antibiotic therapy and has prompt hospital discharge¹⁹. PCR has been proved to be a key method to detect viral infections of the CNS²⁸. Molecular biology analysis offers advantages over cellular culture, because several days are taken for viral growth and identification using the latter method³⁰. The use of PCR based on the search of specific gene sequence 5'-UTR, a highly conserved region for different serotypes for enterovirus, demonstrates the advantage of this methodology since the results can be obtained in hours providing support in clinical practice and patient management, e.g. the use of antibiotics or the benefits of hospitalization³¹. Herpesviruses analysis by PCR has been found to present high sensitivity and sensibility and has definitely revolutionized virological diagnostic^{28,31-33}. Prior to PCR, the viral diagnosis of CNS infection was based on cellular culture of the CSF, brain biopsy or the demonstration of intrathecal antibody response, so that, in many instances the positive diagnostic was retrospective³⁴. PCR has been largely used for CSF diagnosis with great advantages since it is possible to recover, amplify and detect low copies of the virus genome. In this study, DNA/RNA extraction was done as soon as possible to preserve and recover the viral genome from the CSF samples under PCR analysis. The material used for the procedure was DNA/RNAse free to minimize the inhibition factors for PCR analysis.

The high number of CSF samples negative for the viruses studied suggests other causes of lymphomonocytary meningitis, although less prevalent, could be associated. Other infectious agents such as *Mycobacterium tuberculosis*, *Treponema pallidum*, *Cryptococcus neoformans*, *Lysteria monocytogenes*, *Brucella spp.*, Mycoplasma, neurocysticercosis, leptospirose or non infectious conditions including autoimmune diseases and carcinomatous meningitis must be considered²⁹. Also other viruses not detected by PCR included adenovirus, HIV, measles, rubella and mumps.

Additionally, the implication of the many inhibiting

factors in PCR results is well known. In CSF samples collected less than 3 days after the onset of neurological symptoms only 12% showed positive results. A higher positivity was observed on the fourth or fifth day³⁴. In our study most of the CSF samples were collected on the second day (median 12hs) which could explain the high number of negative samples. Additionally, the elevated number of RBC cells (p=0.0071), due to traumatic lumbar puncture, in the CSF's samples, as well as the presence of hemoglobin, probably had an inhibitory effect showing false negative results. We excluded from this analysis CSF samples with predomination of neutrophils, as the aim of this study was lymphomonocytary meningitis. This could have little impact in the number of positive PCR samples as the samples included in this study were collected with a mean of 12 hrs.

Finally, PCR analysis must include strict criteria in order to assure an accurate diagnosis. The standardization of the molecular technique confirms the validity of the PCR method for the rapid detection of RNA enterovirus and DNA herpesvirus in CSF. Enterovirus PCR takes eight hours and pan-herpes PCR takes four and a half hour. Further analysis of other agents and the standardization of biology molecular methodologies such as Real Time PCR are future proposition to improve the laboratorial diagnosis of viral meningitis in Brazil.

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