



Assessment of the anti-leukemic and antioxidant potential of the methanol extract of a wild, edible, and novel mushroom, *Astraeus hygrometricus*, and unraveling its metabolomic profile

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ABSTRACT

Mushrooms are enriched with a plethora of bioactive molecules that play a vital role in the prevention of human diseases. The balance between ROS generation and cancer growth is one of the main prerequisites for efficient cancer treatment. In this study, to testify the aforesaid theory, five wild edible mushrooms were initially screened based on their anti-proliferative efficiency, and the best mushroom extract was selected for the assessment of their antioxidant potentialities *in vitro* in various artificially generated free radicals such as DPPH, ABTS+, and by FRAP experiment. The reason behind the antiproliferative potentiality and antioxidant capability of the most potent extract was also correlated by profiling its metabolites through GC-MS analysis. The study reveals, that the methanolic extract of *Astraeus hygrometricus* is the most potent anti-leukemic extract (IC₅₀ 22.7 ± 0.23 µg/mL) followed by *Serpula* sp. (75.7 ± 0.44 µg/mL), *Phallus* sp. (60.53 ± 0.36 µg/mL), *Tricholoma* sp. (53.76 ± 0.46 µg/mL), *Lentinus* sp. (58 ± 0.13 µg/mL) against the Jurkat cell line with negligible effect on normal PBMC cells isolated from healthy donors. The assessment of the antioxidant profile *Astraeus hygrometricus* reveals its moderate antioxidant efficacy against several artificially generated free radicals, such as DPPH (76.9 ± 0.16 µg/mL), ABTS+ (142 ± 0.66 µg/mL) and moderate iron chelating efficacy (32.37 ± 2.31 µM). The GC-MS analysis of both methanol and ethyl acetate extracts of *Astraeus hygrometricus* have found 53 and 52 compounds respectively, with wide diverse ranges of chemically classified biomolecules such as alkane, alcohol, fatty acid, organic acid, polycyclic and heterocyclic compounds, amino acid, vitamin, and hormone, etc. with a wide array of biological activity such as anticancer and antioxidant potentiality. In conclusion, it can be said that these wild edible mushroom *Astraeus hygrometricus* are a repository of novel biomolecules that can be used in the treatment of Leukemia in the future.

INTRODUCTION

The basic prerequisite for all living organisms is energy which is produced by the oxidation process, and the generation of reactive oxygen species (ROS) as a byproduct is an inevitable outcome of it. Though the indigenous antioxidant mechanism to counter

oxidative stress is present but the continuous production of these uncontrolled free radicals *in vivo* may lead to various health disorders and even lead to developing variously non-communicable diseases like cancer [1–3]. Today's cancer treatment is mostly confined to the use of different combinations of chemotherapeutic drugs, but most of the commercially

available chemotherapeutic drugs show various degrees of side effects, most commonly the overproduction of ROS, and thus leads to oxidative stress. To counteract the negative consequences of the phenomenon, the various synthetic antioxidant molecules such as butylated hydroxyanisole (BHA), butylated hydroxytoluene (BHT), *tert*-butylated hydroxyquinine (TBHQ), etc. have been used as an adjuvant in chemotherapy, but these synthetic antioxidant drugs in most the cases show various side effects [4,5]. So, to allay the negative consequences of the existing drugs, there is an urgent need for the development of safe anticancer and antioxidant exogenous drugs preferably from natural sources.

In recent years, the mushroom-derived foods, as it contains a wide range of secondary metabolites have attracted attention as a natural reservoir of effective commercial antioxidant and anticancer molecules [6-8]. Mushroom contains a wide variety of bioactive compounds such as diverse phenol and flavonoid molecule, polysaccharides, glycosides, tocopherols, carotenoids, alkaloids, volatile oils, organic acids and has the potentiality to serve as an alternative reservoir for naturally occurring antioxidants, anti-inflammatory, cardiovascular, anti-microbial, immunomodulation, anticancer, and anti-diabetic drug [3,9-11]. Low molecular weight substances isolated from wood degrading fungus *Cerrena unicolor* has showed dual efficiency, as anticancer molecule towards two breast cancer cell lines MDA-MB-231, MCF7 and one prostatic carcinoma cells PC3 with IC₅₀ value less than 7 mg/mL and also as antioxidant molecule against artificially generated free radicle 1,1-diphenyl-2-picryl-hydrazyl, with promising half-maximal inhibitory concentration ranging in between 20.39 µg/mL to 64.14 µg/mL [12]. Another impotent β-Glucan, lentinan isolated from edible mushroom *Lentinus edodes*, also used as both anticancer and antioxidant agent [13].

Profiling of total primary and secondary metabolome has been considered as one of the reliable methods to investigate primarily the possible group of molecules responsible for the desired bioactivity. In order to look at the metabolomic profile, GC-MS method has been widely used to detect several abundant molecules such as steroids, essential oil, triterpenoid, diterpenoid, phytosterol group of compounds etc. that are responsible for bioactivity of any extract are individually detected reliably, even they are present in heterogenous extract [14]. The metabolomic profiling, through the GC-MS analysis of methanolic extract of a wild mushroom *Phallus* sp. has revealed presence of

different derivatives of anticancer sterol ergo sterol thus directly point out its anti-cancer efficacy of *Phallus* sp. [4].

Therefore, the present study was carried out to compare the anticancer potentiality between methanol and ethyl acetate extracts of five wild edible mushrooms *Astraeus hygrometricus*, *Phallus* sp., *Lentinus* sp., *Tricholoma* sp., and *Serpula* sp. and the most effective extract of best mushroom was selected for antioxidant assessment to investigate the relationship between anti-cancer activity and ROS management *in vitro*. Consequently, to point out the probable bioactive components responsible behind this anticancer and antioxidant efficacies, the metabolomics profile of potent extracts also conducted.

MATERIALS AND METHODS

Sample collection and identification

Mushroom fruit bodies were collected from various locations of West Bengal, the eastern state of India in between the months of May to August between 2015 – 2019. The mature fruit body of *Phallus* sp. was collected from Lolegaon (27.0206°N, 88.5650°E) Kalimpong District positioned under Himalayan foothills, *Tricholoma* sp. and *Serpula* sp. from Mathurapur (22.1203° N, 88.3943° E) hinterland of Bay of Bengal South 24-Parganas District, *Astraeus hygrometricus* from Jamboni (22.4502° N, 86.8998° E), *Lentinus* sp. From Sigram (22.4479° N, 86.8967° E) belonged to the laterite region of Jhargram District (Figure 1a-e). All possible morpho-organoleptic characteristics such as color, size, and the shape were tested immediately after collection.

Identification of the mushroom

Regarding the identification of mushrooms, both classical methods (morphological and anatomical methods) and molecular identification methods were applied. *Astraeus hygrometricus*, *Serpula* sp., was identified by Prof. Krishnendu Acharya and *Lentinus* sp. was identified by Prof. Santanu Paul with the consultation of various standard published keys and the specimen sample was preserved in Calcutta University Herbarium with voucher number CUH/AM/681 (*Astraeus hygrometricus*) and CUH/FN/JH/SP/04 (*Lentinus* sp.). Identification of the other two samples was validated by molecular identification, by amplifying its conserved internal

transcribed spacer regions (ITS1, 2, 5.8s rDNA) along with conventional identification methods.

Molecular identification of the mushroom-

DNA extraction- Genomic DNA of the mushroom was extracted from air-dried samples in accordance with the manufacturer protocol of fungal gDNA Mini Kit (Xcelris Genomics, Ahmedabad, India).

Polymerase Chain Reaction (PCR) -Amplification of its specific Internal transcribed spacer regions (ITS1, 2, 5.8s rDNA), was done by selecting a pair of primer sequence ITS1 (forward primer 5' TCC GTA GGT GAA CCT GCGG 3') and ITS4 (reverse primer 5' TCC TCC GCT TAT TGA TAT GC3') [15]. The PCR reaction mixture was prepared as per the method described by Ray *et.al.* 2020[4]. The amplification reaction was designed as initial denaturation for 4 min at 94°C followed by 35 cycles consisting of 1 minute at 94°C (denaturation) then 1 min in 56°C for annealing, 1 min at 72°C (extension), and finally one cycle for 10 min at 72°C for final elongation [4].

Sequencing- Automated DNA sequencing method, performed on an ABI3730XL-15104-028DNA Analyzer (Applied Bio systems, USA) was used for the sequencing of the ITS regions. The same sets of primers identical with amplicons for the ITS rDNA region were used for the sequencing reaction. The newly generated sequences were deposited in GenBank (www.ncbi.nlm.nih.gov).

Preparation of mushroom extracts

The collected mushroom samples were dried and ground to powder. Three types of solvents hexane, ethyl acetate, and methanol were selected based on the polarity for the preparation of mushroom extracts [8,16]. In brief, initially, 50 g of mushroom powder was taken and then percolated in hexane for three days to remove the fatty substances and the hexane extract was prepared through filtration followed by evaporation in a rotary evaporator. Repetition of the same procedure was performed to prepare the Ethyl Acetate (EAE) and Methanol Extract (ME) respectively. All the extracts were stored at 4°C for further experiments. The methanol extract was selected for further evaluation of the antioxidant and antiproliferative properties of selected mushrooms.

Chemicals

All chemicals that were used for the experiments were analytical grade and freshly prepared. DMSO, Gallic acid (Merck), Sodium Carbonate (Merck), FolinCiocalteau (Merck), Quercetin (SRL), Sodium nitrite (Himedia), Aluminium chloride (Merck), Sodium hydroxide (SRL), 2, 2-diphenyl-1-picrylhydrazyl (DPPH) (SRL), Methanol (Merck), Acetic acid (Merck), Thiobarbituric acid, n-Butanol (Himedia), ABTS, FeSO₄·7H₂O (MERK), acetate buffer, Sodium phosphate buffer, Phosphate buffer.

Cell line and cell culture

In order to assay the efficacy of the anti-proliferative potentiality of the crude extracts, one leukemic cell line, Jurkat (T Acute Lymphoblastic Leukemia) was selected and the cell line was maintained in the RPMI medium supplemented with 10% FBS, 100U/mL penicillin, and 100 U/mL streptomycin incubated at 37°C in a humidified atmosphere containing 5% CO₂.

MTT assay

The anti-proliferative effect of both of the crude extracts ME and EAE against Jurkat was determined by the MTT dye uptake method in accordance with Gopal *et.al.* 2014 [17]. Briefly, 1X10⁴ cells were seeded in a 96-well plate with various concentrations of mushroom extracts 10, 25, 50 100 µg/mL for 24 hours at 37°C. Thereafter, 25µL MTT solution (5 mg/mL in PBS) was added to each well and the optical density (OD) was optically measured at 590. To measure 100% lysis of cell, after the incubation of 2.5 hours at 37°C, 0.1 mL extraction buffer (20% SDS) was added and incubated overnight at 37°C, and finally, the OD was recorded at 590 mn. The percentage of cell viability was calculated according to the following equation: % cell viability = (O.D. sample – O.D. 100%lysis) / (O.D. 0%lysis – O.D.100%lysis) × 100 [17].

Isolation of peripheral blood mononuclear cells (PBMC)

PBMC from normal volunteers was isolated as per Gopal *et al*, 2014 [17].

Measurement of the antioxidant potentiality of the ME

Determination of DPPH radical scavenging activity

DPPH scavenging activity of methanolic extract of mushrooms was evaluated according to the method of Hajra *et al.* 2018 and Pal *et al.* 2019 [18,8] with slight modification. Briefly, 800 μ L of DPPH solution (0.1mM/mL methanol) was added to 200 μ L of ME extract ranging from 100 μ g/mL-1000 μ g/mL and incubated for one hour in dark, the absorption was finally spectrophotometrically recorded at 517 nm. The experiment was performed in duplicates for each concentration. The percent of scavenging of DPPH radical was calculated according to the following equation: % DPPH reduction = $[(Ac-As)/Ac] \times 100$ (Where; As is the absorbance of sample, Ac is the absorbance of control) [8].

Determination of ABTS•+radical scavenging activity

The ABTS•+ assay was performed with accordance of the method of Ray *et al.* 2020 [4] with minute modifications. Briefly, the working solution of ABTS•+ was prepared by allowing the reaction of the two stock solutions (7.4 mM ABTS and 2.6 mM potassium persulfate solution) in equal quantities for 12 hours at room temperature in the dark. After that the final reaction solution of the ABTS radical was prepared by diluting the aforesaid solution in such a way, that obtain a spectrometric measurement of the absorbance of 0.7 (\pm 0.2) at 734 nm. ME of each mushroom with the concentration range 100 μ g/mL to 1000 μ g/mL was allowed to react with 2.85 mL of the ABTS solution for 2 hours in a dark condition, finally the absorbance was taken at 734 nm using the spectrophotometer. The percent of scavenging of the ABTS•+radical was calculated according to the following equation: % ABTS•+radical reduction = $[(Ac-As)/Ac] \times 100$ (Where; As is the absorbance of sample, Ac is the absorbance of control) [4].

Determination of percentage inhibition of lipid peroxidation

Thiobarbituric acid-reactive species (TBARS) assay was considered to measure the percentage inhibition of lipid peroxidation by ME in accordance with the protocol of Tokuret *al.* 2006 [19] with some modifications. Briefly, 20% v/v egg homogenate and 0.1 mL ME (concentration range 100 μ g/mL-1000 μ g/mL)

along with 0.07 M FeSO₄ was incubated for 30 minutes. After that, acetic acid (pH 3.5), TBA in 1% SDS, was added to the reaction mixture and was vortexed and heated at 95°C for an hour. Finally, Butanol was added to each tube after the cooling down of the reaction mixture, and the tubes were centrifuged at 3000 rpm for 10 min. The upper organic layer was taken for spectroscopic measurement at 532 nm. Percentage inhibition of lipid peroxidation by the crude methanolic extract was calculated according to the following equation: $[(Absorbance\ Control - Absorbance\ Sample) / Absorbance\ Control] \times 100$.

Determination of FRAP activity

FRAP activity of the ME was evaluated by following the protocol of Benzie *et al.* 1996 [20] with minor modification. In brief, the working solution of FRAP was prepared by mixing 300 mM acetate buffer, TPTZ, and FeCl₃, 6H₂O solution (10:1:1), and then warmed at 37°C before use. Mushroom extracts (1.5 mL) were allowed to react with 2.85 ml of the FRAP solution for 30 minutes in dark conditions and finally, the resulted colored product [ferrous tripyridyltriazine complex] was measured at 593 nm by spectrophotometer. The biological property of any mushroom extract depends on the phytochemical compounds that present in the extracts. The phenol and flavonoid are considered the major phytochemical components present in biological mixtures. So, the total phenol and flavonoid content were considered and also quantitatively evaluated.

Determination of total phenol content

The total phenol content of the methanol extract of the five mushrooms was evaluated according to the method of Mridha *et al.* 2017 and Ray *et al.* 2020 by using the Folin-Ciocalteu reagent [4,21]. Briefly, an equal volume of the ME and 0.2 N Folin-Ciocalteu reagent was incubated for 3 minutes, followed by the addition of 10% Sodium carbonate and the mixture was vortexed and incubated in dark at room temperature for 40 minutes and finally, it was spectrophotometrically measured at 760 nm. Gallic acid was considered as standard phenol and a concentration range of Gallic acid (100 μ g/mL-1000 μ g/mL) was prepared as a standard curve. The experimental replica was prepared in a duplicate manner. The result was expressed as mg of gallic acid equivalents (GAE) per gram of methanolic extract.

Determination of total flavonoid content

The total flavonoid content of the methanolic extracts of mushrooms was evaluated according to the method of Kamtekar *et al.*, 2014 [22]. Briefly, 1 mL of the crude extract, 4 mL of distilled water and 5% sodium nitrite solution were incubated for 5 minutes, followed by the addition of 10% aluminum chloride and 1 M NaOH and the whole mixture was vortexed and finally spectrophotometrically measured at 510 nm. Different concentrations of Quercetin (10 µg/mL-100 µg/mL) were considered to prepare a standard curve of standard flavonoid. The experimental replica was prepared in a duplicate manner. The result was expressed as mg of Quercetin equivalents (QE) per gram of methanolic extract.

Identification by gas chromatography mass spectrometry (GC-MS) analysis

The GC-MS profile of both EAE and ME of *Astraeus hygrometricus* was performed by gas chromatographic detectors 7890A (Agilent-Technologies) coupled with a mass spectrometer (MSD 7000). A capillary non-polar column HP-5MS (5% Phenyl Methyl Silox: Agilent 19091S-433) with the dimension of 30 m X 250 µm, along with film thickness: 0.25 µm was considered to perform gas chromatographic analysis and gas chromatographic analysis and helium mobile phase with a constant flow rate of 2.25 mL min⁻¹. 1 µL of the sample with a split ratio of 2:1 at 320 °C was considered for the injection and the oven was programmed as initially at 60 °C, then raised to 320 °C for 12 min at 6 °C min⁻¹ rates. The mass spectrometric analysis was programmed in a full scan module, with an electron impact of 70 eV in a range of 50-550 (m/z). The phytochemicals were identified by analyzing and comparing their related fragmentation profile with spectra present in the library NIST version 2.2 [4,14].

Statistical analysis

All the experimental data, that was obtained from three independent experiments represented as mean ± standard deviation and created by GraphPad Prism, version 7.0 (GraphPad Software Inc, San Diego, CA, USA).

RESULTS

Identification of the mushroom

DNA barcoding is a new emerging and reproducible molecular tool frequently used for the identification and taxonomic revision of biological specimens. *Phallus* sp. was confirmed by our previous publication and reference to the Genbank accession number was MT00752 [4]. *Tricholoma* sp. was verified through molecular identification method through DNA barcoding, the newly generated sequence of 661 bp long, was submitted to Genbank with accession number MT103100. The closest hit of MT103100 is *Tricholoma* sp. BAB-4947 (GeneBank accession number KR155037) with sequence identity 661/662 (99%); gaps 1/662 (0%).

Extraction yield

The extraction yield from both ME and EAE of five mushrooms (*Lentinus* sp., *Astraeus hygrometricus*, *Serpula* sp., *Tricholoma* sp., *Phallus* sp.) was depicted in Table 1. As the percentage of the yield was much higher in ME than EAE, it can be interpreted that there was more polar compound in mushrooms than non-polar compounds. There was no significant difference (P<0.05) in the extraction yield between the extracts of the two mushrooms.

Table 1. Percentage yield of the methanolic and ethyl acetate extract of the collected mushrooms.

Sample	Extract Type	Initial Weight (gm)	Extract Weight (gm)	Yield (%)
<i>Tricholoma</i> sp.	Methanol	50	8.2875	16.575±0.43
	Ethyl acetate	50	0.504	1±0.65
<i>Serpula</i> sp.	Methanol	50	5.8735	11.747±0.42
	Ethyl acetate	50	1.29	2.58±0.48
<i>Astraeus hygrometricus</i>	Methanol	50	1.181	14.354±0.68
	Ethyl acetate	50	7.177	2.362±0.63
<i>Phallus</i> sp.	Methanol	50	6.4905	12.981±0.75
	Ethyl acetate	50	0.4355	0.871±0.66
<i>Lentinus</i> sp.	Methanol	50	6.5905	13.181±0.68
	Ethyl acetate	50	0.4355	0.871±0.63

Antiproliferative assay

MTT result indicated that the two different extracts of the five different genera of mushrooms showed dose-dependent differential anti-proliferative efficacy against Jurkat cells (Figure 1 f, g) and the result was also found to be statistically significant ($p < 0.05$). The efficacy of the methanolic extract (ME) of the five different mushrooms were evaluated and it has been found that *Astraeus hygrometricus* showed the highest anti-proliferative activity with IC₅₀ value of $22.7 \pm 0.23 \mu\text{g/mL}$ followed by *Serpula* sp. $75.7 \pm 0.44 \mu\text{g/mL}$, *Phallus* sp (Pha) $60.53 \pm 0.36 \mu\text{g/mL}$, *Tricholoma* sp. (Tricho) $53.76 \pm 0.46 \mu\text{g/mL}$, *Lentinus* sp. $58 \pm 0.13 \mu\text{g/mL}$ against the Jurkat cell line. On the other hand, we also checked the anti-proliferative activity of the ethyl acetate (EAE) of the same mushrooms (Figure 1g), in that case, all ethyl acetate extracts showed much less

efficacy than ME extracts. So, the result indicated that among five different mushrooms, in all cases the ME extracts performed well than EAE extracts, in particular, the ME extract of *Astraeus hygrometricus* showed robust anti-proliferative activity on leukemic cells (Jurkat) in both cases with promising IC₅₀ values than its counterpart EAE (IC₅₀ $68.9 \pm 0.33 \mu\text{g/mL}$). Both of the extracts, ME and EAE of *Astraeus hygrometricus* showed very minimal effect on human PBMC cells which signifies its specificity towards Jurkat cells.

This result led us to carry the metabolic profiling of both methanolic and ethylacetate extracts, which was performed through GC-MS. In order to decipher the link between ROS generation and Cancer we checked the antioxidant potentiality of the methanolic extracts of all the mushrooms.

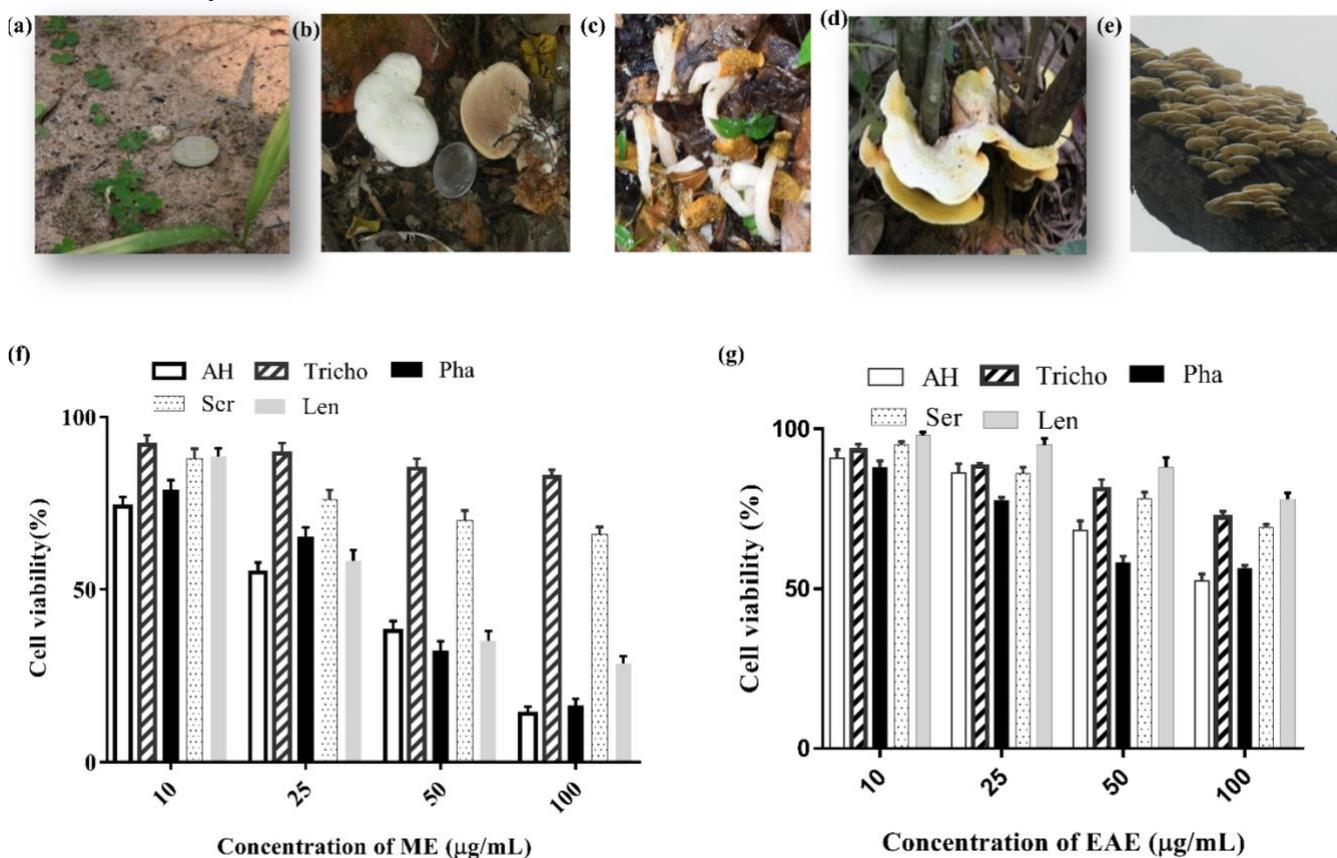


Figure 1. Habitat of the collected mushrooms morphology in its habitat: a. *Astraeus hygrometricus* (AH), b. *Tricholoma* sp. (Tricho), c. *Phallus* sp. (Pha), d. *Serpula* sp. (Ser), e. *Lentinus* sp. (Len). Selective anti-proliferative activity of ME (1f) and EA (1g) of five different mushrooms on Jurkat cell line (f,g).

DPPH radical scavenging assay

The result of the DPPH radical scavenging assay was represented (Figure 2a) in respect of its free radical scavenging potentiality which was carried out with five

different concentrations (100, 200, 500, 800, and 1000 µg/mL) of methanolic extract (ME) of five mushrooms *Lentinus* sp. (Len), *Astraeus hygrometricus* (AH), *Serpula* sp. (Ser), *Tricholoma* sp. (Tricho), *Phallus* sp. (Pha). Results indicated that the free radical scavenging potentiality of ME of five mushrooms occurred in a

concentration-dependent manner. In respect of the IC_{50} value of the standard antioxidant molecule Ascorbic acid $9.9 \pm 0.62 \mu\text{g/mL}$, *Astraeus hygrometricus* (AH) and *Phallus* sp. (Pha) have shown promising results with

IC_{50} value $76.9 \pm 0.16 \mu\text{g/mL}$ and $632.91 \pm 0.32 \mu\text{g/mL}$ respectively. The IC_{50} value of *Lentinus* sp. (Len) was less than $100 \mu\text{g/mL}$ and *Tricholoma* sp. (Tricho) and *Serpula* sp. (Ser) was more than $1000 \mu\text{g/mL}$.

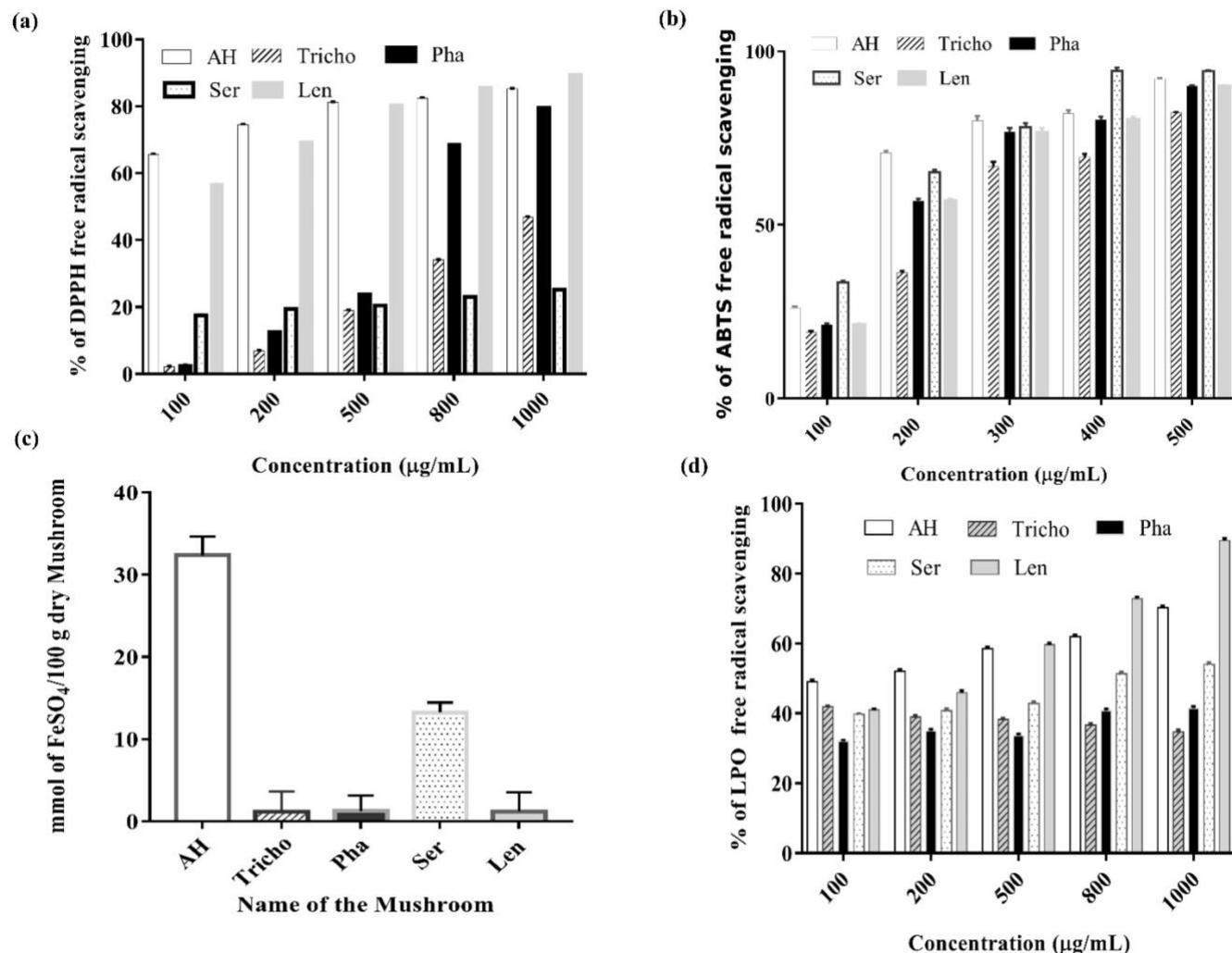


Figure 2. (a-d) Comparative study of antioxidant properties of the methanolic extract of five mushrooms *Astraeus hygrometricus* (AH), b. *Tricholoma* sp (Tricho), c. *Phallus* sp (Pha), and d. *Serpula* sp. (Ser) against DPPH free radical (2 a), ABTS+ free radical (2 b), Lipid peroxidation assay (2 d), and FRAP assay (2 c).

ABTS•+ free radical scavenging assay

ABTS•+ free radical scavenging assay was performed with a concentration range (100-500 $\mu\text{g/mL}$) of the ME of five mushrooms (Figure 2b). The percentage of scavenging of ABTS•+ free radical proportionally increased with the increase of the concentration of mushroom extract in a linear manner. The maximum scavenging percentage of ME at the concentration of 500 $\mu\text{g/mL}$ was ranging between 80% to 92%. The IC_{50} value of the five mushrooms *Astraeus hygrometricus* (AH), *Phallus* sp. (Pha), *Lentinus* sp. (Len), *Tricholoma* sp. (Tricho), and *Serpula* sp (Ser). AH, Tricho, Pha, Ser, Len

against the ABTS•+ free radical was $142 \pm 0.66 \mu\text{g/mL}$, $227.27 \pm 0.72 \mu\text{g/mL}$, $178.5 \pm 0.32 \mu\text{g/mL}$, $153.8 \pm 0.48 \mu\text{g/mL}$ and $178.5 \pm 0.36 \mu\text{g/mL}$ respectively. This result indicated that antioxidant molecules present in ME extract of mushrooms were more effective on DPPH free radical than ABTS•+ free radical.

Lipid peroxidation assay

The Lipid Peroxidation inhibition (LPO) potentiality of any biological extract is a depiction of its antioxidant property. The LPO of ME of five mushrooms with a concentration range of (100-1000 $\mu\text{g/mL}$) is represented

in (Figure 2d). The IC₅₀ value of five mushrooms as follows *Astraeus hygrometricus* (AH) 192.2±0.25 µg/mL, *Tricholoma* sp. (Tricho) 1440±0.32 µg/mL, *Phallus* sp. (Pha) 1219 ± 0.65 µg/mL, *Serpula* sp. (Ser) 925 ±0.65 µg/mL and *Lentinus* sp. (Len) 561.79 ± 0.48µg/mL with respect to the IC₅₀ value of the standard antioxidant Ascorbic acid 29.63 ±0.22 µg/mL.

FRAP

The result of the FRAP assay the antioxidant activities of the mushrooms is represented in (Figure 2c) and expressed as the concentrations of antioxidants having a ferric reducing ability equivalent to that of 1 mM of FeSO₄. The extracts showed a considerable antioxidant effect, *Astraeus hygrometricus* (AH) 32.37±2.31µM, *Tricholoma* sp. (Tricho) 1.15±2.50 µM, *Phallus* sp. (Pha) 1.27±1.9 µM, *Serpula* sp. (Ser) 13.23±1.23µM, *Lentinus* sp. (Len) 1.22 µM ±2.3 and equivalent of FeSO₄/100 g dry mushroom. All the mushroom samples possessed antioxidant activity, while *Astraeus hygrometricus* (AH) and *Serpula* sp (Ser), showed significantly the highest results better than standard ascorbic acid, 2.017±2.9 µM of FeSO₄/100 g equivalent.

Total phenol content

The total phenolic content of ME of five mushrooms was determined by the Folin-Ciocalteu method and is expressed as mg of gallic acid equivalents per gram of extract represented in (Figure 3e). The total phenol content of the mushroom extracts was calculated from the equation $y=0.0002x-0.0125$ (x = gallic acid concentration, y =absorbance of ME, obtained from the standard curve of different concentration of gallic acid, $R^2=0.9974$) and the result indicated that the *Astraeus hygrometricus*, *Tricholoma* sp., *Phallus* sp., *Serpula* sp. and *Lentinus* sp., (AH, Tricho, Pha, Ser, and Len) contained 3.427 ± 0.33 mg, 1.237 ± 0.65 mg, 1.367±0.9 mg, 1.512 ±0.4 mg and 2.4±0.62 mg of GAE per g of mushroom dry weight.

Total flavonoid content

The total flavonoid content of ME of five mushrooms was determined by Aluminium chloride assay and the result is expressed as milligram of quercetin equivalents flavonoid present per gram of mushroom extract. Total flavonoid content of the mushroom extracts (Figure 3f) was calculated from the equation

$y=0.001x+0.002$, $R^2=0.953$ (x = quercetin concentration, y =absorbance of each mushroom methanolic extract)) and the result indicated that the *Astraeus hygrometricus*, *Tricholoma* sp., *Phallus* sp., *Serpula* sp., and *Lentinus* sp. (AH, Tricho, Pha, Ser, and Len) contained 74±0.4 mg, 49 ±0.33 mg, 31±0.90 mg, 14±0.65mg, and 53 ±0.25 mg of GAE per g of mushroom dry weight.

Flavonoid and phenolic compounds are considered as most impactful naturally occurring bioactive compounds which depict the antioxidant profile of any biological extracts [4]. The ME has shown higher flavonoid content than phenol content, so it may be predicted that the flavonoids are a major factor behind the significant antioxidant activity of the ME extracts.

Metabolomic profiling of methanolic extract of *Astraeus hygrometricus*

Among the five mushrooms, the *Astraeus hygrometricus* (AH) had shown antiproliferative activity against T cell leukemia cell line, Jurkat, and also exhibited moderate antioxidant activity, that provoked us to execute the GC-MS analysis for a comprehensive assessment of the metabolomic profile of both the ME and EAE extracts. On the other hand, in order to address the less anti-proliferative potentiality of ethyl acetate extract of AH we have compared the metabolomic profiling of both extracts. In this study, compounds obtained from the EAE and ME of AH were identified by comparing the mass spectra with pre-existing molecular library NIST version 2.2. The number of compounds present in each sample at different retention times and percentages of the area is depicted in (Tables 2 and 3). GC-MS analysis of the ethyl acetate (EAE) extract exhibited a total of 52 compounds (Table 2 and Figure 3 a,c) at different retention times. The total compounds in each extract are divided into major and minor compounds. Major compounds are those having a peak area percentage of more than two whereas minor compounds are less than two. These major and minor compounds of the two extracts are compared on the basis of peak area percentage (Figure 4). The major components of both ethyl acetate and methanolic extract are enlisted on the basis of the presence or absence of bioactivity of the compound in (Table 4). Total compounds were categorized into broad chemical groups, in the case of EAE, the compounds were categorized into twelve broad groups viz. alkane (7), alcohol(4), fatty acid(10), aromatic group of compounds (5), secondary metabolites (4), organic acid (11), polycyclic compound (4), amino acid (1), a heterocyclic group-containing

compound (1), vitamin (2), polymer (1) and hormone (2). The percentage of the area of respective compounds in accordance with resulted GC peak is represented as a pie chart in (Figure 3b). On other hand, the methanolic extracts (ME) of AH showed a total number of 53 compounds (Figure 3d) along with retention time and area percentage are presented in (Table 3). The compounds are categorized in the same manner of EAE

into eleven categories of broad chemical groups viz. monocyclic compound (4), alkane (8), alcohol (4), fatty acid (7), aromatic (6), secondary metabolites (4), organic acid (8), polycyclic (7), heterocyclic (2), vitamin (1) and sugar (2). The percentage of the area of respective compounds in accordance with resulted GC peak is represented as a pie chart in (Figure 3d).

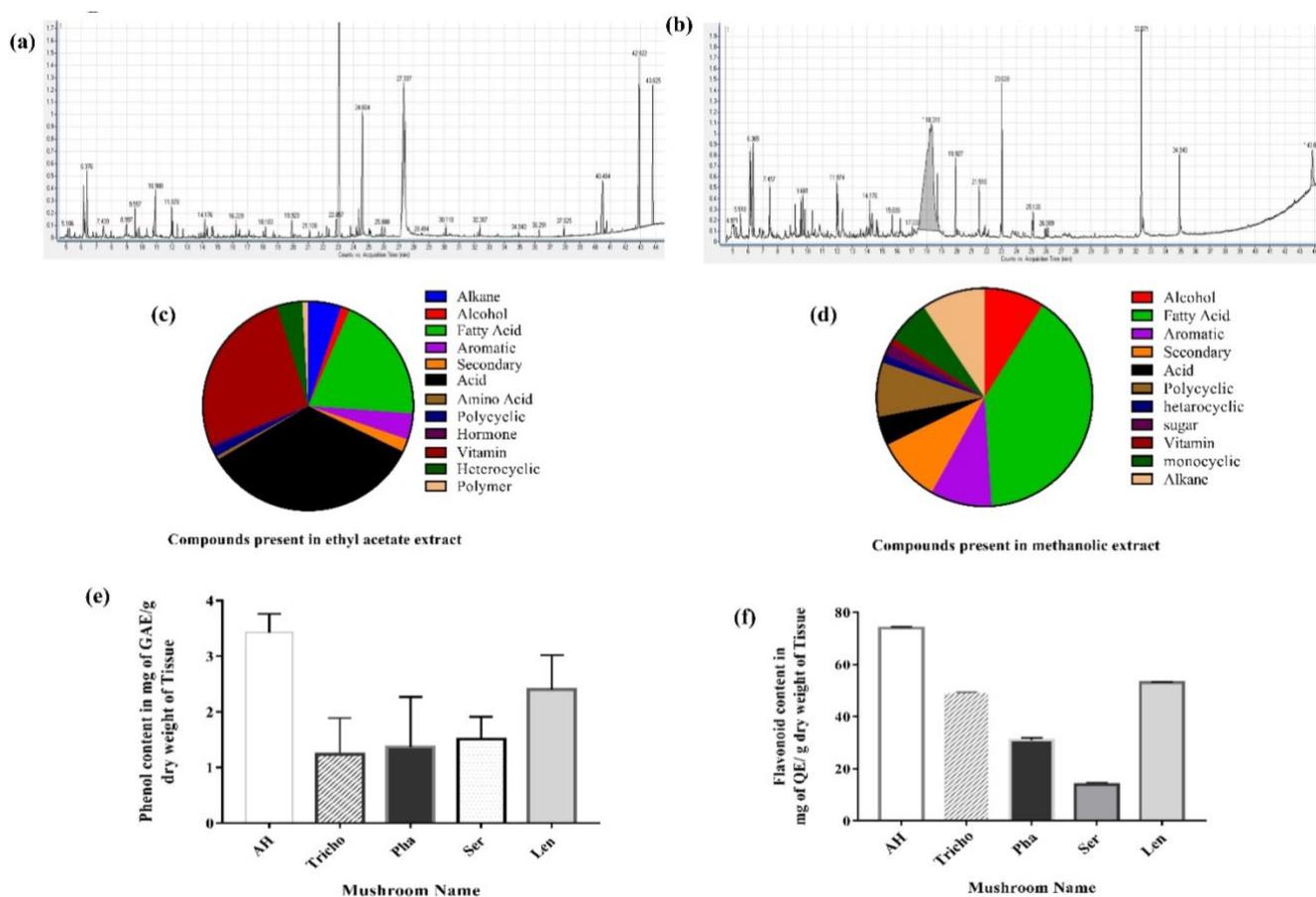


Figure 3. (a-d) Metabolomic profiling of ethyl acetate and methanolic extract of *Astraeus hygrometricus*(AH). GC-MS chromatogram Ethyl acetate (EA) fraction of *Astraeus hygrometricus* (3a). GC-MS chromatogram Methanolic fraction of *Astraeus hygrometricus*(3b). Pie chart that has different sectors. Each sector represents a large group. Each extract of *Astraeus hygrometricus* has many compounds. Now compounds with similar chemical functional or structural structures are categorized under large groups. Each compound has its individual area percentage at a particular retention time. The area percentage of similar compounds in the particular large group represents the area percentage of a large group, this cumulative area percentage of each large group presents an angular sector in the pie chart. (3c) represents different large groups of compounds of ethyl acetate extract of *Astraeus hygrometricus*. There are 12 large groups present. (3d) represents different large groups of compounds of methanolic extract of *Astraeus hygrometricus*. There are 11 large groups present. (3e) Total Phenol content in respect to GAE/g of the methanolic extract of the collected mushrooms AH, Tricho, Pha, Ser, Len. (3f) Total Flavonoid content in respect to quercetin of the methanolic extract of AH, Tricho, Pha, Ser, Len.

Table 2. List of compounds identified in ethyl acetate extract (EAE) of *Astraeus hygrometricus* in GCMS analysis.

List of Compound Present	Molecular Weight	Molecular Formulae	Retention Time	% Area	Cas no.
Alkane					
Limonene	136.234	C ₁₀ H ₁₆	6.227	0.79	138-86-3
Dodecane	170.3348	C ₁₂ H ₂₆	9.83	0.41	112-40-3
1H-Indene, 1-ethylidene-	142.1971	C ₁₁ H ₁₀	11.978	1.02	2471-83-2
Tridecane	184.3614	C ₁₃ H ₂₈	12.036	0.44	629-50-5
Tetradecane	198.388	C ₁₄ H ₃₀	14.176	0.65	629-59-4
Hexadecane	226.4412	C ₁₆ H ₃₄	18.183	0.38	544-76-3
1-Nonadecene	266.505	C ₁₉ H ₃₈	21.703	0.22	18435-45-5
Alcohol					
α-Terpineol	154.2493	C ₁₀ H ₁₈ O	9.688	0.25	98-55-5
Triethylene glycol	150.173	C ₆ H ₁₄ O ₄	10.336	0.33	112-27-6
2,4-Di-tert-butylphenol	206.3239	C ₁₄ H ₂₂ O	16.48	0.24	96-76-4
n-Pentadecanol	228.414	C ₁₅ H ₃₂ O	18.041	0.21	629-76-5
Fatty acid					
Hexanoic Acid(S)	116.1583	C ₆ H ₁₂ O ₂	5.106	0.45	142-62-1
Hexanoic Acid 2 ethyl	144.2114	C ₈ H ₁₆ O ₂	7.963	0.29	149-57-5
Pentadecanoic acid	242.3975	C ₁₅ H ₃₀ O ₂	22.234	0.54	1002-84-2
Palmitoleic acid	254.4082	C ₁₆ H ₃₀ O ₂	24.2	0.57	373-49-9
Eicosanoic acid, ethyl ester	340.5836	C ₂₂ H ₄₄ O ₂	25.044	0.19	18281-05-5
cis-10-Heptadecenoic acid	268.4348	C ₁₇ H ₃₂ O ₂	25.888	0.47	29743-97-3
Octadecanoic acid	284.4772	C ₁₈ H ₃₆ O ₂	26.074	0.35	57-11-4
9,12-Octadecadienoic acid (Z,Z)-	280.4455	C ₁₈ H ₃₂ O ₂	27.337	11.56	60-33-3
9-Hexadecenoic acid	254.4082	C ₁₆ H ₃₀ O ₂	30.118	0.49	2091-29-4
Octadecanoic acid, 2,3-dihydroxypropyl ester	358.5558	C ₂₁ H ₄₂ O ₄	34.94	0.23	123-94-4
Aromatic Compounds					
Phenol	94.1112	C ₆ H ₆ O	5.211	0.50	108-95-2
Benzene 1,2,4 tri methyl	120.1916	C ₉ H ₁₂	5.557	0.24	95-63-6
Benzene 1methyl 3(1-methylethyl)	134.2182	C ₁₀ H ₁₄	6.143	2.04	535-77-3
Biphenyl	154.2078	C ₁₂ H ₁₀	13.794	0.20	92-52-4
Benzophenone	182.2179	C ₁₃ H ₁₀ O	18.802	0.14	119-61-9
Secondary Metabolites					
Phenol	94.1112	C ₆ H ₆ O	5.211	0.50	108-95-2
2,4-Di-tert-butylphenol	206.3239	C ₁₄ H ₂₂ O	16.48	0.24	96-76-4
Squalene	410.718	C ₃₀ H ₅₀	36.291	0.33	111-02-4
Ergosta-5,7,9(11),22-tetraen-3-ol, (3β,22E)-	394.6325	C ₂₈ H ₄₂ O	37.925	0.45	516-85-8
Organic acid					
5-Benzolypentanoic acid	206.24	C ₁₂ H ₁₄ O ₃	6.98	0.25	4144-62-1
1,2,3 Propanetriol 1 acetate	134.1305	C ₅ H ₁₀ O ₄	7.439	0.18	160-61-6
Benzoic acid	122.1213	C ₇ H ₆ O ₂	8.997	0.94	65-85-0
1,2,3-Propanetriol, 1-acetate	134.1305	C ₅ H ₁₀ O ₄	10.751	0.31	106-61-6
Benzeneacetic acid	136.1479	C ₈ H ₈ O ₂	10.9	3.49	103-82-2
Hydrocinnamic acid	150.1745	C ₉ H ₁₀ O ₂	12.724	0.43	501-52-0
2-Propenoic acid, 3-phenyl-	148.1586	C ₉ H ₈ O ₂	14.642	0.47	621-82-9
Benzeneacetic acid, 4-hydroxy-	152.1473	C ₈ H ₈ O ₃	17.084	0.30	156-38-7
Dodecanoic acid, 1-methylethyl ester	242.3975	C ₁₅ H ₃₀ O ₂	18.707	0.23	10233-13-3
Tetradecanoic acid	228.3709	C ₁₄ H ₂₈ O ₂	21.106	0.31	544-63-8
1,2-Benzenedicarboxylic acid, bis(2-methylpropyl) ester	278.3435	C ₁₆ H ₂₂ O ₄	23.046	19.36	84-69-5
Amino acid					
Hydrocinnamic acid(ana-Phe)	150.1745	C ₉ H ₁₀ O ₂	12.724	0.43	501-52-0
Polycyclic compounds					
Naphthalene, 2-ethyl-	156.2237	C ₁₂ H ₁₂	14.107	0.18	939-27-5
Naphthalene, 2,7-dimethyl-	156.2237	C ₁₂ H ₁₂	14.332	0.47	582-16-1
Naphthalene, 1,7-dimethyl-	156.2237	C ₁₂ H ₁₂	14.715	0.35	575-37-1
Naphthalene, 1,4-dimethyl-	156.2237	C ₁₂ H ₁₂	15.046	0.14	571-58-4
Hormone					
Indoleacetic acid	174.176	C ₁₀ H ₈ NO ₂	23.228	0.18	87-51-4
Indole, 3-methyl-	131.1745	C ₉ H ₉ N	13.921	0.20	83-34-1

Vitamin					
Retinoic acid, methyl ester	314.4617	C ₂₁ H ₃₀ O	42.922	12.83	339-16-2
Ergosterol	396.6484	C ₂₈ H ₄₄ O	40.494	7.30	57-87-4
Heterocyclic compounds					
2-Pyrrolidinone 1- methyl	99.1311	C ₅ H ₉ NO	6.376	3.02	872-50-4
Polymer					
dodecyl prop-2-enoate	240.3816	C ₁₅ H ₂₈ O ₂	19.923	0.69	2156-97-0

Table 3. List of compounds identified in methanolic (ME) of *Astraeus hygrometricus* in GCMS analysis.

List of Compound Present	Molecular Weight	Molecular Formulae	Retention Time	% Area	Cas no.
Monocyclic compounds					
Benzene-1 methyl-4-(1-methylethenyl)	132	C ₁₀ H ₁₂	7.457	2.69	1195-32-0
4-Butoxyphenylacetone nitrile	189.2536	C ₁₂ H ₁₅ NO	21.51	2.29	38746-93-9
Morpholine 4[2-(4-tert-butyl benzene sulfoxyl cyclo hexyl)]	365	O(CH ₂ CH ₂) ₂ NH	15.686	0.96	
Benzene-n -butyl	134	C ₁₂ H ₂₁ N	6.769	0.17	104-51-8
Alkane					
Limonene	136	C ₁₀ H ₁₆	6.22	5	138-86-3
Tetradecane	198.388	C ₁₄ H ₃₀	14.176	1.71	629-59-4
3[2-Methyl propenyl]	170	C ₁₀ H ₁₆ O	16.225	0.63	
Octane-5-ethyl-2methyl	156	C ₁₁ H ₂₄	7.628	0.28	62016-18-6
2 Acetamide -2-deoxy-d-manno lactone	217	C ₈ H ₁₃ NO ₆	12.833	0.26	
1-Dodecanethiol	202.4	C ₁₂ H ₂₆ S	16.374	0.16	112-55-0
Hexadecane	226.4412	C ₁₆ H ₃₄	16.163	0.14	544-76-3
1,8,15,22-Tricosatetrayne	308.5002	C ₂₃ H ₃₂	25.128	0.82	4641-78-5
Alcohol					
1-Hexadecanol, 2-methyl-(1S,3S,4S,5R)-1-Isopropyl-4-methylbicyclo[3.1.0]hexan-3-ol	256.4671	C ₁₇ H ₃₆ O	19.927	4.1	2490-48-4
Benzenmethanol-α-(2-nitrocyclopentyl)	154.2493	C ₁₀ H ₁₈ O	9.681	3.19	7712-79-0
zz-2,5-Penta decadein-1-ol	221	C ₁₄ H ₁₀ O ₂	12.029	1.12	103130-20-5
	224	C ₁₅ H ₂₈ O	7.73	0.13	139185-79-8
Fatty acid					
Dodecanoic acid, 1-methylethyl ester	242.3975	C ₁₅ H ₃₀ O ₂	18.311	20.32	10233-13-3
Hexadecanoic acid, 2-hydroxy-1-(hydroxymethyl)ethyl ester	330.5026	C ₁₉ H ₃₈ O ₄	32.371	11.54	23470-00-0
Octadecanoic acid,2,3-dihydroxypropyl ester	358.5558	C ₂₁ H ₄₂ O ₄	34.94	4.69	123-94-4
2,5-Octadecadiynoic acid, methyl ester	290.4403	C ₁₉ H ₃₀ O ₂	11.344	0.41	57156-91-9
10,12- Tricosa diyanoic acid	418	C ₂₃ H ₃₈ O ₂	10.5	0.41	
Hexanoic acid 2-ethyl	144	C ₈ H ₁₆ O ₂	8.505	0.28	
10,13 octadecadiynoic acid methyl ester	290	C ₁₉ H ₃₄ O ₂	13.204	0.16	18202-24-9
Aromatic compounds					
1,2-Benzenedicarboxylic acid, bis(2-methylpropyl) ester	278.3435	C ₁₆ H ₂₂ O ₄	23.028	7.68	84-69-5
Acetophenone	120	C ₈ H ₈ O	6.977	0.3	98-86-2
Biphenyl	154	C ₁₂ H ₁₀	13.794	0.23	
Glycerol 1,2-diacetate	176.1672	C ₇ H ₁₂ O ₅	13.099	0.23	102-62-5
Diphenyl ether	170.2072	C ₁₂ H ₁₀ O	14.281	0.14	101-84-8
Benzeneaceto nitrile 4 Hydroxy	133	C ₈ H ₇ NO	15.042	0.13	1491-95-8
Secondary metabolites					
Eicosane, 7-hexyl-	366.707	C ₂₆ H ₅₄	43.854	6.21	55333-99-8
1,6- Octadecine 2,7 dimethyl(terpenoid)	138	C ₁₀ H ₁₈	5.51	2.02	40195-90-3
2(1H)Naphthalenone octahydro 4a Phenyl-trans	228	C ₁₆ H ₂₀ O	26.118	0.48	22844-36-6
1- Benzoyl-2-(pyrrolidinomethyl)piperine	272	C ₁₇ H ₂₄ N ₂ O	13.557	0.44	
Fatty acid					
Undecanoic acid	201	C ₁₁ H ₂₂ O ₂	9.171	1.81	2432-99-7
4-Hydroxy phenylacetamide	151	C ₈ H ₉ NO ₂	25.918	0.51	30316-13-3

Bicyclo(2,2,1)heptane-3-acetic acid,2 methoxy carbonyl,methyl ester	226	C ₉ H ₁₄ O ₂	21.892	0.49	58987-46-5
Butyric acid 3-(2,5-dimethyl benzoyl) methyl ester	234	C ₁₄ H ₁₈ O ₃	26.009	0.47	
Benzeneacetic acid, 3-hydroxy-, methyl ester	166.1739	C ₉ H ₁₀ O ₃	22.132	0.29	42058-59-3
Amino acetamide	209	C ₂ H ₆ N ₂ O	20.047	0.21	124045-52-9
3 Hydroxy 2 methyl thio-3 phenyl propanoic acid	212	C ₉ H ₁₀ O ₃	4.971	0.17	70760-18-8
Benzeneacetic acid, hexyl ester	220.3074	C ₁₄ H ₂₀ O ₂	10.806	0.12	5421-17-0
Polycyclic compounds					
4a,8a-Methaniminomethano-naphthalene 9,11 dione 10-phenyl	275	C ₁₈ H ₁₃	9.546	2.26	69915-10-2
1H-Indene, 1-ethylidene-	142.1971	C ₁₁ H ₁₀	12.342	1.09	2471-83-2
Triethylene glycol	150.173	C ₆ H ₁₄ O ₄	10.332	1.24	112-27-6
Naphthalene, 1,7-dimethyl-	156.2237	C ₁₂ H ₁₂	14.336	1.18	575-37-1
1H-Indene, 1-ethylidene-	142.1971	C ₁₁ H ₁₀	12.342	1.09	2471-83-2
Naphthalene, 1,4-dimethyl-	156.2237	C ₁₂ H ₁₂	14.711	0.62	571-58-4
Naphthalene, 2-ethyl-	156.2237	C ₁₂ H ₁₂	14.099	0.15	939-27-5
Heterocyclic compounds					
6,7-Dimethyl 3,5,8,8a-tetra hydro1H-2-benzopyran	164	C ₁₁ H ₁₆ O	8.836	0.59	110028-10-9
2H-Pyran-2-one, 5,6-dihydro-	98.0999	C ₅ H ₆ O ₂	4.607	0.53	3393-45-1
Vitamin					
Niacinamide	122.1246	C ₆ H ₆ N ₂ O	13.968	0.85	98-92-0
Sugar					
Phenyl β D -glucoside	256	C ₁₂ H ₁₆ O ₆	5.208	1.26	1464-44-4
β-D-Glucopyranoside, methyl	194.1825	C ₇ H ₁₄ O ₆	17.033	0.41	709-50-2

Table 4. Bioactivity of major compounds (area >2%) of Ethyl acetate and Methanolic extract of *Astraeus hygrometricus*.

List of Compound Present	Area (%)	Bioactivity	Derived compound/related compound	References
1,6- Octadecine 2,7 dimethyl	2.02	Anti-bacterial, antifungal activity,	1,6- octadien-3-ol, 3,7-dimethyl	[34]
O-Cymene	4	Anti-microbial, antimigration and antiproliferation	anti-oxidant and NR	[35-37]
Limonene	5	Anti-microbial, anti-tumor	pharmacological, NR	[38,39]
2 Pyrrolidinone 1- methyl	5.3	Anti-microbial, antibacterial, anticancer and anticonvulsant	antifungal, NR	[40,41]
Benzene-1 methyl-4-(1-methylethenyl)	2.69	NR	NR	NA
4a,8a-Methaniminomethano)naphthalene 9,11 dione 10-phenyl	2.26	Antimicrobial		[42]
(1S,3S,4S,5R)-1-Isopropyl-4-methylbicyclo[3.1.0]hexan-3-ol	3.19	NR	NR	NA
1H-Indene, 1-ethylidene-	2.03	NR	NR	NA
1-Hexadecanol, 2-methyl-	4.1	Anti-microbial	NR	[43]
4-Butoxyphenylacetone nitrile	2.29	NR	NR	NA
1,2-Benzenedicarboxylic acid, bis(2-methylpropyl) ester	7.68	Cytotoxic	NA	[44]
Hexadecanoic acid, 2-hydroxy-1-(hydroxymethyl)ethyl ester	11.54	NR	NA	NA
Octadecanoic acid, 2,3-dihydroxypropyl ester	4.69	NR	NR	NA
Eicosane, 7-hexyl-	6.21	NR	NR	NA

List of compound present in EA extract

Benzene 1methyl 3(1-methylethyl)	2.04	NR	NR	NA
2-Pyrrolidinone 1- methyl	3.01	Cytotoxicity		[3]
4a,8a-(Methaniminomethano) naphthalene 9,11dione,10-phenyl	1.23	NR	NR	NA
Benzeneacetic acid	3.48	Cytotoxicity		[45]
1H-Indene, 1-ethylidene-	1.01	NR	NR	NA
1,2-Benzenedicarboxylic acid, bis(2-methylpropyl) ester	19.35	Cytotoxic	NR	[46]
Palmitoleic acid	1.35	Cytotoxic, Motility	Gastrointestinal NR	[47,48]
Hexadecanoic acid	9.72	Anti-bacterial	NR	[49]
9,12-Octadecadienoic acid (Z,Z)-	11.56	Anti-bacterial, anti-fungal, anti-inflammatory	NR	[49,50]
Ergosterol	7.30	anti-oxidant, anti-cancer	provitamin D, lumisterol and tachy-sterol	[51]
Retinoic acid, methyl ester	12.83	Immunopellet differentiation, cyto toxicity, anti-cancer	NR	[52]
26-27 Dinoergosta-5-33-den-3ol	6.92	NR	NR	NA

(NR-Not Reported, NA-Not Applicable)

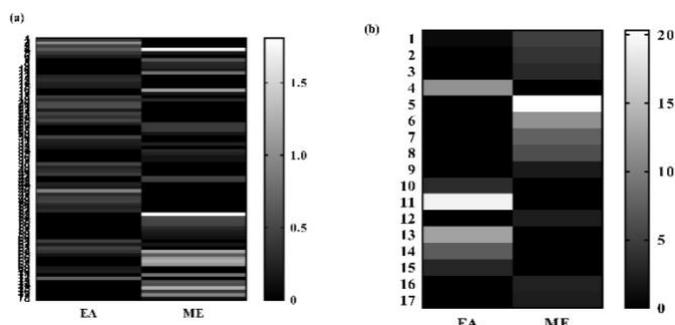


Figure 4. The two columns represent the compounds of ethyl acetate and methanolic extracts of *Astraeus hygrometricus*. The narrow column represents the gradual continuation shades of color black to white. It corresponds to a scale of 0-2 in Fig a, and 0-20 in Fig b. Here the scale denotes area percentage. Total compounds present in both the extract are compared on the basis of the area percentage of each compound. As maximum compounds are restricted to each extract but some are common to both extracts. Though they present in both extracts the common compounds differ in their area percentage. (a) Compounds of *Astraeus hygrometricus* both the extract having area percentage <2. (b) Compounds of *Astraeus hygrometricus* both the extract having area percentage > 2.

DISCUSSION

Mushrooms are rich in the secondary metabolites [23] and proved to be a source of high potential anti-oxidant property and anti-proliferative activity [24]. In this study the five mushrooms that are collected, dried and extracted by solvent percolation method. We have observed that the yielding percentage of methanolic extract of every five mushrooms are appreciable over their respective ethyl acetate extracts. The methanolic extract of *Astraeus hygrometricus* was selected on the basis of the screening studies between methanolic and

ethyl acetate extracts, of five different wild and edible mushrooms viz. *Lentinus* sp. (Len), *Astraeus hygrometricus*(AH), *Serpula* sp. (Ser), *Tricholoma* sp. (Tricho), *Phallus* sp. (Pha) collected from different geographical regions of West Bengal. In all cases of initial screening, we have found that ME of *Astraeus hygrometricus* showed very promising antiproliferative activity by MTT assay that shows indirect cell-mediated cytotoxicity based on hydrolysis of MTT by mitochondrial dehydrogenases of living cells with IC₅₀ value 22.7 ± 0.23 µg/mL [25]. According to many reports Cancer and ROS management are directly related phenomenon [26], so we have also correlated the antioxidant capacity of all the extracts with its antiproliferative efficacy, in this study, in all cases the methanolic extracts of the *Astraeus hygrometricus* has expressed much better antioxidant potentiality that fascinated us to array its metabolic profile. The anti-proliferative assessment study indicated that methanolic extract of *Astraeus hygrometricus* was 3.03-fold potent than its ethyl acetate counterpart and showed negligible anti-proliferative activity against PBMC isolated from normal donors. This initial data upholds the possibility of the up regulation of intermediate cancer signaling molecules such as PI3K/Akt, Wnt, Notch, mTOR, and JaK/Stat pathways, which can be an interesting experimental sphere in future [27].

The volatile compounds in *Astraeus* sp. previously reported indicated that the volatiles found in this Gasteromycetidae were almost exclusively formed from fatty acid [28]. *Astraeus hygrometricus* from

Southwest India reported to contain tannin, phenols and flavonoids [29]. In the present study, the investigation of ethyl acetate and methanolic extracts from fruit body of *Astraeus hygrometricus* revealed the presence of various potent compounds, including Alkane, Alcohol, Fatty acid, aromatic, Secondary metabolites, Acid, Amino acid, Polycyclic, Hormone, vitamin, Polymer, heterocyclic. These bioactive phytoconstituents apart from fatty acid and secondary metabolites are reported as pioneer, could be responsible for the therapeutic ability of methanolic extracts of *Astraeus hygrometricus*. The analysis was carried out by gas chromatography-mass spectrometry (GC-MS), one of the most widely used techniques for separation of phytoconstituents. The GC-MS investigation of methanolic *Astraeus hygrometricus* extracts revealed the presence of 53 compounds, which could contribute to the medicinal properties of this mushroom species. As, the methanolic extract contains much polar compounds, so it can be interpreted that most of the bioactive molecules of the mushroom is polar. By comparing the profile of compounds found in both the methanolic and ethyl acetate extract of *Astraeus hygrometricus*, it can be inferred that some compounds such as limonene, tetradecane, and hexadecane are common in both extracts but present in different percentages. As it was previously reported that C8 compounds, including 1-octen-3-ol, (E)-2-octen-1-ol, 1-octen-3-one, 3-octanone, and 1-octanol, were the main volatile compounds in fresh *Astraeus hygrometricus*. These C8 compounds are characteristic aroma-active compounds in mushrooms, and they contribute greatly to the flavor properties of *Astraeus* sp. whereas it was reported previously that Sulfur-containing compounds (dimethyl sulfone) in the pine mushroom [28] here in this study we had also obtained 2 Acetamide -2-deoxy-d-manno lactone, Hexanoic acid 2-ethyl, Acetophenone, Benzeneaceto nitrile 4 Hydroxy, 4-Hydroxy phenylacetamide as C8 compounds 3[2-Methyl propenyl], Octane-5-ethyl-2methyl, 2 Acetamide -2-deoxy-d-manno lactone, 1-Dodecanethiol, 1,8,15,22-Tricosatetrayne, these are alkanes found only in methanolic extract. Eicosane, 7-hexyl-,1,6- Octadecane 2,7 dimethyl (terpenoid), 2(1H)Naphthalenone octahydro 4a Phenyl-trans, 1-Benzoyl-2-(pyrrolidinomethyl) piperine are secondary metabolites. Benzene-1 methyl-4-(1-methylethenyl), 4-Butoxyphenylacetone nitrile, Morpholine 4[2-(4-tert-butyl benzene sulfoxyl cyclo hexyl)], Benzene-n -butyl are monocyclic compounds. Above these compounds are found only in methanolic extract of *Astraeus*

hygrometricus. Literature survey reveals that most of the listed compounds have cytotoxic, anticancer, anti-inflammatory, anti-diabetic and anti-microbial activity except Dinoergosta-5-33-den-3ol, 1H-Indene, 1-ethylidene-, 4a,8a-Methaniminomethano naphthalene 9,11dione,10-phenyl, and Benzene 1methyl 3(1-methyl ethyl) [30]. Among the compounds of methanolic extract Benzene-1 methyl-4-(1-methyl phenyl), (1S,3S,4S,5R)-1-Isopropyl-4-methylbicyclo [3.1.0] hexan-3-ol, 14-Butyl Phenyl Acetonitrile H-Indene, 1-ethylidene, Hexadecanoic acid, 2-hydroxy-1-(hydroxymethyl) ethyl ester, Octadecanoic acid, 2,3-dihydroxy propyl ester, and Eicosane, 7-hexyl are not reported as a bioactive compound. Rest compounds are mainly anti-microbial inactivity. The predominant molecules that exclusively present on methanolic extracts are Cymene, 2 Pyrrolidinone 1- methyl or 1, 2-Benzenedicarboxylic acid, bis (2-methyl propyl) ester may have a role in inducing robust anti-proliferative activity against Jurkat cell line. Though there are two reports of anti-cancer activity by *Astraeus hygrometricus* on hepatocellular cell lines [31,32] and one tumor regression [33], we are the first to report its anti-leukemic potentiality. As, the methanolic extract of *Astraeus hygrometricus* reveals both antioxidant and anti-leukemic potentiality with a very minimal amount of cytotoxicity against PBMC cells, so it has fulfilled our previous testimony and can be selected as an alternative anti-leukemic drug and also as an antioxidant adjuvant.

CONCLUSION

Screening studies between methanolic and ethyl acetate extracts, of five different wild and edible mushrooms viz. *Lentinus* sp. (Len), *Astraeus hygrometricus*(AH), *Serpula* sp. (Ser), *Tricholoma* sp. (Tricho), *Phallus* sp. (Pha) collected from different geographical regions of West Bengal, showed the most robust anti-proliferative activity on Jurkat cell line by the methanolic extract of the *Astraeus hygrometricus*. As the methanolic extract showed much better anti-proliferative efficacy than its counterpart, the ethyl acetate extract, antioxidant potentiality of the methanolic extracts of all five mushrooms were only evaluated. Among the five different mushrooms, the methanol extract of *Astraeus hygrometricus* exhibited appreciable anti-antioxidant efficacy. The anti-proliferative assessment study indicated that the IC₅₀ value of methanolic extract of *Astraeus hygrometricus* was 22.7 ± 0.23 µg/mL more efficient than ethyl acetate extract which is 68.9 ± 0.33

µg/mL. The minimum anti-proliferative effect was observed in the case of PBMC isolated from normal donors.

The comprehensive comparison of the metabolomic profile of both extracts of *Astraeus hygrometricus* was executed through the GC-MS method. GC-MS chromatogram inferred that there were 52 and 53 compounds from ethyl acetate and methanolic extract respectively. These compounds were categorized according to the chemical family, the compounds detected from the ethyl acetate extract were divided into twelve groups whereas in methanolic extract the compounds were categorized into eleven chemical groups. The compounds of each extract were compared and inferred that the Naphthalene, 2-ethyl-, Naphthalene, 1,7-dimethyl-,Naphthalene, 1,4-dimethyl-,Biphenyl, Hexanoic acid 2-ethyl, Octadecanoic acid,2,3-dihydroxy propyl ester, Limonene, Tetradecane, and Hexadecane were common in both the extract but had different percentage area and retention time. On analyzing the bioactivity of the major compounds of each extract it was found that mainly the compounds were reported as antimicrobial, anti-cancer even anti-inflammatory.

In a nutshell it can be concluded that the methanolic extract of *Astraeus hygrometricus* composed of novel biomolecules that have the efficacy of inducing robust anti-leukemic activity. Further, the extract exhibits appreciable antioxidant activity, which may be a promising potent anti-cancer agent in the future.

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AUTHOR CONTRIBUTIONS

Prof. Santanu Paul designed the entire experiment. Dr. Krishnendu Acharya identified mushrooms. Ribhu Ray collected mushrooms and has performed molecular identification. Amrita Pal has collected and dried the mushrooms, prepared extracts, and performed all the laboratory tests. Prof. Santanu Paul and Amrita Pal analyzed data and wrote the manuscript. All authors approved the final manuscript.

CONFLICTS OF INTEREST

There is no conflict of interest among the authors.

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