



# Corrigendum: Denatonium as a Bitter Taste Receptor Agonist Modifies Transcriptomic Profile and Functions of Acute Myeloid Leukemia Cells

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**Keywords:** acute myeloid leukemia, bitter taste receptors, denatonium benzoate, bone marrow microenvironment, bitter compounds

## A Corrigendum on

### Denatonium as a Bitter Taste Receptor Agonist Modifies Transcriptomic Profile and Functions of Acute Myeloid Leukemia Cells

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In the original article, there was an error in the GEO database accession number. The correct GEO database accession number for gene expression data of denatonium-treated cells is GSE149548.

A correction has been made to the "Methods" section, paragraph 2, and the Data Availability Statement:

## GENE EXPRESSION PROFILING (GEP)

TAS2R expression was analyzed in 61 AML, 49 from a published dataset (32) and 12 new cases. As validation set, we used also 183 AML samples downloaded from The Cancer Genome Atlas (TCGA) ([https://gdc.cancer.gov/about-data/publications/laml\\_2012](https://gdc.cancer.gov/about-data/publications/laml_2012))(33). GEP after DEN treatment was performed in 5 newly diagnosed AML samples and THP-1 and OCI-AML3 cell lines. Three independent replicates of each condition were hybridized to Human Clariom S Arrays (Thermo Fisher Scientific) according to the manufacturer's recommendations. Data quality control, normalization (signal space transformation robust multiple-array average), and supervised analysis were carried out by Expression Console and Transcriptome Analysis Console software, respectively (Thermo Fisher

Scientific). For AML cells, data were normalized on vehicle-treated cells before comparison. Genes with a 1.5 fold difference and  $p \leq 0.05$  were considered for enrichment analyses. Downstream analyses were performed as reported in (32,34), and with Thomson Reuter's MetaCore software suite (Clarivate Analytics, Philadelphia, PA, USA). Gene expression data of denatonium-treated cells will be publicly available on the GEO database under the accession number GSE149548.

## DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and

accession number(s) can be found below: the NCBI Gene Expression Omnibus (GSE149548).

The authors apologize for these errors and state that this do not change the scientific conclusions of the article in any way. The original article has been updated.

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