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Correction for Sandai et al., The Evolutionary Rewiring of Ubiquitination Targets Has Reprogrammed the Regulation of Carbon Assimilation in the Pathogenic Yeast *Candida albicans*

Doblin Sandai
University of Aberdeen

Zhikang Yin
University of Aberdeen

Laura Selway
University of Aberdeen

David Stead
University of Aberdeen

Janet Walker
University of Aberdeen

Sandai, Doblin; Yin, Zhikang; Selway, Laura; Stead, David; Walker, Janet; Leach, Michelle D.; Bohovych, Iryna; Ene, Iuliana V.; Kastora, Stavroula; Budge, Susan; Munro, Carol A.; Odds, Frank C.; Gow, Neil A.R.; and Brown, Alistair J.P., "Correction for Sandai et al., The Evolutionary Rewiring of Ubiquitination Targets Has Reprogrammed the Regulation of Carbon Assimilation in the Pathogenic Yeast *Candida albicans*" (2015). *Biochemistry -- Faculty Publications*. 265.
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Authors

Doblin Sandai, Zhikang Yin, Laura Selway, David Stead, Janet Walker, Michelle D. Leach, Iryna Bohovych, Iuliana V. Ene, Stavroula Kastora, Susan Budge, Carol A. Munro, Frank C. Odds, Neil A.R. Gow, and Alistair J.P. Brown

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Doblin Sandai,* Zhikang Yin, Laura Selway, David Stead, Janet Walker, Michelle D. Leach,* Iryna Bohovych,* Iuliana V. Ene, Stavroula Kastora, Susan Budge, Carol A. Munro, Frank C. Odds, Neil A. R. Gow, Alistair J. P. Brown

School of Medical Sciences, University of Aberdeen, Institute of Medical Sciences, Foresterhill, Aberdeen, United Kingdom

* Present address: Doblin Sandai, Institut Perubatan & Pergigian Termaju, Universiti Sains Malaysia, Pulau Pinang, Malaysia; Michelle D. Leach, Department of Molecular Genetics, University of Toronto, Toronto, Canada; Iryna Bohovych, Nebraska Redox Biology Center, University of Nebraska-Lincoln, Lincoln, Nebraska, USA.

Volume 3, no. 6, doi:10.1128/mBio.00495-12, 2012. An error has been identified in Fig. 8B, where the wrong Western blot was used inadvertently. Figure 8B should appear as shown below. This change does not affect the conclusions in any way.

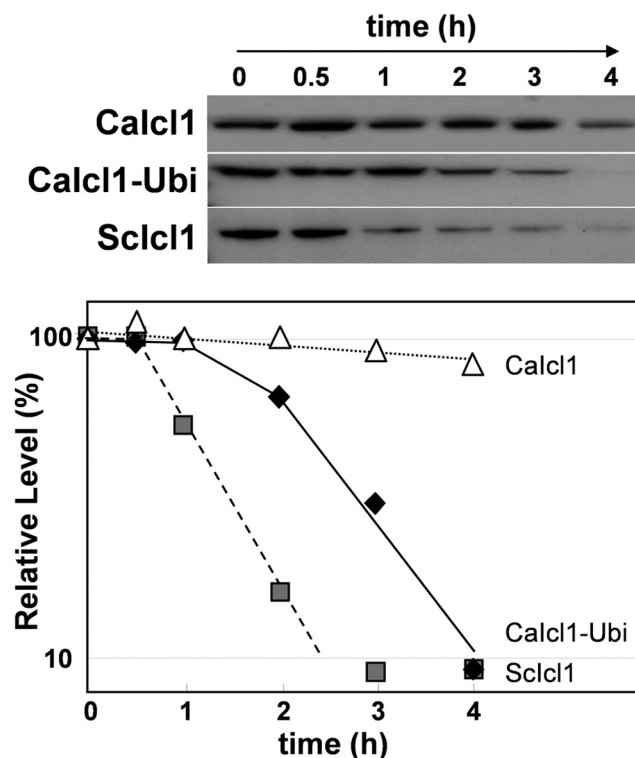


FIG 8 Addition of a consensus ubiquitin site stimulates glucose-accelerated degradation of CaIc1 in *C. albicans*. (B) The carboxy-terminal ubiquitination site from ScIc1 was fused to CaIc1 to create CaIc1-Ubi-Myc in *C. albicans* DSC04 (Table S1). These cells were grown on lactate, and the levels of CaIc1-Ubi-Myc were assayed by Western blotting after glucose addition. As controls, the stabilities of CaIc1-Myc (CA1395; open diamonds) and ScIc1-Myc (DSC01; gray squares) in *C. albicans* were compared under equivalent conditions. CaIc1-Ubi-Myc, ScIc1-Myc, and CaIc1-Myc levels are expressed as a percentage of their abundance at time zero (100%). Similar data were obtained from two independent replicate experiments.

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Address correspondence to Alistair J. P. Brown, brown@abdn.ac.uk.