

Thousand Cankers Disease: New Insights and Future Directions



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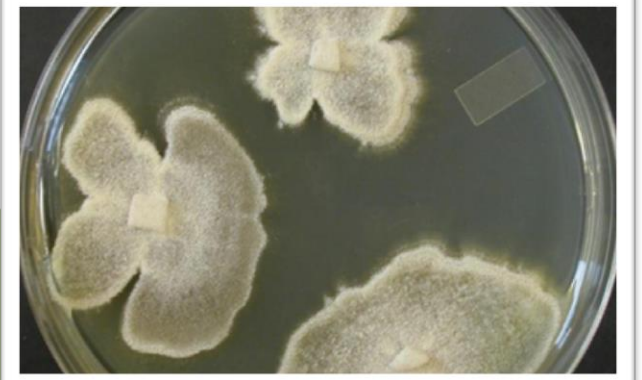
Thousand Cankers Disease



Walnut, Butternut, Wingnut
Juglans and *Pterocarya* spp.

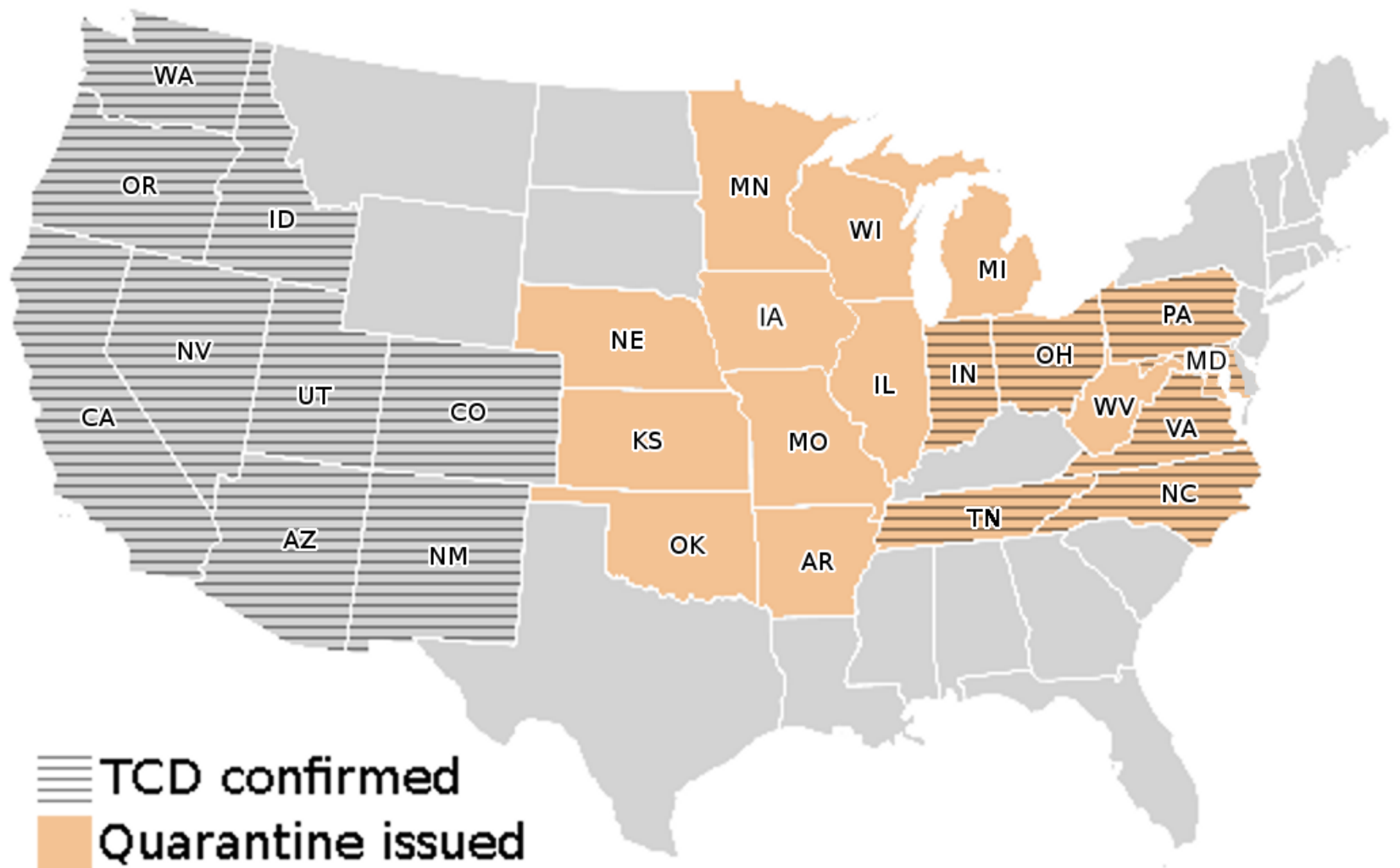


Walnut Twig Beetle (WTB)
Pityophthorus juglandis



Geosmithia morbida

Distribution of Thousand Cankers Disease as of April 20, 2015



Source: www.thousandcankers.com

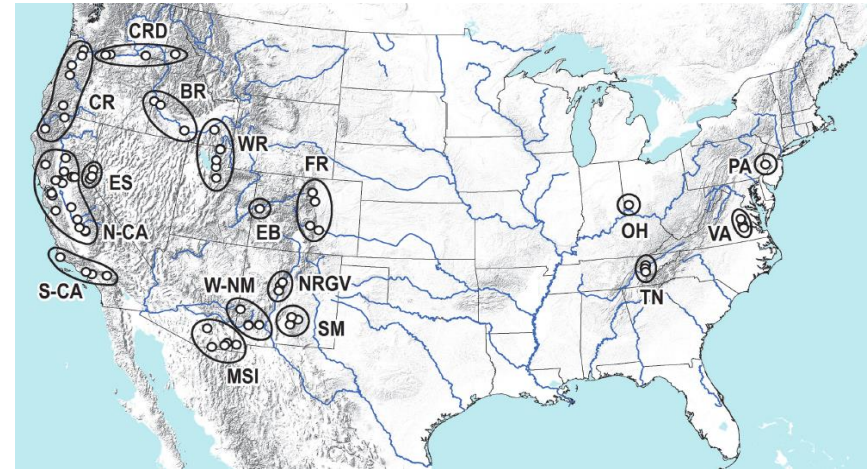
Two new studies on genetic diversity and spread of WTB and *Geosmithia*



Phylogeography of the Walnut Twig Beetle, *Pityophthorus juglandis*, the Vector of Thousand Cankers Disease in North American Walnut Trees

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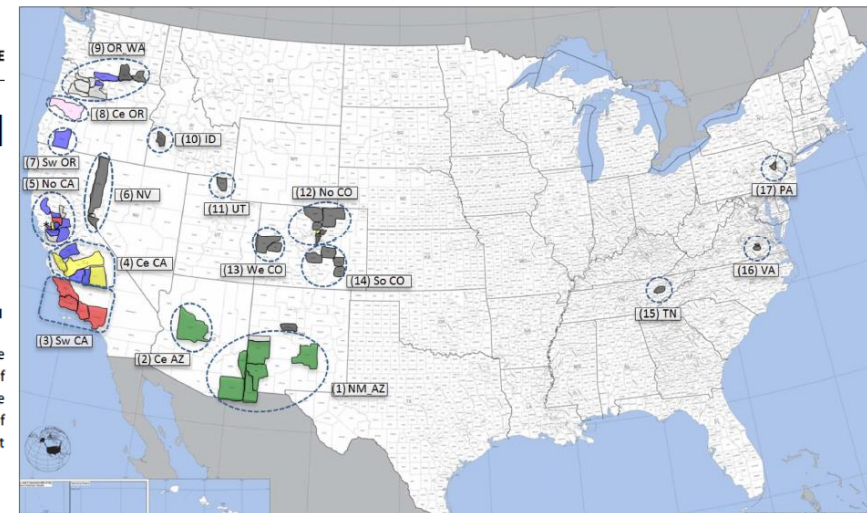
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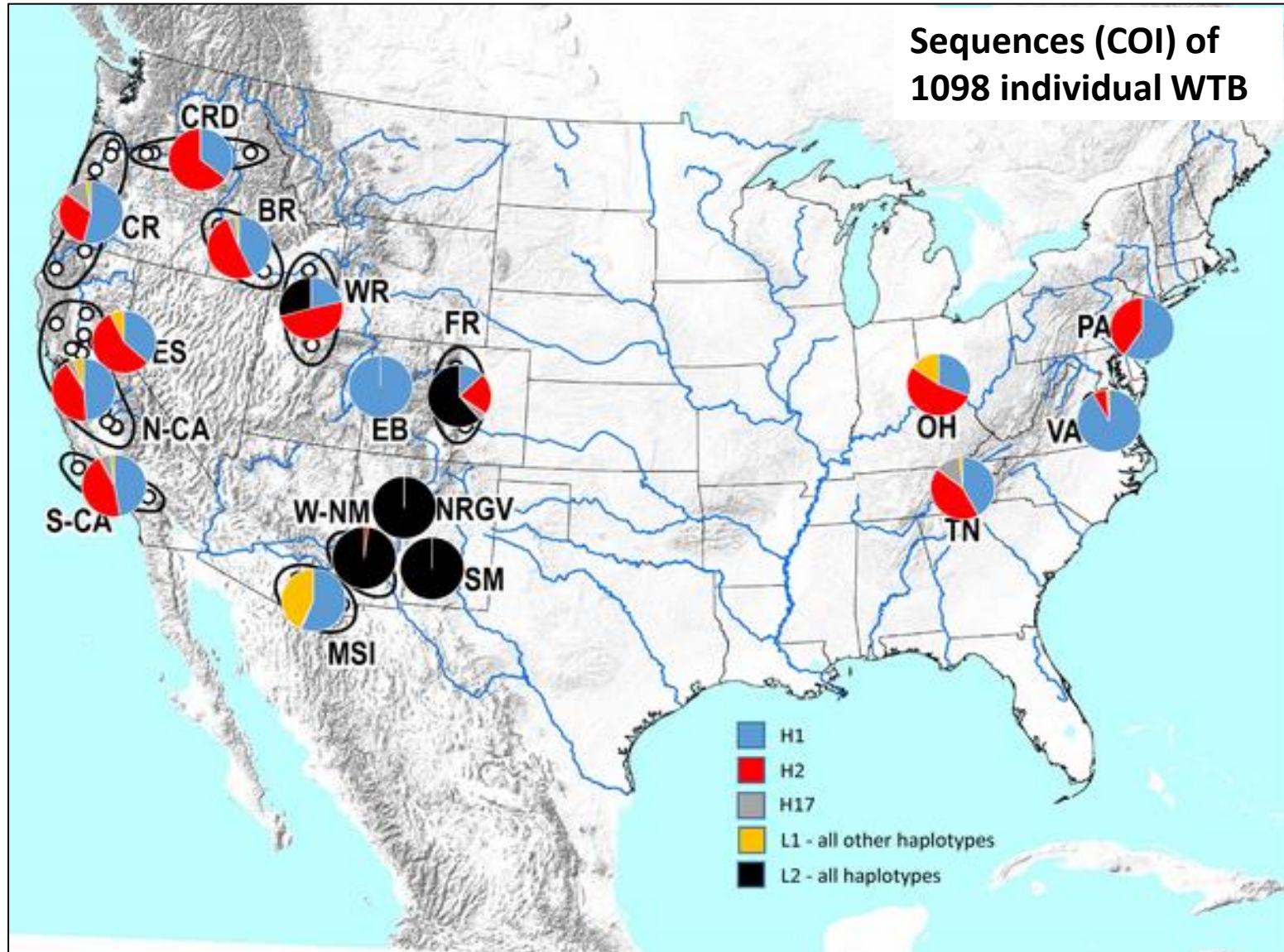
Population Structure of *Geosmithia morbida*, the Causal Agent of Thousand Cankers Disease of Walnut Trees in the United States

Marcelo M. Zerillo^{1,*}, Jorge Ibarra Caballero¹, Keith Woeste², Andrew D. Graves³, Colleen Hartel⁴, Jay W. Pscheidt⁵, Jadelys Tonos⁴, Kirk Broders⁶, Whitney Cranshaw¹, Steven J. Seybold⁷, Ned Tisserat¹

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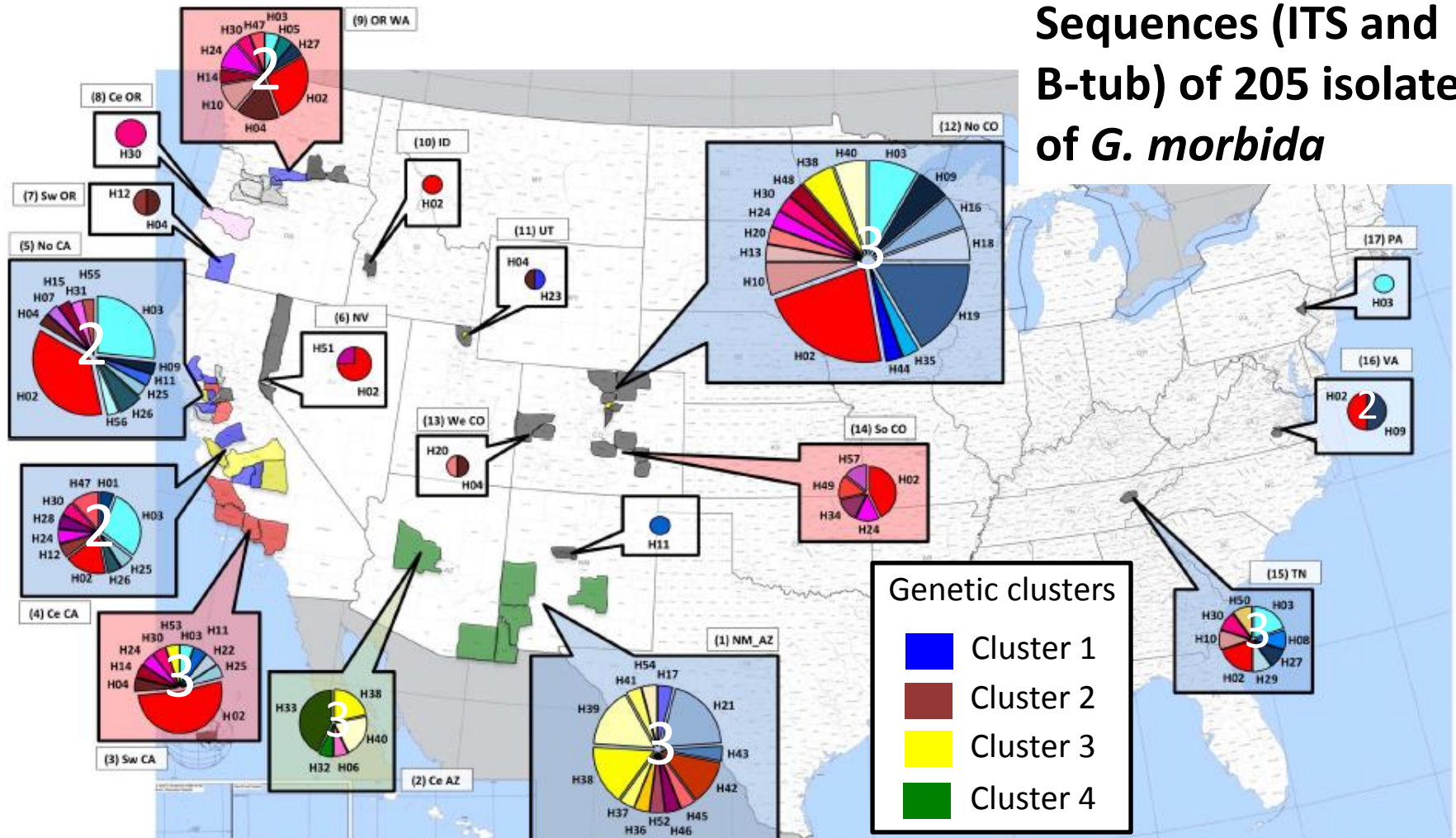


Distribution of 2 lineages of WTB

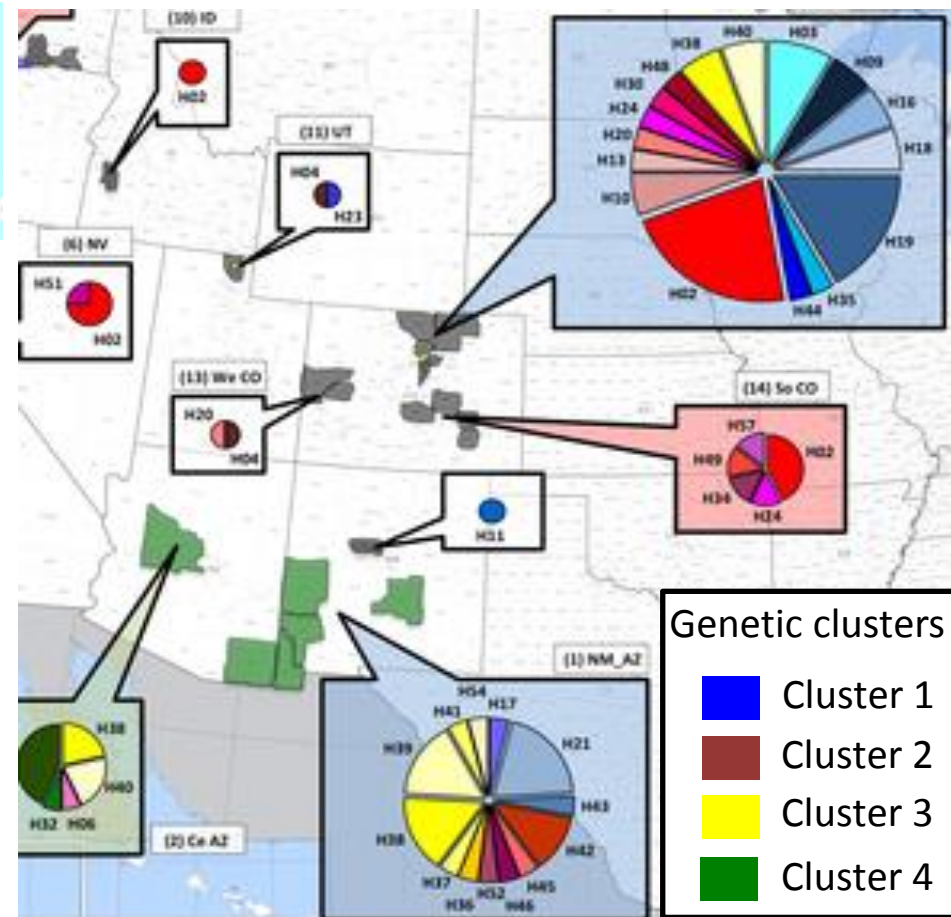
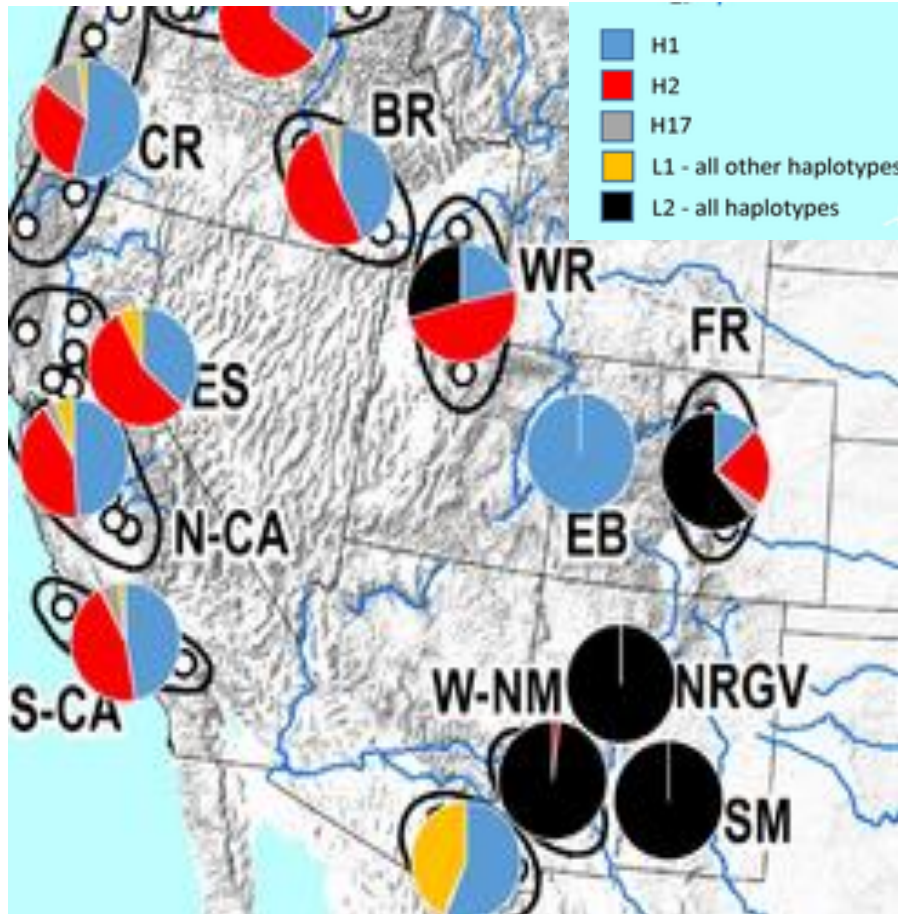


Distribution of 4 lineages of *Geosmithia morbida* across the United States

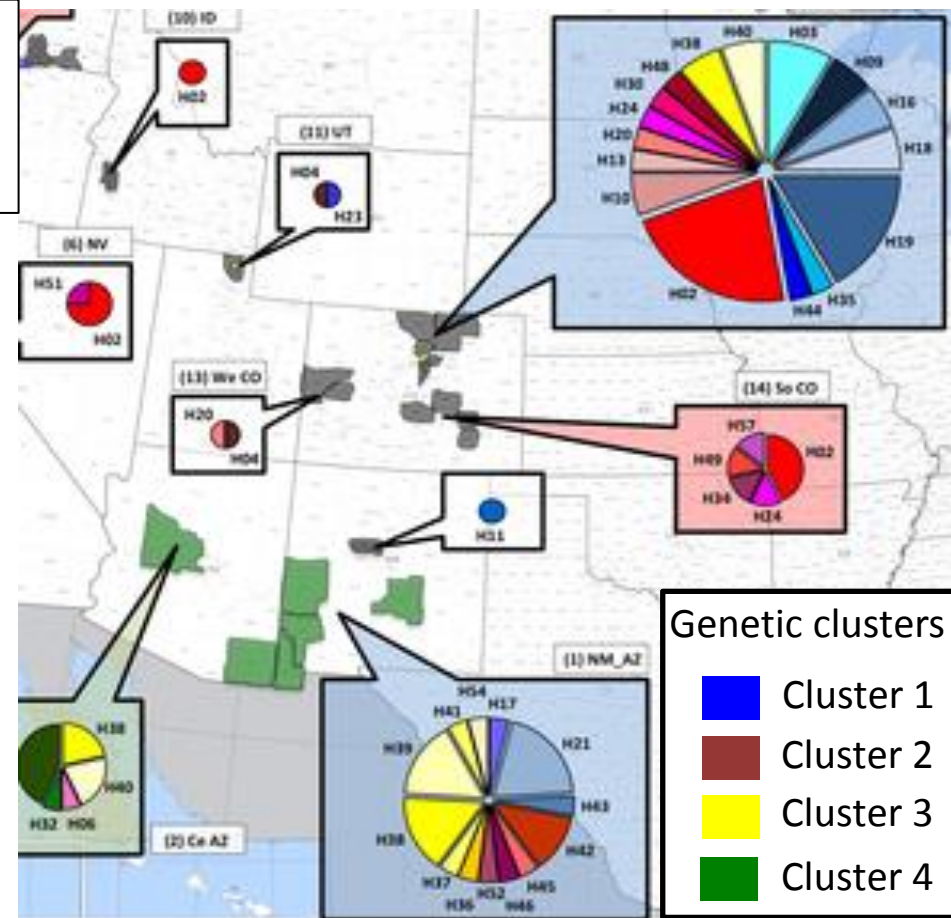
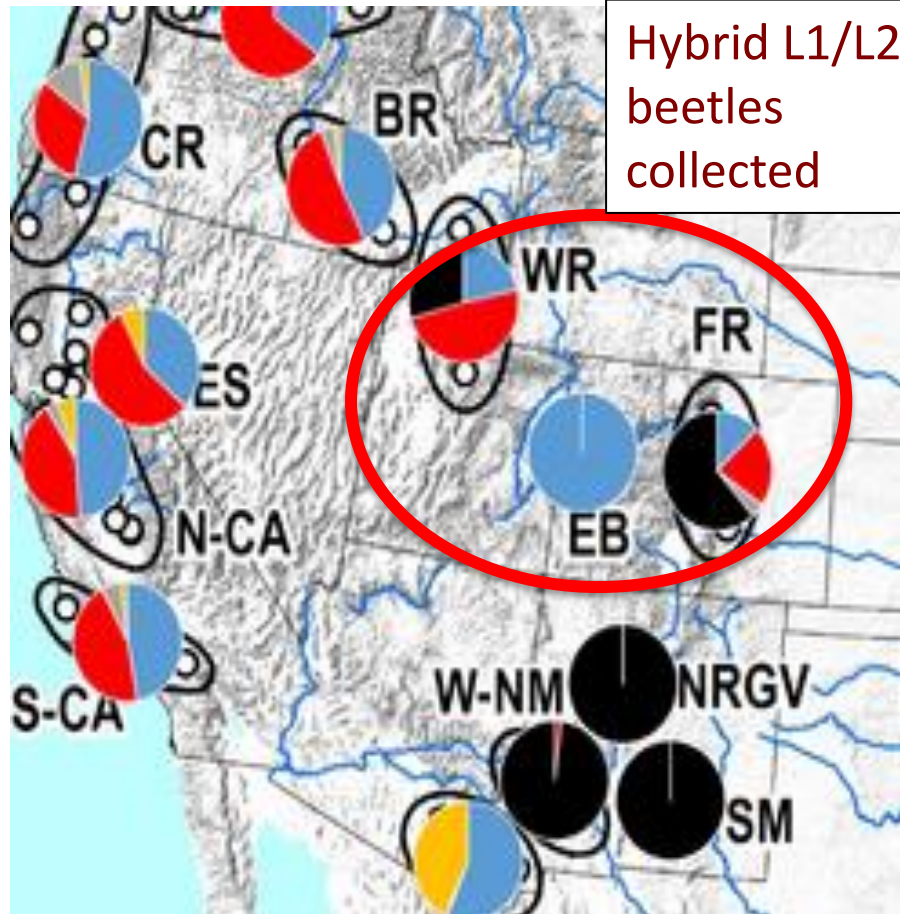
Sequences (ITS and B-tub) of 205 isolates of *G. morbida*



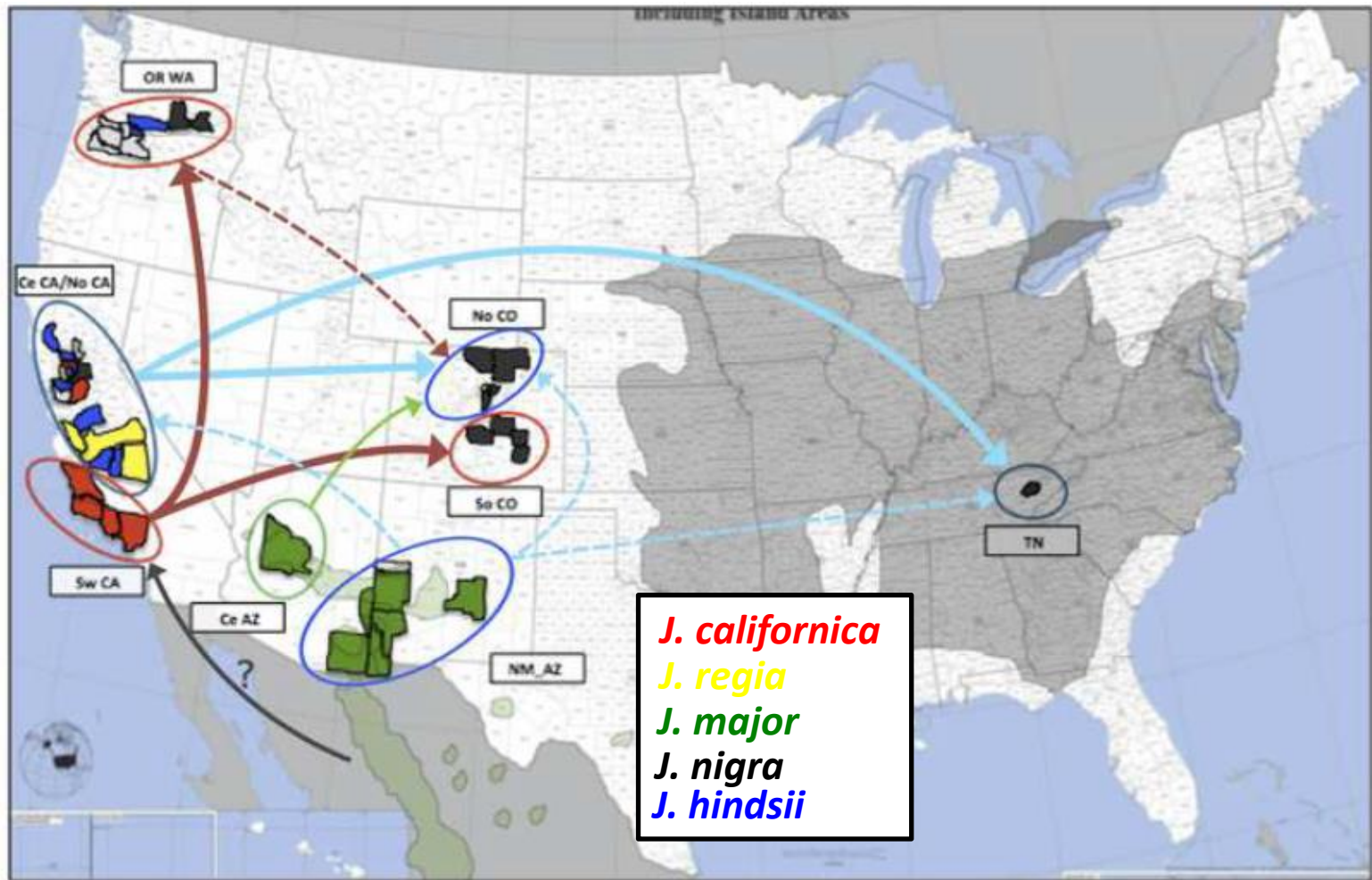
WTB and *G. morbida* most diverse in the SW



Evidence for hybridization of WTB lineages may have spread more aggressive beetle/pathogen complex



Movement and spread of *G. morbida*



Ongoing research at CSU:

- Testing different cultivars of black walnut (different seed sources) for resistance to *G. morbida*
- Testing virulence of *G. morbida* genetic clusters
- Understanding mechanisms of walnut resistance and pathogen virulence by examining RNA expression level
 - *G. morbida* in culture vs. *in planta*
 - Walnut vs. walnut in response to *G. morbida*
- Possibility of hybrids and increased aggressiveness of *G. morbida*
 - Genomic comparisons between *G. morbida* and non-pathogens

