

# **Parallel-PSM**

# **Parallelizing the Population**

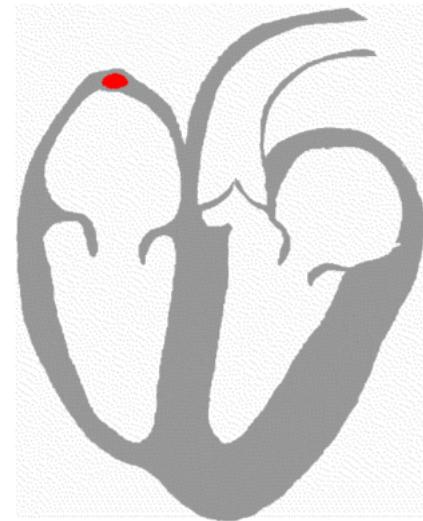
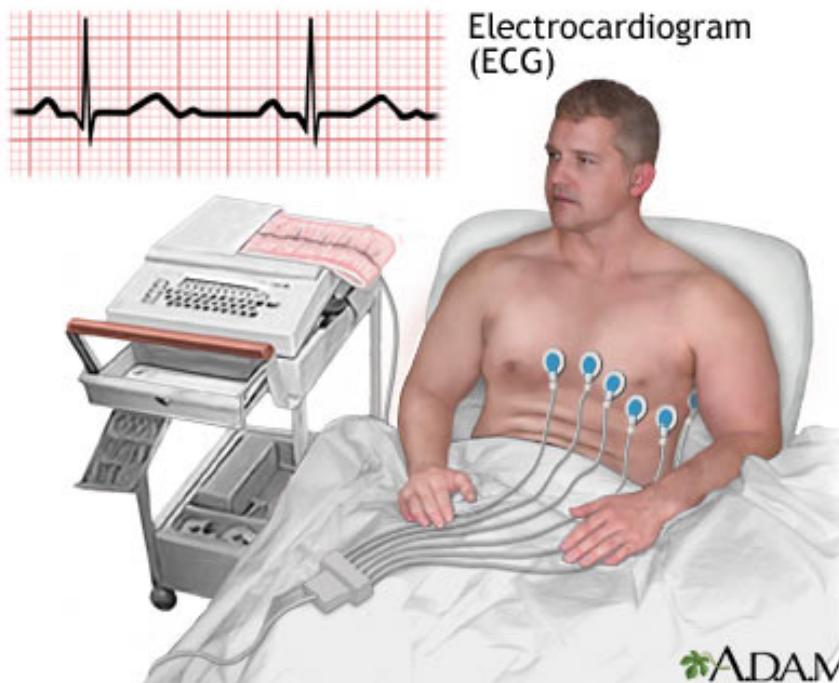
# **Stochastic Modeling R Package**

# **with UNICORE**

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# ECG - Electrocardiogram



# CPR - cardiopulmonary resuscitation





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# Start of the story...

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# Modelling the relationship between ECG characteristics and CPR quality during cardiac arrest

Kenneth Gundersen

Jan Terje Kvaløy

# Introduction

- Measurements easily available during out-of-hospital resuscitation include:
  - Electrocardiogram (ECG)
  - End-Tidal CO<sub>2</sub> (ETCO<sub>2</sub>)
  - **Compression depth** and **force**
  - Thoracic impedance
- Overall aim: a mathematical model for the interaction between intervention and response during VF/VT.
- This work: develop minimal model for CPR quality variables and ECG coarseness (response).

# Introduction

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- This type of model can possibly be used to:
  - Identify the CPR quality variables that indicate its effectiveness.
  - Test different compression techniques against each other.
  - Predict the patients' response to further CPR and thereby to **choose between immediate defibrillation or further CPR.**
  - General framework for testing different hypothesis.

# ECG coarseness – median slope

- The ROSC-predictor median-slope (MS) has high prediction accuracy and is easily computed:

$$MS = \text{median}(|\text{ecg}(n) - \text{ecg}(n-1)|), n = 1, \dots, N$$

- MS combines amplitude and frequency information -> high ECG amplitude and frequency (coarse ECG) indicates a good state of patient.

# ECG coarseness – median slope

Normal



Heart block



# Data

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- Observational data from out-of hospital cardiac arrest episodes.
- 85 patients in baseline data, 27 patients contributed with a total of 118 VF/VT sequences.

# Challenges

- Model fitting extremely computationally intensive.
- Heterogeneous patients -> random effects on many parameters -> many model parameters to fit.
- Limited size of available dataset.
- Noise in ECG recordings (must be manually identified and censored).

# Future research

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- **Speed up model fitting (modify software to enable parallel processing).**
- **Obtain (larger) dataset where ECG noise is minimized.**
- Include ETCO<sub>2</sub> in an extended model.
- Consider excluding "hopeless" patients from analysis.
- Take a step back and develop (idealized) model from animal data?

# In brief...



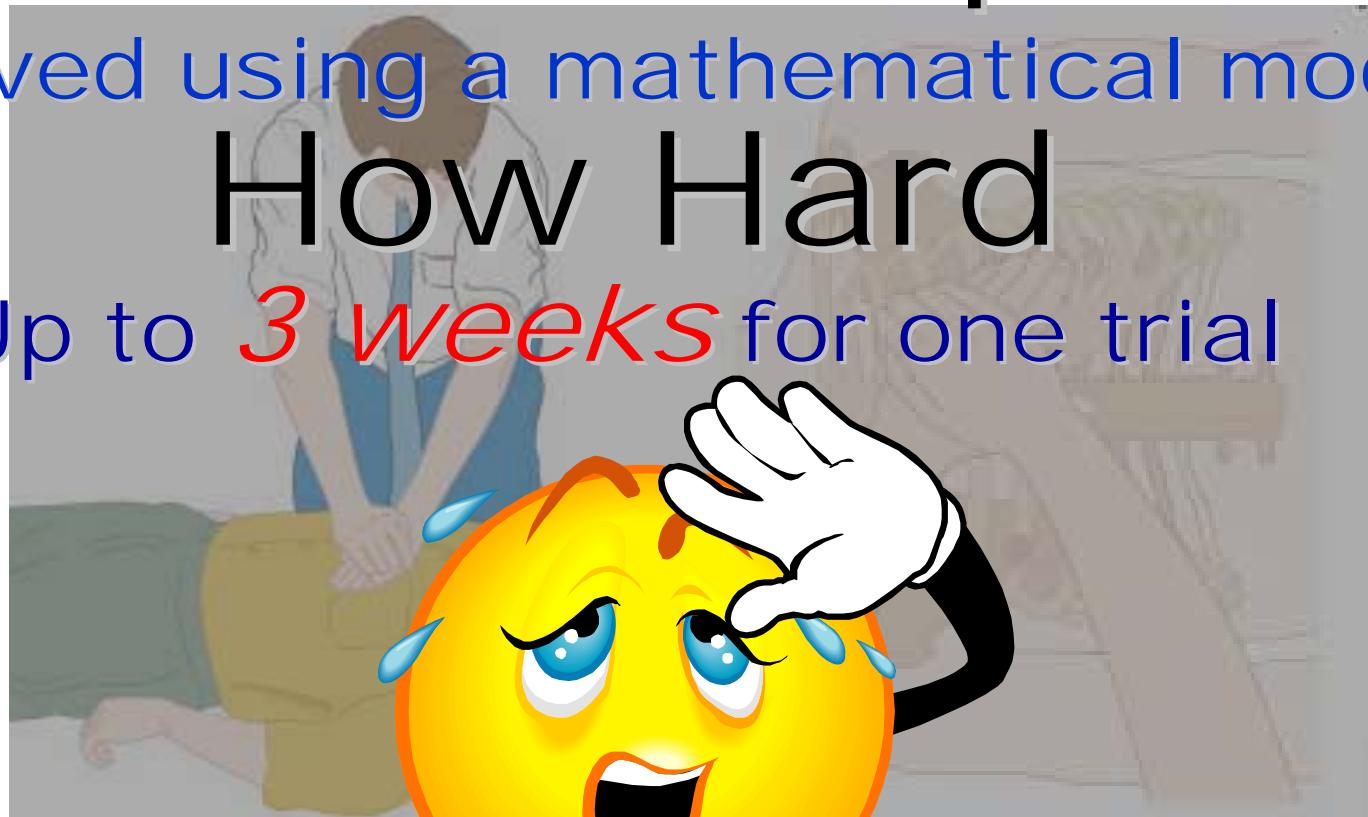
In brief...

# How Deep

Solved using a mathematical model

# How Hard

Up to *3 weeks* for one trial

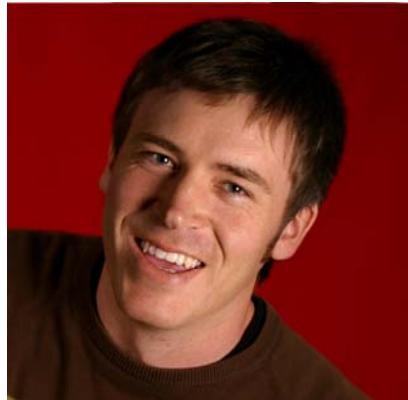




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And then...

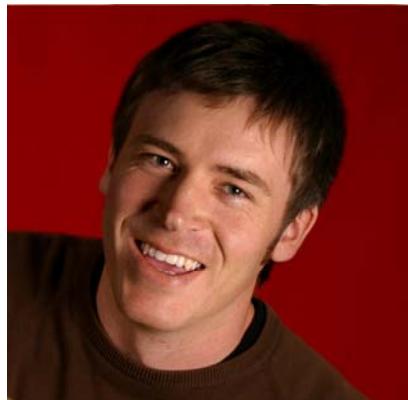
# Scenario...



Hein,  
can you try to do it?



# Scenario...



Hein,  
can you try to do it?

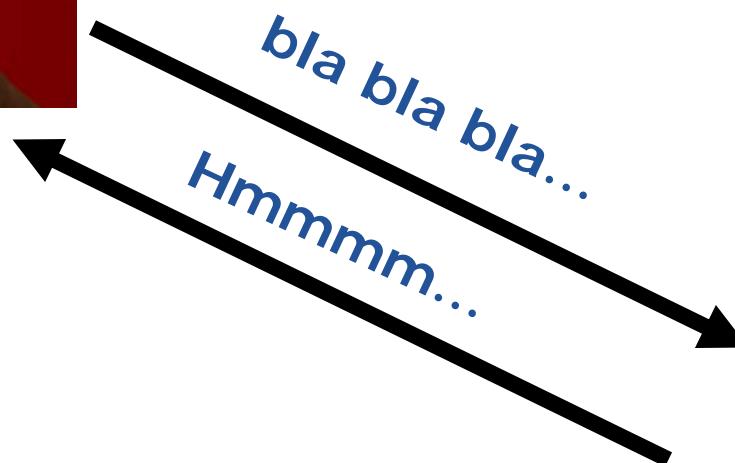
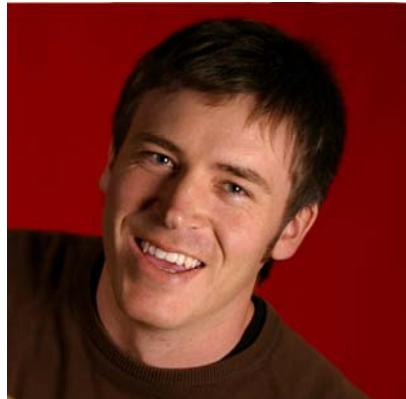


CC:



Azab,  
See what you can do?

# Scenario...



# Team

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Kenneth Gundersen  
Post Doc  
UiS



Hein Meling  
Asc. Professor  
UiS

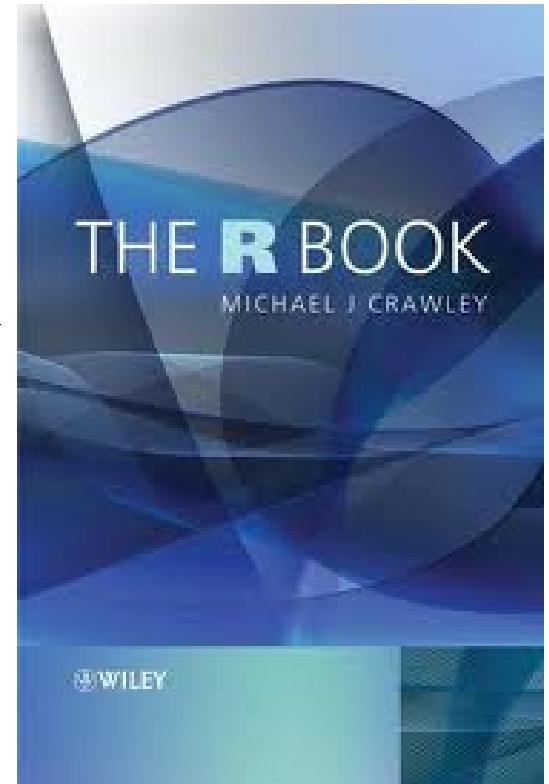


Abdulrahman Azab  
Research Fellow  
UiS

# Scenario...



Learning R



# The *R* language

## Introduction to R

### What is R?

- ▶ A ‘language and environment for statistical computing and graphics’ (implements a dialect of the language ‘S’)
- ▶ Syntax is C-like, but philosophy is functional
- ▶ Focus on matrices and vectors
- ▶ Free, open-source (GPL)
- ▶ Well documented
- ▶ Command line based, but there are GUIs
- ▶ Latest version: 2.13.0 (April 2011)
- ▶ URL: <http://www.r-project.org/>

# The *R* language

## Introduction to R

### R Features

- ▶ Very many analysis methods available
- ▶ Scriptable and extensible
- ▶ Can be used interactively, process batch jobs or run as a script
- ▶ Bindings to many other systems/languages, e.g., Python, Perl, Matlab, \*SQL, Excel
- ▶ Active user community with thousands of contributed packages

# The *R* language

## Introduction to R

### Elementary Data Types

Basic elements: *numbers, strings, logicals*

```
> 42
[1] 42
> "a string"
[1] "a string"
> TRUE
[1] TRUE

> 42 + 13.5
[1] 55.5
> 42 > 13.5
[1] TRUE
> substr("a string", 3, 5)
[1] "str"
> ! TRUE
[1] FALSE
```



# The *R* language

## Introduction to R

### Compound Data Types: Vectors

- ▶ Basic elements collected in *vectors*, *lists*, *matrices* and *data frames*
- ▶ Collection of *equal* types of elements: *vector*:

```
> 1:5
[1] 1 2 3 4 5
> c(42, 1:5)  # "c" for "concatenate"
[1] 42  1  2  3  4  5
> c("three", "small", "things")
[1] "three"  "small"  "things"
```

- ▶ Indexing:

```
> nums <- c(42, 33, 58, 1, 3.2)
> nums[2]
[1] 33
> nums[2:3]
[1] 33 58
> nums[c(1,3)]
[1] 42 58
```



# The *R* language

## Introduction to R

### Compound Data Types: Lists

- ▶ Collection of *different* types of elements: *list*:

```
> c(42, "Mary")      # Probably not what you want
[1] "42"    "Mary"
> list(42, "Mary")   # That's more like it!
[[1]]
[1] 42
```

```
[[2]]
[1] "Mary"
```

- ▶ Indexing: `[]` and `[][]`:

```
> lag <- list(c(3,5), "string", rep(TRUE, 5))
> lag
> lag[2:3]    # sub list
> lag[3]       # still sub list
> lag[[3]]    # element
```

# The *R* language

## Introduction to R

### Compound Data Types: Matrices

- ▶ Two-dimensional collections: *matrices* and *data frames*

- ▶ Same element type (usually number): *matrix*:

```
> A <- matrix(1:9, ncol = 3, nrow = 3)
```

```
> A
```

```
 [,1] [,2] [,3]
```

```
[1,] 1 4 7
```

```
[2,] 2 5 8
```

```
[3,] 3 6 9
```

```
> B <- matrix(6:1, ncol = 2, nrow = 3)
```

```
> A %*% B
```

```
 [,1] [,2]
```

```
[1,] 54 18
```

```
[2,] 69 24
```

```
[3,] 84 30
```

- ▶ Indexing: `A[2,1]`

# The *R* language

## Introduction to R

### Variables

- ▶ Values can be stored in *variables*:

```
> mynum <- 42
> mynum + 13.5
[1] 55.5
> adj <- "interesting"
> sentence <- paste("Very", adj)
> sentence
[1] "Very interesting"
```

- ▶ Tip: use descriptive names; avoid single-character names.
- ▶ List all variables: `ls()`
- ▶ Show value of variable: `mynum` or `print(mynum)`
- ▶ Remove a variable: `rm()`, e.g., `rm(adj)`

# The *R* language

## Introduction to R

### Help!

This is probably the most important slide!

- ▶ `?mean` - help for a function
- ▶ `help.search("regression")` or simply `??regression` - search in your installed R
- ▶ `RSiteSearch("logistic")` - search the R web site
- ▶ `demo()` - list/run demos
- ▶ `vignette()` - list package vignettes
- ▶ `help.start()` - start help centre



# The *R* language

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[http://www.notur.no/notur2011/material/R\\_course\\_notur2011\\_slides.pdf](http://www.notur.no/notur2011/material/R_course_notur2011_slides.pdf)



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# Back to the problem...

# ECG – CPR modeling code (by Kenneth)

## 1. Create the model

```
NLmod.MS <- Model
```

## 2. Generate the simulated heart block reading

```
simdata <- PSM.simulate(NLmod.MS,...)
```

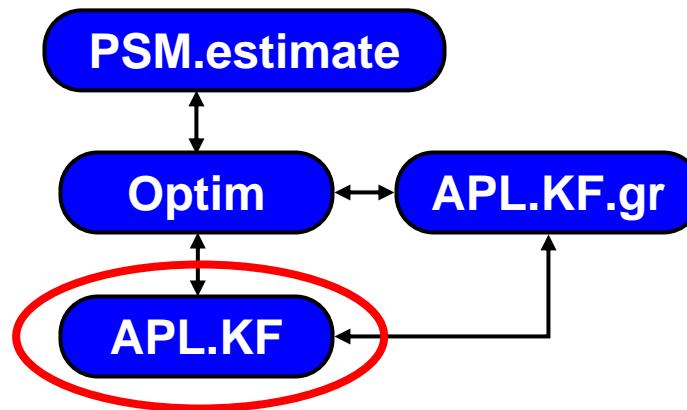


## 3. Estimate the optimum compression force and depth

```
fit_simmod <-
PSM.estimate(NLmod.MS,simdata,...,maxit=1000,...)
```

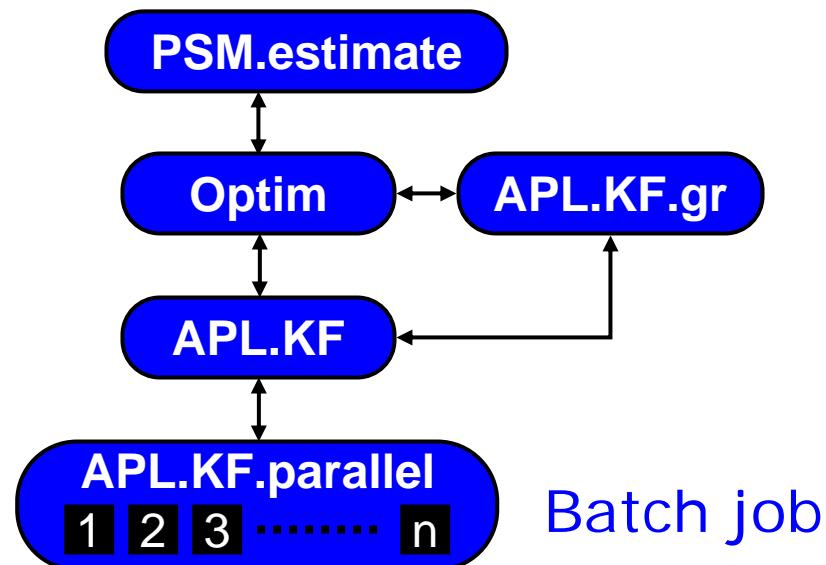
30 minutes per iteration on a single machine 31

# ECG – CPR modeling code (by Kenneth)



Very computational intensive, but parallelizable

# ECG – CPR modeling code (by Kenneth)

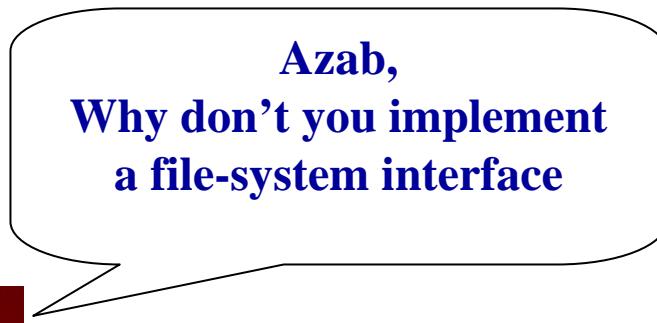


# Challenges

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- Parallel packages for R, e.g. snow and GridR, are either not dependable or faulty.
- R doesn't directly support java. Cannot use HiLA.
- The computation is iterative, cannot be shaped as a workflow.
- It is time consuming to create a new parallel R package, *and I want to finish my PhD.*

# Thinking...



# Thinking...



Azab,  
Why don't you implement  
a file-system interface

Let's try and see..

Do you think it is a good  
solution?



## Stroll: Job submission by file-system `write()`

---

```
#Create the job directory
taskDir<-dir.create("./Parallel.APL.KF")

# Set the job configuration
write("exec=job.R;in=in$(i).RData;target=unicore", file
      = "./config")
```

...Generate input files...

```
# Submit the job
write("SUBMIT -W", file = "./control")
```

...Collect output files...



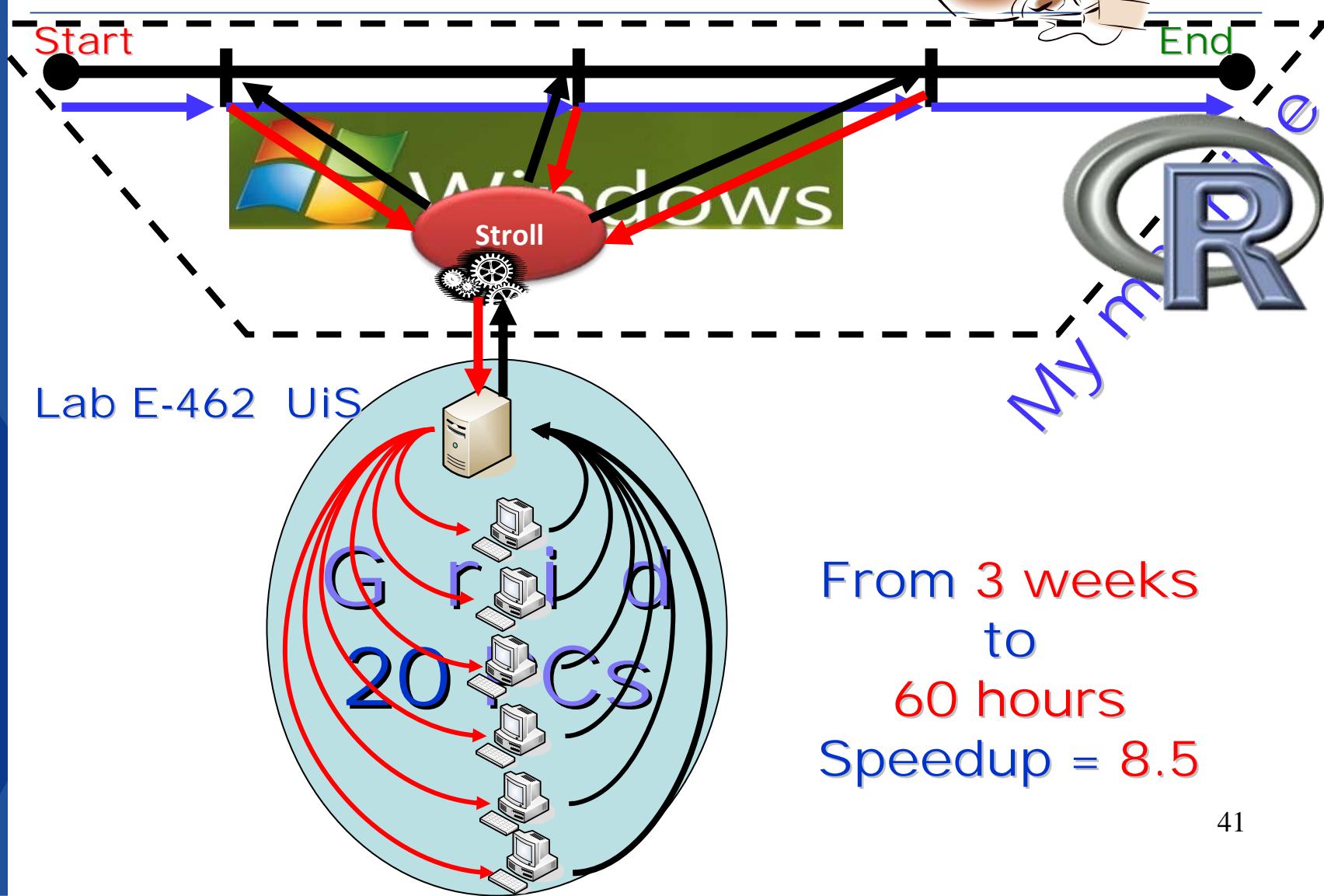
# Stroll: Job submission by file-system `write("SUBMIT")` using Condor

```
> source("simmod_Grid.R")
* Non-linear model (Grid)*
Using logit transformation of parameters
Grid: Starting switch statement
Using Optimizer:      optim
optim: Start
optim: functions defined
optim: 1
APL.KF: Start dimS = 40 , Time: 0.19
Grid: input copied to 40 Workers: elapsed time = 0.91
1 ,2 ,3 ,4 ,5 ,6 ,7 ,8 ,9 ,10 ,11 ,12 ,13 ,14 ,15 ,16 ,17 ,18 ,19 ,20 ,21 ,22 ,23 ,24 ,25 ,26 ,27 ,28 ,29 ,30
,31 ,32 ,33 ,34 ,35 ,36 ,37 ,38 ,39 ,40 ,
Submitting job(s).....
Logging submit event(s).....
40 job(s) submitted to cluster 624.
Waiting for results from workers....
3 ,9 ,5 ,7 ,6 ,11 ,12 ,4 ,8 ,13 ,17 ,10 ,14 ,15 ,2 ,16 ,18 ,19 ,20 ,21 ,22 ,23 ,24 ,25 ,1 ,26 ,27 ,28 ,29 ,30
,31 ,32 ,33 ,34 ,36 ,37 ,38 ,39 ,40 ,35 ,
APL.KF: Ends , Time: 150.5
      a
      -logL = -45.2597115 (2:30.31)
optim: 2
Grid: switch statement ended
[1] Runtime: 2:30.56
Collecting average data colums..
```

# Stroll: Job submission by file-system **write("SUBMIT -W")** using UNICORE

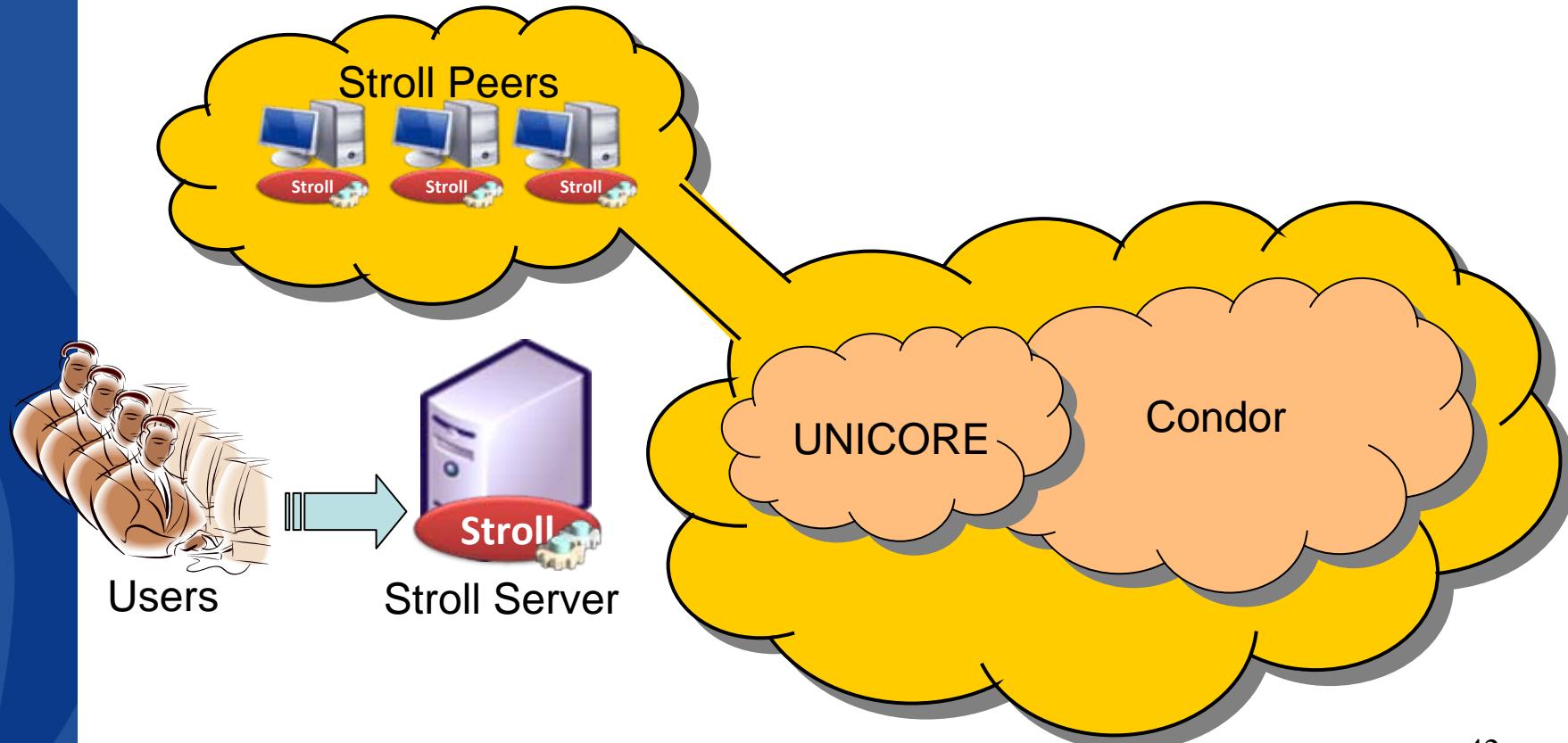
```
> source("simmod_Grid.R")
* Non-linear model (Grid)*
Using logit transformation of parameters
Grid: Starting switch statement
Using Optimizer:      optim
optim: Start
optim: functions defined
optim: 1
APL.KF: Start dimS = 40 , Time: 0.07
Grid: input copied to 40 Workers: elapsed time = 4.21
1 ,2 ,3 ,4 ,5 ,6 ,7 ,8 ,9 ,10 ,11 ,12 ,13 ,14 ,15 ,16 ,17 ,18 ,19 ,20 ,21 ,22 ,23 ,24 ,25 ,26 ,27 ,28 ,29 ,30
,31 ,32 ,33 ,34 ,35 ,36 ,37 ,38 ,39 ,40 ,
Waiting for results from workers....
1 ,2 ,3 ,4 ,5 ,6 ,7 ,8 ,9 ,10 ,11 ,12 ,13 ,14 ,15 ,16 ,17 ,18 ,19 ,20 ,21 ,22 ,23 ,24 ,25 ,26 ,27 ,28 ,29 ,30
,31 ,32 ,33 ,34 ,35 ,36 ,37 ,38 ,39 ,40 ,
APL.KF: Ends , Time: 294.01
      a
-logL = -45.2597115 (4:53.97)
optim: 2
Grid: switch statement ended
[1] Runtime: 4:54.04
Collecting average data colums..
```

## Execution Scenario

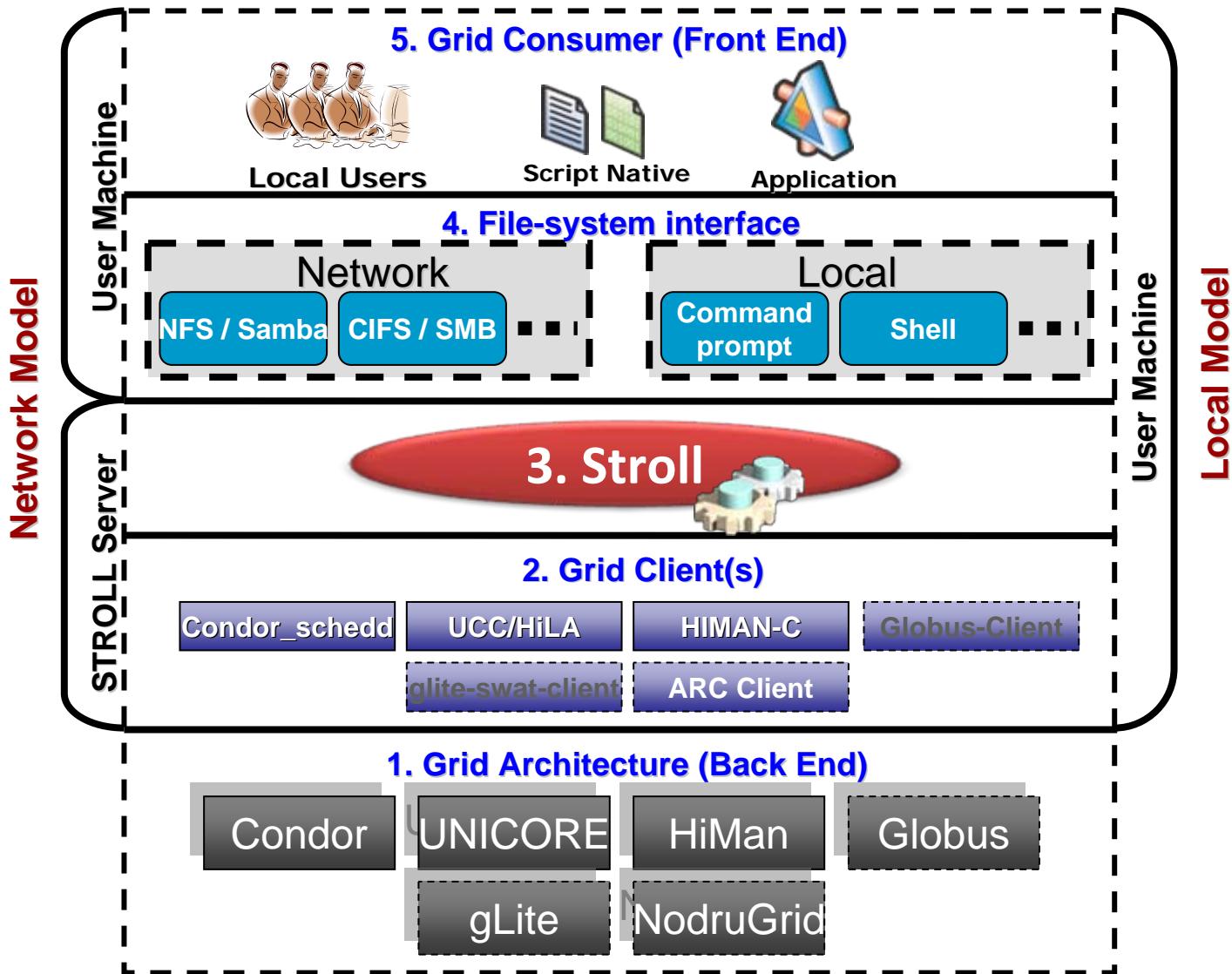


# Setup

- 20 single node UNICORE target systems.
- 40 Condor workers, in 3 labs.

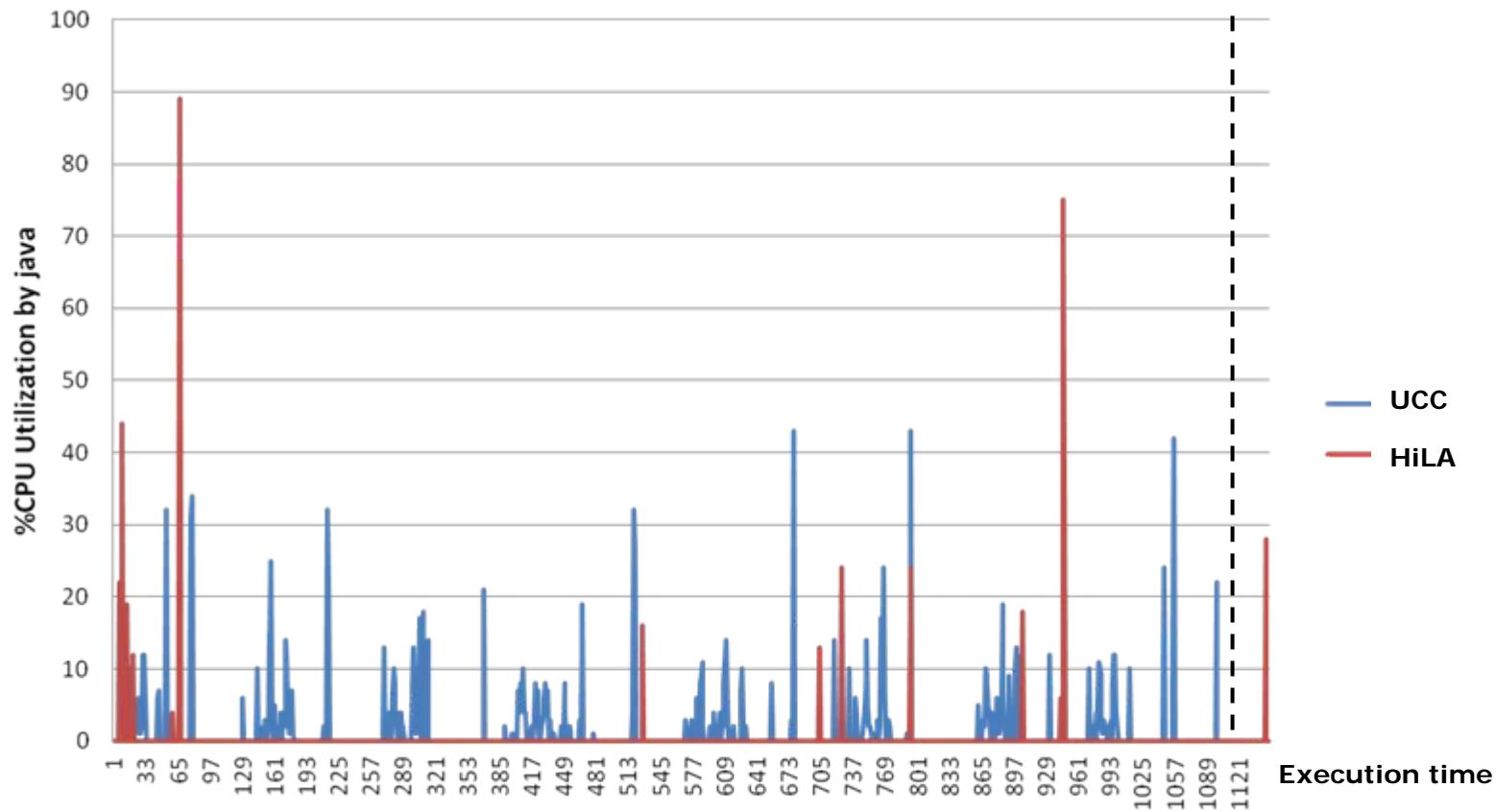


# Stroll Architecture



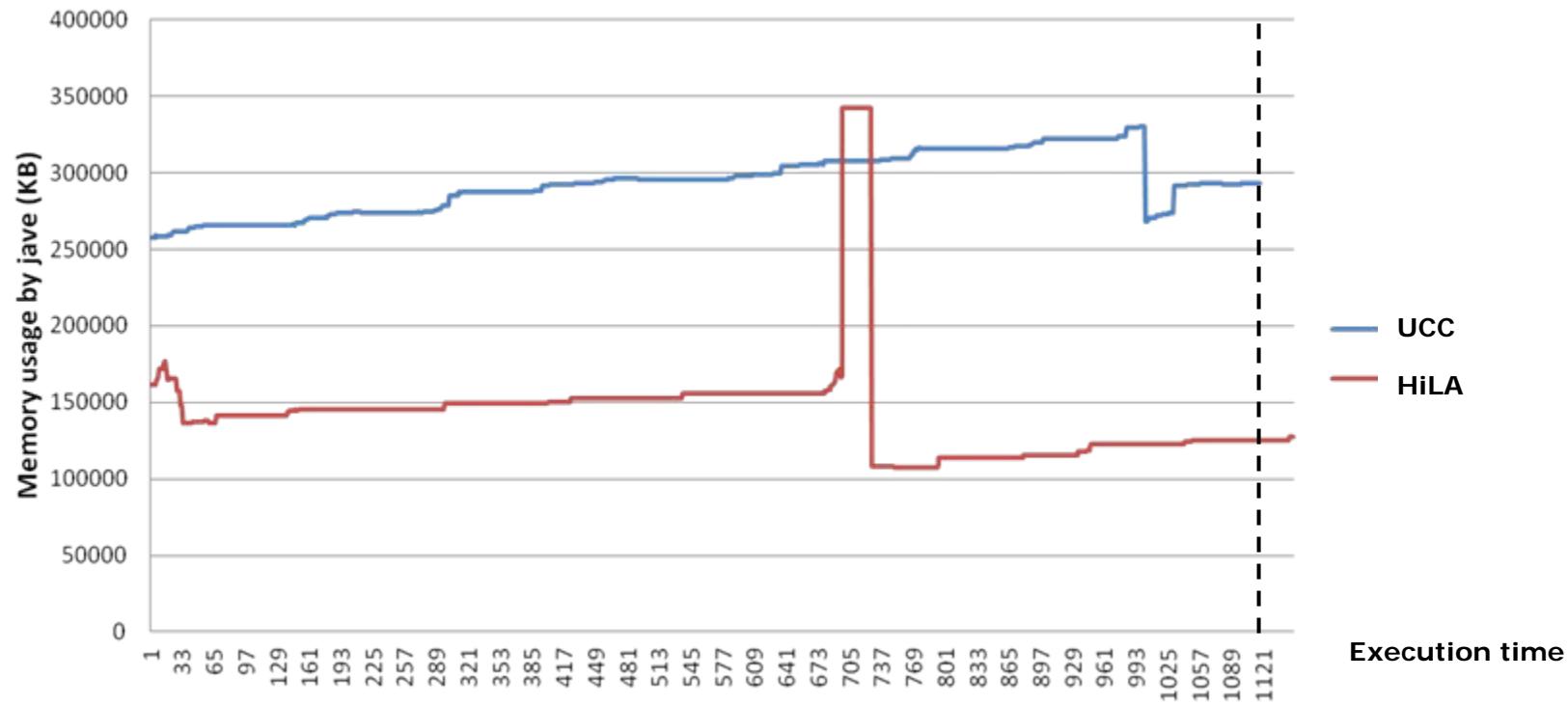
# Results – UNICORE

- PSM.estimate(...,maxit=1,...)



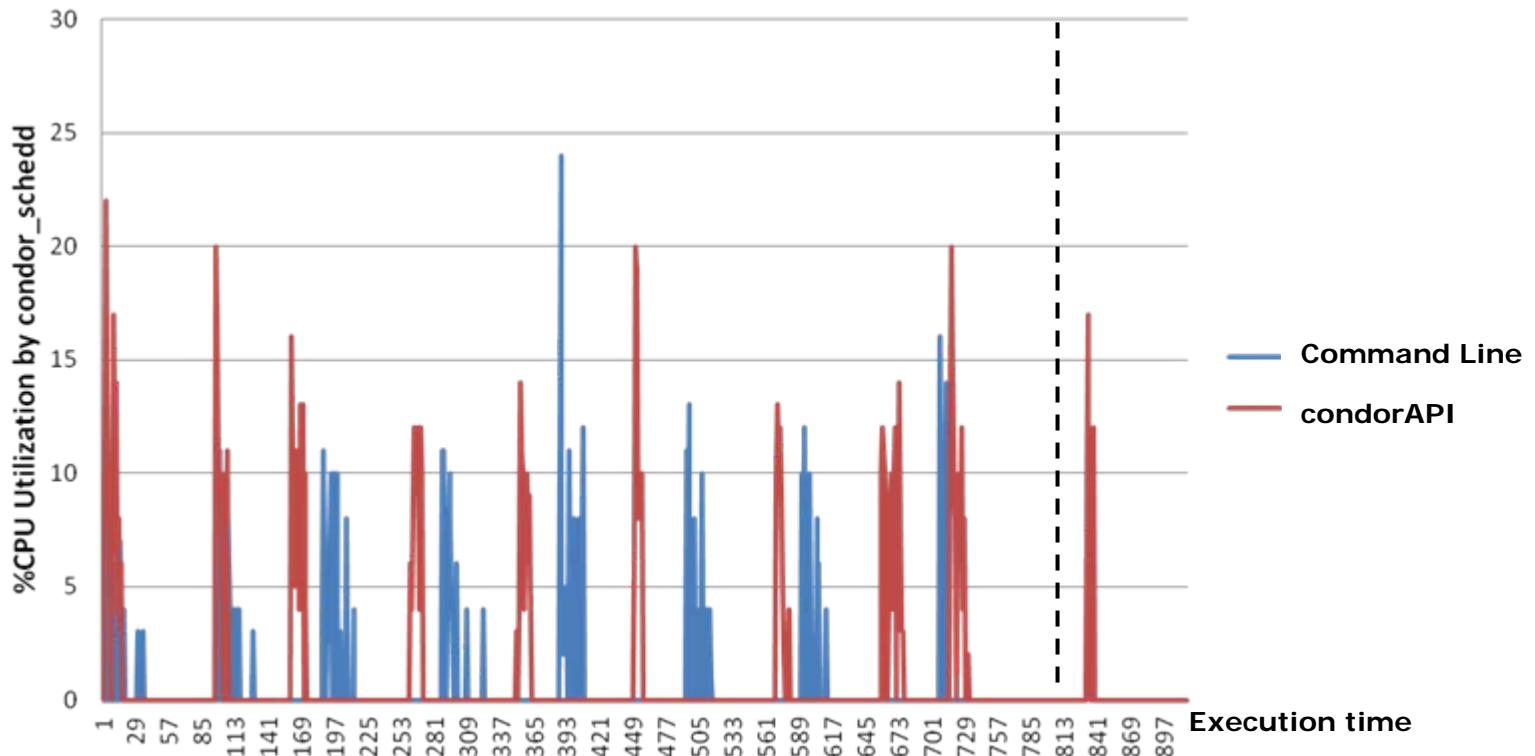
# Results – UNICORE

- PSM.estimate(...,maxit=1,...)



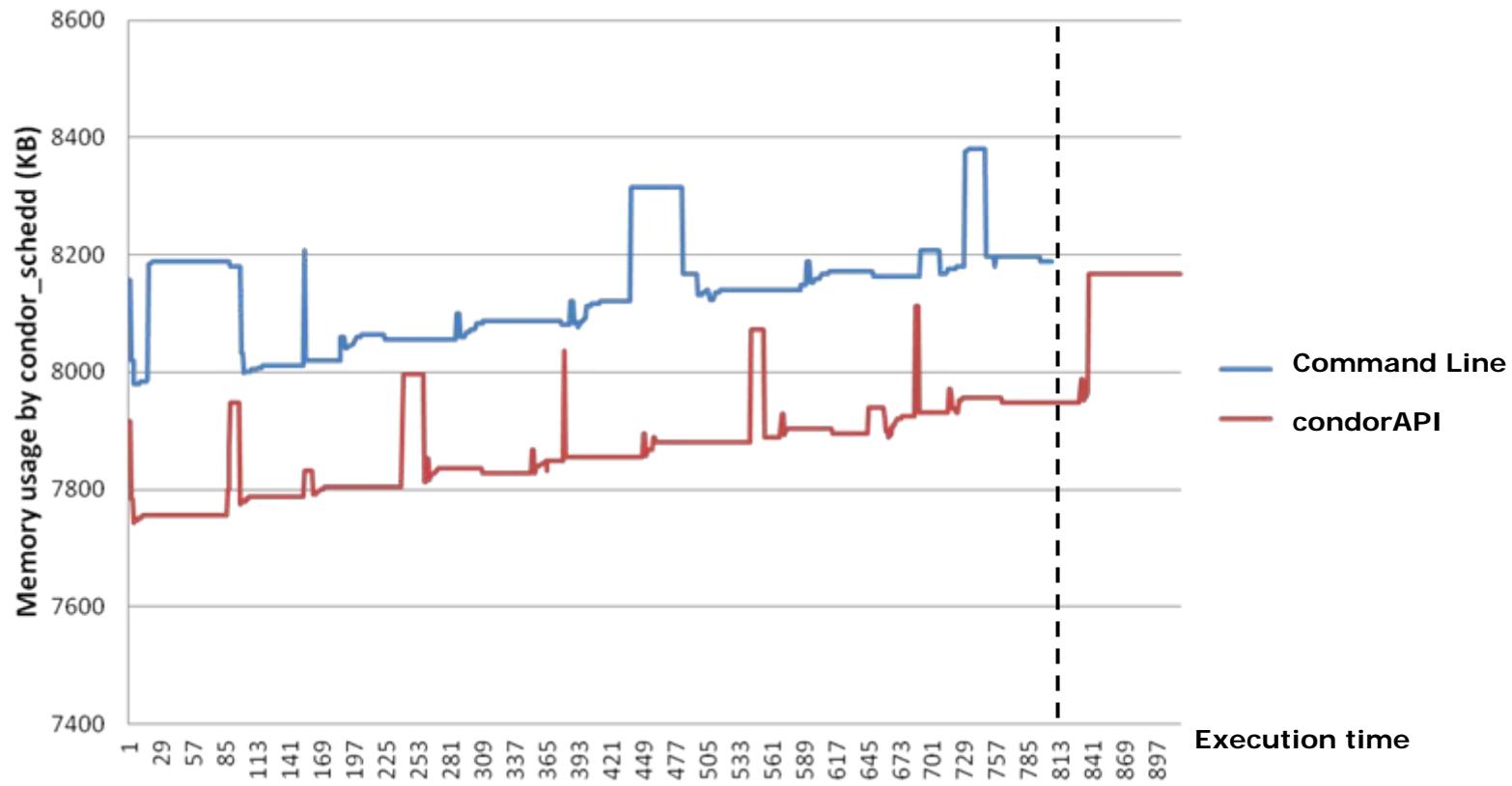
# Results – Condor

- PSM.estimate(...,maxit=1,...)



# Results – Condor

- PSM.estimate(...,maxit=1,...)





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# Questions

