

## Reviewer Report

**Title:** The Gene-Rich Genome of the Scallop *Pecten maximus*

**Version:** Revision 1      **Date:** 3/9/2020

**Reviewer name:** Yi-Jyun Luo

### Reviewer Comments to Author:

The authors addressed all my comments. I have no further comments but a notion from the readme file of AUGUSTUS as follows.

--UTR=on/off

    predict the untranslated regions in addition to the coding sequence. This currently works only for human, *galdieria*, *toxoplasma* and *caenorhabditis*.

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