

## Review Form 1.6

Journal Name:	<a href="#">Journal of Pharmaceutical Research International</a>
Manuscript Number:	Ms_JPRI_82773
Title of the Manuscript:	DESIGN, SYNTHESIS AND MAO INHIBITOR ACTIVITY OF CHROMAN-4- ONE DERIVATIVE
Type of the Article	Original Research Article

### General guideline for Peer Review process:

This journal's peer review policy states that **NO** manuscript should be rejected only on the basis of '**lack of Novelty**', provided the manuscript is scientifically robust and technically sound. To know the complete guideline for Peer Review process, reviewers are requested to visit this link:

(<https://www.journaljpri.com/index.php/JPRI/editorial-policy>)

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### PART 1: Review Comments

	Reviewer's comment	Author's comment (if agreed with reviewer, correct the manuscript and highlight that part in the manuscript. It is mandatory that authors should write his/her feedback here)
<b>Compulsory</b> REVISION comments	<p>The paper is interesting and it could deserve publication but it has serious problems.</p> <ol style="list-style-type: none"><li>1. Nevertheless the English language must be revised by someone knowing it (for example Schiff and not schiff, etc.)</li><li>2. In the Abstract, hMAO is used but not defined.</li><li>3. In point 2.1 please use 1, 2, 3, etc. instead of symbols.</li><li>4. The following points are confusing. Authors must mention what they really calculated (and nothing more). After this they must mention the software used with references and not with a list of what the software does.</li><li>5. Eliminate point 2.2 and simply state that 'the calculations were performed in.. etc.'</li><li>6. Eliminate point 2.3 (see above point 4).</li><li>7. About 2.4.2 Ligand preparation for pharmacophore model development: This point looks like a lecture for beginners. Some points to be clarified:<ul style="list-style-type: none"><li>• Generate Stereoisomers. Why????</li><li>• Remove noncompliant structures. How????</li><li>• Perform an energy minimization. What methodology was used? This is very important.</li></ul></li><li>8. 2.4.3 Generation of the conformers. All the following list:<ul style="list-style-type: none"><li>• No. of steps per rotatable bond-100</li><li>• Pre process minimization steps-100</li><li>• Post process minimization steps-50</li><li>• Force field-OPLS-2005</li><li>• Maximum relative energy difference-10 Kcal/mol</li><li>• RMSD-Cutoff-1Å°</li></ul>Must be placed in only one phrase, such as 'the conditions to generate conformers were: etc.'</li><li>9. Regarding the remaining of the paper, it must be rewritten again. Authors must avoid giving trivial definition of R, SD, etc. because this is a research paper.</li><li>10. Figs. 8 and 9 need to be done with a better software (there are many free scientific programs going better graphs. Do not forget capital letters!)</li></ol>	
<b>Minor</b> REVISION comments		
<b>Optional/General</b> comments		

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**PART 2:**

	<b>Reviewer's comment</b>	<b>Author's comment</b> <i>(if agreed with reviewer, correct the manuscript and highlight that part in the manuscript. It is mandatory that authors should write his/her feedback here)</i>
<b>Are there ethical issues in this manuscript?</b>	<i>(If yes, Kindly please write down the ethical issues here in details)</i>	

**Reviewer Details:**

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