Oregon Sudden Oak Death Management Follow-up: Epidemiology

Hazel Daniels and **Jared LeBoldus**, Department of Botany and Plant Pathology, Oregon State University, Corvallis, OR; hazel.daniels@oregonstate.edu

Phytophthora ramorum, the oomycete pathogen responsible for Sudden Oak Death (SOD) in California and Oregon forests, requires constant vigilance to detect and manage. Prior to 2011, the focus of Oregon-based management was complete eradication of all disease centers through removal of infected plant material. Despite intense eradication procedures, the disease continued to spread. Current Oregon management focuses on reducing the spread of SOD within Curry County, and preventing spread to neighboring counties.

Previous research published in 2015 focused exclusively on the NA1 lineage and its population genomics. The EU1 lineage was first confirmed in a forested site in 2015, in Curry County, OR. Therefore, there's a need to reassess the effectiveness of management techniques with the discovery of this more virulent lineage. Ongoing research looks at the effectiveness of eradication/suppression efforts, including the newly discovered EU1 strain. Additionally, impacts of natural fire on eradication can be studied through the 2017 Chetco Bar Fire, which swept through portions of the Oregon Quarantine Zone and Generally Infested Area.

In addition to assessing management of SOD, a study of the relation between spatial and genetic distance is nearing completion. Using data collected since 2001, both NA1 and EU1 infections were tracked over time and space. A preliminary computational assessment of subsequent new SOD infections relative to previously eradicated infections revealed candidate samples for sequencing. Whole-genome sequencing of spatially related trees and their neighbors will reveal their level of genetic relatedness.

References

Goheen, E.M.; Kanaskie, A.; Navarro, S. and Hansen, E. 2017. Sudden oak death management in Oregon tanoak forests. Forest Phytophthoras. **7:** 45–53.

Kamvar, Z.N.; Larsen, M.M.; Kanaskie, A.M.; Hansen, E.M. and Grünwald, N.J. 2015. Spatial and temporal analysis of populations of the sudden oak death pathogen in Oregon forests. Phytopathology. 105: 982–989.

Grünwald, N.J. et al. 2016. First report of the EU1 clonal lineage of *Phytophthora ramorum* on tanoak in an Oregon forest. Plant Disease. 100: 1024–1024.

Manter, D.K., Kolodny, E.H., Hansen, E.M. and Parke, J.L. 2010. Virulence, sporulation, and elicitin production in three clonal lineages of *Phytophthora ramorum*. Physiological and Molecular Plant Pathology. 74: 317–322.