

## GDSC fitted dose response description

Possible columns in GDSC fitted data results file. Not every listed column is present in every file.

Column	Description	Notes
<b>DATASET_VERSION</b>	Each dataset is processed (curve fitted and ANOVA analysis) as a whole.	
<b>IC50_RESULTS_ID</b>	Identifier for the fitted dose response	
<b>COSMIC_ID</b>	Cell identifier from the COSMIC database	
<b>CELL_LINE_NAME</b>	Primary name for the cell line	
<b>DRUG_ID</b>	Unique identifier for a drug. Used for internal lab tracking	
<b>DRUG_NAME</b>	Primary name for the drug	
<b>PUTATIVE_TARGET</b>	Putative drug target	

<b>MAX_CONC_MICROMOLAR</b>	Maximum micromolar screening concentration of the drug	
<b>MIN_CONC_MICROMOLAR</b>	Minimum micromolar screening concentration of the drug	
<b>LN_IC50</b>	Natural log of the fitted IC50	To convert to micromolar take the exponent of this value, i.e. $\exp(\text{IC50\_nat\_log})$
<b>AUC</b>	Area Under the Curve for the fitted model. Presented as a fraction of the total area between the highest and lowest screening concentration.	
<b>RMSE</b>	Root Mean Squared Error, a measurement of how well the modelled curve fits the data points.	Curves with RMSE > 0.3 are excluded prior to release as part of quality control.
<b>Z_SCORE</b>	Z score of the LN_IC50 ( $x$ ) comparing it to the mean ( $\mu$ ) and standard deviation ( $\sigma^2$ ) of the LN_IC50 values for the drug in question over all cell lines treated.	$Z = \frac{x - \mu}{\sigma^2}$