

# Risk factors for symptoms of infection and microbial carriage among French medical students abroad

Thi Loi Dao, Naomie Canard, van Thuan Hoang, Tran Duc Anh Ly, Tassadit Drali, Laetitia Ninove, Florence Fenollar, Didier Raoult, Philippe Parola, Pierre Marty, et al.

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1	Risk factors for symptoms of infection and microbial carriage among French medical students
2	abroad
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## 20 Abstract

Objectives: To investigate symptoms of infections and their risk factors among French medical
 students undertaking an internship abroad.

Methods: Clinical follow up and qPCR-based respiratory, gastrointestinal, and vaginal pathogen
 carriage were prospectively assessed pre-travel and post-travel, in a cohort of medical students
 departing from Marseille, France.

26 Results: 293 students were included. 63.5%, 35.8% and 3.6% of students reported gastrointestinal, 27 respiratory, and vaginal symptoms, respectively. The acquisition rate of Enteroaggregative 28 Escherichia coli and Enteropathogenic E. coli was 40.9% and 18.6%, respectively. A significant 29 increase was observed for rhinovirus and Streptococcus pneumoniae by comparing the prevalence of 30 pathogens in pre-travel and post-travel samples. Gardnerella vaginalis and Atopobium vaginae 31 acquisition rates were 12.9% and 13.9%, respectively. Being female, primarily travelling to 32 Vietnam, and living in basic accommodation conditions were independent risk factors for reporting 33 respiratory symptoms. Students reporting respiratory symptoms were three times more likely to 34 acquire S. pneumoniae. Travelling primarily to north India and Senegal were independent risk 35 factors for diarrhoea.

36 Conclusion: This study makes it possible to identify the main infectious diseases linked to travel in
37 a group of French medical students undertaking an internship abroad and the risk factors on which
38 to base targeting students for reinforced pre-travel advice.

39 Keywords: respiratory symptoms, travellers' diarrhoea, medical students, Streptococcus
40 pneumoniae, E. coli, risk factor

41

## 42 Introduction

International travellers are exposed to the acquisition of potential pathogens including viruses, bacteria or parasites with the risk of community or hospital spread upon return, whether or not they present health problems during their trip. Therefore, there is a risk of pathogens being imported into France from endemic areas abroad, either from foreign travellers visiting France, from French travellers visiting foreign countries, or via migrants and expatriates treated in France with a risk of indigenous spread. This has been extensively described, by example, among French Hajj pilgrims travelling to Mecca, Saudi Arabia [1].

50 An Australian organisation, Work the World, has enabled 15,000 medical students to take part in 51 internships abroad since 2005 [2]. Developing countries are becoming popular destinations of 52 internship. Medical internships abroad are generally hospital immersion experiences but, young 53 medical students also participate in humanitarian missions that are unrelated to clinical activities 54 such as school renovation for example [3, 4]. In one Australian survey, 64% of students experienced 55 some sort of health problems while taking part in electives abroad, and travellers' diarrhoea was the 56 most common problem (40%) [5]. In our preliminary report on a cohort of 134 French medical 57 students participating in international electives, we showed that 73.9%, 38.8% and 5% of them 58 reported gastrointestinal, respiratory, and vaginal symptoms, respectively [4]. We showed that the 59 acquisition rate of Enteropathogenic *Escherichia coli* (EPEC) and Enteroaggregative *E. coli* (EAEC) 60 was 41% and 53%, respectively. By contrast, the acquisition of respiratory viruses was low but was 61 associated with persistent respiratory symptoms at return. Respiratory bacterial acquisition ranged 62 from 3.3% for Streptococcus pyogenes to 15.0% for Haemophilus influenzae. Atopobium vaginae 63 and Gardnerella vaginalis percentage of acquisition were 14.3% and 7.7%, respectively. So far, to 64 our knowledge, the risk factors for acquisition of pathogens have not been clearly identified among 65 medical students abroad to date. The relationship between symptoms and the carriage of pathogens

also remain poorly understood, making it difficult to distinguish between infection and colonisation.
We aim to conduct this study to investigate the risk factors for symptoms of infections among
French medical students undertaking an internship abroad.

69

## 70 Materials and methods

## 71 Study design

72 A monocentric prospective cohort survey was conducted over two years (2018-2019) among 73 medical students from the Faculty of Medicine in Marseille, France who were planning to take part 74 in an internship abroad during the summer. Recruitment was performed on a voluntary basis, during 75 their vaccination and pre-travel consultation at the Institut Méditerranée Infection which is on the 76 Marseille University medical campus. Participants were asked to complete an inclusion 77 questionnaire including demographic data, history of chronic illness, intended travel dates and 78 destination. All participants received advice regarding the prevention of diarrhoea during travel 79 (hand hygiene, safe food, and water habits) but prophylaxis for traveller's diarrhoea was not 80 prescribed. Because dates of departure and return are different for each student, the samples were 81 not taken at once. The participants were given two sets of "pre-travel" and "post-travel" kits which 82 contained questionnaire and sampling equipment (commercial rigid cotton-tipped swab applicators 83 and viral transport media). They were also instructed how to self-collect samples, as following: 3 cm 84 in the nostril, 5 turns and post wall of the pharynx, 5 streaks for respiratory samples; rectal samples 85 were collected using two methods: rectal self-sampling when having a bowel movement, 3 86 centimetres through the anus, gently rubbing the inner walls of the rectum several times or stool 87 collection after emission; vaginal samples were collected by placing the swab about three 88 centimetres in the vagina and gently rubbing the inner walls several times, avoid touching the skin

89 and vulva with the swab. Samples were self-collected using commercial rigid cotton-tipped swab 90 applicators (Medical Wire & Equipment, Wiltshire, UK) and placed in viral transport media (Sigma 91 Virocul<sup>®</sup>) for further process at our laboratory. A document of instructions for self-sampling was 92 also provided in the kits. During the week before travel, each student was invited to deposit their 93 self-collect samples and a pre-travel questionnaire that collected information about their health 94 problems and antibiotic use if applicable. After their travel, they were invited to self-collect samples 95 during the week following their return to France. Students were also provided with a post-travel 96 questionnaire addressing the exact place of the internship, the type of activities during their stay, 97 including tourism and travel to other countries over the internship period. Accommodation 98 conditions, contact with animals or children, symptoms, onset of symptoms and treatment during 99 their stay were also documented. Influenza-like illness (ILI) was defined as sore throat, cough plus 100 subjective fever [6]. Diarrhoea was defined by at least three loose or liquid stools per 24 hours.

## 101 Microbiological methods

102 The methods for identifying respiratory, gastro-intestinal, and vaginal pathogens by PCR assay are 103 detailed elsewhere [4]. The followed respiratory pathogens were screened: Influenza A and B 104 viruses, Human coronaviruses, Human parainfluenza viruses, Human rhinovirus, Human 105 metapneumovirus, Adenovirus, Respiratory syncytial virus, Staphylococcus aureus, H. influenzae, 106 Streptococcus pneumoniae, Klebsiella pneumoniae, and S. pyogenes. The following gastro-intestinal 107 pathogens were screened for: Norovirus, Rotavirus, Adenovirus, Astrovirus, Entamoeba histolytica, 108 Giardia lamblia, Cryptosporidium spp, Salmonella spp, Shigella spp (EIEC), Enterohemorrhagic E. 109 coli (EHEC), Enteropathogenic E. coli (EPEC), Enteroaggregative E. coli (EAEC), Campylobacter 110 *jejuni* and *Tropheryma whipplei*. The following vaginal pathogens were screened for: *Chlamydia* 111 trachomatis, Mycoplasma genitalium, Neisseria gonorrhoeae, Trichomonas vaginalis, Mycoplasma 112 hominis, A. vaginae and G. vaginalis.

113 PCR were considered positive for virus or bacteria detection when the cycle threshold (CT) value 114 was  $\leq$ 35. Bacterial vaginosis was defined by a *G. vaginalis* DNA load  $\geq$  10<sup>9</sup> copies/mL (CT $\leq$  18)

115 and/or an A. vaginae DNA load  $\geq 10^8$  copies/mL (CT $\leq 21$ ), as previously reported [7].

116 The acquisition of a pathogen was defined as negative before travel and positive when returning to

117 Marseille, France.

## 118 Statistical analysis

119 STATA software version 14.2 was used to conduct statistical analysis. Differences in the 120 proportions were tested by using Fisher's exact or Pearson's chi-square tests, when appropriate. 121 McNemar's test was used to evaluate the potential acquisition of pathogens (prevalence after versus 122 before travel). Clinical symptoms during travel were reported only if onset of symptoms took place 123 during travel. Univariate analysis was used to evaluate unadjusted associations between the 124 prevalence of symptoms during travel and multiple factors. A p-value <0.05 was considered to be 125 statistically significant. Only variables with a prevalence equal or more than 5.0% were considered 126 for statistical analysis. Variables with p-values <0.2 in the univariate analysis were included in the 127 multivariate analysis. Log-binomial regression was used to estimate factors' adjusted risk ratios for 128 symptoms of infections.

129

## 130 Ethics

131 The protocol was approved by our Institutional Review Board (2019-006). It was performed in 132 accordance with the good clinical practices recommended by the Declaration of Helsinki and its 133 amendments. All participants gave their written informed consent.

134

#### 135 **Results**

## 136 Characteristics of study participants

A total of 293 students agreed to participate and answered the post-travel questionnaire. The M/F gender ratio was 0.31 with a median age of 21 years (ranging from 18 to 25 years). Most participants (80.9%) were students in their second year of medical studies and were taking part in a non-medical humanitarian mission. The remaining participants were in their 4<sup>th</sup> year of study and were assigned to different departments of medicine or surgery for clinical training (Supplementary Table S1).

143 The primary travel destinations for internships were in Africa (29.0%), South East Asia (27.7%), 144 South America (21.8%) and South Asia (18.8%). The top five primary destination countries were 145 Vietnam (24.2%), India (18.8%), Peru (16.4%), Tanzania (10.2%) and Madagascar (9.2%). The 146 mean travel duration was 41 days  $\pm$  11.1 days (ranging from 16 to 78 days). Accommodation 147 conditions were judged as being "very clean" by 14.7% (43/293) of students, "clean" by 44% 148 (129/293) and "very basic" by 41.3% (121/293). 42.7% and 79.9% of the participants reported 149 contact with animals and local children, respectively (Supplementary Table S2). During their stays, 150 94.2% also travelled as tourists in the country of primary destination and 50 (17.1%) visited other 151 countries. The top five additional destination countries were Laos (4.8%), Cambodia (4.4%), Bolivia 152 (3.4%), Thailand (1.7%) and Argentina (1.7%).

- 153 A total of 55/293 (18.8%) were vaccinated against influenza and only 2/293 (0.7%) against invasive
- 154 pneumococcal infections at inclusion. In addition, 260/293 (88.7%) and 273/293 (93.2%) student
- 155 were vaccinated against hepatitis A and hepatitis B before travel, respectively.
- Overall, 5.1% took antibiotics in the week before departure and 17.8% took doxycycline as achemoprophylaxis against malaria during their stay.
- 158 **Respiratory infections**

A total of 35.8% (105/293) students reported respiratory symptoms during travel. The median time between arrival at the travel destination and the onset of symptoms was 17 days [ranging from 1 to 58 days]. The most frequent respiratory symptoms were rhinitis (27.1%), sore throat (21.8%) and cough (20.1%), followed by fever (9.6%) and dyspnoea (6.5%) (Figure 1a). 7.9% of students declared persistence of symptoms on their return to France and 5.5% took antibiotics (ATB) for respiratory symptoms during travel.

165 A total of 275 (93.9%) students provided paired nasopharyngeal swabs. 52.4% of students acquired 166 at least one respiratory pathogen. 17.8% of (49/275) students acquired at least one respiratory virus 167 with human rhinovirus (14.6%) being the most frequent. Twenty-three students were still 168 symptomatic after returning to France. Of whom, 16 (69.6%) were positive for at least one 169 pathogens. Bacterial acquisition rates were higher (40.7%), with S. aureus (18.9%) being the most 170 frequent, followed by *H. influenza* (17.1%) (Table 1). A total of 6.2% students acquired a virus-171 bacteria combination and 11.6% a bacteria combination. When comparing the post- versus pre-172 travel prevalence of pathogens, a significant increase was observed for rhinovirus and S. 173 pneumoniae.

## 174 Gastro-intestinal infections

A proportion of 63.5% of students reported at least one gastro-intestinal symptom, all during travel and only one following return. The median time between arrival at the travel destination and the onset of symptoms was 13 days [ranging from 1 to 65 days]. The most frequent symptoms were diarrhoea (48.1%) and abdominal pain (46.4%), followed by nausea (26.3%) and constipation (19.8%) (Figure 1b). 8.9% reported persistent symptoms on return to France and 3.4% took an antibiotic for diarrhoea during their stay. 181 A total of 274 (93.5%) students provided paired rectal swabs. Three students reported 182 gastrointestinal symptoms at day 1 after departure. But all rectal samples pre-travel were negative 183 for gastrointestinal pathogens. 51.5% students acquired at least one gastrointestinal pathogen. Nine 184 students (3.3%) acquired at least one virus (adenovirus, astrovirus and norovirus). Bacterial 185 acquisition rates were higher (49.3%), notably for EAEC (40.9%) and EPEC (18.6%). Additionally, 186 2.6% of individuals acquired Shigella spp/EIEC, 1.5% Salmonella spp and 1.1% G. lamblia (Table 187 2). When comparing the post- versus pre-travel prevalence of pathogens, a significant increase was 188 observed for viruses overall, EAEC, EPEC, Salmonella spp and Shigella spp/EIEC.

## 189 Vaginal infections

190 Of the 224 female students, eight (3.6%) reported vaginal symptoms such as leucorrhoea, during 191 their stay, and two took antibiotics for this purpose. The median time between arrival at the travel 192 destination and the onset of symptoms was 26 days [ranging from 6 to 59 days]. Three students 193 (1.3%) were still symptomatic on returning to France.

194 209 (93.3%) of the participants provided paired vaginal samples. 15.8% of the participants acquired 195 at least one vaginal microorganism with the highest acquisition rate for *A. vaginae* (13.9%) and *G.* 196 *vaginalis* (12.9%) (Table 3). Nine students (4.0%) had molecular criteria for bacterial vaginosis on 197 return. When comparing the post- versus pre-travel prevalence of microorganisms, a significant 198 increase was observed for *G. vaginalis* (CT<18), indicative of vaginosis.

## 199 Risk factors for symptoms of infections

Being female, primarily travelling to Vietnam, and living in basic accommodation conditions were independent risk factors for reporting respiratory symptoms. Students suffering respiratory symptoms were 3 times more likely to acquire *S. pneumoniae* during travel. Travelling primarily to north India and Senegal were independent risk factors for reporting diarrhoea (Table 4). 204

## 205 Discussion

206 Most students in our study travelled to low-income tropical countries for about two months and 207 participated in humanitarian missions in relatively basic conditions of housing and in close contact 208 with local children and animals. Two-thirds of students reported gastrointestinal symptoms (notably 209 diarrhoea, abdominal pain and nausea-vomiting which are suggestive of gastroenteritis) and one-210 third of students reported respiratory symptoms (notably rhinitis, a sore throat and a cough which 211 are suggestive of an upper respiratory tract infection). These symptoms appeared within 2 weeks of 212 their arrival at destination of their internship. Moreover, we observed a low proportion of travel-213 associated vaginal symptoms. Overall, symptoms were relatively mild with fewer than 5% students 214 requiring antibiotics and most symptoms resolved before students came back to France. This result 215 is consistent with other studies realized on medical students abroad, confirming the fact that despite 216 reinforced pre-travel counselling, travel-associated respiratory infections and travellers' diarrhoea 217 were very frequent among medical students who were fully aware of the ways to prevent these 218 illnesses [5, 8-10].

We found a significant acquisition of human rhinovirus and *S. pneumoniae* as reported previously among international travel as well as among Hajj pilgrims [1, 11, 12]. We also observed a high acquisition rate of EAEC (40.9%) and EPEC (18.6%) among health students, as documented in other studies realized in different populations of domestic and international travellers [13-18].

Our results showed that respiratory symptoms were significantly more frequent in female. We have no explanation for this observation. Interestingly, the travel destination was distinctly associated with symptoms. Travel to India and Senegal was a risk factor for diarrhoea while travel to Vietnam was a risk factor for respiratory symptoms. Our results are discordant with those of a previous study 227 on 649 international travellers showing that respiratory infections (sore throat or cough) were 228 significantly increased in travellers returning from the non-tropical regions (7.6%) than those from 229 tropical regions, including Vietnam (2.0%) [19]. Differences by travel destination are also known to 230 be relative to the incidence of travellers' diarrhoea. This result was in line with most other studies 231 that have also found that travelling to the Indian subcontinent was a highest relative risk for 232 diarrhoea, followed by African regions [16, 20-23].

In addition, the observed correlation between respiratory symptoms and very basic accommodation conditions suggests that precarious housing conditions may encourage respiratory infections. A significant association between the acquisition of *S. pneumoniae* and respiratory symptoms was also observed in cohorts of Hajj pilgrims [24].

We observed no significant association between E. coli acquisition and diarrhoea in our study, in 237 238 contrast to other studies where EAEC or EPEC have been reported to be more frequent in travellers 239 with diarrhoea returning from several geographical areas [14, 16, 25]. This may be explained by the 240 onset of gastro-intestinal symptoms occurring early during the trip, while sampling was performed 241 on return, several weeks later. Furthermore, asymptomatic carriage of potential pathogens was also 242 observed in participants. In a study by Adachi et al., EAEC was detected in the stools of 26% of 243 patients with traveller's diarrhoea returning from Mexico, Jamaica, or India [26]. On the other hand, 244 a recent case-controlled study conducted on German and Dutch travellers showed that EAEC 245 detection was not significantly different in diarrheal persons and asymptomatic controls. However, 246 the prevalence of this bacterium among participants suffering from diarrhoea during international 247 travel was high (40.0%) [14]. Such results are in line with ours.

Our study has a few limitations. First, this study was monocentric and conducted on a very specific
population of travellers which impairs generalisation of our findings. Also, qPCR does not

differentiate between dead and viable microorganisms. Finally, sampling was realized during the week preceding departure and during the week following return, samples at onset of symptoms were not available. Among students who reported clinical symptoms early after their arrival abroad, we are not sure whether these students were infected before or after departure, since incubation times of diseases are very different and may vary (Supplementary table S3). Likewise, post-travel samples were collected at a significant time after the onset of symptoms during travel and responsible pathogens may have been partly cleared.

This study makes it possible to identify the main infectious diseases linked to travel in a group of French medical students undertaking an internship abroad and risk factors upon which to base targeting students for reinforced pre-travel advice. Sampling at the time of the onset of symptoms should be carried out in future studies to better understand the relationship between the carriage of pathogens and symptoms.

262

## 263 Transparency declaration

264 The authors declare that there are no conflicts of interest.

265

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270

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Table 1: Prevalence of respiratory pathogens

	Pre-t	ravel	Post-t	ravel	Acquis	ition	
Variables	n = 275 <sup>1</sup>	%	n = 279 <sup>2</sup>	%	n =275 <sup>3</sup>	%	p*
Viruses							
Adenovirus	1	0.4	0	0	0	0	NA
Coronaviruses	1	0.4	6	2.2	6	2.2	0.06
Coronavirus HKU1	0	0	3	1.1	3	1.1	0.08
Coronavirus 229E	0	0	2	0.7	2	0.7	0.16
Coronavirus NL63	1	0.4	1	0.4	1	0.4	1.0
Coronavirus OC43	0	0	0	0	0	0	NA
Influenza A	0	0	1	0.4	1	0.4	0.32
Influenza B	0	0	2	0.7	2	0.7	0.16
Metapneumovirus	0	0	0	0	0	0	NA
Parainfluenza virus	0	0	0	0	0	0	NA
Respiratory syncytial virus	1	0.4	2	0.7	2	0.7	0.56
Rhinovirus	25	9.1	44	15.8	40	14.6	0.015
At least one virus	26	9.5	54	19.6	49	17.8	0.001
Bacteria							
Haemophilus influenzae	122	44.4	119	42.7	47	17.1	0.76
Klebsiella pneumoniae	23	8.4	28	10.0	19	6.9	0.38
Staphylococcus aureus	63	22.9	83	29.8	52	18.9	0.052
Streptococcus pneumoniae	5	1.8	15	5.4	15	5.5	0.025
Streptococcus pyogenes	3	1.1	8	2.9	7	2.6	0.096
At least one bacteria	161	58.6	180	64.5	112	40.7	0.11
Virus and bacteria combination	13	4.7	29	10.4	17	6.2	0.01
H. influenzae-rhinovirus	9	3.3	13	4.7	4	1.5	0.37
K. pneumoniae-rhinovirus	1	0.4	1	0.4	0	0	NA

S. pneumoniae-rhinovirus	0	0	1	0.4	1	0.4	0.32
S. aureus-rhinovirus	6	2.2	12	4.3	8	2.9	0.16
Bacteria combination of two bacteria	61	22.2	87	31.2	32	11.6	<0.000
H. influenza - K. pneumoniae	12	4.4	15	5.4	3	1.1	0.49
H. influenza - S. pneumoniae	4	1.5	11	3.9	5	1.8	0.07
H. influenza - S. aureus	36	13.1	40	14.3	12	4.4	0.55
H. influenzae-S. pyogenes	3	1.1	5	1.8	1	0.4	0.41
K .pneumoniae-S. pneumoniae	0	0	3	1.1	3	1.1	0.08
K. pneumoniae-S. aureus	4	1.5	7	2.5	2	0.7	0.26
K. pneumoniae-S. pyogenes	0	0	0	0	0	0	NA
S. pneumoniae-S. aureus	1	0.4	6	2.1	6	2.2	0.06
S. pneumoniae-S. pyogenes	0	0	0	0	0	0	NA
S. aureus-S. pyogenes	1	0.4	0	0	0	0	NA
At least one pathogen	174	63.3	205	73.5	144	52.4	0.008

\* p-value: pre- versus post-travel, McNemar's test

1, 2: 275 and 279 students provided pre- and post-travel nasopharyngeal swabs, respectively

3: Acquisition of respiratory pathogens was calculated in 275 students who provided both pre- and post-travel nasopharyngeal samples

Dethe anna	Pre-tra	vel	Post-Tra	avel	Acquisi	··· •	
Pathogens	$N = 282^{1}$	%	$N = 283^2$	%	$N = 274^3$	%	p*
Viruses							
Adenovirus	0	0	3	1.1	3	1.1	0.08
Astrovirus	0	0	3	1.1	3	1.1	0.08
Norovirus	0	0	3	1.1	3	1.1	0.08
Rotavirus	0	0	0	0	0	0	NA
At least one virus	0	0	9	3.2	9	3.3	0.003
Bacteria							
Campylobacter jejuni	2	0.7	5	1.8	5	1.8	0.26
Enteroaggregative Escherichia coli	10	3.6	119	42.1	112	40.9	< 0.000
Enterohemorrhagic E. coli	3	1.1	4	1.4	2	0.7	0.65
Enteropathogenic E. coli	25	8.9	66	23.3	51	18.6	< 0.000
Salmonella spp	0	0	4	1.4	4	1.5	0.05
Shigella spp/EIEC	0	0	7	2.5	7	2.6	0.008
Tropheryma whipplei	1	0.4	1	0.4	0	0	NA
At least one bacteria	38	13.5	147	51.9	135	49.3	< 0.000
Parasites							
Cryptosporidium parvum/hominis	0	0	0	0	0	0	NA
Entamoeba histolytica	0	0	0	0	0	0	NA
Giardia lamblia	1	0.4	3	1.1	3	1.1	0.32
At least one gastrointestinal pathogen	39	13.8	153	54.1	143	52.2	< 0.000

\* p-value: pre- versus post- travel, McNemar's test

1, 2: 282 and 283 students provided pre- and post-travel rectal swabs, respectively

3: Acquisition of gastrointestinal pathogens was calculated in 274 students who provided both preand post-travel rectal samples

Missossiana	Pre-travel	Post-travel	Acquisition	
Microorganisms	N =212 <sup>1</sup> (%)	N= 221 <sup>2</sup> (%)	$N = 209^3 (\%)$	p *
Bacteria				
Atopobium vaginae	42 (19.8)	42 (19.0)	29 (13.9)	0.896
Atopobium vaginae (CT<21)	1 (0.5)	1 (0.5)	1 (0.5)	1.0
Chlamydia trachomatis	1 (0.5)	1 (0.5)	1 (0.5)	1.0
Gardnerella vaginalis	59 (27.8)	55 (24.9)	27 (12.9)	0.439
Gardnerella vaginalis (CT<18)	2 (0.9)	9 (4.1)	9 (4.3)	0.035
Mycoplasma genitalium	0	2 (0.9)	2 (0.96)	0.16
Mycoplasma hominis	4 (1.9)	3 (1.4)	2 (0.96)	0.65
Neisseria gonorrhoeae	0	0	0	NA
Parasites				
Trichomonas vaginalis	0	0	0	NA
At least one vaginal pathogen	82 (38.7)	78 (35.3)	33 (15.8)	0.42

Table 3: Prevalence of vaginal microorganisms

\* p-value: pre- versus post-travel, McNemar's test

1, 2: 212 and 221 students provided pre- and post-travel vaginal swabs, respectively

3: Acquisition of vaginal pathogens was calculated in 209 students who provided both pre- and post-travel vaginal samples

Table 4: Risk factors for respiratory symptoms	and diarrhoea during travel
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	Risk fa	ctors for re	espiratory symp	otoms du		Risk facto	r for diarrhoea	for diarrhoea during the travel			
		Univari	ate analyse		Multivariate analyse		Univariate analyse			Multivariate analyse	
	No	Yes	RR		aRR	No	Yes	RR		aRR	
	n (%)	n (%)	[95%CI]	р	р [95%CI], р г	n (%)	n (%)	[95%CI]	р	[95%CI], p	
Gender											
	136	88		0.02		121	103				
Female	(72.3)	(83.8)	1.50		1.55 0.02 [1.08-2.22], p= 0.02	(54.0)	(46.0)	0.77	0.10		
Male	52	17	[1.06-2.11]			31	38	[0.51-1.15]	0.19	-	
Male	(27.7)	(16.2)			P 0.02	(44.9)	(55.1)				
Age	20.7±1.	20.7±1.	t= 0.38	0.70		20.7 ±	20.6±	t-0.70	0.48		
(mean ± SD)	1	1	0.70	0.70		1.1	1.1	t=0.70			
Medical history											
Vaccination	83	22	1.15								
vaccination against influenza	(34.9)	(40.0)	[0.78-1.70] 0.48	0.48		NA	NA	NA	NA		
Primary travel de	estination										
South Foot Acia	71	34	1.26	0.10		104	37	0.93	0.61		
South East Asia	(33.5)	(42.0)	[0.89-1.77]	0.18	-	(49.1)	(45.7)	[0.71-1.22]	0.61		

Vietnam	73 (32.9)	32 (45.1)	1.40 [0.97-2.00]	0.07	1.57 [1.07-2.31], p= 0.02	111 (50.0)	30 (42.3)	0.85 [0.62-1.14]	0.26	
South Asia	86 (36.1)	19 (34.6)	0.96 [0.65-1.40]	0.82		107 (45.0)	34 (61.8)	1.38 [1.07-1.77]	0.02	-
South India	98 (37.4)	7 (22.6)	0.66 [0.41-1.05]	0.08	-	125 (47.7)	16 (51.6)	1.08 [0.75-1.56]	0.68	
North India	93 (34.6)	12 (50.0)	1.53 [0.84-2.79]	0.16	-	123 (45.7)	18 (75.0)	1.64 [1.26-2.14]	<0.01	2.57 [1.13-5.85] p= 0.025
Africa	77 (37.0)	28 (32.9)	0.89 [0.65-1.24]	0.51		105 (50.5)	36 (42.4)	0.84 [0.63-1.11]	0.21	
Tanzania	99 (37.6)	6 (20.0)	0.61 [0.38-0.97]	0.04	-	137 (52.1)	4 (13.3)	0.26 [0.10-0.64]	<0.000	0.37 [0.22-0.60] p <0.000
Madagascar	91 (34.2)	14 (51.9)	1.63 [0.92-2.90]	0.09	-	129 (48.5)	12 (44.4)	0.92 [0.59-1.42]	0.69	
Senegal	98 (35.3)	7 (46.7)	1.37 [0.82-2.83]	0.40		129 (46.4)	12 (80.0)	1.72 [1.30-2.29]]	0.01	3.31 [1.05-10.45] p= 0.04
South America	84	21	0.90	0.569		108	33	1.09	0.53	

	(36.7)	(32.8)	[0.63-1.29]			(47.2)	(51.6)	[0.83-1.44]				
D	87	18	1.06	0.70		114	27	1.21	0.00			
Peru	(35.5)	(37.5)	[0.70-1.58]	0.79		(46.5)	(56.3)	[0.91-1.60]	0.22			
Condition during	travel											
Contact with	18	87	1.20			27	114	1.06				
children	(30.5)	(37.2)	[0.83-1.73]	0.33		(45.8)	(48.7)	[0.78-1.45]	0.69			
Contact with	52	53	1.37	0.046		77	64	1.12				
animals	(31.0)	(42.4)	[1.01-1.86]			0.046	-	(45.8)	(51.2)	[0.88-1.42]	0.36	
Type of accommod	lation											
<b>X</b> 7 1	34	9	c	c	rfr	22	21	c				
Very clean	(79.1)	(20.9)	rfr	rfr		(51.2)	(48.8)	rfr	Rfr			
	84	45	1.49	0.0 <b>-</b>	0.0 <b>7</b>	0.07	0.07		73	56	0.86	0.54
Clean	(65.1)	(34.9)	[0.96-2.29]	0.07	- (56.6) (43.4)	[0.53-1.40]	0.54					
	70	51	1.81		1.43	57	61	1 1 2				
Basic				<0.01	[1.02-1.99],		64	1.13	0.65			
	(57.9)	(42.2)	[1.16-2.82]		p= 0.04	(47.1)	(52.9)	[0.68-1.86]				
Mission during tr	avel											
Humanitarian	94	11	0.60			115	26	1.02				
mission in an orphanage	(39.2)	(20.8)	[0.41-0.87]	<0.01	-	(47.9)	(49.1)	[0.76-1.39]	0.88			

School renovation	49 (31.4)	56 (40.9)	1.29 [0.96-1.75]	0.09 -	79 (50.6)	62 (45.3)	0.89	0.36
Supply of medical equipment and	(31.1) 86 (35.0)	(101 <i>)</i> ) 19 (40.4)	1.16	0.48	(47.6)	24 (51.1)	1.07	0.66
health advice Internships in hospital	(36.3) (36.3)	(10.1) 19 (33.9)	0.94	0.74	(17.3) 112 (47.3)	(51.1) 29 (51.8)	[0.77 1.10] 1.10 [0.82-1.46]	0.54
Internships in surgery department	97 (35.3)	8 (44.4)	1.29 [0.82-1.12]	0.45	130 (47.3)	11 (61.1)	1.29 [0.88-1.91]	0.26
Duration of travel (mean ± SD)	40.5 ±10.7	41.8 ±11.9	t=-0.95	0.34	39.6± 10.7	42.6± 11.4	t=-2.35	0.02 -
Acquisition of resp	piratory p	athogens <sup>2</sup>	75					
Rhinovirus	86 (36.6)	11 (27.5)	0.78 [0.51-1.19]	0.25	NA	NA	NA	NA
At least one respiratory virus	84 (37.2)	13 (26.5)	0.75 [0.50-1.10]	0.14 -	NA	NA	NA	NA
Haemophilus influenzae	79 (34.7)	18 (38.3)	1.10 [0.73-1.67]	0.64	NA	NA	NA	NA
Klebsiella pneumoniae	89 (34.8)	8 (42.1)	1.22 [0.65-2.29]	0.53	NA	NA	NA	NA

Staphylococcus	78	19	1.04	0.83		NI A	NI A	N A	
aureus	(35.0)	(36.5)	[0.70-1.55]	0.85		NA	NA	NA	NA
Streptococcus	87	10	2.70		2.79				
pneumoniae	(33.5)	(66.7)	[1.10-6.62]	0.03	[1.13-6.88], p= 0.03	NA	NA	NA	NA
At least one	56	41	1.06	0.70			NT A	NT A	
respiratory bacteria	(34.4)	(36.6)	[0.78-1.45]	0.70	0.70	NA	NA	NA	NA
Acquisition of dige	estive path	nogens <sup>274</sup>							
Enteroaggregativ	NA NA	NA	NA		78	58	1.08	0.55	
e Escherichia coli	INA	MA	NA	NA		(48.2)	(51.8)	[0.85-1.37]	0.35
Enteropathogenic	NA	NA	NA	NA		110	26	1.03	0.83
E. coli	INA	INA	INA	INA		(49.3)	(51.0)	[0.77-1.40]	0.85
At least one	NT A		NT A	NT A		68	68	1.04	
gastrointestinal bacteria	NA	NA	NA	NA		(48.9)	(50.4)	[0.74-1.47]	0.81
At least one	NT A	NT A	NT A	NT A		64	72	1.04	0.91
gastrointestinal bathogen	NA	NA	NA	NA	NA	(48.9)	(50.4)	[0.74-1.47]	0.81

NA: not applicable

RR: relative risk, aRR: adjusted relative risk, CI: confidence interval, p: p value, rfr: reference

<sup>¥</sup>Number of individuals for whom data were available

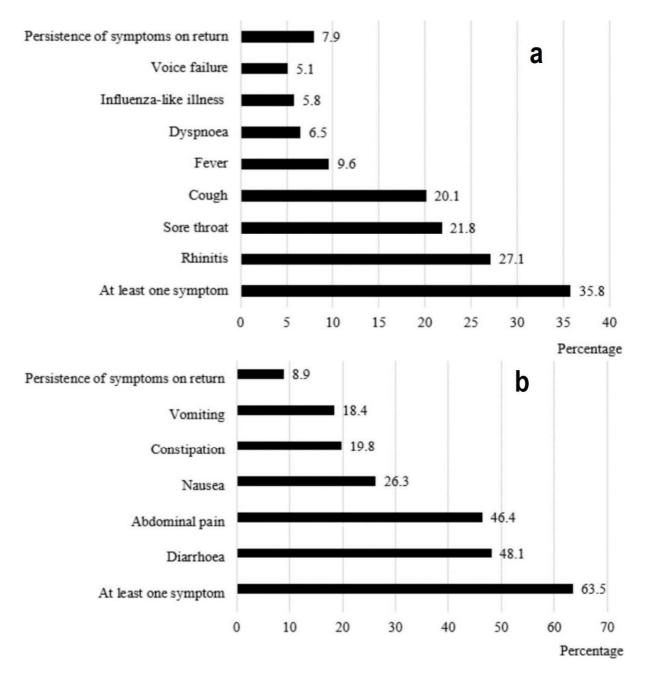


Figure 1: Prevalence of respiratory (a) and gastrointestinal (b) symptoms during travel