Correction

**Feature selection environment for genomic applications**

Fabrício Martins Lopes*1,2, David Corrêa Martins Jr*1,2 and Roberto M Cesar Jr

Address: 1Instituto de Matemática e Estatística, Universidade de São Paulo, Rua do Matão 1010, 05508-090, São Paulo-SP, Brazil and 2COINF, Universidade Tecnológica Federal do Paraná, Av. Alberto Carazzai, 1640, 86300-000, Cornélio Procópio-PR, Brazil

Email: Fabrício Martins Lopes* - fabricio@utfpr.edu.br; David Corrêa Martins* - davidjr@vision.ime.usp.br; Roberto M Cesar - cesar@vision.ime.usp.br

* Corresponding authors

Published: 10 September 2009
Received: 25 August 2009
Accepted: 10 September 2009

BMC Bioinformatics 2009, 10:285 doi:10.1186/1471-2105-10-285

This article is available from: http://www.biomedcentral.com/1471-2105/10/285

© 2009 Lopes et al; licensee BioMed Central Ltd.
This is an Open Access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/2.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

**Correction**

We have found an error after the publication of this work [1]. The error is present in the following sentence: "We have that $q = 1$ is the Shannon's Entropy (extensive entropy), $q < 1$ is the sub-extensive entropy and $q > 1$ is the super-extensive entropy".

Where it says "sub-extensive", should be changed to "super-extensive" and where it says "super-extensive", should be changed to "sub-extensive".

The corrected sentence: We have that $q = 1$ is the Shannon's Entropy (extensive entropy), $q < 1$ is the super-extensive entropy and $q > 1$ is the sub-extensive entropy.

We regret any inconvenience that this inaccuracy in the text might have caused.

**References**

1. Lopes FM, Martins DC Jr, Cesar RM Jr: Feature selection environment for genomic applications. BMC Bioinformatics 2008, 9(451):1-21 [http://www.biomedcentral.com/1471-2105/9/451/].

