

archivist: Managing Data Analysis Results

<https://github.com/pbiecek/archivist>

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About Grupa Wirtualna Polska



The Leader Of The Polish Internet



IT Research and Development

- large-scale online learning
- personalized news article recommendation
- e-mail targeting
- text mining
- web user behavior identification



archivist: What does it do?

Main features

- allows to **store** and **archive** objects in repositories (stored on a local disk or via GitHub/Dropbox)
- provides handy tools facilitating objects' **search** and **recovery**
- ideally performs as **cache**
- supports the philosophy of **reproducible research**



archivist: Why is it useful?

Solves reproducible research problems

- sometimes raw data are large or with limited access
- computations take a lot of time or require specialized hardware
- reproducibility requires specific versions of packages



archivist: Cache Use Case

```
getMaxDistribution <- function(D = rnorm, N = 10,  
                               R = 1000000) {  
  res <- replicate(R, max(D(N)))  
  summary(res)  
}  
  
# or get directly from GitHub  
library(archivist)  
loadFromGithubRepo(md5hash="c", user="MarcinKosinski",  
                   repo="Museum")
```

archivist: Cache Use Case



Using the archivist one can prepare a repository which stores calls and results of the `cache()` function to avoid their re-call in the future.

```
cacheRepo <- tempdir()
createEmptyRepo(cacheRepo)

system.time(cache(cacheRepo, getMaxDistribution,
                  rnorm, 10) )

  user  system elapsed
5.230   0.090   5.315
system.time(cache(cacheRepo, getMaxDistribution,
                  rnorm, 10) )

  user  system elapsed
0.009   0.000   0.008
```

archivist: Retrieving an Object Use Case



Fragment of a poster

This can be reproduced in R with

```
library(RTCGA.mutations);library(RTCGA.rnaseq)
mutationsBox(c('BRCA','HNSC','LUSC','PRAD'),'TP53','ETF1')
```

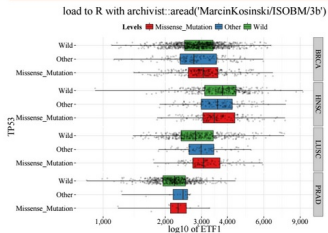


Figure : ggplot object

Using archivist to retrieve an object

```
read('MarcinKosinski/ISOBM/3b')
-> mutationsPlot
plot(mutationsPlot)
```

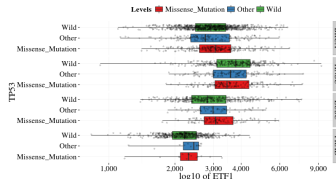


Figure : downloaded object



archivist: Object's Pedigree Use Case

```
createEmptyRepo("FORUM_BI", default = TRUE)
invisible(aoptions("silent", TRUE))
```

```
data(iris)
iris %a%
  dplyr::filter(Sepal.Length < 16) %a%
  lm(Petal.Length ~ Species, data=.) %a%
summary() -> obj
```

```
ahistory(obj)
```

```
iris [ff575c261c949d073b2895b05d1097c3]
-> dplyr::filter(Sepal.Length < 16) [9f7045b7322cdf3a9071377c6fe9c175]
-> lm(Petal.Length ~ Species, data = .) [0a82efeb8250a47718cea9d7f64e5ae7]
-> summary() [671a0b89fccdf02087acb002374a0fcd]
```




archivist: Objects' Exploration Within a Repository

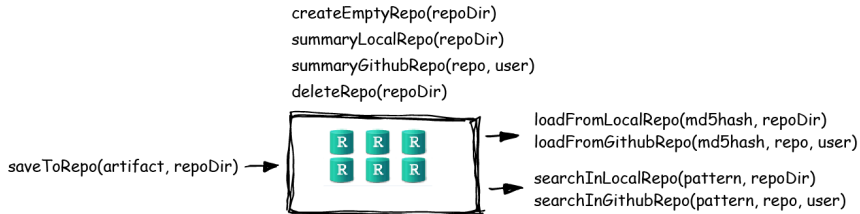
```
models <- asearch("pbiecek/graphGallery",  
  patterns = c("class:lm", "coefname:Sepal.Length"))  
lapply(models, coef)
```

```
[[1]]  
  (Intercept) Sepal.Length  
    -7.101443      1.858433
```

```
[[2]]  
  (Intercept)      Sepal.Length Speciesversicolor Speciesvirginica  
    -1.7023422         0.6321099         2.2101378         3.0900021
```



archivist: How does it work?



Each repository contains a database with objects metadata.

Objects are stored as binary files.

Each object has a unique key - md5 hash.

Metadata, like object class, name, creation date, relations with other objects are useful when searching for an object in a repository.

```
|| library("archivist")
```

archivist: Plans & Prototypes



Automated repository creation on github, commit, push and a return of a hook to an object.

```
archive(iris, "MarcinKosinski",
  "archivist-Museum-RforeverR_last3",
  USER_EMAIL, USER_PASSWORD, app_key, app_secret,
  github_token) -> aread_input
# function returns a hook
archivist::aread("MarcinKosinski/archivist-Museum-
  RforeverR_last3/ff575c261c949d073b2895b05d1097c3")
archivist::aread(aread_input) -> x
digest::digest(x)

[1] "ff575c261c949d073b2895b05d1097c3"

identical(iris,x)

[1] TRUE
```



archivist: Learn More

<http://pbiecek.github.io/archivist/>