WEEKLY REPORT – WASTEWATER-BASED EPIDEMIOLOGICAL SURVEILLANCE OF THE SARS-COV-2 – RESULTS OF 02/11/2022

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In collaboration with:
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1. Summary

In the present work, the circulation of the SARS-CoV-2 virus in the environment is assessed based on three indicators. The analysis of wastewater samples collected in 42 wastewater treatment plants covers 45% of the Belgian population, with increased accuracy in the urban areas. The results of the wastewater surveillance are a source of complementary information to the infection cases number. Indeed, the wastewater results do notably include all asymptomatic persons, and are independent of the testing strategy.

Here are the conclusions based on the latest results of November 02th 2022:

• At the national level: Since two weeks, the Increasing trend indicator is stable, with approximately one quarter of the areas positive. The High Circulation trend indicator still remains at a low level with less than 5 areas positive. Overall, the wastewaters viral loads are at low levels compared to the 5th wave.
• At the regional level: None of the three indicators is positive in each of the three regions.
• At the provincial level: The Increasing Trend indicator is not positive in any province, and was in 1 last week. The Fast Increase indicator is not positive in any province, and was in none last week. The High Circulation indicator is not positive in any province, and was in none last week.
• At the covered areas level: among the 42 areas covered1, the number of treatment plants belonging to the different indicators are 8 for the Increasing Trend indicator, 1 for the Fast Increase indicator, and 0 for the High Circulation indicator.

The wastewater situation can be followed on a weekly basis on:

• The graphics available on the public COVID-19 dashboard
• The Risk Assessment Group (RAG) updating the weekly epidemiological situation through a report validated by the Risk Management Group (RMG) and published every Friday in French and Dutch.
• Further details on the methodology applied for the wastewater surveillance can be found in the Appendix Methodology document (access available online). The methodology was updated on June 20th 2022.

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1 Due to the 2021 flooding events, the treatment plants from Wegnez (Verviers) is temporarily out of order. This area is therefore not screened for SARS-CoV-2. Liège is still well represented within this surveillance, however Verviers is not. Since the 13th of October, the covered area of Soumagne is added to the surveillance. Due to technical issues, the areas of Bass Wavre (Dyle) and Vallée du Hain (L’Orchis) were not sampled this week.
2. Introduction

In mid-September 2020 started the SARS-CoV-2 national wastewater surveillance project. The present report is an outcome of this project aiming to assess weekly the wastewater-based epidemiological situation of Belgium. The surveillance is based on the analysis of water samples collected twice per week from the influent of 42 WasteWater Treatment Plants (WWTPs) spread over Belgium. The evolution of the SARS-CoV-2 wastewater concentrations measured over time in the different treatment plants is analysed at different levels: regional, provincial, and the catchment area covered by the individual treatment plants. Also, three alerting indicators were developed to highlight areas of concern regarding the high circulation, the fast evolution, and the increasing trend of the observed wastewater concentrations.

In this report, the weekly wastewater-based epidemiological situation is presented and discussed at the above-mentioned levels based on the three categories of alerting indicators. Moreover, the remaining sources of uncertainties are discussed together with their expected impacts on the wastewater results interpretation.

3. Methodology

3.1. SAMPLE COLLECTION AND ANALYSIS

The surveillance project, which started in mid-September 2020 covers around 45% of the Belgian population. The population covered is 40% in the Flemish region, 35% in the Walloon region, and nearly 100% in the Brussels region. Figure 1 shows the catchment areas covered by the 42 WWTPs included in the project, which are located in high population density areas. A catchment area is defined by the area delimiting the population covered by a specific wastewater sample, collected at the inlet of the WWTPs. Further coverage details can be found in Table 2 by province (see also Table A1 in the Appendix Methodology document (access available online) ) and on the Sciensano public dashboard. In practice, 24-hour composite samples are collected twice a week on Mondays and Wednesdays from the influent of WWTP and are analysed for the presence of SARS-CoV-2 RNA by three different laboratories. The resulting SARS-CoV-2 concentrations (3 targeted genes) are delivered to Sciensano within 2 days for data analysis purpose. Concretely, the results which are representative of Mondays and Wednesdays are respectively available on Wednesdays and Fridays.
3.2. WASTEWATER RESULTS

Between mid-September 2020 and mid-February 2021, the concentration were not quantified for all treatment plant and were expressed in Ct values. In mid-February 2021, the quantification of SARS-CoV-2 viral concentration (RNA copies/ml) was generalized to all the treatment plants investigated. Until 10th June 2022 the NIBSC standard was used to quantify the SARS-CoV-2 and after this date, the EURM standard was used.

In order to account for possible dilution by rainy events and for the number of people persons living in the catchment of each WWTP, a correction is applied on the viral concentration: the concentration is multiplied by the inlet flow and divided by the number of inhabitant equivalent of its respective WWTP and expressed by 100k inhabitant. The units of viral loads are copy/day/100k inhabitants.

Preliminary results estimated the limit of quantification of the analytical method at 20 copies/ml or $2 \times 10^{12}$ copy/day/100k inhab.

3.3. ALERTING INDICATORS

To highlight the areas of possible concern, the three alerting indicators are assessed twice a week, based on viral loads (RNA copies/day/100k inhabitants) measured for the three targeted gene fragments (E, N1, and N2):

1. The **Increasing Trend** indicator highlights the catchment areas where the viral loads have been increasing for more than 6 days. The indicator is computed based on the moving average on the past two weeks of the viral load.

2. The **Fast Increase** indicator highlights the catchment areas where the viral loads have rapidly increased for the last week. It corresponds to a situation where the moving average on the past
7 days of the viral load has increased faster than 70% per week if being above the estimated limit of quantification. The increasing slope is normalized for each treatment plant.

3. The **High Circulation** indicator highlights the catchment areas where the viral loads are high. It corresponds to a situation where the viral loads exceed half of the highest value recorded during the fifth wave (i.e. from first of January 2022 till first of March 2022).

The indicators were developed in order to be able to track the different phases of an outbreak. Typically, when the viral loads in wastewaters in an area start to increase, the indicator Increasing Trend will be the first fulfilled. If the viral loads increase quickly, the Fast Increase indicator will be fulfilled. Finally, after the initial increasing phase, the viral loads in an area may be sufficiently high to result in the High Circulation to be fulfilled.

### 3.4. CASES RESULTS

The cases number data presented in this report come from the COVID-19 laboratory-confirmed cases database centralized by Sciensano. The cases number, used to compute the 14 days-incidence, only accounts for the physical areas covered by the wastewater project (see Figure 1).

### 3.5. CAUTION POINTS FOR THE RESULTS INTERPRETATION

Only the trends, observed through the alerting indicators, should be assessed for the comparison of different areas. As the viral load values differ from a WWTP to another, notably due to the differences in population sizes covered, absolute values should not be compared. Additionally, the situation comparison between the regions should be considered with caution. The degree of comparability is not yet known and depends on the comparability of the results between the different laboratories performing the analyses (see the Appendix Methodology document (access online)).

Moreover, the wastewater viral loads and the cases numbers presented in this report do not originate from the same population, even though the positive cases are selected only for the municipalities covered by the wastewater surveillance. For instance, the wastewater results account for all infected persons (whatever age or symptomatic condition) while the cases numbers include only the persons clinically tested positive. Likewise, an infected person covered by the wastewater results could be associated with another area in the clinical surveillance as the person’s postal code is used for clinical statistics, i.e. mobility bias. Therefore, the correlation between the wastewater viral load and the cases number varies according to the area considered. The wastewater results are thus complementary and independent information to the results of the cases.

The correlation between the wastewater viral loads and the cases numbers could also be influenced by the vaccination campaign and the circulation of variants.
4. Results

4.1. NATIONAL LEVEL

Table 1 shows the national results allowing to track the changes between the situation as of today (November 02th 2022) and the situation as of last week (October 26th 2022). In this table, any change in indicator fulfilment (i.e. if the value for any indicator has changed from 0 to 1 or from 1 to 0) is indicated in **coloured bold text**.

None of the three indicators is positive at the national level.

<table>
<thead>
<tr>
<th>Nation</th>
<th>Pop. coverage</th>
<th>High</th>
<th>Fast</th>
<th>Incr.</th>
<th>Norm. viral load (%)(^1)</th>
<th>Mean viral load(^2)</th>
<th>Norm. evol. (%/week)(^3)</th>
<th>Incr. days(^4)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Belgium</td>
<td>44%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>4.9</td>
<td>0.74</td>
<td>-22</td>
<td>2</td>
</tr>
</tbody>
</table>

1: the viral load normalized with the maximum viral load measured in the corresponding catchment area during the fifth wave (i.e. from first of January 2022 till the first of march 2022).
2: the viral load computed on the mean of the replicate of the three targeted gene fragments as explained in section “3.2 Wastewater results”. The mean viral load is expressed in \(10^{12}\) copies/day/100k inhabitants.
3: the slope (%/week) of the past 7 days moving average of the viral load if the corresponding concentration is above the estimated limit of quantification.
4: the cumulative number of days of increase of the past 14 days moving average of the viral load.

Figure 2 was developed to show the localization of the covered areas by the wastewater surveillance in Belgium and the state of the three alerting indicators in each of them. Figure 3 was developed to offer a dynamic view of the three alerting indicators over time. For further insights on the dynamic of the different indicators, see Section 3.3.

Since two weeks, the Increasing trend indicator is stable, with approximately one quarter of the areas positive. The High Circulation trend indicator still remains at a low level with less than 5 areas positive. Overall, the wastewaters viral loads are at low levels compared to the 5th wave.
Figure 2: Localisation of the covered areas (grey polygons) by the wastewater surveillance in Belgium. In each area, the fulfillment of the High Circulation, Fast Increase and Increasing Trend indicator is indicated by a pink, blue and yellow slice, respectively. The names of the covered areas with respect of their localization can be found in Figure 1.
Figure 3: The number of areas (among the 42 covered by the wastewater surveillance this week and the 42 normally considered), with positive alerting indicators (black) compared to the total number of areas considered every week in the wastewater surveillance (white). Latest results (last bar) are from November 02\textsuperscript{nd} 2022. The white bars represent the total number of areas considered every week in the wastewater surveillance. Before the date marked by the vertical dashed line, the NIBSC standard was used to quantify the SARS-CoV-2 and after this date, the EURM standard was used.
4.2. REGIONAL LEVEL

Figure 4 represents the quantitative SARS-CoV-2 RNA viral loads in the wastewaters and the 14 days incidence for each region, compiling the incidence data of the area covered by the wastewater surveillance.

Several waves can be seen in Figure 4:

- The 5th wave starting on 27th December 2022.
- The 6th wave starting on 28th February 2022.
- The 7th wave starting on 30th May 2022.

Table 2 shows, for each region, the results associated with the samples of Wednesday November 02th, 2022. It allows to track the changes between the situation as of today (November 02th, 2022) and the situation as of last week (October 26th, 2022). Hereby, two distinct cases are taken into account:

1. If a region has at least one indicator fulfilled this week and it was not the case last week, its **name is displayed in bold** in the table.
2. If a region has at least one indicator fulfilled this week and if it also was the case last week, any change in indicator fulfilment (i.e. if the value for any indicator has changed from 0 to 1 or from 1 to 0) is indicated in **coloured bold text**.

None of the three indicators is positive in each of the three regions.
Table 2: Alerting indicators fulfilled (1) or not (0) on November 02\textsuperscript{nd} 2022. Columns represent the population coverage of the regions (Pop. coverage) and the three alerting indicators High Circulation (High), Fast Increase (Fast) and Increasing Trend (Incr.). The specifications of the four last columns are explained in the footnotes 1-4 below the table. Missing data is indicated with a “/”.

<table>
<thead>
<tr>
<th>Region</th>
<th>Pop. coverage</th>
<th>High</th>
<th>Fast</th>
<th>Incr.</th>
<th>Norm. viral load (%)	extsuperscript{1}</th>
<th>Mean viral load\textsuperscript{2}</th>
<th>Norm. evol. (%/week)	extsuperscript{3}</th>
<th>Incr. days\textsuperscript{4}</th>
</tr>
</thead>
<tbody>
<tr>
<td>Brussels</td>
<td>100%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>3.7</td>
<td>1.8</td>
<td>-58</td>
<td>2</td>
</tr>
<tr>
<td>Flanders</td>
<td>41%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>3.4</td>
<td>0.3</td>
<td>-14</td>
<td>1</td>
</tr>
<tr>
<td>Wallonia</td>
<td>31%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>10.3</td>
<td>0.4</td>
<td>7</td>
<td>3</td>
</tr>
</tbody>
</table>

\textsuperscript{1}: the viral load normalized with the maximum viral load measured in the corresponding catchment area during the fifth wave (i.e. from first of January 2022 till the first of March 2022).
\textsuperscript{2}: the viral load computed on the mean of the replicate of the three targeted gene fragments as explained in section “3.2 Wastewater results”. The mean viral load is expressed in $10^{12}$ copies/day/100k inhabitants.
\textsuperscript{3}: the slope (%/week) of the past 7 days moving average of the viral load if the corresponding concentration is above the estimated limit of quantification.
\textsuperscript{4}: the cumulative number of days of increase of the past 14 days moving average of the viral load.
4.3. PROVINCIAL LEVEL

Table 3 shows, for each Province, the results associated with the samples of Wednesday November 02\textsuperscript{nd} 2022. It allows to track the changes between the situation as of today (November 02\textsuperscript{nd} 2022) and the situation as of last week (October 26\textsuperscript{th} 2022). Hereby, two distinct cases are taken into account:

1. If a province has at least one indicator fulfilled this week and it was not the case last week, its name is displayed in **bold** in the table.
2. If a province has at least one indicator fulfilled this week and if it also was the case last week, any change in indicator fulfilment (i.e. if the value for any indicator has changed from 0 to 1 or from 1 to 0) is indicated in **coloured** **bold text**.

Table 3 shows, for each Province, the results associated with the samples of Wednesday November 02\textsuperscript{nd} 2022, for the three alerting indicators:

- The Increasing Trend indicator is not positive in any province, and was in 1 last week.
- The Fast Increase indicator is not positive in any province, and was in none last week.
- The High Circulation indicator is not positive in any province, and was in none last week.
- Due to technical issues, the areas of Bass Wavre (Dyle) and Vallée du Hain (L’Orchis) were not sampled this week. The situation in the province of Brabant Wallon could hence not be assessed.

Table 3: Alerting indicators fulfilled (1) or not (0) on November 02\textsuperscript{nd} 2022. Columns represent the population coverage of the WWTPs within the Province (Pop. coverage) and the three alerting indicators High Circulation (High), Fast Increase (Fast) and Increasing Trend (Incr.). The specifications of the four last columns are explained in the footnotes 1-4 below the table. Missing data is indicated with a “/”.

<table>
<thead>
<tr>
<th>Province</th>
<th>Pop. coverage</th>
<th>High</th>
<th>Fast</th>
<th>Incr.</th>
<th>Norm. viral load(^1)</th>
<th>Mean viral load(^2)</th>
<th>Norm. evol. (%/w)(^3)</th>
<th>Incr. days(^4)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Antwerpen</td>
<td>41%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0.0</td>
<td>0.00</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Brabant Wallon</td>
<td>33%</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>/</td>
</tr>
<tr>
<td>Brussels</td>
<td>100%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>3.7</td>
<td>1.82</td>
<td>-58</td>
<td>2</td>
</tr>
<tr>
<td>Hainaut</td>
<td>29%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>16.2</td>
<td>0.59</td>
<td>16</td>
<td>5</td>
</tr>
<tr>
<td>Liège</td>
<td>50%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>5.9</td>
<td>0.29</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>Limburg</td>
<td>26%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0.1</td>
<td>0.01</td>
<td>0</td>
<td>2</td>
</tr>
<tr>
<td>Luxembourg</td>
<td>10%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>2.6</td>
<td>0.24</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>Namur</td>
<td>23%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>10.4</td>
<td>0.29</td>
<td>0</td>
<td>6</td>
</tr>
<tr>
<td>Oost-Vlaanderen</td>
<td>38%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>3.7</td>
<td>0.27</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>Vlaams-Brabant</td>
<td>61%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>4.2</td>
<td>0.52</td>
<td>-45</td>
<td>1</td>
</tr>
<tr>
<td>West-Vlaanderen</td>
<td>52%</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>7.8</td>
<td>0.64</td>
<td>0</td>
<td>2</td>
</tr>
</tbody>
</table>

\(^1\) the viral load normalized with the maximum viral load measured in the corresponding catchment area during the fifth wave (i.e. from first of January 2022 till the first of march 2022).

\(^2\) the viral load computed on the mean of the replicates of the three targeted gene fragments as explained in section “3.2 Wastewater results”. The mean viral load is expressed in \(10^{12}\) copies/day/100k inhabitants.

\(^3\) the slope (%/week) of the past 7 days moving average of the viral load if the corresponding concentration is above the estimated limit of quantification.

\(^4\) the cumulative number of days of increase of the past 14 days moving average of the viral load.
4.4. INDIVIDUAL CATCHMENT AREAS LEVEL

Table 4 shows, for each catchment area, the values of the three alerting indicators obtained based on the results of last Wednesday’s sample. Table 4 is a snapshot of the number of areas highlighted by the indicators for the last results obtained. It allows to track the changes between the situation as of today (November 02th 2022) and the situation as of last week (October 26th 2022). Hereby, three distinct cases are taken into account:

1. If an area has at least one indicator fulfilled this week and it was not the case last week, its name is displayed in **bold** in the table.
2. If an area has at least one indicator fulfilled this week and if it also was the case last week, any change in indicator fulfilment (i.e. if the value for any indicator has changed from 0 to 1 or from 1 to 0) is indicated in *coloured bold text*.
3. Any area which had at least one indicator fulfilled last week but not this week is listed below Table 4.

Here are the results associated with the samples of November 02th 2022:

- The Increasing Trend indicator is positive in 8 covered areas. Last week, this indicator was positive in 8 covered areas.
- The Fast Increase indicator is positive in only 1 covered area: Marchienne-au-Pont (127% increase per week) (see Appendix A2 for more details). Last week, this indicator was positive in 2 covered areas.
- The High Circulation indicator is not positive is any covered area. Last week, this indicator was not positive in any covered area.

The wastewater results at the level of the local covered areas can be accessed online for each area on the COVID-19 dashboard.

**Table 4: Alerting indicators fulfilled (1) or not (0) on November 02th 2022. Columns represent the provinces, different WWTPs within the Provinces and the three alerting indicators High Circulation (High), Fast Increase (Fast) and Increasing Trend (Incr.). The specifications of the four last columns are explained in the footnotes 1-4 below the table. Missing data is indicated with a “/”:**

<table>
<thead>
<tr>
<th>Province</th>
<th>WWTP</th>
<th>High</th>
<th>Fast</th>
<th>Incr.</th>
<th>Norm. viral load (%)</th>
<th>Mean viral load</th>
<th>Norm evol. (%/week)</th>
<th>Incr. days</th>
</tr>
</thead>
<tbody>
<tr>
<td>Liège</td>
<td>Amay</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>4.4</td>
<td>0.25</td>
<td>0</td>
<td>7</td>
</tr>
<tr>
<td>Limburg</td>
<td>Genk</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>0.2</td>
<td>0.03</td>
<td>0</td>
<td>7</td>
</tr>
<tr>
<td>Liège</td>
<td>Liège (Grosses Battes)</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>8.4</td>
<td>0.84</td>
<td>5</td>
<td>9</td>
</tr>
<tr>
<td>Hainaut</td>
<td>Marchienne-au-Pont</td>
<td>0</td>
<td>1</td>
<td>1</td>
<td>16.4</td>
<td>0.90</td>
<td>127</td>
<td>7</td>
</tr>
<tr>
<td>Hainaut</td>
<td>Mouscron versant Espierres</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>20.9</td>
<td>1.71</td>
<td>67</td>
<td>9</td>
</tr>
<tr>
<td>Namur</td>
<td>Namur-Brumagne</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>13.6</td>
<td>0.36</td>
<td>0</td>
<td>9</td>
</tr>
<tr>
<td>Liège</td>
<td>Soumagne</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>2.2</td>
<td>0.30</td>
<td>34</td>
<td>16</td>
</tr>
<tr>
<td>Hainaut</td>
<td>Wasmuel</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>17.6</td>
<td>0.36</td>
<td>-4</td>
<td>7</td>
</tr>
</tbody>
</table>

1: the viral load normalized with the maximum viral load measured in the corresponding catchment area during the fifth wave (i.e. from first of January 2022 till the first of march 2022).
2: the viral load computed on the mean of the replicates of the three targeted gene fragments as explained in section “3.2 Wastewater results”; The mean viral load is expressed in 10^12 copies/day/100k inhabitants.
The following areas had the corresponding indicator fulfilled last week but not this week:

- Fast Increase indicator: no area.
- High Circulation indicator: no area.

Further details on covered area without fulfilled indicators can be found in Table A4.
5. Appendix – Areas classified by alerting indicator

Table A1: Covered areas (0 out of 42 on November 02\textsuperscript{th} 2022) characterized as High Circulation sorted in the descending order of importance.

No area was characterized as High Circulation this week

Table A2: Covered areas (1 out of 42 on the November 02\textsuperscript{th} 2022) characterized as Fast Increase sorted in the descending order of importance.

<table>
<thead>
<tr>
<th>Province</th>
<th>WWTP</th>
<th>High</th>
<th>Fast</th>
<th>Incr.</th>
<th>Norm. viral load (%)\textsuperscript{1}</th>
<th>Mean viral load\textsuperscript{2}</th>
<th>Norm evol. (%/we ek)\textsuperscript{3}</th>
<th>Incr. days\textsuperscript{4}</th>
<th>Date Max cc\textsuperscript{5}</th>
</tr>
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<tbody>
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<td>Hainaut</td>
<td>Marchienne-au-Pont</td>
<td>0</td>
<td>1</td>
<td>1</td>
<td>16.4</td>
<td>0.90</td>
<td>127</td>
<td>7</td>
<td>27/07/2022</td>
</tr>
</tbody>
</table>

1: the viral load normalized with the maximum viral load measured in the corresponding catchment area during the fifth wave (i.e. from first of January 2022 till the first of March 2022).
2: the viral load computed on the mean of the replicates of the three targeted gene fragments as explained in section “3.2 Wastewater results”. The mean viral load is expressed in 10\textsuperscript{12} copies/day/100k inhabitants.
3: the slope (%/week) of the past 7 days moving average of the viral load if the corresponding concentration is above the estimated limit of quantification.
4: the cumulative number of days of increase of the past 14 days moving average of the viral load.
5: date at which the measured viral load was the highest since the beginning of 2022. If the date was between the first of January 2022 and the first of March 2022, the date is considered to be during the fifth wave and mentioned as such.

Table A3: Covered areas (8 out of 42 on the November 02\textsuperscript{th} 2022) characterized as Increasing Trend sorted in the descending order of importance.

<table>
<thead>
<tr>
<th>Province</th>
<th>WWTP</th>
<th>High</th>
<th>Fast</th>
<th>Incr.</th>
<th>Norm. viral load (%)\textsuperscript{1}</th>
<th>Mean viral load\textsuperscript{2}</th>
<th>Norm evol. (%/we ek)\textsuperscript{3}</th>
<th>Incr. days\textsuperscript{4}</th>
<th>Date Max cc\textsuperscript{5}</th>
</tr>
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<tbody>
<tr>
<td>Liège</td>
<td>Soumagne</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>2.2</td>
<td>0.29</td>
<td>34</td>
<td>16</td>
<td>28/03/2022</td>
</tr>
<tr>
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<td>Liège (Grosses Battes)</td>
<td>0</td>
<td>0</td>
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<td>0.84</td>
<td>5</td>
<td>9</td>
<td>5\textsuperscript{th} wave</td>
</tr>
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<td>Mouscron versant Espieres</td>
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<td>0</td>
<td>1</td>
<td>20.9</td>
<td>1.71</td>
<td>67</td>
<td>9</td>
<td>27/04/2022</td>
</tr>
<tr>
<td>Namur</td>
<td>Namur-Brumagne</td>
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<td>1</td>
<td>13.6</td>
<td>0.36</td>
<td>0</td>
<td>9</td>
<td>11/07/2022</td>
</tr>
<tr>
<td>Liège</td>
<td>Amay</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>4.4</td>
<td>0.25</td>
<td>0</td>
<td>7</td>
<td>5\textsuperscript{th} wave</td>
</tr>
<tr>
<td>Limburg</td>
<td>Genk</td>
<td>0</td>
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<td>0.2</td>
<td>0.03</td>
<td>0</td>
<td>7</td>
<td>09/03/2022</td>
</tr>
<tr>
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<td>1</td>
<td>1</td>
<td>16.4</td>
<td>0.90</td>
<td>127</td>
<td>7</td>
<td>27/07/2022</td>
</tr>
<tr>
<td>Hainaut</td>
<td>Wasmuel</td>
<td>0</td>
<td>0</td>
<td>1</td>
<td>17.6</td>
<td>0.36</td>
<td>-4</td>
<td>7</td>
<td>13/07/2022</td>
</tr>
</tbody>
</table>

1: the viral load normalized with the maximum viral load measured in the corresponding catchment area during the fifth wave (i.e. from first of January 2022 till the first of March 2022).
2: the viral load computed on the mean of the replicates of the three targeted gene fragments as explained in section “3.2 Wastewater results”. The mean viral load is expressed in 10\textsuperscript{12} copies/day/100k inhabitants.
3: the slope (%/week) of the past 7 days moving average of the viral load if the corresponding concentration is above the estimated limit of quantification.
4: the cumulative number of days of increase of the past 14 days moving average of the viral load.
Table A4: Covered areas (34 out of 42 on the November 02\textsuperscript{th} 2022) in which no alerting indicator are fulfilled.

<table>
<thead>
<tr>
<th>Province</th>
<th>WWTP</th>
<th>High</th>
<th>Fast</th>
<th>Incr.</th>
<th>Norm. viral load (%)</th>
<th>Mean viral load\textsuperscript{2}</th>
<th>Norm. evol. (%/week)\textsuperscript{3}</th>
<th>Incr. days\textsuperscript{4}</th>
<th>Date Max \textsuperscript{cc}\textsuperscript{5}</th>
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</thead>
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<td>10.1</td>
<td>0.58</td>
<td>-83</td>
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<td>0</td>
<td>0</td>
<td>0.0</td>
<td>0.00</td>
<td>0</td>
<td>0</td>
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<td>0</td>
<td>0</td>
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<td>0.00</td>
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<tr>
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<td>/</td>
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<tr>
<td>Vlaams-Brabant</td>
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<td>0.32</td>
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</tr>
<tr>
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<td>Mean viral load</td>
<td>Norm evol. (%/week)</td>
<td>Incr. days</td>
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<td>5th wave</td>
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</tr>
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<td>Turnhout</td>
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<td>0</td>
<td>16/05/2022</td>
</tr>
<tr>
<td>Brabant Wallon</td>
<td>Vallée du Hain (L’Orchis)</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>/</td>
<td>11/07/2022</td>
</tr>
</tbody>
</table>

1: the viral load normalized with the maximum viral load measured in the corresponding catchment area during the fifth wave (i.e. from first of January 2022 till the first of March 2022).
2: the viral load computed on the mean of the replicates of the three targeted gene fragments as explained in section “3.2 Wastewater results”. The mean viral load is expressed in $10^{12}$ copies/day/100k inhabitants.
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5: date at which the measured viral load was the highest since the beginning of 2022. If the date was between the first of January 2022 and the first of March 2022, the date is considered to be during the fifth wave and mentioned as such.