DETECTION OF THE PRESENCE OF QUARANTINE PATHOGENS Xylella fastidiosa IN POTENTIAL HOST PLANTS OF SRI LANKA

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Xylella fastidiosa is a gram negative, xylem-colonizing, vector-transmitted phytopathogenic bacterium that affects around 600 plant species. The European Union has set quarantine regulations for X. fastidiosa and demanded proof of pathogen-free certification before exporting. Though Taiwan is currently the only Asian country with a long-standing presence of X. fastidiosa, there are chances the bacterium to be present to Sri Lanka. Therefore, this research was carried out to determine whether X. fastidiosa was present in possible host plant species in Sri Lanka. In total, 130 composite samples were collected from potential host plant species listed in European Union Regulation 2029/1201 of August 14th 2020, from 09 different districts: Kalutara, Rathnapura, Kurunegala, Kegalle, Jaffna, Galle, Nuwara Eliya, Hambanthota, and Badulla. The specific primers for a conserved genomic region of the polymerase sigma factor of the rpoD gene of 733 bp were used to detect the presence of X. fastidiosa by PCR in the collected samples. The gene obtained from the French collection of Plant Associated Bacteria, CIRM-CFBP was used as the positive control in detecting the PCR products by gel electrophoresis. The results showed that no X. fastidiosa bacterium was present in any of the collected samples.

Keywords: European Union Regulation 2029/1201, Quarantine pathogens, RNA polymerase sigma factor, *rpoD* Gene