## **ERRATUM**

**Lupolova** *et al.*, *Microbial Genomics* 2018;4
DOI 10.1099/mgen.0.000193





## Erratum: Patchy promiscuity: machine learning applied to predict the host specificity of *Salmonella enterica* and *Escherichia coli*

Nadejda Lupolova, <sup>1</sup> Tim J. Dallman, <sup>2</sup> Nicola J. Holden<sup>3</sup> and David L. Gally<sup>4,\*</sup>

Microbial Genomics 2017;3, doi: 10.1099/mgen.0.000135

An error occurred during the publishing process of this article.

There was text inserted in the final paragraph of the Discussion, in the following sentence:

'We consider that machine learning has tremendous potential to interrogate complex seqLineColumnRule IDProbe Message-Node TextNode XpathParent Node Textfatal/var/www/html/\_default/resources/microbio/\_\_package/144333/144333. xmlf002block-formatting check: Entire content of title should not be formatted (Tagging Guidelines)Salmonella entericauence datasets and identify genes/sequences associated with host specificity.'

The sentence should read as follows:

'We consider that machine learning has tremendous potential to interrogate complex sequence datasets and identify genes/ sequences associated with host specificity.'

The Microbiology Society apologizes for any inconvenience caused.

Received 1 June 2018; Accepted 1 June 2018

Author affiliations: <sup>1</sup>University of Edinburgh, Edinburgh, UK; <sup>2</sup>Public Health England, England, UK; <sup>3</sup>James Hutton Institute, Dundee, UK; <sup>4</sup>Division of Immunity and Infection, The Roslin Institute, University of Edinburgh, Easter Bush, Edinburgh EH25 9RG, UK. \*Correspondence: David L. Gally, dgally@ed.ac.uk