



# Phase 2 Study of the HSP-90 Inhibitor AUY922 in Previously Treated and Molecularly Defined Patients with Advanced Non-Small Cell Lung Cancer

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## ABSTRACT

**Introduction:** In this phase 2 study, we evaluated the activity of AUY922 in pretreated patients with stage IV NSCLC.

**Methods:** Patients with advanced NSCLC were divided into molecularly defined strata based on mutations in the *EGFR* gene, the ALK receptor tyrosine kinase gene (*ALK*), the *KRAS* gene, or the wild type of all three. All patients must have received more than two prior lines of therapy, except for those in a fifth stratum for a less pretreated *EGFR* cohort (*EGFR*<2). In the *EGFR*-mutant and *ALK*-rearranged strata, prior platinum therapy was not required. Patients with *EGFR* mutation must have received an *EGFR* tyrosine kinase inhibitor unless they had de novo resistance (e.g., T790M or exon 20 insertions). Eligible patients received weekly intravenous AUY922, 70 mg/m<sup>2</sup>. The primary objective was to estimate efficacy (complete or partial response, or in the absence of complete or partial response, stable disease) at 18 weeks, by the Response Criteria in Solid Tumors.

**Results:** A total of 153 patients from 21 global centers were enrolled from October 2010 to November 2014. The investigator-assessed overall response rate and stable disease rate at 18 weeks were 31.8% and 9.1% in the *ALK*-rearranged stratum, 17.1% and 8.6% in *EGFR*-mutant stratum, 9.7% and 22.6% in the *EGFR*<2 stratum, 0% and 7.1% in *KRAS*-mutant stratum, and 8.8% and 8.8% in wild-type stratum. Biomarker data showed activity of AUY922 in *EGFR*-mutant patients with exon 19 deletion, T790M mutation, and exon 20 insertion. The most common (≥40%) all-causality adverse events were diarrhea, nausea, and decreased appetite. Visual-related disorders were reported in 79.7% of patients (most were grade 1/2). Thirty-five patients (22.9%) reported night blindness.

**Conclusion:** AUY922 is active in patients with NSCLC, particularly among patients with *ALK* rearrangements and *EGFR* mutations.

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**Keywords:** NSCLC; AUY922; HSP90 inhibitor; *EGFR* mutation; *KRAS* mutation; *ALK* rearrangement

## Introduction

Heat-shock protein90 (HSP90) is an adenosine triphosphate-dependent, molecular chaperone involved in the maintenance, maturation, and functioning of several oncogenic client proteins relevant in the pathogenesis of NSCLC.<sup>1</sup> Studies on resected NSCLC tumor specimens have shown that a lower expression of the gene encoding HSP90 correlates with better prognosis.<sup>2</sup> Inhibition of HSP90 has also been shown to induce apoptosis in NSCLC cells.<sup>1</sup>

AUY922 is a highly potent, adenosine triphosphate-competitive, nongeldanamycin HSP90 inhibitor. It potentially inhibits growth in most NSCLC cell lines in vitro and in a variety of tumor xenografts in vivo.<sup>3–6</sup> The first-in-human, phase 1 dose-escalation study of AUY922, which was conducted in patients with advanced solid tumors demonstrated acceptable tolerability at the recommended dose of 70 mg/m<sup>2</sup>.<sup>7</sup> These preclinical and clinical studies provided the rationale for the current phase 2 study of AUY922 in patients with advanced NSCLC with different tumor molecular etiologies.

## Patients and Methods

### Study Design

This was a global, multicenter, open label phase 2 study of AUY922 that was conducted in 21 global centers between October 2010 and November 2014. Patients with advanced NSCLC were divided into molecularly defined strata based on mutations in the *EGFR* gene, the ALK receptor tyrosine kinase gene (*ALK*), the *KRAS* gene, or the wild-type of all three. All patients must have received more than 2 prior lines of therapy, except for those in a fifth stratum for a less pretreated *EGFR* cohort (*EGFR*<2). In the *EGFR*-mutant and *ALK*-rearranged strata, prior platinum therapy was not required. Patients with *EGFR* mutation must have received an *EGFR* tyrosine kinase inhibitor (TKI) unless they had de novo resistance.

The primary objective was to estimate efficacy as assessed by the Response Criteria in Solid Tumors, classified as complete or partial response (overall response rate [ORR]) or in the absence of complete or partial response, as stable disease or no clinical benefit (NCB) for each study stratum at 18 weeks after treatment initiation. NCB is defined as development of progressive disease in within 18 weeks after the patient had started receiving treatment. Secondary objectives included assessment of overall survival (OS), progression-free survival (PFS), safety, and pharmacokinetics. An exploratory objective was included to examine molecular markers of response. Further details are provided in the Supplementary Materials and Methods.

## Results

### Patient Disposition and Characteristics

Of a total of 153 patients, 150 were assigned to five strata, *ALK*-rearranged (n = 22), *EGFR*-mutant (n = 35), *EGFR*<2 (n = 31), *KRAS*-mutant (n = 28), and wild-type (n = 34), and three patients were classified as “unknown” because the tumor tissue from them was not of a quality acceptable for molecular determination. Of the

153 patients, 151 (98.7%) discontinued treatment, most (73.2%) because of disease progression.

Baseline demographics and disease characteristics are presented in Table 1. All patients had received one or more prior antineoplastic medications, including chemotherapy in 92.2% of patients and targeted therapy

in 63.3%. Among these, four patients (2.6%), 31 (20.3%), and 40 (26.1%) received more than four, four, and three previous regimens, respectively, whereas the remaining 78 patients (51%) received one or two previous regimens. At baseline, TKIs were administered to nearly all patients in the *EGFR*-mutant (97.1%) and

**Table 1.** Baseline Demographics and Clinical Characteristics

Characteristic	No. of patients (%)						
	<i>ALK</i> -Rearranged (n = 22)	<i>EGFR</i> -Mutant (n = 35)	<i>EGFR</i> <2-Mutant (n = 31)	Wild-Type (n = 34)	<i>KRAS</i> -Mutant (n = 28)	Unknown (n = 3)	All Patients (N = 153)
Age, y							
Median	53.0	63.0	56.0	62.5	60.0	66.0	59.0
Range	37-73	33-80	40-76	40-73	39-72	48-67	33-80
<65, n (%)	18 (81.8)	21 (60.0)	25 (80.6)	23 (67.6)	20 (71.4)	1 (33.3)	108 (70.6)
≥65, n (%)	4 (18.2)	14 (40.0)	6 (19.4)	11 (32.4)	8 (28.6)	2 (66.7)	45 (29.4)
Female	15 (68.2)	25 (71.4)	19 (61.3)	18 (52.9)	11 (39.3)	0.0	88 (57.5)
Race, n (%)							
White	16 (72.7)	25 (71.4)	16 (51.6)	23 (67.6)	25 (89.3)	2 (66.7)	107 (69.9)
Asian	6 (27.3)	10 (28.6)	13 (41.9)	9 (26.5)	2 (7.1)	1 (33.3)	41 (26.8)
Black	0	0	0	1 (2.9)	0	0	1 (0.7)
Other	0	0	2 (6.5)	1 (2.9)	1 (3.6)	0	4 (2.6)
WHO performance status, n (%)							
0	9 (40.9)	13 (37.1)	11 (35.5)	10 (29.4)	10 (35.7)	1 (33.3)	54 (35.3)
1	11 (50.0)	19 (54.3)	20 (64.5)	21 (61.8)	18 (64.3)	2 (66.7)	91 (59.5)
2	2 (9.1)	3 (8.6)	0.0	3 (8.8)	0.0	0.0	8 (5.2)
Smoking status, n (%)							
Current smoker	2 (9.1)	1 (2.9)	1 (3.2)	2 (5.9)	3 (10.7)	1 (33.3)	10 (6.5)
Ex-smoker	5 (22.7)	12 (34.3)	16 (51.6)	18 (52.9)	21 (75.0)	0.0	72 (47.1)
Never smoked	15 (68.2)	22 (62.9)	14 (45.2)	14 (41.2)	4 (14.3)	2 (66.7)	71 (46.4)
Median time since initial diagnosis to first dose of study drug (range), mo	22.37 (2.9-64.1)	24.61 (7.6-32.8)	19.15 (3.1-65.1)	18.00 (4.0-65.7)	19.91 (5.1-43.5)	18.79 (8.0-27.1)	20.83 (2.9-132.8)
No. of prior regimens, n (%)							
1	2 (9.1)	2 (5.7)	13 (41.9)	0.0	1 (3.6)	0.0	18 (11.8)
2	4 (18.2)	14 (40.0)	18 (58.1)	16 (47.1)	8 (28.6)	0.0	60 (39.2)
3	7 (31.8)	12 (34.3)	0.0	5 (14.7)	14 (50.0)	2 (66.7)	40 (26.1)
4	7 (31.8)	7 (20.0)	0.0	12 (35.3)	4 (14.3)	1 (33.3)	31 (20.3)
>4	2 (9.1)	0.0	0.0	1 (2.9)	1 (3.6)	0.0	4 (2.6)
Therapy type at last regimen, n (%)							
Chemotherapy	13 (59.1)	19 (54.3)	11 (35.5)	25 (73.5)	22 (78.6)	2 (66.7)	92 (60.1)
Hormonal therapy	0	0	1 (3.2)	0	0	0	1 (0.7)
Immunotherapy	0	0	1 (3.2)	1 (2.9)	1 (3.6)	0	3 (2.0)
Targeted therapy	9 (40.9)	18 (51.4)	19 (61.3)	11 (32.4)	6 (21.4)	1 (33.3)	64 (41.8)
Other	0	1 (2.9)	1 (3.2)	1 (2.9)	0	0	3 (2.0)
Tumor histologic/cytologic type, (%)							
Adenocarcinoma	20 (90.9)	32 (91.4)	30 (96.8)	28 (82.4)	23 (82.1)	3 (100.0)	136 (88.9)
Squamous cell carcinoma	0	0	0	2 (5.9)	1 (3.6)	0	3 (2.0)
Adenosquamous cell carcinoma	0	0	0	2 (5.9)	0	0	2 (1.3)
Bronchioalveolar carcinoma	1 (4.5)	1 (2.9)	0	0	2 (7.1)	0	4 (2.6)
Large cell carcinoma	0	1 (2.9)	0	2 (5.9)	1 (3.6)	0	4 (2.6)
Other	1 (4.5)	1 (2.9)	1 (3.2)	0	1 (3.6)	0	4 (2.6)
Key metastatic site of cancer, n (%)							
Bone	11 (50.0)	15 (42.9)	13 (41.9)	15 (44.1)	9 (32.1)	2 (66.7)	65 (42.5)
Brain	4 (18.2)	6 (17.1)	2 (6.5)	5 (14.7)	6 (21.4)	0	23 (15.0)
Liver	5 (22.7)	9 (25.7)	4 (12.9)	8 (23.5)	4 (14.3)	1 (33.3)	31 (20.3)

*ALK*, *ALK* receptor tyrosine kinase gene; *EGFR*<2-Mutant, stratum of patients with *EGFR* mutation and fewer than two prior lines of therapy.

**Table 2.** Summary of Response Assessment by Study Stratum, as per Investigator Assessment and Central Radiological Review

	No. of patients (%)						
Response	<i>ALK</i> -Rearranged (n = 22)	<i>EGFR</i> -Mutant (n = 35)	<i>EGFR</i> <2-Mutant (n = 31)	Wild-type (n = 34)	<i>KRAS</i> -Mutant (n = 28)	Unknown (n = 3)	All patients (N = 153)
Primary assessment: investigator assessment							
ORR (CR + PR) at 18 wk, n (%) (95% CI)	7 (31.8) (13.9-54.9)	6 (17.1) (6.6-33.6)	3 (9.7) (2.0-25.8)	3 (8.8) (1.9-23.7)	0 (0-12.3)	1 (33.3) (0.8-90.6)	20 (13.1) (8.2-19.5)
Stable disease at 18 wk, n (%)	2 (9.1)	3 (8.6)	7 (22.6)	3 (8.8)	2 (7.1)	0	17 (11.1)
NCB, n (%)	13 (59.1)	26 (74.3)	21 (67.7)	28 (82.4)	26 (92.9)	2 (66.7)	116 (75.8)
Secondary assessment: central radiological review							
ORR (CR + PR) at 18 weeks, n (%) (95% CI)	6 (27.3) (10.7-50.2)	5 (14.3) (4.8-30.3)	1 (3.2) (0.1-16.7)	3 (8.8) (1.9-23.4)	0 (0-12.3)	0 (0-70.8)	15 (9.8) (5.6-15.7)
Stable disease at 18 wk, n (%)	1 (4.5)	1 (2.9)	7 (22.6)	4 (11.8)	3 (10.7)	1 (33.3)	17 (11.1)
NCB, n (%)	15 (68.2)	29 (82.9)	23 (74.2)	27 (79.4)	25 (89.3)	2 (66.7)	121 (79.1)

ALK, ALK receptor tyrosine kinase gene; EGFR<2-Mutant, stratum of patients with EGFR mutation and fewer than two prior lines of therapy; ORR, overall response rate; CR, complete response; PR, partial response; CI, confidence interval; NCB, no clinical benefit.

EGFR<2 (90.3%) strata, most commonly erlotinib. Among patients with ALK rearrangement who had received at least one prior regimen, 77.3% had received the anaplastic lymphoma kinase (ALK) inhibitor crizotinib.

### Efficacy

Overall, 20 patients achieved an investigator-assessed tumor response (ORR of 13.1%) and 17 patients (11.1%) had stable disease at 18 weeks. The investigator-assessed ORR and stable disease at 18 weeks among the five strata are reported in Table 2. Most of the patients (75.8%) showed NCB, as they did not have a radiological response per the Response Criteria in Solid Tumors and progressed within 18 weeks of study treatment. Overall, on the basis of central radiological review, 15 patients (9.8%) had a radiological response, 17 patients (11.1%) had stable disease at 18 weeks, and the remainder (79.1%) showed NCB (see Table 2).

Bayesian analysis of the ORR and NCB rates demonstrated (1) a high probability of improved efficacy (on the basis of predefined thresholds) in the ALK-rearranged (0.899) and EGFR-mutant (0.832) strata, (2) a low probability of improved efficacy in the wild-type (0.346) and EGFR<2-mutant strata (0.185), and (3) a high probability of no efficacy in the KRAS-mutant stratum (0.036) (Supplementary Table 1).

Investigator-assessed disease control rate was highest in the EGFR<2-mutant (71.0% [95% confidence interval (CI): 52.0–85.8]), ALK-rearranged (59.1% [95% CI: 36.4–79.3]), and EGFR-mutant (57.1% [95% CI: 39.4–73.7]) strata (Supplementary Table 2). Three of the

seven patients who had a response to treatment had received prior crizotinib-based treatment.

Next-generation sequencing data were successfully generated for 54 patients (35% because of limited tissue availability), including 11 of 31 patients (35%) assigned to the EGFR<2 stratum and 14 of 35 patients (40%) assigned to the EGFR-mutant stratum (Table 3). Among the EGFR<2-mutant patients, seven had two mutations each (exon 19 deletions and T790M mutations [n = 6] or L858R and T790M [n = 1]). Among the EGFR-mutant patients who received more than two lines of therapy,

**Table 3.** Summary of Mutations and Responses in Patients in the Less Pretreated EGFR-Mutant and EGFR-Mutant Strata by Next-Generation Sequencing

Study Stratum	Mutation Type	No. of Patients	No. of Responders (PR/CR)
EGFR<2-mutant	Exon 19 deletion and T790M	6	1
	Exon 19 deletion	3	0
	Exon 21 L858R and T790M	1	0
	Not detected	1	1
	Not sequenced	20	Not applicable
EGFR-mutant	Exon 19 deletion	4	1
	Exon 19 deletion and T790M	4	2
	Exon 20 insertion	2	1
	Exon 21 L858R	1	0
	Exon 21 L858R and S768I	1	0
	Exon 21 L861Q	1	0
	L747P	1	0
	Not sequenced	21	Not applicable

CR, complete response; PR, partial response; EGFR<2-mutant, stratum of patients with EGFR mutation and fewer than two prior lines of therapy.

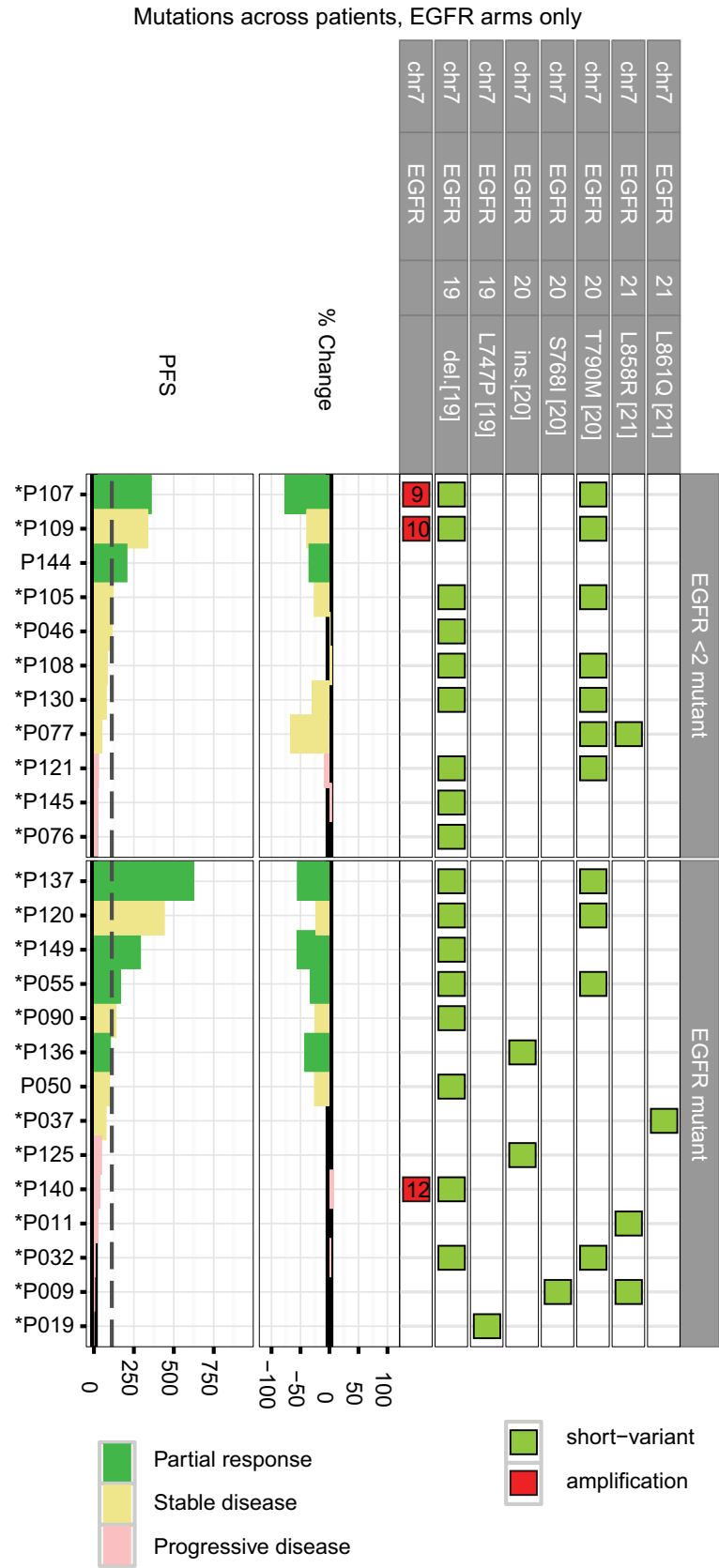
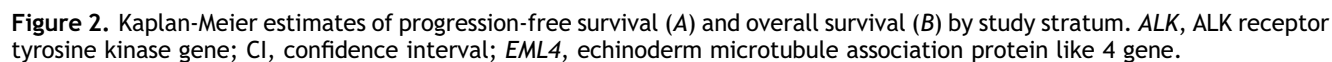


Figure 1. Mutations across patients with clinical variables (EGFR, all patients). PFS, progression-free survival.





**Table 4.** Most Common All-Causality Adverse Events by Study Stratum

Preferred Term	No. of patients (%)						All Patients, n (%) (N = 153)
	ALK-Rearranged, n (%) (n = 22)	EGFR-Mutant, n (%) (n = 35)	EGFR<2-Mutant, n (%) (n = 31)	KRAS-Mutant, n (%) (n = 28)	Wild-Type, n (%) (n = 34)	Unknown, n (%) (n = 3)	
Diarrhea	17 (77.3)	24 (68.6)	24 (77.4)	25 (89.3)	25 (73.5)	3 (100.0)	118 (77.1)
Nausea	10 (45.5)	19 (54.3)	12 (38.7)	16 (57.1)	13 (38.2)	1 (33.3)	71 (46.4)
Decreased appetite	7 (31.8)	15 (42.9)	10 (32.3)	17 (60.7)	12 (35.3)	1 (33.3)	62 (40.5)
Asthenia	10 (45.5)	14 (40.0)	4 (12.9)	13 (46.4)	11 (32.4)	1 (33.3)	53 (34.6)
Headache	13 (59.1)	8 (22.9)	13 (41.9)	8 (28.6)	8 (23.5)	2 (66.7)	52 (34.0)
Fatigue	3 (13.6)	9 (25.7)	15 (48.4)	7 (25.0)	14 (41.2)	0 (0.0)	48 (31.4)
Vomiting	9 (40.9)	14 (40.0)	4 (12.9)	13 (46.4)	4 (11.8)	2 (66.7)	46 (30.1)
Cough	7 (31.8)	8 (22.9)	8 (25.8)	5 (17.9)	9 (26.5)	1 (33.3)	38 (24.8)
Dyspnea	3 (13.6)	13 (37.1)	2 (6.5)	7 (25.0)	9 (26.5)	1 (33.3)	35 (22.9)
Night blindness	5 (22.7)	10 (28.6)	10 (32.3)	3 (10.7)	6 (17.6)	1 (33.3)	35 (22.9)
Photopsia	10 (45.5)	5 (14.3)	6 (19.4)	3 (10.7)	10 (29.4)	0 (0.0)	34 (22.2)
Back pain	3 (13.6)	5 (14.3)	7 (22.6)	8 (28.6)	8 (23.5)	0 (0.0)	31 (20.3)
Vision blurred	6 (27.3)	6 (17.1)	6 (19.4)	5 (17.9)	7 (20.6)	0 (0.0)	30 (19.6)
Visual impairment	6 (27.3)	8 (22.9)	7 (22.6)	6 (21.4)	3 (8.8)	0 (0.0)	30 (19.6)
Constipation	4 (18.2)	6 (17.1)	5 (16.1)	6 (21.4)	5 (14.7)	0 (0.0)	26 (17.0)
Visual acuity reduced	7 (31.8)	6 (17.1)	5 (16.1)	1 (3.6)	7 (20.6)	0 (0.0)	26 (17.0)
Pyrexia	6 (27.3)	3 (8.6)	4 (12.9)	5 (17.9)	4 (11.8)	2 (66.7)	24 (15.7)
Abdominal pain	5 (22.7)	5 (14.3)	2 (6.5)	5 (17.9)	4 (11.8)	0	21 (13.7)
Myalgia	5 (22.7)	4 (11.4)	6 (19.4)	0	4 (11.8)	1 (33.3)	20 (13.1)
Insomnia	5 (22.7)	3 (8.6)	4 (12.9)	0	6 (17.6)	0	18 (11.8)
Anemia	3 (13.6)	3 (8.6)	4 (12.9)	2 (7.1)	5 (14.7)	0	17 (11.1)
Retinal disorder	5 (22.7)	3 (8.6)	7 (22.6)	1 (3.6)	1 (2.9)	0	17 (11.1)
Dry mouth	0	4 (11.4)	1 (3.2)	5 (17.9)	6 (17.6)	0	16 (10.5)
Hypertension	3 (13.6)	5 (14.3)	5 (16.1)	0	3 (8.8)	0	16 (10.5)
Ocular toxicity	3 (13.6)	4 (11.4)	3 (9.7)	4 (14.3)	1 (2.9)	1 (33.3)	16 (10.5)

Note: Occurring in at least 10% of all patients regardless of study drug relationship.

ALK, ALK receptor tyrosine kinase gene; EGFR<2-Mutant, stratum of patients with EGFR mutation and fewer than two prior lines of therapy.

five had two mutations each (exon 19 deletions and T790M mutations [n = 4] and S786I and L858R mutation [n = 1]). Overall in both of the EGFR mutation groups, six patients demonstrated partial response: three patients with both exon 19 deletions and T790M and one patient each with exon 19 deletion or exon 20 insertion (Fig. 1).

In the ALK-rearranged stratum, on the basis of investigator assessment, the median PFS was 2.35 months (95% CI: 0.72–5.09) (Fig. 2A) and the median OS was 9.53 months (95% CI: 4.07–27.73) (Fig. 2B). In the EGFR-mutant stratum, the median PFS was 2.56 months (95% CI: 1.41–3.48) (see Fig. 2A) and the median OS was 8.44 months (95% CI: 5.32–13.31) (see Fig. 2B). Similarly, in the EGFR<2-mutant stratum, the median PFS was 3.68 months (95% CI: 1.64–6.64) (see Fig. 2A) and the median OS was 14.52 months (95% CI: 11.33–25.76) (see Fig. 2B). The median PFS times for the KRAS-mutant

and wild-type strata were 1.38 (95% CI: 0.92–1.71) and 1.25 months (95% CI: 0.99–2.56), respectively (see Fig. 2A). The overall survival times for the KRAS-mutant and wild-type strata were 4.86 (95% CI: 2.76–5.95) and 7.82 (95% CI: 4.76–16.10) months, respectively (see Fig. 2B).

## Safety

Overall, the median duration of exposure was 7.0 weeks, with a longer duration observed in the following strata: ALK-rearranged (median 11.5 weeks, 41% treated past 18 weeks), EGFR<2 (median 12 weeks, 32% treated past 18 weeks), and EGFR-mutant (median 11 weeks, 23% treated past 18 weeks). Most patients (79.7%) did not require a dose reduction, and the rates of reduction were similar across all strata. Overall, 21 patients (13.7%) discontinued treatment because of adverse events (AEs); 10 patients (6.5%) discontinued

treatment because of AEs that were suspected to be related to the study drug.

All patients experienced one or more AE, regardless of causality. The most common all-causality AEs ( $\geq 10\%$  incidence) are summarized in Table 4. Overall, 67.3% reported at least one grade 3 or 4 AE, regardless of causality, most frequently diarrhea (7.2%), dyspnea (7.2%), and asthenia (5.9%). The most frequently reported study drug-related AEs were diarrhea (69.9%), nausea (37.9%), asthenia (24.8%), fatigue (23.5%), night blindness (22.9%), photopsia (22.2%), and vomiting (20.9%).

Almost half of all patients (49.7%) experienced serious AEs, regardless of causality, with 16 of them (10.5%) experiencing a serious AE that was considered related to the study drug. There were no study drug-related deaths.

## Discussion

This study evaluated activity of AUY922 across a broad range of mutations (*ALK*, *EGFR*, and *KRAS*) in pretreated patients with advanced NSCLC. Our results show that AUY922 is active in those with *ALK* rearrangement and *EGFR* mutations. Despite preclinical evidence suggesting that *KRAS*-mutant models are sensitive on the basis of downstream proteins that are clients of HSP90, we did not observe any clinical benefit for patients with *KRAS* mutations.<sup>8</sup>

Development of resistance occurs in most patients treated with *ALK* TKIs.<sup>9,10</sup> AUY922 showed activity in both *ALK* inhibitor-naïve and *ALK* inhibitor-pretreated patients. Similar prior studies with the single-agent HSP90 inhibitors ganetespib and IPI-504 showed responses in patients with NSCLC who had *ALK* rearrangement.<sup>11,12</sup>

Responses in *EGFR*-mutant patients with other HSP90 inhibitors, ganetespib and IPI-504, have been limited.<sup>11,12</sup> In our study, clinical activity was observed in patients with *EGFR* mutations, including exon 19 deletions and exon 20 insertions, and in patients with acquired T790M mutations. Differences in *EGFR* TKI response based on *EGFR* mutation have been reported previously.<sup>13</sup> Similar to our results described here, an ongoing phase 2 study of AUY922 in patients with NSCLC with exon 20 insertions demonstrated an ORR of 32%.<sup>14</sup>

The safety data were similar in patients with advanced solid tumors in the phase 1 trial.<sup>7</sup> Low-grade ocular toxicity was common but effectively managed with dose reductions and dose interruptions. The rate of discontinuation due to AEs was lower in the *ALK*-rearranged and *EGFR*<2 strata than in the other groups, indicating that these patients may not have many disease-related comorbidities or that the AEs may have

been deemed more tolerable in patients who were benefiting from therapy.

In summary, AUY922 had a manageable safety profile and is active in patients with *ALK* rearrangement and *EGFR* mutations, including T790M, which had de novo resistance to *EGFR* TKIs. Future clinical studies are required to address subsequent overlap between different biomarkers to allocate the right therapy to the right group of patients.

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## Supplementary Data

Note: To access the supplementary material accompanying this article, visit the online version of the *Journal of Thoracic Oncology* at [www.jto.org](http://www.jto.org) and at <https://doi.org/10.1016/j.jtho.2017.11.131>.

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