



CORRECTION

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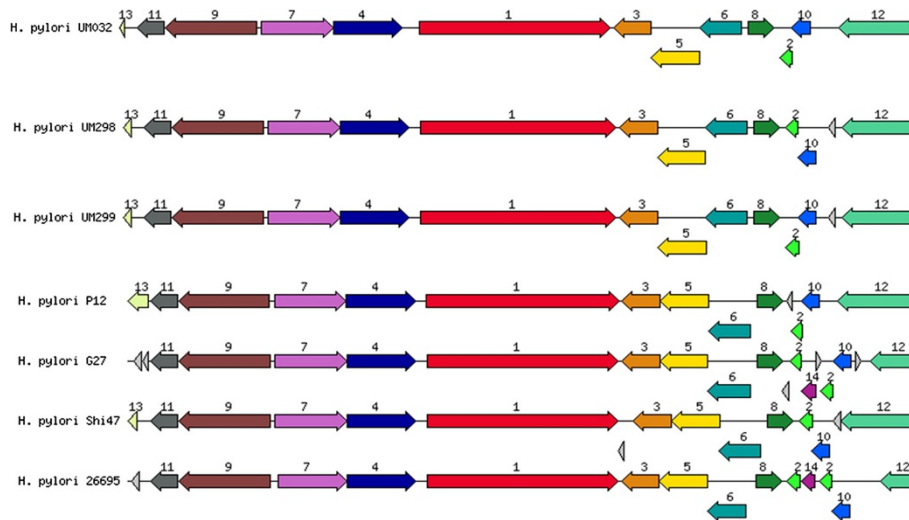
# Correction: Comparing the genomes of *Helicobacter pylori* clinical strain UM032 and mice-adapted derivatives

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### Correction

After publication of this work [1], we discovered that due to an inadvertent mistake, Figures 4 and 5 in the original

article were essentially the same figure. Amendment to Figure 5 (Figure 1 here) has now been added. We offer our sincerest apologies for this oversight.



**Figure 1 Genetic relatedness of *vacA* cluster with closely related bacteria.** 1: vacuolating cytotoxin, 2: hypothetical protein, 3: haemin uptake system ATP-binding protein, 4: cysteinyl-Trna-SYNTHEASE, 5: IRON III, 6: dehydrogenases with different specificities, 7: proposed peptidoglycan lipid, 8: hypothetical protein, 9: hypothetical protein, 10: DNA damage inducible protein J, 11: holliday junction DNA helicase RUUA, 12: putative outer membrane protein, 13: hypothetical protein.

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Received: 28 April 2014 Accepted: 29 April 2014

Published: 8 May 2014

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doi:10.1186/1757-4749-6-11

**Cite this article as:** Khosravi *et al.*: Correction: Comparing the genomes of *Helicobacter pylori* clinical strain UM032 and mice-adapted derivatives. *Gut Pathogens* 2014 **6**:11.

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