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#WCLC25

## UPDATED EFFICACY, SAFETY, AND GENOMIC DATA IN PATIENTS WITH TRK FUSION LUNG CANCER TREATED WITH LAROTRECTINIB

Jessica J. Lin,<sup>1</sup> Victor Moreno,<sup>2</sup> Shivaani Kummar,<sup>3</sup> Daniel S.W. Tan,<sup>4</sup> Damian T. Rieke,<sup>5</sup>  
Biswajit Dubashi,<sup>6</sup> Kunhi Parambath Haresh,<sup>7</sup> Domnita-Ileana Burcoveanu,<sup>8</sup> Natascha Neu,<sup>9</sup>  
Hong Zheng,<sup>10</sup> Kui Shen,<sup>11</sup> Chiara Mussi,<sup>12</sup> Changsong Qi,<sup>13</sup> Alexander Drilon<sup>14</sup>

<sup>1</sup>Department of Medicine, Massachusetts General Hospital & Harvard Medical School, Boston, MA, USA; <sup>2</sup>START MADRID-FJD, Hospital Fundación Jiménez Díaz, Madrid, Spain; <sup>3</sup>Oregon Health & Science University, Portland, OR, USA; <sup>4</sup>Division of Medical Oncology, National Cancer Centre Singapore, Duke-NUS Medical School, Singapore, Singapore; <sup>5</sup>Charité – Universitätsmedizin Berlin, Berlin, Germany; <sup>6</sup>Jawaharlal Institute of Postgraduate Medical Education and Research, Pondicherry, India; <sup>7</sup>All India Institute of Medical Sciences, New Delhi, India; <sup>8</sup>Bayer HealthCare Pharmaceuticals, Inc., Basel, Switzerland; <sup>9</sup>Chrestos GmbH, Essen, Germany; <sup>10</sup>Bayer HealthCare Pharmaceuticals, Mississauga, ON, Canada; <sup>11</sup>Bayer HealthCare Pharmaceuticals, Whippany, NJ, USA; <sup>12</sup>Bayer S.p.A., Milan, Italy; <sup>13</sup>State Key Laboratory of Holistic Integrative Management of Gastrointestinal Cancers, Beijing Key Laboratory of Carcinogenesis and Translational Research, Peking University Cancer Hospital & Institute, Beijing, China;

<sup>14</sup>Memorial Sloan Kettering Cancer Center & Weill Cornell Medical College, New York, NY, USA

CONQUERING LUNG AND OTHER THORACIC CANCERS WORLDWIDE IN THE 21ST CENTURY

# Introduction and study design

- *NTRK* gene fusions are oncogenic drivers in a variety of cancers, including lung cancer.<sup>1</sup>
- Larotrectinib is the first-in-class, highly selective, CNS-active TRK inhibitor approved for tumor-agnostic use in TRK fusion cancer.<sup>2,3</sup>
- We report an additional year of follow-up, as well as a biomarker analysis, in patients with TRK fusion lung cancer.

## Adult phase 1 (NCT02122913)

- Age ≥18 years
- Advanced solid tumors

n=1

## NAVIGATE: adult/adolescent phase 2 “basket” trial (NCT02576431)

- Age ≥12 years
- Advanced solid tumors
- TRK fusion cancer

n=31

32 patients  
with TRK  
fusion lung  
cancer

## Endpoints

### Primary

- ORR per IRC assessment (RECIST v1.1)

### Secondary

- DoR
- PFS
- OS
- Safety

### Exploratory

- Genomic analysis

Data cutoff  
July 20, 2024

Larotrectinib  
100 mg BID

1. Amatu A et al. *Ann Oncol*. 2019;30:viii5–viii15. 2. Bayer. VITRAKVI US PI. 2023. Available at: [https://labeling.bayerhealthcare.com/html/products/pi/vitrakvi\\_PI.pdf](https://labeling.bayerhealthcare.com/html/products/pi/vitrakvi_PI.pdf). Accessed July 22, 2025.

3. Bayer. VITRAKVI SmPC. 2023. Available at: [https://www.ema.europa.eu/en/documents/product-information/vitrakvi-epar-product-information\\_en.pdf](https://www.ema.europa.eu/en/documents/product-information/vitrakvi-epar-product-information_en.pdf). Accessed July 22, 2025.

BID, twice daily; CNS, central nervous system; DoR, duration of response; IRC, independent review committee; ORR, overall response rate; OS, overall survival; PFS, progression-free survival; RECIST, Response Evaluation Criteria in Solid Tumors.

# Baseline Characteristics in Patients with TRK Fusion Lung Cancer

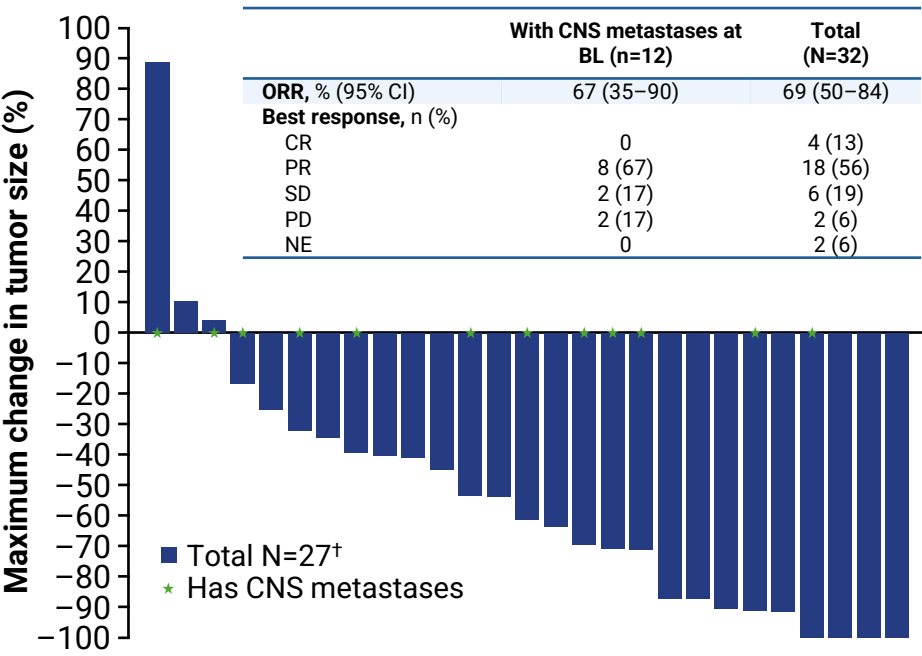
N=32	
Age, years, median (range)	55.5 (25–81)
Sex, n (%)	
Female	19 (59)
Male	13 (41)
<b><i>NTRK</i> gene fusion, n (%)<sup>†</sup></b>	
<i>NTRK1</i>	24 (75)
<i>NTRK3</i>	8 (25)
<b>Tumor histology, n (%)</b>	
Adenocarcinoma	30 (94)
Atypical carcinoid	1 (3)
Neuroendocrine	1 (3) <sup>‡</sup>
<b>Known CNS metastases at baseline, n (%)</b>	
No	20 (63)
Yes	12 (38)

N=32	
<b>Prior therapies, n (%)<sup>§</sup></b>	
Surgery	16 (50)
Radiotherapy	15 (47)
Systemic therapy in the metastatic/unresectable setting	31 (97)
Immunotherapy	13 (41)
<b>Prior systemic therapies in the metastatic/unresectable setting, median (range)</b>	2 (0–8)
<b>Prior systemic therapies in the metastatic/unresectable setting, n (%)</b>	
0	1 (3)
1	12 (38)
2	7 (22)
≥3	12 (38)
<b>Best response to prior systemic therapy, n (%)<sup>  </sup></b>	
Complete response	1 (3)
Partial response	3 (9)
Stable disease	7 (22)
Progressive disease	6 (19)
Other <sup>¶</sup>	14 (44)

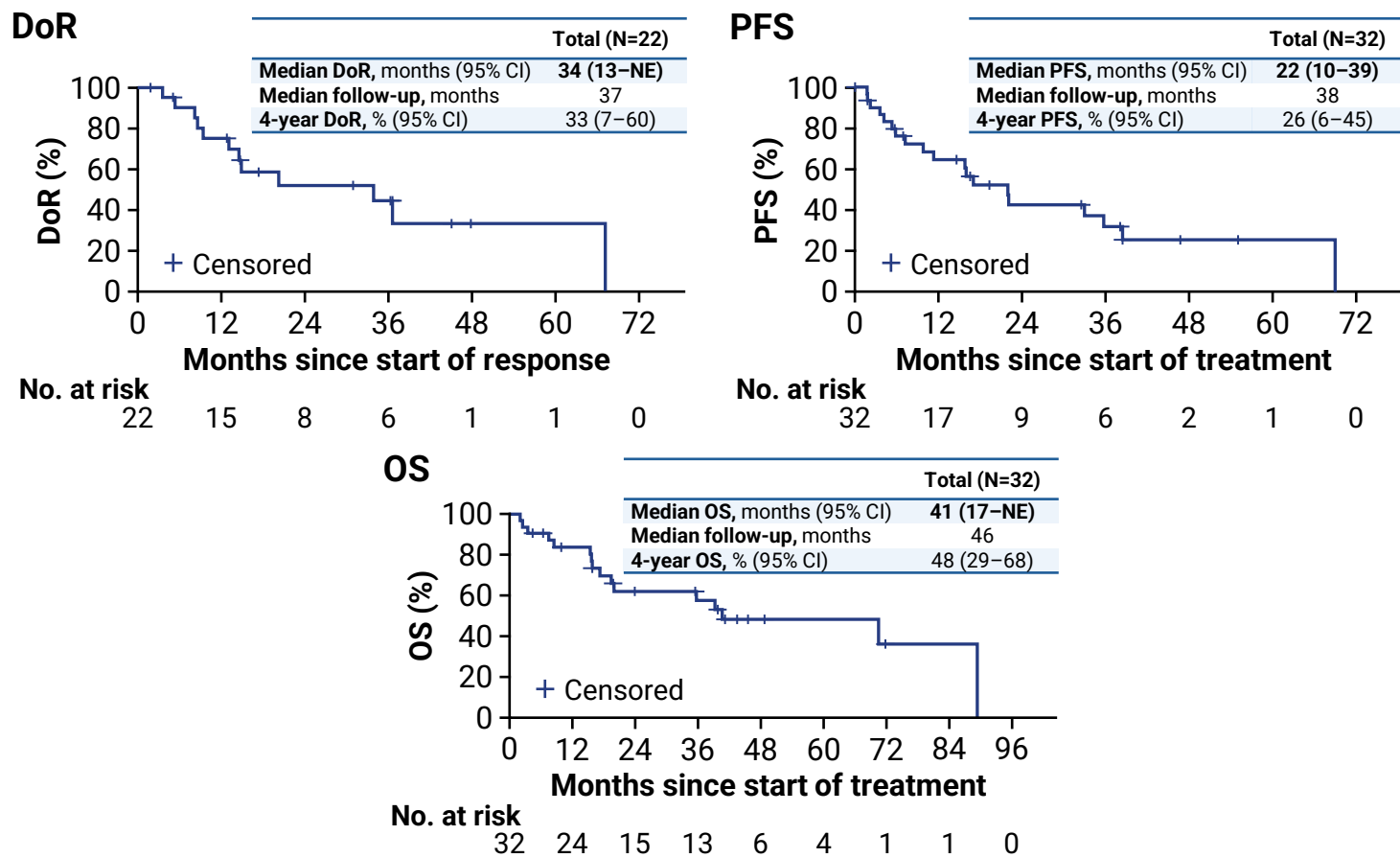
<sup>†</sup>*NTRK* gene fusions were identified by NGS in all patients. <sup>‡</sup>This patient was originally diagnosed with small cell lung cancer that was subsequently assessed as neuroendocrine carcinoma. <sup>§</sup>Patients may be counted in more than 1 row. <sup>||</sup>In the 13 patients with ICI therapy, best overall responses were complete response (n=1), stable disease (n=1), progressive disease (n=4), not evaluable (n=2), and unknown (n=5). <sup>¶</sup>Includes unknown and not evaluable.  
CNS, central nervous system; ICI, immune checkpoint inhibitor; NGS, next-generation sequencing.

# Efficacy: ORR, DoR, PFS, and OS

- ORR was 69% (95% CI 50–84).
- The median time to response was 1.8 months (range 1.5–7.3).

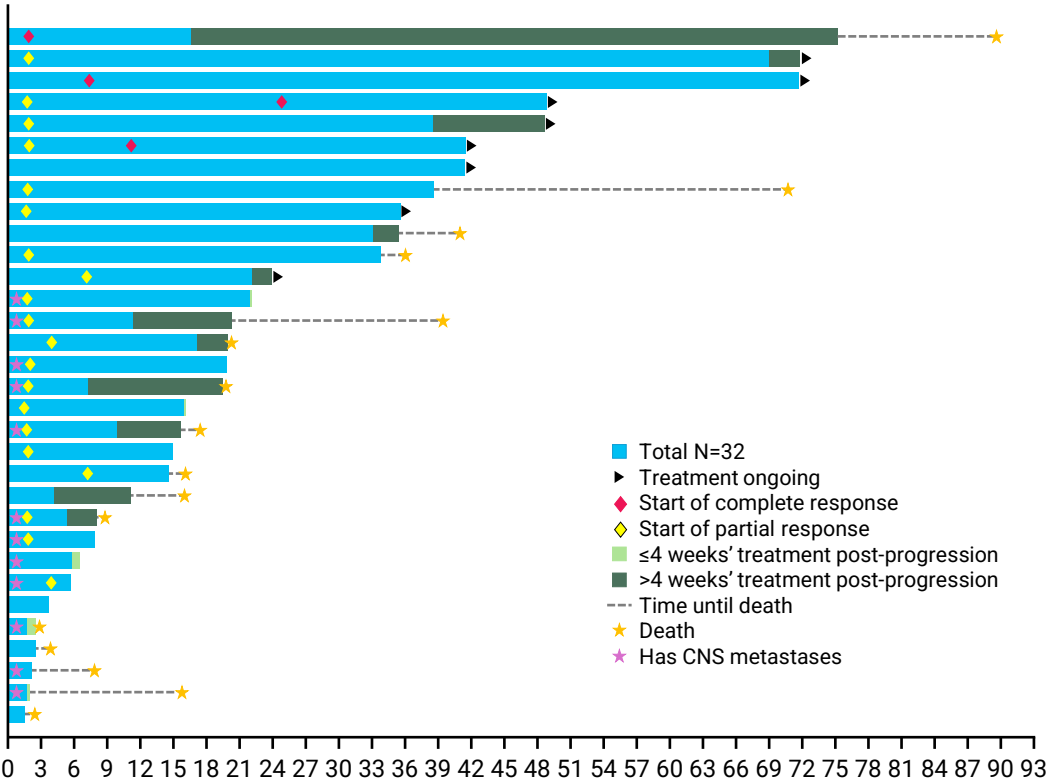


<sup>†</sup>Five patients had no measurable lesions or had missing data as assessed by IRC.  
BL, baseline; CI, confidence interval; CNS, central nervous system; CR, complete response; DoR, duration of response; IRC, independent review committee; NE, not estimable; ORR, overall response rate; OS, overall survival; PD, progressive disease; PFS, progression-free survival; PR, partial response, SD, stable disease.



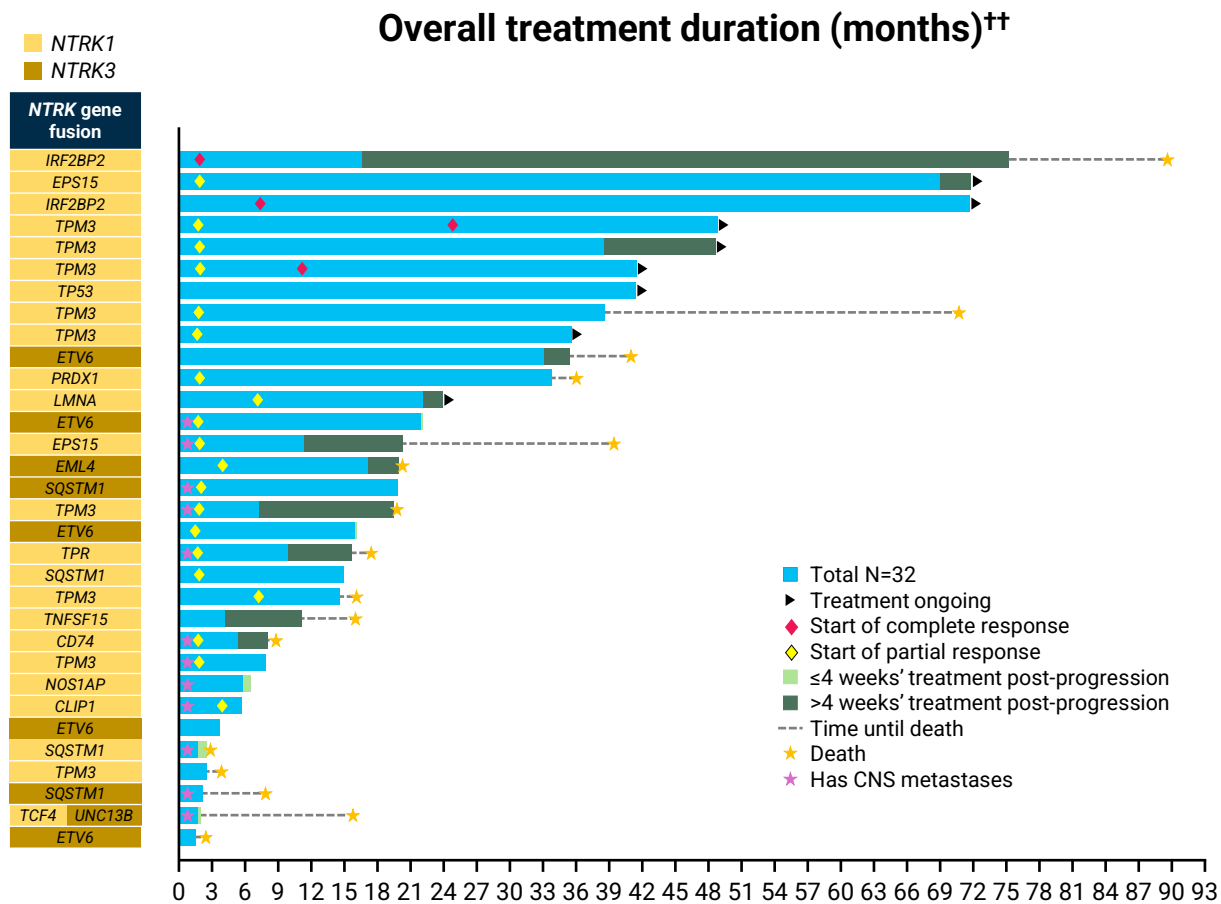
# Biomarker analysis and treatment duration

Overall treatment duration (months)<sup>††</sup>



The CMC project classifies mutations in COSMIC as to their oncogenic potential based on manually curated data. Oncogenic mutations can be classified as known or likely oncogenic (tier 1 or tier 2) as defined by COSMIC v98. <sup>\*</sup>On-target *NTRK* mutations and COSMIC-classified tier 1/2 off-target mutations were identified. Secondary (acquired) resistance was defined as the development of resistance after meaningful clinical benefit (CR/PR/SD ≥4 months). <sup>‡</sup>Mutations in *CBL* and *CTNNB1* were not detected post-baseline. <sup>§</sup>Reasons for no data availability may be sample not available, QC failure, or insufficient sample. <sup>||</sup>Mutations were detected in the baseline ctDNA sample. <sup>¶</sup>Patient had an *ARHGEF11::NTRK1* fusion detected by ctDNA. <sup>††</sup>The median duration of treatment was 20 months (range 2–75). CNS, central nervous system; COSMIC, Catalogue of Somatic Mutations in Cancer; CR, complete response; PR, partial response; SD, stable disease.

# Biomarker analysis and treatment duration

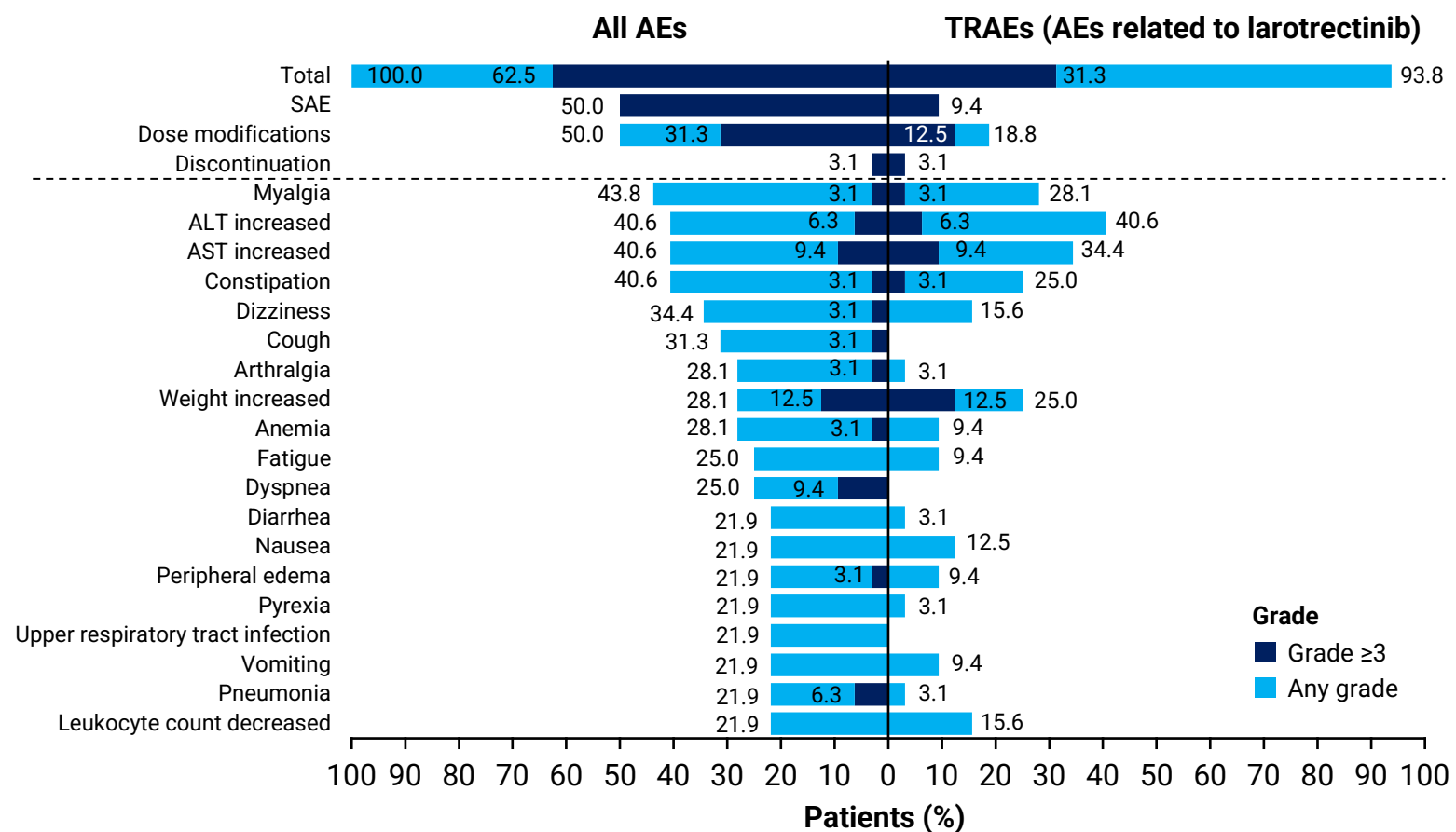


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# AEs in $\geq 20\%$ of patients (N=32)



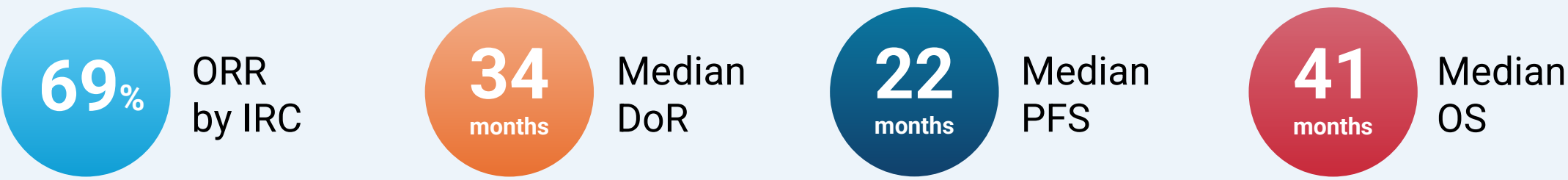
- TRAEs were predominantly Grade 1/2.
- Grade 3/4 TRAEs were reported in 10 (31%) patients.
- One patient discontinued treatment due to TRAEs (increased ALT, AST, and GGT).
- There were no treatment-related deaths.

AE, adverse event; ALT, alanine transaminase; AST, aspartate aminotransferase; GGT, gamma-glutamyl transferase; SAE, serious adverse event; TRAE, treatment-related adverse event.

# Conclusions

- Larotrectinib continues to demonstrate rapid and durable responses, extended survival benefit, and a favorable safety profile in patients with advanced TRK fusion lung cancer.
- These results support the wider adoption of NGS panels that incorporate *NTRK* gene fusion detection to detect patients who may benefit from targeted treatment.
- Post-treatment ctDNA analysis revealed acquired alterations in the TRK kinase domain, *TP53* and *KRAS*.

## In patients with TRK fusion lung cancer:



ctDNA, circulating tumor DNA; DoR, duration of response; IRC, independent review committee; NGS, next-generation sequencing; ORR, overall response rate; OS, overall survival; PFS, progression-free survival.



# Acknowledgments

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- We thank the patients and their families, many of whom travelled long distances to participate in these studies.
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Thank You



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