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Corrigendum: *In silico* serotyping of *E. coli* from short read data identifies limited novel O-loci but extensive diversity of O:H serotype combinations within and between pathogenic lineages

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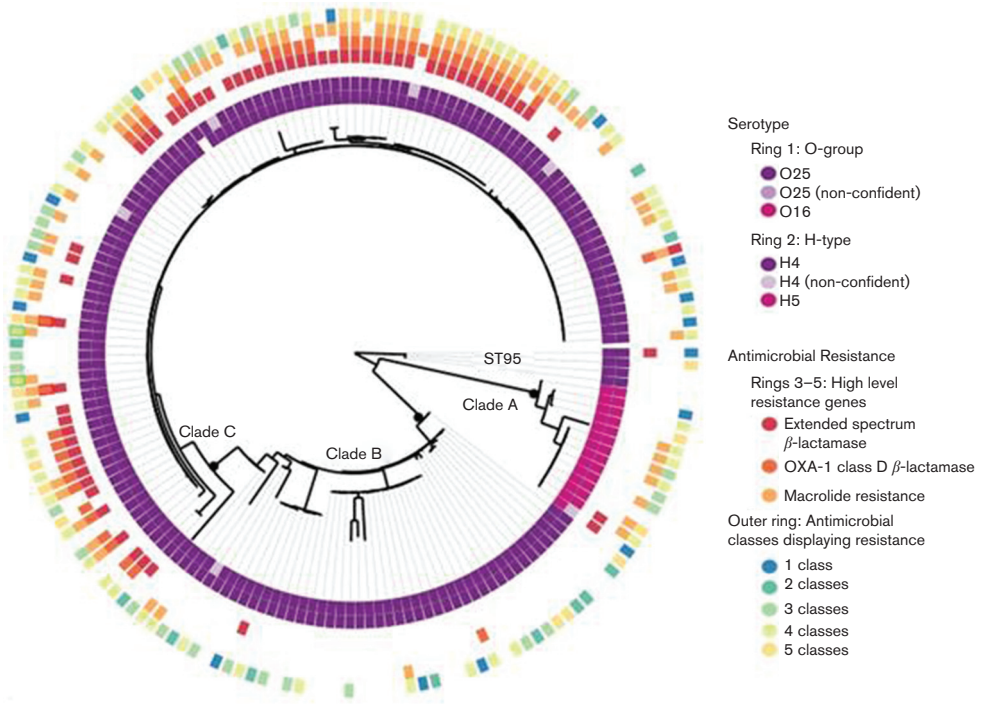
Microbial Genomics 2016;2, doi: 10.1099/mgen.0.000064

There was an error in Fig. 1 in the published article. In the key, under Antimicrobial Resistance, the orange colour indicates the presence of the OXA-1 class D β-lactamase (not Carbapenemases as originally labelled).

The authors apologise for any inconvenience caused.



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