

# Response to: "Response to *Different measures of 'genome-wide' DNA methylation exhibit unique properties in placental and somatic tissues*"

E. Magda Price<sup>1,2</sup> and Wendy P. Robinson<sup>2,\*</sup>

<sup>1</sup>Department of Obstetrics and Gynaecology; University of British Columbia; Vancouver, BC Canada;

<sup>2</sup>Department of Medical Genetics; University of British Columbia; Vancouver, BC Canada

We are responding to a Letter to the Editor addressing the Method section of our paper "Different measures of 'genome-wide' DNA methylation exhibit unique properties in placental and somatic tissues" (Price ME, Cotton AM, Peñaherrera MS, McFadden DE, Kobor MS, Robinson WP. Different measures of "genome-wide" DNA methylation exhibit unique properties in placental and somatic tissues. *Epigenetics* 2012; 7:652-63; PMID: 22531475; <http://dx.doi.org/10.4161/epi.20221>). The letter raised concerns that the protocol for Epigentek's MethylFlash™ kit was followed incorrectly based on the wording of an online publication of our article. We admittedly made an error in the language used to describe the MethylFlash™ protocol in our initial submission and thus this was corrected as soon as it was brought to our attention. However, the error was only in language and not procedure. We are confident that the protocol was followed as stated in the insert provided with the MethylFlash™ Methylated DNA Quantification kit (Colorimetric).

Do not distribute.

\*Correspondence to: Wendy P. Robinson; Email: [wrobinson@cfri.ca](mailto:wrobinson@cfri.ca)

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Comment on: Li W. Response to: "Different measures of 'genome-wide' DNA methylation exhibit unique properties in placental and somatic tissues." *Epigenetics* 2012; 7:796; PMID: 22772086; <http://dx.doi.org/10.4161/epi.21099>