**REVIEW ARTICLE** 

# Overview of ESBL-producing *Enterobacteriaceae* from a Nordic perspective

## Alma Brolund, MScPharm, PhD1,2\*

<sup>1</sup>Department of Microbiology, Tumor and Cell Biology, Karolinska Institutet, Stockholm, Sweden;

Extended spectrum β-lactamases (ESBL) are increasing rapidly worldwide. *E. coli* producing CTX-M type ESBLs are the most common clinically encountered. The majority of *E. coli* ESBL infections are represented by urinary tract infections, but they can also cause severe infections, for example, in the blood stream and central nervous system. Since *E. coli* is a common colonizer of the normal gut microbiota, increasing prevalence of ESBL-producing strains is particularly worrisome. Once disseminated in the community, containment of this resistance type will be challenging. The driver of ESBL-producing *Enterobacteriaceae* (EPE) is debated. Some suggest that the ESBL genes were introduced to particularly successful bacterial clones. Others imply that very successful plasmids drive the rapid dissemination. More research and epidemiological studies of strain types, plasmids and mobile genetic elements are needed for these questions to be answered. In order to combat, or at least slow down, the worrisome trend of increasing numbers of EPE more knowledge is also needed on persistence of EPE in carriers as well as better understanding of how antibiotic treatment and other risk factors affect persistence and further dissemination. This review aims at giving an overview of this global problem from a Nordic perspective.

Keywords: antibiotic resistance; extended spectrum  $\beta$ -lactamase; E. coli; K. pneumoniae; epidemiology

Responsible Editor: Lovisa Svensson, Örebro University, Sweden.

\*Correspondence to: Alma Brolund, Public Health Agency of Sweden, SE-17182 Solna, Sweden, Email: alma.brolund@gmail.com

Received: 6 April 2014; Revised: 23 August 2014; Accepted: 6 September 2014; Published: 1 October 2014

Extended spectrum β-lactamases (ESBLs) are enzymes produced by bacteria that degrade β-lactam antibiotics with extended spectrum (such as thirdgeneration cephalosporins) (1). β-Lactams are extensively used for the treatment of common infections, for example, pneumonia and urinary tract infections (UTI), as well as for severe and life threatening infections such as bloodstream infections (BSI). β-Lactams are also frequently used as prophylaxis treatment before surgery.

Different nomenclatures have been proposed and debated for β-lactamases which includes several hundreds of enzymes (http://www.lahey.org/studies/webt.asp). Also what to include in the ESBL definition is debated. In this review, ESBL<sub>A</sub> as defined by Giske et al. will be used (2). The Giske et al. definition divides ESBL enzymes in three main groups; ESBL<sub>A</sub>, ESBL<sub>M</sub> and ESBL<sub>CARBA</sub>. ESBL<sub>A</sub> includes the most frequently found CTX-M enzymes as well as SHV and TEM enzymes that are horizontally transferrable and degraded by clavulanic acid. ESBL<sub>M</sub> are miscellaneous ESBLs where acquired

AmpC represent the most common type.  $ESBL_{CARBA}$  are enzymes conferring carbapenemase activity, such as metallo  $\beta$ -lactamases and KPC (2).

The CTX-Ms were first discovered in the 1980s and since 1995 they have spread to all continents. The numbers have since increased tremendously. The  $bla_{\rm CTX-M}$  genes are divided into five phylogroups; CTX-M-1, CTX-M-2, CTX-M-9 and CTX-M-25 (3, 4). The CTX-M-1 phylogroup includes  $bla_{\rm CTX-M-15}$  which is the most dominant is found globally.

Enterobacteriaceae is the main bacterial family associated to ESBL production, of which *E. coli* and *K. pneumoniae* are most important.

Enterobacteriaceae are of clinical importance causing infections in the central nervous system, lower respiratory tract, bloodstream, gastrointestinal and urinary tract, but are also common colonizers of the gastrointestinal tract (GIT).

E. coli is the main causative agent of community-acquired UTI (5). Women are more affected than men due to anatomical differences.

<sup>&</sup>lt;sup>2</sup>Department of Microbiology, Public Health Agency of Sweden, Solna, Sweden

K. pneumoniae can cause pneumonia, mostly in immunocompromised humans, and BSI. They are the second most common cause of UTI.

K. pneumoniae are to a larger extent acquired in hospital environments than in the community, for example, in patients with urinary catheters or immunocompromised patients (6).

As the ESBL-carrier rate in the population increases and nosocomial transmission becomes more common, the risk of being infected increases. ESBL-producing *Enterobacteriaceae* (EPE) often displays multidrug-resistant phenotypes further limiting the therapeutic options. Coresistance with fluoroquinolones, aminoglycosides and trimethoprim are often found (7) which are common empirical treatment options for severely ill patients resulting in increased morbidity and mortality.

ESBLs also constitute a burden on health care systems conferring prolonged hospital stay. de Kraker and coworkers describe BSIs in Europe caused by *E. coli* isolates resistant to extended spectrum cephalosporins. They estimate the cost to >2,700 excess deaths and 18.1 million EUR, represented by >120,000 excess days of hospital stay (8).

Patients with severe infections are highly dependent on receiving adequate treatment early in the course of infection for a benign outcome. Many times the diagnostic procedures available are too slow and empirical treatment necessary. Knowledge of the local epidemiology is therefore very important. For a patient with BSI caused by resistant bacteria, such as an ESBL-producer, administration of an ineffective antibiotic can be lethal. Changing to an effective regimen after treatment failure might be too late (9). Peralta and coworkers studied the rate of adequate empirical treatment in BSIs caused by Enterobacteriaceae in 19 Spanish hospitals during 4 years. They found the empirical treatment to be inadequate in 48.8% of the cases. Twenty-four of the 387 patients included in the study died during the first 3 days of inadequate therapy (10).

National and global epidemiology is also important since it provides ways to detect emerging clones or new types of resistant bacteria. A wider perspective can bring valuable information to the practitioners treating patients associated to other parts of the country or the world. Governmental institutions and agencies play an important role gathering data through different surveillance tools.

## **Epidemiology of CTX-M-genes**

Different ESBL genotypes have different substrate specificities for the many  $\beta$ -lactam antibiotics (11), also between CTX-M enzymes. As mentioned  $bla_{\text{CTX-Ms}}$  are found all over the world and  $bla_{\text{CTX-M-15}}$  is the most dominating genotype (12). However, local variations occur. In the Nordic countries, the distribution of

genotypes is homogenous. In Sweden, 50-60% of ESBL-producing *E. coli* harbor  $bla_{\text{CTX-M-15}}$  followed by  $bla_{\text{CTX-M-14}}$  found in 10-15% (7). In Norway and Denmark, about 50% and 20% of the genotypes are  $bla_{\text{CTX-M-15}}$  and  $bla_{\text{CTX-M-14}}$  respectively (13, 14). The ESBL epidemiology in Finland has been sparsely reported, but a predominance of the CTX-M-1 group (including  $bla_{\text{CTX-M-15}}$ ) is found (15, 16).

In other parts of Europe, accumulation of different genotypes is found. In Spain, there is a high level of  $bla_{\text{CTX-M-14}}$  and  $bla_{\text{CTX-M-9}}$ , and in Poland as well as several other eastern European countries  $bla_{\text{CTX-M-3}}$  has been reported as frequent (17, 18).

In the African and Australian continents,  $bla_{\text{CTX-M-15}}$  is dominating (19). In Canada,  $bla_{\text{CTX-M-15}}$  is also the most frequent and increasingly so in the United States of America, where  $bla_{\text{SHV}}$ -phenotypes used to be the most encountered (20). Differences are seen between Asian countries. In Japan, the CTX-M-9 phylogroup is largely dominating (21) and also in China  $bla_{\text{CTX-M-14}}$  is more commonly reported (22). In India,  $bla_{\text{CTX-M-15}}$  is almost exclusively found (23). South America stands out with a wide distribution of the CTX-M 2 phylogroup. Also the CTX-M-8 phylogroup is found in South America which is rare on other continents (12, 19).

## Dissemination of ESBLs

Transmission of genes encoding ESBL enzymes can occur either by emerging bacterial clones or by horizontal gene transfer. In the latter case, plasmids containing resistance genes are spread between bacteria of the same and/or different species.

The gut flora is an ideal reservoir for antibiotic resistance genes, where kilograms of bacteria of different species can interact, most often without causing disease and hence to much interference with the immune system. When antibiotics are used resistant strains gain a selective advantage and are accumulated. This increases the probability for genes important for survival to be further disseminated. Plasmids can contain resistance genes to several classes of antibiotics (24). This can explain the multiresistance often found amongst ESBL-producing bacteria.

## Clonal expansion

Clonal expansion refers to a particular bacterial cell line multiplying and disseminating in a community or causing an outbreak, for example, in a hospital. Rapid detection of clonal expansions is important in order to be able to stop or diminish the source of dissemination. Hence, effective molecular epidemiological strain typing tools are crucial.

An example of clonal expansion is the *E. coli* sequence type 131 (ST131) that is found globally (25). ST131 is highly associated to *bla<sub>CTX-M-15</sub>* (25). In the Nordic countries,

ST131 represents 20–40% of the ESBL-producing E. coli (13, 14, 26). However, ST131 is also found in non-ESBL E. coli, suggesting that ST131 is a dominant E. coli type that has acquired ESBLs which may have contributed to the successful dissemination of these resistance genes (25). Successful STs of K. pneumoniae are also described. An example is the STs of clonal complex 11 (CC11) which have been associated to CTX-M dissemination (4, 27).

Outbreaks caused by EPE have occurred in the Nordic countries. The largest ones occurred in Uppsala, Sweden and Stavanger, Norway. The Uppsala outbreak, in 2005, was due to a bla<sub>CTX-M-15</sub> producing K. pneumoniae that was spread over several ward departments of the University Hospital. In total, 64 patients were affected (28). Another large outbreak of  $bla_{CTX-M-15}$  producing K. pneumoniae occurred in a neonatal ward department of Stavanger University Hospital in 2008-2009. Fifty-eight infants were infected, probably for receiving contaminated breast milk (29).

Although ST131 represent a large fraction of the ESBL-producing E. coli population, still  $\sim 50\%$  of Swedish isolates are genetically unrelated isolates. This indicates that horizontal gene transfer is a major route for disseminating ESBLs and that these resistance genes have a remarkable ability to enter E. coli of many different back-bones (7).

## Horizontal dissemination

Plasmids are double-stranded extra-chromosomal DNA that replicate independently of the chromosome in the bacterial cell hosting it. Plasmids are highly diverse and the plasmid genome is often scattered with mobile genetic elements that can move genes around within the plasmid as well as between chromosome and plasmid(s). Examples of mobile genetic elements found on plasmids are ISelements, transposons, integrons and insertion sequence common regions (ISCR), of which the latter is associated to ESBL genes. Plasmids have developed ways to secure their persistence in the bacterial cell. One example is through toxin-antitoxin systems that act by eliminating cells that have lost the plasmid after cell division (24). Figure 1 pictures a schematic illustration of plasmid conjugation which is a common route for plasmid dissemination.

Plasmids can be classified according to incompatibility (Inc) groups, which are based on the principle that plasmids with the same replicon cannot be propagated stably in the same cell (30). The explanation for this is that similar plasmids compete over common cellular functions involved in, for example, plasmid replication control (31).

A replicon is a highly conserved part of the plasmid where genes encoding replication initiation, control, copy number, etc. are situated (illustrated as the plasmid core genome in Figure 2). Nowadays incompatibility grouping have been replaced by replicon typing which is simpler

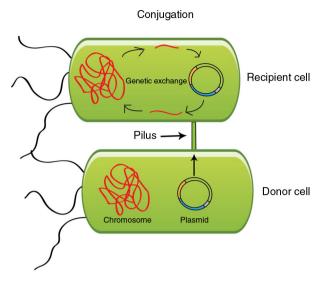


Fig. 1. Illustration of bacterial conjugation.

experimentally and also allows for a subdivision of some of the Inc groups. One example is the IncF1 plasmids that include repFIA, FIB and FIC of which repFIA and FIB plasmids are commonly found to harbor ESBL genes.

The host range of plasmids is largely determined by the Inc type. IncF plasmids are large narrow host range plasmids with low copy number. They are the most common Inc type associated to CTX-M ESBLs. IncA/C plasmids, found associated to ESBL<sub>M</sub> and ESBL<sub>CARBA</sub>, are contrarily broad host range plasmids that are found in a wide variety of bacterial species. Some Inc groups are considered as endemic resistance plasmids in Enterobacteriaceae. These are IncF, IncA/C, IncL/M, IncN and IncI (32).

Among fully sequenced E. coli plasmids harboring bla<sub>CTX-M</sub>-genes in GenBank, the following plasmid families are found: IncF, IncN, IncN2, IncI1, IncHI2, IncL/M, IncA/C, IncK, IncX4, IncU and RCR (33).

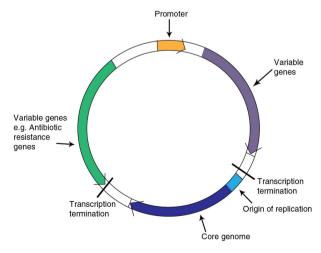


Fig. 2. A schematic illustration of a plasmid.

Plasmid variability within incompatibility groups is high. In Norwegian and Swedish, ESBL-producing E. coli IncF plasmids are almost exclusively found (13, 34). Subtyping by replicon sequence typing RST) however shows a great variability within F-plasmids, and it is not possible to link a specific IncF-plasmid to the successful dissemination of bla<sub>CTX-Ms</sub> (35). IncF plasmids are found also in antibiotic-sensitive Enterobacteriaceae suggesting that the ESBL genes have entered an already wellestablished plasmid type which has been beneficial for their dissemination (36).

## **Epidemiology of EPE**

The Nordic countries are considered low prevalence countries, but the EPE numbers are increasing fast. In Sweden, national surveillance has been performed since 2007 when ESBLs were included in the communicable disease act. The number of cases have increased from 2099 in 2007 (Feb-Dec) to 8131 in 2013.

Several studies pictures the local frequency in different parts of Sweden; Helldal et al. described the prevalence of ESBL-producing E. coli in South Western Sweden during 2004-2008 with a higher increase in nosocomially acquired isolates in comparison to community-acquired, 0.2-2.5\% and 0.2-1.6\%, respectively (37). A study from the county of Östergötland indicated an increased prevalence of EPE from 2002 to 2007, yet the numbers were very low (<1%) (38). An investigation of the fecal carriage of ESBL-producing bacteria in both Primary Health Care Units and University Hospital setting in southern Sweden was conducted in 2008-2010. The only species found was E. coli, and the authors described the prevalence as higher than expected in both groups, 2.1-3\% and 1.8-6.8\%, respectively (39). A recent study from Uppsala investigated the prevalence of EPE in healthy preschool children compared to patients in the same age group. The authors found a carrier rate of 2.9% in healthy children compared to 8.4% in patients of the corresponding age group (40). In Helsinki, Finland, the annual incidence rate of EPE has gone from 0.5/100 000 inhabitants in the year 2000 to 69/ 100 000 in 2004 (15). In Denmark, the rate of invasive EPE has increased from 2.5% in 2006 to 6.2% in 2009 (41). The annual report of antibiotic resistance isolates in Norway, Norm/Norm-Vet, published in 2012 states that EPE is increasing steadily, and the number of EPE in blood cultures has increased from 3.3% in 2011 to 5.5% in 2012 (42).

In Europe, the prevalence of bacteria resistant to extended spectrum cephalosporins differs a lot between countries. They are less frequent in the north and more common in the southern and eastern parts of Europe. The European network of national surveillance systems of antimicrobial resistance summarizes the levels of invasive infections caused by resistant bacteria in the European countries in a yearly report available on the ECDC homepage: http://ecdc.europa.eu/. The results are also illustrated on color-coded maps. In Fig. 3, levels of invasive E. coli infections resistant to third-generation cephalosporins are illustrated over time. However, one must keep in mind that the results are biased by differences in reporting, national monitoring and sampling.

Looking at countries outside Europe the ESBL prevalence also varies a lot. A recent study from the USA presents results from the SENTRY surveillance program including 26 hospitals from 20 states. The resistance levels to cephalosporins and/or aztreonam in invasive Enterobacteriaceae was 6.4% (43).

A Canadian study reported ESBL-production in almost 5% of the E. coli population – numbers similar to northern Europe (44).

The level of resistant bacteria is truly troublesome in other parts of the world. In Thailand, carriage of ESBLproducing bacteria, in as high numbers as 52% of healthy volunteers, has been reported (45). In India, the reports are also alarming with high numbers seen among patients. A study from two hospitals in Ujjain, India, collected consecutive isolates from incoming patients with abscesses, UTI, BSI, etc. and found ESBL production in 69% of E. coli (46).

One factor contributing to the low EPE numbers in the Nordic countries is the restricted antibiotic treatment policies. Antibiotics can only be purchased through prescriptions. But despite the strict antibiotic policy and decreasing antibiotic consumption, EPE is still increasing (47). It has been shown that one entry of EPE to low prevalence countries is through traveling to countries with higher prevalence. In 2010, Tängden et al. studied the acquisition of resistance genes as a result of travel outside northern Europe. Healthy travelers voluntarily participated in taking fecal samples before and after travel. The study resulted in 24% returning as new carriers of EPE after travel (48). Patients admitted to an Infectious Disease department in Aarhus, Denmark, who had been abroad within 3 months were identified and subjected to screening for multidrug resistant bacteria. EPE colonization was found in 12.5% of which > 80%had been abroad for longer than 2 weeks (49).

Many countries outside Scandinavia (e.g. the member states of the European Union) also have antibiotic prescribing policies where prescription is required for antibiotics to be purchased. Other countries lack such regulations and the implementation of the policies differs between countries. Acquisition of antibiotics 'over the counter' in pharmacies or equivalents still occurs in many places. There are also differences in prescribing tradition and culture between countries. The climate affects the frequency of infections and also the rate of dissemination. Economical status, household conditions and population density also clearly play a role.

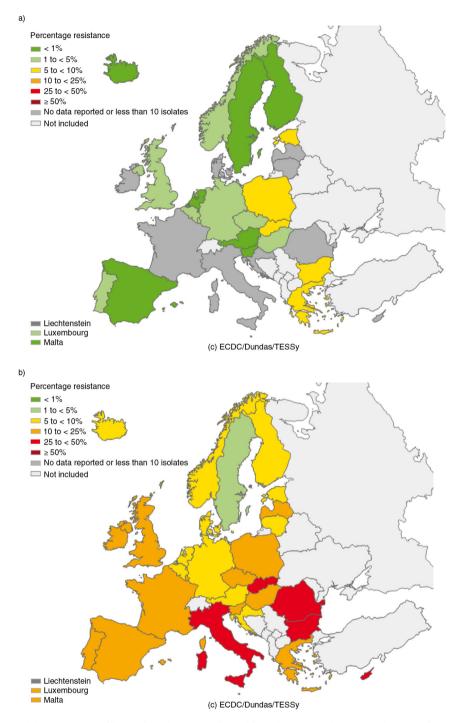


Fig. 3. Maps generated by EARS-Net illustrating the proportion of invasive *E. coli* isolates resistant or intermediately resistant to third-generation cephalosporins from reporting countries in a) 2001 and b) 2012. Figure 3 is reprinted with permission from ECDC. Source: ECDC/EARS-Net 2001 and 2012.

Infection control routines and programs have a large impact on dissemination of bacteria (50–52). Differences in infection control routines are substantial between different countries, also between European countries. Furthermore, the microbiological diagnostic traditions and setup of surveillance programs differ. If awareness of the regional situation is low, preventive measures will also

be. More and more studies show that there is dissemination of bacteria between food, animal (both food-production animals and pets) and humans (53–56). Consequently antibiotic usage and inadequate infection control in veterinary medicine, fish farming, irrigation, etc. also contribute to the increased spread of antibiotic resistance.

### Persistence of ESBLs

Little is known about the persistence of ESBL-producing bacteria in patients. This is mainly due to difficulties in studying persistence. Fecal cultures represent a snapshot of the flora at the time of study, and low numbers of bacteria can be difficult to detect. It has been shown that a person can present several negative samples and then later on still come up with a positive one (57). Another difficulty is that negative samples prior to colonization or confirmed infection seldom exist.

There are however some studies on the persistence of EPE. Most are conducted after detecting nosocomial outbreaks. One example is Löhr et al. who have followed children and their family members for 3 years after an outbreak in the neonatal intensive care unit in Stavanger, Norway. The authors observed carrier length of up to 2 years in some of the children (58). Persistence in neonates is proposed to be longer than in older children and adults since they have not yet established a naturally protective microbial flora. They also have an undeveloped immune system and are more often treated with antibiotics (58).

Alsterlund et al. studied EPE fecal colonization after a hospital outbreak in southern Sweden and found some individuals to still be colonized after 5 years (57). Another Swedish study by Tham et al. describes persistence of ESBL in 10% of patients after 3 years. The authors detected two new ESBL-producing strain types in one of the patients indicating transfer of the resistance phenotype (59). An alternative interpretation would be that this patient was infected with additional strains, although given the low frequency of EPE in Sweden the probability seems low. Low risk of re-colonization is beneficial when studying colonization rates, since the probability of finding the true colonization rate, duration of carriage and migration of plasmids between species increases. Titelman et al. studied persistence of EPE in 61 patients over the course of 1 year. By the end of the study period, 43% of the patients were still colonized (60). This study resulted in different scenarios in different patients. Some of the patients presented EPE of different species and molecular strain types carrying the same resistance gene over time picturing the interaction between the bacteria in the GIT. Colonization in other patients in the same study, however, presented one identical strain type with the same resistance gene over time (60). The factors behind these differences would be highly interesting to identify.

Dissemination and persistence of ESBL-producing bacteria have also been studied within households. A Spanish study reported that 16.7% of people living together with persons with known community-acquired infections also were colonized (61). Another Spanish study found fecal carriage in 27.4% of household members of known carriers. Relatives outside the household had a

15.6% carrier rate. These numbers can be compared with a control group of unrelated persons where the carrier rate was 7.4% (62). A Chinese study investigated ESBL carriage in children 0-5 years of age admitted to a pediatric clinic in Hong Kong and found CTX-M producing bacteria in 43.5%. In 83% of these children, at least one of the household members was also a carrier (63). A French study found intra-family transmission from adoptive children to their new family members at a rate of 23% (64).

A different way of studying persistence is through plasmid dissemination. Cottell et al. investigated a plasmid, pCT, harboring bla<sub>CTX-M-14</sub> (the only resistance gene on that plasmid). They transformed the plasmid into an E. coli strain and found that it provided little or no additional fitness to the bacteria, but still the plasmid persisted successfully in the absence of selective antibiotic pressure. This suggests that persistence of the plasmid is not always dependent on the resistance gene (65).

The studies mentioned show that ESBLs can persist in carriers for several years. They are shared within households and disseminated in the community. More research is however needed to understand how persistence and dissemination can be minimized. Dissemination through plasmids adds a second dimension to this problem, and most probably prolongs the ability for resistance phenotypes to persist.

## Conclusions

The impact of antibiotic treatment on the dissemination and persistence of resistant bacteria cannot be underestimated. In order to use these extremely important drugs as rational as possible, we need to gain full understanding of how resistance arises and all the factors that influence dissemination. Optimized antibiotic treatment is in turn dependent on the development of faster diagnostic tools in order to minimize empirical treatment.

Much research is still needed in this field. Strain and plasmid interaction and dissemination needs further attention in order to be fully understood. The persistence of EPE in the healthy population is also important to learn more about. Such knowledge would contribute to important information regarding how to handle patients colonized with EPE in hospitals and for how long persistence is to be expected under different conditions. It could also give us insight in how to optimize antibiotic treatment to be as effective as possible for patients and yet minimize further antibiotic resistance from arising and disseminating.

For antibiotics to be potent also in the future, promoting rational antibiotic use in all sectors minimizes the selective pressure for resistant bacteria. New antibiotic agents are also needed as well as optimizing the usage of the already available antibiotics. New treatment options can buy us time to address this important issue that constitutes a major public health problem.

## Acknowledgements

This review is based on the doctoral thesis 'Plasmid mediated antibiotic resistance – with focus on extended spectrum β-lactamases (ESBL),' by Alma Brolund, 2013, published by Karolinska Institutet, ISBN 978-91-7549-210-0.

I gratefully acknowledge Karin Tegmark Wisell, Christian G Giske and Öjar Meleford for supervision during my research education. A special thanks also to countries providing data for EARS-Net at ECDC.

## Conflict of interest and funding

The author has no conflict of interest.

#### References

- 1. Paterson DL, Bonomo RA. Extended-spectrum beta-lactamases: a clinical update. Clin Microbiol Rev 2005; 18: 657-86.
- 2. Giske CG, Sundsfjord AS, Kahlmeter G, Woodford N, Nordmann P, Paterson DL, et al. Redefining extended-spectrum beta-lactamases: balancing science and clinical need. J Antimicrob Chemother 2009; 63: 1-4.
- 3. Birkett CI, Ludlam HA, Woodford N, Brown DF, Brown NM, Roberts MT, et al. Real-time TaqMan PCR for rapid detection and typing of genes encoding CTX-M extended-spectrum betalactamases. J Med Microbiol 2007; 56: 52-5.
- 4. D'Andrea MM, Arena F, Pallecchi L, Rossolini GM. CTX-Mtype beta-lactamases: a successful story of antibiotic resistance. Int J Med Microbiol 2013; 303: 305-17.
- 5. Abraham S, Chapman TA, Zhang R, Chin J, Mabbett AN, Totsika M, et al. Molecular characterization of Escherichia coli strains that cause symptomatic and asymptomatic urinary tract infections. J Clin Microbiol 2012; 50: 1027-30.
- 6. Madigan MT, Martinko JM, Parker J. Brock biology of microorganisms. 10th ed. NJ, USA: Pearson Education, Inc., Upper Saddle iver; 2003.
- 7. Brolund A, Edquist PJ, Makitalo B, Olsson-Liljequist B, Soderblom T, Wisell KT, et al. Epidemiology of extendedspectrum beta-lactamase-producing Escherichia coli in Sweden 2007-2011. Clin Microbiol Infect 2014; 20: O344-52.
- 8. de Kraker ME, Davey PG, Grundmann H, BURDEN study group. Mortality and hospital stay associated with resistant Staphylococcus aureus and Escherichia coli bacteremia: estimating the burden of antibiotic resistance in Europe. PLoS Med 2011; 8: e1001104.
- 9. Giske CG, Monnet DL, Cars O, Carmeli Y, ReAct-Action on Antibiotic Resistance. Clinical and economic impact of common multidrug-resistant gram-negative bacilli. Antimicrob Agents Chemother 2008; 52: 813-21.
- 10. Peralta G, Lamelo M, Alvarez-Garcia P, Velasco M, Delgado A, Horcajada JP, et al. Impact of empirical treatment in extended-spectrum beta-lactamase-producing Escherichia coli and Klebsiella spp. bacteremia. A multicentric cohort study. BMC Infect Dis 2012; 12: 245.
- 11. Bonnet R. Growing group of extended-spectrum beta-lactamases: the CTX-M enzymes. Antimicrob Agents Chemother 2004; 48: 1-14.
- 12. Canton R, Coque TM. The CTX-M beta-lactamase pandemic. Curr Opin Microbiol 2006; 9: 466-75.
- 13. Naseer U, Haldorsen B, Tofteland S, Hegstad K, Scheutz F, Simonsen GS, et al. Molecular characterization of CTX-M-15producing clinical isolates of Escherichia coli reveals the spread of multidrug-resistant ST131 (O25:H4) and ST964 (O102:H6) strains in Norway. APMIS 2009; 117: 526-36.

- 14. Olesen B, Hansen DS, Nilsson F, Frimodt-Moller J, Leihof RF, Struve C, et al. Prevalence and characteristics of the epidemic multiresistant Escherichia coli ST131 clonal group among extended-spectrum beta-lactamase-producing E. coli isolates in Copenhagen, Denmark. J Clin Microbiol 2013; 51: 1779-85.
- 15. Forssten SD, Kolho E, Lauhio A, Lehtola L, Mero S, Oksaharju A, et al. Emergence of extended-spectrum betalactamase-producing Escherichia coli and Klebsiella pneumoniae during the years 2000 and 2004 in Helsinki, Finland. Clin Microbiol Infect 2010; 16: 1158-61.
- 16. Nyberg SD, Osterblad M, Hakanen AJ, Huovinen P, Jalava J, Resistance TF. Detection and molecular genetics of extendedspectrum beta-lactamases among cefuroxime-resistant Escherichia coli and Klebsiella spp. isolates from Finland, 2002-2004. Scand J Infect Dis 2007; 39: 417-24.
- 17. Baraniak A, Fiett J, Sulikowska A, Hryniewicz W, Gniadkowski M. Countrywide spread of CTX-M-3 extendedspectrum beta-lactamase-producing microorganisms of the family Enterobacteriaceae in Poland. Antimicrob Agents Chemother 2002; 46: 151-9.
- 18. Livermore DM, Canton R, Gniadkowski M, Nordmann P, Rossolini GM, Arlet G, et al. CTX-M: changing the face of ESBLs in Europe. J Antimicrob Chemother 2007; 59: 165-74.
- 19. Naseer U, Sundsfjord A. The CTX-M conundrum: dissemination of plasmids and Escherichia coli clones. Microb Drug Resist 2011; 17: 83-97.
- 20. Doi Y, Park YS, Rivera JI, Adams-Haduch JM, Hingwe A, Sordillo EM, et al. Community-associated extended-spectrum beta-lactamase-producing Escherichia coli infection in the United States. Clin Infect Dis 2013; 56: 641-8.
- 21. Chong Y, Shimoda S, Yakushiji H, Ito Y, Miyamoto T, Kamimura T, et al. Community spread of extended-spectrum beta-lactamase-producing Escherichia coli, Klebsiella pneumoniae and Proteus mirabilis: a long-term study in Japan. J Med Microbiol 2013: 62: 1038-43.
- 22. Xia S, Fan X, Huang Z, Xia L, Xiao M, Chen R, et al. Dominance of CTX-M-type extended-spectrum beta-lactamase (ESBL)-producing Escherichia coli isolated from patients with community-onset and hospital-onset infection in China. PLoS One 2014; 9: e100707.
- 23. Muzaheed, Doi Y, Adams-Haduch JM, Endimiani A, Sidjabat HE, Gaddad SM, et al. High prevalence of CTX-M-15-producing Klebsiella pneumoniae among inpatients and outpatients with urinary tract infection in Southern India. J Antimicrob Chemother 2008; 61: 1393-4.
- 24. Bennett PM. Plasmid encoded antibiotic resistance: acquisition and transfer of antibiotic resistance genes in bacteria. Br J Pharmacol 2008; 153: S347-57.
- 25. Nicolas-Chanoine MH, Bertrand X, Madec JY. Escherichia coli ST131, an intriguing clonal group. Clin Microbiol Rev 2014; 27:
- 26. Brolund A, Edquist PJ, Makitalo B, Olsson-Liljequist B, Soderblom T, Tegmark Wisell K, et al. Epidemiology of ESBL-producing Escherichia coli in Sweden 2007-2011. Clin Microbiol Infect, 2013; 20(6): O344-52.
- 27. Andrade LN, Vitali L, Gaspar GG, Bellissimo-Rodrigues F, Martinez R, Darini AL. Expansion and evolution of a virulent, extensively drug-resistant (polymyxin B-resistant), QnrS1-, CTX-M-2-, and KPC-2-producing Klebsiella pneumoniae ST11 international high-risk clone. J Clin Microbiol 2014; 52: 2530-5.
- 28. Lytsy B, Sandegren L, Tano E, Torell E, Andersson DI, Melhus A. The first major extended-spectrum beta-lactamase outbreak in Scandinavia was caused by clonal spread of a multiresistant Klebsiella pneumoniae producing CTX-M-15. APMIS 2008; 116: 302-8.

- 29. Rettedal S, Lohr IH, Natas O, Giske CG, Sundsfjord A, Oymar K. First outbreak of extended-spectrum beta-lactamaseproducing Klebsiella pneumoniae in a Norwegian neonatal intensive care unit; associated with contaminated breast milk and resolved by strict cohorting. APMIS 2012; 120: 612-21.
- 30. Novick RP. Plasmid incompatibility. Microbiol Rev 1987; 51: 381-95.
- 31. Couturier M, Bex F, Bergquist PL, Maas WK. Identification and classification of bacterial plasmids. Microbiol Rev 1988; 52:
- 32. Carattoli A. Plasmids in Gram negatives: molecular typing of resistance plasmids. Int J Med Microbiol 2011; 301: 654-8.
- 33. Carattoli A. Plasmids and the spread of resistance. Int J Med Microbiol 2013: 303: 298-304.
- 34. Brolund A, Franzen O, Melefors O, Tegmark-Wisell K, Sandegren L. Plasmidome-analysis of ESBL-producing Escherichia coli using conventional typing and high-throughput sequencing. PLoS One 2013; 8: e65793.
- 35. Villa L, Garcia-Fernandez A, Fortini D, Carattoli A. Replicon sequence typing of IncF plasmids carrying virulence and resistance determinants. J Antimicrob Chemother 2010: 65:
- 36. Bengtsson S, Naseer U, Sundsfjord A, Kahlmeter G, Sundqvist M. Sequence types and plasmid carriage of uropathogenic Escherichia coli devoid of phenotypically detectable resistance. J Antimicrob Chemother 2012; 67: 69-73.
- 37. Helldal L, Karami N, Floren K, Welinder-Olsson C, Moore ER, Ahren C. Shift of CTX-M genotypes has determined the increased prevalence of extended-spectrum beta-lactamaseproducing Escherichia coli in south-western Sweden. Clin Microbiol Infect 2013; 19: E87-90.
- 38. Ostholm-Balkhed A, Tarnberg M, Nilsson M, Johansson AV, Hanberger H, Monstein HJ, et al. Prevalence of extendedspectrum beta-lactamase-producing Enterobacteriaceae and trends in antibiotic consumption in a county of Sweden. Scand J Infect Dis 2010; 42: 831-8.
- 39. Stromdahl H, Tham J, Melander E, Walder M, Edquist PJ, Odenholt I. Prevalence of faecal ESBL carriage in the community and in a hospital setting in a county of Southern Sweden. Eur J Clin Microbiol Infect Dis 2011; 30: 1159-62.
- 40. Kaarme J, Molin Y, Olsen B, Melhus A. Prevalence of extendedspectrum beta-lactamase-producing Enterobacteriaceae in healthy Swedish preschool children. Acta Paediatr 2013; 102: 655-60.
- 41. Hansen F, Olsen SS, Heltberg O, Justesen US, Fuglsang-Damgaard D, Knudsen JD, et al. Characterization of thirdgeneration cephalosporin-resistant Escherichia coli from bloodstream infections in Denmark. Microb Drug Resist 2014: 20: 316-24.
- 42. NORM/NORM-VET 2012. Usage of antimicrobial agents and occurrence of antimicrobial resistance in Norway. Tromsø/Oslo 2013.
- 43. Castanheira M, Farrell SE, Deshpande LM, Mendes RE, Jones RN. Prevalence of beta-lactamase-encoding genes among Enterobacteriaceae bacteremia isolates collected in 26 U.S. hospitals: report from the SENTRY Antimicrobial Surveillance Program (2010). Antimicrob Agents Chemother 2013; 57: 3012 - 20.
- 44. Simner PJ, Zhanel GG, Pitout J, Tailor F, McCracken M, Mulvey MR, et al. Prevalence and characterization of extendedspectrum beta-lactamase- and AmpC beta-lactamase-producing Escherichia coli: results of the CANWARD 2007-2009 study. Diagn Microbiol Infect Dis 2011; 69: 326-34.

- 45. Oteo J, Perez-Vazquez M, Campos J. Extended-spectrum [beta]lactamase producing Escherichia coli: changing epidemiology and clinical impact. Curr Opin Infect Dis 2010; 23: 320-6.
- 46. Pathak A, Marothi Y, Kekre V, Mahadik K, Macaden R, Lundborg CS. High prevalence of extended-spectrum betalactamase-producing pathogens: results of a surveillance study in two hospitals in Ujjain, India. Infect Drug Resist 2012; 5: 65-73
- 47. SWEDRES-SVARM 2012. Use of antimicrobials and occurrence of antimicrobial resistance in Sweden. 2012.
- 48. Tangden T, Cars O, Melhus A, Lowdin E. 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. Antimicrob Agents Chemother 2010: 54: 3564-8.
- 49. Lausch KR, Fuursted K, Larsen CS, Storgaard M. Colonisation with multi-resistant Enterobacteriaceae in hospitalised Danish patients with a history of recent travel: a cross-sectional study. Travel Med Infect Dis 2013; 11: 320-3.
- 50. March A, Aschbacher R, Pagani E, Sleghel F, Soelva G, Hopkins KL, et al. Changes in colonization of residents and staff of a long-term care facility and an adjacent acute-care hospital geriatric unit by multidrug-resistant bacteria over a four-year period. Scand J Infect Dis 2014; 46: 114-22.
- 51. Jadhav S, Sahasrabudhe T, Kalley V, Gandham N. The microbial colonization profile of respiratory devices and the significance of the role of disinfection: a blinded study. J Clin Diagn Res 2013; 7: 1021-6.
- 52. Deege MP, Paterson DL. Reducing the development of antibiotic resistance in critical care units. Curr Pharm Biotechnol 2011; 12: 2062-9.
- 53. Trott D. β-Lactam resistance in gram-negative pathogens isolated from animals. Curr Pharm Des 2013; 19: 239-49.
- 54. Reuland EA, Al Naiemi N, Raadsen SA, Savelkoul PH, Kluytmans JA, Vandenbroucke-Grauls CM. Prevalence of ESBL-producing Enterobacteriaceae in raw vegetables. Eur J Clin Microbiol Infect Dis 2014; [Epub ahead of print].
- 55. Egervarn M, Borjesson S, Byfors S, Finn M, Kaipe C, Englund S, et al. Escherichia coli with extended-spectrum beta-lactamases or transferable AmpC beta-lactamases and Salmonella on meat imported into Sweden. Int J Food Microbiol 2014; 171: 8-14.
- 56. Tschudin-Sutter S, Frei R, Stephan R, Hachler H, Nogarth D, Widmer AF. Extended-spectrum beta-lactamase (ESBL)producing Enterobacteriaceae: a threat from the kitchen. Infect Control Hosp Epidemiol 2014; 35: 581-4.
- 57. Alsterlund R, Axelsson C, Olsson-Liljequist B. Long-term carriage of extended-spectrum beta-lactamase-producing Escherichia coli. Scand J Infect Dis 2012; 44: 51-4.
- 58. Lohr IH, Rettedal S, Natas OB, Naseer U, Oymar K, Sundsfjord A. Long-term faecal carriage in infants and intrahousehold transmission of CTX-M-15-producing Klebsiella pneumoniae following a nosocomial outbreak. J Antimicrob Chemother 2013; 68: 1043-8.
- 59. Tham J, Walder M, Melander E, Odenholt I. Duration of colonization with extended-spectrum beta-lactamase-producing Escherichia coli in patients with travellers' diarrhoea. Scand J Infect Dis 2012; 44: 573-7.
- 60. Titelman E, Hasan CM, Iversen A, Naucler P, Kais M, Kalin M, et al. Faecal carriage of extended-spectrum betalactamase-producing Enterobacteriaceae is common 12 months after infection and is related to strain factors. Clin Microbiol Infect 2014; [Epub ahead of print].

- 61. Valverde A, Grill F, Coque TM, Pintado V, Baquero F, Canton R, et al. High rate of intestinal colonization with extended-spectrum-beta-lactamase-producing organisms in household contacts of infected community patients. J Clin Microbiol 2008; 46: 2796-9.
- 62. Rodriguez-Bano J, Lopez-Cerero L, Navarro MD, Diaz de Alba P, Pascual A. Faecal carriage of extended-spectrum betalactamase-producing Escherichia coli: prevalence, risk factors and molecular epidemiology. J Antimicrob Chemother 2008; 62: 1142-9.
- 63. Lo WU, Ho PL, Chow KH, Lai EL, Yeung F, Chiu SS. Fecal carriage of CTXM type extended-spectrum beta-lactamase-

- producing organisms by children and their household contacts. J Infect 2010: 60: 286-92.
- 64. Tande D, Boisrame-Gastrin S, Munck MR, Hery-Arnaud G, Gouriou S, Jallot N, et al. Intrafamilial transmission of extendedspectrum-beta-lactamase-producing Escherichia coli and Salmonella enterica Babelsberg among the families of internationally adopted children. J Antimicrob Chemother 2010; 65: 859-65.
- 65. Cottell JL, Webber MA, Piddock LJ. Persistence of transferable extended-spectrum-beta-lactamase resistance in the absence of antibiotic pressure. Antimicrob Agents Chemother 2012; 56: 4703-6.