Just tired of endless loops!
or `parallel`: Stata module for parallel computing

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Agenda

- Both computation speeds and size of data are ever increasing
- Often our work is easily broken down into independent chunks
- But, implementing parallel computing even for these "embarrassingly parallel" problems is not easy, most of this due to lack of (user-friendly) statistical computing tools.
- `parallel` aims to make a contribution to these issues.
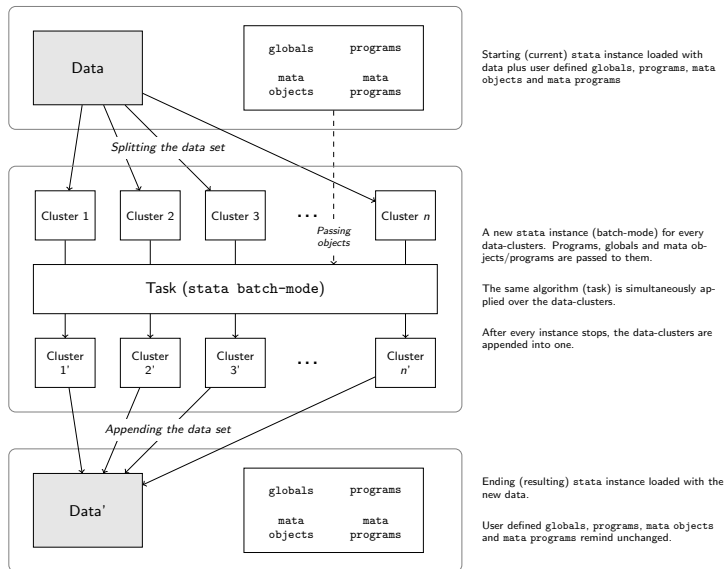
- ▶ NB: StataMP is available but only parallelizes some internal commands. Therefore a user-provided solution is needed.
- ▶ Inspired by the R package "snow" (several other examples exists: StataMP, HTCondor, Matlab's Parallel Toolbox, etc.)
- ▶ Is designed to be used in multiprocessor machines (e.g. simultaneous multi-threading, multiple cores, multiple sockets).
- ▶ It implements parallel computing methods through an OS's shell scripting (using Stata in batch mode).
- ▶ On Windows we have a compiled plugging allowing
  - ▶ Functionality when the parent Stata is in batch-mode
  - ▶ Seamless user experience by launching the child programs in a hidden desktop (otherwise GUI for each steals focus)
- ▶ Depending on the task, can reach near to (or over) linear speedups proportional to the number of physical cores of the computer.
- ▶ Thus having a quad-core computer can lead to a 400% speedup.

What is and how does it work
How does it work?

Starting (current) stata instance loaded with data plus user defined globals, programs, mata objects and mata programs

A new stata instance (batch-mode) for every data-clusters. Programs, globals and mata objects/programs are passed to them.

The same algorithm (task) is simultaneously applied over the data-clusters.

After every instance stops, the data-clusters are appended into one.

Ending (resulting) stata instance loaded with the new data.

User defined globals, programs, mata objects and mata programs remind unchanged.

- Method is *split-apply-combine* like MapReduce
- Straightforward usage when there is observation or group-level work
- If each iteration needs the entire dataset, then can be reformulated as
  1. Save real dataset to file
  2. Generate in-memory data as the list of tasks. For example,
     - List of seeds for each bootstrap resampling
     - List of parameter values for simulations
  3. Child process reads its list of tasks, loads the real dataset, and does it's own looping over tasks
- If the list of tasks is not initially definable then the "nodata" alternative mechanism allows for more flexibility.

When the user enters

```
parallel:  gen x2 = x*x
```

parallel takes the command and writes something like this

```
  cap clear all
  cd ~
1 set seed 34815
  set memory 16777216b
  cap set maxvar 5000
  cap set matsize 400
2 local pll_instance 1
  local pll_id efcql2tspr
  capture {
  noisily {
3 use __pllefcql2tsprdataset if _efcql2tsprcut == 1
  gen x2 = x*x
  }
  }
4 save __pllefcql2tsprdta1, replace
  local result = _rc
  cd ~
5 mata: write_diagnosis(st_local("result"),
  >"__pllefcql2tsprfinito1")
```

```
  cap clear all
  cd ~
1 set seed 98327
  set memory 16777216b
  cap set maxvar 5000
  cap set matsize 400
2 local pll_instance 2
  local pll_id efcql2tspr
  capture {
  noisily {
3 use __pllefcql2tsprdataset if _efcql2tsprcut == 2
  gen x2 = x*x
  }
  }
4 save __pllefcql2tsprdta2, replace
  local result = _rc
  cd ~
5 mata: write_diagnosis(st_local("result"),
  >"__pllefcql2tsprfinito2")
```

Serial fashion

```
do mydofile.do
```

Parallel fashion

```
parallel do mydofile.do
```

Figure: mydofile.do

```
local size = _N
forval i=1/`size' {
        qui replace x = ///
            1/sqrt(2*`c(pi)')*exp(-(x^2/2)) in `i'
}
```

Table: Serial replacing using a loop on a Linux Server (16 clusters)

	100,000	1,000,000	10,000,000
CPU	1.43	16.94	144.68
Total	0.34	3.20	12.49
Setup	0.00	0.00	0.00
Compute	0.32	3.07	11.54
Finish	0.02	0.12	0.95
Ratio (compute)	4.50	5.51	12.53
Ratio (total)	4.22 (26%)	5.30 (30%)	11.58 (72%)

Tested on a Intel Xeon X470 (hexadeca-core) machine

Benchmarks
Monte Carlo simulation (Windows Machine)

Serial fashion

```
do myexperiment.do
```

Parallel fashion

```
parallel do myexperiment.do, nodata
```


Figure: myexperiment.do

```
local num_of_intervals = 50
if length("`pll_id'") == 0 {
    local start = 1
    local end = `num_of_intervals'
}
else {
    local ntot = floor(`num_of_intervals'/$PLL_CLUSTERS)
    local start = (`pll_instance' - 1)*`ntot' + 1
    local end = (`pll_instance')*`ntot'
    if `pll_instance' == $PLL_CLUSTERS local end = 10
}
local reps 10000
forval i=`start'/`end' {
    qui use census2, clear
    gen true_y = age
    gen x_factor = region
    sum x_factor, meanonly
    scalar xmu = r(mean)
    qui {
        gen y1 = .
        gen y2 = .
        local c = `i'
        set seed `c'
        simulate c=r(c) mu1=r(mu1) se_mu1 = r(se_mu1) ///
                 mu2=r(mu2) se_mu2 = r(se_mu2), ///
                 saving(`c1', replace) nodots reps(`reps'): ///
                 mcsimul1, c(`c')
    }
}
```

Table: Monte Carlo Experiment on a Windows Machine (4 clusters)

	2	4
CPU	111.49	114.13
Total	58.02	37.48
Setup	0.00	0.00
Compute	58.02	37.48
Finish	0.00	0.00
Ratio (compute)	1.92	3.04
Ratio (total)	1.92 (96%)	3.04 (76%)

Tested on a Intel i3 2120 (dual-core) machine

Benchmarks
Monte Carlo simulation (Unix Machine)

Serial fashion

```
do myexperiment.do
```

Parallel fashion

```
parallel do myexperiment.do, nodata
```

Table: Monte Carlo Experiment on a Linux Server (16 clusters)

	2	4	8	16
CPU	164.79	164.04	162.84	163.89
Total	69.85	34.28	19.00	10.78
Setup	0.00	0.00	0.00	0.00
Compute	69.85	34.28	19.00	10.78
Finish	0.00	0.00	0.00	0.00
Ratio (compute)	2.36	4.78	8.57	15.21
Ratio (total)	2.36 (118%)	4.78 (120%)	8.57 (107%)	15.21 (95%)

Tested on a Intel Xeon X470 (hexadeca-core) machine

Serial fashion

```
reshape wide tipsolic rutemp opta derecho ngiros, ///
        i(id) j(time)
```

Parallel fashion

```
parallel, by(id) :reshape wide tipsolic rutemp opta derecho ngiros, ///
        i(id) j(time)
```

Table: Reshaping wide a large database on a Linux Server (8 clusters)

	100,000	1,000,000	5,000,000
CPU	5.51	72.70	392.97
Total	2.33	17.46	86.44
Setup	0.00	0.00	0.00
Compute	1.83	12.42	57.93
Finish	0.50	5.04	28.51
Ratio (compute)	3.01	5.85	6.78
Ratio (total)	2.37 (29%)	4.16 (52%)	4.55 (57%)

Tested on a Intel Xeon X470 (hexadeca-core) machine

Benchmarks
Bootstrap with `parallel bs`

```
sysuse auto, clear expand 10

// Serial fashion
bs, rep($size) nodots: regress mpg weight gear foreign

// Parallel fashion parallel setclusters $number_of_clusters
parallel bs, rep($size) nodots: regress mpg weight gear foreign
```

Problem size	Serial	2 Clusters	4 Clusters
1,000	2.93s	1.62s	1.09s
	×2.69	×1.48	×1.00
2,000	5.80s	3.13s	2.03s
	×2.85	×1.54	×1.00
4,000	11.59s	6.27s	3.86s
	×3.01	×1.62	×1.00

Table: Absolute and relative computing times for each run of a basic bootstrap problem. For each given problem size, the first row shows the time in seconds, and the second row shows the relative time each method took to complete the task relative to using parallel with four clusters. Each cell represents a 1,000 runs.

Benchmarks
Simulations with `parallel sim`

```
prog def mysim, rclass
    // Data generating process
    drop _all
    set obs 1000
    gen eps = rnormal()
    gen X = rnormal()
    gen Y = X*2 + eps

    // Estimation
    reg Y X
    mat def ans = e(b)
    return scalar beta = ans[1,1]
end

// Serial fashion
simulate beta=r(beta), reps($size) nodots: mysim

// Parallel fashion
parallel setclusters $number_of_clusters
parallel sim, reps($size) expr(beta=r(beta)) nodots: mysim
```

Problem size	Serial	2 Clusters	4 Clusters
1000	2.19s	1.18s	0.73s
	×3.01	×1.62	×1.00
2000	4.36s	2.29s	1.33s
	×3.29	×1.73	×1.00
4000	8.69s	4.53s	2.55s
	×3.40	×1.77	×1.00

Table: Absolute and relative computing times for each run of a simple Monte Carlo exercise. For each given problem size, the first row shows the time in seconds, and the second row shows the relative time each method took to complete the task relative to using parallel with four clusters. Each cell represents a 1,000 runs.

Code for replicating this is available at
https://github.com/gvegayon/parallel

Syntax and Usage

Setup

parallel setclusters # [, <u>f</u>orce]

By syntax

parallel [, by(*varlist*) <u>p</u>rograms <u>m</u>ata <u>s</u>eeds(*string*) <u>r</u>andtype(*random.org|datetime*)
<u>p</u>rocessors(*integer*) <u>nod</u>ata]: *stata_cmd*

Do syntax

parallel do *filename*
[, by(*varlist*) <u>p</u>rograms <u>m</u>ata <u>s</u>eeds(*string*) <u>r</u>andtype(*random.org|datetime*)
<u>p</u>rocessors(*integer*) <u>nod</u>ata]

Bootstrap syntax

parallel bs [, by(*varlist*) <u>p</u>rograms <u>m</u>ata <u>s</u>eeds(*string*) <u>r</u>andtype(*random.org|datetime*)
<u>p</u>rocessors(*integer*) <u>nod</u>ata]: *stata_cmd*

Simulations syntax

parallel sim [, by(*varlist*) <u>p</u>rograms <u>m</u>ata <u>s</u>eeds(*string*) <u>r</u>andtype(*random.org|datetime*)
<u>p</u>rocessors(*integer*) <u>nod</u>ata]: *stata_cmd*

Append syntax

parallel append [, by(*varlist*) <u>p</u>rograms <u>m</u>ata <u>s</u>eeds(*string*) <u>r</u>andtype(*random.org|datetime*)
<u>p</u>rocessors(*integer*) <u>nod</u>ata]: *stata_cmd*

`parallel suits ...`

- ► Monte-Carlo simulation.
- ► Extensive nested control flow (loops, while, ifs, etc.).
- ► Bootstrapping/Jackknife.
- ► Simulations in general.

`parallel doesn't suit ...`

- ► (already) fast commands.
- ► Regressions, ARIMA, etc.
- ► Linear Algebra.
- ► Whatever StataMP does better.
- ► (Currently) Tasks that already take up all of RAM.

Concluding Remarks

- Brings parallel computing to many more commands than StataMP
- its major strengths/advantages are in simulation models and non-vectorized operations such as control-flow statements.
- Depending on the proportion of the algorithm that can be de-serialized, it is possible to reach near to linear scale speedups.
- `parallel` can be incorporated into other user commands as an optional speedup.
- Caveat: Has not been tested on Stata 15.
- Contribute, find help, and report bugs at `http://github.com/gvegayon/parallel`

Thank you very much!

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