CORRIGENDUM

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Branched tricarboxylic acid metabolism in *Plasmodium falciparum*

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The samples used for histone proteomics described in this Letter were inadvertently switched, such that the U-^{13}C-glucose and U-^{13}C-^{15}N-glutamine data were inverted. The plots in Fig. 2b and the spectra in Supplementary Fig. 3 have been modified to reflect this. The corrected results demonstrate that ^{13}C-labelling of histone acetyl groups occurs only in cells grown on ^{13}C-glucose and not on ^{13}C-glutamine. Therefore, glucose is the primary source of the acetyl units used for both amino sugar biosynthesis and nuclear protein acetylation. Although U-^{13}C-^{15}N-glutamine does give rise to labelled acetyl-CoA, its localization and function remain unclear. The model presented in Fig. 4 has been modified to reflect these facts, which do not alter the paper's main conclusions about TCA cycle architecture. The corrected Figs 2b and 4 are shown below. The authors apologize for this error.

Supplementary Information is linked to the online version of the paper at www.nature.com/nature.

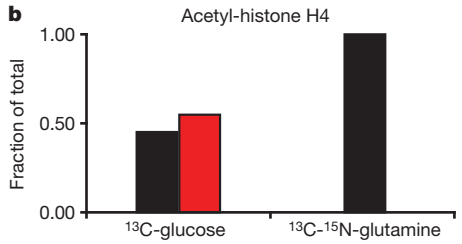

Figure 2

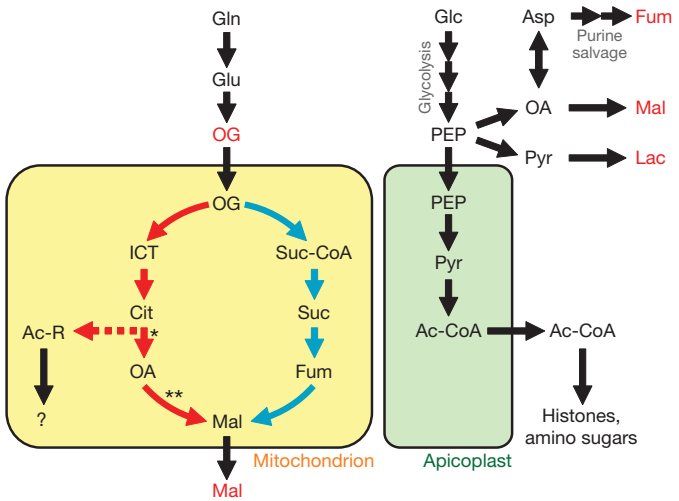

Figure 4