



S1 Figure. Correlation analysis of gene expression experiments (RNAseq or DNA microarray) using AtCAST. Experiments showing a significant correlation (Spearman's correlation coefficient (SCC) > 0.3) with "*max1-1 (max1-1 mutants compared with wild-type)*" were searched. Red arrows indicate strong positive relationships (SCCs > 0.7); blue arrows indicate strong negative relationships (SCCs < -0.65); pink arrows, moderate relationships (SCCs > 0.5); and light blue arrows indicate moderately negative relationships (SCCs < -0.5). Arrowhead directions indicate whether the interactions were uni- or bidirectional.