

Plant Pathology Seminar Series

“Novel plant diseases and their potential threats”

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Abstract

Novel plant-pathogen interactions have the potential to develop into new diseases when the pathogens or plant species are introduced to regions outside their native range (9). Human induced factors like introduction of planting materials or pathogens, changes in the natural landscape, and farming techniques influence the natural equilibrium between host-pathogen interactions and facilitate the emergence of new diseases (1,11). Different groups of plant pathogens (i.e. fungi, bacteria, viruses, and nematodes) respond differently to these factors and may emerge as the causal agents of new diseases (1). Many economically important diseases such as rice blast and potato late blight are thought to result from the interactions of the pathogens on new hosts during different historic time periods (4,11). Many novel plant diseases have become economic problems as the result of the expansion of agriculture and the advent of international trade (7,9). In addition, pathogens have adapted different evolutionary mechanisms to infect new hosts (6,8,10,11). Three different models, host shift, hybridization, and horizontal gene transfer, have been suggested as the evolutionary mechanisms of novel plant diseases (11). Recently, the rice blast pathogen (*Magnaporthe grisea*) was demonstrated to be pathogenic to wheat through host shift in Brazil (4,12,13). Subsequently, the wheat blast pathogen has spread to all wheat growing areas and caused severe yield losses in Brazil (12,13). Utilizing the mechanism of horizontal gene transfer, *Pyrenophora tritici-repentis* that was previously described as a saprophyte or weak pathogen of wheat became more virulent around 1940 and has caused tan spot on wheat. The fungal pathogen acquired a host specific toxin gene, *ToxA*, from *Stagonospora nodorum*, the causal agent of wheat glume blotch (5). Hybridization between *Phytophthora cambivora* and a close taxon of *P. fragariae* resulted in a new species that causes *Phytophthora* disease of alder, a new disease first reported in 1993 in Great Britain (2,3). Agro-ecosystems homogenize the environment, which is favorable not only for more frequent disease epidemics but also for the development of new diseases as compared to natural ecosystems. These new diseases can pose serious threats to food security, biodiversity, and human health and welfare (1,9,11).

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References:

1. Anderson P. K., Cunningham A. A., Patel N. G., Morales F.J., Epstein P. R., and Daszak, P. 2004. Emerging infectious diseases of plants: pathogen pollution, climate change and agrotechnology drivers. *Trend Ecol. Evol.* 19: 535-544.
2. Brasier, C. M., Cooke, D. E. L., and Duncan, J. M. 1999. Origin of a new *Phytophthora* pathogen through interspecific hybridization. *Proc. Natl. Acad. Sci.* 96:5878-5883.
3. Brasier, C. M., Kirk, S. A., Delcan, J., Cooke, D. E., Jung, T., and Man In't Veld, W. A. 2004. *Phytophthora alni* sp. nov. and its variants: designation of emerging heteroploid hybrid pathogens spreading on *Alnus* trees. *Mycol. Res.* 108:1172-1184.
4. Couch, B. C., Fudal, I., Lebrun, M. H., Tharreau, D., Valent, B., Kim, P. V., Notteghem, J. L., and Kohn, L. M. 2005. Origins of host-specific populations of the blast pathogen *Magnaporthe oryzae* in crop domestication with subsequent expansion of pandemic clones on rice and weeds of rice. *Genetics* 170:613-630.
5. Friesen, T. L., Stukenbrock, E. H., Liu, Z., Meinhardt, S., Ling, H., Faris, J. D., Rasmussen, J. B., Solomon, P. S., McDonald, B. A., and Oliver, R. P. 2006. Emergence of a new disease as a result of interspecific virulence gene transfer. *Nature Genet.* 38:953-956.
6. Gabriela Roca, M., Read, N. D., and Wheals, A. E. 2005. Conidial anastomosis tubes in filamentous fungi. *FEMS Microbiol. Lett.* 249:191-198.
7. Matson, P. A., Parton W. J., Power A. G., and Swift, M. J. 1997. Agricultural intensification and ecosystem properties. *Science* 277:504-509.
8. Nikoh, N., and Fukatsu, T. 2000. Interkingdom host jumping underground: phylogenetic analysis of entomoparasitic fungi of the genus *Cordyceps*. *Mol. Biol. Evol.* 17:629-638.
9. Parker I. M., and Gilbert G. S. 2004. The evolutionary ecology of novel plant-pathogen interactions. *Annu. Rev. Ecol. Evol. Syst.* 35:675-700.
10. Rosewich, U. L., and Kistler, H. C. 2000. Role of horizontal gene transfer in the evolution of fungi. *Annu. Rev. Phytopathol.* 38:325-363.
11. Stukenbrock E. H., and McDonald B. A. 2008. The origins of plant pathogens in Agro-ecosystems. *Annu. Rev. Phytopathol.* 46:75-100.
12. Urashima, A. S., Hashimoto, Y., Don, L. D., Kusaba, M., Tosa, Y., Nakayashiki, H., and Mayama, S. 1999. Molecular analysis of the wheat blast [*Pyricularia oryzae*] population in Brazil with a homolog of retrotransposon MGR583. *Ann. Phytopathol. Soc. Jpn.*
13. Urashima, A. S., Igarashi, S., and Kato, H. 1993. Host range, mating type, and fertility of *Pyricularia grisea* from wheat in Brazil. *Plant Dis.* 77:1211-1216.