

Coupling wastewater-based epidemiological surveillance and modelling of SARS-CoV-2/ COVID-19: Practical applications at the Public Health Agency of Canada – Supplemental material

Meong Jin Joung, Chand S Mangat, Edgard M Mejia, Audra Nagasawa, Anil Nichani, Carol Perez-Iratxeta, Shelley W Peterson, David Champredon

List of supplemental figures and text

Figure S1: Inferred reportable incidence in five Canadian cities

Figure S2: Wastewater-based effective reproduction number in five Canadian cities

Laboratory analysis of SARS-CoV-2 concentration

Figure S1: Inferred reportable incidence in five

Canadian cities^a

Table S1: Conventional wastewater parameters

Laboratory processing of wastewater sample for SARS-CoV-2 ribonucleic acid detection

Figure S3: Laboratory processing of wastewater sample for identification of SARS-CoV-2

Vancouver Vancouver Bergen Descriptions Descriptions

Abbreviations: NML, National Microbiology Laboratory; PCR, polymerase chain reaction ^a For each city, the top panel shows the reported COVID-19 clinical cases by the municipality (black curve) and the inferred reportable incidence using wastewater epidemic model (WEM) (blue curve represents the mean estimate, and the blue shade its 95% CI) calibrated on the viral concentration in wastewater measured in that city (bottom panel, brown curve). Note the wastewater-based incidence is labelled "reportable" because we use a constant ratio of the reported cases over viral concentration. We do not attempt to estimate the "true" (and unobserved) incidence because we do not have any data to support the estimation of the ratio of true infections in the community over measured viral concentration. In all five cities, the wastewater signal and the clinical surveillance were in agreement until December 2021, when the Omicron variant emerged (top panels, black and blue curves are similar). After Omicron, we observe a divergence between the clinical and wastewater-based incidence: the clinical surveillance systematically reports fewer cases than what is implied by the wastewater signal (top panels, black curve below the blue one).

Figure S2: Wastewater-based effective reproduction number in five Canadian cities^a



Abbreviation: Eff. reprod. num., effective reproduction number ^a Each panel shows the mean wastewater-based effective reproduction number estimated using the wastewater epidemic model (WEM)

Laboratory analysis of SARS-CoV-2 concentration

Sampling frequency

As most wastewater surveillance activities are limited, daily sampling is often not possible. Typically, labs collect 2–3 samples per week which are often sent in-batch to a testing laboratory. To most accurately describe the trends of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) concentration in wastewater, these samples should be collected in well-spaced intervals throughout the week.

Sample handling, transportation, storage and quality control

By their nature, wastewater samples are very "active", i.e. there is a high degree of biological activity that will cause the nature of the sample to change fairly rapidly. After collection, samples should be cooled to 4°C as quickly as possible, then shipped cold using natural ice or ice packs via the most available rapid transportation. Upon arrival at the laboratory, they should be extracted as soon as possible, keeping the samples refrigerated until extraction. Standard Methods (1) recommends extraction within seven days of collection when analyzing for trace contaminants such as semi-volatile organics. Wastewater samples also contain suspended solids, which are an integral part of the matrix. As such, samples must be shaken frequently and thoroughly during any sub-sampling in the field or laboratory.

Composite and grab sampling both include the use of consumables (tubing, bottles) and reusable containers and equipment. These containers and equipment must be tested to ensure that the sampling system is not introducing contaminants into the samples. Laboratory-grade water can be used to create Equipment Blanks by simulating a composite or grab sampling event that includes sample tubing, pump tubing, collection containers and sub-sampling containers.

Wastewater treatment plan metadata and context

As discussed above, a wastewater study must be designed in the context of the collection and treatment system realities and details. Wastewater samples should always be characterized for conventional parameters to provide the context of wastewater strength and effectiveness of the treatment process. These parameters are listed in **Table S1**.

Table S1: Conventional wastewater parameters

Parameter	Comments	
Temperature—process	Indicator of microbial conditions for treatment	
Temperature—sample	Confirmation of target sampling temperature	
рН	Indicator for general chemistry and microbiology	
Alkalinity	Indicator of buffering capacity and nitrification	
Total suspended solids	Empirical gravimetric test, indicator of wastewater strength and treatment effectiveness, can be correlated with some chemical and microbiological constituents	
Chemical oxygen demand	Measure of material amenable to oxidation under strong chemical conditions, indicator of wastewater strength and treatment effectiveness	
Biochemical oxygen demand	Measure of material amenable to oxidation under specific biological conditions, indicator of wastewater strength and treatment effectiveness	
Total organic carbon	Measure of total organic (reduced) carbon, indicator of wastewater strength and treatment effectiveness	
Total Kjeldahl nitrogen	Measure of total organic (reduced) nitrogen	
Ammonia nitrogen	Measure of nitrogen available for nitrification	
Nitrate+nitrite	Measure of oxidized nitrogen, indicator of nitrification or denitrification	
Measured average daily flow	Available from the wastewater treatment plants, indicates the size of the system and confirms dry weather conditions or influence of storm events	

Summary

Any study of wastewater constituents requires a thorough understanding of the collection and treatment system in order to design a sample collection process that will answer the study questions. Sampling locations should be confirmed in consultation with wastewater treatment plants (WWTP) operators and described in detail in all reports and publications (e.g. collection of raw influent or after screening process). Likewise, sampling techniques (composite or grab) should be described in sufficient detail. Wastewater samples must be stored, transported, and handled appropriately to maintain their integrity.



Laboratory processing of wastewater sample for SARS-CoV-2 ribonucleic acid detection

Once samples are received at the lab, the temperature should be taken and recorded for future reference. Samples should be processed within 24 hours of receiving them in the lab. Effective wastewater surveillance that aims to detect the emergence of infection relies on rapid data collection and testing. For future use, store unused portions/aliquots of collected samples at -70° C. Since the strength of the viral ribonucleic acid (RNA) signal decreases after freezing, consequently more than one freeze-thaw cycle should be avoided. **Figure S3** describes the laboratory processing steps employed to detect a SARS-CoV-2 RNA signal from a wastewater sample.

Figure S3: Laboratory processing of wastewater sample for identification of SARS-CoV-2



Abbreviations: MHV-A59, Mouse Hepatitis Virus A59; RNA, ribonucleic acid; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2 $\,$

A wastewater test for SARS-CoV-2 is composed of three major steps (Figure S3):

- Viral concentration
- RNA extraction
- Molecular detection

Currently, the Public Health Agency of Canada (PHAC)/ National Microbiology Laboratory (NML) employs a variant of an ultrafiltration-based viral concentration method previously described by the Wigginton group (2) in addition to an assay directed at the "solids" or insoluble fraction of wastewater followed by molecular detection of SARS-CoV-2 by real-time polymerase chain reaction (RT-qPCR) directed at the N1 and N2 targets as developed by the United States (US) Centers for Disease Control and Prevention (CDC). Previous studies have shown that SARS-CoV-2 is roughly partitioned equally between the solid and liquid phase of wastewater. Volumetrically, the solids fraction is the major component of wastewater that holds the virus (3–6). The solids fraction can be easily collected and extracted to yield genetic material suitable for testing.

There is no consensus or gold standard test for SARS-CoV-2 detection in wastewater and inter-laboratory comparisons of methods have shown that most perform comparably. In collaboration with the Canadian Water Network, PHAC/ NML participated in an inter-laboratory study of SAR-CoV-2 wastewater detection methods (7). The sample was drawn from the largest of three Winnipeg WWTP and there were 85 clinical cases across the city at the time of collection. SARS-CoV-2 concentrations from most labs were within a 1-log band of each other. The Water Research Foundation (8) performed a similar inter-laboratory comparison amongst US laboratories. Grab samples from two WWTPs servicing Los Angeles County (~30K cases reported in the previous 14 days) were distributed to 36 laboratories for analysis. Despite methodological differences, range of reported results from most laboratories were within a 2-log band. Importantly, the above studies show that there was no consensus in method amongst participating laboratories, suggesting that most methods performed comparably.

The structure of the Canadian study revealed important considerations for laboratory methods. Eight laboratories received three sample types; samples spiked with inactivated SARS-CoV-2 at a high and low concentration (1,800 cp/mL vs. 20 cp/mL), and an unspiked sample. First, only laboratories that processed the insoluble or "solids" fraction of wastewater were able to derive signal from the unspiked sample. Because of the low clinical case count at the time of collection, this suggests that the insoluble fraction of wastewater is more effective for delivering early-warning indicators from wastewater surveillance. Studies of primary sludge and fractionation of wastewater influent have confirmed that the majority of the SARS-CoV-2 viral signal resides in the insoluble fraction. Secondly, the SARS-CoV-2 spiked into wastewater did not appreciably partition to the insoluble or "solids" fraction of wastewater. This suggests that the use of surrogate virus controls to monitor the overall efficiency of laboratory methods may not report on natural viral signal (see Controls, below).

Laboratory methods

Viral concentration

As SARS-CoV-2 is found at low levels within wastewater, concentration is required for accurate analysis, especially during the initial outbreak phases when the viral load is low (Figure S1). Concentration and extraction are widely considered as the most influential steps in directing the overall performance of the assay. Ahmed *et al.* have elaborated comparatively on the different methods (9). There are a variety of concentration methods, each with their own advantages and disadvantages as described in **Table S2**.

Table S2: Three common concentration methods

Method	Advantages	Disadvantages
Ultrafiltration Viral particles are concentrated by the use of a centrifugal filter device	Easy to use Short turnaround time Higher throughput than most methods Doesn't access solids component	Co-concentrations of inhibitory compounds Cost of labware (~\$30 per device) Supply of reagent may vary Requires centrifuge (to 4K x g) Filters can clog when sample turbidity is high
Electronegative filtration Viral particles are captured on a charged membrane by vacuum filtration	Low cost Low carryover of inhibitory compounds Low lab overhead to install test	Low-throughput High hands on time Requires extensive RNA extraction/ clean-up
PEG precipitation A precipitating agent is added to samples and viral particles are recovered by centrifugation	Low cost High supply of reagents	Low throughput, some methods require overnight incubation Requires centrifugation (to 10K x g) Requires extensive RNA extraction/ clean-up

Abbreviations: PEG, polyethylene glycol; RNA, ribonucleic acid

Some investigators employ direct extraction schemes on whole wastewater that skip the concentration step altogether (Figure S1). Typically, this involves processing about 1 mL of whole influent, which is both practical and amenable to highthroughput. Direct extraction avoids sample losses associated with concentration and could improve overall yield. However, PHAC/NML cautions against using direct extraction schemes because of their unknown performance during periods of low viral load, especially when testing low volumes of wastewater.

As discussed above, the majority of the viral signal is associated with the insoluble fraction of wastewater. As such, collection and processing of wastewater solids could improve recovery dramatically. PHAC/NML has found that the viral level in wastewater solids is equal to, or greater than, the liquids fraction.

RNA extraction

There are a variety of commercial RNA extraction kits available and each should be chosen based on the type on input material. Samples with high-solids content require mechanical disruption and extensive wash steps to remove inhibitory compounds; "soil" or "microbiome" extraction kits are well suited for this purpose. General RNA extraction kits can be employed when the input material is clarified by centrifugation. Commercial clean-up kits can improve detection in inhibited samples. RNA is unstable once extracted, therefore molecular detection should be performed the day of extraction.

Molecular detection

Detection of the viral signal from SARS-CoV-2 is by RT-qPCR as indicated in Figure S3. Specific primers amplify the SARS-CoV-2 genome and an intervening fluorescent probe is concomitantly consumed in this process. The viral signal is monitored by the increase in fluorescence associated with the consumption of this probe. Quantitation is achieved by measuring the number of cycles (cycle threshold or Ct) required for the fluorescence detection of consumed probe over a baseline value "threshold", which is compared to a standard curve of known input quantities. A consistent "threshold" value should be used for all samples and the "auto-thresholding" function of the RT-qPCR instrument should be disabled. The threshold is specific to the RT-qPCR instrument and primers/probes chosen for analysis require optimization to reduce noise between replicates.

Sequences for SARS-CoV-2 primers, probes, controls, and sequences required for quantification above are provided in (3–6). There are several established primer/probe sets used to detect SARS-CoV-2 in wastewater and there is no current consensus as to which molecular targets are best. Indeed, investigators have reported contradictory performances of the same primer/probe combinations. PHAC/NML has evaluated the E_Sarbeco (10) and US-CDC N1/N2 targets and found N1/N2 to be the most sensitive and consistent. NML recommends using two targets to mitigate the risk of mutation.

Controls

The complex and variable nature of wastewater requires three controls run alongside the molecular detection of SARS-CoV-2 to account for variations in the composition of wastewater and evaluate overall efficiency of the process.



Process control

To account for varying efficiencies of the RNA extraction from wastewater a spike-in control of whole-viral material is added to wastewater prior to concentration as described in the main article Figure 3. A parallel concentration/extraction is run in the phosphate buffered saline (PBS) and the overall concentration and extraction efficiency is inferred by comparing the relative recovery. The process control or surrogate is ideally a coronavirus of the same genus as SARS-CoV-2 and thus physically structured similarly to SARS-CoV-2 to best report on its recovery. Common process controls are murine hepatitis virus (Mouse Hepatitis Virus A59, MHV-A59), bovine coronavirus or one of the seasonal human coronaviruses. PHAC/NML currently adds MHV-A59 as a process control in its assays. PHAC/NML has found that cultured MHV-A59 does not appreciably partition to the solids phase of wastewater and thus is not reflective of the natural state of the virus (as discussed above). A similar lack of solids phase partitioning has been observed from cell-culture-produced SARS-CoV-2. Therefore, the utility of surrogates is likely more suited to methods that process only the liquid fraction of wastewater.

Fecal control

The fecal load of wastewater can vary across wastewater collection systems. Surface water, ground water, and varying industrial and institutional inputs can dilute wastewater and introduce variance to the SARS-CoV-2 signal. A test specific for fecal load is applied to account for the varying composition of wastewater. PHAC/NML currently directs a RT-qPCR reaction against the Pepper Mild Mottle Virus (PMMoV), a naturally occurring virus that is found abundantly in edible peppers and reports on fecal load (11). Other fecal indicators of note are the HF183 and crAssphage (12,13).

Inhibition control

Wastewater contains contaminants that are known to inhibit PCR assays. To detect the presence of inhibitors, purified RNA from a source that is not found in wastewater is added to wastewater RNA extracts or alternatively to the wastewater concentrates. The signal intensity of this reaction is compared to the inhibition indicator material tested alone. The Water Research Foundation inter-laboratory study suggests that a shift of \geq 1 Ct suggests absence of PCR inhibition (8). An alternative approach when purified RNA is not available is to dilute a wastewater RT-qPCR reaction and compare the resultant Ct value with the expected value (14). When inhibition is outside of one Ct of the expected range, it is suggested to dilute the wastewater prior to extraction and/or to flag the results prior to reporting.

Negative controls

Good practice with RT-qPCR based experiments is to run a RT-qPCR reaction without the addition of template. Any signal observed in this control would indicate the presence of contaminants in the RT-qPCR reagents. Mock concentration/ extractions using buffer or water alone should be run periodically to identify contaminated labware or reagents.

Normalizing techniques

Following quantification, the SARS-CoV-2 signal and associated controls are expressed in copies per volume of the processed wastewater (e.g. cp/mL). Adjustments should be made for wastewater losses over concentration (e.g. unrecoverable dead volume in centrifugal filter devices) and/or dilution of samples prior to extraction (e.g. to mitigate inhibition if observed). Estimation of the dead-volume can be made by determining the weight difference of the centrifugal filter device before the application of the sample and after sample recovery, assuming a density of 1 g/mL.

Quantified viral targets are normalized to the quantified fecal indicator and this value alongside the un-normalized data should be considered minimal for reporting in Equation 1.

Equation 1:

$$\frac{Viral target \left(\frac{copies}{mL}\right)}{Fecal Indicator \left(\frac{copies}{mL}\right)} = Fecal normalized (copies/mL)$$

Further adjustments to the reported value can be made by incorporating the yield of the process control to overall yield. First, overall yield is calculated in Equation 2.

Equation 2:

 $\frac{Process \ control \ copies \ recovered}{Process \ control \ copies \ in} = Process \ control \ yield$

To adjust for yield process recovery, apply the following formula to calculate the yield adjusted viral load in Equation 3.

Equation 3:

$$\frac{1}{Process \ control \ yield} \ * \ Fecal \ normalized \ \left(\frac{copies}{mL}\right) = Yield \ adjusted \ viral \ load$$

As described above, the process control may not accurately report on the overall SARS-CoV-2 yield, especially where the solids fraction is primary target for extraction. Where possible, flow-based normalization is the preferred method to account for dilution effects, which can be a major contributor to the loss of signal—particularly during wet weather events.

References

- Baird R, Bridgewater L. Standard methods for the examination of water and wastewater. 23rd ed. Washington, D.C.: American Public Health Association; 2017.
- Ye Y, Ellenberg RM, Graham KE, Wigginton KR. Survivability, partitioning, and recovery of enveloped viruses in untreated municipal wastewater. Environ Sci Technol 2016;50(10): 5077–85. DOI PubMed
- Kitamura K, Sadamasu K, Muramatsu M, Yoshida H. Efficient detection of SARS-CoV-2 RNA in the solid fraction of wastewater. Sci Total Environ 2021;763:144587. DOI PubMed
- Westhaus S, Weber FA, Schiwy S, Linnemann V, Brinkmann M, Widera M, Greve C, Janke A, Hollert H, Wintgens T, Ciesek S. Detection of SARS-CoV-2 in raw and treated wastewater in Germany - Suitability for COVID-19 surveillance and potential transmission risks. Sci Total Environ 2021;751:141750. DOI PubMed
- Hokajärvi AM, Rytkönen A, Tiwari A, Kauppinen A, Oikarinen S, Lehto KM, Kankaanpää A, Gunnar T, Al-Hello H, Blomqvist S, Miettinen IT, Savolainen-Kopra C, Pitkänen T. The detection and stability of the SARS-CoV-2 RNA biomarkers in wastewater influent in Helsinki, Finland. Sci Total Environ 2021;770:145274. DOI PubMed
- Graham KE, Loeb SK, Wolfe MK, Catoe D, Sinnott-Armstrong N, Kim S, Yamahara KM, Sassoubre LM, Mendoza Grijalva LM, Roldan-Hernandez L, Langenfeld K, Wigginton KR, Boehm AB. SARS-CoV-2 RNA in Wastewater Settled Solids Is Associated with COVID-19 Cases in a Large Urban Sewershed. Environ Sci Technol 2021;55(1):488–98. DOI PubMed
- Chik AH, Glier MB, Servos M, Mangat CS, Pang XL, Qiu Y, D'Aoust PM, Burnet JB, Delatolla R, Dorner S, Geng Q, Giesy JP Jr, McKay RM, Mulvey MR, Prystajecky N, Srikanthan N, Xie Y, Conant B, Hrudey SE; Canadian SARS-CoV-2 Inter-Laboratory Consortium. Comparison of approaches to quantify SARS-CoV-2 in wastewater using RT-qPCR: results and implications from a collaborative inter-laboratory study in Canada. J Environ Sci (China) 2021;107:218–29. DOI PubMed

- Pecson BM, Darby E, Haas CN, Amha YM, Bartolo M, Danielson R, Dearborn Y, Di Giovanni G, Ferguson C, Fevig S, Gaddis E, Gray D, Lukasik G, Mull B, Olivas L, Olivieri A, Qu Y; SARS-CoV-2 Interlaboratory Consortium. Reproducibility and sensitivity of 36 methods to quantify the SARS-CoV-2 genetic signal in raw wastewater: findings from an interlaboratory methods evaluation in the U.S. Environ Sci (Camb) 2021;7:504–20. DOI PubMed
- Ahmed W, Bertsch PM, Bivins A, Bibby K, Farkas K, Gathercole A, Haramoto E, Gyawali P, Korajkic A, McMinn BR, Mueller JF, Simpson SL, Smith WJ, Symonds EM, Thomas KV, Verhagen R, Kitajima M. Comparison of virus concentration methods for the RT-qPCR-based recovery of murine hepatitis virus, a surrogate for SARS-CoV-2 from untreated wastewater. Sci Total Environ 2020;739:139960. DOI PubMed
- Corman VM, Landt O, Kaiser M, Molenkamp R, Meijer A, Chu DK, Bleicker T, Brünink S, Schneider J, Schmidt ML, Mulders DG, Haagmans BL, van der Veer B, van den Brink S, Wijsman L, Goderski G, Romette JL, Ellis J, Zambon M, Peiris M, Goossens H, Reusken C, Koopmans MP, Drosten C. Detection of 2019 novel coronavirus (2019-nCoV) by realtime RT-PCR. Euro Surveill 2020;25(3):2000045. DOI PubMed
- Haramoto E, Kitajima M, Kishida N, Konno Y, Katayama H, Asami M, Akiba M. Occurrence of pepper mild mottle virus in drinking water sources in Japan. Appl Environ Microbiol 2013;79(23):7413–8. DOI PubMed
- Stachler E, Kelty C, Sivaganesan M, Li X, Bibby K, Shanks OC. Quantitative CrAssphage PCR Assays for Human Fecal Pollution Measurement. Environ Sci Technol 2017;51(16):9146–54. DOI PubMed
- Green HC, Haugland RA, Varma M, Millen HT, Borchardt MA, Field KG, Walters WA, Knight R, Sivaganesan M, Kelty CA, Shanks OC. Improved HF183 quantitative real-time PCR assay for characterization of human fecal pollution in ambient surface water samples. Appl Environ Microbiol 2014;80(10):3086–94. DOI PubMed
- King C, Debruyne R, Kuch M, Schwarz C, Poinar H. A quantitative approach to detect and overcome PCR inhibition in ancient DNA extracts. Biotechniques 2009;47(5):941–9. DOI PubMed