

# libFM 1.4 - Manual

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<http://www.libfm.org/>

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*Note: advanced chapters are marked with \*.*

# 1 Installation

- **Windows:** download the compiled executable from <http://www.libfm.org/libfm-1.40.windows.zip>. You can skip section 1.1.
- **Linux:** compile LIBFM by following the instructions in section 1.1.
- **MacOS X:** compile LIBFM by following the instructions in section 1.1.

## 1.1 Compiling

LIBFM has been tested with the GNU compiler collection and GNU make. Both should be available in Linux and MacOS X.

With the following steps, you can build LIBFM:

1. Download libFM source code: <http://www.libfm.org/libfm-1.40.src.tar.gz>
2. Unzip and untar: e.g. `tar -xzvf libfm-1.40.src.tar.gz`
3. Enter the directory `libfm-1.40.src` and compile the tools: `make all`

### Overview of files

- `license.txt`: license for usage of LIBFM
- `history.txt`: version history and changes
- `readme.pdf`: this manual
- `Makefile`: compiles the executables using `make`
- `bin`: the folder with the executables<sup>1</sup>
  - `libFM`: the LIBFM tool
  - `convert`: a tool for converting text-files into binary format
  - `transpose`: a tool for transposing binary design matrices
- `scripts`
  - `triple_format_to_libfm.pl`: a Perl script for converting comma/tab-separated datasets into LIBFM-format.
- `src`: the source files of LIBFM and the tools

---

<sup>1</sup>The executables have to be built with `make`, see sec. 1.1.

## 2 Data Format

LIBFM supports two file formats for input data: a text format and a binary format. Working with the text format is easier and recommended for new LIBFM-users.

### 2.1 Text Format

The data format is the same as in SVMlite [3] and LIBSVM [1]<sup>2</sup> : Each row contains a training case  $(\mathbf{x}, y)$  for the real-valued feature vector  $\mathbf{x}$  with the target  $y$ . The row states first the value  $y$  and then the non-zero values of  $\mathbf{x}$ . For binary classification, cases with  $y > 0$  are regarded as the positive class and with  $y \leq 0$  as the negative class.

#### Example

```
4 0:1.5 3:-7.9
2 1:1e-5 3:2
-1 6:1
...
```

This file contains three cases. The first column states the target of each of the three case: i.e. 4 for the first case, 2 for the second and -1 for the third. After the target, each line contains the non-zero elements of  $\mathbf{x}$ , where an entry like 0:1.5 reads  $x_0 = 1.5$  and 3:-7.9 means  $x_3 = -7.9$ , etc. That means the left side of INDEX:VALUE states the index within  $\mathbf{x}$  whereas the right side states the value of  $x_{\text{INDEX}}$ , i.e.  $x_{\text{INDEX}} = \text{VALUE}$ .

In total the data from the example describes the following design matrix  $X$  and target vector  $y$ :

$$X = \begin{pmatrix} 1.5 & 0.0 & 0.0 & -7.9 & 0.0 & 0.0 & 0.0 \\ 0.0 & 10^{-5} & 0.0 & 2.0 & 0.0 & 0.0 & 0.0 \\ 0.0 & 0.0 & 0.0 & 0.0 & 0.0 & 0.0 & 1.0 \end{pmatrix}, \quad \mathbf{y} = \begin{pmatrix} 4 \\ 2 \\ -1 \end{pmatrix} \quad (1)$$

#### 2.1.1 Converting Recommender Files

In recommender systems often a file format like `userid itemid rating` is used. A Perl script to convert such datasets (and also more complex ones like in context-aware settings) to the libFM file format is included in the `scripts` directory. E.g. for converting the `ratings.dat` file from the Movielens 1M<sup>3</sup> dataset into LIBFM format, call:

```
./triple_format_to_libfm.pl -in ratings.dat -target 2 -delete_column 3 -separator "::-"
```

The output will be written to a file with extension `.libfm`. In the example the output is written to `ratings.dat.libfm`.

If a single dataset consists of multiple files, e.g. a train and test split, then the conversion script should be called with both files (several files for the `-in` option):

```
./triple_format_to_libfm.pl -in train.txt,test.txt -target 2 -separator "\t"
```

Warning: If you would run the conversion script for each file separately, the variables (ids) would not match. E.g. the  $n$ -th variable in the first file is different from the  $n$ -th variable in the second file.

### 2.2 Binary Format\*

Besides the standard text format, LIBFM supports a binary data format. The advantages of the binary format are: (1) faster reading, (2) if your data does not fit into main memory, the binary data format supports caching data on hard disc and keeping only a small portion in memory (use the `--cache_size` in LIBFM), (3) if you work with ALS and MCMC, you can precompute the transposed design matrix which saves time when the data set is read.

To convert a file in LIBFM text format to the binary data format, use the tool `convert` in the `bin` folder:

<sup>2</sup>Note: LIBFM supports files where the variable index starts with 0. In SVMlite and LIBSVM, the index has to start with 1. LIBFM supports both, i.e. files that work with SVMlite or LIBSVM can be used directly in LIBFM.

<sup>3</sup><http://www.grouplens.org/node/12>

## Convert

Version: 1.4.0  
Author: Steffen Rendle, [steffen.rendle@uni-konstanz.de](mailto:steffen.rendle@uni-konstanz.de)  
WWW: <http://www.libfm.org/>  
License: Free for academic use. See [license.txt](#).

---

-help            this screen  
-ifile          input file name, file has to be in binary sparse format  
                 [MANDATORY]  
-ofilex        output file name for x [MANDATORY]  
-ofiley        output file name for y [MANDATORY]

**Example** To convert the Movielens dataset from the example above to binary format:

```
./convert --ifile ratings.dat.libfm --ofilex ratings.x --ofiley ratings.y
```

The output will consist of two files: (1) a file containing the design matrix  $X$ , i.e. the predictor variables and (2) a file containing the prediction targets  $y$ . It is recommended that these files have the ending `.x` for the design matrix and `.y` for the targets.

### 2.2.1 Transpose data

For MCMC and ALS learning, a transposed design matrix is used. If you use the text format, the data is automatically transposed by LIBFM internally. However if you use binary format, the transposed data has to be present in binary format as well. To transpose a design matrix to binary format use the tool `transpose`.

## Transpose

Version: 1.4.0  
Author: Steffen Rendle, [steffen.rendle@uni-konstanz.de](mailto:steffen.rendle@uni-konstanz.de)  
WWW: <http://www.libfm.org/>  
License: Free for academic use. See [license.txt](#).

---

-cache\_size    cache size for data storage, default=200000000  
-help          this screen  
-ifile         input file name, file has to be in binary sparse format  
                 [MANDATORY]  
-ofile         output file name [MANDATORY]

**Example** To transpose the Movielens dataset from the example above:

```
./transpose --ifile ratings.x --ofile ratings.xt
```

The output will be a transposed copy of the design matrix. It is recommended that the transposed file has the ending `.xt`.

### 3 libFM

The LIBFM tool trains a Factorization Machine (FM) [4] model from training data `-train` and predicts the test data `-test`. LIBFM has the following options:

```
-----
libFM
  Version: 1.40
  Author:  Steffen Rendle, steffen.rendle@uni-konstanz.de
  WWW:    http://www.libfm.org/
  License: Free for academic use. See license.txt.
-----

-cache_size    cache size for data storage (only applicable if data is
               in binary format), default=infty
-dim           'k0,k1,k2': k0=use bias, k1=use 1-way interactions,
               k2=dim of 2-way interactions; default=1,1,8
-help          this screen
-init_stdev    stdev for initialization of 2-way factors; default=0.1
-iter          number of iterations; default=100
-learn_rate    learn_rate for SGD; default=0.1
-meta          filename for meta information about data set
-method        learning method (SGD, SGDA, ALS, MCMC); default=MCMC
-out           filename for output
-regular       'r0,r1,r2' for SGD and ALS: r0=bias regularization,
               r1=1-way regularization, r2=2-way regularization
-relation      BS: filenames for the relations, default=''
-rlog          write measurements within iterations to a file;
               default=''
-task          r=regression, c=binary classification [MANDATORY]
-test          filename for test data [MANDATORY]
-train         filename for training data [MANDATORY]
-validation    filename for validation data (only for SGDA)
-verbosity     how much infos to print; default=0
```

#### 3.1 Mandatory Parameters

- The first mandatory parameter to specify is the `-task` which is either classification (`-task c`) or regression (`-task r`).
- Secondly training data (`-train`) and test data (`-test`) has to be present. You can use here a data file in LIBFM-text format or binary format (see sec. 3.2.2 for details about using binary files).
- Third, the dimensionality of the factorization machine has to be specified with `-dim`. This argument consists of three numbers:  $k_0, k_1, k_2$ .
  - $k_0 \in \{0, 1\}$  determines if the global bias term  $w_0$  should be used in the model (see [4] for details).
  - $k_1 \in \{0, 1\}$  determines if one-way interactions (bias terms for each variable), i.e.  $\mathbf{w}$  should be used in the model (see [4] for details).
  - $k_2 \in \mathbb{N}_0$  gives the number of factors that are used for pairwise interactions, i.e. the  $k$  of  $V \in \mathbb{R}^{p \times k}$  (see [4] for details).

**Example** An FM for a regression task using bias, 1-way interactions and a factorization of  $k = 8$  for pairwise interactions:

```
./libFM -task r -train ml1m-train.libfm -test ml1m-test.libfm -dim '1,1,8'
```

## 3.2 Optional Parameters

### 3.2.1 Basic Parameters

- **out**: After training is finished, you can write all predictions of the test dataset to the file specified by **out**. The out-file is in text format, has as many lines as the test dataset and the  $i$ -th line states the prediction of the  $i$ -th test case. Note that for classification the output is the probability that the case has the positive class.
- **rlog**: A logfile with statistics about each iteration is generated. The file is in CSV format using TAB separated fields. Note that it depends on the learning method which fields are reported.
- **verbosity**: With the argument **-verbosity 1**, LIBFM prints more information. This is useful to check if your data is read correctly, and to find errors.

### 3.2.2 Advanced Parameters\*

**Grouping** You can group input variables using the **meta** option. Grouping can be used for MCMC, SGDA and ALS to define a more complex regularization structure. Every group can have an individual regularization parameter. To use grouping, the **meta** parameter expects the name of a textfile with as many lines as there are input variables (columns of the design matrix). Each line specifies the group ID of the corresponding input variable. Please note that IDs for groups should be numerical and start with 0. E.g. a grouping file for the design matrix of the example in eq. (1) (which has 7 columns; the largest ID is 6) could be:

```
2
2
0
1
1
1
1
0
```

Which would read: in total there are three groups. The first two variables (columns in the design matrix) have the same group, the third and last have the same group and the fourth to sixth variable have the same group.

**Binary Data and Caching** In sec. 2.2 it was stated that the filenames for binary files should end with **.x** for the design matrix, **.y** for the target and **.xt** for transposed data. If you want to use binary data in LIBFM, the filenames for the command line arguments of training, test and validation should be specified *without* the **.x**, **.y**, **.xt** ending. I.e. if you have compiled training (**ml1m-train.x**, **ml1m-train.y**, **ml1m-train.xt**) and test data (**ml1m-test.x**, **ml1m-test.y**, **ml1m-test.xt**) call:

```
./libFM -task r -train ml1m-train -test ml1m-test -dim '1,1,8'
```

LIBFM will automatically append the proper file extensions and load the data files that are necessary for the learning algorithm.

If your data does not fit into memory, you can specify how much of a file LIBFM is allowed to keep in memory:

```
./libFM -task r -train ml1m-train -test ml1m-test -dim '1,1,8' -cache_size 100000000
```

In this example, 100,000,000 Bytes (100 MB) would be used as cache for each **.x** or **.xt** file. Note that the **.y** files are always read completely into memory.

If the argument **cache\_size** is not specified, all data is loaded into memory. Note: you should use caching only if the data does not fit into memory because caching uses the harddisc which will be much slower than memory access.

### 3.3 Learning Methods

By default MCMC inference is used for learning because MCMC is the most easiest to handle (no learning rate, no regularization). In LIBFM you can choose from the following learning methods: SGD, ALS, MCMC and SGDA. For all learning methods, the number of iterations `iter` has to be specified.

#### 3.3.1 Stochastic Gradient Descent (SGD)

Use `-method sgd` for parameter learning with SGD [4]. For stochastic gradient descent the following parameters have to be chosen:

- `-learn_rate`: the learning rate aka step size of SGD which should have a non-zero or positive value.
- `-regular`: the regularization parameters which should have zero or positive value. For SGD you can specify the regularization values the following way:
  - One value (`-regular value`): all model parameters use the same regularization value.
  - Three values (`-regular 'value0,value1,value2'`): 0-way interactions ( $w_0$ ) use `value0` as regularization, 1-way interactions ( $\mathbf{w}$ ) use `value1` and pairwise ones ( $V$ ) use `value2`.
  - No value: if the parameter `-regular` is not specified at all, this corresponds to no regularization, i.e. `-regular 0`.
- `-init_stdev`: the standard deviation of the normal distribution that is used for initializing the parameters  $V$ . You should use a non-zero, positive value here.

Please choose these arguments carefully as the prediction quality largely depends on good choices.

#### Example

```
./libFM -task r -train ml1m-train.libfm -test ml1m-test.libfm -dim '1,1,8' -iter 1000  
-method sgd -learn_rate 0.01 -regular '0,0,0.01' -init_stdev 0.1
```

#### 3.3.2 Alternating Least Squares (ALS)

Use `-method als` for parameter learning with ALS [8]. The following parameters have to be chosen:

- `-regular`: the regularization parameters which should have zero or positive value. For ALS you can specify the regularization values the following way:
  - One value (`-regular value`): all model parameters use the same regularization value.
  - Three values (`-regular 'value0,value1,value2'`): 0-way interactions ( $w_0$ ) use `value0` as regularization, 1-way interactions ( $\mathbf{w}$ ) use `value1` and pairwise ones ( $V$ ) use `value2`.
  - Group specific values (`-regular 'value0,value1g1,...,value1gm,value2g1,...,value2gm'`), i.e. for  $m$  groups there are  $1 + 2m$  many regularization values: if the input variables are grouped, for each group and 1-way and 2-way interaction an individual regularization value can be used.
  - No value: if the parameter `-regular` is not specified at all, this corresponds to no regularization, i.e. `-regular 0`.
- `-init_stdev`: the standard deviation of the normal distribution that is used for initializing the parameters  $V$ . You should use a non-zero, positive value here.

Please choose these arguments carefully as the prediction quality largely depends on good choices.

#### Example

```
./libFM -task r -train ml1m-train.libfm -test ml1m-test.libfm -dim '1,1,8' -iter 1000  
-method als -regular '0,0,10' -init_stdev 0.1
```

### 3.3.3 Markov Chain Monte Carlo (MCMC)

Use `-method mcmc` for parameter learning with MCMC [2]. The following parameters have to be chosen:

- `-init_stdev`: the standard deviation of the normal distribution that is used for initializing the parameters  $V$ . You should use a non-zero, positive value here.

Please choose this argument carefully as the speed of convergence depends on a good choice.

#### Example

```
./libFM -task r -train ml1m-train.libfm -test ml1m-test.libfm -dim '1,1,8' -iter 1000  
-method mcmc -init_stdev 0.1
```

### 3.3.4 Adaptive SGD (SGDA)

Use `-method sgda` for parameter learning with adaptive SGD [6]. With adaptive SGD, regularization values (per group and layer) are found automatically. You have to specify a validation set that is used to tune the regularization values:

- `-validation`: a data set containing training examples used as a validation set to tune the regularization values. This set should be distinct from the training dataset.
- `-learn_rate`: the learning rate aka step size of SGD which should have a non-zero or positive value.
- `-init_stdev`: the standard deviation of the normal distribution that is used for initializing the parameters  $V$ . You should use a non-zero, positive value here.

Please choose these arguments carefully as the prediction quality largely depends on good choices.

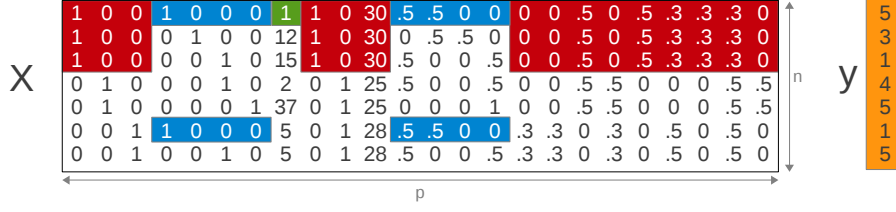
#### Example

```
./libFM -task r -train ml1m-train.libfm -test ml1m-test.libfm -dim '1,1,8' -iter 1000  
-method sgda -learn_rate 0.01 -init_stdev 0.1 -validation ml1m-val.libfm
```



## 4 Block Structure (BS) Extension\*

(a) Training Data (*Design Matrix*)



(b) Block Structure Representation of Design Matrix

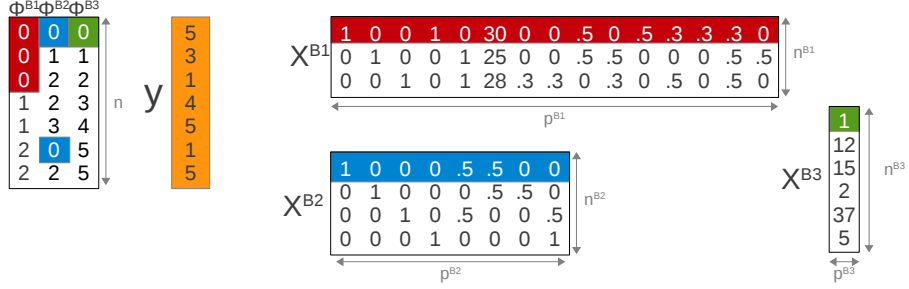


Figure 1: (a) LibFM data files (=representation of the design matrix  $X$ ) might contain large blocks of repeating patterns. (b) The BS extension of LIBFM allows to use a more compact representation of the data files where repeating patterns are described just once. (figure adapted from [7])

In relational settings, design matrices might contain large blocks of repeating patterns (see Fig. 1a). This can result in a very large design matrix which makes learning slow and uses lots of memory. The BS extension of LIBFM allows to define and make use of the block structure in design matrices. Both runtime and memory consumption will be linear in the BS data size instead of the original data size. For details about large design matrices from relational data, see [7].

### 4.1 Data Format

The BS extension allows to define blocks (e.g.  $B1$ ,  $B2$ ,  $B3$  in Fig. 1) and to use them in libFM. Each block definition consists of:

- The design matrix (libFM-file) of the block (e.g.  $X^{B1}$  in Fig. 1).
- The mapping from train (or test) cases to the rows in the block (e.g.  $\phi^{B1}$  in Fig. 1).
- Optional grouping of variables in the design matrix (compare section 3.2.2).

For each block, the following files are expected:

Filename	Description
<code>&lt;blockname&gt;.x</code>	The design matrix of the block in binary format (e.g. $X^{B1}$ in Fig. 1). The <code>convert</code> tool can be used to generate this from a file in LIBFM text format (see section 3.2.2).
<code>&lt;blockname&gt;.xt</code>	The transpose of <code>&lt;blockname&gt;.x</code> in binary format. The <code>transpose</code> tool can be used to generate this file from <code>&lt;blockname&gt;.x</code> (see section 3.2.2).
<code>&lt;blockname&gt;.train</code>	The mapping from train rows to block rows (e.g. $\phi^{B1}$ in Fig. 1). The expected file is in text format with as many lines as the training dataset (in the <code>--train</code> parameter). In each row, the file states the index <sup>4</sup> of the row in $X^{B1}$ to which this training case links. E.g. if the entry in the 5th row is 21, then the 5th train case uses the data in the 22nd row of $X^{B1}$ .
<code>&lt;blockname&gt;.test</code>	Same as <code>&lt;blockname&gt;.train</code> but now the lines in the test dataset are linked to $X^{B1}$ .
<code>&lt;blockname&gt;.groups</code>	Optional file for grouping predictor variables. The same format as in section 3.2.2 is expected.

## 4.2 Running libFM with BS data

The blocks are passed in the command line parameter `--relation`. Assuming two blocks (`rel.user` and `rel.item`) have been defined, the call would be:

```
./libFM -task r -train ml1m-train -test ml1m-test -dim '1,1,8' --relation rel.user,rel.item
```

Note that for each block the files listed above have to be present (i.e. `rel.user.x`, `rel.user.xt`, `rel.user.train`, `rel.user.test`, (`rel.user.groups`), `rel.item.x`, `rel.item.xt`, etc.).

## 4.3 Notes about BS Usage in libFM

- BS is only supported by MCMC and ALS/CD.
- Even when using BS, the `--train` and `--test` parameters are still mandatory and files have to be specified here. The LIBFM files passed in `--train` and `--test` can have predictor variables as well but might also be empty<sup>5</sup>. The files can be in binary or text format.
- The namespaces of variable ids in BS design matrices are distinct. E.g. a variable with index 7 in  $X^{B1}$  as well as in  $X^{B2}$  and the main train/test file  $X$  is considered to be different each. Internally, LIBFM adds the offset of the largest variable id of the preceding block  $X^{B_{i-1}}$  to each id in block  $X^{B_i}$ . Thus it is recommended to start indexing all blocks with variable id 0 to avoid wasting memory.
- The namespaces of the groups in BS files are distinct. Each group file can have groups starting from 0 – overlaps are resolved the same way as with predictor ids.
- If no group files are passed, each block is automatically assumed to have a different group.

<sup>4</sup>All indices are 0-based. I.e. the index of the first row is 0, the index of the second row is 1, etc.

<sup>5</sup>The target should be still specified. The predictor variables can be empty.

## 5 License

Please see `license.txt` for details. If you use LIBFM in your work, please cite the following paper:

```
@article{rendle:tist2012,  
  author = {Rendle, Steffen},  
  title = {Factorization Machines with {libFM}},  
  journal = {ACM Trans. Intell. Syst. Technol.},  
  issue_date = {May 2012},  
  volume = {3},  
  number = {3},  
  month = May,  
  year = {2012},  
  issn = {2157-6904},  
  pages = {57:1--57:22},  
  articleno = {57},  
  numpages = {22},  
  publisher = {ACM},  
  address = {New York, NY, USA},  
}
```

## References

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