Correction



Correction: Cysteine modification reveals an allosteric inhibitory site on the CAL PDZ domain

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After acceptance and the publication of the Accepted 3, follow-up experiments revealed unexpected variability in the binding constants obtained by the mixed- K_D analysis of FP binding data as shown in Table 2 in the Accepted Manuscript. Further investigation using non-reducing SDS–PAGE found previously undetected PDZ–PDZ dimerization triggered by extended incubation with the small-molecule compounds under stringent labeling conditions. This dimerization is not accounted for by our mixed- K_D model. The final Version of Record instead reports apparent K_D values determined from FP titrations with PDZ domains treated to minimize PDZ–PDZ dimerization while preserving high levels of small-molecule adduct formation, together with associated changes in text.

This has been approved by the Editorial Board of *Bioscience Reports*.

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