

Erratum for Ben Zakour et al., Sequential Acquisition of Virulence and Fluoroquinolone Resistance Has Shaped the Evolution of *Escherichia coli* ST131

 Nouri L. Ben Zakour,^{a,b}  Areej S. Alsheikh-Hussain,^{a,b}  Melinda M. Ashcroft,^{a,b}  Nguyen Thi Khanh Nhu,^{a,b}  Leah W. Roberts,^{a,b}  Mitchell Stanton-Cook,^{a,b}  Mark A. Schembri,^a  Scott A. Beatson^{a,b}

Australian Infectious Diseases Research Centre, School of Chemistry and Molecular Biosciences, The University of Queensland, Brisbane, Australia^a; Australian Centre for Ecogenomics, School of Chemistry and Molecular Biosciences, The University of Queensland, Brisbane, Australia^b

A.S.A.H., M.M.A., and N.T.K.N. contributed equally to this article.

The article cited as reference 17 (N. Stoesser et al.) was originally available as a preprint in BioRxiv (<http://dx.doi.org/10.1101/030668>) on 6 November 2015. It was subsequently published in mBio while our article was in proof.

Published 28 June 2016

Citation Ben Zakour NL, Alsheikh-Hussain AS, Ashcroft MM, Khanh Nhu NT, Roberts LW, Stanton-Cook M, Schembri MA, Beatson SA. 2016. Erratum for Ben Zakour et al., Sequential acquisition of virulence and fluoroquinolone resistance has shaped the evolution of *Escherichia coli* ST131. mBio 7(3):e00958-16. doi:10.1128/mBio.00958-16.

Copyright © 2016 Ben Zakour et al. This is an open-access article distributed under the terms of the [Creative Commons Attribution 4.0 International license](https://creativecommons.org/licenses/by/4.0/).

Address correspondence to Mark A. Schembri, m.schembri@uq.edu.au, or Scott A. Beatson, s.beatson@uq.edu.au.