

Structural basis of sialidase in complex with geranylated flavonoids as potent natural inhibitors. Corrigendum

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A correction is made to the article by Lee *et al.* [(2014) *Acta Cryst. D* **70**, 1357–1365].

The authors of Lee *et al.* (2014) have made a careful re-analysis of all of data sets used in their structure determination of the putative complex between the human pathogen *Clostridium perfringens* sialidase and the natural inhibitory compound, geranylated flavonoid diplacone, described in the article. They now no longer believe that their structure (PDB accession number: 4l2e) is supported by X-ray diffraction data and thus have asked for the 4l2e entry to be made obsolete in the Protein Data Bank. This does not affect the validity of the biochemical characterization of specific interactions of the sialidase with diplacone and other naturally occurring compounds reported in the paper.

References

Lee, Y., Ryu, Y. B., Youn, H.-S., Cho, J. K., Kim, Y. M., Park, J.-Y., Lee, W. S., Park, K. H. & Eom, S. H. (2014). *Acta Cryst. D* **70**, 1357–1365.