



# *Invitro* and *Insilico* analysis of antioxidant activity of selected medicinal plants

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## Abstract

Medicinal plants are the important essential sources given by the god for all the living creatures. The usage of the medicinal plants are increased rapidly day by day. Antioxidants are substances that delay or prevent the oxidation of cellular oxidizable substrates. The present study deals with the three different medicinal plants for phytochemical analysis and molecular docking studies. The analysis was undergone and which showed the results like carbohydrates, flavonoids, alkaloids, and the molecular docking studies shows the antioxidant compound present in all the selected samples.

**Key words:** Medicinal plants, phytochemical analysis, Molecular docking, compounds, and antioxidant activity.

## Introduction

Plants play a vital role in our lives more than animals mainly due to their extraordinary array of diverse classes of biochemical with a variety of biological activities. The usage of the medicinal plant are increased rapidly day by day. Antioxidants are substances that delay or prevent the oxidation of cellular oxidizable substrates. They exert their effect by scavenging reactive oxygen species, activating a battery of detoxifying proteins or preventing the generation of reactive oxygen species. In recent years, there has been an increasing interest in finding natural antioxidants, which can protect the human body from free radicals and retard the progress of many chronic diseases, (Rashmi and Linu Mathew, 2012).

The main of the study is to go through with the morphological and medicinal properties of the selected medicinal plants. To conduct phytochemical screening for the evaluation of compounds in selected medicinal plants like *Justicia tranquebariensis* L.f., *Adhatoda vasica* Linn, and *Eclipta prostrata*, L., and to identify the primary metabolites in the methanolic leaf extract of the selected samples. The antioxidant properties also present in the selected medicinal plants through analyzing the Molecular Docking studies.

## Materials and Methods

### Study area

Tamil Nadu lies in the southernmost part of the Indian peninsula. Coimbatore is the city in Tamilnadu, South India. It is the capital city Kongunadu region and is often referred to as the Manchester of south India.

Kinathukadavu is situated in the district of Coimbatore, which has a pleasant climate due to the presence of forests to the north and cool winds blowing through the Palghat gap in the Western Ghats.

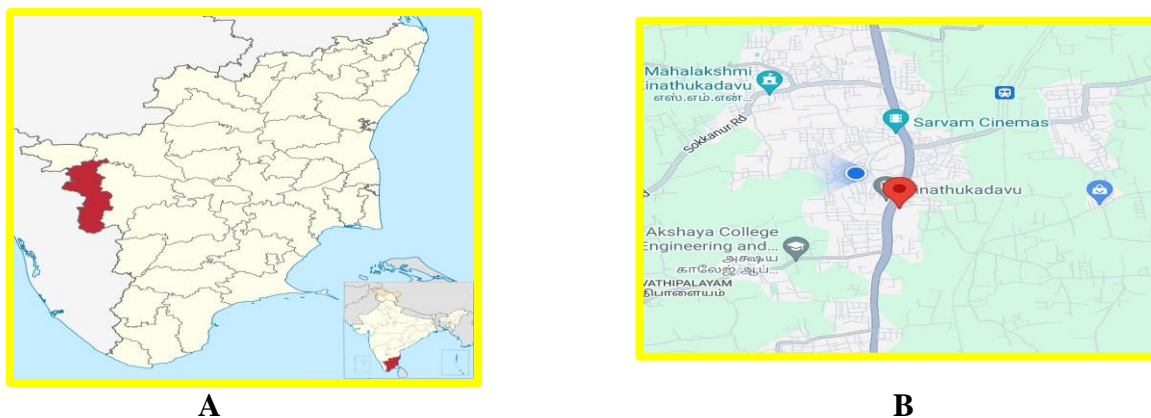


Fig-1: A) Study Area B) Location map

### Collection of the selected samples

For the present study the samples were collected and washed with running tap water to remove adhering dust and then dried under the shade. Then the dried samples were powdered with the help of a Pulverizer. The powdered samples were stored in an airtight container for further analysis.

### Sample-1 Systematic Position:

Division : Plantae  
 Class : Dicotyledonous  
 Order : Lamiales  
 Family : Acanthaceae  
 Genus : *Justica*  
 Species : *J. tranquebariensis* L.f.



Fig-2: A) Habit of *J. tranquebariensis* L.f. B) Powder of *J. tranquebariensis* L.f.

### Plant description

*Justicia tranquebariensis* L.f. is a small annual herb common to tropical countries. It is distributed throughout the hotter parts of India and Australia, often found in waste places along the roadsides. It is usually erect, slender-stemmed; spreading up to 80 cm tall, though sometimes it can be seen lying down. The plant is an annual broad-leaved herb that has a hairy stem with branches from the base to the top. The leaves are opposite, elliptical, oblong, or oblong-lanceolate with a faintly toothed margin and darker on the upper surface. Flowers are small, sexual, solitary, numerous and crowded together in axillary cymes about 1 cm in diameter. The fruits are yellow, three-celled capsules with wrinkled seeds. The stem and leaves reduce a white or milky juice that is frequently seen occupying open waste spaces, banks of watercourses, grasslands, road sides, and pathways. *Justicia tranquebariensis* L.f. is widely used as a traditional medicine herb in all the tropical countries (Begum *et al.*, 2011).

**Uses**

It is a popular herb amongst practitioners of traditional medicine, widely used as a decoction or infusion to treat various ailments including intestinal parasites, diarrhea, peptic ulcers, heartburn, vomiting, amoebic dysentery, asthma, bronchitis, fever, laryngeal spasms, emphysema, coughs, colds, kidney stones, menstrual problems, sterility, and venereal diseases.

**Sample-2 *Adhatoda vasica*, Linn.****Systematic Position:**

Kingdom : Plantae  
 Class : Dicotyledonous  
 Order : Lamiales  
 Family : Acanthaceae  
 Genus : *Adhatoda*  
 Species : *A. vasica*, Linn.

**A****B****Fig-3: A) Habit of *A. vasica*, Linn. B) Powder of *A. vasica*, Linn.,****Plant description**

*Adhatoda vasica* Linn, is a shrub with 10-20 lance-shaped leaves 8-9 centimeters in length by four wide. They are oppositely arranged, smooth-edged, and borne on short petioles. When dry they are of a dull brownish-green color. They are bitter- tasting. When a leaf is cleared with chloral hydrate and examined microscopically the oval stomata can be seen. They are surrounded by two crescent-shaped cells at right angles to the ostiole. The epidermis bears simple one- to three-celled warty hairs, and small glandular hairs. Cystoliths occur beneath the epidermis of the underside of the blade, (Wantkhem Sandhya chanu and Kanambala sarangthem, 2014).

**Uses**

The leaves are containing phytochemicals such as alkaloids, tannins, saponins, phenols and flavonoids. The most important is vasicine, and a quinazoline alkaloid.

**Sample - 3 *Eclipta prostrata*, L.,**

Kingdom : Plantae  
 Class : Angiosperms  
 Order : Asterales  
 Family : Asteraceae  
 Genus : *Eclipta*  
 Species : *E. prostrata*, L.,





A

Fig-3: A) Habit of *E. prostrata.L.*

B

B) Powder of *E. prostrata.L.*,

### Plant description

It is widely distributed throughout India, Nepal, China, Thailand, Bangladesh, and Brazil, In India it is widely used as a cholagogue and deobstruent in hepatic enlargement. Plant has cylindrical, grayish roots. Solid, circular, purplish stems with white fine hairs 0.8m. Leaves arranged in opposite pairs, hairy in two-sided, lanceolate, serrated 2–12.5 cm long, 5-35 mm wide. The solitary flower heads are 6-8 mm (0.24 -0.31 in) in diameter, with white florets. This species grows commonly in moist places in warm temperate to tropical areas worldwide (Shipam Sinha and Richa Raghuvanshi, 2016).

### Uses:

Contains a wide range of active phytoconstituents, which includes coumestan derivatives, triterpene saponins, steroidal saponins, triterpene, steroids, steroidal alkaloids, flavonoids, phenolic acids, thiophene derivatives and many other compounds.

### I. Preliminary phytochemical analysis - (Malathi *et al.*, 2018.)

#### Qualitative phytochemical tests are performed as preliminary tests.

For the preliminary phytochemical screening of methanolic leaf extracts of selected samples were analyzed by the standard methods and showed the presence of various phytochemical constituents such as saponins, phenols, alkaloids, proteins, tannins, carbohydrates and terpenoids.

#### Test of saponins

2 ml of extract was shaken vigorously with 5 ml distilled water to obtain stable persistent foam. The formation of emulsion indicates the presence of saponins.

#### Test for phenols

To 1 ml extract, add 5 ml distilled water followed by drops of 10% ferric chloride, the formation of blue or black color indicates the presence of phenolic groups.

#### Test for alkaloids

1 ml of extract in a test tube was treated with a few drops of Mayer's reagent Creamy white precipitate indicating the presence of alkaloids.

#### Test for protein

To 1 ml extract, add 2 ml water followed by a few drops of conc. HNO<sub>3</sub>. The formation of yellow color indicates the presence of protein.

#### Test for tannins

To 2ml extract, 1 ml of distilled water and 1-2 drops of ferric chloride solution was added and observed for brownish green or a blue-black coloration.

#### Test for flavonoids

A few drops of 1% NH<sub>3</sub> solution was added to 2 ml of extract in a test tube. A yellow coloration was observed for the flavonoids.

#### Test for carbohydrates and reducing sugars

5-8 drops of Fehling solution were added to 2 ml extract. The mixture was heated in a boiling water bath for 5 minutes. A brick red precipitate shows the presence of reducing sugars.

### Test for Quinone

2ml extract was mixed with 2 ml of chloroform and 1 ml of 10% ammonia solution was added. The presence of a pink, red or violet color indicates the anthraquinones.

### Test for terpenoids

2ml extract was mixed with 2 ml chloroform in a test tube. To this, 3 ml cone. H<sub>2</sub>SO<sub>4</sub> was carefully added along the wall of the test tube to form a layer. An interface with a reddish-brown coloration the presence of terpenoids.

### Test for glycosides

1 ml of 50% H<sub>2</sub>SO<sub>4</sub> was added to the 2 ml of extract in a boiling tube. The mixture was heated in a boiling water bath for 5 minutes. 10 ml of Fehling's solution was added and boiled. A brick red precipitate indicated the presence of glycosides.

## II. Molecular Docking- (Tomasz J. Guzik *et al.*, 2006)

The 2D structures of the compounds were retrieved from PubChem (Kim *et al.*, 2023) in SMILES format. The Smiles format was translated into 3D structure PDB format using online Smiles Translator tool. The Molecular Weight, Hydrogen Bond Donors, Hydrogen Bond Acceptors and Log P, the Lipinski parameters along with Drug Likeness Score were calculated using MuleSoft Online Software that can be accessed at, the receptor for the docking was retrieved from Protein Data Bank (Berman *et al.*, 2000). Molecular Docking of the identified receptor and synthesized ligands was performed using Autodesk Tools (ADT) 1.5.6 (Morris *et al.*, 2009) by employing the following method:

### Preparation of Receptor and Ligand files

Autodock entails both the receptor and ligand in PDBQT format for assessing the binding affinity between them. PDBQT format restrains the atomic coordinates, partial charges and atom types. Initially, the receptor file in PDB format obtained from Protein Databank was accessed in Autodock Workspace. The water molecules in the receptor file were removed and implicit Hydrogen atoms were added. Finally, partial charges were added and the receptor file was saved in PDBQT format. Similarly, the ligand files in PDB format was retrieved by Autodock and saved in PDBQT format.

### Preparation of Grid and Dock Parameter files

Autogrid 4.2 program in ADT was used to perform the grid computation. The grids maps with a dimension of 60X60X60 and spacing of 0.375 Å were centered along the ligand binding site. The receptor and ligand files in PDBQT format along with the grid maps were saved as the grid parameter file to execute the Autogrid program. After the Autogrid calculation, Autodock parameter file was created with the receptor, ligand and selection of Autodock parameters.

### Docking

Docking was performed using Lamarckian Genetic Algorithm with 10 independent runs per ligand with an initial population of 150 randomly placed ligand on the receptor binding site. A maximum of  $2.5 \times 10^5$  evaluations on the energy will be carried out for  $27 \times 10^3$  generations with a mutation rate of 0.02 and a cross over rate of 0.80. The local-energy-minimization algorithm was limited to 100 steps for 6% of the population. To explore the conformational space of ligands, the overall translation steps was set to 0.2 Å, and the overall rotation and torsion rotation step were set to 5 in the docking studies. The auto dock 4.0 program in ADT was executed and the docking scores were reported using binding free energy energies in kcal/mol.

### Molecular Interactions Visualization using PyMOL

The bound complex with the receptor and ligand was visualized using PyMOL molecular Visualization Software (Schrodinger and DeLano, 2020). PyMOL is a user-sponsored molecular visualization system on an open-source foundation, maintained and distributed by Schrödinger. PyMOL, a cross-platform molecular graphics tool, has been widely used for three-dimensional (3D) visualization of proteins, nucleic acids, small molecules, electron densities, surfaces, and trajectories. It is also capable of editing molecules, ray tracing, and making movies.

## Results and Discussions

I. Phytochemical Analysis of the sample (1) *Justicia tranquebariensis* L.f. contains Alkaloids, Phenols, Tannins, Flavonoids, Carbohydrate, Steroids, Glycosides and Protein, sample (2) *Adhatoda vasica*, Linn., results with the presence of Alkaloids, carbohydrate, protein, flavonoids and terpenoids and the sample (3) *Eclipta prostrata*, L., reveals the presence of Alkaloids, Phenols, Tannins, Flavonoids, Carbohydrate, Steroids, terpenoids and Protein.

S. No	Compounds	<i>Justicia tranquebariensis</i> L.f.	<i>Adhatoda vasica</i> , Linn.,	<i>Eclipta prostrata</i> , Linn.,
1.	Carbohydrate	++	+	++
2.	Proteins	-	++	+
3.	Reducing sugars	-	-	-
4.	Steroids	++	-	++
5.	Glycosides	++	-	-
6.	Flavonoids	++	++	++
7.	Alkaloids	++	++	++
8.	Tannins	+	-	++
9.	Saponins	-	-	++
10.	Terpenoids	-	++	++

**Table - 1 Phytochemical Analysis of the selected medicinal plant**

++ Present + Partial present – Absent

## II. Molecular Docking

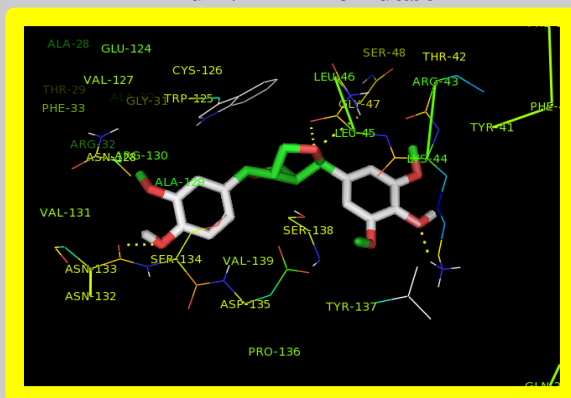
### Sample - 1 *Justicia tranquebariensis* L.f

Potential target for anti-oxidant activity is NADPH oxidase (Nox enzymes), that play a central role because they can regulate other enzymatic sources of ROS (Tomasz and David, 2006). Hence the Drug Target was chosen as NADPH oxidase and the structure was downloaded from PDB with PDB Id, 7u8g.

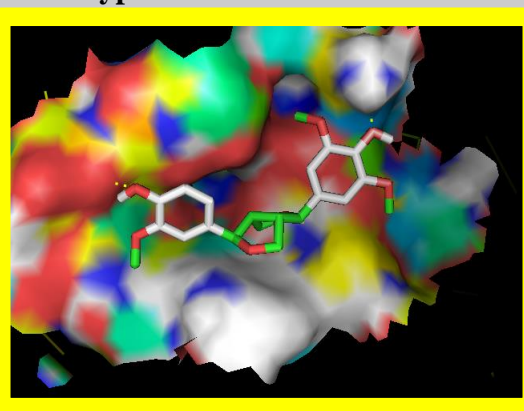
The binding energy between Isolariciresinol and NADPH oxidase, was -4.78 kcal/mol with a ligand Efficiency of -0.18 kcal/mol. 314.02  $\mu$ M of Isolariciresinol is required to acquire half maximum Inhibition in NADPH oxidase. There are 2 Hydrogen bonds formed between Leucine 45 and Glycine 47 of NADPH oxidase and Oxygen atoms of Isolariciresinol. The binding energy between 4-[(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c]furan-6-yl]-2,6-dimethoxyphenol and NADPH oxidase, was -5.98 kcal/mol with a ligand Efficiency of -0.21 kcal/mol. 41.48  $\mu$ M of **4-[(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c]furan-6-yl]-2,6-dimethoxyphenol** is required to acquire half maximum Inhibition in NADPH oxidase. There are 2 Hydrogen bonds formed between Asparagine 133 and Lysine 33 of NADPH oxidase and Hydrogen and Oxygen atoms of 4-[(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c]furan-6-yl]-2,6-dimethoxyphenol respectively.

The Docking of the Phytocompounds was compared with the Standard Ascorbic acid and the binding energy between Ascorbic acid and NADPH oxidase, was -4.64 kcal/mol with a ligand Efficiency of -0.39 kcal/mol. 397.52  $\mu$ M of Ascorbic acid is required to acquire half maximum Inhibition in NADPH oxidase. There are four Hydrogen bonds formed between O of Arginine 43 and Oxygen of Ascorbic acid and another Hydrogen bond between Glycine 47 and Oxygen atom of Ascorbic acid. Among the compounds that were docked both the Compounds, Isolariciresinol and 4-[(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c]furan-6-yl]-2,6-dimethoxyphenol showed better affinity than the Standard Ascorbic acid due to its least binding energy. Also, the IC<sub>50</sub> value 4-[(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c]furan-6-yl]-2,6-dimethoxyphenol was 41.48  $\mu$ M which is much less than the Standard Ascorbic Acid as well as Isolariciresinol, thus can be a potential candidate for anti-oxidant activity

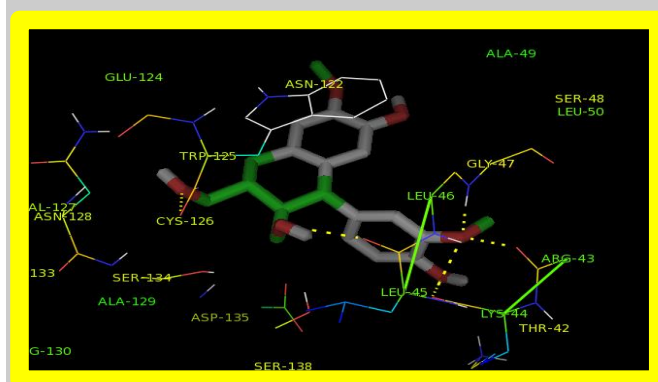
**Docked conformation between 4-[(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c]furan-6-yl]-2,6-dimethoxyphenol And NADPH oxidase**



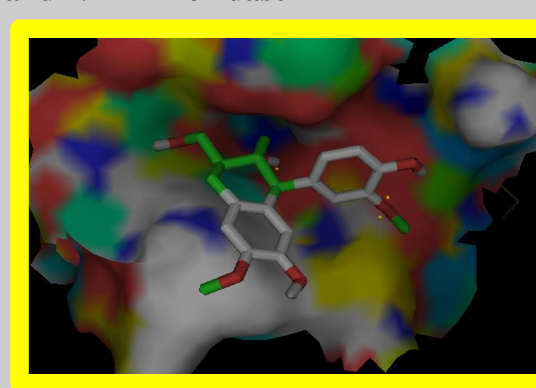
**Surface representation of Docked conformation between 4-[(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c]furan-6-yl]-2,6-dimethoxyphenol and NADPH oxidase**



**Docked conformation between Isolariciresinol and NADPH oxidase**

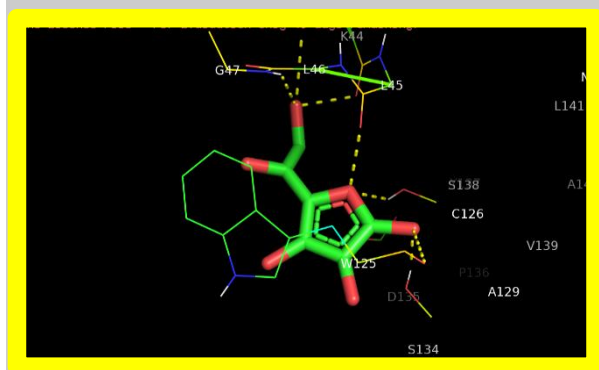


**Surface representation of Docked conformation between Isolariciresinol and NADPH oxidase**





### Docked Conformation of Ascorbic Acid and NADPH Oxidase



### Surface representation of Docked Conformation of Ascorbic Acid and NADPH Oxidase

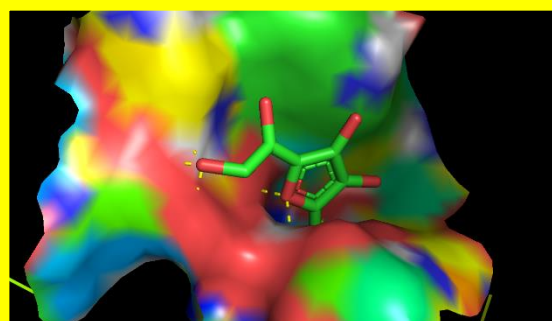


Table - 2 Anti-Oxidant Activity of *Justicia tranquebariensis* L.f.

S.No	Compound	Binding energy (KJ/mol)	Ligand efficiency (KJ/mol)	Inhibitory constant	No. Of hydrogen bonds	Hydrogen bond
1.	Isolariciresinol	-4.78	-0.18	314.02 $\mu$ M	2	Lig:H::Leu45:O Gly47:N::Lig:O
2.	4-[(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c]furan-6-yl]-2,6-dimethoxyphenol	-5.98	-0.21	41.48 $\mu$ M	2	Lig:H::Asn133:O Lys33:H::Lig:O
3.	Ascorbic acid	-4.64	-0.39	397.52 $\mu$ M	4	Arg43:O::LigO Gly47:NH::Lig:O Ser134:HG::Lig:O Leu45:O::Lig:O

### Sample - 2 *Adhatoda vasica* Linn,

Potential target for anti-oxidant activity is NADPH oxidase (Nox enzymes), that play a central role because they can regulate other enzymatic sources of ROS (Tomasz and David, 2006). Hence the Drug Target was chosen as NADPH oxidase and the structure was downloaded from PDB with PDB Id, 7u8g.

The binding energy between Luteolin and NADPH oxidase, was -5.73 kcal/mol with a ligand Efficiency of -0.27 kcal/mol. 62.85  $\mu$ M of Luteolin is required to acquire half maximum Inhibition in NADPH oxidase. There are three Hydrogen bonds formed between Aspartic acid 135, Serine 138 and Lysine 44 of NADPH oxidase and Oxygen/Hydrogen atoms of Luteolin.

The binding energy between **Kaempferol** and NADPH oxidase, was -5.86 kcal/mol with a ligand Efficiency of -0.28 kcal/mol. 50.26  $\mu$ M of **Kaempferol** is required to acquire half maximum Inhibition in NADPH oxidase. There are 4 Hydrogen bonds formed between Aspartic acid 135, Lysine 44, Asparagine 133 and Serine 138 of NADPH oxidase and Oxygen atoms of **Kaempferol**.

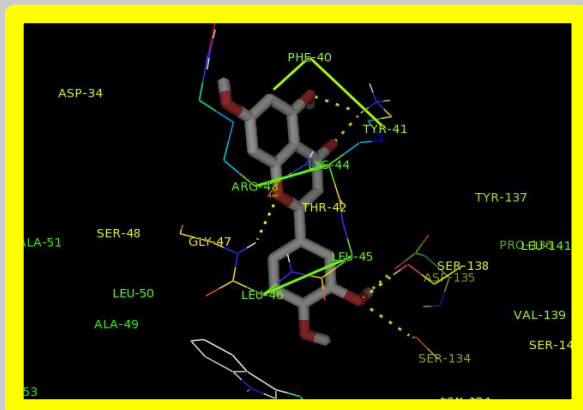
The Docking of the Phytocompounds was compared with the Standard Ascorbic acid and the binding energy



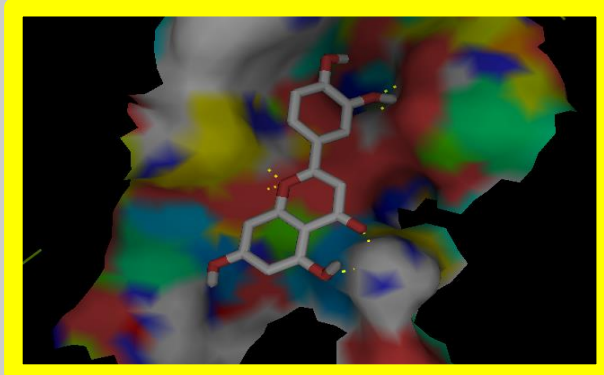
between Ascorbic acid and NADPH oxidase, was  $-4.64$  kcal/mol with a ligand Efficiency of  $-0.39$  kcal/mol.  $397.52$   $\mu\text{M}$  of Ascorbic acid is required to acquire half maximum Inhibition in NADPH oxidase. There are four Hydrogen bonds formed between O of Arginine 43 and Oxygen of Ascorbic acid and another Hydrogen bond between Glycine 47 and Oxygen atom of Ascorbic acid. The third Hydrogen bond is formed between Serine 134 of NADPH Oxidase and Oxygen of the Standard Ascorbic acid and the last hydrogen bond between Leucine 45 and Oxygen of Ligand.

Among the compounds that were docked both the Compounds, Luteolin and **Kaempferol** showed better affinity than the Standard Ascorbic acid due to its least binding energy. Also, the IC<sub>50</sub> value **Kaempferol** was  $50.26$   $\mu\text{M}$  which is much less than the Standard Ascorbic Acid as well as Luteolin, thus can be a potential candidate for anti-oxidant activity.

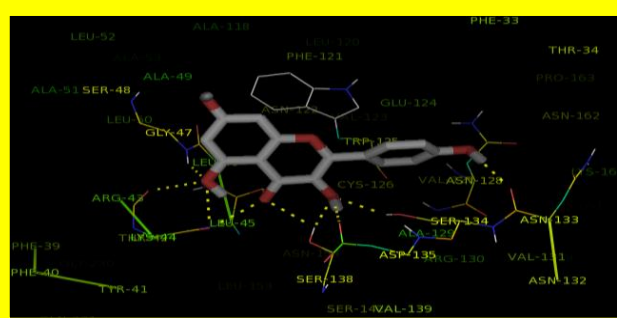
**Docked conformation between Luteolin and NADPH oxidase**



**Surface representation of Docked conformation between Luteolin and NADPH oxidase**

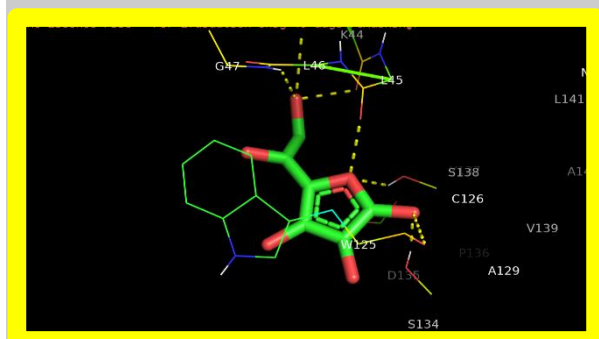
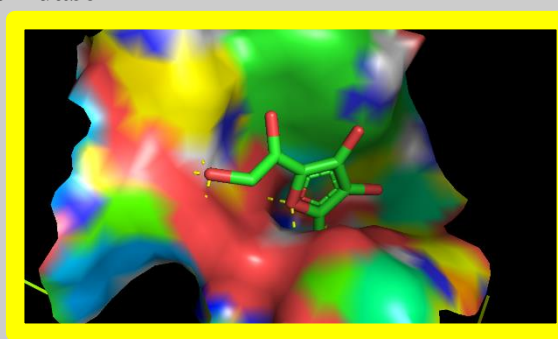


**Docked conformation between Kaempferol and NADPH oxidase**



**Surface representation of Docked conformation between Kaempferol and NADPH oxidase**



**Docked Conformation of Ascorbic Acid and NADPH Oxidase****Surface representation of Docked Conformation of Ascorbic Acid and NADPH Oxidase****Table - 3 Anti-oxidant Activity of *Adhatoda vasica* Linn,**

S. No	Compound	Binding energy (KJ/mol)	Ligand efficiency (KJ/mol)	Inhibitory constant	No. Of hydrogen bonds	Hydrogen bond
1.	Luteolin	-5.73	-0.27	62.85 $\mu$ M	3	Lig:H::Asp135:O Ser138:H::Lig:O Lys44:H::Lig:O
2.	Kaempferol	-5.86	-0.28	50.26 $\mu$ M	4	Lig:H::Asp135:O Lig:H::Lys44:O Lig:H::Asn133:O Ser138:H::Lig:O
3.	Ascorbic acid	-4.64	-0.39	397.52 $\mu$ M	4	Arg43:O::Lig:O Gly47:NH::Lig:O Ser134:HG::Lig:O Leu45:O::Lig:O

**Sample - 3 *Eclipta prostrata*.L**

Potential target for anti-oxidant activity is NADPH oxidase (Nox enzymes), that play a central role because they can regulate other enzymatic sources of ROS (Tomasz and David, 2006). Hence the Drug Target was chosen as NADPH oxidase and the structure was downloaded from PDB with PDB Id, 7u8g.

The binding energy between **Diosmetin** and NADPH oxidase, was -5.96 kcal/mol with a ligand Efficiency of -0.27 kcal/mol. 43.02  $\mu$ M of Diosmetin is required to acquire half maximum Inhibition in NADPH oxidase. There are three Hydrogen bonds formed between Lysine 44, Tryptophan 125 and Serine 134 of NADPH oxidase and Oxygen atoms of Diosmetin.

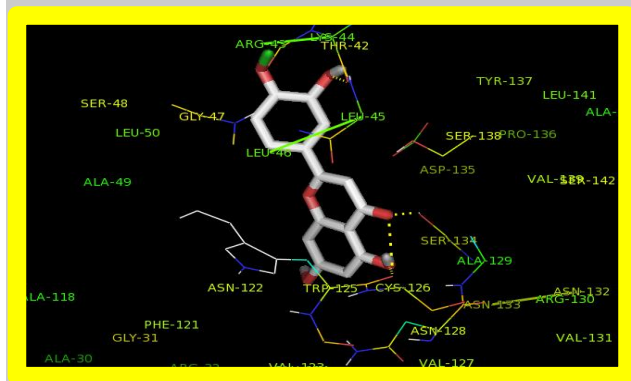
The binding energy between alpha-Terthienylmethanol and NADPH oxidase, was -5.66 kcal/mol with a ligand Efficiency of -0.33 kcal/mol. 71.44  $\mu$ M of alpha-Terthienylmethanol is required to acquire half maximum Inhibition in NADPH oxidase. There is a Hydrogen bond formed between Asparagine 133 of NADPH oxidase and Oxygen atom of alpha- Terthienylmethanol.

The Docking of the Phytocompounds was compared with the Standard Ascorbic acid and the binding energy between Ascorbic acid and NADPH oxidase, was -4.64 kcal/mol with a ligand Efficiency of -0.39 kcal/mol. 397.52  $\mu$ M of Ascorbic acid is required to acquire half maximum Inhibition in NADPH oxidase. There are four Hydrogen bonds formed between O of Arginine 43 and Oxygen of Ascorbic acid and another Hydrogen bond between Glycine 47 and Oxygen atom of Ascorbic acid. The third Hydrogen bond is formed between Serine 134 of NADPH Oxidase and Oxygen of the Standard Ascorbic acid and the last hydrogen bond between Leucine 45 and Oxygen of Ligand.

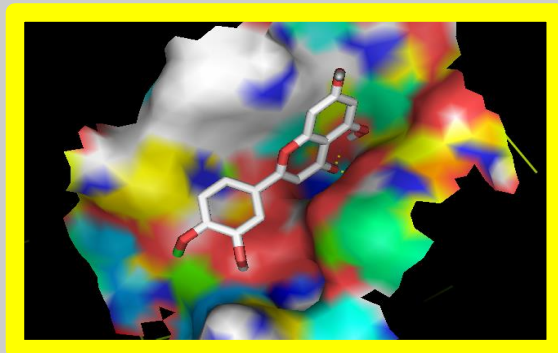
Among the compounds that were docked both the Compounds, **Diosmetin** and alpha- Terthienylmethanol showed better affinity than the Standard Ascorbic acid due to its least binding energy. Also, the IC50 value Diosmetin was 43.02  $\mu$ M which is much less than the Standard Ascorbic Acid as well as alpha-Terthienylmethanol, thus can

be a potential candidate for anti-oxidant activity.

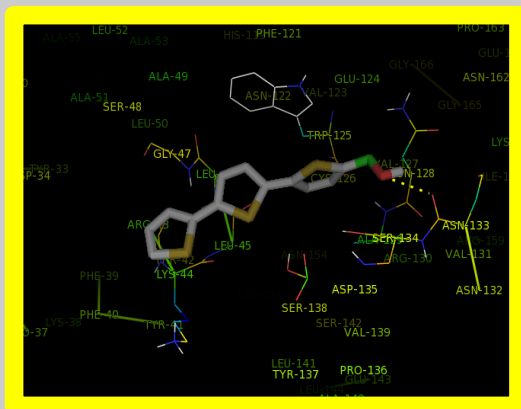
**Docked conformation between Diosmetin and NADPH oxidase**



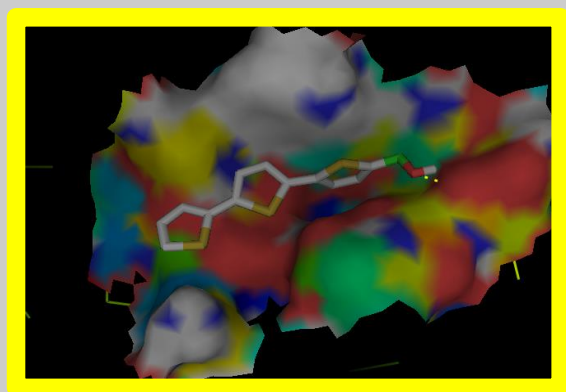
**Surface representation of Docked conformation between Diosmetin and NADPH oxidase**



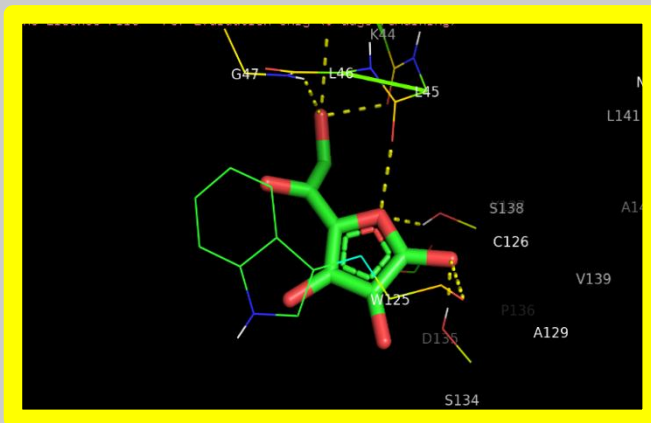
**Docked conformation between alpha-Terthienylmethanol and NADPH oxidase**



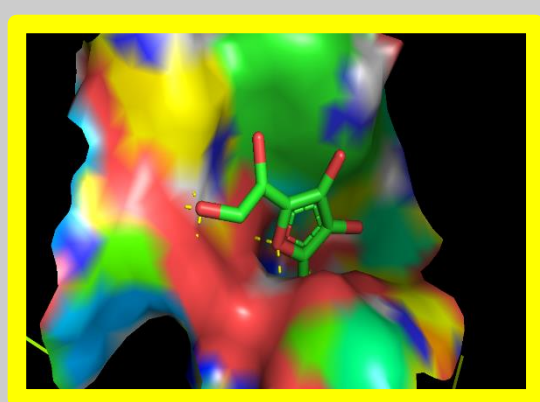
**Surface representation of Docked conformation between alpha-Terthienylmethanol and NADPH oxidase**



**Docked Conformation of Ascorbic Acid and NADPH Oxidase**



**Surface representation of Docked Conformation of Ascorbic Acid and NADPH Oxidase**



**Table -2 Anti-oxidant Activity of *Eclipta prostrata*, L.,**

S. No	Compound	Binding energy (KJ/mol)	Ligand efficiency (KJ/mol)	Inhibitory constant	No of hydrogen bonds	Hydrogen bond
1.	Diosmetin	-5.96	-0.27	43.02 $\mu$ M	3	Lig:O::Lys44:O Lig:H::Trp125:O Ser134:H::Lig:O
2.	alpha-Terthienylmethanol	-5.66	-0.33	71.44 $\mu$ M	1	Lig:H::Asn133:O
3.	Ascorbic acid	-4.64	-0.39	397.52 $\mu$ M	4	Arg43:O::Lig:O Gly47:NH::Lig:O Ser134:HG::Lig:O Leu45:O::Lig:O

### Summary and Conclusion

All the selected samples have the activities such as antibacterial, anticancer, antifungal, antioxidant, and antiproliferative activities. As the result of phytochemical analysis of the selected samples (*Justicia tranquebariensis* L.f., *Adhatoda vasica* Linn and *Eclipta prostrata*.L) *Eclipta prostrata*.L., have more number of compounds among the selected samples. And the molecular docking studies of the selected sample shows the different compounds like *Justicia tranquebariensis* L.f., contains Isolariciresinol, 4- [(3R,3aS,6R,6aS)-3-(4-hydroxy-3-methoxyphenyl)-1,3,3a,4,6,6a-hexahydrofuro[3,4-c] furan-6-yl]-2,6-dimethoxyphenol and Ascorbic acid. *Justicia adhatoda*, Linn., contains Luteolin, Kaempferol and Ascorbic acid. *Eclipta prostrata*.L contains Diosmetin, alpha-Terthienylmethanol and Ascorbic acid. Alkaloids directly act on the Central Nervous System in the human body. The compound tannins is useful in the therapy of diseases related to the oxidative stress such as colorectal cancer and heart and liver damage. Carbohydrate acts as an energy source helps in control blood glucose, insulin metabolism and helps in fermentation. Glycolysis helps in the sodium potassium metabolism. Protein reduces the appetite and gives strength to the bones and muscles and lowers the blood pressure. Flavonoids contains ant oxidative activity and good for heart diseases. Steroids will reduce the swelling and inflammation in our human body. Reducing sugar is good in weight gain. Terpenoids compounds are enrich in anti-inflammatory, anti-diabetic and used as a diuretic. Saponins decrease blood lipids, lower cancer risks and lower blood sugar level.

Among these compounds the selected medicinal plants contain carbohydrates, flavonoids, alkaloids in all the three plants. Hence these plants are used as a folk medicine for various diseases like fever, anemia, cold, cough and digestive diseases.

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