

Sono utili i test genomici per la valutazione della prognosi?

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Disclosures



Scientific advisory board, meeting, congress:

Celgene,
Lilly,
Novartis,
Roche,
Pfizer,
Astra Zeneca

My Outline

- * **How to show the relationship between Genomic Test and Prognosis?**
- * **From predictive test to prognostic test: is it possible?**
- * **Point on locally advanced disease and prognosis by genomic test**
- * **Metastatic disease and genomic: towards a response through biomarkers**

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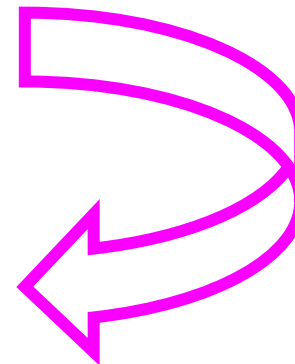
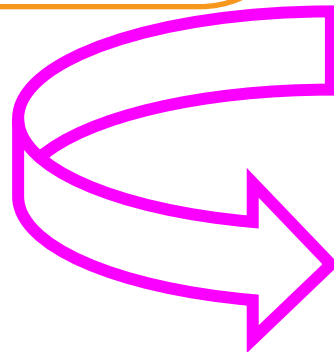
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Prognostic Versus Predictive Value

Prognostic Test/Biomarker

A prognostic test/biomarker provides information on a cancer outcome

(disease recurrence, disease progression, death for cancer)



Adjuvant Treatment Decisions Are Driven by Both Prognostic and Predictive Factors

Prognostic factors: information on outcomes
(eg, recurrence rate)

Predictive factors: degree of response to a specific therapy

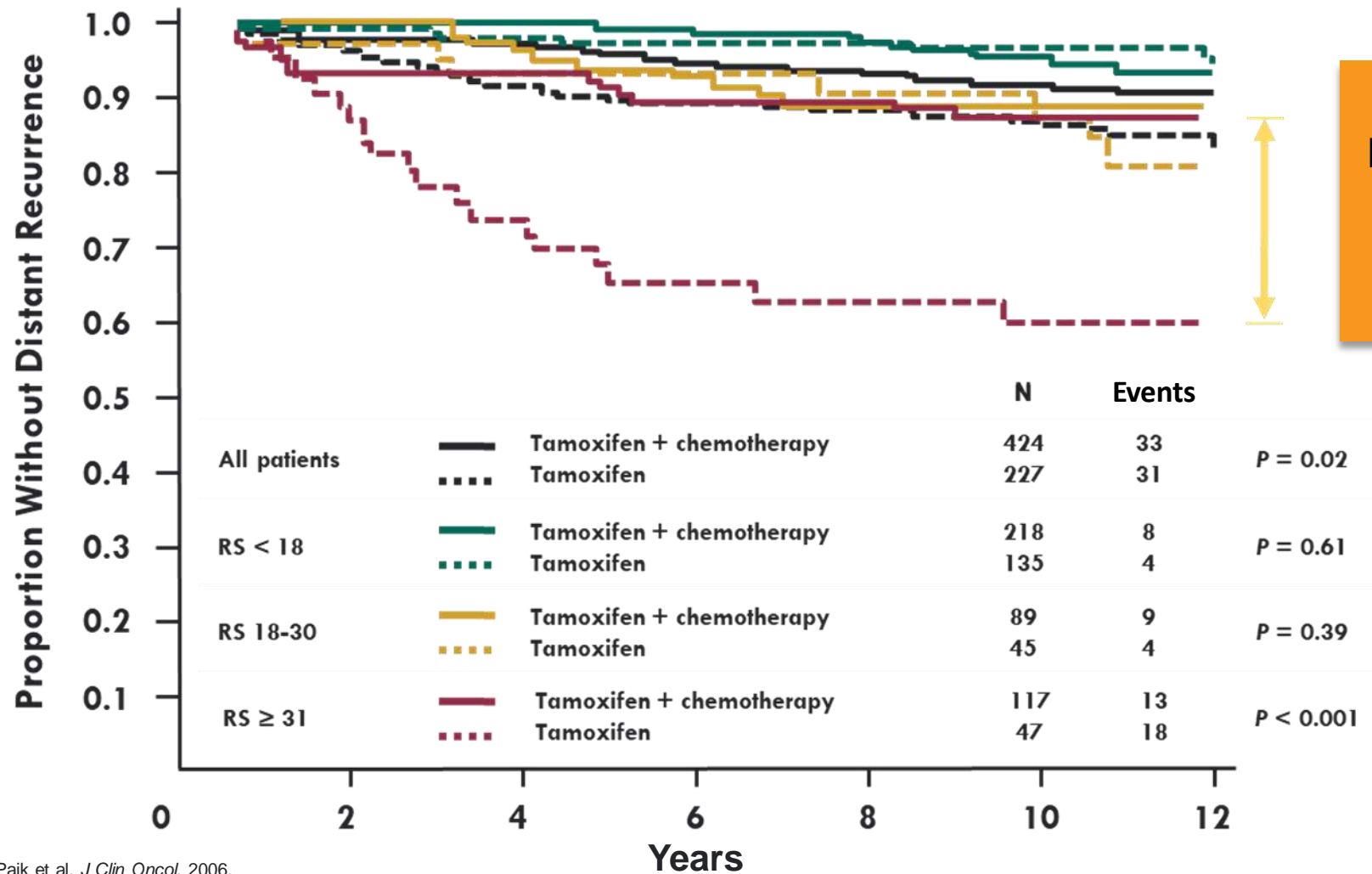
- Age
- Nodal status
- Tumor size
- Tumor Grade
- HER2
- ER/PR

- ER
- HER2
-

What do we have over?

BRS Test Predicts Those Patients Who Do and Do Not Derive Benefit From Chemotherapy

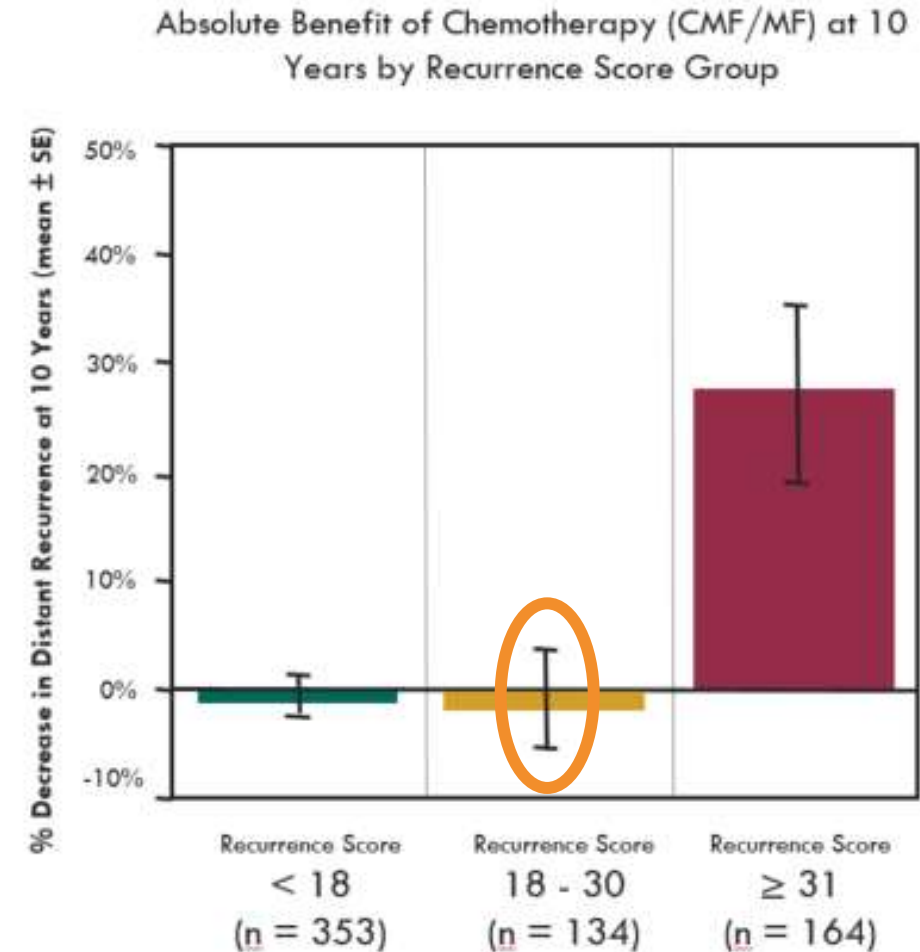
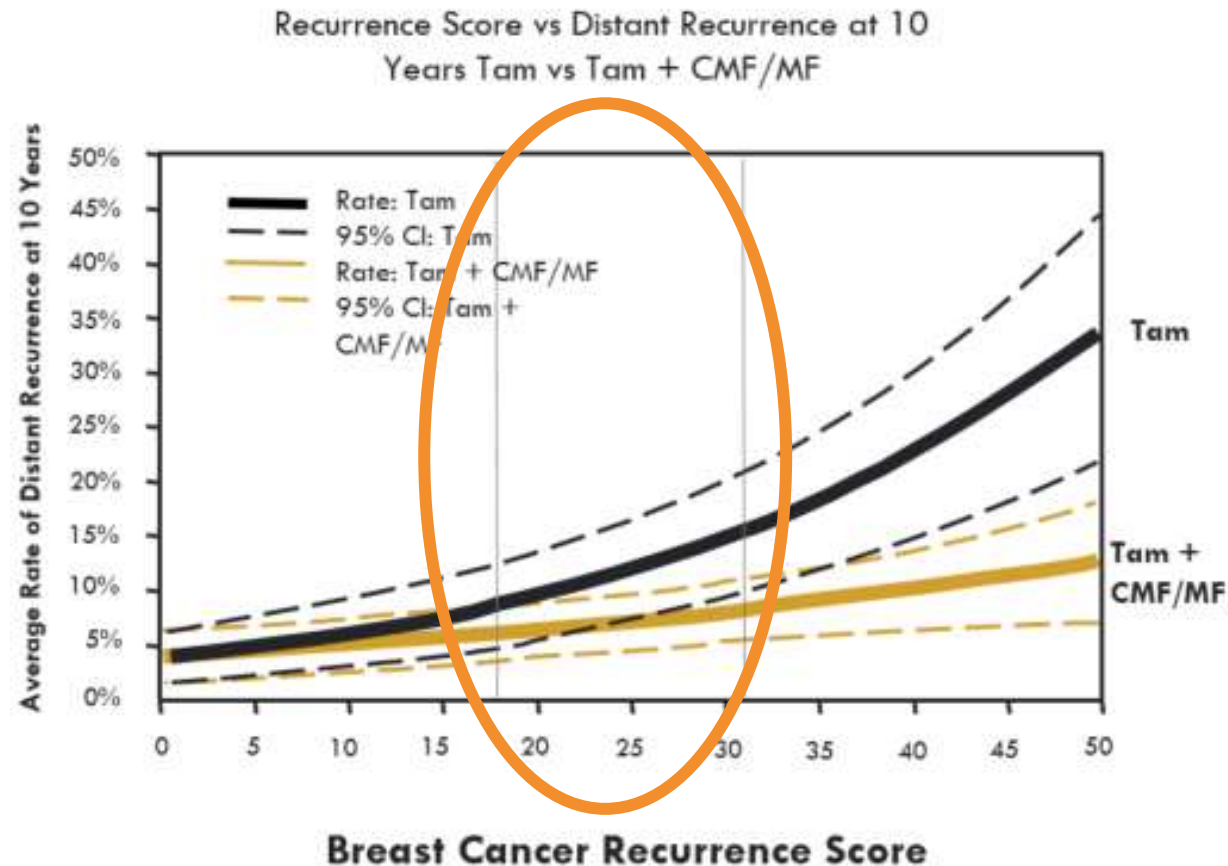
NSABP B-20: Validation Study for Prediction in Node-Negative Patient Population



**PATIENTS WITH HIGH RS ≥ 31
28% absolute benefit from
tamoxifen + chemotherapy**

Interaction $P = 0.038$

Rationale for Investigating Chemotherapy Benefit in Intermediate Oncotype DX Breast RS

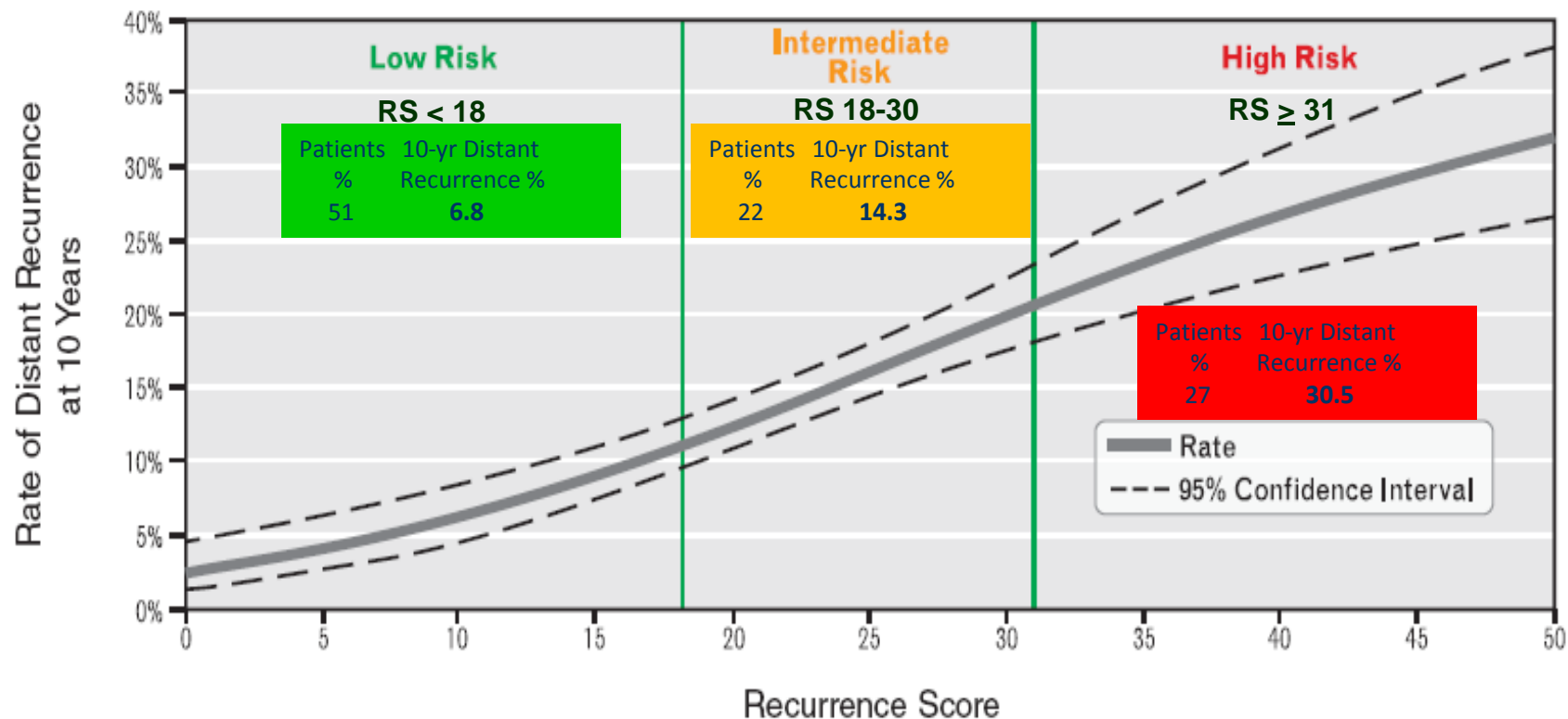


The RS is a continuous predictor of the Risk of Distant Recurrence

NSABP-14 CLINICAL VALIDATION

668 pts (stage I-II, N-, ER+ treated with 5 yrs of TAM)

Recurrence Score as Continuous Predictor

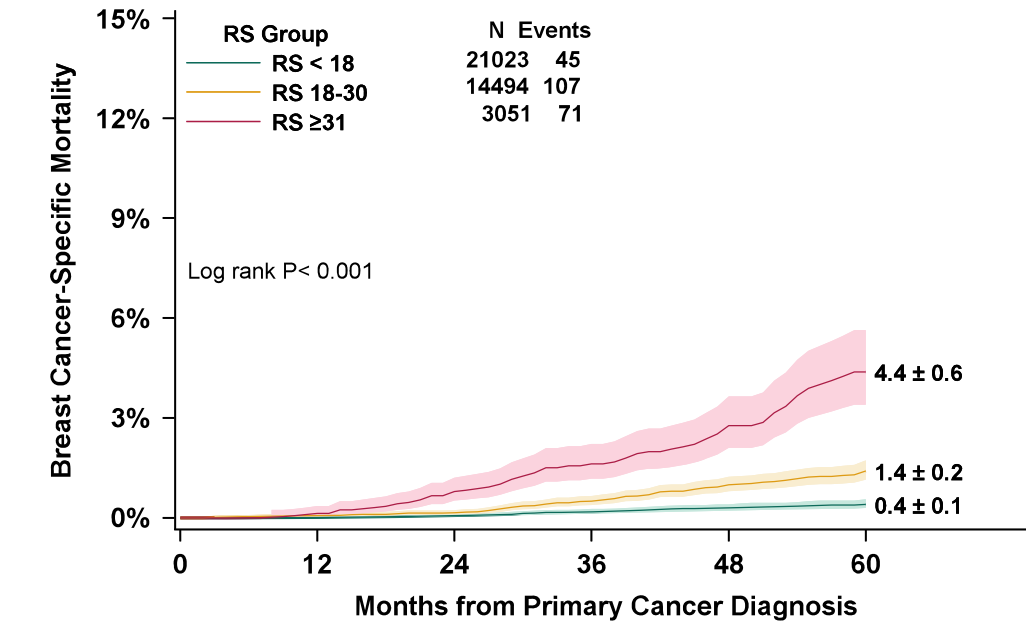


Paik S, NEJM 351(27):2817, 2004

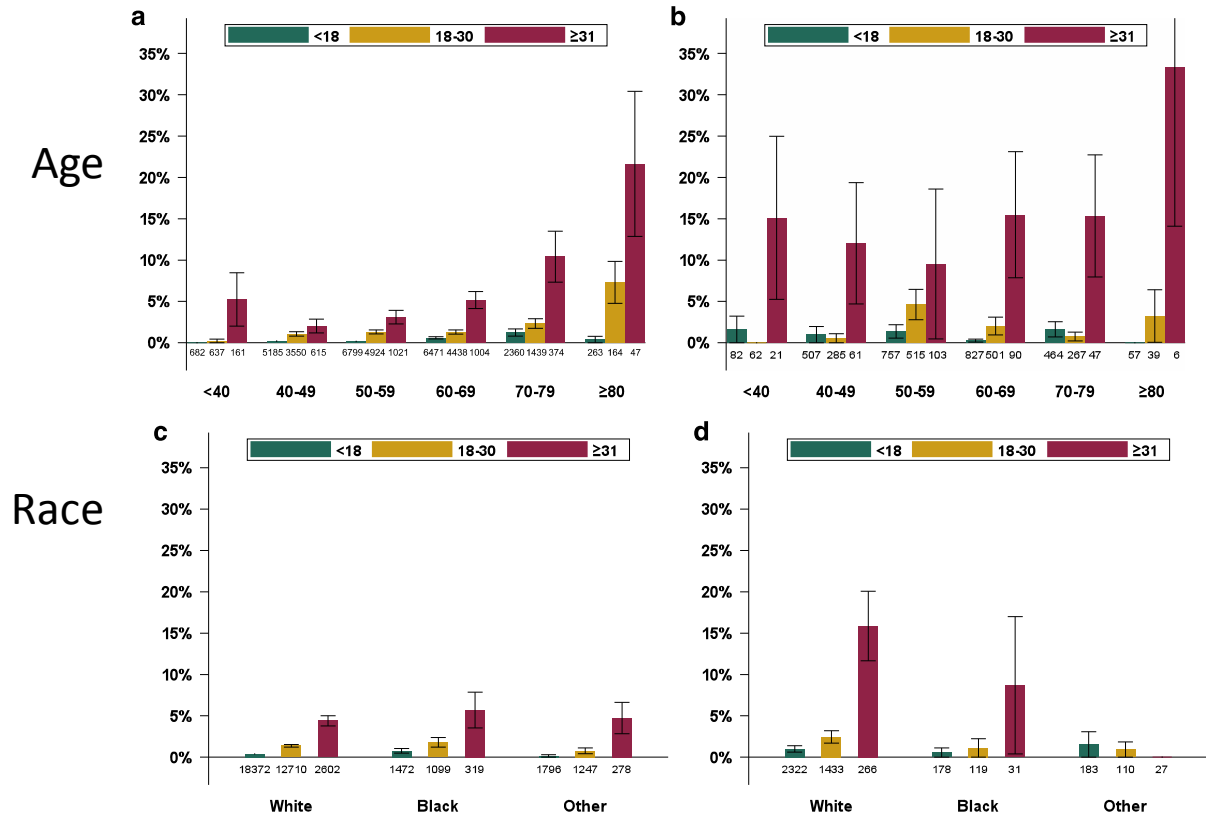
2019 carcinoma mammario: i traguardi raggiunti e le nuove sfide

Recurrence Score group was significantly prognostic *BC specific mortality*

38.568 pts



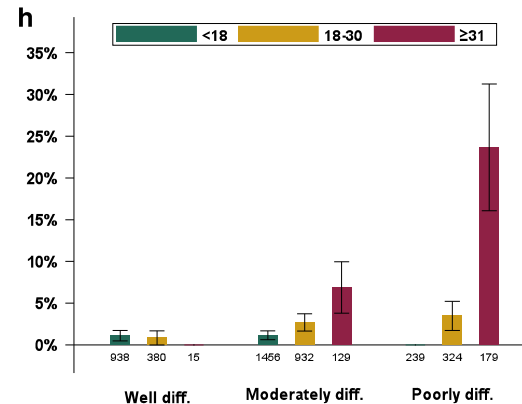
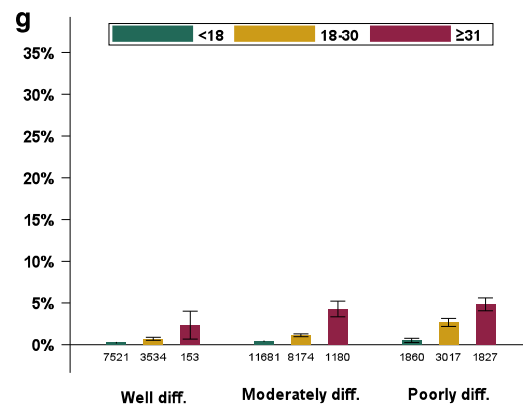
RS < 18	21023	20481	15685	11543	7551	4200
RS 18-30	14494	14138	11011	8247	5624	3369
RS ≥ 31	3051	2979	2313	1731	1153	670



Petkov et al, npj Breast 2016

Recurrence Score group was significantly prognostic *BC specific mortality* 38,568 pts

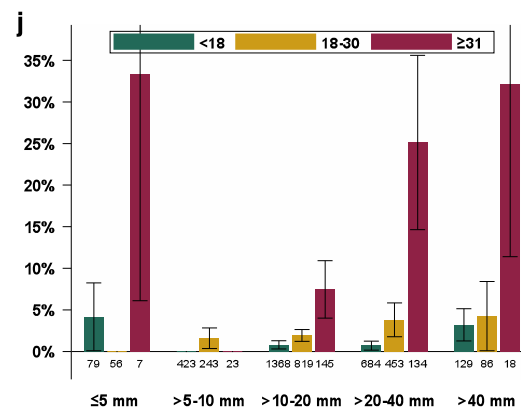
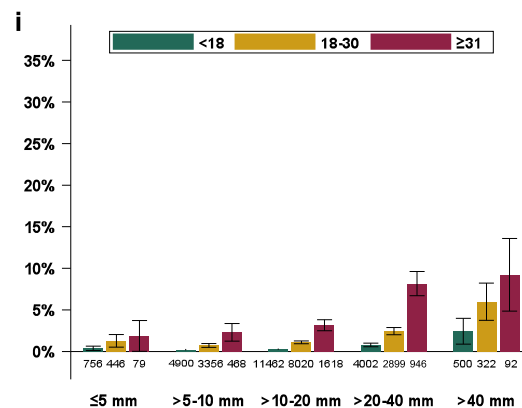
Grade



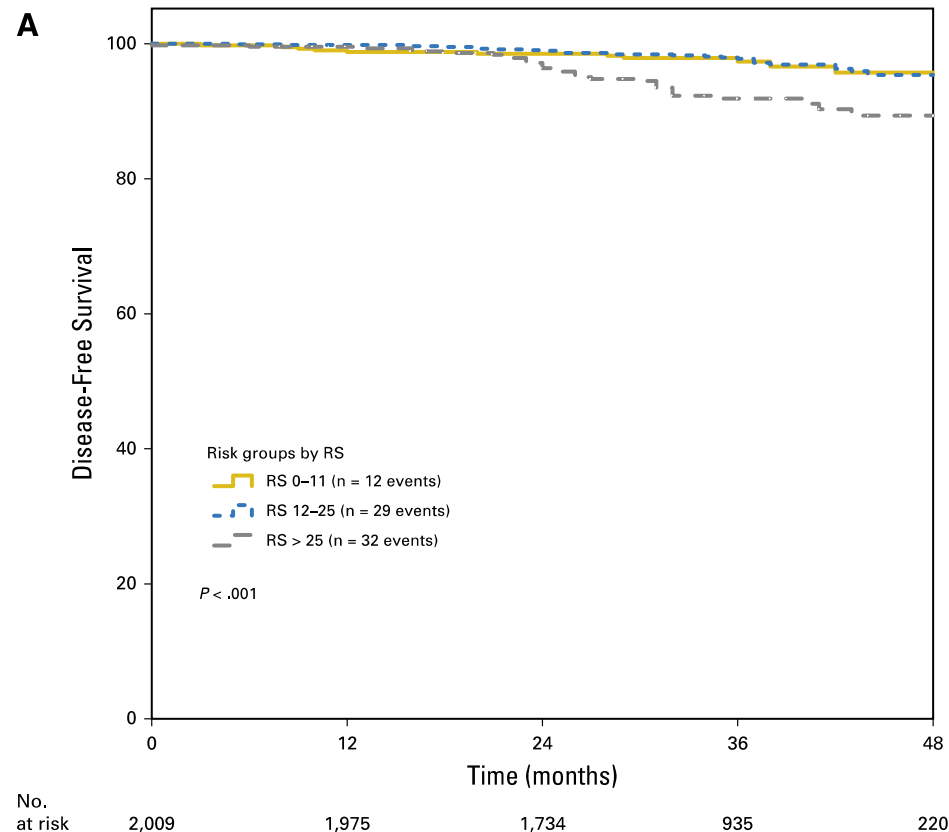
Characteristic	Effect	Unadjusted HR (95% CI)	Unadjusted P value ^a	Adjusted HR (95% CI)	Adjusted P value ^a
RS group (versus RS < 18)	RS 18-30	3.1 (2.3, 4.3)	< 0.001	3.0 (2.1, 4.2)	< 0.001
	RS ≥31	11.0 (7.8, 15.5)		7.8 (5.3, 11.6)	

RS > 31 higher probability of BCSM

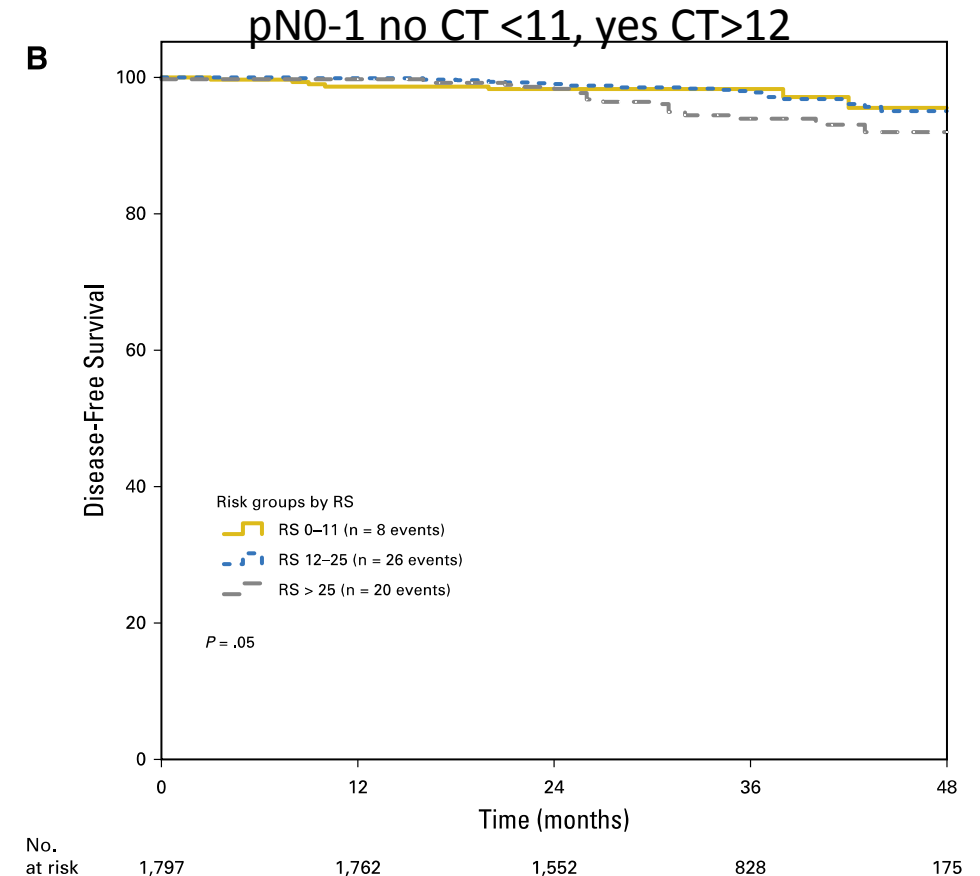
Size
Tumor



Petkov et al, npj Breast 2016



3-year DFS was **92%** (95% CI, 89.0% to 94.8%) in patients with high RS versus **98%** (95% CI, 96.8% to 98.8%) in pts with intermediate RS and **97%** in patients with RS < 11 (95% CI, 95.6% to 99.1%; $p = .001$)



3-year DFS was **95%** within the RS > 25 group (95% CI, 91.4% to 98.4%) versus **97.5%** (95% CI, 95.9% to 99.0%) within RS 12 to 25 group and **98%** (95% CI, 97.0% to 99.8%) within the RS < 11 group ($P = .05$ for RS > 25 v others)

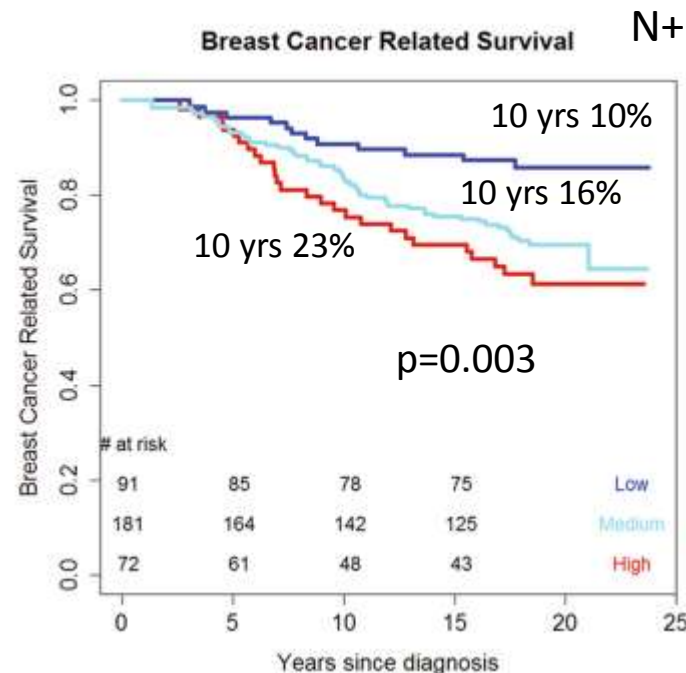
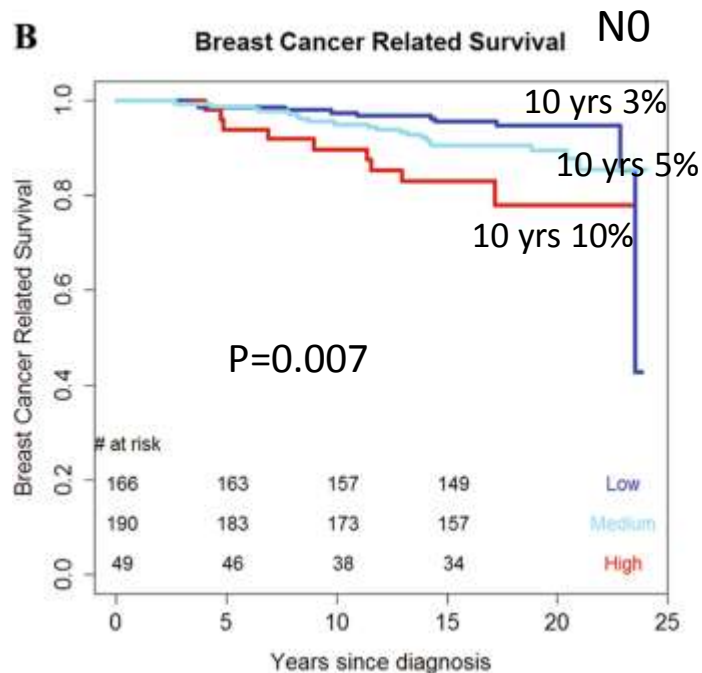
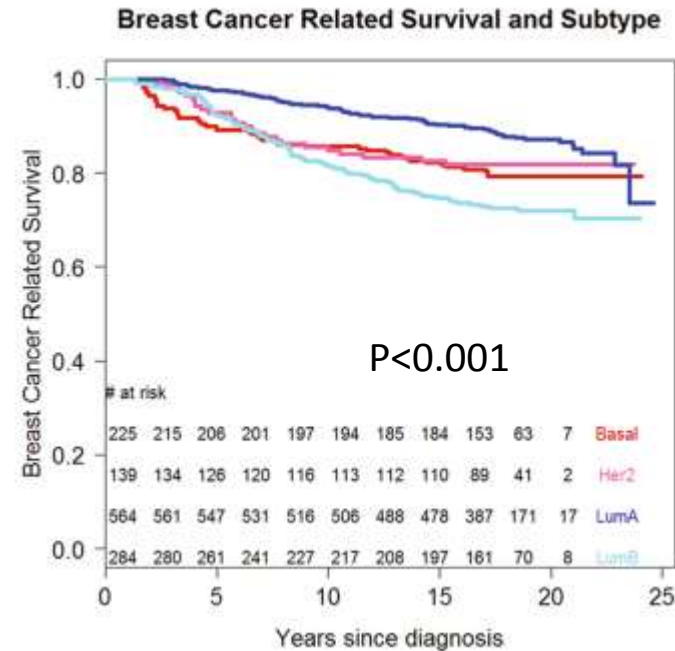
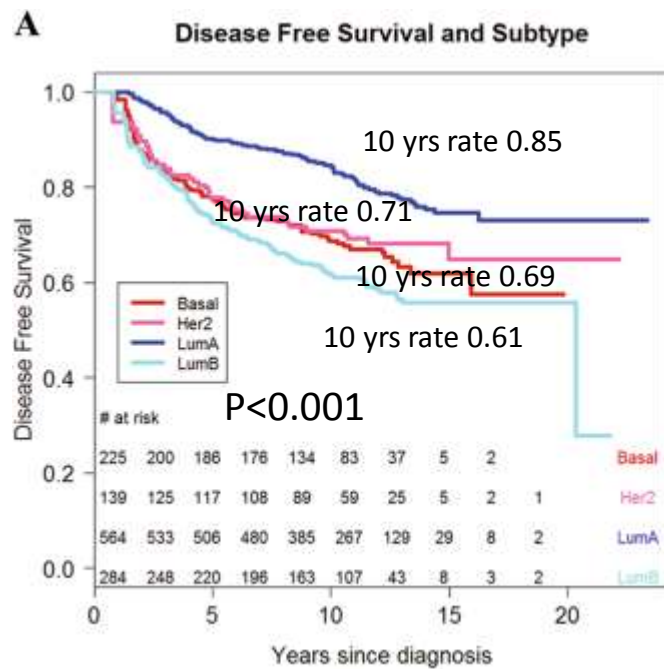
PAM50 signature and long-term breast cancer survival

Distribution of PAM50 subtypes by clinical characteristics (15 yrs follow up)

		45%	23%	18%	11%	3%	
		Luminal A %	Luminal B %	Basal-like %	Her2%	Normal %	P
	N	564	284	225	139	41	
Cancer stage							<0.0001
I	453	55.6	17.7	15.9	8.2	2.6	
IIA	432	42.4	22.9	20.1	11.1	3.5	
IIB	144	34.7	26.4	20.1	15.3	3.5	
IIIA	166	37.4	31.3	13.3	13.9	4.2	
IIIC	58	29.3	25.9	25.9	15.5	3.5	
Tumor grade							<0.0001
Well-differentiated	159	80.5	12	2.5	1.3	3.8	
Moderately-diff	496	57.3	26.6	4.8	7.9	3.4	
Poorly diff	497	17.9	23.9	37.4	18.1	2.6	
Unspecified	101	62.4	13.9	10.9	7.9	5	
Mean age at diagnosis (SE)		52.8 (0.4)	50.8 (0.5)	48.2 (0.6)	50.5 (0.8)	49.6 (1.2)	<0.0001
Menopausal status at diagnosis							0.02
Pre-menopausal		40.3	25	19.9	10.9	3.9	
Post-menopausal		49.7	20.3	16.3	11.3	2.7	

ER+/Her2-
=
ER+ tumors

ER+/Her2+ split
across Her2-
enriched (34%)
Luminal A (29%)
Luminal B (31%)



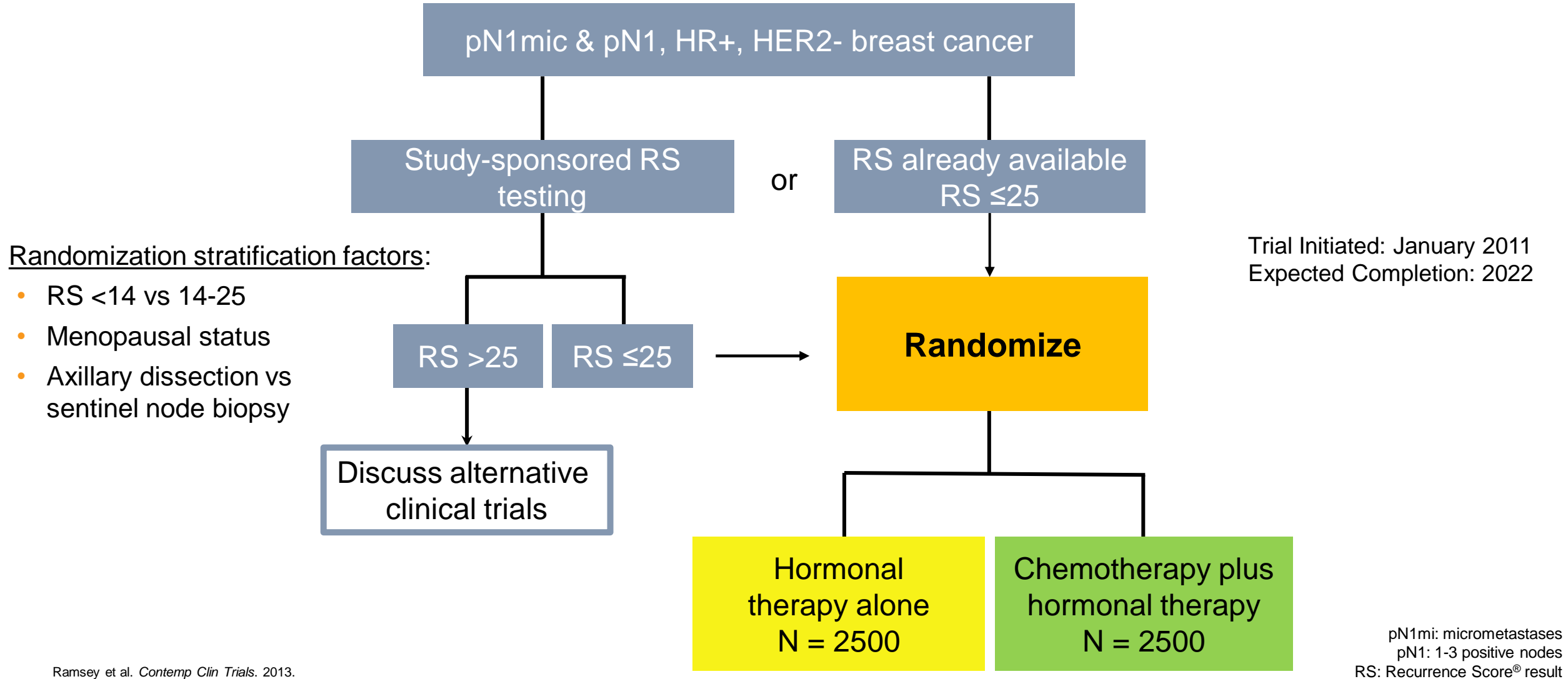
1253 pts

Prognostic value of PAM50 subtypes over clinical factors in an independent breast cancer cohort with long-term follow-up

Pam50 intrinsic subtype is independently prognostic for long-term breast cancer survival, irrespective of menopausal status

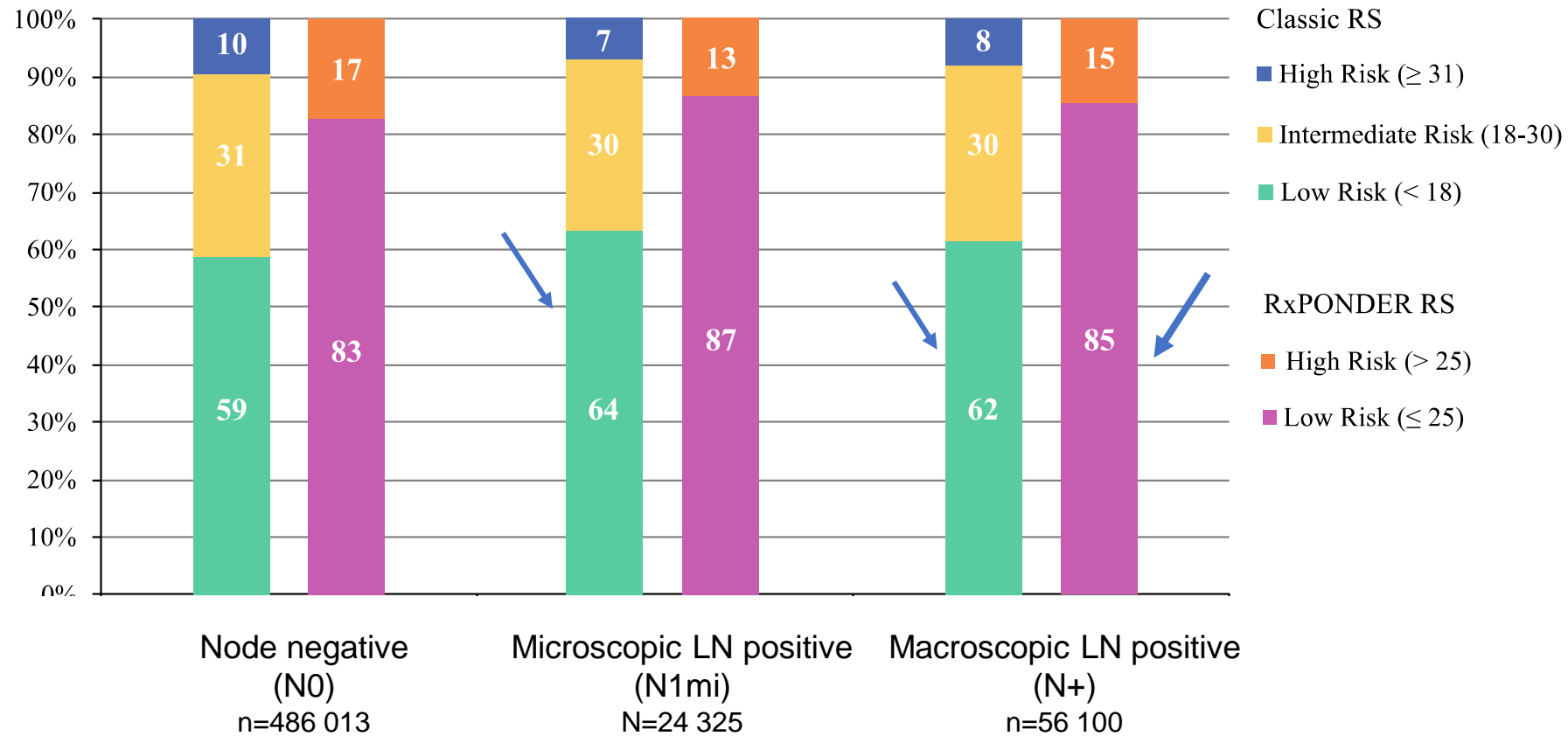
Pu et al Br Cancer Res Treat 2019

Node-Positive Disease: RxPONDER Trial Schema



Ramsey et al. *Contemp Clin Trials*. 2013.

Lymph Node Status Does Not Predict Tumor Biology (2004-2017), N=610.350



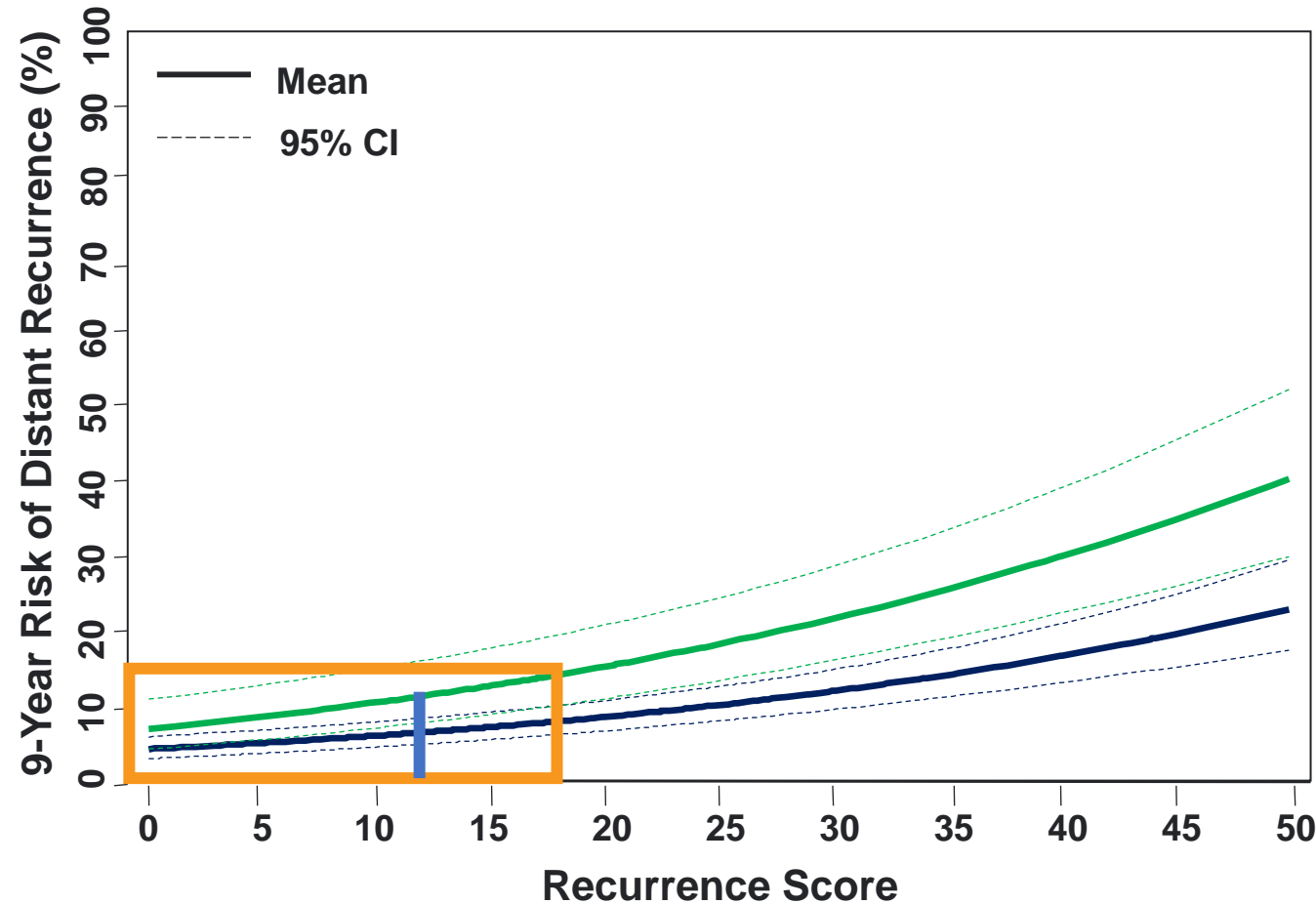
- With classic low risk cutoff RS 0-17, 64% N1mi and 62% of N1 patients can be spared chemotherapy
- If RxPONDER shows no chemotherapy benefit with RS ≤ 25 , 87% N1mi and 85% N1 patients can be spared chemotherapy

Node-Positive (N1mi/1-3 LN+)

Level IA Evidence	Study	Type of Study	N	Study Design	Endpoints
	transATAC	Prospective—retrospective; validation	306 (all N+)	ANA vs TAM vs ANA+TAM	9-year proportion DR-free
	SWOG 8814	Prospective—retrospective; validation	367 (all N+)	TAM vs CAF→T vs CAFT	10-year DFS in TAM-alone arm
	WSG PlanB	Prospective outcomes	930 (1-3 N+)	RS < 12: ET RS 12-25: ET vs CT RS ≥ 26: CT	5-year DFS 5-year DDFS
	Clalit	Prospective outcomes	709 (all N+)	Population-based registry	5-year DR 5-year BCSD
	SEER	Prospective outcomes	6,483 (N1mi/1-3 N+)	Population-based registry	5-year BCSM

Good Outcomes in Patients With Low-Risk RS Results Without Chemotherapy

TransATAC



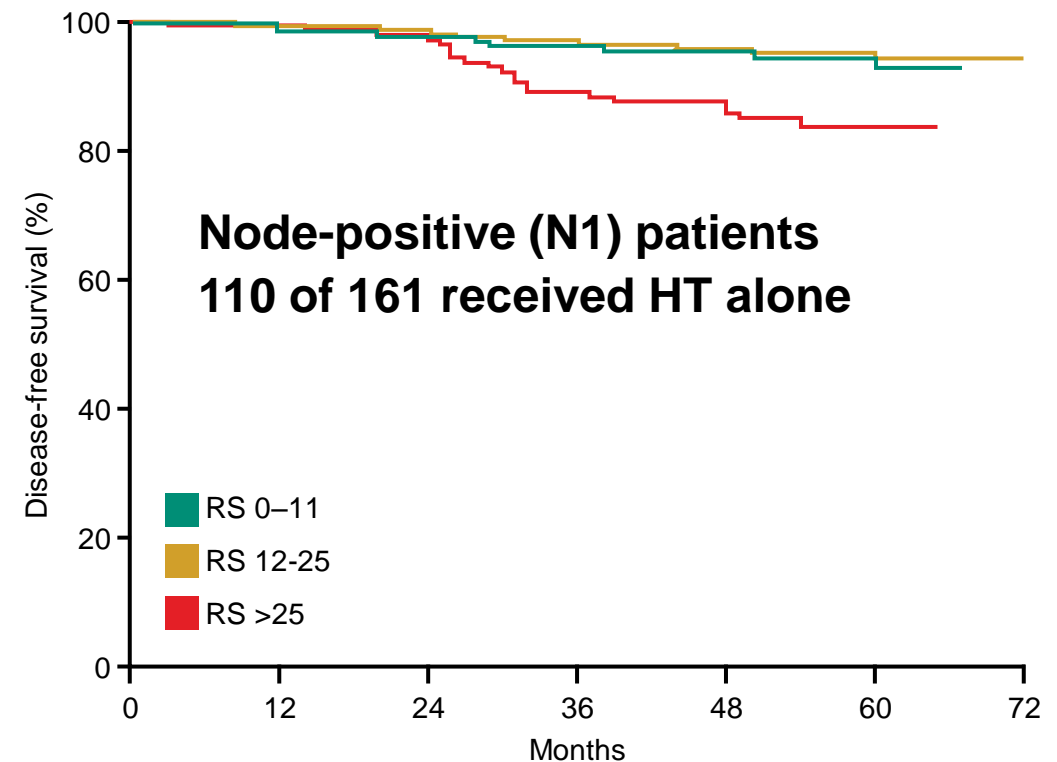
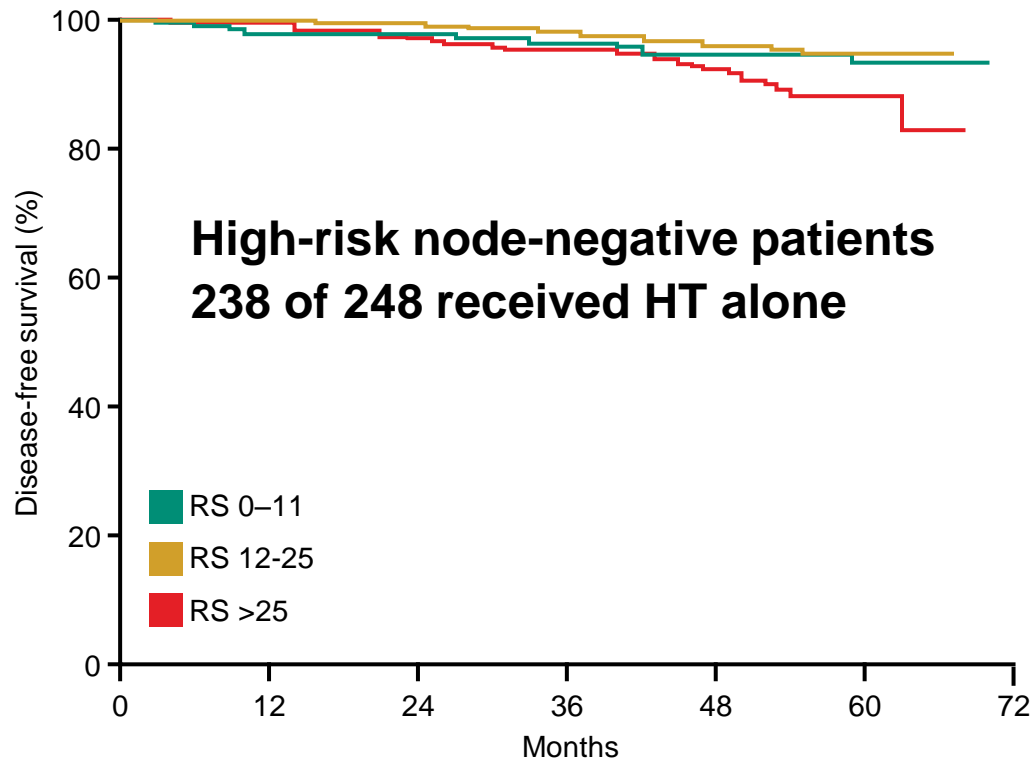
A low Recurrence Score result (<18) indicates a low risk of recurrence for patients with 1-3 positive nodes

**1-3 Positive nodes
n = 243**

**Node-negative
n = 872**

RS Result Risk-Stratifies Node-Positive Patients Using Hormone Therapy Alone

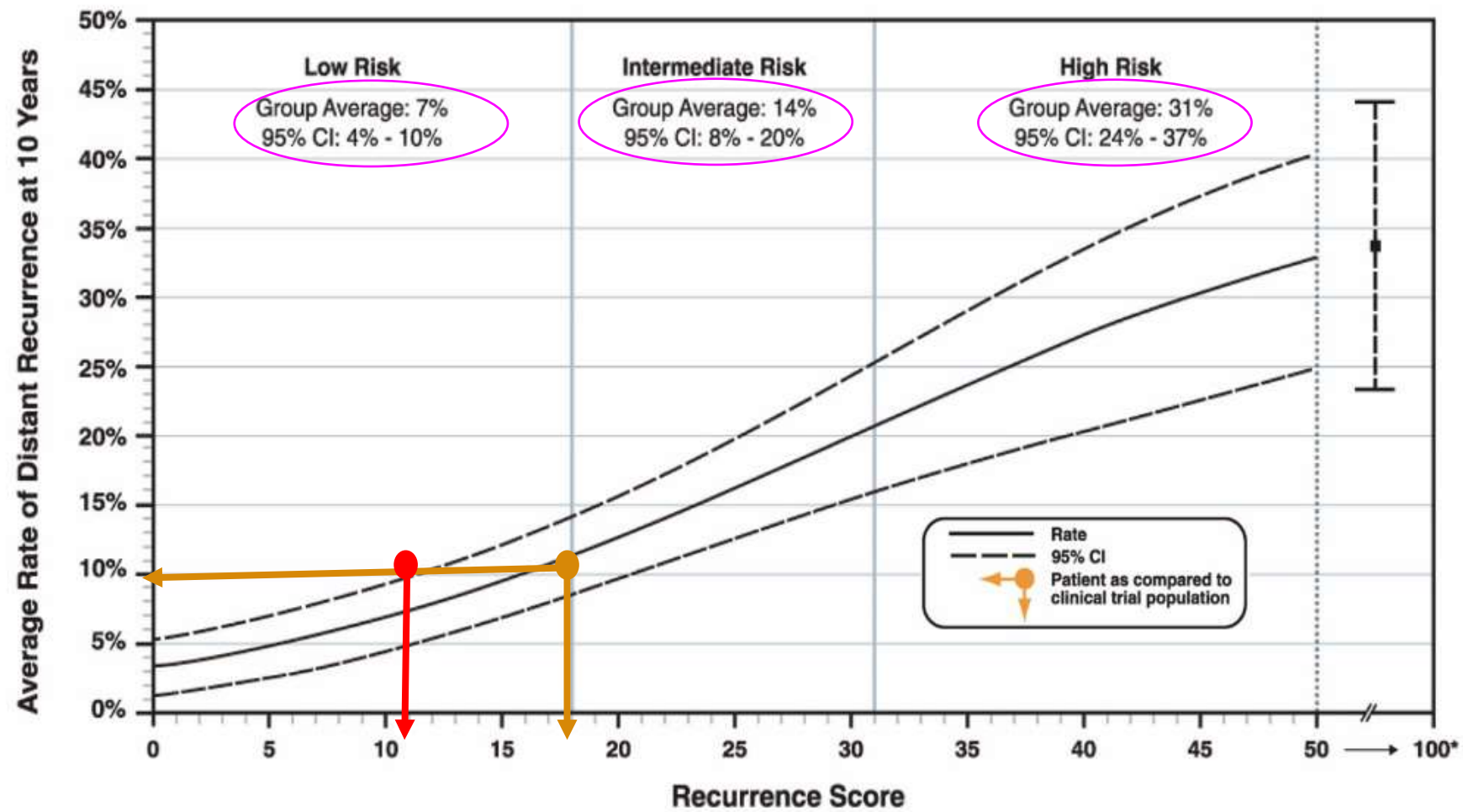
West German Study Group PlanB Trial: High-Risk N0 and N1 Patients With RS 0-11 Do Equally Well With ET Alone



5-year disease-free survival was 94% in high-risk N0 and N1 patients with Recurrence Score results 0-11 and treated with hormone therapy alone

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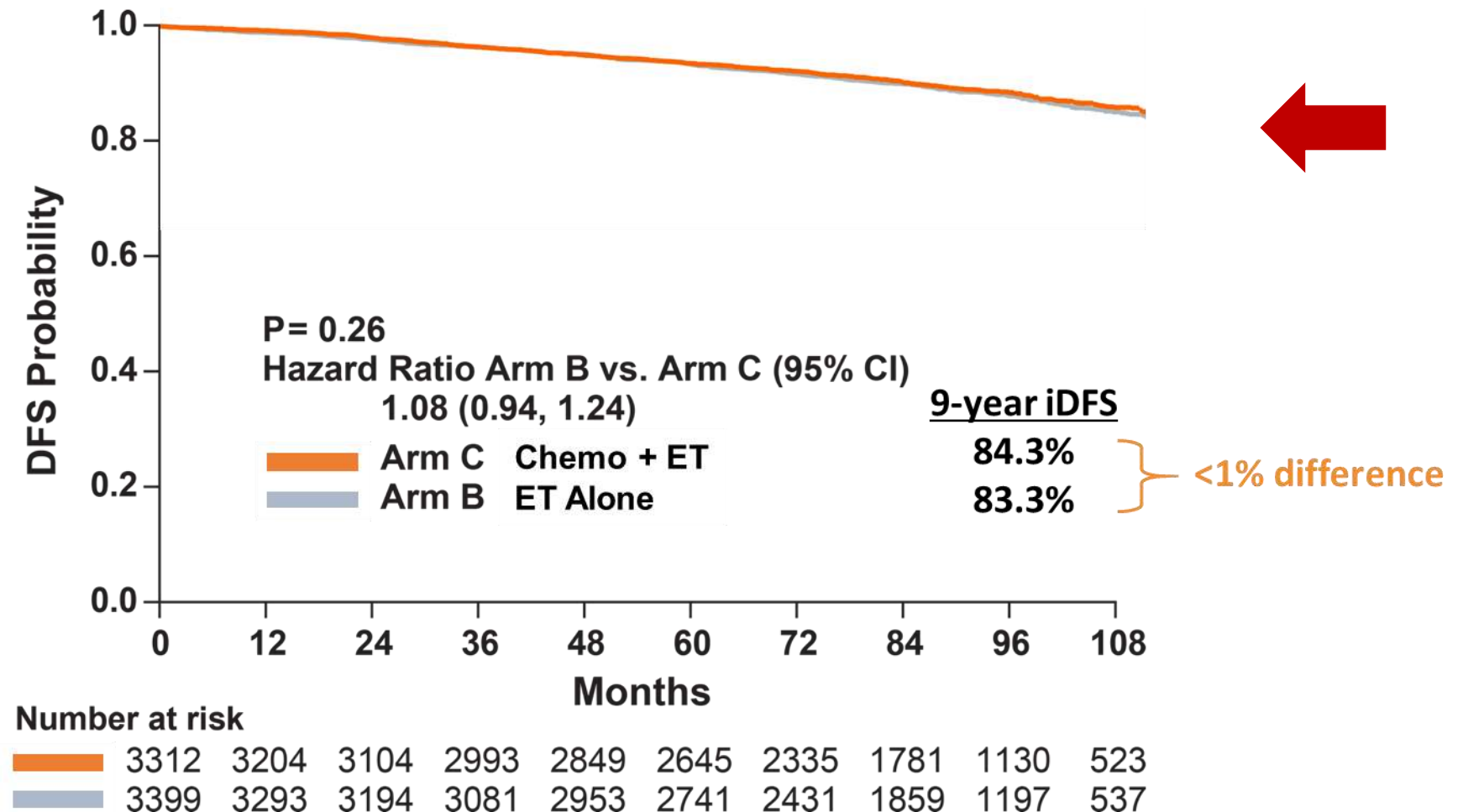


*For Recurrence Scores >50, group average rate of distant recurrence and 95% CI shown

TAILORx

ET Alone Was Not Inferior to CT/ET in RS 11-25

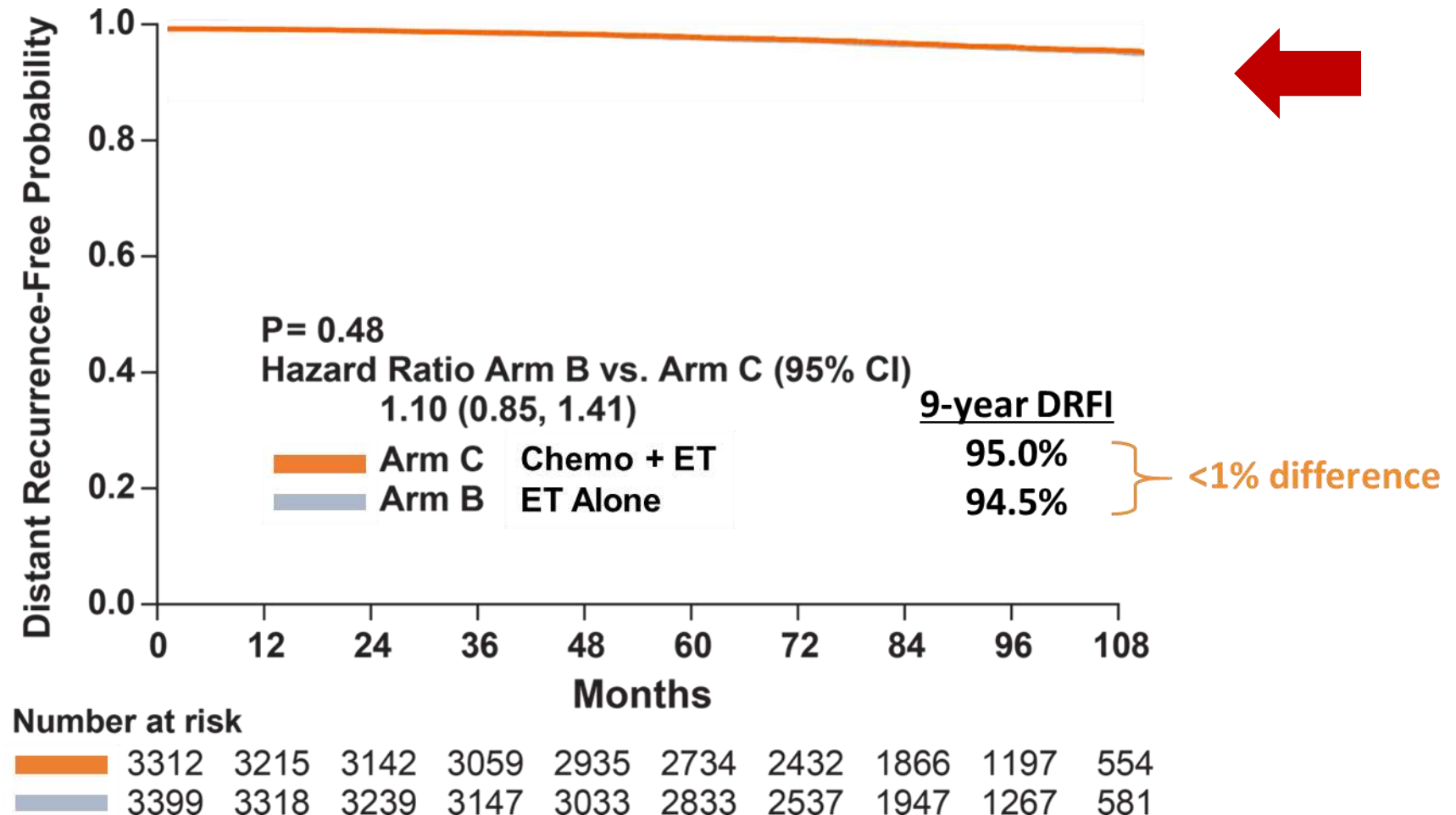
836 iDFS events after median follow-up of 7.5 years



TAILORx

Very Low Risk of Distant Recurrence in RS 11-25

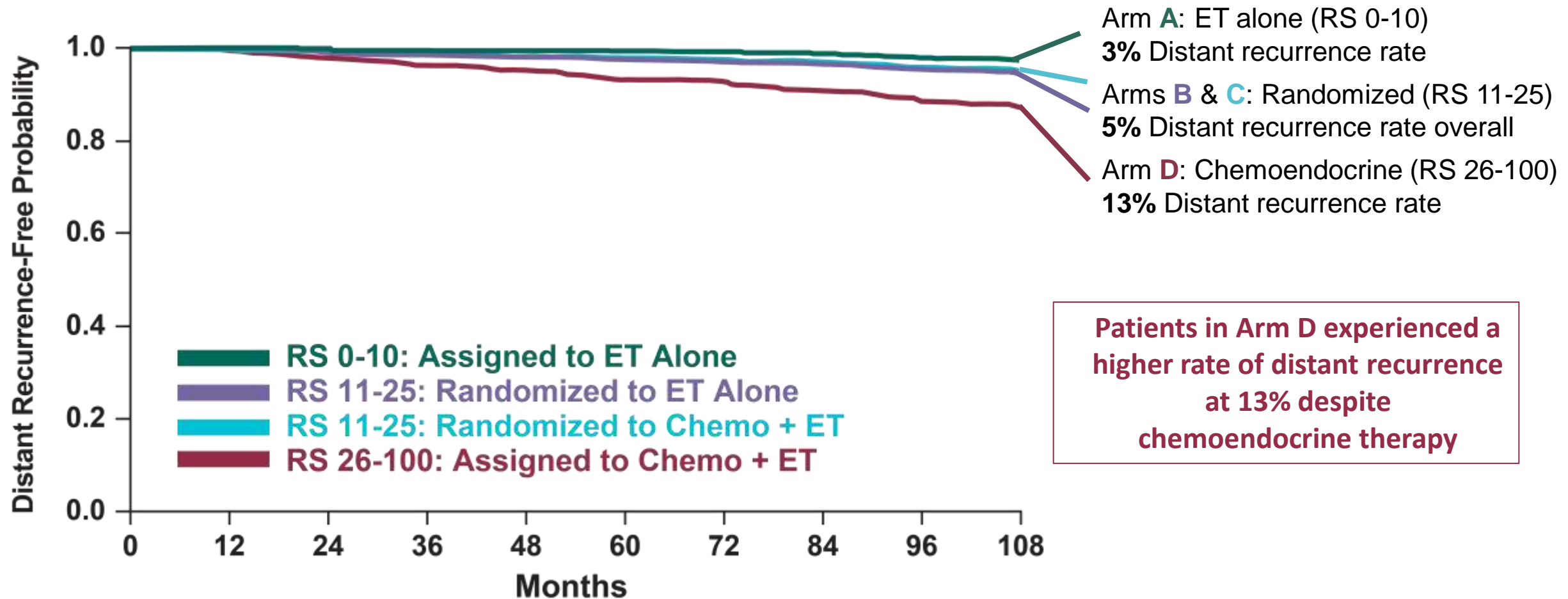
199 of 836 (23.8%)
were distant
recurrences



TAILORx

Arms A, B & C With RS 0-25 Have $\leq 5\%$ Risk of Distant Recurrence at 9 Years

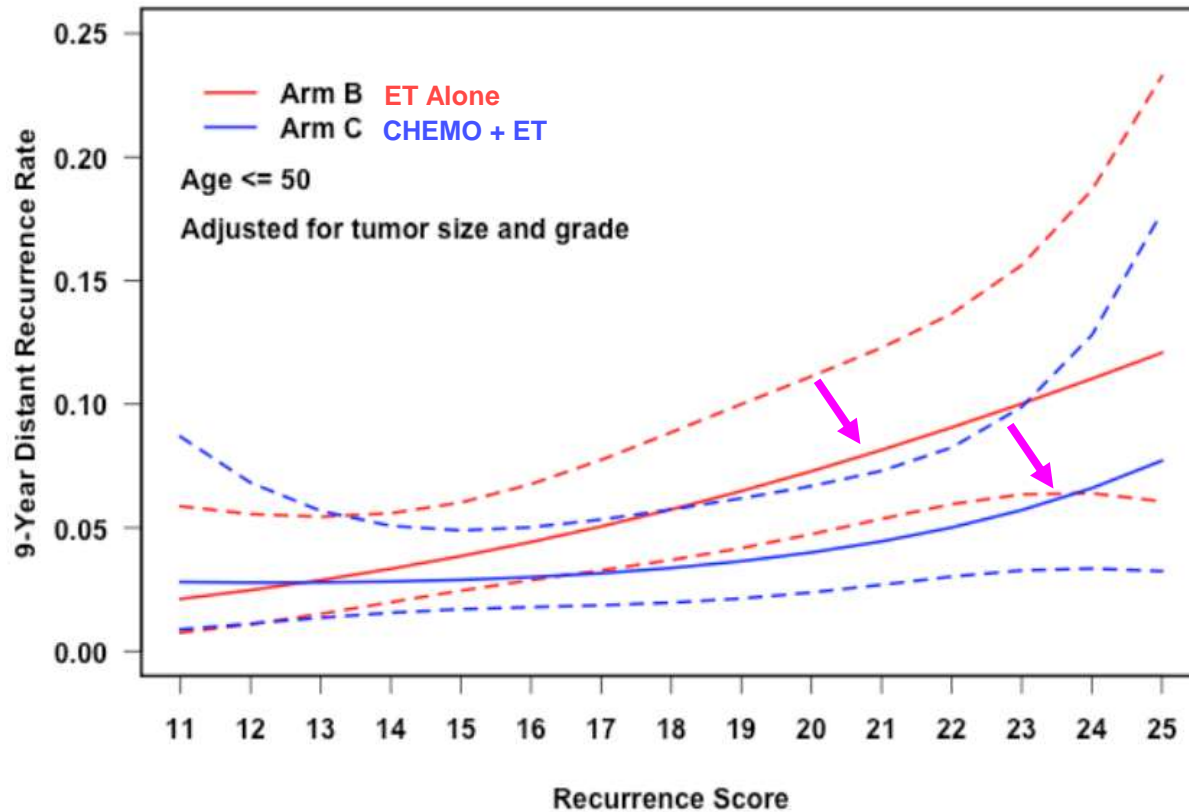
9-Year Event Rates – ITT Population: All Arms



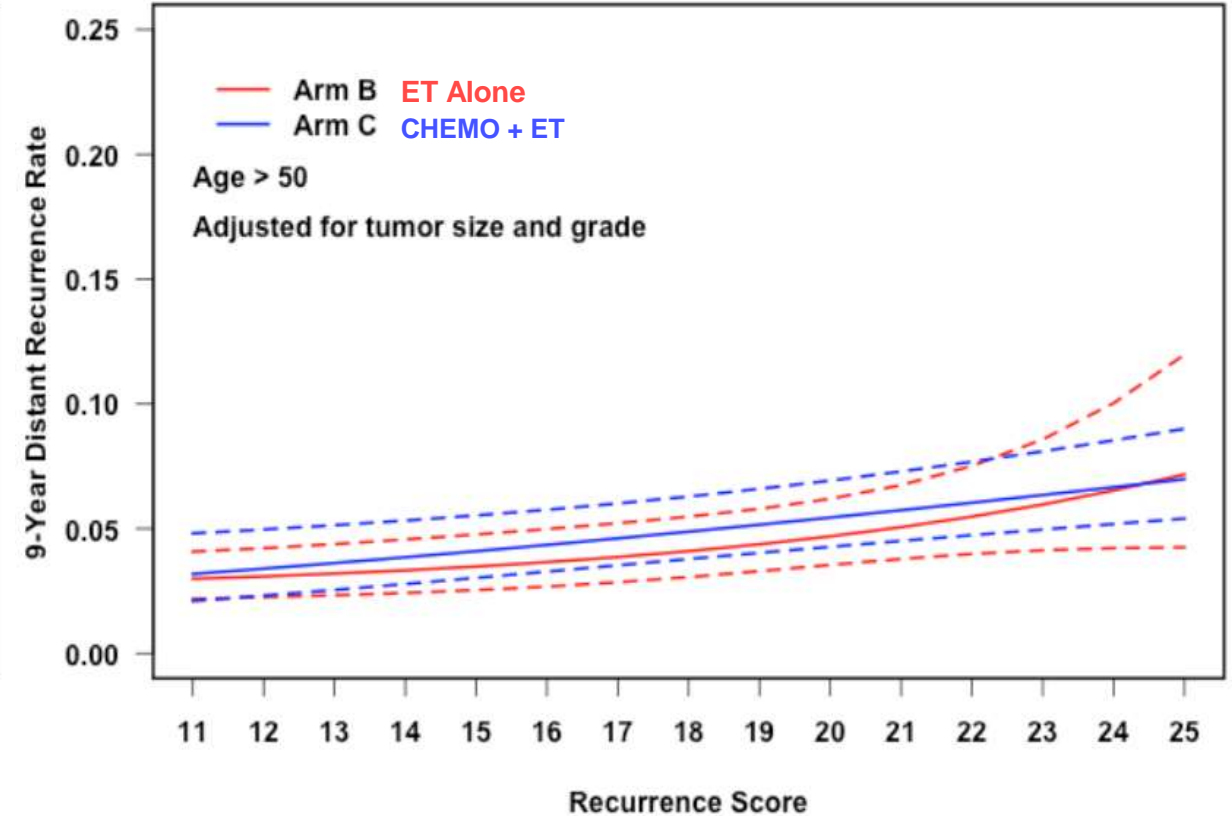
Sparano et al. *N Engl J Med*. 2018.

TAILORx and the Age

≤50 Years (n=2216)



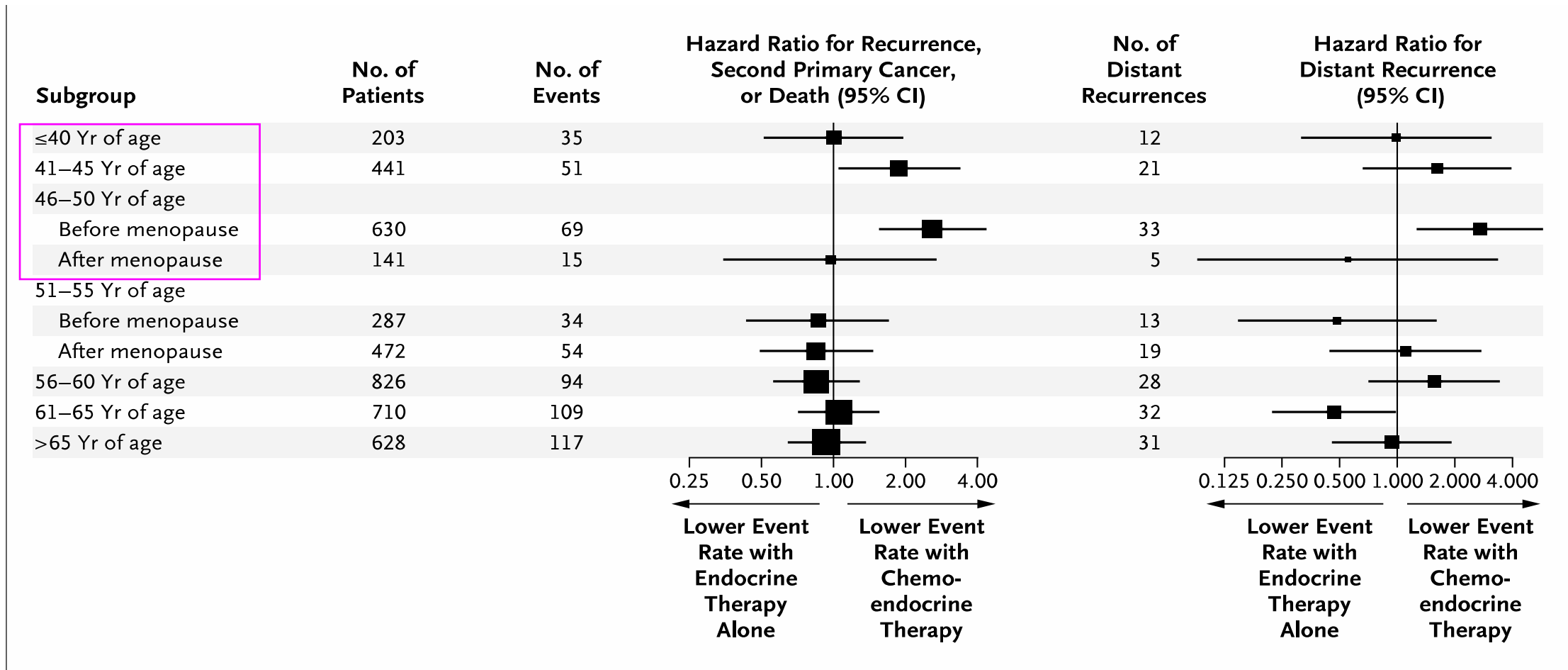
>50 Years (n=4495)



The magnitude of chemotherapy benefit in patients ≤50 years increases with increasing Recurrence Score result, but was not statistically significant

Effect of Age and Menopausal Status on Chemotherapy Benefit

RS 16-25



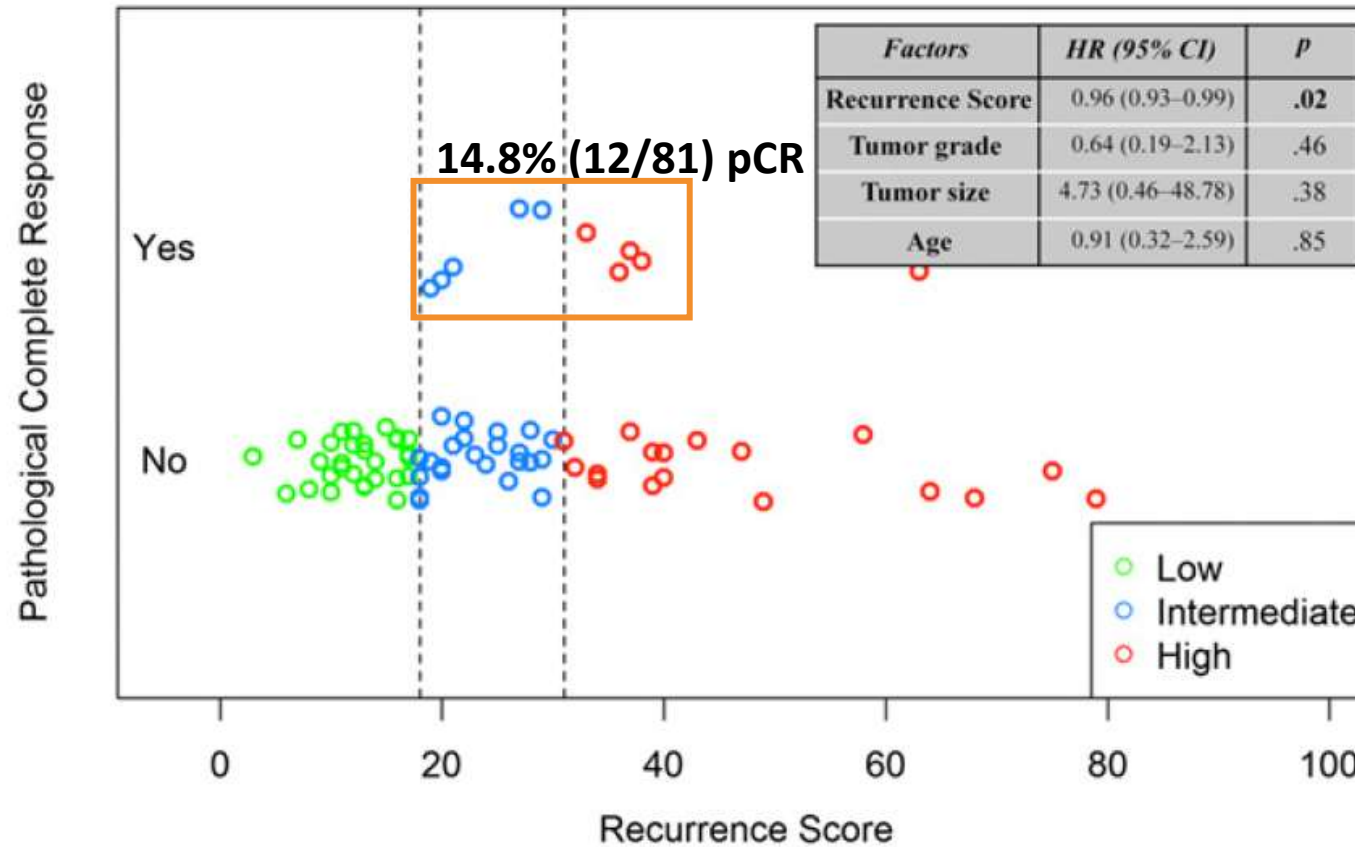
Sparano et al, NEJM 2019

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What about prognostic value of Genomic Testing in Neoadjuvant Setting?

pCR and RS



RS was the only significant predictor of pCR

Neoadjuvant Studies Supporting Chemotherapy Benefit with RS Group 26-100

Neoadjuvant Chemotherapy

Study	Type of Study	N	pCR Rate	
			RS 0-25	RS 26-100
Gianni et al.	Neoadjuvant CT	89	0%	12%
Zelnak et al.	NACT vs NAHT	46	0%	22%
Yardley et al.	Neoadjuvant CT	108	0%	26%
Bear et al.	NACT vs NAHT	64	0%	14%

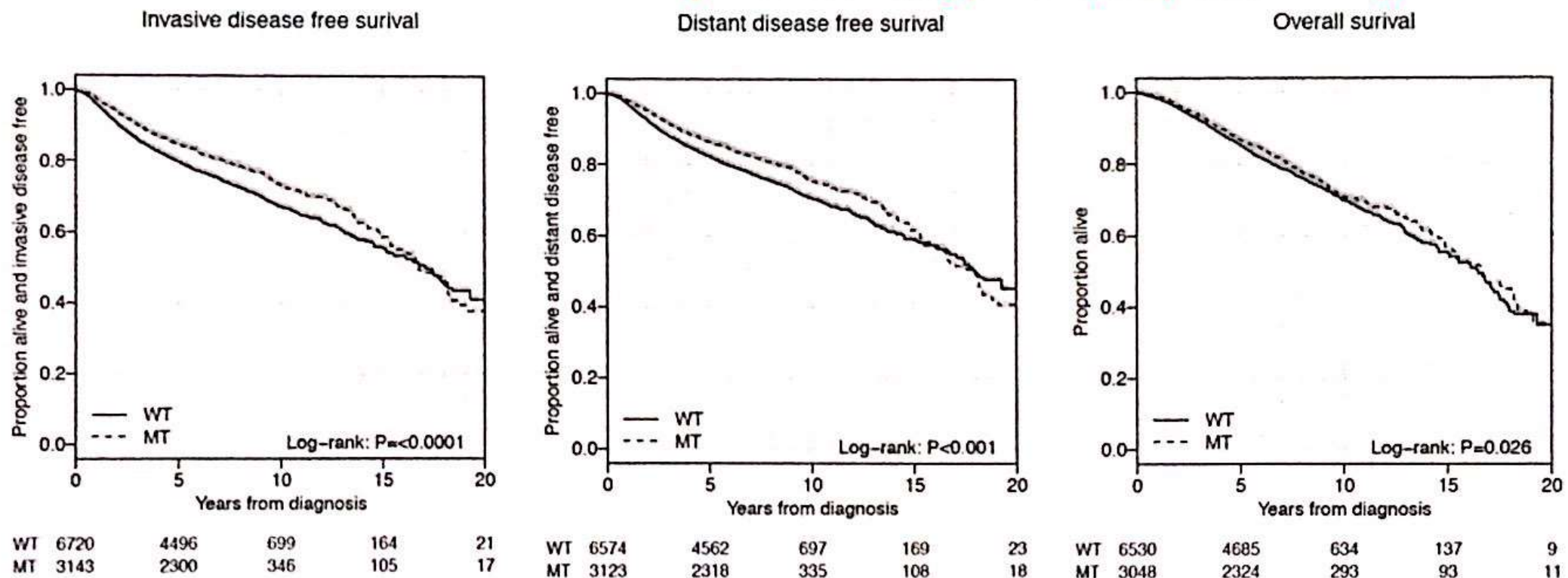
Sparano et al. *J Clin Oncol.* 2008; Gianni L, et al. *J Clin Oncol.* 2005; Chang JC, et al. *Breast Cancer Res Treat.* 2008; Zelnak AB, et al. *J Clin Oncol.* 2013; Yardley DA, et al. *Breast Cancer Res Treat.* 2015; Bear HD, et al. *J Surg Oncol.* 2017

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PIK3CA mt BCs & prognosis in early stage disease n=10,319

PIK3CA mutants had better outcomes than WT HR = 0.77 [95%CI: 0.71 – 0.84], $p < 0.001$, independent of subtype and location



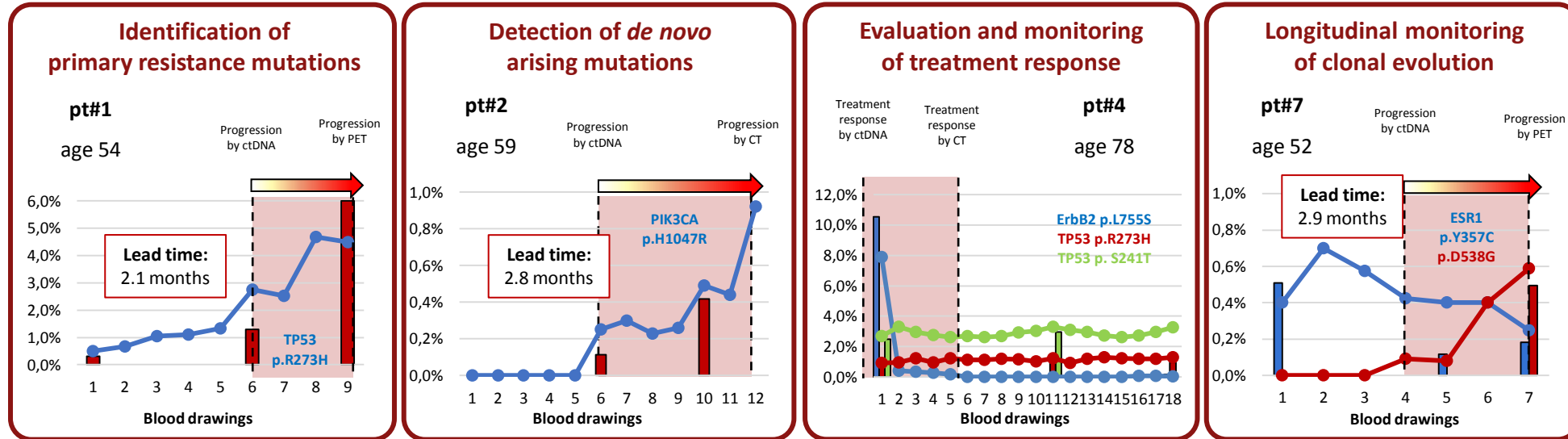
Pooled analysis of 10,319 patients from 19 studies, Median OS FU 7yrs .

Zardavas et al, submitted

LiquERB: the GIM21 project when prediction become prognosis

LiqBreastTrack trial: preliminary results (II)

VAF (blood)



ctDNA present at time 0,
and slowly going up

ctDNA **NOT** present at time 0,
but de novo appearing some time
after the beginning of treatment

ultra-fast clearance

intersecting ctDNA
trajectories

resistance (primary)

resistance (acquired/adaptive)

*sensitivity
(best responders)*

*sensitivity &
resistance
(bi-clonal ear-
marking)*



NGS



dPCR

Fabi, oral communication

THANK YOU for Your Attention

