Development and validation of a composite biomarker (luminal A, *ERBB2*, and PR) predictive of palbociclib + endocrine treatment benefit in early breast cancer: PENELOPE-B and PALLAS trials

















# Objective



To define and independently validate a predictive biomarker that identifies patients with EBC who could preferentially benefit from the addition of palbociclib to ET in the adjuvant setting.

## Conclusions



- The composite predictive biomarker (LumA with ERBB2-high and/or LumA ER+/PR-) was defined from PENELOPE-B.
- Results from a prospectively defined retrospective analysis of a subset of intermediate/high risk patients with EBC selected from PALLAS validated the candidate predictive biomarker.
- Despite differences in the percent of patients who received prior chemotherapy in the PENELOPE-B and PALLAS HTG Sets, patients from both studies with biomarker-positive surgical resection tumor samples showed a significant improvement in iDFS with addition of palbociclib to ET.
- This predictive biomarker may be used for patient stratification and can potentially be applied to enrich future adjuvant clinical trials for the treatment of HR+/HER2- EBC.

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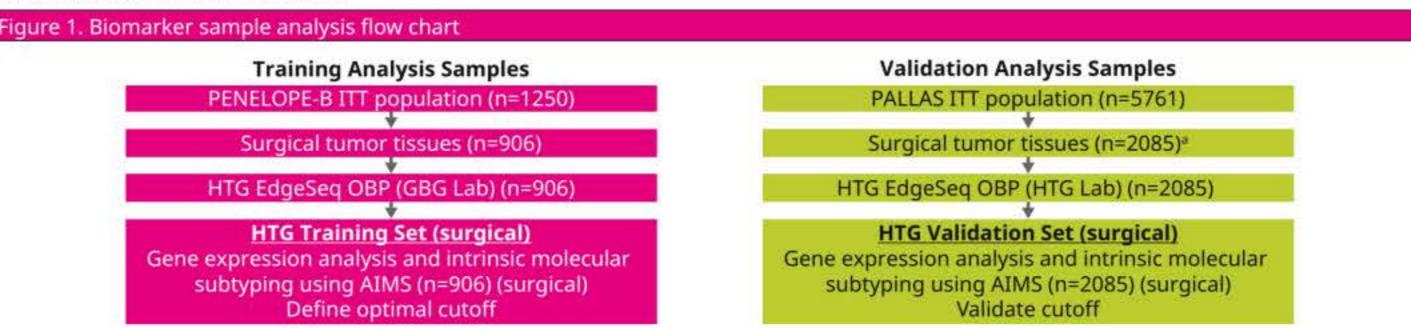
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## Introduction

- The large prospective, randomized, phase 3 PENELOPE-B (NCT01864746)1 and PALLAS (NCT02513394)2 trials evaluated adjuvant palbociclib + endocrine therapy (ET) versus ET in patients with hormone receptor-positive (HR+)/human epidermal growth factor receptor 2-negative (HER2-) early breast cancer (EBC); most patients in these studies were classified as high or intermediate risk.
- However, neither trial met the primary endpoint of improving invasive disease-free survival (iDFS).1-
- Based on evidence generated from biomarker analyses from PALOMA-2 and PALOMA-3, we identified 3 key elements for further investigation: 1. luminal subtypes,<sup>3,4</sup> 2. the *ERBB2* pathway,3 and 3. estrogen receptor (ER)/progesterone receptor (PR)
- Each of these biomarkers had prognostic value individually; we therefore hypothesized that a composite biomarker may be more precise in identifying the patients with EBC who could derive the most benefit from palbociclib treatment.

## Materials and Methods

#### ANALYSIS POPULATION



AIMS, absolute intrinsic molecular subtyping; CPS-EG, clinical pathologic stage and estrogen receptor status and histologic grade; GBG, German Breast Group;

A total of 2830 PALLAS patient samples were selected (resection samples were required; if resection sample was not available, biopsy sample was used). Sample selection nposed of ~50% "PENELOPE-like" patients (ie, received prior chemotherapy and had calculated CPS-EG score ≥3 or CPS-EG score 2 and ypN+) and ~50% randomly selected atient tissue samples so that biomarker validation results could be generalized to the PALLAS population. Gene expression data for N=2604 samples (n=2085 resection samples and n=519 biopsy samples) was generated. Only surgical samples (n=2085) were used for the current analysis

#### GENE EXPRESSION AND STATISTICAL ANALYSES

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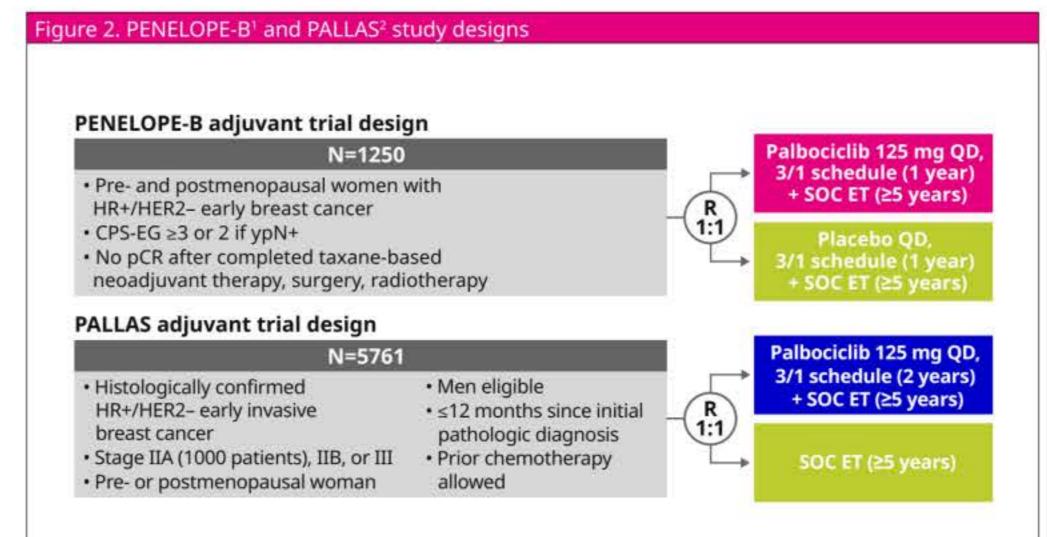
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- Gene expression analyses were conducted on the patient tumor samples using the HTG EdgeSeq Oncology Biomarker Panel.
- HTG gene expression analysis of the PENELOPE-B HTG Training Set was conducted at GBG Labs and was used to develop a composite biomarker by using a biologically driven, targeted gene interrogation.
- HTG absolute intrinsic molecular subtyping (HTG-AIMS) classifications of LumA, LumB, HER2-enriched, basal-like, and normal-like were provided by the GBG Team for PENELOPE-B.7
- A outcome-oriented approach based on iDFS utilizing a selection procedure that maximized the log-rank statistic was
- applied to estimate a standard Z score-based optimal cutoff for ERBB2 gene expression.
- These data were coupled with PR immunohistochemistry (IHC) data and analyzed to refine the composite biomarker.
- The biomarker-positive population was defined to consist of patients with tumors that were LumA with ERBB2-high ("high" as defined by the Z score-based optimal cutoff) and/or LumA ER+/PR-; the biomarker-negative population was all
- iDFS rates at 3 years were estimated using the Kaplan-Meier method, and iDFS distributions between the treatment groups were compared using a log-rank test. Hazard ratios (HRs) and 95% confidence intervals (CIs) were estimated using Cox proportional hazards models. Interaction between treatment and composite biomarker status (positive vs negative) was assessed
- Multivariable analyses based on the trial stratification factors were performed to adjust for potential baseline confounders.

## Results

### BASELINE AND CLINICAL CHARACTERISTICS



PS-EG, clinical pathologic stage and estrogen receptor status and histologic grade; ET, endocrine therapy; HER2, human epidermal growth factor receptor 2; HR, hormone receptor; pCR, pathological complete response; R, randomized; SOC, standard of care; QD, daily.

- No differences in iDFS were observed between treatment groups in the ITT population and HTG Set PENELOPE-B
- ITT population (N=1250), treatment effect HR: 0.92 (95% CI: 0.73–1.17), P=0.50
- HTG Training Set (N=906), treatment effect HR: 0.88 (95% CI: 0.68–1.14), P=0.35
- ITT population (N=5761), treatment effect HR: 0.96 (95% CI: 0.81–1.14), P=0.65 HTG Validation Set (N=2085), treatment effect HR: 1.02 (95% CI: 0.79-1.33), P=0.86
- Table 1. Baseline characteristics in the HTG Sets were representative of the ITT populations in PENELOPE-B1 and PALLAS2 DENIEL ODE D. DALLAC

Characteristics n (%)	PENELOPE-B HTG Training Set (N=906)		PALLAS HTG Validation Set (N=2085)	
	Palbociclib + ET (n=464)	Placebo + ET (n=442)	Palbociclib + ET (n=1070)	ET alone (n=1015)
Age, years ≤50 >50	266 (57.3) 198 (42.7)	237 (53.6) 205 (46.4)	425 (40) 644 (60)	418 (41) 597 (59)
Race Asian Black or African American White Other Unknown	45 (9.7) 5 (1.1) 361 (77.8) 53 (11.4)	36 (8.1) 5 (1.1) 349 (79.0) 52 (11.8)	36 (3.4) 22 (2.1) 963 (90) 4 (0.4) 45 (4.2)	37 (3.6) 24 (2.4) 909 (90) 9 (0.9) 36 (3.5)
Geographical region Asia Pacific Europe North America Other	74 (15.9) 332 (71.6) 58 (12.5)	57 (12.9) 331 (74.9) 54 (12.2)	72 (6.7) 528 (49) 460 (43) 10 (0.9)	62 (6.1) 502 (49) 442 (44) 9 (0.9)
Menopausal status Postmenopausal Premenopausal Not applicable Unknown	236 (50.9) 228 (49.1) - -	226 (51.1) 216 (48.9) - -	636 (59) 427 (40) 6 (0.6) 1 (<0.1)	590 (58) 419 (41) 5 (0.5) 1 (<0.1)

#### able 1. Baseline characteristics in the HTG Sets were representative of the ITT populations in PENELOPE-B1 and PALLAS2 (continued) PENELOPE-B PALLAS HTG Validation Set (N=2085) Palbociclib + ET | Placebo + ET | ER+/PR-ER+/PR+ 796 (74) 368 (79.3) 337 (76.2) 788 (78) 2 (0.2) ER-/PR+ 2 (0.5) 0 (0) 1 (0.2) 1 (<0.1) ER-/PR-0(0) Nodal status at surgery 224 (50.7) 212 (45.7) ypN 2-3 252 (54.3) 218 (49.3) 722 (67) N0/N0(i+)/N1/NX 198 (20) 224 (21) 116 (11) 123 (11) 1 (<0.1) 0 (0) 4 (0.4) 1 (0.2) 3 (0.3) 239 (22) 246 (24) 29 (6.3) 35 (7.9) 342 (34) 185 (39.9) 195 (44.1) 352 (33) 249 (53.7) 211 (47.7) 424 (42) 474 (44) 0 (0) 1 (<0.1) Tumor grading at surgery 15 (3.4) 135 (13) 233 (50.2) 207 (46.8) 646 (60) 626 (62) 219 (50.0) 206 (20) 233 (22) 221 (47.6) 48 (4.7) 1 (0.2) 1 (0.2) 3 (0.3) 0(0) Central Ki-67 at surgery 132 (28.5) 133 (30.1) 0(0) (0.9)CPS-EG score ≥3 282 (60.8) 182 (39.2) 179 (40.5) CPS-EG score 2 and ypN+ Prior neoadjuvant or adjuvant chemotherapy 435 (43) 580 (57) 438 (41) 442 (100) 632 (59) 444 (44) 474 (44) 116 (11) 20 (2)

Other included Native Hawaiian or other Pacific Islander and American Indian or Alaska Native. bClinical stage in

CPS-EG, clinical pathologic stage and estrogen receptor status and histologic grade; ER, estrogen receptor; T, endocrine therapy; ITT, intention-to-treat; PR, progesterone receptor.

AIMS, absolute intrinsic molecular subtyping; HER2, human epidermal growth factor receptor 2.

Table 2. HTG-AIMS intrinsic molecular subtype distributions were similar between PENELOPE-B and PALLAS HTG Sets, as were the subtype prognostic profiles8 (data not shown) PENELOPE-B HTG Training Set **PALLAS HTG Validation Set** 

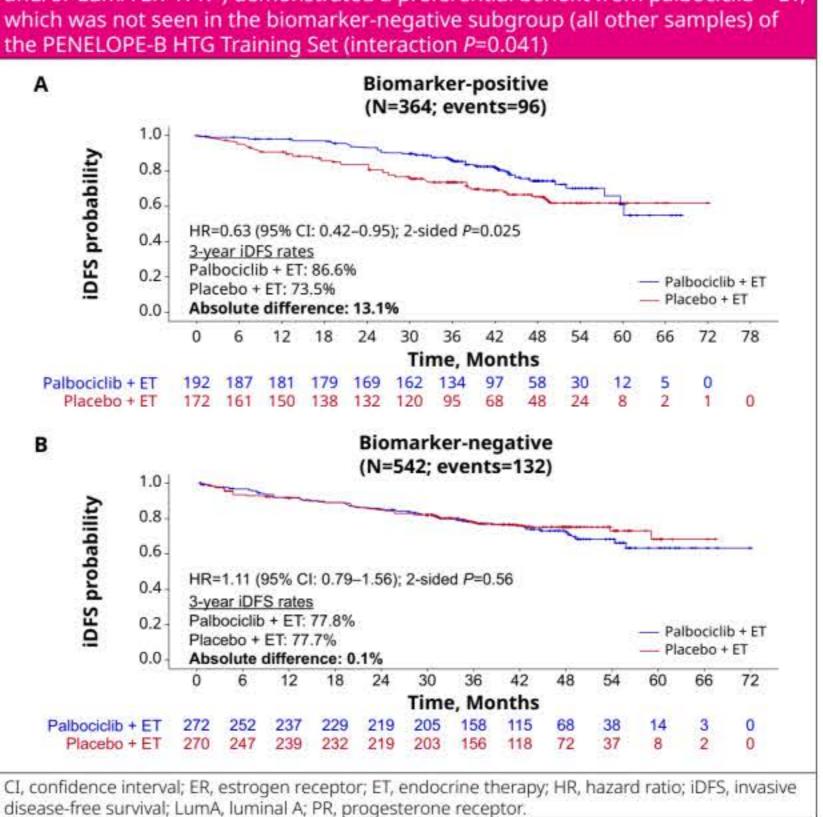
(N=2085) (N=906) (HTG-AIMS), n (%) 28 (3.1) 49 (2.5) HER2-enriched 1516 (72.7) 663 (73.2) uminal A 172 (8.2) Luminal B 311 (13.6) 135 (14.9) Normal-like

## COMPOSITE BIOMARKER IDENTIFICATION AND VALIDATION

#### PENELOPE-B HTG TRAINING SET (N=906)

of Texas MD Anderson Cancer Center, Houston, TX, USA

#### Figure 3. The biomarker-positive subgroup (composite of LumA with ERBB2-high and/or LumA ER+/PR-) demonstrated a preferential benefit from palbociclib + E1 which was not seen in the biomarker-negative subgroup (all other samples) of e PENELOPE-B HTG Training Set (interaction P=0.041)



The number of patients from the PENELOPE-B HTG Training Set with different components of the biomarker was n=298 LumA with ERBB2-high (32.9%), n=122 LumA with ER+/PR- (13.5%), n=485 ERBB2-high and/or ER+/PR- (53.5%) and n=364 LumA with ERBB2-high and/or LumA ER+/PR- (40.2%).

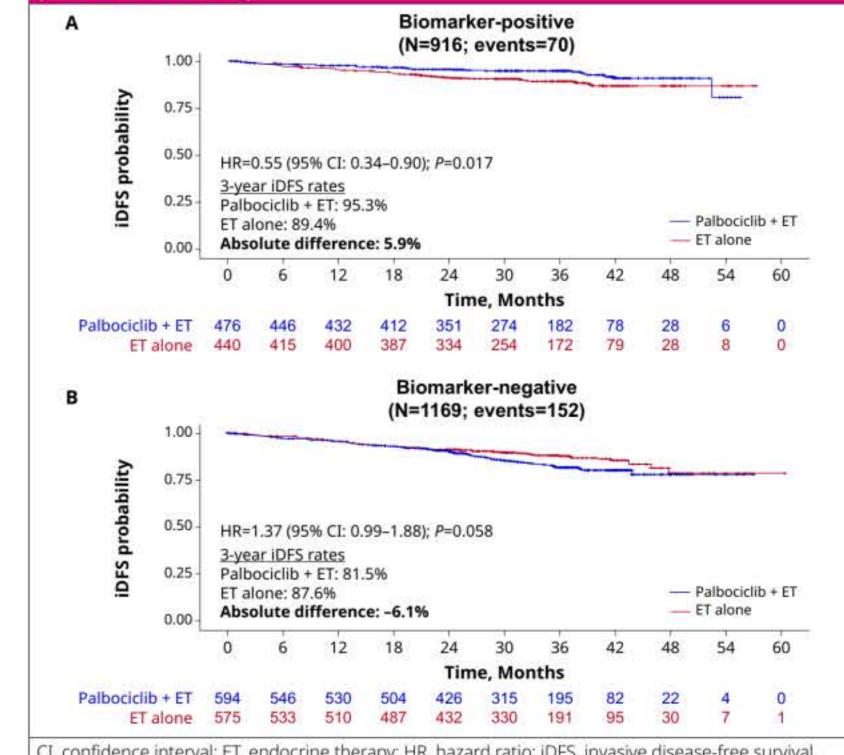
## Table 3. Significant treatment effect remained in the PENELOPE-B biomarkerositive subgroup (n=364) after adjusting for potential confounding from the

	PENELOPE-B Biomarker-positive subgroup		
Variable	HR (95% CI)	P-value	
Treatment Palbociclib + ET vs placebo + ET	0.63 (0.42-0.95)	0.03	
Age at first diagnosis (years) ≤50 vs >50	0.99 (0.65-1.50)	0.96	
Central Ki67 ≤15% vs >15%	0.67 (0.43-1.03)	0.07	
Global region Asian vs non-Asian	0.68 (0.27-1.70)	0.41	
Histological lymph node status at surgery ypN 0-1 vs ypN 2-3	0.95 (0.63-1.44)	0.81	
Risk status CPS-EG 2 and ypN+ vs CPS-EG ≥3	0.75 (0.49-1.15)	0.19	

### PALLAS HTG VALIDATION SET (N=2085)

## Figure 4. Independent validation of the biomarker with tumor samples from the

PALLAS HTG Validation Set confirmed significant benefit from palbociclib + ET in the biomarker-positive subgroup, but not in the biomarker-negative subgroup



confidence interval; ET, endocrine therapy; HR, hazard ratio; iDFS, invasive disease-free survival.

The number of patients from the PALLAS HTG Validation Set with different components of the biomarker was n=763 LumA with ERBB2-high (36.6%), n=294 LumA with ER+/PR- (14.1%), n=1206 ERBB2-high and/or ER+/PR- (57.8%) and n=916 LumA with ERBB2-high and/or LumA ER+/PR- (43.9%).

Table 4. Significant treatment effect remained in the PALLAS biomarker-positive subgroup (n=916) after adjusting for potential confounding from the stratification actors used at randomization (multivariate Cox regression analysis)

	PALLAS Biomarker-positive subgroup		
Variable	HR (95% CI)	<i>P</i> -value	
Treatment Palbociclib + ET vs ET alone	0.55 (0.34-0.89)	0.02	
Age at randomization (years) ≤50 vs >50	0.97 (0.60-1.58)	0.91	
Anatomic stage IIA vs IIB/III	0.67 (0.35-1.30)	0.24	
Neo/adjuvant chemotherapy No vs yes	0.48 (0.27-0.82)	0.01	
Geographic region Europe vs North America Other vs North America Europe vs other	0.60 (0.36-0.98) 0.85 (0.36-2.04) 0.70 (0.29-1.69)	0.04 0.72 0.42	

confidence interval; ET, endocrine therapy; HR, hazard ratio. This presentation is the intellectual property of the author/presenter. Contact them at Sibylle.Loibl@gbg.de for permission to reprint and/or distribute. Copyright ©2022. All rights reserved.