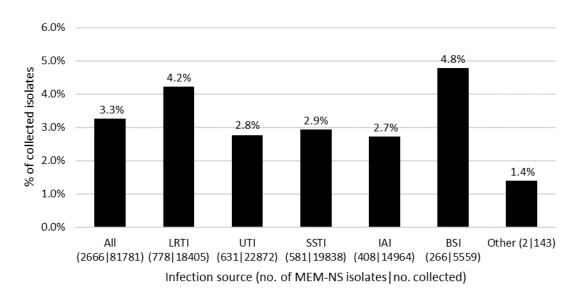
AAC02000-20R1

2 Supplementary material

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- 4 Figure S1. Distribution of meropenem-nonsusceptible Enterobacterales isolates collected
- 5 globally from 2012 to 2017, by infection source



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- 7 LRTI, lower respiratory tract infection; UTI, urinary tract infection; SSTI, skin and soft tissue
- 8 infection; IAI, intra-abdominal infection; BSI, bloodstream infection; MEM-NS, meropenem-
- 9 nonsusceptible.

Figure S2. Species composition of meropenem-nonsusceptible Enterobacterales isolates

collected globally from 2012 to 2017

1.2% 0.7% 0.3% 3.0% _ 2.6% 3.8% ■ K. pneumoniae (n=2046) ■ E. cloacae (n=177) 5.1% ■ E. coli (n=136) ■ Klebsiella spp. (n=101) 6.6% ■ Citrobacter spp. (n=79) ■ Proteeae (n=70) ■ S. marcescens (n=31) ■ Enterobacter spp. (n=18) ■ Raoultella spp. (n=8) 76.7%

15 Klebsiella spp. were composed of Klebsiella aerogenes (n=51), Klebsiella oxytoca (n=47), and

Klebsiella variicola (n=3). Citrobacter spp. were composed of Citrobacter freundii (n=64),

17 Citrobacter koseri (n=7), Citrobacter farmeri (n=4), Citrobacter amalonaticus (n=2),

Citrobacter braakii (n=1), and Citrobacter, species not determined (n=1). The Proteeae were

composed of Providencia rettgeri (n=27), Proteus mirabilis (n=21), Providencia stuartii (n=19),

Proteus hauseri (n=2), and Proteus vulgaris (n=1). Enterobacter spp. were composed of

Enterobacter asburiae (n=15) and Enterobacter kobei (n=3). Raoultella spp. were composed of

Raoultella ornithinolytica (n=5) and Raoultella planticola (n=3).

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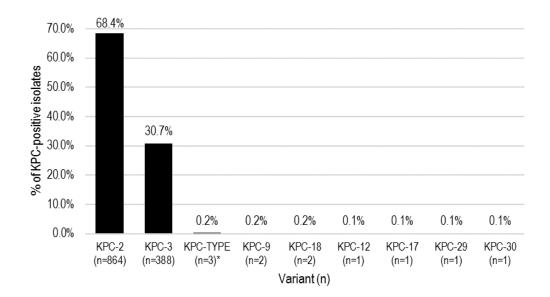
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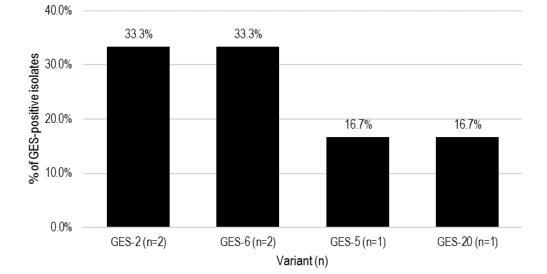
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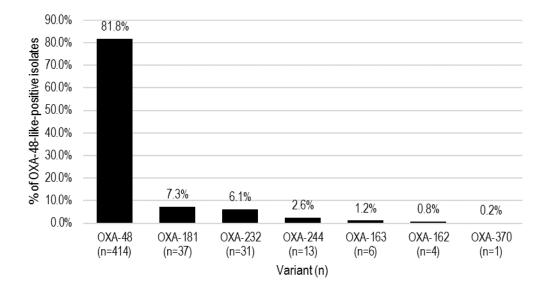
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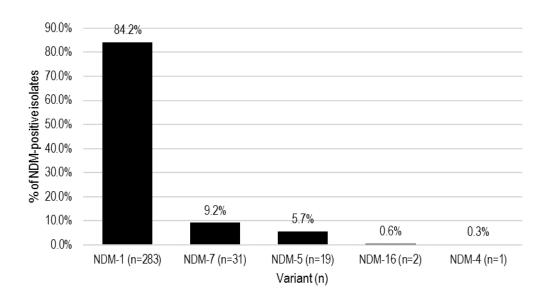
Figure S3. Diversity and relative prevalence of sequence variants carried by meropenem-

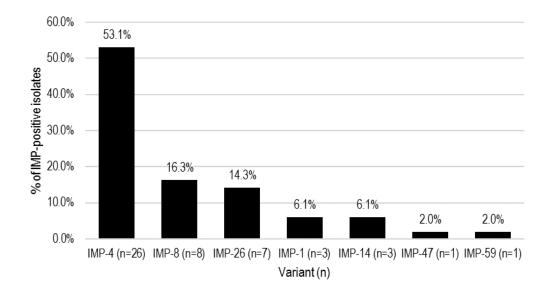
nonsusceptible Enterobacterales isolates collected globally from 2012 to 2017

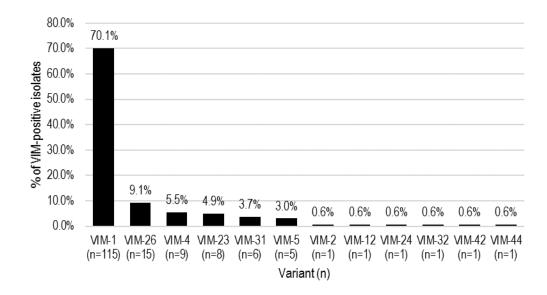






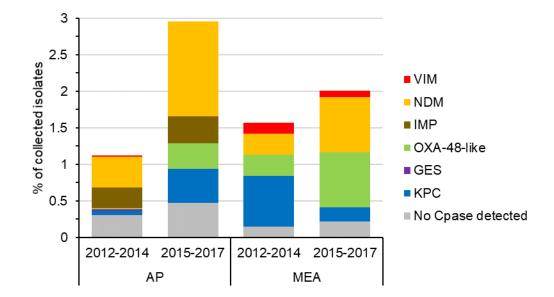






*KPC-type; the bla_{KPC} could not be amplified for sequencing using flanking primers that hybridized entirely outside of the coding sequence and instead was amplified using primers that overlap the ends of the gene. For this reason, a sequence variant was not assigned.

Figure S4. Comparison of the distribution of carbapenem-resistance mechanisms identified in meropenem-nonsusceptible Enterobacterales isolates collected in countries in Asia/South Pacific and Middle East/Africa that participated in all 6 years surveyed



Region (countries; no. of isolates collected in 2012-2014|no. of isolates collected in 2015-2017):

AP, Asia/South Pacific (Australia, Japan, the Philippines, South Korea, Taiwan and Thailand;

5,268|6,831); MEA, Middle East/Africa (Israel, Kuwait, and South Africa; 2,739|3,690).

Countries that were removed from analysis (years of participation): China (2012-2013), Hong

Kong (2012-2014), and Malaysia (2013-2015) (AP); Kenya (2013-2014) and Nigeria (2013-2014) (MEA). No Cpase detected, no gene encoding a carbapenemase was detected by PCR.

Isolates carrying multiple carbapenemases were counted for each individual carbapenemase type.

		Years of participation ^a					
Region	Country	2012	2013	2014	2015	2016	2017
Europe	Austria	X	X	X	X	X	_
	Belgium	X	X	X	X	X	X
	Czech Republic	X	X	X	X	X	X
	Denmark	X	X	X	X	X	X
	France	X	X	X	X	X	X
	Germany	X	X	X	X	X	X
	Greece	X	X	X	X	X	X
	Hungary	X	X	X	X	X	X
	Italy	X	X	X	X	X	X
	Netherlands	X	X	X	X	X	X
	Poland		X	X	X	X	X
	Portugal	X	X	X	X	X	X
	Romania	X	X	X	X	X	X
	Russia	X	X	X	X	X	X
	Spain	X	X	X	X	X	X
	Sweden	X	X	X	_	_	X
	Turkey	X	X	X	X	X	X
	United Kingdom	X	X	X	X	X	X
Latin America	Argentina	X	X	X	X	X	X
-	Brazil	X	X	X	X	X	X
	Chile	X	X	X	X	X	X
	Colombia	X	X	X	X	X	X
	Mexico	X	X	X	X	X	X
	Venezuela	X	X	X	X	X	X
Asia/ South	Australia	X	X	X	X	X	X
Pacific	Mainland China	X	X	NA^b	NA^b	NA^b	NAb
	Hong Kong ^c	X	X	X	_	_	_
	Japan	X	X	X	X	X	X
	Malaysia		X	X	X	_	_
	Philippines	X	X	X	X	X	X
	South Korea	X	X	X	X	X	X
	Taiwan	X	X	X	X	X	X
	Thailand	X	X	X	X	X	X
Middle East/	Israel	X	X	X	X	X	X
Africa	Kenya	_	X	X			_
	Kuwait	X	X	X	X	X	X
	Nigeria	_	X	X		_	_
	South Africa	X	X	X	X	X	X
North America	United States	X	X	X	X	X	

a "X" denotes participation during the indicated year. "—" denotes no participation during the indicated year.

^b "NA", isolates were collected from patients in mainland China but were not available due to export restrictions.

- 60 ^c Isolates were obtained from patients in Hong Kong during the first three years of the study (2012-2014). Hong
- Kong is listed separately from mainland China for this reason.