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Supplemental information

Multi-omics analysis of early

leaf development in Arabidopsis thaliana

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Figure S1. Metabolite changes across early leaf development. (A) Number of differential metabolites calculated between two consecutive time-points (e.g., between day 8 and day 9) using *t*-Test. **(B)** Number of metabolites with the highest abundance measured at a given day (axis x). **(A, B)** Analysis was restricted to the metabolites assigned as differential using one-way ANOVA. Complete data can be found in **Supplementary Dataset S3.**



Figure S2. Pearson correlation matrix between differential proteins and metabolites (red to blue). Also given max normalized accumulation pattern across developmental series (blue to yellow). Complete data can be found in Supplementary Dataset S4.