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Prevalence, risk factors for and trends in faecal colonisation by extended-spectrum beta-lactamase-producing Enterobacterales (ESBL-PE) in the community in Spain: A cross-sectional study and meta-analysis



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ABSTRACT

Objectives: To determine the prevalence of and risk factors for extended-spectrum beta-lactamase-producing Enterobacterales (ESBL-PE) faecal carriage among healthy volunteers from Barcelona, and to estimate the pooled prevalence in the community in Spain.

Methods: University students were asked to complete a questionnaire and provide a rectal swab, which was tested for ESBL-producing, ciprofloxacin- and trimethoprim–sulfamethoxazole-resistant Enterobacterales. Risk factors for carriage of antimicrobial resistance were identified by multivariate logistic regression. To place these results in the appropriate context, a systematic literature search was conducted to retrieve articles containing data on the prevalence of ESBL-PE faecal carriage in the community in Spain. To obtain the pooled prevalence, a random-effects meta-analysis was performed.

Results: One hundred and thirty-five of 214 participants were included in the analysis. Faecal carriage of Escherichia coli/Klebsiella pneumoniae (E/K) resistant to at least one of the antibiotics tested was found in 32 participants (23.7%). Fourteen subjects carried ESBL-E/K (10.4%), with the CTX-M type being the most prevalent (85.7%). Risk factors for ESBL carriage were travel to a high-risk region in the past 3 years (OR 5.66; 95% CI 1.07–29.9) and living in a crowded city district (OR 6.91; 95% CI 1.22–39.08). Thirteen articles covering 21,760 individuals from Spain were included in the meta-analysis, giving a pooled prevalence rate for ESBL-PE carriage in the community of 5.8% (95% CI 4.1–7.8%), and a steady increase per year. Conclusions: The faecal colonisation prevalence by ESBL-PE among healthy individuals in Spain is high. It is associated with international travel and living in crowded city districts.

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Prevalencia, factores de riesgo y tendencias en la colonización fecal por Enterobacterales productores de beta-lactamasas de espectro extendido en la comunidad en España: un estudio transversal y metaanálisis

RESUMEN

Palabras clave:
Betalactamasas de especto extendido (BLEE)
Prevalencia
Factores de riesgo
Comunidad
Población sana
Metaanálisis

Objetivos: Determinar la prevalencia y los factores de riesgo de la colonización fecal por ESBL-PE entre voluntarios sanos de Barcelona, así como estimar la prevalencia en la comunidad en España.

Métodos: Estudiantes universitarios completaron un cuestionario y proporcionaron un frotis rectal, que fue analizado para detectar Enterobacterales productores de betalactamasas de espectro extendido (BLEE-PE) y resistentes a ciprofloxacino y a trimetoprima-sulfametoxazol. Los factores de riesgo se identificaron mediante regresión logística multivariada. Para situar estos resultados en el contexto adecuado, se realizó una búsqueda sistemática de la literatura para recuperar artículos con datos sobre la prevalencia de la colonización fecal por BLEE-PE en la comunidad en España. Para obtener la prevalencia combinada, se realizó un metaanálisis de efectos aleatorios.

Resultados: Ciento treinta y cinco de los 214 participantes fueron incluidos en el análisis. Se encontró colonización fecal de *E. coli/K. pneumoniae* (E/K) resistente a al menos uno de los antibióticos analizados en 32 participantes (23,7%). Catorce sujetos portaban BLEE-E/K (10,4%), siendo el tipo CTX-M el más prevalente (85,7%). Los factores de riesgo para la colonización por BLEE fueron viajar a una región de alto riesgo en los últimos 3 años (OR: 5,66; IC 95%: 1,07-29,9) y vivir en un distrito urbano densamente poblado (OR: 6,91; IC 95%: 1,22-39,08). En el metaanálisis se incluyeron 13 artículos que abarcaban a 21.760 individuos de España, resultando una tasa de prevalencia combinada de colonización por BLEE-PE en la comunidad del 5,8% (IC 95%: 4,1-7,8%), con un con un aumento anual constante.

Conclusiones: La prevalencia de la colonización fecal por BLEE-PE entre individuos sanos en España es alta. Está asociada con viajes internacionales y vivir en distritos urbanos densamente poblados.

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Introduction

Antibiotic resistance in community-acquired infections is becoming a major public health threat. In this context, Enterobacterales, mainly those producing extended-spectrum beta-lactamases (ESBL), are a major concern: patients often require carbapenems and treatment options are limited. ²

Genes conferring ESBL activity are easily transmitted and thus disseminated.³ Furthermore, plasmids carrying ESBLs often encode other antibiotic resistance genes such as plasmid-mediated quinolone resistance determinants or sulphonamide resistance genes, conferring on ESBL-producing Enterobacterales (ESBL-PE) a much broader spectrum of resistance.

ESBL-PE faecal carriage rates in the community are increasing worldwide. 1,3,4 And while the presence of this gut microbiota reservoir is usually asymptomatic, many studies have shown a worse outcome in patients with ESBL-PE infections. 5

In Europe, the reported community carriage prevalence is 6% (data from studies conducted between 2004 and 2019).^{1,4} In Spain, a few studies have assessed ESBL-PE carriage rates in the community,^{6–8} but a pooled prevalence has not been reported.

Hypothesised attributable sources of human exposure to ESBLs include animals, food, the environment, and other human beings. ^{9,10} However, it remains unclear which transmission route or risk factor for acquisition antibiotic-resistant bacteria predominates

The aim of the present study was to estimate the prevalence of faecal carriage of ESBL-producing, ciprofloxacin-(CIP) and trimethoprim-sulfamethoxazole (SXT)-resistant Enterobacterales in healthy volunteers from Barcelona, Spain, and to assess the risk factors associated with colonisation status. To place these results in the appropriate geographical and temporal context, a comprehensive literature review and meta-analysis of published community carriage rates in Spain was undertaken.

Methods

This study was reported following the Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) checklist for cross-sectional studies and the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) 2009 checklist.

Participant data collection

A total of around 240 healthy first- and second-year medical students living in Barcelona or surrounding areas were invited to participate by direct face-to-face recruitment between January 2020 and September 2020. After providing informed consent, participants received material to self-collect a rectal swab with the indications of inserting it about 3–5 cm into the anal canal and rubbing it gently for about 5–10 s against the walls of the rectum. Participants were also asked to fill in a detailed questionnaire about possible risk factors for acquiring drug-resistant bacteria (demographics, use of medication, contact with animals, diet, hygiene, and travel history). Regions where >25% of *Escherichia coli* clinical isolates are estimated to be ESBL producers⁹ were defined as high-risk regions.

Microbiology

Rectal swabs were taken using Deltalab Amies agar gel transport swabs and streaked on chromIDTM ESBL, CARBA/OXA and Mac-Conkey agar plates (bioMérieux, France). A first reading was taken after 24 h and a second one after 48 h of incubation at 37 °C. These selective and differential media were used for presumptive identification of *E. coli* (pink colonies) and *Klebsiella pneumoniae* (green colonies). On MacConkey agar, both colonies are lactose-positive and the difference is established by the consistency of bacterial isolates: dry (*E. coli*) and mucous (*K. pneumoniae*). Definitive bacterial

identification was performed using matrix-assisted laser desorption/ionisation time-of-flight (MALDI-TOF) mass spectrometry.

Antibiotic susceptibility testing was performed by the disc diffusion method on Mueller-Hinton agar plates, including a synergy test to detect/confirm ESBL production and paper disks impregnated with ciprofloxacin and SXT to detect resistance to these antibiotics. Results were interpreted following European Committee on Antimicrobial Susceptibility Testing (EUCAST) clinical breakpoints (Version 9.0).¹¹

The ESBL-producing isolates were subjected to nucleotide sequencing to determine their molecular type. DNA extraction was performed using DNeasy Ultra Clean Microbial kit (Qiagen) as described by manufacturer. Genome DNA was quantified using Qubit 3.0 (Thermofisher). Library preparation and Whole Genome Sequencing were performed by NovoGene (Novogene Sequencing Europe) on an Illumina NovaSeq instrument with Illumina 2×150 cycles sequencing. Reads were quality trimmed using fastp v0.23.2 (available at: https://github.com/OpenGene/fastp); trimmed reads were de novo assembled using Unicycler v0.5.0, with Spades v3.15.5 (available at: https://github.com/rrwick/Unicycler/releases and https://github.com/ablab/spades, respectively). Fasta files were then submitted to ResFinder and CARD to betalactamse resistance prediction. Sequence types (ST) were determined using two schemes for E. coli (Achtman and Pasteur MLST) and one scheme for K. pneumoniae (Pasteur MLST).

Metanalysis data sources and search terms

A systematic literature search was conducted in PubMed to retrieve relevant articles published between 1 January 2000 and 23 December 2020. Five groups of search terms were used: (i) Escherichia coli OR E. coli OR Enterobacterales; (ii) extended-spectrum beta-lactamase OR ESBL; (iii) faecal OR faeces OR stool OR intestinal OR gastrointestinal tract; and (iv) community OR community-acquired OR healthy OR volunteers OR at admission; (v) Spain OR Barcelona OR Madrid. These groups were then connected by the Boolean operator 'AND' to find papers that contained these terms anywhere in the article.

Statistical analyses

For descriptive statistics, quantitative variables were expressed as mean and standard deviation. Qualitative variables were expressed as frequencies (number and percentage). To check the association of continuous and categorical variables with ESBLproducing, CIP- and SXT-resistant Enterobacterales carriage, the Student's t-test and Chi-squared test (or Fisher's exact in the case of expected numbers < 5) were performed, as appropriate. Logistic regression was used to study the risk factors associated with the carriage of antibiotic-resistant E. coli or K. pneumoniae. Four logistic regression models were developed (one for each type of resistance (ESBL-producing or resistance to CIP or SXT) and an additional one combining the three resistance types assessed). All variables with a p value < 0.05 in univariate analysis, as well as other variables considered clinically relevant (age, place of residence, travel history and meat consumption) were included in multivariate models. Results are presented as odd ratio (OR) with 95% confidence interval (CI). A p value < 0.05 was used to determine significance. All analyses were performed using IBM SPSS Statistics 25.

The meta-analysis was performed using a random-effects method to obtain a pooled prevalence and estimate the national trend of faecal ESBL-PE carriage. The Freeman-Tukey arcsine methodology was used to stabilise the variance of raw proportions, and no studies with 0% or 100% proportions were excluded. The l^2 statistic was used to test heterogeneity. Following this meta-analysis, we performed a univariate meta-regression taking the

year of publication as predictor variable. Results from this metaregression were expressed by means of β (coefficient of year) and plotting a bubble plot (prevalence by year).

Ethics

This study was approved by the Parc de Salut Mar-IMIM ethics committee in Barcelona (no. 2019/8971/I). All participants provided written informed consent. The present study was carried out in accordance with current regulations and the basic principles of the protection of the rights and dignity of the human being, as stated in the Helsinki Declaration (64th General Assembly, Brazil, October 2013).

Results

Observational study

A total of 214 participants were enrolled in the study, 135 of which were included in the analysis (Fig. S1). Seventy-nine participants were excluded, either because they did not respond to the questionnaire (n=1), or their rectal swabs were lost (n=46) or not well taken (defined as non-bacterial growth in MacConkey agar plates after 48 h of incubation at $37\,^{\circ}$ C) (n=30), or the samples were well taken but neither *E. coli* nor *K. pneumoniae* were identified on MacConkey agar plates (n=2). Comparison of baseline characteristics of participants not included in the analysis because of not well taken swab samples and those finally included is shown in Table S1.

Faecal carriage of *E. coli/K. pneumoniae* (E/K) with at least one of the phenotypes of interest (ESBL-producing or resistance to CIP or SXT) was found in 32 participants (23.7 \pm 7.17%, 95% CI): 14 subjects carried ESBL-E/K (10.4 \pm 5.14%, 95% CI), 20 carried co-trimoxazole (SXT)-resistant E/K (14.8 \pm 5.99%, 95% CI) and 14 carried ciprofloxacin (CIP)-resistant E/K (10.4 \pm 5.14%, 95% CI). CIP and/or SXT co-resistance in ESBL-E/K was found in 9 out of 14 (64.3%) of the ESBL-E/K identified.

The genotypic characterisation of the 14 ESBL-E/K producers isolated is shown in Table 1. Eleven out of 14 (78.5%) isolates were *E. coli* and three out of 14 (21.4%) were *K. pneumoniae*. Multidrug resistance (resistance to at least three antibiotic families) was observed in 92.8% of the ESBL-producing strains. Ten out of 14 (71.4%) isolates had more than one type of beta-lactamase. The CTX-M-type was the most prevalent (85.7%) followed by the TEM-type (42.9%) and the SHV-type (28.5%). Discrepancies between phenotypic and genotypic ESBL confirmation tests were only seen with *E. coli* isolate ST453, as TEM-1 beta-lactamase is not considered an ESBL; a false positive result probably explained by TEM-1 hyperproduction combined with altered permeability, a mechanism argued in EUCAST guidelines. All the *K. pneumoniae* isolates belonged to the same sequence type: ST469. *E. coli* isolates belonged to different sequence types: ST69, ST131, ST540, ST453, ST93 and ST227.

The characteristics of the participants are presented in Tables 2a and 2b. Table 3 shows the multivariate analysis of potential risk factors for the carriage of antibiotic-resistant E/K. Antibiotic therapy in the previous 3 months (OR 4.10; 95% CI 1.14–14.83), travel to a high-risk region in the past 3 years (OR 2.95; 95% CI 1.04–8.36) and living in Eixample (a crowded district in the Barcelona centre) (OR 4.61; 95% CI 1.13–18.82) were found to be independently associated with carriage of Enterobacterales resistant to any of the three antimicrobials assessed.

Specific risk factors for ESBL-E/K carriage were travel to a highrisk region in the past 3 years (OR 5.66; 95% CI 1.07–29.9) and living in Eixample district (OR 6.91; 95% CI 1.22–39.08). Risk factor for SXT-resistant E/K carriage was previous antibiotic therapy. No spe-

Table 1Sequence type (ST) and resistome of 14 ESBL-E. coli/K. pneumoniae strains isolated.

Strain	Beta-lactamase	Aminogly- cosides	Fosfomycin	MLSb	Quinolones	Sulpho- namides	Trime- thoprim	Tetra- cyclines	Chloram- phenicol
E. coli ST69/ST3	blaCTX-M- 55 + blaTEM-1	aph(3')-Ia; aph(3")-Ib; aph(6)-Id; aadA1; aadA2; aadA5	fosA3	erm(B); mph(A)		sul1; sul2; sul3	dfrA12; dfrA17	tet(A)	cml (96.6%; 929/909) ^b
E. coli ST69/ST3	blaCTX-M- 14+blaTEM-1	aac(3)-Iid; aadA5		mph(A)		sul1	dfrA17	tet(B)	
E. coli ST69/ST3	blaCTX-M- 2 + blaTEM-1	aph(3")-Ib; aph(6)-Id; aac(3)-Iia; aac(3)-Via; aadA1			qnrB19	sul1; sul2	dfrA7	tet(A) + tet(D)	
E. coli ST131/ST506	blaCTX-M-15	not found		erm(B); mph(A)			dfrA14		
E. coli ST131/ST43 ^a	blaCTX-M-27	aph(6)-Id; aph(3")-Ib; aadA5		erm(B); mph(A)		sul1; sul2	dfrA17	tet(A)	
E. coli ST131/ST506	blaCTX-M-27	aph(6)-Id; aph(3")-Ib; aadA5		mph(A)		sul1; sul2	dfrA17	tet(A)	
E. coli ST4980 ^a /new	blaSHV-12	aadA1; aadA2b				sul3		tet(A)	cmlA1
E. coli ST540/ST716	blaCTX-M- 15 + blaTEM-1	aph(3")-Ib; aph(6)-Id			qnrS1	sul2		tet(A)	
E. coli ST453/ST86	blaTEM-1	aph(3")-Ib; aph(6)-Id				sul2	dfrA17	tet(A)	
E. coli ST93/ST17	blaCTX-M- 32+blaTEM-1	aadA12				sul1		tet(A)	
E. coli ST227/ST638	blaCTX-M-15	not found			qnrS1				
K. pneumoniae ST469	blaCTX-M- 15+blaSHV11	aph(6)-Id; aadA2 aph(3″)-Ib	fosA	mph(A)	qnrS1; OqxA/OqxB	sul1: sul2	dfrA12		
K. pneumoniae ST469	blaCTX-M- 15+blaSHV11	aph(6)-Id; aadA2 aph(3")-Ib	fosA	mph(A)	qnrS1; OqxA/OqxB	sul1: sul2	dfrA12		
K. pneumoniae ST469	blaCTX-M- 15 + blaSHV11	aph(6)-Id; aadA2 aph(3")-Ib	fosA	mph(A)	qnrS1; OqxA/OqxB	sul1: sul2	dfrA12		

To determine the ST for E. coli, two schemes were used: MLST Achtman which includes loci adk, fumC, gyrB, icd, mdh, purA, recA, and MLST Pasteur which includes loci dinB, icdA, pabB, polB, putP, trpA, trpB, uidA.

cific risk factors for ciprofloxacin-resistant E/K carriage were found. In our models, meat consumption (white or red) did not appear as a risk factor.

Literature systematic research and metanalysis

Twenty-four articles indexed in PubMed were retrieved for further screening. Nine articles included in two meta-analyses on this topic $^{3.4}$ conducted in Spain were also retrieved for screening (Fig. S2). For more information on how the studies were selected and which data were extracted, see supplementary material and Table S2. The presence of publication bias was assessed using Egger's test (Fig. S3). The 13 studies $^{12-20}$ analysed in the meta-analysis covered a total of 21,760 individuals from different Spanish regions between 2001 and 2020. This gave a national pooled prevalence of community intestinal carriage of ESBL-PE of 5.8% (95% CI 4.1–7.8%) (Fig. 1). A goodness-of-fit analysis comparing the national pooled prevalence derived from the meta-analysis with the ESBL carriage rate reported in the present study (10.4%) showed a statistically significant difference (p = 0.027).

Results from linear meta-regression analysis, which included the present study, showed that year of study was positively associated with prevalence (β =0.0057; p-value=0.031). Results are shown in Fig. 2.

Discussion

Our study shows that the prevalence of faecal colonisation by multidrug-resistant Enterobacterales (23.7%) and ESBL-PE (10.4%) among healthy individuals in Spain is not negligible. The prevalence of ESBL-PE confirms and exceeds the results of previous studies

performed in our country, showing an overall pooled prevalence of 5.8% and a steady increase per year. The reported ESBL-PE prevalence among participants included in our study did not differ from the estimated one by 2020 via linear meta-regression analysis of the time trend of ESBL-PE carriage in Spain. These results strengthen confidence in the methods used and reduce the risk of selection or information bias.

ESBL-PE prevalence being positively correlated with year of study, coincides with the existing evidence informed by other meta-analyses of an annual steady increase in the overall global pooled carriage rate. The worst projections estimate an ESBLE-PE prevalence of 43% globally by 2030 (with an annual increase of 1.5%). There is limited data from other European countries for comparison. In the Netherlands, a study including 4177 participants from the general population reported an ESBL-PE carriage rate of 5.0% (95% CI: 3.4–6.6%), less than half the prevalence observed in our study.

Given the poorer prognosis of patients with ESBL-PE infection, 5 these increasing rates, together with the high co-resistance detected in ESBL-PE, offer a future scenario in which, if no action is taken, carbapenems may frequently have to be used as empirical treatment of systemic or organ-affecting bacterial infections due to Enterobacterales, and not only for patients with recognised risk factors for multi-drug resistant Enterobacterales carriage. According to the projection made in this study, this assumption could be a reality in Spain in 2050, when ESBL-PE carriage rates in the community will have reached 20%. This threshold has been used by guidelines to advise against empirical treatment with fluoroquinolones in pyelonephritis, 3 since resistance prevalences as high as 50% have been reported in community-onset healthcare-associated urinary tract infections in this country. 4

a In these cases, the closest ST was indicated. For ST43, locus trpB and for ST4980, locus adk were new alleles in the MLST Pasteur scheme.

^b The similarity coefficient (%) represents the proportion of compared nucleotides/gene nucleotides.

Table 2aDescriptive characteristics of faecal carriers of the ESBL-producing or CIP or SXT resistant bacteria vs non-carriers.

	ESBL, CI		
	Carriers [n = 32 (%)]	Non-carriers [n = 103 (%)]	p value
Female Age-mean ± SD (years)	23(71.9) 20.63 ± 2.67	65 (63) 21.65 ± 3.57	0.403 0.136
Health Hospitalized in the last year ATB treatment in the last 3 months Close contacts ^a recently treated with ATB Close contacts hospitalized in the last year	1 (3.1) 7 (21.9) 5 (15.6) 5 (15.6)	6(5.8) 9(8.7) 20(19.4) 13(12.6)	1.000 0.060 0.796 0.766
Source of products of animal origin ^b Store 1 Store 2	19(59.4) 6(18.8)	62(60.2) 35 (34)	1 0.126
International travel (last 3 years) Been outside Europe Been to a high-risk region Number of trips made mean ± SD	$11(35.5) \\ 22(68.8) \\ 3.88 \pm 3.13$	41(39.8) 54(52.9) 4.13 ± 3.36	0.834 0.153 0.708
Animals and pets Regular contact with farms Living in a rural environment Pet owner	4(12.5) 2(6.3) 11(34.4)	4(3.9) 10(9.7) 46(44.7)	0.090 0.731 0.413
Place of residence ^b Barcelona city	19(61.3)	42(41.1)	0.064
Barcelona district Eixample (centre)	8(26.7)	9(9.2)	0.027*
Handwashing after WC No Yes, with water Yes, with soap and water Yes, with alcohol	1 (3.1) 5 (15.6) 26(81.3) 0 (0)	6(5.8) 14(13.6) 80(77.7) 3(2.9)	1
Source of drinking water Tap water Purified tap water Bottled water	13(40.6) 9(28.1) 10(31.3)	51 (49.5) 19 (18.4) 33 (32.0)	0.485
Diet Vegetarian White meat servings per week – mean ± SD Red meat servings per week – mean ± SD Seafood servings per week – mean ± SD Dairy servings per week – mean ± SD Volume of drinking water per day – mean ± SD (litres)	$2(6.3)$ 3.13 ± 1.90 2.75 ± 1.78 2.22 ± 1.39 6.72 ± 3.24 1.51 ± 0.59	$11 (10.7)$ 2.51 ± 1.66 2.21 ± 1.73 2.35 ± 1.74 6.97 ± 2.67 1.58 ± 0.52	0.733 0.078 0.126 0.768 0.658 0.514
Contacts Roommates/family members – mean ± SD	2.56 ± 1.19	2.70 ± 1.05	0.534

CIP: ciprofloxacin. SXT: co-trimoxazole. ATB: antibiotic.

With regard to the molecular characterisation of beta-lactamase genes, our results coincide with those reported in other studies conducted in Spain, in which CTX-M type enzymes predominate in healthy individuals, with the $bla_{\text{CTX-M-15}}$ variant being the most prevalent.²⁵ As for sequence types, the most frequently detected E coli isolates (ST131 and ST69) in our study have also been described as prominent infection isolates globally.²⁶ In addition, three K pneumoniae isolates belonged to the same sequence type, suggesting some direct relation among carriers which could not be further confirmed.

Consistent with previous reports,⁴ previous antibiotic consumption and travel to regions with high ESBL-PE prevalence were found to be associated with carriage of antimicrobial resistant bacteria. These results could be related to selective pressure on antibiotics prescribed in the community, and increased faecal-oral, human-to-human transmission associated with poor hygiene in regions where ESBL-PE prevalence is high (such as Vietnam, Laos

or China).¹ Spanish guidelines for the treatment of multidrugresistant Gram-negative bacterial infections suggest that recent travel to a region of high ESBL-PE prevalence should be considered when deciding whether ESBL coverage with carbapenems is needed.²² However, this risk factor has not yet been included in any of the predictive scores currently used in clinical practice for infection due to ESBL-PE at the time of hospital admission.

Living in the city centre of Barcelona (specifically, the Eixample district, which is densely populated) was also found to be a risk factor associated with carriage of antibiotic resistant bacteria and, specifically ESBL-producing Enterobacterales. This result fits the theory that more crowded areas may facilitate human-to-human transmission.²⁷

Although the use of antibiotics in veterinary practices raised concerns that animal-derived food products represent an important route of acquisition, meat consumption (white or red) did not appear to be a risk factor in our study. Overall, these results

^a Close contacts defined as roommates or family members.

b Additional variables on the source of animal-derived products and place of residence can be found in supplementary information (Table S3).

^{*} Significant at p < 0.05.

Table 2bDescriptive characteristics of faecal carriers of each type of resistance (ESBL-producing or resistance to CIP or SXT) vs non-carriers.

	ESBL			CIP			SXT		
	Carriers [<i>n</i> = 14 (%)]	Non-carriers [n = 121 (%)]	p value	Carriers [n = 14 (%)]	Non-carriers [n = 121 (%)]	p value	Carriers [n = 20 (%)]	Non-carriers [n = 115 (%)]	p value
Female Age – mean ± SD (years)	11 (78.6) 20.50 ± 2.53	78 (63.6) 21.51 ± 3.48	0.378 0.243	11 (78.6) 20.43 ± 2.86	77 (63.6) 21.52 ± 3.45	0.378 0.256	15 (75) 20.05 ± 2.67	73 (63.5) 21.64 ± 3.47	0.447 0.053
Health									
Hospitalized in the last year	0(0)	7(5.8)	1	0(0)	7(5.8)	1	1 (5)	6(5.2)	1
ATB treatment in the last 3 months	1(7.1)	15(12.4)	1	1(7.1)	15(12.4)	1	6 (30)	10(8.7)	0.015*
Close contacts ^a recently treated with	2(14.3)	23 (19)	1	3(21.4)	22(18.2)	0.723	3 (15)	22(19.1)	1
ATB									
Close contacts hospitalized in the	2(14.3)	16(13.2)	1	3(21.4)	15(12.4)	0.400	3 (15)	15 (13)	0.731
last year									
Source of animal-derived products ^b									
Store 1	6(42.9)	75 (62)	0.248	9(64.3)	72 (59.5)	0.782	10 (50)	71 (61.7)	0.334
Store 2	5(35.7)	36(29.8)	0.760	3(21.4)	38(31.4)	0.551	4 (20)	37(32.2)	0.429
International travel (last 3 years)									
Been outside Europe	6(42.9)	46(38.3)	0.777	3(23.1)	49(40.5)	0.369	8(42.1)	44(38.3)	0.802
Been to a high-risk region	11(78.6)	65 (54.2)	0.095	10(71.4)	66 (55)	0.271	12 (60)	64(56.1)	0.811
Number of trips mean \pm SD	4.50 ± 3.84	4.02 ± 3.25	0.606	3.57 ± 2.17	4.12 ± 3.41	0.555	3.95 ± 2.69	4.09 ± 3.40	0.865
Animals and pets	2/1/2)	C (F)	0.105	1 (7.1)	7(5.0)	0.504	1 (5)	7(6.1)	
Regular contact with farms	2(14.3)	6(5)	0.195	1(7.1)	7(5.8)	0.594	1 (5)	7(6.1)	1
Living in a rural environment	2(14.3)	10(8.3)	0.360	1(7.1)	11(9.1)	1	0(0)	12(10.4)	0.213
Pet owner	4(28.6)	53 (43.8)	0.393	3(21.4)	54(44.6)	0.152	7 (35)	50(43.5)	0.625
Place of residence ^b									
Barcelona city	8(61.5)	53(44.2)	0.256	8(61.5)	53(43.4)	0.250	13 (65)	48 (42.5)	0.088
Barcelona district									
Eixample (centre)	5(38.5)	12(10.4)	0.015*	4(28.6)	13(11.4)	0.092	3(15.8)	14(12.8)	0.718
Handwashing after WC			0.762			0.658			0.872
No	1(7.1)	6 (5)	0.702	0(0)	7(5.8)	0.036	0(0)	7(6.1)	0.072
Yes, with water	3(21.4)	16(13.2)		3(21.4)	16(13.2)		3 (15)	16(13.9)	
Yes, with soap and water	10(71.4)	96(79.3)		11(78.6)	95 (78.5)		17 (85)	89(77.4)	
Yes, with alcohol	0(0)	3(2.5)		0(0)	3(2.5)		0(0)	3(2.6)	
	0 (0)	3(2.3)		0 (0)	3(2.3)		0(0)	3(2.0)	
Source of drinking water			0.747			0.094			0.950
Tap water	6(42.9)	58(47.9)		6(42.9)	58(47.9)		9 (45)	55(47.8)	
Purified tap water	4(28.6)	24(19.8)		6(42.9)	22(18.2)		4 (20)	24(20.9)	
Bottled water	4(28.6)	39(32.2)		2(14.3)	41 (33.9)		7 (35)	36(31.3)	
Diet									
Vegetarian	1(7.1)	12(9.9)	1	2(14.3)	11(9.1)	0.626	1 (5)	12(10.4)	0.691
White meat servings per week –	2.93 ± 2.17	2.62 ± 1.68	0.536	2.86 ± 2.35	2.63 ± 1.65	0.648	2.80 ± 1.54	2.63 ± 1.77	0.689
mean ± SD					,			,,	
Red meat servings per week –	2.36 ± 1.60	2.33 ± 1.77	0.962	2.50 ± 1.70	2.32 ± 1.76	0.712	3.10 ± 1.71	2.20 ± 1.73	0.034*
mean ± SD									
Seafood servings per week –	2.36 ± 1.65	2.32 ± 1.66	0.941	2.50 ± 1.40	2.31 ± 1.69	0.680	2.05 ± 1.47	2.37 ± 1.69	0.422
mean ± SD									
Dairy servings per week - mean ± SD	6.86 ± 2.91	$\boldsymbol{6.92 \pm 2.80}$	0.940	6.36 ± 2.71	$\boldsymbol{6.98 \pm 2.82}$	0.436	7.30 ± 3.11	6.84 ± 2.75	0.503
Volume of drinking water per day -	1.54 ± 0.73	1.58 ± 0.51	0.783	1.57 ± 0.70	1.56 ± 0.51	0.952	1.43 ± 0.55	1.59 ± 0.53	0.218
mean ± SD (litres)									
Contacts									
Roommates/family members -	2.79 ± 0.98	2.65 ± 1.09	0.664	2.93 ± 1.14	2.64 ± 1.07	0.339	2.30 ± 1.26	2.73 ± 1.04	0.100
$mean \pm SD$									

CIP: ciprofloxacin. SXT: co-trimoxazole. ATB: antibiotic.

could support faecal-oral human-to-human transmission as the main route of ESBL-PE acquisition in the community, as Mughini-Gras L et al. recently demonstrated. In their study, 60% of community-acquired ESBL *E. coli* was attributable to human-to-human transmission, whereas food accounted for about 20%. The same conclusion was reached by a recent study from the UK, which found few resistance genes common to both livestock and humans. Food has also been posited as the most important non-human source for the acquisition of ESBL-PE. Direct exposure to products colonised by ESBL-PE could be one explanation. Another suggested mechanism is the ingestion of antibiotic residues that promote the emergence and selection of resistance genes in the

human microbiome.²⁹ Significant amounts of these antibiotic residues have been found, mainly in animal source foods.³⁰

In terms of limitations, the study population is not representative of healthy young adults living in Barcelona, as only university students were included, and consequently subjects from less favourable socioeconomic backgrounds were less likely to be selected. However, unlike other well-known bacteria, such as *Helicobacter pylori*, ESBL-PE carriage has not been linked to lower socioeconomic groups.²¹ For this reason, we consider it unlikely that the real prevalence among healthy young adults was underestimated. The risk of measuring hospital-acquired bacterial carriage was reduced by including only medical students in years 1 and 2

^a Close contacts defined as roommates or family members.

b Additional variables on the source of animal-derived products and place of residence can be found in supplementary information (Table S3).

^{*} Significant at p < 0.05.

Table 3Multivariate analysis models of risk factors for different antimicrobial resistance profiles.

ESBL, CIP-R and SXT-R model (1)				ESBL m			
Variable	Multivariate OR	95% CI	p value	Variable	Multivariate OR	95% CI	p value
Age	0.87	0.73-1.03	0.114	Age	0.86	0.66-1.13	0.272
Recent ATB use	4.10	1.14-14.83	0.031*	Recent ATB use	0.81	0.09-7.62	0.854
Been to a high-risk region	2.95	1.04-8.36	0.042*	Been to a high-risk region	5.66	1.07-29.9	0.041*
Place of residence: Barcelona city	1.39	0.48-4.04	0.544	Place of residence: Barcelona city	1.02	0.22 - 4.75	0.976
District of residence: Eixample	4.61	1.13-18.82	0.033*	District of residence: Eixample	6.91	1.22-39.08	0.029*
White meat consumption	1.12	0.84-1.50	0.424	White meat consumption	1.08	0.76-1.53	0.660
Red meat consumption	1.22	0.92-1.62	0.170	Red meat consumption	0.99	0.67-1.47	0.969
CIP-R model (3)				SXT-R model (4)			
Variable	Multivariate OR	95% CI	p value	Variable	Multivariate OR	95% CI	p value
Age	0.89	0.71-1.11	0.293	Age	0.89	0.74-1.07	0.208
Recent ATB use	0.45	0.05 - 3.89	0.466	Recent ATB use	3.81	1.07-13.59	0.039*
Been to a high-risk region	1.86	0.54-6.45	0.327	Been to a high-risk region	1.23	0.42-3.61	0.707
Place of residence: Barcelona city	1.62	0.52 - 5.04	0.408	Place of residence: Barcelona city	2.77	0.94-8.18	0.065
White meat consumption	1.07	0.76-1.50	0.717	White meat consumption	0.90	0.64-1.25	0.525
Red meat consumption	1.06	0.74-1.50	0.755	Red meat consumption	1.35	0.99-1.82	0.051

ESBL: extended-spectrum beta-lactamases. ATB: antibiotic. CIP-R: ciprofloxacin resistant. SXT-R: co-trimoxazole resistant.

All variables with a *p* value < 0.05 in univariate analysis were included in the multivariate models (district of residence: eixample for models 1 and 2, read meat consumption and recent ATB use in model 4). All other variables were included because there were considered clinically relevant (age, place of residence, travel history and meat consumption).

^{*} Significant at p < 0.05.

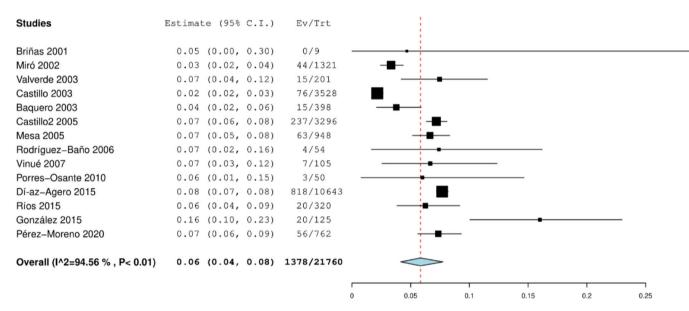


Fig. 1. Forest plot showing the prevalence of ESBL carriage for each study included in the meta-analysis. Plot created with OpenMeta (Analyst).

(without contact with medical care facilities). Moreover, the prevalence seen within young adults may be similar in other age groups as ESBL carriage status has not been associated with this variable. Exposure assessment is not likely to be greatly affected by recall bias, as the subjects answered the questionnaire without knowing their colonisation status, and there is no reason to think that participants carrying resistant bacteria would report differently from non-carriers. Even so, real differences between groups could have been overlooked due to low statistical power.

Our results, together with the existing evidence, may support the development of updated predictive scores for ESBL-PE infection at hospital admission, which should include recent travel to a region of high ESBL-PE prevalence, and living in crowed city districts as additional criteria for suspected ESBL-PE infections.

Further investigations with larger sample sizes should address the role of dietary intake as a possible acquisition pathway in more detail, focusing on non-animal food products such as vegetables

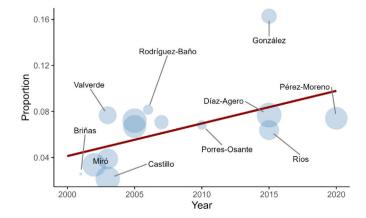


Fig. 2. Results from meta-regression representing the evolution of ESBL prevalence across the years (β = 0.0057; p-value = 0.031). Plot created with R package metafor.

and fruit, which could represent an important reservoir for resistant bacteria. Measuring the levels of antibiotics present in food products, as well as in biological samples from subjects, may help determine whether long-term low dose antibiotic exposure contributes to this increasing prevalence of resistance prevalence.

Finally, given the increasing rates of antimicrobial resistant and ESBL-PE carriage among healthy individuals in Spain, general measures like community surveillance plans and antimicrobial stewardship programmes should be implemented in an effort to slow the progression and reduce the potential deleterious effects of this concerning issue.

Conflicts of interest

None.

Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.gastrohep.2024.502236.

References

- Bezabih YM, Sabiiti W, Alamneh E, Bezabih A, Peterson GM, Bezabhe WM, et al. The global prevalence and trend of human intestinal carriage of ESBL-producing Escherichia coli in the community. J Antimicrob Chemother. 2021;76:22–9, http://dx.doi.org/10.1093/jac/dkaa399.
- Paterson DL, Isler B, Harris PNA. PRO: carbapenems should be used for ALL infections caused by ceftriaxone-resistant Enterobacterales. JAC Antimicrob Resist. 2021;3:dlab013, http://dx.doi.org/10.1093/jacamr/dlab013.
- 3. Woerther PL, Burdet C, Chachaty E, Andremont A. Trends in human fecal carriage of extended-spectrum β-lactamases in the community: toward the globalization of CTX-M. Clin Microbiol Rev. 2013;26:744–58, http://dx.doi.org/10.1128/cmr.00023-13.
- Karanika S, Karantanos T, Árvanitis M, Grigoras C, Mylonakis E. Fecal colonization with extended-spectrum beta-lactamase-producing Enterobacterales and risk factors among healthy individuals: a systematic review and metaanalysis. Clin Infect Dis. 2016;63:310–8, http://dx.doi.org/10.1093/cid/ciw283.
 Massart N, Camus C, Benezit F, Moriconi M, Fillatre P, Le Tulzo Y. Incidence and
- Massart N, Camus C, Benezit F, Moriconi M, Fillatre P, Le Tulzo Y. Incidence and risk factors for acquired colonization and infection due to extended-spectrum beta-lactamase-producing Gram-negative bacilli: a retrospective analysis in three ICUs with low multidrug resistance rate. Eur J Clin Microbiol Infect Dis. 2020:39:889–95. http://dx.doi.org/10.1007/s10096-019-03800-y.
- 2020;39:889–95, http://dx.doi.org/10.1007/s10096-019-03800-y.

 6. Valverde A, Coque TM, Sánchez-Moreno MP, Rollán A, Baquero F, Cantón R. Dramatic increase in prevalence of fecal carriage of extended-spectrum beta-lactamase-producing Enterobacterales during nonoutbreak situations in Spain. J Clin Microbiol. 2004;42:4769–75, http://dx.doi.org/10.1128/jcm.42.10.4769-4775.2004.
- Vinué L, Sáenz Y, Martínez S, Somalo S, Moreno MA, Torres C, et al. Prevalence and diversity of extended-spectrum β-lactamases in faecal *Escherichia coli* isolates from healthy humans in Spain. Clin Microbiol Infect. 2009;15:954–7, http://dx.doi.org/10.1111/1469-0691.2009.02803.x.
- Pérez-Moreno MO, Moral-Parras P, Domenech-Spanedda MF, Casacuberta-Barberà N, Bas-García I, Centelles-Serrano MJ, et al. Extended-spectrum β-lactamase- and carbapenemase-producing Enterobacterales intestinal carriage among outpatients: microbiological and epidemiological differences between private dwelling residents and nursing home residents. Microb Drug Resist. 2021;27:879–88, http://dx.doi.org/10.1089/mdr.2020.0201.
- Doi Y, Iovleva A, Bonomo RA. The ecology of extended-spectrum β-lactamases (ESBLs) in the developed world. J Travel Med. 2017;24 Suppl. 1:S44–51, http://dx.doi.org/10.1093/jtm/taw102.
- Mighini-Gras L, Dorado-García A, van Duijkeren E, van den Bunt G, Dierikx CM, Bonten MJM, et al. Attributable sources of community-acquired carriage of *Escherichia coli* containing β-lactam antibiotic resistance genes: a population-based modelling study. Lancet Planet Health. 2019;3:e357–69, http://dx.doi.org/10.1016/S2542-5196(19)30130-5.
- 11. The European Committee on Antimicrobial Susceptibility Testing. Breakpoint tables for interpretation of MICs and zone diameters. Version 9.0, 2019. http://www.eucast.org.
- 12. Briñas L, Zarazaga M, Sáenz Y, Ruiz-Larrea F, Torres C. Beta-lactamases in ampicillin-resistant *Escherichia coli* isolates from foods, humans, and healthy animals. Antimicrob Agents Chemother. 2002;46:3156–63, http://dx.doi.org/10.1128/AAC.46.10.3156-3163.2002.
- 13. Miró E, Mirelis B, Navarro F, Rivera A, Mesa RJ, Roig MC, et al. Surveillance of extended-spectrum beta-lactamases from clinical samples and

- faecal carriers in Barcelona, Spain. J Antimicrob Chemother. 2005;56:1152-5, http://dx.doi.org/10.1093/jac/dki395.
- 14. Castillo García FJ, Seral García C, Pardos De la Gandara M, Millán Lou MI, Pitart Ferré C. Prevalence of fecal carriage of ESBL-producing Enterobacterales in hospitalized and ambulatory patients during two non-outbreak periods. Eur J Clin Microbiol Infect Dis. 2007;26:77–8, http://dx.doi.org/10.1007/s10096-006-0242-8.
- 15. Baquero F, Cercenado E, Cisterna R, de la Rosa M, García-Rodríguez JA, Gobernado M, et al. Patrones de sensibilidad a antimicrobianos de Enterobacterales causantes de infecciones intraabdominales en España: resultados del estudio SMART 2003. Rev Esp Quimioter. 2006;19:51–9.
- 16. Mesa RJ, Blanc V, Blanch AR, Cortés P, González JJ, Lavilla S, et al. Extended-spectrum beta-lactamase-producing Enterobacterales in different environments (humans, food, animal farms and sewage). J Antimicrob Chemother. 2006;58:211–5, http://dx.doi.org/10.1093/jac/dkl211.
- Rodríguez-Baño J, López-Cerero L, Navarro MD, Díaz de Alba P, Pascual A. Faecal carriage of extended-spectrum beta-lactamase-producing *Escherichia coli*: prevalence, risk factors and molecular epidemiology. J Antimicrob Chemother. 2008;62:1142–9, http://dx.doi.org/10.1093/jac/dkn293.
- Porres-Osante N, Sáenz Y, Somalo S, Torres C. Characterization of betalactamases in faecal Enterobacterales recovered from healthy humans in Spain: focusing on AmpC polymorphisms. Microb Ecol. 2015;70:132–40, http://dx.doi.org/10.1007/s00248-014-0544-9.
- Díaz-Agero Pérez C, López-Fresneña N, Rincon Carlavilla AL, Hernandez Garcia M, Ruiz-Garbajosa P, Aranaz-Andrés JM, et al. Local prevalence of extendedspectrum beta-lactamase (ESBL) producing Enterobacterales intestinal carriers at admission and co-expression of ESBL and OXA-48 carbapenemase in Klebsiella pneumoniae: a prevalence survey in a Spanish University Hospital. BMJ Open. 2019;9:e024879, http://dx.doi.org/10.1136/bmjopen-2018-024879. E024879.
- 20. Ríos E, López MC, Rodríguez-Avial I, Culebras E, Picazo JJ. Detection of *Escherichia coli* ST131 clonal complex (ST705) and *Klebsiella pneumoniae* ST15 among faecal carriage of extended-spectrum β-lactamase- and carbapenemase-producing Enterobacterales. J Med Microbiol. 2017;66:169–74, http://dx.doi.org/10.1099/jimm.0.000399.
- van den Bunt G, van Pelt W, Hidalgo L, Scharringa J, de Greeff SC, Schürch AC, et al. Prevalence, risk factors and genetic characterisation of extended-spectrum beta-lactamase and carbapenemase-producing Enterobacterales (ESBL-E and CPE): a community-based cross-sectional study, the Netherlands, 2014–2016. Euro Surveill. 2019;24:1800594, http://dx.doi.org/10.2807/1560-7917.ES. 2019.24.41.1800594.
- Rodríguez-Baño J, Cisneros JM, Cobos-Trigueros N, Fresco G, Navarro-San Francisco C, Gudiol C, et al. Diagnosis and antimicrobial treatment of invasive infections due to multidrug-resistant Enterobacterales. Guidelines of the Spanish Society of Infectious Diseases and Clinical Microbiology. Enferm Infecc Microbiol Clin. 2015;33, http://dx.doi.org/10.1016/j.eimc.2014.11.009, 337.e1-e21.
- de Cueto M, Aliaga L, Alós JI, Canut A, Los-Arcos I, Martínez JA, et al. Executive summary of the diagnosis and treatment of urinary tract infection: guidelines of the Spanish Society of Clinical Microbiology and Infectious Diseases (SEIMC). Enferm Infecc Microbiol Clin. 2017;35:314–20, http://dx.doi.org/10.1016/j.eimc.2016.11.005.
- 24. Horcajada JP, Shaw E, Padilla B, Pintado V, Calbo E, Benito N, et al. Healthcare-associated, community-acquired and hospital-acquired bacteraemic urinary tract infections in hospitalized patients: a prospective multicentre cohort study in the era of antimicrobial resistance. Clin Microbiol Infect. 2013;19:962–8, http://dx.doi.org/10.1111/1469-0691.12089.
- 25. González D, Gallagher E, Zúñiga T, Leiva J, Vitas Al. Prevalence and characterization of β -lactamase-producing Enterobacterales in healthy human carriers. Int Microbiol. 2020;23:171–7, http://dx.doi.org/10.1007/s10123-019-00087-z.
- Manges AR, Geum HM, Guo A, Edens TJ, Fibke CD, Pitout JDD. Global extraintestinal pathogenic *Escherichia coli* (ExPEC) lineages. Clin Microbiol Rev. 2019;32, http://dx.doi.org/10.1128/CMR.00135-18, e00135-18.
- 27. Low M, Neuberger A, Hooton T, Green M, Raz R, Balicer R, et al. Association between urinary community-acquired fluoroquinolone-resistant *Escherichia coli* and neighbourhood antibiotic consumption: a population-based case-control study. Lancet Infect Dis. 2019;19:419–28, http://dx.doi.org/10.1016/S1473-3099(18)30676-5.
- Day MJ, Hopkins KL, Wareham DW, Toleman MA, Elviss N, Randall L, et al. Extended-spectrum B-lactamase-producing Escherichia coli in human-derived and foodchain-derived samples from England, Wales, and Scotland: an epidemiological surveillance and typing study. Lancet Infect Dis. 2019;19:1325–35, http://dx.doi.org/10.1016/S1473-3099(19)30273-7.
- Ben Y, Fu C, Hu M, Liu L, Wong M, Zheng C. Human health risk assessment of antibiotic resistance associated with antibiotic residues in the environment: a review. Environ Res. 2019;169:483–93, http://dx.doi.org/10.1016/j.envres.2018.11.040.
- Yamaguchi T, Okihashi M, Harada K, Konishi Y, Uchida K, Do M, et al. Antibiotic residue monitoring results for pork, chicken, and beef samples in Vietnam in 2012–2013. J Agric Food Chem. 2015;63:5141–5, http://dx.doi.org/10.1021/jf505254y.