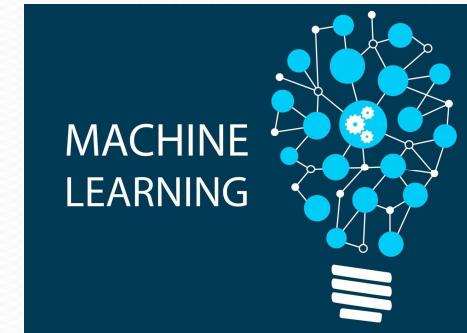


# Local PLS with package Jchemo



matthieu.lesnoff@cirad.fr



SensorFINT 08/09/2022 Sète, France

<https://julialang.org>



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# The Julia Programming Language

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## Julia in a Nutshell

### Fast

Julia was designed from the beginning for [high performance](#). Julia programs compile to efficient native code for [multiple platforms](#) via LLVM.

### Dynamic

Julia is [dynamically typed](#), feels like a scripting language, and has good support for [interactive use](#).

### Reproducible

[Reproducible environments](#) make it possible to recreate the same Julia environment every time, across platforms, with [pre-built binaries](#).

### Composable

Julia uses [multiple dispatch](#) as a paradigm, making it easy to express many object-oriented and [functional programming](#) patterns. The talk on the [Unreasonable Effectiveness of Multiple Dispatch](#) explains why it works so well.

### General

Julia provides [asynchronous I/O](#), [metaprogramming](#), [debugging](#), [logging](#), [profiling](#), a [package manager](#), and more. One can build entire [Applications](#) and [Microservices](#) in Julia.

### Open source

Julia is an open source project with over 1,000 contributors. It is made available under the [MIT license](#). The source code is available on [GitHub](#).

# Download Julia

 Star 40,198

Please star us [on GitHub](#). If you use Julia in your research, please [cite us](#). If possible, do consider [sponsoring us](#).

## Current stable release: v1.8.0 (August 17, 2022)

Checksums for this release are available in both [MD5](#) and [SHA256](#) formats.

<b>Windows</b> <a href="#">[help]</a>	64-bit (installer), 64-bit (portable)	32-bit (installer), 32-bit (portable)
<b>macOS x86 (Intel or Rosetta)</b> <a href="#">[help]</a>	64-bit (.dmg), 64-bit (.tar.gz)	
<b>macOS ARM (M-series Processor)</b> <a href="#">[help]</a>	64-bit	
<b>Generic Linux on x86</b> <a href="#">[help]</a>	64-bit (glibc) (GPG), 64-bit (musl) <sup>[1]</sup> (GPG)	32-bit (GPG)
<b>Generic Linux on ARM</b> <a href="#">[help]</a>	64-bit (AArch64) (GPG)	
<b>Generic FreeBSD on x86</b> <a href="#">[help]</a>	64-bit (GPG)	
<b>Source</b>	Tarball (GPG)	Tarball with dependencies (GPG) <a href="#">GitHub</a>

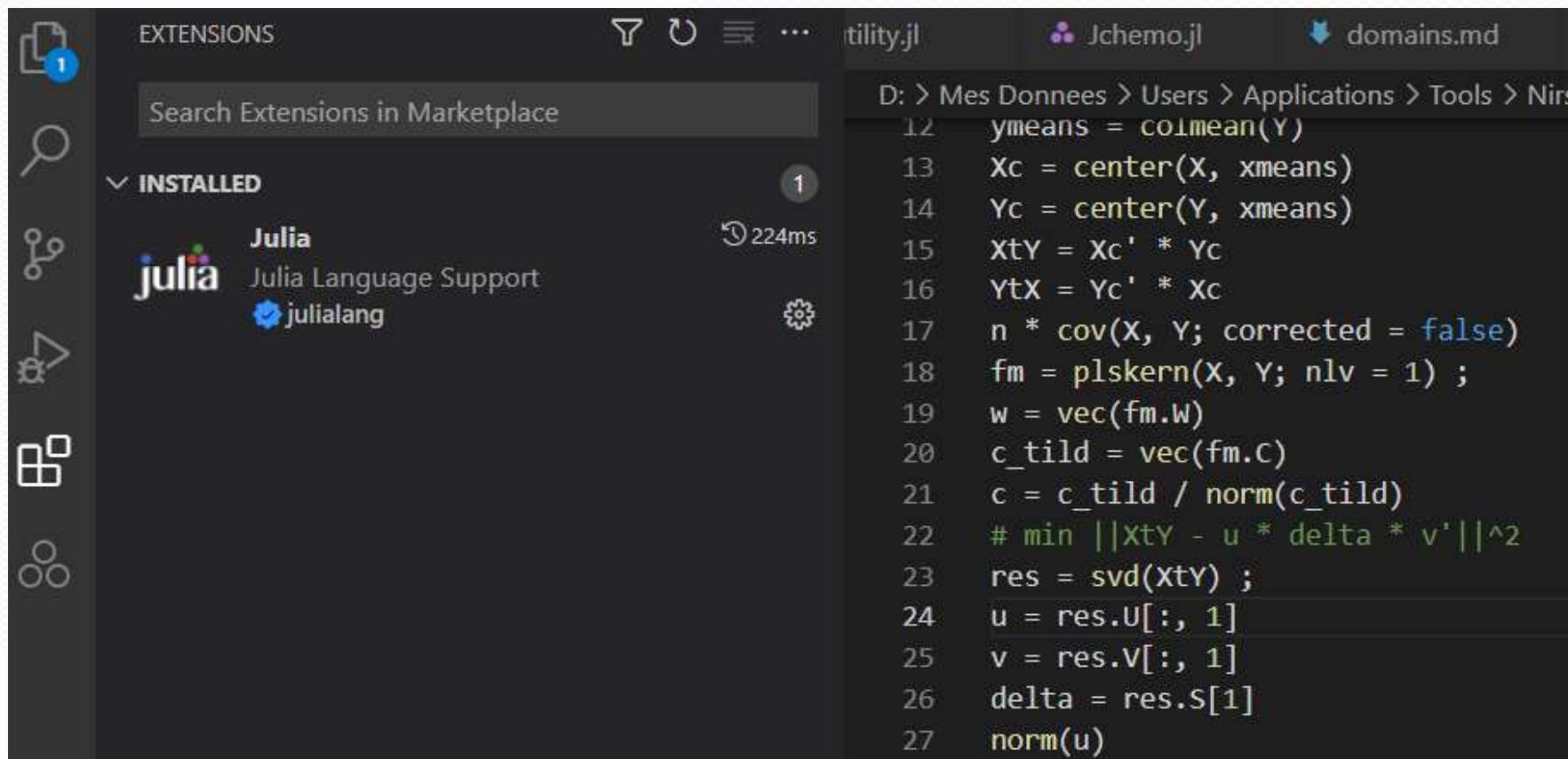
<https://code.visualstudio.com>

The screenshot shows the official Visual Studio Code website. At the top, there's a navigation bar with links for Visual Studio Code, Docs, Updates, Blog, API, Extensions, FAQ, Learn, a search bar labeled "Search Docs", and a "Download" button.

The main content area features a large heading "Code editing. Redefined." with the subtext "Free. Built on open source. Runs everywhere." Below this is a "Download for Windows" button for the "Stable Build". There's also a link to "Web, Insiders edition, or other platforms". A note below the download button states: "By using VS Code, you agree to its [license](#) and [privacy statement](#)".

To the right, a screenshot of the Visual Studio Code interface is displayed. It shows the code editor with several tabs open: "App.js", "index.js", and "serviceWorker.js". The "serviceWorker.js" tab contains code related to service workers. The left sidebar shows the "EXTENSIONS: MARKETPLACE" section with various extensions listed, such as Python, GitLens, C/C++, ESLint, Debugger for Chrome, Language Support, vscode-icons, Vetur, and C#. The bottom of the screenshot shows the terminal window with the command "node" and some status indicators like "master" and "Ln 43, Col 19".

# Install Julia extension under VsCode



# Package Jchemo

<https://github.com/mlesnoff/Jchemo.jl>

master ▾ 2 branches 0 tags Go to file Add file ▾ Code ▾

mlesnoff	Update plsr_avg.jl	✓ 877b7e8 16 hours ago	⌚ 253 commits
📁 .github/workflows	Update ci.yml	3 months ago	
📁 docs	0.0.24	3 days ago	
📁 src	Update plsr_avg.jl	16 hours ago	
📁 test	0.0.17	3 months ago	
📄 LICENSE.md	v0.0.0	15 months ago	
📄 Project.toml	0.0.24	3 days ago	
📄 README.md	0.0.24	3 days ago	

☰ README.md

## Jchemo.jl

Dimension reduction, Regression and Discrimination for Chemometrics

Version 0.0.24

docs dev CI passing repo status Active

About

Julia package for regression and discrimination, with focus on chemometrics and high-dimensional data

machine-learning kernel random-forest  
svm fda julia-language xgboost  
pca preprocessing knn  
chemometrics discrimination  
local-regression ridge  
one-class-classification multiblock plsda  
lwplsrl plsr lwplsda

Readme View license 1 star 1 watching 1 fork

Releases

No releases published Create a new release

Packages

# Documentation

Jchemo.jl

Search docs

Home

Available methods

- PCA
- RANDOM PROJECTIONS
- REGRESSION
- DISCRIMINATION ANALYSIS (DA)
- TUNING MODELS
- DATA MANAGEMENT
- PLOTTING
- UTILITIES

Index of functions

News

Version dev

Non linear

- **rpmat\_ll** Sparse random projection matrix

## REGRESSION

### Linear models

*Multiple linear regression (MLR)*

- **mlr** QR algorithm
- **mlrchol** Normal equations and Choleski factorization
- **mlrpinv** Pseudo-inverse
- **mlrpinv\_n** Normal equations and pseudo-inverse
- **mlrvec** Simple linear regression (Univariate x)

Anova

- **aov1** One factor ANOVA

### Partial least squares (PLSR)

- **plskern** "Improved kernel #1" *Dayal & McGregor 1997*
- **plsnipals** NIPALS
- **plsrosa** ROSA *Liland et al. 2016*
- **plssimp** SIMPLS *de Jong 1993*

*Variants*

- **cglsr** Conjugate gradient for the least squares normal equations (CGLS)
- **covselr** MLR on variables selected from partial correlation or covariance (Covsel)
- **pcr** SVD factorization

## Benchmark

```
n = 10^6 ; p = 500 ; q = 10
```

```
X = rand(n, p)
```

```
Y = rand(n, q)
```

```
nlv = 25
```

```
@time plskern(X, Y[:, 1]; nlv = nlv) ;
```

**7.1 seconds** (50 allocations: 4.001 GiB, 3.45% gc time)

```
@time plskern(X, Y; nlv = nlv) ;
```

**7.3 seconds** (9.76 k allocations: 4.130 GiB, 7.71% gc time, 0.17% compilation time)

## Examples of syntax

### Fitting a model

```
using Jchemo  
  
n = 150 ; p = 200 ; q = 2 ; m = 50  
Xtrain = rand(n, p) ; Ytrain = rand(n, q) ;  
Xtest = rand(m, p) ; Ytest = rand(m, q) ;  
  
nlv = 5  
fm = plskern(Xtrain, Ytrain; nlv = nlv) ;  
pnames(fm)  
  
summary(fm, Xtrain, Ytrain)  
  
transform(fm, Xtest)  
transform(fm, Xtest; nlv = 1)  
  
coef(fm)  
coef(fm; nlv = 2)  
  
res = predict(fm, Xtest) ;  
res.pred  
rmsep(res.pred, Ytest)  
mse(res.pred, Ytest)
```

Fictive data

Model fitting

LVs computations  
(projections)

b-coefs

Predictions

Error rates (scores)

# Tuning models by grid search

Runs over all the combinations of the model parameters  
and compute the error rate (RMSEP etc.)  $\Rightarrow$  Model selection

```
par1 = [.1 ; .5 ; 1]  
par2 = ["a" ; "e"]
```

```
par1 par2  
0.1  "a"  
0.5  "a"  
1.0  "a"  
0.1  "e"  
0.5  "e"  
1.0  "e"
```

Function mpar  
builds the grid

```
pars = mpar(par1 = par1, par2 = par2)
```

- Functions **gridscore**, **gridscorelv**, **gridscorelb**

Train = Cal + Val

`gridscore(Xcal, Ycal, Xval, Yval ; ....)`

- Functions **gridcv**, **gridcvlv**, **gridcvlb**

Cross-validation

```
segm = segmkf(ntrain, 5; rep = 5)      # Replicated K-fold cross-validation  
#segm = segmts(ntrain, 30; rep = 5)    # Replicated test-set validation
```

`gridcv(Xtrain, Ytrain ; ....)`

- With gridscore

```
using Jchemo, CairoMakie

n = 150 ; p = 200 ; m = 50
Xtrain = rand(n, p) ; ytrain = rand(n)
Xval = rand(m, p) ; yval = rand(m)

nlv = 0:10
pars = mpar(nlv = nlv)
res = gridscore(
    Xtrain, ytrain, Xval, yval;
    score = rmsep, fun = plskern, pars = pars)

plotgrid(res.nlv, res.y1,
    xlabel = "Nb. LVs", ylabel = "RMSEP").f

u = findall(res.y1 .== minimum(res.y1))[1]
res[u, :]
fm = plskern(Xval, yval; nlv = res.nlv[u]) ;
res = Jchemo.predict(fm, Xval) ;
rmsep(res.pred, yval)

## For PLSR models, using gridscorelv is much faster than gridscore!!!

res = gridscorelv(
    Xtrain, ytrain, Xval, yval;
    score = rmsep, fun = plskern, nlv = nlv)
```

- With gridcv

```

using Jchemo

n = 150 ; p = 200 ; m = 50
Xtrain = rand(n, p) ; ytrain = rand(n)
Xval = rand(m, p) ; yval = rand(m)

segm = segmkf(n, 5; rep = 5)      # Replicated K-fold cross-validation
#segm = segmts(n, 30; rep = 5)    # Replicated test-set validation

nlv = 0:10
pars = mpar(nlv = nlv)
zres = gridcv(
    Xtrain, ytrain; segm,
    score = rmsep, fun = plskern, pars = pars) ;
pnames(zres)
res = zres.res

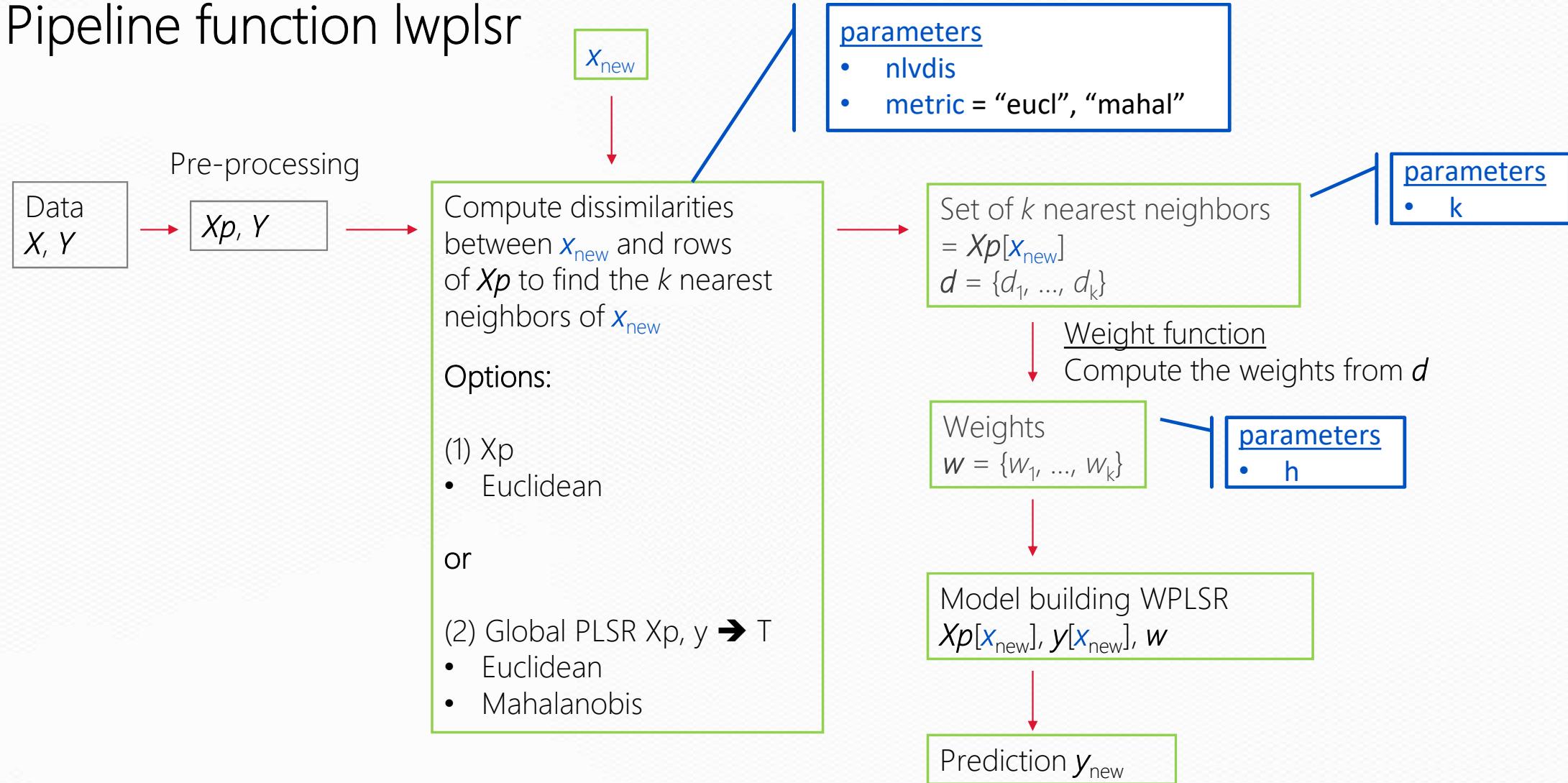
plotgrid(res.nlv, res.y1,
         xlabel = "Nb. LVs", ylabel = "RMSEP").f

u = findall(res.y1 .== minimum(res.y1))[1]
res[u, :]
fm = plskern(Xval, yval; nlv = res.nlv[u]) ;
res = Jchemo.predict(fm, Xval) ;
rmsep(res.pred, yval)

## For PLSR models, using gridcvlv is much faster than gridcv!!!
zres = gridcvlv(
    Xtrain, ytrain; segm,
    score = rmsep, fun = plskern, nlv = nlv) ;
zres.res

```

# Pipeline function lwplsR



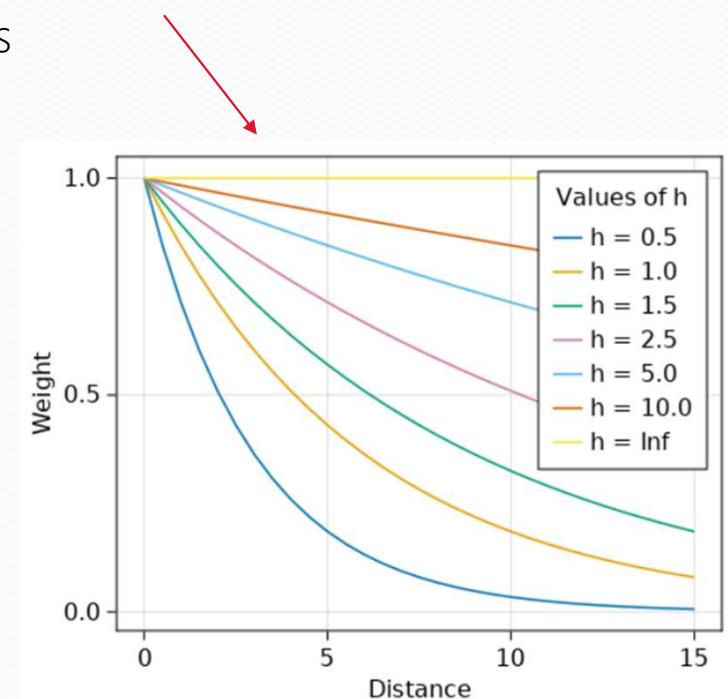
## SYNTAX

```
nlvdis = 20 ; metric = "mahal"      # Mahalanobis distance on 20 global PLS scores  
h = 1; k = 100                      # k neighbors, weight function h = 1  
nlv = 15                            # Nb. LVs in local models
```

```
fm = lwpls(Xtrain, ytrain; nlvdis = nlvdis,  
           metric = metric, h = h, k = k, nlv = nlv) ;
```

```
pred = predict(fm, Xtest).pred
```

```
rmsep(pred, ytest)
```



- **GRIDSCORE**

```
nlvdis = [25] ; metric = ["mahal"]  
h = [1; 2; 5] ; k = [100; 250; 500]
```

```
pars = mpar(nlvdis = nlvdis, metric = metric, h = h, k = k)      # Build the grid  
nlv = 0:25
```

```
res = gridscorelv(Xcal, ycal, Xval, yval; score = rmsep,  
    fun = lwpls, nlv = nlv, pars = pars)  
u = findall(res.y1 .== minimum(res.y1))[1]
```

```
fm = lwpls(Xtrain, ytrain; nlvdis = res.nlvdis[u],  
    metric = res.metric[u], h = res.h[u], k = res.k[u],  
    nlv = res.nlv[u], verbose = true) ;
```

```
pred = predict(fm, Xtest).pred
```

```
rmsep(pred, ytest)
```

- **GRIDCV**

```
nlvdis = [25] ; metric = ["mahal"]
h = [1; 2; 5] ; k = [100; 250; 500]

pars = mpar(nlvdis = nlvdis, metric = metric, h = h, k = k)      # Build the grid
nlv = 0:25

m = 100 ; segm = segmts(ntrain, m; rep = 6)      # Test-set CV
#K = 3 ; segm = segmkf(ntrain, K; rep = 2)      # K-fold CV

res = gridcvlv(Xtrain, ytrain; segm = segm, score = rmsep,
    fun = lwpls, nlv = nlv, pars = pars).res
u = findall(res.y1 .== minimum(res.y1))[1]
res[u, :]

fm = lwpls(Xtrain, ytrain; nlvdis = res.nlvdis[u],
    metric = res.metric[u], h = res.h[u], k = res.k[u],
    nlv = res.nlv[u]) ;

pred = predict(fm, Xtest).pred

rmsep(pred, ytest)
```

# Practical work

## forages2

NIRS data from mixed forages (dried and grounded): stems, leaves etc. Origin: mainly tropical African areas. FOSS NiRS System Instruments 1100-2498 nm (step = 2 nm). Data being private, spectra have been preprocessed with Savitzky-Golay ( $d = 2$ ). Response variables:

- DM: dry matter content
- NDF: fibers content

Source: CIRAD, [Selmet research unit](#)

```
julia> Y = dat.Y
```

```
485x4 DataFrame
```

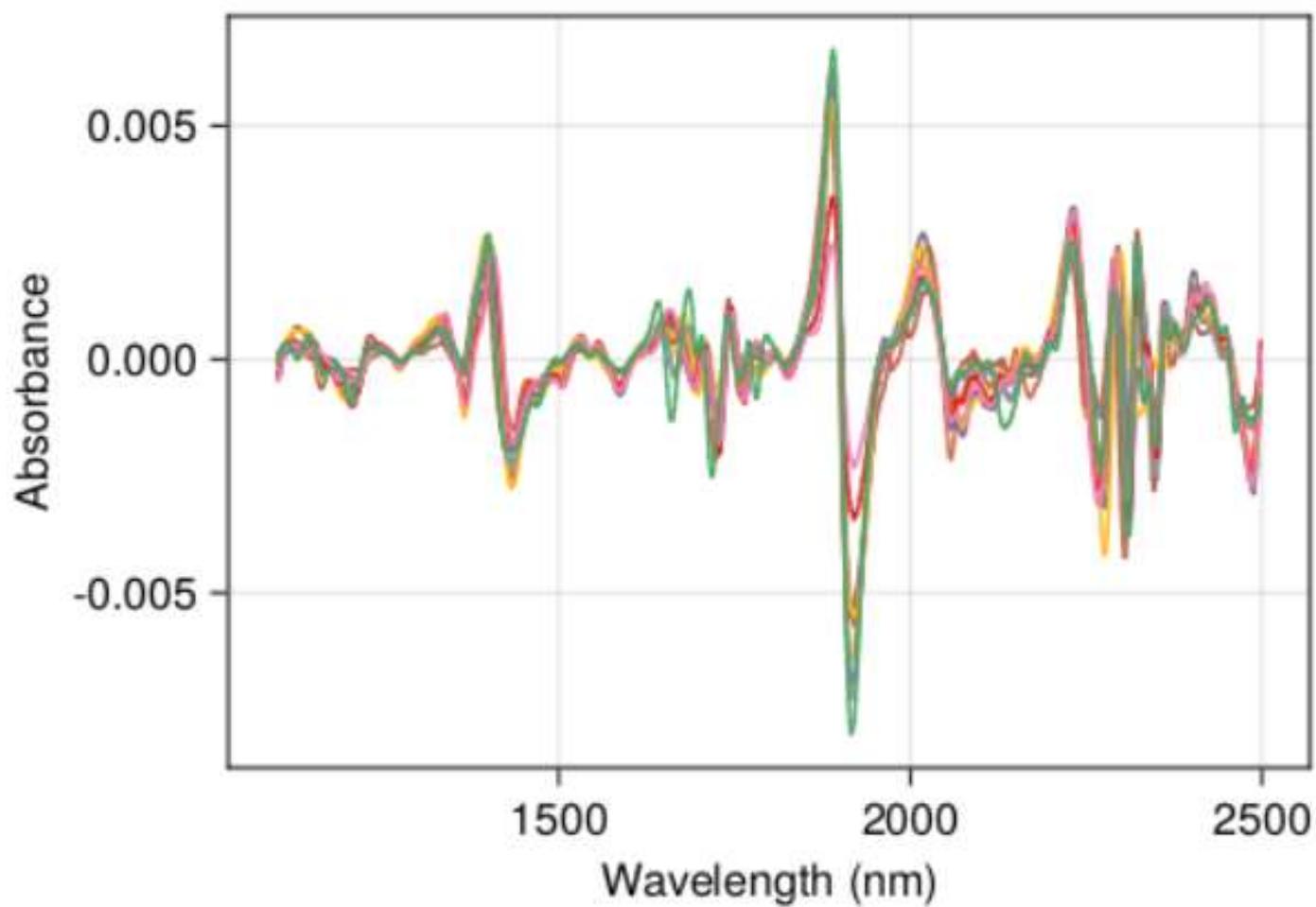
Row	dm	ndf	typ	test
	Float64?	Float64?	String	Int64
1	92.23	37.58	Legume forages	1
2	93.26	49.6462	Legume forages	0
3	92.9	63.2939	Forage trees	0
4	94.44	51.6413	Cereal and grass forages	0
5	93.16	46.5114	Legume forages	0
6	93.44	40.8176	Legume forages	0
:	:	:	:	:
481	93.7303	55.5358	Legume forages	1
482	93.2967	68.9812	Legume forages	0
483	93.1954	39.1038	Legume forages	0
484	93.17	48.8868	Legume forages	0
485	93.2842	34.6457	Legume forages	1
474 rows omitted				

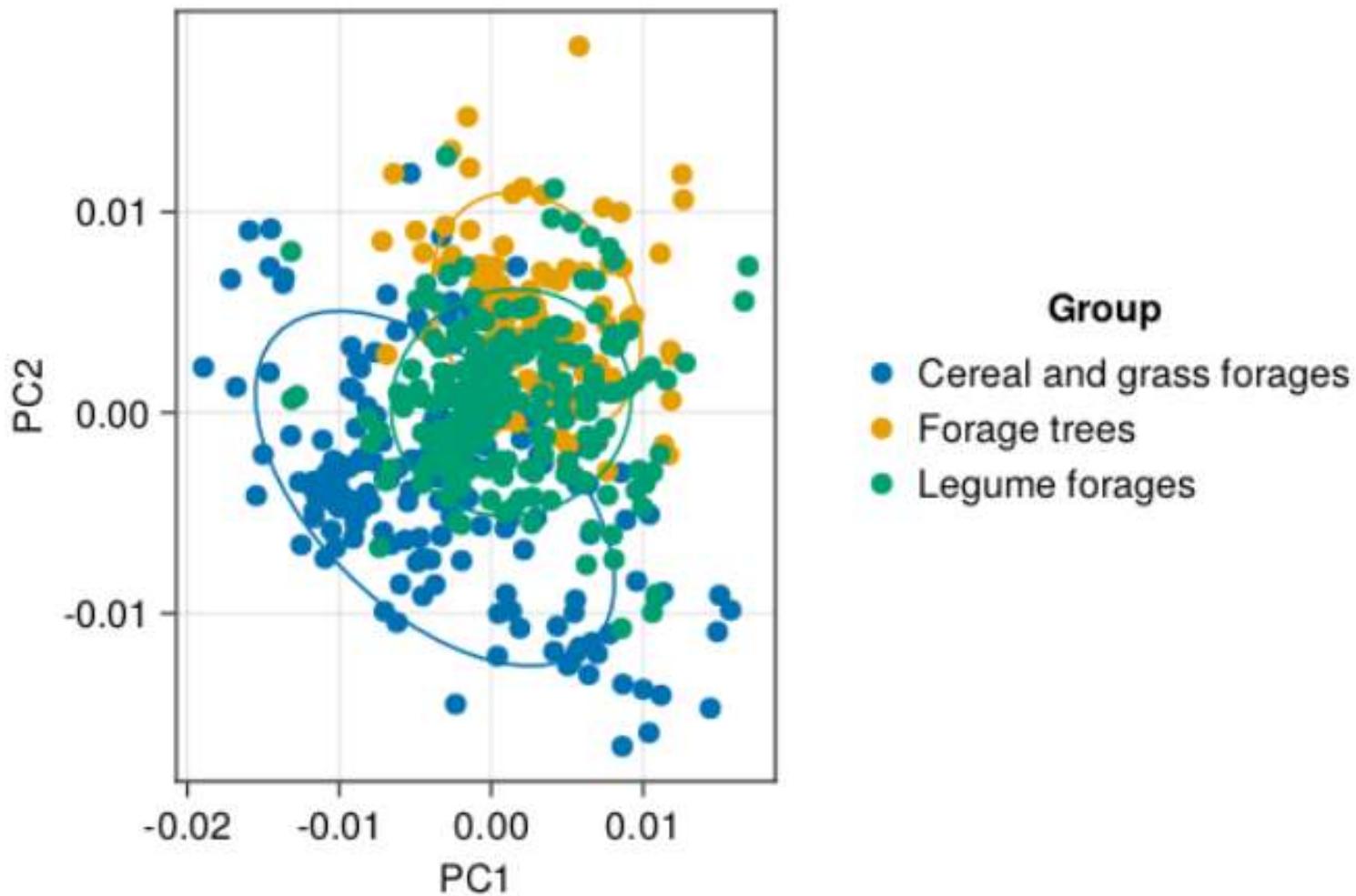
```
julia> X = dat.X
```

```
485x700 DataFrame
```

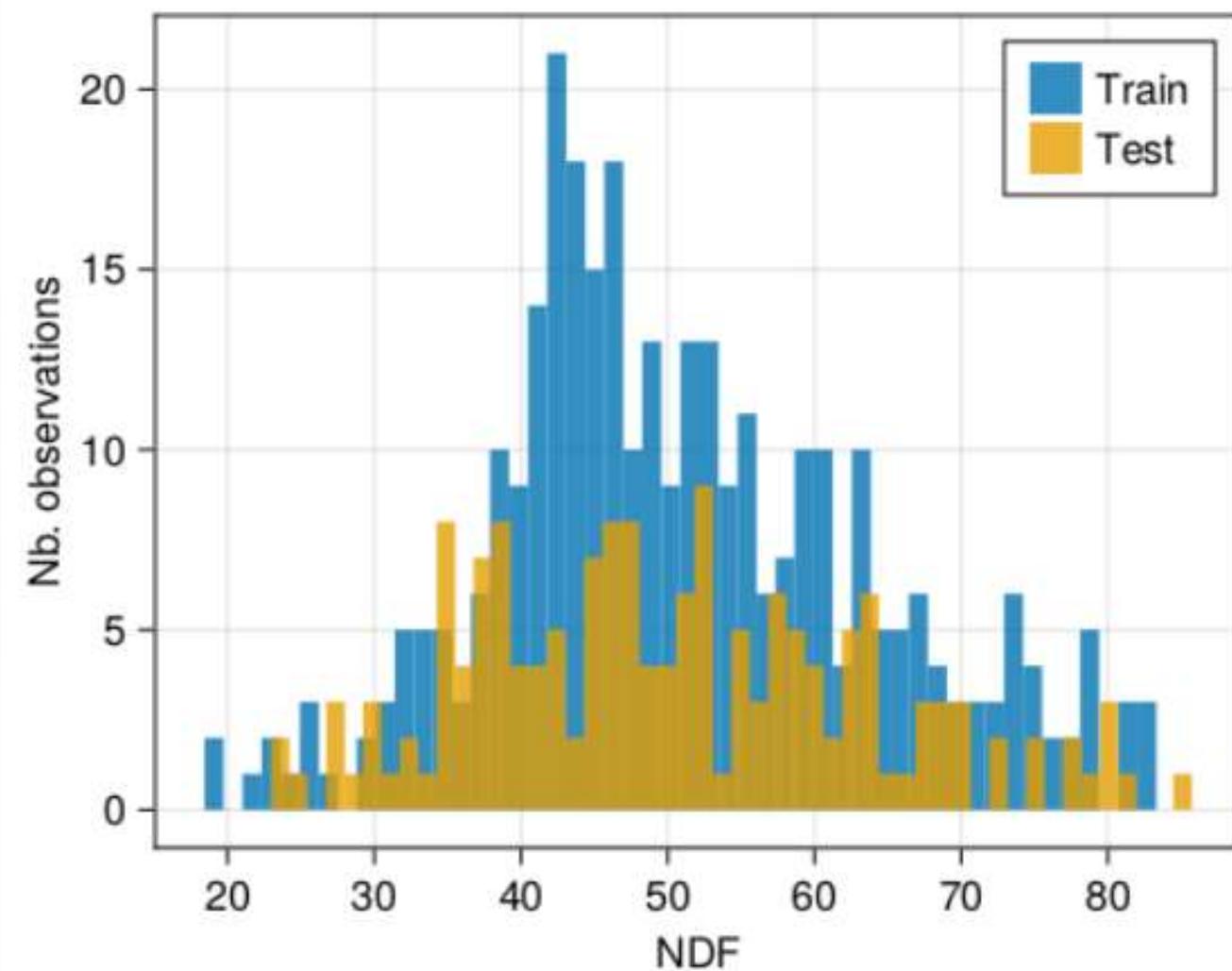
Row	1100	1102	1104	1106
	Float64	Float64	Float64	Float64
1	-0.000231591	-0.000175945	-8.48176e-5	2.05217e-5
2	-9.66352e-5	-3.30928e-5	5.64966e-5	0.000154135
3	-0.000131769	-7.8398e-5	7.92223e-7	8.90044e-5
4	-5.27109e-5	-1.38838e-5	3.73562e-5	8.87785e-5
5	-4.13284e-5	1.46607e-5	8.7976e-5	0.000161825
6	-0.000562333	-0.000451182	-0.000256137	-1.91839e-5
:	:	:	:	:
481	-0.000790809	-0.000707089	-0.000525597	-0.000272477
482	-0.000866662	-0.000767647	-0.000559022	-0.000274144
483	-0.000788774	-0.000703783	-0.000521761	-0.000270395
484	-0.00033791	-0.00028799	-0.000190344	-5.73166e-5
485	-0.000790261	-0.000703107	-0.000519605	-0.000268478

SNV + SAV-GOL d=2





## Predict NDF



## Script files

forages2\_descri.jl

gridscore\_lwplsrforges2.jl

gridcv\_lwplsrforges2.jl