



IPSN FACTSHEETS

Xylella fastidiosa



Introduction

The bacterium *Xylella fastidiosa* is considered one of the most threatening plant pathogens of the world, affecting more than 600 plant species including species of high economic value such as olive, coffee, almond, lavender, etc. Pierce's Disease of grapevine, caused by *X. fastidiosa*, was first described in 1892. However, it took almost 80 years for scientists to identify *X. fastidiosa* as the causal pathogen behind outbreaks of the disease.

The main distribution area of *X. fastidiosa* is on the American continent, from Canada to Argentina, but the bacteria is now found outside its endemic area. The presence of the bacteria has been confirmed in Taiwan, Iran, Lebanon and Israel. In 2013, it was found for the first time in Europe, in Italy. Since then outbreaks have been observed in France, Spain and Portugal [see distribution].

There are four known subspecies of *Xylella* and within each subspecies multiple strains exist (also known as sequence types or STs). The symptoms and their severity vary depending on the strain, host and environment, with some plant species showing latent infection. However, even in this latent state, these plants can serve as inoculum sources. *X. fastidiosa* is transmitted by xylem-feeding insects, primarily spittlebugs, sharpshooters, and leafhoppers. Insect vectors transmit the bacteria over short distances (their flight range is around 100 m, but they can travel longer distances aided by the wind), and the main pathway over long distances is the trade of contaminated plants.

Host

Xylella fastidiosa causes serious disease in a wide range of plants, including important crops and ornamental plants [see list of hosts].

High risk plants include: Polygala myrtifolia, Olea europaea (olive), Salvia rosmarinus officinalis (rosemary), Lavandula spp. (lavender), Prunus spp. (plums, cherries, almonds etc.), Nerium oleander (oleander), Coffea spp. (coffee). Other important hosts are: Acer spp. (maple), Cistus spp., Citrus spp., Fraxinus spp. (ash), Hebe spp., Platanus spp. (plane), Quercus spp. (oak), Ulmus spp. (elm), Vaccinium spp. (blueberries), Vitis spp. (grapevine).

Biology

Xylella fastidiosa is a gram-negative, aerobic bacteria, with its optimal growth range between 26-28°C.

Xylella fastidiosa is naturally transmitted from one plant to another with the assistance of vector insects belonging to the Hemiptera order which feeds on xylem. Confined within the xylem of plants, the bacteria spreads along the veins and multiplies inside the plant vessels. As the bacteria invades the xylem vessels, blockages form that obstruct the flow of raw sap. Because the movement of water and nutrients are interrupted, the symptoms often resemble water deficiency or nutrient deprivation.

Symptoms

For details of the symptoms, scan or click on the QR code to access the accompanying poster.

More information

- EPPO Global Database: https://gd.eppo.int/taxon/XYLEFA
- POnTe: https://www.ponteproject.eu/category/symptom-xylella/
- BeXyl project: https://www.bexylproject.org/
- European Commission: https://food.ec.europa.eu/plants/plant-health-and-biosecurity/legislation/control-measures/xylella-fastidiosa
- UK Plant Health: https://planthealthportal.defra.gov.uk/pests-and-diseases/high-profile-pests-and-diseases/xylella/



This factsheet was updated by Itxaso Quintana, BGCI, & Adam Bryning, Fera, May 2024, from a previous version written by IPSN & Fera.

