

Corrections

CELL BIOLOGY

Correction for “Core epithelial-to-mesenchymal transition interactome gene-expression signature is associated with claudin-low and metaplastic breast cancer subtypes,” by Joseph H. Taube, Jason I. Herschkowitz, Kakajan Komurov, Alicia Y. Zhou, Supriya Gupta, Jing Yang, Kimberly Hartwell, Tamer T. Onder, Piyush B. Gupta, Kurt W. Evans, Brett G. Hollier, Prahlad T. Ram, Eric S. Lander, Jeffrey M. Rosen, Robert A. Weinberg, and Sendurai A. Mani, which appeared in issue 35, August 31, 2010 of *Proc Natl Acad Sci USA* (107:15449–15454; first published August 16, 2010; 10.1073/pnas.1004900107).

The authors note that on page 15453, right column, fifth full paragraph, sentence 2, “Microarray data for HMLE Gsc, Snail, Twist, TGF- β 1, and vector control has been deposited in GEO under accession number GSE9691” should instead appear as “Microarray data for HMLE Gsc, Snail, Twist, TGF- β 1, and vector control has been deposited in GEO under accession number GSE24202.”

www.pnas.org/cgi/doi/10.1073/pnas.1015095107

MEDICAL SCIENCES

Correction for “Detection of MLV-related virus gene sequences in blood of patients with chronic fatigue syndrome and healthy blood donors,” by Shyh-Ching Lo, Natalia Pripuzova, Bingjie Li, Anthony L. Komaroff, Guo-Chiuan Hung, Richard Wang, and Harvey J. Alter, which appeared in issue 36, September 7, 2010, of *Proc Natl Acad Sci USA* (107:15874–15879; first published August 23, 2010; 10.1073/pnas.1006901107).

The authors note that the GenBank Accession Numbers for the *gag* gene are HM630557–HM630562, and GenBank Accession Numbers for the *env* gene are HQ157342–HQ157343.

www.pnas.org/cgi/doi/10.1073/pnas.1015107107