

Exploratory Analysis on Big Data using the MEDA Toolbox Present and Future

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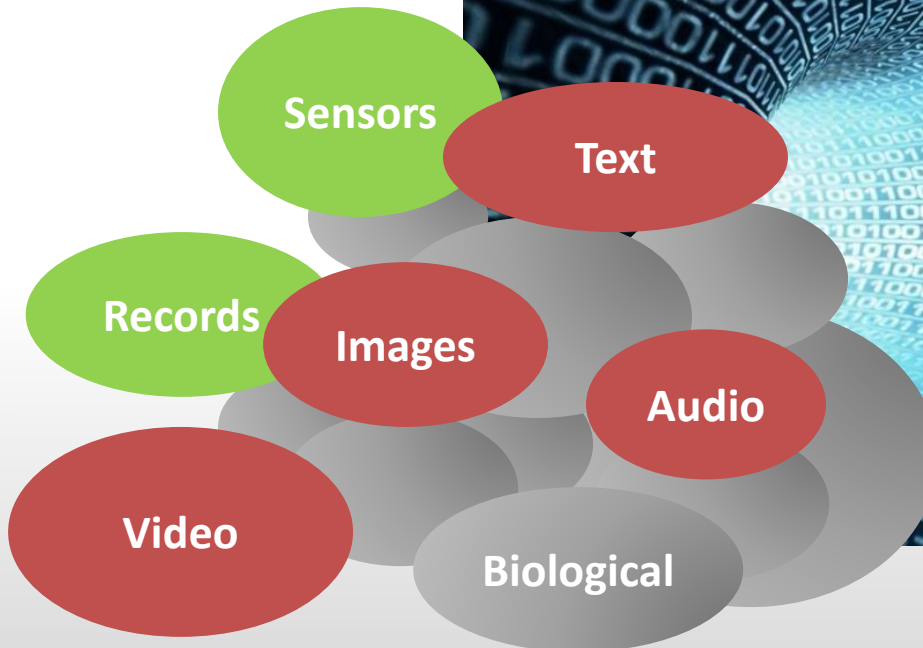


Network Engineering & Security Group
<http://nesg.ugr.es>

Velocity



Big Data



Variety

VOLUME

 Data-Driven Documents



Visual Analytics

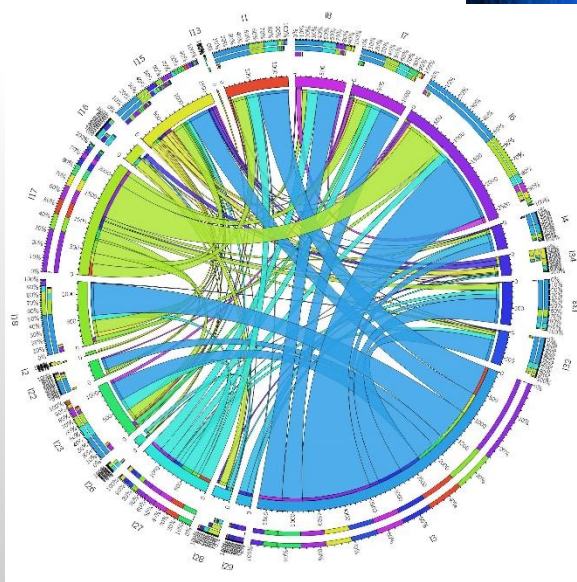
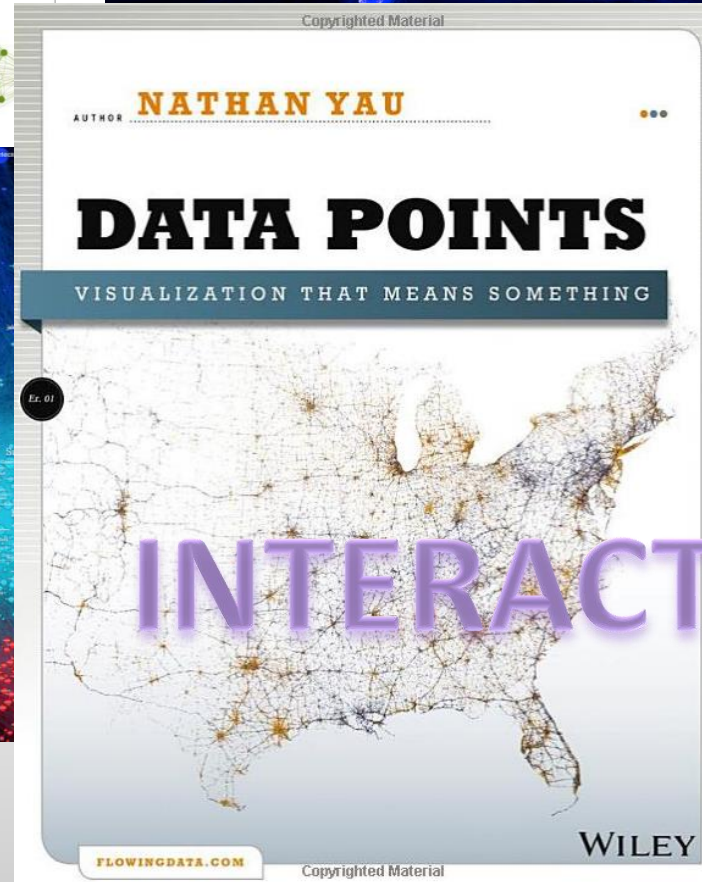
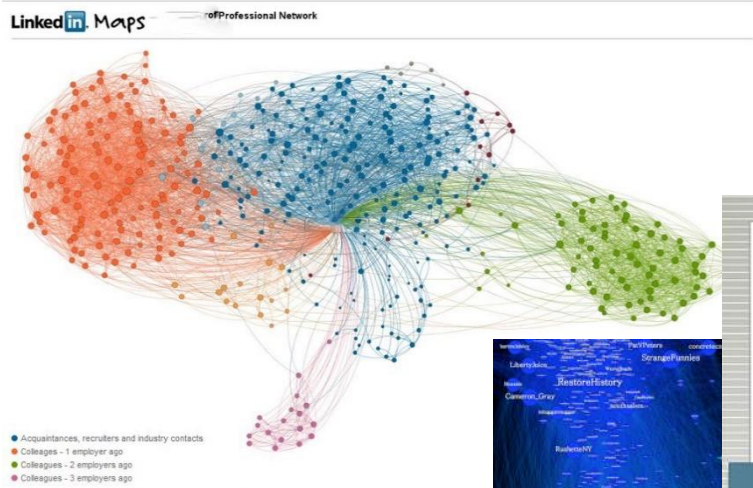
BIG DATA ECOSYSTEM

Storage

Parallel Processing

Cluster of Computers

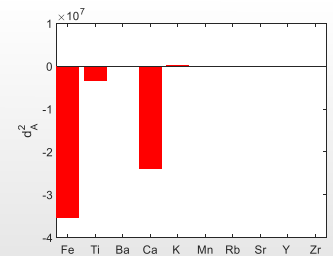
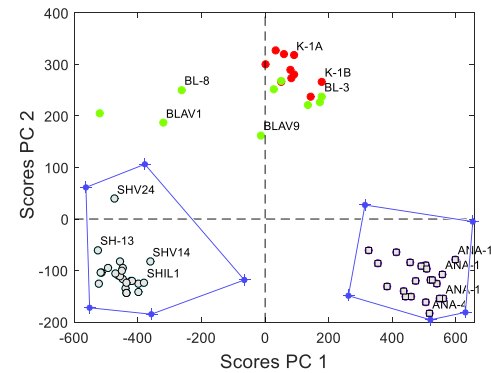




MEDA Toolbox

<https://github.com/josecamachop/MEDA-Toolbox>

- ✓ Models: PCA, PLS-DA, SPLS, GPCA, GPLS
- ✓ Dimensionality:
 - Scree plots
 - CV & D-CV
 - SVI Plots
- ✓ Structure in Variables:
 - Loading plots
 - MEDA plots
- ✓ Distribution of Observations
 - Score plots
 - MSPC: D-st, Q-st
 - Covariance MSPC: ADICOV
- ✓ Observations vs Variables
 - oMEDA plots
- ✓ Data simulation
 - [simuleMV](#)



ChemoLab, (2015) 143: 49

<https://github.com/josecamachop/MEDA-Toolbox>

josecamachop / MEDA-Toolbox Unwatch 9 Star 5 Fork 6

Code Issues 22 Pull requests 0 Projects 0 Wiki Pulse Graphs Settings

Multivariate Exploratory Data Analysis Toolbox for Matlab — Edit

251 commits 6 branches 2 releases 4 contributors GPL-3.0

Branch: master New pull request Create new file Upload files Find file Clone or download

Commit	Message	Time
josecamachop	Minor correction in gia when returning 0 states	Latest commit 5206a94 16 days ago
BigData	Updating copyright	a month ago
Examples	Addition of double crossvalidation in PLS	4 months ago
GUI	Updating copyright	a month ago
Technical Reports	Big Data functionality added	2 years ago
.gitattributes	Added .gitattributes & .gitignore files	2 years ago
.gitignore	More on new version	8 months ago
ADICOV.m	Updating copyright	a month ago
ADindex.m	Updating copyright	a month ago
ADindex2.m	Updating copyright	a month ago
GUIDELINES.txt	Adding SPLS, CV and DCV	a month ago
LICENSE.txt	first commit	2 years ago
README.txt	Adding SPLS, CV and DCV	a month ago

➔ Extensions for Big Data

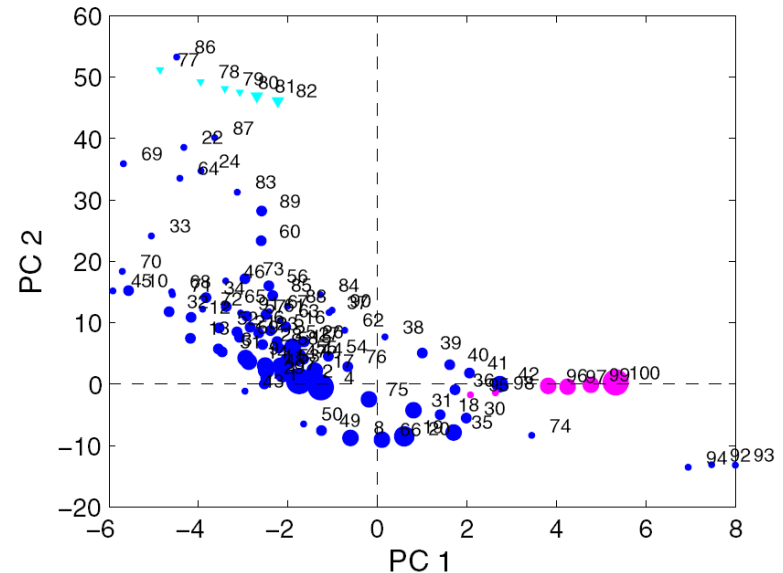
- ✓ For variables ➔ (linear) kernel calibration with EWMA update

$$(X'X)_t = \lambda \cdot (X'X)_{t-1} + \tilde{X}_t' \cdot \tilde{X}_t$$

- Scalable to any size
- PCA/PLS, MEDA, oMEDA
- GPCA, GPLS
- ADICOV MSPC

- ✓ For observations ➔ Clustering

- Scalable to any size
- Compressed Score Plots
- Compressed MSPC



ChemoLab, (2014) 135: 110

```

clear
load kdd

Lmodel = Lmodel_ini; % Initialization
Lmodel.update = 2; % Change this to 1 for EWMA and 2 for Iterative
Lmodel.type = 2; % Change this to 1 for PCA and 2 for PLS
Lmodel.lv = 3; % Initial number of LVs
Lmodel.prep = 2; % X-block prepr. 0: None, 1: Mean-center, 2: Auto-scaling
Lmodel.prepy = 2; % Y-block prepr. 0: None, 1: Mean-center, 2: Auto-scaling
Lmodel.nc = 100; % Number of clusters

lambda = 1-1e-4; % Forgetting factor in EWMA
step = 0.01;

%% Model building (EWMA or Iterative)

if Lmodel.update == 1
    Lmodel = update_ewma(short_list, '', Lmodel, lambda, step, 1); % EWMA
else
    Lmodel = update_iterative(short_list, '', Lmodel, 20, step, 0, '', 1); % Iterative
end

%% Data Analysis

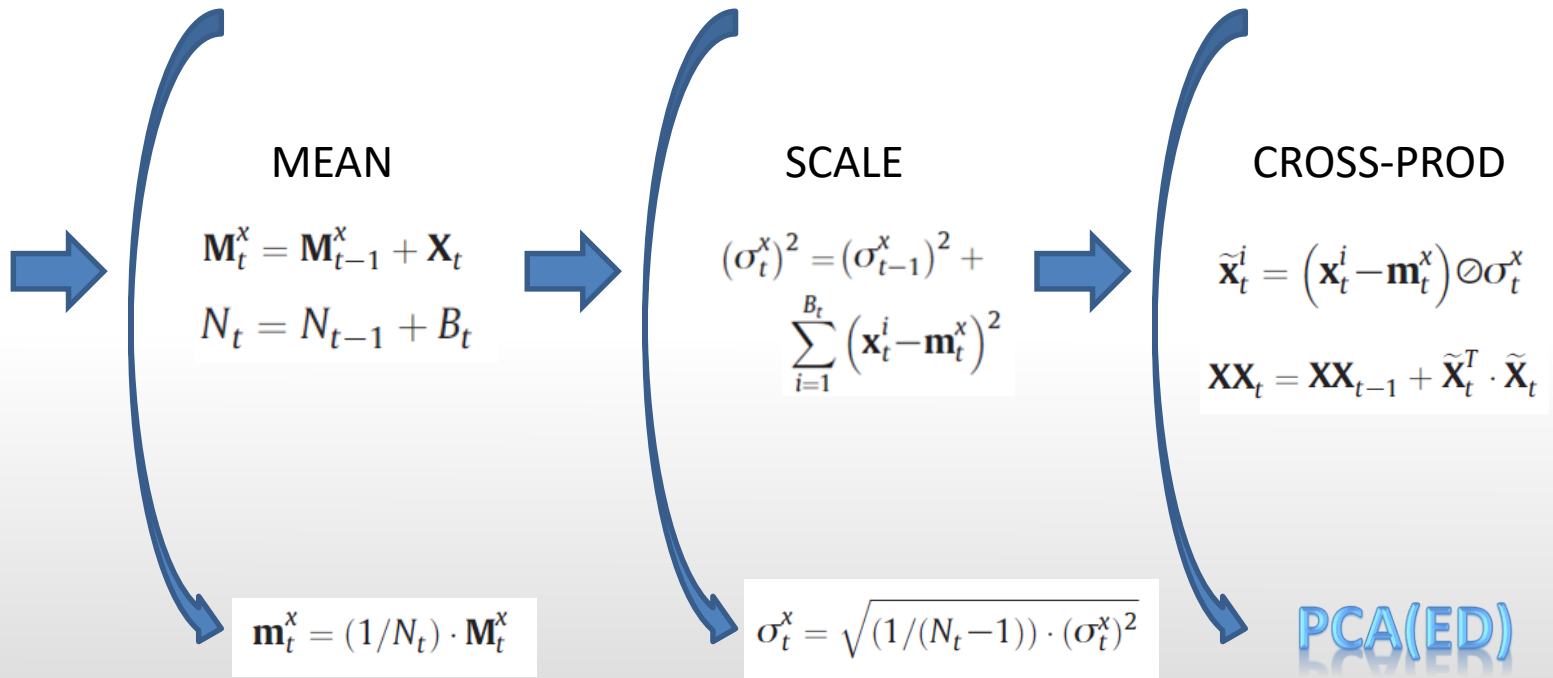
if Lmodel.type==2, % for PLS

    % Score plot
    scores_Lpls(Lmodel, 1:2);

    % MEDA
    map = meda_Lpls(Lmodel, 1:2, 0, 3);
    
```


BIG DATA SETS

```
lmodel = update_iterative(short_list, '', lmodel, 20, step, 0, '', 1); % Iterative
```



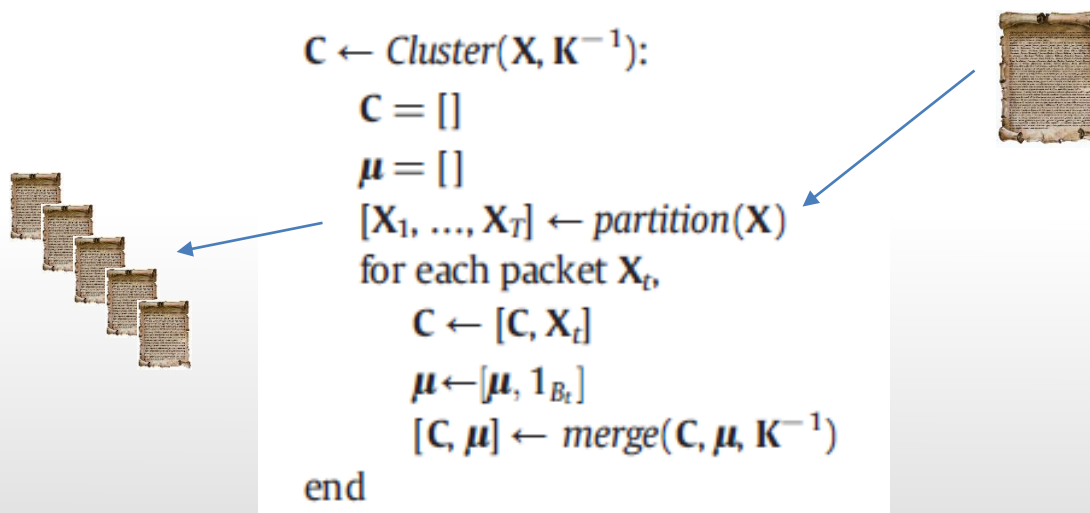
ChemoLab, (2014) 135:110

MEDA, Loading plots,
CV, ...

BIG DATA SETS

```
Lmodel = update_iterative(short_list, '', Lmodel, 20, step, 0, '', 1); % Iterative
```

Compressed Scores



ChemoLab, (2014) 135:110

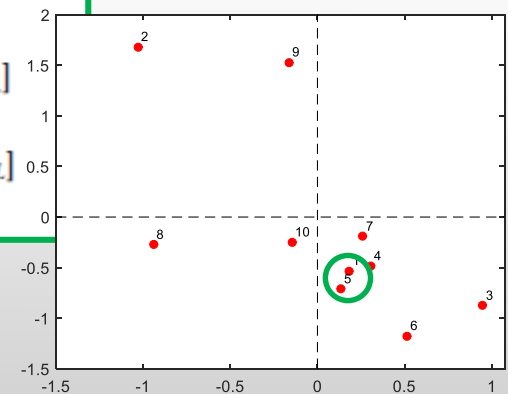
BIG DATA SETS

```
lmodel = update_iterative(short_list, '', lmodel, 20, step, 0, '', 1); % Iterative
```

Compressed Scores

```
C ← Cluster(X, K-1):
  C = []
  μ = []
  [X1, ..., XT] ← partition(X)
  for each packet Xt,
    C ← [C, Xt]
    μ ← [μ, 1|Xt|]
    [C, μ] ← merge(C, μ, K-1)
  end
```

```
[C, μ] ← merge(C, μ, K-1):
  L := # (C)
  C := [c1, ..., cL]
  μ := [μ1, ..., μL]
  while (# (C) > Lend),
    [ci, cj] ← min_dist(C, K-1)
    ci ← centroid(μi · ci, μj · cj)
    C ← [c1, ..., cj-1, cj+1, ..., cL]
    μi ← μi + μj
    μ ← [μ1, ..., μj-1, μj+1, ..., μL]
  end
```



ChemoLab, (2014) 135:110

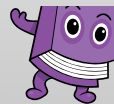
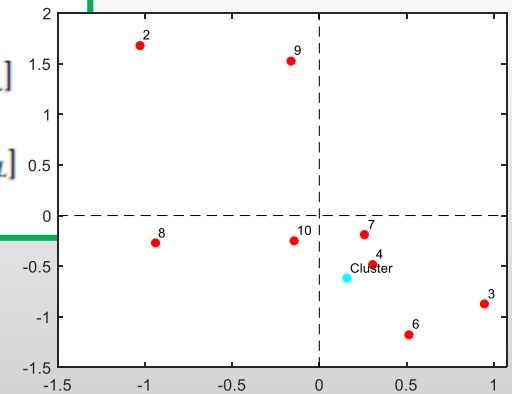
BIG DATA SETS

```
Lmodel = update_iterative(short_list, '', Lmodel, 20, step, 0, '', 1); % Iterative
```

Compressed Scores

```
C ← Cluster(X, K-1):
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  [C, μ] ← merge(C, μ, K-1)
end
```

```
[C, μ] ← merge(C, μ, K-1):
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    μi ← μi + μj
    μ ← [μ1, ..., μj-1, μj+1, ..., μL]
  end
```



ChemoLab, (2014) 135:110

BIG DATA SETS

```
Lmodel = update_iterative(short_list, '', Lmodel, 20, step, 0, '', 1); % Iterative
```

Compressed Scores

```
C ← Cluster(X, K-1):
  C = []
  μ = []
  [X1, ..., XT] ← partition(X)
  for each packet Xt,
    C ← [C, Xt]
    μ ← [μ, 1pt]
  [C, μ] ← merge(C, μ, K-1)
end
```

```
[C, μ] ← merge(C, μ, K-1):
  L := # (C)
  C := [c1, ..., cL]
  μ := [μ1, ..., μL]
  while (# (C) > Lend),
    [ci, cj] ← min_dist(C, K-1)
    ci ← centroid(μi · ci, μj · cj)
    C ← [c1, ..., cj-1, cj+1, ..., cL]
    μi ← μi + μj
    μ ← [μ1, ..., μj-1, μj+1, ..., μL]
  end
```



$$d_K(x_i, x_j) = \|x_i - x_j\|_K = \left((x_i - x_j)^T K^{-1} (x_i - x_j) \right)^{1/2}$$

$$K^{-1} = P \cdot K_{PCA}^{-1} \cdot P^T$$

$$K^{-1} = R \cdot K_{PLS}^{-1} \cdot R^T$$

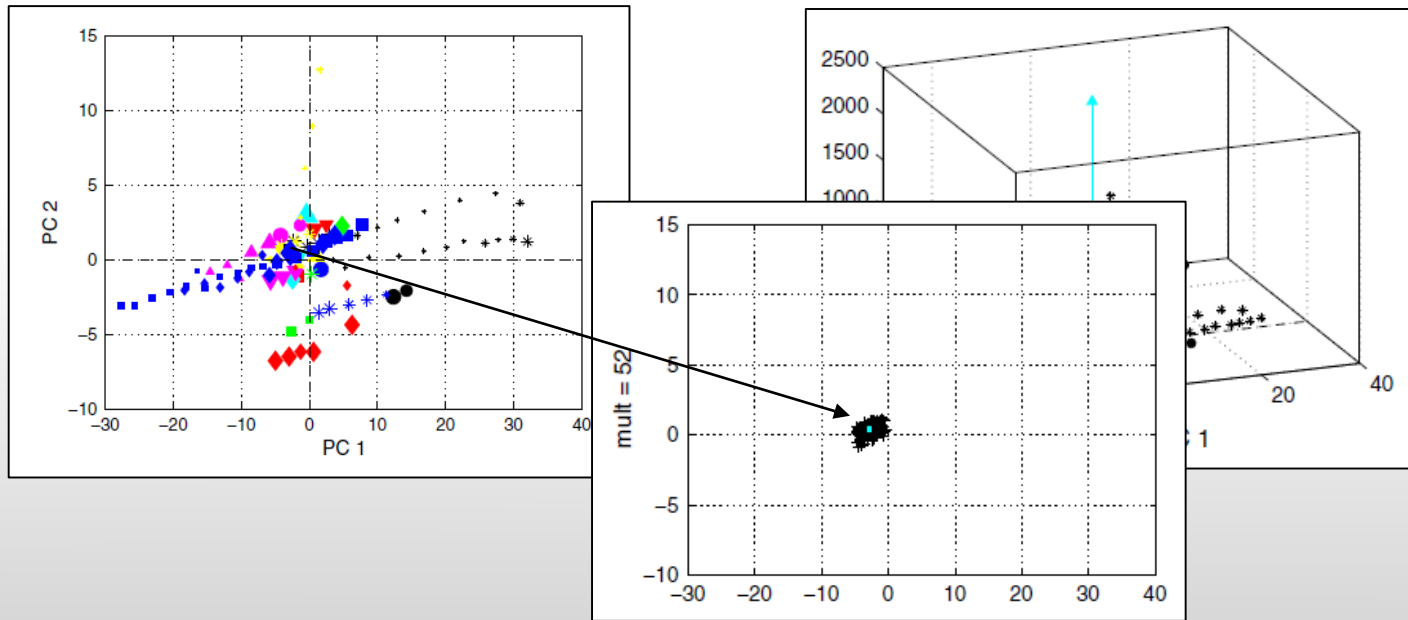


ChemoLab, (2014) 135:110

BIG DATA SETS

```
Lmodel = update_iterative(short_list, '', Lmodel, 20, step, 0, '', 1); % Iterative
```

Compressed Score Plot (CSP)

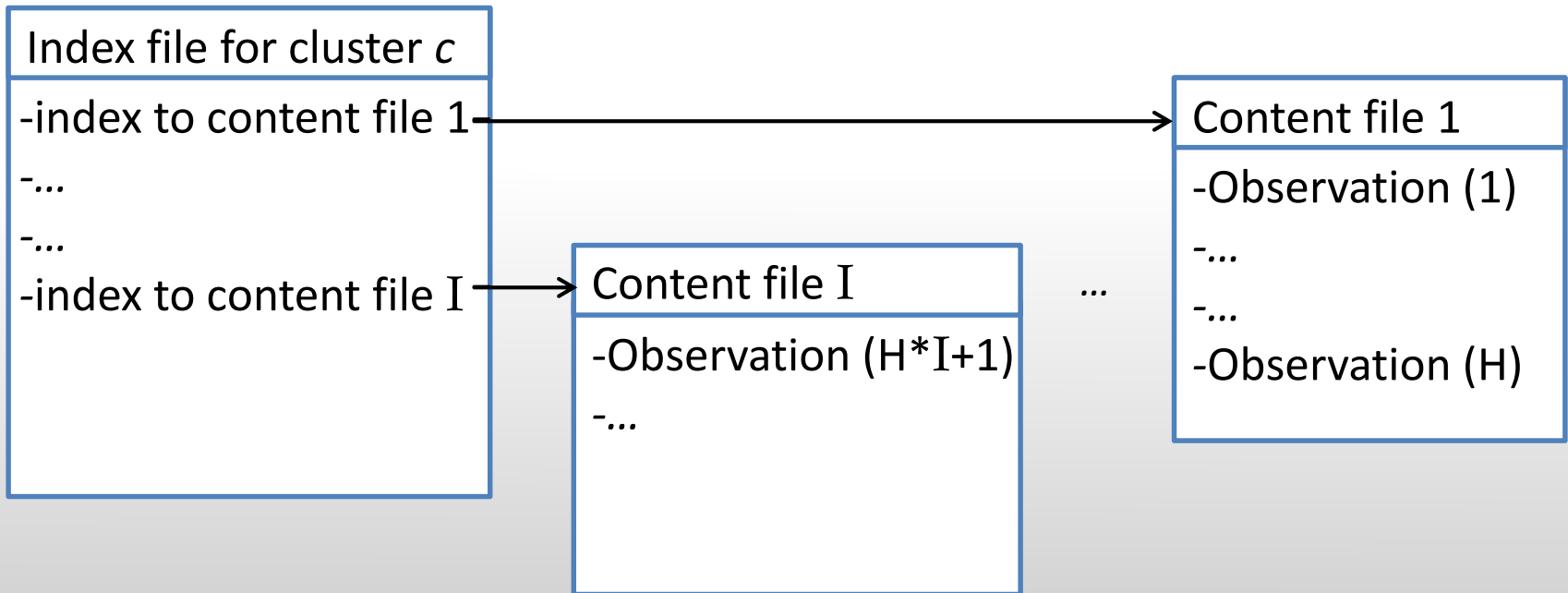


ChemoLab, (2014) 135:110

BIG DATA SETS

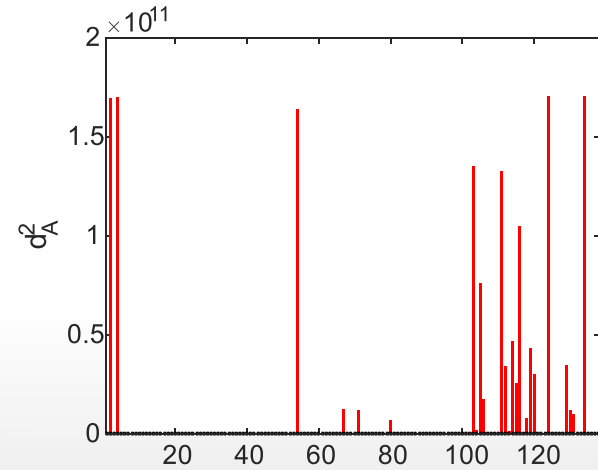
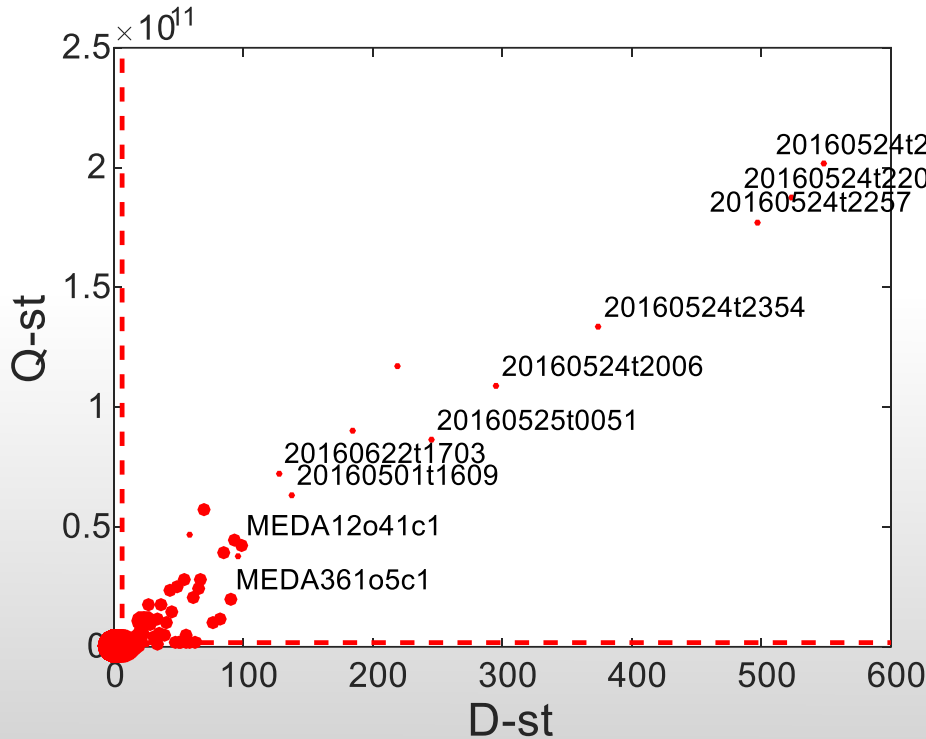
```
Lmodel = update_iterative(short_list, '', Lmodel, 20, step, 1, './output/', 1); % Iterative
```

Compressed Score Plot (CSP)



BIG DATA SETS

Example: vw PCA-MSPC in Big Data (Networkmetrics)



SPAM ATTACK

Data Streams

```
Lmodel = update_ewma(short_list, '', Lmodel, lambda, step, 1); % EWMA
```



$$\mathbf{M}_t^x = \lambda \cdot \mathbf{M}_{t-1}^x + \mathbf{X}_t$$

$$\mathbf{m}_t^x = (1/N_t) \cdot \mathbf{M}_t^x$$

$$N_t = \lambda \cdot N_{t-1} + B_t$$

$$(\sigma_t^x)^2 = \lambda \cdot (\sigma_{t-1}^x)^2 + \sum_{i=1}^{B_t} (\mathbf{x}_t^i - \mathbf{m}_t^x)^2$$

$$\sigma_t^x = \sqrt{(1/(N_t-1)) \cdot (\sigma_t^x)^2}$$



$$\tilde{\mathbf{x}}_t^i = (\mathbf{x}_t^i - \mathbf{m}_t^x) \oslash \sigma_t^x$$

$$\mathbf{XX}_t = \lambda \cdot \mathbf{XX}_{t-1} + \tilde{\mathbf{X}}_t^T \cdot \tilde{\mathbf{X}}_t$$

PCA(ED)

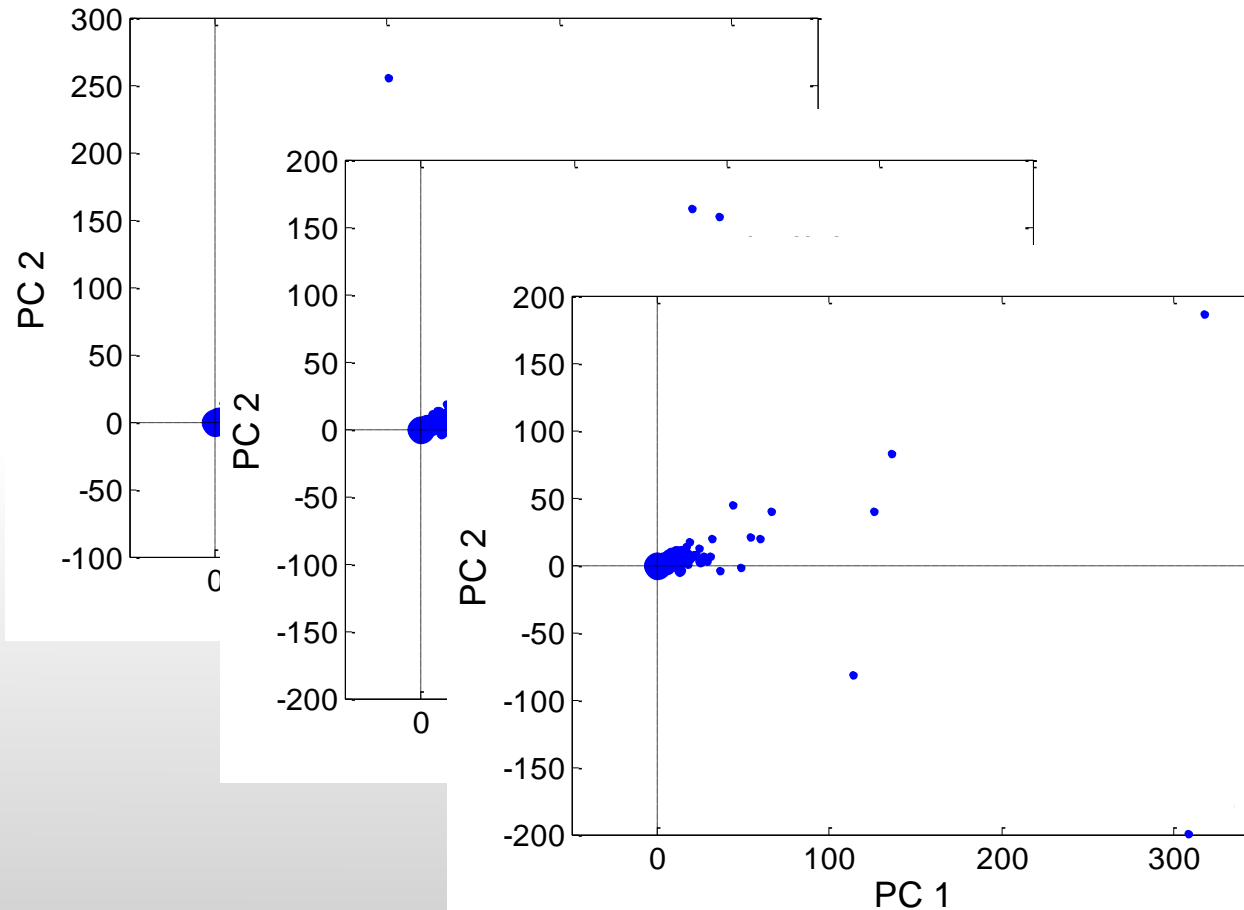
PLS (XX,XY)

MEDA,
Loading plots,
CV, ...



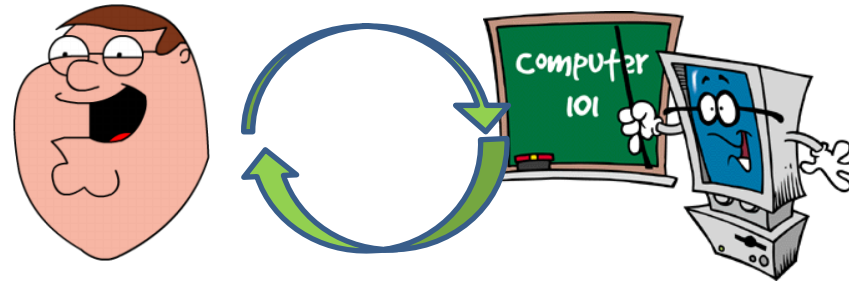
J.P.C., (1997) 7:169
ChemoLab, (2014) 135:110

Data Streams



$$Lmodel \text{ at } t: (X'X)_t = 0.9 \cdot (X'X)_{t-1} + \tilde{X}_t' \cdot \tilde{X}_t$$

➔ Data mining / Machine learning



➔ Chemometrics / Exploratory Data Analysis



➔ EDA + Visual Analytics



iMEDA Dashboard 1.1

Interactive visualization for EDA using the MEDA-Toolbox.

Please select the initial number of principal components:

Min 2 - Max 30.

Please select the pre-processing method from the following:

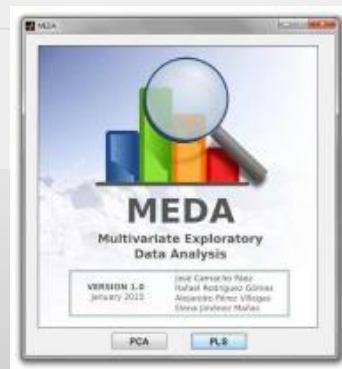
- No pre-processing
- Mean centering
- Auto-scaled

Please select the dataset from the following:

- Selwood
- Spectral

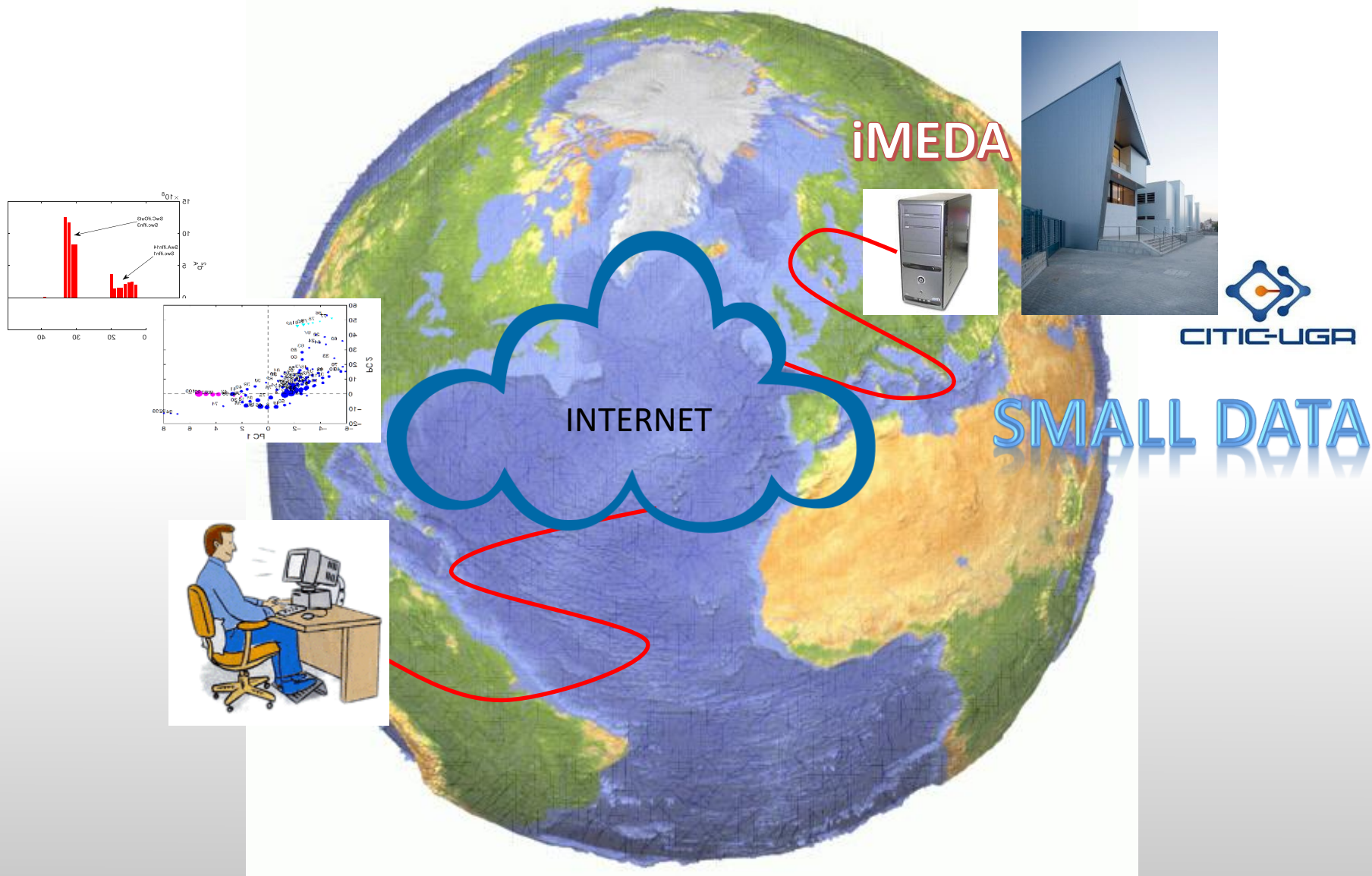
Submit

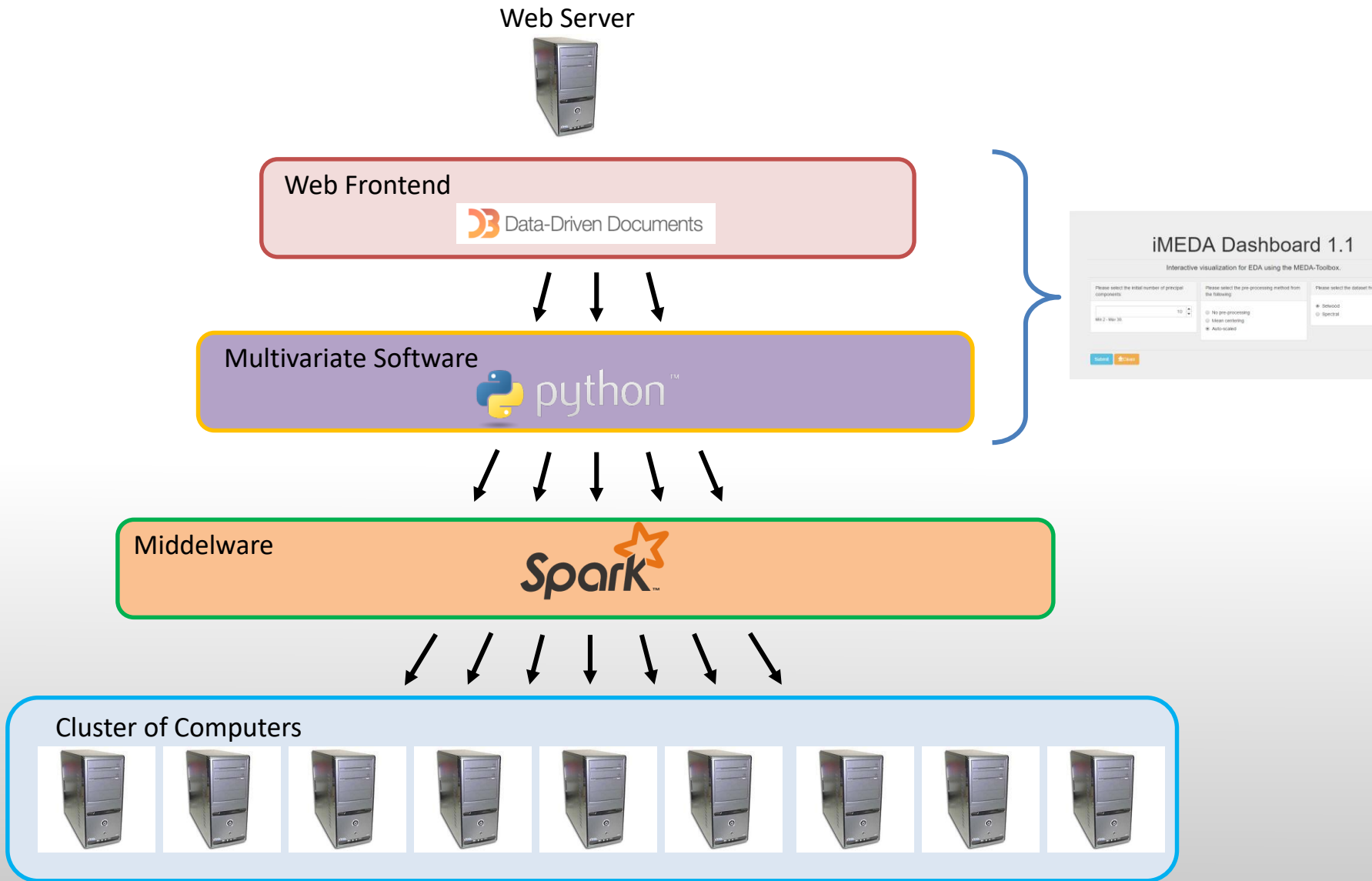
Clean

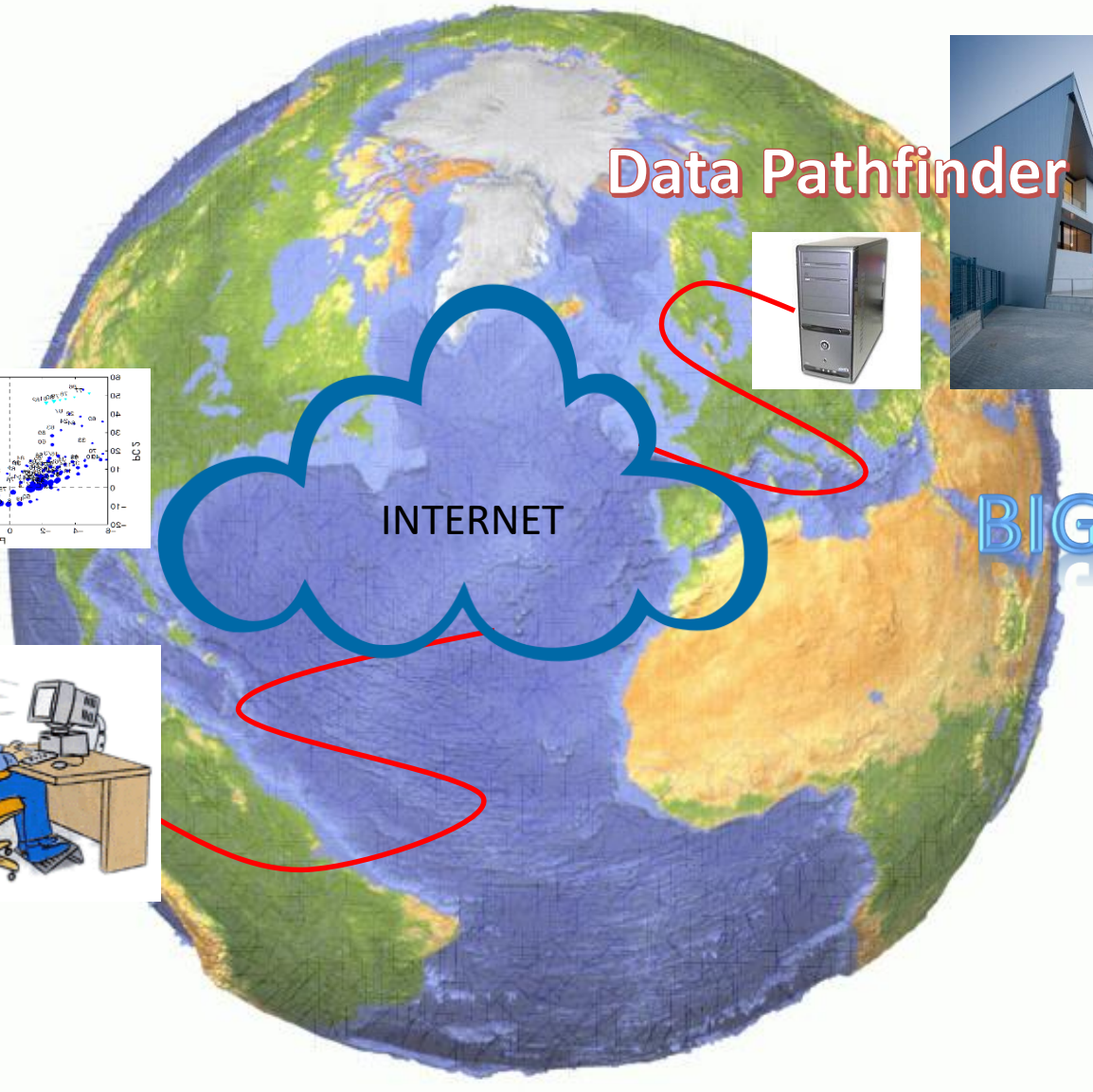
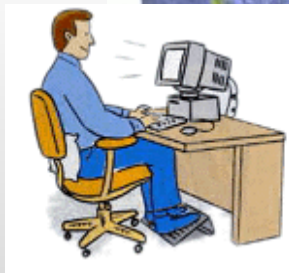
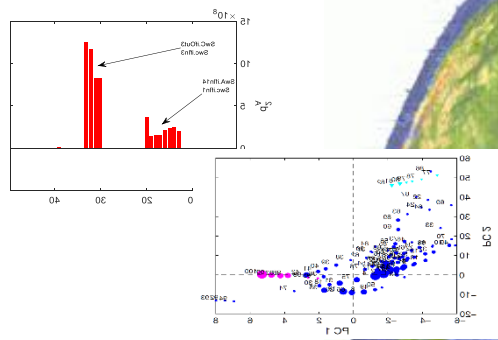


OCTAVE









Data Pathfinder



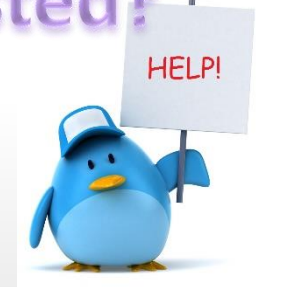
CITIC-UGR

BIG DATA



Is this the way
for MA + BD?

Interested?



IS IT TO ACHIEVE
FULL DATA INTERACTIVITY?

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