

Correction: Novel Hybrid Phenotype Revealed in Small Cell Lung Cancer by a Transcription Factor Network Model That Can Explain Tumor Heterogeneity

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In the original version of this article (1), Fig. 1 was incorrectly replaced with Fig. 2. The error has been corrected in the latest online HTML and PDF versions of the article. The publisher regrets this error.

Reference

1. Udyavar AR, Wooten DJ, Hoeksema M, Bansal M, Califano A, Estrada L, et al. Novel hybrid phenotype revealed in small cell lung cancer by a transcription factor network model that can explain tumor heterogeneity. *Cancer Res* 2017;77:1063–74.

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