MORPHOLOGICAL AND MOLECULAR CHARACTERISTICS OF *Panax* sp. (Araliaceae) FROM PHU XAI LAI LENG MOUNTAIN, NGHE AN PROVINCE, VIETNAM

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ABSTRACT: Seven sterile and young specimens of *Panax* sp. were collected in Phu Xai Lai Leng silicat mountain, the highest mountain peak in central Vietnam. Morphologically, they resemble *P. stipuleanatus* and *P. bipinnatifidus*. However, they have stipules and leaflets that are not bipinnatifid. The ITS-DNA sequences of *Panax* sp. from Phu Xai Lai Leng are identical and exhibit a close relationship with *P. stipuleanatus* from Tam Duong district, Lai Chau province with strong bootstrap support (98%). Noticeably, these two taxa are different by 2 nucleotides. A collection of standard specimens including inflorescence, infructescence, flowers and ripe fruits of *Panax* from Phu Xai Lai Leng is needed to determine its taxonomic status.

Keywords: Panax, ITS-rDNA, morphology, Phu Xai Lai Leng silicate mountain.

INTRODUCTION

In the world, the genus *Panax* comprises 6 species distributing in North America and East Asia [5, 6]. The Plant List included 12 valid species of *Panax* [8]. In China, 6-7 species *Panax* are recognized [9]. In Vietnam, this genus consists of three known [1, 2, 3, 4], namely, *P. bipinnatifidus* Seem., *P. stipuleanatus* and *P. vietnamensis*. The occurrence of three other species, namely, *P. pseudoginseng* [9], *P. zingiberensis* and *P. notoginseng* [9] are doubtful and should be confirmed by voucher specimens.

During a recent survey of *Panax* plants on the north-eastern slope of Phu Xai Lai Leng mountain, western Nghe An Biosphere Reserve, north central Vietnam, we collected 7 specimens belonging to this genus with leaves similar to those of *P. pseudoginseng*, *P. bipinnatifidus* and *P. stipuleanatus*. In this paper, we provide morphological and molecular characteristics of the collected specimens.

MATERIALS AND METHODS

Morphological characteristics

Seven *Panax* specimens, mostly sterile and young (except specimen *NTL 101* which is

adult) were collected in Phu Xai Lai Leng Mountain (table 1). Morphological characteristics were described from the fresh specimens in the field and laboratory. Photographs were taken using a "Canon EOS 200" camera with "Canon" and "Cosina" lenses of various focal distances. Voucher herbarium specimens were deposited in the herbaria of the Vietnam National Museum of Nature (VNMN) and of the Hanoi University of Science (HNU).

Molecular characteristics

Total DNA was extracted from dried leaves using the DNeasy Plant Minikit (Oiagen, Singapore). The ITS-rDNA regions were amplified using the forward primer PaITSF 5'-CAC TGA ACC TTA TCA TTT AG AG -3' and the reverse primer PaITSR 5'-CTT ATT GAT ATG CTT AAA CTC AG-3' was designed based on the ITS sequence of the genus Panax obtained from GenBank. After purification, DNA fragments were sequenced with a BigDye Terminator Cycle Sequencing Ready Reaction kit (PE Applied Biosystems, Foster City, CA, USA) and run on an ABI PRISM 3100 Genetic Analyzer (PE Applied Biosystems). All sequences were submitted to Genbank with the accession number as follow: KP091694 (NTL101), KP091695 (NTL106), KP091698 (NTL102), KP091699 (NTL104), KP091696 (NTL105), KP091697 (NTL103), KP091700 (NTL107).

Table 1. Panax sp. specimens collected in Phu Xai Lai Leng mountain, western Nghe An province

N°	Code	Latitude, N	Longitude, E	Altitude a.s.l.	Habit
1	NTL 101	19°13′16"	104°11'59"	1599 m	Adult plant, 20-30 cm tall with fallen inflorescence
2	NTL 102	19°13'23"	104°11'57"	1595 m	Young plant, 20-30 cm tall
3	NTL 103	19°13'22"	104°11'58"	1599 m	Young plant, 20-25 cm tall
4	NTL 104	19°13'23"	104°11'57"	1597 m	Young plant, 20-25 cm tall
5	NTL 105	19°13'18"	104°11'59"	1627 m	Young plant, 20-25 cm tall
6	NTL 106	19°13'18"	104°11'59"	1627 m	Young plant, 20 cm tall
7	NTL 107	19°13'18"	104°11'59"	1627 m	Young plant, 17 cm tall

Sequences of 7 *Panax* specimens from Phu Xai Lai Leng mountain were aligned using Clustal X 1.64 [9] with sequences of *Panax* spp., according to Phan et al. [4].

Equally weighted maximum parsimony (MP) analyses were performed using PAUP* (4.0 beta version) [7]. A heuristic search procedure was used with the following settings: ten replicates of random taxon addition, tree-bisection reconnection branch swapping, multiple trees retained, no steepest descent, and accelerated transformation. Gaps were treated as missing data. Bootstrap analysis was carried out with 100 replicates.

RESULTS AND DISCUSSION

Morphological characteristics

According to the leaf morphology, the 7 specimens of *Panax* collected from Phu Xai Lai Leng can be identified as two groups: group PXLL-1 with divided leaflets and group PXLL-2 with undivided leaflets.

Groups of specimens PXLL-1: Leaflets from slightly lobed to pinnatifid (specimens NTL 101, NTL 102 & NTL 103) (fig. 1).

Herb, perennial, 0.2-0.3 m tall (without inflorescence/infructescence). Rootstock moniliform-mounded, usually horizontal, resembling that of species of *Zingiber*. All plant parts glabrous except for scattered setulose hairs on veins of adaxial leaf surface, very rare on abaxial leaf surface; trichomes 1.5-2 mm long. Leaves of adult plants usually 3-4, verticillate at

apex of stem, palmately compound; petioles 3-9 cm long, base with stipules or stipule-like appendages, deltoid-cuspidate, less than 2 mm long. Leaflets 5 (-7), membranous, from slightly lobed to pinnatifid, margin serrulate; central leaflet largest, up to 17 × 6 cm, elliptic, rarely oblanceolate in outline, acute at base, cuspidate or long cuspidate, rarely acuminate at apex with leaf blade 2.4-2.8 as long as wide, petiolules 0.8-1.5 cm long, at base sometimes with some trichomes similar to the ones on nerves of leaf blades: lateral leaflets smaller: the lowermost pair of leaflets smallest, rhombic-ovate or obovate, asymmetric, base broadly cuneate or subrounded, margin serrate. Inflorescence and infructescence, flowers and fruits unknown.

The specimens differ from *Panax stipuleanatus* by the smaller vegetative parts, not up to 0.4-0.6 m, petiole base with stipules or stipule-like appendages and leaflets not 2-pinnatifid. Description of rhizome of this species made by Xiang & Lowry II (2007) [9] as fusiform was inaccurate, but rather "thick, elongate, finally zigzag" as observed in the isotype specimen *Feng 13694* (KUN 0448506!).

The specimens differ from *Panax bipinnatifidus* by the leaflets which are not bipinnatifid, adaxially sparsely setose, and base of petiole with stipules or stipule-like appendages.

Groups of specimens PXLL-2: Leaflets undivided (NTL 104, NTL 105, NTL 106 & NTL 107). All specimens are young plants (fig. 1)

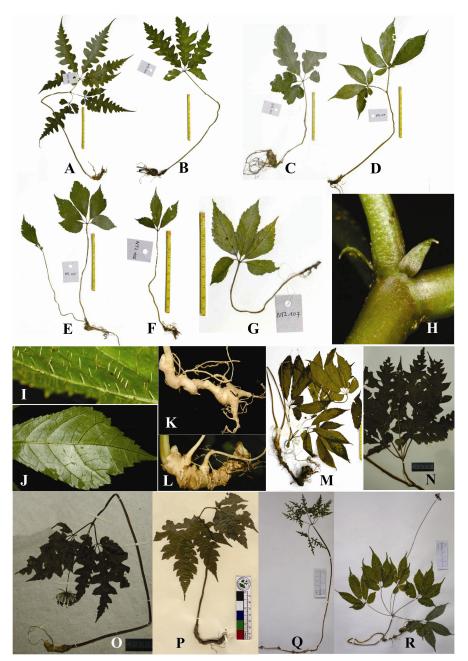


Figure 1. A- NTL 101; B- NTL 102; C- NTL 103; D- NTL 104; E- NTL 105; F- NTL 106; G- NTL 107; H- Stipules; I- Scattered setulose hairs on veins on adaxial leaf surface; J- Abaxial leaf surface; K- Rhizome of NTL 106; L- Rhizome of NTL 101; M- Panax stipuleanatus H.T.Tsai & K.M.Feng, Vietnam: Lai Chau, Tam Duong, Khun Ha, P 11172 (VNMN!, HNU!); N- Panax stipuleanatus, Vietnam: Lao Cai, Sa Pa, Sino-Viet. Exp. 344 (KUN 0560456!); O- Panax stipuleanatus, China, 23014 (KUN 0560447!); P- Panax stipuleanatus-Type: China: Yunnan, Feng 13694 (Iso- KUN 0448506!); Q- Panax bipinnatifidus Seem., forma 2-pinnatifid. China: Xizang (After J. Wen, 1539155!); R- Panax bipinnatifidus Seem., forma undivided. China: Yunnan (After J. Wen, 1410528!).

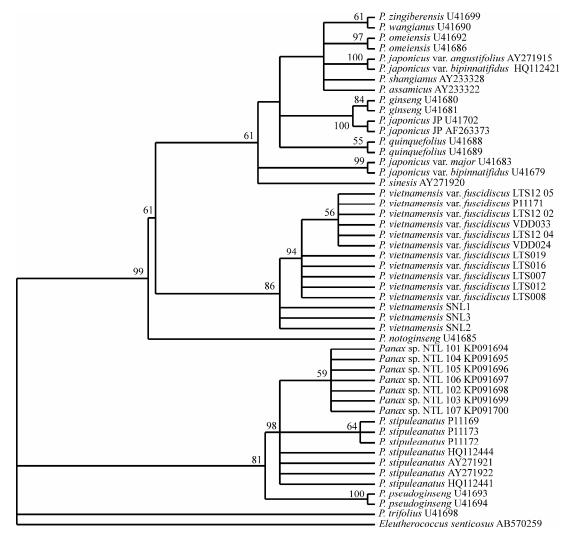


Figure 2. The molecular relationship of *Panax* sp. collected from Phu Xai Lai Leng Mountain, Western Nghe An Province with other *Panax* spp. based on MP analysis. *Eleutherococcus senticosus* AB570259 is the outgroup taxon.

This group is close to PXLL-1 but differs only by leaflets undivided, not lobed nor divided.

Herb, perennial, 0.17-0.25 m tall (without inflorescence/infructescence). Rootstock moniliform-mounded, usually horizontal, resembling that of species of *Zingiber*. All plant parts glabrous except for scattered setulose hairs on veins of adaxial leaf surface, very rare on abaxial leaf surface; trichomes 1.5-2 mm long. Plant young, having two leaves opposite or one leaf on top of stem; petiole base with stipules deltoid-cuspidate or stipule-like appendages.

Leaves palmately compound; petiole ca. 3.5-5 cm long; leaflets (3-)5(-7), oblanceolate in outline, acute at base, cuspidate (1.5-2 cm), rarely acuminate at apex, membranous, undivided, margin serrulate. Median leaflet biggest, up to $6\text{-}11 \times 1.5\text{-}3.5$ cm including petiolule with leaf blade $2.7\text{-}2.9 \times$ as long as wide and petiolules 0.2-0.5 cm long; lateral leaflets smaller, two lowest leaflets smallest, asymmetric, blades entire, ca. 1.5×0.9 cm, serrulate at top. Inflorescence and infructescence, flowers and fruits unknown. Rare.

The length of the *Panax* sp. ITS-rDNA

sequence for alignment with an outgroup taxon

was 587 bp and all of them are identical. MP

analysis of this alignment indicated that among 588 characters, 69 were parsimony informative.

In the consensus MP tree (fig. 1), the *Panax* sp.

is clustered with the clade of P. stipuleanatus

and P. pseudoginseng with a moderate bootstrap

support (81%) and they have sister relationship

with P. stipuleanatus (KJ418187, KJ418197,

KJ418198) from Tam Duong, Lai Chau with

The specimens differ from Panax stipuleanatus by the smaller vegetative parts, petiole base with stipules or stipule-like appendages and leaflets not bipinnatifid.

specimens differ The from Panax bipinnatifidus in the undivided leaflets not lobed nor divided as in Panax bipinnatifidus and by the adaxially sparsely setose leaflets and the base of petiole with stipules or stipule-like appendages.

Molecular relationship with other species

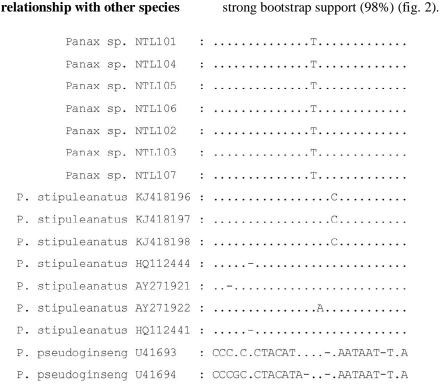


Figure 3. The nucleotide variation in the ITS-rDNA region of the P. stipuleanatus clade

Sequence variation

The ITS-rDNA sequence of Panax sp. was 587 bp, the same as for *P. stipuleanatus* from Tam Duong, Lai Chau (KJ418187, KJ418197, KJ418198) and P. stipuleanatus AY271922; longer than that of P. stipuleanatus (HQ112444, AY271921, HQ112441) and P. pseudoginseng (U41693, U41694). The ITS-rDNA sequences of Panax sp. differed from sister clade of P. stipuleanatus (KJ418187, KJ418197, KJ418198) by 2 nucleotides (fig. 3).

Although the specimens of *Panax* sp. collected from Phu Xai Lai Leng mountain, western Nghe An Province were divided to two groups, PXLL-1 and PXLL-2 based on the morphology of leaflets, the ITS-rDNA sequences of all these specimens were identical. Therefore they should be regarded as the same infraspecific taxon.

CONCLUSION

According to the morphology of leaflets and molecular data, the Panax sp. collected from Phu Xai Lai Leng mountain, western Nghe An Province could be considered as a new variety of *Panax*. Additional collection including inflorescence, infructescence, flowers and ripe fruits of the *Panax* sp. from Phu Xai Lai Leng is needed to determine its taxonomic status.

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ĐẶC ĐIỂM HÌNH THÁI VÀ PHÂN TỬ CỦA *Panax* sp. (Araliaceae) THU Ở NÚI PHU XAI LAI LENG, TỈNH NGHỆ AN, VIỆT NAM

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TÓM TẮT

Bài báo trình bày kết quả phân tích 7 mẫu không mang cơ quan sinh sản và còn non của Sâm *Panax* sp. được thu thập trên núi đá silicát Phu Xai Lai Leng, núi cao nhất Bắc Trung bộ Việt Nam, ở vị trí 19°13' vĩ Bắc, 104°12' kinh Đông, độ cao 1600 m so với mặt nước biển. Về hình thái, chúng rất gần với loài *P. stipuleanatus* và *P. bipinnatifidus*. Tuy nhiên, chúng có lá kèm và lá chét không xẻ lông chim 2 lần. Trình tự vùng gen ITS-DNA của *Panax* sp. ở Phu Xai Lai Leng giống hệt nhau, và có mối quan hệ gần gũi với *P. stipuleanatus* thu ở Tam Đường, Lai Châu với bootstrap lên đến 98%. Tuy nhiên, giữa chúng có sự sai khác ở 2 nucleotide. Việc thu thập mẫu có đầy đủ cụm hoa, cụm quả, hoa và quả chín của sâm *Panax* sp. ở Phu Xai Lai Leng là ưu tiên hàng đầu để xác định vị trí phân loại của loài.

Từ khóa: Panax, ITS-rDNA, hình thái, núi đá xilicát, Phu Xai Lai Leng.

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