



## Maine CDC Infectious Disease Program Quarterly Case Counts: Quarters 1-2, 2020

	Androscoggin	Aroostook	Cumberland	Franklin	Hancock	Kennebec	Knox	Lincoln	Oxford	Penobscot	Piscataquis	Sagadahoc	Somerset	Waldo	Washington	York	Total
2019 Novel Coronavirus (2019-nCoV)	478	23	1714	40	15	145	24	24	35	108	3	34	28	56	3	521	3251
Absenteeism	1	2	5	4	4	3	0	4	3	3	0	1	3	1	2	5	41
Air Outbreak	17	1	55	2	0	4	0	0	0	4	0	1	0	1	0	13	98
<i>Anaplasma phagocytophilum</i>	15	0	20	0	14	15	17	29	14	8	0	7	5	23	0	22	189
Babesiosis	2	0	2	0	0	0	2	2	0	0	0	0	0	1	0	0	9
<i>Borrelia miyamotoi</i>	0	0	1	0	0	1	1	0	0	0	0	0	0	0	0	0	3
CP-CRE, <i>Enterobacter</i> spp	0	0	1	0	0	0	0	0	0	1	0	0	0	0	0	0	2
CP-CRE, <i>Klebsiella</i> spp	0	0	2	0	0	1	0	0	0	0	0	0	0	0	0	1	4
Campylobacteriosis	5	2	8	1	2	7	2	4	2	9	2	0	5	0	1	7	57
Carbapenem-resistant <i>Enterobacteriaceae</i> (CRE)	4	2	11	1	0	9	2	4	1	3	0	2	1	2	0	9	51
<i>Chlamydia trachomatis</i> infection	192	74	403	34	56	155	23	30	48	189	22	32	52	33	40	201	1584
Cryptosporidiosis	0	2	0	0	0	4	0	1	0	7	0	0	4	1	1	0	20
Ehrlichiosis, chaffeensis	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1
Ehrlichiosis/Anaplasmosis, undetermined	0	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1
Emerging Infection	0	0	3	0	0	1	0	0	0	0	0	0	0	0	0	0	4
GI Illness Outbreak	1	0	7	0	1	4	0	0	1	1	0	0	2	0	0	5	22
Giardiasis	2	3	11	1	7	6	4	0	1	7	2	0	3	4	2	5	58
Gonorrhea	36	7	88	3	5	20	2	0	3	15	0	4	6	3	2	28	222
Group A <i>Streptococcus</i> , invasive	2	2	6	0	4	1	0	0	2	8	0	1	0	0	0	10	36
HIV	1	0	5	0	1	1	0	1	1	1	0	0	1	0	0	0	12

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<i>Haemophilus influenzae</i> , invasive	0	1	2	0	1	1	0	0	0	0	0	0	0	0	0	1	6
Hepatitis A, acute	0	0	3	2	2	3	1	0	1	19	0	0	11	4	2	7	55
Hepatitis B, acute	3	1	8	0	0	1	1	0	2	2	0	0	0	0	1	2	21
Hepatitis B, chronic	6	2	28	1	1	5	1	0	5	8	0	0	0	3	1	3	64
Hepatitis C, acute	9	4	22	6	2	10	3	2	2	21	1	0	9	2	2	8	103
Hepatitis C, chronic	60	22	168	20	25	58	30	14	33	92	11	7	26	15	19	94	694
Hepatitis C, perinatal infection	0	0	1	0	0	1	0	0	0	0	0	0	0	0	0	1	3
Hepatitis Delta co- or super-infection, acute	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1
ILI Related Outbreak	4	3	19	4	2	6	2	4	5	5	0	2	2	2	1	14	75
Influenza Associated Pediatric Mortality	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	1
Invasive Pneumococcal Disease	10	5	9	0	0	4	5	2	5	9	2	0	1	2	1	6	61
Legionellosis	0	0	0	0	1	0	0	0	0	1	0	0	0	0	1	0	3
Lyme disease**	16	2	51	2	32	27	37	18	10	30	2	8	9	21	5	41	311
Malaria	0	0	2	0	0	0	0	0	0	0	0	0	0	0	0	0	2
<i>Neisseria meningitidis</i> , invasive (Mening. disease)	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	0	1
Pertussis	1	0	10	0	2	0	1	0	0	0	0	0	1	4	0	10	29
Q fever, Acute	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	1
Rabies PEP	2	0	8	0	2	7	1	5	2	1	0	11	2	0	0	12	53
Rabies, animal	4	1	3	1	0	4	2	1	2	1	0	10	1	3	1	4	38
<i>S. aureus</i> , coag+, meth- or oxi- resistant (MRSA)	3	8	23	1	3	16	6	3	2	17	1	5	4	3	4	14	113
<i>S. aureus</i> , vancomycin intermediate susc (VISA)	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	1

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Salmonellosis (excluding S.typhi and paratyphi)	2	4	6	2	0	2	0	0	1	2	1	1	3	1	2	6	33
Shiga toxin-producing Escherichia coli (STEC)	0	0	1	0	0	0	0	0	0	3	0	0	0	0	0	2	6
Shigellosis	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	1	2
Streptococcal toxic-shock syndrome	1	0	2	0	0	0	0	0	0	1	0	0	0	0	0	2	6
Syphilis, early non-primary, non-secondary	1	0	3	0	0	0	0	0	0	1	0	0	0	0	0	2	7
Syphilis, primary	0	0	1	0	0	0	0	0	0	0	0	0	0	0	0	3	4
Syphilis, secondary	0	0	9	0	0	1	0	0	0	0	0	0	1	0	0	2	13
Syphilis, unknown duration or late	0	0	0	0	0	5	0	0	0	0	0	0	0	0	0	2	7
Tuberculosis	0	0	6	0	1	0	0	0	0	0	0	0	0	0	0	4	11
Varicella (Chickenpox)	1	0	4	1	5	0	3	0	1	3	0	0	1	3	0	2	24
Vibriosis (non-cholera Vibrio species infections)	0	0	2	0	0	0	0	0	1	0	0	0	0	0	0	0	3

This includes investigations initiated during Quarters 1 and 2, MMWR Year 2020 (December 29, 2019 through June 30, 2020). Only investigations with a case status are included in this report. For all outbreak conditions confirmed, probable, and suspect case statuses are included. For non-outbreak conditions confirmed and probable case statuses are included. If a condition is not listed that means no cases met the case definition during this time.

\*Outbreaks that are part of a national cluster are classified as out of state. If the total number does not match the county breakdown, the remaining outbreaks are out of state.

\*\*Lyme disease surveillance is a passive system and can take several months for cases to be counted. These numbers may change dramatically as providers return case report forms and the system is updated.