Review of: "Expansion of the antifungal activities through in silico docking study of compounds from Albizia lebbeck fruits"

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Potential competing interests: No potential competing interests to declare.

The manuscript is about the antimicrobial activity of a series of molecules against the Candida albicans target. Docking studies with two crystal structures of sterol 14-alpha demethylase (CYP51) were done to support the experimental results.

In my opinion, the work is interesting. However, the docking protocol is not clear.

1) What are the dimensions and coordinates of the docking box?

2) Was there redocking of the ligands in complex in the crystal to obtain the docking parameters?

3) What is the criterion for choosing the docking poses?

All this information needs to be explained in the Methodology section.

In Table 1, were the other compounds besides 2 and 7 tested? Did they not show activity or, in fact, were they not tested? This information is confusing.