

# Data Analysis and Modeling Techniques

## Microarray Technology

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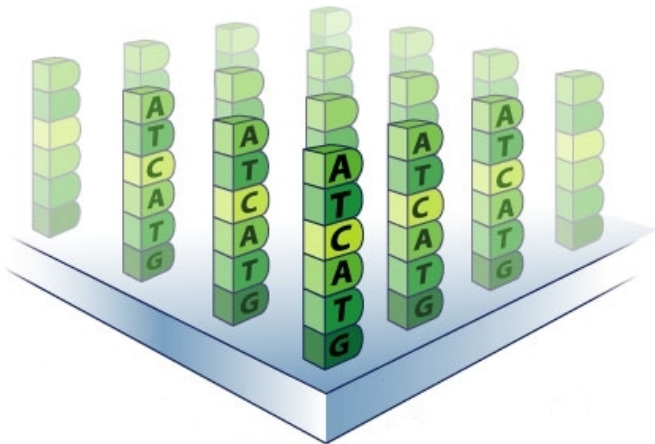
A *microarray* is composed of

- DNA fragments fixed on a solid support
- ordered position of probes
- principle of hybridization to a specific probe of complementary sequence
- molecular labeling

➔ Simultaneous detection of thousands of sequences in parallel

# Microarray

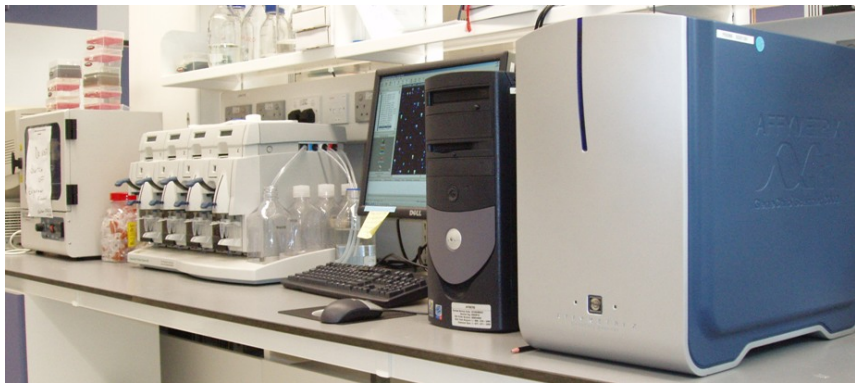
## Probes



There exist several high-throughput methods to simultaneously measure the expression of a large number of genes :

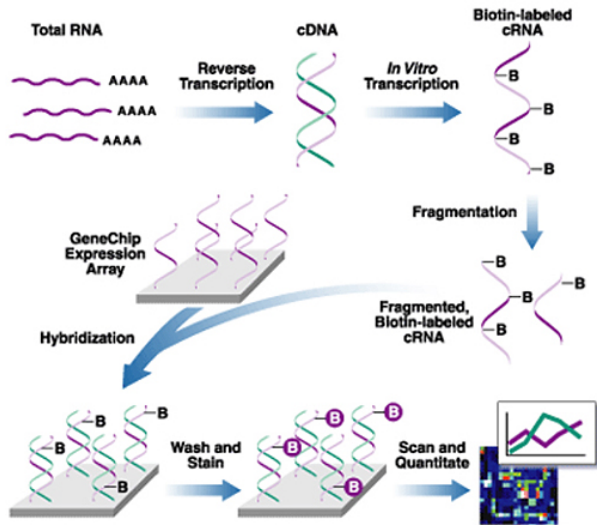
- cDNA microarray
- oligonucleotide microarray
  - ▶ short oligonucleotide ([AFFYMETRIX<sup>©</sup>](#))
  - ▶ long oligonucleotide ([AGILENT<sup>©</sup>](#), [CODELINK<sup>©</sup>](#))
- multiplex quantitative RT-PCR

We will present the [AFFYMETRIX<sup>©</sup>](#) platform in order to overview the main characteristics of microarray technology.



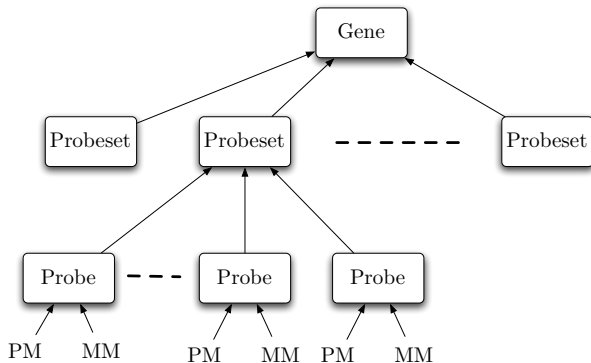


# AFFYMETRIX<sup>©</sup> Design



# AFFYMETRIX<sup>©</sup> GeneChip Structure

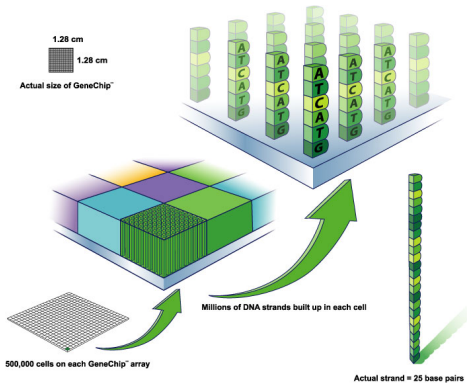
- 1 gene is represented by 1 or more probe sets
- 1 probe set includes 11 to 20 probe pairs
- 1 probe pair includes a Perfect Match (PM) value and a MisMatch value (MM)





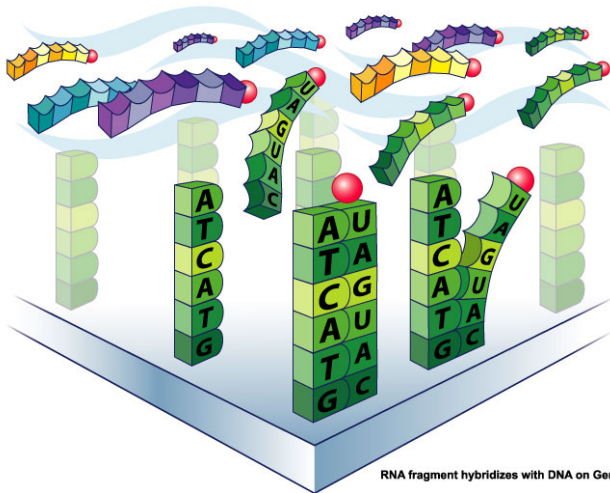
# AFFYMETRIX<sup>©</sup> GeneChip Structure

Suite



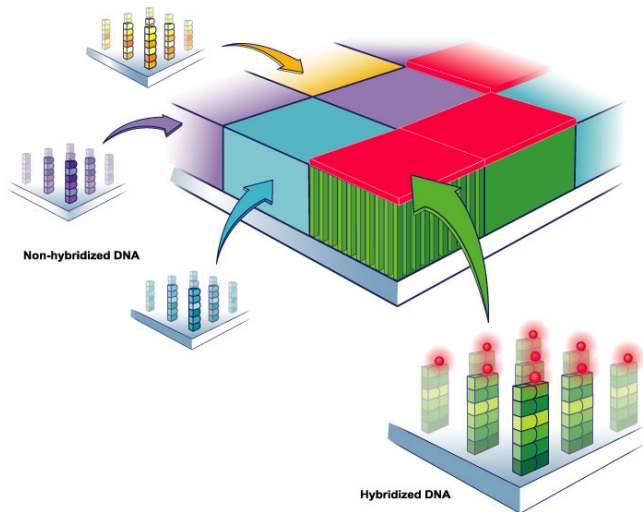
# AFFYMETRIX<sup>©</sup> Hybridization

RNA fragments with fluorescent tags from sample to be tested



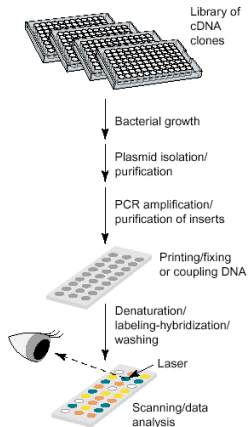
RNA fragment hybridizes with DNA on GeneChip

Shining a laser light at GeneChip causes tagged DNA fragments that hybridized to glow

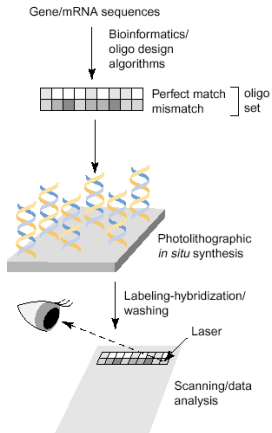


# Microarray Comparison

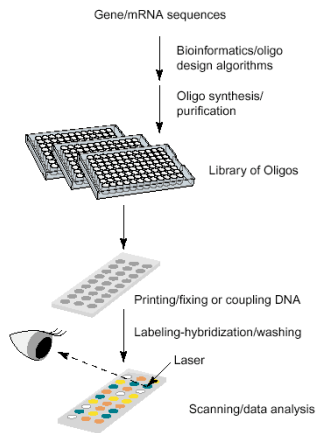
## cDNA



## short oligonucleotide



## long oligonucleotide



AFFYMETRIX<sup>©</sup> advantages :

- commercially available for several years (strong manufacturing)
- large number of published studies (generally accepted method)
- no reference sample → possible comparison between studies

AFFYMETRIX<sup>©</sup> disadvantages :

- cost of the devices and the chips (but easy use)
- changes in probe design is hard (but new program permits to create his own design)
- short oligos → several oligos per gene, specificity/sensitivity trade-off (complex methods to get gene expression)

- Course web page : [http://www.bioinfomaster.ulb.ac.be/cursus/index\\_html/en#DATANA](http://www.bioinfomaster.ulb.ac.be/cursus/index_html/en#DATANA)
- Personal homepage : <http://www.ulb.ac.be/di/map/bhaibeka/>
- This presentation : [http://www.ulb.ac.be/di/map/bhaibeka/bioinfo\\_courses/microarray\\_pres\\_hkb.pdf](http://www.ulb.ac.be/di/map/bhaibeka/bioinfo_courses/microarray_pres_hkb.pdf)

**Thank you for your attention.**