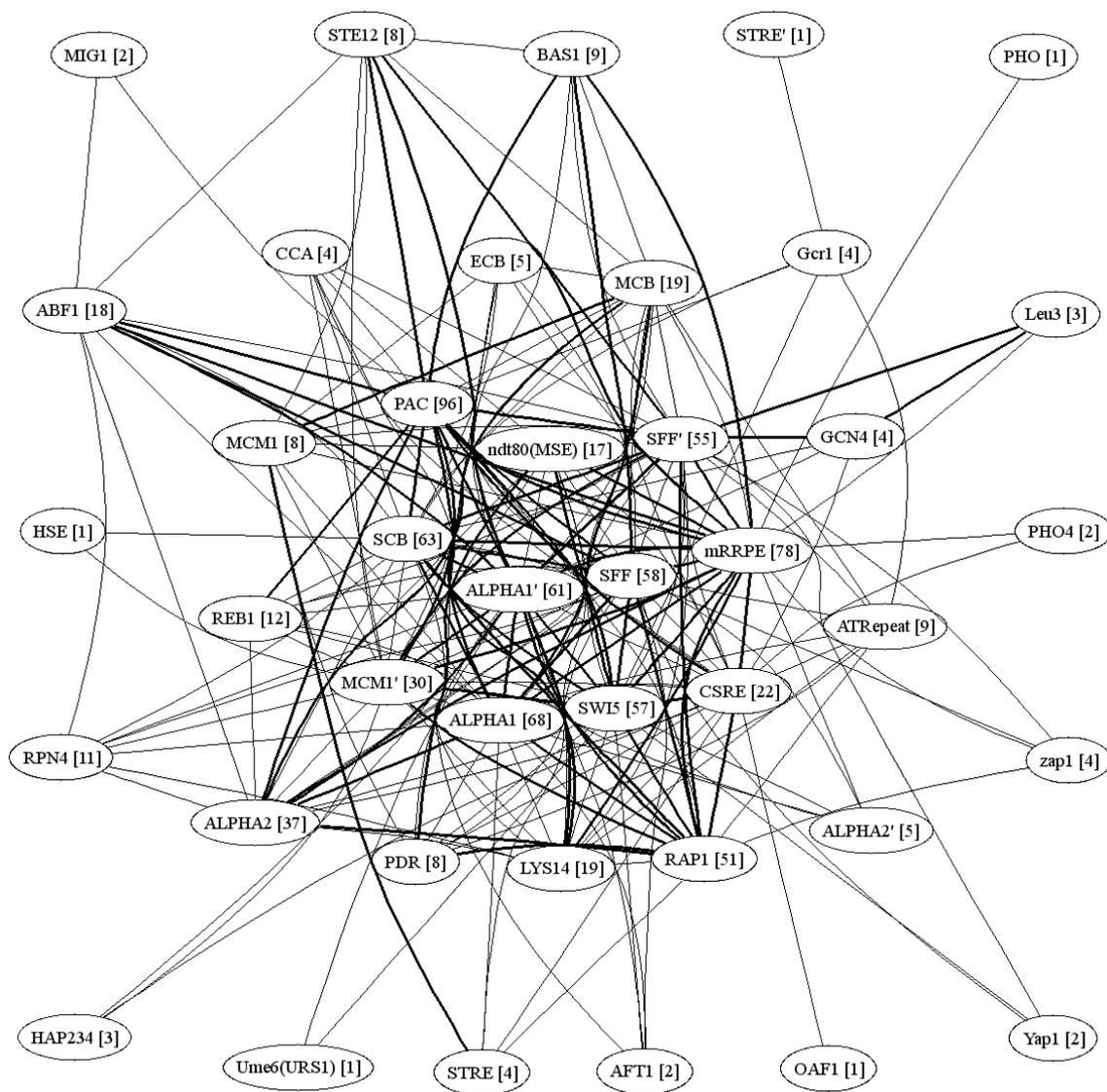
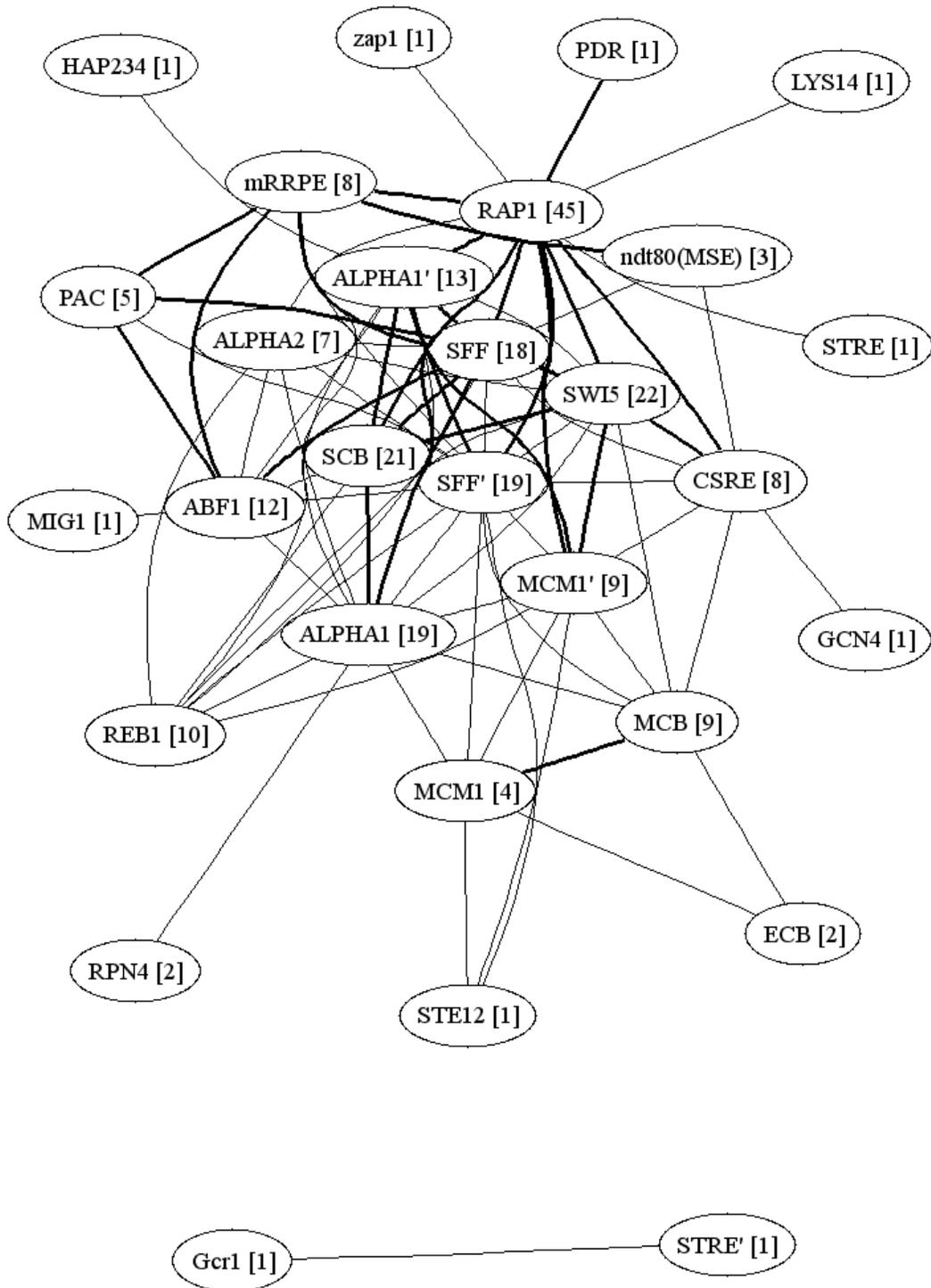


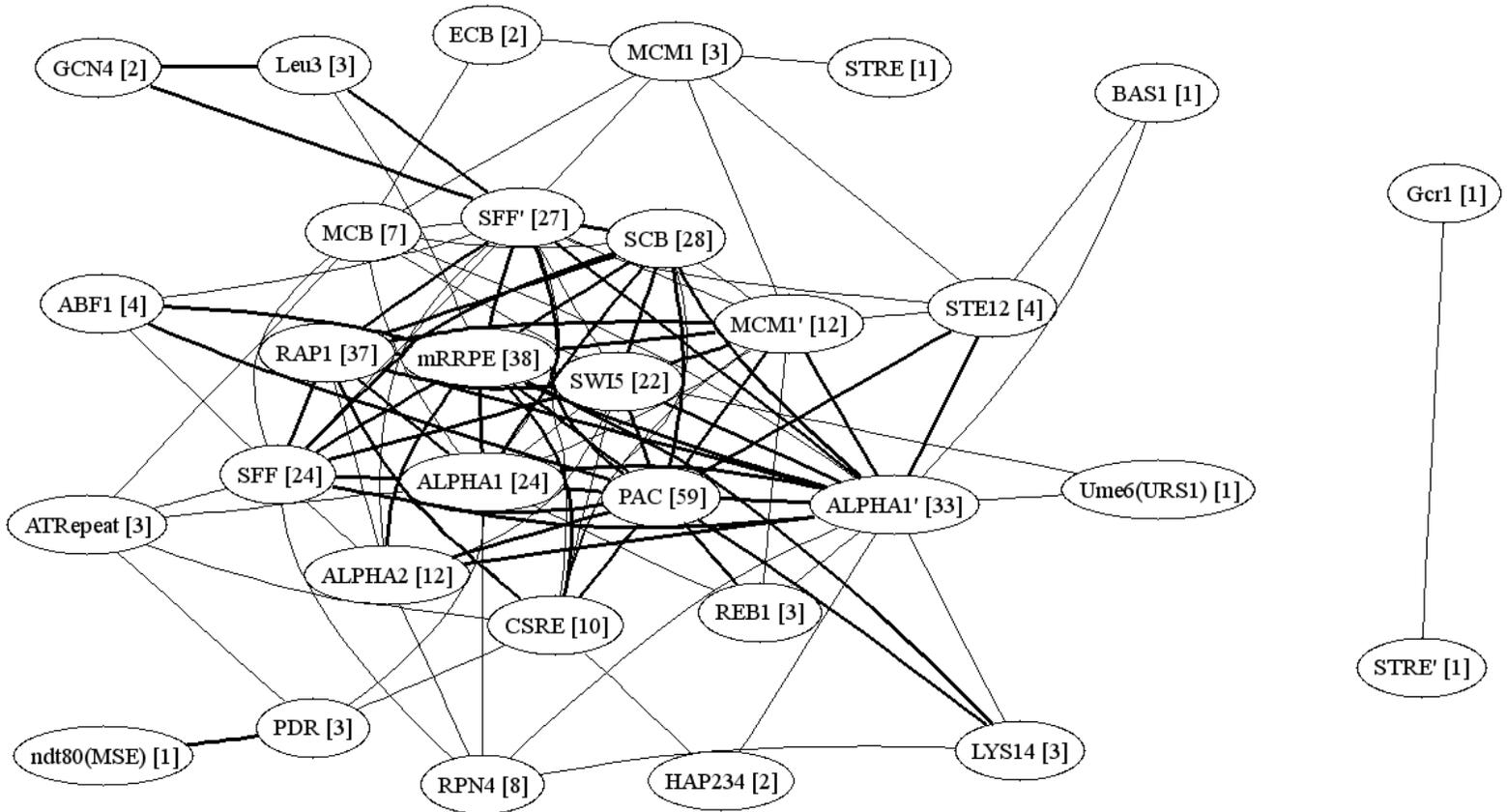
Figure 4 (EXTENDED). Graph showing which binding site pairs participate in the same binding site modules as hypothesized by our rules. Nodes are the binding sites and there is an edge between any two binding sites if they appear in the same rule (number of rules including a particular binding site is given in brackets). Bold edges indicate that the two binding sites appear in a rule that was induced from more than one expression dataset. The graph includes 41 of the 43 known binding sites. GAL and MET31-32 were not found in any rule. Corresponding graphs constructed using only significant rules according to each part of Gene Ontology and the binding data by Lee et al. (2002) are also shown.



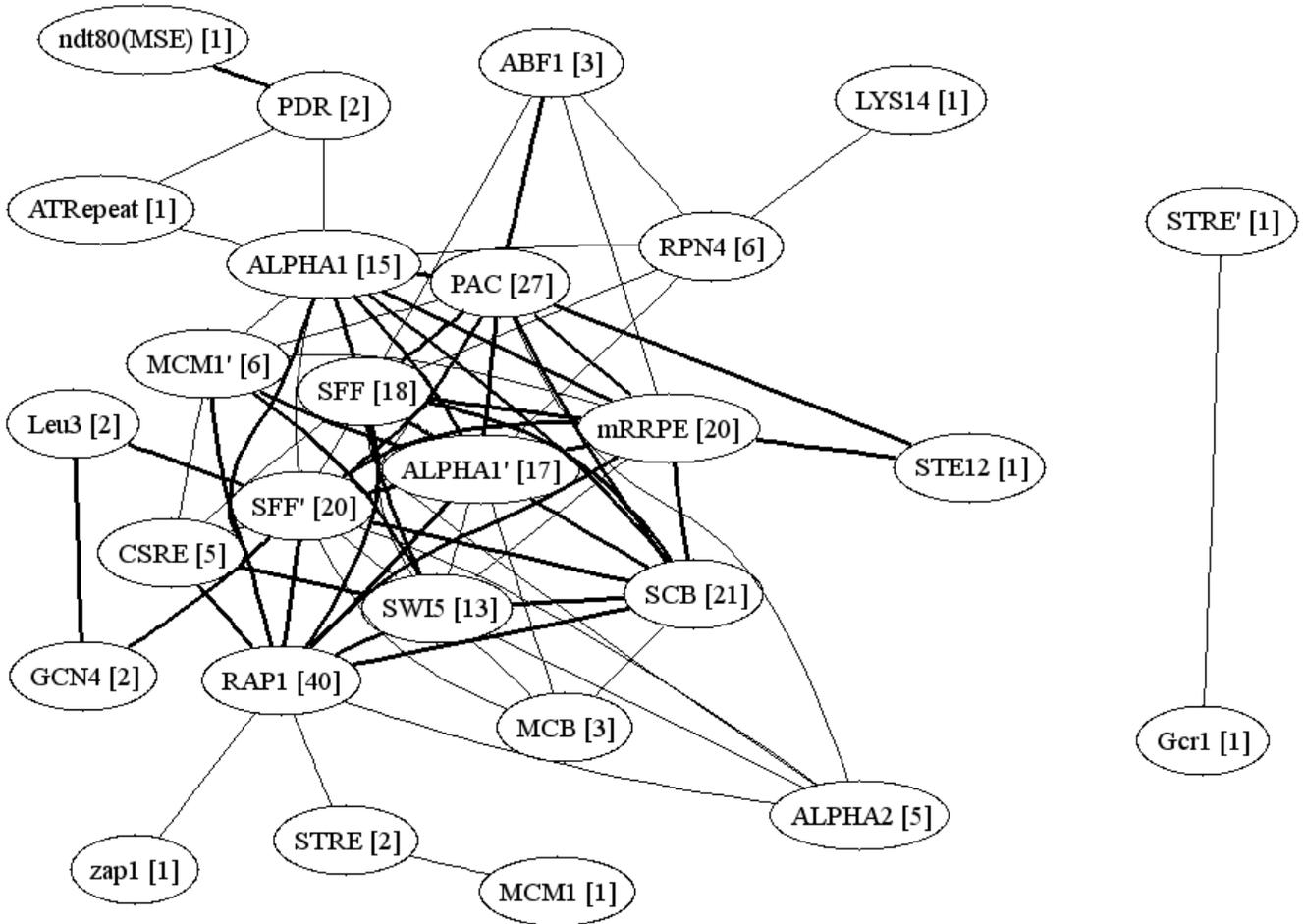
From significant rules according to the binding data by Lee et al. (2002):



From significant rules according to Gene Ontology biological process:



From significant rules according to Gene Ontology molecular function:



From significant rules according to Gene Ontology cellular component:

