Corrigendum

RepeatsDB 2.0: improved annotation, classification, search and visualization of repeat protein structures

Lisanna Paladin^{1,†}, Layla Hirsh^{1,2,†}, Damiano Piovesan¹, Miguel A. Andrade-Navarro³, Andrey V. Kajava^{4,5,6} and Silvio C.E. Tosatto^{1,7,*}

¹Dept. of Biomedical Sciences, University of Padua, 35121 Padova, Italy, ²Departamento de Ingeniería, Pontificia Universidad Católica del Perú, 32 Lima, Perú, ³Institute of Molecular Biology, Faculty of Biology, Johannes Gutenberg University of Mainz, 55128 Mainz, Germany, ⁴Centre de Recherches de Biochimie Macromoléculaire, CNRS, Université Montpellier, 34293 Montpellier, France, ⁵Institut de Biologie Computationnelle (IBC), 34293 Montpellier, France, ⁶Institute of Bioengineering, University ITMO, 197101 St. Petersburg, Russia and ⁷CNR Institute of Neuroscience, 35121 Padova, Italy

Nucl. Acids Res. (2016) doi: 10.1093/nar/gkw1136

The Authors wish it to be known that, in their opinion, the first two authors should be regarded as joint First Authors.

^{*}To whom correspondence should be addressed. Tel: +39 0498276269; Fax: +39 0498276260; Email: silvio.tosatto@unipd.it

These authors contributed equally to the paper as first authors.

[©] The Author(s) 2016. Published by Oxford University Press on behalf of Nucleic Acids Research.