



Erratum: Identification of Breast Cancer Stem Cell Related Genes Using Functional Cellular Assays Combined With Single-Cell RNA Sequencing in MDA-MB-231 Cells

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An Erratum on

Identification of Breast Cancer Stem Cell Related Genes Using Functional Cellular Assays Combined With Single-Cell RNA Sequencing in MDA-MB-231 Cells

Jonasson E, Ghannoum S, Persson E, Karlsson J, Kroneis T, Larsson E, Landberg G and Ståhlberg A (2019). *Front. Genet.* 10:500. doi: 10.3389/fgene.2019.00500

The Data Availability Statement was missing in the original article. A correction has therefore been made, and the statement should read:

“RNA sequencing data have been deposited in NCBI’s Gene Expression Omnibus (Edgar et al., 2002) database with accession number GSE124989.”

The original version of this article has been updated.

REFERENCE

Edgar, R., Domrachev, M., and Lash, A. E. (2002). Gene Expression Omnibus: NCBI gene expression and hybridization array data repository. *Nucleic Acids Res.* 30, 207–210. doi: 10.1093/nar/30.1.207

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