



Figure S3. A phylogenetic tree constructed of 16S rRNA gene sequences from phytoplasma by the neighbor-joining method using MEGA 6. Numbers on branches are the bootstrap values (1000 replicates). GenBank accession numbers are given in parentheses, and *Acholeplasma Laidlawii* was the outgroup. The bold represented the F2nR2 regions sequences of 16Sr RNA gene amplified from SPLDaP in SPLD leaf samples (-A) and corresponding mealybugs samples (-M) from Hainan (HN), Guangdong (GD) and Guangxi (GX).