

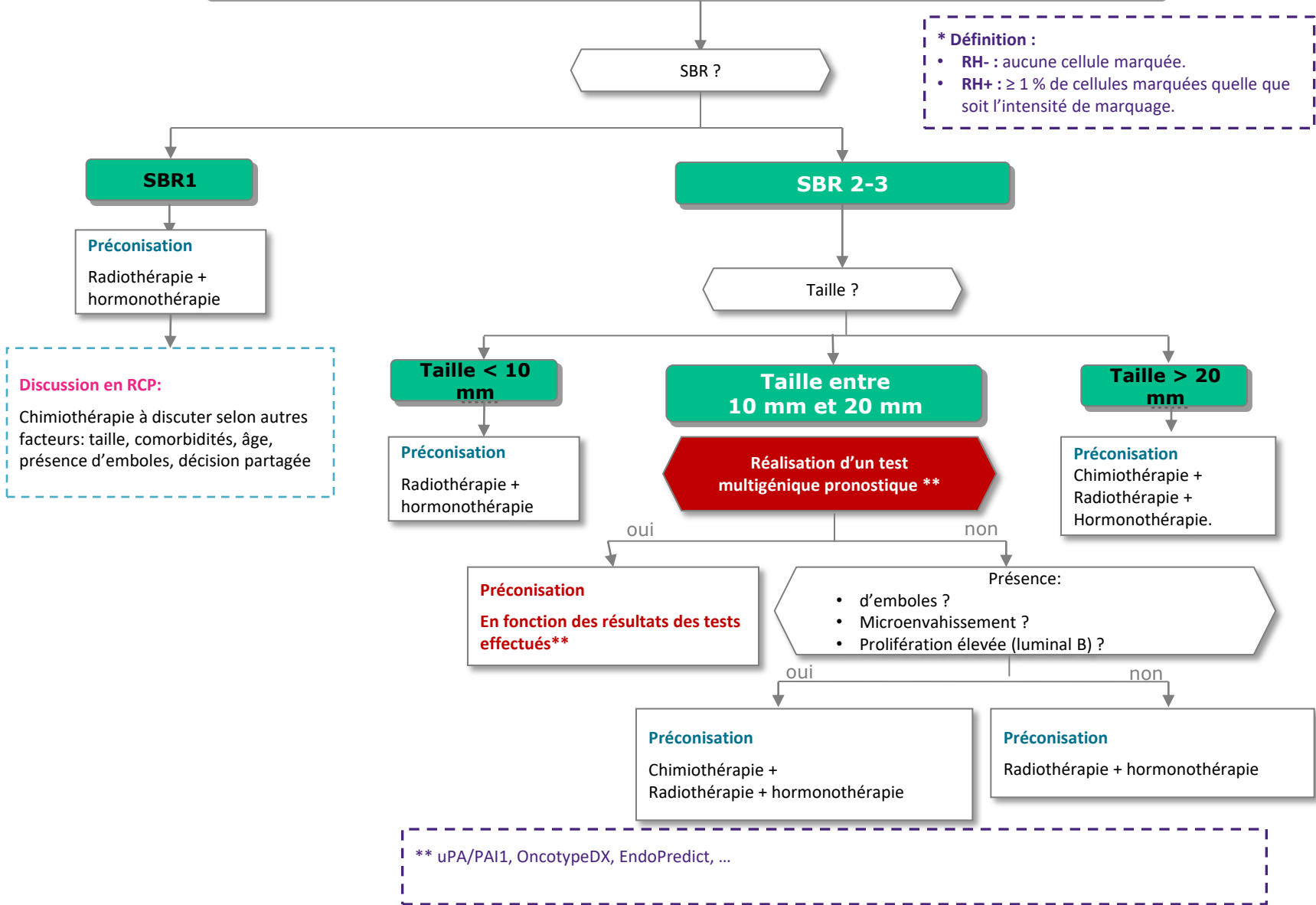


SOCIÉTÉ FRANÇAISE
DE SÉNOLOGIE
ET DE PATHOLOGIE
MAMMAIRE

Quand utiliser les signatures moléculaires pronostiques ?

Olivier Trédan
Centre Léon Bérard

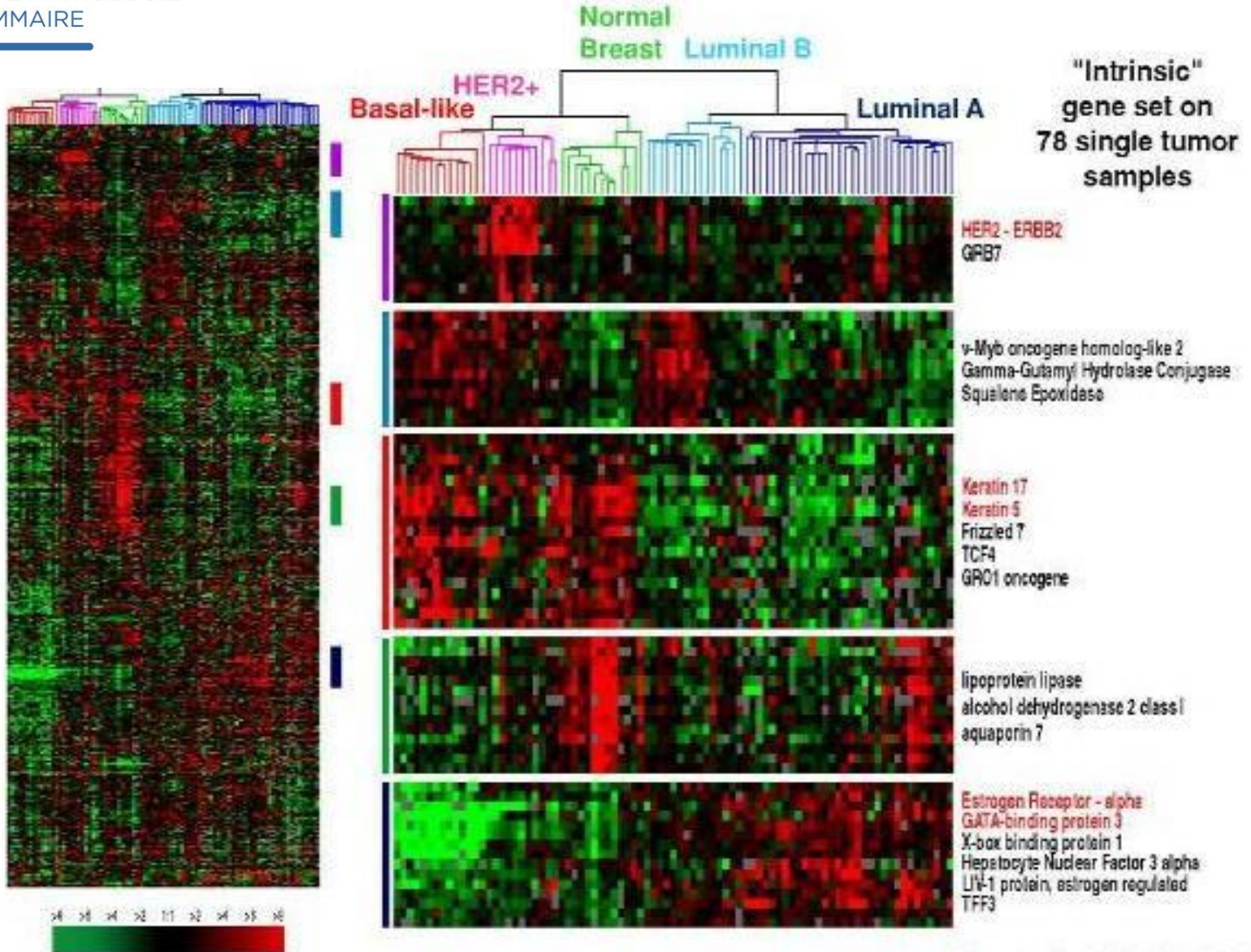
Cancer du sein non métastatique – traitement adjuvant des tumeurs pN0 ou micro envahissement ganglionnaire chez une patiente > 35 ans et < 70 ans et Her2- et RH+*

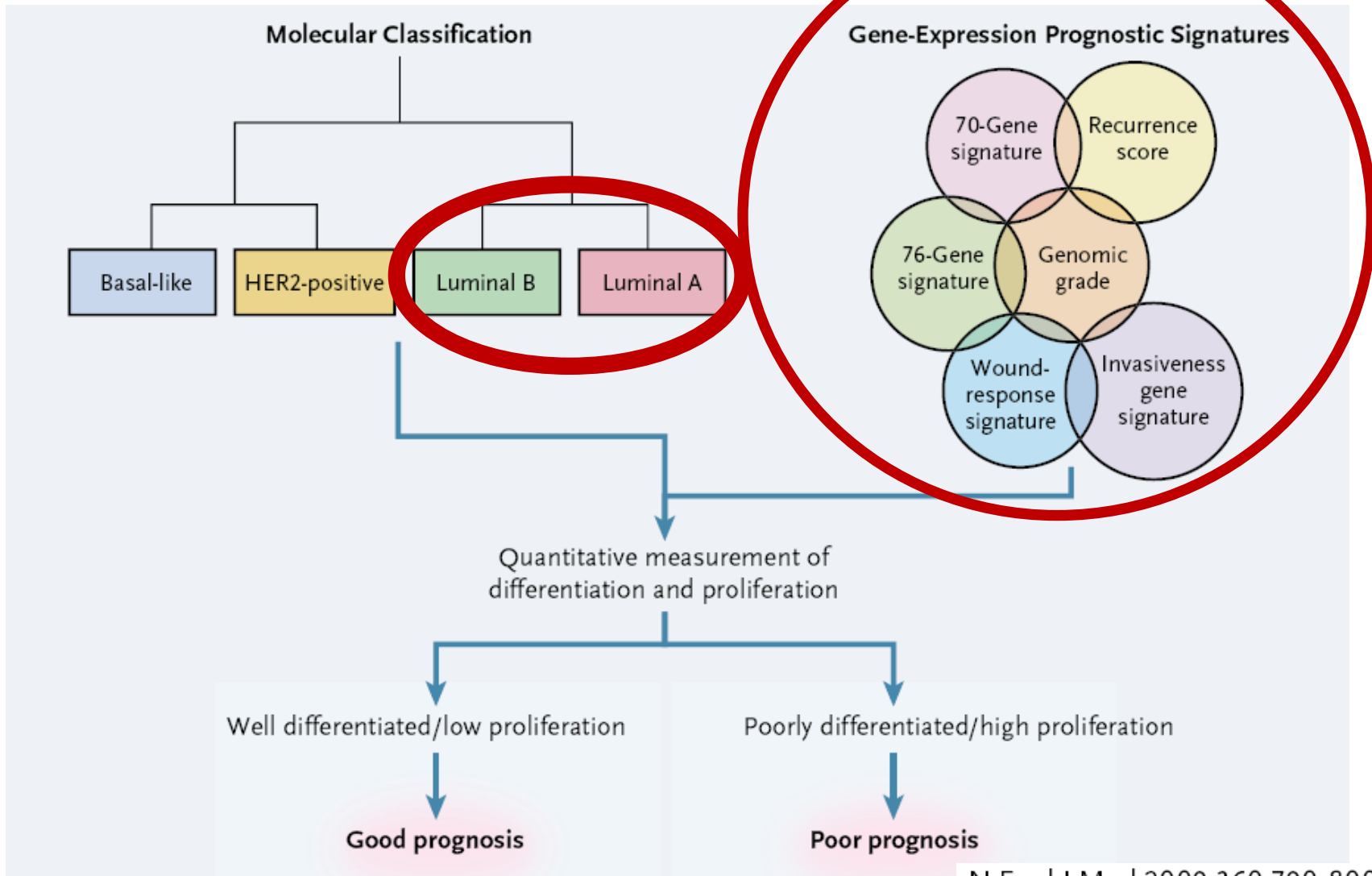


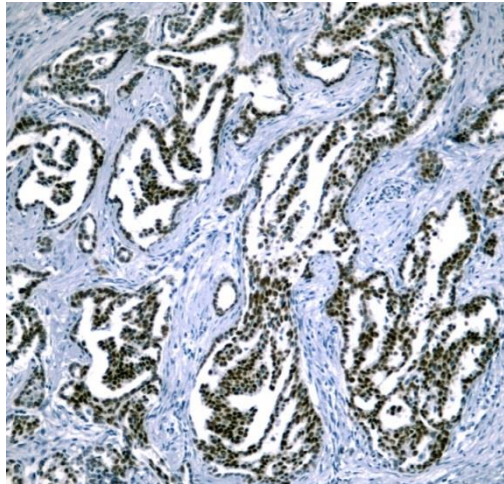
* Définition :

- RH- : aucune cellule marquée.
- RH+ : ≥ 1 % de cellules marquées quelle que soit l'intensité de marquage.

** uPA/PAI1, OncotypeDX, EndoPredict, ...



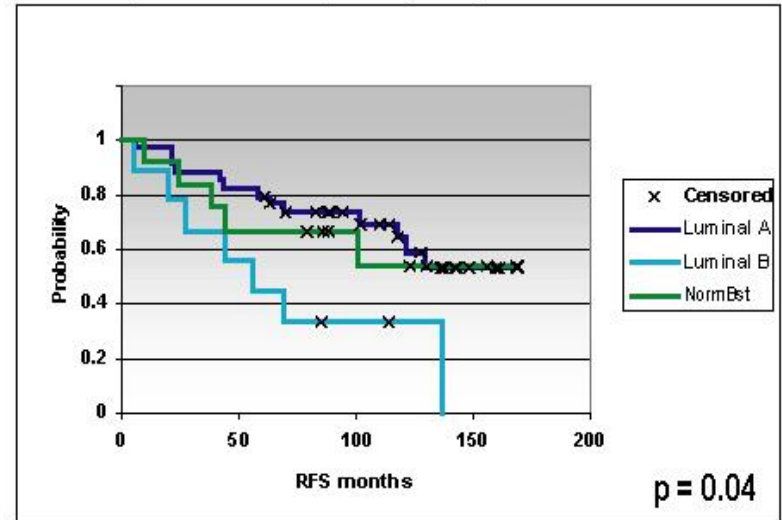




Luminal A

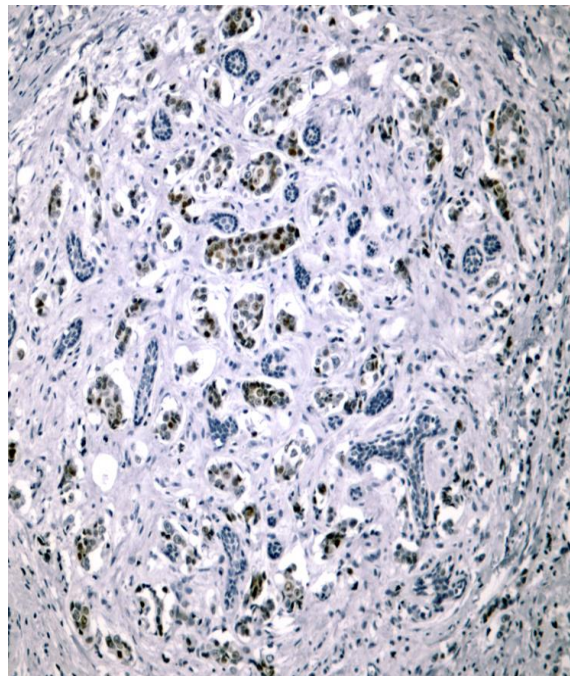
- RE+
- Et tous
 - RP +
 - HER2 -
 - Ki67 bas
 - bas risque molec

60 Sample ER+ Tamoxifen-Treated Test Set
Ma et al., Cancer Cell 5, 1-10 (2004).



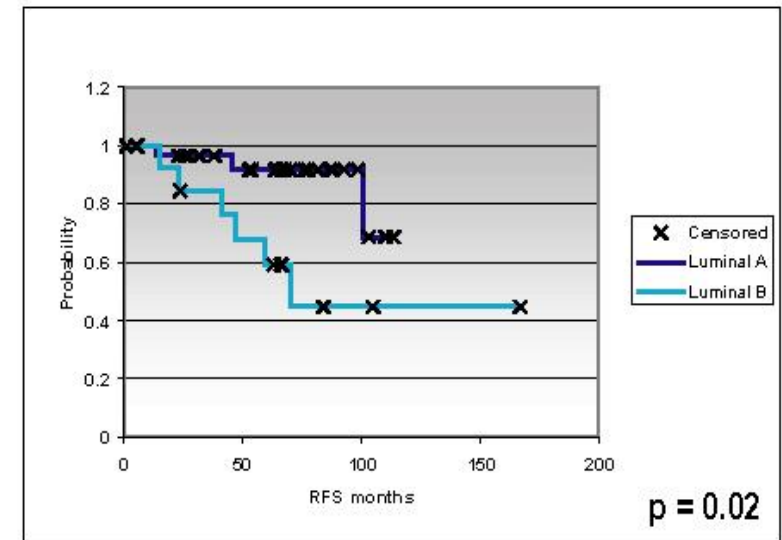
Luminal B

- RE+
- Et au moins
 - RP faible
 - Ki67 élevé
 - Haut risque molec



Luminal B HER2 +

- RE+, HER2 3+
- Quelque soit RP
- Quelque soit Ki67



45 Tamoxifen Treated Test Set #2

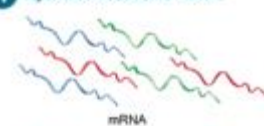
Chang et al., PNAS 102, 3738-43 (2005) + UNC



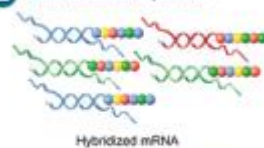
SOCIÉTÉ FRANÇAISE
DE SÉNOLOGIE
ET DE PATHOLOGIE
MAMMAIRE

Prosigna™ labels genes with a barcode

1 Hybridize CodeSet to mRNA.



2 Remove excess reporters.



3 Bind hybridized reporter to surface of cartridge.



4 Immobilize and align hybridized reporter.



5 Collect data.

Barcode	Gene	Counts
	LumA	3
	LumB	2
	Her2e	1
	Basal	1

Prosigna (PAM50) (NanoString Technology, USA)

RECONNAISSANCE DES
SOUS-TYPES « MOLECULAIRES »
(LumA, LumB, HER2-enrichi, Basal)

tissu fixé, inclus en paraffine
puce ADN avec le signal numérique
(1 gène = 1 code bar)

50 GENES SELECTIONNES

TEST « LOCAL »
(l'analyse se fait dans les laboratoires
avec équipements spécifique)

SIGNATURE
RECHUTE TARDIVE
(à 10 ans)

BAS RISQUE (ROR)
+ HORMONOTHERAPIE / - CHIMIOOTHERAPIE
RISQUE INTERMEDIAIRE (ROR)

HAUT RISQUE (ROR)
+ HORMONOTHERAPIE / + CHIMIOOTHERAPIE

5 min HANDS-ON
Day 1

1 Hybridize

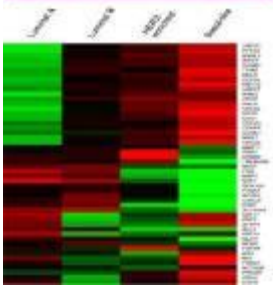
5 min HANDS-ON
Day 2
AUTOMATED

2 Purify

5 min HANDS-ON
Day 2
AUTOMATED

3 Count

Subtypes have distinct gene expression



$ROR = aR_{LumA+}$

bR_{LumB+}

cR_{Her2e+}

dR_{Basal+}

$eP+$

fT

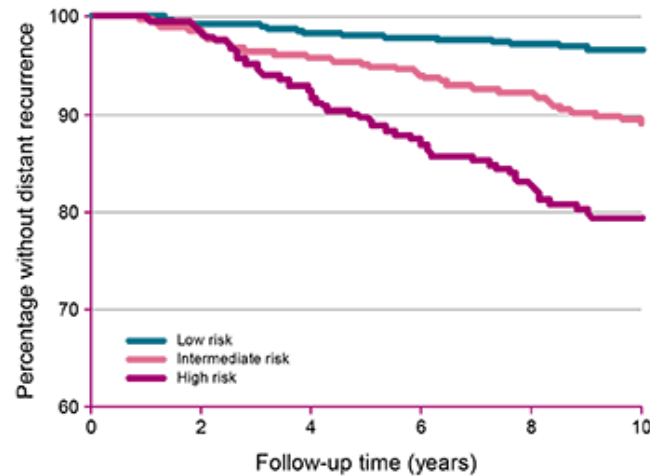
Pearson's correlation to centroids*

Proliferation score (19 genes)

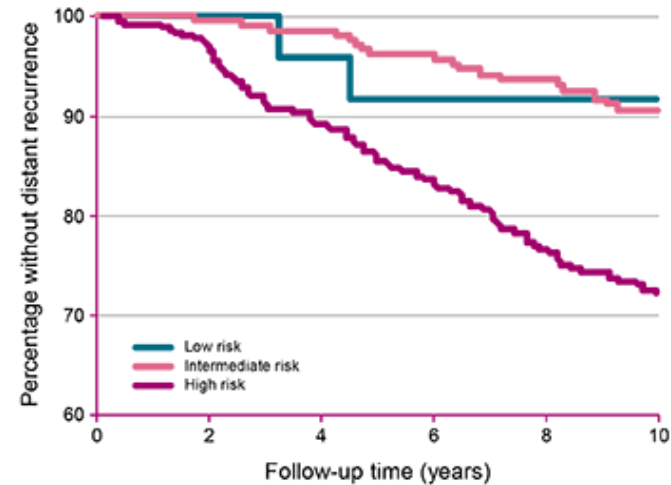
Tumor size



Prosigna (PAM50) (NanoString Technology, USA)



DRFS in node-negative patients¹



DRFS in patients with 1 to 3 positive nodes¹

RISQUE BAS : >95% DE SURVIE SANS RECHUTE (Dg + 10 ans)
RISQUE HAUT : <80% DE SURVIE SANS RECHUTE (Dg + 10 ans)

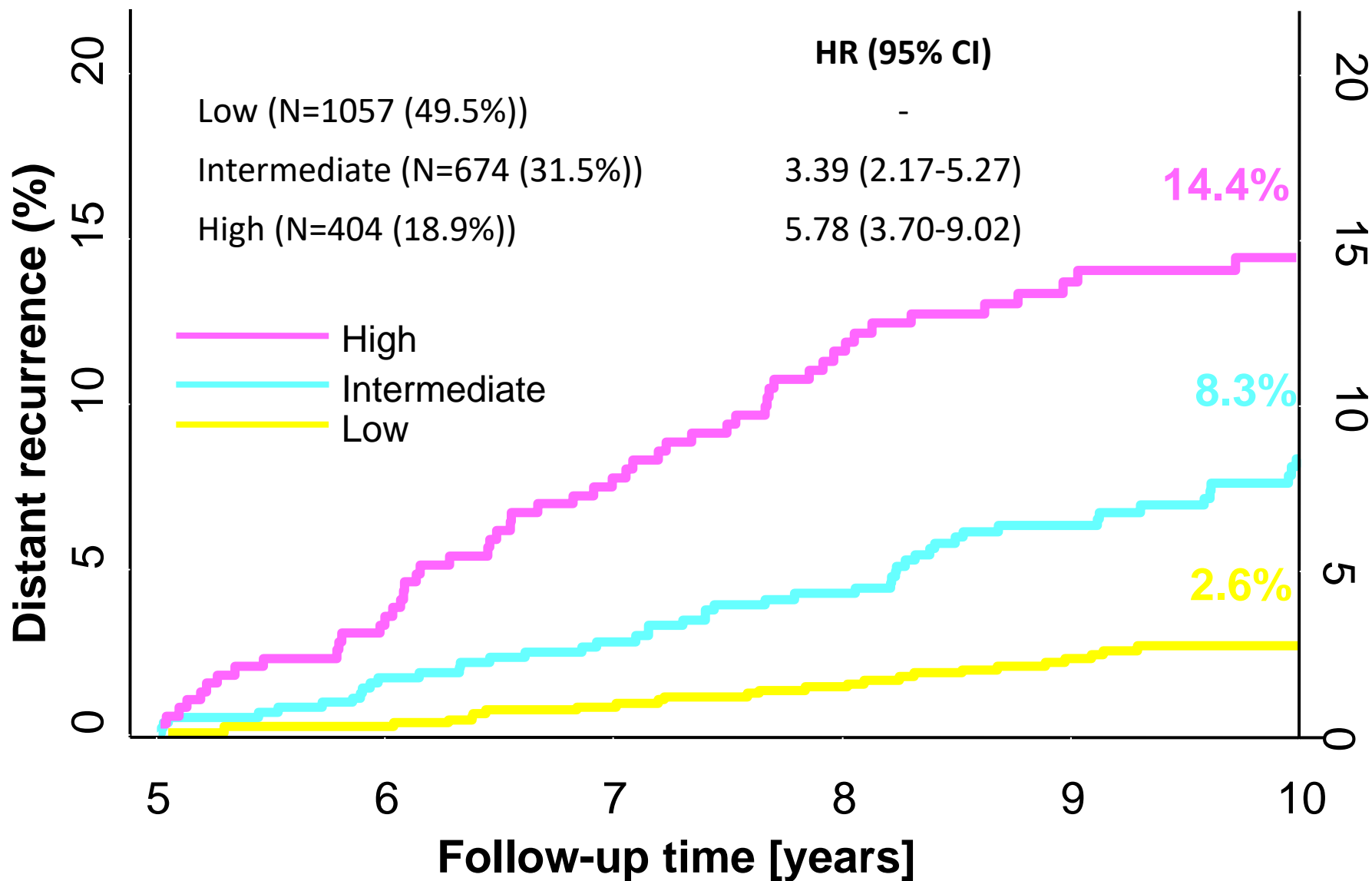
N0

TransATAC + ABCSG-8 : 2400 pts

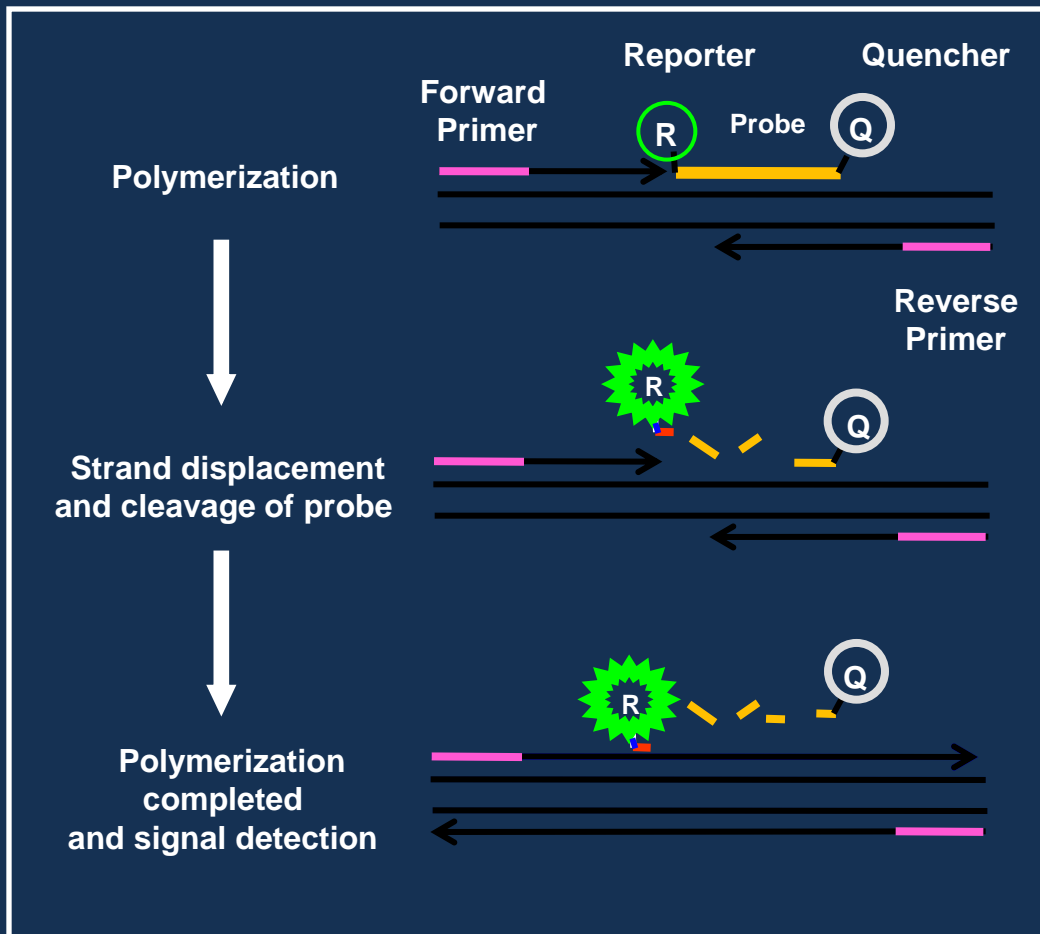
**RH+ / HER2- T1-2 N0/N+ postménop 5 ans d'anti-œstrogènes adj
tous les groupes de risque
tous les sous-types moléculaires**



Prosigna (PAM50) pour les rechutes tardives



Oncotype DX[®] Uses RT-PCR Technology and is Analytically Validated



- RT-PCR works well with RNA from formalin-fixed paraffin-embedded tissue
- RT-PCR provides >65,000-fold range of measurement
- RT-PCR reactions can be repeated with high quantitative precision
- Standardized RT-PCR optimized for use of small RNA fragments and various sources of pre-analytical variability

Oncotype DX[®] Recurrence Score

16 CANCER RELATED GENES

Estrogen	Proliferation	HER2	Invasion	Others
ER PR Bcl2 SCUBE2	Ki-67 STK15 Survivin Cyclin B1 MYBL2	GRB7 HER2	Stromelysin 3 Cathepsin L2	CD68
				GSTM1
				BAG1

5 REFERENCE GENES

Beta-actin	GAPDH	RPLPO	GUS	TFRC
------------	-------	-------	-----	------

Onco^{type} DX[®] Recurrence Score Calculation and Risk Categories

Recurrence Score =

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

Risk Group

Recurrence Score

Low risk

<18

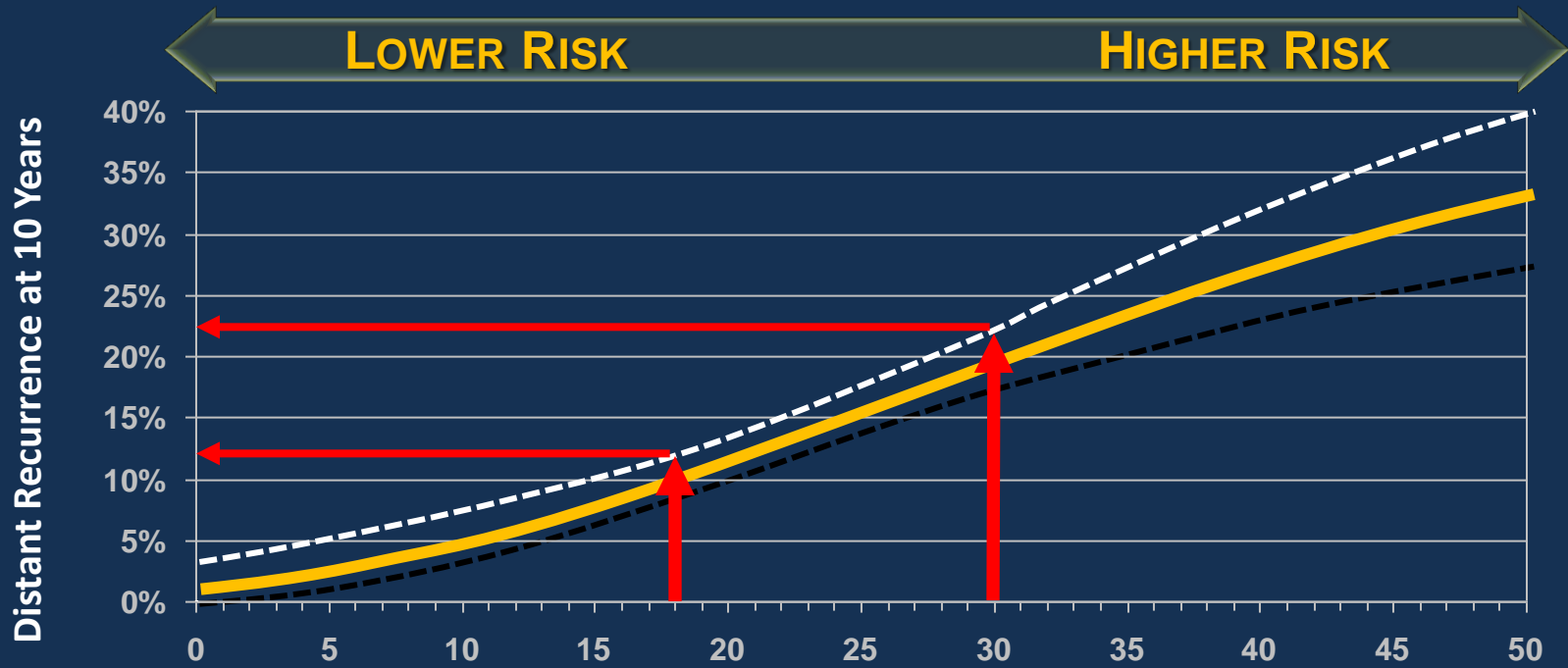
Intermediate risk

18 - 30

High risk

≥31

Onco^{type} DX[®] Recurrence Score Calculation and Risk Categories



Low risk

Recurrence Score <18

Intermediate risk

18 - 30

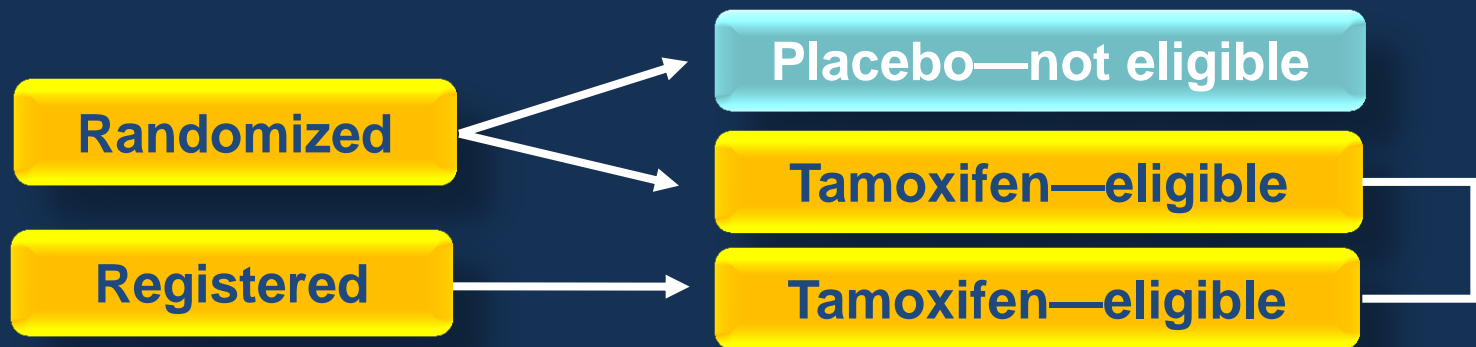


High risk

≥31

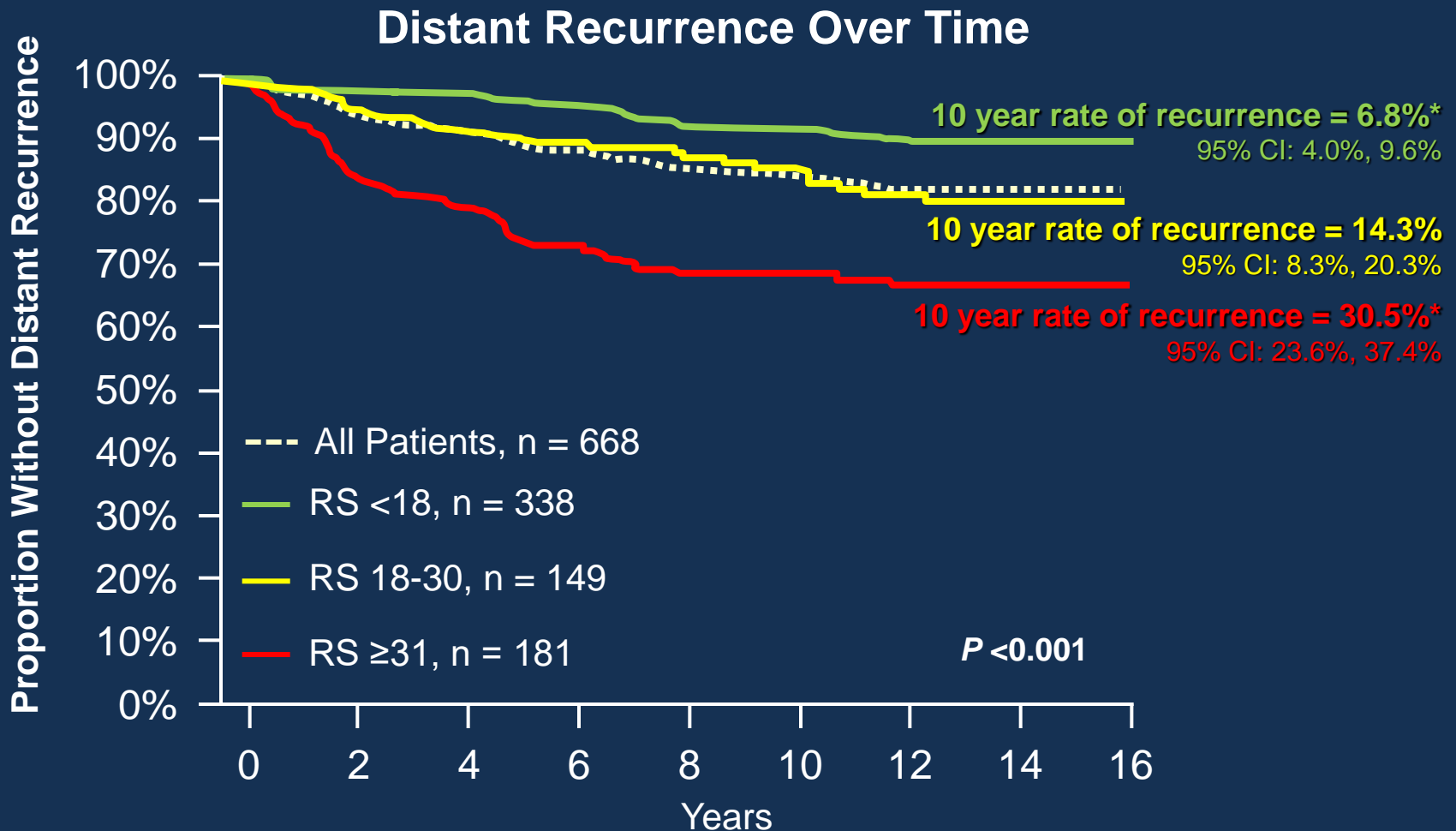
Onco^{type} DX[®] Clinical Validation: NSABP B-14

- Objective: Prospectively validate Recurrence Score as predictor of distant recurrence in N⁻, ER⁺ patients



--- All Patients, n = 668

Onco^{type} DX[®] Clinical Validation: NSABP B-14 – Distant Recurrence

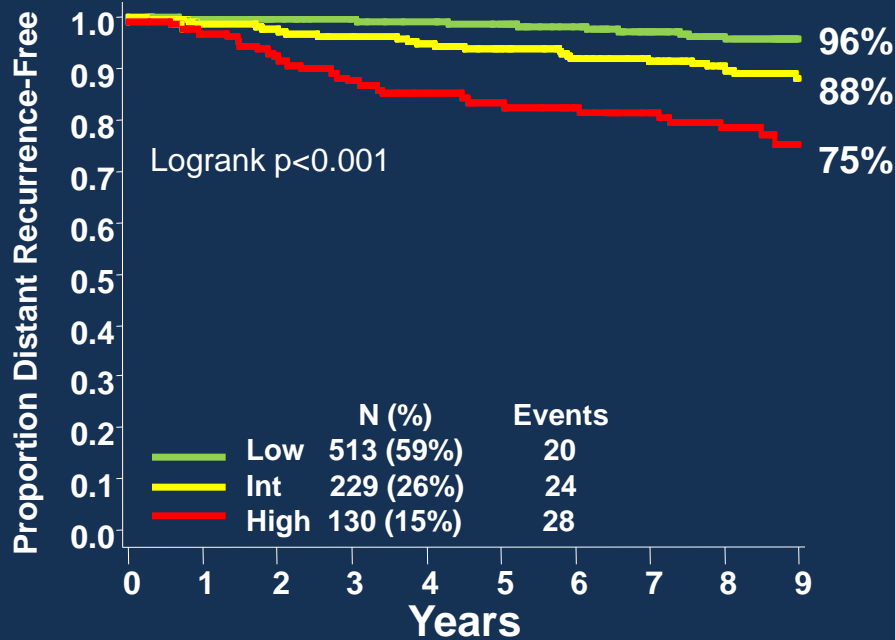


*10-year Distant Recurrence comparison between low-and high-risk groups: $P < 0.001$

Paik et al. *N Engl J Med.* 2004;351:2817-2826.

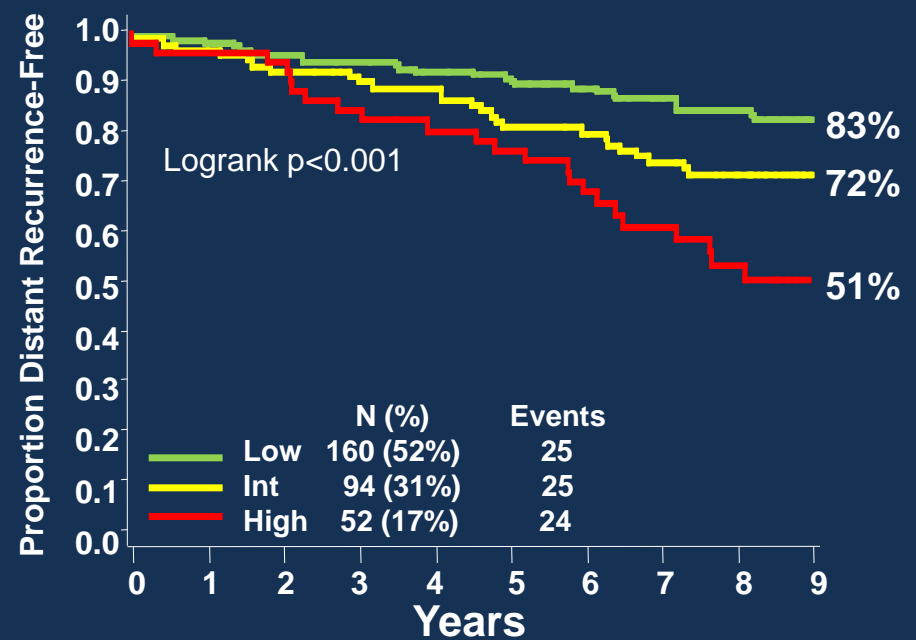
Oncotype DX[®] Clinical Validation: TransATAC

**Node Negative (n=872)
(both treatment arms)**



RS Group	HR* (95% CI)
High vs Low	5.2 (2.7 – 10.1)
Int vs Low	2.5 (1.3 – 4.5)

**Node Positive (n=306)
(both treatment arms)**

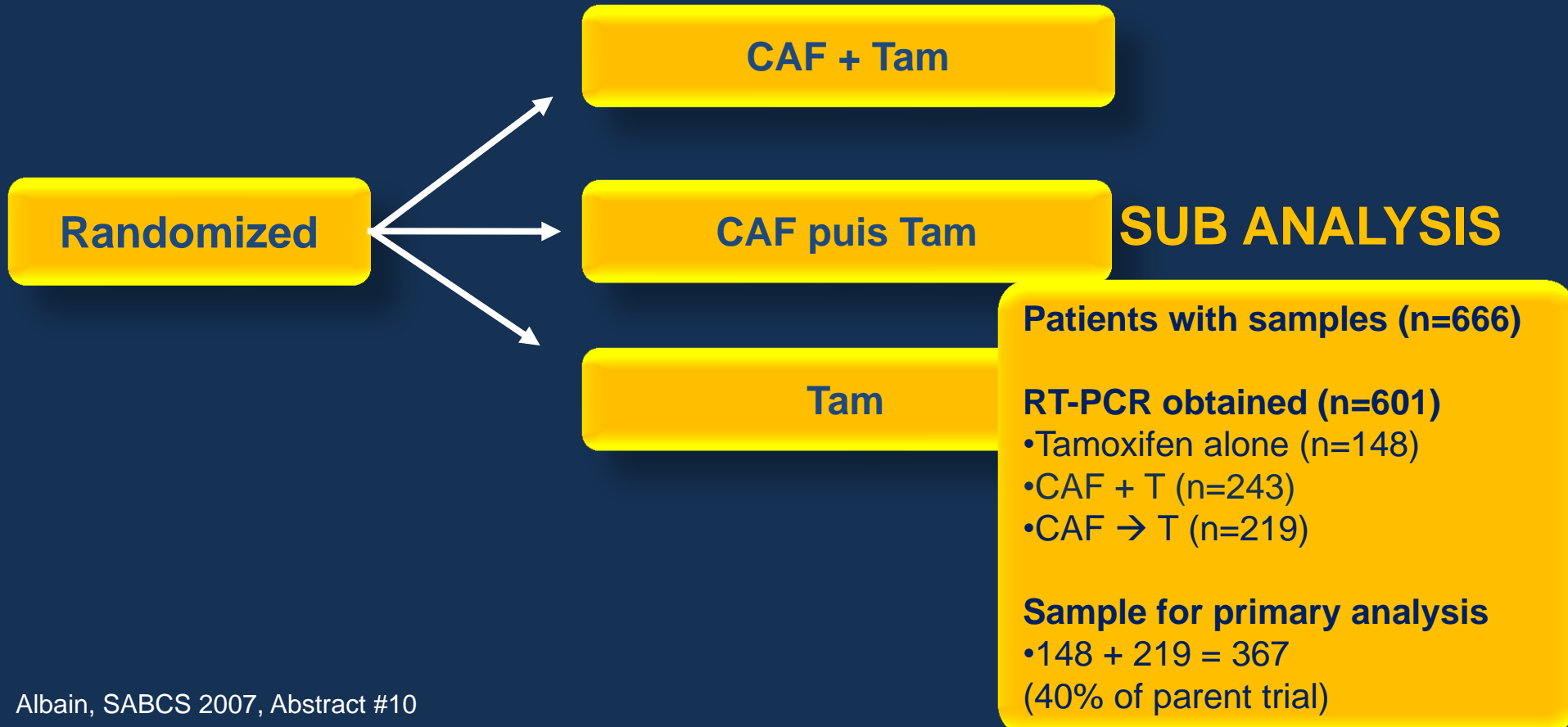


RS Group	HR* (95% CI)
High vs Low	2.7 (1.5 – 5.1)
Int vs Low	1.8 (1.0 – 3.2)

*Hazard Ratio for RS Group adjusted for tumor size, grade, age and treatment

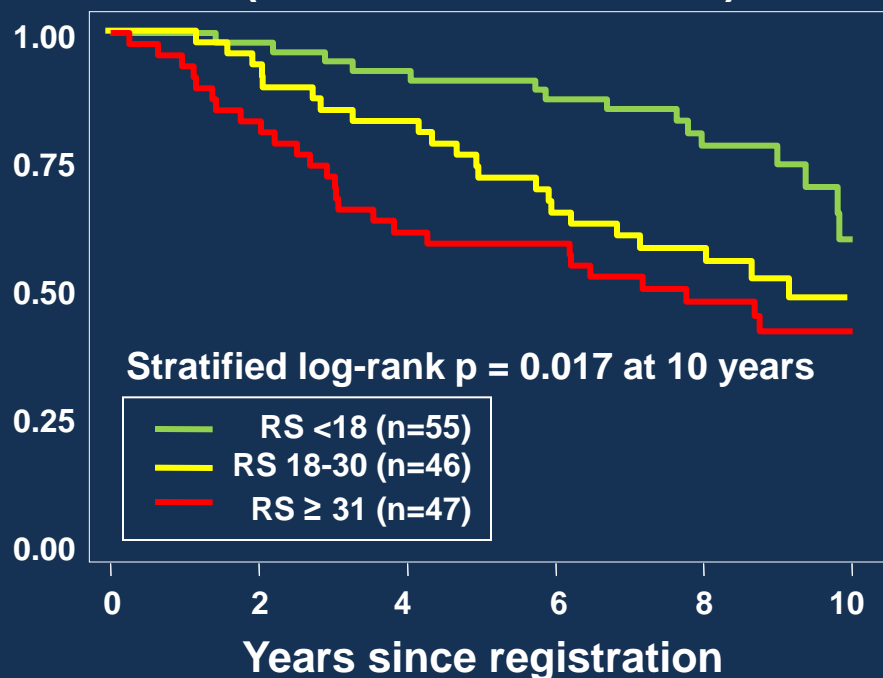
Oncotype DX[®] Clinical Validation: SWOG 8814 sub-analysis

- Objective: To determine the relationship between RS and chemotherapy benefit in N+, ER+ patients



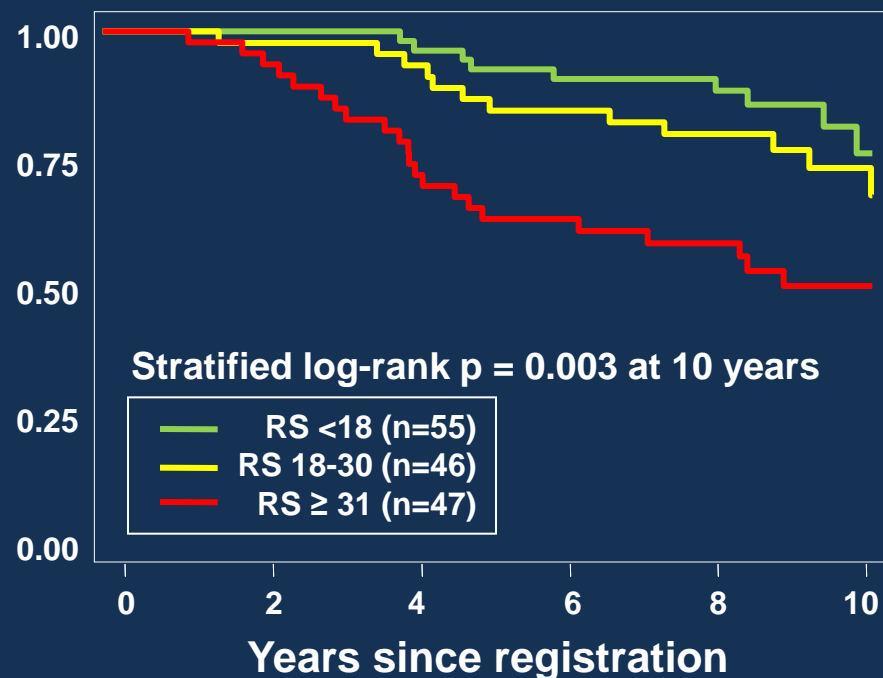
Oncotype DX[®] Clinical Validation: SWOG 8814 sub-analysis for N+ pts on Tam

**DFS by Risk Group
(tamoxifen alone arm)**



10-year DFS: 60%, 49%, 43%

**OS by Risk Group
(tamoxifen alone arm)**



10-year OS: 77%, 68%, 51%

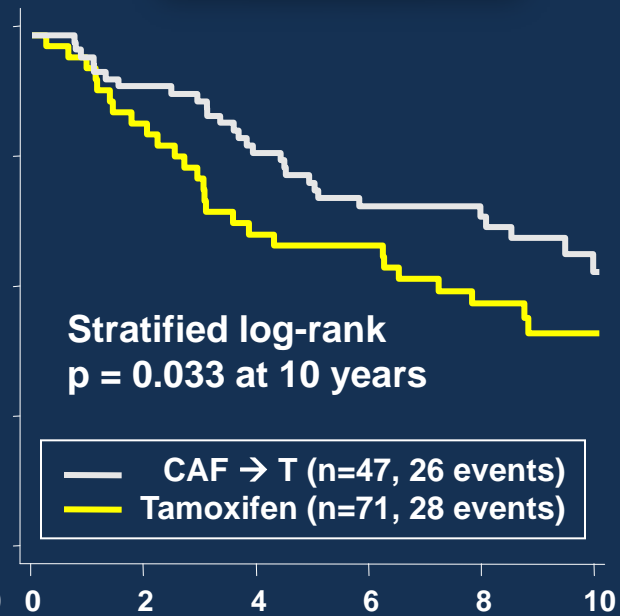
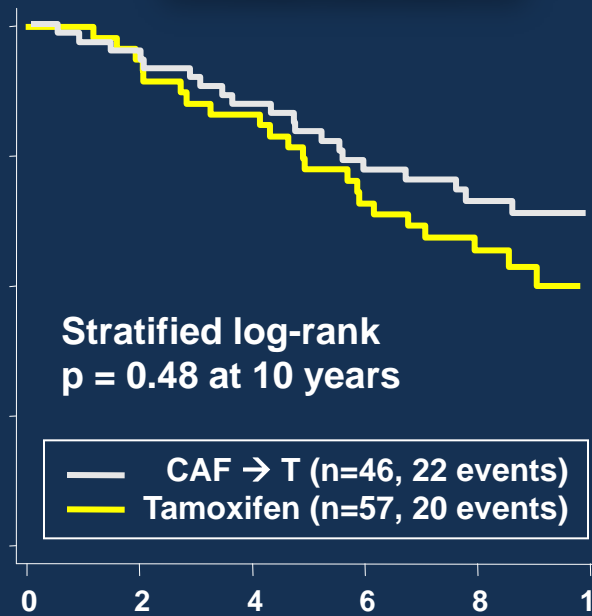
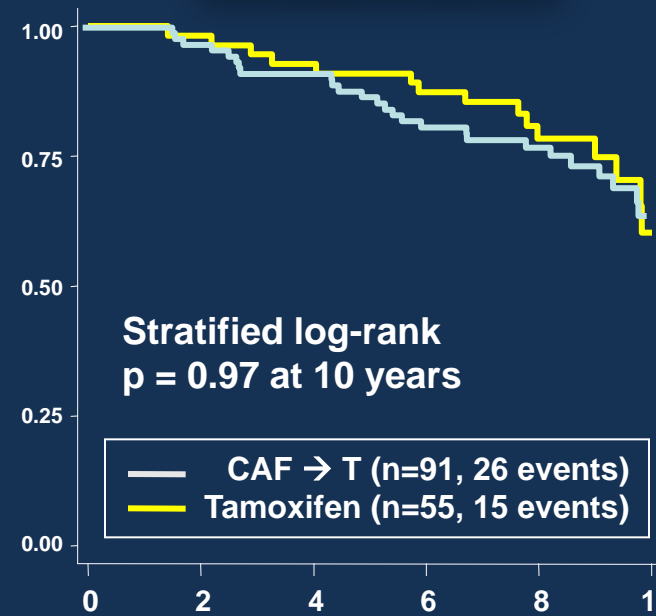
High Recurrence Score Predictive of Chemotherapy Benefit in Node-Positive Patients

DFS BY TREATMENT & RS GROUP

RS < 18

RS 18-30

RS ≥31



Years since registration

Years since registration

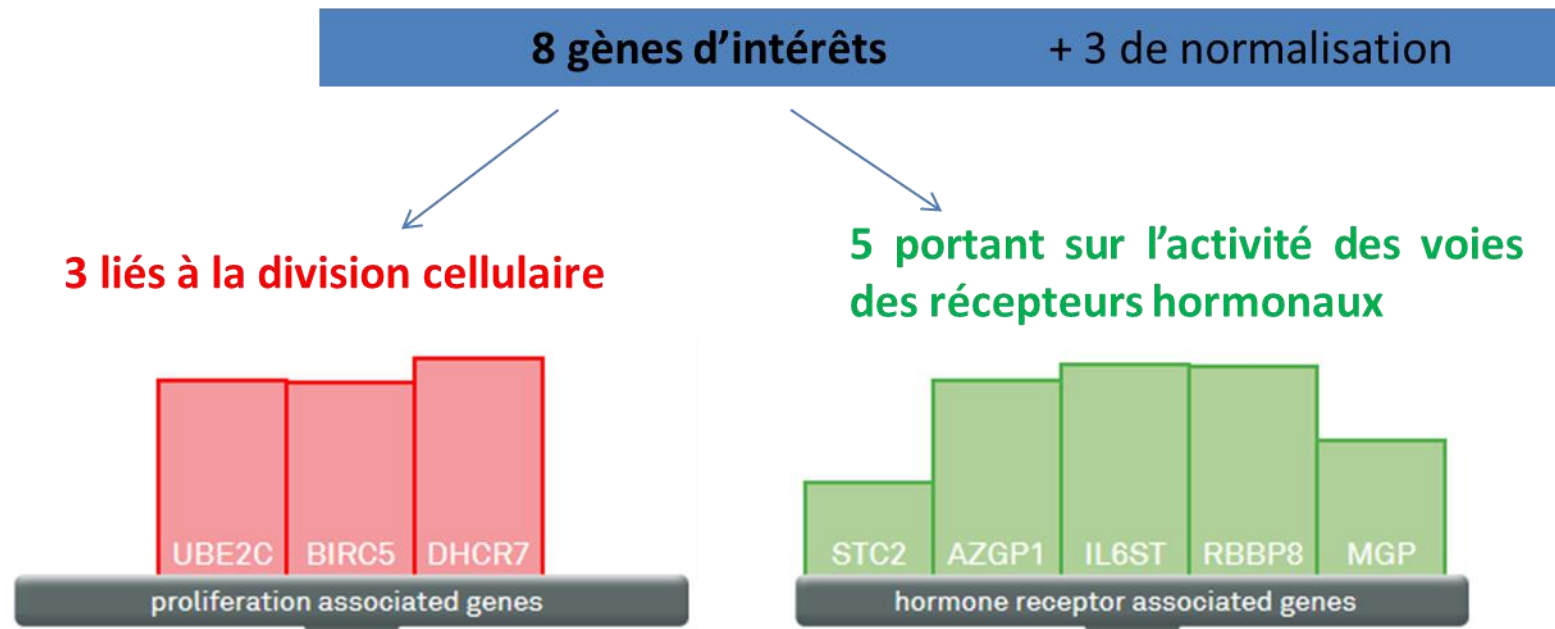
Years since registration

**No benefit to CAF
over time if low RS**

**Strong benefit if
high RS**

Test EndoPredict

- Il s'agit d'un test d'expression génique par RT-PCR à partir d'un échantillon tumoral fixé dans de la paraffine
- 12 gènes spécifiques sont testés





Test EndoPredict

- Classification du **risque selon le EP score**
 - Bas
 - Elevé
- Classification du **risque selon le EPclin score**
(EP score + T + N)
 - Bas
 - Elevé
- Détermination de la **probabilité d'une rechute à distance à 10 ans** selon le EPclin score
 - 0-100 %



EndoPredict® Report

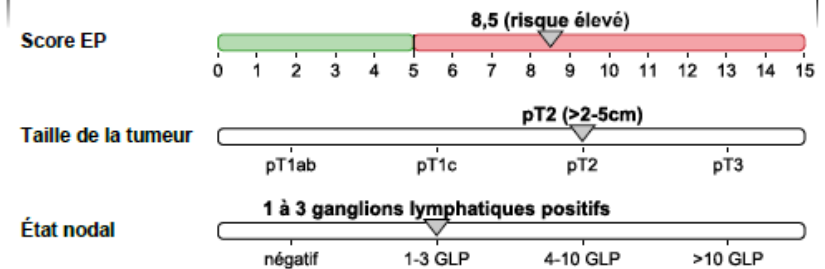


ID d'échantillon : 15F1045/250035022

Note :

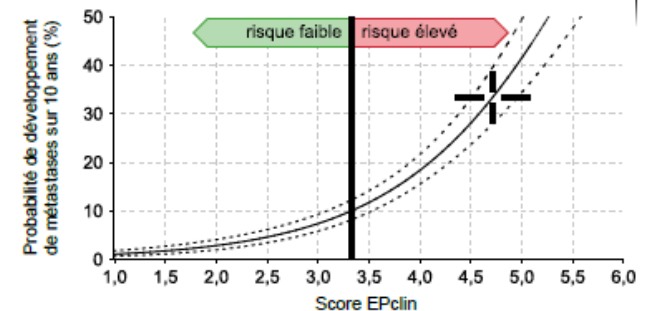
2 oct. 2015 08:30 (CEST)

Caractéristiques de la tumeur



Évaluation du risque selon EPclin

EPclin combine le score EP, la taille de la tumeur et le nombre de ganglions lymphatiques positifs pour définir un score au pouvoir prédictif supérieur.



Score EPclin

4,7

Risque 10a EPclin*

33%

Classe EPclin

risque élevé

* La probabilité de développement d'une métastase à distance sur 10 ans chez les patientes ayant suivi une thérapie endocrinienne de 5 ans est de 33%.

Rapport type EndoPredict®



EndoPredict® Report

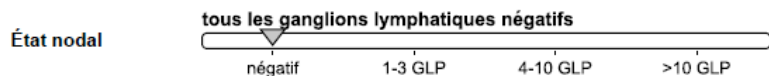
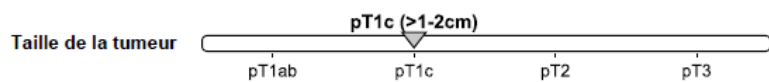


ID d'échantillon :

Note :

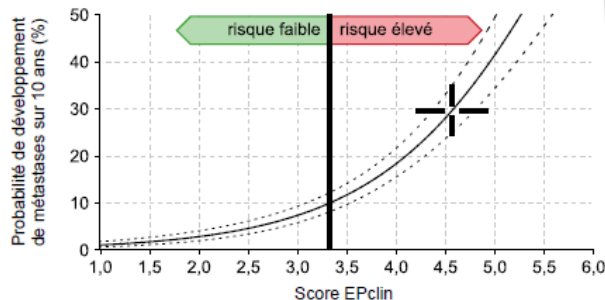
9 juin 2016 14:32 (CEST)

Caractéristiques de la tumeur



Évaluation du risque selon EPclin

EPclin combine le score EP, la taille de la tumeur et le nombre de ganglions lymphatiques positifs pour définir un score au pouvoir prédictif supérieur.



Score EPclin

4,6

Risque 10a EPclin*

30%

Classe EPclin

risque élevé

* La probabilité de développement d'une métastase à distance sur 10 ans chez les patientes ayant suivi une thérapie endocrinienne de 5 ans est de 30%.

Validation par le pathologiste

Les contrôles indiqués ont été effectués et le résultat du test est valide.

Signature autorisée

Basé sur : Filipits et al. (2011): A new molecular predictor of distant recurrence in ER-positive, HER2-negative breast cancer adds independent information to conventional clinical risk factors. Clinical Cancer Research 17: 8012-8020.



EndoPredict® Report

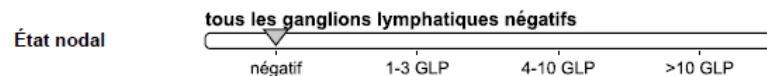
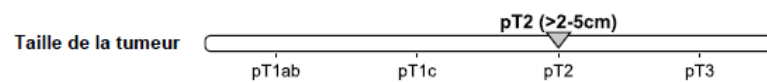
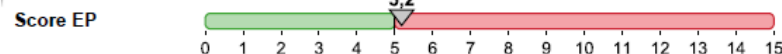


ID d'échantillon :

Note :

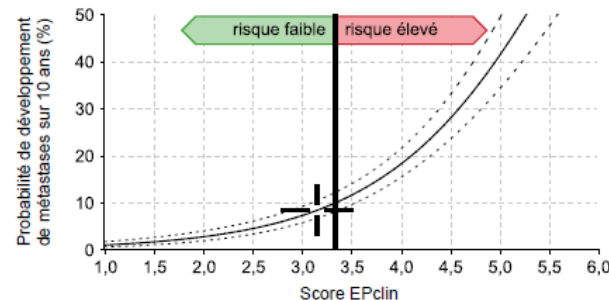
9 juin 2016 14:32 (CEST)

Caractéristiques de la tumeur



Évaluation du risque selon EPclin

EPclin combine le score EP, la taille de la tumeur et le nombre de ganglions lymphatiques positifs pour définir un score au pouvoir prédictif supérieur.



Score EPclin

3,1

Risque 10a EPclin*

8%

Classe EPclin

risque faible

* La probabilité de développement d'une métastase à distance sur 10 ans chez les patientes ayant suivi une thérapie endocrinienne de 5 ans est de 8%.

Validation par le pathologiste

Les contrôles indiqués ont été effectués et le résultat du test est valide.

Signature autorisée

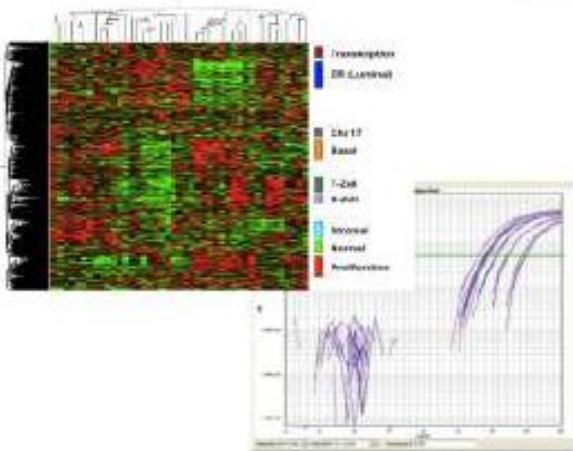
Basé sur : Filipits et al. (2011): A new molecular predictor of distant recurrence in ER-positive, HER2-negative breast cancer adds independent information to conventional clinical risk factors. Clinical Cancer Research 17: 8012-8020.



Test EndoPredict

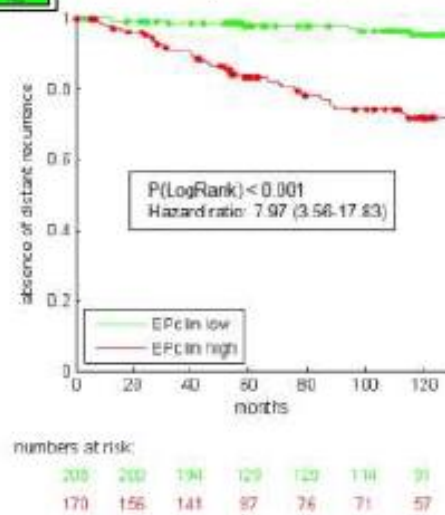
Training

**Multicenter
TAM Monotherapy
(n=964)**



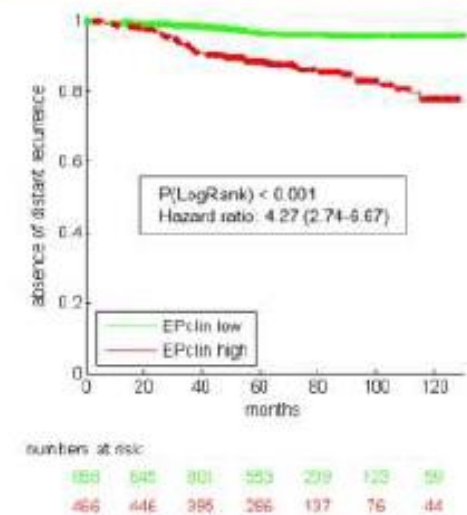
Validation I

**ABCSG 6
TAM
(n=378)**



Validation II

**ABCSG 8
TAM vs. TAM/ANA
(n=1.324)**

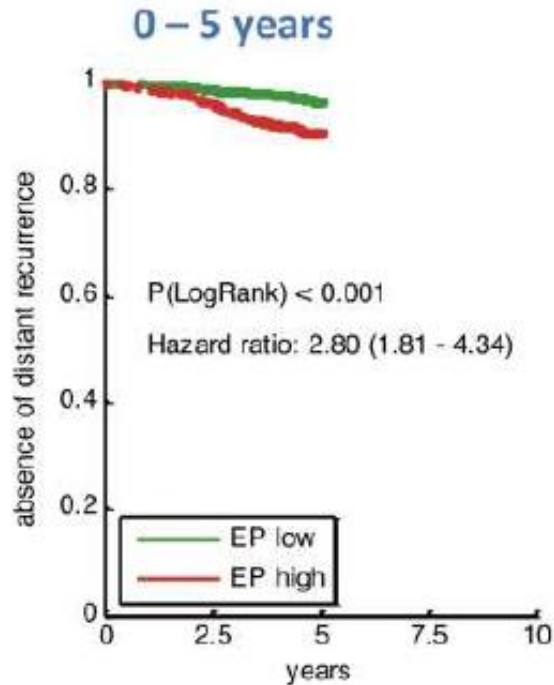


- Validated in two independent prospective-retrospective studies
- Evidence level: Ib (according to Simon et al., J Natl Cancer Inst, 2009)

Filipits et al., Clin Cancer Res. 2011
Dubsky et al., Ann Oncol. 2012

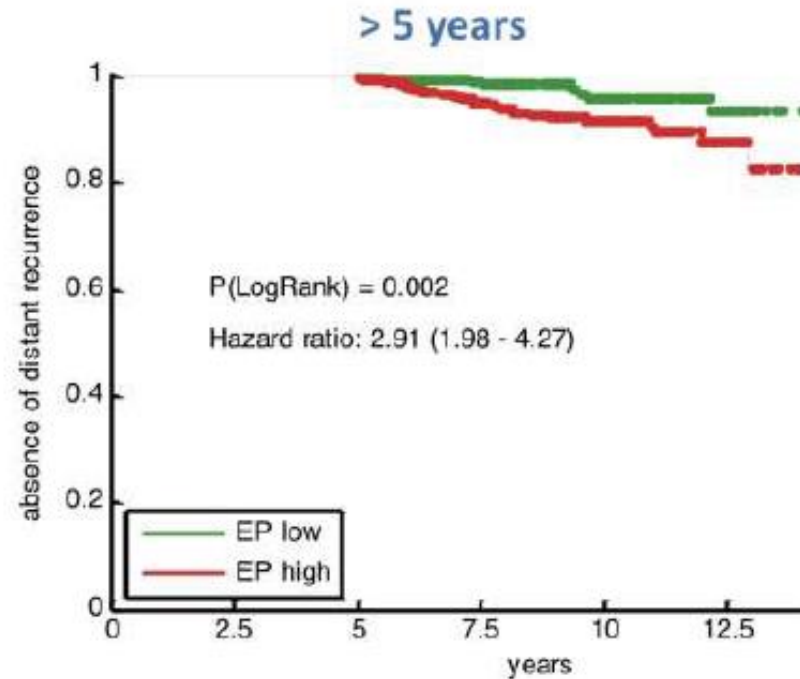
Test EndoPredict

Results



numbers at risk:

832	796	540
870	805	515



numbers at risk:

503	236	122	28
495	235	129	25

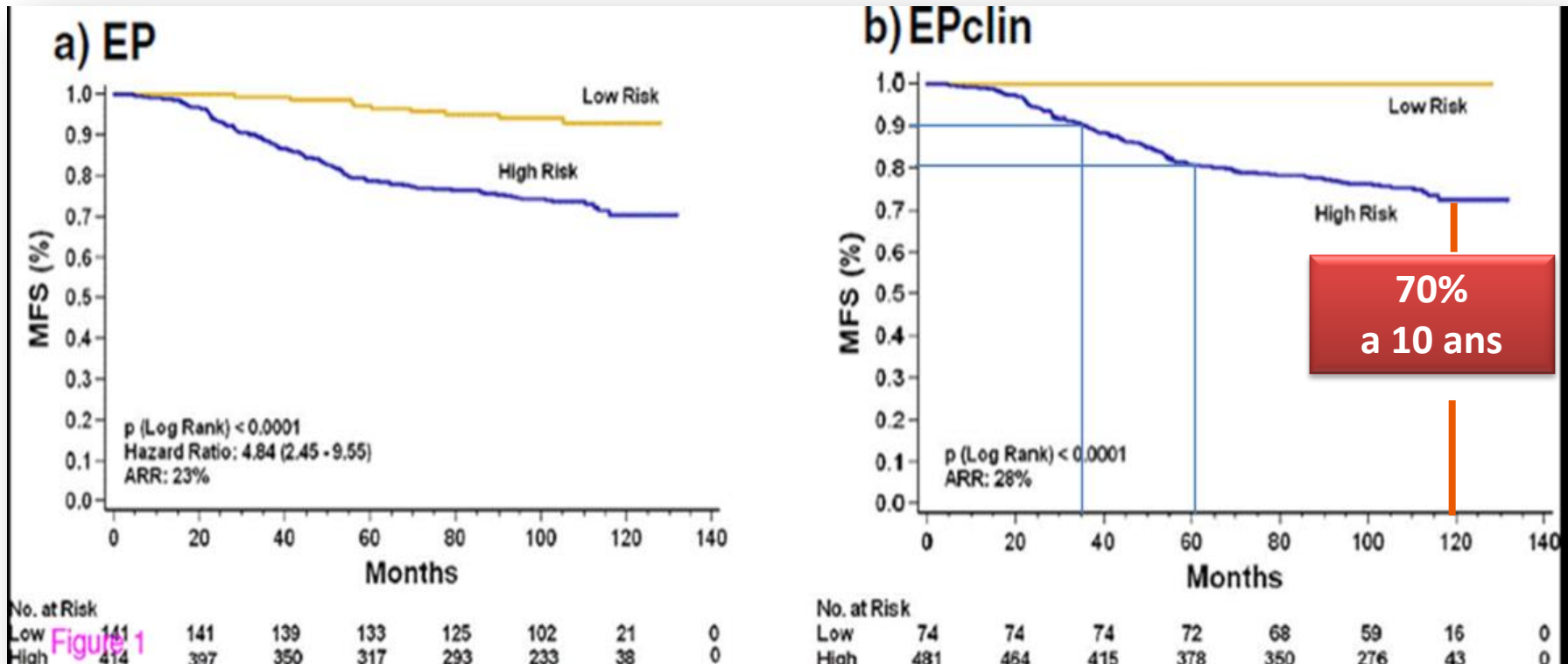
- The EndoPredict low-risk group (49%) had a significantly improved clinical outcome before and after 5 years of follow-up. 96.3% were free of distant metastases between 5 and 10 years in the EP low risk group.



Test EndoPredict

Clinical validation of the EndoPredict test in node-positive chemotherapy-treated ER+/HER2- breast cancer patients: results from the GEICAM/9906 trial

Breast Cancer Research 2014, 16:R38 doi:10.1186/bcr3642



EP low vs high risk

EPclin low vs high risk



JAMA Oncology | **Original Investigation**

Comparison of the Performance of 6 Prognostic Signatures for Estrogen Receptor–Positive Breast Cancer A Secondary Analysis of a Randomized Clinical Trial

Ivana Sestak, PhD; Richard Buus, PhD; Jack Cuzick, PhD; Peter Dubsy, MD; Ralf Kronenwett, MD;
Carsten Denkert, MD; Sean Ferree, PhD; Dennis Sgroi, MD; Catherine Schnabel, PhD;
Frederick L. Baehner, MD; Elizabeth Mallon, PhD; Mitch Dowsett, PhD

- Patient group:
 - primary breast cancer; ER+, HER2-; node negative and positive; postmenopausal
- Treatment:
 - 5 years endocrine therapy only (tamoxifen or anastrozole monotherapy)
- Number of patients: 774
 - N0: 591 (76%)
 - N+ 1-3: 183 (24%)
- Primary endpoint: Time to distant recurrence
- Median follow-up: 10 years



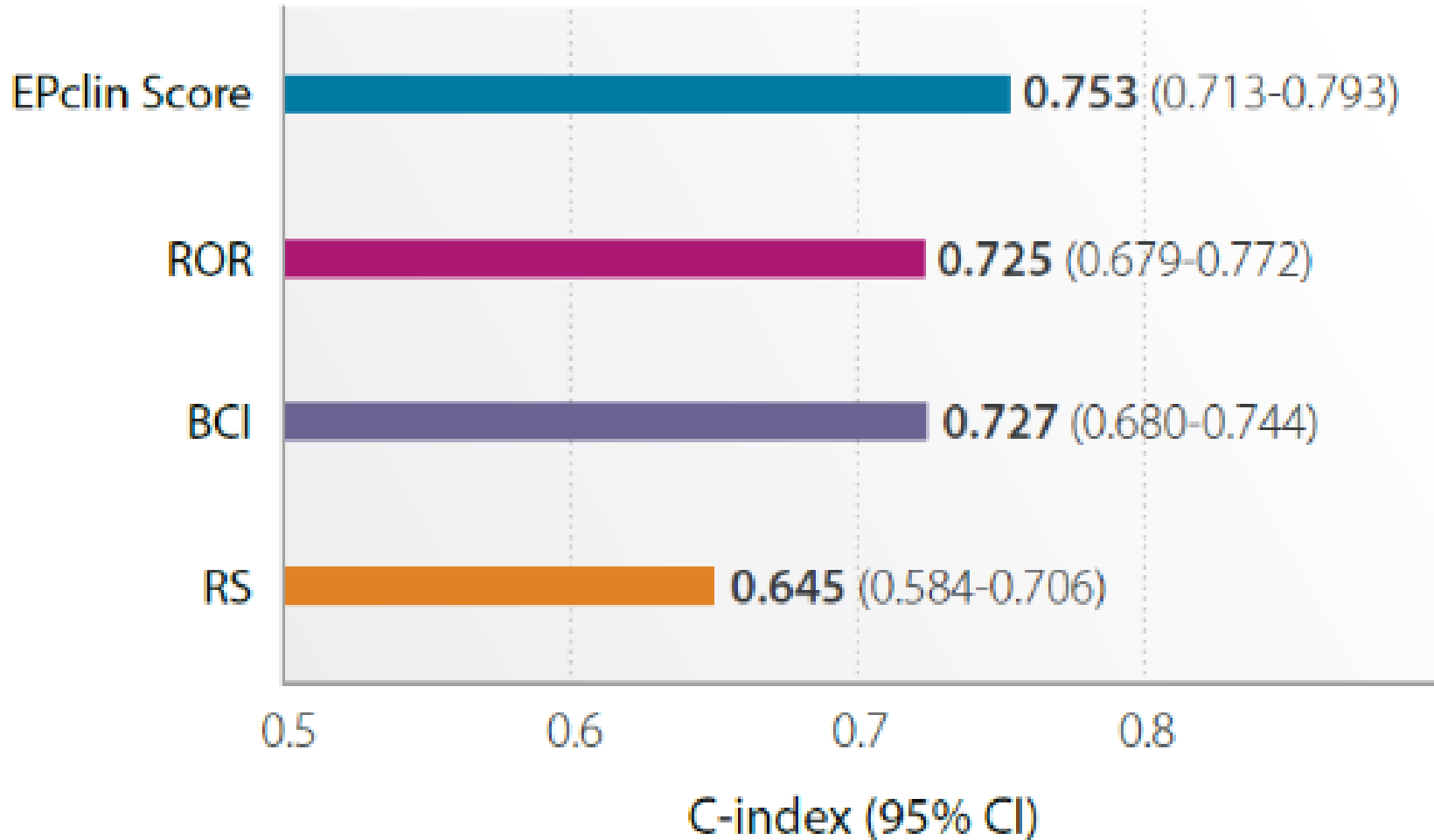
eTable 1. Baseline Characteristics According to Nodal Status

	Node-negative (N=591)	Node-positive (N=183)
Mean age, years (SD)	63.4 (7.9)	66.4 (8.3)
Mean BMI, kg/m ² (SD)	27.3 (4.9)	26.7 (4.7)
Grade		
1	23.2%	21.9%
2	59.7%	58.5%
3	17.1%	19.7%
Mean tumour size, mm (SD)	17.6 (8.5)	24.2 (12.2)
Distant recurrence		
0-10 years	58 (9.8%)	40 (21.9%)
5-10 years	34/535 (6.4%)	21/154 (13.6%)

SD=Standard Deviation, kg-kilogram, m=metre, mm=millimetre

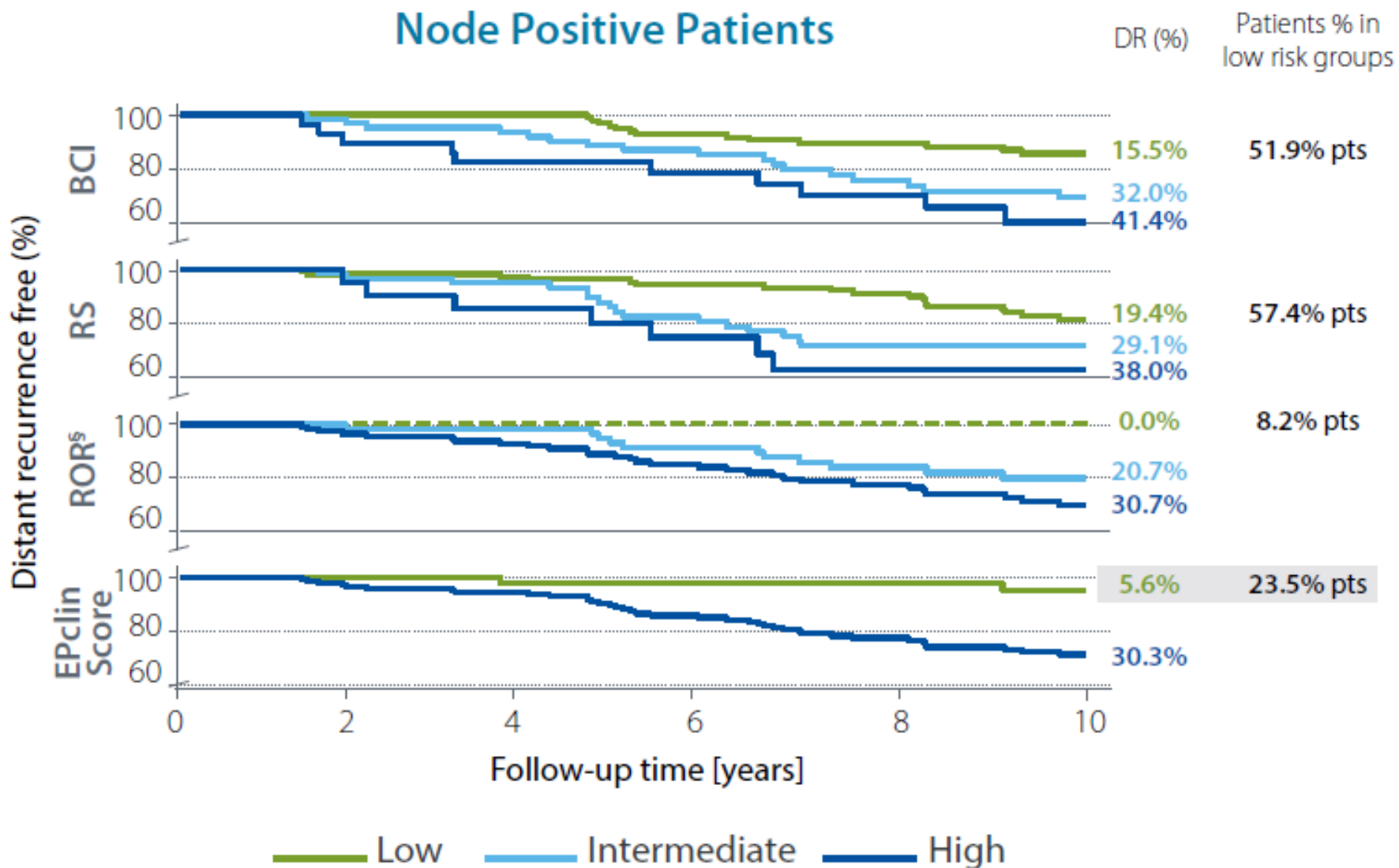


All Patients - Years 0-10



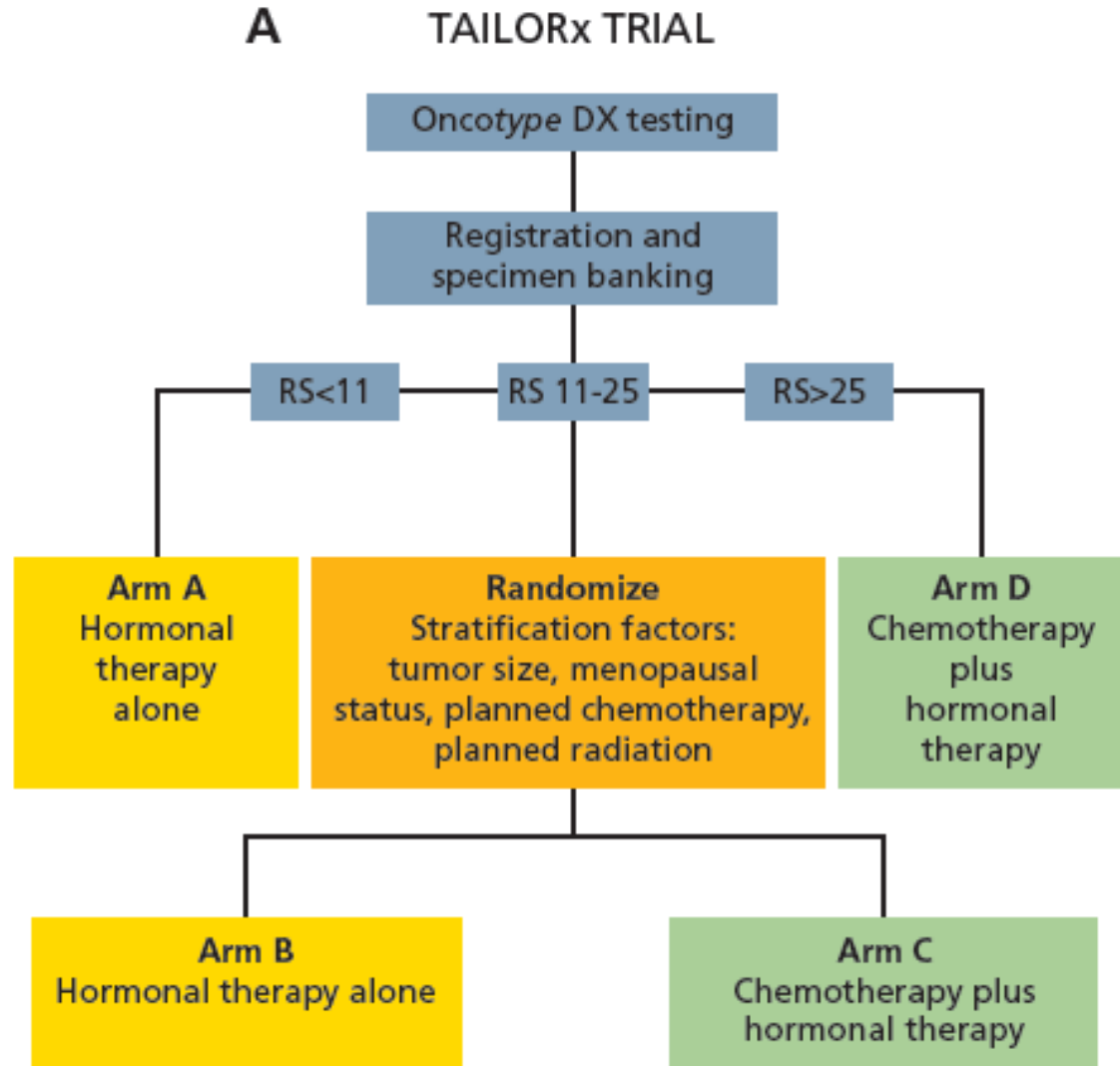


Node Positive Patients



Essai prospectif TAILORx

- N-
- RH+
- HER2-
- Pre et Post Menop.



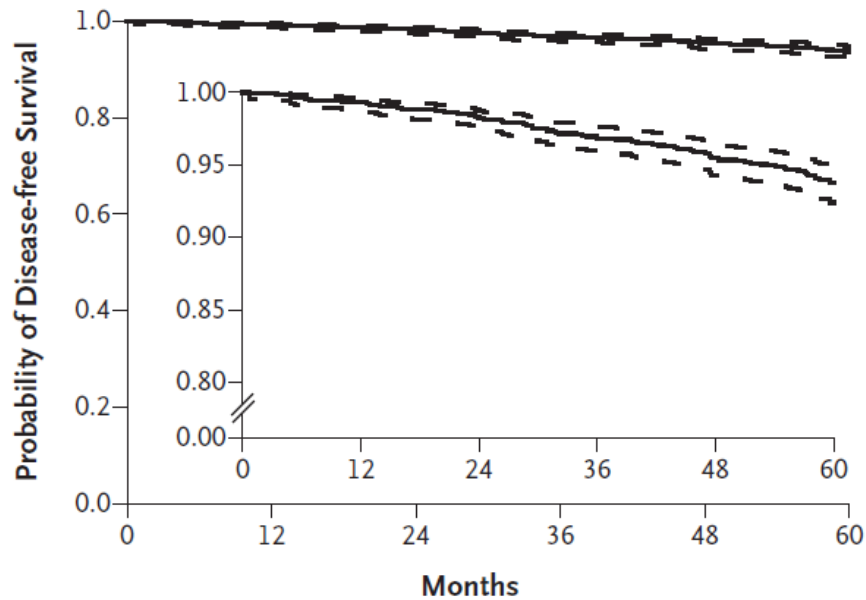
TAILORx: résultats de la cohorte bas risque (ce n'est pas l'obj principal)

Table 1. Characteristics of the Patients at Baseline, According to Recurrence-Score Cohort.*

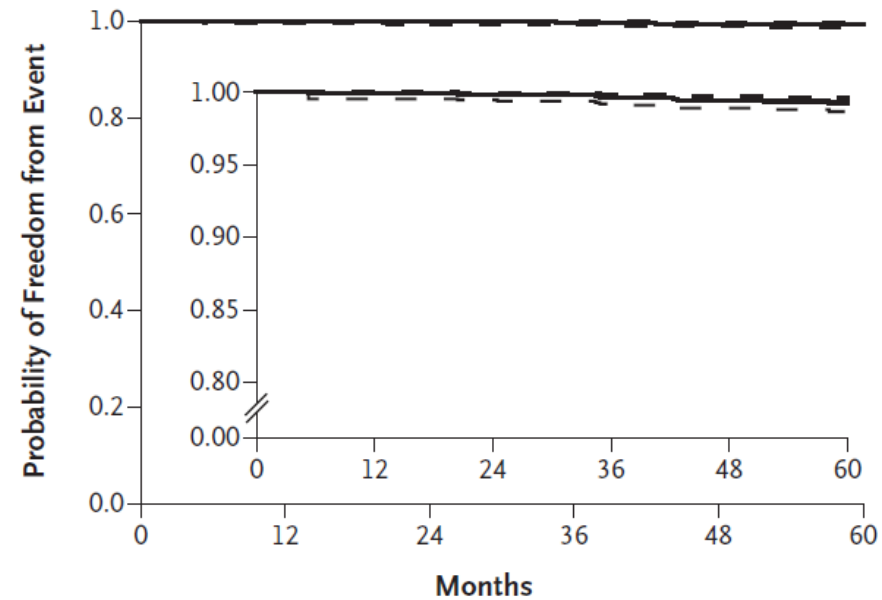
Characteristic	Recurrence Score, 0–10 (N = 1626)	Recurrence Score, 11–25 (N = 6897)	P Value
Tumor size in the greatest dimension			
Median (interquartile range) — cm	1.5 (1.2–2.0)	1.5 (1.2–2.0)	0.31
Mean — cm	1.74±0.77	1.71±0.79	0.23
Distribution — no./total no. (%)			0.42
<1.0 cm	128/1626 (8)	568/6883 (8)	
1.0–1.9 cm	993/1626 (61)	4270/6883 (62)	
2.0–2.9 cm	366/1626 (23)	1543/6883 (22)	
3.0–3.9 cm	104/1626 (6)	358/6883 (5)	
≥4.0 cm	35/1626 (2)	144/6883 (2)	
Histologic grade of tumor — no./total no. (%)			<0.001
Low	530/1578 (34)	1941/6665 (29)	
Intermediate	937/1578 (59)	3812/6665 (57)	
High	111/1578 (7)	912/6665 (14)	

TAILORx: résultats de la cohorte bas risque RS < 11

A Invasive Disease-free Survival



B Freedom from Recurrence of Breast Cancer at Distant Site



IDFS @ 5 ans : 93.8%

MFS @ 5 ans : 99.3%

MINDACT

« **Microarray for node negative disease may avoid chemotherapy »**

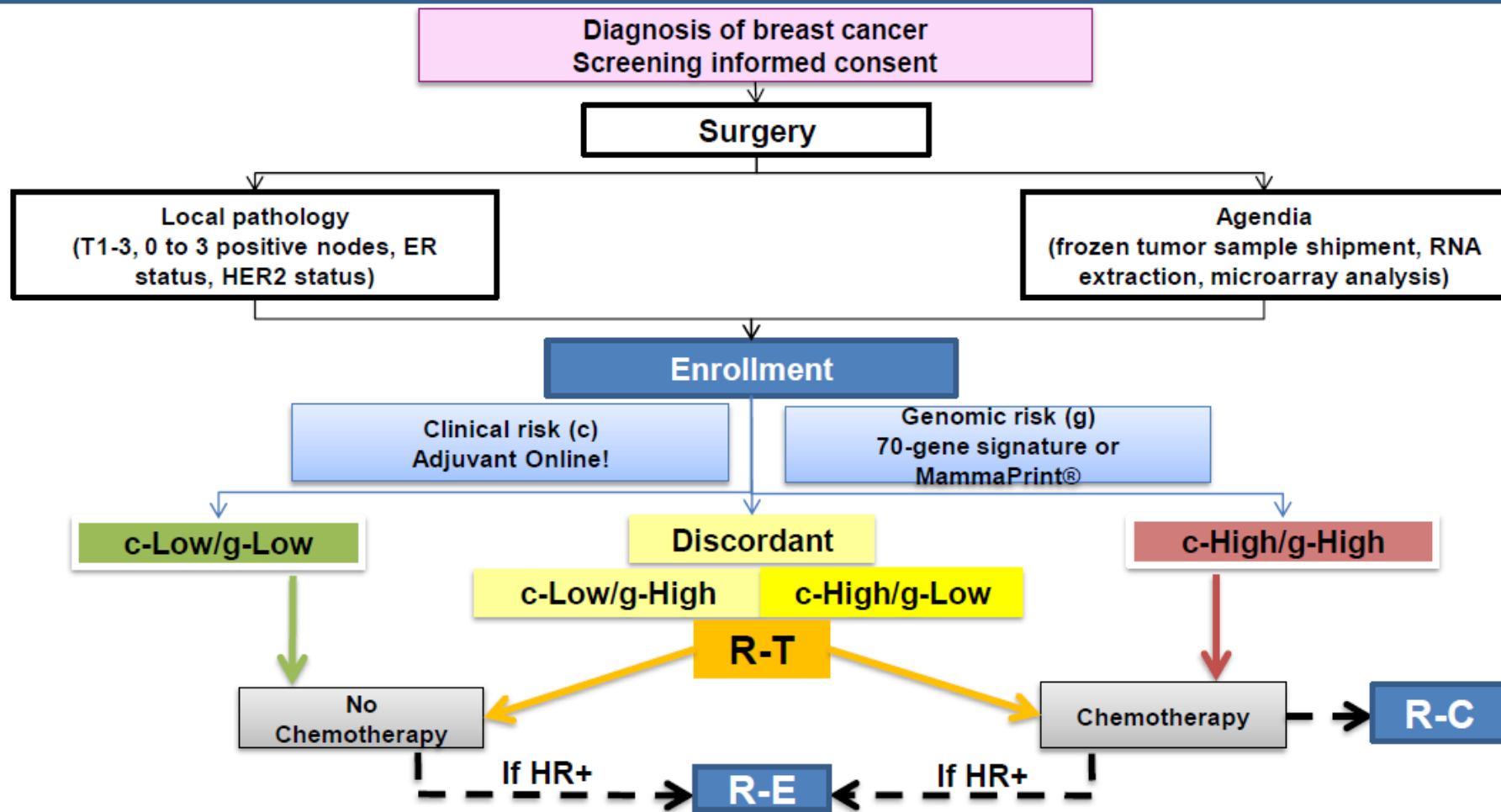
**Primary analysis of the EORTC 10041/ BIG 3-04 MINDACT study:
A prospective, randomized study evaluating the clinical utility of the 70-gene
signature (MammaPrint®) combined with common clinical-pathological criteria
for selection of patients for adjuvant chemotherapy in breast cancer
with 0 to 3 positive nodes**

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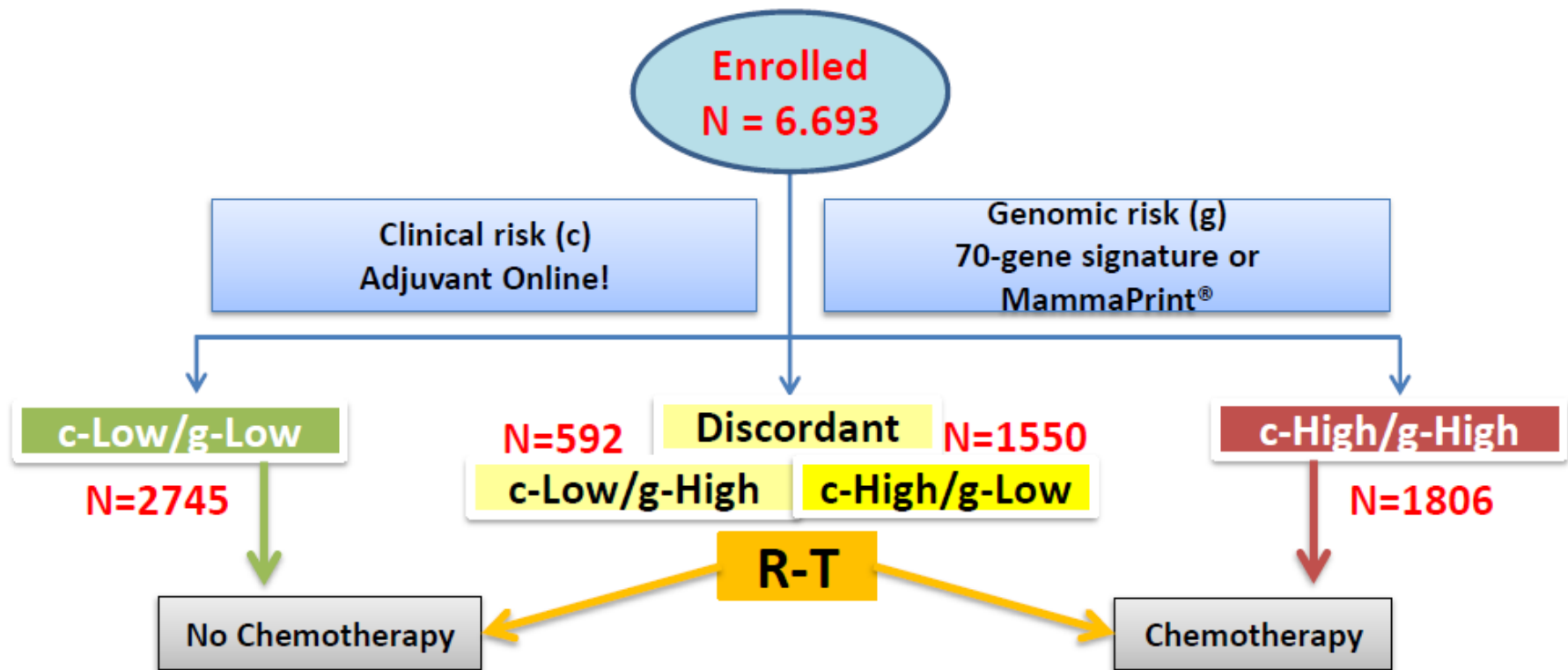
***On behalf of the European Commission supported TRANSBIG consortium
and MINDACT investigators***

Presented at AACR, April 18, 2016

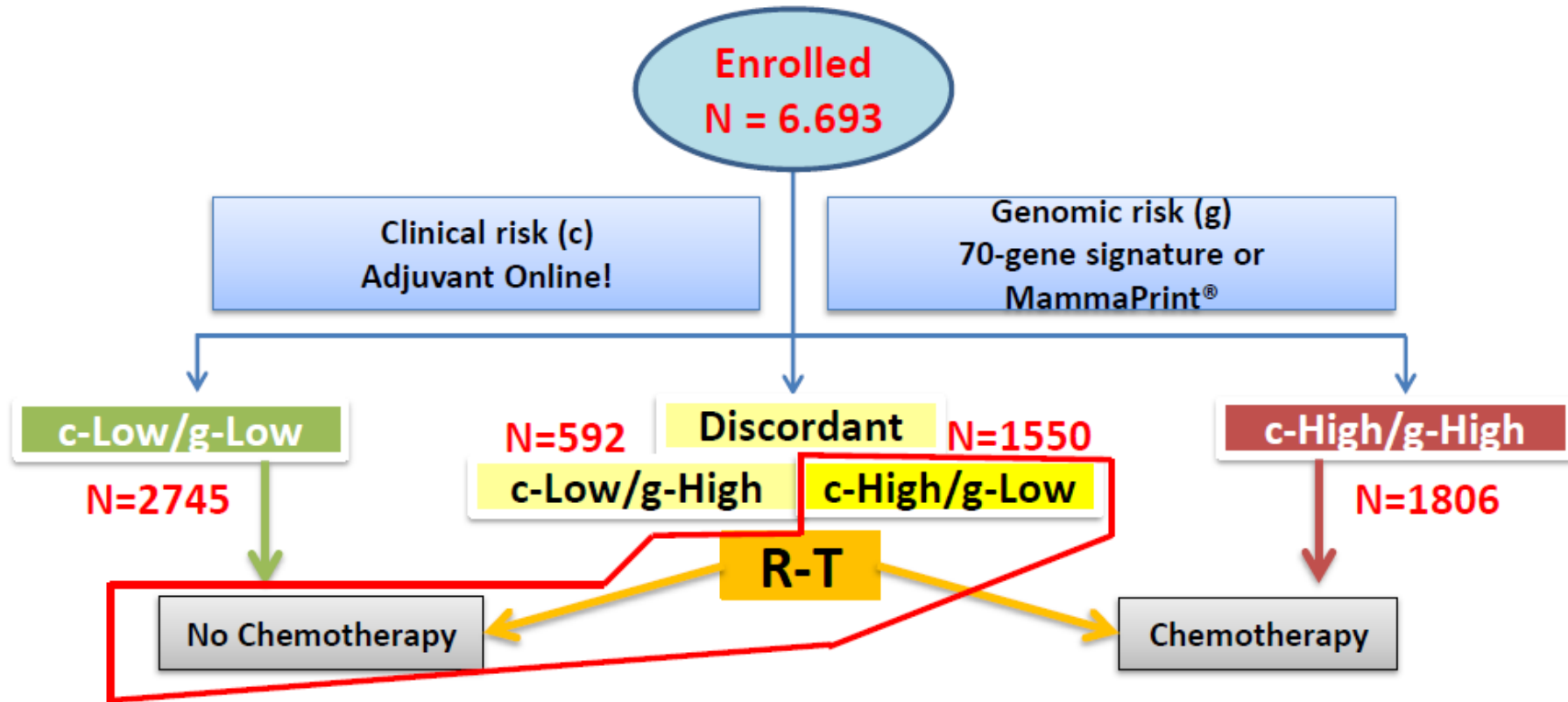
The MINDACT study design



The MINDACT study: Patient enrollment

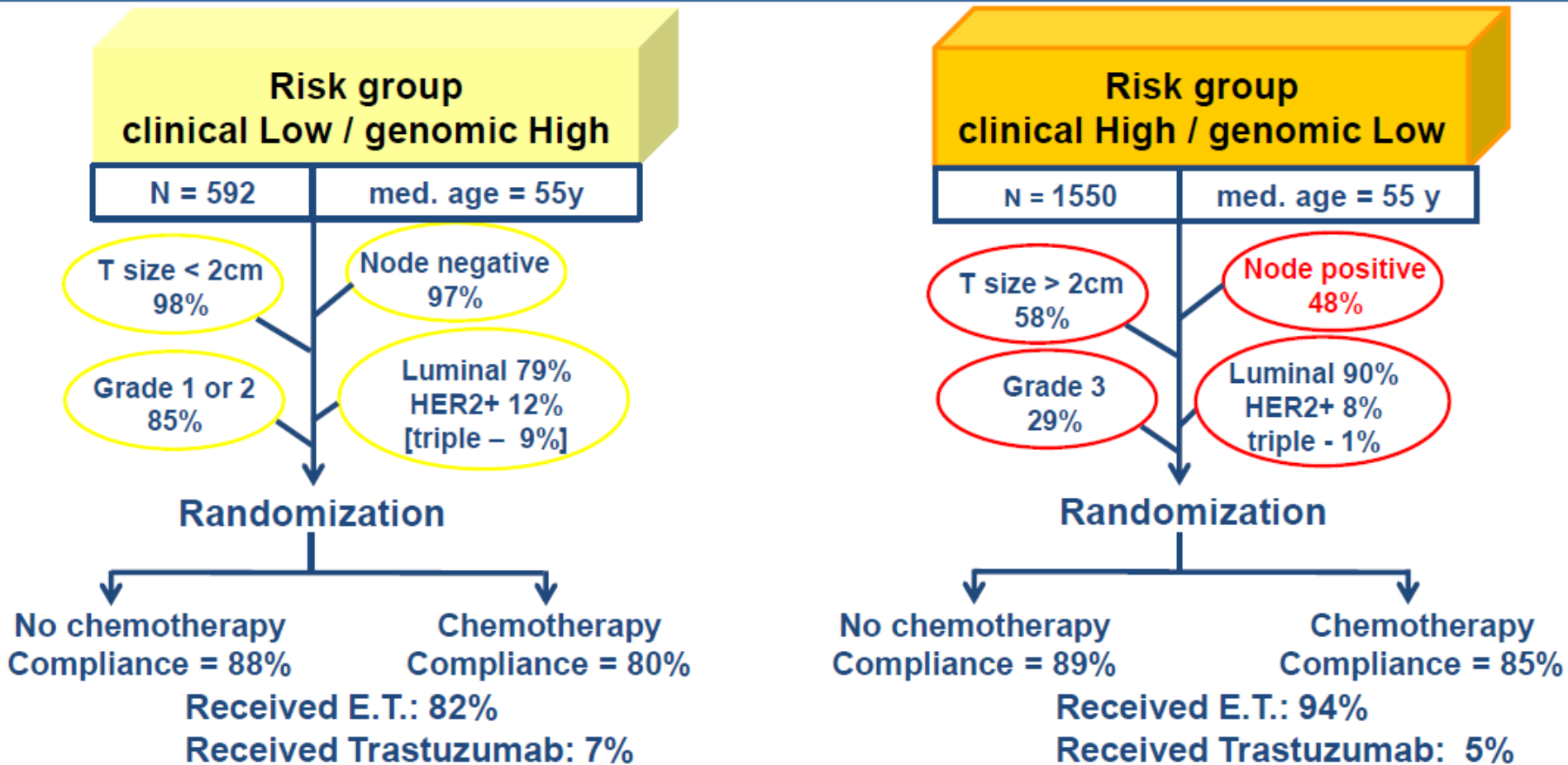


The primary analysis population



N = 644

MINDACT: patient demographics and compliance with assigned therapy



B) DISCORDANT RISK GROUPS: PRIMARY TEST

The primary analysis population

Discordant risks

c-Low /g-High

c-High/g-Low

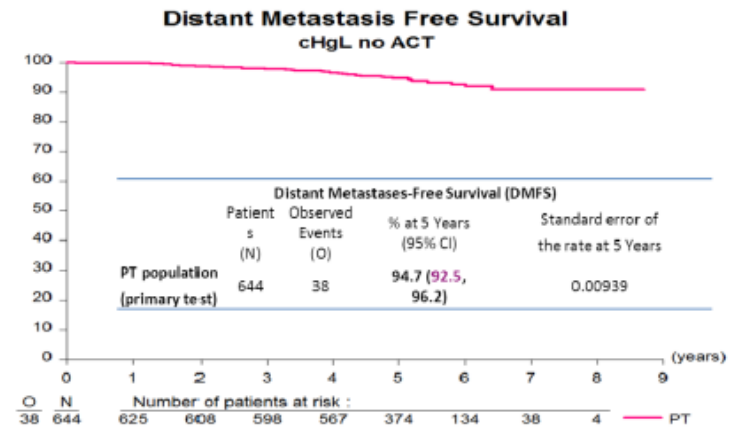
RANDOMIZATION

No chemotherapy
N = 748

CT

No change in risk
post enrollement
and no CT received
N = 644

The primary statistical test (DMFS at 5Y)



Null Ho set à 92%

Observed 5Y DMFS = 94.7%

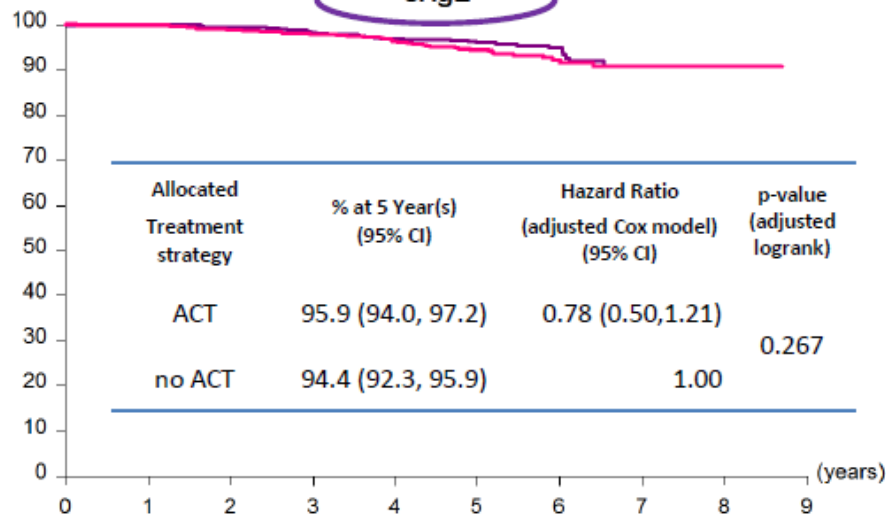
95% CI ≈ 92.5 – 96.2% excludes 92% !!!

Efficacy: ACT vs no ACT in discordant risk groups

Intent-to-treat analysis

Distant Metastasis Free Survival

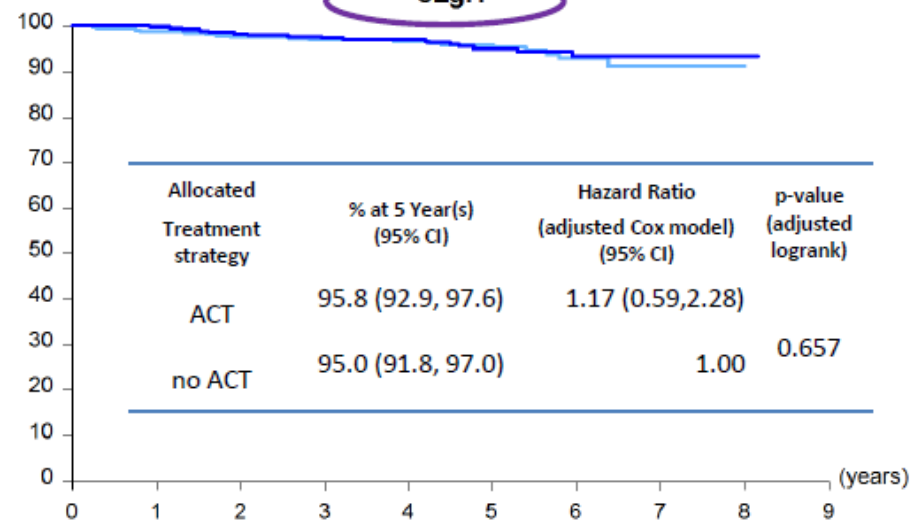
cHgL



O	N	Number of patients at risk :								Allocated to:	
34	749	714	698	677	611	346	145	41	3	—	ACT
46	748	727	708	696	655	424	160	41	4	—	no ACT

Distant Metastasis Free Survival

cLgH



O	N	Number of patients at risk :									Allocated to:	
18	344	321	316	306	281	179	81	22	0	—	ACT	
17	346	336	327	319	291	178	82	24	3	—	no ACT	

Aucun bénéfice du test pour les c-Low !

c-Low/g-High CT vs no CT per protocol population						
	Treatment received	Pts	Observed Events	% at 5 Year(s) (95% CI)	Hazard Ratio (adjusted Cox model) (95% CI)	p-value (adjusted logrank)
DMFS	CT	224	11	96.1 (92.4, 98.1)	0.90 (0.40,2.01)	0.798
	no CT	254	14	93.9 (89.6, 96.5)		
DFS	CT	224	17	92.7 (87.9, 95.7)	0.74 (0.40,1.39)	0.355
	no CT	254	25	90.5 (85.7, 93.8)		
OS	CT	224	5	98.1 (94.9, 99.3)	0.72 (0.23,2.24)	0.572
	no CT	254	8	97.0 (93.8, 98.6)		

Intérêt discutable du test pour les c-High...

c-High/g-Low CT vs no CT per protocol population						
	Treatment received	Patients	Observed Events	% at 5 Year(s) (95% CI)	Hazard Ratio (adjusted Cox model) (95% CI)	p-value (adjusted logrank)
DMFS	CT	592	22	96.7 (94.7, 98.0)	0.65 (0.38,1.10)	0.106
	no CT	636	37	94.8 (92.6, 96.3)	1.00	
DFS	CT	592	39	93.3 (90.7, 95.2)	0.64 (0.43,0.95)	0.026
	no CT	636	66	90.3 (87.6, 92.4)	1.00	
OS	CT	592	10	98.8 (97.4, 99.5)	0.63 (0.29,1.37)	0.245
	no CT	636	18	97.3 (95.6, 98.4)	1.00	



Cas clinique 1

- Patiente de 55 ans sans ATCD non ménopausée
- Dépistage: mammo normale , écho : sur rayon 2h à 4 cm du mamelon, masse hypoéchogène atténuante ACR4c de 9 mm du sein gauche , aires ganglionnaires sans gg suspect
- Biopsie sous écho: lobulaire infiltrant SBR2, ER+,PR+, HER2 neg.
- Palpation mammaire et des aires ganglionnaires normale
- IRM mammaire: lésion unique gauche ACR6 de 10 mm sans gg suspect
- Tumorectomie après repérage écho et ganglion sentinelle:
- Carcinome infiltrant de type lobulaire pléomorphe mesurant 7x6 mm de grade 2 (ER+,PR+,HER2-),pas de CIS, pas d'embolie vasculaire
- Marges chirurgicales: résection in sano avec marge minimale de 2 mm en externe
- Ganglion sentinelle: 1 ganglion siège d'une micrométastase de 0,5mm sans rupture capsulaire et 2 ganglions lymphatiques histologiquement normaux

Que faites-vous si la signature multigénique est en faveur d'un risque de rechute faible ? chimio ou pas ?



Cas clinique 2

- Patiente de 60 ans ménopausée sans ATCD
- Dépistage retrouvant sur la mammographie une opacité correspondant en écho à une image ACR5 de 10 mm QSI sein droit , aires ganglionnaires axillaires normales en écho
- Palpation retrouvant un nodule de 10 mm du QSI du sein droit sans adénopathie axillaire palpable
- La biopsie retrouve un carcinome infiltrant SBR1 ER+PR+HER2-
- Tumorectomie avec ganglion sentinelle: carcinome infiltrant SBR1 ER+++100%, PR++80%,HER2-,pas d'embolie, exérèse in sano avec marges à 3mm en interne.
- 2 ganglions sentinelles dont 1 sur les 2 présente une macrométastase de 3mm sans effraction capsulaire

Que faites-vous si la signature multigénique est en faveur d'un risque de rechute faible ? chimio ou pas ?