Erratum

Increased yields of duplex sequencing data by a series of quality control tools

Gundula Povysil^{1,*,†}, Monika Heinzl^{1,†}, Renato Salazar¹, Nicholas Stoler², Anton Nekrutenko² and Irene Tiemann-Boege^{1,*}

¹Institute of Biophysics, Johannes Kepler University, 4020 Linz, Austria and ²Graduate Program in Bioinformatics and Genomics, The Huck Institutes for Life Sciences, The Pennsylvania State University, University Park, PA 16802, USA

In the original published version of the article 'chr17:' was not used consistently throughout the text. All instances of 'chr17-' or 'chr17-g-' have now been replaced by 'chr17:'

The colours in Table 5 were removed. Line 1 (starting with chr17: 7669456-C-A) of Table 5 should have been coloured in blue and line 9 (starting with chr17:7676325-C-Del) should have been coloured in green.

Corrections have been made to the published article.

^{*}To whom correspondence should be addressed. Irene Tiemann-Boege. Tel: +43 732 2468 7620; Fax: +43 732 2468 27620; Email: irene.tiemann@jku.at Correspondence may also be addressed to Gundula Povysil. Tel: +1 212 305 0754; Email: povysil@bioinf.jku.at †The authors wish to be known that, in their opinion, the first two authors should be regarded as Joint First Authors. Present address: Gundula Povysil, Institute for Genomic Medicine, Columbia University Irving Medical Center, New York, NY 10032, USA.