Can foodborne illness estimates from different countries be legitimately compared?: case study of rates in the UK compared with Australia, Canada and USA

Darren Holland, Robin Clifford, Nazmina Mahmoudzadeh, Sarah O'Brien, Guy Poppy, Zebulah Meredith, Harry Grantham-Hill

ABSTRACT
Objective Mathematical models have gained traction when estimating cases of foodborne illness. Model structures vary due to differences in data availability. This begs the question as to whether differences in foodborne illness rates internationally are real or due to differences in modelling approaches.

Difficulties in comparing illness rates have come into focus with COVID-19 infection rates being contrasted between countries. Furthermore, with post-EU Exit trade talks ongoing, being able to understand and compare foodborne illness rates internationally is a vital part of risk assessments related to trade in food commodities.

Design We compared foodborne illness estimates for the United Kingdom (UK) with those from Australia, Canada and the USA. We then undertook sensitivity analysis, by recreating the mathematical models used in each country, to understand the impact of some of the key differences in approach and to enable more like-for-like comparisons.

Results Published estimates of overall foodborne illness rates in the UK were lower than the other countries. However, when UK estimates were adjusted to a more like-for-like approach to the other countries, differences were smaller and often had overlapping credible intervals. When comparing rates by specific pathogens, there were fewer differences between countries. The few large differences found, such as virus rates in Canada, could at least partly be traced to methodological differences.

Conclusion Foodborne illness estimation models are country specific, making international comparisons problematic. Some of the disparities in estimated rates between countries can be shown to be attributed to differences in methodology rather than real differences in risk.

INTRODUCTION
Foodborne illness (FBI), also referred to as foodborne disease or food poisoning, is caused by consuming contaminated food or drink. The majority of over 250 known FBIs are due to pathogenic micro-organisms. Typical symptoms include nausea, vomiting, stomach cramps and diarrhoea. The WHO estimates that just 51 foodborne diseases cause as many as 600 million cases of illness worldwide each year. That is equivalent to almost 1 in 10 people on the planet becoming

WHAT IS ALREADY KNOW ABOUT THIS SUBJECT
⇒ In recent years, several countries have used mathematical models to produce estimates of foodborne illness. The models vary by country and are dependent on the source data available. These estimates were developed to aid within country policy development and prioritisation.
⇒ Since EU Exit the results from these models have been used to make comparisons of foodborne illness rates between countries as part of risk assessments related to trade in food commodities.
⇒ Published estimates of total infectious intestinal disease and total foodborne illness rates are lower in the UK than Australia, Canada and USA.

WHAT ARE THE NEW FINDINGS
⇒ When UK estimates of total infectious intestinal disease and foodborne illness rates were adjusted to a more similar approach to the other three countries, differences were smaller and often had overlapping credible intervals.
⇒ Differences in estimates in foodborne illness rates for individual pathogens, while generally smaller, can in some instances also be demonstrated to be affected by differences in methodology.

HOW MIGHT THIS IMPACT OF CLINICAL PRACTICES IN THE FORESEEABLE FUTURE
⇒ This study highlights the uncertainty in estimates of foodborne illness and how differences in methodology can impact results making comparisons between countries problematic. However, whichever methodology is used, foodborne illness remains a significant issue in the UK.
ill annually. These illnesses lead to around 420,000 deaths, with children under 5 years of age bearing the brunt (125,000 deaths in this age group). Diarrheal diseases alone, caused by bacteria, viruses, and parasites, are associated with 550 million illnesses and 230,000 deaths. Norovirus and Campylobacter spp are the most common viral and bacterial causes of FBI, respectively. Non typhoidal salmonellas, Salmonella Typhi, Taenia solium, hepatitis A virus and aflatoxin are the most common causes of death.

Coverage of COVID-19 has highlighted the difficulties of comparing infectious disease case rates between countries. There are similar problems when comparing intestinal disease (IID) and FBI rates. Following the result of the 2016 United Kingdom (UK) European Union (EU) membership referendum one of the many questions raised was whether changes in the UK’s trading partners might increase food safety risks. International trade in food is big business with 28% of food and drink consumed in the UK coming from the EU and a further 19% from third countries. To try to answer this question, several commentators have quoted publicly available FBI rates. In particular, media focus has been on differences between the UK and USA due to the potential of a substantial trade deal between the two nations. The US Centers for Disease Control and Prevention estimate that 48 million people get ill from FBI each year—a rate of one in six people getting FBI a year. For comparison, in the UK, the Food Standards Agency (FSA) estimates 2.4 million cases of FBI a year—a rate of 1 in 28 people getting FBI a year. On the face of it, this would support the assertion that the risk of acquiring FBI in the USA is over four and half times greater than in the UK. However, such top-level figures disguise differences in the ways in which baseline data are collected, the pathogens included, the manner in which overall disease burden is attributed to foodborne transmission, and other important nuances in calculations. Furthermore, all estimates provide credible intervals to show uncertainty. These intervals can be quite wide but are seldom reported in the press or by others. Without a full understanding of the context and methods involved in producing estimates, comparisons can be misleading at best, or at worst, be used to distort the facts deliberately.

Following discussion in the press and elsewhere on the relative risks of UK and US food in relation to potential trade deals, we re-analysed the estimates from the UK and the USA to determine what evidence there is for real differences in FBI rates between the two countries, considering both overall FBI levels and estimates for individual pathogens. We added studies from additional countries to the analysis based on a literature review, commissioned by the UK’s FSA, of the different methodologies used to estimate foodborne disease in the UK and in other countries. This review found 33 studies between 1994 and 2017. Of these studies, those for Australia and Canada used very similar approaches to the study for the USA, with all three providing estimates for at least 12 of the 13 individual pathogens covered in the UK study.

We, therefore, decided to add the Australian and Canadian studies to the analysis, to test whether arguments for the differences between the UK and USA would hold for them. For total IID, we also included research from Netherlands and Sweden, which were the only other two studies referenced in the literature review that, like the UK, used cohort studies. Unfortunately, these two studies did not include estimates for individual pathogens or for FBI so could not be included in the rest of the analyses.

**METHODS**

This re-analysis was based on data and results from studies for the UK, Australia, Canada and USA. The main features of the studies in the four countries are summarised in the online supplemental table 1.

The studies for each country produced the following estimates (see also online supplemental table 2):

1. Total IID: this was based on the specific case definition of IID (or acute gastroenteritis illnesses as it is often referred to outside the UK) used in each study. For Australia, Canada and USA, telephone surveys with 28-day recall were used, while for the UK, three different estimates were produced. These were based on a cohort study and two telephone surveys using 7-day and 28-day recall, respectively.

2. Total cases for specific pathogens: for Australia, Canada and USA, the studies used two different approaches depending on pathogen and data availability. The first used surveillance data scaled up for underreporting and underdiagnosis. They were then adjusted for cases acquired abroad. The second used an estimate of the proportion of IID due to each pathogen and applied this to the total IID in table 1.

For the UK, estimates were based on a cohort study undertaken in 2009: where data were sufficiently robust, the UK had subsequently adjusted estimates for some pathogens based on changes in confirmed laboratory reports to reflect changes between 2009 and 2018. As well as producing estimates for pathogens that cause IID, all four countries produced estimates for one or more non-IID pathogen (pathogens that do not cause diarrhoea or vomiting) using approach 1. For Australia, the IID and non-IID pathogens were reported separately and only total FBI figures for the IID pathogens are included in this paper. For the other three countries, the impact of excluding non-IID pathogens is explored later.

3. Cases of FBI caused by specific pathogens: for all four countries, these were based on the estimates from all sources (2 above) multiplied by the proportion estimated to be due to food. This is with the exception of norovirus in the UK where the estimate was based on the Norovirus Attribution study,13 14 which assessed the contribution made by the food chain to the burden of UK-acquired norovirus infection.

4. Total FBI: for all four countries, these estimates were based on subtracting the IID estimates for each specific
IID pathogen (2 above) from total IID (1 above) to get an estimate of unattributed cases (also known as unspecified agents or cases of unknown aetiology). These are cases where no specific pathogen was identified, either because there was insufficient data to attribute it to a specific pathogen, or the specific pathogen was not one of the ones included in the study. This figure was then multiplied by the foodborne proportion, estimated by taking the weighted foodborne proportion of all the specific IID pathogens. The foodborne estimate for the unattributed cases was then added to all the foodborne estimates for the specific pathogens (with the exception of Australia, these included one or more non-IID pathogen) to get total FBI. These unattributed cases make a large contribution of overall FBI, accounting for around 60% of total FBI in the UK and Canada, 80% in Australia and the USA.

Estimates from 2 to 4 all used Monte Carlo simulations to model the uncertainty in the estimates (note while the USA study used estimates under 2 to produce 3 and 4, the figures were not published. We, therefore, recreated these estimates based on the information in the technical annex 3 of the study).

Using the data available from these studies, we compared the estimates of overall FBI and IID as well as the estimates for individual pathogens. Estimates were converted into rates per 1 million person-years. To compare results between countries, more easily we re-ran the UK models to produce estimates of the mean and 90% credible intervals (UK published results gave the median with 95% credible intervals), so that the outputs were consistent. We recreated the Monte Carlo simulation models for each country based on information in the published papers and relevant technical appendices to provide comparable outputs. This allowed us to produce comparable results for all countries and to try some sensitivity analysis under different assumptions. Each model, including the one for the UK, was produced using @risk V.7.6 (http://www.palisade.com/). Each simulation was run for 100000 iterations. As the reported results were from simulations using random numbers, it was not possible to get complete agreement; however, differences between our results and those published were small.

For comparisons between countries, we compared 90% credible intervals.

RESULTS

Overall IID rates
Figure 1 (see online supplemental table 3) shows 90% credible intervals for the overall IID rates. The studies for Australia, Canada and USA used telephone surveys with 28-day recall to estimate total IID. The UK produced three estimates, a cohort study plus two telephone studies with 7-day and 28-day recall, respectively. We have also added two cohort studies from Netherland and Sweden for comparison (cohort studies tend to be uncommon, largely due to their complexity and expense).

Differences between countries, using similar telephone surveys with 28-day recall, are smaller than the differences between the three UK approaches, with the telephone surveys giving larger IID estimates than the cohort study. The cohort studies from Netherland and Sweden give estimates that are closer to the UK cohort study.

Overall FBI rates
Figure 2 (see online supplemental table 4) shows 90% credible intervals for the overall FBI rates for all four countries. The highest mean rate is for Australia with 193000 cases per 1 million person-years. This compares with the UK cohort rate of 36000 cases per 1 million person-years. We re-ran the UK model using the figures from the UK telephone surveys with 7 and 28-day recall for a fairer comparison. Figure 2 shows the mean rate from the 28-day recall is two times the rate from the UK cohort model, but it is only about a third of that from the 7-day recall. The mean estimate for the UK using 28-day recall
is still less than the other three countries. However, its 90% credible interval overlaps with those of Australia, Canada and the USA (note the cohort surveys in Netherlands and Sweden mentioned previously did not provide foodborne disease rates so could not be included).

**Sensitivity of FBI rates to the inclusion of certain pathogens**

Each study used a slightly different set of pathogens to produce estimates of total FBI. We, therefore, reproduced estimates for each country based on the 10 IID pathogens common to all four studies plus estimates for unattributed cases. The 10 common pathogens were *C. perfringens*, *Campylobacter*, *Salmonella* non typhoidal, *Shigella*, Cryptosporidium, Giardia, astrovirus, norovirus, rotavirus and sapovirus. These estimates were then compared with those published in the individual papers (see online supplemental figure S1).

Despite the large variation in the number of pathogens included in each study, the differences between the new estimates and those in the papers were relatively small and certainly much smaller than differences in rates seen between countries. There are a few reasons for this. First, these 10 pathogens contribute a large percentage of the total estimates from the known pathogens in each country, that is, 98.4% for the UK, 64.7% for Australia (96% of the rest is from other pathogenic *E. coli*), 91.0% for Canada and 91.7% for USA (see online supplemental table 5). Second, when the additional IID pathogens are removed from the calculations, this does not reduce overall IID, it just moves it from known pathogens to unspecified agents. Third, the proportion of IID from unspecified agents that is attributed to food is based on the weighted proportion for each of the known pathogens included in the calculation. As previously stated, the majority of cases are from the 10 common IID pathogens, so the weighting used only changes by a small amount when the other IID pathogens are excluded (see online supplemental table 6 for percentages). Finally, the number of cases attributed to non-IID pathogens is very small, making up 0.02% of the total for the UK, 0.6% for Canada and 1.1% for USA (for Australia only IID pathogens were included in the estimates in this paper). All four of these effects, even in combination, only make a small difference to overall estimates.

To further analyse the impact of using a cohort study compared with a telephone survey with either 7-day or 28-day recall, we produced estimates for IID for known pathogens only, as these estimates do not use any of these studies. As the different studies included, different known pathogens in their estimates of FBI, we estimated the total IID rate for the 10 common IID pathogens used in all four studies, as listed above, plus *Listeria monocytogenes* which was also included in all four studies. The total IID rates for USA and Australia for these 11 pathogens (see online supplemental table 7 and figure S2) are much closer to those in the UK, with large overlaps in the credible intervals. However, the IID rate for the 11 pathogens is two times as high in Canada than in the UK.

A key reason for this difference to Canada is the approach used to estimate virus rates. As no Canadian virus data were available, the Canadian study calculated these proportions using the UK IID2 study data. To do this, they used the UK estimate for each virus divided by the overall estimate of IID from the UK cohort study to get a proportion of the total IID caused by the virus. The Canadian estimate of total IID obtained by their telephone survey was then multiplied by this proportion. As the Canadian estimate of total IID (based on a telephone survey with 28-day recall) was 2.3 times greater than the UK cohort study, this means their estimate for each virus was also greater. If the study had used the UK’s telephone survey with 28-day recall instead as the estimate of total IID in the UK, then the Canadian virus estimates would have decreased by 51%. The Canadian estimate for *C. perfringens* was estimated in the same way as the viruses and so has the same issue. This is explored further in the comparisons for these individual pathogens later on.
We similarly considered just the 11 known pathogens common to all studies and estimated their contribution to foodborne disease. FBI rates on this basis are still lower in the UK than in the other three countries (see online supplemental table 8 and figure S3). Again, the large difference between the UK and Canadian figures is largely due to the way viruses and \textit{C. perfringens} were estimated in the Canadian study as discussed above.

Figure 3  Infectious Intestinal Disease rates for individual pathogens per 1 million population per annum (90% credible intervals shown with the mean).
IID rates for individual pathogens

Figure 3 (see online supplemental table 9) shows the 90% credible intervals for the estimated IID rates for each pathogen and country. These are for the 13 pathogens included in the UK study. For most pathogens, the 90% credible intervals largely overlap between countries, although there are some exceptions. One pathogen where there are differences is *E. coli* O157. In the UK models, it is assumed that there is no under-reporting due to the seriousness of the illness, which explains the much lower estimates compared with other countries, where adjustments for under-reporting have been included in the models. A second example is rotavirus, where estimates in UK and Australia are lower than the other two countries. For the UK at least, a large reason for this difference will be the years on which the estimates are based. In the UK, estimates are based on 2018 data (Australia on 2010) while Canadian and US estimates are based on 2006 data (see online supplemental table 1). This difference in timescales is particularly important due to roll out rotavirus vaccination programmes in infants over this period, which has decreased overall rates of the illness.18

As previously mentioned, no virus or *C. perfringens* data were available for Canada, so estimates were based on UK data using the proportion of IID associated with each pathogen. The UK IID figures used were those from the cohort study. If they were based on the telephone survey with 28-day recall, which is 2.3 times larger (see figure 1), then the proportion of IID attributed to each pathogen would reduce as would the Canadian estimates. As the estimates of IID for Canada also used a telephone survey with 28-day recall, this seems a reasonable adjustment. Figure 4 demonstrates the impact of this change. It shows estimates for Canada for *C. perfringens*, adenovirus, astrovirus, norovirus, rotavirus and sapovirus. New estimates have been calculated using the proportion of the IID attributed to each pathogen based on the UK telephone survey with 28-day recall. These estimates are then compared with those from the UK and the previous estimates for Canada based on the cohort study.

As is seen in figure 4 (see online supplemental table 10), the estimates for Canada are now much more similar to those of the UK. This is with the exception of rotavirus where UK rates are still lower due to the infant vaccination programme mentioned earlier.

FBI rates for individual pathogens

Figure 5 (see online supplemental table 11) shows the 90% credible intervals for the estimated FBI rates for each pathogen and country.

As with estimates of IID, for most pathogens, the 90% credible intervals largely overlap between countries, with *E. coli* O157 again an outlier. Estimates for viruses are
more similar than for IID. The reason for this narrowing of the gap is that although estimates of the proportion of IID attributed to food are low for the UK for viruses, they are still higher than for some of the other countries. Estimates of the proportion of cases attributed to food by pathogen by country are found in online supplemental table 12 and figure S4. Notable differences in these estimates between countries include those for Campylobacter and Giardia. For Campylobacter, the mean estimate for

**Figure 5** Foodborne illness rates for individual pathogens per 1 million population per annum (90% credible intervals shown with the mean).
rates based on the UK telephone survey with 28-
gastroenteritis.19  
are closer to the estimates in the comparator countries.  
differences on estimates of FBI.  
looking at a common symptom-
of the latter has been explored more fully in a study  
mation of individual pathogen rates may explain some  
three countries. Methodological differences in the esti-

produced on a like-
data were missing. Arguably then, it is inappropriate to  
ated to a similar basis as the  

DISCUSSION  
Overall FBI  
The three different IID estimates from the UK IID2 study  
methodology. Using the estimates of IID from either of  
the telephone surveys rather than the cohort study will  
lead to large increases in the FBI estimate, everything  
else being equal. An earlier study has previously shown  
the impact of such differences on estimates of IID,18  
including the effect of length of the recall period of  
phone surveys, but here we also show the impact of such  
differences on estimates of FBI.  
As shown in figures 1 and 2, the total UK IID and FBI  
rates based on the UK telephone survey with 28-day recall  
are closer to the estimates in the comparator countries.  
However, mean estimates are still less than the other  
three countries. Methodological differences in the esti-
mation of individual pathogen rates may explain some  
of the variation, as can differences in the case definition  
of IID (see online supplemental table 1). The impact  
of the latter has been explored more fully in a study  
looking at a common symptom-based case definition of  
gastroenteritis.19  
Future studies are likely to include online surveys which  
may introduce further methodological differences.  

Individual pathogens  
Researchers in each country have estimated cases based  
on the best data available to them. This varies consider-
ably between countries and pathogens. For instance,  
the UK estimates of IID are based on a cohort study, while  
the other three main studies reviewed used one of two  
different models for each pathogen depending on the  
data availability. Even then, further tweaks were needed.  
These included adjusting for partial geographical surveil-
ance for some pathogens in Australia and the USA, and  
using proxy parameters from other pathogens where  
data were missing. Arguably then, it is inappropriate  
to compare estimates for individual pathogens that are not  
produced on a like-for-like basis.  
A further consideration is the way the foodborne  
proportion is estimated for in each study, as the accuracy  
of this clearly makes a difference to the results. Both the  
Australia and Canadian studies used expert elicitation to  
estimate these proportions,20 21 while the UK and USA use  
outbreak data to estimate what proportion of cases from  
outbreaks was foodborne (with the exception of noro-
virus in the UK where foodborne estimates are based on  
a quantitative risk assessment model). This is an area of  
major uncertainty due to the difficulty of attributing cases  
to specific sources, meaning estimates are often based on  
limited data. Consequently, a few percentage point differ-
ence in proportion can lead to a large difference in the  
final estimate, particularly when the proportion is small.  
This is illustrated with sapovirus, where the estimated  
proportion attributed to food in the UK is 2.6%, which  
is fivefold higher than the Canadian and USA figures of  
0.5%. Where the proportion of cases attributed to food  
is higher, a similar percentage point difference will have  
more impact, for example, a change from 90% to 92.1%  
would be only a 2.3% increase.  
For Salmonella non typhoidal, the USA and Canada  
had significantly higher rates than the UK. We could find  
no obvious methodological reason for this difference.  
A real difference may be due to the mass vaccination of  
 poultry in the UK from the late 1990s.22  
Following this, human cases of Salmonella non typhoidal decreased by  
40% between 2000 and 2009 in the UK (Food Standards  
Agency, 2018).23  
A general point is that the studies on which these  
estimates are based are resource intensive and so are  
infroneglected and undertaken. This means that the compar-
isons made are based on research undertaken several  
years ago and at different periods. The study for the UK  
used data from 2008 to 2009 and 2018, Australia 2006  
The number of cases for each pathogen will vary by year  
anyway, but there may also been more fundamental long-
term changes over time, which could explain differences.  

CONCLUSIONS  
Estimates of FBI are important for policymakers in  
making decisions on resource allocation and prioritising  
interventions. Recently, such estimates have been used  
by various commentators to make statements on rela-
tive food safety risks in different countries. This paper  
shows that such comparisons need to be undertaken with  
considerable caution and appropriate caveats. Compar-
isons for individual pathogens between the four studies  
are possible but need very careful interpretation. While  
we have tried to control for methodological differences in  
approach where possible, there are other sources of vari-
atations between countries which we were unable to adjust  
for which may also have an impact on estimates. These  
include surveillance data, geographical coverage, access  
to healthcare and laboratory testing practices.  
For overall estimates of FBI, comparing published rates  
between countries can be misleading due to major differ-
ces in methodology and data sources. Our sensitivity  
analysis adjusted UK estimates to a similar basis as the  
other three countries. Then the differences in estimates,
which initially appear large, are much smaller, often with overlapping credible intervals.

By far the biggest impact on estimates for overall IID and FBI is the type of study used, for example, cohort study or telephone survey. In the case of the latter, the length of the recall period is also a major contributor to differences. We, therefore, recommend that only estimates using the same approach are compared, and ideally undertaken at a similar period of time. Differences in the number of pathogens included tend to have less impact, as most studies all include the major pathogens which contribute most of the total burden.

For individual pathogens, one big consideration is the year the estimates are for, which can make a large difference, particularly where interventions directed at specific pathogens have been undertaken between the different time periods. Another big factor we found is when estimates are based on the proportion of IID caused by a specific pathogen based on figures from a different country, particularly when that country uses a different method to estimate total IID.

Finally, differences between FBI rates are only part of the story in terms of whether changes in trading partners may change risks for the UK. These risks are more likely to be dependent on which specific foods are imported from each country and the relative risks of these compared with the equivalent domestically produced food, rather than overall FBI rates. Genuine differences in FBI rates between countries are also likely to be influenced by factors such as consumer preferences in terms of which foods are consumed and how they are prepared, diet, hygiene practices both in food business and at home and wider environmental factors such as climate.

REFERENCES


Table 1. Main features of studies of foodborne illness in the UK, Australia, Canada and USA

<table>
<thead>
<tr>
<th>Feature</th>
<th>UK</th>
<th>Australia</th>
<th>Canada</th>
<th>USA</th>
</tr>
</thead>
<tbody>
<tr>
<td>Year to which outputs relate</td>
<td>2009 &amp; 2018</td>
<td>2010</td>
<td>2006</td>
<td>2006</td>
</tr>
<tr>
<td><strong>Data sources on which estimates of overall burden of illness are based.</strong></td>
<td></td>
<td>National Gastroenteritis Survey II – a nationally representative, retrospective cross-sectional telephone survey conducted in 2008–2009 with 4-week recall.</td>
<td>National Studies on Acute Gastrointestinal Illness (NSAGI) population surveys conducted in Ontario and British Columbia via retrospective, cross-sectional telephone surveys with 28-day recall.</td>
<td>Cross-sectional, retrospective telephone survey of self-reported acute diarrheal illness with 28-day recall in Foodborne Diseases Active Surveillance Network (FoodNet) at 10 US sites extrapolated to the whole of the US.</td>
</tr>
<tr>
<td><strong>Case definitions.</strong></td>
<td></td>
<td>Gastroenteritis defined as ≥3 loose stools in 24 h or 2 or more episodes of vomiting within a 24-h period during the preceding 4 weeks in the absence of a non-infectious cause for illness.</td>
<td>Acute diarrheal defined as ≥3 loose stools in 24 h with duration lasting &gt;1 day not resulting from pregnancy, medication, food allergy, and/or medical conditions previously diagnosed by a doctor (e.g. colitis, diverticulitis, Crohn's disease, irritable bowel syndrome).</td>
<td>Acute diarrheal illness defined as ≥3 loose stools in 24 hours lasting more than 1 day or resulting in restricted daily activities.</td>
</tr>
<tr>
<td><strong>Adjustment for under-diagnosis?</strong></td>
<td>Estimates for 2009 did not need adjustments as they were measured as part of the IID2 study. 10 2018 updates used adjustments based on this study and the change in confirmed laboratory for Campylobacter, E. coli O157, Salmonella, Shigella, Cryptosporidium, Giardia and rotavirus. For 2018, C. Perfringens, adenovirus, astrovirus, norovirus,</td>
<td>Yes</td>
<td>Yes</td>
<td>Yes</td>
</tr>
</tbody>
</table>

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and sapovirus were scaled up based on population from the IID2 study, so were not adjusted for under-diagnosis. *E. coli* O157 and *Listeria monocytogenes* were assumed to have no under-diagnosis.

**Estimating proportion of illnesses that are foodborne**

<table>
<thead>
<tr>
<th>Methodology to produce estimates by country</th>
<th>UK</th>
<th>Australia</th>
<th>Canada</th>
<th>USA</th>
</tr>
</thead>
<tbody>
<tr>
<td>Total IID (1)</td>
<td>Cohort Study</td>
<td>Telephone survey with 28 day recall</td>
<td>Telephone survey with 28 day recall</td>
<td>Telephone survey with 28 day recall</td>
</tr>
<tr>
<td>Total cases for individual pathogens (2)</td>
<td>Estimates based largely on cohort study. Of the 13 pathogens covered in detail in this paper, <em>Campylobacter</em>, <em>Salmonella non-typhoidal</em>, <em>Shigella</em>, <em>Cryptosporidium</em>, <em>Giardia</em>, norovirus and rotavirus were scaled up based on confirmed laboratory reports and underreporting rates from the cohort study. For <em>C. perfringens</em>, adenovirus, astrovirus, norovirus and sapovirus cases</td>
<td>Of the 12 pathogens covered in detail in this paper <em>C. perfringens</em>, <em>Campylobacter</em>, <em>Listeria monocytogenes</em>, <em>Salmonella non-typhoidal</em>, <em>Shigella</em>, <em>Cryptosporidium</em> and <em>Giardia</em> scaled up from surveillance data. Adenovirus, astrovirus, norovirus, rotavirus and sapovirus scaled down from Total IID (1)</td>
<td>Of the 13 pathogens covered in detail in this paper <em>Campylobacter</em>, <em>E. coli O157</em>, <em>Listeria monocytogenes</em>, <em>Salmonella non-typhoidal</em>, <em>Shigella</em>, <em>Cryptosporidium</em> and <em>Giardia</em> scaled up from surveillance data. <em>C. perfringens</em>, adenovirus, astrovirus, norovirus, rotavirus and sapovirus scaled down from Total IID (1)</td>
<td>Of the 12 pathogens covered in detail in this paper <em>C. perfringens</em>, <em>Campylobacter</em>, <em>E. coli O157</em>, <em>Listeria monocytogenes</em>, <em>Salmonella non-typhoidal</em>, <em>Shigella</em>, <em>Cryptosporidium</em> and <em>Giardia</em> scaled up from surveillance data. Astrovirus, norovirus, rotavirus and sapovirus scaled down from Total IID (1)</td>
</tr>
</tbody>
</table>

**Table 2: Methodology to produce estimates by country**
rates per 1,000 person-years as given in the cohort study were used. For *E. coli* O157 and *Listeria monocytogenes* confirmed laboratory numbers were used as there was assumed to be no underreporting.

- **FBI for individual pathogens (3)**: Estimates of total cases for each pathogen (2) multiplied by proportion attributed to food. Exception is norovirus which is based on a Quantitative Microbial Risk Assessment model.

- **Total FBI (4)**: Total of FBI for all individual pathogens (3) plus foodborne estimate for unattributed cases. IID estimate for unattributed cases is Total IID (1) minus sum of IID for individual IID pathogens (2). Foodborne proportion for unattributed cases is based on weighted foodborne proportions for individual IID pathogens.

### Table 3: Infectious Intestinal Disease per 1 million population per annum

<table>
<thead>
<tr>
<th>Country and approach</th>
<th>Annual rate per 1 million population Mean</th>
<th>Lower 90% Cred.</th>
<th>Upper 90% Cred.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Australia - telephone survey 28 day recall</td>
<td>746.479</td>
<td>643,192</td>
<td>845,070</td>
</tr>
<tr>
<td>Canada - telephone survey 28 day recall</td>
<td>630.500</td>
<td>594,973</td>
<td>666,432</td>
</tr>
<tr>
<td>USA - telephone survey 28 day recall</td>
<td>612,000</td>
<td>581,786</td>
<td>642,214</td>
</tr>
<tr>
<td>UK - cohort study</td>
<td>274,512</td>
<td>257,230</td>
<td>292,506</td>
</tr>
<tr>
<td>UK - telephone survey 7 day recall</td>
<td>1,548,929</td>
<td>1,178,609</td>
<td>1,985,067</td>
</tr>
<tr>
<td>UK - telephone survey 28 day recall</td>
<td>542,370</td>
<td>393,532</td>
<td>722,424</td>
</tr>
<tr>
<td>Sweden - cohort study</td>
<td>360,000</td>
<td>326,000</td>
<td>395,000</td>
</tr>
<tr>
<td>Netherlands - cohort study</td>
<td>283,000</td>
<td>252,000</td>
<td>315,000</td>
</tr>
</tbody>
</table>
Table 4: Foodborne Illness per 1 million population per annum

<table>
<thead>
<tr>
<th>Country and approach</th>
<th>Annual rate per 1 million population</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
</tr>
<tr>
<td>Australia - telephone survey 28 day recall</td>
<td>193,005</td>
</tr>
<tr>
<td>Canada - telephone survey 28 day recall</td>
<td>123,077</td>
</tr>
<tr>
<td>USA - telephone survey 28 day recall</td>
<td>159,802</td>
</tr>
<tr>
<td>UK - cohort study</td>
<td>35,979</td>
</tr>
<tr>
<td>UK - telephone survey 7 day recall</td>
<td>202,985</td>
</tr>
<tr>
<td>UK - telephone survey 28 day recall</td>
<td>71,079</td>
</tr>
</tbody>
</table>

Figure S1: FBI rates per 1 million population per annum – sensitivity analysis (90% credible intervals shown with the mean)

The baseline is based on the models as described in the published papers. As the models have been re-run there are slight differences to previously published figures. Sensitivity analysis is based on using just the 10 IID pathogens common to all four studies.
Table 5: Proportion of estimated FBI from known pathogens by IID pathogens common to all 4 studies, other IID pathogens (those included in some studies) and non-IID pathogens

<table>
<thead>
<tr>
<th>% FBI by pathogen category</th>
<th>UK</th>
<th>Australia</th>
<th>Canada</th>
<th>USA</th>
</tr>
</thead>
<tbody>
<tr>
<td>% FBI from 10 common IID pathogens</td>
<td>98.4%</td>
<td>64.7%</td>
<td>91.0%</td>
<td>91.7%</td>
</tr>
<tr>
<td>% FBI from other IID pathogens</td>
<td>1.6%</td>
<td>35.3%</td>
<td>8.4%</td>
<td>7.2%</td>
</tr>
<tr>
<td>% FBI from non-IID pathogens</td>
<td>0.02%</td>
<td>NA</td>
<td>0.6%</td>
<td>1.1%</td>
</tr>
</tbody>
</table>
Table 6: Estimated proportion of cases attributable to food for unspecified agents

<table>
<thead>
<tr>
<th>Proportions based on:</th>
<th>UK</th>
<th>Australia</th>
<th>Canada</th>
<th>USA</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>Lower 90% Cred. Int.</td>
<td>Upper 90% Cred. Int.</td>
<td>Mean</td>
</tr>
<tr>
<td>IID pathogens used in individual country's study</td>
<td>13.1%</td>
<td>10.4%</td>
<td>16.3%</td>
<td>25.5%</td>
</tr>
<tr>
<td>10 IID Pathogens common to all 4 studies</td>
<td>14.4%</td>
<td>11.4%</td>
<td>18.0%</td>
<td>26.3%</td>
</tr>
</tbody>
</table>
Table 7: Infectious Intestinal Disease per 1 million population per annum for 11 pathogens where estimates are available for all four countries

<table>
<thead>
<tr>
<th>Country</th>
<th>Annual rate per 1 million population</th>
<th>Mean</th>
<th>Lower 90% Cred. Int.</th>
<th>Upper 90% Cred. Int.</th>
</tr>
</thead>
<tbody>
<tr>
<td>UK</td>
<td>94,855</td>
<td>84,808</td>
<td>105,689</td>
<td></td>
</tr>
<tr>
<td>Australia</td>
<td>100,860</td>
<td>82,505</td>
<td>121,332</td>
<td></td>
</tr>
<tr>
<td>Canada</td>
<td>220,306</td>
<td>201,342</td>
<td>240,021</td>
<td></td>
</tr>
<tr>
<td>USA</td>
<td>118,152</td>
<td>91,577</td>
<td>149,176</td>
<td></td>
</tr>
</tbody>
</table>

Figure S2: IID rates per 1 million population per annum for the 11 pathogens common to all 4 studies (90% credible intervals shown with the mean)

Note: Figures are based on the sum of the estimates for the 11 pathogens and do not use total IID estimates from either the telephone surveys or cohort studies.
Table 8: Foodborne Illness per 1 million population per year for 11 pathogens where estimates are available for all four countries

<table>
<thead>
<tr>
<th>Country</th>
<th>Annual rate per 1 million population</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
</tr>
<tr>
<td>UK</td>
<td>13,676</td>
</tr>
<tr>
<td>Australia</td>
<td>26,450</td>
</tr>
<tr>
<td>Canada</td>
<td>45,798</td>
</tr>
<tr>
<td>USA</td>
<td>28,752</td>
</tr>
</tbody>
</table>

Figure S3: FBI rates per 1 million population per annum for the 11 pathogens common to all 4 studies (90% credible intervals shown with the mean)

Note: Figures are based on the sum of the estimates for the 11 pathogens and do not use total IID estimates from either the telephone surveys or cohort studies.
Table 9: Infectious Intestinal Disease per 1 million population per year

<table>
<thead>
<tr>
<th>Pathogen</th>
<th>UK Lower</th>
<th>UK Upper</th>
<th>Australia Lower</th>
<th>Australia Upper</th>
<th>Canada Lower</th>
<th>Canada Upper</th>
<th>USA Lower</th>
<th>USA Upper</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>90% Cred.</td>
<td>Mean</td>
<td>90% Cred.</td>
<td>Mean</td>
<td>90% Cred.</td>
<td>Mean</td>
<td>90% Cred.</td>
</tr>
<tr>
<td>Bacteria</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>C. perfringens</td>
<td>1,695</td>
<td>671</td>
<td>3,366</td>
<td>2,465</td>
<td>5,446</td>
<td>3,221</td>
<td>642</td>
<td>8,306</td>
</tr>
<tr>
<td>Campylobacter</td>
<td>9,748</td>
<td>6,535</td>
<td>13,809</td>
<td>7,131</td>
<td>7,131</td>
<td>5,502</td>
<td>6,735</td>
<td></td>
</tr>
<tr>
<td>E. coli O157</td>
<td>13</td>
<td>13</td>
<td>13</td>
<td>NA</td>
<td>520</td>
<td>309</td>
<td>78</td>
<td></td>
</tr>
<tr>
<td>Listeria monocytogenes</td>
<td>3</td>
<td>3</td>
<td>3</td>
<td>7</td>
<td>7</td>
<td>5</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>Salmonella non typhoidal</td>
<td>716</td>
<td>145</td>
<td>1,914</td>
<td>2,650</td>
<td>3,366</td>
<td>3,366</td>
<td>5,903</td>
<td></td>
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<tr>
<td>Shigella</td>
<td>108</td>
<td>108</td>
<td>108</td>
<td>140</td>
<td>205</td>
<td>1,399</td>
<td>4,016</td>
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</tr>
<tr>
<td>Parasites</td>
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<td></td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cryptosporidium</td>
<td>745</td>
<td>151</td>
<td>1,990</td>
<td>850</td>
<td>779</td>
<td>2,248</td>
<td>6,482</td>
<td></td>
</tr>
<tr>
<td>Giardia</td>
<td>1,310</td>
<td>311</td>
<td>3,295</td>
<td>1,560</td>
<td>3,339</td>
<td>2,248</td>
<td>6,482</td>
<td></td>
</tr>
<tr>
<td>Viruses</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Adenovirus</td>
<td>10,379</td>
<td>7,187</td>
<td>14,342</td>
<td>4,150</td>
<td>23,009</td>
<td>NA</td>
<td>NA</td>
<td></td>
</tr>
<tr>
<td>Astrovirus</td>
<td>5,528</td>
<td>3,290</td>
<td>8,540</td>
<td>3,150</td>
<td>11,768</td>
<td>10,340</td>
<td>7,859</td>
<td></td>
</tr>
<tr>
<td>Norovirus</td>
<td>47,078</td>
<td>40,436</td>
<td>54,369</td>
<td>72,500</td>
<td>104,000</td>
<td>69,414</td>
<td>99,402</td>
<td></td>
</tr>
<tr>
<td>Rotavirus</td>
<td>1,712</td>
<td>1,221</td>
<td>2,311</td>
<td>2,100</td>
<td>26,161</td>
<td>10,340</td>
<td>7,858</td>
<td></td>
</tr>
<tr>
<td>Sapovirus</td>
<td>26,212</td>
<td>20,987</td>
<td>32,188</td>
<td>3,800</td>
<td>61,217</td>
<td>10,340</td>
<td>7,858</td>
<td></td>
</tr>
</tbody>
</table>
Table 10: Estimates of Infectious Intestinal Disease for certain pathogens for Canada and UK per 1 million population per year. Based on using different estimates of overall IID for Canada.

<table>
<thead>
<tr>
<th>Pathogen</th>
<th>UK Lower Mean</th>
<th>90% Cred. Int.</th>
<th>UK Upper Mean</th>
<th>90% Cred. Int.</th>
<th>Canada based on UK Cohort study as in paper Lower Mean</th>
<th>90% Cred. Int.</th>
<th>Upper Mean</th>
<th>90% Cred. Int.</th>
<th>Canada based on UK Cohort Study re-modelled Lower Mean</th>
<th>90% Cred. Int.</th>
<th>Upper Mean</th>
<th>90% Cred. Int.</th>
<th>Canada based on UK Telephone Survey with 28 day recall Lower Mean</th>
<th>90% Cred. Int.</th>
<th>Upper Mean</th>
<th>90% Cred. Int.</th>
</tr>
</thead>
<tbody>
<tr>
<td>C. perfringens</td>
<td>1,695</td>
<td>671</td>
<td>3,366</td>
<td></td>
<td>5,446</td>
<td>2,647</td>
<td>8,784</td>
<td>4,439</td>
<td>1,861</td>
<td>7,661</td>
<td>2,353</td>
<td>933</td>
<td>4,142</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Adenovirus</td>
<td>10,379</td>
<td>7,187</td>
<td>14,342</td>
<td></td>
<td>23,010</td>
<td>17,047</td>
<td>29,533</td>
<td>24,476</td>
<td>17,618</td>
<td>32,100</td>
<td>12,990</td>
<td>8,362</td>
<td>18,230</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Astrovirus</td>
<td>5,528</td>
<td>3,290</td>
<td>8,540</td>
<td></td>
<td>11,768</td>
<td>7,573</td>
<td>16,558</td>
<td>13,237</td>
<td>8,317</td>
<td>18,874</td>
<td>7,018</td>
<td>4,041</td>
<td>10,487</td>
<td></td>
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</tr>
<tr>
<td>Sapovirus</td>
<td>26,212</td>
<td>20,987</td>
<td>32,188</td>
<td></td>
<td>58,417</td>
<td>48,622</td>
<td>69,040</td>
<td>61,077</td>
<td>49,235</td>
<td>73,945</td>
<td>32,433</td>
<td>22,768</td>
<td>43,299</td>
<td></td>
<td></td>
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</tr>
</tbody>
</table>

Note this table gives estimates for Canada based on the UK study as they appear in the published paper and the those re-created using the models. Both figures have been provided to illustrate that the models for Canada re-created for this study give similar but not identical to those published. Only the published figures are given in figure 6 in the main paper.
Table 11: Foodborne Illness per 1 million population per year

<table>
<thead>
<tr>
<th>Pathogen</th>
<th>UK Mean</th>
<th>Lower 90%</th>
<th>Upper 90%</th>
<th>Australia Mean</th>
<th>Lower 90%</th>
<th>Upper 90%</th>
<th>Canada Mean</th>
<th>Lower 90%</th>
<th>Upper 90%</th>
<th>USA Mean</th>
<th>Lower 90%</th>
<th>Upper 90%</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>Int.</td>
<td>Int.</td>
<td>Mean</td>
<td>Int.</td>
<td>Int.</td>
<td>Mean</td>
<td>Int.</td>
<td>Int.</td>
<td>Mean</td>
<td>Int.</td>
<td>Int.</td>
</tr>
<tr>
<td><strong>Bacteria</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
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</tr>
<tr>
<td>C. perfringens</td>
<td>1,443</td>
<td>562</td>
<td>2,885</td>
<td>765</td>
<td>130</td>
<td>2,350</td>
<td>5,445</td>
<td>2,930</td>
<td>8,313</td>
<td>3,231</td>
<td>643</td>
<td>8,238</td>
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<tr>
<td>Campylobacter</td>
<td>4,706</td>
<td>2,247</td>
<td>7,831</td>
<td>8,400</td>
<td>5,050</td>
<td>13,650</td>
<td>4,472</td>
<td>2,944</td>
<td>6,553</td>
<td>2,856</td>
<td>1,127</td>
<td>5,388</td>
</tr>
<tr>
<td>E. coli O157</td>
<td>7</td>
<td>5</td>
<td>9</td>
<td>NA</td>
<td>NA</td>
<td>NA</td>
<td>395</td>
<td>161</td>
<td>700</td>
<td>211</td>
<td>59</td>
<td>500</td>
</tr>
<tr>
<td>Listeria monocytogenes</td>
<td>2</td>
<td>2</td>
<td>3</td>
<td>7</td>
<td>3</td>
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<td>4</td>
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<td>5</td>
<td>2</td>
<td>11</td>
</tr>
<tr>
<td>Salmonella non typhoidal</td>
<td>647</td>
<td>131</td>
<td>1,731</td>
<td>1,850</td>
<td>1,000</td>
<td>3,350</td>
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<tr>
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<td>3</td>
<td>67</td>
<td>16</td>
<td>6</td>
<td>40</td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cryptosporidium</td>
<td>47</td>
<td>7</td>
<td>139</td>
<td>80</td>
<td>57</td>
<td>320</td>
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<td>31</td>
<td>139</td>
<td>193</td>
<td>40</td>
<td>558</td>
</tr>
<tr>
<td>Giardia</td>
<td>285</td>
<td>42</td>
<td>821</td>
<td>175</td>
<td>35</td>
<td>490</td>
<td>239</td>
<td>146</td>
<td>365</td>
<td>257</td>
<td>171</td>
<td>367</td>
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<tr>
<td><strong>Viruses</strong></td>
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<td></td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Adenovirus</td>
<td>213</td>
<td>60</td>
<td>450</td>
<td>80</td>
<td>25</td>
<td>215</td>
<td>115</td>
<td>42</td>
<td>197</td>
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<td>NA</td>
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<tr>
<td>Astroviruses</td>
<td>45</td>
<td>11</td>
<td>101</td>
<td>60</td>
<td>20</td>
<td>160</td>
<td>59</td>
<td>21</td>
<td>105</td>
<td>52</td>
<td>19</td>
<td>89</td>
</tr>
<tr>
<td>Norovirus</td>
<td>5,768</td>
<td>5,768</td>
<td>5,768</td>
<td>12,920</td>
<td>3,620</td>
<td>26,300</td>
<td>32,238</td>
<td>20,910</td>
<td>44,125</td>
<td>18,267</td>
<td>10,793</td>
<td>27,791</td>
</tr>
<tr>
<td>Rotavirus</td>
<td>35</td>
<td>10</td>
<td>74</td>
<td>40</td>
<td>15</td>
<td>95</td>
<td>131</td>
<td>48</td>
<td>222</td>
<td>52</td>
<td>19</td>
<td>89</td>
</tr>
<tr>
<td>Sapovirus</td>
<td>669</td>
<td>466</td>
<td>915</td>
<td>700</td>
<td>350</td>
<td>1,150</td>
<td>292</td>
<td>109</td>
<td>483</td>
<td>52</td>
<td>19</td>
<td>89</td>
</tr>
</tbody>
</table>

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Table 12: Proportion of cases attributed to food by country by pathogen

<table>
<thead>
<tr>
<th>Pathogen</th>
<th>UK Mean</th>
<th>Lower 90%</th>
<th>Upper 90%</th>
<th>Lower 90% Cred.</th>
<th>Upper 90% Cred.</th>
<th>Australia Mean</th>
<th>Lower 90%</th>
<th>Upper 90%</th>
<th>Lower 90% Cred.</th>
<th>Upper 90% Cred.</th>
<th>Canada Mean</th>
<th>Lower 90%</th>
<th>Upper 90%</th>
<th>Lower 90% Cred.</th>
<th>Upper 90% Cred.</th>
<th>USA Mean</th>
<th>Lower 90%</th>
<th>Upper 90%</th>
<th>Lower 90% Cred.</th>
<th>Upper 90% Cred.</th>
</tr>
</thead>
<tbody>
<tr>
<td>Bacteria</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
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<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>C. perfringens</td>
<td>85%</td>
<td>73%</td>
<td>95%</td>
<td>98%</td>
<td>86%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
<td>100%</td>
</tr>
<tr>
<td>Campylobacter</td>
<td>48%</td>
<td>26%</td>
<td>71%</td>
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Figure S4: Proportion of cases attributed to food by pathogen by country (90% credible intervals shown with the mean)

Proportion of cases attributed to food

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