

## Reviewer Report

**Title:** Metaphor - A workflow for streamlined assembly and binning of metagenomes

**Version:** Original Submission    **Date:** 5/15/2023

**Reviewer name:** Po-Yu Liu

### Reviewer Comments to Author:

The Metaphor is a workflow with high completeness for short-read-based metagenomic analysis. I look forward to its compatibility with long-read platforms (ONT and PacBio). This work is worth publishing. However, it is still a bioinformatic knowledge and skill-required toolkit. If the Metaphor can be integrated into a web-based platform, such as Galaxy or Kbase, it would be more user-friendly for much more users.

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