Correction to:

*Impact of Specimen Heterogeneity on Biomarkers in Repository Samples from Patients with Acute Myeloid Leukemia: A SWOG Report,*

by Pogosova-Agadjanyan, EL, et al.


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In the February 2018 issue of Biopreservation and Biobanking (vol. 16, no. 1, pp. 42–52), in the article titled *Impact of Specimen Heterogeneity on Biomarkers in Repository Samples from Patients with Acute Myeloid Leukemia: A SWOG Report* by Era L. Pogosova-Agadjanyan et al., the authors identified several errors in the published article.

One of the errors was in Figure 2, on page 46, in which the images in the A and B panels were switched.

There was also a minor error on page 46, first column, dealing with the $p$ values for the BM and PB, although the interpretation remains the same. The following was corrected under the *Results* section: (BM: 95% CI 1.2–2.6, $p < 0.001$, PB: 95% CI 0.4–2.6, $p = 0.009$; Fig. 2).

The other error stemmed from a shift in columns in Supplementary Table S11 (the revised Suppl Table S11 has been reuploaded), which impacted a paragraph in the first column on Page 47, where the following changes were made:

*BAALC, CEBPA, CD34, ERG1, EVI1, KIT,* and *MN1* were significantly associated with normal cytogenetics in MNCs and AML blasts. Half of the biomarkers were significantly associated with normal cytogenetics in both unsorted MNCs and AML blasts, with *RUNX1 and JAG1* being significantly associated with normal cytogenetics in MNCs. *CCNA, ERG1, GATA2, IL3RA, RUNXI,* and *WT1* were positively correlated with PB blast percentages in unsorted MNCs and AML blasts. *EVI1, FLT3,* and *KIT* in unsorted blasts were significantly correlated with PB blast percentage. Similarly, *BAALC, CCNA1, CD34, ERG1, IL3RA, KIT,* and *MN1* were significantly correlated with WBC in MNCs and blasts, whereas *EVI1* was significantly correlated with WBC in unsorted MNCs only (Supplementary Table S11).

The online version has been corrected.

The authors apologize for these errors.