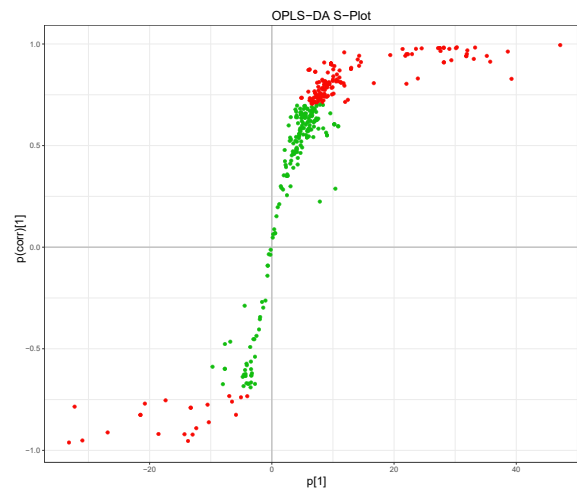
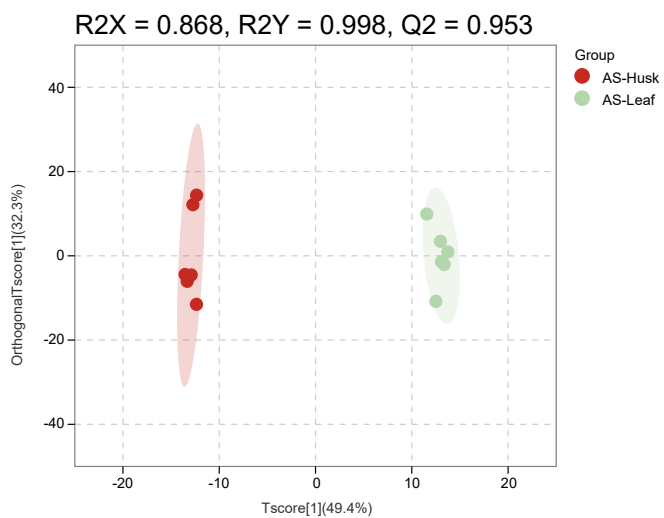
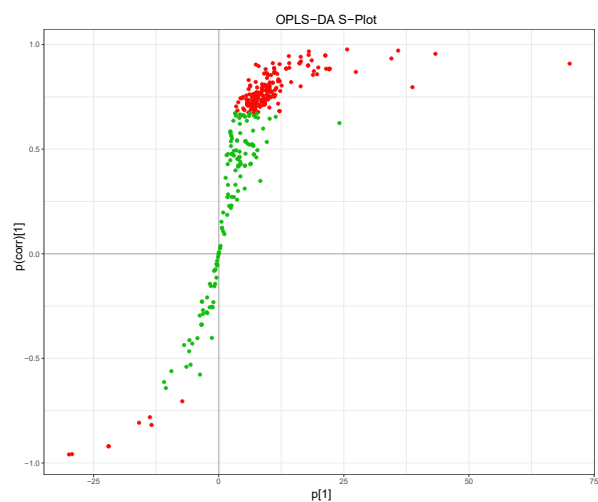
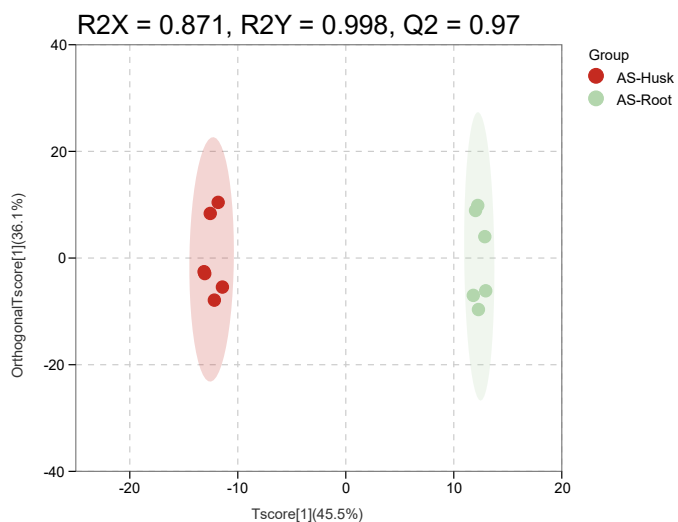
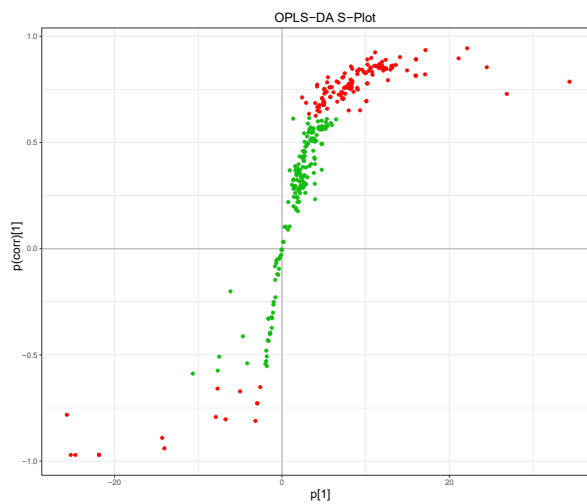
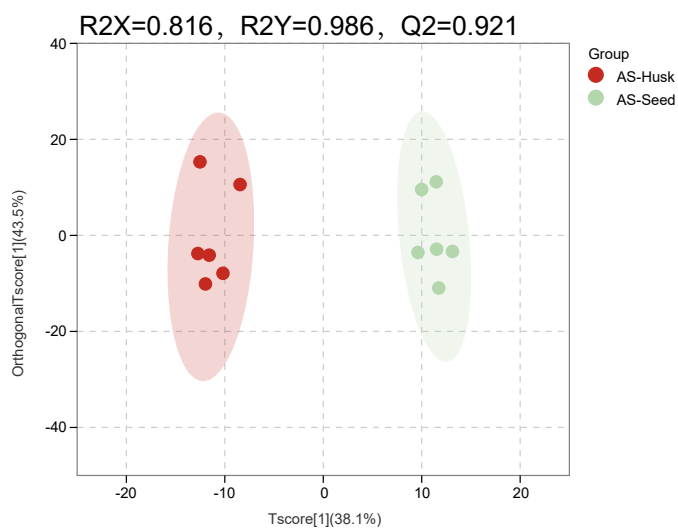
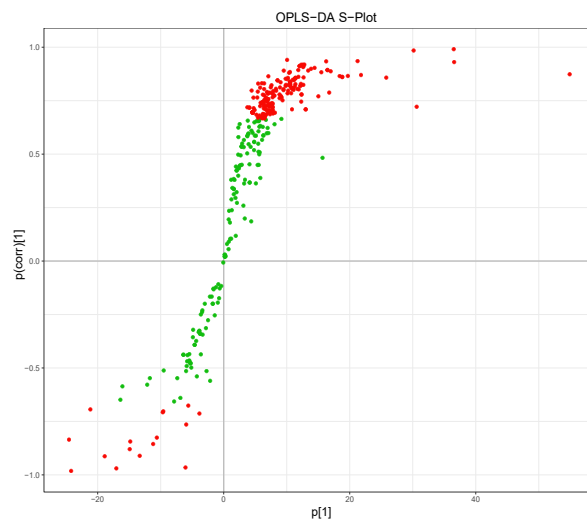
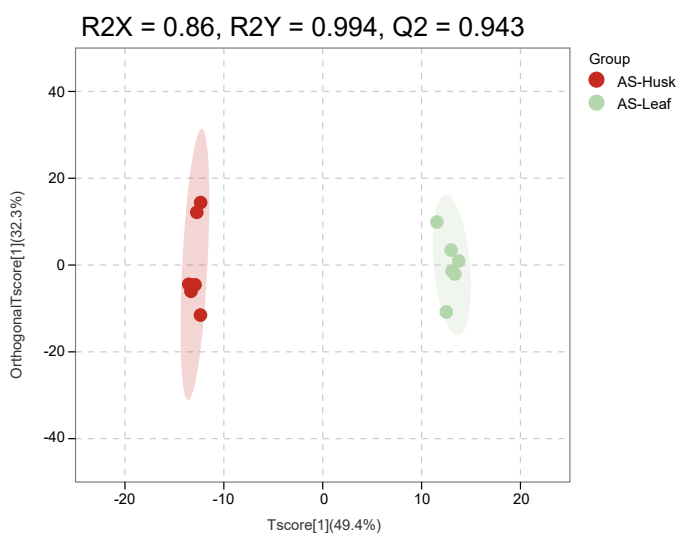
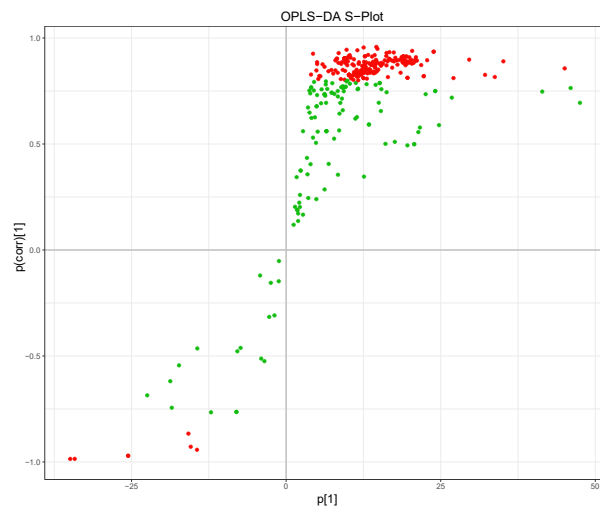
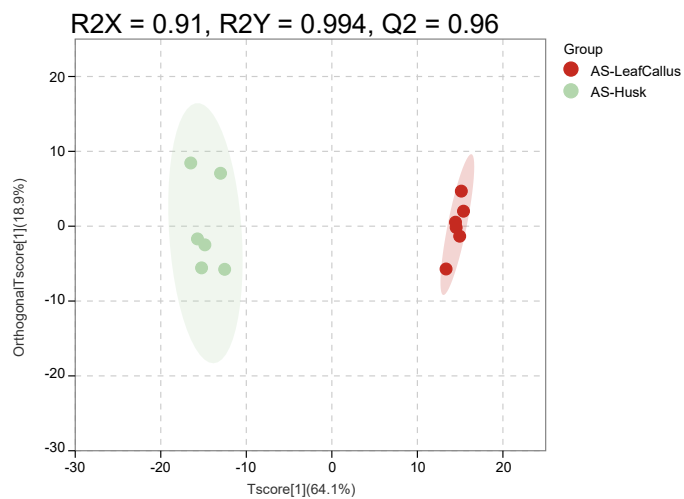
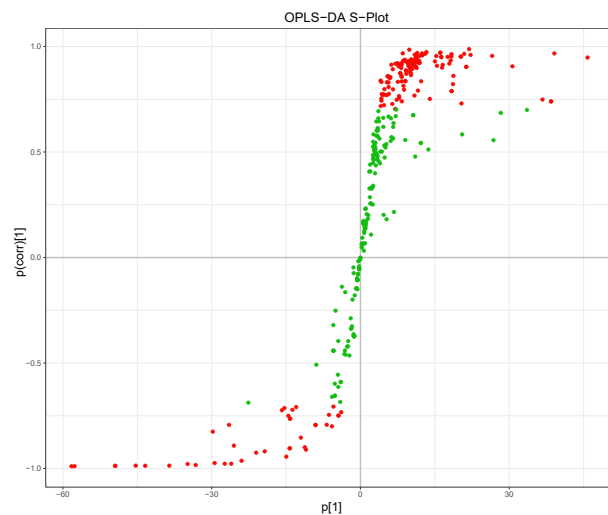
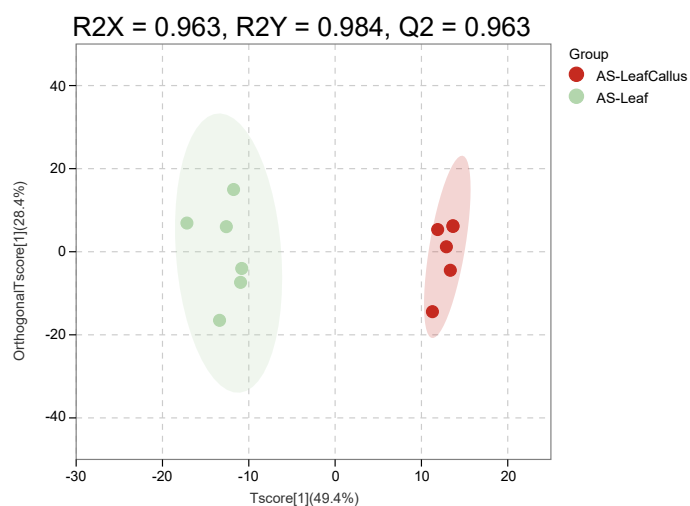
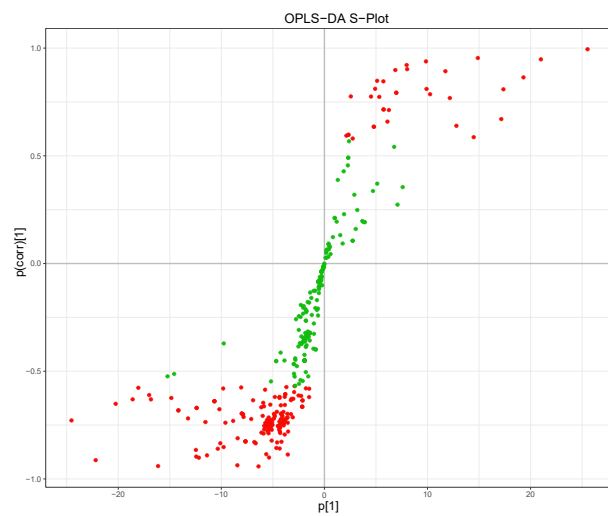
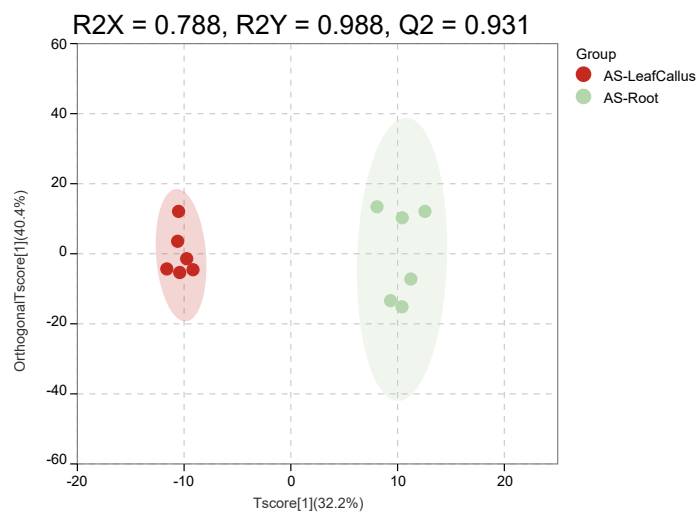
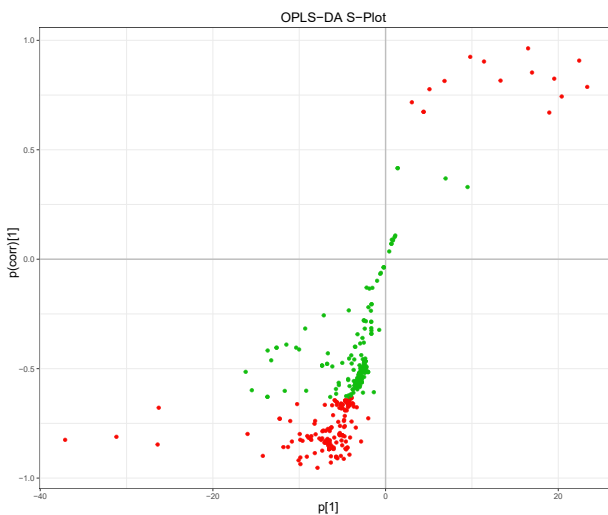
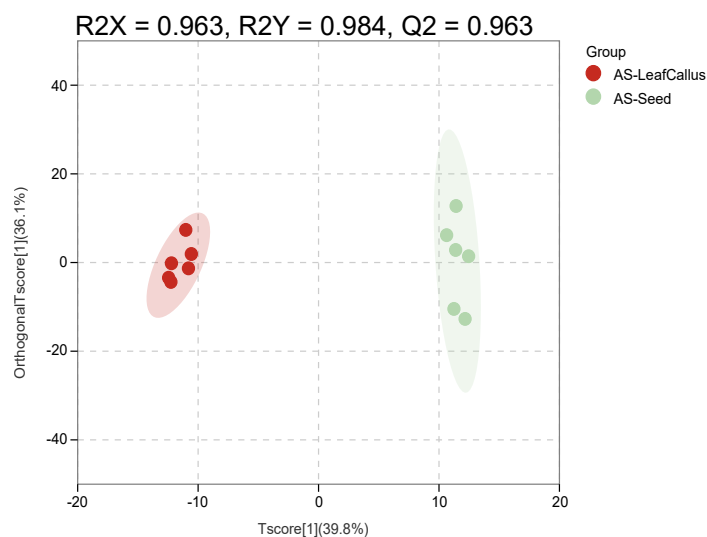
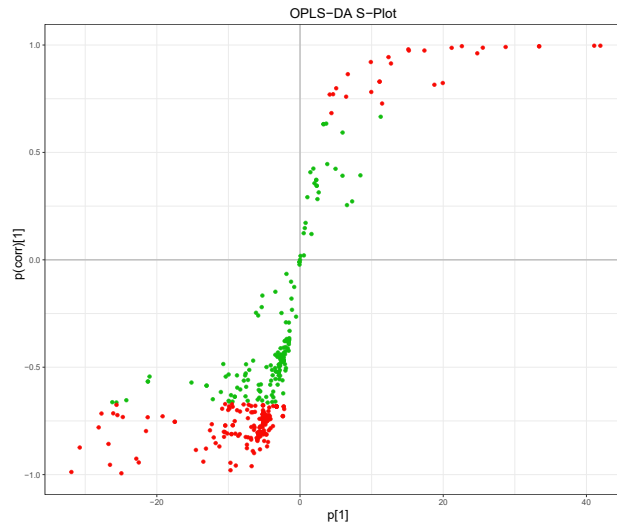
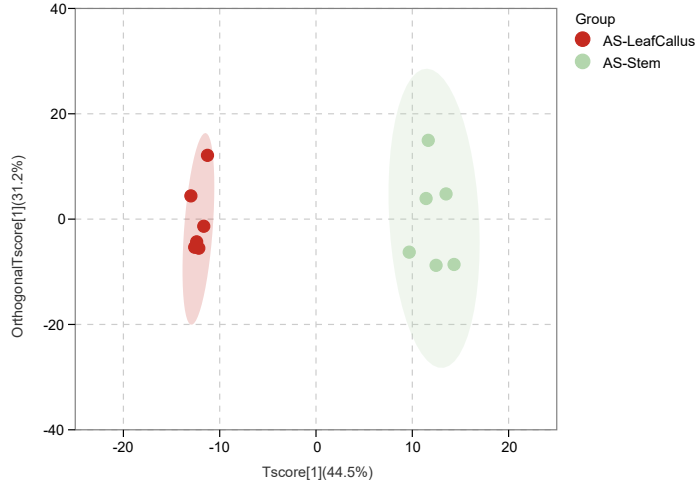


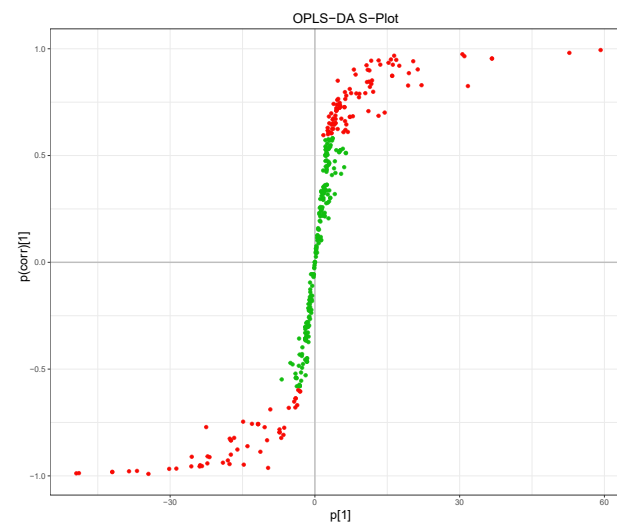
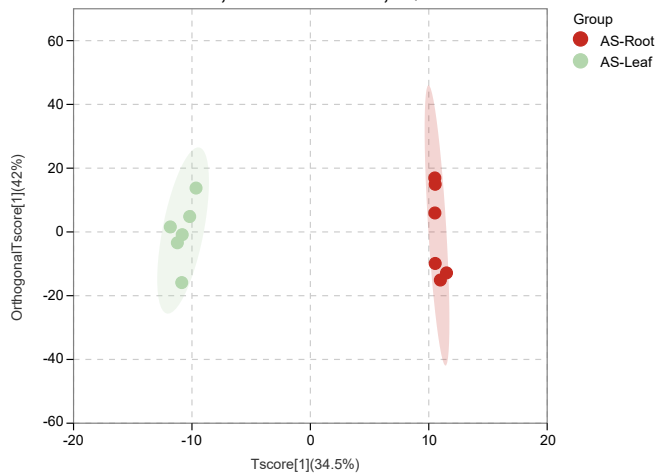
A**B****C****D**

E**F****G****H**

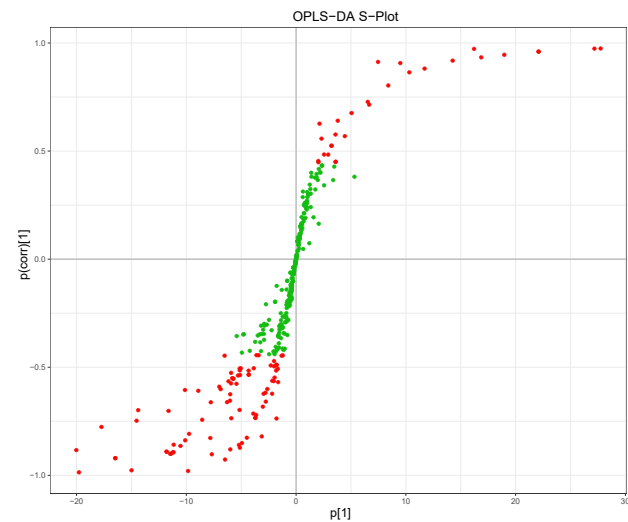
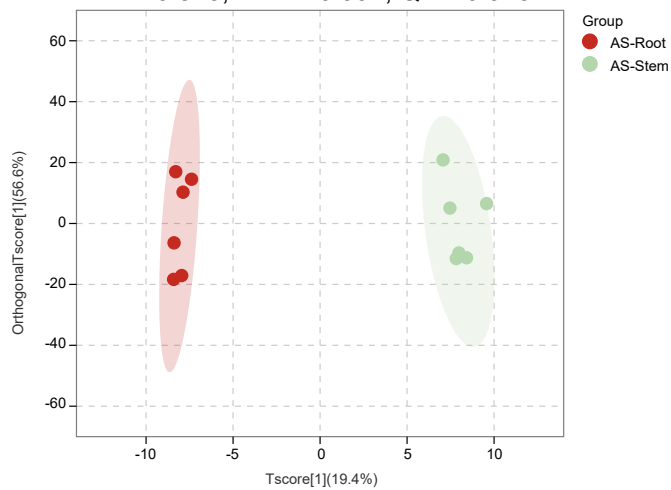
I R2X = 0.757, R2Y = 0.991, Q2 = 0.953



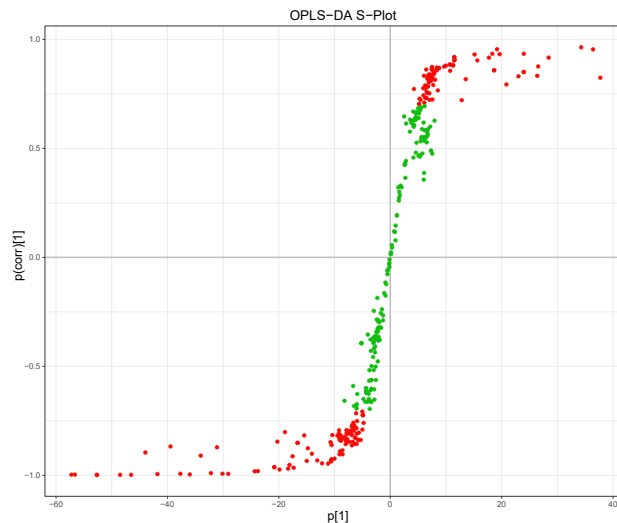
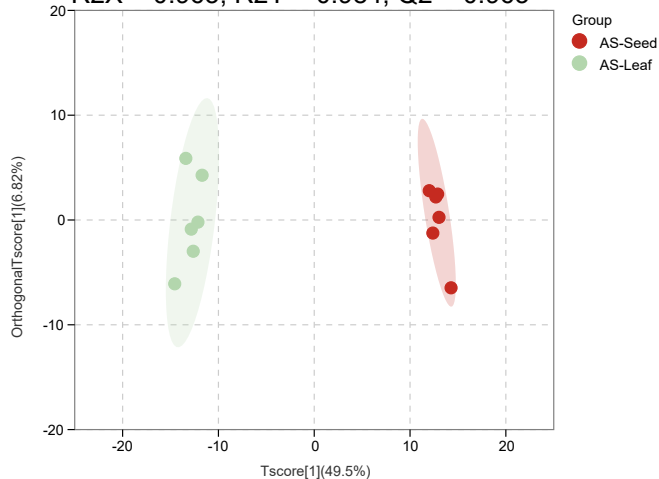
J R2X = 0.839, R2Y = 0.997, Q2 = 0.979



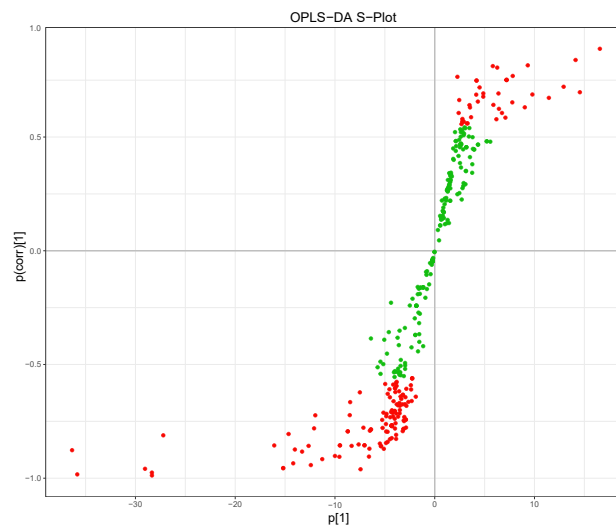
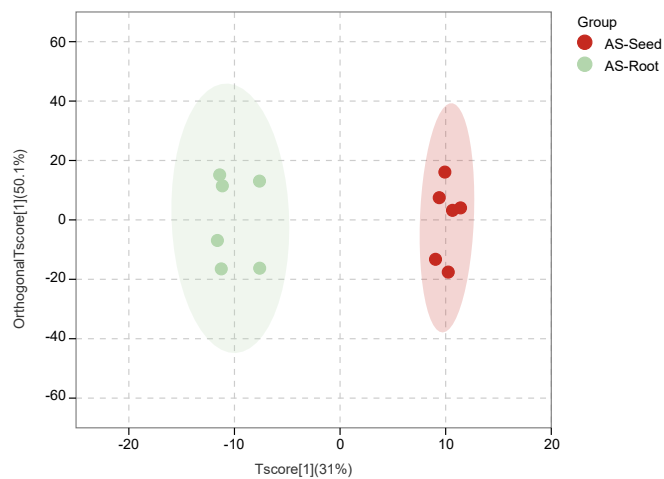
K R2X = 0.813, R2Y = 0.994, Q2 = 0.945



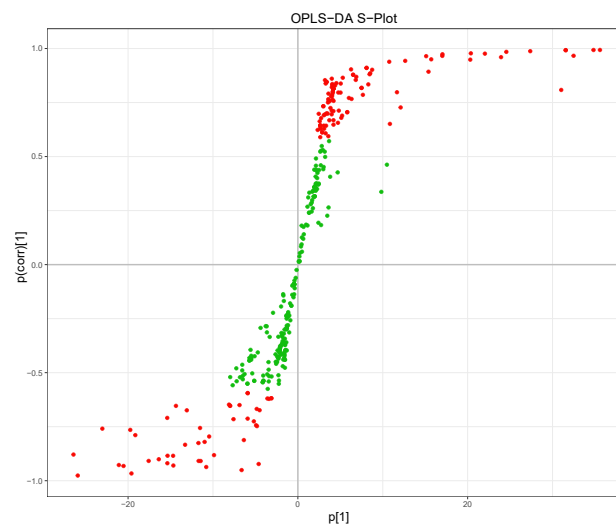
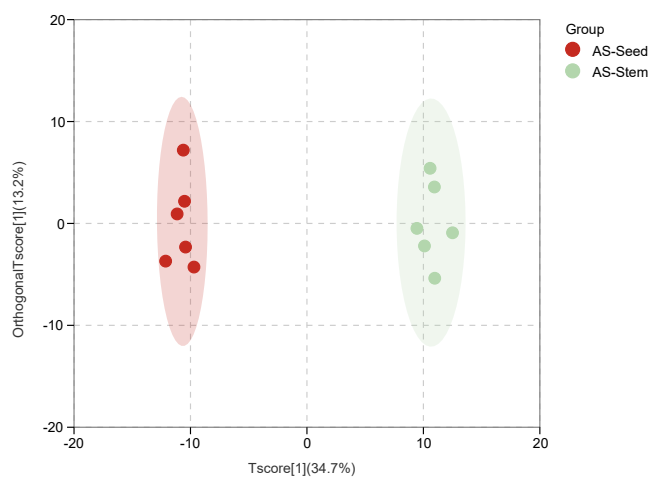
L R2X = 0.963, R2Y = 0.984, Q2 = 0.963



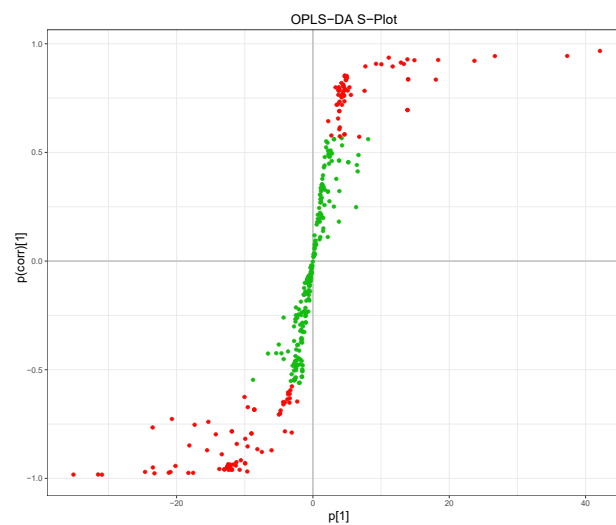
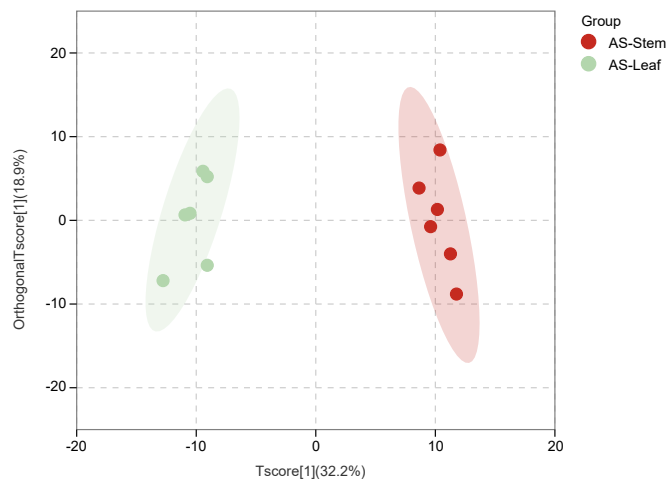
M



N



O



Supplementary Figure S2. Orthogonal partial least squares-discriminant analysis (OPLS-DA) scores and OPLS-DA S-plot models between six different tissue sites. (A) AS-Husk_vs_AS-Leaf, (B) AS-Husk_vs_AS-Root, (C) AS-Husk_vs_AS-Seed, (D) AS-Husk_vs_AS-Stem, (E) AS-LeafCallus_vs_AS-Husk, (F) AS-LeafCallus_vs_AS-Leaf, (G) AS-LeafCallus_vs_AS-Root, (H) AS-LeafCallus_vs_AS-Seed, (I) AS-LeafCallus_vs_AS-Stem, (J) AS-Root_vs_AS-Leaf, (K) AS-Root_vs_AS-Stem, (L) AS-Seed_vs_AS-Leaf, (M) AS-Seed_vs_AS-Root, (N) AS-Seed_vs_AS-Stem, (O) AS-Stem_vs_AS-Leaf.