Reviewer’s report

Title: Targeted capture-based NGS is superior to multiplex PCR-based NGS for hereditary BRCA1 and BRCA2 gene analysis in FFPE tumor samples

Version: 1 Date: 30 Jan 2019

Reviewer: Matthias Nees

Reviewer's report:

I carefully checked the work and comments the authors put into this revised document, and how they have addressed the multiple reviewers' recommendations and criticism. There is little to be criticized at this point any more, and I think the manuscript should be accepted.

As promised, the author have also done a Grammar check to the entire document and I could not find major flaws.

Figure 2, which I suggested, reflects the advantages of the method described here in the case of sub-optimal DNA quality; which is a common feature with FFPE samples.

I believe this is ready to go…

Are the methods appropriate and well described?
If not, please specify what is required in your comments to the authors.

Yes

Does the work include the necessary controls?
If not, please specify which controls are required in your comments to the authors.

Yes

Are the conclusions drawn adequately supported by the data shown?
If not, please explain in your comments to the authors.

Yes

Are you able to assess any statistics in the manuscript or would you recommend an additional statistical review?
If an additional statistical review is recommended, please specify what aspects require further assessment in your comments to the editors.

I am able to assess the statistics
Quality of written English
Please indicate the quality of language in the manuscript:

Acceptable

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None

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