Corrigendum

Genome Research 28: 396–411 (2018)

Corrigendum: SQANTI: extensive characterization of long-read transcript sequences for quality control in full-length transcriptome identification and quantification

Manuel Tardaguila, Lorena de la Fuente, Cristina Marti, Cécile Pereira, Francisco Jose Pardo-Palacios, Hector del Risco, Marc Ferrell, Maravillas Mellado, Marissa Macchietto, Kenneth Verheggen, Mariola Edelmann, Iakes Ezkurdia, Jesus Vazquez, Michael Tress, Ali Mortazavi, Lennart Martens, Susana Rodriguez-Navarro, Victoria Moreno-Manzano, and Ana Conesa

The authors would like to correct the omission of a funding source in the Acknowledgments section which was inadvertently excluded from the initial publication of this article. The additional text is as follows and has been updated online:

“This project has received funding from the Marie Curie International Research Staff Exchange Scheme within the 7th European Community Framework Program under grant agreement No 612583-DEANN.”

The authors apologize for the inconvenience.

doi: 10.1101/gr.239137.118

Corrigendum: SQANTI: extensive characterization of long-read transcript sequences for quality control in full-length transcriptome identification and quantification

Manuel Tardaguila, Lorena de la Fuente, Cristina Marti, et al.

Genome Res. 2018 28: 1096

Access the most recent version at doi:[10.1101/gr.239137.118](https://doi.org/10.1101/gr.239137.118)

Related Content **SQANTI: extensive characterization of long-read transcript sequences for quality control in full-length transcriptome identification and quantification**
Manuel Tardaguila, Lorena de la Fuente, Cristina Marti, et al.
[Genome Res. March , 2018 28: 396-411](#)

Open Access Freely available online through the *Genome Research* Open Access option.

License

Email Alerting Service Receive free email alerts when new articles cite this article - sign up in the box at the top right corner of the article or [click here](#).

**Affordable, Accurate
Sequencing.**

To subscribe to *Genome Research* go to:
<https://genome.cshlp.org/subscriptions>