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Edited and reviewed by:

Matt Arthur Baker,
University of New South
Wales, Australia

***Correspondence:**

Morgan Beeby
mbeeby@imperial.ac.uk

†ORCID:

Josie L. Ferreira
orcid.org/0000-0002-4411-6131

Izaak Coleman
orcid.org/0000-0003-4697-6079

Tobias Zachs
orcid.org/0000-0002-0836-0989

Bonnie L. Quigley
orcid.org/0000-0003-3787-0993

Morgan Beeby
orcid.org/0000-0001-6413-9835

‡Present address:

Josie L. Ferreira,
Centre for Structural Systems Biology,
Heinrich-Pette-Institut, Leibniz-Institut
für Experimentelle Virologie, Hamburg,
Germany

Izaak Coleman,
Department of Systems Biology,
Columbia University, Irving Cancer
Research Center, New York, NY,
United States

Max L. Addison,
Warwick Medical School, Microbiology
and Infection, University of Warwick,
Coventry, United Kingdom

Tobias Zachs,
Institute of Molecular Biology and
Biophysics, Eidgenössische
Technische Hochschule Zürich,
Zurich, Switzerland

Bonnie L. Quigley, Provectus Algae
Pty Ltd., Noosaville, QLD, Australia

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Corrigendum: The “Jack-of-all-Trades” Flagellum From *Salmonella* and *E. coli* Was Horizontally Acquired From an Ancestral β -Proteobacterium

Josie L. Ferreira^{1†}, Izaak Coleman^{1†}, Max L. Addison^{1‡}, Tobias Zachs^{1†‡},
Bonnie L. Quigley^{1†‡}, Kristin Wuichet² and Morgan Beeby^{1*†}

¹ Department of Life Sciences, Imperial College London, London, United Kingdom, ² Department of Biomedical Informatics, Vanderbilt University Medical Center, Nashville, TN, United States

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A Corrigendum on

The “Jack-of-all-Trades” Flagellum From *Salmonella* and *E. coli* Was Horizontally Acquired From an Ancestral β -Proteobacterium

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In the original article, there was a mistake in **Figure 1** as published. **We inadvertently uploaded an outdated version of this figure.** The corrected **Figure 1** appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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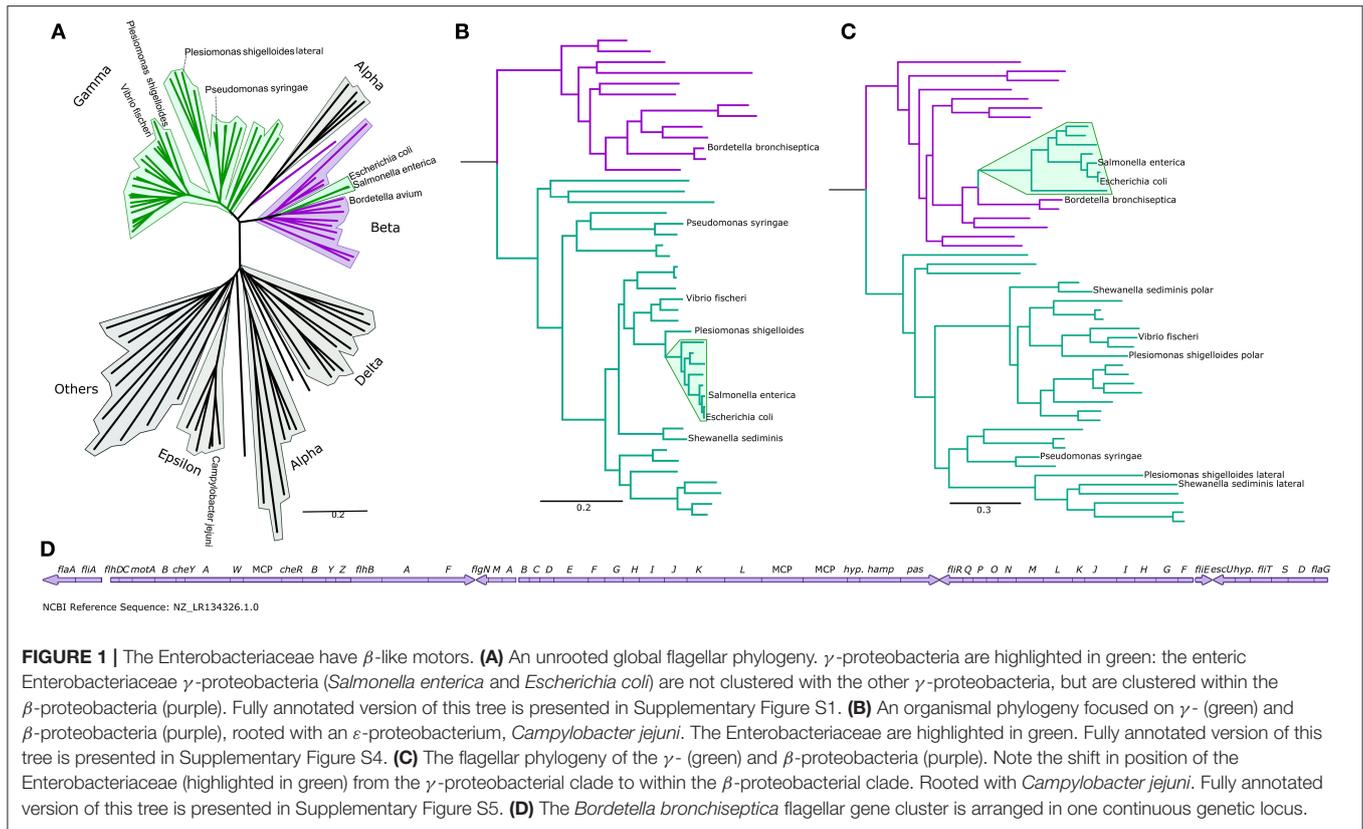


FIGURE 1 | The Enterobacteriaceae have β -like motors. **(A)** An unrooted global flagellar phylogeny. γ -proteobacteria are highlighted in green: the enteric Enterobacteriaceae γ -proteobacteria (*Salmonella enterica* and *Escherichia coli*) are not clustered with the other γ -proteobacteria, but are clustered within the β -proteobacteria (purple). Fully annotated version of this tree is presented in Supplementary Figure S1. **(B)** An organismal phylogeny focused on γ - (green) and β -proteobacteria (purple), rooted with an ϵ -proteobacterium, *Campylobacter jejuni*. The Enterobacteriaceae are highlighted in green. Fully annotated version of this tree is presented in Supplementary Figure S4. **(C)** The flagellar phylogeny of the γ - (green) and β -proteobacteria (purple). Note the shift in position of the Enterobacteriaceae (highlighted in green) from the γ -proteobacterial clade to within the β -proteobacterial clade. Rooted with *Campylobacter jejuni*. Fully annotated version of this tree is presented in Supplementary Figure S5. **(D)** The *Bordetella bronchiseptica* flagellar gene cluster is arranged in one continuous genetic locus.