

Review of: "Expansion of the Experimental Antifungal Activities Through in Silico Docking Study of Compounds From Albizia Lebbeck"

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Potential competing interests: No potential competing interests to declare.

I have thoroughly checked the manuscript entitled, "Expansion of the Experimental Antifungal Activities Through in Silico Docking Study of Compounds From Albizia Lebbeck."

In my opinion, this article could be interesting for publication, but before that, the following points have to be revised.

In subsection 3. Results and Discussion:

- 1- Authors should mention the length of the hydrogen bonds, at least for the best ligand-protein conformations,
- 2- π - π is not a bond; it is a stacking interaction, so authors should modify it accordingly.
- 3- It is better if the authors also analyze π-π and X•••H-π interactions because the ligands have multiple rings .
- 4- As a computational paper, some information about the docking protocol should be included, like the minimization energy, receptor preparation, and glide grid box dimensions,
- 5- Authors should write the full term for each abbreviation, like extra precision (XP).
- 6- The authors used one of the best insilico software (Schrödinger), which provides excellent figures, but they presented only 2D visualization, so they should present 3D visualization of the best conformation.

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