

## Corrigendum

# Multi-omic and multi-view clustering algorithms: review and cancer benchmark

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Our paper included a benchmark comparing nine methods for multi-omics clustering, including PINS (1). We reported that PINS crashed when run on a breast cancer dataset. We performed all the benchmarks on a 64-bit computer, using the 32-bit version of R. In later tests we observed that PINS did not crash on 64-bit R, and it only crashed on 32-bit R due to insufficient memory. The clustering that PINS obtained on the breast cancer dataset had 4 enriched clinical parameters, and the *P*-value for the logrank test on that clustering was 0.05.

### REFERENCES

1. Nguyen,T., Tagett,R., Diaz,D. and Draghici,S. (2017) A novel approach for data integration and disease subtyping. *Genome Res*, **27**, 2025–2039.

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