



SYBERIA: A DEVELOPMENT FRAMEWORK FOR R

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- Packages work well for recording abstract solutions to problems, but not for large projects maintained by multiple users tied to solving problems in a specific domain

- R workflows are typically loosely organized collections of scripts
- Packages work well for recording abstract solutions to problems, but not for large projects maintained by multiple users tied to solving problems in a specific domain
- Test-driven development is difficult for modeling work that is intended for real-time systems



Machine Learning: The High Interest Credit Card of
Technical Debt (2014) NIPS 2014 Workshop proceedings
D. Sculley, Gary Holt, Daniel Golovin, et al

“Risk factors include boundary erosion, entanglement, hidden feedback loops, undeclared consumers, data dependencies, changes in the external world, and a variety of system-level anti-patterns.”

Solution: Think like a developer

Developers are good at simplifying work that needs to be done down to its core abstractions



Syberia is a framework for building complex projects in R.

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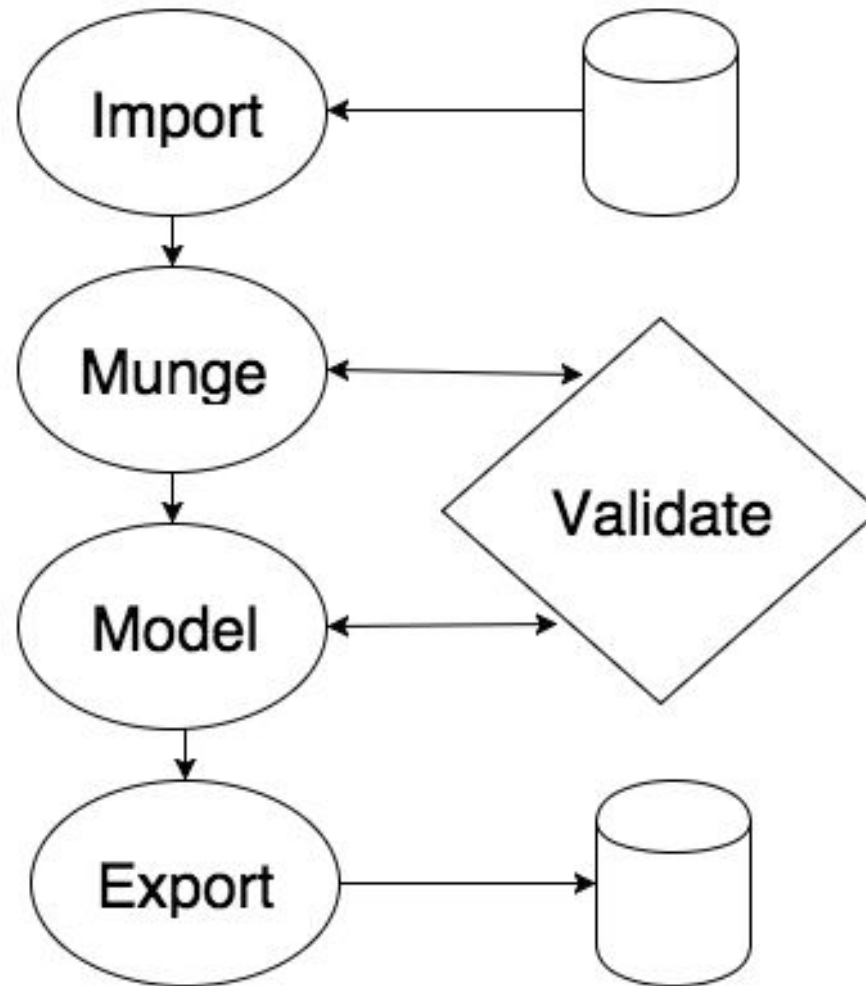
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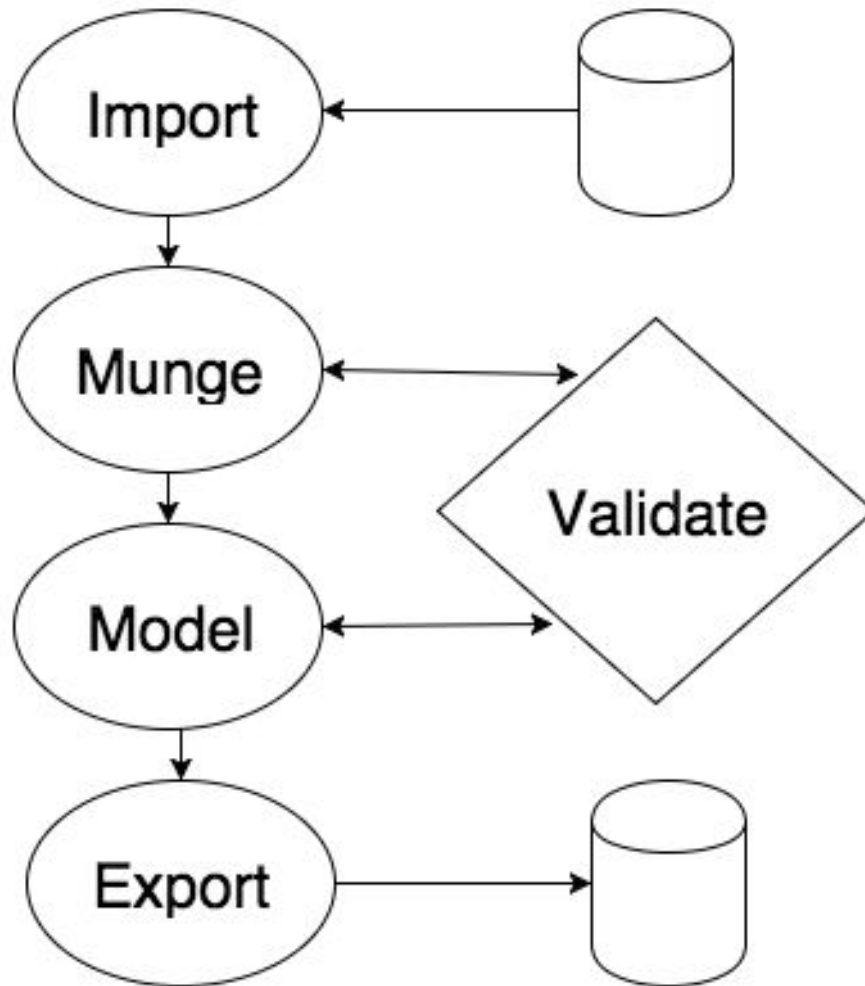
The modular design unit is an **engine**.

Today we are releasing the **modeling engine** for building and deploying production-ready machine learning products in R.

How Does It Work?



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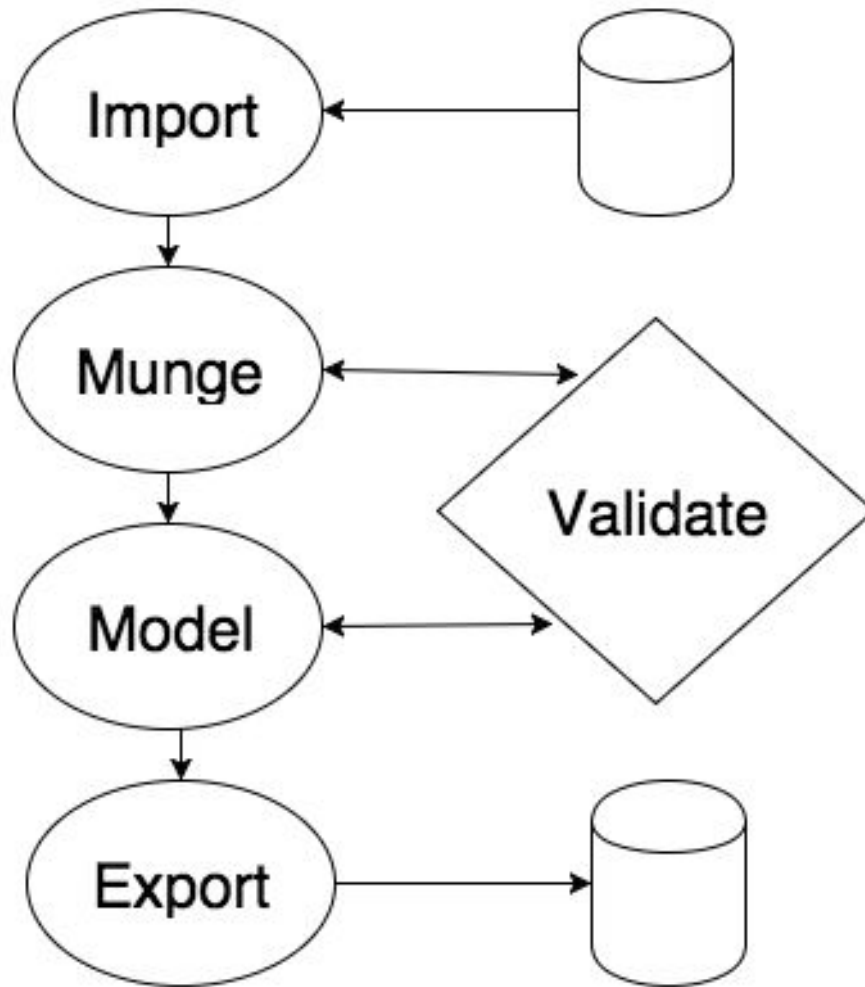


The modeling engine is the boss.

Mungebits package powers feature engineering.

Tundra package powers model containers.

How Does It Work?



The modeling engine is the boss.

Director package holds the project together.

Stagerunner package is the workflow and execution system.

```
source("my_R_script.R")
```

- Script-driven workflow makes it harder to re-use components
- Testing is not built in unless you move to a package

```
resource ("lib/adapters/s3")
```

- **Basic idea:** Everything is a resource
- All resources must be tested
(test/lib/adapters/s3)
- Each resource type can have its own “grammar”

Task: Let's import some data
using Syberia.

An Example

```
# lib/adapters/s3.R
read <- function(name) {
  s3mpi::s3read(name)
}

write <- function(object, name) {
  s3mpi::s3store(object, name)
}

# package: github.com/robertzk/s3mpi
```

An Example

```
# config/routes.R
list(
  "config/global"      = "globals",
  "lib/adapters"      = "adapters",
  "lib/classifiers"    = "classifiers",
  "lib/mungebits"      = "mungebits",
  "models"             = "models",
  "test/models"        = "test/models",
  "data"               = "data")
```

The Basic Structure

```
a <- resource("lib/adapters/s3")
```

```
a$write(iris, "tmp/iris")
```

```
# From a new R session
```

```
a <- resource("lib/adapters/s3")
```

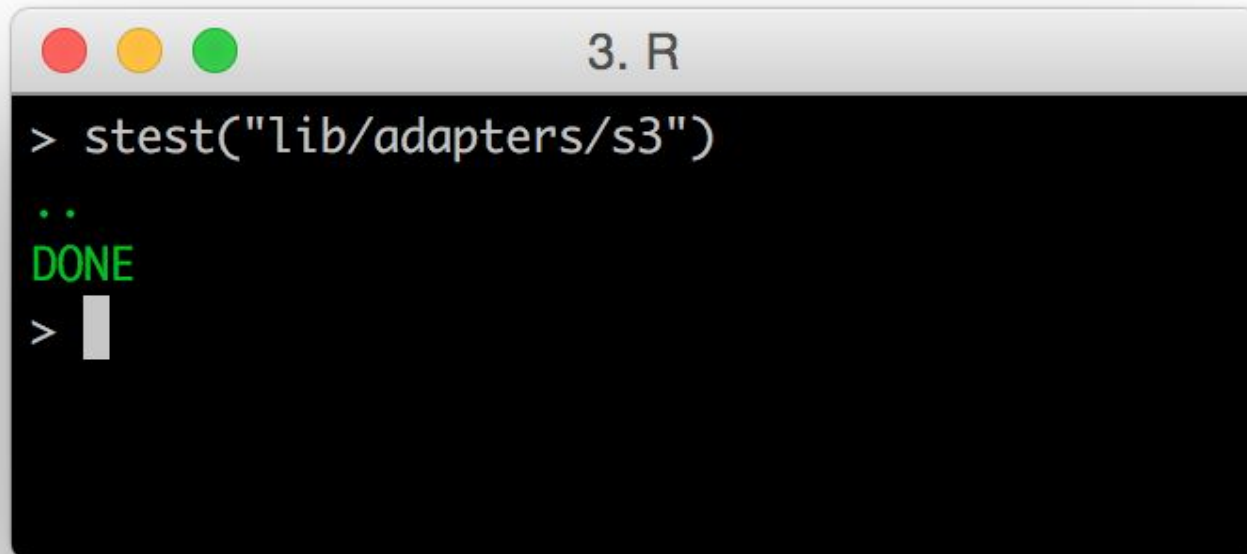
```
identical(  
  a$read("tmp/iris"),  
  iris  
)
```

An Example

```
# test/lib/adapters/s3.R
```

```
test_that("it can write a data set to S3", {  
  env <- new.env()  
  package_stub("s3mpi", "s3store", function(...) { env[[..2]] <- ..1 }, {  
    adapter <- resource()  
    adapter$write(iris, "test_key", prefix = "")  
    expect_identical(env$test_key, iris,  
      info = "iris should have been stored in the test_key in env")  
  })  
})  
  
test_that("it can read a data set from S3", {  
  env <- list2env(list(test_key = iris))  
  package_stub("s3mpi", "s3read", function(...) { env[[..1]] }, {  
    adapter <- resource()  
    expect_identical(adapter$read("test_key", prefix = ""), env$test_key,  
      info = "iris should have been read from the test_key in env")  
  })  
})
```

An Example



```
> stest("lib/adapters/s3")  
..  
DONE  
> |
```

An Example

```
# config/routes.R
list(
  "config/global"      = "globals",
  "lib/adapters"      = "adapters",
  "lib/classifiers"    = "classifiers",
  "lib/mungebits"      = "mungebits",
  "models"             = "models",
  "test/models"        = "test/models",
  "data"               = "data")
```

An Example

```
# lib/controllers/adapters.R
function(input) {
  adapter_class <- function(r, w) {
    list(read = r, write = w)
  }

  # Construct the adapter object.
  adapter_class(
    input$read,
    input$write
  )
}
```


An Example

```
# lib/adapters/s3.R
read <- function(name) {
  s3mpi::s3read(name)
}

write <- function(object, name) {
  s3mpi::s3store(object, name)
}

# package: github.com/robertzk/s3mpi
```

Think of how you might write adapters for other storage backends:

- Reading from and writing to a CSV file
- Reading from and writing to a database
- Reading from and writing to a JSON service
- Et cetera

An Example

- Defining “adapters” abstracted away the storage backend from the underlying implementation
- Each adapter has the same interface

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- Each adapter has the same interface

```
adapter <- resource("lib/adapters/s3")  
adapter$read("some_key")  
adapter$write(object, "some_key")
```

- Defining “adapters” abstracted away the storage backend from the underlying implementation
- Each adapter has the same interface

```
adapter <- resource("lib/adapters/s3")  
adapter$read("some_key")  
adapter$write(object, "some_key")
```

```
adapter <- resource("lib/adapters/file")  
adapter$read("some_file.rds")  
adapter$write(iris, "some_file.csv")
```

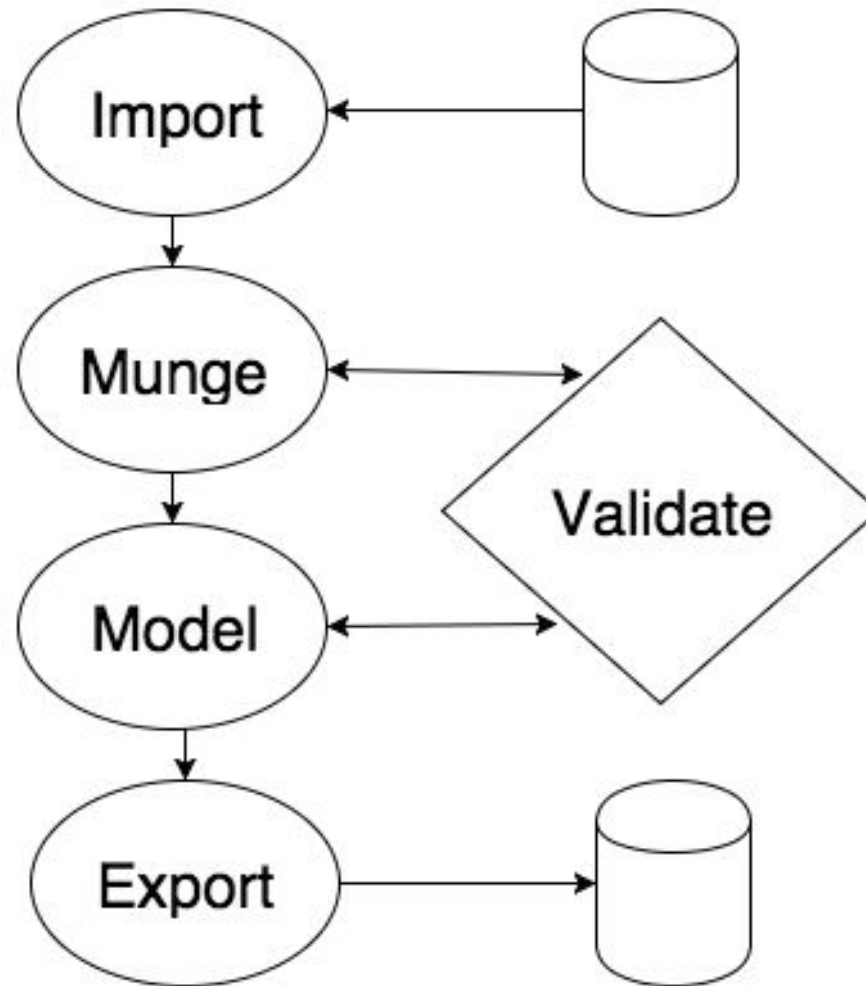
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- Each resource produces a single R object
- Encourages finding common interfaces and abstractions
- Work becomes easily re-usable instead of locked away in scripts

The Modeling Engine



The Modeling Engine

```
list(  
  # Titanic dataset is fairly popular. Here we're downloading it from a public github repo  
  
  import = list(url = "https://raw.githubusercontent.com/haven-jeon/.../master/bicdata/data/titanic.csv"),  
  
  data = list(  
    # The left-hand side defines the informal name of a mungebit that you will see when you run this model.  
    # The right-hand side is the mungebit invocation.  
    "Factor to character" = list(column_transformation(as.character), is.factor)  
    , "has paren in name" = list(multi_column_transformation(function(name) grepl("(", fixed = TRUE, name)), "name", "has_paren")  
    , "Name length variable" = list(new_variable, function(name) nchar(name), "name_length")  
    # ~40 removed  
    , "Restore levels" = list(restore_categorical_variables)  
    , "Rename dep_var" = list(renamer, c("survived" = "dep_var"))  
  ),  
  
  model = list('gbm'  
    , .id_var = 'X'  
    , distribution = 'bernoulli'  
    , number_of_trees = 3000  
    , shrinkage_factor = 0.005  
  ),  
  
  export = list(R = 'titanic')  
)
```

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export = list(R = 'titanic')  
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```
# config/routes.R
list(
  "config/global"      = "globals",
  "lib/adapters"       = "adapters",
  "lib/classifiers"    = "classifiers",
  "lib/mungebits"      = "mungebits",
  "models"             = "models",
  "test/models"        = "test/models",
  "data"               = "data")
```

Sort data in ascending order by a given column.

```
# lib/mungebits/orderer.R
train <- predict <- function(dataframe, col) {
  dataframe[order(dataframe[[col]]), ]
}

# From R console
m <- resource("lib/mungebits/orderer")
stopifnot(all.equal(
  m$run(iris, 1), iris[order(iris[[1]]), ]
))
```

Mean imputation for one column.

```
# lib/mungebits/simple_impute.R
train <- function(dataframe, col) {
  input$col <- col
  input$mean <- mean(dataframe[[col]], na.rm=T)
  dataframe[is.na(dataframe[[col]]), col] <-
    input$mean
  dataframe
}
predict <- function(dataframe, ...) {
  col <- input$col
  dataframe[is.na(dataframe[[col]]), col] <-
    input$mean
  dataframe
}
```

```
# In R console
m <- resource("lib/mungebits/simple_impute")
iris2 <- iris; iris2[1, 1] <- NA
m$run(iris2, 1)
stopifnot(all.equal(
  m$run(iris2)[[1]],
  c(mean(iris[-1, 1]), iris2[-1, 1])
))
```


Mungebits

```
# lib/controllers/mungebits.R
function(input) {
  if (isTRUE(input$column_transformation)) {
    mungebits2::mungebit$new(
      mungebits2::column_transformation(input$train),
      mungebits2::column_transformation(input$predict))
  } else {
    mungebits2::mungebit$new(
      input$train, input$predict)
  }
}
```

Classifiers

```
# lib/controllers/classifiers.R
```

```
function(input) {  
  force(input)  
  function(munge_procedure = list(), default_args = list(), internal = list()) {  
    input <- lapply(as.list(input), full_deflate)  
  
    container <- tundra::tundra_container$new(resource, input$train, input$predict,  
      munge_procedure, full_deflate(default_args), full_deflate(internal))  
  
    container$hooks <- lapply(container$hooks, function(fn) {  
      environment(fn) <- globalenv(); fn  
    })  
  
    if (!is.null(input$read) ||  
        !is.null(input$write)) {  
      attr(container, "s3mpi.serialize") <- list(read = input$read, write = input$write)  
    }  
    container  
  }  
}
```

The Modeling Engine

```
list(  
  # Titanic dataset is fairly popular. Here we're downloading it from a public github repo  
  
  import = list(url = "https://raw.githubusercontent.com/haven-jeon/.../master/bicdata/data/titanic.csv"),  
  
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    , "Name length variable" = list(new_variable, function(name) nchar(name), "name_length")  
    # ~40 removed  
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    , "Rename dep_var" = list(renamer, c("survived" = "dep_var"))  
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  model = list('gbm'  
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    , distribution = 'bernoulli'  
    , number_of_trees = 3000  
    , shrinkage_factor = 0.005  
  ),  
  
  export = list(R = 'titanic')  
)
```

```
list(  
  # Import and data stage  
  model = list('gbm'  
    , .id_var          = 'X'  
    , distribution      = 'bernoulli'  
    , number_of_trees   = 3000  
    , shrinkage_factor  = 0.005  
  )  
  # Export stage  
)
```

The Models Controllers

lib/controllers/models/models.R

```
construct_stage_runner <- Ramd::define("construct_stage_runner")[[1]](resource)
preprocessor <- Ramd::define("preprocessor")[[1]]
function(args, resource, output, director, modified, any_dependencies_modified) {
  parent.env(parent.env(environment(construct_stage_runner))) <- environment()
  if (is.element("raw", names(args))) return(output)
  require(objectdiff)
  message("Loading model: ", resource)
  tests <- file.path('test', resource)
  has_tests <- director$exists(tests)
  has_tests <- FALSE
  if (has_tests) {
    testrunner <- stageRunner$new(new.env(), director$resource(tests))
    testrunner$transform(function(fn) {
      library(testthat); force(fn)
      function(after) fn(cached_env, after)
    })
  }
  model_version <- gsub("^\\w+/", "", resource)
  if (isTRUE(args$fresh) || !identical(resource, director$cache_get("last_model"))) {
    stagerunner <- construct_stage_runner(output, model_version)
  } else if (modified || any_dependencies_modified) {
    message(crayon::yellow("Copying cached environments..."))
    stagerunner <- construct_stage_runner(output, model_version)
    stagerunner$coalesce(director$cache_get("last_model_runner"))
  } else if (!director$cache_exists("last_model_runner")) {
    stagerunner <- construct_stage_runner(output, model_version)
  } else {
    stagerunner <- director$cache_get("last_model_runner")
  }
  if (has_tests) stagerunner$overlay(testrunner, "tests", flat = TRUE)
  director$cache_set("last_model", resource)
  director$cache_set("last_model_runner", stagerunner)
  stagerunner
}
```

../models/preprocessor.R

```
preprocessor <- function(resource, director, source_env) {
  source_env$extending <- function(model_version, expr) {
    eval.parent(substitute(within(resource(file.path("models/", model_version), raw = TRUE), { expr })))
  }

  source_env$model_version <- version <- gsub("^[/]+\\[/]+\\/", "", resource)
  source_env$model_name <- basename(version)
  source_env$output <-
    function(suffix = "", create = TRUE, dir = file.path(director$root(), "tmp")) {
      filename <- file.path(dir, version, suffix)
      if (create && !file.exists(dir <- dirname(filename)))
        dir.create(dir, recursive = TRUE)
      filename
    }
  lexicals <- director$resource("lib/shared/lexicals")
  for (x in ls(lexicals)) source_env[[x]] <- lexicals[[x]]
  director$resource("lib/shared/source_mungebits")(source_env, director)
  model <- source()
  if (nzchar(Sys.getenv("CI"))) {
    model$import <- NULL
  }
  }
  model
}
```

../models/construct_stage_runner.R

```
model_env <- function() {
  if (identical(getOption("environment_type"), "environment")) {
```

- Other useful abstractions with their own grammar:
 - **Modules:** Nested bundles of related R functions.
 - **Indicators:** $y \sim x1 + x2$ tied to an ETL backend
 - **Stages:** Execution tasks for our stage runner.
 - **Jobs:** Build reports, monitoring checks, etc.
 - **Queries:** “Object-relational mapper” for R

Running A Model

```
run("example") # Runs models/dev/example.R
```

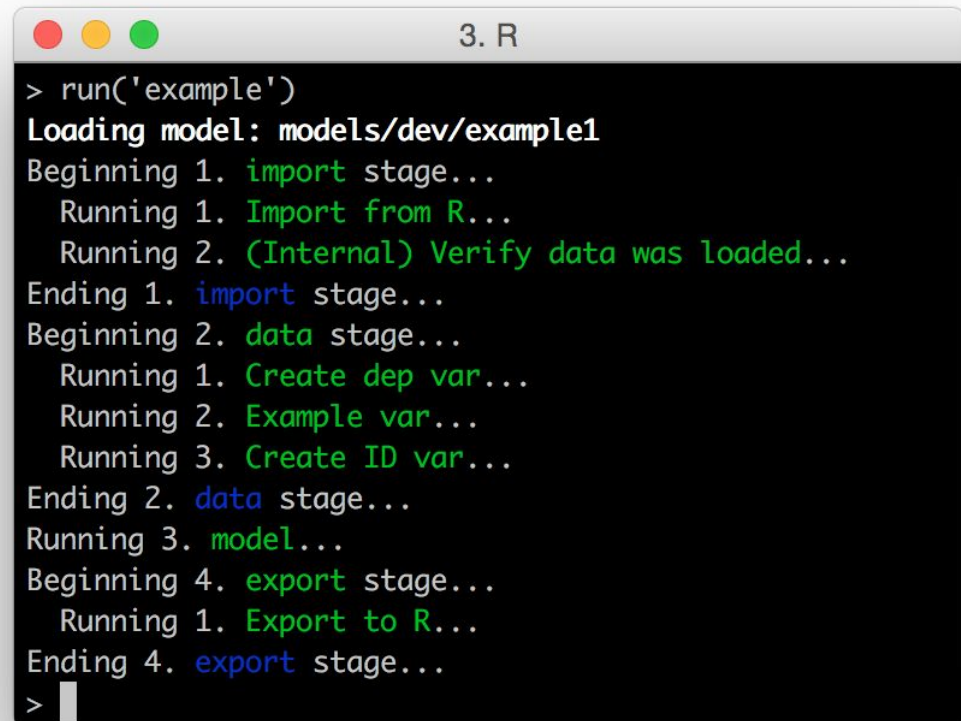
```
list(  
  import = list(R = "iris"),  
  data = list(  
    "Create dep var" = list(renamer,  
      c("Sepal.Length" = "dep_var")),  
    "Example var" = list(example),  
    "Create ID var" = list(  
multi_column_transformation(seq_along),  
      "dep_var", "id")  
  ),  
  model = list("lm", .id_var = "id"),  
  export = list(R = "model")  
)
```

```
3. R  
> run('example')  
Loading model: models/dev/example1  
Beginning 1. import stage...  
  Running 1. Import from R...  
  Running 2. (Internal) Verify data was loaded...  
Ending 1. import stage...  
Beginning 2. data stage...  
  Running 1. Create dep var...  
  Running 2. Example var...  
  Running 3. Create ID var...  
Ending 2. data stage...  
Running 3. model...  
Beginning 4. export stage...  
  Running 1. Export to R...  
Ending 4. export stage...  
> 
```

Running A Model

```
run("example") # Runs models/dev/example.R
```

```
list(  
  import = list(R = "iris"),  
  data = list(  
    "Create dep var" = list(renamer,  
      c("Sepal.Length" = "dep_var")),  
    "Example var" = list(example),  
    "Create ID var" = list(  
multi_column_transformation(seq_along),  
      "dep_var", "id")  
  ),  
  model = list("lm", .id_var = "id"),  
  export = list(R = "model")  
)
```



```
3. R  
> run('example')  
Loading model: models/dev/example1  
Beginning 1. import stage...  
  Running 1. Import from R...  
  Running 2. (Internal) Verify data was loaded...  
Ending 1. import stage...  
Beginning 2. data stage...  
  Running 1. Create dep var...  
  Running 2. Example var...  
  Running 3. Create ID var...  
Ending 2. data stage...  
Running 3. model...  
Beginning 4. export stage...  
  Running 1. Export to R...  
Ending 4. export stage...  
> 
```

```
model$predict(iris) # [1] 5.005 4.757 4.890 ...
```


Running A Model

```
run(, "data/3") # Re-runs one munge step
```

```
list(  
  import = list(R = "iris"),  
  data = list(  
    "Create dep var" = list(renamer  
      c("Sepal.Length" = "dep_var")),  
    "Example var" = list(example),  
    "Create ID var" = list(  
multi_column_transformation(seq_along),  
      "dep_var", "id")  
  ),  
  model = list("lm", .id_var = "id"),  
  export = list(R = "model")  
)
```

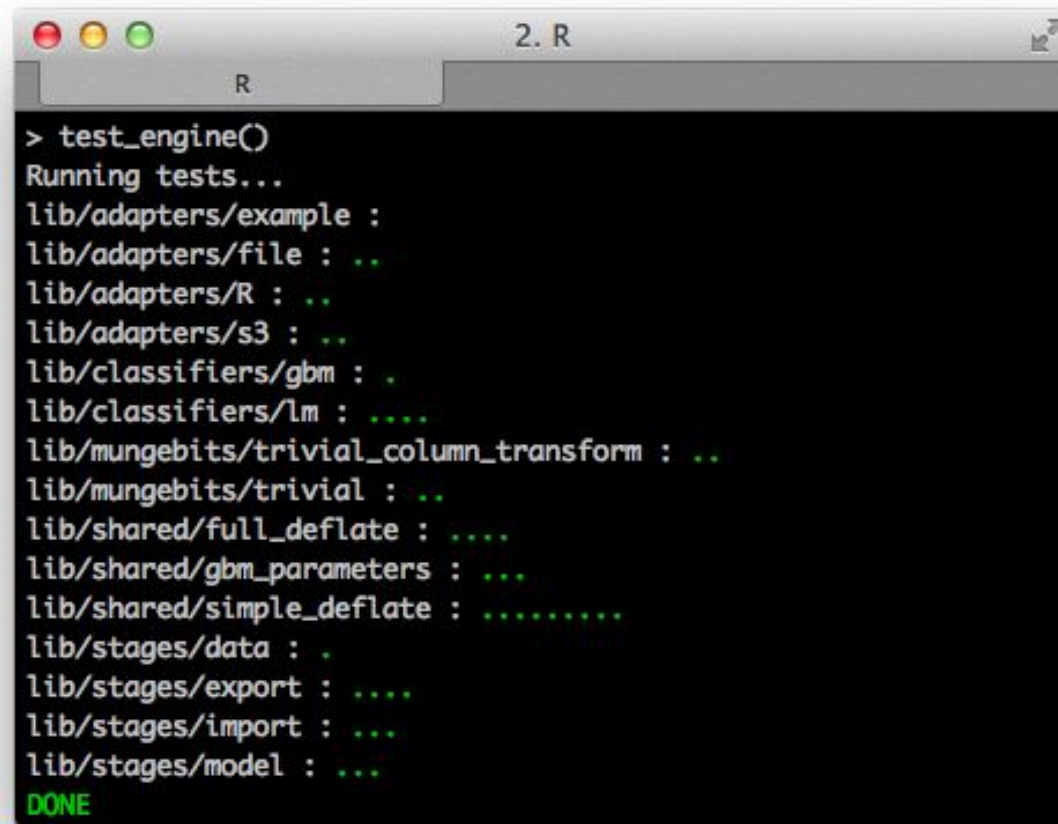


```
> run("data/3")  
Loading model: models/dev/example1  
Beginning 2. data stage...  
  Running 3. Create ID var...  
Ending 2. data stage...  
> dim(B)  
[1] 150  5  
> dim(A)  
[1] 150  6  
> setdiff(ls(A), ls(B))  
[1] "id"  
> 
```

```
setdiff(ls(B), ls(A)) # [1] "id"
```

Each resource requires an accompanying test

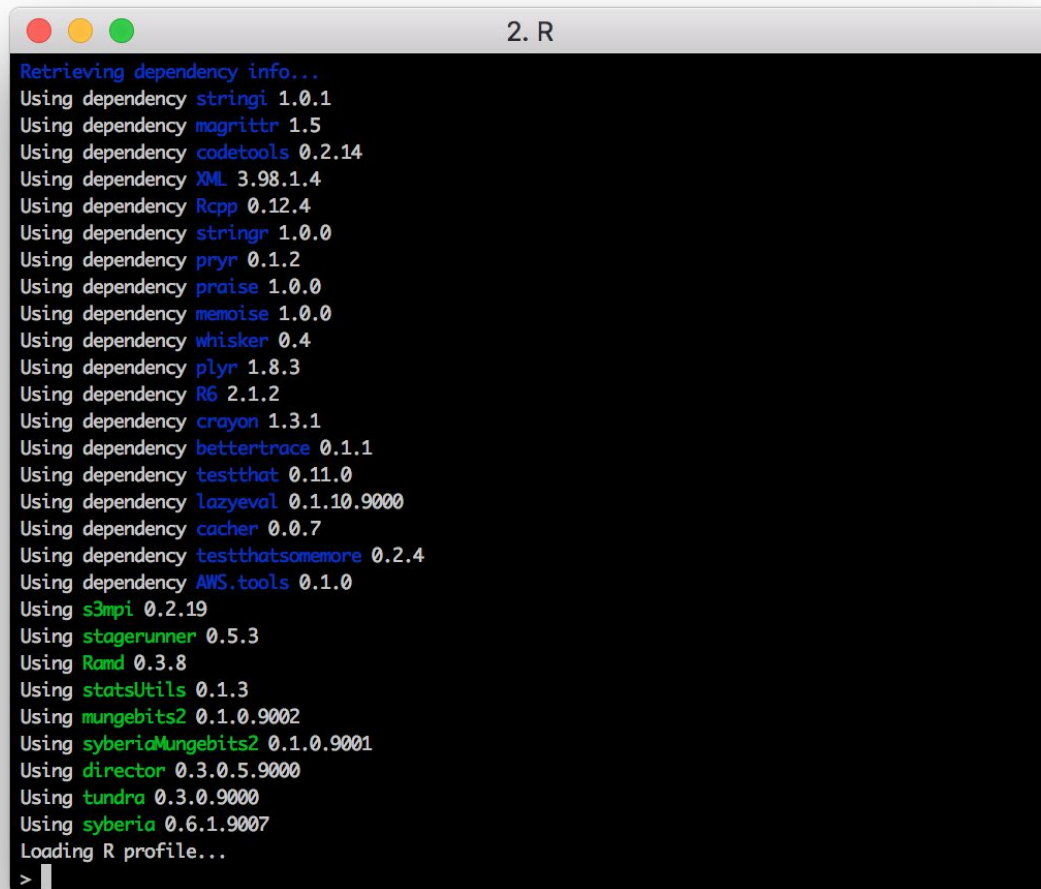
`test_project()`



```
> test_engine()
Running tests...
lib/adapters/example : 
lib/adapters/file : ..
lib/adapters/R : ..
lib/adapters/s3 : ..
lib/classifiers/gbm : .
lib/classifiers/lm : .....
lib/mungebits/trivial_column_transform : ..
lib/mungebits/trivial : ..
lib/shared/full_deflate : .....
lib/shared/gbm_parameters : ...
lib/shared/simple_deflate : .....
lib/stages/data : .
lib/stages/export : .....
lib/stages/import : ...
lib/stages/model : ...
DONE
```

Dependency Management

Everyone working on the project has the same set of dependencies

A terminal window titled "2. R" with a dark background and light-colored text. It displays the output of an R command to retrieve and install dependencies. The output lists 34 packages and their versions, each preceded by "Using dependency". The packages include stringi, magrittr, codetools, XML, Rcpp, stringr, pryr, praise, memoise, whisker, plyr, R6, crayon, bettertrace, testthat, lazyeval, cachier, testthatsomemore, AWS.tools, s3mpi, stagerunner, Rcmdr, statsUtils, mungebits2, syberiaMungebits2, director, tundra, and syberia. The process concludes with "Loading R profile..." and a prompt character ">".

```
2. R
Retrieving dependency info...
Using dependency stringi 1.0.1
Using dependency magrittr 1.5
Using dependency codetools 0.2.14
Using dependency XML 3.98.1.4
Using dependency Rcpp 0.12.4
Using dependency stringr 1.0.0
Using dependency pryr 0.1.2
Using dependency praise 1.0.0
Using dependency memoise 1.0.0
Using dependency whisker 0.4
Using dependency plyr 1.8.3
Using dependency R6 2.1.2
Using dependency crayon 1.3.1
Using dependency bettertrace 0.1.1
Using dependency testthat 0.11.0
Using dependency lazyeval 0.1.10.9000
Using dependency cachier 0.0.7
Using dependency testthatsomemore 0.2.4
Using dependency AWS.tools 0.1.0
Using s3mpi 0.2.19
Using stagerunner 0.5.3
Using Rcmdr 0.3.8
Using statsUtils 0.1.3
Using mungebits2 0.1.0.9002
Using syberiaMungebits2 0.1.0.9001
Using director 0.3.0.5.9000
Using tundra 0.3.0.9000
Using syberia 0.6.1.9007
Loading R profile...
> 
```

Dependency Management



syberia/lockbox

```
# lockfile.yml
```

```
packages:
```

```
-
  name: devtools
  version: 1.12.0
  repo: hadley/devtools
  ref: v1.12.0
-
  name: checkr
  version: 0.1.4
  repo: syberia/checkr
-
  name: s3mpi
  version: 0.2.40
  repo: robertzk/s3mpi
-
  name: objectdiff
  version: 0.2.3.9003
  repo: robertzk/objectdiff
-
  name: stagerunner
  version: 0.5.6
  repo: syberia/stagerunner
-
  name: Ramd
  version: 0.3.8
  repo: robertzk/Ramd
-
  name: statsUtils
  version: 0.1.4
  repo: robertzk/statsUtils
-
  name: mungebits2
  version: 0.1.0.9014
  repo: syberia/mungebits2
-
  name: syberiaMungebits2
  version: 0.1.2.9002
  repo: syberia/syberiaMungebits2
-
  name: director
  version: 0.3.0.5.9000
  repo: syberia/director
-
  name: tundra
  version: 0.3.0.9000
  repo: syberia/tundra
-
  name: syberia
  version: 0.6.1.9009
  repo: syberia/syberia
  ref: 0.6.1.9009
```

Scales to large teams of contributors working on **thousands of R models**

 **25,778** commits



 **517** branches



 **34** contributors



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