Identification and Characterization of Berberine in *Tinospora cordifolia* by Liquid Chromatography Quadrupole Time of Flight Mass Spectrometry (LC MS/MS Q-tof) and Evaluation of its anti Inflammatory Potential

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**ABSTRACT**

**Introduction:** *Tinospora cordifolia* (Willd.) Miers ex Hook F and Thomas commonly called as gudduchi or amrita is a widely used plant in traditional medicinal system of Ayurveda. A UPLC MS/MS Q-tof method for the identification and characterization of berberine in *Tinospora cordifolia* (Willd.) Miers. ex Hook.F and Thomas, and to evaluate the anti inflammatory potential of bioactive fraction. **Materials and Methods:** The presence of berberine in *Tinospora cordifolia* was determined by HPLC and was subsequently isolated by HPTLC. The anti inflammatory property of the fraction containing berberine was demonstrated to have an inhibitory activity on 5 lipoxygenase, an enzyme involved in inflammatory pathway and its IC$_{50}$ value was obtained. The binding interactions between berberine and 5-LOX were demonstrated by docking studies. **Result:** The presence of berberine in *Tinospora cordifolia* methanolic extract was identified by HPLC and HPTLC analysis and confirmed by UPLC MS/MS Q-tof. The fraction containing berberine inhibited 5-LOX with an IC$_{50}$ of 0.041± 0.0003µg/mL as compared to that of NDGA (positive control) which showed an IC$_{50}$ of 2.75 ± 0.05 µg/mL. Molecular docking of berberine with 5-LOX showed a binding energy of -8.942 ± 0.039665 kcal/mol and Ki of 273.16 ± 3.026 nM as compared to the NDGA which has a binding energy of -7.186 ± 0.170503 kcal/mol and Ki 5.604± 1.618 µM. **Conclusion:** *Tinospora cordifolia* can be used as a source of berberine and possible anti inflammatory activity of *Tinospora cordifolia* may be attributed to the presence of berberine.

**Key words:** *Tinospora cordifolia*, Mass Spectrometry, 5 Lipoxygenase, Berberine, Molecular Docking.

**INTRODUCTION**

*Tinospora cordifolia* (Willd.) Miers ex Hook F and Thomas commonly called as gudduchi or amrita is a widely used plant in traditional medicinal system of Ayurveda. The plant has been used for centuries for the treatment of various ailments. It has been reported to have anti-allergic, anti-arithmetic, anti-diabetic, anti-inflammatory, anti-oxidant, anti-spasmodic, anti-periodic, radio protective properties. Various compounds have been isolated and identified from the plant.

Berberine is an isoquinoline alkaloid found in various plants and has been used in traditional Indian and Chinese medicines having wide range of pharmacological actions against gastroenteritis, diabetes, hyperlipidemia, obesity, fatty liver, coronary artery diseases, hypertension, diabetes and metabolic syndrome, polycystic ovary, Alzheimer's disease and in cancer. HPLC determination of berberine in plants extracts have been described earlier. LC MS/MS multiple mass spectrometry is an highly effective tool in structural characterization of low molecular weight compounds. 5-Lipoxygenase is a key enzyme involved in pathophysiology of inflammation. Biomolecules which can inhibit lipoxygenase have potential therapeautic role in controlling various inflammatory diseases including cancer.

The study has led to identification of berberine in *Tinospora cordifolia* by LC MS/MS Q-tof and the bioactivity of the fraction containing berberine was evaluated for anti inflammatory potential by inhibition of 5-LOX.

**MATERIALS AND METHODS**

HPLC analysis was done using Shimadzu LC20AP HPLC system using Enable C18 G250 X 4.6mm column having particle size 5µm and pore size 120Å and SPD20A UV-VIS detector (Shimadzu, Japan), HPTLC was from Camag, Switzerland. HPTLC system consisting of Linomet TLC sample applicator to spot the sample and Wincats software were used for spectral scanning of TLC plates. Aluminum coated with silica gel 256GF TLC plates (mesh size 60Å, layer thickness of 200 µm and particle size of 8-12 µm) was purchased from Sigma –Aldrich, USA. UPLC MS/MS Q-tof analysis was carried out in Waters Xevo G2-Q-Q mass spectrometer with ACQUITY UPLC BEH C18 column hav-
Berberine standard was from Sigma–Aldrich, MO, USA. LC-MS grade solvents were purchased from Fischer Scientific (New Jersey, USA). All the solvents used were of HPLC grade and was purchased from Sigma-Aldrich, MO, USA. 5-Lipoxygenase and NDGA was procured from Cayman chemicals, MI, USA. Xylenol orange, ferrous sulphate and sulphuric acid was from Himedia chemicals, India. The structure of 5 LOX was downloaded from protein data base (http://www.rcsb.org/pdb/explore.do?structureId=3v99) with PDB ID: 3V99.

**Plant material**

The plant materials were collected from parts of Kottayam region of Kerala, India and were identified by taxonomist at Department of Botany, St.Thomas College, and Pala, India. A Voucher specimen was deposited and voucher number (JA17151) was obtained. The plant stem was cleaned, dried and powdered and stored at -20°C.

**Extraction and isolation**

The shade dried stem was powdered, weighed and 20g was used for further extraction. First, the powder was defatted with hexane, followed by soxhlet extraction for 24 hrs with methanol. After extraction, methanol was removed by rotary evaporation and the remaining extract having a percentage yield of 6.94 % was lyophilized and stored at -20°C. The lyophilized samples were re-dissolved in methanol. The sample was filtered through 0.22µm syringe filter.

For reverse phase HPLC analysis, 20µl of sample was injected into Shimadzu HPLC system. The solvent system was methanol (A) and 0.1% formic acid in water (B) with a gradient time program from 0-60 min with B ratio ranging from 5-70%. The flow rate was kept constant at 1.5ml/min with a maximum pressure of 300kg/cm². Berberine standard dissolved in methanol were run in similar conditions. The peaks obtained for T. cordifolia extract and that for berberine standard was compared.

The extract was spotted onto silica gel 256GF TLC plates along side of berberine. The plate was developed in presaturated camag twin trough chamber with solvent system toluene, acetone and water in the ratio 5:1:5:1 up to 80mm. The plate was removed, dried, viewed and scanned at 256nm and 366nm using Camag TLC scanner with help of WinCats™ software.

The region of the TLC plates in sample spots having similar Rf to that of standard Berberine was scrapped off and mixed thoroughly with ethanol and centrifuged. The isolated fraction was lyophilized (4.2mg) and was further used for UPLC MS/MS Q-tof and in inhibition studies.

**Characterization of Berberine by UPLC ESI MS/MS Q-tof**

The isolated fraction obtained from TLC was subjected to UPLC MS/MS Q-tof as described earlier. A 10µl of fraction and berberine standard was injected. A mixture of 0.1% formic acid in water (A) and acetonitrile (B) was used as solvents with gradient time program from 0-60 min :10-95% B, 6.5-9min: 95-10% B with a flow rate of 0.3ml/min at 30°C. Mode of ionization was positive electrospray ionization with capillary voltage set at 3.5kV and cone voltage of 30V. Nitrogen was used as both the sheath gas and drying gas. The flow rate for sheath gas was 900 L/hr and for drying gas it was 50 L/hr and was set at temperature of 350°C and 135°C respectively. The back pressure was set at 1500psi. The Q-tof data was collected for a range of m/z 50 and m/z 1500. The data was collected using MS² centroid technology which uses two collision energies (6eV and 20-30eV) to produce precursor and fragment ions. Mass calibration was done using leucine-enkephalin and the residual error was set at 2ppm. The accurate mass and composition of parent and daughter ions was calculated using Mass Lynx V 4.1 software.

**Inhibition of 5-Lipoxygenase**

The fraction isolated by HPTLC, was lyophilized, quantified and re-dissolved in methanol. Efficacy of different concentrations of the isolated fraction and standard drug NDGA on inhibition of 5-LOX were determined using FOX assay with modifications. Briefly, 20µl of 5-LOX and 20µl of different concentrations of the fraction were mixed together and incubated at RT for 20 min. To this 140µM of linoleic acid was added, mixed and incubated in dark for 20min. FOX reagent (130µl) was added, mixed and incubated again in dark for 30 min. Absorbance was taken at 560nm. The percentage inhibition was calculated as (Absorbance of control-Absorbance of test)/ (Absorbance of control) × 100. The IC₅₀ values were calculated using GraphPad Prism 5.0 software.

**Docking studies**

The structure of human 5- lipoxygenase was downloaded from RCSB protein data bank (PDB ID: 3V99) (http://www.rcsb.org). The structure of berberine was drawn using Chemsketch (www.acdlabs.com/resources/freeware/chemsketch). The structure of NDGA used as standard was downloaded from Chemspider (http://www.chemspider.com/Chemical-Structure.4375.html). Protein modification was performed using Swiss-PDB viewer and by AutoDock 4.0. During protein preparation all heteroatoms was removed except iron which is required for enzyme activity, polar hydrogen atoms added and Kollman charges was assigned to the protein macromolecule. For the binding of ligand with the protein, the binding site was determined having a dimension of 60 A³ around the active site with amino acid coordinates 17.650, -28.623, -26.503 in B chain of the enzyme. During docking the protein was set rigid and docked by applying Lamarckian genetic algorithm with a maximum of 2500000 energy evaluations and RMSD tolerance of about 2.0 Å. The docked protein was visualized by using Discovery Studio 4.5 visualizer. The results were determined based on RMSD(root mean square deviation) and binding energy.

**Statistical analysis**

The results were expressed as mean ± standard deviation of three independent experiments. Statistical analysis was done using unpaired t-test using GraphPad prism 5.0™. A p value ≤ 0.05 were considered as statistically significant.

**RESULT AND DISCUSSION**

In UPLC total ion chromatogram a clean peak was obtained for berberine standard at 3.79min and a peak at 3.73min was obtained (Figure 2).

In HPLC chromatogram of berberine as standard showed a broad peak with retention time at 31.898 min at a wavelength of 350nm. *Tinospora cordifolia* methanolic extract chromatogram showed a peak of 31.477 corresponding to that of berberine standard under similar run conditions. Statistical analysis was done using pairwise t-test using GraphPad Prism 5.0. A p value ≤ 0.05 were considered as statistically significant.
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Figure 1: HPLC chromatogram of (A) Berberine standard and (B) T.cordifolia methanolic extract. HPLC conditions were same for both berberine and T.cordifolia extract.

Figure 2: HPTLC chromatogram of Berberine standard and T.cordifolia. An Rf of 0.16 was observed for T.cordifolia methanolic extract which corresponds to the Rf of standard berberine.

Figure 3: Total ion chromatogram obtained for (A) standard berberine and (B) T.cordifolia methanolic extract. A retention time of 3.79cm and 3.73cm was obtained for berberine and T.cordifolia methanolic extract respectively.

Figure 4: MS spectra for (A) standard berberine and (B) T.cordifolia methanolic extract. Berberine standard showed a molecular mass of 336.1240g/mol and T.cordifolia methanolic extract showed a compound having molecular mass of 336.1243g/mol which corresponds to the molecular mass of berberine.
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**Figure 5:** MS/MS fragmentation pattern for (A) Berberine standard and (B) *T.cordifolia* methanolic extract bioactive fraction.

**Figure 6:** A) and (B). Docking of Berberine with 5 LOX active site showing different types of interactions. The interactions are shown as dotted lines.

**Table 1:** IC$_{50}$ value and Binding energies of berberine were compared with that of NDGA. Better inhibition was observed by berberine than for NDGA towards 5 LOX. Binding of berberine was higher than NDGA as shown by its binding energy and Ki values. *In silico* docking of 5 LOX (PDB ID: 3V99) was in accordance to the *in vitro* data obtained. Values expressed as mean ± S.D. *p value < 0.05, # p value <0.001.

<table>
<thead>
<tr>
<th>Sl No</th>
<th>Compound</th>
<th>IC$_{50}$ (µg/mL)</th>
<th>Binding Energy (kcal/mol)</th>
<th>Ki</th>
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<tr>
<td>1</td>
<td>Berberine</td>
<td>0.041 ± 0.0003*</td>
<td>-8.942 ± 0.039665*</td>
<td>0.273 ± 0.003 µM*</td>
</tr>
<tr>
<td>2</td>
<td>NDGA</td>
<td>2.75 ± 0.05</td>
<td>-7.186 ± 0.170503</td>
<td>5.604 ± 1.618 µM</td>
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obtain MS/MS fragmentation spectrum. MS/MS chromatogram showed similar fragmentation pattern for both standard berberine and compound from *Tinospora cordifolia* methanolic extract (Figure 5A, B). The accurate mass and the structure were elucidated by using Mass Lynx™ v4.1 software.

5LOX is a major enzyme involved in synthesis of pharmacologically important leukotriene and lipoxins which are mediators of inflammation.¹⁵ Leukotriens are involved in chemotaxis of granulocytes, its attachment to vascular cell walls, stimulate secretion of antibodies IgE, IgG and IgM, increased production of interleukin and plays role in pathogenesis of inflammatory arthritis, psoriasis, asthma,¹⁶ allergic rhinitis, inflammatory bowel and skin disease, osteoporosis, cancer and cardiovascular disease.¹¹ *Tinospora cordifolia* have been used in Indian traditional medical system of ayurveda for treatment of various ailments and has reported to inhibit 5-LOX.¹⁰ Inhibition of 12-lipoxygenase by fractions containing protoberberine has described earlier.¹⁰ Berberine have shown to be anti-inflammatory and yet the pathway for the control of inflammation has to be elucidated.¹¹ The fraction containing berberine inhibited 5-LOX with an IC$_{50}$ of 0.041± 0.0003µg/mL as compared to that of NDGA (positive control ) which showed an IC$_{50}$ of 2.75 ± 0.05 µg/mL.

Docking of berberine with 5 Lox active site was done to determine the different types of interactions involved and the strength of binding was expressed as binding energy. A lower binding energy represents high binding efficiency of ligand with the receptor.²² Berberine showed a binding energy of -8.942 ± 0.039665 kcal/mol and the estimated inhibition constant, Ki towards 5 LOX was found to be 273.16 ± 3.026 nM as compared to the NDGA which has a binding energy of -7.186 ± 0.170503 kcal/mol and Ki 5.604± 1.618 µM (Table 1). Docking revealed a strong binding of berberine than NDGA. Hydrogen bond was seen between berberine and amino acids Tyr 558, Asn554, Gln557 of A chain of 5 LOX. Hydrophobic interactions were seen between A chain His372, Ile406, Phe555 and Leu607 (Figure 6A, B). Inhibitory constant Ki for berberine was less than NDGA which is in accordance to the IC$_{50}$ obtained. Statistical analysis revealed berberine to be a better inhibitor than NDGA used as positive control.

**CONCLUSION**

The study has confirmed the presence of berberine in *Tinospora cordifolia* by means of high end mass spectrometric analysis. The fraction containing berberine has been shown to inhibit 5-1ipoxygenase which is involved in inflammatory pathway and interactions were demonstrated by means of docking. *Tinospora cordifolia* can be used as a source of berberine and...
possible anti-inflammatory activity of *Tinospora cordifolia* may be attributed to the presence of berberine.

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**CONFLICT OF INTEREST**

The authors declare no conflict of interest.

**ABBREVIATIONS USED**

HPLC: High performance liquid chromatography; HPTLC: High performance thin layer chromatography; UPLC ESI MS Qtof: Ultra perfor-

**REFERENCES**


7. Yong, Y., Kai, H., Bao-Shun, Z. & Xue-Gang, L. Study on interaction between plasmid DNA and berberine derivatives with aliphatic chain by fluorescence


14. Waslidge N B, Hayes DJ. A colorimetric method for the determination of lipo-

15. Kumar M, Dagar A, Gupta VK, Sharma A. In silico docking studies of bioac-


17. Aparoy P, Reddy RN, Guruprasad L, Reddy MR, Reddanna P. Homology mod-


**SUMMARY**

- Presence of berberine in *Tinospora cordifolia* was confirmed by means of high end mass spectrometry analysis (UPLC MS/MS Qtof). The fraction containing berberine was checked for anti-inflammatory activity which showed an inhibition on activity of 5-lipoxygenase (IC50. 0.041 ± 0.0003g/ml). The interaction between berberine and 5-lipoxygenase was demonstrated by molecular dock-

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Mohind C. Mohan: He is currently involved in identifying and characterizing anti inflammatory biomolecules from Ayurvedic medicines.

Anu P. Abhimannue: Has carried out works in determining possible anti inflammatory and anti carcinogenic lead molecules from medicinal plants.

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