### **Reviewer Report**

Title: rCASC: reproducible Classification Analysis of Single Cell sequencing data

Version: Revision 1 Date: 4/29/2019

**Reviewer name: Nils Eling** 

#### **Reviewer Comments to Author:**

The authors have now extensively revised the manuscript and added a number of useful functions to the existing rCASC package. By doing so, the authors addressed the majority of my previous comments. However, there remain a few minor comments that should be addressed prior to publication.

- 1) Relating to the previous major comment 1: The publication associated to the griph package can now be cited as: Serra et al., Self-organization and symmetry breaking in intestinal organoid development. Nature (2019).
- 2) Relating to the previous major comment 7: It would be useful to provide more details on which variability measure (variance, squared coefficient of variation or a regression residual) is used to rank genes based on their variability in the topx function while setting type="variance".
- 3) Relating to the previous major comment 10: The legend of Figure 2 in the main manuscript has not been corrected for spelling and phrasing mistakes yet.

#### **Level of Interest**

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# **Quality of Written English**

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