Emergence of *Escherichia coli* carrying carbapenemase genes, European Union/European Economic Area, August 2012 to May 2024

## **Supplementary material**

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## **SUPPLEMENTARY FIGURE.** Time distribution of isolates of *Escherichia coli* ST131, including its single locus variants (SLVs) within clusters by reporting country and cluster, $2014-2024^{a}$ (n = 350)



Number of isolates

- AT: Austria; CZ: Czechia; DE: Germany; DK: Denmark; FI: Finland; FR: France; HU: Hungary; IE: Ireland; LU: Luxembourg; NL: the Netherlands; NO: Norway; PT: Portugal; SE: Sweden; TR: Türkiye. AD: allelic difference; NA: not applicable, even though the ST13730 isolates branch out within ST131 clade A; ST: sequence type.
- The plot includes only isolates within clusters (grey) with the largest clusters containing more than ten isolates coloured by reporting country in panels above. Within cluster diversity is described as median allelic differences and range.
- <sup>a</sup> Three isolates without information on year of sampling were not marked in the plots.
- <sup>b</sup> *bla*<sub>KPC-31</sub> listed in Beta-Lactamase DataBase (http://bldb.eu/, accessed 28 October 2024) as carbapenemase, but reported in literature as extended spectrum beta-lactamase not conferring carbapenem resistance [Haidar G, Clancy CJ, Shields RK, Hao B, Cheng S, Nguyen MH. Mutations in *bla*<sub>KPC-3</sub> that confer ceftazidimeavibactam resistance encode novel KPC-3 variants that function as extended-spectrum β-lactamases. Antimicrob Agents Chemother. 2017;61(5):e02534-16. <u>http://dx.doi.org/10.1128/AAC.02534-16</u> <u>PMID:28223379</u>].

<sup>c</sup> An isolate from Türkiye clustered within 2024-08.ECOLI.59.ST131, but did not carry *bla*<sub>OXA-48</sub>.