



Figure S1: Distribution of meropenem minimum inhibitory concentrations (MIC mg/L) by Flank Pattern (FP). Circle size indicates the number of observations per category and the red hashed line denotes the EUCAST breakpoint (>8mg/L). Isolates from the EuSCAPE dataset (short read only sequencing, n=226 isolates with linkable phenotype data) were assigned to a Flank Pattern based on the top mash containment hit). No isolates were assigned to FPs 4,8 or 17 and the one isolate assigned to FP10 had no linkable phenotypic data.

Table S1 – NCBI project accession numbers for sequencing data used

Description	Project Accession	Number of isolates
Complete plasmid assemblies containing <i>bla_{KPC2/3}</i>	PRJEB33308 (EuSCAPE)	42
	ERP118777 (Dutch CPE)	8
Complete plasmid assemblies containing <i>bla_{OXA-48}</i>	PRJEB33308 (Dutch CPE)	9
	PRJNA591727 (EuSCAPE)	42
Short read Illumina fastq files for isolates containing <i>bla_{OXA-48}</i> *	PRJEB10018 (EuSCAPE)	425
Short read Illumina fastq files for isolates containing <i>bla_{KPC-2/3}</i> *	PRJEB10018 (EuSCAPE)	442

*Full list of accession numbers used and associated metadata is available at <https://doi.org/10.6084/m9.figshare.14074250.v1>

Table S2 – Flank Pattern (FP), MEFinder and TETyper calls for blaKPC-2/3 flanking regions. The isolate column gives the contig name within the assembly, the next three columns give FP assignments at 500/5000/7200bp upstream of the gene. Columns 5 to 8 give the MEFinder/TETyper calls for the 500bp/5000bp upstream flanking regions whereas the final column gives TETyper calls when TETyper was given the whole plasmid assembly as input.

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFYD010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
MT560073.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYQ010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFZV010000 006.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGAG01000 0002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYI0100000 02.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGAO01000 0002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFYF010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYH010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
MT560060.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGAI010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABGAZ010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYZ010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGAQ010000 0009.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGBS010000 004.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFYU010000 005.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGAR010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
MT560075.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFZA010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFX010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFZG010000 005.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYK010000 005.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFXS010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFY010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGAY01000 0002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGBH010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYC010000 004.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFYE010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGAK01000 0004.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYG010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGAX01000 0009.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2203 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFXR010000 002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYB010000 005.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYS010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYL010000 004.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFYT010000008.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
MT560078.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABGBM010000006.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYN010000003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
CABFZZ010000 003.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFZH010000 004.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
CABFYM01000 0002.1	1	1	1	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6602 70 20- 7118 72 02- 10006	1- 2202 70 20- 7118 72 02- 10006	Tn4401a
MT560080.1	2	4	5	No hits	Tn1000 1 X6020 0.1	1- 7127 72 02- 10006	1- 7127 72 02- 10006	unknown

Contig Name	Flanker 500bp	Flanker 5000 bp	Flanker 7200 bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
MT560061.1	2	4	4	No hits	Tn1000 1 X6020 0.1	1- 7127 72 02- 10006	1- 7127 72 02- 10006	unknown
MT560063.1	2	3	6	No hits	Tn1000 1 X6020 0.1	1- 7127 72 02- 10006	1- 7127 72 02- 10006	unknown
CABFYR010000 005.1	3	2	2	No hits	Tn6296 1 FJ628 167	1- 7127 72 02- 10006	1- 7127 72 02- 10006	Tn4401_tru ncC
CABGBR010000 009.1	3	2	2	No hits	Tn6296 1 FJ628 167	1- 7127 72 02- 10006	1- 7127 72 02- 10006	Tn4401_tru ncC
MT560066.1	3	5	3	No hits	No hits	1- 7127 72 02- 10006	1- 7127 72 02- 10006	Tn4401_tru ncC
CABFYJ010000 003.1	4	1	8	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6633 70	1- 2233 70	unknown

Contig Name	Flanker 500bp	Flanker 5000bp	Flanker 7200bp	MEFinder 500bp	MEFinder 5000bp	TETyper 500bp	TETyper 5000bp	TETyper Tn4401 whole
						08- 7075 72 02- 10006	08- 7075 72 02- 10006	
CABFZC010000 002.1	4	1	7	Tn4401 1 KT37 8596.1	Tn4401 1 KT37 8596.1	1- 6633 70 08- 7075 72 02- 10006	1- 2233 70 08- 7075 72 02- 10006	Tn4401d

Flank Pattern	N
FP1	1
FP2	37
FP3	8
FP4	0
FP5	4
FP6	230

Flank Pattern	N
FP7	2
FP8	0
FP9	5
FP10	1
FP11	4
FP12	2
FP13	9
FP14	1
FP15	13
FP16	108
FP17	0

Table S3 – Number of bla_{OXA-48} Flank Patterns (FPs) observed in the EuSCAPE short read assembly dataset. For each FP identified by Flanker in the hybrid assembly dataset (2200bp upstream of the gene), one random representative was chosen (see main methods). EuSCAPE short read assemblies (n=425 total) were screened for containment of these using mash and the FP was taken to be the match with greatest containment.

	Group 1	Group 2	Group 7	Group 8
Austria	0	0	1	0
Belgium	0	0	0	0

	Group 1	Group 2	Group 7	Group 8
Croatia	0	0	0	0
France	0	0	0	0
Germany	0	0	2	1
Greece	0	0	0	3
Ireland	0	0	0	0
Israel	0	0	19	0
Italy	0	0	1	0
Luxembourg	0	0	0	0
Macedonia	0	0	0	0
Poland	0	0	2	0
Portugal	0	0	34	0
Romania	0	0	0	0
Slovakia	0	0	0	0
Spain	0	5	0	0
United Kingdom (England, Wales & N. Ireland)	0	0	0	2

Table S4 – Geographical distribution of bla_{KPC-2/3} flanking patterns observed in the EuSCAPE short read assembly dataset (n=313 with linkable geographic data). For each FP identified by Flanker in the hybrid assembly dataset (7200bp upstream of the gene), one random representative was chosen (see main methods). EuSCAPE short read assemblies were screened for containment of these using mash and the FP was taken to be the match with greatest containment. There were no isolates assigned to FPs3/4/5/6 that had linkable geographic data.

Meropenem MIC (mg/L)	FP 1	FP 2	FP 7	FP 8
≤0.06	0	0	0	0
0.12	0	1	0	0
0.25	0	0	0	0
0.5	0	0	0	0
1	0	0	0	0
2	0	1	1	0
4	0	1	1	0
8	0	1	0	1
16	0	0	3	3
32	0	0	6	1
>32	0	1	14	1

Table S5 – Distribution of meropenem minimum inhibitory concentrations (MIC) by blaKPC-2/3 7200bp upstream flanking region in the EuSCAPE dataset (n= 274 with linkable phenotypic data). For each FP identified by Flanker in the hybrid assembly dataset (7200bp upstream of the gene), one random representative was chosen (see main methods). EuSCAPE short read assemblies were screened for containment of these using mash and the FP was taken to be the match with greatest containment. There were no isolates assigned to FPs3/4/5/6 that had linkable phenotypic data.