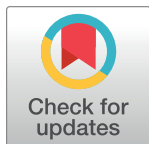


CORRECTION

Correction: Mitochondrial genome sequence of *Phytophthora sansomeana* and comparative analysis of *Phytophthora* mitochondrial genomes

The PLOS Staff

[Fig 3](#) is incorrect. The authors have provided a corrected version here. The publisher apologizes for the error.



OPEN ACCESS

Citation: The PLOS Staff (2020) Correction: Mitochondrial genome sequence of *Phytophthora sansomeana* and comparative analysis of *Phytophthora* mitochondrial genomes. PLoS ONE 15(7): e0236076. <https://doi.org/10.1371/journal.pone.0236076>

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Species	Location	Predicted protein sequence
<i>P. infestans</i>	Ia 10230..10457	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
	Ib 10266..10493	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
	IIa 12177..12404	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
	IIb 12148..12375	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. andina</i>	Ia 10222..10449	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
	Ic 10215..10442	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. ipomoeae</i>	10230..10457	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. mirabilis</i>	10198..10425	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. phaseoli</i>	10248..10475	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. nicotianae</i>	9379..9582	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. sojae</i>	13409..13182	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. polonica</i>	7376..7149	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. ramorum</i>	12818..12591	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF
<i>P. sansomeana</i>	12823..12596	MLLQASKFLGAGLATLGLIGAGIGIGNVFSGSLIIGISRNPQLQELMRTAILGFALTEAIALFCLMMAFLILFAF

Fig 3. Deduced protein sequences of *atp9* gene in the mitochondrial genomes of 10 *Phytophthora* species. Amino acids differ from the consensus are shaded in grey. A deletion at the 3' end of this gene in *P. nicotianae* results in the replacement of the last 13 amino acids with five non-homologous ones (boxed in red).

<https://doi.org/10.1371/journal.pone.0236076.g001>

Reference

1. Cai G, Scofield SR (2020) Mitochondrial genome sequence of *Phytophthora sansomeana* and comparative analysis of *Phytophthora* mitochondrial genomes. PLoS ONE 15(5): e0231296. <https://doi.org/10.1371/journal.pone.0231296> PMID: 32407378