

Research Article

Molecular identification and proximate composition of wild edible wood growing mushrooms of Mizoram, North East India for food safety and livelihood

VL Thachunglura

Mizoram University, Department of Environmental Science, 796004, Tanhril Aizawl, Mizoram India

John Zothanzama

Mizoram University, Department of Biotechnology, 796004, Tanhril Aizawl, Mizoram, India

Zohmangaiha Chawngthu

Mizoram University, Department of Environmental Science, 796004, Tanhril Aizawl, Mizoram India

Lallawmkima Bochung

Mizoram University, Department of Environmental Science, 796004, Tanhril Aizawl, Mizoram India

R. Vanlalmalsawmi

Mizoram University, Department of Environmental Science, 796004, Tanhril Aizawl, Mizoram India

Prabhat Kumar Rai*

Mizoram University, Department of Environmental Science, 796004, Tanhril Aizawl, Mizoram India

*Corresponding author. E-mail: pkraimzu@gmail.com

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Abstract

Exploring the nutritional attributes of wild edible mushrooms can be vital in addressing global food security. However, there is a dearth of studies on wild edible mushrooms regarding their identification, nutritional status, and molecular characterization in Mizoram, North East India, an integral landscape of the Indo-Burma hotspot region. In the present study, samples of wild edible wood-growing mushrooms were collected from the Mamit and Champhai District of Mizoram. The present study investigated the identities and phylogenetic relationships of the eight fungal species using molecular approaches. The mushrooms were identified up to the species level based on fungal sequences with known identities in GenBank, viz. *Auricularia delicata*, *Laetiporus sulphureus*, *Panus roseus*, *Lentinus squarrosulus*, *Pleurotus cystidiosus*, *Pleurotus djamor*, *Pleurotus pulmonarius* and *Schizophyllum commune*. Further, the identified mushrooms were analysed for macro and micro-nutritional values to assess their suitability for human dietary intake. Results revealed that the collected wild edible mushrooms were rich sources of protein (17.79 - 36.46 g/100 g), carbohydrates (58.14 - 33.77 g/100 g) and fibres (5.63 - 8.32 g/100 g), while the amount of fat (1.93 - 2.77 g/100 g) was low. In addition to bio-molecules, the mushroom samples contained appreciable amounts of essential minerals. Therefore, the selected mushrooms can be used as a potential foodstuff and may also be used as a supplementary protein diet to maintain human health and help achieve food security and sustainable development goals. The present results encourage bio-prospecting, and the need to explore wild edible mushrooms further, especially in the least explored global biodiversity hotspots for sustainable livelihood.

Keywords: Epixylic, Food, Molecular identification, Nutrients, Wild edible mushrooms

INTRODUCTION

Wild edible mushrooms are a promising food source that can address global issues like food security and

sustainability. These wild edible mushrooms can be cultivated in various environments, yielding high yields per unit area by simple utilization of organic waste materials. Their high nutritional properties contribute to

improved health and well-being, combating malnutrition. Cultivation of wild edible mushrooms aligns with UNSDGs such as zero hunger (SDG 2) and health and well-being (SDG 3) (Pandey *et al.*, 2018; El-Ramady *et al.*, 2022). WEM and the fruiting body of higher fungi have been used as a food source by human society since ancient times (Mattila *et al.*, 2001; Karunarathna *et al.*, 2024). There has been an increasing trend in the consumption of wild edible mushrooms due to their rich nutritional value (Gunasekara *et al.*, 2021; Thachunglura *et al.*, 2023), along with both low and high-molecular-weight carbohydrates (Zhou *et al.*, 2016), making them a favored delicacy in numerous countries. Edible mushrooms have been valued for their role in sustaining health, primarily due to their significant protein content, essential amino acids, fibers, carbohydrates, vitamins, and minerals (Li *et al.*, 2021; Wang and Zhao, 2023). Mushrooms provide a range of health benefits, such as antioxidants, antimicrobial properties, anticancer effects, cholesterol regulation, and immune system stimulation (Hyde *et al.*, 2019; Galappaththi *et al.*, 2022; Maaloul *et al.*, 2023). Edible mushrooms, in particular, contain a high level of beneficial compounds like polyphenols, polysaccharides, vitamins, and carotenoids, which make them a great source of antioxidants. Both edible and medicinal mushrooms are highly valued for their natural abundance of antioxidants, which contribute significantly to overall health and vitality (Kozarski *et al.*, 2015; Mwangi *et al.*, 2022; Khumlianlal *et al.*, 2022). Mushrooms are also one of the most significant natural sources of food consumed by the tribal people in India (Agrahar-Murugkar and Subbulakshmi, 2005; Sargunam *et al.*, 2012; Zothanzama *et al.*, 2018; Khumlianlal *et al.*, 2024).

Mizoram, an ecologically relevant state within the Indo-Burma biodiversity hotspot, showcases rich biodiversity, with diverse flora and fauna, including many endemic species (Rai and Lalramnghinglova, 2011; Lalarzovi and Lalnunluanga, 2022). The high humidity during the monsoon season in Mizoram creates an ideal atmospheric environmental condition that supports the abundant growth and diversity of edible macrofungi (Lalrinawmi *et al.*, 2017; Chawngthu *et al.*, 2024; Thachunglura *et al.*, 2024a). In various parts of the world, the study of wild edible mushrooms for their nutritional properties has unveiled diverse, sustainable food sources and health benefits, enriching global dietary knowledge (Bernas *et al.*, 2006; Ouzouni *et al.*, 2009; Atri *et al.*, 2013; Sudheep and Sridhar, 2014; Paloi *et al.*, 2023). Past studies have identified several wild edible mushrooms based on their morphology and molecular characteristics (Lallawmsanga *et al.*, 2016; Zothanzama *et al.*, 2018; Zohmangaiha *et al.*, 2023; Thachunglura *et al.*, 2024b); however, such studies are scant in the context of Mizoram, North East India. To

this end, data and knowledge on the nutritional value of wild edible mushrooms in Mizoram are very scarce, fragmentary, and quiet compared to the higher plants. Most of the frequently used edible wild mushrooms are collected by the marginal section of society and sold in local and city markets (Zothanzama *et al.*, 2018). However, a large number of wild edible mushrooms remain unidentified in Mizoram, and explicit characterization of their nutritional value and reliable identification is urgently required.

In recent times, considering the increasing interest of people in the dietary consumption of mushrooms, the bio-prospecting of mushrooms in the Indo-Burma biodiversity hotspot is extremely essential in terms of their assessment for nutritional quality and suitability for consumption. The reduction in fertile arable land and increased food insecurity raises the marketing value of wood-growing mushrooms and Mizoram is no exception. Therefore, wild edible mushrooms require serious research to provide knowledge and awareness to the people and other stakeholders about mushroom cultivation. In light of these knowledge gaps, the present study aimed to identify the molecular characteristics and determine the proximate chemical composition of Mizoram's wild edible wood-growing mushroom fungal species to assess their nutritive potential.

MATERIALS AND METHODS

Sample collection, transportation and processing

The samples of wood-growing wild edible mushrooms were collected in the fields from Mamit and Champhai Districts of Mizoram. Mamit District lies between 23.8377° N 92.5396° E in northwestern Mizoram. It is characterized by hilly terrain and lush green forests, while Champhai District in the east is marked by its mountainous landscapes, steep slopes, and proximity to the India-Myanmar border, and lies between 23.6357° N, 93.1780° E (Fig. 1). The collected samples were cleaned with a plastic knife to remove forest debris. The collected mushrooms were identified *in situ* if possible and morphological observations such as pileus, lamellae, stipe, spore print and color were assessed from fresh samples (Largent and Stuntz, 1977; Lodge *et al.*, 2004; Zothanzama, 2011) or else the specimens were retained for later identification and photographs were taken in the field. The samples were carefully labelled and kept in air-tight containers before transporting to the laboratory. The fleshy collected mushroom samples were oven-dried at 45° C for 3 days to obtain a constant weight and ground into fine powder for proximate analysis.

Molecular analysis of specimens

DNA was extracted using the Cetyltrimethylammonium

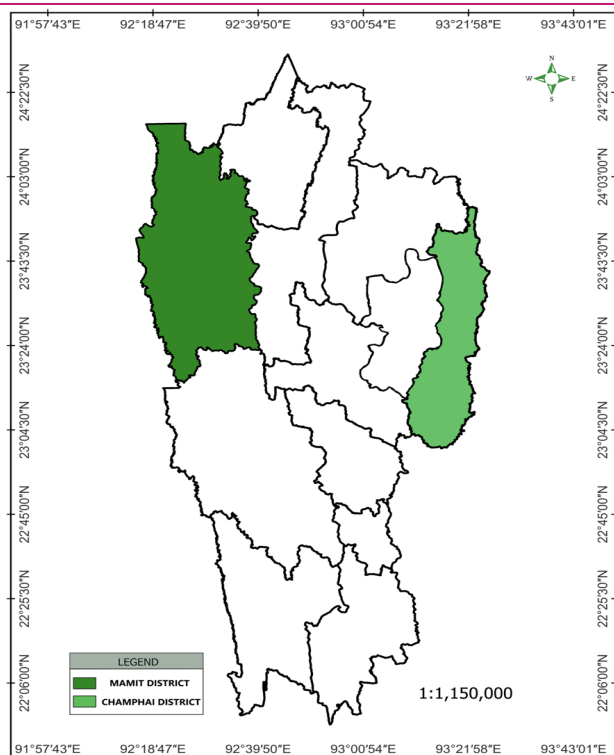


Fig. 1. Map of Mizoram indicating the study site Mamit District and Champhai District

Bromide (CTAB) method (Zothanzama *et al.*, 2018). A small amount of fungal tissue was homogenized in CTAB lysis buffer, incubated at 65°C for cell lysis, and purified with chloroform. The supernatant was transferred to a new tube, mixed with isopropanol, and incubated at -20°C to precipitate DNA. After centrifugation, the DNA pellet was washed with 70% ethanol, air-dried, and resuspended in 100 µL of sterile water.

Polymerase Chain Reaction (PCR) amplification targeted the Internal transcribed spacer (ITS) region using primers ITS1-F and ITS4 (White *et al.*, 1990). Reactions (25.5 µL) contained GoTaq Green MasterMix (Promega), primers (1 µL each, 5 µM), template DNA (1 µL), and nuclease-free water. Thermocycling conditions included initial denaturation (94°C, 5 min), 35 cycles of denaturation (94°C, 1 minute), annealing (52°C, 1 minute), extension (72°C, 1 minute), and final elongation (72°C, 5 minute). Amplicons were visualized via agarose gel electrophoresis and sequenced bidirectionally using Sanger sequencing. Consensus sequences were aligned in BioEdit (Hall, 1999) and compared to GenBank entries via BLASTn (Altschul *et al.*, 1990). Phylogenetic trees were constructed in rAxML GUI using ClustalW-aligned sequences (Larkin *et al.*, 2007).

Phylogenetic analysis

Phylogenetic analysis of the eight fungal isolates was performed using ITS gene data with Maximum Likelihood (ML) and Neighbor-Joining (NJ) in rAxML GUI. Alignment gaps were treated as missing data, and NJ

trees were based on total character differences with 1,000 bootstrap replicates.

Analytical methods

Moisture, protein, fat, crude fiber and ash content of the identified mushroom samples were determined on a dry weight basis using the standard methods of the Association of Official Analytical Chemists (AOAC, 2000). The moisture content was determined by drying the sample at 105 °C, and 2 g was taken in a porcelain crucible and placed into a hot air oven (HAO) at 105 °C for 3 h. The sample was cooled in a desiccator, and the difference between the sample weight before and after drying was used to measure the moisture content percentage. Total nitrogen (N) was determined by the microKjeldahl method, and protein content was calculated as total N × 6.25. Fat content was determined by extracting the sample using the Soxhlet apparatus with petroleum ether for 14 h; crude fibre was determined by acid and alkali digestion. The ash content was determined by incinerating the known weight of samples in porcelain crucibles in a muffle furnace at 550 °C for 6 h.

The total carbohydrate was calculated by difference (total carbohydrates (%) = 100 - [moisture (%) + protein (%) + fiber (%) + fat (%) + ash (%)] (Raghuramulu *et al.*, 2003) and energy value was calculated by following the equation given by Crisan and Sands (1978) conversion factors: [Energy value (kcal/100 g) = (2.62 × % protein) + (8.37 × % fat) + (4.2 × % carbohydrate)] Eq.1.

Minerals such as calcium (Ca), iron (Fe), magnesium (Mg), manganese (Mn), potassium (K), sodium (Na) and zinc (Zn) were evaluated using an Atomic absorption spectrophotometer (Shimadzu AA-7000; Shimadzu Corporation, Japan) after dry ashing of the sample (Isildak *et al.*, 2004) with detection of limits ranges between 0.01 to 0.09 ppm.

Statistical analysis

The analysis of all the mushrooms was done in triplicate on a dry weight basis. All the experimental results were the mean of three replicates, and the data were expressed as Mean ± standard deviation. Statistical analysis was conducted using the GraphPad Prism 5.0 (GraphPad Software, Inc., USA). Statistical analysis was done by one-way analysis of variance (ANOVA) with Tukey's multiple comparisons test (P < 0.05).

RESULTS AND DISCUSSION

Identification of specimen

The nucleotide sequences of the mushrooms blasted against sequences from GenBank database revealed the identification of eight representative fungal isolates

Table 1. Showing list of mushroom species, their collection site, family, and order

Species	Collection Site	Family	Order
<i>Auricularia delicata</i> (Mont. ex Fr.) Henn.	Mamit District	Auriculariaceae	Auriculariales
<i>Laetiporus sulphureus</i> (Bull.) Murrill	Champhai District	Laetiporaceae	Polyporales
<i>Panus roseus</i> (Karun., K.D. Hyde & Zhu L. Yang) N. Vinjusha & T.K.A. Kumar	Mamit District	Panaceae	Polyporales
<i>Lentinus squarrosulus</i> Mont. Singer	Mamit District	Polyporaceae	Polyporales
<i>Pleurotus cystidiosus</i> O.K. Mill	Mamit District	Pleurotaceae	Agaricales
<i>Pleurotus djamor</i> (Rumph. ex Fr.) Boedijn	Mamit District	Pleurotaceae	Agaricales
<i>Pleurotus pulmonarius</i> (Fr.) Quél.	Champhai District	Pleurotaceae	Agaricales
<i>Schizophyllum commune</i> Fr.	Champhai District	Schizophyllaceae	Agaricales

from Mizoram. According to the blast, samples mostly resembled with *Auricularia delicata* (JZT-VL/016), *Laetiporus sulphureus* (JZT-VL/011), *Panus roseus* (JZT-VL/012), *Lentinus squarrosulus* (JZT-VL/018), *Pleurotus cystidiosus* (JZT-VL/019) *Pleurotus djamor* (JZT-VL/023), *Pleurotus pulmonarius* (JZT-VL/009) and *Schizophyllum commune* (JZT-VL/007). The fruiting bodies of the collected sample are represented in Fig. 2, and their phylogenetic tree is given in Fig. 3.

Among the identified species, some species (*A. delicata*, *La. sulphureus*, *L. squarrosulus*, *Pl. djamor*, *Pl. pulmonarius*, and *S. commune*) are recognized as having high nutritional value and are widely consumed in various parts of the world (Mortimer *et al.*, 2014; Bandara *et al.*, 2019; Torres-Martínez *et al.*, 2022; Wang *et al.*, 2023). However, it appears that many of these species are not traditionally consumed by the people of Mizoram. On the other hand, *L. squarrosulus* and *S. commune* are only known to be commonly consumed by the indigenous people of Mizoram. Species like *A. delicata*, *P. roseus* and *Pl. pulmonarius* are consumed only by a small population in the region. Limited knowledge of mycology restricts the consumption of wild edible mushrooms in Mizoram (Zothanzama *et al.*, 2018). The collection site and their respective family and order are mentioned in Table 1.

The ITS1-5.8S-ITS2 DNA sequences of the eight fungal isolates were compared with 34 reference sequences of fungal species from the database. The

species, voucher numbers, GenBank accession numbers, and localities of the 34 corresponding sequences used for the analysis to construct the phylogenetic tree are presented in Table 2. In the Maximum Likelihood tree generated, the specimens of the different species of fungi clustered with their related species with high support values. Clade A included *La. sulphureus* alongside the specimen JZT-VL/011. Clade B comprised *Lentinus* and *Panus* species, which were closely related in the phylogenetic tree. Clade C was formed by *A. delicata*, while clade D contained *Pleurotus* species. Clade E grouped *S. commune* with the present species, exhibiting strong support. *Amanita spissacea* was used as an outgroup. Blast search showed that each isolate is <96% identity with query coverage of <96%.

Proximate composition of wild edible mushrooms

The results of the nutritive value of selected wild edible wood-growing mushrooms are shown in Table 3. In this context, it can be noted that the growth characteristics, stage of development and post-harvest condition of edible mushrooms may influence their nutritional value (Klomklung *et al.*, 2014; Valverde *et al.*, 2015; Dawadi *et al.*, 2022).

Edible mushrooms had approximately 90% of moisture and 10% of dry matter, while fresh and dried mushrooms had approximately 90% of dry matter and 10% of moisture (Kurtzman, 1975; Crisan and Sands,

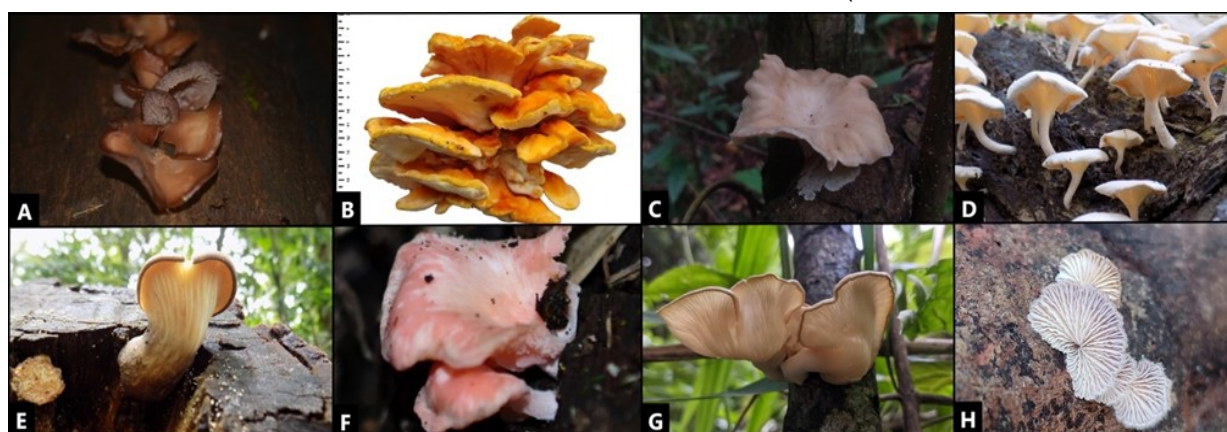


Fig. 2. Fruiting body of (A) *Auricularia delicata* (B) *Laetiporus sulphureus* (C) *Panus roseus* (D) *Lentinus squarrosulus* (E) *Pleurotus cystidiosus* (F) *Pleurotus djamor* (G) *Pleurotus pulmonarius* (H) *Schizophyllum commune*

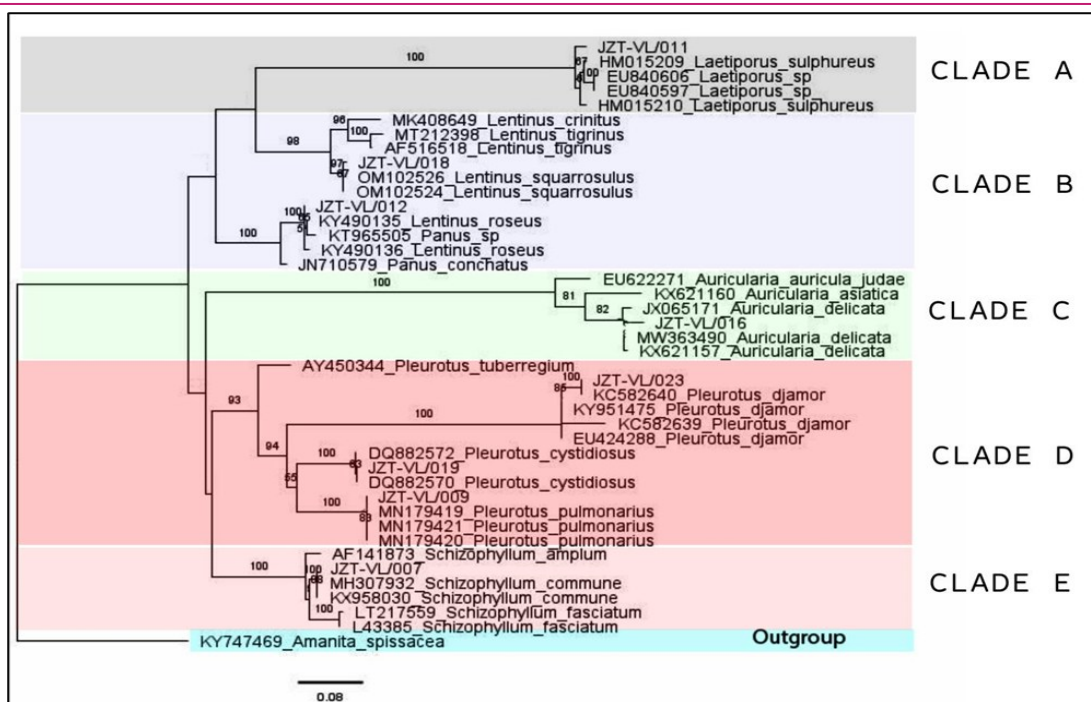


Fig. 3. Phylogenetic tree - Evolutionary history inferred by using the Maximum Likelihood method based on the GTR-GAMMA model. Tree is drawn to scale, with branch lengths measured in the number of substitutions per site. Analysis involved 42 nucleotide sequences. Evolutionary analyses were conducted in raxmlGUI 2.0

1978). Moisture content is vital to sustain mushrooms' physical and chemical properties (Shams *et al.*, 2022). There was considerable variation in the moisture content of the mushrooms analysed in the present study, ranging from 8.06% in *S. commune* to 12.39% in *A. delicata*.

The protein contents of the studied wild edible mushrooms were generally high and varied from 17.79 g/100 g in *A. delicata* to 36.46 g/100 g in *P. roseus*. Particularly, *P. roseus*, *L. squarrosulus*, *Pl. cystidiosus*, *Pl. djamor* and *Pl. pulmonarius* exhibited protein content ranging from 32.96 to 34.12 g/100g, which is consistent with the 19–35% protein range reported by several authors (Chang and Buswell, 1996; Rugolo *et al.*, 2023; Sousa *et al.*, 2023). Analysis of *La. sulphureus* by Teke *et al.* (2021) showed 8.62 g/100 g crude protein, which is substantially low in comparison to the protein content (21.29 g/100 g) obtained in the Indian sample of *La. sulphureus*. However, it aligns with the report by Saha *et al.* (2014) on *La. sulphureus* (21 g/100 g). It is known that protein content varies widely according to the type of mushrooms, the level of nutrient nitrogen and location (Bernas *et al.*, 2006; Kaur *et al.*, 2022).

Fats contents were relatively low in the studied mushrooms, ranging from 1.93 g/100 g in *S. commune* to 2.77 g/100 g in *Pl. pulmonarius*. Interestingly, the fat content values of *L. squarrosulus* (2.67 g/100 g) in the present study were lower when compared with the study of Nwanze *et al.* (2006), who noted considerably higher fat content in *L. squarrosulus* stipe (6.01 g/100

g) and pileus (6.56 g/100 g). However, the present results of *S. commune* (1.21 g/100 g) and *L. squarrosulus* (2.31 g/100 g) fat contents are consistent with Gunasekara *et al.* (2021) for *L. squarrosulus* (2.31 g/100 g) and *S. commune* (1.21 g/100 g). Mushrooms are known to be low in fats, and the lower fat content values of the present study were also in agreement with several studies on different mushroom species (Agrahar-Murugkar and Subbulakshmi, 2005; Pushpa and Purushothama, 2010; Lallawmsanga *et al.*, 2016). Mushrooms, characterized by their low fat content and abundant essential unsaturated fatty acids, are ideal for the human diet and health (Nakalembe *et al.*, 2015). Maximum value of fiber was recorded for *L. squarrosulus* (8.32 g/100 g), while minimum was noted in the case of *S. commune* (5.63 g/100 g). These results on fiber contents were in agreement with earlier reports of Silva *et al.* (2002) for *Pl. pulmonarius*, Rai and Sohi (1988) for *Lentinus* and *Panus* species, Raman *et al.* (2020) for *Pl. djamor* and Luangharn *et al.* (2014) for *La. sulphureus*. Fibers in mushrooms are vital as they can promote healthy digestion, aid in maintaining a healthy weight, and support heart health. The ash content of the mushrooms consists of unburnable inorganic salts, with phosphorus and potassium being the main components. However, ash content does not provide direct information about the macronutrient value of mushrooms, such as proteins, fats and carbohydrates (Mattila *et al.*, 2001; Ouzouni *et al.*, 2009; Sifat *et al.*, 2020). It is an indication of the mineral content of the mushrooms. The ash content

Table 2. List of the mushroom species used for phylogenetic analysis, along with their voucher numbers, GenBank accession numbers, and localities

No	Voucher	Accession Number	Species Name	Locality
1	FUM-093	KY951475	<i>Pleurotus djamor</i>	Malaysia
2	ABM1049204	KC582640	<i>Pleurotus djamor</i> var. <i>roseus</i>	Malaysia
3	ABM1049203	KC582639	<i>Pleurotus djamor</i> var. <i>roseus</i>	Malaysia
4	CBS 665.85	EU424288	<i>Pleurotus djamor</i>	China
5	OR1214	KY747469	<i>Amanita cf. spissacea</i>	Belgium
6	A.S-E 2018-03-04.1	MH307932	<i>Schizophyllum commune</i>	Luxembourg
7	15R-5-F01	KX958030	<i>Schizophyllum commune</i>	China
8	CBS267.60.	LT217559	<i>Schizophyllum fasciatum</i>	Spain
9	CBS 267.60	L43385	<i>Schizophyllum fasciatum</i>	Mexico
10		AF141873	<i>Schizophyllum amplum</i>	Sweden
11	Bla0 clone A 1	DQ882570	<i>Pleurotus cystidiosus</i>	Japan
12	S396 clone A	DQ882572	<i>Pleurotus cystidiosus</i>	Japan
13		AY450344	<i>Pleurotus tuberregium</i>	USA
14	LE(BIN)0861 SBI	AF516518	<i>Lentinus tigrinus</i>	USA
15	SR1352	MT212398	<i>Lentinus tigrinus</i>	India
16	LSQBot	OM102524	<i>Lentinus squarrosulus</i>	Philippines
17	LSQOs	OM102526	<i>Lentinus squarrosulus</i>	Philippines
18	EF58b	MK408649	<i>Lentinus crinitus</i>	Brazil
19	HKAS 94714	KY490136	<i>Lentinus roseus</i>	China
20	HKAS 94715	KY490135	<i>Lentinus roseus</i>	China
21	X1234	JN710579	<i>Panus conchatus</i>	Finland
22	FBD167	KT965505	<i>Panus</i> sp.	Vietnam
23	CNSBlitz0098	JX065171	<i>Auricularia delicata</i>	USA
24	clone rlc-526	MW363490	<i>Auricularia delicata</i>	China
25	MFLU162117	KX621157	<i>Auricularia delicata</i>	Thailand
26	BBH895	KX621160	<i>Auricularia asiatica</i>	Thailand
27	NW486	EU622271	<i>Auricularia auricula-judae</i>	China
28	NSK 1014215	MN179421	<i>Pleurotus pulmonarius</i>	Russia
29	NSK 1014280	MN179420	<i>Pleurotus pulmonarius</i>	Russia
30	NSK 1014279	MN179419	<i>Pleurotus pulmonarius</i>	Russia
31	LAE10-LS04	HM015210	<i>Laetiporus sulphureus</i>	Poland
32	LAE09-LS03	HM015209	<i>Laetiporus sulphureus</i>	Poland
33	KATRIN-2	EU840606	<i>Laetiporus</i> sp.	Sweden
34	RVP2	EU840597	<i>Laetiporus</i> sp.	Sweden

was highest in *P. roseus* (7.74 g/100 g) and lowest in *Pl. cystidiosus* (5.84 g/100 g). The recorded values of ash content were similar to the report of Khan *et al.* (2013) for *Pl. djamor*, Okwulehie *et al.* (2014) for *Pl. pulmonarius*, Ao and Deb (2019) for *S. commune* and *La. sulphureus*. Wide variation in ash content of the present investigation might be due to the fact that the relatively young mushrooms' fruiting bodies were noted to have higher ash content than the mature ones (Kalmis *et al.*, 2011).

Carbohydrate content was found to be abundant in all tested samples, constituting mushroom's main nutritional value, ranging from 33.77 g/100 g in *P. roseus* to 58.14 g/100 g in *S. commune*. In the present nutritional analysis on carbohydrate content, the results of *S. commune* was reasonably higher as compared with previous studies of Kumar *et al.* (2013) for *S. commune* (32.43 g/100 g), while it was quite similar to those reported by Rampinelli *et al.* (2010) for *Pl. djamor* and Vishwakarma *et al.* (2017) for *Pl. cystidiosus*.

Table 3. Showing proximate composition of wild edible wood growing mushrooms on dry weight basis

Mushroom Species	Moisture (g/100 g)	Protein (g/100 g)	Fat (g/100 g)	Fiber (g/100 g)	Ash (g/100 g)	CHO (g/100 g)	Energy Kcal/100 g
<i>Auricularia delicata</i>	12.39 ± 0.18 ^a	17.79 ± 1.01 ^c	1.96 ± 0.16 ^b	6.52 ± 0.20 ^{abc}	6.31 ± 0.07 ^{bc}	54.73 ± 1.46 ^a	293.3 ± 2.53 ^{ab}
<i>Laetiporus sulphureus</i>	9.25 ± 0.19 ^c	21.29 ± 0.50 ^c	2.05 ± 0.14 ^{ab}	7.01 ± 0.32 ^{abc}	6.58 ± 0.31 ^{bc}	54.17 ± 0.53 ^a	300.5 ± 1.96 ^a
<i>Panus roseus</i>	11.83 ± 0.12 ^{ab}	36.46 ± 0.50 ^a	2.20 ± 0.09 ^{ab}	7.99 ± 0.34 ^a	7.74 ± 0.18 ^a	33.77 ± 0.56 ^c	255.8 ± 1.49 ^d
<i>Lentinus squarrosulus</i>	10.24 ± 0.11 ^{bc}	34.10 ± 0.87 ^{ab}	2.67 ± 0.17 ^{ab}	8.32 ± 0.54 ^a	7.18 ± 0.14 ^{ab}	36.58 ± 1.00 ^c	268.7 ± 4.67 ^{cd}
<i>Pleurotus cystidiosus</i>	10.19 ± 0.11 ^{bc}	33.54 ± 1.34 ^{ab}	2.15 ± 0.19 ^{ab}	6.03 ± 0.19 ^b	5.84 ± 0.12 ^c	42.24 ± 1.14 ^b	278.0 ± 9.35 ^{bc}
<i>Pleurotus djamar</i>	10.21 ± 0.04 ^{bc}	32.96 ± 0.50 ^b	2.71 ± 0.16 ^{ab}	8.29 ± 0.24 ^a	7.04 ± 0.41 ^{ab}	38.35 ± 0.43 ^{bc}	270.1 ± 3.45 ^{cd}
<i>Pleurotus pulmonarius</i>	11.36 ± 1.11 ^{ab}	34.12 ± 0.88 ^{ab}	2.77 ± 0.13 ^a	8.21 ± 0.14 ^a	5.98 ± 0.12 ^{bc}	38.19 ± 0.64 ^{bc}	272.8 ± 1.36 ^{bcd}
<i>Schizophyllum commune</i>	8.06 ± 0.05 ^d	19.59 ± 0.47 ^c	1.93 ± 0.21 ^b	5.63 ± 0.67 ^c	6.99 ± 0.19 ^{ab}	58.14 ± 1.45 ^a	310.7 ± 2.53 ^a

Each value is expressed in mean ± SD, (n = 3); In each column, different letters mean significant differences between species (p < 0.05)

Therefore, wild edible mushrooms are a good source of carbohydrates and their contents vary greatly and generally range from 28.38 g/100 g to 84.48 g/100g on a dry weight basis (Wang *et al.*, 2001; Pushpa and Purushothama, 2010; Shin *et al.*, 2007). Carbohydrates are an essential part of a healthy diet and provide an important energy source for the human body. Energy values ranged from 255.8 Kcal/100 g in *P. roseus* to 310.7 Kcal/100 g in *S. commune*. This was in agreement with the results of Manjunathan and Kaviyaran (2010) and Nakalembe *et al.* (2015). The average nutritional values across eight samples indicated 10.44 ± 1.4 (g/100 g) of moisture, 26.34 ± 12.46 (g/100 g) of proteins, 2.31 ± 0.35 (g/100 g) of fat, 7.25 ± 1.10 (g/100 g) of fibre, 6.71 ± 0.65 (g/100 g) of ash, 44.52 ± 9.60 (g/100 g) of carbohydrates, and an energy value of 281.24 ± 18.5 (Kcal/100 g).

Table 4 presents the mineral content of eight wild edible mushrooms. Among these minerals, potassium was identified as the most abundant, followed by iron and calcium, while manganese was the least abundant mineral found in the mushrooms. *L. squarrosulus* exhibited the highest potassium (1925.50 mg/100 g), calcium (221.22 mg/kg), iron (216.91 mg/kg), magnesium (115.62 mg/kg), and zinc (221.61 mg/kg) levels, while *A. delicata* recorded the lowest contents of potassium (1204.09 mg/100 g), calcium (57.44 mg/kg), iron (65.06 mg/kg) and magnesium (23.97 mg/kg). Regarding zinc, *S. commune* had the lowest content (36.97 mg/kg). Lastly, *S. commune* showed the highest sodium (215.58 mg/kg) and manganese (23.37 mg/kg) levels, while *P. roseus* had the lowest sodium (115.83 mg/kg) content, and *Pl. cystidiosus* had the lowest manganese content (6.25 mg/kg). The analyzed mushrooms fulfil a significant portion of the daily requirements for various essential minerals, aligning with findings from several studies in the literature (Isildak *et al.*, 2004; Genççelep *et al.*, 2009; Teke *et al.*, 2021).

The mineral contents in wild edible mushrooms can vary significantly due to several factors, including the species of mushroom, the environmental conditions in which they are grown, and the substrate they feed on (Liu *et al.*, 2022; Mleczek *et al.*, 2024). Nutrient availability in the soil, climate, and the presence of specific microorganisms can all influence the mineral uptake and accumulation in mushrooms. Moreover, harvesting methods and post-harvest processing may also affect the final mineral content of these fungi, highlighting the importance of understanding the ecological context in which wild mushrooms grow. To our knowledge, the present study provides an initial insight into the nutritional chemistry of wild edible mushrooms in a global biodiversity hotspot. Elucidating the nutritional value of wild edible mushrooms bears a tight interrelationship with food security and the livelihood of local indigenous people.

Table 4. Showing contents of essential minerals in wild edible mushrooms (mg/kg or mg/100g in dw)

Mushroom Species	Ca (mg/kg)	Iron (mg/kg)	Magnesium (mg/kg)	Manganese (mg/kg)	Potassium (mg/100g)	Sodium (mg/kg)	Zinc (mg/kg)
<i>Auricularia delicata</i>	57.44 ±0.98 ^h	65.06 ±0.878 ^h	23.97 ±0.37 ^h	22.78 ±0.58 ^{ab}	1204.09 ±3.33 ^h	179.28 ±0.73 ^b	75.68 ±0.57 ^g
<i>Laetiporus sulphureus</i>	179.61 ±0.77 ^c	203.74 ±0.64 ^b	45.74 ±0.39 ^d	15.96 ±0.64 ^d	1752.13 ±4.07 ^c	157.58 ±0.68 ^d	165.42 ±0.97 ^c
<i>Panus roseus</i>	98.78 ±0.41 ^f	114.82 ±0.83 ^f	38.93 ±0.45 ^f	14.45 ±0.23 ^{ef}	1252.33 ±1.43 ^g	115.83 ±0.39 ^h	181.95 ±0.82 ^b
<i>Lentinus squarrosulus</i>	221.22 ±0.72 ^a	216.91 ±1.28 ^a	115.62 ±0.84 ^a	22.99 ±0.44 ^{ab}	1925.50 ±0.7 ^a	148.12 ±0.68 ^e	221.61 ±1.07 ^a
<i>Pleurotus cystidiosus</i>	195.16 ±0.92 ^b	174.04 ±0.66 ^c	57.07 ±0.79 ^b	6.25 ±0.27 ^g	1457.28 ±1.32 ^e	124.19 ±0.29 ^g	134.95 ±0.29 ^e
<i>Pleurotus djamor</i>	144.43 ±0.68 ^e	157.06 ±0.65 ^d	48.11 ±0.74 ^e	13.29 ±0.56 ^f	1422.68 ±0.68 ^f	145.6 ±0.64 ^f	162.1 ±0.29 ^d
<i>Pleurotus pulmonarius</i>	163.08 ±0.78 ^d	143.34 ±0.55 ^e	53.4 ±0.24 ^c	21.95 ±0.38 ^b	1821.74 ±1.4 ^b	166.25 ±0.82 ^c	85.41 ±1.18 ^f
<i>Schizophyllum commune</i>	78.41 ±0.95 ^g	70.22 ±0.88 ^g	34.47 ±0.71 ^g	23.37 ±0.55 ^a	1642.39 ±2.22 ^d	215.58 ±0.84 ^a	36.97 ±0.63 ^h

Each value is expressed in mean ± SD, (n = 3); In each column, different letters mean significant differences between species (p < 0.05)

Conclusion

Molecular analysis in the present study confirmed the presence of eight wild edible mushroom species in Mizoram. The study also evaluated the nutritional chemistry and phylogenetic insight of eight species of wild edible mushrooms belonging to five different families. The wild edible mushrooms were observed to be an excellent source of macronutrients and micronutrients. These mushrooms were found to be rich in macronutrients and micronutrients, making them a valuable source of human nutrition. Additionally, they support immune health and contribute to food security by addressing malnutrition and diversifying diets, especially in regions with limited food availability. More research is warranted on the antioxidant properties of the eight wild edible mushrooms to provide a better understanding of their potential health benefits and food safety implications. This will help expand the knowledge of wild edible mushrooms and increase their market value for food security/-safety.

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Conflict of interest

The authors declare that they have no conflict of interest.

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