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

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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.

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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The Royal Society of Chemistry apologises for these errors and any consequent inconvenience to authors and readers.

