

Assessing risk factors of sporadic *Campylobacter* infection: a case-control study in Arizona

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SUMMARY

Case-control studies of sporadic *Campylobacter* infections have predominately been conducted in non-Hispanic populations. In Arizona, rates of campylobacteriosis have been historically higher than the national average, with particularly high rates in Hispanics. In 2010, health departments and a state university collaborated to conduct a statewide case-control study to determine whether risk factors differ in an ethnically diverse region of the United States. Statistically significant risk factors in the final multivariate model were: eating cantaloupe [odds ratio (OR) 7·64], handling raw poultry (OR 4·88) and eating queso fresco (OR 7·11). In addition, compared to non-Hispanic/non-travellers, the highest risk group were Hispanic/non-travellers (OR 7·27), and Hispanic/travellers (OR 5·87, not significant). Results of this study suggest Hispanics have higher odds of disease, probably due to differential exposures. In addition to common risk factors, consumption of cantaloupe was identified as a significant risk factor. These results will inform public health officials of the varying risk factors for *Campylobacter* in this region.

Key words: *Campylobacter*, community epidemics, foodborne infections.

INTRODUCTION

Campylobacteriosis is the leading cause of bacterial gastroenteritis in the developed world [1]. In the United States it causes an estimated 2 million infections every year, but is highly underreported with only 43 000 cases confirmed annually [2]. This is probably due to the generally mild clinical symptoms that are frequently seen and the sporadic nature of transmission [3–5]. Data from FoodNet, the Center for Disease Control and Prevention's enhanced surveillance system, estimated the US rate to be 12·63

cases/100 000 people in 2008 [6]. In Arizona, which is not part of FoodNet, rates have historically been higher and in 2008 the reported rate was 15·4 cases/100 000 [7]. Racial disparities are consistent across the United States and Arizona, with Hispanics having higher rates of disease than non-Hispanic whites (United States, 10·73 vs. 8·07; Arizona, 14·3 vs. 7·5) [6, 8]. Younger ages are also at increased risk with children aged <4 years having very high rates of disease [9]. In Arizona, children aged <4 years experience rates over 2·5 times the overall incidence (39·3 cases/100 000) [10].

Identification of risk factors for *Campylobacter* has largely been conducted using case-control studies of sporadic cases as outbreaks are rare. Common risk factors identified from these studies are foreign travel

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[3, 4, 11–13], consumption of poultry, both cooked and undercooked [13, 14], as well as raw dairy products [15], barbeque meat [16], contact with puppies [4, 17, 18] or animals with diarrhoea [17, 19], and contact or consumption of untreated water [20]. The majority of these studies have been conducted in Northern European countries and across FoodNet sites which are not generally representative of a more diverse ethnic profile.

The Arizona Department of Health Services conducted a collaborative statewide case-control study to determine what common or unique risk factors might be associated with reported *Campylobacter* infections in Arizona and whether the risk factors varied by ethnicity.

METHODS

Overview

The study was designed as a matched case-control study where cases were matched to controls (1:2) on age group, gender, and residential location. All cases were identified from the state's routine passive surveillance system.

Study population

Cases. Individuals with a positive culture for *Campylobacter jejuni* reported through the state's routine laboratory surveillance system were included in the study if they were (a) not part of a recognized outbreak, (b) reported to the health department as a laboratory-confirmed case from 1 April 2010 to 31 December 2010 and (c) able to be interviewed over the telephone within 3 weeks of the report date. Five attempts on various days and times were made for each case before being considered as a non-contact. A representative sample of 20% of cases from Maricopa County was randomly selected for recruitment due to the county's large population size and number of cases (selection took place each week of the study period). In all other 14 counties, recruitment was attempted for all reported cases [7, 21].

Controls. The goal was to match cases to controls (1:2) on age group (0–11 months, 1–9, 10–19, 20–29, 30–59, ≥ 60 years), gender and residential location. Reverse lookup was used to identify controls closest to the case residence. Households with home phone numbers on the same street within two blocks either direction were first called and, if no one was

available, the households on neighbouring streets of the case street were then called. Controls were excluded if they (a) had a *Campylobacter* infection in the last 30 days, (b) experienced diarrhoea or abdominal pain with a fever in the past 30 days, or (c) were not able to be interviewed within 2 weeks of the case interview. Students at the University of Arizona who were part of the SAFER (Student Aid for Field Epidemiology Response) team conducted interviews with those cases identified from Maricopa County and their associated controls. All other interviews were conducted by county and state health department epidemiologists. Subjects were excluded if they did not speak either English or Spanish.

Questionnaire design

Individuals with a laboratory-confirmed case of *Campylobacter* are reportable to county and state health departments under Arizona law within 5 days of confirmation (A.A.C. R9-6-202). Cases reported through this system are contacted and interviewed by the county health department in which the case resides. The standard questionnaire covers some of the commonly associated risk factors associated with *Campylobacter* infection reported in the literature. A more extensive questionnaire was designed for this study using a *Campylobacteriosis* questionnaire from the World Health Organization's 'Control and Prevention of *Campylobacter* Infections' [22] as a guide. The questionnaires collected demographic information, 2-week travel history (both domestic and international), 2-week food history asking about specific foods based on previous studies of both sporadic and outbreak-related infections, kitchen and food-handling practices, as well as animal and water exposure information. For cases, additional information was collected on symptoms and any medical care received. This document was also the basis for the age groups used in control recruitment.

Data management and analyses

A standardized database and data entry protocol were created and disseminated to all participating sites conducting interviews. Data from all sites were merged and binary variables were generated for analysis. All analyses were conducted at The University of Arizona using Stata v. 11.0 (Stata Corporation, USA).

The distribution of demographic factors for all cases and controls was calculated. Differences in factors between unmatched cases (cases without at least one matched control) and matched cases (cases with at least one matched control) were determined using Fisher's exact test or Student's *t* tests.

For all foods and many environmental exposures, the time period participants were asked to recall was 2 weeks prior to illness onset for cases and 2 weeks prior to the interview for controls. Participants were asked if they could definitively recall (yes/no response), or were not sure (do not remember), or if they frequently ate that item or had a particular exposure but could not say for certain. Ultimately, only 'yes' responses were used in the regression models to ensure a higher degree of confidence in the exposures of interest.

As only 34% of cases were successfully matched to at least one control, random-effects logistic regression modelling was used. This method was chosen because it allows for the inclusion of unmatched cases, in addition to the matched case/control pairs [23]. When generating these models, a random-effect variable was first created that identified each matched set [a case and its matched control(s) received the same number] and unmatched cases received a separate unique identifier. This variable was then included in each model to adjust for the variability in correlation introduced if both unmatched and matched sets were included. Univariate random-effects logistic regression was utilized to estimate magnitude of effect for each risk factor. Age and gender were included in all models to adjust for the original matching factors. Odds ratios (OR) and associated 95% confidence intervals (95% CI) for each risk factor were calculated. Both conditional and random-effects logistic regressions were performed. The magnitudes of effect were the same although the sample size was much smaller for the conditional analyses, resulting in wider confidence intervals. Results from the random-effects models are presented.

A goal of this analysis was to build a final model that included all statistically significant risk factors that were associated with campylobacteriosis in Arizona. The strategy to select risk factors for potential inclusion included variables previously identified in the literature and variables identified in the Arizona database that were associated with disease at $P \leq 0.10$. Backward, stepwise regression was then used to develop the final predictive model; the variable with the highest *P* value was dropped from the model

and the fit of each model was compared ($P < 0.05$) using the likelihood ratio test. As variables were removed from the models their impact on variables remaining in the model were assessed for confounding. If the parameter estimates changed by $>10\%$, the variable was kept in the model. Concordance among variables in the final model was tested to ensure there were no high levels of collinearity between risk factors. Specific interaction terms were evaluated between key behavioural risk factors and were included if statistically significant. Age and gender were forced into the final model to adjust for the matching in the study design. Age, rather than age group, was included as a continuous variable to account for any residual confounding that may have been introduced due to the broad age groups.

Initial analyses found increased crude ORs for both Hispanic ethnicity and travel. To further explore whether Hispanic (an uncommonly reported risk factor) was simply a surrogate marker for travel or an independent risk factor, a joint variable was created and included in the models: Hispanic/travellers, Hispanic/non-travellers, non-Hispanic/travellers and non-Hispanic/non-travellers (reference group).

Finally, as a validation step and to evaluate the potential effects from these response and control matching problems, a subsample of the data collected at The University of Arizona was analysed (23 controls/23 cases, 14 matched, 9 unmatched, all cases and controls from Maricopa County). This subsample represented the 'best' matching site in the study, meaning a higher percentage of controls were recruited to cases. Both random-effects and exact logistic regression models were run on this smaller dataset.

RESULTS

Demographics and symptoms

During the study period, 781 laboratory culture-confirmed cases of *C. jejuni* were reported to the state health department [24]; 424 cases were selected as eligible for the study [20% of Maricopa cases selected at random ($N = 90$), all cases from other counties ($N = 334$)]. Of these 110 (26%) cases participated. Of the non-Maricopa County cases, 20.4% were interviewed and included in the study ($N = 68$). Of the Maricopa County cases, 42 (46.7%) were interviewed.

A total of 61 control interviews were completed, resulting in 37 matched sets (a set consisted of a case and either one or two matched controls).

Table 1. Demographic characteristics for *Campylobacter* cases and controls

	All controls (<i>N</i> = 61) <i>n</i> (%)	All cases (<i>N</i> = 110) <i>n</i> (%)	Matched cases* (<i>N</i> = 37) <i>n</i> (%)	Unmatched cases (<i>N</i> = 73) <i>n</i> (%)	<i>P</i> value†
Gender					
Male	26 (43)	44 (40)	15 (41)	29 (40)	0.9
Female	35 (57)	65 (60)	22 (59)	44 (60)	
Age, years					
Mean (min-max)	44.2 (0.25–80)	32.4 (0.75–82)	42.6 (0.72–82)	27.2 (0.9–81)	0.003
0–11 mos.	1 (1.6)	2 (1.8)	1 (2.7)	1 (1.4)	0.63
1–9	11 (18)	30 (27)	7 (19)	23 (32)	0.15
10–19	1 (1.6)	14 (13)	1 (2.7)	13 (18)	0.02
20–29	2 (3.3)	10 (9.1)	1 (2.7)	9 (12)	0.105
30–59	27 (44)	32 (29)	18 (49)	14 (19)	0.001
≥ 60	18 (30)	22 (20)	9 (24)	13 (18)	0.45
Youth (<20 yr)	13 (21)	46 (42)	9 (24)	37 (51)	0.008
Ethnicity					
Hispanic	9 (15)	53 (48)	7 (19)	46 (63)	0.000
Non-Hispanic	49 (80)	48 (44)	24 (65)	24 (33)	0.001
Residence					
Urban	14 (23)	32 (29)	9 (24)	23 (32)	0.38
Suburban	18 (30)	25 (23)	13 (35)	12 (16)	0.24
Town	9 (15)	31 (28)	8 (22)	23 (32)	0.27
Rural/farm	12 (20)	16 (15)	6 (16)	10 (14)	0.78
Healthcare provider	2 (3)	7 (6)	2 (5)	5 (7)	0.80
Daycare provider	2 (3)	6 (6)	1 (3)	5 (7)	0.40

* Matched cases only included if at least one respective matched control interviewed.

† *t* test or Fisher's exact test of the difference between matched and unmatched cases.

Demographics for all controls and cases are shown in Table 1 as well as for matched and unmatched cases; 40% of the cases were male with an overall mean age of 32.4 years (median 39.5 years); 57% of all cases resided in either an urban or suburban area.

Symptoms for all cases and stratification by matched status are given in Table 2. All cases experienced diarrhoea and many reported abdominal pain (75%), fever (68%), nausea (55%) and headache (46%). At least 27% of cases reported bloody diarrhoea and 35% were hospitalized, indicating that these reported cases were more severe than average compared to the national estimated rate of hospitalization for *Campylobacter* of 17% [25]. The average duration of illness was 13.9 days (mode 7 days), with longer term effects (symptoms lasting >2 weeks) reported by 40% of cases. Medical risk factors, including chronic conditions and medications taken prior to the illness, were also enquired of, with 43% of cases reporting a pre-existing chronic condition. Only 10% and 13% reported taking antibiotics and antacids prior to illness, respectively, both reported risk factors for *Campylobacter* infection [11, 26, 27]. The only two conditions that differed between matched cases and

non-matched cases were vomiting ($P = 0.002$) and pre-existing chronic conditions ($P < 0.001$).

Risk factors for *Campylobacter* in Arizona

Table 3 reports results for various risk factors related to ethnicity and probable environmental and behavioural exposures. The largest OR was for Hispanic ethnicity (OR 6.01, 95% CI 2.7–13.5). The only behavioural risk factor found to be strongly associated with the odds of developing disease was travel history in the last 2 weeks (OR 3.9, 95% CI 1.3–11.8) and more specifically, travel to Mexico (OR 4.2, 95% CI 1.2–14.8). Contact with untreated water, by 'swimming in a river, lake or pond', was elevated but not statistically significant. A history of handling raw chicken was associated with infection (OR 2.5, 95% CI 1.2–5.3) (attributable risk 71.8%). While a history of washing cutting boards (OR 0.4, 95% CI 0.2–0.8) and counter tops with soap and water following raw meat preparation (OR 0.5, 95% CI 0.3–1.08) were both found to be protective against disease.

The odds of disease associated with specific food items are given in Table 4. During the study period,

Table 2. Characteristics of illness in *Campylobacter* cases

	All cases (<i>N</i> = 110) <i>n</i> (%)	Matched cases* (<i>N</i> = 37) <i>n</i> (%)	Unmatched cases (<i>N</i> = 73) <i>n</i> (%)
Reported symptoms			
Diarrhoea	110 (100)	36 (97)	73 (100)
Bloody stools	30 (27)	8 (22)	22 (30)
Nausea	61 (55)	17 (46)	44 (60)
Vomiting*	43 (39)	9 (24)	34 (47)
Fever	75 (68)	26 (70)	49 (67)
Abdominal pain	82 (75)	26 (70)	56 (77)
Chills	15 (14)	6 (16)	10 (14)
Headache	51 (46)	13 (35)	38 (52)
Hospitalized	39 (35)	12 (32)	27 (37)
Duration of illness, days, mean (range) [median]	13.9 (1–74) [10]	14.6 (1–74) [10]	13.6 (2–53) [10]
Lost work days, mean (range) [median]	3.7 (0–16) [1]	2.2 (0–16) [2]	4.2 (0–5) [0]
Received diagnosis from doctor	75 (68)	24 (65)	51 (70)
Number of visits to medical provider to receive diagnosis, mean (range) [median]	1.57 (1–7) [1]	1.44 (1–4) [1]	1.64 (0–7) [1]
Before infection			
Ulcer medication	3 (2.7)	1 (2.7)	2 (2.7)
Antibiotics	11 (10)	4 (11)	7 (10)
Antacids	14 (13)	5 (14)	9 (12)
Chronic condition(s)*	52 (43)	26 (70)	26 (36)

* Statistically different difference between matched and unmatched cases, $P < 0.05$.

only consumption of cantaloupe (OR 2.8, 95% CI 1.2–5.7) (AR 69.4%) and queso fresco, a traditional, often unpasteurized, white cheese (OR 4.4, 95% CI 1.4–13.3) (AR 84%) were statistically significant at the $\alpha = 0.05$ level. Consumption of some foods were found to be ‘protective’ against the disease including delicatessen chicken, roast beef, raw peas, tomatoes, blueberries and pasteurized dairy.

The final multivariate model is shown in Table 5. The largest risk factors were a history of eating cantaloupe (OR 7.64, 95% CI 2.0–28.6), handling raw poultry (OR 4.88, 95% CI 1.50–15.9) and eating queso fresco (OR 7.11 95% CI 0.9–55.2). Protective risk factors included a history of washing the cutting board following use for raw meat (OR 0.14, 95% CI 0.04–0.45), and a history of consuming blueberries (OR 0.15, 95% CI 0.04–0.60) or roast beef (OR 0.10, 95% CI 0.02–0.49). This model also included the joint relationship between Hispanic and travel history. Compared to non-Hispanic/non-travellers, the highest risk group were Hispanic/non-travellers (OR 7.27, 95% CI 1.7–31.5), and Hispanic/travellers (OR 5.87, 95% CI 0.8–43.5). Non-Hispanic/travellers had decreased odds (OR 0.59, 95% CI 0.3–11.7) although the sample size for this group was very small ($n = 4$).

Not shown in the table, Hispanic and travel history were analysed individually in separate models (not

using dummy variables) along with inclusion and exclusion of interaction terms. Being Hispanic was an elevated risk factor, but travel and the interaction between travel and Hispanic were not. No other interaction terms between ethnicity or travel with other risk factors were significant.

For the subsample (Maricopa data interviewed at The University of Arizona), few results were different from the full dataset (data not shown). Consumption of cantaloupe remained statistically significant and had an OR of 7.5 in the reduced set compared to an OR of 3.6 in the full dataset for the entire state.

DISCUSSION

Overall, the results of this study are similar to previous case-control studies of *Campylobacter* with two notable exceptions. First, the role of ethnicity has not been examined in other published studies. Most *Campylobacter* case-control studies have been performed in Northern European countries [12, 16, 19], Australia [14] and New Zealand [4]. In the one large case-control study in the United States only 8% [3] of the cases were Hispanic compared to 48% in this Arizona study.

While initially hypothesized that Hispanics were at greater risk due to more frequent travel to Mexico, it

Table 3. Association between selected univariate behavioural exposures for *Campylobacter* in Arizona and case-control status

Exposure	Controls (<i>N</i> = 61) <i>n</i> (%)	Cases (<i>N</i> = 110) <i>n</i> (%)	Random effects OR (95% CI)
Hispanic	9 (15)	53 (48)	6.01 (2.7–13.5)
Age <20 years	46 (42)	13 (21)	2.7 (1.3–5.5)
Attended a recent gathering	21 (34)	24 (22)	1.1 (0.9–1.3)
History of any travel in last 2 weeks (includes Mexico)	4 (7)	24 (22)	3.9 (1.3–11.8)
Recent travel to Mexico	3 (5)	20 (18)	4.2 (1.2–14.8)
Urban/suburban residence (vs. rural/farm)	32 (52)	57 (52)	0.80 (0.4–1.6)
Water contact in the last 2 weeks			
Drink well water	11 (18)	9 (8)	0.4 (0.2–1.2)
Recent history of swimming in untreated water (river, lake or pond)	1 (2)	9 (8)	3.7 (0.4–30.2)
Kitchen practices in the last 2 weeks			
Handled raw poultry	11 (18)	39 (35)	2.5 (1.2–5.3)
Handled raw red meat	15 (25)	25 (23)	0.9 (0.43–1.9)
Handled raw meat (other)	2 (3.3)	14 (13)	4.3 (0.94–19.6)
Handled raw eggs	20 (33)	36 (33)	1.0 (0.5–1.9)
Nibbled raw meat	2 (3.3)	4 (4)	1.2 (0.2–6.6)
Used cutting board for any raw meat	47 (77)	72 (66)	0.8 (0.4–1.8)
Used plastic cutting board	31 (47)	44 (40)	0.7 (0.4–1.3)
Used wood cutting board	8 (13)	25 (23)	2.1 (0.9–5.0)*
Used same board for raw meat and vegetables	24 (39)	47 (43)	1.5 (0.75–3.1)*
Cutting board			
Unused after contact with any raw meat	2 (3)	11 (10)	3.3 (0.7–15.3)
Washed with soap	44 (72)	59 (54)	0.4 (0.2–0.9)
Knives			
Unused after contact with any raw meat	2 (3.3)	12 (11)	3.6 (0.8–16.7)
Washed with soap	46 (75)	70 (64)	0.6 (0.3–1.1)
Counter			
Washed with soap	35 (57)	45 (41)	0.5 (0.3–0.96)
Wiped with water	3 (5)	11 (10)	2.1 (0.6–8.0)
Sprayed with cleaner	14 (23)	20 (18)	0.7 (0.3–1.6)
Hands			
Washed with soap	50 (82)	79 (72)	0.6 (0.3–1.2)
History of animal contact in the last 2 weeks			
Puppy	9 (15)	21 (19)	1.4 (0.6–3.2)
Any animal contact (domestic pets, farm or wild animals)	42 (69)	70 (64)	0.8 (0.4–1.5)
Contact with ill animal	3 (5)	12 (11)	1.0 (0.9–1.1)
Farm	4 (7)	15 (14)	2.3 (0.7–7.4)
Zoo	0	5 (5)	Omitted
Petting zoo	2 (3)	5 (5)	1.4 (0.3–7.7)
Fair	3 (5)	2 (2)	0.4 (0.1–2.3)
Contact with manure	5 (8)	15 (14)	1.8 (0.6–5.4)
History of eating in restaurants in the last 2 weeks			
Ate at sit-down restaurant	35 (57)	54 (49)	0.7 (0.4–1.3)
Fast food	35 (57)	56 (51)	0.8 (0.4–1.4)
Cafeteria	9 (15)	8 (7)	0.5 (0.2–1.2)
Delicatessen	7 (12)	9 (8)	0.7 (0.2–1.9)
Street vendor	1 (2)	11 (10)	6.7 (0.8–52.9)*
Concession stand	5 (8)	4 (4)	0.4 (0.1–1.6)
Snack bar	2 (3)	1 (0.9)	0.3 (0.02–3.0)
Gas station	4 (7)	14 (13)	2.1 (0.7–6.6)
Ready-to-eat foods from grocery store	11 (18)	12 (11)	0.6 (0.2–1.4)
Coffee house	1 (2)	7 (6)	4.1 (0.5–34.0)

OR, Odds ratio; CI, confidence interval.

Odds ratios in **bold** are significant at the 0.05 alpha level; those with an * indicate a value 0.05 < *P* value ≤ 0.1.

Table 4. Food specific risk factors for *Campylobacter* in Arizona

Food item	Controls (N = 61) n (%)	Cases (N = 110) n (%)	Random effects OR (95% CI)
Poultry			
Purchased for home	50 (82)	88 (80)	0.9 (0.4–2.2)
Raw and fresh	20 (33)	45 (41)	1.6 (0.8–3.2)
Raw and frozen	20 (33)	32 (29)	0.9 (0.4–1.7)
Pre-cooked	27 (44)	44 (40)	0.9 (0.4–1.7)
Poultry (any)	52 (85)	90 (82)	1.5 (0.8–3.0)
Poultry away from home	52 (85)	94 (86)	1.3 (0.7–2.6)
Chicken wings	13 (21)	20 (18)	1.1 (0.8–1.3)
Chicken breast	37 (61)	58 (53)	1.4 (0.9–2.1)*
Roasted chicken	22 (36)	24 (22)	1.02 (0.8–1.3)
Chicken stir-fry	8 (13)	15 (14)	1.4 (0.8–2.3)
Chicken nuggets or strips	8 (13)	23 (21)	1.2 (0.9–1.5)
Chicken salad	8 (13)	9 (8)	1.3 (0.8–1.9)
Grilled chicken	16 (26)	24 (22)	1.3 (0.9–1.8)
Delicatessen chicken	12 (20)	5 (4.5)	0.2 (0.08–0.7)
Chicken (other)	1 (2)	11 (10)	1.18 (0.9–1.6)
Roasted turkey	9 (15)	15 (14)	1.3 (0.9–1.8)
Delicatessen turkey	15 (25)	23 (21)	1.2 (0.9–1.8)
Poultry other	1 (2)	2 (2)	1.7 (0.5–5.3)
Meat			
Purchased for home	56 (92)	88 (80)	0.3 (0.1–1.2)
Raw and fresh	35 (57)	59 (54)	1.2 (0.6–2.5)
Raw and frozen	14 (23)	30 (27)	1.6 (0.7–3.3)
Pre-cooked	22 (41)	39 (36)	1.2 (0.6–2.4)
Meat (any)	56 (92)	99 (90)	1.4 (0.4–5.5)
Ground beef	44 (72)	71 (65)	0.7 (0.3–1.7)
Roast beef	18 (30)	10 (9.1)	0.2 (0.1–0.6)
Steak	32 (53)	44 (40)	0.6 (0.3–1.2)
Beef jerky	5 (8)	4 (4)	0.4 (0.1–1.7)
Ham	21 (34)	47 (43)	1.5 (0.8–3.0)
Pork chops	17 (28)	26 (24)	0.8 (0.4–1.6)
Bacon	28 (46)	43 (39)	0.8 (0.4–1.5)
Ribs	11 (18)	14 (23)	0.7 (0.3–1.6)
Pork other	7 (12)	7 (12)	0.5 (0.2–1.6)
Sausage	18 (30)	35 (32)	1.1 (0.5–2.1)
Wild game	0	4 (4)	Omitted
Meat other	1 (2)	10 (9.1)	5.9 (0.7–47.1)*
Barbeque	33 (54)	51 (46)	0.7 (0.4–1.4)
Dairy, fruits and vegetables			
Eggs	49 (80)	89 (81)	1.4 (0.8–2.6)
Eggs: raw or runny	7 (11)	19 (17)	1.7 (0.6–4.3)
Eggs: purchase for home	46 (75)	86 (78)	1.0 (0.4–2.7)
Any raw fruit or vegetables	58 (95)	103 (93)	1.2 (0.2–7.3)
Lettuce	43 (71)	66 (60)	0.7 (0.3–1.4)
Pre-packaged lettuce	26 (43)	33 (30)	0.6 (0.3–1.2)
Spinach	14 (23)	20 (18)	0.8 (0.4–1.7)
Alfalfa sprouts	0	5 (5)	Omitted
Bean sprouts	0	4 (4)	Omitted
Raw peas	9 (15)	4 (4)	0.3 (0.07–0.8)
Raw carrots	32 (52)	45 (41)	0.7 (0.4–1.3)
Cilantro	17 (28)	39 (36)	1.5 (0.8–3.1)
Tomatoes	49 (80)	61 (55)	0.3 (0.1–0.7)
Raw mushrooms	12 (20)	19 (17)	0.9 (0.4–2.0)
Other raw veggies	36 (59)	38 (35)	0.4 (0.2–0.7)
Strawberries	32 (53)	44 (40)	0.7 (0.4–1.4)

Table 4 (cont.)

Food item	Controls (N = 61) n (%)	Cases (N = 110) n (%)	Random effects OR (95% CI)
Blueberries	20 (33)	18 (16)	0·4 (0·2–0·94)
Cantaloupe	11 (18)	36 (33)	2·6 (1·2–5·7)
Other raw fruit	47 (77)	65 (59)	0·4 (0·2–0·9)
Potato salad	10 (16)	18 (16)	1·0 (0·4–2·4)
Unpasteurized dairy	3 (5)	3 (3)	0·5 (0·1–2·8)
Pasteurized dairy	54 (89)	72 (65)	0·2 (0·1–0·6)
Milkshake	13 (21)	19 (17)	0·8 (0·4–1·8)
Ice cream	37 (61)	53 (48)	0·7 (0·4–1·3)
Yogurt	26 (43)	56 (51)	1·5 (0·8–2·8)
Queso fresco	4 (7)	25 (23)	4·4 (1·4–13·3)
Unpasteurized juice	3 (4·9)	2 (1·8)	0·3 (0·05–2·01)
Salsa	21 (35)	26 (24)	0·7 (0·3–1·3)

OR, Odds ratio; CI, confidence interval.

Odds ratios in **bold** are significant at the 0·05 alpha level, those with an * indicate a value 0·05 < P value ≤ 0·1.

was found that while travel was a general risk factor (crude OR 3·9, 95% CI 1·3–11·8), compared to non-Hispanic/non-travellers, Hispanics who did not travel actually had the highest odds of disease (OR 7·27) followed by Hispanic/travellers (OR 5·87). This differential among Hispanics could possibly be due to increased exposure and higher levels of immunity in this Arizona Hispanic population that travel to Mexico frequently; the possible development of immunity to *Campylobacter* has been reported in other highly exposed populations [28]. Questions about the frequency of travel and length of time residing in the United States would be important in future studies to determine what role, if any, this factor plays for populations in this region. However, these analyses indicate that Hispanics in Arizona have a higher risk of disease that is not related to travel history.

Differences in ethnicity due to dietary practices such as the consumption of queso fresco and cantaloupe, the younger age distribution of the population and kitchen maintenance practices might explain some of these differences, although none of these variables were statistically significant when stratified analyses were conducted (data not shown). Future studies should examine additional factors such as food preparation techniques and the role of socioeconomic status as possible explanations for these differences by ethnicity.

The second result that has not been reported in other studies was the increased risk associated with cantaloupe consumption. Most studies reported consumption of fresh fruits and vegetables to be protective [13], although it was not reported if any of these studies asked specifically about cantaloupe.

Potentially, the source of cantaloupe in Arizona might differ from other regions of the country and be more vulnerable to contamination. Only one reported outbreak in 1985 of *C. jejuni* had been linked to cantaloupes [29], although more recent outbreaks of *Listeria* [30] and *Salmonella* [31] were both linked to the fruit. History of consumption was found to have statistically significant elevated ORs in the univariate and multivariate models.

This current study identified many risk factors that were similar to other studies. Those statistically significant risk factors included youth (aged <20 years) [14, 18], travel history [3, 4, 11–13], handling raw poultry [4, 32], and consumption of queso fresco [4, 13, 15, 32, 33].

For the observed 'protective' exposures, washing a cutting board following meat preparation follows other findings that washing cutting boards can decrease cross-contamination and risk of disease [34]. Consuming roast beef may simply be an effect of choosing beef over chicken, which is known to be a riskier food exposure [20, 35, 36]. For blueberries, this may also be an indicator of general dietary practices. Another possibility is that antibacterial properties against other enteric pathogens such as *Salmonella* have been identified for blueberries [37]. The three strongest risk factors in this study, handling raw poultry, consumption of cantaloupe and consumption of queso fresco all had high attributable risk percentages (72%, 69%, 84%, respectively). This indicates that focusing on behaviour modification surrounding these few risk factors could have a large impact on reducing the burden of disease overall in this population.

Table 5. Final multivariate model* for risk factors associated with *Campylobacter* infections in Arizona

Risk factor	OR†‡	95% CI	P value
History of eating cantaloupe	7.64	2.04–28.6	0.003
Handling raw poultry	4.88	1.50–15.9	0.005
History of eating queso fresco	7.11	0.91–55.2	0.006
History of reporting ‘wash cutting board with soap and water following use for raw meat’	0.14	0.04–0.45	0.001
History of eating blueberries	0.15	0.04–0.60	0.007
History of eating roast beef	0.10	0.02–0.49	0.004
Hispanic/travel status			
Hispanic travellers	5.87	0.8–43.5	0.083
Hispanic non-travellers	7.27	1.7–31.5	0.008
Non-Hispanic travellers (small sample size, n = 4)	0.59	0.3–11.7	0.73
Non-Hispanic non-travellers	Ref.		

OR, Odds ratio; CI, confidence interval.

* Variables included in the initial model, but not found to be statistically significant include: History of eating raw peas, tomatoes, other raw vegetables, other raw fruit, pasteurized milk, unpasteurized milk, chicken breast, other meat, washing kitchen counters after preparing raw meat, using the same board for raw meat and vegetables, using a wood cutting board and eating from a street vendor.

† Random effects logistic regression of all cases and all controls, backwards stepwise regression.

‡ Adjusted for age and gender.

Limitations

Logistical barriers impeded the ability of the various recruitment sites to acquire matches for all cases. Lack of dedicated resources at several interview sites led to differential matching success. Recruitment of Hispanic controls could have been improved by using only bilingual interviewers but that option was not possible given the unfunded nature of the study. While differences in recruiting Hispanic cases (48%) and controls (15%) may appear marked at first, by matching on the other three variables, the controls were still well matched to cases.

The next two limitations were related to gender and age. Only 40% of those interviewed were male compared to 52% of reported cases during the study period statewide being male [38], although cases and controls

recruited into the study were matched well on gender. Matching on gender may have affected control recruitment; however, this step likely ensured that we had a higher number of men than we would have had without this as part of the protocol. Moreover, there was a substantial difference between cases and controls by age group. For cases aged >30 years, the matching to controls was successful; however, recruitment in the younger groups was problematic. For younger children, parents were more often home and able to answer questions. For school-age children, parents were less often available during the day, and for children aged >14 years it was difficult to convince both the child and the parent to answer the 30-min questionnaire. For the control group aged 20–29 years, the greatest obstacle was the protocol-prescribed use of only landline numbers. National survey estimates during this time period reported that 23% of adults did not have a landline and these rates were much higher for younger adults [39]. Given that low level of landline use, it was not surprising that few controls in the 20–29 years age group were recruited [39]. Future studies would need to review the benefits of matching by neighbourhood to the limitations of using only listed landlines numbers, or determine how to get access to cell phone numbers by address [39].

A final limitation is due to the number of potential risk factors (over 100) assessed and the possibility that some of our results are due to chance. However, given the large ORs and highly significant results of the more novel risk factors identified in the full model, namely being Hispanic ($P = 0.018$) and consumption of cantaloupe ($P = 0.003$), we are confident these results would hold in a larger study or under more conservative multiple comparison methods.

Strengths

This study is one of a few statewide population-based case-control studies of enteric disease in the United States. All steps in the process from design, implementation and data analysis and interpretation were completed by a collaborative partnership of local, county, state, and university public health professionals and numerous departments and jurisdictions. This study utilized a more extensive questionnaire than has typically been utilized for *Campylobacter* providing enhanced information for broader characterization of risk factors within this region. In addition, despite the length of the questionnaire, overall the study had very complete

data, with <20% missing data from any one variable, most around $\leq 10\%$. Finally, Arizona represents a unique and highly diverse population that has experienced higher rates of infection, making it important to determine the contributing factors.

The results of the smaller subset analyses also helped to validate the findings of the overall study. In addition, it was unlikely that differences in the cases had any effect on the ability to recruit a matched control. There were very few differences in symptomatology by matched and unmatched cases and the differences by chronic condition were not due to age group and gender.

Many public health case-control studies suffer the consequences from low response rates due to lack of incentives for participation, particularly in community-wide studies. For studies that choose to match as part of their study design, low response rates yield very low numbers of paired matches, thus decreasing study power and the likelihood of observing statistically significant results. In these circumstances, using a random-effects model allows all of the cases, both matched and unmatched, to be included in the analyses which increases the precision of the estimated fixed effects and also may strengthen the generalizability of the results to a broader community setting.

CONCLUSION

In this statewide study of *C. jejuni*, results suggest there are some unique risk factors that may be contributing to the higher rates of disease seen in Arizona. It is clear that the high rate of reported infections in Hispanics should be investigated further. Future studies should focus on understanding the role ethnicity might have on either exposure frequency, symptoms or possible differential reporting of disease. The result from this study will help to inform public health officials of the varying risk factors for *Campylobacter* in this region and beyond and can be used to modify the routine questionnaires to address these risks.

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DECLARATION OF INTEREST

None.

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