Correction: Inferring Loss-of-Heterozygosity from Unpaired Tumors Using High-Density Oligonucleotide SNP Arrays

Rameen Beroukhim, Ming Lin, Yuhyun Park, Ke Hao, Xiaojun Zhao, Levi A. Garraway, Edward A. Fox, Ephraim P. Hochberg, Ingo K. Mellinghoff, Matthias D. Hofer, Aurelien Descazeaud, Mark A. Rubin, Matthew Meyerson, Wing Hung Wong, William R. Sellers, Cheng Li

In *PLoS Computational Biology,* volume 2, issue 5:

In the subsection "Initial probabilities" of the Results section, the first sentence was incorrect and should say:

These probabilities, denoted by $P_0(\text{RET})$ and $P_0(\text{LOSS}) = 1 - P_0(\text{RET})$, specify the probabilities of RET and LOSS for the p-terminal marker on a chromosome.

The second line of Equation 1 was incorrect. The correct equation is:

$$P(\text{LOSS}|\text{LOSS}) = \theta \cdot P_0(\text{LOSS}) + (1 - \theta)$$
$$\text{and } P(\text{LOSS}|\text{RET}) = \theta \cdot P_0(\text{LOSS})$$

Correction: The Genome: An Outsider's View

Carl Zimmer

In *PLoS Computational Biology,* vol 2, issue 12:

In the Acknowledgments section of this Perspective, the spelling of the name of Pavel Pevzner was incorrect.