

**Nature strikes back!**





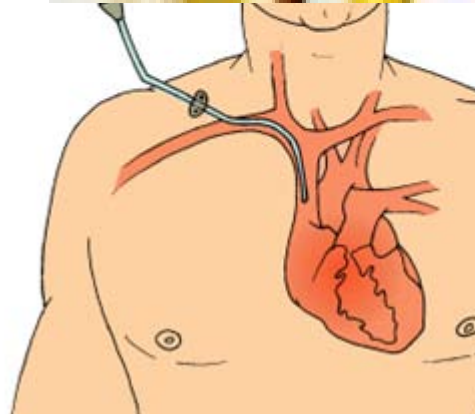
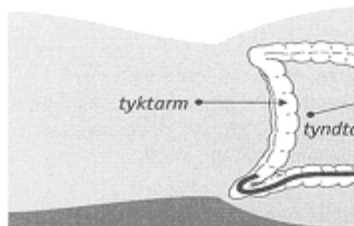


<b>Bakterie</b>	<b>Antal stammar (%) i Norge</b>	<b>Antal stammar (%) i Malmö</b>
<i>E. coli</i>	131 (31,0)	231 (26,4)
<i>S. pneumoniae</i>	69 (16,4)	46 (5,3)
<i>S. aureus</i>	59 (14,0)	110 (12,6)
<i>Klebsiella spp.</i>	29 (6,9)	54 (6,2)
<i>Enterokock spp.</i>	20 (4,7)	61 (7,0)
<i>S. pyogenes</i>	18 (4,3)	50 (5,7)
<i>Viridans streptokocker</i>	17 (4,0)	69 (7,9)
<i>P. mirabilis</i>	15 (3,6)	17 (1,9)
<i>P. aeruginosa</i>	9 (2,1)	21 (2,4)
<i>Neisseria spp.</i>	8 (1,9)	1
<i>S. agalactiae</i>	8 (1,9)	0
<i>Enterobacter spp.</i>	7 (1,7)	47 (5,4)
Övrigt	32 (7,6)	165 (18,9)



Numerous studies have shown that the acquisition and infection of *C. difficile* is often associated with antibiotic use.

Antibiotics for *C. difficile*



## ORIGINAL ARTICLE

Extended-spectrum beta-lactamase-producing *Escherichia coli* in patients withEur J Clin Microbiol Infect Dis  
DOI 10.1007/s10096-011-1202-5

## ARTICLE

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and in aH. Strömdahl  
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Scandinavian Journal of Infectious Diseases, 2012; Early Online, 1–5

## Abstract

The identification of patients at risk of colonization with extended-spectrum beta-lactamase (ESBL)-producing *Escherichia coli* (ESBL-*E. coli*) is important to investigate the occurrence of ESBL-producing *E. coli* in patients with diarrhoea. The aim of this study was to identify ESBL-producing *E. coli* in patients with diarrhoea and to determine the prevalence of ESBL-producing *E. coli* in patients with diarrhoea. The study was conducted in a tertiary care hospital in Sweden in 2007–2010. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea was 100%. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea was 100%. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea was 100%.Received: 24 November 2011  
© Springer-Verlag**Abstract** The prevalence of ESBL-producing *E. coli* in a group of patients with diarrhoea was 100%. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea was 100%. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea was 100%. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea was 100%. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea was 100%.

## Introduction

The occurrence of ESBL-producing *E. coli* has become a worldwide phenomenon, posing a great threat to public health. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea has increased in recent years [1]. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea has increased in recent years [1]. The prevalence of ESBL-producing *E. coli* in patients with diarrhoea has increased in recent years [1].Antibiotic use for colonization of Enterobacteriaceae with severe illness, prolonged hospital attendance, haemodialysis and recent surgery practices increase the prevalence of ESBL-producing *E. coli*.

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## ORIGINAL ARTICLE

Duration of colonization with extended-spectrum beta-lactamase-producing *Escherichia coli* in patients with travellers' diarrhoeaJOHAN THAM<sup>1</sup>, MATS WALDER<sup>2</sup>, EVA MELANDER<sup>2,3</sup> & INGA ODENHOLT<sup>1</sup>From the <sup>1</sup>Infectious Diseases Unit, Department of Clinical Sciences, Lund University, Malmö, <sup>2</sup>Medical Microbiology, Department of Laboratory Medicine, Lund University, Malmö, and <sup>3</sup>Department of Infection Control, Laboratory Medicine, Skåne County, Sweden

## Abstract

**Background:** Resistant Enterobacteriaceae have become a worldwide epidemic during the last decade and are a great threat to health care worldwide. International travel is a major risk factor for becoming colonized with extended-spectrum beta-lactamase (ESBL)-producing bacteria. Data on the persistence of colonization with ESBL-producing bacteria in the faecal flora are limited. **Methods:** A prospective cohort study was performed between October 2007 and October 2010. Fifty patients with faecal carriage of ESBL-producing *Escherichia coli* from a previous study of patients with travellers' diarrhoea were included. **Results:** Forty-one of the patients had a complete follow-up. Ten of these patients (24%) carried ESBL-producing *E. coli* at the first follow-up point (3–8 months), of whom 4 had a new ESBL strain. At the 3-year follow-up, 10 patients carried ESBL (10%), of whom 1 had 2 new ESBL strains. **Conclusions:** The long duration of ESBL carriage is worrisome. These carriers may be an important source of the spread of ESBLs in the population and this has implications for public health.

# Extended-spectrum beta-lactamase-producing *Escherichia coli* in patients with travellers' diarrhoea

242 patients with travellers' diarrhoea

## ESBL-screen

Medium selective for cephalosporin resistance  
(ChromID ESBL, BioMerieux)



## Synergy testing

With disks containing ceftazidime  
and cefotaxime  
amoxicillin/clavulanic acid



# Extended-spectrum beta-lactamase-producing *Escherichia coli* in patients with travellers' diarrhoea





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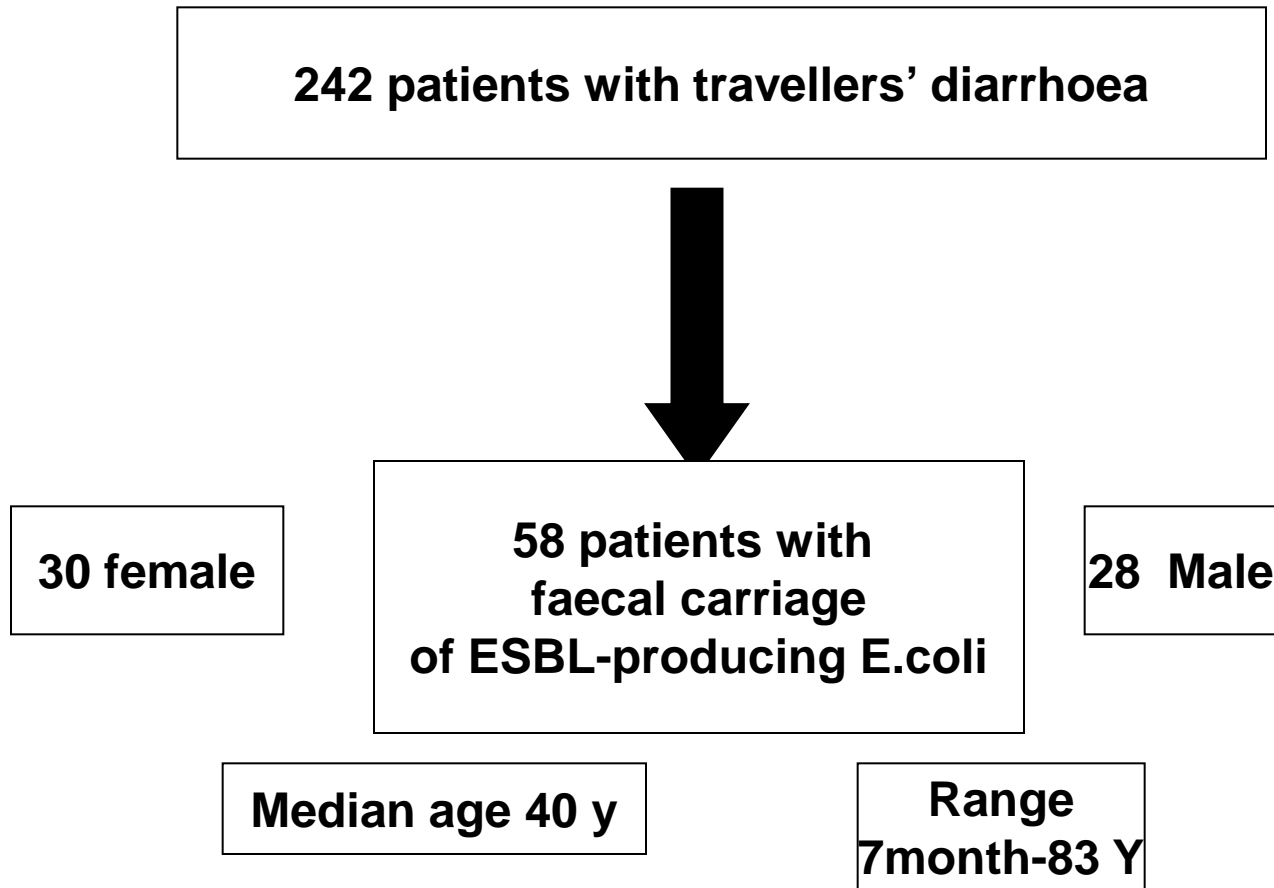
## Synergy testing

with disks containing ceftazidime and cefotaxime and amoxicillin/clavulanic acid

58 patients  
ESBL-producing  
*E.coli*



# Extended-spectrum beta-lactamase-producing *Escherichia coli* in patients with travellers' diarrhoea



# Regions and countries involved in the study

Table II. Regions and countries involved in the study: Europe (Bosnia, Bulgaria, Denmark, UK, France, Germany, Greece, Hungary, Ireland, Italy, Kosovo, Romania, Spain, Turkey and Ukraine), Middle East (Kurdistan, Lebanon, Morocco, Iraq, Oman, Saudi Arabia, Syria and Tunisia), Africa (Gambia, Ghana, Guinea, Kenya, Tanzania and unspecified), Southeast Asia (Afghanistan, Australia, Bangladesh, Cambodia, China, Pakistan, Papua New Guinea, Philippines, Singapore and Tahiti), America (Argentina, Bolivia, Caribbean, Chile, Mexico and unspecified parts of America).

Region	ESBL-positive ( <i>n</i> )	ESBL-negative ( <i>n</i> )	Total ( <i>n</i> )	Proportion positive	95% CI	<i>p</i> -Value compared to Europe
World	58	184	242	(58/242)=0.24	0.19–0.30	
World excl. Europe and unspecified	50	88	138	(50/138)=0.36	0.29–0.45	<0.0001
Europe excl. Sweden	2	61	63	(2/63)=0.03	0.004–0.11	
Egypt	19	19	38	(19/38)=0.50	0.33–0.67	<0.0001
Thailand	8	28	36	(8/36)=0.22	0.10–0.39	0.0042
India	11	3	14	(11/14)=0.79	0.49–0.95	<0.0001
Middle East	4	6	10	(4/10)=0.40	0.12–0.74	0.0025
Southeast Asia incl. Australia	5	8	13	(5/13)=0.38	0.14–0.68	0.0012
Africa excl. Egypt	2	15	17	(2/17)=0.12	0.015–0.36	0.1965 (NS)
America incl. West Indies	1	9	10	(1/10)=0.10	0.0025–0.44	0.3615 (NS)
Unspecified	6	35	41	(6/42)=0.15	0.06–0.29	0.0550 (NS)

ESBL, extended-spectrum beta-lactamase; CI, confidence interval; NS, not significant.



## Pathogens found in the stool samples. All of these isolates were ESBL-negative

Pathogen	ESBL-negative	ESBL-positive (E. coli)	Total
<i>Campylobacter jejuni/coli</i>	31	3	34
<i>Salmonella enteritidis</i>	2	0	2
<i>Salmonella</i> group 04	2	1	3
<i>Salmonella</i> group 07	1	0	1
<i>Salmonella</i> group 08	1	0	1
<i>Salmonella senftenberg</i>	0	1	1
<i>Shigella flexneri</i>	1	0	1
<i>Shigella sonnei</i>	2	1	3
<i>Shigella boydii</i>	0	1	1
All pathogens (SSYC)	40	7	47

ESBL, extended-spectrum beta-lactamase.



# Antibiotic resistance

Antibiotic	Clinical E. coli isolates with ESBLs (%)	Study E. coli isolates with ESBLs (%)
Tobramycin	42	54
Ciprofloxacin	62	68
Piperacillin–tazobactam	22	8
Mecillinam	6	0
Trimethoprim	80	91
Trimethoprim–sulfamethoxazole	87	75
Nitrofurantoin	8	5



# Enzyme typing and rep PCR results

- 90% CTX-M group
- CTX-M 1 68% ( The only group found in India)
- CTX-M 9 24%
- The others were TEM or SHV and some isolates both TEM and SHV



# CTX-M typing and rep PCR results

- rep PCR fingerprint pattern:
  - the strains from the same geographical region displayed no genetic similarity
  - were also different from Swedish E. coli isolates studied earlier



# Limitations

- Our patients were not cultured for ESBL-producing bacteria before going abroad
- Lack of other epidemiologic information
- Low number of patients with travellers' diarrhoea from some parts of the world





**Summary Objectives:** Extended-spectrum  $\beta$ -lactamase (ESBL)-producing *Escherichia coli* have emerged as significant causes of community-onset disease. We sought to identify risk factors for acquiring community-onset ESBL-producing *E. coli*.

**Methods:** Prospective, population-based surveillance for ESBL-producing *E. coli* was performed in the Calgary Health Region (population 1.2 million), Canada during a two-year period.

**Results:** 247 patients were identified; 177 (72%; 7.6 per 100,000/year) were community acquired, and 70 (28%; 3.0 per 100,000/year) were healthcare associated. The acquisition risk increased with advancing age. Females were at higher risk as compared to males [relative risk (RR) 4.3; 95% confidence interval (CI), 3.1–6.1] as were urban as compared to rural residents (RR 2.2; 95% CI, 1.4–3.6). A number of co-morbidities increased risk (RR; 95% CI) including requirement for hemodialysis (56.3; 15.1–147.4), urinary incontinence (21.7; 15.0–30.9), cancer (11.1; 7.0–17.0), heart disease (6.5; 4.3–9.7), and diabetes (4.4; 2.6–7.1). Overseas travel overall increased the risk (5.7; 4.1–7.8) and was highest in travelers to India (145.6; 77.7–252.1), the Middle East (18.1; 8.1–35.2), and Africa (7.7; 2.8–17.2).

Accepted 24 September 2008  
Available online 5 November 2008



# Foreign Travel Is a Major Risk Factor for Colonization with *Escherichia coli* Producing CTX-M-Type Extended-Spectrum $\beta$ -Lactamases: a Prospective Study with Swedish Volunteers<sup>∇</sup>

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Received 15 February 2010/Returned for modification 17 April 2010/Accepted 8 June 2010

Foreign travel has been suggested to be a risk factor for the acquisition of extended-spectrum beta-lactamase (ESBL)-producing *Enterobacteriaceae*. To our knowledge, this has not previously been demonstrated in a prospective study. Healthy volunteers traveling outside Northern Europe were enrolled. Rectal swabs and data on potential travel-associated risk factors were collected before and after traveling. A total of 105 volunteers were enrolled. Four of them did not complete the study, and one participant carried ESBL-producing *Escherichia coli* before travel. Twenty-four of 100 participants with negative pretravel samples were colonized with ESBL-producing *Escherichia coli* after the trip. All strains produced CTX-M enzymes, mostly CTX-M-15, and some coproduced TEM or SHV enzymes. Coresistance to several antibiotic subclasses was common. Travel to India was associated with the highest risk for the acquisition of ESBLs (88%;  $n = 7$ ). Gastroenteritis during the trip was an additional risk factor ( $P = 0.003$ ). Five of 21 volunteers who completed the follow-up after 6 months had persistent colonization with ESBLs. This is the first prospective study demonstrating that international travel is a major risk factor for colonization with ESBL-producing *Enterobacteriaceae*. Considering the high acquisition rate of 24%, it is obvious that global efforts are needed to meet the emergence and spread of CTX-M enzymes and other antimicrobial resistances.

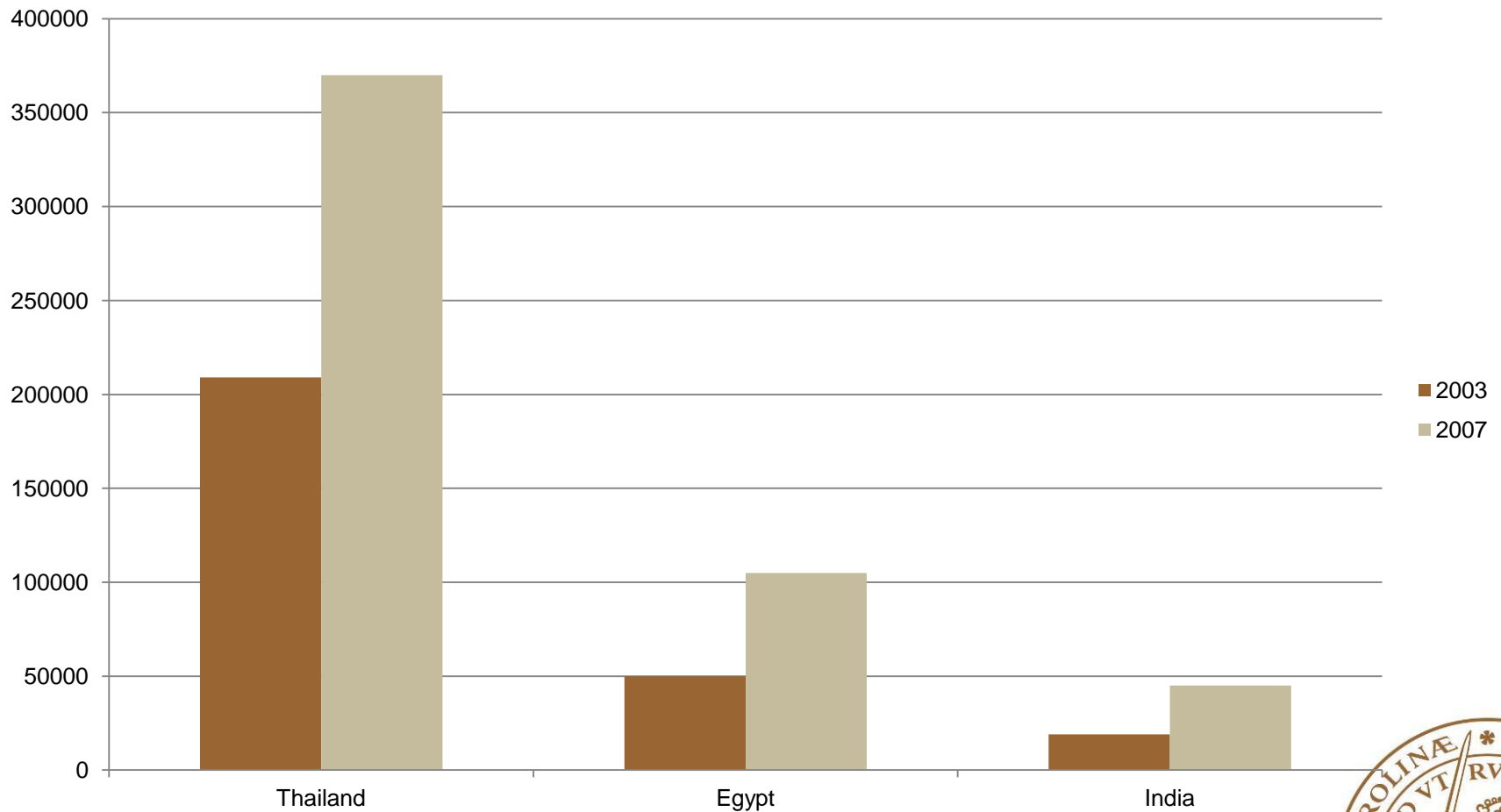


## **Tängden et al** (ANTIMICROBIAL AGENTS AND CHEMOTHERAPY, Sept. 2010, p. 3564–3568)

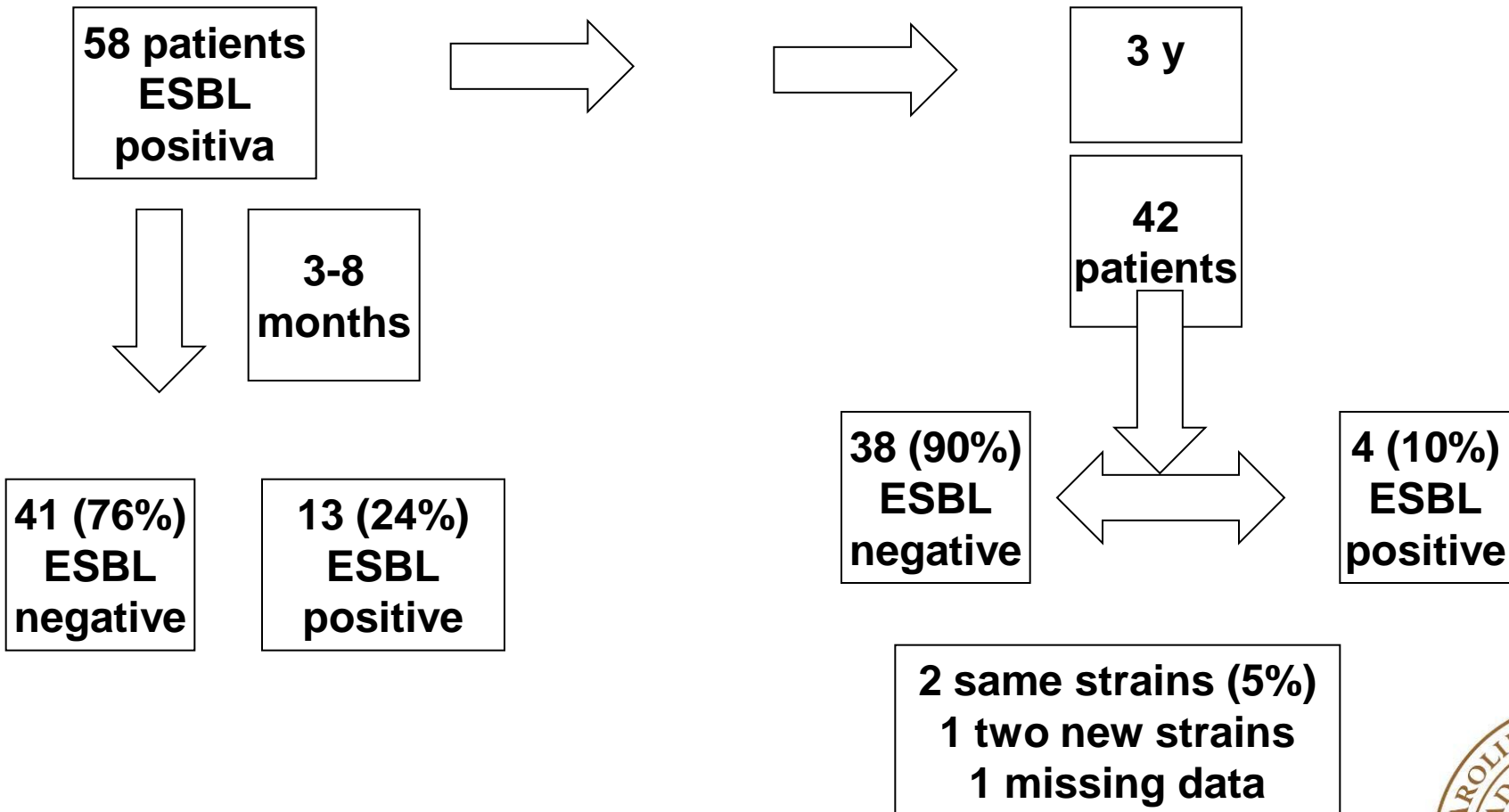
- **Twenty-four of 100 participants with negative pretravel samples were colonized with ESBL-producing *Escherichia coli* after the trip**
- **All strains produced CTX-M enzymes, mostly CTX-M-15, and some coproduced TEM or SHV enzymes**
- **Coresistance to several antibiotic subclasses was common**
- **Gastroenteritis during the trip was an additional risk factor**
- **Five of 21 (24%) volunteers who completed the follow-up after 6 months had persistent colonization with ESBLs**



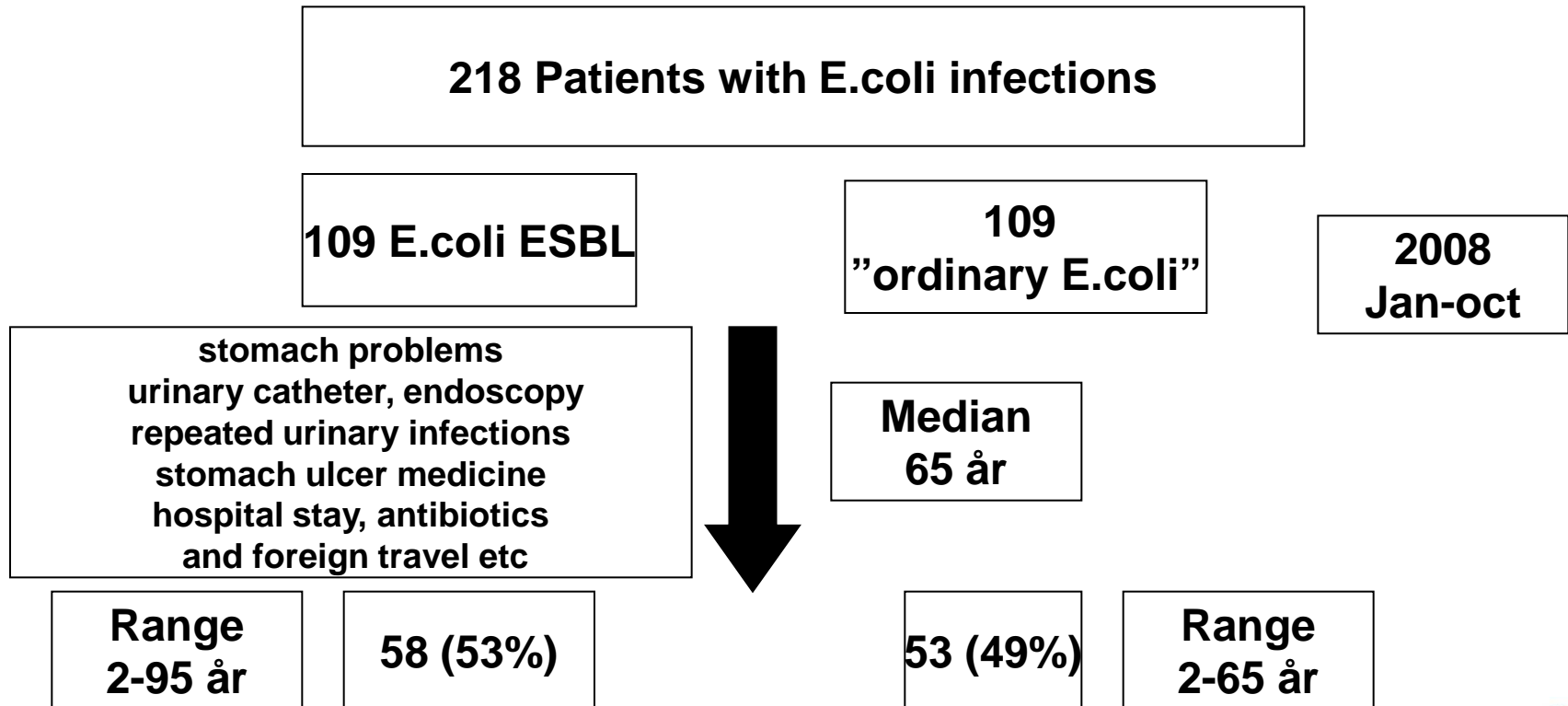
# Foreign travel is increasing



# Duration of colonization with Extended-spectrum beta-lactamase producing Escherichia coli in patients with travellers' diarrhoea



# Risk factors for infections with Extended-spectrum beta-lactamase producing Escherichia coli



# Risk factors for infections with Extended-spectrum beta-lactamase producing *Escherichia coli*



**Hospital stay (n=8)  
> 1 month  
P<0,01**

**Foreign travel (n=14)  
Asia, Middle East  
P=0,02**





## ORIGINAL ARTICLES

## Colonization of Returning Travelers With CTX-M-Producing *Escherichia coli*

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and Johann D.D. Pitout, MD<sup>\*#¶</sup>

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DOI: 10.1111/j.1708-8305.2011.00548.x

**Background.** We previously identified foreign travel as a risk factor for acquiring infections due to CTX-M (active on cefotaxime first isolated in Munich) producing *Escherichia coli*. The objective of this study was to assess the prevalence of extended-spectrum  $\beta$ -lactamase (ESBL)-producing *E coli* among stool samples submitted from travelers as compared to non-travelers (a non-traveler had not been outside of Canada for at least 6 months before submitting a stool specimen).

**Methods.** Once a travel case was identified, the next stool from a non-traveler (not been outside of Canada for at least 6 months) was included and cultured on the chromID-ESBL selection media. Molecular characterization was done using polymerase chain reaction and sequencing for *bla*<sub>CTX-Ms</sub>, *bla*<sub>TEMs</sub>, *bla*<sub>SHVs</sub>, plasmid-mediated quinolone-resistant determinants, O25-ST131, phylogenetic groups, pulsed-field gel electrophoresis (PFGE), and multilocus sequencing typing.

**Results.** A total of 226 individuals were included; 195 (86%) were negative, and 31 (14%) were positive for ESBL-producing *E coli*. Notably, travelers were 5.2 (95% CI 2.1–31.1) times more likely than non-travelers to have an ESBL-producing *E coli*





# Colonization of Returning Travelers With CTX-M-Producing *Escherichia coli*

- stool samples submitted from travelers as compared to non-travelers
- travelers were 5.2 (95% CI 2.1–31.1) times more likely than non-travelers to have an ESBL-producing *E coli*
- Confirms that foreign travel, especially to the Indian subcontinent and Africa, represents a major risk for rectal colonization with CTX-M-producing *E coli* and contributed to the Worldwide spread of these bacteria



Scand J Infect Dis. 1983;15(4):367-73.

## **Changes in serotype and resistance pattern of the intestinal *Escherichia coli* flora during travel. Results from a trial of mecillinam as a prophylactic against travellers' diarrhoea.**

Stenderup J, Orskov I, Orskov F.

### **Abstract**

The changes in the intestinal *Escherichia coli* flora during travel has been studied by serological methods. A group of 74 tourists visiting Egypt and the Far East were given mecillinam or placebo in a randomized double-blind study. In all but 3 participants, 2 in the placebo group and 1 in the mecillinam group, a complete change in the *E. coli* flora occurred after a few days, and changes continued to occur during the 25 days of travel. The percentage of multiresistant strains rose from 8% in the pretravel samples to 50-60% in the posttravel samples. Less than 5% of the pretravel *E. coli* strains were resistant to mecillinam, whereas in the posttravel samples 42.9% of the *E. coli* strains in the mecillinam group and 19.1% in the placebo group were resistant to mecillinam. Of the 30 mecillinam resistant *E. coli* strains from the diarrhoeal samples only 6 showed transferable mecillinam resistance.

PMID: 6318304 [PubMed - indexed for MEDLINE]



# What about the patient?

- What shall we do?
- Which antibiotic should we choose?









- Comparisons among the fully sequenced genomes of nonpathogenic and pathogenic strains have revealed an average genome size of approximately 5000 genes, but only approximately 2200 of these are shared among all *E. coli* strains. Most of the pathogens have larger genomes than do the nonpathogenic strains.[\[26.99\]](#) Furthermore, many of the genes that are not found in the nonpathogenic strain are specific to particular strains or pathotypes. It is estimated that the total “pangenome” of *E. coli* consists of more than 13,000 genes.[\[99\]](#)

