



FIGURE 7.24. Tree of 205 core proteins from γ -proteobacterial genomes. The topology shown agrees with almost all individual gene alignments. The same tree is obtained after removing the two genes showing evidence for lateral gene transfer. The position of the root corresponds to the one obtained repeatedly using SSU (small subunit) rRNA.

7.24, redrawn from Lerat E. et al., *PLoS Biol.* 3: E130, © 2005 Public Library of Science