

# Reproducible Research for OMNeT++ Based on Python and Pweave

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## Outline

- Reproducible Research
- Python and Pweave
- Reproducible Research for OMNeT++
- Example: OMNeT++ FIFO Simulation

# Reproducible Research

## Reproducible Research

- Reproducible research is a key to any scientific method and ensures repeating an experiment and the results of its analysis in any place with any person.
- A study can be truly reproducible when it satisfies at least the following three criteria:
  - All experimental methods are fully reported.
  - All data and files used for the analysis are (publicly) available.
  - The process of analyzing raw data is well reported and preserved.
- Reproducible research is to ensure
  - Same data + Same script = Same results

## Why Do We Need Reproducible Research: Two Examples

- LIGO - Gravitational Wave Detection
- Schön scandal - Molecular Computing

## LIGO - Gravitational Wave Detection

- The [Laser Interferometer Gravitational-Wave Observatory \(LIGO\)](#) is a large-scale physics experiment and observatory to detect cosmic gravitational waves.

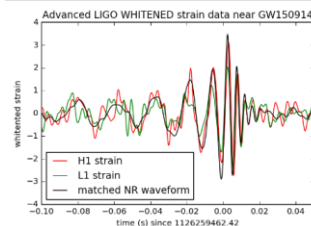
- The detection of gravitational wave was reported in *Physical Review Letters* in Feb. 2016, together with [ipython notebook](#) with analysis code and data.



```
In [9]: # We need to suppress the high frequencies with some bandpassing:
bb, ab = butter(4, [20.*2./fs, 300.*2./fs], btype='band')
strain_H1_whitenbp = filtfilt(bb, ab, strain_H1_whiten)
strain_L1_whitenbp = filtfilt(bb, ab, strain_L1_whiten)
NR_H1_whitenbp = filtfilt(bb, ab, NR_H1_whiten)

# plot the data after whitening:
# first, shift L1 by 7 ms, and invert. See the GW150914 detection paper.
strain_L1_shift = -np.roll(strain_L1_whitenbp, int(0.007*fs))

plt.figure()
plt.plot(time-tevent, strain_H1_whitenbp, 'r', label='H1 strain')
plt.plot(time-tevent, strain_L1_shift, 'g', label='L1 strain')
plt.plot(hitime+0.002, NR_H1_whitenbp, 'k', label='matched NR waveform')
plt.xlim([-0.1, 0.05])
plt.ylim([-4, 4])
plt.xlabel('time (s) since '+str(tevent))
plt.ylabel('whitened strain')
plt.legend(loc='lower left')
plt.title('Advanced LIGO WHITENED strain data near GW150914')
plt.savefig('GW150914_strain_whitened.png')
```



## Schön Scandal - Molecular Computing

- No records found for his groundbreaking experimental results, including lab notebook, experimental samples and data, hard disk drives.
- During the investigation, he kept repeating **"I clearly observed them in the Lab but ..."**

**Bell Labs launches inquiry into allegations of data duplication**

**Misconduct finding at Bell Labs shakes physics community**

**Bell Labs inquiry spreads to superconductors**

**Is a bell tolling for Bell Labs?**

**commentary**

**Paul Davis**

There is a small but growing number of scientists who are skeptical of the results of the investigation into the Schön scandal. They are not alone in their skepticism. The investigation into the Schön scandal has been ongoing for several months, and the results have been mixed. Some scientists believe that the investigation was thorough and that the results are reliable. Others believe that the investigation was biased and that the results are unreliable. The investigation into the Schön scandal has been a major event in the history of science, and it has led to a re-examination of the scientific process. The investigation has shown that there are many ways in which scientists can be misled, and it has led to a re-examination of the scientific process. The investigation has also led to a re-examination of the role of the media in science. The media has played a major role in the investigation, and it has helped to bring the story to the attention of the public. The investigation into the Schön scandal has been a major event in the history of science, and it has led to a re-examination of the scientific process. The investigation has shown that there are many ways in which scientists can be misled, and it has led to a re-examination of the scientific process. The investigation has also led to a re-examination of the role of the media in science. The media has played a major role in the investigation, and it has helped to bring the story to the attention of the public.

## Python and Pweave

## R/Sweave to Python/Pweave

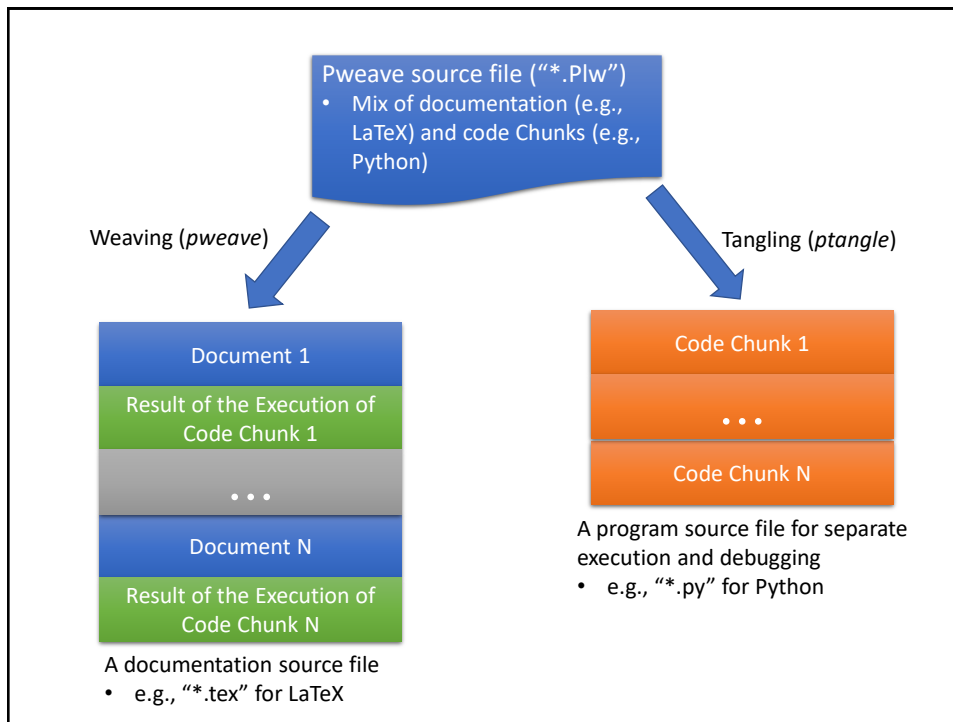
- Until recently, R was the language of choice for statistical processing and data analysis.
  - Still, R has the largest code base for a wide variety of statistical and graphical techniques.
- Like *ipython* (now *jupyter*), R provides a nice tool called *Sweave* (now replaced by *knitr*) to weave documentation and the results of the execution of R code chunks into one source file for integrated documentation.
- Python — one of the most popular languages in scientific computing, including artificial intelligence & machine learning — recently takes over R in statistical processing and data analysis as well.
  - Thanks to [pandas](#) implementing DataFrame object similar to R and [Pweave](#), python can replace R for most statistical and data analysis tasks, while retaining its many advantages over R (i.e., fully-featured programming language with easy syntax and higher speed).

```
### customize
old <- theme_set(theme_bw())
pt_size <- 3.5

## generate summary plots for reference architecture with N=1
rf_N1.data <- paste(rf_N1.wd, paste(rf_N1.base, "data", sep="."), sep="/")
df <- read.csv(rf_N1.data, header=TRUE)
## df <- df[order(df$N, df$dr, df$br, df$repetition), ] # order data frame
df <- sort_df(df, vars=c("N", "n", "dr", "br", "repetition")) # sort data frame
rf_N1.df <- ddply(df, c(.n), .(dr), function(df) {return(GetMeansAndCiWidths(df))})
rf_N1.plots <- list()
for (.i in 1:7) {
  df <- subset(rf_N1.df, select = c(1, 2, (.i*2+1):((.i+1)*2)))
  names(df)[3:4] <- c("mean", "ci.width")
  limits <- aes(ymin = mean - ci.width, ymax = mean + ci.width)
  p <- ggplot(data=df, aes(group=dr, colour=factor(dr), x=n, y=mean)) + geom_line() + scale_y_continuous(limits=limits)
  p <- p + xlab("Number of Users per ONU (n)") + ylab(paste("Line Rate", .i, "Mbps"))
  ## p <- p + geom_point(aes(group=dr, colour=factor(dr), x=n, y=mean), size=pt_size)
  p <- p + geom_point(aes(group=dr, shape=factor(dr), x=n, y=mean), size=pt_size) + scale_shape_manual(values=c(1, 2, 3, 4, 5, 6, 7))
  p <- p + geom_errorbar(limits, width=0.1) + scale_colour_discrete("Line Rate")
  rf_N1.plots[[.i]] <- p
}
```

Snippets of R Source Code  
and Sweave File for LaTeX

```
\subsection{Hybrid PON}
%% tables for dedicated access
%%
<<echo=F,results=tex>>=
df <- subset(hp.df, select=c(1:8))
names(df)[3:8] <- c(
  "dly.mean", "dly.ci.width",
  "thr.mean", "thr.ci.width",
  "trf.mean", "trf.ci.width"
)
tabledf <- xtable(df, caption="Performance measures of FTTP traffic
digits(tabledf)[2:9] <- c(0, 1, rep(-4, 6))
print(tabledf,
  tabular.environment="longtable", caption.placement="top",
  include.rownames=FALSE, floating=FALSE, NA.string="NA")
@
```



## Weaving Example: Automatic Table Generation

The following Python code chunk can automatically generate a long table over multiple pages from a pandas dataframe<sup>1</sup>:

```
<<echo=False, results='raw'>> =
import numpy as np
df = fifo_df.filter(regex="^(?! (r|R)un).*$") # exclude columns starting with run/Run
print(df.to_latex(longtable=True))
@
```

Weaving & LaTeXing

The following Python code chunk can automatically generate a long table over multiple pages from a pandas dataframe<sup>1</sup>:

```
<<echo=False, results='raw'>> =
import numpy as np
df = fifo_df.filter(regex="^(?! (r|R)un).*$") # exclude columns starting with run/Run
print(df.to_latex(longtable=True))
@
```

	File	Module	Name	Unnamed: 19
0	Fifo1-st=0.01-#0.sca	_runattrs_	st	0.010000
1	Fifo1-st=0.01-#0.sca	FifoNet.fifo	queueingTime:mean	0.000262
2	Fifo1-st=0.01-#0.sca	FifoNet.fifo	queueingTime:max	0.031311
3	Fifo1-st=0.01-#0.sca	FifoNet.fifo	busy:timeavg	0.049941
4	Fifo1-st=0.01-#0.sca	FifoNet.fifo	qlen:timeavg	0.001308
5	Fifo1-st=0.01-#0.sca	FifoNet.fifo	qlen:max	4.000000
6	Fifo1-st=0.01-#0.sca	FifoNet.sink	lifetime:mean	0.010262

Continued on next page

<sup>1</sup>Note that a space is inserted between 'r' and 's' to prevent Pweave from weaving the code; it seems that there is no way to escape Pweave chunk code markers.

# Reproducible Research for OMNeT++

## How to Deal with Simulation Input Files

- Include them in the document.
  - OK for small simulations
- Use a snapshot of the whole configurations.
  - e.g., git commit hashes

```
// This file is part of an OMNeT++/OMNEST simulation example.
// Copyright (C) 1992-2015 Andrea Varga
// This file is distributed WITHOUT ANY WARRANTY. See the file
// "license" for details on this and other legal matters.
//
// Simple queueing network: generator + FIFO + sink.
//
network FifoNet
{
  submodules:
  {
    gen: Source {
      parameters:
        @display("p=85,100");
    }
    fifo: Fifo {
      parameters:
        @display("p=205,100");
    }
    sink: Sink {
      parameters:
        @display("p=325,100");
    }
  }
  connections:
    gen.out --> fifo.in;
    fifo.out --> sink.in;
}
```

```
commit 857ae37cd233914fd7271584afc4be10bcf75a61
Author: Kyeong Soo (Joseph) Kim <kyeongsso.kim@gmail.com>
Date: Mon Feb 27 08:59:31 2017 +0000

    Add ini file.

commit f1e7fad0265068d906efd02026e774076c00297
Author: Kyeong Soo (Joseph) Kim <kyeongsso.kim@gmail.com>
Date: Mon Feb 27 08:56:07 2017 +0000

    Remove README.rst; only the markdown version of README

commit 8765336f9e2f5543fea8c4f37a0cf894da7f4c8e
Author: Kyeong Soo (Joseph) Kim <kyeongsso.kim@gmail.com>
Date: Sun Oct 2 17:32:02 2016 +0000

    Change simulation time.
```

Listing 1: 'FifoNet.ned' for FIFO sample model.

## How to Guarantee Match Between Input Files and Output Data


- Online generation of results
  - Include simulation execution code within a document
    - Refer to the provided sample Pweave file.
  - OK for smaller simulations, but not for larger simulations.
- Use a snapshot of the whole configurations and data
  - e.g., git commit hashes
  - Version controlling output data together with source code and input configuration files, however, may greatly increase the size of a repository.

## How to Present and Analyze Output Data

- Unstacking of stacked DataFrame
  - Use *pivot* function (see the example shown here).
- Aggregated processing of measurement data over independent variables
  - Use *pivot\_table* function.
  - Useful for the calculation of mean and confidence intervals over multiple iterations.
- Online calculation of confidence intervals
  - Confidence intervals (CIs) can be calculated by assigning a custom function for CI to *aggfunc* parameter of *pivot\_table* function.
  - Now pandas support error bars in its own plot functions.

```
In [1]: df
Out[1]:
```

	date	variable	value
0	2000-01-03	A	0.469112
1	2000-01-04	A	-0.282863
2	2000-01-05	A	-1.509059
3	2000-01-03	B	-1.135632
4	2000-01-04	B	1.212112
5	2000-01-05	B	-0.173215
6	2000-01-03	C	0.119209
7	2000-01-04	C	-1.044236
8	2000-01-05	C	-0.861849
9	2000-01-03	D	-2.104569
10	2000-01-04	D	-0.494929
11	2000-01-05	D	1.071804



```
In [3]: df.pivot(index='date', columns='variable', values='value')
Out[3]:
```

variable	A	B	C	D
date				
2000-01-03	0.469112	-1.135632	0.119209	-2.104569
2000-01-04	-0.282863	1.212112	-1.044236	-0.494929
2000-01-05	-1.509059	-0.173215	-0.861849	1.071804



## Demo: OMNeT++ FIFO Simulation