



## Isolation of stigmasterol from hexane extract of leaves of *Pisonia grandis* R.Br. in vitro anti-diabetic and its molecular docking studies

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### ABSTRACT

*Pisonia grandis* R.Br belonging to the family *Nyctaginaceae* is a widely distributed evergreen tree in India known for its medicinal uses. The study was aimed to investigate the anti-diabetic property in the leaves of *Pisonia grandis* R.Br. The isolation and purification were performed by the conventional column chromatography and the resultant yield was found to be a white crystalline powder, which was further subjected for characterisation through IR, <sup>1</sup>H NMR, <sup>13</sup>C NMR and mass spectroscopy. From the characterisation data, the isolated compound was identified as stigmasterol; it was first time isolated from the hexane extract of the leaves. The  $\alpha$ -amylase inhibitory activity of stigmasterol from the hexane extract of the leaves of *Pisonia grandis* R.Br showed high potent activity with an IC<sub>50</sub> value of 46  $\mu$ g/mL. The anti-diabetic activity of the compound against  $\alpha$ -amylase and four other diabetic enzymes-  $\alpha$ -glucosidase, acid phosphatase, endo- $\beta$ -N-acetylglucosaminidase and  $\beta$ -glucuronidase were further investigated by molecular docking studies and proved that stigmasterol can be a potential anti-diabetic agent.



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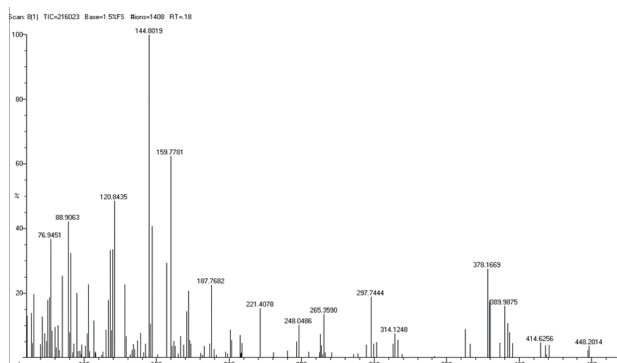
### INTRODUCTION

Drugs from natural sources have proved to be effective in the treatment of human diseases. Throughout history, plant material has served as a reservoir of potential sources for new drugs. A great example

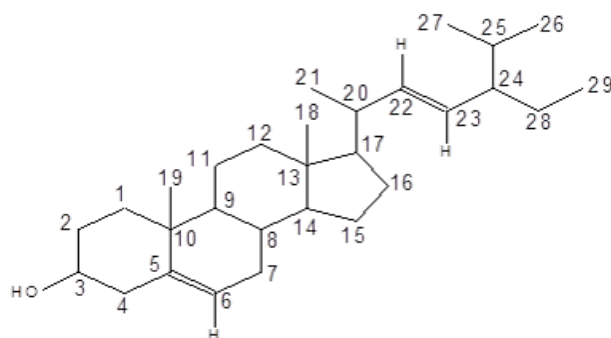
is that of vincristine and vinblastine, which are used in the treatment of cancerous cells. Diabetes mellitus is a disease that has been affecting humanity globally. The number of casualties with diabetes is estimated to rise from 171 million in 2000 to around 366 million in 2030 (Wild *et al.*, 2004). According to the International Diabetic Federation, there was nearly 73 million diabetic's diagnosis in India in 2017. WHO has predicted that by 2030, diabetes will be the seventh among causes of death.

*Pisonia grandis* is an evergreen tree native to tropical islands in the Indian and Pacific Oceans, and adjoining coastal areas (CSIR. 1969). It belongs to the family of *Nyctaginaceae*. *Pisonia grandis* has been extensively used in Indian traditional medicine for the treatment of inflammatory diseases (Anbalagan *et al.*, 2002; Elumalai *et al.*, 2012a), ulcer (Mabberley, 1997), arthritis (Elumalai and Prakash, 2012),

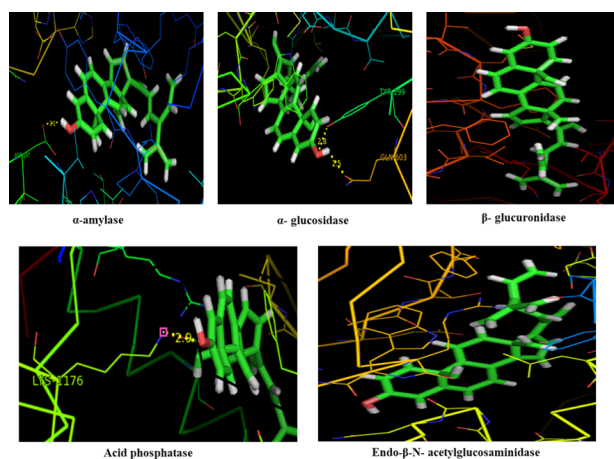




**Figure 4: Mass spectral data of the isolated compound**



**Figure 5: Structure of the isolated compound, stigmasterol**



**Figure 6: PyMol visualization of the enzymes**

a colorimetric method using acarbose as the reference compound. The starch solution (0.5% w/v) was obtained by stirring and boiling 0.25 g of potato starch in 50 mL distilled water for 15 minutes. The enzyme solution (0.5 units/mL) was prepared by mixing 0.001 g of alpha-amylase in 100 mL of 20 mM phosphate buffer (pH-6.9) containing 6.7 mM sodium chloride. Phenol (stock solution 1 mg/mL) was dissolved and made up with DMSO and from the solution, different concentrations (two-fold dilutions) 50, 100, 200, 400 and 800  $\mu\text{g/mL}$  were prepared. The indicator prepared was comprised of

96 mM 3, 5-dinitro salicylic acid (20 mL), 5.31 mM potassium tartrate in 2M sodium hydroxide (8 mL) and distilled water (12 mL). 1 mL of phenol and 1 mL enzyme solution were mixed and incubated at 25 °C for 30 minutes. To this, 1 mL of the coloring reagent was added and closed, which was further placed into a water bath maintained at 85 °C. 15 minutes later, the reaction mixture was removed from the water bath, cooled and diluted with 9 mL distilled water. The generation of maltose was quantified by the reduction of 3, 5-dinitro salicylic acid to 3-amino-5-nitro salicylic acid and detected at an absorbance of 540 nm. The readings were compared with the control (acarbose), containing buffer instead of the sample. The percentage inhibition of alpha-amylase was assessed by the given formula below,

Percentage Inhibition

$$= \left[ \frac{\text{Absorbance of control} - \text{Absorbance of test}}{\text{Absorbance of control}} \right] \times 100$$

### Molecular Docking Studies

Molecular docking study (Eissa et al., 2009) was carried out to study the binding mechanism of the isolated compound in the active site of  $\alpha$ -amylase.

Additionally, four other enzymes associated with diabetes mellitus, namely  $\alpha$ -glucosidase, acid phosphatase, endo- $\beta$ -N-acetylglucosaminidase and  $\beta$ -glucuronidase were selected to assess the binding efficacy and inhibitory effects of the isolated compound against the active enzymes in the *in silico* method. The protein sequences were taken from NCBI (National Center for Biotechnology Information) and were converted into FASTA format. The sequences were then allowed into the BLAST (Basic Local Alignment Search Tool) database. The protein structure files were taken from Protein Data Bank. Water molecules and other heterogeneous atoms were removed and hydrogen atoms were added to the structure. PyMOL software was used to view the structure and calculate the length of the hydrogen bond. Docking was performed using Argus Lab.exe. The ligand molecules (standard) were obtained from PubChem and generated by using Chem Sketch ACD Lab.

## RESULTS AND DISCUSSION

### Characterisation of the isolated compound

The column chromatography of the hexane extract of *Pisonia grandis* leaves over silica gel led to the isolation of a white crystalline compound (48 mg). The compound was recovered and subjected to thin-layer chromatography for identification.  $R_f$  value of

**Table 1: NMR spectral data of isolated compound**

Atom	Carbon, $\delta$ , ppm	Hydrogen, $\delta$ , ppm
1	31.88	1.02 – 1.61 (m, 2H)
2	37.16	1.02 – 1.61 (m, 2H)
3	71.08	3.59 (m, 1H)
4	43.80	1.73 – 2.05 (m, 2H)
5	139.58	—
6	117.47	5.03 (m, 1H)
7	31.49	1.73 – 2.05 (m, 2H)
8	38.00	1.02 – 1.61 (m, 1H)
9	49.47	1.02 – 1.61 (m, 1H)
10	39.48	—
11	25.40	1.02 – 1.61 (m, 2H)
12	40.28	1.02 – 1.61 (m, 2H)
13	55.06	—
14	55.92	1.02 – 1.61 (m, 1H)
15	29.65	1.02 – 1.61 (m, 2H)
16	29.65	1.02 – 1.61 (m, 2H)
17	55.14	1.02 – 1.61 (m, 1H)
18	19.83	0.53 – 1.03 (s, 3H)
19	26.22	0.53 – 1.03 (s, 3H)
20	40.83	1.73 – 2.05 (m, 2H)
21	19.01	0.53 – 1.03 (s, 3H)
22	129.46	5.15 (m, 1H)
23	138.17	5.18 (m, 1H)
24	51.26	1.73 – 2.05 (m, 1H)
25	34.23	1.73 – 2.05 (m, 1H)
26	21.56	0.53 – 1.03 (s, 3H)
27	21.38	0.53 – 1.03 (s, 3H)
28	28.51	1.02 – 1.61 (m, 2H)
29	13.05	0.53 – 1.03 (s, 3H)

**Table 2:  $\alpha$ -amylase inhibitory activity**

S/No	Concentration ( $\mu\text{g/mL}$ )	% Relative Enzyme Inhibitory activity (R.E.I.)	% Inhibitory Activity (100-R.E.I.)	IC <sub>50</sub> ( $\mu\text{g/mL}$ )
1	50	68.98	31.02	46
2	100	18.76	81.24	
3	150	18.12	81.88	
4	200	12.60	87.40	

**Table 3: Molecular docking scores**

S/No	Enzyme	Docking Score	Amino Acid	Hydrogen Bond Length
1	Endo- $\beta$ -N-acetylglucosaminidase	-12.6655 kcal/mol	-	-
2	$\alpha$ -glucosidase	-13.1196 kcal/mol	Glutamine-603 Tyrosine-299	2.5 2.8
3	$\alpha$ -amylase	-13.6192 kcal/mol	Asparagine-197	2.1
4	Acid phosphatase	-14.1174 kcal/mol	Lysine-1176	2.9
5	$\beta$ -glucuronidase	-16.1866 kcal/mol	-	-

0.26 was found using the mobile phase system, i.e., hexane: ethyl acetate (4:1).

The  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectrum of the isolated product (Table 1) indicates that a multiplet at 3.59 ppm (Figure 2) is due to the CH bond connected to oxygen (C-O) in proton NMR and at 71.08 ppm (Figure 3) in carbon NMR. This is confirmed by the IR spectrum stretching appearing at  $1375\text{ cm}^{-1}$  (Figure 1). The signal which appeared in the range of 0.53-2.05 ppm (Figure 2) was assigned to  $\text{CH}_3$ ,  $\text{CH}_2$  and CH groups and in carbon NMR signals, which appeared in the range of 13.05 ppm to 55.92 ppm (Figure 3). This is further confirmed by IR stretching at 2943,  $2868\text{ cm}^{-1}$  (Figure 1). A peak at 5.03 to 5.18 ppm (Figure 2) in proton and 117.47 and 138.17 ppm (Figure 3) in carbon NMR, corresponds to  $-\text{C}=\text{CH}$ -group. The peak at  $3317\text{ cm}^{-1}$  (Figure 1) in the IR spectrum corresponds to the hydroxyl (OH) group. The actual molecular weight of the compound is 412 and in the mass spectrum, it shows a peak at 448 ( $\text{M}+2\text{NH}_4$ ) $^+$ , as seen in Figure 4. All the above spectral data confirm the structure of the compound to be stigmaterol (Poongothai and Sripathi, 2018) (Figure 5). Salkowski and Liebermann-Burchard tests were carried out (Njoku and Obi, 2009; Kandati et al., 2012) and stigmaterol tested positive for both confirming the presence of steroidal moiety (Rambeloso et al., 2014; Pierre and Moses, 2015; Kirtikar and Basu, 2012).

#### **In Vitro $\alpha$ -amylase Inhibitory Assay**

The  $\text{IC}_{50}$  value of stigmaterol was found to be  $46\text{ }\mu\text{g/mL}$  (Table 2). The plant claimed to have anti-diabetic activity as the leaves contain stigmaterol, which mimics the action of acarbose.

#### **Molecular Docking Studies**

The PyMol visualization of the interaction between stigmaterol and the enzymes is given in Figure 6. At the end of the docking studies, it was found that  $\alpha$ -amylase and  $\alpha$ -glucosidase gave good docking scores of  $-13.6192\text{ kcal/mol}$  and  $-13.1196\text{ kcal/mol}$ , respectively (Table 3).  $\alpha$ -amylase bonded with isoleucine and lysine. A bond length

of 2.1 with asparagine signifies a close interaction between  $\alpha$ -amylase and stigmaterol (ligand).  $\alpha$ -glucosidase showed hydrogen bond lengths of 2.5 and 2.8, respectively, with glutamine and tyrosine, which also signifies a close interaction with the ligand. The interaction could mean an inhibitory effect. Acid phosphatase gave a bond length of 2.9 with lysine. Endo- $\beta$ -N-acetylglucosaminidase and  $\beta$ -glucuronidase did not show any bonding with stigmaterol. These enzymes are associated with diabetes mellitus.

#### **CONCLUSION**

The study aimed to isolate, characterise, screen and perform molecular docking studies of the isolated phytosterol present in the hexane extract of the leaves of *Pisonia grandis* R.Br. The isolated compound was physically characterised and was found to have a melting point in the range of  $164\text{--}166\text{ }^\circ\text{C}$ . The characterised compound was structurally confirmed to be stigmaterol by spectroscopic methods like IR,  $^1\text{H}$  NMR,  $^{13}\text{C}$  NMR and mass spectroscopy. The compound, when subjected to biological screening test ( $\alpha$ -amylase inhibitory activity), gave an  $\text{IC}_{50}$  value of  $46\text{ }\mu\text{g/mL}$ . This signifies that stigmaterol has an inhibitory activity on  $\alpha$ -amylase enzyme and can be of use in the treatment of diabetes mellitus. Finally, molecular docking studies were carried out with stigmaterol and enzymes associated with diabetes mellitus which further established that stigmaterol can be a promising anti-diabetic agent. The furtherance of this research work will involve the semi-synthetic modification of this ligand using the computer-aided drug designing tools and subsequently evaluating its in-vitro activity for anti-diabetic activity.

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