Figure S1. Single gene trees inferred from ML analyses with 1,000 replicates. Bootstrap scores for all nodes are displayed above the branches. Those taxa different from the combined tree of 232 genes (Figure 1) are bolded. *Lb. delbrueckii subsp. bulgaricus* 1 = *Lb. delbrueckii subsp. bulgaricus* ATCC BAA-365; *Lb. delbrueckii subsp. bulgaricus* 2 = *Lb. delbrueckii subsp. bulgaricus* ATCC 11842. a, *uvrB* gene; b, *polC* gene; c, *phpB* gene.