



Figure S3. Heat map for the two-dimensional genome scan with a two-QTL model in the KD-F2 plants.

Top left triangle: the maximum LOD scores for the interaction model. Bottom right triangle: the maximum LOD scores for the full model (two QTLs plus an interaction). The color scale indicates the values for the interaction model on the left (LOD threshold = 6.52) and the full model on the right (LOD threshold = 9.03).