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CORRECTION



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Correction: Proteomic analysis of the copper resistance of *Streptococcus pneumoniae*

Zhong Guo, Junlong Han, Xiao-Yan Yang, Kun Cao, Ke He, Gaofei Du, Guandi Zeng, Liang Zhang, Guangchuang Yu, Zhenghua Sun, Qing-Yu He* and Xuesong Sun*

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Correction for 'Proteomic analysis of the copper resistance of *Streptococcus pneumoniae*' by Zhong Guo *et al., Metallomics*, 2015, **7**, 448–454.

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The sentence on p. 1 beginning 'Copper resistance in...' should read 'Copper resistance in *S. pneumoniae* is mainly mediated by a single operon encoding a copper-dependent repressor protein (CopY),¹¹ a copper transporting ATPase (CopA)^{7,12} and a copper metallochaperone (CupA).⁷

The sentence on p. 1 beginning 'The third gene...' should read 'The third gene encodes a structurally characterized, membrane anchored cupredoxin-fold protein, designated CupA,¹ which binds two Cu(i) ions with high affinity and is essential for maintenance of cellular copper resistance.⁷

References

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- 12 A. Reyes, A. Leiva, V. Cambiazo, M. A. Mendez and M. Gonzalez, *Biol. Res.*, 2006, **39**, 87–93. The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

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Key Laboratory of Functional Protein Research of Guangdong Higher Education Institutes, Institute of Life and Health Engineering, College of Life Science and Technology, Jinan University, Guangzhou 510632, China. E-mail: tqyhe@jnu.edu.cn; Fax: +86-20-85227039; Tel: +86-20-85227039