

Combined Analysis of the HER2DX Genomic Tool in Adjuvant APT and ATEMPT Trials



Paolo Tarantino^{1,2,3}, Guillermo Villacampa⁴, Noah T. Graham¹, Adrienne G. Waks^{1,2}, Patricia Villagrasa⁴, Fara Braso-Maristany⁴, Esther Sanfeliu⁴, Patricia Galvan⁴, Laia Pare⁴, Michelle Demeo¹, Ann H. Partridge^{1,2}, Harold J. Burstein^{1,2}, Ian E. Krop⁵, Nabihah Tayob^{1,2}, Eric P. Winer⁵, Aleix Prat^{4,6}, Sara M. Tolaney^{1,2}

Affiliations: ¹Dana-Farber Cancer Institute, Boston, USA; ²Harvard Medical School, Boston, USA, ⁶IDIBAPS, Barcelona, Spain; ⁵Yale Cancer Center, New Haven, USA, ⁶IDIBAPS, Barcelona, Spain

BACKGROUND

In the phase 2 APT trial, adjuvant paclitaxel and trastuzumab (TH) was associated with excellent long-term outcomes for patients with small, nodenegative HER2+ breast cancer. In the randomized phase 2 ATEMPT trial, adjuvant trastuzumab emtansine (T-DM1) also demonstrated outstanding outcomes in patients with stage 1 HER2+ breast cancer. HER2DX risk-score was associated with survival outcomes in both trials, separately.

METHODS

We conducted a retrospective analysis combining patients included in the APT (n=406) and ATEMPT (n=497) trials with available HER2DX data. The co-primary endpoints of the study were associations of HER2DX with relapse-free interval (RFI) and invasive disease-free survival (iDFS).

		Overall
		(n=471)
Study, n (%)	APT	284 (60.3)
	ATEMPT	187 (39.7)
Clinical tumor stage, n (%)	T1	440 (93.3)
	T2	31 (6.6)
Clinical nodal stage, n (%)	NO	462 (98.1)
	N1mic	9 (1.9)
Hormone receptor, n (%)	Negative	136 (28.9)
	Positive	335 (71.1)
Treatment, n (%)	TH	324 (68.8)
	T-DM1	147 (31.2)
HER2DX risk groups	Low	445 (94.5)
(original cut-off 50), n (%)	High	26 (5.5)
HER2DX risk groups	Low	385 (81.7)
(cut-off 32), n (%)	High	86 (18.3)
HER2DX pCR score groups, n (%)	Low	45 (9.6)
	Med	86 (18.3)
	High	340 (72.2)

Table 1 – Clinicopathologic characteristics and HER2DX scores for the patients included in the pooled analysis of APT/ATEMPT

The HER2DX risk-score was evaluated i) as a continuous variable (0 - 100), ii) using the predefined cut-off (50), and iii) using an exploratory optimal cutoff (32) derived by maximizing the log-rank statistic for the recurrence-free interval endpoint in the APT trial. The Kaplan-Meier method and stratified Cox models were used to estimate hazard ratios (HRs) to evaluate the association between HER2DX and survival outcomes.

RESULTS

Overall, 471 patients receiving TH (n=324) or T-DM1 (n=147) were included in the analysis, most having stage I (n=432) and HR+ disease (n=335) (Table 1). The median follow-up was 6.7 years (10.8 and 5.8 for APT and ATEMPT, respectively).

The median HER2DX risk-score was 13.9 (IQR 4.7-27.0), with 5.5% and 18.3% of the patients having HER2DX high-risk disease according to the predefined and optimal cut-offs, respectively. Most tumors showed a high expression of HER2 (55.6%), a high expression of proliferation signatures (55.6%), IgG signatures (52.9%), and luminal signatures (42.9%). (Figure 1)

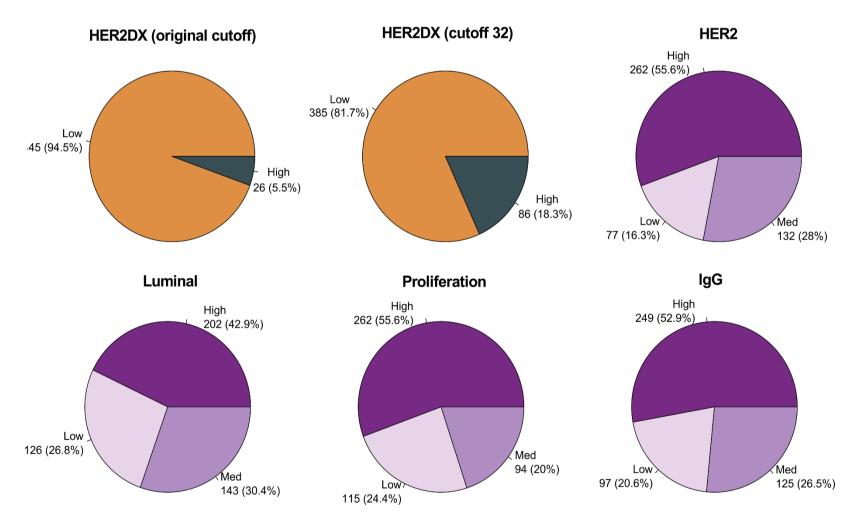


Figure 1 – HER2DX risk scores and expression of HER2, luminal, proliferation and IgG signatures among patients in APT/ATEMPT with available HER2DX data

HER2DX risk score as a continuous variable was found significantly associated with RFI (HR per 10-units: 1.39, 95%CI: 1.09-1.78; p=0.009) but not with iDFS (HR per 10-units: 1.18, 0.98-1.42; p=0.09).

Using the predefined cut-off (50), patients with HER2DX high-risk disease had higher RFI risk (HR: 7.33, 2.29-23.47, p<0.001), but the effect on iDFS was nonsignificant (HR: 2.78, 0.97-7.95, p=0.057) (**Figure 2, A-B**). The optimal cut-off (32) identified patients with low-risk disease (RFI at 7 years of 98.2%; 95% CI: 96.7%-99.6%) from those with high-risk (RFI at 7 years of 88.7%; 95% CI: 80.4%-97.8%) [delta of 9.5%], with a significant difference in both iDFS and RFI. (Figure 2, C-D)

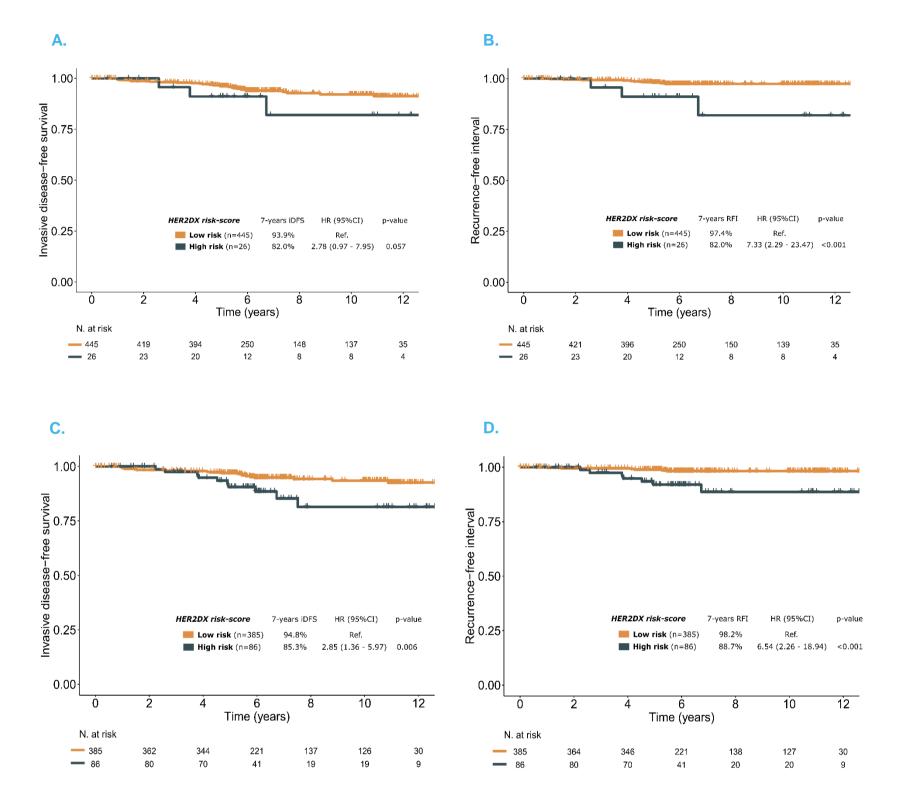


Figure 2 – 7-year iDFS and 7-year RFI by HER2DX score using the pre-defined risk score cutoff (50, curves A and B) or an exploratory optimal cutoff (32, curves C and D)

In the multivariable analysis, HER2DX group-risk maintained statistical significance after adjusting by HR status and tumor stage in RFI (HR: 6.87, 2.22-21.27, p<0.001) and iDFS (HR: 2.81, 1.26-6.23, p=0.01) (Table 2). Tumor size and HR expression were not found associated with RFI nor with iDFS (Figure 3).



CORRESPONDENCE TO: paolo_tarantino@DFCI.Harvard.edu

TWITTER: @PTarantinoMD

PRESENTER'S DISCLOSURES: PT has served as advisor/consultant for AstraZeneca, Daiichi Sankyo, Gilead, Eli Lilly and Roche.



Abstract #719

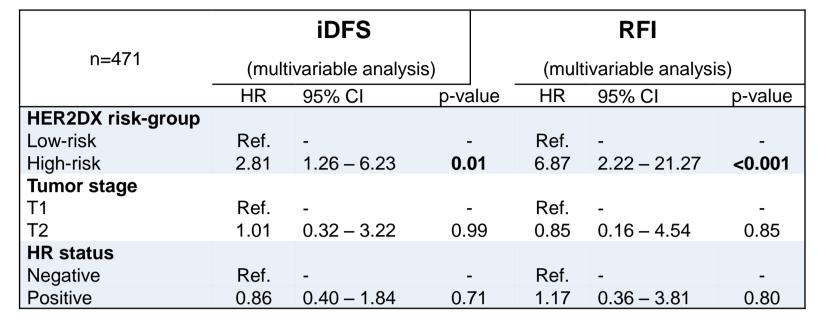


Table 2 – Multivariable analysis of iDFS and RFI in APT/ATEMPT by HER2DX score

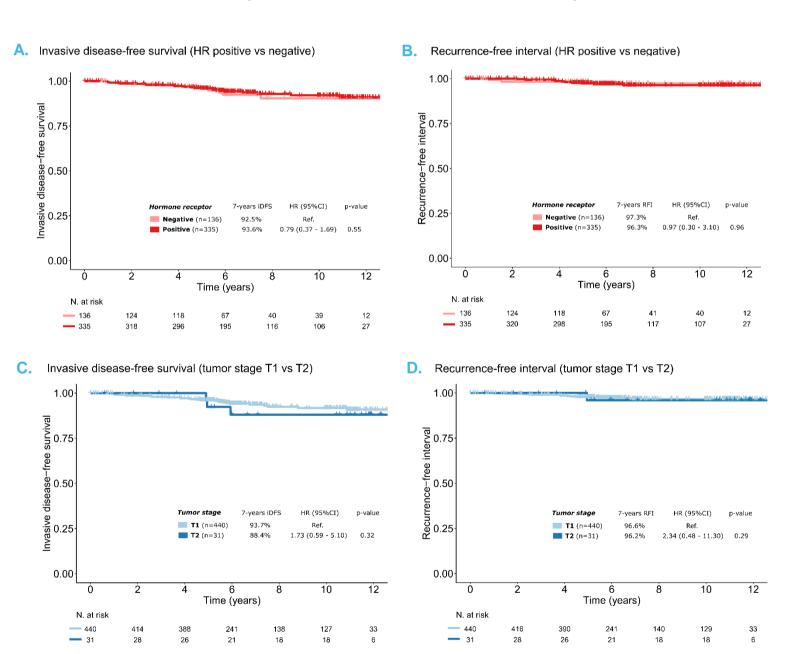


Figure 3 – iDFS and RFI by HR status (A-B) and tumor size (C-D) in the pooled analysis

CONCLUSION

The HER2DX score is associated with the risk of recurrence among patients with small, node negative HER2+ breast tumors, and, if validated, may help in tailoring treatments for this disease.