

# CRESTONE: Initial Efficacy and Safety of Seribantumab in Solid Tumors Harboring NRG1 Fusions

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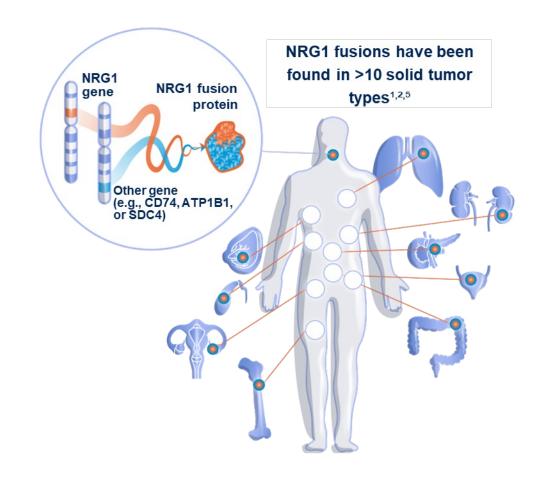






### **NRG1 Gene Fusions in Solid Tumors**

- NRG1 gene fusions are:
  - Rare genomic alterations resulting from the fusion of NRG1 with a partner gene¹
  - NRG1 fusion proteins bind to and activate HER3¹
  - Often mutually exclusive of other known oncogenic alterations<sup>2-4</sup>
  - Found in 0.2% of all solid tumors;
    - Enrichment has been observed in KRAS wild-type PDAC and invasive mucinous adenocarcinoma of the lung<sup>3–6</sup>
- Due to the large intronic regions of the gene fusion,
   RNA-based sequencing is the gold standard for detecting NRG1 fusions<sup>6–8</sup>
- Patients with tumors harboring an NRG1 fusion have poor outcomes with standard therapies, including chemotherapy and immunotherapy<sup>3,9</sup>
- There are currently no approved targeted therapies for tumors harboring NRG1 fusions<sup>6,10</sup>



NRG1, neurequlin-1; PDAC, pancreatic ductal adenocarcinoma

Sources: 1. Jonna S et al. Clin Cancer Res. 2019;25:4966–4972; 2. Drilon A et al. Cancer Discov. 2018;8:686–695; 3. Chang JC et al. Clin Cancer Res. 2021;27:4066–4076; 4. Jones MR et al. Clin Cancer Res. 2019;25:4674–4681; 5. Jonna S et al. J Clin Oncol. 2020;38(15\_suppl):3113–3113; 6. Laskin J et al. Ann Oncol. 2020;31:1693–1703; 7. Russo A et al. Precis Cancer Med. 2020;3:14; 8. Liu S. Lung Cancer. 2021;158:25–28; 9. Drilon A et al. J Clin Oncol. 2021;39:2791–2802; 10. Lyu H et al. Acta Pharm Sin B. 2018;8:503–510.

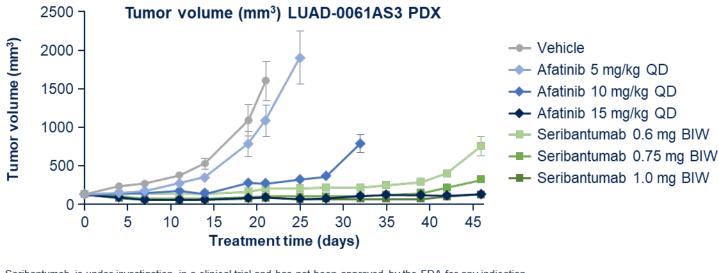


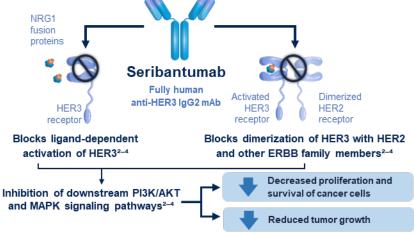


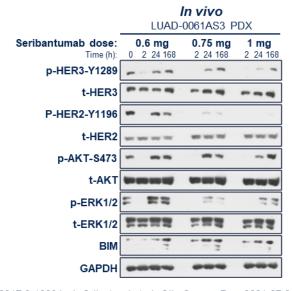


### **Seribantumab Inhibits NRG1 Fusion Tumor Growth**

- Seribantumab is a fully human anti-HER3 lgG2 monoclonal antibody<sup>1,2</sup>
  - Competes with NRG1 to bind to HER3<sup>2,4</sup>
  - Prevents dimerization and phosphorylation of HER3 with other HER family members<sup>2–4</sup>
  - Inhibits downstream PI3K/AKT and MAPK/ERK pathways to inhibit tumor cell growth and proliferation<sup>2–4</sup>
- Seribantumab inhibited tumor growth and induced tumor regression in preclinical models at clinically achievable concentrations<sup>4</sup>







Seribantumab is under investigation in a clinical trial and has not been approved by the FDA for any indication LUAD, lung adenocarcinoma; PDX, patient-derived xenograft

**Sources:** 1. Schoeberl B et al. Sci Signal. 2009;77:1–14; 2. Schoeberl B et al. Cancer Res. 2010;70:2485–2494; 3. Schoeberl B et al. NPJ Syst Biol Appl. 2017;3:16034; 4. Odinstov T et al. Clin Cancer Res. 2021;27:3154–3166.







### **Key Inclusion Criteria**

- ✓ Patients with locally advanced or metastatic solid tumors harboring an NRG1 gene fusion (by local testing)
- ✓ Minimum 1 prior systemic therapy
- ✓ No other oncogenic alterations (except for Cohort 3)

NCT04383210

Seribantumab

3q IV QW<sup>†</sup>

### Cohort 1 n ≥55

No prior pan-ERBB, HER2, or HER3 targeted therapy AND NRG1 gene fusion centrally confirmed<sup>‡</sup>

### **Primary Endpoint:**

 ORR by independent/central radiologic review (per RECIST v1.1)

### **Secondary Endpoints:**

- Safety
- · ORR by investigator assessment
- DOR, PFS, OS, CBR (CR, PR, SD, ≥24 weeks)

### EXPLORATORY Cohort 2 n ≥10

R/R following pan-ERBB, HER2, or HER3 targeted therapy

### EXPLORATORY Cohort 3 n ≥10

NRG1 fusion without EGF-like domain or NRG1 fusion with other molecular alteration or insufficient tissue for confirmatory testing

<sup>†</sup>A safety run-in phase evaluated seribantumab as induction, consolidation, and maintenance dosing: seribantumab 3g QW selected as the optimized RP2D for patients with solid tumors harboring an NRG1 fusion. Patients from the safety run-in who transitioned to 3g QW after induction/reinduction will be included in the primary efficacy analysis per the SAP:

\*Patients are enrolled and treated based on local NRG1 fusion testing result with post-enrollment confirmation by central RNA-based NGS assay.







Safety Population

Primary Efficacy Population

# Patient Demographics and Disease Characteristics

Demographic	Cohort 1 <sup>†</sup> (N=15)	Safety Population <sup>‡</sup> (N=35)
Age		
Median (range)	61 (44, 76)	65 (19, 76)
Sex; n (%)		
Female	10 (67)	24 (69)
Male	5 (33)	11 (31)
Race; n (%)		
American Indian or Alaska Native	1 (7)	1 (3)
Asian	2 (13)	7 (20)
Black or African American	2 (13)	3 (9)
White	10 (67)	23 (66)
Other	0	1 (3)
ECOG PS; n (%)		
0	5 (33)	18 (51)
1	10 (67)	17 (49)
2	0	0

†Includes patients in Cohort 1 who received the optimized RP2D of seribantumab 3g QW including patients in the safety run-in who received seribantumab 3g QW beyond induction/reinduction; ‡Includes all patients who received at least 1 dose of seribantumab in all cohorts; §Colon, endometrial, esophagus, and head and neck; BAG4, FUT10, IL1RL2, ITGB1, POMK, PTN, RBPMS, RNF169, TMPRSS3, VTCN1; ARequired for Cohort 1 only post-enrollment; ACentral RNA-based NGS assay pending/quality not sufficient for testing/results not available. Note: Percentages may not add up to 100% due to rounding

Visit cut-off: 18 April 2022

Disease Characteristic	Cohort1 <sup>†</sup> (N=15)	Safety Population <sup>‡</sup> (N=35)
Primary Tumor Type; n (%)		
Biliary Tract/cholangiocarcinoma	0	2 (6)
Breast	0	4 (11)
NSCLC	14 (93)	20 (57)
Pancreas	1 (7)	5 (14)
Other <sup>§</sup>	0	4 (11)
NRG1 Fusion Partners; n (%)		
ATP1B1	1 (7)	2 (6)
CD74	6 (40)	11 (31)
SDC4	2 (13)	2 (6)
SLC3A2	5 (33)	6 (17)
AGRN	0	2 (6)
APP	0	2 (6)
Other∥	1 (7)	10 (29)
Central NRG1 Fusion Status^; n (%)		
Confirmed	14 (93)	
Unconfirmed	0	٨
Unknown <sup>^^</sup>	1 (7)	
Prior Systemic Therapies		
Median (range)	1 (1, 5)	2 (1, 6)









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# Safety Summary of Seribantumab Monotherapy

### Adverse events reported in ≥15% of patients

	Treatment-emergent AEs (N=35); n (%)			Treatment-related AEs (N = 35); n (%)				
Preferred Term	Any Grade	Grade 1	Grade 2	Grade≥3 <sup>†</sup>	Any Grade	Grade 1	Grade 2	Grade≥ 3 <sup>‡</sup>
Patients with ≥1 AE	35 (100)	8 (23)	10 (29)	17 (49)	30 (86)	17 (49)	11 (31)	2 (6)
Diarrhea	17 (49)	11 (31)	4 (11)	2 (6)	14 (40)	10 (29)	3 (9)	1 (3)
Fatigue	14 (40)	7 (20)	7 (20)	0	10 (29)	5 (14)	5 (14)	0
Rash§	11 (31)	9 (26)	2 (6)	0	9 (26)	7 (20)	2 (6)	0
Hypokalemia	10 (29)	6 (17)	3 (9)	1 (3)	3 (9)	3 (9)	0	0
Nausea	10 (29)	7 (20)	1 (3)	2 (6)	6 (17)	5 (14)	1 (3)	0
Abdominal pain <sup>  </sup>	8 (23)	4 (11)	2 (6)	2 (6)	3 (9)	1 (3)	2 (6)	0
Decreased appetite	8 (23)	4 (11)	3 (9)	0	3 (9)	1 (3)	2 (6)	0
Headache	8 (23)	7 (20)	1 (3)	0	1 (3)	1 (3)	0	0
Hypomagnesemia	8 (23)	6 (17)	1 (3)	0	2 (6)	2 (6)	0	0
Cough	7 (20)	5 (14)	2 (6)	0	1 (3)	1 (3)	0	0
Anemia <sup>^</sup>	6 (17)	4 (11)	1 (3)	1 (3)	1 (3)	1 (3)	0	0
Dysuria	6 (17)	6 (17)	0	0	0	0	0	0

- Safety profile of 35 patients with tumors harboring NRG1 fusions who received at least 1 dose of seribantumab in the CRESTONE study
  - One DLT (Grade 2 fatigue resulting in dose reduction by the Investigator in the safety run-in)
  - 27 (77%) patients received the optimized RP2D of seribantumab 3g QW
- Majority (80%) of TRAEs were Grade 1 or 2
- Two (6%) patients received dose reductions for AEs per the Investigator
  - One patient for Grade 1 ALT increase
  - One patient for Grade 2 fatigue
- No patients discontinued seribantumab for AEs

†2 Grade 5 TEAEs (unrelated to seribantumab) of lung infection (n=1 patient) and malignant neoplasm progression (n=1 patient); ‡No Grade 4 or 5 TRAEs reported; 2 Grade 3 TRAEs of diarrhea (n=1 patient) and vomiting (n=1 patient); §Includes preferred term (PT) of rash and maculo-papular rash; Includes PT of abdominal pain, abdominal pain upper, abdominal distention; Alncludes PT of anemia and iron deficiency anemia.

AE, adverse event; ALT, alanine transaminase; DLT, dose-limiting toxicity; TRAE, treatment-related adverse event Visit cut-off: 18 April 2022







# Clinical Activity of Seribantumab in Tumors Harboring NRG1 Fusions

Investigator-assessed (INV) Response, %	Cohort 1 Primary Efficacy Population <sup>†</sup> (n=12 <sup>‡</sup> )	Cohort 1 - NSCLC Primary Efficacy Population <sup>†</sup> (n=11 <sup>‡</sup> )
Objective response rate; n (%)	4 (33)	4 (36)
Complete response; n (%)	2 (17)	2 (18)
Partial response; n (%)	2 (17)	2 (18)
Stable disease; n (%)	7 (58)	6 (55)
Progressive disease; n (%)	1 (8)	1 (9)
Disease control rate; n (%)	11 (92)	10 (91)

†Defined in the study protocol and Statistical Analysis Plan as Cohort 1 patients with centrally confirmed NRG1 fusion status who received at least 1 dose of seribantumab at 3g QW dosing regimen (starting with Protocol Version 3.0 and later). Patients in the safety-run in (enrolled under Protocol Version 2.0 or earlier) are included if they received seribantumab at 3g QW beyond induction/reinduction;

<sup>‡3</sup> patients are not included for efficacy analysis 2 patients are not for evaluable response (1 patient recently enrolled and not yet eligible for post-baseline scan; 1 patient died due to sepsis, unrelated to disease progression or seribantumab and no post-baseline tumor measurements recorded) AND 1 patient with unknown central NRG1 fusion status as of visit cut-off date.

All responses are confirmed ≥4 weeks per RECIST v1.1. All patients have tumors harboring NRG1 fusion by local testing Note: Percentages may not add up to 100% due to rounding

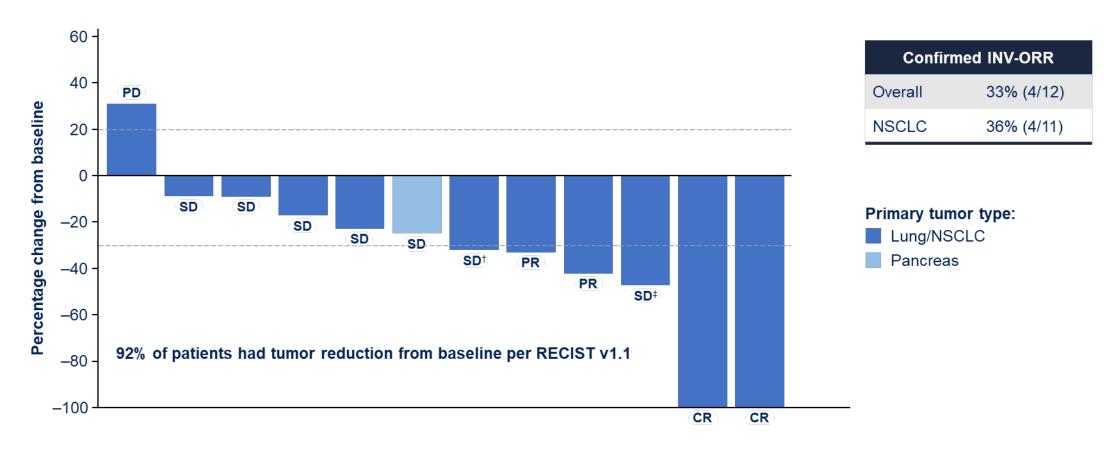
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# Efficacy of Seribantumab in Tumors Harboring NRG1 Fusions



†Unconfirmed PR, unable to be confirmed as subsequent scans showed patient in SD;

‡Unconfirmed PR, patient died due to lung infection (history of COVID-19 infection) before confirmatory scan was able to be completed, no evidence of clinical disease progression at time of death.

INV-ORR, investigator-assessed objective response rate; RECIST v1.1, Response Evaluation Criteria in Solid Tumors version 1.1

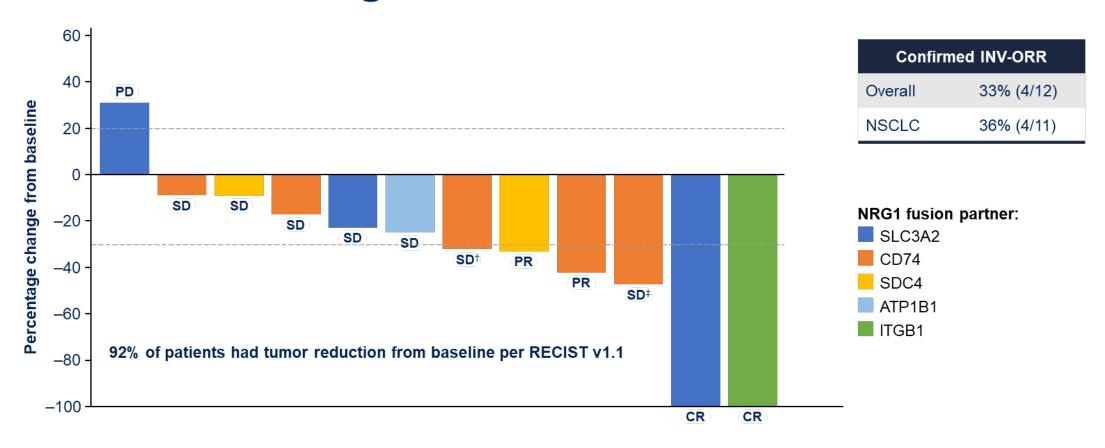
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# Efficacy of Seribantumab in Tumors Harboring NRG1 Fusions Regardless of Fusion Partner



†Unconfirmed PR, unable to be confirmed as subsequent scans showed patient in SD;

†Unconfirmed PR, patient died due to lung infection (history of COVID-19 infection) before confirmatory scan was able to be completed, no evidence of clinical disease progression at time of death.

INV-ORR, investigator-assessed objective response rate; RECIST v1.1, Response Evaluation Criteria in Solid Tumors version 1.1

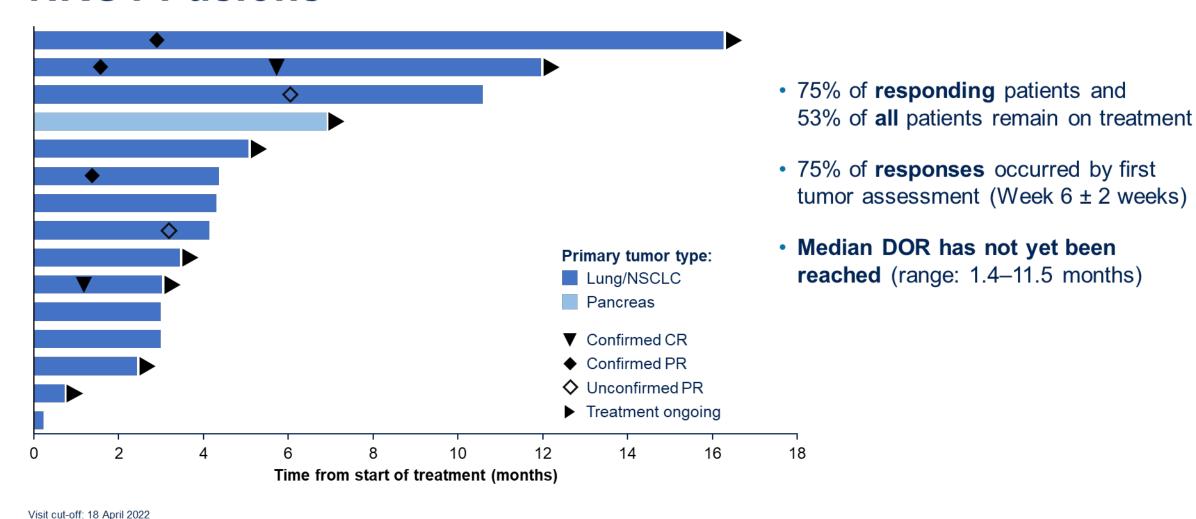
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# **Duration of Seribantumab Therapy in Tumors Harboring NRG1 Fusions**

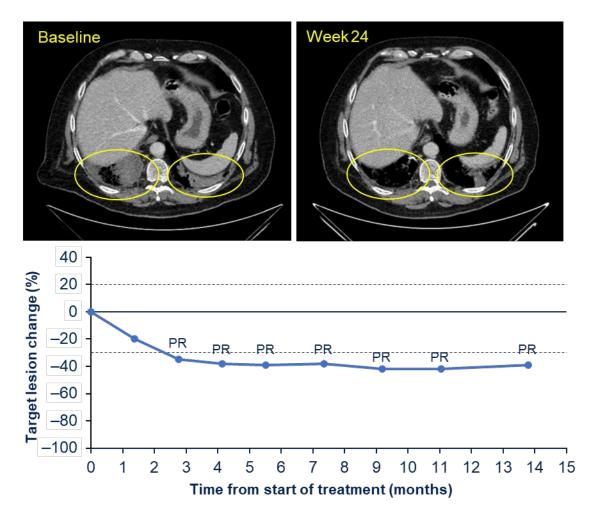








### **Durable Response in CD74-NRG1 NSCLC**



- 70-year-old male with NSCLC adenocarcinoma
- Three prior lines including platinum-based chemotherapy, immunotherapy (IO), and investigational therapy (IO and targeted agent)
- PR (35% tumor reduction) at Week 12; maximum tumor reduction of 42%
- Duration of response 11.5 months (ongoing)
- Seribantumab treatment ongoing for 16.0 months
  - Initiated seribantumab treatment under safety run-in
  - Transitioned to seribantumab 3g QW after induction/consolidation

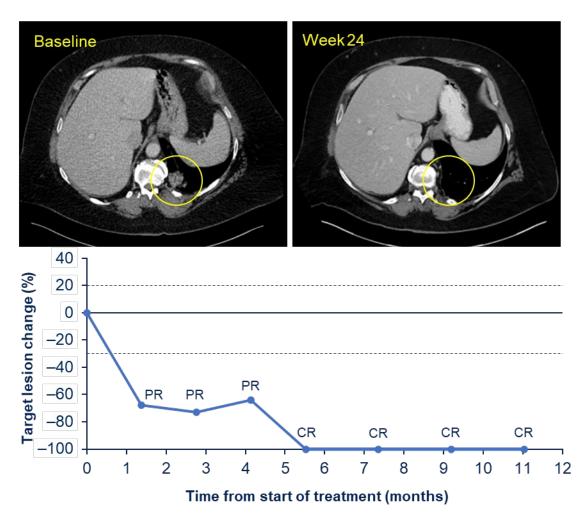
Images courtesy of Tejas Patil, MD; University of Colorado Visit cut-off: 18 April 2022







## Deep and Durable Response in ITGB1-NRG1 NSCLC



- 60-year-old female with NSCLC adenocarcinoma
- Three prior lines including platinum-based chemotherapy and immunotherapy
- PR (68% tumor reduction) at Week 6
- Deepening of response to CR at Week 24
- Duration of response 9.7 months
  - CR ongoing for 5.6 months
- Treatment ongoing for 11.7 months

Images courtesy of Daniel R. Carrizosa, MD; Levine Cancer Institute Visit cut-off: 18 April 2022







### **Conclusions**

- Initial data support the ability of seribantumab to produce deep and durable benefit for patients with previously treated solid tumors harboring NRG1 fusions
  - INV-ORR: 33% (2 PRs, 2 CRs); NSCLC INV-ORR: 36% (2 PRs, 2 CRs)
  - Durable ongoing responses with DOR of 9.7 and 11.5 months
  - Disease control rate of 92%
- Seribantumab was generally well tolerated, and the safety profile was consistent with observations from previous studies evaluating seribantumab<sup>1–4</sup>
  - Majority of AEs were Grade 1 or 2
  - No AEs led to discontinuation of seribantumab treatment
- These data support the continued evaluation of seribantumab in the ongoing Phase 2 CRESTONE (NCT04383210) study as a potential new standard of care for patients with solid tumors harboring NRG1 fusions
  - Comprehensive genomic profiling, particularly RNA-based NGS, will be important to identify patients with gene fusions

INV-ORR, investigator-assessed objective response rate

Sources: 1. Denlinger C et al. Invest New Drugs. 2021;39:1604–1612; 2. Liu JF et al. J Clin Oncol. 2016;20;34:4345–4353; 3. Cleary JM et al. Invest New Drugs. 2017;35:68–78; 4. Sequist LV et al. Oncologist. 2019;24:1095–1102.







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