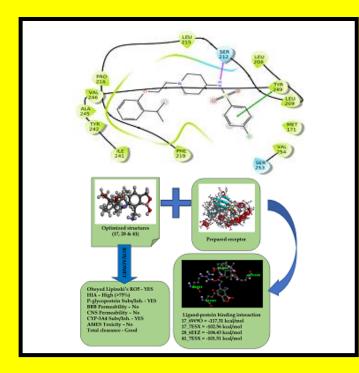
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Volume 2 Issue 1 June 2022



Pharmacophore Based Virtual Screening and Docking of Different Aryl Sulfonamide Derivatives of 5HT7R Antagonist

In Silico Approach: Effect of the Oxidation Iron State (Heme-Group) in Steroidogenesis Pathways



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### **EDITORIAL WORDS**

apt. Mohammad Rizki Fadhil Pratama, S.Farm., M.Si. Editor in Chief J Mol Docking

#### Assalamu'alaikum Wr. Wb.

Alhamdulillahirabbil 'alamin. After a long wait for almost 1 years from the first planned, finally the new scientific journal of the Department of Pharmacy Universitas Muhammadiyah Palangkaraya can be published. This scientifc journal was named **Journal of Molecular Docking** (*J Mol Docking*), inspired by one of the most popular *in silico* methods in computer-aided drug design. Journal of Molecular Docking published every 6 months (2 issues/year) every June and December.

This edition contains five articles consisting of writings from three countries including India, Mexico, and Nigeria. The authors come from several institutions, including Maulana Azad National Urdu University, Osmania University, Osmania University College for Women, Vignan's Foundation for Science, Technology & Research, Ahmadu Bello University, Benemérita Universidad Autónoma de Puebla, Universidad Autónoma de Guerrero, The Tamil Nadu Dr. M.G.R. Medical University, and Anna University.

Editorial boards are fully aware that there are still room for improvement in this edition, hence with all humility willing to accept constructive suggestions and feedback for improvements to the publication for the next editions. The editorial board would like to thank all editors and reviewers, and contributors of the scientific articles who have provided the repetoire in this issue. We hope that all parties, especially the contributors of the articles, could re-participate for the the publication in the next edition on December 2022.

Wassalamu'alaikum Wr. Wb.

Palangka Raya, June 2022

Editor-in-Chief

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# Author Guidelines JOURNAL OF MOLECULAR DOCKING

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- 1. Writing is the result of original research and literature review in the field of pharmacy, chemistry, bioinformatics, and others using *Molecular Docking Simulation* as its main analysis method which have never been published in other scientific publication media.
- 2. Manuscripts that have been published in the form of a preprint on several platforms such as arXiv, SSRN, and ResearchGate are welcome to publish.
- 3. The manuscript is written in English with Book Antiqua font 10 pt, two-column, density 1.15 spaces, on A4 paper with a top-down margin of 2.5 cm and right-left margins 1.5 cm.
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- 5. The manuscript is typed strictly with the following rules:
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    - c. Abstract, written in English which is a summary of the article. The abstract is created in one paragraph and a maximum of 250 words with Book Antiqua font 9 pt, single column, and space 1. Points that must be listed in the abstract include the research objectives, methods, results, conclusions, and keywords. The number of keywords is between three and five keywords.
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    - g. Conclusion, made in a single paragraph without the citation contains the final conclusion and suggestions for advanced research.
    - h. References, a list of recently selected topics is published last five years (minimum 75% of at least 10 references). The bibliography is written in appearance order chronologically with the serial number (Vancouver System) and suggested using a reference management application such as EndNote, Mendeley, Zotero, and other applications.
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- 7. Authors will get full access to the article published on e-journal of Universitas Muhammadiyah Palangkaraya at the website https://journal.umpr.ac.id/index.php/jmd.

