



Figure S4. Identification of DGEs in transcriptome by qRT-PCR.

The kale seedlings were treated with 10mg·L⁻¹ SeMet for 5 d, and the leaves were sampled with H₂O treatment as the CK. The red line represents the FPKM of RNA-seq, the histogram represents the relative mRNA expression determined by qRT-PCR. Data are the means of three replicates (±SDs). Different letters indicate the significant differences between samples according to the T-test (* representing P<0.05 and ** representing P<0.01).