



Karolinska  
Institutet

## Is AmpC-producing *Escherichia coli* an increasing problem?

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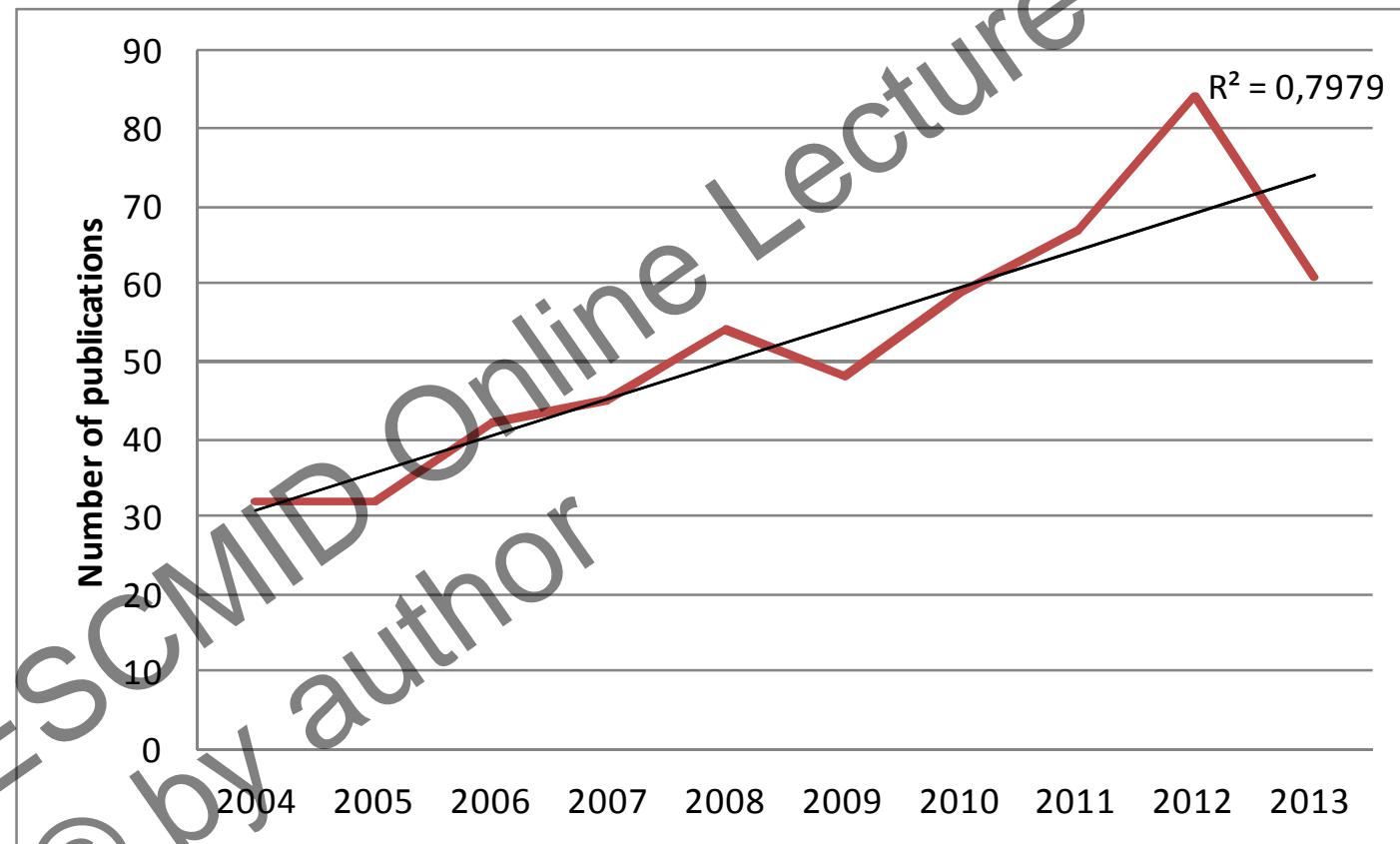
Clinical microbiology

Karolinska University Hospital

Barcelona, 21 November 2013

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## PubMed: AmpC AND plasmid

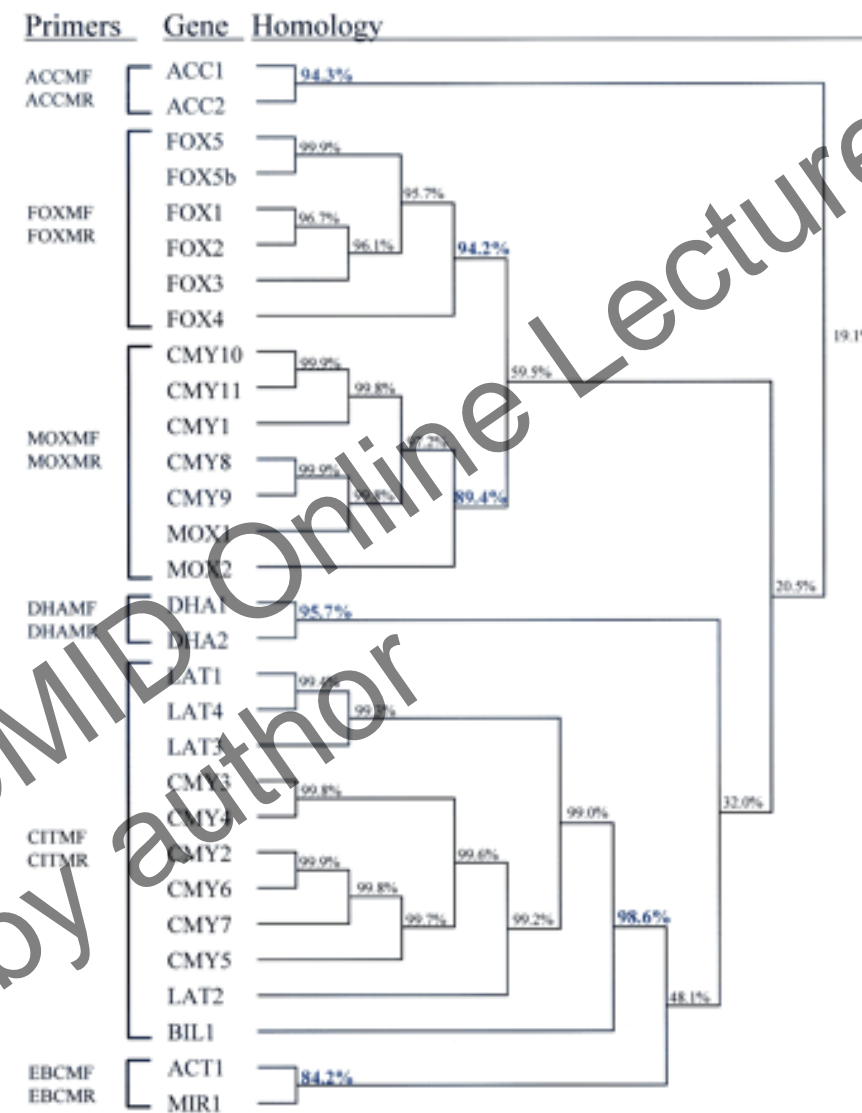


**Introductory remarks on plasmid-mediated  
AmpC (pAmpC)**

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## What are the genes to consider?



Perez-Perez and  
Hanson. JCM 2002

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## Recent reports on plasmid AmpC



● Countries that have reported pAmpC in 2012 or 2013

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## CMY and linkage to plasmids

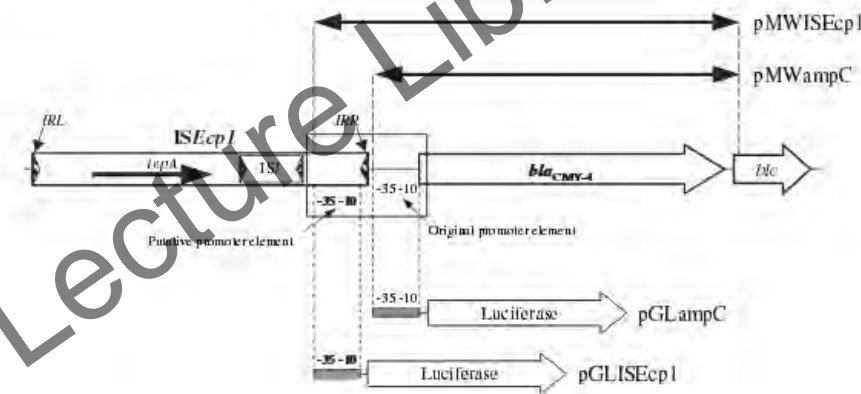
Enzyme	Replicon(s)	Species	No. of plasmids	Country(ies)	Source(s)
CMY-2	A/C	Escherichia coli, S. enterica	155	Canada, France, Honduras, Iraq, Ireland, UK, USA	Humans, cattle, pigs, poultry
CMY-2	I1	E. coli, S. enterica	30	Canada, France, Gambia, Italy, UK, USA	Humans, cattle, horses, dogs, pigs, poultry
CMY-2	FIA-FIB	E. coli	1	UK	Humans
CMY-2	NT	E. coli, S. enterica	6	Canada, UK	Cattle, poultry, pigs
CMY-4	A/C	S. enterica	1	UK	Humans
CMY-7	I1	E. coli	11	Pakistan, UK	Humans
CMY-8 (CTX-M-3)	HI2	Klebsiella pneumoniae	1	Taiwan	Humans
CMY-21	I1	E. coli	1	UK	Humans
CMY-31	ColE	S. enterica	1	USA	Humans
CMY-36	ColE	K. pneumoniae	1	Greece	Humans

Carattoli. AAC 2009

## Link of CMY-2 to ISEcp1

<i>bla</i> <sub>AmpC</sub>	ST	PG	Genetic support	Plasmid number	Size AmpC plasmid	Replicon typing
CMY-2	117 D		ISEcp1	2	80 kb	A/C
CMY-2	127 B2		ISEcp1	2	80 kb	I1
CMY-2	131 B2		ISEcp1	3	80 kb	-
CMY-2	131 B2		ISEcp1	3	80 kb	I1
CMY-2	131 B2		ISEcp1	4	140 kb	I1
CMY-2	131 B2		ISEcp1	2	80 kb	I1
CMY-2	359 A		ISEcp1	2	80 kb	A/C-FII
CMY-2	636 B2		ISEcp1	2	80 kb	I1
CMY-2	976 A		ISEcp1	1	80 kb	I1-A/C-FII
CMY-2	10 A		ISEcp1	2	80 kb	I1-FII
CMY-2	10 A		ISEcp1	2	80 kb	I1-FII
CMY-7	167 A		ISEcp1	2	80 kb	I1
CMY-2	155 A		ISEcp1	4	130 kb	-
CMY-2	155 B1		-	1	80 kb	I1-FII
CMY-2	93 A		ISEcp1	3	80 kb	-
DHA-1	23 A		<i>qacΔ1</i> <i>su1</i>	3	NT	-
CMY-2	88 A		ISEcp1	1	130 kb	A/C-FII
CMY-2	88 B1		ISEcp1	3	80 kb	-
CMY-2	393 D		ISEcp1	NT	NT	-
CMY-2	38 D		ISEcp1	1	NT	C <sup>β</sup>
CMY-2	778 D		ISEcp1	1	NT	C <sup>β</sup>
CMY-2	778 D		ISEcp1	1	NT	C <sup>β</sup>
CMY-2	778 D		ISEcp1	2	NT	C <sup>β</sup>
CMY-2	778 D		ISEcp1	3	NT	C <sup>β</sup>
CMY-2	963 D		-	1	80 kb	I1
CMY-2	981 D		-	1	80 kb	I1
CMY-2	405 D		ISEcp1	1	NT	C <sup>β</sup>
CMY-2	448 B1		ISEcp1	2	NT	C <sup>β</sup>
CMY-2	448 A		ISEcp1	1	NT	C <sup>β</sup>
CMY-2	448 A		ISEcp1	3	NT	C <sup>β</sup>
CMY-2	448 B1		ISEcp1	3	NT	C <sup>β</sup>
CMY-2	448 B1		ISEcp1	2	140 kb	FII
DHA-1	977 B1		-	-	NT	-
CMY-2	420 D		ISEcp1	3	80 kb	-
CMY-2	648 D		ISEcp1	3	80 kb	FII
CMY-2	69 D		ISEcp1	3	130 kb	A/C-FII
CMY-2	453 A		ISEcp1	2	140 kb	A/C-FII
CMY-2	979 B2		ISEcp1	3	250 kb	A/C

A



- *ISEcp1* provides the promoter regions that drive high-level expression of *bla*<sub>CMY-4</sub>
- Nakano et al. J Infect Chemother 2007

Naseer. CMI 2010



**Some recent data on international  
prevalence**

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## EARS-Net 2012: *E. coli* resistant to 3rd gen cephs

### Percentage resistance

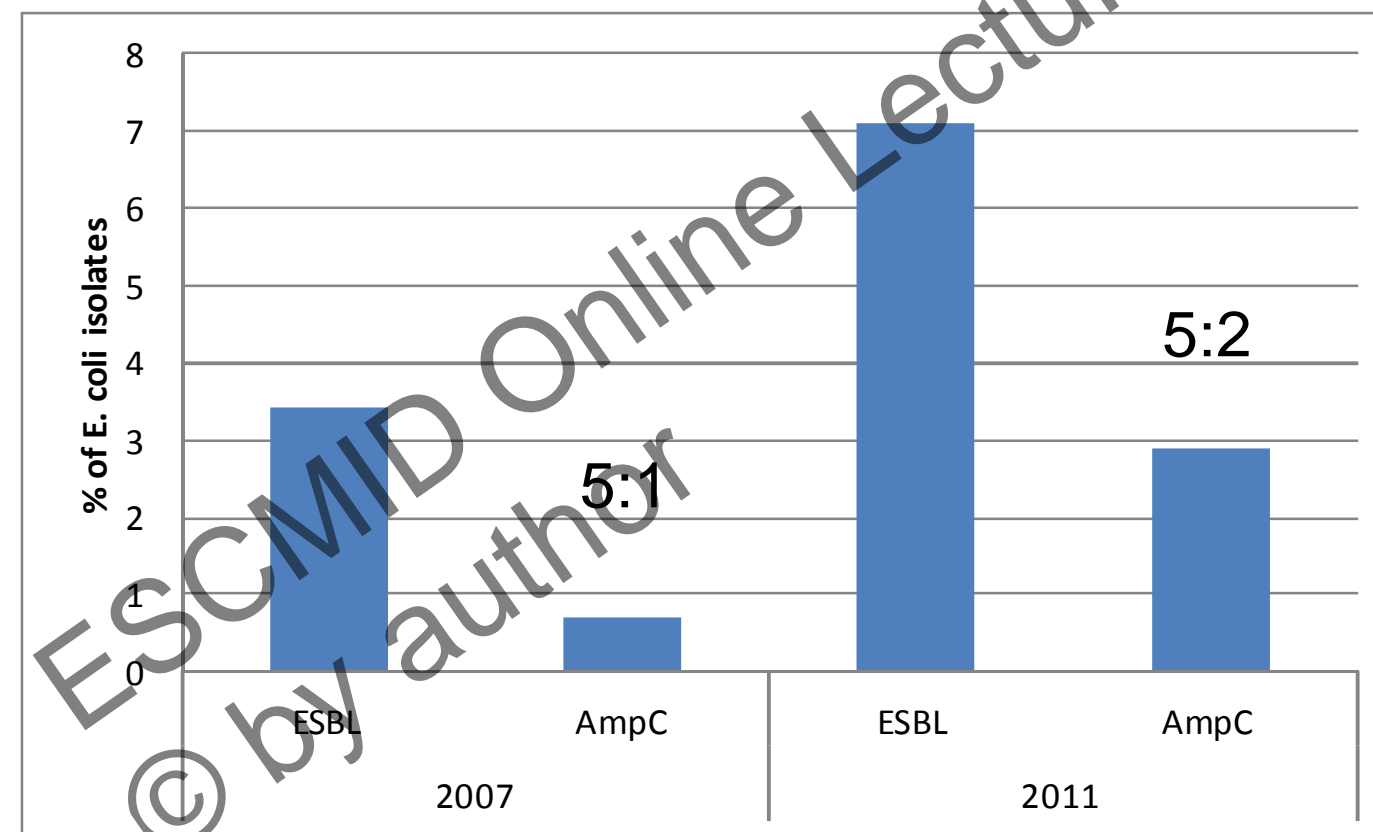
- < 1%
- 1 to < 5%
- 5 to < 10%
- 10 to < 25%
- 25 to < 50%
- ≥ 50%
- No data reported or less than 10 isolates
- Not included

- Liechtenstein
- Luxembourg
- Malta

(C) ECDC/Dundas/TESSy

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## CANWARD national surveillance study 2007-2011 (5,451 *E. coli*)

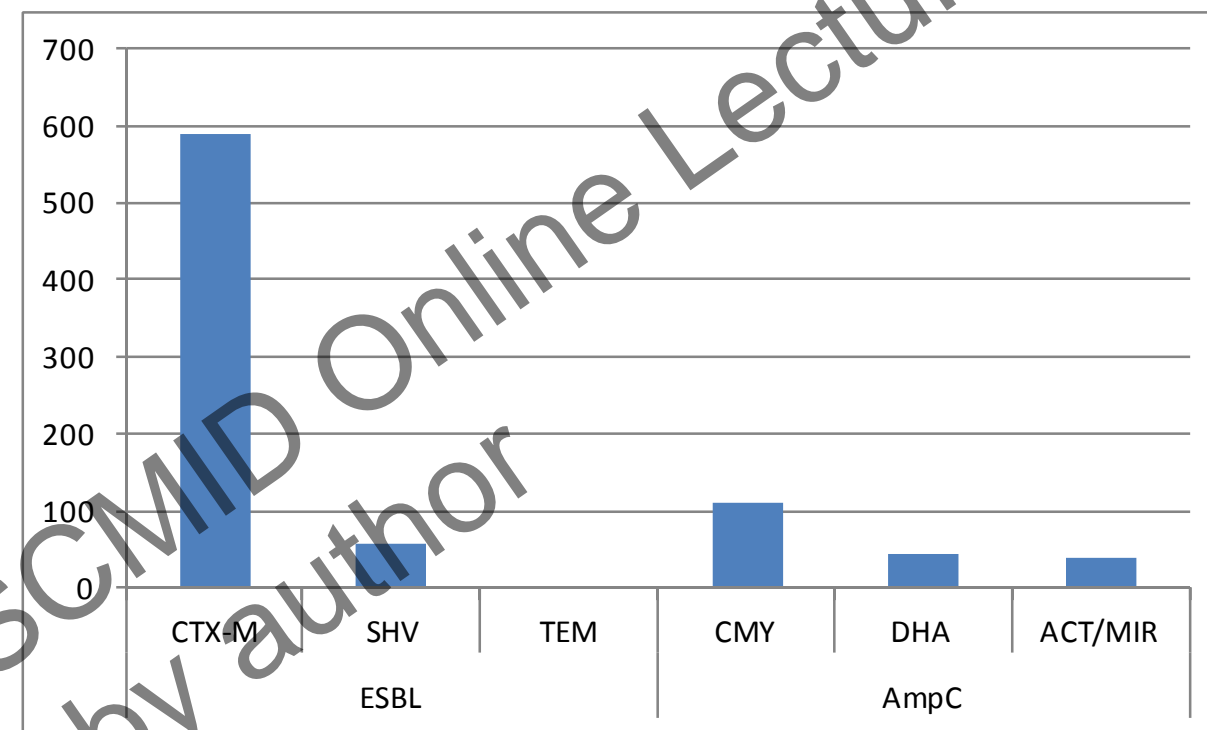


ST131:  
55.8% of  
ESBL vs  
28.7% of  
AmpC  
( $p=0.001$ )

55.8%  
CMY-2

Denisuik JAC 2013

## SMART study – intraabdominal infections Asia-Pacific 2008-2009



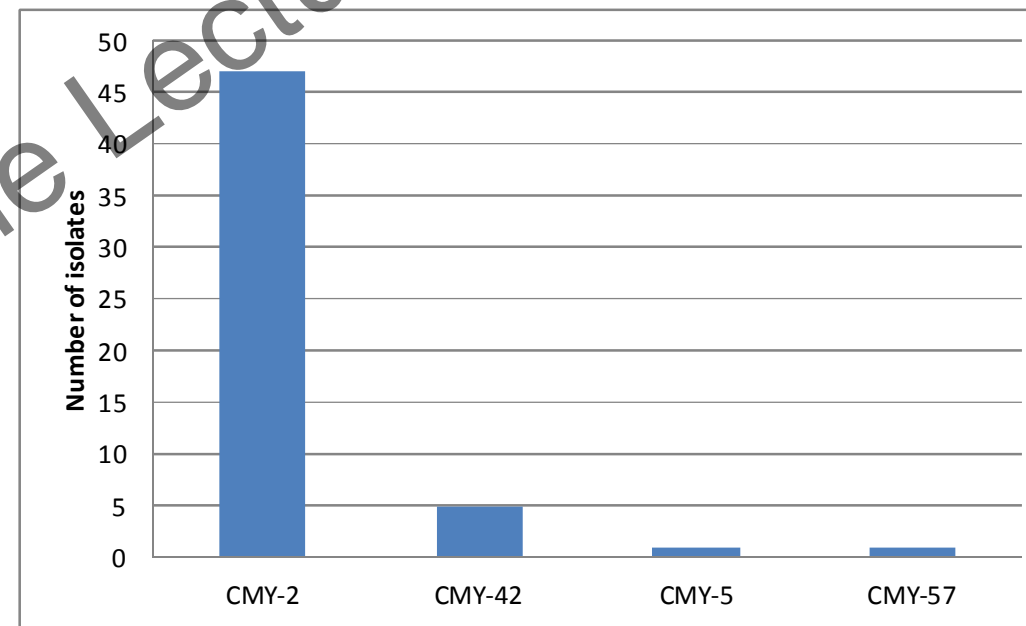
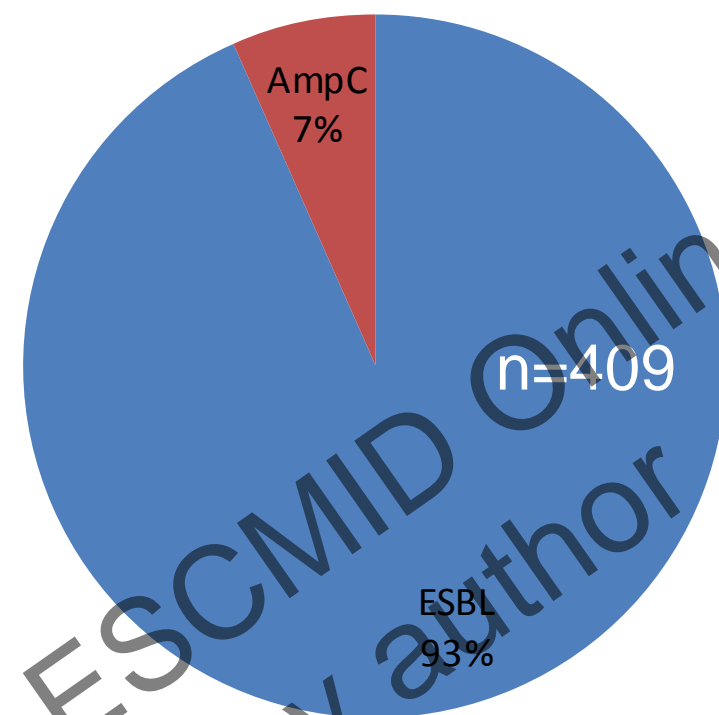
N=699 isolates with extended-spectrum cephalosporin resistance  
Sheng et al. AAC 2013

## SENTRY U.S. 2010

- 3,061 Enterobacteriaceae bloodstream isolates were consecutively collected from 26 hospitals located in 20 U.S. states during 2010 as part of the SENTRY Program
- 0.5% of the isolates had plasmid-mediated AmpC (Castanheira AAC 2013)

CMY-2, CTX-M-14	1	New York (1)	<i>E. coli</i> (1)
CMY-2, SHV-33	1	Vermont (1)	<i>K. pneumoniae</i> (1)
CMY-2, TEM-1	5	Colorado (1), Kentucky (1), Michigan (1), New Jersey (1), Texas (1)	<i>E. coli</i> (5)
CMY-like, TEM-1	1	Iowa (1)	<i>E. coli</i> (1)
DHA-1, TEM-1	1	Hawaii (1)	<i>P. mirabilis</i> (1)
FOX-5, PSE-like	3	Florida (1), Massachusetts (1), Virginia (1)	<i>E. cloacae</i> (1), <i>K. pneumoniae</i> (1), <i>P. mirabilis</i> (1)
FOX-5, SHV-11	1	Wisconsin (1)	<i>K. pneumoniae</i> (1)
FOX-5, PSE-like, SHV-11	1	Kentucky (1)	<i>K. pneumoniae</i> (1)

## National Swedish point prevalence study



Brolund A et al. CMI 2013 Epub



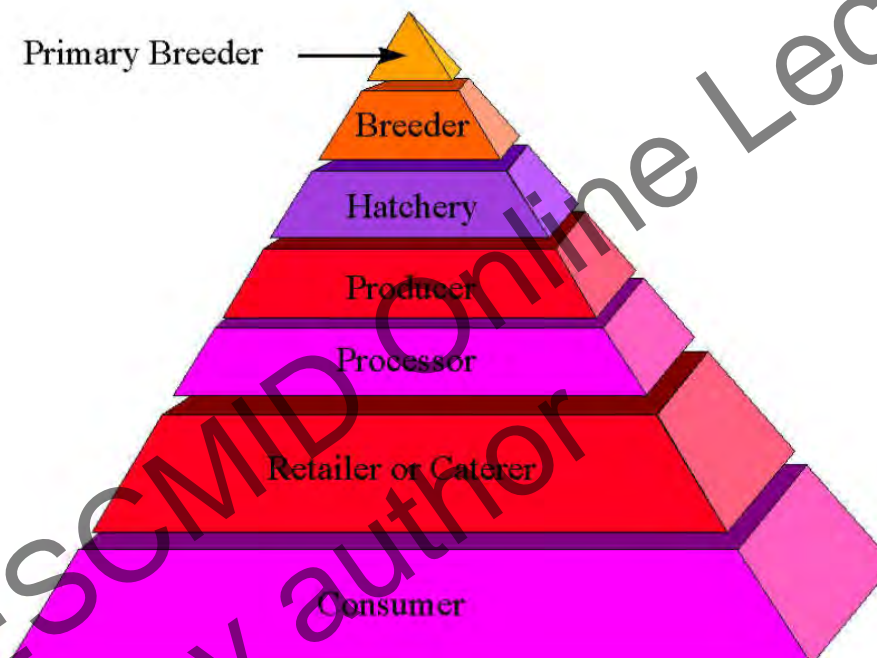
## Plasmid-mediated AmpC among animals

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## Poultry breeding pyramid



British Veterinary Poultry Association

- Apex: very small population of elite breeding birds
- 1 male selected by the primary breeder could give rise to 20 million offspring
- Purpose: rapid dissemination of genetic progress

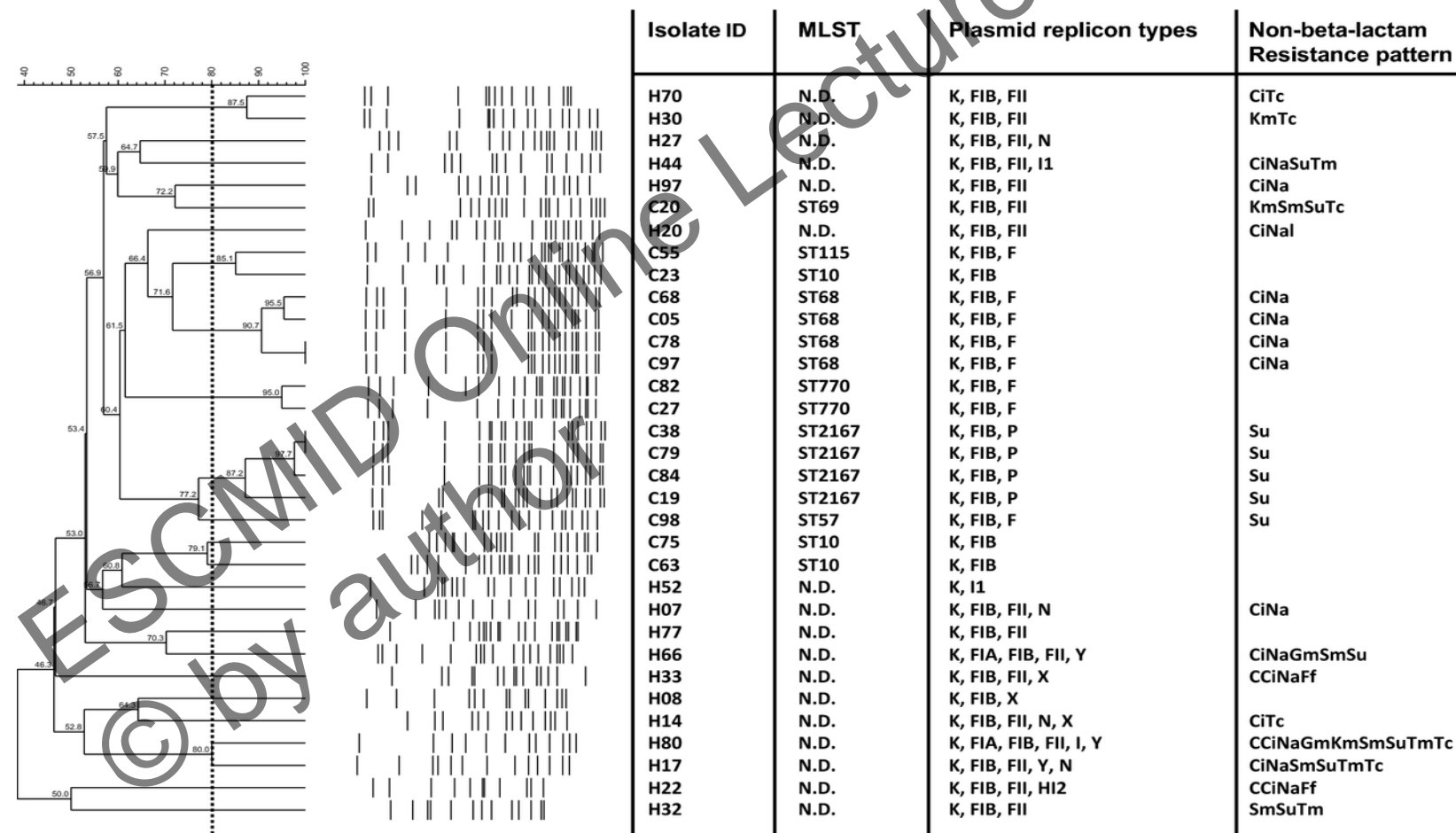
## AmpC and ESBLs in the broiler production, The Netherlands

1 Upon arrival in NL 2 days				2 Rearing farm 18 wks <sup>†</sup>				3 Production farm 31 wks			
Code	Sex	Caeca (n)	Prevalence ESBL/AmpC, % (95%CI)*	Poultry house	GPS chickens (n)	Cloaca swabs (n)	Prevalence ESBL/AmpC, % (95%CI) (phenotypes)	Poultry house	GPS chickens (n)	Cloaca swabs (n)	Prevalence ESBL/AmpC, % (95%CI) (phenotypes)
11-1	♂	10	20 (3-56)	1	2100 ♂	41	0 (0-9)	2	400 ♂ (PH 1) & 4000 ♀ (PH 4)	41	2 (0-13) (1 ESBL)
11-2	♂	10	0 (0-31)								
22-1	♀	10	10 (0-45)	4	4200 ♀	41	0 (0-9)	4	535 ♂ (PH 2b) & 5050 ♀ (PH 3)	41	24 (12-40) (10 AmpC)
22-2	♀	10	70 (35-93)								
44-1	♀	10	60 (26-88)	3 <sup>†</sup>	8250 ♀	25	100 (86-100) (24 AmpC, 1 ESBL)	3	380 ♂ (PH 2b) & 500 ♀ (PH 2a) & 3100 ♀ (PH 3)	41	27 (14-43) (8 AmpC, 3 ESBL)
33-1	♂	10	0 (0-31)	2b	3600 ♀	41	44 (28-60) (18 ESBL)	1	230 ♂ (PH 2b) & 2200 ♀ (PH 2a)	41	7 (2-20) (3 ESBL)
33-2	♂	10	10 (0-45)	2a	2730 ♀	41	2 (0-13) (1 ESBL)				
44-2	♀	10	10 (0-45)								

Arrows show how broilers were mixed between poultry houses with time  
 Only a few primary breeding companies at the top of the production pyramid,  
 and pAmpC is present already at this level (Dierikx et al. PloSOne 2013)

## AmpC among Swedish broilers

- 2010 Sweden: AmpC (CMY-2) and ESBL (CTX-M-1) among 34% of Swedish broilers, most likely related to imported breeding stocks
- No evidence of clonal spread between animals and humans



## Prevalence of AmpC and ESBL in healthy and diarrheic cats and dogs

ESBL/AmpC coding gene	Dog		Cat	
	Healthy	Diarrheic	Healthy	Diarrheic
No. of animals carrying				
<i>bla</i> <sub>CTX-M-1</sub>	4	2	–	–
<i>bla</i> <sub>CTX-M-15</sub>	–	–	–	1
<i>bla</i> <sub>CTX-M-14</sub>	–	1	–	–
<i>bla</i> <sub>CTX-M-1</sub> + <i>bla</i> <sub>CTX-M-14</sub>	–	1	–	–
<i>bla</i> <sub>CTX-M-1</sub> + <i>bla</i> <sub>TEM-52-StPaul</sub>	–	–	–	1
<i>bla</i> <sub>CTX-M-1</sub> + <i>bla</i> <sub>CMY-2</sub>	–	–	–	1
<i>bla</i> <sub>CTX-M-1</sub> + <i>bla</i> <sub>CMY-2</sub> + <i>ampC</i> type-3	–	–	–	1 <sup>f</sup>
<i>bla</i> <sub>CTX-M-1</sub> + <i>bla</i> <sub>CMY-2</sub> + <i>bla</i> <sub>TEM-35</sub>	–	1 <sup>a</sup>	–	–
<i>bla</i> <sub>CTX-M-1</sub> + <i>bla</i> <sub>CMY-2</sub> + unknown <sup>b</sup>	1	–	–	–
<i>bla</i> <sub>CTX-M-1</sub> + <i>ampC</i> type-3	–	1	–	–
<i>bla</i> <sub>SHV-12</sub> + <i>bla</i> <sub>CMY-2</sub>	–	–	–	1
<i>bla</i> <sub>CMY-2</sub>	2	2	–	–
<i>ampC</i> type-3 <sup>c</sup>	1	1	–	–
<i>ampC</i> type-new <sup>d</sup>	0	1	–	–
Unknown <sup>b</sup>	1	1	–	–
>1 R gene <sup>e</sup>	1	3	0	4
<i>E. coli</i>	9	11	0	4
<i>E. coli</i> + <i>P. mirabilis</i>	0	0	0	1 <sup>f</sup>

<sup>a</sup>This animal carried one *E. coli* harboring both a *bla*<sub>TEM-35</sub> gene and a *bla*<sub>CMY-2</sub> gene. The *bla*<sub>CTX-M-1</sub> gene and another *bla*<sub>CMY-2</sub> gene were found in separate *E. coli*.

<sup>b</sup>Isolates negative on Check-MDR CT-103 array, and no *ampC* type-3.

<sup>c</sup>Classification as previously described (Mulvey et al., 2005).

<sup>d</sup>New *ampC* variant compared to previously described variants (Mulvey et al., 2005).

<sup>e</sup>Number of animals with more than one type of ESBL/pAmpC gene.

<sup>f</sup>This animal carried a *P. mirabilis* harboring a *bla*<sub>CMY-2</sub> gene. All other genes reported in this table were found in *E. coli*.

Hordijk. Front Microbiol 2013

**How do you detect plasmid-mediated  
AmpC?**

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# EUCAST guidelines



EUCAST guidelines for detection of resistance mechanisms and specific resistances of clinical and/or epidemiological importance

Version 1.0  
November 2013

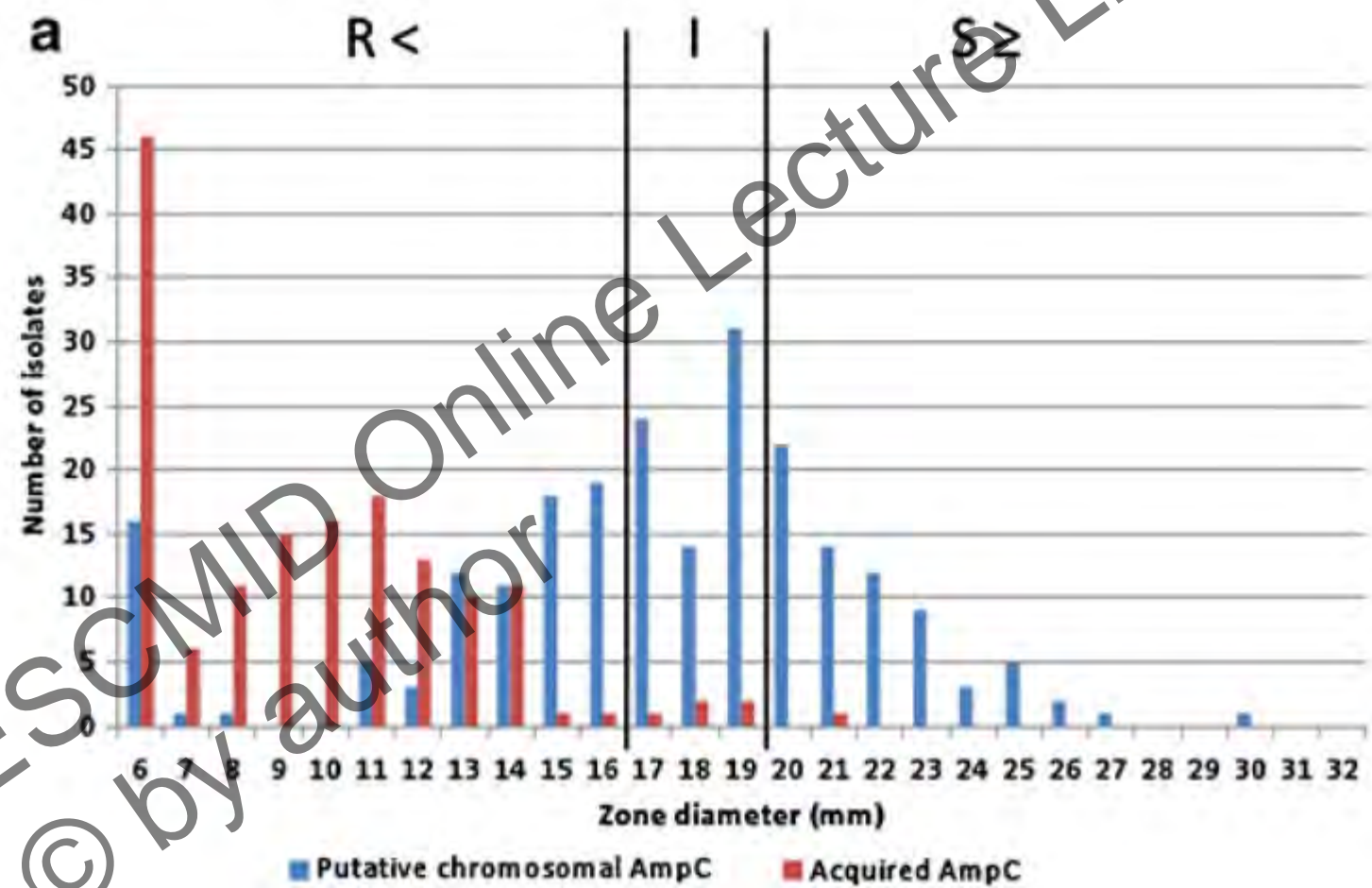


## Contents

Section	Page
1. Introduction	3
2. Carbapenemase-producing Enterobacteriaceae	4
3. Extended-spectrum $\beta$ -lactamase-producing Enterobacteriaceae	11
<u>4. Acquired AmpC <math>\beta</math>-lactamase-producing Enterobacteriaceae</u>	<u>20</u>
5. Methicillin resistant <i>Staphylococcus aureus</i>	24
6. Glycopeptide non-susceptible <i>Staphylococcus aureus</i>	27
7. Vancomycin resistant <i>Enterococcus faecium</i> and <i>Enterococcus faecalis</i>	31
8. Penicillin non-susceptible <i>Streptococcus pneumoniae</i>	36
9. Transparency declaration	39

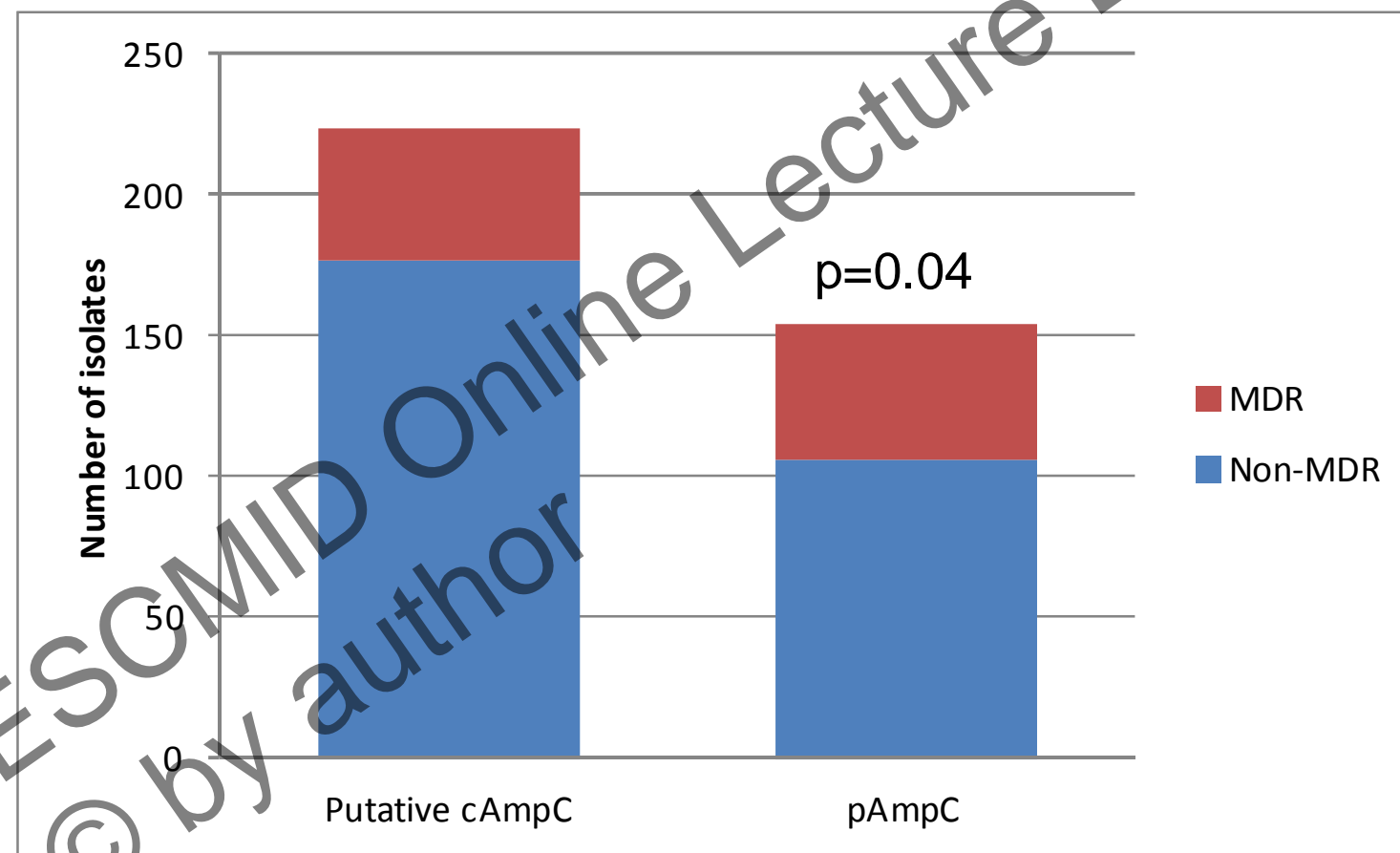
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## EUCAST disk diffusion – CTX 5 $\mu\text{g}$



Edquist P et al. Eur J Clin Microbiol Infect Dis 2013

## Prevalence of MDR (2/3 of GEN, CIP, TSU)



Edquist P et al. Eur J Clin Microbiol Infect Dis 2013

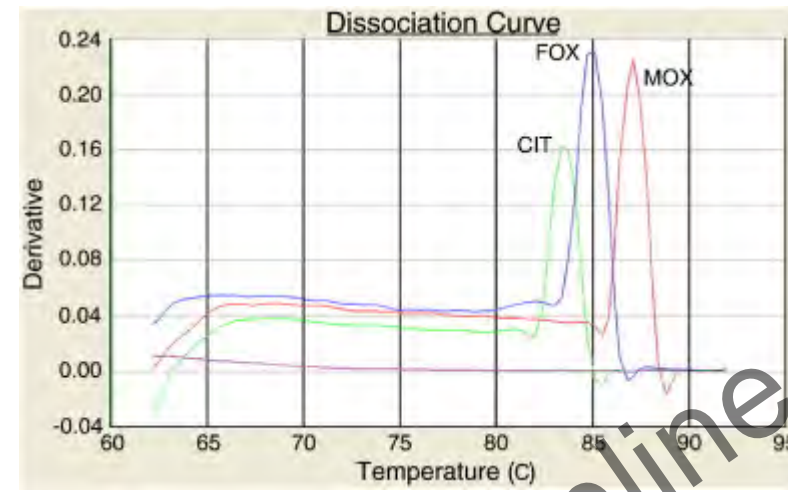


## Comparison of methods

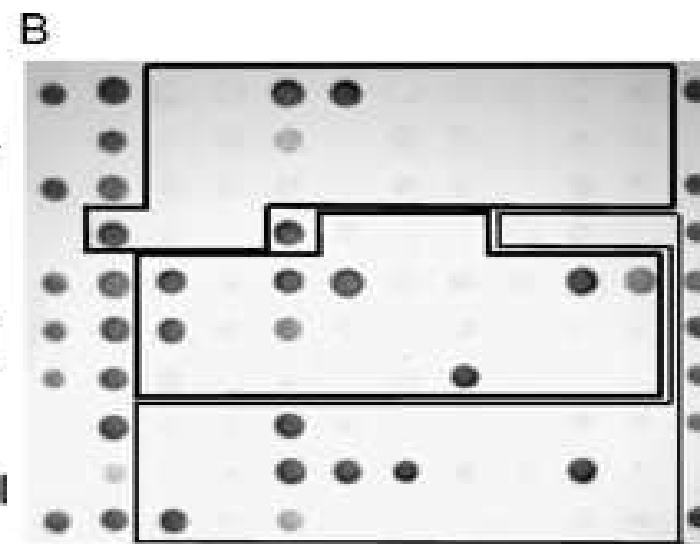
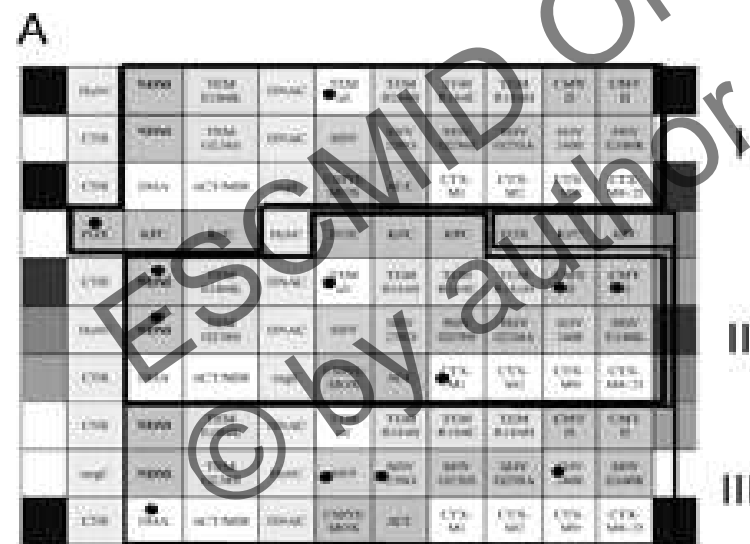
Method	Sensitivity	Specificity*
In-house FOX+cloxacillin	98%	52%
Rosco CTX and CAZ+cloxacillin	97%	54%
Etest cefotetan + cloxacillin	93%	69%

\* Specificity calculated as specificity for pAmpC-detection

# Genotypic detection



Brolund A.  
J Microbiol  
Methods 2012



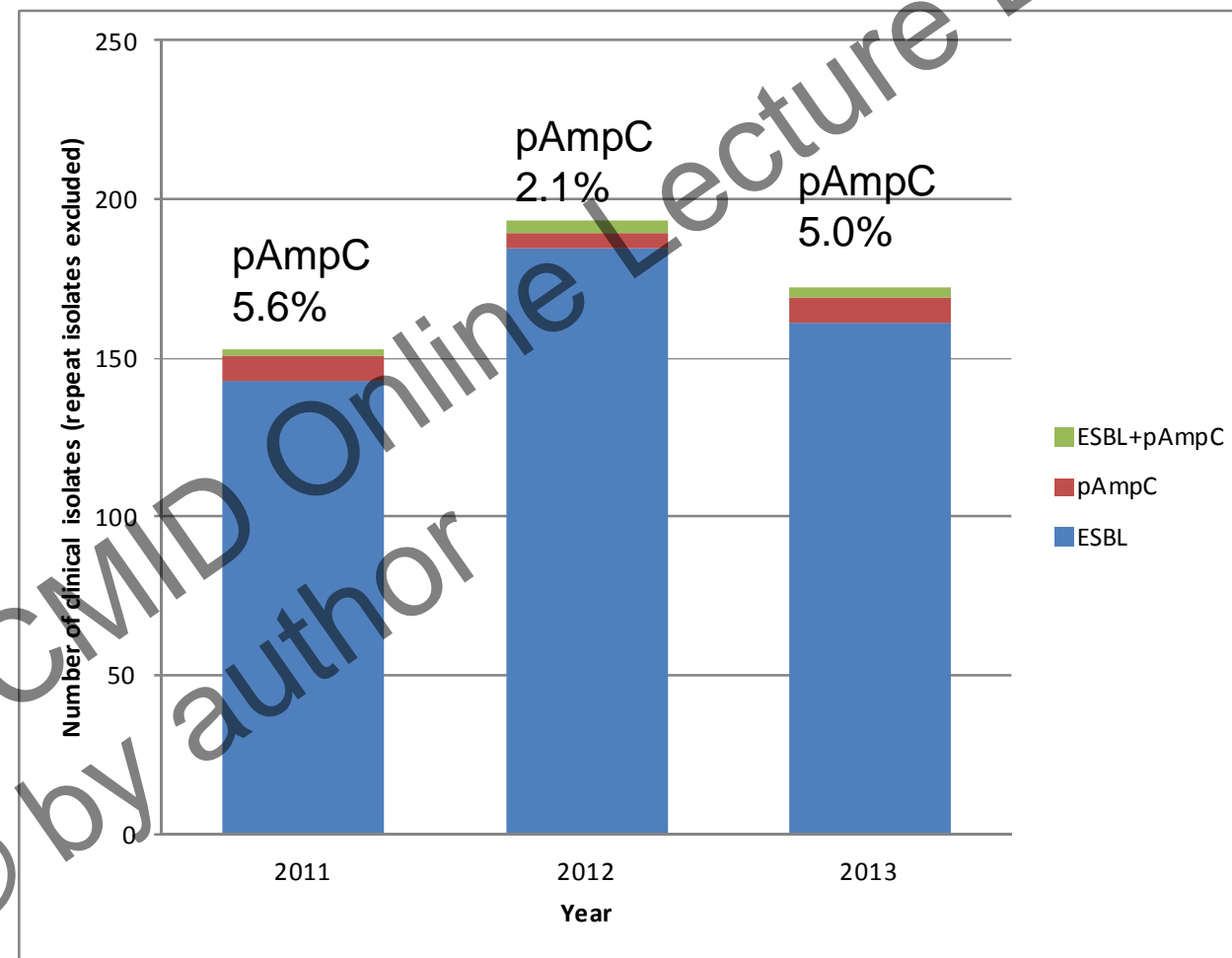
Bogaerts P.  
AAC 2011

**What do you find when you search beyond  
point prevalence studies?**

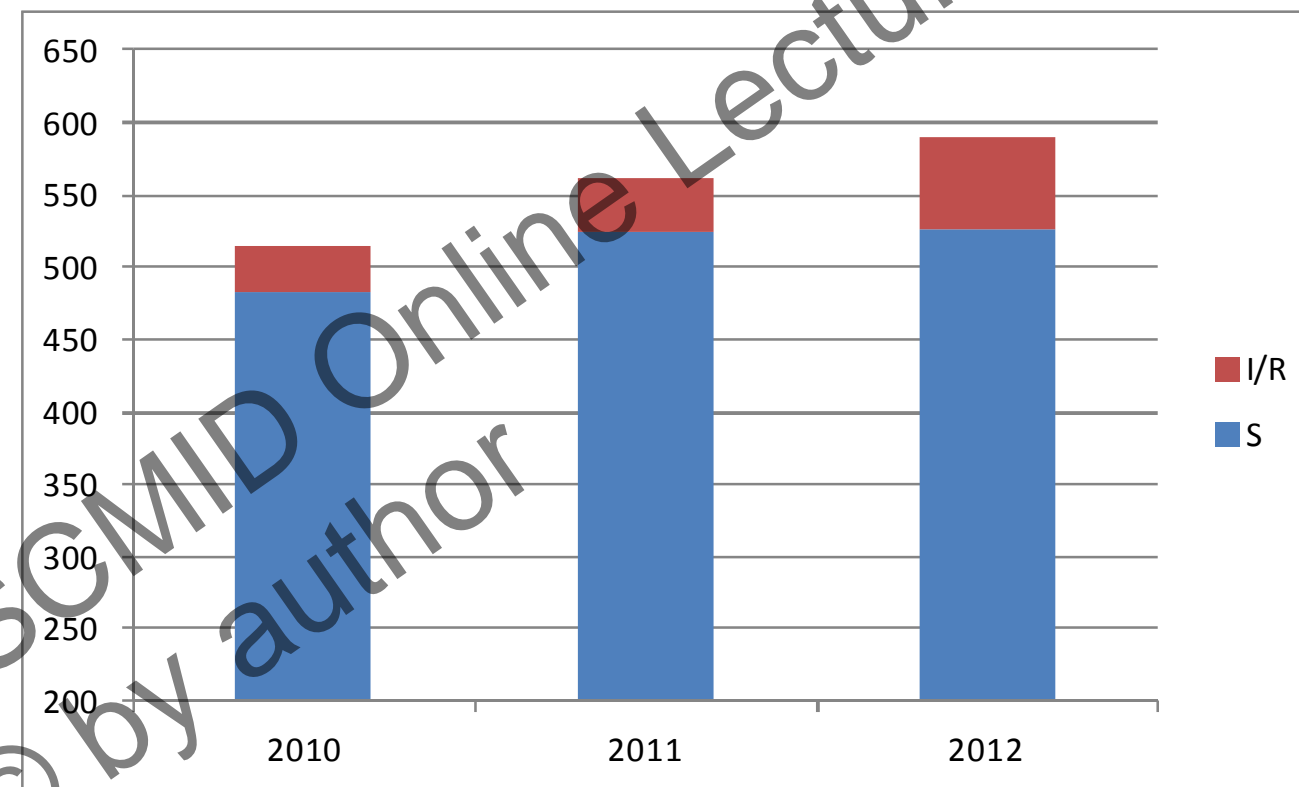
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## Transferable cephalosporin resistance in *E. coli* at Karolinska 2011-2013



## 3rd gen ceph R *E. coli* are increasing the burden of disease; blood cultures at Karolinska



## Conclusions

- No evidence supporting a rapid increase of pAmpC in *E. coli*
  - Slow increase observed in most areas where these enzymes are actively searched for
  - In many areas pAmpC are far more common than the minor ESBLs (SHV- and TEM-ESBLs)
  - CMY-2 common among food-producing and companion animals
  - CMY-2 has all the hallmarks of a successful enzyme, including the link to ISEcp1, broad-range plasmids, and to ST131
  - pAmpC-testing in blood culture isolates of *E. coli* with extended-spectrum cephalosporin resistance/negative ESBL-test may be warranted
-

Thank you for your attention!



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