

Erratum: ABC ATPase signature helices in Rad50 link nucleotide state to Mre11 interface for DNA repair

Gareth J Williams, R Scott Williams, Jessica S Williams, Gabriel Moncalian, Andrew S Arvai, Oliver Limbo, Grant Guenther, Soumita SilDas, Michal Hammel, Paul Russell & John A Tainer
Nat. Struct. Mol. Biol. **18**, 423–431 (2011); published online 27 March 2011; corrected after print 21 September 2011

In the version of this article initially published, panels in Figures 2b (middle row, Dex-LWH), 3b (0.1 μM CPT) and 3d (0.1 μM CPT) were mistakenly replaced with duplicates of adjacent panels. The errors have been corrected in the HTML and PDF versions of the article.

Corrigendum: miRNA-mediated deadenylation is orchestrated by GW182 through two conserved motifs that interact with CCR4–NOT

Marc R Fabian, Maja K Cieplak, Filipp Frank, Masahiro Morita, Jonathan Green, Tharan Srikumar, Bhushan Nagar, Tadashi Yamamoto, Brian Raught, Thomas F Duchaine & Nahum Sonenberg
Nat. Struct. Mol. Biol. **18**, 1211–1217 (2011); published online 7 October 2011; corrected after print 8 November 2011

In the version of this article initially published, several acknowledgments were missing. The end of the section should read as follows: "...N.S., who was also supported by a Howard Hughes Medical Institute (HHMI) International Scholarship. Postdoctoral support was from the Terry Fox Foundation of the Canadian Cancer Society to M.R.F. T.F.D. was supported through a CIHR grant. M.K.C. was supported by the Foundation for Polish Science International PhD Projects. F.F. was supported by a Boehringer Ingelheim Fonds PhD fellowship." The error has been corrected in the HTML and PDF versions of the article.

Corrigendum: Transcriptome-wide sequencing reveals numerous APOBEC1 mRNA-editing targets in transcript 3' UTRs

Brad R Rosenberg, Claire E Hamilton, Michael M Mwangi, Scott Dewell & F Nina Papavasiliou
Nat. Struct. Mol. Biol. **18**, 230–236 (2011); published online 23 January 2011; corrected after print 14 December 2011

In the version of this article initially published, financial support from the Starr Foundation was not acknowledged. The error has been corrected in the HTML and PDF versions of the article.

Erratum: Crystal structure of a monomeric retroviral protease solved by protein folding game players

Firas Khatib, Frank DiMaio, Foldit Contenders Group, Foldit Void Crushers Group, Seth Cooper, Maciej Kazmierczyk, Miroslaw Gilski, Szymon Krzywda, Helena Zabranska, Iva Pichova, James Thompson, Zoran Popović, Mariusz Jaskolski & David Baker
Nat. Struct. Mol. Biol. **18**, 1175–1177 (2011); published online 18 September 2011; corrected after print 20 December 2011

In the version of this article initially published, the location for the Academy of Sciences of the Czech Republic was given as Poznan. The Academy is in Prague. The error has been corrected in the HTML and PDF versions of the article.

Corrigendum: Human mitochondrial transcription factor A induces a U-turn structure in the light strand promoter

Anna Rubio-Cosials, Jasmin F Sidow, Nereida Jiménez-Menéndez, Pablo Fernández-Millán, Julio Montoya, Howard T Jacobs, Miquel Coll, Pau Bernadó & Maria Solà
Nat. Struct. Mol. Biol. **18**, 1281–1289 (2011); published online 30 October 2011; corrected after print 20 December 2011

In the version of this article initially published, the name of the second author was misspelled. The error has been corrected in the HTML and PDF versions of the article.