



# Two new records of *Alpinia* Roxb. (Zingiberaceae) in Sumatra, Indonesia and phylogenetic relationship to their allied species

Rayfiqa Maulidah<sup>1</sup>, Suci Erta Fitri<sup>1</sup>, Nurainas<sup>2</sup>, Syamsuardi<sup>1</sup>, Dayar Arbain<sup>3</sup>

**1** Department of Biology, Faculty of Mathematics and Natural Science, Andalas University, Padang 25163, West Sumatra. **2** Herbarium Universitas Andalas (ANDA), Department of Biology, Andalas University, Padang 25163, West Sumatra. **3** Laboratory of Sumatran Biota, Andalas University, Padang 25163, West Sumatra.

**Corresponding author:** Nurainas, [nas\\_herb@yahoo.com](mailto:nas_herb@yahoo.com); [nurainas@fmipa.unand.ac.id](mailto:nurainas@fmipa.unand.ac.id)

## Abstract

*Alpinia* is one genus of Zingiberaceae that is distributed throughout the tropical regions. Nine species of *Alpinia* have been recorded in Sumatra. In this article, we report 2 new record species for Sumatra, which are *Alpinia submutica* Roxb. and *A. denticulata* (Ridl.) Holttum. We also registered 3 sequences of them as new data in GenBank under accession number: MH087456, MH087457, and MH218561. Phylogenetic analysis using a molecular approach with their allied species in Sumatra is also presented.

## Key words

Ginger, molecular analysis, new occurrences, Sumatra, taxonomy.

**Academic editor:** Nik Fadzly N Rosely | Received 16 November 2018 | Accepted 3 January 2019 | Published 1 February 2019

**Citation:** Maulidah R, Fitri SE, Nurainas, Syamsuardi, Arbain D (2019) Two new records of *Alpinia* Roxb. (Zingiberaceae) in Sumatra, Indonesia and phylogenetic relationship to their allied species. Check List 15 (1): 109–117. <https://doi.org/10.15560/15.1.109>

## Introduction

*Alpinia* Roxb. is a genus with the highest diversity in Zingiberaceae family that is widespread from Sri Lanka to northern New South Wales (Smith 1990, Larsen et al. 1998). Some species of this genus, like *A. galanga* (L.) Willd., *A. malaccensis* (Burm.f.) Roscoe, and *A. nutans* (L.) Roscoe have been used as medicine, rhizome of *A. galanga* (L.) Willd. is also used as a spice, and the beautiful *A. purpurata* (Vieill.) K.Schum. flowers are known as ornamental flowers (Heyne 1987, Larsen et al. 1999, van Valkenburg and Bunyaphatsara 2001, Nurainas 2007, Namsa et al. 2009).

*Alpinia* is a native genus of Southeast Asia and related information about the species has been reported in some publications. Twenty-five *Alpinia* species are noted from the Malay Peninsula region. Seven species

are distributed throughout Java. Borneo region has 9 species, out of which, 6 endemic, Sarawak, which is the closest area to Sumatra, has 8 species. The latest report for Sumatra has five species: *A. aquatica* (Retz.) Roscoe, *A. capitellata* Jack, *A. galanga* (L.) Willd., *A. mutica* Roxb., *A. sumatrana* (Miq.) K.Schum. (Newman et al. 2004, Poulsen 2006, Lamb et al. 2013).

Exploration of the Sumatran Zingiberaceae including *Alpinia* has been carried out quite thoroughly since 2006. In that period, several new discoveries have been reported, such as the new species, namely, *Zingiber engganoensis* Ardiyani, *Zingiber alba* Nurainas, and *Scaphoclamys perakensis* Holtt. which is a new record for Sumatra (Ardiyani 2016, Nurainas and Arbain 2017). Information related to the distribution or morphological description of *Alpinia* in Sumatra has not been recorded properly, and there is a big chance for new findings.

Based on a survey conducted in Sijunjung and Harau, West Sumatra, several populations of *Alpinia* have been found. They have morphological similarities with *A. submutica* which was last found on Java and *A. denticulata* which was reported previously in Perak, Terengganu, Peninsular Malaysia. The objective of this study is to report the distribution and morphological description as well as the phylogenetic relationship of 2 new records of *Alpinia* in Sumatra with their allied species.

## Methods

**Taxonomy work.** The surveys and exploration were carried out in the field from December 2016 until September 2017 followed by samples collection at various sites in Sumatra, which referred to Herbarium ANDA specimen information. Sample collection and preservation techniques referred to Smith (1981) and character note referred to Smith (1990). The description follows de Vogel (1987), where the description starts from general to specific characters. Collected specimens were deposited as voucher specimens at ANDA Herbarium of Andalas University.

Identification process, information related to taxonomy and geographic distribution to confirm species identities were obtained from the related literature (Schumann 1904b, Holttum 1950, Newman et al. 2004). Additional specimen information based on Herbarium Bogoriense (BO) Cibinong, virtual online database Plantlist (<http://www.theplantlist.org/>), Global Biodiversity Information Facility (<https://www.gbif.org/>) and Kew's Herbarium (<http://www.kew.org/herbarcat>).

**Molecular work.** The genomic DNAs were extracted from fresh or silica dried tissue using some modification of Doyle and Doyle (1987) CTAB (hexadecyltrimethylammonium bromide) method. The amplification

of Internal Transcribed Spacer (ITS) was accomplished using primer pair namely ITS4–ITS5 and the *trnH-psbA* intergenic spacer was amplified with *trnHf\_05-psbA3\_f* (White et al. 1990, Kress et al. 2005b). We used MyTaq™ Red Mix for amplification according to the manufacturer's directions and following the same conditions as described in (Kress et al. 2005a, Kress and Erickson 2007), with annealing temperatures of 54–58 °C. Consensus sequences were aligned using Multiple Sequence Alignment tool Clustal X v. 2.0 (Thompson et al. 1997) and concatenation was done using Geneious Prime 2019.0.4 (<http://www.geneious.com>, Kearse et al. 2012).

Maximum Parsimony (MP) and Maximum Likelihood (ML) analyses of the ITS and *trnH-psbA* sequence data were carried out using MEGA 7 with Tamura 3-Parameter model (Kumar et al. 2016), with equally weighted characters and 1000 random-sequence-additional-replicates. A total of 13 ITS and *trnH-psbA* sequences from GenBank were used as ingroups and 3 sequences of *Curcuma longa* L., *Hedychium coccineum* Buch.-Ham. ex Sm., *Zingiber mioga* (Thunb.) Roscoe were chosen as an outgroup. Bootstrap support was categorized according to criteria by Kress et al. (2002), i.e., strong (> 85%), moderate (70–85%), weak (50–70%), or poor (< 50%) support.

## Results

***Alpinia submutica* K.Schum.** *Alpinia submutica* K. Schum., Bot. Jahrb. Syst. 27 (1899) 280. — Type: J.E. Teijsmann 2040 (holo BO!).

Figure 1

**New record.** Indonesia. Sumatra: West Sumatra, Payakumbuh, Lima Puluh Kota regency, Harau village, Batang Harau riverside near Aka Barayun waterfall,

**Table 1.** Sources, and voucher specimens of ingroup and outgroup were obtained from GenBank.

No.	Species	GenBank accession number		Voucher	
		ITS	trnH-psbA	ITS	trnH-psbA
Ingroup					
1	<i>Alpinia aquatica</i>	KJ507884	JN043815	N9	—
2	<i>Alpinia conchigera</i>	AF478712	JN043826	Kress #00-6706 US	ZL305-3
3	<i>Alpinia pusilla</i>	KT280465	KT280463	WI 97	WI 97
4	<i>Alpinia macrocrista</i>	KT280464	KT280462	WI 96	WI 96
5	<i>Alpinia officinarum</i>	AF478718	GU180428	Kress #00-6614 US	PS0519MT04
6	<i>Alpinia zerumbet</i>	KX018020	JN043874	—	ZL355-1
7	<i>Alpinia japonica</i>	AF254474	KF694867	LM	OUT2
8	<i>Alpinia mutica</i>	KJ507924	—	N84	—
9	<i>Alpinia blepharocalyx</i>	AF478709	JN043817	Kress #98-6136 US	ZL302-3
10	<i>Alpinia foxworthyi</i>	AF478714	JN043836	Kress #98-6293 US	Kress #94-5539
11	<i>Alpinia oxyphylla</i>	AY742372	JN043863	—	—
12	<i>Alpinia polyantha</i>	AY745692	JN043873	Kress #94-3744 US	—
13	<i>Alpinia calcarata</i>	KJ871912	JN043820	86558	—
Outgroup					
1	<i>Hedychium coccineum</i>	AY424758	KC597934	—	ZMN06
2	<i>Curcuma longa</i>	JQ409956	FJ687416	JLS 73222	—
3	<i>Zingiber mioga</i>	KJ025068	GQ435047	TKM-1-000049	PS0528MT01





**Figure 1.** *Alpinia submutica* K.Schum. (S.E. Fitri et al. 04-Sc AL HR, ANDA). **A.** Habit. **B.** Inflorescence. **C.** Flower. **D.** Ligule. **E.** Leaf. **F.** Fruits. **G.** Dissected perianth (from left): flower, calyx, corolla lobes, labellum, stamen, pistil, fruit. Scale bars: B = 2 cm; C = 2 cm; F = 2 cm; G = 1 cm. Photographed by S.E. Fitri.

(00°06'21.82" S, 099°88'56.4" E), 500 m a.s.l., 23-IV-2016, fl., S.E. Fitri et al. 04-Sc AL (ANDA); Sijunjung regency, Muaro Sijunjung village, Kulampi customary forest, (00°39' S, 100°58' E), 150 m a.s.l., 2-IV-2016, fl.&fr., S.E. Fitri et al. 07 Al SC SJJ (ANDA).

**Description.** Herbs 1–2.5 m tall, rhizome subsurface, ca 2.1 cm in diameter, cream inside, fleshy, aromatic. Pseudostem 2.0–3.0 cm in diameter, erect, thick. Leaves 16–21 in 1 plant, petioles 0.5–4 cm, glabrous, ligule 0.5–0.10 × 0.5–1.0 mm long, green with red pattern; laminae 14.0–34 × 4–6.0 cm long, lanceolate, green, with hairy lower side and midrib, upper side glabrous, apex caudate, base unequal, margins entire. Inflorescences terminal, erect or drooping, rachis 10.5–16.0 cm, 6–25 cincinni,

primary bract absent. Flower ca 5.5 cm long, 2–4 flowers open at a time; bracteole absent; calyx white, obovate; corolla tube shorter than calyx; dorsal lobe white, ovate, apex rounded, 0.1 × 2.5 cm, glabrous; lateral lobe white, like ship, 0.5 × 3.0 cm; labellum obovate, 2.5 × 3.5 cm, basally concave with red spots and lines, apex bilobed, yellow, with wrinkled margin throughout; stamen ca 3.5 cm long; anther 0.5 × 1.0 mm, cream, not crested; style glabrous, white, 3–4.0 cm long; ovary green, ca 1 cm long, hairy; epigynous glands ca 4 mm long. Fruit 2.0–3.0 cm in diameter, tricoccous, round, red when ripe, wavy surface, calyx persistent, 7–21 fruit in one stem; 35–36 seeds in 1 fruit, seed black with aril.

**Distribution.** Java and Sumatra

**Ecology.** *Alpinia submutica* is found in riverside areas of secondary forest.

**Additional Information.** Previously, *Alpinia submutica* was identified as *A. mutica* Roxb. but differs in its fruit surface. *Alpinia submutica* has a wavy surface, while *A. mutica* has a flat surface. This result was confirmed based on specimen of Herbarium Bogoriense *Alpinia submutica* K.Schum. Bot. Jahrb. Syst. 27 (1899) 280 [Type: J.E. Teijsmann 2040 (holo BO!)] and Archip. Ind. Sumatra Sidjoendjoeng. There was no sequence data for this species in GenBank, and we registered it for the first time on 20 March 2018 under accession number MH087457.

***Alpinia denticulata* (Ridl.) Holttum**, *Alpinia denticulata* (Ridl.) Holttum, Gard. Bull. Singapore 13 (1950) 143. — Basionym: *Hedygium denticulatum* Ridl., J. Straits Branch Roy. Asiat. Soc. 32 (1899) 102. — *Odontochium denticulatum* (Ridl.) K. Schum., Pflanzenr. IV, 46 (1904) 60. — Lectotype: H.N. Ridley 9455 (SING!), designated by Turner, 2000), Perak.

Figure 2

**New record.** Indonesia. Sumatra: West Sumatra, Payakumbuh, Lima Puluh Kota regency, Harau village, (00°06'19.55" E, 100°40'23.88" S), ca 720 m a.s.l., 23-IV-2016, fl.&fr., S.E. Fitri et al. 01-Sc Al HR (ANDA).

**Description.** Herbs 1–2 m tall, rhizome subsurface, 1–1.5 cm in diameter, cream inside, fleshy, aromatic. Pseudostem 1–3 cm in diameter, erect, thick. Leaves 11–13 in 1 plant, petioles 0.5–1.5 cm, green, ligule 0.5–1.0 × 0.1–0.5 cm long, green, hairy; laminae 7.5–32 × 1–5.5 cm long, lanceolate, apex caudate, base unequal, entire margins, green, hairy. Inflorescences terminal, erect, rachis 13.0–14.0 cm, 26–35 cincinni (1–3 flowers per cincinni), 2 primary bracts, 3.5–6 cm and 8–12 cm long. Flower ca 5.0 cm long, 1–6 flowers open at a time; bracteole obovate, brown; calyx tubular, tube 1.0 × 1.5 cm long, apex acute, brown, stiff; corolla: dorsal lobe acute, ca 1.5 cm long, red, apex acute, thick, glabrous; lateral lobe 1.0 × 1.5 cm, brownish-cream, apex rounded, thick, glabrous; labellum ca 2.6 cm long, apex trilobed, from base to middle red, thick, glabrous, middle to top light green; stamen ca 4 cm; staminodes ca 9 mm long, like long horn; anther 8 mm long, reddish-cream, not crested; style ca 2.6 cm long, white; ovary ca 10 mm long, light green, glabrous; epigynous glands ca 2 mm long. Fruits round, trilocular, green when unripe, hairy, ca 6 in 1 stem.

**Distribution.** Perak, Terengganu and Harau Valley, West Sumatra.

**Ecology.** Found in limestone hills at ca 720 m a.s.l. in the Harau Valley. Vernacular name “Langkueh Rimbo”.

**Additional Information.** *Alpinia denticulata* was collected in Lumut, Perak in the period 1896–1898 by Ridley and collected in Bauk Hill, Terengganu by Lewis

in 1977 (Holttum 1950, Royal 2018). There was no sequence data for this species in GenBank, and we registered it for the first time on 20 March 2018 under the accession numbers MH087456 and MH218561.

Twenty-three sequences of ITS and *trnH-psbA* were obtained from GenBank and taken as ingroup and outgroup in this study. *Alpinia submutica* from *trnH-psbA* region is not included in this analysis because of the poor sequencing result. The phylogenetic tree building methods Maximum Parsimony (MP) and Maximum Likelihood (ML) produced similar ITS and *trnH-psbA* tree topologies (Figs 3, 4). Based on the phylogenetic analysis, it was seen that the Internal Transcribed Spacer had a total aligned length of 572 bp with the mean GC content of 55.67%, meanwhile, *trnH-psbA* had a total aligned length of 509 bp with 29.87% mean GC content. Combined alignments of ITS and *trnH-psbA* produce a total aligned length of 1081 bp with mean GC content of 44.13%.

## Discussion

Seven species of *Alpinia* species that have already been reported are spread across several regions in Sumatra. These results increase the number of species reported from Sumatra to 11 species, with 2 new records, namely *A. submutica* (Fig. 1) and *A. denticulata* (Fig. 2). They were not initially recorded in Sumatra region, where *A. submutica* was reported to be distributed in Java (Newman et al. 2004), while *A. denticulata* was last found in Perak and Terengganu-Malaysia (Holttum 1950).

In 2009, *A. submutica* (Fig. 1) was collected in Harau, Lima Puluh Kota District, West Sumatra, and the specimens were stored in ANDA Herbarium, and identified as *A. mutica* (Nelvita 2009). The difference in fruit surface of these 2 species caused doubts about the results of the identification. After tracing and recollecting this species in the same area, it was re-identified and the similarities were found with that of the specimen of Herbarium Bogoriense *Alpinia submutica* K.Schum. Bot. Syst. 27 (1899) 280. — Type: J.E. Teijsmann 2040 (holo! BO!) And Archip Ind Sumatra Sidjoendjoeng, with the same wavy fruit surface and this species can be confirmed as *A. submutica* (Fig. 1). In this study, *A. submutica* was also found in Kulampi Customary Forest, Muaro Sijunjung, Sijunjung District, West Sumatra.

*Alpinia denticulata* was collected in Lumut, Perak in the period 1896–1898 by Ridley and collected in Bauk Hill, Terengganu by Lewis in 1977 (Holttum 1950, Royal Botanic Garden Edinburgh 2018). Initially, this species was identified as *Hedygium* because of its narrowed ligule and long staminodes. These characters did not have similarities with *Alpinia* in general. Schumann (1904b) was doubtful of that result because of the similarity of characters of inflorescences and fruit with that of *Dieramalpinia* section of *Alpinia* prepared by him.





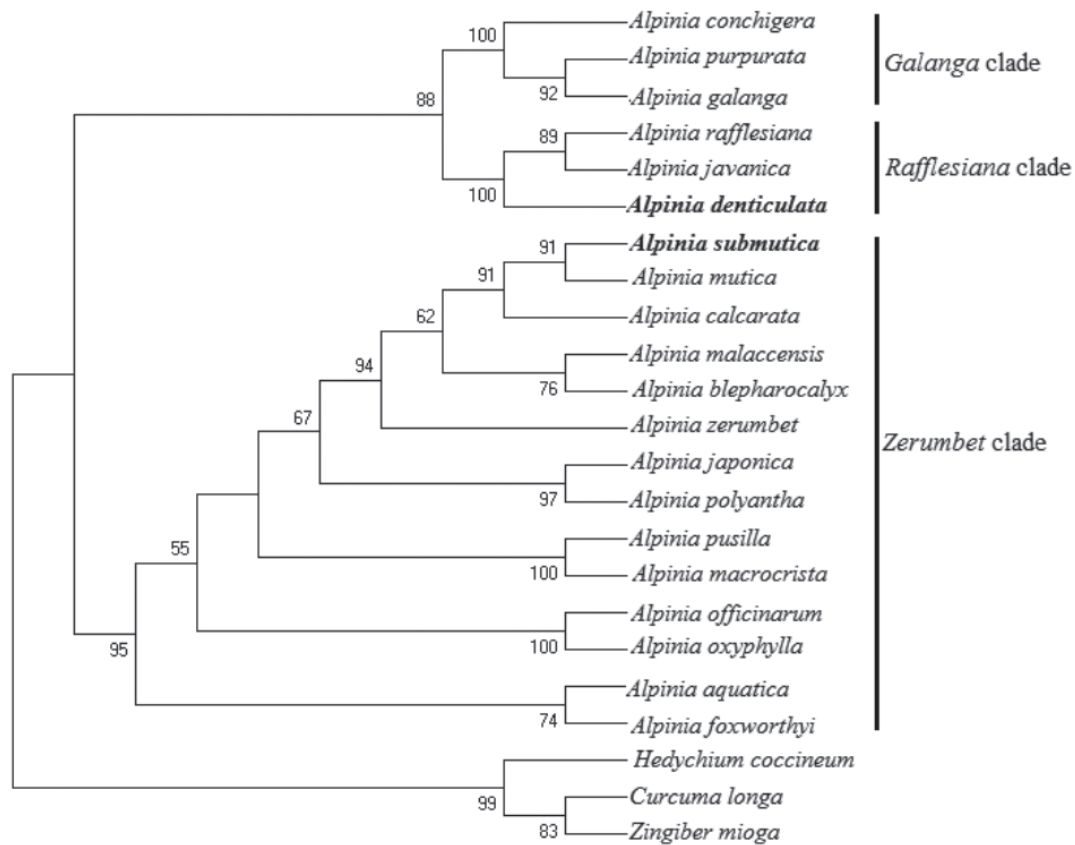
**Figure 2.** *Alpinia denticulata* (Ridl.) Holttum (S.E. Fitri et al. 01–SC Al HR, ANDA). **A.** Habit. **B.** Inflorescence. **C.** Ligule. **D.** Young leaf. **E.** Flower. **F.** Fruits. **G.** Dissection of flower (from left): flower, calyx, bract, corolla lobes, labellum, stamen, pistilum. Scale bars: B = 2 cm; C = 2 mm; F = 2 cm; G = 1 cm. Photographed by S.E. Fitri.

Staminodes of this species are not much different from some species of *Alpinia*, only the curved shape of ligule makes a striking distinguishing character compared to other *Alpinia*. These results were referred to Kew Herbarium specimen (<http://specimens.kew.org/herbarium/K000292387>) and illustration of *A. denticulata* (Schumann, 1904a). The existence of these 2 species in new locations that have not been previously reported is possible, this can be attributed to the distribution area in the Malay Peninsula which is geographically very close to Sumatran Island.

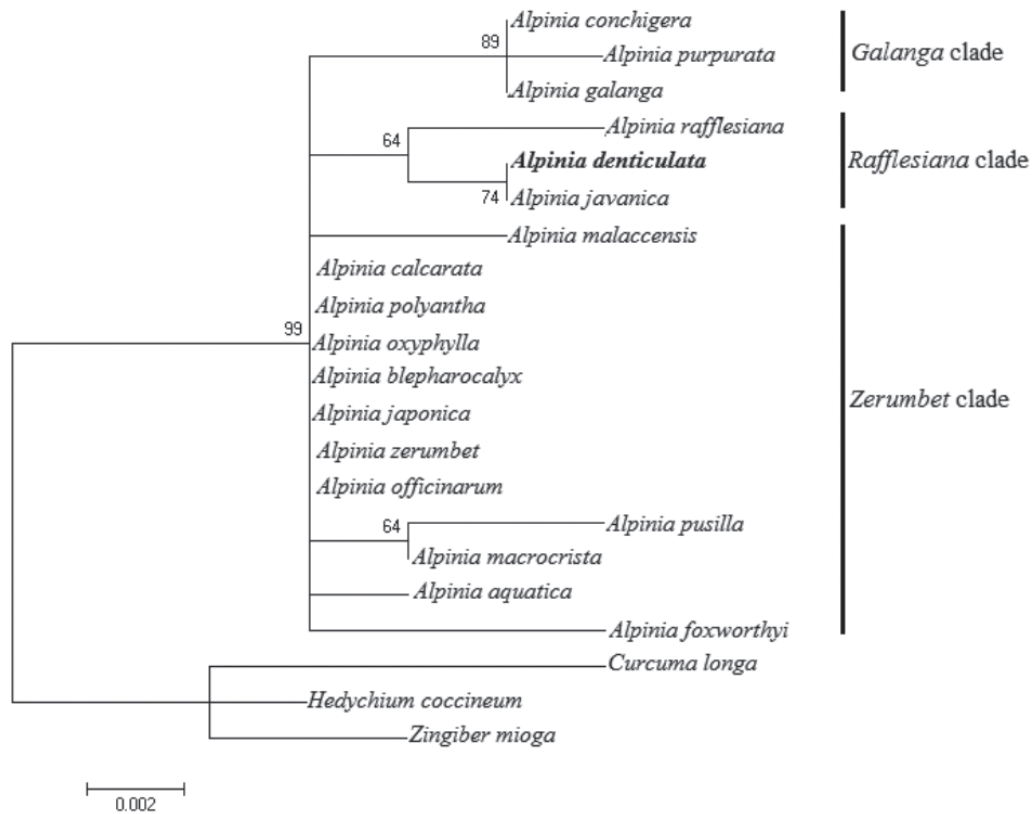
Maximum Likelihood (ML) analysis, which is

commonly used, provides the most robust result and Maximum Parsimony (MP) that uses character-based method, is chosen to construct the phylogenetic tree. The phylogenetic analysis carried out shows the position of the 2 new records of *Alpinia* along with other existing species (Figs 3–6). The ITS+trnH-psbA combination gave a better result than single marker analysis of ITS and trnH-psbA (Figs 5, 6). This result can be seen from the percentage of bootstrap value.

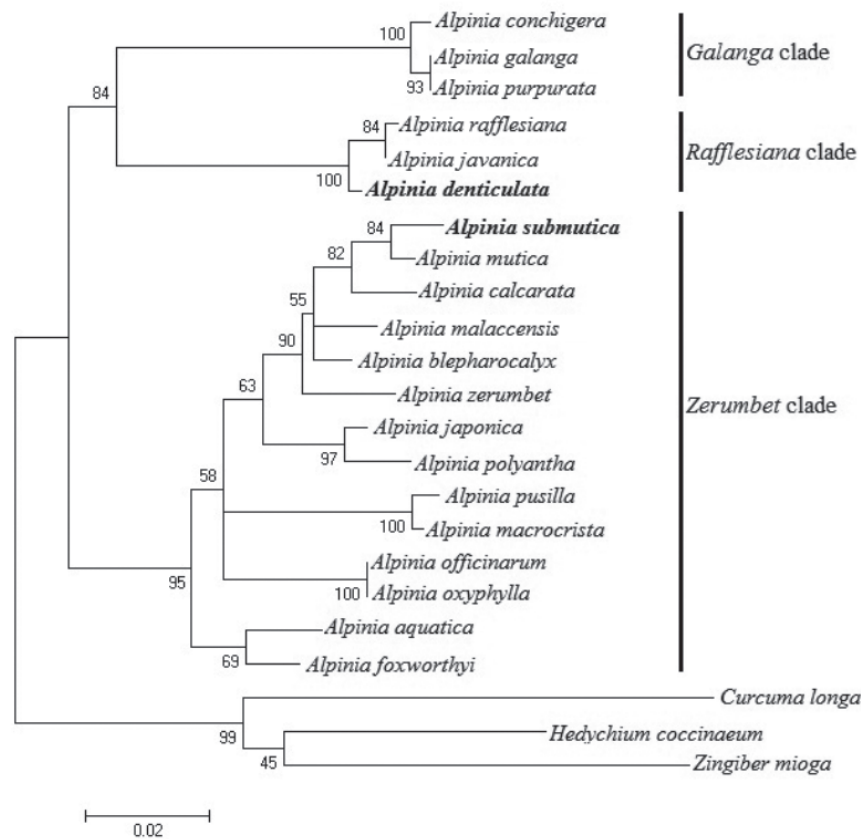
Based on 7 species that were sequenced, the results of sequence divergence range are 0.2–12.3%. *Alpinia denticulata* and *A. submutica* were added for the first time



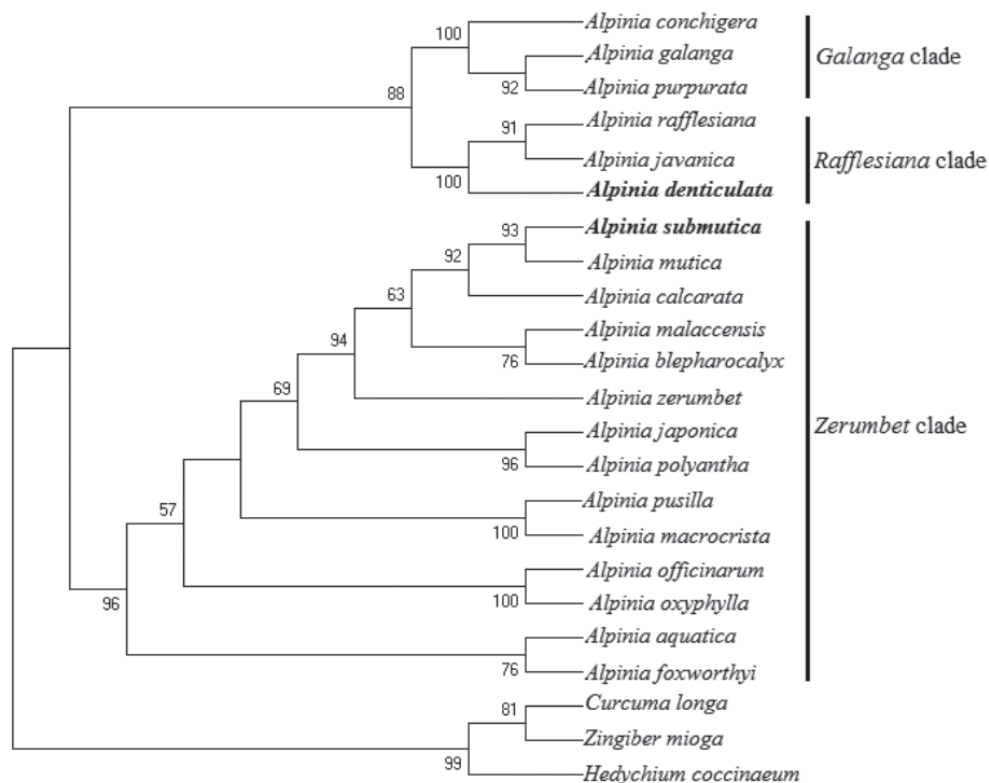
**Figure 3.** Maximum Parsimony phylogenetic tree (length = 245) based on complete sequences of Internal Transcribed Spacer (ITS) region of *Alpinia* from this study and GenBank data sequences with bootstrap values ( $\geq 50\%$ ). Cluster name based on Kress et al. (2005a).



**Figure 4.** Maximum Likelihood phylogenetic tree ( $-\log$  likelihood =  $-848.30$ ) based on complete sequences of *trnH-psbA* Intergenic Spacer region of *Alpinia* from this study and GenBank data sequences with bootstrap values ( $\geq 50\%$ ). Cluster name based on Kress et al. (2005a).

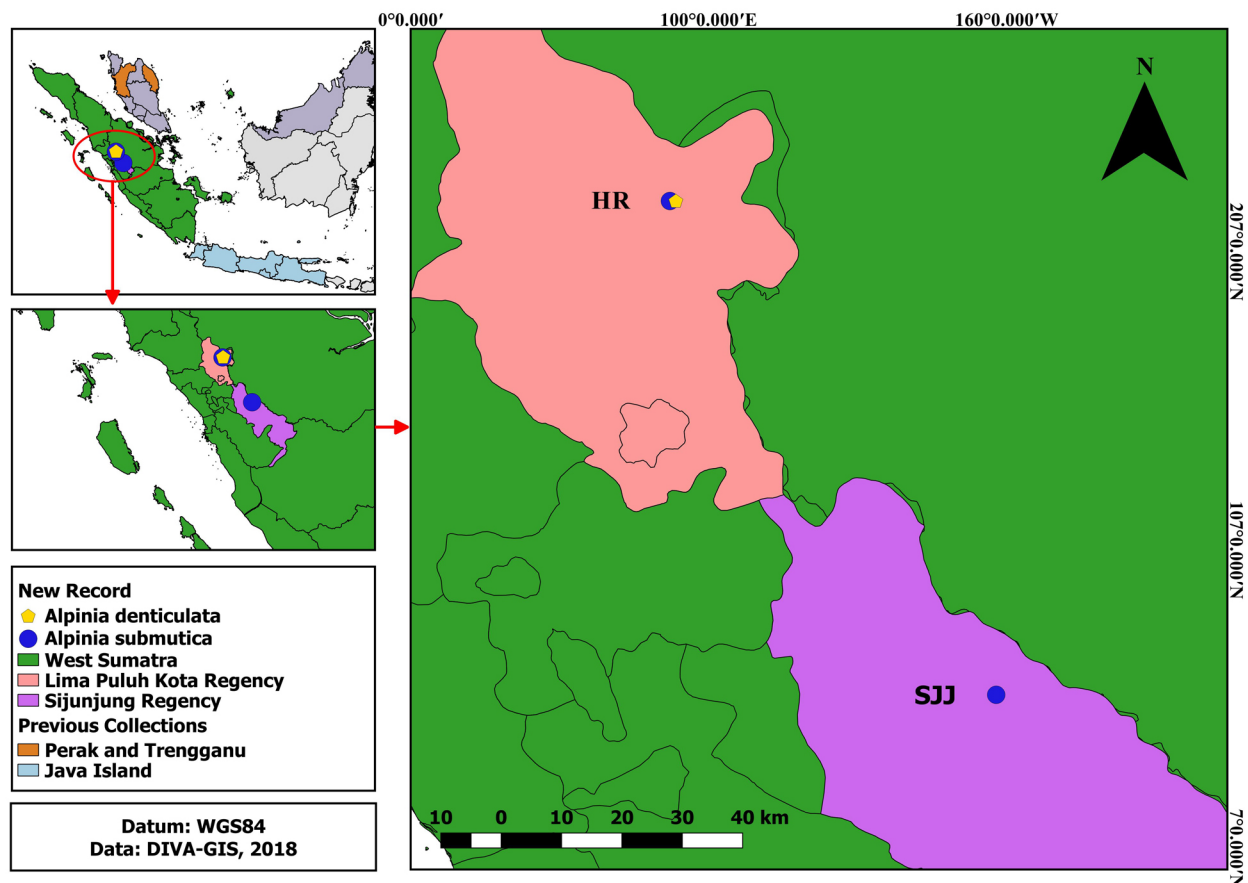


**Figure 5.** Maximum Likelihood phylogenetic tree ( $-\log$  likelihood =  $-1988.44$ ) based on combined ITS and *trnH-psbA* Inter-genic Spacer region from this study and GenBank sequences data with bootstrap values ( $\geq 50\%$ ). Cluster name based on Kress et al. (2005a).



**Figure 6.** Maximum Parsimony phylogenetic tree (length = 245) based on combined ITS and *trnH-psbA* Inter-genic Spacer region from this study and GenBank sequences data with bootstrap values ( $\geq 50\%$ ). Cluster name based on Kress et al. (2005a).





**Figure 7.** Collection side of 2 new records in Sumatra and previous collections of them based on Royal (2018) database and Schumann (1904b).

in the grouping of the the *Alpinia supertree* by Kress et al. (2005a). Based on its position, *A. submutica* is a sister taxon of *A. mutica* with a sequence divergence value of 1.2% and percentage bootstrap value more than 84% for ITS and combined ITS+*trnH-psbA*. They have similar morphological characteristics, but in previous studies were considered as distinct species because of the differences in fruit morphology. From this morphological and molecular character, it can be ascertained that both are different species.

Meanwhile, *A. denticulata* is in the same cluster with *A. rafflesiana* and *A. javanica* in Rafflesian Clade (Figs 3, 4). The sequence divergence value range of this cluster is 0.8–1% with strong bootstrap support of 100% for ITS and combined ITS+*trnH-psbA*, but weak bootstrap support (64%) for *trnH-psbA*. Rafflesian Clade group is not based on similarity of morphological characters, because *A. denticulata* has a small flowers and cylindrical ovary, while *A. rafflesiana* and *A. javanica* have large flowers with the square ovary. This cluster is clearly formed more on the similarity of distribution area in the Malay Peninsula (Kress et al. 2005a).

**Conclusion.** *Alpinia denticulata* and *A. submutica* are newly recorded species of *Alpinia* in Sumatra, supported by their morphological descriptions and a phylogenetic analysis.

## Acknowledgements

We thank Herbarium Andalas University (ANDA) and Herbarium Bogoriense (BO) for the facilities provided to conduct this research. We also thank Andalas University for Professor's Grant 2017 (to DA). We also thank Dr Dewi Imelda Rosma for the laboratory facilities provided related to molecular work and Dr Marlina Ardiani for discussion on molecular data.

## References

- Ardiyani M (2016) A new species of Zingiber (Zingiberaceae) from Enggano Island, Indonesia. *Reinwardtia* 14: 307–310. <https://doi.org/10.14203/reinwardtia.v14i2.1676>
- de Vogel E (1987) *Manual of Herbarium Taxonomy*, UNESCO, Jakarta, 164 pp.
- Doyle JJ, Doyle JL (1987) A rapid procedure for DNA purification from small quantities of fresh leaf tissue. *Phytochemical Bulletin* 19: 11–15.
- Heyne K (1987) *Tumbuhan berguna indonesia*. Badan Penelitian dan Pengembangan Kehutanan, Departemen Kehutanan 2: 1188–1189.
- Holttum RE (1950) The Zingiberaceae of the Malay Peninsula. *Gardens' Bulletin Singapore*. 13: 1–249.
- Kearse M, Moir R, Wilson A, Stones-Havas S, Cheung M, Sturrock S, Buxton S, Cooper A, Markowitz S, Duran C (2012) Geneious Basic: an integrated and extendable desktop software platform for the organization and analysis of sequence data. *Bioinformatics* 28: 1647–1649. <https://doi.org/10.1093/bioinformatics/bts199>.



- Accessed on: 2018-12-27.
- Kress WJ, Erickson DL (2007) A two-locus global DNA barcode for land plants: the coding *rbcl* gene complements the non-coding *trnH-psbA* spacer region. *PLoS ONE* 2: e508. <https://doi.org/10.1093/bioinformatics/bts199>
- Kress WJ, Liu AZ, Newman M, Li QJ (2005a) The molecular phylogeny of *Alpinia* (Zingiberaceae): a complex and polyphyletic genus of gingers. *American Journal of Botany* 92: 167–178. <https://doi.org/10.3732/ajb.92.1.167>
- Kress WJ, Prince LM, Williams KJ (2002) The phylogeny and a new classification of the gingers (Zingiberaceae): evidence from molecular data. *American Journal of Botany* 89: 1682–1696. <https://doi.org/10.3732/ajb.89.10.1682>
- Kress WJ, Wurdack KJ, Zimmer EA, Weigt LA, Janzen DH (2005b) Use of DNA barcodes to identify flowering plants. *Proceedings of the National Academy of Sciences* 102: 8369–8374. <https://doi.org/10.1073/pnas.0503123102>
- Kumar S, Stecher G, Tamura K (2016) MEGA7: molecular evolutionary genetics analysis version 7.0 for bigger datasets. *Molecular Biology and Evolution* 33: 1870–1874. <https://doi.org/10.1093/molbev/msw054>
- Lamb A, Gobilik J, Ardiyana M, Dalberg-Poulsen A (2013) A Guide to Gingers of Borneo. Natural History Publications, Borneo, 24–35.
- Larsen K, Ibrahim H, Khaw S, Saw L (1999) Gingers of Peninsular Malaysia and Singapore. Natural History Publications, Borneo, 57–58.
- Larsen K, Lock J, Maas H, Maas P (1998) Zingiberaceae. In: Kubitzki K (Ed), *The Families and Genera of Vascular Plants*. Springer-Verlag, Germany, 474–495.
- Namsa ND, Tag H, Mandal M, Kalita P, Das A (2009) An ethnobotanical study of traditional anti-inflammatory plants used by the Lohit community of Arunachal Pradesh, India. *Journal of Ethnopharmacology* 125: 234–245. <https://doi.org/10.1016/j.jep.2009.07.004>
- Nelvita R (2009) Zingiberaceae in Several Limestone Areas of West Sumatra. Andalas University, West Sumatra, 136 [Unpublished].
- Newman M, Lhuillier A, Poulsen AD (2004) Checklist of the Zingiberaceae of Malesia, Nationaal Herbarium Nederland, Universiteit Leiden Branch, Leiden, 166.
- Nurainas (2007) Wild Ginger (Zingiberaceae) Diversity from Rimbo Panti Nature Reserved Forest in West Sumatra. Young lectures research report, Andalas University, West Sumatra, 38 [unpublished].
- Nurainas N, Arbain D (2017) A new species and a new record of Zingiberaceae from Sumatra, Indonesia. *Taiwania* 62: 294–298. <http://doi.org/10.6165%2ftai.2017.62.294>
- Poulsen AD (2006) Gingers of Sarawak. Natural History Publication, Kota Kinabalu, 18 pp.
- Royal Botanic Garden Edinburgh (2018) Royal Botanic Garden Edinburgh Herbarium (E). Occurrence dataset. <https://www.gbif.org/occurrence/574924177>. Accessed on: 2018-10-28. <https://doi.org/10.15468/ypoir>
- Schumann K (1904a) Zingiberaceae. Das pflanzenreich, Zingiberaceae 4: 46 [Heft 20], p. 354, fig. 42G. [http://plantillustrations.org/illustration.php?id\\_illustration=179724](http://plantillustrations.org/illustration.php?id_illustration=179724). Accessed on: 2018-10-07.
- Schumann K (1904b) Zingiberaceae mit 355 Einzelbild. in 52 Fig. In: Engler A (Ed) *Pflanzenreich. Regni vegetabilis conspectus*. Im Auftrage der Königl. preuss. Akademie der Wissenschaften. Verlag von Wilhelm Engelmann, Leipzig, 329.
- Smith R (1981) Synoptic keys to the genera of Zingiberaceae pro parte. Royal Botanical Garden Edinburgh, Departmental Publication series 2: 1–28.
- Smith R (1990) *Alpinia* (Zingiberaceae): a proposed new infrageneric classification. *Edinburgh Journal of Botany* 47: 1–75. <https://doi.org/10.1017/S0960428600003140>
- Thompson JD, Gibson TJ, Plewniak F, Jeanmougin F, Higgins DG (1997) The CLUSTAL\_X windows interface: flexible strategies for multiple sequence alignment aided by quality analysis tools. *Nucleic Acids Research* 25: 4876–4882. <https://doi.org/10.1093/nar/25.24.4876>
- van Valkenburg J, Bunyapraphatsara N (Eds) (2001) Medicinal and Poisonous Plants 2. In: *Plant Resources of South-East Asia*, Volume 12. Backhuys Publisher, Leiden, 52–61.
- White TJ, Bruns T, Lee S, Taylor J (1990) Amplification and direct sequencing of fungal ribosomal RNA genes for phylogenetics. In: Innis MA, Gelfand DH, Sninsky JJ, White TJ (Eds) *PCR protocols: A Guide to Methods and Applications*. Academic Press, New York, 315–322.