

**Table 2** Genes, genomic location, size and percentage of polymorphic sites among mitochondrial and nuclear loci examined from *Phytophthora infestans*

Fragment	Gene	Genomic Location	n	Size of sequenced fragment (bp)	Size of Unambiguously aligned region (bp)	No (%) of polymorphic nucleotide sites		No (%) of polymorphic nucleotide sites	
						Potato	<i>Anarrhichomenum</i>	Potato	<i>Anarrhichomenum</i>
Intron Ras	Non-coding	Nucleus	88	349	224	5 (2.23)	15 (6.69)	0	ND
Ras	<i>PiYpt1</i>	Nucleus	79	600	542	7 (1.29)	-----	0	ND
B-Tubulin	<i>β-tubulin</i>	Nucleus	78	542	490	1 (0.20)	-----	0	ND
P3	<i>rpl14</i> , <i>rp15</i> <i>tRNAs</i>	Mitochondria	66	1308	1180	9 (0.76)	17 (1.44)	4	7
P4	<i>cox 1</i>	Mitochondria	79	964	832	4 (0.48)	12 (1.44)	0	0

n = number of isolates sequenced

Potato isolates include wild *Solanum* species and tomato isolates and section *Anarrhichomenum* includes isolates of the Ic haplotype of *P. andina* from *Solanum tetrapetalum*.

ND = Not determined

Two synonymous substitutions were found (Glutamine and Asparagine) in amino acids for Ras

One synonymous substitution was found (Histidine) in amino acid for B-Tubulin