



Detection of SARS-CoV-2 RNA with a Simple Concentration Method in Wastewater in Turkey: A Pilot Study in Çorum

Türkiye’de Atık Suda SARS-CoV-2 RNA’nın Basit bir Konsantrasyon Yöntemiyle Tespit Edilmesi: Çorum’da Pilot Çalışma

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ABSTRACT

Introduction: Many studies have shown the advantages of monitoring wastewater in the evaluation of microbiological pathogens circulating in the community. It was aimed to detect of SARS-CoV-2 RNA with a simple concentration method in wastewater in this study.

Materials and Methods: In our study, 7 wastewater samples were investigated, which were collected from the Urban Wastewater Treatment Plant (WWTP) of Çorum, between October to November 2020. Sorbent bags were left in water for 24 hours. Then they were used to trap and concentrate the virus. Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) RNA detected by using reverse transcription-quantitative polymerase chain reaction (RT-qPCR) assays.

Results: As a result, 3 of the 7 samples taken were positive for N and ORF1ab target gene regions.

Conclusion: This is the first study reporting the detection of SARS-CoV-2 RNA in wastewater with different concentration and capture method.

Key Words: COVID-19; SARS-CoV-2; Wastewater

ÖZ

Türkiye’de Atık Suda SARS-CoV-2 RNA’nın Basit bir Konsantrasyon Yöntemiyle Tespit Edilmesi: Çorum’da Pilot Çalışma

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Giriş: Toplumda dolaşan mikrobiyolojik patojenlerin izlenmesinde atık suyu izlemenin avantajları birçok çalışmayla gösterilmiştir. Bizde çalışmamızda basit bir konsantrasyon metoduyla atık suda SARS-CoV-2 RNA tespit etmeyi amaçladık.

Materyal ve Metod: Çalışmamızda Ekim-Kasım 2020 tarihleri arasında Çorum Kentsel Atık Su Arıtma Tesisi’nden toplanan 7 atık su numunesi incelendi. Sorbent torbalar 24 saat suda bırakıldı. Torbalar virüsü yakalamak ve konsantre etmek için kullanıldı. SARS CoV-2 RNA, ters transkripsiyon-kantitatif polimeraz zincir reaksiyonu (RT-qPCR) testleri kullanılarak saptandı.

Bulgular: Sonuç olarak, alınan 7 numuneden 3’ü N ve ORF1ab hedef gen bölgeleri için pozitif.

Sonuç: Çalışmamız, farklı konsantrasyon ve yakalama yöntemi ile atık sularda SARS-CoV-2 RNA tespitini bildiren ilk çalışmadır.

Anahtar Kelimeler: COVID-19; SARS-CoV-2; Atık su

INTRODUCTION

The novel coronavirus (SARS-CoV-2), which was first detected in December 2019 in Wuhan, China, caused a Public Health Emergency of International Concern about the new coronavirus disease (COVID-19) and global pandemic was officially declared by the World Health Organization (WHO)^[1]. Since that time, SARS-CoV-2 has spread to almost all countries and regions of the world, with 116.874.912 confirmed cases and 2.597.381 confirmed deaths, according to WHO^[2]. Turkey is among the countries affected by the pandemic.

Severe acute respiratory syndrome-related coronaviruses in the subgenus Sarbecovirus of the family Coronaviridae are a large family of viruses with single-stranded RNA, enveloped, and protein protrusions on the surface^[3]. Coronaviruses are responsible for mild to moderately severe respiratory diseases as well as severe diseases such as MERS (Middle East respiratory syndrome), SARS (Severe acute respiratory syndrome) and

COVID-19 (Novel coronavirus disease). However, SARS-CoV-2 is known to have a higher transmission rate and infectivity than those of SARS-CoV and MERS-CoV^[4].

The respiratory disease called COVID-19 typically causes fever, dry cough, dyspnea, or severe pneumonia^[5]. It is known that the main transmission route of the disease is inhalation from person-to-person aerosol transmission and direct contact, or fomite-to-hand contamination. Both live SARS-CoV-2 and viral RNA are found in a various specimens especially respiratory secretions, mucus, sputum and saliva. Currently, the most common and standard method used for COVID-19 diagnosis is the detection of SARS-CoV-2 in upper and lower respiratory specimens, including nasopharyngeal swabs, oropharyngeal swabs, sputum, aspirates, and bronchoalveolar lavage by real-time reverse transcription polymerase chain reaction (RT-PCR). The number and proportion of people infected with COVID-19 are mainly determined based on individual testing. On the

other hand, the virus can be detected in serum, urine and faeces^[6,7]. Patients can excrete the virus even if they are asymptomatic or pre-symptomatic and the pharyngeal samples obtained are negative, the stool may remain positive for SARS-CoV-2 for weeks by PCR^[8,9].

Sewage or wastewater, which are important sampling points, have been used for detection of pathogens in the community for years and its offer a practical approach to identify pathogens excreted in the faeces^[10]. Wastewater-based epidemiology (WBE) is not as specific as clinical surveillance, but it provides opportunities to estimate the prevalence of viruses^[11-13]. After the outbreak of SARS-CoV-2, presence of viruses RNA in wastewater was searched by researchers, and the virus was detected in many studies^[14-30]. Table 1 summarizes these researches. All of these studies have shown that wastewater monitoring can help understand the circulation of SARS-CoV-2 in human population.

According to our research, there is only one study indicating the presence of SARS-CoV-2 in wastewater in Turkey. Kocamemmi et al. have conducted the study in İstanbul, and RdRp gene has been used for virus detection^[28]. Our study is the first study using the modified sorbent bag method on detecting SARS-CoV-2 in wastewater.

MATERIALS and METHODS

Between October to December 2020, 7 samples were taken (Table 2) from the Wastewater Treatment Plant (WWTP) in Corum, Turkey (Figure 1,2). Samples were taken weekly and, the concentration were performed using bags with sorbent method to trap viruses recommended by the WHO Guidelines for environmental surveillance of poliovirus circulation with some modifications^[31]. A sorbent bag was fixed on the field using fishing line. Thus, the bag was left in the stream of water. After exposure for 24 hours, the sorbent-bag was placed in a sterile flask and transported to the laboratory in a cold box. The bag with sorbent was placed in a sterile Petri dish and washed out with sterile distilled water (about 5 ml) using a pipette and, eluate was collected and investigated. After RNA extraction with viral nucleic acid buffer (Bio-Eksen), RT-qP-

CR was performed using the SARS-CoV-2 Double Gene RT-qPCR kit (Bio-Speedy). Nucleocapsid (N) and ORF1ab gene region were used to detect the presence of SARS-CoV-2 by RT-qPCR assays from wastewater. As other studies, samples were considered positive for Ct below 40^[26,27,29].

RESULTS

We measured SARS-CoV-2 RNA by RT-qPCR using N and ORF1ab primer sets. Of all RT-qPCR quantification cycle (Cq) values, 42.8% (3) were below 40, and all of these samples had a Cq value less than 37. As a result, 3 out of 7 twenty-four hours raw wastewater samples were considered positive (Table 2).

DISCUSSION

Sewage systems are places where viruses are exposed to various temperatures and chemicals after being excreted in the faeces. Since virus concentration is also diluted with other waters and there are some inhibitors in these systems, it is very important even if viruses are detected in low titers in wastewater^[32]. Wastewater monitoring is known to have resulted in successful public health interventions in the past. The most successful example of this has been implemented within the polio virus eradication program, and the number of infected cases has decreased^[33].

The main focus of the studies carried out in sewage systems related to SARS-CoV-2 is the isolation and detection of RNA in wastewater. SARS-CoV-2 is known to have poor stability in wastewater and is more sensitive to disinfectants than non-enveloped viruses^[34]. However, there are many variables that can affect the detection and quantification of SARS-CoV-2 RNA. These are sampling method, sampling frequency, sampling types, sample storage conditions, pretreatment, virus concentration methods, RNA extraction methods, PCR format, target gene region and PCR performance characteristics^[35].

This study investigated the presence of SARS-CoV-2 RNA in untreated wastewater samples and also evaluated the effectiveness of an alternative concentration method. For the detection of SARS-CoV-2 in wastewater Polyethylene glycol (PEG) precipitation, ultracentrifugation and ultrafiltration methods were most frequently used for virus but

Table 1. Summary of SARS-CoV-2 studies in wastewater conducted in 2020

Reference	Country (in alphabetic order)	Dates of sampling	Type of sample	Concentration Method	RT-qPCR assay or target gene	Number of positive sample/ Total
[14]	Australia	March to April 2020	Untreated waste- water	Adsorbition- extraction and ultrafiltration	N_Sarbeco, NIID_2019- nCOV_N	2/9 0/9
[15]	Brazil	30 th October, 2019 until 4 th March, 2020	Raw sewage	PEG precipitation, centrifugation	N1, S, RdRp	4/6
[16]	Brazil	15 th April	Raw sewage	Ultracentrifugation	N2	5/12
[17]	Czech Republic	April to June 2020	Untreated wastewater	Direct flocculation	TGEV genome	13/112
[18]	Germany	April 8 th , 9 th	Untreated wastewater	Ultrafiltration and centrifugation	M gene RdRp	9/9
[19]	France	May 7 th , 18 th , 26 th Jun 4 th , 15 th , 25 th July 20 th 2020	Influent wastewater	Centrifugation and filter membran	N1, N3	Corelation Raw and corrected cases SARS-CoV-2 levels
[20]	France	March 5 th to April 23 th 2020	Raw wastewater	Centrifugation	E, RdRp	All processed samples
[21]	India	May 8 th and 27 th 2020	Untreated wastewater	Centrifugation, filtration, PEG pre- cipitation	ORF1ab, N, S	2/2
[22]	Italy	3 rd February-2 nd April 2020	Untreated wastewater	PEG dextran method	ORF1ab, Spike protein, RdRp	6/12 2/12
[23]	Italy	9 th October- 28 th February 2020	Untreated wastewater	PEG dextran method	ORF1ab, E gene, RdRp	18/40 26/40
[24]	Italy	14 th and 22 th April 2020	Untreated wastewater	Membrane filtration	ORF1ab, N, E gene	4/8
[25]	Japan	17 th March to 7 th May 2020	Influent waste water	Electronegative membrane-vortex	N_Sarbeco, NIID_2019- nCOV_N, CDC-N1 CDC-N2	0/5 1/5
[26]	Spain	February 12 th to April 14 th 2020	Untreated wastewater	Adsorbition- precipitation	CDC N1 CDC N2	12/15 13/15
[27]	The Netherland	5 th , 6 th , 7 th February and 4 th , 5 th , 15 th , 16 th March 2020	Untreated wastewater	Ultrafiltration	N1, N2, N3, E gene	18/29
[28]	Turkey	21 st and 25 th April 2020	Raw sewage from WWTP and manholes	Ultrafiltration, PEG adsorption	RdRp	5/7 2/2
[29]	USA	18 th March- 25 th of March 2020	Untreated wastewater	PEG precipitation	CDC N1 CDC N2 CDC N3	10/10

Table 1. Summary of SARS-CoV-2 studies in wastewater conducted in 2020 (continue)

Reference	Country (in alphabetic order)	Dates of sampling	Type of sample	Concentration Method	RT-qPCR assay or target gene	Number of positive sample/ Total
[30]	USA	January 13 th , February 3 rd , March 2 nd , April 8 th and 29 th 2020	Untreated wastewater	Centrifugation and ultrafiltration	CDC N1 CDC N2	2/15
*	This study	October 22 th and 27 th ; 3 th , 11 th , 17 th , 25 th November; 3 th December 2020	Untreated wastewater	Modified sorbent bags	N and ORF1ab	3/7

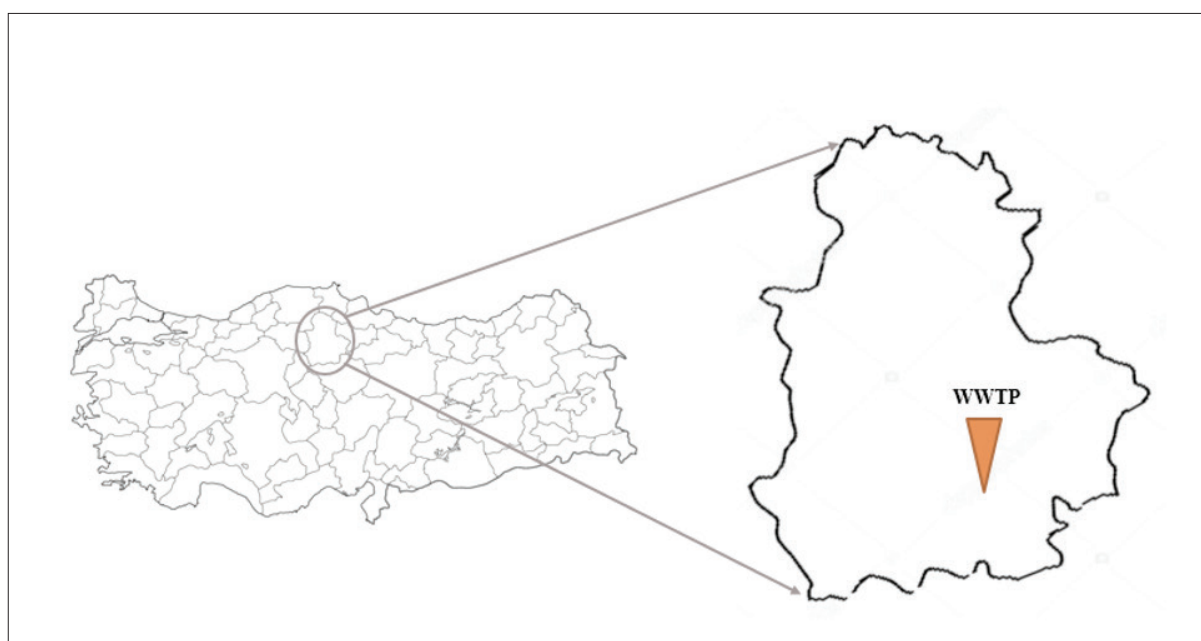


Figure 1. Çorum Province and WWTP on Turkey map.

WWTP: Wastewater treatment plant.

Table 2. Results of SARS-CoV-2 detection in the study period

Week	Sampling date	Types of sample	RT-PCRassays (Cq value)
1	22/10/2020	24 hours primary inlet wastewater	36.93
2	27/10/2020	24 hours primary inlet wastewater	36.77
3	3/11/2020	24 hours primary inlet wastewater	36.36
4	11/11/2020	24 hours primary inlet wastewater	-
5	17/11/2020	24 hours primary inlet wastewater	-
6	25/11/2020	24 hours primary inlet wastewater	-
7	03/12/2020	24 hours primary inlet wastewater	-



Figure 2. Wastewater treatment plant in Çorum.

studies show that there is lack of a standard method of concentration^[36,37]. In this study, the sorbent bag method recommended in the WHO's World Health Organization's Guidelines on Environmental Surveillance of the Poliovirus Circulation Protocol was modified and a new concentration method was adapted to capture and concentrate viruses in wastewater^[31]. Our study is the first and only study in the literature in which SARS-CoV-2 was detected using the modified sorbent bag method.

The first SARS-CoV-2 PCR tests were developed against the E and N gene, and these genes were used in numerous studies analyzing wastewater^[24-27,29,30]. Also, PCR targeting SARS-CoV-2 RdRP was used to a minor proportion^[18,20,22,28]. In this study, molecular detection of SARS-CoV-2 RNA was carried out using the test kit based on detecting the N and Orf1b genes. The other study in Turkey on the detection of SARS-CoV-2 effluent samples were target of RdRp gene^[28]. To our knowledge, our study is the first to detect SARS-CoV-2 RNA in environmental samples by targeting N and Orf1ab. In this study, 42.8% (3/7) of the wastewater samples showed positive results for SARS-CoV-2 RNA. Although the detection rates of the virus vary in studies, they all show that

the virus can be detected in wastewater.

Monitoring the spread of SARS-CoV-2 disease in the community is very important to take appropriate precautions. But since most people are asymptomatic, the virus is very difficult to track. In financial terms, it is not possible to do clinical tests on all individuals. Under these conditions, with a passive but effective sewage or wastewater monitoring method, the circulation of SARS-CoV-2 in the population can be monitored, especially when clinical testing is limited. In this way, the number of infected individuals can be estimated. Most of these findings are the results of research studies. Only the Netherlands uses environmental surveillance as part of a routine COVID-19 monitoring package^[38]. With a similar approach, sewage monitoring has been initiated in Germany, Australia and New Zealand^[38]. The study is important in terms of showing the detectability of the virus in wastewater in Turkey. Virus concentration method can serve as the basis for further research.

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ETHICS COMMITTEE APPROVAL

For this study, it is not necessary to obtain the ethical approval because the study is conducted with the samples of wastewater (untreated sewage water).

CONFLICT of INTEREST

None of the authors had conflict of interest.

AUTHORSHIP CONTRIBUTIONS

Concept and Design: SA, DK, ATÖ

Data Collection or Processing: SA, ÜS, ASG, GAA, MA, BG, ÇK, AA, NK

Analysis/Interpretation: SA, DK, GAA, ATÖ

Literature Search: SA, ATÖ, LX

Writing: SA, ATÖ, LX

Final Approval: All of authors

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