


BMJ Open Prevalence and risk factors for SARS-CoV-2 infection in children with and without symptoms seeking care in Managua, Nicaragua: results of a cross-sectional survey

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ABSTRACT

Objective This study aimed to capture key epidemiological data on SARS-CoV-2 infection in Nicaraguan children (≤ 18 years) seeking medical care, between 6 October and 16 November 2020.

Design In this cross-sectional study, 418 children were recruited: 319 with symptoms characteristic of COVID-19 and 99 with no symptoms of illness. Children were tested for SARS-CoV-2 RNA using loop-mediated isothermal amplification. A questionnaire was employed to identify symptoms, risk factors, comorbidities and COVID-19 prevention measures.

Setting Research was carried out in four hospitals and two clinics in Managua, Nicaragua, where schools and businesses remained open throughout the COVID-19 pandemic.

Participants Children were enrolled into a possible COVID-19 group if presenting with clinical symptoms. A comparison group included children lacking any COVID-19 symptoms attending routine check-ups or seeking care for issues unrelated to COVID-19.

Results A high prevalence (43%) of SARS-CoV-2 infection was found, which was relatively equivalent in symptomatic and non-symptomatic children. Age distribution was similar between symptomatic and non-symptomatic children testing positive for SARS-CoV-2. Symptomatic children who tested positive for SARS-CoV-2 were 2.7 times more likely to have diarrhoea (26.7% in positive vs 12.0% in negative; OR=2.7 (95% CI 1.5 to 4.8), $p=0.001$) and were 2.0 times more likely to have myalgia (17.8% in positive vs 9.8% in negative; OR=2.0 (95% CI 1.0 to 3.8), $p=0.04$). Children with COVID-19 symptoms, who tested positive for SARS-CoV-2, were more likely to be under age 5 years and to have a pre-existing comorbid condition than children who tested positive but did not have symptoms.

Conclusions This is the first paediatric study to provide laboratory-confirmed data on SARS-CoV-2 infection in Nicaragua, crucial for paediatric health services planning and a successful COVID-19 response. The high prevalence of the virus suggests widespread and sustained community transmission, underscoring the urgent need for robust data on the true extent of SARS-CoV-2 infection throughout Nicaragua.

Strengths and limitations of this study

- Primary data from low-income countries are urgently needed. This is the first study providing data obtained from testing of SARS-CoV-2 infection in the Nicaraguan paediatric population, presenting evidence that fills a major research gap.
- We accomplished this by using a simple, highly sensitive detection technique, loop-mediated isothermal amplification, which may be used in developing countries both as a detection method and for epidemiological surveillance.
- Our study indicates widespread and sustained community transmission in Nicaragua, particularly among children, an issue that merits additional urgent attention to improve the overall response to COVID-19 disease.
- This research is critical, considering the lack of effective pandemic response and of credible COVID-19 statistics in Nicaragua.
- The study's data may not reflect the national prevalence of SARS-CoV-2 among Nicaraguan children since most children were from the capital of Nicaragua, and it is possible that some individuals were more likely to participate if they had a history of close contact with confirmed cases.

INTRODUCTION

The Latin American and Caribbean regions continue to face serious difficulties in containing the COVID-19 pandemic, with more than 21.8 million confirmed COVID-19 cases and 689 967 deaths.¹ The situation may be more difficult in the countries of Central America, such as Nicaragua, where insufficient health services and limited infrastructure are the norm. The Nicaraguan Ministry of Health confirmed the first COVID-19 case in Nicaragua on 18 March 2020. However, more than a year later, not much is known about the true COVID-19 situation in Nicaragua, and it is almost impossible to assess

the growth of the pandemic, as testing is largely unavailable. Official data lack accuracy and transparency, resulting in little public confidence. As of 24 February 2021, the official government count admits to only 6445 confirmed cases and 173 deaths,² but the Citizens Observatory of COVID-19, an independent research group, estimated a cumulative number of cases of 13 140 and with at least 2976 deaths from COVID-19, making it one of the highest COVID-19 mortality rates in Latin America.³ Furthermore, official death tolls undercount the total number of fatalities and the government-run health centres seldom report COVID-19 as the cause. Between March and August of 2020, the number of deaths exceeded the average number of deaths in previous years by 7600, a figure that was 47% higher than the expected number for that period and almost 50 times the deaths officially attributed solely to COVID-19.⁴

In direct contrast to most Latin American countries, the Nicaraguan authorities refused to implement any significant public health or social interventions in response to the pandemic.⁵ While more than a hundred countries had closed in-person school attendance by mid-March, 2020,⁶ the Nicaraguan government kept public schools open all through the pandemic, and there were no restrictions imposed on transport and mobility. However, school attendance was irregular, varying by grade and geographical area, and some parents decided not to send their children to school. These closures and irregularities confound the understanding of children susceptibility to SARS-CoV-2 infection.

Clinical manifestations of COVID-19 are generally milder in children compared with adults,⁷ although some children do require hospitalisation and intensive care.⁸ Recent studies suggest that children are just as likely as adults to become infected with SARS-CoV-2 but have fewer symptoms and lower case fatality rates.^{9,10} Considering public health policy implications, more studies on the epidemiology of SARS-CoV-2 infections are needed along with data on the clinical outcome of COVID-19 in children¹¹ and the role of children in SARS-CoV-2 transmission. However, most data come primarily from higher income countries where rates of childhood obesity and diabetes may be higher, leading to disproportionately severe outcomes. The public health implications of the pandemic in small countries with limited resources remain underinvestigated and under-reported. The aim of this study was to capture key epidemiological data on SARS-CoV-2 infection in Nicaraguan children and adolescents 18 years and younger (≤ 18 years) seeking medical care to inform physicians, healthcare workers and decision makers on the impact of the pandemic in children and to provide a resource for paediatric health services planning and overall COVID-19 infection response.

MATERIALS AND METHODS

Study population and data collection

The Pediatrics COVID-19 Network was established to conduct this research in Managua, Nicaragua. This network is composed of 33 paediatricians from four hospitals and two clinics across the city. From 6 October

to 16 November 2020, study staff at participating hospitals and clinics recruited children and adolescents (≤ 18 years) at the outpatient units by inviting parents/guardians to enrol their children in the study. Age was the only inclusion or exclusion criteria. All parents were approached for participation. Parents who showed interest in volunteering their children for the study were informed of the study's purpose and were asked to sign a form indicating informed consent. Children were recruited into a possible COVID-19 group based on symptoms, as judged by the individual clinician. Any children presenting with COVID-19 symptoms were generally considered possible COVID-19 cases. Clinicians looked for the following most commonly associated COVID-19 symptoms: fever, cough, sore throat, diarrhoea, headache, fatigue and myalgia. The full list of symptoms used to classify as probable COVID-19 case is in the study questionnaire (online supplemental file 1). A comparison group included children lacking any COVID-19 symptoms, who were all attending routine check-ups or seeking care for issues unrelated to COVID-19. The screening settings were the same for both the possible COVID-19 group and the comparison group. None of the study participants were asked about their possible exposure to the SARS-CoV-2 virus prior to recruitment. A total of 418 children were recruited. All children were tested for SARS-CoV-2 RNA through the loop-mediated isothermal amplification (LAMP) detection method as described previously.¹²

Survey tool

A questionnaire was employed to identify: (1) possible exposures or risk factors for infection, (2) current symptoms and (3) factors associated with vulnerability to severe illness and complications (age, sex and chronic medical conditions). The questionnaire also gathered data on school attendance, use of face masks, handwashing habits and presence of a family member with COVID-19 symptoms at home. Paediatricians filled out the questionnaires while interviewing parents and the children (online supplemental file 1).

Specimen collection and testing

Between 1 and 4 mL of saliva were self-collected by each study participant age 3 years and over in flasks containing 2 mL of sample buffer¹³ (1× phosphate buffered saline, pH 7) for LAMP detection of SARS-CoV-2. LAMP detection has previously demonstrated 97.5% and a specificity of 99.7% for detection of SARS-CoV-2 infections. For children under 3 years of age, we used mouth swabs (Fisherbrand) designed to painlessly collect saliva for analysis. After collection, specimens were kept on ice and transported immediately to the laboratory for the detection of nucleic acids from SARS-CoV-2. Testing assays were performed on saliva samples without an RNA purification step within 1–2 hours following collection, as previously described.¹² All LAMP reactions were performed following New England Biolab's (NEB) published protocol using WarmStart Colorimetric LAMP 2× Master

Mix (NEB, M1800L).^{14 15} We used 20 µL reactions consisting of 10 µL of 2× master mix, 2 µL of 10× primer mix targeting the viral genes N and E, 5 µL nuclease-free water and 3 µL samples (NEB E2019 COVID-19 LAMP kit). LAMP reactions were incubated at 65°C using an Applied Biosystems 2720 Thermal Cycler for 45 min. For quality control, we performed 20 additional saliva tests on individuals who had been diagnosed by quantitative reverse transcription PCR (qRT-PCR) using nasopharyngeal swabs at the central government laboratory. There was 100% concordance with the LAMP results. These 20 volunteers (12 SARS-CoV-2 positives and 8 SARS-CoV-2 negatives) were identified among university staff. Results were communicated to paediatricians within 6 hours via phone calls, who in turn conveyed the results, along with appropriate medical advice, to the corresponding parents or legal guardians.

Clinical features in children

Following the USA National Institutes of Health (NIH) COVID-19 Treatment Guidelines,¹⁶ the illness severity of patients with COVID-19 was stratified into the following categories: asymptomatic or presymptomatic infection, mild, moderate, severe and critical illness. Asymptomatic or presymptomatic infection: individuals who test positive for SARS-CoV-2 using a virological test but who have no symptoms that are consistent with COVID-19. No symptoms: individuals who showed none of the key COVID-19 symptoms at the moment of testing. Participating paediatricians collected clinical information on cases using the survey tool, recording age, sex, underlying disease, date of diagnosis, risk factors for exposure, symptoms, laboratory test results, treatment and outcome. Twenty-five of those testing positive were followed for 21 days to determine clinical outcome.

Statistical analysis

All data were entered into a Microsoft Excel spreadsheet including results of SARS-CoV-2 detection using the LAMP assay, along with data obtained from the questionnaire. Prior to statistical analyses, all data were transferred to a spreadsheet without linkage to the original database, eliminating all personal identification data. We identified the proportion positive in the two recruitment groups: symptoms/no symptoms. Participants were categorised by group: symptomatic and no symptoms and further divided by SARS-CoV-2 infection result from the LAMP assay. For some analyses, all four strata were maintained, and in others, we grouped by infection status. All analyses were conducted using SAS V.9.4.

Proportion of positive tests were compared by age, sex and symptom presentation. Pearson χ^2 tests were conducted to identify differences in the distribution of age, sex and presentation (symptoms/no symptoms) in children with positive versus negative test results. ORs were estimated to: (1) compare symptoms between children with negative versus positive tests overall and by age group, (2) identify ORs of symptomatic versus children

with no symptoms who tested positive and (3) to examine associations between social factors for those with positive versus negative LAMP tests. ORs were estimated using logistic regression modelling. Models of social factors associated with positive and negative tests were stratified by children with and without symptoms and were adjusted for age and sex.

Patient and public involvement

Patients and members of the public were not directly involved in the design and conduct of this study. The publication of this study will be shared with all participants, and links to this work will be included on the institutional website.

RESULTS

In this cross-sectional study, we included 418 children and adolescents under the age of 18 years with a nearly equal sample of males (206/49%) and females (212/51%). More than half of the participating children were ≤5 years of age. Children were recruited into two distinct groups: children presenting with COVID-19 symptoms and children without COVID-19 symptoms and further divided by SARS-CoV-2 infection result from the LAMP assay (figure 1). The demographic characteristics of study participants are summarised in table 1.

SARS-CoV-2 prevalence

A total of 181 children (43.3%) tested positive for SARS-CoV-2. A percentage of 74.6 (135 of 181) were symptomatic and 25.4% (46 of 181) were asymptomatic carriers.

We examined infection rates among participants in three different age groups: 0–5, 6–11 and 12–18 years. Within the ≤5 age group, 109 children (45.6%) were SARS-CoV-2 positive, while in the 6–11 and 12–18 age groups, 46 (40%) and 26 (40.6%) tested positive, respectively (table 1). In the comparison group of children with no symptoms, 46 children (46.5%) were SARS-CoV-2 positive. The youngest SARS-CoV-2 patient identified was 21 days old. Girls (105 of 212 (49.5%)) had higher prevalence of SARS-CoV-2 infection than boys (76 of 206 (36.9%); $p=0.009$) (table 1).

Clinical presentation of children

The most common symptoms reported by those testing positive for SARS-CoV-2 were fever ($n=83$; 61.5%) and cough ($n=82$; 60.7%) (table 2). Symptomatic children who tested positive for SARS-CoV-2 were 2.7 times more likely to have diarrhoea (26.7% in positive vs 12% in negative; $OR=2.7$ (95% CI 1.5 to 4.8), $p=0.001$) and were 2.0 times more likely to have myalgia (17.8% in positive vs 9.8% in negative; $OR=2.0$ (95% CI 1.0 to 3.8), $p=0.04$). They were also less likely to have sore throat (20% in positive vs 29.4% in negative; $OR=0.6$ (95% CI 0.4 to 1.0), $p=0.06$). Analysis of symptoms associated with a positive SARS-CoV-2 test across the age groups demonstrated similar patterns. Diarrhoea presented more commonly in

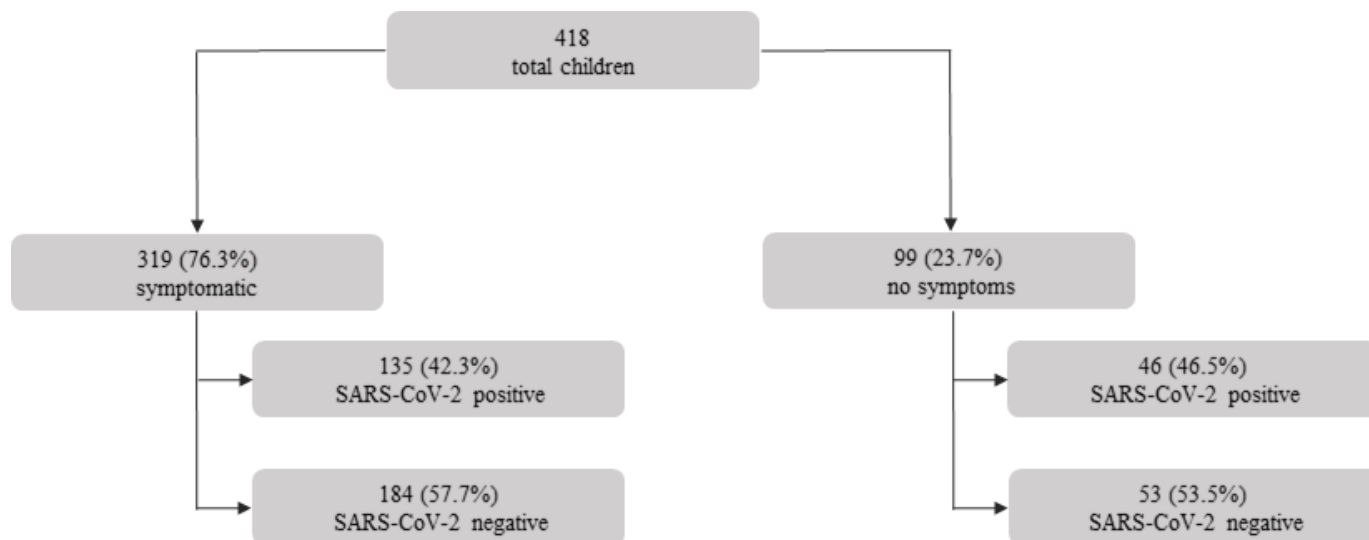


Figure 1 Flow chart of study participants and SARS-CoV-2 infection status.

all age groups for those positive for SARS-CoV-2 (table 3) though this did not reach significance in the oldest age group ($p=0.19$). Conjunctivitis was positively associated with a positive SARS-CoV-2 result in children under age 5 years, but not in other age groups. Children who tested positive for SARS-CoV-2 that had symptoms were more likely to be under age 5 years ($OR=11.1$ (95% CI 4 to 31.1), $p<0.0001$) and have a pre-existing comorbid condition ($OR=2.6$ (95% CI 1.0 to 6.7), $p=0.04$ (table 4).

In our study, 97 (23.2%) out of the 418 enrolled children had at least one pre-existing chronic underlying disease and several had an association of two or more chronic conditions. Among the 319 symptomatic children, the most common comorbidity was respiratory disease ($n=40$; 12.5%) (table 2). Other important chronic diseases were heart disease ($n=12$; 3.8%) and obesity ($n=6$; 1.9%). Children with pre-existing cardiovascular disease that presented with symptoms were 7.3 times more likely to test positive for SARS-CoV-2 ($OR=7.3$ (95% CI 1.6 to

33.8), $p=0.01$). Furthermore, data in table 4 indicate that within positive children, those who had comorbidities were more likely to be in the group with symptoms $OR=2.6$ (95% CI 1.0 to 6.7) $p=0.04$.

Clinical course of the disease

Among all SARS-CoV-2 confirmed cases, the paediatrics network was able to follow-up the clinical course of the disease for 25 (14% of test positive) children (online supplemental file 2). Out of these 25 children, severity of illness was reported as mild for 12 (48%), moderate for 12 (48%) and severe for 1 (4%). Fifteen of these children presented underlying medical conditions that put them at increased risk for severe illness from COVID-19. The youngest group (≤ 5) represented the majority of mild (67%) and moderate (75%) cases, whereas the only severe case reported was within the 6–11 years age group. Overall, seven (4%) children were hospitalised

Table 1 Demographic data of children enrolled in the study and SARS-CoV-2 test results, $n=418$

Category	Total	Positive test $n=181$, n (%)	Negative test $n=237$, (%)	χ^2 (df, N)=, p value
Age, years				
≤ 5	239	109 (45.6)	130 (54.4)	χ^2 1.2, p value=0.54
6–11	115	46 (40)	69 (60)	
12–18	64	26 (40.6)	38 (59.3)	
Sex				
Male	206	76 (36.9)	130 (63.1)	χ^2 6.8, p value=0.009
Female	212	105 (49.5)	107 (50.5)	
Symptom history				
No symptoms	99	46 (46.5%)	53 (53.5)	χ^2 0.53, p value=0.47
Symptoms	319	135 (42.3%)	184 (57.7)	

Bold values are considered statistically significant.

*Pearson χ^2 and p values compares the distribution of age, sex and symptoms by positive and negative test results.

Table 2 Symptoms and comorbidities of children and SARS-CoV-2 test results for those presenting with COVID-19 like symptoms

	Total n=319 (76.3%)	Positive n=135 (42.3%)	Negative n=184 (57.7%)	OR (95% CI)	P value for OR
Screening symptoms distribution, n (%)					
Fever	190 (59.6)	83 (61.5)	107 (58.2)	1.1 (0.8 to 1.4)	0.55
Cough	190 (59.6)	82 (60.7)	108 (58.7)	1.1 (0.7 to 1.7)	0.71
Rhinorrhoea	125 (39.2)	57 (42.2)	68 (37.0)	1.0 (0.6 to 1.6)	0.95
Sore throat	81 (25.4)	27 (20)	54 (29.4)	0.6 (0.4 to 1.0)	0.06
Diarrhoea	58 (18.2)	36 (26.7)	22 (12.0)	2.7 (1.5 to 4.8)	0.001
Headache	58 (18.2)	22 (16.3)	36 (19.6)	0.8 (0.4 to 1.4)	0.46
Asthenia	53 (16.6)	22 (16.3)	31 (16.8)	1.0 (0.5 to 1.7)	0.90
Myalgia	42 (13.2)	24 (17.8)	18 (9.8)	2.0 (1.0 to 3.8)	0.04
Vomiting	21 (6.6)	9 (6.7)	12 (6.5)	1.1 (0.3 to 3.8)	0.83
Difficulty breathing or shortness of breath	16 (5.0)	7 (5.2)	9 (4.9)	1.1 (0.4 to 2.9)	0.90
Loss of taste	5 (1.6)	3 (2.2)	2 (1.1)	2.1 (0.3 to 12.6)	0.42
Loss of smell	3 (0.9)	2 (1.5)	1 (0.5)	2.8 (0.2 to 30.7)	0.41
Other symptoms					
Skin rash	13 (4.1)	7 (5.2)	6 (3.3)	1.6 (0.5 to 4.9)	0.39
Conjunctivitis	10 (3.1)	7 (5.2)	3 (1.6)	3.3 (0.8 to 13.0)	0.08
Chest pain or pressure	10 (3.1)	4 (3.0)	6 (3.3)	0.9 (0.3 to 3.3)	0.88
Loss of colour on the fingers of the hands or feet	3 (0.9)	1 (0.7)	2 (1.1)	0.7 (0.1 to 7.6)	0.75
Loss of speech and movement	1 (0.3)	1 (0.7)	0 (0)	nd	
Comorbidities					
Respiratory diseases	40 (12.5)	21 (15.6)	19 (10.3)	1.6 (0.8 to 3.1)	0.17
Cardiovascular diseases	12 (3.8)	10 (7.4)	2 (1.1)	7.3 (1.6 to 33.8)	0.01
Obesity	6 (1.9)	3 (2.2)	3 (1.6)	1.4 (0.3 to 6.9)	0.70
Cancer	1 (0.3)	0 (0)	1 (0.5)	nd	
Kawasaki disease	1 (0.3)	1 (0.7)	0 (0)	nd	

P values for the bivariate models are indicated in the rightmost column.

Bold values are considered statistically significant.

*ORs generated using logistic regression modelling with 95% CI.

nd, not determined.

with COVID-19, including two (1%) that were admitted to the intensive care unit (ICU). No patient died.

Contact with suspected COVID-19 cases

Seventy-nine (19%) out of the 418 enrolled children reported having at least one family member with COVID-19 symptoms at home (table 5). Of those, 48 were females (61%) and 31 males (39%). In children without symptoms, those with a family member currently sick at home were 2.8 times more likely to test positive for SARS-CoV-2 (OR=2.8 (95% CI 1.1 to 6.7), p=0.03). Out of those that reported at least one family member with COVID-19 symptoms, 43.0% (n=34) resulted SARS-CoV-2 positive, 14 symptomatic and 20 without any symptoms. Among females who reported having a family member with COVID-19 symptoms, 45.8% (n=22) resulted SARS-CoV-2

positive, including 8 symptomatic and 14 asymptomatic females, whereas among males, 38.7% (n=12) resulted positive, six symptomatic and six asymptomatic males. Nearly half of families reported having had a family member with suspected COVID-19 in the past (41%) with fewer reporting knowing individuals sick at school (19%), work (21%) or in the neighbourhood (22%).

Outside activities and SARS-CoV-2 infection

One hundred and thirty-three children (32%) were reported attending school in person during the study period (table 5), 73 females (55%) and 60 males (45%). Out of the 73 females that attended school, 28 (38%) resulted SARS-CoV-2 positive. Among the 60 males reported to attend school in person, 18 (30%) resulted SARS-CoV-2 positive. In children who presented with

Table 3 Symptoms associated with a positive test by age group (reduced subset limited to factors p<0.20)

	Total, n (%)	Positive, n (%)	Negative, n (%)	OR (95% CI)	P value for OR
Under age 5 years	n=216	n=100 (46%)	n=116 (54%)		
Cough	133 (62)	68 (68)	65 (56)	1.7 (0.6 to 2.9)	0.07
Diarrhoea	38 (18)	24 (24)	14 (12)	2.3 (1.1 to 4.7)	0.02
Myalgia	24 (11)	15 (15)	9 (8)	2.1 (0.9 to 5.0)	0.09
Conjunctivitis	7 (3)	5 (5)	2 (2)	3.0 (0.6 to 15.8)	0.19
5–12 years	n=68	n=22 (32%)	n=46 (68)		
Cough	40 (59)	10 (45)	30 (65)	0.4 (0.2 to 1.3)	0.12
Rhinitis	21 (31)	4 (18)	17 (37)	0.4 (0.1 to 1.3)	0.12
Sore throat	32 (47)	7 (22)	25 (54)	0.4 (0.1 to 1.1)	0.09
Diarrhoea	11 (16)	7 (31)	4 (9)	4.9 (1.3 to 19.1)	0.02
Age >12 years	n=35	n=13 (37)	n=22 (63)		
Cough	17 (49)	4 (31)	13 (59)	0.3 (0.1 to 1.3)	0.11
Rhinitis	8 (23)	1 (8)	7 (32)	0.2 (0.02 to 1.7)	0.13
Diarrhoea	9 (26)	5 (38)	4 (18)	2.8 (0.6 to 13.3)	0.19
Myalgia	9 (26)	5 (38)	4 (18)	2.8 (0.6 to 13.3)	0.19

Bold values are considered statistically significant.

symptoms, those who went to school were less likely to test positive for SARS-CoV-2 than those not in school (OR=0.8 (95% CI 0.6 to 1.0), p=0.08). This association was not noted in children who were presenting without symptoms (OR=1.2 (95% CI 0.8 to 2.0), p=0.40).

Regarding the use of transportation to get to school, 117 (80.7%) children reported using private family vehicles, 12 (8.3%) walked to school and 16 (11%) children used public transportation. Among those children transported to school in private family vehicles, 37 (31.6%) were SARS-CoV-2 positive, including 28 symptomatic and

9 asymptomatic children. Among children who walked to school, three (25%) were SARS-CoV-2 positive. Out of 16 children that used public transportation to get to school, 12 (75%) were SARS-CoV-2 positive (six symptomatic and six asymptomatic children). In children who presented with symptoms, those who rode in the family car or walked were 0.1 time as likely to test positive for SARS-CoV-2 when compared with those who rode public or school buses (OR=0.1 (95% CI 0.1 to 0.8), p=0.03) (table 5).

Table 4 Factors associated with having symptoms in those that have a SARS-CoV-2 positive test

	Symptoms (n=135)	No symptoms (n=46)	OR (95% CI)	P value
Sex, n (%)				
Male	59 (77)	17 (23)	1.3 (0.7 to 2.6)	0.42
Female	76 (72)	29 (28)	ref	
Age group (years), n (%)				
≤5	100 (92)	9 (8)	11.1 (4.0 to 31.1)	<0.0001
5–11	22 (48)	24 (52)	0.9 (0.4 to 2.4)	0.86
12–18	13 (50)	13 (50)	ref	
Comorbidities, n (%)				
Any comorbidity	38 (86)	6 (14)	2.6 (1.0 to 6.7)	0.04
Respiratory diseases	21 (88)	3 (13)	2.6 (0.7 to 9.3)	0.13
Cardiovascular diseases	10 (100)	0 (0)	nd	
Obesity	3 (100)	0 (0)	nd	

P values for the bivariate models are indicated in the rightmost column.

Bold values are considered statistically significant.

*OR generated using logistic regression models with 95% CI.

N, sample size; nd, not determined.

Table 5 Social factors related to positive tests for children presenting with symptoms and those without symptoms

Exposure variable (n=total, asymptomatic, symptomatic)	No symptoms			Symptoms			
	Reporting exposure (n, %)	SARS-CoV-2 positive (n, %)	SARS CoV-2 negative (n, %)	OR* (95% CI)	SARS CoV-2 positive (n, %)	SARS CoV-2 negative (n, %)	OR* (95% CI)
Suspected contact with individuals infected with COVID-19							
Family member currently sick (n=418 [†] , 319 [§] , 99 [¶])	79 (19)	20 (43)	11 (21)	2.8 (1.1 to 6.7), p=0.03	14 (10)	34 (18)	0.5 (0.4 to 1.0), p=0.05
Family member history of suspected of COVID (n=418 [†] , 319 [§] , 99 [¶])	172 (41)	24 (52)	29 (55)	0.9 (0.4 to 1.9), p=0.69	55 (41)	64 (35)	1.4 (0.9 to 2.2), p=0.19
Knows sick people at school (n=330 [†] , 247 [§] , 83 [¶])	62 (19)	9 (23)	12 (28)	0.7 (0.2 to 1.9), p=0.46	12 (12)	29 (20)	0.6 (0.3 to 1.3), p=0.19
Knows sick people at work (n=347 [†] , 266 [§] , 81 [¶])	74 (21)	11 (28)	5 (12)	2.9 (0.9 to 9.6), p=0.07	22 (20%)	36 (23)	0.9 (0.5 to 1.7), p=0.77
Knows sick people in neighbourhood (n=365 [†] , 276 [§] , 89 [¶])	82 (22)	10 (24)	15 (32)	0.6 (0.2 to 1.7), p=0.37	22 (19)	35 (22)	0.9 (0.5 to 1.7), p=0.88
Household characteristics							
Family size (n=399 [†] , 305 [§] , 94 [¶])							
1–3 people	55 (13)	7 (15)	12 (23)	ref	11 (8)	25 (14)	ref
4–6 people	263 (63)	27 (59)	34 (64%)	1.3 (0.4 to 3.8) p=0.64	85 (63)	117 (64)	1.8 (0.8 to 4.0), p=0.12
>6 people	100 (24)	12 (26)	7 (13)	2.6 (.7 to 9.9) p=0.16	39 (29)	42 (23)	2.2 (0.9 to 5.1), p=0.07
Any children attending school (n=364 [†] , 277 [§] , 87 [¶])	147 (40)	16 (40)	16 (34)	1.4 (.5 to 3.5), p=0.51	38 (33)	77 (48)	0.6 (0.4 to 1.0), p=0.04
Outside activities							
Child attends school (n=412 [†] , 313 [§] , 99 [¶])	133 (32)	16 (35)	15 (28)	1.2 (0.8 to 2.0) p=0.40	30 (23)	72 (40)	0.8 (0.6 to 1.0), p=0.08
Family school transport (n=145 [†] , 111 [§] , 34 [¶])							
Car/walking versus Bus (public or school)	129 (89)	11 (65)	15 (88)	0.2 (0.1 to 1.5), p=0.13	29 (83)	74 (97)	0.1 (0.1 to 0.8), p=0.03
Number of people who work outside home (n=383 [†] , 291 [§] , 92 [¶])							
0–1	172 (45)	22 (52)	26 (48)	1.0 (0.2 to 4.6), p=1.0	48 (38)	79 (48)	0.6 (0.3 to 1.0), p=0.05
2	149 (39)	20 (48)	21 (5%)	0.8 (0.2 to 3.8), p=0.80	41 (33)	62 (38)	0.6 (0.3 to 1.1), p=0.09

Continued

Table 5 Continued

Exposure variable (n=total, asymptomatic, symptomatic)	Reporting exposure (n, %)	No symptoms			Symptoms		
		SARS-CoV-2 positive (n, %)	SARS CoV-2 negative (n, %)	OR* (95% CI)	SARS CoV-2 positive (n, %)	SARS CoV-2 negative (n, %)	OR* (95% CI)
3 or more	62 (16)	1 (1)	0 (0)	ref	37 (29)	24 (15)	ref
Many activities outside home (n=329 ^t , 245 ^s , 84 ⁿ)	162 (49)	15 (35)	16 (39)	1.1 (0.4 to 2.8), p=0.82	58 (55)	73 (53)	1.1 (0.7 to 1.9), p=0.66
Prevention behaviours							
Child wears a mask (n=366 ^t , 277 ^s , 89 ⁿ)	216 (59)	29 (67)	37 (80)	0.5 (0.2 to 1.4), p=0.19	49 (41)	101 (64)	0.4 (0.2 to 0.7), p=0.001
Family wears a mask (n=394 ^t , 300 ^s , 94 ⁿ)	173 (44)	20 (44)	31 (63)	0.5 (0.2 to 1.1), p=0.10	51 (40)	71 (41)	1.0 (0.6 to 1.6), p=0.90
Tutors wears a mask (n=407 ^t , 310 ^s , 97 ⁿ)	376 (92)	44 (98)	48 (92)	4.2 (0.4 to 40.2), p=0.21	116 (87)	168 (95)	0.3 (0.1 to 0.7), p=0.005
Visitors wear a mask (n=397 ^t , 299 ^s , 98 ⁿ)	207 (52)	25 (54)	33 (63)	0.6 (0.3 to 1.5), p=0.18	64 (51)	85 (49)	1.0 (0.6 to 1.6), p=0.91
Family physically distances when visits (n=351 ^t , 264 ^s , 87 ⁿ)	124 (35)	21 (55)	27 (55)	1.0 (0.4 to 2.4), p=0.98	31 (12)	45 (30)	0.9 (0.5 to 1.6), p=0.78
Visitors physically distance when visiting (n=351 ^t , 264 ^s , 87 ⁿ)	162 (46%)	23 (59%)	29 (60%)	0.9 (0.4, to 2.2), p=0.83	44 (41%)	66 (42%)	1.0 (0.6 to 1.6), p=0.87
Talks about COVID-19 with neighbours (n=383 ^t , 295 ^s , 88 ⁿ)	187 (49%)	23 (55%)	29 (63%)	0.7 (0.3 to 1.7), p=0.42	58 (46%)	77 (46%)	1.0 (0.6 to 1.7), p=0.88

Bold values are considered statistically significant.

*Age-adjusted and sex-adjusted ORs with 95% CI, % use sample size as denoted in column 1 (t=total sample size for that variable, s=sample size for the symptomatic children, n=sample size for the non-symptomatic children), proportion is for the total that answered 'yes' to the factor of interest or the categories as noted. ORs for a positive test were calculated using the 'no' response as reference.

Out of the 418 enrolled children, 219 children (113 (51.6%) females and 106 (48.4%) males), reported having activities outside of their homes, including visiting families and friends, birthday parties, shopping and attending church activities. Ninety-four (42.9%) of those reporting activities outside the home tested SARS-CoV-2 positive; 52 (55.3%) females versus 42 males (44.7%) resulted positive for COVID-19. Furthermore, out of the 162 children that reported having frequent activities outside their homes, 73 (45%) tested SARS-CoV-2 positive, 58 (55%) symptomatic and 15 (35%) non-symptomatic children. Though there was no significant association between reporting having many activities outside the home and testing positive for SARS-CoV-2 (table 5).

Prevention strategies

A total of 216 (59.0%) children were reported using face masks frequently (104 males and 112 females). Among females who reported wearing face masks frequently, 49 (43.8%) tested SARS-CoV-2 positive. In the case of males, 29 (27.9%) of those who used face masks frequently tested SARS-CoV-2 positive. Among children who demonstrated symptoms of illness, children who wore a face mask were 0.4 times as likely to test positive for SARS-CoV-2 (OR=0.4 (95% CI 0.2 to 0.7), p=0.001); a similar trend was seen in those who had no symptoms, but it did not reach significance (OR=0.5 (95% CI 0.2 to 1.4), p=0.19). Mask wearing was reported less frequently for the entire family with 44% of families overall wearing masks. Tutors were

reported as wearing masks most frequently, with 92% of tutors reported as wearing a mask overall. Having a tutor wear a mask was associated with a reduced risk of SARS-CoV-2 infection in children with symptoms (OR=0.3 (95% CI 0.1 to 0.7) $p=0.005$). There was no association in children without symptoms. Visitors were reported as wearing masks about half the time (52%) and was not associated with children testing positive in either group. Physical distancing was reported less frequently than wearing a mask with 124 (35%) of individuals indicating family physically distances when visiting. This was somewhat higher for visitors with 162 (46%) physically distancing when visiting. Approximately half ($n=187$, 49%) of participants indicated that they talk with their neighbours about COVID-19.

DISCUSSION

Principal findings

There are five significant findings from our study: (1) there was a high prevalence of SARS-CoV-2 infection during the study period; SARS-CoV-2 was detected in 43.3% of the total number of enrolled children, including in 46.5% of the children without symptoms, suggesting that the infection may be widespread among Nicaraguan children; (2) the younger age group of ≤ 5 years showed higher prevalence (45.6%) of SARS-CoV-2 infection than the other two age groups (6–11 and 12–18 years), and similarly, girls had a higher prevalence than boys (49.5% vs 36.9%) throughout all age groups; (3) out of those that reported at least one family member with COVID-19 symptoms, 43.0% resulted SARS-CoV-2 positive; (4) about half (46.5%) of the children were asymptomatic and 23% had at least one pre-existing chronic underlying disease; and (5) social distancing measures and wearing face masks showed a strong protective effect in significantly reducing infection in children.

Comparison with other studies

The high level of prevalence of SARS-CoV-2 infection among children in our study follows a similar pattern often detected during peaks in the pandemic curve and may be associated with higher incidence of COVID-19 in the general population, as children are as likely to become infected as adults.⁹ Data from a high-incidence setting in India showed positivity rates ranged between 33% and 40%.¹⁷ Cases there had a younger age distribution compared with higher income countries. High COVID-19 prevalence relative to the population size was also estimated for Peru (31%), Mexico (27%) and Brazil (22%).¹⁸

Several studies have indicated that very young children may experience more severe illness.^{19 20} Although it is hard to know for sure why the younger children in our study had the highest prevalence and were more likely to have symptoms of illness, it may be explained from children's habits at home. Infected household members are a major risk factor for paediatric COVID-19 cases^{21 22} and

was associated with a higher risk of infection in children in this study as well. Furthermore, the sources of infection for most children and youths appear to be family members and not schools, as demonstrated in a Hong Kong cross-sectional study.²³

A significant percentage (46.5%) of the children were asymptomatic carriers, which is consistent with the available evidence. Recent publications suggest that approximately 20% of SARS-CoV-2 infected children are asymptomatic, although the prevalence may be as high as 50%.^{24–26} However, it is difficult to establish the exact incidence of asymptomatic SARS-CoV-2 infection, especially when testing is limited.²⁶ Surprisingly, the symptom most associated with having SARS-CoV-2 infection in the children who presented with symptoms was diarrhoea and myalgia, not respiratory symptoms. A study of symptomology in the UK also indicated that diarrhoea was a key predictor of SARS-CoV-2 infection in children,²⁵ and common gastrointestinal symptoms have been reported in COVID-19 infected children.²⁷

Our results do not conflict with current findings by others that COVID-19 is largely a mild disease in children under 19 years, including infants.^{19 20 28–30} However, comorbidities are considered a significant risk factor for requiring ICU admission.²⁰ In our study, 23% of the enrolled children had at least one comorbidity, the most common being respiratory illness (12.5%), which was slightly higher among SARS-CoV-2 positive children (15.6%). Heart disease appears to be an important risk factor for testing positive for SARS-CoV-2 (7.4% vs 1.1%, OR=7.3 (95% CI 1.6 to 33.8), $p=0.01$). Individuals with comorbidities who tested positive were also more likely to be in the symptomatic group of children.

There was a significant association between taking public transportation or school buses to school and infection. Consequently, the use of public transportation may have contributed to the spread of the virus among children. In this study, we found a strong protective effect of masks against SARS-CoV-2 infection. However, children may have also been exposed at home during visits from extended family and friends who could have been SARS-CoV-2 positive and who did not always use face masks nor practice social distancing to limit virus transmission. Accurate recall of consistent physical distancing may be challenging, particularly if interactions are frequent and is further complicated if interactions were indoors or outdoors,³¹ something that was not asked in the current study.

Interpretation and potential implications of findings

This is the first study providing data obtained from testing of SARS-CoV-2 infection in the Nicaraguan paediatric population, presenting evidence that fills a major research gap. We accomplished this by using a simple, highly sensitive detection technique (LAMP), which may be used in developing countries both as a detection method and for epidemiological surveillance, with a growing field of application variations.^{32–37} LAMP is faster and less cumbersome than



qRT-PCR, and when used as a detection technique, LAMP has shown excellent sensitivity of up to 97.5% and specificity of up to 99.7% compared with qRT-PCR.^{38,39} Such a degree of accuracy, with a fast turnaround time, is important for low-income countries such as Nicaragua that rely on a single government-controlled testing laboratory. Although it should be noted that, for infections with very low viral loads below the limit of detection, the LAMP method may return indeterminate or false-negative results.⁴⁰ Furthermore, using saliva as the specimen to detect SARS-CoV-2 is a reliable method as it correlates well with clinical and immunological profiles.⁴¹ It is also less aerosol generating as compared with nasopharyngeal swabs or aspirate.

The high prevalence of SARS-CoV-2 infection found in children may point to widespread and sustained community transmission in Nicaragua. Our data highlight the importance of screening for SARS-CoV-2 in children, particularly in those who have an ill family member at home. In contrast to the rest of Central America, Nicaraguan authorities chose a path in complete opposition to the recommendations of the WHO, that is, leaving borders open, refusing to scale up testing and contact tracing and leaving all business and universities open.^{42,43} All through the pandemic, the Nicaraguan government kept public schools open, and there were no restrictions imposed on transport and mobility. Preventive measures such as wearing face masks, social distancing and avoiding direct contact with suspected COVID-19 cases were not actively promoted.^{43,44} It is critical that health authorities start implementing widespread frequent testing along with containment measures to diminish the widespread community transmission of SARS-CoV-2.

Limitations of the study

The study's data may not reflect the national prevalence of SARS-CoV-2 among Nicaraguan children since most children were from Managua, the capital of Nicaragua, although all individuals visiting these practices were eligible for inclusion and provide a fairly representative sample of children attending the participating clinics. Also, it is possible that some individuals were more likely to participate if they had a history of close contact with confirmed cases. However, a minority of people reported being in contact with individuals who were ill, suggesting this was not the primary motivating factor for participation. Results related to risk factors and exposures may be limited by the cross-sectional design. Children who had no symptoms at the time of clinic visit, who tested positive, may have developed symptoms later. Participant responses to surveys are subject to misclassification and recall bias. Recall bias is minimised, however, as neither the provider nor the participants were aware of their child's infection status at time of interview. Given the cross-sectional nature of the study, it is not possible to know if some children had lower chance of infection by attending school or if they had already been infected and recovered. While acknowledging these limitations, we conclude, based on the currently available data, that the prevalence of SARS-CoV-2 among Nicaraguan children was high during the time of this study.

Conclusions

To our knowledge, this is the first study to highlight the significantly high SARS-CoV-2 prevalence among Nicaraguan children, suggesting widespread and sustained community transmission, which further underscores the need for producing robust data on the true extent of SARS-CoV-2 infection in Nicaragua. With no available vaccine for children, the lack of testing is a key barrier to the fight against the virus in Nicaragua. We provide important evidence that is of global interest for policy implications to help inform optimum paediatric health service planning and effective non-pharmaceutical interventions to reduce the rate of infection and to improve the overall COVID-19 disease response.

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