

Erratum: “Role of Tumor Pericytes in the Recruitment of Myeloid-Derived Suppressor Cells” by JongWook Hong, Nicholas P. Tobin, Helene Rundqvist, Tian Li, Marion Lavergne, Yaiza Garcia-Ibanez, Hanyu Qin, Janna Paulsson, Manuel Zeitelhofer, Milena Z. Adzemovic, Ingrid Nilsson, Pernilla Roswall, Johan Hartman, Randall S. Johnson, Arne Ostman, Jonas Bergh, Mirjana Poljakovic, and Guillem Genove. *J Natl Cancer Inst* 2015; 107(10): djv209 doi:10.1093/jnci/djv209. In the Methods section under the subheading Patients and Specimens, there is an inaccurate statement. The corrected paragraph is as follows; the corrected text is in bold.

Our study population consisted of the Uppsala breast cancer cohort (253 patients), previously described in detail (18,19). The ethics committee at Karolinska Institute approved the gene expression study (Stockholm, Sweden). **The present gene expression data were derived from previously published patient cohorts, with approvals from the ethical committee at Karolinska University hospital and the committee at Karolinska Institutet. The tissue micro arrays were derived from biobank samples, with individual signed informed consents for biobanking.** The distribution of tumor stage at diagnosis was as follows: Stage 1, 41.0%; Stage 2, 24.6%; Stage 3, 34.4%; and mean age at diagnosis was 62.13 years (range = 28–93 years). Raw data files are publically available (NCBI Geo GSE3494).

The authors regret the error.

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Erratum: “Enhancing Breast Cancer Recurrence Algorithms Through Selective Use of Medical Record Data” by Candyce H. Kroenke, Jessica Chubak, Lisa Johnson, Adrienne Castillo, Erin Weltzien, and Bette J. Caan. *J Natl Cancer Inst* 2016; 108(3): djv336 doi:10.1093/jnci/djv336. There were minor errors in Table 1 that have been corrected and republished in the March 2016 online issue. The errors did not appear in the print issue. The authors regret the errors.

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