SDI FINAL EVALUATION FORM 1.1

PART 1:

Journal Name:	**Journal of Applied Life Sciences International**
Manuscript Number:	**Ms_JALSI_43532**
Title of the Manuscript:	**In-Silico Structural Annotation of Hypothetical Protein Zm00014a_031618 (*Zea mays*)**
Type of Article:	**Original Research Article**

PART 2:

FINAL EVALUATOR'S comments on revised paper (if any)	Authors' response to final evaluator's comments
The authors tried well. However, some important changes/addition are require: As the manuscript is totally computer based study so its presentation and discussion are important for readers of this paper. 1. Write few sentences (3-4 lines) about "Methylthioadenosine Nucleosidase" at the end of introduction section. 2. Remove section 2.4 (line 68-70), this is already covered in section 2.1. **3.** Combine section 2.5 and 2.6 and give a heading that cover both sections. 4. I did't see any proper discussion regarding structure and ligand interaction. I think, if the interaction present through PyMOL software, it may enhance work quality and attract readers.e.g. 5. "Fig. 5. Predicted Binding Residues of Methylthioadenosine Nucleosidase Protein", It should be "Protein ligand interaction of Methylthioadenosine Nucleosidase enzyme". Try to show only active regions of protein in PyMOL (Please see the samples attached file)	

Reviewer Details:

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