

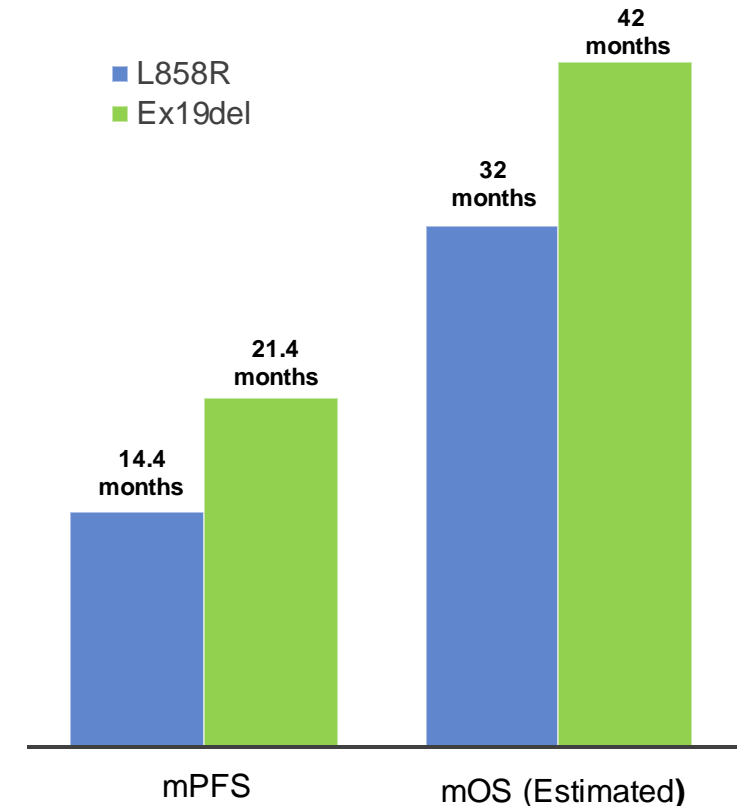
BLU-945 monotherapy and in combination with osimertinib in previously treated patients with advanced *EGFR*-mutant NSCLC in the phase 1/2 SYMPHONY study

Yasir Elamin, MD,¹ Misako Nagasaka, MD, PhD,² Elaine Shum, MD,³ Lyudmila Bazhenova, MD,⁴ D. Ross Camidge, MD, PhD,⁵ Byoung Chul Cho, MD, PhD,⁶ Enriqueta Felip, MD, PhD,⁷ Koichi Goto, MD, PhD,⁸ Chia-Chi Lin, MD, PhD,⁹ Zofia Piotrowska, MD,¹⁰ David Planchard, MD, PhD,¹¹ Julia Rotow, MD,¹² David R. Spigel, MD,¹³ Daniel S. W. Tan, MD, PhD,¹⁴ Tatsuya Yoshida, MD, PhD,¹⁵ Anna Minchom, MD,¹⁶ Adrianus Johannes de Langen, MD,¹⁷ Terufumi Kato, MD,¹⁸ Alena Zalutskaya, MD, PhD,¹⁹ Karen L. Reckamp, MD²⁰

Introduction

- In EGFRm-positive NSCLC, on-target and off-target treatment resistance eventually develops with third generation EGFR TKIs, presenting a patient population that is challenging to treat
- Optimization of EGFR pathway inhibition with combination treatment is likely to be more successful in front-line due to patient genomic homogeneity
- While EGFR TKI-TKI combinations have been explored, a high rate of EGFR WT toxicity has limited their clinical utility¹
- **BLU-945 is an investigational, next-generation, oral TKI uniquely selective against EGFR WT:**
 - nM potency on EGFR-activating (L858R, ex19del) and T790M and C797X resistance mutations
 - Large EGFR WT window makes BLU-945 a combination partner with reduced risk for unacceptable EGFR WT toxicity

Osimertinib outcomes in first-line treatment of NSCLC^{2,3}



EGFR, epidermal growth factor receptor; ex19del, exon 19 deletion; mOS, median overall survival; mPFS, median progression free survival; NSCLC, non-small cell lung cancer; TKI, tyrosine kinase inhibitor, WT, wild type.

1. Rotow J, et al. *J Clin Oncol.* 2020;38(15):9507-9507. 2. Souria JC, et al. *New Engl J Med.* 2018;378:113-125. 3. Ramalingam SS, et al. *J Thorac Oncol.* 2022;17(9):S67-S68.

SYMPHONY (NCT04862780) study design and patient characteristics

| Key eligibility criteria | Phase 1 (dose escalation) | |
|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------|
| | Part 1A (N=112) BLU-945 monotherapy BOIN design Starting dose: 25 mg QD ^a Initiated May 2021 | Part 1B (N=55) BLU-945 + osimertinib (80 mg) Starting dose: BLU-945 200 mg QD ^a Initiated June 2022 |
| <ul style="list-style-type: none"> Adults with metastatic <i>EGFR</i>m NSCLC No other known oncogenic tumor drivers ECOG status 0-1 Prior treatment with ≥ 1 EGFR TKI with activity against T790M; progression on osimertinib as last therapy (part 1B only) | All combination patients received osimertinib as last line of therapy without a washout period | |
| | Primary endpoints MTD, RP2D, safety | |

| Characteristic | BLU-945 | |
|-----------------------------------|-------------------------------------|------------------------------------|
| | Monotherapy ^b (n=112) | Combination ^c (n=55) |
| Age, years, median (min, max) | 63 (34, 84) | 62 (28, 87) |
| Age group, n (%) | | |
| <65 years | 63 (56.3) | 32 (58.2) |
| ≥ 65 years | 49 (43.8) | 23 (41.8) |
| Female, n (%) | 74 (66.1) | 34 (61.8) |
| CNS metastases at baseline, n (%) | 43 (38.4) | 17 (30.9) |
| Prior LOT, median (min, max) | 3.5 (1, 13) | 2 (1, 7) |

- Patients enrolled in the phase 1 dose escalation were heavily pretreated
- 94% of monotherapy and 89% of combination patients had an additional EGFR and/or detectable additional genetic alteration
- Combination dose escalation is ongoing

^aBID dosing was also evaluated. ^b25–600 mg QD; 100–300 mg BID. ^c200–400 mg QD; 100–200 mg BID with OSI 80 mg QD.
 BID, twice daily; BOIN, Bayesian optimal interval; ECOG PS, Eastern Cooperative Oncology Group performance status; ex19del, exon 19 deletion; LOT, line of therapy; MTD, maximum tolerated dose; QD, every day; RP2D, recommended phase 2 dose; TKI, tyrosine kinase inhibitor.

BLU-945 monotherapy was generally well tolerated

| TRAEs, N=112 | | |
|----------------------------------------------------|-----------|------------------|
| TRAEs, n (%) Safety population | Any grade | Grade $\geq 3^a$ |
| Any TRAE | 86 (76.8) | 37 (33.0) |
| EGFR-related TRAEs (all patients) | | |
| Rash | 11 (9.8) | 0 |
| Diarrhea | 7 (6.3) | 0 |
| Dry skin | 4 (3.6) | 0 |
| Paronychia | 2 (1.8) | 0 |
| TRAEs in $\geq 25\%$ of patients | | |
| ALT | 41 (36.6) | 25 (22.3) |
| Nausea | 38 (33.9) | 3 (2.7) |
| AST | 37 (33.0) | 12 (10.7) |
| Headache | 31 (27.7) | 0 |
| Vomiting | 30 (26.8) | 1 (>1) |

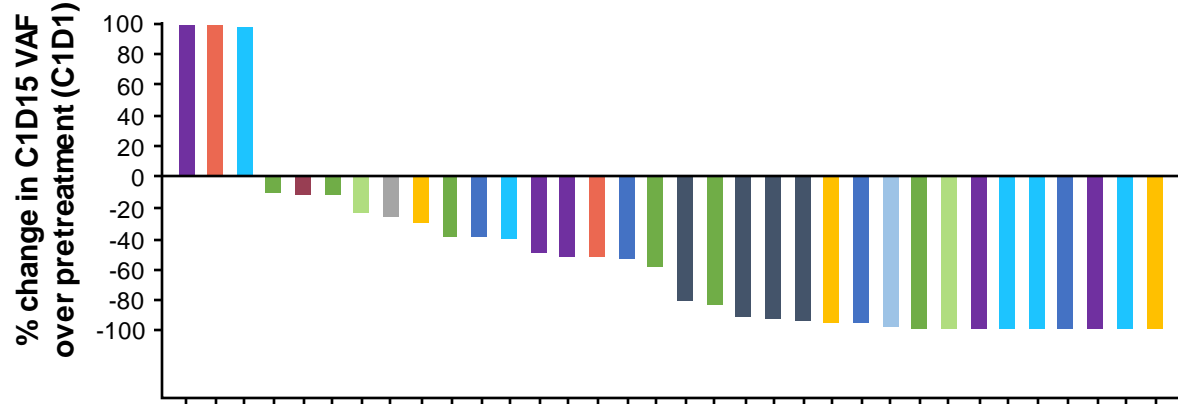
- Majority of TRAEs were low grade (NCI CTCAE Grade 1–2)
- There were 12 patients with DLTs across 400 mg–600 mg total daily doses (QD and BID), with the most common DLTs being Grade 3 ALT and AST elevation
- EGFR-WT associated AEs were low grade and infrequent (<10%)

ALT, alanine aminotransferase; AST, aspartate aminotransferase; BID, twice daily; NCI CTCAE, National Cancer Institute Common Terminology Criteria for Adverse Events; DLT, dose limiting toxicity; EGFR, epidermal growth factor receptor; QD, once daily; TRAE, treatment-related adverse event; WT, wild type.

^aTwo patients (1.8%) experienced Grade 5 AE possibly related to BLU-945 as assessed by an investigator: pneumonitis at 300 mg BID and intracranial bleeding at 100 mg BID in a patient with suspected brain metastases.

BLU-945 monotherapy resulted in dose-dependent reduction and clearance of EGFR T790M and EGFR C797X ctDNA at Cycle 1 Day 15

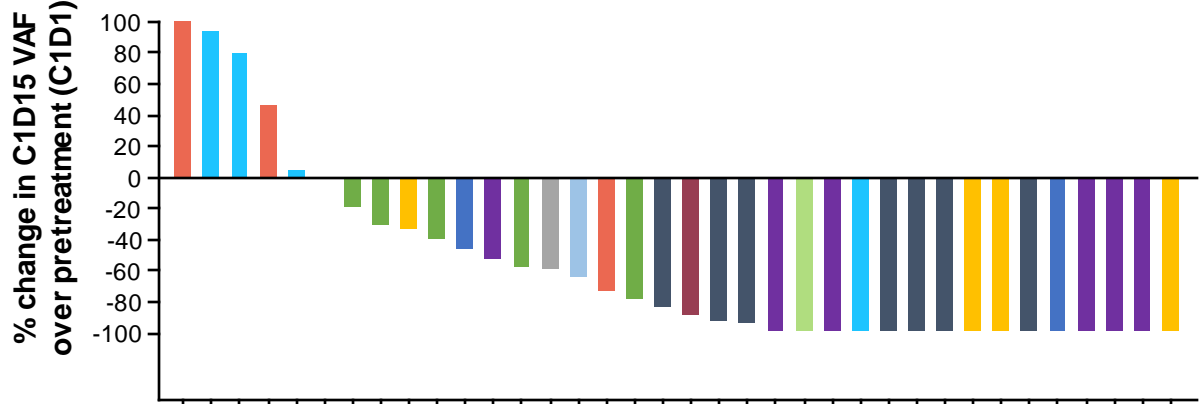
Longitudinal profiling of EGFR T790M mutation allele levels^a



| EGFR mutational profile | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
|-------------------------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| EGFR-ex19del | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | | | |
| EGFR-L858R | X | | | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | | | |
| EGFR-T790M | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X |
| EGFR-C797x | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X |

- 25 mg QD
- 50 mg QD
- 100 mg QD
- 200 mg QD
- 400 mg QD
- 500 mg QD
- 600 mg QD
- 100/100 mg BID
- 200/200 mg BID
- 250/250 mg BID
- 300/300 mg BID

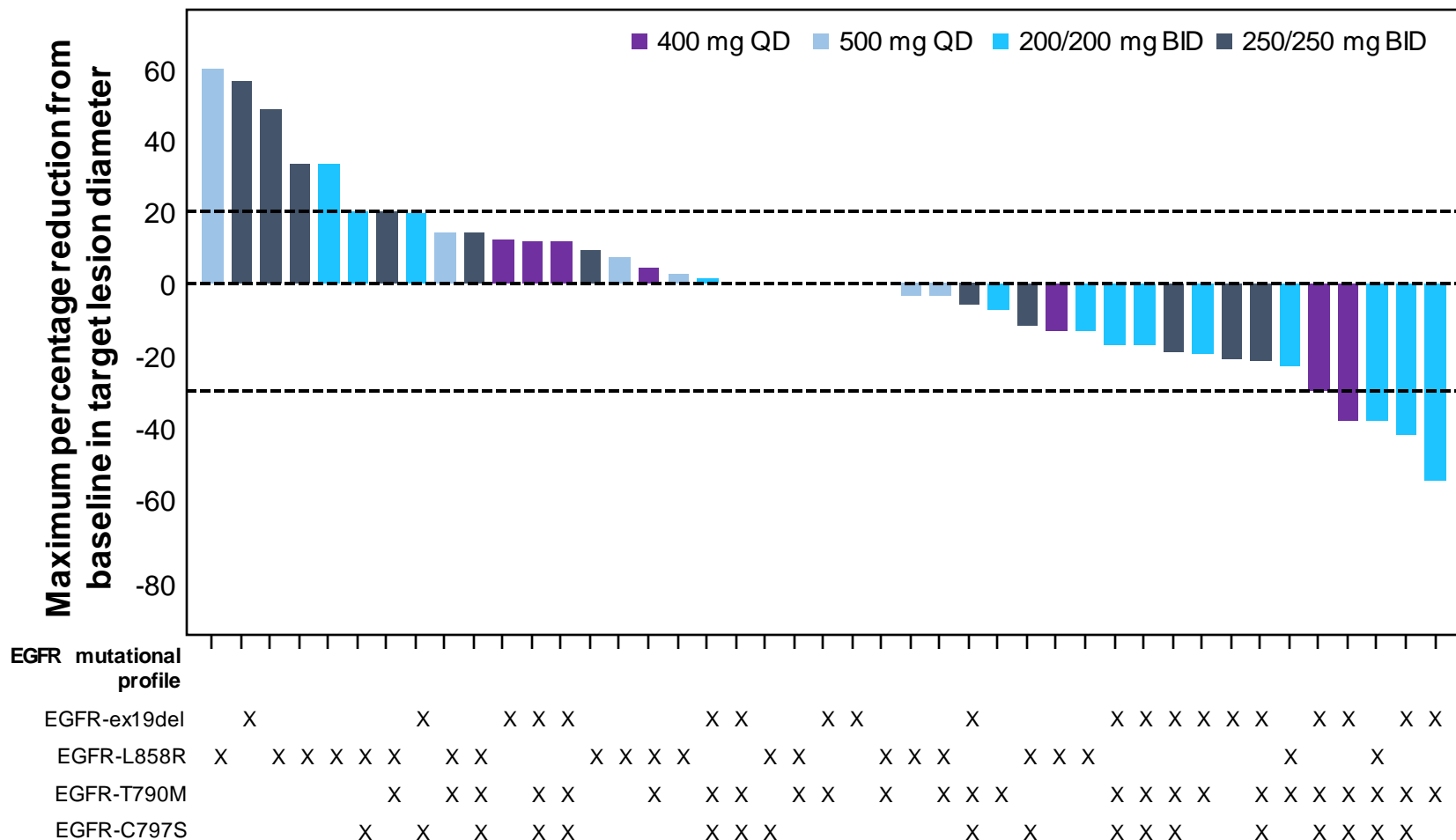
Longitudinal profiling of EGFR C797X mutation allele levels^a



| EGFR mutational profile | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
|-------------------------|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| EGFR-ex19del | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | | | | | | | |
| EGFR-L858R | | | | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | | | | | |
| EGFR-T790M | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | | | |
| EGFR-C797x | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X | X |

^aPercent change greater than 100% are displayed as 100% in the figure. EGFR mutational profile based on results from Foundation One Liquid CDx(F1CDx) baseline (C1D1) analysis
 Note: Patient with multiple mutations for EGFR C797S in the same specimen are shown as a different colored X in the EGFR mutational profile.
 BID, twice daily; EGFR, epidermal growth factor receptor; QD, once daily

BLU-945 monotherapy antitumor activity^a



- Heavily pretreated population resulting in disease heterogeneity
- Tumor reduction and two confirmed partial responses were observed at higher dose levels of BLU-945 monotherapy
- Limited durability of clinical benefit observed, likely due to late-line disease heterogeneity and off-target resistance

^aPatients with EGFR-mutant positive NSCLC were enrolled based on local mutation assessment of tumor biopsy or blood ctDNA (displayed) with a follow-up central ctDNA assessment at C1D1. Patients were counted only once.

BID, twice daily; EGFR, epidermal growth factor receptor; PD, progressive disease; PR, partial response; QD, once daily; SD, stable disease.

BLU-945 + osimertinib combination is well tolerated with limited EGFR WT AEs

| TRAEs, (N=55) | | |
|-----------------------------------|-----------|----------|
| TRAEs, n (%) Safety population | Any grade | Grade ≥3 |
| Any TRAEs | 52 (94.5) | 6 (10.9) |
| EGFR-associated TRAEs | | |
| Diarrhea | 16 (29.1) | 0 |
| Dry skin | 9 (16.4) | 0 |
| Dermatitis acneiform | 8 (14.5) | 1 (1.8) |
| Paronychia | 6 (10.9) | 0 |
| TRAEs in ≥10% of patients | | |
| Headache | 19 (34.5) | 0 |
| Nausea | 19 (34.5) | 0 |
| Fatigue | 12 (21.8) | 1 (1.8) |
| Decreased appetite | 7 (12.7) | 0 |
| Vomiting | 6 (10.9) | 0 |

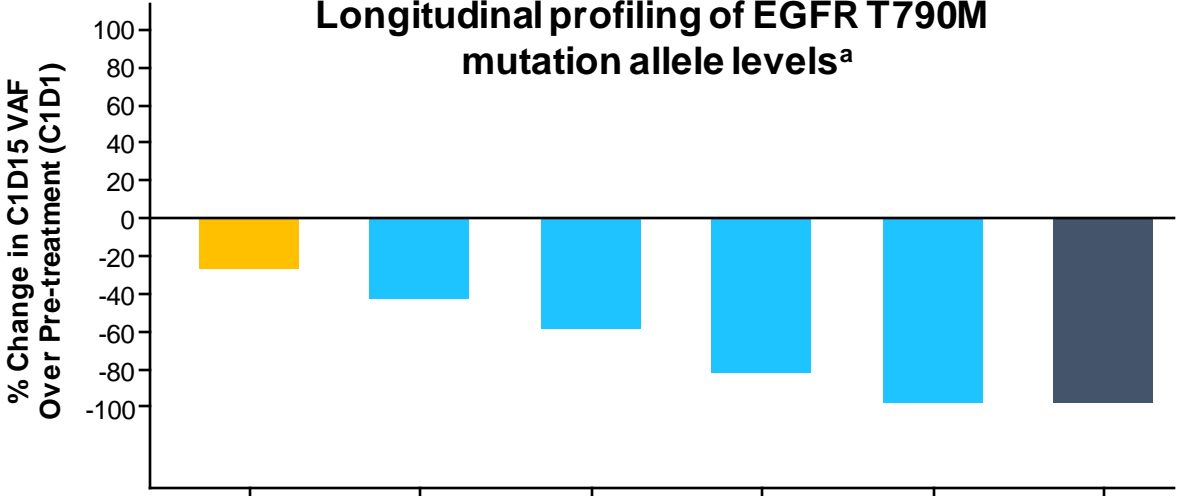
- Exposure of BLU-945 and osimertinib when coadministered are comparable to PK data from BLU-945 given alone and published osimertinib data^{1,2}
- EGFR-WT associated AEs were infrequent, and the majority were Grade 1
- Three patients had DLTs across 200 – 400 mg total daily doses
 - 100 mg BID + 80 mg osi, Grade 3 acute respiratory failure
 - 300 mg QD + 80 mg osi - Grade 4 pneumonitis
 - 400 mg QD + 80 mg osi- Grade 3 dermatitis acneiform
- Two patients (3.6%) discontinued due to TRAEs
- There were no treatment-related deaths
- Dose escalation is on-going with MTD/RP2D yet to be determined

AE, adverse event; BID, twice daily; DLT, dose limiting toxicity; EGFR, epidermal growth factor receptor; MTD, maximum tolerated dose; QD, once daily; RP2D, recommended phase 2 dose; TRAE, treatment-related adverse event; WT, wild type.

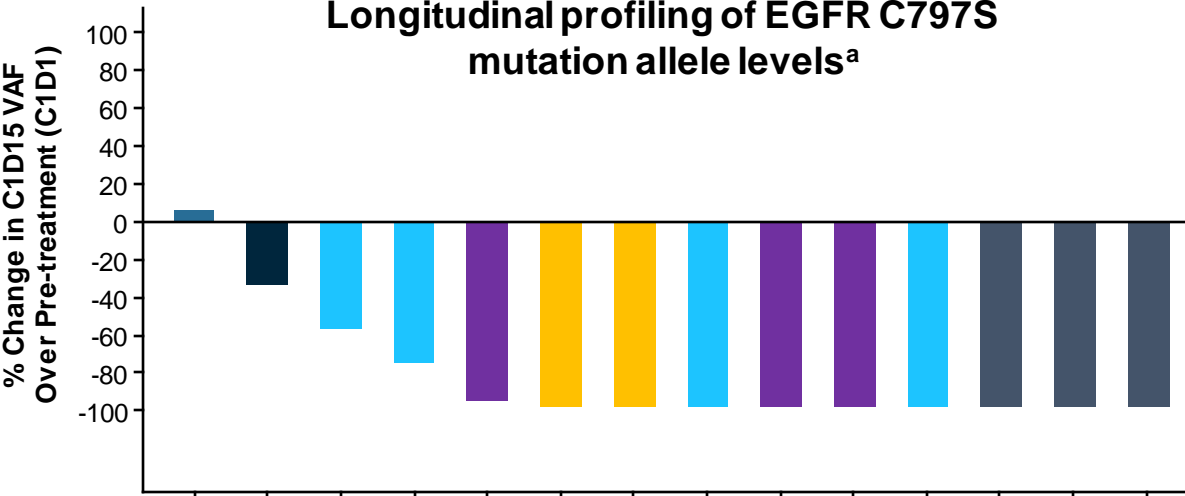
1. Brown K, et al. *Br J Clin Pharmacol.* 2017;83(6):1216-1226. 2. Planchard D, et al. *Cancer Chemother Pharmacol.* 2016;77:767-776.

BLU-945 + osimertinib combination therapy resulted in dose-dependent reduction of EGFR T790M and EGFR C797S mutant allele levels at Cycle 1, Day 15

Longitudinal profiling of EGFR T790M mutation allele levels^a



Longitudinal profiling of EGFR C797S mutation allele levels^a



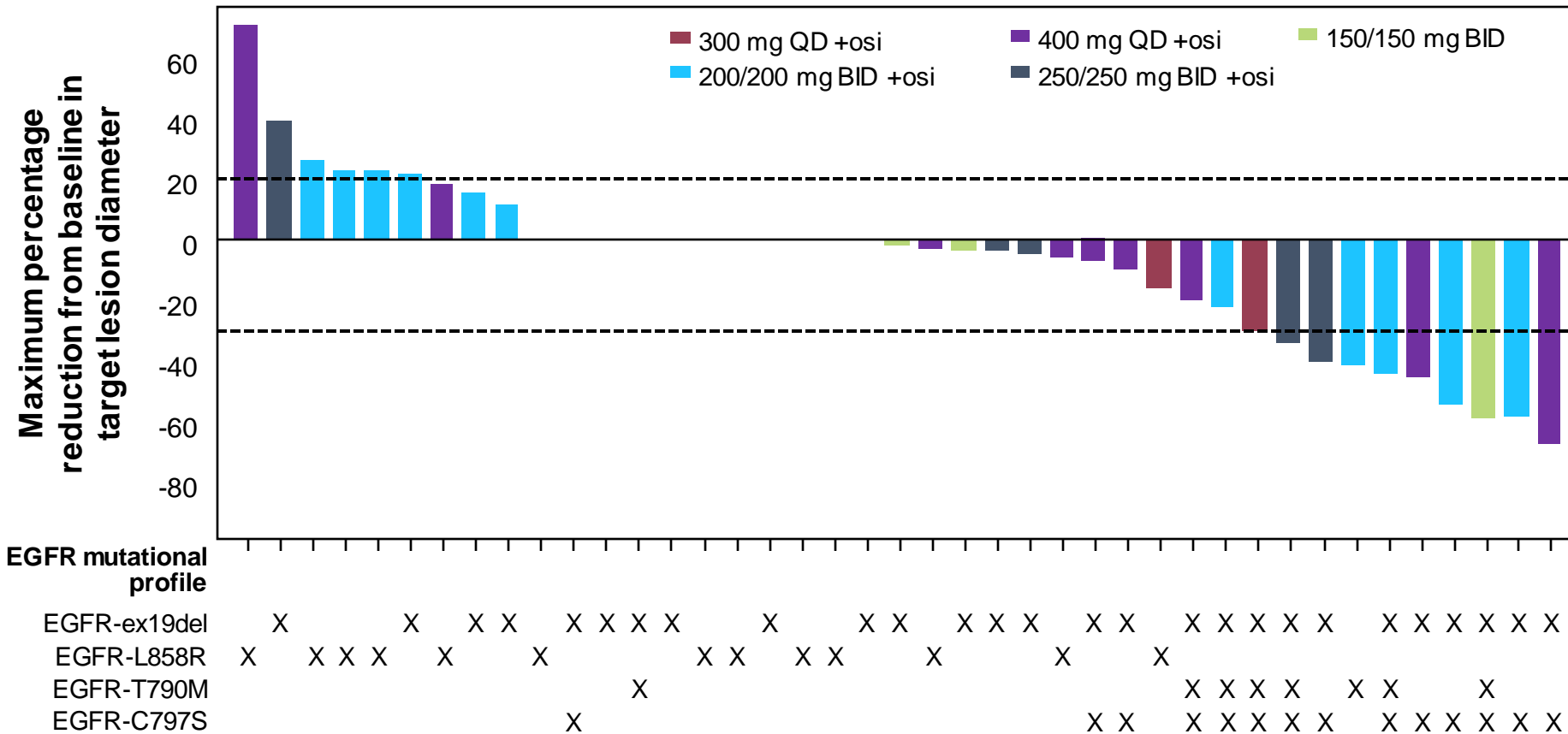
| EGFR mutational profile | 1 | 2 | 3 | 4 | 5 | 6 |
|-------------------------|---|---|---|---|---|---|
| EGFR-ex19del | X | X | X | X | X | X |
| EGFR-L858R | | | | | | |
| EGFR-T790M | X | X | X | X | X | X |
| EGFR-C797S | X | X | X | X | X | X |

| EGFR mutational profile | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 10 | 11 | 12 | 13 | 14 |
|-------------------------|---|---|---|---|---|---|---|---|---|----|----|----|----|----|
| EGFR-ex19del | X | X | X | X | X | X | X | X | X | X | X | X | X | X |
| EGFR-L858R | | | | | | | | | | | | | | |
| EGFR-T790M | X | X | X | X | X | X | X | X | X | X | X | X | X | X |
| EGFR-C797S | X | X | X | X | X | X | X | X | X | X | X | X | X | X |

■ 200 mg QD +osi
 ■ 300 mg QD +osi
 ■ 400 mg QD +osi
 ■ 100/100 mg BID +osi
 ■ 150/150 mg BID +osi
■ 200/200 mg BID +osi
 ■ 250/250 mg BID +osi

^aPercent change greater than 100% are displayed as 100% in the figure. EGFR mutational profile based on results from Foundation One Liquid CDxbaseline (C1D1) analysis. Note: Patient with multiple mutations for EGFR C797S in the same specimen are shown as a different colored X in the EGFR mutational profile.

Early BLU-945 + osimertinib antitumor activity^a



• In the ongoing dose-escalation, tumor shrinkage, including 4 confirmed PRs, was observed in patients who had progressed on osimertinib as the last therapy line

^aPatients with EGFR-mutant positive NSCLC were enrolled based on local mutation assessment of tumor biopsy or blood ctDNA with a follow-up central ctDNA assessment at C1D1. Patients were counted only once. BID, twice daily; EGFR, epidermal growth factor receptor.

Conclusions

- In heavily pretreated EGFR-mutant NSCLC patients, BLU-945 monotherapy was active and well-tolerated; however, due to genomic heterogeneity, responses were not durable
- Emerging BLU-945 + osimertinib combination data demonstrated clinical activity post progression on osimertinib and was well tolerated with infrequent EGFR WT toxicity
- A correspondence between reduction of the resistance mutation alleles by ctDNA and tumor shrinkage was observed in both cohorts
- Phase 1 data support BLU-945 + osimertinib as a differentiated, fully oral, novel combination for treatment of EGFR-mutant NSCLC, warranting further clinical development
 - Combination escalation is ongoing with RP2D/MTD yet to be established

EGFR, epidermal growth factor receptor; MTD, maximum tolerated dose; NSCLC, non-small cell lung cancer; RP2D, recommended phase 2 dose; WT, wild type.

Medical writing support was provided by Maureen Wallace-Nadolski, PhD, of Round Hill, a Lockheed company (Stamford, CT, USA), and was supported by Blueprint Medicines Corporation, Cambridge, MA, USA, according to Good Publication Practice guidelines.