

| <b>Table S2.</b> Allelic diversity metrics for each locus of clone-corrected <i>Phytophthora ramorum</i> data in Curry County, Oregon between 2001-14 causing sudden oak death. |                |            |             |            |
|---|----------------|------------|-------------|------------|
| <b>locus</b>  | <b>alleles</b> | <b>1-D</b> | <b>Hexp</b> | <b>E.5</b> |
| PrMS6   | 2              | 0.4963     | 0.9927      | 0.9927     |
| Pr9C3   | 2              | 0.495      | 0.99        | 0.9901     |
| PrMS39  | 5              | 0.6229     | 0.7786      | 0.7817     |
| PrMS45  | 3              | 0.5058     | 0.7587      | 0.9533     |
| PrMS43  | 18             | 0.8932     | 0.9457      | 0.7837     |
| mean  | 6              | 0.6026     | 0.8931      | 0.9003     |
| alleles = number of observed alleles  |                |            |             |            |
| 1-D = Simpson's Index   |                |            |             |            |
| Hexp = Nei's 1978 expected heterozygosity   |                |            |             |            |
| E.5 = Evenness  |                |            |             |            |