RESEARCH

Open Access



Comparable detection of nasopharyngeal swabs and induced sputum specimens for viral nucleic acid detection of suspected novel coronavirus (SARS-Cov-2) patients in Fayoum governorate, Egypt

Doaa Y. Ali^{1*}, Rasha A. Hussein², Shahira Morsy ELshafie¹, Reem Amgad Mohamed¹ and Fadwa Abd El Reheem¹

Abstract

Background The most commonly utilized samples for severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) detection using real-time quantitative reverse transcriptase-polymerase chain reaction (RT-qPCR) are nasopharyngeal swabs (NPS) and oropharyngeal swabs. However, there are some drawbacks. For SARS-CoV-2 detection, induced sputum might be analyzed and may be equivalent to pharyngeal swabs. This study was done to assess the potential superiority of induced sputum over NPS for SARS-CoV-2 detection. Sixty symptomatic COVID-19 patients who attended Fayoum University Hospitals in Fayoum Governorate, Egypt, were included in this cross-sectional descriptive study. Paired NPS and induced sputum samples were collected from each subject on the third and tenth days after symptoms began for RT-qPCR SARS-COV2 diagnosis.

Results At day 3, 52 (86.7%) of NPS and 48 (80.00%) of induced sputum specimens had positive RT-qPCR results with a significant statistical difference (P=0.001). At day 10, 41 induced sputum samples (68.3%) were negative, while 19 (31.7%) were positive. Only three (5.0%) of the 19 positive induced sputum samples tested positive for NPS. NPS samples had a higher viral load than induced sputum samples at day 3 [25 (41.7%) vs. 23 (38.3%)]. At day 10, induced sputum samples had a higher viral load than NPS [9 (15.0%) vs. 6 (10.0%)]. A statistically significant positive correlation between the viral load value of the NPS and the induced sputum sample at day 3 (r=0.497, p=0.00) denoting similarity in the results of the two types of samples. By ROC analysis, the highest area under the curve for the overall CT value of the induced sputum was (0.604), with a statistically significant difference (p value=0.0418).

Conclusion In the early stages of the disease, induced sputum and NPS tests had comparable results, but NPS yielded more false negative results later in the disease course than an induced sputum sample, which yielded higher sample positivity and viral load than NPS. Furthermore, induced sputum collection is a straightforward, non-invasive, and risk-free method. As a result, induced sputum could be useful for COVID-19 confirmation in patients with radio-logically or epidemiologically suspected COVID-19 who have a negative NPS or in difficult-to-diagnose COVID-19 patients.

*Correspondence: Doaa Y. Ali younisdoaa@yahoo.com Full list of author information is available at the end of the article



© The Author(s) 2023. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/.

Keywords Induced sputum, Nasopharyngeal swab, SARS-CoV-2, RT-qPCR, Viral load

1 Background

The World Health Organization (WHO) has identified the '2019 new coronavirus,' or 'COVID-19,' as the primary cause of the current pneumonia epidemic, which began in early December 2019 in Wuhan, Hubei Province, China [1]. On February 14, 2020, Egypt reported its first case of coronavirus disease 2019 (COVID-19) [2].

Coronaviruses belong to the Nidovirales order, the Coronaviridae family, and the Coronavirinae subfamily of the Nidovirales order. Based on serological data, the Coronavirinae are divided into four families: alphacronavirus, betacoronavirus, gammacoronavirus, and delta coronavirus [3]. Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), the agent responsible for COVID-19, is a species of the genus Betacoronavirus [3].

The gold standard approach for detecting SARS-CoV-2 has always been probe-based Real-time quantitative reverse transcriptase-polymerase chain reaction (RT-qPCR), which the Centers for Disease Control and Prevention (CDC) and WHO recommend for population screening globally [4].

The RNA-dependent RNA polymerase (RdRp) gene, envelope (E) gene, spike (S) gene, nucleocapsid (N) gene, and ORF8 or ORF1b portions of the SARS-CoV-2 genome were all targeted using RT-qPCR assays [5].

For a Covid-19 diagnosis, the upper respiratory tract is sampled from the nasopharyngeal (NPS), oropharyngeal swab (OPS), NP wash, or saliva, while the lower respiratory tract is sampled from sputum, tracheal aspirate, bronchoalveolar lavage fluid (BAL), or bronchoscopic brushing [6, 7]. Which are considered to be even more sensitive than upper respiratory tract samples [6].

The most commonly utilized samples for COVID-19 detection using RT-qPCR are NPS and OPS. However, there are some drawbacks, such as the false-negative findings in upper respiratory samples obtained from asymptomatic patients or mild illnesses, as well as the necessity for repeat sampling and testing [7]. Negative findings do not rule out the possibility of COVID-19 infection [8]. Nasopharyngeal RT-PCR positivity is thought to decline within one week of symptom onset; therefore, a positive test late in the course of the disease is expected to be from sputum, BAL [9]. So, the need for appropriate specimen selection is critical for increasing SARS-Cov-2 detection using the RT-qPCR technique and lowering existing false-negative detection. As a result, searching for alternative specimen types with greater precision and detection accuracy is required.

Sputum is a helpful, non-invasive technique for the detection of SARS-CoV-2. It might identify SARS-CoV-2 at a greater rate than NPS or throat swabs [10, 11]. A possible drawback of testing sputum samples for COVID-19 diagnosis is that not all patients infected with COVID-19 can expectorate sputum, and it is restricted to patients who can introduce sputum. According to certain research, low sputum production is a prevalent symptom in COVID-19 patients, making it difficult to get sputum in these individuals [12, 13]. So, for SARS-CoV-2 detection, induced sputum or a self-collected deep cough specimen might be analyzed, especially in those who are uncapable of introducing sputum, and may be equivalent to pharyngeal swabs [14, 15]. Sputum induction using a hypertonic saline solution is one of the most commonly used methods for studying airway secretions in patients with lung disorders [16], such as chronic obstructive pulmonary disease, bronchial asthma, and pulmonary hypertension [17]. Recently, in Mycoplasma pneumoniae and infantile tuberculosis [18].

The safety and effect of induced sputum for SARS-CoV-2 detection had previously been reported in some multi-center cross-sectional studies [19, 20]. Therefore, we conducted this study to further assess the potential superiority of induced sputum over NPS for SARS-CoV-2 detection.

2 Method

2.1 Study design

This is a cross-sectional descriptive study that was conducted on sixty symptomatic COVID-19 patients who attended Fayoum University Hospitals, Fayoum Governorate, Egypt, from April 30 to October 30, 2021.

The study follows the Declaration of Helsinki guidelines. In addition, informed written consent was obtained from the legal guardians of all subjects participating in this study. The severity classification of our cases was according to WHO interim guidance into mild, moderate, severe, and critical cases [21].

We included in our study all suspected COVID-19 patients of both genders. While, we excluded asymptomatic patients, those diagnosed with SARS-CoV-2 infection via a single type of specimen (induced sputum or NPS), and any ventilated patients who could not give induced sputum samples.

All patients were subjected to a full medical history and clinical examination including the presence of any respiratory distress signs. On the third day after the onset of symptoms, all patients had a computed tomography of the chest (CT-Chest). Depending on the CT result, the patients were classified according to the coronavirus disease 2019 (COVID-19) Reporting and Data System (CO-RADS) classification from CO-RAD1 to CO-RAD6 and according to the total severity score (TSS) into minimal, mild, moderate, and severe degrees [22].

2.2 Laboratory investigations

The following laboratory investigations were done on all patients on the first day of the onset of the symptom: Full blood examination including: total leucocytic count (TLC), platelets count, and hemoglobin (HB) (All were done on Sysmex XN 1000, Canada), C-reactive protein (CRP) was done by automated CRP instrument (CoaD-ATA 4004 instrument,Germany), D-dimer was measured using a chemiluminescent enzyme immunoassay quantitative technique by (Path fast compact immune-analyzer, Japan).

2.3 Samples

Each subject was asked to give paired samples of NPS and induced sputum at the 3rd and 10th days of the beginning of symptoms for SARS- COV-2 laboratory diagnosis by RT-q PCR. NPS were collected using specialized dacron, rayon, or calcium alginate-tipped collection swabs with plastic or non-aluminum wire shafts. Samples were collected in specific vials containing viral transport media (VTM) (STOR-F (DNA technology, Russia).

Each patient inhaled a 3% saline solution by nebulizer to induce sputum. In order to decrease the risk of oral contamination before induction of sputum, a saline solution was used to rinse the mouth, and then the patients were asked to give a deep cough and introduce sputum into a screw-capped sterile container. We followed the CDC's guidelines for collecting, handling, and testing COVID-19 different clinical specimens [23].

2.4 RT-qPCR detection of SARS-COV-2

The molecular diagnosis of COVID-19 was done in the molecular biology unit in the Clinical and Chemical Pathology Department at Fayoum University Hospitals. Covid-19 RNA extraction was done by a nucleic acid extraction kit ("DNA- Technology ' made PREP-NA DNA/RNA) using Lab Turbo 48 C automated extraction system (Tiagen Bioscience Corporation, Taiwan). The SARS-CoV-2/SARS-CoV Multiplex REAL-TIME PCR Detection Kit is used to amplify and detect the SARS-COV-2 target region (N gene and ORF gene).

The following were RT-qPCR thermocycling conditions: for twenty minutes at 35 °C and for five minutes at 95 °C, fifty amplification cycles at 94 °C for ten seconds and 64 °C for fifteen seconds, then for one minute at 80 °C using DT Lite thermocycler (DNA Technology Research & Production, LLC, Russia). The quantification of viral nucleic acid in patient samples was done by measuring the RT-PCR cycle threshold (CT) of the ORF gene. The CT represents the number of replication cycles required to produce a fluorescent signal. The viral load is reflected by the CT number. CT value less than 25 indicates high viral load; CT value 25 to 35 indicates moderate viral load; and CT value greater than 35 indicates low viral load. To ensure validation of the result, an internal control is measured parallel to each sample, and positive and negative controls were used in each PCR run.

2.5 Statistical analysis

Our study data was statistical analysis using SPSS (Social Science version 28.00). Descriptive analyses as frequency and percent for qualitative data and for quantitative data as median and interquartile range (IQR). Non-parametric quantitative data analysis was done using the Kruskal–Wallis test and the Mann–Whitney test. A qualitative analysis of associations between variables was compared with the Chi-square test. The receiver operating characteristic (ROC) and Spearman correlation coefficient (r) between the two variables were done. Statistical significance level at a *p* value ≤ 0.05 .

3 Results

3.1 Demographic, clinical, and radiological data of the studied patients

This descriptive cross-sectional study included 60 clinically and laboratory-suspected COVID-19 patients, whose ages ranged from 15 to 72 years with a median age of 32 years. Twenty-one patients (35.0%) were male. Thirty-nine patients (65.0%) were female. Regarding COVID-19 risk factors, there were 7 patients (11.7%) with both diabetic mellitus (D.M.) and hypertension (HTN), while only 2 patients (3.3%) presented with chronic liver disease. The median of O2 saturation was 96.0%; it ranged from 80.0% to 99.0%. Thirty six patients (60.0%) had an abnormal CT finding. According to TSS, there were 7 cases (11.7%) in each mild and sever degree, and 10 cases (16.7%) had a moderate degree. The clinical data of our cases were detailed in (Table 1).

According to the COVID-19 severity classification, we classified patients into four groups: 35 cases (58.3%) were mild, 16 cases (26.7%) were moderate, and 9 cases (15.0%) were severe. No critical cases were presented in our study.

Regarding the laboratory results in our study, the medians of TLC, lymphocytic count, HB, platelet count, CRP, and D-dimer were (6.22, 2.10, 12.55, 234.50, 5.50, and 0.30) respectively (Table 2).

Age	Range	15.00-72.00	15.00-72.00 32.00 (25.00 45.00)		
	Median (IQR)				
		Number	Percent (%)		
Sex	Male	21	35.0		
	Female	39	65.0		
HCW	Non HCW	30	50.0		
	HCW	30	50.0		
HTN	Non-HTN	53	88.3		
	HTN	7	11.7		
DM	Non DM	53	88.3		
	DM	7	11.7		
CLD	Non-hepatic patients	58	96.7		
	Hepatic patients	2	3.3		
Clinical symptoms		Number	Percent (%)		
Fever	Absent	11	18.3		
	Present	49	81.7		
Cough	Absent	13	21.7		
	Present	47	78.3		
Dyspnea	Absent	45	75.0		
	Present	15	25.0		
Anosmia	Absent	46	76.7		
	Present	14	23.3		
Diarrhea	Absent	54	90.0		
	Present	6	10.0		
CT. Finding classification according to TSS	No CT. finding	36	60.0		
	Mild	7	11.7		
	Moderate	10	16.7		
	Severe	7	11.7		
O2 saturation%	Range	80.0–99.0%			
	Median (IQR)	96.00% (94.00–97.00)			

Table 1 Demographic, clinical, and radiological data of the studied patients

Total Number = 60

HCW healthcare workers, DM diabetes mellitus, HTN hypertension, CLD chronic liver disease, C.T computed tomography, TSS total severity score, IQR Inter quartile range.

At day 3 of the beginning of the symptoms, the majority of the patients had positive RT-qPCR results, with 52 (86.7%) of their NPS and 48 (80.0%) of their induced sputum specimens being positive (Table 3). However, six patients (10.0%) showed positive NPS and negative induced sputum samples, while only two patients (3.3%) showed negative NPS and positive induced sputum samples (Table 4). The findings showed that positive rates displayed a highly significant statistical difference (P=0.001) between NPS and induced sputum specimens at day 3.

While, at day 10, 41 induced sputum samples (68.3%) were negative, while 19 (31.7%) were positive. Fourteen

cases (23.3%) were positive by NPS (Table 3). From 19 positive induced sputum samples, only 3 cases (5.0%) were positive by NPS (Table 4). There was a statistically non-significant difference (P=0.347) between the RT-qPCR results of NPS and induced sputum specimens at day 10.

The median (IQR) for the NPS CT value at day 3 and at day 10 were (26.25 and 42.00), respectively, and for induced sputum samples at day 3 and day 10 were (27.25 and 41.00), respectively. The cycle threshold (CT) values of the NPs and induced sputum samples were non-significantly different at days 3 and 10 (p = 0.308 and 0.551, respectively) (Figs. 1, 2).

Table 2 Laboratory investigations of the studied patients

TLC ($\times 10^{9}$ per L)	
Range	2–25.9
Median (IQR)	6.22 (4.72–7.77)
Lymphocyte count (× 10 ⁹ per L)	
Range	0.55-4.50
Median (IQR)	2.10 (1.29–2.50)
Hb (g/dL)	
Range	9.00-16.30
Median (<i>IQR</i>)	12.55 (11.50–13.57)
Platelet (× 10^9 per L)	
Range	83.00-610.00
Median (IQR)	234.500 (200.00–292.50)
CRP (mg/L)	
Range	0.500-134.90
Median (<i>IQR</i>)	5.50 (3.00–36.22)
D-dimer (µg/mL)	
Range	0.10-5.00
Median (<i>IQR</i>)	0.30 (0.20–0.57)

TLC total leucocytic count, *H.B* hemoglobin, *CRP* C-reactive protein, *IQR* interquartile range

 Table 3
 Distribution of RT-qPCR results on NPS and induced sputum samples

RT-qPCR results	Number	Percent (%)
NPS at day 3		
Negative	8	13.3
Positive	52	86.7
Induced sputum samples at day 3		
Negative	12	20.0
Positive	48	80.0
NPS at day 10		
Negative	46	76.7
Positive	14	23.3
Induced sputum samples at day 10		
Negative	41	68.3
Positive	19	31.7

Total N = 60

RT-qPCR real-time quantitative reverse transcriptase-polymerase chain reaction, NPS nasopharyngeal swabs

Regarding the viral load results, NPS samples had a higher viral load than induced sputum samples at day 3 [25 (41.7%) vs. 23 (38.3%)]. In contrast, at day 10, the results of the high viral load of induced sputum samples

were superior to those of NPS [9 (15.0%) vs. 6 (10.0%)] (Table 5).

The receiver operating characteristic (ROC) curve was drawn to compare the overall diagnostic performance of NPS and induced sputum specimens during the study. By ROC analysis, we found that the highest area under the curve (AUC) for the overall CT value of the induced sputum samples was (0.604), which was greater than that of NPS (0.548) (Fig. 3), with a statistically significant difference (p value = 0.0418). The sensitivity of induced sputum samples was 64.0% (95% CI, 49.2–77.1), the specificity was 61.43% (95% CI, 49.0–72.8), the positive predictive value (NPV) was 70.5%. (Table 6; Fig. 3).

By applying Spearman's correlation test, there was a statistically significant positive correlation between viral load according to the CT value of the NPS and the induced sputum samples at day 3 (r=0.497, p=0.000), denoting similarity in the results of the two types of samples (Fig. 4).

4 Discussion

The real-time reverse transcription-polymerase chain reaction assay (RT-q PCR) is the most important method for laboratory confirmation of SARS-CoV2. Nasopharyngeal swab remains the most common samples for SARS-COV-2 diagnosis but it has some limitations and easy to miss the diagnosis [7, 8]. Currently, there are more instances of false negative nucleic acid testing findings and released patients who turn positive again are increasing [24]. As a result, it may be necessary to reconsider whether the patients are virally free. SARS-CoV-2 detection through RT-qPCR might be enhanced by improving specimen selection, which is important to minimize the number of false negative results.

Therefore, looking for other specimen types with greater accuracy and detection efficiency is necessary. So, we conducted this research to assay the potential superiority and the accuracy of induced sputum over NPS for viral nucleic acid detection of novel coronavirus (SARS-Cov-2). This study was done at Fayoum University Hospitals and included 60 patients who fulfilled the inclusion criteria and formed the study population. We measured the viral load by nucleic acid assays (RT-qPCR) in paired samples (Nasopharyngeal swabs and induced sputum samples) early on day 3 to ensure having samples with a high viral load, and on day 10 to confirm that patients are free from infection and can return to their work; this is according to the "symptom-based strategy" of the CDC [25].

Regarding detection of the viral nucleic acid by RTqPCR in NPS and induced sputum samples on the

Table 4 Comparison of RT-qPCR results between NPS and induced sputum samples at day 3 and day 10

	Induced Sp	outum at day 3			Chi-square	<i>P</i> -value
	Negative		Positive	Positive		
	N	%	N	%		
NPS at day 3						
Negative	6	10.0	2	3.3	17.452	0.001 HS
Positive	6	10.0	46	76.7		
	Induced Sp	outum at day 10			Chi-square	P-value
	Negative		Positive			
	N	%	N	%		
NPS at day 10						
Negative	30	50	16	26.7	0.885	0.347 NS
Positive	11	18.3	3	5.0		

NPS nasopharyngeal swabs, P-value < 0.05 was considered statistically significant

NS non-significant at p-value > 0.05, HS high-significant



Fig. 1 A boxplot comparing NPS and induced sputum samples at day 3 based on the cycle threshold (CT) value. Legend: Data are expressed as a box plot. The median (Inter quartile range) for the cycle threshold (CT) value of NPS at day 3 was (26.25) and for induced sputum samples was (27.25)



Fig. 2 A boxplot comparing NPS and induced sputum samples at day 10 based on the cycle threshold (CT) value. Legend: Data are expressed as a box plot. The median (Inter quartile range) for the cycle threshold (CT) value of NPS at day 10 was (42.00) and for induced sputum samples was (41.00)

third day of symptoms, 86.7% of cases had positive NPS, and 80.0% had positive induced sputum samples. A few patients (10.0%) had positive NPS and negative induced sputum samples. The positive rate of sputum samples on the 3rd day was lower than that of NPS, with a highly significant statistical difference (P=0.001).

This finding mismatched with Lin et al. [26] who found that the positive rate of sputum specimens was higher than that of nose and throat swabs (76.9% and 44.2% respectively).

This may be explained by the slightly high percentage of mild cases (58.3%) in our study who presented most

 Table 5
 Viral load classification of NPS and induced sputum samples at day3 and day10

	Number	Percent (%)
Viral load of NPS at day 3		
High viremia	25	41.7
Moderate viremia	27	45.0
Low viremia	8	13.3
Viral load of induced sputum samples at day3		
High viremia	23	38.3
Moderate viremia	24	40.0
Low viremia	13	21.7
Viral load of NPS at day 10		
High viremia	6	10.0
Moderate viremia	8	13.3
Low viremia	46	76.7
Viral load of induced sputum samples at day10		
High viremia	9	15.0
Moderate viremia	10	16.7
Low viremia	41	68.3

NPS nasopharyngeal swabs

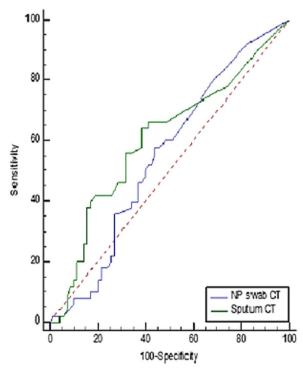


Fig. 3 Receiver operating characteristic (ROC) curve for overall cycle threshold (CT) value of NPS and induced sputum samples. Legend: By ROC analysis, the area under the curve (AUC) for the overall CT value of the induced sputum samples was 0.604, with the best cutoff value at \leq 32, sensitivity = 64.0% (95% Cl, 49.2–77.1), specificity = 61.43% (95% Cl, 49.0–72.8), and a *p* value = 0.0418. The AUC for the overall CT value of NPS was 0.548, with the best cutoff value at \leq 31.9, sensitivity = 58.0% (95% Cl, 43.2–71.8), specificity = 55.71% (95% Cl, 43.3–67.6), and *p* value = 0.3651

frequently with upper respiratory symptoms of COVID-19 infection rather than lower respiratory ones. Although sputum is one of the lower respiratory tract samples, it gives a better positive result in a patient with lower respiratory tract infections. Also, in the Lin et al. [26] study, the average age of the study population was 57.3 years. As a result, elderly patients presented with more severe forms of the disease, including lower respiratory tract infections, while the median age of our study population was 32 years old.

While the results of viral nucleic acid by RT-qPCR in NPS and induced sputum samples on the 10th day of symptoms showed a higher positive rate in induced sputum samples than NPS [19 (31.7%), 14 (23.3%)], only three (5.0%) of the 19 positive induced sputum sample cases were positive by NPS, while the remaining 16 (26.7%) had positive induced sputum and negative NPS results. This is most likely owing to the greater prevalence of angiotensin converting enzyme-2 (ACE2) receptors in pneumocytes and epithelial cells of the lower respiratory airway relative to upper airway epithelial cells. That had been recognized as the SARS-CoV-2 functional receptor [27, 28]. This finding agrees with Liu et al. [29] who found longer SARS-CoV-2 detection times in sputum samples compared to NPS samples. According to Zhang et al. [30] the findings were also similar.

Early in the disease, results of high viremia were found more in NPS than in induced sputum samples (41.7% vs. 38.3%, respectively). In contrast, at day 10, more cases with high viral loads were seen in induced sputum than in NPS (15.0% vs. 10.0%, respectively). The length of time between virus shedding may vary from part of the respiratory tract to another [30, 31]. For example, in the upper respiratory tract specimens, peak levels of SARS-CoV-2 were seen very early in the course of the disease [31]. Moreover, another study reported that viral shedding occurs over a longer period of time in lower respiratory tract secretions and that the maximum viral loads appear about 2 weeks after the onset of symptoms [32]. In these regions, the different expression levels of ACE2, the putative cell entry receptor of SARS-CoV-2, may partly explain this phenomenon [27, 28]. Meanwhile, the lower respiratory tract communicates less with the outside world than the upper respiratory tract, exacerbating viral retention [33, 34].

Regarding the median CT value on the 3rd day, we found an insignificant difference between NPS results and induced sputum results (p value = 0.308). Our result showed agreement with Liu et al. [29] who compared SARS-CoV-2 CT values in sputum and NPS samples with and without underlying diseases, and the p values were 0.65 and 0.22, respectively [29].

Cutoff point	AUC (95% Cl)	P-value	Sensitivity (95% Cl)	Specificity (95% Cl)	PPV (%)	NPV (%)
Overall CT of NPS						
≤31.9	0.548 (0.444–0.651)	0.3651	58.0% (43.2-71.8)	55.71% (43.3–67.6)	48.3	65.0
Overall CT of indu	iced sputum samples					
≤32	0.604 (0.499–0.709)	0.0418	64.0% (49.2–77.1)	61.43% (49.0–72.8)	54.2	70.5

Table 6 Overall diagnostic performance of NPS and induced sputum samples

AUC area under the curve, CT cycle threshold, CI confidence interval, PPV Positive predictive value, NPV negative predictive value

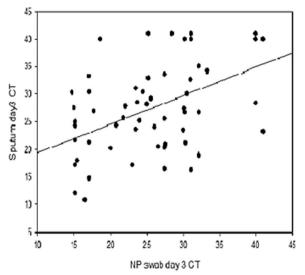


Fig. 4 Linear correlation between the cycle threshold (CT) value of NPS and induced sputum samples at day 3. Legend: A correlation analysis was done between CT values of NPS and induced sputum samples at day 3, with a significant positive correlation (r = 0.497, p = 0.000)

By using the ROC curve, we found that the overall diagnostic performance of induced sputum was relatively greater than NPS with a statistically significant difference (*p* value=0.0418), The sensitivity, specificity, PPV, and NPV of induced sputum were 64.0%, 61.43%, 54.2%, and 70.5%, respectively, compared to NPS (58.0%, 55.71%, 48.3%, and 65.0%, respectively). This was matched with the findings of Lai et al. [35], who discovered that induced sputum samples outperformed oropharyngeal swabs in terms of sensitivity and specificity (85.5% and 79.1%, respectively). There was a statistically significant positive correlation between viral load according to the CT values of both samples at day 3 (r=0.497, p=0.000).

5 Conclusion

Induced sputum and NP swab tests had comparable results. In terms of diagnosing COVID-19, NPS was slightly superior to sputum early in the disease course, yielding more positive results. Later in the disease course, an induced sputum sample yielded higher sample positivity and viral load than NPS. Furthermore, induced sputum collection is a straightforward, non-invasive, and risk-free method.

We believe that sputum might aid in identifying COVID-19 in sputum-producing patients. Therefore, induced sputum could be useful for COVID-19 confirmation in patients with radiologically or epidemiologically suspected COVID-19 who have a negative NPS or in difficult-to-diagnose COVID-19 patients. Negative induced sputum should be used as a criterion for hospital discharge for COVID-19 recovering patients and release from quarantine. However, this study does not fully invalidate the need for nasopharyngeal sampling but rather recommends a noninvasive, relatively simple collection method with apparent advantages, such as induced sputum samples.

The small sample size, single-center study, and restriction of sample types to NPS and induced sputum samples were the study's limitations, as other samples would have been better tested. As an example, saliva is simple, quick, and safe to collect. Moreover, new research has indicated that saliva has some advantages in COVID-19 diagnosis [14, 15, 36].

Abbreviations

CDC	Centers for Disease Control and Prevention
COVID-19	Novel coronovirus disease for 2019
CRP	C-reactive protein
CT	Cycle threshold
HB	Hemoglobin
NPS	Nasopharyngeal swabs
ROC	The receiver operating characteristic
RT-qPCR	Real-time quantitative reverse transcriptase-polymer-
	ase chain reaction.
SARS-CoV-2	Severe acute respiratory syndrome coronavirus 2
TLC	Total leucocytic count
WHO	The World Health Organization

Acknowledgements

The authors are highly thankful to the chest disease doctors at Fayoum University Hospitals for their help in the clinical diagnosis of our cases. Also, we wish to thank the radiological disease specialists at Fayoum University Hospitals who interpret the radiological data of our study group and the anonymous reviewers for their perceptive comments on the manuscript.

Author contributions

All authors contributed to the study conception and design. Material preparation, data collection, and analysis were performed by DYA, RAH, RAM, FA. SM supervised data. DYA was a major contributor in writing the manuscript. All authors read and approved the final manuscript.

Funding

No funding is obtained for this study.

Availability of data and materials

All data generated or analyzed as part of this study are included in this published article.

Declarations

Ethics approval and consent to participate

Approval by the ethical committee of the Faculty of Medicine, Fayoum University, was obtained before the commencement of the study (the approval number for the ethical committee is M535; Session: 81; Date: April 11, 2021). Informed written consent was obtained from the legal guardians of all subjects participating in this study.

Consent for publication

Not applicable.

Competing interests

The author declares that there is no conflict of interest.

Author details

¹Department of Clinical and Chemical Pathology, Faculty of Medicine, Fayoum University, Fayoum, Egypt. ²Department of Medical Microbiology and Immunology, Faculty of Medicine, Zagazig University, Zagazig, Egypt.

Received: 1 March 2023 Accepted: 24 April 2023 Published online: 02 May 2023

References

- 1. Hui DS, Azhar El, Madani TA, Ntoumi F, Kock R, Dar O, Ippolito G, Mchugh Z, Memish TD, Drosten A, Zumla A (2020) The continuing 2019-nCoV epidemic threat of novel coronaviruses to global health. The latest 2019 novel coronavirus outbreak in Wuhan, China. Int J Infect Dis 91:264–266. https://doi.org/10.1016/j.ijid.2020.01.009
- WHO (2020) Health Emergency Dashboard. https://covid19.who.int/ region/emro/country/eg. Publisher
- Fehr AR, Perlman S (2015) Coronaviruses: an overview of their replication and pathogenesis. Methods Mol Biol 1282:1–23. https://doi.org/10.1007/ 978-1-4939-2438-7_1
- Zhang H, Hong C, Zheng Q, Zhou P, Zhu Y, Zhang Z, Bi Q (2020) Ma T (2020) A multi-family cluster of COVID-19 associated with asymptomatic and pre-symptomatic transmission in Jixi City, Heilongjiang, China. Emerg Microbes Infect 9(1):2509–2514. https://doi.org/10.1080/22221751.2020. 1837015
- Reusken CBEM, Broberg EK, Haagmans B, Meijer A, Corman VM, Papa A, Charrel R, Drosten C, Koopmans M, Leitmeyer K, On Behalf of Evd-LabNet And Erli-Net (2020) Laboratory readiness and response for novel coronavirus (2019-nCoV) in expert laboratories in 30 EU/EEA countries, January 2020. Euro Surveill 25(6):2000082. doi:https://doi.org/10.2807/1560-7917. ES.2020.25.6.2000082
- Bwire GM, Majigo MV, Njiro BJ, Mawazo A (2021) Detection profile of SARS-CoV-2 using RT-PCR in different types of clinical specimens: a systematic review and meta-analysis. J Med Virol 93(2):719–725. https:// doi.org/10.1002/jmv.26349
- Mawaddah A, Gendeh HS, Lum SG, Marina MB (2020) Upper respiratory tract sampling in COVID-19. Malays J Pathol 42(1):23–35
- -Torretta S, Zuccotti G, Cristofaro V, Ettori J, Solimeno L, Battilocchi L, D'Onghia A, Bonsembiante A, Pignataro L, Marchisio P, Capaccio P (2021) Diagnosis of SARS-CoV-2 by RT-PCR using different sample sources: review of the literature. Ear Nose Throat J 100(2_suppl):131S–138S. https://doi.org/10.1177/0145561320953231
- 9. Sethuraman N, Jeremiah SS, Ryo A (2020) Interpreting diagnostic tests for SARS-CoV-2. JAMA 323(22):2249–2251. https://doi.org/10.1001/jama. 2020.8259

- Wang W, Xu Y, Gao R, Lu R, Han K, Wu G, Tan W (2020) Detection of SARS-CoV-2 in Different Types of Clinical Specimens. JAMA 323(18):1843–1844. https://doi.org/10.1001/jama.2020.3786
- Guo WL, Jiang Q, Ye F, Li SQ, Hong C, Chen LY, Li SY (2020) Effect of throat washings on detection of 2019 novel coronavirus. Clin Infect Dis 71(8):1980–1981. https://doi.org/10.1093/cid/ciaa416
- -Argenziano MG, Bruce SL, Slater CL, Tiao JR, Baldwin MR, Barr RG, Chang BP, Chau KH, Choi JJ, Gavin N, Goyal P, Mills AM, Patel AA, Romney MS, Safford MM, Schluger NW, Sengupta S, Sobieszczyk ME, Zucker JE, Asadourian PA, Bell FM, Boyd R, Cohen MF, Colquhoun MI, Colville LA, de Jonge JH, Dershowitz LB, Dey SA, Eiseman KA, Girvin ZP, Goni DT, Harb AA, Herzik N, Householder S, Karaaslan LE, Lee H, Lieberman E, Ling A, Lu R, Shou AY, Sisti AC, Snow ZE, Sperring CP, Xiong Y, Zhou HW, Natarajan K, Hripcsak G, Chen R (2020) Characterization and clinical course of 1000 patients with coronavirus disease 2019 in New York: retrospective case series. BMJ 369:m1996. https://doi.org/10.1136/bmj.m1996
- -Huang C, Wang Y, Li X, Ren L, Zhao J, Hu Y, Zhang L, Fan G, Xu J, Gu X, Cheng Z, Yu T, Xia J, Wei Y, Wu W, Xie X, Yin W, Li H, Liu M, Xiao Y, Gao H, Guo L, Xie J, Wang G, Jiang R, Gao Z, Jin Q, Wang J, Cao B (2020) Clinical features of patients infected with 2019 novel coronavirus in Wuhan, China. [published correction appears in Lancet. 2020 Jan 30:]. Lancet 395(10223):497–506. https://doi.org/10.1016/S0140-6736(20)30183-5. Epub 2020 Jan 24. Erratum in: Lancet. 2020 Jan 30
- Pasomsub E, Watcharananan SP, Boonyawat K, Janchompoo P, Wongtabtim G, Suksuwan W, Sungkanuparph S, Phuphuakrat A (2021) Saliva sample as a non-invasive specimen for the diagnosis of coronavirus disease 2019: a cross-sectional study. Clin Microbiol Infect 27(2):285.e1-285. e4. https://doi.org/10.1016/j.cmi.2020.05.001
- Vaz SN, Santana DS, Netto EM, Pedroso C, Wang WK, Santos FDA, Brites C (2020) Saliva is a reliable, non-invasive specimen for SARS-CoV-2 detection. Braz J Infect Dis 24(5):422–427. https://doi.org/10.1016/j.bjid.2020. 08.001. (Epub 2020 May 15)
- Lai CKC, Chen Z, Lui G, Ling L, Li T, Wong MCS, Ng RWY, Tso EYK, Ho T, Fung KSC, Ng ST, Wong BKC, Boon SS, Hui DSC, Chan PKS (2020) Prospective study comparing deep throat saliva with other respiratory tract specimens in the diagnosis of novel coronavirus disease 2019. J Infect Dis 222(10):1612–1619. https://doi.org/10.1093/infdis/jiaa487
- Guo XX, Deng NS, Chen QH, Yu HY, Ding XH, Hu SP, Nie HX (2019) Application of inflammatory markers in induced sputum in stable chronic obstructive pulmonary disease patients with positive bronchodilation tests. Curr Med Sci 39(4):560–567. https://doi.org/10.1007/ s11596-019-2074-7
- Chew MY, Ng J, Lim TK (2019) Diagnosing pulmonary tuberculosis by pooling induced sputum. J Clin Tuberc Other Mycobact Dis 15:100100. https://doi.org/10.1016/j.jctube.2019.100100
- Han H, Luo Q, Mo F, Long L, Zheng W (2020) SARS-CoV-2 RNA more readily detected in induced sputum than in throat swabs of convalescent COVID-19 patients Lancet Infect Dis 20(6):655–656. https://doi. org/10.1016/S1473-3099(20)30174-2. Erratum in: Lancet Infect Dis. 2020;20(5):e79
- Lai T, Xiang F, Zeng J, Huang Y, Jia L, Chen H, Wu J, Xie J, Liu S, Deng W, Zheng W, Huang Y, Zhang Q, Luo Q, Mo F, Long L, Zhang W, Chen W, Han H (2020) Reliability of induced sputum test is greater than that of throat swab test for detecting SARS-CoV-2 in patients with COVID-19: a multicenter cross-sectional study. Virulence 1:1394–1401. https://doi.org/10. 1080/21505594.2020.1831342
- World Health Organization (2020) Clinical management of COVID-19: interim guidance. World Health Organization. https://apps.who.int/iris/ handle/10665/332196.License:CCBY-NC-SA3.0IGO
- Chung M, Bernheim A, Mei X, Zhang N, Huang M, Zeng X, Cui J, Xu W, Yang Y, Fayad ZA, Jacobi A, Li K, Li S, Shan H (2020) CT imaging features of 2019 novel coronavirus (2019-nCoV). Radiology 295(1):202–207. https:// doi.org/10.1148/radiol.2020200230
- 23. CDC's Interim Guidelines for Collecting, Handling, and Testing Clinical Specimens from Persons for Coronavirus Disease 2019", 2021
- Zhang JF, Yan K, Ye HH, Lin J, Zheng JJ, Cai T (2020) SARS-CoV-2 turned positive in a discharged patient with COVID-19 arouses concern regarding the present standards for discharge. Int J Infect Dis 97:212–214. https://doi.org/10.1016/j.ijid.2020.03.007

- CDC. Return-to-work criteria for healthcare workers. Updated April 30, 2020. Accessed 3 May 2020. https://www.cdc.gov/coronavirus/2019ncov/hcp/return-to-work.html
- Lin C, Xiang J, Yan M, Li H, Huang S, Shen C (2020) Comparison of throat swabs and sputum specimens for viral nucleic acid detection in 52 cases of novel coronavirus (SARS-Cov-2)-infected pneumonia (COVID-19). Clin Chem Lab Med 58(7):1089–1094. https://doi.org/10.1515/ cclm-2020-0187
- Bosch BJ, Smits SL, Haagmans BL (2014) Membrane ectopeptidases targeted by human coronaviruses. Curr Opin Virol 6:55–60. https://doi. org/10.1016/j.coviro.2014.03.011
- Hamming J, Timens W, Bulthuis ML, Lely AT, Navis G, van Goor H (2004) Tissue distribution of ACE2 protein, the functional receptor for SARS coronavirus: a first step in understanding SARS pathogenesis. J Pathol 203(2):631–7. https://doi.org/10.1002/path.1570
- Liu R, Yi S, Zhang J, Lv Z, Zhu C, Zhang Y (2020) Viral load dynamics in sputum and nasopharyngeal swab in patients with COVID-19. J Dent Res 99(11):1239–1244. https://doi.org/10.1177/0022034520946251
- Zhang ZL, Hou YL, Li DT, Li FZ (2020) Laboratory findings of COVID-19: a systematic review and meta-analysis. Scand J Clin Lab Invest 80(6):441– 447. https://doi.org/10.1080/00365513.2020.1768587
- 31. Guan WJ, Ni ZY, Hu Y, Liang WH, Ou CQ, He JX, Liu L, Shan H, Lei CL, Hui DSC, Du B, Li LJ, Zeng G, Yuen KY, Chen RC, Tang CL, Wang T, Chen PY, Xiang J, Li SY, Wang JL, Liang ZJ, Peng YX, Wei L, Liu Y, Hu YH, Peng P, Wang JM, Liu JY, Chen Z, Li G, Zheng ZJ, Qiu SQ, Luo J, Ye CJ, Zhu SY, Zhong NS; China Medical Treatment Expert Group for Covid-19 (2020) Clinical characteristics of coronavirus disease 2019 in China. N Engl J Med 382(18):1708–1720. https://doi.org/10.1056/NEJMoa2002032
- 32. Zheng S, Fan J, Yu F, Feng B, Lou B, Zou Q, Xie G, Lin S, Wang R, Yang X, Chen W, Wang Q, Zhang D, Liu Y, Gong R, Ma Z, Lu S, Xiao Y, Gu Y, Zhang J, Yao H, Xu K, Lu X, Wei G, Zhou J, Fang Q, Cai H, Qiu Y, Sheng J, Chen Y, Liang T (2020) Viral load dynamics and disease severity in patients infected with SARS-CoV-2 in Zhejiang province, China, January–March 2020: retrospective cohort study. BMJ 369:m1443. https://doi.org/10. 1136/bmj.m1443
- 33. Wölfel R, Corman VM, Guggemos W, Seilmaier M, Zange S, Müller MA, Niemeyer D, Jones TC, Vollmar P, Rothe C, Hoelscher M, Bleicker T, Brünink S, Schneider J, Ehmann R, Zwirglmaier K, Drosten C, Wendtner C (2020) Virological assessment of hospitalized patients with COVID-2019. Nature 581(7809):465–469. https://doi.org/10.1038/s41586-020-2196-x. Epub 2020 Apr 1. Erratum in: Nature. 2020;588(7839): E35
- Huang Y, Chen S, Yang Z, Guan W, Liu D, Lin Z, Zhang Y, Xu Z, Liu X, Li Y (2020) SARS-CoV-2 viral load in clinical samples from critically ill patients. Am J Respir Crit Care Med 201(11):1435–1438. https://doi.org/10.1164/ rccm.202003-0572LE
- Lai T, Xiang F, Zeng J, Huang Y, Jia L, Chen H, Wu J, Xie J, Liu S, Deng W, Zheng W, Huang Y, Zhang Q, Luo Q, Mo F, Long L, Zhang W, Chen W, Han H (2020) Reliability of induced sputum test is greater than that of throat swab test for detecting SARS-CoV-2 in patients with COVID-19: a multicenter cross-sectional study. Virulence 11(1):1394–1401. https://doi.org/ 10.1080/21505594.2020.1831342
- Sri Santosh T, Parmar R, Anand H, Srikanth K, Saritha M (2020) A review of salivary diagnostics and its potential implication in detection of covid-19. Cureus 12(4):e7708. https://doi.org/10.7759/cureus.7708

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Submit your manuscript to a SpringerOpen[®] journal and benefit from:

- Convenient online submission
- ► Rigorous peer review
- Open access: articles freely available online
- High visibility within the field
- Retaining the copyright to your article

Submit your next manuscript at > springeropen.com