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), a copper transporting ATPase (CopA) and a copper metallochaperone (CupA).’

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, which binds two Cu(II) ions with high affinity and is essential for maintenance of cellular copper resistance.’

References


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