

Correction



Correction: Zhang, Y., *et al.* Clones of *FeSOD*, *MDHAR*, *DHAR* Genes from White Clover and Gene Expression Analysis of ROS-Scavenging Enzymes during Abiotic Stress and Hormone Treatments. *Molecules* 2015, 20, 20939–20954

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Received: 27 November 2015 ; Accepted: 30 December 2015 ; Published: 11 December 2015 Academic Editor: Derek J. McPhee

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The authors wish to make the following correction to this paper [1]. Due to mislabeling, the following figures:



Figure 4. Quantitative real-time RT-PCR was used to analyze of ROS-scavenging enzyme genes expression during drought stress and normalized to β -*actin*. White clover leaves were sampled after 0, 3, 6, 12, 24, 48 and 72 h treatment. Data represent means of three replicates. Error bars representing standard errors and the different letters above the bars represent significant difference (*p* < 0.05).



Figure 6. Quantitative real-time RT-PCR was used to analysis of ROS-scavenging enzyme genes expression during heavy metal stress and normalized to β -*actin*. White clover leaves were sampled after 0, 3, 6, 12, 24, 48 and 72 h treatment. Data represent means of three replicates. Error bars representing standard errors and the different letters above the bars represent significant difference (p < 0.05).



Figure 7. Quantitative real-time RT-PCR was used to analyze of ROS-scavenging enzyme genes expression treatment with ABA and normalized to β -*actin*. White clover leaves were sampled after 0, 3, 6, 12, 24, 48 and 72 h treatment. Data represent means of three replicates. Error bars representing standard errors and the different letters above the bars represent significant difference (*p* < 0.05).



Figure 8. Quantitative real-time RT-PCR was used to analyze of ROS-scavenging enzyme genes expression treatment with Spd and normalized to β -*actin*. White clover leaves were sampled after 0, 3, 6, 12, 24, 48 and 72 h treatment. Data represent means of three replicates. Error bars representing standard errors and the different letters above the bars represent significant difference (*p* < 0.05).



should be replaced by:

Figure 4. Quantitative real-time RT-PCR was used to analyze of ROS-scavenging enzyme genes expression during drought stress and normalized to β -*actin*. White clover leaves were sampled after 0, 3, 6, 12, 24, 48 and 72 h treatment. Data represent means of three replicates. Error bars representing standard errors and the different letters above the bars represent significant difference (*p* < 0.05).



Figure 6. Quantitative real-time RT-PCR was used to analysis of ROS-scavenging enzyme genes expression during heavy metal stress and normalized to β -*actin*. White clover leaves were sampled after 0, 3, 6, 12, 24, 48 and 72 h treatment. Data represent means of three replicates. Error bars representing standard errors and the different letters above the bars represent significant difference (p < 0.05).



Figure 7. Quantitative real-time RT-PCR was used to analyze of ROS-scavenging enzyme genes expression treatment with ABA and normalized to β -*actin*. White clover leaves were sampled after 0, 3, 6, 12, 24, 48 and 72 h treatment. Data represent means of three replicates. Error bars representing standard errors and the different letters above the bars represent significant difference (*p* < 0.05).



Figure 8. Quantitative real-time RT-PCR was used to analyze of ROS-scavenging enzyme genes expression treatment with Spd and normalized to β -*actin*. White clover leaves were sampled after 0, 3, 6, 12, 24, 48 and 72 h treatment. Data represent means of three replicates. Error bars representing standard errors and the different letters above the bars represent significant difference (p < 0.05).

Some of the letters which represented significance were not correct, and accordingly the changes in the figures are as follows: In Figure 4 *FeSOD*, the letters at 6 and 12 h should be changed to b and a, in *Cu/Zn SOD*, the letters at 3 and 6 h should be changed to ab and a. In Figure 6 *FeSOD*, the letters of all treatments except for 6 h should be changed to a. In Figure 7 *Cu/Zn SOD*, the letters of all treatments should be changed to d, c, bc, ab, a, ab and cd, respectively, in *MnSOD*, the letters of all treatments except for 72 h should be changed to a, in *POD*, the letters at 3 and 6 h should be changed to a and 6 h should be changed to b and a, in *APX*, the letters at 24 and 48 h should be changed to a and b, in *DHAR*, the letters at 48 and 72 h should be changed to b and d. In Figure 8 *APX*, the letters at 3 and 6 h should be changed to b and a, in *GR*, the letters at 72 h should be changed to c. The authors would like to apologize for any inconvenience caused to the readers by these changes.

Reference

Zhang, Y.; Li, Z.; Peng, Y.; Wang, X.; Peng, D.; Li, Y.; He, X.; Zhang, X.; Ma, X.; Huang, L.; *et al.* Clones of *FeSOD*, *MDHAR*, *DHAR* Genes from White Clover and Gene Expression Analysis of ROS-Scavenging Enzymes during Abiotic Stress and Hormone Treatments. *Molecules* 2015, 20, 20939–20954. [CrossRef] [PubMed]



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