

CORRIGENDUM

InterPro in 2011: new developments in the family and domain prediction database

Sarah Hunter, Philip Jones, Alex Mitchell, Rolf Apweiler, Teresa K. Attwood, Alex Bateman, Thomas Bernard, David Binns, Peer Bork, Sarah Burge, Edouard de Castro, Penny Coggill, Matthew Corbett, Ujjwal Das, Louise Daugherty, Lauranne Duquenne, Robert D. Finn, Matthew Fraser, Julian Gough, Daniel Haft, Nicolas Hulo, Daniel Kahn, Elizabeth Kelly, Ivica Letunic, David Lonsdale, Rodrigo Lopez, Martin Madera, John Maslen, Craig McAnulla, Jennifer McDowall, Conor McMenamin, Huaiyu Mi, Prudence Mutowo-Muellenet, Nicola Mulder, Darren Natale, Christine Orengo, Sebastien Pesseat, Marco Punta, Antony F. Quinn, Catherine Rivoire, Amaia Sangrador-Vegas, Jeremy D. Selengut, Christian J. A. Sigrist, Maxim Scheremetjew, John Tate, Manjulapramila Thimmajanarthanan, Paul D. Thomas, Cathy H. Wu, Corin Yeats and Siew-Yit Yong

Nucleic Acids Res. 2011; **40**, D306–D312. doi: 10.1093/nar/gkr948

Gene3D: a domain-based resource for comparative genomics, functional annotation and protein network analysis

Jonathan Lees, Corin Yeats, James Perkins, Ian Sillitoe, Robert Rentzsch, Benoit H. Dessailly and Christine Orengo

Nucleic Acids Res. 2011; **40**, D465–D471. doi: 10.1093/nar/gkr1181

The authors apologize for not acknowledging the following funding bodies in these articles.

Federal funds from the National Institute of Allergy and Infectious Diseases, National Institutes of Health, Department of Health and Human Services (HSN272200700058C to C.Y.).

This correction does not influence the validity of the results and the conclusions of these articles.