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Erratum: Large-Scale Gene Expression Signatures Reveal a Microbicidal Pattern of Activation in *Mycobacterium leprae*-Infected Monocyte-Derived Macrophages With Low Multiplicity of Infection

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An erratum on

Large-Scale Gene Expression Signatures Reveal a Microbicidal Pattern of Activation in *Mycobacterium leprae*-Infected Monocyte-Derived Macrophages With Low Multiplicity of Infection by Leal-Calvo T, Martins BL, Bertoluci DF, Rosa PS, Camargo RM, Germano GV, Brito de Souza VN, Pereira Latin AC and Moraes MO (2021). *Front. Immunol.* 12:647832. doi: 10.3389/fimmu.2021.647832

Due to a production error, there was a mistake in **Figure 2** as published. The filled circles in the dot plot did not export correctly. The corrected **Figure 2** appears below. The publisher apologizes for this mistake.

The original version of this article has been updated.

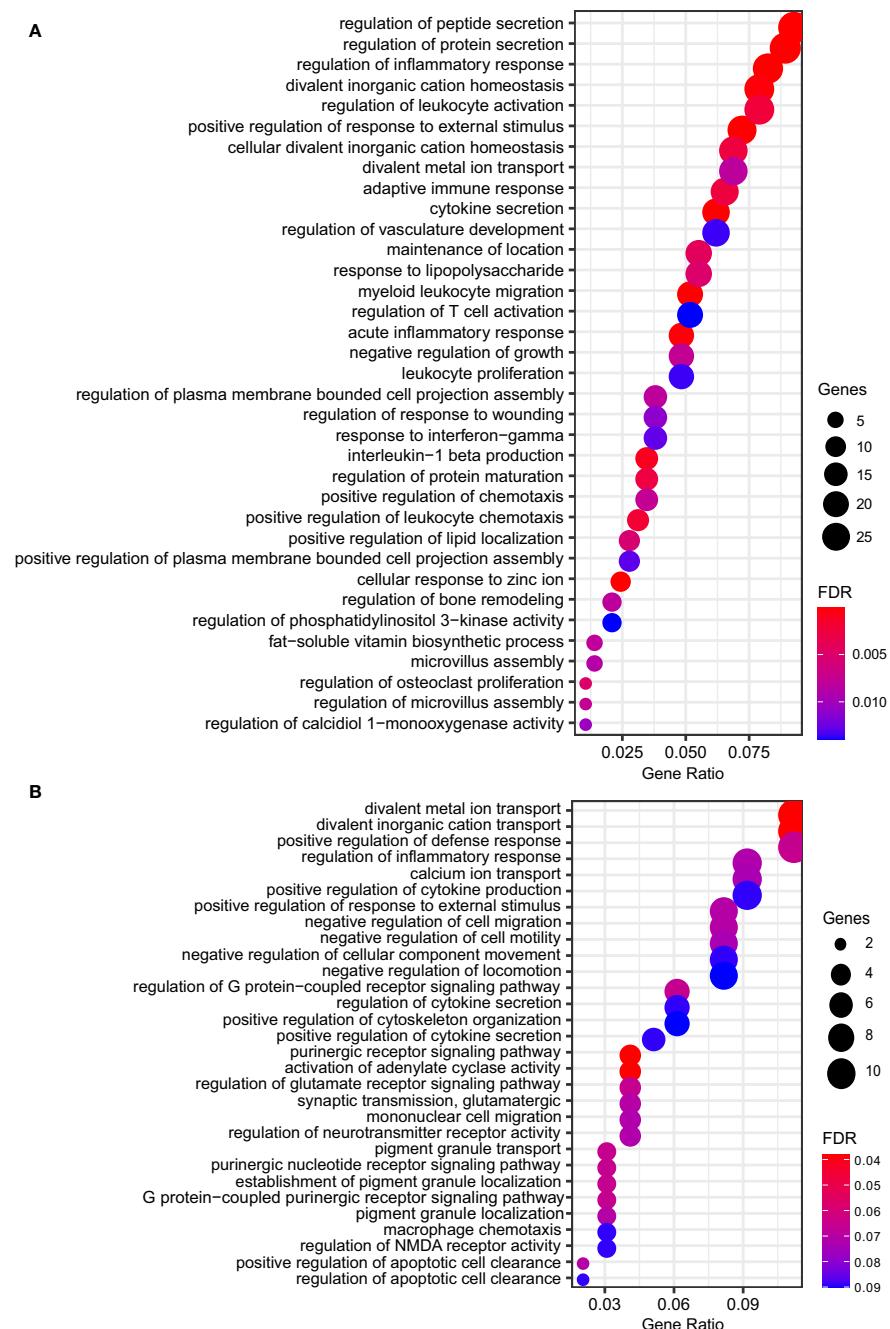


FIGURE 2 | Dot plot showing the top significant GO biological processes enriched from ORA of genes (A) upregulated ($n = 35$) or (B) repressed ($n = 30$) by *M. leprae* infection. Gene ratio is the fraction of genes belonging to an ontology over the total number of modulated genes. Circle size shows the number of modulated genes per biological process. FDR, false discovery rate.

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