Correction: The genomic and clinical landscape of fetal akinesia

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Genetics in Medicine (2020) 22:1426; <https://doi.org/10.1038/s41436-020-0839-9>

Correction to: *Genetics in Medicine* 22:2020; <https://doi.org/10.1038/s41436-019-0680-1>; published online 04 November 2019

In the original version of this Article there was a nomenclature error in Table 2 and Supplementary Table S1. For patient 30 it should read c.[2863C>T];[3082+1G>C] and not c.[2863C>T];[3082+1C>G]. This has now been corrected in both the PDF and HTML versions of the Article as well as the supplementary information file.

Published online: 26 May 2020

Correction: An approach to integrating exome sequencing for fetal structural anomalies into clinical practice

Neeta L. Vora, MD, Kelly Gilmore, MS, CGC, Alicia Brandt, MS, Chelsea Gustafson, BS, Natasha Strande, PhD, Lori Ramkissoo, PhD, Emily Hardisty, MS, CGC, Ann Katherine M. Foreman, MS, CGC, Kirk Wilhelmsen, MD, PhD, Phillips Owen, BS, Karen E. Weck, MD, Jonathan S. Berg, MD, PhD, Cynthia M. Powell, MD, MS and Bradford C. Powell, MD, PhD

Genetics in Medicine (2020) 22:1426; <https://doi.org/10.1038/s41436-020-0870-x>

Correction to: *Genetics in Medicine* 22:2020; <https://doi.org/10.1038/s41436-020-0750-4>, published online 24 January 2020

The original version of this Article contained an error in the spelling of the TP63 gene “NM_003722.4:c.1028G>C.; [p.Arg343Pro]” in Table 2, which was incorrectly given as “1028G>A”. This has now been corrected in both the PDF and HTML versions of the Article.

Published online: 18 June 2020

Correction: Phenotate: crowdsourcing phenotype annotations as exercises in undergraduate classes

Willie H. Chang, MSc, Pouria Mashouri, MSc, Alexander X. Lozano, MSc, Brittney Johnstone, MSc, CGC, Mia Husić, MSc, Annie Olry, PhD, Sylvie Maiella, PhD, Tugce B. Balci, MD, Sarah L. Sawyer, PhD, MD, Peter N. Robinson, MD, MSc, Ana Rath, MD and Michael Brudno, PhD

Genetics in Medicine (2020) 22:1427; <https://doi.org/10.1038/s41436-020-0866-6>

Correction to: *Genetics in Medicine* 2020; <https://doi.org/10.1038/s41436-020-0812-7>, published online 05 May 2020

The original version of this Article contained an incorrect supplementary file in the Excel file format. This has now been replaced in the HTML version of the Article.

Published online: 18 June 2020

Open

Correction: Alternative mRNA splicing can attenuate the pathogenicity of presumed loss-of-function variants in *BRCA2*

Romy L. S. Mesman, MSc, Fabienne M. G. R. Calléja, MSc, Miguel de la Hoya, PhD, Peter Devilee, PhD, Christi J. van Asperen, MD, PhD, Harry Vrieling, PhD and Maaïke P. G. Vreeswijk, PhD

Genetics in Medicine (2020) 22:1427–1428; <https://doi.org/10.1038/s41436-020-0883-5>

Correction to: *Genetics in Medicine* 2020; <https://doi.org/10.1038/s41436-020-0814-5>, published online 13 May 2020

The original version of this Article omitted an essential reference (“Meulemans L, Mesman RLS, Caputo SM, et al. Skipping nonsense to maintain function: the paradigm of *BRCA2* exon 12. *Cancer Res.* 2020;80(7):1374–1386. DOI: 10.1158/0008-5472.CAN-19-2491”). This has now been added in both the PDF and HTML versions of the Article.

In addition, the original version of this Article contained a typo in the sentence “However, two variants in respectively the splice acceptor site (c.517–2A>G) and donor site (c.631+2T>G) of exon 7 were classified as pathogenic based...” that was incorrectly given as “However, two variants in respectively the splice donor site (c.517–2A>G) and acceptor site (c.631+2T>G) of exon 7 were classified as pathogenic based...”. Where “donor” should be “acceptor” and vice versa. This has now been corrected in both the PDF and HTML versions of the Article.

Finally, the original version of this Article contained an incorrect supplementary file. This has now been replaced in the HTML version of the Article.

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