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Research Article

Standardization parameters, Internal Transcribed
Spacer nucleotide sequence and their anti-malarial
activity of *Eurycoma longifolia* Jack

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ABSTRACT

Determination of pharmacognostic parameters plays a significant role for standardization of medicinal plants. The objective of recent study aims to examine the standardization parameters of *Eurycoma longifolia* Jack in Thailand, the pharmacognostic specifications of foreign matter, acid insoluble ash, total ash, ethanol-soluble content and water content of *Eurycoma longifolia* Jack were evaluated according to World Health Organization guideline. The ITS region was amplified by PCR and sequenced. Dried root of *Eurycoma longifolia* Jack was exhaustively extracted with ethanol and the obtained dried extract were analyzed with thin layer chromatography. The *in vitro* activity of the extract against *P. falciparum* strain 3D7 and K1 were also evaluated using SYBR green I-based assay. The results indicated that the ethanolic extract of dried root showed a great antimalarial activity against both *P. falciparum* 3D7 and K1 strain with the IC₅₀ of 2.16 µg/ml and 1.79 µg/ml respectively. The PCR amplification of ITS region generated the PCR product approximately 750 base pairs (bp) in size. The alignment of ITS sequences showed 98-99% homology, 12 single nucleotide polymorphisms and 23 Indels were detected. The Combination of standardization parameters, biomolecular method and biological activity are useful for correct identification and quality control of *Eurycoma longifolia* Jack.

Keywords: *Eurycoma longifolia* Jack, standardization parameters, nucleotide sequence, internal transcribed spacer (ITS) region, antimalarial

1. INTRODUCTION

The increasing demand in herbal medicine has stimulated the improvement in analysis and quality control of their herbal materials and products. To ensure reproducible quality of any herbal plants, requirements and methods for research and evaluation are essential. *Eurycoma longifolia* Jack is a popular medicinal plant in the family Simaroubaceae consists of 200 species. It is indigenous to Southeast Asia including Thailand. All parts of *E. longifolia* Jack has been traditionally used for its antimalarial, antipyretic, antidiabetic, Aphrodisiac, and antimicrobial activities¹⁻⁵. Roots extracts have also been reported for enhancing testosterone levels. In modern dietary supplements, *E. longifolia* can be use to improve sexual function, restore hormonal balance, and improve physical and

mental energy due to their chemical composition that have different effects in the body⁶⁻⁷. A wide range of chemical components have been isolated and characterized from *Eurycoma longifolia*⁸⁻¹⁰. Although these plants have been widely used in traditional medicine, there is still limited scientific report undertaken in their standardization for safety reason. Moreover, many closely related species have similar morphology and growth conditions as well as many other variables such as the storage condition, environmental condition, and the harvest processing may affect the analysis and quality control of the herbal materials and products. Therefore, additional of molecular method have been developed for medicinal plants authentication. Sequence variation of the Internal transcribe spacer (ITS) region in the

nuclear DNA has been applied widely for identification in species level and can be further applied for investigation the adulteration of herbal drugs¹¹.

Standardization of dried root of *E. longifolia* Jack was carried out in the present study. Combination of several characteristics such as standardization, biological activity and DNA based analysis have been introduced in this study for correct identification and quality control of this Thai herbal plant. This study aims to examine the standardization parameters, ITS region sequence variation and their antimalarial activity of *Eurycoma longifolia* Jack in Thailand. The information from this study will provide data which is useful in the correct identification and authentication and may help in preventing its adulteration.

2. MATERIAL AND METHODS

2.1 Plants collection

Whole plants of *E. longifolia* Jack were collected from several localities in Thailand. Plants were authenticated by botanical expert and comparing with the herbarium specimens at Forest Herbarium Thailand (BKF). All samples are deposited at College of Public Health Sciences, Chulalongkorn University.

2.2 Standardization parameters

Dried root of *E. longifolia* Jack were collected from 15 Thai traditional drug stores in 11 provinces in Thailand and evaluated according to World Health Organization (WHO) quality control method for medicinal plant materials¹². The pharmacognostic specifications of foreign matter, acid insoluble ash, total ash, ethanol-soluble content and water content were determined and presented as mean \pm SD. Thin layer chromatography (TLC) of *E. longifolia* Jack roots ethanolic extract was also evaluated. The TLC chromatogram was captured by a digital scanner (Hewlett Packard Deskjet F2280).

2.3 Antimalarial sensitivity assay

Dried root of plant material was grounded to powder and exhaustively extracted with ethanol using Soxhlet apparatus. The extract was filtered through filter paper then concentrated and stored at -20°C. The *in vitro* activity of the extracts were evaluated against *P. falciparum* strain 3D7 (chloroquine sensitive) and K1 (chloroquine resistant) using SYBR green I-based assay¹³. The extract was dissolved in 50% ethanol to give the concentration of 1 mg/ml and then diluted with complete media to obtain the concentration range 0-100 μ g/ml. Dihydroartemisinin (DHA) was used as positive control. Fifty microlitre parasite suspension and 50 μ l of different concentration of the

extracts were mixed together in 96 well plate and incubated at 37°C under a gas mixture of 50% N₂, 5% O₂ and 5% CO₂ for 2 days. One hundred microlitre of fluorescent haemolysis reagent (0.01% of fluorescent dye SYBR Green I in lysis buffer) was added to each well and incubated the plates in the dark for 1 hours. Fluorescence intensity was determined at the excitation and emission wave lengths of 485 and 530 nm, respectively. Mean values were calculated from the triplicate results. The percent growth inhibition was calculated and the IC₅₀ was determined from a log dose-response analysis using the CalcuSynTM computer program.

2.4 DNA extraction and amplification of ITS region

Genomic DNA was individually extracted from the fresh young leaves of *E. longifolia* Jack using a modified CTAB technique¹⁴. The ITS region was amplified using a pair of universal primer ITS4 and ITS5¹⁵. Amplifications were carried out in 20 μ l reaction mixture containing of 1x PCR buffer [100mM Tris-HCl (pH 8.8), 500mM KCl, 0.8% (v/v) Nonidet P40], 2.5mM MgCl₂, 0.1mM dNTPs, 0.1 μ M of each primer, 0.5 Unit of *Taq* polymerase (Fermentas) and 1 μ l of DNA template. The PCR amplification reaction conditions were as following: an initial denaturation for 5 min at 95°C, 30 cycles of denaturation at 95°C for 30 second, annealing at 55°C for 30 second and extension at 72°C for 30 second, and then follow by a final 5 min extension at 72°C. The PCR products were evaluated by 1.5% agarose gel electrophoresis, stain with ethidium bromide and visualized under UV transilluminator. The amplified PCR products were then purified by PCR purification kit (QIAGEN) prior sequencing of both sense and antisense strand. ClustalW program was used to multiple aligned the sequence of the ITS region.

3. RESULTS

3.1 Standardization parameters

The pharmacognostic study of *E. longifolia* Jack was determined from 15 Thai traditional drug stores from 11 provinces of Thailand (**Table 1**) and TLC fingerprinting of ethanolic extract was showed in **Figure 1**.

3.2 Antimalarial activity

The susceptibility of *P. falciparum* strain K1 and 3D7 to the ethanolic extract of *E. longifolia* Jack root was investigated *in vitro* based on SYBR Green I assay. The IC₅₀ value (drug concentration that inhibits the parasite growth by 50%) was shown in **Table 2**. Dihydroartemisinin was used as positive control.

Table 1
The pharmacognostic constant values determination in *E. longifolia* Jack

Specification	Mean \pm SD	Min - Max
Foreign matter	0.168 \pm 0.179	0 - 0.705
Acid-insoluble ash	0.696 \pm 0.064	0.503 - 0.889
Total ash	1.973 \pm 0.039	1.857 - 2.088
Ethanol-soluble extractive	1.733 \pm 0.178	1.199 - 2.267
Water-soluble extractive	4.888 \pm 0.376	3.760 - 6.017
Loss on drying	7.515 \pm 0.252	6.760 - 8.271
Volatile oil content	0	0
Water content	13.352 \pm 0.839	10.836 - 15.869

Table 2
 IC_{50} of *Eurycoma longifolia* Jack extracts against *P. falciparum* strain 3D7 and K1

	IC_{50} (μ g/ml)	
	3D7	K1
<i>E. longifolia</i> Jack root ethanolic extracts	2.16	1.79
Dihydroartemisinin	1.97	1.46

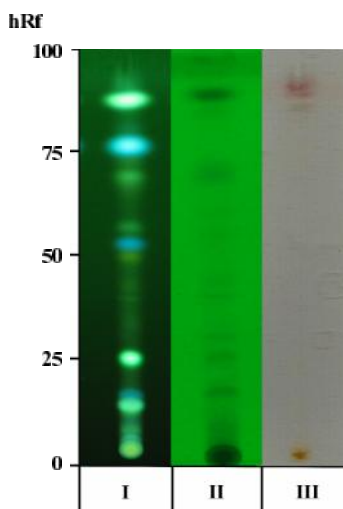


Figure 1
TLC fingerprinting of ethanolic extract from dried root of *E. longifolia* Jack using solvent system; chloroform: methanol = 9:1
Panel I: Detection under UV light (366 nm), Panel II: Detection under UV light (254 nm), and Panel III: Detection with Anisaldehyde.

EL-NK	CGTA-CAAGGTT-CCGTAGGGGA---CTGC-GGAGGA-CATTGTCGAAACCTGCCAGC	53
EL-AB	CGTAA-CAAGGTTTCCGTAGGTGAA-CCTGC-GGAAGGATCATTGTCGAAACCTGCCAGC	58
EL-PB	CGTAA-CAAGGTTTCCGTAGGTGAAACCTGCAGGAAGGATCATTGTCGAAACCTGCCAGC	60
	*	
EL-NK	AGAACGACCCCGGAACGAGTGATGAAAACCGG-AGTGGTGGGGCGTGC GGGTGTCGCTCG	112
EL-AB	AGAACGACCCCGGAACGAGTGATGAAAACCGG-AGTGGTGGGGCGTGC GGGTGTCGCTCG	117
EL-PB	AGAACGACCCCGGAACGAGTGATGAAAACCGG-AGTGGTGGGGCGTGC GGGTGTCGCTCG	120
EL-NK	CTCGCGTCTCCCTCCTCGCTGCCTGGTGGGAGGGCCCCGTCCGCGGGACCCCTGCCCGGG	172
EL-AB	CTCGCGTCTCCCTCCTCGCTGCCTGGTGGGAGGGCCCCGTCCGCGGGACCCCTGCCCGGG	177
EL-PB	CTCGCGTCTCCCTCCTCGCTGCCTGGTGGGAGGGCCCCGTCCGCGGGACCCCTGCCCGGG	180
EL-NK	GAGCAAAAACGAACCCCGGCGGGACTGCGCCAAGGAAAACAAAACGAGAGAGCATGCCCT	232
EL-AB	GAGCAAAAACGAACCCCGGCGGGACTGCGCCAAGGAAAACAAAACGAGAGAGCATGCCCT	237
EL-PB	GAGCAAAAACGAACCCCGGCGGGACTGCGCCAAGGAAAACAAAACGAGAGAGCATGCCCT	240
EL-NK	CGCCGCCCCGGACACGGTGTGCGCGGGGGTGTGCCTTCTTTTAAATAAAGTCTATAACG	292
EL-AB	CGCCGCCCCGGACACGGTGTGCGCGGGGGTGTGCCTTCTTTTAAATAAAGTCTATAACG	297
EL-PB	CGCCGCCCCGGACACGGTGTGCGCGGGGGTGTGCCTTCTTTCAAATAAAGTCTATAACG	300
	*	
EL-NK	ACTCTCGGCAATGGATATCTCGGCTCTCGCATCGATGAAGAACGTAGCGAAATGCGATAC	352
EL-AB	ACTCTCGGCAATGGATATCTCGGCTCTCGCATCGATGAAGAACGTAGCGAAATGCGATAC	357
EL-PB	ACTCTCGGCAATGGATATCTCGGCTCTCGCATCGATGAAGAACGTAGCGAAATGCGATAC	360
EL-NK	TTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGCAAGTTGCGCCCCAA	412
EL-AB	TTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGCAAGTTGCGCCCCAA	417
EL-PB	TTGGTGTGAATTGCAGAATCCCGTGAACCATCGAGTCTTTGAACGCAAGTTGCGCCCCAA	420
EL-NK	GCCGTTAGGCCGAGGGCAGCTCTGCCTGGGTGTACGCATCGTCGCCCCCCCTCCTCCTC	472
EL-AB	GCCGTTAGGCCGAGGGCAGCTCTGCCTGGGTGTACGCATCGTCGCCCCCCCTCCTCCTC	477
EL-PB	GCCGTTAGGCCGAGGGCAGCTCTGCCTGGGTGTACGCATCGTCGCCCCCCCTCCTCCTC	480
EL-NK	GCCTCCCACCCGAGGCGAGCCGGAGCCGCGGGGCGGATACTGGCCTCCCGTGCCTCCCC	532
EL-AB	GCCTCCCACCCGAGGCGAGCCGGAGCCGCGGGGCGGATACTGGCCTCCCGTGCCTCCCC	537
EL-PB	GCCTCCCACCCGAGGCGAGCCGGAGCCGCGGGGCGGATACTGGCCTCCCGTGCCTCCCC	540
	*	
EL-NK	GCTCGCGGTTGGCCCAAATTCGAGTCTCGGCGGCGGTGCGCCGCGACGATCGGTGGCGAA	592
EL-AB	GCTCGCGGTTGGCCCAAATTCGAGTCTCGGCGGCGGTGCGCCGCGACGATCGGTGGCGAA	597
EL-PB	GCTCGCGGTTGGCCCAAATTCGAGTCTCGGCGGCGGTGCGCCGCGACGATCGGTGGCGAA	600
EL-NK	ATTTTCTTTTCATCGAGTTCCTCGCGAGCGCCCGCCCCGGAACGGGGTCTCCTCGGAC	652
EL-AB	ATTTTCTTTTCATCGAGTTCCTCGCGAGCGCCCGCCCCGGAACGGGGTCTCCTCGGAC	657
EL-PB	ATTTTCTTTTCATCGAGTTCCTCGCGAGCGCCCGCCCCGGAACCTTGGGCTCCTCGGAC	660
	* * *	
EL-NK	CCTGATGCGCCGTTTT-CTTCGGCG-TTCGCCTT-GCGACCCAGG--TCAGGCGGGAT	707
EL-AB	CCTGATGCGCCGTTTT-CTTCGGCG-TTCGCCTT-GCGACCCAGG--TCAGG-CGGGAT	711
EL-PB	CCTGATGCGCCGTTTT-CTTCGGCG-TTCGCCTT-TGCGACCCAGGTTTCAGGCGGGAT	720
EL-NK	TACCCG--CTGAGTTT--AAGCAT-ATC--ATAAGGG--GGAGGAA	744
EL-AB	TACCCG--CTGAGTTT--AAGCAT-ATC--ATAAGCC--GGAGGAA	749
EL-PB	TACCCGCGCTGAGTTTTTAAGCATTATTCGATAAAGGCCGGGATGGA	767
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Figure 2
The ITS nucleotide sequence of *E. longifolia* Jack.
(Gray box indicated indels,* indicated SNP)

3.3 Variation of ITS region

An approximately 750 base pair in size of PCR product was obtained from the ITS region amplification from 3 location (EL-NK, EL-AB, and EL-PB). Analysis of sequence alignment of *E. longifolia* Jack showed 98-99% sequence homology (Figure 2). There are 12 single nucleotide polymorphisms (SNPs) and 23 Indels.

4. DISCUSSION AND CONCLUSION

Medicinal plants have been long history used for treatment of various diseases. Correct identification and quality control of the starting material is essential for herbal preparation. However, the major problem of employing medicinal plant for traditional usages is the lack of adequate supporting scientific information on their safety, quality and efficacy. Standardization

is a system to ensure the safety and quality of medicinal product. Determination of pharmacognostic parameters such as determination of ash residues, extractive values, and water content plays a significant role for standardization of medicinal plants. Based on the method recommended by World Health Organization, physicochemical parameters of 15 *E. longifolia* Jack were evaluated as present in Table 1. The water content value was 13.352 ± 0.839 , the less value of water content could prevent the medicinal plant from microorganism growth through storage¹⁶. The ash value used to find out quality and purity of medicinal plant. The total ash and acid insoluble ash value were 1.973 ± 0.039 and 0.696 ± 0.064 respectively. The solubility value of *E. longifolia* Jack in water extraction is higher (4.888 ± 0.376) when compared with ethanolic extraction (1.733 ± 0.178). The extractive values can be estimate the solubility of the chemical constituents in the particular solvent. According to the results, ethanolic extracted of *E. longifolia* Jack showed active antimalarial activity both 3D7 and K1 with the IC₅₀ of 2.16 µg/ml and 1.79 µg/ml respectively, suggesting that this medicinal plant has a potential to be an antimalarial agent when compare to dihydroartemisinin, a standard antimalarial drug. Previous study indicated that extraction of *E. longifolia* showed antimalarial activity against multidrug resistant strain of *Plasmodium falciparum* under *in vitro* conditions¹⁷⁻¹⁸. *E. longifolia* root extracts inhibited six *P. falciparum* isolates at 1.25-5.00 µg/ml¹⁹. However, studies on antimalarial activities of *E. longifolia in vivo* are still limited. Therefore, additional of DNA technologies have been applied for medicinal plant authentication and combinations of various methods have been employed for herbal drug technology development. Identifying of specific regions in the DNA can be used for different level of authentication²⁰ as the genetic composition is unique for each species and is not affected by physical and environmental factors. Further study should be performing for investigation sequence variation in this plant for more information and developing the genetic marker for species identification of their substitution and adulterants materials.

In conclusion, combination of various methods such as standardization, biological activity, and DNA based method is an effective tool for determination the identity of this herbal plants; *E. longifolia* Jack.

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