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.

It is not clear in the manuscript whether all the bound substances (ligands and co-factors) and solvent molecules were removed from the protein molecule or not. Is there any importance in choosing the crystal structures of sterol 14-alpha demethylase (CYP51) from Candida albicans in complex with the tetrazole-based antifungal drug candidate and the crystal structure of sterol 14-alpha demethylase (CYP51) from the pathogenic yeast Candida albicans in complex with the antifungal drug posaconazole? Do the tetrazole-based antifungal drug candidate and posaconazole play any significant role in the study?