

1 **International travel as a risk factor for gastrointestinal infections in residents of North**
2 **East England**

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19 **Abstract**

20 International travel is thought to be a major risk factor for developing gastrointestinal illness for
21 UK residents. Here we present an analysis of routine laboratory and exposure surveillance data
22 from North East England, describing the destination-specific contribution that international travel
23 makes to the regional burden of gastrointestinal infection.

24 Laboratory reports of common notifiable enteric infections were linked to exposure data for
25 cases reported between 1 January 2013 and 31 December 2022. Demographic characteristics
26 of cases were described and rates per 100,000 visits determined using published estimates of
27 overseas visits from the Office for National Statistics International Passenger Survey.

28 34.9% of cases reported international travel during their incubation period between 2013 and
29 2022, although travel associated cases were significantly reduced (>80%) during the COVID-19
30 pandemic. Between 2013-2019, half of *Shigella spp* and non-typhoidal *Salmonella* infections,
31 and a third of *Giardia sp*, *Cryptosporidium spp* and Shiga-Toxin producing *Escherichia coli*
32 infections were following travel. Rates of illness were highest in travellers returning from Africa
33 and Asia (107.8 and 61.1 per 100,000 visits), with high rates also associated with tourist resorts
34 like Turkey, Egypt and the Dominican Republic (386.4-147.9 per 100,000 visits).

35 International travel is a major risk factor for the development of gastrointestinal infections. High
36 rates of illness were reported following travel to both destinations typically regarded as high risk
37 and common tourist resorts. This work highlights the need to better understand risks while
38 travelling to support the implementation of guidance and control measures to reduce the burden
39 of illness in returning travellers.

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43 **Introduction**

44 Gastroenteritis is a common cause of morbidity, with estimates suggesting up to 17 million
45 cases annually in the UK (1). While many cases of are relatively mild and short-lived,
46 particularly those caused by viral pathogens such as norovirus, others can result in more
47 prolonged or severe illness and may require hospitalization or lead to death. Bacterial and
48 parasitic pathogens, which are more commonly associated with severe outcomes, are usually
49 acquired through foodborne or waterborne routes, as opposed to viral pathogens which are
50 generally acquired through person-to-person transmission (2). In high income countries,
51 international travel is thought to be a major risk factor for gastrointestinal illness, particularly for
52 bacterial and parasitic pathogens. Risk is often associated with destination country, with
53 pathogens often endemic in lower- to middle- income (LMIC) destination countries, where
54 sanitation and hygiene is more often compromised.

55 Estimates suggest that up to 60% of international travellers will develop diarrhoea (3, 4), with
56 morbidity highest in those visiting LMICs. However, many studies are conducted within travel
57 clinic settings, which may bias findings towards travellers at greater risk of developing illness
58 due to the nature of their travel plans. Incidence of gastrointestinal illness associated with travel
59 is thought to have decreased over the last 20 years, particularly in travellers to countries that
60 were previously high risk but have seen considerable economic improvement, such as areas of
61 East Asia and South America (5). However, gastrointestinal illnesses remain one of the most
62 common health complaints reported by travellers, with areas such as South Asia and Africa
63 consistently reported as being associated with a higher risk of illness (3).

64 While destination of travel is thought to be the biggest risk factor, other factors influence the
65 likelihood of developing a gastrointestinal illness while travelling. These include type of travel,
66 with backpacking and visiting family thought to higher risk activities, and food choices taken (6).
67 In addition, certain groups have been shown to have increased susceptibility, including
68 individuals at extremes of age, those with immunosuppression, and those with gastrointestinal
69 conditions such as inflammatory bowel disease (5). Furthermore, international travel is a known
70 risk factor for acquisition of resistant organisms into the gut microbiota. Studies have shown that
71 a higher proportion of multidrug resistant gastrointestinal pathogens are isolated from patients
72 reporting recent travel outside the UK (7, 8).

73 Having a better understanding of travel associated enteric pathogens could help to improve pre-
74 travel guidance and support public health actions, which could ultimately lead to a reduction in

75 travel associated GI infections and, potentially, the importation of AMR, and a reduction in the
76 overall burden of GI infections in settings such as the UK. In England, all laboratory confirmed
77 cases of notifiable enteric infections are reported to UKHSA from all national health service
78 (NHS) laboratories via England's main infectious disease laboratory surveillance system, the
79 Second-Generation Surveillance System (SGSS). North East (NE) England is unique in that it
80 has its own surveillance system, EpiNorth3, which links routinely collected SGSS data,
81 laboratory typing data and exposure data from standardised exposure questionnaires. Here we
82 describe the epidemiology of gastrointestinal infections in residents of North East England
83 providing insight into the contribution that international travel makes to the overall and
84 pathogen-specific burden of gastrointestinal infection in the region.

85

86 **Methods**

87 Definitions and exclusions

88 Exposure questionnaires are undertaken with all North East residents notified with laboratory-
89 confirmed *Cryptosporidium* spp, *Giardia* sp, Hepatitis A, *Salmonella* spp (typhoidal and non-
90 typhoidal), *Shigella* spp, Shiga-toxin producing *Escherichia coli* (STEC; O157 and certain non-
91 O157 serotypes), *Vibrio* spp and *Yersinia* spp infections. Campylobacteriosis cases are
92 excluded from this study as exposure questionnaires are not routinely performed. Listeriosis
93 cases were also excluded from this study to avoid deductive disclosure due to low numbers.

94 Data on enteric infections reported to UKHSA between 1 January 2013 and 31 December 2022
95 were extracted from EpiNorth3 in January 2023. Cases were defined as being associated with
96 international travel if the case had a completed exposure questionnaire and reported travel
97 outside of the UK during the standardized incubation period specified in the exposure
98 questionnaire (7 days prior to onset: non-typhoidal *Salmonella* spp, *Shigella* spp, STEC,
99 *Yersinia* spp; 14 days prior to onset: *Cryptosporidium* spp and *Giardia* sp; 60 days prior to
100 onset: typhoidal *Salmonella* spp; 8 weeks prior to onset: Hepatitis A). UK acquired cases were
101 defined as cases with a completed exposure questionnaire who did not report travel outside of
102 the UK during the standardised incubation period. Cases without an exposure questionnaire
103 were defined as having an unknown travel status and were excluded from analyses unless
104 otherwise stated. Given the reduction in international travel reported in England during 2020 and
105 2021 as a result of the COVID-19 pandemic response, cases reported in 2020 and 2021
106 (pandemic years) were also excluded from analyses unless otherwise stated.

107 Analysis

108 All analyses were performed using R studio version 4.2.0. Demographic data including ethnicity,
109 sex and age were extracted from EpiNorth3. Deprivation and urban/rural classification of
110 residence were derived from postcode of residence recorded in EpiNorth3 using the publicly
111 available English indices of deprivation 2019 dataset (9) and the 2011 rural-urban classification
112 (RUC2011) dataset (10). Directly standardised rates of illness per 100,000 population were
113 calculated for age and ethnic group with denominator data on the North East England
114 population taken from the 2021 census and 2021 mid-year population estimates (11), with 95%
115 confidence intervals calculated using the Dobson Method. Chi squared tests were performed for
116 categorical variables.

117 Destination countries reported in exposure questionnaires were extracted from EpiNorth3.
118 Destinations reported as resorts or cities and incorrectly spelled destinations were recoded.
119 Where multiple locations were recorded during an incubation period, the location was recoded
120 to 'Multiple/unspecified'. Countries were recoded based on nomenclature used in the UK Office
121 for National Statistics (ONS) International Passenger Survey (IPS) Travepac dataset, to
122 account for sovereignty (12). Within the EpiNorth3 dataset there was no distinction between
123 Northern Cyprus and the Republic of Cyprus, therefore both are reported as Cyprus.

124 Using published estimates from the ONS IPS, it was possible to establish the most common
125 travel destinations for North East England residents. Using visits as a denominator, rates of
126 illness were determined by destination. Countries were grouped into global regions (Africa, Asia,
127 America and Caribbean, Europe, Middle East and Rest of World) as specified in the Travepac
128 dataset. Rates per 100,000 visits were calculated using the total number of visits to each
129 country or country group between 2013 and 2019 calculated using the 'Final weight' variable in
130 the Travepac dataset for 2013-2019 and the total number of cases reporting travel to the
131 location between 2013 and 2019. At the time of analysis, Travepac data was unavailable for
132 2020-2022.

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137 Results

138 Between 2013 and 2022, 9,358 laboratory confirmed cases of gastrointestinal illness resulting
139 from infection with *Cryptosporidium* spp, *Giardia* sp, Hepatitis A, *Salmonella* spp (typhoidal and
140 non-typhoidal), *Shigella* spp, Shiga-toxin producing *Escherichia coli* (STEC; O157 and certain
141 non-O157 serotypes), *Vibrio* spp and *Yersinia* spp were reported in North East England
142 residents. Routine exposure questionnaires were completed for 7,909 cases (84.5%), of which
143 2,764 cases (34.9%) reported international travel during their incubation period.

144

145 Travel as a risk factor over time

146 The proportion of cases associated with international travel remained consistent between 2013
147 and 2019 (average 38.0%; 95% CI: 35.9-40.1%; range 33.6-41.6%; Chi2: p=0.96; **Figure 1**).

148 During England's COVID-19 pandemic response in 2020 and 2021, total gastrointestinal
149 infections (travel associated, UK-acquired and unknown exposures; n=480 in 2020 and n=654
150 in 2021) were significantly lower than historic figures (2013-2019 average: 1,038; 95% CI:947-
151 1,129). Reductions in travel associated infections were greater than reductions in UK acquired
152 infections (travel associated infections; -82.5% change in 2020 and -86.6% in 2021 vs. UK
153 acquired infections; -42.9% change in 2020 and -16.3% change in 2021). In 2022,
154 gastrointestinal infection reports returned to pre-pandemic levels predominantly because of
155 increases in travel associated cases (total n=956; travel associated: n= 303; UK acquired: n=
156 493), with the proportion of cases reporting travel comparable to pre-pandemic years (38.1%).

157 In non-pandemic years, where exposure was known (n=7,026; 2,660 reporting travel; 37.9%),
158 infections with *Vibrio* species and Typhoidal *Salmonella* were exclusively associated with
159 international travel, while around half of infections with Hepatitis A, *Shigella* spp and non-
160 Typhoidal *Salmonella* were travel acquired (**Table 1** and **Supplementary Figure 1**). Infections
161 caused by *Giardia* sp, *Cryptosporidium* spp and O157 STEC were less commonly associated
162 with travel (31.7%, 28.1% and 20.8% of infections respectively). Although average annual
163 numbers of infections associated with travel were relatively low for some pathogens (*Vibrio* spp:
164 n=<5; Typhoidal *Salmonella*: n=8; **Table 1**), others contributed considerably to annual
165 gastrointestinal morbidity in the region (*Salmonella*: n=159).

166 Between 2012 and 2019 the percentage of total cases associated with travel remained
167 consistent for most pathogens apart from *Shigella* spp, (Chi2 p=0.02) where an increase in UK

168 acquired cases has been observed since 2013, and STEC O157 (Chi2 $p < 0.001$) where an
169 increase in internationally acquired cases was reported in 2019 (**Supplementary Figure 2**).

170

171 Demographic characteristics

172 The demographic characteristics of cases diagnosed with common gastrointestinal infections
173 following international travel were compared with individuals who acquired their infection in the
174 UK (**Table 2**). The proportions of male (38.5%) and female (37.2%) reporting travel was similar
175 ($p = 0.27$). The percentage of infections acquired in the UK was significantly higher than
176 infections associated with travel for all age groups; however, children aged under 9 years and
177 adults aged over 60 years were significantly more likely to have acquired their infection in the
178 UK. (**Table 2, Supplementary Figure 3 and Supplementary Table 1**). Ethnicity was poorly
179 completed; however, where available, individuals of Asian ethnicity were more likely to have
180 acquired their infection during international travel (acquired abroad: 66.7%, $n = 114$ vs. 33.3%,
181 $n = 57$ acquired in UK), with the rate of reported travel associated infection in those of Asian
182 ethnicity (152.8; 95% CI: 126.1 - 183.6) significantly higher than the rate for those of White
183 ethnicity (59.1; 95% CI: 56.1 - 62.2).

184

185 Temporal distribution of travel associated cases

186 Travel associated cases were highest in the summer with average reported cases in August and
187 September significantly higher than other months (**Figure 2**). The number of travel associated
188 cases were significantly lower than the number of UK acquired cases for all months except
189 between June and September. The monthly distribution of cases was dependent on
190 geographical region of travel (**Supplementary Figure 4**). There was less variability in the
191 monthly number of UK acquired cases; however, the number of cases reported in September
192 and October were significantly higher than numbers reported in other months. Travel associated
193 cases corresponded with visits abroad, which were highest in August (333,054 visits; 95%
194 CI: 282,456-383,652) and September (290,153 visits; 95% CI: 241,662-338,643. However, when
195 taking visits into account, rates of illness per 100,000 visits remained highest in August (20.8)
196 and September (22.9) and were lowest in February (8.0).

197

198 Destination of travel

199 Between 2013 and 2019, 2,357 cases had a country of travel reported (100.0 % of cases
200 reported between 2013 and 2019). Of these, 2,284 reported travel to a single country (96.9%).
201 The most common destination country reported by cases was Spain (n=510), followed by
202 Turkey (n=322), India (n=145) and Egypt (n=131). 47.0% of cases reported travel to one of
203 these four countries (n=1108). Between 2013 and 2019, Spain (including the Balearic Islands)
204 was the most frequently visited destination for North East England residents with an estimated
205 4,548,582 visits made over the period (649,797 average annual visits; **Supplementary Figure**
206 **5**). France (1,226,916 total and 175,274 average annual visits), the Canary Islands (1,109,696
207 total and 158,528 average annual visits) and the USA (771,945 total and 110,278 average
208 annual visits) were also common destinations. All destinations with over 100,000 average
209 annual visits were within Europe or the USA.

210 Rates of illness per 100,000 visits across the period were highest in travellers who visited Africa
211 (107.8 per 100,000 visits; 311 cases) and Asia (61.1 per 100,000 visits; 441 cases) and lowest
212 in travellers visiting European countries (excluding UK; 9.4 per 100,000 visits; 1,149 cases).
213 Rates of hepatitis A and typhoidal salmonella were highest in travellers to Asia and rates of
214 vibrio were comparable for travellers to both Africa and Asia (**Table 3**). Rates for all other
215 pathogens were highest in travellers returning from Africa. The likelihood of acquiring shigella in
216 travellers to Africa was 109 times higher than in travellers to Europe, while the rate of acquiring
217 non-Typhoidal salmonella was 527 times higher in travellers to Asia when compared to
218 travellers to Europe (**Table 3**).

219 Of the 20 countries reporting a rate of illness of over 100 cases per 100,000 visits (classified
220 here as high risk), only Turkey (147.9 per 100,000), India (110.6 per 100,000) and Tunisia
221 (101.5 per 100,000) had more than 10,000 visitors annually (**Table 4**). Of note, high rates of
222 illness were also associated with tourist destinations such as Egypt (386.4 per 100,000 visits)
223 and the Dominican Republic (244.2 per 100,000 visits), which receive fewer than 10,000 visitors
224 annually but like Turkey are also popular tourist destinations. The highest rate of illness was
225 reported from travellers to Nepal (769.4 per 100,000 visits), but less than 250 North East
226 residents were estimated to visit Nepal each year. Rates of illness were high from countries in
227 South Asia and Africa, including Kenya (400.9 per 100,000), Pakistan (252.0 per 100,000) and
228 Cambodia (113.7 per 100,000). Several countries in South and Central America also had high
229 rates of illness per 100,000 visits (Colombia 208.6; Ecuador 169.5 and Peru 139.1).

230 Of the 2,404 individuals with routinely collected exposure data it was possible to identify the
231 type of accommodation used while travelling for 1,868 cases (77.7%). 92.5% of cases visiting
232 Europe, 86.2% visiting the Americas, 84.7% visiting Africa and 83.8% visiting Asia stayed in
233 hotels. Staying with family and friends while travelling was less commonly reported; 5% of cases
234 reporting travel to Africa, 4.2% of cases travelling to Asia, 3% of cases travelling to the
235 Americas and 2.1% of cases travelling to Europe. 1,233 cases reported named premises of
236 which 1,058 premises were unique and were only reported by one case (85.8%). The remaining
237 premises were associated with clusters of between 2 and 13 cases (median: 2, IQR: 1).

238 Clusters, defined as two or more cases, were most commonly associated with salmonella
239 ($n=54$) and *Cryptosporidium spp* ($n=41$), fewer than 10 clusters were reported for each of
240 *Giardia*, *Shigella* or STEC (O157) or STEC (non-O157). Salmonella outbreaks were
241 predominantly associated with travel to Turkey ($n=19$ clusters, $n=42$ cases), Egypt ($n=11$
242 clusters; $n=23$ cases) and Mexico ($n=6$; 13 cases). *Cryptosporidium* outbreaks were
243 predominantly associated with Spain ($n=17$; 39 cases), Turkey ($n=6$; $n=20$ cases), the Canary
244 Islands ($n=4$; $n=16$ cases) and Egypt ($n=4$; $n=9$ cases). Overall, eleven hotels had clusters
245 reported on two separate years and 4 hotels reported clusters on three separate years.

246

247 Discussion

248 Through this analysis of laboratory and exposure data for cases of notifiable gastrointestinal
249 infections in North East England we show that international travel is a major risk factor,
250 contributing substantially to the burden of infection in the region. Furthermore, as there has
251 been no reduction in the proportion of travel associated infections in non-pandemic years since
252 2013 this work highlights the need to better understand the risk factors associated with
253 developing gastrointestinal illness while travelling.

254 The considerable decline in gastrointestinal infections observed during the COVID-19 pandemic
255 was likely driven by a reduction in travel associated infections. This suggests the overall burden
256 of GI illness could be reduced if improvements were made to the number of individuals
257 acquiring an illness while travelling abroad, particularly as returning travellers may be seeding
258 illness and on-going transmission across the wider population within the UK (13). Pathogen
259 specific reductions in GI infections were also observed in England overall during the COVID-19
260 pandemic, with diagnoses of pathogens such as salmonella and cryptosporidium, which are

261 commonly associated with foreign travel, remaining lower than infections with pathogens such
262 as STEC which are often UK acquired (13, 14).

263 The strength of this study is that it used denominator data for international travel for the North
264 East England population allowing rates to be determined. Country specific case numbers may
265 correlate with the volume of travel to a destination, which makes it challenging to draw
266 conclusions on the destination specific risks. For example, Spain was the most commonly
267 reported travel destination of cases, but was also the most common destination of travel for
268 North East England residents, with the rate of illness per visit similar to that reported for other
269 European countries. Conversely, travel to countries in Africa and Asia was less common for
270 North East England residents, but it was associated with a high risk of illness.. With
271 globalisation, changes in travel patterns and an increasing non-UK born population in the North
272 East England, it is possible that visits to high risk destinations will increase (15).

273 Travel to high-risk countries to visit friends and relatives is a known risk factor for
274 gastrointestinal infections (16), with 75% of enteric fever cases occurred in individuals travelling
275 to visit friends and relatives and high rates observed among individuals of Pakistani or South
276 Asian ethnicity (17). In our study, where ethnicity was completed, those of Asian ethnicity were
277 more likely to have acquired their infection during international travel, with the rate of
278 international travel associated with Asian ethnicity significantly higher than for those of white
279 ethnicity. Due to small numbers, there was insufficient data available to demonstrate that higher
280 rates of illness in those of Asian ethnicity were the result of travel to visit friends and relatives,
281 but the study did demonstrate that a higher proportion of cases reported as visiting countries in
282 Asia were staying with friends or family. However, it has also been shown that residents from
283 ethnic minorities in high-income countries have lower health literacy with language proficiency
284 and lower social support identified as key barriers (18). Future work looking at infections across
285 England overall could provide further evidence as to why rates of illness are higher in those of
286 Asian ethnicity.

287 While the findings of this study do not indicate absolute risk associated with travel to specific
288 areas, they do allow for comparisons in patterns of illness between countries. High rates of
289 illness were reported following travel to countries or regions which were documented in other
290 studies and in travel guidance to be 'high risk' for travel associated GI infections (3, 4, 19). This
291 study additionally highlights increased rates of illness associated with 'all-inclusive' holiday
292 destinations including the Dominican Republic, Turkey, and Egypt, with rates per 100,000 visits

293 as high as destinations commonly categorised as 'high risk' (12). This has also been reported in
294 other studies with the Dominican Republic shown to have the 3rd highest number of all-pathogen
295 travel related diagnoses in returning travellers reported in the United States GeoSentinel
296 Network between 2012 and 2021, after Mexico and India (19). All-inclusive travel to low- and
297 middle-income countries may be perceived as lower risk as this type of travel and companies
298 offering it are often mainly associated with lower risk destinations such as high-income countries
299 in western Europe. Higher rates of illness reported from Turkey and Egypt may also be
300 associated with outbreak activity at hotel resorts. Over the period, 175 hotels were associated
301 with more than one case with clusters more commonly reported in travellers to Turkey and
302 Egypt.

303 As travel associated infections are only included if diagnosed following return to the North East
304 England, this may lead to an underestimation of infections, particularly those that may be short-
305 lived or less severe (20). Conversely, there may be an overestimation of travel as a cause of
306 illness with primary care physicians often more often arranging stool testing for individuals
307 reporting international travel than for those with similar symptoms without a history of travel (21).
308 It has also been shown previously that travel as a risk factor may be overestimated, with cases
309 associated with domestic transmission misclassified as travel associated when shorter
310 incubation period durations are taken into account (22). A further limitation is that denominators
311 are estimates based on survey data and may not fully reflect travel patterns of North East
312 England residents.

313 This study highlights that international travel remains a common risk factor for enteric infections.
314 However, it was not possible to explore in detail the risks while travelling using secondary
315 analysis of routinely collected data due to the unstructured nature of data collected. Given the
316 large proportion of diagnosed cases acquiring their infection abroad we recommend that further
317 studies are undertaken to collect structured travel specific data from cases diagnosed with
318 gastrointestinal infections following travel, and that this be considered within routine
319 surveillance, to help inform public health messages aimed at prevention and reduction of travel
320 associated gastrointestinal illness in travellers.

321

322 **Ethical statement:** This study was conducted under the provisions of Section 251 of the NHS
323 Act 2006 and therefore did not require individual patient consent. The authors affirm that the
324 manuscript is an honest, accurate, and transparent account of the study being reported; that no

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327

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334

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336

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340

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342

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344 D.W. contributed to the interpretation of the results. N.L. wrote the manuscript with input from all
345 authors.

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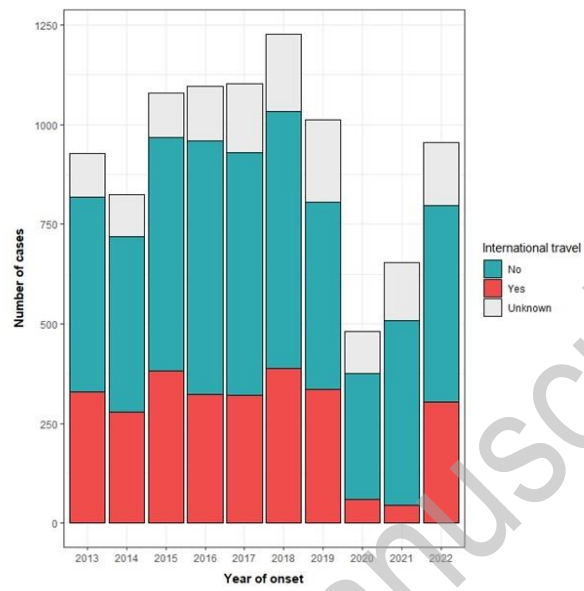
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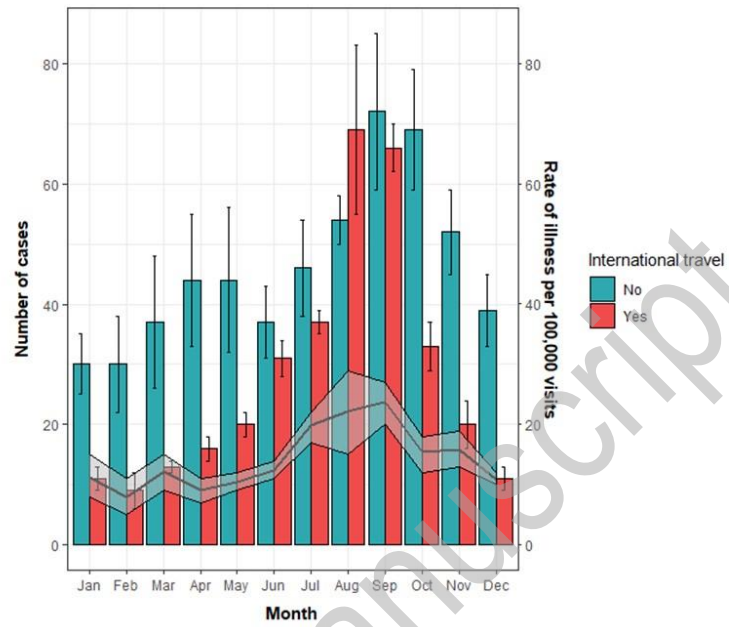
425 Figure 1



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430 **Table 1 – Gastrointestinal infections reported in North East residents in (2013-2019 average) by pathogen and travel exposure status.**

Pathogen	Exposure duration (days prior to onset)	Average annual number of infections	Annual rate per 100,000 population	Exposures recorded		International travel reported		UK acquired	
				Number of cases with travel exposure completed	(% with travel exposure completed)	Number of cases with reported travel	(% cases with reported travel)	Number of UK acquired infections, with travel exposure completed	(% UK acquired, with travel exposure completed)
Cryptosporidium spp	14	299	11.5	263	88.0	74	28.1	189	71.9
Giardia sp	14	226	8.7	180	79.6	57	31.7	123	68.3
Hepatitis A	8 weeks	8	0.3	<5	50.0	<5	50.0	<5	50.0
Non-typhoidal Salmonella	60	372	14.3	326	88.2	159	48.5	169	51.5
Shigella spp	7	49	1.9	43	87.8	20	46.3	23	53.5
STEC Non-O157	7	16	1.9	7	43.8	<5	28.6	5	71.4
STEC O157	7	49	0.6	48	98.0	10	20.8	38	79.2
Typhoidal Salmonella	7	7	0.3	7	100.0	7	100.0	0	0.0
Vibrio spp		6	0.2	<5	66.7	<5	100.0	0	0.0
Yersinia spp	7	5	0.2	<5	80.0	<5	25.0	<5	75.0

431

432 **Table 2 – Demographic characteristics of North East residents diagnosed with gastrointestinal infections between 2013 and 2019 with**
 433 **travel exposure information available**

Demographic characteristic		International travel reported			UK acquired		Prevalence Ratio	P value
		Number of cases with travel exposure completed	Number of cases with reported travel	(% cases with reported travel)	Number of UK acquired infections, with travel exposure completed	(% UK acquired, with travel exposure completed)		
Sex	Male	3391	1306	38.5	2085	61.5	0.63	0.27
	Female	3635	1354	37.2	2281	62.8	0.59	
Age group	0 – 9 years	1600	443	27.7	1157	72.3	0.38	<0.001
	10 – 19 years	555	241	43.4	314	56.6	0.77	
	20 – 29 years	1067	459	43.0	608	57.0	0.75	
	30 – 39 years	1127	449	39.8	678	60.2	0.66	
	40 – 49 years	841	358	42.6	483	57.4	0.74	
	50 – 59 years	818	365	44.6	453	55.4	0.81	
	>60 years	1018	345	33.9	673	66.1	0.51	
Ethnicity	Asian	171	114	66.7	57	33.3	2.00	<0.001
	Black	23	10	43.5	13	56.5	0.77	
	Mixed	42	18	42.9	24	57.1	0.75	
	Other	30	14	46.7	16	53.3	0.88	
	White	3753	1463	39.0	2290	61.0	0.64	
	1	1125	376	30.4	749	66.6	0.50	
	2	861	259	30.1	602	69.9	0.43	

Demographic characteristic	International travel reported				UK acquired		Prevalence Ratio	P value		
	Number of cases with travel exposure completed	Number of cases with reported travel	(% cases with reported travel)	Number of UK acquired infections, with travel exposure completed	(% UK acquired, with travel exposure completed)					
Index of multiple deprivation	3	757	286	37.8	471	62.2	0.61	<0.001		
	4	699	250	35.8	449	64.2	0.56			
	5	505	196	38.8	309	61.2	0.63			
	6	401	152	37.9	249	62.1	0.61			
	7	484	196	40.5	288	59.5	0.68			
	8	498	229	46.0	269	54.0	0.85			
	9	508	231	45.5	277	54.5	0.83			
	10	392	182	46.4	210	53.6	0.87			
	Rural/Urban residence	Rural	1063	363	34.1	700	65.9		0.52	0.007
		Urban	5167	1994	38.5	3173	61.5		0.63	
Duration of illness	Median (IQR)	8 days (7)	10 days (7)		8 days (6)			<0.001		
Hospital admission	Yes	1131	370	32.7	761	67.3	0.49	<0.001		
	No	4575	1760	38.5	2815	61.5	0.63			

435 **Table 3 - Rates of illness per 100,000 visits by pathogen and geographical region of travel**

436

Pathogen	Europe		Africa		America and Caribbean		Asia		Middle East		Rest of world	
	Rate	RR	Rate	RR	Rate	RR	Rate	RR	Rate	RR	Rate	RR
Cryptosporidium	3.1	ref	18.4	6.0	3.9	1.3	5.0	1.6	2.5	0.8	0.8	0.3
Giardia	1.4	ref	15.6	11.4	4.9	3.6	13.2	9.6	1.6	1.2	5.6	4.1
Hepatitis A	0.1	ref	-	-	0.2	2.6	0.7	9.9	-	-	-	-
Salmonella	4.3	ref	56.8	13.2	12.2	2.8	29.3	6.8	8.3	1.9	8.1	1.9
Shigella	0.1	ref	14.2	109.2	1.4	10.5	8.2	62.9	-	-	2.01	15.5
STEC Non-O157	0.07	ref	1.4	19.9	-	-	-	-	-	-	0.4	5.7
STEC O157	0.4	ref	3.1	7.4	0.3	0.6	0.3	0.7	0.63	1.5	0.4	1.0
Typhoidal Salmonella	0.01	ref	0.4	35.0	0.2	18.0	5.3	527.0	0.63	63.0	-	-
Vibrio	0.02	ref	2.1	104.0	0.3	13.5	2.2	111.0	-	-	0.4	20.0
Yersinia	0.02	ref	0.7	34.5	-	-	0.1	7.0	0.32	16.0	-	-

437

438 Rates per 100,000 visits were calculated using the total number of visits to each country or country group
 439 between 2013 and 2019 calculated using the 'Final weight' variable in the Travelpac dataset for 2013-
 440 2019 and the total number of cases reporting travel to the location between 2013 and 2019.

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442

443 **Table 4 - Total cases per 100,000 visits by destination country indicating average**
 444 **annual number of visitors per country**

Cases per 100,000 visits	Destination country
< 10	Australia, Austria, Belgium, Canary Islands, China (excl Taiwan)/Tibet, Croatia, Czech Republic, France/Corsica, Germany, Holland, Hungary, Iceland, Irish Republic, Israel, Italy/Sardinia, Kuwait, Latvia, Madeira/Azores, Malaysia, New Zealand, Norway, Poland, Portugal, Romania, Switzerland, USA
10.1 - 20	Greece/Crete/Rhodes, Hong Kong, Iran, Japan, Kazakhstan, Maldives, Russia, Slovakia, South Africa, Spain, Trinidad & Tobago, United Arab Emirates
20.1 - 30	Argentina, Bolivia, Brazil, Cyprus, Democratic Republic of Congo, Lebanon, Libya, Malta, Mauritius, Nigeria, Philippines, Qatar
30.1 - 40	Azerbaijan, Barbados, Bulgaria, Iraq, Jordan, Nevis/St Kitts, Singapore
40.1 - 50	Gambia, Jamaica, Mongolia, Saudi Arabia, Sri Lanka
50.1 - 60	Costa Rica, Montenegro, Thailand
60.1 - 70	Mexico, Morocco, Serbia, Vietnam
70.1 - 80	Antigua, Malawi, Uzbekistan
80.1 - 90	Bangladesh, Cape Verde Islands
90.1 - 100	Bali/Borneo/Indonesia, Cuba, Uganda, Zambia
100.1 - 200	Afghanistan, Burkina Faso, Cambodia/Kampuchea, Ecuador, Equatorial Guinea, Ghana, India, Namibia, Peru, Tanzania, Tunisia, Turkey
200.1 - 300	Colombia, Dominican Republic, Ethiopia, Madagascar, Pakistan
>300.1	Egypt, Kenya, Nepal, North Sudan, Somalia

445
 446 Rates per 100,000 visits were calculated using the total number of visits to each country or country group
 447 between 2013 and 2019 calculated using the 'Final weight' variable in the Travepac dataset for 2013-
 448 2019 and the total number of cases reporting travel to the location between 2013 and 2019.

449 Countries in blue have <10,000 visitors annually, those in purple have between 10,000 and 50,000 visits
 450 annually and those in Red have more than 50,000 visits annually.

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452

453 **Table 5 - Rate ratios for travel destinations compared to Spain (reference country)**
 454 **indicating average annual number of visitors per country**

RR	Destination country
< 1.00	Australia, Austria, Belgium, Canary Islands, China (excl Taiwan)/Tibet, Croatia, Czech Republic, France/Corsica, Germany, Holland, Hungary, Iceland, Irish Republic, Israel, Italy/Sardinia, Kuwait, Latvia, Madeira/Azores, Malaysia, New Zealand, Norway, Poland, Portugal, Romania, Switzerland, USA
1.00	Spain (reference)
1.01 – 2.00	Greece/Crete/Rhodes, Hong Kong, Iran, Japan, Kazakhstan, Maldives, Russia, Slovakia, South Africa, Trinidad & Tobago, United Arab Emirates
2.01 – 3.00	Argentina, Bolivia, Brazil, Cyprus, Democratic Republic of Congo, Lebanon, Libya, Malta, Mauritius, Nigeria, Philippines, Qatar, Singapore
3.01- 4.00	Azerbaijan, Barbados, Bulgaria, Iraq, Jordan, Mongolia, Nevis/St Kitts
4.01 – 5.00	Gambia, Jamaica, Saudi Arabia, Sri Lanka
5.01 – 6.00	Costa Rica, Montenegro, Thailand, Vietnam
6.01 – 7.00	Mexico, Morocco, Serbia
7.01- 8.00	Antigua, Cape Verde Islands, Malawi, Uzbekistan
8.01 – 9.00	Bali/Borneo/Indonesia, Bangladesh, Uganda
9.01 – 10.00	Cuba, Tunisia, Zambia
10.01 – 20.00	Afghanistan, Burkina Faso, Cambodia/Kampuchea, Ecuador, Equatorial Guinea, Ghana, India, Namibia, Peru, Tanzania, Turkey
20.01 – 30.00	Colombia, Dominican Republic, Ethiopia, Madagascar, Pakistan
30.01 – 40.00	Egypt, Kenya, Somalia
>40.01	Nepal, North Sudan

455

456 Rates per 100,000 visits were calculated using the total number of visits to each country or country group
 457 between 2013 and 2019 calculated using the 'Final weight' variable in the Travelpac dataset for 2013-
 458 2019 and the total number of cases reporting travel to the location between 2013 and 2019. Rates were
 459 compared to a reference country (Spain), which was the most common destination of travel for North East
 460 residents between 2013 and 2019.

461 Countries in blue have <10,000 visitors annually, those in purple have between 10,000 and 50,000 visits
 462 annually and those in Red have more than 50,000 visits annually.

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