

PROTEIN DETERMINATION THROUGH BRADFORD'S METHOD OF NEPALESE MUSHROOM

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Abstract: The Wild edible mushrooms are one of the most important non timber forest products. Due to its vigorous growth in the rainy season, delicious taste and nutritional value the mycophagus group consume them. In this paper an attempt has been made to determine the protein of these 35 species through Bradford's method. Among them thirty three species of wild mushroom collected from different altitude (200m – 4200m), phytogeographical habitat of central Nepal and two species *Agaricus bisporous* & *Pleurotus sajorokaju* cultivated sample from Balambu farm. The highest amount of protein 1.576mg/ml in *Cantharellus subscibarius* and least 0.131 mg/ml in *Cordyceps sinensis* were found.

Key words: Wild mushroom; Protein; Bradford's method.

INTRODUCTION

Protein is the most critical component contributing to the nutritional value of food. The determination of crude protein was done by the procedure known as Micro Kjeldahl's process (Sawhney, Singh, 2000). However, this process is both time consuming and sample consuming (Young, 1963).

Bradford's Method (Bradford, 1976) is a rapid, simple and sensitive method for estimation of proteins in a sample extract. The color development is virtually complete in 2 minutes and the color is stable for about 1 hour. Unlike Lowry's method (Lowry et al., 1951) metal ions such as NH_4^+ , Na^+ , K^+ , phenol and carbohydrate such as sucrose do not interfere in this assay. The procedure is based on interaction of a dye, Coomassie Brilliant Blue with protein.

Wild edible mushrooms are one of the important minor forest products, which are locally traded in local market of the country. In Nepal the mushroom collection and consumption have been continuing since time immemorial by different ethnic groups. Due to the lack of scientific knowledge in ethnic groups, they only utilized few species that could be identified from their own traditional knowledge. Various mycophagus groups like Sherpa, Tamang, Gurung, Tharu, Danuwar, Chepang and Newar are directly concerned with the collection and consumption of mushrooms since long due to its delicious taste as well as vigorous growth during rainy season in appropriate habitat in the forest of the country. Due to their high content of vitamin, protein and mineral, mushrooms are considered as "Poor man's Protein" (Pandey 2004).

The Greeks and Romans described mushrooms as "Food for

the Gods", and were served only on celebrations. Reference to mushrooms is found in Vedas (Chaube, 1995; Adhikari, 2000, 2004). Mushrooms can be used for the food to solve the malnutrition problem (Manandhar, 2003). Mushrooms have good nutritional value particularly as a source of protein that can enrich human diets especially in some developing countries where animal protein may not be available and are expensive. The protein content of fresh mushroom is 3.7% stated by Food Agriculture Organization's publication in 1978. The edible and medicinal mushrooms can be used on human welfare in the 21st century (Chang, 1999).

Nowadays the marketing of wild mushroom is decreasing due to new policy of forest conservation applied by Nepal Government yet some delicious mushrooms were also supplied in the reputed hotels through different means. Wild edible mushrooms are also called "Rajashi Khana (Food for Royal Palace)" due to delicious taste (Pandey, 2004). More than twenty species of mushrooms have been cultivated commercially around the world (Manandhar, 1994).

OBJECTIVES

The main objective of this research is

- To collect and identify mushroom samples in the study area.
- To find out the amount of protein present in different species of mushrooms.

MATERIALS AND METHODS

The mushrooms needed for the research were collected from the Langtang, Kathmandu valley and adjoining area. Identification of different collected samples were done on

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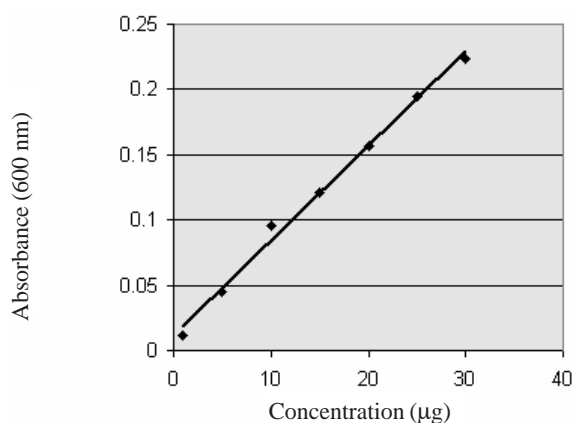


Figure 1: BSA Calibration curve for Bradford Method.

the basis of different important mushrooms characters and techniques such as habit and habitat, morphological structures, spore print color, microscopic observations, relevant monograph, recent literatures and books (Adhikari, 1990, 2000; Arora, 1986; Pandey, 2004; Pandey, 2006 a,b,c; Pacioni, 1981; Singer, 1986, Rea, 1922; Fries, 1838; Millor, 1984. Internet as well as cross checking with the voucher specimen. The proteins were determined by Bradford's method (Bradford, 1976).

An amount of 2 gm of mushroom samples were cut into pieces with a scissor and grinded in mortar with 5ml of phosphate

buffer (pH 7.6) and was then transformed to the centrifuge tubes. The homogenate was centrifuged at 8000 rpm for 20 minutes. The supernatant of different mushroom samples were put in separate tubes. The volume of all of the samples in tubes were then made equal by adding phosphate buffer solution and the extraction were stored in the refrigerator at 4°C for further analysis.

After extraction, 30µl of different mushroom samples were taken out in separate tubes and were mixed with 70µl of distilled water separately. In all of these separate sample tubes 2.9 ml of Coomassie Brilliant Blue solution was then added and mixed thoroughly. The Total volume now was 3ml in each tube. All these tubes were incubated for 5 minutes at room temperature and absorbance at 600(595) nm was recorded against the reagent blank. A standard curve of Absorbance (600 nm) versus Concentration (µg) of protein was plotted as shown in Figure 1.

Protein content in the extracted samples was determined from the standard curve and the amount of protein in mg/ml was calculated. The results were illustrated in the Table 1.

RESULTS

The highest amount of protein was observed in *Cantharellus subcibarius* 1.567 mg/ml followed by *Russula Virescens* 1.427 mg/ml, *Lactarius piperatus* 1.38 mg/ml and least amount of protein was found in *Cordycep sinensis* 0.131 mg/ml. (Table 1)

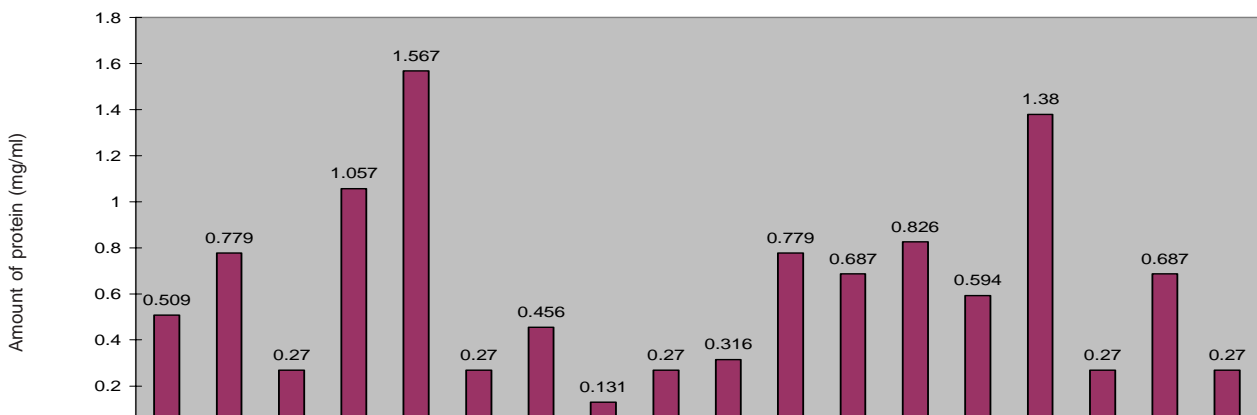
Table 1

Sample	Coll.No	Place of coll.	Abs(600nm)	Conc. in 30µl	protein mg/ml
<i>Agaricus bisporus</i>	Market	Baneshwor	0.1	19.22222	0.509
<i>Agaricus campestris</i>	24536	Kirtipur	0.18	23.38889	0.779
<i>Amanita vaginata</i>	24609	Matatirtha	0.1	12.27778	0.27
<i>Cantharellus cibarius</i>	24630	Suryabinayak	0.24	31.72222	1.057
<i>Cantharellus subcibarius</i>	24606	Matatirtha	0.35	47	1.567
<i>Clavaria rosea</i>	24584	Sundarijal	0.07	8.111111	0.27
<i>Coprinus comatus</i>	21108	Langtang	0.11	13.66667	0.456
<i>Cordycep sinensis</i>	23455	Langtang	0.04	3.944444	0.131
<i>Coriolus hirsitus</i>	25645	Matatirtha	0.07	8.111111	0.27
<i>Ganoderma tsugae</i>	23440	Langtang	0.08	9.5	0.316
<i>Hypholoma capsonoid</i>	25653	Langtang	0.18	23.38889	0.779
<i>Laccaria laccata</i>	25629	Suryabinayak	0.16	20.61111	0.687
<i>Laccaria laccata</i>	24598	Matatirtha	0.19	24.77778	0.826
<i>Laccaria laccata</i>	24526	Godawari	0.14	17.83333	0.594
<i>Lactarius piperatus</i>	22158	Suryabinayak	0.31	41.44444	1.38
<i>Lactarius volemus</i>	24570	Tistung Palung	0.09	10.88889	0.27
<i>Laetiporus sulphureus</i>	24541	Langtang	0.16	20.61111	0.687
<i>Marasmius maximus</i>	24572	Tistung Palung	0.09	8.111111	0.27
<i>Morchella conica</i>	23461	Langtang	0.14	17.8333	0.594
<i>Mycena Sp</i>	25651	Langtang	0.16	20.61111	0.687
<i>Omphalotus olearis</i>	24576	Baneshwor	0.16	20.61111	0.687
<i>Oudemansiella radicata</i>	24617	Baneshwor	0.14	17.8333	0.594
<i>Pahelo chyau</i>	24573	Baneshwor	0.1	12.27778	0.409
<i>Pleurotus sajorkaju</i>	Market	Tistung Palung	0.15	12.27778	0.64
<i>Pleurotus Cornucopiae</i>	23469	Matatirtha	0.16	20.61111	0.687

Abbreviations: Sp = Species, Conc = Concentration, Coll = Collection, No = Number.

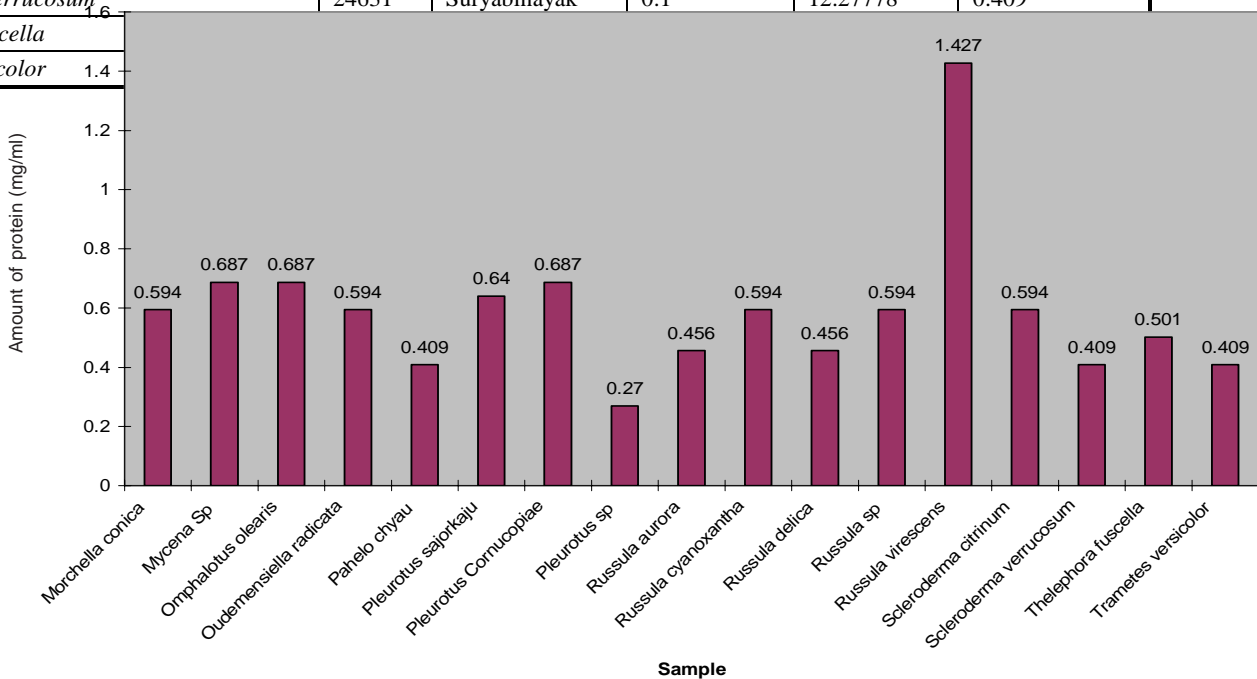
Sample Versus Amount of protein in mg/ml Charts:

Sample Versus Amount of protein in mg/ml Chart



<i>Pleurotus sp</i>	24505	Tistung Palung	0.07	8.111111	0.27
<i>Russula aurora</i>	24801	Matatirtha	0.08	13.66667	0.456
<i>Russula cyanoxantha</i>	24605	Matatirtha	0.32	42.83333	0.594
<i>Russula delica</i>	24571	Tistung Palung	0.1	13.66667	0.456
<i>Russula sp</i>	24604	Matatirtha	0.14	17.83333	0.594
<i>Russula virescens</i>	25681	Dakchinkali	0.32	42.83333	1.427
<i>Scleroderma citrinum</i>	25412	Nuwakot Tigaon	0.14	17.83333	0.594
<i>Scleroderma verrucosum</i>	24631	Suryabinayak	0.1	12.27778	0.409

Sample Versus Amount of protein in mg/ml Chart (Continued)



DISCUSSION AND CONCLUSION

Large amount of protein is found in mushrooms. Hence it is regarded as an ideal protein source for vegetarian as well as for old age people who are unable to chew meat. It was found that the amount of protein varies from species to species in the same genus as for example two different species of genus *Cantharellus*, i.e., *Cantharellus cibarius* contents 1.057 mg/ml of protein while *Cantharellus subcibarius* contents 1.567 mg/ml of protein. Similarly the three different species of the genus *Pleurotus*, four species of the genus *Russula* and two species of the genus *Scleroderma* have different protein concentration as shown in the Table1.

In our research, wild edible mushrooms *Laccaria laccata* collected from three different places namely Godawari, Matatirtha and Suryabinayak were found to have 0.594, 0.862 and 0.687 mg/ml amount of protein. Clearly, it was seen that the amount of protein in mg/ml is different even in the mushrooms of the same species collected at the three different places. Hence it was also concluded from the research that the amount of protein varied even in the same species according to phyto-graphical condition.

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