

1 Wastewater Surveillance for Monkeypox Virus in Nine California Communities

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21 Abstract

22 Background: Wastewater represents a composite biological sample from the entire contributing
23 population. People infected with monkeypox virus (MPXV)¹ may excrete viral DNA into wastewater
24 via multiple ways such as in feces, urine, skin lesions, and/or saliva. We describe results from rapid
25 establishment of wastewater surveillance in selected regions in California within a month of the first
26 reported case of monkeypox in the United States. Methods: PCR assays targeting genomic DNA from
27 MPXV were deployed in an ongoing wastewater surveillance program in California. MPXV DNA
28 concentrations were measured daily in settled solids samples from nine wastewater plants. Results
29 over a four-week period were validated across different MPXV assays, compared using influent and
30 solids samples, and correlated using non-parametric methods (Kendall's tau) with the number of
31 monkeypox cases reported from each sewershed. Results: MPXV DNA was detected at all nine sites
32 between June 19 and August 1, 2022; 5 of 9 sites detected MPXV DNA prior to or within a day of the
33 first case identified in the source sewershed. At the four sites with >10 positive detections, we
34 observed a positive, statistically significant correlation ($p < 0.001$) between MPXV DNA in wastewater
35 solids and incidence rate of reported cases. Conclusions: Our findings suggest wastewater can be
36 used to effectively detect the introduction of MPXV and monitor its circulation in the community to
37 inform public health and clinical response. This flexible wastewater surveillance infrastructure may be
38 rapidly leveraged to monitor other pathogens of public health importance that are shed into
39 wastewater.

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42 Introduction

43 Monkeypox virus (MPXV), of the *Orthopoxvirus* genus in the family *Poxviridae*, is endemic in Western
44 and Central Africa where infection has been linked to transmission from infected animals or humans.
45 Sporadic cases and outbreaks linked to travel or imported animals have been recognized in non-
46 endemic countries since the first identification of human disease in 1970¹. In May 2022, cases of
47 monkeypox infection without association to endemic areas were reported in multiple European
48 countries¹. Within weeks, cases were identified in the United States and a dozen countries around the
49 world. By late July, the World Health Organization declared monkeypox a public health emergency of
50 international concern². Unlike previous outbreaks, this global outbreak is driven by human-to-human
51 transmission and cases without association to each other have been identified, suggesting additional
52 undetected community transmission^{3,4}.

53
54 Since recognition of the global outbreak, there has been a rapid scale-up of public health response,
55 including substantial increases in testing and efforts to educate clinicians and the public to mitigate
56 spread⁵. However, surveillance is dependent on practical access to and utilization of testing, limited
57 by awareness of a disease that is novel to the general public and most clinicians, stigma of a disease
58 that has to date primarily been diagnosed in gay, bisexual, and other men who have sex with men,
59 and the potential for asymptomatic cases^{6,7}. Alternative public health surveillance approaches
60 independent of individual testing, such as wastewater surveillance, provide an attractive means to
61 detect and track emerging monkeypox transmission and provide situational awareness for public
62 health agencies and clinicians.

63
64 Use of wastewater surveillance to monitor trends in infectious diseases has grown rapidly.
65 Wastewater represents a composite biological sample combining inputs from community members
66 connected to a sewer network and many pathogens are shed in ways that reliably end up in
67 wastewater, including in urine, feces, oral and nasal secretions, and sloughing of skin. Wastewater
68 surveillance has been reliably used by public health agencies throughout the coronavirus disease
69 2019 (COVID-19) pandemic to monitor for SARS-CoV-2, the causative virus of COVID-19.
70 Concentrations of SARS-CoV-2 viral RNA are strongly correlated with COVID-19 case incidence⁸⁻¹⁰,
71 and recent studies show this is also the case for other respiratory viruses such as Influenza A and
72 respiratory syncytial virus^{11,12}.

73
74 Recent small observational studies of monkeypox infection have confirmed the presence of viral DNA
75 for some individuals in saliva, feces, urine, semen, and/or skin lesions^{13,14}. Although concentrations of
76 viral DNA in specimens were not reported, Cq values (a measurement of how many cycles are needed
77 to detect a signal) from quantitative polymerase chain reaction (PCR) measurements were low,
78 suggesting high concentrations of MPXV DNA even 16 days after symptom onset¹³. These data are
79 consistent with reports of MPXV shedding from previous outbreaks in humans¹⁵ and experimental
80 studies in animals¹⁶⁻¹⁹. Related viruses including smallpox have been shown to be excreted in urine²⁰.
81 Together, this evidence suggests that MPXV DNA is likely to appear in wastewater. Based on a
82 systematic review of the literature²¹, no study to date has documented the persistence of
83 orthopoxviruses in wastewater. However, Vaccinia virus has been reported to persist for days in raw
84 freshwater and marine waters²¹, suggesting orthopoxviruses may persist as wastewater transits from
85 homes to nearby wastewater treatment plants.

86

87 In a collaborative effort between researchers running a wastewater surveillance program and the
88 California Department of Public Health (CDPH), we rapidly adapted and deployed an assay for
89 surveillance of MPXV DNA in multiple sewersheds in California. We describe results from this work,
90 including the establishment of monitoring within a month of the first reported case of monkeypox in
91 the US and an exploration of the relationship between MPXV DNA concentrations in wastewater and
92 cases in the community.

93

94 Methods

95

96 *MPXV Molecular Assay.* To detect MPXV DNA in wastewater, we used assays developed by the United
97 States Centers for Disease Control and Prevention (CDC)²². The G2R_G assay, which targets a region
98 of the OPG002 gene common to all MPXV sequences, was used on all samples. A subset of samples
99 (described further below) was also assayed for MPXV DNA using the G2R_WA assay which targets a
100 separate region of the OPG002 gene that is specific to Clade II. The choice of assays was confirmed by
101 alignment with sequences from the 2022 outbreak.

102

103 *Wastewater samples for MPXV surveillance.* Each day between June 19, 2022 and August 1, 2022,
104 settled solids were collected from nine publicly owned treatment works (POTWs) in California in the
105 Greater San Francisco Bay and Sacramento areas (Table 1, S1). The POTWs treat wastewater from
106 between 66,622 and 1,480,00 people; details of the POTWs and specific sample collection processes
107 are provided in Wolfe et al.⁸ Samples were couriered to a laboratory the same day they were
108 collected and processed immediately, unless otherwise specified (Table S2), with results available
109 within 24 hours of sample collection. A total of 407 solids samples were included in this study.

110

111 At the laboratory, nucleic acids were extracted and purified from the solids using previously
112 published methods^{8,23-25} (see SI for additional details). Nucleic acids were used undiluted as template
113 in digital droplet PCR wells; 10 replicate wells were run per sample and results from the wells were
114 merged to determine the concentration of the G2R_G target. PCR cycling conditions and further
115 details of digital droplet PCR (ddPCR) are provided in the SI. We also measured pepper mild mottle
116 virus (PMMoV) RNA gene concentrations; as well as recovery of spiked-in bovine coronavirus (BCoV)
117 RNA following methods outlined elsewhere²³⁻²⁵. Additionally, negative and positive extraction
118 controls, and negative and positive PCR controls were included in each plate. Gene fragments were
119 used as controls for the G2R_G assay; other controls are described elsewhere⁸. Nucleic acids were
120 not stored prior to analysis, with the exception of some from a POTW serving the part of San
121 Francisco (SEP) which were stored at -80°C less than two weeks and subjected to a freeze thaw prior
122 to G2R_G analysis (Table S2).

123

124 A subset of samples with the highest concentrations of G2R_G detection from the two POTWs with
125 the most frequent G2R_G (OSP and SEP, Table S2) detections were processed for the second MPXV
126 genomic target, G2R_WA, using the same replication and QA/QC as described for G2R_G. A gene
127 fragment was used as the G2R_WA positive control. PCR cycling conditions are provided in the SI.

128

129 *Liquid influent samples.* Samples of 24-h composited influent were collected in sterile containers on
130 ~7 consecutive days (Table S2) from two POTWs (OSP and SEP) that had the highest rates of detection
131 of the G2R_G target to compare the concentrations obtained using liquid wastewater to those

132 obtained using settled solids. Methodological details are in the SI. In brief, viral targets were
133 concentrated from wastewater influent using Nanotrap particles (Ceres Nanosciences, Manassas, VA)
134 before nucleic acids were extracted, purified, and used as template in digital droplet PCR to measure
135 concentrations of G2R_G and G2R_WA targets, as well as PMMoV and BCoV recovery following the
136 same protocol as described for solids.

137
138 *Case Data.* Counts of incident cases of monkeypox, defined in this study as people with lesion swabs
139 PCR positive for orthopoxvirus or MPXVs, were recorded as a function of episode date (symptom
140 onset date, or if not available, the earliest of laboratory result or case record creation date) and
141 report date (laboratory result date or, if not available, case record create date). Cases were
142 aggregated within sewersheds based on georeferenced home addresses, delineated using POTW-
143 specific GIS shape files. To obtain daily incidence rates of monkeypox, a 7-day rolling average of
144 incident cases per 100,000 population was calculated using the estimated population served in each
145 sewershed (Table S1). To explore incidence of cases of monkeypox at the time MPXV DNA was first
146 detected in wastewater at each sewershed, we compared incidence during periods with no MPXV
147 detection (any 7-day period without detection in wastewater) with the period of first detection (first
148 7-day period with at least 2 detections in wastewater). Weekly incidence of cases of monkeypox was
149 estimated for each of these periods (all cases with an episode date within 7-day periods of no
150 detection of MPXV DNA, or all cases within the 7-day period leading up to the date of first detection
151 of MPXV DNA, divided by total sewershed population). Since the goal was to describe upper limits of
152 incidence of monkeypox cases in each sewershed when MPXV DNA was not detected in wastewater,
153 for sewersheds with multiple 7-day periods without detection of MPXV DNA in wastewater, the single
154 highest weekly incidence of cases of monkeypox was used.

155
156 *Statistical analysis.* We assessed the association between the moving 5-day trimmed (highest and
157 lowest of the 5 values excluded) average of MPXV DNA concentrations in wastewater solids and 7-
158 day moving averages of monkeypox daily incidence using Kendall's tau for the subset of sites for
159 which both variables had more than 10 values (out of ~44) greater than 0 to ensure both variables
160 had sufficient variance. Tests were repeated using the MPXV DNA concentrations in wastewater
161 normalized by PMMoV, and for both the episode date and report date associated with cases. Non-
162 parametric methods were chosen as data were not normally distributed (Shapiro-Wilk test). We also
163 assessed the relationship between MPXV DNA concentrations in measurements taken from liquid and
164 solids samples and the relationship between results from the G2R_G and G2R_WA assays on the
165 same sample using Kendall's tau. A paired Wilcoxon Sign Rank test was then used to assess the
166 difference in results between liquids and solids, and between the G2R_G and G2R_WA assays used on
167 the same samples. All statistical analysis was performed in RStudio (version 2021.09.2).

168 ab

169 This activity was reviewed by CDC and was conducted consistent with applicable federal law and CDC
170 policy.§

171

172 Results

173 MPXV DNA was detected in wastewater samples across all sites (9/9) monitored during the study
174 period from June 19 to August 1, 2022 (Table 1, Fig 1, Fig 2, Fig S4). Concentrations of MPXV DNA
175 (G2R_G target) ranged from non-detect to 24,113 copies/g dry weight of wastewater solids (more
176 about lowest detectable concentration in SI). Positive and negative controls were all positive and

177 negative respectively, BCoV recoveries were higher than 10%, and PMMoV concentrations were
178 within an expected range for each POTW. This indicates assays performed efficiently and
179 demonstrated no evidence of contamination. A subset of samples was run with a second assay
180 (G2R_WA, specific to Clade II of MPXV) and no significant difference between measurements of
181 G2R_WA and G2R_G were observed (details in SI). A comparison of results from liquid and solids
182 samples showed a significant association (Kendall's tau = 0.52, n = 28, p = 0.00012), with significantly
183 higher concentration in solids (about 10^3 higher) on a per mass basis (Wilcoxon signed rank test, n =
184 28, p<0.001) (Fig 3). Further details are in the SI.

185
186 The first clinical monkeypox case in any of the study areas was reported in the Sacramento
187 sewershed on May 23, 2022; wastewater monitoring for MPXV began on June 19 (Table 1). The first
188 wastewater samples to test positive were June 20 samples from each of the two facilities serving San
189 Francisco (OSP, SEP). Daily samples from these sites were sporadically positive for the following 1-2
190 weeks, after which samples were consistently positive in increased concentrations (Fig 1, Fig 2, Fig
191 S4). The next sites to test positive were SJ (June 25), and SAC (July 3). These sites also had a pattern of
192 sporadic positives followed by increasing rates of positive samples and concentrations. Samples from
193 SUN, PA, Gil, Dav, and SVCW also tested positive; SVCW and PA had some increase in rates of positive
194 samples.

195
196 During the study period, the total number of cases in each sewershed ranged from 0 (Dav) to 494
197 (SEP). In SJ, Gil, and Dav, a first positive detection in wastewater preceded the report of a first case of
198 monkeypox. During periods with no detection in wastewater, the highest weekly incidence recorded
199 at each sewershed was an average of 0.68 cases (median 0.65, range 0-1.28) per 100,000. The
200 average weekly incidence when a sewershed first had at least 2 detections in 7 days was 1.14
201 (median 0.95, range 0.89-2.01) per 100,000.

202
203 At the four sites with >10 positive samples and days with cases, there was a significant, positive
204 association between the 5-day trimmed average concentration of viral DNA in wastewater and the 7-
205 day average monkeypox incidence rate in the corresponding sewershed both when cases were
206 compared using episode date (Kendall's tau = 0.59, p <0.001, n = 176) and report date (Kendall's tau =
207 0.66, p <0.001, n = 130). Results were similar when wastewater data were normalized by PMMoV and
208 with raw, daily concentrations from wastewater (Table S4).

209 210 Discussion

211 Surveillance for infectious diseases is a core function to inform public health, clinical, and general
212 public understanding of risk and strategies for prevention of disease and is especially important for
213 emerging infectious diseases. In the current global monkeypox outbreak, localized detection of
214 disease introduction and circulation informs public health and clinical response, allows appropriate
215 allocation of scarce testing, therapeutic and vaccine resources, and is important for risk messaging to
216 the public. Traditional case surveillance for monkeypox is dependent on confirmatory PCR diagnostic
217 testing of cases. While such testing is important for clinical decision making and is the backbone of
218 disease surveillance, limitations include variable disease recognition, variable care-seeking due to
219 disease severity and stigma, testing availability, and clinician awareness. Complementary strategies
220 that can overcome these limitations and provide rapid population level awareness, like wastewater
221 surveillance, can be vital for public health response.

222

223 We found that an established wastewater surveillance infrastructure could be rapidly leveraged to
224 detect and monitor MPXV DNA to inform public health response. MPXV DNA was consistently
225 detectable in wastewater from sewersheds with confirmed cases, even in locations with few
226 identified cases. In some places, detections in wastewater preceded identification of the first cases in
227 the community. We also found that the level of viral DNA in wastewater correlated with monkeypox
228 incidence rate, suggesting that wastewater surveillance is a viable methodology to monitor trends in
229 monkeypox disease activity. These findings suggest that wastewater surveillance can be adopted
230 where feasible as an adjunct public health tool in the current global outbreak to monitor monkeypox
231 disease activity, including in areas without known cases.

232

233 It is not yet possible to translate the concentration of MPXV DNA in wastewater to a predicted
234 number of cases in the sewershed. One important reason is the unclear understanding of the true
235 case incidence in each sewershed. However, comparisons between the weekly case incidence at the
236 time of first consistent detections in wastewater (at least two detections within a 7-day period), and
237 the weekly incidence during 7-day periods of no detection suggest that the beginning of consistent
238 detections in wastewater correspond to a weekly incidence of at least 0.89-2.01 cases per 100,000.
239 This is a likely underestimate, with true cases under-reported due to the reasons discussed above.
240 One sewershed (Dav) had one positive wastewater sample but no reported cases; this likely reflects
241 unidentified case(s) in residents or visitors.

242

243 There was a higher association between wastewater concentration and case incidence by report
244 date, as compared to episode date (primarily reflecting symptom onset date). This suggests both that
245 wastewater surveillance results appear timely relative to when cases are known to public health, as
246 well as a lag between symptom onset and detection of MPXV DNA in wastewater, potentially
247 reflecting delayed viral shedding into wastewater. Further research on MPXV shedding is necessary to
248 describe how viral shedding may affect lag and vary by disease severity, and to improve estimates of
249 the number of cases based on a wastewater concentration.

250

251 Wastewater samples regularly collected for routine public health monitoring can be tested for new
252 targets with minimal change in processes. Our findings suggest wastewater can be used to effectively
253 monitor for the introduction of MPXV and track its circulation for public health, clinical, and public
254 awareness. The rapid adaptation of wastewater surveillance infrastructure to effectively monitor a
255 non-enteric, non-respiratory virus such as MPXV, is promising for the future use of this tool for
256 emerging infectious diseases of public health concern. Increased and sustained investment is needed
257 to build wastewater surveillance infrastructure and for it to be ready to adapt rapidly to meet new
258 and emerging public health needs.

259

260

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271

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276 *Disclaimer:* The findings and conclusions in this report are those of the authors and do not necessarily
277 represent the official positions of the California Department of Public Health or the California Health
278 and Human Services Agency.

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281 represent the official positions of the Centers for Disease Control and Prevention.

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283 §See e.g., 45 C.F.R. part 46, 21 C.F.R. part 56; 42 U.S.C. §241(d); 5 U.S.C. §552a; 44 U.S.C. §3501 et
284 seq.

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Table 1. First cases of monkeypox and first detection of monkeypox virus (MPXV) DNA in wastewater by sewershed, California.

Sewersheds serviced by publicly owned treatment works (POTWs): SJ (San Jose), PA (Palo Alto), Gil (Gilroy), Sun (Sunnyvale), SVCW (Silicon Valley Clean Water), OSP (Oceanside, San Francisco), SEP (Southeast, San Francisco), Sac (Sacramento), Dav (Davis). Symptom onset date of the first case in each sewershed was available and thus reported. Total cases using episode date (symptom onset date, or if not available, the earliest of laboratory result or case record creation date) and report date are noted separately for each sewershed within the study period (June 19 to August 1, 2022).

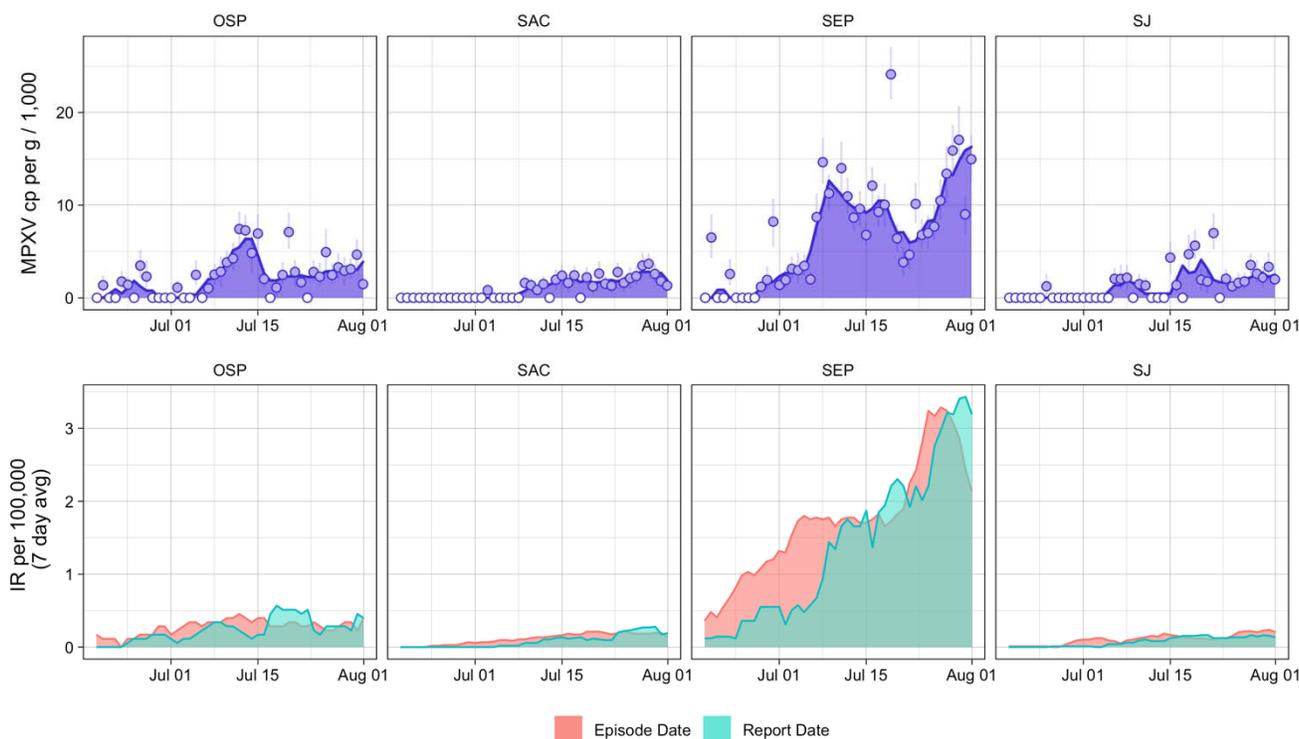
Sewershed	Symptom onset date of first case	Report date of first case	First detection date in wastewater	Total cases, by episode date	Total cases, by report date
SJ	6/22/22	6/29/22	6/25/22	74	59
PA	6/16/22	6/22/22	7/15/22	6	6
Gil	7/3/22	7/11/22	6/27/22	1	1
Sun	7/2/22	7/8/22	7/9/22	5	5
SVCW	7/20/22	7/20/22	7/21/22	5	4
OSP	5/29/22	6/10/22	6/20/22	36	30
SEP	5/30/22	6/3/22	6/20/22	494	442
SAC	5/12/22	5/23/22	7/3/22	82	67
Dav	n/a	n/a	6/27/22	0	0

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309 **Figure 2. Wastewater monkeypox virus (MPXV) concentrations and incidence of monkeypox cases**
310 **by sewershed.**

311 Top row: Time series of wastewater concentrations (concentration of MPXV DNA normalized by
312 concentration of PMMoV RNA) at select publicly owned treatment works (POTWs) with >10 positive
313 detections during the study time period: SEP (Southeast, San Francisco), OSP (Oceanside, San
314 Francisco), Sac (Sacramento), and SJ (San Jose). The area under the curve represents the 5 day
315 trimmed average of MPXV DNA cp/g over PMMoV cp/g in wastewater. Points represent daily values;
316 open circles indicate non-detects. Error bars represent standard deviations and include Poisson error
317 and variability among the 10 replicates (68% confidence intervals reported by the instrument
318 software as “total error”). Bottom row: Daily incidence rate (IR) or monkeypox cases, averaged over
319 7-days, using episode date (red) and report date (green).

320

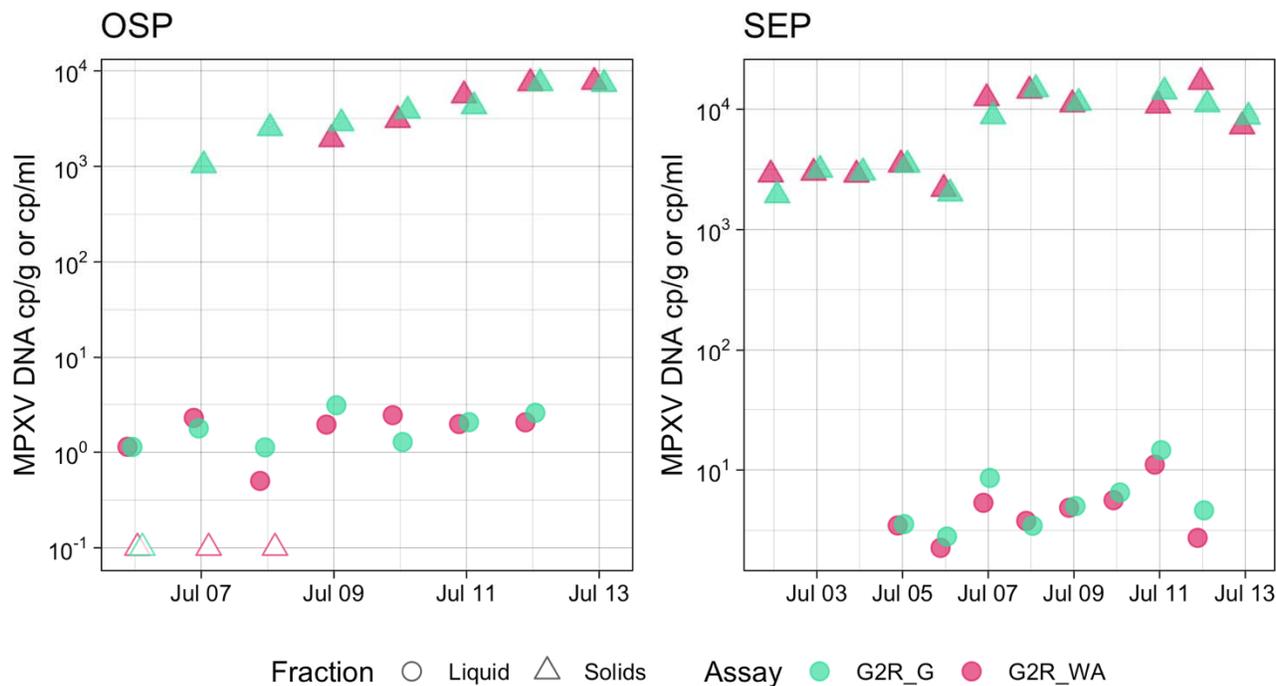


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324 **Figure 3. Concentrations of monkeypox virus (MPXV) in wastewater solids and liquid influent by**
325 **assay.**

326 Concentrations of G2R_G and G2R_WA in wastewater solids and liquid influent at OSP (Oceanside,
327 San Francisco) and SEP (Southeast, San Francisco) POTWs. Units are per gram for measurements
328 using wastewater solids, and per mL for measurements from liquid wastewater.
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334

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340 [health-regulations-\(2005\)-\(ihr\)-emergency-committee-regarding-the-multi-country-outbreak-of-](https://www.who.int/news/item/23-07-2022-second-meeting-of-the-international-health-regulations-(2005)-(ihr)-emergency-committee-regarding-the-multi-country-outbreak-of-monkeypox)
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