SURVEILLANCE OF INFECTIOUS INTESTINAL (IID), ZOONOTIC AND **VECTORBORNE DISEASE, AND OUTBREAKS of INFECTIOUS DISEASE IN IRELAND**







A quarterly report by the Health Protection Surveillance Centre in collaboration with the Departments of Public Health

Quarter 3–2018

December 2018

This is the third quarterly report for 2018 produced by the Gastroenteric Unit of the Health Protection Surveillance Centre.

The production of this quarterly report would not be possible without the valuable input and commitment from the Directors of Public Health, Specialists in Public Health Medicine, Surveillance Scientists, Clinical Microbiologists, General Practitioners, Hospital Clinicians, Infection Control, Environmental Health and laboratory personnel, and other professionals who provide the data for the HPSC's surveillance systems.

Note: Data are collected and analysed using the Computerised Infectious Disease Reporting (CIDR) system. The data in this report are provisional and will not be regarded as final until all returns are received and data have been validated.

OUTBREAK SURVEILLANCE

| | Table 1. General outbreaks of infectious intestinal disease (IID) in Q3, 2018 | | | | | | | | | | |
|-------|---|---------------------------|-----------|--------------|------------|---------------------------------|----------------------------|--|--|--|--|
| Month | HSE area | Location | No. ill * | No. Hosp. | Date Onset | Suspect mode of transmission | Disease | | | | |
| Jul | NW | Nursing home | 4 | 0 | 02/07/2018 | P-P | AIG | | | | |
| Jul | HPSC | Community outbreak | 86 | 44 | 04/06/2018 | Unknown | VTEC | | | | |
| Jul | S | Childcare facility | | | 23/06/2018 | P-P | VTEC | | | | |
| Jul | W | Childcare facility | 20 | 4 | 20/06/2018 | Unknown | VTEC | | | | |
| Jul | SE | Residential institution | 2 | 0 | 06/07/2018 | P-P & AB | AIG | | | | |
| Jul | S | Childcare facility | | | 10/07/2018 | P-P | Noroviral infection | | | | |
| Jul | S | Childcare facility | 7 | 0 | 12/06/2018 | P-P | VTEC | | | | |
| Jul | S | Childcare facility | 5 | | 01/07/2018 | Not Specified | VTEC | | | | |
| Jul | NE | Nursing home | 21 | 2 | 23/07/2018 | P-P | Noroviral infection | | | | |
| Jul | Е | Nursing home | 14 | | 28/07/2018 | P-P | AIG | | | | |
| Aug | Е | Other | 2 | | 01/05/2018 | Unknown | Giardiasis | | | | |
| Aug | NE | Childcare facility | 2 | | 15/07/2018 | Unknown | VTEC | | | | |
| Aug | Е | Nursing home | 11 | | 19/07/2018 | P-P | AIG | | | | |
| Aug | MW | Comm. Hosp/Long-stay unit | 14 | | 29/07/2018 | P-P & AB | Noroviral infection | | | | |
| Aug | NW | Nursing home | 13 | 2 | 04/08/2018 | P-P | AIG | | | | |
| Aug | М | Nursing home | 1 | 0 | 31/07/2018 | Unknown | VTEC | | | | |
| Aug | SE | Comm. Hosp/Long-stay unit | 30 | | 08/08/2018 | P-P | Noroviral infection | | | | |
| Aug | М | Hospital | 3 | | | Unknown | Noroviral infection | | | | |
| Aug | Е | Hospital | 6 | 6 | 20/08/2018 | P-P | Noroviral infection | | | | |
| Aug | М | Hotel | 3 | 0 | | Unknown | Noroviral infection | | | | |
| Aug | S | Nursing home | | | 18/08/2018 | P-P | Noroviral infection | | | | |
| Aug | NW | Residential institution | 3 | | 30/08/2018 | P-P | AIG | | | | |
| Aug | W | Hotel | 2 | 0 | | FB | Campylobacter infection | | | | |
| Sep | W | Nursing home | 22 | 0 | 02/09/2018 | P-P | Noroviral infection | | | | |
| Sep | Е | Nursing home | 37 | 0 | 31/08/2018 | P-P & AB | Rotavirus infection | | | | |
| Sep | М | Comm. Hosp/Long-stay unit | 14 | | | Not Specified | Noroviral infection | | | | |
| Sep | S | Residential institution | 4 | 0 | 04/09/2018 | P-P | AIG | | | | |
| Sep | Е | School | 15 | | 12/09/2018 | P-P | AIG | | | | |
| Sep | NW | Nursing home | 6 | 0 | 06/09/2018 | P-P | Noroviral infection | | | | |
| Sep | NE | Residential institution | 4 | 0 | 09/09/2018 | P-P | AIG | | | | |
| Sep | S | Residential institution | 7 | 0 | 16/09/2018 | P-P | AIG | | | | |
| Sep | E | Comm. Hosp/Long-stay unit | 2 | | 24/09/2018 | P-P | Noroviral infection | | | | |
| Sep | М | Comm. Hosp/Long-stay unit | 4 | | 18/09/2018 | Unknown | Noroviral infection | | | | |
| Sep | NE | Residential institution | 2 | | | P-P | AIG | | | | |
| Sep | Е | Nursing home | 11 | | 21/09/2018 | P-P | Noroviral infection | | | | |

P-P denotes Person-to-Person transmission, FB denotes foodborne, WB denotes waterborne; AB denotes airborne; AIG denotes Acute Infectious Gastroenteritis (unspecified); VTEC denotes infection with Verotoxigenic *E. coli;* NK=unknown * Total numbers ill does not include asymptomatic cases

| Month | HSE area | Family outbreak Location | No. ill * | No. Hosp. | Date Onset | Suspect mode of | Disease |
|------------|-------------|----------------------------|--------------|--------------|------------|-------------------------|---------------------|
| Jul | М | Private house | 1 | 0 | 11/06/2018 | transmission Unknown | VTEC |
| Jul | М | Private house | | | | Unknown | VTEC |
| Jul | М | Private house | 1 | 1 | | Unknown | VTEC |
| Jul | М | Private house | 1 | 1 | 26/02/2018 | Unknown | VTEC |
| Jul | М | Private house | | | | Unknown | VTEC |
| Jul | М | Private house | 1 | 0 | 02/07/2018 | Unknown | VTEC |
| Jul | NW | Private house | 3 | 1 | 28/06/2018 | P-P | VTEC |
| Jul | М | Private house | 1 | 1 | 04/07/2018 | Unknown | VTEC |
| Jul | S | Private house | 2 | 0 | 12/06/2018 | P-P | VTEC |
| Jul | M | Travel related | 1 | 0 | 01/07/2018 | Unknown | VTEC |
| Jul | M | Private house | 1 | 0 | 06/07/2018 | WB & Animal | VTEC |
| Jul | E | Private house | | Ŭ | 25/06/2018 | P-P & Animal | VTEC |
| Jul | M | Private house | 2 | 0 | 13/07/2018 | Unknown | VTEC |
| Jul | S | Private house | 2 | 0 | 10/01/2010 | P-P | VTEC |
| Jul | W | Private house | 2 | 0 | 04/07/2018 | P-P & FB | VTEC |
| Jul | M | Private house | 1 | 1 | 14/07/2018 | Unknown | VTEC |
| Jul | M | Private house | 1 | 0 | 02/07/2018 | Unknown | VTEC |
| Jul | SE | Other | 4 | 0 | 14/07/2018 | Unknown | AIG |
| Jul | W | Private house | 2 | 0 | 28/06/2018 | Unknown | VTEC |
| Jul | M | Private house | 1 | 0 | 23/07/2018 | Unknown | VTEC |
| Jul | M | Private house | 1 | 0 | 20/07/2018 | Not Specified | VTEC |
| | M | Private house | 2 | 0 | 23/07/2018 | Unknown | VTEC |
| Aug Aug | SE | Private house | 3 | 0 | 23/06/2018 | P-P | Shigellosis |
| - | M | Private house | 1 | 0 | 23/00/2010 | Unknown | VTEC |
| Aug | S | | | | 25/06/2019 | P-P | VTEC |
| Aug | | Private house | 2 | 0 | 25/06/2018 | | |
| Aug | SE | Private house | 2 | 0 | 15/06/2018 | P-P & FB | VTEC |
| Aug | M | Not Specified | 0 | 0 | 44/07/0040 | Not Specified | VTEC |
| Aug | W | Private house | 2 | 0 | 11/07/2018 | Not Specified | Cryptosporidiosis |
| Aug | E | Private house | 3 | 2 | 24/07/2018 | Unknown | VTEC |
| Aug | M | Private house | | | | Unknown | VTEC |
| Aug | М | Private house | 1 | 1 | 24/07/2018 | Unknown | VTEC |
| Aug | NW | Private house | 5 | 3 | 31/07/2018 | P-P | Rotavirus infection |
| Aug | E | Private house | 3 | | | Unknown | Giardiasis |
| Aug | S | Private house | 2 | 1 | 30/07/2018 | P-P | VTEC |
| Aug | М | Private house | 1 | 1 | | Unknown | VTEC |
| Aug | MW | Private house | 1 | 0 | 01/12/2017 | P-P | VTEC |
| Aug | MW | Private house | 1 | 0 | 27/03/2018 | P-P | VTEC |
| Aug | W | Private house | 2 | 0 | 17/08/2018 | Unknown | VTEC |
| Aug | W | Private house | 2 | 0 | 05/08/2018 | Unknown | VTEC |
| Aug | М | Private house | 1 | | 03/08/2018 | Unknown | VTEC |
| Aug | М | Private house | 2 | 0 | 14/08/2018 | Unknown | VTEC |
| Aug | Е | Private house | 1 | | | Unknown | VTEC |
| Aug | М | Travel related | 2 | 0 | 24/07/2018 | WB | Giardiasis |
| Aug | М | Private house | 1 | | | Not Specified | VTEC |
| Sep | М | Not Specified | | | | Unknown | VTEC |
| Sep | М | Private house | 1 | | 30/08/2018 | Unknown | VTEC |
| Sep | NE | Private house | 1 | | 21/07/2018 | P-P | VTEC |

| | | | 1 | - | | | |
|-----|----|---------------|---|---|------------|-------------------|-------------------|
| Sep | E | Private house | 2 | | 22/08/2018 | Unknown | VTEC |
| Sep | Μ | Not Specified | | | 30/08/2018 | Not Specified | VTEC |
| Sep | Μ | Not Specified | | | 18/08/2018 | Unknown | VTEC |
| Sep | Μ | Private house | 1 | 1 | 21/08/2018 | Unknown | VTEC |
| Sep | Μ | Private house | 1 | 0 | 24/08/2018 | Not Specified | VTEC |
| Sep | W | Private house | 2 | | 13/08/2018 | P-P & Animal | VTEC |
| Sep | MW | Private house | 3 | 0 | 15/08/2018 | Unknown | VTEC |
| Sep | S | Private house | 2 | 0 | 18/08/2018 | P-P & FB | VTEC |
| Sep | MW | Private house | | | 31/08/2018 | P-P | VTEC |
| Sep | NW | Private house | 2 | 2 | 25/08/2018 | Unknown | Cryptosporidiosis |
| Sep | Е | Unknown | 3 | | 10/08/2018 | Unknown | Giardiasis |
| Sep | W | Private house | 2 | 1 | 30/08/2018 | P-P | VTEC |
| Sep | Μ | Private house | 1 | | 13/09/2018 | Unknown | VTEC |
| Sep | Μ | Private house | | | 11/09/2018 | Unknown | VTEC |
| Sep | W | Private house | 2 | 0 | 31/07/2018 | P-P | VTEC |
| Sep | W | Private house | 2 | | 15/08/2018 | P-P | VTEC |
| Sep | SE | Private house | 1 | 1 | 12/09/2018 | P-P | VTEC |
| Sep | S | Private house | | | 04/09/2018 | Animal contact | VTEC |

P-P denotes Person-to-Person transmission, FB denotes foodborne, WB denotes waterborne; AB denotes airborne; AIG denotes Acute InfectiousGastroenteritis; VTEC denotes infection with Verotoxigenic *E. coli* NK denotes unknown * Total numbers ill does not include asymptomatic cases

| | Table 3. Non-IID outbreaks in Q3, 2018 | | | | | | | | | | |
|-------|--|------------------|-------------------------------|----------------|--------------|------------|---------------------------------|--------------------------------|--|--|--|
| Month | HSE area | Type of outbreak | Location | No. ill * | No. Hosp. | Date Onset | Suspect mode of transmission | Organism | | | |
| Jul | Е | General | Hospital | 2 colonised | | 20/06/2018 | P-P | CPE | | | |
| Jul | S | Family | Extended family | | | 15/05/2018 | P-P | Pertussis | | | |
| Jul | S | General | Comm. Hosp/Long- stay unit | 12 | | 08/07/2018 | P-P | Acute respiratory infection | | | |
| Jul | NE | General | Hospital | 2 colonised | | | P-P | Enterobacter cloacae oxa-48 | | | |
| Jul | Е | General | Other | 17 | 2 | 05/07/2018 | P-P & AB | Measles | | | |
| Jul | SE | General | Hospital | 2 colonised | | | P-P | CPE | | | |
| Jul | Е | General | Hospital | 2 colonised | | | P-P | CPE | | | |
| Jul | W | General | Childcare facility | 15 | 1 | 04/07/2018 | Unknown | Adenovirus | | | |
| Jul | SE | General | Hospital | 6 colonised | | | Environmental / Fomite | CPE | | | |
| Jul | SE | General | Hospital | 4 | | 16/07/2018 | P-P | Acute respiratory infection | | | |
| Aug | E | General | Hospital | 3 | 3 | | Environmental / Fomite | Aspergillus | | | |
| Aug | E | General | Hospital | 2 colonised | | | P-P | CPE | | | |
| Aug | MW | General | Community outbreak | 6 | | 25/07/2018 | P-P & AB | Mumps | | | |
| Aug | Е | General | Hospital | 3 colonised | 3 | | P-P | CPE | | | |
| Aug | S | General | Hospital | 2 colonised | | | Not Specified | CPE | | | |
| Aug | MW | General | Hospital | 3 | 3 | 23/07/2018 | P-P | MRSA | | | |
| Aug | SE | General | Workplace | 3 | 1 | 31/07/2018 | P-P | Measles | | | |
| Aug | S | General | Unknown | 7 | 7 | | P-P | Viral meningitis | | | |
| Sep | NE | Family | Private house | 2 | 2 | | P-P | Influenza | | | |
| Sep | NE | General | Hospital | 2 colonised | | | P-P | CPE | | | |
| Sep | W | General | Nursing home | 7 | | | P-P | Clostridium difficile | | | |
| | Page 4 of 14 | | | | | | | | | | |

| Month | HSE area | Type of outbreak | Location | No. ill * | No. Hosp. | Date Onset | Suspect mode of transmission | Organism |
|-------|-------------|---------------------|-------------------------------|----------------|--------------|------------|------------------------------|------------------------------------|
| | | | | | | | | infection |
| Sep | Е | General | Hospital | 4 | 4 | 05/09/2018 | P-P | Clostridium difficile infection |
| Sep | Е | General | Hospital | 2 | 2 | 15/09/2018 | P-P | Clostridium difficile infection |
| Sep | E | General | Comm. Hosp/Long- stay unit | 2 colonised | | 06/09/2018 | P-P | CPE |

P-P denotes Person-to-Person transmission, WB denotes waterborne; AB denotes airborne; NK denotes unknown; CPE denotes Carbapenemresistant Enterobacteriaceae; RSV denotes Respiratory syncytial virus; Pts denotes Patients.

* Total numbers ill does not include asymptomatic cases.

Since July 2001, outbreaks have been reported to HPSC. Preliminary information is provided by a public health professional when the outbreak is first notified. Further information is provided by the lead investigator once more complete data are available.

The data requested includes information on the source of reporting of the outbreak, the extent of the outbreak, mode of transmission, location, pathogen involved, laboratory investigation, morbidity and mortality data, suspect vehicle and factors contributing to the outbreak. The data provided are crucial in providing information on the reasons why the outbreak occurred, the factors that lead to the spread of disease and the lessons that can be learnt to prevent further such outbreaks.

Since the 1st January 2004, with the amendment to the Infectious Diseases Regulations (2003), there is a statutory requirement for medical practitioners and clinical directors of a diagnostic laboratory to notify to the medical officer of health 'any unusual clusters or changing patterns of any illness, and individual cases thereof, that may be of public health concern'.

Tables 1 and 2 present a line listing of all general and family outbreaks of IID reported to HPSC in the third quarter of 2018. There were 35 general and 65 family IID outbreaks reported during this period, resulting in at least 456 people being ill.

Norovirus (n=14) and Acute infectious gastroenteritis (n=11) were responsible for the most general outbreaks of IID (71%).

Twenty-four general IID outbreaks were transmitted person-to-person/person-to-person & airborne (68%). Twenty-five general IID outbreaks (71%) were reported to have occurred in healthcare settings, i.e. hospitals or residential institutions, during this period.

The most common cause of family outbreaks of IID was VTEC (n=57) [88%]. Other pathogens responsible for family outbreaks in Q3 2018 were AIG, cryptosporidiosis, giardiasis, rotavirus and shigellosis (Table 2).

There were twenty-four non-IID outbreaks reported during Q3 2018 (Table 3). The most common cause of non IID outbreaks was during this period was CPE (Carbapenemase Producing Enterobacteriaceae) (n=9) [38%]. All CPE colonisations reported in Q3 2018 occurred in healthcare settings.

Table 4 outlines the outbreak rate per HSEarea for outbreaks notified during Q3 2018.

Table4.Numberofinfectiousdiseaseoutbreaks by HSE Area, Q3 2018

| HSE Area | No. of outbreaks | Rate per 100,000 population |
|----------|---------------------|--------------------------------|
| E | 23 | 1.3 |
| Μ | 37 | 13.0 |
| MW | 7 | 2.0 |
| NE | 8 | 2.0 |
| NW | 7 | 3.0 |
| SE | 10 | 2.0 |
| S | 17 | 2.4 |
| W | 14 | 3.0 |
| Total | 123 | 2.6 |

NOTIFICATIONS OF INFECTIOUS INTESTINAL, ZOONOTIC AND VECTORBORNE DISEASE

The number of notifications of infectious intestinal, zoonotic and vectorborne disease by HSE-Area for the third quarter of 2018 is shown in Table 5.

| Table 5. Infectious intestinal, zoonotic and | d vecto | rborne | disease | notific | ations | Q3, 201 | 8 by HS | SE-Area | 3 |
|---|---------|--------|---------|---------|--------|---------|---------|---------|-------|
| Infectious Intestinal Disease | Е | М | MW | NE | NW | SE | S | W | Total |
| Bacillus cereus foodborne infection/intoxication | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Botulism | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Campylobacter infection ¹ | 308 | 49 | 59 | 77 | 33 | 93 | 127 | 74 | 820 |
| Cholera | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Clostridium perfringens (type A) food-borne disease | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Cryptosporidiosis | 18 | 13 | 5 | 4 | 4 | 9 | 12 | 16 | 81 |
| Giardiasis | 25 | 4 | 4 | 0 | 1 | 13 | 14 | 6 | 67 |
| Listeriosis | 4 | 1 | 0 | 0 | 0 | 3 | 0 | 2 | 10 |
| Noroviral infection ² | 115 | 16 | 17 | 22 | 3 | 2 | 34 | 13 | 222 |
| Paratyphoid | 2 | ~ | ~ | ~ | ~ | ~ | ~ | ~ | 2 |
| Rotavirus infection ³ | 56 | 12 | 8 | 19 | 7 | 17 | 39 | 14 | 172 |
| Salmonellosis | 49 | 12 | 11 | 29 | 7 | 12 | 21 | 8 | 149 |
| Shigellosis | 19 | 0 | 7 | 2 | 1 | 5 | 3 | 0 | 37 |
| Staphylococcal food poisoning | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Typhoid | 1 | ~ | ~ | ~ | ~ | ~ | ~ | ~ | 4 |
| Verotoxigenic Escherichia coli infection | 87 | 59 | 47 | 46 | 13 | 65 | 93 | 90 | 500 |
| Yersiniosis | 2 | 0 | 0 | 1 | 0 | 0 | 0 | 0 | 3 |
| Zoonotic Disease | | | | | | | | | |
| Anthrax | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Brucellosis | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Echinococcosis | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 0 | 1 |
| Leptospirosis | 1 | 0 | 0 | 0 | 0 | 1 | 1 | 0 | 3 |
| Plague | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Q Fever | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Rabies | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Toxoplasmosis | 3 | 0 | 1 | 0 | 0 | 1 | 0 | 1 | 6 |
| Trichinosis | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Vectorborne Disease | | | | | | | | | |
| Chikungunya disease | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Dengue | 4 | 0 | 1 | 0 | 0 | 0 | 0 | 0 | 5 |
| Lyme disease (neuroborreliosis) | 0 | 0 | 0 | 0 | 0 | 2 | 2 | 0 | 4 |
| Malaria | 7 | 1 | 0 | 1 | 0 | 4 | 2 | 1 | 16 |
| Typhus | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| West Nile fever | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Zika Virus Infection | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |

¹ From August 2017, campylobacter notifications from HSE-East re based on laboratory testing results rather than patient episodes. Notifications from HSE-E may also refer to area of laboratory testing rather than area of patient residence.

² Between March 2013 and July 2017, norovirus notifications from HSE-East were based on laboratory testing results rather than patient episodes. Notifications from HSE-E may also refer to area of laboratory testing rather than area of patient residence.

³ Between March 2013 and July 2017, rotavirus notifications from HSE-East were based on laboratory testing results rather than patient episodes. Notifications from HSE-E may also refer to area of laboratory testing rather than area of patient residence.

SALMONELLA ENTERICA

Human salmonellosis (*S. enterica*) is a notifiable disease. The National *Salmonella*, *Shigella* and *Listeria* Reference Laboratory (NSSLRL) in Ireland was established in 2000 in the Dept. of Medical Microbiology, University College Hospital, Galway. This laboratory accepts *S. enterica* isolates from all clinical and food laboratories in Ireland for serotyping, phage typing and antimicrobial sensitivity testing. Table 6 shows the number of salmonellosis notifications by HSE-Area and month for the third quarter of 2018. Comparison of trends with previous years is shown in Figure 1.

Table 6. Salmonellosis notifications by HSE-Area and month, Q3 2018

| Month | Е | М | MW | NE | NW | SE | s | W | Total |
|-------|----|----|----|----|----|----|----|---|-------|
| Jul | 12 | 3 | 7 | 4 | 4 | 1 | 7 | 0 | 38 |
| Aug | 19 | 4 | 2 | 4 | 0 | 4 | 11 | 4 | 48 |
| Sep | 18 | 5 | 2 | 21 | 3 | 7 | 3 | 4 | 63 |
| Total | 49 | 12 | 11 | 29 | 7 | 12 | 21 | 8 | 149 |



Figure 1. Seasonal distribution of human salmonellosis notifications, 2015 to end Q3 2018

Table 7 shows the serotypes for the Salmonella isolates typed by the NSSLRL in the third quarter of 2018 by HSE area (n=147). The commonest human serotypes reported this quarter were S. Typhimurium (n=45, 31%) and S. Enteritidis (n=31, 21%).

| Table 7. Seroty | pes of | human | Salmo | onella i | isola | tes |
|-----------------|---------------|--------|-------|----------|-------|-----|
| referred to NSS | SLRL Q | 3 2018 | | | | |
| | | | | | | |

| referred to NSS | | | | | | | | | |
|------------------|----|----|----|----|----|----|----|---|-------|
| Serotype | Е | Μ | MW | NE | NW | SE | S | W | Total |
| 4,[5],12:i:- | 7 | 3 | 1 | 11 | 0 | 1 | 5 | 0 | 28 |
| Agama | 0 | 0 | 1 | 0 | 1 | 0 | 0 | 0 | 2 |
| Agona | 2 | 0 | 0 | 0 | 0 | 0 | 1 | 0 | 3 |
| Ajiobo | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Bareilly | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Bochum | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 0 | 1 |
| Bovismorbificans | 0 | 0 | 0 | 0 | 0 | 2 | 0 | 0 | 2 |
| Braenderup | 1 | 1 | 1 | 1 | 0 | 0 | 0 | 0 | 4 |
| Bredeney | 1 | 0 | 0 | 1 | 0 | 0 | 0 | 0 | 2 |
| Chester | 0 | 0 | 0 | 1 | 0 | 0 | 1 | 0 | 2 |
| Coeln | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Derby | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Enteritidis | 7 | 6 | 3 | 5 | 1 | 3 | 4 | 2 | 31 |
| Goldcoast | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 1 |
| Heidelberg | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Idikan | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| IIIb 61:c:z35 | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Infantis | 2 | 0 | 0 | 0 | 0 | 0 | 1 | 0 | 3 |
| Isangi | 0 | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 1 |
| Java | 0 | 0 | 0 | 0 | 0 | 0 | 3 | 0 | 3 |
| Kentucky | 4 | 0 | 0 | 0 | 0 | 0 | 1 | 0 | 5 |
| Kingston | 2 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 2 |
| Litchfield | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| London | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Mbandaka | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 0 | 1 |
| Newport | 3 | 0 | 0 | 1 | 2 | 0 | 2 | 1 | 9 |
| Panama | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 0 | 1 |
| Paratyphi A | ~ | ~ | ~ | ~ | ~ | ~ | ~ | ~ | 2 |
| Rissen | 1 | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 2 |
| Saintpaul | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 0 | 1 |
| Schwarzengrund | 2 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 2 |
| Stanley | 3 | 0 | 0 | 0 | 0 | 0 | 0 | 1 | 4 |
| Teko | 0 | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Telelkebir | 0 | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 1 |
| Thompson | 0 | 0 | 0 | 0 | 1 | 0 | 0 | 0 | 1 |
| Typhi | ~ | ~ | ~ | ~ | ~ | ~ | ~ | ~ | 3 |
| Typhimurium | 4 | 1 | 3 | 1 | 1 | 3 | 2 | 2 | 17 |
| Unnamed | 1 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 1 |
| Virchow | 2 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 2 |
| Total | 53 | 12 | 9 | 21 | 8 | 13 | 24 | 7 | 147 |

Data Source: NSSLRL

^{*}Includes 28 cases of monophasic *S*.Typhimurium 4,5,12:i:-

Table 8. Confirmed Salmonella notifications byserotype and travel status, Q3 2018 [n(%)]

| Serotype | Indigenous | Travel- associated | Unk/not specified | Total |
|-----------------|------------|-----------------------|----------------------|---------------|
| S. Enteritidis | 8 (18%) | 19 (26%) | 6 (22%) | 33 (23%) |
| S. Typhimurium* | 19 (41%) | 19 (26%) | 8 (30%) | 46 (32%) |
| Other | 17 (37%) | 32 (45%) | 11 (41%) | 60 (41%) |
| Salmonella spp | 2 (4%) | 2 (3%) | 2 (7%) | 6 (4%) |
| Total | 46 (100%) | 72 (100%) | 27 (100%) | 145 (100%) |

Note: Data source CIDR. Travel status is inferred from *Country of Infection* variable on CIDR.

Note excludes probable notifications

Includes monophasic S.Typhimurium 4,5,12:i:-

Table 8 shows the serotype distribution of confirmed *Salmonella* cases by travel status this quarter among salmonellosis notifications on CIDR. 49% (n=72) were travel-associated, 32%% (n=46) were indigenous and for 27 cases, the country of infection was unknown/not specified.

S. Typhi and S. Paratyphi

There were four cases of typhoid reported in Q3 2018 – one associated with travel to East Africa and three associated with travel to the Indian Sub-Continent.

There were two cases of Paratyphoid notified this quarter, both associated with travel to India.

Outbreaks of S. Typhi and S. Paratyphi

There were no outbreaks of typhoid or paratyphoid notified in Q3 2018.

Outbreaks of salmonellosis

There were no outbreaks of salmonellosis notified in Q3 2018. (Tables 1 & 2).

VEROTOXIGENIC E. COLI (VTEC)

Verotoxigenic *E. coli* (VTEC) became a notifiable disease on January 1st 2012. Previously, VTEC were notified under the category of Enterohaemorrhagic *E. coli* between 2004 and 2011.

Five hundred cases of VTEC were notified this quarter, the regional distribution of which is shown in Table 9. This compares with 353 VTEC cases notified in Q3 2017 and 318 in Q3 2016 (figure 2).

Table 9 shows the number of VTEC cases reported by case classification and HSE-area and Table 10 shows the number of VTEC cases by serogroup and month, Q3 2018.

Table 9. Number VTEC notified by caseclassification and HSE-area, Q3 2018

| Case classification | Е | Μ | мw | NE | NW | SE | S | W | Total |
|------------------------|----|----|----|----|----|----|----|----|-------|
| Confirmed | 85 | 46 | 42 | 38 | 13 | 63 | 92 | 77 | 456 |
| Probable | 2 | 13 | 5 | 8 | 0 | 2 | 1 | 13 | 44 |
| Possible | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 | 0 |
| Total | 87 | 59 | 47 | 46 | 13 | 65 | 93 | 90 | 500 |

Table 10. VTEC notified by serogroup and month, Q3 2018

| Month | O157 | O26 | Other | Total |
|-------|------|-----|-------|-------|
| Jul | 78 | 70 | 67 | 215 |
| Aug | 18 | 38 | 84 | 140 |
| Sep | 40 | 34 | 71 | 145 |
| Total | 136 | 142 | 222 | 500 |

Twenty-four VTEC cases notified this quarter were reported as having developed HUS – ten O26, nine O157, one O146, one O146 and three ungroupable strains.



Figure 2. Seasonal distribution of VTEC cases notified 2015 to end Q3 2018

The HSE-DML Public Health Laboratory at Cherry Orchard Hospital, Dublin provides a national *E. coli* O157 and non-O157 diagnostic service for clinical samples, including *E. coli* serotyping, verotoxin detection and VTEC molecular typing. Table 11 shows the *vt* types of VTEC cases notified in Q3 2018.

Table 11. Verotoxin typing profiles of *E. coli*referred to the HSE DML Public HealthLaboratory, Cherry Orchard Hospital in Q32018

| Serogroup | vt1 | vt2 | vt1+vt2 | Not spec. | Total |
|-----------|-----|-----|---------|--------------|-------|
| 0157 | 0 | 54 | 74 | 8 | 136 |
| O26 | 24 | 1 | 113 | 4 | 142 |
| Other | 59 | 93 | 37 | 33 | 222 |
| Total | 83 | 148 | 224 | 45 | 500 |

Data Source: PHL Cherry Orchard

Outbreaks of VTEC infection

There were seven general and fifty-seven family outbreaks of VTEC infection reported during this quarter (Tables 1 & 2).

CAMPYLOBACTER

Human campylobacteriosis became a notifiable disease on January 1st 2004. Prior to this, human campylobacter infection was notified under the category of 'Food Poisoning (bacterial other than Salmonella)'. The notifications for the third quarter of 2018 are shown in Table 12. There were 820 cases of campylobacteriosis notified in Q3 2018 compared to 797 in the same period in 2017 and 759 in Q3 2016 (Figure 3).

From August 2017, campylobacter notifications from HSE-East are based on laboratory testing results rather than patient episodes. Notifications from HSE-E may also refer to area of laboratory testing rather than area of patient residence.

Table 12. Campylobacter notifications byHSE-Area and month, Q3 2018

| Month | Е | М | MW | NE | NW | SE | s | W | Total |
|-------|-----|----|----|----|----|----|-----|----|-------|
| Jul | 92 | 24 | 30 | 23 | 12 | 36 | 39 | 27 | 283 |
| Aug | 120 | 14 | 15 | 31 | 11 | 39 | 56 | 28 | 314 |
| Sep | 96 | 11 | 14 | 23 | 10 | 18 | 32 | 19 | 223 |
| Total | 308 | 49 | 59 | 77 | 33 | 93 | 127 | 74 | 820 |

Human cryptosporidiosis became a notifiable disease on January 1st 2004. Prior to this, cryptosporidiosis was notifiable in Ireland only in young children under the category 'Gastroenteritis in Children Under 2'. In Q3 2018, 81 cases of cryptosporidiosis were notified (Table 13), compared to 108 in the same period in 2017 and 73 in Q3 2016 (Figure 4).

Table 13. Cryptosporidiosis notifications byHSE-Area and month, Q3 2018

| Month | Е | М | MW | NE | NW | SE | S | W | Total |
|-------|----|----|----|----|----|----|----|----|-------|
| Jul | 4 | 2 | 2 | 2 | 1 | 2 | 6 | 5 | 24 |
| Aug | 5 | 6 | 2 | 1 | 0 | 3 | 4 | 6 | 27 |
| Sep | 9 | 5 | 1 | 1 | 3 | 4 | 2 | 5 | 30 |
| Total | 18 | 13 | 5 | 4 | 4 | 9 | 12 | 16 | 81 |

Outbreaks of cryptosporidiosis

There were two family outbreaks of cryptosporidiosis reported in quarter 3 2018 (Tables 1 and 2).

Outbreaks of Campylobacter infection

There was one general outbreak of campylobacteriosis reported in Q3 2018 (Tables 1 and 2).





CRYPTOSPORIDIUM



Figure 4. Seasonal distribution of cryptosporidiosis notifications 2015 to end Q3 2018

NOROVIRUS

Human noroviral infection became a notifiable disease on January 1st 2004. Since March 2013, norovirus notifications from HSE-East are based on laboratory testing results rather than patient episodes. Notifications from HSE-E may also refer to area of laboratory testing rather than area of patient residence.

There were 222 cases notified in the third quarter of 2018 (Table 14). These data are certainly an under-ascertainment of the true burden of disease due to this pathogen.

Table 14. Norovirus notifications by HSE-Area and month, Q3 2018

| Month | Е | М | MW | NE | NW | SE | S | w | Total |
|-------|-----|----|----|----|----|----|----|----|-------|
| Jul | 32 | 0 | 2 | 5 | 1 | 0 | 16 | 6 | 62 |
| Aug | 53 | 9 | 10 | 12 | 0 | 2 | 11 | 2 | 99 |
| Sep | 30 | 7 | 5 | 5 | 2 | 0 | 7 | 5 | 61 |
| Total | 115 | 16 | 17 | 22 | 3 | 2 | 34 | 13 | 222 |

Norovirus outbreaks

Norovirus or suspect viral aetiology is the commonest cause of outbreaks of acute

Prior to 2004, rotavirus cases were notified under the "Gastroenteritis in children under two years" disease category. From 2004 to 2010, rotavirus was notifiable in all age groups under the "Acute Infectious Gastroenteritis" (AIG) disease category, until it became notifiable as a disease in its own right under the Infectious Diseases (Amendment) Regulations 2011 (S.I. No. 452 of 2011). Between March 2013 and July 2017, rotavirus notifications from HSE-East were based on laboratory testing results rather than patient episodes.

Rotavirus notifications for the third quarter of 2018 are shown in Table 15 and Figure 6.

Table 15. Rotavirus infection by HSE-Areaand month, Q3 2018

| Month | Е | Μ | MW | NE | NW | SE | S | W | Total |
|-------|----|----|----|----|----|----|----|----|-------|
| Jul | 21 | 8 | 4 | 6 | 1 | 10 | 17 | 5 | 72 |
| Aug | 14 | 3 | 3 | 8 | 3 | 7 | 17 | 2 | 57 |
| Sep | 21 | 1 | 1 | 5 | 3 | 0 | 5 | 7 | 43 |
| Total | 56 | 12 | 8 | 19 | 7 | 17 | 39 | 14 | 172 |

gastroenteritis in Ireland. In the third quarter of 2018, there were fourteen outbreaks confirmed as being caused by this virus, involving at least 136 people becoming ill, as outlined in tables 1 & 2. The seasonal trend is outlined in figure 5.





ROTAVIRUS INFECTION



Figure 6. Seasonal distribution of rotavirus notifications, 2015 to end Q3 2018

Outbreaks of rotavirus

There was one family and one general outbreak of rotavirus notified this quarter (Table 2).

On January 1st 2004, infection with *Shigella* spp. became notifiable as 'Shigellosis'. Prior to this, it was notifiable as 'Bacillary Dysentery'.

During Q3 2018, thirty-seven cases of shigellosis were notified (Table 5). This compares with thirty-six cases notified in Q3 2017 and twenty-one in Q3 2016.

Eighteen cases were travel related and the country of infection was reported as Ireland for a futher ten cases. The country of infection was reported as unknown/not specified for the remaining nine cases.

Twenty-one isolates were referred for typing to NSSLRL this quarter (Table 16).

Table 16: Species and serotype distribution of human *Shigella* isolates referred to the NSSLRL in Q3 2018

| Serotype | Number of isolates |
|-----------------------------|--------------------|
| Shigella flexneri 1c | 1 |
| Shigella flexneri 2a | 3 |
| Shigella flexneri 6 | 1 |
| Shigella flexneri Y variant | 1 |
| Shigella sonnei | 15 |

Data Source: NSSLRL

Outbreaks of shigellosis

There was one family outbreak of shigellosis notified in Q2 2018 (Table 2).

GIARDIA

Human giardiasis became a notifiable disease on January 1st 2004. Prior to this, giardiasis was notifiable in Ireland only in young children under the category 'gastroenteritis in children under 2 years'.

During Quarter 3, 2018, sixty-seven cases of giardiasis were notified (Table 5); this compares with 70 cases notified in Q3 2017 and 44 in Q3 2016.

Fourteen cases were reported to have acquired their illness abroad. Country of infection was reported as Ireland for thirteen cases and 'not specified' or 'unknown' for the remaining forty cases.

Outbreaks of giardiasis

There was one general and four family outbreaks of giardiasis notified in Q3 2018 (Table 2).

LISTERIA

Human listeriosis became a notifiable disease on January 1st 2004. Prior to this, listeriosis was notified under the category of 'Food Poisoning (bacterial other than Salmonella)' or 'Bacterial Meningitis' as appropriate.

There were eight adult and two pregnancy related cases of listeriosis notified in Q3 2018, compared to two cases in quarter 3 2017 and four in quarter 3 2016.

Ten isolates were referred for typing to NSSLRL this quarter (Table 17).

Table 17: Serotypes of human Listeriaisolates referred to the NSSLRL in Q3 2018

| Serotype | Number of isolates |
|----------|--------------------|
| 4b | 6 |
| 1/2a | 3 |
| 1/2b | 1 |

Data Source: NSSLRL

Outbreaks of listeriosis

There were no outbreaks of listeriosis notified in Q3 2018 (Table 2).

FOODBORNE INTOXICATIONS

Bacillus cereus foodborne infection/intoxication, botulism, *Clostridium perfringens* (type A) foodborne disease and staphylococcal food poisoning became notifiable diseases on January 1st 2004. Prior to this, these diseases were notified under the category of 'Food Poisoning (bacterial other than Salmonella)'.

There were no cases of foodborne infection/intoxication reported in Q3 2018.

NON-IID ZOONOTIC DISEASES

Non-IID zoonoses now notifiable include: anthrax, brucellosis, echinococcosis, leptospirosis, plague, Q fever, toxoplasmosis, trichinosis and rabies. The Q3 2018 notifications of these zoonotic diseases are reported by HSE-Area in Table 5.

Six cases of toxoplasmosis were notified in this quarter. This compares with eight cases notified in the same period in 2017 and eight cases in Q3 2016.

There were three cases of leptospirosis notified in Q3 2018. This compares with nine cases in Q3 2017 and seven cases in Q3 2016. Two leptospirosis cases this quarter are believed to have acquired their infection occupationally while one case is associated with travel to South East Asia.

There were no cases of brucellosis notified in Q3 2018. This compares with none in Q3 2017 and none in the same period in 2016.

There was one case of echinococcosis reported in Q3 2018. This compares with no cases in the same period in 2017 and one case in Q3 2016.

There were no cases of trichinosis or Q Fever notified this quarter.

MALARIA

Malaria has been a notifiable disease for many years. The Q3 2018 notifications are reported in Table 5 by HSE-Area.

Sixteen cases of malaria were notified in Q3 2018. This compares with twenty-nine cases reported in Q3 2017 and forty-five in Q3 2016.

Fifteen cases this quarter were reported as *P. falciparum* and the organism was not specified for one case.

Eight cases were exposed in Sub-Saharan Africa. Country of infection is unknown/not specified for the remaining seven cases this quarter.

Six cases cited 'visiting family in country of origin' as their reason for travel while two cases reported business/professional travel. Travel information was not specified/unknown for the remaining seven cases this quarter.

OTHER NOTIFIABLE VECTORBORNE DISEASES

Under Infectious Diseases (Amendment) Regulations 2011 (S.I. No. 452 of 2011) (Sept 2011), Chikungunya disease, Dengue, Lyme disease (neuroborreliosis) and West Nile fever were made notifiable. Zika virus infection is a notifiable disease in Ireland under the Infectious Diseases (Amendment) Regulations 2016 (S.I. No. 276 of 2016).

The Q3 2018 notifications are reported in Table 5 by HSE-Area.

There were four cases of Lyme disease (neuroborreliosis) reported in Q3 2018. This

compares with four cases in the same period in 2017 and ten cases in Q3 2016.

There were five cases of Dengue fever notified in Q3 2018. This compares with four cases in the same period in 2017 and four cases in Q3 2016.

One case this quarter was associated with travel to the Caribbean. Country of infection was not specified for the remaining four cases.

There were no notifications of Chikungunya disease, West Nile or Zika virus infection fever this quarter.

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