

## Corrections

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Cyclostreptin (FR182877), an Antitumor Tubulin-Polymerizing Agent Deficient in Enhancing Tubulin Assembly Despite Its High Affinity for the Taxoid Site, by Michael C. Edler, Rubén M. Buey, Rick Gussio, Adam I. Marcus, Christopher D. Vanderwal, Erik J. Sorensen, J. Fernando Díaz, Paraskevi Giannakakou, and Ernest Hamel\*, Volume 44, Number 34, August 30, 2005, pages 11525–11538.

In this paper, we had intended to use the actual residue positions of  $\beta$ -tubulin (1), as opposed to the maximum homology positions used in the deposited tubulin structure (2) in PDB entry 1TUB. However, we neglected to take into account the 8-position shift after Pro-258 (1), termed Pro-260 (2) in 1TUB. Therefore, on pages 11535 and 11536, including Figure 10 and its legend, and in the table of contents graphic, all notations of Arg-367, Gly-368, and Leu-369 should be Arg-359, Gly-360, and Leu-361, respectively. In addition, on page 11536, right column, three paragraphs from the bottom, Phe-228 should be deleted since the actual amino acid residue at that position, leucine, is already noted in the same sentence.

### REFERENCES

1. Krauhs, E., Little, M., Kempf, T., Hofer-Warbinek, R., Ade, W., and Ponstingl, H. (1981) Complete amino acid sequence of  $\beta$ -tubulin from porcine brain, *Proc. Natl. Acad. Sci. U.S.A.* 78, 4156–4160.
2. Nogales, E., Wolf, S. G., and Downing, K. H. (1998) Structure of the  $\alpha\beta$  tubulin dimer by electron crystallography, *Nature* 391, 199–203.

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