

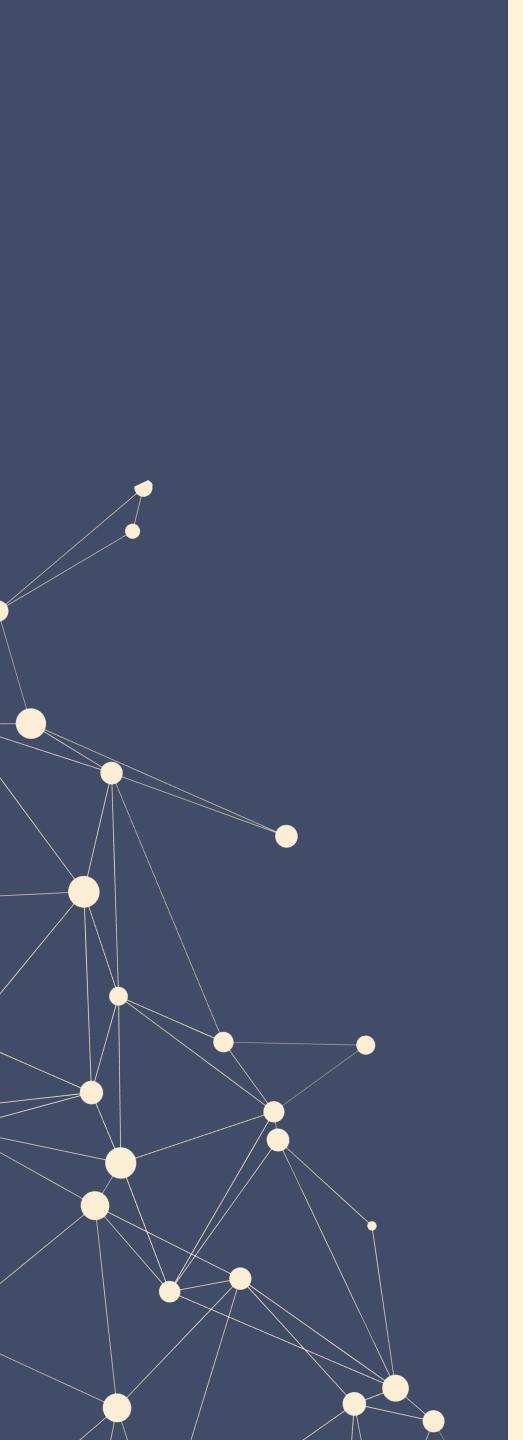
## A PHASE IA/IB STUDY OF BOTENSILIMAB, A NOVEL INNATE/ADAPTIVE IMMUNE ACTIVATOR, PLUS BALSTILIMAB FOR THE TREATMENT OF PATIENTS WITH SARCOMA

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# **Ctos**®





## DISCLOSURES

- Consultant/Advisory Role: Adaptimmune, Adcendo,
- Daiichi Sankyo, Deciphera, Epizyme, Polaris, Springworks
- Institutional Coordinating PI: Agenus







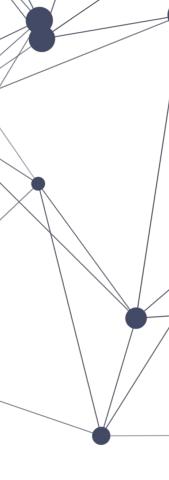
## RECENT I-O TRIALS IN SARCOMA

Summary of Responses in All-comer Soft-Tissue Sarcoma Patients

Study Agents	Study N Range	Median Prior Treatments Range	ORR Range (All-Comers)	Median OS Range
PD-(L)1 as Monotherapy <sup>1-3</sup>	40-79	2	16-37%	11.3-13.0 months
PD-1 + CTLA-4 <sup>4-7</sup>	16-48	2-3	12-16%	13.1-NR months
Chemotherapy Combinations <sup>8-11</sup>	12-88	0	19-37%	14.0-27.6 months
Alkylating Agent Combinations <sup>12</sup>	50	2	2%	5.6-NR months
<b>Receptor Tyrosine Kinase Inhibitor</b> <b>Combinations</b> <sup>13-14</sup>	33-58	0-2	21-25%	24 months

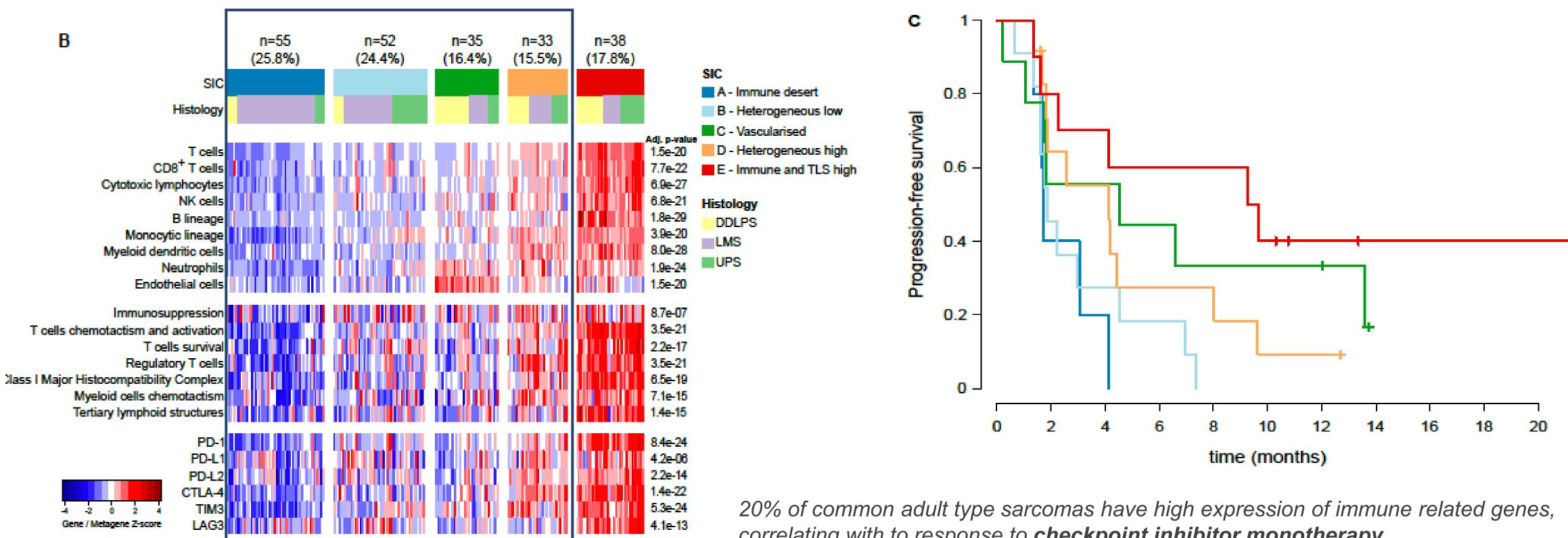
Tawbi HA, et al. *Lancet Oncol.* 2017;18(11):1493-1501. 2. Burgess MA, et al. *J Clin Oncol.* 2019;37(15)\_suppl:11015-11015. 3. Naqash AR, et al. *J Clin Oncol.* 2021;39(15)\_suppl:11519-11519.
 Somaiah N, et al *Lancet Oncol.* 2022;23(9):1156-1166. 5. D'Angelo SP, et al. *Lancet Oncol.* 2018;19(3):416-426. 6. Chen JL, et al. *J Clin Oncol.* 2020;38(15)\_suppl:11511-11511.
 Wagner MJ, et al. *J Immunother Cancer.* 2021;9:e002990. 8. EM Gordon, et al. *J Clin Oncol.* 2022;40(16)\_suppl:11573-11573. 9. Pollack SM, et al. *JAMA Oncol.* 2020;6(11):1778-1782.
 Livingston MB, et al. *Clin Cancer Res.* 2021;27(23):6424-6431. 11. Rosenbaum E, *J Clin Oncol.* 2022;40(16)\_suppl:11516-11516. 12. Toulmonde M, et al. *JAMA Oncol.* 2018;4(1):93-97.
 Martin-Broto J, et al. *J Immunother Cancer.* 2020;8:e001561. 14. Wilky BA, et al. *Lancet Oncol.* 2019;20(6):837-848.







## STANDARD ICIS INEFFECTIVE FOR MAJORITY OF SARCOMA PATIENTS



- 1. Petitprez F, et al. Nature. 2020;577:556-560.
- 2. Progression-free survival graph based on the multicenter Phase 2 clinical trial of pembrolizumab in soft-tissue sarcoma (SARC028).

correlating with to response to **checkpoint inhibitor monotherapy** 



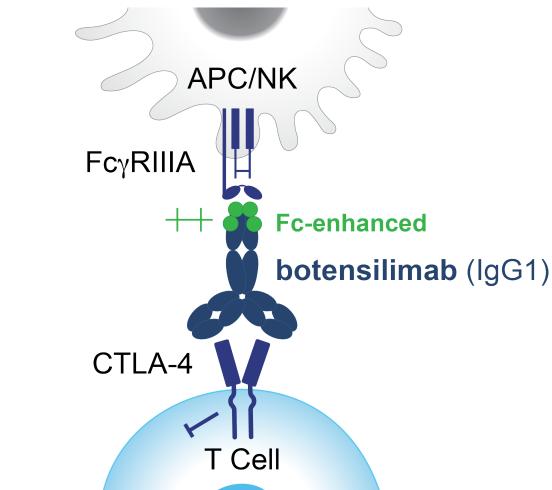




## NOVEL IMMUNOTHERAPY AGENTS

## botensilimab

#### **Fc-enhanced CTLA-4 Inhibitor**

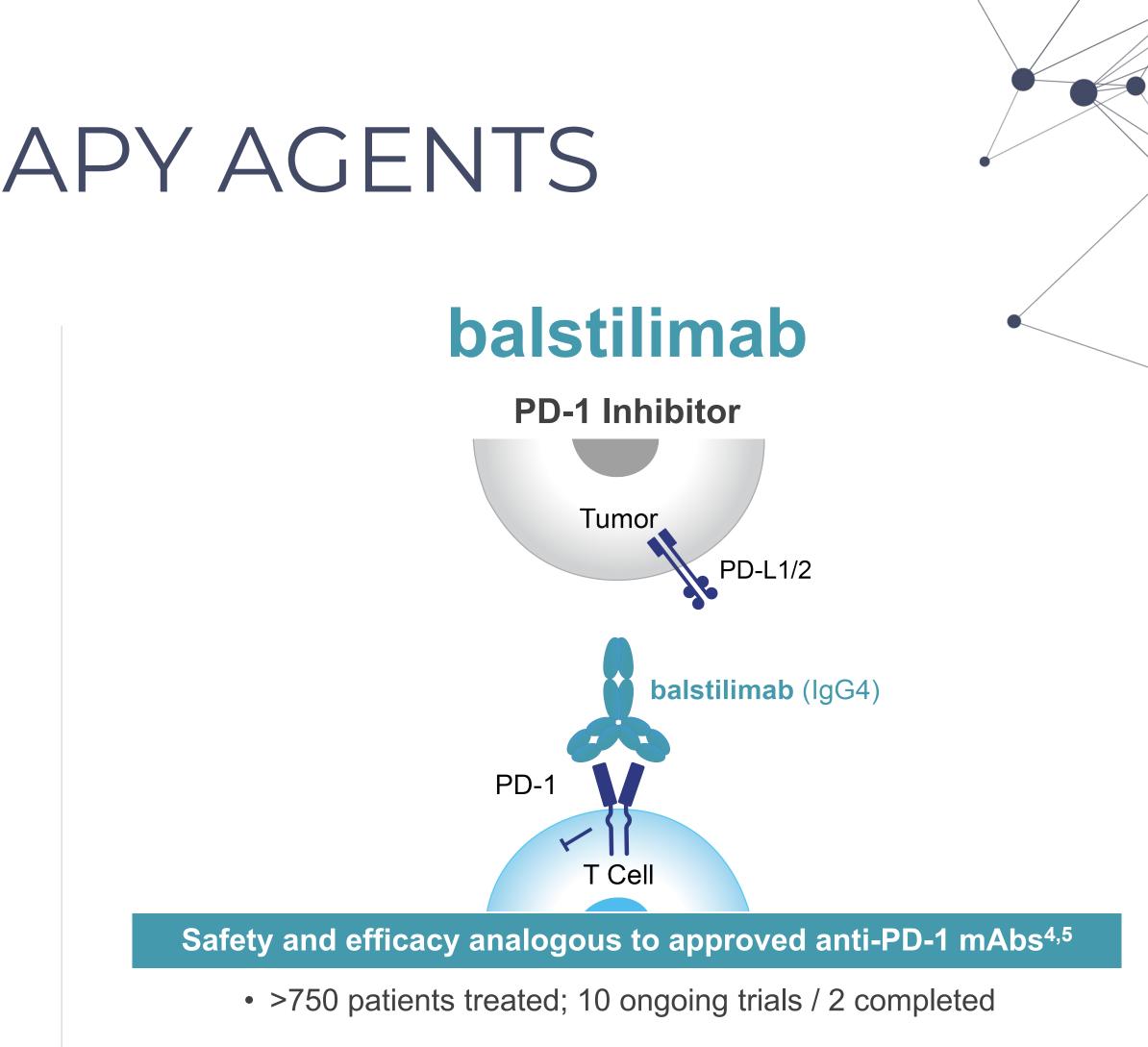


#### Active in cold and I-O refractory tumors<sup>1</sup>:

- $\uparrow$  T cell priming, expansion, memory<sup>2,3</sup>

- UCC Complement mediated toxicity

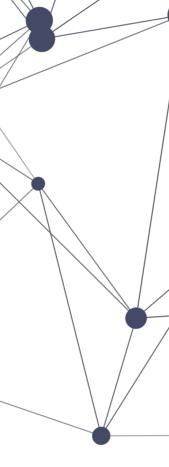
1. El-Khoueiry AB, et al. SITC 2021 Annual Meeting. Poster #479. 2. Waight et al. Cancer Cell. 2018;33(6): 1033-1047. 3. Data on File. Agenus, Inc. November 2022. 4. O'Malley, et al. Gynecol Oncol. 2021; 163: 274-280. 5. O'Malley et al, J Clin Oncol. 2022; 40(7): 762-771.



• Enhanced T cell activation and effector function

• Complete blocker of PD-1-PD-L1/2 interactions







## C-800-01 STUDY

#### POPULATION

### **Key Eligibility**

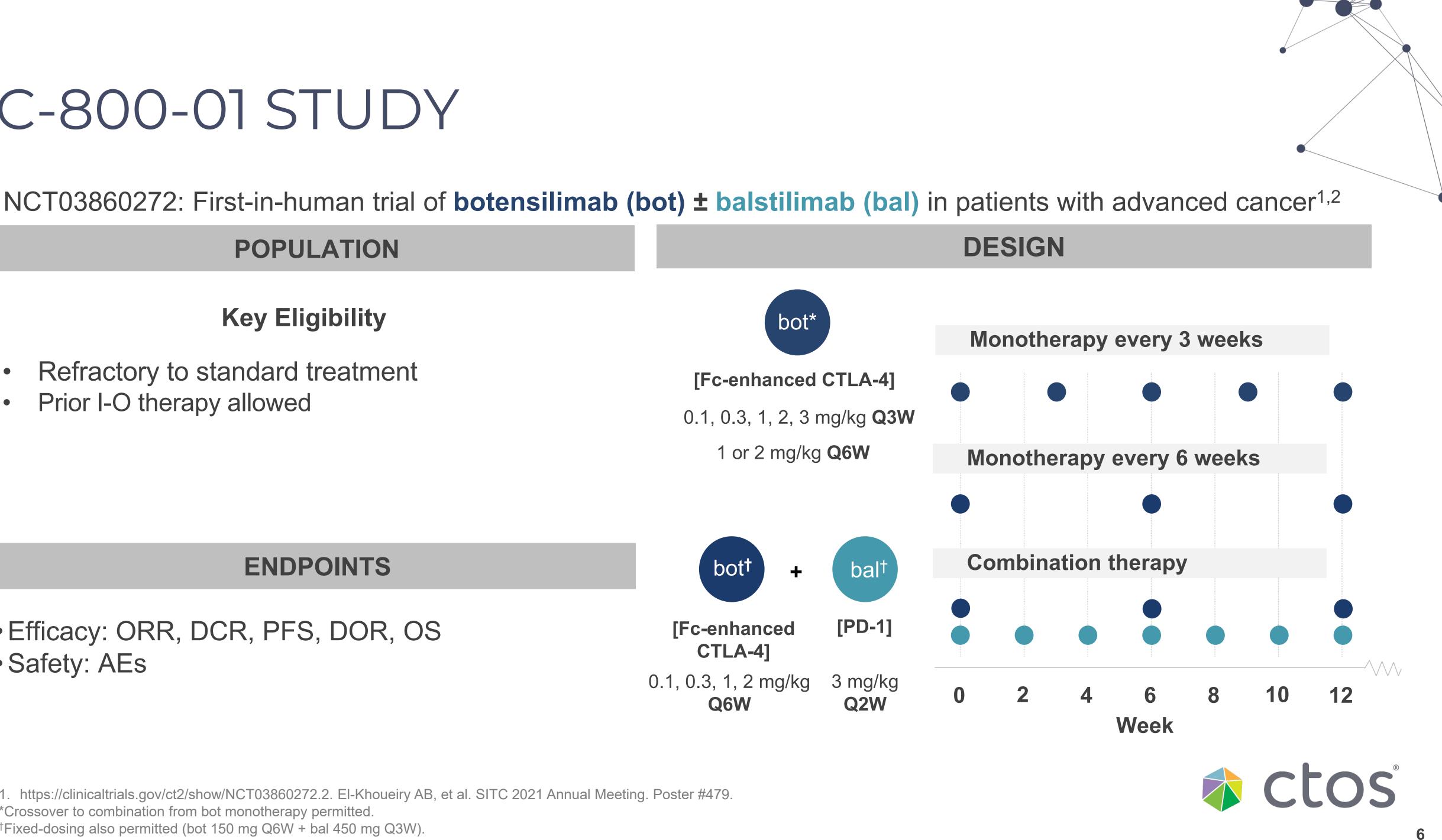
- Refractory to standard treatment
- Prior I-O therapy allowed

## **ENDPOINTS**

• Efficacy: ORR, DCR, PFS, DOR, OS • Safety: AEs

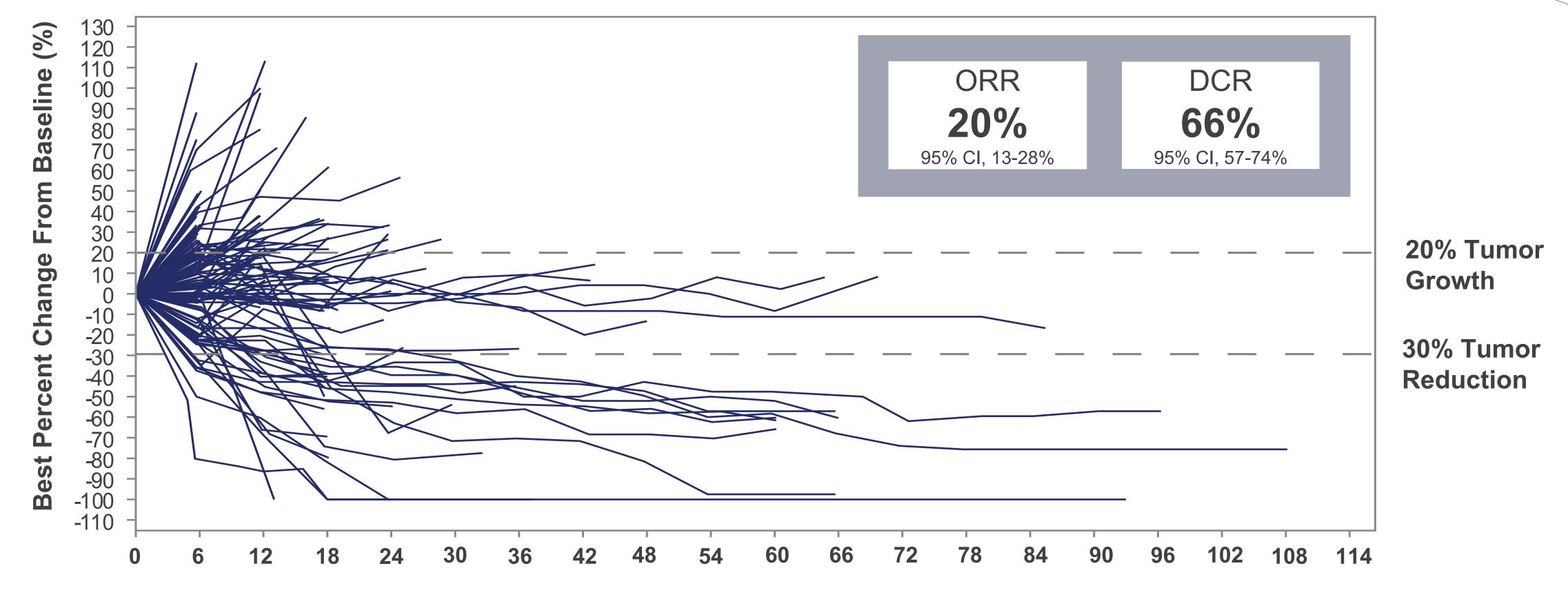
1. https://clinicaltrials.gov/ct2/show/NCT03860272.2. El-Khoueiry AB, et al. SITC 2021 Annual Meeting. Poster #479. \*Crossover to combination from bot monotherapy permitted.

<sup>†</sup>Fixed-dosing also permitted (bot 150 mg Q6W + bal 450 mg Q3W).





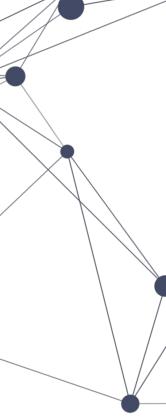
## BOTENSILIMAB/BALSTILIMAB EFFICACY IN COLD AND I-O RESISTANT TUMORS (N=125)\*



Wilky BA, et al. SITC 2022 Annual Meeting. Oral Presentation #778.

\*Evaluable patients were treated with bot + bal as of 19 May,2022, and had  $\geq$ 1 Q6W imaging assessment. Includes combination patients who received 1 or 2 mg/kg bot Q6W plue 450 mg bet Q plus 3 mg/kg bal Q2W, fixed dose patients (150 mg bot Q6W plus 450 mg bal Q3W), and rescue patients who crossed over from monotherapy to combination therapy.

#### Weeks on Study







## CLINICAL RESPONSES IN SELECT EXPANSION COHORTS

	MSS CRC (n=59)
ORR, %	22% (95% CI, 12-35%)*
BOR, n (%)	
CR	1 (2)
PR	12 (20)
SD	30 (51)
PD	16 (27)
DCR (CR + PR + SD), %	<b>73%</b> (95% CI, 60-84%)
Median DOR, months	NR (95% CI, 2.8-NR)
Median F/U, months	6.2 (Range, 1.6-29.3)

Wilky BA, et al. SITC 2022 Annual Meeting. Oral Presentation #778. \*Includes unconfirmed responses.

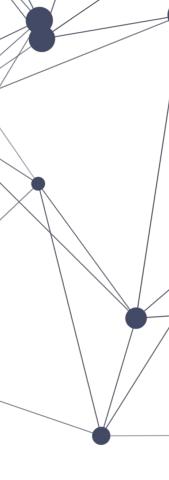
<sup>†</sup>Includes one additional evaluable patient since the data cutoff (ongoing confirmed PR).

Select Expansion Cohorts	
Ovarian (n=19)	Anti-PD-(L)1 R/R NSCLC (n=5)
26% (95% CI, 9-51%)*	<b>60%</b> <sup>†</sup>
1 (5)	0 (0)
4 (21)	3 (60)‡
7 (37)	1 (20)
7 (37)	1 (20)
63% (95% CI, 38-84%)	80%‡
NR (95% CI, 4.2-NR)	NR (95% CI, 4.5-NR)

6.5 (Range, 2.0-24.0)

5.3 (Range, 3.3-16.2)







## C-800-01 SARCOMA ANALYSIS POPULATION

### NCT03860272: First-in-human trial of **bot ± bal** in patients with advanced cancer<sup>1,2</sup>

#### POPULATION

#### **Evaluable Combination Patients\***

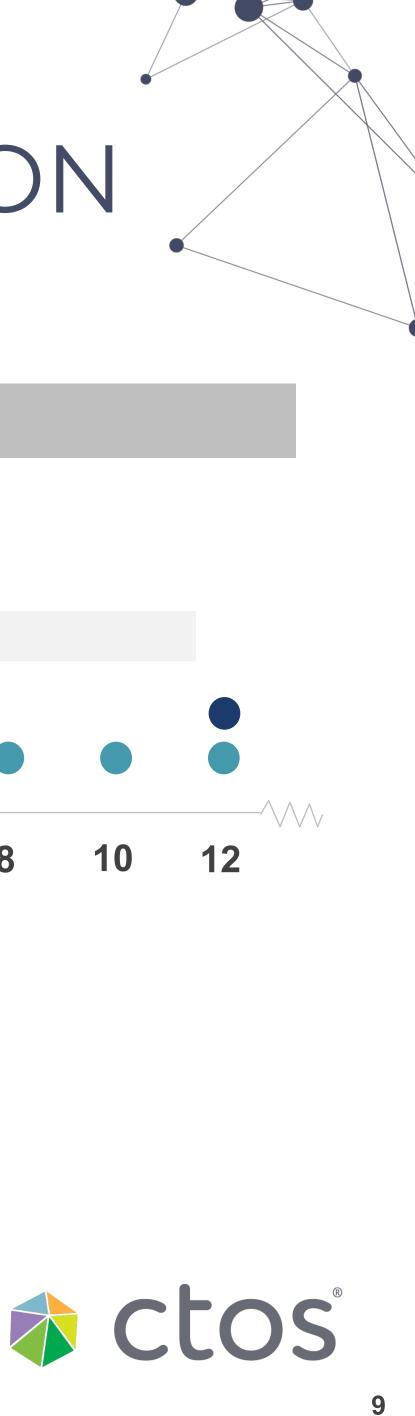
**13** treated with 1 or 2 mg/kg bot + bal as of 16 June 2022 with ≥1 Q6W imaging assessment

- **11** *de novo* combination patients
- 2 rescue patients who received combination therapy after bot monotherapy

1. https://clinicaltrials.gov/ct2/show/NCT03860272. 2. El-Khoueiry AB, et al. SITC 2021 Annual Meeting. Poster #479. \*ITT population (N=20): Seven additional patients were treated as of 16 June 2022 but were not included in the analysis as they did not have at least one 6-week imaging scan either due to progression (n=5), withdrawal of consent (n=1), or due to coming off study due to an unrelated AE (n=1). <sup>†</sup>Fixed-dosing also permitted (bot 150 mg Q6W + bal 450 mg Q3W).



DESIGN

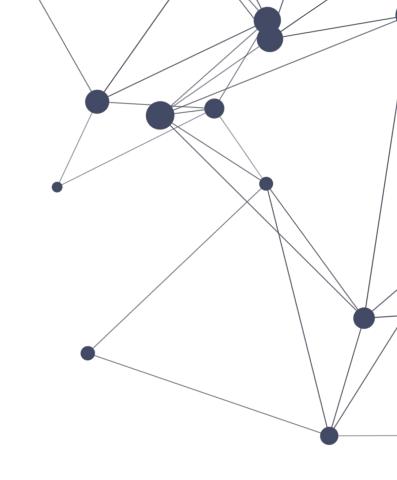


## PATIENT CHARACTERISTICS (N=13)

Age, median (range)       54 (41-80)         Sex, n (%)          Male       7 (54)         Female       6 (46)         ECOG PS at baseline, n (%)          0       9 (69)         1       4 (31)         Prior lines of therapy, n (%)          Median (range)       3 (0-4)         2       1 (8)         3       4 (31)         Prior I-O, n (%)'       4 (31)         Microsatellite stable status, n (%)†       11/11 (100)         TMB >10 mutations/Mb, n/N (%)†       2/11 (18)         PD-L1 (≥1%), n/N (%)†       5/11 (45)         Sarcoma subtype, n (%)          Cutaneous angiosarcoma       5 (38)         Visceral angiosarcoma (leiomyosarcomatous differentiation)       1 (8)         Liposarcoma (leiomyosarcomatous differentiation)       1 (8)         Leiomorphic sarcoma       2 (15)         Pleiomorphic sarcoma       1 (8)	Characteristic	Overall (N=13)
Male       7 (54)         Female       6 (46)         ECOG PS at baseline, n (%)       9 (69)         0       9 (69)         1       4 (31)         Prior lines of therapy, n (%)       3 (0-4)         2       1 (8)         3       4 (31)         4       4 (31)         Prior I-O, n (%)*       4 (31)         Microsatellite stable status, n (%)*       11/11 (100)         TIMB >10 mutations/Mb, n/N (%)*       2/11 (18)         PD-L1 (≥1%), n/N (%)*       5/11 (45)         Sarcoma subtype, n (%)       Cutaneous angiosarcoma         Visceral angiosarcoma       5 (38)         Visceral angiosarcoma       4 (31)         Liposarcoma (leiomyosarcomatous differentiation)       1 (8)         Leiomyosarcoma       2 (15)	Age, median (range)	54 (41-80)
Female       6 (46)         ECOG PS at baseline, n (%)       9 (69)         1       4 (31)         Prior lines of therapy, n (%)       3 (0-4)         2       1 (8)         3       4 (31)         4       4 (31)         Prior I-O, n (%)*       4 (31)         Microsatellite stable status, n (%)†       11/11 (100)         TMB >10 mutations/Mb, n/N (%)†       2/11 (18)         PD-L1 (≥1%), n/N (%)†       5/11 (45)         Sarcoma subtype, n (%)       5 (38)         Visceral angiosarcoma       5 (38)         Visceral angiosarcoma       4 (31)         Liposarcoma (leiomyosarcomatous differentiation)       1 (8)         Leiomyosarcoma       2 (15)	Sex, n (%)	
ECOG PS at baseline, n (%)       9 (69)         1       4 (31)         Prior lines of therapy, n (%) $3 (0-4)$ 2       1 (8)         3       4 (31)         4       4 (31)         Prior I-O, n (%)*       4 (31)         Microsatellite stable status, n (%)†       11/11 (100)         TMB >10 mutations/Mb, n/N (%)†       2/11 (18)         PD-L1 (≥1%), n/N (%)†       5/11 (45)         Sarcoma subtype, n (%)       Cutaneous angiosarcoma         Cutaneous angiosarcoma       5 (38)         Visceral angiosarcoma       4 (31)         Liposarcoma (leiomyosarcomatous differentiation)       1 (8)         Leiomyosarcoma       2 (15)	Male	7 (54)
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Prior lines of therapy, n (%)         Median (range) $3 (0-4)$ 2       1 (8)         3       4 (31)         4       4 (31)         Prior I-O, n (%)*       4 (31)         Microsatellite stable status, n (%)†       11/11 (100)         TMB >10 mutations/Mb, n/N (%)†       2/11 (18)         PD-L1 (≥1%), n/N (%)†       5/11 (45)         Sarcoma subtype, n (%)       5 (38)         Cutaneous angiosarcoma       5 (38)         Visceral angiosarcoma       4 (31)         Liposarcoma (leiomyosarcomatous differentiation)       1 (8)         Leiomyosarcoma       2 (15)	0	9 (69)
Median (range) $3 (0-4)$ 2       1 (8)         3       4 (31)         4       4 (31)         Prior I-O, n (%)*       4 (31)         Microsatellite stable status, n (%)†       11/11 (100)         TMB >10 mutations/Mb, n/N (%)†       2/11 (18)         PD-L1 (≥1%), n/N (%)†       5/11 (45)         Sarcoma subtype, n (%)       5 (38)         Visceral angiosarcoma       5 (38)         Visceral angiosarcoma       4 (31)         Liposarcoma (leiomyosarcomatous differentiation)       1 (8)         Leiomyosarcoma       2 (15)	1	4 (31)
2       1 (8)         3       4 (31)         4       4 (31)         Prior I-O, n (%)*       4 (31)         Microsatellite stable status, n (%)†       11/11 (100)         TMB >10 mutations/Mb, n/N (%)†       2/11 (18)         PD-L1 (>1%), n/N (%)†       5/11 (45)         Sarcoma subtype, n (%)       5 (38)         Cutaneous angiosarcoma       5 (38)         Visceral angiosarcoma       4 (31)         Liposarcoma (leiomyosarcomatous differentiation)       1 (8)         Leiomyosarcoma       2 (15)	Prior lines of therapy, n (%)	
3       4 (31)         4       4 (31)         Prior I-O, n (%)*       4 (31)         Microsatellite stable status, n (%)†       11/11 (100)         TMB >10 mutations/Mb, n/N (%)†       2/11 (18)         PD-L1 (≥1%), n/N (%)†       5/11 (45)         Sarcoma subtype, n (%)       5 (38)         Cutaneous angiosarcoma       5 (38)         Visceral angiosarcoma       4 (31)         Liposarcoma (leiomyosarcomatous differentiation)       1 (8)         Leiomyosarcoma       2 (15)	Median (range)	3 (0-4)
4 $4 (31)$ Prior I-O, n (%)* $4 (31)$ Microsatellite stable status, n (%)* $11/11 (100)$ TMB >10 mutations/Mb, n/N (%)* $2/11 (18)$ PD-L1 (≥1%), n/N (%)* $5/11 (45)$ Sarcoma subtype, n (%) $5 (38)$ Cutaneous angiosarcoma $5 (38)$ Visceral angiosarcoma $4 (31)$ Liposarcoma (leiomyosarcomatous differentiation) $1 (8)$ Leiomyosarcoma $2 (15)$	2	1 (8)
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Microsatellite stable status, n (%) <sup>†</sup> $11/11 (100)$ TMB >10 mutations/Mb, n/N (%) <sup>†</sup> $2/11 (18)$ PD-L1 (>1%), n/N (%) <sup>†</sup> $5/11 (45)$ Sarcoma subtype, n (%) $5 (38)$ Cutaneous angiosarcoma $5 (38)$ Visceral angiosarcoma $4 (31)$ Liposarcoma (leiomyosarcomatous differentiation) $1 (8)$ Leiomyosarcoma $2 (15)$	4	4 (31)
TMB >10 mutations/Mb, n/N (%)† $2/11 (18)$ PD-L1 (≥1%), n/N (%)† $5/11 (45)$ Sarcoma subtype, n (%) $5 (38)$ Cutaneous angiosarcoma $5 (38)$ Visceral angiosarcoma $4 (31)$ Liposarcoma (leiomyosarcomatous differentiation) $1 (8)$ Leiomyosarcoma $2 (15)$	Prior I-O, n (%)*	4 (31)
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Liposarcoma (leiomyosarcomatous differentiation)1 (8)Leiomyosarcoma2 (15)	Cutaneous angiosarcoma	5 (38)
Leiomyosarcoma 2 (15)	Visceral angiosarcoma	4 (31)
	Liposarcoma (leiomyosarcomatous differentiation)	1 (8)
Pleiomorphic sarcoma 1 (8)	Leiomyosarcoma	2 (15)
	Pleiomorphic sarcoma	1 (8)

\*Including prior PD-(L)1 and/or CTLA-4 inhibitors, and PD-1/IL-2.

<sup>†</sup>MSS, TMB, and PD-L1 status was unknown in two patients (TPS or CPS ≥1%).









## EFFICACY (N=13)

ORR,	%
BOR,	n (%)
CR*	

PR

SD

PD

**DCR (CR + PR + SD)**, %

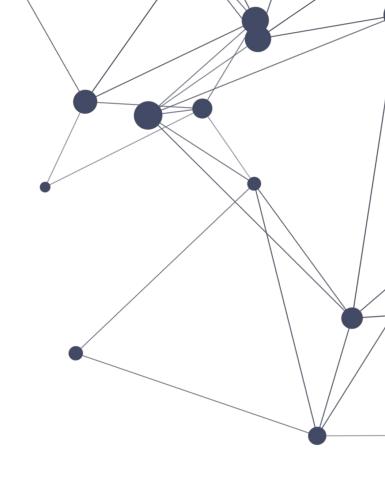
#### Median DOR, months

#### Median OS, months

12-months OS, months

## Median F/U, months

\*One patient with an unconfirmed CR. All other responses confirmed.



#### Overall (N=13)

**46%** (95% CI, 19-75%)

1 (8)

5 (38)

3 (23)

4 (31)

**69%** (95% CI, 39-91%)

NR (95% CI, 1.9-NR)

NR (95% CI, 3.2-NR)

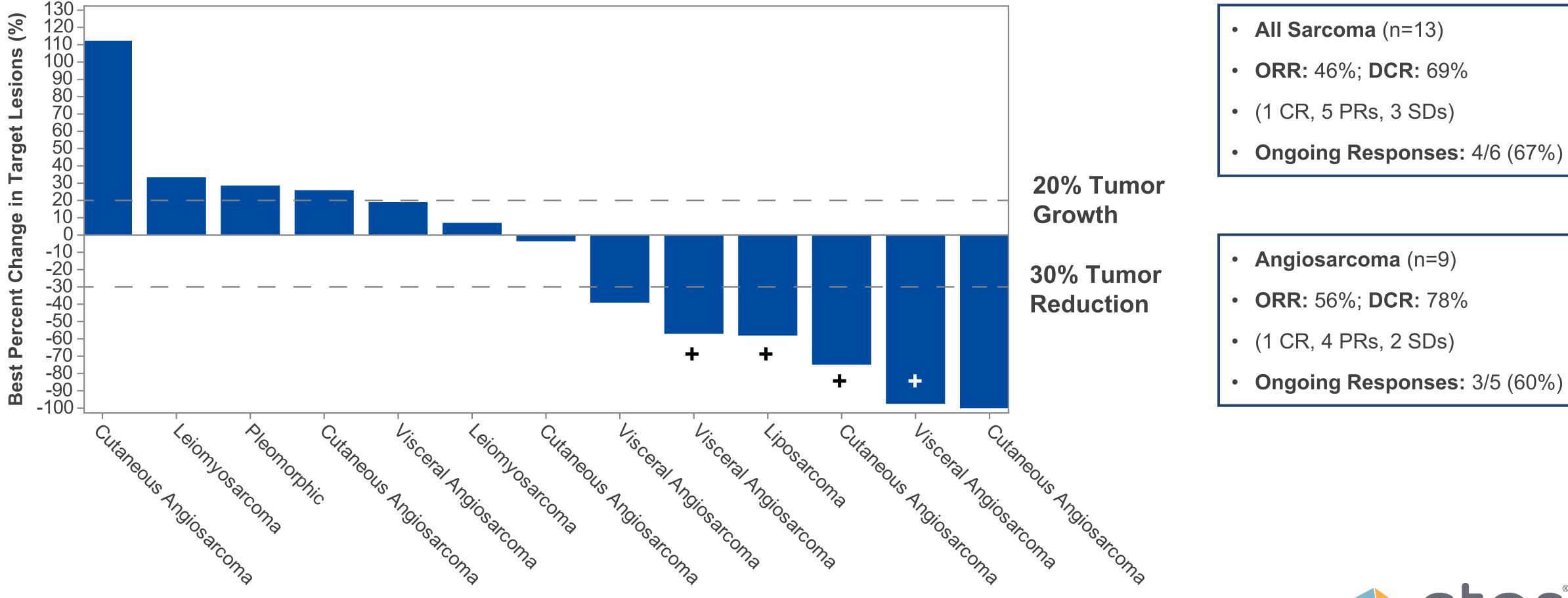
77% (95% CI, 35-94%)

4.2 (Range, 1.1-18.9)



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# BOTENSILIMAB PROMOTES RESPONSES IN SUBTYPES UNRESPONSIVE TO I-O (N=13)

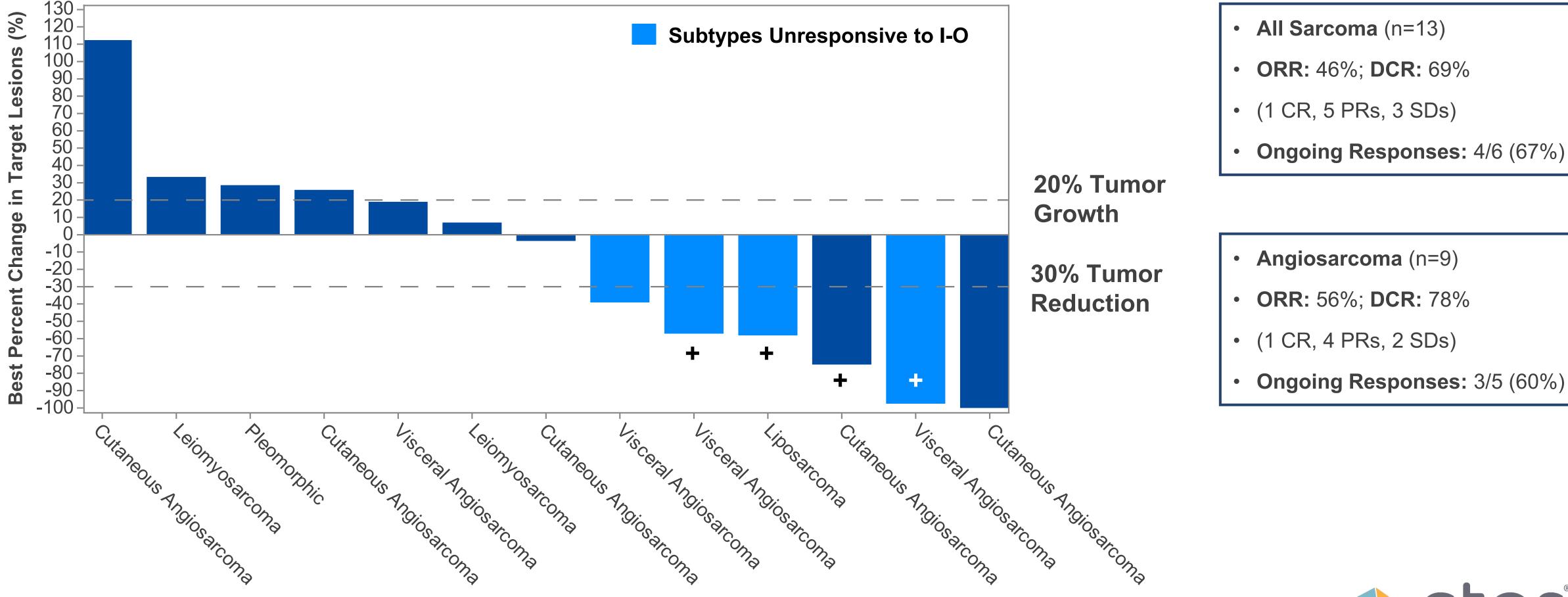




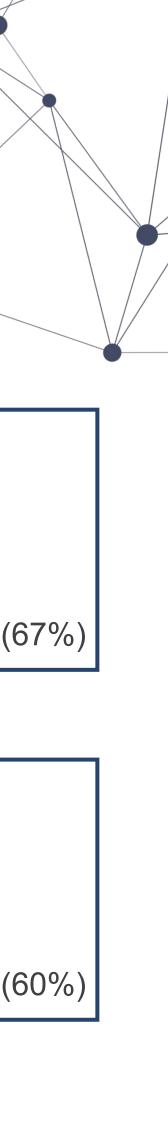




# BOTENSILIMAB PROMOTES RESPONSES IN SUBTYPES UNRESPONSIVE TO I-O (N=13)

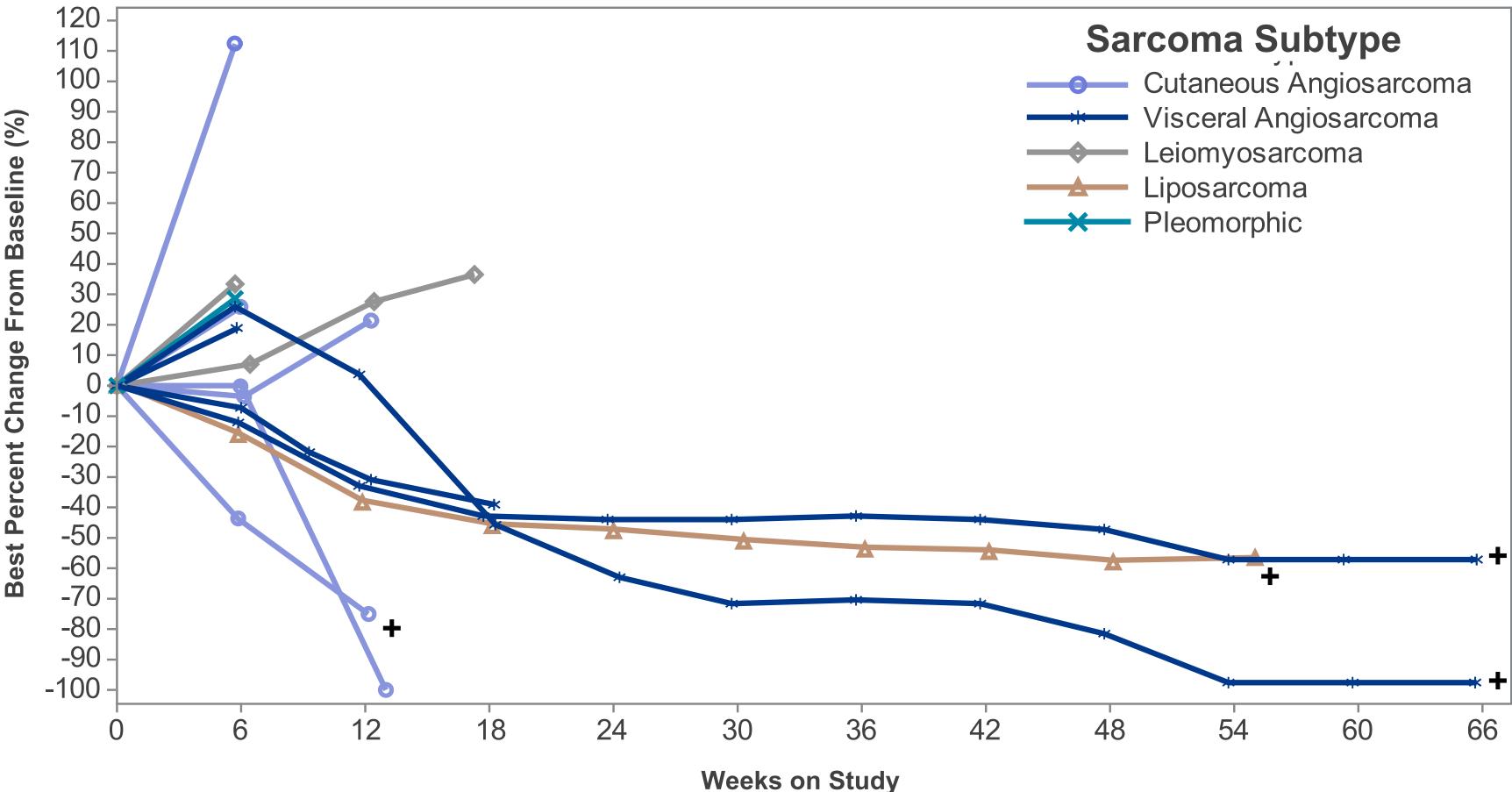








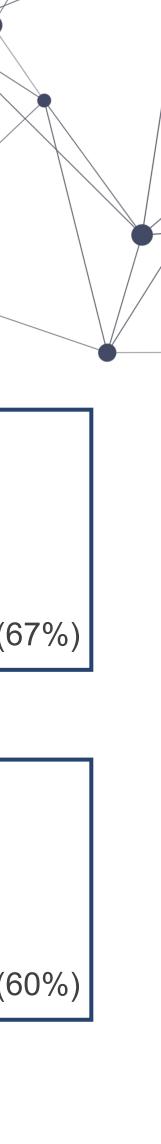
## BOTENSILIMAB PROMOTES DURABLE RESPONSES IN MULTIPLE SUBTYPES (N=13)



+ = Ongoing response.

- All Sarcoma (n=13)
- ORR: 46%; DCR: 69%
- (1 CR, 5 PRs, 3 SDs)
- Ongoing Responses: 4/6 (67%)
- Angiosarcoma (n=9)
- ORR: 56%; DCR: 78%
- (1 CR, 4 PRs, 2 SDs)
- Ongoing Responses: 3/5 (60%)





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## SAFETY TRAESOF ANY GRADE IN ≥2 PATIENTS (N=13)

Characteristic	All Grade
Any TRAE	12 (92)
GASTROINTESTINAL	
Diarrhea/colitis	6 (46)
Nausea	3 (23)
CONSTITUTIONAL	
Fatigue	5 (38)
Chills	4 (31)
Decreased appetite	3 (23)
Pyrexia	2 (15)
MUSCULOSKELETAL	
Myalgia	5 (38)
↑ CPK	2 (15)
SKIN	
Rash	5 (38)
ENDOCRINE	
Hypothyroidism/hyperthyroidism	2 (15)
BLOOD	
Lymphopenia	2 (15)
EYE	
Eye pain	2 (15)
RESPIRATORY	
Cough	2 (15)

Grade 3	
3 (23)	
2 (15) 0 (0)	
4 (0)	
1 (8) 0 (0) 0 (0) 0 (0)	Similar safety profile to the larger trial overall
	No hypophysitis, pneumonitis or
1 (8) 0 (0)	myocarditis
1 (8)	No grade 4 or 5 TRAEs
0 (0)	Discontinuation due to a TRAE:
1 (0)	8% bot only
1 (8)	<ul> <li>8% bot and bal</li> </ul>
1 (8)	
O(0)	

0 (0)

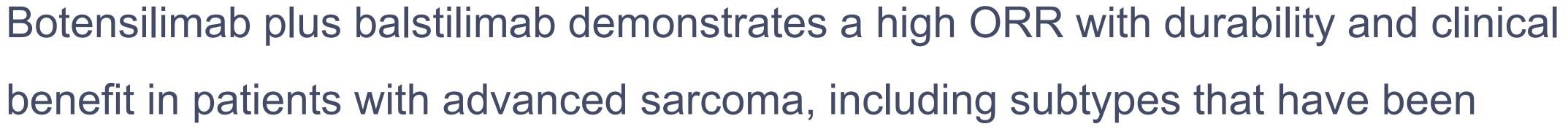






## SUMMARY

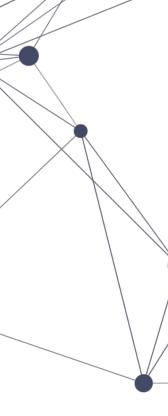
- unresponsive to I-O
- The combination is well tolerated and appears differentiated from first-gen CTLA-4based regimens, with less high-grade visceral toxicity outside of the GI tract, consistent with its molecular design<sup>1,2</sup>
- cohort
- A Phase 2 trial in sarcoma is planned for 2023



The current Phase 1 C-800-01 study is actively enrolling an expansion sarcoma









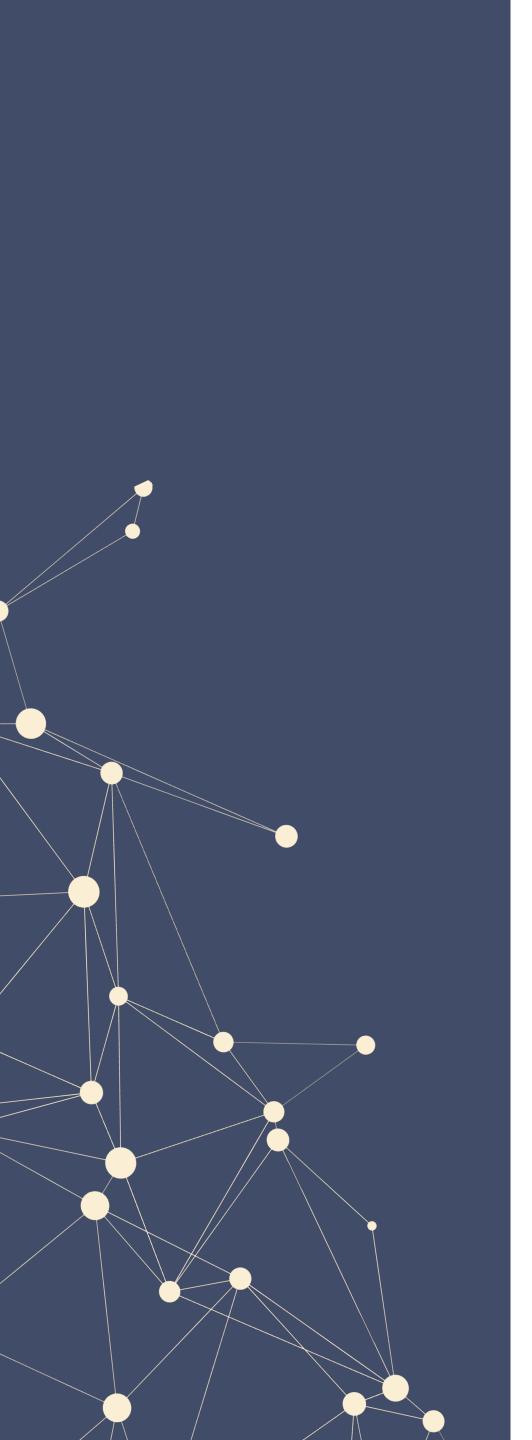
## ACKNOWLEDGEMENTS

Agenus Inc. funded and is the sponsor of this study.

The authors would like to thank the patients and their families for participating in the C-800-01 study, as well as the trial coordinators and investigators for their contributions.







## ABBREVIATIONS

AE, adverse event **APC**, antigen presenting cell **bal**, balstilimab **BOR**, best overall response **bot**, botensilimab **CPS**, combined positive score **CR**, complete response CRC, colorectal cancer **CPK**, creatine phosphokinase CTLA-4, cytotoxic T-lymphocyte and **DC**, dendritic cell **DCR**, disease control rate **DOR**, duration of response ECOG, Eastern Cooperative Oncold **Fc**, fragment crystallizable FcyRIIIA, Fragment crystallizable ga **F/U**, follow-up **ICI**, immune checkpoint inhibitor **IgG**, immunoglobulin G **I-O**, immunotherapy

	irAE, immune-related adverse event
	ITT, intention-to-treat
	mAb, monoclonal antibody
	MSS, microsatellite stable
	NK, natural killer
	NR, not reached
	ORR, objective response rate
	<b>OS</b> , overall survival
	PD-1, programmed death receptor-1
ntigen-4	PD-L1/2, programmed death-ligand 1/2
	PFS, progression-free survival
	PR, partial response
	PS, performance status
logy Group	QXW, every X weeks
	R/R, relapsed/refractory
gamma receptor IIIA	SD, stable disease
	TMB, tumor mutational burden
	<b>TPS</b> , tumor proportion score
	TRAE, treatment-related adverse event
	Treg, regulatory T cell

