

rwolf2 Implementation and Flexible Syntax

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Abstract

This is a short document that details the updated more flexible syntax of the `rwolf2` Stata command. It lays out the syntax, shows how this syntax is used to replicate the previous implementations in the original `rwolf` command, and describes cases where it allows for a much easier implementation of the Romano & Wolf multiple hypothesis correction in Stata.

Introduction

`rwolf2` is a new flexible implementation of the Romano and Wolf (2005a,b, 2016) multiple hypothesis correction written for Stata. A previous implementation of this procedure is available as `rwolf` (Clarke, 2016). Both are available for download from Stata's SSC using:

```
ssc install ado
```

where `ado` should be replaced with `rwolf` or `rwolf2`. Both return the same results (provided the same seed is set) when the same multiple hypothesis correction is implemented. However `rwolf2` has a much more flexible syntax (in line with Stata's `sureg` syntax) implying that any type of estimation procedures can be used, and these estimation procedures can be mixed within a single multiple correction procedure, provided that they return a point estimate and standard error. This includes standard Stata estimation procedures such as `reg`, `xtreg`, `areg` and so forth, but also other user-written libraries such as `rdrobust` (Calonico et al., 2017), `ivreg2` (Baum et al., 2002) and `reghdfe` (Correia, 2014) among others.

In this document I provide a brief description of the new more flexible syntax, document its implementation, and show that it replicates exactly the results of `rwolf` from examples shown in Clarke, Romano,

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and Wolf (2020). I note the things which can now be implemented very easily (in one line) which were previously complicated to implement in `rwolf`.

New Syntax

The syntax in `rwolf2` allows for the concatenation of all multiple tests to be implemented as a series of estimation commands, and requires that an option indicates the independent variables of interest (for which the correction should be implemented). Specifically, the syntax is:

```
rwolf2 (model1 depvar1 varlist1 [if] [in] [weight], [options1])
      (model2 depvar2 varlist2 [if] [in] [weight], [options2])
      ...
      (modelN depvarN varlistN [if] [in] [weight], [optionsN]),
indepvars(vars1, vars2, ..., varsN) [other_options]
```

Above `model1`, `model2`, ..., `modelN` refer to estimation commands, `depvar1`, `depvar2`, ..., `depvarN` refer to dependent variables and `varlist1`, `varlist2`, ..., `varlistN` refer to independent variables in each model. These independent variables can be syntax structures such as `(endog=iv) control` in the case that the model is `ivregress`. The `indepvars()` option specifies lists of independent variables in each model for which the Romano-Wolf correction is required. The help file of the command provides a full description of its syntax and implementation. As an example, a simple implementation of the command may be:

```
rwolf2 (reg y1 Treat1 Treat2, robust)
      (reg y2 Treat1 Treat2, robust),
indepvars(Treat1 Treat2, Treat1 Treat2)
```

where corrections for four tests are made (two treatment variables with two regression outcomes). However, more complicated cases can be easily incorporated such as:

```
rwolf2 (regress y1 Treat1 control1a control1b, robust)
      (ivregress 2sls y2 (Treat1=Assignment1) control2, robust),
indepvars(Treat1, Treat1)
```

In this second option, a correction is made for two multiple hypothesis tests (on the variable `Treat1` in both cases), but note that one model is estimated by IV, while the other is estimated by linear regression. Any

other combinations can be easily implemented. A complicated example may include regression discontinuity models, IV models high-dimensional fixed-effect models, and so forth. For example:

```
rwolf2 (regress y1 Treat1 control1a control1b, robust)
      (ivreg2 y2 (Treat1=Assignment1) control2, robust),
      (rdrobust y3 runningvar, covs(control3)),
      (reghdfe y4 Treat1 Treat2, absorb(state year)),
indepvars(Treat1, Treat1, RD_Estimate, Treat1 Treat2)
```

where an OLS, IV, regression discontinuity and high-dimensional fixed effect models are all implemented, and multiple hypothesis corrections are implemented for 5 variables of interest (specified in the indepvars option). This multiple hypothesis correction was previously complicated to implement with `rwolf`, but is very easy with `rwolf2`.

Changes

The `rwolf2` command makes it much easier than in `rwolf` to:

- Include any estimation commands for underlying models, as well as different estimation commands between models, including user written commands like `reghdfe`, `ivreg2`, `rdrobust` and so forth.
- Adjust for multiple and differing treatment variables within and across models
- Include different controls in each model (or estimate with different weighting schemes, estimation options, sub-samples, etc.)

Consistency Across Versions

In cases where `rwolf2` and `rwolf` can be implemented to conduct the same multiple hypothesis test, they will return exactly the same results, provided that the same seed is set. For example, see below for the IV regression example from Clarke et al. (2020). The first implementation uses `rwolf2` with the new syntax, while the second version implements the same tests with the old syntax of `rwolf`. The output is the same in each case.

```

. use "http://www.stata-press.com/data/r13/hsng", clear
(1980 Census housing data)

. #delimit ;
delimiter now ;
. rwolf2 (ivregress 2sls rent (hsngval=faminc i.region) pcturban)
>       (ivregress 2sls popden (hsngval=faminc i.region) pcturban)
>       (ivregress 2sls popgrow (hsngval=faminc i.region) pcturban)
>       (ivregress 2sls hsng (hsngval=faminc i.region) pcturban), graph
> indepvars(hsngval, hsngval, hsngval, hsngval) reps(10000) seed(2) nodots;
Bootstrap replications (10000). This may take some time.

```

Romano-Wolf step-down adjusted p-values
Number of resamples: 10000

	Model p-value	Resample p-value	Romano-Wolf p-value
rent	0.0000	0.0169	0.0169
popden	0.0654	0.0845	0.0845
popgrow	0.0019	0.0107	0.0169
hsng	0.0236	0.0106	0.0460

```

. rwolf rent popden popgrow hsng, indepvar(hsngval) method(ivregress)
> iv(faminc i.region) reps(10000) graph nodots controls(pcturban) seed(2);
Bootstrap replications (10000). This may take some time.

```

Romano-Wolf step-down adjusted p-values

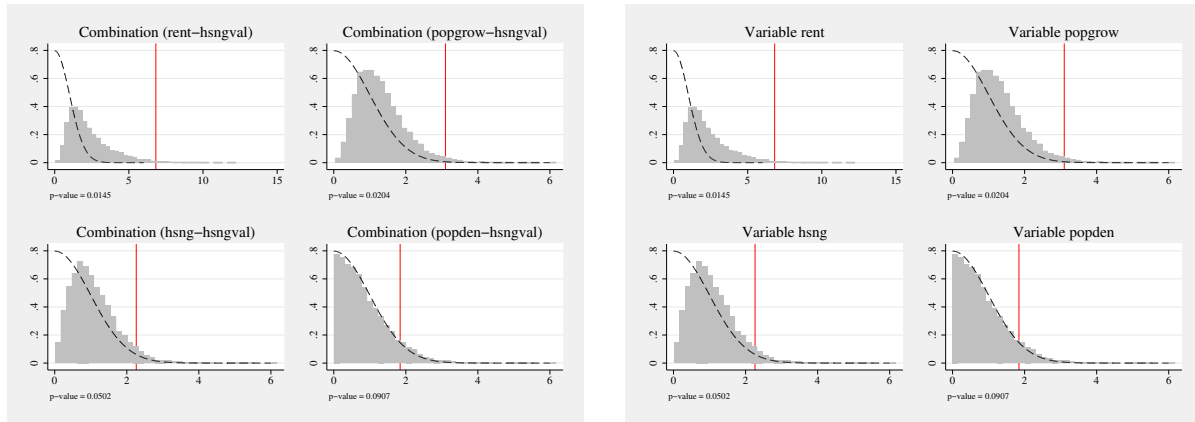
Independent variable: hsngval
Outcome variables: rent popden popgrow hsng

Number of resamples: 10000

Outcome Variable	Model p-value	Resample p-value	Romano-Wolf p-value
rent	0.0000	0.0169	0.0169
popden	0.0654	0.0845	0.0845
popgrow	0.0019	0.0107	0.0169
hsng	0.0236	0.0106	0.0460

The only change is in terms of graphical output if the graph option is used. While they are identical in terms of the displayed results, the new `rwolf2` output renames null distribution graphs more clearly for cases where dependent variables may be constant across models, so that both dependent and independent variable names are used to caption graphs.

Figure 1: Graphical Output from `rwolf2` and `rwolf`



(a) Output from `rwolf2`

(b) Output from `rwolf`

References

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