Review of: "EOESGC: Predicting miRNA−disease Associations Based on Embedding of Embedding and Simplified Graph Convolutional Network"

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This paper proposed EOESGC, a embedding of embedding and simplified graph convolutional network method for miRNA-disease prediction. However, I have some concerns below.

1. The description of the motivation is not sufficient; authors should introduce the motivation more depth from a methodological point of view. There are a lot of miRNA-disease prediction methods at present, what are their limitations, and what problems we aim to solve.

2. The HMDD database used in the article already has version 3.0, why are the authors still using version 2.0?

3. The experimental results are not fully presented. More baselines based on graph convolutional network should be included. And authors should use more evaluation criterions in "Comparisons with the state−of−the−art methods" section.

4. Authors should carefully check the information of references and ensure the format of references is consistent.