

# Amivantamab in Wild-type Advanced Non-small Cell Lung Cancer (NSCLC) After Disease Progression on Checkpoint Inhibition and Chemotherapy: Results from the Phase 1b CHRYSALIS Study

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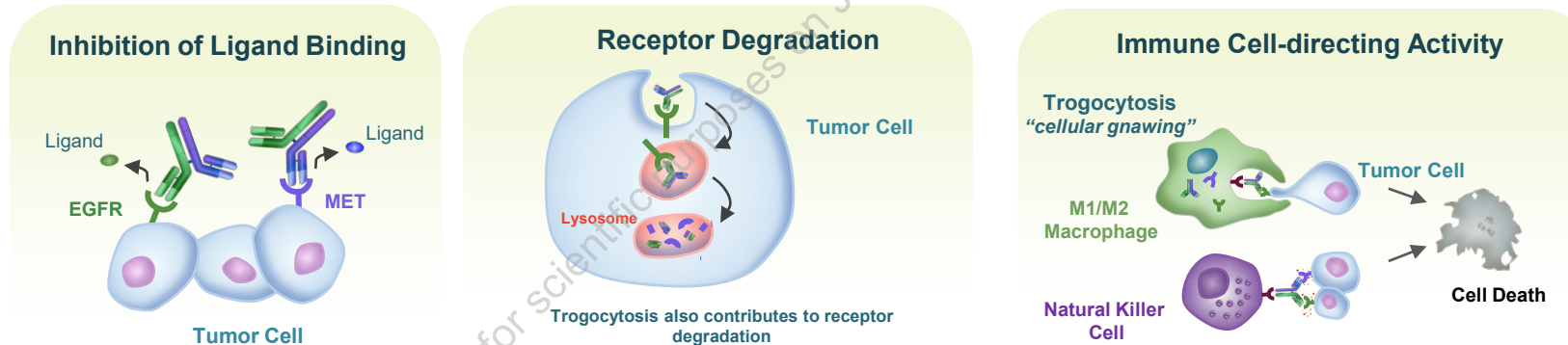
# Declaration of Interests: Byoung Chul Cho

- **Stock or stock options:** TheraCanVac Inc, Gencurix Inc, BridgeBio Therapeutics, KANAPH Therapeutic Inc, Cyrus Therapeutics, Interpark Bio Convergence Corp., J INTS BIO
- **Royalties/intellectual property/patent beneficiary:** Champions Oncology, Crown Bioscience, Imagen
- **Research funding:** MOGAM Institute, LG Chem, Oscotec, Interpark Bio Convergence Corp, GInnovation, GI-Cell, Abion, Abbvie, AstraZeneca, Bayer, Blueprint Medicines, Boehringer Ingelheim, Champions Oncology, CJ Bioscience, CJ Blossom Park, Cyrus, Dizal Pharma, Genexine, Janssen, Lilly, MSD, Novartis, Nuvalent, Oncternal, Ono, Regeneron, Dong-A ST, Bridgebio Therapeutics, Yuhan, ImmuneOncia, Illumina, Kanaph Therapeutics, Therapex, J INTS BIO, Hanmi, CHA Bundang Medical Center
- **Advisory board:** KANAPH Therapeutic Inc, Bridgebio Therapeutics, Cyrus Therapeutics, Guardant Health, Oscotec Inc
- **Consultant:** Abion, BeiGene, Novartis, AstraZeneca, Boehringer-Ingelheim, Roche, BMS, CJ, CureLogen, Cyrus Therapeutics, Ono, Onegene Biotechnology, Yuhan, Pfizer, Eli Lilly, GI-Cell, Guardant, HK Inno-N, Imnewrun Biosciences Inc., Janssen, Takeda, MSD, Medpacto, Blueprint Medicines, RandBio, Hanmi
- **Founder:** DAAN Biotherapeutics (Founder)
- **Member of the board of directors:** Interpark Bio Convergence Corp., J INTS BIO



# Background

- Amivantamab is an EGFR-MET bispecific antibody with immune cell-directing activity that has shown antitumor clinical activity across a range of *EGFR*- and *MET*-driven disease<sup>1-4</sup>
- Overexpression of *EGFR* and *MET* have been observed in previously untreated wild-type adenocarcinoma and squamous NSCLC and are associated with cancer progression<sup>5,6</sup>
- Amivantamab has demonstrated anti-tumor activity through multiple mechanisms of action, with ligand binding identified as the primary mechanism in wild-type NSCLC patient-derived xenograph models<sup>7</sup>
  - Additionally, preclinical inhibition of ligand binding by amivantamab led to reduced *EGFR* and *MET* signaling in wild-type NSCLC<sup>1</sup>



We evaluated amivantamab monotherapy activity in patients with wild-type NSCLC after disease progression on platinum-based chemotherapy and anti-PD-1 or PD-L1 therapy

2L, second-line; ADCC, antibody-dependent cellular cytotoxicity; *EGFR*, epidermal growth factor receptor; Ex20ins, exon 20 insertions; HGF, hepatocyte growth factor; *MET*, mesenchymal-epithelial transition factor; *MET*Ex14, MET exon 14 skipping mutation; NSCLC, non-small cell lung cancer; WT, wild-type.

1. Moores S, et al. *Cancer Res.* 2016;76(13):3942-3953. 2. Vijayaraghavan S, et al. *Mol Cancer Ther.* 2020;19(10):2044-2056. 3. Yun J, et al. *Cancer Discov.* 2020;10(8):1194-1209. 4. RYBREVANT (amivantamab-vmjw) injection, for intravenous use [prescribing information]. Horsham, PA: Janssen Biotech, Inc.; 2021. 5. Huang L, et al. *J Thorac Oncol.* 2014;9: 725-728. 6. Xu N, et al. *Oncotarget.* 2016;7(4):3884-96. 7. Henley B, et al. *Cancer Research.* 2021;81(13\_Supplement):Abstract nr 953.



# Methods

## Dose-escalation phase

**RP2D was identified:**  
Amivantamab 1050 mg IV  
(1400 mg if  $\geq 80$  kg)

## Eligibility Criteria for WT Cohorts

- Patients with advanced NSCLC that did not have *EGFR*, *ALK*, or *METex14* activating mutations (wild-type NSCLC)
- Adenocarcinoma (for **WT-Ad**) or squamous cell carcinoma (for **WT-Sq**) histology
- Confirmed *EGFR* and/or *MET* expression by IHC<sup>a</sup>
- Previously treated with anti-PD-1/PD-L1 and chemotherapy

## Dose-expansion cohorts

**Cohort A:** Post-any *EGFR* TKI (T790M+, C797S+)

**Cohort B:** Post-any *EGFR* TKI (T790M-, C797S-)

**Cohort C:** Post-osimertinib (C797S+)

**Cohort D:** *EGFR* Ex20ins<sup>b</sup>

**Cohort MET-1:** Post-any *EGFR* TKI (*MET* amplified)

**Cohort MET-2:** *METex14*<sup>c</sup>

**Cohort WT-Ad<sup>d</sup>:** *EGFR* wild-type status adenocarcinoma

**Cohort WT-Sq<sup>d</sup>:** *EGFR* wild-type status SCC

**Focus of this presentation**

*CHRYSALIS* also included 2 additional combination cohorts:

- *Amivantamab* + *lazertinib*
- *Amivantamab* + *carboplatin-pemetrexed*

**All cohorts in the *CHRYSALIS* study are now closed**

## Endpoints<sup>e</sup>

- Objective response rate (primary)
- Duration of response
- Clinical benefit rate<sup>f</sup>
- Progression-free survival
- Overall survival
- Adverse events
- Biomarker analyses (exploratory)

Plasma samples were collected pre-treatment, with ctDNA analyzed by Guardant Health (Redwood City, CA)

<sup>a</sup>Patients had to have 1+ staining (defined as  $\geq 50\%$  positive 1+/2+/3+) using archival tissue submitted during prescreening or fresh tissue during screening. <sup>b</sup>Cohort D was presented at WCLC 2020 (Sabari, JK. *J Thorac Oncol.* 2021;16(3):S108-109. OA04.04). <sup>c</sup>Primary *METex14* mutation was confirmed locally; all patients must have failed or be ineligible for standard of care therapy. Data for patients with *METex14* (including those in cohorts other than Cohort MET-2) was presented at WCLC 2023 (Leighl, N. *J Thorac Oncol.* 2023;18(11S):S93-94. OA21.04). <sup>d</sup>Due to slow enrollment progress, both cohorts were closed to further enrollment once one cohort (WT-Ad) had reached the protocol-defined interim analysis. <sup>e</sup>Response was assessed by the investigator per RECIST v1.1. <sup>f</sup>Percentage of patients with confirmed response or SD of  $\geq 11$  weeks duration.

*EGFR*, epidermal growth factor receptor; ex20ins; exon 20 insertion mutation; IHC, immunohistochemistry; IV, intravenous; *MET*, mesenchymal-epithelial transition factor; *METex14*, *MET* exon 14 skipping mutation; NSCLC, non-small cell lung cancer; RP2D, recommended phase 2 dose; SCC, squamous cell carcinoma; SD, stable disease; TKI, tyrosine kinase inhibitor; WT, wild-type.



# Results: Demographic and Baseline Characteristics

Characteristic, n (%)	WT-Ad cohort (n=41)	WT-Sq cohort (n=14)
<b>Median age, years (range)</b>	62 (35–86)	71 (60–81)
<b>Male / female</b>	26 (63) / 15 (37)	8 (57) / 6 (43)
<b>Race</b>		
White	8 (20)	3 (21)
Asian	15 (37)	6 (43)
Black or African American	1 (2)	0
Not reported	17 (41)	5 (36)
<b>ECOG PS</b>		
0	7 (17)	2 (14)
1	33 (80)	12 (86)
2	1 (2)	0
<b>Median number of prior lines (range)</b>	2 (1–5)	2 (1–3)

- In the **WT-Ad cohort**, 41 patients received amivantamab, with a median follow-up of 6.2 months
  - 30 patients had ≥1 post-baseline disease assessment
- In the **WT-Sq cohort**, 14 patients received amivantamab, with a median follow-up of 6.3 months
  - 12 patients had ≥1 post-baseline disease assessment



# Results: Safety Profile

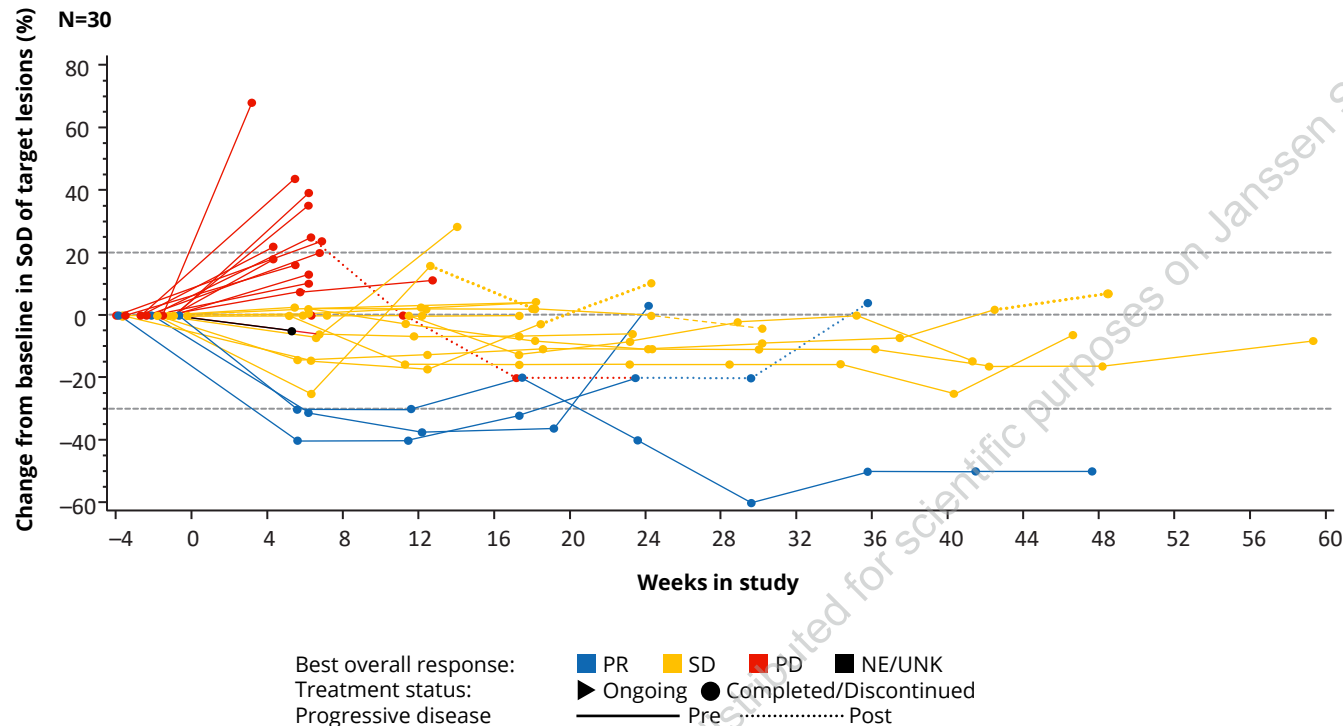
AEs (≥15%) by preferred term, n (%)	WT-Ad cohort (n=41)		WT-Sq cohort (n=14)	
	All grades	Grade ≥3	All grades	Grade ≥3
<b>Associated with EGFR inhibition</b>				
Rash	13 (32)	0	4 (29)	0
Dermatitis acneiform	13 (32)	0	2 (14)	0
Paronychia	11 (27)	0	3 (21)	0
<b>Associated with MET inhibition</b>				
Hypoalbuminemia	16 (39)	1 (2)	4 (29)	1 (7)
Peripheral edema	6 (15)	0	4 (29)	0
<b>Other</b>				
Infusion-related reactions	26 (63)	4 (10)	8 (57)	1 (7)
Dyspnea	11 (27)	3 (7)	2 (14)	0
Constipation	10 (24)	1 (2)	3 (21)	0
Decreased appetite	9 (22)	0	2 (14)	0
Nausea	9 (22)	1 (2)	1 (7)	0
Asthenia	9 (22)	0	1 (7)	1 (7)
Fatigue	8 (20)	1 (2)	1 (7)	0
Pneumonia	6 (15)	5 (12)	6 (43)	3 (21)
Gamma-glutamyltransferase increased	6 (15)	1 (2)	2 (14)	0
Headache	2 (5)	0	3 (21)	0

- Most common AEs were EGFR- or MET-related, primarily grade 1-2
- Treatment-related grade ≥3 AEs:
  - **WT-Ad** cohort: 17%
  - **WT-Sq** cohort: 21%
- Discontinuations of amivantamab due to treatment-related AEs
  - **WT-Ad** cohort: 4 (10%) patients
  - **WT-Sq** cohort: 2 (14%) patients
- There was 1 event of interstitial lung disease, of grade 2 and occurred in the **WT-Ad** cohort



# Results: Amivantamab in WT-Ad Cohort

Percent change from baseline in Sum of Diameters of target lesions<sup>a</sup>



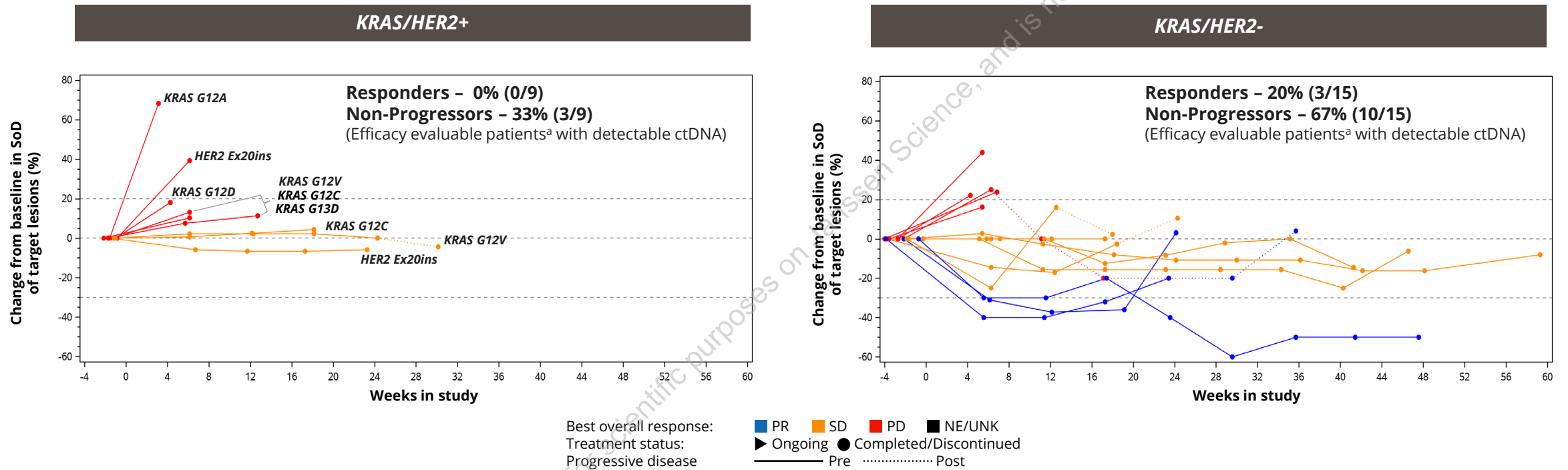
Investigator-assessed response (n=41)	
<b>Median follow-up</b>	6.2 mo (range, 0–18)
<b>ORR</b>	7% (95% CI, 1.5–20)
Median DoR	4.2 mo (95% CI, 4.1–NE)
DoR ≥6 months, n (%)	1 (33)
<b>Best overall response, n (%)</b>	
CR	0
PR	3 (7)
SD	12 (29)
PD	22 (54)
NE/unknown	4 (10)
<b>CBR<sup>b</sup></b>	29% (95% CI, 16.1–45.5)
<b>Median PFS</b>	1.6 mo (95% CI, 1.4–3.6)
<b>Median OS</b>	9.5 mo (95% CI, 5.7–12.0)

<sup>a</sup>30 patients had ≥1 post-baseline disease assessment, 11 patients discontinued due to adverse events (n=5), disease progression (n=4), or physician decision (n=2). <sup>b</sup>CBR is defined as the percentage of patients achieving confirmed complete or partial response, or durable stable disease (duration of at least 11 weeks).

Ad, adenocarcinoma; DoR, duration of response; CBR, clinical benefit rate; CI, confidence interval; mo, months; NE, not evaluable; ORR, objective response rate; OS, overall survival; PD, progressive disease; PFS, progression-free survival; PR, partial response; SD, stable disease; SoD, sum of diameters; UNK, unknown; WT, wild-type.



# Results: Amivantamab Activity in Biomarker-defined Subgroups



- Among patients with detectable ctDNA (n=24), 9 patients were *KRAS/HER2+*
- ORR was **0% (0/9)** for *KRAS/HER2+* and **20% (3/15)** for *KRAS/HER2-*
- Median PFS was shorter for *KRAS/HER2+* (**1.4 months** [95% CI, 0.7–NE]) vs **4.2 months** [95% CI, 1.2–9.4] for *KRAS/HER2-*

<sup>a</sup>Patients were efficacy evaluable if they received at least one dose of study drug and have undergone at least 1 scheduled post-baseline disease assessments or discontinued treatment for any reason.

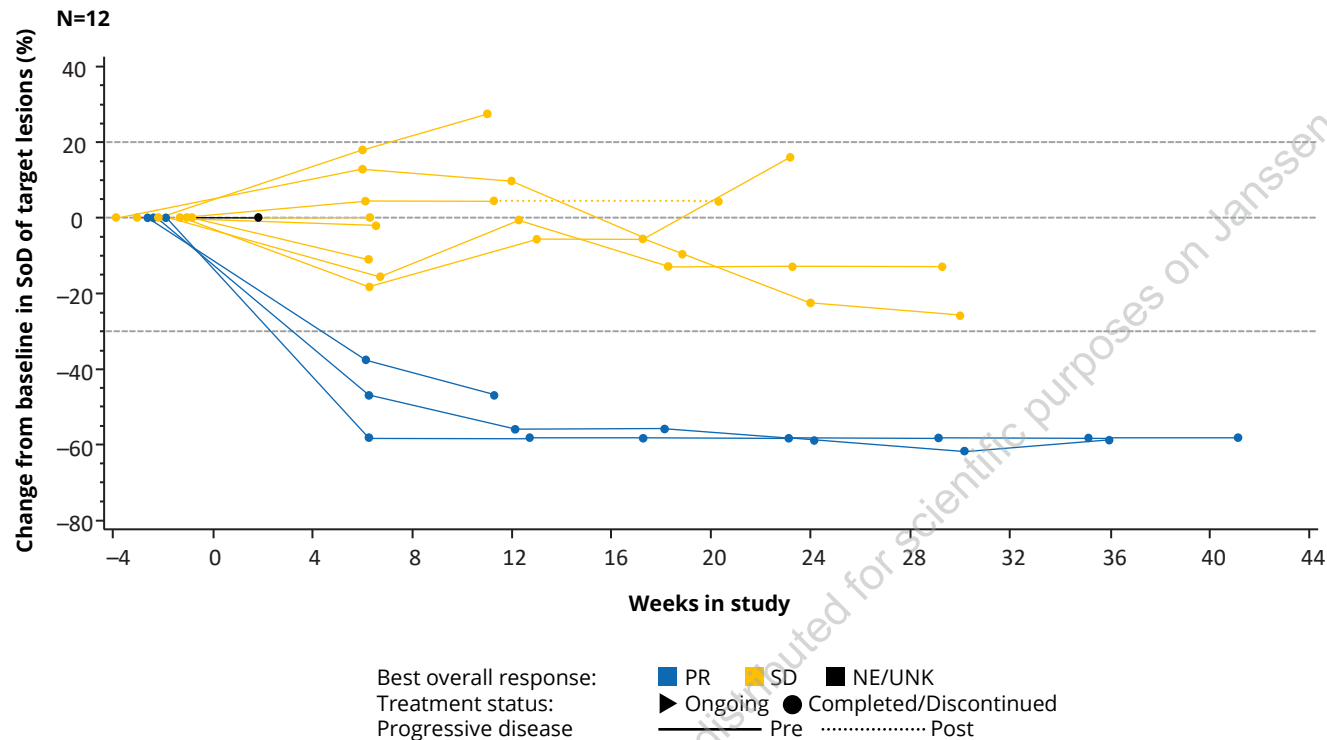
CI, confidence interval; ctDNA, circulating tumor DNA; *KRAS/HER2+*, patients with *KRAS/HER2* mutations; *KRAS/HER2-*, patients without *KRAS/HER2* mutations; NE, not evaluable; ORR, objective response rate; PD, progressive disease; PFS, progression-free survival; PR, partial response; SD, stable disease; SoD, sum of diameters; UNK, unknown.





# Results: Amivantamab in WT-Sq Cohort

Percent change from baseline in SoD of target lesions<sup>a</sup>



Investigator-assessed response (n=14)	
<b>Median follow-up</b>	6.2 mo (range, 1-19)
<b>ORR</b>	21% (95% CI, 5-51)
Median DoR	NE
DoR ≥6 months, n (%)	2 (67%)
<b>Best overall response, n (%)</b>	
CR	0
PR	3 (21)
SD	8 (57)
PD	0
NE/unknown	3 (21)
<b>CBR<sup>b</sup></b>	43% (95% CI, 17.7-71.1)
<b>Median PFS</b>	4.0 mo (95% CI, 2.2-7.3)
<b>Median OS</b>	NE

<sup>a</sup>12 patients had ≥1 post-baseline disease assessment, 2 patients discontinued due to an AE (IRR) or refusal of further study treatment. <sup>b</sup>CBR is defined as the percentage of patients achieving confirmed complete or partial response, or durable stable disease (duration of at least 11 weeks).

DoR, duration of response; CBR, clinical benefit rate; CI, confidence interval; NE, not evaluable; ORR, objective response rate; OS, overall survival; PD, progressive disease; PFS, progression-free survival; PR, partial response; SD, stable disease; SoD, sum of diameters; Sq, squamous cell carcinoma; UNK, unknown; WT, wild-type.



# Conclusions

- ✓ Amivantamab demonstrated antitumor activity in patients with refractory WT adenocarcinoma or refractory WT squamous cell NSCLC
  - No progressive disease was seen in the WT squamous cell carcinoma cohort
- ✓ Amivantamab activity was stronger in patients whose tumors lacked *KRAS/HER2* mutations
  - No response was seen in patients with *KRAS/HER2* mutations



# Key Takeaways

- Amivantamab monotherapy demonstrated antitumor activity in patients with wild-type squamous and adenocarcinoma NSCLC after disease progression on platinum-based chemotherapy and anti-PD-1 or PD-L1 therapy
- No new safety signals of amivantamab monotherapy were identified



# Acknowledgements

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