

**joinby** — Form all pairwise combinations within groups

<a href="#">Description</a>	<a href="#">Quick start</a>	<a href="#">Menu</a>
<a href="#">Syntax</a>	<a href="#">Options</a>	<a href="#">Remarks and examples</a>
<a href="#">Acknowledgment</a>	<a href="#">Reference</a>	<a href="#">Also see</a>

## Description

`joinby` joins, within groups formed by *varlist*, observations of the dataset in memory with *filename*, a Stata-format dataset. By *join* we mean to form all pairwise combinations. *filename* is required to be sorted by *varlist*. If *filename* is specified without an extension, `.dta` is assumed.

If *varlist* is not specified, `joinby` takes as *varlist* the set of variables common to the dataset in memory and in *filename*.

Observations unique to one or the other dataset are ignored unless `unmatched()` specifies differently. Whether you load one dataset and join the other or vice versa makes no difference in the number of resulting observations.

If there are common variables between the two datasets, however, the combined dataset will contain the values from the master data for those observations. This behavior can be modified with the `update` and `replace` options.

## Quick start

Form pairwise combinations of observations from `mydata1.dta` in memory with those from `mydata2.dta` using all common variables and drop unmatched observations

```
joinby using mydata2
```

As above, but join on `v1`, `v2`, and `v3`

```
joinby v1 v2 v3 using mydata2
```

As above, but include unmatched observations only from `mydata2.dta` and add `_merge` indicating whether the variable was in both datasets or only the using dataset

```
joinby v1 v2 v3 using mydata2, unmatched(using)
```

As above, but include unmatched observations only from `mydata1.dta`

```
joinby v1 v2 v3 using mydata2, unmatched(master)
```

As above, but name the variable indicating the source of the observation `newv`

```
joinby v1 v2 v3 using mydata2, unmatched(master) _merge(newv)
```

Replace missing data in `mydata1.dta` with values from `mydata2.dta`

```
joinby v1 v2 v3 using mydata2, update
```

Replace missing and conflicting data in `mydata1.dta` with values from `mydata2.dta`

```
joinby v1 v2 v3 using mydata2, update replace
```

## Menu

Data > Combine datasets > Form all pairwise combinations within groups

## Syntax

```
joinby [varlist] using filename [, options]
```

<i>options</i>	Description
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### Options

When observations match:

<code>update</code>	replace missing data in memory with values from <i>filename</i>
<code>replace</code>	replace all data in memory with values from <i>filename</i>

When observations do not match:

<code>unmatched(none)</code>	ignore all; the default
<code>unmatched(both)</code>	include from both datasets
<code>unmatched(master)</code>	include from data in memory
<code>unmatched(using)</code>	include from data in <i>filename</i>
<code>_merge(varname)</code>	<i>varname</i> marks source of resulting observation; default is <code>_merge</code>
<code>nolabel</code>	do not copy value-label definitions from <i>filename</i>

*varlist* may not contain `strLs`.

## Options

### Options

`update` varies the action that `joinby` takes when an observation is matched. By default, values from the master data are retained when the same variables are found in both datasets. If `update` is specified, however, the values from the using dataset are retained where the master dataset contains missing.

`replace`, allowed with `update` only, specifies that nonmissing values in the master dataset be replaced with corresponding values from the using dataset. A nonmissing value, however, will never be replaced with a missing value.

`unmatched(none | both | master | using)` specifies whether observations unique to one of the datasets are to be kept, with the variables from the other dataset set to missing. Valid values are

<code>none</code>	ignore all unmatched observations (default)
<code>both</code>	include unmatched observations from the master and using data
<code>master</code>	include unmatched observations from the master data
<code>using</code>	include unmatched observations from the using data

`_merge(varname)` specifies the name of the variable that will mark the source of the resulting observation. The default name is `_merge` (`_merge`). To preserve compatibility with earlier versions of `joinby`, `_merge` is generated only if `unmatched` is specified.

`nolabel` prevents Stata from copying the value-label definitions from the dataset on disk into the dataset in memory. Even if you do not specify this option, label definitions from the disk dataset do not replace label definitions already in memory.

## Remarks and examples

The following, admittedly artificial, example illustrates joinby.

### ▷ Example 1

We have two datasets: `child.dta` and `parent.dta`. Both contain a `family_id` variable, which identifies the people who belong to the same family.

```
. use http://www.stata-press.com/data/r15/child
(Data on Children)
. describe
Contains data from http://www.stata-press.com/data/r15/child.dta
  obs:          5          Data on Children
  vars:         4          11 Dec 2016 21:08
  size:        30
```

variable name	storage type	display format	value label	variable label
family_id	int	%8.0g		Family ID number
child_id	byte	%8.0g		Child ID number
x1	byte	%8.0g		
x2	int	%8.0g		

Sorted by: family\_id

```
. list
```

	family~d	child_id	x1	x2
1.	1025	3	11	320
2.	1025	1	12	300
3.	1025	4	10	275
4.	1026	2	13	280
5.	1027	5	15	210

```
. use http://www.stata-press.com/data/r15/parent
(Data on Parents)
. describe
Contains data from http://www.stata-press.com/data/r15/parent.dta
  obs:          6          Data on Parents
  vars:         4          11 Dec 2016 03:06
  size:        84
```

variable name	storage type	display format	value label	variable label
family_id	int	%8.0g		Family ID number
parent_id	float	%9.0g		Parent ID number
x1	float	%9.0g		
x3	float	%9.0g		

Sorted by:

## 4 joinby — Form all pairwise combinations within groups

```
. list, sep(0)
```

	family~d	parent~d	x1	x3
1.	1030	10	39	600
2.	1025	11	20	643
3.	1025	12	27	721
4.	1026	13	30	760
5.	1026	14	26	668
6.	1030	15	32	684

We want to join the information for the parents and their children. The data on parents are in memory, and the data on children are posted at <http://www.stata-press.com>. `child.dta` has been sorted by `family_id`, but `parent.dta` has not, so first we sort the parent data on `family_id`:

```
. sort family_id
. joinby family_id using http://www.stata-press.com/data/r15/child
. describe
```

Contains data

```
obs:      8                      Data on Parents
vars:     6
size:    136
```

variable name	storage type	display format	value label	variable label
family_id	int	%8.0g		Family ID number
parent_id	float	%9.0g		Parent ID number
x1	float	%9.0g		
x3	float	%9.0g		
child_id	byte	%8.0g		Child ID number
x2	int	%8.0g		

Sorted by: family\_id

Note: Dataset has changed since last saved.

```
. list, sepby(family_id) abbrev(12)
```

	family_id	parent_id	x1	x3	child_id	x2
1.	1025	12	27	721	1	300
2.	1025	12	27	721	4	275
3.	1025	12	27	721	3	320
4.	1025	11	20	643	3	320
5.	1025	11	20	643	1	300
6.	1025	11	20	643	4	275
7.	1026	14	26	668	2	280
8.	1026	13	30	760	2	280

1. `family_id` of 1027, which appears only in `child.dta`, and `family_id` of 1030, which appears only in `parent.dta`, are not in the combined dataset. Observations for which the matching variables are not in both datasets are omitted.
2. The `x1` variable is in both datasets. Values for this variable in the joined dataset are the values from `parent.dta`—the dataset in memory when we issued the `joinby` command. If we had `child.dta` in memory and `parent.dta` on disk when we requested `joinby`, the values for `x1`

would have been those from `child.dta`. Values from the dataset in memory take precedence over the dataset on disk.



## Acknowledgment

`joinby` was written by Jeroen Weesie of the Department of Sociology at Utrecht University, The Netherlands.

## Reference

Baum, C. F. 2016. *An Introduction to Stata Programming*. 2nd ed. College Station, TX: Stata Press.

## Also see

[D] [append](#) — Append datasets

[D] [cross](#) — Form every pairwise combination of two datasets

[D] [fillin](#) — Rectangularize dataset

[D] [merge](#) — Merge datasets

[D] [save](#) — Save Stata dataset

[U] [22 Combining datasets](#)