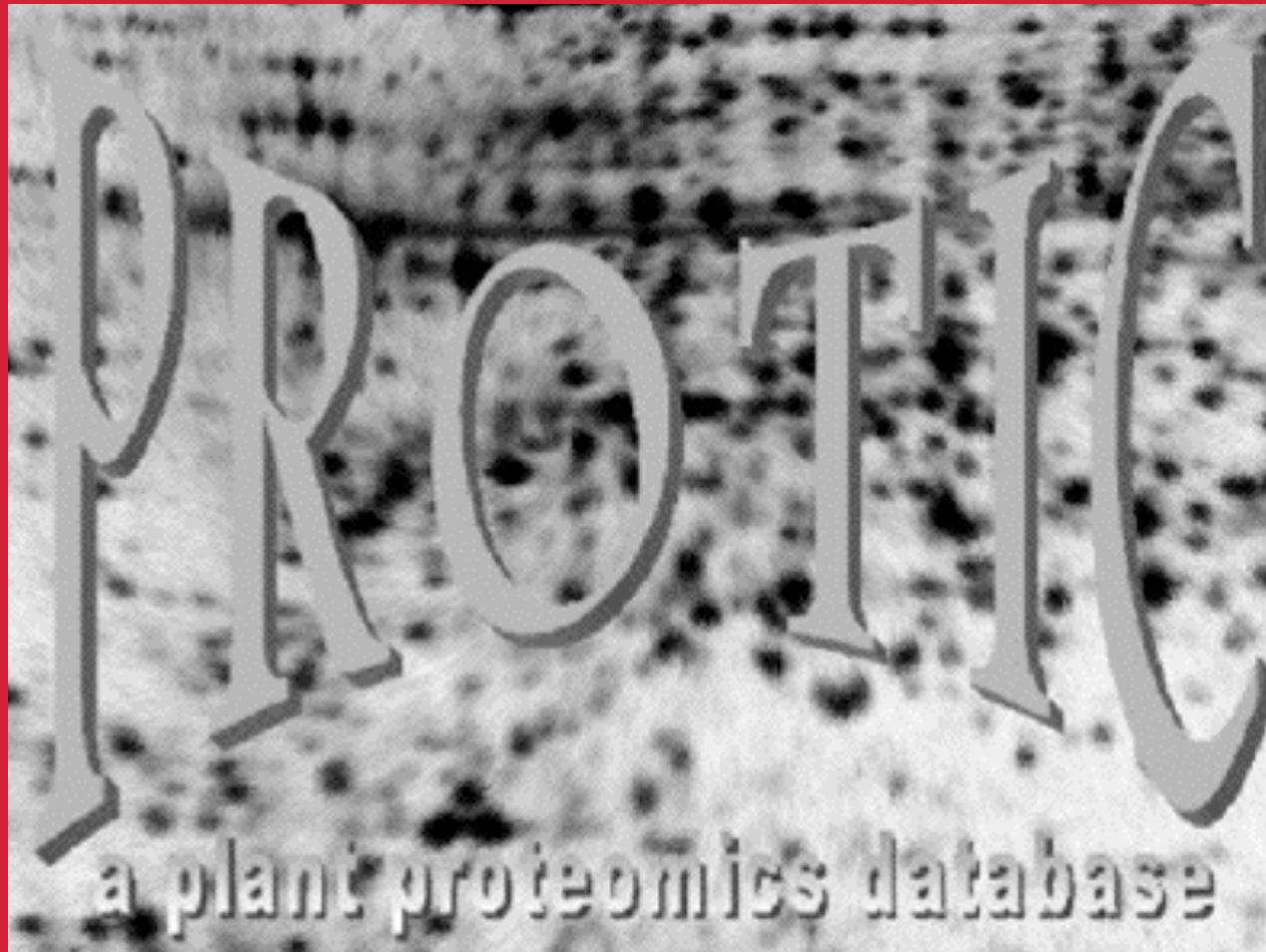
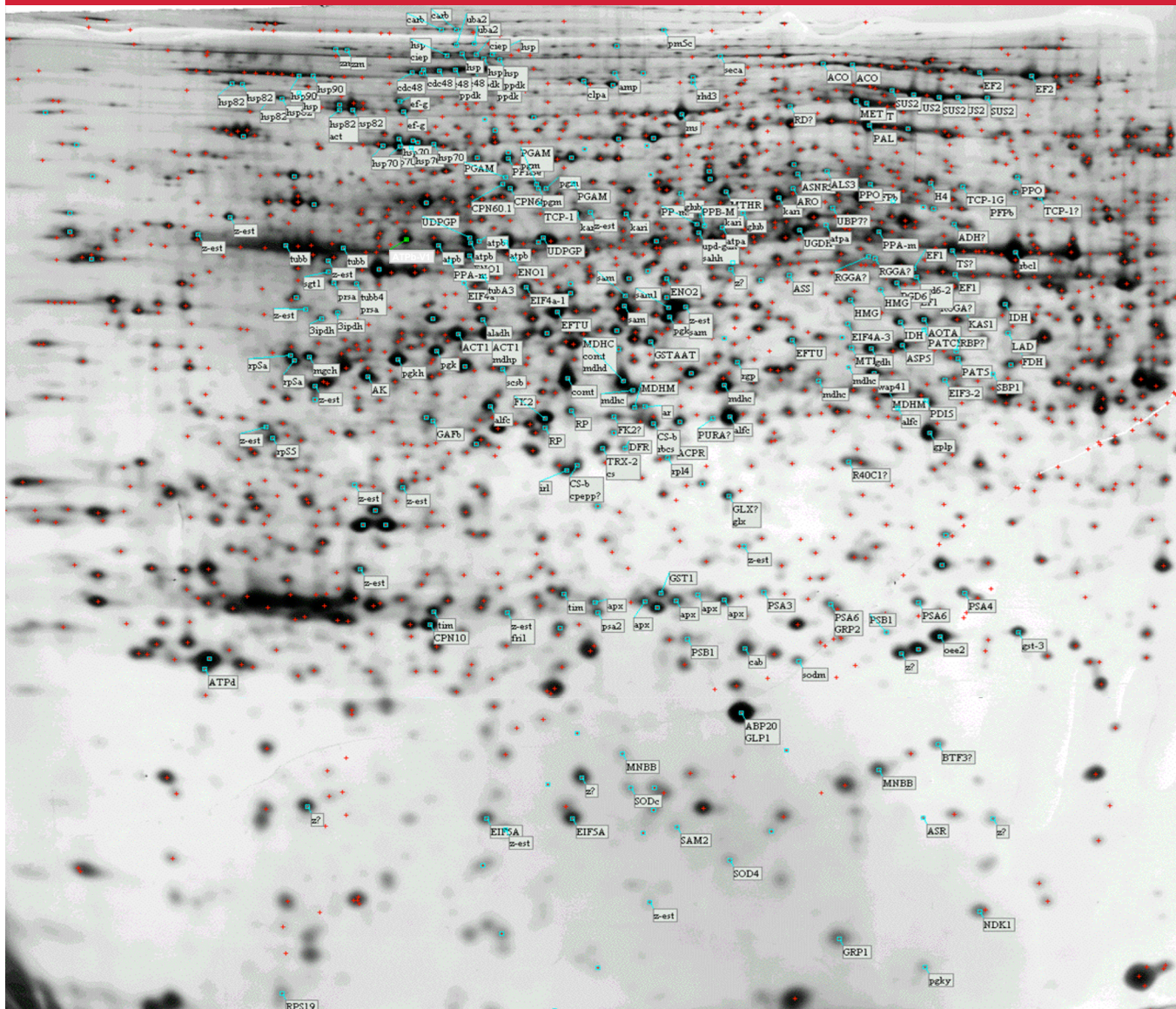


PROTICdb : a PROteomic database



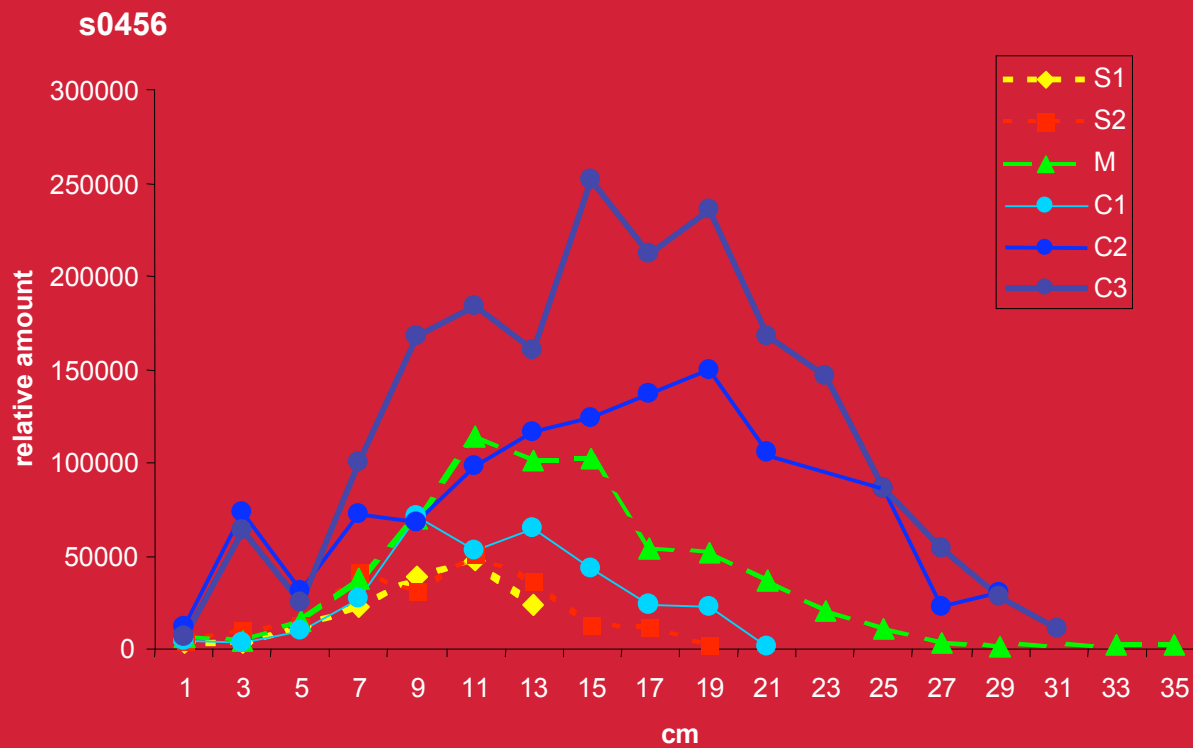
Some proteomic results on maize leaves



Protein
identification

720 MS analyses
80% identifications

Some proteomic results on maize leaves



Drought stress

- Control
- Mild deficit
- Strong deficit

Several stages

- growing leaves
- mature leaves

Different leaf areas

- elongation zone
- differentiation zone

Different projects

On different targets:

drought

kernel filling

ploidy,

wood formation

On different species:

Maize

Rape, Cabbage

Arabidopsis

Pine

On different organs:

leaves

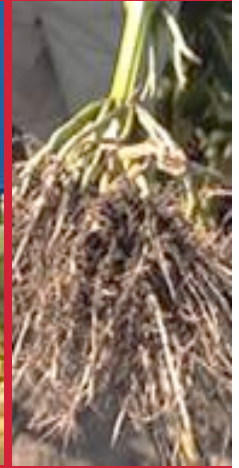
seeds

roots

stem (wood)



2n
4n



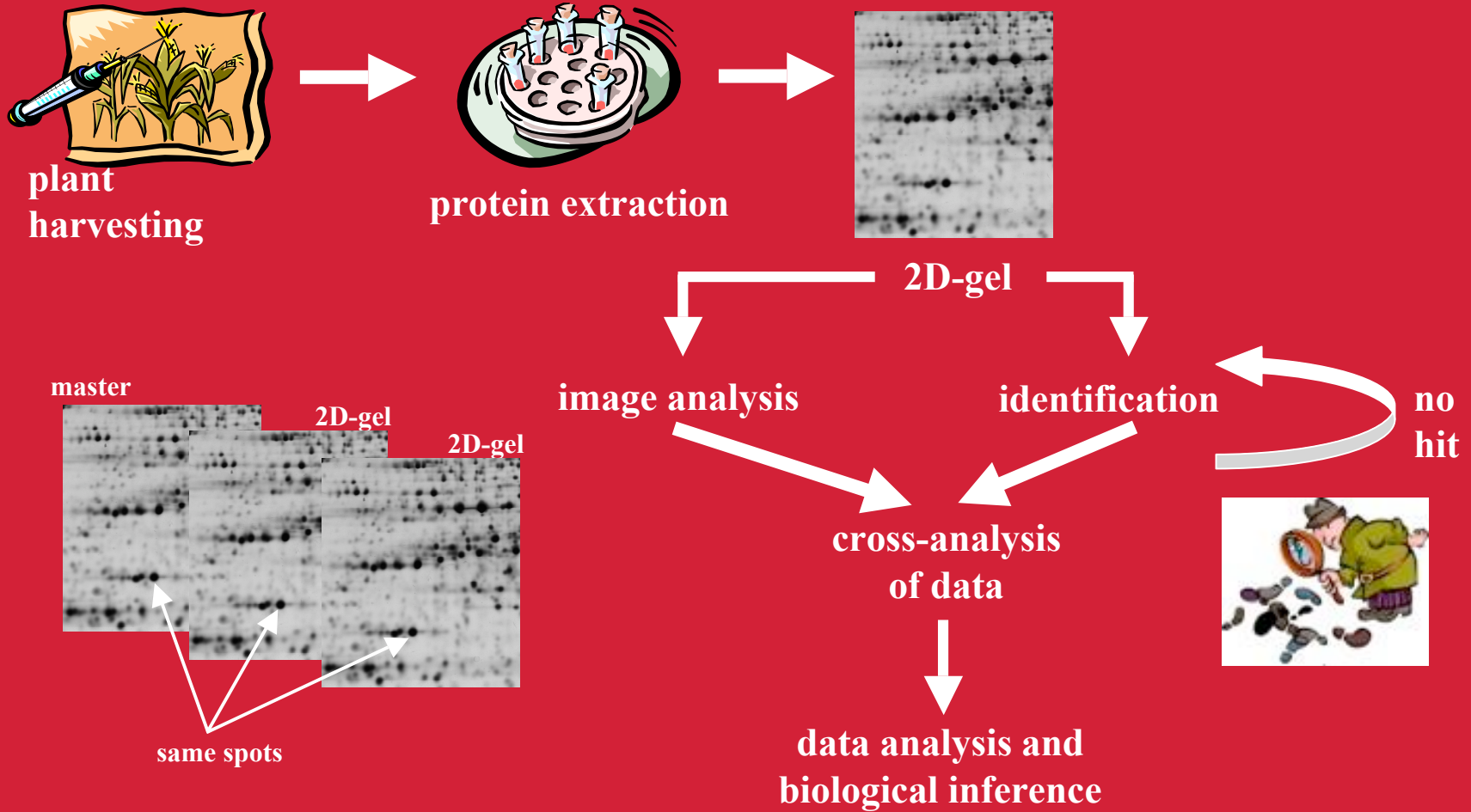
Objectives

Highly heterogeneous to store in an organized way and to manage in order to extract information.

The objectives are :

- 1- Data storing, management, visualization and mining
- 2- Data export to analysis software
- 3- Knowledge management
- 4- Web publication

Experimental data flow



Data structuring

SpotGel : spot detected with a specific detection method on an image and bearing an unique number in a given number system.

Spot 35

Spot : group of SpotGels with the same number in a given number system.

SpotGel 35

SpotGel 125

SpotGel 198

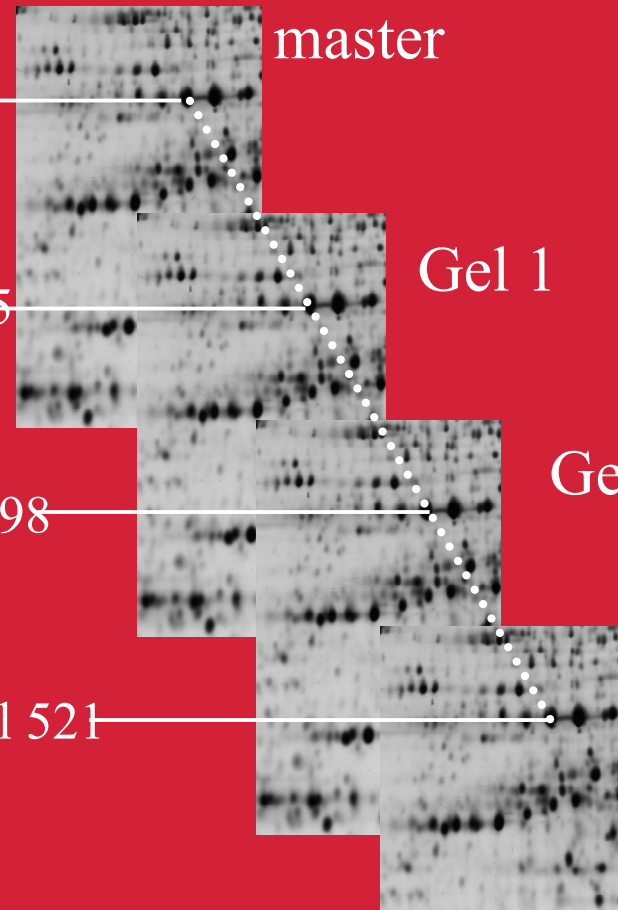
SpotGel 521

master

Gel 1

Gel 2

Gel n ..



Data structuring

same spot

post-translational mod.

phosphorylation

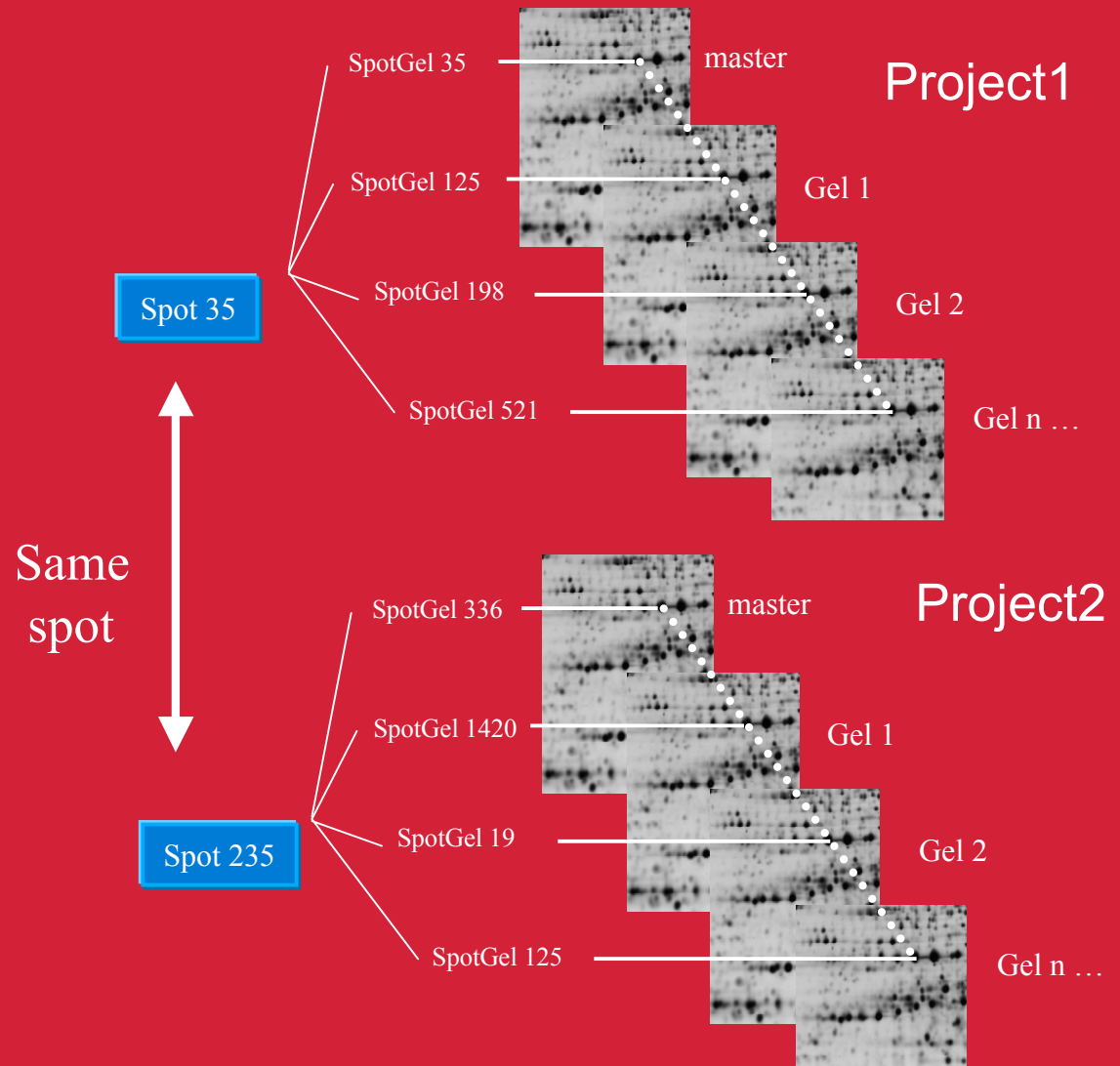
glycosylation

other post. mod.

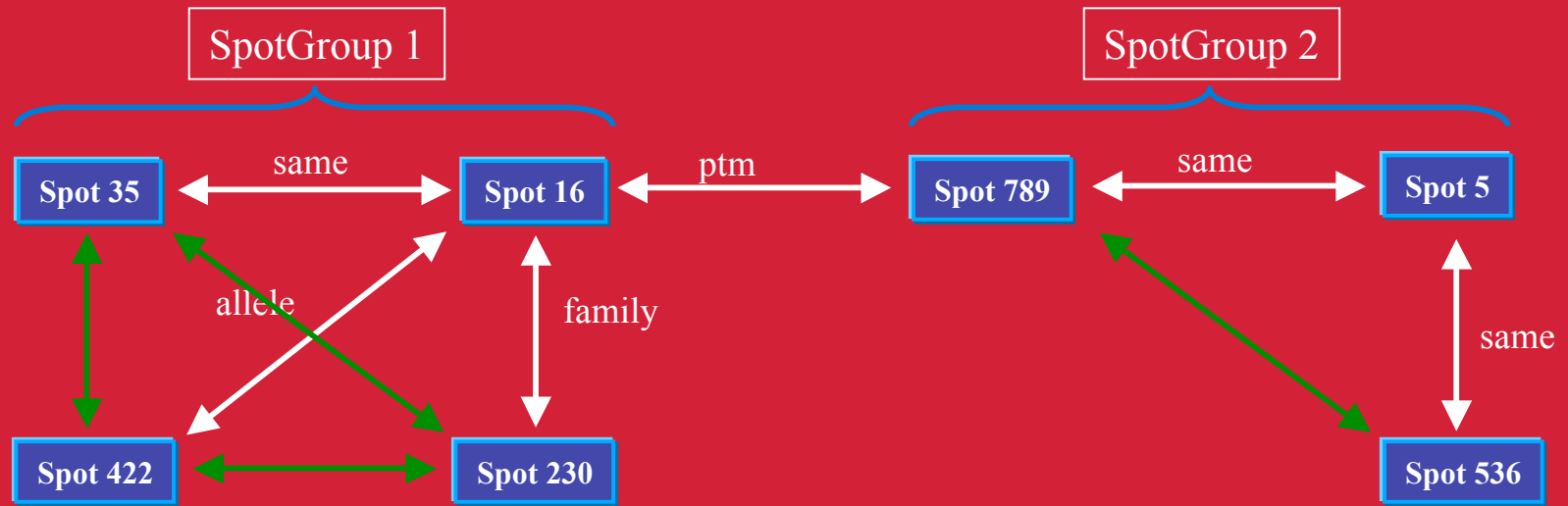
allelic form

isoform

multigenic family



Data structuring

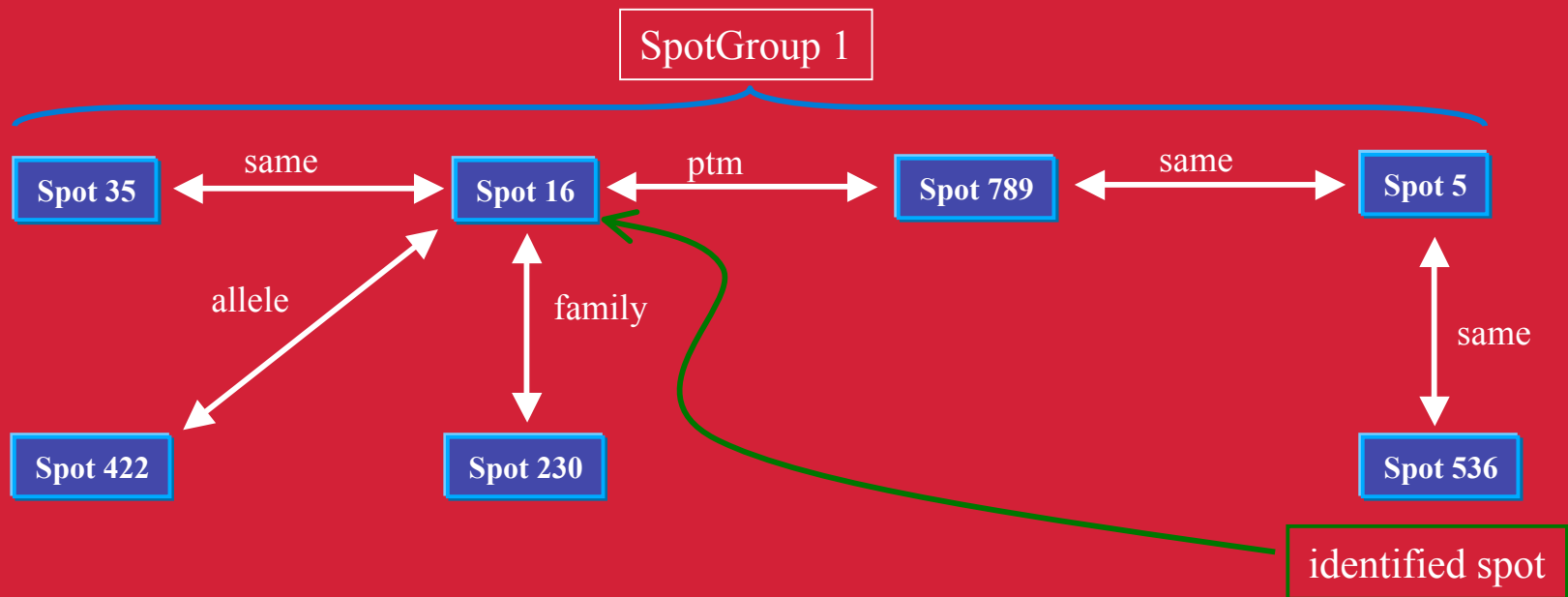


↔ automatically inferred relationships

The relationships are evidenced by the users.

The SpotGroups are created by the database.

Data structuring



The SpotGroups are up-dated by the database.

The information is spread from one spot to another.



[Plant](#) [Plant sample](#) [Protein sample](#) [Gel](#) [Gel image](#)

User: THIERRY
Project: PLANT
RESPONSE TO
DROUGHT

- Logout
- Help
- Project selector
- Administration

- Controlled vocabulary
- Methods

- Gel browser
- Form based feeding
- File based feeding

- Credits



Plant sample update

plant sample :

Plant sample update

Sample name*

Plant

plant
(population/number):

Harvesting stage

plant stage:

organ :

rank definition:

rank number:

date:

Data status

public private

Comment



User: THIERRY
Project PLANT
RESPONSE TO
DROUGHT

Logout

Help

Project selector

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Controlled vocabulary

Methods

Gel browser

Form based feeding

File based feeding

Credits

New sample organ reference

[Plant](#) [Plant sample](#) [Protein sample](#) [Gel](#) [Gel image](#)

[numerisation calibration](#) [concentration unit](#) [picture format](#) [quantity unit](#)

[protein sample quality](#) [harvesting stage](#) [rank definition](#) [sample organ](#)

[tissue](#) [scanner name](#) [detection software](#) [scanner software](#)

[subcellular compartment](#) [genotype](#) [plant test](#) [spot numerotation system](#)

- 6TH LEAF
- 8TH LEAF
- ADULT LEAF
- ADULT STEM
- EAR LEAF
- GRAIN
- [create a new reference for sample organ](#)

Create a reference for sample organ

sample organ reference name :





[plant2image \(csv file\)](#) [virtual spot detection \(update\)](#) [spot detection \(new/update\)](#)

[MS identification request \(in\)](#) [MS identification request \(out\)](#) [identification results](#)

User: THIERRY
Project: PLANT
RESPONSE TO
DROUGHT

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Project selector

Administration

Controlled vocabulary

Methods

Gel browser

Form based feeding

File based feeding

Credits

Plant to image file upload

[here is the excel template file](#)

Choose your
experiment :

CHOU RC HAPLOIDE, DIPLOIDE ET TETRAPLOIDE

new/update

Select your file :

Parcourir...

Submit

Gel image file upload

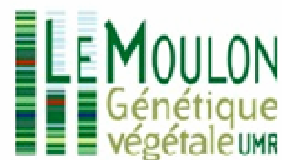
image file format

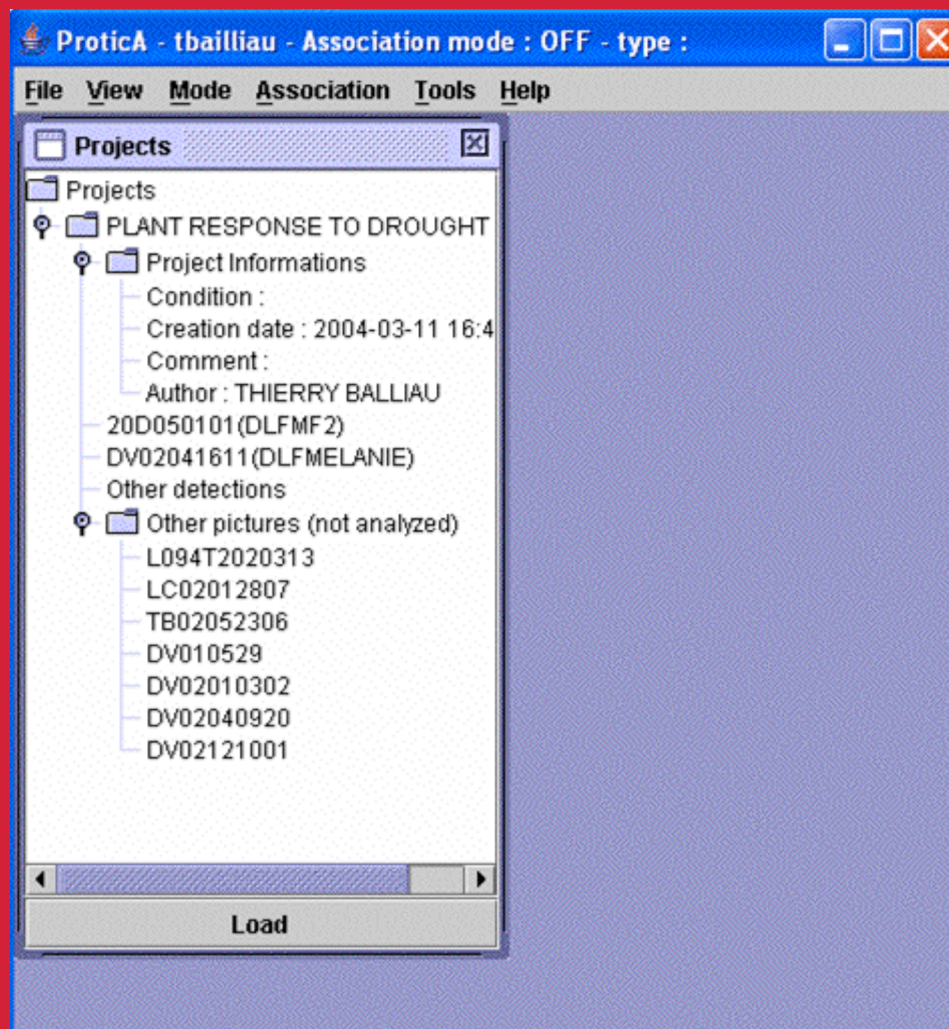
—

Select your gel
image file :

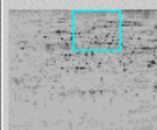
Parcourir...

Submit

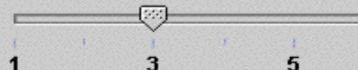




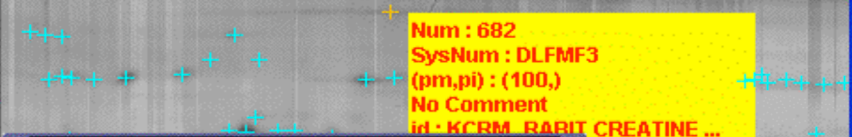
20D050101



Zoom level : 3



Gel informations :
 Plant name : F2
 Organ name : 6TH LEAF
 Project name : PROTIC
 Number of identified sp
 Number of detected sp



Spot Information

- Identification
 - validated
 - MS details
 - Ident 0(3)
 - gij125307|sp|P00563
 - KCRM_RABIT CREATINE KIN.
 - 125307|entrez (link)
 - P00563|swiss prot (link)
 - Peptid (TDLNHENLK)
 - Peptid (GQSIDDM*IPAQK)
 - gij23491144|gb|EAA22750.1
 - gij21672846|ref|NP_660911.1
- Quantification
 - Area :
 - Area unit :
 - Volume : 3577
 - Volume unit :
 - Pic height : 10
 - Pic height unit :
 - Sx : 8

[\[Entry info\]](#) [\[Name and origin\]](#) [\[References\]](#) [\[Comments\]](#) [\[Cross-references\]](#) [\[Keywords\]](#) [\[Features\]](#) [\[Sequence\]](#) [\[Tools\]](#)

Note: most headings are clickable, even if they don't appear as links. They link to the [user manual](#) or [other documents](#).

Entry information

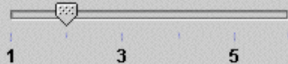
Entry name	KCRM_RABIT
Primary accession number	P00563
Secondary accession numbers	None
Entered in Swiss-Prot in	Release 01, July 1986
Sequence was last modified in	Release 01, July 1986
Annotations were last modified in	Release 43, March 2004

Name and origin of the protein

Protein name	Creatine kinase, M chain
Synonyms	EC 2.7.3.2

MASTER

Zoom level : 2



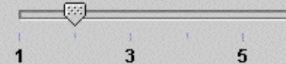
Gel informations :

Plant name : CORSE
Organ name : BOIS EN DL...
Project name : PINUS XY...
Number of identified spot : 251
Number of detected spot : 1039



ETE 1

Zoom level : 2



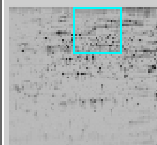
Gel informations :

Plant name : CORSE
Organ name : BOIS EN DL...
Project name : PINUS XY...
Number of identified spot : 0
Number of detected spot : 557



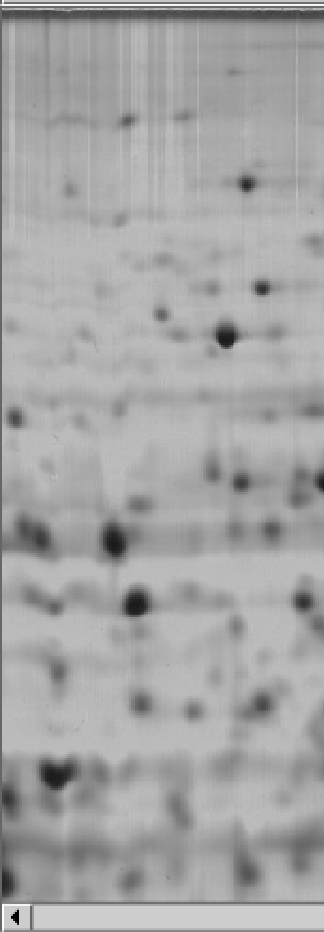
Num : 419
SysNum : NONE
(x,y) : (1850,1265)
No Comment
Relation : isoform
242 (MASTER) - 212 (MASTER)
Relation : match
419 (ETE 1) - 242 (MASTER)

20D050101



Zoom level

1

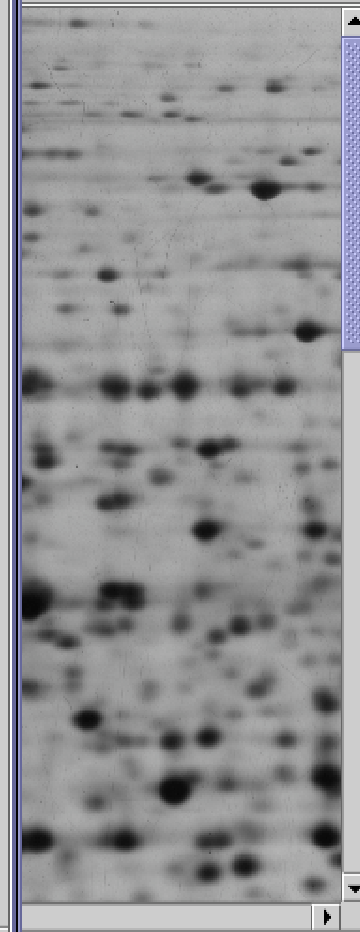


Exported data

spot_num:spot_soft_num (volume)	11D170101:PROCBILLESEGM:NO	17D301100:PROCBILLESEGM:NO
991:194	4267	0
145:1168	35716	0
396:1216	15536	0
484:702	12077	0
1571:14	0	42
231:705	0	2615
380:1014	0	81762
174:827	17604	0
1164:1008	9625	0
231:1340	30295	0
359:386	8069	0
1199:513	1918	0
164:326	7660	0
1689:507	0	72
977:1323	2244	0
676:772	0	3197
87:770	73313	0
525:312	4894	0
112:902	62745	0
1580:539	0	591
139:1290	25343	0
557:979	5239	0
244:404	0	596
343:684	0	33880
704:679	24156	0
800:257	4524	0
1685:761	0	505
133:117	0	1856
1533:1017	8255	0
962:196	0	669
671:552	0	3582
673:1626	12384	0
828:1638	10002	0
399:1628	28840	0
892:1309	6993	0
370:414	10830	0
1245:409	1307	0
1009:576	0	3888
132:356	28085	0

Gel informations :

Plant name : F2
 Organ name : 6TH LEAF
 Project name : PROTICDE...
 Number of identified spot : 0
 Number of detected spot : 2046



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