S1006M07
Integrated Pest Management

S1006

A simulation model to predict the potential spread of the pine wood nematode, Bursaphelenchus xylophilus over a range of scales

<u>Christelle Robinet</u>¹, Jianghua Sun², Richard Baker³, Hugh Evans⁴, Alain Roques⁵

¹INRA, France, ²Chinese Academy of Sciences, China, ³Food and Environment Research Agency, United Kingdom, ⁴Forestry Commission Research Agency, United Kingdom, ⁵INRA, France

The pine wood nematode, Bursaphelenchus xylophylus, is the causal agent of the pine wilt disease. The species is native to North America where it is only a minor pest, mainly on non-native tree species. It has been introduced to several Asian countries (Japan, China, Taiwan, Korea) and to Europe (Portugal with some records in Spain). In most cases, the nematode has spread very rapidly despite control measures, and caused extensive damage to forests. If tree species are susceptible and climate is suitable, then the pine wood nematode can cause pine wilt and trees can die within a few weeks. Within forests and woodlands, the nematode is carried from one tree to another by adult beetles of the genus Monochamus whereas it can spread over long distances as a result of accidental transportation of infested materials. It is therefore important to understand the dispersal mechanism of the nematode and to determine the areas at risk. Here we present a spread model which combines short distance spread (to describe the dispersal of the vector) and long distance spread (to describe human transportation). This model was successfully used to predict the potential range expansion of the pine wood nematode and pine wilt disease in China. It also provides indications of which European ports are the most at risk for nematode introduction arising from international trade. The potential spread in Europe from the present invaded area in Portugal will be determined more accurately in the frame of the EU project REPHRAME (2011-2014).

Keywords: Pine wood nematode, Bursaphelenchus xylophylus, Biological invasion, Monochamus, Spread model, Potential range expansion

All abstracts are subject to approval once submitted with the attendance certification issued by ICE2012