

Table S2. Base pairs (bp) differences across *Btub*, ITS, *cox1*, *nadh1* sequences showing inter- and intraspecific variation of *Phytophthora heterospora* (HET), *P. palmivora* (PAL) and *P. taxon palmivora*-like (LIK).

	Length of alignment (bp)	Differences PAL/HET (bp)	Differences PAL/LIK (bp)	Differences HET/LIK (bp)	Differences within PAL (bp)	Differences within HET (bp)
<i>Btub</i>	918	11-16	6-14	7-9	0-16	0-2
ITS	788	3-4	0-1	3	0-1	0
<i>cox1</i>	1346	1-2	2	0	0-2	0
<i>nadh1</i>	796	6	0-6	0	0-6	0
Total	3848	21-28	8-21	10-19	0-25	0-2