

June 11, 2020

## Big Sky Wastewater Testing Results

### Result Summary: Big Sky samples were negative but trending toward positive

#### Sample Description:

A composite sample of wastewater (1.5 L total) inflow to the Big Sky treatment plant was captured on 6/9/2020 using an auto-sampler over the previous 24-hour period. Referred to below as “Inflow” samples.

#### Testing Information and Raw Data:

Testing for the presence and abundance of the SARS-CoV2 genome in the above samples was performed using a kit designed by the US Centers for Disease Control and Prevention (CDC 2019-Novel Coronavirus (2019-nCoV), Real-Time RT-PCR Diagnostic Panel). Importantly, this test kit was originally designed to detect the virus in human samples and NOT wastewater or other kinds of environmental samples. The test was used here to determine whether a detectable amount of virus was present. Results need to be interpreted with caution, as described below.

Each of the above samples were split and processed as three replicates. Two tests were performed on each replicate and two independent locations on the SARS-CoV2 genome were targeted (N1 and N2). RNA was isolated from inactivated/concentrated samples, reverse-transcribed to DNA and used as template in quantitative PCR reactions as per kit instructions. Results were recorded as cycle threshold (Ct) numbers. All Ct numbers above (greater than) 40 cycles were highlighted in the table. Based on test interpretation guidelines described by the CDC (see below) Ct numbers greater than 40 should not be considered positive.

Results were as follows:

| Big Sky   |              |        |         | Potential Genomes | per |
|-----------|--------------|--------|---------|-------------------|-----|
| Sample ID | Replicate ID | Target | Ct      | liter             |     |
| Inflow_1  | Inflow_1.1   | N1     | NA      | NA                |     |
| Inflow_1  | Inflow_1.1   | N2     | 39.6113 | 573               |     |
| Inflow_1  | Inflow_1.2   | N1     | 37.4776 | 404               |     |
| Inflow_1  | Inflow_1.2   | N2     | 41.0313 | 207               |     |
| Inflow_2  | Inflow_2.1   | N1     | 37.3724 | 439               |     |
| Inflow_2  | Inflow_2.1   | N2     | 39.8631 | 478               |     |
| Inflow_2  | Inflow_2.2   | N1     | 38.115  | 242               |     |
| Inflow_2  | Inflow_2.2   | N2     | 40.617  | 279               |     |
| Inflow_3  | Inflow_3.1   | N1     | 37.9026 | 287               |     |
| Inflow_3  | Inflow_3.1   | N2     | 40.258  | 360               |     |
| Inflow_3  | Inflow_3.2   | N1     | 37.7093 | 335               |     |
| Inflow_3  | Inflow_3.2   | N2     | 41.6319 | 135               |     |

#### Interpretation:

We detected a concerning signal increase in all but one of the replicate wastewater inflow samples, particularly the N1 target, which unlike previous weeks of testing, was consistently detectable (i.e. above limit of detection). Signal from the N2 target was present but not consistently below the CDC recommended cut-off value of  $\leq 40$  Ct numbers. Such results might be expected if the amount of virus in wastewater approached the assay detection limit. Given our experience with testing environmental and wastewater samples and results from previous weeks of testing, we feel these results are consistent with an increasing trend in virus level that should be monitored closely.

#### Relevant text from CDC guidelines:

“...a specimen is considered positive for 2019-nCoV if all 2019-nCoV marker (N1, N2) cycle threshold growth curves cross the threshold line within 40.00 cycles ( $< 40.00$  Ct).”

“When all controls exhibit the expected performance and the cycle threshold growth curve for any one marker (N1 or N2 but not both markers) crosses the threshold line within 40.00 cycles ( $< 40.00$  Ct) the result is inconclusive.”