Open Access

https://doi.org/10.48130/sif-0024-0001

Studies in Fungi **2024**, 9: e001

Author Correction: *Alternaria*: update on species limits, evolution, multi-locus phylogeny, and classification

Jun-Fu Li¹, Hong-Bo Jiang^{1,2,3}, Rajesh Jeewon⁴, Sinang Hongsanan^{5,6,7,8}, Darbhe Jarayama Bhat^{9,10}, Song-Ming Tang^{2,3}, Saisamorn Lumyong^{5,6,11}, Peter Edward Mortimer¹, Jian-Chu Xu^{1,12}, Erio Camporesi^{13,14,15}, Timur S. Bulgakov¹⁶, Gao-Juan Zhao¹, Nakarin Suwannarach^{5,6*} and Rungtiwa Phookamsak^{1,5,12*}

- ¹ Honghe Center for Mountain Futures, Kunming Institute of Botany, Chinese Academy of Sciences, Honghe County 654400, Yunnan, P.R. China
- ² Center of Excellence in Fungal Research, Mae Fah Luang University, Chiang Rai 57100, Thailand
- ³ School of Science, Mae Fah Luang University, Chiang Rai 57100, Thailand
- ⁴ Department of Health Sciences, Faculty of Medicine and Health Sciences, University of Mauritius, Reduit, Mauritius
- ⁵ Research Center of Microbial Diversity and Sustainable Utilization, Faculty of Sciences, Chiang Mai University, Chiang Mai 50200, Thailand
- ⁶ Department of Biology, Faculty of Science, Chiang Mai University, Chiang Mai 50200, Thailand
- ⁷ Department of Entomology and Plant Pathology, Faculty of Agriculture, Chiang Mai University, Chiang Mai 50200, Thailand
- ⁸ Guangdong Provincial Key Laboratory for Plant Epigenetics, Shenzhen Key Laboratory of Microbial Genetic Engineering, College of Life Science and Oceanography, Shenzhen University, Shenzhen 518060, P.R. China
- ⁹ No. 128/1-J, Azad Housing Society, Curca, P.O. Goa Velha-403108, India
- ¹⁰ Distinguished Scientist Fellowship Programme, College of Science, King Saud University, Riyadh-11451, Saudi Arabia
- ¹¹ Academy of Science, The Royal Society of Thailand, Bangkok 10300, Thailand
- ¹² CIFOR-ICRAF China Program, World Agroforestry (ICRAF), Kunming 650201, Yunnan, P.R. China
- ¹³ A.M.B, Circolo Micologico 'Giovanni Carini', C.P. 314, Brescia, Italy
- ¹⁴ A.M.B. Gruppo, Micologico Forlivese 'Antonio Cicognani', Via Roma 18, Forlí, Italy
- ¹⁵ Società per gli Studi Naturalistici della Romagna, C.P. 143, Bagnacavallo, RA, Italy
- 16 Department of Plant Protection, Federal Research Centre the Subtropical Scientific Centre of the Russian Academy of Sciences, Sochi 354002, Krasnodar Region, Russia
- * Corresponding authors, E-mail: suwan_461@hotmail.com; jomjam.rp2@gmail.com

Correction to: Studies in Fungi https://doi.org/10.48130/SIF-2023-0001, published online 31 January 2023.

Since the publication of this article, the authors have noticed that the sequence data of primers of LSU, Alt-a1 and GADPH which were listed in Table 2 are wrong. The correct sequence data of primers of LSU, Alt-a1 and GADPH should be:

LROR: GTA CCC GCT GAA CTT AAG C;

LR5: ATC CTG AGG GAA ACT TC;

Alt-for: ATG CAG TTC ACC ACC ATC GC;

Alt-rev: ACG AGG GTG AYG TAG GCG TC;

GPD-1: CAA CGG CTT CGG TCG CAT TG;

GPD-2: GCC AAG CAG TTG GTT GTG C.

Pre-revision Table 2:

Table 2. Gene loci and primers used in this study.

Gene loci	Primers	Sequence 5'-3'	References
Internal transcribed spacer region (ITS, including the 5.85 gene)	ITS5	GGA AGT AAA AGT CGT AAC AAG G	[139]
	ITS4	TCC TCC GCT TAT TGA TAT GC	
28S large subunit rDNA (LSU)	LROR	TCC TGA GGG AAA CTT CG	[140]
	LR5	ACC CGC TGA ACT TAA GC	
18S small subunit rDNA (SSU)	NS1	GTA GTC ATA TGC TTG TCT C	[139]
	NS4	CTT CCG TCA ATT CCT TTA AG	
Alternaria major allergen (<i>Alt-a1</i>)	ALT-F	CAY CCW GGY TTY ATC AAG AA	[49]
	ALT-R	CCN GCD ATN TCR TTR TCC ATR TA	
Glyceraldehyde 3-phosphate Dehydrogenase (<i>GAPDH</i>)	GDP-1	GGT AAC CAA ATC GGT GCT GCT TTC	[141]
	GDP-2	ACC CTC AGT GTA GTG ACC CTT GGC	
Plasma membrane ATPase (<i>ATPase</i>)	ATPDF1	ATC GTC TCC ATG ACC GAG TTC G	[14]
	ATPDR1	TCC GAT GGA GTT CAT GAT AGC C	
The second largest subunit of RNA polymerase II (RPB2)	fRPB2-5f	GAY GAY MGW GAT CAY TTY GG	[142]
	fRPB2-7cR	CCC ATR GCT TGY TTR CCC AT	
Translation elongation factor 1- $lpha$ (TEF1- $lpha$)	EF1-983F	GCY CCY GGH CAY CGT GAY TTY AT	[143]
	EF1-2218R	ATG ACA CCR ACR GCR ACR GTY TG	
	EF1-728F	CATCGAGAAGTTCGAGAAGG	[144]
	EF1-986R	TACTTGAAGGAACCCTTACC	



Post-revision Table 2:

Table 2. Gene loci and primers used in this study.

Gene loci	Primers	Sequence 5′–3′	References
Internal transcribed spacer region (ITS, including the 5.8S gene)	ITS5	GGA AGT AAA AGT CGT AAC AAG G	[139]
	ITS4	TCC TCC GCT TAT TGA TAT GC	
28S large subunit rDNA (LSU)	LROR	GTA CCC GCT GAA CTT AAG C	[140]
	LR5	ATC CTG AGG GAA ACT TC	
18S small subunit rDNA (SSU)	NS1	GTA GTC ATA TGC TTG TCT C	[139]
	NS4	CTT CCG TCA ATT CCT TTA AG	
Alternaria major allergen (<i>Alt-a1</i>)	Alt-for	ATG CAG TTC ACC ACC ATC GC	[49]
	Alt-rev	ACG AGG GTG AYG TAG GCG TC	
Glyceraldehyde 3-phosphate Dehydrogenase (<i>GAPDH</i>)	GDP-1	CAA CGG CTT CGG TCG CAT TG	[141]
	GDP-2	GCC AAG CAG TTG GTT GTG C	
Plasma membrane ATPase (<i>ATPase</i>)	ATPDF1	ATC GTC TCC ATG ACC GAG TTC G	[14]
	ATPDR1	TCC GAT GGA GTT CAT GAT AGC C	
The second largest subunit of RNA polymerase II (RPB2)	fRPB2-5f	GAY GAY MGW GAT CAY TTY GG CCC	[142]
	fRPB2-7cR	ATR GCT TGY TTR CCC AT	
Translation elongation factor 1- $lpha$ (TEF1- $lpha$)	EF1-983F	GCY CCY GGH CAY CGT GAY TTY AT	[143]
	EF1-2218R	ATG ACA CCR ACR GCR ACR GTY TG	
	EF1-728F	CATCGAGAAGTTCGAGAAGG	[144]
	EF1-986R	TACTTGAAGGAACCCTTACC	

The authors would like to apologize for the error.

The original article has been corrected in the HTML and PDF versions.

Published online 8 March 2024 https://doi.org/10.48130/sif-0024-0001



Copyright: © 2024 by the author(s). Published by Maximum Academic Press, Fayetteville, GA. This article is an open access article distributed under Creative Commons Attribution License (CC BY 4.0), visit https://creativecommons.org/licenses/by/4.0/.