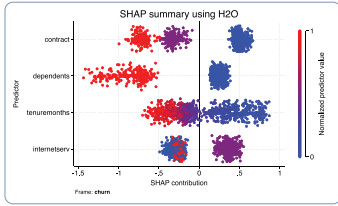
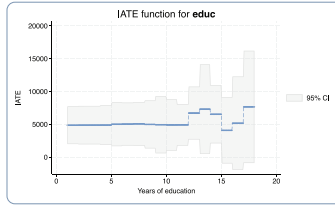


New in STATA® 19



Machine learning via H2O: Ensemble decision trees

Employ machine learning to uncover insights from the data when traditional statistical models fall short. Use gradient boosting machine (GBM) and random forest to perform regression or classification.



Conditional average treatment effects

Go beyond estimating overall treatment effects in your causal analysis to estimating individualized and group-specific ones. Compare different interventions and policies. Explore treatment-effects heterogeneity.

```
regress, absorb(z1 z2 z3)
xtreg, absorb(z1 z2 z3) fe
ivregress 2sls, absorb(z1 z2 z3)
```

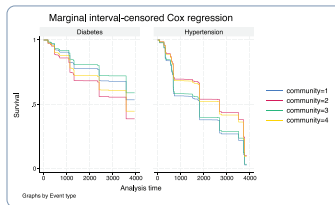
High-dimensional fixed effects

Absorb not just one but multiple high-dimensional categorical variables in your linear, fixed-effects linear, and instrumental-variables linear models, and enjoy remarkable speed gains!

Bayesian variable selection		MCMC iterations = 12,500	
Microsplit-shrinkage and Gibbs sampling	Mean = 2,200	MCMC sample size = 10,000	
Global-local shrinkage coefficient priors (Korachoson1)	Number of obs = 423	Acceptance rate = .8633	
	Efficiency ratio = .1516	avg = .44	
	max = .1		
Log marginal-likelihood = -475.56099			
		Equal-tailed	
		Posterior mean	
diabetes	Mean	Std. dev.	95% conf. interval
seumal	-.236779	.0671549	-.3102866 -0.1631979
bmi	-.3284698	.0414561	-.4004504 -.247525
bp	-.1909207	.0411339	-.2598225 -.1218276
sex	-.1278497	.0398828	-.2046245 -.0510712
seumal	-.125876	.1273556	-.0327762 -.4472664
seumal	-.0282577	.0782906	-.101311 -0.0551919
seumal	-.048893	.0733476	-.1749584 -.2131995
seumal	-.0182026	.1207191	-.0223902 -.0140119
seumal	-.0300323	.0361144	-.0506945 -.0109577
age	-.001922	.1282981	-.0222981 -.0014765

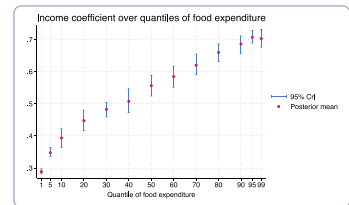
Bayesian variable selection

Select variables in a linear regression and account for variable-selection uncertainty with Bayesian variable selection. Choose between global-local shrinkage and spike-and-slab priors for regression coefficients, and perform Bayesian inference.



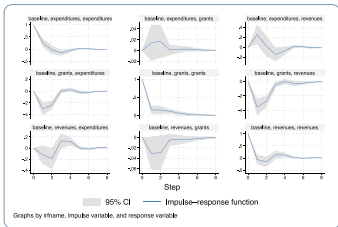
Interval-censored multiple-event Cox model

Need to analyze event times from multiple events, such as the onsets of diabetes and hypertension?
Don't know the exact event times?
Use the marginal interval-censored multiple-event Cox model.



Bayesian quantile regression

Use Bayesian quantile regression to obtain full posterior distributions of quantile regression coefficients for comprehensive inference, including model-based "standard errors".



Panel-data VAR model

Fit a panel-data vector autoregressive (VAR) model to analyze the trajectories of related variables when you observe multiple units or panels over time.

```
xtreg, re + xtreg, fe = xtreg, cre
```

Correlated random-effects model

Want coefficient estimates of time-invariant covariates in your panel-data model? Fit an RE model. Want to allow for correlation between covariates and unobserved panel-level effects? Fit an FE model. Want both? With **xtreg, cre**, you can now fit a CRE model.

Bayesian bootstrap		Observation prior: Inorganic			
Linear regression		Number of obs = 74			
		Replications = 50			
		Wald chi2(1) = 189.23			
		Prob > chi2 = 0.0000			
		R-squared = 0.6515			
		Adj R-squared = 0.6487			
		Root MSE = 2.4388			
	Observed	Bayesian	Normal-based		
reg	coefficient	std. err.	z	P> z	[95% conf. interval]
weight	-.0000007	.000575	-18.45	0.000	-.0021356 -.0008657
_cons	39.44028	1.860948	21.13	0.000	35.78113 43.09943

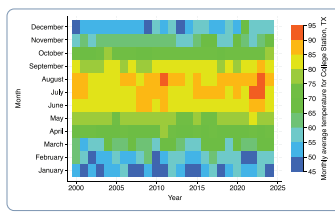
Bayesian bootstrap

Perform Bayesian bootstrap to obtain more precise parameter estimates in small samples and incorporate prior information when sampling observations. Use it with official commands or community-contributed commands.

cfprogsa rest pcturban (hanyal = fmainc l.region, interact(fmainc))						
Control-function linear regression						
Number of obs = 50						
Wald chi2(1) = 85.18						
Prob > chi2 = 0.0000						
R-squared = 0.9963						
Root MSE = 22.2851						
Endogenous variable model:						
Slinear hanyal						
vars	Coefficients	Std. err.	z	P> z	[95% conf. interval]	
rest	hanyal	2.155381	.3437284	6.27	0.000	1.461686 2.849076
	pcturban	.4794597	.2362242	2.03	0.042	-.0164689 .9424596
	_cons	86.15959	13.49558	7.08	0.000	59.57261 125.263
r.rest	cf(hanyal)	10.66765	3.619462	2.95	0.003	3.573873 17.76163
cf(hanyal)	fmainc	-.5610651	.1743049	-3.22	0.001	-.9026665 -.2184338

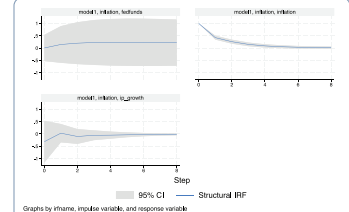
Control-function linear and probit models

Fit control-function linear and probit models, with continuous, binary, fractional, and count endogenous variables. Easily test for endogeneity.



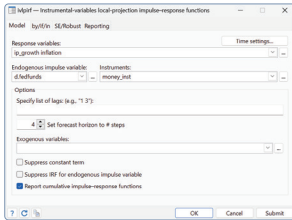
Graphics: Bar graph CIs, heat maps, ...

Enjoy exciting new graphical features such as heat maps, spike and capped-spike range plots, bar graphs with CIs, and more!



SVAR models via instrumental variables

Use instruments instead of short-run constraints to estimate dynamic causal effects.



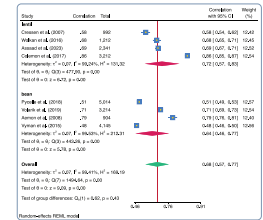
IV local-projection IRFs

Account for endogeneity when using local projections to estimate dynamic causal effects.

Classes	BIC	DMR (P<DMR)	Class	marginal probabilities (SE)
1	1,189.80	1.00 (0.00)		
2	1,057.31	75.55 (0.001)	0.72 (0.06)	0.28 (0.06)
3	1,081.86	2.25 (0.087)	0.16 (0.33)	0.63 (0.194) 0.21 (0.37)

Latent class model-comparison statistics

Easily compare latent class models with varying numbers of latent classes. Construct and export publication-quality tables comparing models.



Meta-analysis for correlations

Perform meta-analysis for correlations just like you already do for two-sample means and proportions. Use all standard features such as forest plots and subgroup analysis.

```
. estat weakrobust
Weak-instrument-robust test
Model VCE: Robust

(1) hsgval = 0

Cond. likelihood-ratio (CLR) test = 5.48
Prob > CLR = 0.0253

Notes: CLR test reported by default because
model is overidentified.
p-value computed by simulation
(25,000 replications).
```

Inference robust to weak instruments

Do you have weak instruments in your instrumental-variables regression? Use tests robust to weak instruments to perform reliable inference on endogenous regressors.

```
RE? FE? CRE?

. xtreg ...
. mundlak
```

Mundlak specification test

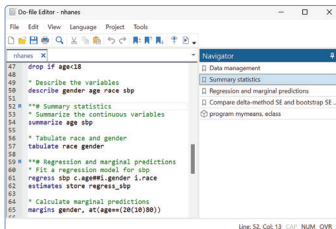
Use the Mundlak specification test to choose between random-effects and fixed-effects or correlated random-effects models, even with cluster-robust, bootstrap, or jackknife standard errors.

Bayesian asym. Laplace (quantile) regression	NMC iterations =	15,000
Random-walk Metropolis-Hastings sampling	Burn-in =	5,000
	NMC sample size =	10,000
	Number of obs =	887
	Acceptance ratio =	0.339
	Efficiency: min =	0.1428
	avg =	0.037
	max =	0.034

Log marginal-likelihood	Mean	Std. dev.	MCSE	Median	195% cred. interval
math3_q25					
math3	-781409	-0.033464	-0.001375	-7818769	-7714074
u1	0	0	0	1	1
const	27.11278	-0.114259	-0.0442	27.09278	26.33384
math3_q75					
math3	-6021812	-0.0285305	-0.013	-6029619	-344931
u2	0	0	0	1	1
const	34.21954	-0.2754823	-0.01821	34.22474	33.46647
math3	-1.507848	-0.073112	-0.00087	-1.507952	-1.488466
var_u1	7.729257	0.2641174	-0.01128	7.5467193	6.529766
var_u2	2.256973	-0.7847156	-0.04895	2.153597	1.128309

Bayesian asymmetric Laplace model

Go beyond classic quantile regression by fitting Bayesian simultaneous, multilevel, and nonlinear quantile regression models.



Do-file Editor: Autocompletion, templates, ...

Navigator panel, file templates, code folding improvements, word and selection highlighting, more autocompletion, temporary bookmarks, and much more.

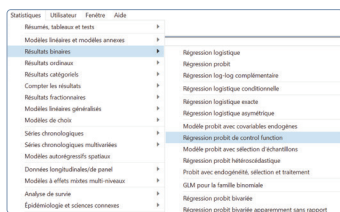
Tables: Easier tabulations, exporting, ...

Create a table, customize it with a title and notes, and export it in one command. Easily collect and customize tabulations with measures of association, tabulations of survey data, and ANOVA tables.

```
frames modify, add(myframe)
frames modify, drop(myframe)
```

Multiple datasets: Modify a set of frames

If you work with multiple datasets in memory or frames, you can now modify a frameset file without loading it into memory: add frames in memory to it or drop frames from it.



Stata in French

All of Stata's interface—all menus and all dialogs—is now available in French.

More

- Alternative at-risk table for survival graphs
- PyStata enhancements
- Log-scale analysis of bioequivalence
- Robust standard errors for VAR models
- Bayesian predictions in user-defined evaluators
- Mata functions for least-squares solvers, labels, and more

