





A number of different diagnostic protocols to detect and/or quantify the most important quarantine and emerging pathogens of strawberry in Europe were tested in test performance studies involving the laboratories of the participating countries; a pronounced variation in the percentage of correctly detected samples (56-96%) among the participating labs and between the tests was observed. False negative results could be attributed to a reduced sensitivity due to processes of lyophilisation or vacuum concentration of primers and/or extracted DNA from samples, which were performed to simplify transportation of the material tested. Contamination during the rehydration of samples and/or primers or handling with the PCR mix may have led to false positive results. However, these assumptions would have to be examined in more detail. Under optimal conditions, using freshly extracted DNA and primers, all tests should be suitable to detect the selected diseases directly from diseased strawberry plants.

However, the test performance study pointed out that for implementation of these molecular methods in different laboratory conditions, some optimization is necessary in order to obtain robust diagnostic tests capable to provide reproducible results using different equipment, reagents and laboratory set up.

Further research is needed to get the adequate overview of the pathogens, especially the soil-borne (ie. nematodes) occurrence. RT-PCR is the best suitable detection method to date, but there is a need for a more universal diagnostic approach as several diseases and conditions often form a complex.

Project ID: Assessment and testing of strawberry pathogens (SPAT).