

Supplementary Materials Table S1: Sequence read counts from the ITS metabarcoding for the 30 soil samples which were selected for amplification with the nested RPS10 metabarcoding approach.

Sample ID	ITS sequence reads						
	<i>Phytophthora cinnamomi</i>	<i>Phytophthora agathidicida</i>	<i>Phytophthora multivora</i>	<i>Phytophthora</i> sp (<i>P. europaea</i> -like)	<i>Phytophthora cactorum/ P. aleatoria</i>	<i>Phytophthora pseudocryptogea</i>	<i>Phytophthora kernoviae</i>
	ASV_1	ASV_2	ASV_7	ASV_10	ASV_11	ASV_30	ASV_32
5016	37222	6229	0	0	0	0	0
5017	12732	0	0	502	0	0	0
5021	0	0	0	22062	0	0	0
5059	0	0	0	850	0	0	0
5068	0	0	0	760	0	0	0
5080	0	0	6878	12285	0	0	0
5147	0	0	0	8797	0	0	0
5159	4397	0	0	11057	0	0	0
5174	0	0	0	4037	0	6659	0
5186	0	0	0	0	36025	0	0
5232	0	0	0	205	0	0	0
5273	0	0	0	433	0	0	0
5276	0	0	0	472	0	0	0
5305	7969	1360	0	0	0	2754	0
5330	0	0	0	0	10358	0	0
5362	0	0	0	0	0	0	5381
5382	0	0	268	0	0	0	0
5385	5002	0	0	0	10692	0	0
5405	0	0	0	450	0	0	0
5414	0	0	0	671	0	0	0
5441	0	0	0	409	0	0	0
5448	0	0	0	221	0	0	0
5462	0	0	0	584	0	0	0
5499	14381	0	0	0	1547	0	0
5556	0	0	0	493	0	0	0
5560	99	1093	0	302	0	0	0
5577	0	0	0	252	0	0	0
5611	0	0	0	330	0	0	0
5678	0	0	0	0	0	0	0
5725	0	0	0	0	0	0	12304