

HPSC - Weekly Outbreak Report



Statutory Notifications of Outbreaks* reported in Ireland via the Computerised Infectious Disease Reporting (CIDR) system for:

Week 18, 2025

Notification Period: 27/04/2025 - 03/05/2025

Data are provisional

Report produced by the Health Protection Surveillance Centre on 13/05/2025

*COVID-19 outbreaks are not included here but published at:

<https://respiratoryvirus.hpsc.ie/pages/covid-19>



Table 1: Number of Outbreaks* and Total Number Ill by HSE Region in Week 18, 2025

| HSE Region | Number of Outbreaks | Sum of Total number ill |
|--------------|---------------------|-------------------------|
| E | 6 | 49 |
| HPSC | 0 | 0 |
| M | 1 | 5 |
| MW | 1 | 4 |
| NE | 2 | 8 |
| NW | 1 | 3 |
| S | 2 | 11 |
| SE | 3 | 10 |
| W | 0 | 0 |
| Total | 16 | 90 |

Table 2: Number of Norovirus Outbreaks and Total Number Ill in Week 18, 2025 and for the year to date in 2025 compared with the same period last year

| | Week Ending | 2025 | 2024 |
|--|-------------|-----------|-----------|
| | 03/05/2025 | Wk 1 - 18 | Wk 1 - 18 |
| Number of outbreaks of noroviral infection | 2 | 151 | 101 |
| Total number ill | 13 | 1,837 | 1,769 |



Table 3: Summary Line Listing of Outbreaks* in Week 18, 2025

| Outbreak ID | Disease/Organism | HSE area | Outbreak type | Outbreak transmission mode* | Outbreak location | Total Lab Confirmed | Total Number ill |
|-------------|--|----------|---------------|-----------------------------|----------------------------------|---------------------|------------------|
| 42880 | Acute infectious gastroenteritis (unspecified) | E | General | P-P | Residential institution | | 10 |
| 42888 | Acute infectious gastroenteritis (unspecified) | NE | General | P-P and FB | Residential institution | | 4 |
| 42896 | Acute infectious gastroenteritis (unspecified) | NE | General | P-P | Hospital | | 4 |
| 42876 | Acute infectious gastroenteritis (unspecified) | S | General | P-P | Residential institution | | 7 |
| 42891 | Campylobacter infection | SE | Family | P-P | Private house | 4 | 3 |
| 42889 | Influenza | M | General | P-P and Airborne | Hospital | | 5 |
| 42871 | Influenza | E | General | P-P and Airborne | Nursing home | 1 | 24 |
| 42882 | Influenza | E | General | P-P and Airborne | Community Hosp. / Long-stay unit | | 2 |
| 42872 | Measles | NW | Family | P-P and Airborne | Private house | 2 | 3 |
| 42870 | Noroviral infection | E | General | P-P | Hospital | | 9 |
| 42879 | Noroviral infection | S | General | P-P | Hospital | 4 | 4 |
| 42892 | Scabies | MW | Family | P-P | Private house | | 4 |
| 42883 | Scabies Clinical Diagnosis | SE | General | P-P | Nursing home | | 5 |
| 42895 | Verotoxigenic Escherichia coli infection | E | Family | P-P | Private house | 1 | 2 |



| Outbreak ID | Disease/Organism | HSE area | Outbreak type | Outbreak transmission mode* | Outbreak location | Total Lab Confirmed | Total Number ill |
|-------------|--|----------|---------------|-----------------------------|-------------------|---------------------|------------------|
| 42894 | Verotoxigenic Escherichia coli infection | E | Family | Animal contact | Private house | 2 | 2 |
| 42881 | Verotoxigenic Escherichia coli infection | SE | Family | P-P | Private house | 2 | 2 |



Table 4: Summary Line Listing of Outbreaks* from Week 15, 2025 to Week 17, 2025

Week 15

| Outbreak ID | Disease/Organism | HSE Region | Type | Transmission mode* | Location | Total Lab Confirmed | Total Number ill |
|-------------|--|------------|---------|--------------------|-------------------------|---------------------|------------------|
| 42746 | Fusarium proliferatum complex | E | General | Unknown | Hospital | | 4 |
| 42758 | Noroviral infection | NE | General | P-P | Hospital | | 5 |
| 42783 | Noroviral infection | NE | General | P-P | Hospital | 2 | 5 |
| 42787 | Acute infectious gastroenteritis (unspecified) | NE | General | P-P and FB | Nursing home | | 13 |
| 42743 | Varicella suspected | M | General | P-P and Airborne | Other | | 6 |
| 42745 | Respiratory syncytial virus infection | E | General | P-P and Airborne | Hospital | | 2 |
| 42780 | Influenza | E | General | P-P and Airborne | Nursing home | | 38 |
| 42751 | Clostridium difficile infection | E | General | P-P | Hospital | 2 | 4 |
| 42762 | Noroviral infection | E | General | P-P | Hospital | 3 | 4 |
| 42763 | Varicella | E | General | P-P and Airborne | Residential institution | 1 | 4 |
| 42774 | Influenza | E | General | P-P and Airborne | Residential institution | | 3 |
| 42775 | Clostridium difficile infection | M | General | P-P and Airborne | Hospital | 2 | 14 |
| 42776 | VRE | M | General | P-P | Hospital | | 3 |
| 42747 | Acute infectious gastroenteritis (unspecified) | SE | General | P-P | Nursing home | | 16 |
| 42754 | Acute infectious gastroenteritis (unspecified) | E | General | P-P | Nursing home | | 3 |
| 42748 | Influenza | E | General | P-P and Airborne | Nursing home | 8 | 13 |
| 42755 | Noroviral infection | E | General | P-P | Nursing home | 3 | 7 |
| 42766 | Scabies Clinical Diagnosis | SE | General | P-P | Nursing home | | 10 |
| 42779 | Clostridium difficile infection | E | General | P-P | Hospital | 2 | 2 |
| 42781 | Acute infectious gastroenteritis (unspecified) | E | General | P-P | Residential institution | | 10 |
| 42784 | Influenza | SE | General | P-P and Airborne | Nursing home | 1 | 4 |
| 42785 | Noroviral infection | SE | General | P-P | Hospital | 1 | 3 |
| 42757 | Influenza | NW | General | P-P and Airborne | Hospital | 18 | 18 |
| 42768 | Noroviral infection | NW | General | P-P | Hospital | 5 | 7 |
| 42744 | Pertussis | S | Family | Not Specified | Private house | 1 | 3 |



| Outbreak ID | Disease/Organism | HSE Region | Type | Transmission mode* | Location | Total Lab Confirmed | Total Number ill |
|-------------|--|------------|---------|--------------------|-------------------------|---------------------|------------------|
| 42756 | Acute infectious gastroenteritis (unspecified) | S | General | P-P | Residential institution | | 5 |
| 42764 | Pertussis | S | Family | P-P | Private house | 1 | 2 |
| 42769 | scabies- clinical dx. by GP | S | Family | P-P | Private house | | 4 |
| 42761 | Verotoxigenic Escherichia coli infection | W | Family | P-P | Private house | 2 | 3 |
| 42772 | Noroviral infection | W | General | P-P | Hospital | 1 | 6 |
| 42777 | Scabies | W | General | P-P | Nursing home | | 16 |
| 42778 | Pertussis | W | Family | P-P and Airborne | Private house | 1 | 5 |

Week 16

| Outbreak ID | Disease/Organism | HSE Region | Type | Transmission mode* | Location | Total Lab Confirmed | Total Number ill |
|-------------|--|------------|---------|--------------------|--------------------------|---------------------|------------------|
| 42814 | Hepatitis A (acute) | E | General | P-P | Private house | 3 | 3 |
| 42797 | Influenza | E | General | P-P | Hospital | 8 | 9 |
| 42799 | Noroviral infection | E | General | P-P | Nursing home | | 15 |
| 42807 | Noroviral infection | E | General | P-P | Nursing home | 3 | 12 |
| 42808 | Acute respiratory infection | NE | General | P-P | Nursing home | | 9 |
| 42790 | Noroviral infection | NE | General | P-P | Hospital | | 13 |
| 42809 | Acute infectious gastroenteritis (unspecified) | M | General | P-P | Residential institution | | 3 |
| 42821 | Noroviral infection | E | General | P-P | Hospital | 1 | 19 |
| 42822 | Noroviral infection | E | General | P-P | Hospital | 1 | 5 |
| 42791 | Respiratory syncytial virus infection | E | General | P-P and Airborne | Other healthcare service | | 3 |
| 42789 | Noroviral infection | SE | General | P-P | Nursing home | 1 | 11 |
| 42792 | Salmonellosis | SE | Family | P-P | Private house | 2 | 2 |
| 42815 | Cryptosporidiosis | SE | General | P-P | Hotel | 3 | 3 |
| 42798 | Noroviral infection | SE | General | P-P | Hospital | | 4 |
| 42802 | Carbapenemase-producing Enterobacteriaceae (CPE) infection or colonisation | E | General | P-P | Hospital | | 0 |
| 42816 | Noroviral infection | SE | General | P-P | Hotel | 2 | 5 |
| 42812 | Acute respiratory infection | E | General | P-P and Airborne | Nursing home | | 6 |



| Outbreak ID | Disease/Organism | HSE Region | Type | Transmission mode* | Location | Total Lab Confirmed | Total Number ill |
|-------------|--|------------|---------|--------------------|---------------|---------------------|------------------|
| 42804 | Verotoxigenic Escherichia coli infection | MW | Family | P-P | Private house | 2 | 4 |
| 42795 | Cryptosporidiosis | NW | Family | P-P | Private house | 2 | 2 |
| 42810 | Rotavirus infection | NW | General | P-P and Airborne | Hospital | 4 | 4 |
| 42805 | Cryptosporidiosis | S | Family | P-P | Private house | 2 | 2 |
| 42817 | Respiratory syncytial virus infection | S | General | P-P | Nursing home | 6 | 7 |
| 42788 | Noroviral infection | W | General | P-P | Nursing home | 2 | 6 |
| 42793 | Acute infectious gastroenteritis (unspecified) | W | General | P-P | Nursing home | 1 | 9 |
| 42801 | Noroviral infection | W | General | P-P | Nursing home | 3 | 17 |

Week 17

| Outbreak ID | Disease/Organism | HSE Region | Type | Transmission mode* | Location | Total Lab Confirmed | Total Number ill |
|-------------|--|------------|---------|--------------------|--------------------------|---------------------|------------------|
| 42829 | Acute respiratory infection | E | General | P-P | Nursing home | | 13 |
| 42852 | Tuberculosis | E | Family | P-P | Private house | 2 | 3 |
| 42865 | Acute infectious gastroenteritis (unspecified) | NE | General | P-P and FB | Other healthcare service | | 2 |
| 42824 | Influenza | E | General | P-P and Airborne | Hospital | | 4 |
| 42843 | Respiratory syncytial virus infection | M | General | P-P and Airborne | Nursing home | | 10 |
| 42850 | Influenza | M | General | P-P and Airborne | Hospital | | 11 |
| 42853 | Tuberculosis | M | Family | P-P and Airborne | Private house | 2 | 2 |
| 42854 | Influenza | E | General | P-P and Airborne | Hospital | | 9 |
| 42859 | Mumps | M | Family | P-P | Private house | | 4 |
| 42864 | Acute infectious gastroenteritis (unspecified) | E | General | P-P | Nursing home | | 4 |
| 42844 | Cryptosporidiosis | M | Family | Not Specified | Private house | 2 | 2 |
| 42831 | Noroviral infection | SE | General | P-P | Residential institution | 2 | 8 |
| 42827 | vancomycin-resistant enterococci | SE | General | P-P | Hospital | | 0 |
| 42828 | vancomycin-resistant enterococci | SE | General | P-P | Hospital | | 0 |
| 42838 | Noroviral infection | E | General | P-P | Nursing home | 1 | 19 |
| 42845 | Verotoxigenic Escherichia coli infection | SE | Family | P-P | Private house | 1 | 2 |



| Outbreak ID | Disease/Organism | HSE Region | Type | Transmission mode* | Location | Total Lab Confirmed | Total Number ill |
|-------------|--|------------|---------|--------------------|---------------------------------|---------------------|------------------|
| 42856 | Influenza | SE | General | P-P and Airborne | Residential institution | | 3 |
| 42867 | Varicella clinical Diagnosis | E | General | P-P | Residential institution | | 2 |
| 42832 | Noroviral infection | NW | General | P-P and Airborne | Hospital | 4 | 8 |
| 42846 | Cryptosporidiosis | NW | Family | P-P | Private house | 2 | 2 |
| 42847 | Influenza | NW | General | P-P and Airborne | Hospital | 3 | 3 |
| 42861 | Acute respiratory infection | NW | General | P-P and Airborne | Nursing home | | 11 |
| 42866 | Influenza | NW | General | P-P and Airborne | Nursing home | 4 | 5 |
| 42836 | Influenza | S | General | P-P | Nursing home | 2 | 6 |
| 42840 | Outbreak | W | Family | P-P | Private house | 2 | 2 |
| 42851 | Verotoxigenic Escherichia coli infection | W | Family | P-P | Private house | 2 | 2 |
| 42858 | Acute respiratory infection | W | General | P-P | Community Hosp. /Long-stay unit | | 3 |
| 42862 | blaCTX-M-15 gene (ESBL) Detected | W | General | P-P | Hospital | | 6 |
| 42868 | Cryptosporidiosis | W | Family | P-P | Private house | 2 | 2 |

Guidance notes:

1. An outbreak of infection or foodborne illness may be defined as two or more linked cases of the same illness, or the situation where the observed number of cases exceeds the expected number, or a single case of disease caused by a significant pathogen (e.g. diphtheria or viral haemorrhagic fever). Outbreaks may be confined to some of the members of one family or may be more widespread and involve cases either locally, nationally or internationally.
2. Data for this report were extracted from Computerised Infectious Disease Reporting (CIDR) on 13 May 2025, and were correct at the time that data were extracted. Information from previous weeks is updated on an ongoing basis in CIDR, therefore data in this report may be updated in future reports.
3. Data are collated and analysed using the CIDR system. Data are provisional in this report and are subject to change until final validation at the end of an outbreak.
4. The counties covered by each HSE area are as follows:
 - HSE East (E): Dublin, Kildare & Wicklow;
 - HSE Midlands (M): Laois, Longford, Offaly & Westmeath;
 - HSE Midwest (MW): Clare, Limerick & N. Tipperary;
 - HSE Northeast (NE): Cavan, Louth, Meath & Monaghan;
 - HSE Northwest (NW): Donegal, Leitrim & Sligo;
 - HSE South (S): Kerry & Cork;
 - HSE Southeast (SE): Carlow, Kilkenny, S. Tipperary, Waterford & Wexford;
 - HSE West (W): Galway, Mayo & Roscommon.
5. The following abbreviations are used when describing mode of transmission:
 - P-P : Person-to-Person transmission;
 - FB : Foodborne transmission;
 - WB : Waterborne transmission.
6. Total number ill does not include asymptomatic cases
7. COVID-19 outbreaks are not included here but published at:
[Respiratory Virus Notification Data Hub \(hpsc.ie\)](https://hpsc.ie)