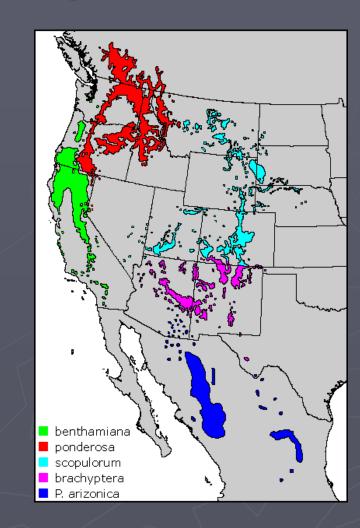
Nesting microsatellite genotype data within low-copy nuclear gene trees to interpret the complex evolutionary relationships of *Pinus* washoensis (*Ponderosae*; Pinaceae)

Brandon Linz (Hendrix College) Ann Willyard (Hendrix College) Aaron Liston (Oregon State University), Richard Cronn (USDA Forest Service PNW)





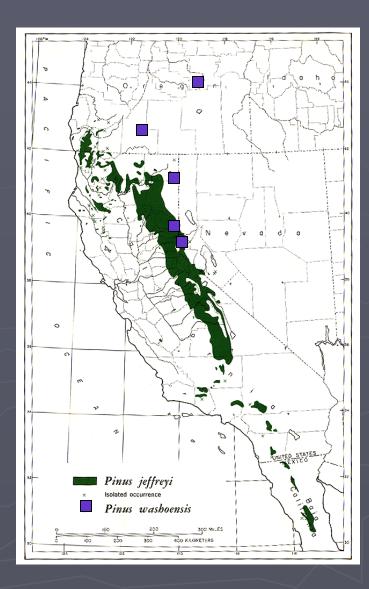
ponderosa





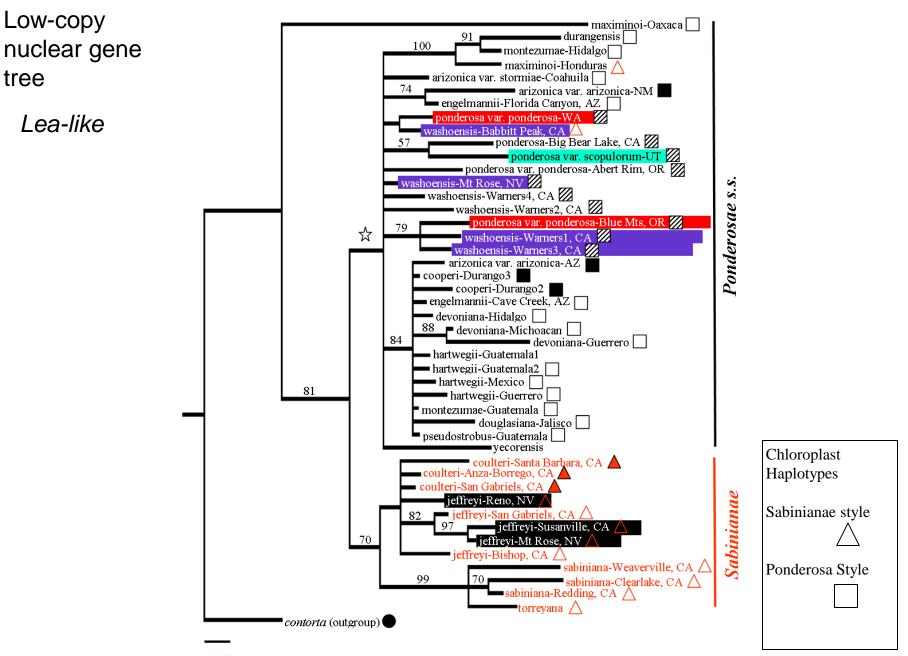
What is Washoe pine?

>Only known from three locations >Higher elevation than Jeffrey >Looks different than nearby Pacific race ponderosas Suggested that outlying populations of North Plateau race

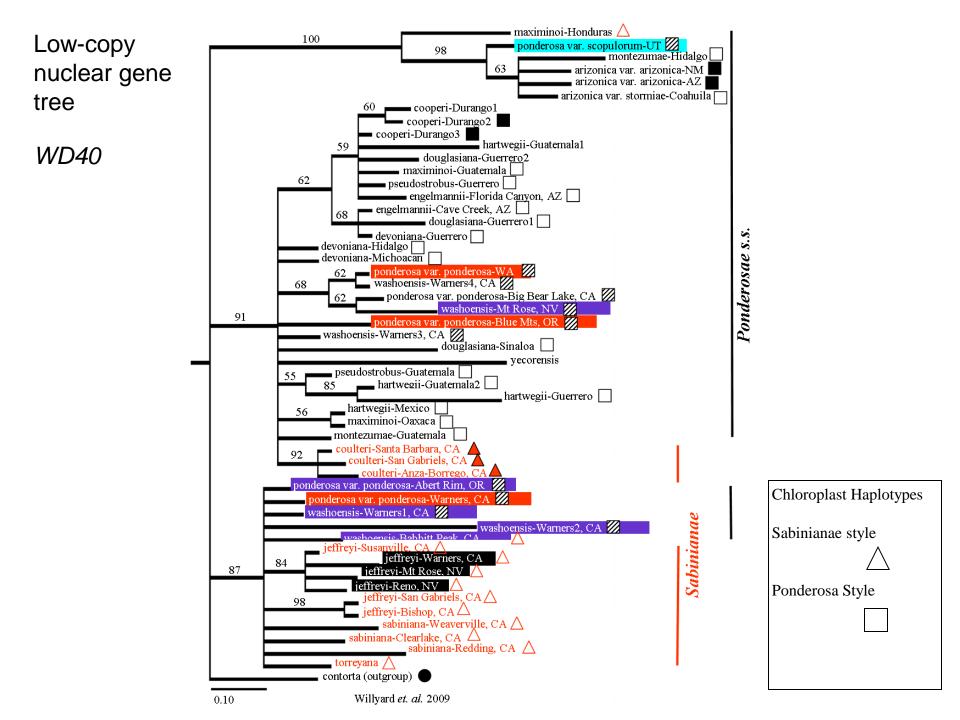


Washoe pines

 High elevation Washoe resemble other high elevation ponderosa and grow better
 Low ponderosa resemble high Washoe pops



Willyard et. al. 2009

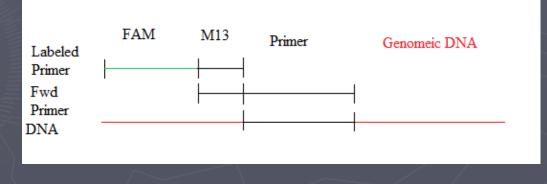


Microsats/SSRs/VNTRs

Repetitive sequences
Non-coding
High mutation rate
Usefull in phylogenetics and systematics

Amplification & Genotyping

PCR Amplify specific region 3 primer system





Hypothesis

Is *P. washoensis* unique or part of the wideranging *P. ponderosa?*



Three Alternatives:
A. *P. washoensis* distinct: high-elevation Oregon population + traditional locations
B. *P. washoensis* distinct but confined to traditional locations
C. *P. washoensis* encompassed within the diversity of ponderosa pine

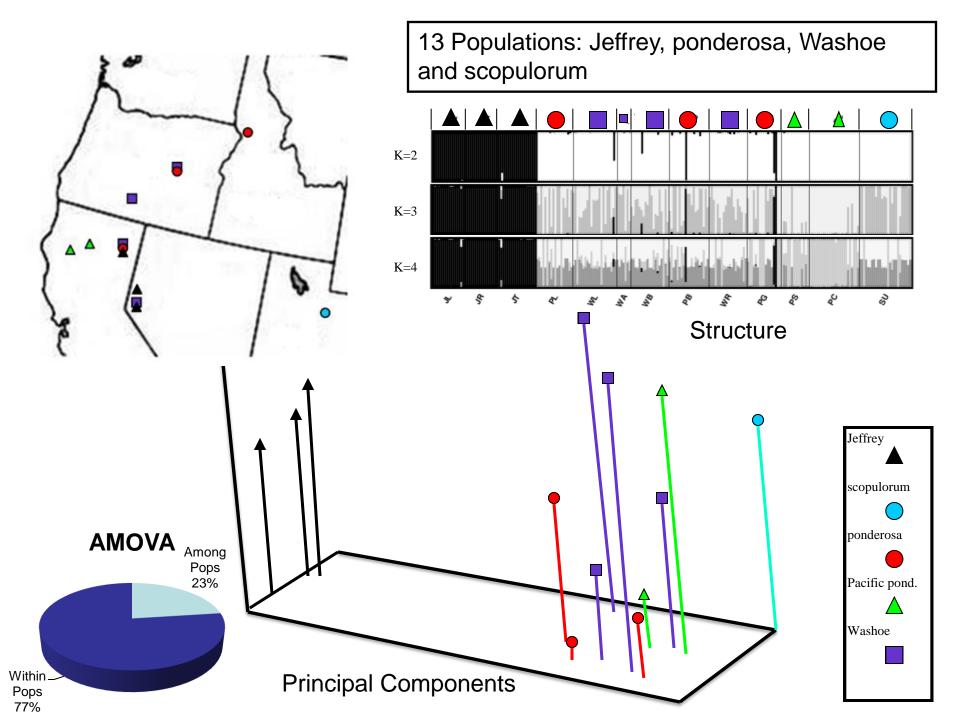
Lab Methods

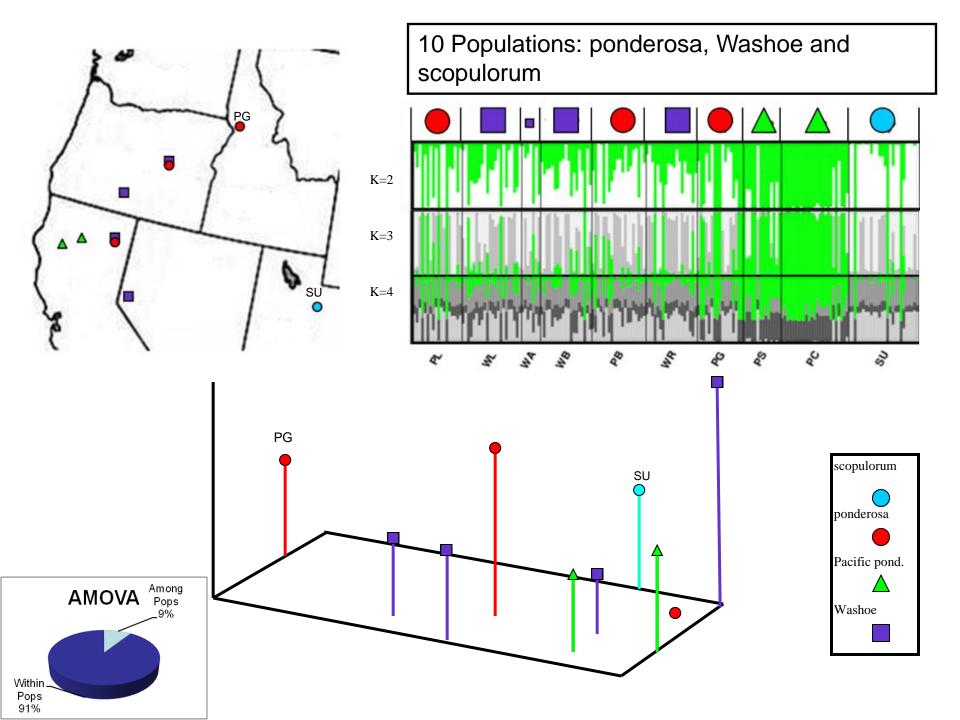
▶ Using 17 nSSRs from *P. taeda* (subg. *Pinus*) Selected 9 single-locus and useable amplifications From 13 populations and 249 individuals (mean=19.2 per population) Amplified 6 polymorphic loci Genotyped on ABI 3130xe No. of Locus Alleles 239 individuals with no missing data LOP5 23 ▶ 70 alleles from 6 loci PtTX2123 **PtTX2128** STRUCTURE (Pritchard) **PtTX3025** Principal Component Analysis 19 PtTX3030 PtTX3098

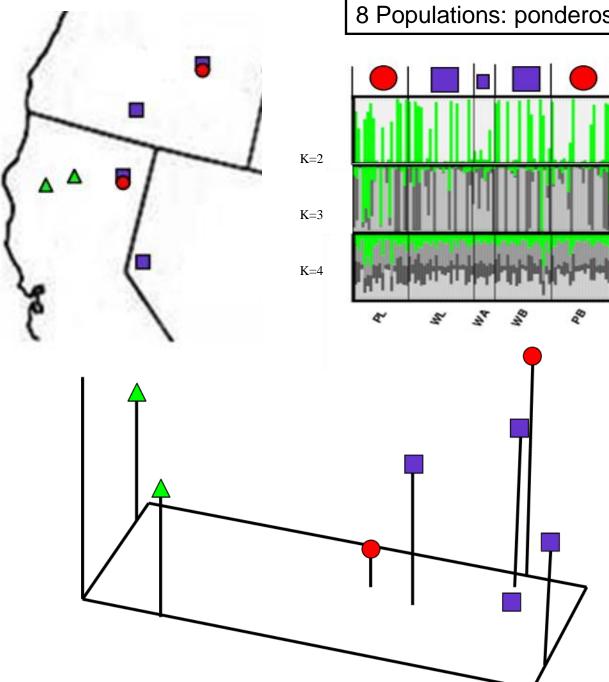
8

70

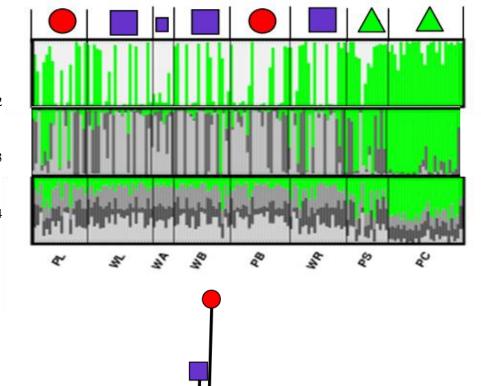
Total

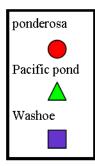


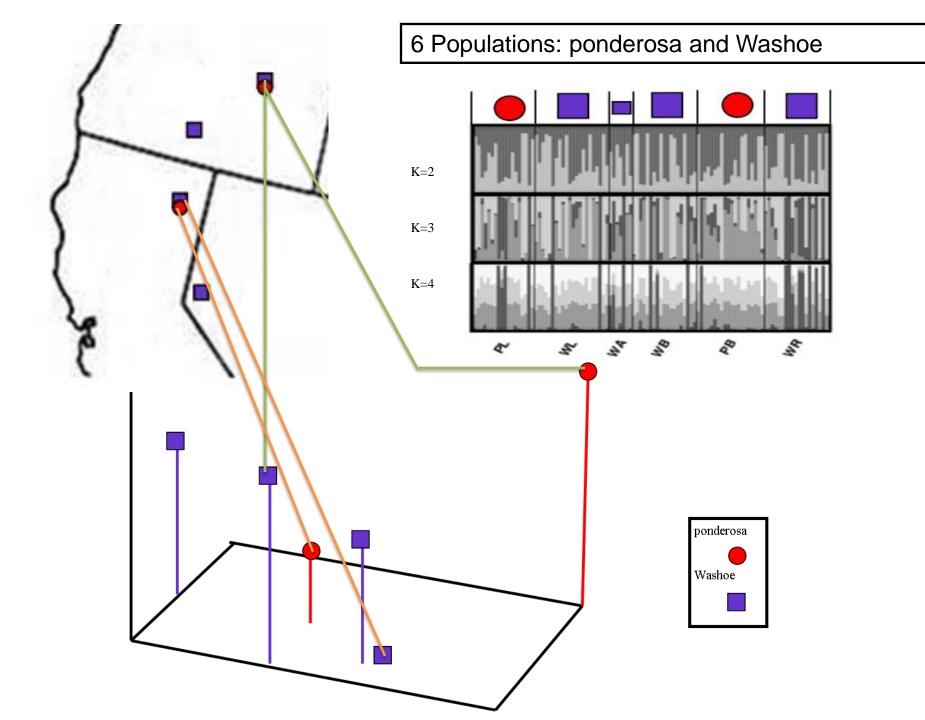




8 Populations: ponderosa and Washoe







Conclusions

Reject hypothesis that *P. washoensis* is distinct from *P. ponderosa* var. *ponderosa* Support that Washoe fits within North Plateau race





Acknowledgements:

David S. Gernandt John R. Haller Kristen Finch, Trang Nguyen, and Nicole Segear Drs. Duina, Murray, Kaushal, & Dearolf

NSF DEB 0317103 to AL & RC USDA Forest Service, Pacific Northwest Research Station Leslie and Vera Gottlieb Research Fund in Plant Evolutionary Biology