

**mi update** — Ensure that mi data are consistent[Description](#)[Menu](#)[Syntax](#)[Remarks and examples](#)[Also see](#)

## Description

`mi update` verifies that `mi` data are consistent. If the data are not consistent, `mi update` reports the inconsistencies and makes the necessary changes to make the data consistent.

`mi update` can change the sort order of the data.

## Menu

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## Syntax

```
mi update
```

## Remarks and examples

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Remarks are presented under the following headings:

*Purpose of mi update*

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### Purpose of mi update

`mi update` allows you to

- change the values of existing variables, whether imputed, passive, regular, or unregistered;
- add or remove missing values from imputed variables (or from any variables);
- drop variables;
- create new variables;
- drop observations; and
- duplicate observations (but not add observations in other ways).

You can make any of or all the above changes and then type

```
. mi update
```

and `mi update` will handle making whatever additional changes are required to keep the data consistent. For instance,

```
. drop if sex==1
(75 observations deleted)
. mi update
(375 m>0 obs dropped because of dropped obs in m=0)
```

In this example, we happen to have five imputations and are working with `flongsep` data. We dropped 75 observations in  $m = 0$ , and that still left  $5 \times 75 = 375$  observations to be dropped in  $m > 0$ .

The messages `mi update` produces vary according to the style of the data because the changes required to make the data consistent are determined by the style. Had we been working with `flong` data, we might have seen

```
. drop if sex==1
(450 observations deleted)
. mi update
(system variable _mi_id updated because of changed number of obs)
```

With `flong` data in memory, when we dropped `if sex==1`, we dropped all  $75 + 5 \times 75 = 450$  observations, so no more observations needed to be dropped; but here `mi update` needed to update one of its system variables because of the change we made.

Had we been working with `mlong` data, we might have seen

```
. drop if sex==1
(90 observations deleted)
. mi update
(system variable _mi_id updated because of changed number of obs)
```

The story here is very much like the story in the `flong` case. In `mlong` data, dropping `if sex==1` drops the 75 observations in  $m = 0$  and also drops the incomplete observations among the 75 in  $m = 1, m = 2, \dots, m = 5$ . In this example, there are three such observations, so a total of  $75 + 5 \times 3 = 90$  were dropped, and because of the change, `mi update` needed to update its system variable.

Had we been using wide data, we might have seen

```
. drop if sex==1
(75 observations deleted)
. mi update
```

`mi update`'s silence indicates that `mi update` did nothing, because after dropping observations in wide data, nothing more needs to be done. We could have skipped typing `mi update` here, but do not think that way because changing values, dropping variables, creating new variables, dropping observations, or creating new observations can have unanticipated consequences.

For instance, in our data is variable `farmincome`, and it seems obvious that `farmincome` should be 0 if the person does not have a farm, so we type

```
. replace farmincome = 0 if !farm
(15 real changes made)
```

After changing values, you should type `mi update` even if you do not suspect that it is necessary. Here is what happens when we do that with these data:

```
. mi update
(12 m=0 obs now marked as complete)
```

Typing `mi update` was indeed necessary! We forgot that the `farmincome` variable was imputed, and it turns out that the variable contained missing in 12 nonfarm observations; `mi` needed to deal with that.

Running `mi update` is so important that `mi` itself is constantly running it just in case you forget. For instance, let's "forget" to type `mi update` and then convert our data to wide:

```
. replace farmincome = 0 if !farm
(15 real changes made)
. mi convert wide, clear
(12 m=0 obs now marked as complete)
```

The parenthetical message was produced because `mi convert` ran `mi update` for us. For more information on this, see [MI] [noupdate option](#).

## What mi update does

- `mi update` checks whether you have changed  $N$ , the number of observations in  $m = 0$ , and resets  $N$  if necessary.
- `mi update` checks whether you have changed  $M$ , the number of imputations, and adjusts the data if necessary.
- `mi update` checks whether you have added, dropped, registered, or unregistered any variables and takes the appropriate action.
- `mi update` checks whether you have added or deleted any observations. If you have, it then checks whether you carried out the operation consistently for  $m = 0, m = 1, \dots, m = M$ . If you have not carried it out consistently, `mi update` carries it out consistently for you.
- In the `mlong`, `flong`, and `flongsep` styles, `mi update` checks system variable `_mi_id`, which links observations across  $m$ , and reconstructs the variable if necessary.
- `mi update` checks that the system variable `_mi_miss`, which marks the incomplete observations, is correct and, if not, updates it and makes any other changes required by the change.
- `mi update` verifies that the values recorded in imputed variables in  $m > 0$  are equal to the values in  $m = 0$  when they are nonmissing and updates any that differ.
- `mi update` verifies that the values recorded in passive variables in  $m > 0$  are equal to the values recorded in  $m = 0$ 's complete observations and updates any that differ.
- `mi update` verifies that the values recorded in regular variables in  $m > 0$  equal the values in  $m = 0$  and updates any that differ.
- `mi update` adds any new variables in  $m = 0$  to  $m > 0$ .
- `mi update` drops any variables from  $m > 0$  that do not appear in  $m = 0$ .

## mi update is run automatically

As we mentioned before, running `mi update` is so important that many `mi` commands simply run it as a matter of course. This is discussed in [MI] [noupdate option](#). In a nutshell, the `mi` commands that run `mi update` automatically have a `noupdate` option, so you can identify them, and you can specify the option to skip running the update and so speed execution, but only with the adrenaline rush caused by a small amount of danger.

Whether you specify `noupdate` or not, we advise you to run `mi update` periodically and to always run `mi update` after dropping or adding variables or observations, or changing values.

## Also see

[MI] [Intro](#) — Introduction to `mi`

[MI] [noupdate option](#) — The `noupdate` option

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