



Figure S4. OOMTs, AADCs, and PARs amino acid sequences analysis. (a) Domain analysis of OOMTs amino acid sequences from *R. chinensis*. The green box indicates the dimerization domain. The orange boxes represent the AdoMet_MTases superfamily domain. (b) Phylogenetic analysis of AADCs amino acid sequences from *R. chinensis* and other plants. Protein IDs were shown in Table S6. (c) Motif analysis of PARs amino acid sequences from *R. chinensis* and other plants.