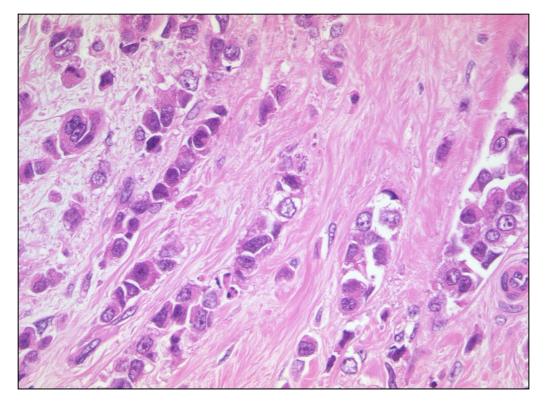
Techniques in pathology



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Berlin, Germany 7.5.14 IMPAKT Meeting, Brussels



Conflict of interest statement

- Research funding, honoraria, shareholder: Sividon Diagnostics
- Research funding: Siemens Medical Solutions

Outline – techniques in pathology

Introduction

- research strategies in pathology
- strengths and weaknesses
- FFPE tissue
 - H&E
 - immunohistochemistry
 - digital imaging
 - RNA analysis
 - DNA analysis
 - NGS sequencing
- workflow options

Options for research in pathology

- Why are we doing the research project?
- What are the aims?
- Options for pathologists:
 - *— "*traditional approach" definition and description of new tumor entities
 - hypothesis-generating research
 - predictive biomarker-focussed research
 - practice changing research

Level of evidence for biomarker studies

Type of tumor marker study		Definition	Possible level of evidence	
A	Prospective	clinical trial designed to address tumormarker	1	validation preferred, but not required
В	Prospective using archived samples "prospective- retrospective"	prospective biomarker design, existing samples collected in clinical trial	1	two studies with identical results
			2	only one study
С	Prospective observational	prospective registry and sample collection, no standardized treatment and follow-up	2	two studies with identical results
			3	only one study
D	Retrospective observational	collection of samples from archive, no standardized treatment	4-5	hypothesis generating, no clinical utility

Simon, Paik, Hayes JNCI, 2009

JOURNAL OF CLINICAL ONCOLOGY

Publication of Tumor Marker Research Results: The Necessity for Complete and Transparent Reporting Lisa M. McShane and Daniel F. Hayes

REVIEW ARTICLE

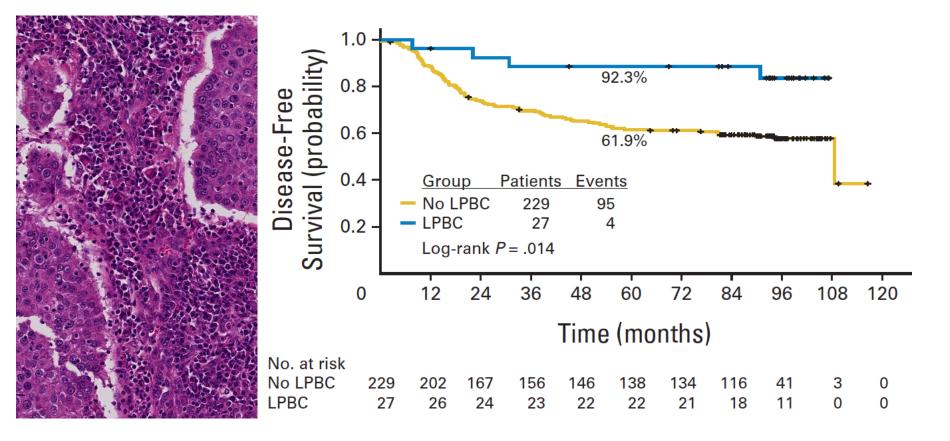
- Table 1. Requirements for a Marker-Based Test to Reach Level IB Evidence of Clinical Utility Based on Prospective-Retrospective Studies
- Adequate amounts of archived specimen must be available from enough patients from a prospective trial (which for predictive factors should generally be a randomized design) for analyses to have adequate statistical power and for the patients included in the evaluation to be clearly representative of the patients in the trial.
- The marker-based test should be analytically and preanalytically validated for use with archived specimens.
- The plan for marker evaluation should be completely specified in writing before the performance of marker assays on archived specimens and should be focused on evaluation of a single completely defined markerbased test.
- The results from archived specimens should be validated using specimens from one or more similar, but separate, studies.

NOTE. Guidelines adapted.²²

Requirements for pathology research 2014

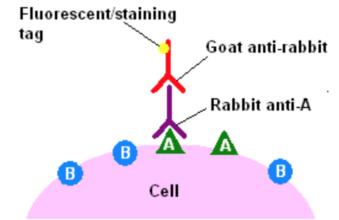
- large sample cohorts (aim: >1000 samples, but difficult to reach)
 - only possible for FFPE tissue
 - heterogenous cohorts vs tumor-type specific cohorts ?
 - prognostic markers may simply be markers of luminal differentiation
 - very difficult to find markers in TNBC cohorts
 - additional cohorts for validation
- prespecified analysis plan as a document
- methods validation

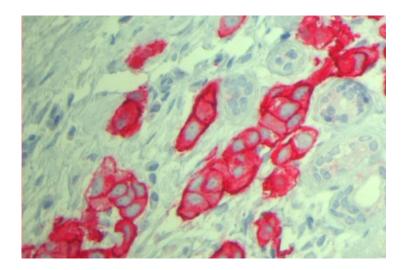
H&E based studies - Prognosis of TNBC – increased lymphocytic infiltrate defines a good prognosis group Loi et al, JCO 2013 BIG2-98 study (total n=2009, TNBC n=256)



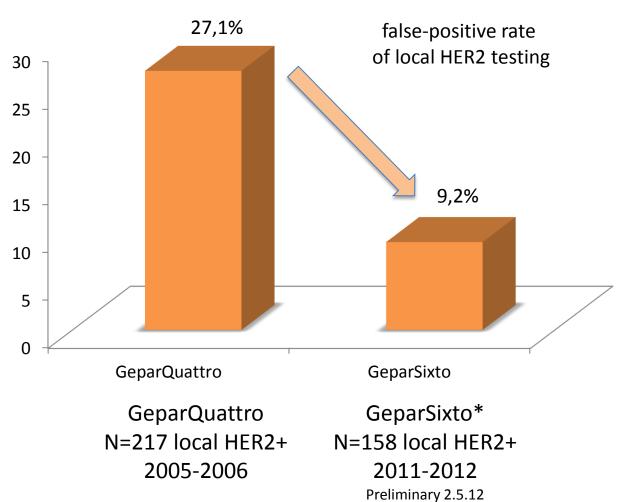
Immunohistochemistry = "in situ proteomics"

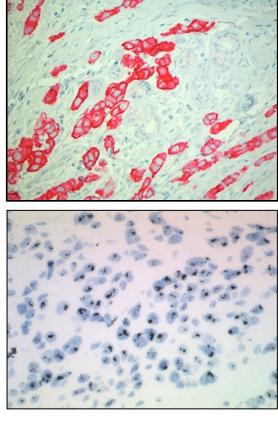
- antibody-based detection of molecular markers on tissue slides
- Advantages:
 - easy, useful on FFPE tissue
 - combined molecular and morphological information
 - type of cells, localisation in cells
- Disadvantages:
 - standardisation issues
 - staining intensity
 - percentage of of positive cells





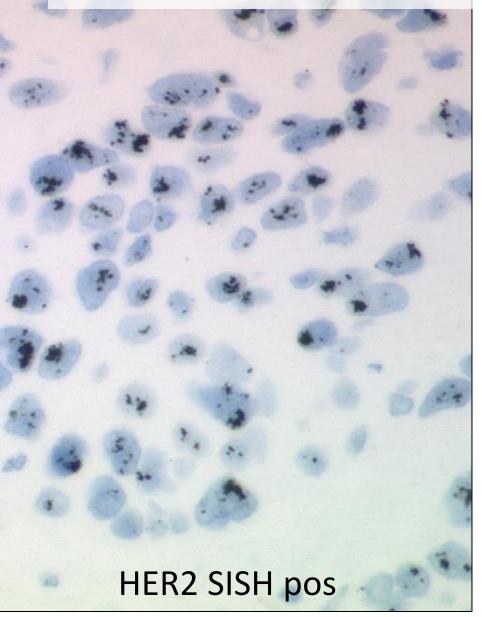
Comparison of local and central HER2 in GBG trials

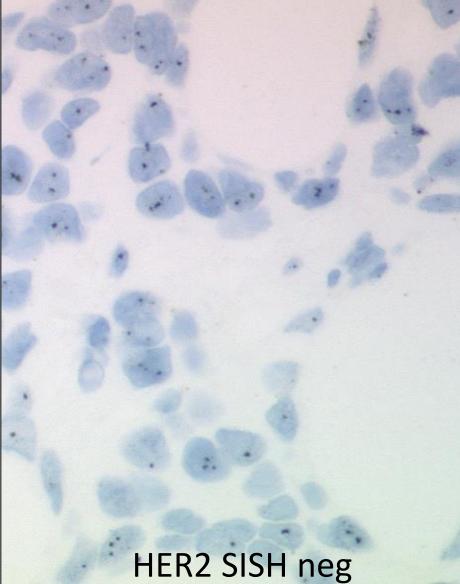






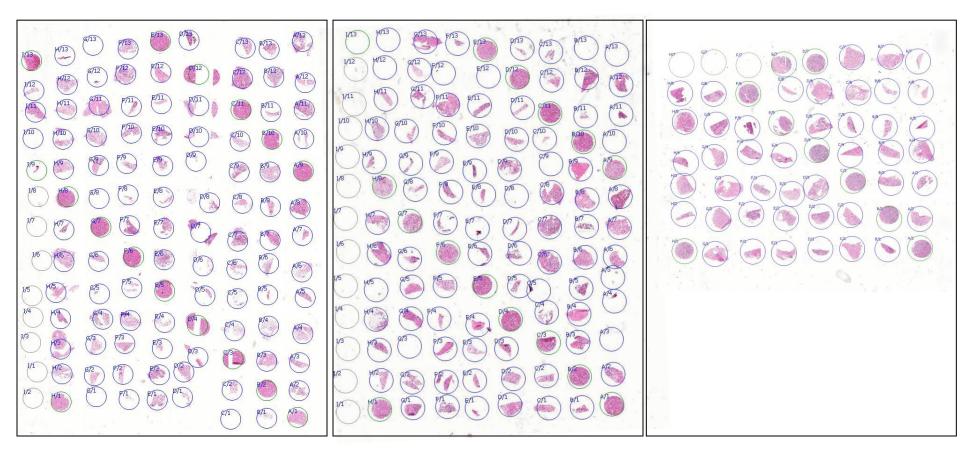
in-situ hybridisation





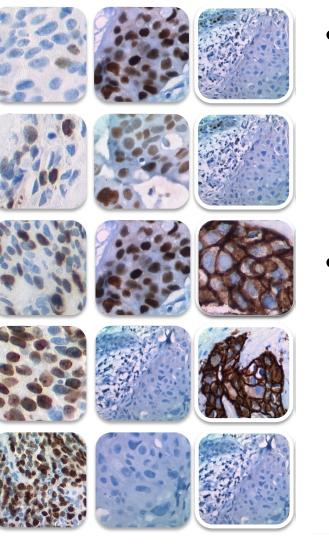
TMA – tissue microarray

TMA from 227 pretherapeutic core biopsies from HER2-positive Tumors from the GeparQuattro-trial



TMA 1

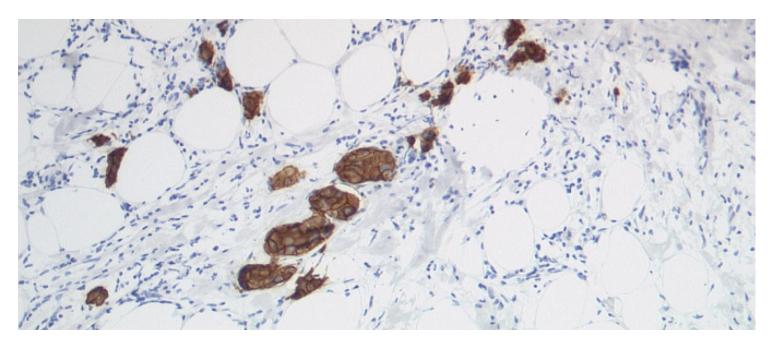
Limitations of immunohistochemistry

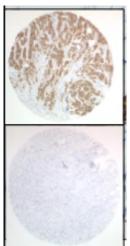


- Technical issues
 - Quantification of markers and cutoffs, e.g. Ki67
 - Interobserver variability
 - Assay standardisation
- More complex questions:
 - Endocrine Tx vs. Chemoendocrine Tx
 - Different types of anti-HER2 therapy
 - Response to anti-angiogenic Tx

Immunohistochemistry - standardisation

- use of Autostainers
- internal / external controls
- quantitative markers image analysis

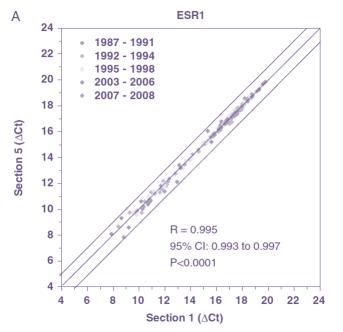




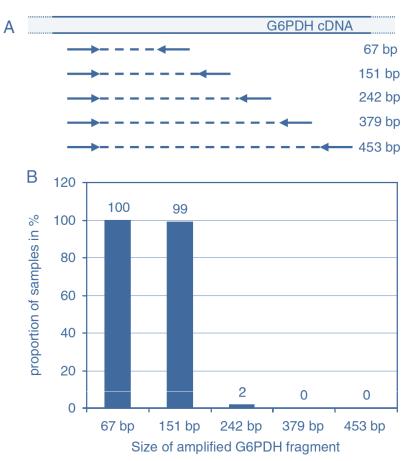
mRNA biomarkers in FFPE tissue

- mRNA isolation is feasible from FFPE tissue ("Formalin is an RNA-protective substance")
- 2. mRNA analysis can be used to assess breast cancer biomarkers
- 3. EndoPredict / Oncotype Dx routine mRNA expression analysis in breast cancer

RNA Isolation from FFPE tissue is feasible

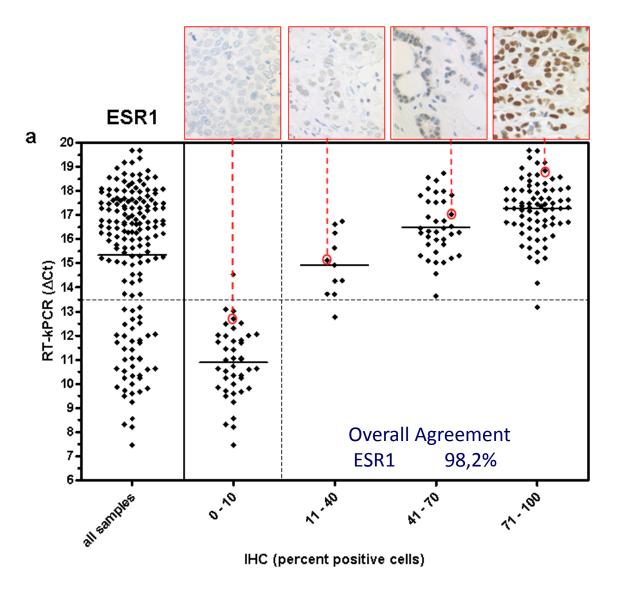


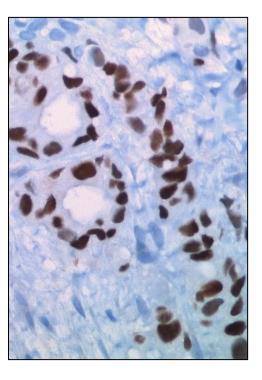
- 167 FFPE samples, age up to 21 years
- 501 RNA isolations
- Fragment length: ca 150bp
- RNA: ca. 1ug/10um section
- = ca. 100 PCR reactions
- High concordance of consecutive sections



Müller et al., 2011

Concordance for ESR1 measured by RNA analysis and immunohistochemistry

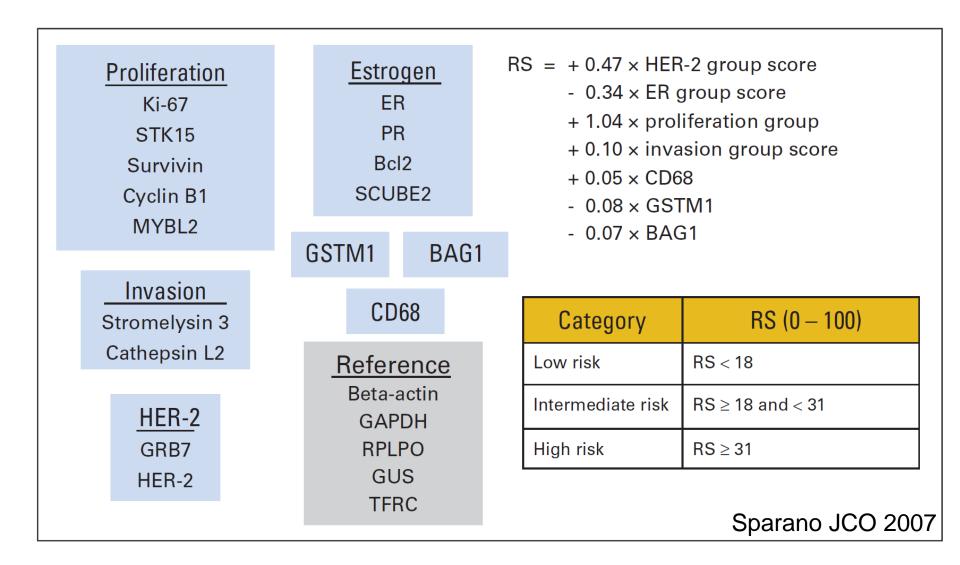




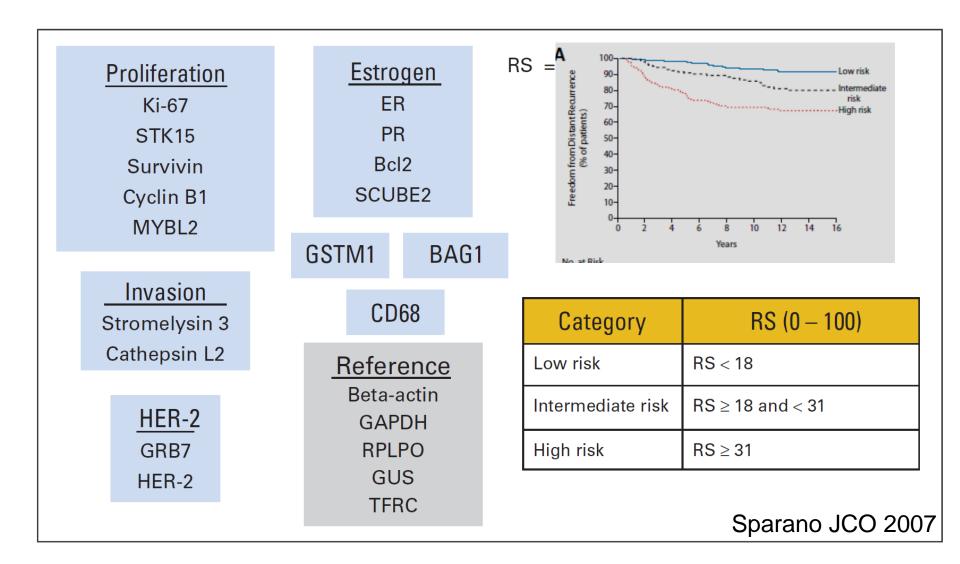
Müller et al., 2011

Recurrence Score - Oncotype DX

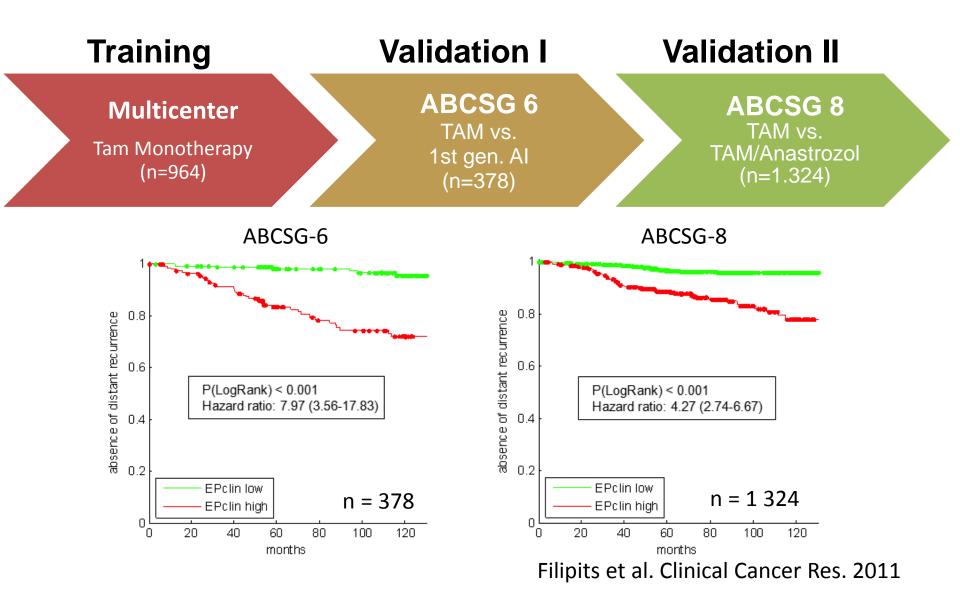
16 genes and 5 control genes



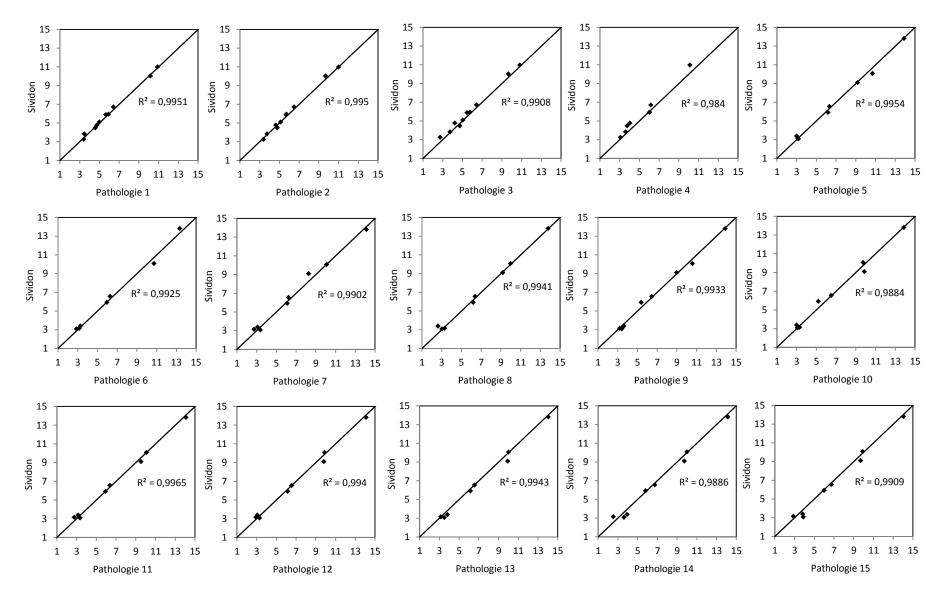
Recurrence Score - Oncotype DX 16 genes and 5 control genes



Development of the Endopredict assay



Endopredict Test – 15 different laboratories



DNA markers and mutation analysis

- DNA is more stable than mRNA
- can be isolated from FFPE tissue
- fragmentation occurs focus on analysis of small fragments

- classical Sanger sequencing
- NGS sequencing

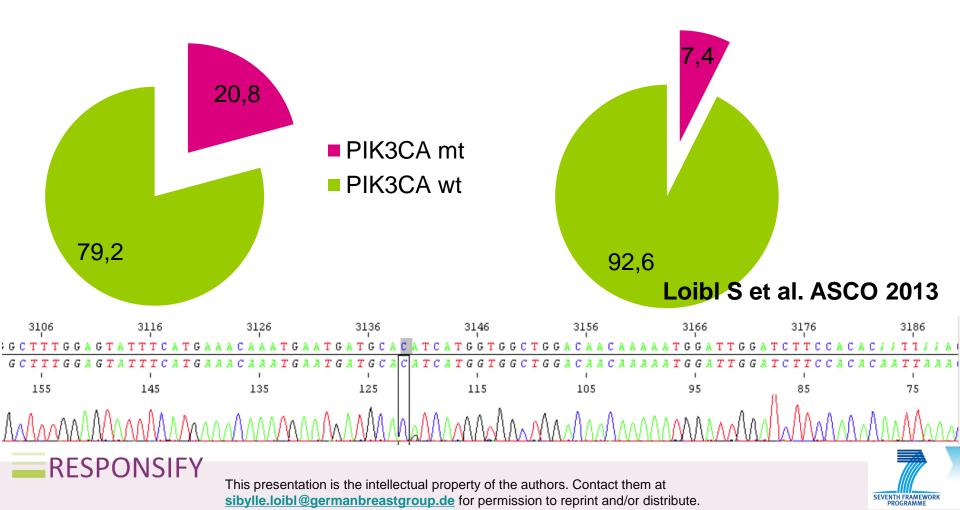


San Antonio Breast Cancer Symposium - Cancer Therapy and Research Center at UT Health Science Center – December 10-14, 2013



PIK3CA Mutation analysis (Exon 9 & 20) in FFPE samples of the GeparQuinto and GeparSixto study

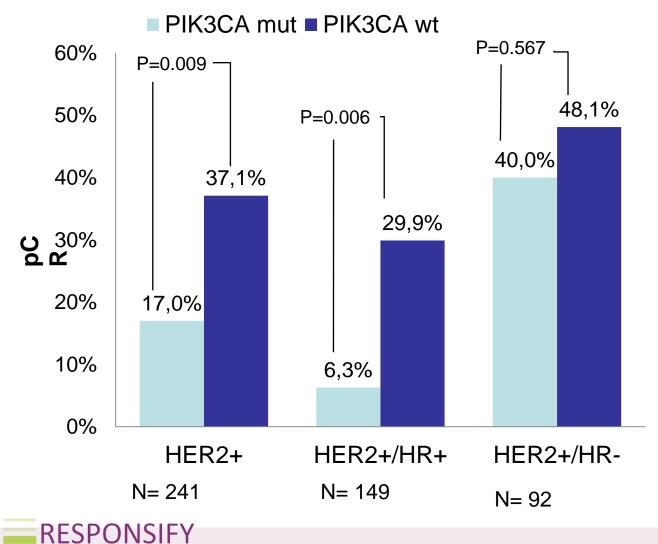
HER2+ tumours n=360 triple-negative tumours n=285







pCR rate according to PIK3CA mutation status in GeparSixto study



Multivariate

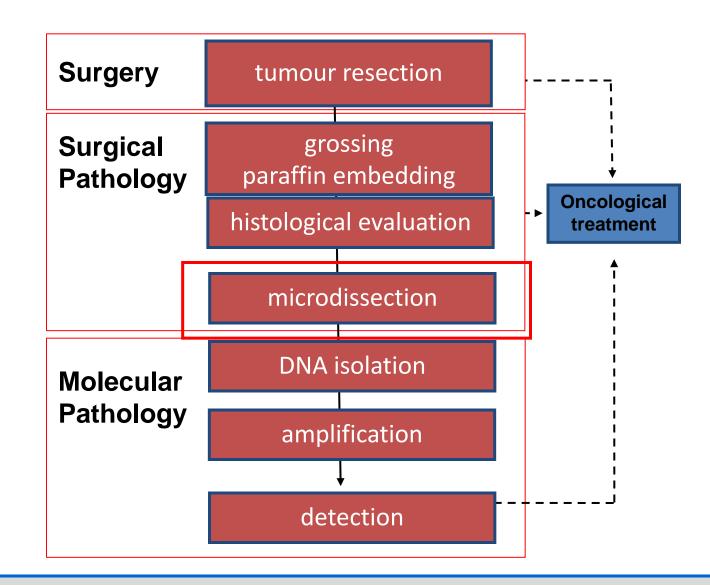
		Odds <u>ratio</u>	P- value
HR statu s	neg	1.00	0.006
	pos	0.44 (0.24-0.79)	
PIK3 CA	wt	1.00	0.007
	mut	0.29 (0.12-0.71)	

adjusted for therapy, age, tumour and nodal status, histotype and grading



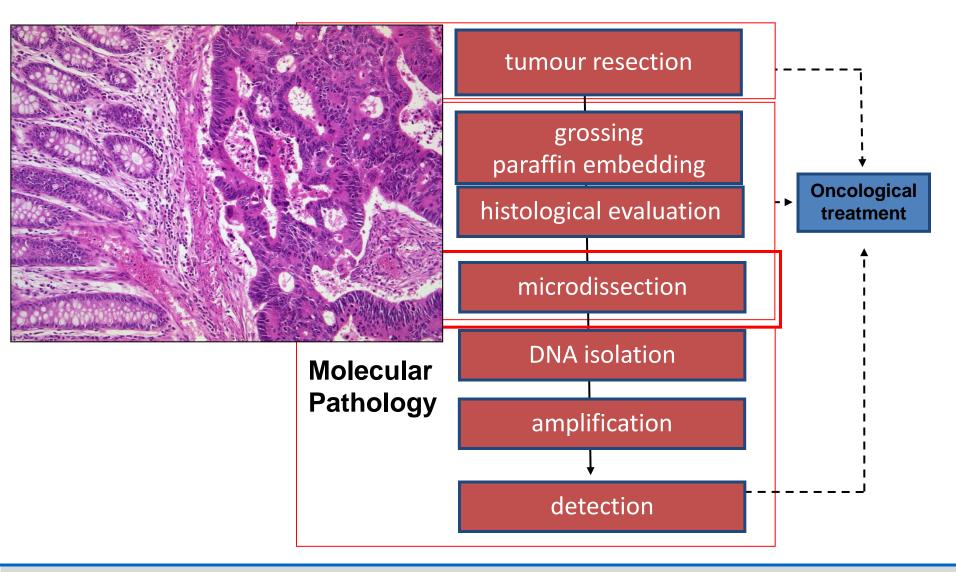
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Workflow in molecular pathology





Workflow in molecular pathology



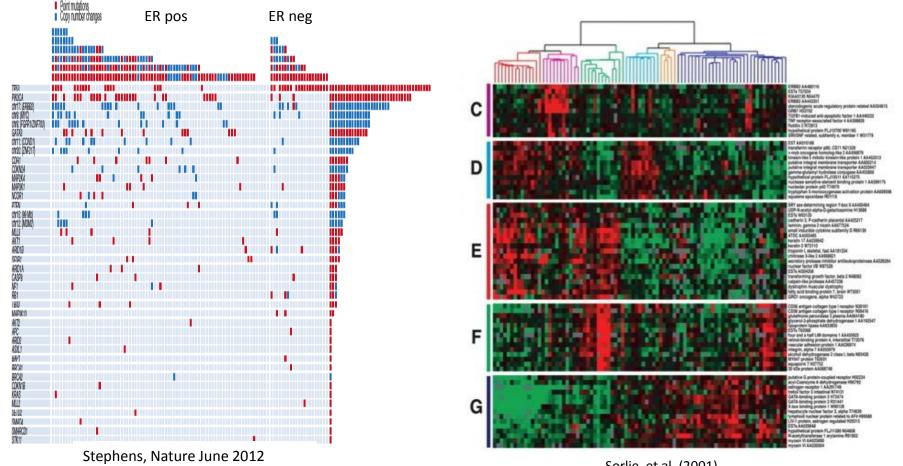


Next generation sequencing

- NGS: all genes / transcripts in a tumor are measured
- overview on all genetic alterations in a tumor
 - Mutations
 - Copy number variations
 - Amplifikations
 - Deletions
- applicable to routine pathology: targeted exome sequencing

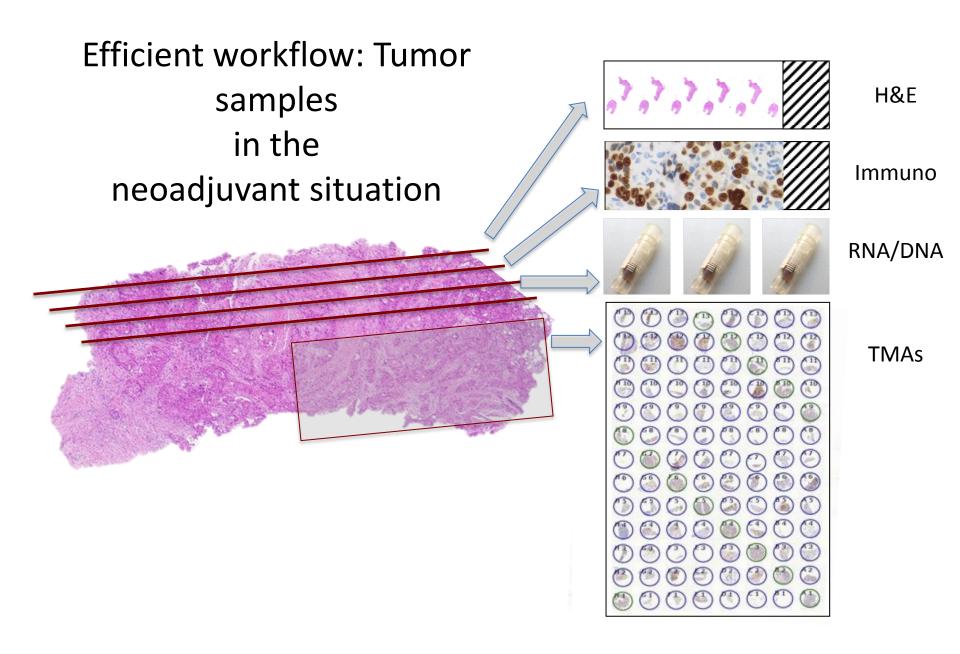


High throughput technologies - mRNA and DNA alterations in breast cancer



Sorlie, et al. (2001) Proc. Natl. Acad. Sci. USA 98, 10869-10874





GBG GERMAN BREAST GROUP

Conclusion – methods in pathology

- FFPE tissue is suitable for analysis of protein, mRNA and DNA markers
- targeted exome sequencing as an upcoming method in routine pathology
- critical parameters are:
 - standardisation and quantification for immunohistochemistry
 - selection of primers for mRNA / DNA analysis
 - selection of tissue area for mRNA / DNA analysis
- requirements for practice changing research in pathology: high level of evidence, "prospectiveretrospetive" studies, large sample collections

