

Comparative analysis and *in-silico* studies of *Oecophylla smaragdina* collected from the states of Chhattisgarh and Maharashtra, India

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A. Collection of O.smaragdina from

areas

and

B. Drying the ants, eggs and grinding

of

Bastar,

Gadchiroli,

forested

in motor and pestle

Chhattisgarh

Maharashtra

the



- Collection of O.smaragdina from the forested areas of Bastar, Chhattisgarh and Gadchiroli, Maharashtra.
- Comparative analysis of biochemical and antimicrobial properties against microbes.
- To determine the elements (Crude protein, Zinc and calcium), Total phenolic content and DPPH activity.
- Determination of Anti-inflammatory properties and Cytotoxicity properties
- In-silico analysis and Protein docking studies of the proteins and genes found ant.

Methodology

Soxhlet based extraction in methanol solvent



BIOCHEMICAL CHARACTERIZATION

ANTIMICROBIAL ANALYSIS



The genes and proteins were retrieved from the NCBI database, protein 3D structure were modelled using SWISSmodel server, and the modelled 3D structures were assessed for their ProSA and Ramchandran Plot analysis. The 3D structure optimized using GROMACS package and P-P docking was carried out with the drug target of the gastrointestinal respiratory, Malarial, Dengue, Jaundice and COVID-19 (SARS CoV-2) infectious pathogens using the computational tool, Hex 8.0.0.

Results

Total Phenol content DPPH Radical scavenging 70.00%

MTT Assay % Survival





Antimicrobial activity of the methanolic extract against E. coli, S. aureus, S. pyogenes and K. pneumoniae

- The proximate analysis revealed a significant presence of Zinc in both the samples.
- The DPPH assay revealed former to have higher % inhibition and lower antioxidant activity i.e., 54.35 mg/AAE ml whereas Sample 2 showed lower % inhibition & higher antioxidant activity i.e., 86.9 mg/AAE ml.
- The total phenolic contents (Catechol equivalents, mg/g) in the samples were calculated to be 3.9853 and 5.228 mg Catechol eq/ml, respectively.
- Sample 2 demonstrated significant results and clearer zone of inhibition against all the four bacteria.
- The percentage survival of the cell was recorded as 76.76% (Sample 1) and 78.26% (Sample 2), with IC50 values of 23.5111± 0.4474 & 23.0719 ± 0.4425 respectively.
- No hyperactivation of other cells in blood culture were recorded, indicating that these samples do not alleviate any hypercell activity and are safe for utilization on human blood cells.

In-silico analysis

Ramachandran Plot data

	Highly	
rotein	preferred	Preferred
ytochrome b (partial)	96.71%	2.82%
ytochrome oxidase subunit 1, partial	97.02%	2.39%
ng wavelength rhodopsin, partial	98.69%	0.67%
ingless gene (wg)	91.54%	6.92%
ongation factor 1_ alpha 2 gene	91.78%	4.11%
rginine kinase gene	97.22%	2.78%
NA topoisomerase 1 gene	99.25%	0.76%
ytochrome C oxidase subunit 1, partial	98.18%	1.82%
ilk Fibroin	92.68%	2.44%



Protein-Protein Docking

75%

50%



Protein-Protein complex from Elongation factor protein from *S.aureus* and the EF-1 α -2 protein from *O. smaragdina*.





Protein-Protein complex of Hepatitis B virus and the Silk fibroin protein from O. smaragdina.



 RBC membrane stabilization demonstrated stabilization potentials for both the sample inflammatory potential of these samples. 	46.36% & 63.12% s, indicating the anti-	Protein-Protein complex of Cytochrome b oxidase from <i>P. falciparum</i> and the cytochrome b oxidase from <i>O. smaragdina</i> .	Protein-Protein complex of dengue virus corresponding genomic protein and the EF-1 α-2 from <i>O. smaragdina</i>	Protein- Protein complex of SARS Cov-2'O ribose methyltransferase protein from SARS Cov-2 and the Silk fibroin from <i>O. smaragdina</i> .		
 Red weaver ants are great reservoir for proteins, calcium and zinc which were obtained in significant amounts in both the samples. Free polyphenols with higher radical scavenging activity was observed in the sample 2. Due to the presence of high formic acid, the samples exhibited increased anti-microbial potential. The cytotoxicity assay and the anti-inflammatory studies suggests the consumption of red weaver ants are SAFE for humans. Protein-protein docking studies reported a strong affinity between both the proteins. From the <i>in-silico</i> analysis, here we predict the plausible interaction between the proteins from the red weaver ants that interacts to the proteins known as drug target from the selected pathogen and hence may be able to inhibit their pathogenicity. 						
No Conflict of Interest: The authors have no conflict of interest Acknowledgements: The authors thank the School or Biotechnology and Bioinformatics, D Y Patil Deemed to be University, India for all the facilities provided for research	References: References: Wheeler DE, Martínez T. Storage proteins in ants. Comp Biochem Physiol B Biochem Mol Biol. 1995, 112(1):15-9. doi: 10.1016/0305 0491(95)00035-7. Shanmugam Alagappan, et al 2021. Nutritional analysis, volatile composition, antimicrobial and antioxidant properties of Australian green and (O. smaragdina), Future Foods, 3,100007,https://doi.org/10.1016/j.fufo.2020.100007.					

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