ERRATUM

## Erratum to: Computational structural analysis of proteins of *Mycobacterium tuberculosis* and a resource for identifying off-targets

Sameer Hassan • Abhimita Debnath • Vasantha Mahalingam • Luke Elizabeth Hanna

Published online: 5 July 2012 © Springer-Verlag 2012

## Erratum to: J Mol Model DOI 10.1007/s00894-012-1412-5

The original version of this article unfortunately contained a mistake. On page 3, line 15 - the sentence "The  $R^2$  statistic can range from -1 to 1, with 1 representing perfect positive correlation and -1 representing perfect negative correlation" should be corrected to "The values of R squared range from 0 to 1 and the R correlation can range from -1 to 1, with 1 representing perfect positive correlation and -1 representing perfect negative correlation generated to "The values of R squared range from 0 to 1 and the R correlation can range from -1 to 1, with 1 representing perfect positive correlation and -1 representing perfect negative correlation".

The online version of the original article can be found at http://dx.doi.org/ 10.1007/s00894-012-1412-5.

S. Hassan · A. Debnath · L. E. Hanna (⊠) Department of Biomedical Informatics, National Institute for Research in Tuberculosis, Chetpet, Chennai, India e-mail: hanna@trcchennai.in

V. Mahalingam Department of Statistics, National Institute for Research in Tuberculosis, Chetpet, Chennai, India

L. E. Hanna Department of Clinical Research, National Institute for Research in Tuberculosis, Chetpet, Chennai 600031, India