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EDITED AND REVIEWED BY Wolfgang Holnthoner, AUVA Research Centre, Austria

*CORRESPONDENCE Juan C. Briceño, ⊠ jbriceno@uniandes.edu.co

RECEIVED 07 April 2025 ACCEPTED 09 April 2025 PUBLISHED 24 April 2025

CITATION

Rodríguez-Soto MA, Riveros-Cortés A, Orjuela-Garzón IC, Fernández-Calderón IM, Rodríguez CF, Suárez Vargas N, Ostos C, Muñoz Camargo C, Cruz JC, Kim S, D'Amore A, Wagner WR and Briceño JC (2025) Corrigendum: Redefining vascular repair: revealing cellular responses on PEUU—gelatin electrospun vascular grafts for endothelialization and immune responses on *in vitro* models. *Front. Bioeng. Biotechnol.* 13:1607125.

doi: 10.3389/fbioe.2025.1607125

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Corrigendum: Redefining vascular repair: revealing cellular responses on PEUU—gelatin electrospun vascular grafts for endothelialization and immune responses on *in vitro* models

María A. Rodríguez-Soto¹, Alejandra Riveros-Cortés¹, Ian C. Orjuela-Garzón¹, Inés María Fernández-Calderón¹, Cristian F. Rodríguez¹, Natalia Suárez Vargas¹, Carlos Ostos², Carolina Muñoz Camargo¹, Juan C. Cruz¹, Seungil Kim³, Antonio D'Amore³, William R. Wagner³ and Juan C. Briceño^{1.4*}

¹Department of Biomedical Engineering, Universidad de los Andes, Bogotá, Colombia, ²Instituto de Química, Facultad de Ciencias Exactas y Naturales, Universidad de Antioquia, Medellín, Colombia, ³McGowan Institute for Regenerative Medicine and Department of Bioengineering, University of Pittsburgh, Pittsburgh, PA, United States, ⁴Department of Congenital Heart Disease and Cardiovascular Surgery, Fundación CardioInfantil Instituto de Cardiología, Bogotá, Colombia

KEYWORDS

tissue engineered vascular grafts, regenerative medicine, biomaterials, inflammatory response, immunomodulation, M1/M2 macrophage polarization, endothelialization, cell signaling

A Corrigendum on

Redefining vascular repair: revealing cellular responses on PEUU—gelatin electrospun vascular grafts for endothelialization and immune responses on in vitro models

by Rodríguez-Soto MA, Riveros-Cortés A, Orjuela-Garzón IC, Fernández- Calderón IM, Rodríguez CF, Vargas NS, Ostos C, Camargo CM, Cruz JC, Kim S, D'Amore A, Wagner WR and Briceño JC (2024). Front. Bioeng. Biotechnol. 12:1410863. doi: 10.3389/fbioe.2024.1410863

In the published article, there was an error in Figure 9 as published. The image labeled as "2D control Day 1 panel of Phalloidin (AF 488)" was mistakenly similar to the image labeled as "ML + P + P Lumen Day 7 panel of Phalloidin (AF 488) due to a mislabeling during file organization." The corrected Figure 9 and its caption "Endothelialization potential of ML + P + P with HUVECs seeded on the luminal surface. (A) Phalloidin staining at days 1 and 7 compared with a 2D control on a glass slide. (B) SEM images of Endothelial cell lining. Black arrows highlight cells and cell nuclei, yellow arrows indicate cell boundaries, and red arrows correspond to cracks in the fixed cell monolayer resulting from sample processing; beneath this layer, electrospun fibers are visible. (C) Percentage of covered surface area by HUVECs,



FIGURE 9

Endothelialization potential of ML + P + P with HUVECs seeded on the luminal surface. (A) Phalloidin staining at days 1 and 7 compared with a 2D control on a glass slide. (B) SEM images of Endothelial cell lining. Black arrows highlight cells and cell nuclei, yellow arrows indicate cell boundaries, and red arrows correspond to cracks in the fixed cell monolayer resulting from sample processing; beneath this layer, electrospun fibers are visible. (C) Percentage of covered surface area by HUVECs, data normalized from with 2D control. (D) DNA quantification at 7 days. (E) RNA expression profile. (E) VEGF and NO release. (F) Intracellular ROS production. (Mean \pm SD) where, ns = no significant *p \leq 0.001, ***p \leq 0.001, ****p \leq 0.001, ****p \leq 0.001.

data normalized from with 2D control. (D) DNA quantification at 7 days. (E) RNA expression profile. (E) VEGF and NO release. (F) Intracellular ROS production. (Mean \pm SD) where, ns = no significant *p \leq 0.05, **p \leq 0.01, ***p \leq 0.001, ****p \leq 0.0001." appear 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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