

Version 1.0

WebApp for Dose-Response Analysis of Multiple Experiments

User's guide

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The web application MDRA performs the dose-response analysis from multiple experiments.

• The four-parameter log-logistic (4pLL) model is used to fit dose-response data from each individual experiment. The 4pLL model function is given by

$$response = f(dose) = \phi^{(c)} + \frac{\phi^{(d)} - \phi^{(c)}}{1 + \exp\{\phi^{(b)}[\log(dose) - \log(\phi^{(e)})]\}}$$

The parameters $\phi^{(c)}$ and $\phi^{(d)}$ correspond to the lower and upper horizontal limits for mean response, respectively. The parameter $\phi^{(e)}$ is the effective concentration EC50. The parameter $\phi^{(b)}$ determines the slope of the curve. The natural logarithm ('log') is used in this model function.

- The DORES (dose-response screening) plot provides visualization of multiple dose-response experimental data sets in a single plot. It displays information from studies involving a large number of experiments: which concentration levels and ranges are chosen for each experiment; whether all experiments exhibit complete and biologically relevant dose-response relationships; and what the individual ECx estimates and their confidence intervals are. For details about the DORES plot, see Jiang, Xiaoqi and Annette Kopp-Schneider. Statistical strategies for averaging EC50 from multiple dose-response experiments. *Archives of Toxicology*, (2014).
- The meta-analysis approach is applied to average ECx estimates from multiple dose-response experiments. For details, see: Jiang, Xiaoqi, and Annette Kopp-Schneider. Summarizing EC50 estimates from multiple dose-response experiments: A comparison of a meta-analysis strategy to a mixed-effects model approach. *Biometrical Journal*, 56(3): 493-512 (2014).
- The 4pLL model is fitted using the R package drc (Ritz, C., Streibig, J.C., 2005. Bioassay analysis using R. *Journal of Statistical Software*. 12: 1–22.). The R package nlme (Pinheiro, J. C. and Bates, D. M. (2000). The R package metafor (Viechtbauer W., 2010. Conducting meta-analyses in r with the metafor package. *Journal of Statistical Software* 36:1-48.) is used for computing the heterogeneity estimators in meta-analysis.

1. Getting started with MDRA

Enter <u>http://biostatistics.dkfz.de/mdra/</u> in a browser window to access the WebApp (Web Application) MDRA. The browsers Chrome and Firefox are recommended.

To follow this user's guide, please first enter the webpage 'About' and download the exemplary data file "Example_DataSet.csv".

2. Data Upload

Only the CSV-File format is supported for uploading your data to the WebApp MDRA. CSV-Files can be easily created from Excel data by *saving as* type "(*.csv)".

- (1) Click on the Browse... button at the top of the left sidebar to choose the data file that you want to upload.
- (2) Tick the Header item to use the first row of the file for column titles. Make sure the column titles in the first row of your data do not contain any special characters or spaces. Short column titles are recommended. If there are no column titles shown in the first row of your CSV data, please untick the Header item and then your data will be imported with column titles V1, V2, V3...
- (3) Select the column delimiter and decimal point appropriate for your data file.

Column delimiter:	Decimal point:
Comma	Dot
Semicolon	Comma
🔘 Tab	

In the exemplary data file "Example_DataSet.csv", both the column delimiter and decimal point are commas.

On the preview page 'Data Set' –at the top of the webpage- you can see how your data is affected by changing the above items.

Before going to the next step, make sure that your data set has been correctly imported by MDRA. The numeric variables must be displayed with a dot ('.') as a decimal point or in scientific (e-) notation, whether the decimal point in your CSV-File is a dot or comma.

A correctly imported data set should look like this:

	Chemical	LabID	ExperimentID	Unit	Concentration	Response
1	Sodium_valproate	С	5	М	1e-04	0.981404 90255796
2	Sodium_valproate	С	5	М	1e-04	1.03256332906318
3	Sodium_valproate	С	5	М	1e-05	1.00942187693706
4	Sodium_valproate	С	5	М	1e-04	0.979999173519567
5	Sodium_valproate	С	5	М	1e-04	0.987768089590479
6	Sodium_valproate	С	5	М	1e-05	0.949336749452457
7	Sodium_valproate	E	7	М	0	0.991691341700312
8	Sodium_valproate	E	7	М	0	0.860742663385681
-		_	-		-	

3. Selecting variables for multiple experimental data for dose-response analysis

In the drop-down boxes, you will see all column titles from your data set, which can be selected as variable names for analyses in the next step.

*Select variable 'dose'	*Select variable 'response'
Concentration	▲
	Select additional identifier (e.g. Lab IDs)
Chemical	_
LabID	····
ExperimentID	er is optional, if this information is available.
Unit	
Concentration	
Response	
THE QUALITIE OF LONG/LUND.	

To perform a dose-response analysis from multiple experimental data, the variables 'dose' and 'response' and an identifier for 'experiment' (IDs) are required fields. An additional identifier is optional, e.g., it can be used when experiments are carried out in different laboratories and this information is available. An additional identifier can also be used for any other potential factors that may influence experimental results, such as technicians and experiment dates. In the exemplary data, the variables "Concentration" and "Response" have been selected for dose-response analysis, and the variables "ExperimentID" and "PartnerID" act as identifiers for 'experiment' and 'lab'.

*Select variable 'dose'	*Select variable 'response'
Concentration •	Response •
*Select identifier for 'experiment'	Select additional identifier (e.g. Lab IDs)
ExperimentID •	LabID 🗸
* = Required Fields. The additional identif	ier is optional if this information is available

4. Choice of conditions for dose-response analysis

(1) Choose the name of a measure used to describe the potency or toxicity of a chemical

Choose the name of a measure of toxicity (potency):

EC
IC
ED
ID

- a. EC: effective concentration (increasing curve)
- b. IC: inhibition concentration (decreasing curve)
- c. ED: effective dose (increasing curve)
- d. ID: inhibition dose (decreasing curve)

Note that the four options do not result in different estimated values. The selected name is used in results tables and plots. These four terms can be defined in two ways: relative and absolute. The relative ECx (ICx, EDx or IDx) refers to the concentration or dose that causes x% change of the maximal effect, which is the most common definition. We only used this relative definition in the dose-response analysis.

(2) Choose the quantile of the measure.

For example, enter "50" in the input box to calculate EC50 for the exemplary data:

The quantile of EC/IC/ED/ID:					
50					
e.g., enter 50 for estimation of EC50/IC50/ED50/ID50.					

(3) Define the criteria for the experiment having no biologically relevant effect. The lack of relevance can be quantitatively assessed upon inspection of the difference

between the lower and upper limits $\phi^{(c)}$ and $\phi^{(d)}$. Specifically, for a predefined percentage r%, if $|(\phi^{(d)} + \phi^{(c)})/\phi^{(d)}| < r$ %, then the observed dose-response relationship may be considered as not biologically relevant. In the exemplary data set, our choice was r=40, so we then entered 0.4 in the input box.



- e.g., the maximum change of mean observed effect is less than 40% (0.4),
- (4) Restrictions on the limits for lower $\phi^{(c)}$ and upper $\phi^{(d)}$ plateaus of a 4-parameter loglogistic curve can be chosen, e.g., the lower limit could be restricted to a positive value (i.e., enter 0 for lower limit for lower plateau). Note that the model parameters $\phi^{(c)}$ and $\phi^{(d)}$ are estimated, even when their limit values are restricted.

Restriction on 4-parameter log-logistic curve:

Lower limits for lower plateau	Upper limits for upper plateau	
Screening experiments	Averaging analysis	

If restrictions are not given (default setting), the 4-parameter log-logistic curve is allowed to run from negative infinity to positive infinity or the opposite. In case response values are normalized between 0 and 100%, lower limits for the lower plateau could be restricted to 0 or upper limits for the upper plateau could be restricted to 1. This may improve the estimation of individual ECx values, but only when the highest concentration (dose) level does not cover the lower or upper plateau of the dose-response curve

5. Screening dose-response experimental data

Click on the Screening experiments button at the end of the right sidebar, choose the webpage 'DORES plot', and you will get an overview of all your dose-response data from multiple experiments. The title and x-axis label can be changed in the corresponding input boxes to the right of the DORES plot. Click on the **Source** button or right mouse click the button and choose "Save Link as" to download the DORES plot.



The resulting table displayed in the webpage 'Single Experiment Analysis' provides this information about the fitted curve and estimated measure of toxicity for each single experiment:

- the four estimated parameters of the log-logistic curve, i.e., 'Hill Slope', 'lower asymptote', 'upper asymptote' and 'natural logEC50' (natural logarithm of EC50);
- estimated measure of toxicity (e.g. EC50) and its 95% confidence interval.

About Data Set DORES Plot Single Experiment Analysis Averaging Results

🛓 Results Table Download

The following result talbe is saved as CSV-file for Download

La Single Curve Download

The dose-response curve plotted for each single experimental data is saved as PDF-file for Download

Experiment ID	b (=Hill Slope)	c (=lower asymptote)	d (=upper asymptote)	e (=natural logEC50)	EC50	Lower 95%-CL	Upper 95%-CL
1	-7.98e+00	8.90e-01	9.84e-01	-1.32e+01	1.85e-06	4.95e-23	6.93e+10
2	1.32e-01	1.91e-01	1.00e+00	-1.50e+00	2.22e-01	3.64e-29	1.36e+27
3	5.29e+00	8.85e-01	1.02e+00	-1.87e+01	7.54e-09	1.75e-10	3.24e-07
4	5.84e-01	-2.30e-03	1.01e+00	-3.40e+00	3.35e-02	1.08e-24	1.04e+21
5	4.23e-05	1.03e+00	1.00e+00	-5.04e+01	1.25e-22	0.00e+00	Inf
6	2.32e+00	4.57e-03	9.76e-01	-4.37e+00	1.26e-02	1.05e-02	1.52e-02
7	1.69e+00	-3.63e-02	9.75e-01	-5.42e+00	4.41e-03	3.96e-03	4.91e-03
8	8.44e-01	6.64e-01	9.98e-01	-7.95e+00	3.52e-04	1.93e-05	6.45e-03
9	-5.41e-01	1.01e+00	3.25e+00	-1.30e+00	2.74e-01	NA	NA
10	2.25e+00	-7.35e-03	9.39e-01	-5.36e+00	4.71e-03	3.84e-03	5.78e-03
11	-5.26e-05	1.00e+00	7.50e-01	-3.10e+00	4.49e-02	0.00e+00	Inf
12	2.96e+00	1.39e-03	1.04e+00	-4.79e+00	8.33e-03	6.91e-03	1.00e-02
13	1.56e+00	-1.30e-02	9.95e-01	-5.73e+00	3.23e-03	2.46e-03	4.24e-03
14	1.85e+00	2.18e-03	9.67e-01	-5.65e+00	3.52e-03	2.62e-03	4.72e-03
15	6.45e+00	1.79e-03	9.60e-01	-4.21e+00	1.49e-02	1.42e-02	1.56e-02
16	4.03e+00	-2.69e-02	9.55e-01	-4.18e+00	1.54e-02	1.45e-02	1.63e-02
17	2.97e+01	2.51e-02	9.85e-01	-3.48e+00	3.10e-02	2.98e-02	3.22e-02
18	2.60e+00	3.75e-03	9.64e-01	-4.40e+00	1.23e-02	1.13e-02	1.34e-02
19	1.39e+01	-1.46e-02	1.19e+00	-4.17e+00	1.54e-02	1.12e-02	2.13e-02
20	-5.81e+00	9.33e-01	1.04e+00	-1.36e+01	1.29e-06	2.22e-09	7.55e-04
21	4.13e-05	1.10e+00	1.00e+00	-3.16e-02	9.69e-01	0.00e+00	Inf
22	-4.39e-06	1.00e+00	6.42e-01	1.04e+01	3.34e+04	NA	NA

In this step, the input of conditions for dose-response analysis can still be changed. After inputs have been changed, click on the Screening experiments button again, and the DORES plot and resulting table will be updated. For example, enter 0 and 1 in the input box of the lower and upper limits for plateaus.

ower limits for lower plateau	Upper limits for upper pla
Screening experiments	Averaging analysis

Upper 95%-CL	Lower 95%-CL	EC50	e (=natural logEC50)	d (=upper asymptote)	c (=lower asymptote)	b (=Hill Slope)	Experiment ID
Inf	0.00e+00	2.77e+02	5.62e+00	8.19e-01	1.00e+00	-1.16e-05	1
1.39e+25	1.89e-27	1.62e-01	-1.82e+00	1.00e+00	2.11e-01	1.36e-01	2
3.36e-07	1.86e-10	7.91e-09	-1.87e+01	1.00e+00	8.85e-01	5.40e+00	3
NA	NA	1.18e-02	-4.44e+00	1.00e+00	0.00e+00	8.83e-01	4
NA	NA	1.29e-22	-5.04e+01	1.00e+00	1.03e+00	2.44e-05	5
1.52e-02	1.05e-02	1.26e-02	-4.37e+00	9.76e-01	4.57e-03	2.32e+00	6
4.72e-03	3.79e-03	4.23e-03	-5.47e+00	9.68e-01	0.00e+00	1.87e+00	7
6.46e-03	1.92e-05	3.53e-04	-7.95e+00	9.98e-01	6.64e-01	8.44e-01	8
Inf	0.00e+00	2.63e+10	2.40e+01	1.00e+00	1.04e+00	-3.47e-01	9
5.73e-03	3.82e-03	4.68e-03	-5.36e+00	9.37e-01	0.00e+00	2.31e+00	10
Inf	0.00e+00	1.20e-02	-4.42e+00	7.50e-01	1.00e+00	-7.76e-05	11
1.04e-02	7.08e-03	8.56e-03	-4.76e+00	1.00e+00	5.50e-03	3.22e+00	12
4.15e-03	2.42e-03	3.17e-03	-5.76e+00	9.94e-01	0.00e+00	1.60e+00	13
4.72e-03	2.62e-03	3.52e-03	-5.65e+00	9.67e-01	2.18e-03	1.85e+00	14
1.56e-02	1.42e-02	1.49e-02	-4.21e+00	9.60e-01	1.79e-03	6.45e+00	15
1.61e-02	1.44e-02	1.52e-02	-4.19e+00	9.48e-01	0.00e+00	4.42e+00	16
3.12e-02	3.11e-02	3.12e-02	-3.47e+00	9.85e-01	1.06e-02	2.53e+02	17
1.34e-02	1.13e-02	1.23e-02	-4.40e+00	9.64e-01	3.74e-03	2.60e+00	18
2.34e-02	1.11e-02	1.61e-02	-4.13e+00	1.00e+00	9.10e-03	3.00e+01	19
7.43e-04	1.77e-09	1.15e-06	-1.37e+01	1.00e+00	9.33e-01	-6.09e+00	20
Inf	0.00e+00	9.69e-01	-3.16e-02	1.00e+00	1.10e+00	1.89e-05	21
Inf	0.00e+00	4.30e+04	1.07e+01	6.44e-01	1.00e+0	-2.41e-05	22

Click on the **Levent Stable Download** button or right mouse click the button and choose

"Save Link as" to download the resulting table saved in a CSV-File.

If you want to see a fitted log-logistic curve for each experiment data, click on the

Single Curve Download button or right mouse click the button and choose "Save Link as" to download plots saved in a PDF-File.



6. Averaging dose-response data from multiple experiments

Experimental data sets not containing useful information for further averaging analysis should be excluded, i.e., data sets for which no EC50 estimate could be obtained or where EC50 estimates have very large confidence intervals.

Under the webpage 'DORES plot', all the experiment IDs are listed at top of this webpage:



The elements of experiment ID are extracted from the identifier for 'experiment' that you select in the step 'Uploading data'.

For the exemplary data, the experiment IDs: 1,2,3,4,5,8,9,11,20,21,22 are excluded from the averaging analysis. Thus, tick these elements in the check box list and click on the

Screening experiments b

button again, and the DORES plot will be updated.

Upload complete		Exclusion of 'no-effects'	experiments from averaging analysis: 2 🔽 3 🔽 4 🔽 5 🔲 6 🛄 7 🖾 8 🖉 9 🚞	10 🗹 11 📄 12 📄 13 📄 14
Header		16 17 18	19 🔽 20 🖳 21 💌 22	
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Semicolon	Comma	6 - •····	••••••••••••••••••••••••••••••••••••••	 D
1 TAD				
You can see how your data is affected	in the preview page 'Data Set'.	19 - •	·····••···••••••••••••••••••••••••••••	F
Select variable 'dose'	*Select variable 'response'	18 - •		F
Concentration	▼ Response ▼	14 - •	······································	E
Select identifier for 'experiment'	Select additional identifier (e.g. Lab IDs)	13 - •	······································	E
ExperimentID	▼ LabID ▼	12 - •	······································	E
* = Required Fields. The additional ide	ntifier is optional, if this information is available.	10 - •	······································	E
Choose the name of a measure of toxic	city (potency):	7 - •	•••••••••••••••••••••••••••••••••••••••	E
Note that the four options do not result	in different estimated values but are different			
names a measure later used in results	tables and plots.	1/ - •	• • • • • • • • • •	A
The quantile of EC/IC/ED/ID:		16 - •	• • • • • • • • •	Α
50				
e.g., enter 50 for estimation of EC50//C	50/ED50/ID50	15 - •	····· • • • • • • • • • • • • • • • • •	Α
efine the experiment having no biolog	rical relevant effect:			
		/ <i>/</i>		<u> </u>
0.4		0	1e-04 0.001 0.01 0.1	1 10
e.g., the maximum change of mean obs	served effect is less than 40% (0.4).		Concentration [M]	
Restriction on 4-parameter log-logi	stic curve:	+: increasing	curve; -: decreasing curve; N: model convergence failed	I
ower limits for lower plateau	upper limits for upper plateau	*: no biologic	ally relevant effect	
×				
	Automatica construite			
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Finally, click on the Averaging analysis button at the end of the left sidebar and choose the webpage 'Averaging Results' to average dose-response data from multiple experiments. The results of the averaging analysis are displayed in a forest plot. The title can be changed in the input box to the right of the forest plot. Click on the Download button or right mouse click the button and choose "Save Link as" to download the forest plot.



Forest plot

Experiment ID:		Sodium valpro	ate	EC50 [95% CI]
6		⊢ •-+		1.26e-02 [1.05e-02 , 1.52e-02]
7		⊢ •+		4.41e-03 [3.96e-03 , 4.91e-03]
10		⊢		4.71e-03 [3.84e-03 , 5.78e-03]
12		⊢•-+		8.33e-03 [6.91e-03 , 1.00e-02]
13	⊢ −•			3.23e-03 [2.46e-03 , 4.24e-03]
14	H	- −-+		3.52e-03 [2.62e-03 , 4.72e-03]
15		H		1.49e-02 [1.42e-02 , 1.56e-02]
16		 ■ 		1.54e-02 [1.45e-02 , 1.63e-02]
17		I T		3.10e-02 [2.98e-02 , 3.22e-02]
18		+ •+		1.23e-02 [1.13e-02 , 1.34e-02]
19		⊢ •+		1.54e-02 [1.12e-02, 2.13e-02]
Meta-analysis:				Weighted mean [95% Cl]
Hunter-Schmidt:		⊢		9.13e-03 [5.57e-03 , 1.50e-02]
Hedges:		⊢		9.10e-03 [5.55e-03 , 1.49e-02]
DerSimonian-Laird:		⊢ →		9.11e-03 [5.55e-03 , 1.49e-02]
Sidik-Jonkman:		⊢		9.10e-03 [5.55e-03 , 1.49e-02]
ML:		⊢		9.10e-03 [5.55e-03 , 1.49e-02]
REML:		⊢ →		9.10e-03 [5.55e-03 , 1.49e-02]
Empirical Bayes:		⊢		9.10e-03 [5.55e-03 , 1.49e-02]
				Unweighted mean [95% Cl]
		⊢ →		9.05e-03 [5.52e-03 , 1.48e-02]
г // —				
0	0.001	0.01	0.1	

Overall average EC50 estimate