

# Predictors of Illness Severity in COVID-19 Cases in Saudi Arabia

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**Purpose:** Multiple studies worldwide have reported the clinical and epidemiological features of coronavirus disease 2019 (COVID-19), with limited reports from the Middle East. This study describes the clinical and epidemiological features of COVID-19 cases in the Eastern Province of Saudi Arabia and identified factors associated with the severity of illness.

**Patients and Methods:** This was an observational study of 341 COVID-19 cases. These cases were reported in the first three months after the first case in the country was identified. Clinical and demographic data were analyzed and described to identify the effects of age, sex, and ethnicity on illness severity. In addition, the duration of viral shedding and cycle threshold (Ct) values of real-time PCR were evaluated as predictors of severity.

**Results:** The median age was 45 years. Males were twice as likely to be infected than females ( $p < 0.0001$ ). The duration of viral shedding ranged from 9 to 36 days. The most common clinical presentations include fever, shortness of breath, cough, myalgia, sore throat, vomiting, and headache. Critical cases were significantly higher in men (23% vs 8.7%), senior adults (>65 years), individuals of Bengali ethnicity, and in patients with comorbidities including diabetes, hypertension, and dyslipidemia ( $p = 0.001$ ). The case fatality rate was found to be 10%. The fatality was significantly higher in males than females (13.8% vs 2.6%), and in Asians (17.9%) than Arabs (6%) and Africans (0) ( $p = 0.002$ ). No association was found between viral load, represented by the RT-PCR cycle threshold (Ct) values, and severity of illness.

**Conclusion:** Age, sex, and ethnicity are important predictors of COVID-19 severity. The cycle threshold (Ct) of the SARS-CoV-2 RT-PCR test cannot be used as a predictor of the criticality of illness.

**Keywords:** factors, ethnicity, Ct value, Arabs, virus shedding

## Introduction

In late December 2019, a cluster of pneumonia of unknown etiology emerged in Wuhan, China<sup>1</sup> that was believed to be a zoonotic disease<sup>2,3</sup> linked to a local animal market.<sup>4</sup> A few weeks later, the disease showed sustained transmissibility among humans<sup>5</sup> and the etiology was identified as a novel member of the *Coronaviridae*, later referred to as the severe acute respiratory syndrome coronavirus-2 (SARS-CoV-2) due to its genetic homology (82%) to the previously identified member of the coronavirus family, SARS-CoV.<sup>4</sup> Soon after, the virus spread rapidly beyond China, signaling an impending pandemic. On January 30, 2020, the World Health Organization declared the novel coronavirus outbreak as a Public Health Emergency of International Concern (PHEIC).

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The disease caused by the virus was named the coronavirus disease-2019 (COVID-19).<sup>6</sup> Individuals infected with SARS-CoV-2 may be asymptomatic. More commonly, individuals present with a wide spectrum of clinical manifestations, ranging from mild illness to life-threatening pneumonia with a median incubation period of 6.4 days.<sup>7</sup> Symptomatic patients commonly present with fever, cough, fatigue, shortness of breath,<sup>8</sup> and sometimes gastrointestinal manifestations such as diarrhea, anorexia, nausea, vomiting, and abdominal pain.<sup>9</sup> Children have been shown to manifest milder features of the disease compared to adults.<sup>10</sup> However, some studies have suggested an increase in the incidence of multisystem inflammatory disease in children with COVID-19.<sup>11,12</sup> In addition, pregnant women and their newborns seem to be mildly affected by COVID-19 with no reports of teratogenicity or severe illness.<sup>13</sup> On November 26, 2020, the World Health Organization indicated that COVID-19 cases exceeded 59.8 million cases. Furthermore, multiple studies demonstrated that the virus was gradually evolving<sup>14,15</sup> and this genetic evolution was thought to impact the severity and infectivity of COVID-19.<sup>16,17</sup>

This study is the only study that described the epidemiological and clinical features of COVID-19 in the first 341 cases diagnosed at a university teaching hospital in the Eastern Province of Saudi Arabia over the first three months of the pandemic in the country. We identified the spectrum of clinical presentation in different age groups. Demographic data and comorbidities were also analyzed and correlated to identify predictors of severity and prognosis. Understanding and documenting the features of COVID-19 at the start of the spread of the pandemic in the kingdom is vital for better understanding of the effect of genetic evolution on the behavior of the virus, a story that must be told.

## Materials and Methods

A combined retrospective and prospective single-centered observational study was conducted at a university teaching hospital with 550 beds that serve patients from the Eastern Province of Saudi Arabia, as assigned by the Ministry of Health during the first wave of the COVID-19 pandemic. This study described the epidemiological and clinical features of COVID-19 for the initial 341 SARS-CoV-2 confirmed cases in the hospital (March 10 - June 13, 2020).

During the study period, 3146 individuals were tested for SARS-CoV-2 infection using nucleic acid detection tests, of which 341 were positive. The indications for testing included clinical suspicion of symptomatic COVID-19,

contact with confirmed cases, healthcare worker screening, and screening for all elective hospital admissions.

Testing for SARS-CoV-2 infection was performed using either the RealStar SARS-CoV-2 RT-PCR kit (Altona Diagnostics, Hamburg, Germany) or the Xpert Xpress SARS-CoV-2 kit (Cepheid, California, USA) according to the availability and following the manufacturer's recommendations. Samples from all positive cases were sent to the National Health Laboratory for confirmation. Cycle thresholds were retrieved from the designated software, as shown in the final report.

All laboratory-confirmed COVID-19 cases were included in this study. Clinical, demographic, and SARS-CoV-2 test results for all patients were obtained from a computerized hospital data system. Patients were stratified by age into groups: infants ( $\leq 1$  year), children ( $> 1$  years–19 years), adults (20–65 years), and senior adults ( $> 65$  years), for analysis according to the WHO classification. Patients were also grouped based on clinical presentation as described by the Saudi Ministry of Health into four groups: asymptomatic, mild to moderate, severe, and critical cases based on respiratory distress, oxygen requirements, and organ dysfunction. Moderate disease was defined by the presence of pneumonia. Severe cases were defined by respiratory rates measuring more than 30/min and 40/min, in adults and children, respectively, or arterial oxygen levels less than 90%. Critical cases were signified by the need for mechanical ventilation, or the occurrence of acute respiratory distress syndrome (ARDS), sepsis, or septic shock.<sup>18</sup> Serial RT-PCR results were recorded for all confirmed cases that were followed up for clearance using a test-based strategy. Clearance duration was defined as the duration between the first positive RT-PCR and the first negative RT-PCR sample, when the negative sample was confirmed by another negative RT-PCR sample collected  $\geq 24$  h later.

## Statistical Analysis

All data were tabulated using Microsoft Excel spreadsheets. Statistical analyses were performed using SPSS, version 24 (IBM Corp, Armonk, NY, USA). Multivariate linear regression analysis was first performed (Table 1), followed by post hoc analysis for each of the variables.

## Results

### Epidemiological Features of COVID-19

The total number of patients with laboratory-confirmed COVID-19 infections was 341. The male-to-female ratio

**Table 1** Multivariate Analysis of the Age Groups and Disease Severity with Other Demographic and Clinical Variables

	Value	F	Hypothesis Degrees of Freedom	Error Degrees of Freedom	Significance	Partial Eta Squared	Observed Power
Age groups	0.262	1.752	54.000	987.000	0.001	0.087	1.000
Disease severity	0.413	3.257	42.000	858.000	0.000	0.138	1.000

was 2:1 (225:116) ( $p < 0.0001$ ). The mean age of patients was  $44.19 \pm 18.03$  years (median = 45 years, range = 4 months to 96 years). There was no significant difference in the mean age between men and women ( $p = 0.958$ ).

Regarding the mode of acquisition of infection, 67.2% of patients ( $n = 229$ ) did not recall contact with a suspected or confirmed COVID-19 case, nor was there any clear epidemiological link with the source of infection. In contrast, 20.2% ( $n = 69$ ) reported positive household contact with COVID-19 cases, and 3.2% ( $n = 11$ ) reported contact with suspected cases not confirmed by laboratory tests. Only 7.3% ( $n = 25$ ) and 2.1% ( $n = 7$ ) of patients reported positive contact at work or travel history, respectively.

The case fatality rate (CFR), which is the proportion of deaths among cases, was 10% ( $n = 34$ ). Male patients exhibited a higher mortality rate (13.8%,  $n = 31$ ) than their female counterparts (2.6%,  $n = 3$ ), and this difference was statistically significant ( $\text{Chi}^2 = 10.68$ ,  $p = 0.001$ ). Furthermore, the CFR was higher among patients of Asian descent ( $n = 21$ , 17.9%) than in Arabs ( $n = 13$ , 6%) and Africans ( $n = 0$ ) ( $p = 0.002$ ). The mortality rate was highest among patients older than 65 years (24.32%), followed by patients aged between 41 and 65 years (12.84%).

### Clinical Features of COVID-19

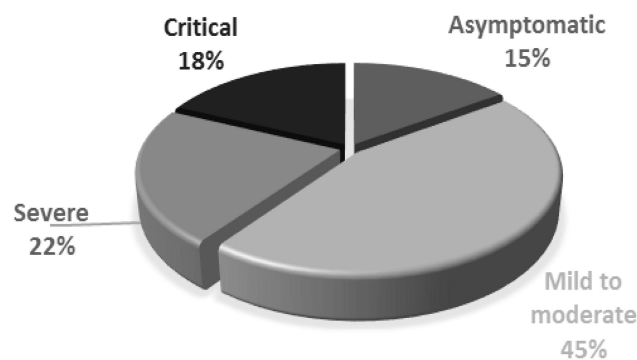
Of the 341 COVID-19 cases, 52 (15.2%) were asymptomatic. Of the symptomatic cases, 153 (44.9%) had mild to moderate disease, 74 (21.7%) had severe disease, and 62 (18.2%) were critically ill (Figure 1). Symptomatic patients presented with a wide variety of clinical manifestations, including systemic manifestations in the form of fever, headache, and myalgia, as well as respiratory symptoms, such as cough, shortness of breath, and rhinorrhea. Three patients reported a loss of smell and taste. Some patients also reported gastrointestinal manifestations, such as nausea, vomiting, anorexia, and diarrhea. Overall, the most common clinical features were fever, shortness of breath, cough, myalgia, vomiting, sore throat, and headache (Table 2).

All infant cases described in this study (5/5) showed fever, which was significantly higher in infants than in all other age groups.

COVID-19 was found to be more severe in males, senior adults, and patients of Bangladeshi nationality. On the contrary, the disease was less severe in women, infants, children, and individuals of Saudi and Egyptian nationalities (Table 3). Furthermore, patients with diabetes mellitus, hypertension, and dyslipidemia were more likely to have severe illnesses (Table 3). No significant association was found between bronchial asthma, heart disease, cancer, or sickle cell disease and COVID-19 severity (Table 3).

### Viral Shedding Dynamics and Predictors of Severe COVID-19

Of the 341 cases, one hundred and thirty-five (39.6%) confirmed cases underwent clearance following a test-based strategy, with other cases being followed on a time-based clearance. The duration of viral shedding ranged between 9 and 36 days, with a median of 13.5 days  $\pm$  5.66 (Figure 2). The association between the duration of shedding and severity of illness was not possible because asymptomatic cases followed time-based clearance with no repeated testing. No association was found between the cycle threshold (Ct values) of the Xpert Xpress SARS-CoV2 test and the severity of illness (Figure 3).

**Figure 1** The spectrum of the severity of illness among the 341 COVID-19 cases.

**Table 2** Clinical Features of COVID-19 Cases Between March and June 2020 According to Age Groups

Clinical Features	Infant ( $\leq 1$ Year)		Children ( $>1-19$ Years)		Adults ( $>19-65$ )		Senior Adults ( $65+$ )		Total N= 341	P- value*
	N=5	%	N=18	%	N=280	%	N= 38	%		
Fever	5	2.16	12	5.19	189	81.82	25	10.82	231	0.296
Shivering	0	0.00	0	0.00	22	95.65	1	4.35	23	0.936
Shortness of Breath	2	1.31	4	2.61	127	83.01	20	13.07	153	0.106
Chest pain	0	0.00	1	7.14	12	85.71	1	7.14	14	0.598
Wheezes	0	0.00	0	0.00	3	75.00	1	25.00	4	0.683
Cough	1	0.53	6	3.19	158	84.04	23	12.23	188	0.038
Sputum production	0	0.00	0	0.00	17	80.95	4	19.05	21	0.179
Hemoptysis	0	0.00	1	25.0	2	50.00	1	25.00	4	0.511
Rhinorrhea	0	0.00	3	30.0	6	60.00	1	10.00	10	0.057
Loss of taste/smell	0	0.00	0	0.00	3	100.00	0	0.00	3	0.470
Sore Throat	0	0.00	3	9.68	26	83.87	2	6.45	31	0.336
Headache	0	0.00	3	10.00	24	80.00	3	10.00	30	0.576
Myalgia	0	0.00	3	5.56	41	75.93	10	18.52	54	0.122
Vomiting	0	0.00	4	11.76	30	88.24	0	0.00	34	0.035
Diarrhea	1	2.63	1	2.63	32	84.21	4	10.53	38	0.823
Abdominal pain	0	0.00	1	14.29	6	85.71	0	0.00	7	0.168
Nausea	0	0.00	0	0.00	7	77.78	2	22.22	9	0.376
Anorexia	0	0.00	0	0.00	1	16.67	5	83.33	6	<0.001

**Note:** \*Significance is defined as  $p < 0.05$ .

## Discussion

This study of the first COVID-19 cases diagnosed in a university teaching hospital in eastern Saudi Arabia showed that out of 3146 individuals tested for SARS-CoV-2, 341 were positive. The overall detection rate was 10.8%. The majority of COVID-19 cases were detected in adults (aged 19–65 years), followed by senior adults (>65 years), which most probably reflects the frequency of COVID-19 diagnosis in the described settings rather than the incidence in various age groups. Furthermore, males were more commonly affected than females, which is consistent with a recent multivariate analysis study that identified male sex as an independent risk factor for the acquisition of COVID-19 infection.<sup>19</sup> However, this could also be attributed to local social factors, with males being more exposed by leaving homes to work and family commitments during the lockdown periods.

The mode of acquisition was not known in most cases, indicating the huge impact of asymptomatic spreaders of the disease and the need to improve contact tracing measures in Saudi Arabia. This may also indicate the possibility of contracting the virus through means other than close contacts, such as long-distance migrating micro-aerosols. Such transmission is gaining growing acceptance among experts in the field.<sup>20</sup> Our results support this mode of transmission as only one-fifth of our cases reported

household contact with COVID-19 confirmed cases and only 3% reported contact with suspected cases. Population-level studies are still limited, and most data regarding the source of infection available to date represent specific institutional experiences.

The case fatality rate (CFR) was found to be 10%, which is double that reported from Wuhan, China (5.6%).<sup>21</sup> There could be multiple reasons for the higher CFR, including the study setting. Our study involved a university hospital that was authorized to receive COVID-19 cases from other regional hospitals. This may lead to selection bias in the severity of cases included. Heterogeneity in ethnicity and other demographics may have also contributed to this difference. Importantly, the fatality rate does not take into consideration all individuals affected by SARS-CoV2 but only those who presented for testing (symptomatic individuals, contacts with known cases, or healthcare workers). This is also evident by the proportion of asymptomatic cases (15.2%) in our cohort, which is less than that described in many similar studies.<sup>22</sup> Thus, the overall infection mortality rate among individuals with COVID-19 infections is likely to be considerably lower. It has been estimated by the CDC and WHO to be around 0.5% - 1%.<sup>23,24</sup> On the other hand, the reported case fatality rates may have been underestimated considering that fatal COVID-19 infections may remain undiagnosed.<sup>25</sup>

**Table 3** The Distribution of COVID-19 Cases Based on Gender, Age, Co-Morbidities, and Nationality Relative to the Disease Severity

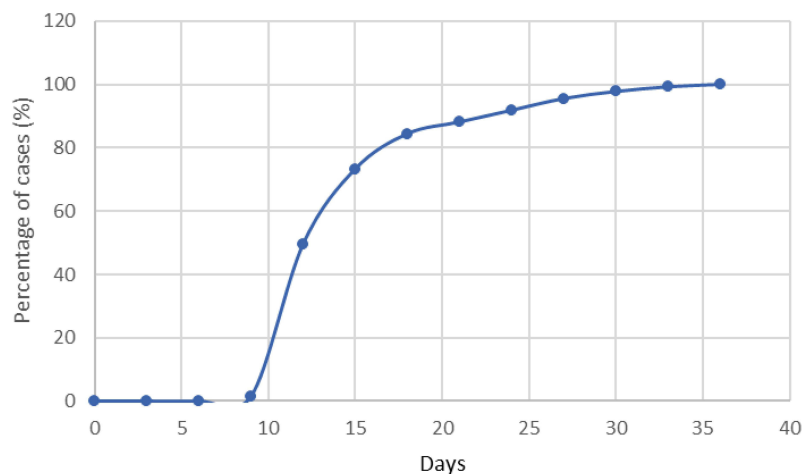
Variables	Asymptomatic		Mild to Moderate		Severe		Critical		Total	p-value*
	N 52	%	N 153	%	N 74	%	N 62	%		
<b>Gender</b>										
Female	22	19.30	60	52.63	22	19.30	10	8.77	114	<0.001
Male	29	12.83	93	41.15	52	23.01	52	23.01	226	
<b>Age groups</b>										
Infant (<=1 year)	2	40.00	2	40.00	1	20.00	0	0.00	5	0.08
Children (>1–19 years)	4	21.05	12	63.16	3	15.79	0	0.00	19	0.017
Adults (>19–65)	43	15.36	128	45.71	59	21.07	50	17.86	280	0.52
Senior adults (>65)	3	8.11	11	29.73	11	29.73	12	32.43	37	0.004
<b>Co-morbidity</b>										
Diabetes Mellitus	4	4.04	26	26.26	32	32.32	37	37.37	99	<0.001
Hypertension	5	6.17	23	28.40	32	39.51	21	25.93	81	<0.001
Smoking	0	0.00	1	20.00	3	60.00	1	20.00	5	0.23
Heart Failure	1	12.50	2	25.00	5	62.50	0	0.00	8	0.96
Cancer	1	16.67	3	50.00	0	0.00	2	33.33	6	0.97
Dyslipidemia	1	3.23	8	25.81	12	38.71	10	32.26	31	<0.001
CAD	1	14.29	4	57.14	1	14.29	1	14.29	7	0.55
CKD	5	14.71	8	23.53	8	23.53	12	35.29	34	0.23
Bronchial Asthma	0	0.00	4	57.14	3	42.86	0	0.00	7	0.84
SCD	0	0.00	4	80.00	1	20.00	0	0.00	5	0.44
<b>Nationality</b>										
Saudi	37	19.9	93	50.0	36	19.4	20	10.8	186	<0.001
Bahraini	0	0.0	0	0.0	1	100.0	0	0.0	1	–
Egyptian	7	43.8	4	25.0	4	25.0	1	6.3	16	0.025
Jordanian	0	0.0	0	0.0	0	0.0	2	100.0	2	0.05
Yemeni	1	11.1	4	44.4	1	11.1	3	33.3	9	0.56
Bengali	2	5.1	8	20.5	12	30.8	17	43.6	39	<0.001
Indian	1	2.5	23	57.5	7	17.5	9	22.5	40	0.26
Pakistani	0	0.0	10	52.6	4	21.1	5	26.3	19	0.18
Philippine	3	13.6	8	36.4	8	36.4	3	13.6	22	0.8
Somalian	0	0.0	1	100.0	0	0.0	0	0.0	1	–
Sudanese	1	20.0	1	20.0	1	20.0	2	40.0	5	0.52

**Note:** \*Significance is defined as  $p < 0.05$ .

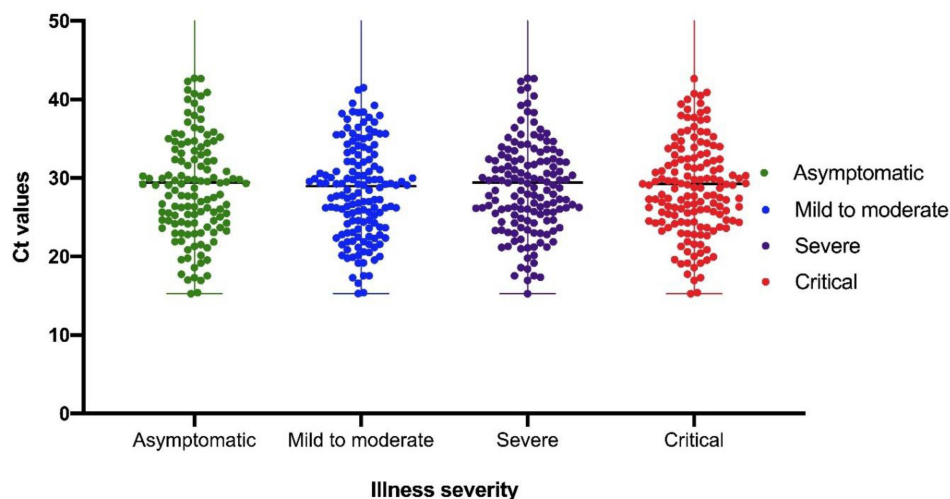
**Abbreviation:** SCD, sickle cell disease.

In our study, case fatality was significantly higher among males, which can be explained by a possible genetic predisposition and justified by the random and heterogeneous nature of the study sample. In contrast, nationality was a significant confounding factor similar to earlier reports.<sup>25,26</sup> In addition, fatality was also found to be affected by age. Higher fatality rates were detected in the older age groups. By ethnicity, fatality was higher in Asians, particularly Bengalis, compared to Arabs (17.9% to 6%) ( $p = 0.002$ ).<sup>27</sup> Similarly, the risk of severe disease and mortality has been shown to be higher in the African

American population in the United States, and attributed to the disproportional social determinants of health.<sup>28–30</sup> The impact of racial and ethnic disparities has been a concern during the COVID-19 pandemic. This is evident among healthcare workers (HCWs), who are at higher risk of acquiring the infection due to the nature of their jobs.<sup>31,32</sup> For instance, Hispanics constitute 17.6% of the total workforce in the United States, but they represent 25% of the healthcare aides who directly face continuous exposure.<sup>33</sup> Furthermore, living conditions, including air pollution, overcrowding, and delayed access to medical



**Figure 2** Dynamics of SARS-CoV2 viral clearance of 135 cases based on SARS-CoV-2 RNA detection.



**Figure 3** Frequency of cycle threshold (Ct) values for the E and N2 gene targets relative to COVID-19 severity (p value 0.65).

care among socially disadvantaged groups may contribute to this finding.<sup>34,35</sup>

Clinically, COVID-19 confirmed patients were classified into four categories based on the local national protocol.<sup>18</sup> The categories included asymptomatic, mild to moderate, severe, and critical. This classification was based on clinical features, evidence of pneumonia, oxygen requirements, and the presence of serious complications. In addition to respiratory symptoms, patients also reported gastrointestinal symptoms, such as vomiting, diarrhea, abdominal pain, nausea, and anorexia, which is in line with reports from different countries.<sup>36</sup> Fever was also significantly associated with age. In particular, fever was reported in 100% of infants, a feature that has been well documented in other studies.<sup>37</sup> It is interesting

that only three patients reported a lack of smell/taste in the study. Whether this is a feature of the disease in the region or a lack of reporting requires further investigation.

The severity of COVID-19 was significantly associated with male sex, senior adult age group, and Bengali ethnicity. Saudis and Egyptians (Arab ethnic groups) demonstrated a significantly lower severity of illness compared to the other groups. The severity of the disease in various patient populations needs to be interpreted with caution, taking into account the full burden of the pandemic, including delayed care and social determinants of health. Severity is also affected by comorbidities, particularly diabetes mellitus, hypertension, and dyslipidemia, as found in other studies.<sup>38</sup>

The cycle threshold (Ct) refers to the number of cycles in a PCR assay required to amplify viral nucleic acids to a detectable level. To the best of our knowledge, no clinical studies have validated the use of Ct to estimate the disease burden of COVID-19. Although the cycle threshold (Ct value) of the RT-PCR has often been considered a surrogate semi-quantitative measure for viral load estimation,<sup>39</sup> the correlation between the period of infectivity and culture positivity was not verified, and isolation of the virus from clinical samples was possible even in patients with high Ct values, indicating that infectivity does not correlate with Ct values.<sup>40,41</sup> In our study, the initial Ct value on presentation did not correlate with the severity of illness, which concurred with other studies.<sup>42</sup> This can be attributed to variability in the timing of sample collection concerning the onset of clinical features, since many cases with severe illness commonly present late. In contrast, asymptomatic contacts commonly present within a few days of exposure and demonstrate variable Ct values.<sup>42</sup> Furthermore, the Ct values are not standardized across various diagnostic platforms. Therefore, results cannot be compared between different assays. Given the uncertainty of the role of Ct values in estimating the relative viral RNA level in a specimen, our study does not support its use for guiding clinical decisions.

The minimum duration for viral clearance from respiratory samples in our cohort was 9 days, and the maximum reported duration was 36 days. The duration of viral shedding has been reported to be variable and influenced by the severity of the illness.<sup>43–46</sup> Nevertheless, detectable viral RNA does not necessarily correlate with the presence of live infectious virus, as more accumulative evidence suggests that viral RNA detected after 8 days is unlikely to be infectious.<sup>43</sup> Additionally, the virus has been detected in other types of specimens even after respiratory clearance, although this is unlikely to be of clinical or infection control significance.<sup>47,48</sup> Therefore, time-based strategies for clearance have been widely adopted in various institutions. Nonetheless, recent reports have described viral transmission 10 days later from severely infected patients. Therefore, caution should be considered in vulnerable patient populations.<sup>44,49</sup> Further data are needed to conclude the period of infectivity in cases of critical and severe illness.

Our study is not without limitations. Nevertheless, it contributes to the growing evidence of variable COVID-19 presentation and illness severity in relation to other factors

and provides insight into the usefulness of Ct values as a predictive clinical tool.

## Conclusion

Patients with COVID-19 manifest a wide range of clinical presentations, ranging from asymptomatic infection to severe and critical cases. Adult males were found to be twice as affected as females. The severity of COVID-19 was significantly associated with male sex, senior age group, and Asian ethnicity, particularly Bengali ethnicity. On the other hand, for the first time, milder forms of the disease were found to be significantly associated with Arab ethnic groups (Saudi and Egyptian), demonstrating the impact of ethnicity on disease severity. Furthermore, CFR was also found to be three times higher in Asians than in Arabs. The cycle threshold (Ct) of Xpert Xpress SARS-CoV2, a surrogate measure of viral load, was not found to correlate with the severity of illness and therefore, is not recommended for use as a predictor tool.

## Data Sharing Statement

All data analyzed in the study are available upon reasonable request from the corresponding author.

## Ethics Approval and Informed Consent

Ethical approval for the study was obtained from the Institutional Review Board at Imam Abdulrahman Bin Faisal University (IRB-2020-01-150). Written patient consent was not requested by the IRB committee due to the retrospective nature of the study and that no patients identifiable information is presented. Patients' confidentiality is maintained throughout the study and the study doesn't carry any additional risk to patients. Furthermore, participation in the study did not interfere with patients management and all investigations included and analyzed were part of patients management as per the hospital protocols.

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## Disclosure

The authors report no conflicts of interest in this work.

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