Supplementary Figure S1. Phylogenetic tree of *Peronosporales* and *Pythiales* species from the minimum evolution analysis based on *cytochrome c oxidase* subunit I (*cox*1) mtDNA sequences. Bootstrapping values (minimum evolution BP/maximum likelihood BP) higher than 70 % were given above or below the branches (1,000 replicates). The strains obtained from this study are shown in bold. The scale bar equals the number of nucleotide substitutions per site.