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AUTHOR CORRECTION OPEN Author Correction: Breast-cancer-specific mortality in patients treated based on the 21-gene assay: a SEER population-based study

Valentina I Petkov¹, Dave P Miller², Nadia Howlader¹, Nathan Gliner², Will Howe³, Nicola Schussler³, Kathleen Cronin¹, Frederick L Baehner^{2,4}, Rosemary Cress⁵, Dennis Deapen⁶, Sally L Glaser^{7,8}, Brenda Y Hernandez⁹, Charles F Lynch¹⁰, Lloyd Mueller¹¹, Ann G Schwartz¹², Stephen M Schwartz¹³, Antoinette Stroup^{14,15}, Carol Sweeney¹⁶, Thomas C Tucker¹⁷, Kevin C Ward¹⁸, Charles Wiggins¹⁹, Xiao-Cheng Wu²⁰, Lynne Penberthy¹ and Steven Shak²

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In the original version of the published article, line three of the third paragraph of the methods stated "Excluding patients with micrometastatic disease, the 5-year BCSM for patients with Recurrence Score results <18 and 1–3 positive nodes (n = 2,617) was 1.3% (95% CI, 0.6–2.9%)." To improve clarity this statement has been replaced with "Excluding patients with micrometastatic disease, there were 2,617 patients with 1–3 positive nodes. Of these, 1,487 also had Recurrence Score results <18 with 5-year BCSM of 1.3% (95% CI, 0.6–2.9%)." The original version of the published article also contained an error in the second sentence of the Figure 2 legend describing the mutation status of the patient population examined. The sentence in the original published version of the Article stated "Patients with HR+, HER2-positive, node-negative..." this has been changed to "Patients with HR+,

HER2-negative, node-negative...". This has been corrected in the PDF and HTML versions of this paper.

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