

- < テーマ > 治療効果予測と予後予測～乳がんのターゲット治療の模索～
乳がんにおける予後及び治療効果の予測検査法について、欧・米・東アジアからの招聘者と共に検討する国際セミナーです。
- < 場所 > 東京国際フォーラム ホールB5



The Current Status and the Future Prospects of Multigene testing in Europe

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NKI-AVL



St. Gallen Recommendations 2009
= 'Europe' (?)

St Gallen Recommendations 2007 for Adjuvant Treatment

	Highly endocrine responsive	Incompletely endocrine responsive	Endocrine non-responsive
HER2-negative	ET (consider adding CT according to risk)	ET (consider adding CT according to risk)	CT
HER2-positive	ET + Trastuzumab + CT	ET + Trastuzumab + CT	ET + Trastuzumab + CT

endocrine highly and incompletely responsive HER2neg patients consider adding chemotherapy according to risk

ET: endocrine therapy
CT: chemotherapy



Intermediate Risk Treatment Advice

	Highly endocrine responsive	Incompletely endocrine responsive	Endocrine non-responsive
HER2-negative	ET (consider adding CT according to risk)	ET (consider adding CT according to risk)	CT
HER2-positive	ET + Trastuzumab + CT	ET + Trastuzumab + CT	ET + Trastuzumab + CT

- St Gallen 2009: The panel accepts the use of validated molecular based tools, if readily available, as an adjunct to high quality standard histopathologic assessment in patients with ER+ breast cancer when the doctor and the patient are uncertain or ambivalent about the administration of adjuvant chemotherapy. Optimally the test should be used in clinical trials.

- Yes 80%
- No 18%
- unknown 3%



Guidelines: St Gallen International Expert Consensus 2009

Table 3. Chemoendocrine therapy in patients with ER-positive, HER2-negative disease

<i>Clinicopathological Features</i>			
	Relative Indications for Chemoendocrine therapy	Factors Not Useful for Decision	Relative Indications for Endocrine Therapy Alone
ER, PgR	Lower ER and PgR level		Higher ER and PgR level
Histological Grade	Grade 3	Grade 2	Grade 1
Proliferation	High ^a	Intermediate ^a	Low ^a
Nodes	Node positive (4 or more involved nodes)	Node positive (1-3 involved nodes)	Node negative
Peritumoral Vascular Invasion (PVI)	Presence of extensive PVI		Absence of extensive PVI
pT-size	> 5cm	2.1 – 5 cm	≤ 2cm
Patient Preference	Use all available treatments		Avoid side effects
<i>Multi-gene Assays</i>			
Gene Signature^b	High score	Intermediate score	Low score

St. Gallen Recommendations on March 14th, 2009



MammaPrint Accepted into St. Gallen's Oncology Guidelines for Early Stage Breast Cancer Treatment

“The Panel accepts the use of validated molecularly based tools if readily available as an adjunct to high-quality standard histopathologic assessment in patients with ER+ breast cancer when the doctor and patient are uncertain or ambivalent about the administration of adjunctive chemotherapy.”

In addition, the Panel felt “intermediate” results were of little clinical value.

St. Gallen Guideline Consensus, Annals of Oncology, 2009

Guidelines: St Gallen International Expert Consensus 2009

Table 3. Chemoendocrine therapy in patients with ER-positive, HER2-negative disease

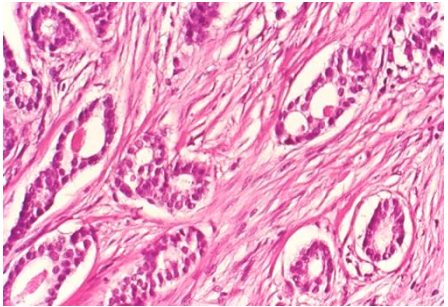
<i>Clinicopathological Features</i>			
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<i>Multi-gene Assays</i>			
Gene Signature^b 21 recurrence score	High score	Intermediate score	Low score

70 gene prognosis signature

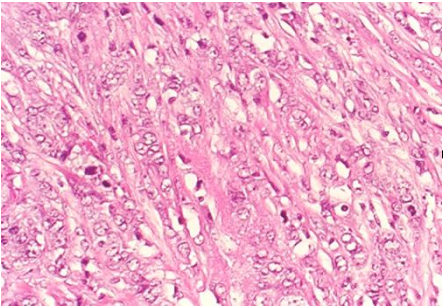
Contrast of Appearance and Expression Phenotyping



Microscope



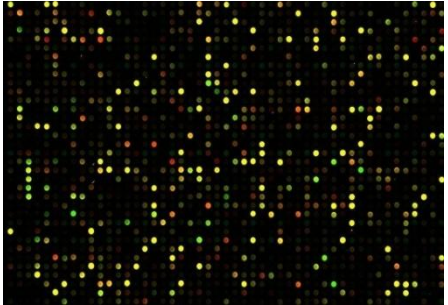
Low Grade



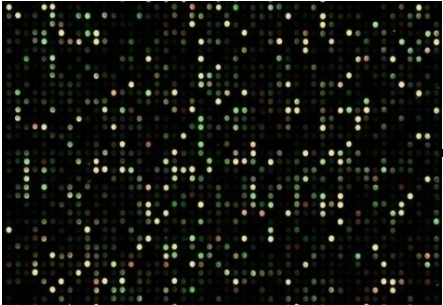
High Grade



Microarray



Low Risk



High Risk

Treatment Advice

21-Gene Recurrence Score (RS) Assay Oncotype DX (Genomic Health)

16 Cancer and 5 Reference Genes From 3 Studies

PROLIFERATION

Ki-67
STK15
Survivin
Cyclin B1
MYBL2

ESTROGEN

ER
PR
Bcl2
SCUBE2

GSTM1

CD68

BAG1

INVASION

Stromolysin 3
Cathepsin L2

HER2

GRB7
HER2

REFERENCE

Beta-actin
GAPDH GUS
RPLPO TFRC

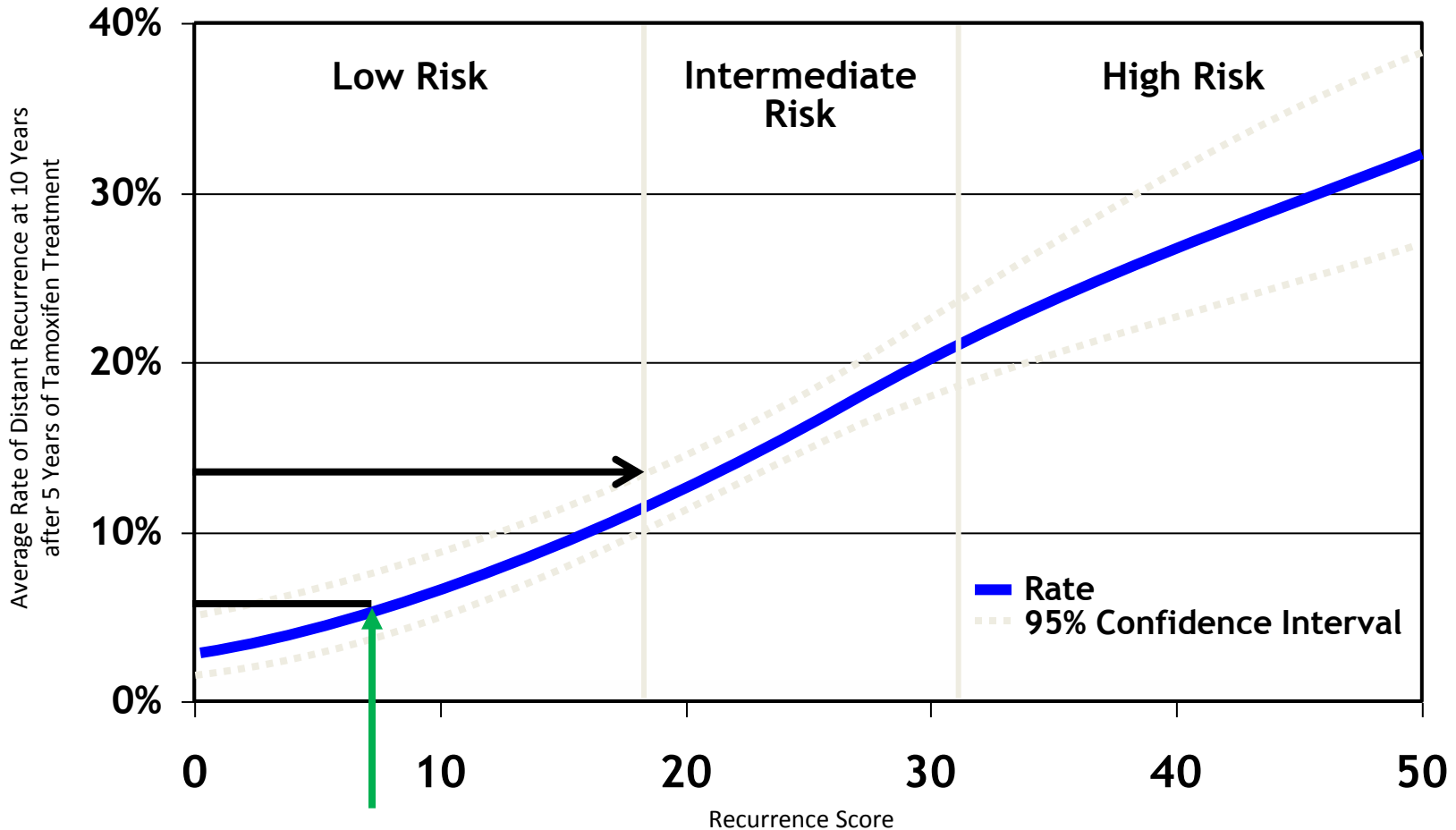
RS Weighting:

+ 0.47 x HER2 Group
- 0.34 x ER Group
+1.04 x Proliferation Group
+ 0.10 x Invasion Group
+ 0.05 x CD68
- 0.08 x GSTM1
- 0.07 x BAG1

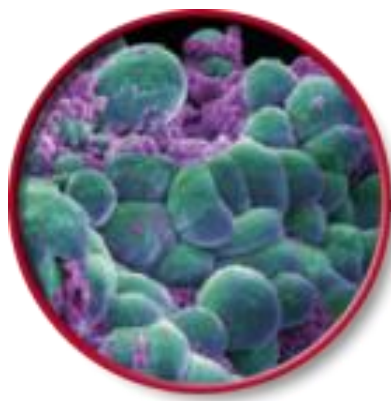
Category	RS (0 - 100)
Low risk	RS < 18
Intermediate risk	RS ≥ 18 and < 31
High risk	RS ≥ 31

The Recurrence Score

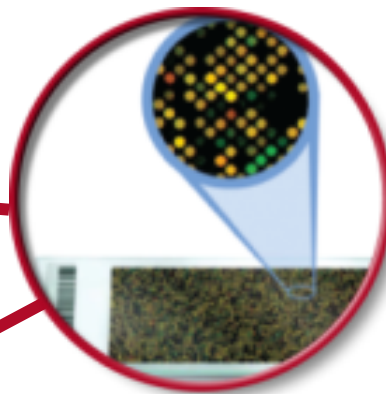
The following results are from a clinical validation study with prospectively-defined endpoints involving 668 patients. The patients enrolled in the study were female, stage I or II, node-negative, ER-positive, and treated with tamoxifen. *N Engl J Med* 2004; 351:2817-26.



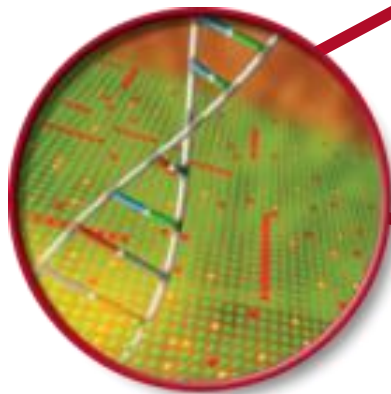
MammaPrint[®]: Enhanced Scientific Guidance Through Unbiased Gene Selection



Untreated Patient
Breast Tumor Samples

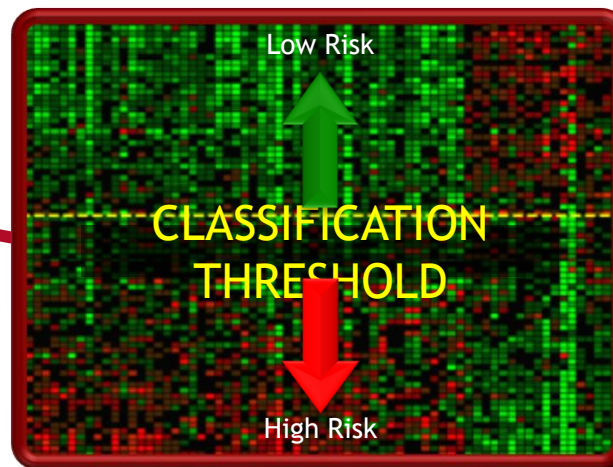


Gene Expression Analysis
of Entire Human Genome
~25,000 Genes



231 Prognostic Breast
Cancer Genes Identified

70 Most
Prognostic Genes



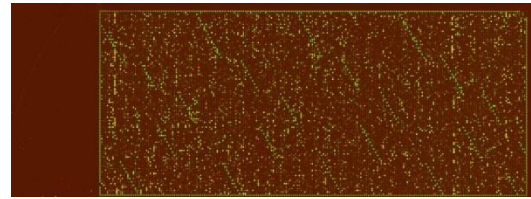
High Risk or Low Risk,
No Intermediates

70-gene MammaPrint

- Is not just another prognostic factor
- Is designed from the beginning to tell you the metastatic potential of an individual breast cancer

Development of 70 gene prognosis signature

Tumor samples of known
clinical outcome



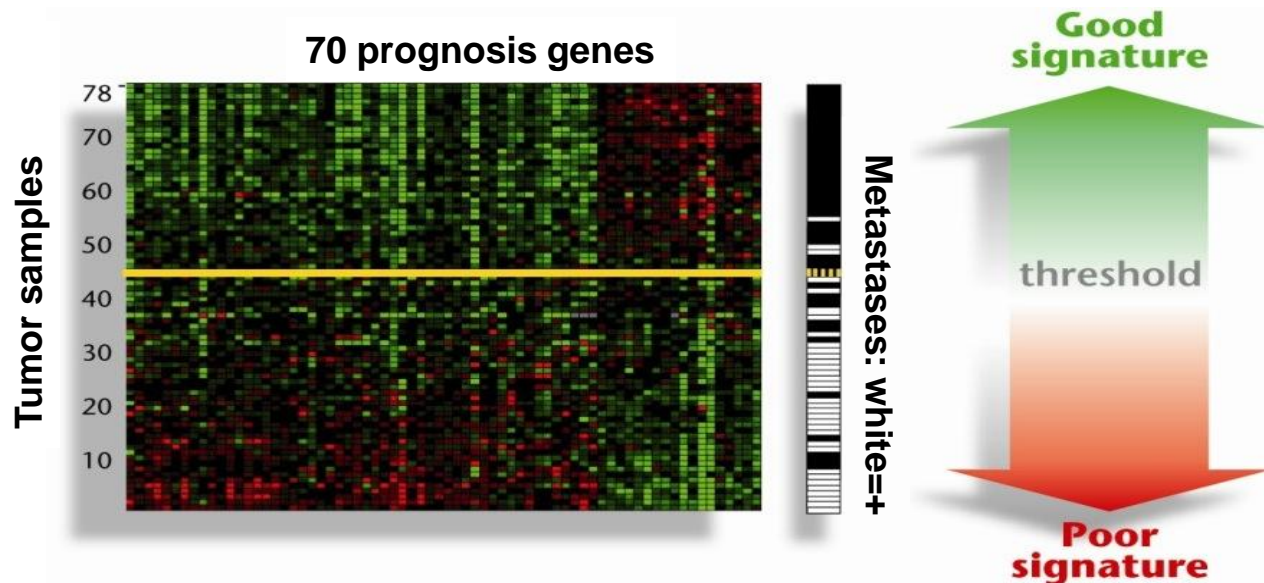
Unbiased full genome
gene expression
analysis

Prognosis reporter genes

Distant metastases
group

No distant metastases
group

b
















70-gene MammaPrint

- Function of majority of genes is identified and are all related to the process of dissemination

Prognostic value of the 70-gene assay

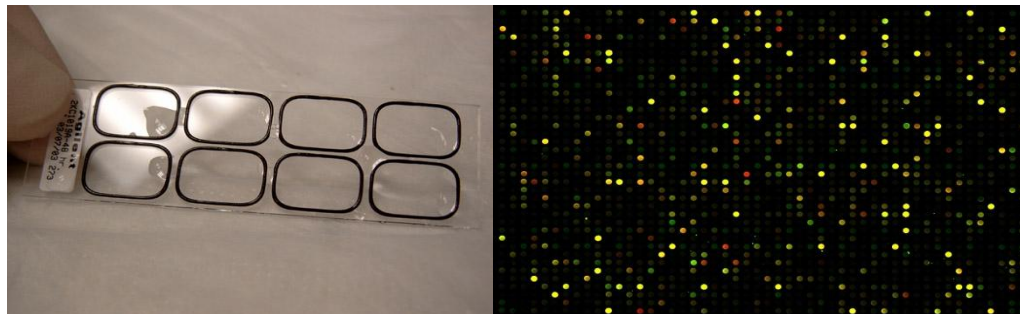
- Biologically plausible
- Better compared to conventional criteria (multivariate-analysis)
- Validated in 8 independent series

MammaPrint validation studies > 2500 patients

Validierungsstudie	Land	Reference	Jahr				
			2006	2007	2008	2009	2010
Independent European study		Buyse et al J NCI 17	302				
Prospective Study		de Mesquita et al. Lancet Oncology		427			
Dutch patient cohort		de Mesquita Breast Cancer Res Treat			123		
Core Needle biopsies		Mayordomo et al. ESMO Meeting			35		
Validation in US patients		Wittner et al. Clin Cancer Res 14			100		
Validation 1-3 LN+ patients		Mook et al. Breast Cancer Res Treat.			241		
Postmenopausal patients		Mook et al. Breast Cancer Res Treat			148		
Patients treated w Tamoxifen		Kok et al. (submitted)				192	
German patient cohort		Kunz et al. St. Gallen Conference				140	
Japanese patient cohort		Ishitobi et al. Jap J Clin Oncology				118	
Validation 4-9 LN+ patients		Saghastchian et al. St. Gallen Conf				167	
Neoadjuvant predictive study		Straver et al. Breast Cancer Res Treat				162	
Predictiveness study		Knauer et al. Breast Cancer Res Treat					541

MammaPrint from Research to Diagnostics

- Retrospective validation
 - Prospective Technology assessment
 - Diagnostic test
 - Laboratory
 - Diagnostic test
 - Diagnostic test
 - Diagnostic test and clinical use
 - Treatment Recommendations
 - Treatment Recommendations
- Completed
 - Utility & Cost-effectiveness
 - International CE marked
 - CLIA registered
 - ISO17025 certified
 - CAP accredited
 - FDA approved, IVD MIA feb07
 - Dutch Guidelines 08
 - StGallen International Guidelines 09



Reproducibility
Test Result >98%
Success rate >95%

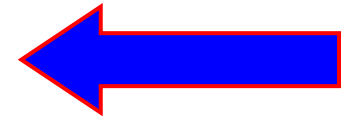
Glas et al,
BMC Genomics 200

Validation 1: N = 151

vd Vijver et al, N Engl J Med 347: 1999-2009, 2002

TABLE 4. MULTIVARIABLE PROPORTIONAL-HAZARDS ANALYSIS OF THE RISK OF DISTANT METASTASES AS A FIRST EVENT.

VARIABLE	HAZARD RATIO (95% CI)*	P VALUE
Poor-prognosis signature (vs. good-prognosis signature)	4.6 (2.3–9.2)	<0.001
Age (per 10-yr increment)	0.73 (0.50–1.06)	0.10
Lymph-node status (per positive node)	1.13 (1.03–1.24)	0.01
Diameter of tumor (per cm)	1.56 (1.22–2.0)	<0.001
Tumor grade		0.54
Grade 2 (vs. grade 1)	1.35 (0.61–3.0)	
Grade 3 (vs. grade 1)	1.03 (0.44–2.4)	
Vascular invasion		0.05
1–3 Vessels (vs. 0 vessels)	0.66 (0.30–1.44)	
>3 Vessels (vs. 0 vessels)	1.65 (0.98–2.8)	
Estrogen-receptor expression (per point)†	0.86 (0.56–1.31)	0.48
Mastectomy (vs. breast-conserving therapy)	1.27 (0.79–2.0)	0.32
Chemotherapy (vs. no chemotherapy)	0.37 (0.20–0.66)	<0.001
Hormonal treatment (vs. no hormonal treatment)	0.62 (0.29–1.34)	0.23



Validation 2: N = 307

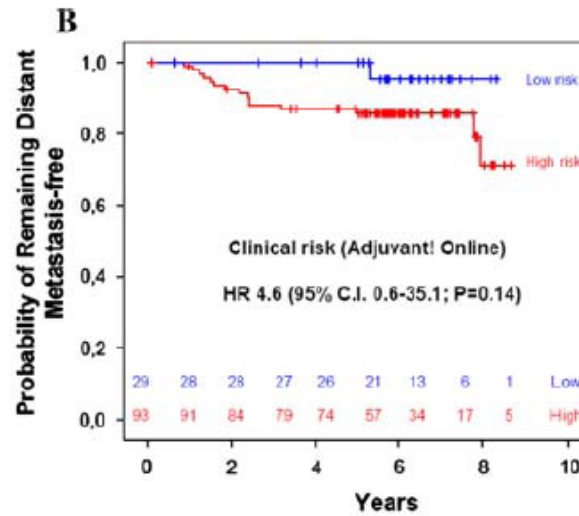
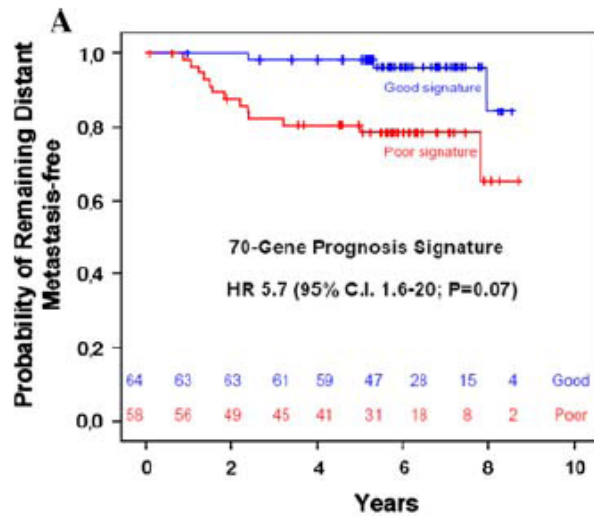
Buyse M, et al.: J Natl Cancer Inst 98: 1183-92, 2006

Risk factor or classification	Time to distant metastases
Age (≤ 50 y versus > 50 y)	0.86 (0.54 to 1.37) $P = .52$
Tumor size (T2 versus T1)	1.42 (0.90 to 2.23) $P = .14$
Tumor grade (good versus intermediate versus poor differentiation)	0.76 (0.54 to 1.07) $P = .12$
Estrogen receptor status (negative versus positive)	2.18 (1.37 to 3.48) $P = .001$
Adjuvant! software (high risk versus low risk)	1.68 (0.92 to 3.07) $P = .092$
Nottingham Prognostic Index (high risk versus low risk) \ddagger	1.65 (1.02 to 2.66) $P = .043$
St Gallen criteria (high risk versus low risk) \S	2.22 (0.70 to 7.08) $P = .18$
Gene signature (high risk versus low risk) \parallel	2.32 (1.35 to 4.00) $P = .002$



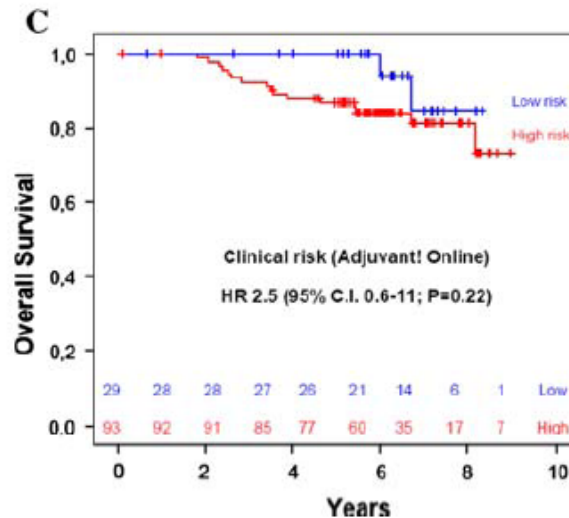
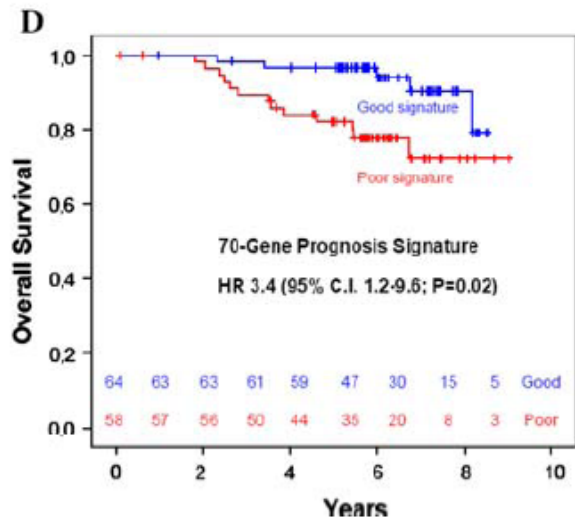
Validation 3: N = 123

Bueno-de-Mesquita JM: Breast Cancer Res Treatm 2008



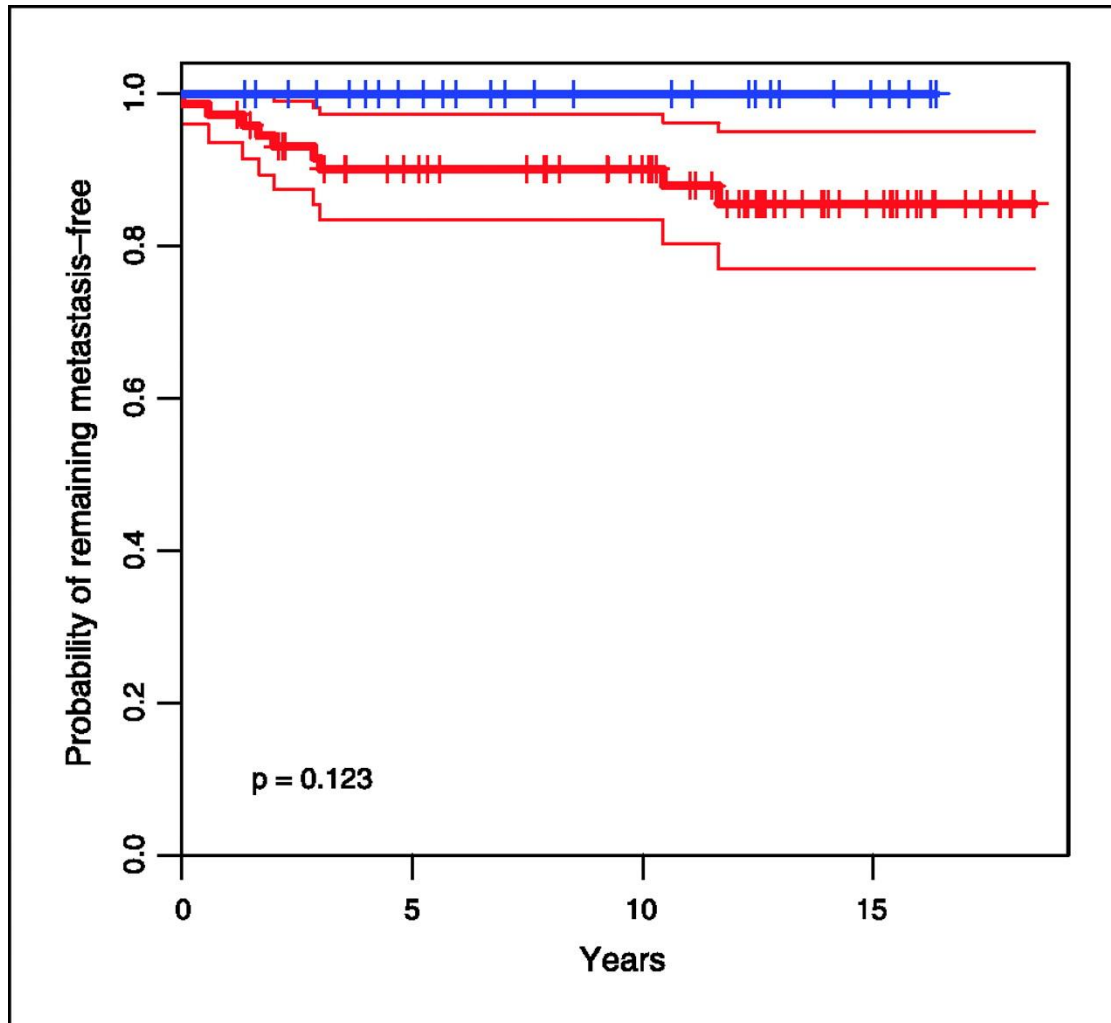
The 70-gene Signature Outperforms:

- Adjuvant Online
- St Gallen criteria
- Nottingham PI
- CBO guidelines



Validation 4: N = 100

Wittner et al., Clin Cancer Res 14: 2988, 2008



**MGH series,
Boston;**

Time to metastasis

St.Gallen RISK Categories

1696 patients analyzed from pooled database

	All endocrine responsive patients	Highly/incompletely responsive	All patients
Low risk	145	141	145
Intermediate risk	1020	773	1287
High risk	91	85	190
Total	1256/1696 = 74%	999/1696 = 59%	1622/1696 = 96%

Median follow-up 7.08 years \pm 5.02 (0.01 – 25.22)



Added value to assess risk in intermediate category by MammaPrint

Are they all at 'intermediate' risk?

A large meta analysis

Pooled analysis MammaPrint

1696 patients with MammaPrint

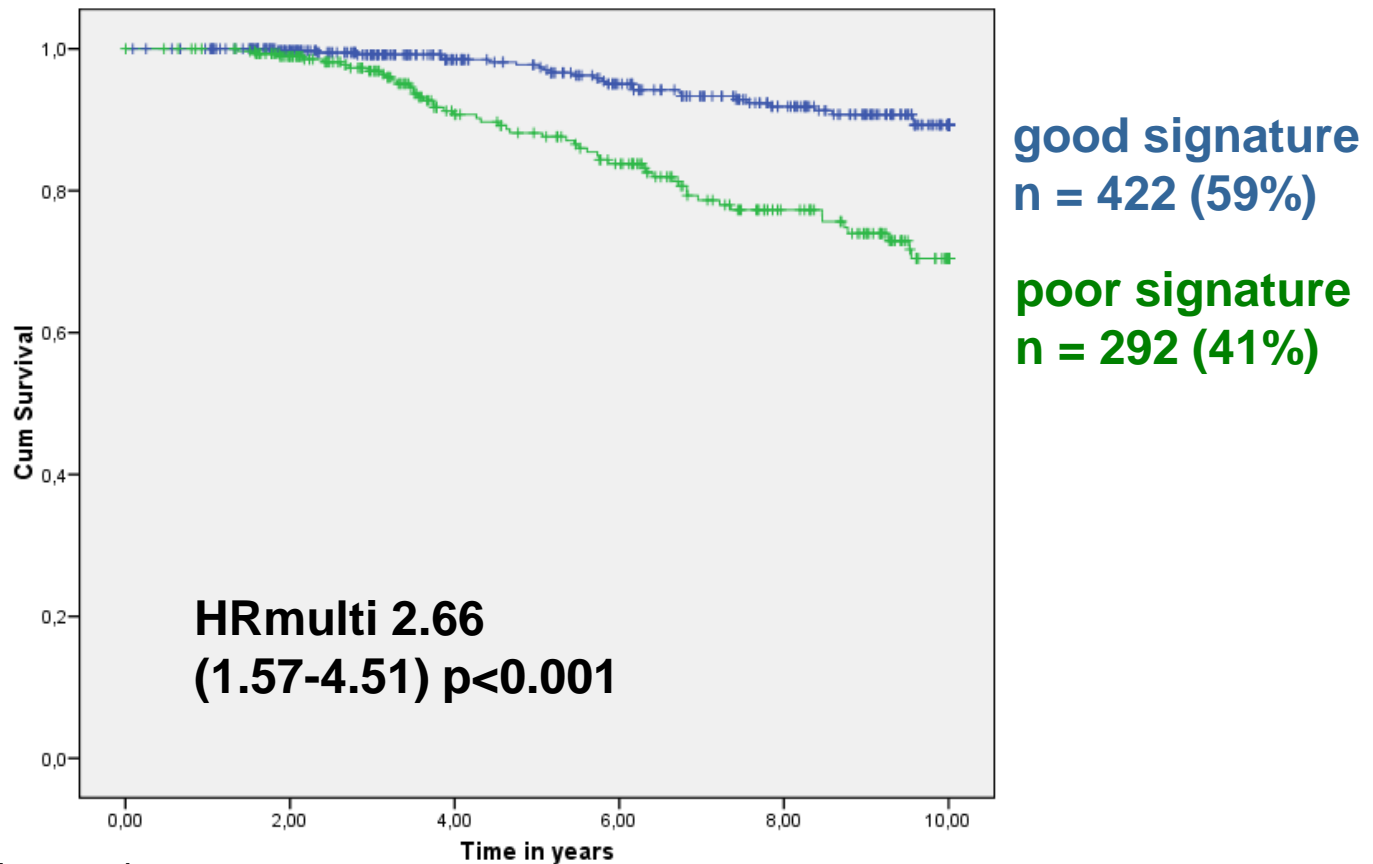
7 studies:

295 ptn van de Vijver et al	(<53, LN0/LN+)	(NEJM, 2002)
302 ptn Buyse et al.	(Transbig Int, <60, LN0)	(JNCI, 2006)
427 ptn Bueno et al.	(RASTER prosp, <60, LN0)	(Lancet Oncol, 2007)
123 ptn Bueno et al.	(Recent, LNneg)	(Br Can Res Tr, 2008)
241 ptn Mook et al.	(1-3 LNpos)	(Br Can Res Tr, 2008)
148 ptn Mook et al.	(age 55-70)	(SABCS 2007, #1063)
160 ptn Kok et al.	(adj tamoxifen)	(unpublished)

Median follow-up 7.08 years (0.01 – 25.22)

Intermediate Risk by MammaPrint

*Breast Cancer Specific Survival
for highly and incomplete endocrine responsive HER2 neg patients
n = 714*



Note: untreated and treated
Knauer et al, abstracts StGallen, ASCO and submitted 2009

Issues in early breast cancer

- Is good pathology as good?
- Small cancers good prognosis?
- Her 2 overexpression: chemo?
- MammaPrint and chemo-effect

Issues in early breast cancer

- Is good pathology as good?

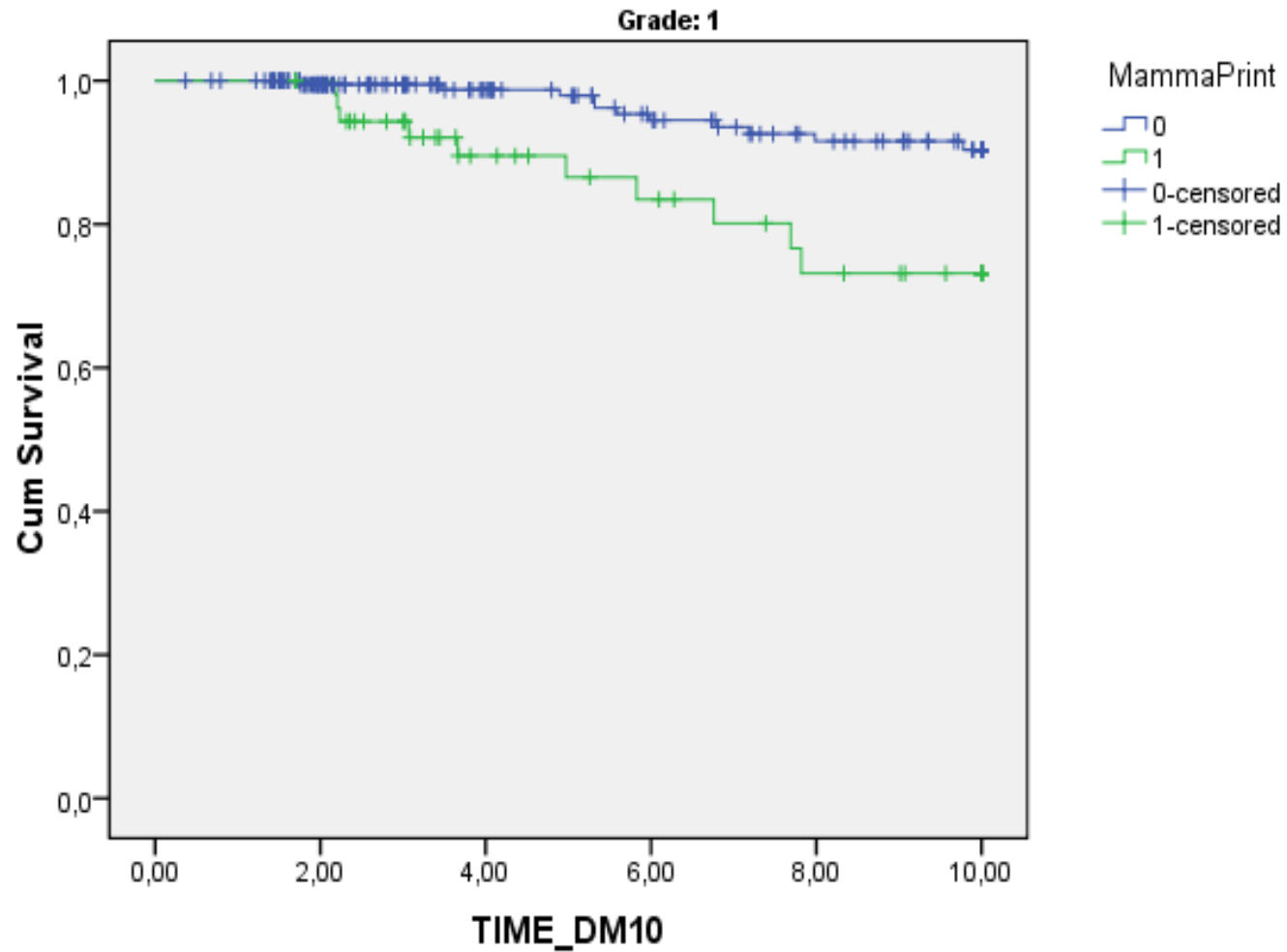
Is Grading the golden standard....

- Or can we do better?

Patients (n=965)	Characteristic	n (%)
Age	≤ 50 years	509 (53%)
	> 50 years	456 (47%)
Tumor size	T1a/b	140 (14%)
	T1c	825 (86%)
Lymph node status	Node negative	716 (74%)
	Node positive	241 (25%)
	n.a.	8 (1%)
Histological grade	Grade 1	280 (29%)
	Grade 2	412 (43%)
	Grade 3	262 (27%)
	n.a.	11 (1%)
Estrogen receptor status	Positive (≥10%)	808 (84%)
Progesterone receptor status	Positive (≥10%)	554 (57%)
Her2-status	Positive	91 (9%)
Adjuvant treatment	No adjuvant therapy	562 (59%)
	Endocrine therapy	182 (19%)
	Chemotherapy	100 (10%)
	Both	117 (12%)

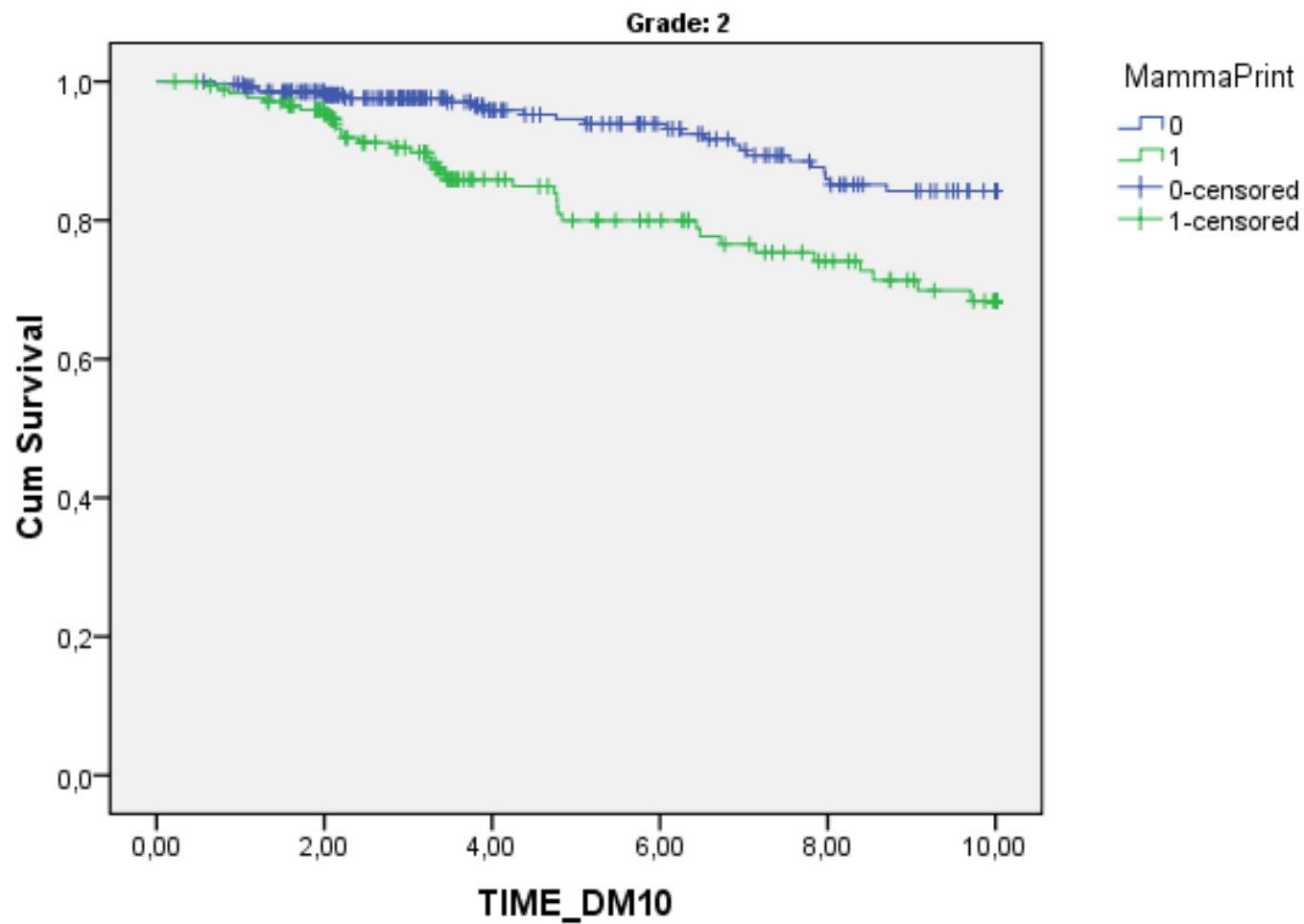
DDFS N0

Survival Functions



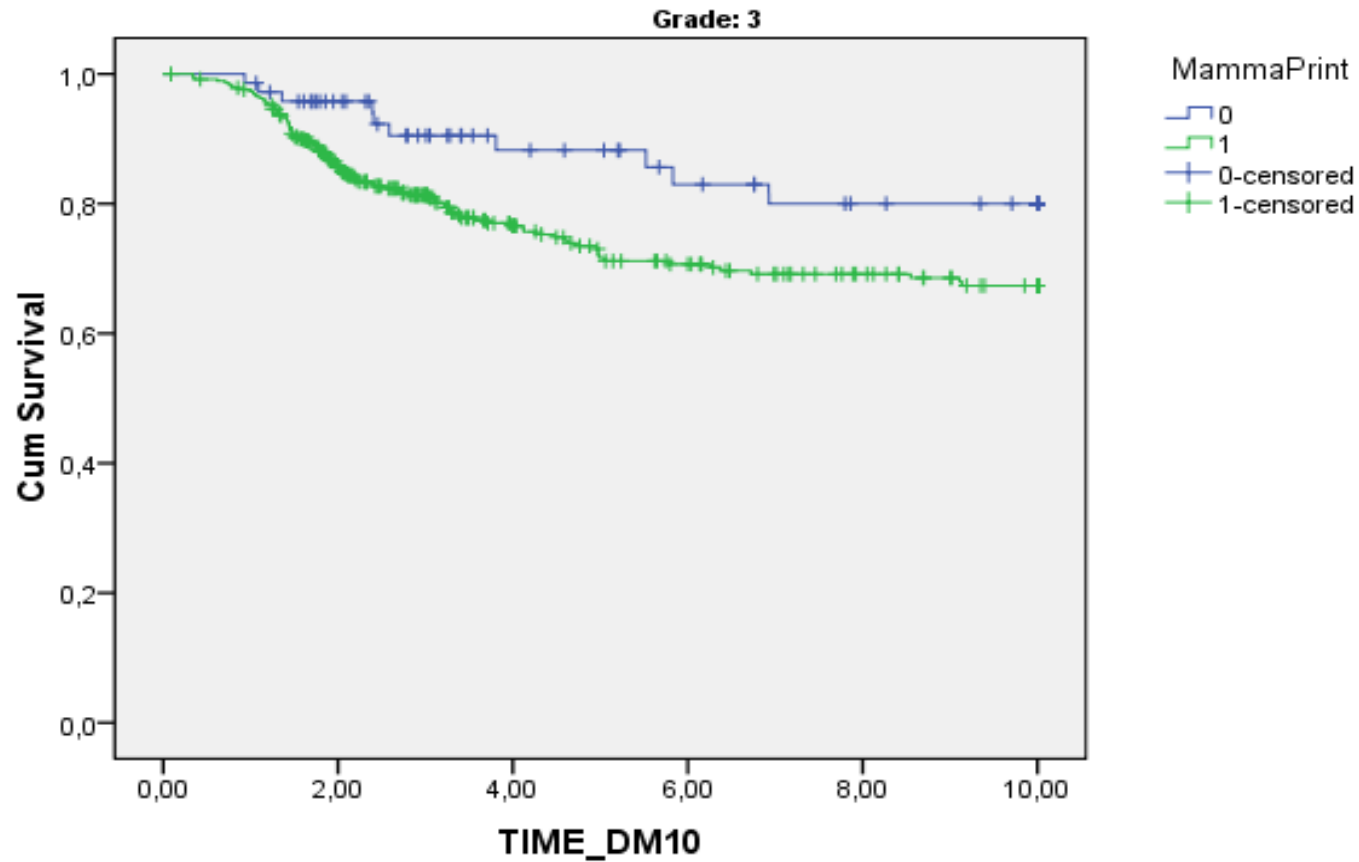
DDFS N0

Survival Functions

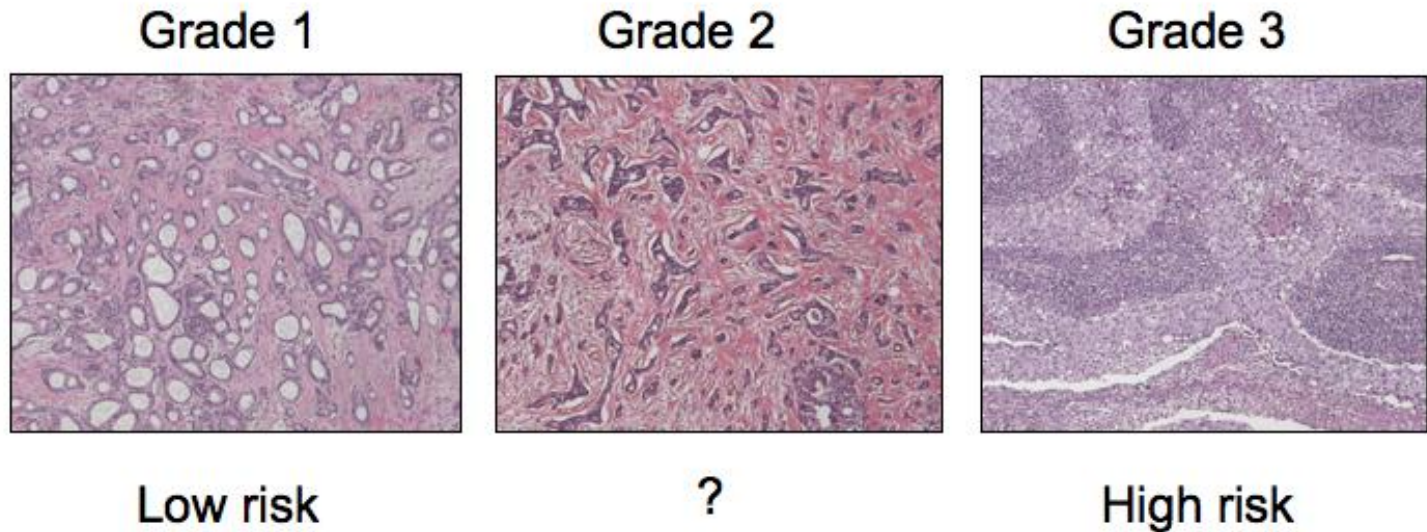


DDFS N0

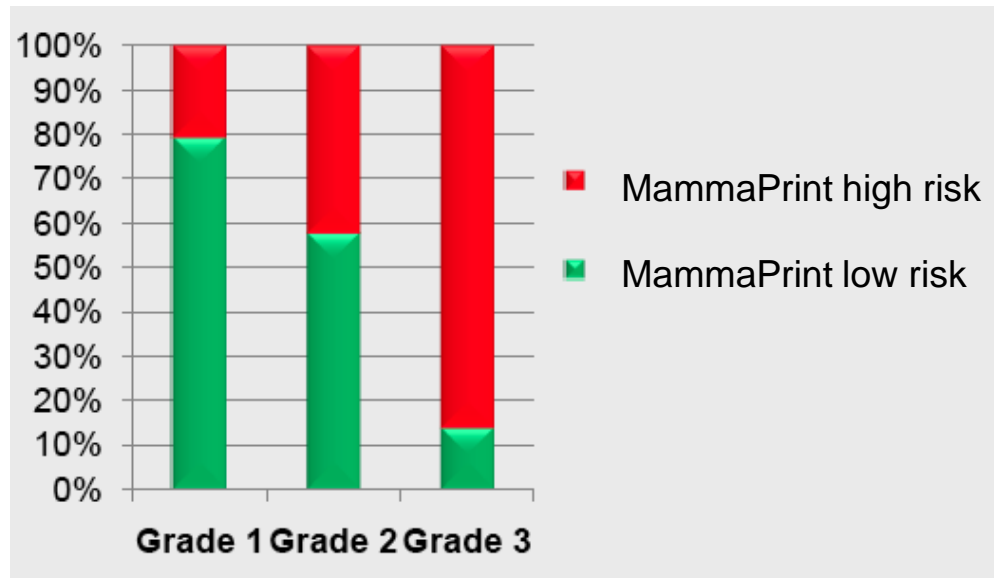
Survival Functions



MammaPrint adds to grading of breast cancer



764 of 1630 patients (47%) were classified as good prognosis and 866 (53%) as poor prognosis by MammaPrint
Histological grading was centrally reviewed for all patients



Issues in early breast cancer

- Small cancers good prognosis?

Patient inclusion criteria:

T1 breast cancer

Irrespective of

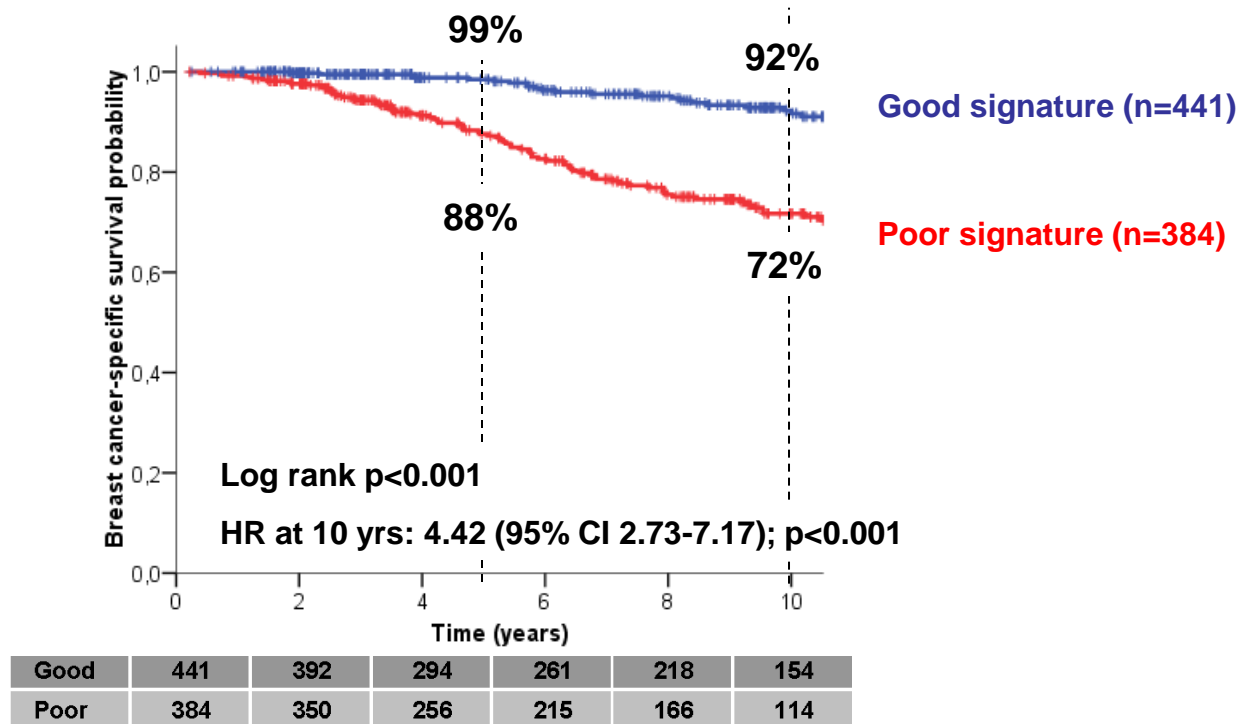
- Age
- Nodal status
- ER, PR, Her2-status

=> 965 patients

Median follow-up 7.1 years (0.2-25.2)

MammaPrint and Tumorsize T1c BCSS

11 – 22 mm Tumors

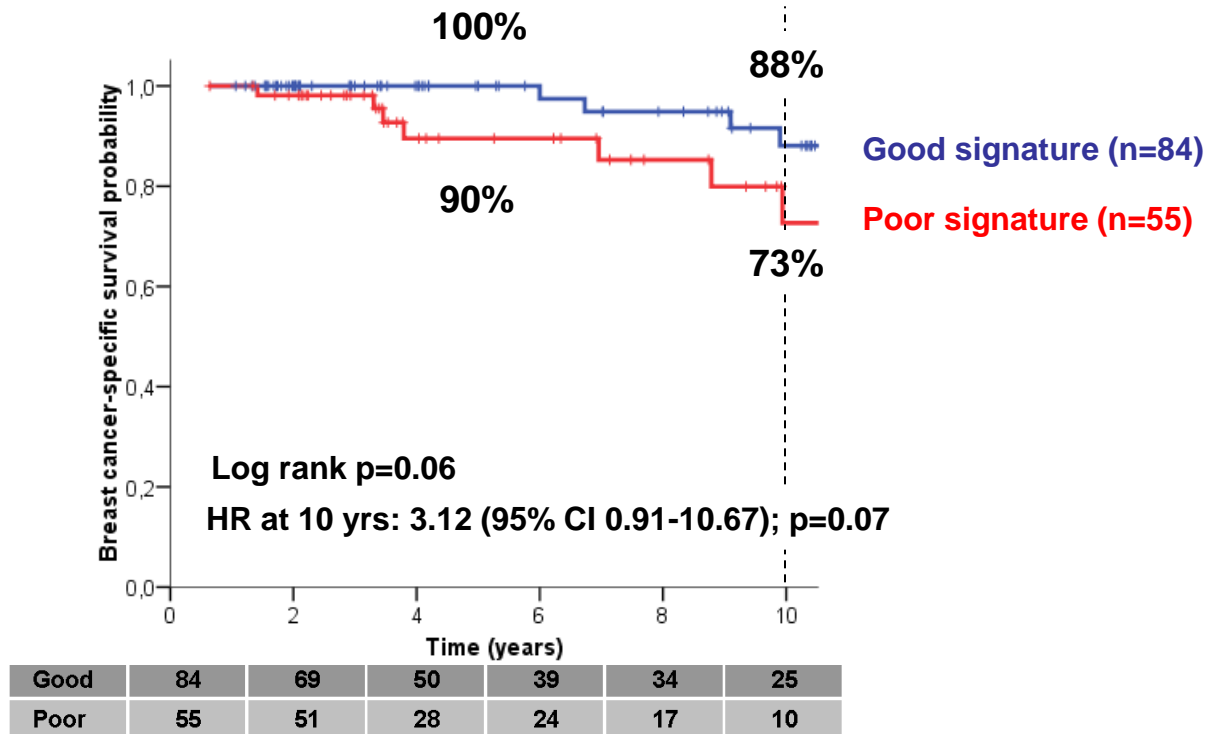


T1c tumors derived from pooled database of all MammaPrint validation studies (all, n=1696)

Mook et al, Ann Surg Oncol, 2010

MammaPrint and Tumorsize T1ab BCSS

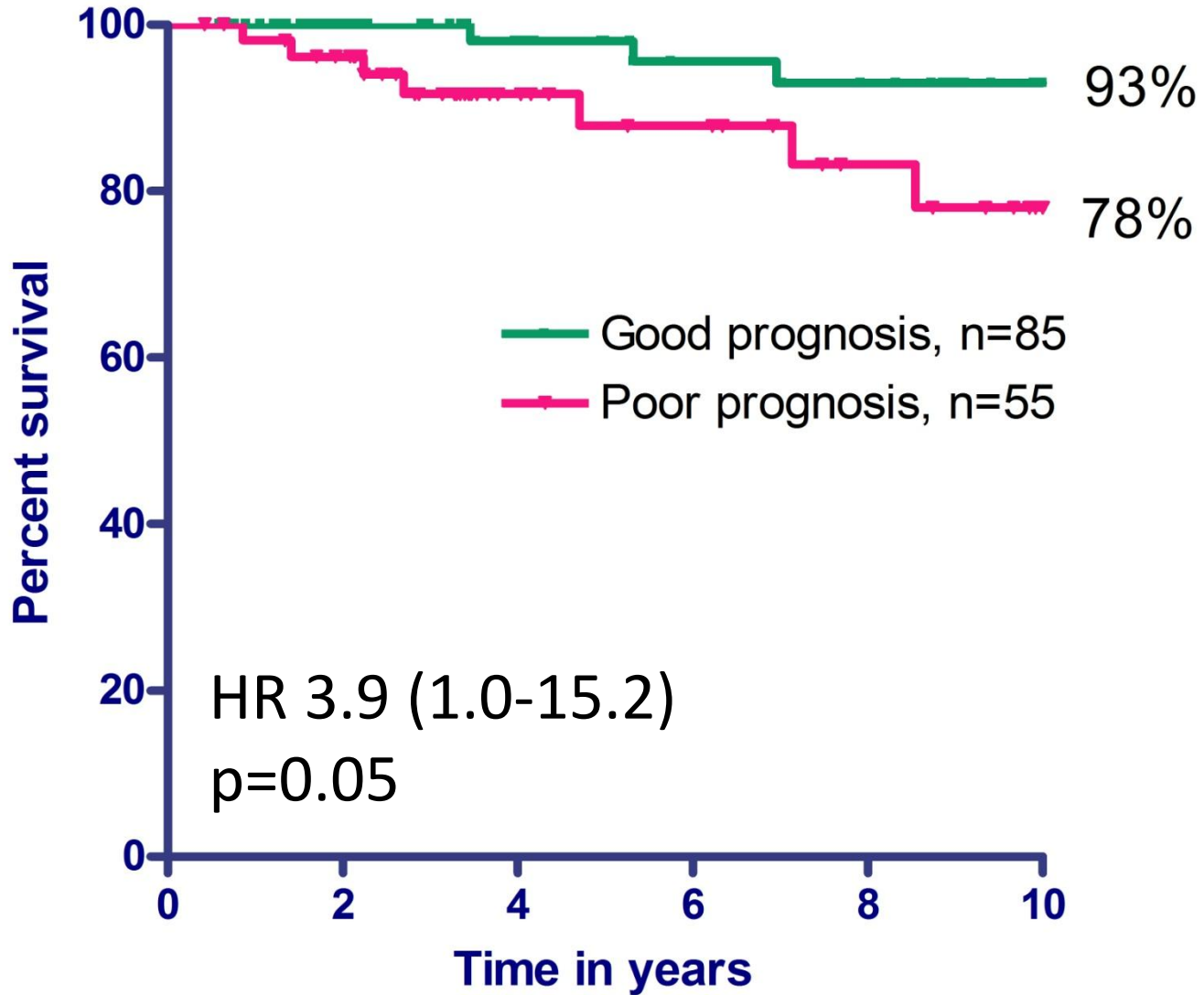
0 – 10 mm Tumors



T1ab tumors derived from pooled database of all MammaPrint validation studies (all, n=1696)

Mook et al, Ann Surg Oncol 2010

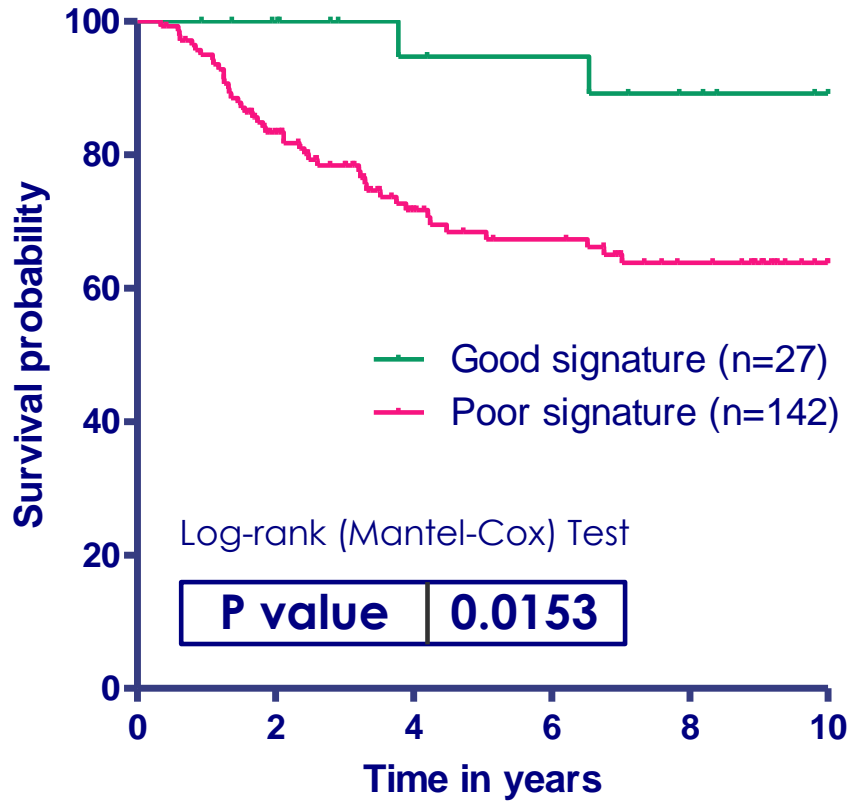
**DDFS: T1 a/b
(n=140)**



And Her-2 positive BC....

- Always poor prognosis?

DDFS: All patients



BCSS: All patients

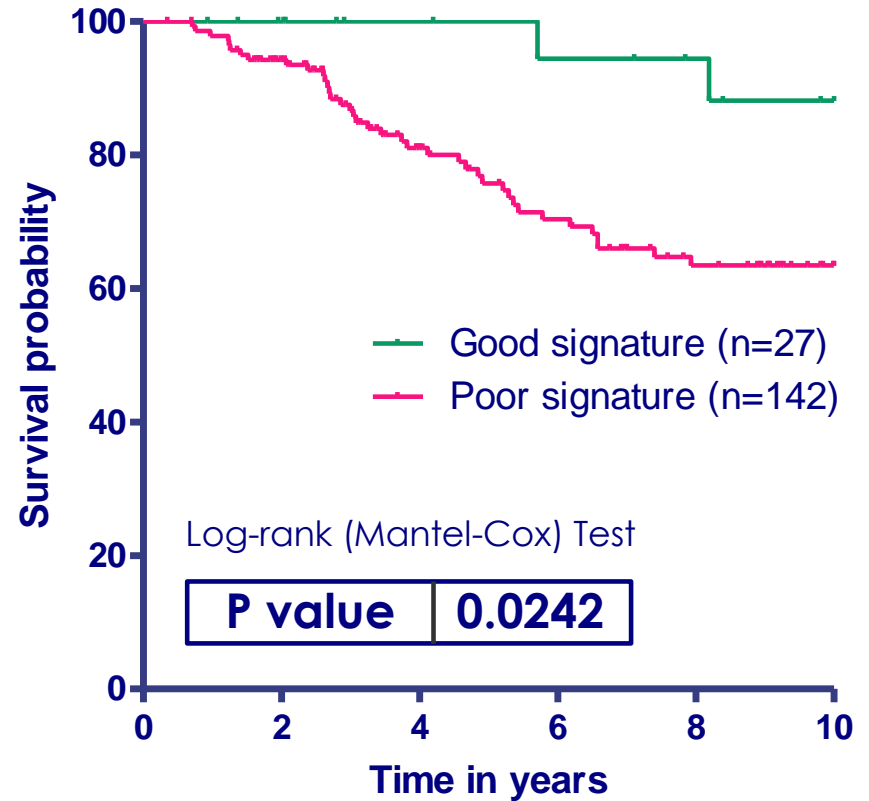
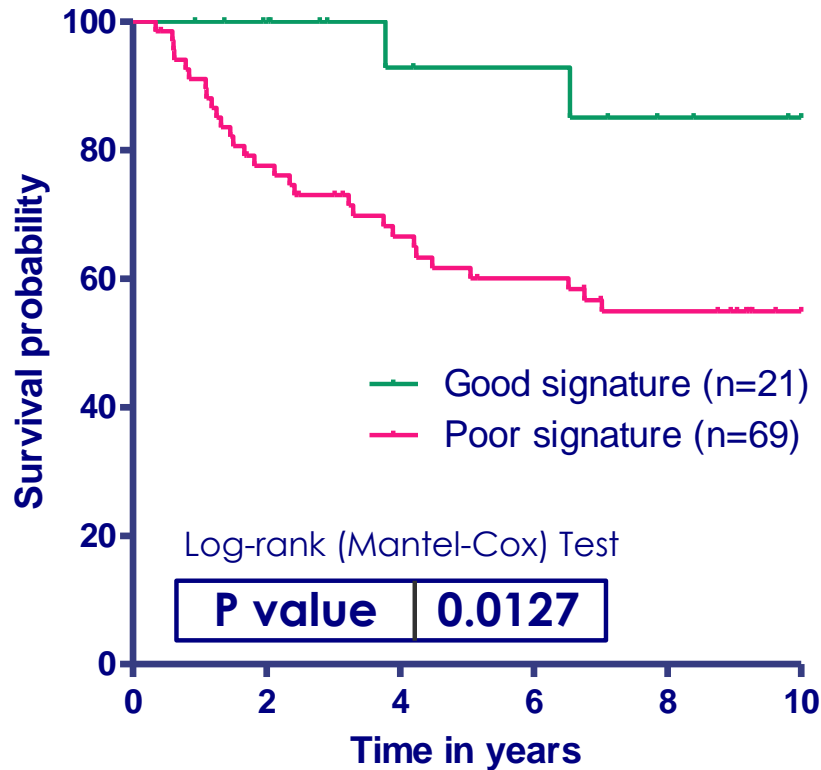


Figure 1:

Distant disease-free survival (LEFT) and breast cancer-specific survival (RIGHT) according to the 70-gene signature for all 169 Her2-positive breast cancer patients.

DDFS: without chemotherapy/trastuzumab



BCSS: without chemotherapy/trastuzumab

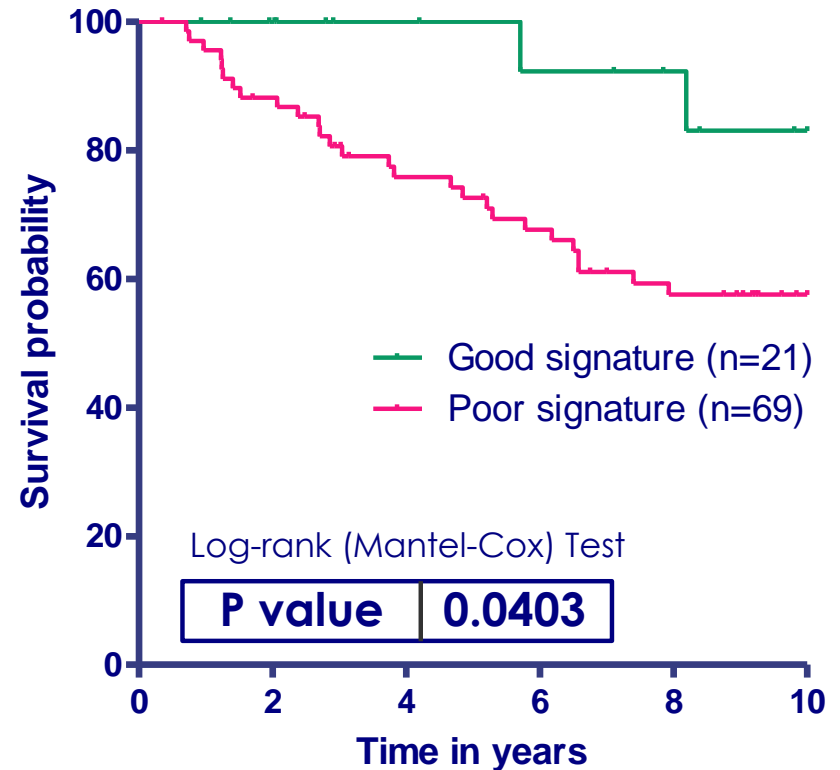
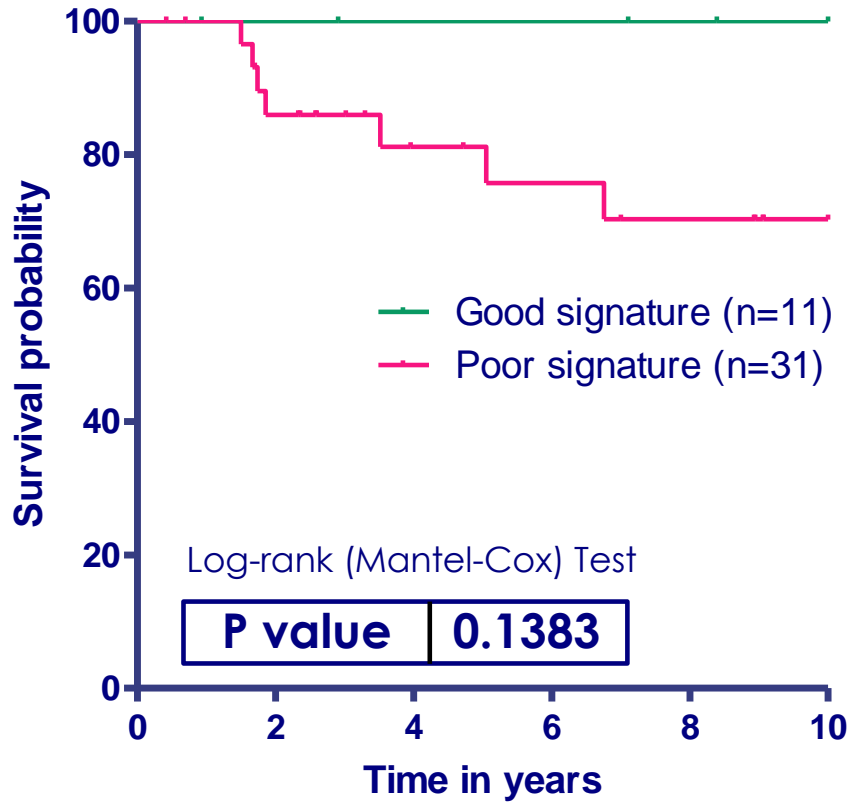


Figure 2:

Distant disease-free survival (LEFT) and breast cancer-specific survival (RIGHT) according to the 70-gene signature for 90 patients without adjuvant chemotherapy or trastuzumab.

DDFS: highly endocrine responsive



BCSS: highly endocrine responsive

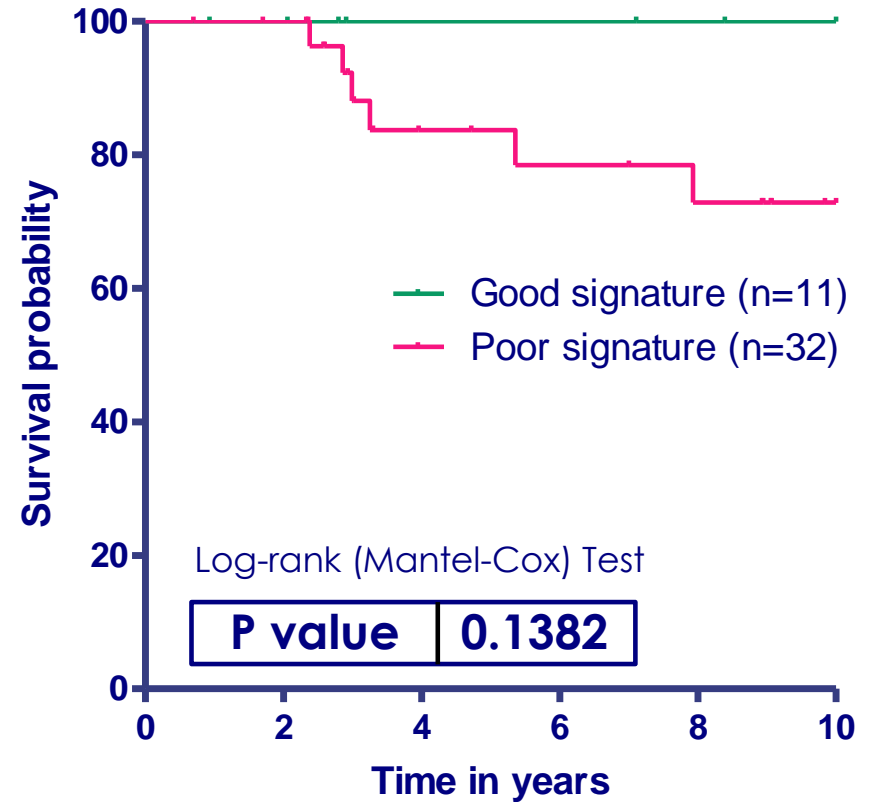


Figure 3:

Distant disease-free survival (LEFT) and breast cancer-specific survival (RIGHT) according to the 70-gene signature for 42 patients with highly endocrine-responsive tumors according to the St.Gallen criteria. Out of 11 low risk patients, 7 were untreated, 4 received chemotherapy and one of those received trastuzumab.

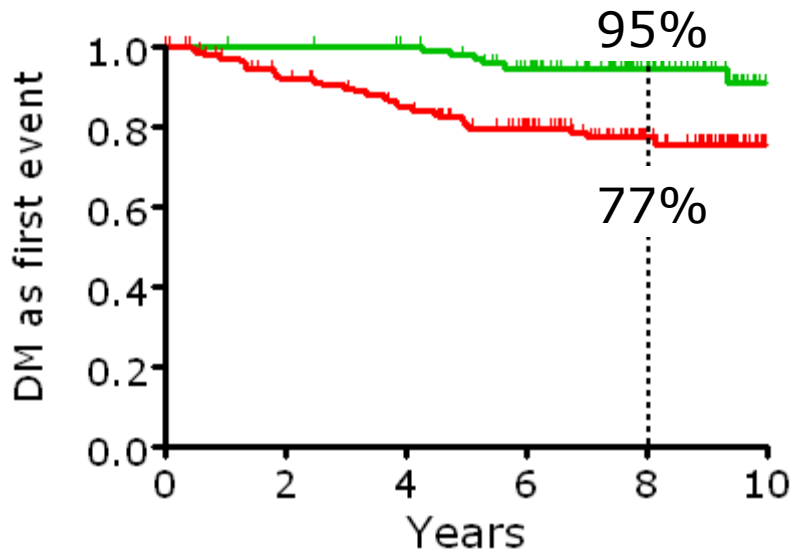
And Node +ve Breast Cancer?

- Always indication for chemotherapy?

70-gene Profile and Prognosis in Breast Cancer with 1-3 Axillary Lymph Node Metasases
S. Mook et al., *Breast Cancer Res Treatm* 2008.

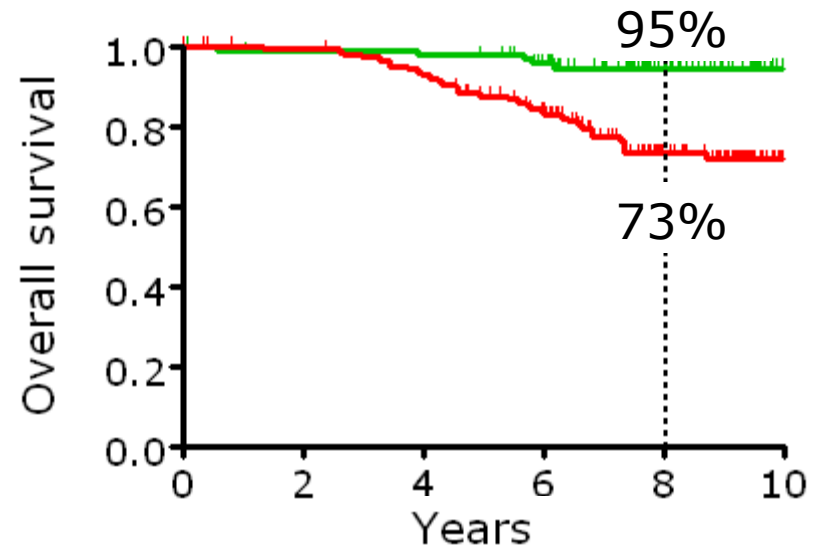
— Good profile (n=99)
— Poor profile (n=142)

Distant metastases as first event



HR 4.1
(95%CI 1.7 – 10.0), p=0.002

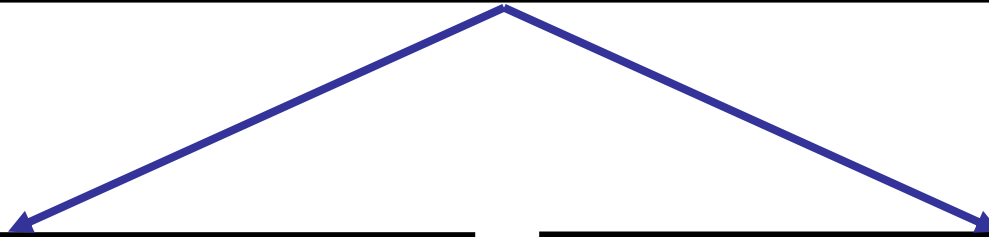
Overall survival



HR 5.4
(95%CI 2.1 – 13.8), p<0.001

Good profile: sufficiently low risk?

1-3 positive nodes, all patients (n=241)	8-yr OS	SE
Good profile (n=99)	95%	2
Poor profile (n=142)	73%	4



Chemotherapy (n=128)	8-yr OS	SE
Good profile (n=39)	95%	4
Poor profile (n=89)	69%	6

No chemotherapy (n=101)	8-yr OS	SE
Good profile (n=57)	94%	3
Poor profile (n=44)	77%	7

Adjuvant Chemotherapy and 70 gene prognosis signature Clinical Utility and Clinical Benefit

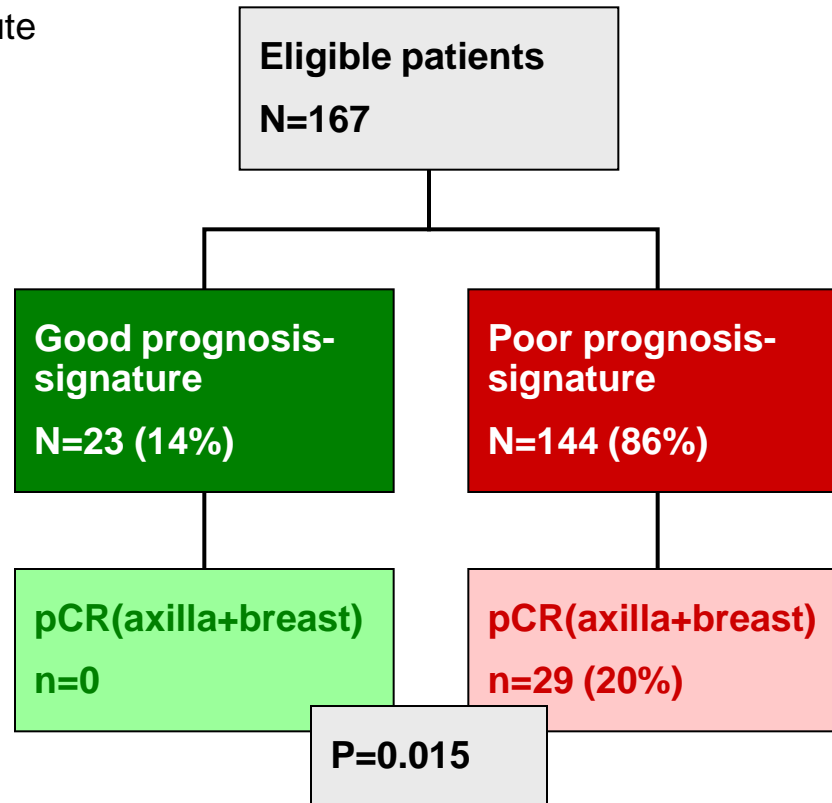
- 70 gene Low Risk Signature group has low risk for recurrence, and does not show significant chemo benefit
- 70 gene High Risk signature patients show significant neo-adjuvant chemo-sensitivity
- 70 gene High Risk Signature Patients show substantial Clinical Benefit of Adjuvant Chemotherapy (Cave: not a randomized trial)

Issues in early breast cancer

- MammaPrint and chemo-effect

Benefit of neo-adjuvant chemotherapy for MammaPrint high risk patients

- Netherlands Cancer Institute
- 2 clinical trials
- T-stage >3 cm and/or LNplus (SNB/FNA)
- ultrasound guided 14 gauge biopsies
- MRI imaging
- Pathology



- Antracycline-like
- Antracyclin-Taxane
- Taxane

- pCR:
- pathological
- complete remission

MammaPrint low risk signature → no benefit of chemotherapy

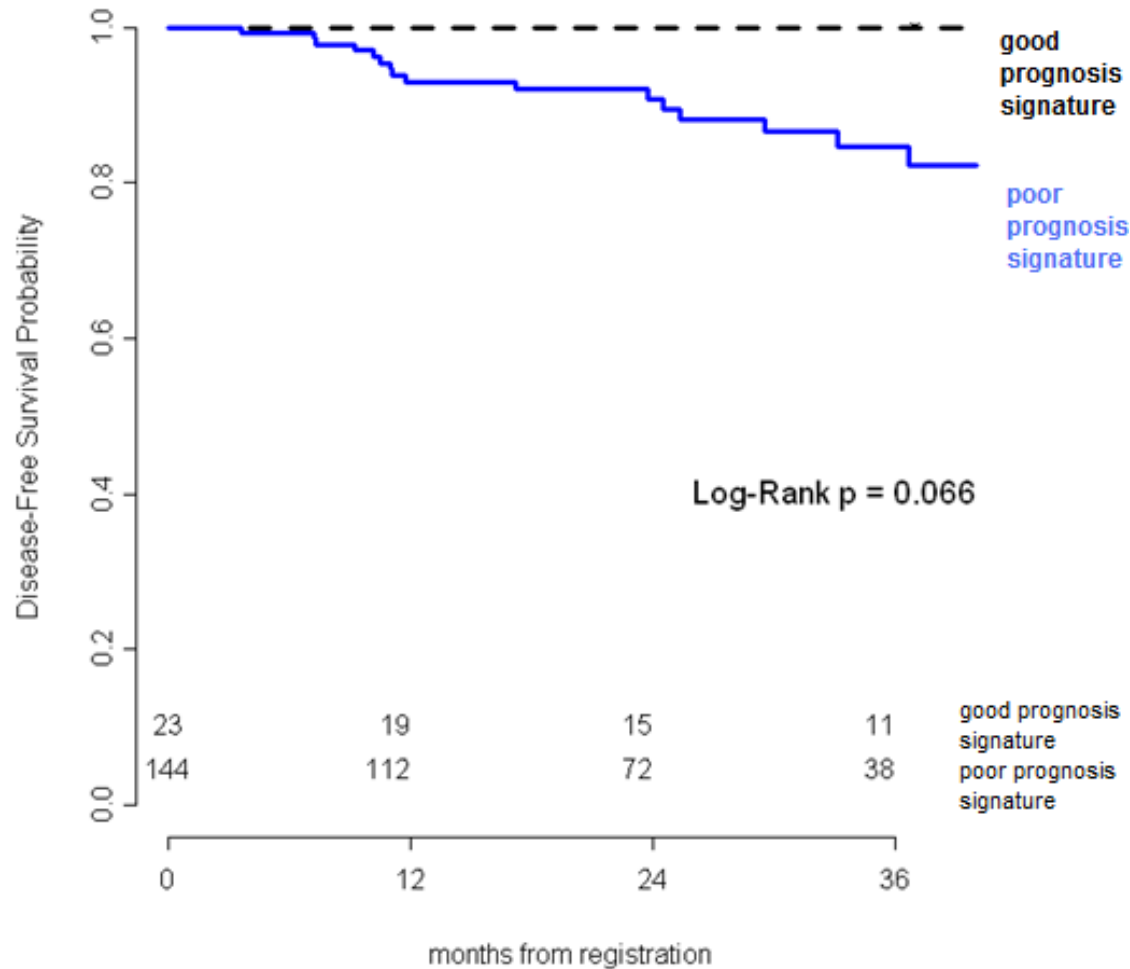
MammaPrint high risk signature → benefit of chemotherapy

Neo-adjuvant Standard Chemotherapy and MammaPrint Clinical Benefit

- 70 gene MammaPrint High Risk Signature patients show significantly higher chemosensitivity
- All pCR are found in the High Risk Signature group

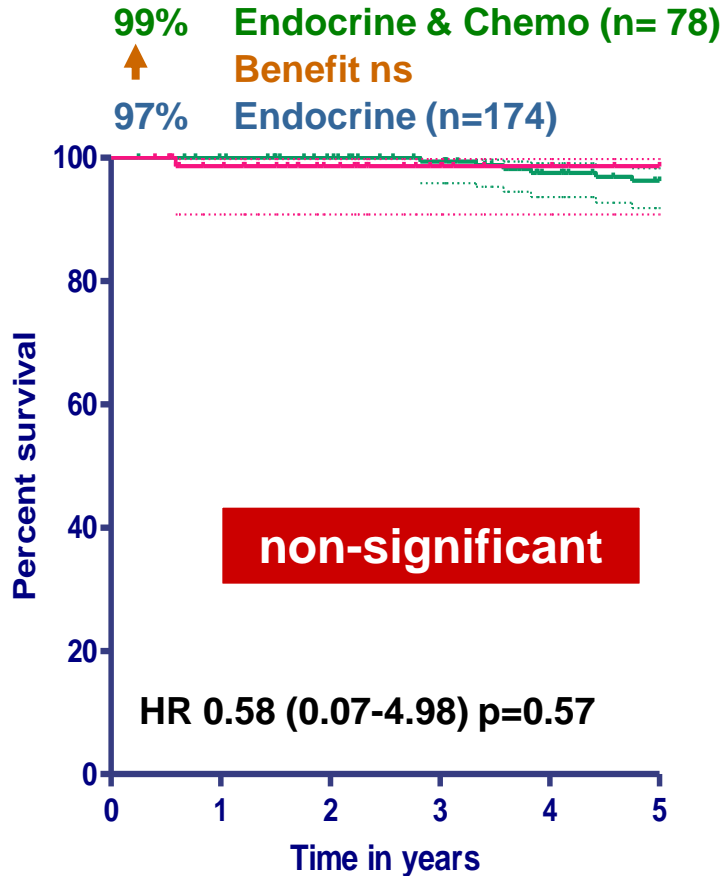
***High Risk Signature Patients show
Clinical Benefit of Chemotherapy***

So far, no recurrences in MammaPrint good prognosis group

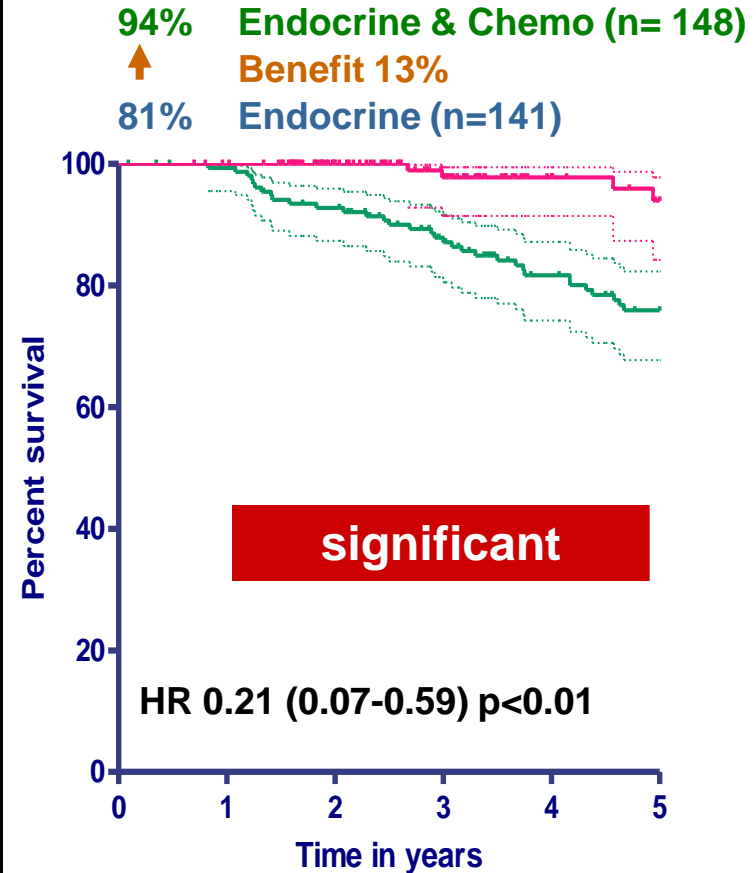


Breast Cancer Specific Survival (5 yrs) Endocrine vs Endocrine-Chemo within 70 gene low and high risk signature (n=575)

70 gene signature Low risk (n=268)



70 gene signature High risk (n=307)



Interaction term for differential effect p=0.45; Cave: not a randomized trial

Cox multivariate analysis: Backward Stepwise DDFS at 10 years for ET vs. ET + CT

Variable	p-value	HR	95% CI
Age	0.307	1.02	0.99-1.05
Tumor-Diameter	0.011	1.03	1.01-1.06
LN-status	0.136	1.17	0.95-1.44
Grade	0.024	1.78	1.08-2.95
ER-status	0.670	0.86	0.43-1.72
PR-status	0.012	0.48	0.27-0.85
Her2-status	0.398	1.33	0.69-2.55
ET vs. ET + CT	0.011	0.26	0.09-0.74

Adjuvant Standard Chemotherapy and MammaPrint Clinical Benefit

- MammaPrint High Risk signature patients show significant chemo-sensitivity
- MammaPrint Low Risk Signature group does not show significant chemo benefit

High Risk Signature Patients show substantial Clinical Benefit of Adjuvant Chemotherapy

(Cave: not a randomized trial)

Moving forward

Genomics in Breast Cancer

- Science: Prospective evaluation 70 gene MammaPrint signature and therapy benefit, 6000 patients (TRANSBIG-MINDACT)
- Science: Provide comprehensive biobank and standard molecular biological assays to integrate knowledge on tumor type, germline status (TRANSBIG-MINDACT)

MINDACT trial

- Is not to validate the prognostic value of the 70-gene MammaPrint
- Will tell us if chemotherapy is rightfully withheld to patients with a MP good prognosis who would have been advised chemotherapy on the basis of current clinical criteria within strict limits: 5 yrs breast cancer specific survival of 93-95%.

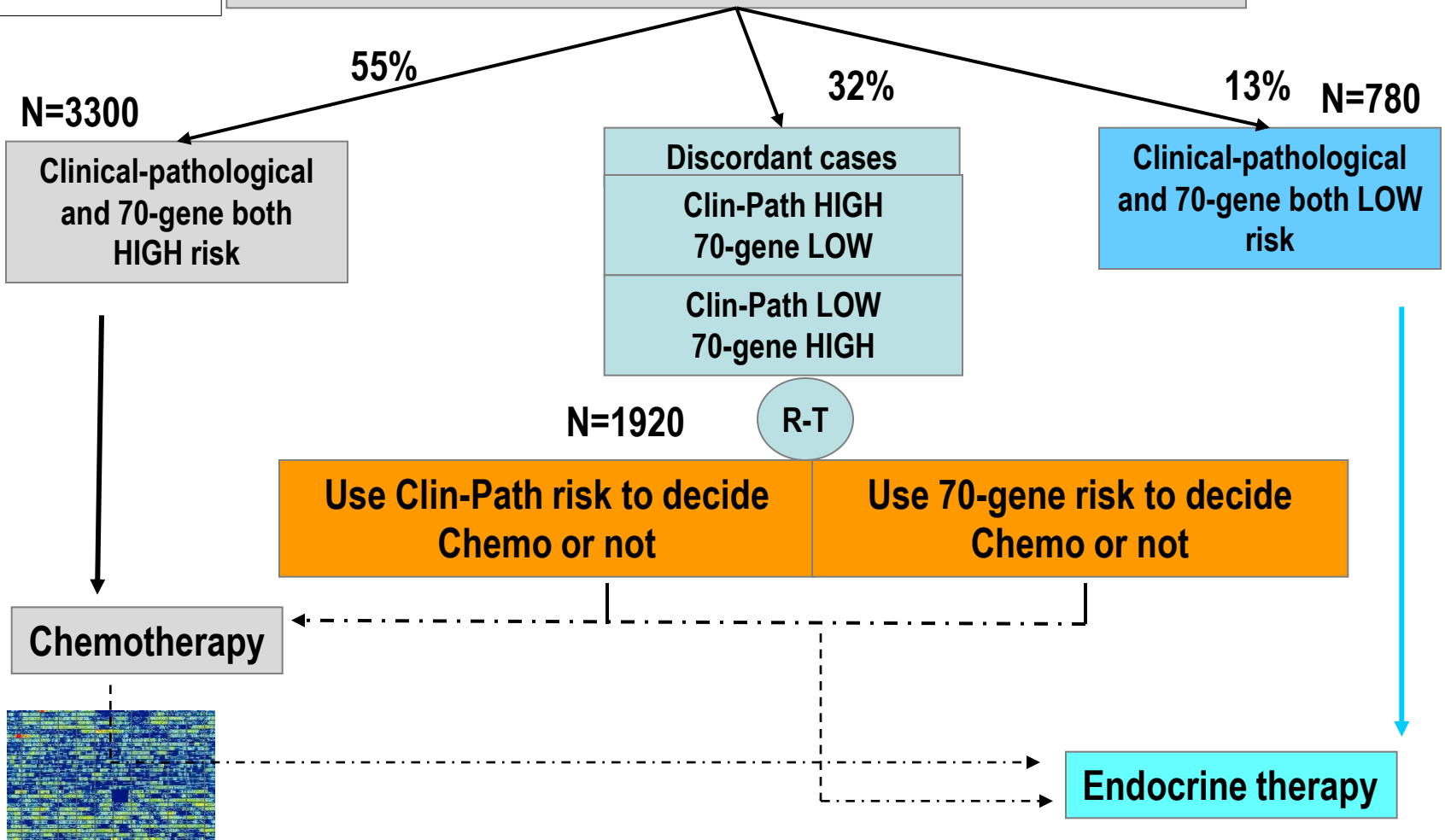
EORTC 10041 BIG 3-04 trial: MINDACT

6,000 Node neg & 1-3 N+ women



PI's: Martine Piccart, Emiel Rutgers, Fatima Cardoso

Evaluate Clinical-Pathological risk and 70-gene signature risk

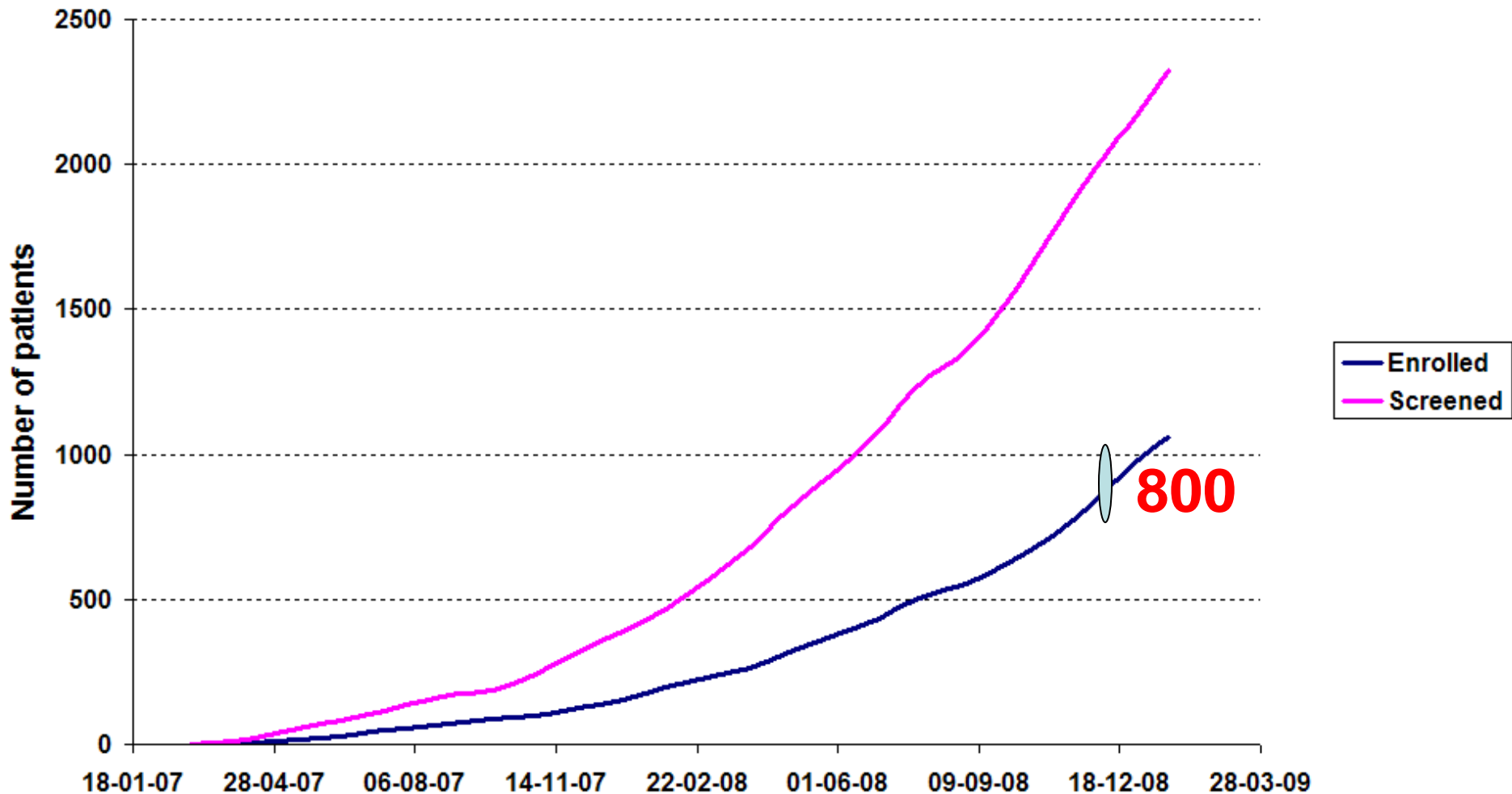


potential chemotherapy sparing in 10-15% pts, without affecting survival

PRE-SPECIFIED PILOT PHASE FIRST 800 PTS TO ENSURE:

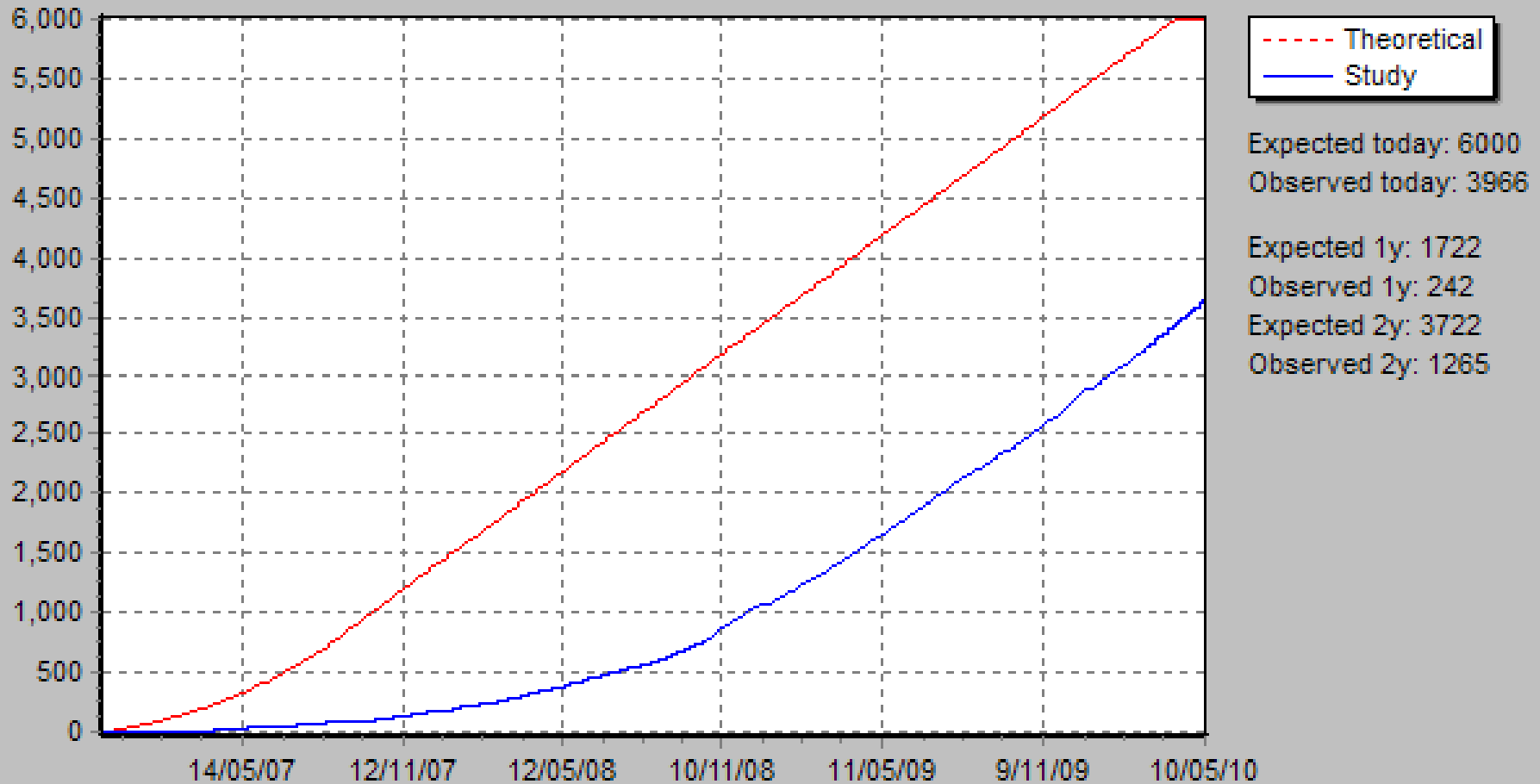
- **Logistically feasible**
- **Unbiased patient recruitment**
- **Less CT in genomic-based risk group than in clinical-based risk group**
- **Compliance**

MINDACT RECRUITMENT IN THE PILOT PHASE



MINDACT ENROLLED BY July 26 2010 after registration of 7183 pts.

Accrual of study 10041



Unbiased recruitment?

The patient population in the trial as compared to the patients from the validation series (all node -)

Older

Fewer ER-, fewer HER2+

Tumor size smaller

Tumor grade comparable

- **Expected fraction of clinical high risk patients 77% vs. observed 42.4%.**
- **Change in biology of breast cancer and/or more screening detected?**

RISK CROSS-TABLE

Clinical risk by 70 gene risk, at enrollment (overall % ages)			
	Clinical risk at enrollment		
	Low risk (N=461) N (%)	High risk (N=339) N (%)	Total (N=800) N (%)
70 gene risk at enrollment			
Low risk	386 (48.3)	141 (17.6)	527 (65.9)
High risk	75 (9.4)	198 (24.8)	273 (34.1)

Discordant cases: 27%

Estimates:

$p_C = 339/800$ (42%)

$p_G = 273/800$ (34%)

$p_C - p_G = 8.25\%$, 95% CI = 4.69% - 11.81% ($p < 0.0001$)

WAS CHEMO ACTUALLY GIVEN? COMPLIANCE!

Chemotherapy administration (best current knowledge) by assignment to chemotherapy			
	Treatment decision outcome		
	chemo (N=309)	no chemo**** (N=491)	Total (N=800)
	N (%)	N (%)	N (%)
Chemo received			
No	21* (6.8)	472 (96.1)	493 (61.6)
Yes	268 (86.7)	19** (3.9)	287 (35.9)
Unknown	20*** (6.5)	0 (0.0)	20 (2.5)

92% assigned to CT, received

100% assigned to no CT, did not receive

Overall no significant difference in toxicity
between different chemo regimens

CONCLUSIONS

- 1 Logistics **feasible** (>4500 pts enrolled!)
- 2 More low risk cancers: **Change in biology** of breast cancer and/or due to screening. This does **not affect discordancy** rates required for primary aim randomization
- 3 Clinicians & patients **comply** with the protocol in the “70-gene signature /genomic arm”.
- 4 Statistically significant difference in **reduction in Chemotherapy administration.**

The 70-Gene Signature:

1. Has been validated in terms of prognostication and adds to conventional criteria
2. It has also been validated for T₁₋₃N₁ tumors and for elderly patients
3. The assay is stable and reliable (FDA)
4. There are strong indications that the good-prognosis profile is associated with decreased chemotherapy benefit

Using a good-prognosis 70-Gene Signature to withhold adjuvant chemotherapy in clinically low- to moderate risk patients is therefore justified

- < テーマ > 治療効果予測と予後予測～乳がんのターゲット治療の模索～
乳がんにおける予後及び治療効果の予測検査法について、欧・米・東アジアからの招聘者と共に検討する国際セミナーです。
- < 場所 > 東京国際フォーラム ホールB5

Thank you for your attention & discussion & inviting me
JCCNB, Dr Seigo Nakamura



Paik et al, JCO, 2006 OTDX Predictive Data based on NSABP B-20 Archival FFPE Blocks

Multigene Predictor for Chemotherapy Response in Breast Cancer

Table 1. Kaplan-Meier Estimates of the Proportion of Patients Free of Distant Recurrence at 10 Years for Tamoxifen-Treated Patients and Tamoxifen Plus Chemotherapy-Treated Patients

Group	No. of Patients	Tamoxifen			Tamoxifen Plus Chemotherapy		
		10-Year DRF (%)	95% CI	No. of Patients	10-Year DRF (%)	95% CI	No. of Patients
All patients	651	87.8	83.3% to 92.3%	227	92.2	89.4% to 94.9%	424
Low risk (RS < 18)	353	96.8	93.7% to 99.9%	135	95.6	92.7% to 98.6%	218
Intermediate risk (RS 18-30)	134	90.9	82.5% to 99.4%	45	89.1	82.4% to 95.9%	89
High risk (RS ≥ 31)	164	60.5	46.2% to 74.8%	47	88.1	82.0% to 94.2%	117

NOTE. Results are given for all patients and for the pre-specified Recurrence Score risk categories. Abbreviations: DRF, distant recurrence free; RS, recurrence score.

The "High Risk" Tam only arm contained only 47 patients whose OS and DDFS was the metric for comparison with Tam plus Chemorx benefit—a very small group.

TargetPrint, single gene read out ER/PR/HER2

ER	IHC pos	IHC neg	PR	IHC pos	IHC neg	HER2	IHC pos	IHC neg
MA pos	309	16	MA pos	194	48	MA pos	50	10
MA neg	9	62	MA neg	29	108	MA neg	8	342

N=~400	ER	PR	HER2
Accuracy	0.94	0.80	0.96
Sensitivity	0.97	0.87	0.86
Specificity	0.80	0.69	0.97

Conclusions:

- Very high concordance between microarray and IHC or FISH for ER and HER2
- Accurate read-out of ER, PR and HER2 on microarray

reference