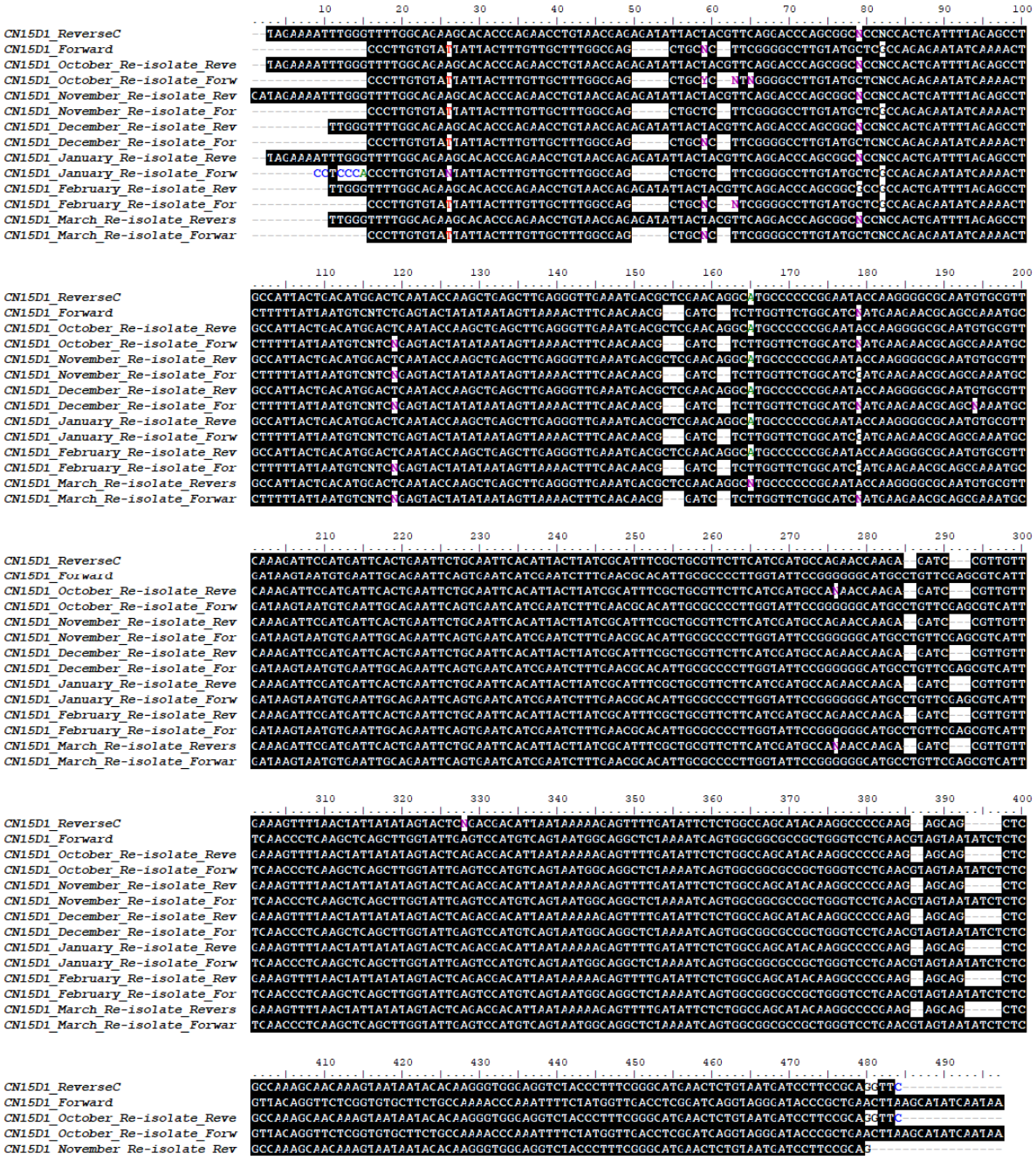

Figure S2. ITS sequences to identify and confirm the pathogen identity of the original CN15D1 isolates and re-isolates from plantings inoculated with mycelia sub- cultured from CN15D1 isolate. The multiple seaquence alignment was generated via ClustalW in BioEdit.