



Fig. 2. Alignment of *Candidatus* *Phytoplasma solani* DNA sequences from the GenBank NCBI database (<http://www.ncbi.nlm.nih.gov/BLAST>) with those from DNA sequencing of infected grape samples with primers to *SecY* gene. Alignments were generated in the Unipro UGENE program (Russia) version 1.31.1, the attachment zone for the forward primer *BN_F* is highlighted with a white frame. *Phytoplasma solani* 1-6 — samples collected in six different habitats.