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(Accessible Version: https://wonder.cdc.gov/nndss/static/2020/annual/2020-table3.html)

Disease	Jan.	Feb.	Mar.	Apr.	May	June	July	Aug.	Sept.	Oct.	Nov.	Dec.	Unknown	Total
Anthrax												1		1
Arboviral diseases														
Chikungunya virus disease	10	8	1	2	3	_	1	_	3	2	2	5		37
Eastern equine encephalitis virus disease														
Neuroinvasive	—		—	_	_	_	2	5	5	1	—	_	_	13
Non-neuroinvasive	—									_				
Jamestown Canyon virus disease														
Neuroinvasive					1	2	5	1		1				10
Non-neuroinvasive					1	1			1					3
La Crosse virus disease														
Neuroinvasive	—	1	—			1	16	32	26	8				84
Non-neuroinvasive							2	2		_				4
Powassan virus disease														
Neuroinvasive	—			2	3	6	3			1	2	2		19
Non-neuroinvasive								1						· 1
St. Louis encephalitis virus disease														
Neuroinvasive	—			1		1	3	2	3	1	2	1		14
Non-neuroinvasive	—		—	_	_	_	—	1	1	_		_	_	2
West Nile virus disease														
Neuroinvasive	—	1		1	4	6	29	107	266	94	20	30		558
Non-neuroinvasive			2		4	4	11	41	67	21	8	14		172
Western equine encephalitis virus disease														
Neuroinvasive	—		—							_				
Non-neuroinvasive										_				·
Babesiosis														
Total	14	10	12	11	39	231	586	517	162	86	46	106		1,820
Confirmed	12	6	4	8	35	210	553	461	133	74	44	91		1,631
Probable	2	4	8	3	4	21	33	56	29	12	2	15		189
Botulism														
Total	11	15	9	9	17	14	13	22	18	19	26	16		189
Foodborne					3	_		1	2	1	1			8
Infant	9	13	8	7	12	12	11	15	14	15	18	12		146
Other (wound & unspecified)	2	2	1	2	2	2	2	6		3				35
Brucellosis		6					5	7						87
Campylobacteriosis	4,010	4,888	3,044	2,193	4,165	4,817	4,985	6,218	4,643	5,165	3,200	4,436		51,764
<i>Candida auris</i> , clinical <sup>†</sup>	10	17	11	10	13	12	38	38	27	27	23	29		255
Carbapenemase-producing carbapenem-resistant Enterobacteriaceae	166	197	129	99	126	142	122	158	246	189	117	157	_	1,848
Chancroid							_	_	_	_	_	_	_	· _
Chlamydia trachomatis	136,161	181,488	115,313	74,501	114,731	115,809	120,653	163,561	128,965	165,558	118,472	144,625	_	1,579,837
Cholera				_	_	_	_	_	1	_	<u> </u>	_	_	1
Coccidioidomycosis	1,435	2,005	1,329	857	1,262	1,194	1,154	1,765	1,709	2,131	1,959	2,420	_	19,220
Coronavirus Disease 2019 (COVID-19)														
Total	718	1,393	237,279	828,259	786,191	879,911	1,760,192	1,649,270	1,143,635	2,244,093	4,425,759	7,138,430	54,601	21,149,731
Confirmed	587										3,932,456			19,179,356
Probable <sup>§</sup>	131	177	3,636	12,107	14,620	15,159	38,011	67,143	88,008	213,948	493,303	1,018,070	6,062	1,970,375

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Confirmed         512         588         326         291         477         474         580         852         527         510         288         418            Probable         191         179         115         112         112         117         128         128         537         699         222         85         85         103          Cyclosprissis         12          Cyclosprissis         12          Cyclosprissis         12          Cyclosprissis         12   <	Disease	Jan.	Feb.	Mar.	Apr.	May	June	July	Aug.	Sept.	Oct.	Nov.	Dec.	Unknown	Total
Probable         191         179         115         112         165         131         174         200         148         144         76         132            Cyclosporisdis         12         6         17         15         55         815         555         699         262         85         600	Total	703	765	441	403	642	605	754	1,112	675	654	344			7,648
Cyclosportalsis         12         6         17         15         55         815         535         699         262         85         88         103         —           Dengue virus infections*         4         -         1         -															5,861
Dengue virus infections         Image: Participation infection         Image: Participation infection <thimage: infection<="" partin="" th="">         Image: Partin infectio</thimage:>															1,787
Dengue         85         57         40         6         7         4         19         47         47         40         43         46            Dengue ike illness         4         -         1         1         -         1         1         1         6         6         2         1         -         3         -		12	6	17	15	55	815	535	699	262	85	85	103		2,689
Dengue-like liness         4         -         1         1         -         1         1         2         2         2         20         21         17         7         537         220         218         211         231         -         -         -         1         1         2         6         11         10         3         5         6         3         2         -         -         -         -         1         1         1         1         1         1         1         1	Dengue virus infections ¶														
Swere dengue         -         1         1         6         6         2         1         -         -         -         1         1         2         6         1         10         3         5         6         3         2         -         <		85	57	40		7	4	19	47	47	40	43	46		441
Diphtheria         1         -         1         -         -         1         1         0         2         2         20         11         0         3         5         6         3         2         -         -         -         -         1         1         2         2         20         11         10         3         5         6         3         21         23	Dengue-like illness	4	_	1	1	_		_			1	1	1		9
Ehrlichlosis and Anaplesmosis         Intervision         Int	Severe dengue							_							
Anaplasmosis       Image of the second	Diphtheria	1						_							1
phagocycophilum infection         17         14         22         94         395         961         717         537         220         218         211         231            Britichia chaffeensis infection         15         16         20         24         126         252         290         191         99         67         21         57            Britichia evingiinfection         -         1         1         2         6         11         10         6         6         2         1          3            Gonorrhea         49.926         62.609         41.929         33.142         50.922         50.316         52.768         73.103         59.677         76.904         56.754         69.701          6           Haemophilus influenze, invasive disease           1         14         10         2         4          5         6         5         8            Age -5 years           1         4         2         1         1         1          -         4         1         1         1          -															
infection         15         16         20         24         126         252         290         191         99         67         21         57         —           Britichia ewingii infection         -         1         -         -         1         1         6         6         2         1         -         3         -           Gendiaisi         884         1029         749         393         580         608         731         1278         948         10.18         548         677          6           Genorica         49926         62.00         41.929         31.42         50.922         50.316         52.768         73.103         59.677         76.904         56.754         69.701         -         6           Haemophilus influenzee, invasive disease         -         -         -         -         1         14         175         154         119         124         139         137         149         134         220         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -	, phagocytophilum	17	14	22	94	395	961	717	537	220	218	211	231	_	3,637
Undetermined enhichosis/anaplasmosis         -         1         1         2         6         11         10         3         5         6         3         2         -           Gardiasis         894         1,029         749         333         580         688         731         1,278         948         1,018         548         677         -         6           Gonorrhea         49,926         62,609         41,929         33,142         50,922         50,316         52,768         73,103         59,677         76,904         56,754         69,701         -         6           Haresonybuiks influenzae, invasive disease         -         -         -         -         1         4         2         1         1         -         -         -         1         4         2         1         1         -         -         -         1         4         2         1         1         -         -         -         -         1         1         1         -         -         -         -         -         -         -         -         1         1         1         2         1         1         1         -         -		15	16	20	24	126	252	290	191	99	67	21	57	_	1,178
ehrlichiosis/anaplasmosis         -         1         1         2         6         11         10         3         5         6         3         2         -           Giardiais         884         1.029         749         333         580         608         731         1.278         948         1.018         548         677         -         6           Gonorrhea         49.926         62.609         41.929         33.142         50.922         50.316         52.768         73.103         59.677         76.904         56.754         69.701         -         6           All ages, all serotypes         577         654         414         175         154         119         124         139         137         149         134         220         -         1         4         2         1         1         1         -         1         4         2         1         1         1         -         1         4         20         1         1         1         1         -         1         1         1         1         1         1         1         1         1         1         1         1         1         1         1	Ehrlichia ewingii infection	_	1	_	_	1	1	6	6	2	1		3	_	21
Contrictions/stanaplasmosts       Non-beside       Sequence       Se		_	1	1	2	6	11	10	3	5	6	3	2	_	50
Gonorrhea         49,926         62,609         41,922         33,142         50,922         50,316         52,768         73,103         59,677         76,904         56,754         69,701         —         6           Haemophilus influenzae, invasive disease         Image: Construction of the constr															
Haemophilus influenze, invasive disease         Image Solution         Solution </td <td></td> <td>9,453</td>															9,453
invasive disease       Image is all serotypes       577       654       414       175       154       119       124       139       137       149       134       220          Age <5 years  <		49,926	62,609	41,929	33,142	50,922	50,316	52,768	73,103	59,677	76,904	56,754	69,701		677,751
Age -5 years       Image -1	, , , ,														
Serotype b       2       2       1         1       4       2       1       1       1          Non-b serotype       11       14       10       2       4        5       6       5       6       5       8          Non-b serotype       14       28       17       8       10       5       6       4       1       7       8       13          Unknown serotype       14       28       17       8       10       5       6       4       1       7       8       13          Hansen's disease       7       10       3       13       2       6       4       5       6       3       5       4          Hantavirus pulmonary syndrome **       -       -       -       -       1       -       1       -	All ages, all serotypes	577	654	414	175	154	119	124	139	137	149	134	220		2,996
Non-b serotype         11         14         10         2         4         -         5         6         5         6         5         8         -           Nontypeable         19         34         11         6         6         5         5         -         5         3         3         7         -           Unknown serotype         14         28         17         8         10         5         6         4         1         7         8         13         -           Hansen's disease         7         10         3         13         2         6         4         5         6         3         5         4         -           Hantavirus pulmonary syndrome         -         -         -         -         1         -         1         -	Age <5 years														
Nontypeable         19         34         11         6         6         5         -         5         3         3         7         -           Unknown serotype         14         28         17         8         10         5         6         4         1         7         8         13         -           Hansen's disease         7         10         3         13         2         6         4         5         6         3         5         4         -           Hantavirus infection, non- hantavirus pulmonary syndrome **         -         -         -         -         1         -         -         -         -         1         -         -         -         -         -         -         -         -         1         -	Serotype b	2	2	1	—	—	_	1	4		1	1	1		15
Unknown serotype       14       28       17       8       10       5       6       4       1       7       8       13       -         Hansen's disease       7       10       3       13       2       6       4       5       6       3       5       4       -         Hantavirus infection, non-hantavirus pulmonary syndrome **       -       -       -       -       1       -	Non-b serotype	11	14	10	2	4	_	5	6	5	6	5	8		76
Hansen's disease       7       10       3       13       2       6       4       5       6       3       5       4       -         Hantavirus infection, non- hantavirus pulmonary syndrome **	Nontypeable	19						5		5		3	7		104
Hantavirus infection, non-hantavirus pulmonary syndrome **       -       -       -       -       -       1       -       1       -       -       -       -       -       -       -       -       -       -       1       -       1       - </td <td>Unknown serotype</td> <td>14</td> <td>28</td> <td>17</td> <td>8</td> <td>10</td> <td>5</td> <td>6</td> <td>4</td> <td>1</td> <td>7</td> <td>8</td> <td>13</td> <td></td> <td>121</td>	Unknown serotype	14	28	17	8	10	5	6	4	1	7	8	13		121
hantavirus pulmonary syndrome **          1        1   <	Hansen's disease	7	10	3	13	2	6	4	5	6	3	5	4		68
Hantavirus pulmonary syndrome       1       -       -       1       1       1       2       2       2       1       -       3       -         Hemolytic uremic syndrome post-diarrheal       6       9       11       12       13       12       20       19       15       18       9       23       -         Hepatitis, Viral Disease <sup>++</sup> -       -	hantavirus pulmonary	_	_	_	_	_	1	_	1	_	_	_	_	_	2
syndrome post-diarrheal       6       9       11       12       13       12       20       19       15       18       9       23       —         Hepatitis, Viral Disease <sup>++</sup> </td <td>Hantavirus pulmonary</td> <td>1</td> <td>_</td> <td></td> <td>1</td> <td>1</td> <td>1</td> <td>2</td> <td>2</td> <td>2</td> <td>1</td> <td></td> <td>3</td> <td></td> <td>14</td>	Hantavirus pulmonary	1	_		1	1	1	2	2	2	1		3		14
Hepatitis A       889       1,121       690       659       899       710       685       931       807       1,027       720       808       —         Hepatitis B       Image: Constraint of the state of the stat		6	9	11	12	13	12	20	19	15	18	9	23	_	167
Hepatitis B       Image: Second	Hepatitis, Viral Disease <sup>++</sup>														
Hepatitis B       Image: Second		889	1,121	690	659	899	710	685	931	807	1,027	720	808	_	9,946
Acute       189       260       147       155       189       152       155       195       171       185       106       251       —         Perinatal infection       1       —       1       —       2       —       1       1       3       1       —       —       —         Hepatitis C       Image: Confirmed state       520       758       466       349       481       521       453       524       432       553       351       617       —       …															
Hepatitis C         Image: Constraint of the state		189	260	147	155	189	152	155	195	171	185	106	251	_	2,155
Acute         520         758         466         349         481         521         453         524         432         553         351         617         —           Confirmed         440         641         360         275         377         401         358         419         332         448         266         481         —           Probable         80         117         106         74         104         120         95         105         100         105         85         136         —           Perinatal infection         22         21         13         7         10         10         9         15         16         18         10         14         —	Perinatal infection	1	_	1	_	_	2	_	1	1	3	1	_		10
Confirmed         440         641         360         275         377         401         358         419         332         448         266         481         —           Probable         80         117         106         74         104         120         95         105         100         105         85         136         —           Perinatal infection         22         21         13         7         10         10         9         15         16         18         10         14         —	Hepatitis C														
Probable         80         117         106         74         104         120         95         105         100         105         85         136         —           Perinatal infection         22         21         13         7         10         10         9         15         16         18         10         14         —           Human immunodeficiency         Immunodeficiency <t< td=""><td>Acute</td><td>520</td><td>758</td><td>466</td><td>349</td><td>481</td><td>521</td><td>453</td><td>524</td><td>432</td><td>553</td><td>351</td><td>617</td><td>_</td><td>6,025</td></t<>	Acute	520	758	466	349	481	521	453	524	432	553	351	617	_	6,025
Perinatal infection         22         21         13         7         10         10         9         15         16         18         10         14         —	Confirmed	440	641	360	275	377	401	358	419	332	448	266	481		4,798
Human immunodeficiency	Probable	80	117	106	74	104	120	95	105	100	105	85	136		1,227
Human immunodeficiency	Perinatal infection	22	21	13	7	10	10	9	15	16	18	10	14		165
virus diagnoses 3,091 2,672 2,208 1,596 1,930 2,315 2,416 2,266 2,285 2,200 1,522 506 —	Human immunodeficiency virus diagnoses	3,091	2,672	2,208	1,596	1,930	2,315	2,416	2,266	2,285	2,200	1,522	506		25,007
Influenza-associated 43 69 26 9 12 3 1 2 - 5 - 1 -		43	69	26	9	12	3	1	2		5	_	1	_	171
Invasive pneumococcal disease <sup>§§</sup>	Invasive pneumococcal														
		2 323	2.648	1.660	720	545	388	391	469	458	708	612	1.024	_	11,946
															11,718
Probable         44         35         20         15         19         6         6         11         17         21         10         24         —															228
Age <5 years         98         117         85         37         22         18         16         32         28         38         30         40															561
Age of years         36         177         35         37         22         18         10         32         28         36         30         40         40           Confirmed         93         108         82         35         22         18         16         32         27         36         29         38															536
Probable         5         9         3         2         -         -         -         1         2         1         2         -															25

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Disease	Jan.	Feb.	Mar.	Apr.	May	June	July	Aug.	Sept.	Oct.	Nov.	Dec.	Unknown	Total
Legionellosis ¶¶	439	482	307	245	312	481	637	796	773	628	444	766		6,310
Leptospirosis	8	6	5	2	3	4	4	6	3	7	1	2	_	51
Listeriosis ***														
Total	37	54	38	36	54	57	74	99	80	106	57	88		780
Confirmed	36	53	35	35	53	55	71	99	77	103	56			754
Probable	1	1	3	1	1	2	3		3	3		7	_	26
Lyme disease		'										,		20
Total	712	925	506	389	1,169	2,884	3,575	3,217	1,365	1,284	805	1,169		18,000
Confirmed	447	587	300	249	744	1,960	2,606	2,222	892	856	519	741		12,123
Probable	265	338	206	140	425	924	969	995	473	428	286	428		5,877
Malaria	129	130	41	22	18	14	22	46	30	42	44	65		603
Measles ***	125	150			10			10	50					005
Total	4	7									1			12
		5												6
Indigenous	3	2				_								6
Imported	3	2									1			6
Meningococcal disease				4-	40	40	40			40	10	40		242
All serogroups	41 15	55 19	33 8	17	16 2	12 8	13 5	8	6	18	10			242
Serogroups ACWY	7	19	8	4	3	8	2	1	3	6		5		80 - 40
Serogroup B				-			2	1				1		
Other serogroups	4 15	6 15	2	1 11	1 10	1				1				16
Unknown serogroup	164	252	17 107	41	27	3 18	6	6 18	3 13	7	-			106
Mumps	164	252	107	41	27	18	9	18	13	16	12	17		694
Novel Influenza A virus infections	_	_	_	—	_	_	_	_		-	_	-	_	·
Pertussis	1,523	1,867	1,018	511	340	192	110	116	94	103	97	153		6,124
Plague <sup>§§§</sup>		.,	1			1	2	4	1				_	9
Poliomyelitis, paralytic														
Poliovirus infection,														
nonparalytic	_	—	—	—	—	—	—	—	_	-	-		-	·
Psittacosis	_	_	_		3	_	_	2	2	_	1		_	8
O fever														-
Total	8	18	7	5	13	12	11	9	10	6	10	11	_	120
Acute	7	14	5	3	9	8	8	8	7	5			_	90
Chronic	1	4	2		4	4	3	1	3		2			30
Rabies														
Animal	270	274	393	364	430	441	445	559	500	360	230	191	_	4,457
Human	_	_	_		_	_		_					_	
Rubella	2	_	_	1	_	1	_	1	_	_	_	1	_	6
Rubella, congenital syndrome	_	_			_	_	_	_	_		_		_	· _
Salmonella Paratyphi														
infection ¶¶¶	15	19	18	4	1	5	3	3	2	3		2		75
<i>Salmonella</i> Typhi infection ****	32	45	30	9	7	5	6	8	9	11	8	12		182
Salmonellosis (excluding <i>S.</i> Typhi infection and <i>S.</i> Paratyphi infection) <sup>++++</sup>	2,264	2,653	1,774	1,623	3,145	3,929	5,446	6,995	5,362	5,357	3,145	3,749	_	45,442
Severe acute respiratory syndrome-associated coronavirus disease							_	_	_		_		_	· _
Shiga toxin-producing	764	932	583	335	676	936	1,094	1,273	962	998	605	764		9,922
Escherichia coli (STEC)	1,207	1 6 1 1	1,086	426	522	452	506	770	669	760	476	623		
Shigellosis	1,207	1,611	1,080		522	452	anc		609	/60	4/6			9,108
Smallpox								_						
Spotted fever rickettsiosis			~		400	201	470	45-	100					
Total	36	46	31	53	135	201	179	157	139	65	70			1,175
Confirmed	1	2	1	2	8	9	13	9	11	4	-			64
Probable	35	44	30	51	127	192	166	148	128	61	67	62	_	1,111

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Disease	Jan.	Feb.	Mar.	Apr.	May	June	July	Aug.	Sept.	Oct.	Nov.	Dec.	Unknown	Total
Streptococcal toxic shock syndrome	35	46	30	21	19	14	9	12	8	9	5	16	_	224
Syphilis														
Total, all stages <sup>§§§§</sup>	10,852	14,484	9,725	6,753	10,247	9,740	9,646	13,190	10,936	14,447	10,639	13,274	_	133,933
Congenital ¶¶¶¶	186	165	138	162	154	151	211	202	197	195	202	185	_	2,148
Primary and secondary	3,216	4,132	2,744	2,277	3,481	3,207	3,022	4,114	3,455	4,474	3,340	4,192	_	41,654
Tetanus	_	1	_	_	3	3	3	1	_	3	2	1	_	17
Toxic shock syndrome (other than Streptococcal)	1	3	3	1	3	1	_	6	1	3	1	1	_	24
Trichinellosis	—	_	_	_	_	_	—	—	—	—	_	—	_	
Tuberculosis	397	707	566	477	603	571	510	631	514	712	509	977	_	7,174
Tularemia	_	9	6	3	21	21	20	24	13	17	5	11		150
Vancomycin-intermediate <i>Staphylococcus aureus</i>	4	4	1	1	4	5	6	4	5	2	1	8	_	45
Vancomycin-resistant <i>Staphylococcus</i> <i>aureus</i> *****	1	_	_				_	_		_	_	_	_	1
Varicella morbidity	498	648	290	128	111	167	114	151	190	240	184	206	_	2,927
Varicella mortality	1	_	_	_	_	_	_	_	_	_	_	_	_	1
Vibriosis														
Total	87	117	77	50	101	128	210	391	236	245	99	111	_	1,852
Confirmed	35	44	34	24	50	78	149	272	170	156	61	60	_	1,133
Probable	52	73	43	26	51	50	61	119	66	89	38	51	_	719
Viral hemorrhagic fevers														
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 <sup>+++++</sup>	_	_	_	_	_	_	_	_	_	_	_	_	_	
Zika virus disease, non- congenital	_	_	1	1	1	_	_	_	_	1	_	_		4
Zika virus infection, congenital	_	_	_	_	_	_	_	_	_	_	_	_		_
Zika virus infection, non- congenital	_	4	3	4	1	_	1	1	2	1	_	2		19

-: No reported cases — The reporting jurisdiction did not submit any cases to CDC.

\* Month is defined using MMWR week (https://ndc.services.cdc.gov/wp-content/uploads/2021/02/MMWR\_Week\_overview.pdf). MMWR week calendars can be found at https://ndc.services.cdc.gov/event-codes-other-surveillance-resources/.

+ Candida auris colonization/screening cases are not included in this table. These data are available on the Mycotic Diseases Branch's Tracking Candida auris page (https://www.cdc.gov/fungal/candida-auris/tracking-c-auris.html)

§ Please see Note #11.

¶ Counts include confirmed and probable dengue cases.

\*\* Case counts may include Old World hantavirus infections, such as Seoul virus.

<sup>++</sup> 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, 2020, published online by CDC's Division of Viral Hepatitis, available at https://www.cdc.gov/hepatitis/statistics/SurveillanceRpts.htm. §§ 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.</p>

¶¶ In 2020, the CSTE case definition changed such that cases diagnosed by PCR were classified as confirmed, whereas previously those cases were classified as suspect and did not meet the publication/print criteria.

\*\*\* 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.

<sup>+++</sup> 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 2020, confirmed and probable plague cases are being combined and published. In 2019, only confirmed plague cases were 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.

§§§§ Includes the following categories: primary; secondary; early non-primary non-secondary (includes cases previously reported as early latent); unknown duration or late (includes cases previously reported as late latent syphilis and cases previously reported as late syphilis with clinical manifestations) and congenital syphilis.

¶¶¶¶ 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.

\*\*\*\*\* 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 2020.

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

#### Notes:

- 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 2020 annual tables were officially closed on September 27, 2022.
- 2. The list of national notifiable Infectious diseases and conditions for 2020 and their national surveillance case definitions are available by navigating to the Surveillance Case Definitions | CDC web page, selecting "2020" for the notifiable condition list year, checking "Infectious" conditions, and clicking "Get Notifiable List by Year". This list incorporates the Council of State and Territorial Epidemiologists (CSTE) position statements approved in 2019 by CSTE for national surveillance that were implemented in January 2020. Revised case definitions were implemented for the following conditions; plague, legionellosis, acute hepatitis C, spotted fever rickettsiosis, and pertussis. In addition, CSTE adopted the first coronavirus disease 2019 (COVID-19) national surveillance case definition on April 5, 2020, and they approved a revision to the COVID-19 national surveillance case definition, effective August 5, 2020. Publication criteria for the finalized 2020 data are available at https://wonder.cdc.gov/nndss/documents/2020\_NNDSS\_Publication\_Criteria\_03162022.pdf. See also Guide to Interpreting Provisional and Finalized NNDSS Data.
- 3. Population estimates for incidence rates are July 1st, 2020, estimates obtained from the National Center for Health Statistics (NCHS) postcensal estimates of the resident population of the United States for April 1, 2010, to July 1, 2020, by year, county, single year of age (range: 0 to 85 years), bridged-race (white, black or African American, American Indian or Alaska Native, Asian, or Pacific Islander), Hispanic ethnicity (not Hispanic or Latino, Hispanic or Latino), and sex (Vintage 2020), prepared under a collaborative arrangement with the U.S. Census Bureau. Population estimates for states released September 21, 2021, are available at https://www.cdc.gov/nchs/nvss/bridged\_race/data\_documentation.htm. Population estimates for territories are the 2020 mid-year estimates from the U.S. Census Bureau International Data Base, accessed on March 15, 2022, at https://www.census.gov/data-tools/demo/idb/#/country?YR\_ANIM=2022. 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.
- 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 (once in CDC Stacks, select "Annual Reports" in the "Genre" box to the left).
- 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, denominator is <24 months)

Perinatal hepatitis C infection (age restriction in numerator is ≤36 months, denominator is <36 months).

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 2020 (National Center for Health Statistics Natality 2020, 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).
- 9. The following reporting areas may have incomplete data, due to technical or programmatic challenges while reconciling data during the COVID-19 pandemic: California, Guam, and Minnesota.
- 10. The following reporting areas may have incomplete data due to updates made to their data after the 2020 reconciliation period ended and there was not sufficient time before publication of the annual tables to confirm the updated counts: Idaho, Kansas, Maryland, Vermont, and Virgin Islands.
- 11. Of the reporting areas that submitted 2020 aggregate COVID-19 data to CDC, three did not submit probable cases. New York (excluding New York City) and Utah did not collect probable cases. U.S. Virgin Islands collected probable cases, but did not report them to CDC.
- 12. Disease data presented in the 2020 tables reflect impacts of the COVID-19 pandemic, such as changes in exposure-related behavior, healthcareseeking behavior, disease reporting, and public health investigations.

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• CDC acknowledges the local, state, and territorial health departments that collected the data from a range of case ascertainment sources (e.g., healthcare providers, hospitals, laboratories) and reported these data to CDC's National Notifiable Diseases Surveillance System.

National Notifiable Diseases Surveillance System

Provided by CDC WONDER