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Original Article

### ANTIOXIDANT POTENTIAL AND FATTY ACID COMPOSITION OF *LACTARIUS SANGUIFLUUS*

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#### ABSTRACT

Edible mushrooms have been valued as a source of food and traditional medicines since ancient times. Mushrooms have drawn the attention of numerous researchers in recent decades since they contain a variety of bioactive compounds with antibacterial, antiallergic, antidiabetic, antitumoral, and antioxidant activities. As a member of the Russulaceae family *Lactarius sanguifluus* (Paulet) Fr. (1838) (also known as bloody milk cap) is an ectomycorrhizal edible mushroom distributed in Asia, the Mediterranean shores and mid-Europe. In this study it was aimed to identify the mushroom sample both by physical traits according to the current literature and by molecular techniques. The antioxidant capability of *L. sanguifluus* methanol extract was evaluated by stable radical scavenging activity and metal ion reducing capability methods. The content of total phenolic compounds and the fatty acid composition of the mushroom sample were also investigated. The mushroom sample was identified both by comparing the physical traits with the current literature and by comparing the sequence data of the ITS region of rDNA with GenBank. The methanol extract of the dried mushroom sample was used to perform the analysis to determine the antioxidant activity and content of total phenolic compounds. It was found that *L. sanguifluus* methanol extract contained  $3.677 \pm 0.207$  mg GAE/g DW phenolic compounds and exhibited radical scavenging activities as  $211.652 \pm 0.537$  mg TE/g DW against DPPH and  $73.389 \pm 1.003$  mg TE/g DW against ABTS. *L. sanguifluus* methanol extract also had metal ion reducing capabilities with ferric reducing and cupric reducing antioxidant capacities of  $191.903 \pm 0.659$  mg AAE/g DW and  $379.598 \pm 7.527$  mg TE/g DW, respectively. The GC-MS analysis of the hexane extract of the mushroom sample revealed that palmitic acid, linoleic acid, oleic acid and stearic acid were the predominant fatty acids found in *L. sanguifluus*.

**KEYWORDS:** *Lactarius sanguifluus*, Molecular identification, Antioxidant activity, Total phenolics, Fatty acids.

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#### 1. Introduction

A fungus is a type of eukaryotic heterotroph organism made up of cells called hyphae that resemble tubes. Fungi obtain their nourishment from dissolved substances in their surroundings through the secretion of extracellular digesting enzymes or from the host plants through the symbiotic relationship called mycorrhiza. Generally found above ground, where the fungal mycelia are located, mushrooms are spore-bearing fruiting bodies of fungus with fleshy structures [1].

People have valued edible mushrooms as a source of food and traditional medicines since ancient times. Mushrooms have drawn the attention of numerous researchers in recent decades since they contain a variety

of bioactive compounds with antibacterial, antiallergic, antidiabetic, antitumoral, and antioxidant activities [2]. As a member of the Russulaceae family *Lactarius sanguifluus* (Paulet) Fr. (1838) (also known as bloody milk cap) is an ectomycorrhizal edible mushroom distributed in Asia, the Mediterranean shores and mid-Europe.

The focus of studies with wild *Lactarius* species such as *L. deliciosus*, *L. volemus*, *L. sanguifluus*, *L. semisanguifluus* and *L. piperatus* is on the investigation of their biological properties (antioxidant, antimicrobial, cytotoxicity and neuroprotective activity) and the determination of total phenols [3].

Mushrooms have the potential to be used as sources of antioxidants because they contain bioactive substances such as polysaccharides, vitamins, minerals, polyphenols,

and carotenoids [4-6]. As reactive oxygen species build up from normal metabolic processes, they can lead to oxidative stress, which may accelerate cell aging through lipid peroxidation and cause serious diseases, including Parkinson's disease, cancer, and Alzheimer's disease. Hence, considerable research has been conducted to identify new antioxidant substances that can be used as supplements to prevent oxidative stress while possessing the fewest possible negative effects [7].

Fatty acids have important metabolic functions and are molecules composed of carbon chains with methyl and carboxyl groups at either end. They perform a variety of functions, including energy storage and transit, insulation against mechanical, thermal, and electrical forces, signal transmission, and gene regulation, and are used as components of cell membranes [8].

An abundance of mushroom species can be found in Türkiye because of its rich biodiverse environment [9]. However, little research has been conducted on the biological activities of indigenous mushrooms [10]. In this study it was aimed to determine the content of total phenolic compounds, free radical scavenging activity, metal ion reducing capacity, and fatty acid composition following molecular identification of *Lactarius sanguifluus* collected from Niğde.

## 2. Materials and Methods

### 2.1. Material

Macrofungi sample was obtained in May 2019 from a local bazaar in the Ulukışla district of Niğde, Türkiye. The current literature [11-13] was used to identify the sample based on its morphological features. After being identified, the sample was freeze-dried and stored in laboratory conditions until use.

### 2.2. Molecular identification

Additional molecular identification was carried out to support morphological identification using the Internal Transcribed Spacer (ITS) region of the rDNA, which is often utilized to discriminate fungal species [14].

With a few minor adjustments to the supplier's instructions, DNA was extracted from the samples using the Macherey-Nagel Nucleospin Plant II DNA extraction kit. Polymerase Chain Reaction (PCR) was used to amplify the ITS regions using the primers ITS-1 (5'-TCCGTAGGTGAACCTGCGG-3') and ITS-4 (5'-TCCTCCGCTTATTGATATGC-3') [15]. The PCR product was delivered to BM Labosis (Ankara, Türkiye) for sequence analysis after agarose gel electrophoresis. The macrofungus sample was identified by using the Basic Local Alignment Search Tool (BLAST) to compare data from sequence analysis with the NCBI GenBank database.

### 2.3. Extract preparation

Dried fungal samples were mashed into a powder using a mortar and pestle. Then, 45 mL of methanol and 2 g of powdered material were combined and the mixture was homogenized for 5 min. at 8000 rpm using a Daihan HG-15D homogenizer. The mixture was then incubated overnight at ambient temperature and for 30 min at 30°C in an ultrasonic bath (Sonorex, Bandelin). Following incubation, the mixture was filtered with Macherey Nagel filter paper and dried using a rotary evaporator (Hei-Vap

Value, Heidolph). To create an extract with a concentration of 100 mg/mL, the residue was resuspended in methanol. The antioxidative characteristics of the fungal sample and the total phenolic compounds were analyzed using this extract.

### 2.4. Total phenolic compounds

The total phenolic content in the methanol extract was determined using Folin-Ciocalteu's phenol reagent method [16]. The measurements were conducted in triplicate and the results were expressed as mg GAE/g DW.

### 2.5. DPPH assay

A commonly employed technique to examine the fungal samples' antioxidant activity is the 2,2-diphenyl-1-picrylhydrazyl (DPPH) scavenging capacity. The highest absorbance of DPPH, a dark-colored powder containing stable free radicals, was observed at 517 nm [17]. DPPH radical scavenging assay was performed as reported by Blois [18]. A standard curve was established using the DPPH scavenging values of Trolox solutions and the obtained data were expressed in mg TE/g DW.

### 2.6. ABTS assay

Another commonly used stable radical for assessing the ability of samples with antioxidant qualities to inhibit radicals is 2-azino-bis-(3-ethylbenzothiazoline-6-sulphonic acid) (ABTS). The ABTS radical scavenging activity of the macrofungus extract was evaluated according to the method modified by Cam et al. [19]. The inhibitory percentage values of Trolox solutions at different concentrations were used to create a standard curve, and the findings were represented as mg TE/g DW.

### 2.7. PFRAP assay

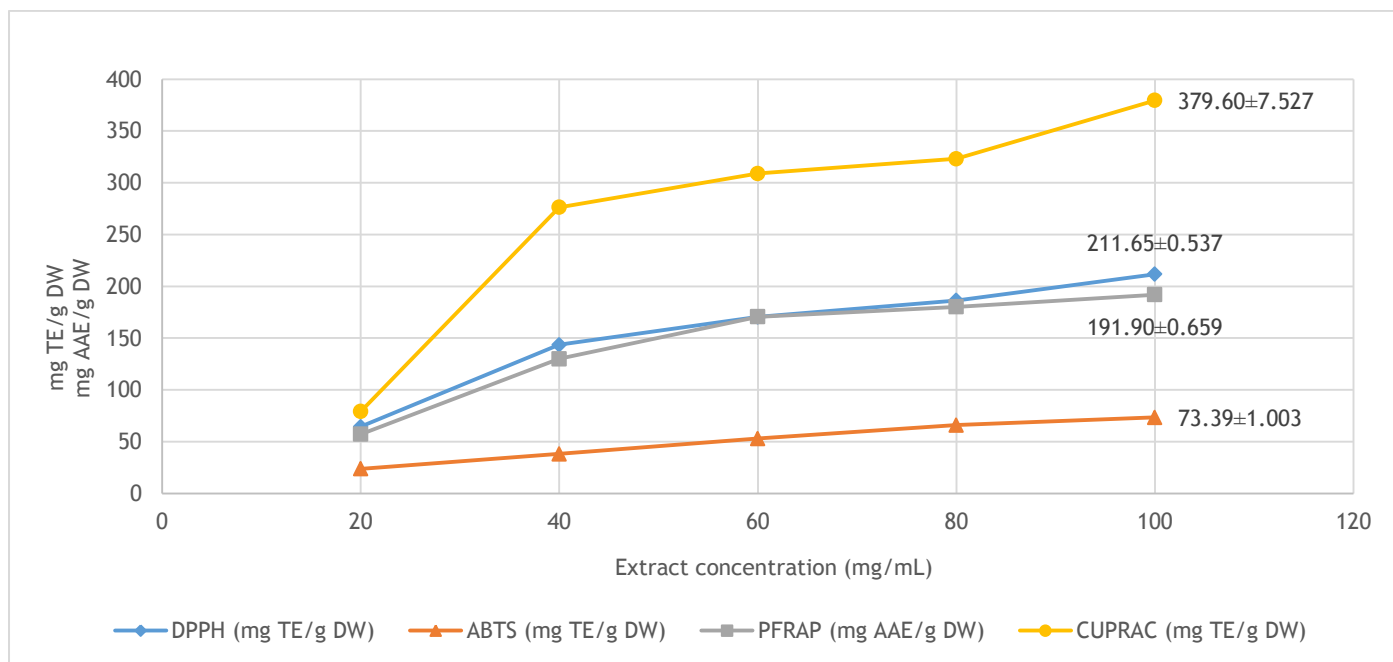
An increase in the absorbance of ferric ferrocyanide which is a blue-colored complex with a maximum absorbance at 700 nm, can be related to the antioxidative activity of the sample which forms potassium ferrocyanide ( $K_4[Fe(CN)_6]$ ) by reacting potassium ferricyanide ( $K_3[Fe(CN)_6]$ ) [20]. The PFRAP assay was performed according to current literature [21,22] with slight modifications. The ascorbic acid standard curve was used for data evaluation, and the results were represented as mg AAE/g DW.

### 2.8. CUPRAC assay

The color change of neocuproine and  $Cu(II)Cl_2$  mixture from bright yellow to orange in the presence of compounds acting as electron donors indicates the antioxidant capacity of a sample by reducing cupric ( $Cu^{2+}$ ) ions to cuprous ( $Cu^+$ ) ions. The CUPRAC assay was performed according to the method originally reported by Apak et al. [23], with slight modifications. The absorbance of all samples was measured at 450 nm against a blank solution. The Trolox standard curve was established to evaluate the antioxidant capacity of mushroom samples with different concentrations and the obtained data was expressed as mg TE/g DW.

### 2.9. Determination of fatty acid composition

The fatty acid composition of fungus sample was determined by gas chromatography-mass spectrometry (GC-MS). One g of powdered mushroom sample was mixed



**Fig. 1.** Free radical scavenging activity and metal ion reducing capacity of *L. sanguifluus* methanol extract. (DPPH: DPPH scavenging activity, mg TE/g DW; ABTS: ABTS scavenging activity, mg TE/g DW; PFRAP: Potassium ferricyanide reducing antioxidant power, mg AAE/g DW; CUPRAC: Cupric ion reducing antioxidant capacity, mg TE/g DW; TE: Trolox equivalent; AAE: Ascorbic acid equivalent; DW: Dry weight).

with 20 mL hexane and homogenized for 5 min. at 8000 rpm. The mixture was then incubated in an ultrasonic bath for 30 min at 30°C. After filtering the homogenate the solvent was evaporated and the residue was resuspended with 5 mL of hexane. Methyl esters of the fatty acids were derived from the hexane extract with 2 M KOH in methanol and 1 N HCl solution. The clear upper layer after the phase separation was dried with anhydrous Na<sub>2</sub>SO<sub>4</sub> and passed through a 0.45 µm syringe filter. An aliquote of 1 µL sample was analyzed by a Shimadzu QP2010 Ultra GC-MS system equipped with Restek Rxi-5MS column. Helium was used as the carrier gas at a flow rate of 1 mL/min. The fatty acids were characterized by comparing the obtained spectra with those from the Wiley (W9N11) mass spectra library and Flavor and Fragrance Natural and Synthetic Compounds (FFNSC 1.2) library [24].

### 3. Results and Discussion

#### 3.1. Identification of macrofungus

The macrofungus sample investigated in this study was identified both in terms of morphological characteristics according to the current literature [11-13] and by comparison of sequence data with those deposited in the GenBank database via BLAST. Most of the data in GenBank comprise sequence data of ITS fragments belonging to fungal samples. According to results the macrofungus sample was identified as *Lactarius sanguifluus* (Paulet) Fr. (1838) with a similarity rate of 99.85%.

#### 3.2. Antioxidant activity

The antioxidant capability of the *L. sanguifluus* methanol extract was investigated using four different

methods, along with the content of total phenolic compounds. The results are shown in Fig. 1 and Table 1.

According to Fig. 1, *L. sanguifluus* methanol extract exhibited high DPPH radical scavenging activity with 211.65±0.537 mg TE/g DW and 73.39±1.003 mg TE/g DW ABTS radical scavenging activity and high metal ion reducing activities with 191.90±0.659 mg AAE/g DW potassium ferricyanide reducing activity and 379.60±7.527 mg TE/g DW cupric reducing activity. These findings indicate that the free radical scavenging activity and metal ion reducing capacity of *L. sanguifluus* collected from Türkiye are similar to those reported in the literature [25-27].

**Table 1.** Biochemical properties of *L. sanguifluus* methanol extract.

| TPC             | DPPH             | ABTS            | PFRAP            | CUPRAC           |
|-----------------|------------------|-----------------|------------------|------------------|
| 3.677<br>±0.207 | 211.65<br>±0.537 | 73.3<br>9±1.003 | 191.90<br>±0.659 | 379.60<br>±7.527 |

(TPC: Total phenolic compounds, mg GAE/g DW; DPPH: DPPH scavenging activity mg TE/g DW; ABTS: ABTS scavenging activity, mg TE/g DW; PFRAP: Potassium ferricyanide reducing antioxidant power, mg AAE/g DW; CUPRAC: Cupric ion reducing antioxidant capacity, mg TE/g DW; GAE: Gallic acid equivalent TE: Trolox equivalent; AAE: Ascorbic acid equivalent; DW: Dry weight)

The phenolic compounds are generally correlated with the antioxidant capability since they can act as hydrogen donors, reducing agents, and singlet oxygen scavengers [28]. The content of total phenolic compounds was determined as 3.667±0.207 mg GAE/g DW for *L. sanguifluus*. Yıldız and Gürgen [29] stated that the content of total phenolic compounds of three *Lactarius* species (*L. deciosus*, *L. insulsus* and

**Table 2.** Fatty acid composition of *L. sanguifluus* determined by GC-MS.

| No. | Compound  | RT     | MW+CF   | RC    | SI |
|-----|---|--------|---|-------|----|
| 1   | Myristic acid, methyl ester   | 32.761 | 242 (C <sub>15</sub> H <sub>30</sub> O <sub>2</sub> ) | 0.54  | 87 |
| 2   | Palmitic acid methyl ester  | 37.912 | 270 (C <sub>17</sub> H <sub>34</sub> O <sub>2</sub> ) | 12.31 | 94 |
| 3   | 3,5-Bis(1,1-dimethylethyl)-4-hydroxy-benzenepropanoic acid methyl ester | 38.396 | 292 (C <sub>18</sub> H <sub>28</sub> O <sub>3</sub> ) | 1.28  | 86 |
| 4   | Linoleic acid, methyl ester   | 41.614 | 294 (C <sub>19</sub> H <sub>34</sub> O <sub>2</sub> ) | 6.69  | 97 |
| 5   | Oleic acid, methyl ester  | 41.736 | 296 (C <sub>19</sub> H <sub>36</sub> O <sub>2</sub> ) | 7.86  | 95 |
| 6   | Stearic acid, methyl ester  | 42.244 | 298 (C <sub>19</sub> H <sub>38</sub> O <sub>2</sub> ) | 25.03 | 97 |
| 7   | 2-Propenoic acid, tetradecyl ester                                      | 45.471 | 268 (C <sub>17</sub> H <sub>32</sub> O <sub>2</sub> ) | 2.06  | 88 |
| 8   | Arachidic acid methyl ester   | 46.044 | 326 (C <sub>21</sub> H <sub>42</sub> O <sub>2</sub> ) | 0.48  | 85 |

\*RT: Retention time, min; MW: Molecular weight; CF: Chemical formula; RC: Relative concentration, %; SI: Similarity index, %

*L. vellereus*) varied between 1.124±0.032 and 2.247±0.086 mg GAE/g. *L. vellereus*, which had the highest phenolic compounds with a value of 2.247±0.086 mg GAE/g, also demonstrated the highest antioxidant activity with a FRAP value of 5.668±0.042 µmol FeSO<sub>4</sub>/g. These findings supported the correlation between the phenolic compounds and the antioxidant activity.

Erdoğan et al. determined the DPPH scavenging activity and the amount of phenolic compounds as 1283.51 µmol TE/100 g DW and 1133.12 mg GAE/100 g DW, respectively. Researchers have stated that the possible mechanism of antioxidant activity of mushroom extracts occurs through the hydrogen bonding capacity and oxidation of peroxy radicals [30]. Stankovic et al. examined four species of mushrooms in a 2022 study: *L. deliciosus*, *L. volemus*, *L. sanguifluus*, *L. semisanguifluus* and *L. piperatus* [31]. The mushroom species *L. sanguifluus* exhibited the highest levels of total phenolic compounds and the greatest antioxidant activity as measured by ABTS and CUPRAC assays. It has been stated that the *Lactarius* genus is not only a food source but also contains components with high antioxidant potential.

The factors affecting the difference in the content of total phenolic compounds and antioxidant capabilities between this study and the literature can be considered to be the geographical position and altitude of the sampling area, environmental and meteorological conditions, growth stage of the macrofungus, extraction method and solvent.

### 3.3. Determination of fatty acid composition

The fatty acid composition of *L. sanguifluus* was also investigated by GC-MS and the results are shown in Table 2.

Stearic acid (25.03%, RT: 42.244 min), palmitic acid (12.31%, RT: 37.912 min), oleic acid (7.86%, RT: 41.736 min), and linoleic acid (6.69%, RT: 41.614 min) are found to be the predominant fatty acids in *L. sanguifluus*. Although the saturated fatty acid concentrations were found higher than the unsaturated fatty acids, which are considered healthy components for human diet, it was reported that stearic acid did not affect the increase in total cholesterol and low density lipoproteins in human blood [32]. Myristic acid (0.54%, RT: 32.761 min), which was associated with increased cholesterol levels, along with palmitic and lauric acids [33], was also determined. 2-Propenoic acid (2.06%, RT: 45.471 min), 3,5-Bis(1,1-dimethylethyl)-4-hydroxy-benzenepropanoic acid (1.28%,

RT: 38.396 min) and arachidic acid (0.48%, RT: 46.044 min) were also detected by GC-MS analysis of fatty acids in *L. sanguifluus* collected from Niğde. In this study, stearic acid was the major fatty acid detected in *L. sanguifluus*, and it is one of the most important essential fatty acids. Mushrooms are rich in fat and fatty acids, which are necessary for human metabolism. Myristic, palmitic, stearic, oleic, and linoleic fatty acids are commonly found in *Lactarius* species [4,34-36]. Supporting our findings, it was reported that palmitic, stearic, oleic, and linoleic acids were determined as predominant fatty acids in the study conducted with three *Lactarius* species, namely *L. piperatus*, *L. quietus* and *L. vellereus* [37]. Erbiai et al. [25] reported that *Lactarius sanguifluus* is rich in organic compounds and contains high amounts of phenolic acids, ascorbic acid, flavonoids, tannins, sugars, fatty acids. The researchers stated that the high amounts of these chemical compounds in the extracts of *L. sanguifluus* may be linked to the significant biological activities observed [25].

### 4. Conclusions

In conclusion, the results of this study provide valuable insights into the total phenolic compound content, antioxidant activity and fatty acid composition of *L. sanguifluus* collected from Niğde, highlighting its potential as a promising source of bioactive compounds for various applications. Future studies should focus on investigating the bioactive compounds present in *L. sanguifluus* and their potential health benefits as well as exploring the antioxidant activity and fatty acid composition of other mushroom species.

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