

GenePaint.org

Derk Geene
Harmen van der Spek

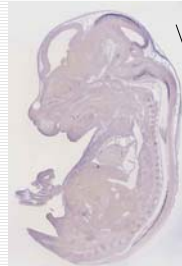
Outline

- What is GenePaint.org
 - In situ hybridization (ISH)
 - Data generation and processing
 - Data storage
 - Querying the database
 - Data visualization
-

About GenePaint.org

- **Representing gene expression patterns in mice**
- Max Plack Institute of Experimental Endocrinology
- Baylor College of Medicine, Department of Biochemistry and Molecular Biology
- According to article: GenePaint.org: An atlas of gene expression patterns in the mouse embryo

About GenePaint.org



E14.5 mouse
Dmbx1:
diencephalon/mesencephalon-expressed
brain homeobox 1

In situ hybridization

- Pairing labeled probe with mRNA
 - Labeling indicates amount of mRNA
 - Amount of mRNA indicates amount of gene expression
 - All done on cells, not on isolated material
-

Data generation

- Non-radioactive ISH on frozen sections of mouse embryos and mouse brains
 - Labeling technique makes 5 mRNA copies per cell visible (estimate)
 - Section of E14.5 embryo requires 195 exposures, each 500x668 pixels, 24 bit-color.
 - After cropping image 100-200 MB
-

Data processing

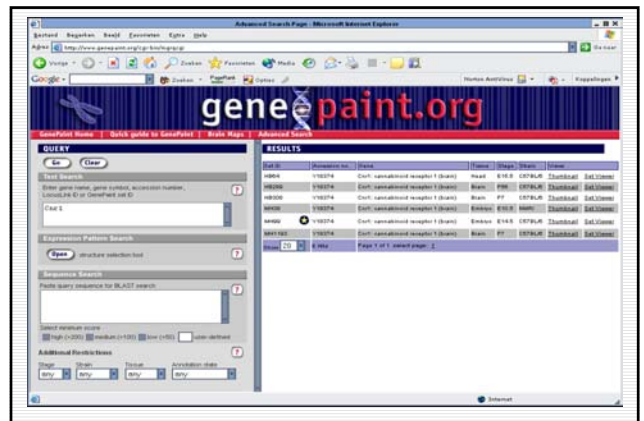
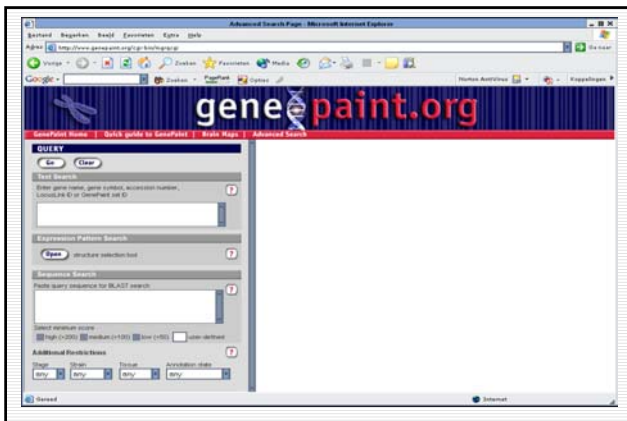
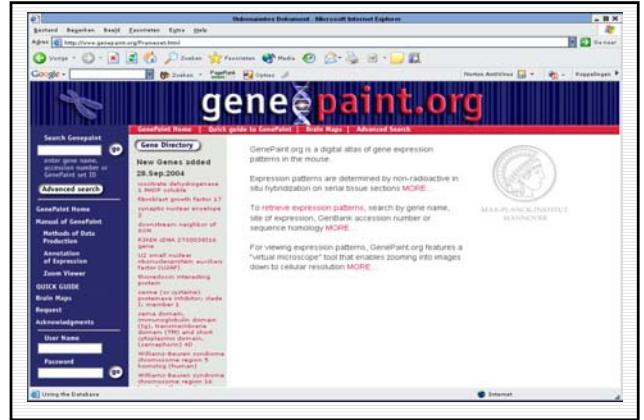
- Meta data:
 - Specimen identifier
 - Sequence of template
 - Hybridization conditions
 - Quality control
 - Template sequence validation
 - Tissue quality assessment
 - On failure, data are recollected
-

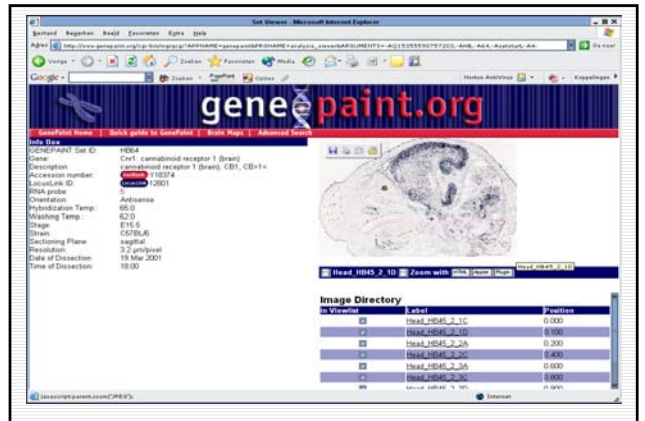
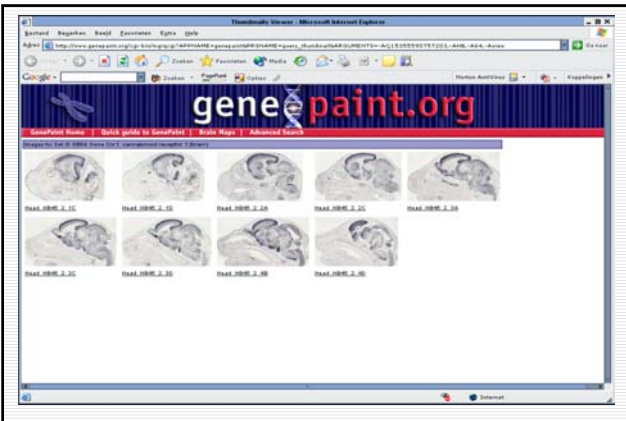
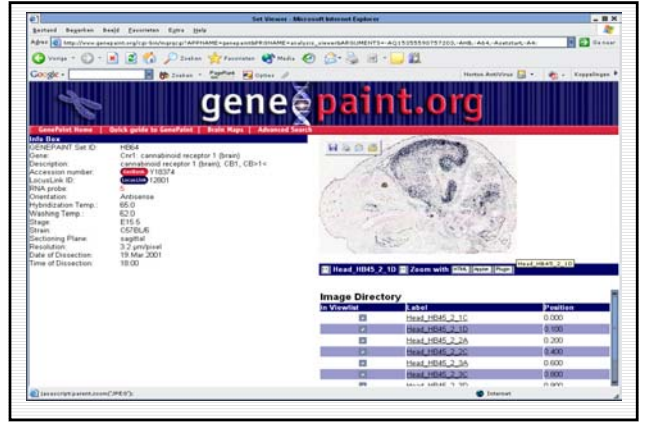
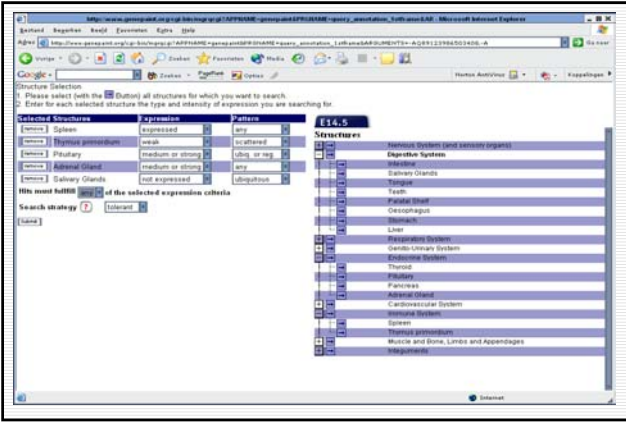
Data storage

- Relational DB2 database
 - Public part
 - Private part: Content management system
 - Nomenclature
 - LocusLink (Results link to this site)
 - Mouse genome database
-

Querying genepaint.org

- Public component
- Laboratory management system
 - Document experimental steps
 - Parameters for dataproduction
- Retrieving and viewing gene expression patterns





Set Viewer - Microsoft Internet Explorer

Address: http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=Nucleotide&cmd=Retrieve&list_uids=118374

Enter the Nucleotide - Microsoft Internet Explorer

NCBI Nucleotide

Search [Nucleotide] for 118374

Links: Summary | History | Clipboard | Details

1 118374
Mus musculus gene encoding cannabinoid CB1 receptor
g4480370[emb]118374 1[M]118374(480370)

NCBI Nucleotide

MESEQUEST Set ID: 118374
Gene: CB1 receptor
Description: cannabinoid recep
Accession number: 118374
LocusLink ID: 12801
5
Antenna
Hybridization Temp: 65.0
Washing Temp: E15
Stage: C50L6
Sectioning Plane: sagittal
Resolution: 3.2 micromet
Date of Dissection: 19 Mar 2001
Time of Dissection: 18:00

Enter the Nucleotide
Help | FAQ

Enter Tools
Check sequence
revision history
LinkOut

Internet

0.000
0.000
0.000
0.000
0.000

Set Viewer - Microsoft Internet Explorer

Address: http://www.ncbi.nlm.nih.gov/LocusLink/LocusLink.cgi?id=12801

LocusLink Report - Microsoft Internet Explorer

NCBI LocusLink

PubMap | Entrez | BLAST | OMIM | Map Viewer | Taxonomy | Structure

Search LocusLink | Display | Draft | Organism: AB

Query: [] [] []

View: Min Col | One of 1 List | Save All List

ABCDEFGHIJKLMNOPQRSTUVWXYZ
LocusLink will be replaced by Entrez Gene. Check Entrez Gene for current information.

Click to Display mRNA, Oligos, Alignments (updating 202598 hits)

Gene | Pub | Unpub | Map | RefSeq | Homolo | Ref | AF

UCSC

NCBI LocusLink Home

Entrez Index
Top of Page
Nomenclature
Overview
Function
Relationships
Map
RefSeq
Related Seqs
Links

LocusLink: Collaborators | Download | FAQ | Help | Index

Overview

Locus Type: gene with protein product, function known or inferred
Product: cannabinoid receptor 1 (brain)
Alias: CB1, CB1C

Set Viewer - Microsoft Internet Explorer

Address: http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?db=Gene&cmd=Retrieve&list_uids=118374

GenePanel Set ID: HB64 Gene: Cur 1: cannabinoid receptor 1 (brain) Stage: E15.5 - Microsoft Internet Explorer

GenePanel Set ID: HB64 Gene: Cur 1: cannabinoid receptor 1 (brain) Stage: E15.5

GenePanel Set ID: HB64 Gene: Cur 1: cannabinoid receptor 1 (brain) Stage: E15.5

Position: 0.000, 0.500, 1.000, 1.500, 2.000, 2.500, 3.000, 3.500, 4.000

SEE

Set Viewer - Microsoft Internet Explorer

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SEE

