



# Phase I study of alpelisib (BYL-719) and trastuzumab emtansine (T-DM1) in HER2-positive metastatic breast cancer (MBC) after trastuzumab and taxane therapy

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## Abstract

**Purpose** Activation of the phosphoinositide 3-kinase (PI3K) pathway is an important resistance mechanism to anti-HER2 therapies. This study aimed to assess the safety and activity of alpelisib (a PI3K $\alpha$  isoform-specific inhibitor) with T-DM1 in trastuzumab- and taxane-resistant HER2-positive MBC.

**Methods** Patients with HER2-positive MBC that had progressed on trastuzumab-based therapy were treated with alpelisib daily and T-DM1 3.6 mg/kg every 3 weeks. The dose-limiting toxicity (DLT), maximum tolerated dose (MTD), adverse events, overall response rate (ORR), and clinical benefit rate (CBR = CR + PR + SD > 6 months) were assessed with descriptive statistics. Progression-free survival (PFS) was calculated by the Kaplan–Meier method.

**Results** Seventeen patients were enrolled with a median of 3 prior therapies for metastatic disease. The DLT was a maculopapular rash and MTD was 250 mg alpelisib daily. The most frequently occurring toxicities included fatigue, rash, gastrointestinal side effects, thrombocytopenia, anemia, elevated liver enzymes, and hyperglycemia. Fourteen patients were evaluable for response with an ORR of 43%. In patients with prior treatment and progression on T-DM1 ( $n = 10$ ), the ORR was 30%. The CBR was 71% in evaluable patients and 60% in those with prior T-DM1. The median PFS was 8.1 months.

**Conclusions** The combination of alpelisib and T-DM1 is tolerable and demonstrates activity in trastuzumab-resistant HER2-positive MBC. Furthermore, activity was observed in T-DM1-resistant disease. These data suggest that PIK3CA inhibition targets an important resistance pathway to anti-HER2 therapy, providing rationale for further study of PI3K inhibition in refractory HER2-positive MBC to validate these results.

**Keywords** Alpelisib · HER2-positive · T-DM1 · PI3-Kinase · PIK3CA

## Background

The human epidermal growth factor receptor 2 neu (HER2) is an oncogene overexpressed in 20% of breast cancers. HER2 activation leads to downstream signaling along the PI3K/AKT/mTOR and RAS/RAF/MAPK pathways and

promotes cell motility, survival and proliferation, and resistance to apoptosis [1]. HER2 overexpression confers a more aggressive breast cancer phenotype, which prior to the introduction of trastuzumab, was associated with a poor prognosis [2–4]. The development of HER2-targeted therapies has resulted in significant improvements in the median progression-free and overall survival (PFS and OS), which is now over 4 years in HER2-positive MBC [5–8]. The EMILIA and TH3RESA studies demonstrated improvement in PFS and OS with the antibody–drug conjugate T-DM1 when compared to capecitabine and lapatinib or treatment of physician’s choice, establishing T-DM1 as a standard second-line treatment for HER2-positive MBC [9–12]. Unfortunately, therapy for HER2-positive metastatic disease is not curative and despite an initial response to HER2-targeting, patients

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eventually develop resistance. In fact, the median PFS for second-line T-DM1 is only 6 months [9, 11].

While there are multiple resistance mechanisms to anti-HER2 therapy, the PI3K/AKT pathway is downstream from the site of action of HER2-targeted drugs, making its activation an important resistance pathway [13]. Activation of the PI3K/AKT pathway is frequently seen in breast cancers that progress on HER2-targeted therapy [14–20]. Recent clinical trials demonstrated hyperactivity of the PI3K pathway in 47 and 41% of HER2-positive MBC in the trastuzumab-naïve and trastuzumab-resistant setting, respectively. Mutations in *PIK3CA* were noted in 30% of cases [20]. Furthermore, tumors with *PIK3CA* mutations are less responsive to chemotherapy and anti-HER2 therapy [21]. This data as well as preclinical studies have raised significant interest in targeting the PI3K/AKT/mTOR pathway in HER2-positive breast cancer [22–25].

The BOLERO 1 and 3 trials evaluated the mTOR inhibitor everolimus in HER2-positive metastatic breast cancer (MBC), but showed only a small benefit at the cost of significant toxicity [26, 27]. Thus, developing an isoform-specific PI3K inhibitor targeting PI3K $\alpha$  which is encoded by *PIK3CA* with the goal of maximizing activity but limited off-target toxicities is of interest. Alpelisib (BYL-719) is an oral selective PI3K $\alpha$  isoform inhibitor [28, 29]. Clinical trials in patients with advanced solid tumors with *PIK3CA* alterations and advanced hormone receptor-positive, HER2-negative breast cancer have shown safety and tolerability of alpelisib [30, 31].

We were interested in evaluating alpelisib in trastuzumab-resistant MBC given the compelling data demonstrating the frequency and relevance of PI3K pathway activation in this setting. T-DM1 was chosen as the companion anti-HER2 therapy because of its established role as second-line therapy with more limited activity of trastuzumab in this setting [32, 33]. We conducted a phase I clinical trial to determine the maximum tolerated dose (MTD), safety, and activity of alpelisib with T-DM1 in HER2-positive MBC. We hypothesized that the combination of a specific PI3K $\alpha$  inhibitor alpelisib with T-DM1 would have acceptable toxicities and demonstrate activity in trastuzumab-refractory HER2-positive breast cancer patients.

## Patients and methods

### Patient population

Patients had histologically confirmed HER2-positive inoperable locally advanced or MBC. Locally advanced disease must be inoperable and have progressed within 6 months of trastuzumab and/or taxane treatment and MBC must have progressed during or after trastuzumab and taxane therapy.

Other key inclusion criteria included ECOG performance status  $\leq 2$  and adequate bone marrow, renal, and hepatic functions. Patients were not required to have measurable disease. Key exclusion criteria included clinically manifested diabetes (fasting plasma glucose  $> 140$  mg/dL or a history of steroid-induced diabetes), known history of liver disease, or other active medical problems.

Approval was obtained from the ethics committee at Northwestern University and regulatory authorities. All patients gave informed consent. The study followed the Declaration of Helsinki and Good Clinical Practice guidelines.

### Study design

This was a phase I, open-label, single-arm, single center, dose escalation study of alpelisib in combination with T-DM1 using a standard 3 + 3 design with a dose expansion cohort at the maximum tolerated dose (MTD). Patients received T-DM1 3.6 mg/kg (based on the FDA-approved dose) on day 1 of each cycle  $\pm 3$  days (1 cycle = 21 days). The first cohort received alpelisib at 300 mg oral daily (25% below the single-agent MTD). Patients were instructed to take alpelisib about 1 h after a light meal at approximately the same time daily. The planned alpelisib escalation tiers were 350 and 400 mg daily and de-escalation tiers were 250, 200, and 150 mg daily.

The primary endpoints were MTD and frequency and severity of toxicities of alpelisib in combination with T-DM1. All patients who received at least 1 dose of the study medication and T-DM1 were considered evaluable for the primary endpoint. The secondary endpoints were preliminary efficacy by assessment of PFS and objective response rate (ORR). All patients who received at least 3 cycles of therapy and had imaging at first response assessment were considered evaluable for the secondary endpoint.

Toxicities were defined using the NCI's CTCAE version 4.03. Dose-limiting toxicity (DLT) was defined as a clinically significant grade  $\geq 3$  toxicity being at least possibly related to the study medication, which occurred  $\leq 21$  days following the first dose of alpelisib or an adverse event grade  $< 3$  that led to dose interruption of alpelisib for greater than 7 days. The MTD was defined as the highest dose at which no more than one-third of patients experienced any DLTs during the first cycle. Patients were treated until disease progression, unacceptable toxicity, investigator decision, or withdrawal of consent.

### Safety and efficacy assessments

Patient assessment by clinical and laboratory evaluation was conducted at baseline and on day 1 of each 21-day cycle. ECG and hemoglobin (Hgb) A1c were assessed at baseline and every 3 cycles. PFS was defined as time from the first

study treatment to the first occurrence of progression or death. ORR was assessed every 3 cycles with imaging by CT of the chest, abdomen, and pelvis and bone scan or Positron Emission Tomography (PET)/CT. ORR (partial or complete response) was defined as the best response by RECIST v1.1 criteria, and by clinical examination for subjects with only skin disease (followed with serial photographs).

### Statistical analysis

The target sample size was 19–28 patients depending on occurrence of DLTs. This 3 + 3 design had a 91% chance of dose escalating when the true toxicity rate for that dose was 10%. Response was analyzed using exact binomial methods and 95% confidence intervals. PFS was analyzed using Kaplan–Meier curves. Exploratory objectives were analyzed by performing a descriptive analysis of the frequency of specific genetic alterations.

### Correlative studies

Expression of PTEN and AKT were assessed by immunohistochemistry of archival tumor specimen (from a banked specimen or optional biopsy). PIK3CA gene mutation status was recorded when known from prior next generation sequencing of a primary or metastatic tumor.

## Results

### Study population

Between May 2014 and February 2015, 17 patients with HER2-positive MBC were enrolled and received at least 1 dose of alpelisib (Table 1). The median age was 53 (range 40–67). The median number of metastatic sites was 3 (1–5) and 82% ( $n = 14$ ) of patients had visceral disease. The most common sites of metastases were bone (59%), liver (53%), lymph nodes (47%), lung (41%), and brain (29%). Nine patients (53%) had hormone receptor-negative disease. Eight patients (47%) initially presented with de novo metastatic disease. In the metastatic setting patients had received a median of 3 lines (range 0–12) of prior therapy. All patients had received prior trastuzumab and taxane (94% for metastatic disease and 6% in the (neo)adjuvant setting). For treatment of metastatic disease, 94% received pertuzumab, 47% received lapatinib, and 59% received T-DM1. Five (50%) of patients with prior T-DM1 had previously experienced a clinical response to T-DM1. All patients with prior T-DM1 ( $n = 10$ ) had progression of disease on T-DM1 (after median 8 cycles). T-DM1 was the most recent therapy in 4 patients.

### DLT and MTD

Of the 17 patients who received alpelisib, 6 patients received a starting dose of 300 mg daily (cohort 1) and 11 received a starting dose of 250 mg daily (de-escalation cohort – 1). In cohort 1, 1 patient was deemed not evaluable due to death from rapidly progressive disease after only 1 dose of alpelisib. Of the initial 3 evaluable patients receiving 300 mg daily, 1 patient developed grade 3 thrombocytopenia that was attributed to T-DM1 (the patient had thrombocytopenia with prior T-DM1 therapy and this toxicity resolved upon stopping T-DM1). The data monitoring committee recommended enrollment of 3 additional patients at 300 mg daily to ensure safety of the combination. However, the next 2 patients experienced DLTs (both grade 3 rash). Therefore, de-escalation cohort – 1 was opened and enrolled 3 patients at a starting dose of 250 mg daily, in which no DLTs occurred. Accordingly, the MTD and recommended phase II dose for alpelisib in combination with T-DM1 was determined to be 250 mg daily with rash as the DLT at the 300 mg dose. A dose expansion cohort of an additional 8 patients was enrolled for a total of 11 patients who received alpelisib 250 mg daily as the starting dose. Three (27%) of these patients developed a grade 3 rash and 1 (9%) developed grade 3 hyperglycemia during cycle 1 (Table 2).

### Adverse events (AE)

The most common treatment-related non-hematologic toxicities were elevation in transaminases ( $n = 13$ , 76%), hyperglycemia ( $n = 9$ , 53%), fatigue ( $n = 9$ , 53%), rash ( $n = 8$ , 47%), and nausea ( $n = 7$ , 41%) (Table 3). Hematologic toxicities included thrombocytopenia ( $n = 9$ , 53%), anemia ( $n = 7$ , 41%), and leukopenia ( $n = 5$ , 29%). Elevation in transaminases was always grade 1 or 2, was thought to be at least partially related to T-DM1, and did not require any treatment modifications or other management. T-DM1 was thought to be responsible for most hematologic toxicity, particularly in this heavily pre-treated population (Table 4).

Ten patients (59%) experienced a  $\geq$  grade 3 treatment-related toxicity. Grade 3 treatment-related AEs included rash ( $n = 7$ ), hyperglycemia ( $n = 3$ ), anorexia ( $n = 2$ ), hypertension ( $n = 2$ ), thrombocytopenia ( $n = 1$ ), weight loss ( $n = 1$ ), QTC prolongation ( $n = 1$ ), pancreatitis ( $n = 1$ ), and anemia/abnormal uterine bleeding ( $n = 1$ ). One Grade 4 AE occurred (thrombocytopenia) likely due to T-DM1. One grade 5 AE occurred on study, which was thought to be unrelated to the study drug in a patient who received only 1 dose of the study drugs. The patient developed hypoxic respiratory failure with a CT chest showing marked progression of metastatic disease with likely superimposed infection.

**Table 1** Patient characteristics

Entire cohort ( <i>n</i> = 17)	<i>N</i> (%) unless otherwise specified
Median age	53 years (range 40–67)
PS	
0	2 (12%)
1	12 (71%)
2	3 (18%)
Hormone receptor status	
Positive	8 (47%)
Negative	9 (53%)
Median number of metastatic sites [median (range)]	3 (1–5)
Visceral disease	14 (82%)
Lymph nodes	8 (47%)
Bone	10 (59%)
Liver	9 (53%)
Lung	7 (41%)
Brain	5 (29%)
Skin	4 (24%)
Other	3 (18%)
Median time since diagnosis	34 months (range 16–132 months)
Setting of last trastuzumab treatment	
Neoadjuvant/adjuvant	1 (6%)
Metastatic	16 (94%)
Lines of prior systemic therapy in the metastatic setting	3 (range 0–12)
Trastuzumab	16 (94%)
Pertuzumab	16 (94%)
Lapatinib	8 (47%)
T-DM1	10 (59%)
Taxane	16 (94%)
Gemcitabine	3 (18%)
Capecitabine	8 (47%)
Vinorelbine	6 (35%)
Eribulin	6 (35%)
Other chemotherapy	3 (18%)
Patients with prior T-DM1 ( <i>n</i> = 10)	
Median age	53.5 (range 40–64)
HR-positive	4 (40%)
Prior lines of therapy in the metastatic setting including T-DM1	7 (2–12)
Median cycles of T-DM1 received prior to enrollment (approximate)	8 (3–18)
T-DM1 as most recent therapy	4 (40%)
Best clinical response to prior T-DM1	
PR	5 (50%)
SD	2 (20%)
PD	1 (10%)
Unknown	2 (20%)
Prior progression on T-DM1	10 (100%)

### Management of serious adverse events

The DLT and most common serious adverse event was a maculopapular rash (*n* = 7, 41%). No standard premedication was recommended by the study protocol initially. The

rash developed cycle 1, days 10 to 15 in 6 of 7 patients with a grade 3 rash. This occurred in both the 300 mg alpelisib and 250 mg alpelisib cohort. The grade 3 rash resolved in a median of 8 days (range 5–15) with alpelisib interruption and dose reduction. Topical steroid cream, oral antihistamines,

**Table 2** Dose-limiting toxicities and dose modifications of apelisib

	Any dose	300 mg	250 mg	200 mg
Dose-limiting toxicity				
Rash	5	2	3	
Hyperglycemia	1		1	
Thrombocytopenia <sup>a</sup>	1	1		
Toxicity resulting in apelisib dose modification				
Rash	5	1	4	
Hyperglycemia	3	1	1	1
Anorexia/weight loss	2	1	1	
Prolonged QTC	1		1	

<sup>a</sup>Thrombocytopenia thought to be related to TDM1

**Table 3** Apelisib- and TDM1-related adverse events

	Grade 1/2	Grade 3/4	Any grade
General clinical symptoms			
Fatigue	9 (53)	0	9 (53)
Maculopapular rash	1 (6)	7 (41)	8 (47)
Dizziness	2 (12)	0	2 (12)
Cough	2 (12)	0	2 (12)
Dyspnea	2 (12)	0	2 (12)
Peripheral sensory neuropathy	2 (12)	0	2 (12)
Gastrointestinal and metabolic			
Elevated transaminases	13 (76)	0	13 (76)
Hyperglycemia	6 (35)	4 (24)	10 (59)
Elevated alkaline phosphatase	7 (41)	0	7 (41)
Nausea	7 (41)	0	7 (41)
Hypokalemia	6 (35)	0	6 (35)
Anorexia	2 (12)	2 (12)	4 (24)
Elevated bilirubin	4 (24)	0	4 (24)
Low albumin	3 (18)	0	3 (18)
Diarrhea	3 (18)	0	3 (18)
Weight loss	2 (12)	1 (6)	3 (18)
Dry mouth	3 (18)	0	3 (18)
Pancreatitis	0	1 (6)	1 (6)
Cardiovascular			
Hypertension	1 (6)	2 (12)	3 (18)
QTc prolongation	2 (12)	1 (6)	3 (18)
Hematologic			
Thrombocytopenia	6 (35)	3 (18)	9 (53)
Anemia	6 (35)	1 (6)	7 (41)
Leukopenia	5 (29)	0	5 (29)
Lymphopenia	4 (24)	0	4 (24)
Neutropenia	4 (24)	0	4 (24)
Abnormal uterine bleeding	0	1 (6)	1 (6)

oral steroids, and dermatology consultation were ordered as needed. Oral steroids (prednisone 30–80 mg daily) were only used in cohort – 1 ( $n=4$ ) over a 5- to 21-day course that

included a taper for some patients. The rash was biopsied in one patient and demonstrated perivascular lymphocytic dermatitis, suggestive of dermal hypersensitivity reaction such as a drug eruption. After improvement in the grade 3 rash, some patients experienced a grade 1 or 2 rash that was managed with as needed topical steroid cream and oral antihistamines per protocol.

Other grade 3/4 toxicities were managed as follows: hyperglycemia was managed with diet modification, anti-hyperglycemic medications, and endocrinology consultations when necessary; anorexia and weight loss were managed with nutritional supplements and dose reduction; hypertension reaching the 160 s mm Hg systolic, was asymptomatic, and resolved without intervention; QTC prolongation in a patient with a history of prolonged QTC resolved with dose reduction; grade 3 pancreatitis was thought to be possibly apelisib related and was managed as per hospital protocol; and hematologic toxicities (thrombocytopenia, anemia, and abnormal uterine bleeding) were managed by dose modification or holding T-DM1 and transfusion if needed.

## Tolerability

The median number of cycles of treatment in patients who completed at least 1 cycle ( $n=14$ ) was 11 (3–19). Three patients in cohort 1 and 5 patients in cohort – 1 required dose reductions in apelisib for toxicity. Toxicities requiring dose reductions were rash, hyperglycemia, anorexia/weight loss, and prolonged QTC (Table 2). At the final analysis, all patients had discontinued the drug. Of the 17 patients who received at least 1 dose of the drug, treatment was discontinued due to progression of disease in 14 patients and difficulty complying with study procedures and follow-up in 2 patients. One patient was taken off the study due to recurrent hyperglycemia after non-adherence to recommended interventions.

## Clinical activity

Fourteen patients were evaluable for response (Table 3). Three patients received less than 1 cycle and were not evaluable for response. Two of these patients discontinued treatment due to difficulty complying with study requirements and follow-up visits and one patient died of progressive disease after taking only one dose of the study medication. The ORR was 43%. Exploratory analysis found that clinical benefit rate [CBR; complete response (CR) + partial response (PR) + > 6 mo stable disease (SD)] was experienced by 10 patients (71%).

In the subgroup of 10 patients with prior progression on T-DM1, the ORR was 30% and the CBR was 60%. Only 2 patients, both with prior TDM1 exposure, had progressive

**Table 4** Efficacy in patients evaluable for response

	All patients ( <i>n</i> = 14)	Without prior TDM1 exposure ( <i>n</i> = 4)	With prior TDM1 exposure ( <i>n</i> = 10)
Best response, <i>n</i> (%)			
ORR	6 (43%)	3 (75%)	3 (30%)
CR	1 (7%)	1 (25%)	0
PR	5 (36%)	2 (50%)	3 (30%)
SD	6 (43%)	1 (25%)	5 (50%)
PD	2 (14%)	0	2 (20%)
Clinical benefit rate (CR + PR + > 6mo SD)	10 (71%)	4 (100%)	6 (60%)
Median number of cycles (range)	11 (3–19)	13.5 (12–19)	8 (3–19)
Median PFS	8.1 mo (3.9–10.8 mo)	10.8 mo (8.1–12.5 mo)	6.2 (1.6–10.5 mo)

disease as their best response, of which one had asymptomatic progression of CNS metastases with stable systemic disease.

The median follow-up was 11.6 months (0.3–19.5). Among all patients evaluable for response, the median PFS (Fig. 1) was 8.1 months (95% CI 3.9–10.8). In the 10 patients with prior T-DM1, the median PFS was 6.3 months (95% CI 1.6–10.5) and in the 4 patients without prior T-DM1 median PFS was 10.8 months (95% CI 8.1–12.5).

### PI3K pathway aberrations

Tumor samples were available from 7 patients, all of whom had a sample from the primary site and 4 had tissue from a metastatic site (Fig. 2). Immunohistochemistry (IHC) showed PTEN loss in 2 patients and AKT increased expression in 3 patients. Six patients had next generation sequencing (NGS) done of a metastatic tumor sample prior to therapy of which 4 had mutations in *PIK3CA*. In total, 9 of 11 patients that had a tumor sample from which IHC and/or

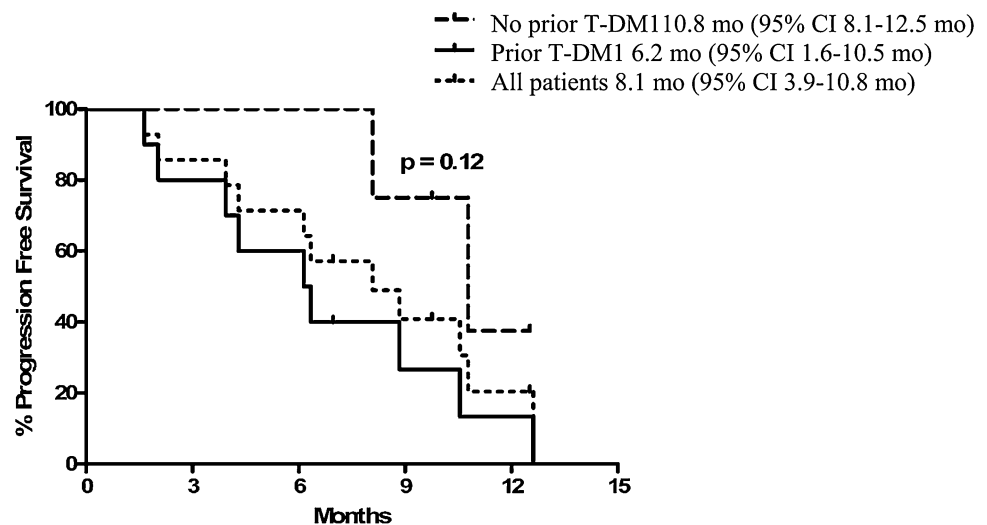
NGS demonstrated aberrations in the PI3K pathway (PTEN loss, AKT overexpression, or *PIK3CA* mutation).

Of patients with an overall response, two had evaluable tissue for IHC assessment with one showing PTEN loss and the other showing AKT overexpression. Of patients with clinical benefit and evaluable samples (*n* = 6), 5 had pathway aberrations. In patients with prior T-DM1 exposure and evaluable tissue (*n* = 8), 6 had pathway aberrations.

### Discussion

The results demonstrate the safety and tolerability of the alpelisib and T-DM1 combination with a DLT of rash and a recommended phase II dose of alpelisib 250 mg daily with standard T-DM1 dosing. The study showed activity of the combination, including a median PFS of 8.1 months, ORR of 43%, and a CBR of 71% despite a heavily pre-treated population. Most impressively, even patients with progression on prior T-DM1 demonstrated response (30%) and clinical benefit (60%) with a median PFS of 6.2 months.

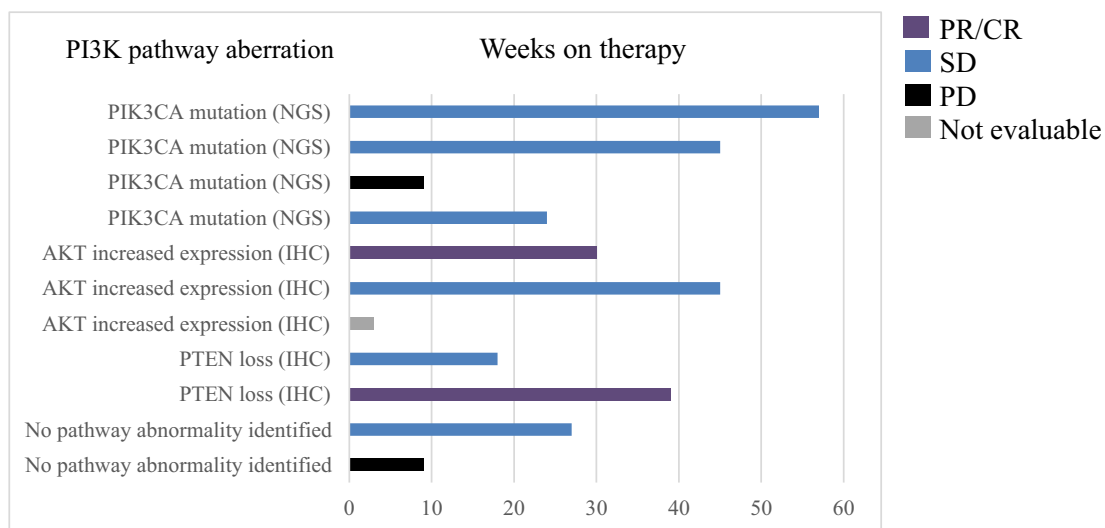
**Fig. 1** Progression-free survival for all patients and by prior T-DM1 exposure





**a** PI3K pathway aberrations by IHC of NGS in patients with evaluable tissue

Patient	PTEN loss (IHC)	AKT increased expression (IHC)	PIK3CA mutation (NGS)	PI3K pathway aberration	Prior T-DM1	Best response
102	Yes	No	-	Yes	No	CR
103	No	No	-	No	Yes	PD
105	No	Yes	-	Yes	No	NE
106	No	Yes	No	Yes	No	SD (>6mo)
201	No	No	Yes	Yes	Yes	SD
202	Yes	No	-	Yes	Yes	SD
203	No	Yes	-	Yes	Yes	PR
204	-	-	Yes	Yes	Yes	PD
205	-	-	Yes	Yes	Yes	SD (>6mo)
208	-	-	Yes	Yes	Yes	SD (>6mo)
210	-	-	No	No	Yes	SD (>6mo)

**b** Duration of response in patients with evaluable tissue

**Fig. 2** Aberrations in PI3K pathway and response. **a** PI3K pathway aberrations by IHC of NGS in patients with evaluable tissue. **b** Duration of response in patients with evaluable tissue

In this study although toxicities were common (59% of patients experienced  $\geq$  grade 3 toxicity), they were manageable, with evaluable patients receiving a median of 11 cycles of therapy. Only one patient was taken off study due to recurrent hyperglycemia; however, the patient did not comply with treatment recommendations. The alpelisib-related DLTs of a maculopapular rash occurred in 7 patients at both the 250 and 300 mg dose levels and hyperglycemia occurred in 1 patient. These toxicities were managed with additional medication, dose interruption, and/or dose reduction as described above, and did not preclude further tolerance of or receiving subsequent benefit from the alpelisib and T-DM1 combination. The consistent timing and presentation of the maculopapular rash, may allow for implementation of

strategies to mitigate it and improve alpelisib tolerability in future studies.

The study population experienced frequent but mild gastrointestinal toxicities (anorexia, weight loss, nausea, transaminase elevation) and fatigue that was likely due to both alpelisib and T-DM1 and did not result in medication discontinuation. One patient developed grade 3 pancreatitis which has not been previously reported with PI3K inhibitors, but has been reported with T-DM1 [34]. Although hematologic toxicities, particularly thrombocytopenia, are expected with T-DM1 given it contains a cytotoxic agent, the prevalence was higher with this combination than described with T-DM1 monotherapy [9, 30]. This may be related to evaluation in a heavily pre-treated population and is less likely

related to alpelisib given that these hematologic toxicities are not common with alpelisib monotherapy [30].

Alpelisib when given with letrozole was tolerated at a higher dose (MTD alpelisib 300 mg daily) with toxicities that included rash (all grades 42%,  $\geq$  grade 3 8%), hyperglycemia (all grades 62%,  $\geq$  grade 3 15%), nausea (all grades 62%,  $\geq$  grade 3 4%), fatigue (all grades 54%,  $\geq$  grade 3 0%), and transaminase elevation (all grades 19%,  $\geq$  grade 3 4%) [30, 31]. Similar toxicities are noted with other PI3K inhibitors, with pan-PI3K also associated with depression and anxiety [35–40]. Toxicities of single-agent T-DM1 include thrombocytopenia (all grade 29%,  $\geq$  grade 3 10%), increased AST (all grades 23%,  $\geq$  grade 3 4%), fatigue (all grade 45%,  $\geq$  grade 3 3%) [41].

The combination of a PI3K inhibitor and TDM1, when compared with the studies described above, may have some overlapping low-grade toxicities resulting in higher rates of gastrointestinal toxicity and fatigue than each medication alone. Only one other trial that has reported the use of alpelisib with a cytotoxic agent (paclitaxel), but toxicity prevented further evaluation of this combination [42]. Studies of the pan-PI3K inhibitors buparlisib and pictilisib with chemotherapy showed increased toxicities requiring dose modifications [43–45]. It is difficult to assess whether the increased specificity of alpelisib led to a more favorable side effect profile than pan-PI3K inhibitors given its combination with a cytotoxic agent and evaluation in a heavily pre-treated population [35–40, 46, 47]. An ongoing phase III study of alpelisib with fulvestrant in HR-positive HER2-negative patients (SOLAR-1) may provide additional insight into the incidence and severity of alpelisib toxicities and how they compare to those of pan-PI3K inhibitors [47, 48].

T-DM1 is the preferred standard therapy after progression on trastuzumab given its superior PFS, OS, and toxicity profile [9]. The safety or efficacy of combining T-DM1 with inhibitors of the PI3K pathway has not been previously reported. This study supports further investigation of combination T-DM1 and alpelisib in extending or enhancing benefit from T-DM1 alone. It is unclear whether the activity demonstrated in this study is primarily from alpelisib or from alpelisib resensitizing the tumor to HER2-targeting with T-DM1 by inhibiting a resistance pathway. Although resistance pathways for T-DM1 are not as well described as those for trastuzumab, their shared binding to HER2 suggests that activation of the downstream PI3K pathway is a possible resistance mechanism to T-DM1 [49].

Inhibitors of the PI3K/AKT/mTOR pathway are currently only investigational in HER2-positive MBC. Adding everolimus (mTOR inhibitor) to trastuzumab and chemotherapy in trastuzumab- and taxane-refractory disease showed modest improvement in PFS in the BOLERO-3 trial, however no benefit was seen in the first-line setting in BOLERO-1 [26, 27]. Early phase studies evaluating PI3K inhibitors

(buparlisib and pictilisib) with trastuzumab in HER2-positive disease have suggested activity, safety, and tolerability [24, 25, 35, 36].

The toxicities of PI3K pathway inhibition and the development of other active therapies for HER2-positive disease highlight the importance of identifying biomarkers that might select for patients likely to derive the greatest benefit from therapy. In two large phase III studies of inhibitors of the PI3K pathway in HER2-negative MBC, patients with PI3K pathway aberrations had improved treatment responses [47, 50]. Additionally, subgroup analysis from BOLERO-1 and –3 suggested only patients with a hyperactive PI3K pathway derived benefit from the addition of everolimus [20]. Our limited correlative studies show that this trastuzumab-resistant population is enhanced for PI3K pathway activation. However, the limited patient samples collected cannot provide definite value in predicting response to alpelisib and T-DM1 therapy. Rigorous evaluation of the PI3K pathway should be an important component of future studies involving PI3K inhibitors.

These are several limitations in this study. A small sample size and single-arm design make it difficult to determine the true impact of clinical or pathologic features on the likelihood of benefit from the combination. Additionally, subgroup analysis of T-DM1 exposed and non-exposed cohorts and calculation of clinical benefit rate was not pre-specified. There was no standardized premedications or a specific protocol for management of the alpelisib-associated rash. If prophylactic measures are initiated in future studies, it potentially could mitigate the DLT. Tissue for correlative studies was only available from a subset of patients. Finally, samples for these correlatives were from archival tissue obtained at variable times during a patient's course so we do not know if patients had PI3K pathway activation at the time of treatment. Future studies may include biopsy or circulating tumor DNA prior to initiating therapy to more consistently evaluate PI3K pathway abnormalities as a biomarker.

This study provides important evidence for the safety, tolerability, and preliminary efficacy of alpelisib and T-DM1 in trastuzumab-refractory HER2-positive MBC patients. The activity of this combination in trastuzumab- and T-DM1-refractory patients supports further study of the combination with proactive strategies for managing toxicities in phase II studies and highlights the potential of PI3K inhibition in T-DM1-resistant disease.

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## Compliance with ethical standards

**Conflict of interest** Dr. Cristofanilli is a consultant for Novartis, received funding and remuneration from Pfizer. Dr. Santa-Maria received research funding from Medimmune and Pfizer. The remaining authors declare they have no conflicts of interest.



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