



Figure S2. Highly distributed metabolites and highly expressed genes in different parts of *D. nobile*. Highly accumulated metabolites in roots (A), stems (C), leaves (E), flowers (G), and fruits (I). Highly expressed genes in roots (B), stems (D), leaves (F), flowers (H), and fruits (J). Only the metabolites with more than 1% distribution portion and the genes with more than 0.5% expression portion were shown here.