Metabolic Signatures in Response to Abscisic Acid (ABA) Treatment in *Brassica napus* Guard Cells Revealed by Metabolomics

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Supplementary information

Supplemental Figure S1. Mapping of identified metabolites from *B. napus* guard cells into KEGG metabolic pathways. Black dots represent mapped metabolites.

Supplemental Figure S2. Principal component analysis using all detected features from the combined GC-MS/MS and LC-MS/MS data sets.

Supplemental Table 1. Metabolites of *B. napus* guard cells identified from GC-MS/MS and LC-MS/MS platforms.

Supplemental Table 2. ABA-responsive metabolites in *B. napus* guard cell revealed by non-targeted metabolomics.

Supplemental Table 3. Pathway enrichment analysis report generated from MetaboAnalyst.



Supplemental Figure S1. Mapping of identified metabolites from *B. napus* guard cells into KEGG metabolic pathways. Black dots represent mapped metabolites.



Supplemental Figure S2. Principal component analysis using all detected features from the combined GC-MS/MS and LC-MS/MS data sets.