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Whole-genome Comparison of Urinary Pathogenic *Escherichia coli* and Faecal Isolates of UTI Patients and Healthy Controls

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Short running Title: Whole-genome comparison of faecal and urinary *E. coli*

Abstract

The faecal flora is a common reservoir for urinary tract infection (UTI), and *E. coli* is frequently found in this reservoir without causing extraintestinal infection. We investigated these *E. coli*

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