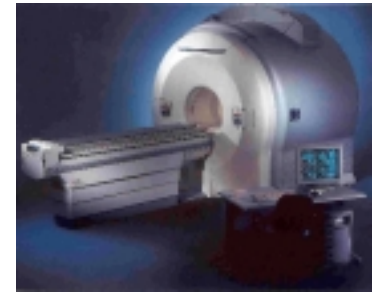


Introduction to Software Package

Andrzej MATERKA,

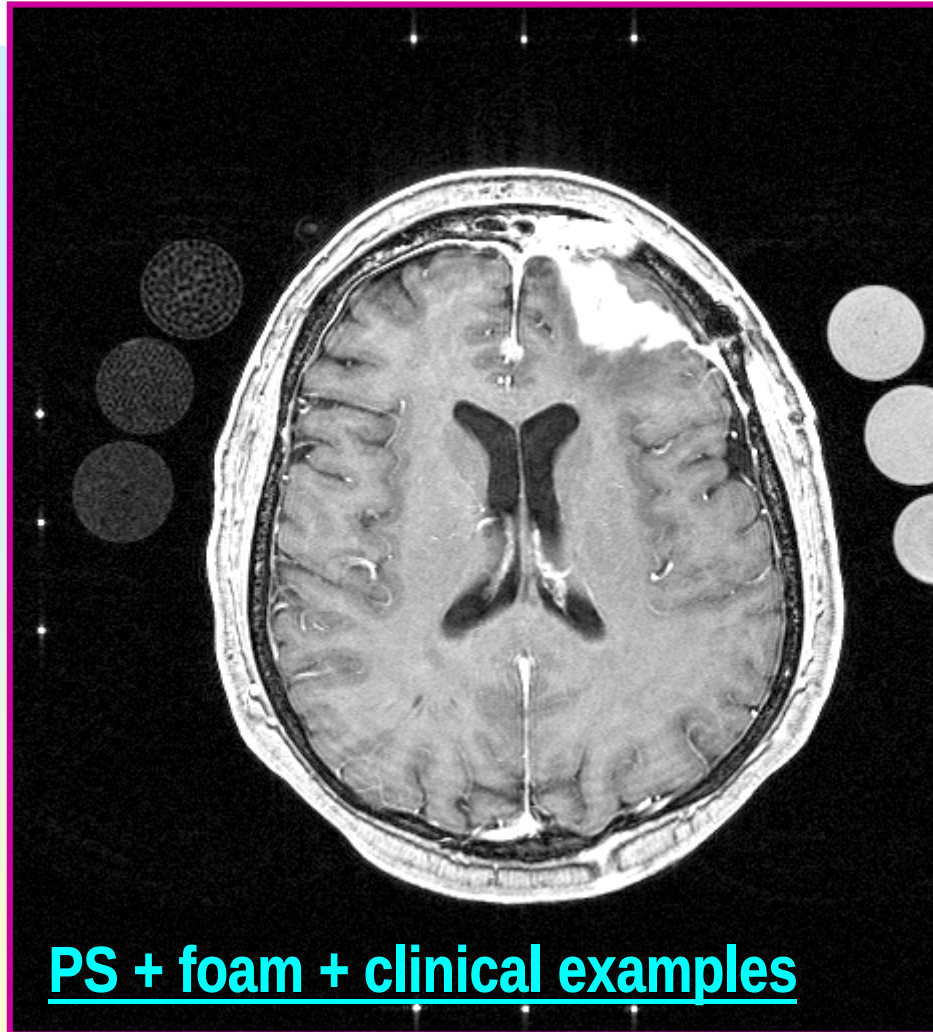
Technical University of Lodz, Poland



Goals

- To develop PC MS Windows computer programs with appropriate GUI and agreed set of functions.
- To develop efficient techniques of image processing and pattern recognition, adequate for MRI quantitative texture analysis.

Material (on CD)

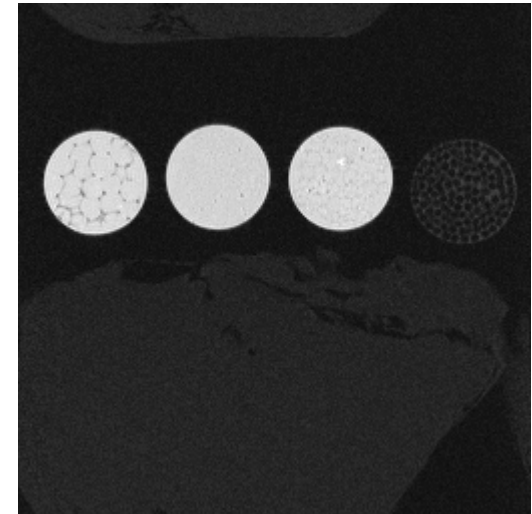


PS + foam + clinical examples

d:\brussels2000\lothar*.*		
Name	↑Ext	Size
↑...[...]		<DIR>
↑[clinical_examples]		<DIR>
↑[phantoms]		<DIR>

Name
↑...[...]
↑[733-0]
↑[733-6]
↑[733-7]
↑[733-8]
↑[735-57]
↑[735-6]
↑[735-61]
↑[735-62]
↑[735-63]
↑[735-64]
↑[737-0]
↑[737-2]
↑[737-4]
↑[737-5]
↑[737-6]
↑[763-0]
↑[763-2]
↑[763-3]
↑[763-4]
↑[771-0]
↑[771-59]
↑[771-60]

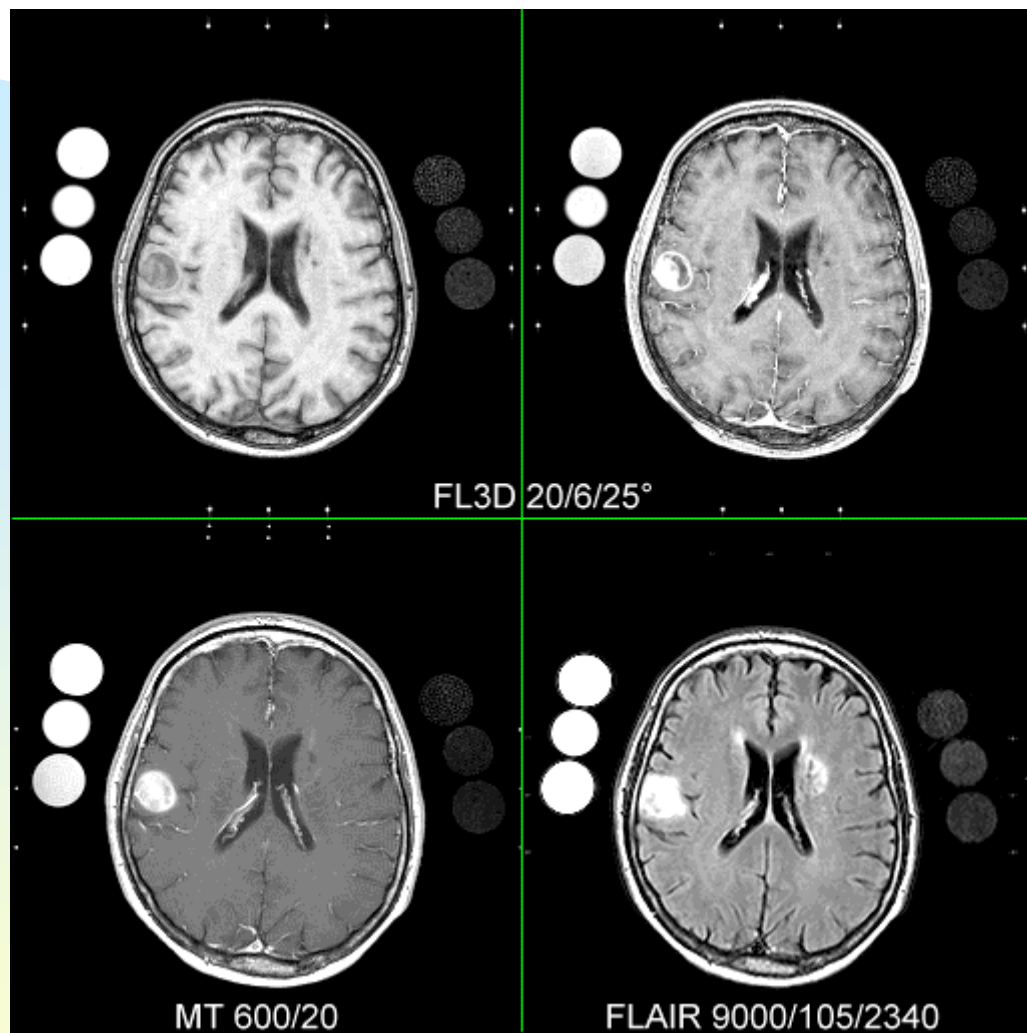
d:\brussels2000*.*		
Name	↑Ext	Size
↑...[...]		<DIR>
↑[hans]		<DIR>
↑[heidelberg]		<DIR>
↑[lothar]		<DIR>
↑[software]		<DIR>



Phantoms

Name
↑...[...]
↑[FOV100thick2]
↑[FOV100thick4]
↑[FOV200thick2]
↑[FOV200thick4]

Material (on CD)

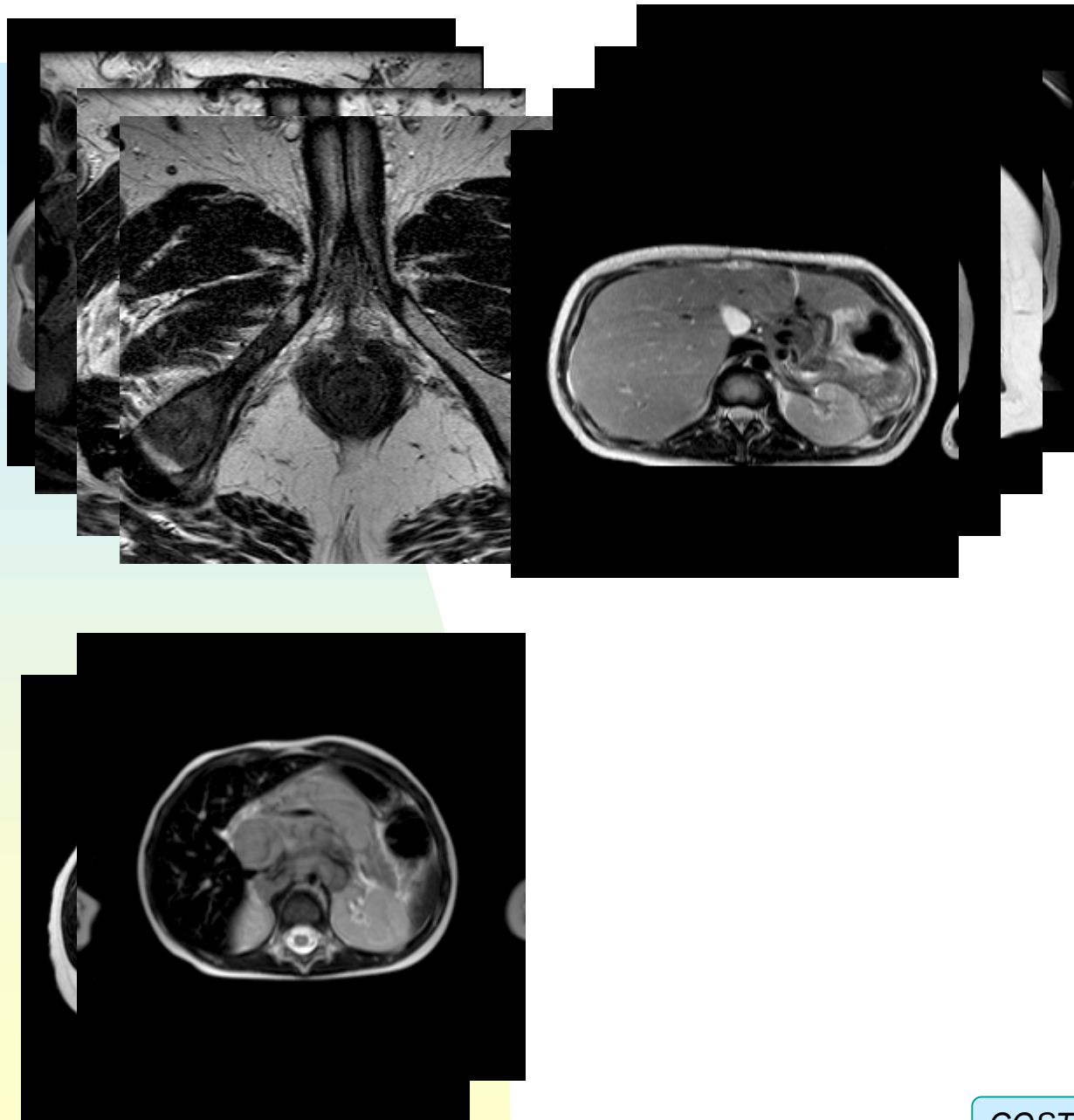


Patient data

d:\brussels2000*.*		
Name	↑Ext	Size
⚡...[.]		<DIR>
📁 [hans]		<DIR>
📁 [heidelberg]		<DIR>
📁 [lothar]		<DIR>
📁 [software]		<DIR>

Name	↑Ext	Size
⚡...[.]		<DIR>
📁 [clinical_examples]		<DIR>
📁 [phantoms]		<DIR>
📁 patient #1	eml	2 728 096
📁 patient #2	eml	2 916 439
📁 patient #3	eml	4 721 313
📁 patient #4(missed ima..	eml	729 197
📁 patient #4	eml	3 267 032
📁 patient #5	eml	2 728 015
📁 patient #6	eml	3 994 162
📁 patient #7 (last one)	eml	3 994 173

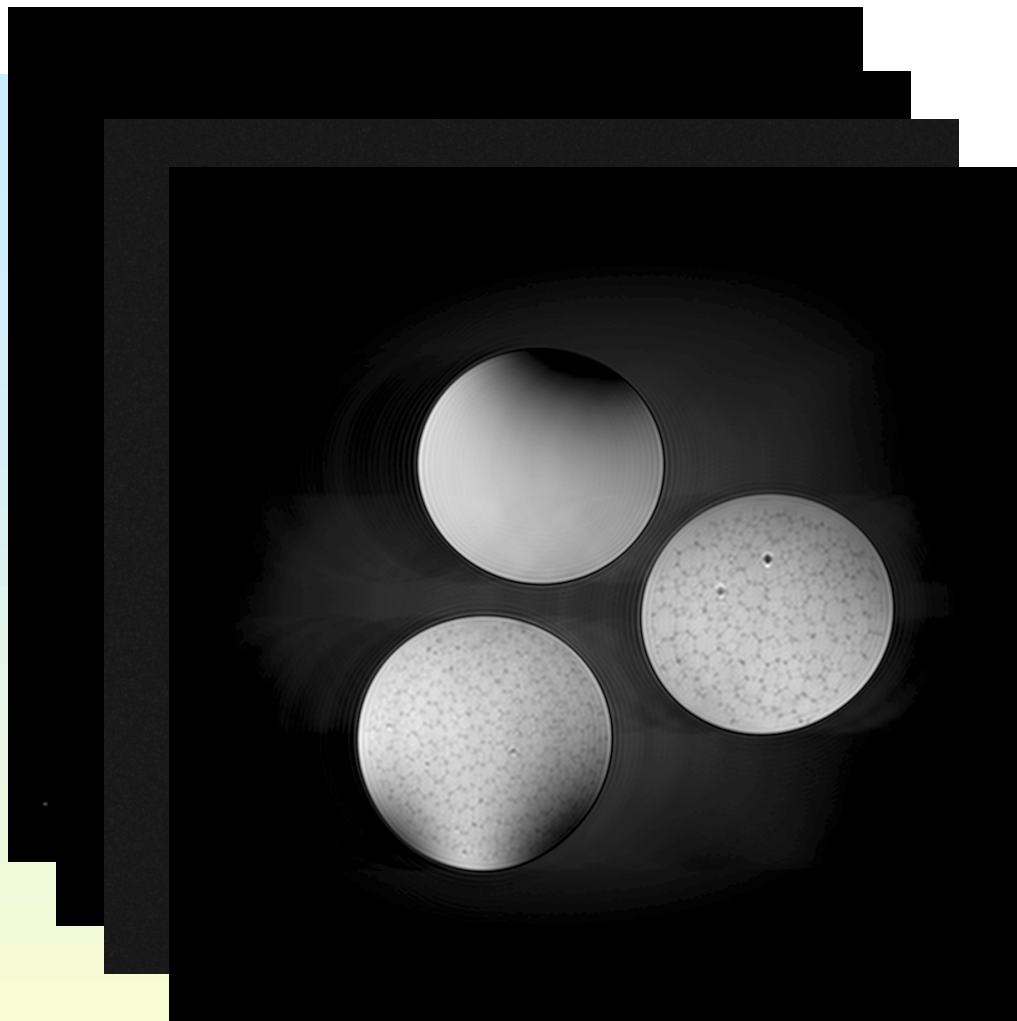
Material (on CD)



d:\brussels2000*. *		
Name	↑Ext	Size
↑...[.]		<DIR>
↑[hans]		<DIR>
↑[heidelberg]		<DIR>
↑[lothar]		<DIR>
↑[software]		<DIR>

d:\brussels2000\hans\aarhus_cost		
Name	↑Ext	Size
↑...[.]		<DIR>
↑[c_prostata.1]		<DIR>
↑[c_prostata.2]		<DIR>
↑[c_prostata.3]		<DIR>
↑[c_prostata.4]		<DIR>
↑[l_cirrous]		<DIR>
↑[l_cirrous_pancratitis.1]		<DIR>
↑[l_cirrous_pancratitis.2]		<DIR>
↑[l_cirrous_pancratitis.3]		<DIR>
↑[l_tomour]		<DIR>
↑[neuroblastom]		<DIR>
↑[phantoms]		<DIR>
↑[pregnant]		<DIR>
↑[ratimages]		<DIR>
↑readme	txt	1 401

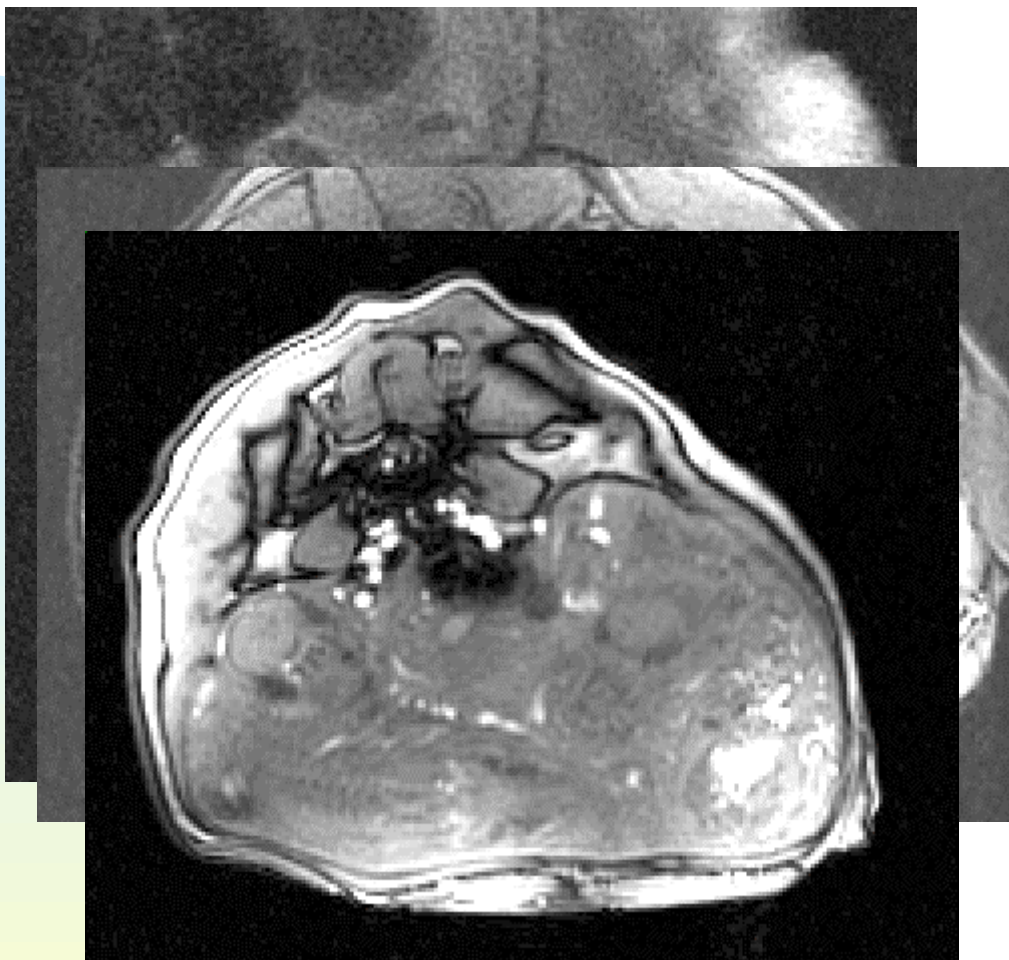
Material (on CD)



d:\brussels2000*.*		
Name	↑Ext	Size
↑...[.]		<DIR>
[hans]		<DIR>
[heidelberg]		<DIR>
[lothar]		<DIR>
[software]		<DIR>

d:\brussels2000\hans\aarhus_cost		
Name	↑Ext	Size
↑...[.]		<DIR>
[c_prostata.1]		<DIR>
[c_prostata.2]		<DIR>
[c_prostata.3]		<DIR>
[c_prostata.4]		<DIR>
[l_cirrous]		<DIR>
[l_cirrous_pancratitis.1]		<DIR>
[l_cirrous_pancratitis.2]		<DIR>
[l_cirrous_pancratitis.3]		<DIR>
[l_tomour]		<DIR>
[neuroblastom]		<DIR>
[phantoms]		<DIR>
[pregnant]		<DIR>
[ratimages]		<DIR>
readme	txt	1 401

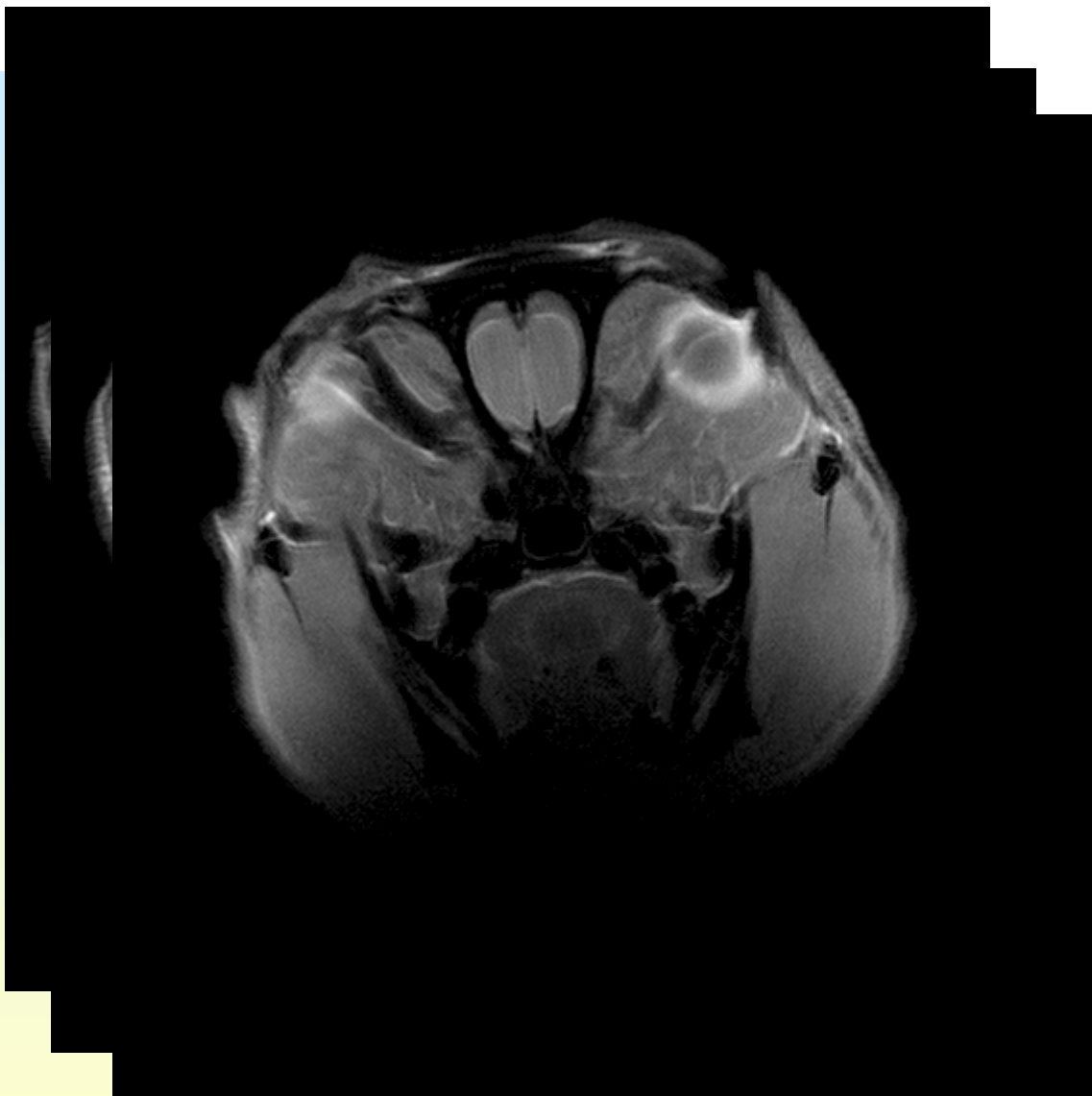
Material (on CD)



d:\brussels2000*.*		
Name	↑Ext	Size
↑...[..]		<DIR>
[hans]		<DIR>
[heidelberg]		<DIR>
[lothar]		<DIR>
[software]		<DIR>

d:\brussels2000\hans\aarhus_cost		
Name	↑Ext	Size
↑...[..]		<DIR>
[c_prostata.1]		<DIR>
[c_prostata.2]		<DIR>
[c_prostata.3]		<DIR>
[c_prostata.4]		<DIR>
[l_cirrous]		<DIR>
[l_cirrous_pancratitis.1]		<DIR>
[l_cirrous_pancratitis.2]		<DIR>
[l_cirrous_pancratitis.3]		<DIR>
[l_tomour]		<DIR>
[neuroblastom]		<DIR>
[phantoms]		<DIR>
[pregnant]		<DIR>
[ratimages]		<DIR>
readme	txt	1 401

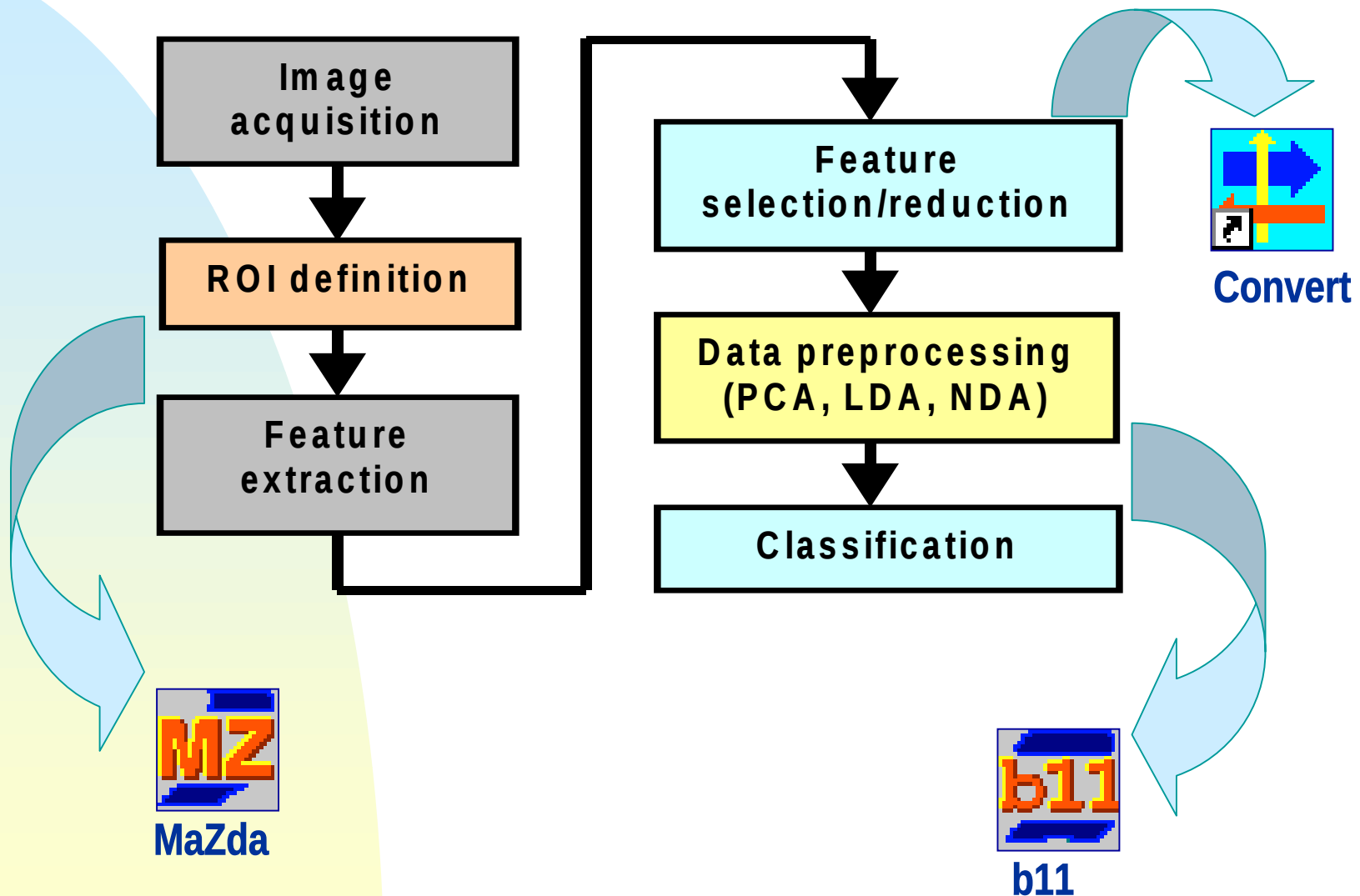
Material (on CD)

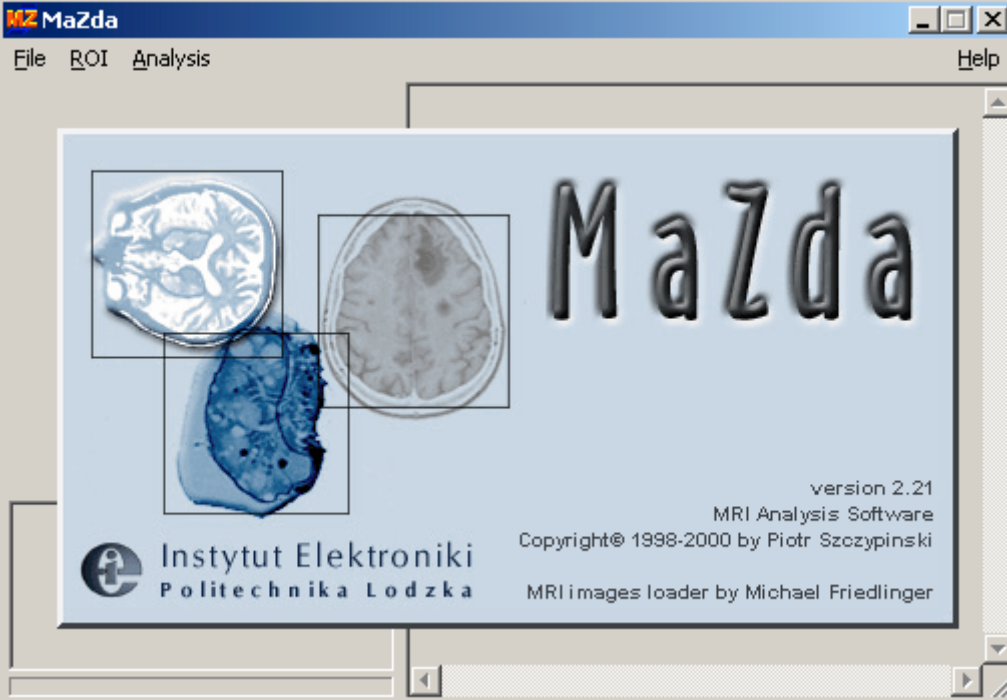


d:\brussels2000*. *		
Name	↑Ext	Size
↑...[..]		<DIR>
[hans]		<DIR>
[heidelberg]		<DIR>
[lothar]		<DIR>
[software]		<DIR>

d:\brussels2000\hans\aarhus_cost		
Name	↑Ext	Size
↑...[..]		<DIR>
[c_prostata.1]		<DIR>
[c_prostata.2]		<DIR>
[c_prostata.3]		<DIR>
[c_prostata.4]		<DIR>
[l_cirrous]		<DIR>
[l_cirrous_pancratitis.1]		<DIR>
[l_cirrous_pancratitis.2]		<DIR>
[l_cirrous_pancratitis.3]		<DIR>
[l_tomour]		<DIR>
[neuroblastom]		<DIR>
[phantoms]		<DIR>
[pregnant]		<DIR>
[ratimages]		<DIR>
readme	txt	1 401

Methods





MaZda, ver. 2.21

Feature extraction

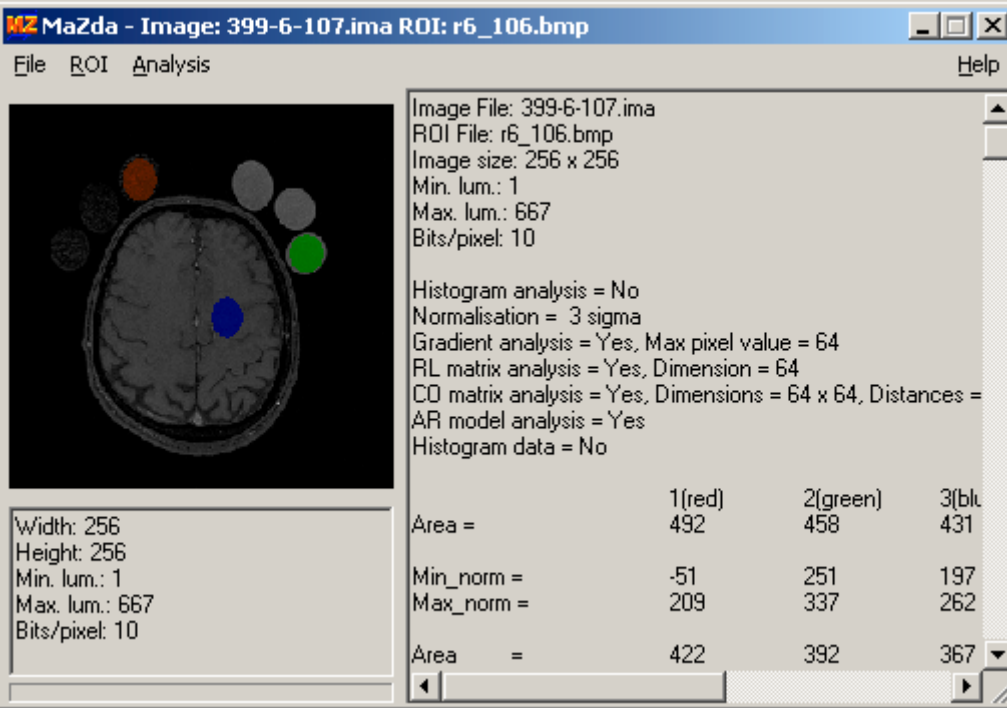
Input: images in 11 file formats

Methods

histogram, gradient, CO matrix,
AR model (total of 259 features)

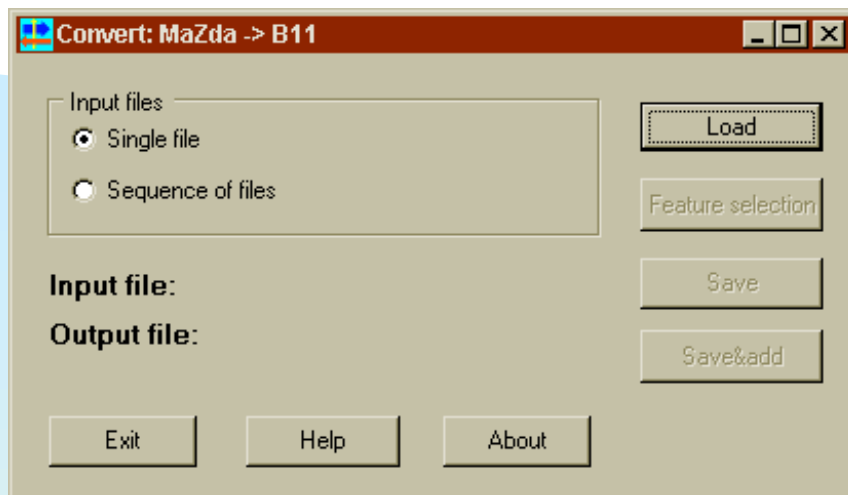
Output:

- (*.par) text file,
- (*.bmp) histograms
- (*.bmp) feature maps



Convert

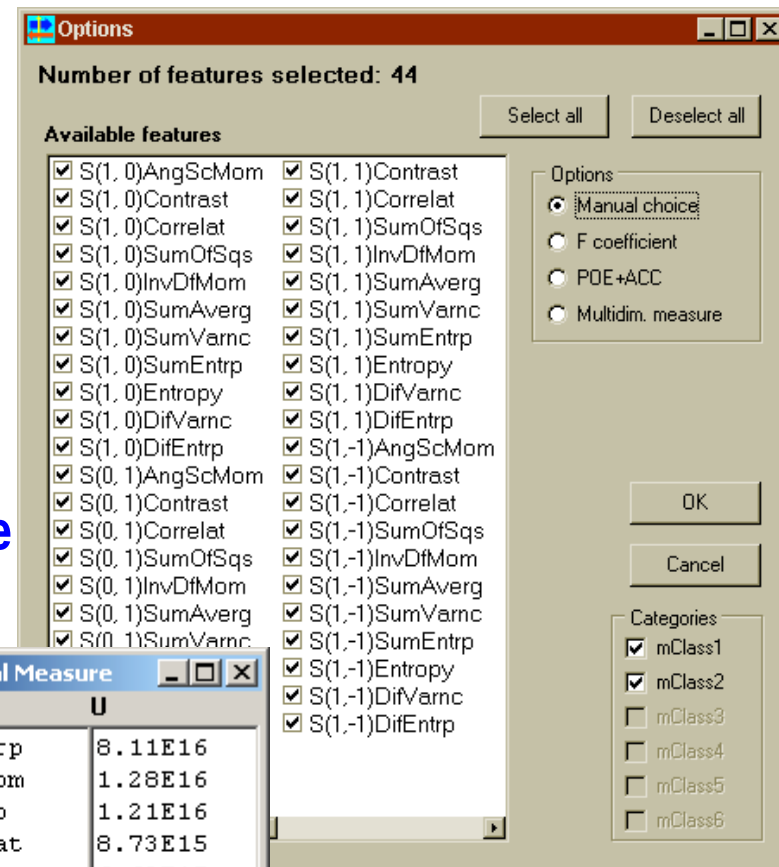
Feature selection



Input: (*.par) files from MaZda

Methods: Fisher coefficient, POE+ACC, multidimensional analysis of variance, manual choice

Output: (*.sel) files



Fisher coefficient

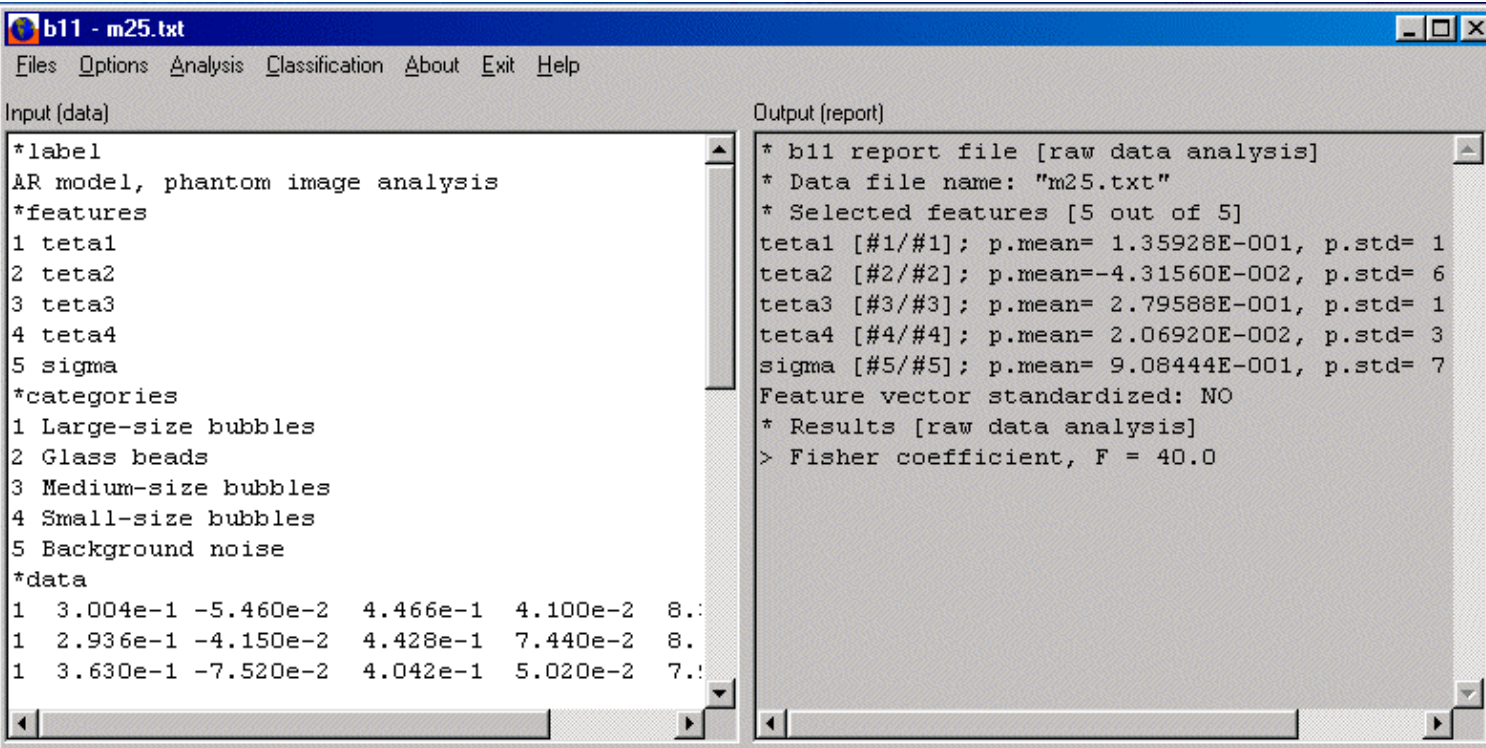
Feature name	F
S(1, 1)Correlat	44.00
S(0, 1)Correlat	43.52
S(1, 0)Correlat	37.59
S(1, 1)SumVarnC	36.19
S(0, 1)Contrast	35.72
S(1, 0)Contrast	33.66
S(1, 1)Contrast	33.01
S(1, 0)DifEntrp	31.47
Sigma	29.65
S(1, -1)Correlat	27.96

POE+ACC

Feature name	POE+ACC
GrMean	0.00
S(0, 1)SumVarnC	0.02
S(4, 4)DifVarnC	0.06
Teta4	0.10
S(0, 5)Correlat	0.10
S(4, -4)SumAverg	0.10
S(5, 0)DifVarnC	0.11
S(1, 0)DifEntrp	0.10
S(1, 0)SumVarnC	0.12
S(1, -1)InvDfMom	0.12

Multidimensional Measure

Feature name	U
S(0, 1)SumEntrp	8.11E16
S(0, 4)AngScMom	1.28E16
45dgr_ShrtrEmp	1.21E16
S(4, -4)Correlat	8.73E15
S(4, 0)Entropy	6.63E15
S(0, 4)DifEntrp	6.42E15
S(5, -5)InvDfMom	4.63E15
S(0, 4)InvDfMom	3.43E15
S(4, 0)AngScMom	3.38E15
S(1, -1)SumEntrp	3.24E15



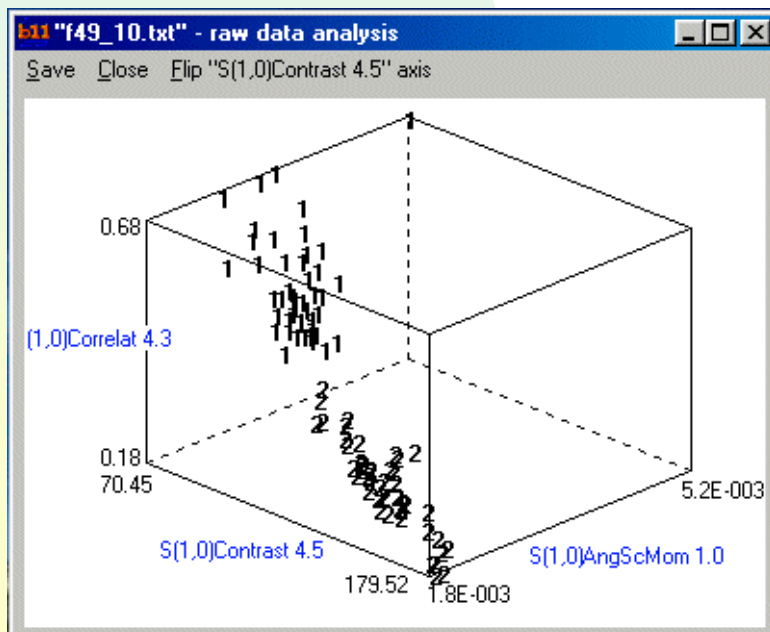
B11

Input

- (*.sel) files

Output

- text files,
- BMP graphs



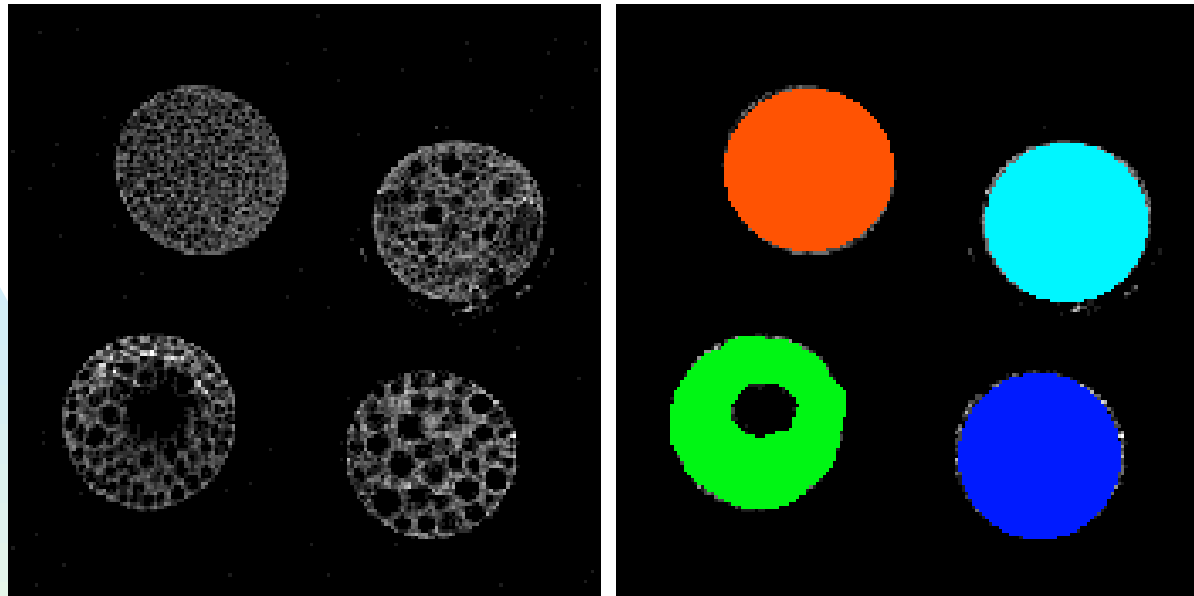
**Analysis,
classification**

Methods

- data analysis: raw, PCA, LDA, NDA
- classification: k-NN, neural network

Example - PS phantom image analysis

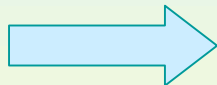
22 images



- class 1
- class 2
- class 3
- class 4



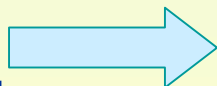
MaZda



22 (*.par) files, each contains 259 texture parameters



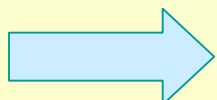
Convert



2 (*.sel) files, each of 10 parameters (F, POE)



b11

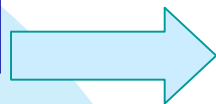


Exploratory data analysis and classification

Example - PS phantom image analysis

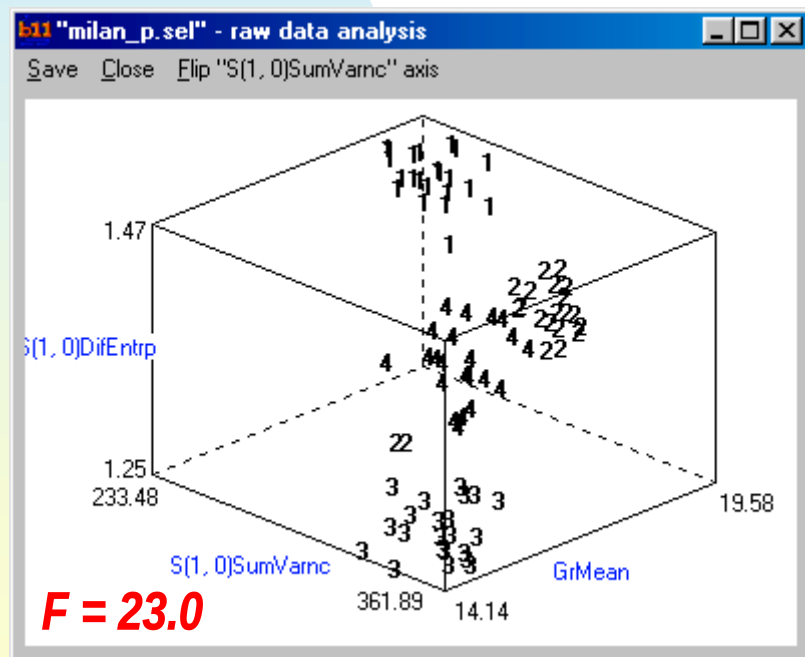


b11



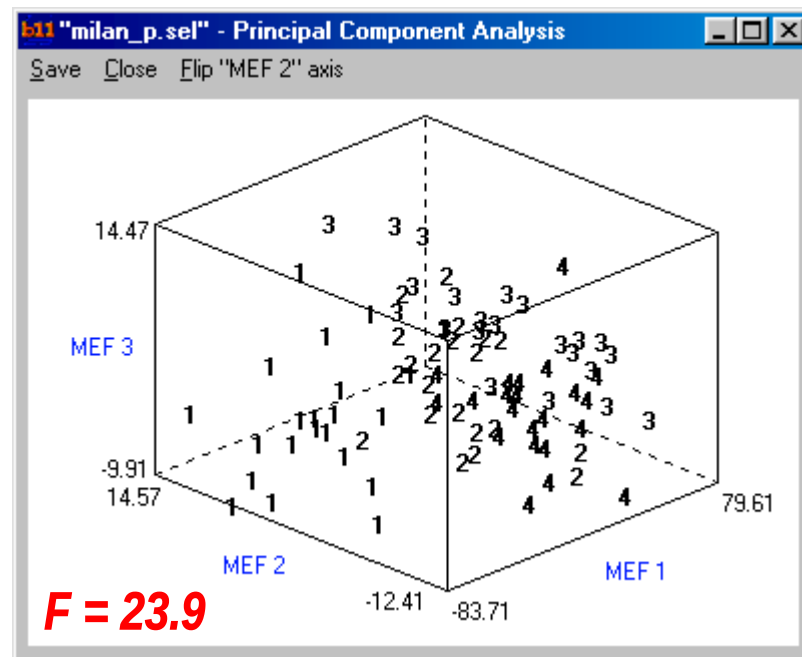
Exploratory data analysis and classification

Raw data



1-NN: 8 classification errors

First 3 PCAs



1-NN: 10 classification errors

Example - PS phantom image analysis

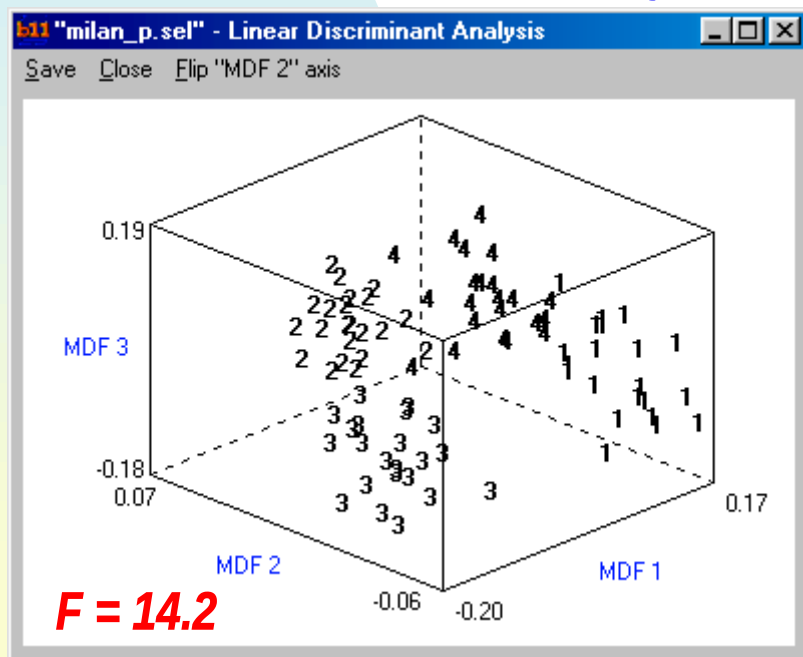


b11



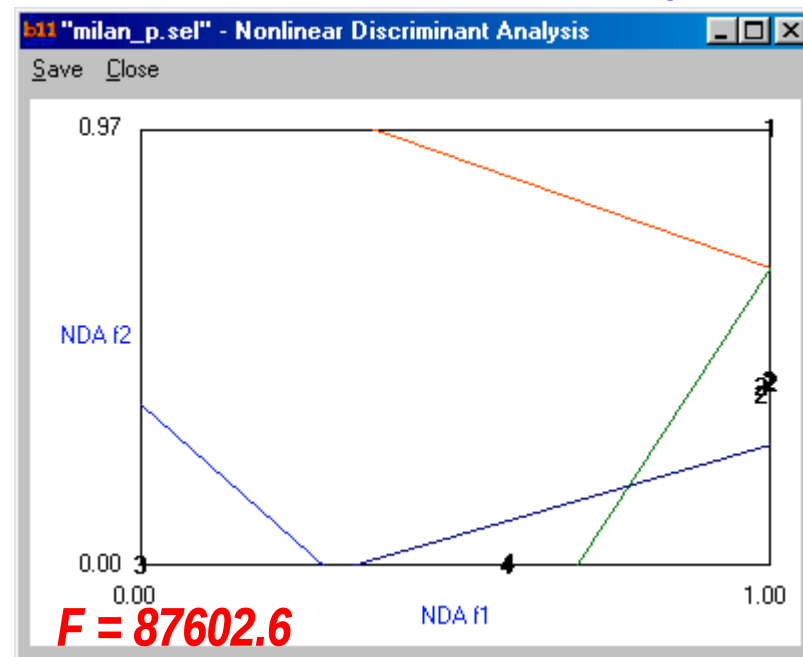
Exploratory data analysis and classification

Linear Discriminant Analysis



1-NN: 0 classification errors

Nonlinear Discriminant Analysis



0 classification errors

Projects under way

- MaZda: ROI editor (to replace MS Paint). 
- MaZda: new texture features (**wavelets** and **mathematical morphology**).
- MaZda: reading header of image files, supporting new image file standards.
- MaZda: image brightness scaling (optionally manual/auto).
- MaZda: integration of tools (ROEdit, BMPView).
- Convert: feature reduction and selection techniques.
- B11: testing on separate data sets (PCA, LDA, ANN).
- Extensive experiments using test object and biological MRI.