CORRECTION

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Correction to: *Neisseria lactamica* Y92–1009 (Correction to: *Neisseria lactamica* Y92–1009)

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Correction

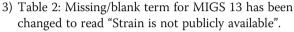
After publication of the original article [1] it was identified that an incorrect version of the manuscript has been published. This was caused by a technical error which led to a discrepancy between the editorially-accepted version of the manuscript, and the published version.

A list of Corrections is included below:

- The following abbreviations have been corrected throughout the article: *Nla* changed to *N. lactamica Nme* changed to *N. meningitides N.go* changed to *N. gonorrhoeae Nci* changed to *N. cinerea*
- 2) Table 1: 'Term' and 'Evidence code' columns of the first 7 rows have been updated to the following:

Domain Bacteria	TAS [37]
Phylum Proteobacteria	TAS [38]
Class Betaproteobacteria	TAS [39, 40]
Order Neisseriales	TAS [39, 41]
Family - Neisseriaceae	TAS [42]
Genus Neisseria	TAS [42]
Species Neisseria lactamica	TAS [42, 43]

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- 4) 2nd paragraph under heading "Insights from the genome sequence": Sentence containing "... approximately a third (33.5%) of all open reading frames (ORFs) in this genome..." has been shortened to "approximately a third (33.5%) of all ORFs in this genome..."
- 5) 3rd paragraph under heading "Insights from the genome sequence": Bracketed abbreviation '(DUS)' has been removed.
- 6) 3rd paragraph under heading "Insights from the genome sequence": Bracketed abbreviation '(CREEs)' has been removed
- Bracketed link for PHAST '(http:// phast.wishartlad.com/index.html)' changed to reference 35.
- 8) Reference 14 initially missing from article, has now been added and correctly referenced
- 9) Reference 33 was removed, as it was a duplicate of reference 30.
- 10) References 36–40 removed, and replaced with updated reference list (36–46):

36. Field D, Garrity G, Gray T, Morrison N, Selengut J, Sterk P, et al. The minimum information about a genome sequence (MIGS) specification. Nat Biotechnol. 2008;26:541–7.

37. Woese CR, Kandler O, Wheelis ML. Towards a natural system of organisms: proposal for the domains *Archaea, Bacteria,* and *Eucarya.* Proc Natl Acad Sci. 1990;87:4576–9. https://doi.org/10.1073/pnas.87.12.4576.

38. Garrity GM, Holt JG. The Road Map to the Manual. In: Bergey's Manual[®] of Systematic Bacteriology. 2001. p. 119–66. https://doi.org/10.1007/978-0-387-21609-6_15.

39. Euzeby. J. List of new names and new combinations previously effectively, but not validly, published. Int J Syst Evol Microbiol. 2006;56:1–6.



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40. Garrity GM, Bell JA, Lilburn T. Class II. *Betaproteobacteria* class. Nov. In: Bergey's Manual[®] of Systematic Bacteriology. 2005. p. 575–922. https://doi.org/10.1007/0-387-29298-5_2.

41. Tønjum T. *Neisseriales* ord. Nov. In: Bergey's Manual of Systematics of Archaea and Bacteria. John Wiley & Sons, Ltd.; 2015. https://doi.org/10.1002/9781118960608.obm00079.

42. SKERMAN VBD, McGOWAN V, SNEATH PHA. Approved Lists of Bacterial Names. Int J Syst Evol Microbiol. 1980;30:225–420.

43. Hollis DG, Wiggins GL, Weaver RE. *Neisseria lactamicus* sp. n., a lactose-fermenting species resembling Neisseria meningitidis. Appl Microbiol. 1969;17:71–7.

44. Aho EL, Keating a M, McGillivray SM. A comparative analysis of pilin genes from pathogenic and nonpathogenic Neisseria species. Microb Pathog. 2000;28:81–8.

45. Lennette, Edwin H. Manual of Clinical Microbiology. 4th edition. Washington DC: American Society for Microbiology; 1985.

46. Neisseria MLST website. https://pubmlst.org/ neisseria/. Accessed 2 Feb 2017.

The Publisher apologises for this error.

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Reference

 Pandey AK, Cleary DW, Laver JR, Maiden MCJ, Didelot X, Gorringe A, Read RC. *Neisseria lactamica* Y92–1009 complete genome sequence. Stand Genomic Sci. 2017;12:41. https://doi.org/10.1186/s40793-017-0250-6.