# MFIT v1.0.0 – Quick Start User Guide

### Jacques Bodin, 2019

## Introduction

MFIT is a Windows-based computer software package for the analytical modeling of concentration breakthrough curves (BTCs) from tracer experiments in fractured and karst aquifers. The software integrates four transport models built on a multi-flow approach which assumes that the spatial heterogeneity of the aquifer can be approximated by a combination of independent one-dimensional channels. The four transport models are referred to as MDMi (Multi Dispersion Model, instantaneous injection), MDMed (Multi Dispersion Model, exponentially decaying injection), MDP-SFDM (Multi-Double Porosity, Single-Fracture Dispersion Model), and MDP-2RNE (Multi-Double Porosity, two-region non-equilibrium model). The conceptual and mathematical description of these models are given in the companion article to this manual and will not be represented here. The MDMi, MDMed, MDP-SFDM and MDP-2RNE models were implemented as independent codes which can be run either as console applications or using the MFIT program as a Graphical User Interface (GUI). MFIT provides a GUI for (i) the importation and graphic visualization of user-provided BTC data, (ii) the parameterization, direct run, and graphical output of the analytical transport models, (iii) the inversion (automatic calibration) of model parameters for optimal curve fitting, and (iv) the assessment of the uncertainty of calibrated parameter values. Both the optimization and uncertainty analysis of model parameters are carried out using routines from the PEST software package (<http://www.pesthomepage.org/>).

## Direct (forward) transport simulations

### Using the programs MDMi.exe, MDMed.exe, MDP\_SFDM.exe, and MDP\_2RNE.exe as console applications

The model input parameters are read from a file named « input.txt », which can be built using any text editor program. The format of the input file format differs slightly depending on the model, see below.

#### Structure of the input parameter file (input.txt) for the program MDMi.exe

Lower limit of the simulation time range (Tmin), real number

Upper limit of the simulation time range (Tmax), real number

Number of time-concentration BTC points simulated by the model, integer

Total system flow rate (Q), real number

Number of channels, integer

Solute mass flowing through the first channel, real number

Mean transit time (T0) in the first channel, real number

Peclet number (Pe) of the first channel, real number

Solute mass flowing through the second channel, real number

Mean transit time (T0) in the second channel, real number

Peclet number (Pe) of the second channel, real number

etc.

#### Structure of the input parameter file (input.txt) for the program MDMed.exe

Lower limit of the simulation time range (Tmin), real number

Upper limit of the simulation time range (Tmax), real number

Number of time-concentration BTC points simulated by the model, integer

Initial/maximum injection concentration at the inflow boundary of the system (C0), real number

Number of channels, integer

Flowrate contribution ratio of the first channel, real number

Mean transit time (T0) in the first channel, real number

Peclet number (Pe) of the first channel, real number

Gamma coefficient of the first channel, real number

Flowrate contribution ratio of the first channel, real number

Mean transit time (T0) in the second channel, real number

Peclet number (Pe) of the second channel, real number

Gamma coefficient of the second channel, real number

etc.

#### Structure of the input parameter file (input.txt) for the program MDP\_SFDM.exe

Lower limit of the simulation time range (Tmin), real number

Upper limit of the simulation time range (Tmax), real number

Number of time-concentration BTC points simulated by the model, integer

Total system flow rate (Q), real number

Number of channels, integer

Solute mass flowing through the first channel, real number

Diffusion parameter (Beta coefficient) of the first channel, real number

Mean transit time (T0) in the first channel, real number

Peclet number (Pe) of the first channel, real number

Solute mass flowing through the second channel, real number

Diffusion parameter (Beta coefficient) of the second channel, real number

Mean transit time (T0) in the second channel, real number

Peclet number (Pe) of the second channel, real number

etc.

#### Structure of the input parameter file (input.txt) for the program MDP\_2RNE.exe

Lower limit of the simulation time range (Tmin), real number

Upper limit of the simulation time range (Tmax), real number

Number of time-concentration BTC points simulated by the model, integer

Total system flow rate (Q), real number

Number of channels, integer

Solute mass flowing through the first channel, real number

Length of the first channel, real number

Mean transit time (T0) in the first channel, real number

Peclet number (Pe) of the first channel, real number

Psi coefficient of the first channel, real number

Omega coefficient of the first channel, real number

Solute mass flowing through the second channel, real number

Length of the second channel, real number

Mean transit time (T0) in the second channel, real number

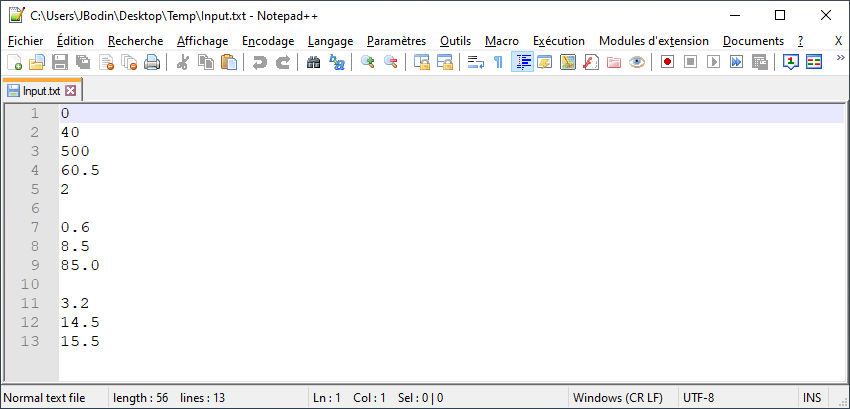
Peclet number (Pe) of the second channel, real number

Psi coefficient of the second channel, real number

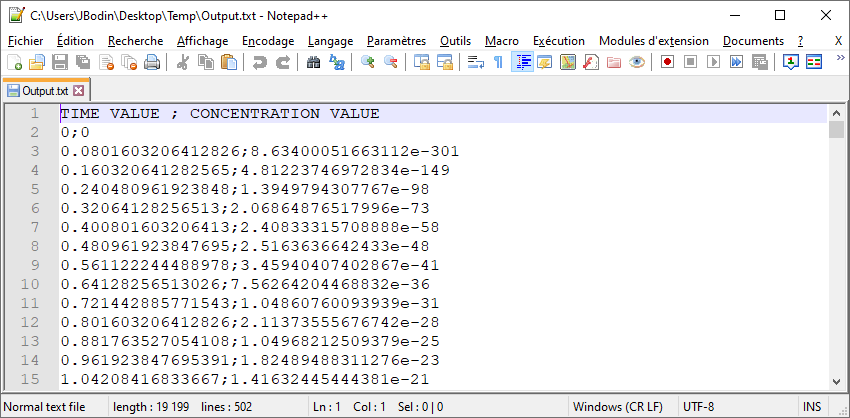
Omega coefficient of the second channel, real number

etc.

By running the transport model code, a series of time-concentration values are computed and saved in a text file named « output.txt ». Care must be taken to use consistent units for all the variables in the input file, which also determine the time and concentration units of the output file (e.g., concentrations are in g/m3 if the the system flowrate has been expressed in m3/s, the solute-mass in g, and the mean transit times in s). Fig. 1 shows an example of input file for a two-channel MDMi model. The related output file written by the MDMi program is shown in Fig. 2.



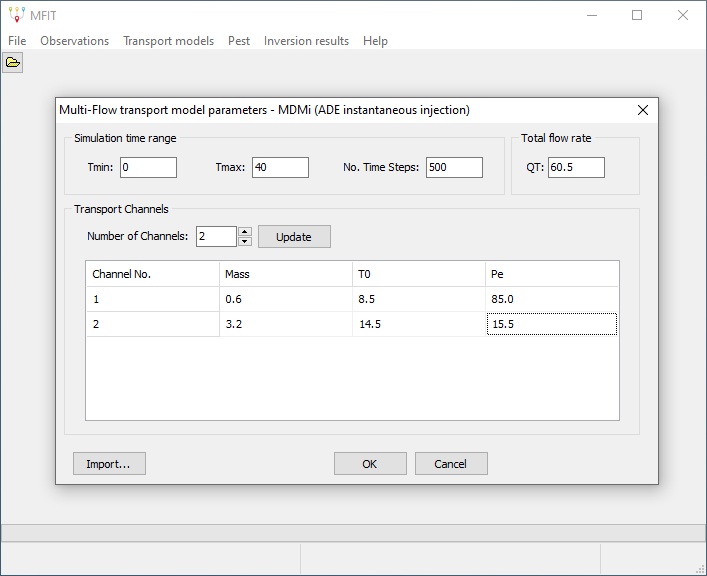
**Fig. 1.** Example of input file for a two-channel MDMi model.



**Fig. 2.** Ouput file written by the MDMi program from the input file shown in Fig. 1. Only the first lines of the output file (which contains 500 time-concentration lines) are displayed.

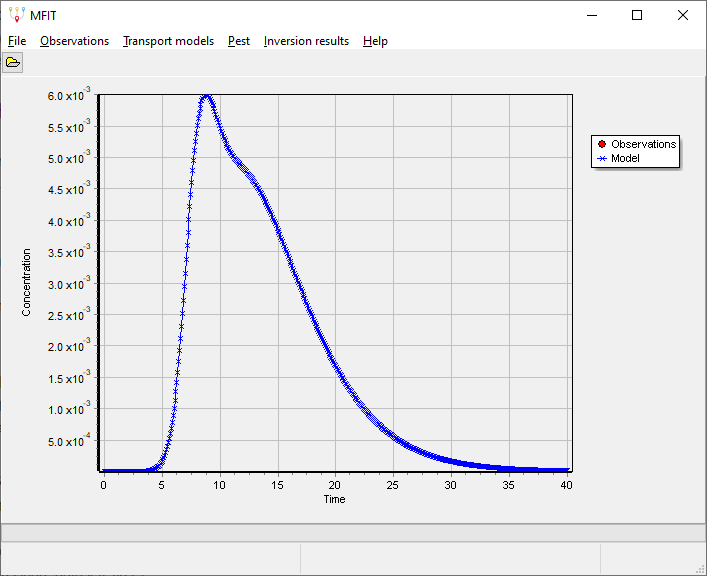
### Using MFIT as a GUI for the programs MDMi.exe, MDMed.exe, MDP\_SFDM.exe, and MDP\_2RNE.exe

1. Select **File | New** from the main menu.
2. Select **Transport models | Working Directory** and pick a folder for the writing of the working (temporary) files and model output files.
3. Select **Transport models**, choose one of the four proposed transport models, and then select **Parameters**. A dialog will be displayed that allows you to set the simulation parameters. The example shown in Fig. 3 is similar to the two-channel MDMi model input.txt file of Fig. 1. The dialog can also be filled by importing a pre-existing input.txt file (or any other text file conforming to the same structure) using the Import button. When finished, click the OK button.



**Fig. 3.** GUI dialog for the setting of the MDMi model parameters.

1. Select **Transport models | Compute BTC**. The simulated BTC will be displayed (in blue, see Fig. 4). By clicking on the bottom- of left-axis black line, a dialog appears which allows to change the X- and Y-axis ranges.



**Fig. 4.** Example of MFIT-simulated BTC (two-channel MDMi model).

1. Optionally, select **File | Save As** for saving the simulation parameters and the path of the working directory into a MFIT file (extension: .mfi). The MFIT files can be edited with any text editor and can be reopened using the **File | Open** menu command.

## Importing BTC data

If you already have some tracer-test data and if you want to use the MFIT software for modeling the observed concentration BTC, the first step is to import your BTC data in MFIT. The steps are as follows:

1. Start the MFIT software and select **File | New** from the main menu.
2. Select **Observations** from the main menu and fill the Time-Concentration-Weight table dialog. The « weight » value that must be assigned to each line is only relevant if PEST is going to be used for the automatic calibration of a MFIT transport model against the user-provided BTC data (see section 5.1 below). If you don’t plan to use PEST, just apply any constant weight value to all the lines. The BTC data (including weights) can be either filled manually (use « Insert Row » or « Add Rows » for adding as many rows needed), or copy and pasted from a spreadsheet (e.g. Excel), or imported from a CSV text file (separator : comma).
3. Large BTC datasets can be resampled by applying a filter routine accessible from the « Filter » button. A general rule in BTC-data analysis is to seek the best compromise between simplification and preservation of the information content of the BTC, especially in the case of multiple local concentration peaks. A number of observations between 50 and 200 is appropriate in most cases.
4. Click on the Ok button to display (in red) the observed BTC.
5. Optionally, select **File | Save As** for saving the Time-Concentration-Weight table into a MFIT file (extension: .mfi). The MFIT files can be edited with any text editor and can be reopened using the **File | Open** menu command.

## Model-curve fitting of the observed BTC

### Manual trial-and-error procedure

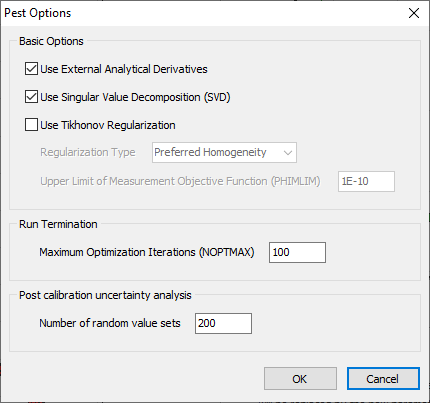
The procedure described in section 2.2 above can be used for simulating a model BTC that will be plotted in the same graph as the observed BTC. A better agreement between the model and observed BTCs can be seeked from a trial-and-error procedure which consists in changing the model parameters, runing again the model, and evaluating (either by « eye » or by computing some error criteria between the model and observed BTCs) if the new set of model parameters is better (or worse) than the preceeding one.

### PEST model inversion

PEST is a model-independent optimizer that can be used for seeking the set of model parameters that allows the « best fit » between the simulated and observed BTCs. As detailed in the PEST documentation (<http://www.pesthomepage.org/Downloads.php>), a number of settings may influence the way PEST operates. A decision was made to narrow the MFIT-PEST GUI dialog to the most important settings in order to keep a simple (i.e. user friendly) but effective program interface. Other PEST control variables were fixed in the MFIT code to expected optimum settings, based on recommandations from the PEST user manual and/or experience knowledge. Additional GUI dialogs can be easily added to MFIT for access to any of these control variables, a modified version of MFIT can be made available upon request.

The objective function that is minimized by PEST may be expressed as a « measurement objective function » only, or as the sum of a « measurement objective function » and a « regularization objective function ». The measurement objective function is defined as the sum of squared weighted residuals between the observed and the simulated c(t) values. The weight value assigned to each time-concentration line in the observation table (see Section 3) has a direct influence on the computation of the measurement objective function and therefore on the optimization results. If the same weight-value (e.g., 1.0) is set for each line, PEST will pay the same attention to each concentration value and will search for the best overall model-curve fitting of the tracer BTC. If you prefer to promote the model fit on a specific part of the tracer BTC (e.g., the concentration peak), use higher relative weight values for the related lines (or lower relative weight values for the other lines). The regularization objective function acts as a penalty function for deviations from some preferred parameter conditions. Two options have been implemented in MFIT. The first option, referred to as « preferred homogeneity », promotes a solution of minimum variance for the model parameters pertaining to the different channels. With the second option, referred to as « preferred value », the optimization process seeks the solution that is the closest to some prior estimates of the model parameters. The steps for optimizing a MFIT transport model are as follows:

1. Import BTC data (see Section 3)
2. Choose and run one of the four MFIT transport problems, using trial-and-error as a first optimization method (see Section 2.2).
3. Select **Pest | Options** from the main menu. A dialog will be displayed that allows to set the PEST options, see Fig. 5.



**Fig. 5.** Pest option GUI dialog.

The first and second checkboxes should remain checked in most cases. The first checkbox is used to enable or disable the analytical computation of the Jacobian matrix derivatives. Unchecking this checkbox will force PEST to compute all the derivatives using finite differences, which is both time consuming and less accurate. The second checkbox activates (or deactivates) the use of the Singular Value Decomposition (SVD) method for solving the inverse problem. Using SVD is recommended as it guarantees the numerical stability of the inversion process.

The third checkbox allows to add some « preferred homogeneity » or « preferred value » Tikhonov regularization constraints to the optimization process. If the second option is selected, PEST will use the parameter values specified in the model dialog (Fig. 3) as « preferred values ». As discussed below, two options are also available for the number of channels to be considered in the optimization procedure. The first option, referred to as « User-specified channels », considers a fixed number of channels as specified in the model dialog. With the second option, referred to as « Automatic multiple channels », a series of automatic tracer BTC fitting is performed for a decreasing number of channels ranging from Nmax (upper bound) to 1. If both the the « preferred value » Tikhonov regularization and « Automatic multiple channels » options are selected, and if several channels were specified in the model dialog, only the parameters pertaining to the first channel are used as preferred values and are applied to all the channels during the optimization.

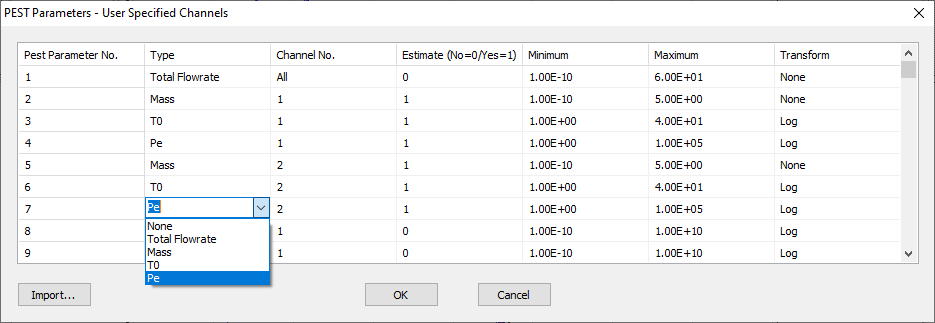
The variable PHIMLIM controls the trade-off between the model curve-fitting of the observed BTC and the regularization constraints, i.e. adherence to preferred homogeneity or preferred value conditions. As discussed in the PEST documentation, a suitable value for PHIMLIM can be determined by first running PEST without Tikhonov regularization, determining the measurement objective function achieved by PEST, and then setting PHIMLIM 5 to 10 percent higher to this value.

The NOPTMAX variable sets the maximum allowed number of optimization iterations for each PEST run. A minimum value of 100 is recommended for this variable to ensure that PEST terminates its execution because a convergence criteria has been reached rather than because of the NOPTMAX limit.

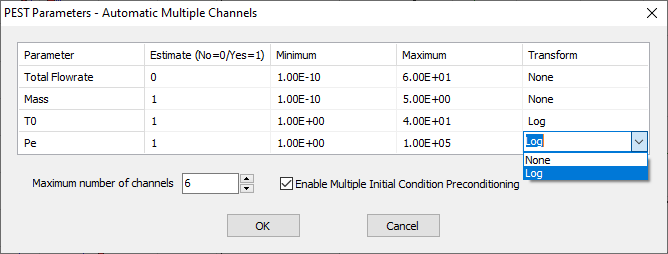
The last edit control on the PEST option window dialog allows to set the number of parameter sets that will used for assessing the post-calibration parameter uncertainty, if requested by the user after completion of a « User-specified channels » optimization (see section # below). This number should be set as high as possible within computational time constraints.

When the PEST options have been configured, click the OK button.

1. Select **Pest | Parameters** from the main menu and then select either **User-Specified Channels** or **Automatic Multiple Channels**. A different dialog (Fig. 6 and Fig. 7) will be displayed depending on the selected option and also on the selected MFIT model. Each of these dialogs allows to specify the parameters to be estimated by PEST, their lower and upper bounds, and allows to choose whether the optimization should be conducted on the parameters themselves or on the logarithms of the parameters. The lower and upper parameter bounds must reflect as accurately as possible the range of prior parameter uncertainty as they are used both in the optimization process (internal parameter scaling « boundscale » option, see the PEST manual) and for the assessment of post-calibration parameter uncertainty (see section # below). The log-transformation is recommended for the parameters that are non-linearly related to simulated concentration values (e.g. T0, Pe, Gamma, Beta, Psi, …) as a means to promote the numerical stability of the inversion process. The parameters that are linearly related to the model’s response, i.e. the total system flowrate Q and the solute mass in each channel for the MDMi, MDP-SFDM and MDP-2RNE models, the initial source concentration C0 and the flowrate contribution ratio of each channel in the MDMed model, should be let untransformed.

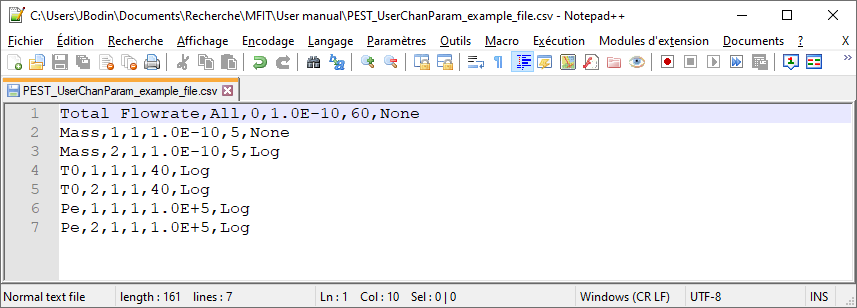


**Fig. 6.** GUI dialog for the selection and setting of model parameters to be optimized with the « User-Specified Channels » option.



**Fig. 7.** GUI dialog for the selection and setting of model parameters to be optimized with the « Automatic Multiple Channels » option.

The « Import » button in the lower left corner of the User-Specified Channels PEST parameter dialog allows to import a CSV text file (separator: comma, see example Fig. 8) for implementing the PEST parameter table.



**Fig. 8.** Example of CSV text file than can be imported into the PEST parameter table of Fig. 6.

While the « User-Specified Channels » option considers a fixed number of channels as specified in the model dialog (Fig. 3), the selection of the « Automatic Multiple Channels » option leads to a series of PEST model calibrations using a decreasing number of channels ranging from Nmax (upper bound) to 1. The maximum number of channels, Nmax, can be specified using the edit control located below the PEST parameter table (see Fig. 7). The checkbox at the right of the Nmax edit control activates (or deactivates) the use of a multi-start optimization method which consists in multiplying the PEST runs using different sets of initial parameter values. This method, wich is detailed in the companion article to this manual, can be computationally expensive but significantly reduces the risk for PEST to be trapped in a local minima of the parameter space. It is therefore advisable to use this method.

When the setting of the PEST parameter is complete, click the OK button to validate.

## Contacts and technical support

If you have a question or problem, feel free to send me an email ([jacques.bodin@univ-poitiers.fr](mailto:jacques.bodin@univ-poitiers.fr)). I will also appreciate any comment or suggestion about further improvements of the MFIT software.