

# Weekly Report for Influenza, Enterovirus, Respiratory Syncytial Virus (RSV), Norovirus, SARS-CoV-2 and Poliovirus in Wastewater: 6 March 2024

Samples taken up to and including: 28 February 2024

## Background

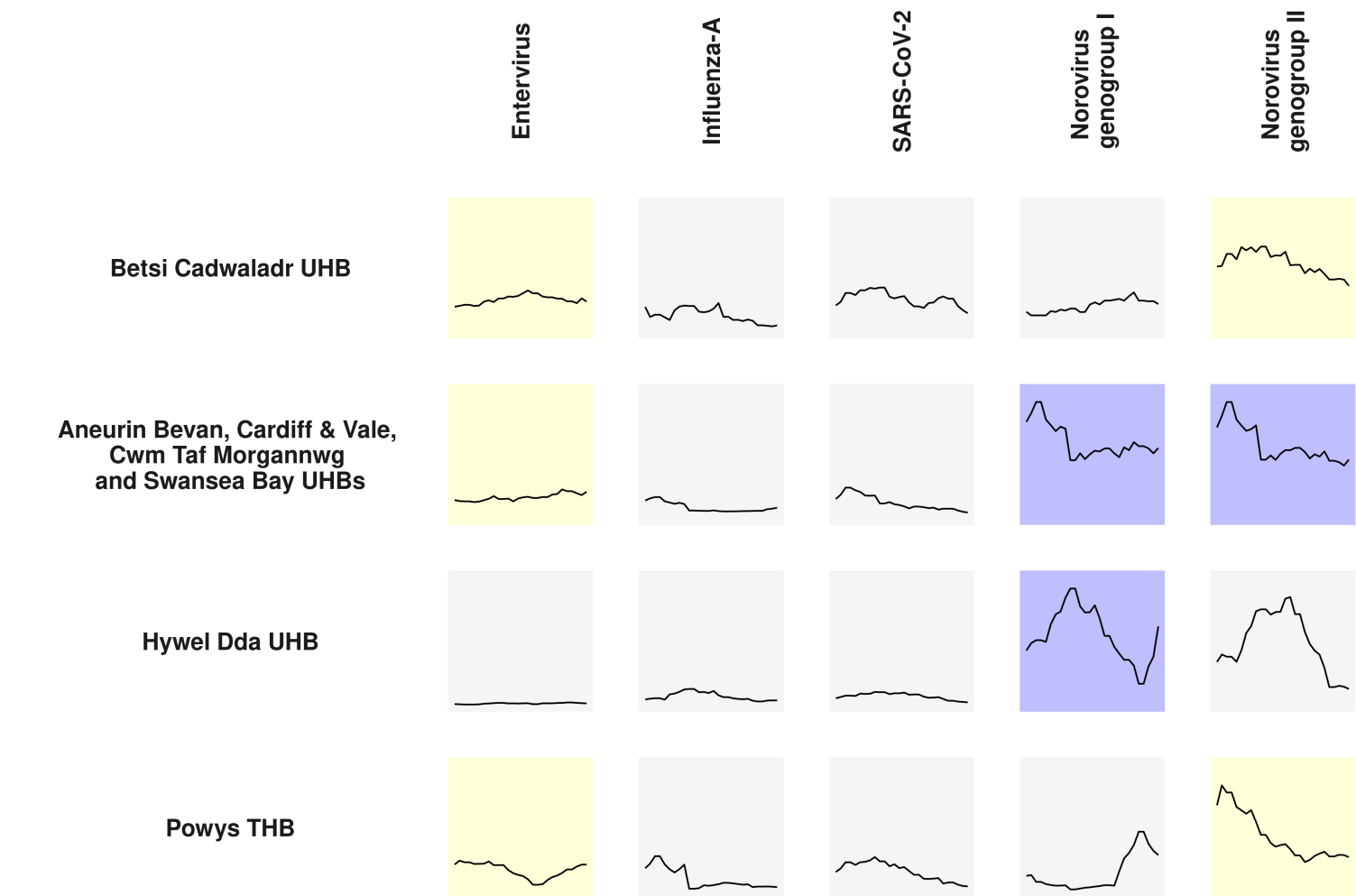
Wastewater samples are collected from 11 wastewater treatment works located across Wales (Monday to Wednesday) each week. Samples are representative of communities in each of the Welsh Health Boards. The concentrations of SARS-CoV-2 and other viruses are then quantified in the samples, as well as SARS-CoV-2 variants detected.

## Guidance

In this report, Cardiff & Vale, Aneurin Bevan, Cwm Taf Morgannwg and Swansea Bay University Health Boards are grouped together. Signal for all markers is given as a daily rate of gene copies per 100,000 people (gc/day per 100k individuals). Please check site level plots for greater detail. For plots showing enterovirus signal, the beige block represents a period where two enterovirus assays were being used (EV1 and EV2), with two lines being displayed. Currently, only the EV2 assay is being used as it provides better sensitivity. No samples were taken in the period between July 2023 and November 2023.



**Figure 1.** Map showing the 11 wastewater treatment plants (WwTPs) monitored by the programme, along with their respective health boards.



**Graphic 1.** Background shades indicate the average signal (%) for the most recent week: light grey (<20%), mid yellow (20%–40%), and dark blue (>40%). The average signal is calculated relative to the highest ever observed virus value per area. Lines represent the smoothed signal during the most recent 4 week period, derived from a 10-day rolling average, presented as a percentage of the highest observed virus value per area.

## Poliovirus

No poliovirus was detected in this sampling period. Poliovirus assays are run on samples from the first week of each month.

SARS-CoV-2

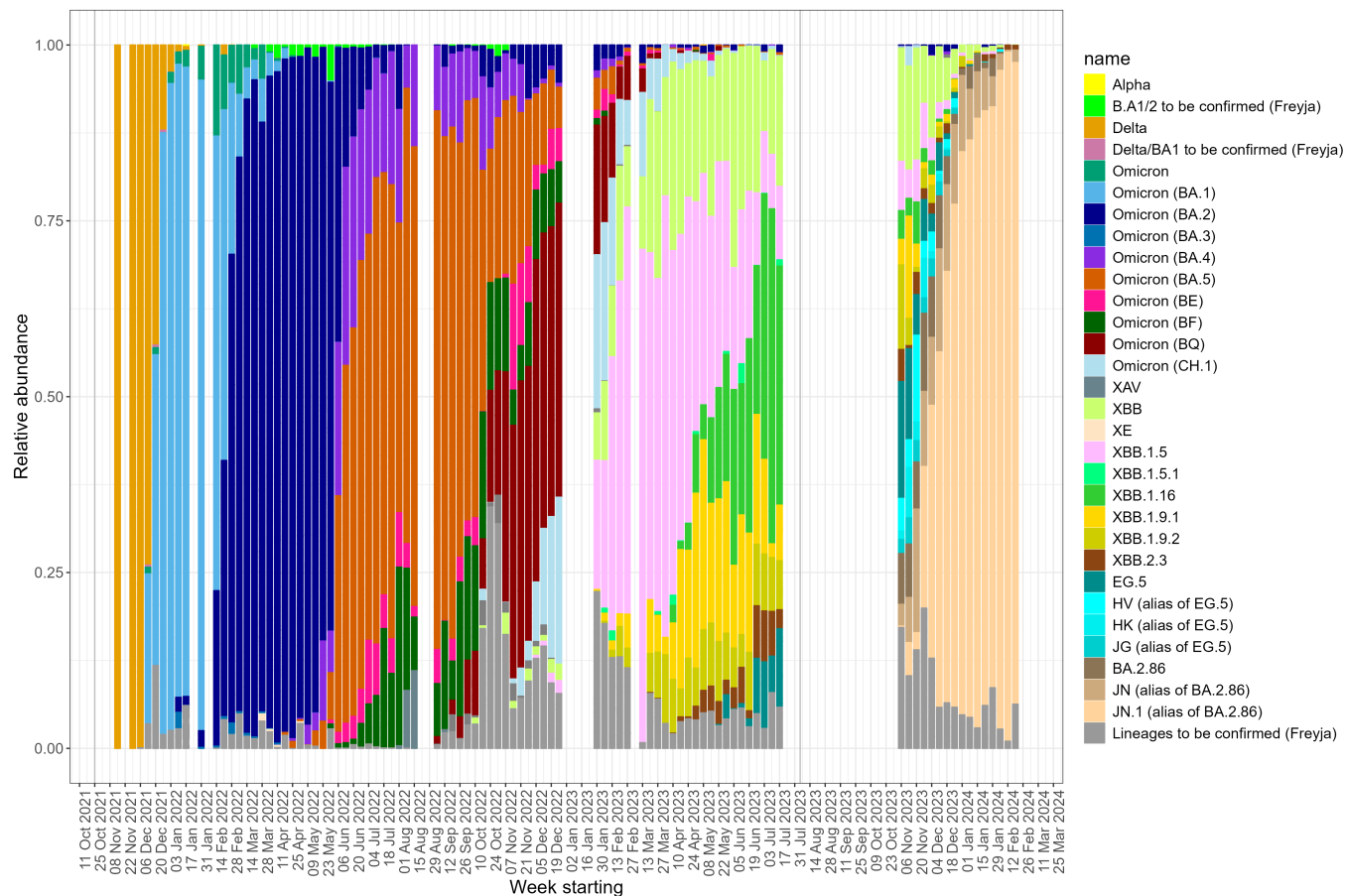
National Summary



**Figure 2.** Level of SARS-CoV-2 given as a 10 day rolling mean at the national (bold line) and healthboard (faint lines) level. An upper limit has been placed on the y axis, which obscures a high healthboard value but, allows better visualisation of the national trend.

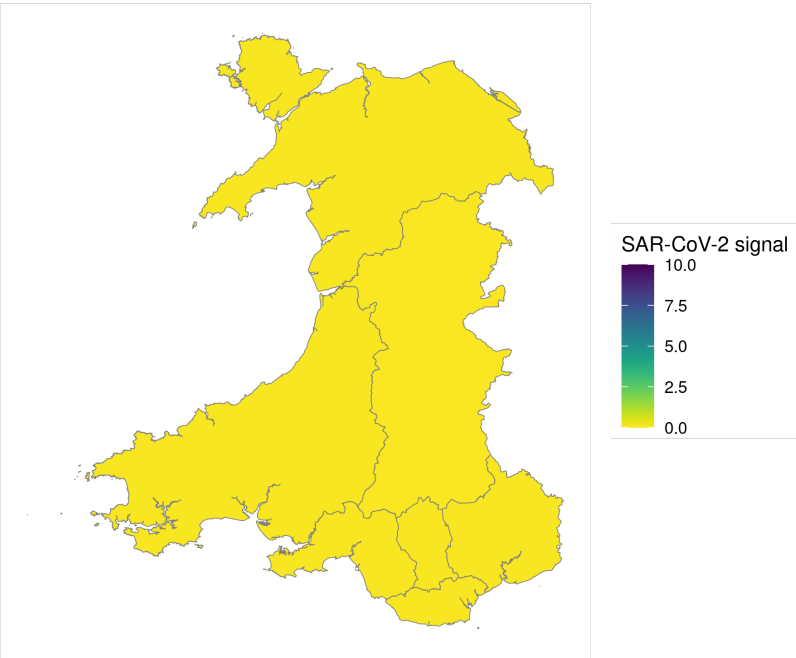


Variants of Concern and/or Interest



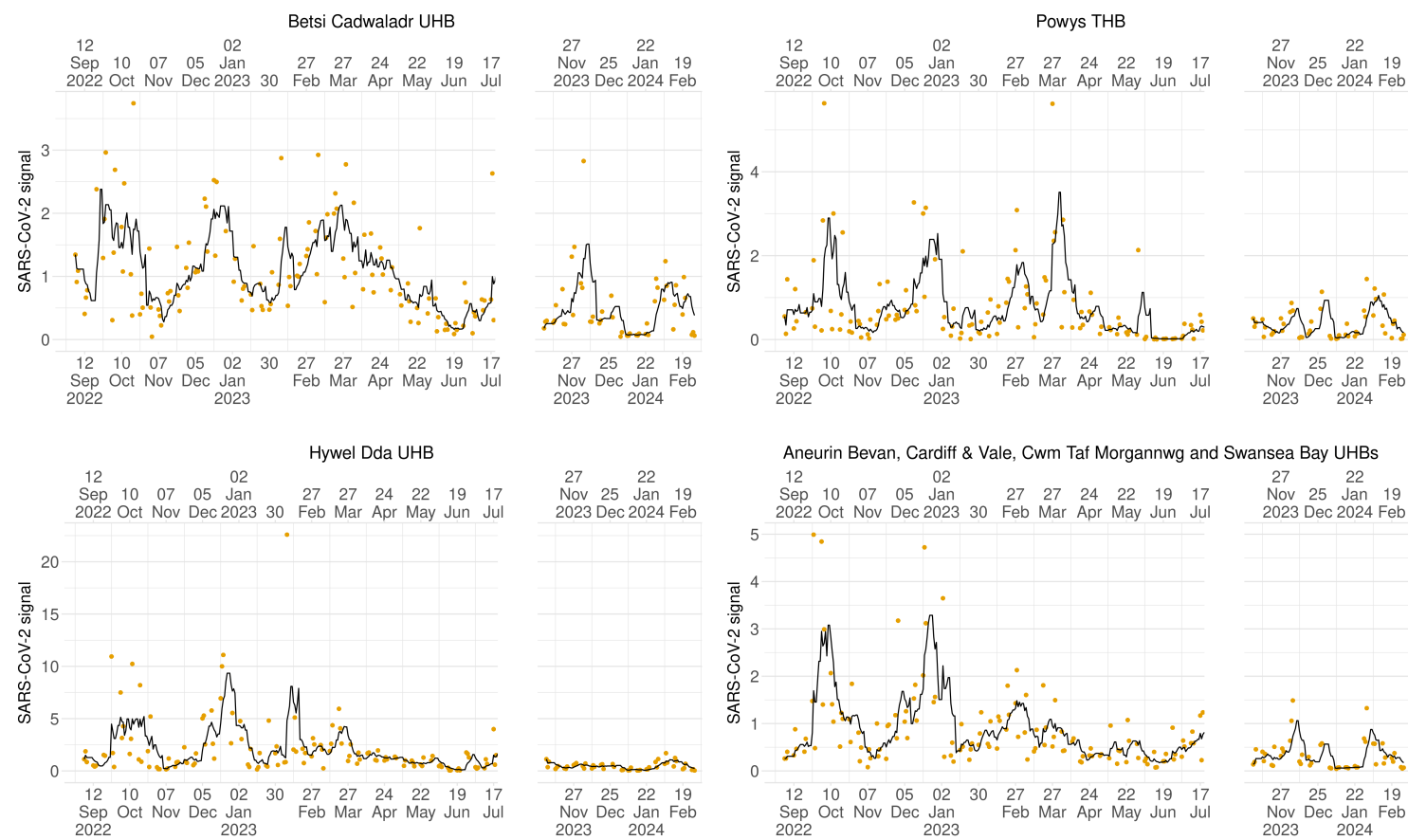
**Figure 3.** SARS-CoV-2 variants identified at the national level (plotting proportion of sequence reads assigned to each variant lineage using the **Freyja programme**; see figure legend). Limitations of the Freyja tool in reconstructing and quantifying SARS-CoV-2 lineages from wastewater metabarcoding sequencing should be considered. All sites contribute equally to the national lineage proportions, but a weighted national lineage plot can be found in the assets archive.

Heat Map



**Figure 4.** National heat map showing mean SARS-CoV-2 gc/day per 100k for each healthboard in the most recent sampling week.

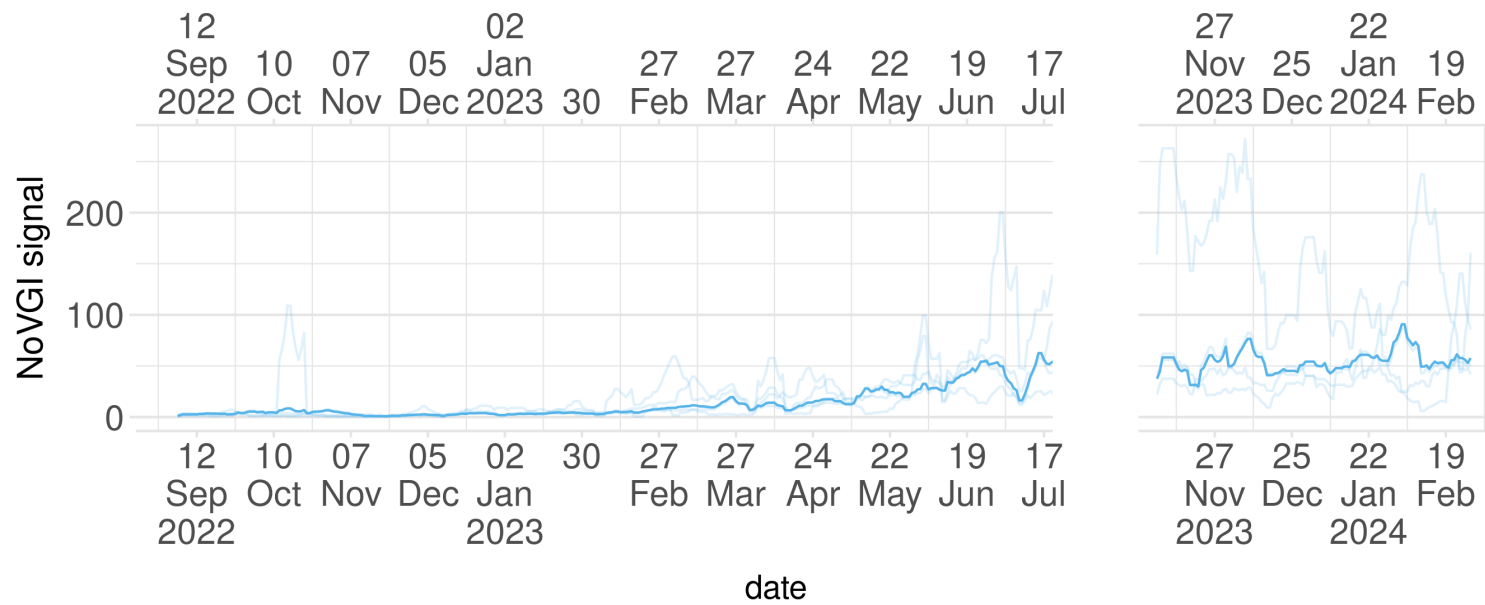
**Healthboard Summaries**



**Figure 5.** Level of SARS-CoV-2 given as a 10 day rolling mean at the healthboard level (bold line) and per day (dots).

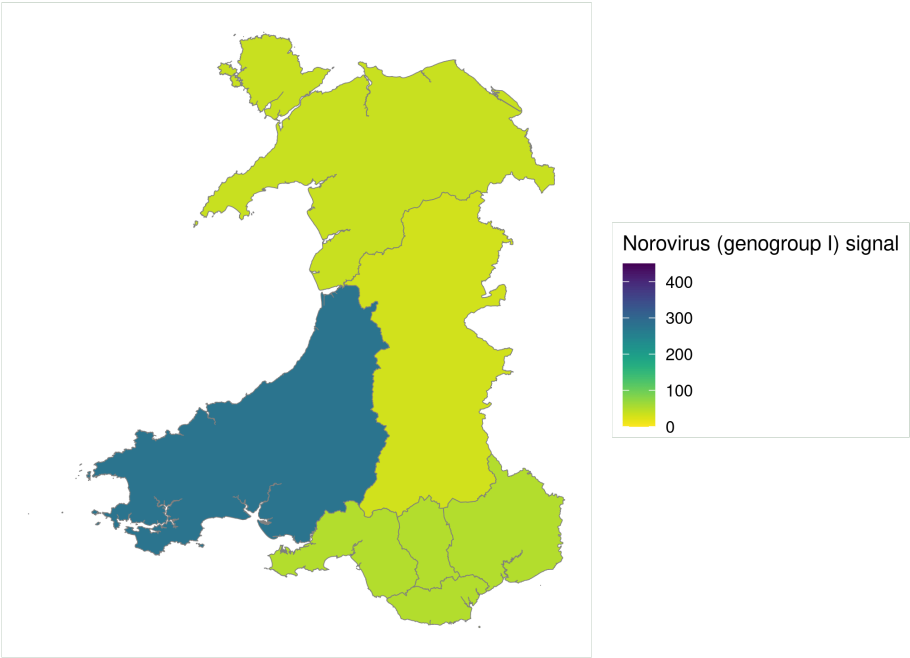
# Norovirus genogroup I (NoVGI)

## National Summary



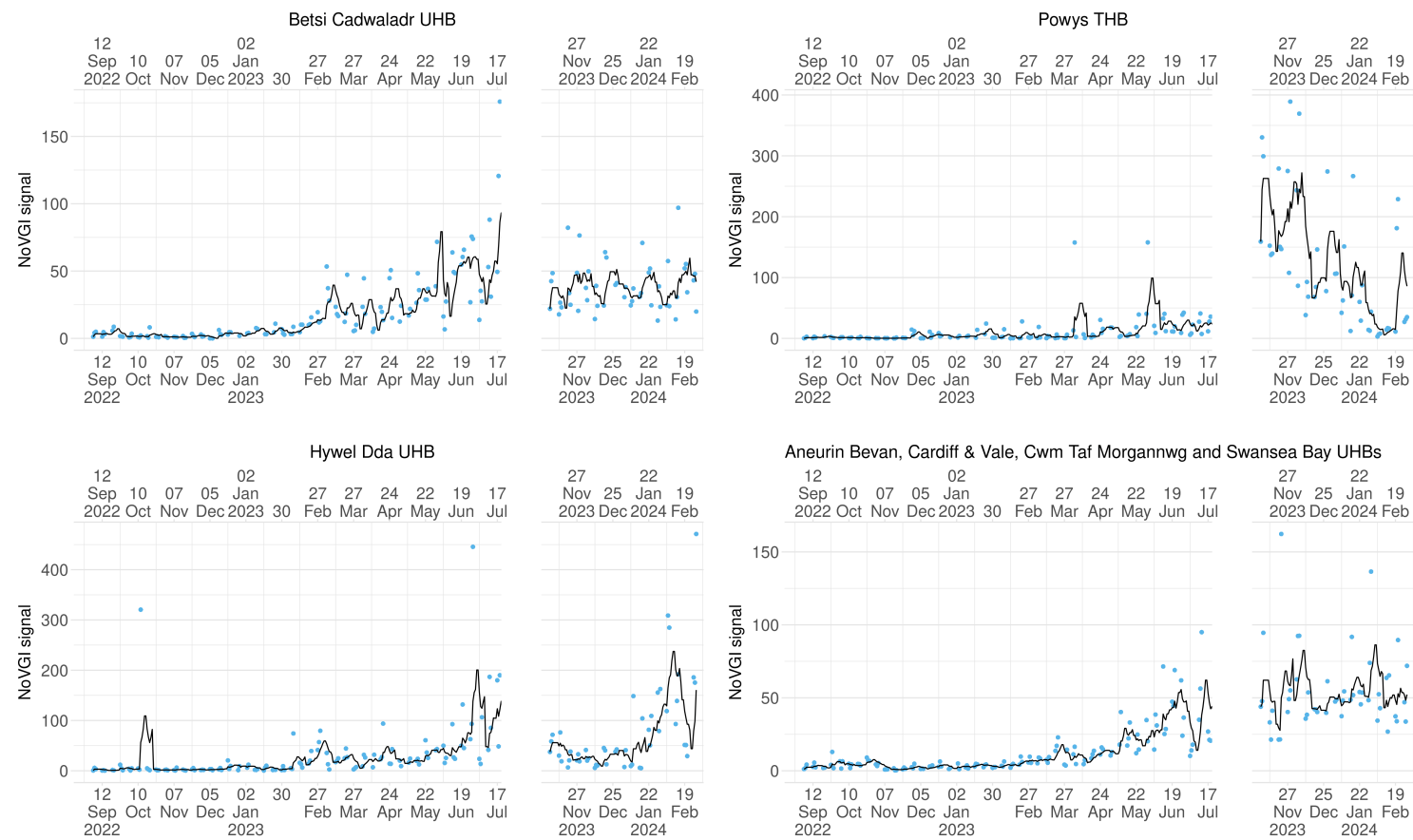
**Figure 6.** Level of norovirus genogroup I (NoVGI) given as a 10 day rolling mean at the national (bold line) and healthboard (faint lines) level.

## Heat Map



**Figure 7.** National heat map showing mean norovirus genogroup I gc/day per 100k for each healthboard in the most recent sampling week.

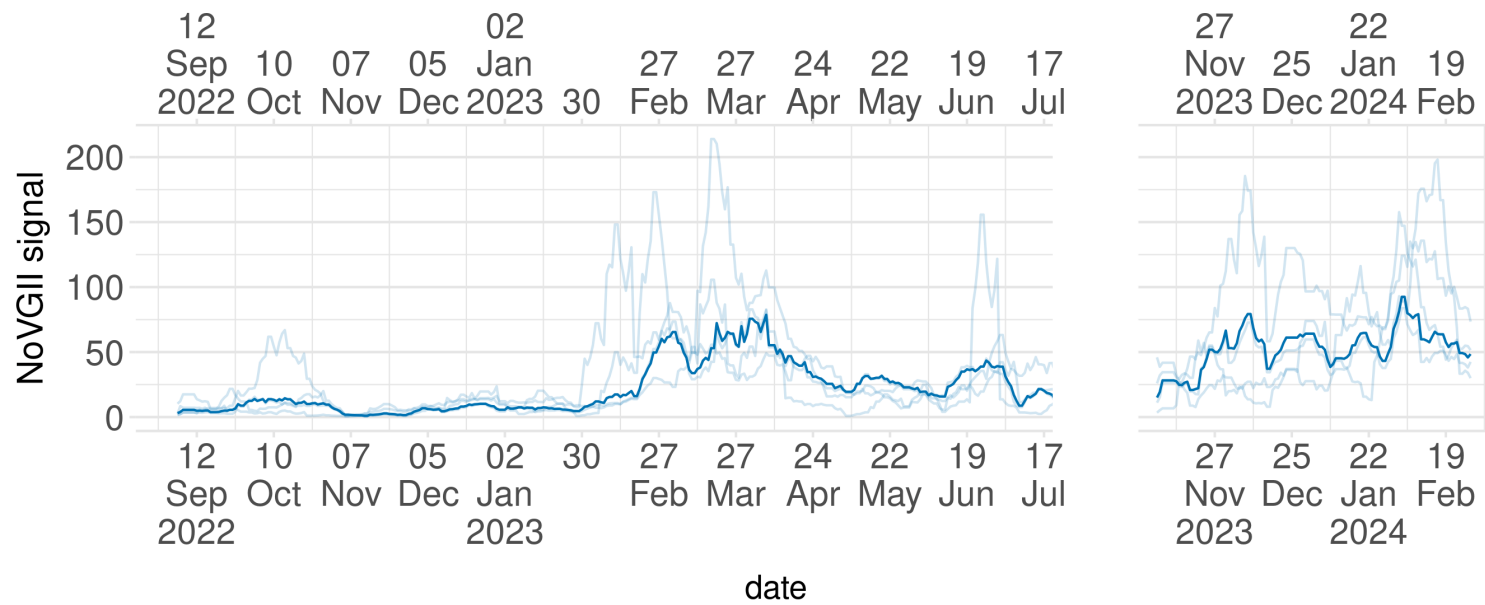
Healthboard Summaries



**Figure 8.** Level of norovirus genogroup I (NoVG1) given as a 10 day rolling mean at the healthboard level (bold line) and per day (dots).

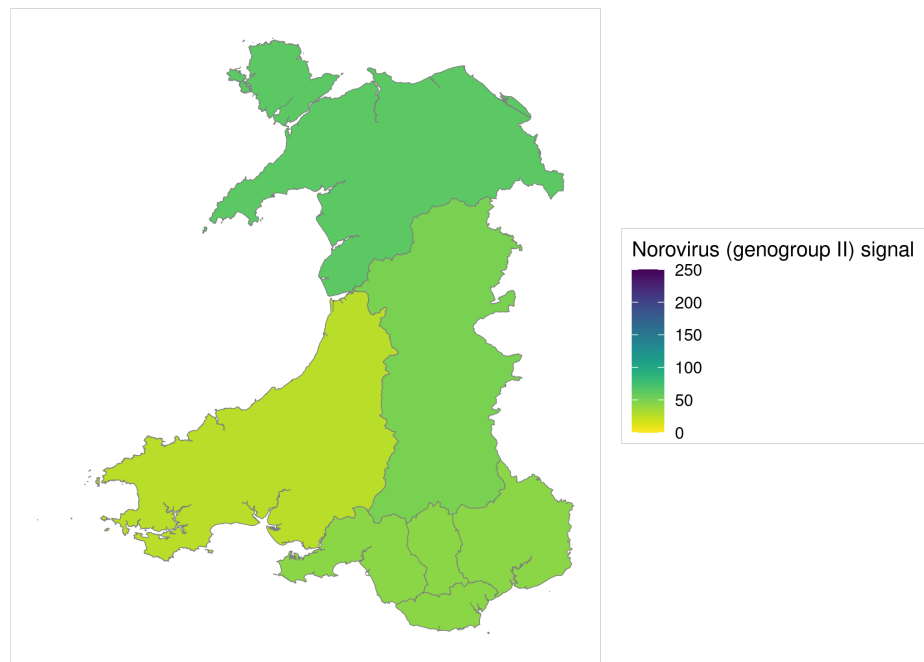
# Norovirus genogroup II (NoVGII)

## National Summary



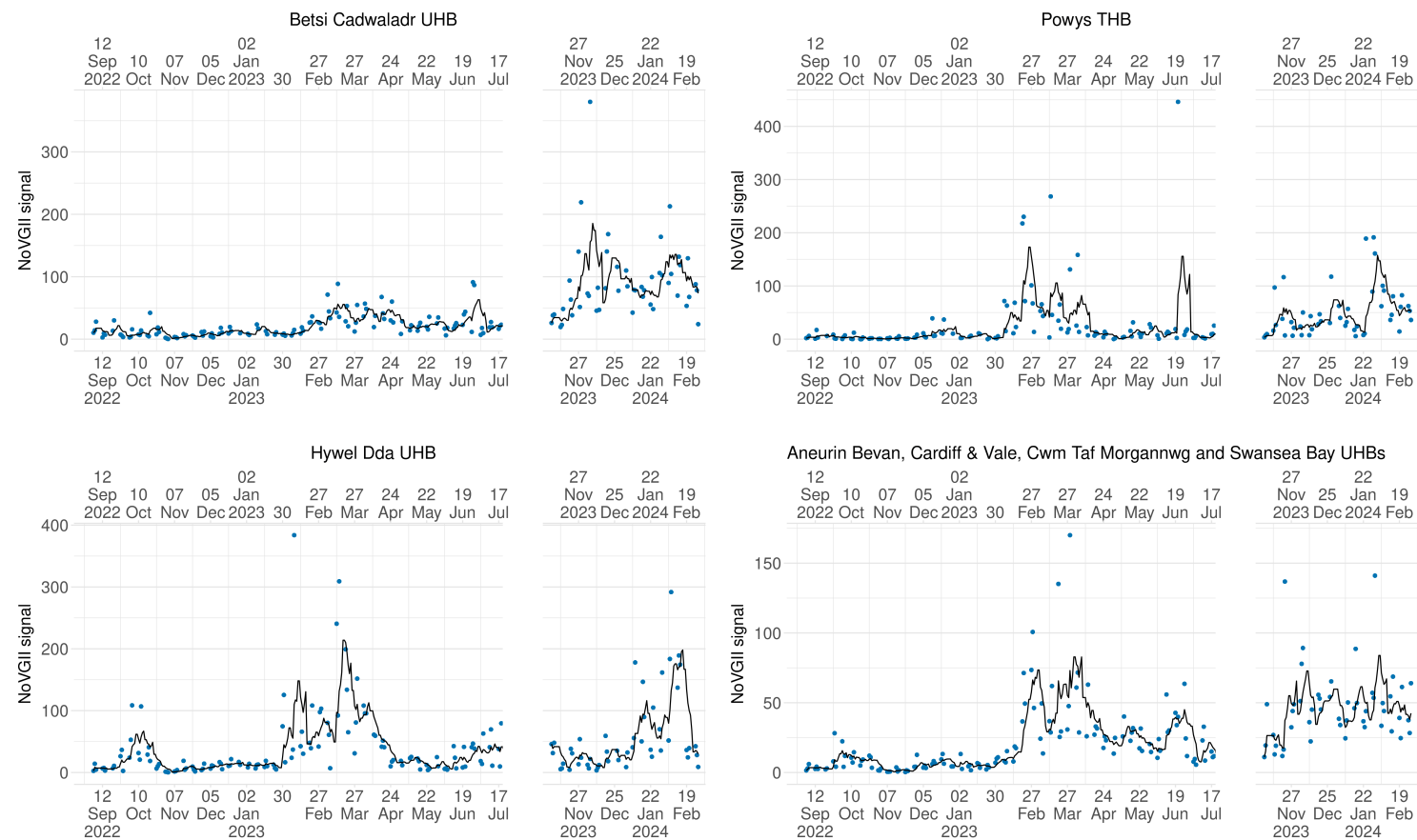
**Figure 9.** Level of norovirus genogroup II (NoVGII) given as a 10 day rolling mean at the national (bold line) and healthboard (faint lines) level.

## Heat Map



**Figure 10.** National heat map showing mean norovirus genogroup II gc/day per 100k for each healthboard in the most recent sampling week.

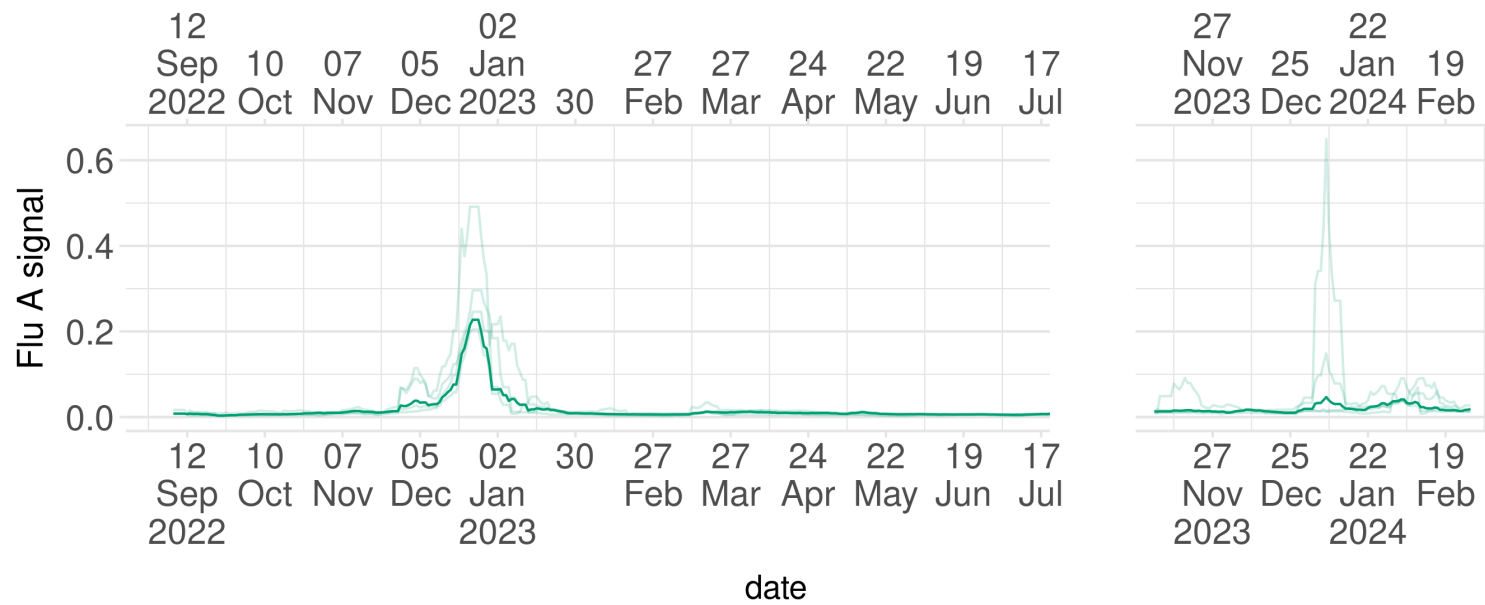
Healthboard Summaries



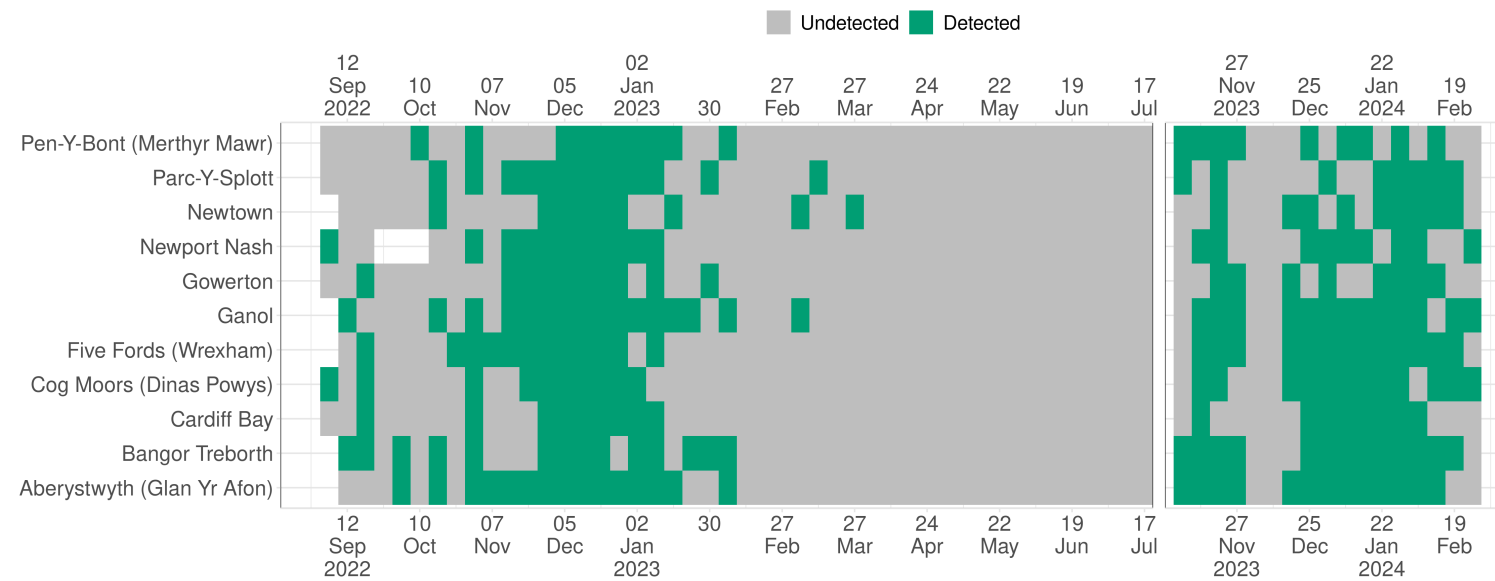
**Figure 11.** Level of norovirus genogroup I (NoVGII) given as a 10 day rolling mean at the healthboard level (bold line) and per day (dots).

# Influenza A (flu-A) virus

## National Summary



**Figure 12.** Level of influenza A (flu-A) virus given as a 10 day rolling mean at the national (bold line) and healthboard (faint lines) level.



**Figure 13.** Weekly aggregated presence and absence of influenza A (flu-A) virus. The limit of detection for the flu-A assay is 303 gene copies per litre of wastewater.

Heat Map

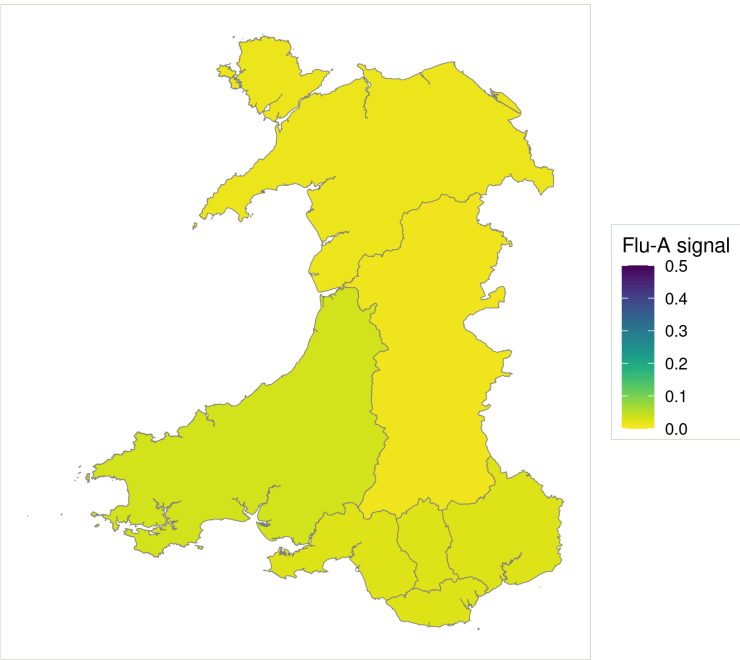


Figure 14. National heat map showing mean flu-A virus gc/day per 100k for each healthboard in the most recent sampling week.

Healthboard Summaries

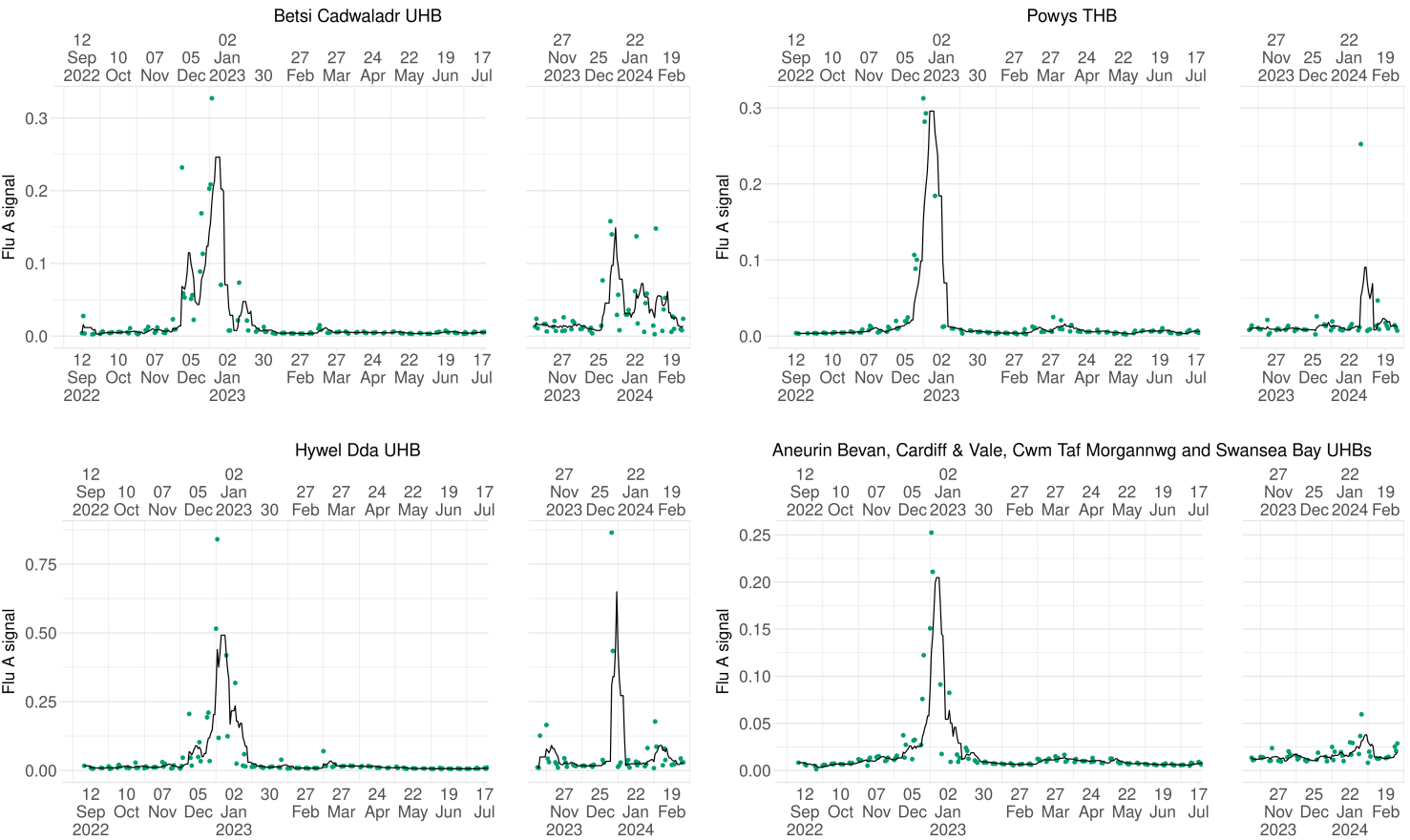
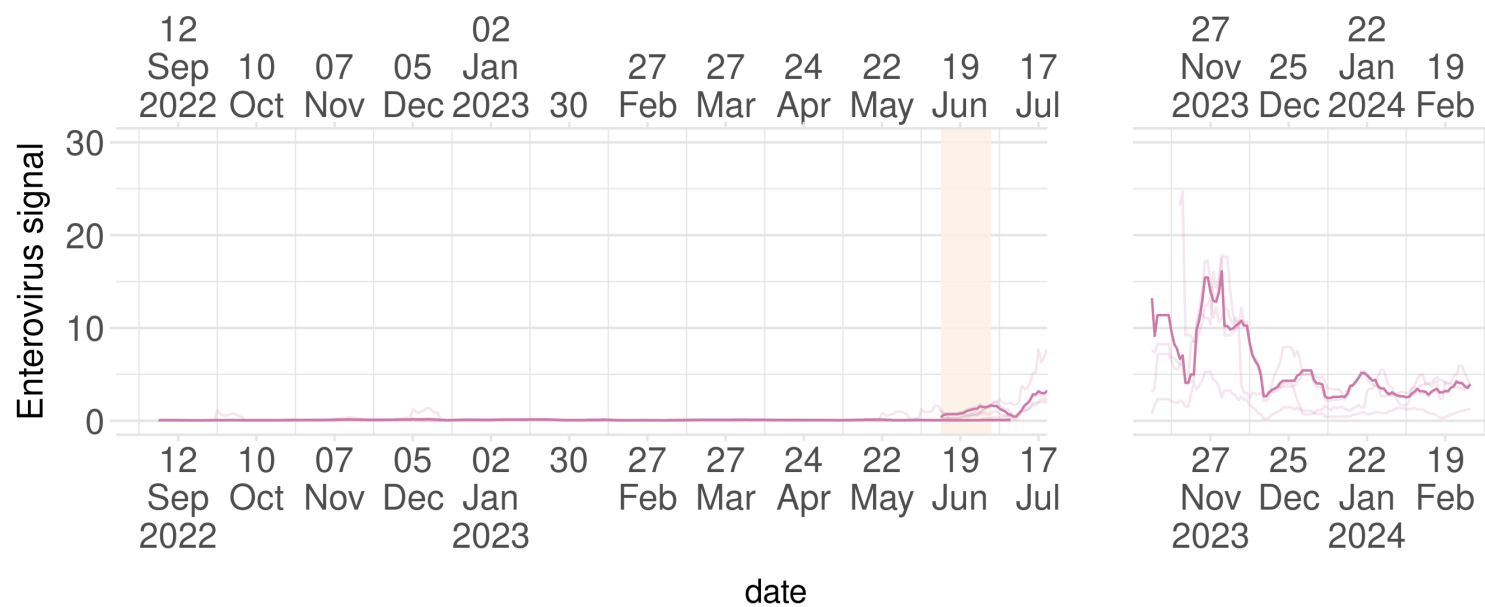


Figure 15. Level of influenza A (flu-A) virus given as a 10 day rolling mean at the healthboard level (bold line) and per day (dots).



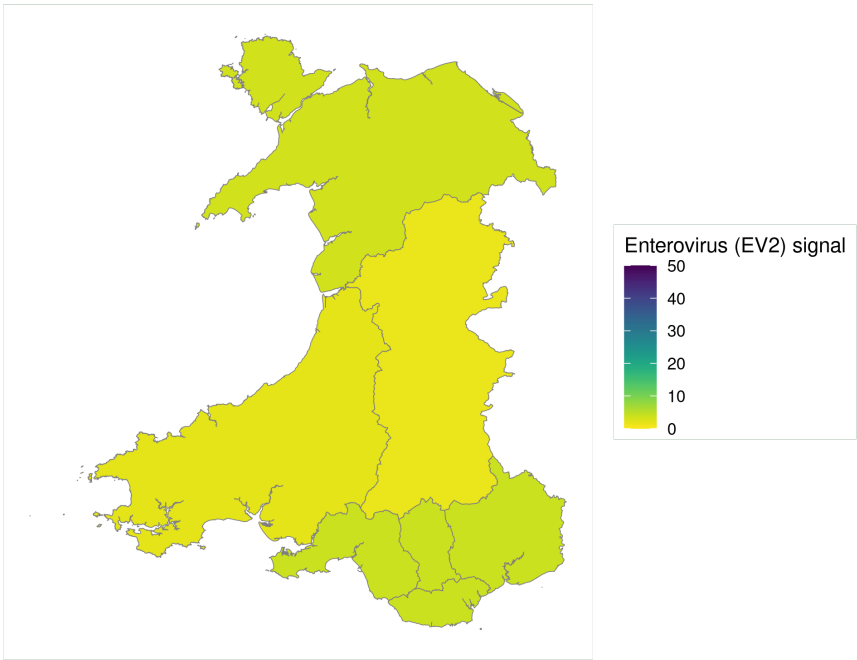
# Enterovirus spp.

## National Summary



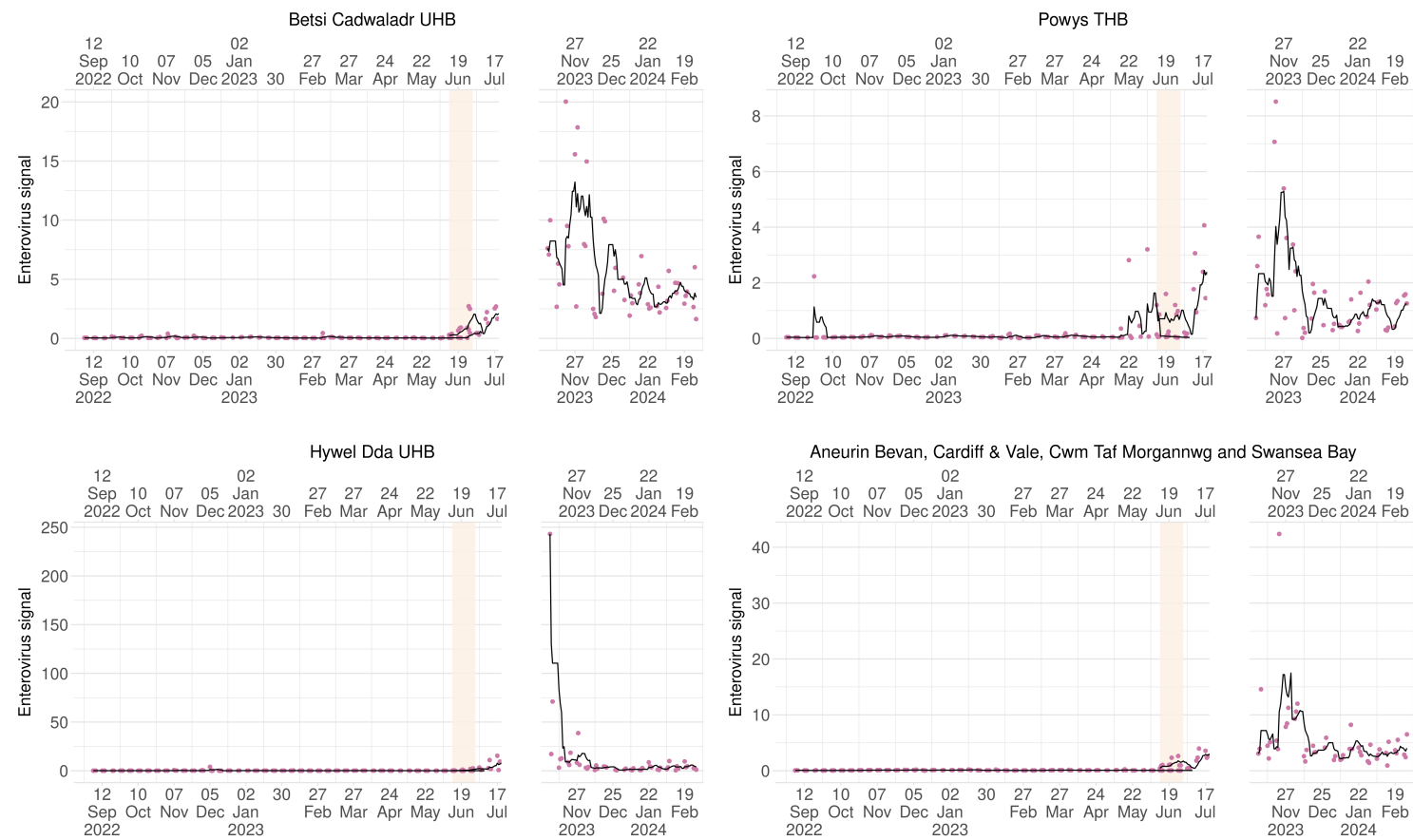
**Figure 16.** Level of Enterovirus spp. (EV1 & EV2) given as a 10 day rolling mean at the national (bold line) and healthboard (faint lines) level. An upper limit has been placed on the y axis, which obscures a high healthboard value, but allows better visualisation of the national trend.

## Heat Map



**Figure 17.** National heat map showing mean Enterovirus spp. gc/day per 100k for each healthboard in the most recent sampling week.

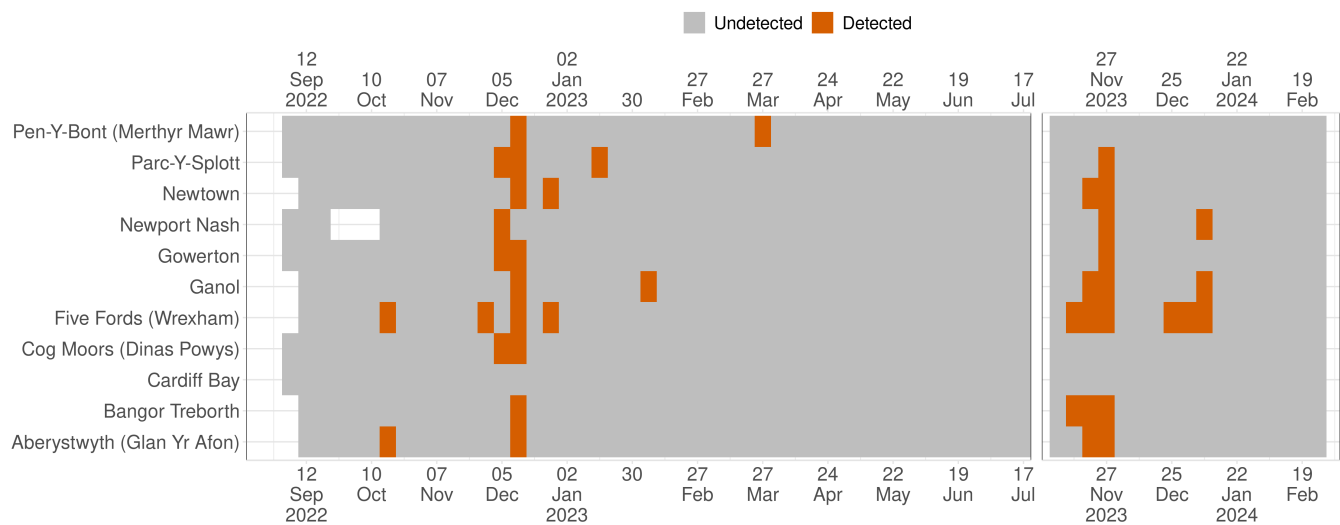
Healthboard Summaries



**Figure 18.** Level of Enterovirus spp. given as a 10 day rolling mean at the healthboard level (bold line) and per day (dots).

# Respiratory Syncytial Virus (RSV)

## National Summary



**Figure 19.** Weekly aggregated presence and absence of Respiratory Syncytial Virus (RSV). The limit of detection for the RSV assay is 703 gene copies per litre of wastewater.

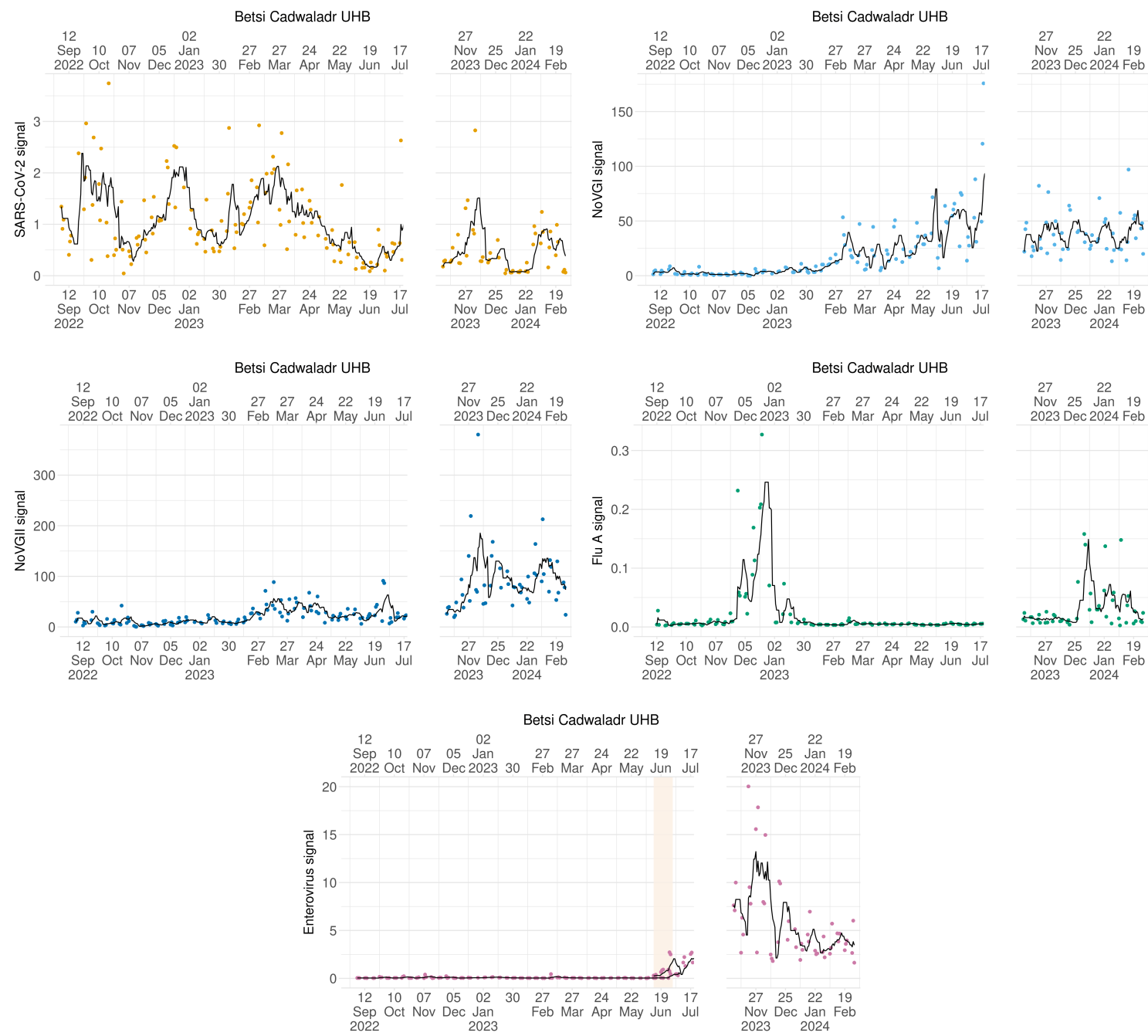
## Heat Map



**Figure 20.** National weekly aggregated presence and absence of Respiratory Syncytial Virus (RSV) for each healthboard in the most recent sampling week.

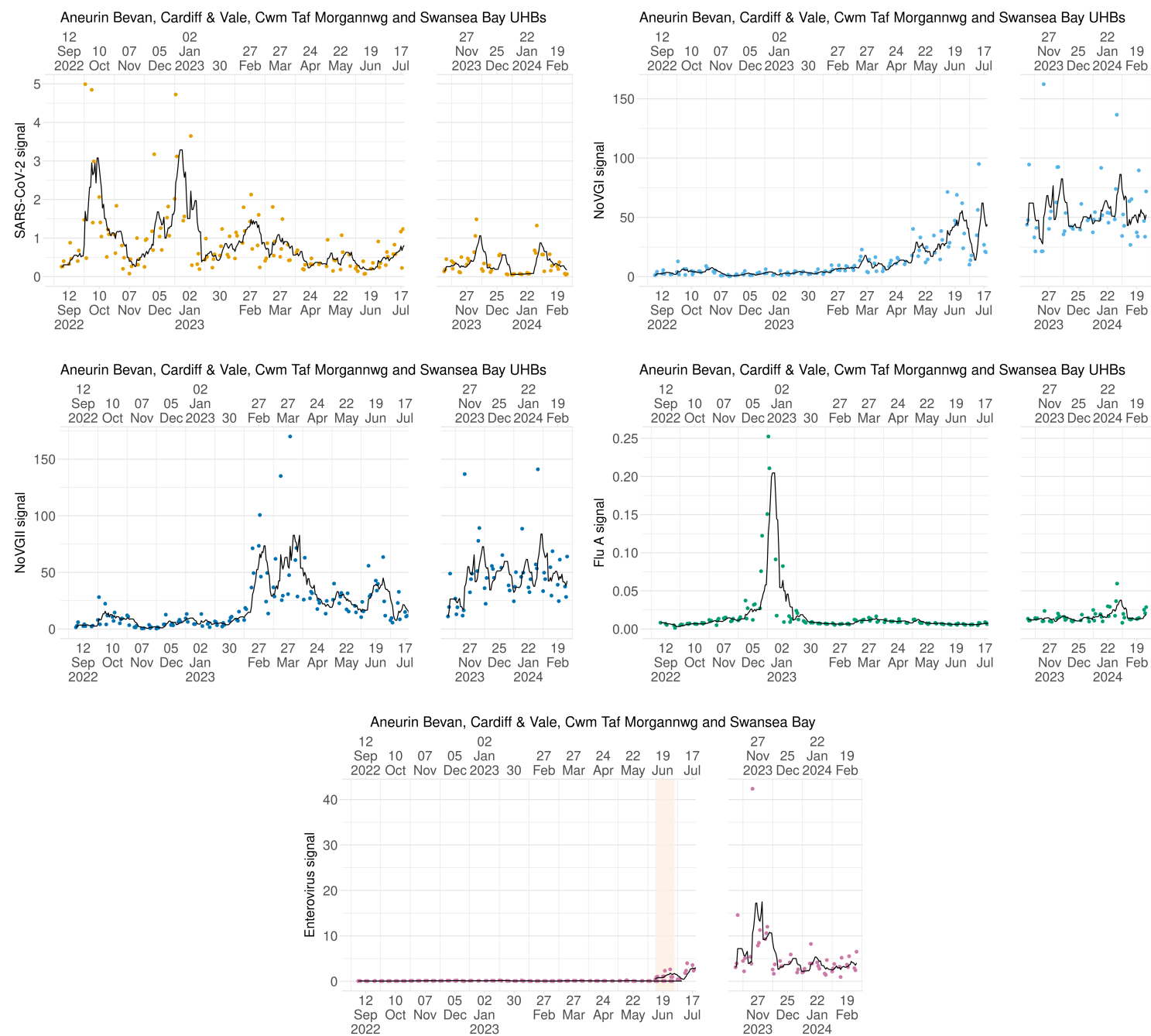
# Health Board Summaries

## Betsi Cadwaladr UHB



**Figure 21.** Viral levels (gc/day per 100k) given as a 10 day rolling mean for Betsi Cadwaladr UHB.

Cardiff & Vale, Aneurin Bevan, Cwm Taf Morgannwg and Swansea Bay UHBs



**Figure 22.** Viral levels (gc/day per 100k) given as a 10 day rolling mean for Cardiff & Vale, Aneurin Bevan, Swansea Bay and Cwm Taf Morgannwg UHBs.

Hywel Dda UHB



Figure 23. Viral levels (gc/day per 100k) given as a 10 day rolling mean for Hywel Dda UHB.



**Figure 24.** Viral levels (gc/day per 100k) given as a 10 day rolling mean for Powys THB.

# Appendices

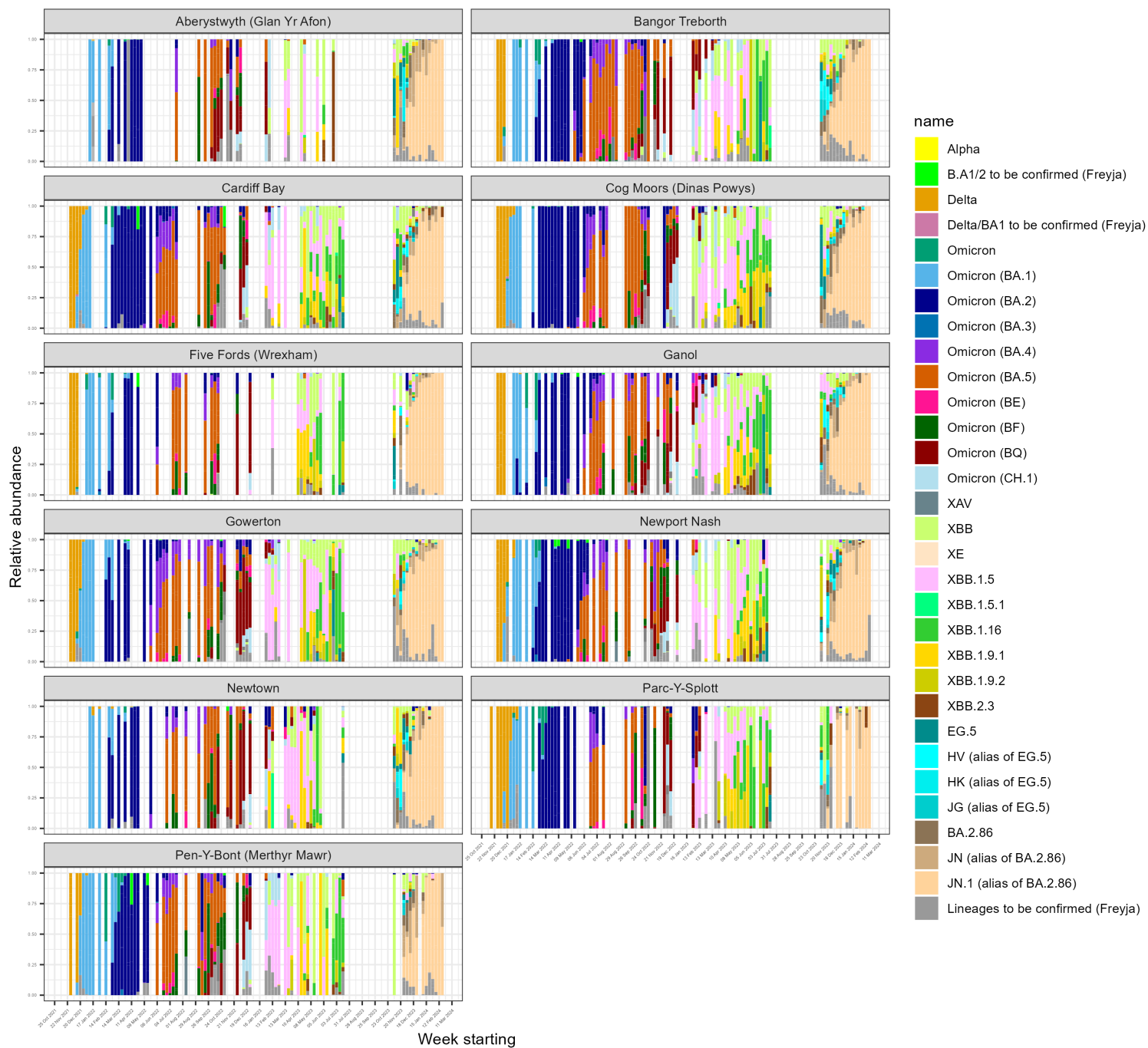
## Appendix 1

Health Board	Sampling Sites	Major City/Town
Aneurin Bevan, Cardiff & Vale, Cwm Taf Morgannwg, Swansea Bay UHBs	Newport Nash	Newport
	Cardiff Bay	Cardiff
	Cog Moors (Dinas Powys)	Barry and Vale of Glamorgan
	Pen-Y-Bont (Merthyr Mawr)	Bridgend
	Gowerton	Swansea
Betsi Cadwaladr UHB	Bangor Treborth	Bangor
	Five Fords (Wrexham)	Wrexham
	Ganol	Llandudno
	Aberystwyth (Glan Yr Afon)	Aberystwyth
Hywel Dda UHB	Parc-Y-Splott	Carmarthen
Powys THB	Newtown	Newtown



## Appendix 2: Site-level summaries

### SARS-CoV-2



**Figure 25.** Proportions of SARS-CoV-2 variants identified at the site level (across 11 WwTPs covered by the Programme).



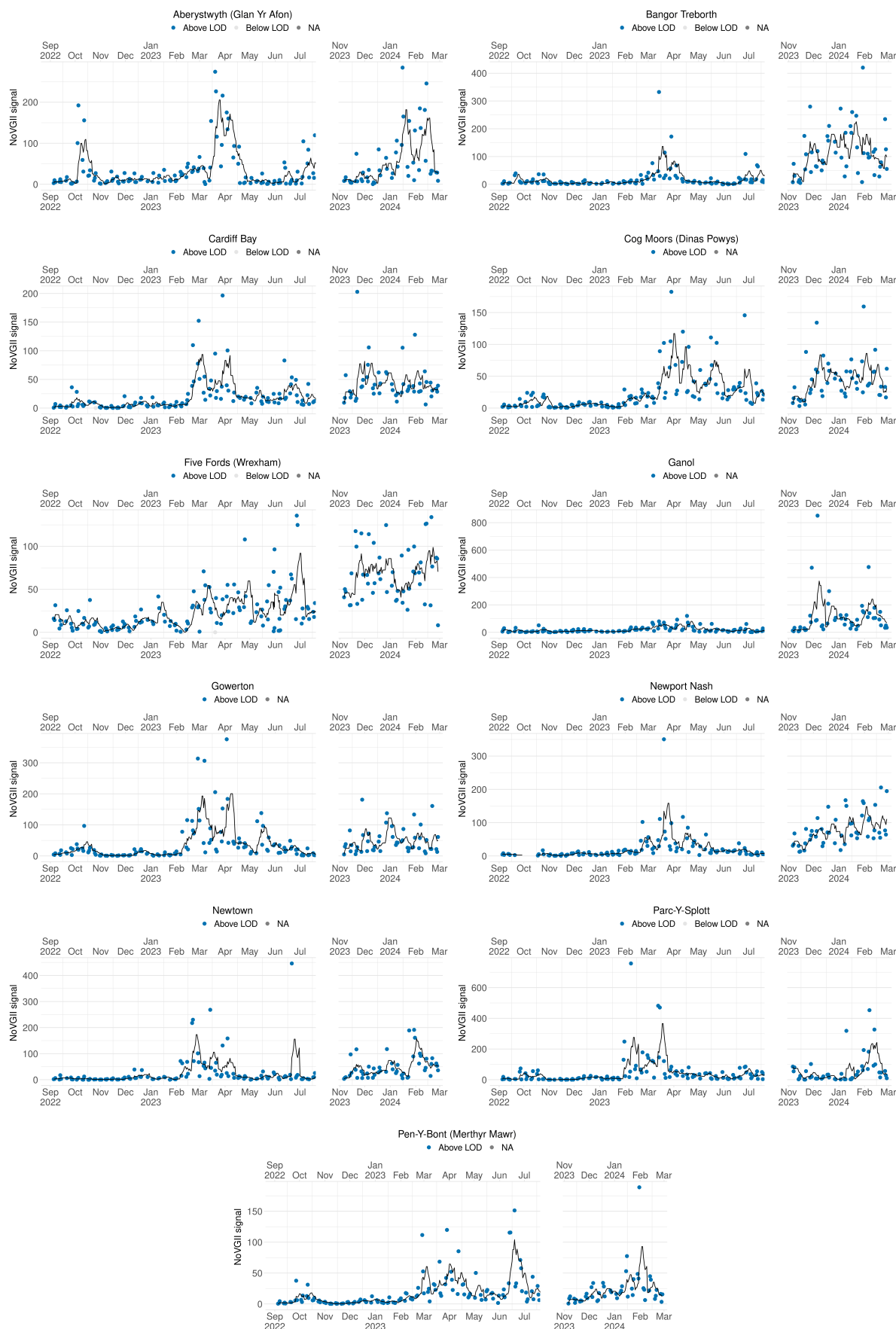
**Figure 26.** Levels of SARS-CoV-2 identified at the site level (across 11 WwTPs covered by the Programme). Each point represents a sample and the line is a 10 day rolling average.

Norovirus genogroup I



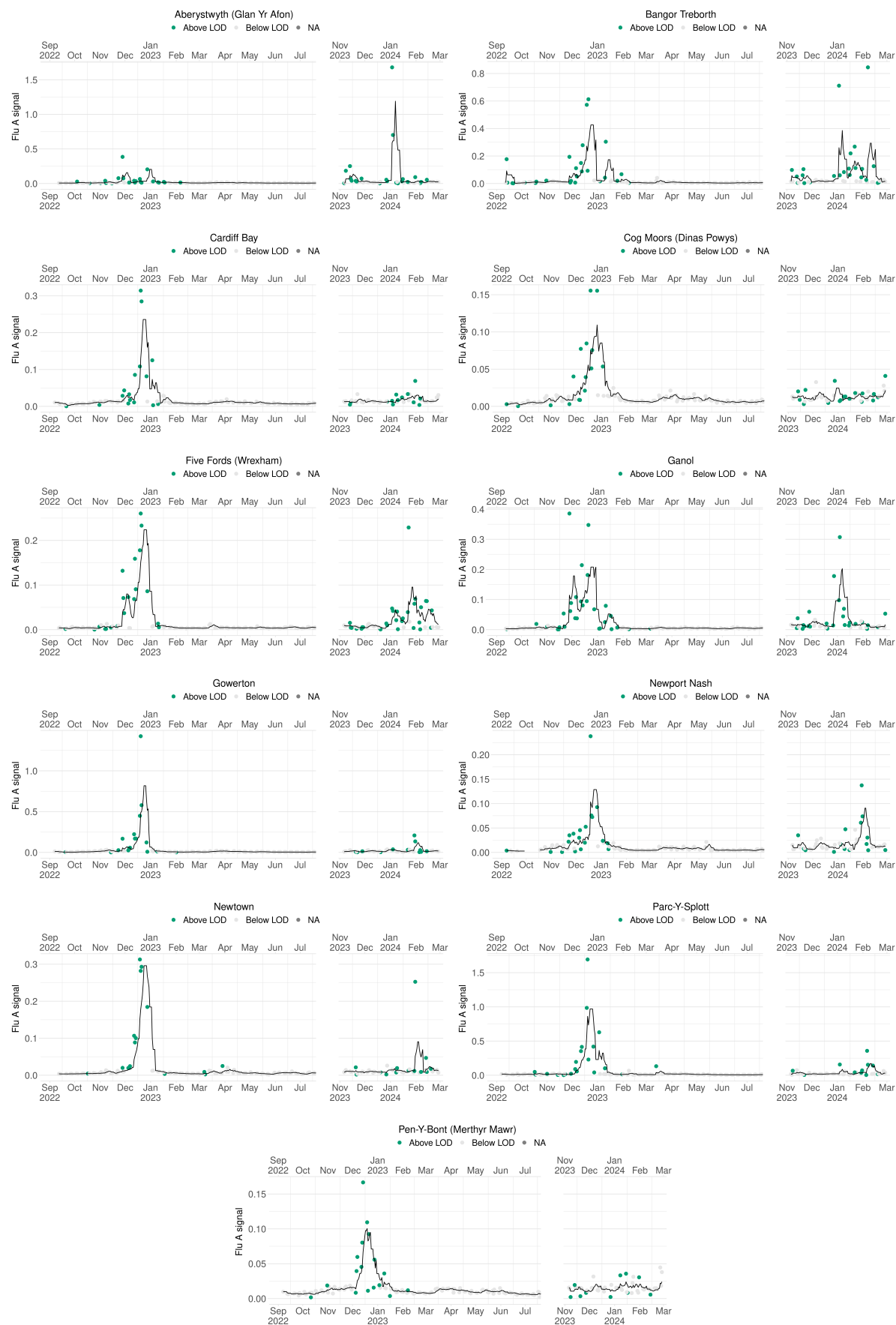
**Figure 27.** Levels of norovirus genogroup I (NoVGI) identified at the site level (across 11 WwTPs covered by the Programme). Each point represents a sample and the line is a 10 day rolling average.

## Norovirus genogroup II



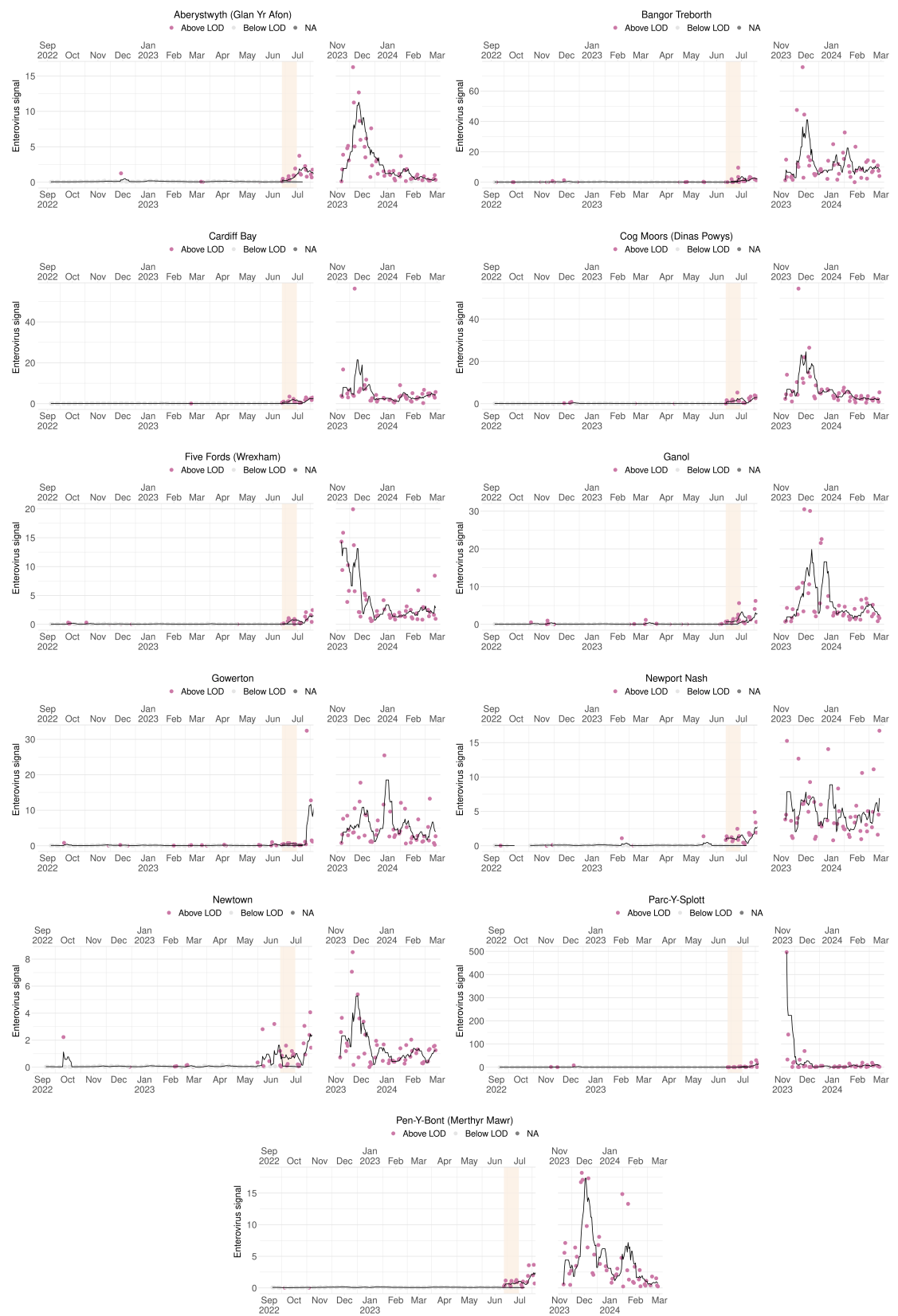
**Figure 28.** Levels of norovirus genogroup II (NoVGII) identified at the site level (across 11 WwTPs covered by the Programme). Each point represents a sample and the line is a 10 day rolling average.

Influenza A (flu-A) virus



**Figure 29.** Levels of influenza A (flu-A) virus identified at the site level (across 11 WwTPs covered by the Programme). Each point represents a sample and the line is a 10 day rolling average.

Enterovirus spp.



**Figure 30.** Levels of enterovirus (EV1 & EV2) identified at the site level (across 11 WwTPs covered by the Programme). Each point represents a sample and the line is a 10 day rolling average.