

Supplementary Figure S2: Matrix of 1133 out of 4891 alleles; not taking into account missing values of *K. pneumoniae* ST15 producing *bla*_{VIM-1} and *bla*_{CTX-M-15}, the 13 contemporary genomes of ST15 isolates from other centers in the region and the 2 genomes of VIM-1-producing ST15 from Spanish regions other than Andalusia.

