CORRECTION

Open Access

Correction: Synergistic anti-AML effects of the LSD1 inhibitor T-3775440 and the NEDD8-activating enzyme inhibitor pevonedistat via transdifferentiation and DNA rereplication

Y. Ishikawa¹, K. Nakayama¹, M. Morimoto¹, A. Mizutani¹, A. Nakayama¹, K. Toyoshima¹, A. Hayashi¹, S. Takagi¹, R. Dairiki¹, H. Miyashita¹, S. Matsumoto², K. Gamo¹, T. Nomura¹ and K. Nakamura¹

Correction to: *Oncogenesis*

<https://doi.org/10.1038/oncsis.2017.76>

Published online 11 September 2017

The FAB classification of OCI-M2, GF-D8, CMK-11-5, and THP-1 cell lines in Table 1 was wrong. This error has been fixed in the revised table.

Published online: 11 September 2019

Correspondence: K. Nakamura (kazuhide.nakamura@takeda.com)

¹Oncology Drug Discovery Unit, Pharmaceutical Research Division, Takeda Pharmaceutical Company Limited, Fujisawa, Japan

²Integrated Technology Research Laboratories, Pharmaceutical Research Division, Takeda Pharmaceutical Company Limited, Fujisawa, Japan

These authors contributed equally: Y. Ishikawa, K. Nakayama

© The Author(s) 2019

Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit <http://creativecommons.org/licenses/by/4.0/>.

Table 1 Synergy score of T-3775440/pevonedistat combination in an AML cell panel

Cell Line	Meaning	Blending Synergy	Combination Index	Incubation (h)	FAB
TF-1a	Synergy	36.5	0.3	72	M6
TF-1a/Ara-C	Synergy	34.9	NA	72	M6
NB4	Synergy	33.5	NA	120	M3
Kasumi-1	Synergy	31.8	0.45	120	M2
MOLM16	Synergy	28.4	NA	120	M7
HL-60/MX2	Synergy	23.3	0.42	120	M2
HL-60	Synergy	19.6	0.41	120	M2
HEL92.1.7	Additivity	14.6	NA	72	M6
OCI-M2	Additivity	10.9	0.81	72	M6
GF-D8	Additivity	8.6	NA	168	M1
CMK11-5	Additivity	4.3	0.96	72	M7
THP-1	Additivity	-5.5	NA	144	M5
OCI-AML3	Additivity	-7.9	NA	168	M4
EOL-1	Additivity	-17.3	NA	120	Eosinophilic
CMK-86	Subadditivity	-27.6	1.73	72	M7

NOTE: Combination index (CI) values in the range 0–0.7 and 0.7–1.3 are classified as synergy and additivity, respectively. When CI values were not associated, nonlinear blending values greater than 20 and between –20 and +20 were classified as synergy and additivity, respectively. Heat maps are color-coded based on the combination effects: Green, synergy; orange, additive; red, subadditive.