**Reviewer #1:**

1. **The manuscript needs to have a clearer summary of how SWIFTMODELLER 2.0 differs from SWIFTMODELLER.**

The results section has been modified to clearly specify the changes made to the previous version. It also summarizes our work.

1. **The exact license under which the software is available should be given.**

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1. **The manuscript should mention that there is a commercial GUI for MODELLER, distributed by Accelrys.**

The point has been added to the manuscript.

1. **The manuscript should stress that the interface lowers the technical barrier to running MODELLER, not the scientific barrier. Where it describes 'lowering the skill level', it should stress that these are technical skills, not the scientific skills one still needs in order to produce and evaluate a reasonable model.**

The word technical has been added to the introduction of swift modeller. The rest of the paragraph implies that the technical barrier is overcome through the software. The paragraph reads, “It automates the tiresome steps such as formatting of inputs, template protein database searching, writing and execution of python scripts and meaningful data extraction from output files, thus saving an enormous amount of time and human effort for performing homology modeling with MODELLER. In addition it makes pasting of the protein sequence as the only prerequisite, thereby lowering the technical skill level required for using MODELLER”

1. **The authors should confirm the versions of MODELLER with which the interface is compatible.**

SWIFT MODELLER is compatible with all versions that MODELLER provides and thus the reference to MODELLER release notes where the user can see the modeller versions along with the OSs (graphics supported) supported

1. **The manuscript overplays the difficulties of using modeller, for example the words 'extensive' and 'enormous' should be deleted.**

Swift Modeller has been developed keeping the average user in mind with no prior knowledge of using the software and as often is the case, of python too. Thus the software provides a quick solution and can speed up the research by overcoming the barriers of using modeller and scripting in python.

1. **The entire Results section should be deleted.**

The results section’s title has been changed and currently summarizes our work by clearly stating the changes made in the architecture of the previous verion

1. **The figure is not needed and should be deleted.**

The figure (Ramachandran Plot) represents a significant addition from the version 1 of the software and thus has been added

1. **There are a number of sentences that need minor adjustments to English and grammar. However, the errors do not cloud the understanding of the paper and could be picked up at the editorial stage.**

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1. **I am not convinced that Reference 7 is the most appropriate.**

The reference 7 has been removed from the manuscript.

**Reviewer #2:**The Results section has been reorganized to summarize our work and clearly denote the changes made to previous version for the betterment. More snapshots are not added as they were already present in the previous version’s paper. GUI flow wise Ramachandran Plot is the addition and thus its snapshot is made available. Rest of the changes have been made to cater a larger user pool and to improve significantly the previous version in terms of robustness.