

Article

The Impact of Heat Stress on Morpho-Physiological Response and Expression of Specific Genes in the Heat Stress-Responsive Transcriptional Regulatory Network in *Brassica oleracea*

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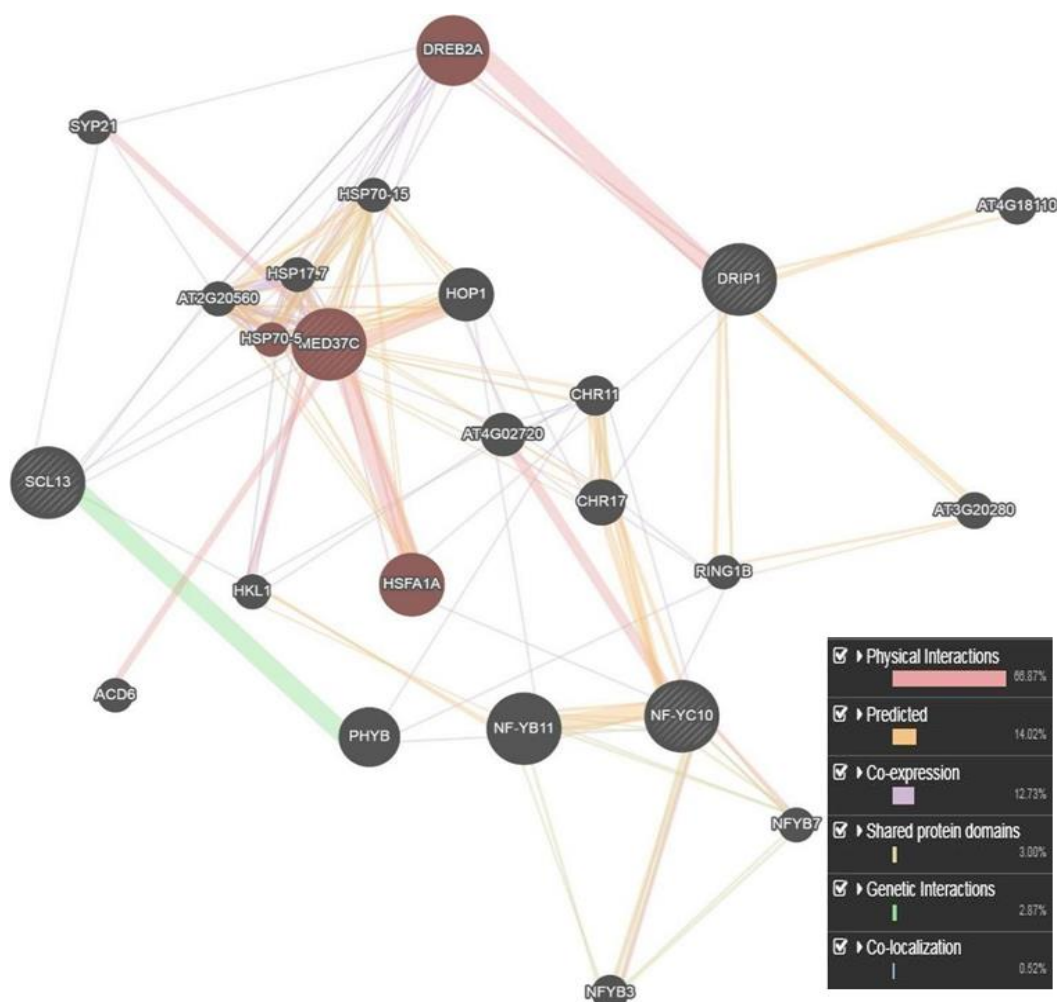


Figure S1. Gene interaction network in *Arabidopsis* for the genes of interest used for the gene expression study in this work. The network was generated by the GeneMANIA prediction server [44]. The right panel presents the different types of interactions with respective color coding, depicted with lines connecting genes in the network: physical interactions (pink), predicted (orange), co-expression (purple) and shared protein domains. Red-filled circles considered heat-shock protein binding.