



FIGURE 3. The results of a SGA screen are enriched for genes functionally related to the query gene, *BIM1*. Shown is a set of genetic lethal/sick interactions determined by SGA analysis with a *BIM1* gene deletion query mutation (Tong et al. 2004). The genes are colored according to the Gene Ontology terms in the SGD database. Annotation of the protein–protein interactions is based on the DIP, MINT, BIND, and IntAct databases. SGA genetic interaction data are enriched for proteins that physically interact with each other or are in the same protein complex or pathway.

Protein–Protein Interactions: A Molecular Cloning Manual, 2nd Ed., © 2005 by Cold Spring Harbor Laboratory Press, Chapter 40, Figure 3.