

SCIENTIFIC REPORTS

OPEN

Corrigendum: Motivational, proteostatic and transcriptional deficits precede synapse loss, gliosis and neurodegeneration in the B6.*Htt*^{Q111/+} model of Huntington's disease

Robert M. Bragg, Sydney R. Coffey, Rory M. Weston, Seth A. Ament, Jeffrey P. Cante, Shawn Minnig, Cory C. Funk, Dominic D. Shuttleworth, Emily L. Woods, Bonnie R. Sullivan, Lindsey Jones, Anne Glickenhau, John S. Anderson, Michael D. Anderson, Stephen B. Dunnett, Vanessa C. Wheeler, Marcy E. MacDonald, Simon P. Brooks, Nathan D. Price & Jeffrey B. Carroll

Scientific Reports 7:41570; doi: 10.1038/srep41570; published online 08 February 2017; updated on 28 March 2017

This Article contains a typographical error in the Methods section, under the subheading “Library construction, RNA Sequencing and RNASeq analysis”, where:

“The fastq files were aligned using the default parameters of SNAPR⁵⁰ (<https://github.com/PriceLab/snapr>) against the GRCh38 genome assembly, along with the transcriptome assembly gtf file from Ensembl, GRCh38.75”.

should read:

“The fastq files were aligned using the default parameters of SNAPR⁵⁰ (<https://github.com/PriceLab/snapr>) against the GRCm38 genome assembly, along with the transcriptome assembly gtf file from Ensembl, GRCm38”.



This work is licensed under a Creative Commons Attribution 4.0 International License. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in the credit line; if the material is not included under the Creative Commons license, users will need to obtain permission from the license holder to reproduce the material. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>

© The Author(s) 2017