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Characterization of Plant-Based Biotin from SesZenBio[™] and In silico Studies to Prove Its Potential in Hair Care

Mihir Gadani^a, Ratna Upadhyay^a, Supriya Raut^a and Kavita Pandey^{b*}

^a Zenherb Labs, Jogeshwari, Mumbai, India. ^b Department of FYIC-Bioanalytical Sciences, G. N. Khalsa College (Matunga), India.

Authors' contributions

This work was carried out in collaboration among all authors. All authors read and approved the final manuscript.

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ABSTRACT

A standardized extract from *Sesbania agati* leaves (SESAL) was developed by Zenherb labs Pvt. Ltd., Mumbai, and is one of its kind plant-based biotin supplements. The current work aimed at characterization of SESAL using chromatographic techniques (like HPLC & HPTLC) and DNA authentication. Additionally, an in-silico approach (docking) was also adopted to predict the molecular interactions and probable mechanisms involved in potential role of SESAL in hair care products range. Drug likeliness was assessed on the basis of ADMET properties and Lipinski's Rule of 5. DNA fingerprinting followed by HPTLC confirmed the identification and authentication. The biotin content in SesZenBio was estimated to be 0.5% by validated HPLC method. Mechanistically, biotin along with 2 other cofactors in *SesZenBioTM* is predicted to stimulate the hair follicle growth and differentiation, thus improving hair health. *SesZenBioTM* might hold 2x potential as against synthetic biotin in hair care applications.

Keywords: Cosmeceutical formulation; hair follicle; DNA barcoding; HPTLC profiling.

1. INTRODUCTION

India has a well-established, traditional medicinal systems which have been accepted globally.

Natural drugs are known to be safer with fewer side effects and this has fuelled a huge demand for herbal medicines, across the world [1]. Additionally, we are one of the major contributors

*Corresponding author: E-mail: kavita.pandey@gnkhalsa.edu.in;

in world market of alternative medicines, owing to the rich biodiversity of medicinal plants in India [2]. Last decade has witnessed a surge in plant extracts in formulations for not only pharmaceuticals but also nutraceutical and cosmeceutical applications.

Sesbania, the plant under study, is one such genus (with over 22 species), explored for medicinal, food and cosmetics potentials [3]. Sesbania grandiflora and S. sesban are the most thoroughly studied amongst all species. S. grandiflora is commonly known as agati and belongs to family Caesalpiniaceae Qualitatively, the extracts contain alkaloids, carbohydrates, polyphenols, saponins, tannins, phytosterols and flavonoids. Leaves and flowers of agati show abundance of vitamins (like A, C riboflavin, biotin etc.) and minerals (like zinc, selenium etc.), while barks and flowers are known to be rich in polyphenols [4,5]. There are scare reports of applications of agati leaves extract in cosmetics. Agati contains biotin, which is known for roles in skin health [6], and has gained attention for cosmeceutical potential.

Biotin plays roles in protein synthesis, and, helps the body produce elastin and keratin to maintain healthy hair, skin, and nails. By helping the body use important enzymes, biotin also supports a healthy metabolism and nervous system. Biotin cannot be stored in the body, hence, regular supplementation is advised. Foods like nuts, legumes, whole grains, egg yolk etc are natural sources of biotin. and is also produced by the gut microbiota. Due its relatively low cost and easy availability in cosmetic products, biotin supplementation is trending in the looks conscious consumers.

Increased use of herbal medicines has created huge demand for the natural products. To assure the quality of the final product, from handling raw materials with good manufacturing practices to continuous quality control tests using modern and conventional analytical methods, become imperative. These analytical techniques aid in identification, quality evaluation and assessment of relative potencies of the crude drugs. Some of the commonly employed techniques include HPLC, TLC, HPTLC, GC, FTIR, MS etc. [7]. Phytochemical profiling, standardization of herbal finger print development formulations. quantitation is a part of development of plantbased formulations [8].

Substitution of the plant raw material or adulteration are the most common challenges

posed in the quality assurance of herbal formulations [9]. Thus, newer and reliable methods are needed for authentication of the raw material. Advancements in genomic technologies aid in molecular identification of the plant raw material, using markers. DNA barcoding is one of the most powerful methods to confirm the authentication of plant materials [1, 10].

The efficacy of the drug depends primarily on the interactions occurring at the molecular level. With the advent of computational biological tools, the drug discovery process has got a boost. Molecular docking aims at studying the interactions of ligands that bind to a specific receptor and the identification of its preferred, energetically most favourable, binding pose.

Zenherb labs, Mumbai has developed a proprietary formulation containing Sesbania agati leaves extract named SesZenBio[™]. In the current study, characterization of the formulation was carried out using analytical (HPLC & HPTLC) and genetic methods (DNA barcoding). Also, mechanistic evaluation of the benefits of biotin consumption on hair health has been carried out using computational studies (molecular docking), to gain а deeper understanding of the molecular mechanisms in play.

2. MATERIALS AND METHODS

2.1 Sample

The sample under study is a plant-based biotin, standardized extract from *S. agati* leaves (SESAL). The product is a proprietary product of the Zen herb lab and finds applications in hair, skin and nails health. The characterization and identification of the extract components were carried out with SESAL.

2.2 Biotin Quantitation

2.2.1 HPLC

Biotin content in *SesZenBioTM* was quantified using HPLC (Agilent, model-1200). Column was a reverse phase C 18 column (Thermo Accucore, 2.6µm,4.6mm x 150 mm). Mobile phase included solutions A (Water containing 0.025 % TFA:Acetonitrile(90:10v/v)) and B (Acetonitrile) flowing at the rate of 0.7ml /min. Detector-UV detector- 204 nm/210nm.

Sample and standards were prepared as per AOAC official method 2016.2 (Official Methods of

Analysis (2016) 20th Ed., AOAC INTERNATIONAL, Rockville, MD, Method 2016.02). Standard used was Biotin - Purity ≥99% (Cat. No. B4501; Sigma Chemical Co., St. Louis, MO , USA, or equivalent). Calibration curve was plotted using standard and test samples for quantification of Biotin.

2.2.2 HPTLC

A sensitive HPTLC method was developed for biotin estimation. CAMAG HPTLC used was the with the Scanner 3 for Densitometric Evaluation of Thin-Layer Chromatograms. Two grams of plant powder was added to 20ml of methanol and sonicated at 45°C for 30min. The content was centrifuged at 1500rpm and filtered. Extract so obtained was used as the sample for further analysis. Sample loaded per track was 4µl and developed in mobile phase containing Toluene: Ethyl acetate: Formic acid (6:3:1 v/v/v). Densitometric scan was carried out at 254nm to quantify biotin.

2.3 DNA Authentication

Small pieces of *S. agati* leaves were used for DNA authentication. Fish DNA barcoding method, suggested by FDA was adopted for DNA extraction [11]. DNA was then amplified using standardized PCR and separated by agarose gel electrophoresis. The desired band was eluted and processed for DNA sequencing. DNA Bold and NCBI BLAST programs were used for identification of the plant material.

2.4 Molecular Docking

significant Biotin $(C_{10}H_{16}N_2O_3S,$ а phytoconstituent from S. agati was docked against 3 proteins involved in hair health, based on thorough literature search (Table 1). PubChem (https://pubchem.ncbi.nlm.nih.gov/) was employed to procure the 3-dimensional structures of the biotin in .SDF format and later converted to pdbgt format. Three target proteins, involved in hair health were chosen for the molecular docking, namely- 2HFP, 2P54, and 3LCK (Table 2). Three dimensional structures were downloaded from RCSB (www.rcsb.org). Heteroatoms water molecules and were removed, while Kollman's charges and hydrogen bonds were added to ready the proteins for docking. The protein structures were saved in the pdbgt format. AutoDock Vina 1.5.6. was used to carry out the molecular docking studies in order to understand the interactions of biotin with target proteins that are involved in hair health. To predict the bioavailability of ligand, ADMET profiling was carried out, based on compliance to Lipinski's rule of 5. Bioactivity scores were predicted using the Molinspiration tool (www.molinspiration.com). Biotin was evaluated for its drug-likeliness.

3. RESULTS AND DISCUSSION

Natural, plant-based source of Biotin *SesZenBioTM* standardized to 0.5% Biotin is a uniquely developed, proven all natural, free from toxic residual solvents ingredient derived from authenticated *Sesbania agati* leaves.

3.1 BIOTIN Estimations

HPLC analysis carried out to quantify the biomarkers present in the $SesZenBio^{TM}$ extract, wherein 2 more cofactors were found (Fig. 1). These 2 cofactors from $SesZenBio^{TM}$ extract are believed to probably contribute towards its 2X effectivity in hair and skin health.

Thin-layer chromatography and HPTLC is one of the important separation chromatographic techniques used for detecting adulteration for assessing the quality of the plants through fingerprint profile of the drug [12]. If the drug is adulterated there might be the appearance of the other compounds, in turn may increase the no of spots. On the other hand, the exhausted or deteriorated drugs may lose the component, and the number of spots appeared might be less . HPTLC confirmed the quantitation of biotin as 0.5% (Fig. 2).

3.2 DNA Authentication

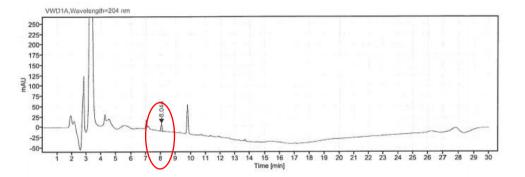
Conventionally, taxonomic identifications were carried out based morphological on characteristics, microscopic evaluation, analytical or phytochemistry based methods etc. [13] DNA barcoding has emerged as a powerful technique for molecular identification of an organism. Barcoding involves 3 steps: DNA extraction, PCR amplification and DNA sequencing followed by interpretation. The most common DNA barcodes used in plant identification are fragments from the genes like RuBisCo large subunit (rbcL) and maturase K (matK) [14,15]. In the current study, barcoding was carried out using matK and Internal transcribed spacer 2 (ITS2) markers [16] (Fig. 3). ITS 2 gene of nuclear ribosomal DNA serves as a unique barcode for identification of medicinal plants from a broad range of plant taxa [17,18,19].

3.3 Molecular Docking

Biotin (Pubchem ID-171548) was docked against 3 target proteins involved in roles in hair health namely-2HFP, 2P54 and 3LCK. The binding energies for the 3 were -6.5, -6.5 and -7.3 kcal/ mol. It complied with all the parameters of Lipinski's rule of 5, indicative of their good bioavailability (Table 1).

Peroxisome proliferator-activated receptors (PPARs) are nuclear hormone receptors abundantly expressed in tissues that display high fatty acid metabolism, including human skin and its appendages [20]. Three different isoforms of PPARs exist—PPAR- α , PPAR- β/δ and PPAR- γ , which demonstrate diverse functions. PPAR- γ -mediated signalling plays a significant role in the hair follicle (HF) development. PPAR activates the keratinocyte differentiation and accelerates

skin barrier recovery. PPAR-y signalling also promotes the mitochondrial energy metabolism in humans [21]. Ramot and co-workers have coworkers, have summarized the biological functions of PPAR-y in skin and hair health, in a review article. Biotin enhances the protein expression levels of PPAR, thus it finds cosmeceutical applications. Continuous stimulation of PPAR-y by endogenous ligands like biotin, may be required for the maintenance of HF energy metabolism, for intrafollicular PPAR-y expression and activity and is crucial for the HF physiology [22]. Biotin deficiency enhances the kinase activity of T CD4 T lymphocytes, which further produces proinflammatory molecules. Bioactivity scores (Table 2), displays the effectivity of biotin based on the typical receptors the ligand will encounter in the cell. The score allows efficient identification of active and inactive molecules.



NAME	Compound Amount Unit	RT [min]	Area	Compound Amount	Peak Tail Factor	PEAK THEORETICAL PLATES USP
Biotin	mcg/kg	8.044	93.19	818.545	0.81	62531.27669
MESSAL SA	1	Sum	93.19			



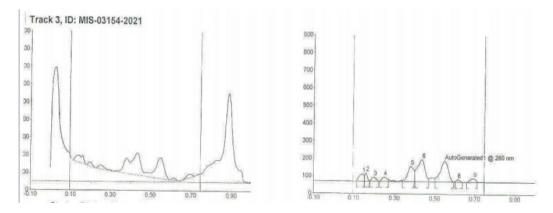


Fig. 2. Quantitation of biotin using HPTLC

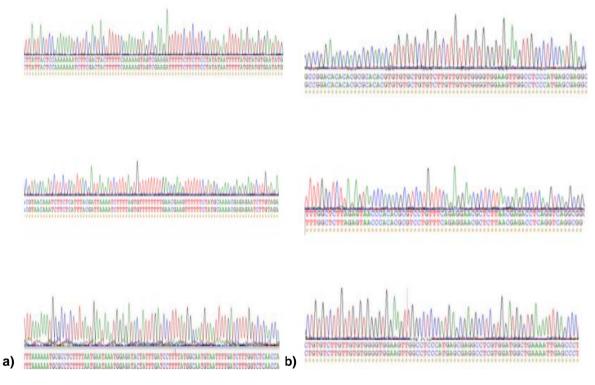


Fig. 3. DNA Barcode Markers from S. agati : a) matK and b) ITS2

Table 1. Evaluation of drug-likeliness of biotin based on the Lipinski's rule of 5

Ligand	Parameters of lipinski's rule's of five							
	miLogP	TPSA	natom	MW	nON	nOHNH	nrotb	nvio
Biotin	0.76	78.42	16	244.32	5	3	5	0

Table 2. Bioactivity scores of biotin

Ligand	Bioactivity scores							
-	Gpcr	lon Channel	Kinase Inhibitor	Nuclear Receptor	Protease Inhibitor	Enzyme inhibitor		
Biotin	-0.10	-0.57	-0.62	- 0.63	0.27	0.25		

4. CONCLUSION

SesZenBioTM has been found to be mechanistically double in hair and skin care potential. This might be attributed to the higher bioactives in the proprietary extract. The extract is derived from green process, hence free from toxic residual solvents. *SesZenBioTM* is proven to have no carbon footprints hence, 100% naturally derived.

Polyphenols are active on 4 receptors,viz.-3LCK,4GYY,5YLC,1MFR(in order of binding scores). Biotin is active on 2R53 receptor-(both synthetic & natural) and complies with the Lipinski's rule of 5, thus is bioavailable and a potential candidate molecule. It is noteworthy that the natural plant-based biotin contains NLT 2% polyphenols which could act on the 4 additional receptors leading to hair follicle stimulation. Hence SeszenBioTM is way ahead of synthetic biotin in therapeutic efficacy which is because of its efficiency to interact with 5 receptors as against synthetic biotin, that binds to only 1 receptor. The analytical techniques confirmed the concentration of biotin in the final product. DNA barcoding identified the plant as S. grandiflora, using matK and ITS2 markers.

DISCLAIMER

The products used for this research are commonly and predominantly used products in our area of research and country.

There is absolutely no conflict of interest between the authors and producers of the products.

COMPETING INTERESTS

Authors have declared that no competing interests exist.

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