

Open Access: Full open access to this and thousands of other papers at http://www.la-press.com.

Evolutionary Bioinformatics

Correction to: "In Silico Detection of Virulence Gene Homologues in the Human Pathogen *Sphingomonas* Spp."

Amr TM Saeb¹, Satish Kumar David² and Hissa Al-Brahim¹

¹Biotechnology Department, Strategic Center for Diabetes Research, King Saud University, Riyadh, Saudi Arabia. ²Information Technology Department, Strategic Center for Diabetes Research, King Saud University, Riyadh, Saudi Arabia.

CITATION: Saeb et al. Correction to: "In Silico Detection of Virulence Gene Homologues in the Human Pathogen Sphingomonas Spp.". Evolutionary Bioinformatics 2015:11 1 doi: 10.4137/EBO.S23536.

TYPE: Corrigendum

COPYRIGHT: © the authors, publisher and licensee Libertas Academica Limited. This is an open-access article distributed under the terms of the Creative Commons CC-BY-NC 3.0 License.

CORRESPONDENCE: dr.amrsaeb@gmail.com

Corrigendum

The first author of Saeb A, David SK and Al-Brahim H. In silico Detection of Virulence Gene Homologues in the Human Pathogen *Sphingomonas* Spp. *Evolutionary Bioinformatics*. 2014:10 229–238 doi: 10.4137/EBO.S20710. has brought to the attention of the editor an error in the funding section of

the paper. The authors wish to disclose funding from College of Medicine Research Center, the Deanship of Scientific Research, College of Medicine at King Saud University, and extend their appreciation for that funding. The authors confirm that the funder had no influence over the study design, content of the article, or selection of this journal.