



Corrigendum: Age-Related Differences in the Expression of Most Relevant Mediators of SARS-CoV-2 Infection in Human Respiratory and Gastrointestinal Tract

OPEN ACCESS

Approved by:

Frontiers Editorial Office, Frontiers Media SA, Switzerland

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Specialty section:

This article was submitted to Pediatric Gastroenterology, Hepatology and Nutrition, a section of the journal Frontiers in Pediatrics

Received: 06 October 2021 Accepted: 15 October 2021 Published: 10 November 2021

Citation:

Berni Canani R, Comegna M, Paparo L, Cernera G, Bruno C, Strisciuglio C, Zollo I, Gravina AG, Miele E, Cantone E, Gennarelli N, Nocerino R, Carucci L, Giglio V, Amato F and Castaldo G (2021) Corrigendum: Age-Related Differences in the Expression of Most Relevant Mediators of SARS-CoV-2 Infection in Human Respiratory and Gastrointestinal Tract. Front. Pediatr. 9:790285. doi: 10.3389/fped.2021.790285 Roberto Berni Canani^{1,2,3,4*}, Marika Comegna^{2,5}, Lorella Paparo^{1,2}, Gustavo Cernera^{2,5}, Cristina Bruno^{1,2}, Caterina Strisciuglio⁶, Immacolata Zollo^{2,5}, Antonietta Gerarda Gravina⁷, Erasmo Miele¹, Elena Cantone⁸, Nicola Gennarelli⁹, Rita Nocerino^{1,2}, Laura Carucci^{1,2}, Veronica Giglio^{1,2}, Felice Amato^{2,5} and Giuseppe Castaldo^{2,5}

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Keywords: COVID-19, angiotensin-converting enzyme 2, transmembrane serine protease-2, neuropilin-1, healthy subjects

A Corrigendum on

Age-Related Differences in the Expression of Most Relevant Mediators of SARS-CoV-2 Infection in Human Respiratory and Gastrointestinal Tract

by Berni Canani, R., Comegna, M., Paparo, L., Cernera, G., Bruno, C., Strisciuglio, C., Zollo, I., Gravina, A. G., Miele, E., Cantone, E., Gennarelli, N., Nocerino, R., Carucci, L., Giglio, V., Amato, F., and Castaldo, G. (2021). Front. Pediatr. 9:697390. doi: 10.3389/fped.2021.697390

In the original article, there was a mistake in the order of **Figures 1** and **2** as published. The figures are given in the correct order below.

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

1



FIGURE 1 | qPCR analysis of Angiotensin I Converting Enzyme (*ACE1*), Angiotensin II Converting Enzyme (*ACE2*), Transmembrane Serine Protease 2 (*TMPRSS2*), and Neuropilin-1 (*NRP1*) genes in nasal epithelium from children and adult subjects. Comparative expression of *ACE1*, *ACE2*, *TMPRSS2*, and *NRP1* in nasal epithelium of children (n = 15) and adult subjects (n = 15). Data analysis was performed using the comparative threshold cycle (CT) method and expressed as 2[^]-delta CT. Gene expression was normalized against the expression of the reference gene hypoxanthine phosphoribosyltransferase 1 (*HPRT*). Data are expressed as median \pm SD, the X in the bars indicates mean values. *P*-value are reported in the graphs; significant differences are indicated as *p < 0.01; **p < 0.005.



FIGURE 2 | qPCR analysis of Angiotensin I Converting Enzyme (*ACE1*), Angiotensin II Converting Enzyme (*ACE2*), Transmembrane Serine Protease 2 (*TMPRSS2*), and Neuropilin-1 (*NRP1*) genes in small intestine from children and adult subjects. Comparative expression of *ACE1*, *ACE2*, *TMPRSS2*, and *NRP1* in small intestine of children (n = 15) and adult subjects (n = 15). Data analysis was performed using the comparative threshold cycle (CT) method and expressed as 2[^]-delta CT. Gene expression was normalized against the expression of the reference gene hypoxanthine phosphoribosyltransferase 1 (*HPRT*). Data are expressed as median \pm SD, the X in the bars indicates mean values. *P*-value are reported in the graphs; significant differences are indicated as *p < 0.01; **p < 0.005.

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