Publication criteria for nationally notifiable conditions reported to the National Notifiable Diseases Surveillance System, 2018.

<table>
<thead>
<tr>
<th>Code</th>
<th>Notifiable condition</th>
<th>Publication criteria* , †, §</th>
</tr>
</thead>
<tbody>
<tr>
<td>11090</td>
<td><em>Anaplasma phagocytophilium</em></td>
<td>Confirmed and probable</td>
</tr>
<tr>
<td>10350</td>
<td>Anthrax</td>
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</tr>
<tr>
<td>12010</td>
<td>Babesiosis</td>
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</tr>
<tr>
<td>10530</td>
<td>Botulism, foodborne</td>
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</tr>
<tr>
<td>10540</td>
<td>Botulism, infant</td>
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</tr>
<tr>
<td>10550</td>
<td>Botulism, other (includes wound)</td>
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</tr>
<tr>
<td>10548</td>
<td>Botulism, other (unspecified)</td>
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</tr>
<tr>
<td>10549</td>
<td>Botulism, wound</td>
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<td>10020</td>
<td>Brucellosis</td>
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<tr>
<td>11020</td>
<td>Campylobacteriosis</td>
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</tr>
<tr>
<td>10273</td>
<td>Chancroid</td>
<td>All reports</td>
</tr>
<tr>
<td>10073</td>
<td>Chikungunya virus diseases</td>
<td>Confirmed and probable</td>
</tr>
<tr>
<td>10274</td>
<td><em>Chlamydia trachomatis</em></td>
<td>All reports</td>
</tr>
<tr>
<td>10470</td>
<td>Cholera (toxigenic <em>Vibrio cholera</em> O1 or O139)</td>
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</tr>
<tr>
<td>11900</td>
<td>Coccioidiomycosis</td>
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<tr>
<td>11640</td>
<td>Crimean-Congo hemorrhagic fever</td>
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</tr>
<tr>
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<td>Cryptosporidiosis</td>
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<td>Cyclosporiasis</td>
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<tr>
<td>10680</td>
<td>Dengue</td>
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<tr>
<td>11705</td>
<td>Dengue, severe</td>
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<tr>
<td>11704</td>
<td>Dengue-like illness</td>
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<td>Diphtheria</td>
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<td>Eastern equine encephalitis virus disease, neuroinvasive</td>
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<tr>
<td>10062</td>
<td>Eastern equine encephalitis virus disease, nonneuroinvasive</td>
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<td>11630</td>
<td>Ebola hemorrhagic fever</td>
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<tr>
<td>11088</td>
<td><em>Ehrlichia chaffeensis</em></td>
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<td>11089</td>
<td><em>Ehrlichia ewingii</em></td>
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<tr>
<td>11091</td>
<td>Ehrlichiosis/Anaplasmosis, undetermined</td>
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<tr>
<td>11570</td>
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<td>Gonorrhea</td>
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<td>Guanarito hemorrhagic fever</td>
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<td><em>Haemophilus influenzae</em>, invasive disease</td>
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<td>10380</td>
<td>Hansen’s disease (Leprosy)</td>
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<td>11610</td>
<td>Hantavirus infection, non-Hantavirus pulmonary syndrome</td>
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<td>Hantavirus pulmonary syndrome (HPS)</td>
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<td>Hemolytic uremic syndrome, postdiarrheal (HUS)</td>
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<td>Hepatitis B, acute</td>
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<tr>
<td>Code</td>
<td>Notifiable condition</td>
<td>Publication criteria*⁺,†,§</td>
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<td>10104</td>
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<td>Hepatitis C, perinatal infection</td>
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<td>HIV diagnoses</td>
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<td>11061</td>
<td>Influenza-associated pediatric mortality</td>
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<td>11723</td>
<td>Invasive pneumococcal disease (Streptococcus pneumoniae, invasive disease) (all ages)</td>
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<td>10078</td>
<td>Jamestown Canyon virus disease, neuroinvasive</td>
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<td>Jamestown Canyon virus disease, non-neuroinvasive</td>
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<td>Junin hemorrhagic fever</td>
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<td>La Crosse virus disease, neuroinvasive</td>
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<td>11632</td>
<td>Lassa fever</td>
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<td>Legionellosis</td>
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<td>Leptospirosis</td>
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<td>11644</td>
<td>Lujo virus</td>
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<td>Machupo hemorrhagic fever</td>
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<td>Malaria</td>
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<td>11631</td>
<td>Marburg fever</td>
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<td>10140</td>
<td>Measles (rubeola), total</td>
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<td>Meningococcal disease (Neisseria meningitidis)</td>
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<td>Mumps</td>
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<td>Novel influenza A virus infections, initial detections of</td>
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<td>50236</td>
<td>Paratyphoid fever (caused by Salmonella serotypes Paratyphi A, Paratyphi B [tartrate negative], and Paratyphi C)</td>
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<tr>
<td>10190</td>
<td>Pertussis (Whooping Cough)</td>
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<td>Poliomyelitis, paralytic</td>
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<td>Poliovirus infection, nonparalytic</td>
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<td>Powassan virus disease, neuroinvasive</td>
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<td>Powassan virus disease, non-neuroinvasive</td>
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<td>Psittacosis (Ornithosis)</td>
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<td>Q fever, chronic</td>
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<td>10460</td>
<td>Rabies, human</td>
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<tr>
<td>Code</td>
<td>Notifiable condition</td>
<td>Publication criteria* † §</td>
</tr>
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<tr>
<td>10200</td>
<td>Rubella</td>
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<td>Rubella, congenital syndrome (CRS)</td>
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<td>Sabia-associated hemorrhagic fever</td>
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<td>50242</td>
<td>Salmonellosis, (excluding paratyphoid fever and typhoid fever )</td>
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<tr>
<td>10575</td>
<td>Severe acute respiratory syndrome-associated coronavirus (SARS-CoV) disease</td>
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<td>11563</td>
<td>Shiga toxin-producing <em>Escherichia coli</em> (STEC)</td>
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<td>11010</td>
<td>Shigellosis</td>
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<tr>
<td>11800</td>
<td>Smallpox</td>
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<td>10250</td>
<td>Spotted fever rickettiosis</td>
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<td>10051</td>
<td>St. Louis encephalitis virus disease, neuroinvasive</td>
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<tr>
<td>10064</td>
<td>St. Louis encephalitis virus disease, nonneuroinvasive</td>
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<tr>
<td>11700</td>
<td>Streptococcal toxic shock syndrome</td>
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<tr>
<td>10316</td>
<td>Syphilis, congenital</td>
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<td>Syphilis, early non-primary, non-secondary</td>
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<tr>
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<td>Syphilis, primary</td>
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<tr>
<td>10312</td>
<td>Syphilis, secondary</td>
<td>All reports</td>
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<tr>
<td>10310</td>
<td>Syphilis, total primary and secondary</td>
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</tr>
<tr>
<td>10320</td>
<td>Syphilis, unknown duration or late</td>
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<td>10210</td>
<td>Tetanus</td>
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<td>Toxic shock syndrome (other than streptococcal) (TSS)</td>
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<tr>
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<td>Trichinellosis</td>
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<tr>
<td>10230</td>
<td>Tularemia</td>
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<tr>
<td>10240</td>
<td>Typhoid fever (caused by <em>Salmonella typhi</em>)</td>
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<tr>
<td>11663</td>
<td>Vancomycin-intermediate <em>Staphylococcus aureus</em> (VISA)</td>
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<tr>
<td>11665</td>
<td>Vancomycin-resistant <em>Staphylococcus aureus</em> (VRSA)</td>
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<tr>
<td>10030</td>
<td>Varicella morbidity (Chickenpox)</td>
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<tr>
<td></td>
<td>Varicella mortality</td>
<td>Confirmed and probable</td>
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<tr>
<td>11545</td>
<td>Vibriosis (any species of the family <em>Vibrionaceae</em>, other than toxigenic <em>Vibrio cholerae O1 or O139</em>)</td>
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</tr>
<tr>
<td>10056</td>
<td>West Nile virus disease, neuroinvasive</td>
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<tr>
<td>Code</td>
<td>Notifiable condition</td>
<td>Publication criteria* † §</td>
</tr>
<tr>
<td>--------</td>
<td>-----------------------------------------------------------</td>
<td>---------------------------</td>
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<tr>
<td>10049</td>
<td>West Nile virus disease, nonneuroinvasive</td>
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<tr>
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<td>Western equine encephalitis virus disease, neuroinvasive</td>
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<td>Confirmed and probable</td>
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<tr>
<td>10660</td>
<td>Yellow fever</td>
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<tr>
<td>50224</td>
<td>Zika virus disease, congenital</td>
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<tr>
<td>50223</td>
<td>Zika virus disease, non-congenital</td>
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</tr>
<tr>
<td>50222</td>
<td>Zika virus infection, congenital</td>
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</tr>
<tr>
<td>50221</td>
<td>Zika virus infection, non-congenital</td>
<td>Confirmed and probable</td>
</tr>
</tbody>
</table>

* An unknown case classification status is used when a reporting jurisdiction sends aggregate counts of cases or when the surveillance information system of a reporting jurisdiction does not capture case classification data. In both situations, cases are verified to meet the case classification (e.g., confirmed, probable, and suspected) specified in the publication criteria.

† Publication criteria for the National Notifiable Diseases Surveillance System (NNDSS): for a case report of a nationally notifiable disease to be published in the tables, the reporting state or territory must have designated the disease reportable in their state or territory for the year corresponding to the year of report to CDC. After this criterion is met, the disease-specific criteria listed in the above table are applied. When the above-listed table indicates that all reports will be earmarked for publication, this means that cases designated with unknown or suspect case confirmation status will be published just as probable and confirmed cases will be published. Because CSTE position statements customarily are not finalized until July of each year, NNDSS data for the newly added conditions usually are not available from all reporting jurisdictions until January of the year following the approval of the CSTE position statement.

§ Based on case classification status.