

ERRATUM

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# Erratum to: Expression and methylation patterns partition luminal-A breast tumors into distinct prognostic subgroups

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## Erratum

After publication of this article [1], the authors requested to add an official acknowledgement for using the TCGA Breast Cancer dataset in their article. The following information has therefore been added to the Acknowledgements section: "The results published here are based upon data generated by The Cancer Genome Atlas managed by the NCI and NHGRI. Information about TCGA can be found at <http://cancergenome.nih.gov>". The amended Acknowledgements section has been included in this erratum accordingly.

## Acknowledgements

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## References

1. Netanelly D, et al. Expression and methylation patterns partition luminal-A breast tumors into distinct prognostic subgroups. *Breast Cancer Res.* 2016; 18:74. doi:10.1186/s13058-016-0724-2.

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