

**Supplementary Materials Table S2:** Sequence read counts from the RPS10 metabarcoding for the 30 soil samples which were selected for amplification with the nested RPS10 metabarcoding approach.

Sample ID	RPS10 sequence reads				
	<i>Phytophthora</i> sp. ( <i>P. europaea</i> -like)	<i>Phytophthora</i> sp. ( <i>P. europaea</i> -like)	<i>Phytophthora agathidicida</i>	<i>Phytophthora cinnamomi</i>	<i>Phytophthora pseudocryptogea</i>
	ASV_4	ASV_10	ASV_33	ASV_42	ASV_80
5016	0	0	87	479	0
5017	0	6318	0	0	0
5021	99	0	0	0	0
5059	0	0	0	0	0
5068	0	0	0	0	0
5080	0	0	0	0	0
5147	0	182	0	0	0
5159	0	0	0	0	0
5174	0	0	0	0	72
5186	0	0	0	0	0
5232	0	0	0	0	0
5273	590	0	0	0	0
5276	889	0	0	0	0
5305	0	0	0	0	0
5330	0	0	0	0	0
5362	0	0	0	0	0
5382	0	0	0	0	0
5385	0	0	0	4	0
5405	127	0	0	0	0
5414	3922	0	0	0	0
5441	199	0	0	0	0
5448	0	0	0	0	0
5462	3378	0	0	0	0
5499	0	0	0	0	0
5556	0	0	0	0	0
5560	147	0	0	0	0
5577	11	0	0	0	0
5611	1783	0	0	0	0
5678	0	0	1091	0	0
5725	0	0	0	0	0

NB. ASV4 and ASV10 differ by 1 bp