

REVIEWER'S REPORT

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3D PERCEPTION OF MAXIMUM DENSITY ZONE ON RAMACHANDRAN PLOTS FOR ZIKA VIRUS PROTEIN STRUCTURES

Recommendation:

Accept as it is
Accept after minor revision (Yes)
 Accept after major revision
 Do not accept (*Reasons below*)

Rating	Excel.	Good	Fair	Poor
Originality		×		
Tech. Quality		×		
Clarity	×			
Significance		×		

Reviewer Name: Dr. Mohamed Ahmed El-Esawi

Date: 12 May 2016

Comments (*Use additional pages, if required*)

Reviewer's Comment / Report

This manuscript discusses Zika virus including the proposed method and implementation. Using the high-fidelity dataset of Zika Virus here, the 3D-plot allows several interesting observations. Most obvious, is the titanic and sharp peak resulting from residues in α -helices. This very sharp peak towers over every other portion of the Ramachandran plot, including the other portions of the classically defined alpha-region. This salient observation suggests that the classically defined alpha-region does not behave as a unit. Moreover it is possible to visualize the maximum density zone in the 3D plot marked with #1 rather than large set of scatter point cluster in the 2D graph. The colour codes being graded gives us a clear picture of how the data in protein data bank files are packed over a narrow region. This work represents a good collection to know more significant information about Zika virus.