

cloudUtil: Cloud Utilization Visualizations

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Contents

1	Recent changes and updates	2
2	Introduction	2
3	Data preparation	2
4	Analysis	3

1 Recent changes and updates

'vignettes' directory has been migrated.

2 Introduction

`cloudUtil` is a package for creating comparison plots for

Cluster, Grid and Cloud utilization data. Under utilization data we understand collected accounting data measuring the job execution time in the above mentioned environments.

The idea behind this package is to create a single visualization of such data that has the following main features:

- gives an overview over the compute system utilization within a certain time frame
- allows the comparison of job lengths between different platforms giving thus hints on how well the respective job queues function e.g. how efficient the queue of Sun Grid Engine is performing
- allows the integration of replicates within the same visualization
- allows the comparison on both absolute and relative timescales

The functionality of `cloudUtilPlot` function was first used in [3].

3 Data preparation

The package includes sample accounting data for demonstration purposes. These data were collected by comparing the running times of several hundred compute jobs: each one of these jobs performs peptide-spectrum matching in proteomics (data published in [1]).

The fragment below shows a random extract from the dataset provided in the package:

```
> library(cloudUtil)
> data(cloudms2)
> cloudms2[sort(sample(nrow(cloudms2), 10)), c(1, 5, 6, 15)]
```

	CLOUD	BEGIN_PREPROCESS	END_PREPROCESS	id
918	EC2_1	1263521108	1263521118	1408
1703	UZH1	1261651508	1261651510	1747
3890	UZH1	1261637008	1261637030	865
4158	UZH1	1261640717	1261640737	1028
4183	FGCZ2	1263423621	1263423662	900
4248	FGCZ2	1263336489	1263336519	227
5191	UZH2	1263429160	1263429172	1056
7733	FGCZ2	1263443894	1263443907	1109
8677	EC2_2	1263580372	1263580378	1462
10646	FGCZ2	1263304577	1263304598	100

The attributes of interest are CLOUD, BEGIN_PREPROCESS, END_PREPROCESS, and id. Additionally, it is also possible to use accounting data collected from other sources e.g. Sun Grid Engine accounting data [2].

4 Analysis

The code extract below creates a plot of the data shown in Section 3:

```
> hist(cloudms2$END_PREPROCESS - cloudms2$BEGIN_PREPROCESS, 100)
> ##
> boxplot((cloudms2$END_PROCESS-cloudms2$BEGIN_PROCESS)/3600~cloudms2$CLOUD,
+   main="process time",
+   ylab="time [hours]")
> ##
> throughput<-cloudms2$MZXMLFILESIZE*10^-6/
+ (cloudms2$END_COPYINPUT-cloudms2$BEGIN_COPYINPUT)
> boxplot(throughput~cloudms2$CLOUD,
+   main="copy input network throughput",
+   ylab="MBytes/s")
> ##
>
> cloudUtilPlot(begin=cloudms2$BEGIN_PROCESS,
+   end=cloudms2$END_PROCESS,
+   id=cloudms2$id,
+   group=cloudms2$CLOUD)
```

Transparency through alpha blending allows furthermore to compare several plots with each other. An example is given in the code fragment below:

```
> #green
> col.amazon<-rgb(0.1, 0.8, 0.1, alpha=0.2)
> col.amazon2<-rgb(0.1, 0.8, 0.1, alpha=0.2)
> #blue
> col.fgcz<-rgb(0.1, 0.1, 0.8, alpha=0.2)
> col.fgcz2<-rgb(0.1, 0.1, 0.5, alpha=0.2)
> #red
> col.uzh<-rgb(0.8, 0.1, 0.1, alpha=0.2)
> col.uzh2<-rgb(0.5, 0.1, 0.1, alpha=0.2)
> cm<-c(col.amazon, col.amazon2, col.fgcz, col.fgcz2, col.uzh, col.uzh2)
> jpeg("cloudms2Fig.jpg", 640, 640)
> op<-par(mfrow=c(2, 1))
> cloudUtilPlot(begin=cloudms2$BEGIN_PROCESS,
+   end=cloudms2$END_PROCESS,
+   id=cloudms2$id,
+   group=cloudms2$CLOUD,
+   colormap=cm,
+   normalize=FALSE,
```

```
+     plotConcurrent=TRUE);
> cloudUtilPlot(begin=cloudms2$BEGIN_PROCESS,
+     end=cloudms2$END_PROCESS,
+     id=cloudms2$id,
+     group=cloudms2$CLOUD,
+     colormap=cm,
+     normalize=TRUE,
+     plotConcurrent=TRUE,
+     plotConcurrentMax=TRUE)
> dev.off()
```

pdf
2

The output of the above listed R session is shown in Figure 1.

References

- [1] E. Brunner, C. H. Ahrens, S. Mohanty, H. Baetschmann, S. Loevenich, F. Potthast, E. W. Deutsch, C. Panse, U. de Lichtenberg, O. Rinner, H. Lee, P. G. Pedrioli, J. Malmstrom, K. Koehler, S. Schrimpf, J. Krijgsveld, F. Kregenow, A. J. Heck, E. Hafen, R. Schlapbach, and R. Aebersold. A high-quality catalog of the *Drosophila melanogaster* proteome. *Nat. Biotechnol.*, 25(5):576–583, May 2007. [DOI:10.1038/nbt1300] [PubMed:17450130].
- [2] Rayson Ho and Ron Chen. Open grid scheduler. <https://sourceforge.net/projects/gridscheduler>, 2013.
- [3] Aleksandar Markovic. Investigation of economical and practical aspects of commercial cloud computing for life sciences. Master's thesis, 2010.

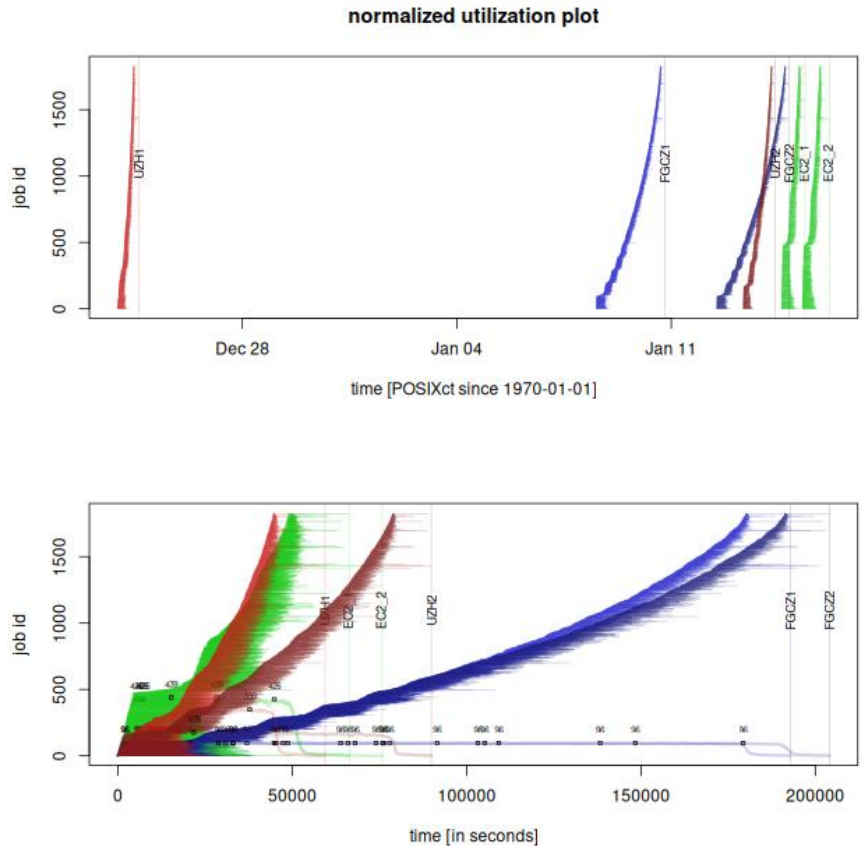


Figure 1: `cloudUtilPlot` visualization for the `cloudms2` data set. On the graphics each horizontal line indicates the start and the end of one single job. Color is used for classifying the different groups. On the upper plot the time of each group was not normalized. The visualization on the bottom on the other side uses normalized time scales which help to compare the compute systems. Tranparent colors were used to dial with the overplotting. The solid lines on the bottom plot show the total number of concurrently running jobs. The squares on the solid lines indicate the maxima on the respective system. The user can make use of all R graphic devices.