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Whole-genome sequencing of *Campylobacter jejuni* isolated from Danish routine human stool samples reveals surprising degree of clustering

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1 Whole-genome sequencing of Campylobacter jejuni isolated from Danish routine human stool samples 2 reveals surprising degree of clustering 3 4 Joensen KG<sup>1</sup>8, Kuhn KG<sup>2</sup>8\*, Müller L<sup>2</sup>, Björkman JT<sup>1</sup>, Torpdahl M<sup>1</sup>, Engberg J<sup>3</sup>, Holt HM<sup>4</sup>, Nielsen HL<sup>5</sup>, 5 Petersen AM<sup>6,7</sup>, Ethelberg S<sup>2</sup>, Nielsen EM<sup>1</sup>. 6 1. Department of Bacteria, Parasites & Fungi, Statens Serum Institut, Copenhagen, Denmark 7 2. Department of Infectious Disease Epidemiology, Statens Serum Institut, Copenhagen, Denmark 8 3. Department of Clinical Microbiology, Slagelse Hospital, Slagelse, Denmark. 9 4. Department of Clinical Microbiology, Odense University Hospital, Odense, Denmark 10 5. Department of Clinical Microbiology, Aalborg University Hospital, Aalborg, Denmark 11 6. Department of Clinical Microbiology, Hvidovre University Hospital, Hvidovre, Denmark 12 7. Department of Gastroenterology, Hvidovre University Hospital, Hvidovre, Denmark 13 8. These authors contributed equally as first authors 14

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