

Table S1. Mean allelic diversity metrics of clone-corrected populations of *Phytophthora ramorum* sampled in Curry County, Oregon between 2001-14 causing sudden oak death.

Population	MLG	alleles	1-D	Hexp	Evenness
ChetcoMain	7	3	0.5837	0.9655	0.9432
Coast	12	3.8	0.5944	0.9733	0.9597
HunterCr	4	2.6	0.5562	0.9615	0.9526
JHallCr	30	4.4	0.572	0.9145	0.8908
NFChetHigh	35	4.8	0.5984	0.8961	0.894
PistolRSF	2	2.4	0.55	1	1
Winchuck	9	3.4	0.563	0.9387	0.9165
2001	10	3.6	0.564	0.937	0.9099
2002	9	3	0.5642	0.9443	0.9306
2003	20	4.2	0.5646	0.9025	0.8931
2004	8	2.8	0.5031	0.8441	0.8548
2005	2	2.4	0.5	0.9	0.918
2006	1	2	0.5	1	1
2010	1	2	0.5	1	1
2011	6	3	0.5833	0.9767	0.9638
2012	7	3.2	0.5755	0.9588	0.9498
2013	47	5.4	0.5979	0.8918	0.8987
2014	17	3.8	0.5909	0.9703	0.9347
pooled	70	6	0.6026	0.8931	0.9003
MLG = Number of Multilocus Genotypes					
alleles = Average number of alleles across 5 loci					
1-D = Simpson's Index averaged across 5 loci					
Hexp = Nei's 1978 expected heterozygosity					
Evenness = Evenness averaged across 5 loci					