



Correction for Volume 14, p. 414

Genome Res. 2004 14: 786

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A horizontal banner advertisement with a teal background. On the left, the text reads 'CRISPR and RNAi Genetic Screening. Your new superpower.' in white. In the center, there is a white-bordered box containing the words 'LEARN MORE' in black. On the right, there is a photograph of a woman wearing a red superhero mask and a red cape over a white top. To the right of the photo is the Cellecta logo, a green molecular structure, and the word 'CELLECTA' in white capital letters.

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Errata

Genome Research 13: 103–107 (2003)

Human–Mouse Alignments With BLASTZ

Scott Schwartz, W. James Kent, Arian Smit, Zheng Zhang, Robert Baertsch, Ross C. Hardison, David Haussler, and Webb Miller

In the second column on page 103, the Web address for axt-Best-processed BLASTZ alignments on the human and mouse genomes contains a typo. The correct Web address is:

<http://genome.cse.ucsc.edu/goldenPath/28jun2002/vsMm2>

The authors apologize for any confusion this may have caused.

Genome Research 14: 414–425 (2004)

Parallel Genotyping of Over 10,000 SNPs Using a One-Primer Assay on a High-Density Oligonucleotide Array

Hajime Matsuzaki, Halina Loi, Shoulian Dong, Ya-Yu Tsai, Joy Fang, Jane Law, Xiaojun Di, Wei-Min Liu, Geoffrey Yang, Guoying Liu, Jing Huang, Giulia C. Kennedy, Thomas B. Ryder, Gregory A. Marcus, P. Sean Walsh, Mark D. Shriver, Jennifer M. Puck, Keith W. Jones, and Rui Mei

The Abstract for this article contained a typo. The sentence should have read:

“The selection of SNPs was primarily determined by computer-predicted lengths of restriction fragments containing the SNPs, and was further driven by strict empirical measurements of accuracy, reproducibility, and average call rate, which we estimate to be >99.5%, >99.9%, and >95%, respectively.”

The editors apologize for any confusion this may have caused.