

CORRECTION OPEN



Correction: Integrative analysis identifies an older female-linked AML patient group with better risk in ECOG-ACRIN Cancer Research Group's clinical trial E3999

Franck Rapaport, Kenneth Seier, Yaseswini Neelamraju, Duane Hassane, Timour Baslan, Daniel T. Gildea, Samuel Haddox, Tak Lee, H. Moses Murdock , Caroline Sheridan, Alexis Thurmond, Ling Wang, Martin Carroll, Larry D. Cripe, Hugo Fernandez, Christopher E. Mason, Elisabeth Paietta, Gail J. Roboz, Zhuoxin Sun, Martin S. Tallman, Yanming Zhang, Mithat Gönen, Ross Levine , Ari M. Melnick , Maria Kleppe and Francine E. Garrett-Bakelman

© The Author(s) 2023

Blood Cancer Journal (2023)13:103; <https://doi.org/10.1038/s41408-023-00862-2>

Correction to: *Blood Cancer Journal* <https://doi.org/10.1038/s41408-022-00736-z>, published online 23 September 2022

Following the publication of this article, the authors noted an error in sample reporting. Eleven specimens included in the original data were follow up specimens taken from patients after initial therapy, and not diagnostic specimens sampled prior to initial therapy.

To correct for this error, and to accurately report on findings in pre-therapy samples which have prognostic value, data from the 11 samples have now been excluded and the data re-analyzed.

Corrections and unchanged results after reanalysis are detailed as follows:

1. The overall number of genes with recurrent oncogenic and likely oncogenic mutations in the study cohort was unchanged (Fig. 1A, B, Supplementary Fig. 2).
2. The comparison to BEAT AML data and the findings of a subset of mutations being enriched in the older AML patients remains unchanged (Fig. 1C, D).
3. Cytogenetics data was not originally available for the specimens excluded, thus these analyses and published results are unchanged.
4. Conclusions from the somatic event co-occurrence analyses were unchanged (Fig. 1E, Supplementary Fig. 5).
5. Due to the smaller sample size, the threshold for inclusion in the regression tree analysis for association with overall survival was increased to keep the balance between false

positive and false negative findings. After re-analysis, sex remains a classifying parameter for overall survival and the somatic events that contribute to the final model are unchanged (Fig. 2A, B).

6. Due to the smaller sample size, the assessment for higher frequency of achieving complete remission between G2 and G3 patient groups was changed from a fisher exact test to a chi-square test. With sufficient numbers in the 2 x 2 table (more than five in each cell) a chi-square test provides Type I control at the nominal level while supplying more power. We found a significant difference in achievement of complete remission between the groups as reported originally; the two groups also had a difference in overall survival, however, this difference was no longer significant (Supplementary Fig. 10; text lines 118–120).
7. The finding that the novel risk group reclassified most patients in the group from poor or intermediate ELN2017 to better risk was unchanged (Fig. 2C).
8. The findings when comparing the patterns of mutations in G2 and G3 risk groups were unchanged (Fig. 2D and Supplementary Fig. 11).
9. The finding of a better and worse risk group in the older AML patients remains unchanged (Fig. 2E).
10. Our findings regarding sex being a classifying parameter for achievement of complete remission in the study cohort remains unchanged (Supplementary Fig. 15).

To reflect the change in specimen count included, all figure panels listed above and associated text in the main paper, Supplementary Information and Supplementary Tables have been updated.

Fig. 1

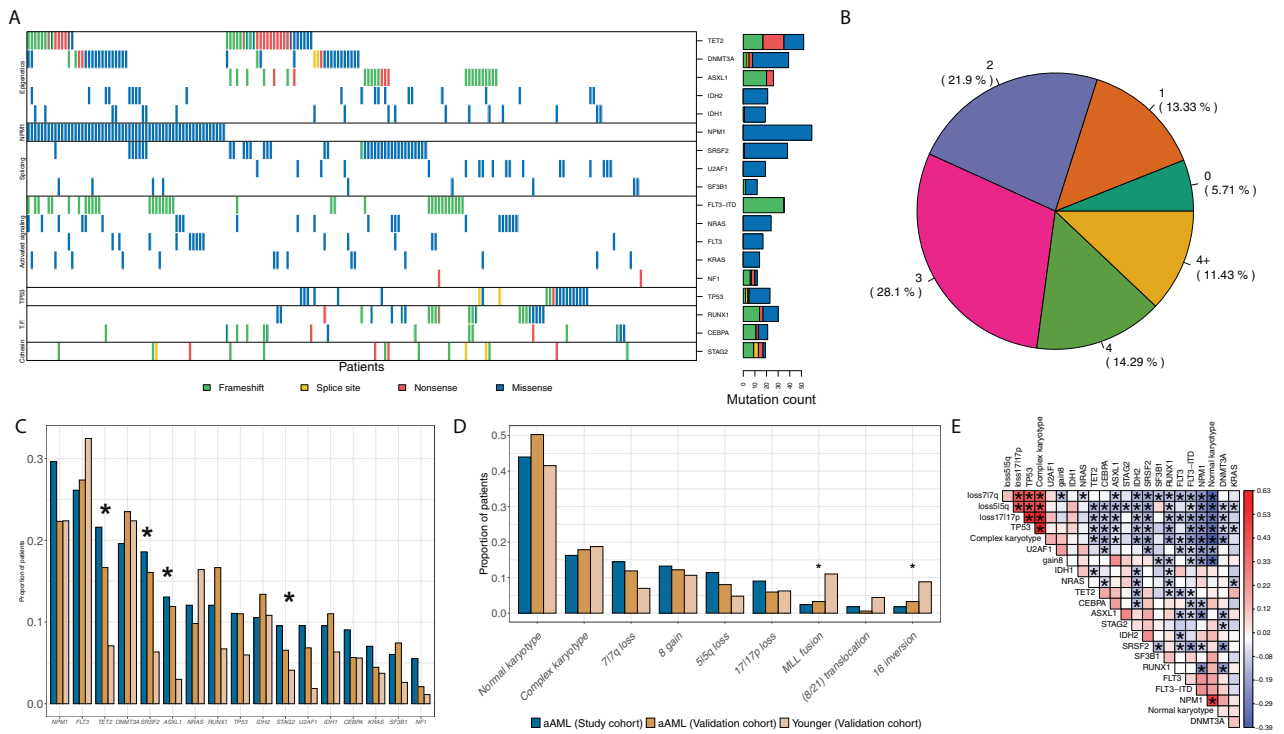
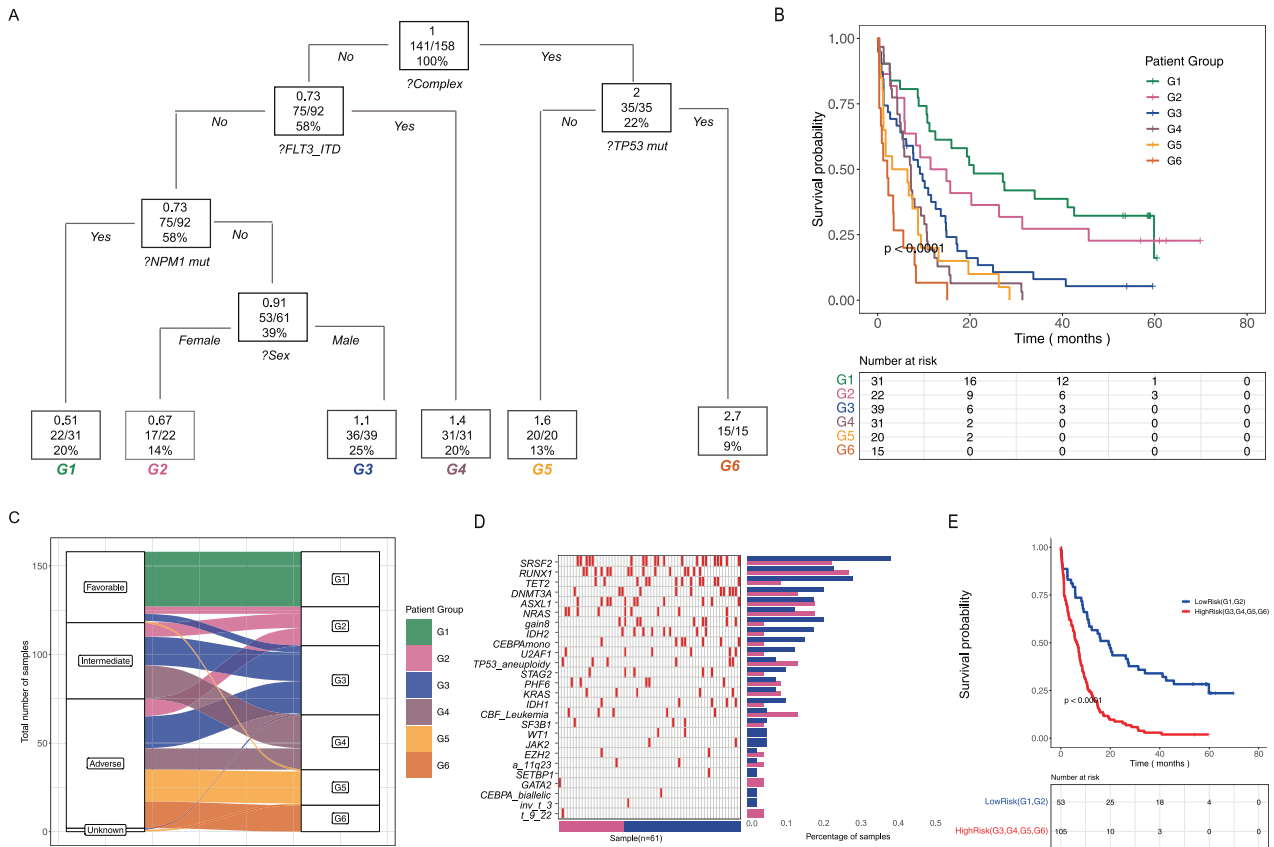


Fig. 2



The original article has been corrected.



Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

© The Author(s) 2023