



**FIGURE 1.** AutoDockTools is a graphical user interface for preparing, running, and analyzing docking experiments with AutoDock. In this figure, the researcher is defining the extent of the affinity grid around the target molecule, trypsin, shown in red. Coordinates were taken from entry 2ptc at the Protein Data Bank (<http://www.pdb.org>).

*Protein-Protein Interactions: A Molecular Cloning Manual*, 2nd Ed., © 2005 by Cold Spring Harbor Laboratory Press, Chapter 45, Figure 1.