

NDM-1 Introduction in Portugal through a ST11 KL105 *Klebsiella pneumoniae* Widespread in Europe

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Table S1. List of *Klebsiella pneumoniae* ST11-KL105 genomes deposited on NCBI public databases used for phylogenetic analysis.

Strain Name	Accession number	Country	Isolation Date
L176	GCF_002262735.1	China	2016
L37	GCF_002262415.1	China	2016
L36	GCF_002262385.1	China	2016
L50	GCF_002262115.1	China	2016
L26	GCF_002262455.1	China	2016
L519	GCF_002261945.1	China	2016
L447	GCF_002262145.1	China	2016
L23	GCF_002262565.1	China	2016
L196	GCF_002262635.1	China	2016
L84	GCF_002261575.1	China	2016
L526	GCF_002261935.1	China	2016
L8	GCF_002261685.1	China	2016
L442	GCF_002262175.1	China	2016
L70	GCF_002261655.1	China	2016
L81	GCF_002261615.1	China	2016
L63	GCF_002261785.1	China	2016
XL-1	GCF_001939845.1	China	2015
KPC40	GCF_002209465.1	Brazil	2015
K606	SAMN16522901	Portugal	2018
K624	SAMN16522902	Portugal	2019
KP38731	GCF_002009075.1	USA	2015
EuSCAPE_RO073	GCF_900504255.1	Romania	2014
EuSCAPE_RO046	GCF_900504005.1	Romania	2013
EuSCAPE_RO054	GCF_900504085.1	Romania	2014
GEN000186	GCF_004146185.1	France	2017
EuSCAPE_PT025	GCF_900502825.1	Portugal	2014
H1523	GCA_009746735.1	Portugal	2011
H642	GCA_009746705.1	Portugal	2006
EuSCAPE_SK022	GCF_900510755.1	Slovakia	2014
EuSCAPE_HU027	GCF_900516865.1	Hungary	2014
EuSCAPE_PL054	GCF_900510385.1	Poland	2014
EuSCAPE_PL012	GCF_900511375.1	Poland	2013

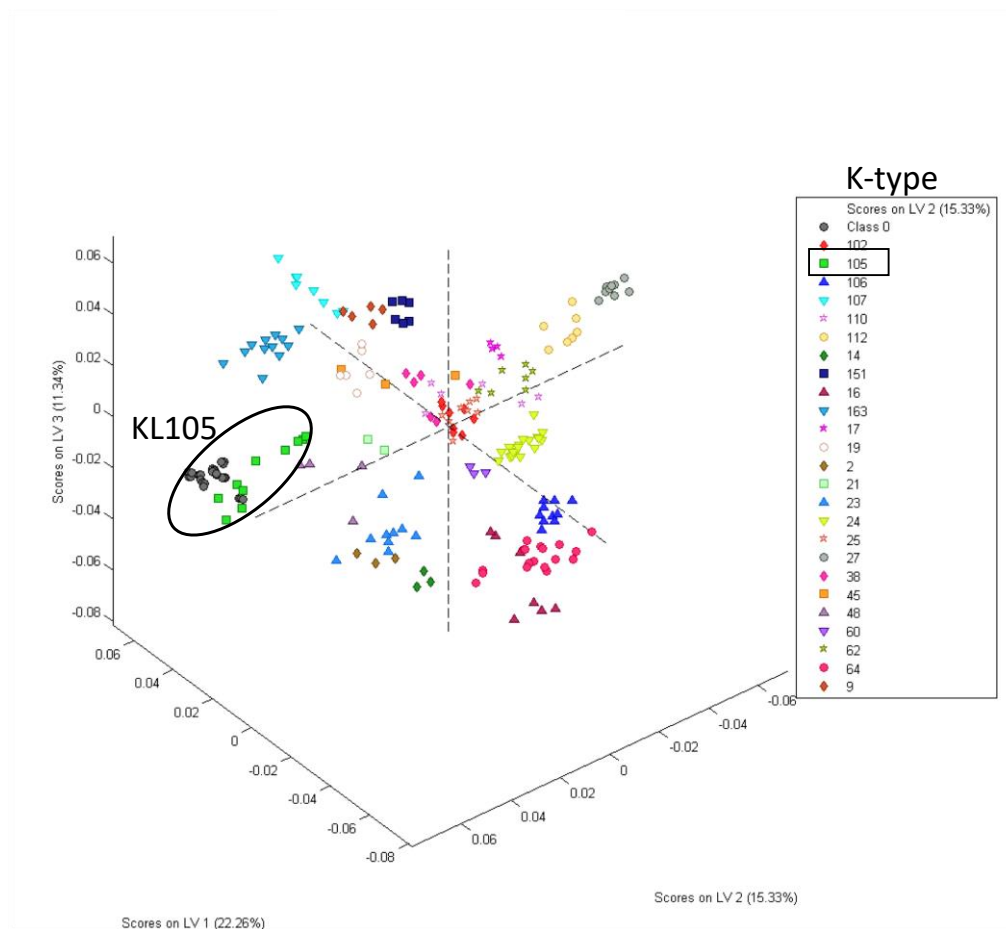


Figure S1. Projection of FT-IR spectra from outbreak *K. pneumoniae* isolates (grey dots) in the in-house partial-least squares discriminant model (PLSDA) discriminating 25 K-types.

0

Not yet implemented

0

100

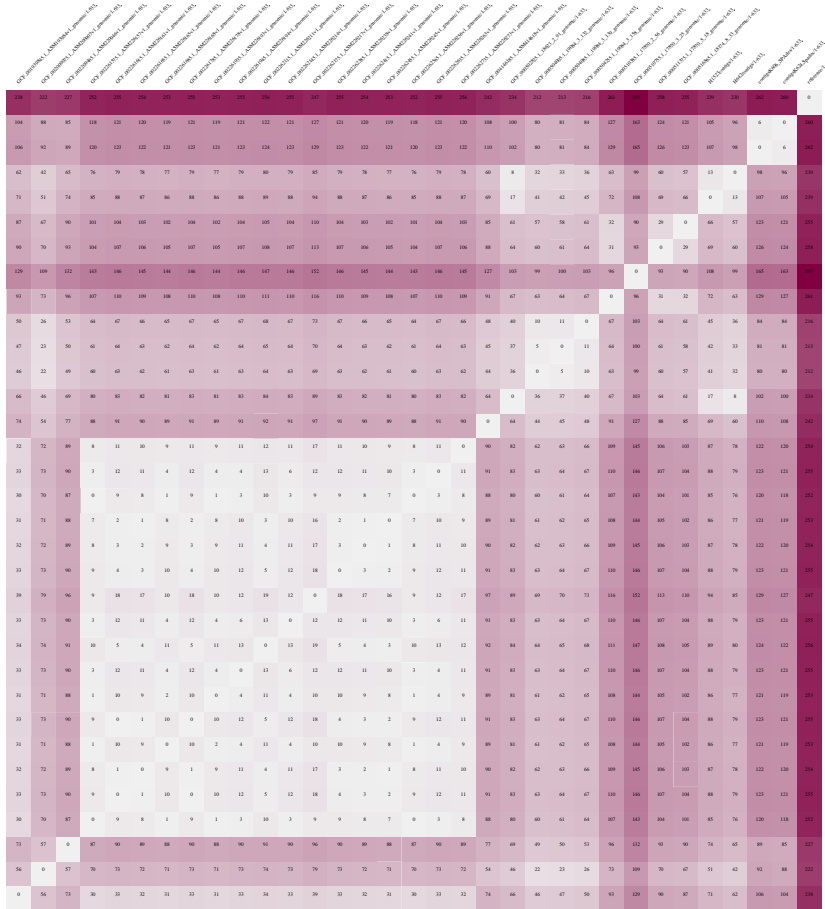


Figure S2. SNP matrix obtained from core-genome DNA comparison between *K. pneumoniae* ST11 KL105 genomes supporting phylogenetic tree.