Corrigendum

Corrigendum to “Salvia miltiorrhiza and the Volatile of Dalbergia odorifera Attenuate Chronic Myocardial Ischemia Injury in a Pig Model: A Metabonomic Approach for the Mechanism Study”

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In the article titled “Salvia miltiorrhiza and the Volatile of Dalbergia odorifera Attenuate Chronic Myocardial Ischemia Injury in a Pig Model: A Metabonomic Approach for the Mechanism Study” [1], the authors identified an error in Figure 4, where (b) and (f) were mistakenly duplicated. Figure 4(f) had been corrected as follows:

Additionally, the P-AMPK bands in Figure 8(a) and the GLUT-4 bands in Figure 8(b) were found to be duplicated. Figure 8(b) has been corrected as follows:
Figure 4: Continued.
Figure 4: The metabolic profiles of OPLS-DA between the model and SM-DOO groups based on HPLC-Q-TOF-MS: (a) the score plot of OPLS-DA in a negative mode; (b) the corresponding validation plot based on 200 times permutation tests demonstrated the robustness of the OPLS-DA model in a negative mode; (c) the S-plot of OPLS-DA in a negative mode; (d) the VIP of OPLS-DA in a negative mode; (e) the score plot of OPLS-DA in a positive mode; (f) the corresponding validation plot based on 200 times permutation tests demonstrated the robustness of the OPLS-DA model in a positive mode; (g) the S-plot of OPLS-DA in a positive mode; (h) the VIP of OPLS-DA in a positive mode.

Figure 8: Effects of SM-DOO on the AMPK/GLUT4 pathway in heart tissues of pigs. p-AMPK, AMPK, GLUT4, and β-actin protein expressions were measured. The protein signals were quantitated by densitometry, and the graph shows their relative levels. All values are presented as the mean ± SD. ##P < 0.01 vs. sham group; **P < 0.01 vs. model group.
References