



Corrigendum: Dysregulation of MicroRNAs and Target Genes Networks in Peripheral Blood of Patients With Sporadic Amyotrophic Lateral Sclerosis

Maria Liguori^{1*}, Nicoletta Nuzziello^{1†}, Alessandro Introna^{2†}, Arianna Consiglio¹, Flavio Licciulli¹, Eustachio D'Errico², Antonio Scarafino², Eugenio Distaso² and Isabella L. Simone²

¹ National Research Council, Institute of Biomedical Technologies, Bari Unit, Bari, Italy, ² Department of Basic Sciences, Neurosciences and Sense Organs, University of Bari, Bari, Italy

Keywords: sporadic amyotrophic lateral sclerosis, microRNA, target genes, peripheral blood markers, high throughput next-generation sequencing (HT-NGS), clinical parameters, bioinformatics, pathway analysis

OPEN ACCESS A Corrigendum on

Edited and reviewed by:

Giuseppe Filomeni, Danish Cancer Society, Denmark

> *Correspondence: Maria Liguori maria.liguori@cnr.it; maria.liguori@ba.itb.cnr.it

[†]These authors have contributed equally to this work

Received: 30 November 2018 Accepted: 14 December 2018 Published: 10 January 2019

Citation:

Liguori M, Nuzziello N, Introna A, Consiglio A, Licciulli F, D'Errico E, Scarafino A, Distaso E and Simone IL (2019) Corrigendum: Dysregulation of MicroRNAs and Target Genes Networks in Peripheral Blood of Patients With Sporadic Amyotrophic Lateral Sclerosis. Front. Mol. Neurosci. 11:488. doi: 10.3389/fnmol.2018.00488

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by Liguori, M., Nuzziello, N., Introna, A., Consiglio, A., Licciulli, F., D'Errico, E., et al. (2018). Front. Mol. Neurosci. 11:288. doi: 10.3389/fnmol.2018.00288

In the original article, there was a mistake in Figure 1 as published. We used the Volcano Plot, which was automatically generated using the Expression Suite v1.0.3 software (Thermo Fisher Scientific). This software visualizes the *p*-values and fold change values after each qRT-PCR run. A log2 (Fold Change) greater than or equal to one, is the standard boundary field.

Therefore, targets with a fold less than the lower boundary were automatically colored in green. However, as stated in the article, according to published guidelines, we validated the miRNAs with p-values < 0.05 (dot over the blue line) independently from their fold change. We have therefore adjusted the representation in the figure to avoid confusion.

The corrected **Figure 1** appears below. The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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FIGURE 1 Volcano plot of validated miRNAs. Green dots represent the 38 differentially expressed miRNAs obtained from the comparison between sALS and HC subjects by qRT-PCR (p < 0.05). All black dots below the blue line did not discriminate sALS from HC. The Y-axis represents the log10 of the p-value and the X-axis represents log fold change of miRNA expression in the sALS versus HC.