

Selected Parallel and Scalable Methods for Scientific Big Data Analytics



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Research Group Leader, Juelich Supercomputing Centre

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*ZIH Kolloquium , 21th May 2015
Technical University of Dresden*



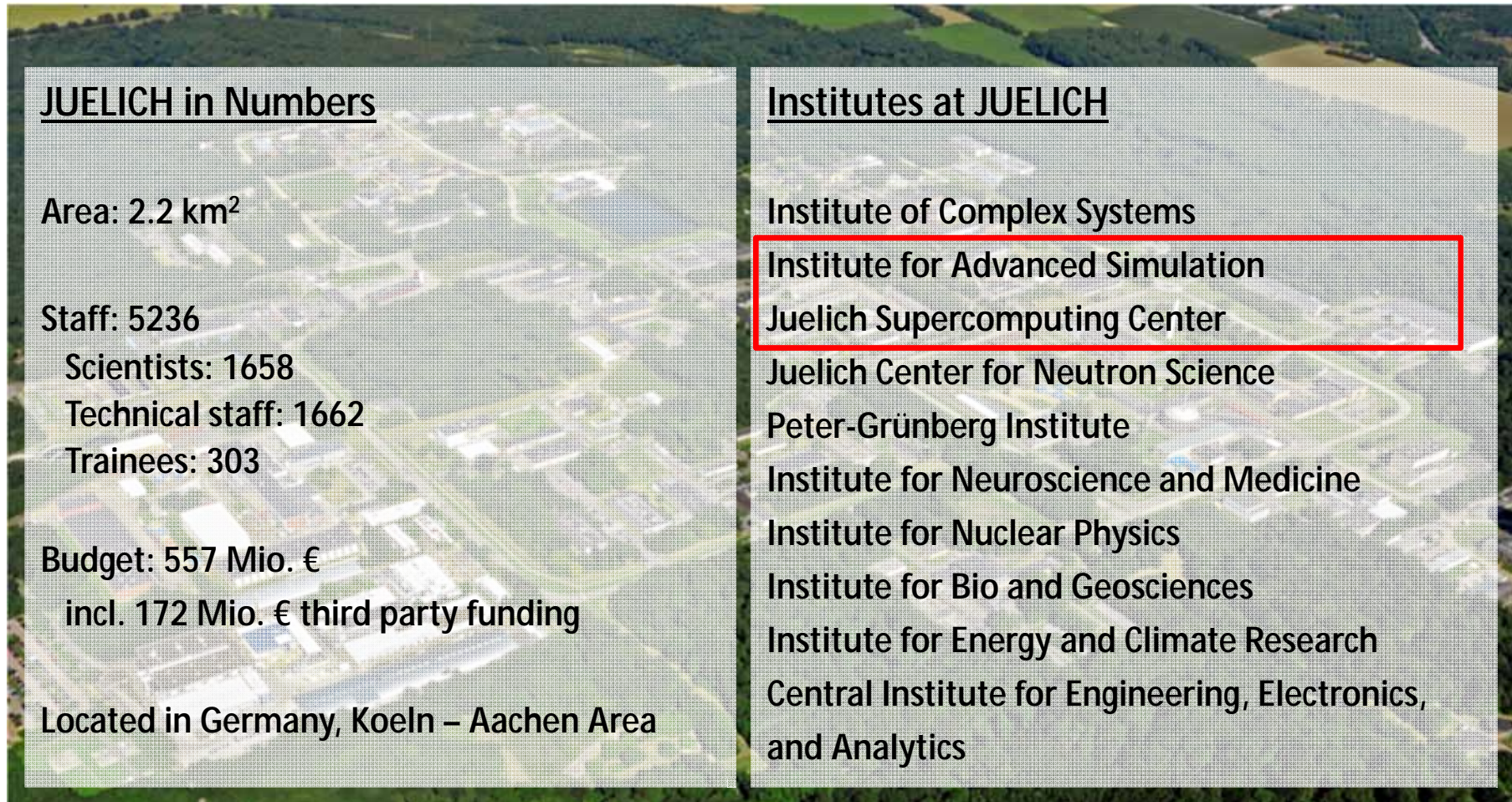
Federated Systems and Data Division

Research Group

High Productivity Data Processing



Research Centre Juelich



JUELICH in Numbers

Area: 2.2 km²

Staff: 5236

 Scientists: 1658

 Technical staff: 1662

 Trainees: 303

Budget: 557 Mio. €

 incl. 172 Mio. € third party funding

Located in Germany, Koeln – Aachen Area

Institutes at JUELICH

Institute of Complex Systems

Institute for Advanced Simulation

Juelich Supercomputing Center

Juelich Center for Neutron Science

Peter-Grünberg Institute

Institute for Neuroscience and Medicine

Institute for Nuclear Physics

Institute for Bio and Geosciences

Institute for Energy and Climate Research

Central Institute for Engineering, Electronics, and Analytics

Research for generic key technologies of the next generation

Scientific & Engineering Application-driven Problem Solving

University of Iceland

Schools of the University

School of Education

School of Humanities

School of Engineering and Natural Sciences

School of Social Sciences

School of Health Sciences

Interdisciplinary Studies

Full programmes taught in English

Staff: ~ 1259

Students: ~14.000

Located in Reykjavik Capital Center, Iceland

Faculties of the School

Civil and Environmental Engineering

Earth Sciences

Electrical and Computer Engineering

Industrial Engineering

Mechanical Engineering

Computer Science

Life and Environmental Sciences

Physical Sciences

Teaching of key technologies in engineering & sciences

University Courses: Statistical Data Mining & HPC-A/B

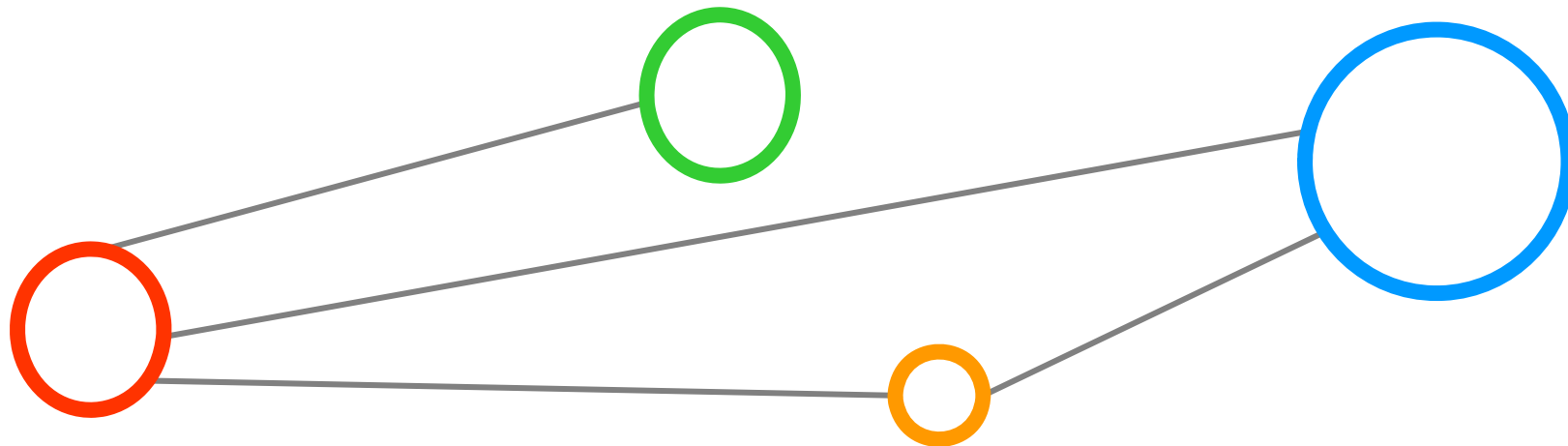


UNIVERSITY OF ICELAND

SCHOOL OF ENGINEERING AND NATURAL SCIENCES

FACULTY OF INDUSTRIAL ENGINEERING,
MECHANICAL ENGINEERING AND COMPUTER SCIENCE

Outline

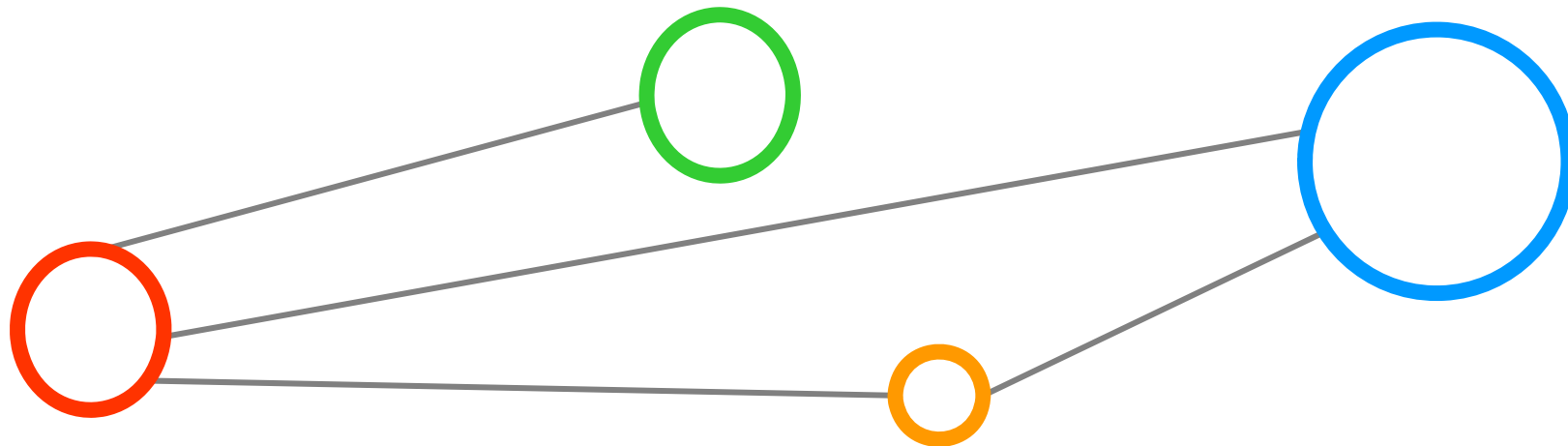


Outline

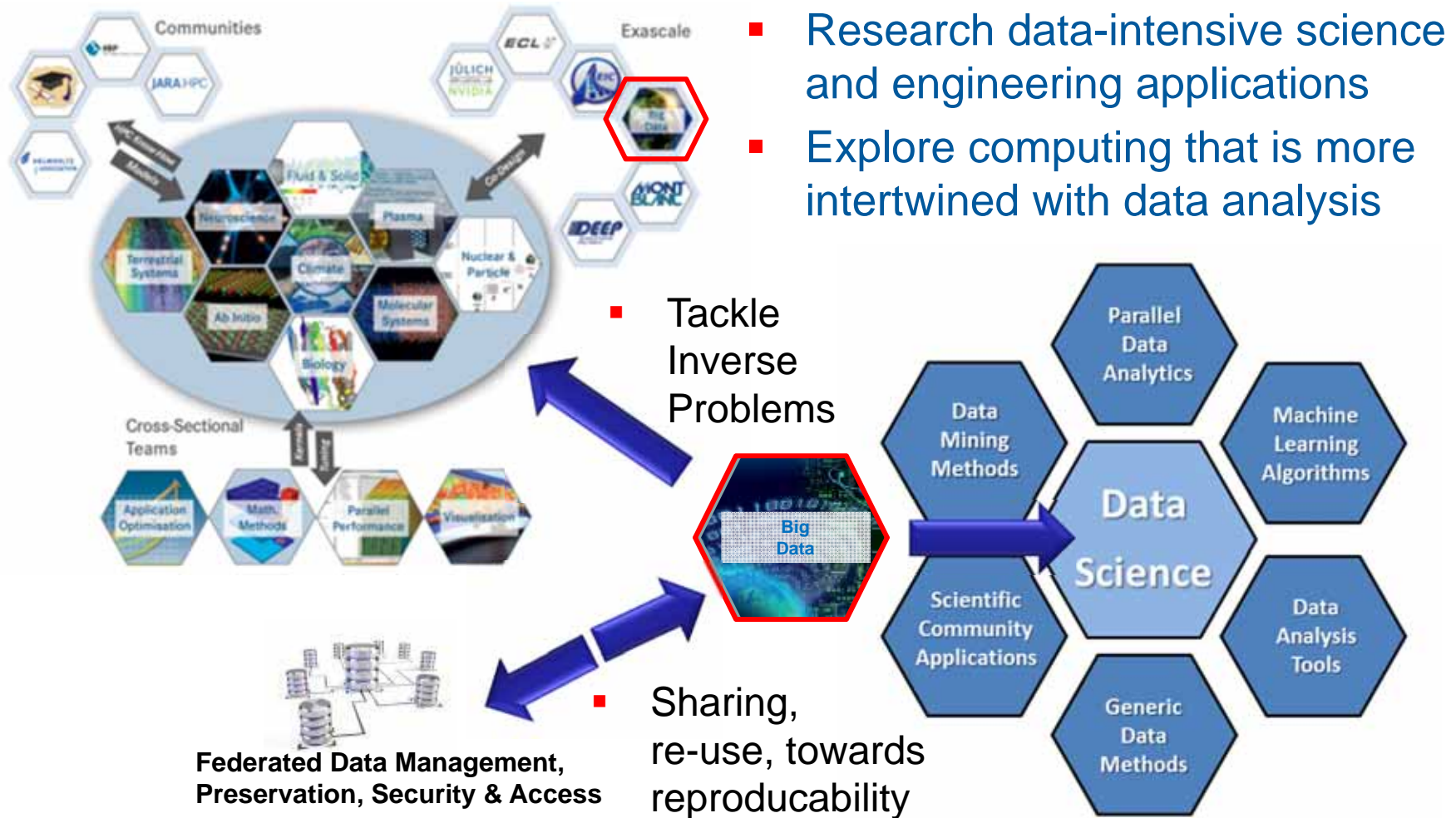
- **Data Analytics @ Juelich**
 - Driven by Scientific & Engineering Demands
 - Understanding of Terms & Key Focus
- **Scalable & Parallel Tools**
 - Clustering – DBSCAN
 - Classification – SVM
 - Scientific Applications in Context
- **Recent Research Directions**
 - ‘Brain Analytics’
 - Deep Learning
- **Conclusions**
- References & Backup Slides



Data Analytics @ Juelich



Data Analytics – Context JSC

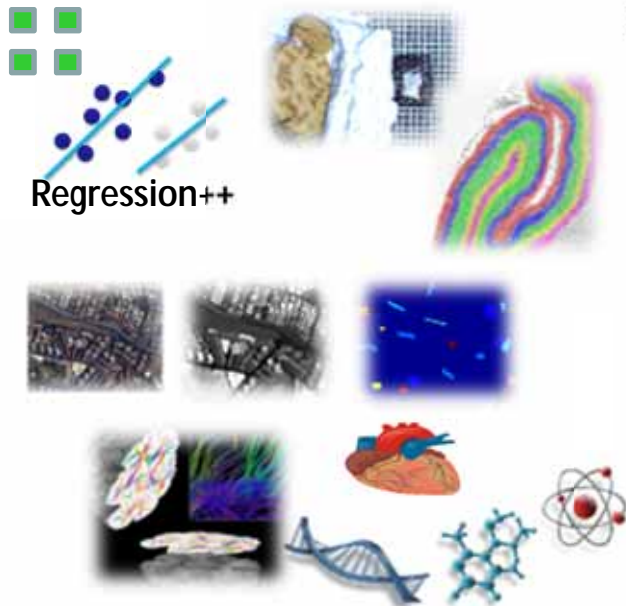


Data Analytics – Term Clarification

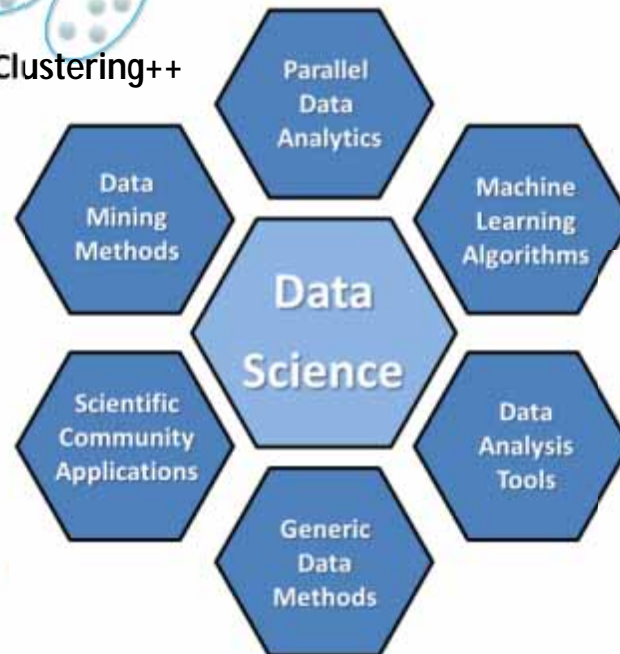
‘Data Analytics’ is an ‘interesting mix’ of different approaches

- Analytics: Whole methodology; Analysis: data investigation process itself
- ‘Big’ requires **scalable processing methods** and **underlying infrastructure**

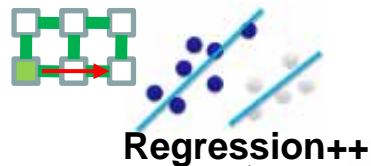
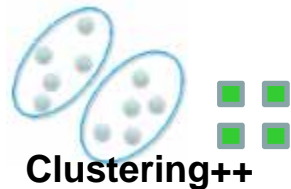
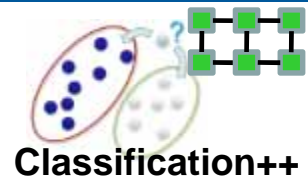
■ Concrete ‘big data’:
large medical data



■ Concrete ‘big data’:
large earth science data



Data Analytics – Research Key Focus

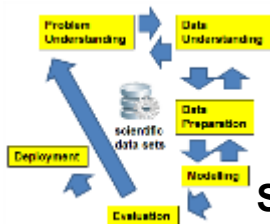


Scientific Computing

Scientific Applications using ‘Big Data’
 Traditional Scientific Computing Methods
 HPC and HTC Paradigms & Parallelization
 Emerging Data Analytics Approaches
 Optimized Data Access & Management
Statistical Data Mining & Machine Learning
 Inverse Problems & their Research Questions

‘Big Data’ Methods

Scientific Applications



Systematic & Automated Analytics guided by CRISP – DM

‘(Big) Data from various data science applications’



Data Analytics – Selected Research Group Activities

John von Neumann Institute for Computing (NIC)

- Peer-review of scientific big data analytics (SBDA) proposals
- Jointly work with SBDA users (first projects starting, prototyping process)

John von Neumann - Institut für Computing



Research Data Alliance (RDA)

- Chairing activities of the Big Data Analytics Interest Group
- Collaboration with a variety of EU and US partners
- Geoffrey Fox, UoIndiana (map-reduce), Kuo Kwo-Sen (NASA, SciDB)



Smart Data Innovation Lab (SDIL)

- Driving activities in the personalised medicine community (with Bayer)
- Collaboration with partners from industry (e.g. IBM, SAP, Siemens, etc.)

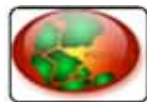


Data Analytics – Selected Research Expertise

Key expertise making algorithms parallel & scalable for ‘big data’

- **Driven by scientific and engineering cases**, e.g. understanding the human brain, remote sensing applications, marine measurements analysis, ...
- **Automate and/or support the data analysis process**
- **Example codes:** Density-based Spatial Clustering of Applications with Noise (DBSCAN), Support Vector Machines (SVMs),

Parallel & Scalable DBSCAN clustering tool



Problem: Automatic outlier detection for data quality

- ✓ Tailor solution for community
- ✓ Scalability towards Big Data
- ✓ Design and improve automatic data analytics approaches



[1] R. Huber, M. Riedel et al., 'Research data enters scholarly communication and big data analysis', EGU2014

Parallel & Scalable SVM classification tool

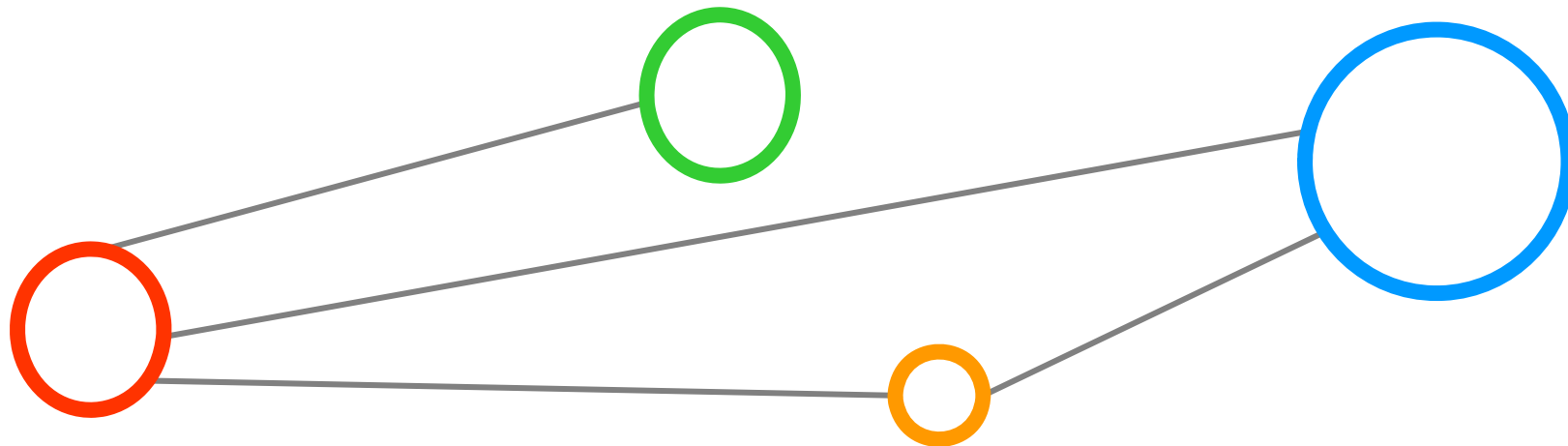
Problem: Classification of buildings from multi-spectral images

- ✓ Enable smooth transition from 'manual Matlab SVM scripts'
- ✓ Research on parallel SVM methods (map-reduce, HPC)



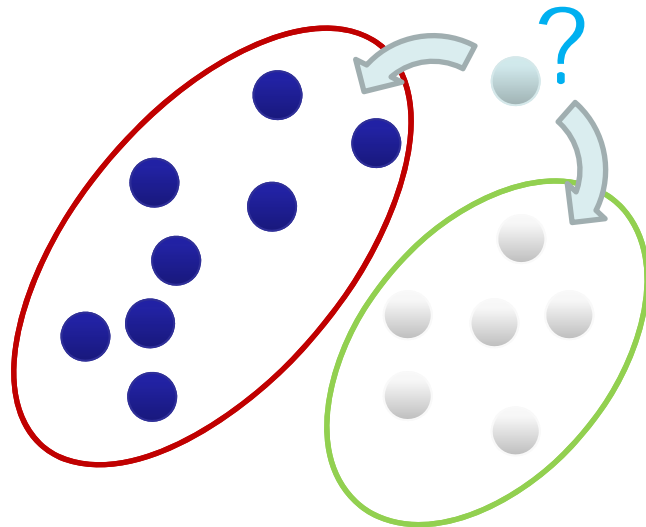
[2] G. Cavallaro, M. Riedel et al., 'Smart Data Analytics for Remote Sensing Applications', IGARSS2014

Scalable & Parallel Tools: Clustering



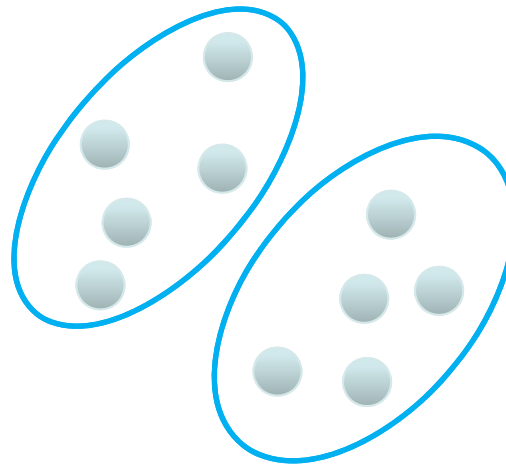
Learning From Data – Clustering Technique

Classification



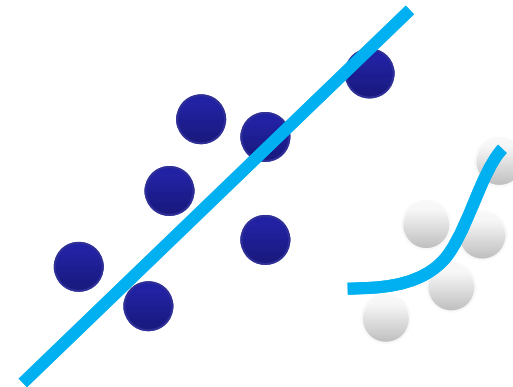
- Groups of data exist
- New data classified to existing groups

Clustering



- No groups of data exist
- Create groups from data close to each other

Regression



- Identify a line with a certain slope describing the data

Selected Clustering Methods

K-Means Clustering – Centroid based clustering

- Partitions a data set into K distinct clusters (centroids can be artificial)

K-Medoids Clustering – Centroid based clustering (variation)

- Partitions a data set into K distinct clusters (centroids are actual points)

Sequential Agglomerative hierarchic nonoverlapping (SAHN)

- Hierarchical Clustering (create tree-like data structure → 'dendrogram')

Clustering Using Representatives (CURE)

- Select representative points / cluster; as far from one another as possible

Density-based spatial clustering of applications + noise

(DBSCAN) Reasoning: density similarity measure helpful in our driving applications

- Assumes clusters of similar density or areas of higher density in dataset

Technology Review of Open & Available Tools

| Technology | Platform Approach | Analysis |
|--|-------------------|---|
| HPDBSCAN (authors implementation) | C; MPI; OpenMP | Parallel, hybrid, DBSCAN |
| Apache Mahout | Java; Hadoop | K-means variants, spectral, no DBSCAN |
| Apache Spark/MLlib | Java; Spark | Only k-means clustering, No DBSCAN |
| scikit-learn | Python | No parallelization strategy for DBSCAN |
| Northwestern University PDSDBSCAN-D | C++; MPI; OpenMP | Parallel DBSCAN |

*M. Goetz, M. Riedel et al., 6th Workshop on Data Mining in Earth System Science,
International Conference of Computational Science (ICCS), Reykjavik, to be published*

Parallel & Scalable DBSCAN MPI/OpenMP Tool (1)

DBSCAN Algorithm

- Introduced 1996 by Martin Ester et al. *[4] Ester et al.*
- Groups number of similar points into clusters of data
- Similarity is defined by a distance measure (e.g. *euclidean distance*)



Unclustered
Data

Distinct Algorithm Features

- Clusters a variable number of clusters
- Forms arbitrarily shaped clusters
- Identifies outliers/noise



Clustered
Data

Understanding Parameters for MPI/OpenMP tool

- Looks for a similar points within a given search radius
→ **Parameter *epsilon***
- A cluster consist of a given minimum number of points
→ **Parameter *minPoints***

[3] M.Goetz & C. Bodenstern, HPDBSCAN Tool

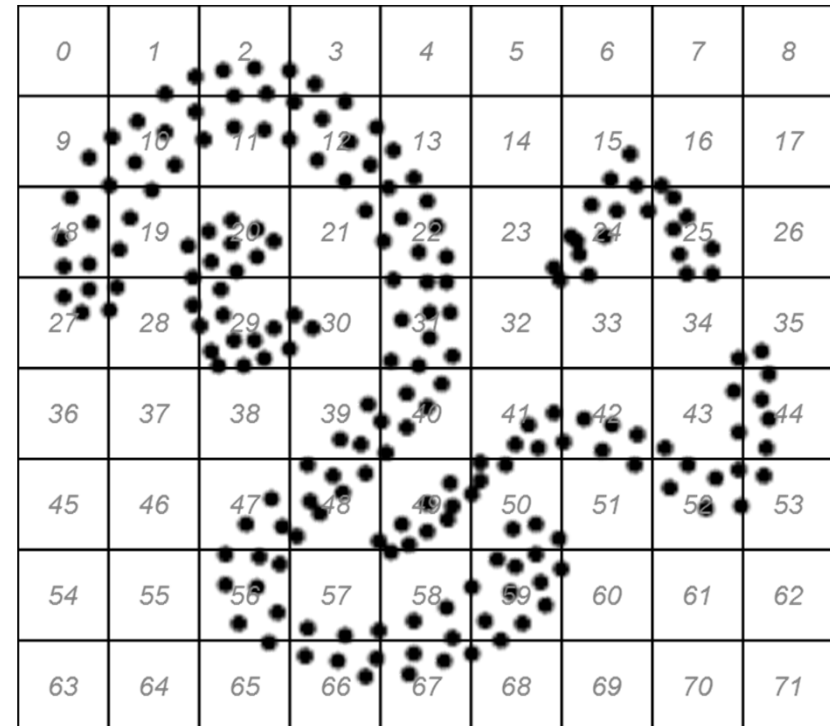
Parallel & Scalable DBSCAN MPI/OpenMP Tool (2)

Parallelization Strategy

- Smart 'Big Data' Preprocessing into Spatial Cells
- OpenMP standalone
- MPI (+ optional OpenMP hybrid)

Preprocessing Step

- Spatial indexing and redistribution according to the point localities
- Data density based chunking of computations



Computational Optimizations

- Caching of point neighborhood searches
- Cluster merging based on comparisons instead of zone reclustering

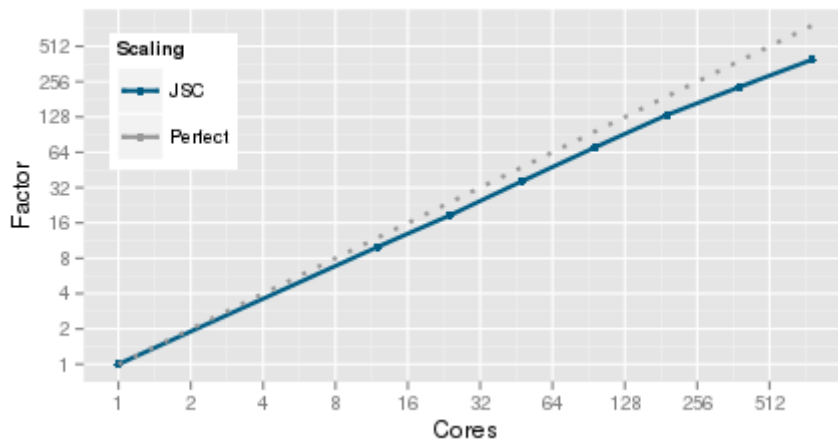
[3] M.Goetz & C. Bodenstein, HPDBSCAN Tool

Parallel & Scalable DBSCAN MPI/OpenMP Tool (3)

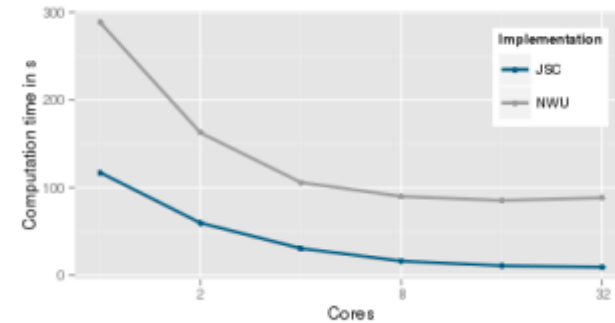
Performance Comparisons

- With another open-source parallel DBSCAN implementation (aka 'NWU') *[5] Patwary et al.*
- 3.7056.351 data points (2 dimensions)
- Use of Hierarchical Data Format (HDF) v.5 for scalable input/output of 'big data'

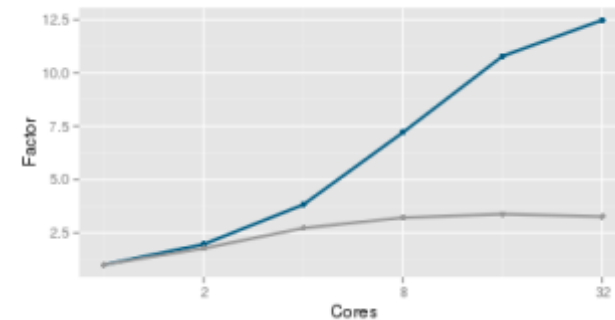
Speedup



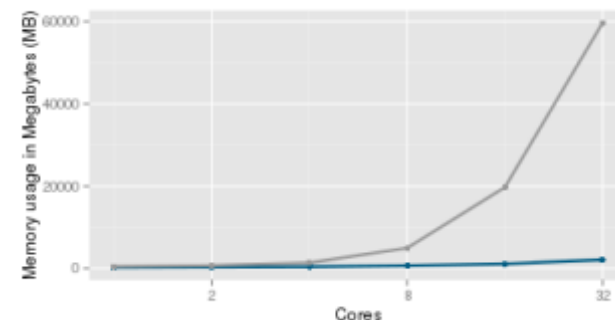
Computation time comparison



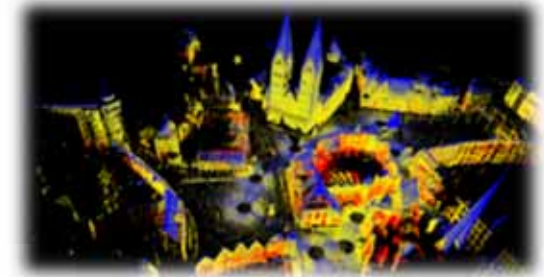
Speedup



Memory usage



Parallel & Scalable DBSCAN MPI/OpenMP Tool (4)



Selected 'Big Data' Applications

- London twitter data
(goal: find density centers of tweets)
- Bremen thermo point cloud data
(goal: noise reduction)
- PANGAEA earth science datasets
(goal: automated outlier detection)



[6] Open PANGAEA Earth Science Data Collection

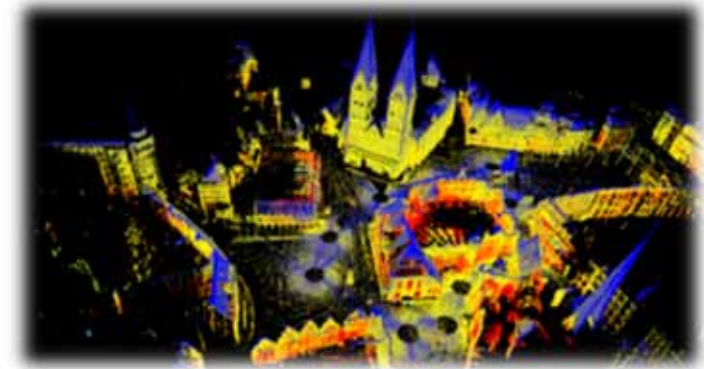
| Computation time | Cores | | | | | |
|------------------|------------|------------|------------|------------|-----------|-----------|
| | 1 | 2 | 4 | 8 | 16 | 32 |
| JSC-HPDBSCAN | 117,18 s | 59,64 s | 30,68 s | 16,25 s | 10,86 s | 9,39 s |
| NWU-PDSDBSCAN | 288,35 s | 162,47 s | 105,94 s | 89,87 s | 85,37 s | 88,42 s |
| Speed-Up | | | | | | |
| JSC-HPDBSCAN | 1,00 x | 1,96 x | 3,82 x | 7,21 x | 10,79 x | 12,48 x |
| NWU-PDSDBSCAN | 1,00 x | 1,77 x | 2,72 x | 3,21 x | 3,38 x | 3,26 x |
| Memory | | | | | | |
| JSC-HPDBSCAN | 251,064 MB | 345,276 MB | 433,340 MB | 678,248 MB | 1,101 GB | 2,111 GB |
| NWU-PDSDBSCAN | 500,512 MB | 725,104 MB | 1,370 GB | 4,954 GB | 19,724 GB | 59,685 GB |

Parallel & Scalable DBSCAN MPI/OpenMP Tool (5)

Free tool available

- Public bitbucket account – open-source
- Tool Website with more information
- Maintained on best effort basis

[3] M.Goetz & C. Bodenstern, HPDBSCAN Tool



3D Point Cloud of Bremen/Germany

→ Usage via simple jobscripts

Usage

- module load hdf5/1.8.13
- `mpiexec -np 1 ./dbscan -e 300 -m 100 -t 12 bremenSmall.h5`

Parameter
epsilon

Parameter
minPoints

Parallel & Scalable DBSCAN MPI/OpenMP Tool (6)

Usage via jobscript

- Using MOAB job scheduler
- Important: **module load hdf5/1.8.13**
- Important: library **gcc-4.9.2/lib64**
- np = number of processors
- t = number of threads



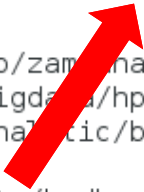
JUDGE @ Juelich

```
mriedel@judge:/homeb/zam/analytic/bigdata/hpdbscan/jsc_mpi/mriruns> more datajobscript.sh
#!/bin/bash
#MSUB -N HPDBSCAN_BremenSmall_1_12
#MSUB -l nodes=1:ppn=12:gpus=0:performance
#MSUB -l walltime=00:03:00
#MSUB -M m.riedel@fz-juelich.de
#MSUB -m abe
#MSUB -v tpt=12
#MSUB -l vmem=64gb
#MSUB -q devel

module load hdf5/1.8.13
export LD_LIBRARY_PATH=/homeb/zam/analytic/bigdata/hpdbscan/gcc-4.9.2/lib64:$LD_LIBRARY_PATH
DBSCAN=/homeb/zam/analytic/bigdata/hpdbscan/jsc_mpi/dbscan
SMALLBREMENDATA=/homeb/zam/analytic/bigdata/hpdbscan/jsc_mpi/mriruns/bremenSmall.h5

cd /homeb/zam/analytic/bigdata/hpdbscan/jsc_mpi/mriruns
mpiexec -np 1 $DBSCAN -e 300 -m 100 -t 12 $SMALLBREMENDATA
```

DBSCAN
Parameters



Parallel & Scalable DBSCAN MPI/OpenMP Tool (7)

Output with various information

- Run-times of different stages
- Clustering task information (e.g. number of identified clusters)
- Noise identification
- Data volume (small Bremen): ~72 MB
- Data volume (large Bremen): ~1.9 GB



JUDGE @ Juelich

```
mriedel@judge:/home/zam/analytic/bigdata/hpdbscan/jsc_mpi/mriruns> more HPDBSCAN_BremenSmall_1_12.o2208066
Calculating Cell Space...
  Computing Dimensions... [OK] in 0.011853
  Computing Cells...      [OK] in 0.073445
  Sorting Points...       [OK] in 0.124476
  Distributing Points...  [OK] in 0.000000
DBSCAN...
  Local Scan... I am ready 0
in 90.606330      [OK] in 90.606364
  Merging Neighbors...   [OK] in 0.000000
  Adjust Labels ...      [OK] in 0.004972
  Rec. Init. Order ...   [OK] in 1.255420
  Writing File ...       [OK] in 0.019120
Result...
  65      Clusters
  2973821 Cluster Points
  26179   Noise Points
  2953129 Core Points
Took: 92.214843s
```

Output results written in same input data:

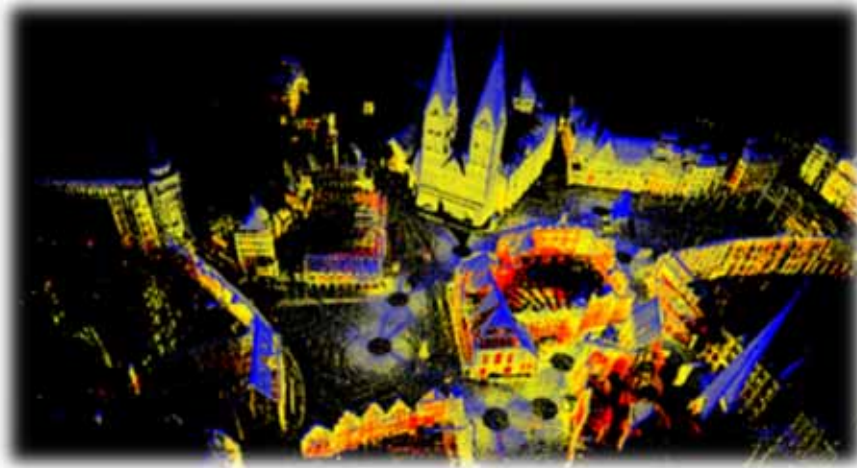
cluster number & noise label
(depends on parameters)



Parallel & Scalable DBSCAN MPI/OpenMP Tool (8)

Visualization Example

- Using Point Cloud Library (PDL) toolset
- Transformation of Data to PCD format (python script on the right)



Usage

- `python H5toPCD.py bremenSmall.h5`
- `pcl_viewer bremenClustered.pcd`

```
import h5py as h5
import numpy as np
import sys

if len(sys.argv) < 2:
    INPUT="bremen.h5"
else:
    INPUT = sys.argv[1]
FILE = "bremenClustered.pcd"

print"loading H5"
bremen = h5.File("bremenSmall.h5")
points = bremen["DBSCAN"]
clusters = bremen["Clusters"]
colors = bremen["COLORS"]

print "Transform to numpy"
points = np.array(points)
clusters = np.array(clusters)
colors = np.array(colors)

#print "Remove Noise"
#points = points[clusters!=0]
#clusters = clusters[clusters!=0]

#data = np.concatenate((points,colors.reshape((-1,1))),axis=1)
data = np.concatenate((points,clusters.reshape((-1,1))),axis=1)

clusters[clusters!=0]=1
data = np.concatenate((data,clusters.reshape((-1,1))),axis=1)

print "Write PCD"
with open(FILE, "w+") as out:
    out.write("""# .PCD v0.7 - Point Cloud Data file format
VERSION 0.7
FIELDS x y z rgb noise
SIZE 4 4 4 4 4
TYPE F F F F F
COUNT 1 1 1 1 1
WIDTH %d
HEIGHT 1
VIEWPOINT 0 -50000 -50000 1 0 0 0
POINTS %d
DATA ascii
""")
    np.savetxt(out, data)
```

H5toPCD.py
python
script

Take advantage
of NumPy library

Parallel & Scalable DBSCAN MPI/OpenMP Tool (9)

Earth Science Application

‘Automated outlier detection in time series‘

- Collaboration with MARUM, Bremen (work in progress)
- Example: water quality data of Koljoefjords
- Connected underwater device
- Measurements: oxygen, temperature, salinity, ...



Use of HPBSCAN algorithm

- Detect outliers and anomalies/events (e.g. water mixing)
- Compare with manually annotated data by domain-scientist
- Needs automation

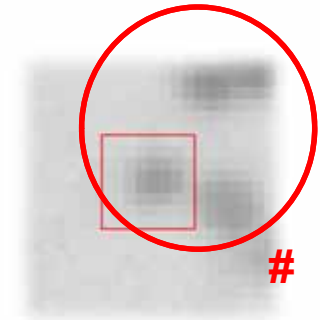


Parallel & Scalable DBSCAN MPI/OpenMP Tool (10)

Neuroscience Application

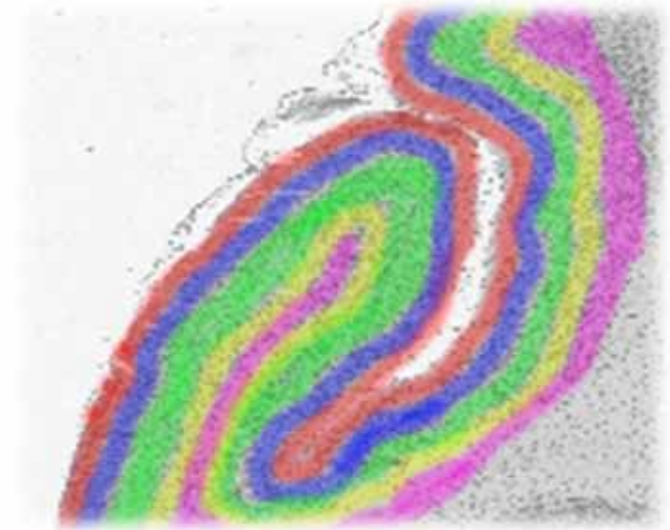
‘Cell nuclei detection and tissue clustering’

- Scientific Case: Detect various layers (colored)
- Layers seem to have different density distribution of cells
- Extract cell nuclei into 2D/3D point cloud
- Cluster different brain areas by cell density



Use of HPBSCAN algorithm

- First 2d results detect various clusters
- Work in progress, not very good results
- Approach: Several iterations (with 3D) with potentially different parameter values
- Investigate other methods (e.g. OPTICS)

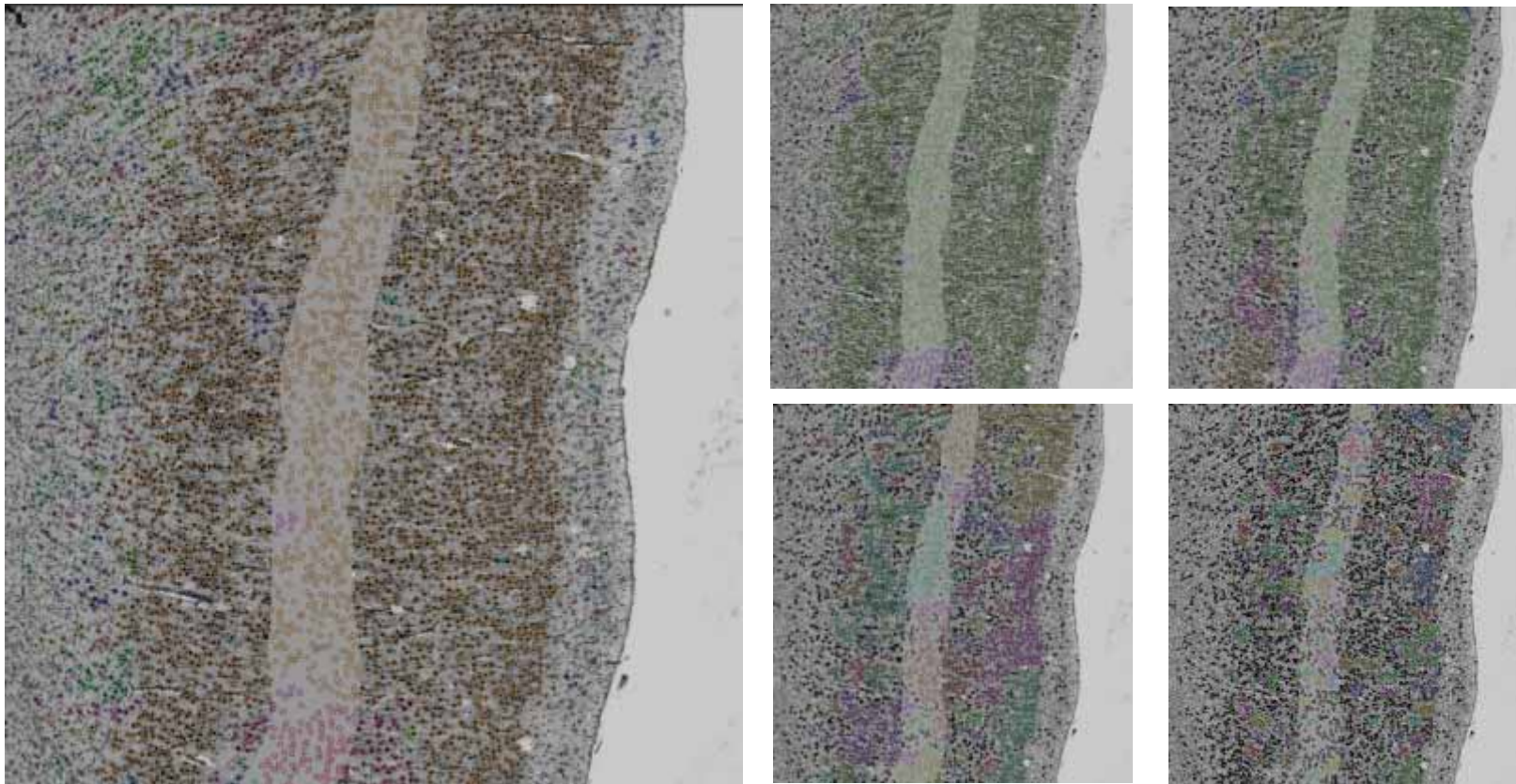


➤ Research activities jointly with T. Dickscheid et al. (Juelich Institute of Neuroscience & Medicine)

Parallel & Scalable DBSCAN MPI/OpenMP Tool (11)

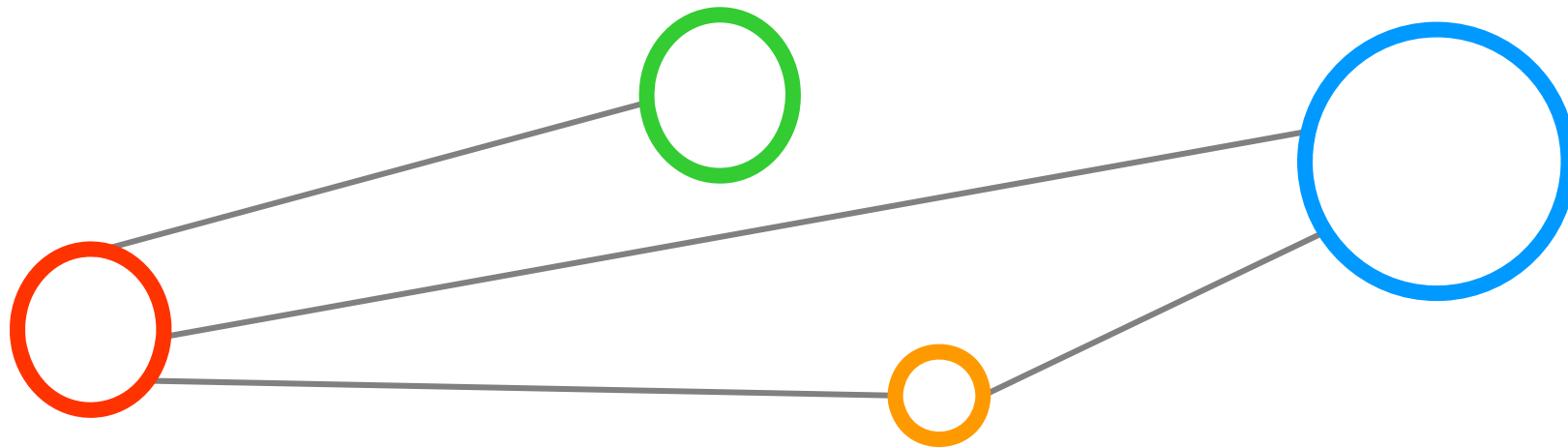
Neuroscience Application – Work in progress (e.g. 3120x3288)

‘Cell nuclei detection and tissue clustering’ – varying parameters



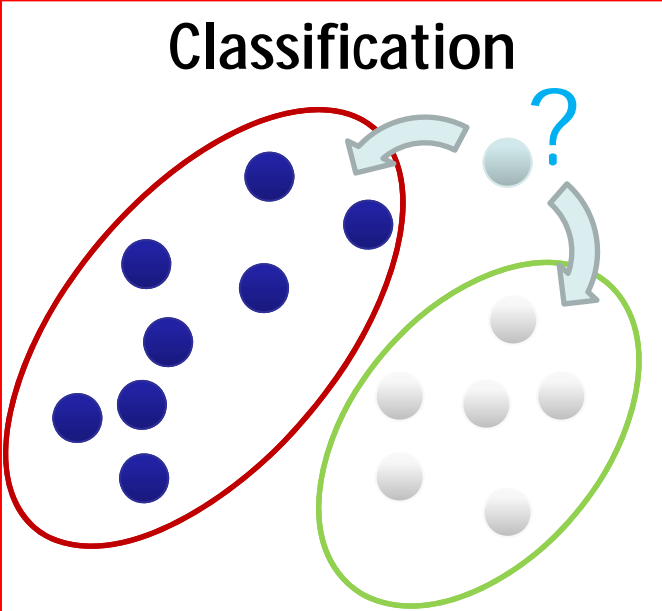
➤ Research activities jointly with T. Dickscheid et al. (Juelich Institute of Neuroscience & Medicine)

Scalable & Parallel Tools: Classification



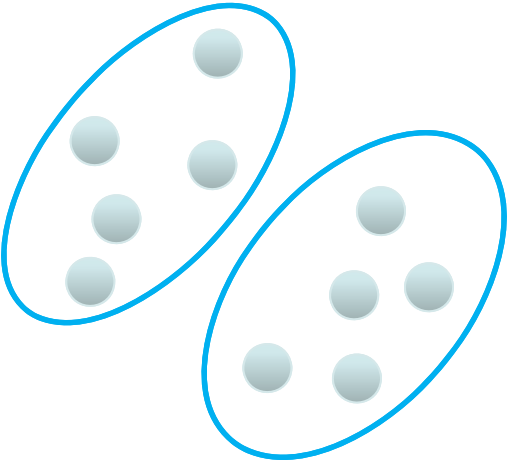
Learning From Data – Classification Technique

Classification



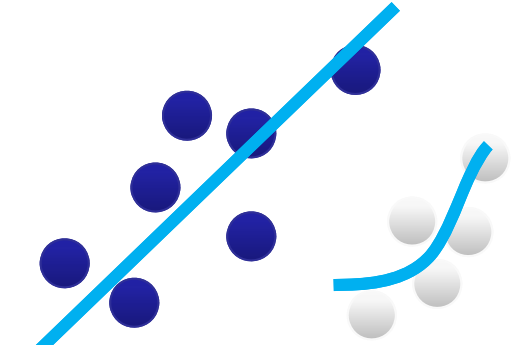
- Groups of data exist
- New data classified to existing groups

Clustering



- No groups of data exist
- Create groups from data close to each other

Regression



- Identify a line with a certain slope describing the data

Selected Classification Methods

Perceptron Learning Algorithm – simple linear classification

- Enables binary classification with ‘a line’ between classes of separable data

Support Vector Machines (SVMs) – non-linear (‘kernel’) classification

- Enables non-linear classification with maximum margin (best ‘out-of-the-box’)

Reasoning: achieves often better results than other methods in remote sensing application

Decision Trees & Ensemble Methods – tree-based classification

- Grows trees for class decisions, ensemble methods average n trees

Artificial Neural Networks (ANNs) – brain-inspired classification

- Combine multiple linear perceptrons to a strong network for non-linear tasks

Naive Bayes Classifier – probabilistic classification

- Use of the Bayes theorem with strong/naive independence between features

Technology Review of Open & Available Tools

| Technology | Platform Approach | Analysis |
|-----------------------|---------------------------|--|
| Apache Mahout | Java; Hadoop | No parallelization strategy for SVMs |
| Apache Spark/MLlib | Java; Spark | Parallel linear SVMs (no multi-class) |
| Twister/ParallelSVM | Java; Twister; Hadoop 1.0 | Parallel SVMs, open source; developer version 0.9 beta |
| scikit-learn | Python | No parallelization strategy for SVMs |
| piSVM 1.2 & piSVM 1.3 | C; MPI | Parallel SVMs; stable; not fully scalable |
| GPU LibSVM | CUDA | Parallel SVMs; hard to programs, early versions |
| pSVM | C; MPI | Parallel SVMs; unstable; beta version |

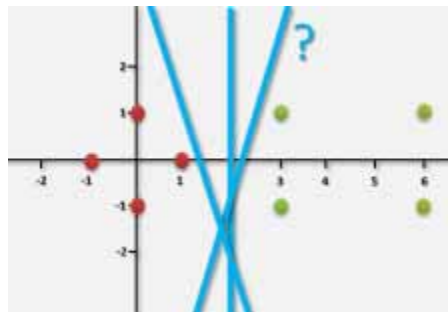
M. Goetz, M. Riedel et al., 6th Workshop on Data Mining in Earth System Science, International Conference of Computational Science (ICCS), Reykjavik, to be published

Parallel & Scalable SVM MPI Tool (1)

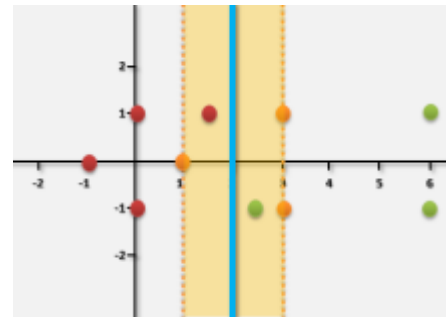
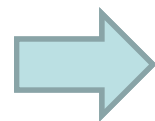
SVM Algorithm Approach

[7] C. Cortes and V. Vapnik et al.

- Introduced 1995 by C. Cortes & V. Vapnik et al.
- Creates a 'maximal margin classifier' to get future points ('more often') right
- Uses quadratic programming & Lagrangian method with $N \times N$



(linear example)



('maximal margin classifier' example)

(use of soft-margin approach for better generalization)

$$\min_{w, \xi_i, b} \left\{ \frac{1}{2} \|\mathbf{w}\|^2 + C \sum_i \xi_i \right\}$$

$$y_i(\mathbf{w} \cdot \mathbf{x}_i - b) \geq 1 - \xi_i, \quad \xi_i \geq 0$$

(maximizing hyperplane turned into optimization problem, minimization, dual problem)

$$\mathcal{L}(\alpha) = \sum_{n=1}^N \alpha_n - \frac{1}{2} \sum_{n=1}^N \sum_{m=1}^N y_n y_m \alpha_n \alpha_m \mathbf{x}_n^T \mathbf{x}_m$$

(max. hyperplane → dual problem, using quadratic programming method)

$$k(\mathbf{x}_i, \mathbf{x}_j) = \langle \phi(\mathbf{x}_i), \phi(\mathbf{x}_j) \rangle$$

$$0 \leq \alpha_i \leq C$$

$$\begin{bmatrix} y_1 y_1 x_1^T x_1 & y_1 y_2 x_1^T x_2 & \dots & y_1 y_N x_1^T x_N \\ \dots & \dots & \dots & \dots \\ y_N y_1 x_N^T x_1 & y_N y_2 x_N^T x_2 & \dots & y_N y_N x_N^T x_N \end{bmatrix}$$

(kernel trick, quadratic coefficients – Computational Complexity & Big Data Impact)

Parallel & Scalable SVM MPI Tool (2)

- True Support Vector Machines are Support Vector Classifiers combined with a non-linear kernel
- Non-linear kernels exist - mostly known are polynomial & Radial Basis Function (RBF) kernels

[8] *An Introduction to Statistical Learning*

Understanding the MPI tool parameters

- Selecting non-linear kernel function K type as RBF → parameter -t 2
- Setting RBF Kernel configuration parameter γ → e.g. parameter -g 16
- Setting SVM allowed errors parameter → e.g. parameter -c 10000

Major benefit of Kernels: Computing done in original space

- Linear Kernel
$$K(x_i, x_{i'}) = \sum_{j=1}^p x_{ij} x_{i'j} \quad (\text{linear in features})$$
- Polynomial Kernel
$$K(x_i, x_{i'}) = \left(1 + \sum_{j=1}^p x_{ij} x_{i'j}\right)^d \quad (\text{polynomial of degree } d)$$
- RBF Kernel
$$K(x_i, x_{i'}) = \exp\left(-\gamma \sum_{j=1}^p (x_{ij} - x_{i'j})^2\right) \quad (\text{large distance, small impact})$$

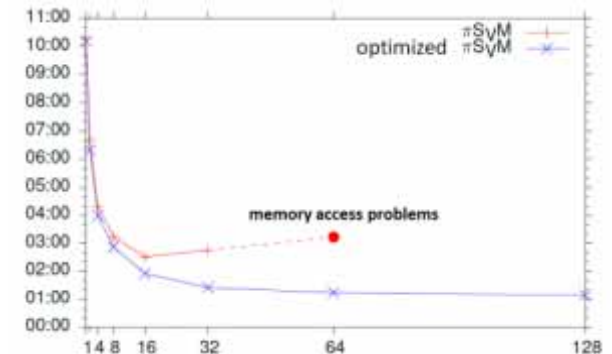
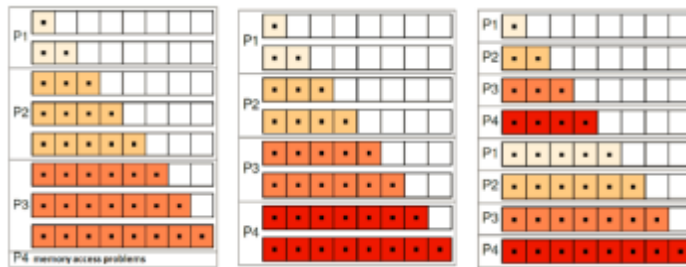
Parallel & Scalable SVM MPI Tool (3)

Original parallel piSVM tool 1.2

- Open-source and based on libSVM library, C, 2011
- Message Passing Interface (MPI)
- New version appeared 2014-10 v. 1.3 (no major improvements)
- Lack of 'big data' support (memory, layout, etc.)

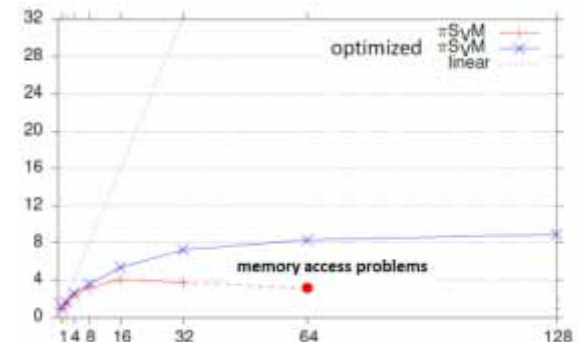


[9] piSVM Website, 2011/2014 code

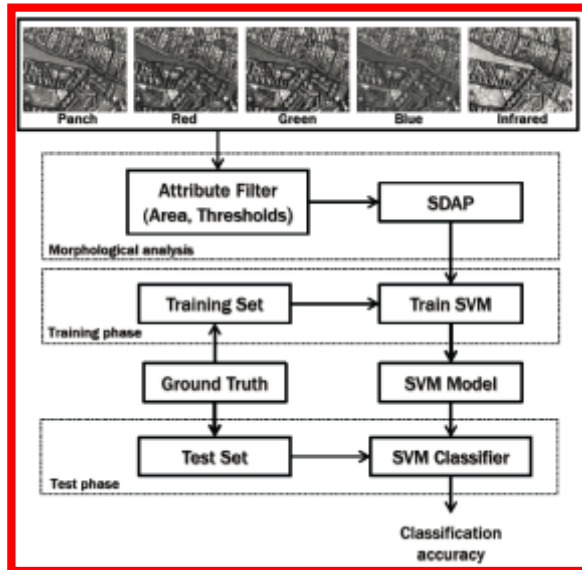
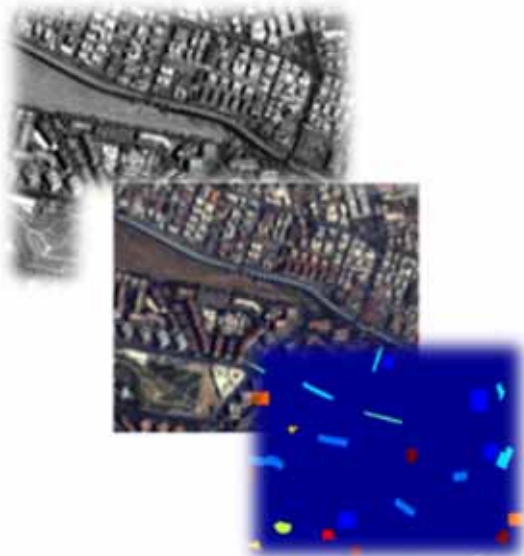


Tuned scalable parallel piSVM tool 1.2.1

- Open-source (repository to be created)
- Based on piSVM tool 1.2
- Optimizations: load balancing; MPI collectives
- Contact: m.richerzhagen@fz-juelich.de



Parallel & Scalable SVM MPI Tool (4)



| Class | Training | Test |
|--------------|--------------|---------------|
| Buildings | 18126 | 163129 |
| Blocks | 10982 | 98834 |
| Roads | 16353 | 147176 |
| Light Train | 1606 | 14454 |
| Vegetation | 6962 | 62655 |
| Trees | 9088 | 81792 |
| Bare Soil | 8127 | 73144 |
| Soil | 1506 | 13551 |
| Tower | 4792 | 43124 |
| Total | 77542 | 697859 |

Sattelite Data (Quickbird)

Parallel Support Vector Machines (SVM)

HPC / MPI

Classification Study of Land Cover Types

„Reference Data Analytics“ for reusability & learning

CRISP-DM Report



Openly Shared Datasets



Running Analytics Code



[10] Rome Image dataset

Parallel & Scalable SVM MPI Tool (5)

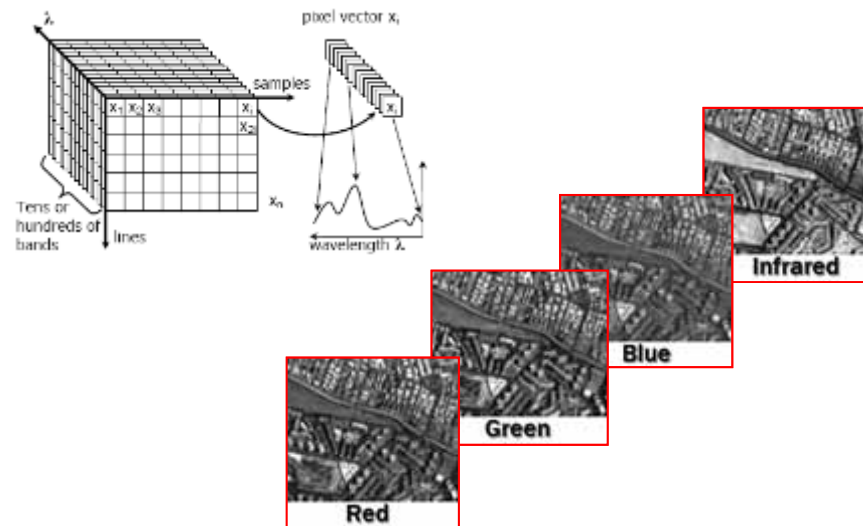
Example dataset: Geographical location: Image of Rome, Italy

- Remote sensor data obtained by Quickbird satellite

High-resolution (0.6m)
panchromatic image



Pansharpened (UDWT) low-resolution
(2.4m) multispectral images



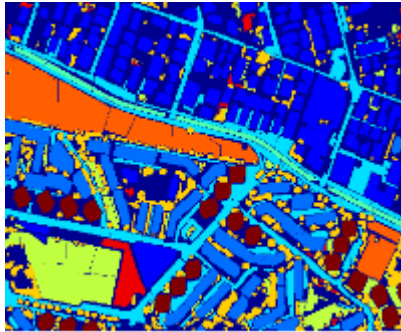
[10] Rome Image dataset



Parallel & Scalable SVM MPI Tool (6)

Labelled data available for train/test data

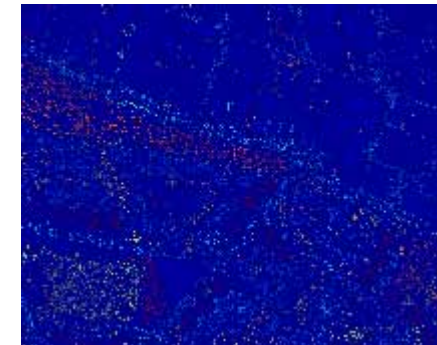
- Groundtruth data of 9 different land-cover classes available



| Class | Training | Test |
|-------------|----------|--------|
| Buildings | 18126 | 163129 |
| Blocks | 10982 | 98834 |
| Roads | 16353 | 147176 |
| Light Train | 1606 | 14454 |
| Vegetation | 6962 | 62655 |
| Trees | 9088 | 81792 |
| Bare Soil | 8127 | 73144 |
| Soil | 1506 | 13551 |
| Tower | 4792 | 43124 |
| Total | 77542 | 697859 |

Data preparation

- We generated a set of training samples by randomly selecting 10% of the reference samples (with labelled data)
- Generated set of test samples from the remaining labels (labelled data, 90% of reference samples)



Training Image
(10% pixels/class)

[10] Rome Image dataset



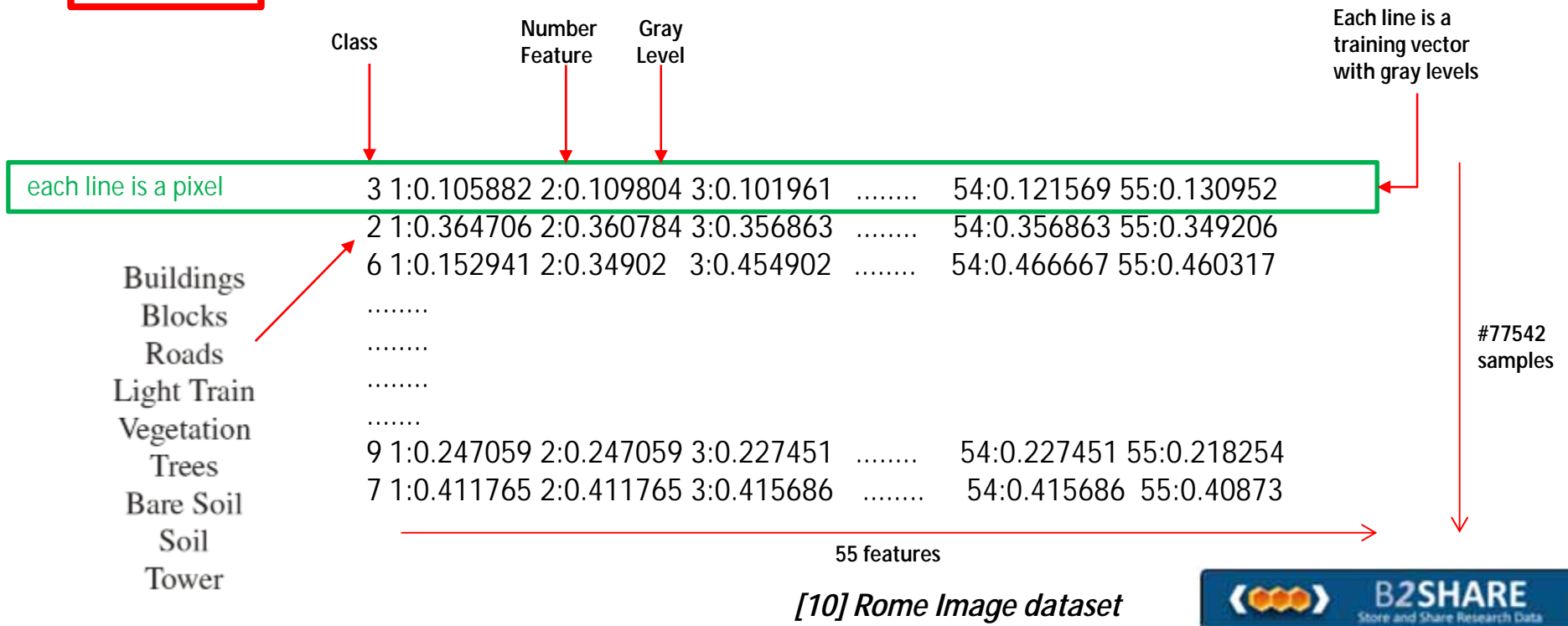
Parallel & Scalable SVM MPI Tool (7)

Based on 'LibSVM data format' (using feature extraction method)

- Add 'Self-Dual Attribute Profile (SDAP) on Area' on all images training file



[11] G. Cavallaro, M. Mura, J.A. Benediktsson, L. Bruzzone et al.



Parallel & Scalable SVM MPI Tool (8)

Usage via jobscript

- Using MOAB job scheduler
- np = number of processors; o/q partitioning

```
#!/bin/bash
#MSUB -N Train-tune-rec86-4-16-32
#MSUB -l nodes=4:ppn=16:performance
#MSUB -l walltime=03:00:00
#MSUB -M m.riedel@fz-juelich.de
#MSUB -m abe
#MSUB -W x=naccesspolicy:singlejob
#MSUB -v tpt=2
#MSUB -q devel
```

```
### jobscript
```

```
cd $PBS_O_WORKDIR
echo "workdir: $PBS_O_WORKDIR"
```

```
NSLOTS=32
```

```
echo "running on $NSLOTS cpus..."
```

```
### location
```

```
PISVM=/homeb/zam/mriedel/pisvm-1.2/pisvm-1.2/pisvm-train
```

```
TRAINDATA=/homeb/zam/mriedel/bigdata/86-romeok/sdap_area_all_training.el
```

```
### submit
```

```
mpirexec -np $NSLOTS $PISVM -o 1024 -q 512 -c 10000 -g 16 -t 2 -m 1024 -s 0 $TRAINDATA
```

→ Usage via simple jobscripts

SVM
Parameters

[12] Rome Analytics Results & job scripts



JUDGE @ Juelich



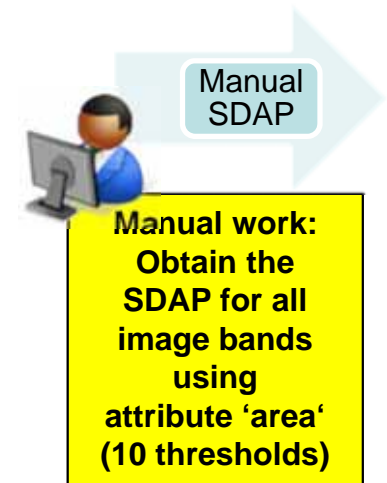
Parallel & Scalable SVM MPI Tool (9)

Training speed-up is possible when number of features is 'high'

- Serial Matlab: ~1277 sec (~21 minutes)
- Parallel (16) Analytics: 220 sec (3:40 minutes)
- Accuracy remains

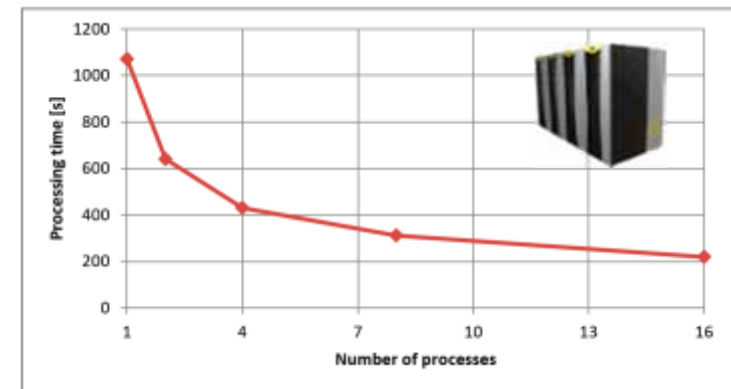
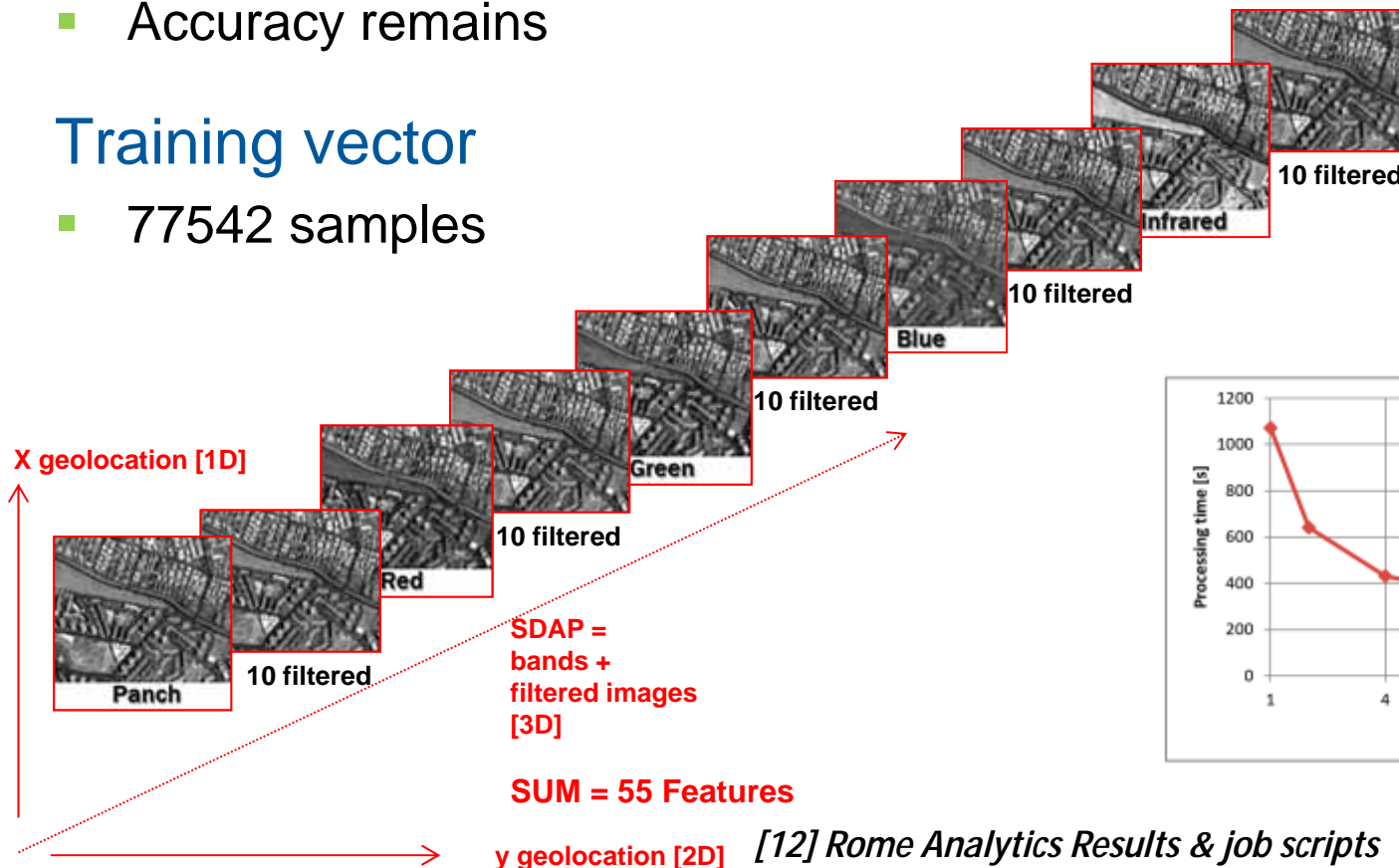
Training vector

- 77542 samples



Manual SDAP

Manual work:
Obtain the SDAP for all image bands using attribute 'area' (10 thresholds)



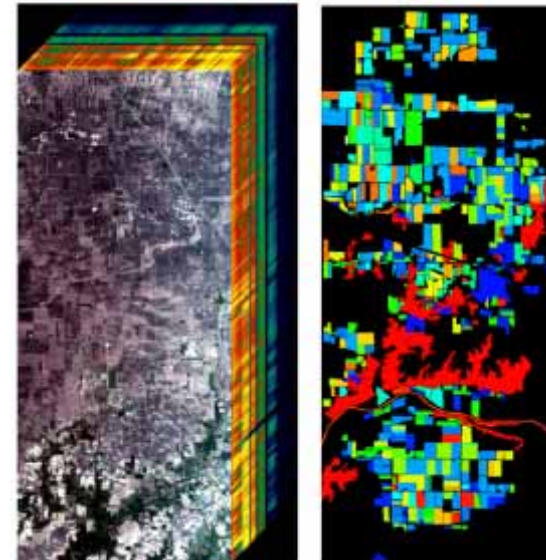
Parallel & Scalable SVM MPI Tool (10)

Another more challenging dataset: high number of classes

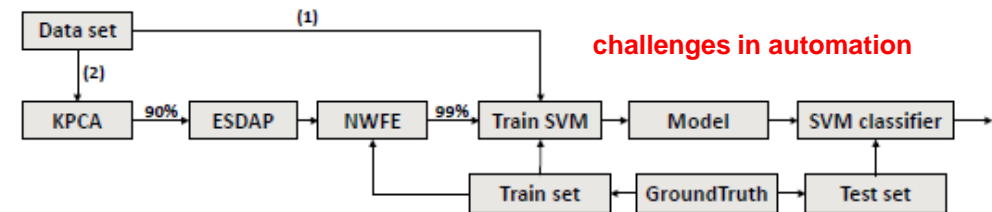
- Parallelization challenges: unbalanced class representations

| Class | | Number of samples | | Class | | Number of samples | |
|--------|-----------------------------|-------------------|-------|--------|----------------------------|-------------------|-------|
| number | name | training | test | number | name | training | test |
| 1 | Buildings | 1720 | 15475 | 27 | Pasture | 1039 | 9347 |
| 2 | Corn | 1778 | 16005 | 28 | pond | 10 | 92 |
| 3 | Corn? | 16 | 142 | 29 | Soybeans | 939 | 8452 |
| 4 | Corn-EW | 51 | 463 | 30 | Soybeans? | 89 | 805 |
| 5 | Corn-NS | 236 | 2120 | 31 | Soybeans-NS | 111 | 999 |
| 6 | Corn-CleanTill | 1240 | 11164 | 32 | Soybeans-CleanTill | 507 | 4567 |
| 7 | Corn-CleanTill-EW | 2649 | 23837 | 33 | Soybeans-CleanTill? | 273 | 2453 |
| 8 | Corn-CleanTill-NS | 3968 | 35710 | 34 | Soybeans-CleanTill-EW | 1180 | 10622 |
| 9 | Corn-CleanTill-NS-Irrigated | 80 | 720 | 35 | Soybeans-CleanTill-NS | 1039 | 9348 |
| 10 | Corn-CleanTilled-NS? | 173 | 1555 | 36 | Soybeans-CleanTill-Drilled | 224 | 2018 |
| 11 | Corn-MinTill | 105 | 944 | 37 | Soybeans-CleanTill-Weedy | 54 | 489 |
| 12 | Corn-MinTill-EW | 563 | 5066 | 38 | Soybeans-Drilled | 1512 | 13606 |
| 13 | Corn-MinTill-NS | 886 | 7976 | 39 | Soybeans-MinTill | 267 | 2400 |
| 14 | Corn-NoTill | 438 | 3943 | 40 | Soybeans-MinTill-EW | 183 | 1649 |
| 15 | Corn-NoTill-EW | 121 | 1085 | 41 | Soybeans-MinTill-Drilled | 810 | 7288 |
| 16 | Corn-NoTill-NS | 569 | 5116 | 42 | Soybeans-MinTill-NS | 495 | 4458 |
| 17 | Fescue | 11 | 103 | 43 | Soybeans-NoTill | 216 | 1941 |
| 18 | Grass | 115 | 1032 | 44 | Soybeans-NoTill-EW | 253 | 2280 |
| 19 | Grass/Trees | 233 | 2098 | 45 | Soybeans-NoTill-NS | 93 | 836 |
| 20 | Hay | 113 | 1015 | 46 | Soybeans-NoTill-Drilled | 873 | 7858 |
| 21 | Hay? | 219 | 1966 | 47 | Swampy Area | 58 | 525 |
| 22 | Hay-Alfalfa | 226 | 2032 | 48 | River | 311 | 2799 |
| 23 | Lake | 22 | 202 | 49 | Trees? | 58 | 522 |
| 24 | NotCropped | 194 | 1746 | 50 | Wheat | 498 | 4481 |
| 25 | Oats | 174 | 1568 | 51 | Woods | 6356 | 57206 |
| 26 | Oats? | 34 | 301 | 52 | Woods? | 14 | 130 |

remote sensing cube & ground reference



G. Cavallaro, M. Riedel et al., *Remote Sensing Journal – Big Data Special Issue*, to be published

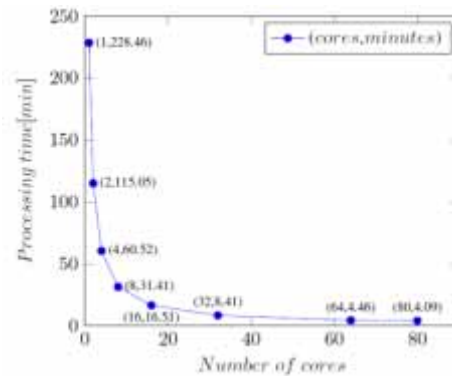
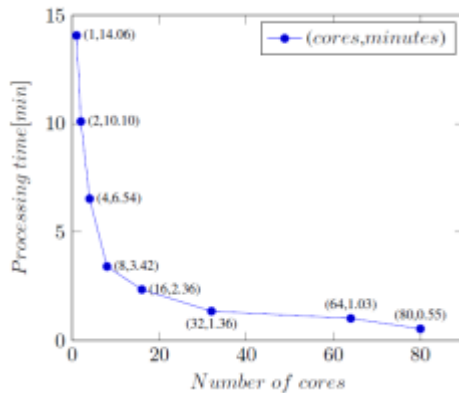


[20] Indian pines dataset, processed and raw

Parallel & Scalable SVM MPI Tool (11)

Another example dataset: high number of classes

- Parallelization benefits: major speed-ups, ~interactive (<1 min) possible

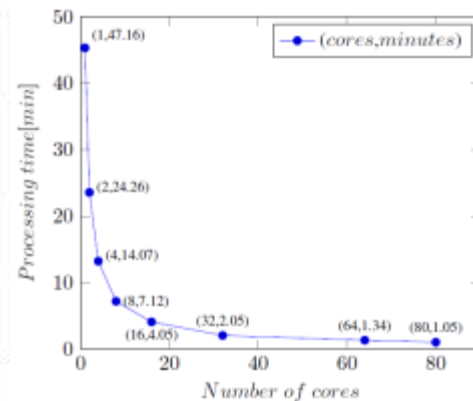
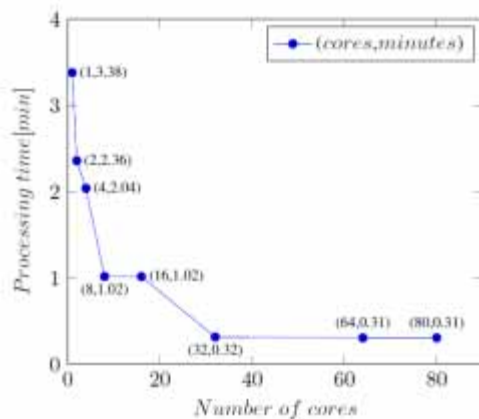


manual & serial activities (in minutes)

| | kpca | esdap | nwfe | 10x CSV | Training | Test | Total |
|--------------|------|-------|------|--------------------|----------|-------|--------------------|
| (1) Scenario | 0 | 0 | 0 | 4.47×10^3 | 10,45 | 71,08 | 4.55×10^3 |
| (2) Scenario | 5 | 15.38 | 1 | 529.55 | 1.37 | 23.25 | 575.55 |

'big data' is not always better data

| | (1) Scenario | (2) Scenario |
|----------------------|--------------|--------------|
| Number of features | 200 | 30 |
| Overall Accuracy (%) | 40,68 | 77,96 |



Manual WORK

Manual work:
Trade-off between raw data processing and using feature extraction methods

Can we automate feature extraction mechanism to some degree?

G. Cavallaro, M. Riedel et al., Remote Sensing Journal – Big Data Special Issue, to be published



[21] Analytics Results (raw)

[22] Analytics Results (processed)

Parallel & Scalable SVM MPI Tool (12)

2x benefits of parallelization (shown in n-fold cross validation)

- Evaluation between Matlab (aka serial) and parallel piSVM
- 10x cross-validation (RBF kernel parameter and C, gridsearch)

raw dataset (serial)

| γ / C | 1 | 10 | 100 | 1000 | 10000 |
|--------------|----------------|-----------------------|----------------|----------------|----------------|
| 2 | 27.30 (109.78) | 34.59 (124.46) | 39.05 (107.85) | 37.38 (116.29) | 37.20 (121.51) |
| 4 | 29.24 (98.18) | 37.75 (85.31) | 38.91 (113.87) | 38.36 (119.12) | 38.36 (118.98) |
| 8 | 31.31 (109.95) | 39.68 (118.28) | 39.06 (112.99) | 39.06 (190.72) | 39.06 (872.27) |
| 16 | 33.37 (126.14) | 39.46 (171.11) | 39.19 (206.66) | 39.19 (181.82) | 39.19 (146.98) |
| 32 | 34.61 (179.04) | 38.37 (202.30) | 38.37 (231.10) | 38.37 (240.36) | 38.37 (278.02) |

processed dataset (serial)

| γ / C | 1 | 10 | 100 | 1000 | 10000 |
|--------------|---------------|---------------|----------------------|---------------|---------------|
| 2 | 48.90 (18.81) | 65.01 (19.57) | 73.21 (20.11) | 75.55 (22.53) | 74.42 (21.21) |
| 4 | 57.53 (16.82) | 70.74 (13.94) | 75.94 (13.53) | 76.04 (14.04) | 74.06 (15.55) |
| 8 | 64.18 (18.30) | 74.45 (15.04) | 77.00 (14.41) | 75.78 (14.65) | 74.58 (14.92) |
| 16 | 68.37 (23.21) | 76.20 (21.88) | 76.51 (20.69) | 75.32 (19.60) | 74.72 (19.66) |
| 32 | 70.17 (34.45) | 75.48 (34.76) | 74.88 (34.05) | 74.08 (34.03) | 73.84 (38.78) |

raw dataset (parallel, 80 cores)

| γ / C | 1 | 10 | 100 | 1000 | 10000 |
|--------------|--------------|---------------------|--------------|---------------|---------------|
| 2 | 27.26 (3.38) | 34.49 (3.35) | 39.16 (5.35) | 37.56 (11.46) | 37.57 (13.02) |
| 4 | 29.12 (3.34) | 37.58 (3.38) | 38.91 (6.02) | 38.43 (7.47) | 38.43 (7.47) |
| 8 | 31.24 (3.38) | 39.77 (4.09) | 39.14 (5.45) | 39.14 (5.42) | 39.14 (5.43) |
| 16 | 33.36 (4.09) | 39.61 (4.56) | 39.25 (5.06) | 39.25 (5.27) | 39.25 (5.10) |
| 32 | 34.61 (5.13) | 38.37 (5.30) | 38.36 (5.43) | 38.36 (5.49) | 38.36 (5.28) |

processed dataset (parallel, 80 cores)

| γ / C | 1 | 10 | 100 | 1000 | 10000 |
|--------------|--------------|--------------|---------------------|--------------|--------------|
| 2 | 75.26 (1.02) | 65.12 (1.03) | 73.18 (1.33) | 75.76 (2.35) | 74.53 (4.40) |
| 4 | 57.60 (1.03) | 70.88 (1.02) | 75.87 (1.03) | 76.01 (1.33) | 74.06 (2.35) |
| 8 | 64.17 (1.02) | 74.52 (1.03) | 77.02 (1.02) | 75.79 (1.04) | 74.42 (1.34) |
| 16 | 68.57 (1.33) | 76.07 (1.33) | 76.40 (1.34) | 75.26 (1.05) | 74.53 (1.34) |
| 32 | 70.21 (1.33) | 75.38 (1.34) | 74.69 (1.34) | 73.91 (1.47) | 73.73 (1.33) |

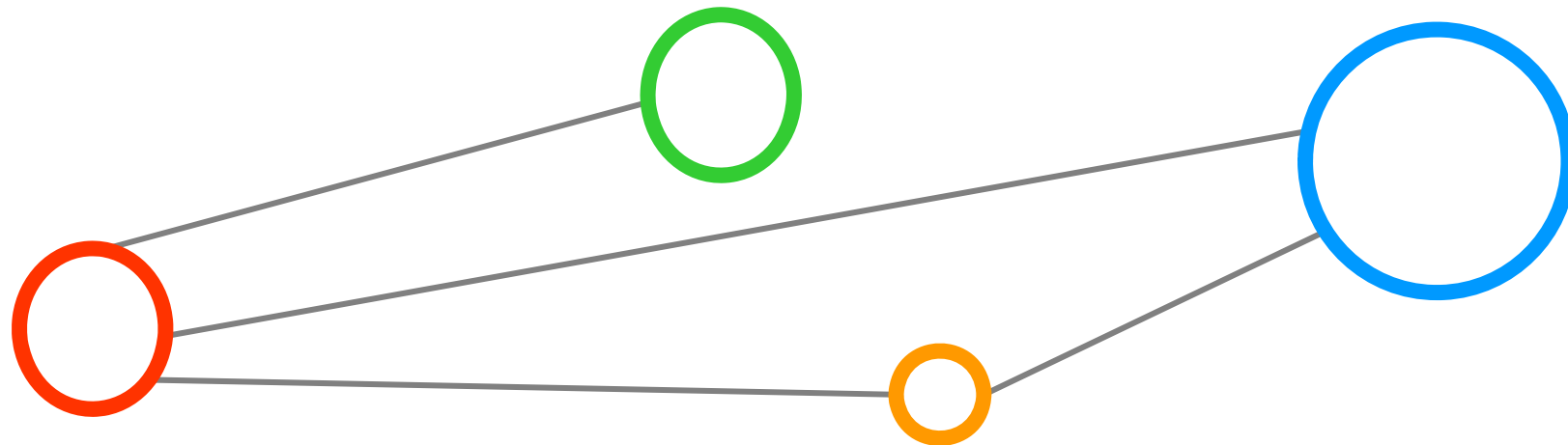
G. Cavallaro, M. Riedel et al., *Remote Sensing Journal – Big Data Special Issue, to be published*



[23] Analytics 10 fold cross-validation Results (raw)

[24] Analytics 10 fold cross-validation Results (processed)

Recent Research Directions



Recent Research Directions – Brain Data Classification

- Build 'reconstructed brain (one 3d volume) that matches with sections & block images
- Understanding the 'sectioning of the brain' and support automation of reconstruction

1. Some 'pattern' exists

- Image content classification (e.g. SVMs, RandomForst, etc.)

Smart Data
Innovation Lab

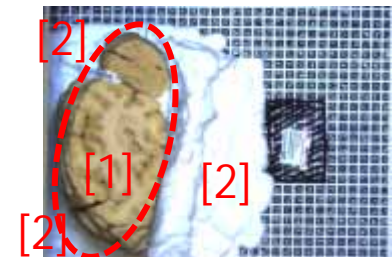


2. No exact mathematical formula exists

- No precise formula for 'contour of the brain'

3. Dataset (next: 5 brains, >100.000 pixels, 2PB raw)

- Block face images (of frozen brain tissue)
- Every 20 micron (cut size), resolution: 3272 x 2469
- ~ 14 MB / RGB image
- ~ 8 MB / corresponding mask image ('groundtruth')
- ~700 images → ~40 GB dataset



➤ Research activities jointly with T. Dickscheid et al. (Juelich Institute of Neuroscience & Medicine)

Recent Research Directions – Deep Learning

Investigate a pipeline for cell nuclei detection and tissue clustering

1. Some 'pattern' exists

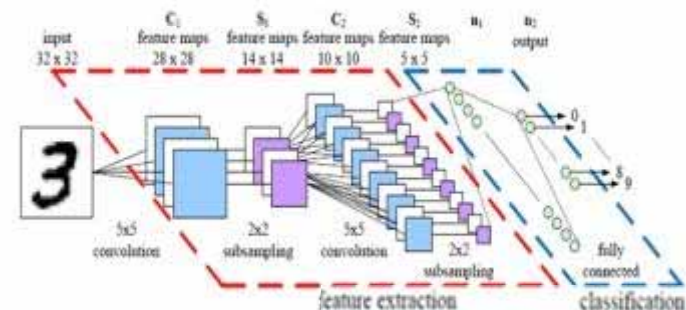
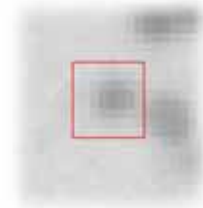
- Image content classification & clustering

2. No exact mathematical formula exists

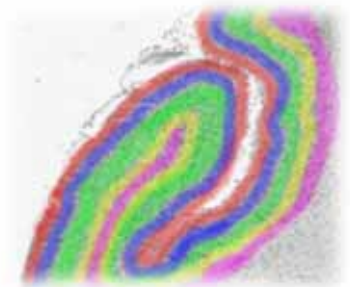
- No precise formula for 'brain layers'

3. Dataset – raw images exist

- Needs to be properly prepared
- Generate labeled data to learn from (manual tool supporting scientists)
- Use Deep Learning (deep convolutional neural network, GPGPUs) to classify cell nuclei
- Extract cell nuclei into 2D/3D point cloud
- Cluster different brain areas by cell density (parallel DBSCAN)

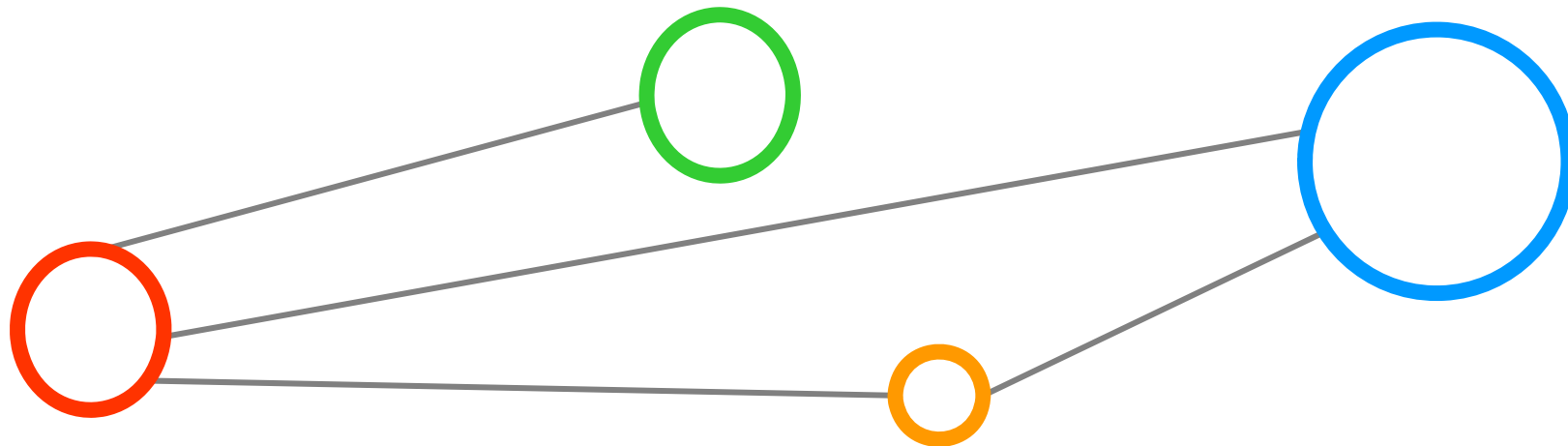


[13] Deep Learning Architecture



➤ Research activities jointly with T. Dickscheid et al. (Juelich Institute of Neuroscience & Medicine)

Conclusions



Conclusions

Scientific Peer Review is essential to progress in the field

- Work in the field needs to be guided & steered by communities
- NIC Scientific Big Data Analytics (SBDA) first step (learn from HPC)
- Towards enabling reproducibility by uploading runs and datasets

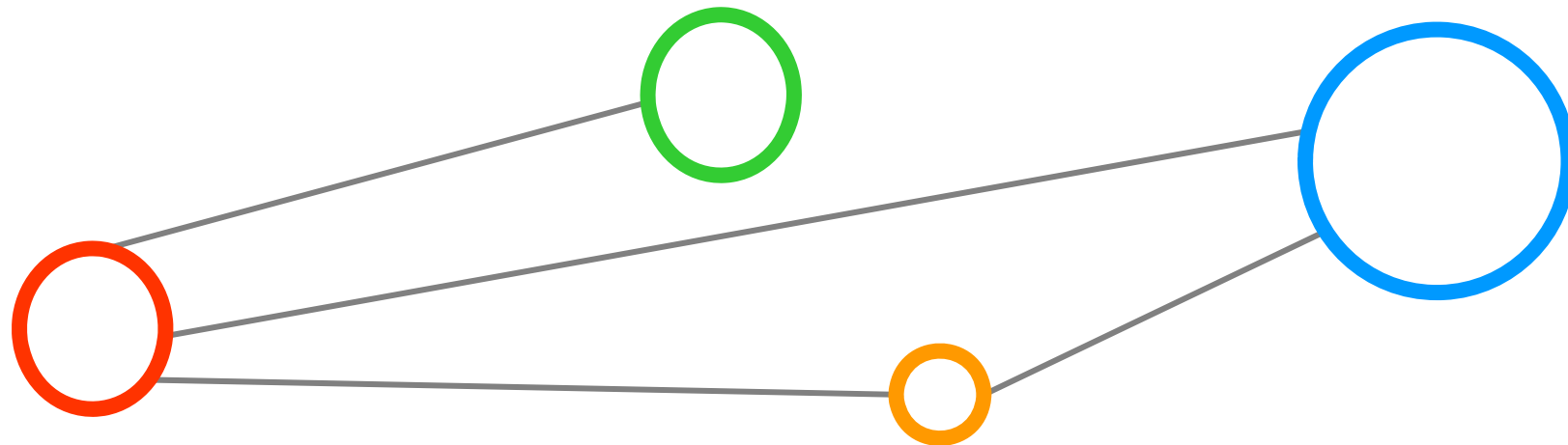
Selected SBDA benefit from parallelization

- Statistical data mining techniques able to reduce 'big data' (e.g. PCA, etc.)
- Benefits in n-fold cross-validation & raw data, less on preprocessed data
- Two codes available to use and maintained @JSC: HPDBSCAN, piSVM

Number of Data Analytics et al. Technologies incredible high

- Thorough analysis and evaluation hard (needs different infrastructures)
- (Less) open source & working versions available, often paper studies
- Still evaluating approaches: HPC, map-reduce, Spark, SciDB, MaTex, ...

References



References (1)

- [1] R. Huber, M. Riedel et al., 'PANGAEA – Data Publisher for Earth & Environmental Science - Research data enters scholarly communication and big data analysis', presentation at EGU 2014, Vienna
- [2] G. Cavallaro, J.A. Benediktsson and M. Riedel et al. 'Smart Data Analytics Methods for Remote Sensing Applications', in proceedings of IGARSS 2014
- [3] M.Goetz & C. Bodenstern, Clustering Highly Parallelizable DBSCAN Algorithm, JSC, Online: http://www.fz-juelich.de/ias/jsc/EN/Research/DistributedComputing/DataAnalytics/Clustering/Clustering_node.html
- [4] Ester, Martin, et al. "A density-based algorithm for discovering clusters in large spatial databases with noise." Kdd. Vol. 96. 1996.
- [5] Patwary, Md Mostofa Ali, et al. "A new scalable parallel dbscan algorithm using the disjoint-set data structure." High Performance Computing, Networking, Storage and Analysis (SC), 2012 International Conference for. IEEE, 2012
- [6] PANGAEA Earth Science Data Collection, Online: <http://www.pangaea.de/>
- [7] C. Cortes and V. Vapnik, "Support-vector networks," Machine Learning, vol. 20(3), pp. 273–297, 1995
- [8] An Introduction to Statistical Learning with Applications in R,
Online: <http://www-bcf.usc.edu/~gareth/ISL/index.html>
- [9] Original piSVM tool, online: <http://pisvm.sourceforge.net/>
- [10] B2SHARE data collection Rome Dataset,
online: <http://hdl.handle.net/11304/4615928c-e1a5-11e3-8cd7-14feb57d12b9>
- [11] G. Cavallaro, M. Mura, J.A. Benediktsson, L. Bruzzone 'A Comparison of Self-Dual Attribute Profiles based on different filter rules for classification', IEEE IGARSS2014, Quebec, Canada
- [12] B2SHARE data collection piSVM1.2 Analytics JUDGE Cluster Rome Images 55 Features, online: <http://hdl.handle.net/11304/6880662c-1edf-11e4-81ac-dcbd1b51435e>

References (2)

- [13] Deep Learning Architecture, Online: <http://parse.ele.tue.nl/cluster/2/CNNArchitecture.jpg>
- [14] UNICORE, online: <http://www.unicore.eu>
- [15] DOE ASCAC Report, 'Synergistic Challenges in Data-Intensive Science and. Exascale Computing'
- [16] M. Riedel and P. Wittenburg et al. 'A Data Infrastructure Reference Model with Applications: Towards Realization of a ScienceTube Vision with a Data Replication Service', 2013
- [17] Shearer C., 'The CRISP-DM model: the new blueprint for data mining', J Data Warehousing (2000); 5:13—22.
- [18] Jeremy Ginsburg et al., 'Detecting influenza epidemics using search engine query data', Nature 457, 2009
- [19] David Lazer, Ryan Kennedy, Gary King, and Alessandro Vespignani, 'The Parable of Google Flu: Traps in Big Data Analysis', Science Vol (343), 2014
- [20] B2SHARE data collection, remote sensing indian pines images, Online: <http://hdl.handle.net/11304/7e8eec8e-ad61-11e4-ac7e-860aa0063d1f>
- [21] B2SHARE data collection, piSVM remote sensing indian pines analytics results (raw), Online: <http://hdl.handle.net/11304/c06a8c7e-fe6c-11e4-8a18-f31aa6f4d448>
- [22] B2SHARE data collection, piSVM remote sensing indian pines analytics results (processed), Online: <http://hdl.handle.net/11304/c528998e-ff7c-11e4-8a18-f31aa6f4d448>
- [23] B2SHARE data collection, Analytics 10 fold cross-validation (raw), Online: <http://hdl.handle.net/11304/163ba8e8-fe60-11e4-8a18-f31aa6f4d448>
- [24] B2SHARE data collection, Analytics 10 fold cross-validation (processed), Online: <http://hdl.handle.net/11304/5bba8e36-fe63-11e4-8a18-f31aa6f4d448>

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Selected Members of the Research Group on High Productivity Data Processing

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Mohammad Shahbaz Memon
Markus Goetz
Christian Bodenstein
Philipp Glock
Matthias Richerzhagen

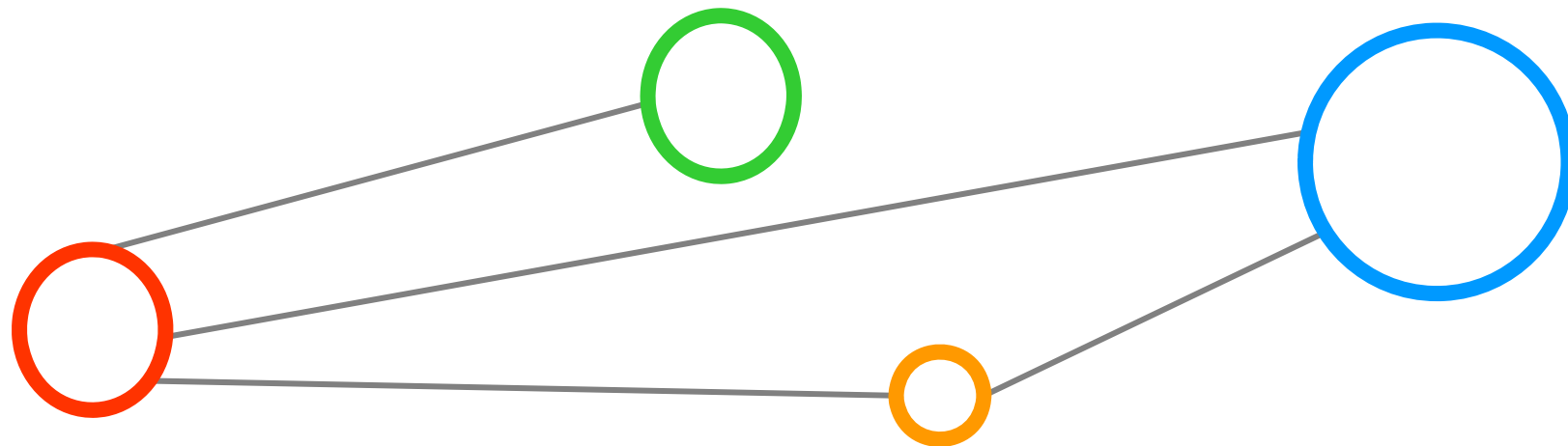


Thanks



Slides available at <http://www.morrisriedel.de/talks>

Selected Backup Slides for Discussions

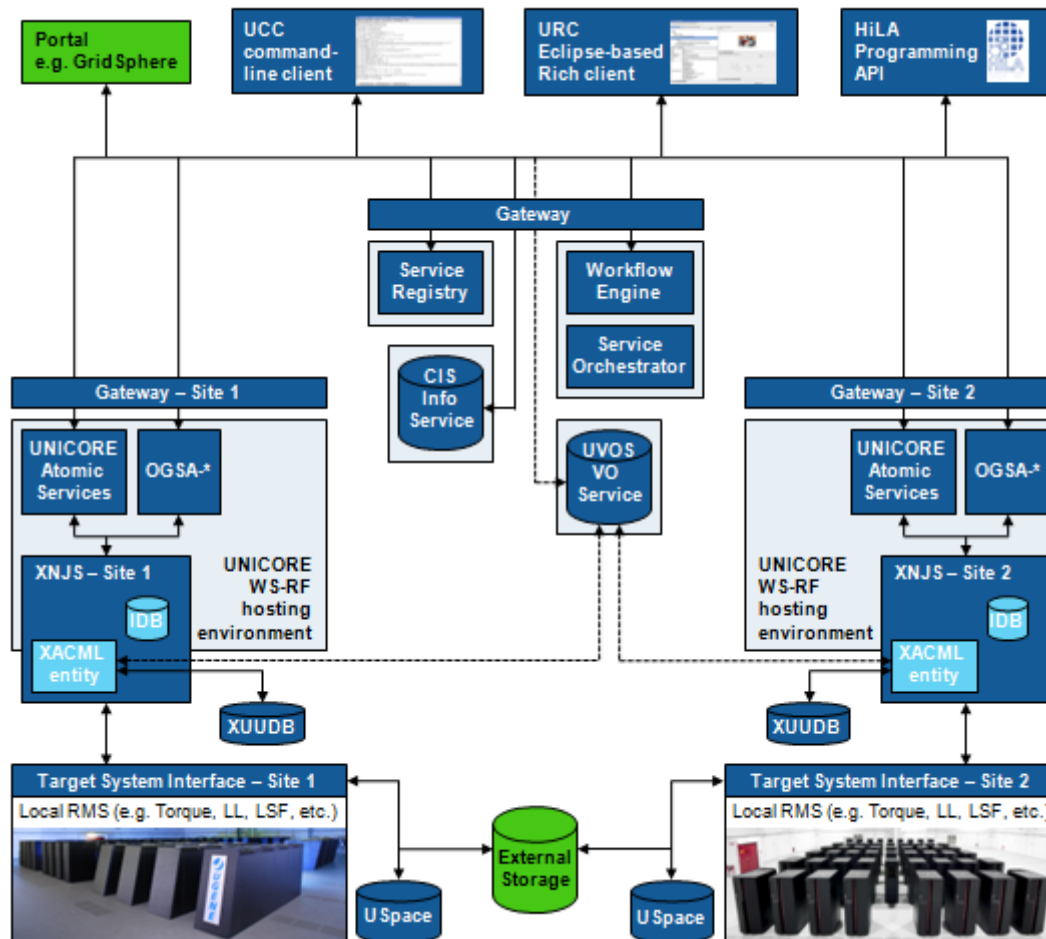


Distributed Large-scale Data Management

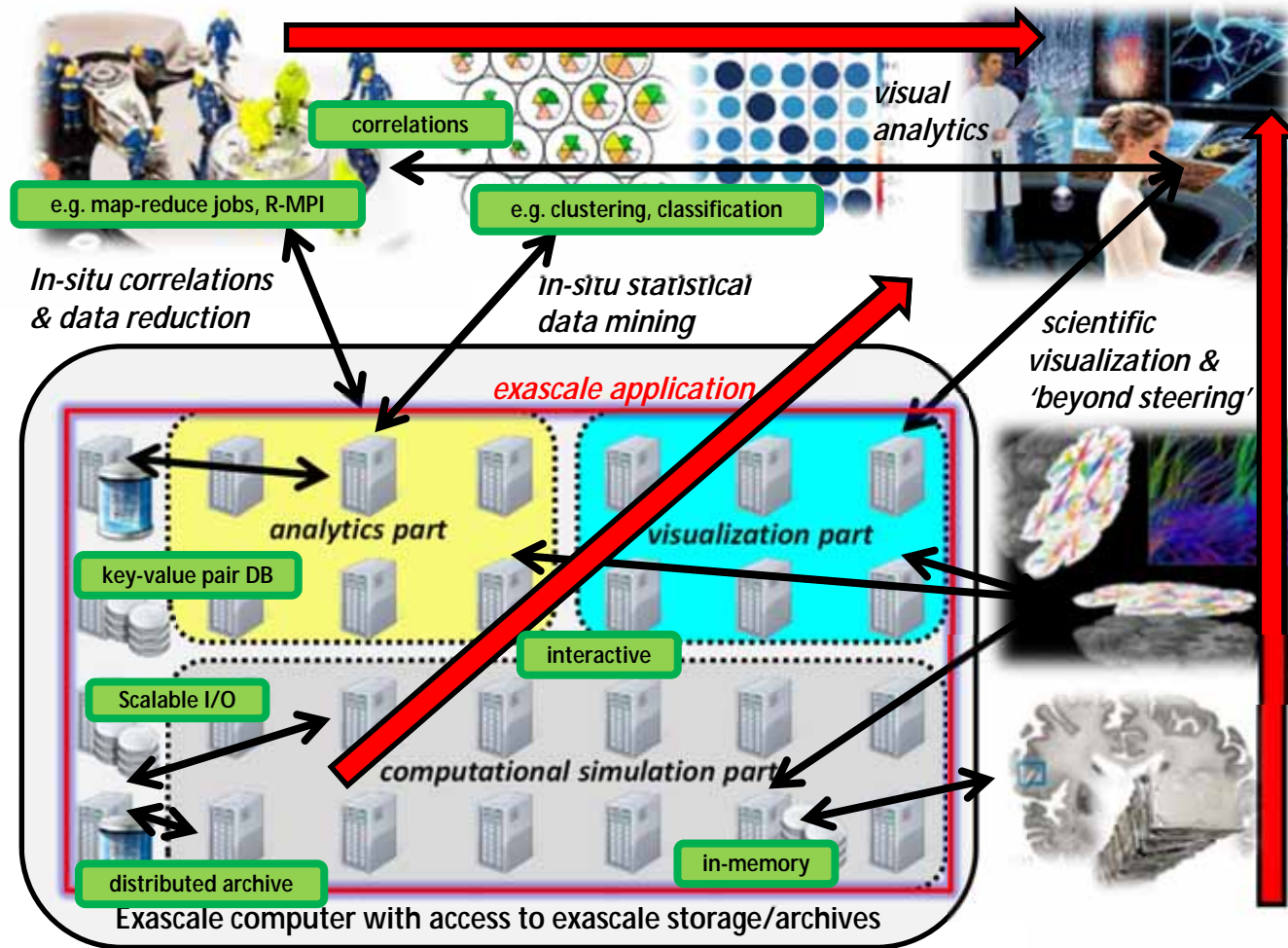


UNICORE

[14] UNICORE.eu

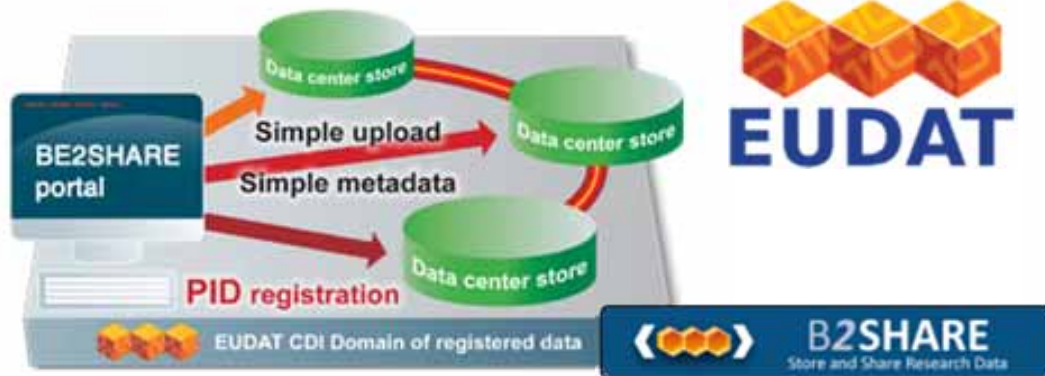


In-Situ Analytics for HPC & Exascale



[15] Inspired by ASCAC DOE report

Tools for Large-scale Distributed Data Management

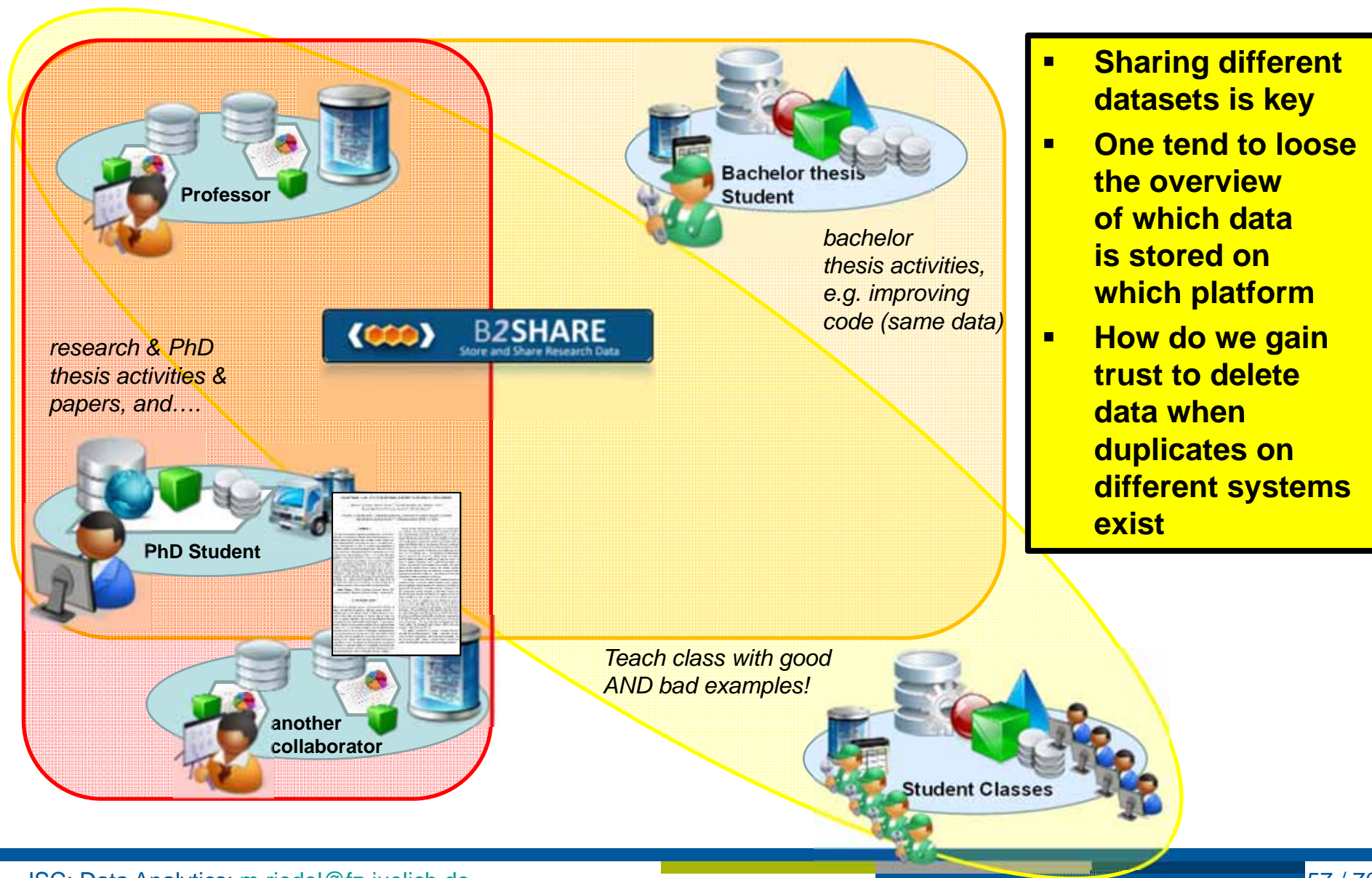


[16] M. Riedel & P. Wittenburg et al.

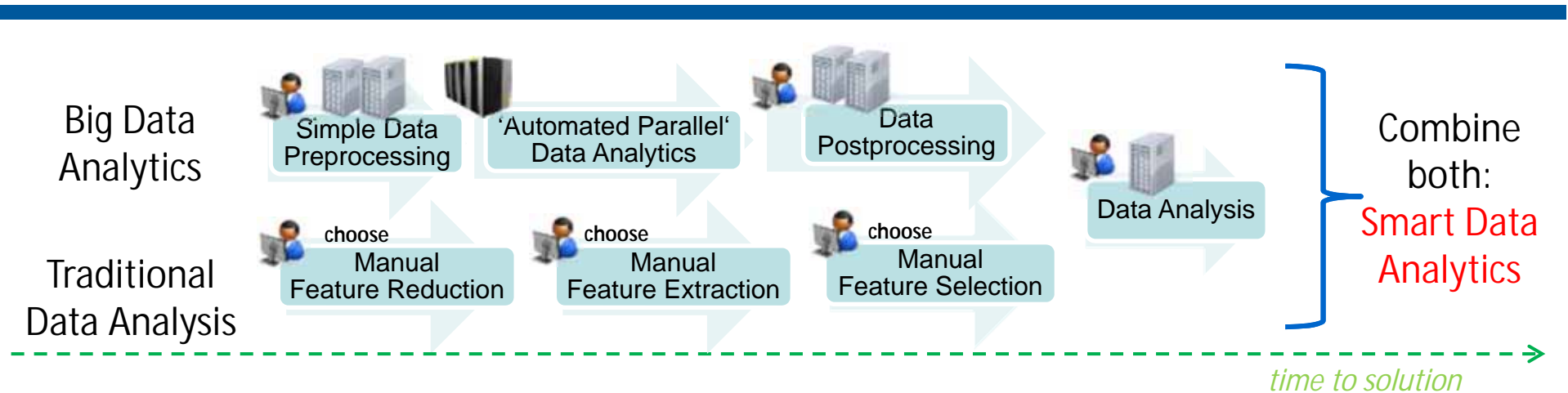
- Useful tools for data-driven scientists & HPC users



Need for Sharing & Reproducibility in HPC – Example



Smart Data Analytics Process



Concrete Datasets (& source/sensor)

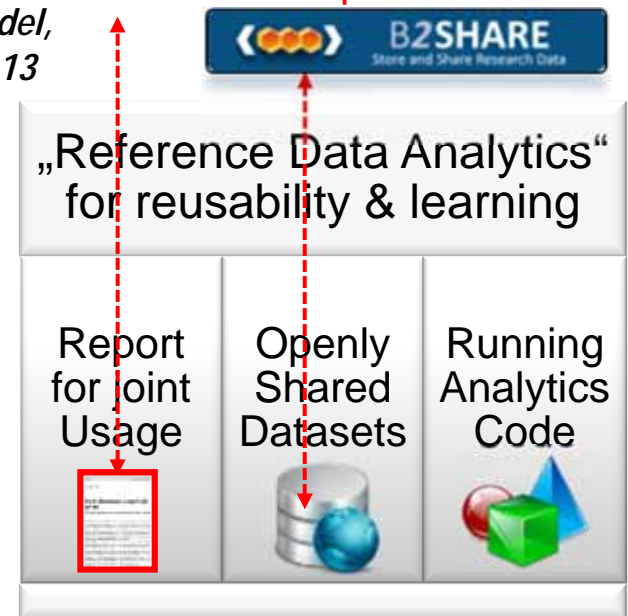
(parallel) Algorithms & Methods

Technologies & Resources

Scientific Data Applications

[17] C. Shearer, CRISP-DM model, Journal Data Warehousing, 5:13

CRISP-DM report

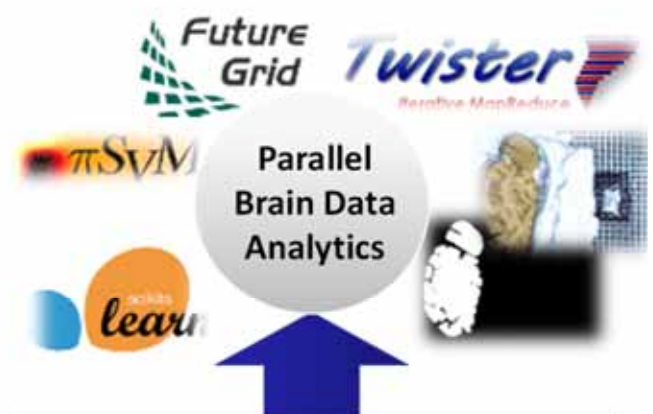


Selected Research Data Alliance (RDA) Activities

▪ **Big Data Analytics Interest Group – Establish something like UCI machine learning repository, but for big data analytics...**



[2] G. Cavallaro and M. Riedel et al. 'Smart Data Analytics Methods for Remote Sensing Applications', IGARSS 2014



Sattelite Data(Quickbird)

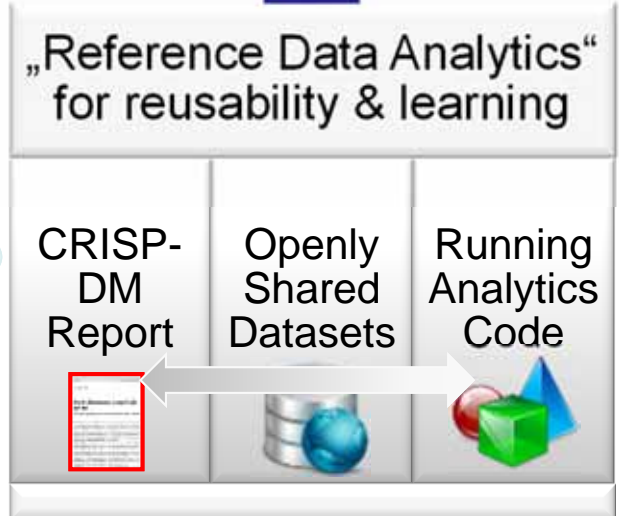
Parallel Support Vector Machines (SVM)

HPC & MPI

Classification Study of Land Cover Types

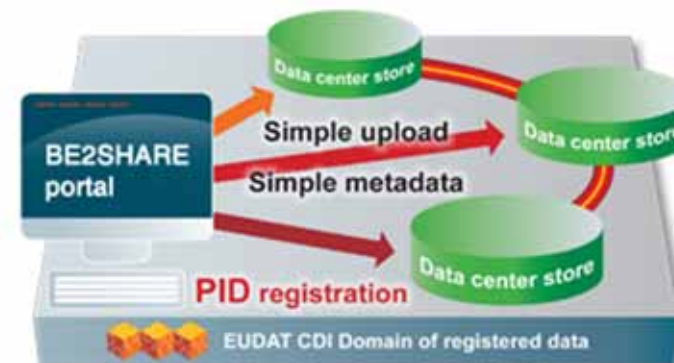
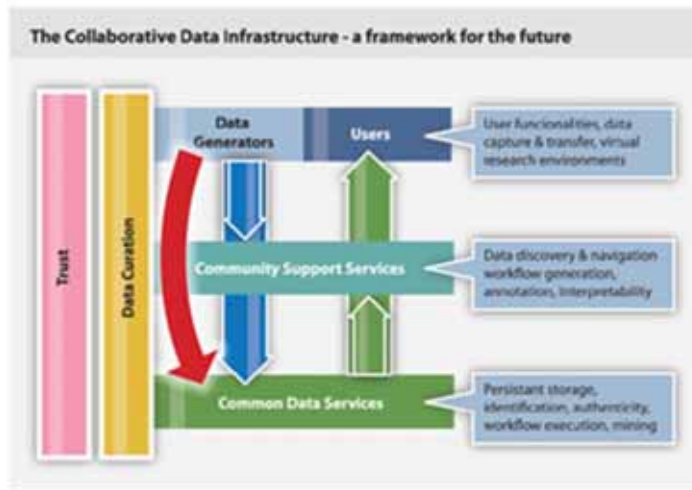
'Best Practices'

Classification++



➤ Research activities with Gabriele Cavallaro (PhD thesis, Uolceland) on Self Dual Attribute Profile

Reproducibility Example in Data-driven Science (1)



- Having this tool available on the Web helps tremendously to save time for no research tasks
- Using the tool enables to focus better on the research tasks

Reproducibility Example in Data-driven Science (2)

- Sharing pre-processed data
- LibSVM format
- Training and Testing Datasets
- Different setups for analysis (SDAP on All or SDAP on Panchromatic)

The screenshot displays the B2SHARE interface for a record titled "Rome data set OK". The page includes navigation tabs for "Information", "Comments", "Reviews", and "Usage statistics". The record details show it was created on 22 May 2014 and is available at <http://b2share.eudat.eu>. The abstract area is currently blank. Below the abstract, it states "The record appears in these collections: Generic".

The "Files" section contains a table with the following data:

| Name | Date | Size | Download |
|-----------------------------|-------------|----------|----------|
| sdap_area_panch_training.el | 22 May 2014 | 12.7 MB | Download |
| sdap_area_all_training.el | 22 May 2014 | 46.7 MB | Download |
| sdap_area_panch_test.el | 22 May 2014 | 114.8 MB | Download |
| sdap_area_all_test.el | 22 May 2014 | 420.0 MB | Download |

The "Export" section offers options to export the record as BibTeX, MARC, MARCXML, DC, EndNote, NLM, or RefWorks. The "Metadata" section provides the following information:

- PID:** <http://hdl.handle.net/11304/4615928c-e1a5-11e3-8cd7-14f6b57d12b9>
- Publication:** <http://b2share.eudat.eu>
- Publication Date:** 2014-05-22
- Uploaded by:** cavallero.gabriele@gmail.com
- Domain:** generic
- Checksum:** 15ba6c2e80c98859d0f9c044c73d1ec976b68c788e0681e3932d91e5a2a029c1

The browser address bar shows the URL <https://b2share.eudat.eu/record/86>.

Reproducibility Example in Data-driven Science (3)

Simple download from http using the wget command



```
mriedel@judge:~/bigdata> ls -al
total 640
drwxrwxrwx 21 mriedel zam 32768 2014-09-17 22:20 .
drwxr-xr-x 19 mriedel zam 32768 2014-09-18 11:49 ..
drwxr-xr-x 2 mriedel zam 32768 2014-06-19 07:17 102-salinasindian
drwxr-xr-x 2 mriedel zam 512 2014-06-19 20:11 107-salinasrescaled
drwxr-xr-x 2 mriedel zam 512 2014-07-08 17:14 111-romemultispectral
drwxr-xr-x 2 mriedel zam 512 2014-07-10 11:46 112-romeoriginalbands
drwxr-xr-x 2 mriedel zam 512 2014-09-17 22:31 120-indianpine
drwxr-xr-x 2 mriedel zam 512 2014-09-17 22:14 121-salinas
drwxr-xr-x 2 mriedel zam 512 2014-09-17 22:19 122-salinas2
drwxr-xr-x 2 mriedel zam 512 2014-09-17 22:24 123-indianpine2
drwxr-xr-x 2 mriedel zam 512 2014-07-09 11:03 86-romeok
drwxr-xr-x 2 mriedel zam 32768 2014-06-10 18:51 bigindianpines
drwxr-xr-x 2 mriedel zam 32768 2014-05-28 10:59 indian
drwxr-xr-x 2 mriedel zam 32768 2014-06-10 20:48 indianpinesreduced
drwxr-xr-x 2 mriedel zam 512 2014-07-28 17:53 mnist-576-rbf
drwxr-xr-x 2 skoehnen inml 512 2014-07-29 16:35 bli-blockface
drwxr-xr-x 2 mriedel zam 32768 2014-06-25 11:09 rome-ok
drwxr-xr-x 2 mriedel zam 512 2014-07-08 13:29 rome-ok-copy
drwxr-xr-x 2 mriedel zam 32768 2014-06-03 14:24 salinas
drwxr-xr-x 2 mriedel zam 32768 2014-06-16 16:50 salinasindianrev
drwxr-xr-x 2 mriedel zam 32768 2014-06-10 15:47 salinas-new
```

*...other
open
B2SHARE
datasets*

*...before adopting
B2SHARE regularly*

- Simple Download from http using wget
- Well defined directory structures

Reproducibility Example in Data-driven Science (4)

Make a short note in your directory linking back to B2SHARE

```
mriedel@judge:~/bigdata> cd 86-romeok/  
mriedel@judge:~/bigdata/86-romeok> ls -al  
total 580320  
drwxr-xr-x  2 mriedel zam      512 2014-07-09 11:03 .  
drwxr-xr-x 21 mriedel zam    32768 2014-09-17 22:20 ..  
-rw-r--r--  1 mriedel zam       35 2014-07-09 11:01 b2share.txt  
-rw-r--r--  1 mriedel zam 419974873 2014-05-22 13:36 sdap_area_all_test.el  
-rw-r--r--  1 mriedel zam 46652874 2014-05-22 13:36 sdap_area_all_training.el  
-rw-r--r--  1 mriedel zam 114763982 2014-05-22 13:36 sdap_area_panch_test.el  
-rw-r--r--  1 mriedel zam 12745692 2014-05-22 13:36 sdap_area_panch_training.el  
mriedel@judge:~/bigdata/86-romeok> more b2share.txt  
https://b2share.eudat.eu/record/86  
mriedel@judge:~/bigdata/86-romeok>
```

- Enables the trust to delete data if necessary (working against big data)
- Link back to B2SHARE for quick checks and file that links back fosters trust

Reproducibility Example in Data-driven Science (5)



```
mriedel@judge:~> ls -al
total 111840
drwxr-xr-x 19 mriedel zam      32768 2014-09-18 11:49 .
drwxr-xr-x 214 root    sys      32768 2014-09-12 09:02 ..
-rw-r--r-- 1 mriedel zam 113233920 2014-08-08 10:35 115-RunsMatthiasStable.tar ... a bachelor project
drwxr-xr-x 3 mriedel zam      32768 2014-06-03 16:24 ann-0.1
drwxr-xr-x 3 mriedel zam      32768 2014-06-03 17:02 ann-0.2
drwxr-xr-x 3 mriedel zam      32768 2014-06-04 14:42 ann-0.3
drwxr-xr-x 2 mriedel zam      32768 2014-06-16 19:12 ann-0.4
drwxr-xr-x 2 mriedel zam      32768 2014-06-16 19:24 ann-0.4-orig
drwxr-xr-x 2 mriedel zam      32768 2014-06-19 08:38 ann-0.5
drwxr-xr-x 6 mriedel zam      32768 2014-06-25 00:52 ann-0.6
drwxr-xr-x 4 mriedel zam      32768 2014-06-19 16:31 ann-0.6-scal
drwxr-xr-x 2 mriedel zam      32768 2014-06-24 17:02 ann-0.7
-rw----- 1 mriedel zam      1797 2014-05-12 13:51 .bashrc
drwxrwxrwx 21 mriedel zam      32768 2014-09-17 22:20 bigdata
drwxr-xr-x 3 mriedel zam      512 2014-06-19 09:34 .config
drwxr-xr-x 3 mriedel zam      32768 2014-06-03 14:38 .emacs.d
-rw----- 1 mriedel zam      1864 2014-05-12 13:51 .kshrc
drwxr-xr-x 3 mriedel zam      32768 2014-05-12 14:56 pivm-1.2
drwxr-xr-x 5 mriedel zam      32768 2014-09-18 11:49 pivm-1.2.1
drwxr-xr-x 3 mriedel zam      512 2014-07-09 14:51 pivm-1.2-refactored
-rw----- 1 mriedel zam      2686 2014-05-12 13:51 .profile
-rw----- 1 mriedel zam     22490 2014-09-18 11:51 .sh_history
drwx----- 2 mriedel zam      32768 2014-05-12 14:38 .ssh
drwxr-xr-x 2 mriedel zam      32768 2014-05-12 14:39 transfers
-rw----- 1 mriedel zam     19526 2014-09-18
-rw----- 1 mriedel zam      204 2014-09-17
mriedel@judge:~> █
```

... different versions of a parallel neural network code (another classification technique)

... different versions of a parallel support vector machine code

▪ True reproducibility needs: (1) datasets; (2) technique parameters (here for SVM); and (3) correct versions of algorithm code

Deep Learning (1)

Classical Machine Learning

Dealing with Big Data in traditional Machine Learning

- Define Features to learn from ?!
- Transform data into supported format ?!
- How to reduce dimensions ?!
- How to parallelize ?!



Deep Learning (2)

Deep Learning

Dealing with Big Data in Deep Learning

- Define Features to learn from
 - → Automatically learn how to define features
- Transform data into supported format
 - → Adopt the model to your data
- How to reduce dimensions
 - → Automatically reduce dimensions in every hidden layer
- How to parallelize
 - → Naturally the brain is parallel, so Artificial Neural Networks are!



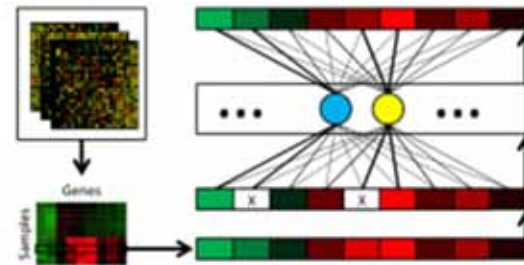
- A. Ng, Google Brain

Deep Learning (3)

Deep Learning in Computational Biomedicine

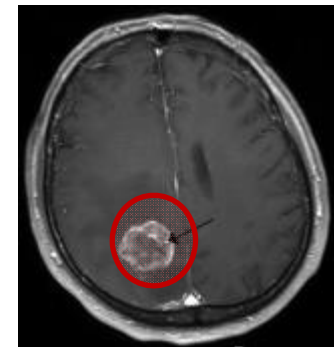
Genome Analysis

- Find high level features on low level –omics data



Medical Image Analysis

- Use 2D (or 3D) structure of the data for classification



Unstructured Data Analysis

- Use DL for text analysis to classify patient data, drug recommendations by users, ...



Etc...

Deep Learning (4)

Deep Learning Packages

There exists several frameworks for deep neural networks

- **Pylearn2**
 - Python tool on the top of the Theano python library
 - Easy configuration of data, model, learning via YAML files
 - CUDA support for accelerated calculations
 - Jobman for parallel cross validation
- **Caffe**
 - C++ implementation with python & matlab wrappers
 - CUDA acceleration
- **DL4J**
 - Java implementation of Deep Learning
 - CUDA + Hadoop support

Chances and Pitfalls for ‘Scientific Big Data Analytics’

~2009 – H1N1 Virus Made Headlines

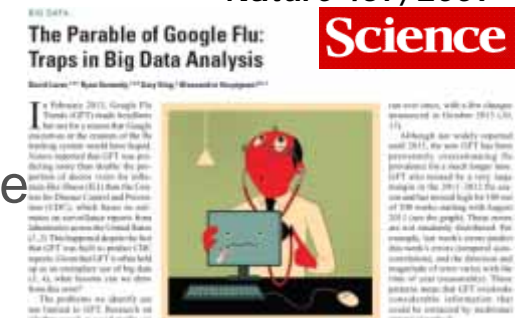
- Nature paper from Google employees
- Explains how Google is able to predict fast winter flus
- Not only on national scale, but down to regions
- Possible via logged big data – ‘search queries’



[18] Jeremy Ginsburg et al.,
'Detecting influenza epidemics
using search engine query data',
Nature 457, 2009

~2014 – The Parable of Google Flu

- Large errors in flu prediction & lessons learned
- (1) Dataset: Transparency & replicability impossible
- (2) Study the algorithm since they keep changing
- (3) It's not just about size of the data



[19] David Lazer, Ryan Kennedy,
Gary King, and Alessandro Vespignani,
'The Parable of Google Flu: Traps in Big Data Analysis',
Science Vol (343), 2014

■ Big data is not always better data – Think about difference of causality vs. correlation

Location-based Social Network-based Health Analytics

Scientific Domain Area

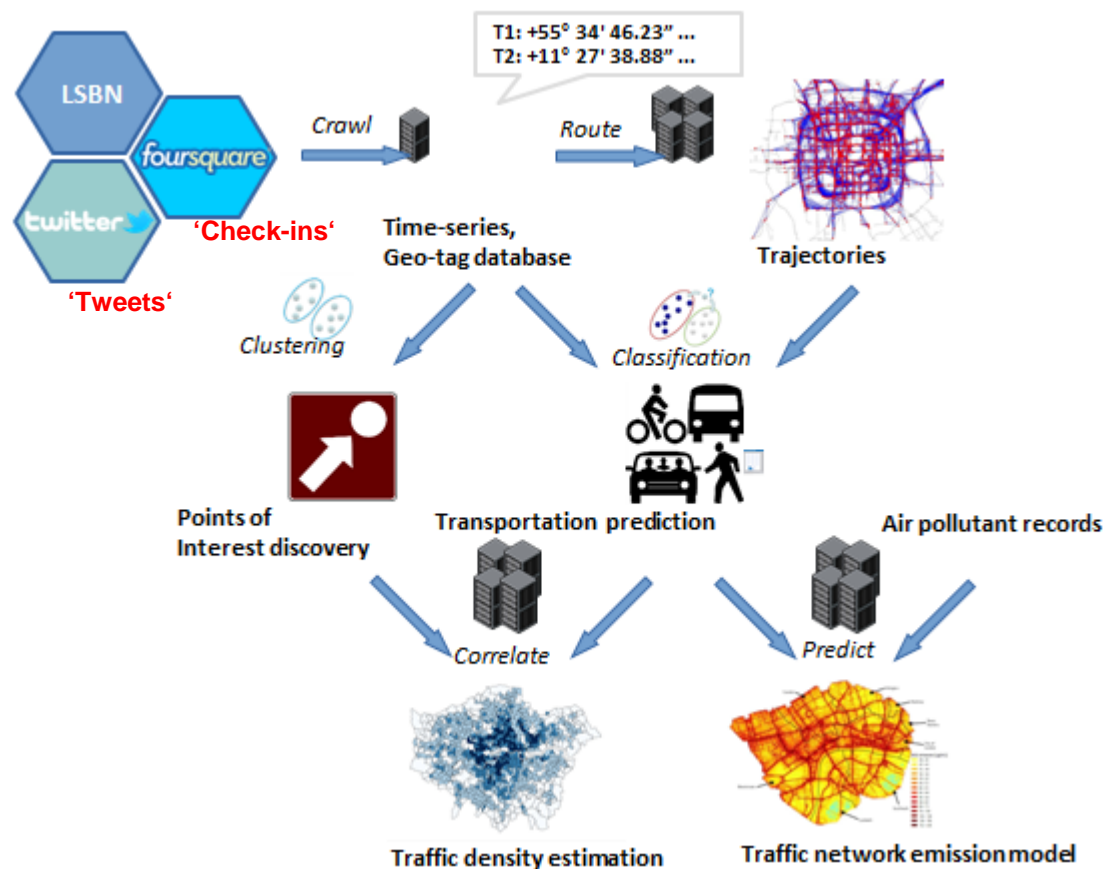
- Smart Cities approaches combined with Health Analytics Research

Scientific Outcome

- Traffic density estimation
- Network emission model

Location-based Social Networks (LBSN) Data

- Open data sources: Twitter & Foursquare
- Plan: Validation with real measurements in cities



➤ Research activities with Markus Goetz (PhD thesis) – Juelich Supercomputing Centre, Uolceland