

Acknowledgements

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Introduction

Foodborne diseases cause an estimated 48 million illnesses each year in the United States, including 9.4 million caused by known pathogens (1-2). Though only a small proportion of these illnesses occur in the setting of an outbreak, data collected during outbreak investigations can provide valuable insight into the pathogens and foods that cause illness. Public health officials, regulatory agencies, and the food industry can use this information to create targeted control strategies along the farm-to-table continuum to address specific pathogens and foods.

A foodborne disease outbreak is defined as the occurrence of two or more cases of a similar illness resulting from ingestion of a common food. Foodborne disease outbreaks are a nationally notifiable condition (http://c. ymcdn.com/sites/www.cste.org/resource/resmgr/ CSTENotifiableConditionListA.pdf). CDC conducts foodborne disease outbreak surveillance in the United States through the Foodborne Disease Outbreak Surveillance System. Public health agencies in all 50 states, the District of Columbia, U.S. territories, and Freely Associated States voluntarily submit reports of outbreaks investigated by their agencies using a web-based reporting platform, the National Outbreak Reporting System (NORS) (http://www.cdc.gov/nors/). NORS also collects reports of enteric disease outbreaks caused by other modes of transmission, including person-toperson contact, animal contact, water, environmental contamination, and unknown mode of transmission.

Investigating and reporting agencies use a standard outbreak reporting form (http://www.cdc.gov/nors/pdf/NORS_CDC_5213.pdf) to report foodborne disease outbreaks. Data requested for each outbreak include the reporting state; date of first illness onset; number of illnesses, hospitalizations and deaths; etiology; implicated food and ingredients; locations of food preparation; and factors contributing to food contamination (see appendix). Outbreaks that are excluded from the Foodborne Disease Outbreak Surveillance System include those that occurred on

cruise ships with both US and international ports and those in which the food was eaten outside the United States, even if the illness occurred in the United States.

This report includes foodborne disease outbreaks that were reported to the Foodborne Disease Outbreak Surveillance System by October 17, 2013, in which the first illness occurred during 2012. Etiologic agents were reported as confirmed if they met pre-defined confirmation criteria (3); otherwise, they were reported as suspected. For outbreaks caused by a single confirmed or suspected etiology, etiologies were grouped as bacterial, chemical and toxin, parasitic, or viral. Multistate outbreaks were defined as outbreaks in which exposure to the implicated food occurred in more than one state or territory. Population-based outbreak reporting rates were calculated for each state using United States Census estimates of the 2011 state populations (http://www.census.gov/popest); multistate outbreaks were included in population-based outbreak reporting rates by assigning one outbreak to each state that reported cases in the outbreak. Implicated foods were classified into one of 24 single food categories if a single contaminated ingredient was identified or if all ingredients belonged to that category (4). Outbreaks attributed to foods that could not be assigned to one of these single food categories, or for which the report contained insufficient information for category assignment, were not attributed to any category.

Findings: Foodborne disease outbreaks, United States, 2012

During 2012, 831 foodborne disease outbreaks were reported (Table 1), resulting in 14,972 illnesses, 794 hospitalizations, and 23 deaths (Table 1). Outbreaks were reported by public health officials from all 50 states, the District of Colombia, and Puerto Rico (Figure 1). The median rate was 3.4 foodborne disease outbreaks per 1 million population; rates ranged from 1.0 outbreak per 1 million population in Mississippi to 11.4 outbreaks per 1 million population in Rhode Island.

A single confirmed or suspected etiologic agent was identified in 579 (69%) outbreaks (423 confirmed and 156 suspected) (Table 1). Among the 423 outbreaks with a single confirmed etiologic agent, bacteria caused the most outbreaks (208 outbreaks, 49%), followed by viruses (175, 41%), chemicals and toxins (36, 9%), and parasites (4, 1%). Norovirus was the most common cause of confirmed, single-etiology outbreaks and illnesses, accounting for 172 (41%) outbreaks and 4,679 (45%) illnesses. Salmonella was next, accounting for 106 (25%) outbreaks and 3,366 (33%) illnesses. Among the 101 confirmed Salmonella outbreaks with a serotype reported, Enteritidis was the most common (26 outbreaks, 26%), followed by Typhimurium (13, 13%), Newport (10, 10%), Javiana (7, 7%), and Heidelberg (6, 6%). Shiga toxin-producing *Escherichia coli* (STEC) caused 29 confirmed, single-etiology outbreaks, of which 24 (83%) were caused by serogroup O157; 2 outbreaks were caused by serogroup O145 and 1 outbreak each by O45, O111, and O121.

Of the 10,319 outbreak-related illnesses caused by a single confirmed etiologic agent, 701 (7%) resulted in hospitalization (Table 1). Among confirmed, single-etiology outbreaks, *Salmonella* caused the most outbreak-related hospitalizations (449 hospitalizations, 64%), followed by STEC (98, 14%) and norovirus (46, 7%). Outbreaks caused by *Clostridium botulinum* resulted in the highest proportion of ill persons hospitalized (100%). Among the 23 deaths reported, 17 (74%) were attributed to bacterial etiologies (*Listeria monocytogenes* [6], *Campylobacter* [4], STEC serogroups O157 [3] and O145 [1], *Salmonella* [2], and *Clostridium botulinum* [1]); 4 deaths were attributed to mycotoxins, 1 to another chemical or toxin, and 1 to norovirus.

A food vehicle was reported for 334 (40%) outbreaks. In 192 (57%) of these outbreaks, the food vehicle could be classified into one of the 24 single food categories (Table 2a); the categories most commonly implicated were fish (31 outbreaks, 16%), vegetable row crops (23, 12%), and dairy (19, 10%). Among the 17 dairy-associated outbreaks for which pasteurization information was reported,

12 (71%) involved unpasteurized products. The categories associated with the most outbreak-related illnesses were fruits (858 illnesses, 21%), fish (515, 12%), and chicken (509, 12%).

The pathogen-food category pairs responsible for the most outbreaks were scombroid toxin (histamine fish poisoning) in fish (17 outbreaks), Campylobacter in unpasteurized dairy (10), and Vibrio parahaemolyticus in mollusks (10) (Table 2b). The pathogen-food category pairs responsible for the most outbreak-related illnesses were Salmonella in fruits (446 illnesses), Salmonella in fish (425), and Salmonella in chicken (345). The pathogen-food category pairs responsible for the most hospitalizations were Salmonella in chicken (109 hospitalizations), Salmonella in fruits (55), and Salmonella in fish (55). Deaths were reported for the following pathogen-food category pairs: Listeria in dairy (5 deaths) and sprouts (1), Campylobacter in chicken (4), STEC in vegetable row crops (1), Salmonella in eggs (1) and fruits (1), other chemical or toxin in fish (1), Clostridium botulinum in root or other underground vegetable (1).

Twenty-four multistate outbreaks (3% of all outbreaks) were reported. They involved a median of 8 states (range: 3 – 23). Fifteen outbreaks were caused by Salmonella. Reported Salmonella serotypes were Typhimurium (4 outbreaks); Javiana, Newport, and multiple serotypes (2 each); Bredeney, Cubana, Enteritidis, Heidelberg, and Schwarzengrund (1 each). The remaining 9 outbreaks were caused by STEC (5 outbreaks; serogroups O157 [2], O45 [1], O121 [1], and O145 [1]), Listeria (2), Campylobacter (1), and Vibrio parahaemolyticus (1). Fourteen (93%) of 15 multistate outbreaks caused by Salmonella implicated a food vehicle; the foods were cantaloupes (4 outbreaks); ground beef and chicken (2 each); cucumbers, mangos, peanut butter, sprouts, romaine lettuce, and tuna (1 each). In the fifteenth, a food vehicle was not identified, but food was consumed at restaurants of a Mexican-style chain. Five multistate outbreaks were caused by STEC; it was transmitted in romaine lettuce (serogroup O157 [1]), prepackaged leafy greens (O157 [1]), an unspecified type of lettuce (O145 [1]),

frozen meals (O121 [1]), and sandwiches (O45 [1]). Two multistate outbreaks were caused by *Listeria* in pasteurized ricotta salata cheese and sprouts (1 outbreak each). Chicken livers were implicated in the *Campylobacter* outbreak and oysters were implicated in the *Vibrio parahaemolyticus* outbreak.

Among the 719 outbreaks and 12,657 illnesses with a reported single location where food was prepared, 433 outbreaks (60%) and 5,174 illnesses (41%) were associated with foods prepared in a restaurant (Table 3a). Among these outbreaks, sit-down dining-style was the type of restaurant most commonly reported as the location where food was prepared (358 outbreaks, 83%).

Twenty outbreaks resulted in product recalls; the foods recalled were ground beef (3 outbreaks); oysters and unpasteurized milk (2 each); multiple types of shellfish, raw tuna scrape and strips, pasteurized ricotta salata cheese, pasteurized milk and chocolate products, hog head cheese, pre-packaged leafy greens, cantaloupes, mangos, multiple fruits, peanut butter and other peanut products, tempeh, unpasteurized apple cider, and frozen meals (1 each).

Limitations

The findings in this report are subject to at least four limitations. First, only a small proportion of foodborne illnesses reported each year are identified as being associated with outbreaks. The extent to which the distribution of food vehicles and locations of preparation implicated in foodborne disease outbreaks reflect the same vehicles and locations as sporadic foodborne illnesses is unknown. Similarly, not all outbreaks are identified, investigated, or reported. Second, many outbreaks had an unknown etiology, an unknown food vehicle, or both, and conclusions drawn from outbreaks with a confirmed or suspected

etiology or food vehicle might not apply to outbreaks with an unknown etiology or food vehicle. Third, CDC's outbreak surveillance system is dynamic; agencies can submit new reports and can change or delete previous reports as new information becomes available. Therefore, the results of this analysis might differ from those published earlier or from future reports. Finally, because of changes in the surveillance system implemented in 2009 and the use of a new food categorization scheme in 2011, comparisons with preceding years should be made with caution.

Main Findings and Comments

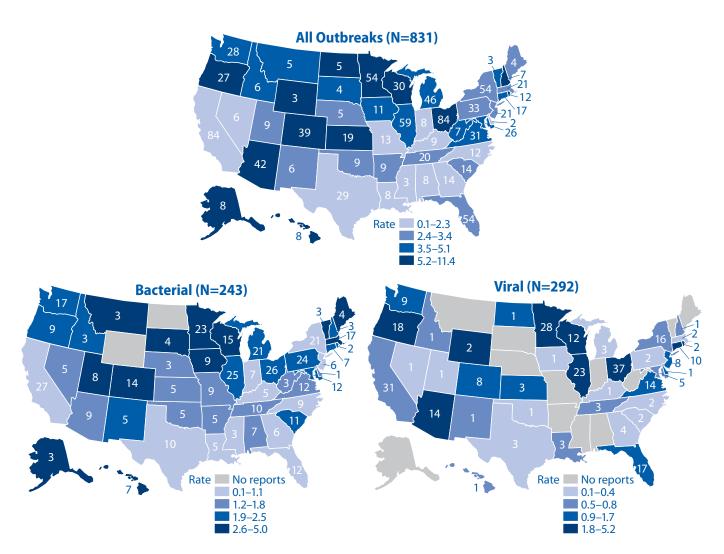
- In 2012, 831 foodborne disease outbreaks were reported, resulting in 14,972 illnesses, 794 hospitalizations, 23 deaths, and 20 food recalls.
- The number of outbreaks caused by Campylobacter increased from 25 in 2010 to 30 in 2011 and 37 in 2012. Among the 19 Campylobacter outbreaks with a known food vehicle in 2012, 10 (53%) were attributed to unpasteurized dairy products.
- Fish, vegetable row crops, and dairy were the most commonly implicated single food categories in outbreaks in which a single food category could be implicated.
- Restaurants, specifically sit-down dining-style restaurants, were most commonly reported as the location of food preparation.

Public health, regulatory, and food industry professionals use foodborne disease outbreak surveillance data to target prevention efforts related to pathogens and foods that cause foodborne disease outbreaks. Additional information on outbreaks and the Foodborne Disease Outbreak Online Database are available at http://www.cdc.gov/foodsafety/fdoss/.

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Figure: Rate of reported foodborne disease outbreaks per 1 million population* and number of outbreaks,† by state‡ and major etiology group§—Foodborne Disease Outbreak Surveillance System, United States, 2012



^{*} Cutpoints for outbreak rate categories determined using quartiles. Legend differs for each map.

[†] Number of reported outbreaks in each state. In addition to the 50 states, Puerto Rico reported 14 outbreaks and the District of Columbia reported 1 outbreak.

[‡] Includes 24 multistate outbreaks (i.e., outbreaks in which exposure occurred in more than one state) assigned as an outbreak to each state involved. Multistate outbreaks involved a median of 8 states (range: 3 – 23).

[§] Includes outbreaks caused by both confirmed and suspected etiologies.

Table 1: Number of reported foodborne disease outbreaks, outbreak-associated illnesses, and hospitalizations, by etiology (confirmed or suspected)*—Foodborne Disease Outbreak Surveillance System, United States, 2012

		No. Ou	tbreaks			No. III	nesses		N	o. Hospi	talization	IS
Etiology	CE	SE	Total	%	CE	SE	Total	%	CE	SE	Total	%
Bacterial												
Salmonella†	106	7	113	20	3,366	28	3,394	28	449	5	454	61
Campylobacter‡	30	7	37	6	434	42	476	4	27	2	29	4
Escherichia coli, Shiga toxin-producing (STEC)§	29	0	29	5	500	0	500	4	98	0	98	13
Clostridium perfringens	18	7	25	4	989	73	1,062	9	4	0	4	1
Vibrio parahaemolyticus	8	3	11	2	54	12	66	1	4	1	5	1
Clostridium botulinum	5	1	6	1	19	2	21	0	19	2	21	3
Listeria monocytogenes	4	1	5	1	19	23	42	0	17	21	38	5
Staphylococcus aureus, enterotoxin	2	3	5	1	26	123	149	1	1	3	4	1
Bacillus cereus	1	1	2	0	10	14	24	0	0	0	0	0
Shigella¶	2	0	2	0	18	0	18	0	3	0	3	0
Yersinia enterocolitica	1	1	2	0	2	4	6	0	0	0	0	0
Streptococcus, Group A	1	0	1	0	18	0	18	0	0	0	0	0
Vibrio other	1	0	1	0	3	0	3	0	1	0	1	0
Other	0	4	4	1	0	12	12	0	0	0	0	0
Subtotal	208	35	243	42	5,458	333	5,791	48	623	34	657	88
Chemical and toxin												
Scombroid toxin/Histamine	18	1	19	3	53	2	55	0	0	0	0	0
Ciguatoxin	7	1	8	1	17	3	20	0	6	3	9	1
Mycotoxins	6	0	6	1	21	0	21	0	17	0	17	2
Paralytic shellfish poison	2	0	2	0	11	0	11	0	4	0	4	1
Other	3	2	5	1	13	11	24	0	2	0	2	0
Subtotal	36	4	40	7	115	16	131	1	29	3	32	4
Parasitic												
Giardia	2	0	2	0	7	0	7	0	0	0	2	0
Trichinella	2	0	2	0	4	0	4	0	2	0	2	0
Subtotal	4	0	4	1	11	0	11	0	2	0	2	0
Viral												
Norovirus	172	115	287	50	4,679	1,330	6,009	50	46	12	58	8
Sapovirus	3	0	3	1	56	0	56	0	1	0	1	0
Astrovirus	0	1	1	0	0	22	22	0	0	0	0	0
Other	0	1	1	0	0	9	9	0	0	0	0	0
Subtotal	175	117	292	50	4,735	1,361	6,096	51	47	12	59	8
Single etiology**	423	156	579	70	10,319	1,710	12,029	80	701	49	750	94
Multiple etiologies	2	5	7	1	77	35	112	1	10	0	10	1
Unknown etiology ^{††}	0	245	245	29	0	2,831	2,831	19	0	34	34	4
Total**	425	406	831	100	10,396	4,576	14,972	100	711	83	794	100

Abbreviations: No. = Number; CE = confirmed etiology; SE = suspected etiology.

^{*}If at least one etiology was laboratory-confirmed, the outbreak was considered to have a confirmed etiology. If no etiology was laboratory-confirmed, but an etiology was reported based on clinical or epidemiologic features, the outbreak was considered to have a suspected etiology.

[†] Salmonella serotypes causing more than five outbreaks were Enteritidis (27 outbreaks), Typhimurium (13), Newport (10), Javiana (7), and Heidelberg (6).

[‡] Campylobacter jejuni (25 outbreaks), Campylobacter multiple species (2), Campylobacter other species (1), and Campylobacter unknown species (8).

^{\$}STEC serogroups O157 (24 outbreaks; 23 were H7 and 1 was nonmotile), O145 (2), O45 (1), O111:nonmotile (1), and O121 (1).

[¶] Shigella sonnei (2 outbreaks).

^{**}The denominator for the etiology percentages is the single etiology total. The denominator for the single etiology, multiple etiologies, and unknown etiology is the total. Because of rounding, numbers might not add up to the single etiology total or the total.

 $^{\ \, \}dagger\dagger\ \, \text{An etiologic agent was not confirmed or suspected based on clinical, laboratory, or epidemiologic information.}$

Table 2a: Number of reported foodborne disease outbreaks and outbreak-associated illnesses, by food category*—Foodborne Disease Outbreak Surveillance System, United States, 2012

	No. Ou	tbreaks	No. IIIr	nesses
Food Category*	Total	%	Total	%
Aquatic animals				
Crustaceans	1	1	3	0
Mollusks [†]	13	7	83	2
Fish	31	16	515	12
Other aquatic animals	0	0	0	0
Subtotal	45	23	601	15
Land animals				
Dairy [‡]	19	10	356	9
Eggs	8	4	85	2
Beef	11	6	232	6
Pork	10	5	340	8
Other meat (sheep, goat, etc.)	0	0	0	0
Chicken	18	9	509	12
Turkey	8	4	202	5
Other poultry	2	1	7	0
Game	2	1	5	0
Subtotal	79	41	1,736	42
Plants				
Oils and sugars	1	1	7	0
Fungi	5	3	15	0
Sprouts	2	1	25	1
Root and other underground vegetables§	5	3	34	1
Seeded vegetables ¹	3	2	206	5
Herbs	0	0	0	0
Vegetable row crops**	23	12	377	9
Fruits ^{††}	16	8	858	21
Grains and beans#	8	4	190	5
Nuts and seeds ^{§§}	1	1	42	1
Subtotal	63	33	1754	42
Other	5	3	49	1
Food reported, attributed to a single food category ¹¹	192	23	4,140	28
Food reported, not attributed to a single food category	142	17	3,212	21
No food reported	497	60	7,620	51
Total**	831	100	14,972	100

$\textbf{Abbreviations:} \ \mathsf{No.} = \mathsf{Number.}$

§§Nuts (1 outbreak).

^{*}Interagency Food Safety Analytics Collaboration (IFSAC) food categorization scheme: http://www.cdc.gov/foodsafety/ifsac/projects/completed.html †Bi-valve mollusks (13 outbreaks).

[‡]Unpasteurized dairy products (12 outbreaks), pasteurized dairy products (5), and pasteurization unknown (2).

[§]Tubers (4 outbreaks) and root vegetables (1).

[¶]Solanaceous seeded vegetables (2 outbreaks) and vine-grown seeded vegetables (1).

^{**}Leafy vegetables (20 outbreaks), stem vegetables (1), and vegetable row crops not further classified (1).

^{††}Melons (4 outbreaks), small fruits (2), tropical fruits (1), and fruits not further classified (9).

^{##}Grains (4 outbreaks), beans (2), and grains-beans not further classified (2).

^{¶¶}The denominator for the food category percentages is the "food reported, attributed to a single food category" total. The denominator for the "food reported, attributed to a single food category", "food reported, not attributed to a single food category", and "No food reported" is the total. Because of rounding, numbers might not add up to the "food reported, attributed to a single food category" total or the total.

Table 2b: Most common pathogen-food category pairs resulting in outbreaks, outbreaks-associated illnesses, hospitalizations, and deaths—Foodborne Disease Outbreak Surveillance System, United States, 2012

Top 5 pathogen-food category pairs resu	ılting in outbreaks				
Etiology	Food Category*	No. Outbreaks	No. Illnesses	No. Hospitalizations	No. Deaths
Scombroid toxin/Histamine	Fish	17	50	0	0
Campylobacter	Dairy	10	201	13	0
Vibrio parahaemolyticus	Mollusks	10	60	5	0
Salmonella	Chicken	9	345	109	0
Norovirus	Vegetable row crops	8	154	2	0
Escherichia coli, Shiga toxin-producing (STEC)	Vegetable row crops	8	136	34	1
Salmonella	Eggs	7	58	11	1
Ciguatoxin	Fish	7	18	9	0

Top 5 pathogen-food category pa	irs resulting in outbreak-	-associated illnes	ses		
Etiology	Food Category*	No. Outbreaks	No. Illnesses	No. Hospitalizations	No. Deaths
Salmonella	Fruits	6	446	55	1
Salmonella	Fish	1	425	55	0
Salmonella	Chicken	9	345	109	0
Clostridium perfringens	Pork	4	286	1	0
Norovirus	Fruits	4	210	0	0

Top 5 pathogen-food category pairs resu	ılting in outbreak-a	issociated hospi	talizations		
Etiology	Food Category*	No. Outbreaks	No. Illnesses	No. Hospitalizations	No. Deaths
Salmonella	Chicken	9	345	109	0
Salmonella	Fruits	6	446	55	1
Salmonella	Fish	1	425	55	0
Salmonella	Seeded vegetables	2	151	50	0
Escherichia coli, Shiga toxin-producing (STEC)	Vegetable row crops	8	136	34	1
Listeria monocytogenes	Dairy	1	23	21	5

Pathogen-food category pairs resulting	in outbreak-associ	ated deaths			
Etiology	Food Category*	No. Outbreaks	No. Illnesses	No. Hospitalizations	No. Deaths
Listeria monocytogenes	Dairy	1	23	21	5
Campylobacter	Chicken	4	37	3	4
Escherichia coli, Shiga toxin-producing (STEC)	Vegetable row crops	8	136	34	1
Salmonella	Eggs	7	58	11	1
Salmonella	Fruits	6	446	55	1
Other chemical or toxin	Fish	3	13	2	1
Listeria monocytogenes	Sprouts	1	6	4	1
Clostridium botulinum	Root and other underground vegetables	1	3	3	1

 $\textbf{Abbreviations:} \ \mathsf{No.} = \mathsf{Number.}$

^{*} Interagency Food Safety Analytics Collaboration (IFSAC) food categorization scheme: http://www.cdc.gov/foodsafety/ifsac/projects/completed.html.

Table 3a: Number of reported foodborne disease outbreaks and outbreak-associated illnesses, by location of food preparation—Foodborne Disease Outbreak Surveillance System, United States, 2012

	No. Ou	tbreaks	No. Illr	nesses
Location of food preparation	Total	%	Total	%
Restaurant	433	60	5,174	41
Sit-down dining	358	50	4,518	36
Fast-food	53	7	425	3
Other or unknown type	20	3	204	2
Multiple types	2	0	27	0
Catering or banquet facility	100	14	3,000	24
Private home	90	13	1,577	12
Other institutional location	31	4	1,497	12
School	12	2	211	2
Prison or jail	11	2	1,119	9
Workplace cafeteria	4	1	59	0
Camp	2	0	44	0
Day care	1	0	54	0
Workplace, not cafeteria	1	0	10	0
Other location	26	4	466	4
Other commercial location	22	3	372	3
Grocery store	13	2	154	1
Fair, festival, or temporary mobile service	5	1	76	1
Farm or dairy	4	1	142	1
Hospital or nursing home	11	2	201	2
Nursing home	8	1	114	1
Hospital	3	0	87	1
Other private location	6	1	370	3
Place of worship	6	1	370	3
Single location*	719	87	12,657	85
Multiple locations	34	4	1,093	7
Unknown location	78	9	1,222	8
Total*	831	100	14,972	100

 $\textbf{Abbreviations:} \ \mathsf{No.} = \mathsf{Number.}$

^{*}The denominator for the location percentages is the single location total. The denominator for the single location, multiple locations, and unknown location is the total. Because of rounding, numbers might not add up to the single location total or the total.

Table 3b: Number of reported foodborne disease outbreaks and outbreak-associated illnesses, by etiology (confirmed and suspected)* and location of food preparation[†]—Foodborne Disease Outbreak Surveillance System, United States, 2012

	ban	ing or quet ility	Resta	aurant	comn	her iercial tion	nur	ital or sing me	institu	her utional ation		vate ome	pri	her vate ation		her ition
Etiology	NO	NI	NO	NI	NO	NI	NO	NI	NO	NI	NO	NI	NO	NI	NO	NI
Bacterial																
Salmonella	3	131	40	536	5	68	2	24	4	596	26	1,000	1	33	6	139
Campylobacter	4	99	9	63	4	142	‡	_	2	10	8	55	1	3	2	40
Escherichia coli, Shiga toxin-producing (STEC)	_	_	7	59	1	17	_	_	1	8	7	101	_	_	1	16
Clostridium perfringens	10	514	7	44	1	20	1	6	1	108	2	91	2	270	1	9
Vibrio parahaemolyticus	_	_	7	26	_	_	_	_	_	_	3	12	_	_	_	_
Clostridium botulinum	_	_	_	_	_	_	_	_	2	12	3	7	_	_	1	2
Listeria monocytogenes	_	_	1	3	_	_	1	7	_	_	_	_	_	_	_	_
Staphylococcus aureus, enterotoxin	2	119	2	24	_	_	_	_	_	_	1	6	_	_	_	_
Bacillus cereus	_	_	2	24	_	_	_	_	_	_	_	_	_	_	_	_
Shigella	_	_	1	11	_	_	_	_	1	7	_	_		_		_
Yersinia enterocolitica	_	_	_	_	_	_	_	_	_	_	2	6	_	_	_	_
Streptococcus, Group A	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_
Vibrio other	_	_	_	_	_	_	_	_	_	_	1	3	_	_	_	_
Other	_	_	4	12	_	_	_	_	_	_	_	_	_	_	_	_
Subtotal	19	863	80	802	11	247	4	37	11	741	53	1,272	4	306	11	206
Chemical and toxin																
Scombroid toxin/Histamine	_	_	15	43	_	_	_	_	_	_	1	2	_	_	3	10
Ciguatoxin	_	_	2	5	_	_	_	_	_	_	6	15	_	_	_	_
Mycotoxins	_	_	_	_	_	_	1	6	_	_	4	13	_	_	_	_
Paralytic shellfish poison	_	_	_	_	_	_	_	_	_	_	1	7	_	_	_	_
Other	_	_	1	6	_	_	_	_	1	5	1	2	_	_	2	11
Subtotal	_	_	18	54	_	_	1	6	1	5	13	39	_	_	5	21
Parasitic																
Giardia	_	_	_	_	1	4	_	_	_	_	_	_	_	_	1	3
Trichinella	_	_	1	2	_	_	_	_	1	2	_	_	_	_	_	_
Subtotal	_	_	1	2	1	4	_	_	1	2	_	_	_	_	1	3
Viral																
Norovirus	51	1,497	184	3,229	1	10	4	73	12	527	9	99	1	25	3	97
Sapovirus	1	9	1	22	_	_	_	_	_	_	_	_	_	_	1	25
Astrovirus	1	22	_	_	_	_	_	_	_	_	_	_	_	_	_	_
Other	_		_	_	_	_	_	_	_	_	_		_	_	_	_
Subtotal	53	1,528	185	3,251	1	10	4	73	12	527	9	99	1	25	4	122
Single etiology	72	2,391	284	4,109	13	261	9	116	25	1,275	75	1,419	5	331	21	352
Multiple etiologies	_	_	4	24	_	_	_	_	_	_	1	21	_	_	_	_
Unknown etiology [§]	28	609	145	1,041	9	111	2	85	6	222	14	137	1	39	5	114
Total	100	3,000	433	5,174	22	372	11	201	31	1,497	90	1,577	6	370	26	466

Abbreviations: NO = number of outbreaks; NI = number of illnesses.

^{*}If at least one etiology was laboratory-confirmed, the outbreak was considered to have a confirmed etiology. If no etiology was laboratory-confirmed, but an etiology was reported based on clinical or epidemiologic features, the outbreak was considered to have a suspected etiology.

[†]Reported locations were grouped as follows: catering or banquet facility, restaurant, other commercial location, hospital or nursing home, other institutional location, private home, other private location, and other location (see Table 3a).

^{*}No outbreaks in the data reported fall into this category.

[§]An etiologic agent was not confirmed or suspected based on clinical, laboratory, or epidemiologic information.

Appendix: Number of reported foodborne disease outbreaks, by etiology (confirmed and suspected)* and contributing factors†—Foodborne Disease Outbreak Surveillance System, United States, 2012

Etiology Bacterial Salmonella Campylobacter Escherichia coli, Shiga toxin-producing (STEC) Clostridium perfringens Vibrio parahaemolyticus Clostridium botulinum Listeria monocytogenes Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other Other	* *	C2				19	4				C11	C12	C13	C14	C15	≥1 factor reported	with reported contributing factors	Total No. outbreaks
Salmonella Campylobacter Escherichia coli, Shiga toxin-producing (STEC) Clostridium perfringens Vibrio parahaemolyticus Clostridium botulinum Listeria monocytogenes Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other	_# 		_ _ _ _	_ _ _	_ _ _			_	9	_								
Campylobacter Escherichia coli, Shiga toxin-producing (STEC) Clostridium perfringens Vibrio parahaemolyticus Clostridium botulinum Listeria monocytogenes Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other	* 		_ _ _ _	_ _ _	_ _ _			_	9	_								
Escherichia coli, Shiga toxin-producing (STEC) Clostridium perfringens Vibrio parahaemolyticus Clostridium botulinum Listeria monocytogenes Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other		_ _ _ _	_ _ _ _	_ _ _	_ _	5	0		,	/	2	3	—	—	3	36	44	113
Clostridium perfringens Vibrio parahaemolyticus Clostridium botulinum Listeria monocytogenes Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other		_ _ _ _	_ _ _	_	_		9	_	4	_	_	_	_	1	_	17	20	37
Vibrio parahaemolyticus Clostridium botulinum Listeria monocytogenes Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other	_ _ _ _	_ _ _	_	_		2	7	_	1	_	_	—	—	—	_	10	10	29
Clostridium botulinum Listeria monocytogenes Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other	_ _ _	_	_		_	2	_	_	1	_	_	_	_	_	5	6	20	25
Listeria monocytogenes Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other	_ 	_		_	_	—	8	_	1	_	_	—	—	_	_	8	8	11
Staphylococcus aureus, enterotoxin Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other	_		_	_	_	1	1	_	_	_	_	_	_	_	_	2	6	6
Bacillus cereus Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other	_	_	_	_	_	—	_	_	1	_	_	—	—	—	1	2	2	5
Shigella Yersinia enterocolitica Streptococcus, Group A Vibrio other		_	_	_	_	_	_	_	_	2	_	2	_	_	_	3	3	5
Yersinia enterocolitica Streptococcus, Group A Vibrio other	_	_	_	_	_	_	_	_	1	_	_	—	_	_	_	1	2	2
Streptococcus, Group A Vibrio other	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	2
Vibrio other	1	_	_	_	_	_	_	_	_	_	_	—	_	_	_	1	1	2
	_	_	_	_	_	_	_	_	_	1	1	_	_	_	_	1	1	1
Other	_	_	_	_	_	_	1	_	_	_	_	_	_	_	_	1	1	1
Other	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	2	4
Subtotal	1	_	_	_	_	30	30	_	19	10	3	5	_	1	9	89	121	243
Chemical and toxin																		
Scombroid toxin/Histamine	9	_	1	_	_	1	3	_	_	_	_	_	_	_	_	13	15	19
Ciguatoxin	7	_	_	_	_	_	_	_	_	_	_	_	_	_	_	7	7	8
Mycotoxins	5	_	_	_	_	_	_	_	_	_	_	—	_	_	_	5	5	6
Paralytic shellfish poison	1	_	_	_	_	_	_	1	_	_	_	_	_	_	_	1	1	2
Other	2	_	1	1	_	_	_	_	_	_	_	_	_	_	1	4	5	5
Subtotal	24	_	2	1	_	1	3	1	_	_	_	_	_	_	1	30	33	40
Parasitic																		
Giardia	_	_	_	_	_	_	1	_	_	_	_	_	_	_	_	1	1	2
Trichinella	1	_	_	_	_	1	_	_	_	_	_	_	_	_	_	2	2	2
Subtotal	1	_	_	_	_	1	1	_	_	_	_	_	_	_	_	3	3	4
Viral							_											
Norovirus	_	_	_	_	_	_	_	_	7	55	33	34	8	6	11	113	116	287
Sapovirus	_	_	_				_	_	_	_	_	1	_			1	1	3
Astrovirus	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	_	1
Other		_	_				_		_			_	_			_	_	1
Subtotal	_	_	_	_	_	_	_	_	7	55	33	35	8	6	11	114	117	292
Single etiology	26		2	1	_	31	34	1								235	274	579
Multiple etiologies	_	_	_	_	_	1	_	_	_	_	_	_	_	_	_	1	3	7
Unknown etiology ¹																I		
Total	2	_	_	_	_	5	3	_	6	16	9	8	1	6	6	44	68	245

Appendices' footnotes are on page 12.

Appendix: Number of reported foodborne disease outbreaks, by etiology (confirmed and suspected)* and contributing factors†—Foodborne Disease Outbreak Surveillance System, United States, 2012

P1 P2 P3 P4 P5 P6 P7 P8 P9 P10 P11 P12 P14 P15 P	
Salmonella	37 29 25 11 6 5 5 2 2 2 2 1 1 4
Campylobacter	37 29 25 11 6 5 5 2 2 2 2 1 1 4
Escherichia coli, Shiga toxin-producing (STEC)	29 25 11 6 5 5 2 2 2 1 1
Clostridium perfringens	25 11 6 5 5 2 2 2 1 1 4
Vibrio parahaemolyticus	11 6 5 5 2 2 2 2 1 1 4
Clostridium botulinum	6 5 5 2 2 2 2 1 1 4
Listeria monocytogenes	5 5 2 2 2 2 1 1 4
Staphylococcus aureus, enterotoxin 3	5 2 2 2 1 1 4
Bacillus cereus	2 2 2 1 1 4
Shigella	2 2 1 1 4
Yersinia enterocolitica 1 — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — —	2 1 1 4
Streptococcus, Group A 1 — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — <td>1 1 4</td>	1 1 4
Vibrio other — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — — <t< td=""><td>1 4</td></t<>	1 4
Other 1 1 1 1 1 1 1 1 1 1 1 1 2 1 8 1 1 5 12 73 121 Chemical and toxin Scombroid toxin/Histamine 1 1 1 1 - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - - -	4
Subtotal 37 15 6 3 15 2 15 8 1 1 5 12 73 121 Chemical and toxin 1 1 1 1 2 5 15 15 Ciguatoxin 2 5 5 15 Mycotoxins 2 2 5 5 15 Paralytic shellfish poison 1 - 1 5 5 Subtotal 1 1 1 1 2 1 2 8 33 Parasitic	
Chemical and toxin Scombroid toxin/Histamine 1 1 1 2 5 15 Ciguatoxin 2 2 5 Mycotoxins 2 2 5 Paralytic shellfish poison 1 - 1 5 Other 1 2 8 33 Parasitic Giardia	243
Scombroid toxin/Histamine 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 <td></td>	
Ciguatoxin — — — — — — — — — — — — — — — — — — —	
Mycotoxins — — — — — — — — 2 — — — — 2 5 Paralytic shellfish poison — — — — — — — — — — — — — — — — — — —	19
Paralytic shellfish poison — — — — — — — — — — — — — — — — — — —	8
Other — — — — — — — — — 1 1 5 Subtotal 1 1 1 — — — 1 2 8 33 Parasitic Giardia — — — — — — — — — — — — — — 1 1 1 1	6
Other — — — — — — — — — 1 1 5 Subtotal 1 1 1 — — — 1 2 8 33 Parasitic Giardia — — — — — — — — — — — — — — 1 1 1 1	2
Parasitic — — — — — — — — — — 1	5
Giardia — — — — — — — — 1	40
	2
Trichinella	2
Subtotal	4
Viral	
Norovirus 2 — — 1 3 — 2 1 — — — 2 6 116	287
Sapovirus	3
Astrovirus	1
Other	1
Subtotal 2 — — 1 3 — 2 1 — — — 2 6 117	292
Single etiology 40 16 7 4 18 2 17 11 1 1 6 16 87 274	579
Multiple etiologies 1 1 - 1 2 - - 2 3	7
Unknown etiology ¹ 8 11 4 6 16 1 14 13 3 — 2 43 68	245
Total 49 28 11 10 35 3 32 26 4 1 6 18 132 344	

Appendices' footnotes are on page 14.

Appendix: Number of reported foodborne disease outbreaks, by etiology (confirmed and suspected)* and contributing factors†—Foodborne Disease Outbreak Surveillance System, United States, 2011.

	Survival Factors‡						No. outbreaks with reported	Total No.
Etiology	S 1	S2	S 3	S4	S 5	≥1 factor reported	contributing factors	outbreaks
Bacterial								
Salmonella	13	1	‡	2	7	21	44	113
Campylobacter	4	_	_	_	2	6	20	37
Escherichia coli, Shiga toxin-producing (STEC)	1	_	_	_	1	2	10	29
Clostridium perfringens	3	7	_	_	2	10	20	25
Vibrio parahaemolyticus	_	_	_	_	_	_	8	11
Clostridium botulinum	1	2	_	1	3	5	6	6
Listeria monocytogenes	_	_	_	_	_	_	2	5
Staphylococcus aureus, enterotoxin	_	2	_	_	_	2	3	5
Bacillus cereus	_	_	_	_	_	_	2	2
Shigella	_	_	_	_	_	_	_	2
Yersinia enterocolitica	1	_	_	_	_	1	1	2
Streptococcus, Group A	_	_	_	_	1	1	1	1
Vibrio other	_	_	_	_	_	_	1	1
Other	_	_	_	_	_	_	2	4
Subtotal	23	12	_	3	16	48	121	243
Chemical and toxin					1			
Scombroid toxin/Histamine	_	_	1	_	_	1	15	19
Ciguatoxin	_	_	_	_	_	_	7	8
Mycotoxins	_	_	_	_	_	_	5	6
Paralytic shellfish poison	_	_	_	_	_	_	1	2
Other	_	_	_	_	_	_	5	5
Subtotal	_	_	1	_	_	1	33	40
Parasitic								
Giardia	_	_	_	_	1	1	2	2
Trichinella	_	_	_	_	_	_	1	2
Subtotal	_	_	_	_	_	_	1	1
Viral								
Norovirus	2	1	_	3	5	9	116	287
Sapovirus	_					_	1	3
Astrovirus	_	_	_	_	_	_	_	1
Other	_	_	_	_	_	_	_	1
Subtotal	2	1	_	3	5	9	117	292
Single etiology	27	13	1	6	21	60	274	579
Multiple etiologies	1	2	_	_	_	3	3	7
Unknown etiology ¹	5	6	_	2	10	22	68	245
Total	33	21	1	8	31	85	344	831

Appendices' footnotes are on page 14.

Appendix Footnotes:

* If at least one etiology was laboratory-confirmed, the outbreak was considered to have a confirmed etiology. If no etiology was laboratory-confirmed, but an etiology was reported based on clinical or epidemiologic features, the outbreak was considered to have a suspected etiology.

[†] Contributing factors are defined as risk factors that either enable an outbreak to occur, or amplify an outbreak caused by other means. Contributing factors are classified into three categories: contamination factors (factors that introduce or otherwise permit contamination), proliferation / amplification factors (factors that allow proliferation or growth of the etiologic agent), and survival factors (factors that allow survival or fail to inactivate a contaminant) (Bryan FL, Guzewich JJ, Todd EC. Surveillance of Foodborne Diseases III. Summary and Presentation of Data on Vehicles and Contributory Factors: Their value and limitations. J Food Prot 1997;60(6):701 – 14). More than one contributing factor might be reported per outbreak.

[‡]Contributing factors:

- C1: toxic substance part of the tissue
- C2: poisonous substance intentionally / deliberately added
- C3: poisonous substance accidentally / inadvertently added
- C4: addition of excessive quantities of ingredients that are toxic in large amounts
- C5: toxic container
- C6: contaminated raw product food that was intended to be consumed after a kill step
- C7: contaminated raw product food was intended to be consumed raw or undercooked / under-processed
- C8: foods originating from sources shown to be contaminated or polluted (such as a growing field or harvest area)
- C9: cross-contamination of ingredients (cross-contamination does not include ill food workers)
- C10: bare-handed contact by a food handler / worker / preparer who is suspected to be infectious
- C11: glove-handed contact by a food handler / worker / preparer who is suspected to be infectious
- C12: other mode of contamination (excluding cross-contamination) by a food handler / worker / preparer who is suspected to be infectious
- C13: foods contaminated by non-food handler / worker / preparer who is suspected to be infectious
- C14: storage in a contaminated environment
- C15: other source of contamination
- P1: food preparation practices that support proliferation of pathogens (during food preparation)
- P2: no attempt was made to control the temperature of implicated food or the length of time food was out of temperature control (during food service or display of food)
- P3: improper adherence of approved plan to use Time as a Public Health Control
- P4: improper cold holding due to malfunctioning refrigeration equipment
- P5: improper cold holding due to an improper procedure or protocol
- P6: improper hot holding due to malfunctioning equipment
- P7: improper hot holding due to improper procedure or protocol
- P8: improper / slow cooling
- P9: prolonged cold storage
- P10: inadequate modified atmospheric packaging
- P11: inadequate processing (acidification, water activity, fermentation)
- P12: other situations that promoted or allowed microbial growth or toxin production
- S1: insufficient time and / or temperature control during initial cooking / heat processing
- S2: insufficient time and / or temperature during reheating
- S3: insufficient time and / or temperature control during freezing
- S4: insufficient or improper use of chemical processes designed for pathogen destruction
- S5: other process failures that permit pathogen survival

[§] No outbreaks in the data reported fall in this category.

An etiologic agent was not confirmed or suspected based on clinical, laboratory, or epidemiologic information.

