

# **gputools: an R package for GPU computing**

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# Outline

- Contents of gputools
- Usage
- Performance
- Other R packages for the GPU

# CONTENTS OF gputools

A handful of selected R functions implemented with CUDA C for use on a GPU:

- Choose your device:

gputools function	CPU analog	Same usage?
chooseGpu()	none	NA
getGpuId()	none	NA

- Linear algebra:

gputools function	CPU analog	Same usage?
gpuDist()	dist()	no
gpuMatMult()	%*% operator	no
gpuCrossprod()	crossprod()	yes
gpuTcrossprod()	tcrossprod()	yes
gpuQr()	qr()	almost
gpuSolve()	solve()	no
gpuSvd()	svd()	almost

- Simple model fitting:

<b>gputools function</b>	<b>CPU analog</b>	<b>Same exact usage?</b>
gpuLm()	lm()	yes
gpuLsfit()	lsfit()	yes
gpuGlm()	glm()	yes
gpuGlm.fit()	glm.fit()	yes

- Hypothesis testing:

<b>gputools function</b>	<b>CPU analog</b>	<b>Same exact usage?</b>
gpuTtest()	t.test()	no
getAucEstimate()	???	???

- Other routines:

<b>gputools function</b>	<b>CPU analog</b>	<b>Same exact usage?</b>
gpuHclust()	hclust()	no
gpuDistClust()	hclust(dist())	no
gpuFastICA()	fastICA() (fastICA package)	yes
gpuGranger()	grangertest() (lmtest package)	no
gpuMi()	???	???
gpuSvmPredict()	See <a href="http://www.jstatsoft.org/v15/i09/paper">www.jstatsoft.org/v15/i09/paper</a>	no
gpuSvmTrain()	See <a href="http://www.jstatsoft.org/v15/i09/paper">www.jstatsoft.org/v15/i09/paper</a>	no

## **getAucEstimate()**

Estimates the area under a receiver operating characteristic (ROC) curve.

Used to evaluate the performance of a hypothesis test in a multiple testing scenario.

Reference:

Hand, David J. and Till, Robert J. (2001). A simple generalisation of the area under the ROC curve for multiple class classification problems. Machine Learning. 45, 171-186.

## **gpuHclust()**

Performs hierarchical clustering on a set of points.

The distances among the points must be given in an object of class "dist".

## **gpuDistClust()**

Given a set of points, computes all pairwise distances and then performs hierarchical clustering on the points.

Both steps are done on the GPU.

# gpuFastICA

Performs Independent Component Analysis (ICA) and Projection Pursuit.

ICA, like principle component analysis, is a linear decomposition of a design matrix.

The authors of fastICA claim that, unlike PCA, ICA “unmixes” the underlying sources of variability in the data by assuming a non-Gaussian structure.

This function is exactly like the ICA implementation in the **fastICA** package except that the **gpuTools** version uses **gpuSvd()** instead of **svd()**.

References:

A. Hyvärinen and E. Oja (2000) Independent Component Analysis: Algorithms and Applications, *Neural Networks*, 13(4-5):411-430. <http://www.cis.hut.fi/aapo/>

A. Hyvärinen. Independent Component Analysis: Recent Advances. *Philosophical Transactions of the Royal Society A*, in press. <http://www.cs.helsinki.fi/u/ahyvarin/papers/PTRSA12.pdf>.

## **gpuGranger()**

Performs the Granger Causality Test, which tests how well one time series forecasts another.

Reference:

Hacker R.S. and Hatemi-J A. (2006) "Tests for causality between integrated variables using asymptotic and bootstrap distributions: theory and application", Applied Economics, Vol. 38(13), pp. 1489-1500.

## **gpuMi()**

Estimates the mutual information for pairs of vectors using a B spline approach.

Reference:

Carten O. Daub, Ralf Steuer, Joachim Selbig, and Sebastian Kloska. 2004. Estimating mutual information using B-spline functions - an improved similarity measure for analysing gene expression data. BMC Bioinformatics. 5:118. Available from <http://www.biomedcentral.com/1471-2105/5/118>

# **gpuSvmPredict()**

Classifies points in a data set using a support vector machine.

In machine learning, support vector machine (SVM) is a learning model used for classification and regression.

Reference:

Carpenter, Austin. cuSVM: a cuda implementation of support vector classification and regression.

<http://patternsonascreen.net/cuSVM.html>

## **gpuSvmTrain()**

Trains a support vector machine.

Reference:

Carpenter, Austin. cuSVM: a cuda implementation of support vector classification and regression.  
<http://patternsonascreen.net/cuSVM.html>

## USAGE

- gputools is already installed on impact1.stat.iastate.edu, ready to load with `library(gputools)` in R.
- For other GPU systems, download gputools from CRAN with a simple `install.packages("gputools")` in R.
  - **WARNING:** installation will fail on non-GPU systems since the CUDA C compiler doesn't exist
- Documentation:
  - <http://brainarray.mbnl.med.umich.edu/Brainarray/Rgpgpu/>
  - <http://cran.r-project.org/web/packages/gputools/index.html>
  - <http://cran.r-project.org/web/packages/gputools/gputools.pdf>
- Requirements:
  - R ( $\geq$  version 2.8.0)
  - Nvidia's CUDA toolkit ( $\geq$  version 2.3)

## MANAGING YOUR DEVICES: chooseGpu() AND getGpuId()

Impact1 has four GPUs, each with a unique index from 0 to 3. To see this for yourself, log into impact1 and run the following:

```
[landau@impact1 ~]$ cd /usr/local/NVIDIA_GPU_Computing_SDK/C/bin/linux/release  
[landau@impact1 release]$ ./deviceQuery
```

Here are some pieces of the (quite verbose) output of  
`./deviceQuery`:

```
[deviceQuery] starting...
```

```
./deviceQuery Starting...
```

```
CUDA Device Query (Runtime API) version (CUDART static linking)
```

```
Found 4 CUDA Capable device(s)
```

```
Device 0: "Tesla M2070"
```

CUDA Driver Version / Runtime Version	4.1 / 4.1
CUDA Capability Major/Minor version number:	2.0
Total amount of global memory:	5375 MBytes (5636554752 bytes)
(14) Multiprocessors x (32) CUDA Cores/MP:	448 CUDA Cores
GPU Clock Speed:	1.15 GHz
Memory Clock rate:	1566.00 Mhz
Memory Bus Width:	384-bit
L2 Cache Size:	786432 bytes

**Device 1: "Tesla M2070"**

CUDA Driver Version / Runtime Version	4.1 / 4.1
CUDA Capability Major/Minor version number:	2.0
Total amount of global memory:	5375 MBytes (5636554752 bytes)
(14) Multiprocessors x (32) CUDA Cores/MP:	448 CUDA Cores
GPU Clock Speed:	1.15 GHz
Memory Clock rate:	1566.00 Mhz
Memory Bus Width:	384-bit
L2 Cache Size:	786432 bytes

**Device 2: "Tesla M2070"**

CUDA Driver Version / Runtime Version	4.1 / 4.1
CUDA Capability Major/Minor version number:	2.0
Total amount of global memory:	5375 MBytes (5636554752 bytes)
(14) Multiprocessors x (32) CUDA Cores/MP:	448 CUDA Cores
GPU Clock Speed:	1.15 GHz
Memory Clock rate:	1566.00 Mhz
Memory Bus Width:	384-bit
L2 Cache Size:	786432 bytes

Device 3: "Tesla M2070"	
CUDA Driver Version / Runtime Version	4.1 / 4.1
CUDA Capability Major/Minor version number:	2.0
Total amount of global memory:	5375 MBytes (5636554752 bytes)
(14) Multiprocessors x (32) CUDA Cores/MP:	448 CUDA Cores
GPU Clock Speed:	1.15 GHz
Memory Clock rate:	1566.00 Mhz
Memory Bus Width:	384-bit
L2 Cache Size:	786432 bytes

Things to note:

- Device 3 is a GPU
- “Tesla M2070” is the name of the model of the GPU.
- Device 3 contains multiple cores, or “sub-processors”. From the output, it has 448 CUDA-capable cores.

# nvidia-smi: CHECK GPU USAGE BEFORE chooseGpu()

```
[landau@impact1 ~]$ nvidia-smi
Thu Sep 13 09:37:05 2012
+-----+
| NVIDIA-SMI 2.290.10    Driver Version: 290.10      |
+-----+
| Nb. Name           | Bus Id     Disp. | Volatile ECC SB / DB |
| Fan   Temp          Power Usage /Cap | Memory Usage | GPU Util. Compute M. |
+=====+=====+=====+=====+=====+=====+=====+=====+
| 0. Tesla M2070        | 0000:0B:00.0 Off | 0% 9MB / 5375MB | 0% Default 0 0 |
| N/A   N/A P8  Off / Off | 0% 9MB / 5375MB | 0% Default 0 0 |
+-----+
| 1. Tesla M2070        | 0000:0C:00.0 Off | 0% 9MB / 5375MB | 0% Default 0 0 |
| N/A   N/A P8  Off / Off | 0% 9MB / 5375MB | 0% Default 0 0 |
+-----+
| 2. Tesla M2070        | 0000:0D:00.0 Off | 0% 9MB / 5375MB | 0% Default 0 0 |
| N/A   N/A P8  Off / Off | 0% 9MB / 5375MB | 0% Default 0 0 |
+-----+
| 3. Tesla M2070        | 0000:0E:00.0 Off | 0% 9MB / 5375MB | 0% Default 0 0 |
| N/A   N/A P8  Off / Off | 0% 9MB / 5375MB | 0% Default 0 0 |
+-----+
| Compute processes:                                     GPU Memory |
| GPU PID      Process name                           Usage   |
+-----+
| No running compute processes found                   |
+-----+
[landau@impact1 ~]$
```

```
[clandau@impact1 ~]$ nvidia-smi -i 0 -q
=====
NVSMI LOG=====

Timestamp : Thu Sep 13 09:37:54 2012
Driver Version : 290.10
Attached GPUs : 4

GPU 0000:0B:00.0
Product Name : Tesla M2070
Display Mode : Disabled
Persistence Mode : Disabled
Driver Model
    Current : N/A
    Pending : N/A
Serial Number : 0323111076435
GPU UUID : GPU-63911bd22733e078-94bd6965-7a0cbc1f-29f7d33f-489fcc8d5229600c5a45b88a
VBIOS Version : 70.00.3E.00.03
Inforom Version
    OEM Object : 1.0
    ECC Object : 1.0
    Power Management Object : 1.0
```

**PCI**

Bus	:	0x0B
Device	:	0x00
Domain	:	0x0000
Device Id	:	0x06D210DE
Bus Id	:	0000:0B:00.0
Sub System Id	:	0x083010DE
GPU Link Info		
PCIe Generation		
Max	:	2
Current	:	2
Link Width		
Max	:	16x
Current	:	16x
Fan Speed	:	N/A
Performance State	:	P8
Memory Usage		
Total	:	5375 MB
Used	:	9 MB
Free	:	5365 MB
Compute Mode	:	Default
Utilization		
Gpu	:	0 %
Memory	:	0 %

ECC Errors

Volatile

    Single Bit

        Device Memory : 0

        Register File : 0

        L1 Cache : 0

        L2 Cache : 0

        Total : 0

    Double Bit

        Device Memory : 0

        Register File : 0

        L1 Cache : 0

        L2 Cache : 0

        Total : 0

Aggregate

    Single Bit

        Device Memory : N/A

        Register File : N/A

        L1 Cache : N/A

        L2 Cache : N/A

        Total : 0

    Double Bit

Temperature	
Gpu	: N/A
Power Readings	
Power Management	: N/A
Power Draw	: N/A
Power Limit	: N/A
Clocks	
Graphics	: 270 MHz
SM	: 540 MHz
Memory	: 1566 MHz
Max Clocks	
Graphics	: 573 MHz
SM	: 1147 MHz
Memory	: 1566 MHz
Compute Processes	: None

[landau@impact1 ~]\$ |

## EXAMPLE: MATRIX MULTIPLICATION

Now, suppose I want to do a giant matrix multiplication on Device 3. I'm automatically set to Device 0:

```
> getGpuId()
[1] 0
```

So I change to Device 3:

```
> chooseGpu(3)
[[1]]
[1] 3
```

and then if I want, I can verify the change:

```
> getGpuId()
[1] 3
```

Now, I define the matrices that I want to multiply on Device 3:

```
> A <- matrix(runif(1e+7), nrow = 1e+4)
> B <- matrix(runif(1e+7), ncol = 1e+4)
```

Then, I tell the device to multiply A and B using the GPU hardware:

```
> ptm <- proc.time(); C <- gpuMatMult(A, B); proc.time() - ptm  
    user  system elapsed  
2.959   2.190   5.159
```

Compare the run time to that of the analogous CPU run on impact1:

```
> ptm <- proc.time(); D <- A %*% B; proc.time() - ptm  
    user  system elapsed  
116.389   0.166 116.503
```

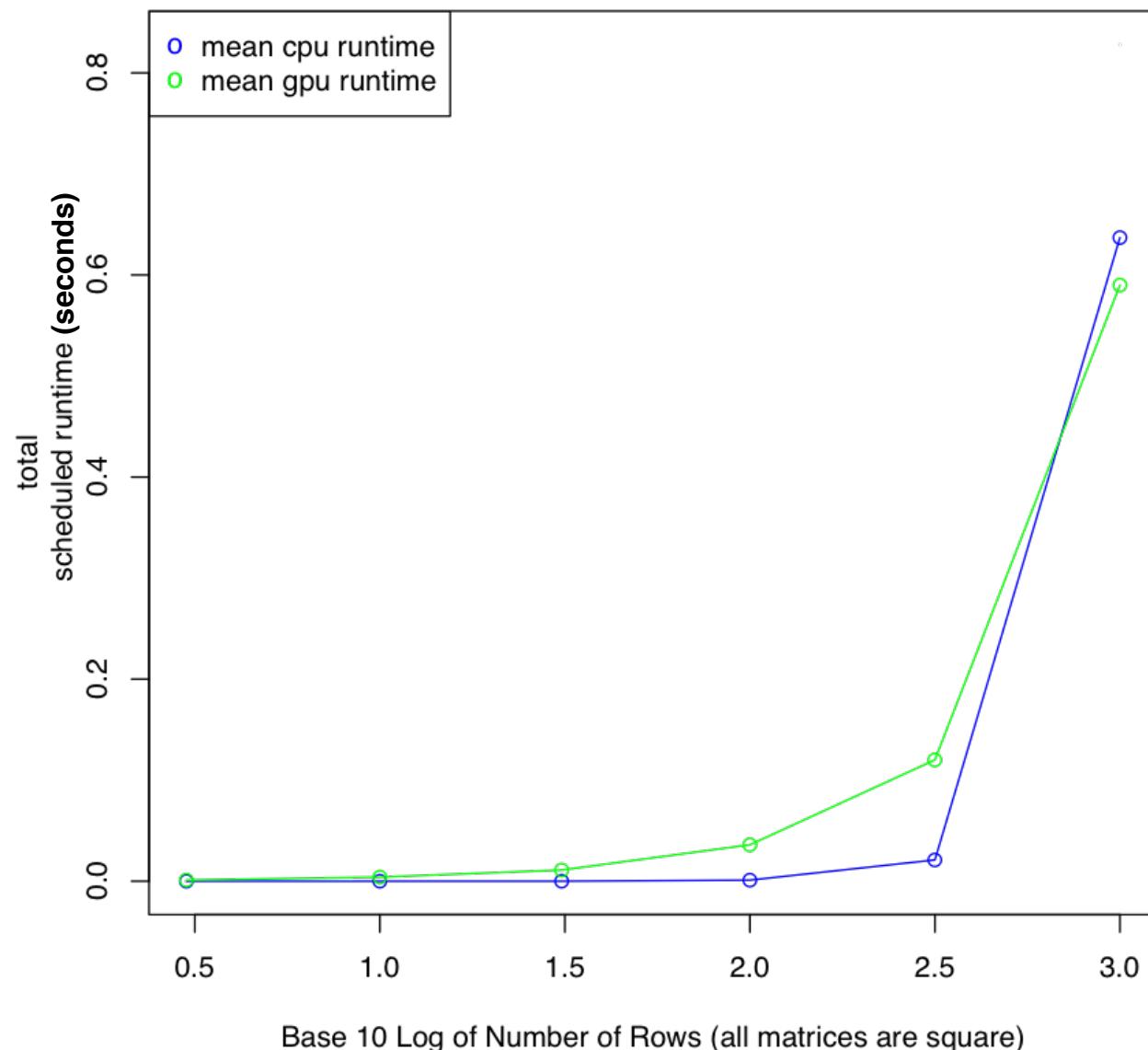
# **PERFORMANCE**

## A COMPARISON OF `gpuQr()` AND `qr()`

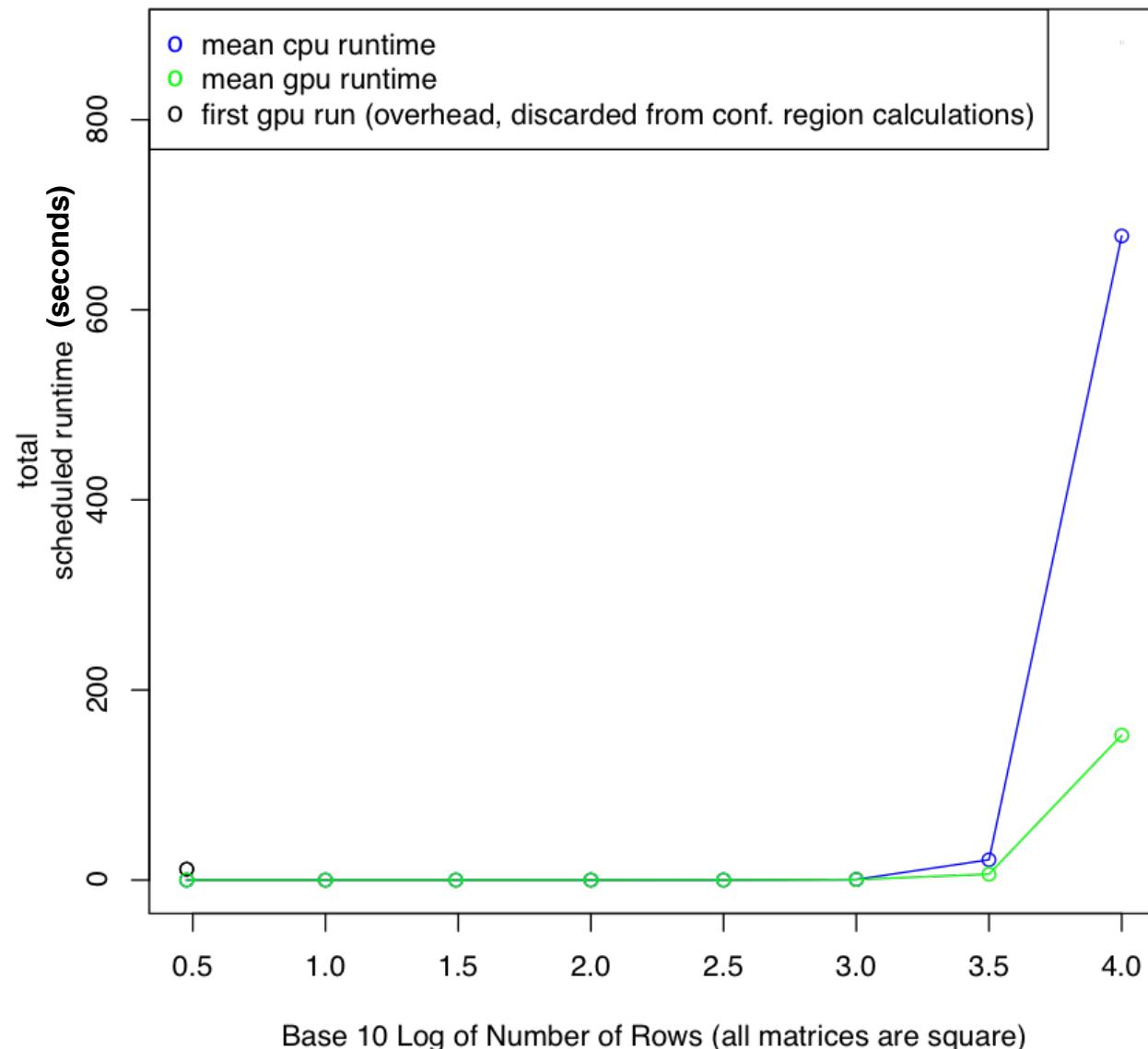
The R script, `gpuQr.r`, compares the performance of `gpuQr(arg)` and `qr(arg)` for square matrices `arg` of varying sizes.

See the results on the next few slides.

**total  
scheduled runtime:  
 $qr()$  vs  $gpuQr()$**



**total  
scheduled runtime:  
 $qr()$  vs  $gpuQr()$**

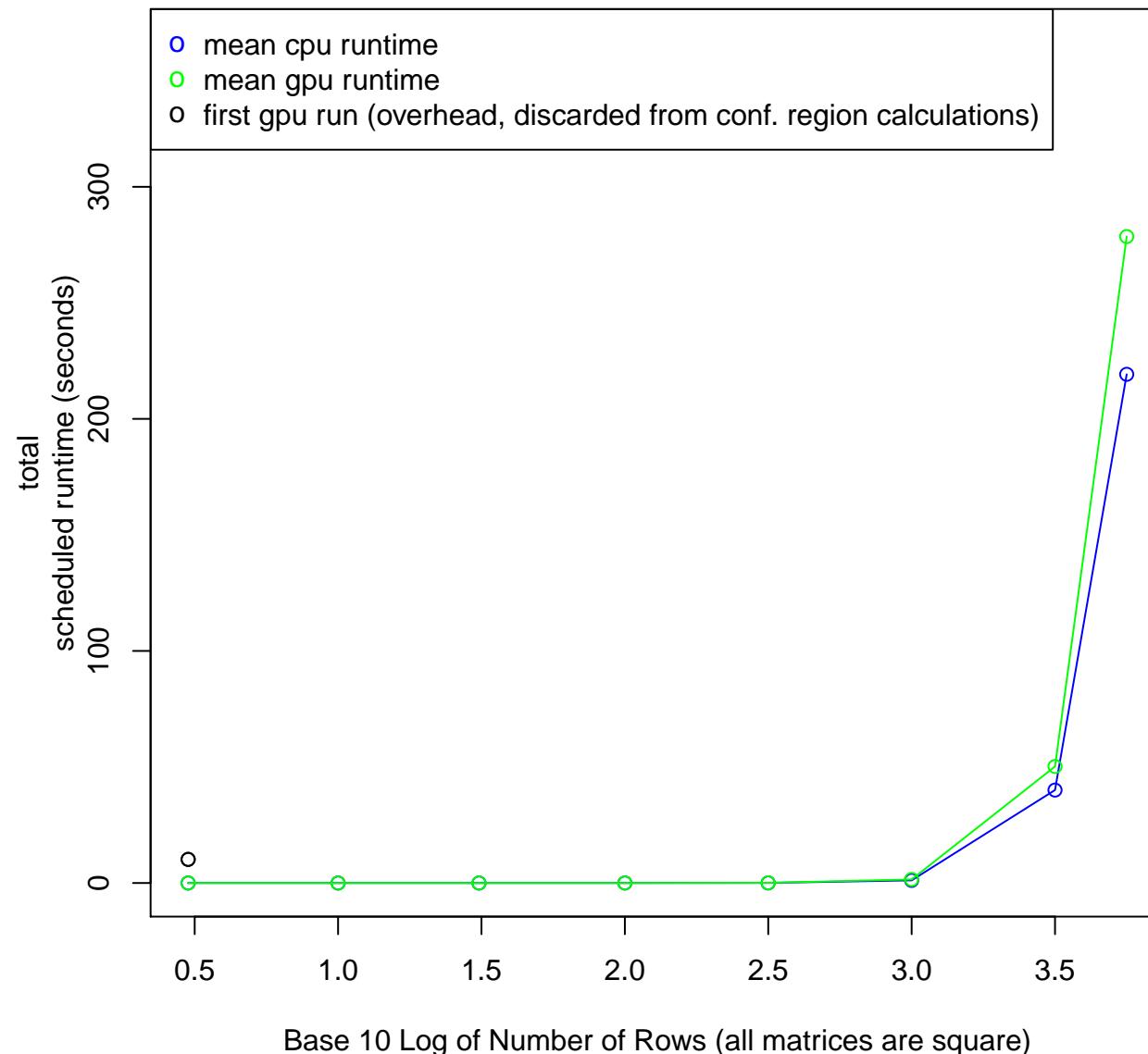


## A COMPARISON OF `gpuSolve()` AND `solve()`

The R script, `gpuSolve.r`, compares the performance of `gpuSolve(arg)` and `solve(arg)` for square matrices `arg` of varying sizes.

See the results on the next slide.

**total  
scheduled runtime (seconds):  
solve() vs gpuSolve()**



## A COMPARISON OF `gpuLm()` AND `lm()`

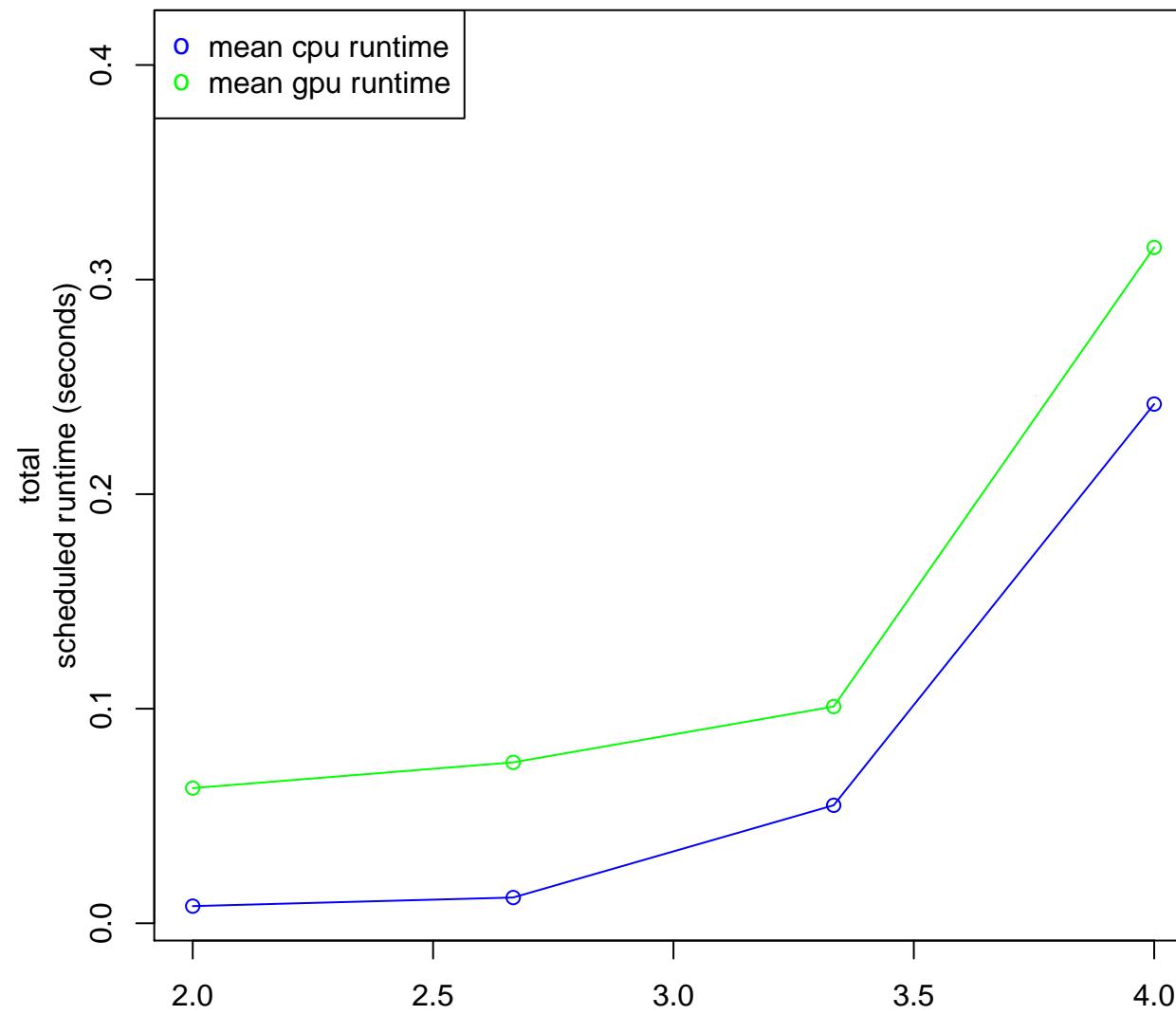
The R script, `gpuLm.r`, compares the performance of `gpuLm(y ~ X)` and `lm(y ~ X)`, where:

- `y` is a random vector of observations.
- `X` is a random design matrix with `length(y)` rows and 100 columns.

The script times each function with varying `length(y)` and `nrow(X)`.

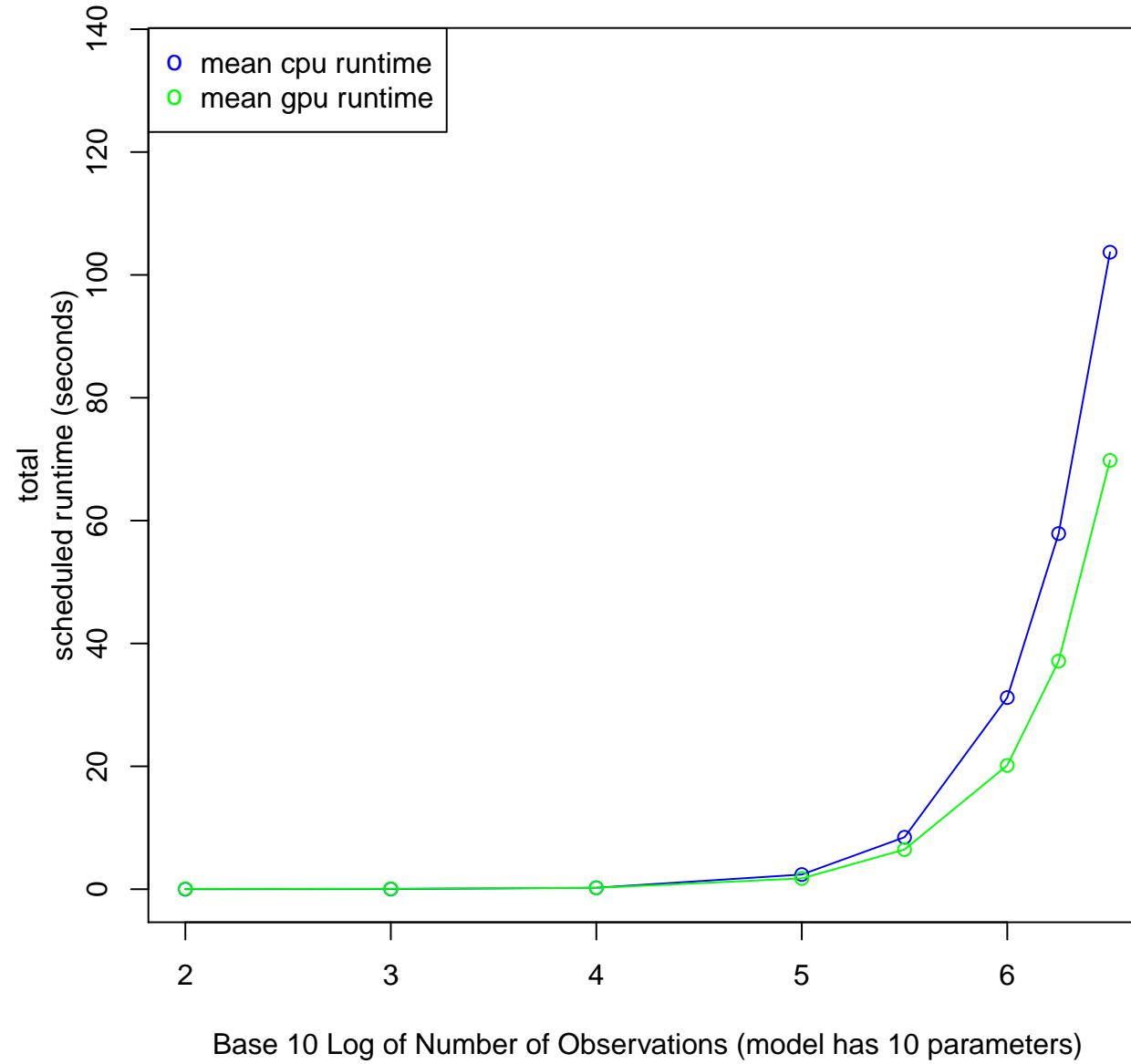
See the results on the next slides.

**total  
scheduled runtime (seconds):  
lm() vs gpuLm()**



Base 10 Log of Number of Observations (model has 10 parameters)

**total**  
**scheduled runtime (seconds):**  
**lm() vs gpuLm()**



# A COMPARISON OF `gpuGlm()` AND `glm()`

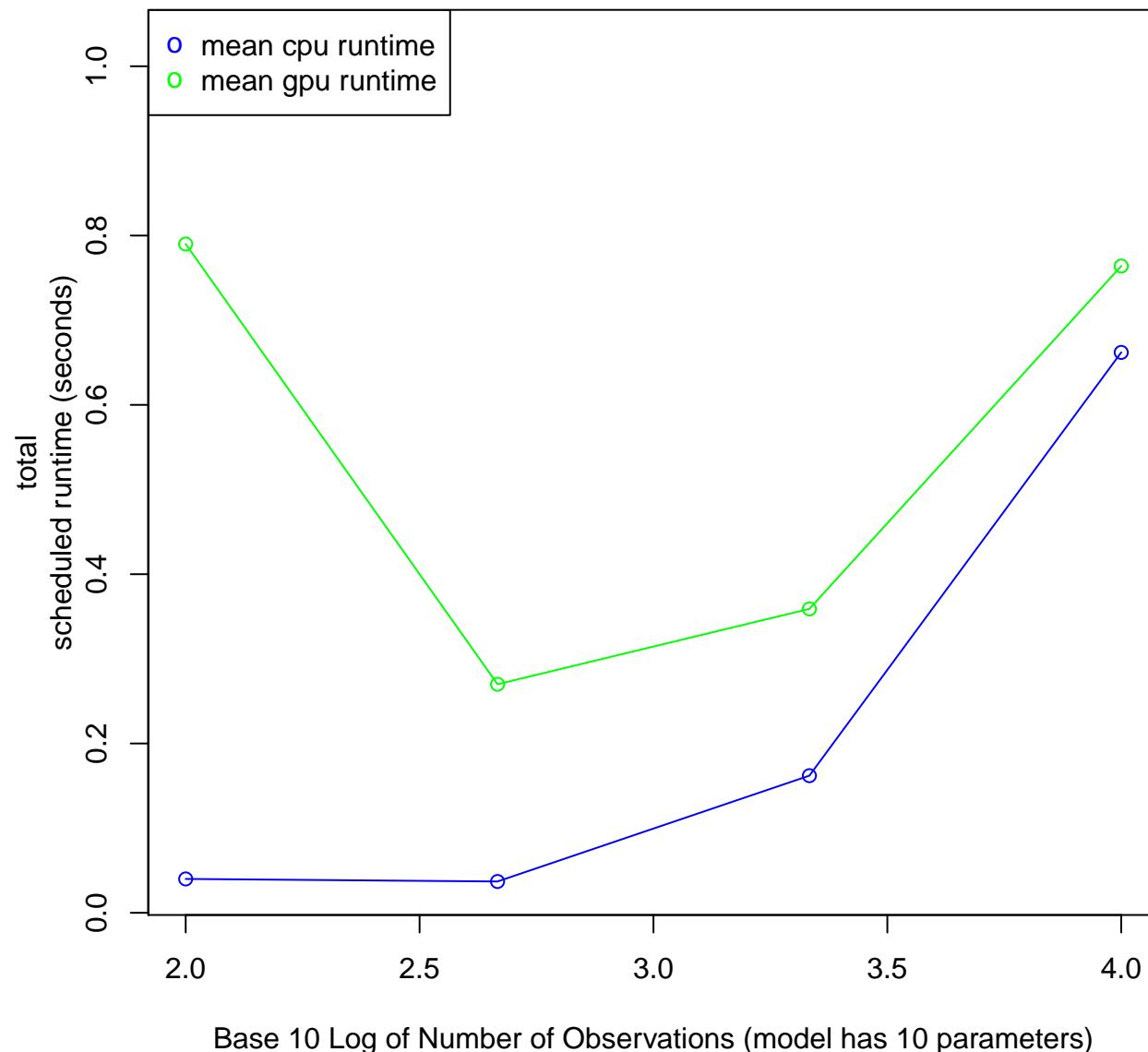
The R script, `gpuGlm.r`, compares the performance of `gpuGlm(y ~ X, family = poisson())` and `glm(y ~ X, family = poisson())`, where:

- `y` is a random vector of observations.
- `X` is a random design matrix with `length(y)` rows and 100 columns.

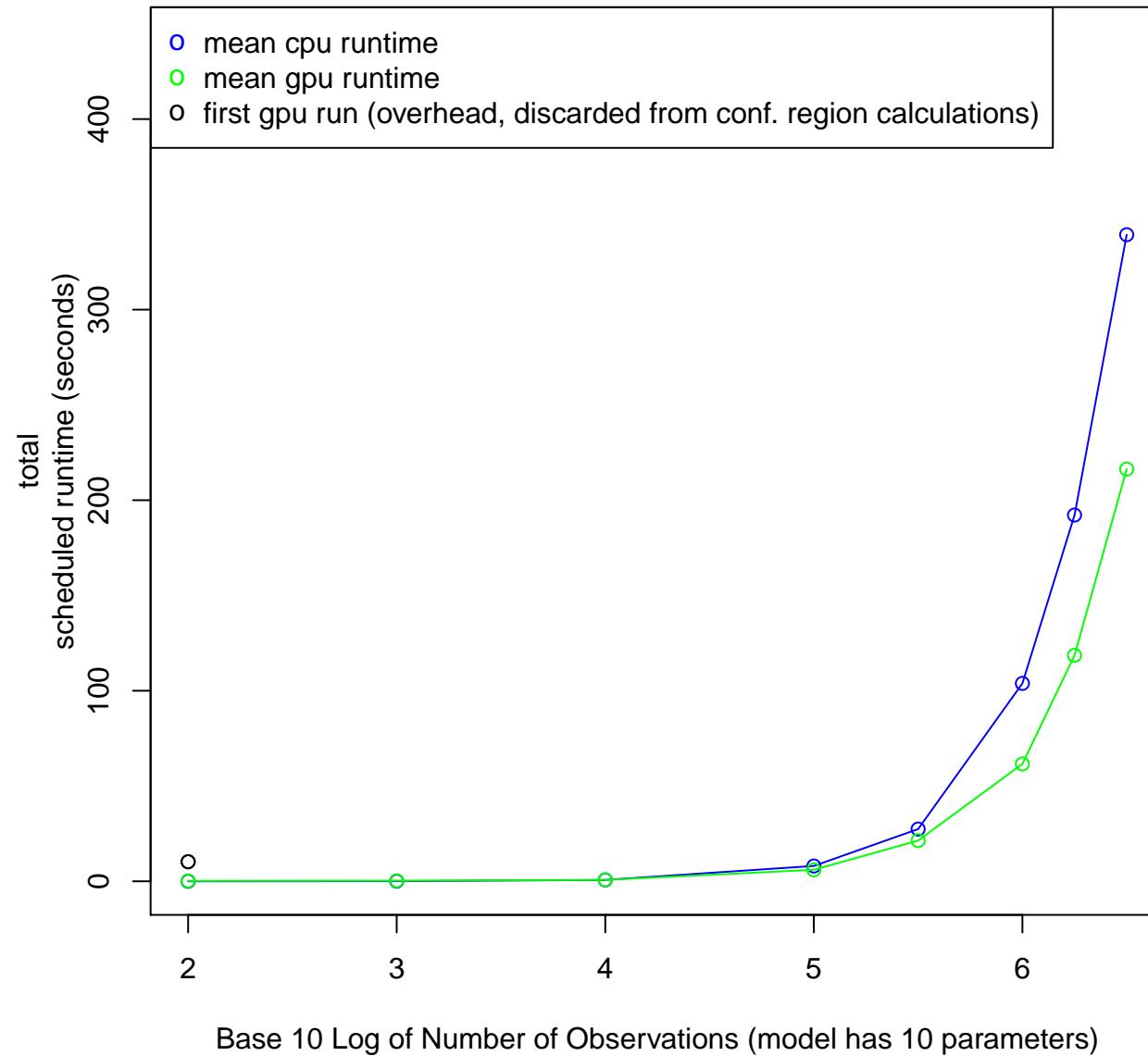
The script times each function with varying `length(y)` and `nrow(X)`.

See the results on the next slides.

**total  
scheduled runtime (seconds):  
glm() vs gpuGlm()**



**total**  
**scheduled runtime (seconds):**  
**glm() vs gpuGlm()**

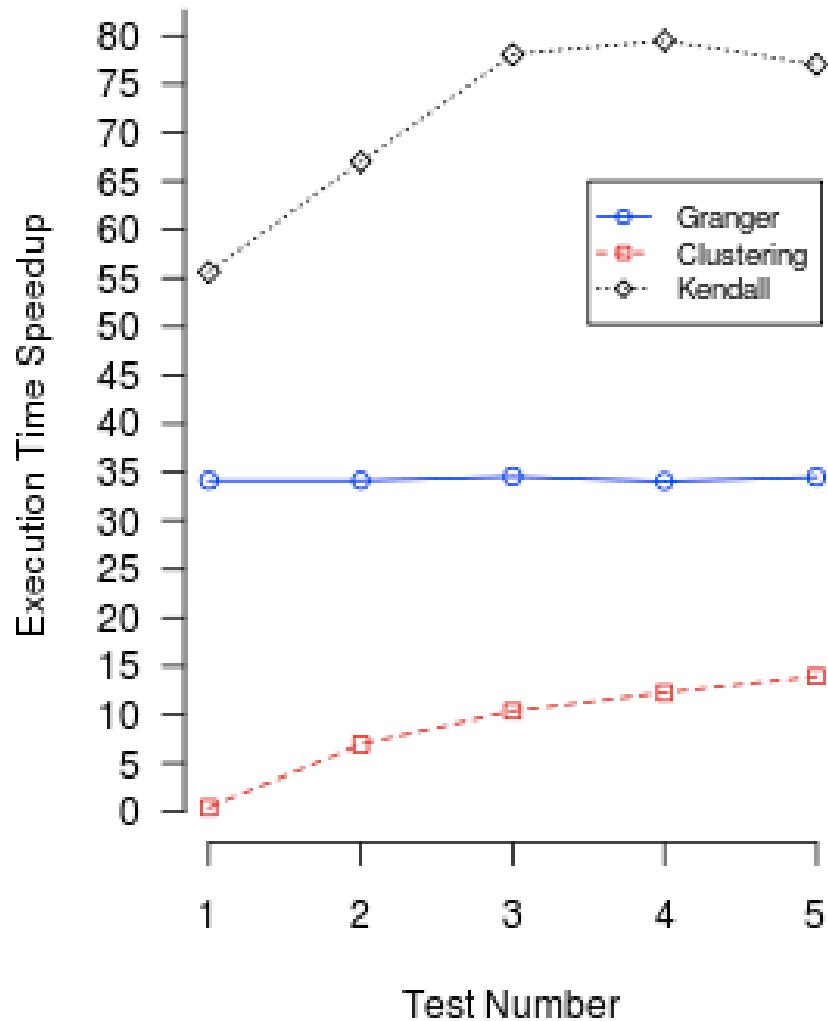


# CLAIMS FROM THE AUTHORS OF gputools

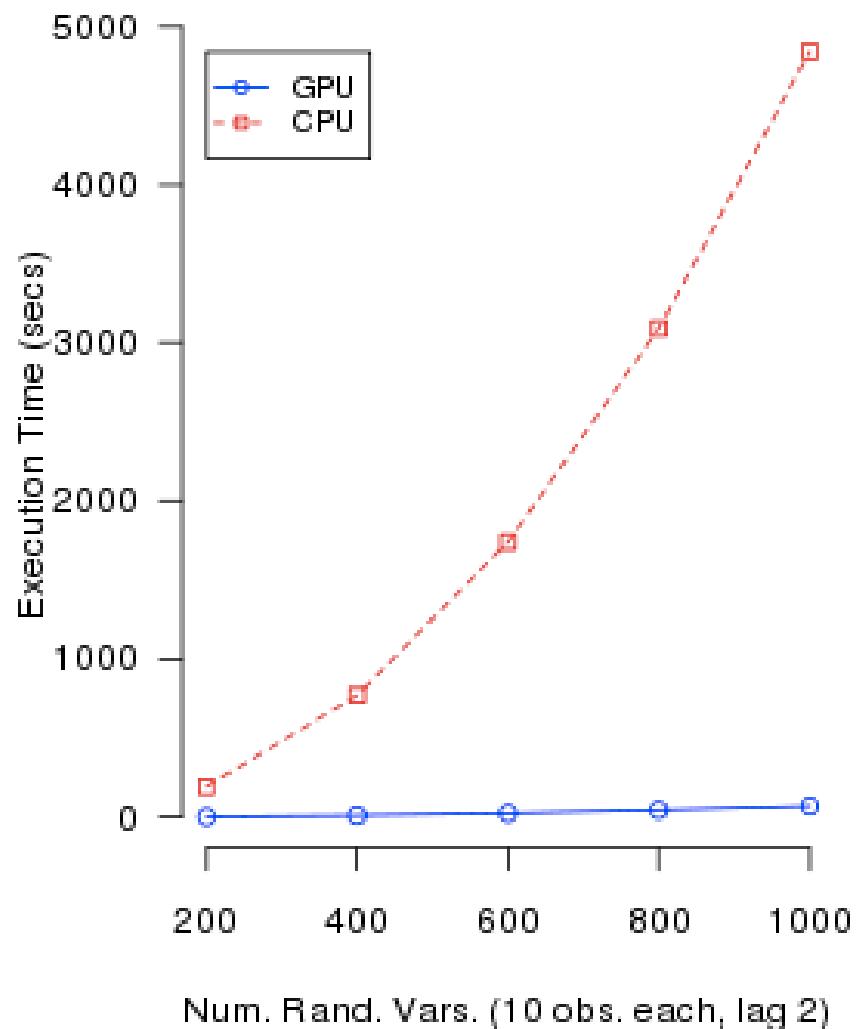
(All of the following is from  
<http://brainarray.mbnl.med.umich.edu/Brainarray/Rgpgpu/.>)

“Tested on a subset of GSE6306, non-GPU enabled fastICA took over four hours while gpuFastICA took just 80 seconds!”

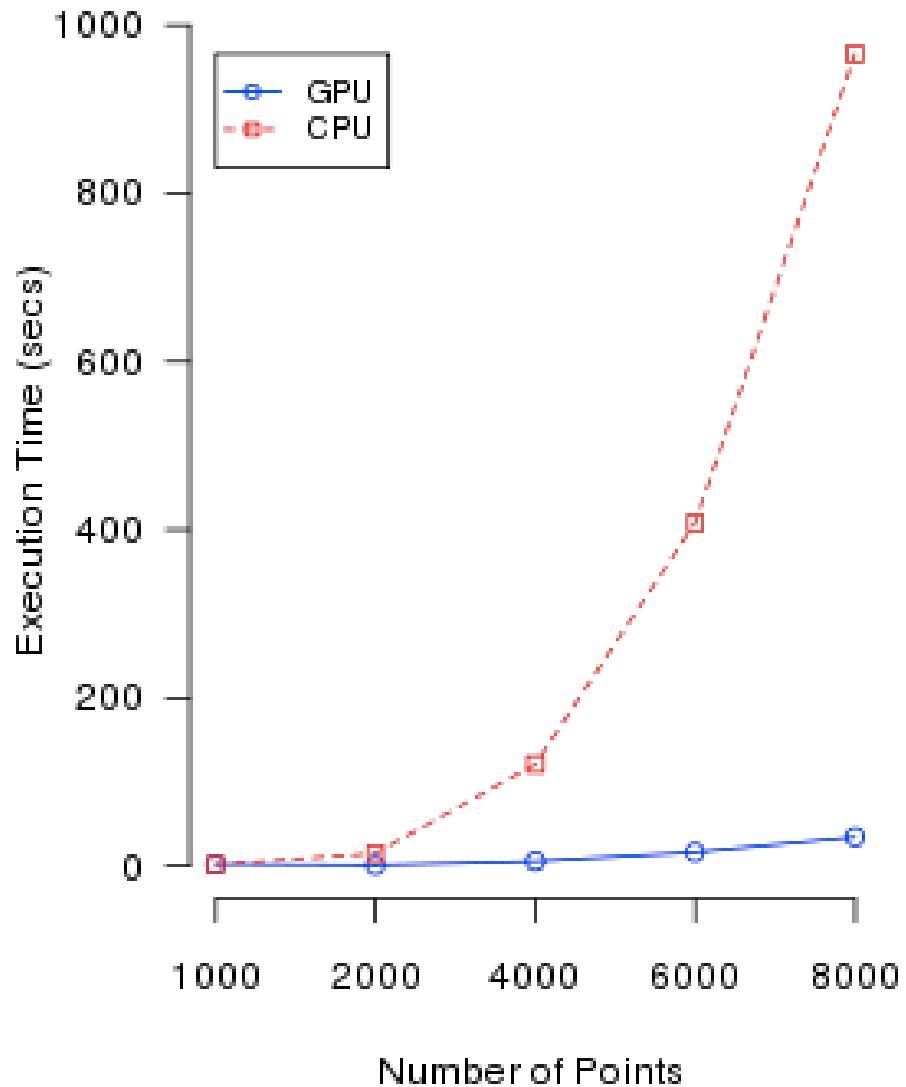
*Fig. 1: Speedup (R GPU vs. CPU)*



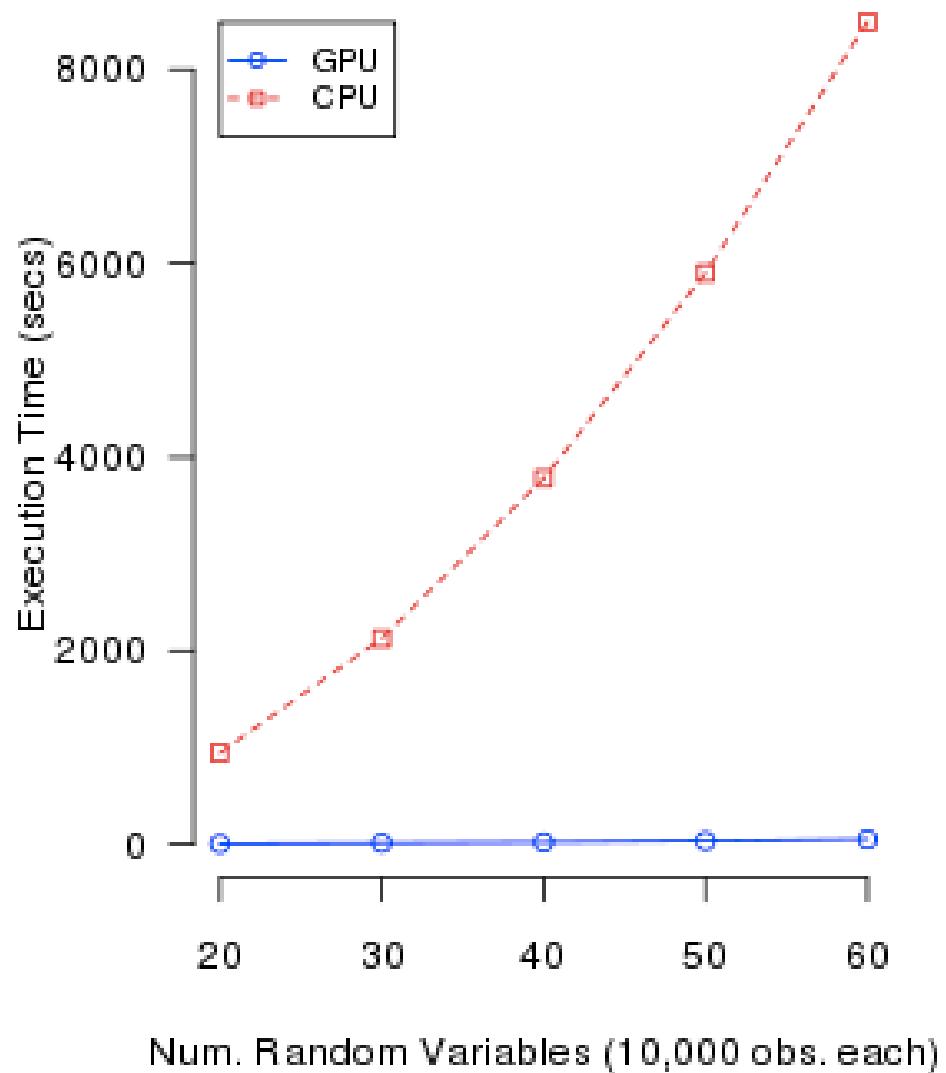
**Fig. 2: Granger Times**



*Fig. 3: Cluster Times*



*Fig. 4: Kendall Times*



## OTHER R PACKAGES FOR THE GPU

- WideLM - used to quickly fit a large number of linear models to a fixed design matrix and response vector.
- magma - a small linear algebra with implementations of backsolving and the LU factorization.
- cudaBayesreg - implements a Bayesian model for fitting fMRI data.
- gcbd - a Debian package for “benchmarking” linear algebra algorithms such as the QR, SVD and LU factorizations.

# Outline

- Contents of gptools
- Usage
- Performance
- Other R packages for the GPU

# GPU SERIES MATERIALS

These slides, a tentative syllabus for the whole series, and code are available at:

<https://github.com/wlandau/gpu>.

After logging into you home directory on impact1, type:

```
git clone https://github.com/wlandau/gpu
```

into the command line to download all the materials.

# REFERENCES

- Josh Buckner, Mark Seligman, Justin Wilson. “R+GPU”.  
<http://brainarray.mbnl.med.umich.edu/Brainarray/Rgpgpu/#introduction>.
- Carten O. Daub, Ralf Steuer, Joachim Selbig, and Sebastian Kloska. 2004. Estimating mutual information using B-spline functions - an improved similarity measure for analysing gene expression data. BMC Bioinformatics. 5:118. Available from  
<http://www.biomedcentral.com/1471-2105/5/118>
- Carpenter, Austin. cuSVM: a cuda implementation of support vector classification and regression. <http://patternsonascreen.net/cuSVM.html>
- Dirk Eddelbuettel. “Package gcbd”.  
<http://cran.r-project.org/web/packages/gcbd/gcbd.pdf>.
- Hacker R.S. and Hatemi-J A. (2006) ”Tests for causality between integrated variables using asymptotic and bootstrap distributions: theory and application”, Applied Economics, Vol. 38(13), pp. 1489-1500.
- Hand, David J. and Till, Robert J. (2001). A simple generalisation of the area under the ROC curve for multiple class classification problems. Machine Learning. 45, 171-186.

A. Hyvarinen and E. Oja (2000) Independent Component Analysis: Algorithms and Applications, Neural Networks, 13(4-5):411-430. <http://www.cis.hut.fi/aapo/>

A. Hyvarinen. Independent Component Analysis: Recent Advances. Philosophical Transactions of the Royal Society A, in press.  
<http://www.cs.helsinki.fi/u/ahyvarin/papers/PTRSA12.pdf>.

Mark Seligman, Chris Fraley. “Package WideLM”.  
<http://cran.r-project.org/web/packages/WideLM/WideLM.pdf>.

Brian J Smith. “Package cudaBayesreg”.  
<http://cran.r-project.org/web/packages/cudaBayesreg/cudaBayesreg.pdf>.