Corrigendum

Gene3D: Extensive prediction of globular domains in proteins

Tony E. Lewis^{1,†}, Ian Sillitoe^{1,†}, Natalie Dawson¹, Su Datt Lam^{1,2}, Tristan Clarke¹, David Lee³, Christine Orengo¹ and Jonathan Lees^{1,4,*}

¹Institute of Structural and Molecular Biology, Division of Biosciences, University College London, Gower Street, London WC1E 6BT, UK, ²School of Biosciences and Biotechnology, Faculty of Science and Technology, University Kebangsaan Malaysia, 43600 Bangi, Selangor, Malaysia, ³Bristol Life Sciences Building, University of Bristol, Bristol Life Sciences Building, Bristol, BS8 1TQ, UK and ⁴Oxford Brookes University, Faculty of Health and Life Sciences, Oxford, Oxfordshire, UK

Nucleic Acids Research (2017), https://doi.org/10.1093/nar/gkx1069

The Authors would like to apologise for some small content errors in their text, including a mention on page 2 that the number of S35 domain sequences had been expanded from '2155 to over 32 000'. This should be '21 155 to over 32 000'. A spelling error in the name of Tony E. Lewis was also adapted. The paper has now been corrected online.

*To whom correspondence should be addressed. Tel: +44 2076793890; Fax: +44 2076797193; Email: jonathan.lees@ucl.ac.uk [†]These authors contributed equally to the paper as first authors.

© The Author(s) 2017. Published by Oxford University Press on behalf of Nucleic Acids Research. This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/4.0/), which permits unrestricted reuse, distribution, and reproduction in any medium, provided the original work is properly cited.