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The samples collected were processed and sequenced through High Throughput Sequencing (VANA strategy) and have contributed to increase the knowledge on the epidemiology of the viruses. The collaboration led to first reports, in several participating countries, not only on findings of little cherry viruses (Tahzima *et al.*, 2017, 2019a), but also on several co-infecting viruses (Tahzima *et al.*, 2019 b, c) or associations with phytoplasmas (Salem *et al.*, 2019). Additionally, a test performance study was organized on the tests for the detection of LChV-1 (Bajet *et al.*, 2008; Glasa *et al.*, 2015) and LChV-2 (Eastwell and Bernardy 2001; Rott and Jelkmann 2001) that are routinely used by the participating laboratories. The results revealed a good analytical specificity and sensitivity, as well as high percentages for repeatability and reproducibility for both tests and for both viruses. The project is currently in its final phase and full results/report will be available after the final meeting in October 2019.

Project ID: Unravelling the Little Cherry disease complex at European scale to improve transnational diagnostics and management of the disease (EURAVELCH)

#### References:

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