Nationally Notifiable Infectious Diseases and Conditions, United States: Annual Tables

TABLE 1. Annual reported cases of notifiable diseases and rates per 100,000, United States, excluding U.S. Territories and Non-U.S. Residents, 2023

Disease	Case Count	Rate
Anthrax	—	—
Arboviral diseases		
Chikungunya virus disease	150	0.04
Eastern equine encephalitis virus disease		
Neuroinvasive	7	0.00
Non-neuroinvasive	—	—
Jamestown Canyon virus disease		
Neuroinvasive	20	0.01
Non-neuroinvasive	7	0.00
La Crosse virus disease		
Neuroinvasive	34	0.01
Non-neuroinvasive	1	0.00
Powassan virus disease		
Neuroinvasive	47	0.01
Non-neuroinvasive	2	0.00
St. Louis encephalitis virus disease		
Neuroinvasive	14	0.00
Non-neuroinvasive	7	0.00
West Nile virus disease		
Neuroinvasive	1,791	0.53
Non-neuroinvasive	839	0.25
Western equine encephalitis virus disease		
Neuroinvasive	—	—
Non-neuroinvasive	—	—
Babesiosis		
Confirmed	3,116	1.08
Probable	470	0.16
Total	3,586	1.24
Botulism		
Foodborne	17	0.01
Infant	174	4.77
Other (wound & unspecified)	14	0.00
Total	205	0.06
Brucellosis	145	0.04
Campylobacteriosis	81,547	24.35
Candida auris		
Clinical*	3,200	1.04
Screening†	4,366	1.30
Carbapenemase-producing organisms (CPO), total‡, §	6,910	2.50
Chancroid	4	0.00
Chlamydia trachomatis infection	1,648,444	492.20
Cholera	19	0.01
Coccidioidomycosis¶		
Confirmed	20,685	14.65
Probable	327	0.23
Total	21,012	14.88
Coronavirus Disease 2019 (COVID-19)		
Confirmed	4,896,251	1706.69

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Disease	Case Count	Rate
Probable#	2,474,171	862.43
Total	7,370,422	2569.12
Cryptosporidiosis		
Confirmed	10,584	3.16
Probable	2,726	0.81
Total	13,310	3.97
Cyclosporiasis	4,463	1.42
Dengue virus infections**		
Dengue	1,944	0.58
Dengue-like illness	23	0.01
Severe dengue	53	0.02
Diphtheria	2	0.00
Ehrlichiosis and Anaplasmosis		
Anaplasma phagocytophilum infection	7,279	2.26
Ehrlichia chaffeensis infection	1,907	0.59
Ehrlichia ewingii infection	26	0.01
Undetermined ehrlichiosis/anaplasmosis	89	0.03
Giardiasis	16,178	6.08
Gonorrhea††	601,262	179.53
Haemophilus influenzae, invasive disease		
Age <5 years		
Non-b serotype	139	0.75
Nontypeable	255	1.38
Serotype b	34	0.18
Unknown serotype	228	1.23
All ages, all serotypes	6,827	2.04
Hansen's disease	88	0.03
Hantavirus infection, non-hantavirus pulmonary syndrome‡‡	1	0.00
Hantavirus pulmonary syndrome	26	0.01
Hemolytic uremic syndrome post-diarrheal	276	0.09
Hepatitis, Viral Disease§§		
Hepatitis A, acute	1,643	0.49
Hepatitis B, acute	2,212	0.66
Hepatitis B, perinatal infection	7	NC
Hepatitis C, acute		
Confirmed	4,964	1.52
Probable	574	0.18
Total	5,538	1.69
Hepatitis C, perinatal infection	235	NC
Human immunodeficiency virus diagnoses	38,850	11.60
Influenza-associated pediatric mortality	142	0.19
Invasive pneumococcal disease		
Age <5 years		
Confirmed	1,114	7.43
Probable	67	0.45
Total	1,181	7.87
All ages		
Confirmed	20,551	8.18
Probable	600	0.24

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Disease	Case Count	Rate
Total¶¶	21,151	8.42
Legionellosis	8,101	2.42
Leptospirosis	94	0.04
Listeriosis##		
Confirmed	934	0.28
Probable	64	0.02
Total	998	0.30
Lyme disease***		
Confirmed	3,530	1.06
Probable	85,938	25.77
Total	89,468	26.83
Malaria	2,606	0.78
Measles†††		
Imported	22	0.01
Indigenous	42	0.01
Total	64	0.02
Melioidosis‡‡‡	6	0.00
Meningococcal disease		
All serogroups	438	0.13
Other serogroups	33	0.01
Serogroup B	44	0.01
Serogroups ACWY	260	0.08
Unknown serogroup	101	0.03
Mpox	1,730	0.52
Mumps	433	0.13
Novel Influenza A virus infections	3	0.00
Pertussis	7,063	2.11
Plague§§§	2	0.00
Poliomyelitis, paralytic	—	—
Poliovirus infection, nonparalytic	—	—
Psittacosis	1	0.00
Q fever		
Acute	181	0.05
Chronic	38	0.01
Total	219	0.07
Rabies		
Animal	3,709	1.11
Human	—	—
Rubella	11	0.00
Rubella, congenital syndrome	1	0.03
Salmonella Paratyphi infection¶¶¶	181	0.05
Salmonella Typhi infection###	535	0.16
Salmonellosis (excluding *S. Typhi* infection and *S. Paratyphi* infection)****	60,934	18.19
Severe acute respiratory syndrome-associated coronavirus disease	—	—
Shiga toxin-producing *Escherichia coli* (STEC)	18,844	5.63
Shigellosis	21,769	6.50
Smallpox	—	—
Spotted fever rickettsiosis		
Confirmed	56	0.02

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Disease	Case Count	Rate
Probable	1,148	0.35
Total	1,204	0.36
Streptococcal toxic shock syndrome	891	0.41
Syphilis		
Congenital††††	3,882	106.40
Primary and secondary	52,988	15.82
Total‡‡‡	209,119	62.44
Tetanus	18	0.01
Toxic shock syndrome (other than Streptococcal)	39	0.02
Trichinellosis	1	0.00
Tuberculosis	9,633	2.88
Tularemia	196	0.06
Vancomycin-intermediate *Staphylococcus aureus*	59	0.02
Vancomycin-resistant *Staphylococcus aureus*§§§§	1	0.00
Varicella morbidity¶¶¶¶	6,896	2.47
Varicella mortality	5	0.00
Vibriosis		
Confirmed	1,542	0.47
Probable	1,880	0.57
Total	3,422	1.04
Viral hemorrhagic fevers		
Chapare virus	—	—
Crimean-Congo hemorrhagic fever virus	—	—
Ebola virus	—	—
Guanarito virus	—	—
Junin virus	—	—
Lassa virus	—	—
Lujo virus	—	—
Machupo virus	—	—
Marburg virus	—	—
Sabia virus	—	—
Yellow fever	—	—
Zika virus		
Zika virus disease, congenital####	—	—
Zika virus disease, non-congenital	6	0.00
Zika virus infection, congenital####	—	—
Zika virus infection, non-congenital	1	0.00

—: No reported cases — The reporting jurisdiction did not submit any cases to CDC.
NC: Not Calculated — There is insufficient data available to support this statistic.

* Beginning in 2023, only confirmed cases of *Candida auris* clinical are published to align with CSTE position statement 22-ID-05, whereas in previous years, confirmed and probable case classification statuses were published. Additionally, there may be case count discrepancies of *Candida auris* clinical cases reported by the NNDSS and the CDC's Mycotic Diseases Branch due to differences in data sources, reporting and aggregation methods. Please refer to the Mycotic Diseases Branch's Tracking C. auris | Candida auris (C. auris) | CDC for *Candida auris* case data reported by jurisdictions. These data are submitted to the CDC separately of NNDSS by jurisdictions and are published by location of the facility. Please also see Note #8.
† Beginning in 2023, *Candida auris* screening was added as a nationally notifiable condition, and confirmed cases are published to align with the approved CSTE position statement 22-ID-05. Additionally, there may be case count discrepancies of *Candida auris* screening cases reported by the NNDSS and the CDC's Mycotic Diseases Branch due to differences in data sources, reporting and aggregation methods. Please refer to the Mycotic Diseases Branch's Tracking C. auris | Candida auris (C. auris) | CDC for *Candida auris* case data reported by jurisdictions. These data are submitted to the CDC separately of NNDSS by jurisdictions and are published by location of the facility. Please also see Note #8.
‡ Beginning in 2023, cases began to be reported as carbapenemase-producing organisms, clinical or carbapenemase-producing organisms, screening. In previous years, only carbapenemase-producing carbapenem-resistant *Enterobacteriaceae* (CP-CRE) cases were reported.
§ CPO total is the sum of CP-CRE cases reported in MMWR year 2023, CPO, clinical cases, and CPO, screening cases.
¶ For coccidioidomycosis surveillance reporting purposes, jurisdictions are grouped into high- and low-incidence categories. Probable cases are only reported from low-incidence jurisdictions, while confirmed cases are reported from both high- and low-incidence jurisdictions.
# Of the reporting areas that submitted 2023 aggregate COVID-19 data to CDC, four did not submit probable cases. American Samoa, Commonwealth of Northern Mariana Islands, New York (excluding New York City), and U.S. Virgin Islands did not collect probable cases.
** Counts include confirmed and probable dengue cases.
†† Beginning in 2023, confirmed and probable cases of gonorrhea are published to align with the approved CSTE position statement 22-ID-03, whereas in previous years, all case classification statuses were published. This change may cause a decrease in published case counts when compared to previous years.
‡‡ Case counts may include Old World hantavirus infections, such as Seoul virus.
§§ Chronic hepatitis B and chronic hepatitis C data are not included in NNDSS tables but reported case counts are included in the annual Viral Hepatitis Surveillance Report, 2023, published online by CDC's Division of Viral Hepatitis, available at https://www.cdc.gov/hepatitis/php/statistics-surveillance/.

¶¶ Counts include drug resistant and susceptible cases of Invasive Pneumococcal Disease. This condition was previously named *Streptococcus pneumoniae* invasive disease and cases were reported to CDC using different event codes to specify whether the cases were drug resistant or in a defined age group, such as <5 years.

Before 2019, probable cases were not reported, and cases in neonates ≤60 days of age were counted as one case in a mother-infant pair. Beginning in 2019, confirmed and probable cases are being reported, and maternal and neonatal cases are being counted separately.

*** For Lyme surveillance reporting purposes, jurisdictions are grouped into high- and low-incidence categories. Confirmed cases are only reported from low-incidence jurisdictions; however, probable cases are reported from both high- and low-incidence jurisdictions. For more information on jurisdiction classifications, visit https://www.cdc.gov/lyme. Currently high-incidence jurisdictions include Connecticut, Delaware, Maine, Maryland, Massachusetts, Minnesota, New Hampshire, New Jersey, New York (excluding New York City), New York City, Pennsylvania, Rhode Island, Vermont, Virginia, West Virginia, Wisconsin, and the District of Columbia.

††† Measles is considered imported if the disease was acquired outside of the United States and is considered indigenous if the disease was acquired anywhere within the United States or it is not known where the disease was acquired.

‡‡‡ Beginning in 2023, melioidosis was added as a nationally notifiable condition, and confirmed and probable cases are published to align with the approved CSTE position statement 22-ID-08.

§§§ Beginning in 2020, confirmed and probable plague cases began to be combined and published.

¶¶¶ Beginning in January 2019, cases began to be reported as *Salmonella* Paratyphi infection. In 2018, cases were reported as paratyphoid fever. Prior to 2018, cases of paratyphoid fever were considered salmonellosis.

Beginning in January 2019, cases began to be reported as *Salmonella* Typhi infection. In previous years, cases were reported as typhoid fever.

**** Beginning in January 2019, cases began to be reported as salmonellosis (excluding *Salmonella* Typhi infection and *Salmonella* Paratyphi infection). In 2018, cases were reported as salmonellosis (excluding paratyphoid fever and typhoid fever). Prior to 2018, cases of paratyphoid fever were considered salmonellosis.

†††† Congenital syphilis cases are usually assigned to the mother's state of residence at the time of delivery. Data for congenital syphilis are aggregated by the infant's year of birth.

‡‡‡‡ Includes the following categories: primary; secondary; early non-primary non-secondary; unknown duration or late; and congenital syphilis.

§§§§ Vancomycin-resistant *Staphylococcus aureus* cases reported in this table may not have been verified by CDC. CDC verified 0 vancomycin-resistant *Staphylococcus aureus* cases in 2023.

¶¶¶¶ Beginning in 2023, varicella mortality will have an age restriction of <50 years applied to the numerator (i.e., case counts).

Data reported to ArboNET using the national surveillance case definition for congenital Zika virus infection (CSTE Position Statement 16-ID-01).

Notes:

1. These are **annual** cases of selected infectious national notifiable diseases from the National Notifiable Diseases Surveillance System (NNDSS). NNDSS data reported by the 50 states, New York City, the District of Columbia, and the U.S. territories are collated and published. Cases are reported by state health departments to CDC weekly. Because source datasets may be updated as additional information is received, statistics in publications based on that source data may differ from what is presented in these tables. Source datasets for the 2023 annual tables were officially closed on November 7, 2024.

2. The list of national notifiable Infectious diseases and conditions for 2023 and their national surveillance case definitions are available by navigating to the Surveillance Case Definitions | CDC web page, selecting "2023" for the notifiable condition list year, checking "Infectious" conditions, and clicking "Get Notifiable List by Year". Publication criteria for the finalized 2023 data are available at https://www.cdc.gov/nndss/infectious-disease/notice-to-data-users.html. See also Guide to Interpreting Provisional and Finalized NNDSS Data.

3. Population estimates for incidence rates are July 1st, 2023 postcensal estimates of the resident population of the United States for July 1, 2020, to July 1, 2023, by year, county, single year of age (range: 0 to 85+ years), bridged-race (American Indian or Alaska Native, Asian or Pacific Islander, Black or African American, White), Hispanic ethnicity (Hispanic or Latino, not Hispanic or Latino), and sex (Female, Male), prepared under a collaborative arrangement with the U.S. Census Bureau and the National Cancer Institute (NCI). The "Vintage 2023" population estimates for years 2020–2023 were released February 2025 by the National Cancer Institute at https://seer.cancer.gov/popdata/. For more information, see https://seer.cancer.gov/popdata/singleages.html. The choice of population denominators for incidence is based on the availability of population data at the time of publication preparation.

4. Annual tables for 2016 and later years are available on CDC WONDER's NNDSS Annual Summary Data Query.

5. Annual summary reports from 1993–2015 are available as published in the *Morbidity and Mortality Weekly Report*.

6. NNDSS annual tables since 1952 are available at CDC Stacks. To find them, search for "NNDSS" under Collections. Once in NNDSS Collections, navigate to the "Resource Type Specific" section on the left-hand side and select "Annual Reports" (1952–2015) or "Annual Tables" (2016–present).

7. For most conditions, national incidence rates are calculated as the number of reported cases for each infectious disease or condition divided by the U.S. resident population for the specified demographic population or the total U.S. resident population, multiplied by 100,000. When a national notifiable infectious condition is associated with a specific age restriction, the same restriction was applied to the population in the denominator of the incidence rate calculation. In addition, population data from reporting jurisdictions in which the disease or condition was not reportable or not available were excluded from the denominator of the incidence rate calculations.

Age restrictions in the numerator and denominator are applied for the following childhood conditions:

Zika virus disease, congenital (age restriction in numerator and denominator is <1 year)
Zika virus infection, congenital (age restriction in numerator and denominator is <1 year)
Haemophilus influenzae, invasive disease <5 years (age restriction in numerator and denominator is <5 years)
Invasive pneumococcal disease <5 years (age restriction in numerator and denominator is <5 years)
Influenza associated pediatric mortality (age restriction in numerator and denominator is <18 years)
Infant botulism (age restriction in numerator and denominator is <1 year)
Congenital rubella syndrome (age restriction in numerator and denominator is <1 year)
Perinatal hepatitis B infection (age restriction in numerator is ≤24 months; however, rates will not be calculated due to population estimates not being available for the specific sub-population required to align with the defined age restriction criteria)
Perinatal hepatitis C infection (age restriction in numerator is ≤36 months; however, rates will not be calculated due to population estimates not being available for the specific sub-population required to align with the defined age restriction criteria).

Data for congenital syphilis are aggregated by the infant's year of birth. The rate for congenital syphilis is based upon the number of reported cases per 100,000 live births, using natality data for 2023 (National Center for Health Statistics Natality 2023, as compiled from data provided by the Vital Statistics Cooperative Program). Congenital syphilis cases are usually assigned to the mother's state of residence at the time of delivery. The mother's race and ethnicity are used for race- and ethnicity-specific rates of congenital syphilis cases.

8. Surveillance data reported by other CDC programs might vary from data reported in these tables because of differences in 1) the date used to aggregate the data, 2) the timing of reports, 3) the source of the data, 4) surveillance case definitions, and 5) policies regarding case jurisdiction (i.e., which jurisdiction should submit the case notification to CDC).

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National Notifiable Diseases Surveillance System