OPEN **Author Correction:** Targeted metabolomics identifies high performing diagnostic and prognostic biomarkers for COVID-19

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The original version of this Article contained an error in Figure 1b, where the labels of ‘Group 1’ and ‘Group 2’ were mistakenly interchanged. The label for ‘Group 1’ now reads ‘Group 2’ and the label for ‘Group 2’ now reads ‘Group 1’. The error does not alter the results and conclusions.

The original Figure 1 and accompanying legend appears below.

The original Article has been corrected.

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Figure 1. Multivariate analysis from plasma metabolome profile of G1 versus G2 patients. **(a)** Score scatter plot based on the PLS-DA models to explain the diagnosis (green for G1 and yellow for G2); **(b)** rank of the different metabolites (the top 15) identified by the PLS-DA according to the VIP coefficient on the x-axis. The most discriminating metabolites are shown in descending order of their coefficient scores. The color boxes indicate whether metabolite concentration is increased (red) or decreased (blue) in G1 vs G2; **(c)** ROC curve of the demographic/clinical data model; **(d)** ROC curve of the metabolite-only model; **(e)** ROC curve of the metabolite + demographic/clinical data model. The figures were drawn via MetaboAnalyst software v 4.0 (<https://www.metaboanalyst.ca/>).

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