

CORRECTION

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Publisher Correction: Frugal alignment-free identification of FLT3-internal tandem duplications with FiLT3r

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The original article can be found online at <https://doi.org/10.1186/s12859-022-04983-6>.

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Following publication of the original article [1], the authors identified that the files in Additional file 1 were not separately listed in additional files. This has been corrected since then.

The original article [1] has been corrected.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12859-022-05079-x>.

Additional file 1. supp file 1: filt3r-suppl.pdf.

Additional file 2. supp file 2: results_sra.xlsx.

Additional file 3. supp file 3: accessions.csv.

Additional file 4. supp file 4: results-different-k.xlsx.

Additional file 5. supp file 5: results-small-k.xlsx.

Additional file 6. supp file 6: results-different-perc.xlsx.

Additional file 7. supp file 7: results-subsamples.xlsx.

Additional file 8. supp file 8: genescan_SRR15006459.pdf.

Additional file 9. supp file 9: genescan_SRR15006372.pdf.

Additional file 10. supp file 10: accessions-CCLE.csv.

Additional file 11. supp file 11: results_simulated_150bp.xlsx.

Additional file 12. supp file 12: results_simulated_250bp_hq.xlsx.

Additional file 13. supp file 13: results_simulated_250bp_lc.xlsx.

Additional file 14. supp file 14: results_simulated_250bp_lq.xlsx.

Additional file 15. supp file 15: results_simulated_250bp.xls.

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Reference

1. Boudry A, et al. Frugal alignment-free identification of FLT3-internal tandem duplications with FILT3r. *BMC Bioinform.* 2022;23:448. <https://doi.org/10.1186/s12859-022-04983-6>.

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