



## $\beta$ -Amyrin and Benzene-1,2,4-trimethyl from *Euphorbia hirta* L. and *Nauclea latifolia* (Smith) Leaves Induce Dauer Diapause via Antagonist Inhibition of daf-12 Receptor

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### ABSTRACT

Drug development begins with medicinal plants as valuable materials. *Euphorbia hirta* (EH) and *Nauclea latifolia* (NL) are important medicinal plants having applications in the management of infectious diseases. This research aimed at evaluating EH and NL leaves for their anthelmintic property and establishment of probable dauer diapause effect induced via interaction with daf-12 nuclear receptor in worms using standard procedures. Powdered plant samples were extracted using cold maceration for 72h. Qualitative, quantitative phytochemicals and antioxidants were determined using standard procedures. Metal concentrations were determined using Atomic Absorption Spectrophotometer. Anthelmintic assay was done by observing paralysis and death times of worms. Molecular docking of compounds isolated from plants was done using PyRx-Virtual Screening Tool workspace. SwissADME of the most promising compound was done with online tool. Molecular Dynamics simulation done using UCSF Chimera and graphic processor unit (GPU) version of the PMEMD.CUDA having AMBER package. Different concentrations (10, 30, and 50 mg/mL) of methanol and hexane leaf extracts of EH and NL displayed appreciable anthelmintic activity against *Pheritima pasthuma*, *Taenia solium* and *Fasciola gigantica* in the following order, *P. pasthuma* > *F. gigantica* > *T. solium*. The antagonist effect of Beta amyryn and Benzene-1,2,4-trimethyl on the key Amino acid residues in the binding site of Daf-12 nuclear receptor confer a pronounced dauer diapause effect in EH and NL. *In silico* assay showed Beta-amyryn having promising activity and appreciable stability in the binding site of the target and low bioavailability. This study justifies the reasons behind EH and NL usage as anthelmintic drugs locally

**Keywords:** Anthelmintic, daf-12 receptor, Dauer diapause, *Euphorbia hirta*, *Nauclea latifolia quadrifida*

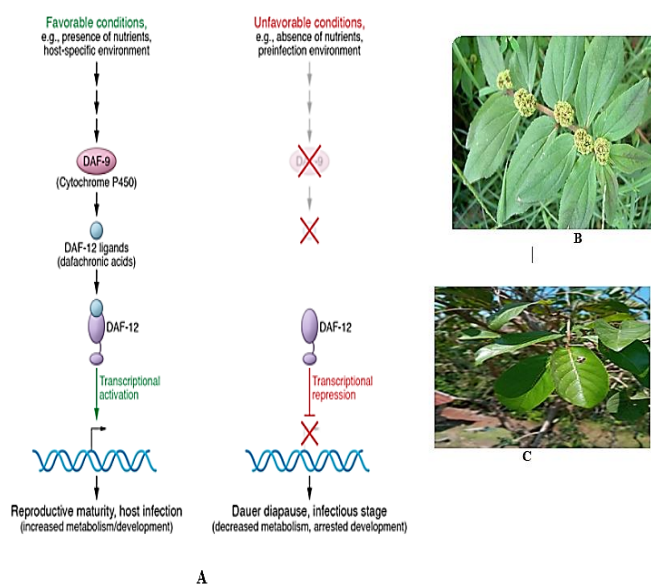
### Introduction

Helminths are a prevalent class among the various classes of parasitic organisms and infect over a billion people worldwide. They cause problems ranging from malnutrition, physical disabilities, mental retardation, stunted growth and eventually death. Studies have shown that over half of the over 40,000 presumed nematode species are parasitic in nature.<sup>1,2</sup>

Unfortunately, not many treatment regimens exist to combat these parasites or treat host patients to the parasites. The continuous use of anthelmintics for long periods has resulted in resistance over a wide range of time which is now common in livestock infected with parasitic nematodes.<sup>1</sup> As alternative therapies e.g. vaccines are lacking, novel drugs for the treatment of these infections are urgently needed, hence, the choice of medicinal plants,<sup>3,4</sup> or plant based bioactive principle(s) targeting a novel receptor known as nuclear receptor (daf-12) Figure 1A (2–6).

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**Figure 1:** A - Role of daf-12 in reproduction and development of parasite in host organism, B - *Euphorbia hirta*, C - *Nauclea latifolia* in their natural habitats

Ligand-gated transcription factors regulating various biological processes such as metabolism, development and reproduction are known as nuclear receptors.<sup>6</sup>

Nuclear receptors are now attractive targets for developing orally available small-molecule drugs because of their ligands' lipophilic nature and the expression of multiple genes being modulated by them within the same pathway.<sup>6,10</sup> A novel therapeutic strategy different from current anthelmintics, that focus on tubulins (Benzimidazoles), ion channels (Ivermectin and Diethylcarbamazine), or symbiotic bacteria of parasites (Doxycycline) could be developed by targeting nuclear receptors,<sup>5</sup> Figure 1A.8

In addition, there are numerous medicinal plants with ethnomedicinal usage in the treatment of helminths, most important of them are *Euphorbia hirta* and *Nuclea latifolia*.<sup>7,8</sup>

*Euphorbia hirta* (Family Euphorbiaceae) (Figure 1B) is an anthropogenic herb that is commonly found in open waste spaces widely throughout West Tropical Africa and it is well known for its antibiotics, diuretic and purgative properties.<sup>11,13,14</sup>

*Nauclea latifolia* (Family Rubiaceae) (Figure 1C), a straggling shrub or small tree, native to Tropical Africa and Asia is used in treating diarrhoea, fever and as an anti-parasitic drug.<sup>12</sup> However, this study was therefore designed to evaluate the anthelmintic properties of *E. hirta* and *N. latifolia* leaf extracts and to determine the mechanism of action of the bioactive principle(s) from both plants on nuclear receptor (daf-12).

## Materials and Methods

### Collection of plant materials

*Euphorbia hirta* fresh leaves were collected from Ikere-Ekiti market while the fresh leaves of *Nauclea latifolia* were collected in Ado-Ekiti (Global Positioning System (GPS) location of 7° 37' 15.9996" N and 5° 13' 17.0004" E), both in Ekiti State in June/July, 2020. The plants (leaves) were identified and authenticated by a plant taxonomist at the herbarium section of the Department of Plant Science and Biotechnology, Ekiti State University, Ado-Ekiti, Nigeria and voucher specimens of the two plants were deposited at the herbarium, with *Euphorbia hirta* and *Nauclea latifolia* having herbarium numbers UHAE 2020061 and UHAE 2020078, respectively.

### Preparation of plant extracts

The two plant samples were air dried at room temperature (about 20°C) for 14 days. The dried leaves of the two plants were powdered separately and stored in air tight bottles until needed. Hundred (100) grams of each plant powdered samples was soaked in a container with 1000 mL of methanol and hexane separately. The samples were shaken vigorously and kept for 72h using cold maceration method, this was further concentrated in vacuo employing a rotary evaporator (Serial Number: R000100958, Bibby Scientific Limited, Stone, Staffordshire, ST15 OSA, UK) at 45°C. Well labelled airtight bottles were used in storing the resulting extracts and kept in the refrigerator at 5°C till further use.<sup>9,10</sup>

### Phytochemicals Qualitative procedures

Preliminary phytochemical screening was done following standardized procedures as described.<sup>9, 10</sup>

### Phytochemical Quantitative Procedures

#### Determination of Total Phenolic Content

Five Hundred microliter (500 µL) of extract was placed into a test tube with Five Hundred microliter (500 µL) Folin-Ciocalteu solution with the addition of one milliliter (1 mL) Sodium carbonate solution and eight milliliter (8 mL) distilled water; the samples remained at room temperature for 30 minutes. The spectrophotometer at 760 nm wavelength was used and the total phenol content was expressed as milligrams of Tannic acid equivalents per gram of extract (mg TAE/g).<sup>11</sup>

#### Determination of Tannins Content

Five hundred (500 mg) milligram of casein was transferred into a 25 mL Erlenmeyer flask after which 5 mL each of the extracts and distilled

water were added. After two hours i.e. time required for the tannins to be complexed to the total protein, the filtered extracts were poured separately into a 10 mL volumetric flask and volume adjusted with distilled water. Five Hundred microliter (500 µL) extract was placed into a test tube, with the addition of Folin-Ciocalteu solution (500 µL), Sodium carbonate solution (1 mL) and distilled water (8 mL) with the samples remaining at room temperature for 30 minutes. The spectrophotometer (Spectrumlab 752s, Gallenkomp, England) was read at 760 nm wavelength and total phenol content expressed as milligrams of tannic acid equivalents per gram of extract (mg TAE/g).<sup>12,13</sup>

#### Determination of Flavonoids Content

Leaf extracts (500 µL) each was transferred into separate test tubes with the addition of Acetic acid solution (500 µL), Pyridine solution (2 mL), Aluminium chloride solution (1 mL) and 80% Methanol (6 mL); while samples were kept for 30 minutes at room temperature and the absorbance read at 420 nm. The flavonoid content was expressed as milligrams of quercetin equivalents per gram of extract (mg QE/g).<sup>14-16</sup>

#### Determination of alkaloids content

Each extract (2 g) of the leaves of *E. hirta* and *N. latifolia* were placed in a 250 mL beaker; 10% Acetic acid (80 mL) in Ethanol was added, covered and allowed to stand for 4 hours after which it was filtered and concentrated on a water bath (HH-S Water Baths, China) to about a quarter of the initial volume. Drops of concentrated Ammonium were added to the extracts until precipitation was completely achieved. The solution was allowed to settle with the precipitate collected, washed with dilute Ammonium hydroxide and filtered. The residue was dried and weighed and that is the alkaloid.<sup>17</sup>

#### Determination of saponins content

Two grams (2 g) of each powdered sample were weighed separately into conical flasks, with 20 mL of 20% aqueous Ethanol added. It was then heated over water bath for 4 hours while stirring continuously at 55°C. This mixture was filtered and the residue re-extracted with another 200 mL of 20% Ethanol. These extracts were reduced to 40 mL separately over water bath at about 90°C. Twenty millilitres (20 mL) of Diethyl ether was added to the concentrates inside a 250 mL separator funnel and shaken vigorously. The aqueous layer was recovered while the ether layer was discarded. There purification process was repeated with an addition of 60 mL n-butanol. Ten millilitres (10 mL) of 5% aqueous Sodium chloride was used in washing the combined n-butanol extracts twice; while the remaining solution was heated over the water bath. After evaporation, the samples were dried in the oven (DHG - 9035A, Gallenkomp, England) to a constant weight to give the saponins which was then calculated.<sup>18,19</sup>

$$\text{Saponins contents} = (W1 - W2) / W3$$

Where,

W1 = Weight of conical flasks + sample before drying

W2 = Weight of conical flasks + sample after drying

W3 = Weight of sample

#### Determination of Anthraquinones Content

Into a 100 mL volumetric flask was added 10mL of the sample which was adjusted to the required volume with solvent and analyzed at 325 nm wavelength using UV-Vis spectrophotometer (Spectrumlab 752s, Gallenkomp, England). Determination of the concentration was by checking the dilution factor of each sample in comparison to the standard anthraquinone solution.<sup>10</sup>

#### Determination of Steroids Content

To each of the test extracts (1 mL) in volumetric flasks (100 mL) was added Sulphuric acid (4N, 2 mL) and Iron (III) chloride (0.5% w/v, 2 mL), after which Potassium hexacyanoferrate (III) solution (0.5% w/v, 0.5 mL) was added. The mixture was heated in a water bath for 30 minutes with temperature at 70±2°C, shaken occasionally and diluted to reach the mark assigned in the volumetric flask with distilled water. Absorbance against the reagent blank was measured at 780nm.<sup>14</sup>

#### Determination of Terpenoids Content

Ten millilitres (10mL) Petroleum ether was used to partition one gram (1 g) of each extract in a separating funnel. The ether extract in pre-weighed glass vials was separated and waited for until completely dried. The yield (%) of total terpenoids contents was measured by the formula below after Petroleum ether was evaporated.<sup>20</sup>

$$\text{Yield (\% of total terpenoids)} = (W1 - W2) / W1 \times 100$$

Where,

W1 = Weight of Ether extract before drying

W2 = Weight of Ether extract after drying

#### Determination of Glycosides Content

Each sample (10 g) was treated with Lead acetate solution for tannins, proteins, colouring matter and other non-glycoside parts to be precipitated. Filtration of the precipitate formed was done and Hydrogen sulphide gas was made to pass through the filtrate to precipitate excess Lead as Lead sulphides, which was removed by filtration. Evaporation of the filtrate to dryness was done on a water bath (HH-S Water Baths, China) and the residue dried, collected and weighed to get the total glycoside content.<sup>20</sup>

#### Determination of Reducing Sugar

Each sample (100 mg) was weighed and the sugar extracted with hot Ethanol (80%). The supernatant after collection was evaporated at 80°C on a water bath. Distilled water (10 mL) was added to dissolve the sugars while 0.1mL was separated into test tubes using a pipette.

Preparation of working standard (0.2, 0.4, 0.6 and 0.8 mL) into series of test tubes was done. Both samples were made up to 2 mL each with distilled water in standard tubes. Two milliliters (2 mL) of distilled water was separated as blank and to each standard tube was added 1mL of alkaline Copper tartrate reagent. The two standard tubes were left for 10 minutes in a water bath. It was allowed to cool down and Arsenomolybdic acid (1 mL) was added to each of the test tubes and made up with distilled water to 10 mL. The absorbance was read at 620nm when the colour turned blue at the end of 10 minutes. The quantity of reducing sugars present was calculated from the graph drawn.<sup>10,21</sup>

#### Metal/Mineral Analysis

Spectrophotometry was the method used for analysing metal concentration using Atomic Absorption Spectrophotometer (AAS) Buck Scientific model 211 VGP employing the calibration plot method. The three processes involved were; preparation of standard, calibration of equipment and analysis of sample. The instrument was auto-zeroed using the blank (distilled water) for each element. The standard from the lowest to the highest concentrations was then aspirated into the flame. The graph of absorbance against concentration was plotted after the corresponding absorbance was obtained by the instrument. Analysis of the samples was done with the concentration of the metals present expressed in parts per million (ppm) having extrapolated from the standard curve.<sup>22</sup>

Radical scavenging activity of extracts using Diphenyl-2-Picryl-Hydrazyl (DPPH) method

The free radical scavenging capacity of the plant extracts against 1, 1 – Diphenyl – 2 – picryl-hydrazyl (DPPH) free radical was measured using the modified method;<sup>23</sup> DPPH (0.1 mM) was added to different concentrations of each sample. Vitamin C (positive control) was also used as standard drug at same concentration range (20 - 100 µg/mL), followed by incubation in an incubator (Model NO. 9052A, Gallenlomp, England) for one hour in the dark and optical density (OD) of the samples was measured at 517 nm against a blank and percentage (%) inhibition was calculated from the absorbance values.

$$\% \text{ DPPH inhibition} = (A - B) / A \times 100$$

Where A is the absorbance of pure DPPH in oxidized form while B is the absorbance of sample taken after 60 minutes of reaction with DPPH.<sup>11</sup>

#### Determination of Ferric Reducing Antioxidant Property (FRAP)

The reducing property of the extracts was determined by.<sup>11</sup> Each extract of about 0.25mL was mixed with 0.25 mL of 200 mM of Sodium phosphate buffer (pH 6.6) and 0.25mL of 1% Potassium Ferrocyanide (KFC). Incubation of the mixture was done for 20 minutes at 50°C, then 0.25mL of 10% TCA was added and centrifuged at 2000 rpm for 10 minutes; the supernatant (1mL) was mixed with 1mL of distilled water and 0.2 mL of 1% Ferric chloride and the absorbance was measured at 700 nm.<sup>1</sup>

#### Anthelmintic assay

##### Collection and authentication of worms

Taenia solium (Tapeworm) 2.2–2.6 g and Fasciola gigantica (Liverfluke) 0.04–0.08 g were obtained from a slaughtered cow at the Irasa Abattoir in Ado-Ekiti metropolis (M7W4+RG5, Ado-Iworoko road, 362103, Ilokun, Ekiti) by exactly 7:00 am in the morning. Pheritima pashuma (Earthworm, 0.09–0.8 g) were collected from a waterlogged area of Ekiti State University Farm with GPS location of N 7° 42' 50.8032", E 5° 15' 36.1332 at exactly 8:30am. The worms were identified at the Zoology and Environmental Science Department of Ekiti State University, Ado-Ekiti, Nigeria.

##### Bioassay

A worm was placed in each petri dish containing the extracts of E.hirta and N.latifolia using three different concentrations (10, 30 and 50 mg/mL in distilled water) respectively. This was carried out in duplicates for each worm type. The average (mean) times for paralyses (P, in minutes) were taken when no movement was observed, except when there was vigorous shaking of the worms. Death times of worms (D, in minutes) were recorded after no movement of worms either on their own or when vigorously shaken was ascertained. Antepar (10 mg/mL) was used as a standard drug i.e. reference compound, while distilled water served as the control.<sup>24-27</sup>

Molecular docking of the compounds isolated from E. hirta and N. latifolia on nuclear receptor daf-12

##### System Preparation

##### Retrieval and Preparation of Protein

The protein target preparation was accomplished through the 3D crystalline structure of worm nuclear receptor (PDB ID: 3UP3, Resolution: 1.25 Å) bound with its co-crystallized antagonists (25s)-cholestenic acid (CID: 10001991), obtained from Protein Data Bank (PDB). The protein crystal structure was expressed from Ancylostom acetylanicum, Homosapiens and was prepared before docking using UCSF Chimera 1.14 tool via removal of ions, water, and bound ligands which are non-standard residues. The structural minimization of the proteins was carried out at 200 steepest descent steps, 0.02 steepest descent steps size (Å), ten conjugate gradient steps, and 0.02 conjugate gradient steps size (Å) 10 update intervals using the structure editing wizard. There was removal of solvents, addition of Hydrogen bonds, allocation of charges through the Gasteiger force field and protonation state through setting of Histidine. Selenomethionine (MSE) present was changed to Methionine (MET), Bromo-UMP (5BU) to UMP (U), MethylselenylUMP (UMS) to UMP (U) and Methylsulfonyl-dCMP (CSL) to CMP (C).<sup>28</sup>

##### Ligand Selection and Preparation

This study made use of ligands that are 3D chemical structures of compounds isolated from E. hirta and N. latifolia. The structure data file (SDF) format of the compounds was retrieved from the PubChem database; PDB coordinates of all compounds were optimized by the UCSF Chimera tool, and charges were computed and added to the ligands through ANTECHAMBER.<sup>29</sup>

##### Molecular Docking

The prepared proteins and ligands were uploaded to the PyRx-Virtual Screening Tool workspace. A blind docking was initially carried out where the ligands were docked to the whole surface of the protein, followed by a site-specific docking to the co-factor and substrate binding pocket as identified in a previous study.<sup>30,31</sup> Using the Autodock Vina of PyRx software, the grid box was set for docking

based on the dimensions of the protein structure; each ligand-receptor binding.<sup>30, 31</sup>

#### Molecular interaction

Bovia Discovery 2020 Client was used to carry out molecular interaction (visualization) between the pharmacophore of the ligands and amino acid residues in the pocket of the indicated target.<sup>32</sup>

#### Swiss ADME of most promising compound

The online tool, namely: SwissADME (www.swissdock.ch) was used for detailed in silico physicochemical and pharmacokinetics studies of the compounds.<sup>33</sup>

#### Molecular Dynamics (MD)

The ligands and the receptor were prepared for the Molecular Dynamics simulation using UCSF Chimera. The MD simulation enquiries were done for the protein and complexes with the most potent compound (Apo+ligand) and cholestenic acid (Apo+control) with the aids of the graphic processor unit (GPU) version of the PMEMD.CUDA engine equipped with the AMBER package, FF18SB variant of the AMBER forcefield used in defining the protein. The stability of the complexes was estimated by computing MD simulations studies. The GENERAL AMBER Force field (GAFF) and the restrained electrostatic potential (RESP) methods aids in the addition of partial charges to the ligands and were improved using ANTECHAMBER. Neutralization and solvation were done with the assistance of LEAP module in AMBER 18 which helps in adding charges; H<sup>+</sup>, Na<sup>+</sup> and Cl counter ions to the unbound protein and ligand-bound systems. Also, all-atoms explicit solvation was carried out in an orthorhombic TIP3P box of water molecules sized 10 Å. The initial minimization (2500 steps) and complete minimization (5000 steps) were performed through a conjugate algorithm. Heating of the systems from 0 to 300 K were conducted for 50 ps. The total length of the simulation time was 200 ns with a time scale of 2 fs using a Langevin thermostat, at 300 K, and 1 bar constant pressure. The coordinates of the Apo-protein, Apo+control and Apo+ligand complexes were saved every 1 ps. All trajectories were studied for RMSD, RMSF and radius of gyration using CPTRAJ and PTRAJ module in AMBER 18/GPU. The investigates of post-MD structures and visualization were observed by using the graphical user interface of UCSF Chimera and Discovery studio 2021 client to enable the study of the mechanism of binding of the ligand-bound systems and MicroCAL Origin 6.0 software for data analysis was used to plot the data.<sup>34-36</sup>

#### Statistical analysis

Microsoft excel and Graph pad prism 7 (version 8) were used for the analyses of all data collected from the entire study. Presentation of the pertinent points was done using relevant tables, charts, and descriptive statistics. The mean  $\pm$  SEM of three replicates was used to express data for the in vitro model. One Way Analysis of Variance (ANOVA) complemented with Krystal-Wallis's test was used to statistically analyze data.

## Results and Discussion

#### Result of qualitative phytochemical screening

Different classes of phytochemicals (secondary metabolites) were observed in the two plants and these classes of compounds include: alkaloids, phenols, tannins, saponins, flavonoids, steroids, anthraquinones, cardiac glycosides, terpenoids and reduced sugar compounds (Table 1).

#### Quantitative phytochemistry of plant samples

According to the quantitative analysis shown in Table 2, *Euphorbia hirta* was observed to having high level of alkaloids of about 1243 mg/100 g of crude extracts, followed by saponins (787 mg/100g) with low amount of reduced sugar (0.4 mg/100 g). *Nauclea latifolia* was observed with high level of flavonoids (746 mg/100 g) and low level of reduced sugar of about 1 mg/100 g, which is higher than that of *Euphorbia hirta*.

#### Results of mineral analysis of plant samples

*Euphorbia hirta* and *Nauclea latifolia* exhibited high level of Potassium of about 2896 ppm and 314.8 ppm respectively and low amounts of Lead of 0.007 ppm and 0.002 ppm as shown in the Table 3.

**Table 1:** Preliminary phytochemical screening of *E. hirta* and *N. latifolia*

Phytochemicals	<i>Euphorbia hirta</i>	<i>Nauclea latifolia</i>
Alkaloids	+	+
Phenolics	+	+
Tannins	+	+
Saponins	+	+
Flavonoids	+	+
Steroids	+	-
Anthraquinones	+	+
Cardiac glycosides	+	-
Terpenoids	+	+
Reducing sugar	-	-

Keys: + = Present; - = Absent

**Table 2:** Amounts of Phytochemicals in *Euphorbia hirta* and *Nauclea latifolia* leaves

Phytochemicals (mg/100g)	<i>Euphorbia hirta</i>	<i>Nauclea latifolia</i>
Alkaloids	1243	388
Phenols (GAE/g)	46	68
Tannins	978	532
Saponins	787	532
Flavonoids	528	746
Steroids	182	108
Anthraquinones	113	87
Cardiac glycosides	35	13
Terpenoids	168	377
Reducing sugar	0.4	1

**Table 3:** Mineral Analysis of *Euphorbia hirta* and *Nauclea latifolia* leaf extracts

Parameters (ppm)	<i>Euphorbia hirta</i>	<i>Nauclea latifolia</i>
Na	161.55	108.35
Ca	140.60	92.25
K	2896.00	314.80
Mg	7.203	41.076
Fe	13.002	0.396
Mn	2.690	0.446
Pb	0.007	0.002
Zn	9.420	6.18
Cu	0.530	0.742
Cr	0.100	0.031
Se	0.108	0.079

**Antioxidant activity of the plant samples**

The effective concentration required for radical scavenging of *Euphorbia hirta* (EH) and *Nauclea latifolia* (NL) is 1.13 µg/mL and 1.63 µg/mL respectively, however from the result it was noted that the effective concentration of EH for DPPH is better than that of NL. Also, Ferric reducing antioxidant power (FRAP) assay concentration of EH was observed to be better when compared to NL. However, the Oxygen Radical Absorbance Capacity (ORAC) percentage inhibition showed that NL inhibited a larger amount of Peroxyl radical induced oxidation compared to EH as shown in Table 4.

**Anthelmintic studies of the plant extracts**

A decrease in time of paralysis (P) and death (D) was observed generally for the methanol extract of the two plants examined at different concentrations in a concentration dependent pattern. However, methanol extract of *E. hirta* at 50 mg/mL showed more potent activity against *P. pasthuma* with a time of paralysis and death of  $7.00 \pm 0.20$  and  $41.00 \pm 0.30$  respectively relative to the negative control (distilled water) at  $p < 0.05$  among the parasites examined as shown in Table 5. It is noteworthy to state that all the concentrations of methanol extract of EH had a significant effect on the time of paralysis and death of *T. solium*, *P. pasthuma* and *F. gigantica* relative to the negative control  $p < 0.05$ .

However, significant observation was noted, when an increase in the concentration of NL (10 – 50 mg/mL) influenced a decrease in both times of paralysis and death in *P. pasthuma* relative to other organisms. *Nauclea latifolia* at 10 and 50 mg/mL produced  $16.00 \pm 0.50$  (P);  $55.00 \pm 0.20$  (D) and  $6.00 \pm 0.10$  (P);  $30.00 \pm 0.20$  (D) respectively which is more potent than  $18.00 \pm 0.20$  (P);  $52.00 \pm 0.1$  (D) observed with 10 mg/mL Antepar (positive control) and distilled water (negative control) at  $p < 0.05$ .

A similar pattern of results observed above (Table 5) was noted for n-hexane especially concentration increase (10 – 50 mg/mL) induced a significant reduction in the time of paralysis (P) and death of *P.*

*pasthuma* relative to other worms (10 mg/mL –  $19.00 \pm 0.30$  (P);  $62.00 \pm 0.10$  (D) while 50 mg/mL –  $10.00 \pm 0.40$  (P);  $43.00 \pm 0.20$  (D) relative to 10 mg/mL antepar (positive control) and distilled water (negative control) at  $p < 0.05$ . In addition, the hexane extract of NL also exhibits a significant and promising effect on *T. solium* and *P. pasthuma* in a concentration-dependent manner with 50 mg/mL of the sample regarded as the most effective for the two parasites  $p < 0.05$  ( $13.00 \pm 0.10$  (P),  $40.00 \pm 0.50$  (D) and  $8.00 \pm 0.30$  (P);  $32.02 \pm 0.20$  (D) for *T. solium* and *P. pasthuma* respectively as shown in Table 6.

**Molecular docking of compounds isolated from *E. hirta* (EH) with Daf-12 receptor**

Compounds isolated from EH were subjected to molecular docking with daf-12 nuclear receptor present in all the worms examined, this resulted in beta-amyrin having the lowest binding affinity of -11.90 relative to -12.5 observed with (25s)-cholestenoic acid (crystallized ligand) as shown in Figure 2.

In addition, in silico screening of all compounds from NL showed that benene-1,2,4-Trimethyl exhibited the lowest binding energy of -10.7 among all the compounds examined as shown in Figure 3.

**Table 4:** *In vitro* Antioxidant Properties of *Euphorbia hirta* and *Nauclea latifolia* leaves

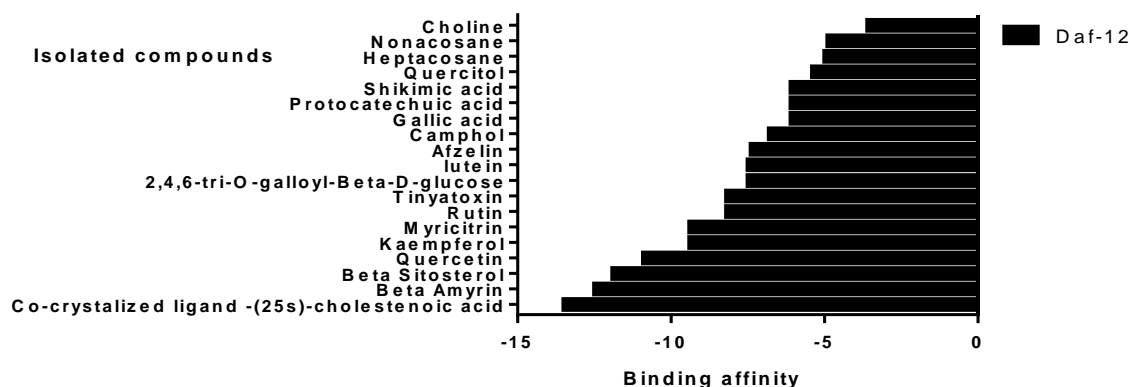
Parameters	<i>Euphorbia hirta</i>	<i>Nauclea latifolia</i>
DPPH (EC <sub>50</sub> )(µg/mL)	1.13	1.63
FRAP (mMFe <sup>++</sup> /Kg)	0.22	0.34
ORAC (% Inhibition)	55.26	61.63

Keys:

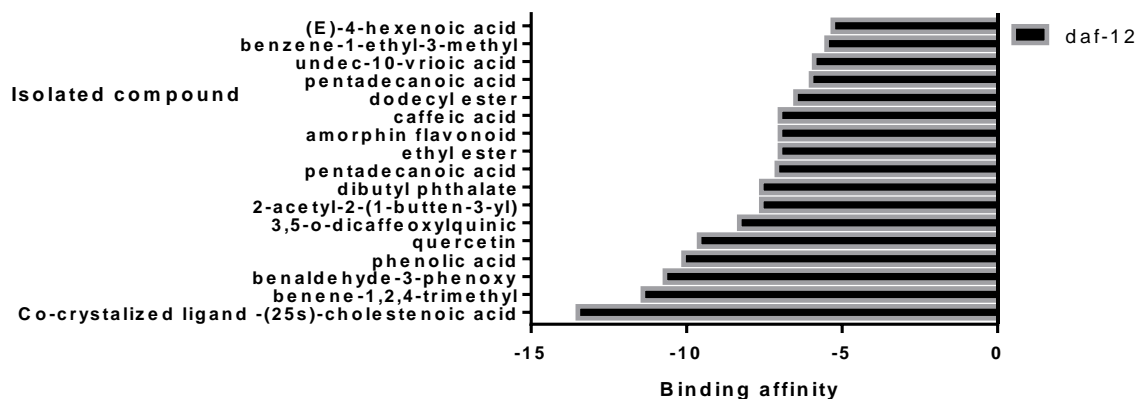
DPPH: 2,2-Diphenyl-1-Picrylhydrazyl

FRAP: Ferric Reducing Antioxidant Power

ORAC: Oxygen Radical Absorbance Capacity



**Figure 2:** Binding affinity of compounds isolated from *E. hirta* with Daf-12 receptor



**Figure 3:** Binding affinity of compounds isolated from *N. latifolia* with Daf-12 receptor

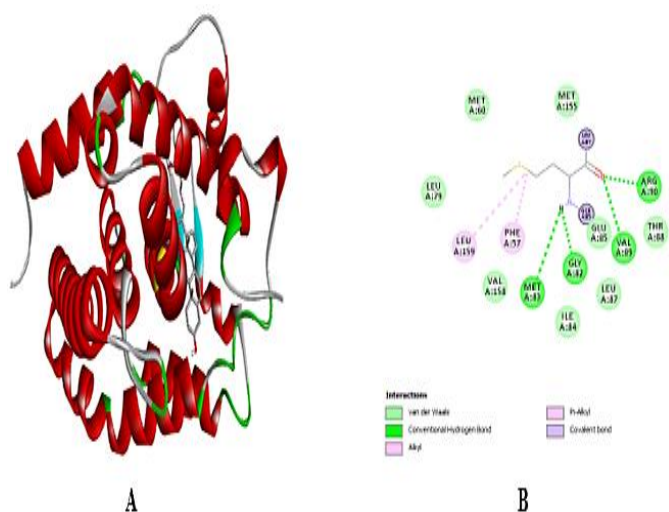


**Table 7:** Physicochemical properties of beta-amyrin isolated (hit compound) from *E. hirta*

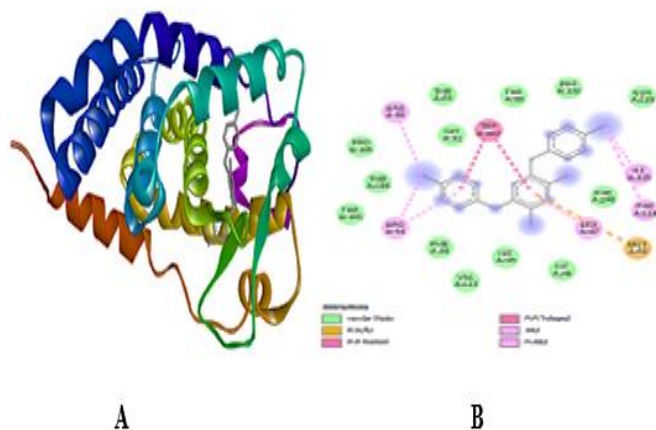
S/N	Compound Name	Molecular Weight g/mol	Heavy Atoms	TPSA	Rotatable Bonds	H-Bond Acceptor(s)	H-Bond Donor(s)	Molecular Refractivity	Fraction CSp <sup>3</sup>	Aromatic Heavy atom
1	$\beta$ -Amyrin	426.72	31	20.23	0	1	1	134.88	0.93	0

**Table 8:** Aqueous solubility of beta-amyrin isolated (hit compound) from *E. hirta*

S/N	Compound Name	Log S (ESOL)	Class	Log S (Ali)
1	$\beta$ -Amyrin	-8.25	Poorly soluble	-9.47

**Figure 5A - B:** Molecular interaction of (25s)-cholestenic (co-crystallized ligand) with nuclear receptor Daf-12

A - 3D interaction of (25s)-cholestenic (co-crystallized ligand) with daf-12; B - 2D interaction of (25s)-cholestenic (co-crystallized ligand) with daf-12

**Figure 6A - B:** Molecular interaction of Benene-1, 2,4-Trimethyl with nuclear receptor Daf-12

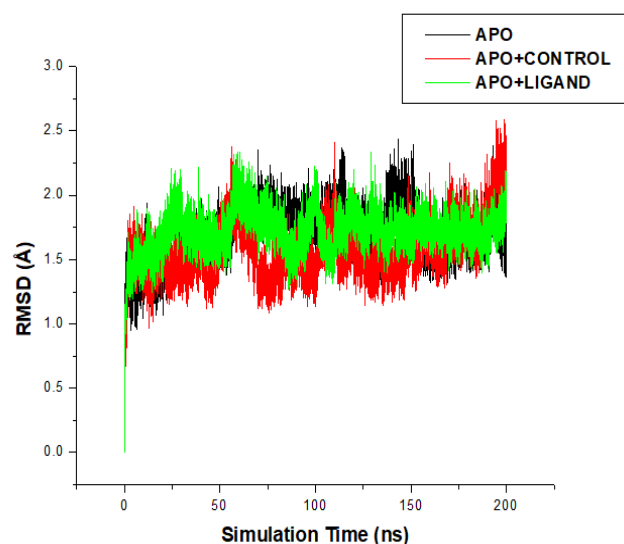
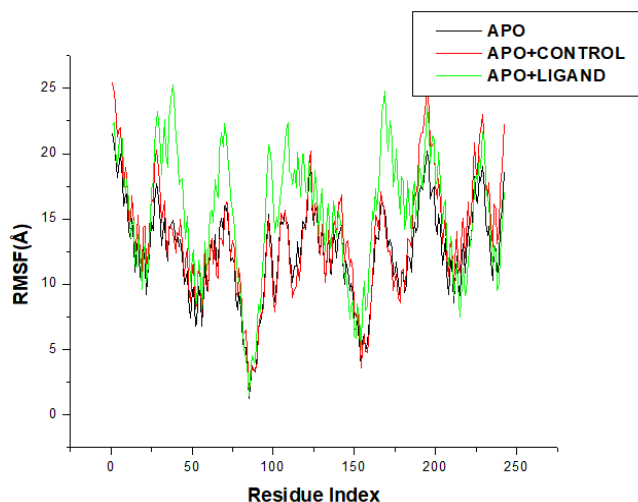
A - 3D interaction of benene-1,2,4-trimethyl with daf-12; B - 2D interaction of benene-1,2,4-trimethyl with daf-12

#### Molecular Dynamics (MD) Simulations

##### Root Mean Square Deviation (RMSD)

Root Mean Square Deviation (RMSD) helps in the determination of the union and stability of the two complexes and the unbound protein.<sup>45</sup> Long MD simulations time is essential for a complex system to attain

stability. An increase in the values of RMSD indicates that there is more atomistic deviation, while a low RMSD value indicates complex with a minimal structural abnormality and highly stable<sup>46</sup>; as shown from Figure 7 the estimated average RMSD values were 1.57 Å, 1.70 Å, and 1.72 Å for control-bound, beta amyrin-bound and the unbound protein complexes respectively. The results displayed that all the systems exhibited an average RMSD up to ~1.7Å at maximum, signifying a comparatively stable simulation. Similarly, the systems attained convergence virtually throughout the entire 200 ns simulation period. The results of the RMSD revealed that the binding of the control had a more stabilizing effect than Ligand and unbound-conformation. The highly stable binding pocket residues detected in the control bound could have favoured a steadier residue interaction with control and probably stronger binding to the protein relative to ligand.<sup>47</sup>

**Figure 7:** The RMSD for the complexes**Figure 8:** The RMSF of the complexes

**Root Mean Square Fluctuation (RMSF)**

Root Mean Square Fluctuation (RMSF) analysis gives understanding of the elasticity of the complex structure by evaluating the fluctuation of each amino acid existing in the protein. The conformational dynamics of the residues of the protein perform a significant role in the function of the protein.<sup>48</sup> The analysis of RMSF offers a better understanding of the fluctuations in conformation upon ligand binding and illustrates the implication of molecular dynamics at the atomistic level. The MD simulations trajectories in attaining the RMSF disclosed the alterations in the residues' flexibilities with or without a ligand. The RMSF was calculated for the residues that were established in the protein upon binding of the control and ligand to observe their effect on the conformational dynamics. The elevated values of RMSF showed more flexible activities, whereas the lower fluctuation values demonstrated lesser conformational changes upon the simulation.<sup>45</sup> The average RMSF values of control-bound, ligand-bound and the unbound protein complexes were 13.5 Å, 15.5 Å and 12.4 Å respectively (Figure 8). As shown, unbound protein had the least residue variability compared to control and ligand, suggesting increased protein residue variability upon control and ligand binding. Furthermore, the lower RMSF value for unbound protein compared to both control and ligand complexes may be indicated in the system's compactness.

**Radius of Gyration (RoG)**

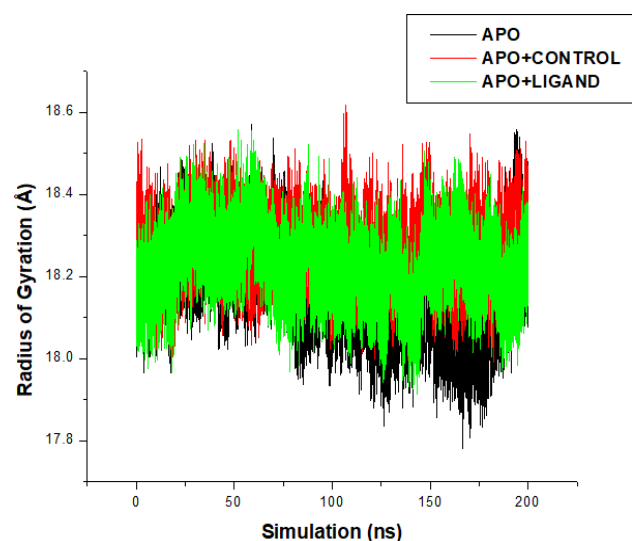
RoG helps to estimate the compactness of the entire protein structure during the course of the simulation.<sup>48</sup> The rigidity or slackness of the protein structure has the ability to interfere with its biological properties. It has been reported that the relationship between C- $\alpha$  RoG and compactness of protein structure revealed that forfeiture in structural compactness is as a result of high RoG value, and structurally compact protein is as a result of a low RoG value.<sup>49</sup> We also predicted the effects of binding of control and ligand on the compactness of the protein active site across the period of MD simulations. The average RoG values of control-bound, Ligand-bound and the unbound protein complexes were 18.3 Å, 18.2 Å, and 18.1 Å respectively (Figure 9), showing very similar compactness for the three systems. On the other hand, the results showed the atomic allocations in the unbound protein to indicate low RoG value and low compactness, which is correlative with the low RMSF value obtained. The high RoG observed in the conjugated system indicates that control and ligand binding resulted in a slightly looser protein conformation compared to the unconjugated system.

**Snapshot**

The Snapshot of interactions at different simulation times of both the control and the ligands with the protein are shown below. It was observed that there are differences in the residue's interaction in both the control and ligand which may account for their stability and compactness.

Plant antioxidants, in recent years have become an object of renewed interest, probably because of side effects of some commercial antioxidants which is undesirable. Medicinal plants have been endowed with diverse quantities of various kinds of bioactive chemical compounds having antioxidant properties which play a major role in stopping free radical chain reactions from being generated. Diphenyl Picryl Hydrazine (DPPH), a stable synthetic free radical has been used widely for determining natural compounds' antioxidant activities.<sup>11,40,41</sup> The DPPH is an ideal method due to it being easy, consistent and fast; water, methanol or ethanol do not also disintegrate

it. Hence, in this study, *E. hirta* and *N. latifolia* as potential antioxidant agents were tested for their radical scavenging and antioxidant activities by DPPH.<sup>11,16</sup> The DPPH scavenging assay finds its basis on the methanol DPPH solution's reduction when Hydrogen donating antioxidant is present leading to a non-radical form DPPH-H (Diphenyl Picryl Hydrazine) been formed. A change in colour from purple to yellow occurs which is proportional to the concentration and scavenging potential of the extract in terms of Hydrogen donating ability.<sup>11</sup> The EC<sub>50</sub> was the value calculated been the amount of antioxidant present in the sample needed to decrease the initial DPPH concentration by 50%. Antioxidant activity is higher when the EC<sub>50</sub> value is lower. *Euphorbia hirta* leaf extract had the lowest EC<sub>50</sub> value due to their high antioxidant activities. In the present study, *N. latifolia* leaf extracts exhibited higher percentage inhibition than *Euphorbia hirta* leaf extracts. Three comparable, complementary antioxidant assays were used in studying the quantitative antioxidant activities of *E. hirta* and *N. latifolia* leaf samples which were DPPH Free Radical Scavenging ability, Ferric Reducing Antioxidant Power (FRAP) and Oxygen Radical Absorbance Capacity (ORAC). *Nauclea latifolia* leaf samples exhibited a higher DPPH value (1.63 µg/mL) compared to *E. hirta* (1.13 µg/mL) (Table 4). This corroborates the quantitative antioxidant assay in the study carried out by.<sup>8, 42</sup> Results indicated that *N. latifolia* leaf sample was a better antioxidant agent compared to *E. hirta* leaves in its capacity to scavenge free radicals in the DPPH assay. The Ferric Reducing Antioxidant Power (FRAP) of *N. latifolia* and *E. hirta* were 3.43 mMFe<sup>++</sup>/Kg and 2.22 mMFe<sup>++</sup>/Kg respectively, still confirming *N. latifolia* as a better antioxidant agent. Oxygen Radical Absorbance antioxidant activities Capacity (ORAC) values of *N. latifolia* and *E. hirta* leaves were 61.6 % and 55.26 % respectively further proving *N. latifolia* leaves as a better antioxidant agent having a higher antioxidant activity (Table 4). Oxygen Radical Absorbance Capacity (ORAC) and FRAP values of *E. hirta* and *N. latifolia* leaves were significantly different at p<0.05 while DPPH values for the two plant leaf samples showed no significant difference at p<0.05.



**Figure 9:** The RoG of the complexes

**Table 9:** Lipophilicity of beta-amyrin isolated (hit compound) from *E. hirta*

S/N	Compound Name	iLog P	XLog P <sub>3</sub>	WLog P (Log Po/w)	MLog P	Silicos-ITLog P	Concensus (Log Po/w)
1	$\beta$ -Amyrin	4.74	9.15	8.17	6.92	6.92	7.18

**Table 10:** Pharmacokinetic Study of Compound Isolated from *E. hirta*

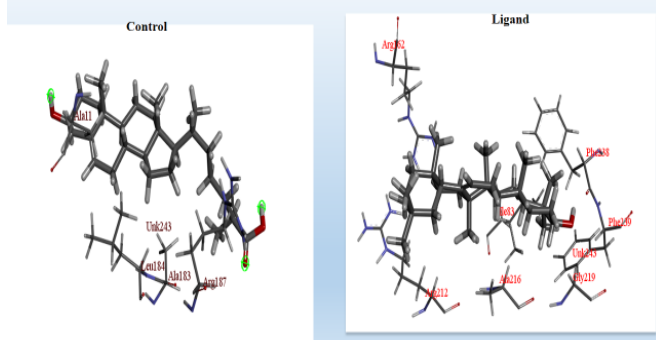
S/N	Compound Name	GI absorption	BBB Permeant	P-gp substrate	CYP1A2 inhibitor	CYP2C19 inhibitor	CYP2C9 inhibitor	CYP2D6 inhibitor	CYP3A4 inhibitor	Log Kp (cm/s)
1	$\beta$ -Amyrin	Low	No	No	No	No	No	No	No	-2.41



The mineral analyses of *E. hirta* and *N. latifolia* leaves as presented in Table 3 showed the presence of micro and macro nutrients in reasonable quantities in the two plants. In this study, it was observed that *E. hirta* was a richer source of most essential minerals as shown in Table 3 with Calcium value being 140.60 ppm compared to 92.25 ppm in *N. latifolia*; Potassium in *E. hirta* was 2896.00 ppm as compared to 314.80 ppm in *N. latifolia*; Sodium in *E. hirta* was 161.55 ppm compared to 108.35 ppm in *N. latifolia*; Zinc in *E. hirta* amounted to 9.42 ppm while 6.128 ppm was found in *N. latifolia*; Iron in *E. hirta* amounted to 13.02 ppm while 0.396 ppm was found in *N. latifolia*; Copper in *E. hirta* had a value of 0.530 ppm as compared to 0.742 ppm in *N. latifolia*; Manganese in *E. hirta* amounted to 2.69 ppm while 0.446 ppm of Manganese was found in *N. latifolia*; Chromium in *E. hirta* amounted to 0.10 ppm while 0.031 ppm was found in *N. latifolia*. Sodium and Potassium as the major extra cellular and intracellular minerals are involved in body water balance and acid-base balance. Hyponatremia is the result of failure to consume sufficient Sodium when fluid and Sodium losses are high. They are also involved in the transport of some non-electrolytes. Serious disorders like headache, tiredness, muscle cramps in the body can result from the maintenance of osmotic equilibrium by Sodium (Na).<sup>22</sup> Calcium is majorly found in the skeleton; the two plant leaves are rich in Calcium as it was present at high values in them. Bone forming and maintenance, muscle contraction and blood clotting are done by Calcium. A large number of enzymes are formed by Zinc, many of which function in wound healing and energy metabolism.<sup>22</sup> DNA synthesis, storage, release, function of insulin and the development of sexual organs and bones are also achieved by Zinc. Iron is necessary for the formation of haemoglobin. Poor Oxygen-carrying capacity, a condition causing endurance problems in athletes, is a characteristic of Iron deficiency or Anaemia. Bone and teeth formation need Phosphorus as an essential element. Manganese, a trace mineral is involved in bone formation, antioxidant activity, immune function and carbohydrate metabolism. Paralysis and convulsion may result as its deficiency. Potassium and Calcium were the most abundant mineral nutrients in the leaves of both plants used for this study. However, the elemental concentrations in both plants may be a reflection of the elemental composition of their parent soils.<sup>15</sup> The statistical results of the mineral analyses showed that Sodium, Calcium, Potassium, Magnesium, Iron, Manganese, Zinc, Copper and Chromium of *E. hirta* and *N. latifolia* leaves were significantly different at  $p < 0.05$  while Lead and Selenium showed no significant difference at  $p < 0.05$ .

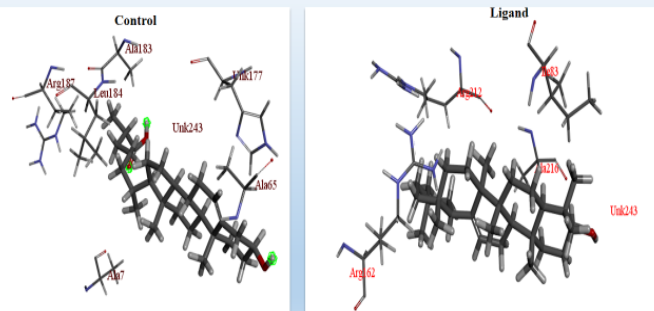
Most worm expellers like antepar functions to paralyze worms so that they are expelled in the faeces of both man and animals. The extracts, apart from causing paralysis to worms used, also caused their death, mostly at 50 mg/mL being the highest concentration used. As shown in Tables 5 and 6, the three concentrations of methanol and hexane leaf extracts of *E. hirta* and *N. latifolia* used (10, 30 and 50 mg/mL) exhibited profound anthelmintic activities. The anthelmintic activities of the plant extracts were dependent on concentrations for the three worm types (*Pheritima pashuma*, *Taenia solium* and *Fasciola gigantica*) used for this study; with the shortest time of paralysis (P) and death (D) achieved at 50 mg/mL for the three worm types used. This result corroborates some scientific findings.<sup>24, 25, 27</sup>

### Snapshot at 1ns



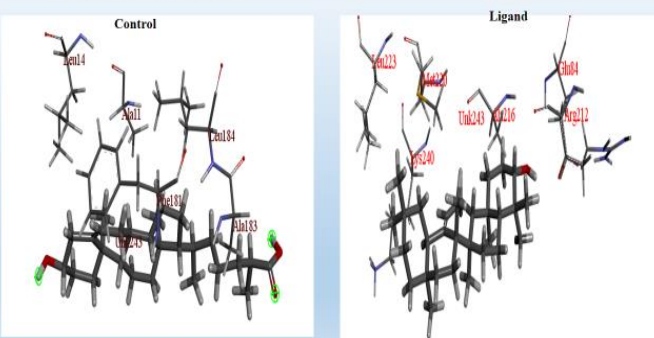
**Figure 10A:** The snapshot of the control and the ligands at 1ns

### Snapshot at 5ns



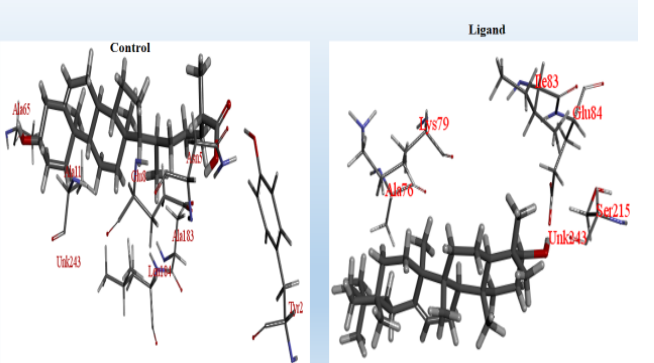
**Figure 10B:** The snapshot of the control and the ligands at 5ns

### Snapshot at 40ns



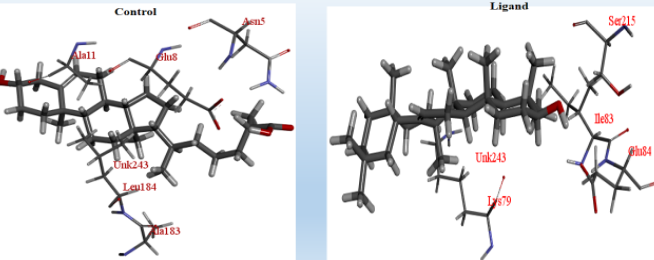
**Figure 10C:** The snapshot of the control and the ligands at 40ns

### Snapshot at 80ns

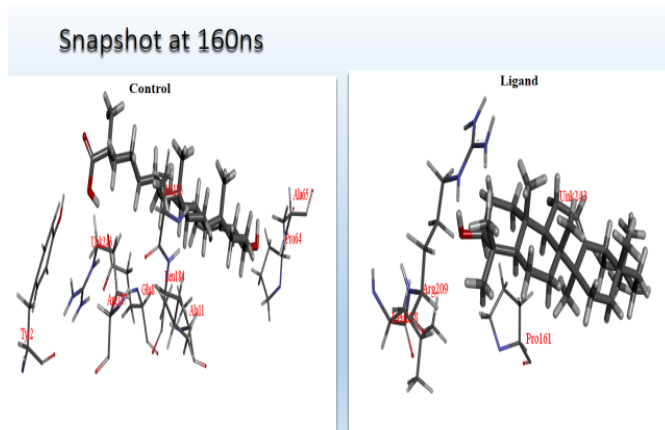


**Figure 10D:** The snapshot of the control and the ligands at 80ns

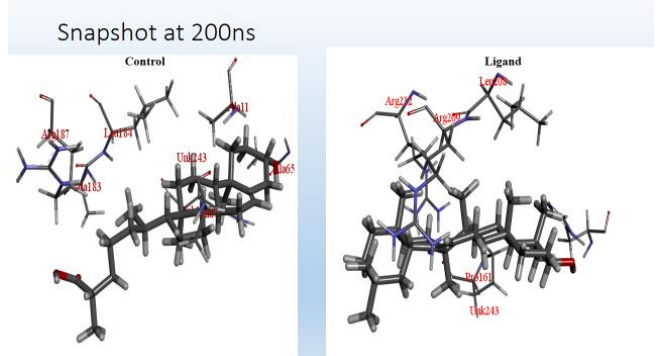
### Snapshot at 120ns



**Figure 10E:** The snapshot of the control and the ligands at 120ns



**Figure 10F:** The snapshot of the control and the ligands at 160ns



**Figure 10G:** The snapshot of the control and the ligands at 200ns

Tables 5 and 6 showed that methanol leaf extract of *N. latifolia* had a higher anthelmintic activity than the methanol leaf extract of *E. hirta* for the three worm types used; while hexane leaf extract of *N. latifolia* showed a higher anthelmintic activity than the hexane leaf extract of *E. hirta* for the three types of worms used. Generally, methanol leaf extract of *N. latifolia* exhibited the highest activity than other extracts used in the study. *Pheritima pasthuma* was most susceptible (sensitive) to the methanol leaf extract of *N. latifolia* (Table 5). It was paralysed within 6 minutes and died 30 minutes after the administration of *N. latifolia* methanol leaf extract; while paralysis (P) and death (D) times of *P. pasthuma* with the standard drug (Antepar) were 18 and 52 minutes respectively. *Pheritima pasthuma* was not easily killed by *E. hirta* hexane leaf extract (Table 6).

*Euphorbia hirta* methanol leaf extracts exhibited appreciable anthelmintic properties with *T. solium*. *Taenia solium* were paralyzed or died within 15-43 minutes of plant extracts administration while the worms were paralyzed or died within 5-43 minutes of the administration of 10 mg/mL dose of Antepar, a standard anthelmintic drug used as control. Another control group (worms in distilled water), lived for about 48 hours (Tables 5 and 6). Hexane leaf extract of *N. latifolia* showed highest anthelmintic activity against *P. pasthuma* with time of paralysis (P) = 8 mins and time of death (D) = 32mins while paralysis and death times for the standard or reference drug (Antepar) were 18 and 52 mins respectively (Table 6). *Pheritima pasthuma* was most sensitive to *N. latifolia* leaf extracts concentrations of 10, 30 and 50 mg/ml with time of paralysis (P) = 18±0.3, 13±0.5 and 8±0.3 minutes and death (D) = 56±0.1, 50±0.4 and 32±0.2 minutes respectively for hexane leaf extracts of *N. latifolia*; and with time of paralysis (P) = 16±0.5, 10±0.8 and 6±0.1 minutes and death (D) = 55±0.1, 48±0.1 and 30±0.2 minutes respectively for methanol leaf extracts of *N. latifolia* (Table 6). Generally, *P. pasthuma* was the most susceptible to the two plant leaf extracts, especially when compared to the standard drug, Antepar (10 mg/mL) at 50 mg/mL, paralysis time (P) for the earthworms was between 8–10 mins and death time (D) also in between 32–60 minutes. The extracts were more effective in causing paralysis of

*T. solium* (tapeworms) at 50 mg/ml rather than death. This result is also in line with the results.<sup>27</sup>

Times of paralysis and death were 18-45 mins for *E. hirta* hexane leaf extract; and 13-40 mins for *N. latifolia* hexane leaf extract; while for the reference drug, paralysis time (P) and death (D) was 8-41 mins (Table 6). The methanol extracts of *N. latifolia* showed reasonable anthelmintic properties with *F. gigantica* which became paralyzed and eventually died between 10–42mins; while *F. gigantica* in *E. hirta* methanol leaf extract was paralyzed and died between 9-25mins at a concentration of 50mg/mL. Antepar also paralyzed and killed the worms between 3-5 mins. Control (worms in distilled water) survived for a period of 48h (Table 5). Both *E. hirta* and *N. latifolia* hexane leaf extracts at 50 mg/mL concentration was also worthy of note; as *Fasciola gigantica* was paralyzed and died between 11-48 mins with *E. hirta* hexane leaf extract while they were paralyzed and died between 12-30 mins with *N. latifolia* hexane leaf extract. Antepar paralyzed and killed the worms around 3-5 mins. *Fasciola gigantica* in distilled water (control) lived for a period of 42h (Table 6).

However, molecular docking of compounds reportedly isolated from *E. hirta* and *N. latifolia* against daf-12 nuclear receptor showed that beta-Amyrin from *E. hirta* had a promising effect with the lowest binding affinity of -11.90 (Figure 3). The compound also had Hydrogen bond, van der Waals and pi-sigma interactions with some Amino acid residues in the binding site of the receptor (Figure 4A – B). Also, Benzene-1,2,4-trimethyl from *N. latifolia* showed a promising result (-10.7), which is higher than binding affinity of beta amyryn (-11.70), hence the lower activity of benzene-1,2,4-trimethyl relative to beta amyryn (Figures 2, 5A and 5B).

Apart from antioxidant, anti-inflammatory, antinociceptive, literature search showed no information relating beta-Amyrin to anthelmintics or daf-12 nuclear receptor, based on this promising role of beta-Amyrin, the compound can be a good drug candidate but might have challenges of gastrointestinal absorption as shown in Tables 7.<sup>10, 33,43</sup>

The structures and dynamics of biomolecules are vital facts provided by the approach of MD simulations and also help in the design of a potent drug. The arrangement of a protein structure is an important factor for its physiological activity; consequently, the arrangements could lead to positive or negative outcome on the activities of such protein when their structural integrity undergoes alteration.<sup>49</sup> As a result of this effect, co-crystallized-bound, beta amyryn-bound and the unbound protein complexes undergoes MD simulations of about 200 ns to investigate the energetics of each complex and confirm the convergence of dynamic stability. Trajectories obtained from the simulated systems were further examined for RMSD and RMSF to understand the structural dynamics and equilibrium during interaction and binding. The RoG and snapshot were also analysed for further understanding of the complexes as shown in Figures 7 – 10. The RMSD results provide information that the binding of the control process more stable effect on the binding domain than ligand (beta amyryn) and protein alone (unbound-conformation).<sup>50,51</sup> The favourable stability of binding pocket of Amino acids was obtained in the (25s)-cholestenoic (control/native ligand) bound could be ideal for tightly bounded residue interaction with control and perhaps stronger binding to the protein relative to ligand (Figure 7). RMSF (Root Mean Square Variation) results demonstrate data demonstrating lower amino acid variation for unbound protein compared to (25s)-cholestate (control/native ligand) and ligand (β-amyryn), indicating that the protein undergoes an increase in conformation in the course of binding controls and ligands (Figure 8). Furthermore, the lower RMSF values of the unbound protein compared to controls and ligand complexes could be due to the system's compactness.<sup>52</sup> However, the RoG result revealed that the atomic distributions in the unbound protein have a low RoG value (Figure 9) and low compactness, which correlated with the low RMSF value obtained in Figure 8. The high RoG was observed in the bound (beta amyryn and native ligand)-complexes indicated that the binding of control and ligand led to a little looser protein conformation relative to the unbound protein.<sup>53,54</sup>

## Conclusion

Conclusion therefore can be drawn that *E. hirta* and *N. latifolia* exhibited reasonable anthelmintic properties which back up their uses in ethnobotany (phytomedicine) as anthelmintic recipes. The dauer diapause effect on all the worms examined in the research might be due to antagonistic effect induced by multiple interactions between beta-Amyrin from *E. hirta* and Benzene-1,2,4-trimethyl from *N. latifolia* and some key Amino acids in the binding site of daf-12 nuclear receptor..

## Conflict of Interest

The authors declare no conflict of interest.

## Authors' Declaration

The authors hereby declare that the work presented in this article is original and that any liability for claims relating to the content of this article will be borne by them.

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