

Fig. 1 Suppl. Germination index (A), shoot length (B), root length (C), and adventitious root number (D) of five different rice cultivars in response to different graphene oxide (GO) dosages for 16 d. Means \pm SDs, $n = 3$. Lower-case letters indicate significant differences (Duncan's multiple range test, $P < 0.01$) between genotypes under the control or particular GO treatment.

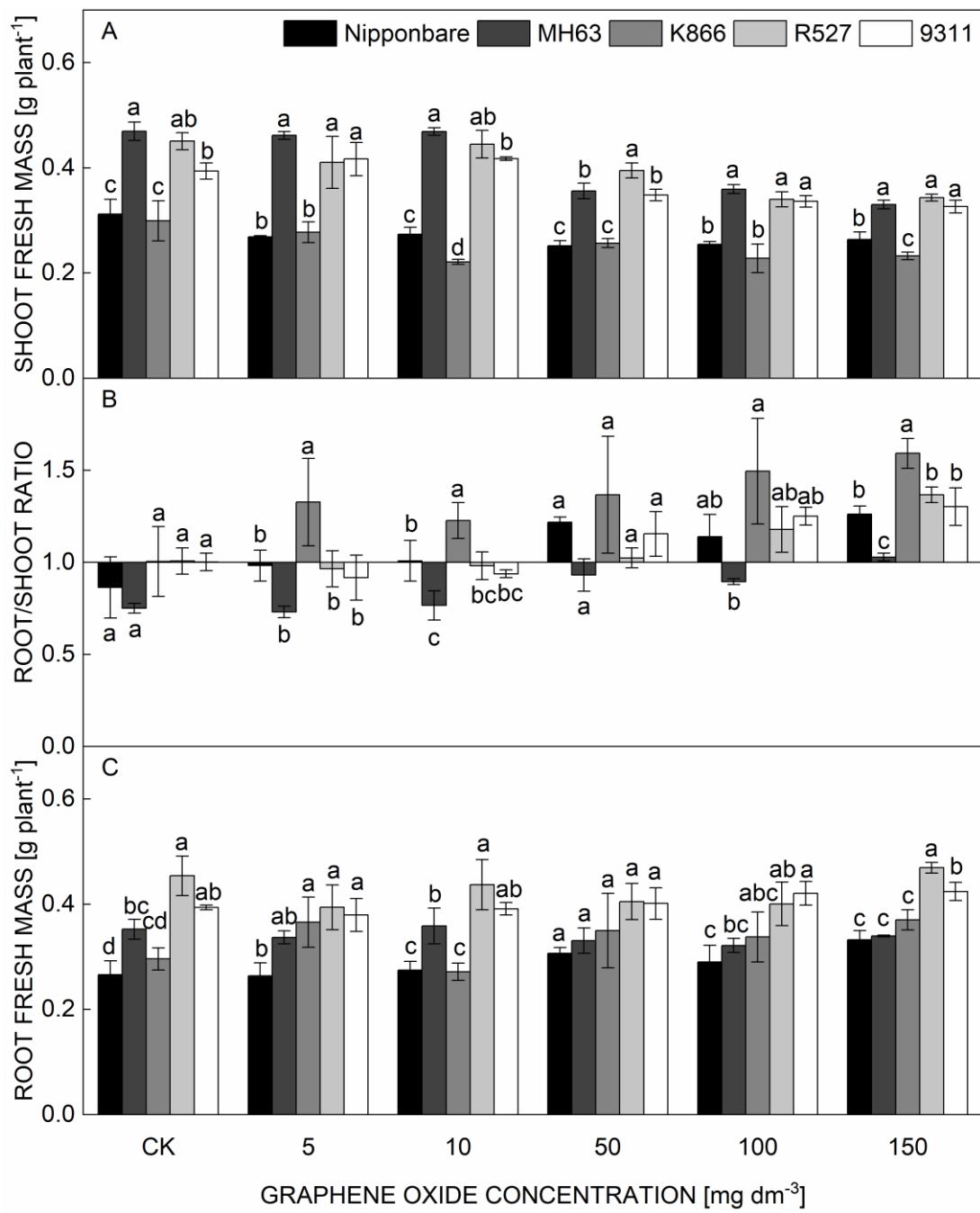


Fig. 2 Suppl. Shoot (A) and root (C) fresh masses, and root/shoot ratio (B) of five different rice cultivars in response to different graphene oxide (GO) dosages for 16 d. Means \pm SDs, $n = 3$. Lower-case letters indicate significant differences (Duncan's multiple range test, $P < 0.01$) between genotypes under the control or particular GO treatment.

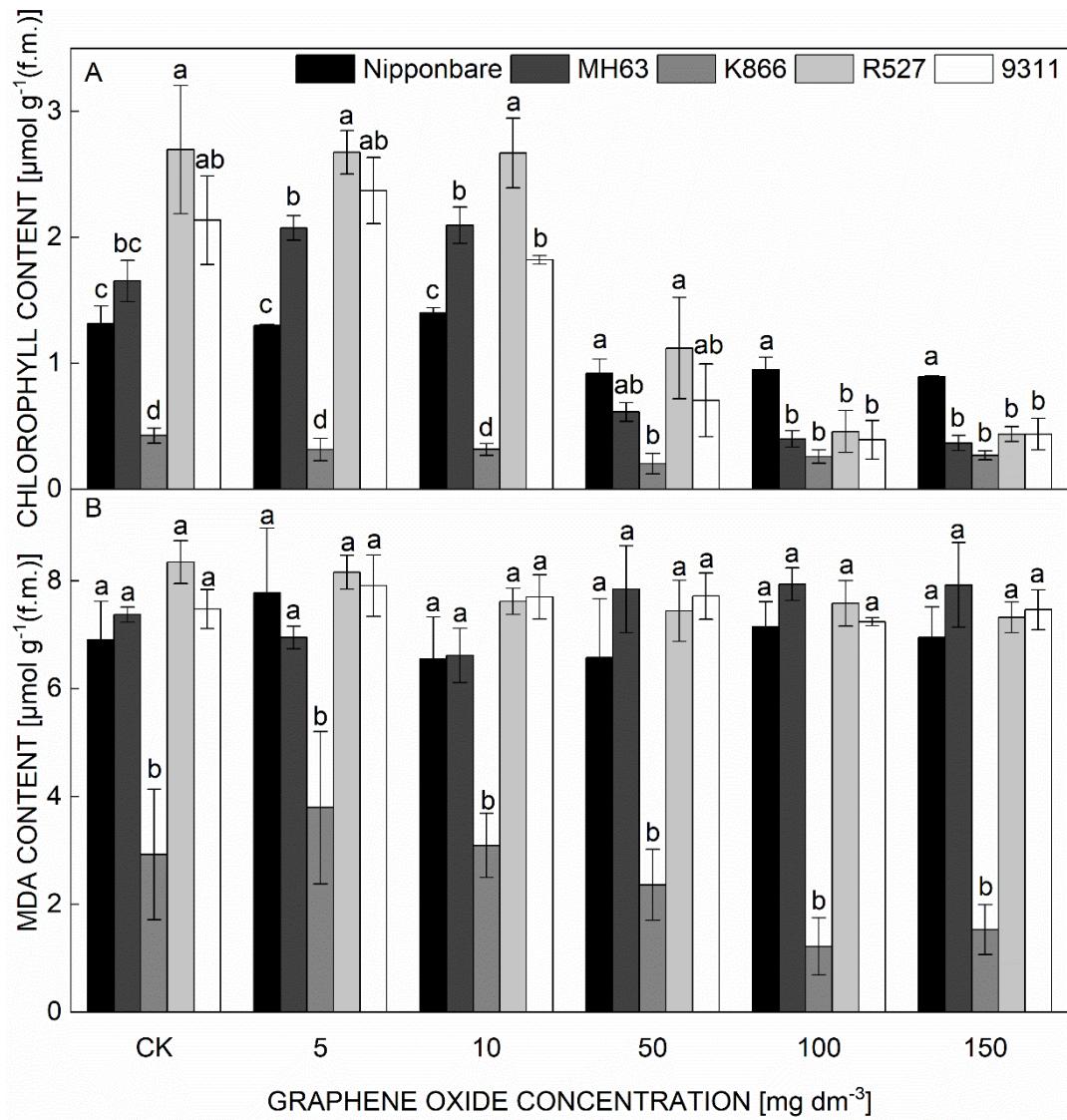


Fig. 3 Suppl. Chlorophyll (A) and malondialdehyde (MDA) content of five different rice cultivars in response to different graphene oxide (GO) dosages for 16 d. Means \pm SDs, $n = 3$. Lower-case letters indicate significant differences (Duncan's multiple range test, $P < 0.01$) between genotypes under the control or particular GO treatment.

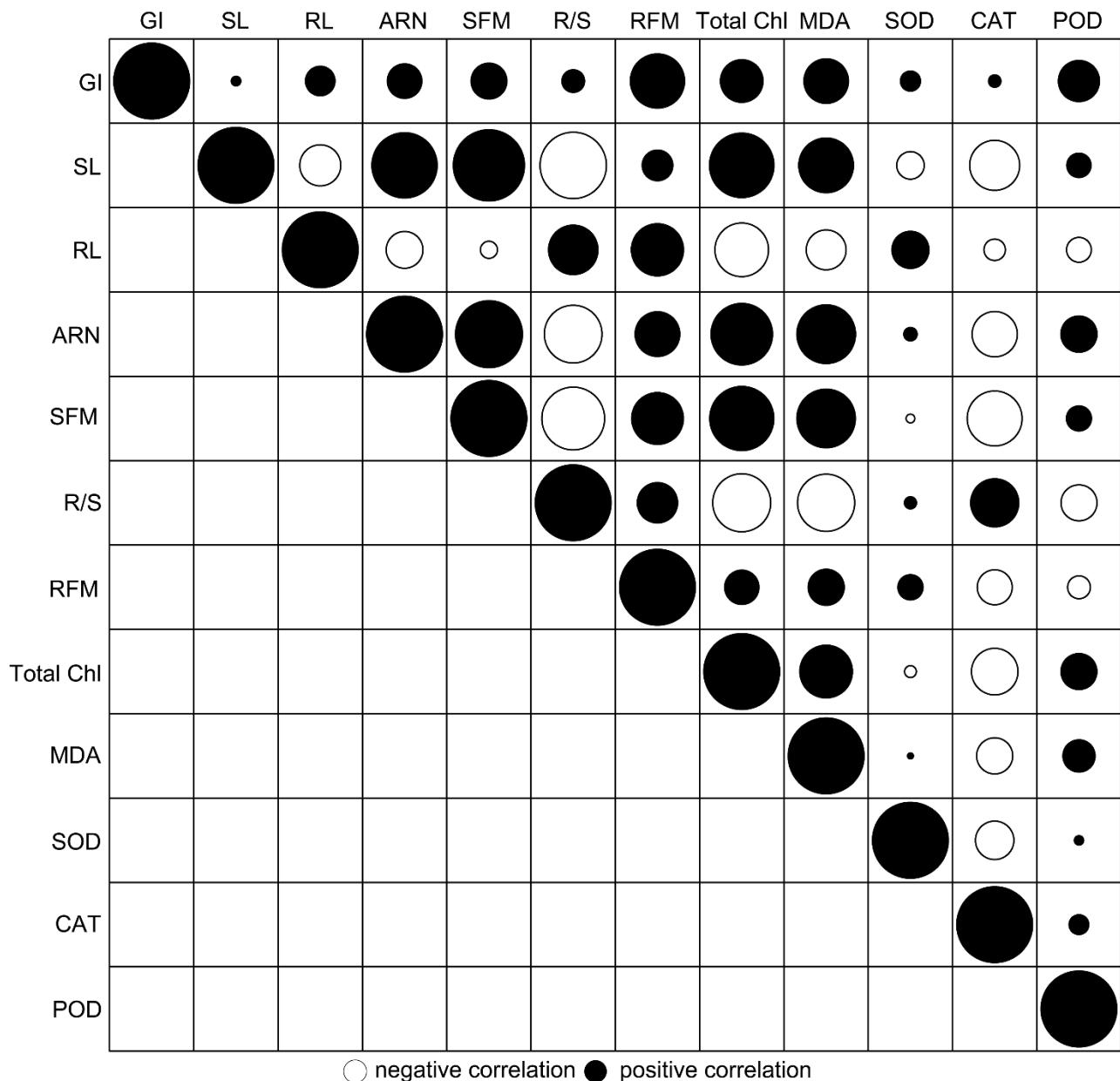


Fig. 4 Suppl. Visualization of a correlation matrix based on Pearson correlation coefficients of five different rice cultivars in response to different graphene oxide dosages for 16 d. *Empty and solid circles* indicate negative and positive correlations, respectively. The size of a circle represents the correlation intensity. GI - germination index, SL - shoot length, RL - root length, ARN - adventitious root number, SFM - shoot fresh mass, RFW - root fresh mass, R/S - root/shoot ratio, total Chl - total chlorophyll content, MDA - malondialdehyde content, SOD - superoxide dismutase activity, CAT - catalase activity, POD - peroxidase activity.