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Corrigendum to: Poisson-Tweedie mixed-effects model: a flexible approach for the analysis of longitudinal RNA-seq data (Aug, 10.1177/1471082X20936017, 2020)

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Corrigendum to Poisson–Tweedie mixed-effects model: A flexible approach for the analysis of longitudinal RNA-seq data

“Poisson–Tweedie mixed-effects model: A flexible approach for the analysis of longitudinal RNA-seq data” by Mirko Signorelli, Pietro Spitali and Roula Tsonaka was published in *Statistical Modelling*, Onlinefirst 24 August 2020, DOI: 10.1177/1471082X20936017.

The authors have recently identified two mistakes in the R code that they used to estimate the Poisson-Tweedie mixed model (ptmixed) in simulations C and D, whose results are presented in Section 3.3 of the OnlineFirst version of the article. Therefore, they have proceeded to rerun such simulations with the corrected code, and to update the results of Section 3.3 accordingly.

The amended results of simulations C and D will be published in the onlinefirst version of the article and the subsequent issue in which it is published.

