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A REVIEW ON BIOCHEMICAL AND THERAPEUTIC ASPECTS OF ARGINASE

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ABSTRACT

One of the prime candidates in the treatment of debilitating human cancers includes group referred to as anticancer enzymes which acts by nutritional starvation therapy. The antitumor activity of these enzymes finds effective in countering wide range of cancers. Nutritional starvation of cancer cells have been tried by many researchers and have been found to be successful in a few cases. *Arginase* comes under this category. *Arginase* was proved to be effective in treatment of human hepatocellular carcinoma, melanoma and prostatic cancer. Apart from anticancer activity, arginase was found to be useful in the treatment of acute neurological disorder, allergic asthma and rheumatoid arthritis. The present review focuses on the isolation, production, assay and media optimization of *arginase* enzyme and its therapeutic uses.

1. INTRODUCTION

Cancer is a disease characterized by abnormal proliferation of cells. It is one of the leading causes of death in the world, and it is of serious problem to people living in both the developing and developed nations ^{1,2,3}. Both benign and malignant tumors manifest as uncontrolled proliferation, but the latter are distinguished by their capacity to dedifferentiate, their invasiveness and their ability to metastasise. Cancer can be cured without much problem if it is diagnosed at the earlier stages. The treatments include chemotherapy, radiation and other forms of therapy. Every therapy has its own advantages and disadvantages. Nutritional starvation of cancer cells have been tried by many researchers and have been found to be successful in a few cases. Cancer cells have absolute requirements for nutrition, a few of its nutrients it can synthesize and for a few others it depends on the host cell. Amino acids like methionine and arginine come under these categories ^{4,5}. If the cancer cells are deprived of these amino acids, they starve to death, since they can't synthesize these amino acids ⁶. Nutritional starvation can be done by two ways, one by controlling the dietary intake of these amino acids and the other by decreasing the serum concentration of these amino acids. A number of enzymes are available which act by the same mechanisms to treat cancers. Examples : *asparaginases*, *glutaminases*, *arginases*, *arginine deaminase* etc. Microorganisms are considered as potential sources of these enzymes . Microbial anticancer enzymes are of considerable interest in view of their specific activity and stability at biological pH. The present review focusses on the isolation ,purification, assay and optimization of anticancer enzymes from microbial sources.

L-Arginase: Arginase (EC 3.5.3.1) is a 105 kD homotrimeric enzyme which requires an intact binuclear manganese cluster for its catalytic activity ⁷. *Arginase* is the terminal enzyme of the urea cycle among the six other enzymes. Interest in the *arginases* as possible regulatory enzymes is growing because of their potential for regulating the availability of arginine for the synthesis of NO, polyamines, agmatine, proline and glutamate ⁸.

Arginase was discovered in mammalian liver by Kossel & Dakin (1904) who showed that the products of its action are ornithine and urea. Since then, the enzyme has attracted a great deal of interest from many points of view. The enzyme has been found to exist in two forms that have evolved with differing tissue distributions, metabolic functions and subcellular locations in mammals ⁹. The cytosolic form, *Arginase I* is found predominantly in the liver or hepatic cells and is important in ureogenesis. *Arginase II* is a mitochondrial enzyme that is extrahepatic and more widely distributed in numerous tissues, for example, kidney, and skeletal muscle ¹⁰. It may be found at lower levels in macrophages, lactating mammary glands, and brain. Important roles of *Arginase II* have been reported to be the biosynthesis of polyamines, the amino acid ornithine, proline and glutamate and in the inflammatory process. Genetic “knockout” experiments suggest that *arginase II* functions in L- arginine homeostasis by regulating L-arginine concentrations for cellular biosynthetic reactions such as nitric oxide biosynthesis. The human type I and type II *arginases* are related by 58% sequence identity and are immunologically distinct.

Mechanism of Action: The *arginases* catalyse the divalent cation-dependent hydrolysis of L-arginine to form the non-protein amino acid L-ornithine and urea ^{11,12}.

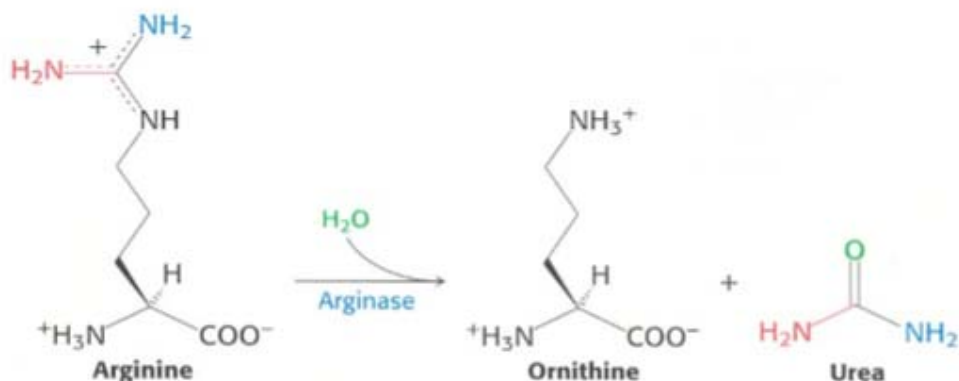
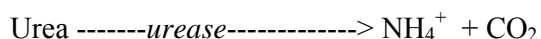


Figure 1 : The arginase reaction.

In the organisms containing *urease*, urea is further converted to ammonia and carbondioxide.



OCCURRENCE AND DISTRIBUTION:

Sources of *L-arginase* : *Arginase* apart from being ubiquitously present in mammalian tissues has also been characterized from various worms, molluscs, fishes, bacteria, fungi, yeast, actinomycetes, algae and plants.

Table no: 1

Bacteria	<i>Mycobacteria</i>	13
	<i>Proteus spp</i>	14
	<i>T.aquaticus</i>	15
	<i>Agrobacterium-Rhizobium group</i>	16
	<i>Cyanobacterium Aphanocapsa</i>	17
	<i>Cyanobacteria,</i>	18
	<i>Bacillus licheniformis</i>	18
	<i>Bacillus subtilis 168</i>	19
	<i>Streptomyces spp.</i>	20
	<i>Streptomyces calvuligerus</i>	21
	<i>Streptomyces calvuligerus</i>	22
	<i>Cyanobacterium Anabaena cycadeae</i>	23
	<i>Thermophile Bacillus caldovelox</i>	24
	<i>Rhodobacter capsulatus E1F1</i>	25
	<i>Bacillus anthracis</i>	26
	<i>Chlamydia pneumoniae</i>	27
Protozoa:	<i>Leishmania, Crithidia and Blastocridhidia</i>	28
	<i>Entamoeba histolytica</i>	29
	<i>Plasmodium falciparum</i>	30

Fungi	<i>Fungi: Neurospora crassa</i>	31
	<i>Aspergillus nidulans</i>	32
	<i>Trichoderma sp.</i>	33
	<i>Agaricus bisporus</i>	34
Lichens	<i>Evernia prunastri</i>	35
	<i>Evernia prunastri and Xanthoria parietina thalli</i>	36
	<i>Peltigera canina</i>	37
	<i>Leptogium corniculatum</i>	38
Yeast	<i>Saccharomyces cerevisiae</i>	39
	<i>Schizosaccharomyces</i>	40
Plant sources	<i>Lathyrus sativus</i>	41
	<i>pumpkin seeds</i>	42
	<i>grapevine- Vitis vinifera</i>	43
	<i>Arachis hypogea</i>	44
	<i>Canavalia lineata</i>	45
	<i>Soybean</i>	46
	<i>Jack bean (Canavalia ensiformis)</i>	47
	<i>kiwifruit vines- Actinidia deliciosa</i>	48
	<i>, ginseng (Panax ginseng</i>	49
	<i>loblolly pine (Pinus taeda L.)</i>	50
	<i>Saccharum officinarum cv. Mayari</i>	51
	<i>tobacco</i>	52
	<i>Lycopersicon esculatum (tomato)</i>	53
	<i>Quercus ilex</i>	54
Insects	<i>Vigna catjang</i>	55
	<i>Hyalomma dromedarii</i>	56
	<i>Drosophila</i>	57
Sea organisms	<i>Bombyx mori- the silk worm</i>	58
	<i>Larva of Phoronis pallida</i>	59
	<i>African snail-Achatina fulica</i>	60
	<i>Protopterus aethiopicus and Protopterus annectens</i>	61
	<i>South American fish pacu (Piaractus mesopotamicus)</i>	61
	<i>South American fish pacu (Piaractus mesopotamicus)</i>	62
	<i>Sparus aurata</i>	63
	<i>Antarctic fish Notothenia rossii and Notothenia neglecta</i>	64
	<i>Pacific spiny dogfish shark Squalus acanthias</i>	65
Mammals	<i>Salamandra salamandra</i>	66
	<i>Genypterus maculatus</i>	67
	<i>Mus booduga</i>	68
	<i>Raja erinacea</i>	69
	<i>(Felis catus)</i>	70
	<i>Rana temporaria</i>	71

Production, Characterisation & Purification of *arginase* from various sources: Numerous reports regarding the characterization of *arginase* & its purification ranging from microbial sources to vertebrate and invertebrate sources are accessible in scientific literature.

Production of *arginase* : Arginine has been reported to be utilized as a nitrogen source in *Neurospora crassa*⁷². Mora et al., grew *Neurospora crassa* in minimal medium supplemented with 1.5% sucrose and reported that arginine synthesised from exogenous citrulline was not effectively utilized as exogenous arginine due to feedback inhibition of *arginase* in vivo and in vitro by arginine pool formed from citrulline⁷³. *Saccharomyces cerevisiae* ATCC 9763 was cultured aerobically at 30°C in defined media containing arginine as a sole source of nitrogen. After 5 hrs of growth, *arginase* was extracted by sonication and partially purified by gel filtration⁷⁴. Most of the arginine pool in *Neurospora* is available inside the vesicle and lesser arginine concentration is found to be present in cytosol in cells growing in minimal medium⁷⁵. In arginine supplemented medium, the cytosolic pool dramatically increases resulting in induction of catabolic enzyme *arginase* as well as rapid catabolism of arginine. Vaca and Mora tested the effect of various nitrogen sources on arginase production in a *N. crassa* mutant ure-1 lacking urease activity and found that arginase was hyperinduced with arginine as the nitrogen source whereas in the wild-type strain the induction was completely repressed by glutamine⁷⁶. In *Bacillus licheniformis* the *arginase* pathway enzymes are subject to strong catabolite repression during growth on glucose while glutamine antagonizes the induction of *arginase* pathway. This effect occurs only in a media containing a good carbon source. In this organism glutamine is a better nitrogen source than arginine, glutamate or ammonia. Broman et al., demonstrated that *Bacillus licheniformis* has two pathways of arginine catabolism and in well aerated cultures the *arginase* route is present, and levels of catabolic ornithine carbamoyl transferase were low⁷⁷. They also demonstrated that an *arginase* pathway deficient mutant, BL196, failed to grow on arginine as a nitrogen source under these conditions.

The intensity of carbon source (glucose, citrate) catabolite repression of *arginase* seems to be much weaker in *B. subtilis*. In the *arginase* pathway in the *B. subtilis* strain used by Baumberg and Harwood repression of the pathway was by glutamine and, in addition, ammonia while ornithine and citrulline were useful in induction of the enzyme²⁰. In the genus *Agrobacterium*, arginine may be used as a nitrogen source by an inducible *arginase* and a constitutive urease. Many *Agrobacterium* strains are also able to use arginine and ornithine as carbon source. However, many *Agrobacterium* strains unable to grow on arginine or ornithine as a carbon source display this ability when they harbor a wild-type octopine or nopaline Ti plasmid⁷⁸. *Synechocystis* sp. PCC 6803 were grown axenically in BG11 (nitrate containing) medium⁷⁹. Cultures were grown 30°C in light with shaking (80-90 rpm) for liquid cultures and supplemented with 1mM filter sterilized L-arginine. Drainas and Weiss (1980) studied the effect of various carbon sources on *arginase* activity in *Neurospora crassa* and found that basal and induced levels of *arginase* were reduced in the following order sucrose, acetate, glycerol and ethanol with regard to carbon sources used⁸⁰. They also reported that arginine pools were similar regardless of carbon source in mycelia grown in arginine-supplemented medium and arginine degradation was proportional to level of *arginase* in both sucrose and glycerol grown mycelia suggesting a possible carbon metabolite effect on arginine metabolic enzyme genes in the fungus. Schreier et al., (1982) studied the regulation of inducible nitrogen catabolic enzyme *arginase* in *B. licheniformis* stating that although levels of enzyme were nitrogen source (Ammonia, Glutamine, Urea, Arginine, Ornithine) dependent its induction could still occur even in presence of preferred nitrogen sources⁸¹. *Neurospora crassa* strain was grown at 30°C in Vogel's minimal medium (VM) with 1.5% sucrose as carbon source. Initially it was supplied with 5mM arginine, 1mM histidine or NH₄NO₃,

later these were replaced with 5mM arginine which acted as nitrogen source. Liquid cultures were inoculated with 106 conidia per ml of media in baffled flasks and shaken at 250rpm⁸². Two forms of arginine with differing molecular weights i.e., 36 kDa and 41 kDa were produced in unsupplemented and arginine-supplemented media respectively.

Purification and kinetic characterization of arginase: Rabbit liver *arginase* was purified by Breitburd and Orth by chromatographic techniques of DEAE cellulose and Sephadex G-200⁸³. It was strongly inhibited by EDTA and lost activity in absence of Mn²⁺ ions. SDS-PAGE showed that enzyme had a subunit molecularweight of 36,500, kDa increasing probability of its tetrameric nature. Molecular characteristics of chicken liver *arginase* were accounted by Grazi and Magri⁸⁴. Purification and properties of rat kidney *arginase* were studied by Kaysen and Strecker showing the enzyme was highly stable at high temperatures such as 60°C and had a Km value of 18 mM and was strongly inhibited by borate and L-ornithine⁸⁵. Tarrab et al., studied molecular isoforms of rabbit liver *arginase* by purifying them in a sequential manner by CM cellulose chromatography⁸⁶. Purification studies of *arginases* from human-leukemic lymphocytes and granulocytes were done by Reyer and Dorner employing purification procedure of acetone extraction, ammonium sulphate precipitation, DEAE-cellulose, CM-Sephadex chromatography and gel filtration on Bio-Gel A⁸⁷. Both the *arginase* proteins were strongly basic with pI values between 9.25 and 9.35. Rat small intestinal *arginase* was purified by Sephadex -100 filtration and shown to have optimum pH of 10.0 and Km of 19 mM by Fujimoto et al⁸⁸. The enzyme was almost completely inactivated by treatment with EDTA. The isoforms differed in their kinetic and pH behavior. *Arginase* from human liver and erythrocytes was isolated by Beruter et al., and was purified using DEAE-cellulose chromatography followed by gel filtration on Sephadex G-200⁸⁹. The human liver *arginase* had a molecular weight of was 107, 000 and its Km for arginine was 10.5 mM. The properties of human heart *arginase* studied by Baranczyk et al., revealed its Km to be 5 mM and molecular weight of about 30, 000 kDa with the enzyme having strong dependence on Mn²⁺ ions for its activity⁹⁰.

Gopalakrishna and Agarajan reported the purification of arginase from rat fibrosarcoma by heat treatment followed by DEAE-cellulose chromatography and gel filtration on Sephadex G-100 and accounted that the Km for arginine was 11 mM and had a pH optimum of 10⁹¹. Purification and physical properties of *Arginase* from *Xenopus laevis* liver were studied by Peiser and Balinsky using procedures of heat treatment, acetone fractionation, and isoelectric focusing and its molecular subunit weight was determined by gel filtration on Sephadex G200 and found to be 76,000 daltons⁹². *Arginase* was obtained from Iris hollandica bulbs and purified approximately using DEAE-Sephacel chromatography, aminohexyl-Sepharose 4B chromatography and gel filtration on Ultrogel AcA 34 by Boutin (1982)⁹³. Constitutive *arginase* from *Evernia prunastri thallus* was purified 920-fold and is activated by endogenous L-arginine⁹⁴. Borkovich and Weiss in 1987 reported that *Neurospora crassa arginase* has a subunit weight of 38,300 determined by SDS-PAGE³¹. The enzyme exhibited hyperbolic kinetics at pH 9.5 with an apparent Km for arginine of 131mM. Antiserum was prepared against the purified enzyme and two proteins in the extracts of *S. cerevisiae* were detected that were weakly cross- reactive with the antiserum. Two forms of *arginase* were isolated from human erythrocytes and their immunological properties were studied by Kedra et al⁹⁵. Later, Ikemoto et al., (1989) developed an efficient method involving hydrophobic chromatography and immunoaffinity chromatography for purification of human erythrocyte *arginase*⁹⁶. Patil et al., (1990) gave a high recovery protocol for purification of ox erythrocyte *arginase* involving heat treatment, CM and DEAE-Sepharose chromatography, arginine AH-Sepharose chromatography and molecular sieving through Biogel P-150⁹⁷. The optimum pH was 11.5 and temperature was 55°C for purified enzyme. Soyabean Glycine max

axes *arginase* was purified by Kang and Cho and its properties were studied ⁴⁶. The enzyme purification procedure followed was ammonium sulphate precipitation followed by chromatography on Sephadex G- 200, DEAE Sephacel, Hydroxypatite and Arginine-Sepharose 4B affinity column. The enzyme had a K_m of 83 mM and pH optimum of 9.5 with a molecular weight of 240, 000 obtained by pore-gradient electrophoresis. Purification of recombinant human liver *arginase* expressed in *E. coli* was described by Ikemoto et al., by chromatographies on CM-Sephadex G-150, DEAE-cellulose and Sephadex G-150 followed by preparative gel electrophoresis ⁹⁸. The purified enzyme was a monomer of molecular weight 35,000 kDa. Singh and Singh, purified liver *arginase* from teleostean fish-*Clarias batrachus* having K_m of 15.38 mM for arginine and an optimum pH of 9.5 ⁹⁹. Ornithine and leucine act as competitive inhibitors whereas valine and isoleucine act as non-competitive inhibitors with respect to L-arginine as substrate. Purification of *arginase* from *S. cerevisiae* was done by Sepharose chromatography and TSK-G3000 gel filtration followed by its characterization (Green et al., 1989) which revealed a K_m of 15.7 mM for arginine and requirement of Mn^{2+} ions for activity. CD spectra analysis showed significant spectral changes from removal of bound metal and dialysis against EDTA ¹⁰⁰.

A thermostable *arginase* from the extreme thermophile *Bacillus caldovelox* was purified by chromatographic techniques and its kinetic properties were studied in 1991 by Patchett et al ²⁴. Its activity is optimal at pH 9 and temperature of 60°C. The K_m for arginine is 3.4 mM. *Arginase* obtained from the phototrophic bacterium- *Rhodobacter capsulatus* E1F1 was purified and studied for its properties. The molecular parameters and kinetic constants of *Rhodobacter arginase* resembled the *Saccharomyces cerevisiae* enzyme rather than those of bacterial *arginases* ²⁵. Extrahepatic arginase (ArgII) was isolated from rat mammary gland and its properties were investigated by Jenkinson and Grigor (1994) ¹⁰¹. This enzyme had a pH optima of 10 and K_m of 12-14mM. Purification of human hepatic *arginase* and its manganese (II) - dependent and pH-dependent interconversion between active and inactive forms was described by Kuhn et al., (1995) who showed that both the maximal velocity of catalysis and the K_m toward arginine were markedly pH dependent in the physiological range ¹⁰². In situ characterization of *Helicobacter pylori arginase* employing NMR spectroscopy, spectrophotometry, radiotracer analysis and protein purification techniques was done by Mendz et al ¹⁰³. A K_m of 22+/- 33mM was determined for the enzyme activity and differences of V_{max} were observed between strains. Divalent cations stimulated *arginase* activity & most potent activators were $Co^{+2} > Ni^{+2} > Mn^{+2}$.

Recombinant human arginase II was expressed in *E. coli* and purified to homogeneity by Colleluori et al., (2001) ¹⁰⁴. The enzyme reportedly had a K_m of 4.8 mM at physiological pH and exists primarily as a trimer. Borate was a non-competitor inhibitor of the enzyme and ornithine which is an inhibitor of *Arginase-I* is not an inhibitor of the type-II enzyme showing that isozyme selectivity occurs between both forms with regard to substrate and product as well as inhibitor binding. *Pinus taeda arginase* was purified by chromatographic separation on DE-52 cellulose, Matrex Green and arginine-linked Sepharose 4B and had a molecular mass of 140 kDa deduced by FPLC while the subunit size was shown to be 37 kDa observed by SDS-PAGE analysis ⁵⁰. *D-arginase* obtained from *Arthrobacter sp KIJ 8602* was purified by anion exchange chromatography using DEAE-Toyopearl 650M followed by hydroxypatite chromatography and gel filtration chromatography with Superdex 200 ¹⁰⁵. The purified enzyme was proposed to be a homohexamer with each subunit of approximately 40, 000 determined by SDS-PAGE. It had an optimum pH of 9.5 and needed Zn^{2+} for activation instead of Mn^{2+} . Bovine liver *arginase* as reported by Wheatley et al., (2003) is a remarkably heat resistant enzyme with a very long life on storage at 4°C in lyophilised form and is occasionally marginally more active at pH 7.2 than at pH

9.9¹³⁷. *Helicobacter pylori arginase* expressed in *E. coli* was loaded onto polypropylene columns containing nickel- nitrilotriacetic acid agarose resin for purification (McGee et al., 2004)¹⁰⁶. The purified enzyme had significant activity with cobalt as cofactor and had acidic pH optima of 6.1. It was inhibited by low concentrations of reducing agents. Dabir et al., (2005) described the purification, properties and alternate substrate specificities of the enzyme from two different sources: *Vigna catjang* cotyledon and buffalo liver⁵⁵. The Michealis-Menten constant for cotyledon arginase and hepatic arginase were found to be 42 mM & 2 mM respectively and their pH-optima were 10 & 9.2 respectively. Munder et al., (2005) have reported that arginase I is constitutively expressed in human granulocytes and participates in fungicidal activity¹⁰⁷. Two *arginases* Arg I and Arg II were separated from *Fasciola gigantica* by purification on a DEAE-Sephacrose column as reported by Mohamed et al., (2005)[108]. Further purification was carried out for Arg II and it was shown that the enzyme was activated by Mn²⁺ and inhibited by Fe²⁺, Ca²⁺, Hg²⁺, Ni²⁺, Co²⁺ and Mg²⁺ ions. Purification of recombinant *Plasmodium falciparum arginase* was done on Superdex S-200 column and its activity was found to be dependent on Mn²⁺. Ni²⁺ when replaced with Mn²⁺ resulted in four-fold loss of activity³⁰. Viator et al., (2008) reported the purification of recombinant *Bacillus anthracis arginase* expressed in *E. coli* XL1-Blue MRF' on *arginase* affinity columns¹⁰⁹.

Structure related studies

Most of the studies related with the structure of *arginase* are by means of studying the interaction of various inhibitor complexes with the substrate specific sites of the enzyme. Electron paramagnetic resonance (EPR) studies on rat liver arginase by Reczkowski and Ash revealed that fully Mn-activated *arginase* contains two Mn²⁺ ions per subunit and these form EPR spin-coupled binuclear centers¹¹⁰. The high resolution X-ray crystal structure of rat *arginase I* was provided by Kanyo et al., which illustrated that the enzyme is a 105 kDa homotrimer with each subunit of 35 kDa containing a spin-coupled binuclear manganese cluster critical for its activity located at the bottom of a ~15 Å -deep active site cleft¹¹¹. Bewley et al., studied the crystal structure of *arginase* structure of inactivated *Bacillus caldovelox arginase*-L-arginine complex which revealed that the structural basis for substrate and inhibitor specificity are an array of direct and water mediated hydrogen bonds saturating all four acceptor positions on α -carboxylase group and all three donor positions on α -amino group¹¹². Cox et al., noted the binding of N ω -hydroxy-L- arginine (NOHA)-an intermediate of NO biosynthesis and modest inhibitor of the enzyme with rat *arginase I* which revealed that the N ω -hydroxyl group displaces the metal-bridging hydroxide ion and bridges the binuclear manganese cluster and that in the binding of rat *arginase I* and nor-NOHA the N ω -hydroxyl group of the inhibitor displaces the metal bridging hydroxide ion¹¹³. The structure for human kidney II type *arginase* is nearly identical to that of rat *arginase I* due to conservation of all metal ligands between the two sequences and its polypeptide fold is topologically identical to α/β fold of rat *arginase I* and hexameric *arginase* from *Bacillus caldovelox*¹¹⁴. Crystal structure of human arginase I at 1.29- angstrom resolution has been studied by Di-Costanzo et al.,¹¹⁵. Inhibitor complexes of *arginase I* with inhibitors ABH and BEC were studied to know the inhibition of the enzyme in human and urine myeloid cells to study its effect on immune response. The role of the hyper- reactive histidine residue (H141) in rat liver *arginase* was studied by studying the structure of the enzyme by chemical modulation, mutagenesis and X-ray diffraction¹¹⁶. By the study, the H141 has been shown to possess conformational mobility supported by its proton shuttling role and proposed to be an acid/base catalyst by deprotonating the metal-bridging water molecule to generate the metal-bridging hydroxide nucleophile, and by protonating the amino group of the product to facilitate its departure. Di Costanzo et al., determined the X-ray crystal structure of a perdeuterated human *arginase I* complexed with 2(S)-

amino-6- boronohexanoic acid (ABH) at 1.90 Å resolution and found that the perdeuteration did not cause any structural and functional changes when compared with the unlabeled enzyme thus rendering the perdeuterated crystals suitable for neutron crystallographic study¹¹⁷. Dowling et al., (2010) have reported the 2.15 Å resolution crystal structure of arginase from the cerebral malarial parasite *Plasmodium falciparum* in complex with the boronic acid inhibitor 2(S)-amino-6-boronohexanoic acid (ABH) - the first report of crystal structure of a parasitic *arginase*¹¹⁸. Inhibition studies with the enzyme structure lend important insights into the antimalarial therapy against liver-stage infection, and ABH may serve as a lead for the development of inhibitors.

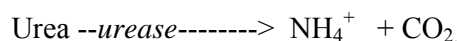
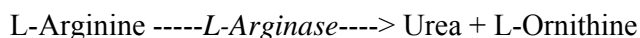
Assay procedures for Arginase :

Numerous methods based on different principles for assaying *arginase* are available. Some of them are listed below:

Colorimetric Method: These methods are based on the colorimetric determination of urea or ornithine- the products of arginase reaction which can be assayed by the development of a colorimetric procedure. In the procedure of urea determination (Foster et al., 1971) - urea is determined using 2,3 – butanedione reagent (BUN reagent)¹¹⁹. This method has been reported for arginase assay (Basch, 1997)¹²⁰. The colour is measured at 460 nm. In the method of Coulombe and Favreau, (1963) urea formed gives red colour on reaction with diacetyl monoxime and thiosemicarbazide in acid medium whose absorbance is measured at 535 nm¹²¹. In the method of Davis and Mora, (1968), the urea formed is assayed by reaction with α- isonitrosopropiophenone¹²². This procedure was originally devised for high accuracy urea determination in blood filtrates and urine by Archibald (1945)¹²³. The absorbance change is read at 540 nm. A novel colorimetric assay was devised by Aminlari (1992) where determination of residual arginine, after its conversion with p-nitrophenyl glyoxal (PNPG) at pH 9.0 in the presence of sodium ascorbate was monitored¹²⁴. The decrease in absorbance in the presence of arginase correlates with the enzyme activity. A colorimetric microplate assay for *arginase* was recently developed by Iyamu et al., (2008) for high-throughput analysis of *arginase* activity in vitro¹²⁵. The method was a modified form of the Chinard reaction of ornithine-ninhydrin reaction.

Electrochemical Method:

Booker and Haslam (1974) used the following reaction sequence for assay of *arginase* by means of a cation NH_4^+ electrode that was used to sense the NH_3 liberated and immobilized urease was used¹²⁶. From 1.6 to 16 U of *arginase* could be assayed in 10 min with a CV of 3%.



Larsen et al. (1975) have described a kinetic method for *Arginase* measuring the NH_3 liberated with an air gap NH_3 electrode^[127]. The above sequential reactions liberate NH_3 whose determination as NH_4^+ is done by the electrode. The determinations were based on monitoring the initial reaction rates of the selective release of ammoniacal nitrogen.

A tenfold excess of *urease* ensured direct proportionality between the rate of NH_3 production and the *arginase* activity. The coefficient of variation (CV) was 2.8%.

Spectrophotometric Method: Ward and Srere reported a novel *arginase* spectrophotometric assay by based on the principle that the absorbancy of *arginase* below 2100 Å is larger than the combined absorbancies of ornithine and urea ¹²⁸. A cleavage of arginine catalyzed by the enzyme thus results in a net decrease in absorbancy at these wavelengths, allowing a rapid and accurate assay for *arginase* activity. Ozer developed a new spectrophotometric assay for *arginase* in which the enzyme is coupled to urease and glutamate dehydrogenase and the decrease in absorbance at 340 nm due to the oxidation of NADPH is followed ¹²⁹. Han and Viola (2001) developed an alternative assay for *arginase* by synthesising an alternative substrate of *arginase* has been synthesized in which the bridging guanidium nitrogen has been replaced with a sulphur ¹³⁰. This thiol compound is a good substrate for *arginase*, leading to urea and 2-amino-5-mercaptovaleric acid which can be monitored with DTNB to produce a disulfide adduct and release 2-nitro-5-thiobenzoate as a chromophoric product.

Fluorimetric Method: Orfanos et al., described a microfluorometric method for assay of *arginase* activity in dried bloodspots on filter paper ¹³¹. The urea formed from the *arginase* reaction is determined fluorimetrically by oxidation of NADH to NAD⁺ in a coupled kinetic reaction.

Conductimetric method: Macholan et al., have constructed a bioelectrode consisting of two parallel noble metal nets and a thin layer of gel- entrapped urease (for urea determination) and microbial cells or tissue slice (for *arginase* determination) for conductimetric measurements in tissue extracts ¹³².

Radioisotopic Method: This method was reported by Carulli et al., with an underlying principle of hydrolysis of labeled substrate (arginine-guanido-14C) ¹³³. The labeled urea produced is hydrolyzed by urease with the measurement of released ¹⁴CO₂ in a liquid scintillation counter as a measure of *arginase* activity. Advantages of the method reported were sensitivity and accuracy derived from the absence of interference by nonradioactive urea in the incubation mixture.

Volke et al., reported a simple and fast HPLC method with radiochemical detection for measurement of arginase activity by use of radiolabeled arginine ¹³⁴. de-Bono et al., (2007) performed radiochemical HPLC detection of arginine metabolism resulting in measurement of nitric oxide synthesis and *arginase* activity in vascular tissue ¹³⁵.

Other methods: Bastone et al., carried out measurement of *arginase* along with argininosuccinate lyase both in liver and erythrocytes, by using a commercial amino acid analyzer ¹³⁶. The two different substrates used by both the enzymes give the similar product ornithine whose measurement is done by this method. Assay kits such as QuantiChrom™ *Arginase* assay kit for determining *arginase* activity are commercially available in market.

ARGINASE AND ITS DEVELOPMENT AS A CHEMOTHERAPEUTIC AGENT:

The use of *Arginase* as a potential chemotherapeutic agent has shown a lot of potential and promise. *Arginase* treatment of cultured HeLa, human diploid fibroblasts and L1210 cells proved to be as efficient as the use of AFM (Arginine Free Medium) by reducing arginine in the medium to micromolar levels within 5-30 minutes resulting in cell death in both the cultures ¹³⁷. Remission of hepatocellular carcinoma was achieved by arginine depletion through endogenous human hepatic *arginase* released from transhepatic arterial embolization ¹³⁸. Using

state-of-the-art DNA technology, researchers of the Hong Kong Polytechnic University (PolyU) produced a human recombinant *arginase* as a novel method in the treatment of liver cancer by arginine depletion ¹³⁹. The combination of the recombinant *arginase* with an anti-neoplastic agent 5 fluorouracil (5FU) for treatment of human malignancies was tested in nude mice bearing an ADI-resistant HCC xenograft and this treatment methodology was shown to be effective for arginine depletion ¹⁴⁰. BCT-100 pegylated recombinant human *arginase* manufactured by large scale fermentation of a recombinant *B. subtilis* strain LLC101 has been reported as a novel anti-melanoma agent for treatment of human melanoma cells as reported by Hsueh et al.¹⁴¹. Pegylation of recombinant human *arginase* (rhArg-peg 5000mw) produced in *B. subtilis* expression system was done and the pegylated enzyme has been shown to have in vitro and in vivo anti-proliferative potential and apoptotic activities in human hepatocellular carcinoma (HCC) by Cheng et al.¹⁴². While studies for conducting clinical trials of PEG BCT-100 are reported to be in process to assess its safety and efficacy in humans (<http://clinicaltrialsfeeds.org/clinical-trials/show/NCT00988195>), recently further studies on pegylation of recombinant *arginase* conducted by the Cheng group by conjugation of rhARG with methoxypolyethylene glycol-succinimidyl propionate (mPEG-SPA 5,000) has been reported as having comparable anti-tumor efficacy to native rhArg1[143]. That the inhibition of human hepatocellular carcinoma by recombinant *arginase* (rhARG1) is by the inducing cell cycle arrest at the G2/M or S phase, possibly mediated by transcriptional modulation of cyclins and/or cyclin dependent kinases (CDKs) has been observed by Lam et al.¹⁴⁴. This group has further proposed the use of rhArg1 alone or in combination with chemotherapeutic drugs for treatment of liver cancer.

In vitro cytotoxicity of human *arginase I* by replacing the two Mn²⁺ ions normally present in the enzyme with Co²⁺ significantly lowered the Km value of the enzyme, increased its serum stability and showed incredible ability to eliminate human hepatocellular carcinoma and melanoma cell lines proving it to be a capable new contender for treatment of L-Arg auxotrophic tumors ¹⁴⁵. A study conducted by Hernandez et al., published just a few months back in the prestigious journal 'Blood' reports of the potential therapeutic role of pegylated *Arginase I* in the treatment of adult patients with acute lymphoblastic T cell leukemia (T-ALL) through arginine depletion ¹⁴⁶.

Recombinant Arginase

Development of recombinant *arginase* has been an intriguing subject of research worldwide. Molecular cloning and nucleotide sequence of cDNA for human liver *arginase* facilitated the investigation of the enzyme and gene structures and helped in elucidating the nature of mutation in argininemia. *Arginase* activity was detected in *Escherichia coli* cells transformed with the plasmid carrying lambda hARG6 cDNA insert by Haraguchi et al ¹⁴⁷. cDNA phARG6 for human liver *arginase* was used for expression of human liver *arginase* in *E. coli* strain KY1436 by Ikemoto et al ⁹⁸. This *E. coli* expressed human liver *arginase* had chemical, immunological and most catalytic properties indistinct from purified human erythrocyte *arginase*. The cloning, expression and crystallization of a thermostable *arginase* from the thermophilic bacterium '*Bacillus caldovelox*' has been discussed by Maria et al ¹⁴⁸. In this

study, the expression of recombinant *arginase* at high levels was achieved in *E. coli* using an inducible T7 RNA polymerase based system. Molecular cloning and nucleotide sequence of the *arginase* gene of *Bacillus brevis*.

TT02-08 and its expression in *Escherichia coli* has been described by Shimotohno et al.¹⁴⁹. The *B. subtilis arginase* encoding rocF gene was cloned and expressed in *E. coli* K-12 for enhanced production of urea by *arginase* pathway by Tuchman et al.⁵⁰. Modulation of mRNA levels of liver *arginase* by insulin and vanadate in experimental diabetes has been studied by Salimuddin et al., (1999) where it was shown that an increase in *arginase* activity and mRNA levels in diabetes and decrease in treated animals may be due to the transcriptional regulation of *arginase* gene¹⁵¹. Kimura et al., transformed *E. coli* strain KY1436 by pTAA12 expression plasmid vector containing human liver-type *arginase* cDNA and developed an ELISA system by using this recombinant enzyme as an antigen for the detection of anti-*arginase* antibodies in patients with Autoimmune Hepatitis (AIH)¹⁵². The expression, purification and characterization of human type II *arginase* involving genetic engineering experiments to develop a recombinant human type II *arginase* expressed in *Escherichia coli* was done by Colleluori et al.¹⁰⁴. The recombinant enzyme was studied for its kinetic properties and inhibitor effects on it. The findings indicated that isozyme selectivity exists between type I and type II *arginases* for binding of substrate and products, as well as inhibitors. The rocF gene encoding *arginase* in *Helicobacter pylori* was purified and expressed in *E. coli* by McGee¹⁰⁶ and was found to confer *arginase* activity to *E. coli*. The transformed cells expressing *arginase* showed enhanced enzyme activity than the native *Helicobacter* enzyme. Recombinant *arginase* was developed by cloning DNA of *Plasmodium falciparum* (*Pfarginase*) into *E. coli* BL21. The recombinant enzyme had a Km of 13±2 mM, required Mn²⁺ for activity and had high thermal stability³⁰. The RocF gene of *Bacillus anthracis* was cloned in *E. coli* for development of a recombinant enzyme. The metal preference of the enzyme was shifted at pH 6 from Ni>Co>Mn to Ni>Mn>Co at pH 9. A novel recombinant *B. subtilis* prophage strain LLC101 was constructed for recombinant enzyme production by Cheng et al.¹³⁸. The recombinant enzyme produced was used for treatment of human malignancies by subsequent pegylation^{138,142,143}. A highly active recombinant *arginase* was obtained by expressing *arginase* gene from *Leishmania (Leishmania) amazonensis* in *E. coli* BL21 (DE3) cells¹⁵³. Antibody against the recombinant protein confirmed a glycosomal cellular localization of the enzyme in promastigotes. RhArg causing significant cytotoxicity in LNCaP, DU-145 and PC-3 prostate cancer cells was shown by Eddy C Hsueh et al.

Modification of Arginase: Immobilization & Chemical Modification

Attempts to increase the activity and stability of the enzyme by various techniques of immobilization and chemical modification have been made. Vanillin polymethacrylates (vanacryls) were used for immobilization of *arginase* by Brown and Joyeau. Carvajal et al., reported the immobilization of human liver *arginase* by attachment to nylon with glytaraldehyde as a cross-linking agent¹⁵⁴. The immobilized enzyme tetrameric enzyme dissociated into monomers by treatment with EDTA. Influence of immobilization on solid matrix on rat liver *arginase* revealed that the immobilized enzyme was more resistant to effect

of inhibitors, denaturing agents such as SDS and chelating agents such as EDTA than free enzyme¹⁵⁵. Bovine liver *arginase* was covalently immobilized by glutaraldehyde method to inner surface of Cuprophane hollow fibers of a conventional hemodialyzer by Rossi et al.¹⁵⁶. The immobilization method did not harmfully affect the physical and mechanical properties of neither hollow fibers nor their hemocompatibility. Aguirre and Kasche, by covalently coupling the enzyme to Sepharose beads and dissociating the resulting matrix-bound tetramer by acid or EDTA treatment, obtained catalytically active monomer forms of immobilized rat liver *arginase*¹⁵⁷. Savoca et al., studied the therapeutic effectiveness of *arginase* in cancer therapy on a Taper liver tumor and the L5178Y murine leukemia by modifying it by covalent attachment of polyethylene glycol. The pegylation of the enzyme increased its stability¹⁵⁸. Beef liver *arginase* was modified by the covalent linking of monomethoxypolyethyleneglycol molecules by Visco et al.¹⁵⁹. The derivative enzyme had more convenient properties for a therapeutic use, such as increased structural stability, decreased digestion by proteolytic enzymes and an expanded clearance time in rats. Veronese et al., discussed the immobilization of *arginase* by radiation induced polymerization of induced monomers¹⁶⁰. Properties of *arginase* immobilized in a fibrin clot were studied by Diez et al.³⁷. That chemical modification of rat liver *arginase* by N-bromosuccinimide leads to its inactivation was reported by Daghighi et al.¹⁶¹. *Arginase* isolated from beef liver was covalently attached to a polyacrylamide bead support bearing carboxylic groups activated by a water-soluble carbodiimide¹⁶². The pH optimum for the catalytic activity was pH 9.5 and apparent temperature maximum was 60° C. Immobilization markedly improved the conformational stability of the enzyme. An enzyme reactor for determination of L-arginine was described by Alonso et al., by immobilization on an epoxy resin matrix¹⁶³. Pegylation has been reported to greatly improve the stability of the enzyme¹³⁷. Pegylated recombinant human *arginase* has in vitro and in vivo anti-proliferative potential and apoptotic activities in human hepatocellular carcinoma by Cheng et al.^{138,142}. For studying thermodynamics of nor NOHA/*arginase* binding and the role of the reactive histidine residue *arginase* was immobilized on a chromatographic support by Bagnost et al.¹⁶⁴. Pegylation of rhARG with methoxypolyethylene glycol-succinimidyl propionate (mPEG-SPA 5,000) proved to be the best for activity retention of the immobilized enzyme for use in arginine depletion as an anti-cancer therapy protocol¹⁴³. *Leishmania arginase* was immobilized on a Ni²⁺ resin and its activation kinetics was studied by varying Mn²⁺ concentrations at a temperature of 23°C. Conformational changes were observed when enzyme interacted with Ni²⁺ present in the column¹⁶⁵. Konst et al., have reported the stabilization and immobilization of *B. subtilis arginase* on commercially available epoxy-activated supports out of which immobilization on Sepabeads EC-EP was most promising¹⁶⁶. The immobilized enzyme was employed in the biobased production of nitrogen-containing chemicals as an alternative to the petrochemical production.

Arginase: The multilateral medical therapeutic

The enzyme has been found to possess profound therapeutic benefits in treatment of various physiological disorders in the body. Measurement of circulating *Arginase I* i.e., serum *arginase* levels have been used experimentally as rapid marker for liver injury¹⁶⁷. *Arginase* has been found to be essential for the treatment of acute neurological disorders¹⁶⁸. Ornithine, produced

by *arginase* is necessary for the production of collagen, is helpful in therapy of rheumatoid arthritis¹⁶⁹. *Arginase* upregulates the synthesis of polyamines and proline via arginine hydrolysis thus, being necessary to provide compounds for cell proliferation and growth¹⁷⁰. *Arginase* competes with NOS for endogenous arginine pools, this way its levels acts as an indirect regulator of penile and vaginal flow thus playing an important role in male and female sexual arousal¹⁷¹. Upregulation of endogenous *arginase I* causes the activation of neural regeneration pathways, the reaction being mediated by polyamines and leading to novel roles of *arginase* in cell survival, regeneration and translation in the central nervous system¹⁷². Munder et al., reported that *arginase-I* is constitutively expressed in human granulocytes and participates in fungicidal activity by a novel antimicrobial effector pathway likely through arginine deprotonation in phagolysosome¹⁷³. A deficiency of the liver enzyme results in hyperargininemia- inherited in an autosomal recessive manner¹⁷⁴. Raised arginase activity in serum of 85% patients suffering from colorectal cancer liver metastases (CRCLM) led to the conclusion that *Arginase* can be a useful marker for diagnosis of CRCLM¹⁷⁵. Human granulocyte *arginase* has been reported to be a promising pharmacologic treatment to reverse unwanted immunosuppression by Munder et al¹⁷⁶. Increase in psychological stress results in raising NO level which results in poor sperm quality. Thus, *arginase* levels are important to keep NO levels (via arginine degradation) in control for maintenance of semen quality¹⁷⁷. *Arginase* has been shown to be effective for treatment of Hepatitis- B¹³⁹. A finding that Hepatitis C virus targets overexpression of *arginase I* in hepatocarcinogenesis suggests that insights into this disease and its control may be through altering arginine metabolism by means of *arginase* activity alterations¹⁷⁸.

The *arginase* levels in NOS related processes prove to be important for pathogenesis of a host of diseases relevant to arginine-NO balance in the body. Due to *arginase* and nitric-oxid synthase (NOS) utilizing the same substrate arginine, a reciprocal regulation and metabolic balance exists between both of them for numerous processes that are dependent on these enzymes, occur inside the body. In rheumatoid arthritis patients a significant correlation between serum concentration of *arginase* protein and rheumatoid factor has been found suggesting that increased *arginase* production may play an important role in pathogenesis of this disease⁴⁹. *Arginase I & II* levels rise remarkably in asthma and lung infection thus, showing the involvement of *arginase* in the pathology of the disease¹⁷⁹. Measurements of plasma *arginase* activity provide a useful marker for underlying metabolic disorder and efficacy of treatment for asthma¹⁸⁰. Increase in levels of arginase during Sickle cell disease makes it a candidate for pathogenesis of this disease¹⁸¹. The pathogenesis of cystic fibrosis has been reported to be *arginase* dependent since *arginase* levels are enhanced leading to depletion of arginine for NOS action to occur contributing to nitric oxide deficiency¹⁸². Because of its role in epidermal hyperproliferation by upregulating CAT-1 expression in psoriatic skin, *arginase* is important in pathophysiology of psoriasis¹⁸³. Similarly, since nitric oxide is a chief vasodilator, high arginase activities have often been associated with heart ailments such as atherosclerosis, myocardial infarction and ischemia. *Arginase* has been shown to modulate myocardial contractility by nitric oxide synthase 1- dependent mechanism¹⁸⁴. Higher *arginase* activity in lower Mg²⁺ ions have been observed in diabetic children by Bjelakovic et al.,

suggesting the pathophysiology of this disease could be understood to be related with increased arginine catabolism due to *arginase* action¹⁸⁵. The *arginase*-NO synthase competition for arginine has led to various effects on airway hyperresponsiveness, bronchial obstruction and lung inflammation in asthma due to increase in *arginase* activity. Munder have reported *arginase* modulation to have a noteworthy role in the mammalian immune system due to L-arginine metabolism being regulated by it leading to immunosuppression by suppression of T-cell immune responses¹⁸⁶. Maarsingh et al., and Munder have reported *arginase* to be a key enzyme in pathophysiology of allergic asthma opening novel therapeutic roles for asthma control through action of *arginase* inhibitors^{187,188}.

Table No: 2. Commercial available microbial arginase:

Name	Category	Company name	Application
Arginase	Arg1	U.S.Life Sciences	Suitable for use in ELISA, Western Blot, Immunoprecipitation and Conjugation. Other applications not tested

CONCLUSION

Microorganisms were considered to be potential sources of number of therapeutic enzymes like anticancer agents. Arginase can be produced commercially using these microorganisms in very less cost. Number of studies should be carried out and further steps has to be taken to promote therapeutic arginase in the market.

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