Reviewer’s report

Title: A streamlined method for analysing genome-wide DNA methylation patterns from low amounts of FFPE DNA

Version: 1 Date: 02 May 2017

Reviewer: Dajun Deng

Reviewer's report:

Because a regular control method was not used in the optimized protocol for the genome-wide methylation profiling by RRBS using low amount of FFPE DNA samples, the exact advantages and novelty are uncertain. Consider the application potential of the reported protocol, it is needed to point out this pitfall and other disadvantages of their study in the discussion section.

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.

No

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.

Not relevant to this manuscript

Quality of written English
Please indicate the quality of language in the manuscript:

Acceptable
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