

Technology meets the patient and
patient meets technology:
*Exploiting the human genome to improve cancer
diagnostics*

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NKI-AVL



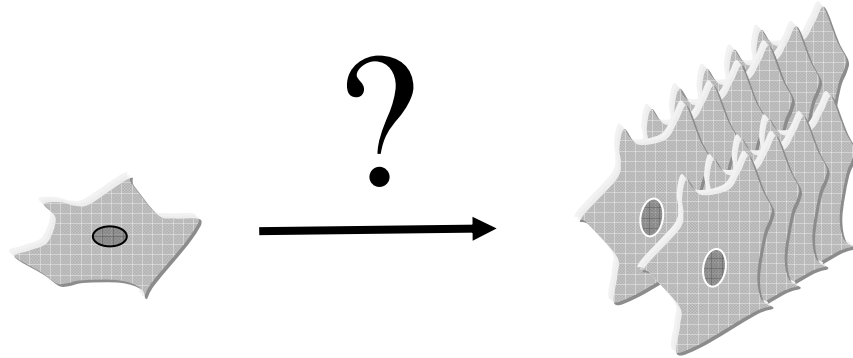
The Netherlands Cancer Institute
Antoni van Leeuwenhoek Hospital



Agendia™

The Agendia logo features a stylized, multi-colored ring (yellow, orange, red, green, blue) above the word "Agendia" in a bold, black, sans-serif font. A small trademark symbol (TM) is located to the right of the word. Below the word "Agendia" is a faint, light-colored shadow.

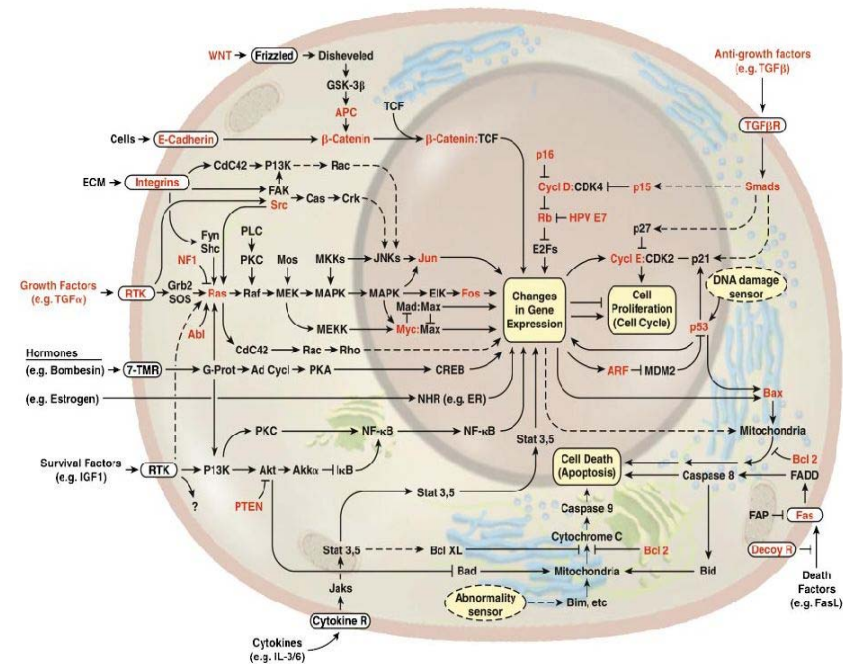
Over 25 years of progress in cancer research



Normal cell

Cancer

1980



2006

In spite of all this progress:

- We still use the light microscope for cancer diagnosis
- We still use many broadly-acting cytotoxic drugs to treat cancer

Genomic technologies can help up to:

- Develop better cancer diagnostics
- Develop more specific cancer therapeutics

The microscope, a major tool for diagnostics for the last 350 years



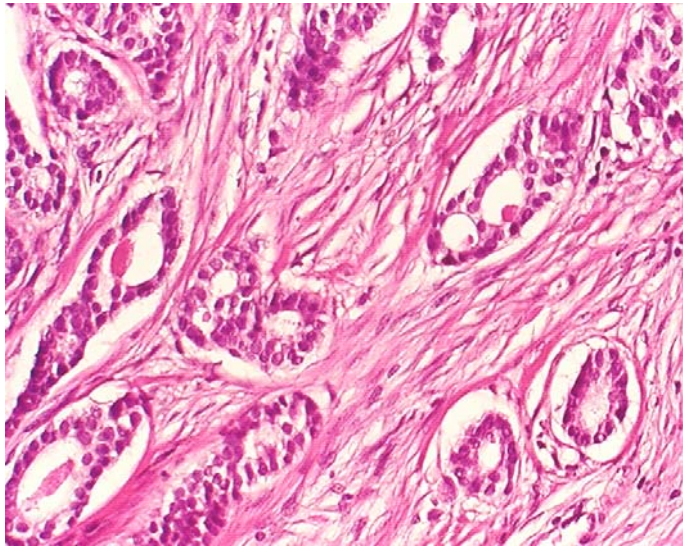
Van Leeuwenhoek microscope, late 1600s



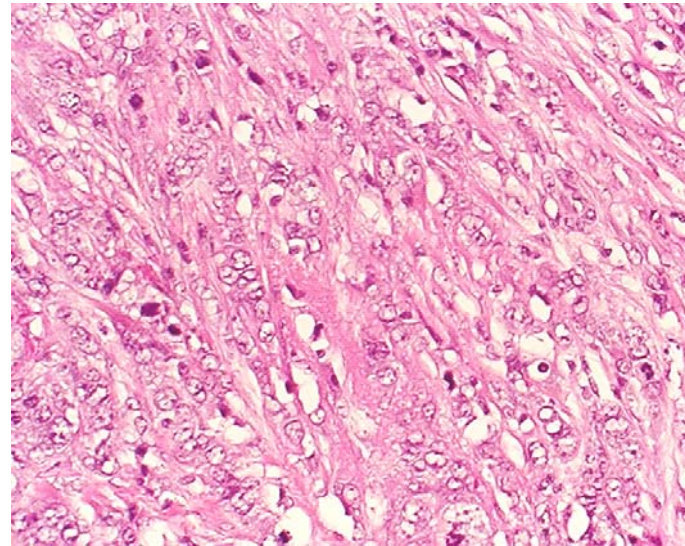
Olympus microscope, late 1900s

Using microscopy to predict disease outcome in cancer

- Grade 1



- Grade 3

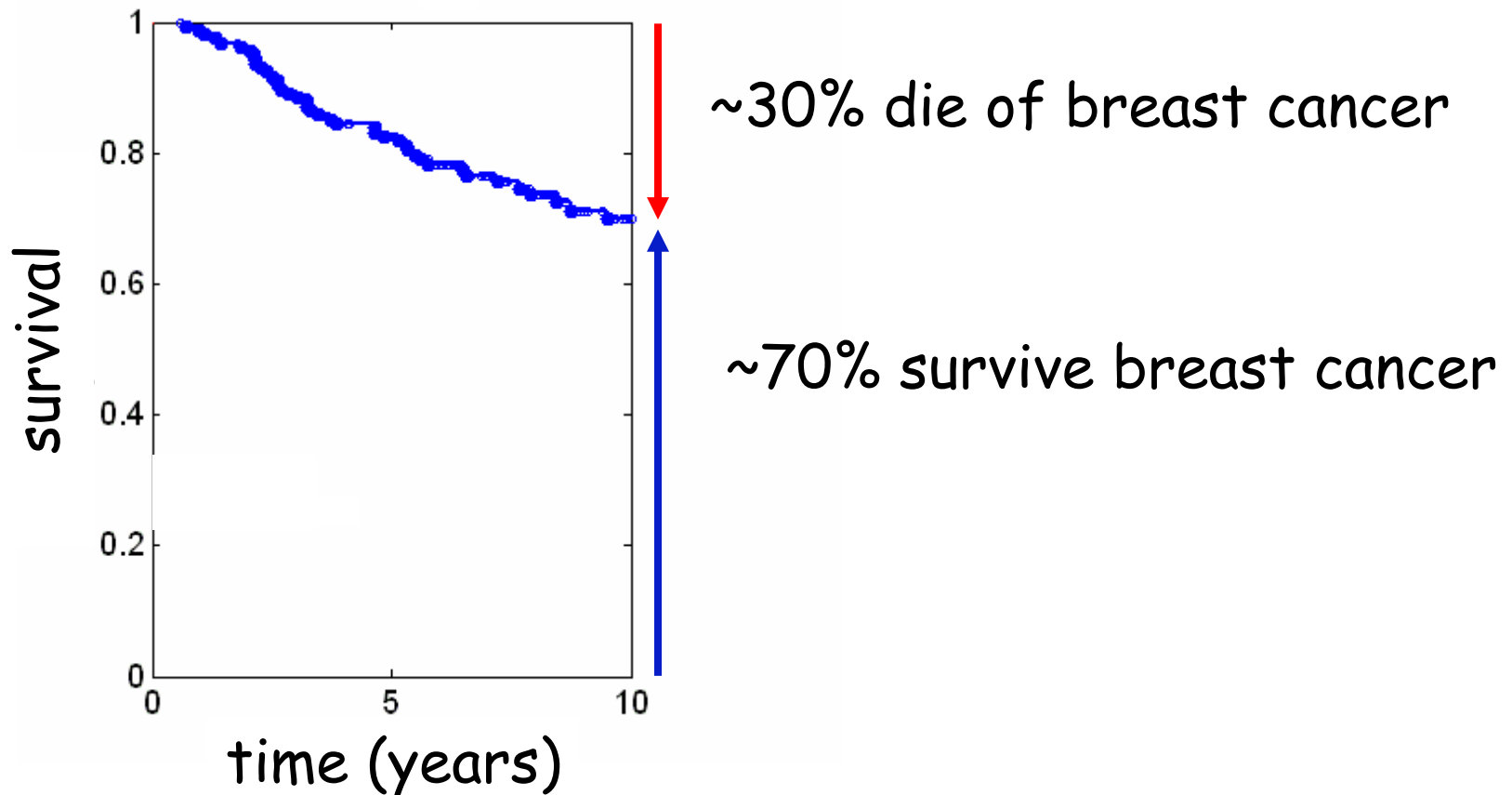


Histological grade

Breast Cancer - Survival

Pre-menopausal patients, lymph node negative

Kaplan-Meier Survival Curve



Breast Cancer - Treatment


premenopausal, lymph node negative

Current adjuvant treatment selection criteria:

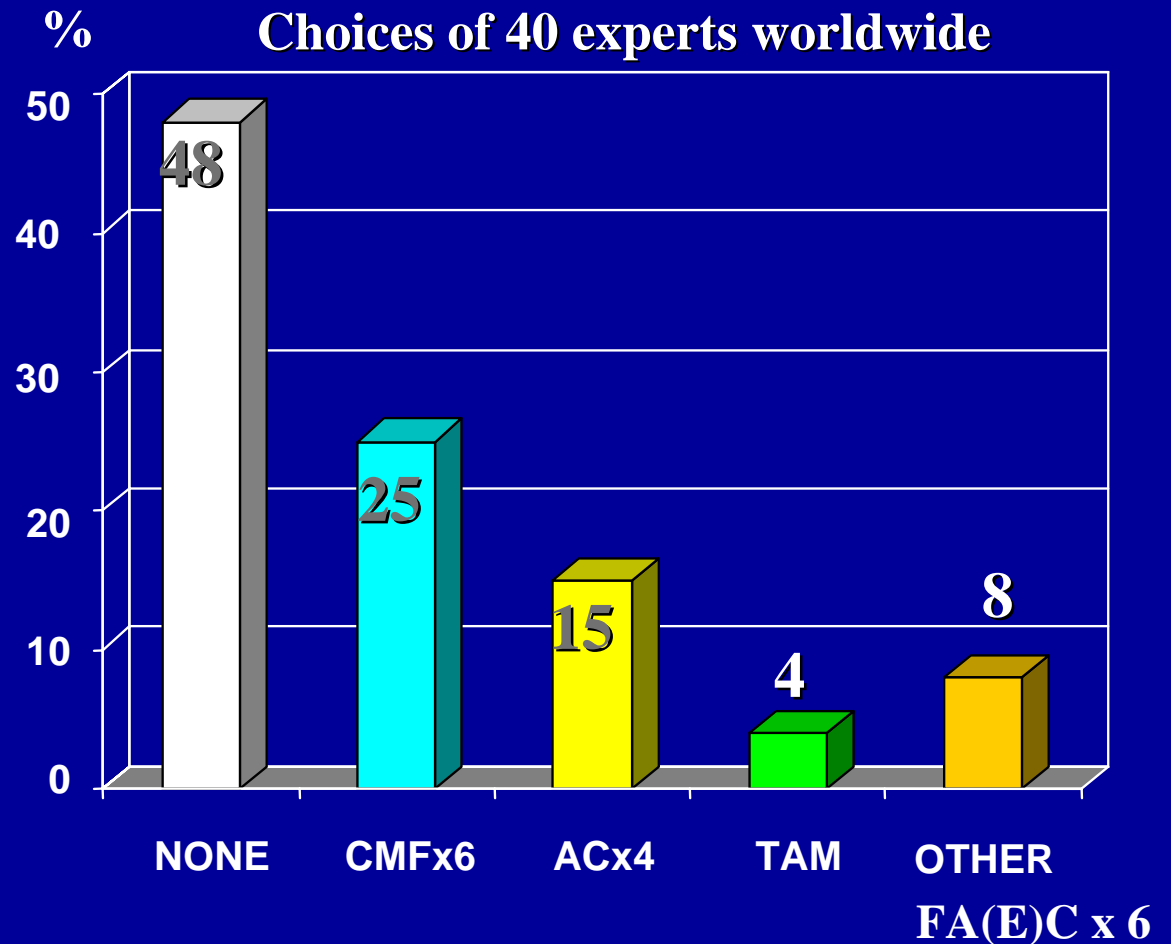
- NIH (US) consensus criteria: > 95%
- St Gallen (EU) consensus criteria: > 80%
receive adjuvant chemo- and hormonal therapy

As only 30% of these patients develop distant metastases, some 50-65% of patients are over-treated with adjuvant (chemo)therapy

SHOULD ONE TREAT A SMALL (<1CM) ENDOCRINE UNRESPONSIVE BREAST TUMOR ?



61 y-old, fit,
postmenopausal
Node negative
pT = 0.9 cm
ductal cancer
ER and PR
negative
HER2 negative
Grade 2



Courtesy of Martine Piccart

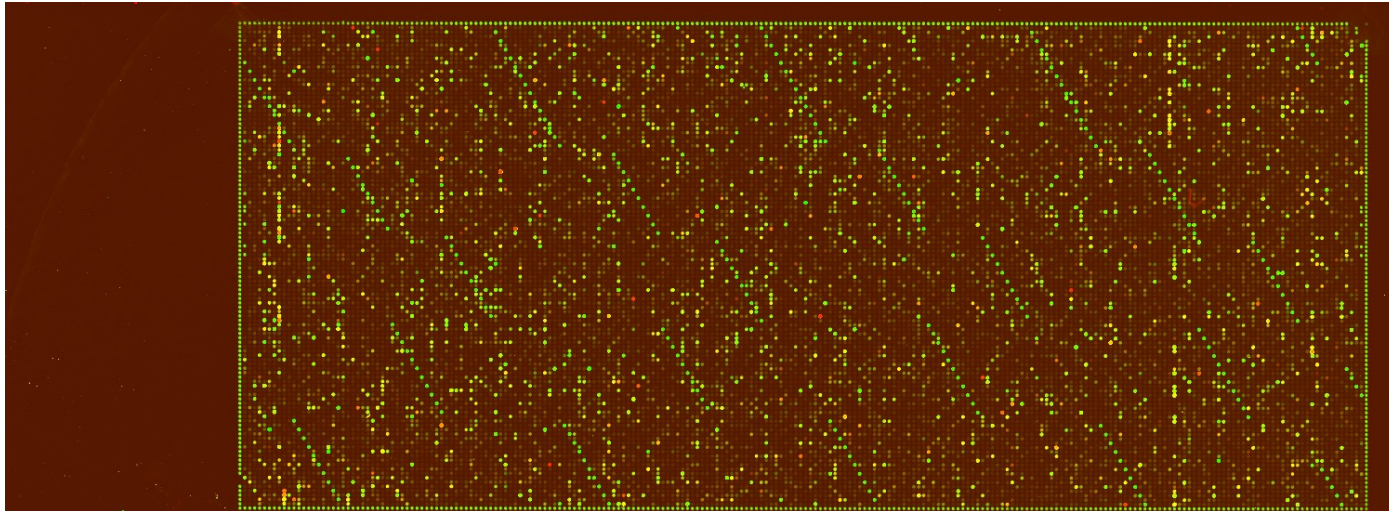
Few data, little information



Tumor cell behavior is determined by the activity of many genes

- The activity of one or a few genes cannot predict tumor cell behavior in a reliable way.
- We need tools to measure the activity of many genes in a single experiment

Gene Expression Profiling



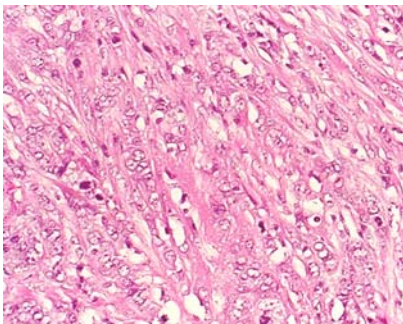
Measure the activity of tens of thousands of genes in a single experiment

"molecular portrait of cancer"

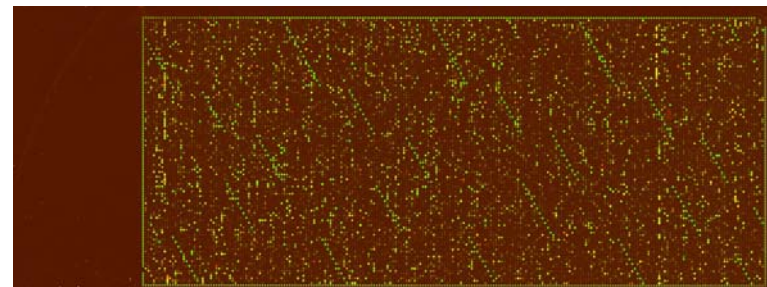
From micro-scope to micro-array



Micro-scope



Micro-array



Can gene expression profiling
be used to more accurately
predict clinical outcome of
disease in breast cancer?

Supervised Classification for prognostic subclasses

78 breast tumors ('83-'94)
patients < 55 years
tumor size < 5 cm
lymph node negative (LNO)
no adjuvant therapy

Prognosis Reporter Genes

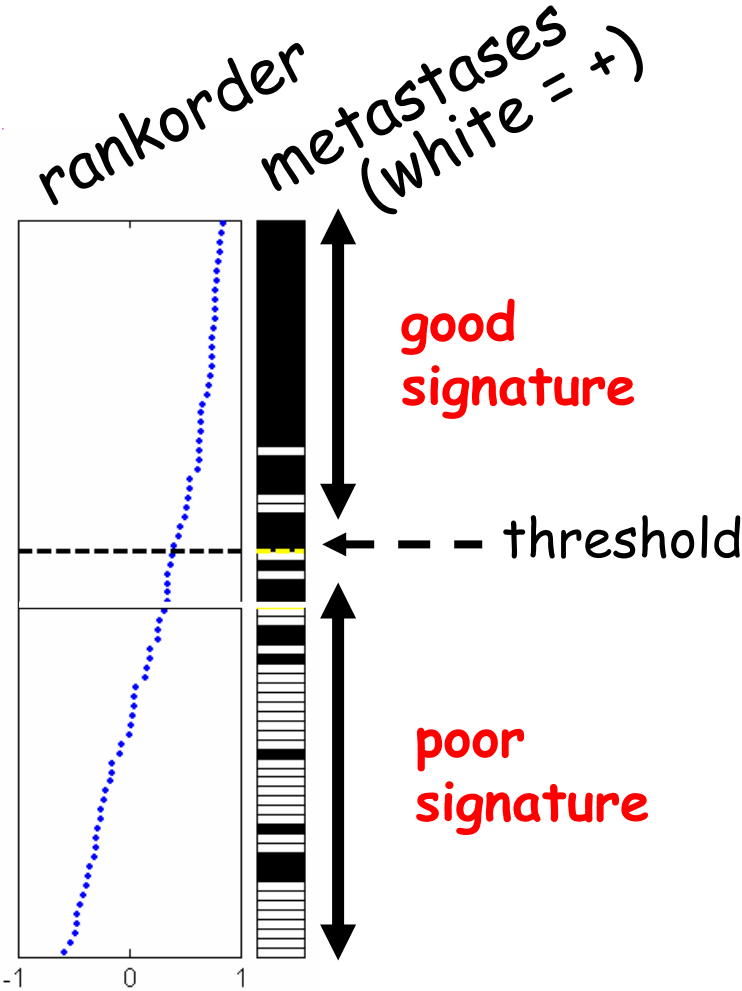
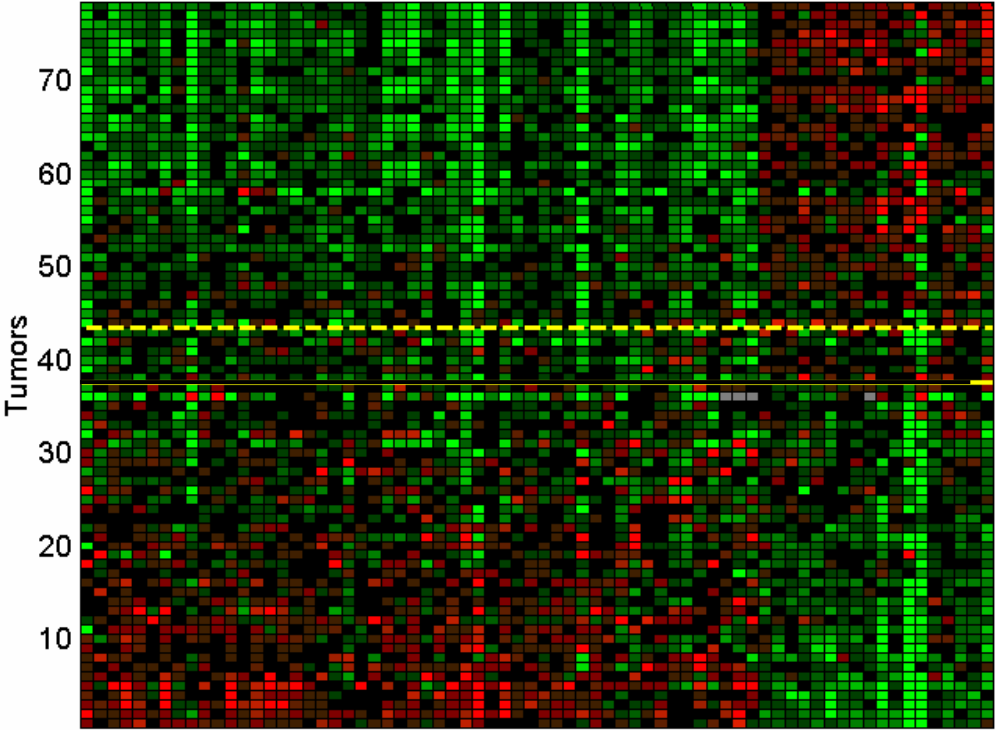
distant metastases
< 5 years (n=34)

no distant metastases
in at least 5 years (n=44)

Microarray-based prognosis of breast cancer

70 prognosis genes

78 tumors

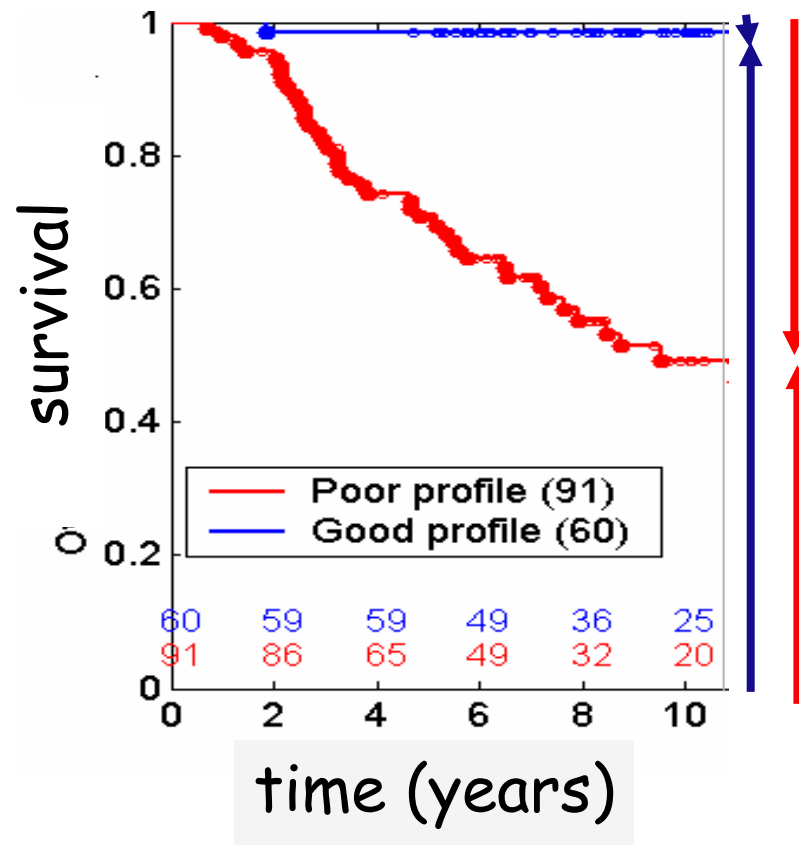


threshold set at 10% false negatives

91 % sensitivity, 73% specificity

Breast Cancer - Survival by profiling

Distinguish in: 40% good profile, 60% poor profile

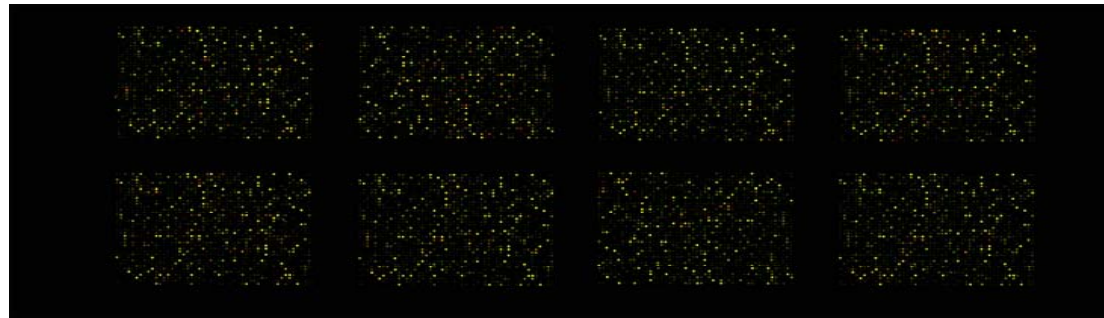
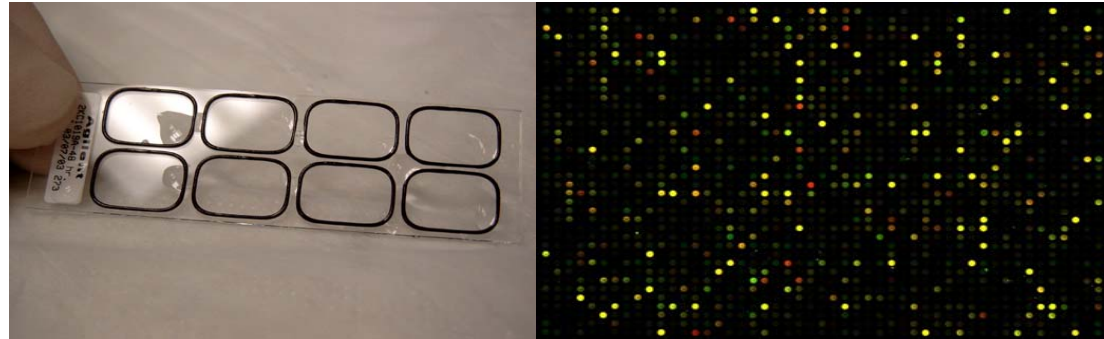


good profile:
~4% die of breast cancer
~96% survive breast cancer

poor profile:
~50% die of breast cancer
~50% survive breast cancer

151 patients, <53, LN0
10 year survival curve

"MammaPrint" Breast cancer prognosis array



Regulatory; QA/QC:

CLIA -registered, ISO 17025-certified,
CE- marked, FDA registration ongoing



Changing attitudes about complex gene tests in clinical practice

2002:

Breast cancer prognosis profile identified:

Not a single diagnostic company interested to commercialize the 70-gene breast cancer prognosis test

2004:

Commercial launch MammaPrint by Agendia

"St. Gallen" breast cancer consensus meeting votes against microarray testing 34 to 0

2006:

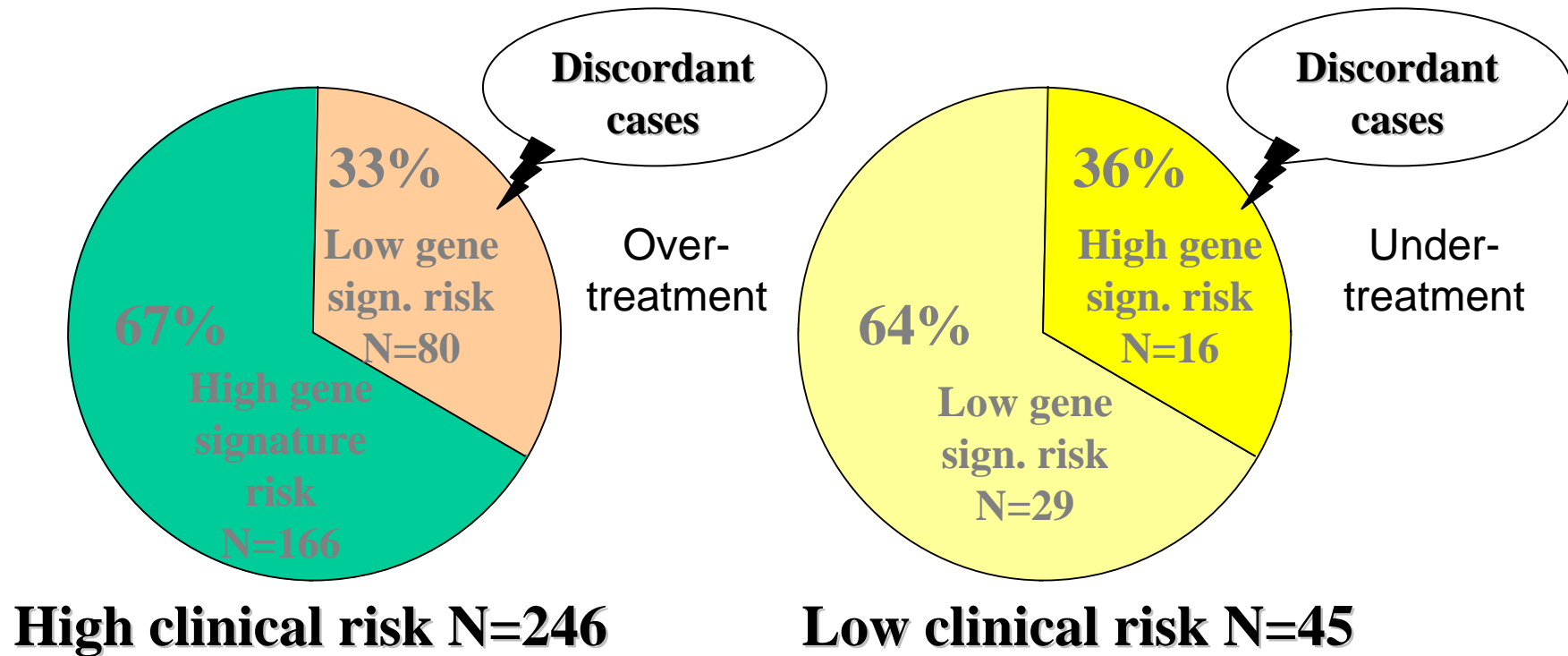
At ESMO, 50% of audience votes in favor of microarray tests for ALL patients, 90% in favor in special cases

In 2006, over 15,000 gene tests will be sold in USA for breast cancer



CONCORDANCE BETWEEN CLINICAL AND GENE SIGNATURE RISK CLASSIFICATION

Threshold for low clinical risk defined as predicted 10-year O.S. > 90%



Discordant cases with other clinical risk classifications

St-Gallen = 35%
NPI = 36%

St-Gallen = 43%
NPI = 54%

Clinical applications of microarrays

Who to treat:

- Prognosis profiles as diagnostic tool
-> improved selection for adjuvant therapy

How to treat:

- Predictive profiles for drug response
-> selection of patients who will benefit most

**WHO
NEEDS
THERAPY?**

Prognostic factors

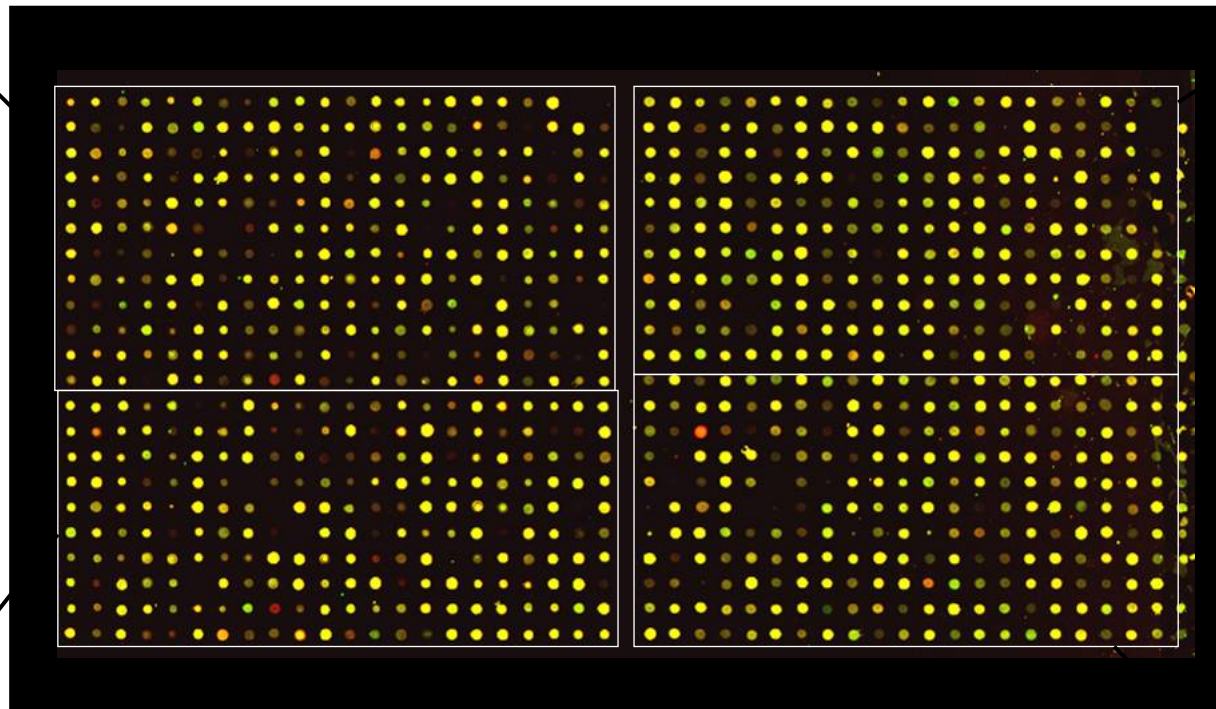
**WHICH
THERAPY WILL
WORK BEST?**

Predictive factors

The (near) future: Multiple predictions made by a single microarray test

Will the tumor come back?

Will the tumor respond radiation?



Will the tumor respond do drug A?

Will the tumor respond do drug B?

Acknowledgements

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