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# Corrigendum: Patterns of within-host spread of *Chlamydia trachomatis* between vagina, endocervix and rectum revealed by comparative genomic analysis

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## KEYWORDS

*Chlamydia trachomatis*, single nucleotide polymorphisms, single variable polymorphisms, sexually transmitted diseases, chlamydiae

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In the published article, there was an error in [Figure 2](#) as published. Some of the colors in the “ompA genotype” column of the heatmap did not match the true genotype of strain in the phylogenetic tree. The corrected [Figure 2](#) and its caption appear 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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