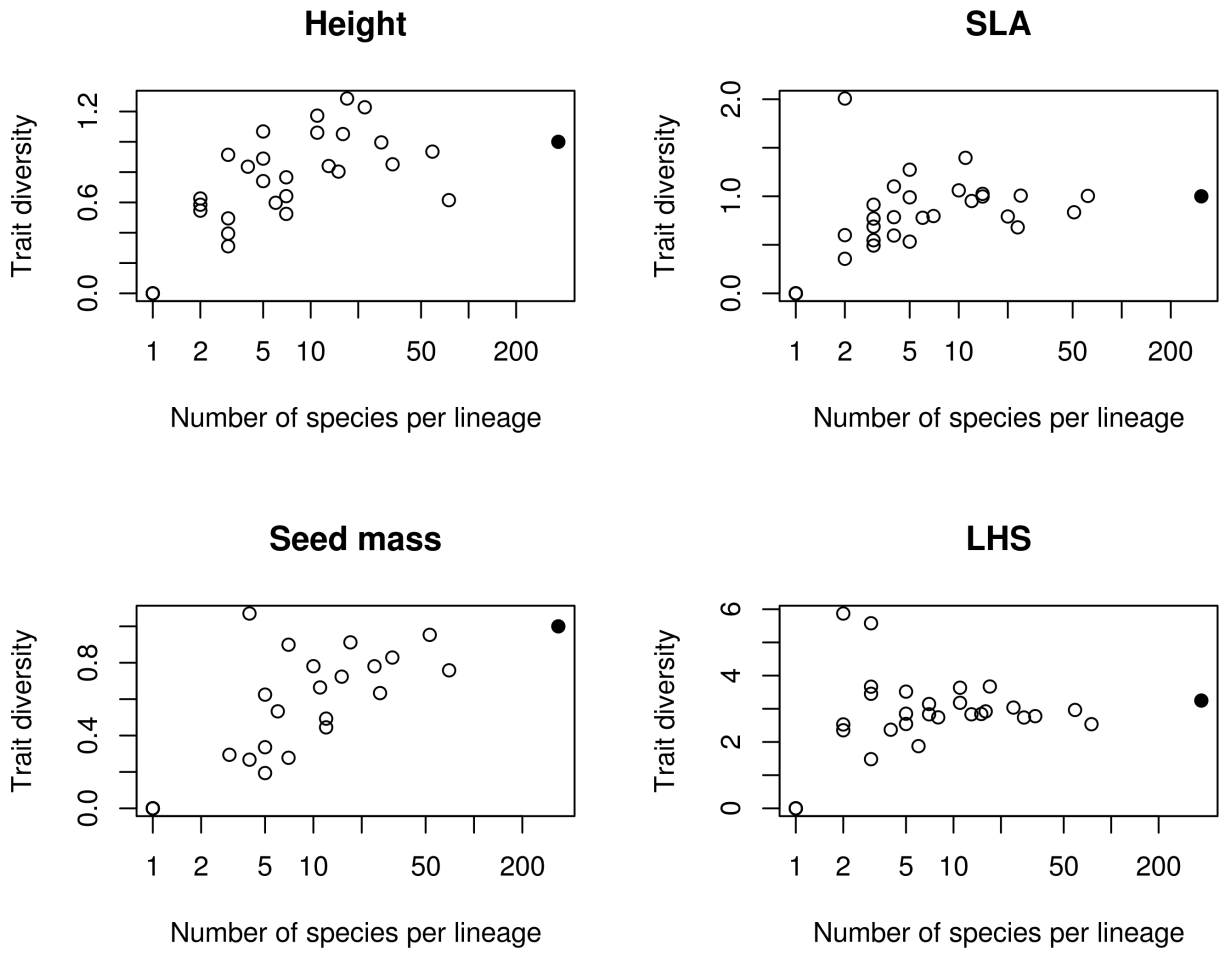Figure S5. Relationship between trait diversity and number of species per lineage (from the reduced species pool at an intermediate evolutionary scale), used to generate the Figure 3. All trait values were log-transformed and standardized. Trait diversity was evaluated as the standard deviation of the trait values within each lineage.. «LHS» (Leaf-Height-Seed) refers to the multi-trait diversity index calculated as the mean Euclidean trait distance between species of a lineage. The black dot represents the diversity of the traits among all species of all lineages.