

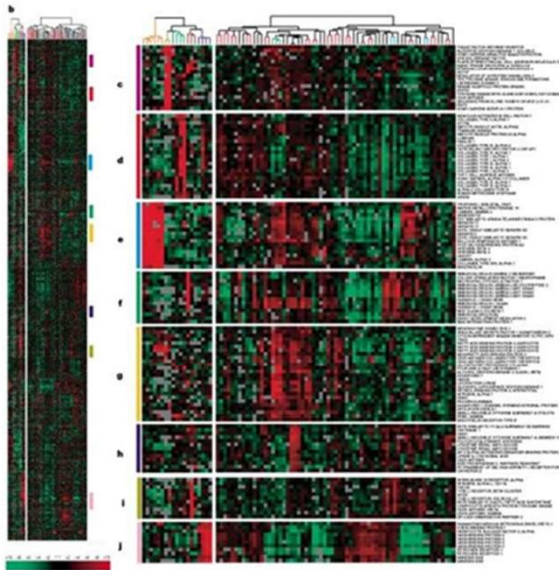


Predictive and Prognostic Biomarkers Using Genomic Profile in Early Breast Cancer

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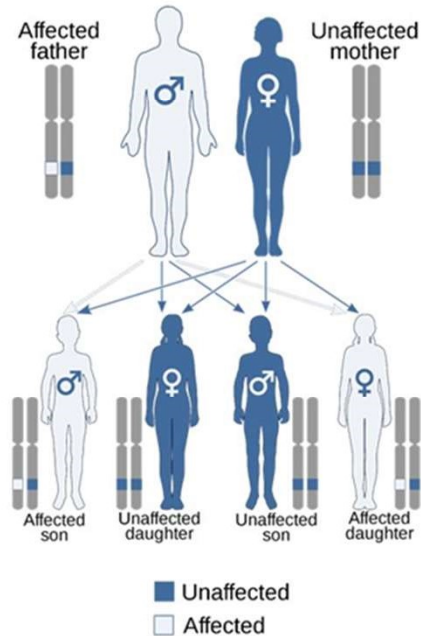


TUMOR TRANSCRIPTOME



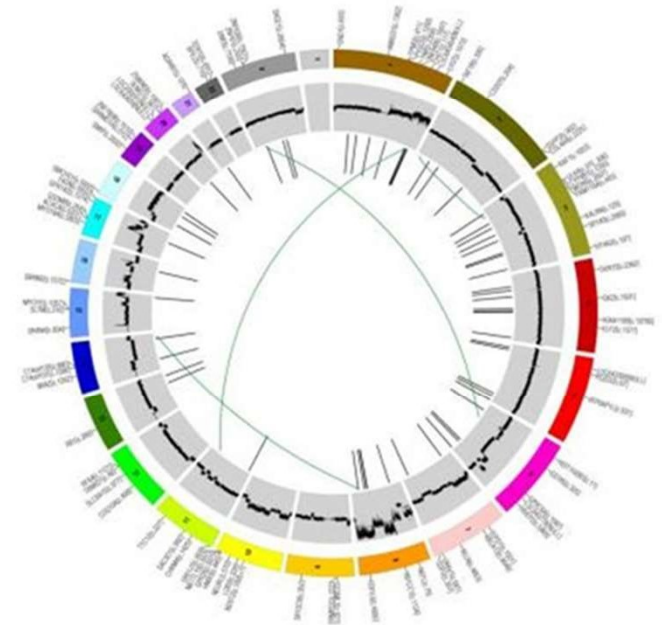
Nucleic Acid: Somatic tumor RNA
 Source: Tumor biopsy
 Timing: baseline diagnosis
 Readout: patterns of gene expression
 # genes: 6 to ~ 100

GERMLINE GENETICS



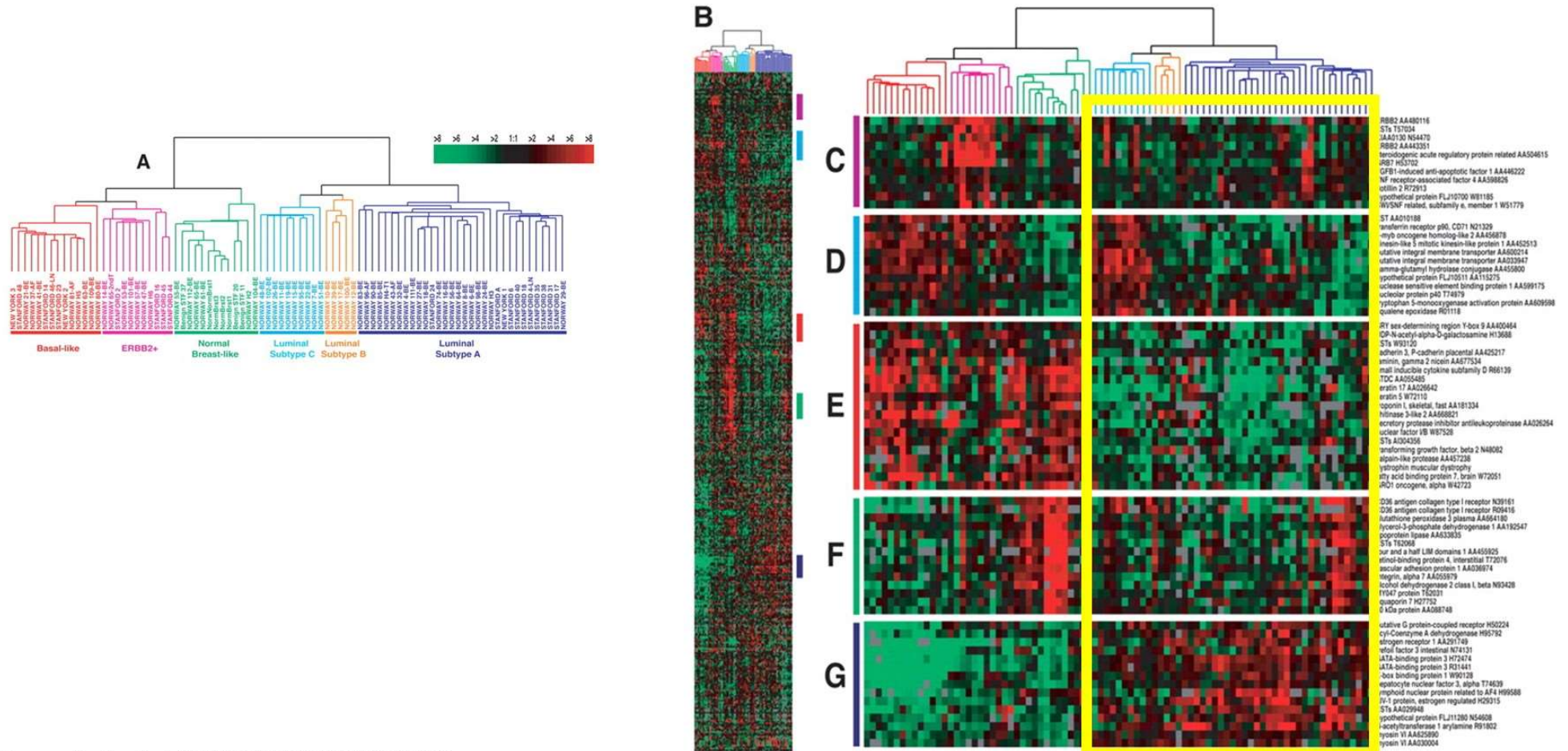
Nucleic Acid: Germline DNA
 Source: "normal" tissue (e.g. blood)
 Timing: At diagnosis, when mutation suggested by FH, or at stage IV
 Readout: mutations in genes predisposing to cancer
 # genes: Variable from 2 (BRCA1/2) to > 25 depending on possible syndromes

TUMOR GENOME

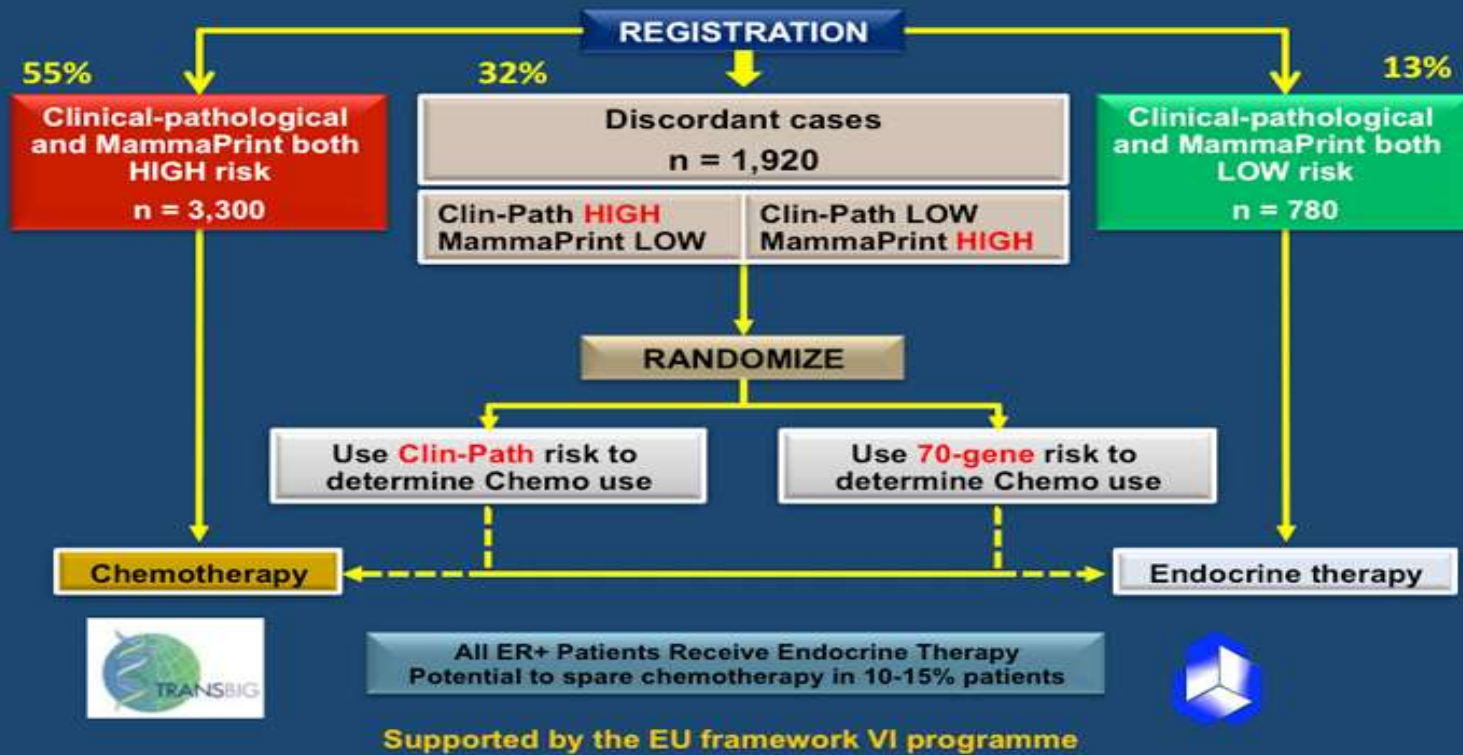


Nucleic Acid: Somatic tumor DNA
 Source: Tumor biopsy or cell-free DNA
 Timing: Metastatic recurrence & serially
 Readout: somatic changes in tumor DNA at baseline and over time
 # genes: ~ 500

Gene expression patterns of 85 experimental samples analyzed by hierarchical clustering using the 476 cDNA intrinsic clone set.



MINDACT Trial Design (n = 6,000) Node negative & 1-3 positive nodes



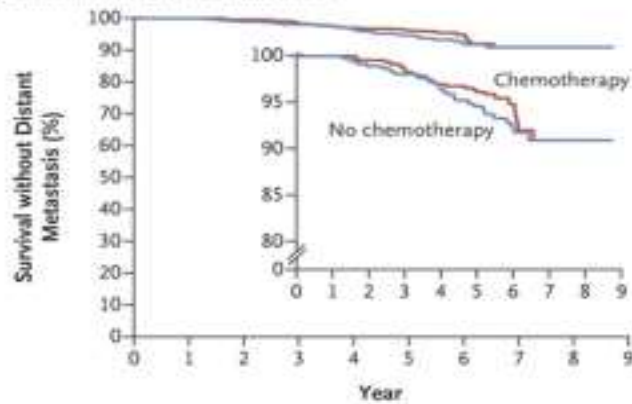
MINDACT RESULTS Cardoso et al. NEJM 2016

Clinical Genomic	Low Low	Low High	High Low	High High
No.	2745	592	1550	1806
N+	6%	3%	48%	26%
T < 2 cm	96%	98%	42%	52%
Grade 3	2%	15%	29%	76%
ER+ Luminal	96%	79%	90%	50%
TNBC	0%	9%	1%	31%
HER2+	4%	12%	8%	19%
5 year DDFS	97.6%	94.8%	95.1%	90.6%
Δ DFS with chemo		2.2% HR .74, p NS	3.0% HR .64, p 0.026	



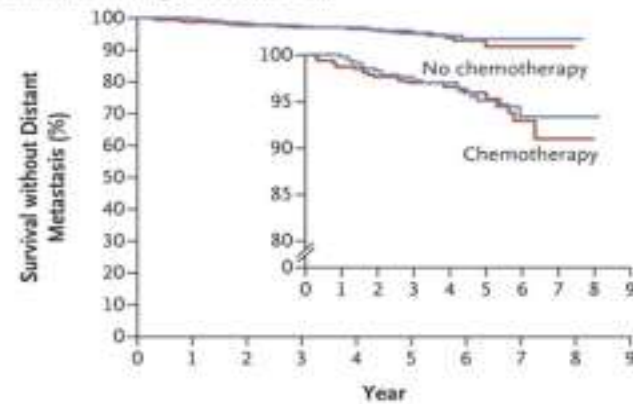
MINDACT: Survival without Distant Metastasis, Disease-free Survival, and Overall Survival in the Two Discordant-Risk Groups, According to Randomized Treatment.

A High Clinical Risk, Low Genomic Risk



No. at risk		0	1	2	3	4	5	6	7	8	9
Chemotherapy	749	714	698	677	611	346	145	41	3		
No chemotherapy	748	727	708	696	655	424	160	41	4		

B Low Clinical Risk, High Genomic Risk



No. at risk		0	1	2	3	4	5	6	7	8	9
Chemotherapy	344	321	316	306	281	179	81	22	0		
No chemotherapy	346	336	327	319	291	178	82	24	3		

Cardoso F et al.
N Engl J Med 2016;375:717-729.



Clinical Risk

Low Clinical Risk:

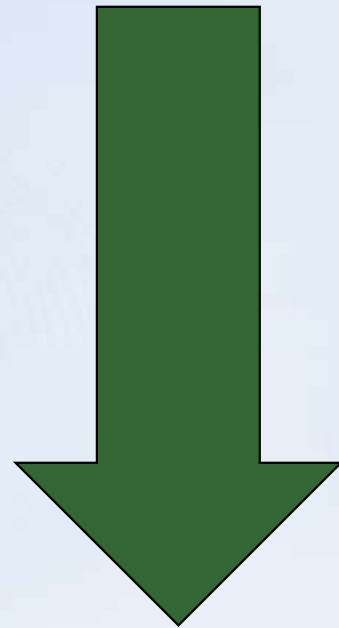
- Tumor <3cm + Low Grade
- Tumor <2cm + Int. Grade
- Tumor <1cm + High Grade

High Clinical Risk:

Those that do not meet
Low risk criteria



IMPACT OF 70 GENE SIGNATURE IN HIGH CLINICAL RISK GROUP

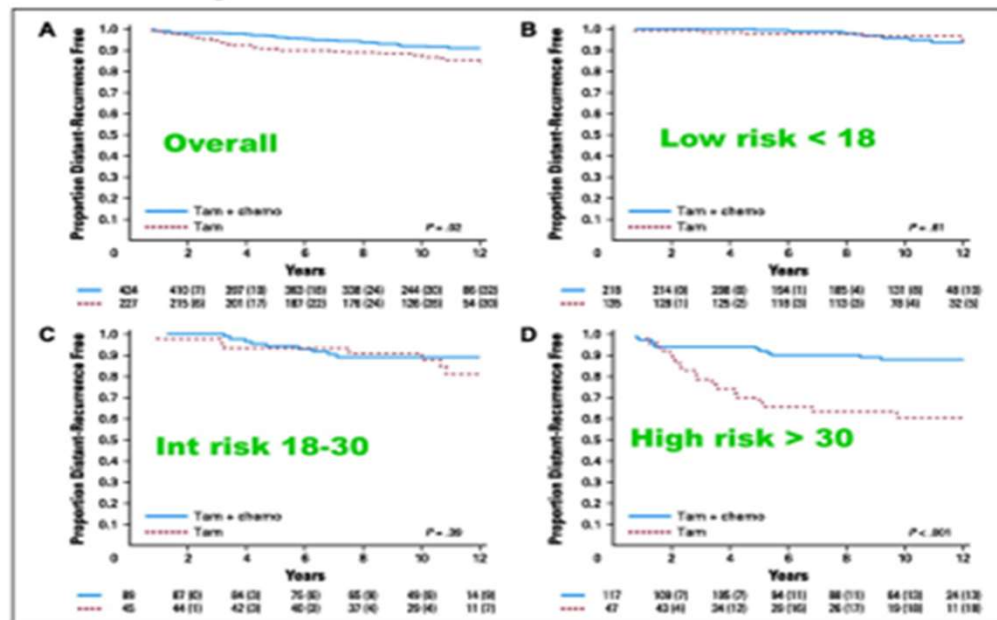
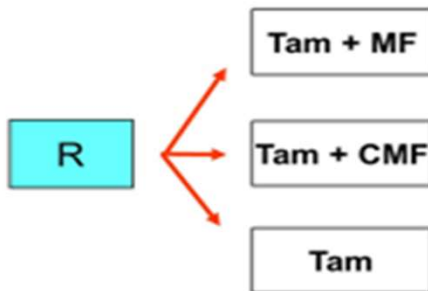


46%

Chemotherapy use



NSABP B-20 Outcome by Recurrence Score



Paik S, et al. *J Clin Oncol*. 2006;24(23):3726-3734.



TAILORx: Treatment Assignment & Randomization

Accrued Between April 2006 - October 2010

Key Eligibility Criteria

- Node-negative
- ER-pos, HER2-neg
- T1c-T2 (high-risk T1b)

Preregister - Oncotype DX RS (N=11,232)

Register (N=10,273)

Statistical Design

- Non-inferiority - IDFS
- HR 1.332 (90 vs. 87% 5-yr DFS)
- Type I 10%, type II 5%
- Full info- 835 IDFS events

ARM A: Low RS 0-10
(N=1629 evaluable)

ASSIGN

Endocrine Therapy (ET)

Mid-Range RS 11-25

(N=6711 evaluable)

RANDOMIZE

Stratification Factors: Menopausal Status, Planned Chemotherapy, Planned Radiation, and RS 11-15, 16-20, 21-25

ARM D: High RS 26-100
(N=1389 evaluable)

ASSIGN

ET + Chemo

ARM B: Experimental Arm

(N=3399)

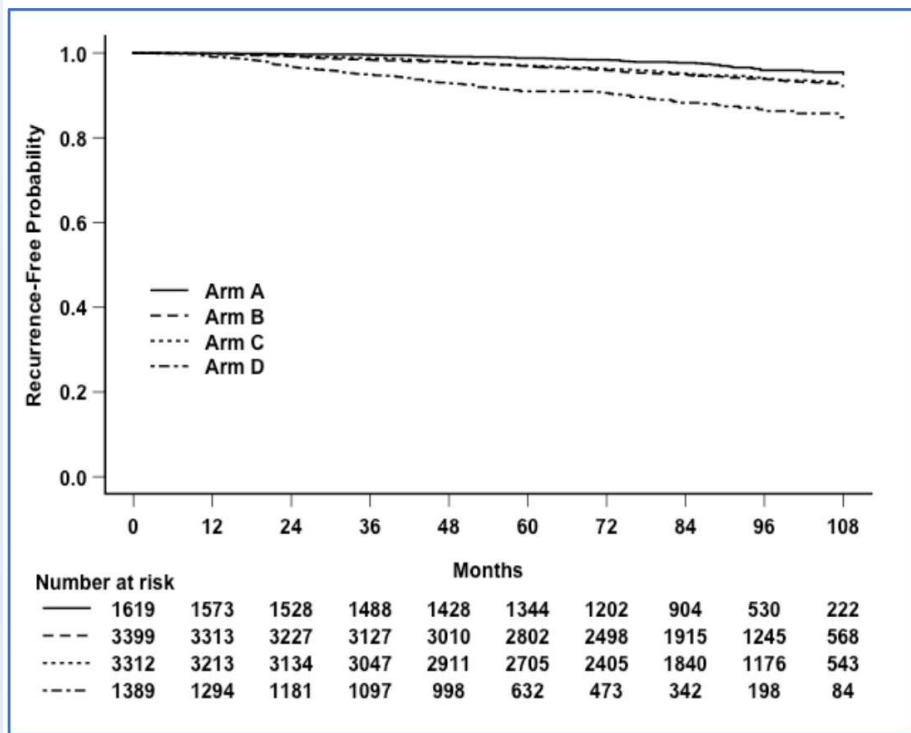
ET Alone

ARM C: Standard Arm

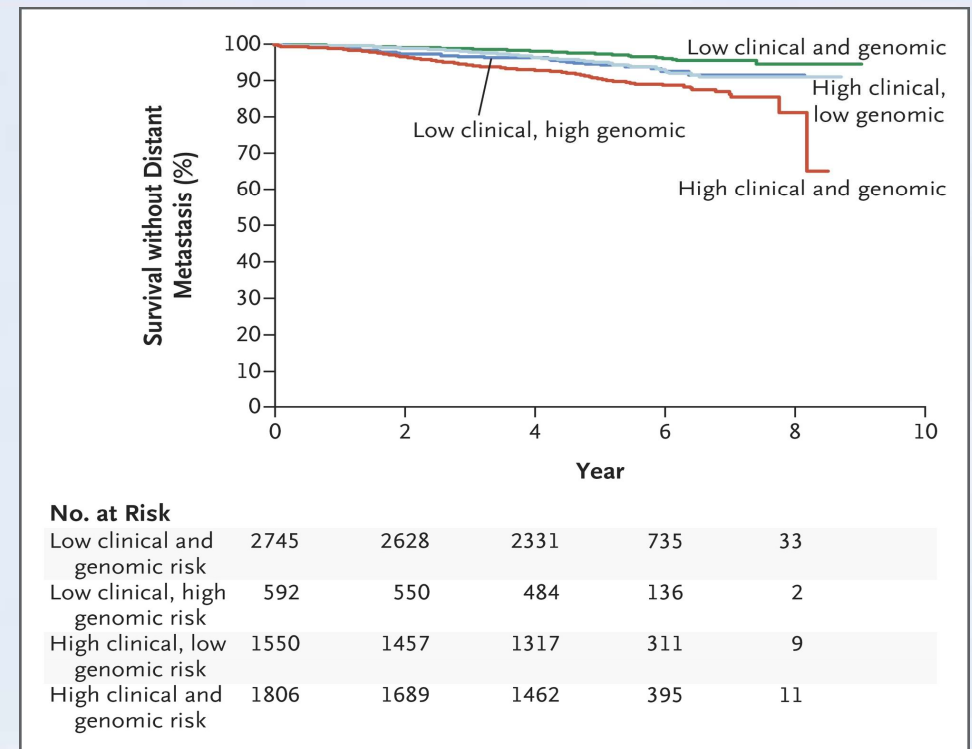
(N=3312)

ET + Chemo

PROSPECTIVE STUDIES IN ENDOCRINE THERAPY ALONE IN LOW GENOMIC RISK, EARLY BREAST CANCER



TAILORx Sparano J, et al. NEJM 2018; 379:111-121



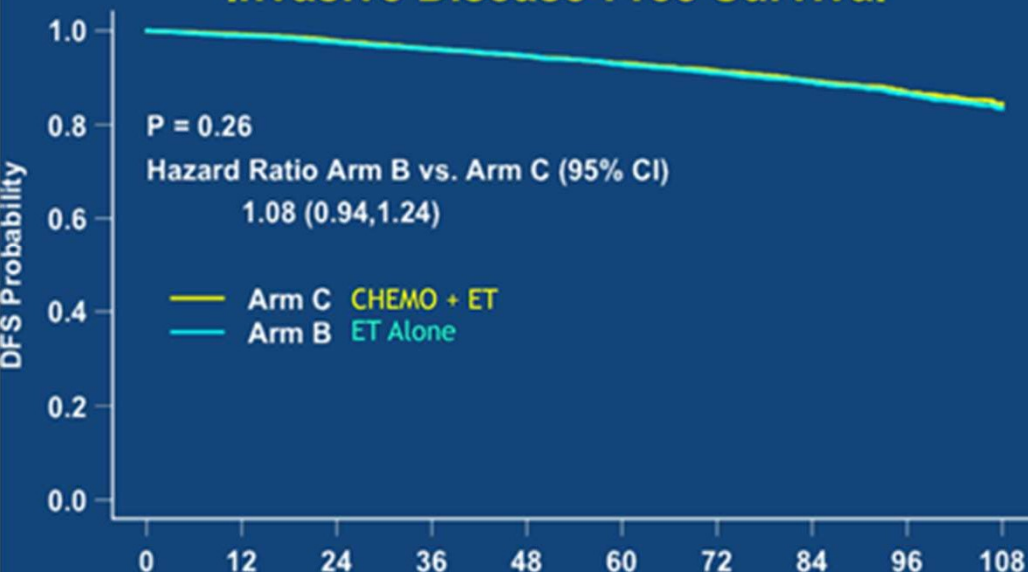
MINDACT Cardoso F, et al. NEJM 2016; 375: 717-729



TAILORx Results - ITT Population: RS 11-25 (Arms B & C)

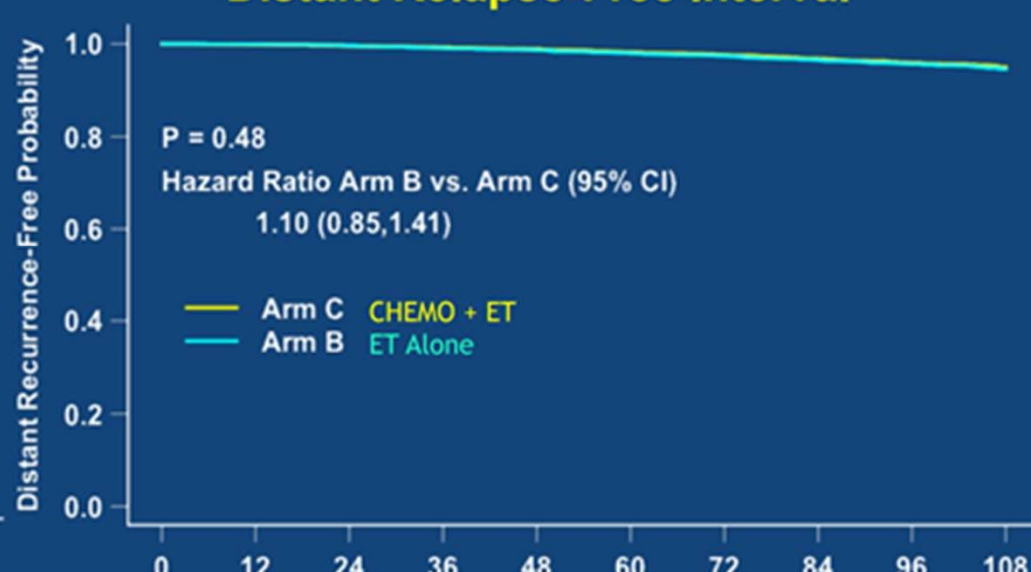
836 IDFS events (after median of 7.5 years), including 338 (40.3%) with recurrence as first event, of which 199 (23.8%) were distant

Primary Endpoint Invasive Disease-Free Survival



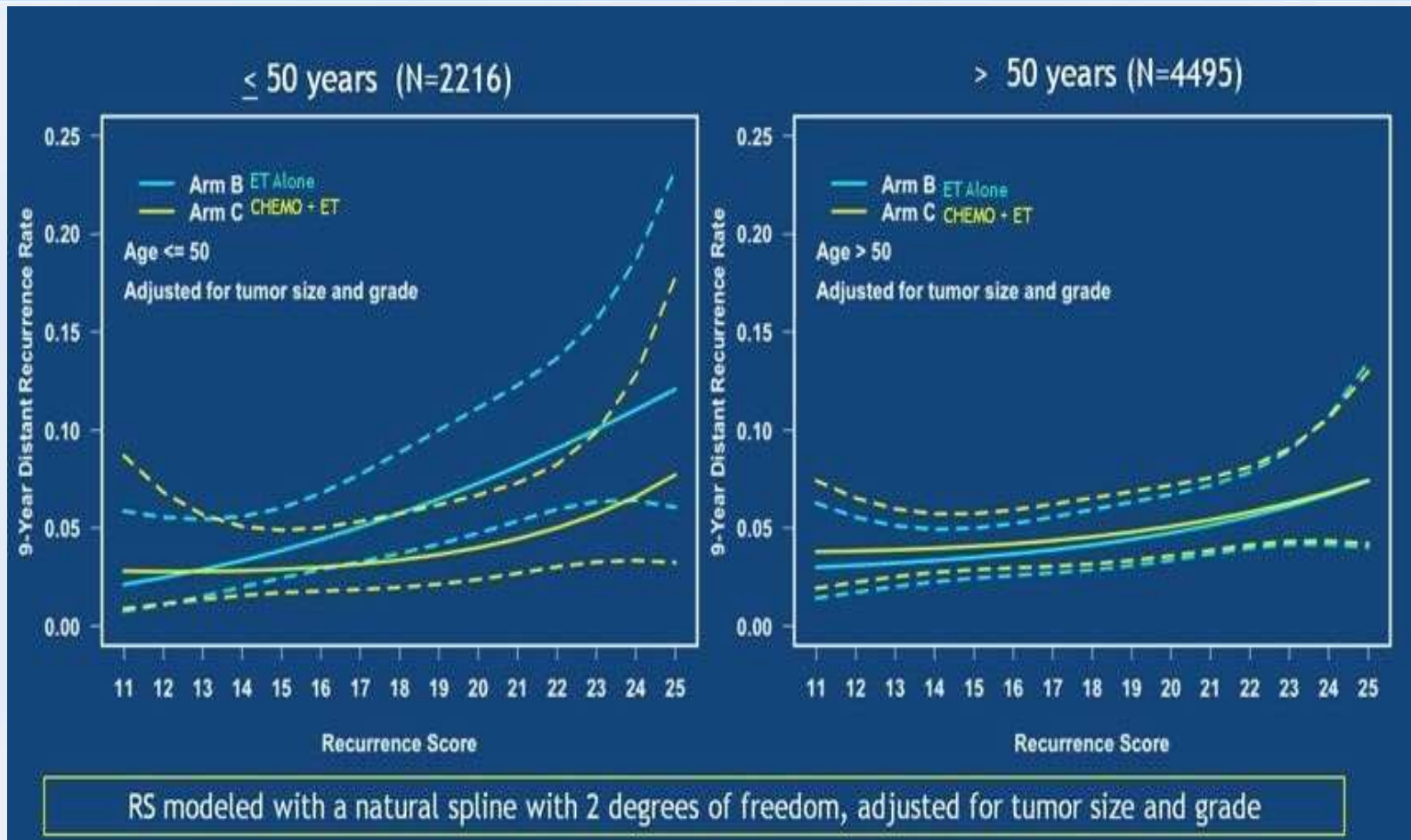
Number at risk		Months									
		0	12	24	36	48	60	72	84	96	108
—	Arm C CHEMO + ET	3312	3204	3104	2993	2849	2645	2335	1781	1130	523
—	Arm B ET Alone	3399	3293	3194	3081	2953	2741	2431	1859	1197	537

Secondary Endpoint Distant Relapse-Free Interval



Number at risk		Months									
		0	12	24	36	48	60	72	84	96	108
—	Arm C CHEMO + ET	3312	3215	3142	3059	2935	2734	2432	1866	1197	554
—	Arm B ET Alone	3399	3318	3239	3147	3033	2833	2537	1947	1267	581

TAILORx Results: Association between continuous RS 11-25 and 9 year distant recurrent rate stratified by age



TAILORx: Impact of Clinical Risk (CR) on Prognosis by RS Group (N=9427) 30% clinical high risk & 70% clinical low risk

Grouped by RS

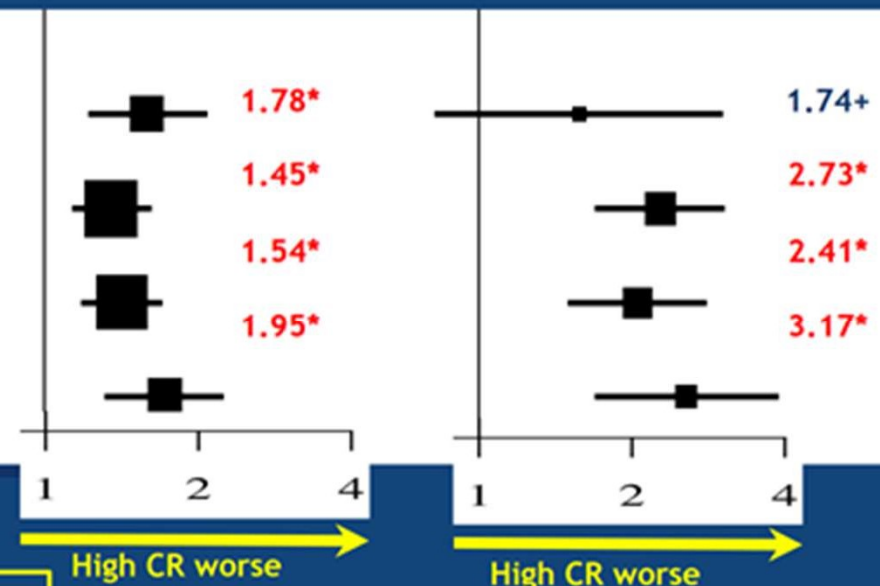
Total #/IDFS/DR events

IDFS Hazard Ratio

DRFI Hazard Ratio

All Patients (n=9427)

RS ≤ 10: Endocrine Therapy	1572/176/ 30
RS 11–25: Endocrine Therapy	3282/422/127
RS 11–25: Chemoendocrine Therapy	3214/389/113
RS > 25: Chemoendocrine Therapy	1359/184/ 96



Multivariate model for distant recurrence in RS 11-25 group:

(N=6496 cases and 240 distant recurrences):

- Clinical risk: HR for high vs. low risk 2.42, $p < 0.001$
- Continuous RS: HR 1.08, $p < 0.001$ (HR for a 1 point higher RS)

Hazard ratio > 1 - high clinical risk worse

+95% CI overlap 1

*95% CI don't overlap 1

Sparano et al. N Engl J Med 2019;380(25):2395-2405.

Clinical Risk

Low Clinical Risk:

- Tumor <3cm + Low Grade
- Tumor <2cm + Int. Grade
- Tumor <1cm + High Grade

High Clinical Risk:

Those that do not meet
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TAILORx: Impact of Clinical Risk (CR) on Prognosis by RS Group (N=9427) 30% clinical high risk & 70% clinical low risk

Grouped by RS

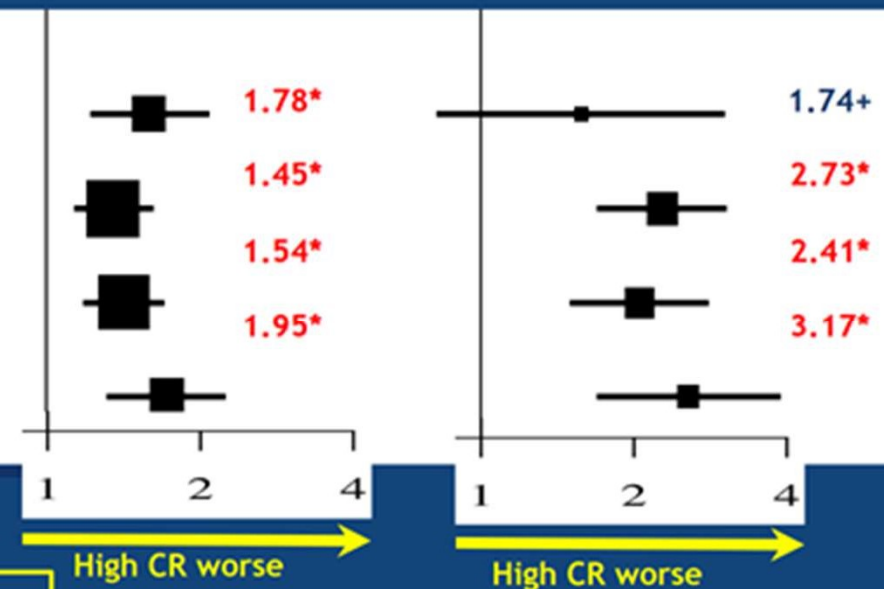
Total #/IDFS/DR events

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DRFI Hazard Ratio

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+95% CI overlap 1

*95% CI don't overlap 1

Sparano et al. N Engl J Med 2019;380(25):2395-2405.

TAILORx: Impact of Clinical Risk (CR) on Prediction of Chemotherapy Benefit by Age in RS 11-25 Group (ET vs. Chemo +ET)

Grouped by Clinical Risk and Age

Total #/#IDFS/DR events

IDFS Hazard Ratio

DRFI Hazard Ratio

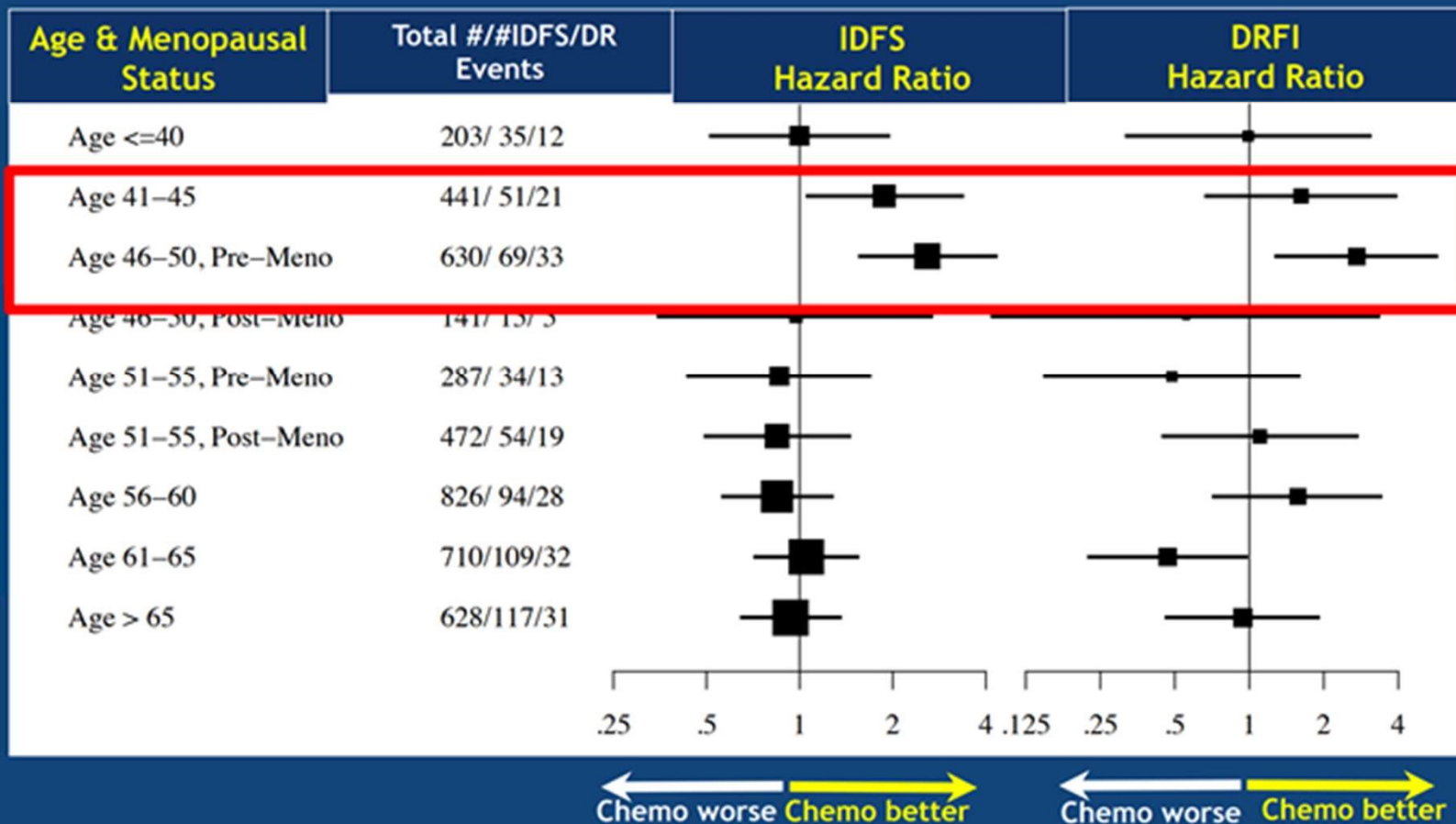
All Patients, Low Clinical Risk	4799/541/129	1.07+	1.03+
All Patients, High Clinical Risk	1697/270/111	1.02+	1.18+
Age > 50, Low Clinical Risk	3173/361/ 80	0.93+	0.90+
Age > 50, High Clinical Risk	1180/204/ 73	0.90+	0.95
Age <= 50, Low Clinical Risk	1626/180/ 49	1.45*	1.28+
Age <= 50, High Clinical Risk	517/ 66/ 38	1.56+	1.80+

Hazard ratio > 1 - chemo better
 +95% CI overlap with 1
 *95% CI don't overlap 1



Sparano et al. N Engl J Med 2019;380(25):2395-2405.

TAILORx: Exploratory Analysis - Impact of Age and Menopausal Status on Chemotherapy Benefit for RS 16-25



Sparano et al. N Engl J Med 2019;380(25):2395-2405.

TAILORx – ITT Population: Potential Chemotherapy benefit in women \leq 50 yrs (N=2216) in RS 11-25

- **RS 16-25 - some chemo benefit**
 - RS 16-20: 9% fewer IDFS events, including 2% fewer distant recurrences
 - RS 21-25: 6% fewer IDFS events, mainly consisting of fewer distant recurrences
- **RS 0-15 - good prognosis with endocrine therapy**
 - 3% distant recurrence with ET alone
 - no evidence for chemo benefit in RS 11-15



TAILORx: CTS5 in All Patients and According to Subgroups

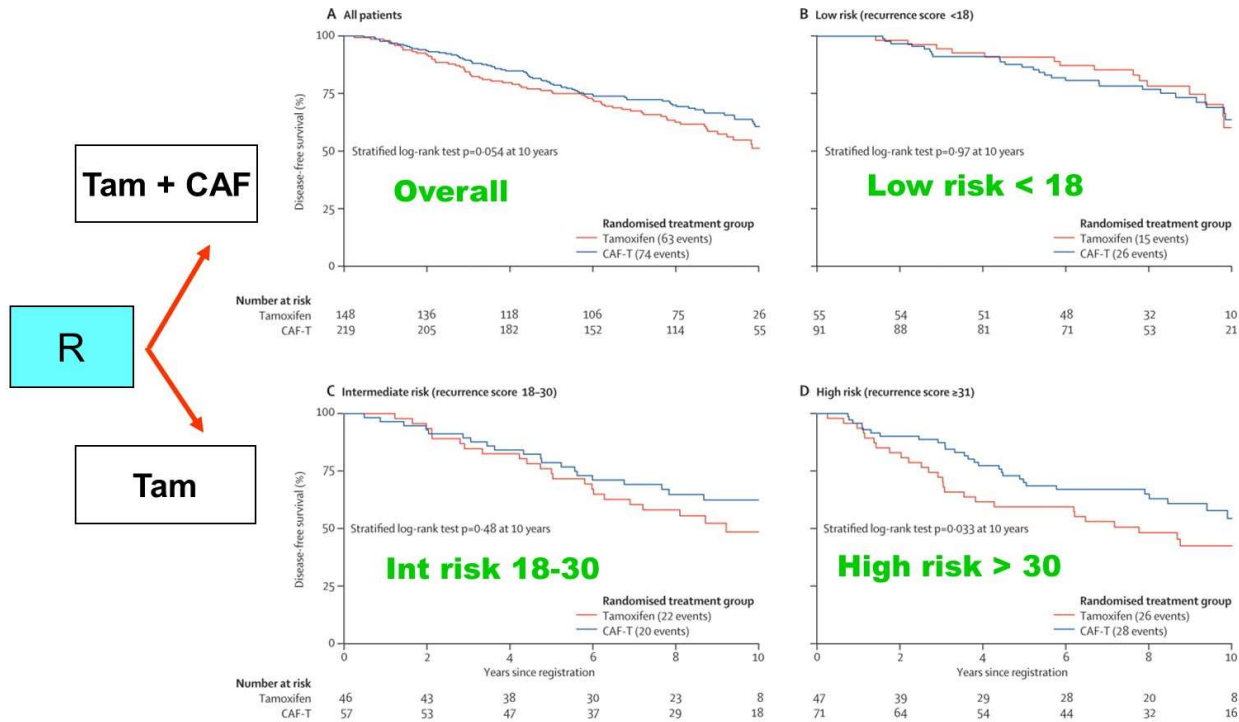
CTS5	RS	Treatment	HR	p-value
All patients (n = 7,353)	0-100	ET/CET	1.57	<0.0001
Arm A (n = 1,323)	0-10	ET	1.34	0.19
Arm B (n = 2,746)	11-25	ET	1.50	0.002
Arm C (n = 2,655)	11-25	CET	1.56	0.0003
Arm D (n = 629)	26-100	CET	1.90	0.004
Age ≤50 years (n = 2,259)	0-100	ET/CET	1.35	0.046
Age >50 years (n = 5,094)	0-100	ET/CET	1.78	<0.0001

ET = endocrine therapy; CET = chemo-ET

- Overall, CTS5 was highly prognostic for late distant recurrence (DR) stratified for assigned chemotherapy arm (HR = 1.57, $p < 0.0001$).
- Looking at each arm separately, CTS5 did not predict late DR in women with RS 0-10 and ET (arm A), but provided strong prognostic information for late DR in arms B (RS 11-25, ET), C (RS 11-25, CET), and D (RS 26-100, CET).
- CTS5 strongly predicted late DR in women >50 years (HR = 1.78, $p < 0.0001$), but to a lesser extent in women aged 50 years or younger (HR = 1.35, $p = 0.046$).

Sestak I et al. San Antonio Breast Cancer Symposium 2019;Abstract GS4-03.

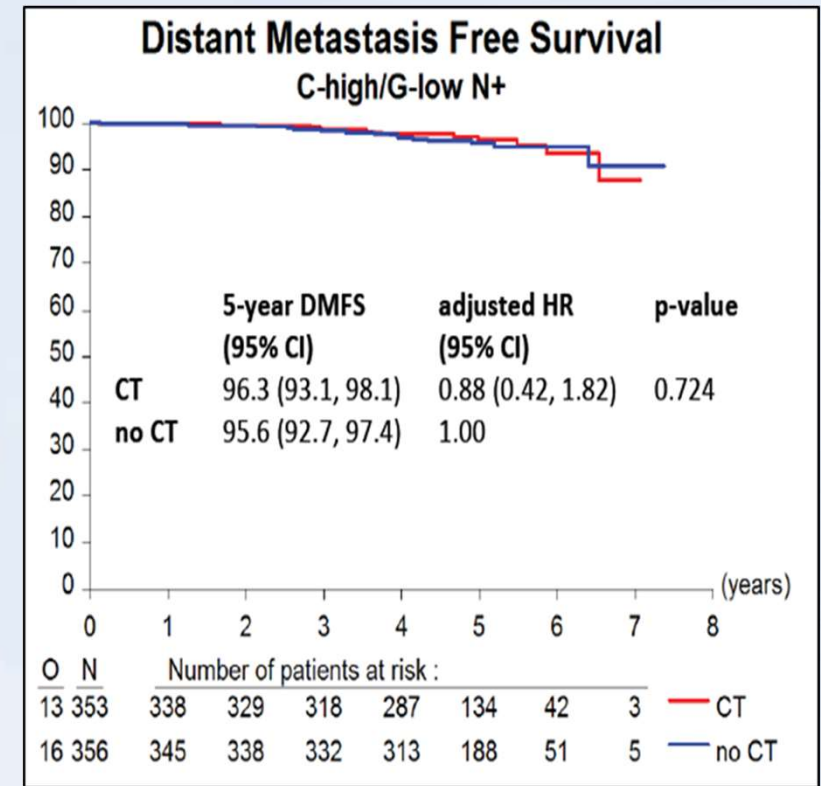
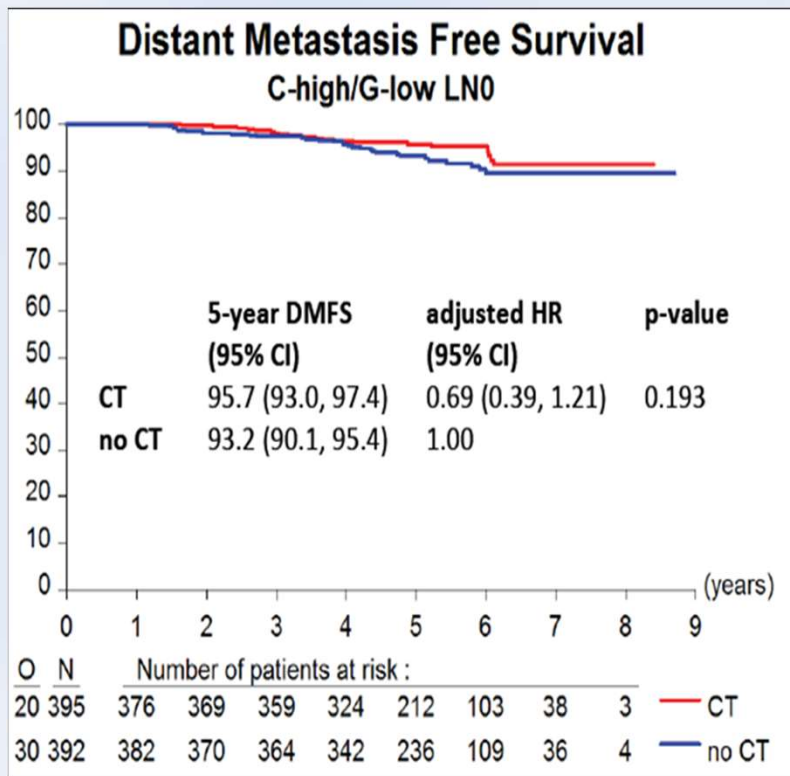
SWOG 8814: ER+ LN+, TAM ± CAF Chemotherapy



Albain KS, et al. *Lancet Oncol.* 2010;11(1):55-65.



MINDACT TRIAL



Take home message

