BMC Bioinformatics

Erratum to: Integrating mean and variance heterogeneities to identify differentially expressed genes

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Erratum

Upon publication of this article [1], an error occurred in the PDF presentation of the hat (WT) formula in 'Concept of MDE genes and mean heterogeneity tests' of the 'Methods' section on page 3. The corrected formula is shown below:

$$\widehat{WT} = \frac{\hat{\mu}_{i1} - \hat{\mu}_{i2}}{\sqrt{\frac{\hat{\sigma}_{i1}^2}{n_1} + \frac{\hat{\sigma}_{i2}^2}{n_2}}}$$

The above formula has been corrected in the original article.

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Reference

1. Ouyang W, et al. Integrating mean and variance heterogeneities to identify differentially expressed genes. BMC Bioinformatics. 2016;17:497. doi:10.1186/s12859-016-1393-y.

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