

## Extraction and quantitative estimation of nucleic acids and proteins from liver tissue of *Heteropneustus fossilis* (fish) and *Caprine barbari* (goat)

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**Abstract:** 2gms of liver tissue was collected from *Heteropneustus fossilis* (fish) and *Caprine barbari* (goat) for extraction and estimation of DNA, RNA and proteins. Nucleoproteins are soluble in solution of high ionic strength but insoluble in solution of low ionic strength. This property was used for extraction and estimation of DNA by diphenylamine. RNA was extracted by trichloroacetic acid and quantitative estimation was made using oriconol reagent. Total proteins were extracted and estimated by biuret method. DNA extracted was 0.42mg, RNA 3.34mg and proteins 13.4mg from 2gms of liver tissue of *Heteropneustes fossilis* (fish) and from 2gms of liver tissue of *Caprine barbari* (goat) DNA estimated was 0.57mg, RNA 2.5mg and proteins were 12.4mg. It is seen that proteins estimated are more than DNA and RNA in given amount of tissue.

**Key words:** Nucleic acids, Proteins, *Heteropneustes fossilis*, *Caprine barbari*.

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### I. Introduction

The living cell is an extraordinarily complicated entity producing thousands of different macromolecules and harboring a genome. The methods of molecular biology depend upon an understanding of the properties of biological macromolecules. The systematic comparison of different animal genomics gives a chance of identifying genetic basis for diversity. We are fast entering a golden era of comparative genome analysis (Watson, 1953). Mice and humans contain roughly same number of genes about 28,000 protein coding genes. A golden era of molecular biology was launched after it became possible to isolate specific DNA segments. Recently it became possible to apply molecular approach to analysis of nucleic acids and protein changes in gene expression causing morphological changes in different groups of animals. There is hope that analysis of DNA will illuminate the mechanisms of evolutionary diversity. Mackey et al (1996) isolated DNA from liver tissue, muscles and plants. Iordi et al (1979) studied DNA component of chicken genome. Scinew et al (1990) studied mitotic chromosome complements and nuclear DNA content of four species of shrimps. Deoxyribonucleic acid is the genetic material and stores genetic information. Eukaryotic cells contain many DNA molecules. Chromatin is a complex of DNA and histone. Nucleic acids are usually associated with proteins to form nucleoproteins. Sakagami (1991) reported high content of DNA protein in liver as compared to kidney, heart and wings. Zang et al (1990) isolated the protein from rat liver using alcohol precipitation and ultra filtration. Misky and Osawa (1991) isolated DNA from nuclei. Konat et al (1996) gave a simple technique for the isolation of higher molecular weight genomic DNA from animal tissue and cell. Pal and Holkar (2013) gave that content of protein is higher than DNA and RNA in same amount of liver tissue of *Clarias batrachus*, *Gallus domesticus* and *Rattus rattus*. Proteomics is the science to determine which proteins are produced in a cell or tissue under a specific set of conditions, Devlin (2006). Goklen et al (1985) studied protein extraction. Gomez-Boustista (1986) studied level of total protein and protein fractions in the serum of rabbit.

### II. Material And Methods

2 gms of liver tissue was collected from *Heteropneustes fossilis* and *Caprine barbari* for each experiment. Samples were collected from fresh animals and experiments for extraction and estimation were repeated ten times. Eight results from each group were taken into consideration. Nucleoproteins are soluble in solution of high ionic strength but is insoluble in solution of low ionic strength. This property was used for extraction and estimation of DNA by diphenylamine. Diphenylamine was prepared by dissolving one gram of diphenylamine in 100 ml of glacial acetic acid and by adding 2.5 ml concentrated sulphuric acid. RNA was extracted by trichloroacetic acid and quantitative estimation was made using orcinol reagent. Total proteins were extracted and estimated by biuret method.

### III. Results And Discussion

Protein are the most abundant macro molecules in cells and constitute over half the dry weight of most organism. Genetic information is expressed by proteins. Each species of animal has a characteristic content of DNA. Eukaryotes vary greatly in DNA content and contain more DNA than prokaryotes. Lin, zang-ping et al. (1979) isolated DNA from plant materials. Misky and Osawa (1991) isolated DNA from nuclei which enabled various quantitative studies to be made. Sakagami et al (1991) in his study showed that mitochondrial DNA decreases in order of liver to testis to placenta. Litjerja (1980) determined the DNA content in the cerebellum of the rat in both male and female. Mackey et al (1996) isolated the DNA from tissue such as liver, muscles and plant. Shotwinska et al (1979) isolated DNA from liver of rats during hepatoma development. Shrinivasan (1988) described the effect of gamma radiation on rat tissue. Chetokov et al (1986) isolated phenylalanine hydroxylase from human liver. Bhattacharya et al (1990) made biochemical studies of liver of rats. Black and Johnson (1984) studied protein structure. Bart et.al(1985) in his studies showed that rat liver contained higher amount of RNA than DNA. More can be added to Molecular biology by studying more of deoxyribonucleic acid, ribonucleic acid and proteins. Zang et al (1990) extracted protein from rat liver.

Table 1:- DNA content in 2gms of liver tissue of *Heteropneustus fossilis*(Fish) and *Caprine barbari*(Goat)

S.no.	Class	Animal	Organ	Amount	Mean(mg)
1.	Pisces	<i>Heteropneustes fossilis</i>	Liver	2gms	0.42
2.	Mammals	<i>Caprine barbari</i>	Liver	2gms	0.57

Table 2:- RNA content from liver tissue of *Heteropneustes fossilis*(Fish) and *Caprine barbari*(Goat)

S.no.	Class	Animal	Organ	Amount	Mean(mg)
1.	Pisces	<i>Heteropneuste sfossilis</i>	Liver	2gms	3.34
2.	Mammals	<i>Caprine barbari</i>	Liver	2gms	2.5

Table 3:- protein content in liver tissue of *Heteropneustes fossilis*(Fish) and *Caprine barbari*(Goat)

S.no.	Class	Animal	Organ	Amount	Mean(mg)
1.	Pisces	<i>Heteropneustes fossilis</i>	Liver	2gms	13.4
2.	Mammals	<i>Caprine barbari</i>	Liver	2gms	12.4

Table 4:- Mean values of DNA, RNA and proteins

S.No	Animal	Mean value DNA (mg)	Milligrams percent (DNA)	Mean value of RNA(mg)	Milligrams percent (RNA)	Mean protein value (mg)	Milligram s percent (proteins)
1.	<i>Heteropneustes fossilis</i>	0.42	0.21	3.34	1.66	13.4	6.4
2.	<i>Caprine barbari</i>	0.57	0.28	2.5	1.25	12.4	6.35

In the present study DNA is 0.42mg, RNA is 3.34mg and proteins are protein are 13.4mg extracted and estimated from 2gms of liver tissue of *Heteropneustes fossilis*(Fish). In *Caprine barbari*(Goat) DNA is 0.57mg, RNA is 2.5mg and proteins are 12.4 mg. extracted and estimated from 2gms of liver tissue. Thus it is seen that in same amount of tissue proteins are more than DNA and RNA. At the molecular level evolution is reflected in protein differences and large number of investigations are directed towards evaluating such differences and research in this area is moving rapidly.

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