Erratum to: Label-Free Quantitative Proteomic Profiling Identifies Potential Active Components to Exert Pharmacological Effects in the Fruit of *Alpinia oxyphylla* by Mass Spectrometry

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Unfortunately Fig. 2 is incorrect. The correct Fig. 2 is given below.

The original article can be found online at <https://doi.org/10.1007/s12374-020-09251-9>.

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Fig. 2 Volcano plots showing the fold change of differential proteins identified between two *A. oxyphylla* fruit groups. **a** The Metaphase group vs the Earliestage group; **b** the Advanced group vs the Earliestage group; **c** the Advanced group vs the Metaphase group