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Short communication

## Syndromic approach to SARS-CoV-2 detection using QIAstat-Dx SARS-CoV-2 panel from clinical samples

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## ABSTRACT

The QIAstat-Dx SARS-CoV-2 panel is a multiplex cartridge based assay based on real time PCR which can detect 17 respiratory viruses, including the novel coronavirus SARS-CoV-2. A syndromic approach is the need of the hour for COVID-19 diagnostics among patients presenting with respiratory symptoms. The present study was done to evaluate 120 archived respiratory clinical specimens for SARS-CoV-2 on the SARS-CoV-2 panel. Further, 27 specimens were tested for other respiratory viruses, in comparison with the BioFire RP1.7 platform. The sensitivity and specificity for SARS-CoV-2 on SARS panel was found to be 90.00 % and 100 % respectively, indicating good diagnostic accuracy. The positive predictive value was found to be 100 %, negative predictive value was found to be 99.93 % and accuracy was 99.93 %. Detection of other respiratory viruses observed a concordance of 77.7 %. Despite advantages of speed, minimal expertise and accurate results; significant costs and discrepancies at Ct >35 remain important limitations of the SARS panel.

### 1. Introduction

The introduction of multiplex, point-of-care tests (POC) for severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) diagnostics are game changing when deciding triaging, isolation and therapy of patients in populous countries like India. At the height of the coronavirus disease 2019 (COVID-19) pandemic, standard testing methods included real time PCR for detection of pan  $\beta$ -Coronavirus (target E gene) and SARS-CoV-2 (target S gene, N gene, RdRp gene, Orf1b gene). (Group ICS et al., 2020) These assays are time, labour intensive and requiring advanced laboratory infrastructure. Further, common respiratory viruses like influenza, rhinovirus and coronavirus 229E are widespread in the community and are often difficult to clinically differentiate from SARS-CoV-2. The recently introduced, QIAstat-Dx Respiratory SARS-CoV-2 panel (SARS-CoV-2 Panel, Qiagen, Hilden, Germany) is a cartridge based nucleic acid amplification test which can be operated in the field, requiring minimal technical expertise with a run time of 70 min. (Boers et al., 2020) It can detect SARS-CoV-2 along with 17 additional respiratory viruses including Influenza A, Influenza B, Parainfluenza, Respiratory syncytial viruses, Coronavirus,

Metapneumoviruses, Adenovirus and Rhino/Enterovirus similar to the QIAstat-Dx Respiratory Panel. The new SARS-CoV-2 Panel detects two genes of the SARS-CoV-2 virus genome (Orf1b/RdRp gene and E genes) with the same fluorescence channel and amplification of either or both target regions leads to a single fluorescence signal. (Leber et al., 2020; Visseaux et al., 2020) The aim of the present study was to evaluate the clinical performance of this assay against the standard Real Time PCR assay for SARS-CoV-2 (RealStar® SARS-CoV-2 RT-PCR Kit, Altona Diagnostics, Hamburg, Germany) and other respiratory viruses with Respiratory panel of multiplex assay, Biofire FilmArray RP1.7. (FilmArray, BioMérieux, France).

### 2. Methods

A retrospective study was conducted over 2 months on archived respiratory specimens (throat and nasopharyngeal swabs, NPS) from 1 November 2020 to 31 December 2020. The study was approved by Institutional Ethics committee and performed as per the Helsinki code. All clinical information were obtained from the hospital information system (HIS).

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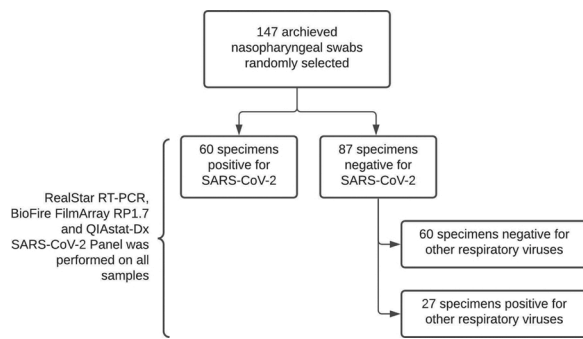


Fig. 1. Flow chart depicting sample selection.

**Table 1**  
Performance evaluation of QIAstat-Dx SARS-CoV-2 Panel when compared to RT-PCR.

	Value	95 % CI
Sensitivity	90.00 %	79.49%–96.24%
Specificity	100.00 %	94.04%–100.00%
Positive Likelihood Ratio	–	–
Negative Likelihood Ratio	0.10	0.05 to 0.21
Positive Predictive Value	100.00 %	–
Negative Predictive Value	99.93 %	99.85%–99.97%
Accuracy	99.93 %	–

\* Disease prevalence.0.70 %.

### 2.1. Specimen selection

The study was done on archived samples kept at  $-80^{\circ}\text{C}$  in the Department repository. The study involved a total of 147 samples that were randomly selected from the database. Group 1 included 120 samples which were selected as per the SARS-CoV-2 RT-PCR results (60 positive and 60 negative) and they all were negative for any other respiratory virus, group 2 included 27 samples that were all negative for SARS-CoV-2 but positive for other respiratory viruses. (Fig. 1) This division was done to evaluate the QIAstat-Dx assay across all the respiratory viral pathogens. All the retrieved samples were tested, in a single freeze thaw cycle, in parallel on QIAstat-Dx and re tested on both

RealStar RT-PCR as well as the Biofire FilmArray RP1.7 to reconfirm the earlier lab results.

#### 2.1.1. Real time PCR (RT-PCR) for SARS CoV-2

The standard test used in our study for the detection of SARS-CoV-2 was RealStar® SARS-CoV-2 RT-PCR. The RealStar RT-PCR targets E gene and S gene for the detection of SARS-CoV-2. RNA extraction was done by QIASymphony DSP Virus/pathogen Mini kit (Qiagen, Germany) and 10  $\mu\text{L}$  of RNA elute, from an original 500  $\mu\text{L}$  of VTM specimen, was required for the PCR. The assay requires 40 cycles of amplification and amplification for both the genes were considered as positive during the inclusion.

#### 2.1.2. QIAstat-dx assay

All 147 specimens were tested in parallel on the SARS-CoV-2 panel. Manufacturers protocol was followed for testing on the QIAstat-Dx Analyzer 1.0. Briefly, 300  $\mu\text{L}$  of clinical sample was added to the main port of the SARS-CoV-2 Panel cartridge and placed within the cartridge port. QIAstat-Dx Analyzer 1.0 Software processes controls, interprets the sample data including  $C_T$  values and provides a final report.

#### 2.1.3. BioFire film array respiratory panel RP 1.7

Detection of additional respiratory viruses was done on the automated multiplex system BioFire Film Array respiratory panel RP1.7. The BioFire is a closed system that performs sample preparation, reverse transcription and PCR in order to detect nucleic acid from multiple respiratory pathogens from a single nasopharyngeal swab.

All samples were processed and handled as per the recommended bio-safety guidelines. Any discrepant results were retested on both detection platforms (RealStar RT-PCR & QIAstat-Dx or BioFire FilmArray & QIAstat-Dx) for confirmation. Statistical analysis was performed using SPSS Ver 22 (IBM Corp., Armonk, NY, USA).

### 3. Results

We evaluated a total of 147 specimens on the SARS-CoV-2 Panel. The median age of the study group was 38 (IQR:15 – 68) years with Male: Female ratio was 3.1:1.

a) Evaluation of SARS-CoV-2 Panel while using RealStar RT-PCR as the reference method for detection of SARS-CoV-2.

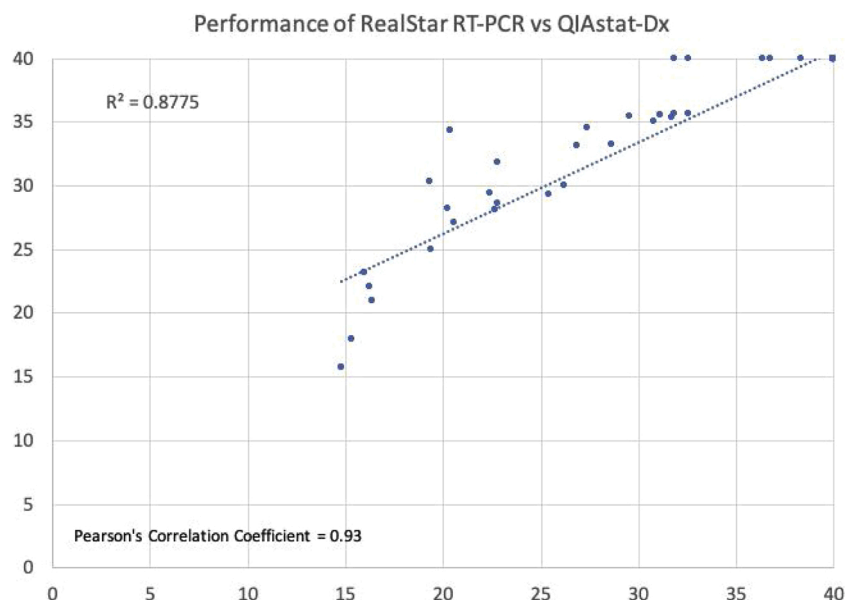


Fig. 2. Correlation between the cycle threshold ( $C_T$ ) values obtained by SARS-CoV-2 Panel on the QIAstat-Dx and by RealStar® RT-PCR for detection of SARS-CoV-2 with Pearson’s correlation coefficient was found to be 0.93. Negative samples have been illustrated as  $C_T$  value of 40.

**Table 2**  
Results description of BioFire RP 1.7 vs SARS-CoV-2 Panel (QIAstat-Dx) along with concordance.

Specimen No.	BioFire RP 1.7	SARS-CoV-2 Panel	Concordant
1	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
2	Coronavirus HKU 1	Coronavirus HKU 1	Yes
3	Influenza AH1 2009	Influenza AH1 2009	Yes
4	Influenza AH3	Influenza AH3	Yes
5	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
6	Influenza AH1 2009	Influenza AH1 2009	Yes
7	Influenza AH1 2009	Influenza AH1 2009	Yes
8	Influenza AH1 2009	Influenza AH1 2009	Yes
9	Coronavirus 229E	Coronavirus 229E	Yes
10	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
11	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
12	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
13	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
14	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
15	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
16	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
17	<b>Human Rhinovirus/ Enterovirus</b>	<b>TND</b>	<b>No</b>
18	<b>Human Rhinovirus/ Enterovirus</b>	<b>TND</b>	<b>No</b>
19	Coronavirus 229E	Coronavirus 229E	Yes
20	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
21	<b>Coronavirus 229E + Human Rhinovirus/ Enterovirus</b>	<b>Human Rhinovirus/ Enterovirus</b>	<b>No</b>
22	Influenza AH3	Influenza A + influenza A H3	Yes
23	Human Rhinovirus/ Enterovirus	Human Rhinovirus/ Enterovirus	Yes
24	<b>Human Rhinovirus/ Enterovirus</b>	<b>TND</b>	<b>No</b>
25	<b>Human Rhinovirus/ Enterovirus</b>	<b>TND</b>	<b>No</b>
26	<b>Coronavirus 229E + human metapneumovirus A/B</b>	<b>Human metapneumovirus A/ B</b>	<b>No</b>
27	Influenza AH3	Influenza A + influenza A H3	Yes

TND – Target Not Detected.

Out of 120 tested samples concordant results were seen in 114, 54 were positive and 66 were negative on the QIAstat-Dx. Taking prevalence of 0.7 % (as reported in India (Group ICS et al., 2020)) into consideration, the sensitivity of SARS-CoV-2 Panel was found to be 90.00 % (95 % CI:79.49%–96.24%) the specificity was found to be 100 % (95 % CI:94.04%–100.00%). (Table 1) Overall agreement of SARS-CoV-2 Panel with RT PCR was 95.0 %. There were 6 discordant samples which were positive on RealStar RT-PCR but negative on SARS-CoV-2 Panel. These discrepant cases were retested on RealStar RT-PCR and it was seen that there was amplification in both the genes but with higher C<sub>T</sub> values (>35).

The performance of the assay was further evaluated and correlation was assessed. Pearson's correlation coefficient was found to be 0.93 and R<sup>2</sup> coefficient of determination was found to be 0.88 (Fig. 2). No other respiratory viruses were detected on the SARS-CoV-2 Panel among these samples.

b) Evaluation of SARS-CoV-2 Panel while using BioFire RP1.7 for non-SARS-CoV-2 respiratory viruses.

The detection of 17 other respiratory viruses performed on 27 known positive clinical samples with various respiratory pathogens. There was 77.7 % concordance between viruses detected on SARS-CoV-2 Panel and BioFire RP1.7. Discrepant results were observed in 6 specimens (Table 2). Among these, 4 specimens were negative on SARS-CoV-2 Panel but detected Human Rhinovirus/Enterovirus on BioFire RP1.7. Further, 2 specimens were found to show infection with dual respiratory viruses on BioFire RP1.7 (Coronavirus 229E + Human Rhinovirus/Enterovirus and Coronavirus 229E + human metapneumovirus A/B). SARS-CoV-2 Panel could not detect Coronavirus 229E in both the specimens.

#### 4. Discussion

The arrival of POC tests for SARS-CoV-2 diagnostics are game changing when deciding triaging, isolation implementation and therapy of patients. (Boers et al., 2020) The SARS-CoV-2 Panel provides a novel sensitive and specific method to diagnose multiple viral infections in at risk patients in the midst of a pandemic. Delay in reporting due to centralised laboratory PCR testing is a significant challenge and SARS-CoV-2 Panel represents an important step to improve detection potential co-infections or superinfections. (Audi et al., 2020; Brendish et al., 2020) Our study found good performance characteristics, similar to other published studies on the QIAstat-Dx, where a positive percent agreement and negative percent agreement with RT-PCR as the reference standard was more than 90 %. (Visseaux et al. (2020); Lebourgeois et al. (2021)) Further, correlation between the C<sub>T</sub> values on the RealStar and QIAstat-Dx platforms were observed. We observed discrepant results on SARS-CoV-2 Panel in 6 cases with C<sub>T</sub> values >35 on the QIAstat-Dx. The higher C<sub>T</sub> values of these cases may indicate presence of degraded viral RNA and viral clearance or possible false positive on RT PCR. (Drew et al., 2020) Now even Indian council of medical research (ICMR) has stated in their guidelines that samples on RT-PCR should be given positive only with Ct values < 35. (Aranha et al., 2021) However, earlier all amplifications in both the genes even if > 35 Ct value were considered as positive while reporting. The SARS-CoV-2 Panel was also evaluated for detection of respiratory viruses other than SARS-CoV-2 using the BioFire RP1.7 as the reference standard. Concordance was found to be 77.7 % between both platforms. This could be due to different primer targets for various viruses on BioFire RP1.7 and SARS-CoV-2 Panel. Thus, a larger prospective study is required to assess the sensitivity of these platforms in the Indian setting. Sensitivity for detection of Human Rhinovirus/Enterovirus and coronavirus 229E on the SARS-CoV-2 Panel appears to be poor in comparison to the Biofire RP 1.7. However, present study could not accurately assess detection of non-SARS-CoV-2 respiratory viruses during the ongoing pandemic as due to the ongoing pandemic and lockdown situation, requests for assessing other respiratory virus were drastically reduced. Present study was further unable to evaluate detection of SARS-CoV-2 by the newer BioFire RP2.1 panel which includes targets for COVID-19 due to non-availability of this revised version of assay at our laboratory. The study is further limited by its small sample size and lack of direct NP/OP swab assessment in the SARS-CoV-2 Panel cartridge. Ease of specimen loading, rapid results and minimal need of technical expertise are in favour of use of the SARS-CoV-2 Panel. However, the decreased sensitivity among SARS-CoV-2 low viral load specimens remains an important limitation of this assay, although with limited clinical relevance.

In conclusion, performance and costs of implementation need to be further evaluated with a larger number of subjects before application in routine investigations in India. Furthermore, evaluation of this point-of-care test in critical care settings is desirable as SARS-CoV-2 becomes a regular differential among respiratory viral infections.

### Author Contribution

AG retrieved samples, performed QIAStat-Dx assay, data analysis and drafted the manuscript, AS performed QIAStat-Dx assay and edited manuscript, SR assisted in retrieving samples and performed QIAStat-Dx assay, DP edited the manuscript, RA designed the study and edited manuscript, EG conceived/designed the study and edited/approved final manuscript.

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None.

### Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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