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## corrections

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### Structure of the *Bacillus subtilis* D-aminopeptidase DppA reveals a novel self-compartmentalizing protease

Han Remaut, Coralie Bompard-Gilles, Colette Goffin, Jean-Maire Frère and Jozef Van Beeumen

*Nature Struct. Biol.* 8, 674–678 (2001).

Part of the reference list was incorrectly printed in this paper. The first ref. 12 in the list should be renumbered as ref. 11; the original ref. 11 in the list should be deleted. The citation of these references in the text corresponds to the corrected numbering scheme. Citations for MLPHARE and DM in footnotes 2 and 3 of Table 1 were also incorrectly printed. The correct citation for both is ref. 27. We apologize for any confusion these may have caused.

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### Structural insights into the hydrolysis of cellular nitric oxide synthase inhibitors by dimethylarginine dimethylaminohydrolase

Judith Murray-Rust, James Leiper, Mark McAlister, John Phelan, Sarah Tilley, Jo Santa Maria, Patrick Vallance and Neil McDonald

*Nature Struct. Biol.* 8, 679–683 (2001).

The genus name for the organism from which dimethylarginine dimethylaminohydrolase is derived was printed incorrectly in this paper. The correct name of the organism is *Pseudomonas aeruginosa*. We apologize for any confusion this may have caused.

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## erratum

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### Structural basis for substrate specificities of cellular deoxyribosenucleoside kinases

Kenneth Johansson, S. Ramaswamy, Catarina Ljungcrantz, Wolfgang Knecht, Jure Piškur, Birgitte Munch-Petersen, Staffan Eriksson and Hans Eklund

*Nature Struct. Biol.* 8, 616–620 (2001).

A production error resulted in several mistakes in the Tables of this paper. In Table 1, the amino acid type of several residues was incorrectly printed; one table heading was incorrectly aligned. For clarity, the corrected Table 1 is reprinted on the next page. The dimension of the a-axis for the native dNK crystal is incorrectly printed in Table 2. The correct value is  $a = 119.5 \text{ \AA}$ . We apologize for any confusion these may have caused.