Jackson et al. BMC Genomics (2016) 17:1018 DOI 10.1186/s12864-016-3379-6

# **BMC Genomics**

ERRATUM Open Access



# Erratum to: Functional variants of human papillomavirus type 16 demonstrate host genome integration and transcriptional alterations corresponding to their unique cancer epidemiology

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### **Erratum**

In the original publication of this article [1], the funding statement omitted the following information, "Work on the NGS platform of the Institut Curie was supported by the Agence Nationale de la Recherche (ANR Investissement d'Avenir) (ANR-10-EQPX-03) and by the France Génomique National infrastructure (ANR-10-INBS-09)."

The final funding statement can be found below:

## **Funding**

This work was supported by Natural Sciences and Engineering Research Council of Canada (NSERC) grants to IZ (#355858-2008, #435891-2013, #RGPIN-2015-03855), NSERC Alexander Graham Bell Canada Graduate Scholarship-Doctoral (CGS-D) to RJ (#454402-2014), NSERC Alexander Graham Bell Canada Graduate Scholarship-Masters (CGS-M) to SC (#442618-2013), and an NSERC Undergraduate Student Research Award (USRA) to JB (#483630-2015). Work on the NGS platform of the Institut Curie was supported by the Agence Nationale de la Recherche (ANR Investissement d'Avenir) (ANR-10-EQPX-03) and by the France Génomique National infrastructure (ANR-10-INBS-09). The funding bodies had no role in study design, data collection, data analysis and interpretation, or preparation of the manuscript.

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Received: 30 November 2016 Accepted: 5 December 2016 Published online: 12 December 2016

# Reference

 Jackson R, et al. Functional variants of human papillomavirus type 16 demonstrate host genome integration and transcriptional alterations corresponding to their unique cancer epidemiology. BMC Genomics. 2016; 17:851. doi:10.1186/s12864-016-3203-3.

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