

Information Systems in the Life Sciences

PERP GeneChip data warehouse;
Implementation of a dynamic time series
query tool with graphical interface

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Overview

- General
- PERP GeneChip data warehouse
- Affymetrix related information
- Focus on one of the tools:
Single Gene Query Tool (SQLT)

General

DNA databases have relatively low dimensionality Genome is a linear code	RNA expression and protein databases need to be able to handle very high dimensional data Time, tissue, cell type and genes are interrelated variables
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RNA & Microarrays

- RNA can be analyzed by Microarray expressions.
- Microarray expression have a high dimensionality of profile data.
- There is currently with no standard experimental platform.
- This complicated the development of web-accessible databases and analytical tools needed for RNA research.

Public Expression Profiling Resource

- PEPR == Public Expression Profiling Resource
- PEPR provides centralized Affymetrix expression profiling data to the public research community
- There is no need for the public researcher to have their own Affymetrix software to use PEPR database
- PEPR claims to be the largest public resource having quality control and standard operating procedures (QC/SOP).
- All future implementations will include these QC/SOP criteria.
- Currently there are 1426 profiles available to the public

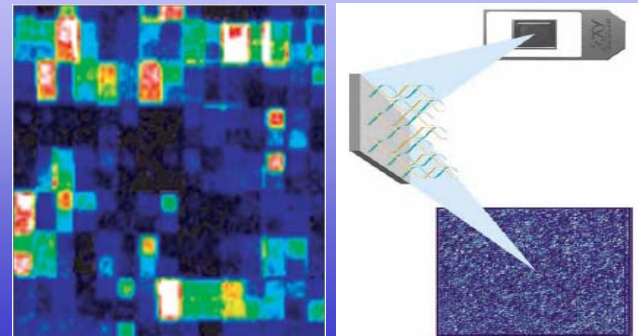
Tools from PEPR

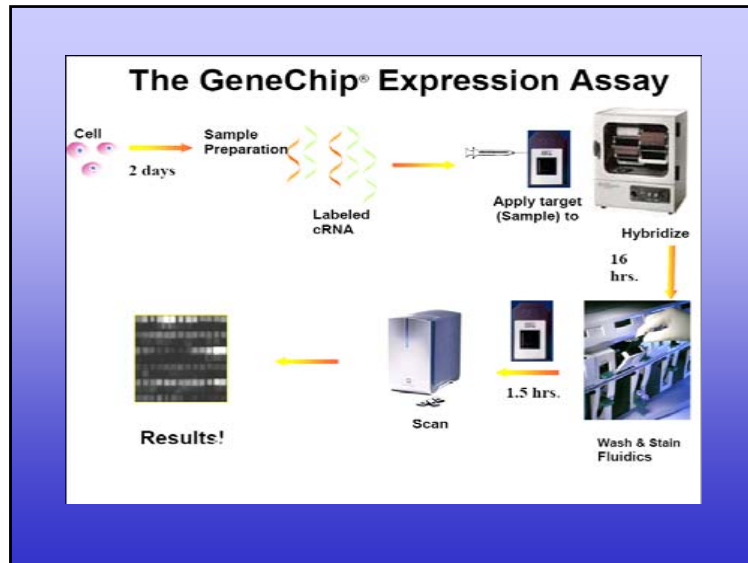
- Oracle based web solutions:
 - Single Gene Query Tool
 - Goal: To query PEPR for all profiles matching the desired queries.
 - PEPR project/file search.
 - Functional Clusters
 - Goal: To provide rapid download of data within a "project"
 - Export of Microarray profile data
- Back end solutions:
 - Direct deposit of data to NCI GEO database
 - Goal: To provide automated link to move profile data to GEO

Affymetrix

- Commercial firm for delivering GeneChips
- Products of Affymetrix : GeneChip Arrays, Assays & Reagents, Instruments and Software
- **Microarray Suite (MAS 5.0)**

How Does It Work





- ## Single Gene Query Tool
- SGQT == Single Gene Query Tool
 - Currently implemented for time series only
 - Goals is to provide:
 - dynamic queries
 - any gene in the genome
 - graphical visualizations
 - spreadsheet downloads

- ## How does the public SGQT work functionally
- Simple project query interface.
 - Gene search function supporting ambiguous queries
 - Graphical visualization showing all replicates
 - Average generated on the fly
 - Mouse-over information for each replicate and average
 - Access to additional database resources with a mouse click
 - Able to download all data in the graph for Spreadsheet use

Access

- Trough URL or using the menu from the main web site “Data” “Search” “Single Gene Query Tool”
- User selects species and “time series”

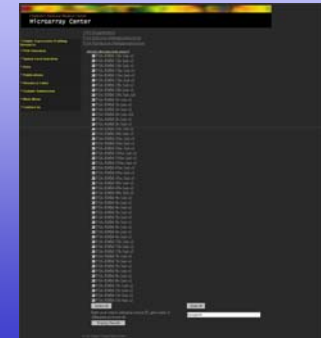
Project Selection

- User selects the list of implemented projects that are shown



Profile Selection

- User select the expanded list of profiles of the selected project
- User types in the requested Gene (search function supports ambiguous queries)



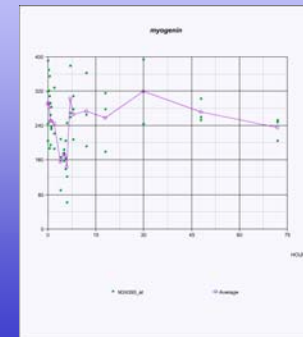
Gene Selection

- User select the required gene from the resulting drop-down menu

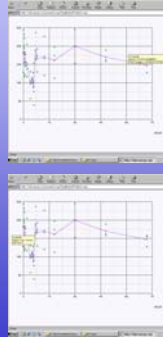


Graphical visualization

- Showing all replicates, with average generated on the fly
- Able to download for all data in the graph for use with a spreadsheet



- Mouse-over for each replicate and average, showing data associated with that data point



Additional resources

- Clickable data points for access to additional database resources for the gene under study

- [Gene Name: myogenin](#)
- [Unigene: Rn 9465](#)
- [Genbank: M24393](#)
- [LocusLink: 29148](#)
- [Affymetrix: M24393_at](#)
- [Download the original dataset: P6A.RMM.Ct.4aA-2](#)

Conclusion

- User needs skills in microarray experiments to use all the available tooling
- SGQT is usable without experience in interpretation of microarray data
- Some of the tools are in the background
- PERP is trying to have reliable project/data using QC/SOP available for the public

Questions?

The End.



References

Websites used:

- www.affymetrix.com
- microarray.cnmcresearch.org

Additional documentation used:

- [pepr2004nar.pdf](#)

Images used from Affymetrix (Must be used according to the licensee agreement):

- [an_actual_array_image.pdf](#)
- [chip_probes_output.pdf](#)
- [gene_expression_assay.pdf](#)