

# Review of: "Inhibition Success of a Virtually Created Molecule: Pseudoeriocitrin and Femtomolar Inhibition"

Hao Chi<sup>1</sup>

<sup>1</sup> Affiliated Hospital of Southwest Medical University

**Potential competing interests:** No potential competing interests to declare.

First and foremost, this paper demonstrates remarkable innovation in the field of molecular design. By exploring the virtually created molecule pseudoesperidin, the authors provide new perspectives for drug design, especially in the context of femtomole-level inhibitory effects. In addition, the study enriches our understanding of potential drug-protein interactions by delving into the interactions between pseudoesperidin and different target proteins through meticulous docking analysis. These findings have important implications for the development of new drugs and may have a positive impact on global health issues.

However, there are some key shortcomings of this study. The most obvious problem is the lack of experimental validation of the efficacy and safety of pseudoesperidin. In addition, the lack of hydrogen donors in the pseudoesperidin structure could lead to potential toxicity issues. The biological mechanisms behind the observed interactions were also not fully explored in the study, which is crucial for understanding the potential side effects and metabolic pathways of the molecule. Additionally, the reproducibility of the data and comparative analysis with existing compounds are aspects of the study that need to be further strengthened.

Overall, while this paper provides seminal insights in virtual drug design and molecular docking, it also highlights the need for more research, especially in experimental validation and toxicity assessment, to fully realize the potential of pseudoesperidin as a therapeutic agent.

Here are my suggestions:

**Advantages:**

**Innovation in Molecular Design:** The study's exploration of a virtually created molecule, pseudoeriocitrin, is innovative. It presents a novel approach to drug design, particularly in the context of femtomolar inhibition.

**Detailed Docking Analysis:** The manuscript offers a comprehensive docking analysis, examining interactions between pseudoeriocitrin and various target proteins. This thorough analysis enhances our understanding of potential drug-protein interactions.

**Significance in Drug Development:** The findings could significantly impact the development of new drugs, especially considering the molecule's femtomolar level of inhibition, which suggests high efficacy at low concentrations.

**Broad Scope of Application:** The study's implications for treating helminth infections are promising, potentially addressing a significant global health issue.

**Advancement in Computational Methods:** The research demonstrates the utility and accuracy of *in silico* methods in drug discovery, which could streamline the drug development process and reduce reliance on *in-vivo* experiments.

**Shortcomings:**

**Lack of Experimental Validation:** While the *in silico* approach is robust, the study lacks experimental validation of pseudoeriocitrin's efficacy and safety, which is crucial for practical application.

**Potential Toxicity Issues:** The absence of hydrogen donors in pseudoeriocitrin's structure, as mentioned in the manuscript, raises concerns about its potential toxicity.

**Limited Understanding of Biological Mechanisms:** The study does not fully explore the biological mechanisms underlying the observed interactions, which is important for understanding the molecule's potential side effects and metabolic pathways.

**Data Reproducibility Concerns:** The manuscript notes structural changes in the molecule during optimization processes. This variability could affect the reproducibility and reliability of the results.

**Absence of Comparative Analysis with Existing Compounds:** The study does not compare pseudoeriocitrin's efficacy and safety profile with that of existing drugs, which is essential to establish its relative advantages or disadvantages.

Overall, while the study presents groundbreaking insights into virtual drug design and molecular docking, it also highlights the need for further research, particularly in experimental validation and toxicity assessment, to fully realize the potential of pseudoeriocitrin as a therapeutic agent.