

Larvicidal Activity and Molecular Docking Studies of Vitexicarpin from *Vitex negundo* Linn

Sandhya Krishnamoorthy¹, Sivakumar Muthusamy^{2*}, Vijayakumar Arumugam Ramamurthy³

¹Faculty of Pharmacy, Sri Ramachandra Institute of Higher Education and Research, Porur, Chennai, Tamil Nadu, INDIA.

²Department of Pharmacognosy, Faculty of Pharmacy, Sree Balaji Medical College and Hospital Campus, Bharath Institute of Higher Education and Research, Chromepet, Chennai, Tamil Nadu, INDIA.

³Department of Pharmacology, Faculty of Pharmacy, Sree Balaji Medical College and Hospital Campus, Bharath Institute of Higher Education and Research, Chromepet, Chennai, Tamilnadu, INDIA.

ABSTRACT

Objectives: To study the larvicidal effect of *Vitex negundo* Linn against mosquito larvae by larvicidal bioassay and *in silico* molecular docking studies.

Methods: The larvicidal bioassay of *Vitex negundo* Linn was assessed by using WHO standard protocol and an *in silico* molecular docking study was performed by using Molegro Virtual Docker against *Culex quinquefasciatus*. The early third instar larvae of *Aedes aegypti*, *Aedes albopictus* and *Culex quinquefasciatus* were exposed to the extract concentrations in three replicates and different temephos concentration were taken as positive control and solvent control was also used. **Results:** The study results shows that of *Vitex negundo* Linn possess the potential larvicidal activity against third instar larvae of *Culex quinquefasciatus* when compared to *Aedes aegypti* L and *Aedes albopictus*. Percentage mortality and half-maximal effective concentration (EC₅₀) value were calculated against the above mentioned three species like wise *Culex quinquefasciatus* (83.77 ppm), *Aedes aegypti* (341.7 ppm), and *Aedes albopictus* (487.9 ppm). The highest larval mortality was found against *Culex quinquefasciatus* of 100 % at 500 parts per million. *In silico* molecular docking studies using Molegro Virtual Docker against Odorant binding protein (PDB: 2L2C) of

Culex quinquefasciatus was performed as it showed better results in *in vitro* study. The results reveal that phytoconstituent like Vitexicarpin showed good docking scores against the Odorant Binding Protein 2L2C and the moldock score of Vitexicarpin was found to be -87.3733, whereas moldock score of standard Azadirachtin was found to be -110.77. **Conclusion:** The present study reveal that the *Vitex negundo* Linn possess the potential larvicidal activity against the *Culex quinquefasciatus* species.

Key words: *Vitex negundo* Linn, *Aedes aegypti* L, *Culex quinquefasciatus*, *Aedes albopictus*, Molegro Virtual Docker 6.0, Odorant binding Protein, Larvicidal.

Correspondence

Dr. M Sivakumar,

Professor, Department of Pharmacognosy, Faculty of Pharmacy, Sree Balaji Medical College and Hospital Campus, Bharath Institute of Higher Education and Research, Chromepet, Chennai-600044, Tamil Nadu, INDIA.

Email id: sivampharma@gmail.com

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INTRODUCTION

Mosquito is a small, midge-like fly that belong to the family of *Culicidae*. Female in most species are ectoparasites, whose tube-like mouthpart pierces skin of the host to consume the blood. The human blood is rich in proteins, amino acids and other nutrients, which makes it the perfect prenatal supplement for growing the mosquito eggs that is a reason female mosquito sucking the blood. The passing from host to host, some of the transmit are extremely harmful infections such as malaria, yellow fever, chikungunya, west Nile virus, dengue fever, filariasis, zikavirus, Japanese encephalitis, schistosomiasis and other arboviruses.¹ The mosquito undergoes four stages of life cycle mainly in egg, larva, pupa, and adult. Approximately, it takes a month to complete its life cycle.^{2,3} There are approximately 3,500 species of mosquitoes in the World and 41 different types of genera, however the majority of mosquitoes fall under three genera categories: *Aedes*, *Anopheles* and *Culex*. *Aedes* causes yellow fever, dengue, malaria and encephalitis. Larvae of *Aedes* are larger when compared to others within 7 to 10 days, larvae of *Aedes* enter the pupa stage. *Culex* causes filariasis, malaria, encephalitis and dengue. Larvae of *Culex* species are present upside down because their respiratory tract is present near the tail. The third instar larvae of three species of *Aedes aegypti*, *Aedes albopictus* and *Culex quinquefasciatus* are used in this experiment. The mosquito larvae for this research work were obtained and authenticated from Dr. Daniel Reegan, Entomologist, Entomology Research Institute, Loyola College, and Chennai. In commercially several synthetic mosquito control repellents available in

the market. All repellents have protected from mosquito bite for several hours and must be applied to all exposed skin areas. The available of common synthetic mosquito repellents is N, N-diethyl-3-methylbenzamide, commonly known as DEET, Picaridin, Permethrin, N, N -diethyl-2-phenyl-acetamide (DEPA), ethyl anthranilate (EA) and Ethyl butyl acetyl amino propionate (IR3535). The side effects like rash, skin eruptions, urticaria and respiratory problem have been encountered on the use of this type of synthetic repellents. The natural mosquito repellents like Citronella, Oil of lemon eucalyptus (OLE), Neem, Peppermint, methyl jasmonate and Lemon grass does not causes any type of skin irritation and allergic reactions as synthetic repellents and this natural type repellents are safe compared to already available synthetic one.

Vitex negundo Linn, generally known as the Chinese chaste tree, five-leaved chaste tree, or horse shoe Vitex or Monk's pepper.⁴ *Vitex negundo* Linn belongs to the family of Verbenaceae. *Vitex negundo* is a large aromatic shrub and with quadrangular, densely whitish, and tomentose branchlets future. It is an erect shrub or small tree growing height approximately 2 to 8 meter.⁵⁻⁹ The leaves are known to possess various flavonoids-chryso-splenol and vitexin, vitexicarpin, Luteolin, and Vitexin rhamnoside. Lignans, sesquiterpenes, flavone glycosides, iridoid glycosides, eucuboside aucubin, stilbenes have been isolated. Chasteberry is evidence that to contain a progesterone-like compounds, vitamin-C and carotene. It contains four phenolic compounds such as salviaplebeiaside, chryso-splenol-D, isovitexin and γ-tocopherol, along

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with β -sitosterol and α -tocoquinone.¹⁰⁻¹⁷ Leaves of this plant have been used as anti-bacterial, anti-tumour, astringent, acne, eczema, anti-pyretic, sedative, treatment of liver disorder, vermifuge, anti-parasitical, insect repellent, anti-inflammatory, catarrhal fever, remedy for cough, anti-ulcer, to treat skin disease, hair tonic to promote hair growth.¹⁸⁻²³

MATERIALS AND METHODS

In vitro studies of *Vitex negundo* Linn

Preparation of Leaves Extract

The leaves of *Vitex negundo* Linn were collected from local area, cleansed and shade dried for a week, milled and defatted using petroleum ether and macerated using ethyl acetate for 72hrs. The extract was filtered and the filtrate was subjected to rotatory evaporator. The residue was collected and stored in the desiccator for further studies.

Larvicidal Bio-Assay Procedure

Larvicidal activities of the ethyl acetate extract of *Vitex negundo* Linn was assessed by using protocol of World Health Organization (WHO) (2005).²³ The early third instar larvae of *Aedes aegypti*, *Aedes albopictus* and *Culex quinquefasciatus* were exposed to the extract concentrations in three replicates such as 31.25, 62.5, 125, 250 and 500 ppm. Temephos of concentration 2.5, 5.0, 7.5 and 10.0 ppm were taken as positive control. Solvent control was also used. The extract with 10 larvae were introduced using half sealed pipette and mortality of the larvae was calculated after 24hrs. Larvae were considered dead, when it sinks at the bottom where it seems to be immobilized or it completely look pitch dark. Thus the same experiment was repeated for three species and percentage mortality was calculated by using following formula to all the three species.

$$\text{Percentage Mortality} = \left(\frac{\text{Number of dead larvae}}{\text{Number of larvae introduced}} \right) * 100$$

An *in silico* molecular docking study using Molegro Virtual Docker 2013 (ver. 6.0) software against *Culex quinquefasciatus* was performed and it showed better results in *in vitro* study.

Molecular docking studies

Preparation of Ligand

The three dimension (3D) structures of the active constituents of *Vitex negundo* Linn which possess larvicidal property (Vitexicarpin, Luteolin, Vitexin and Vitexin rhamnoside) and standard drug (Azadirachtin H)²⁴ were retrieved either from PubChem chemical databases or drawn using Chem Sketch software²⁵ and it has saved in a .mol format file. The study ligand has imported to the workspace and preparation of the ligand was done.

Preparation of Enzyme

The target for docking study is selected as Odorant Binding Protein.²⁶⁻³² The analysis of docking study was done by initially selecting the target for the disease and followed by obtaining the three dimensional structure of Odorant Binding Protein (PDB ID: 2L2C) from protein data bank in .pdb file format.³³ In docking studies the search space of the simulation was exploited and subset region of 25.0 Angstroms around the active side cleft. The water molecules has taken to consideration and replaceable by score of 0.50.

Molecular Docking

Study of Molegro Virtual Docker (MVD) docking search algorithms and scoring functions

Molegro Virtual Docker (version 6.0) is a fast and flexible protein ligand docking simulation program and it allows carrying out docking simulation

in a fully integrated computational study. The MolDock software is a new empirical search algorithm and it combines differential evolution with a cavity prediction algorithm.³⁴ MolDock has an interactive optimization technique inspired by Darwinian Evolution Theory. The generate by new solution by recombination and mutation. Piecewise linear potential (PLP) is scoring function of MolDock, which is a simplified potential to whose parameters has fit to protein-ligand structures and a binding data scoring function.^{35,36} It has further extended in Generic Evolutionary Method for molecule DOCK (GEMDOCK) with a new hydrogen bonding term and charge schemes.³⁷

MolDock Optimizer - Parameter for docking search algorithms

In Molegro Virtual Docker, selected parameters were used for the guided differential evolution algorithm: number of runs = 5 (by checking constrain poses to cavity option), population size=50, maximum interactions = 2000, cross over rate = 0.9, and scaling factor = 0.5. A0, variance-based termination scheme was selected rather than root mean square deviation (RMSD). To ensure the most suitable binding mode in the binding cavity, pose clustering was employed, which may lead to multiple binding modes.

RESULTS

From Table 1, the result has founded that *Culex quinquefasciatus* have higher percentage mortality rate when compared to the other two species like *Aedes aegypti* and *Aedes albopictus* and graph was plotted against percentage (%) mortality versus log concentration. Temephos used as standard and it showed 100 % mortality at 10 ppm to all study species namely *Culex quinquefasciatus*, *Aedes aegypti* and *Aedes albopictus* and the results was showed in Table 2. Half-maximal effective concentration (EC₅₀) value for three species was determined using Graph Pad Prism (version 5.02).

The calculate half maximal effective dose concentration value of the crude extract against *Culex quinquefasciatus* was found to be 83.77 ppm (41.85-167.7 ppm), EC₅₀ value of the crude extract gainst *Aedes aegypti* was found to be 341.7 ppm (209.8-556.6 ppm) and EC₅₀ value of the crude extract against *Aedes albopictus* was found to be 487.9 ppm (319.5-745.1ppm) by using the Graph Pad Prism.

An ability of the *Vitex negundo* contain phytoconstituents to bind with the targets has given in terms of the MolDock Score study. MolDock Score used as a parameter for analysing to the docking results. The plant contain phytoconstituents are ranked according to their binding capability of MolDock score. The ligand possessing the highest molecular docking score shows a strong affinity towards its target protein (Table 3).

In silico docking analysis of phytoconstituents from *Vitex negundo* Linn on Odorant binding protein (PDB ID: 2L2C) which has ranking based on MolDock score, Rerank score and H-Bond is represented in Table 3. In Figure 1, eleven corresponds to the docking pose evaluated and captured by the ligand energy inspector tool in the Molegro virtual docker. The descriptor calculation for the ligands are represented in separate excel file. Moreover, the molecular docking results effective response was found more in Vitexicarpin (Figure 2), Luteolin (Figure 3), Vitexin (Figure 4) and Vitexin rhamnoside (Figure 5), which are present in the leaves of *Vitex negundo* Linn.

DISCUSSION

Vector borne diseases are rapidly increasing due to chloroquine and insecticidal resistance. Synthetic mosquito repellents may trigger undesirable hazardous interaction and leads to harmful effects in human. There are no vaccines or other specific treatments for arbovirus

Table 1: Larvicidal activity using ethyl acetate extract of *V. negundo* Linn against *Culex quinquefasciatus*, *Aedes aegypti* and *Aedes albopictus*.

Ethyl acetate extract of <i>Vitex negundo</i> Linn										Solvent control
31.25 ppm		62.5 ppm		125 ppm		250 ppm		500 ppm		
No of larvae dead	% of Mortality	No of larvae dead	% of Mortality	No of larvae dead	% of Mortality	No of larvae dead	% of Mortality	No of larvae dead	% of Mortality	% of Mortality
<i>Culex quinquefasciatus</i>										
2	20	4	40	5	50	8	80	10	100	10
1	10	3	30	7	70	8	80	10	100	0
1	10	4	40	7	70	9	90	10	100	0
<i>Aedes aegypti</i>										
1	10	1	10	2	20	5	50	6	60	0
0	0	0	0	1	10	4	40	7	70	0
0	0	1	10	3	30	5	50	7	70	10
<i>Aedes albopictus</i>										
0	0	0	0	2	20	4	40	6	60	0
0	0	1	10	2	20	4	40	5	50	0
0	0	0	0	1	10	3	30	4	40	0

Table 2: Larvicidal activity of positive control (Temephos) against different mosquito larvae.

Mosquito species	Treatment	Mortality (%)			
		1.0 ppm	2.5 ppm	5.0 ppm	10 ppm
<i>Culex quinquefasciatus</i>		99	100	100	100
<i>Aedes aegypti</i>	Temephos	79	98	100	100
<i>Aedes albopictus</i>		68	96	99	100

Table 3: In-silico docking analysis of phytoconstituents from *Vitex negundo* Linn on Odorant Binding Protein (PDB ID: 2L2C) ranking based on MolDock Score, Rerank Score and H Bond energy.

Ranking based on MolDock Score				
Name	Ligand	MolDock Score	Rerank Score	H Bond
[00]Azadirachtin H	Azadirachtin H	-110.77	-81.3716	0
[00]Vitexicarpin	Vitexicarpin	-87.3733	-81.9938	-2.5
[00]Luteolin	Luteolin	-85.2923	-78.947	0
[00]Vitexin	Vitexin	-76.2769	-63.8822	-5.72432
[00]Vitexin rhamnoside	Vitexin rhamnoside	-69.9377	-68.3276	-7.65766
Ranking based on Rerank Score				
[00]Vitexicarpin	Vitexicarpin	-87.3733	-81.9938	-2.5
[00]Azadirachtin H	Azadirachtin H	-110.77	-81.3716	0
[00]Luteolin	Luteolin	-85.2923	-78.947	0
[00]Vitexin rhamnoside	Vitexin rhamnoside	-69.9377	-68.3276	-7.65766
[00]Vitexin	Vitexin 2	-76.2769	-63.8822	-5.72432
Ranking based on H Bond energy				
[00]Vitexin rhamnoside	Vitexin rhamnoside	-69.9377	-68.3276	-7.65766
[00]Vitexin	Vitexin	-76.2769	-63.8822	-5.72432
[00]Vitexicarpin	Vitexicarpin	-87.3733	-81.9938	-2.5
[00]Luteolin	Luteolin	-85.2923	-78.947	0
[00]Azadirachtin H	Azadirachtin H	-110.77	-81.3716	0

least docking score demonstrate that high stability of the compound. The docking results suggested ligand Vitexicarpin formed a stable complex with a least docking score of -87.3733 and compared to standard Azadirachtin H (-110.77). The active constituent of the *Vitex negundo* on odorant binding protein was ranked according to their hydrogen bonding, rerank score and MolDock score. The molecular docking score was showed the least docking energy in Vitexicarpin and higher docking energy in Vitexin (-76.2769). This showed interacting residues ALA 88, MET 89, MET 91, VAL 125, GLY 92, LEU 58, LEU 73, LEU 80 of all amino acids contributed to the Vander Waal interactions as compared with standard Azadirachtin H and it is also had Vander Waal interaction contributing to the stability of the complex. Therefore, the present work, natural products are preferred and it has biodegradable, ecologically safe and have to produce the significant toxicity on target species and more safe and protection for human being.

CONCLUSION

The present work, larvicidal activity of *Vitex negundo* Linn against three mosquito species namely *Culex quinquefasciatus*, *Aedes albopictus* and *Aedes aegypti* were screened. The three species larvae were exposed to different concentration of *Vitex negundo* Linn and EC₅₀ value was calculated. Among three mosquito species *Culex quinquefasciatus* showed possible results followed by *Aedes aegypti* and then followed by *Aedes albopictus*. And further *in silico* docking study used in Molegro Virtual Docker (MVD) software against *Culex quinquefasciatus* as it showed better results in *in vitro* study. An *in silico* docking studies suggest that most of the phytoconstituents of *Vitex negundo* Linn possess larvicidal potential as such as standard drug Azadirachtin H. The further *in vivo* molecular pharmacodynamic and lethal toxicity studies may helpful to get a clear idea about the mechanism of action of active constituents of this plant produce larvicidal effect against species of *Culex quinquefasciatus* and possible to commercial purpose in near future.

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