

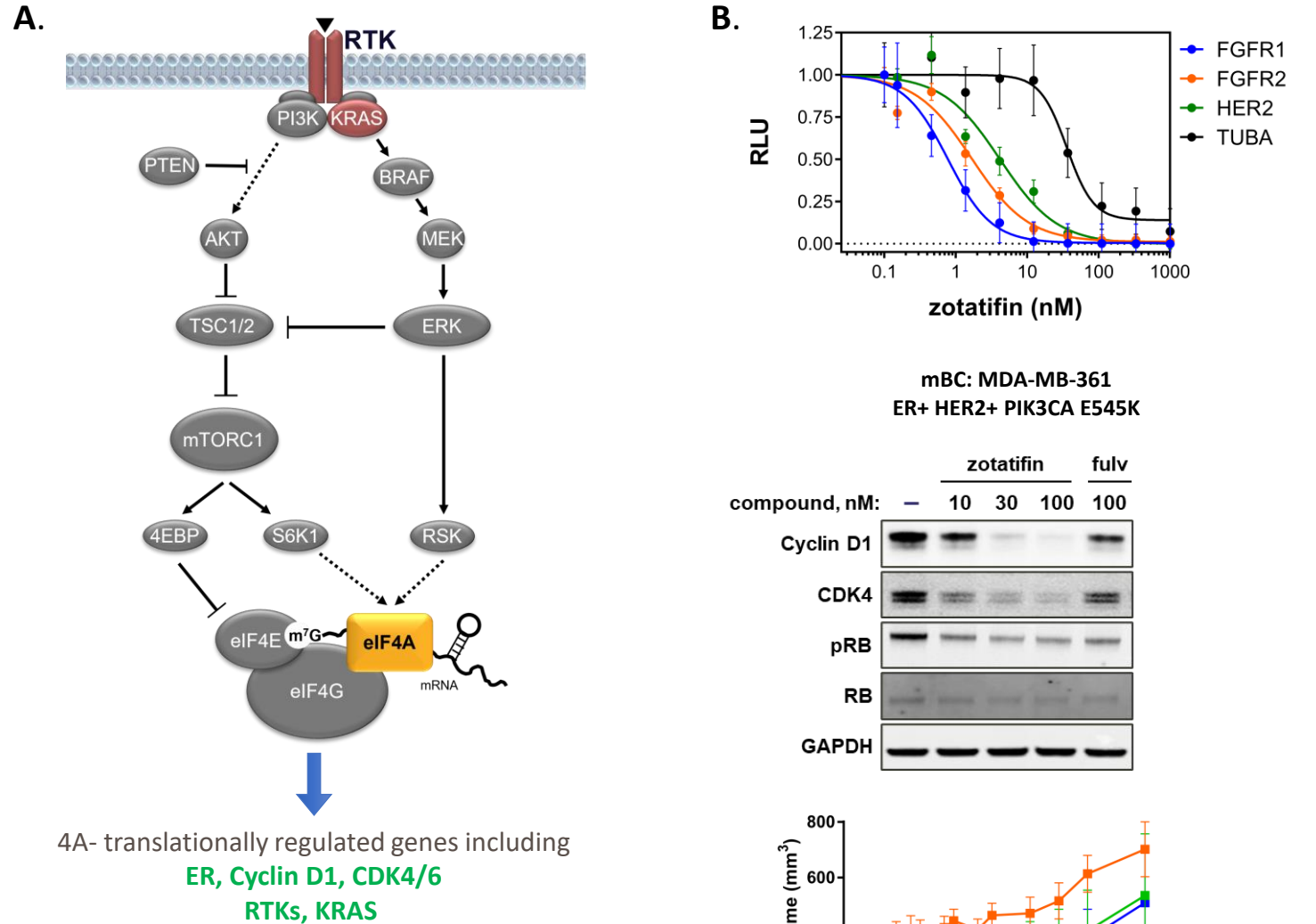
Funda Meric-Bernstam, MD<sup>1</sup>, Manish Sharma, MD<sup>2</sup>, David Sommerhalder, MD<sup>3</sup>, Roland T. Skeel, MD<sup>4</sup>, Anthony B. El-Khoueiry, MD<sup>5</sup>, Jennifer L. Caswell-Jin, MD<sup>6</sup>, Georgina Fulgar, BS<sup>7</sup>, Mark Densel, BS<sup>7</sup>, Samuel Sperry, PhD<sup>7</sup>, Peggy Thompson, PhD<sup>7</sup>, Gary G. Chiang, PhD<sup>7</sup>, Robert Sikorski, MD PhD<sup>7</sup>, Premal Patel, MD PhD<sup>7</sup>, Ezra Rosen, MD PhD<sup>8</sup>

<sup>1</sup> University of Texas MD Anderson Cancer Center, TX <sup>2</sup> START Midwest, MI <sup>3</sup> NEX T Oncology, San Antonio, TX <sup>4</sup> University of Toledo Medical Center / Dana Cancer Center, OH <sup>5</sup> USC Norris Comprehensive Cancer Center, CA <sup>6</sup> Stanford University, CA <sup>7</sup> EFFECTOR Therapeutics, CA <sup>8</sup> Memorial Sloan Kettering Cancer Center, NY

## Background

Zotatifin (eFT226) is a selective inhibitor of eukaryotic translation initiation factor 4A (eIF4A)-mediated translation with a novel mechanism of action. Zotatifin binds selectively to defined nucleotide sequences present in the 5' untranslated region (5'-UTR) of a subset of messenger ribonucleic acids (mRNAs) and concurrently to eIF4A. Zotatifin target sequences are found in the 5'-UTR of the estrogen receptor, Cyclin D1, CDK4, as well as several oncogenes and drivers of cell proliferation (HER2, ERBB3, FGFR1/2, EGFR, and KRAS). Zotatifin selectively blocks translation of these important mRNAs. Tumors with elevated protein levels, activating mutations or gene amplifications in zotatifin-sensitive proteins demonstrate selective downregulation of oncoprotein expression, inhibition of tumor cell proliferation and induction of apoptosis following zotatifin treatment. This mechanism of action results in significant in vivo tumor growth inhibition in multiple tumor models, including models of breast cancer (ER+, HER2 amplified, or FGFR1 overexpressed), non-small cell lung cancer (FGFR1 amplified or KRAS mutant), and colorectal cancer (KRAS mutant).

Collectively, these nonclinical data provided the foundation for initiation of Phase 1 clinical development of zotatifin. The initial development during dose escalation was in a mixed population of tumors with a molecular alteration (an activating mutation, amplification, or fusion) in HER2, ERBB3, FGFR1, FGFR2, EGFR, or KRAS and in cancers with very high prevalence of mutations in 1 of these genes, such as KRAS in pancreatic adenocarcinoma. Based on additional preclinical data and early clinical activity, Phase 2 expansion cohorts are now primarily focused on ER+ breast cancer.



**Figure 1. A. eIF4A is required for protein translation initiation and tumorigenesis. A.** Oncogenic signaling through the RAS and PI3K pathways leads to activation of RNA helicase eIF4A. Activated eIF4A is required for efficient translation of mRNA encoding downstream effector proteins including ER, Cyclin D1, CDK4/6 as well as upstream pathway components including RTKs and KRAS. **B. Pre-clinical efficacy of zotatifin.** Top panel, zotatifin selectively inhibits the translation of eIF4A-dependent mRNAs with nanomolar potency in a cell-based reporter assay. Middle panel, zotatifin selectively inhibits production of cell-cycle proteins including Cyclin D1 and CDK4 in MDA-MB-361 (ER+ HER2+ PIK3CA E545K) cells. Bottom panel, zotatifin inhibits tumor growth in mice bearing MDA-MB-361 xenografts and leads to sustained tumor regression post dosing in combination with palbociclib.

## Objectives

- Primary objectives:**
- Part 1 (Dose escalation)**
    - To define the safety and tolerability of zotatifin as monotherapy in patients with defined, advanced solid tumors
    - To determine the MTD or RP2D for zotatifin as monotherapy
    - To evaluate the PK profile of zotatifin
  - Part 2 (Expansion cohorts)**
    - To evaluate the preliminary antitumor activity of zotatifin as monotherapy and as combination therapy in patients with defined, advanced solid tumors
    - To determine the MTD or RP2D for zotatifin as combination therapy
- Secondary objectives:**
- Part 1 (Dose escalation)**
    - To evaluate the preliminary antitumor activity of zotatifin as monotherapy in patients with defined, advanced solid tumors
  - Part 2 (Expansion cohorts)**
    - To assess the safety of zotatifin as monotherapy and as combination therapy
    - To assess progression free survival (PFS)
    - To evaluate the PK profile of zotatifin in combination therapy
- Exploratory objectives:**
- To explore the effects of zotatifin on pharmacodynamic markers relating to drug mechanism
  - To explore additional biomarkers to further elucidate mechanism of action, predict response to therapy, and understand potential resistance mechanisms

## Methods

**Study design and treatment**

- Open label study in adult patients with advanced solid tumors
- Part 1:** 3+3 dose escalation scheme with weekly intravenous administration of zotatifin in a 21 day cycle
- Part 2:** Expansion at MTD or RP2D as monotherapy or in combination with SOC therapy in tumor types harboring molecular changes that may make them more sensitive to zotatifin therapy as a Simon 2-stage design, with seven patients in stage 1 of each cohort

**Patient eligibility - key inclusion criteria**

Part 1 – Dose escalation	Part 2 – Expansion
<ul style="list-style-type: none"> <li>PDAC: no molecular typing</li> <li>HER2: overexpression</li> <li>HER2, ERBB3, FGFR1, FGFR2, EGFR: Activating mutation, amplification or fusion</li> <li>KRAS: activating mutation</li> </ul>	<ul style="list-style-type: none"> <li>EMBF: ER+ BC, FGFR amp zotatifin monotherapy</li> <li>ECBF: ER+ BC zotatifin + fulvestrant</li> <li>ECBF+ A: ER+ HER2- zotatifin + fulvestrant + abemaciclib</li> <li>ECNS: NSCLC, KRAS G12C zotatifin + sotorasib</li> </ul>
<ul style="list-style-type: none"> <li>Solid tumors progressive after SOC or no potential for cure</li> </ul>	<ul style="list-style-type: none"> <li>Post-endocrine therapy No limit to prior lines</li> <li>Post-endocrine and CDK4/s therapy No limit to prior lines</li> <li>Post-endocrine therapy No limit to prior lines</li> <li>Post-chemo/I/O therapy No prior KRAS-targeted therapy</li> </ul>

**Maximum tolerated dose (MTD) and RP2D**

- MTD: Highest dose level at which  $\geq 6$  patients have been treated and associated with a first cycle DLT rate of  $< 33\%$
- RP2D: May be the MTD or may be a lower dose within the tolerable dose range

**Dose limiting toxicity (DLT)**

- Graded according to NCI CTCAE v5.0 during the 21-day DLT evaluation period and not clearly related to disease progression

Non-hematologic	Hematological
<ul style="list-style-type: none"> <li>Any Grade 2-4 toxicity</li> <li>Grade 3 nausea/vomiting lasting <math>&gt; 48</math> h or any occurrence of Grade 4</li> <li>Grade 3 diarrhea lasting <math>&gt; 48</math> h or any occurrence of Grade 4</li> <li>Grade <math>\geq 3</math> elevation of serum ALT or AST for <math>&gt; 7</math> days or in conjunction with Grade 2 elevation in serum bilirubin</li> </ul>	<ul style="list-style-type: none"> <li>Any Grade 5 toxicity</li> <li>Grade 4 neutropenia lasting <math>&gt; 5</math> days or any febrile neutropenia</li> <li>Grade 4 anemia unexplained by underlying disease</li> <li>Grade 3 thrombocytopenia with bleeding or any Grade 4</li> </ul>

**Pharmacokinetic assessment**

- Non-compartmental methods to assess AUC, CL, Vss, T1/2

**Response assessment**

- Radiological tumor assessment at baseline and every 8 weeks
- Response assessment using RECIST 1.1

**All data reported is interim prior to database lock (data cut-off 3/04/2022)**

## Results

**Table 1. Patient demographic and baseline characteristics**

Characteristic	Part 1 (N=37)	Part 2 (N=17)
Age, median (range), years	62 (36-80)	55 (37-81)
Gender		
• Male	18	0
• Female	19	17
Race		
• White	30	13
• Black or African descent	1	4
• Asian	2	0
• Other	4	
Median number of prior metastatic therapies (range)	3 (0-9)	4 (1-11)
Primary diagnosis/cancer type, n (%)		
• Colorectal	11 (29.7)	
• Pancreatic	6 (16.2)	
• NSCLC	6 (16.2)	1 (5.9)
• Breast	4 (10.8)	16 (94.1)
• Cholangiocarcinoma	2 (5.4)	
• Melanoma	2 (5.4)	
• Other (1 patient/tumor type*)	6 (16.2)	

\*appendiceal, ovarian, sarcoma, small bowel, testicular, unknown

**Table 2. Patient disposition**

	Part 1, N=37 (%)	Part 2, N=17 (%)
Continuing on study	0 (100)	8 (47.1)
Discontinued therapy		
• Disease progression	31 (83.8)	8 (47.1)
• Adverse event	2 (5.4)	1 (5.9)
• Withdrawal of consent	1 (2.7)	0 (0.0)
• Investigator or sponsor decision	2 (5.4)	0 (0.0)
• Other	1 (2.7)	0 (0.0)

## Results (continued)

**Table 3. Safety Summary**

Category	zotatifin, mg/kg									
	Part 1, N (%)							Part 2, N (%)		
Subjects with TEAEs	3 (100)	3 (100)	2 (66.7)	7 (100)	3 (100)	3 (100)	8 (100)	7 (100)	15 (88.1)	
Subjects with TEAEs related to zotatifin <sup>a</sup>	1 (33.3)	3 (100)	2 (66.7)	7 (100)	2 (66.7)	2 (66.7)	7 (87.5)	6 (85.7)	9 (52.9)	
Subjects with DLTs	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	0 (0.0)	0 (0.0)	0 (0.0)	2 (28.6)	1 (5.9)	
Subjects with TSEAEs	1 (33.3)	0 (0.0)	0 (0.0)	3 (42.9)	0 (0.0)	0 (0.0)	1 (12.5)	3 (42.9)	3 (17.6)	
Subjects with TSEAEs related to zotatifin <sup>c</sup>	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	0 (0.0)	0 (0.0)	0 (0.0)	2 (28.6)	0 (0.0)	
Subjects with CTCAE Gr 3/4 TEAEs	2 (66.7)	0 (0.0)	0 (0.0)	5 (71.4)	0 (0.0)	0 (0.0)	1 (33.3)	2 (25.0)	4 (23.5)	
Subjects with CTCAE Gr 5 TEAEs	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	
Subjects with CTCAE Gr 3/4 TEAEs related to zotatifin <sup>c</sup>	0 (0.0)	0 (0.0)	0 (0.0)	5 (71.4)	0 (0.0)	0 (0.0)	0 (0.0)	2 (28.6)	0 (0.0)	
Subjects with CTCAE Gr 5 TEAEs related to zotatifin <sup>c</sup>	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	
Subjects with TEAEs leading to dose interruption or reduction of zotatifin	1 (33.3)	0 (0.0)	0 (0.0)	4 (57.1)	0 (0.0)	0 (0.0)	1 (12.5)	0 (0.0)	1 (5.9)	
Subjects with TEAEs leading to discontinuation of zotatifin	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	0 (0.0)	

After 7 patients had been dosed at 0.035 mg/kg weekly, dosing frequency was reduced to days 1 and 8 of a 21 day cycle to mitigate anemia TEAE, Treatment-Emergent Adverse Event, DLT. Dose Limiting Toxicity; CTCAE, Common Terminology Criteria for Adverse Events (version 5.0). Treatment-emergent adverse events (TEAEs) are AEs that start during or after zotatifin therapy, or AEs with an onset prior to initiating eFT226 therapy that worsen after zotatifin initiation.

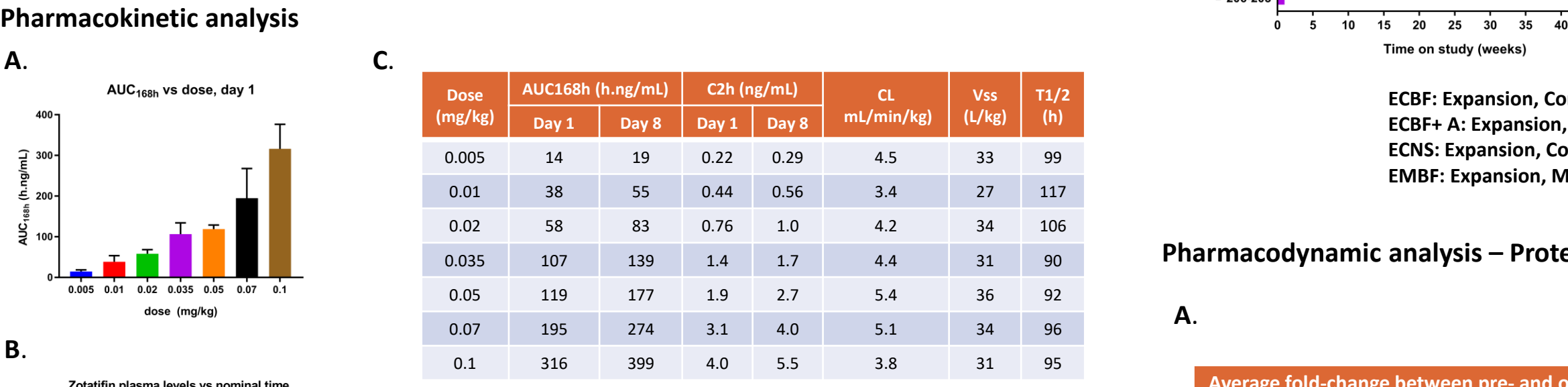
**Table 4. Treatment emergent adverse events (TEAEs) related to zotatifin**

MedDRA term	Grade	zotatifin, mg/kg									
		Part 1, N (%)							Part 2, N (%)		
Fatigue	1-2	1 (33.3)	1 (33.3)	0 (0.0)	4 (57.1)	1 (33.3)	1 (33.3)	1 (12.5)	2 (28.6)	1 (5.9)	
Anemia	1-2	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	
Diarrhea	1-2	1 (33.3)	0 (0.0)	0 (0.0)	3 (42.9)	0 (0.0)	0 (0.0)	3 (37.5)	0 (0.0)	3 (17.6)	
Vomiting	1-2	0 (0.0)	1 (33.3)	0 (0.0)	1 (14.3)	0 (0.0)	0 (0.0)	2 (25.0)	2 (28.6)	3 (17.6)	
Nausea	1-2	0 (0.0)	1 (33.3)	0 (0.0)	1 (0.0)	0 (0.0)	0 (0.0)	1 (12.5)	1 (14.3)	2 (11.7)	

**Table 4. Treatment emergent adverse events (TEAEs) related to zotatifin** (continued)

MedDRA term	Grade	zotatifin, mg/kg									
		Part 1, N (%)							Part 2, N (%)		
Fatigue	3-4	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	
Anemia	3-4	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	1 (14.3)	0 (0.0)	
Diarrhea	3-4	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	
Vomiting	3-4	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	
Nausea	3-4	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	0 (0.0)	

Events listed are  $>10\%$  incidence. Coded with Medical Dictionary for Regulatory Activities (MedDRA) Version 25.0. Percentage is calculated using the number of treated subjects as the denominator. Treatment-emergent adverse events (TEAEs) are AEs that start during or after zotatifin therapy, or AEs with an onset prior to initiating eFT226 therapy that worsen after zotatifin initiation.



**Table 3. Pharmacodynamic analysis - RNA stabilization**

**A. Schematic:** eIF4A1, zotatifin, and target mRNA form a ternary complex, stabilizing the mRNA.

**B. Plot:** Average fold-change from baseline for c-JUN mRNA at various time points (0, 4, 8, 16, 24h) following zotatifin administration (0.005 to 0.1 mg/kg).

**Figure 3. Pharmacodynamic analysis.** Schematic of zotatifin mechanism of action and RPPA analysis results.

**A. Schematic:** eIF4A1, zotatifin, and target mRNA form a ternary complex, stabilizing the mRNA.

**B. RPPA analysis:** Average fold-change from baseline for c-JUN mRNA plotted for each time point.

**Table 5. Pharmacodynamic analysis - Proteins/phosphoproteins**

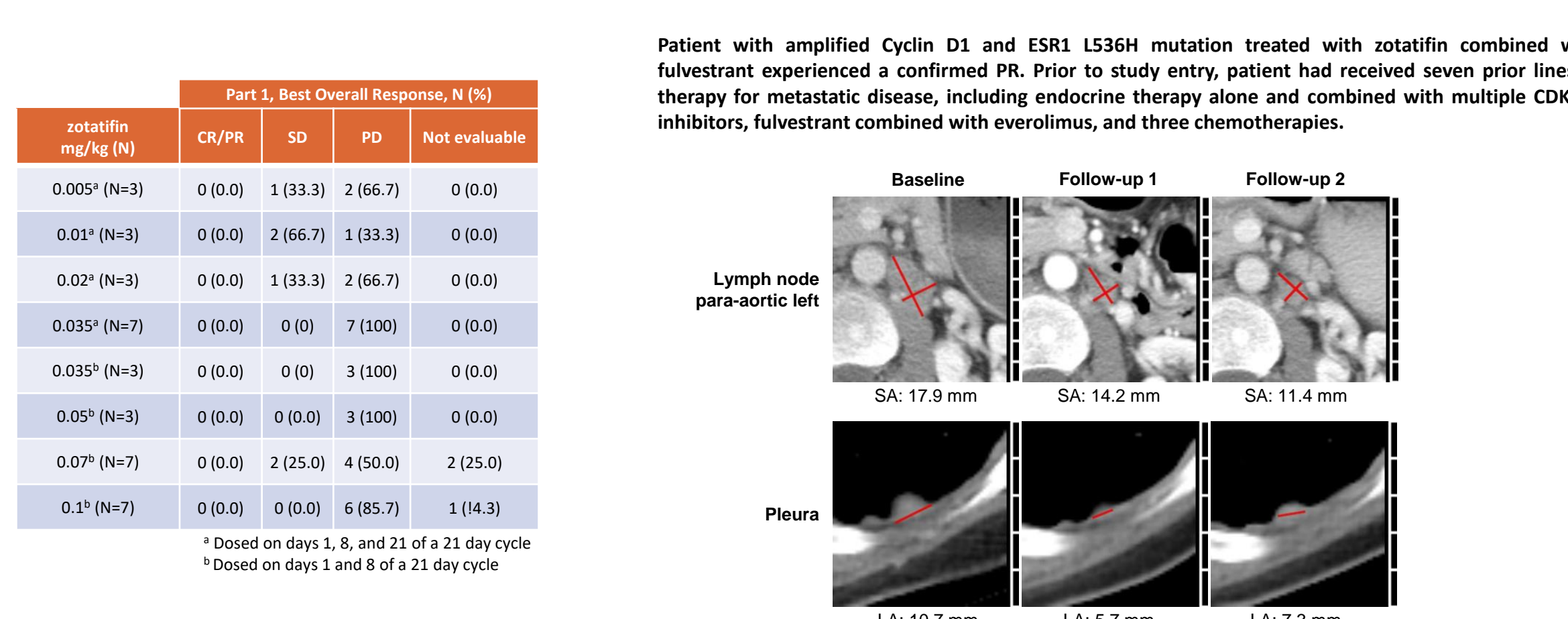
**A. Average fold-change between pre- and on-treatment:**

	Proteins (~389)	Phospho-protein markers (~95)
10 subjects	0.99	0.95
7 BC subjects	0.99	0.91

**B. Before-after plots:** RPPA score and % change from baseline for Cyclin E1, Bcl-2, ATM pS1981, p38 MAPK p180Y182, and PRAS40 pT246 in mBC subjects.

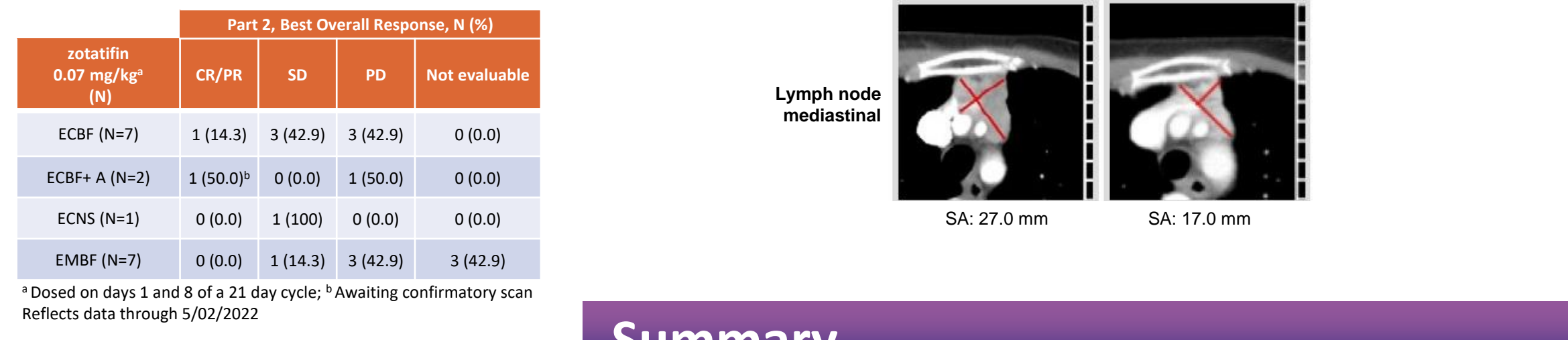
**Figure 3. RPPA analysis.** Paired frozen biopsies from pre-treatment and on-treatment (N=10, including 7 breast cancer patients) were analyzed by reverse phase protein array (RPPA core, MD Anderson) for ~488 protein markers (total or phospho-proteins).

## Efficacy analysis - Part 1, dose escalation in mixed tumors



- During Part 1 dose escalation, no PRs or CRs were observed
- The longest stable disease (23 weeks) was observed in a subject with FGFR1-amplified breast cancer treated at the RP2D who had received three prior lines of treatment for metastatic disease, including palbociclib/anastrozole followed by fulvestrant

## Interim efficacy analysis - Part 2, defined expansion cohorts



## Interim efficacy analysis - Part 2, expansion cohorts (continued)

- Target engagement was demonstrated via stabilization of c-JUN mRNA circulating in blood
- Modulation of translation was shown to be highly selective with less than 1% of protein expression altered
- Reductions in key oncogenic drivers such as Cyclin E and Bcl-2 were observed with the most dramatic effects in patients with the highest pre-treatment signals

## Pharmacodynamic analysis - Proteins/phosphoproteins

**Figure 4. RPPA analysis.** Paired frozen biopsies from pre-treatment and on-treatment (N=10, including 7 breast cancer patients) were analyzed by reverse phase protein array (RPPA core, MD Anderson) for ~488 protein markers (total or phospho-proteins).

**A. Waterfall plot:** Average fold-change between pre- and on-treatment for 10 subjects and 7 BC subjects. Proteins that deviated  $> 2$  standard deviations above or below the mean in breast cancer (BC) subjects are represented as averaged log<sub>2</sub> fold change on the waterfall plot.

**B. Before-after plots:** RPPA score and % change from baseline for selected protein and phospho-protein markers represented as RPPA score (top panels) and % change from baseline (bottom panels).

## Demonstration of a safely tolerated dose/schedule and initial efficacy signals in targeted patients

- Zotatifin was well tolerated at the RP2D with no related Gr 3/4/5 TEAEs in 25 patients
- A confirmed PR was observed in combination with fulvestrant in a patient with amplified Cyclin D1 and an ESR1 mutation whose disease had previously progressed on fulvestrant combined with everolimus
- A not yet confirmed PR was observed in combination with fulvestrant and abemaciclib in a patient with PIK3CA mutations whose disease had progressed on fulvestrant combined with alpelisib and a CDK4/6 inhibitor combined with endocrine therapy

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