

## Supplementary Figures

### **Metabolomic Profiling in Combination with Data Association Analysis Provide Insights about Potential Metabolic Regulation Networks among Non-Volatile and Volatile Metabolites in *Camellia sinensis* cv *Baijiguan***

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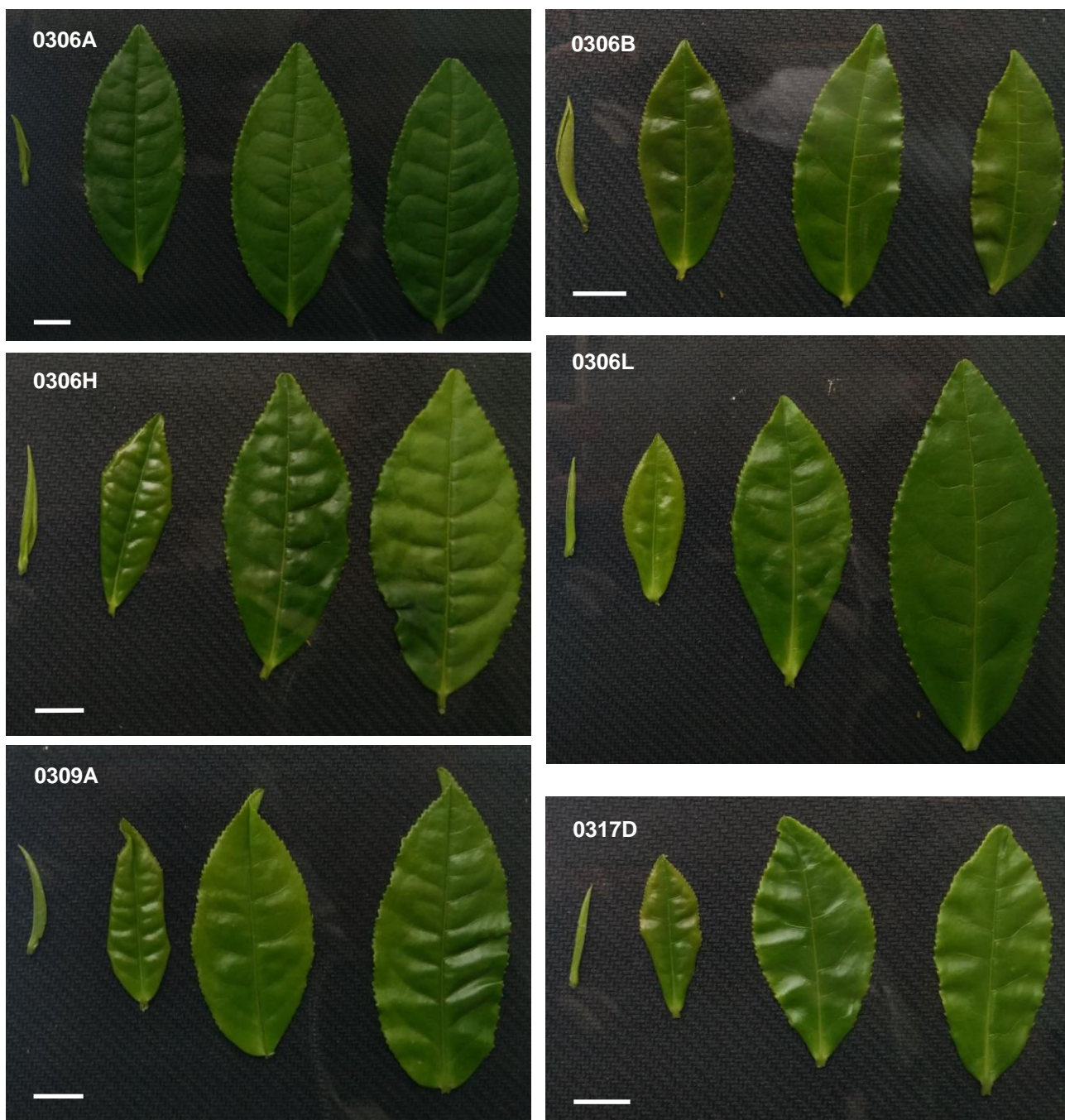
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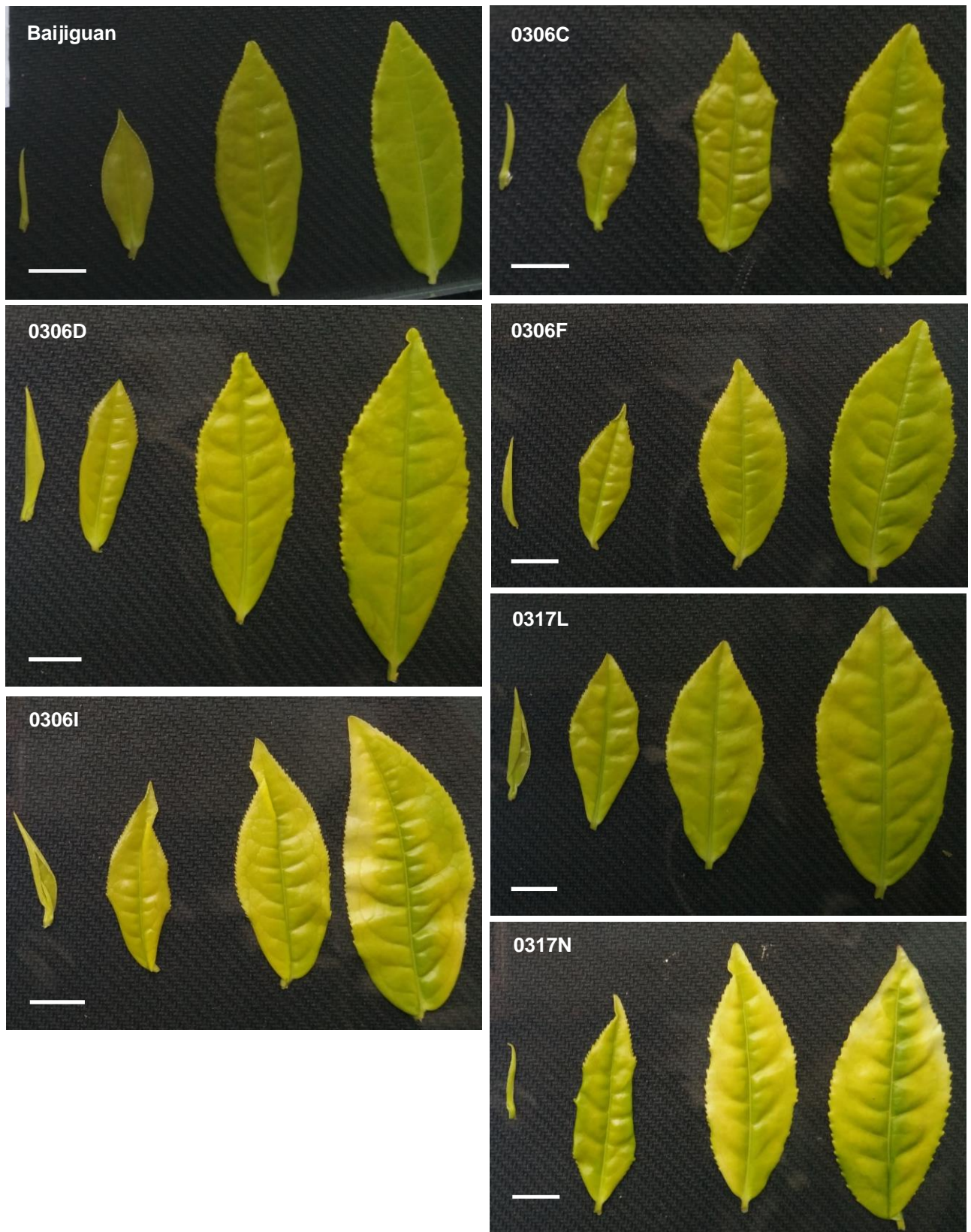
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\*Equal contribution to this work.

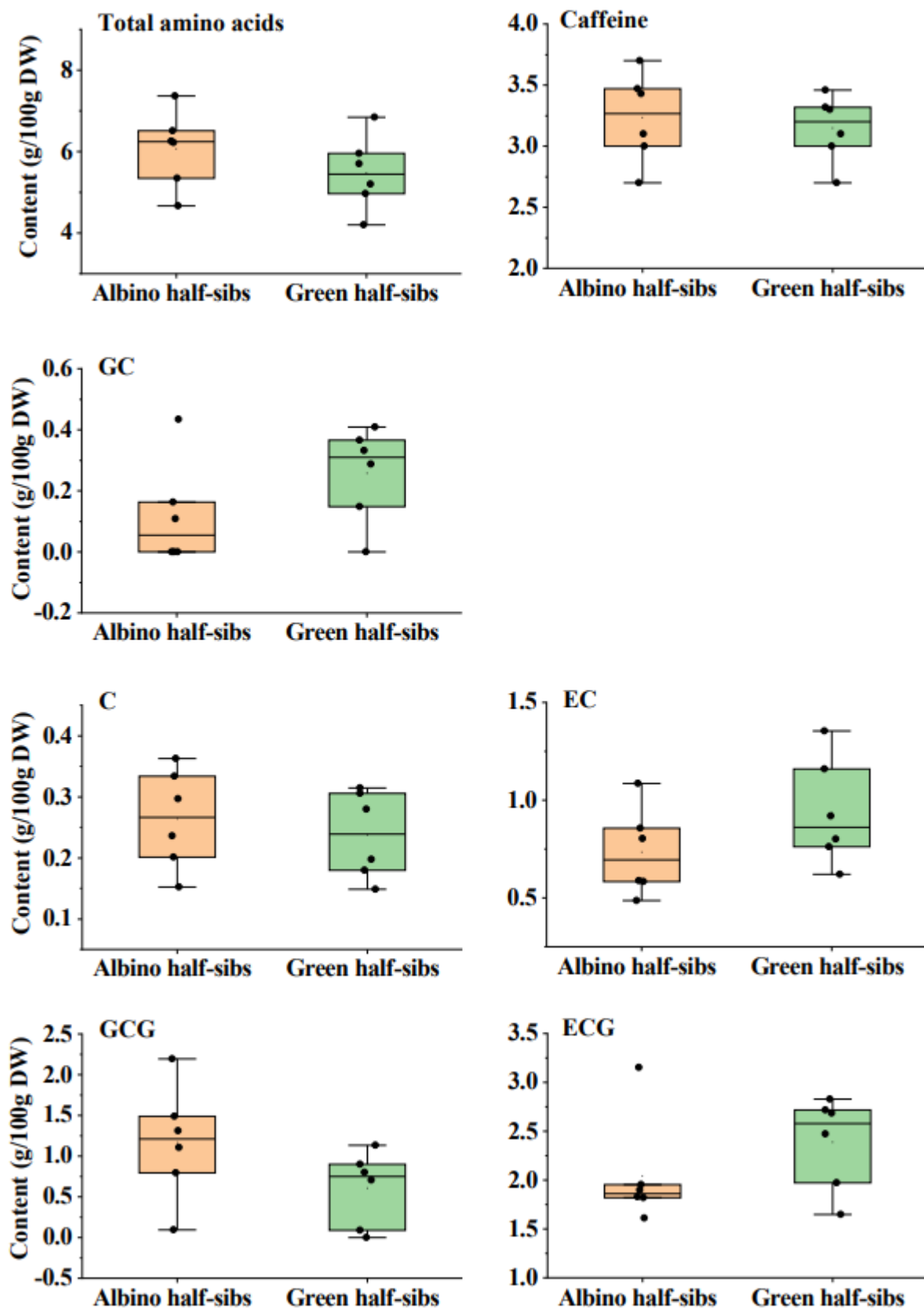


**Figure S1.** One bud and three leaves of the green leaf half-sibs descendent from *Camellia sinensis* cv *Baijiguan*. Bar = 1 cm.

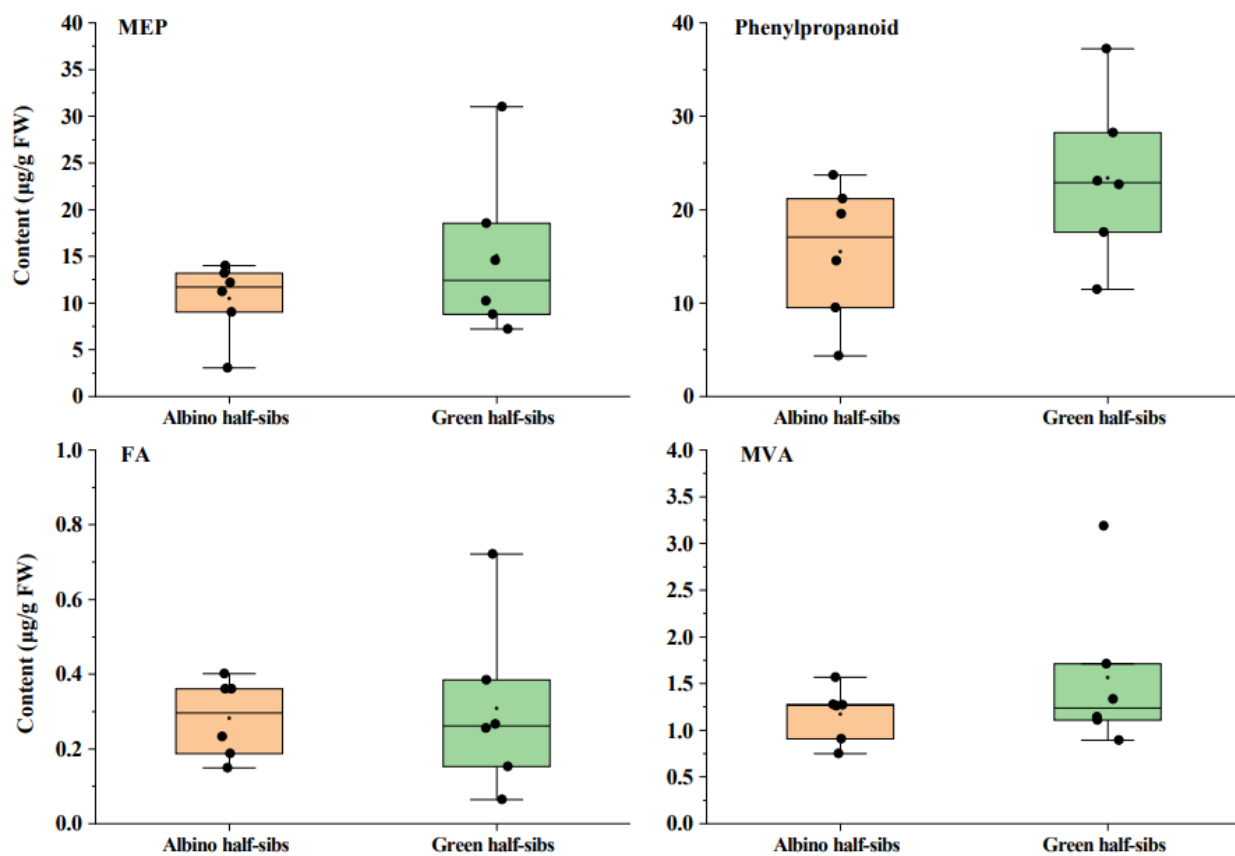




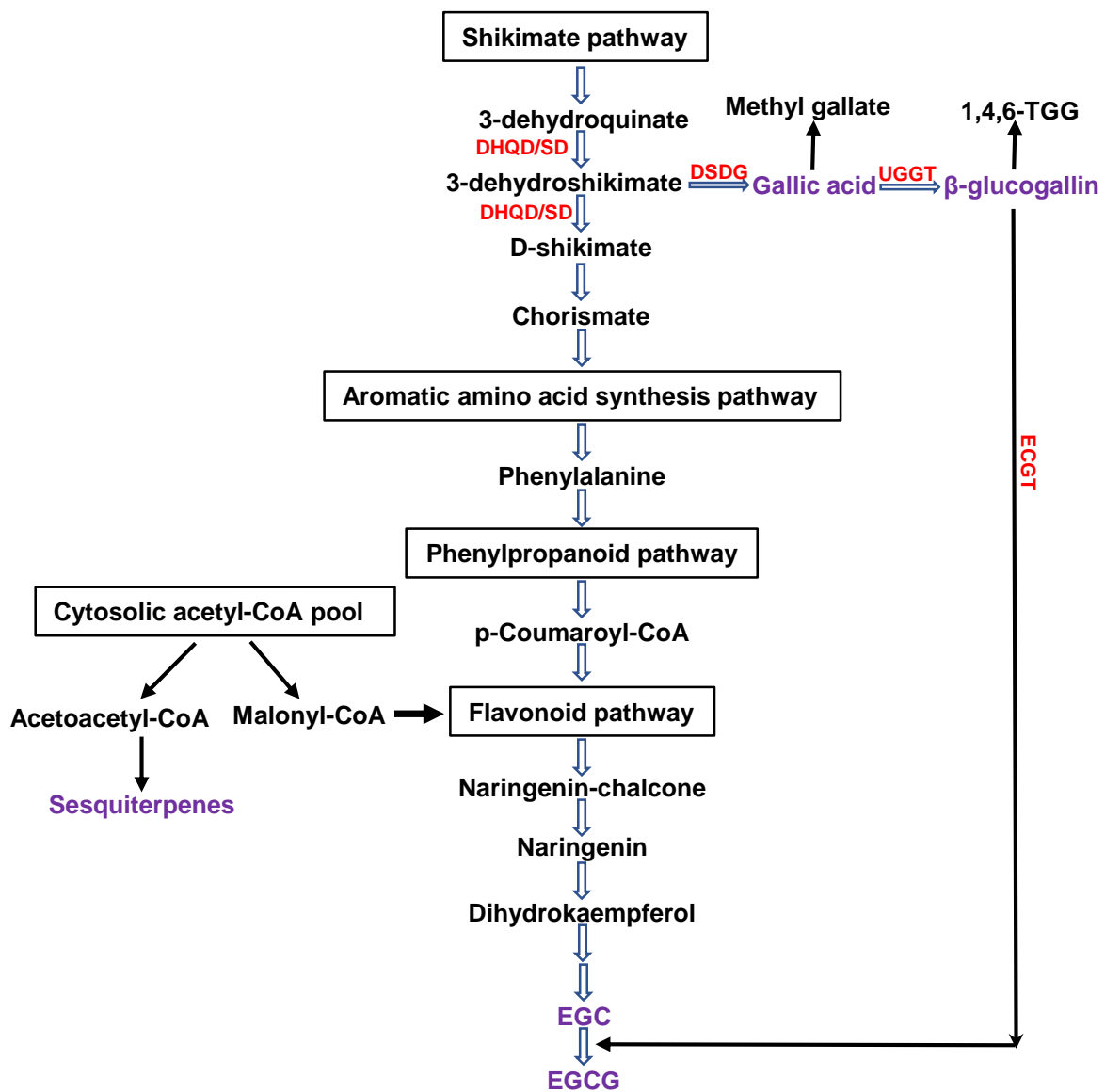
**Figure S2.** One bud and three leaves of the albino half sibs descendent from *Camellia sinensis* cv *Baijiguan*. Bar = 1 cm.



**Figure S3.** The contents of total amino acids, caffeine, and catechins between the albino half-sibs and the green half-sibs



**Figure S4.** The total volatile contents of individual pathway between the albino half-sibs and the green half-sibs



**Figure S5.** A simplified diagram of catechin synthesis pathway from shikimate. DHQD/SD, 3-dehydroquininate dehydratase/shikimate dehydrogenase; DSDG, dehydroshikimate dehydro-genase; UGGT, UDP-glucose:galloyl-1-O-β-D-glucosyl-transferase; ECGT, epicatechin:1-O-galloyl-β-D-glucose O-galloyltransferase; EGC, (-)-epicatechin gallate; EGCG, (-)-epigallocatechin gallate;

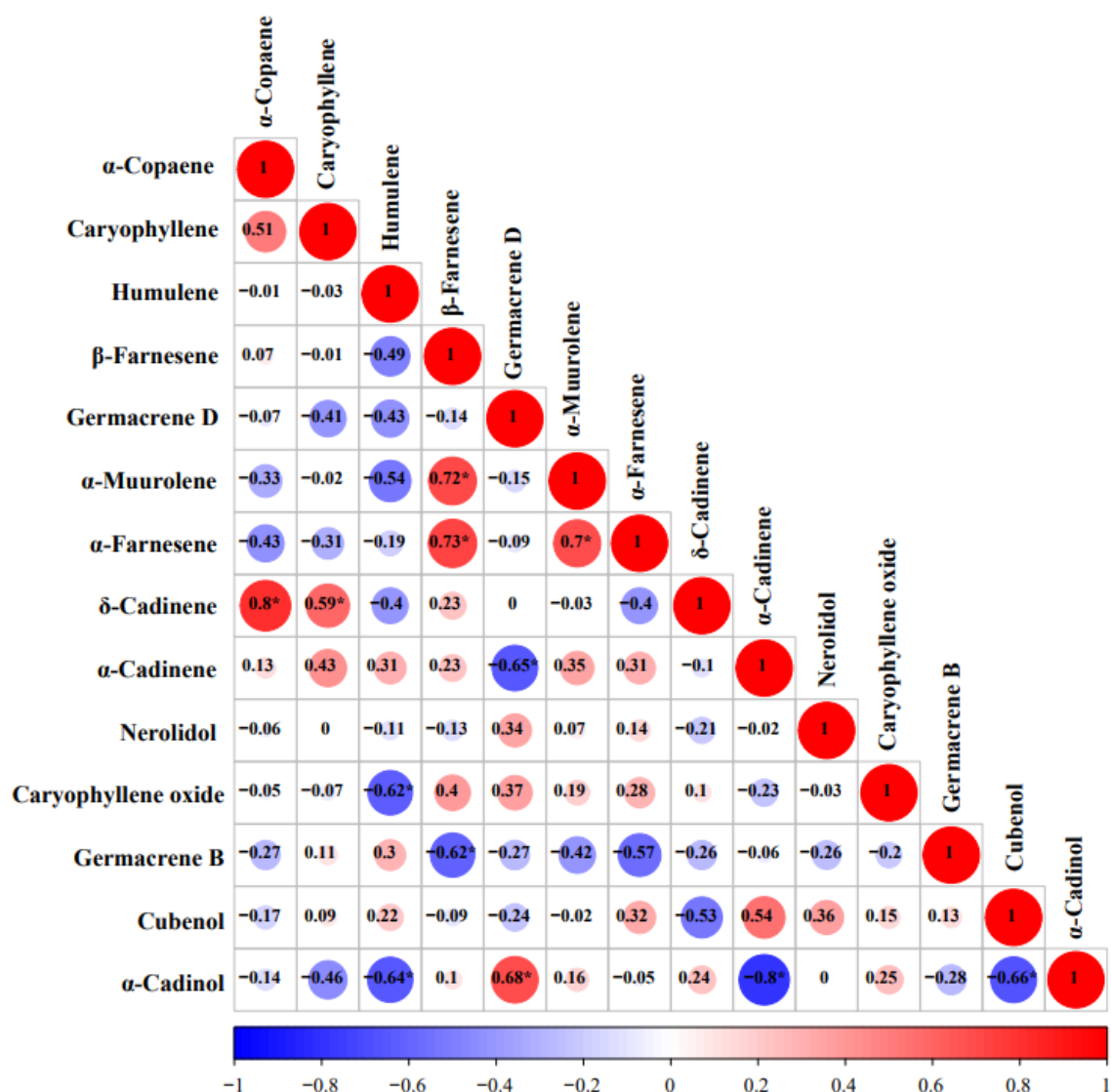
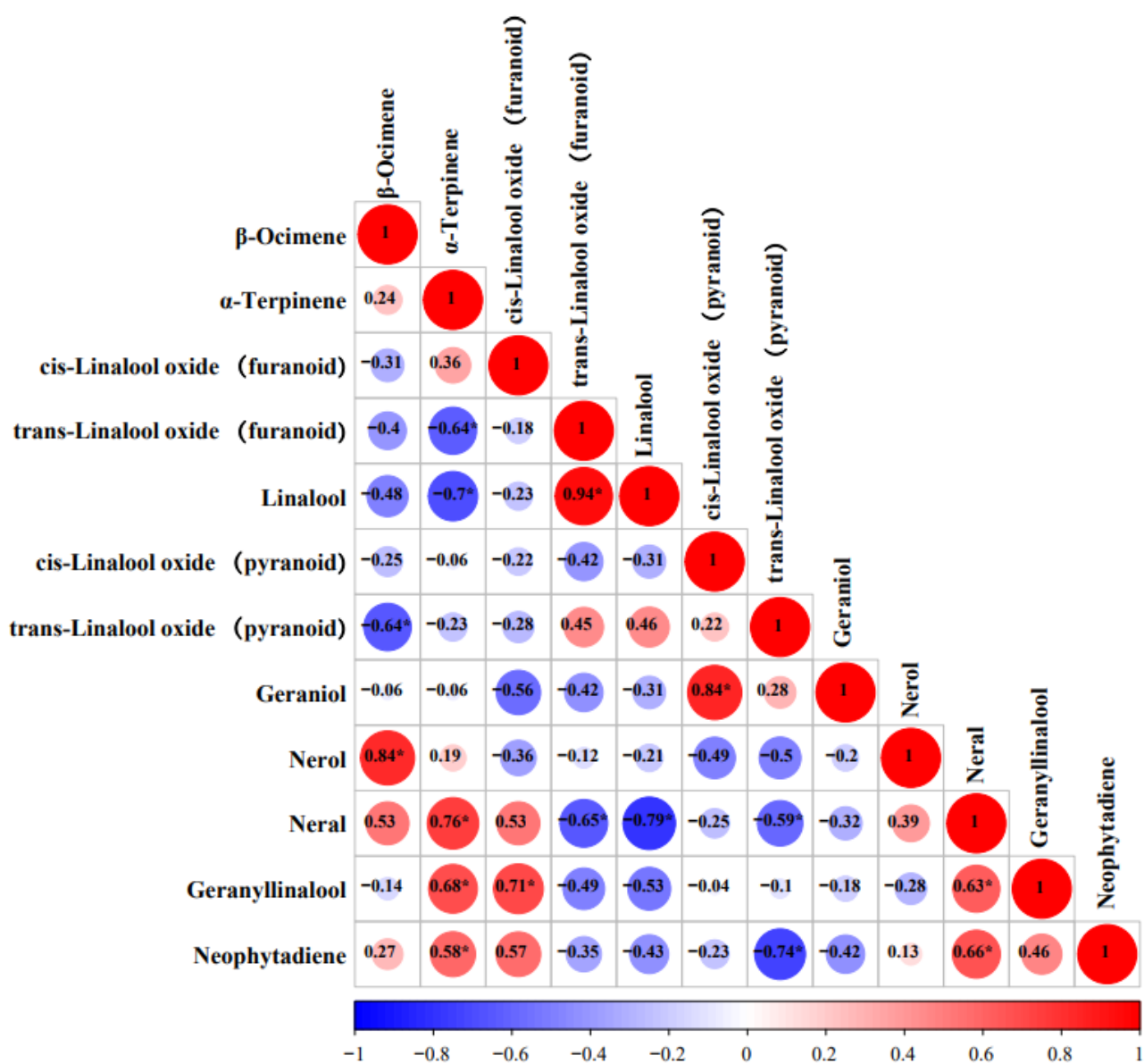
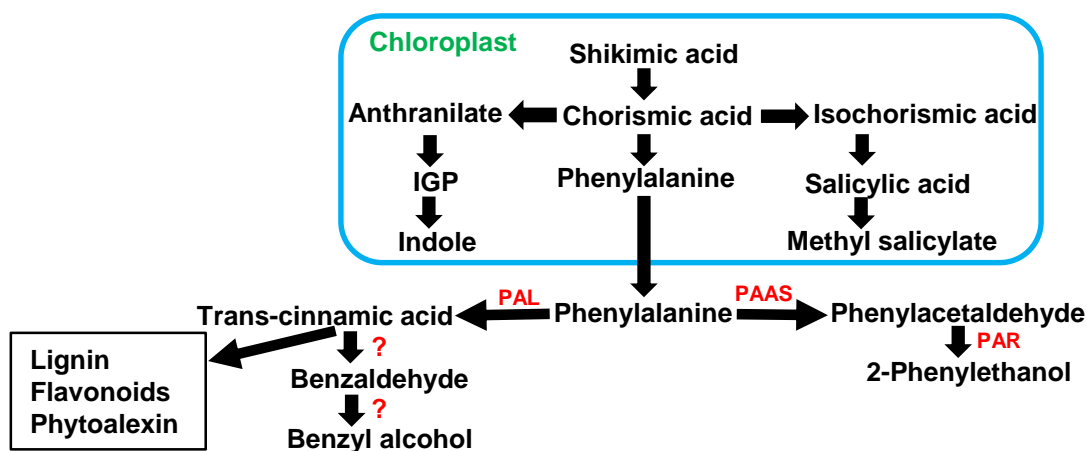


Figure S6. The correlations of volatiles derived from MVA pathway



**Figure S7.** The correlations of volatiles derived from MEP pathway





**Figure S8.** A simplified diagram presentation of shikimate–phenylpropanoid–benzenoid volatile synthesis pathways. PALs, phenylalanine ammonia lyases; PAAS, phenylacetaldehyde synthase; PAR, phenylacetaldehyde reductase