Phase 2 Trial of Brentuximab Vedotin (BV) with Pembrolizumab (Pembro) in Patients with Metastatic Non-Small Cell Lung **Cancer or Metastatic Cutaneous Melanoma** After Progression on Anti-PD-1 Therapy

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Objectives

To present the efficacy, safety, and exploratory biomarker data from an ongoing phase 2 study evaluating the efficacy and safety of BV+pembro in anti-PD-1 refractory solid tumors (SGN35-033).

Conclusions

Treatment with BV+pembro was associated with selective depletion of Tregs and modulation of tumor microenvironment with increased CD8 T cell infiltration in responding patients.

The observed pharmacodynamic activity demonstrates the potential for BV to re-sensitize anti-PD-1 resistant/refractory tumors to PD-1 inhibitors through CD30-directed depletion of Treg cells. Preliminary data in an exploratory analysis suggests that higher response may be observed in patients with tumors with higher PD-L1 expression (data not shown); future analyses are required to confirm this association.

Anti-tumor responses were observed in patients with confirmed PD and prior CPI exposure within 90 days of receiving study treatment.

BV+pembro was well tolerated in patients with R/R metastatic NSCLC and metastatic cutaneous melanoma. No new safety signals were observed.

SGN35-033 is currently enrolling cohorts to investigate the activity of BV+pembro in frontline NSCLC and head and neck cancer (cohorts 5 and 6).

Background

- Anti-PD-1 therapy is the mainstay of frontline treatment for metastatic NSCLC and metastatic cutaneous melanoma.
- Putative mechanisms of resistance to PD-1 inhibitors include activated Tregs and other immunosuppressive cells within the TME.
- BV is hypothesized to selectively deplete a subset of activated effector intratumoral Tregs that express CD30, potentially re-sensitizing tumors to anti-PD-1 therapy. SGN35-033 (NCT04609566) is an ongoing, multiple cohort, multicenter, open-label trial evaluating the efficacy and safety of BV+pembro in anti-PD-1 refractory solid tumors.

Study Design and Endpoints



Exploratory Endpoints

• Biomarker analysis at baseline and during treatment, including T cells in peripheral blood by flow cytometry, Treg and CD8 T cells by IHC in tumor biopsies, and gene expression profiles by RNAseq in tumor biopsies

Primary refractory (cohorts 1 and 3): Progressed without prior objective response during or after prior PD-1 inhibitor therapy within 3 months or SD for <6 months. Patients who progress while on adjuvant anti-PD1 therapy were eligible to enroll. Adjuvant courses of therapy without measurable disease do not apply to the SD for at least 6 months criterion. Secondary refractory (cohorts 2 and 4): Progressed after having developed a prior objective response of CR/PR for at least 3 months or SD for at least 6 months For NSCLC patients only: For patients on CPI treatment for <90 days, iRECIST confirmation of PD at least 4 weeks from the initial PD was required For melanoma patients only: Treatment with a PD-1 CPI containing regimen within 90 days; iRECIST confirmation of PD at least 4 weeks from the initial PD was required Unless otherwise specified, all analyses reflect the FAS and include all subjects who received any amount of study medication (BV or pembrolizumab).

Results

Patient Demographics and Disease Characteristics

	Metastatic NSCLC			Metastatic cutaneous melanoma			
	Primary refractory (n=12)	Secondary refractory (n=43)	Total (n=55)	Primary refractory (n=17)	Secondary refractory (n=41)	Total (n=58)	
Age (years) median (range)	67.5 (49, 80)	67.0 (55, 86)	67.0 (49, 86)	59.0 (23, 83)	65.0 (25, 86)	64.5 (23, 86)	
Sex, n (%)							
Male	11 (92)	24 (56)	35 (64)	11 (65)	25 (61)	36 (62)	
Female	1 (8)	19 (44)	20 (36)	6 (35)	16 (39)	22 (38)	
ECOG performance status, n (%)							
0	6 (50)	16 (37)	22 (40)	13 (76)	26 (63)	39 (67)	
1	6 (50)	27 (63)	33 (60)	4 (24)	15 (37)	19 (33)	
Time from initial diagnosis to first dose (months)							
n	12	40	52	14	34	48	
Median	15.5	22.4	20.0	32.3	59.7	44.1	
Min, max	5, 41	8, 138	5, 138	6, 132	8, 259	6, 259	
Prior therapy, n (%)							
Prior ipilimumab/ anti-CTLA4 exposure	0	6 (14)	6 (11)	15 (88)	32 (78)	47 (81)	

Safety Summary, continued

- No new safety signals were observed.
- TESAEs were reported in 18 (33%) patients with metastatic NSCLC and in 29 (50%) patients with metastatic cutaneous melanoma. The most common TESAEs across all cohorts were vomiting (in 4% of patients) and cerebrovascular accident (in 3% of patients).
- TEAEs leading to treatment discontinuation were reported in 11 (20%) patients with metastatic NSCLC and in 8 (14%) patients with metastatic cutaneous melanoma.
- Treatment-emergent peripheral neuropathy events occurred in 27 (49%) patients with metastatic NSCLC and in 26 (45%) patients with metastatic cutaneous melanoma.
- 29 (26%) patients experienced immune-mediated TEAEs; 13 (12%) patients experienced events that were considered treatment-related. The most common (≥2 patients) treatment-related IMAEs were aspartate aminotransferase increased, hypothyroidism, pneumonitis, and rash maculopapular, all of which occurred in 2 patients each.
- 5 (4%) patients experienced TEAEs that led to death.
- The safety data are generally consistent with the approved safety profiles in the labels of BV and pembrolizumab.

Responding Patients Showed a Trend Toward Higher CD8 and Foxp3 Protein Expression in Baseline Tumor Biopsies



Y: responders (CR or PR); N: Patients did not achieve an objective response

Immune Response Pathway-related Gene Sets are Enriched in Responders as Compared to Non-responders in Baseline **Tumor Biopsies**



Efficacy Analyses

Data cutoff: 10AUG2023

- ORR was 8% to 22% across cohorts.
- DORs appear durable in some patients; however, data are immature and limited by small sample size.
- In an exploratory analysis, numerically higher response rates were observed in patients with tumors that had higher PD-L1 expression (data not shown).
- Most responders in cohorts 3 and 4 received study treatment within 90 days of receiving prior CPI therapy.

Antitumor Activity

	Metastatic NSCLC			Metastatic mela	cutaneous noma
	Primary refractory (n=12)	Secondary refractory (n=43)		Primary refractory (n=17)	Secondary refractory (n=41)
ORR (CR+PR) %,	8	14	iORR (iCR+iPR)ª %,	18	22
(95% CI)	(0.2, 38.5)	(5.3, 27.9)	(95% CI)	(3.8, 43.4)	(10.6, 37.6)
DCR (CR+PR+SD) %,	67	72	iDCR (iCR+iPR+iSD) ^a %,	71	80
(95% CI)	(34.9, 90.1)	(56.3, 84.7)	(95% CI)	(44.0, 89.7)	(65.1, 91.2)

^aOne patient with melanoma experienced pseudoprogression but consolidated to an iCR; hence iRECIST is reported for melanoma cohorts. Data cutoff: 10AUG2023

Duration of Response: Metastatic NSCLC and Metastatic Cutaneous Melanoma

Primary refractory metastatic NSCLC



Secondary refractory metastatic NSCLC





Primary refractory metastatic cutaneous

Response Partial response

Stable disease



melanoma

Primary and Secondary refractory metastatic NSCLC (cohorts 1 and 2): Response Criterion: CR, PR, SD and PD per RECIST v1.1.

Primary and Secondary refractory metastatic cutaneous melanoma (cohorts 3 and 4): Response Criterion:

Exploratory Biomarker Analyses

Transient Decrease in Tregs in Peripheral Blood in Patients Receiving BV+Pembro





The red numbers in each graph (C1 FDR) indicate False Discovery Rate comparing maximum change at C1D8 or C1D15 to C1D1; The numbers underneath each graph indicate the number of patient samples at each time point.

Paired Tumor Biopsies Collected from Responding Patients Suggest Increase in CD8 T Cell Infiltration After Treatment with **BV+Pembro**



CD8 expression increased in 4/4 (100%)

non-responders (iSD or iPD).

responding patients (iCR or iPR) and 6/13 (46%)



Gene Set Enrichment Analysis (GSEA) used ranked log2 fold change between responders (iCR+iPR) and non-responders (iSD+iCPD+iUPD+NE/NA).

Evaluated 15,712 gene sets, including "hallmark", oncogenic gene signature, gene ontology (GO CC, BP, and MF) and human phenotype ontology sets.

HP_NAIL_DYSPLAS

3000

6000

"Collapsed" enriched gene sets with common genes, plots show representative set names.

Antigen Presentation, T Cell Signaling, and Treg Genes in **Baseline Tumor Biopsies May be Associated with a Favorable Immune Environment for Response to BV+Pembro**

54 intersecting leading-edge genes among 58 positively enriched gene sets when comparing iCR+iPR to iSD+iUPD+iCPD +NE/NA using GSEA



iCR, iPR, iSD, iUPD, and iCPD per iRECIST. Indicates the patient with biopsy data shown in biopsy figure. Data cutoff: 10AUG2023

Safety Summary

All TEAEs Occurring at ≥20% Frequency and Grade ≥3 TEAEs **Occurring at ≥5% Frequency**

	Metastatic NSCLC (N=55)		Metastatic cutaneous melanoma (N=58)		
TEAEs, n (%)	Any grade (≥20%)	Grade ≥3 (≥5%)	Any grade (≥20%)	Grade ≥3 (≥5%)	
Any event	51 (93)	31 (56)	56 (97)	32 (55)	
Fatigue	28 (51)	3 (5)	24 (41)	3 (5)	
Nausea	28 (51)	3 (5)	20 (34)	_	
Peripheral sensory neuropathy	20 (36)	—	21 (36)	_	
Diarrhea	16 (29)	—	18 (31)	_	
Constipation	15 (27)	_	17 (29)	_	
Decreased appetite	13 (24)	3 (5)	_	_	
Dyspnea	16 (29)	_	_	_	
Vomiting	11 (20)	_	_	_	
Pruritus	11 (20)	_	_	_	
Acute kidney injury	_	3 (5)	_	_	
Aspartate aminotransferase increased	_	3 (5)	_	_	
Neutropenia	_	3 (5)	_		
Hypokalemia	_	_	_	4 (7)	
Hepatitis	_	_	_	3 (5)	

Note: "–" indicates TEAEs below 20% (all grade) or 5% (grade ≥3) threshold AEs were defined as treatment-emergent if they were newly occurring or worsened following treatment with either brentuximab vedotin or pembrolizumab. Data cutoff: 10AUG2023

Disclosures

SL: Travel expenses from Dava Oncology. MC: Employee of Merck, Inc. HL, JB, KR, SK, and BO are employees of Seagen, Inc.

CD8 T cells increased in 4/5 (80%) responding patients (iCR or iPR) and 7/14 (50%) non-responders (iSD or iPD). Note that some lines overlap.

Paired Biopsies from Patient with Metastatic Cutaneous Melanoma Demonstrate Increase in CD8 and Decrease in Foxp3 **After Treatment with BV+Pembro**



%IC+ = 80%



Baseline and C3D1 biopsy were collected from a melanoma patient who had complete response after pseudo-progression, as indicated by the star on the spider plot for secondary refractory cutaneous melanoma (cohort 4) patients. The brown color indicates cells stained positive for CD8 and Foxp3. The numbers below each graph show pathologist scoring. "% IC" means the percentage of cells out of total immune cells that stained positive.

Connectivity map of 54 leading edge genes showing protein-protein interactions



Abbreviations

N=67

AEs: adverse events; ALK: anaplastic lymphoma kinase; BOR3: best overall response grouped into 3 categories; BP: biological process; BV: brentuximab vedotin; CC: cellular component; CD: cluster of differentiation; CPI: checkpoint inhibitor; CR: complete response; DCR: disease control rate; DOR: duration of response; ECOG: Eastern Cooperative Oncology Group; EGFR: epidermal growth factor receptor; FAS: full analysis set; FDR: false discovery rate; GO: gene ontology; GSEA: gene set enrichment analysis; HP: human phenotype; iCPD: immune confirmed progression; IHC: immunohistochemistry; IMAEs: immune-mediated AEs; iRECIST: immune-RECIST; iUPD: immune unconfirmed progressive disease; LE: leading edge (gene identified using GSEA); MF: molecular function; NCI CTCAE: National Cancer Institute Common Terminology Criteria for Adverse Events; NE: not evaluable; NA: not available subjects including those who discontinued treatment with no post-baseline response; NES: normalized enrichment score; NSCLC: non-small cell lung cancer; ORR: objective response rate; padv: adjusted p-value; PD-L1: programmed cell death ligand 1; pembro: pembrolizumab; PFS: progression-free survival; PR: partial response; pval: p-value; RECIST: Response Evaluation Criteria in Solid Tumors; RNAseq: ribonucleic acid sequencing; R/R: relapsed/refractory; SD: stable disease; TEAE: treatment-emergent adverse events; TESAEs: treatment-emergent serious adverse events; TME: tumor microenvironment; Tregs: regulatory T-cells

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