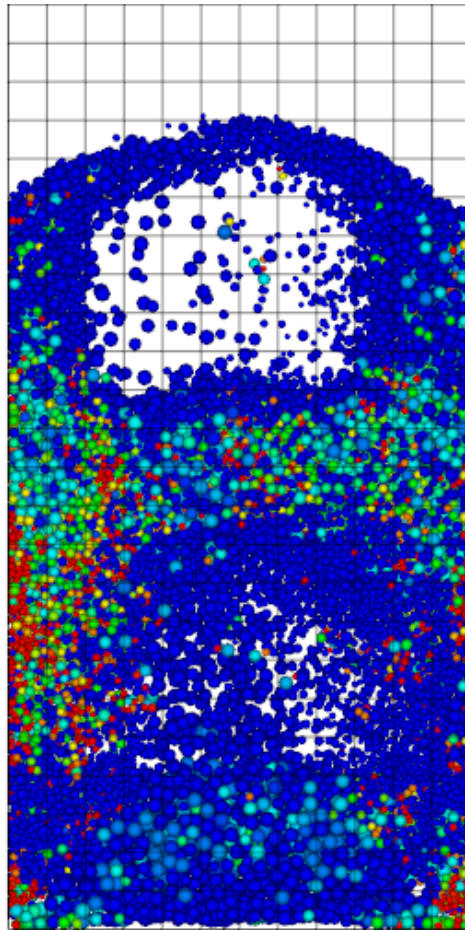


## Multiphase Flow with Interphase eXchanges



Version MFiX-2013-1

February 2013



The front page image shows air fluidization (at about  $2.5 U_{mf}$ ) of a Geldart type A powder with a particle size distribution (from about 40 to 110 microns) where the particles are colored with the magnitude of cohesive forces.

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- The use of MFIX is to be acknowledged in any published paper based on computations using this software by citing the MFIX theory manual. Some of the submodels are being developed by researchers outside of NETL. The use of such submodels is to be acknowledged by citing the appropriate papers of the developers of the submodels.
- The authors would appreciate receiving any reports of bugs or other difficulties with the software, enhancements to the software, and accounts of practical applications of this software.

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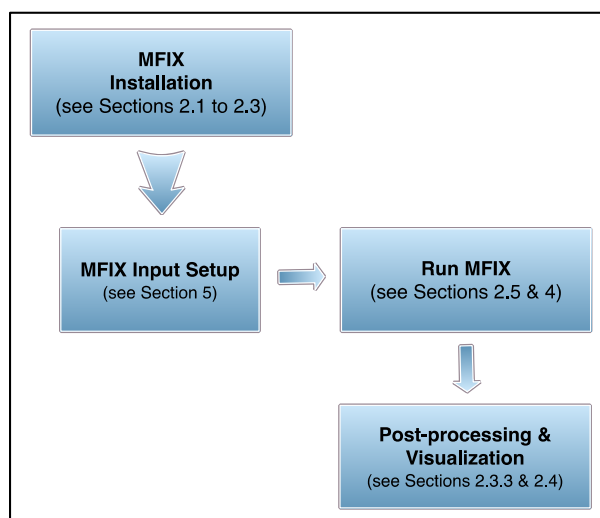
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## 1. Introduction

MFIX is an open-source multiphase flow solver and is therefore free to download and use. A one-time free registration is required prior to downloading the source code. To register, go to the MFIX website at <https://mfix.netl.doe.gov>, click on "Register" at the bottom of the home page or go directly to <https://mfix.netl.doe.gov/registration.php>. Complete the form, read the notice and click on "I Agree" to submit your application. Once your application has been reviewed and accepted, you will receive an email notification, and instructions to download the code.



The above flow chart provides an overview of the work flow and related sections in this document for quick review.

Some Frequently Asked Questions located at <https://mfix.netl.doe.gov/faq.php> may be useful to review before downloading MFIX.

## 2. Setting Up and Running MFIX on a UNIX/LINUX Workstation

Please read this file carefully and if you still have problems with installation or use, send an email to [mfix-help@mfix.netl.doe.gov](mailto:mfix-help@mfix.netl.doe.gov) if you have any. Before sending a question, please subscribe to mfix-help mailing list at <https://mfix.netl.doe.gov/sympa/info/mfix-help>. Once subscribed, please search the archives at <https://mfix.netl.doe.gov/sympa/arc/mfix-help> as your question might be already answered.



MFIX is primarily designed to run on Linux operating system. The setup described in this section is the preferred way of installing and running MFIX. You should be familiar with basic Linux operating system commands and procedures before attempting to use MFIX.

## 2.1. Creating MFIX Directory

It is assumed that you have registered and downloaded MFIX. MFIX is distributed as a tar ball, called **mfix.tar.gz**. After downloading the tar ball, make sure it is located in your home directory.

To install MFIX (version 2013-1) from the tar file, go to the home directory and type

```
gunzip -d mfix.tar.gz
```

```
tar xvf mfix.tar
```

Extracting the tar ball will create the directory mfix, which contains the following subdirectories:

Subdirectory	Contents
model	MFIX source files
post_mfix	Post_mfix source files
tutorials	Several example problems and corresponding pdf files describing some of the tutorials exists, which are useful for a new user
tests	Test simulations used to verify the code during development
visualization_tools	Tools to visualize MFIX results (not supported anymore)
tools	Development tools
doc	MFIX manuals in pdf format

## 2.2. Alias creation

For convenience, create aliases by adding the following lines to the `.login` or the `.cshrc` file (assuming MFIx was installed in the home directory, and you are using the C shell):

```
alias mkmfix sh ~/mfix/model/make_mfix
```

```
alias post ~/mfix/post_mfix/post_mfix
```



Note that makefile should be invoked as “sh make\_mfix.” Erroneously invoking it as “make\_mfix” will cause the user-defined files not to be used for the build. The alias for post\_mfix will be operational when post\_mfix is compiled (see section 2.3.2).

After adding the aliases in the `.login` or `.cshrc` file, the user should logout and login or type: **source .login** so that the aliases are functional. This step needs to be done only once. Then the mfix make file can be invoked from any directory by typing **mkmfix**, and the post-processor can be activated by typing **post**. The creation and utilization of aliases is not required to use MFIx. Aliases provide convenient shortcuts to common commands.

## 2.3. Creation of executable files

Installing MFIx only requires to convert the source code into an executable file. This is called the compilation of the code. You need a Fortran compiler to compile MFIx. Before continuing, make sure you have a working Fortran compiler on your computer. Please contact your system administrator if you are not sure how to properly install and test the compiler.

The distribution of MFIx provides two sets of source codes:

Name	Purpose	Location of source code	Name of executable
<b>MFIX</b>	Multiphase flow solver	~/mfix/model	<b>mfix.exe</b>
<b>POSTMFIX</b>	Simple post processing of the data. This is a text-based program. Data is extracted based on (I,J,K) location in the computational domain.	~/mfix/post_mfix	<b>post_mfix</b>





The open-source visualization tools ParaView and VisIt can be used to visualize and post-process MFIX results. They can be downloaded from:

**ParaView:** <http://www.paraview.org/>

**VisIt:** <https://wci.llnl.gov/codes/visit/home.html>

Please follow the instructions on the above websites to install ParaView or VisIt.

## 2.3.1.MFIX Makefiles

Both MFIX and POSTMFIX are compiled through a makefile, called `make_mfix` for MFIX and `make_post` for POSTMFIX. It is not expected that these files need to be modified, but expert users may edit the makefiles to add compilations options or flag specific to their architecture. The next section focuses on `make_mfix`, due to the availability of several flags. Invoking `make_post` is best described by following the example in section 2.3.3.

### 2.3.1.1. `make_mfix` Usage:

#### Basic usage

Invoke `make_mfix` without arguments to go through the list of options. This is the standard input method where the user manually answers questions to compile MFIX. To invoke the makefile, type

```
sh ~/mfix/model/make_mfix
```

or

```
mkmfix
```

if you created the alias (section 2.2). The following examples assume the alias is created.

#### Compiling with default options

Use the `-default` flag to compile with the default options:

- Serial,optimized code
- Re-compilation of source files is not forced
- GNU Fortran compiler

```
mkmfix -default
```

#### Repeating a compilation

Use the `-repeat` flag to recompile with the same options as the last successful compilation.

```
mkmfix -repeat
```

### Showing all compiler options

Some compilers available on very specific systems are hidden by default. Use the -long flag to display all compiler options.

**mkmfix -long**

### Cleaning-up a previous compilation

Use the -clean flag before doing a fresh compilation. This option will remove all .o, .a and .mod files from the object directory (last compilation). The makefile must be invoked again if a new compilation is needed.

**mkmfix -clean**

**mkmfix**

### Showing a help message

Use the -help flag to display a help message that summarizes available options.

**mkmfix -help**

### Shortcuts

Each flag described above has an equivalent short version for convenience:

Description	Flag	Shortcut
Clean-up a previous compilation	-clean	-c
Default options (serial, optimized, GNU)	-default	-d
Display help	-help	-h
Show all compiler options	-long	-l
Repeat compilation with same options as previous	-repeat	-r

For example, to compile with default options, the following two commands are equivalent:

**mkmfix -default**

**mkmfix -d**

### 2.3.1.2. 2013-1 Release changes to make\_mfix

For MFIx users upgrading to the 2013-1 Release, please note the following make\_mfix changes from the previous (2012-1) release:

1. User input in preparation for MFIx compilation was restructured to be more intuitive and several shortcuts were introduced.
2. Due to some unresolved issues with the SMP mode of execution, this option is currently turned off in the 2013-1 Release, and will be re-activated in a future release. This means that MFIx can be compiled in serial and parallel (DMP) modes only.
3. For DMP compilation, the location of the mpif.h file is automatically detected from mpif90 configuration, and no manual user input is required if the file is found.
4. By default, 3 compilers are shown: GNU (gfortran), Portland Group (pgf90), and Intel (ifort). Additional compilers can be selected by invoking the makefile with the -long flag.
5. All object files are now saved in a dedicated folder to speed-up compilation time when switching back and forth between different options.

### 2.3.2. Compiling MFIx for the first time: A step-by-step example

Before compiling MFIx, you must be sure to have a working Fortran compiler installed on your computer (i.e., can compile a simple Fortran code such as print hello and link to create an executable). In the following example, it is assumed that the gfortran compiler is installed on a 64-bit computer. MFIx will be compiled and run for a simple fluidbed tutorial case. At this point, it is not important to understand the simulation setup. The executable file mfix.exe will be created for serial execution.

1) Go to the tutorial directory:

```
cd ~/mfix/tutorial/fluidbed1
```

2) This folder contains the input file mfix.dat that contains the simulation setup. This file is not required to compile MFIx, but will be used at the end of this example to run the simulation. MFIx can be compiled from any directory, except the /mfix/model directory. The directory where MFIx will be run is called the run directory. Here the run directory is



**fluidbed1**, and currently only contains the mfix.dat file. Display the content of the directory:

**ls**

3) Compile MFIx

3a) Invoke the makefile.

**mkmfix**

3b) You will be prompted to answer 3 questions. Press Enter to keep the default answers to all 3 questions. This will compile MFIx in optimized serial mode.

=====

**Mode of execution:**

=====

**[1] Serial**

**[x] Parallel, Shared Memory (SMP) - Temporarily unavailable**

**[3] Parallel, Distributed Memory (DMP)**

**[x] Parallel, Hybrid (SMP+DMP) - Temporarily unavailable**

Select the mode of execution [1] :

← Press Enter

=====

**Level of Optimization:**

=====

**[0] None (Debug mode)**

**[x] Level 1 (not available)**

**[x] Level 2 (not available)**

**[3] Level 3 (most aggressive)**

Select the level of optimization [3] :

← Press Enter



=====  
**Option to re-compile source files in run directory:**  
=====

- [1] Do not force re-compilation**
- [2] Force re-compilation**

**Select Option to re-compile source files in run directory [1] :**      **← Press Enter**

3c) Next, a list of compilers for which the makefile is configured is displayed. It does not mean they are actually all installed on your machine. Choose a compiler that you know is properly installed on your machine. We will select gfortran as an example:

**64 bit Linux system detected, please select compiler.**

=====  
**MFIX Compilation directives available for following compilers:**  
=====

- [1] GNU (gfortran) version 4.3 and above**
- [2] Portland Group (pgf90) version 11.7 and above**
- [3] Intel (ifort) version 11.1 and above**

**Select the compiler to compile MFIx? [1]**      **← Press Enter**

3d) The compilation should start and may take several minutes to complete. Once the compilation is successful, the executable file mfix.exe is created and ready to use.

\*\*\*\*\*

**Compilation successful: mfix.2013-1 created**

**To run MFIx type: mfix.exe**

\*\*\*\*\*

**4) Run MFIx:**

**./mfix.exe**

As the solution proceeds, a lot of information is displayed on the screen. Please see section 4 for a description of the output. The run should complete in a few minutes.

### 2.3.3.Compiling POSTMFIX

You can use POSTMFIX to extract data at a particular location in the flow field. To compile POSTMFIX, follow the following procedure. Again, it is assumed that gfortran is installed.

1) Go to the post\_mfix folder:

```
cd ~/mfix/post_mfix
```

2) Invoke the make file:

```
sh make_post
```

3) A list of compilers will be displayed, based on your operating system. Again, this list means the makefile is configured to use those compilers, but does not imply they are installed on your computer. In our case, we will select the gfortran compiler. Answering no (n) to the first three questions will select gfortran.

**Linux system with 64 bit processor detected, please select compiler**

**MFIX Compilation directives available for following compilers:**

- Intel Fortran Compiler (IFORT - FCE for 64 bit)
- Portland Group Linux Fortran Compiler (pgf90)
- PathScale compiler (pathf90)
- gfortran

Do you want to compile with Intel Compiler? (y/n) [yes] n      ← Type n and press Enter

Do you want to compile with Portland Group Compiler? (y/n) [yes] n      ← Type n and press Enter

Do you want to compile with PathScale Compiler? (y/n) [yes] n      ← Type n and press Enter

**gfortran selected**

4) Before continuing, delete object files:




**Object files (\*.o, \*.mod, \*.a) in mfix/model will be deleted. Continue? (y/n) [no] y** ← **Type y and press Enter**

5) The compilation will start, and upon successful compilation, the following message will be displayed:

```
*****
* Compilation successful: post_mfix created*
*****
```

Once post\_mfix is successfully compiled, the alias created in Section 2.2 is operational.

## 2.4. Visualizing Data with ParaView

It is assumed you have downloaded and installed a suitable binary version of ParaView. To visualize MFIX results, Launch ParaView, and go to File > Open, (or click on the ) and select the .RES file in the run directory. For example, to visualize the results from the fluidbed1 tutorial, open the BUB01.RES file. Next, click on the green “Apply” button  on the left pane to load all variables. This step may be optional on your version of ParaView. You will see a contour plot of the void fraction (EP\_g). To show an animation of void fraction, press the play button in the animation toolbar . You will see bubbles forming along the left boundary and moving upward. This tutorial is using a 2D cylindrical coordinate system, and the left boundary is the axis of symmetry of the system. Only one side was simulated due to symmetry. To visualize the entire system, go to Filter > Alphabetical > Reflect. In the Object inspector, select Plane X, and Center 0, and click Apply. You should see the complete symmetric system now, with bubbles forming in the center.



## 2.5. Starting a New Run

If you have successfully completed your first compilation of MFiX as described in Section 2.3.1, you are ready to explore other options to compile and run MFiX, with your own simulation setup.

Create a separate subdirectory for each run, referred to as the run directory.



If you want to modify MFiX subroutines, do the following. Copy the subroutine from the *mfix/model* directory into the run directory. Modify the file copied to the run directory. Do not modify the files in the *mfix/model* directory.

Create an MFiX executable file by typing **mkmfix**. The MFiX executable file *mfix.exe* will be created and copied into the run-directory.

First, select the type of executable (currently, only serial and DMP options are available). Regardless of the option chosen here, the executable is always named **mfix.exe**

=====

**Mode of execution:**

=====

**[1] Serial**

**[x] Parallel, Shared Memory (SMP) - Temporarily unavailable**

**[3] Parallel, Distributed Memory (DMP)**

**[x] Parallel, Hybrid (SMP+DMP) - Temporarily unavailable**

**Select the mode of execution [1] :**

Type 1 for serial code or 3 for DMP code and press Enter.

Note: To keep the default answer (1 = serial), just press Enter.



If you intend to compile the DMP version of MFIX, please verify that your MPI installation works properly (see [MPI\\_verification\\_for\\_MFIX.pdf](#) document in /mfix/tools/mfi). Contact your system administrator if you are not sure how to proceed.

Next, select the level of optimization:

=====

**Level of Optimization:**

=====

[0] None (Debug mode)

[x] Level 1 (not available)

[x] Level 2 (not available)

[3] Level 3 (most aggressive)

**Select the level of optimization [3] :**

Type 0 and press Enter for executable in debug mode. This option is useful if you modified some source files and want to test the code for possible bugs before production run. Execution time will be much slower in debug mode than in optimized mode.

Press Enter to keep the default answer (3). This will generate an optimized executable, which will run faster for production run.

=====

**Option to re-compile source files in run directory:**

=====

[1] Do not force re-compilation

[2] Force re-compilation

**Select Option to re-compile source files in run directory [1] :**

Type 2 and press Enter to make sure source files in the run directory are compiled. This option is needed only in the rare instances when the run directory contains a module file (e.g., *run\_mod.f*) and another file (e.g., *usr0.f*), in which the corresponding use statement has been inserted (e.g., use run). The makefile will not be aware of this dependency (*usr0* depends upon *run*). So, in this example, changes made in *run\_mod.f* will not cause the (necessary) recompilation of *usr0.f*, unless this option is selected.

Press Enter to keep the default answer (1).

If you keep all the above default options by pressing Enter for each question, an optimized serial version executable will be produced for that particular platform.

Finally, a list of compilers is proposed for your system:

=====

**MFIX Compilation directives available for following compilers:**

=====

- [1] GNU (gfortran) version 4.3 and above**
- [2] Portland Group (pgf90) version 11.7 and above**
- [3] Intel (ifort) version 11.1 and above**

**Select the compiler to compile MFIX? [1]**

Type the number corresponding to the compiler you want to use, and press Enter. If you want to use the default option (1 = GNU gfortran), just press Enter.

After selecting the compiler, the compilation process will start. At the end of the compilation process, you should have the executable **mfix.exe** in this directory.



If you have compiled MFIX once, you can run `make_mfix` with the same options by invoking the makefile with the flag **-repeat** or **-r**. For example

**mkmfix -r**

will repeat the compilation process with the last known configuration. This is useful if you compile MFIX many times during your own development phase, and do not wish to keep entering the same input over and over.

Write an MFIX data file and name it *mfix.dat*. The key words are listed in section 5. Examples are available in the *mfix/tutorials* directory.



You should run several tutorials and carefully study the corresponding *mfix.dat* before attempting to create your own input file. It is strongly recommended to start with an existing *mfix.dat* from a tutorial that closely matches your setup, and modify this file.

### Serial Execution

When MFIX runs, it displays some output on the screen. There are several options to run MFIX with different ways to manage the screen output:

Command	Purpose
<code>./mfix.exe</code>	Standard way to run MFIX. The output is displayed directly on the screen.
<code>./mfix.exe &amp;</code>	Same as above, with MFIX running in the background.
<code>./mfix.exe &gt; screen.log &amp;</code>	Run in batch mode with the screen output redirected to the file <i>screen.log</i> . To follow the screen output at runtime, type <code>tail -f screen.log</code> . Press Ctrl+C to stop following the screen output. (Note that the file <i>screen.log</i> can become very large.).
<code>./mfix.exe  &amp; tee screen.log</code>	Displays the output on the screen and saves it into the <i>screen.log</i> file. <b>This option is available only for csh or tcshrc shell users.</b>

**Note:** you may be able to use *mfix.exe* instead of *./mfix.exe* (i.e. without the dot and slash) on your computer if you have your *\$PATH* environment variable defined properly for current working directory.



## Parallel Execution

Run the DMP version using `mpirun -np <Number of Processors> mfix.exe`. This command may be different on other machines (e.g., `mpprun -n <Number of Processors> mfix.exe`). Recent versions of MPI library (i.e., MPICH2 on Linux clusters) require mpi daemons running on the compute nodes prior to the launch of any mpi executable. Please make sure the appropriate MPI initialization procedures are followed and the simple MPI examples run successfully to verify MPI setup prior to the launch of MFIX executable in DMP mode.

Note that MFIX results can be retrieved, even while the run is in progress.

To retrieve and manipulate data and to create special restart files, run the post\_processor *post\_mfix* by typing `post`. *post\_mfix* will prompt the user for the run name. Enter the run name (e.g., *BUB01*).

## 2.6. Modifying the Post-Processing Codes

To make *post\_mfix*, first change the directory to the *mfix/post\_mfix* directory. If user-defined post-processing is required, modify the *usr\_post.f* file in the directory. Then type `sh make_post`. Note that the *mfix/model* is required for creating a *post\_mfix* executable.

## 3. Setting Up and Running MFIX on Windows Workstation

### 3.1. Installing MFIX on Windows OS

This is not very well supported; please see our download page (<https://mfix.netl.doe.gov/members/download.php>) for special remarks (<https://mfix.netl.doe.gov/members/wininst.html>) and also earlier postings on mfix-help mailing list.

### 3.2. Installing the MFIX Code using Cygwin on Windows OS

1. Install Cygwin from <http://www.cygwin.com/>. To get more familiar with Linux/Unix - you can do a google search on unix primer (<http://www.google.com/search?q=unix++primer>) or you can see the top hit (<http://bignosebird.com/unix.shtml>)
2. Check that Devel and Editors are installed (To the right of the words will be "default"; Left button click on the word "default" until it says "install")
3. Make sure that make and ex are installed



- `$ which ex`  
`/usr/bin/ex`
  - `$ which make`  
`/usr/bin/make`
4. Now install gfortran (this is now part of gcc)
    - Check that gfortran is available (by typing gfortran on console, you should get 'gfortran: no input files')
  5. Download MFIx from our website, and go to the directory where the MFIx tar ball was saved.
  6. issue this command `"tar -xzf mfix.tar.gz"`
  7. go to tutorials/fluidbed1 by typing `"cd mfix/tutorials/fluidbed1"`
  8. Issue the command `"sh ../../model/make_mfix"`
    - This should start the compilation process and depending on the computer this might take a long time.
    - At the end you should have the executable mfix.exe in this directory. To list the files you can use the 'ls' command
  9. Now issue the command `"nohup ./mfix.exe > out1&"` - this will launch the program in the background.
  10. You can see the output from the out file by issuing command `"tail -f out1"`. This would stop as soon as the program finished executing.
  11. You can visualize the output using ParaView (<http://www.paraview.org>).
  12. Any questions check the archives of mfix-help and if you are still having a problem, email [mfix-help@mfix.netl.doe.gov](mailto:mfix-help@mfix.netl.doe.gov) by providing the following important details in your message after the description of the problem encountered:
    1. MFIx version you are trying to install or run
    2. Some details on your operating system environment (for Linux: copy and paste the response of `uname -a` command, Linux distribution name and version also)
    3. Your compiler name and version number (e.g. `ifort -v` will give the version number for Intel fortran compiler)
    4. Output for your \$PATH environment (in csh type `echo $PATH`)
    5. Your MPI library name and version number (if compilations problem with DMP mode encountered but make sure you can compile and run a simple hello world type MPI program with your current installation) Also please provide hardware details such as number of cores per socket in your system (or send the output for `"cat /proc/cpuinfo"` and how many cores you are trying to utilize.



## 4. MFIX at Run Time

### 4.1. MFIX Output and Messages

MFIX output is stored in nine *\*.SPx* binary files. The restart info is periodically written to the *\*.RES* binary file. The text file *\*.OUT* echos the input, shows the numerical cell distribution, and, if OUT\_DT is defined, prints the field variables at the specified intervals.

The text file *\*.LOG* contains run information. For the DMP version, a LOG file is created for each of the processors and they are numbered as *Name###.LOG*.

Messages about the run are written to the *.LOG* file. The progress of the run will be displayed at the terminal as shown below if the data file specifies FULL\_LOG = .TRUE.

Time = 0.92965 Dt = 0.13930E-02 CPU time left = 54.672 s

Nit	P0	P1	U0	V0	U1	V1	Max Res
1	1.5E-02	2.5E+01	1.3E-03	3.3E-03	3.2E-03	6.3E-03	P0
2	1.	5.0E-02	2.8E-03	6.3E-03	2.5E-03	3.8E-03	P1
3	0.1	2.1E-02	1.2E-03	3.7E-03	1.1E-03	1.8E-03	P0
4	5.2E-02	8.1E-03	6.5E-04	2.1E-03	4.8E-04	8.6E-04	P0
5	3.0E-02	3.9E-03	4.1E-04	1.3E-03	2.5E-04	4.3E-04	P0

The first line shows the time, the time-step, and the CPU time remaining to complete the run. The CPU time remaining is not accurate, especially in the beginning of the run. For each time-step, the normalized residuals for various equations are written out every iteration.

MFIX uses a variable time step, which is automatically adjusted within user-defined limits to reduce the run time. At large values of Dt, the iterations may not converge. When this happens, the time step size is successively reduced until convergence is obtained. Messages about divergence and recovery are displayed on the terminal and in the *.LOG* file, if FULL\_LOG = .TRUE.

The subsequent lines display the iteration number, the normalized residuals for various equations (e.g., gas continuity, solids continuity, x and y gas momentum, and x and y solids momentum), and the equation with the maximum residual. The residuals P0 and P1 are normalized only when Nit>1. The residuals displayed can be selected with the



keyword RESID\_STRING.

MFIX reports errors while reading the data file, while processing input data, and during the run time. Errors in reading the data file and in opening files are reported to the terminal. All other errors are reported in the *.LOG* file.

While reporting errors in reading the data file, MFIX displays the offending line of input, so that the error can be easily detected. The possible causes of error are (1) incorrect format for the name-list input, (2) unknown (misspelt) variable name, or (3) the dimension of the name-list item is too small. For example, if the dimension of DX is set as 5000 (DIM\_I in *param\_mod.f*), and if the input data file contains an entry DX(5001), MFIX will report an input processing error.

While processing the input data, MFIX will report errors if the data specified is insufficient or physically unrealistic. MFIX will supply default values only when it is certain that giving a default value is reasonable.

An occasional input-processing error is the inability to determine the flow plane for a boundary condition. The boundary planes defined in the input data file must have a wall-cell on one side and a fluid-cell on the other side. If the initial condition is not specified for the fluid-cell, MFIX will not recognize the cell as a fluid-cell and, hence, MFIX will be unable to determine the flow plane.

Every NLOG number of time steps, MFIX monitors whether the mass fractions add up to 1.0; the overall reaction rates add up to zero; the viscosities, conductivities, and specific heats are greater than zero; and the temperatures are within the specified limits. A message will be printed out if any errors are encountered. The run may be aborted depending upon the severity of the error. Every NLOG time step, MFIX will print out the number of iterations during the previous time step and the total solids inventory in the reactor.

For the specified mass-outflow condition, after the elapse of time BC\_DT\_0, MFIX prints out time-averaged mass flow rates. For cyclic boundary conditions, MFIX will print out the volume averaged mass fluxes every NLOG time step.

A message is written to the *.LOG* file whenever the *.RES* and *.SPx* files are written. This message also shows an approximate value of the cumulative disk space usage in megabytes.





## 4.2. Restarting a Run

A run is restarted by rerunning MFIX after typing `RUN_TYPE = 'restart_1'` in *mfix.dat*. The old *.OUT* file will be overwritten. The *.LOG* messages will be appended to the old *.LOG* file.

## 4.3. When the Run Does Not Converge

Initial non-convergence: Ensure that the initial conditions are physically realistic. If in the initial time step, the run displays NaN (Not-a-Number) for any residual, reduce the initial time step, since automatic time step reduction will become ineffective. If time step reductions do not help, recheck the problem setup.

Holding the time step constant (`DT_FAC=1`) and ignoring the stalling of iterations (`DETECT_STALL=.FALSE.`) may help in overcoming initial nonconvergence. Often a better initial condition will aid convergence. For example, using a hydrostatic rather than a uniform pressure distribution as the initial condition will aid convergence in fluidized-bed simulations.

If there are computational regions where the solids tend to compact (i.e., solids volume fraction less than `EP_star`), the convergence can be improved by reducing `UR_FAC(2)` below the default value of 0.5.

Convergence is often difficult with higher order discretization methods. First order upwinding may be used to overcome initial transients and then the higher order method may be turned on. Also, higher-order methods such as van Leer and minmod give faster convergence than methods such as superbee and ULTRA-QUICK.

## 5. Keywords in Input Data File (mfix.dat)

[ ] indicates the default value.

The symbols used in the table are as follows:

Dimension	Description	Type	Description
1	Cell number in x, y, or z direction	C	Character
M	Solids-phase number	DP	Double Precision
N	Species number	I	Integer
Ic	Initial condition number	L	Logical
Bc	Boundary condition number		
Is	Internal surface number		
Usr	User-defined output number		

### 5.1. Run Control

Keyword (dimension)	Type	Description
RUN_NAME	C	Name used to create output files. The name should be legal after extensions are added to it; e.g., for run name BUB01, the output files BUB01.LOG, BUB01.OUT, BUB01.RES, etc., will be created.
DESCRIPTION	C	Problem description in 60 characters.
UNITS	C	Units for data input and output.
[CGS]		All input and output in CGS units (g, cm, s, cal).
SI		All input and output in SI units (kg, m, s, J).
RUN_TYPE	C	Type of run.
NEW		New run.
RESTART_1		Normal restart run. Initial conditions from .RES file.
RESTART_2		Start a new run with initial conditions from a .RES file created from another run.
RESTART_3		Continue old run as in RESTART_1, but any input data not given in mfix.dat is read from the .RES file. (Do not use)
RESTART_4		Start a new run as in RESTART_2, but any input data not given in mfix.dat is read from the .RES file. (Do not use)
TIME	DP	Start-time of the run.
TSTOP	DP	Stop-time of the run.
DT	DP	Starting time step. If DT is not defined, a steady-state calculation will be performed.
DT_MAX [1.0]	DP	Maximum time step.
DT_MIN [1E-6]	DP	Minimum time step.

<b>DT_FAC</b> [0.9]	DP	Factor for adjusting time step. Should be less than 1. Use a value of 1 to hold the time step constant.
<b>DETECT_STALL</b> [.TRUE.]	L	Reduce time step if the residuals sum does not decrease.
.FALSE.		Do not reduce time step for stalled iterations.
<b>MODEL_B</b> [.FALSE.]	L	Momentum equations.
.TRUE.		Model A
		Model B
<b>MOMENTUM_X_EQ(m)</b> [.TRUE.]	L	(m=0 indicates gas phase) Solve X-momentum equations of phase m.
.FALSE.		Do not solve X-momentum equations of phase m. Beware of inconsistencies when the momentum equations are turned off; e.g., 2-D developing flow with only Y-momentum should not specify no-slip-walls.
<b>MOMENTUM_Y_EQ(m)</b> [.TRUE.]	L	(m=0 indicates gas phase) Solve Y-momentum equations of phase m.
.FALSE.		Do not solve Y-momentum equations of phase m.
<b>MOMENTUM_Z_EQ(m)</b> [.TRUE.]	L	(m=0 indicates gas phase) Solve Z-momentum equations of phase m.
.FALSE.		Do not solve Z-momentum equations of phase m.
<b>ENERGY_EQ</b> [.TRUE.]	L	Solve energy equations
.FALSE.		Do not solve energy equations
<b>SPECIES_EQ(m)</b>	L	(m=0 indicates gas phase)

[.TRUE.]		Solve species equations of phase m. To solve species equation with no chemical reactions, copy the file mfix/model/rrates.f into run directory and remove the first two executable lines as explained in the comments. No other change is needed in that file.
.FALSE.		Do not solve species equations of phase m.
<b>GRANULAR_ENERGY</b> [.FALSE.]	L	Use the granular energy transport equation (pde) as opposed to the algebraic (alg) equation formulation.
<b>SIMONIN</b> [.FALSE.]	L	Use Simonin model (see ~mfix/doc/Simonin_Ahmadi_Models.pdf for details).
<b>AHMADI</b> [.FALSE.]	L	Use Ahmadi model (see ~mfix/doc/Simonin_Ahmadi_Models.pdf for details).
<b>JENKINS</b> [.FALSE.]	L	Use Jenkins small frictional boundary condition (see ~mfix/doc/Simonin_Ahmadi_Models.pdf for details).
<b>FRICTION</b> [.FALSE.]	L	Use the Schaeffer model when .FALSE., or use the Princeton model when .TRUE.

The combination of the keywords GRANULAR\_ENERGY and FRICTION invokes different solids stress models as shown below:

GRANULAR\_ENERGY = .FALSE.

EP\_g < EP\_star > Schaeffer

EP\_g >= EP\_star > viscous (algebraic)

GRANULAR\_ENERGY = .TRUE.

FRICTION = .TRUE.

EP\_s(IJK,M) > EPS\_f\_min > Princeton + viscous (pde)

EP\_s(IJK,M) < EPS\_f\_min > viscous (pde)

FRICTION = .FALSE.

EP\_g < EP\_star > Schaeffer + viscous (pde)

EP\_g >= EP\_star > viscous (pde)

Keyword (dimension)	Type	Description
SAVAGE	I	For a term appearing in the frictional stress model invoked with FRICTION = .TRUE.
0		Use S:S in the frictional stress model.
[1]		Use an alternate form suggested by Savage.
2		An appropriate combination of the above two forms.
SCHAEFFER	L	
[.TRUE.]		If set to false with FRICTION = .FALSE., then the model will not have any frictional viscosity.
BLENDING_STRESS [.FALSE.]	L	This will turn on the blending function to blend the Schaeffer stresses with that of kinetic theory around $\varepsilon^*$ . The default is hyperbolic tangent function for blending (TANH_BLEND = .TRUE.) and one could also utilize a scaled and truncated sigmoidal function (SIGM_BLEND=.TRUE.).
YU_STANDISH	L	
[.FALSE.]		Use Yu and Standish correlation to compute maximum packing for polydisperse systems.
FEDORS_LANDEL	L	
[.FALSE.]		Use Fedors and Landel correlation to compute maximum packing for a binary (only) mixture of powders.
CALL_USR	L	
.TRUE.		Call user-defined subroutines.
[.FALSE.]		Do not call user-defined subroutines.
NRR [0]	I	The number of user defined chemical reactions stored in the *.SPA file. See Section 4.10 Chemical Reactions.

<b>k-ε Equation for Gas-Phase Turbulence</b>		<p>The numerical parameters (like under-relaxation) are the same as the ones for SCALAR (index = 9)</p> <p>Mu_gmax, which is the maximum value of gas turbulent viscosity must be set in order to use the K-Epsilon model. There is no default for this value and the code will not run if Mu_gmax is not set. A value Mu_gmax =1.E+03 is recommended (it is only used in calc_mu_g.f)</p> <p>All walls must be defined (NSW, FSW or PSW) in order to use standard wall functions. If a user does not specify a wall type, he/she will not obtain the typical turbulent profile in wall-bounded flows.</p>
<b>K_Epsilon</b> [.FALSE. ]	L	<p>When activated the k-ε turbulence model (for single-phase flow) is solved using standard wall functions.</p>
<b>KT_TYPE</b> [UNDEFINED]	C	<p>Solids phase stress model. Models include the kinetic theories of Iddir &amp; Arastoopour (AIChE J, 2005): IA_NONEP; Garzo and Dufty (PRE, 1999): GD_99; Garzo, Hrenya and Dufty (PRE, 2007): GHD. To specify a kinetic theory GRANULAR_ENERGY must be .TRUE. If KT_TYPE is left undefined (default) the viscous model is based on the theory of Lun et al. (1984).</p> <p>The IA theory was modified to include the effects of interstitial fluid on two transport coefficients (conductivity and viscosity) as well as a consistency check to ensure that the stresses of two or more identical solids phases will add to that of a single solids phase. These modifications can be turned off by changing the default value of switch_IA to .FALSE. in constant_mod.f. Note that for the IA theory, the definition of granular temperature includes particle mass (unlike the Lun et al. (1984) theory).</p> <p>GHD theory may only be used with the following drag types: "Wen_Yu" and "HYS".</p>

<b>DRAG_TYPE</b> [SYAM_OBRIEN]	C	<p>Drag models include: SYAM_OBRIEN, GIDASPOW, GIDASPOW_PCF, GIDASPOW_BLEND, GIDASPOW_BLEND_PCF, WEN_YU, WEN_YU_PCF, KOCH_HILL, KOCH_HILL_PCF, BVK, HYS. (see drag_gs.f for details)</p> <p>In the model SYAM_OBRIEN two additional parameters may be specified: drag_c1 and drag_d1 (see section 5.2)</p> <p>The extension _PCF following the specified drag model indicates that the polydisperse correction factor proposed by Beetstra et al. (2007) is implemented with that model.</p> <p>In the HYS drag model there is a lubrication cutoff distance that can be specified by the user (see section 4.2). Details of this drag model can be found in ~/mfix/doc/HYS_drag.pdf</p>
<b>RDF_TYPE</b> [LEBOWITZ ]	C	<p>Radial distribution function at contact for polydisperse systems include: LEBOWITZ, MODIFIED_LEBOWITZ, MANSOORI and MODIFIED_MANSOORI.</p> <p>(Do not specify any RDF for monodisperse systems. "Carnahan-Starling" is the only option available.)</p>
<b>Added_Mass</b> [.FALSE. ]	L	<p>When activated the added (or virtual) mass force effectively acts to increase the inertia of the dispersed phase, which tends to stabilize simulations of bubbly gas-liquid flows.</p>



## 5.2. Physical Parameters

Keyword (dimension)	Type	Description
C(100)	DP	User defined constants.
C_NAME(100)	C	Name of user-defined constants (20 characters long). These character strings are used only to identify user-defined constants ( c ) in the .OUT file.
C_e	DP	Coefficient of restitution for particle-particle collisions. (MFIX 1.94 keyword _e_).
e_w [1.0]	DP	Coefficient of restitution for particle-wall collisions.
PHIP [0.6]	DP	Specularity coefficient associated with particle-wall collisions.
PHIP_OUT_JJ [FALSE]	L	Output the variable specularity coefficient when BC_JJ_M is .TRUE.. The specularity coefficient will be stored in ReactionRates array for post-processing by post-mfix. User needs to set nRR to 1 for this purpose. Be careful with this setting when reacting flow is simulated.
PHIP0 [UNDEFINED]	DP	Specify the value of specularity coefficient when the normalized slip velocity goes to zero when BC_JJ_M is .TRUE.. This variable is calculated internally in the code. Do not modify unless an accurate number is known.
C_f	DP	Coefficient of friction between the particles of two solids phases.
Phi	DP	Angle of internal friction (in degrees). Set this value to zero to turn off plastic regime stress calculations.
Phi_w [0.0]	DP	Angle of internal friction (in degrees) at walls. Set this value to non-zero (phi_w = 11.31 means tan_phi_w = Mu = 0.2) when using Jenkins or BC_JJ_M boundary condition.
eps_f_min [0.5]	DP	Minimum solids fraction above which friction sets in. (when FRICTION = .TRUE.)

EP_S_MAX(MMAX) [1.0 - ep_star]	DP	Maximum solids volume fraction at packing for polydisperse systems (more than one solids phase used). The value of EP_star may change during the computation if solids phases with different particle diameters are specified and Yu_Standish or Fedors_Landel correlations are used.
SEGREGATION_SLO PE_COEFFICIENT [0.0]	DP	Used in calculating the initial slope of segregation: see Gera et al. (2004) - recommended value 0.3. Increasing this coefficient results in decrease in segregation of particles in binary mixtures.
L_scale0 [0.0]	DP	Value of turbulent length initialized. This may be overwritten in specific regions with the keyword IC_L_scale.
Mu_gmax	DP	Maximum value of the turbulent viscosity of the fluid.
V_ex	DP	Excluded volume in Boyle-Massoudi stress.
[0.0]		B-M stress is turned off.
P_ref [0.0]	DP	Reference pressure.
P_scale [1.0]	DP	Scale factor for pressure.
GRAVITY	DP	Gravitational acceleration.
[980.7]		By default, the gravity force acts in the _ve y-direction. Modify file b_force2.inc to change the body force term.
Drag_c1 and Drag_d1 drag_c1 [0.8] drag_d1 [2.65]	DP	Quantities for calibrating Syamlal-O'Brien drag correlation using Umf data. These are determined using the Umf spreadsheet ( <a href="http://www.mfix.org/members/develop/umf.xls">http://www.mfix.org/members/develop/umf.xls</a> ). When these are undefined the default values for these constants are used.
USE_DEF_LAM_HYS [.TRUE.]	L	If set to .TRUE. MFIX will use default lubrication cutoff (LAM_HYS) of 1 $\mu$ m as specified in the drag_gs.f file for the HYS polydisperse drag model. If set to .FALSE. the user must specify a lubrication cutoff distance using the LAM_HYS keyword given in the next table entry.

<p><b>LAM_HYS</b></p> <p>[1<math>\mu</math>m]</p>	<p>DP</p>	<p>If USE_DEF_LAM_HYS is set to .FALSE. the user is able to specify a value for the lubrication cutoff distance (LAM_HYS). In practice this number should be on the order of the mean free path of the gas for smooth particles, or the RMS roughness of a particle if they are rough (if particle roughness is larger than the mean free path).</p>
<p><b>M_AM</b></p>	<p>I</p>	<p>Disperse phase number where the added mass applies.</p>

### 5.3. Numerical Parameters

Keyword (dimension)	Type	Description
MAX_NIT [500]	I	Maximum number of iterations.
NORM_g [Use the residual from the first iteration.]	DP	Factor to normalize the gas continuity equation residual.  Setting Norm_g = 0 invokes a normalization method based on the dominant term in the continuity equation. This setting may speed up calculations, especially near a steady state and for incompressible fluids. But the number of pressure iterations may need to be increased, LEQ_IT(1), to ensure mass balance.
NORM_s [Use the residual from the first iteration.]	DP	Factor to normalize the solids continuity equation residual.  Setting Norm_s = 0 invokes a normalization method based on the dominant term in the continuity equation. This setting may speed up calculations, especially near a steady state. But the number of pressure iterations may need to be increased, LEQ_IT(2), to ensure mass balance.
TOL_RESID [1E-3]	DP	Maximum residual at convergence (continuity+momentum).
TOL_RESID_Th [1E-4]	DP	Maximum residual at convergence (granular energy).
TOL_RESID_T [1E-4]	DP	Maximum residual at convergence (energy).
TOL_RESID_X [1E-4]	DP	Maximum residual at convergence (species balance).
TOL_RESID_Scalar [1E-4]	DP	Maximum residual at convergence (scalar balances.)

<b>TOL_DIVERGE</b> [1E+4]	DP	Minimum residual for declaring divergence. When the fluid is incompressible, the velocity residuals take large values in the second iteration (e.g., 1E+8) and then drop down to a low value in the third iteration (e.g., 0.1). In such cases, it is desirable to increase this setting.
<b>Max_Inlet_Vel_Fac</b> [1]	DP	The code declares divergence if the velocity anywhere in the domain exceeds a maximum value. This maximum value is automatically determined from the boundary values. The user may scale the maximum value by adjusting this scale factor.

The next keywords LEQ\_IT, LEQ\_METHOD, LEQ\_SWEEP, LEQ\_TOL, UR\_FAC, and DISCRETIZE are dimensioned for the nine types of equations:

Index	Equation Type
1	gas pressure
2	solids volume fraction
3	gas and solids u-momentum
4	gas and solids v-momentum
5	gas and solids w-momentum
6	Temperature
7	species mass fractions
8	granular temperature
9	user-defined scalar

For example, LEQ\_IT(3) = 10 will make MFX use 10 linear equation iterations while solving the gas and solids u-momentum equation (Equation Type =3).

Keyword (dimension)	Type	Description
<b>LEQ_IT(9)</b>  [20 for 1 and 2] [5 for 2-5] [15 for 6-9]	I	<p>Number of iterations in the linear equation solver. The nine values are for the nine types of equations noted above.</p> <p>The same convention holds for LEQ_METHOD, UR_FAC, and DISCRETIZE. If the residual of an equation is less than the convergence criterion, MFIX makes LEQ_IT equal to the lesser of 5 and the user-defined value and LEQ_METHOD equal to 1.</p>
<b>LEQ_METHOD(9)</b>  [2 for all equations]	I	<p>The method used in the linear equation solver:</p> <ol style="list-style-type: none"> <li>1. SOR</li> <li>2. BiCGSTAB</li> <li>3. GMRES</li> <li>5. CG</li> </ol>
<b>LEQ_SWEEP(9)</b>  [RSRS]	C	<p>The sweep direction used in the linear equation solver. The sweep direction for preconditioning line relaxation; e.g., if LEQ_SWEEP = "ISIS", 1 sweep with do IK loop followed by send_rcv (repeated twice), or if LEQ_SWEEP = "RSRS", 1 red-black sweep with do IK loop followed by send_rcv (repeated twice), or if LEQ_SWEEP = "ASAS", 1 red-black sweep with do IK loop followed by do JK followed by IJ followed by send_rcv (repeated twice). Only used by BiCGSTAB and CG.</p>
<b>LEQ_TOL(9)</b>  [1.0D-4]	DP	<p>The tolerance, if used, in linear equation solvers. Only used by BiCGSTAB and CG.</p>
<b>LEQ_PC(9)</b>  [LINE]	C	<p>The preconditioner used for the sweeps in the linear solver</p> <p>LINE - Line relaxation  DIAG - Diagonal Scaling  NONE - No preconditioner</p>
<b>UR_FAC(9)</b>  [0.8 for 1, 6, 9] [0.5 for 2, 3, 4, 5, 8] [1.0 for 7]	DP	<p>Under relaxation factors for seven types of equations.</p> <p>Reducing UR_FAC(2) will help convergence in problems in which the solids tend to pack.</p>

<b>DEF_COR</b> [.FALSE.]	L	If true, use deferred correction method for implementing higher order discretization. Otherwise, use down-wind factor method (default).
<b>DISCRETIZE(9)</b>	I	Discretization scheme for seven types of equations.
[0]		First-order upwinding.
1		First-order upwinding (using down-wind factors).
2		Superbee (recommended method).
3		SMART.
4		ULTRA-QUICK.
5		QUICKEST (does not work).
6		MUSCL.
7		van Leer.
8		Minmod.
<b>FPFOI</b> [.FALSE.]	L	Four point fourth order interpolation and is upstream biased. If this scheme is chosen and discretize(*) < 2, discretize(*) is defaulted to 2. If you chose this scheme, set the C_FAC value between 0 and 1.
<b>C_FAC</b> [UNDEFINED]	DP	Factor used in the universal limiter (when FPFOI is set .TRUE.) and can be any value in the set (0,1). The choice of 1 will give (diffusion) first order upwinding and as this value becomes closer to 0 the scheme becomes more compressive.
<b>CN_ON</b> [.FALSE.]	L	Implicit Euler based temporal discretization scheme employed (first order accurate in time).
.TRUE.		Crank-Nicholson based temporal discretization scheme employed (second order accurate in time excluding the restart timestep which is first order).

Chi_Scheme	L	Chi-Scheme, proposed by Darwish and Moukalled (2003), is activated. This scheme guarantees that the set of differenced species mass balance equations has the property that the sum of mass fractions add up to one. When a flux limiter is used with (higher order) spatial discretization schemes it is not guaranteed that the mass fractions add up to one. This problem may be rectified by activating the chi-scheme.
[.FALSE.]		
UR_F_gs	DP	The implicitness calculation of the gas-solids drag coefficient may be underrelaxed by changing UR_F_gs, which takes values between 0 to 1: UR_F_gs = 1 (update F_gs every iteration) UR_F_gs = 0 (update F_gs every time step)
[1]		



## 5.4. Geometry and Discretization

For 2D simulations, the thickness of the third direction specified should be exact if mass or volumetric flow rates, rather than velocities, are specified at the boundaries.

Keyword (dimension)	Type	Description
COORDINATES	C	Coordinates used in the simulation.
CARTESIAN		Cartesian coordinates.
CYLINDRICAL		Cylindrical coordinates.
NO_I	L	(Do not use.)
[.FALSE.]		X (r) direction is considered.
.TRUE.		X (r) direction is not considered.
IMAX	I	Number of cells in the x (r) direction.
DX (I)	DP	Cell sizes in the x (r) direction. Enter values from DX(0) to DX(IMAX-1). (Use uniform mesh size with higher-order discretization methods. Also in cylindrical coordinates DX should be kept uniform for strict momentum conservation.)
XMIN	DP	The inner radius in the simulation of an annular cylindrical region.
XLENGTH	DP	Reactor length in the x (r) direction.
NO_J	L	(Do not use.)
[.FALSE]		y direction is considered.
.TRUE.		y direction is not considered.
JMAX	I	Number of cells in the y direction.
DY (1)	DP	Cell sizes in the y direction. Enter values from DY(0) to DY(JMAX-1). (Use uniform mesh size with second-order discretization methods.)
YLENGTH	DP	Reactor length in the y direction.
NO_K	L	
[.FALSE.]		z ( $\theta$ ) direction is considered.

.TRUE.		z ( $\theta$ ) direction is not considered.
KMAX	I	Number of cells in the z ( $\theta$ ) direction.
DZ (1)	DP	Cell sizes in the z ( $\theta$ ) direction. Enter values from DZ(0) to DZ(IMAX-1). (Use uniform mesh size with second-order discretization methods.)
ZLENGTH	DP	Reactor length in the z ( $\theta$ ) direction.
CYCLIC_X	L	Flag for making the x-direction cyclic without pressure drop. No other boundary conditions for the x-direction should be specified.
[.FALSE.]		No cyclic condition at X-boundary.
.TRUE.		Cyclic condition at X-boundary.
CYCLIC_X_PD	L	Flag for making the x-direction cyclic with pressure drop. If the keyword Flux_g is given a value this becomes a cyclic boundary condition with specified mass flux. No other boundary conditions for the x-direction should be specified.
[.FALSE.]		No cyclic condition at X-boundary.
.TRUE.		Cyclic condition with pressure drop at X-boundary.
DELP_X	DP	Fluid pressure drop across XLENGTH when a cyclic boundary condition with pressure drop is imposed in the x-direction.
CYCLIC_Y	L	Flag for making the y-direction cyclic without pressure drop. No other boundary conditions for the y-direction should be specified.
[.FALSE.]		No cyclic condition at Y-boundary.
.TRUE.		Cyclic condition at X-boundary.
CYCLIC_Y_PD	L	Flag for making the y-direction cyclic with pressure drop. If the keyword Flux_g is given a value this becomes a cyclic boundary condition with specified mass flux. No other boundary conditions for the y-direction should be specified.
[.FALSE.]		No cyclic condition at Y-boundary.

.TRUE.		Cyclic condition with pressure drop at Y-boundary.
DELP_Y	DP	Fluid pressure drop across YLENGTH when a cyclic boundary condition with pressure drop is imposed in the y-direction.
CYCLIC_Z	L	Flag for making the z-direction cyclic without pressure drop. No other boundary conditions for the z-direction should be specified.
[.FALSE.]		No cyclic condition at Z-boundary.
.TRUE.		Cyclic condition at Z-boundary.
CYCLIC_Z_PD	L	Flag for making the z-direction cyclic with pressure drop. If the keyword Flux_g is given a value this becomes a cyclic boundary condition with specified mass flux. No other boundary conditions for the z-direction should be specified.
[.FALSE.]		No cyclic condition at Z-boundary.
.TRUE.		Cyclic condition with pressure drop at Z-boundary.
DELP_Z	DP	Fluid pressure drop across ZLENGTH when a cyclic boundary condition with pressure drop is imposed in the z-direction.
SHEAR	L	If .TRUE. imposes a mean shear on the flow field as a linear function of _x_ coordinate. This feature should only be used when CYCLIC_X=.TRUE. Also, the keyword V-sh needs to be set.
V_sh	DP	Specifies the mean _y_ velocity component at the eastern boundary of the domain (V_sh), and the mean _y_ velocity (-V_sh) at the western boundary of the domain.
Flux_g  [UNDEFINED]	DP	If a value is specified (in units of g/cm <sup>2</sup> .s), the domain-averaged gas flux is held constant at that value in simulations over a periodic domain. A pair of boundaries specified as “periodic with fixed pressure drop” is then treated as “periodic with fixed mass flux.” Even for this case a pressure drop must also be specified, which is used as the initial guess in the simulations.

## 5.5. Gas Phase

Keyword (dimension)	Type	Description
RO_g0	DP	Specified constant gas density. This value may be set to zero to make the drag zero and to simulate granular flow in a vacuum. For this case, users may turn off solving for gas momentum equations to accelerate convergence.
MU_g0	DP	Specified constant gas viscosity.
K_g0	DP	Specified constant gas conductivity.
C_pg0	DP	Specified constant gas specific heat.
DIF_g0	DP	Specified constant gas diffusivity.
MW_AVG	DP	Average molecular weight of gas.
MW_g (n)	DP	Molecular weight of gas species n.

## 5.6. Solids Phase

Keyword (dimension)	Type	Description
<b>MMAX</b> [1]	I	Number of solids phases.
<b>D_p0 (m)</b>	DP	Initial particle diameters, same as the old D_P(m).
<b>RO_s (m)</b>	DP	Particle densities.
<b>MU_s0</b>	DP	Specified constant granular viscosity. If this value is specified, then the kinetic theory calculation is turned off and $P_s = 0$ and $\Lambda_s = -2/3 \text{ MU}_s0$ .
<b>K_s0</b>	DP	Specified constant solids conductivity.
<b>C_ps0</b>	DP	Specified constant solids specific heat.
<b>DIF_s0</b>	DP	Specified constant solids diffusivity.
<b>MW_s (m,n)</b>	DP	Molecular weight of solids phase-m, species n.
<b>EP_star</b>	DP	Packed bed void fraction.
<b>CLOSE_PACKED (m)</b> [.TRUE.]	L	Indicates whether the solids phase forms a packed bed with a void fraction EP_star.

## 5.7. Initial Conditions

Each initial condition (IC) is specified over a rectangular region (or pie-shaped for cylindrical coordinates) that corresponds to the scalar numerical grid. These are 3D regions:  $X_w X_e$ ,  $Y_s Y_n$ , and  $Z_t Z_b$ . The region is defined by the constant coordinates of each of the six faces, which may be specified as the physical coordinates or the cell indices. The

physical coordinates are easier to specify than the cell indices. If cell sizes are not small enough to resolve a region specified using physical coordinates, MFIX will indicate this problem with an error message.

In cylindrical coordinates, when the theta direction crosses the 0 value, split that region into two regions: e.g., Split a region spanning 1.9 pi to 0.1 pi as 1.9 pi to 2 pi and 0 to 0.1 pi.

Two initial condition regions may overlap. When an overlap occurs, MFIX uses the conditions specified for the higher IC number.

Keyword (dimension)	Type	Description
IC_X_w (ic)	DP	x coordinate of the west face.
IC_X_e (ic)	DP	x coordinate of the east face.
IC_Y_s (ic)	DP	y coordinate of the south face.
IC_Y_n (ic)	DP	y coordinate of the north face.
IC_Z_b (ic)	DP	z coordinate of the bottom face.
IC_Z_t (ic)	DP	z coordinate of the top face.
IC_I_w (ic)	I	i index of the west-most wall.
IC_I_e (ic)	I	i index of the east-most wall.
IC_J_s (ic)	I	j index of the south-most wall.
IC_J_n (ic)	I	j index of the north-most wall.
IC_K_b (ic)	I	k index of the bottom-most wall.
IC_K_t (ic)	I	k index of the top-most wall.

IC_TYPE (ic)	C	Type of initial condition. Mainly used in restart runs to overwrite values read from the .RES file by specifying it as _PATCH_. The user needs to be careful when using the _PATCH_ option, since the values from the .RES file are overwritten and no error checking is done for the patched values.
IC_EP_g (ic)	DP	Initial void fraction in the IC region.
IC_P_g (ic)	DP	Initial gas pressure in the IC region. If this quantity is not specified, MFIX will set up a hydrostatic pressure profile, which varies only in the y-direction.
IC_P_star (ic)	DP	Initial solids pressure in the IC region. Usually, this value is specified as zero.
IC_L_scale (ic)	DP	Turbulence length scale in the IC region.
IC_ROP_s (ic, m)	DP	Initial bulk density ( $\text{rop}_s = \text{ro}_s \times \text{ep}_s$ ) of solids phase-m in the IC region. Users need to specify this IC only for polydisperse flow ( $\text{MMAX} > 1$ ). Users must make sure that summation of ( $\text{IC\_ROP}_s(\text{ic}, \text{m}) / \text{RO}_s(\text{m})$ ) over all solids phases is equal to ( $1.0 - \text{IC\_EP}_g(\text{ic})$ ).
IC_T_g (ic)	DP	Initial gas phase temperature in the IC region.
IC_T_s (ic, m)	DP	Initial solids phase-m temperature in the IC region.
IC_Theta_m (ic, m)	DP	Initial solids phase-m granular temperature in the IC region.
IC_GAMA_Rg (ic) [0]	DP	Gas phase radiation coefficient in the IC region. Modify file radtn2.inc to change the source term.
IC_T_Rg (ic)	DP	Gas phase radiation temperature in the IC region.
IC_GAMA_Rs (ic, m) [0]	DP	Solids phase-m radiation coefficient in the IC region. Modify file radtn2.inc to change the source term.
IC_T_Rs (ic, m)	DP	Solids phase-m radiation temperature in the IC region.
IC_U_g (ic)	DP	Initial x-component of gas velocity in the IC region.
IC_U_s (ic, m)	DP	Initial x-component of solids-phase velocity in the IC region.

IC_V_g (ic)	DP	Initial y-component of gas velocity in the IC region.
IC_V_s (ic, m)	DP	Initial y-component of solids-phase velocity in the IC region.
IC_W_g (ic)	DP	Initial z-component of gas velocity in the IC region.
IC_W_s (ic, m)	DP	Initial z-component of solids-phase velocity in the IC region
IC_X_g (ic, n) [0]	DP	Initial mass fraction of gas species n.
IC_X_s (ic, m, n) [0]	DP	Initial mass fraction of solids phase-m, species n.
IC_SCALAR (ic, n) [0]	DP	Initial value of Scalar n.



## 5.8. Boundary Conditions

Boundary conditions (BC) are specified over flow planes or 2D surfaces that are normal to one of the coordinate directions and coincide with a face of the scalar control-volume. The values for one of the three pairs of coordinates are equal. The surface is defined by the constant coordinates of each of the four edges, which can be specified with physical coordinates or cell indices, and the two equal values for the direction normal to the face, which can only be specified with physical coordinates. If cell sizes are not small enough to resolve a surface specified using physical coordinates, MFIx will indicate this problem with an error message.

A flow plane must have a wall cell (or an outside boundary) on one side and a flow cell on the other side.

The BC section is also used to specify obstacles in the flow domain. Obstacles are 3D regions, just as for the IC regions:  $X_w$   $X_e$ ,  $Y_s$   $Y_n$ , and  $Z_t$   $Z_b$ . By default the outside boundary is initialized as no-slip walls. For cylindrical coordinates the axis is initialized as a free-slip wall.

Two boundary surfaces must not intersect. Two obstacle regions may intersect.

Keyword (dimension)	Type	Description
BC_X_w (bc)	DP	x coordinate of the west face or edge.
BC_X_e (bc)	DP	x coordinate of the east face or edge.
BC_Y_s (bc)	DP	y coordinate of the south face or edge.
BC_Y_n (bc)	DP	y coordinate of the north face or edge.
BC_Z_b (bc)	DP	z coordinate of the bottom face or edge.
BC_Z_t (bc)	DP	z coordinate of the top face or edge.
BC_I_w (bc)	I	i index of the west-most cell.
BC_I_e (bc)	I	i index of the east-most cell.
BC_J_s (bc)	I	j index of the south-most cell.
BC_J_n (bc)	I	j index of the north-most cell.
BC_K_b (bc)	I	k index of the bottom-most cell.
BC_K_t (bc)	I	k index of the top-most cell.
BC_TYPE (bc)	C	Type of boundary:

DUMMY		The specified boundary condition is ignored. This is useful for turning off some boundary conditions without having to delete them from the file.
MASS_INFLOW or MI		Mass inflow rates for gas and solids phases are specified at the boundary.
MASS_OUTFLOW or MO		The specified values of gas and solids mass outflow rates at the boundary are maintained, approximately. This condition should be used sparingly for minor outflows, when the bulk of the outflow is occurring through other constant pressure outflow boundaries.
P_INFLOW or PI		Inflow from a boundary at a specified constant pressure. To specify as the west, south, or bottom end of the computational region, add a layer of wall cells to the west, south, or bottom of the PI cells. Users need to specify all scalar quantities and velocity components. The specified values of fluid and solids velocities are only used initially as MFIX computes these values at this inlet boundary.
P_OUTFLOW or PO		Outflow to a boundary at a specified constant pressure. To specify as the west, south, or bottom end of the computational region, add a layer of wall cells to the west, south, or bottom of the PO cells.
FREE_SLIP_WALL or FSW		Velocity gradients at the wall vanish. If BC_JJ_PS is equal to 1, the Johnson-Jackson boundary condition is used for solids. A FSW is equivalent to using a PSW with $h_w=0$ .
NO_SLIP_WALL or NSW		All components of the velocity vanish at the wall. If BC_JJ_PS is equal to 1, the Johnson-Jackson boundary condition is used for solids. A NSW is equivalent to using a PSW with $v_w=0$ and $h_w$ undefined.

PAR_SLIP_WALL or PSW		<p>Partial slip at the wall implemented as <math>dv/dn + h_w (v - v_w) = 0</math>, where <math>n</math> is the normal pointing from the fluid into the wall.</p> <p>The coefficients <math>h_w</math> and <math>v_w</math> should be specified. For free slip set <math>h_w = 0</math>. For no slip leave <math>h_w</math> undefined (<math>h_w = \infty</math>) and set <math>v_w = 0</math>. To set <math>h_w = \infty</math>, leave it unspecified.</p> <p>If BC_JJ_PS is equal to 1, the Johnson-Jackson boundary condition is used for solids.</p>
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### Specifications for WALL boundary conditions:

Keyword (dimension)	Type	Description
<b>Momentum Equation</b>		<p>Partial slip at the wall implemented as <math>dv/dn + h_w (v - v_w) = 0</math>, where <math>n</math> is the normal pointing from the fluid into the wall.</p> <p>If the Johnson and Jackson partial slip boundary condition is <u>not</u> used (i.e., BC_JJ_PS(bc) = 0), the coefficients <math>h_w</math> and <math>v_w</math> should be specified.</p> <p>For free slip set <math>h_w = 0</math>. For no slip leave <math>h_w</math> undefined (<math>h_w = \infty</math>) and set <math>v_w = 0</math>. To set <math>h_w = \infty</math>, leave it unspecified.</p>
BC_hw_g (bc) [ $\infty$ ]	DP	Gas phase $h_w$ for partial slip boundary.
BC_hw_s (bc, m) [ $\infty$ ]	DP	Solids phase $h_w$ for partial slip boundary.
BC_Uw_g (bc)	DP	Gas phase $U_w$ for partial slip boundary.
BC_Uw_s (bc, m)	DP	Solids phase $U_w$ for partial slip boundary.
BC_Vw_g (bc)	DP	Gas phase $V_w$ for partial slip boundary.
BC_Vw_s (bc, m)	DP	Solids phase $V_w$ for partial slip boundary.
BC_Ww_g (bc)	DP	Gas phase $W_w$ for partial slip boundary.
BC_Ww_s (bc, m)	DP	Solids phase $W_w$ for partial slip boundary.

BC_JJ_PS (bc)	I	1: Use Johnson and Jackson partial slip bc. 0: Do not use Johnson and Jackson partial slip bc.
[0]		If granular energy transport equation is not solved: (GRANULAR_ENERGY=.FALSE.).
[1]		If granular energy transport equation is solved: (GRANULAR_ENERGY=.TRUE..
BC_JJ_M [.FALSE.]	L	Use the modified Johnson and Jackson partial slip BC with variable specularity coefficient. Must set e_w and phi_w with this BC.
<b>Granular Energy Equation</b>		<p>The granular energy boundary condition is implemented as <math>dT/dn + h_w (T - T_w) = c</math>, where <math>n</math> is the normal pointing from the fluid into the wall.</p> <p>If the Johnson and Jackson partial slip boundary condition is not used (i.e., BC_JJ_PS(bc) = 0), the coefficients <math>h_w</math> and <math>c</math> should be specified.</p> <p>For specified heat flux set <math>h_w=0</math> and give a value for <math>c</math>. For specified temperature boundary condition leave <math>h_w</math> unspecified (<math>h_w=\infty</math> and give a value for <math>T_w</math>.</p>
BC_Thetaw_m (bc, m)	DP	$T_w$ for granular energy bc.
BC_hw_Theta_m (bc, m) [ $\infty$ ]	DP	$H_w$ for granular energy bc.
BC_C_Theta_m (bc, m)	DP	$c$ for granular energy bc.
<b>Gas and Solids Energy Equations</b>		<p>The thermal boundary condition implemented as <math>dT/dn + h_w (T - T_w) = c</math>, where <math>n</math> is the normal pointing from the fluid into the wall.</p> <p>The coefficients <math>h_w</math>, <math>T_w</math>, and <math>c</math> should be specified. <math>H_w = 0 \Rightarrow</math> specified heat flux; <math>h_w = \infty \Rightarrow</math> specified temperature boundary condition. To set <math>h_w = \infty</math>, leave it unspecified and give a value for <math>T_w</math>.</p>
BC_hw_T_g (bc) [ ]	DP	Gas phase $h_w$ for heat transfer.

BC_hw_T_s (bc, m) [ $\infty$ ]	DP	Solids phase $h_w$ for heat transfer.
BC_Tw_g (bc)	DP	Gas phase $T_w$ for heat transfer.
BC_Tw_s (bc, m)	DP	Solids phase $T_w$ for heat transfer.
BC_C_T_g (bc)	DP	Gas phase C for heat transfer.
BC_C_T_s (bc, m)	DP	Solids phase C for heat transfer.
<b>Gas and Solids Species Equations</b>		<p>The species diffusion boundary condition is implemented as <math>dX/dn + h_w (X - X_w) = c</math>, where <math>n</math> is the normal pointing from the fluid into the wall.</p> <p>The coefficients <math>h_w</math>, <math>X_w</math>, and <math>c</math> should be specified. <math>H_w = 0 \Rightarrow</math> specified species diffusion flux; <math>h_w = \infty \Rightarrow</math> specified species concentration at the boundary. To set <math>h_w = \infty</math>, leave it unspecified and give a value for <math>X_w</math>.</p>
BC_hw_X_g (bc, n) [ $\infty$ ]	DP	<p>The species diffusion boundary condition is implemented as <math>dX/dn + h_w (X - X_w) = c</math>, where <math>n</math> is the normal pointing from the fluid into the wall.</p> <p>The coefficients <math>h_w</math>, <math>X_w</math>, and <math>c</math> should be specified. <math>H_w = 0 \Rightarrow</math> specified species diffusion flux; <math>h_w = \infty \Rightarrow</math> specified species concentration at the boundary. To set <math>h_w = \infty</math>, leave it unspecified and give a value for <math>X_w</math>.</p> <p>Gas phase <math>h_w</math> for mass transfer.</p>

BC_hw_X_s (bc, m, n) [ $\infty$ ]	DP	Solids phase $h_w$ for mass transfer.
BC_Xw_g (bc, n)	DP	Gas phase $X_w$ for mass transfer.
BC_Xw_s (bc, m, n)	DP	Solids phase $X_w$ for mass transfer.
BC_C_X_g (bc, n)	DP	Gas phase C for mass transfer.
BC_C_X_s (bc, m, n)	DP	Solids phase C for mass transfer.
<b>Scalar Transport Equations</b>		<p>The scalar boundary condition is implemented as <math>dS/dn + h_w (S - S_w) = C</math>, where <math>n</math> is the normal pointing from the fluid into the wall.</p> <p>The coefficients <math>h_w</math>, <math>S_w</math>, and <math>c</math> should be specified. <math>H_w = 0 \Rightarrow</math> specified species diffusion flux; <math>h_w = \infty \Rightarrow</math> specified species concentration at the boundary. To set <math>h_w = \infty</math>, leave it unspecified and give a value for <math>S_w</math>.</p>
BC_hw_Scalar (bc, n) [ $\infty$ ]	DP	$h_w$ for scalar transfer at the boundary.
BC_ScalarW (bc, n)	DP	$X_w$ for scalar transfer at the boundary.
BC_C_Scalar (bc, n)	DP	C for scalar transfer at the boundary.

### Specifications for FLOW boundary conditions:

Keyword (dimension)	Type	Description
BC_EP_g (bc)	DP	Void fraction at the BC plane.
BC_P_g (bc)	DP	Gas pressure at the BC plane.
BC_ROP_s (bc, m)	DP	Bulk density of solids phase at the BC plane.
BC_T_g (bc)	DP	Gas phase temperature at the BC plane.
BC_T_s (bc, m)	DP	Solids phase-m temperature at the BC plane.
BC_Theta_m (bc, m)	DP	Solids phase-m granular temperature at the BC plane.

BC_X_g (bc, n) [0]	DP	Mass fraction of gas species n at the BC plane.
BC_X_s (bc, m, n) [0]	DP	Mass fraction of solids phase-m, species n at the BC plane.
BC_U_g (bc)	DP	x-component of gas velocity at the BC plane.
BC_U_s (bc, m)	DP	x-component of solids-phase velocity at the BC plane.
BC_V_g (bc)	DP	y-component of gas velocity at the BC plane.
BC_V_s (bc, m)	DP	y-component of solids-phase velocity at the BC plane.
BC_W_g (bc)	DP	z-component of gas velocity at the BC plane.
BC_W_s (bc, m)	DP	z-component of solids-phase velocity at the BC plane.

For a mass inflow boundary, instead of specifying the normal velocity at a boundary, the gas and solids flow rates may be specified as the volumetric or mass flow rates. If the volumetric or mass flow rate is specified, MFIx will calculate the velocity normal to the boundary. The velocity calculated by MFIx, however, may differ from the velocity calculated based on the physical dimensions of the port because the simulated dimensions may not be exactly equal to the physical dimensions. Specify positive values for all the flow rates. MFIx will assign the correct sign to the computed velocity values.

If the mass or volumetric flow rate is specified for a mass outflow boundary condition, then at every interval BC\_DT\_0, MFIx will adjust the normal velocity so that the average computed-outflow rate is equal to the specified value. The user is cautioned, however, that if unrealistic mass flow rates are specified, the computations may become unstable. It is better to specify the velocity at the mass outflow boundary, if some amount of fluctuation in the mass outflow rate is tolerable.

Keyword (dimension)	Type	Description
BC_VOLFLOW_g (bc)	DP	Gas volumetric flow rate through the boundary.
BC_VOLFLOW_s (bc, m)	DP	Solids volumetric flow rate through the boundary.
BC_MASSFLOW_g (bc)	DP	Gas mass flow rate through the boundary.
BC_MASSFLOW_s (bc, m)	DP	Solids mass flow rate through the boundary.

MFIX allows the specification of a transient jet with its velocity fluctuating between two values. The jet conditions will override the steady condition specified for the normal velocity. Therefore, if there is no transient jet, do not specify any of the following, except BC\_DT\_0, which may be required for mass outflow conditions.

Keyword (dimension)	Type	Description
BC_DT_0 (bc)	DP	The interval at the beginning when the normal velocity at the boundary is equal to BC_Jet_g0. When restarting, run this value and BC_Jet_g0 should be specified such that the transient jet continues correctly. MFIX does not store the jet conditions.  For MASS_OUTFLOW boundary conditions, BC_DT_0 is the time period to average and print the outflow rates. The adjustment of velocities to get a specified mass or volumetric flow rate is based on the average outflow rate.
BC_Jet_g0	DP	Value of normal velocity during the initial interval BC_DT_0.
BC_DT_h (bc)	DP	The interval when normal velocity is equal to BC_Jet_gh.
BC_Jet_gh (bc)	DP	Value of normal velocity during the interval BC_DT_h.
BC_DT_1 (bc)	DP	The interval when normal velocity is equal to BC_Jet_g1.
BC_Jet_g1 (bc)	DP	Value of normal velocity during the interval BC_DT_1.



## 5.9. Internal Surfaces

Internal surfaces (IS) are normal to one of the coordinate directions and coincide with one of the faces of the scalar control volume. One of the three pairs of coordinates is equal. The surface is defined by the constant coordinates of each of the four edges, which can be specified with physical coordinates or cell indices, and the two equal values for the direction normal to the face, which can only be specified with physical coordinates. If cell sizes are not small enough to resolve a surface specified using physical coordinates, MFIX will indicate this problem with an error message.

To specify a large number of internal surfaces in a region, a 3D region may be specified. When IS\_Type is specified for such regions, add a prefix (X\_, Y\_, or Z\_) to indicate the direction of the internal surfaces; e.g., X\_IMPERMEABLE specifies impermeable internal surfaces parallel to the X coordinate.

Internal surfaces act as free-slip walls in stress computations. This default condition cannot be changed.

Keyword (dimension)	Type	Description
IS_X_w (is)	DP	x coordinate of the west face or edge.
IS_X_e (is)	DP	x coordinate of the east face or edge.
IS_Y_s (is)	DP	y coordinate of the south face or edge
IS_Y_n (is)	DP	y coordinate of the north face or edge
IS_Z_b (is)	DP	z coordinate of the bottom face or edge
IS_Z_t (is)	DP	z coordinate of the top face or edge
IS_I_w (is)	I	i index of the west-most cell.
IS_I_e (is)	I	i index of the east-most cell
IS_J_s (is)	I	j index of the south-most cell
IS_J_n (is)	I	j index of the north-most cell
IS_K_b (is)	I	k index of the bottom-most cell
IS_K_t (is)	I	k index of the top-most cell
IS_TYPE (is)	C	Type of internal surface:
IMPERMEABLE or IP		No gas or solids flow through the surface.

SEMIPERMEABLE or SP		Gas flows through the surface with an additional resistance. Solids velocity through the surface is set to zero or to a user-specified fixed value (i.e., solids momentum equation for this direction is not solved)
IS_PC (is, 2) (* , 1) = 1.E32 (* , 2) = 0.0	DP	1: permeability; 2: Inertial resistance coefficient. These values need to be specified for semipermeable surfaces only. The thickness used for pressure drop computation is that of the momentum cell (DX_e, DY_n, or DZ_t). To turn off the resistance, use a large value for permeability (1.E32) and a small value for the inertial resistance coefficient (0.0).
IS_VEL_S (is, m) [0.0]	DP	Value of fixed solids velocity through semipermeable surfaces.

## 5.10. Output Control

Keyword (dimension)	Type	Description
RES_DT	DP	Interval at which restart (.RES) file is updated.
SPX_DT (11)	DP	Interval at which .SPX files are updated.
.SP1		Void fraction (EP_g).
.SP2		Gas pressure, solids pressure (P_g, P_star).
.SP3		Gas velocity (U_g, V_g, W_g).
.SP4		Solids velocity (U_s, V_s, W_s).
.SP5		Solids density (ROP_s).
.SP6		Gas and solids temperature (T_g, T_s1, T_s2).
.SP7		Gas and solids mass fractions (X_g, X-s).
.SP8		Granular temperature (G).
.SP9		User defined scalars.
.SPA		Reaction Rates. (See section 4.11)
.SPB		Turbulence quantities (k and $\epsilon$ ).
OUT_DT	DP	Interval at which standard output (.OUT) file is updated.
USR_DT (5)	DP	Interval at which user-defined outputs are written from the subroutine WRITE_USR1.
NLOG [25]	I	Interval in number of time steps at which .LOG file is written.
FULL_LOG [.FALSE.]	L	If true, display the residuals on the screen and messages about convergence on the screen and in the .LOG file.



RESID_STRING (8)	C	<p>Specify residuals to be printed as 4-character strings.</p> <p>First character specifies the field variable: P - pressure, R - density, U - u velocity, V - v velocity, W - w velocity, T - temperature, X - species mass fraction, G - Granular temperature. The second number specifies the phase (0 for gas). The last two numbers specify the species index; e.g., 'P0' - gas pressure, 'R1' - solids phase 1 density, 'X001' - gas phase, species 1; 'X203' - solids phase 2, species 3; 'K0' - k-<math>\epsilon</math> residuals.</p>
<p>REPORT_MASS_BALANCE_DT</p> <p>[Undefined]</p>	DP	<p>If a value is defined, say 0.1 s, an overall species mass balance is performed and reported in the LOG file. The over all mass balance calculations may slightly slow down the run.</p>

## 5.11. Chemical Reactions

Chemical reactions are specified in the data file (`mfix.dat`) by providing species aliases and chemical equations. Rate expressions are specified in one of two user defined subroutines, `usr_rates.f` and `usr_rates_des.f` respectively. Heats of reaction are automatically calculated. Optionally, users may specify constant heats of reaction in the data file.



An overview of using legacy `rrates.f` files is given at the end of this section. However, this input method is no longer directly supported.

There are five general steps to incorporating chemical reactions into a simulation:

1. Provide species names in the data file.
2. Assign a unique identifier (alias) to each species in the data file.
3. Define chemical reaction parameters in the data file.
4. Define chemical reaction rates in `usr_rates.f` and/or `usr_rates_des.f`.
5. Use `make_mfix` to (re)build the MFiX executable.

### 1. Provide species names in the data file with the following keywords.

Keyword (dimension)	Type	Description
<b>NMAX_g</b> [UNDEFINED_I]	I	Number of species comprising the gas phase
<b>SPECIES_g(n)</b> [UNDEFINED_C]	C	Name of gas phase species n as it appears in the materials database
<b>NMAX_s(m)</b> [UNDEFINED_I]	I	Number of species comprising solids phase m
<b>SPECIES_s(m,n)</b> [UNDEFINED_C]	C	Name of solids phase m, species n as it appears in the materials database



Species names must appear **exactly** as given in the materials database (see *Section 5.12, Thermochemical Properties*). Species names are typically 18 characters, and for some species, trailing spaces are needed.

For reacting discrete element simulations (DES), gas phase species are defined using the above keywords (NMAX\_g, Species\_g). However, DES specific keywords are used to identify the number of discrete solids phase species and material database names.

Keyword (dimension)	Type	Description
DES_NMAX_s(m) [UNDEFINED_I]	I	Number of species comprising discrete solids phase m
DES_SPECIES_s(m,n) [UNDEFINED_C]	C	Name of discrete solids phase m, species n as it appears in the materials database

## 2. Assign a *unique identifier* (alias) to each species with the following keywords.

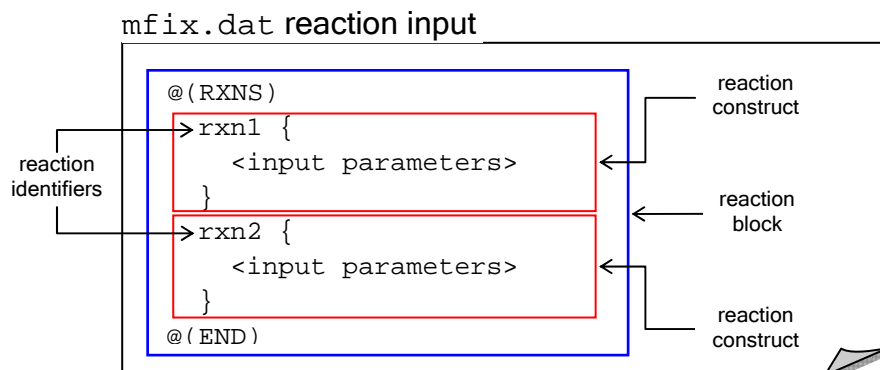
Keyword (dimension)	Type	Description
SPECIES_ALIAS_g(n) [UNDEFINED_C]	C	User defined name for gas phase species n
SPECIES_ALIAS_s(m,n) [UNDEFINED_C]	C	User defined name for solids phase m, species n
DES_SPECIES_ALIAS_s(m,n) [UNDEFINED_C]	C	User defined name for discrete solids phase m, species n

### Alias formatting restrictions:

- Aliases must be unique.
- Aliases are limited to 32 characters and must follow FORTRAN variable naming conventions (i.e., alphanumeric combinations with a letter as the first character).
- Aliases are not case sensitive.
- Aliases cannot conflict with existing MFiX variable names (e.g., a species alias of MU\_g will cause an error when compiling MFiX).

## 3. Define chemical reactions in the data file using species aliases.

Each reaction is identified by a *reaction construct*, and a *reaction block* is used to group reaction constructs in the data file. A reaction construct has the format, rxn\_name{...}, where rxn\_name is a *unique identifier* for the reaction. Reaction identifiers are limited to 32 characters and must follow FORTRAN variable naming convention.





MFIX processes chemical reaction data differently than other input in the data file. A *reaction block* indicates the start and end of the reaction input. A *reaction construct* groups a single reaction's input parameters.

There are two reaction block types:

@ ( RXNS ) ...@ ( END ) - indicates continuum phase chemical reactions (all TFM gas and solids phase reactions and DES homogeneous gas phase reactions).

@ ( DES\_RXNS ) ...@ ( DES\_END ) - indicates heterogeneous DES chemical reactions (particle/gas).



A data file can only contain **one reaction block of each type**, whereas a reaction block must contain **one or more reaction constructs**.

The following keywords are available within a reaction construct.

Keyword (dimension)	Type	Description
CHEM_EQ  [UNDEFINED_C]	C	Chemical equation for the reaction constructed from species aliases. Ex: Char combustion CHEM_EQ = "C + 0.5O2 --> CO"
DH (†Optional) [UNDEFINED]	DP	User provided heat of reaction (cal/ mole for CGS and J/kmole for SI).
fracDH(m) (†Optional) [ZERO]	DP	The fractional amount of DH supplied to phase m.

\* A chemical reaction equation of "NONE" deactivates the reaction during a simulation (e.g., CHEM\_EQ = "NONE").

† By default, heats of reaction are automatically calculated and assigned to the appropriate phase(s). However, users may specify a constant heat of reaction, DH, for one or more reactions to override automated calculations. If DH is given, then fracDH is required. The assigned fractional proportions must sum to one over all phases.

#### Reaction construct formatting notes:

- Chemical reactions are always specified as irreversible with reactants on the left and products on the right. (CHEM\_EQ = "Reactants --> Products")
- An arrow or equals sign can be used to distinguish reactants from products. (Reactants --> Products or Reactants = Products)

- Reversible reactions are specified as two irreversible reactions.  
(see below example, Athermal, gas phase, reversible reaction)
- Chemical equations may span several lines by including an ampersand (&) at the end of the line. As the example below illustrates, each line of the chemical equation is bound in quotation marks and the ampersand is located to the right of the second quotation mark.

```
@(RXNS)                                ! Begin reaction block
  CH4_Combustion {                       ! Reaction 1 construct
    chem_eq = "CH4 + 2O2 --> " &        ! Chemical Reaction Line 1
    "CO2 + 2H2O"                        ! Chemical Reaction Line 2
  }                                       ! End reaction 1 construct
@(END)                                  ! End reaction block
```

- Chemical equations are limited to 512 characters.
- Chemical equations can be bound within single or double quotes.  
(CHEM\_EQ = 'Reactants = Products' or "Reactants = Products")
- Catalytic reactions should contain a species from the catalyst phase in the chemical equation with a coefficient of zero. This insures the proper assignment of the heat of reaction.  
(CHEM\_EQ = 'A + 0.Cat -->3.0\*R' where Cat is a catalyst phase species)
- Catalyst phase species can be listed as a product, reactant, or both.

Several examples illustrating the data file input for several reactions are provided below. Comments are preceded with an exclamation mark (!).

**Example: Methane Combustion:**  $\text{CH}_4(g) + 2\text{O}_2 \rightarrow \text{CO}_2(g) + 2\text{H}_2\text{O}(g)$

Notes: Heat of reaction is automatically calculated (default).

```
NMAX_g = 4                                ! No. of gas phase species
Species_g(1) = "CH4 ANHARMONIC "          ! Methane
Species_g(2) = "O2"                       ! Oxygen
Species_g(3) = "CO2"                      ! Carbon dioxide
Species_g(4) = "H2O"                      ! Water Vapor

Species_Alias_g(1) = "CH4"                ! Methane
Species_Alias_g(2) = "O2"                 ! Oxygen
Species_Alias_g(3) = "CO2"                ! Carbon dioxide
Species_Alias_g(4) = "H2O"                ! Water Vapor

@(RXNS)                                ! Begin reaction block
  CH4_Combustion {                       ! Reaction 1 construct
    chem_eq = "CH4 + 2O2 --> CO2 + 2H2O" ! Chemical Reaction Eq
  }                                       ! End reaction 1 construct
@(END)                                  ! End reaction block
```



**Example: Athermal, gas phase, reversible reaction:**  $A(g) \leftrightarrow R(g)$

Notes: Species database names and aliases are defined on single lines.

The forward and backward reactions are defined separately.

The heats of reaction are defined as zero (athermal) and explicitly assigned to the gas phase.

```
NMAX_g = 2                                ! No. of gas phase species
Species_g(1) = "A" "R"                    ! Database names
Species_Alias_g(1) = "A" "R"              ! Species aliases

@(RXNS)                                    ! Begin reaction block
  fwd_AtoR {                               ! Reaction 1 construct
    chem_eq = "A --> R"                   ! Chemical Reaction Eq
    DH = 0.0                              ! (cal/moles-reacted)
    fracDH(0) = 1.0                       ! Gas phase HoR
  }                                         ! End reaction 1 construct
  rvs_AtoR {                               ! Reaction 2 construct
    chem_eq = "R --> A"                   ! Chemical Reaction Eq
    DH = 0.0                              ! (cal/moles-reacted)
    fracDH(0) = 1.0                       ! Gas phase HoR
  }                                         ! End reaction 2 construct
@(END)                                    ! End reaction block
```

**Example - Char combustion:**  $C(s) + 0.5O_2(g) \rightarrow CO(g)$

Notes: Species database names and aliases are defined on single lines.

The heat of reaction is defined.

The gas phase receives 20% of the heat of reaction.

Solids phase 1 receives 80% of the heat of reaction.

```
NMAX_g = 2                                ! No. gas phase species
Species_g(1) = "O2" "CO"                  ! Database names
Species_Alias_g(1) = "O2" "CO"            ! Species aliases

NMAX_s(1) = 2                              ! No. solids phase species
Species_s(1,1) = "C(GR) REF ELEMENT"      ! Fixed Carbon (graphite)
Species_s(1,2) = "Coal Ash"               ! Coal Ash

Species_Alias_s(1,1) = "C" "Ash"           ! Fixed Carbon and Coal Ash

@(RXNS)                                    ! Begin reaction block
  Char_Combustion {                       ! Reaction 1 construct
    chem_eq = "C + 0.5O2 --> CO"          ! Chemical Reaction Eq
    DH = -52834.0                         ! (cal/moles-reacted)
    fracDH(0) = 0.2                       ! HoR assigned to gas phase
    fracDH(1) = 0.8                       ! HoR assigned to s. phase 1
  }                                         ! End reaction 1 construct
@(END)                                    ! End reaction block
```

**Example - Compound DEM reaction:**

CO combustion:  $\text{CO}(g) + 0.5\text{O}_2(g) \rightarrow \text{CO}_2(g)$

CO<sub>2</sub> gasification:  $\text{C}(s) + \text{CO}_2(g) \rightarrow 2\text{CO}(g)$

Char combustion:  $\text{C}(s) + 0.5\text{O}_2(g) \rightarrow \text{CO}(g)$

Notes: Gas phase species names and aliases are defined on the same line.

DES specific keywords are used to defined DES species data.

Heats of reaction for all reactions are calculated automatically.

A TFM reaction block is used for the gas phase homogeneous reaction.

A DEM reaction block is used for gas/solids reactions.

Reaction constructs are given in one line.

```
! Gas phase species data
NMAX_g = 3
Species_g(1) = "O2"      Species_Alias_g(1) = "O2"
Species_g(2) = "CO"      Species_Alias_g(2) = "CO"
Species_g(3) = "CO2"     Species_Alias_g(3) = "CO2"

! DES solids phase species data
DES_NMAX_s(1) = 2
DES_Species_s(1,1) = "C(GR) REF ELEMENT"
DES_Species_s(1,2) = "Coal Ash"

DES_Species_Alias_s(1,1) = "C"
DES_Species_Alias_s(1,2) = "Ash"

! Homogeneous gas phase reactions
@(RXNS)
  CO_Combustion { chem_eq = "CO + 0.5O2 --> CO2" }
@(END)

! DES Reaction block
@(DES_RXNS)
  CO2_Gasification { chem_eq = "2.0C + O2 --> 2CO" }
  Char_Combustion { chem_eq = "C + CO2 --> 2CO" }
@(DES_END)
```

**Additional comments:**

- Coal Ash is not a species included in the thermochemical database and would require that the properties be given in the data file (see *Section 5.12 Thermochemical properties*).
- One-line reaction constructs are only possible when the heat of reaction is automatically calculated (i.e., the chemical equation is the only input parameter).

#### 4. Define chemical reaction rates in UDF files (`usr_rates.f` and `usr_rates_des.f`).

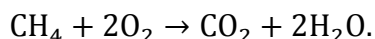
A reaction rate should be given in either `usr_rates.f` or `usr_rates_des.f` for each reaction listed in the data file.

- All TFM gas and solids phase reactions as well as homogeneous gas phase reactions for DEM simulations are to be included in `usr_rates.f`. Reaction rates defined in `usr_rates.f` must have units of reacted moles per time per volume (i.e., moles/sec/cm<sup>3</sup> for CGS units and kmols/sec/m<sup>3</sup> for SI units).
- All discrete phase heterogeneous (particle/gas) reactions are to be included in `usr_rates_des.f` located in the `des` subfolder. Reaction rates defined in `usr_rates_des.f` must have units of reacted moles per time (i.e., moles/sec).



Formation and consumption rates are automatically calculated for each species from the reaction rate and chemical equation.

The rate in terms of reacted moles is related to the rates of formation and consumption through the stoichiometric coefficients. For example, consider homogeneous gas phase reaction of methane combustion:



The rate in terms of reacted moles, *Rate*, is related to the rates of formation and consumption as

$$\begin{aligned} \text{Rate} &= \frac{-r_{\text{CH}_4}}{1} \left( \frac{\text{mol}_{\text{CH}_4}/(\text{s} \cdot \text{cm}^3)}{\text{mol}_{\text{CH}_4}} \right) = \frac{-r_{\text{O}_2}}{2} \left( \frac{\text{mol}_{\text{O}_2}/(\text{s} \cdot \text{cm}^3)}{\text{mol}_{\text{O}_2}} \right) \\ &= \frac{r_{\text{CO}_2}}{1} \left( \frac{\text{mol}_{\text{CO}_2}/(\text{s} \cdot \text{cm}^3)}{\text{mol}_{\text{CO}_2}} \right) = \frac{r_{\text{H}_2\text{O}}}{2} \left( \frac{\text{mol}_{\text{H}_2\text{O}}/(\text{s} \cdot \text{cm}^3)}{\text{mol}_{\text{H}_2\text{O}}} \right), \end{aligned}$$

where  $-r_{\text{CH}_4}$  and  $-r_{\text{O}_2}$  are the rates of consumption of methane and oxygen, and  $r_{\text{CO}_2}$  and  $r_{\text{H}_2\text{O}}$  are the rates of formation of carbon dioxide and water vapor, respectively.

Each reaction rate is assigned to the variable `RATES(rxn_name)`, where `rxn_name` is the reaction identifier used in the reaction construct. To minimize input errors when specifying reaction rates, species aliases (`SPECIES_ALIAS`) defined in the data file should be used in lieu of the associated species index.

For example, if oxygen is defined as gas phase species 2 with an alias of "O2", (e.g., `SPECIES_ALIAS_g(2) = "O2"`), when accessing gas phase species data for oxygen (e.g., molecular weight; `MW_g`), "O2" should be used and not the integer index 2, (e.g., `MW_g(O2)`).

A similar procedure is used for DES reactions with the exception that the reaction rate is assigned to the variable `DES_RATES(rxn_name)`, where `rxn_name` is the reaction identifier used in the reaction construct.

**Example: Methane Combustion:**  $\text{CH}_4(g) + 2\text{O}_2 \rightarrow \text{CO}_2(g) + 2\text{H}_2\text{O}(g)$

Notes: Species database names and alias are defined on the same line.

The fluid cell index (IJK) is passed as a dummy argument.

Global field variables are referenced (RO\_g, X\_g, T\_g, and EP\_g )

Species aliases (O2 and CH4) are used instead of the species indices.

Reaction identifier (CH4\_Combustion) is used in the rates array.

Reaction rate is stored for post processing (see below).

data file input:

```
NMAX_g = 4
Species_g(1) = "CH4 ANHARMONIC " Species_Alias_g(1) = "CH4"
Species_g(2) = "O2" Species_Alias_g(2) = "O2"
Species_g(3) = "CO2" Species_Alias_g(3) = "CO2"
Species_g(4) = "H2O" Species_Alias_g(4) = "H2O"

@(RXNS)
  CH4_Combustion { chem_eq = "CH4 + 2O2 --> CO2 + 2H2O" }
@(END)
```

usr\_rates.f input:

```
SUBROUTINE USR_RATES(IJK, RATES)
  DOUBLE PRECISION, INTENT(IN) :: IJK          ! Fluid Cell Index
  DOUBLE PRECISION, INTENT(OUT) :: RATES(:)    ! Reaction Rates
  :
  DOUBLE PRECISION c_O2 ! Oxygen concentration (mol/cm^3)
  DOUBLE PRECISION c_CH4 ! Methane concentration (mol/cm^3)

  ! Calculate species concentrations:
  c_O2 = (RO_g(IJK) * X_g(IJK,O2))/MW_g(O2)
  c_CH4 = (RO_g(IJK) * X_g(IJK,CH4))/MW_g(CH4)

  ! Methane Combustion
  ! CH4 + 2O2 --> CO2 + 2H2O (reacted moles/sec.cm^3)
  !-----//
  RATES(CH4_Combustion) = 6.7d12 * exp(-2.4358d4/T_g(IJK)) * &
    EP_g(IJK) * (c_O2**1.3) * (c_CH4**0.2)

  ! Store the reaction rate for output/post processing.
  IF(CH4_Combustion <= NRR) &
    ReactionRates(IJK, CH4_Combustion) = RATES(CH4_Combustion)

END SUBROUTINE USR_RATES
```

**Example: Athermal, gas phase, reversible reaction:  $A(g) \leftrightarrow R(g)$**

Notes: Species database names and alias are defined on the same line.

The fluid cell index (IJK) is passed as a dummy argument.

Global field variables are referenced (RO\_g, X\_g, T\_g, and EP\_g )

data file input:

```
NMAX_g = 2                                ! No. of gas phase species
Species_g(1) = "A" "R"                    ! Database names
Species_Alias_g(1) = "A" "R"              ! Species Aliases

@(RXNS)                                   ! Begin reaction block
  fwd_AtoR {                               ! Reaction 1 construct
    chem_eq = "A --> R"                   ! Chemical Reaction Eq
    DH = 0.0                               ! (cal/moles-reacted)
    fracDH(0) = 1.0                       ! Gas phase HoR
  }                                         ! End reaction 1 construct
  rvs_AtoR {                               ! Reaction 2 construct
    chem_eq = "R --> A"                   ! Chemical Reaction Eq
    DH = 0.0                               ! (cal/moles-reacted)
    fracDH(0) = 1.0                       ! Gas phase HoR
  }                                         ! End reaction 2 construct
@(END)                                    ! End reaction block
```

usr\_rates.f input:

```
SUBROUTINE USR_RATES(IJK, RATES)
  DOUBLE PRECISION, INTENT(IN) :: IJK      ! Fluid Cell Index
  DOUBLE PRECISION, INTENT(OUT) :: RATES(:) ! Reaction Rates
  :
  DOUBLE PRECISION c_A ! species A concentration (mol/cm^3)
  DOUBLE PRECISION c_R ! species R concentration (mol/cm^3)

  ! Calculate species concentrations:
  c_A = (RO_g(IJK) * X_g(IJK,A))/MW_g(A)
  c_R = (RO_g(IJK) * X_g(IJK,R))/MW_g(R)

  ! Forward Reaction
  ! A --> R (reacted moles/sec.cm^3)
  !-----//
  RATES(fwd_AtoR) = 1.2d17 * exp(-5.837d3/T_g(IJK)) * &
    EP_g(IJK) * c_A

  ! Reverse Reaction
  ! R --> A (reacted moles/sec.cm^3)
  !-----//
  RATES(rvs_AtoR) = 2.5d41 * exp(-1.4897d4/T_g(IJK)) * &
    EP_g(IJK) * c_R

END SUBROUTINE USR_RATES
```

**Example - Char combustion:**  $C(s) + 0.5O_2(g) \rightarrow CO(g)$ 

Notes: A representative data file input was presented previously.

The fluid cell index (IJK) is passed as a dummy argument.

Algebraic expressions for the rate limiting steps are omitted for brevity.

```

SUBROUTINE USR_RATES(IJK, RATES)
  DOUBLE PRECISION, INTENT(IN) :: IJK          ! Fluid Cell Index
  DOUBLE PRECISION, INTENT(OUT) :: RATES(:)    ! Reaction Rates
  :
! Rate limiting steps:
  DOUBLE PRECISION k_f      ! film diffusion          (cm/sec)
  DOUBLE PRECISION k_a      ! ash layer diffusions    (cm/sec)
  DOUBLE PRECISION k_s      ! chemical kinetics       (cm/sec)
  DOUBLE PRECISION k_eff    ! effective rate          (cm/sec)

! Total surface area of solids phase 1 in IJK
  DOUBLE PRECISION Sa      ! (cm^2/cm^3)

! C + 0.5O2 --> CO (reacted moles/sec.cm^3)
!-----//
! Verify that solids are present
  IF(.NOT.COMPARE(EP_g(IJK),ONE)) THEN
! Calculate film diffusion rate
    k_f = < film diffusion rate expression >      ! (cm/sec)
! Calculate ash diffusion rate
    k_a = < ash diffusion rate expression >        ! (cm/sec)
! Calculate kinetic rate rate
    k_s = < kinetic rate expression >              ! (cm/sec)

! Effective rate (cm/sec)
    k_eff = ONE/(ONE/k_a + ONE/k_f + ONE/k_s)

! Calculate total surface area of solids phase 1
    Sa = 6.0 * EP_s(IJK,1) / D_p0(1)
! Calculate the reaction rate.
    RATES(Char_Combustion) = 2.0 * (Sa * k_eff * Conc(O2))

  ELSE
! No solids --> No reaction
    RATES(Char_Combustion) = ZERO
  ENDIF
END SUBROUTINE USR_RATES

```

See `mfix/model/tutorial/SpoutedBedCombustor` for details on a similar simulation setup.

### Example - DES droplet evaporation: $H_2O(l) \rightarrow H_2O(g)$

Notes: Various algebraic expressions in the sample UDF are omitted for brevity.  
The global particle index (NP), phase index (pM), and fluid cell index (IJK) are passed as dummy arguments.

data file input:

```
NMAX_g = 2                                ! No. of gas phase species
Species_g(1) = "Air" "H2O"                ! Database names
Species_Alias_g(1) = "Air" "Vapor"         ! Species Aliases

DES_NMAX_s(1) = 1                          ! No. of DEM solids phase species
DES_Species_s(1,1) = "H2O(L)"              ! Database names
DES_Species_Alias_s(1,1) = "Liquid"        ! Species Aliases

@(DES_RXNS)
  Evap { Liquid --> Vapor }
@(DES_END)
```

usr\_rates\_des.f input:

```
SUBROUTINE USR_RATES_DES(NP, pM, IJK, DES_RATES)
  DOUBLE PRECISION, INTENT(IN) :: NP      ! Global particle index
  DOUBLE PRECISION, INTENT(IN) :: pM      ! Particle solid phase
  DOUBLE PRECISION, INTENT(IN) :: IJK     ! Fluid Cell Index
  DOUBLE PRECISION, INTENT(OUT) :: DES_RATES(:) ! Reaction Rates
  :
  ! Liquid --> Vapor (reacted moles/sec)
  !-----//
  ! Calculate the concentration gradient (mole/cm^3)
  Cmg_H2O = < expression for calculating gradient >

  IF(Cmg_H2O > ZERO) THEN
    ! Calculate mass transfer coefficient (cm/sec)
    H2O_xfr = < mass transfer coeff calculation >
    ! Calculate droplet surface area (cm^3)
    Sa = Pi * 4.0d0 * (DES_RADIUS(NP)**2)
    ! Calculate the mass transfer rate (moles/sec)
    DES_RATES(Evap) = Sa * H2O_xfr * Cmg_H2O
  ELSE
    DES_RATES(Evap) = ZERO
  ENDIF

  ! Store the reaction rate for post processing.
  IF(Evap <= NRR) ReactionRates(Evap) = &
    ReactionRates(IJK, Evap) + DES_RATES(Evap)

END SUBROUTINE USR_RATES_DES
```

See mfix/tests/dem-tests/evaporation for additional details.

## 5. Use `make_mfix` to (re)build the MFiX executable.

Detailed instructions on building the MFiX executable are given in Section 2.3.1. `make_mfix` should be ran to update the `mfix.exe` executable if any one of the following modifications is made:

- the number, order, or alias of any species is changed in the data file.
- the number, order, or name of any chemical reaction is changed in the data file .
- any change is made to chemical reaction rates in either `usr_rates.f` or `usr_rates_des.f`.



`make_mfix` preprocesses the data file to generate the `species.inc` file which is included within the `usr_rates.f` and `usr_rates_des.f` files as code. Therefore changes in the data file may result in the executable being out of date.

### Additional reaction information::

#### To write out reaction rates to SPx file:

1. In the data file, `mfix.dat`, set NRR to the desired number of reaction rates to be written out to the file `*.SPA`. This number is typically less than or equal to the total number of reactions.
2. In a reaction UDF (`usr_rates.f` or `usr_rates_des.f`) assign the desired reaction information to the variable `ReactionRates`. `ReactionRates` is a two-dimensional array. The first index references the fluid cell, `IJK`, while the second index ranges from 1 to NRR.



If the second index exceeds NRR, a run time error can result from over indexing the array. Using logical checks can eliminate potential errors!

Two of the above examples illustrate using the `ReactionRates` variable:

1. *Methane Combustion*: The calculated reaction rate is directly stored, and logical check is used to prevent over indexing the `ReactionRates` array.
2. *DES droplet evaporation*: The calculated reaction rate is added to the storage array. Adding the calculated data to the storage variable is needed in DES since several discrete particles may exist in a single fluid cell. Again, a logical check is preformed to prevent over indexing the array.



### Using an existing (legacy) `rrates.f` file:

The legacy `rrates.f` file should be copied to the run directory. Additionally, the following keywords should be specified in the data file:

Keyword (dimension)	Type	Description
<b>USE_RRATES</b> [.FALSE]	L	Use the automated reaction rate UDFS; <code>usr_rates.f</code> and <code>usr_rates_des.f</code> .
.TRUE.		Access legacy <code>rrates.f</code> file for reaction rate information. Rates of formation/consumption as well as heats of reaction are provided by the user.
<b>NMAX(m)</b> [UNDEFINED_I]	I	Number of species in phase m. Note that the gas phase is indicated as m=0.
<b>SPECIES_NAME(n)</b> [UNDEFINED_C]	C	Names of gas and solids phase species as it appears in the materials database. The first NMAX(0) are the names of gas species. The next NMAX(1) are the names of solids phase-1 species, etc.



Legacy species keywords , NMAX(m) and SPECIES\_NAME(n) , are **required** when using a legacy `rrates.f` file. Current species keywords NMAX\_g, NMAX\_s, SPECIES\_g, and SPECIES\_s **cannot** be used.



The only modification needed for a legacy `mfix.dat` and `rrates.f` file combination is the inclusion of `USE_RRATES=.TRUE.` in the data file.  
An example of legacy file usage: `mfix/tutorials/reactor1b`

### Additional remarks:

- `make_mfix` requires that the data file, `mfix.dat`, be present in the run directory as the species aliases and reaction identifiers are needed to construct a `species.inc` file.
- Species aliases and reaction identifiers must be unique. `make_mfix` performs a cursory check on the supplied data and exits if non unique entries are identified.
- If any species alias or reaction identifier conflicts with an existing global variable in MFiX, an error will be reported and the code will fail to compile.

## 5.12. Thermochemical Properties

The directory `mfix/model/thermochemical` contains the database of Burcat and Ruscic (2005) and routines for reading the database. With linkage to this database the users need not manually enter data for molecular weight, specific heat, and heats of reactions. Instead the users need to enter the names of the species (keyword `SPECIES_g` and `SPECIES_s`) in the data file. MFIx reads the necessary thermochemical data from files in the following order:

1. `mfix.dat`
2. `BURCAT.THR` file in the run directory
3. `mfix/model/thermochemical/BURCAT.THR`.

The species names are case sensitive and should match the names in `BURCAT.THR` exactly; alternatively aliases can be defined for common species, such as `O2`, in `read_therm.f`. See `mfix/tests/thermo` for a sample case that accesses the database. The format of `BURCAT.THR` file resembles `CHEMKIN` format, but with several notable differences. Thermochemical data must start below a line that starts with `THERMO DATA`.

Example dataset from `BURCAT.THR` with notations:

CAS identifier		valid temperature range				molecular weight						
74-82-8												
comments	CH4 METHANE Same as the Anharmonic but calculated Using the RRHO method rather than the NRRAO2. Max Lst Sq Error Cp @ 6000. K 0.62%.											
	CH4	RRHO	g 8/99C	1.H	4.	0.	0.G	200.000	6000.000	B	16.04246	1
			1.91178600E+00	9.60267960E-03	-3.38387841E-06	5.38797240E-10	-3.19306807E-14					2
			-1.00992136E+04	8.48241861E+00	5.14825732E+00	-1.37002410E-02	4.93749414E-05					3
			-4.91952339E-08	1.70097299E-11	-1.02453222E+04	-4.63322726E+00	-8.97226656E+03					4
species name		high temperature coefficients		low temperature coefficients		formation enthalpy at 298K						

Each entry in the database starts with a unique CAS identifier (74-82-8) for the species, followed by several lines of comments highlighted in green. The data section starts with the species name in columns 1-18 (`CH4 RRHO`). Common species names may be followed by strings (`RRHO`) that identify the method used to determine the coefficients. Additional information follows the species name. The numbers toward the end of the line are the temperature limits (200.000 6000.000) in degrees Kelvin where the property calculation is valid and the molecular weight (16.04246). Unlike `CHEMKIN` the common temperature for the high and low temperature branches are not recorded; it is always 1000 K. The next three lines give the fourteen coefficients (seven coefficients each for the high and low temperature branches) and the formation enthalpy

at 298 K (which is also not included in CHEMKIN format). All the coefficients and the enthalpy of formation are normalized with the gas constant  $R$  (cal/mol/K). The low temperature coefficients ( $a_L$ ) should be used for temperatures in the range  $T_{\text{low}}$  to 1000K and the high temperature coefficients ( $a_H$ ) should be used for temperatures in the range 1000K to  $T_{\text{high}}$ . The coefficients are stored in a fixed format (E15.0) as follows:

$a_H^1$	$a_H^2$	$a_H^3$	$a_H^4$	$a_H^5$
$a_H^6$	$a_H^7$	$a_L^1$	$a_L^2$	$a_L^3$
$a_L^4$	$a_L^5$	$a_L^6$	$a_L^7$	$\Delta H_f^\circ$

where  $\Delta H_f^\circ$  is the formation enthalpy at 298K.

The normalized specific heat is given by

$$C_p/R = a_1 + a_2T + a_3T^2 + a_4T^3 + a_5T^4.$$

#### Additional database comments:

- A number of species in the database have a lower temperature limit of 300K which is 2 degrees above the reference temperature (298 K) used for formation enthalpy calculation. For those species MFIx relaxes the lower limit for  $C_p$  calculations to 298 K to enable heat of reaction calculation.  
see `read_database.f`
- The database reader is set up such that the database is read only if necessary.
- For additional details see the Burcat and Ruscic (2005) report located in the thermo-chemical subdirectory, `mfix/model/thermochemical/intro.pdf`.

### 5.13. User-Defined Subroutines

The user may modify any \*.f or \*.inc file in MFIX. To modify a file, first copy it from the mfix/model directory into the run directory. Modify only this copy in the run directory; do **NOT** modify the original files in mfix/model. Then, invoke the 'sh make\_mfix' command from the run directory. The make\_mfix messages will identify the files from the run directory used to create the MFIX executable. All the (MFIX and non-MFIX) \*.inc files from the run directory will be used to create the MFIX executable. Only MFIX \*.f files from the run directory will be used, however. Non-MFIX \*.f files in the run directory will be ignored. To use new Fortran files, include them in one of the MFIX \*.f files.

The following is a list of MFIX files that are usually modified to include chemical reactions and user defined scalars:

usr_rates.f	Chemical reaction rates.
transport_prop.f	Transport properties.
physical_prop.f	Physical properties.
scalar_prop.f	Properties and source terms in scalar transport equations.
usr_rates_des.f <sup>[1]</sup>	DES chemical reaction rates.

The following routines are used for writing user-defined output:

write_usr0.f	Called once during the run. Can be used for opening user-defined files.
write_usr1.f	Called at intervals defined by USR_DT.

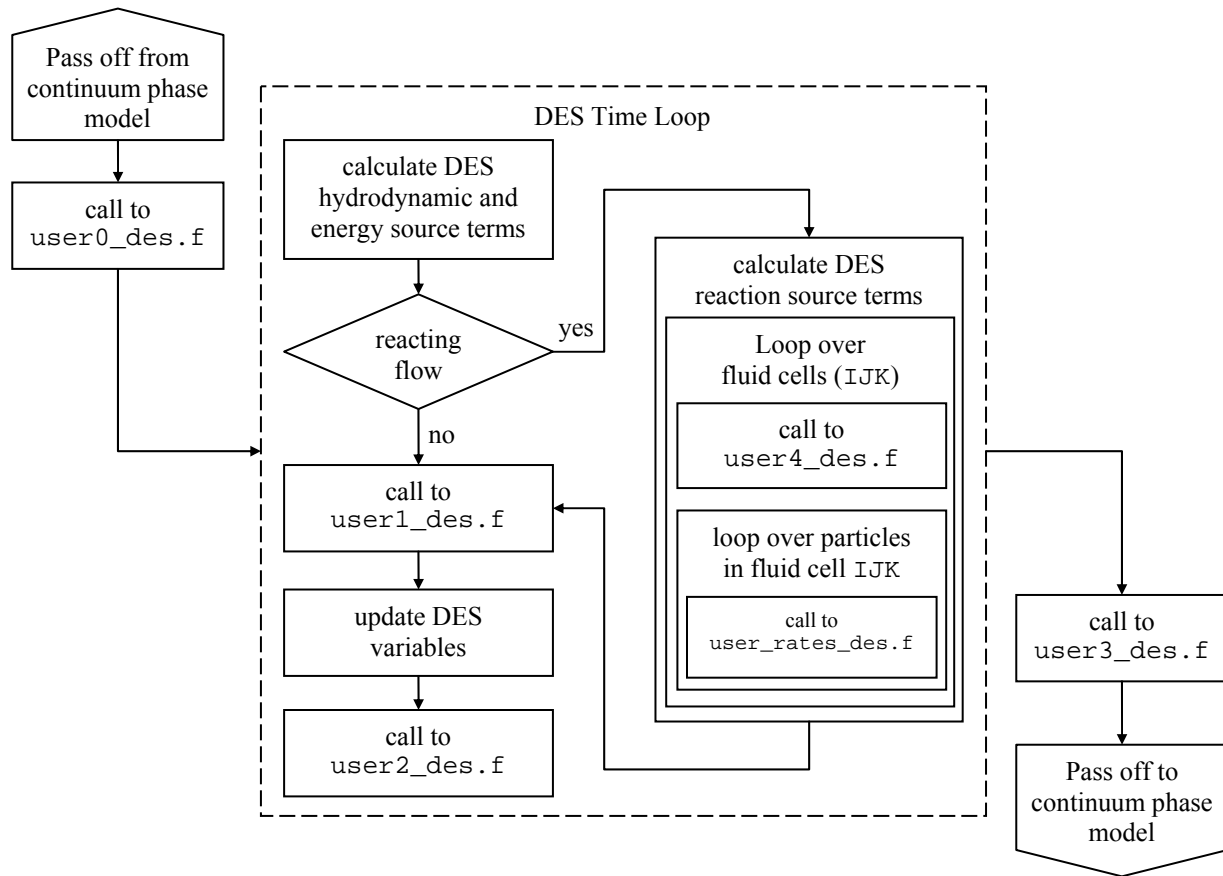
To activate the calls to the following three routines, set call\_usr = .TRUE. in the data file:

usr0.f	A subroutine that is called once every run, just before the time-loop begins.
usr1.f	A subroutine that is called once every timestep.
usr2.f	A subroutine that is called once every iteration.
usr3.f	A subroutine that is called once every run, after the time-loop ends.
usrnlst.inc	List of user-defined keywords. These may be used to enter data through the input data file mfix.dat.

<code>usr_init_namelist.f</code>	Initialize user-defined keywords.
<code>usr_mod.f</code>	User-defined module. Include "Use usr" to use user-defined variables in this module. If allocatable arrays are defined in this module, allocate them in <code>usr0.f</code> .
<code>usr0_des.f</code> <sup>[1]</sup>	A subroutine called before entering the DES time loop.
<code>usr1_des.f</code> <sup>[1]</sup>	A subroutine called every DEM timestep after calculating DES source terms but before source terms are applied to the particles.
<code>usr2_des.f</code> <sup>[1]</sup>	A subroutine called every DES timestep after source terms are applied to the particles.
<code>usr3_des.f</code> <sup>[1]</sup>	A subroutine that is called after completing the DES time loop.
<code>usr4_des.f</code> <sup>[1]</sup>	This subroutine is called from and IJK loop before calculating DEM reaction rates. Used for calculating values that are constant within a fluid cell needed for reaction calculations (e.g., Schmidt Number).

<sup>[1]</sup> Denotes files contained in the DES subfolder (`mfix/model/des/`).

DES User-defined subroutine call structure:



### 5.14. Parallelization Controls

Keyword (dimension)	Type	Description
<b>NODESI</b> [1]	I	Number of grid blocks in x-direction.
<b>NODESJ</b> [1]	I	Number of grid blocks in y-direction.
<b>NODESK</b> [1]	I	Number of grid blocks in z-direction.
<b>MINIMIZE_DOTPRODUCTS</b> [FALSE]	L	The dot products calculations in BiCGStab are minimized and this should only have impact on parallel runs
<b>SOLVER_STATISTICS</b> [FALSE]	L	Print out additional statistics for parallel runs
<b>DEBUG_RESID</b> [TRUE]	L	If set to false, the residuals are grouped into far fewer global collectives and this should only have impact on parallel runs

NODESI \* NODESJ \* NODESK must be the same as the number of processors specified using the mpirun (or equivalent command). Otherwise the code will return with an error.

The parallel performance depends on several things and one has to evaluate different options before choosing the right strategy for the problem at hand. For e.g. if the J direction is the strongest coupled direction, the preconditioning for the linear solver will be poor if there is decomposition in that direction. However, since decomposing in all the directions reduces the processor grid surface area to the volume, the communication cost will be less for the same computational grid. The preconditioners are chosen with the keyword LEQ\_PC. In addition to LINE relaxation, one can choose the "DIAG" or "NONE" preconditioners that reduces interprocessor communications but would increase the number of linear equation solver iterations. The DIAG and NONE choices for preconditioners may be appropriate for all equations except the continuity (or pressure and volume fraction correction) equations. The parallel performance is greatly dependent on the choices stated here, and some trial and error may be required to determine the right combination of decomposition direction and the choice of preconditioners to get the best performance in production runs.

## 5.15. MFIx Execution in Batch Queue Environment

MFIx can be used on systems where code execution is controlled through batch queue submission system instead of interactive or background job type methods shown in the previous section. Usually the user specifies the wall clock time duration of the job and batch queuing system prioritize incoming jobs based on their resource allocation requests. In order for MFIx to terminate cleanly at the end of the batch job session, several keywords need to be entered in `mfix.dat`. Clean termination in environments with batch queue is important as the system may terminate the batch job while MFIx is writing out `*.SP` files, which may corrupt the files or cause loss of data.

For this purpose, MFIx checks if termination criteria specified by user is reached or not at the beginning of each time step. However, to avoid performance bottlenecks on small systems where the user is running their jobs without a batch queue, this feature is disabled by default. In order to enable this feature the following block of commands need to be entered in `mfix.dat`.

```
...  
...  
CHK_BATCHQ_END = .TRUE.      ! Enable the clean termination feature  
BATCH_WALLCLOCK = 3600.0     ! Specify the total wall clock duration  
                             ! of your job in seconds  
TERM_BUFFER = 300.0          ! Specify a buffer time to start  
                             ! clean termination of MFIx
```

Setting `CHK_BATCHQ_END = .TRUE.` in `mfix.dat` will enable the checking of the termination criteria at the beginning of each time step. In the above example, the user has set the total wall clock time for the duration of the batch session to 1 hour (this is specified in seconds in `mfix.dat`) and a buffer of 300 seconds has been set so that MFIx has sufficient time to terminate cleanly by writing out all `*.SP` and `*.RES` files before the batch session terminates. The duration of the buffer is critical for simulations with large files. MFIx will check if `elapsed time >= (BATCH_WALLCLOCK - TERM_BUFFER)` to start clean termination.

Another way to gracefully terminate MFIx as soon as possible is to create an empty file named `MFIx.STOP` (filename all uppercase) in the working directory where MFIx runs.

At the beginning of each time step if `MFIx.STOP` file is detected to exist, then MFIx will terminate gracefully by saving all `*.SP` and `*.RES` files. `CHK_BATCHQ_END` flag must be set to `.TRUE.` in order to activate this feature.

On Linux platforms, the following command could be used to generate the graceful termination trigger:

```
touch MFIx.STOP
```



DO NOT forget to erase the file once MFIX terminates, otherwise next time you run MFIX in the same directory it will terminate immediately.

**rm -f -r ./MFIX.STOP**

Keyword (dimension)	Type	Description
CHK_BATCHQ_END [.FALSE.]	L	Enables clean termination feature.
BATCH_WALLCLOCK [9000.0]	DP	Total wall-clock duration of the job, in seconds.
TERM_BUFFER [180.0]	DP	Buffer time when initiating clean termination, in seconds.

## 5.16. Discrete Element Model (DEM)

The MFIX-DEM main documentation file is titled "Documentation of open-source MFIX-DEM software for gas-solids flows" by R. Garg, J. Galvin, T. Li, and S. Pannala. It is available online at [https://mfix.netl.doe.gov/documentation/dem\\_doc\\_2012-1.pdf](https://mfix.netl.doe.gov/documentation/dem_doc_2012-1.pdf), or in /mfix/doc directory.

The indicated document serves as the user guide for MFIX-DEM and it includes pointers to the DEM code and a discussion on the associated theory.

Keyword (dimension)	Type	Description
<b>DISCRETE_ELEMENT</b> [FALSE]	L	Use discrete particle model for solids. Must be TRUE to do DEM.
<b>MPPIC</b> [FALSE]	L	Use the MPPIC model to represent discrete phase. DISCRETE_ELEMENT flag should also be set to TRUE. Other MPPIC model related flags are explained later in this readme.
<b>DES_CONTINUUM_COUPLED</b> [FALSE]	L	Couple gas and solids flow together.
<b>DES_INTERP_ON</b> [FALSE]	L	Use an interpolation suite to calculate the drag force on each particle based on particle location rather than cell averages.
<b>DES_INTERP_MEAN_FIELDS</b> [FALSE]	L	Use interpolation to compute dispersed phase average fields such as solids volume fraction, solids velocity fields. If false, the average fields are obtained by simple arithmetic averaging.  It is forced to TRUE if MPPIC or cut-cells (Cartesian grid) are used.  If in DEM simulations, drag is interpolated (i.e., DES_INTERP_ON = TRUE), then also it is forced to true for backward compatibility.
<b>DES_INTG_METHOD</b> [EULER]	C	Time stepping scheme EULER - First order Euler scheme ADAMS BASHFORTH - Second order Adams Bashforth scheme.

<b>DIMN</b> [UNDEFINED_I]	I	Specify the dimension of the simulation: 2 or 3. If NO K = '.TRUE.', then DIMN will automatically be set to 2.
<b>GENER_PART_CONFIG</b> [FALSE]	L	Automatically generate an initial particle configuration (position) otherwise use particle input.dat. Also requires setting VOL_FRAC(M), D_P0(M), and DES_EPS_XSTART, DES_EPS_YSTART, and DES_EPS_ZSTART. Once defined this feature will determine the total number of particles in the system and their initial placement. particle input.dat file is ignored.
<b>VOL_FRAC(M)</b> [UNDEFINED]	DP	Only relevant when GENER_PART_CONFIG is T. Volume fraction of the solid phase M for generating articles in the specified domain.
<b>DES_EPS_XSTART</b> [UNDEFINED]	DP	Only needed if GENER_PART_CONFIG. Length of the domain in the x direction wherein particles may be initially placed.
<b>DES_EPS_YSTART</b> [UNDEFINED]	DP	Only needed if GENER_PART_CONFIG. Length of the domain in the y direction wherein particles may be initially placed.
<b>DES_EPS_ZSTART</b> [UNDEFINED]	DP	Only needed if GENER_PART_CONFIG. Length of the domain in the z direction wherein particles may be initially placed.
<b>NFACTOR</b> [10]	I	Only needed if DES_CONTINUUM_COUPLED. Number of times a pure DEM simulation is run before the coupled DEM simulation is started (allows settling).
<b>TSUJI_DRAG</b> [FALSE]	L	Use Tsuji's drag correlation. This correlation is only relevant when invoking Syamlal and O'Brien drag model.
<b>PARTICLES</b> [UNDEFINED_I]	I	Total number of particles.

PVEL_MEAN [0]	DP	Assign initial particle velocities from a Gaussian distribution with the specified mean. Only relevant if PVEL_STDEV is not zero. If used, the assigned velocities will override any other initial settings.
PVEL_STDEV [0]	DP	If not zero, then assign initial particle velocities from a Gaussian distribution with the specified standard deviation. Used with PVEL_MEAN. If used, the assigned velocities will override any other initial settings.
PARTICLES_FACTOR [1.2]	DP	Expand the size of the particle arrays by an arbitrary factor (multiple of the number of particles).

### Boundary Conditions Related Keywords for DEM

Keyword (dimension)	Type	Description
WALLDTSPLIT [F]	L	Treat wall interaction as a two-particle interaction but accounting for the wall properties. Must be TRUE for DEM.
DES_PERIODIC_WALLS [F]	L	Periodic wall boundary condition is imposed on any pair of walls.
DES_PERIODIC_WALLS_X [F]	L	Direction of periodicity: X.
DES_PERIODIC_WALLS_Y [F]	L	Direction of periodicity: Y.
DES_PERIODIC_WALLS_Z [F]	L	Direction of periodicity: Z.

## DES Neighbor Search Related flags needed in DEM model

Keyword (dimension)	Type	Description
DES_NEIGHBOR_SEARCH [1]	I	Neighbor search algorithm. 1=N-square; 2=quadtree (for 2D only); 3=octree (for 3D only); 4=grid based. Options 2 and 3 have not been recently tested. Use them at your own risk.
NEIGHBOR_SEARCH_N [25]	I	Maximum number of steps through a DEM loop before a neighbor search will be performed. (Search may be called earlier).
MN [10]	I	Maximum number of neighbors per particle.
QLM [1]	I	Number of levels to traverse "up" to move a particle to its new quad. Only needed when using octree or quadtree based neighbor search.
QLN [1]	I	Number of levels to traverse "up" to perform particle neighbor search. Only needed when using octree or quadtree based neighbor search.
INIT_QUAD_COUNT [UNDEFINED I]	I	Count to initialize quadtree or octree. Only needed when using octree or quadtree based neighbor search.
MQUAD_FACTOR [1.1]	DP	Factor to create quadtree or octree arrays based on the number of particles. Only needed when using octree or quadtree based neighbor search.
NEIGHBOR_SEARCH_RAD_RATIO [1]	DP	Ratio of the distance (imaginary sphere radius) to particle radius that is allowed before a neighbor search is performed.
FACTOR_RLM [1.2]	DP	Effectively increase the radius of a particle (multiple of the sum of particle radii) for detecting neighbor contacts when using grid based neighbor search or n-square search methods.

## Particle-Particle and Particle-Wall Contact Parameters for soft-spring collision model in DEM

Keyword (dimension)	Type	Description
DES_COLL_MODEL [UNDEFINED_C]	C	Collision model for the soft-sphere approach. By default, the linear spring-dashpot (LSD) model is used (i.e., leave DES_COLL_MODEL undefined for LSD model). Other models include: HERTZIAN. All models require specifying the following parameters: DES_EN_INPUT, DES_EN_WALL_INPUT, MEW, and MEW_W. The default (LSD) model requires: KN, KN_W, KT_FAC, KT_W_FAC, DES_ETAT_FAC, & DES_ETAT_W_FAC. The HERTZIAN model requires: DES_ET_INPUT, DES_ET_WALL_INPUT, E_YOUNG, EW_YOUNG, V_POISSON, & VW_POISSON.
DES_EN_INPUT [UNDEFINED]	DP	The normal restitution coefficient for interparticle collisions that is used to determine the inter-particle normal damping factor. Values are stored as a one dimensional array (see MFIx-DEM doc). So if MAX=3, then 6 values are needed, which are defined as follows: $e_{n11}$ $e_{n12}$ $e_{n13}$ $e_{n22}$ $e_{n23}$ $e_{n33}$ .
DES_ET_INPUT [UNDEFINED]	DP	Tangential restitution coefficient for interparticle collisions. Values are stored as a one dimensional array. Only needed when using the Hertzian collision model.
DES_EN_WALL_INPUT [UNDEFINED]	DP	Normal restitution coefficient for particle wall collisions that is used to determine the particle-wall normal damping factor (see cfassign.f for details). Values are stored as a one dimensional array. So, if MMAX=3, then 3 values are needed, which are defined as follows: $e_{nw1}$ $e_{nw2}$ $e_{nw3}$ .
DES_ET_WALL_INPUT [UNDEFINED]	DP	Tangential restitution coefficient for particle wall collisions. Values are stored as a one dimensional array. Only needed when using the Hertzian collision model.

<b>KN</b> [UNDEFINED]	DP	Normal spring constant for inter-particle collisions. Values are stored as a one dimensional array. Needed when using the default (LSD) collision model.
<b>KT_FAC</b> [2/7]	DP	Ratio of the tangential spring constant to normal spring constant for inter-particle collisions. Use it to specify the tangential spring constant for particle-particle collisions as KT_FAC*KN. Needed when using the default (LSD) collision model.
<b>DES_ETAT_FAC</b> [0.5]	DP	Ratio of the tangential damping factor to the normal damping factor for inter-particle collisions. Needed when using the default (LSD) collision model.
<b>KN_W</b> [UNDEFINED]	DP	Normal spring constant for particle-wall collisions. Needed when using the default (LSD) collision model.
<b>KT_W_FAC</b> [2/7]	DP	Ratio of the tangential spring constant to normal spring constant for particle-wall collisions. Use it to specify the tangential spring constant for particle-wall collisions as KT_W_FAC*KN_W. Needed when using the default (LSD) collision model.
<b>DES_ETAT_W_FAC</b> [0.5]	DP	Ratio of the tangential damping factor to the normal damping factor for particle wall collisions. Needed when using the default (LSD) collision model.
<b>MEW</b> [UNDEFINED]	DP	Particle friction coefficient.
<b>MEW_W</b> [UNDEFINED]	DP	Particle-wall friction coefficient.
<b>E_YOUNG</b> [UNDEFINED]	DP	Young's modulus for the solid phase. Only needed when using the Hertzian collision model.
<b>V_POISSON</b> [UNDEFINED]	DP	Poisson ratio for the solid phase. Only needed when using the Hertzian collision model.
<b>EW_YOUNG</b> [UNDEFINED]	DP	Young's modulus for the wall. Only needed when using the Hertzian collision model.
<b>VW_POISSON</b> [UNDEFINED]	DP	Poisson ratio for the wall. Only needed when using the Hertzian collision model.

## DES Output And Restart Control

Keyword (dimension)	Type	Description
DEBUG_DES [F]	L	Print out additional information from DEM model.
PRINT_DES_DATA [F]	L	Print DEM output to files.
DES_RES_DT [UNDEFINED]	DP	If PRINT_DES_DATA, this is the frequency at which DES.RES and .RES files will be written. This only applies to pure granular simulations, otherwise for coupled simulation the restart frequency is controlled by RES DT.
DES_SPX_DT [UNDEFINED]	DP	IF PRINT_DES_DATA, this is the frequency at which DEM data will be written. This only applies to pure granular simulations, otherwise for coupled simulation the output frequency is controlled by SPX_DT(1).
DEM_OUTPUT_TYPE [UNDEFINED_C]	C	If undefined the default vtp files are written. Other options include: TECPLOT. If 'tecplot' is specified then several .dat files are written including DES_DATA.dat & AVG_EPS.dat. See write_des_data.f for details.

## DEM Mass Inlet/Outlet Specifications:

DEM mass inlet or outlet (bc) are specified over flow planes or 2D surfaces that are normal to one of the coordinate directions and coincide with a face of the scalar control-volume. The values for one of the three pairs of coordinates are equal. The surface is defined by the constant coordinates of each of the four edges and the two equal values for the direction normal to the face. These are specified with physical coordinates. A flow plane must have a wall cell (or an outside boundary) on one side and a flow cell on the other side

For a mass inflow boundary the discrete solids flow rates are specified as the volumetric or mass flow rates. The phase-M flow rates are converted to an equivalent number of phase-m discrete particles enter the system. A uniform inlet velocity is then selected that minimizes the error between the specified bulk density values and the calculated bulk density values. For details see Musser et al., "Development of a discrete mass inflow boundary condition for MFIX", *Journal of Systemics, Cybernetics and Informatics* (JSCI), Volume 9, Number 1, 2011, pg 94-98.



Keyword (dimension)	Type	Description
MAX_PIS [UNDEFINED_I]	I	Maximum number of particles that may exist within a simulation. This quantity is used for calculating the size of arrays for allocation.
DES_BC_X_w(bc) [UNDEFINED]	DP	x coordinate of the west face
DES_BC_X_e(bc) [UNDEFINED]	DP	x coordinate of the east face
DES_BC_Y_s(bc) [UNDEFINED]	DP	y coordinate of the south face
DES_BC_Y_n(bc) [UNDEFINED]	DP	y coordinate of the north face
DES_BC_Z_b(bc) [UNDEFINED]	DP	z coordinate of the bottom face
DES_BC_Z_t(bc) [UNDEFINED]	DP	z coordinate of the top face
DES_BC_TYPE(bc) [UNDEFINED_C]	C	Type of boundary:
MASS_INFLOW or MI		Mass inflow rates for discrete solids phases are specified at the boundary.
MASS_OUTFLOW or MO		This designates the specified region as a discrete mass outflow wherein discrete particles may exit the system. Once a particle center has crossed the mass outflow plane it will continue on that trajectory until it moves fully out of the domain.
DES_BC_VOLFLOW_s(bc, m) [UNDEFINED]	DP	Volumetric flow rate of discrete solids phase M through the mass inlet boundary.
DES_BC_MASSFLOW_s(bc, m) [UNDEFINED]	DP	Mass flow rate of discrete solids phase M through the mass inlet boundary.

<b>DES_ROP_s(bc,m)</b> [UNDEFINED]	DP	Macroscopic bulk density of discrete solids phases at the BC plane.
<b>FORCE_ORD_BC</b> [FALSE]	L	Logical to force the inlet to operate with an ordered boundary condition. This may be useful during long simulations or if the inlet appears to be taking a long time to randomly place particles.
<b>DES_BC_T_s(bc)</b> [UNDEFINED]	DP	Temperature of incoming particles.
<b>DES_BC_X_s(bc,m,n)</b> [UNDEFINED]	DP	Mass fraction of solids phase m species n for incoming particles.

### DEM Initial Conditions:

Each initial condition (IC) is specified over a rectangular region that corresponds to the scalar numerical grid. These are 3D regions:  $X_w$   $X_e$ ,  $Y_s$   $Y_n$ , and  $Z_t$   $Z_b$ . Initial condition regions are used to specify particle properties. The DEM initial conditions are only relevant when solving DEM energy and/or species equations (see below for DEM energy and reactive chemistry related keywords).

<b>Keyword (dimension)</b>	<b>Type</b>	<b>Description</b>
<b>DES_IC_X_w(ic)</b> [UNDEFINED]	DP	x coordinate of the west face
<b>DES_IC_X_e(ic)</b> [UNDEFINED]	DP	x coordinate of the east face
<b>DES_IC_Y_s(ic)</b> [UNDEFINED]	DP	y coordinate of the south face
<b>DES_IC_Y_n(ic)</b> [UNDEFINED]	DP	y coordinate of the north face
<b>DES_IC_Z_b(ic)</b> [UNDEFINED]	DP	z coordinate of the bottom face
<b>DES_IC_Z_t(ic)</b> [UNDEFINED]	DP	z coordinate of the top face

## DEM Energy Equations:

Keyword (dimension)	Type	Description
DES_ENERGY_EQ [.FALSE.]	L	Solve energy equations for DEM.
DES_CONV_EQ [.TRUE.]	L	Include particle-gas convection in DEM heat transfer model.
DES_COND_EQ [.TRUE.]	L	Include particle-particle conduction in DEM heat transfer model.
DES_COND_EQ_PP [.TRUE.]	L	Include particle-particle contact conduction in DEM heat transfer model. (Requires DES_COND_EQ set to true.)
DES_COND_EQ_PFP [.TRUE.]	L	Include particle-fluid-particle conduction in DEM heat transfer model. (Requires DES_COND_EQ set to true.)
DES_RADI_EQ [.TRUE.]	L	Include particle--particle radiation in DEM heat transfer model.
DES_CONV_CORR [RANZ_1952]	C	Specify the Nusselt number correlation used for particle-gas convection. (Only RANZ_1952 is presently included.)
FLPC [@(1.0/5.0)]	DP	Fluid lens proportion constant used to calculate the radius of the fluid lens that surrounds a particle. Used in the particle-fluid-particle conduction model.
DES_K_s0(m) [UNDEFINED]	DP	Specified constant solids thermal conductivity of solids phase m.
DES_C_ps0(m) [UNDEFINED]	DP	Specified constant solids specific heat of solids phase m.
DES_Em(m) [UNDEFINED]	DP	Emissivity of solids phase m.
DES_MIN_COND_DIST [UNDEFINED]	DP	Minimum separation distance between the surfaces of two contacting particles.

## DEM Reactive Chemistry Model:

Keywords for specifying DEM reactions are listed below. A complete description of specifying chemical reactions is provided in Section 5.11.

Keyword (dimension)	Type	Description
DES_SPECIES_EQ(m) [.FALSE.]	L	Solve the species equations for solids phase m.
DES_NMAX_s(m) [UNDEFINED_I]	I	Number of species comprising solids phase m.
DES_SPECIES_s [UNDEFINED_C]	C	Name of solids phase m, species n as it appears in the materials database
DES_SPECIES_ALIAS_s(m,n) [UNDEFINED_C]	C	User defined name of solids phase m, species n
REACTION_MODEL [‘VARIABLE_DENSITY’]	C	Reaction model used to calculate the effects of a gas-solids reaction on a particle.
VARIABLE_DENSITY		Constant particle diameter. Particle diameter changes to accommodate a loss/gain in mass.
SHRINKING_PARTICLE		Constant particle density. Particle’s diameter changes to accommodate a loss/gas in mass.

## DEM Continuum Hybrid Model:

Listed below are the keywords needed to invoke a hybrid scheme wherein solids in the same system can be simultaneously modeled using the Discrete Element and Continuum Models. Note that any and all parameters/quantities that are needed for running a standard, stand-alone continuum model or discrete element model are still required. However, different variables are used to specify the number of discrete solids phases, their diameter and density. At this time all of the discrete and continuum solids phases interact via a solids-solids drag like term adapted from D. Gera, M. Syamlal, T.J. O'Brien, “Hydrodynamics of particle segregation in fluidized beds”, *International Journal of Multiphase Flow*, v30, 2004, p419-428. This term has several parameters that may be controlled including the coefficient of friction (C\_F) and the SEGREGATION\_SLOPE\_COEFFICIENT, which are discussed in the Physical Parameters section.

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## 5.17. MPPIC model:

The MFIX-DEM main documentation file is titled "Documentation of open-source MFIX-PIC software for gas-solids flows" by R. Garg and J. F. Dietiker. It is available online at [https://mfix.netl.doe.gov/documentation/mfix\\_pic\\_doc.pdf](https://mfix.netl.doe.gov/documentation/mfix_pic_doc.pdf), or in /mfix/doc directory.

The indicated document serves as the theory guide for the MPPIC model implementation details in MFIX code. The MFIX code with MPPIC model is referred to as MFIX-PIC in the indicated document and hereinafter. A brief discussion on setting up of input file for MPPIC model assumes reader's familiarity with the MPPIC documentation.



Note that MPPIC model has not been tested with MPI modules. It has only been tested in serial compilation mode. It will work with cut-cell modules in MFIX, but there needs to be work done to ensure conservative coupling between the two phases

The MPPIC model is invoked by setting MPPIC flag to TRUE in conjunction with DISCRETE\_ELEMENT set to TRUE. The initial conditions for the MPPIC model are specified in the same way as continuum model. It is noted that this is a departure from the DEM model in MFIX where the initial condition for solid phase was specified by special DEM related flags. Although the physical region where the parcels are seeded is specified by the same flags as those used in continuum model setup, there is still need to specify the number of parcels per cell and their statistical weights. There are two methods to specify the initial seeding of parcels. In both methods, the user first defines the physical region where the initial solids will be seeded. As discussed earlier, this is done by the same flags that are used in continuum representation of dispersed phase. In the first method, the user specifies the number of parcels per cell by the setting the flag "CONSTANTNPC" to TRUE and specifying number of parcels per cell for each phase by the array "NPC\_PIC". In this case, the user defined number of parcels per cell are randomly seeded in the initial physical region specified by user. The statistical weight is assigned to parcels such that the solid volume fraction implied by parcels equals the user defined solid volume fraction (see the MPPIC documentation).

In the second method, the statistical weight of parcels is fixed by setting the flag "CONSTANTWT" to TRUE along with specifying the statistical weight of parcels belonging to each phase by the array "STATWT\_PIC". The number of parcels per cell is computed by the code such that the solid volume fraction implied by parcels equals the user defined solid volume fraction.

The current implementation of MPPIC model also has the frictional stress model implementation outlined in Snider's [1] paper. This is based on our best understanding of the model from the paper. This model does not simulate very stably.

The minimum gas voidage at maximum packing beyond which the MPPIC frictional stress model gets invoked is still defined by the flag EP\_STAR that is generally used in continuum representation.

Below is the description of MPPIC model related flags needed to set up MFIx input file.

Keyword (dimension)	Type	Description
MPPIC [.FALSE.]	L	Use the MPPIC parcel based representation for dispersed phase.
MPPIC_CONSTANTNPC [.TRUE.]	L	A set number (NPC_PIC) of parcels per cell are initially randomly seeded. Statistical weight is computed by the code.
MPPIC_CONSTANTWT [.TRUE.]	L	Initially parcels are randomly seeded with user defined statistical weight (STATWT_PIC). The number of parcels per cell is computed by the code.
NPC_PIC [UNDEFINED]	I	Desired number of parcels per cell when MPPIC_CONSTANTNPC is true.
STATWT_PIC [UNDEFINED]	DP	Desired statistical weight of parcels when MPPIC_CONSTANTWT is true.
MPPIC_COEFF_EN1 [UNDEFINED]	DP	First frictional coefficient of restitution (see the MPPIC model theory guide).
MPPIC_COEFF_EN2 [UNDEFINED]	DP	Second frictional coefficient of restitution (see the MPPIC model theory guide).
MPPIC_COEFF_EN_WALL [UNDEFINED]	DP	Normal coefficient of friction for parcel-wall collision in MPPIC model
MPPIC_GRAV_TREATMENT [TRUE]	L	To turn on the special case in frictional model when the impulse velocity's direction is collinear with gravity (see the MPPIC model theory guide).
MPPIC_DRAG_IMPLICIT [.FALSE.]	L	For implicit treatment of the drag force term in parcel governing equations.
CFL_PIC [0.1]	DP	CFL number used to decide maximum time step size for parcel's evolution equations.
BC_APPLY_TO_MPPIC	L	Special flag to turn off a pressure outlet boundary condition specifically on the

[.TRUE.]		solids phase. For example, if pressure outlet BC is selected but the user does not want the solids to leave the system, then set this flag to .FALSE.
MPPIC_SOLID_STRESS_SNIDER [.FALSE.]	L	Turn on Snider's version of frictional model. Does not run very stably.
PSFAC_FRIC_PIC [100]	DP	Ps term in the frictional stress model of Snider
FRIC_EXP_PIC [2.5]	DP	Beta term in the frictional stress model of Snider
FRIC_NON_SING_FAC [1E-07]	DP	Non-singularity term (epsilon) in the frictional stress model of Snider



## 5.18. ISAT and Direct Integration for Chemical Reactions

Keyword (dimension)	Type	Description
CALL_DI [F]	L	Variable to decide if chemical reactions are time-split and solved using direct integration (DI) with ODE solver. Do not call time-splitting and DI
CALL_ISAT [F]	L	Variable to decide if chemical reactions are time-split and solved using ISAT. Do not use time-splitting and ISAT.
CALL_GROW [F]	L	Variable to decide if particle growth due to chemical reactions is calculated. Do not do particle growth calculations.
ISATdt	DP	Time step for ISAT simulations (usually the value is less than the average time step in MFIX).

If “CALL\_DI=.TRUE.” or “CALL\_ISAT=.TRUE.”, MFIX uses DI or ISAT to calculate the chemical reactions using the time-splitting method. The ODE solver used here is ODEPACK ([www.netlib.org/odepack](http://www.netlib.org/odepack)).

The algorithm of MFIX using ODEPACK (CALL\_DI = .TRUE.) is shown in Fig. 6 of the ISAT manual in the document directory. By this call, the ODEs will be solved every ISATdt if provided, or every time step of MFIX if ISATdt is not provided by DI.

If “CALL\_ISAT=.TRUE.”, user must have the ISATAB library and link it in the make\_mfix file by specifying ‘-l{path to the library}/libraryname’ at the end of the link options. Figure 7 of the ISAT manual in the document directory shows the flow of MFIX using ISAT. The user must provide ISATdt to keep the high performance of ISAT.

The following is a list of files that are usually modified to include the chemical reactions using ODE solver or ISAT:

mchem_mod.f	Sets the global variables.
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misat_table_init.f	Sets the controlling parameters for ISATAB.
mchem_odepack_init.f	Sets the controlling parameters for ODEPACK.
fex.f	The subroutine provides the source terms for equations in the ODE solver and source terms of reactions rates. The source terms have the following order: $[\rho_g, T_g, X_g], [\epsilon_{s1}, T_{s1}, X_{s1}, d_1], \dots, [\epsilon_{sm}, T_{sm}, X_{sm}, d_m]$ . If the variables are constant during the simulations, the source terms should be set to zero. For example, for isothermal chemical reactions, the source terms of $T_g$ and $T_{sm}$ are zero.
calc_jacobian.f	Provides the Jacobian matrix for ODE solver.
transport_prop.f	Provides transport properties.
physical_prop.f	Provides physical properties.

### Example using ISAT

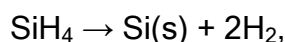
The non-isothermal silane pyrolysis in a fluidized bed (FB) is used as a benchmark case, where detailed chemical reactions are considered. The FB reactor is fed with a mixture of silane ( $\text{SiH}_4$ ) and nitrogen ( $\text{N}_2$ ). First, a reversible gas phase reaction occurs:



The highly reactive  $\text{SiH}_2$  undergoes a gas phase reaction to form  $\text{Si}_2\text{H}_6$ :



Then the heterogeneous decomposition of  $\text{SiH}_4$  and  $\text{SiH}_2$  on alumina ( $\text{Al}_2\text{O}_3$ ) particles is described by two irreversible reactions:



Thus there are five gaseous species and two solid species needed to describe this flow.

The following files are modified to simulate this case:

mchem_mod.f	Sets the global variables.
misat_table_init.f	Controlling parameters for ISATAB are set.
mchem_odepack_init.f	Controlling parameters for ODEPACK are set.

fex.f	<p>The subroutine provides the source terms to ODE solver and sources terms of reactions. For this case, the order of ODEs is</p> <p><math>[\rho_g, T_g, X_{SiH_4}, X_{SiH_2}, X_{H_2}, X_{Si_2H_6}, X_{N_2}], [\epsilon_{s1}, T_{s1}, X_{Si}, X_{Al_2O_3}, d_1]</math>.</p> <p>As written in the code, the user should provide the source terms of reactions, which is <math>RXN\_source\_g(n)</math> for gas phase species <math>n</math> and <math>RXN\_source\_s(m,n)</math> for solid phase <math>m</math> and species <math>n</math>. Then the source terms of the ODEs are calculated. Note that <math>N_2</math> and <math>Al_2O_3</math> are inert species; therefore the source terms are set to zero.</p>
calc_jacobian.f	<p>The subroutine provides the Jacobian matrix for ODE solver.</p> <p>The example case uses the Jacobian matrix generated by ADIFOR (g_derives.f). The ADIFOR can be downloaded from the website (<a href="http://www-unix.mcs.anl.gov/autodiff/ADIFOR">http://www-unix.mcs.anl.gov/autodiff/ADIFOR</a>).</p>
transport_prop.f	Provides the transport properties.
physical_prop.f	Provides the physical properties such as averaged molecular weight.

## 5.19. Direct Quadrature Method of Moments (DQMOM)

Keyword (dimension)	Type	Description
CALL_DQMOM [F]	L	Variable to decide if the population balance equations are solved. Do not invoke DQMOM
Nscalar [0]	I	Number of solid phases to solve the population balance equations.
Aggregation_eff [0.0]	DP	Success-factor for aggregation.
Breakage_eff [0.0]	DP	Success-factor for breakage.

### Example using DQMOM

The example case is a fluidized bed simulation with two solid phases; each has its own particle size. In the code, the population balance equation is turned on. If aggregation dominates, the average particle size will increase. If breakage dominates, the average particle size will decrease. The user can turn off DQMOM by set "Call\_DQMOM = .FALSE.". The ODE solver for the scalar in the MFIx code solves the population balance equation, so "Nscalar" has to be set as the number of solid phases. The initial values for the scalar are set as the initial particle diameter. The aggregation and breakage kernel from kinetic theory is used. The success factor of aggregation and breakage can be changed by setting different values for "Aggregation\_eff" and "Breakage\_eff". An example mfix.dat file is included.

## 5.20. Quadrature Method of Moments (QMOM)

Please refer to the following document for a brief description of the implementation and utilization of the quadrature method of moments (QMOM): A. Passalacqua, and R. O. Fox, "Documentation of open-source MFIX-QMOM software for gas-solids flows", available from [https://mfix.netl.doe.gov/documentation/qmomk\\_doc\\_2012-1.pdf](https://mfix.netl.doe.gov/documentation/qmomk_doc_2012-1.pdf)

## 5.21. Cohesion Model in DEM

Cohesive interparticle forces represent an addition to the discrete-particle simulation. The raw code for this addition was located in the *mfix/model/cohesion* directory. These forces can be implemented using both a square-well model and a Hamaker van der Waals model. Cohesive forces are turned-on by setting the logical variable "USE\_COHESION" equal to true in the *mfix.dat* file.

### Square-Well model

The nature of the square-well model is determined by the well width ( $r_{outer}$ ) and the well depth ( $D$ ). Both of these parameters are specified in the *mfix.dat* file. Furthermore, the simulation is set up to use a separate set of square-well parameters with particle-particle interactions and particle-wall interactions. Generally, the well depth is set to be twice as large for particle-wall interactions to be consistent with the increased surface contact that would exist in such interactions.

Within the square well model, cohesive interactions are implemented as instantaneous momentum impulses ( $J$ ) resulting in a change in particle velocity as shown below:

$$m\vec{v}_{i,post} = m\vec{v}_{i,pre} - J_*\vec{k}_{ij,*} \quad (1)$$

where  $m$  is the particle mass,  $\vec{v}_{i,pre}$  is the pre-interaction velocity,  $\vec{v}_{i,post}$  is the post-interaction velocity and  $\vec{k}_{ij,*}$  is the unit vector connecting the particle centers. This treatment leads to three types of cohesive interactions in the square-well model: approaching cohesive interactions, escaping cohesive interactions and capture cohesive interactions. The momentum impulse during each of these interactions can be calculated by solving the momentum balance along with the stipulation that the total kinetic energy of the two-particle system changes by the depth of the square well ( $D$ ). This treatment leads to the following expressions for the momentum impulse in each type of interaction:

$$J_{coh,app} = \frac{m}{2} \left( \vec{k}_{ij,1} \cdot \vec{v}_{ij,a} - \sqrt{\frac{4D}{m} + (\vec{k}_{ij,1} \cdot \vec{v}_{ij,a})^2} \right) \quad (2)$$

$$J_{coh,esc} = \frac{m}{2} \left( \vec{k}_{ij,3} \cdot \vec{v}_{ij,c} + \sqrt{(\vec{k}_{ij,3} \cdot \vec{v}_{ij,c})^2 - \frac{4D}{m}} \right) \quad (3)$$

$$J_{coh,cap} = m(\vec{k}_{ij,3} \cdot \vec{v}_{ij,c}) \quad (4)$$

Because cohesive interactions are implemented only once on approach and once as the particles depart, the identification of a square-well interaction cannot be done simply using the separation distance. Special book-keeping is employed to ensure that square-well cohesive interactions are implemented only during the time step when the well-widths first become overlapped for approaching interactions and when they first become un-overlapped for departing interactions. This book-keeping involves recording a “list of links” for each particle. Two particles are considered “linked” if their separation distance is less than the width of their square-wells (the square-wells overlap). The list of links is updated at every cohesive interaction. Cohesive square-well interactions are recognized using both the list of links and the particle separation distance at each. For example, an approaching cohesive interaction is only implemented if two particles overlap their square-wells, yet they are not on their respective list of links. The particles will be added to each list during the interaction and at the next time step, they may still have overlapped square-wells, but an interaction will not be implemented because they are already on their respective list of links. A similar algorithm is used to identify departing interactions.

### Hamaker model

The Hamaker model for van der Waals forces predicts the cohesive force between equal-sized, spherical particles according to the following equation [1]:

$$F_{vdW} = \frac{Ar_{inner}}{12H^2} \quad (5)$$

where  $r_{inner}$  is the particle radius;  $A$  is the Hamaker constant, which is specific to a given material and has typical values on the order of  $10^{-20}$  J; and  $H$  is the minimum surface-to-surface separation distance between two particles  $i$  and  $j$ :

$$H = \sqrt{(x_i - x_j)^2 + (y_i - y_j)^2} - 2r_{inner} \quad (6)$$

The corresponding cohesive force between a spherical particle and a flat wall is [1]:

$$F_{vdW,wall} = \frac{A_{wall}r_{inner}}{6H^2} \quad (7)$$

For both of these expressions, the cohesive force approaches infinity as the separation distance approaches zero. This singularity incurred at particle contact is avoided by introducing a “cutoff” distance,  $H_{cut}$ . For separation distances below this cutoff distance,

the interparticle cohesive force is given by a surface adhesion force ( $F_{ad}$ ) model [2].

$$F_{ad} = 2\pi r_{inner}\gamma \quad (8)$$

where  $\gamma$  is a constant surface energy per unit area. The contact surface energy is calculated at the beginning of each simulation based on the specified Hamaker constant and cutoff distance to ensure that the force is continuous at the cutoff distance. Thus, the cohesive force is maintained at a constant value for any separation distances (based on equation 6) below the minimum cutoff distance, which includes any “negative” separation distances that occur during actual particle contact.

### Cohesive Interaction Search

Cohesive interactions are identified using a method separate from the functions used to identify particle contacts in the discrete-particle model. The simulation domain is divided up into search grids in order to reduce the number of particles that must be checked for interactions by each particle. The search grid is made up of several uniform boxes that are defined to be at least as wide as the largest length-scale in the cohesive forces. In this way, particles are only checked against other particles in their box and adjacent boxes. The size of the search boxes ensures that a particle could not have an interaction with particles that are located in boxes outside the adjacent boxes. The arrays recording the indices of the search box for each particle are updated at each time step (*update\_search\_grids.f*). Note that the van der Waals model uses the same search algorithm as for DEM collisions and is fully parallelized unlike the square-well model.

### Cohesion Variables

Keyword (dimension)	Type	Description
MASTER_WELL_DEPTH	DP	Square-well depth read in from mfix.dat for simulations that use same well depth for all particle-particle interactions
MASTER_WALL_WELL_DEPTH	DP	Square-well depth for particle-wall interactions read in from mfix.dat for simulations that use same well width for all particle-wall interactions
RADIUS_RATIO	DP	Ratio of square-well width to particle radius
WALL_RADIUS_RATIO	DP	Ratio of wall square-well depth to particle radius

USE_COHESION	L	Switch to turn cohesion on and off
SQUARE_WELL	L	Switch to turn square well on and off
COHESION_DEBUG	I	Flag to turn on output lines for debugging cohesive simulation
VAN_DER_WAALS	L	Flag to turn on the use Hamaker van der Waals forces
HAMAKER_CONSTANT	DP	Hamaker constant used for particle-particle interactions
VDW_INNER_CUTOFF	DP	Minimum separation distance below which van der Waals forces are calculated using a surface adhesion model
VDW_OUTER_CUTOFF	DP	Maximum separation distance above which van der Waals forces are not implemented
WALL_HAMAKER_CONSTANT	DP	Hamaker constant used in particle-wall interactions
WALL_VDW_INNER_CUTOFF	DP	Minimum separation distance below which van der Waals forces are calculated using a surface adhesion model (particle-wall interactions)
WALL_VDW_OUTER_CUTOFF	DP	Maximum separation distance above which van der Waals forces are not implemented (particle-wall interactions)
Asperities	DP	Mean radius of surface asperities that influence the cohesive force following a model by Rumpf (1990)



## 5.22. Cartesian grid

A new capability, called Cartesian grid cut-cell technique has been implemented in MFIx, which allows the definition of curved or sloping boundaries, instead of the usual stair-step representation. Computational cells are truncated at the wall to conform to the shape of the boundaries. When a face is truncated, the velocity node is moved to the center of the face. The cell truncation introduces an additional face, called the cut face. Face surface areas and cell volumes are updated based on the shape of the cut cell. The contribution of the new cut face is added to the computation. The data can be saved in a vtk file for post-processing purpose.

A detailed user guide named `Cartesian_grid_user_guide.pdf` is located in `mfix/doc` directory. It should be read prior to utilizing the Cartesian grid option to get familiar with this technique, and associated keywords. This file is also available online at [https://mfix.netl.doe.gov/documentation/Cartesian\\_grid\\_user\\_guide.pdf](https://mfix.netl.doe.gov/documentation/Cartesian_grid_user_guide.pdf).



## 6. Mailing lists

Several mailing lists are available to communicate among MFIX users and developers. When your subscription to MFIX is accepted, you are automatically added to the mfix-news mailing list, where important announcements about MFIX are shared with the MFIX community.

The most widely used mailing list is mfix-help, which allows users to post questions and eventually help other users with similar issues.

The mailing list home page is located at <https://mfix.netl.doe.gov/sympa>. Click on the “List of lists” tab to view all available mailing lists. Most of them have a very low bandwidth, and most users only subscribe to the mfix-help list.

Once you subscribe to a list, you can send/receive messages to/from the MFIX community. You can also search archived messages to see if there is already a solution to a common problem.

There are many options to manage your subscription, including subscribing, unsubscribing, and choosing the delivery mode.

Please visit <https://mfix.netl.doe.gov/sympa/help/user> to view the mailing list user guide.

### Mailing list etiquette:

- 1) Please allow sufficient time (say 2 to 3 business days) for MFIX developers and users to reply before posting unanswered questions again.
- 2) Unless prior arrangement has been made with a given MFIX developer, do not send requests directly to the developer, but send the request to the appropriate mailing list instead. This ensures proper archiving of the thread and provides better opportunity for everyone to reply. Follow-up questions should also be sent to the mailing list.
- 3) Prior to submitting help requests regarding MFIX installation or compilation issues, please check the archives of mfix-help and if you are still having a problem, email [mfix-help@mfix.netl.doe.gov](mailto:mfix-help@mfix.netl.doe.gov) by providing the following important details in your message after the description of the problem encountered:
  - a. MFIX version you are trying to install or run
  - b. Some details on your operating system environment (for Linux: copy and paste the response of `uname -a` command, Linux distribution name and version also)



- c. Your compiler name and version number (e.g. ifort -v will give the version number for Intel fortran compiler)
- d. Output for your \$PATH environment (in csh type echo \$PATH)
- e. Your MPI library name and version number (if compilations problem with DMP mode encountered but make sure you can compile and run a simple hello world type MPI program with your current installation) Also please provide hardware details such as number of cores per socket in your system (or send the output for “cat /proc/cpuinfo” and how many cores you are trying to utilize).