

Supplemental Figure S1. Interaction network analysis of CRM genes. A network view of the CRM genes.

Cross-hatched bubbles are the CRM genes while solid bubbles indicate partners identified through network analysis. Connections are color-coded based on the type of interaction. Red, physical interaction; purple, co-expression; orange, predicted; blue, co-localization; turquoise, pathway; green, genetic interactions; yellow, shared protein domains.

