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Investigating locally-relevant risk factors for Campylobacter infection in Australia: protocol for a case-control study and genomic analysis

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SCHOLARONE™ Manuscripts

Investigating locally-relevant risk factors for *Campylobacter* infection in Australia: protocol for a case-control study and genomic analysis

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ABSTRACT

Introduction:

The CampySource project aims to identify risk factors for human *Campylobacter* infection in Australia. We will investigate locally-relevant risk factors and those significant in international studies in a case-control study. Case isolates and contemporaneous isolates from food and animal sources will be sequenced to conduct source attribution modelling, and findings will be combined with the case-control study in a source-assigned analysis.

Methods and analysis:

The case-control study will include 1,200 participants (600 cases and 600 controls) across three regions in Australia. Cases will be recruited from campylobacteriosis notifications to health departments. Only those with a pure and viable *Campylobacter* isolate will be eligible for selection to allow for whole genome sequencing of isolates. Controls will be recruited from notified cases of influenza, frequency matched by sex, age group and geographical area of residence. All participants will be interviewed by trained telephone interviewers using a piloted questionnaire.

We will collect *Campylobacter* isolates from retail meats and companion animals (specifically dogs), and all food, animal and human isolates will undergo whole genome sequencing. We will use sequence data to estimate the proportion of human infections that can be attributed to animal and food reservoirs (source attribution modelling), and to identify spatial clusters and temporal trends. Source-assigned analysis of the case-control study data will also be conducted where cases are grouped according to attributed sources.

Ethics and dissemination:

Human and animal ethics have been approved. Genomic data will be published in online archives accompanied by basic metadata. We anticipate several publications to come from this study.

KEYWORDS

Campylobacter, case-control study, risk factors, Australia, whole genome sequencing, source attribution, source-assigned analysis

ARTICLE SUMMARY

Strengths and limitations of this study

- Case-control study is well-powered to identify locally-relevant risk factors.
- Linking genomic data to the case-control study strengthens the analysis by enabling source attribution and source-assigned analyses to be conducted.
- Case-control questionnaire questions are being validated in a separate study, demonstrating the reliability of participant recall.
- Potential reporting bias due to inaccurate recall of study participants.
- Case-control study lacks efficiency for risk factors with high levels of exposure in the study population.

INTRODUCTION

Campylobacter infection is the most commonly notified cause of foodborne gastroenteritis in Australia, ¹⁻³ as well as a leading cause of bacterial gastroenteritis world-wide. ⁴ At the introduction of Australia's National Notifiable Diseases Surveillance System (NNDSS) in 1991 the incidence rate of notified campylobacteriosis cases was 79.1/100,000 population, ⁵ and despite notification rates plateauing in recent years, incidence had risen to 139.7/100,000 population in Australia in 2015, ⁵ with an estimated 10 cases for every notified case within the community. ⁶ By comparison, the incidence rate of campylobacteriosis in New Zealand in 2014 was 150.3/100,000 population, ⁷ with an estimated 10-30 cases in the community for every notified case. ⁸ Campylobacter notification rates in Australia and New Zealand are still among the highest in the world across high-income countries. Most countries in the European Union consistently report annual campylobacteriosis notification rates below 100/100,000 population. ²

Two species of *Campylobacter—Campylobacter jejuni* and *C. coli*—contribute to approximately 95% of human campylobacteriosis. ⁹ These *Campylobacter* species are commonly detected in sewage and surface water, ¹⁰ reside in the gastrointestinal tract of birds and animals, ¹¹ and are frequently found in raw meat, particularly poultry, and raw milk. ¹² ¹³ Campylobacteriosis is mostly foodborne, with an estimated 77% of cases transmitted via food consumption in Australia. ¹⁴ ¹⁵ Direct and indirect zoonotic transmission can occur via animal contact (direct) or faecally-contaminated water or environments (indirect). Person-to-person transmission is considered rare. ¹⁶ The majority of cases are thought to be sporadic, with outbreaks less commonly detected. ¹⁷ Most outbreaks are linked to the consumption of poultry, raw milk, or contaminated water. ¹⁷ ¹⁸

Targeted control of foodborne bacterial pathogens generally depends on identification of sources and routes of transmission. Since *Campylobacter* are ubiquitous in the environment and most cases are sporadic, identifying sources is difficult. Source attribution methods require isolation of strains from reservoirs to compare *Campylobacter* strain diversity in foods and animals to that in human infections. Beef, sheep and pig meat have a lower prevalence of *Campylobacter* contamination than chicken meat (<5% to 14%), ¹⁹⁻²¹ but a higher prevalence is found in animal offal such as liver, ²² thus making offal a valuable source of host-associated strains of *Campylobacter* in low-prevalence meats.

STUDY RATIONALE

In the United States, evidence from case-control studies have led to policy change, including changes to chicken slaughtering techniques. The incidence of human *Campylobacter* infection has declined in the US since this policy was introduced in 1997.²³ More recently, evidence from source attribution analyses in New Zealand has led to the development of poultry production policies and practices aimed at reducing the risk of *Campylobacter* transmission via poultry food products.²⁴ New Zealand has seen a 74% reduction in the number of campylobacteriosis cases attributed to poultry in the region, as well as a 54% reduction in cases overall.²⁵

Source attribution modelling enables us to determine which foods and animals are the most likely sources of infection with each *Campylobacter* strain type, and the proportion of cases attributed to each source. This can be done with simple proportional similarity index (PSI) calculations, or by using more complex models.²⁴ Source attribution also allows for human campylobacteriosis cases to be grouped by potential source, increasing the specificity of risk factor analyses. These source-assigned analyses combine the epidemiological information gained through the traditional case-control study with source attribution modelling to provide greater explanatory power to investigate locally-relevant risk factors.

OBJECTIVES

This study aims to:

- Identify dietary, environmental and behavioural risk factors for Campylobacter infection in Australia
- 2. Strengthen the epidemiological evidence for previously identified risk factors in Australia
- 3. Identify strain-specific risk factors for infection using Whole Genome Sequencing (WGS) data from case isolates

HYPOTHESES

We will test several hypotheses regarding specific risk factors for *Campylobacter* infection in Australia. The hypotheses are based on exposures which have previously been identified as risk factors for *Campylobacter* infection in Australia as well as internationally.

We hypothesise that:

- 1. Persons who consume undercooked meats, particularly chicken, are at increased risk of infection.
- 2. Persons who consume offal are at increased risk of infection.
- Persons who own companion animals (especially pupples) are at increased risk of infection.
- 4. Poor food hygiene and handling practices in the home increase the risk of infection.
- 5. Most human infections will be attributed to consumption of chicken meat.
- 6. There will be a high level of genetic diversity amongst *Campylobacter* strains.

STUDY DESIGN

We will conduct a case-control study including genomic testing over a two-year period in three sentinel sites: the state of Queensland (QLD), the Australian Capital Territory (ACT), and Hunter New England (HNE) region of New South Wales (Figure 1). Sporadic cases of culture-positive *Campylobacter* infection will be identified either through state notifiable disease registers, from local pathology service databases or local notification databases. An isolate from each case will be paired with epidemiological data from the case interview. One control will be recruited for each case who participates in the study, with trained interviewers conducting telephone interviews with both cases and controls. Participants will be interviewed using a questionnaire that has been specifically designed to collect information on known potential risk factors. This questionnaire will include a selection of questions being validated in a separate study (Liana Varrone, Validation of questions designed for gastroenteritis investigation). For cases, the questions will cover the seven days prior to the onset of illness, while controls will be questioned on the seven days prior to interview. Meanwhile, *Campylobacter* isolates will also be collected from food and animal samples. All human and non-human isolates will undergo whole genome sequencing for comparison in source attribution modelling. Data for this study will be collected from 1st March 2017 to 1st March 2019.

Figure 1. Map of Australian states and territories, showing the Hunter New England region. (Adapted from figure 1 in Eastwood *et al.* 2010)²⁶

Patient and public involvement

To develop the study, we engaged state and territory health departments, food safety agencies and industry to establish research questions and methods. The process involved a dedicated workshop, followed by teleconferences and an iterative process of drafting study documentation. We also established a reference panel, which includes representatives from senior levels of government and industry bodies. No patients or other members of the public were involved in the development of this study.

STUDY POPULATION

The three sentinel sites cover a population of approximately 6.1 million people. Based on notification and diagnostic pathology data, we expect approximately 8,650 *Campylobacter* cases to be notified across these sites during the study period.

DEFINITION AND SELECTION OF CASES

Case definition

We define a case as a person from any of the three participating sites with a recent history of acute diarrhoea and a culture-positive stool result for *Campylobacter*.

SAMPLE SIZE

We used risk factor prevalence data from a previous national *Campylobacter* case-control study in 2001/2002 to estimate sample size for this study. For example, the prevalence of chicken consumption among controls in 2001/2002 was 80%. A sample size of approximately 1,040 subjects (520 cases; 520 controls) would enable the study to detect an association between chicken consumption and illness with an odds ratio of 1.6, at 80% power and α = 0.05, as reported in the previous study. Sample size estimates for other potential risk factors are listed in Table 1.

Table 1. Sample size estimates for an unmatched case-control study

Risk factor	Prevalence of exposure among controls (%)	Prevalence of exposure among cases (%)	Odds ratio	No. of required study subjects
Beef	78	85	1.6	960
Pork	52	60	1.4	1130
Lamb	42	50	1.4	1120
Chicken	80	87	1.6	1040
Offal	2.0	5.0	2.6	1154
Puppies	2.1	5.4	2.7	1040

80% power and $\alpha = 0.05$

From these calculations, we estimate that a study of 1,200 subjects (600 cases; 600 controls) will adequately detect significant associations of these magnitudes for potential risk factors of interest.

Queensland and Hunter New England sites will each enrol at least 250 cases into the study, while ACT will enrol at least 100 cases. Based on the previous Australian case-control study,²⁷ we expect approximately 80% of selected notified cases to be eligible and participate in the study (Table 2).

Table 2. Sampling method for cases in each site

State	Expected number of notified cases during study period	Estimated cases from participating pathology laboratory	Culture +ve cases	Sequential sampling of notified cases	Total no. of cases	Expected no. to be recruited (~ 80% participation rate)
QLD	7000	2800 (40%)	1260 (45% in QLD)	Select every 4 th case	315	250
ACT	600	130	130	Include all notified cases	130	100
NSW (Hunter New England)	~1050	313	313	Include all notified cases	313	250
Total	8650	3243	1703		758	600

In Queensland, we will obtain cases from one private pathology provider reporting approximately 40% of the state's *Campylobacter* notifications. We estimate that this provider will notify 2,800 cases during the study period with an estimated 45% of these being culture-positive (1,260 notified cases). In ACT, approximately 600 *Campylobacter* notifications are expected during the study period; 130 are expected from the participating pathology laboratory. In Hunter New England, approximately 1,050 *Campylobacter* notifications are expected during the study period; 313 of these notifications will be from the participating pathology laboratory.

Enrolment of Cases

We will enrol all cases who meet the eligibility criteria (Table 3). Each site will check for new notifications of culture-positive *Campylobacter* infection daily, with only culture-positive *Campylobacter* cases eligible for this study. If a case refuses to participate in the study, we will select a subsequent case for inclusion. Enrolment of cases will depend on consent from the patient, or in the event of a child aged less than 18 years, consent from either one of the parents or the child's guardian. We will interview cases as soon as possible by telephone, preferably within two weeks of notification from the laboratory. It will be at the parent's or guardian's discretion as to whether a

child aged between 15 and 17 years is interviewed directly. The parent or guardian will be interviewed for cases aged less than 15 years.

Table 3. Eligibility criteria for cases and controls

Criteria	Cases	Controls
Had diarrhoea (≥3 loose bowel movements in	Include	Exclude
24hrs)		
Known date of illness onset	Include	N/A
Household members positive for Campylobacter	Exclude	Exclude (4 weeks prior to
in 4 weeks prior to onset of illness		interview date)
Household members experiencing diarrhoea in 4	Exclude	Exclude (4 weeks prior to
weeks prior to onset of illness		interview date)
Travelled outside of Australia in 2 weeks prior to	Exclude	Exclude (2 weeks prior to
onset of illness		interview date)
Travelled interstate for the entire 2 weeks prior	Exclude	Exclude (2 weeks prior to
to onset of illness		interview date)
Can't speak English	Exclude	Exclude
Not able to answer questions for some other	Exclude	Exclude
reason (e.g. intellectually disabled)		
Not contactable after 6 telephone attempts	Exclude	Exclude
Live outside the catchment areas	Exclude	Exclude
Do not have a telephone number available for	Exclude	Exclude
their primary residence, or a mobile phone		
An enteric pathogen other than Campylobacter	Exclude	N/A
was isolated/detected in their stool (excluding		
Blastocystis hominis and Dientamoeba fragilis)		

DEFINITION AND SELECTION OF CONTROLS

We will recruit controls from notified cases of influenza, frequency matched by sex, age group and geographical area of residence by Statistical Area Level 4 (SA4). These controls will be selected with a delay of at least six months from their influenza infection to ensure that controls have returned to eating their customary diet.

Each participating site (QLD, ACT or HNE) will establish a database of controls (previous influenza cases). All cases of influenza notified to the health department in each site between 1st January and 31st December 2017 will be entered into this control database. The age bands are 0-4 years, 5-14 years, 15-34 years, 35-54 years, 55-74 years, and ≥75 years. An appropriate control will be randomly selected from the database within 30 days of interview of the notified case.

Case and control recruitment

Interviewers trained in computer-assisted telephone interviewing (CATI) will conduct telephone interviews. A maximum of six attempts will be made to contact any one case or control, with no more than three attempts in any one day. Three calls will be attempted between 9:00am and 3:59pm, and three attempts between 4:00pm and 8:00pm. A text message will be sent to the potential participant after three failed call attempts, indicating that Public Health is trying to contact them. This protocol will be continued until the person is enrolled or excluded.

QUESTIONNAIRES

We will use specific case and control questionnaires for all participants (see Appendix 1). Cases will be asked additional questions about the clinical course of their illness and treatment. Interviewers will ask identical questions regarding exposures such as foods consumed, dining locations, water sources, domestic food handling techniques and exposure to animals of cases and controls. Questions on foods consumed, dining locations, water consumed, animal and pet exposures will be asked based on a seven-day history. Questions on international travel will be asked based on a two-week history. Antibiotic and antacid consumption, immunosuppressive treatment and household history of diarrhoea will be based on a four-week history. Questions on food handling and general kitchen practices will be based on usual practices rather than recent history. Demographic information will be collected from cases and controls. Contact information required to conduct interviews will be stored in a password-protected Excel document with only those needing to contact individuals given access. Piloted questionnaires were modified to remove repetitions, improve clarity, and to ensure that interviews could be conducted within 20 minutes.

DATA HANDLING & RISK FACTOR ANALYSIS

We will undertake descriptive reporting of campylobacteriosis incidence by person, place and time. We will also describe the severity of symptoms, treatment, and burden of illness.

Risk factor analysis will involve the examination of two-by-two contingency tables with chi square or exact tests to determine the presence of univariable associations between variables and disease. To measure the strength of an association, we will estimate odds ratios and calculate 95% confidence

intervals in a univariable analysis, followed by multivariable logistic regression modelling to adjust for potential confounders. Risk factors selected for inclusion in the regression model will include age, season and geographic area, variables with a significant univariable association with disease, and variables with a P-value ≤ 0.25 that are biologically plausible and of interest to the research team.

LABORATORY ANALYSES

Human samples

As outlined in Table 2, it is expected that 250 human isolates from Hunter New England, 250 from Queensland, 100 from Victoria and 100 from ACT will be sequenced. The initial isolation and confirmation of *Campylobacter* infection will be performed locally in each State/Territory. Only samples with a pure and viable culture will undergo WGS.

Animal and food samples

We will collect samples from chicken meat (covering the two production methods of continually housed and free range/housed), beef, lamb, pork, and from pet dogs. Given low prevalence of *Campylobacter* in meats other than chicken, samples will be collected from offal (preferably liver) from bovine, ovine and porcine sources to ensure sufficient positive samples are obtained for the study. Given the rising importance of chicken liver pate as a source of outbreaks in Australia, 28 chicken offal will also be sampled. Sample sizes by source are based on data from two states to ensure 50 positive samples per food source, and 30 samples in companion animals (Table 4). We will also contact veterinary clinics and teaching hospitals to ensure sufficient *Campylobacter*-positive samples from dogs. Water samples have been omitted from the genomic aspect of this study due to logistical constraints in sampling untreated water sources across the large geographical area involved in this study, and the complexity of designing an appropriate sampling frame. As there is a lack of evidence implicating municipal drinking water as sources of *Campylobacter* infection in Australia¹²⁷ we excluded water sampling from this study.

Table 4. Sampling to ensure 50 isolates per food source and 30 isolates from companion animals

	Foods					Anima	als	
	Chicken			Beef	Lamb	Pork	Dogs	Total
	Continually	Free-	Offal	Offal	Offal	Offal		
	housed	range						
Assumed	0.7	0.7	0.7	0.14	0.6	0.22	0.2	
prevalence								
Samples	72	72	72	286	100	272	150	1041
required								
Positive	50	50	50	40	60	60	30	330
isolates								

The initial isolation and confirmation of *Campylobacter* will be performed locally at laboratories in each State/Territory, with isolates forwarded to the Microbiological Diagnostic Unit Public Health Laboratory for WGS, except Queensland isolates which will be sequenced at Queensland Health. To detect seasonal and temporal variation in *Campylobacter* genetic types, 1041 non-human samples (estimated to produce 330 *Campylobacter* isolates) will be collected over a period of one year in Queensland, and two years in New South Wales. To assess latitudinal variation in chicken meat samples across eastern Australia, 105 chicken samples (70 chicken meat and 35 chicken offal) will be collected over a six-month period in Victoria. Food samples will be collected monthly from retail premises, using protocols from surveys undertaken in 2014 by partner organisations, with a pilot of 30 isolates in Queensland.

We will also collect an additional 20-30 human isolates from four additional Australian jurisdictions not participating in this case-control study to undergo WGS. This will be done over a two-month period that overlaps with the case-control study sample collection, and is planned to help inform the generalisability of the case-control study.

SEQUENCING AND SEQUENCE DATA PROCESSING

Campylobacter isolates selected for sequencing will be repurified on solid medium and a single colony selected for preparation of genomic DNA. A sequencing library will be prepared from the genomic DNA for sequencing on the Illumina sequencing platform (MiSeq or NextSeq). A sample of the selected colony will be regrown and cryopreserved (resuspended in liquid medium supplemented with 10% Glycerol and stored at -80°C). In some cases, Campylobacter enrichment cultures will be cryopreserved to enable future investigation of the genetic diversity of Campylobacters present. The short-read, paired end dataset produced by the Illumina Instrument from the genomic DNA of each isolate will be processed to produce a draft genome sequence for the isolate using a de novo assembler such as MEGAHIT.²⁹ The draft genome sequence will be annotated using Prokka.³⁰ We will use the draft genome sequence to perform the initial sub-species classification by deriving a multilocus sequence type (MLST) using the "Campylobacter jejuni/coli" typing scheme (pubmlst.org). Again, using the draft genome sequence, further typing e.g. virulence factors (http://www.mgc.ac.cn/VFs/) or antimicrobial resistance genotype (https://cge.cbs.dtu.dk/services/ResFinder/) will be performed using Abricate (https://github.com/tseemann/abricate). We will perform comparative genomics to examine the genetic relationships between selected subgroups of isolates in more detail using Nullarbor (https://github.com/tseemann/nullarbor).

SOURCE ATTRIBUTION MODELLING

We will analyse the epidemiological data within designated MLST groups or other typing groups derived from the genomic sequence data. Source attribution modelling and source-assigned analyses will be conducted.

Source attribution models combine typing data from isolates from food, animal and humans to estimate the proportion of human infections that can be attributed to animal and food reservoirs.³¹ Once inferred MLSTs have been ascertained, the proportional similarity index²⁵ will be used to assess similarities by source. We will then undertake source attribution analyses by adapting the asymmetric island model which has previously been applied to MLST data^{25 33} using Markov Chain Monte Carlo (MCMC) methods³⁴ implemented using the free software WinBUGS.³⁵ These methods will first be applied to MLST data extracted from whole genome sequences (the aforementioned "inferred MLSTs"), and then compared to structured phylogenetic modelling approaches^{36 37} that provide scope to infer inter-host transmission.

We will then group cases according to putative source based on these source attribution methods.³⁸ For example, all isolates attributed to chicken will be grouped together, regardless of differing strains. These cases attributed to chicken will then be compared to all controls in a risk factor analysis to produce a source-assigned analysis.

SPATIAL CLUSTERS AND TEMPORAL TRENDS

We will use newly-designated WGS-based MLSTs to assess heterogeneity in isolates from food sources and companion animals in Queensland and New South Wales, and in isolates from chicken meat and humans across Queensland, New South Wales, Victoria and ACT. A two-year sampling framework in New South Wales, one year of sampling in Queensland, and previous survey work in these states will allow us to assess the extent of seasonal and temporal trends. Postcode-level data associated with human illnesses will be used to detect space-time clusters using a scan statistic implemented in the free software SaTScan, at the Statistical Area 1 level. ³⁹ We will use a retrospective space-time permutation model to detect high risk clusters by comparing the observed number of illnesses to the expected number in that geographic zone and time-period. ⁴⁰

STUDY LINKAGES AND COLLABORATIONS

The CampySource Project Team comprises three working groups and a reference panel. The working groups focus on: food and animal sampling, epidemiology and modelling, and genomics. The reference panel includes expert representatives from government and industry.

The study is supported by the following partner organisations: the Australian National University, Massey University, University of Melbourne, Queensland Health, Queensland Health Forensic and Scientific Services, New South Wales Health, Hunter New England Health, Victorian Department of Health and Human Services, Food Standards Australia New Zealand, Commonwealth Department of Health and AgriFutures Australia – Chicken Meat Program.

CampySource is also supported by collaboration with the following organisations: ACT Health, Sullivan Nicolaides Pathology, University of Queensland, Primary Industries and Regions South Australia, Department of Health and Human Services Tasmania, Meat and Livestock Australia, and New Zealand Ministry for Primary Industries.

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While undertaking studies, LV is supported through an Australian Government Research Training Program (RTP) Scholarship.

DECLARATIONS

Ethics approval and consent to participate

Informed Consent

A suitably trained interviewer will inform potential participants about the purpose, methods and demands of the study. We will obtain verbal consent from all study participants or their guardians.

Persons aged 18 years and older will be interviewed following informed consent. It will be at the parent's or guardian's discretion as to whether a child aged between 15 and 18 years is interviewed directly, following informed parental/guardian consent. Parents/guardians will be interviewed for cases aged less than 15 years, after informed consent is obtained.

Confidentiality

All information and identifiers will be kept confidential. Names and personal identifiers will be collected and entered into computer records but will be password protected. No personal identifiers will be included in any published materials relating to this study. All hard copy questionnaires containing patient identifiers will be stored in locked filing cabinets in a secure location to which only study investigators and interviewers will have access.

Risks and Benefits

Participants will be informed there are no individual benefits associated with the study and that participation is voluntary. Failure to participate or a withdrawal of participation will not affect any future treatment. There is also no risk to the patient, and the only cost is time spent – approximately 20 minutes – being interviewed. They may refuse to answer any of the questions or stop at any time.

Animal Ethics

All procedures involving live animals will be performed in accordance with a protocol approved by the University of Melbourne's Animal Ethics Committee (ethics ID: 1714156).

Consent for publication

Not applicable.

Availability of data and materials

The Illumina read sets produced as part of this study will be published at INSDC (Sequence Read Archive (DDJB/NCBI) or the European Nucleotide Archive (EMBL-EBI))

Competing interests

No authors have any competing interests to declare.

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Author contributions

MDK conceived the original idea for this study. All authors contributed to the study design and analysis plan. LV and RJS wrote the first draft with contributions from all authors. LV, RJS, LS, MDK and KG were involved in multiple revisions. The final version of the manuscript was approved by all authors.

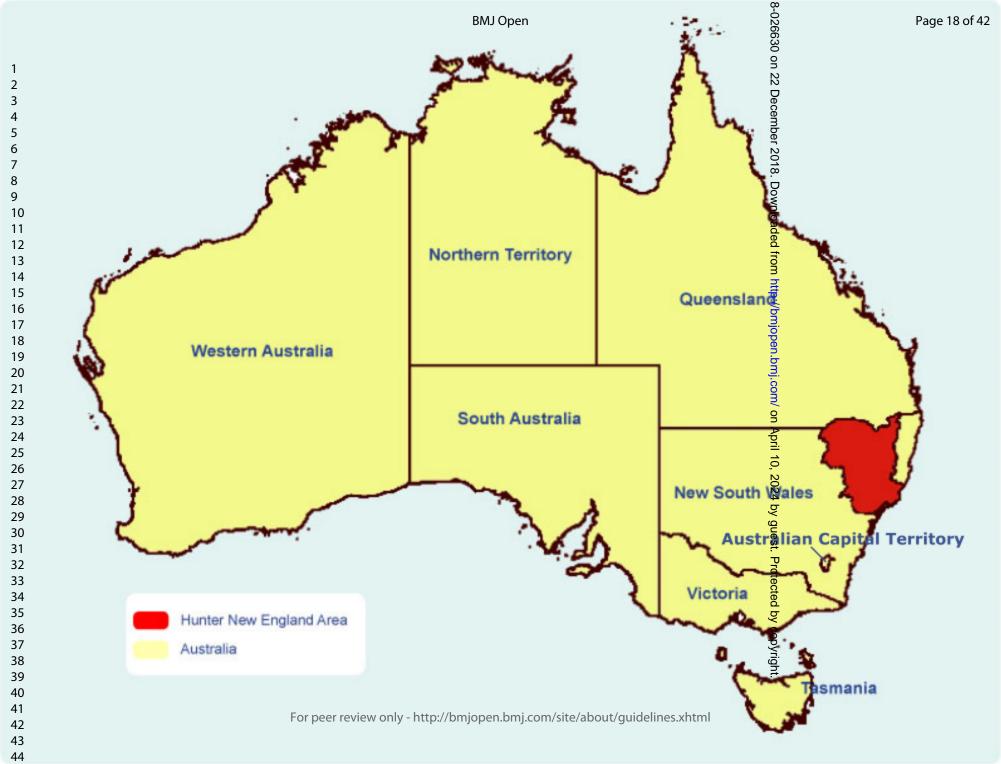
REFERENCES

- 1. The OzFoodNet Working Group. Monitoring the incidence and causes of diseases potentially transmitted by food in Australia: Annual report of the OzFoodNet network, 2011. *Annual Report* 2011; 39(2). http://www.health.gov.au/internet/main/publishing.nsf/Content/cda-cdi3902-pdf-cnt.htm/\$FILE/cdi3902g.pdf.
- 2. Kaakoush NO, Castaño-Rodríguez N, Mitchell HM, et al. Global Epidemiology of Campylobacter Infection. *Clin Microbiol Rev* 2015;28(3):687-720. doi: 10.1128/cmr.00006-15
- 3. Ford L, Kirk MD, Glass K, et al. Sequelae of Foodborne Illness Caused by 5 Pathogens, Australia, Circa 2010. *Emerg Infect Dis* 2014;20(11):1865. doi: 10.3201/eid2011.131316
- 4. Fernandes AM, Balasegaram S, Willis C, et al. Partial Failure of Milk Pasteurization as a Risk for the Transmission of Campylobacter From Cattle to Humans. Clin Infect Dis 2015;61(6):903-09. doi: 10.1093/cid/civ431
- 5. Commonwealth of Australia. National Notifiable Diseases Surveillance System, 2018.
- 6. Hall G, Yohannes K, Raupach J, et al. Estimating Community Incidence of Salmonella, Campylobacter, and Shiga Toxin–producing Escherichia coli Infections, Australia. Emerg Infect Dis 2008;14(10):1601-09. doi: 10.3201/eid1410.071042
- 7. The Institute of Environmental Science and Research Ltd. Notifiable Diseases in New Zealand: Annual Report 2014. Porirua, New Zealand, 2015.
- 8. Cressey P, Lake R. Estimated incidence of foodborne illness in New Zealand: application of overseas models and multipliers, 2011.
- 9. Lastovica AJ, Allos BM. Clinical Significance of Campylobacter and Related Species Other Than Campylobacter jejuni and Campylobacter coli. In: Nachamkin I, Szymanski CM, Blaser MJ, eds. Campylobacter, Third Edition: American Society of Microbiology 2008.
- 10. Fricker CR, Park RWA. A two-year study of the distribution of 'thermophilic' campylobacters in human, environmental and food samples from the Reading area with particular reference to toxin production and heat-stable serotype. *J Appl Bacteriol* 1989;67(6):477-90. doi: 10.1111/j.1365-2672.1989.tb02530.x
- 11. Blaser MJ, Taylor DN, Feldman RA. Epidemiology of Campylobacter jejuni Infections. *Epidemiol Rev* 1983;5(1):157-76.
- 12. Stern NJ, Hernandez MP, Blankenship L, et al. Prevalence and Distribution of Campylobacter jejuni and Campylobacter coli in Retail Meats. *J Food Prot* 1985;48(7):595-99.
- 13. Wood RC, MacDonald KL, Osterholm MT. Campylobacter enteritis outbreaks associated with drinking raw milk during youth activities: A 10-year review of outbreaks in the United States. *JAMA* 1992;268(22):3228-30. doi: 10.1001/jama.1992.03490220072031
- 14. Vally H, Glass K, Ford L, et al. Proportion of illness acquired by foodborne transmission for nine enteric pathogens in Australia: an expert elicitation. *Foodborne Pathog Dis* 2014;11(9):727-33. doi: 10.1089/fpd.2014.1746
- 15. Kirk M, Ford L, Glass K, et al. Foodborne illness, Australia, circa 2000 and circa 2010. *Emerg Infect Dis* 2014;20(11):1857-64. doi: 10.3201/eid2011.131315
- 16. Heymann DL, editor. *Control of Communicable Diseases Manual*. 19th ed. Maryland: United Book Press, Inc., 2008.
- 17. Pebody RG, Ryan MJ, Wall PG. Outbreaks of campylobacter infection: rare events for a common pathogen. *Commun Dis Rep CDR Rev* 1997;7(3):R33-7. [published Online First: 1997/03/07]

- 18. Tauxe Robert V, Hargrett-Bean N, Patton CM. Campylobacter Isolates in the United States, 1982-1986. MMWR Surveill Summ 1988;37(S S-2):1-13.
- 19. Sammarco ML, Ripabelli G, Fanelli I, et al. Prevalence and Biomolecular Characterization of Campylobacter spp. Isolated from Retail Meat. *J Food Prot* 2010;73(4):720-28.
- 20. Whyte P, McGill K, Cowley D, et al. Occurrence of Campylobacter in retail foods in Ireland. *Int J Food Microbiol* 2004;95(2):111-18. doi: 10.1016/j.ijfoodmicro.2003.10.018
- 21. Wong TL, Hollis L, Cornelius A, et al. Prevalence, Numbers, and Subtypes of Campylobacter jejuni and Campylobacter coli in Uncooked Retail Meat Samples. *J Food Prot* 2007;70(3):566-73.
- 22. Noormohamed A, Fakhr MK. A Higher Prevalence Rate of Campylobacter in Retail Beef Livers Compared to Other Beef and Pork Meat Cuts. *Int J Environ Res Public Health* 2013;10(5):2058.
- 23. Samuel M, Reilly K, Shallow S, et al. Burden of Campylobacter infection in the United States and declining trend in California, FoodNet 1996-1998. International Conference on Abstract Emerging Infectious Diseases. Atlanta, GA, 2000.
- 24. French NP, Marshall JC, the Molecular Epidemiology and Public Health Laboratory. Source Attribution Studies for Campylobacteriosis in New Zealand: Massey University, 2014.
- 25. Müllner P, Spencer SEF, Wilson DJ, et al. Assigning the source of human campylobacteriosis in New Zealand: A comparative genetic and epidemiological approach. *Infect Genet Evol* 2009;9(6):1311-19. doi: 10.1016/j.meegid.2009.09.003
- 26. Eastwood K, Durrheim DN, Merritt TD, et al. Field exercises are useful for improving public health emergency responses. *Western Pac Surveill Response J* 2010;1(1):12-18. doi: 10.5365/wpsar.2010.1.1.003
- 27. Stafford Russell J, Schluter PJ, Kirk MD, et al. A multi-centre prospective case-control study of campylobacter infection in persons aged 5 years and older in Australia. *Epidemiol Infect* 2007;135(6):978-88. doi: 10.1017/S0950268806007576
- 28. Merritt TD, Combs B, Pingault N. Campylobacter outbreaks associated with poultry liver dishes. *Commun Dis Intell Q Rep* 2011;35(4):299-300.
- 29. Li D, Liu C-M, Luo R, et al. MEGAHIT: an ultra-fast single-node solution for large and complex metagenomics assembly via succinct de Bruijn graph. *Bioinformatics (Oxford, England)* 2015;31(10):1674-6. doi: 10.1093/bioinformatics/btv033 [published Online First: 2015/01/23]
- 30. Seemann T. Prokka: rapid prokaryotic genome annotation. *Bioinformatics (Oxford, England)* 2014;30(14):2068-9. doi: 10.1093/bioinformatics/btu153 [published Online First: 2014/03/20]
- 31. Glass K, Fearnley E, Hocking H, et al. Bayesian Source Attribution of Salmonellosis in South Australia. *Risk Anal* 2016;36(3):561-70. doi: 10.1111/risa.12444
- 32. Hald T, Vose D, Wegener HC, et al. A Bayesian Approach to Quantify the Contribution of Animal-Food Sources to Human Salmonellosis. *Risk Anal* 2004;24(1):255-69. doi: 10.1111/j.0272-4332.2004.00427.x
- 33. Sears A, Baker MG, Wilson N, et al. Marked Campylobacteriosis Decline after Interventions Aimed at Poultry, New Zealand. *Emerg Infect Dis* 2011;17(6):1007. doi: 10.3201/eid1706.101272
- 34. Müllner P, Jones G, Noble A, et al. Source attribution of food-borne zoonoses in New Zealand: a modified Hald model. *Risk Anal* 2009;29(7):970-84. doi: 10.1111/j.1539-6924.2009.01224.x
- 35. Lunn DJ, Thomas A, Best N, et al. WinBUGS A Bayesian modelling framework: Concepts, structure, and extensibility. *Stat Comput* 2000;10(4):325-37. doi: 10.1023/a:1008929526011
- 36. Dearlove BL, Cody AJ, Pascoe B, et al. Rapid host switching in generalist Campylobacter strains erodes the signal for tracing human infections. *ISME J* 2015;10:721. doi: 10.1038/ismej.2015.149

- 37. Mather AE, Vaughan TG, French NP. Molecular Approaches to Understanding Transmission and Source Attribution in Nontyphoidal Salmonella and Their Application in Africa. *Clin Infect Dis* 2015;61(suppl_4):S259-S65. doi: 10.1093/cid/civ727
- 38. Mughini Gras L, Smid JH, Wagenaar JA, et al. Risk Factors for Campylobacteriosis of Chicken, Ruminant, and Environmental Origin: A Combined Case-Control and Source Attribution Analysis. *PLoS One* 2012;7(8):e42599. doi: 10.1371/journal.pone.0042599
- 39. Kulldorff M, Heffernan R, Hartman J, et al. A Space—Time Permutation Scan Statistic for Disease Outbreak Detection. *PLoS Med* 2005;2(3):e59. doi: 10.1371/journal.pmed.0020059
- 40. Touray K, Adetifa IM, Jallow A, et al. Spatial analysis of tuberculosis in an Urban West African setting: is there evidence of clustering? *Trop Med Int Health* 2010;15(6):664-72. doi: 10.1111/j.1365-3156.2010.02533.x





Refusal Ineligible	
ID Number	
Interview Date Interview Start T	ime
Data entered Date Data checked Date	

Source Attribution of Campylobacter in Australia Study

Case Questionnaire

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INTRODUCTION

Interviewer Note:	If case is less than 15 years of age you will need to speak to parent or guardian most familiar with the eating habits of the child.
	If case is aged between 15–17 years you will need to obtain parent or guardian consent prior to interview.
	Please note that for subjects under the age of 15 years, questions relate to the case, not the person being interviewed unless specified in the body of the questionnaire.

"Hello, my name is <Interviewers Name> and I am calling on behalf of [Queensland Health / ACT Health / Hunter New England Public Health Unit]."

[&]quot;May I please speak with <name of case> or <name of case's mother/father>?"

Interviewer Note:	When the case comes to the phone then repeat the introduction and proceed with the explanatory statement.
	If the case is unavailable then arrange an alternative time for the interview

"The Australian National University in conjunction with the [state health department] is conducting a study of *Campylobacter* infection in Australia to identify possible causes. *Campylobacter* infections are notifiable to health departments in each state and territory throughout Australia. We understand that you recently experienced an illness due to the *Campylobacter* bacteria. We would like to find out more about your *Campylobacter* illness. Your participation is voluntary, all responses are confidential and if there are any questions you do not wish to answer, just say so."

"Would you be prepared to answer some questions about your illness and activities prior to your illness? The questions will take approximately 20 minutes."

Yes No.

"In this study, we will collect information on foods you ate and activities you undertook prior to your illness. Your participation is voluntary and you can stop at any time. No individual information will be presented in any reports or presentations. Partners in this research will comply with the Australian Privacy Act 1988. An information sheet about this research project is available on the ANU website.

"Would you like me to send you a copy or provi	ide you with the link?"	Yes	•	No •
If yes, "Could you give me an address/email?" _				

"This study has been approved by the Australian National University Ethics Committee (Protocol 2016/426). If you have any concerns please direct them to

Human Research Ethics Committee

Research Services Office, Chancellery 10B

The Australian National University, ACT 2601

Tel: 6125 7945 Fax: 6125 4807

Email: Human. Ethics. Officer@anu.edu.au"

"Do you have any further questions about the project?"

Yes • No •

Do you agree to participate in the project?	Yes •	No •
---------------------------------------------	-------	------

If NO, arrange an alternative time to phone back to conduct the interview If YES, continue

[&]quot;Do you have the time right now to answer these questions?"

	for you to have a calendar or diary in front of y	around the time of your illness, it may be helpful ou. Do you need a few minutes to get these?"
	Yes, I will get one no No, I already have one with me Don't have access to a calendar	
	The first few questions we'll be asking you with [your/their] illness.	are about some symptoms that are associated
1.		diarrhoea as 3 or more loose stools or bowel you had your <i>Campylobacter</i> infection, did you
	Yes No Don't know/Not sure	Check ineligible box then END INTERVIE
1a	During this diarrhoeal illness, what was the m you had in any 24 hour period?	aximum number of stools or bowel movements
	0-2	(If response = '0-2', then recode Q.1 as = '2')
	More than 20	5
2.	For how many days did your diarrhoea las	t? DAYS
	Don't know/Not sure	77
	CALCULATE PRIOR TO INTERVIEW	
	Date stool specimen collected	Day Month Year
3.	Could you please let me know what to Day Month Year (If person is unsure of date then prompt with date of sto	he date was when your diarrhoea began?
	Don't know/Not sure	

Interviewer Note:

Refer to your calendar to determine the interval from DATE 4 WEEKS BEFORE

DIARRHOEA BEGAN to DATE 1 DAY BEFORE DIARRHOEA BEGAN.

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		Ş
		Оре
4.	In the 4 weeks before your illness began, that is from <date 4="" before="" diarrhoea="" weeks=""> through <date 1="" before="" began="" day="" diarrhoea="">, did anyone else in your household test positive for Campylobacter?</date></date>	n: first public
	Yes Check ineligible box then END INTERVI No Don't know/Not sure 7	MJ Open: first published as 10.1136/bmjopen-2018-026630 on 22
5.	In the 4 weeks before your illness began, that is from <date 4="" before="" diarrhoea="" weeks=""> through <date 1="" before="" began="" day="" diarrhoea="">, did anyone else in you household have diarrhoea?</date></date>	36/bmjoper
	Yes	EW 7-20
	Don't know/Not sure 2	918-
	Don't known tot sure	026
		630
Intervie	Refer to your calendar to determine the interval from DATE 2 WEEKS BEFORE DIARRHOEA BEGAN to DATE 1 DAY BEFORE DIARRHOEA BEGAN.	on 22 D
		есе
		mbe
6.	In the 2 weeks before your illness began, that is from <date 2="" before="" diarrhoea="" weeks=""> through <date 1="" before="" began="" day="" diarrhoea="">, did you travel overseas or interstate?</date></date>	r 2018. Do
	INTERVIEWER NOTE: IF participant answers "yes", 1. Clarify if the travel was overseas or interstate 2. If travel was interstate: Clarify the length of time spent interstate in the time period just mentioned	December 2018. Downloaded from http://b
	Options to select:	
	A. If the participant has travelled overseas or spent the whole two weeks interstate: (Select option Yes) B. If the participant has travelled interstate only for a portion of the time: (Select option No)	mjopen.bmj.com/ on April 10, 2024 by guest. Protected by copyright.
	Yes	v <u>\$</u> .
	No	con
	Don't know/Not sure	or or
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	HEALTH QUESTIONS
7.	During this illness, did you have any of the following symptoms?
•	Yes No DK/NS
;	a. Fever 1 2 7
1	b. Vomiting 1 2 7
,	c. Stomach cramps 1 2 7
,	d. Blood in your stool 1 2 7
	e. Nausea 1 2 7
	f. Headache 1 2 7
1	g. Muscle/body aches 1 2 2 7
3.	Did you take any antibiotics as a result of this illness?
	Yes
	No
	Don't know/Not sure
).	What antibiotic(s) were you taking? [Ask person to get tablet bottle, if possible]
	Azithromycin
	Ciprofloxacin
	Norfloxacin.
	Erythromycin
	Doxycycline (also known as Doxy or Vibramycin.
	Other (please specify)
	Don't know/Not sure
0.	Were you admitted to hospital overnight because of this illness?
	Yes
	No 2 Go to Q. 12
	Don't know/Not sure
11.	If yes, for how many nights were you hospitalised?
	Don't know/Not sure
Inter	Refer to your calendar to determine the interval from DATE 4 WEEKS BEFORM
	DIARRHOEA BEGAN to DATE 1 DAY BEFORE DIARRHOEA BEGAN.
	"For the next few questions, I would like to ask you about events which may have occurred in the
,	weeks before your illness began, so again that's from <date 4="" before="" began="" diarrhoe="" weeks=""> to <date 1="" before="" began="" day="" diarrhoea="">."</date></date>
]	weeks before your illness began, so again that's from <date 4="" before="" began="" diarrhoe="" weeks=""> to <date 1="" before="" began="" day="" diarrhoea="">."</date></date>
]	weeks before your illness began, so again that's from <date 4="" before="" began="" diarrhoe="" weeks=""> to <date 1="" before="" began="" day="" diarrhoea="">." In those 4 weeks, were you taking any antibiotics?</date></date>
,	weeks before your illness began, so again that's from <date 4="" before="" began="" diarrhoe="" weeks=""> to <date 1="" before="" began="" day="" diarrhoea="">."</date></date>

		ВМЛ	Open		Page 2 4, 0 S
					Open::
Intervie	ewer Note:	If person can't remember the n leave the space blank.	ame of the an	tibiotic(s), check the DK/	Page 2400 NS box and NS box and Pop taking these? IM)
13.	What antibio	tic(s) were you taking? [Ask pe	rson to get tal	olet bottle, if possible]	ed as 10.11:
			DK/NS	What date did you sto	op taking these?
a.	Antibiotic 1		_	(DD/M	IM) 7 DK/NS
b.	Antibiotic 2_		7	(DD/M	IM) \prod_{7} DK/NS $\stackrel{.}{01}_{8}$
c.	Antibiotic 3		_	(DD/N	IM) 7 DK/NS 266
d.	Antibiotic 4		7	(DD/M	IM) 7 DK/NS 9 8
14.	In those 4 we	eks, were you taking any regula	ar medicatio	n that decreases stomach	a acid?
	No	ot sure	<u> </u>	Go to Q. 16 Go to Q. 16	December 2018. Downloaded from http://bi
15.	Did you take	any of the following in the 4 w	eeks prior to	illness?	ownload
<u>H</u>	listamine-2 (H ₂)	Receptor blocker			ded f
a. b. c. d.	Zantac (Raniti Tagamet (Cim Pepcid (Famo	dine)tidine)tidine)	1	No DK/NS 2 7 7 2 7 7 2 7 7	rom http://bmjop
<u>P</u>	roton Pump Inh	ibitor			en.bmj.
a.b.c.d.e.	Nexium (Eson Somac (Panto) Pariet (Rabepi	razole)prazole)prazole)prazole)prazole)prazole)prazole)prazole)	···· 1 ··· 1	2 7 7 7 2 7 7 2 7 7	mjopen.bmj.com/ on April 10, 2024 by guest. Protected by copyright. dition or s disease,
16.	chronic illnes	er been told by a doctor that s in which diarrhoea or vomiti el syndrome, ulcerative colitis,	ng is a major	symptom? (e.g. Crohn'	dition or by guest. I
	No	ot sure.	1 Spe	cify(Protected t
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DIZ/NC

		1 63		110	1	/ 1 % / 1	113
a.	Prednisone						
	or other steroids <u>not</u> used on your skin		1		2		7
b.	Cyclosporine		1		2		7
c.	Chemotherapy		1		2		7
d.	Radiation therapy] 1] ,] ,

	BMJ Open	- ugc 1-8
3.	. EXPOSURES	וואָנ בוואָנ
itervi	Refer to your calendar to determine the interval from the DATE 7 DAYS BEFORE DIARRHOEA BEGAN to the DATE 1 DAY BEFORE DIARRHOEA BEGAN	ā
Ţ.	A. WATER	- 00/DIII
	I'm now going to ask you some questions about water that you consumed in the <u>7 days before your</u> iarrhoea began, that is from [diarr_7_days_prior] to [diarr_1_day_prior].	oben-zo-
	What is your main source of drinking water at home? (select one only)	0-0200
	INTERVIEWER NOTE: Only read out options if they're unsure	90
a.		N C G
b.		<u> </u>
c.		<u> </u>
d.	· · · · · · · · · · · · · · · · · · ·	
e. f.	Municipal water supply (tap water)	
g.		\$
h.		Jack
		<u> </u>
		=
tervi	iewer Note: If person answered "Yes" to "Municipal water supply" or "Purchased bottle water", skip to Q.21	2 (2) 2 (2) Published as 10.1130/bill/open-zo10-ozoooo on zz December zo10. Downloaded nom http://open-zo10-ozoooo on zz December zo10.
	iewer Note: If person answered "Yes" to "Municipal water supply" or "Purchased bottle water", skip to Q.21 Do you usually treat your main source of drinking water before drinking?	an into // binjopen.
	skip to Q.21	an inter/originations.
	Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water Yes	
	Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water	
•	Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water Yes	
	Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water Yes	. July metally controlled to the controlled to
a.	Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water Yes	on industrial objects on April 10, 2024 by guest
a. b.	Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water Yes	on the world bear on the control on which is a second of second on which is a second on secon
a.	Skip to Q.21 Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water Yes	all lite would be a sound on April 10, 2024 by great. Flored
a. b. c.	Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water Yes	on intervenipopen prinjecenia on April 10, 2024 by guesti. Floriected i
a. b. c. d.	Do you usually treat your main source of drinking water before drinking? If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water Yes	

60

21.		Did you <u>drink</u> water from any of the followi (Select all that apply)	ng sourc	es in th	he 7 days before onset of diarrhoe
	0	A rainwater tank	Yes	No	DK/NS
	a. b.	A river or stream	1		2 7
	c.	A private well, bore hole, or spearpoint	1		2 7 7
	d.	A carrier or tank truck			2 7
	e.	Municipal water supply (tap water)			2 7
	f.	Purchased bottle water			2 7
	g.	Other water supply	1		2 7
		Specify ()			
		Specify ()			

B. DINING LOCATIONS

22.	During this time, did you eat any food prepared outside your home, for example takeaway, restaurant, someone else's home?
	Yes
23.	Did you eat any food from the following places?
a. b. c. d.	Kebab shop
23a	period?
	1-2 meals
	3-4 meals
	≥ 5 meals
	Don't know/Unsure

C. DAIRY PRODUCTS

"I would now like to ask you about the dairy products you may have eaten in the 7 days before your diarrhoea began."

24. Did you drink any raw/unpasteurised milk or eat any products made from raw/unpasteurised milk?

INTERVIEWER NOTE:

Cold-pressed milk is pasteurised and is not to be included as "raw/unpasteurised".

a. b.		nilk		No 2 2	DK/NS 7 7
	Specify (0,)		

D. MEAT AND POULTRY

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I	D. MEAT AND POULTRY	MJ Open: first published as 10.1136/bmjopen-2018-026630 on
		t pub
66	I will now ask you some questions about meat and poultry that you may have eaten in the 7 days	lishe
<u>b</u>	pefore your diarrhoea began, that is from <date 7="" before="" began="" days="" diarrhoea=""></date>	das
t	rhrough <date 1="" before="" began="" day="" diarrhoea="">."</date>	10.11
•	During these 7 days, did you eat any of the following deli meats or cold cuts?	36/b
	Yes No DK/NS	mjo O
a	ı. Salami/mettwurst	pen-
	o. Cabanossi/cabana/twiggy sticks 1 2 2 7	201
	c. Ham/chicken/turkey/beef	8-0.
d	d. Devon/frankfurts/cheerios	266
e f		30 (
1	Specify (on Si
	,	22 D
	During these 7 days, did you eat any pate?	December 2018. Downloaded from http://b
	Yes	er 2
	No	018
	Don't know/Not sure	Dov
	Was the pate eaten,	vnload
	Chicken pate	ed fr
	Duck pate 2	om Om
	Pork pate	h tt p
	Another type of pate	_)
	Don't know/Not sure	
	Was this pate homemade or purchased from a store?	en.bm
	Homemade	j.cor
	Store	n/ o
	Don't know/Not sure	ň Ap
	During these 7 days, did you eat any other meat or poultry? Like beef, lamb, chicken etc.	njopen.bmj.com/ on April 10, 2024 by guest. Protected by copyright
	INTERVIEWER NOTE:	024 by
	This does not include eggs	gue
		št. P
	Yes	rote
	No	cte
	Don't know/Not sure	d by
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	BMJ Open	BMJ
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BEE	EF/VEAL	: first
30.	During these 7 days, did you eat any beef or veal?	publis
	Yes	hed
	No	as 1
	Don't know/Not sure	0.1136
31.	During the 7 days prior, did you eat any of the following beef or veal?	5/bmjo _l
		pen-
	DK/N	2018
	DK No Ye	8-02
a.	Minced beef dishes	6630 or
b.	Kebabs/souvlaki 1 2 7	1 22
c.	Offal	ec
d.	Other	ember 2
POR	PK	2018
101		 D
32.	During those 7 days did you get any nouls?	wnl
32.	During these 7 days, did you eat any pork?	bade
	Yes	ă fr
	No	om -
	Don't laile without sale	<u>#</u>
		//brr
33.	During the 7 days prior, did you eat any of the following pork?)jope
		n.br
	DK/N No Yes	nj.co
)m/
a.	Minced pork dishes	on Apri
b.	Kebabs/souvlaki	10,
C.	Offal	— <u>%</u>
d.	Other	24 by gu
LAN	<i>ПВ</i>	BMJ Open: first published as 10.1136/bmjopen-2018-026630 on 22 December 2018. Downloaded from http://bmjopen.bmj.com/ on April 10,-2024 by guest. Protected by copyrigh
34.	During these 7 days, did you eat any lamb/mutton?	ntecte
J 7.		d by
	Yes	cop
	No	yrig
		\neg

	BMJ Open	Page 3‰ f 42 ≦
35.	During the 7 days prior, did you eat any of the following lamb/mutton?	Open: firs
	No Yes	st publish
a.	Minced lamb/mutton dishes	ed as 10.
b. c.	Kebabs/souvlaki. 1 2 7 Offal	of AMJ Open: first published as 10.1136/bmjopen-2018-026630 on age Pa
d.	(eg. casserole, stir fry, steak, fillet, roast, lamb strips)	en-2018-C
<i>GAN</i> 86.	ME MEAT During these 7 days, did you eat any game meat like kangaroo, wallaby, venison or similar?)26630 o
,	Yes	, 22
POU	ULTRY	er 2018. I
37.	How often do you usually consume chicken/poultry meat? 3 or more days per week	December 2018. Downloaded from http://bmjop
38.	Puring the 7 days before your illness began, did you eat any chicken or other poultry? Yes	en.bmj.com/ on Ap
39.	How many meals did you eat that contained chicken or other poultry in the 7 days prior to onset of diarrhoea?	oril 10, 202
	1-2 meals	mjopen.bmj.com/ on April 10, 2024 by guest. Protected by copyright
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	Vac
	Yes
	Don't know/Not sure
	Was the chicken or poultry purchased? (Select all that apply)
	Raw and fresh
	Raw and frozen 2
	Pre-cooked
	Don't know/Not sure 4
	How was it stored before consumption? (Select all that apply)
	INTERVIEWER NOTE:
	(On the bench)
	This is only to be used if they STORE their meat on the bench, this does not include defrosting their meat of the bench.
	In the freezer
	- 1 011
	In the fridge 2
	On the bench
	On the bench
a.	On the bench
	On the bench
a. b. c.	On the bench
b. c.	On the bench
b. c.	On the bench
b. c. d.	On the bench

45.		During this time, on how many days did you eat poultry?
		INTERVIEWER NOTE: 1. A pate is included 2. Eggs are excluded
		Days:
46.		During this time, on how many days did you eat meat (including poultry)?
		INTERVIEWER NOTE: Pate is included
		Days:
47.		During the 7 days prior to illness, did [you/they] eat any meat product, which was raw, rare or appeared undercooked?
		Yes
48.		Which of the following meats did [you/they] eat that was undercooked?
	g.	Ves No DK/NS Chicken/poultry 1 2 7 Beef or veal 1 2 7 Pork 1 2 7 Lamb/mutton 1 2 7 Game meat 1 2 7 Minced meat items 1 2 7 including sausages, hamburger patties) Offal (specify type) 1 2 7
	h.	Other meat

49. How do you *prefer* the following meat to be cooked?

INTERVIEWER NOTE: Raw: Not cooked at all Rare: Mostly red

Medium: Pink through out Well done: Brown through out

INTERVIEWER NOTE:

If participant answers Medium/Rare select the rarer option.. e.g Rare

		<u>Kaw</u>	Kare	<u>Medium</u>	<u>we</u> n done
a.	Chicken/Poultry		2	3	4
b.	Beef/Veal				
		1	2	3	4
c.	Pork	1	2	3	⊢ ⁴
d.	Lamb	1	2	3	4
e.	Hamburgers	1	2	3	4
f.	Minced meat	1	2	3	4
			Ш-		

E	GENE:	$D \Lambda I$	KITC	HEN	DD A	CTICES
12.	OENE.	\mathbf{n}	NIIC	TLIN.	$\Gamma \Lambda A$	CHUES

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Е. С	GENERAL KITCHEN PRACTICES					
R	will now ask you several questions about the way food is usually prepared in your home. emember, your participation is voluntary and you do not have to answer any of the questions if ou don't want to."					
50.	How many times per week do you cook for members of your household? INTERVIEWER NOTE:					
	This section around food prepared in the home refers to the person answering the survey (not necessarily the case or control)					
	0					
	1-5					
	>5					
	Don't know/Not sure					
51.	Did you handle or prepare any raw meats in the kitchen in the 7 days before your					
	diarrhoea began?					
	INTERVIEWER NOTE:					
	Refers to the person answering the survey					
	Yes					
	No					
	Don't know/Not sure					
52.	Did you handle or prepare raw chicken meat or chicken offal in the 7 days before your diarrhoea began? INTERVIEWER NOTE: Refers to the person answering the survey					
	Yes					
	No					
	Don't know/Not sure					
Interv	iewer Note: If person answered "No" to both Q.51 and Q.52 then skip to Q.57					
53.	After a knife is used to cut raw meat or poultry, which of the following options do you usually do? INTERVIEWER NOTE:					
	Refers to the person answering the survey					
	Continue using the knife as is					
	Rinse the knife before continuing to cook 2					
	Wipe the knife before continuing to cook					
	Wash the knife with detergent before continuing					
	Change to another knife					
	Other)				
	No one prepares meat					
ead	Don't know/Not sure					

54.	After a cutting board is used to cut raw meat or poultry, which of the following options do you usual do?
	INTERVIEWER NOTE:
	1. Does not matter if water is hot or cold
	2. Refers to the person answering the survey
	Continue using the cutting board as is
	Rinse the cutting board before continuing to cook
	Wipe the cutting board before continuing to cook
	Wash the cutting board with detergent before continuing
	Change to another cutting board.
	Other
	Specify ()
Don't read	Don't know/Not sure
55.	After handling raw meat or poultry in the kitchen, which of the following would you usually do before continuing to cook?
	INTERVIEWER NOTE:
	Refers to the person answering the survey
	Wipe hands
	Quickly rinse hands under a running tap 2
	Wash hands with soap and water
	Other
	Specify ()
	Don't do anything about hands
Don't read	Don't know/not sure
E (After weaking hands during food proposation, what would you republy dur
56.	After washing hands during food preparation, what would you usually dry your hands on?
	INTERVIEWER NOTE:
	Refers to the person answering the survey
	<u> </u>
	Paper towel
	Sponge/cloth
	Tea-towel /hand towel
	Apron
	Don't dry hands
	Other
D 24 1	Specify (
Don't read	Don't know/Not sure
57.	In the past 3 months, has anyone in the household cook meat on a BBQ?
	Yes
	N- C- 4- O 50
	No

	BMJ Open	Page 3‰f 42 ≧
58.	After cooking on the BBQ, where would the cooked meat most likely be placed?	Open: firs
	Back on the same container. Back on the same container after it has been rinsed with water 2	Q MJ Open: first published as 10.1136/bmjopen-2018-026630 on 22 December 2018. Downloaded from http://bmjopen.bmj.com/ on April 10, 2024 by guest. Protected by copyright.
	Back on the same container after it has been wiped off with a towel	ıs 10.11
	Back on the same container, after the container has been washed with soap and water	36/bmjop
Don't read	On a different container 5 Other 6 Don't know/not sure 7 Specify ()))
		630 on 22
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Is any pet less than

F. ANIMAL AND PET EXPOSURE

Interviewer Note:

"The next few questions are about contact with animals in the 7 days before your diarrhoea began."

59. During this time, did you keep or care for any of the following animals as pets?

INTERVIEWER NOTE:
Not to include one off contact

							o months ola:							
		Yes	;	No	_	DK	/NS		Ye	S	No	DK	/NS	
a.	Cat		1		2		7			1		2		7
b.	Dog		1		2		7			1		2		7
c.	Chickens		1		2		7			1		2		7
d.	Other birds		1		2		7			1		2		7
e.	Other		1		2		7		Sp	ecify				
f.	Do not keep any pets		7 (Go t	o Q.	67								

60.	Do you feed your cat raw meat or bones?			
	Yes		Specify ()
	No		Go to Q. 62	(eg. chicken, beef, kangaroo, lamb etc
	Don't know/Not sure	7	Go to Q. 62	

If person answered No/Don't know to Cat then skip to question 62

Intervie	ewer Note:	If person answered No/Don't know to Dog then skip to question	65
62.	Do you feed	your dog raw meat?	

63.	Do you feed your dog raw bones?				
	Yes	Г	1 1	Specify (
	No		2		
	Don't know/Not sure		7		

		ВМЈ О	oen		Page 46of 42 ≧
					Ј Оре
Intervi	iewer Note:	If person answered No/Don't kn	ow to quest	ions 62-63 then skip to question 65	n: first
64.	How often d	oes your dog get fed raw meat or	bones?		publis
	Weekly Monthly Less often	Not sure	2 3 4		9 MJ Open: first published as 10.1136/bmjopen-2018-026630 on eg P
65.	Did you get	any of your pets in the 4 weeks be	efore your	diarrhoea began?	en-20
	No	Not sure	— •	Pet(s) ()18-026630 on 2
66.	-	your own pets ill with diarrhoea		ys before your diarrhoea began?	22 De
	No	Not sure	1 2 7	Pet(s) (cember 2018.
67.		s before your diarrhoea began, di nure (eg. changing litter boxes or			. Downloade
	No	Not sure	H ,	Pet(s) (December 2018. Downloaded from http://b
68.	Yes	on a farm/hobby farm including a	1 2 7	on acreage 5 acres or over?	mjopen.bmj.com/ o
69.	In the 7 day	s before your diarrhoea began, di	d von visit	a farm or netting zoo?	n April
0).	•			Specify () 10, 1
	No	Not sure	2 7	(eg. private farm, commercial farm, pettin	njopen.bmj.com/ on April 10, 2024 by guest. Protected by copyright
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4. DEMOGRAPHICS

"I would now like to ask you a few final questions. Remember, your participation is voluntary and you do not have to answer any of the questions if you don't want to."

70.	Is any language other than English spoken in your household?
	Yes
	No
Don't read	Don't know/Not sure
Don't read	Refused9
71.	Are you of Aboriginal or Torres Strait Islander origin?
	No
	Aboriginal
	Torres Strait Islander
	Both
Don't read	Don't know/Not sure
Don't read	Refused9
72.	Which of the following places best describe where you live?
,	Inner city or when erec
	Cyleyellore area
	Town
	Rural or remote area community
	Rural or remote area farm or property
Don't read	Don't know/Not sure
Don't read	Refused
Interv	viewer Note: See definitions below.
	ty area:housing close to the centre of a major/capital city
Suburba	an area:housing area further from the centre of the city, which is characterise
TED.	the region being primarily a self-contained residential district.
	r remote area community:community under 2000 r remote area farm or property
<u>Kurar or</u>	Temote area rann or property
73.	Does your occupation involve any of the following?
	Working with raw meat
	(eg. restaurants, butchery, abattoir etc.) Working with animals
	Working with animals
	Other type of occupation
	Retired
	CASE not of working age

Don't know/Unsure....

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		Ј Оре
		en: fire
74.	What is the highest level of education reached by <u>anyone</u> in your household?	st pu
	Schooling to year 10 or below	blished as
Don't read Don't read	University degree—Undergraduate	10.1136/bmjop
		en-2
d	Now I am going to read you a list of income categories. Please stop me when a category be escribes your total household income, before taxes, in the last financial year? That is the total figure all household members." Last year the total income for your household was?	018-026630 on
Don't read	Less than \$25,000	MJ Open: first published as 10.1136/bmjopen-2018-026630 on 22 December 2018. Downloaded from http://b ct con st
Don't read	Refused	Dow
76.	As part of this research we are planning to do a follow-up study. Would you be happy for u you in ~6 months' time?" Yes	
Interv	iewer Note: If person answered No to Q 76 then skip to the end of the questionnaire	njope
Titterv	Details required:	en.bn
	Name:	nj. cor
	Phone number:	n/ or
	Email address:	n Apr
"	Γhat's my last question. Thank you very much for your time and cooperation."	ii 10,
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	Interviewer initials	est. P
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	Fair Average 2 3	
	Good For peer review only http://bmjopen.bmj.com/site/about/guidelines.xhtml Excellent	24

BMJ Open

Investigating locally-relevant risk factors for Campylobacter infection in Australia: protocol for a case-control study and genomic analysis

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SCHOLARONE™ Manuscripts

Investigating locally-relevant risk factors for *Campylobacter* infection in Australia: protocol for a case-control study and genomic analysis

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ABSTRACT

Introduction:

The CampySource project aims to identify risk factors for human *Campylobacter* infection in Australia. We will investigate locally-relevant risk factors and those significant in international studies in a case-control study. Case isolates and contemporaneous isolates from food and animal sources will be sequenced to conduct source attribution modelling, and findings will be combined with the case-control study in a source-assigned analysis.

Methods and analysis:

The case-control study will include 1,200 participants (600 cases and 600 controls) across three regions in Australia. Cases will be recruited from campylobacteriosis notifications to health departments. Only those with a pure and viable *Campylobacter* isolate will be eligible for selection to allow for whole genome sequencing of isolates. Controls will be recruited from notified cases of influenza, frequency matched by sex, age group and geographical area of residence. All participants will be interviewed by trained telephone interviewers using a piloted questionnaire.

We will collect *Campylobacter* isolates from retail meats and companion animals (specifically dogs), and all food, animal and human isolates will undergo whole genome sequencing. We will use sequence data to estimate the proportion of human infections that can be attributed to animal and food reservoirs (source attribution modelling), and to identify spatial clusters and temporal trends. Source-assigned analysis of the case-control study data will also be conducted where cases are grouped according to attributed sources.

Ethics and dissemination:

Human and animal ethics have been approved. Genomic data will be published in online archives accompanied by basic metadata. We anticipate several publications to come from this study.

KEYWORDS

Campylobacter, case-control study, risk factors, Australia, whole genome sequencing, source attribution, source-assigned analysis

ARTICLE SUMMARY

Strengths and limitations of this study

- Case-control study is well-powered to identify locally-relevant risk factors.
- Linking genomic data to the case-control study strengthens the analysis by enabling source attribution and source-assigned analyses to be conducted.
- Case-control questionnaire questions are being validated in a separate study, demonstrating the reliability of participant recall.
- Potential reporting bias due to inaccurate recall of study participants.
- Case-control study lacks efficiency for risk factors with high levels of exposure in the study population.

INTRODUCTION

Campylobacter infection is the most commonly notified cause of foodborne gastroenteritis in Australia, ¹⁻³ as well as a leading cause of bacterial gastroenteritis world-wide. ⁴ At the introduction of Australia's National Notifiable Diseases Surveillance System (NNDSS) in 1991 the incidence rate of notified campylobacteriosis cases was 79.1/100,000 population, ⁵ and despite notification rates plateauing in recent years, incidence had risen to 139.7/100,000 population in Australia in 2015, ⁵ with an estimated 10 cases for every notified case within the community. ⁶ By comparison, the incidence rate of campylobacteriosis in New Zealand in 2014 was 150.3/100,000 population, ⁷ with an estimated 10-30 cases in the community for every notified case. ⁸ Campylobacter notification rates in Australia and New Zealand are still among the highest in the world across high-income countries. Most countries in the European Union consistently report annual campylobacteriosis notification rates below 100/100,000 population. ²

Two species of *Campylobacter—Campylobacter jejuni* and *C. coli*—contribute to approximately 95% of human campylobacteriosis. ⁹ These *Campylobacter* species are commonly detected in sewage and surface water, ¹⁰ reside in the gastrointestinal tract of birds and animals, ¹¹ and are frequently found in raw meat, particularly poultry, and raw milk. ¹² ¹³ Campylobacteriosis is mostly foodborne, with an estimated 77% of cases transmitted via food consumption in Australia. ¹⁴ ¹⁵ Direct and indirect zoonotic transmission can occur via animal contact (direct) or faecally-contaminated water or environments (indirect). Person-to-person transmission is considered rare. ¹⁶ The majority of cases are thought to be sporadic, with outbreaks less commonly detected. ¹⁷ Most outbreaks are linked to the consumption of poultry, raw milk, or contaminated water. ¹⁷ ¹⁸

Targeted control of foodborne bacterial pathogens generally depends on identification of sources and routes of transmission. Since *Campylobacter* are ubiquitous in the environment and most cases are sporadic, identifying sources is difficult. Source attribution methods require isolation of strains from reservoirs to compare *Campylobacter* strain diversity in foods and animals to that in human infections. Beef, sheep and pig meat have a lower prevalence of *Campylobacter* contamination than chicken meat (<5% to 14%), ¹⁹⁻²¹ but a higher prevalence is found in animal offal such as liver, ²² thus making offal a valuable source of host-associated strains of *Campylobacter* in low-prevalence meats.

STUDY RATIONALE

In the United States, evidence from case-control studies have led to policy change, including changes to chicken slaughtering techniques. The incidence of human *Campylobacter* infection has declined in the US since this policy was introduced in 1997.²³ More recently, evidence from source attribution analyses in New Zealand has led to the development of poultry production policies and practices aimed at reducing the risk of *Campylobacter* transmission via poultry food products.²⁴ New Zealand has seen a 74% reduction in the number of campylobacteriosis cases attributed to poultry in the region, as well as a 54% reduction in cases overall.²⁵

Source attribution modelling enables us to determine which foods and animals are the most likely sources of infection with each *Campylobacter* strain type, and the proportion of cases attributed to each source. This can be done with simple proportional similarity index (PSI) calculations, or by using more complex models.²⁴ Source attribution also allows for human campylobacteriosis cases to be grouped by potential source, increasing the specificity of risk factor analyses. These source-assigned analyses combine the epidemiological information gained through the traditional case-control study with source attribution modelling to provide greater explanatory power to investigate locally-relevant risk factors.

OBJECTIVES

This study aims to:

- Identify dietary, environmental and behavioural risk factors for Campylobacter infection in Australia
- 2. Strengthen the epidemiological evidence for previously identified risk factors in Australia
- 3. Identify strain-specific risk factors for infection using Whole Genome Sequencing (WGS) data from case isolates

HYPOTHESES

We will test several hypotheses regarding specific risk factors for *Campylobacter* infection in Australia. The hypotheses are based on exposures which have previously been identified as risk factors for *Campylobacter* infection in Australia as well as internationally.

We hypothesise that:

- 1. Persons who consume undercooked meats, particularly chicken, are at increased risk of infection.
- 2. Persons who consume offal are at increased risk of infection.
- 3. Persons who own companion animals (especially pupples) are at increased risk of infection.
- 4. Poor food hygiene and handling practices in the home increase the risk of infection.
- 5. Most human infections will be attributed to consumption of chicken meat.
- 6. There will be a high level of genetic diversity amongst *Campylobacter* strains.

STUDY DESIGN

We will conduct a case-control study including genomic testing over a two-year period in three sentinel sites: the state of Queensland (QLD), the Australian Capital Territory (ACT), and Hunter New England (HNE) region of New South Wales (Figure 1). Sporadic cases of culture-positive *Campylobacter* infection will be identified either through state notifiable disease registers, from local pathology service databases or local notification databases. An isolate from each case will be paired with epidemiological data from the case interview. One control will be recruited for each case who participates in the study, with trained interviewers conducting telephone interviews with both cases and controls. Participants will be interviewed using a questionnaire that has been specifically designed to collect information on known potential risk factors. This questionnaire will include a selection of questions being validated in a separate study (Liana Varrone, Validation of questions designed for gastroenteritis investigation). For cases, the questions will cover the seven days prior to the onset of illness, while controls will be questioned on the seven days prior to interview. Meanwhile, *Campylobacter* isolates will also be collected from food and animal samples. All human and non-human isolates will undergo whole genome sequencing for comparison in source attribution modelling. Data for this study will be collected from 1st March 2017 to 1st March 2019.

Figure 1. Map of Australian states and territories, showing the Hunter New England region.

Patient and public involvement

To develop the study, we engaged state and territory health departments, food safety agencies and industry to establish research questions and methods. The process involved a dedicated workshop, followed by teleconferences and an iterative process of drafting study documentation. We also established a reference panel, which includes representatives from senior levels of government and industry bodies. No patients or other members of the public were involved in the development of this study.

STUDY POPULATION

The three sentinel sites cover a population of approximately 6.1 million people. Based on notification and diagnostic pathology data, we expect approximately 8,650 *Campylobacter* cases to be notified across these sites during the study period.

DEFINITION AND SELECTION OF CASES

Case definition

We define a case as a person from any of the three participating sites with a recent history of acute diarrhoea and a culture-positive stool result for *Campylobacter*.

SAMPLE SIZE

We used risk factor prevalence data from a previous national *Campylobacter* case-control study in 2001/2002 to estimate sample size for this study. For example, the prevalence of chicken consumption among controls in 2001/2002 was 80%. A sample size of approximately 1,040 subjects (520 cases; 520 controls) would enable the study to detect an association between chicken consumption and illness with an odds ratio of 1.6, at 80% power and α = 0.05, as reported in the previous study. Sample size estimates for other potential risk factors are listed in Table 1.

Table 1. Sample size estimates for an unmatched case-control study

Risk factor	Prevalence of exposure among controls (%)	Prevalence of exposure among cases (%)	Odds ratio	No. of required study subjects
Beef	78	85	1.6	960
Pork	52	60	1.4	1130
Lamb	42	50	1.4	1120
Chicken	80	87	1.6	1040
Offal	2.0	5.0	2.6	1154
Puppies	2.1	5.4	2.7	1040

80% power and $\alpha = 0.05$

From these calculations, we estimate that a study of 1,200 subjects (600 cases; 600 controls) will adequately detect significant associations of these magnitudes for potential risk factors of interest.

Queensland and Hunter New England sites will each enrol at least 250 cases into the study, while ACT will enrol at least 100 cases. Based on the previous Australian case-control study, ²⁶ we expect approximately 80% of selected notified cases to be eligible and participate in the study (Table 2).

Table 2. Sampling method for cases in each site

State	Expected number of notified cases during study period	Estimated cases from participating pathology laboratory	Culture +ve cases	Sequential sampling of notified cases	Total no. of cases	Expected no. to be recruited (~ 80% participation rate)
QLD	7000	2800 (40%)	1260 (45% in QLD)	Select every 4 th case	315	250
ACT	600	130	130	Include all notified cases	130	100
NSW (Hunter New England)	~1050	313	313	Include all notified cases	313	250
Total	8650	3243	1703		758	600

In Queensland, we will obtain cases from one private pathology provider reporting approximately 40% of the state's *Campylobacter* notifications. We estimate that this provider will notify 2,800 cases during the study period with an estimated 45% of these being culture-positive (1,260 notified cases). In ACT, approximately 600 *Campylobacter* notifications are expected during the study period; 130 are expected from the participating pathology laboratory. In Hunter New England, approximately 1,050 *Campylobacter* notifications are expected during the study period; 313 of these notifications will be from the participating pathology laboratory.

Enrolment of Cases

We will enrol all cases who meet the eligibility criteria (Table 3). Each site will check for new notifications of culture-positive *Campylobacter* infection daily, with only culture-positive *Campylobacter* cases eligible for this study. If a case refuses to participate in the study, we will select a subsequent case for inclusion. Enrolment of cases will depend on consent from the patient, or in the event of a child aged less than 18 years, consent from either one of the parents or the child's guardian. We will interview cases as soon as possible by telephone, preferably within two weeks of notification from the laboratory. It will be at the parent's or guardian's discretion as to whether a

child aged between 15 and 17 years is interviewed directly. The parent or guardian will be interviewed for cases aged less than 15 years.

Table 3. Eligibility criteria for cases and controls

Criteria	Cases	Controls
Had diarrhoea (≥3 loose bowel movements in	Include	Exclude
24hrs)		
Known date of illness onset	Include	N/A
Household members positive for Campylobacter	Exclude	Exclude (4 weeks prior to
in 4 weeks prior to onset of illness		interview date)
Household members experiencing diarrhoea in 4	Exclude	Exclude (4 weeks prior to
weeks prior to onset of illness		interview date)
Travelled outside of Australia in 2 weeks prior to	Exclude	Exclude (2 weeks prior to
onset of illness		interview date)
Travelled interstate for the entire 2 weeks prior	Exclude	Exclude (2 weeks prior to
to onset of illness		interview date)
Can't speak English	Exclude	Exclude
Not able to answer questions for some other	Exclude	Exclude
reason (e.g. intellectually disabled)		
Not contactable after 6 telephone attempts	Exclude	Exclude
Live outside the catchment areas	Exclude	Exclude
Do not have a telephone number available for	Exclude	Exclude
their primary residence, or a mobile phone		
An enteric pathogen other than Campylobacter	Exclude	N/A
was isolated/detected in their stool (excluding		
Blastocystis hominis and Dientamoeba fragilis)		

DEFINITION AND SELECTION OF CONTROLS

We will recruit controls from notified cases of influenza, frequency matched by sex, age group and geographical area of residence by Statistical Area Level 4 (SA4). These controls will be selected with a delay of at least six months from their influenza infection to ensure that controls have returned to eating their customary diet.

Each participating site (QLD, ACT or HNE) will establish a database of controls (previous influenza cases). All cases of influenza notified to the health department in each site between 1st January and 31st December 2017 will be entered into this control database. The age bands are 0-4 years, 5-14 years, 15-34 years, 35-54 years, 55-74 years, and ≥75 years. An appropriate control will be randomly selected from the database within 30 days of interview of the notified case.

Case and control recruitment

Interviewers trained in computer-assisted telephone interviewing (CATI) will conduct telephone interviews. A maximum of six attempts will be made to contact any one case or control, with no more than three attempts in any one day. Three calls will be attempted between 9:00am and 3:59pm, and three attempts between 4:00pm and 8:00pm. A text message will be sent to the potential participant after three failed call attempts, indicating that Public Health is trying to contact them. This protocol will be continued until the person is enrolled or excluded.

QUESTIONNAIRES

We will use specific case and control questionnaires for all participants (see Appendix 1). Cases will be asked additional questions about the clinical course of their illness and treatment. Interviewers will ask identical questions regarding exposures such as foods consumed, dining locations, water sources, domestic food handling techniques and exposure to animals of cases and controls. Questions on foods consumed, dining locations, water consumed, animal and pet exposures will be asked based on a seven-day history. Questions on international travel will be asked based on a two-week history. Antibiotic and antacid consumption, immunosuppressive treatment and household history of diarrhoea will be based on a four-week history. Questions on food handling and general kitchen practices will be based on usual practices rather than recent history. Demographic information will be collected from cases and controls. Contact information required to conduct interviews will be stored in a password-protected Excel document with only those needing to contact individuals given access. Piloted questionnaires were modified to remove repetitions, improve clarity, and to ensure that interviews could be conducted within 20 minutes.

DATA HANDLING & RISK FACTOR ANALYSIS

We will undertake descriptive reporting of campylobacteriosis incidence by person, place and time. We will also describe the severity of symptoms, treatment, and burden of illness.

Risk factor analysis will involve the examination of two-by-two contingency tables with chi square or exact tests to determine the presence of univariable associations between variables and disease. To measure the strength of an association, we will estimate odds ratios and calculate 95% confidence

intervals in a univariable analysis, followed by multivariable logistic regression modelling to adjust for potential confounders. Risk factors selected for inclusion in the regression model will include age, season and geographic area, variables with a significant univariable association with disease, and variables with a P-value ≤ 0.25 that are biologically plausible and of interest to the research team.

LABORATORY ANALYSES

Human samples

As outlined in Table 2, it is expected that 250 human isolates from Hunter New England, 250 from Queensland, 100 from Victoria and 100 from ACT will be sequenced. The initial isolation and confirmation of *Campylobacter* infection will be performed locally in each State/Territory. Only samples with a pure and viable culture will undergo WGS.

Animal and food samples

We will collect samples from chicken meat (covering the two production methods of continually housed and free range/housed), beef, lamb, pork, and from pet dogs. Given low prevalence of *Campylobacter* in meats other than chicken, samples will be collected from offal (preferably liver) from bovine, ovine and porcine sources to ensure sufficient positive samples are obtained for the study. Given the rising importance of chicken liver pate as a source of outbreaks in Australia, ²⁷ chicken offal will also be sampled. Sample sizes by source are based on data from two states to ensure 50 positive samples per food source, and 30 samples in companion animals (Table 4). We will also contact veterinary clinics and teaching hospitals to ensure sufficient *Campylobacter*-positive samples from dogs. Water samples have been omitted from the genomic aspect of this study due to logistical constraints in sampling untreated water sources across the large geographical area involved in this study, and the complexity of designing an appropriate sampling frame. As there is a lack of evidence implicating municipal drinking water as sources of *Campylobacter* infection in Australia²⁶ we excluded water sampling from this study.

Table 4. Sampling to ensure 50 isolates per food source and 30 isolates from companion animals

	Foods						Animals	
	Chicken			Beef	Lamb	Pork	Dogs	Total
	Continually	Free-	Offal	Offal	Offal	Offal		
	housed	range						
Assumed	0.7	0.7	0.7	0.14	0.6	0.22	0.2	
prevalence								
Samples	72	72	72	286	100	272	150	1041
required								
Positive	50	50	50	40	60	60	30	330
isolates								

The initial isolation and confirmation of *Campylobacter* will be performed locally at laboratories in each State/Territory, with isolates forwarded to the Microbiological Diagnostic Unit Public Health Laboratory for WGS, except Queensland isolates which will be sequenced at Queensland Health. To detect seasonal and temporal variation in *Campylobacter* genetic types, 1041 non-human samples (estimated to produce 330 *Campylobacter* isolates) will be collected over a period of one year in Queensland, and two years in New South Wales. To assess latitudinal variation in chicken meat samples across eastern Australia, 105 chicken samples (70 chicken meat and 35 chicken offal) will be collected over a six-month period in Victoria. Food samples will be collected monthly from retail premises, using protocols from surveys undertaken in 2014 by partner organisations, with a pilot of 30 isolates in Queensland.

We will also collect an additional 20-30 human isolates from four additional Australian jurisdictions not participating in this case-control study to undergo WGS. This will be done over a two-month period that overlaps with the case-control study sample collection, and is planned to help inform the generalisability of the case-control study.

SEQUENCING AND SEQUENCE DATA PROCESSING

Campylobacter isolates selected for sequencing will be repurified on solid medium and a single colony selected for preparation of genomic DNA. A sequencing library will be prepared from the genomic DNA for sequencing on the Illumina sequencing platform (MiSeq or NextSeq). A sample of the selected colony will be regrown and cryopreserved (resuspended in liquid medium supplemented with 10% Glycerol and stored at -80°C). In some cases, Campylobacter enrichment cultures will be cryopreserved to enable future investigation of the genetic diversity of Campylobacters present. The short-read, paired end dataset produced by the Illumina Instrument from the genomic DNA of each isolate will be processed to produce a draft genome sequence for the isolate using a de novo assembler such as MEGAHIT.²⁸ The draft genome sequence will be annotated using Prokka.²⁹ We will use the draft genome sequence to perform the initial sub-species classification by deriving a multilocus sequence type (MLST) using the "Campylobacter jejuni/coli" typing scheme (pubmlst.org). Again, using the draft genome sequence, further typing e.g. virulence factors (http://www.mgc.ac.cn/VFs/) or antimicrobial resistance genotype (https://cge.cbs.dtu.dk/services/ResFinder/) will be performed using Abricate (https://github.com/tseemann/abricate). We will perform comparative genomics to examine the genetic relationships between selected subgroups of isolates in more detail using Nullarbor (https://github.com/tseemann/nullarbor).

SOURCE ATTRIBUTION MODELLING

We will analyse the epidemiological data within designated MLST groups or other typing groups derived from the genomic sequence data. Source attribution modelling and source-assigned analyses will be conducted.

Source attribution models combine typing data from isolates from food, animal and humans to estimate the proportion of human infections that can be attributed to animal and food reservoirs.³⁰ ³¹ Once inferred MLSTs have been ascertained, the proportional similarity index²⁵ will be used to assess similarities by source. We will then undertake source attribution analyses by adapting the asymmetric island model which has previously been applied to MLST data^{25 32} using Markov Chain Monte Carlo (MCMC) methods³³ implemented using the free software WinBUGS.³⁴ These methods will first be applied to MLST data extracted from whole genome sequences (the aforementioned "inferred MLSTs"), and then compared to structured phylogenetic modelling approaches^{35 36} that provide scope to infer inter-host transmission.

We will then group cases according to putative source based on these source attribution methods.³⁷ For example, all isolates attributed to chicken will be grouped together, regardless of differing strains. These cases attributed to chicken will then be compared to all controls in a risk factor analysis to produce a source-assigned analysis.

SPATIAL CLUSTERS AND TEMPORAL TRENDS

We will use newly-designated WGS-based MLSTs to assess heterogeneity in isolates from food sources and companion animals in Queensland and New South Wales, and in isolates from chicken meat and humans across Queensland, New South Wales, Victoria and ACT. A two-year sampling framework in New South Wales, one year of sampling in Queensland, and previous survey work in these states will allow us to assess the extent of seasonal and temporal trends. Postcode-level data associated with human illnesses will be used to detect space-time clusters using a scan statistic implemented in the free software SaTScan, at the Statistical Area 1 level. We will use a retrospective space-time permutation model to detect high risk clusters by comparing the observed number of illnesses to the expected number in that geographic zone and time-period. 19

STUDY LINKAGES AND COLLABORATIONS

The CampySource Project Team comprises three working groups and a reference panel. The working groups focus on: food and animal sampling, epidemiology and modelling, and genomics. The reference panel includes expert representatives from government and industry.

The study is supported by the following partner organisations: the Australian National University, Massey University, University of Melbourne, Queensland Health, Queensland Health Forensic and Scientific Services, New South Wales Health, Hunter New England Health, Victorian Department of Health and Human Services, Food Standards Australia New Zealand, Commonwealth Department of Health and AgriFutures Australia – Chicken Meat Program.

CampySource is also supported by collaboration with the following organisations: ACT Health, Sullivan Nicolaides Pathology, University of Queensland, Primary Industries and Regions South Australia, Department of Health and Human Services Tasmania, Meat and Livestock Australia, and New Zealand Ministry for Primary Industries.

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The CampySource Project Team consists of: Nigel P. French, Massey University, New Zealand; Mary Valcanis, The University of Melbourne; Dieter Bulach, The University of Melbourne; Emily Fearnley, The Australian National University; Russell Stafford, Queensland Health; John Bates, Queensland Health; Trudy Graham, Queensland Health; Keira Glasgow, Health Protection NSW; Kirsty Hope, Health Protection NSW; Arie H. Havelaar, The University of Florida, USA; Joy Gregory, Department of Health and Human Services, Victoria; James Flint, Hunter New England Health; Simon Firestone, The University of Melbourne; James Conlan, Food Standards Australia New Zealand; James J. Smith, Queensland Health; Sally Symes, Department of Health and Human Services, Victoria; Barbara Butow, Food Standards Australia New Zealand; Liana Varrone, The University of Queensland; Linda Selvey, The University of Queensland; Deborah Denehy, ACT Health; Radomir Krsteski, ACT Health; Natasha Waters, ACT Health; Kim Lilly, Hunter New England Health; Julie Collins, Hunter New England Health; Tony Merritt, Hunter New England Health; Joanne Barfield, Hunter New England Health; Ben Howden, The University of Melbourne; Kylie Hewson, AgriFutures Australia – Chicken Meat Program; Laura Ford, The Australian National University; Liz Walker, The Australian National University; Cameron Moffatt, The Australian National University; Martyn Kirk, The Australian National University; and Kathryn Glass, The Australian National University.

While undertaking studies, LV is supported through an Australian Government Research Training Program (RTP) Scholarship.

DECLARATIONS

Ethics approval and consent to participate

Informed Consent

A suitably trained interviewer will inform potential participants about the purpose, methods and demands of the study. We will obtain verbal consent from all study participants or their guardians.

Persons aged 18 years and older will be interviewed following informed consent. It will be at the parent's or guardian's discretion as to whether a child aged between 15 and 18 years is interviewed directly, following informed parental/guardian consent. Parents/guardians will be interviewed for cases aged less than 15 years, after informed consent is obtained.

Confidentiality

All information and identifiers will be kept confidential. Names and personal identifiers will be collected and entered into computer records but will be password protected. No personal identifiers will be included in any published materials relating to this study. All hard copy questionnaires containing patient identifiers will be stored in locked filing cabinets in a secure location to which only study investigators and interviewers will have access.

Risks and Benefits

Participants will be informed there are no individual benefits associated with the study and that participation is voluntary. Failure to participate or a withdrawal of participation will not affect any future treatment. There is also no risk to the patient, and the only cost is time spent – approximately 20 minutes – being interviewed. They may refuse to answer any of the questions or stop at any time.

Animal Ethics

All procedures involving live animals will be performed in accordance with a protocol approved by the University of Melbourne's Animal Ethics Committee (ethics ID: 1714156).

Consent for publication

Not applicable.

Availability of data and materials

The Illumina read sets produced as part of this study will be published at INSDC (Sequence Read Archive (DDJB/NCBI) or the European Nucleotide Archive (EMBL-EBI))

Competing interests

No authors have any competing interests to declare.

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MDK is supported by a National Health and Medical Research Council Fellowship (GNT1145997).

Author contributions

MDK conceived the original idea for this study. All authors contributed to the study design and analysis plan. LV and RJS wrote the first draft with contributions from all authors. LF was heavily involved in determining timing and logistics in and between all sites. KL assisted in questionnaire design and flow. DB developed the bioinformatics analysis protocol. LV, RJS, LS, MDK and KG were involved in multiple revisions. The final version of the manuscript was approved by all authors.

REFERENCES

- 1. The OzFoodNet Working Group. Monitoring the incidence and causes of diseases potentially transmitted by food in Australia: Annual report of the OzFoodNet network, 2011. *Annual Report* 2011; 39(2). http://www.health.gov.au/internet/main/publishing.nsf/Content/cda-cdi3902-pdf-cnt.htm/\$FILE/cdi3902g.pdf.
- 2. Kaakoush NO, Castaño-Rodríguez N, Mitchell HM, et al. Global Epidemiology of Campylobacter Infection. *Clin Microbiol Rev* 2015;28(3):687-720. doi: 10.1128/cmr.00006-15
- 3. Ford L, Kirk MD, Glass K, et al. Sequelae of Foodborne Illness Caused by 5 Pathogens, Australia, Circa 2010. *Emerg Infect Dis* 2014;20(11):1865. doi: 10.3201/eid2011.131316
- 4. Fernandes AM, Balasegaram S, Willis C, et al. Partial Failure of Milk Pasteurization as a Risk for the Transmission of Campylobacter From Cattle to Humans. *Clin Infect Dis* 2015;61(6):903-09. doi: 10.1093/cid/civ431
- 5. Commonwealth of Australia. National Notifiable Diseases Surveillance System, 2018.
- 6. Hall G, Yohannes K, Raupach J, et al. Estimating Community Incidence of Salmonella, Campylobacter, and Shiga Toxin–producing Escherichia coli Infections, Australia. *Emerg Infect Dis* 2008;14(10):1601-09. doi: 10.3201/eid1410.071042
- 7. The Institute of Environmental Science and Research Ltd. Notifiable Diseases in New Zealand: Annual Report 2014. Porirua, New Zealand, 2015.
- 8. Cressey P, Lake R. Estimated incidence of foodborne illness in New Zealand: application of overseas models and multipliers, 2011.
- 9. Lastovica AJ, Allos BM. Clinical Significance of Campylobacter and Related Species Other Than Campylobacter jejuni and Campylobacter coli. In: Nachamkin I, Szymanski CM, Blaser MJ, eds. Campylobacter, Third Edition: American Society of Microbiology 2008.
- 10. Fricker CR, Park RWA. A two-year study of the distribution of 'thermophilic' campylobacters in human, environmental and food samples from the Reading area with particular reference to toxin production and heat-stable serotype. *J Appl Bacteriol* 1989;67(6):477-90. doi: 10.1111/j.1365-2672.1989.tb02530.x
- 11. Blaser MJ, Taylor DN, Feldman RA. Epidemiology of Campylobacter jejuni Infections. *Epidemiol Rev* 1983;5(1):157-76.
- 12. Stern NJ, Hernandez MP, Blankenship L, et al. Prevalence and Distribution of Campylobacter jejuni and Campylobacter coli in Retail Meats. *J Food Prot* 1985;48(7):595-99.
- 13. Wood RC, MacDonald KL, Osterholm MT. Campylobacter enteritis outbreaks associated with drinking raw milk during youth activities: A 10-year review of outbreaks in the United States. *JAMA* 1992;268(22):3228-30. doi: 10.1001/jama.1992.03490220072031
- 14. Vally H, Glass K, Ford L, et al. Proportion of illness acquired by foodborne transmission for nine enteric pathogens in Australia: an expert elicitation. *Foodborne Pathog Dis* 2014;11(9):727-33. doi: 10.1089/fpd.2014.1746
- 15. Kirk M, Ford L, Glass K, et al. Foodborne illness, Australia, circa 2000 and circa 2010. *Emerg Infect Dis* 2014;20(11):1857-64. doi: 10.3201/eid2011.131315
- 16. Heymann DL, editor. *Control of Communicable Diseases Manual*. 19th ed. Maryland: United Book Press, Inc., 2008.

- 17. Pebody RG, Ryan MJ, Wall PG. Outbreaks of campylobacter infection: rare events for a common pathogen. *Commun Dis Rep CDR Rev* 1997;7(3):R33-7. [published Online First: 1997/03/07]
- 18. Tauxe Robert V, Hargrett-Bean N, Patton CM. Campylobacter Isolates in the United States, 1982-1986. *MMWR Surveill Summ* 1988;37(S S-2):1-13.
- 19. Sammarco ML, Ripabelli G, Fanelli I, et al. Prevalence and Biomolecular Characterization of Campylobacter spp. Isolated from Retail Meat. *J Food Prot* 2010;73(4):720-28.
- 20. Whyte P, McGill K, Cowley D, et al. Occurrence of Campylobacter in retail foods in Ireland. *Int J Food Microbiol* 2004;95(2):111-18. doi: 10.1016/j.ijfoodmicro.2003.10.018
- 21. Wong TL, Hollis L, Cornelius A, et al. Prevalence, Numbers, and Subtypes of Campylobacter jejuni and Campylobacter coli in Uncooked Retail Meat Samples. *J Food Prot* 2007;70(3):566-73.
- 22. Noormohamed A, Fakhr MK. A Higher Prevalence Rate of Campylobacter in Retail Beef Livers Compared to Other Beef and Pork Meat Cuts. *Int J Environ Res Public Health* 2013;10(5):2058.
- 23. Samuel M, Reilly K, Shallow S, et al. Burden of Campylobacter infection in the United States and declining trend in California, FoodNet 1996-1998. International Conference on Abstract Emerging Infectious Diseases. Atlanta, GA, 2000.
- 24. French NP, Marshall JC, the Molecular Epidemiology and Public Health Laboratory. Source Attribution Studies for Campylobacteriosis in New Zealand: Massey University, 2014.
- 25. Müllner P, Spencer SEF, Wilson DJ, et al. Assigning the source of human campylobacteriosis in New Zealand: A comparative genetic and epidemiological approach. *Infect Genet Evol* 2009;9(6):1311-19. doi: 10.1016/j.meegid.2009.09.003
- Stafford Russell J, Schluter PJ, Kirk MD, et al. A multi-centre prospective case-control study of campylobacter infection in persons aged 5 years and older in Australia. *Epidemiol Infect* 2007;135(6):978-88. doi: 10.1017/S0950268806007576
- 27. Merritt TD, Combs B, Pingault N. Campylobacter outbreaks associated with poultry liver dishes. *Commun Dis Intell Q Rep* 2011;35(4):299-300.
- 28. Li D, Liu C-M, Luo R, et al. MEGAHIT: an ultra-fast single-node solution for large and complex metagenomics assembly via succinct de Bruijn graph. *Bioinformatics (Oxford, England)* 2015;31(10):1674-6. doi: 10.1093/bioinformatics/btv033 [published Online First: 2015/01/23]
- 29. Seemann T. Prokka: rapid prokaryotic genome annotation. *Bioinformatics (Oxford, England)* 2014;30(14):2068-9. doi: 10.1093/bioinformatics/btu153 [published Online First: 2014/03/20]
- 30. Glass K, Fearnley E, Hocking H, et al. Bayesian Source Attribution of Salmonellosis in South Australia. *Risk Anal* 2016;36(3):561-70. doi: 10.1111/risa.12444
- 31. Hald T, Vose D, Wegener HC, et al. A Bayesian Approach to Quantify the Contribution of Animal-Food Sources to Human Salmonellosis. *Risk Anal* 2004;24(1):255-69. doi: 10.1111/j.0272-4332.2004.00427.x
- 32. Sears A, Baker MG, Wilson N, et al. Marked Campylobacteriosis Decline after Interventions Aimed at Poultry, New Zealand. *Emerg Infect Dis* 2011;17(6):1007. doi: 10.3201/eid1706.101272
- 33. Müllner P, Jones G, Noble A, et al. Source attribution of food-borne zoonoses in New Zealand: a modified Hald model. *Risk Anal* 2009;29(7):970-84. doi: 10.1111/j.1539-6924.2009.01224.x
- 34. Lunn DJ, Thomas A, Best N, et al. WinBUGS A Bayesian modelling framework: Concepts, structure, and extensibility. *Stat Comput* 2000;10(4):325-37. doi: 10.1023/a:1008929526011
- 35. Dearlove BL, Cody AJ, Pascoe B, et al. Rapid host switching in generalist Campylobacter strains erodes the signal for tracing human infections. *ISME J* 2015;10:721. doi: 10.1038/ismej.2015.149
- 36. Mather AE, Vaughan TG, French NP. Molecular Approaches to Understanding Transmission and Source Attribution in Nontyphoidal Salmonella and Their Application in Africa. *Clin Infect Dis* 2015;61(suppl_4):S259-S65. doi: 10.1093/cid/civ727

- 37. Mughini Gras L, Smid JH, Wagenaar JA, et al. Risk Factors for Campylobacteriosis of Chicken, Ruminant, and Environmental Origin: A Combined Case-Control and Source Attribution Analysis. PLoS One 2012;7(8):e42599. doi: 10.1371/journal.pone.0042599
- 38. Kulldorff M, Heffernan R, Hartman J, et al. A Space-Time Permutation Scan Statistic for Disease
- 39. Touray K, Adetifa IM, Jallow A, et al. Spatial analysis of tuberculosis in an Urban West African



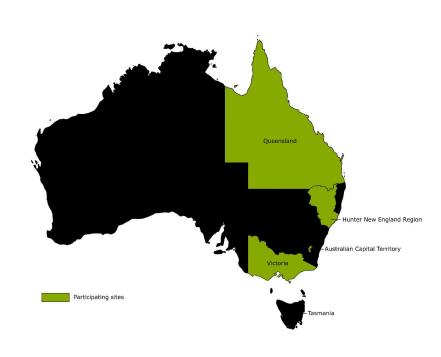


Figure 1. Map of Australian states and territories, showing the Hunter New England region.

INTRODUCTION

erviewer Note:			nt or quardian mag
	If case is less than 15 years of age you will familiar with the eating habits of the child.		it of guardian mos
	If case is aged between 15–17 years you we consent prior to interview.	ill need to obtain paren	nt or guardian
	Please note that for subjects under the age not the person being interviewed unless sp		
Health / Hunter N	is <interviewers name=""> and I am calling on New England Public Health Unit]." eak with <name case="" of=""> or <name case<="" of="" th=""><th></th><th>nd Health / ACT</th></name></name></interviewers>		nd Health / ACT
erviewer Note:	When the case comes to the phone then re explanatory statement.	peat the introduction ar	nd proceed with the
	If the case is unavailable then arrange an a	lternative time for the i	interview
more about you	experienced an illness due to the <i>Campylob</i> or <i>Campylobacter</i> illness. Your participa	acter bacteria. We wou tion is voluntary, al	ıld like to find out ll responses are
that you recently more about you confidential and i "Would you be p illness? The quest "In this study, we illness. Your part be presented in Australian Privace	experienced an illness due to the Campylob	acter bacteria. We would toon is voluntary, all to answer, just say so, our illness and activityes d activities you underty time. No individual this research will o	ald like to find out Il responses are ities prior to your No □ took prior to your I information will comply with the
that you recently more about you confidential and i "Would you be p illness? The quest "In this study, we illness. Your part be presented in Australian Privac ANU website.	experienced an illness due to the Campylobe or Campylobacter illness. Your participal if there are any questions you do not wish repared to answer some questions about you tions will take approximately 20 minutes." will collect information on foods you ate an icipation is voluntary and you can stop at a any reports or presentations. Partners in	acter bacteria. We wou tion is voluntary, al to answer, just say so. our illness and activit Yes d activities you undert ny time. No individual n this research will o	ald like to find out Il responses are ities prior to your No □ took prior to your I information will comply with the
that you recently more about you confidential and i "Would you be p illness? The quest "In this study, we illness. Your part be presented in Australian Privace ANU website."	experienced an illness due to the Campylobe or Campylobacter illness. Your participatif there are any questions you do not wish repared to answer some questions about y tions will take approximately 20 minutes." will collect information on foods you ate an icipation is voluntary and you can stop at a any reports or presentations. Partners in cy Act 1988. An information sheet about the	acter bacteria. We wou tion is voluntary, al to answer, just say so. our illness and activit Yes d activities you undert ny time. No individual n this research will o	ild like to find out Il responses are ties prior to your No took prior to your I information will comply with the available on the
that you recently more about you confidential and i "Would you be p illness? The quest "In this study, we illness. Your part be presented in Australian Privace ANU website. "Would you like "Would you like "This study has b 2016/426). If you Human Research Research Services The Australian N Tel: 6125 7945 Fa	experienced an illness due to the Campylobe or Campylobacter illness. Your participal if there are any questions you do not wish repared to answer some questions about you tions will take approximately 20 minutes." will collect information on foods you ate an icipation is voluntary and you can stop at a any reports or presentations. Partners in cy Act 1988. An information sheet about the me to send you a copy or provide you with u give me an address/email?" Deen approved by the Australian National I have any concerns please direct them to a Ethics Committee is Office, Chancellery 10B lational University, ACT 2601	dacter bacteria. We wou tion is voluntary, al to answer, just say so. our illness and activit Yes d activities you undert ny time. No individual n this research will on his research project is the link?" Yes	ild like to find out Il responses are ties prior to your No took prior to your I information will comply with the available on the
that you recently more about you confidential and i "Would you be p illness? The quest "In this study, we illness. Your part be presented in Australian Privace ANU website. "Would you like the Would you like the Yould you like the Yould you like the Yould you like the Yould you have the Young have the Yould you have the Young have the You	experienced an illness due to the Campylobe or Campylobacter illness. Your participal if there are any questions you do not wish repared to answer some questions about you tions will take approximately 20 minutes." will collect information on foods you ate an icipation is voluntary and you can stop at a any reports or presentations. Partners is cy Act 1988. An information sheet about the me to send you a copy or provide you with u give me an address/email?" been approved by the Australian National thave any concerns please direct them to a Ethics Committee is Office, Chancellery 10B lational University, ACT 2601 ax: 6125 4807	dacter bacteria. We wou tion is voluntary, al to answer, just say so. our illness and activit Yes d activities you undert ny time. No individual n this research will on his research project is the link?" Yes	ild like to find out Il responses are ties prior to your No took prior to your I information will comply with the available on the

If NO, arrange an alternative time to phone back to conduct the interview If YES, continue

[&]quot;Do you have the time right now to answer these questions?"

Interviewer Note:

	"Because I will be asking about specific dates around the time of your illness, it may be helpful
	for you to have a calendar or diary in front of you. Do you need a few minutes to get these?"
	Yes, I will get one no
	The first few questions we'll be asking you are about some symptoms that are associated with [your/their] illness.
l.	For the purposes of this study, we define diarrhoea as 3 or more loose stools or bowel movements in any 24-hour period. When you had your <i>Campylobacter</i> infection, did you have diarrhoea?
	Yes
la	During this diarrhoeal illness, what was the maximum number of stools or bowel movements
	you had in any 24 hour period? 0-2
2.	For how many days did your diarrhoea last?
	Don't know/Not sure
	CALCULATE PRIOR TO INTERVIEW Date stool specimen collected Day Month Year
3.	Could you please let me know what the date was when your diarrhoea began? Day Month Year (If person is unsure of date then prompt with date of stool specimen)
	Don't know/Not sure

Refer to your calendar to determine the interval from DATE 4 WEEKS BEFORE

DIARRHOEA BEGAN to DATE 1 DAY BEFORE DIARRHOEA BEGAN.

4.	In the 4 weeks before your illness began, that is from <date 4="" before="" diarrhoea="" weeks=""> through <date 1="" before="" began="" day="" diarrhoea="">, did anyone else</date></date>							
	in your <u>household</u> test positive for <i>Campylobacter</i> ? Yes							
	No							
	Don't know/Not sure							
5.	In the 4 weeks before your illness began, that is from <date 4="" before<="" th="" weeks=""></date>							
	DIARRHOEA> through <date 1="" before="" began="" day="" diarrhoea="">, did anyone else in your household have diarrhoea?</date>							
	Yes							
	No							
	Don't know/Not sure							
Inter	viewer Note: Refer to your calendar to determine the interval from DATE 2 WEEKS BEFORE							
IIICI	DIARRHOEA BEGAN to DATE 1 DAY BEFORE DIARRHOEA BEGAN.							
6.	In the 2 weeks before your illness began, that is from <date 2="" before<="" td="" weeks=""></date>							
U.	DIARRHOEA> through <date 1="" before="" began="" day="" diarrhoea="">, did you travel overseas or interstate?</date>							
	INTERVIEWER NOTE:							
	IF participant answers "yes",							
	1. Clarify if the travel was overseas or interstate							
	2. If travel was interstate:							
	In the 4 weeks before your illness began, that is from <date 4="" before="" diarrhoea="" weeks=""> through <date 1="" before="" began="" day="" diarrhoea="">, did anyone else in your household test positive for Campylobacter? Yes</date></date>							
	Options to select:							
	A. If the participant has travelled overseas or spent the whole two weeks interstate: (Select option Yes)							
	B. If the participant has travelled interstate only for a portion of the time: (Select option No)							
	Yes							
	No							
	Don't know/Not sure							
	Yes							

During this illness, did you have any of the following symptoms?
Fever
Vomiting
Stomach cramps
Blood in your stool 2 7
Nausea 1 2 7
Headache 1 2 7
Muscle/body aches 1
Did you take any antibiotics as a result of this illness?
Yes
No 2 Go to Q. 10
Don't know/Not sure
What antibiotic(s) were you taking? [Ask person to get tablet bottle, if possible]
Azithromycin
Ciprofloxacin
Norfloxacin
Erythromycin
Doxycycline (also known as Doxy or Vibramycin.
Other (please specify)
Don't know/Not sure
Were you admitted to hospital overnight because of this illness?
Yes
No 2 Go to Q. 12
Don't know/Not sure
If yes, for how many nights were you hospitalised?
Don't know/Not sure
ewer Note: Refer to your calendar to determine the interval from DATE 4 WEEKS BE

"For the next few questions, I would like to ask you about events which may have occurred in the 4 weeks before your illness began, so again that's from <DATE 4 WEEKS BEFORE DIARRHOEA BEGAN> to <DATE 1 DAY BEFORE DIARRHOEA BEGAN>."

12. In those 4 weeks, were you taking any antibiotics?

Yes	1	
No	2	Go to Q. 14
Don't know/Not sure	7	Go to O. 14

ntervi	ewer Note: If person can't remoleave the space blan		ntibiotic(s), check the DK/NS box and
3.	What antibiotic(s) were you takin	ng? [Ask person to get ta	ablet bottle, if possible]
		DK/NS	What date did you stop taking these?
a.	Antibiotic 1	7	(DD/MM) 7 DK/NS
b.	Antibiotic 2	7	(DD/MM) 7 DK/NS
c.	Antibiotic 3	7	(DD/MM)
d.	Antibiotic 4		(DD/MM) 7 DK/NS
4.	In those 4 weeks, were you taking	g any regular medicatio	on that decreases stomach acid?
	Yes	1	
	No	2	Go to Q. 16
	Don't know/Not sure	7	Go to Q. 16
a. b. c. d.	Zantac (Ranitidine)		No DK/NS 2 7 7 2 7 7 2 7 7
		_	
P	roton Pump Inhibitor		
	roton Pump Inhibitor		
a.	Losec (Omeprazole)		2 7
a. b.	Losec (Omeprazole) Nexium (Esomeprazole)	1	2 7
a. b. c.	Losec (Omeprazole) Nexium (Esomeprazole) Somac (Pantoprazole)	1	2 7 7
a. b.	Losec (Omeprazole) Nexium (Esomeprazole)	1 1	2 7
a. b. c. d. e.	Losec (Omeprazole)	octor that you have an	y other long lasting condition or r symptom? (e.g. Crohn's disease,
a. b. c. d.	Losec (Omeprazole)	octor that you have an a or vomiting is a majo	y other long lasting condition or r symptom? (e.g. Crohn's disease,

Don't know/Not sure.....

17.	In the 4	weeks before o	ncet of illness	did you take	or receive any	of the following?
1/.	m me 4	i weeks defore o	mset of miness.	uiu vou take	or receive any	or me ronowing:

INTERVIEWER NOTE:

Cyclosporine ("it's an immunosuppressant")

		Y es		NO	L	/K/ľ	12
a.	Prednisone						
	or other steroids <u>not</u> used on your skin		1		2		7
b.	Cyclosporine		1		2		7
c.	Chemotherapy		1		2		7
	Radiation therapy		1		2		7



3.	EXPOSURES
Intervi	Refer to your calendar to determine the interval from the DATE 7 DAYS BEFORE
	DIARRHOEA BEGAN to the DATE 1 DAY BEFORE DIARRHOEA BEGAN
A	A. WATER
4413	
	m now going to ask you some questions about water that you consumed in the <u>7 days before your</u> <u>arrhoea began,</u> that is from [diarr_7_days_prior] to [diarr_1_day_prior].
<u> </u>	<u> </u>
18.	What is your main source of drinking water at home? (select one only)
	INTERVIEWER NOTE:
	Only read out options if they're unsure
	only read out options it they re unsure
a.	A rainwater tank
b.	A river or stream
C.	A private well, bore hole, or spearpoint A carrier or tank truck
d. e.	Municipal water supply (tap water)
f.	Purchased bottle water
g.	Other water supply
h.	Don't know/Unsure
Intervi	ewer Note: If person answered "Yes" to "Municipal water supply" or "Purchased bottle water",
	skip to Q.21
19.	Do you usually treat your main source of drinking water before drinking?
17.	bo you usuany treat your mam source of armixing water before armixing.
	If Required PROMPT: Some examples are chlorination, filtration, boiling and UV treatment of the water
	Yes
	No
	Don't know/Not sure
20.	Which of the following treatments are in place? (select all that apply)
a.	Chlorination
а. b.	Filtration.
c.	Boiling
d.	UV treatment
e.	Other
f.	Specify () Don't know
1.	Don't know 1 2 7

21.	Did you drink water from any of the fo	llowing sourc	es in tl	he 7 days before onset of diarrhoea:	?
	(Select all that apply)				
		Ves	No	DK/NS	

a. b.	. A river or stream	
c. d.		
u. e.		
f.		
g.		
	Specify ()	
	,	

B. DINING LOCATIONS

		ne next few questions ask about places where you may have eaten food in the <u>7 days before you ess began</u> . So that is from [diarr_7_days_prior] to through [diarr_1_day_prior]"
22.		During this time, did you eat any food prepared outside your home, for example takeaway, restaurant, someone else's home?
		Yes
23.		Did you eat any food from the following places?
1	a. b. c. d.	Café or restaurant
23a		How many meals prepared outside of your home, were eaten during this 7 day period?
		1-2 meals
		3-4 meals
		≥ 5 meals
		Don't know/Unsure

	T A	IDX	DD	TAL	ICTS
(I)A	IK Y	PKI))	

"I would now like to ask you about the dairy products you may have eaten in the 7 days before your diarrhoea began."

24. Did you drink any raw/unpasteurised milk or eat any products made from raw/unpasteurised milk?

INTERVIEWER NOTE:

Cold-pressed milk is pasteurised and is not to be included as "raw/unpasteurised".

Specify (0.)	

"I will now ask you some questions about meat and poultry that you may have eaten in the 7 days

25.	During these 7 days, did you eat any of the fo	ollowing do	eli meats or o	cold cuts?	
		Yes	No	DK/NS	
b. c. d.	Salami/mettwurst Cabanossi/cabana/twiggy sticks Ham/chicken/turkey/beef Devon/frankfurts/cheerios Liverwurst Other Specify ()	1 1 1 1 1 1	2 2 2 2 2 2 2 2 2	7 7 7 7 7	
26.	During these 7 days, did you eat any pate?				
	Yes No Don't know/Not sure	1 2 7	Go to Q.29	•	
27.	Was the pate eaten,				
	Chicken pate Duck pate Pork pate Another type of pate Don't know/Not sure	2 3	Specify (_		
28.	Was this pate homemade or purchased from	a store?			
	Homemade Store Don't know/Not sure	2			
29.	During these 7 days, did you eat any other m	eat or pou	ltry? Like b	eef, lamb, chi	icken etc.
	INTERVIEWER NOTE:				
	This does not include eggs				
	Yes	_ 1			
	No Don't know/Not sure	2	Go to Q.49		

BE	EEF / VEAL	BMJ
30.	During these 7 days, did you eat any beef or veal?	Open:
	Yes	BMJ Open: first published as 10.1136/bmjopen-2018-026630 on 22 December 2018. Downloaded from http://bmjopen.bmj.com/ on April 10, 2024 by guest. Protected by copyright
31.	During the 7 days prior, did you eat any of the following beef or veal?	as 10.113
a	. Minced beef dishes	5/bmjopen-201
	(eg. bolognese sauce, pie, pastie, lasagne, hamburger patties, sausages)	8-026
b c.		630
d	(eg. tripe, liver, tongue)	n 22
	(eg. casserole, stir fry, steak, fillet, roast, beef strips)	Decer
PO	ORK .	nber :
		2018.
32.	During these 7 days, did you eat any pork?	Dow
	Yes) nloac
	Don't know/Not sure	ded from
33.	During the 7 days prior, did you eat any of the following pork?	m http://
		/bmjo
	No DK/N	pen.bn
a	. Minced pork dishes	nj.com/ or
b		ı Apri
C	. Offal	
d	. Other 1 2 7 (eg. casserole, stir fry, steak, fillet, roast, pork strips)	2024 b
LA	ΔMB	y guest. Pro
34.	During these 7 days, did you eat any lamb/mutton?	tected
	Yes Specify () d by (
	No	сору
	Don't know/Not sure	right

35.	During the 7 days prior, did you eat any of the following lamb/mutton?
	·
	$egin{array}{cccccccccccccccccccccccccccccccccccc$
a.	Minced lamb/mutton dishes 1 2 2 7 (eg. bolognese sauce, pie, pastie, lasagne, hamburger patties, sausages)
b.	Kebabs/souvlaki
c.	Offal
d.	Other
	(eg. casserole, stir fry, steak, fillet, roast, lamb strips)
GA	ME MEAT
36.	During these 7 days, did you eat any game meat like kangaroo, wallaby, venison or similar?
	Van
	No
	Don't know/Not sure
PO	ULTRY
37.	How often do you <u>usually</u> consume chicken/poultry meat?
	3 or more days per week
	1-2 days per week
	Once per fortnight
	Less often than once per fortnight
	Never 5 Go to Q.46
	Don't know/Not sure
38.	During the 7 days before your illness began, did you eat any chicken or other poultry?
	Yes
	No
	Don't know/Not sure
39.	How many meals did you eat that contained chicken or other poultry in the 7 days prior to onset of diarrhoea?
	1-2 meals
	3-4 meals
	\geq 5 meals
	Don't know/Not sure
	2511 1110 11/1 100 1010 11 11 11 11 11 11 11 11 11 11

40.	Did you consume any chicken or poultry at home?
	Yes
	No
	Don't know/Not sure
41.	Was the chicken or poultry purchased? (Select all that apply)
	Raw and fresh
	Raw and frozen 2
	Pre-cooked
	Don't know/Not sure
42.	How was it stored before consumption? (Select all that apply)
	INTERVIEWER NOTE:
	(On the bench)
	This is only to be used if they STORE their meat on the bench, this does not include defrosting their meat on
	the bench.
	In the freezer
	In the fridge 2
	On the bench
	Don't know/Not sure
43.	Prior to cooking, was the chicken rinsed or washed under running water?
	Yes
	No
	Don't know/Not sure
44.	During this did time you eat any of the following cooked meats?
77.	
	Yes No DK/NS
	No N
8	a. Chicken mince
ł	b. Chicken kebabs
C	c. Chicken pieces with bones 1 2 7 (i.e. wings, drumsticks, whole chicken)
(d. Chicken pieces without bones.
6	e. Offal
	specify: liver other
f	
٤	g. Turkey 2 7

45.	During this time, on how many days did you eat poultry?
	INTERVIEWER NOTE: 1. A pate is included 2. Eggs are excluded
	Days:
46.	During this time, on how many days did you eat meat (including poultry)?
	INTERVIEWER NOTE: Pate is included
	Days:
47.	During the 7 days prior to illness, did [you/they] eat any meat product, which was raw, rare or appeared undercooked?
	Yes
48.	Which of the following meats did [you/they] eat that was undercooked?
t c c f	Yes No DK/NS
	g. Offal (specify type) Other meat

49. How do you *prefer* the following meat to be cooked?

INTERVIEWER NOTE:

Raw: Not cooked at all Rare: Mostly red

Medium: Pink through out Well done: Brown through out

INTERVIEWER NOTE:

If participant answers Medium/Rare select the rarer option.. e.g Rare

a. b. c. d.	Chicken/Poultry Beef/Veal Pork Lamb	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	Rare 2 2 2	Medium 3 3 3	Well done 4 4 4
e.	Hamburgers		2 2	3 3	4 4
f.	Minced meat		2	3	4

E. GENERAL KITCHEN PRACTICES

yo	ou don't want to."
50.	How many times per week do you cook for members of your household?
	INTERVIEWER NOTE:
	This section around food prepared in the home refers to the person answering the survey (not necessarily the case or control)
	0
	1-5
	>5
	Don't know/Not sure
51.	Did you handle or prepare any raw meats in the kitchen in the 7 days before your
	diarrhoea began?
	INTERVIEWER NOTE:
	Refers to the person answering the survey
	Yes
	No
	Don't know/Not sure
52.	Did you handle or prepare raw chicken meat or chicken offal in the 7 days before your
	diarrhoea began?
	INTERVIEWER NOTE:
	Refers to the person answering the survey
	Yes
	No 2
	Don't know/Not sure
Interv	iewer Note: If person answered "No" to both Q.51 and Q.52 then skip to Q.57
53.	After a knife is used to cut raw meat or poultry, which of the following options do you usually do?
	INTERVIEWER NOTE:
	Refers to the person answering the survey
	Continue using the knife as is
	Rinse the knife before continuing to cook
	Wipe the knife before continuing to cook
	Wash the knife with detergent before continuing
	Change to another knife 5
	Other
	No one prepares meat
read	Don't know/Not sure

1 2	
3 4 54. 5	After a cutting board is used to cut raw meat or poultry, which of the following options do you usually do?
5 7	INTERVIEWER NOTE:
7 8	1. Does not matter if water is hot or cold
9	2. Refers to the person answering the survey
10	
11	Continue using the cutting board as is.
12	Rinse the cutting board before continuing to cook
13	Wipe the cutting board before continuing to cook
14	Wash the cutting board with detergent before continuing
15	Change to another cutting board.
16 17	Other
17 18	Specify ()
19 <i>Don't read</i> 20	Don't know/Not sure
21	
²² 55.	After handling raw meat or poultry in the kitchen, which of the following would you
23 24	usually do before continuing to cook?
25	INTERVIEWER NOTE:
26	Refers to the person answering the survey
27	
28	Wipe hands
29	Quickly rinse hands under a running tap 2
30	Wash hands with soap and water 3
31	Other
32	Specify (
33 34	Specify ()
	Don't do anything about hands
³⁵ Don't read 36	Don't know/not sure
37	
³⁸ 56.	After washing hands during food preparation, what would you usually dry
39	your hands on?
40	INTERVIEWER NOTE:
41	Refers to the person answering the survey
42 43	
44	Paper towel
45	Sponge/cloth.
46	Tea-towel /hand towel
47	A
48	Don't day had do
49	· · · · · · · · · · · · · · · · · · ·
50	Other
51 52 Dani's ward	Specify ()
52 Don't read	Don't know/Not sure
53 54	
55	
56 57.	In the past 3 months, has anyone in the household cook meat on a BBQ?
58	Yes
59	No 2 Go to Q. 59
60	Don't know/Not sure

58.	After cooking on the BBQ, where would the cooked meat most likely be placed?	
	Back on the same container	
	Back on the same container after it has been rinsed with water	
	Back on the same container after it has been wiped off with a towel	
	Back on the same container, after the container has been washed with soap and water	
n't read	On a different container 5 Other 6 Don't know/not sure 7 Specify (

1 F. ANIMAL AND PET EXPOSURE "The next few questions are about contact with animals in the 7 days before your diarrhoea began." 59. During this time, did you keep or care for any of the following animals as pets? **INTERVIEWER NOTE:** Not to include one off contact Is any pet less than 6 months old? Yes DK/NS No DK/NS a. Cat.... 2 2 b. Dog..... 2 2 Chickens c. 2 2 d. Other birds. 2 2 Specify (Other e. f. Do not keep any pets..... 7 Go to Q.67 If person answered No/Don't know to Cat then skip to question 62 **Interviewer Note: 60.** Do you feed your cat raw meat or bones? Yes.... Specify (Go to Q. 62 No..... (eg. chicken, beef, kangaroo, lamb etc.) Don't know/Not sure..... Go to Q. 62 61. How often does your cat get fed raw meat or bones? Daily..... Weekly Monthly..... Less often.... Don't know/Unsure.... If person answered No/Don't know to Dog then skip to question 65

IIItti v	it were two. If person answered two/Don't kno-	w to Dog	then skip to question of	
62.	Do you feed your dog raw meat?			
	Yes No	1	Specify ()
	Don't know/Not sure	7	(eg. chicken, beer, kangaroo, iamb etc.)	
63.	Do you feed your dog raw bones?			
	Yes	<u> </u>	Specify ()
	No	2		
	Don't know/Not sure	7		

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4. How often does your dog get fed raw meat or bones?	
_	
Daily	
Weekly	
Monthly 3 Less often	
Don't know/Not sure	
5. Did you get any of your pets in the 4 weeks before your diarrhoea began?	
Yes)
No	/
Don't know/Not sure	
2 on thin without suite	
6. Were any of your own pets ill with diarrhoea in the <u>7 days before your diarrhoea began</u> ?	
Yes)
No 2	
Don't know/Not sure	
bag)? Yes)
No 2	
Don't know/Not sure	
8. Do you live on a farm/hobby farm including a property on acreage 5 acres or over?	
Yes	
Double Lord and Allest and	
Don't know/Not sure	
9. In the 7 days before your diarrhoea began, did you visit a farm or petting zoo?	
Yes	
No	, petting zoo e
Don't know/Not sure	

3

4. DEMOGRAPHICS

4 [
5 6 "I	would now like to ask you a few final questions. Remember, your participation is voluntary and
,	ou do not have to answer any of the questions if you don't want to."
3	1.
70.	Is any language other than English spoken in your household?
0	
1	Yes
2	No 2
3 Don't read	Don't know/Not sure
4 Don't read	Refused
5	
6	
⁷ 71.	Are you of Aboriginal or Torres Strait Islander origin?
8	<u> </u>
9	No
0	Aboriginal 2
:1	Torres Strait Islander
2	Both
3 4 Don't read	Don't know/Not sure
5 Don't read	Refused
	Telused
16 17	
7 8 72.	Which of the following places best describe where you live?
.6 7 2. .9	which of the following places best describe where you live:
0	Inner city or urban area
1	Suburban area
2	Town
3	Rural or remote area community4
34	Rural or remote area farm or property
³⁵ Don't read	
Don i redu	
Don't read	Refused 9
8	
9	
	lewer Note: See definitions below.
.1	
2	
	y area:housing close to the centre of a major/capital city
4 <u>Suburba</u> 5	n area:
_	the region being primarily a self-contained residential district.
7 <u>10wii.</u>	community over 2000 people
Rural or	remote area community:community under 2000
Rural or	remote area farm or property
0	
1	
2	
73.	Does your occupation involve any of the following?
4	_
5	Working with raw meat
6	(eg. restaurants, butchery, abattoir etc.)
7	Working with animals
8	(eg. farmer, zookeeper, vet/nurse etc.)
9	Other type of occupation
50	Retired14

CASE not of working age.....

Don't know/Unsure.....

74.	What is the highest level of education reached by <u>anyone</u> in your household?	
	Schooling to year 10 or below	
on't read on't read	University degree—Undergraduate	
de fo	Now I am going to read you a list of income categories. Please stop me when a categor escribes your total household income, before taxes, in the last financial year? That is the total or all household members."	-
75.	Last year the total income for your household was?	
	Less than \$25,000	
on't read	Don't know/Not sure	
on't read	Refused	
76.	As part of this research we are planning to do a follow-up study. Would you be happy you in ~6 months' time?" Yes	for us to con
	you in ~6 months' time?" Yes	for us to con
	you in ~6 months' time?" Yes	for us to con
	you in ~6 months' time?" Yes	for us to con
	you in ~6 months' time?" Yes	for us to con
	you in ~6 months' time?" Yes	for us to con
	you in ~6 months' time?" Yes	for us to con
Intervi	you in ~6 months' time?" Yes	for us to con
Intervi	you in ~6 months' time?" Yes	for us to con
Intervi	you in ~6 months' time?" Yes	for us to con
Intervi	you in ~6 months' time?" Yes	for us to con
Intervi	you in ~6 months' time?" Yes	for us to con
Intervi	you in ~6 months' time?" Yes	for us to con
Intervi	you in ~6 months' time?" Yes	for us to con
Intervi	you in ~6 months' time?" Yes	for us to cor
Intervi	you in ~6 months' time?" Yes	for us to con