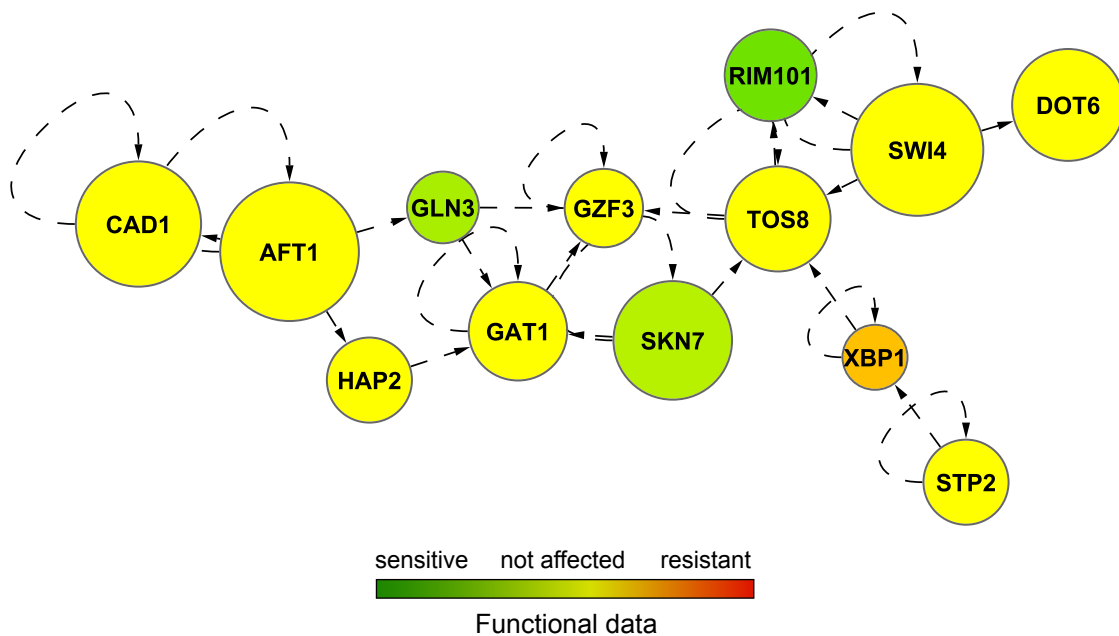


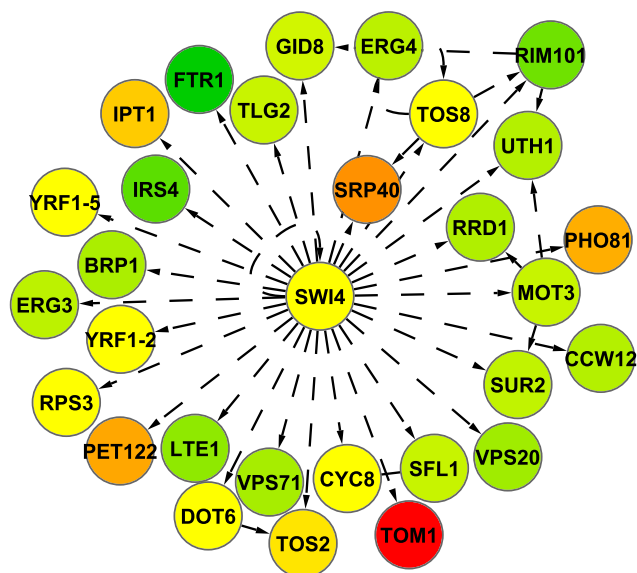
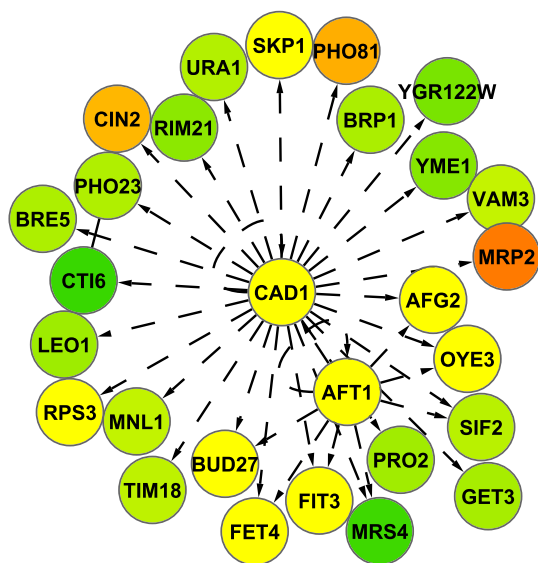
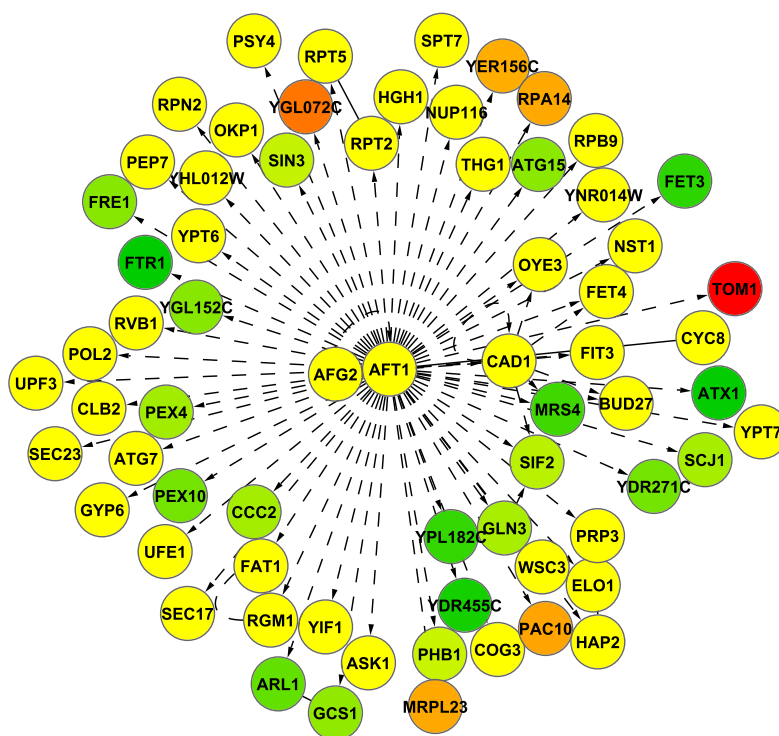
Novel Insights into iron metabolism by integrating deletome and transcriptome analysis in an iron deficiency model of the yeast *Saccharomyces cerevisiae*
Jo, Kim, Oh, *et al.* (2009)

Additional File 5: Functional identification of transcriptional response networks in iron deficiency

Click on transcription factor to visualize interactions

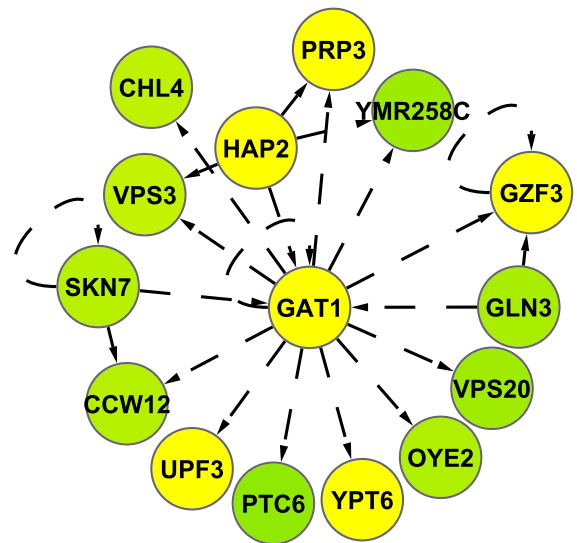
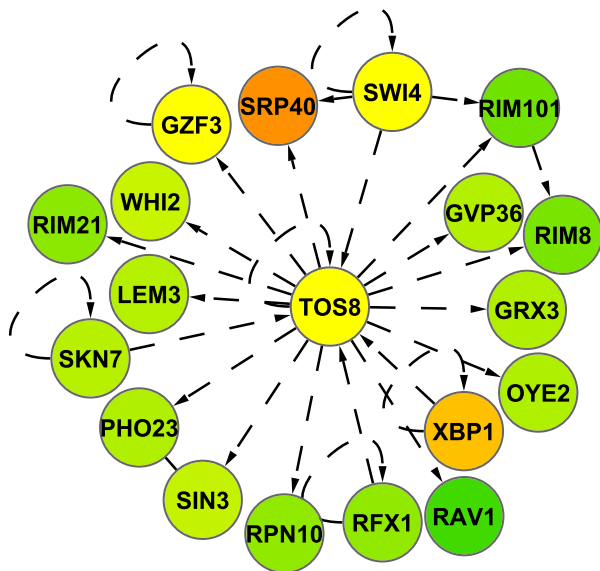
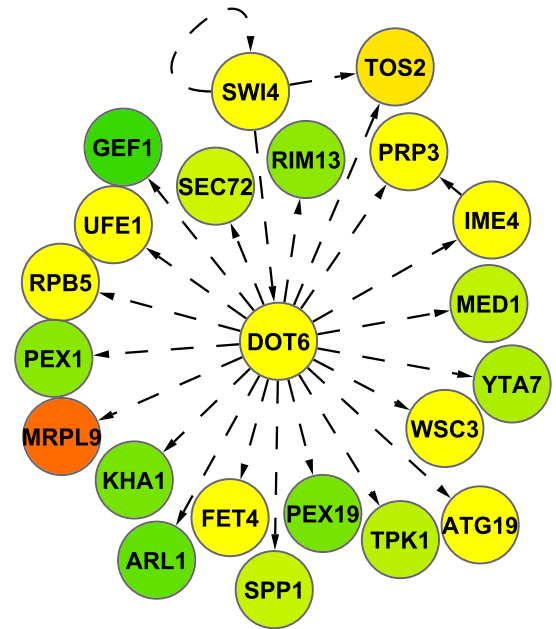
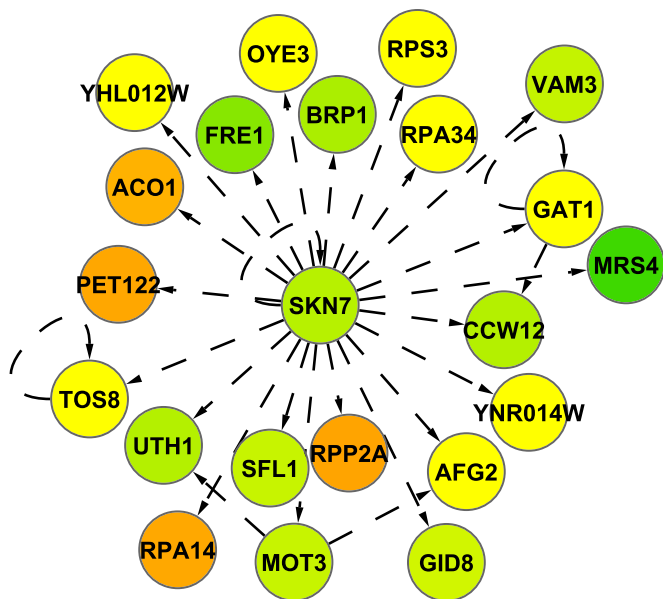


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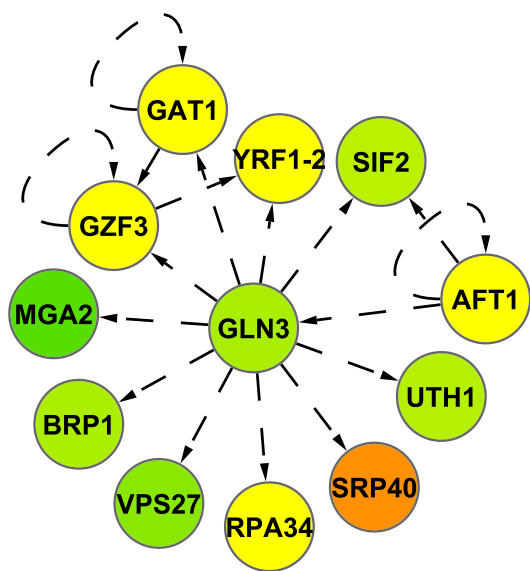
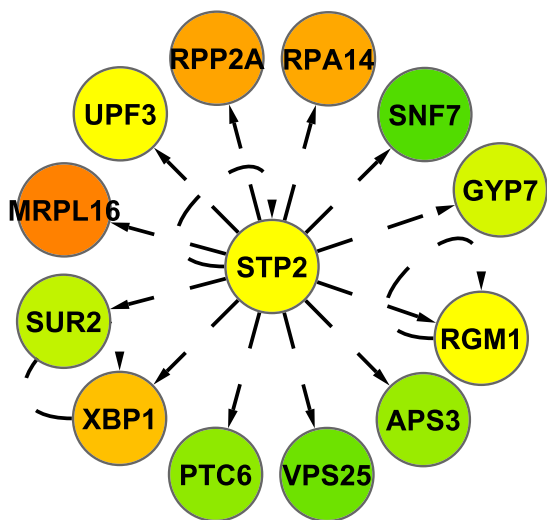
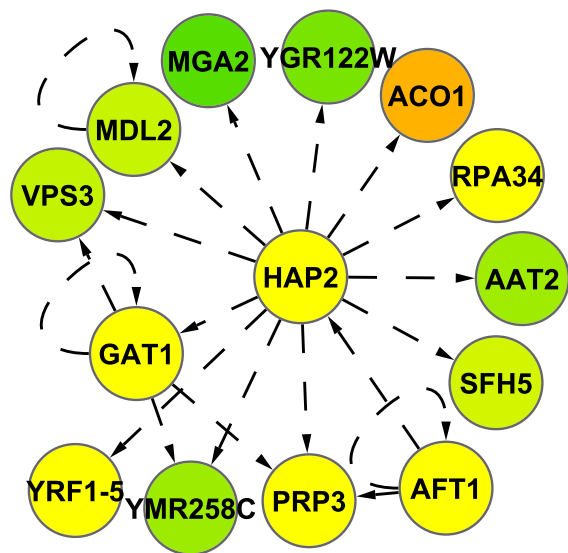
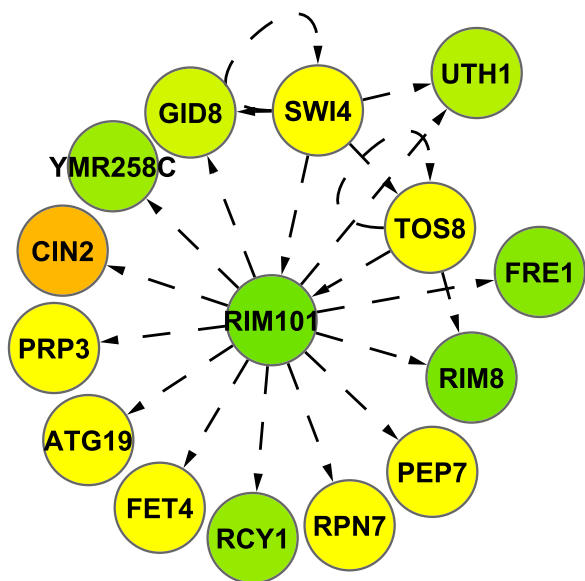

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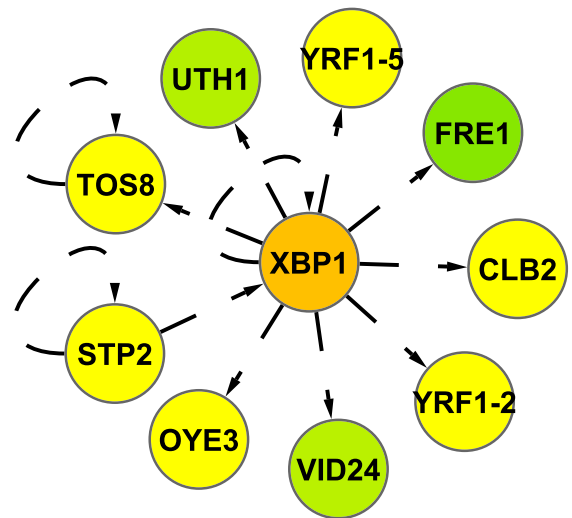
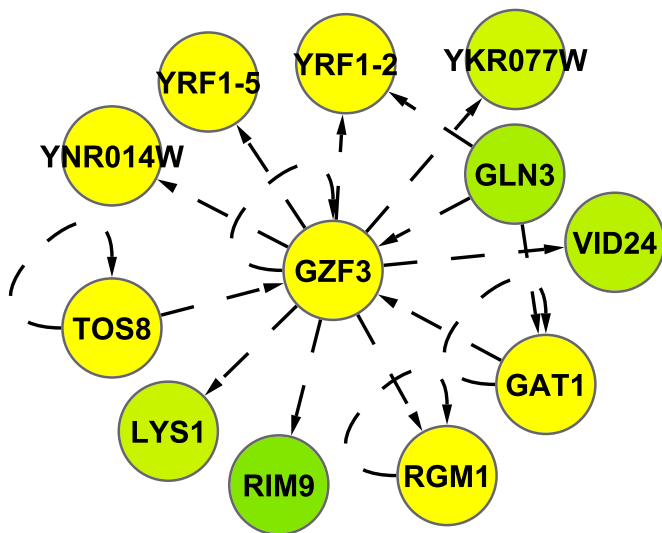

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