Lessons from VISTA: New Strategies to Address an Important Immune Checkpoint

November 21, 2022

Guest Speaker: Robert Schreiber, Ph.D.

Andrew M. and Jane M. Distinguished Professor of Pathology and Immunology; Professor, Molecular Microbiology; and Director of the Bursky Center for Human Immunology and Immunotherapy Programs at the Washington University School of Medicine. He is also co-leader of the Tumor Immunology Program of Washington University's Siteman Comprehensive Cancer Center, an Associate Director of the Scientific Advisory Board to the Cancer Research Institute and Co-editor-in-Chief of the journal Cancer Immunology Research. Schreiber obtained his PhD in Immunology/Biochemistry at the State University of New York in Buffalo, New York, and received his postdoctoral training at The Scripps Research Institute in La Jolla, California. Sensei IOAB Member.

Sensei Presenters:

John Celebi Chief Executive Officer

Dr. Robert Pierce Chief R&D Officer

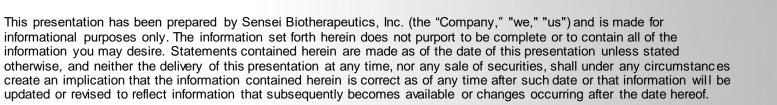
Dr. Ron Weitzman

Dr. Edward van der Horst

SVP, TMAb Antibody Development



Disclaimer



This presentation contains estimates and other statistical data made by independent parties and by us relating to market shares and other data about our industry. This presentation also contains "forward-looking" statements as that term is defined in the Private Securities Litigation Reform Act of 1995 that are based on our management's beliefs and assumptions and on information currently available to management. These forward-looking statements include, without limitation, expectations regarding the development of our product candidates; the potential safety profile of our product candidates; the potential benefits of our product candidates; the availability of data from our preclinical studies; the timing of selection of product candidates; the timing of IND submissions to the FDA; expected clinical development timelines; and our belief that our existing cash and cash equivalents will be sufficient to fund our operations at least into the first quarter of 2025.

When used in this presentation, the words and phrases "designed to," "may," "believes," "intends," "seeks," "anticipates," "plans," "estimates," "expects," "should," "assumes," "continues," "could," "will," "future" and the negative of these or similar terms and phrases are intended to identify forward-looking statements. Forward-looking statements involve known and unknown risks, uncertainties and other factors that may cause our actual results, performance or achievements to be materially different from any future results, performance or achievements expressed or implied by the forward-looking statements. Risks and uncertainties that may cause actual results to differ materially include uncertainties inherent in the development of therapeutic product candidates, such as preclinical discovery and development, conduct of clinical trials and related regulatory requirements, our reliance on third parties over which we may not always have full control, and other risk and uncertainties that are described in our Quarterly Report on Form 10-Q filed with the SEC on November 8, 2022 and our other Periodic Reports filed with the SEC. Forward-looking statements represent our management's beliefs and assumptions only as of the date of this presentation and include all matters that are not historical facts. Our actual future results may be materially different from what we expect. Except as required by law, we assume no obligation to update these forward-looking statements, even if new information becomes available in the future.

Certain information contained in this presentation relates to, or is based on, studies, publications, surveys and other data obtained from third-party sources and the Company's own internal estimates and research. While the Company believes these third-party sources to be reliable as of the date of this presentation, it has not independently verified, and makes no representation as to the adequacy, fairness, accuracy or completeness of, any information obtained from third-party sources. In addition, all of the market data included in this presentation involves a number of assumptions and limitations, and there can be no guarantee as to the accuracy or reliability of such assumptions. Finally, while we believe our own internal research is reliable, such research has not been verified by any independent source.



Ag	en	da
•		

Robert Schreiber, Ph.D. Professor, Washington University School of Medicine Member, Sensei Immuno-Oncology Advisory Board	VISTABiology
Edward van der Horst, Ph.D. SVP, TMAb Antibody Development, Sensei Bio	SNS-101 Overview & Preclinical Data Review
Robert Pierce, M.D. Chief R&D Officer, Sensio Bio	SNS-101 Translational Medicine & Clinical Development Plan
Ron Weitzman, M.D. Interim Chief Medical Officer, Sensei Bio	
Neil Canavan MD, KOL Network, LifeSci Advisors	Fireside Chat



Engineered Selectivity to Extend the Reach of Immuno-oncology Agents





Lack of Selectivity is a Major Obstacle to CI Innovation

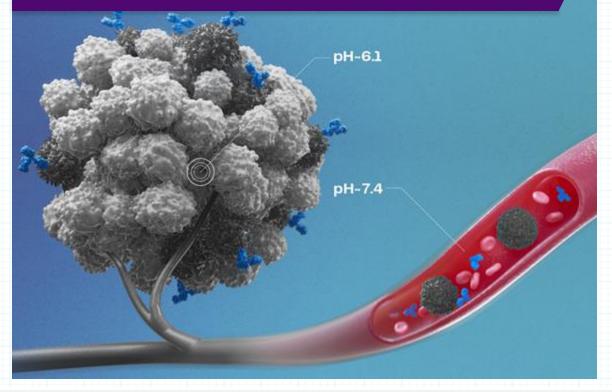
In	dustry Problem		Sensei's Solution
checkpoints t	antibodies target immu hat are highly expresse s, resulting in:	I	Conditionally active antibodies are selectively targeted to the tumor microenvironment, potentially providing:
Pharmacological sink of	due to on-target/off-tumor action effect requires higher and more frequ e to poor PK and dose-limiting toxiciti		Little or no toxicity due to selective on-target/on-tumor action Lower and less frequent doses by avoiding normal tissue binding Powerful activity selectively focused on the tumor microenvironment
Only one new check inhibitor has been ap since the original CT and PD-1/PD-L1 grou	oproved LA-4	Pembrolizumab (anti-PD-1)	Relatlimab (anti-LAG-3)
	2011	2014	2022



pH-sensitive Antibodies Have Potential to Selectively Bind Their Targets in the Low-pH Tumor Microenvironment

TMAb Platform

The tumor microenvironment of pH ~6 is lower than physiological pH of 7.4



Sensei's technology identifies pH-sensitive antibodies designed to bind only at the tumor

- Exploits the tumor microenvironment using pHselective properties
- Intended to alleviate undesirable properties:
 - Dose-limiting toxicities due to on-target/offtumor binding
 - Higher and more frequent dosing due to poor pharmacokinetics (Target-mediated Drug Disposition (TMDD))
- Bolsters specific activities
- Goal is to unlock previously undruggable immune targets



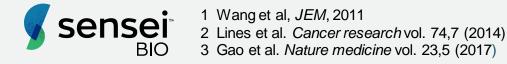
Robert Schreiber, Ph.D. VISTA Biology

VISTA (B7-H5) is recognized an important immune checkpoint and B7 family member that is expressed on myeloid cells, a hub of immunosuppressive activity, and is activated via binding to its receptor on T-cells (PSGL-1) at sub-physiologic pH

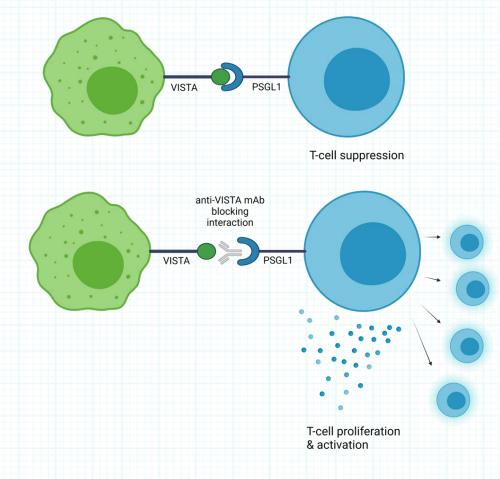


VISTA: A promising immune checkpoint target expressed predominantly on myeloid cells

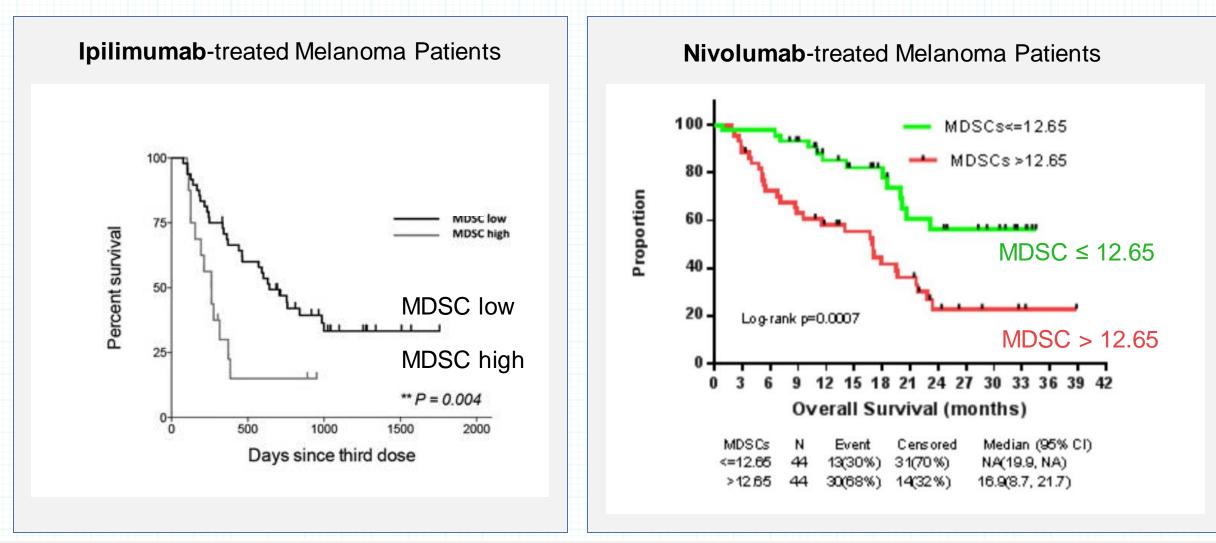
- VISTA (V-domain Ig-containing Suppressor of T cell Activation VISTA), also known as B7-H5 or PD-1H is a B7 family member protein with homology to PD-L1
- VISTA suppresses T cell activation¹
- Highly expressed on myeloid cells including macrophages and neutrophils; NK cells, T-regs and exhausted T cells have been reported to express VISTA²
- VISTA expression appears to be upregulated by hypoxia with a HIF1 α site identified in VISTA's promoter
- Inhibition of VISTA may "convert" myeloid cells to a proinflammatory/immune activating state; VISTA may "reverse signal" and play a role in enforcing myeloid immunosuppressive program
- Excellent therapeutic combinability with CTLA-4 or PD-1/PD-L1 T cell checkpoints³



VISTA is a Negative Regulator of T cell Function

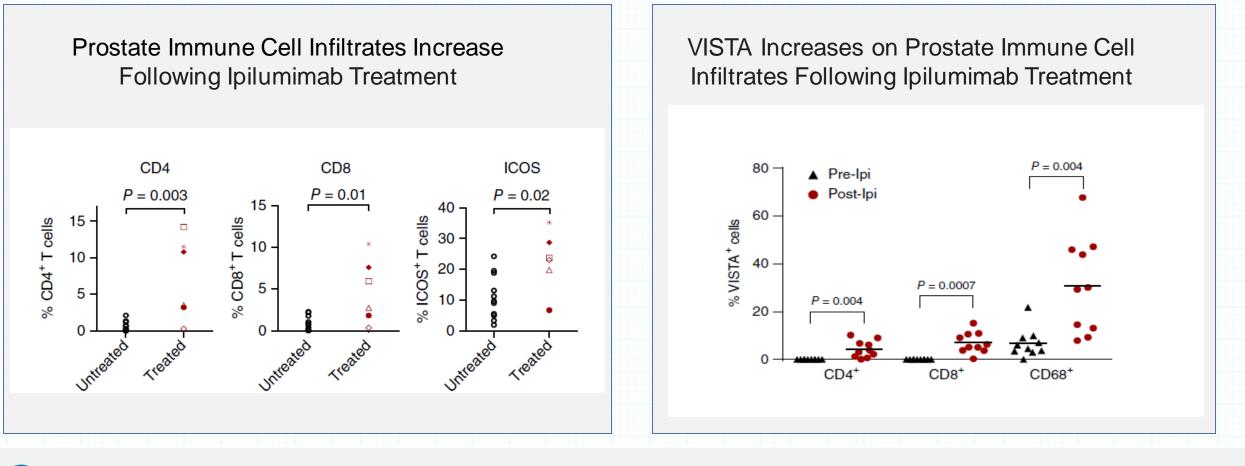


Patients with High Circulating Myeloid Cells Have Shown Lower Overall Survival When Treated with Checkpoint Blockade



VISTA Checkpoint May Be an Important Resistance Mechanism to Checkpoint Blockade

Can targeting VISTA augment T-cell checkpoint blockade in refractory tumors?

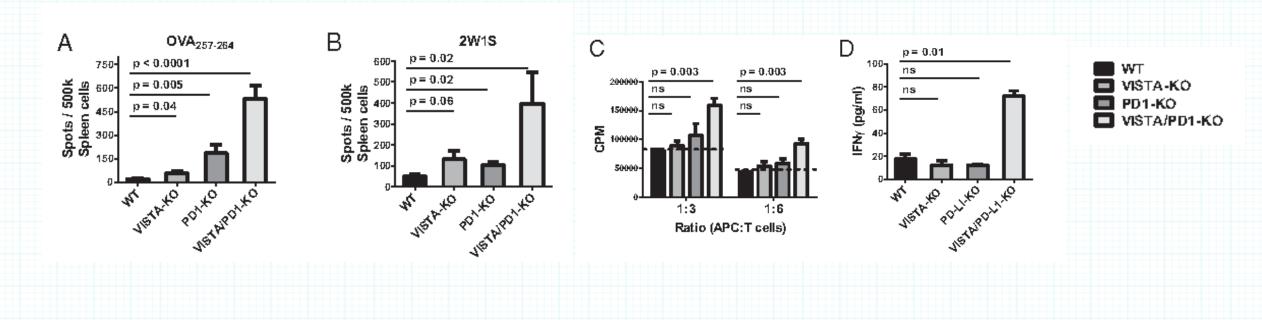




Blocking PD-1 and VISTA together leads to a synergistic increase in antigen-specific T cell responses in Vaccine Models

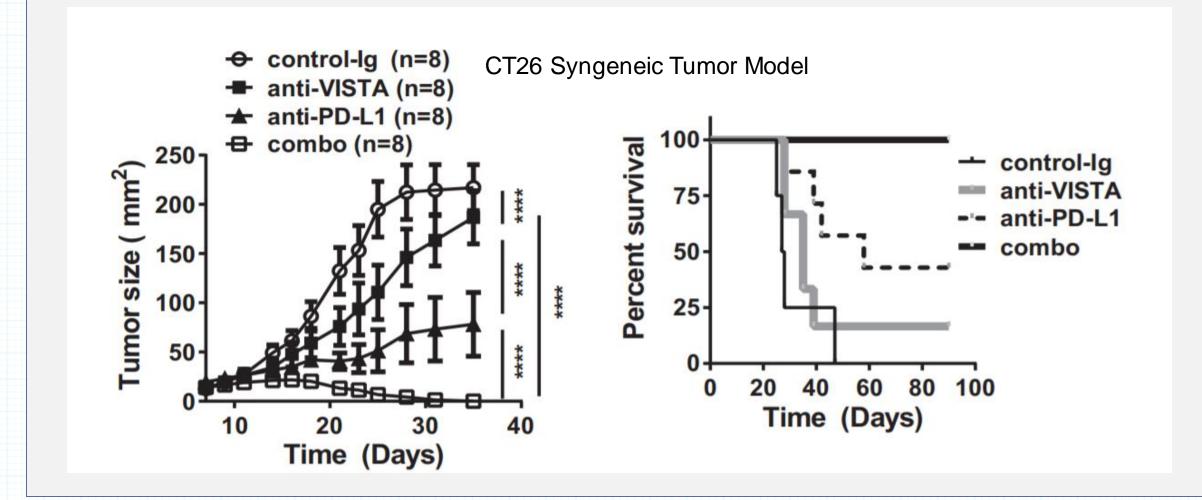
Immune-checkpoint proteins VISTA and PD-1 nonredundantly regulate murine T-cell responses

Jun Liu^{a,b}, Ying Yuan^{a,1}, Wenna Chen^a, Juan Putra^c, Arief A. Suriawinata^c, Austin D. Schenk^d, Halli E. Miller^a, Indira Guleria^e, Richard J. Barth^d, Yina H. Huang^c, and Li Wang^{a,2}





VISTA Blockade Synergizes With PD-1/L-1 Pathway Inhibition in a Tumor Model



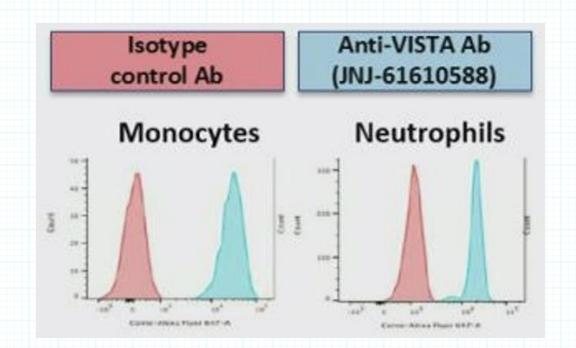
Why Has VISTA Been a Difficult Target?

Development of successful drugs targeting VISTA inhibition <u>had</u> been stymied in the past due to two major factors:

- Strong constitutive expression of VISTA on neutrophils and monocytes in the blood
 - Binding of drugs to VISTA+ cells in the blood at physiologic pH results in target-mediated drug disposition (TMDD) and clearance
 - Binding of mAbs to cells in blood can lead to cellular activation and cytokine release syndrome (CRS), particularly Fc-competent IgG1 antibodies
 - JNJ 61610588, an Fc-competent IgG1 monoclonal antibody, was the first anti-VISTA mAb was tested in the clinic (Phase 1, advanced solid tumors)
 - TMDD was pronounced and CRS was noted at subtherapeutic doses, including a grade 3 CRS at 0.3 mg/kg¹
 - Development was halted by JNJ; this antibody is now licensed to Curis and renamed CI-8893
- Lack of clarity on the identity of the critical counter-receptor responsible for T cell suppression
 - Putative interaction partners included VISTA itself (homotypic interaction), VSIG-3, VSIG-8, Syndecan-2, LRIG-1 and PSGL-1
 - Unclear how to select the right "inhibitor" if you don't know what interaction needs to be blocked
 - From an immunological perspective, PSGL-1 makes sense as it had been previously demonstrated to act as a T cell checkpoint, strongly suppressing T cell activation, concomitantly increasing PD-1 expression and exhaustion

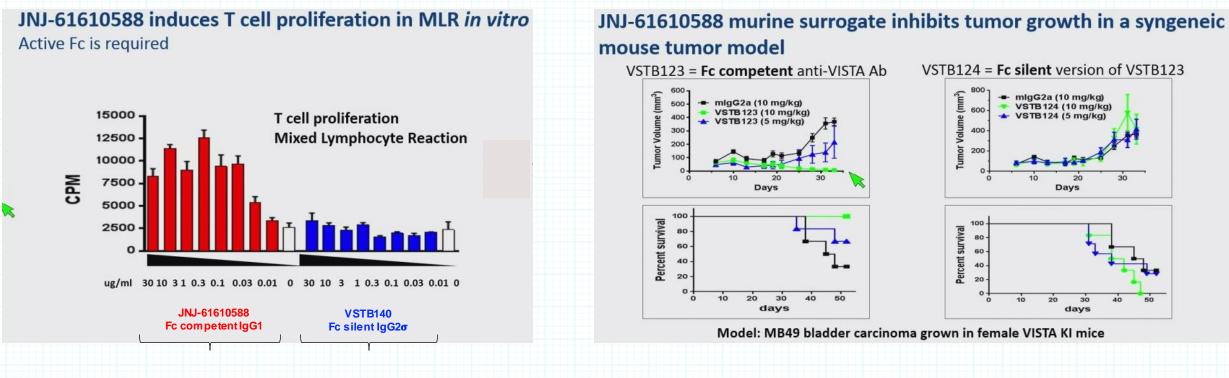
VISTA is Expressed at High Levels on Human Monocytes and Neutrophils

Flow Cytometry Analysis of VISTA Expression on Normal Human Peripheral Immune Cells





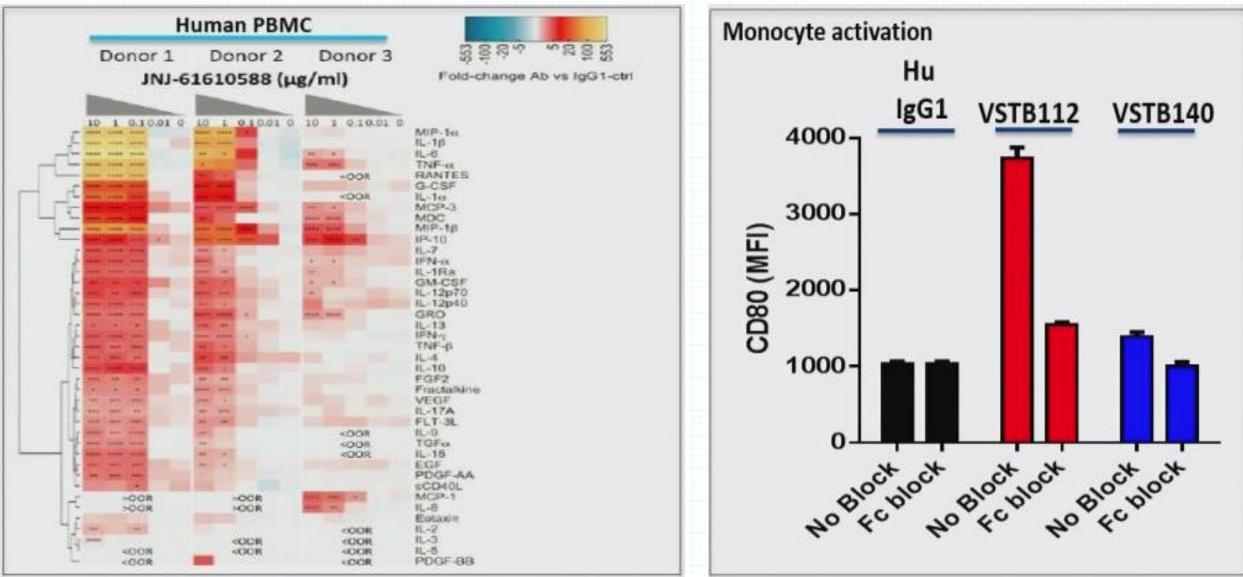
Fc-competent Framework is Required for Optimal Activity of Anti-VISTA Monoclonal Antibodies



Combination of on-target/off-tumor binding to neutrophils and monocytes in the blood and Fcmediated myeloid activation likely caused CRS seen with JNJ 61610588 in Phase 1 study

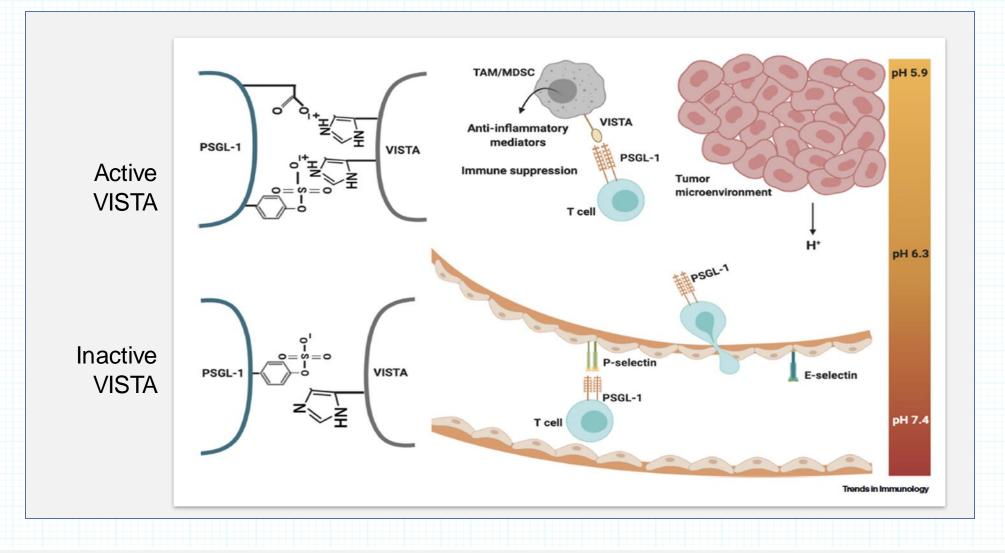


JNJ-61610588 or surrogate induced monocyte activation and cytokine release requires FC receptor cross-linking



sensei BIO

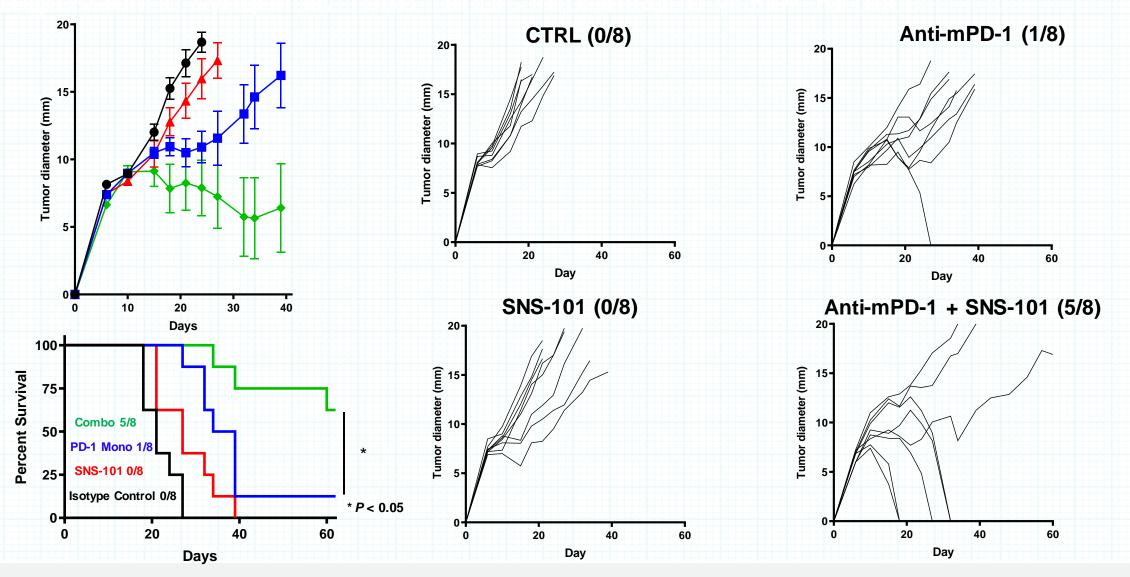
VISTA binding to the PSGL-1, inhibitory T cell receptor, is pH dependent



sensei

BIO

SNS-101 Re-sensitized Anti-PD-1 Insensitive Sarcomas Tumors in 1956 Model in Human VISTA Knock-in Mice





VISTA Biology – Summary Why VISTA Has Been Difficult to Drug Historically

- VISTA is expressed at high levels on monocytes and neutrophils
 - Binding to cells in the blood leads to sub-optimal PK due to target-mediated clearance and ontarget/off-tumor toxicity (CRS)
- Engagement of FcgR may be a prerequisite for optimal activity of anti-VISTA antibodies
 - Fc silent antibodies are not effective at T cell proliferation ex vivo or anti-tumor activity in vivo despite picomolar binding affinity to VISTA
 - CRS likely due to binding to blood monocytes and neutrophils, resulting in FcgR-mediated activation
- BUT the VISTA checkpoint itself is only "ON", when VISTA is protonated under low pH conditions and capable of binding PSGL-1
- Opportunity to develop an antibody with high selectivity for the active/protonated form of VISTA versus the inactive form of VISTA in the blood.
 - Johnston, et al (BMS), demonstrated feasibility of this approach
 - Sensei has developed a conditionally-active, pH-dependent antibody (SNS-101)



Edward van der Horst, Ph.D. SNS-101 Overview & Preclinical Data



SNS-101: Selectively Targeting VISTA with a pH-sensitive Antibody

Key features

- Selectivity for Active VISTA^{pH6} over VISTA^{pH7.4}
- Designed to block VISTA's interaction with PSGL-1 and all other T-cell receptors at pH 6.0
- IgG1 format
- Active Fc

Development milestones

- Preclinical PK, safety and efficacy data presented at conferences throughout 2022
- IND submission planned for 1H23

	рН 6.0	pH 7.4
Monovalent Affinity (K _D) [nM]	0.218	132 (~No binding)



SNS-101 Is a Fully Differentiated Anti-VISTA Antibody

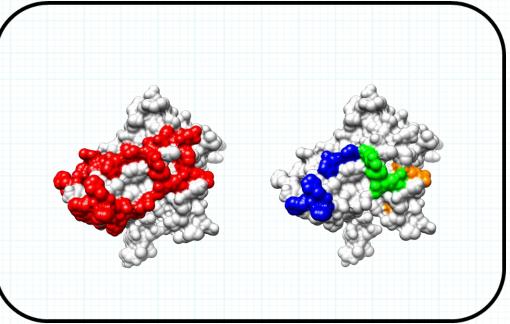
	SNS-101 Sensei	CI-8993; JNJ-61610588 (J&J/Curis)	K01401-020; W0180 (Pierre Fabre)	HMBD-002 (Hummingbird)	KVA12.1 (Kineta)	VISTA.18 (BMS)	(PMC-309) Pharm Abcine
Inhibit PSGL-1 Binding	\bigcirc	\bigcirc	\bigotimes	\bigotimes	\bigotimes	\bigotimes	\bigotimes
pH Sensitive Binding	\bigotimes	\bigotimes	\bigotimes	\bigotimes	\bigotimes	\bigotimes	\bigotimes
Fc Active	(IgG1)	(IgG1)	N/A	\bigotimes	(IgG1)	(IgG4)	(lgG1)
Stage	Preclinical	Phase 1	Phase 1	Phase 1	Preclinical	Preclinical	Preclinical



Johnston et al., Nature 2019; Kineta website; Snyder et al, AACR Annual Meeting 2016; Pierre Fabre website; Hummingbird website; Thakkar et al, J of Immunother Cancer, 2022; PharmAbcine website

SNS-101 Blocks PSGL-1:VISTA Protein Interface

SNS-101



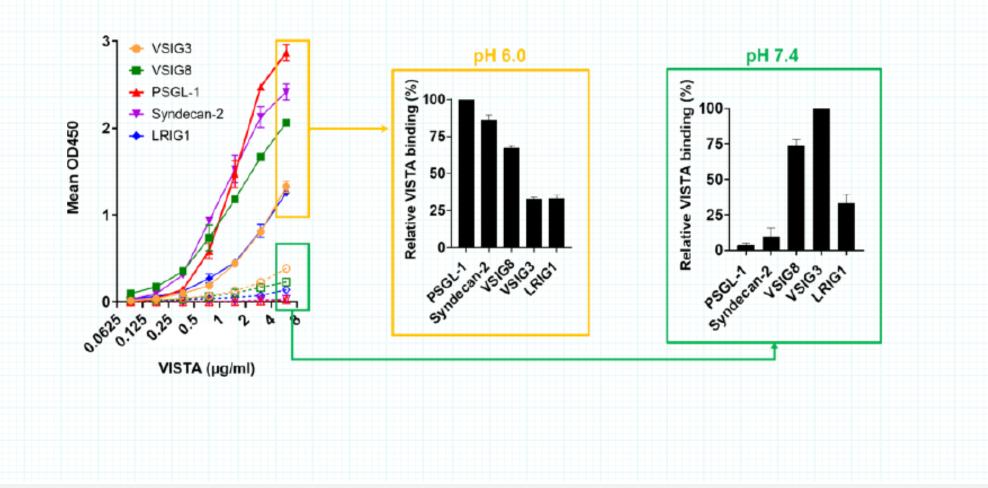
PSGL-1, VSIG-3, and LRIG-1

 VISTA's extracellular domain is uniquely rich in histidines¹

 Histidines are protonated at low pH enabling VISTA to distinguish the active (acidic pH) and inactive (neutral pH) PSGL-1 binding interface

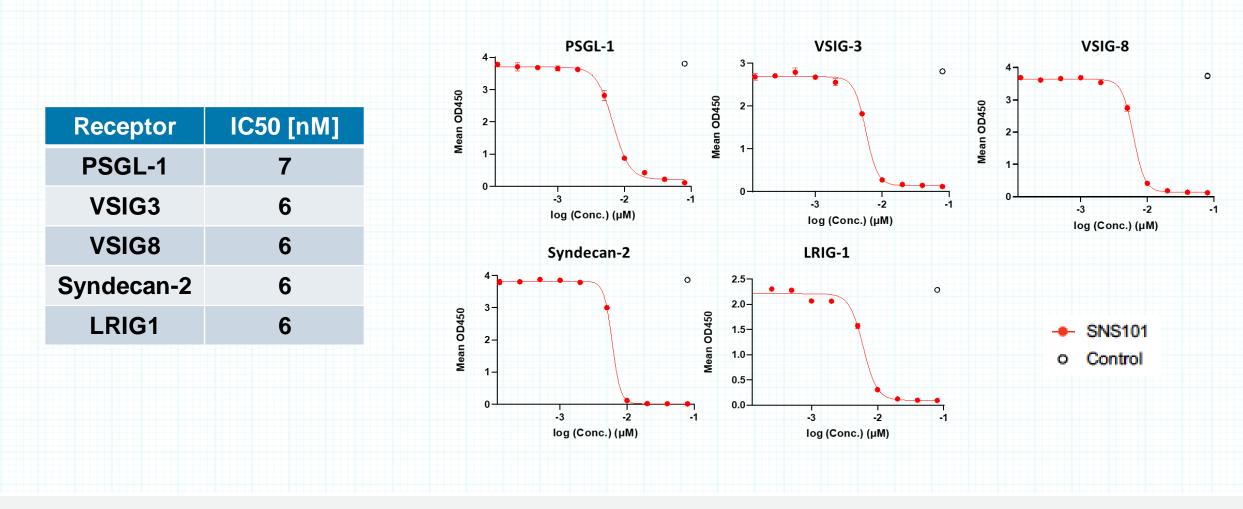


The VISTA:PSGL-1 Interaction is Selective for low pH

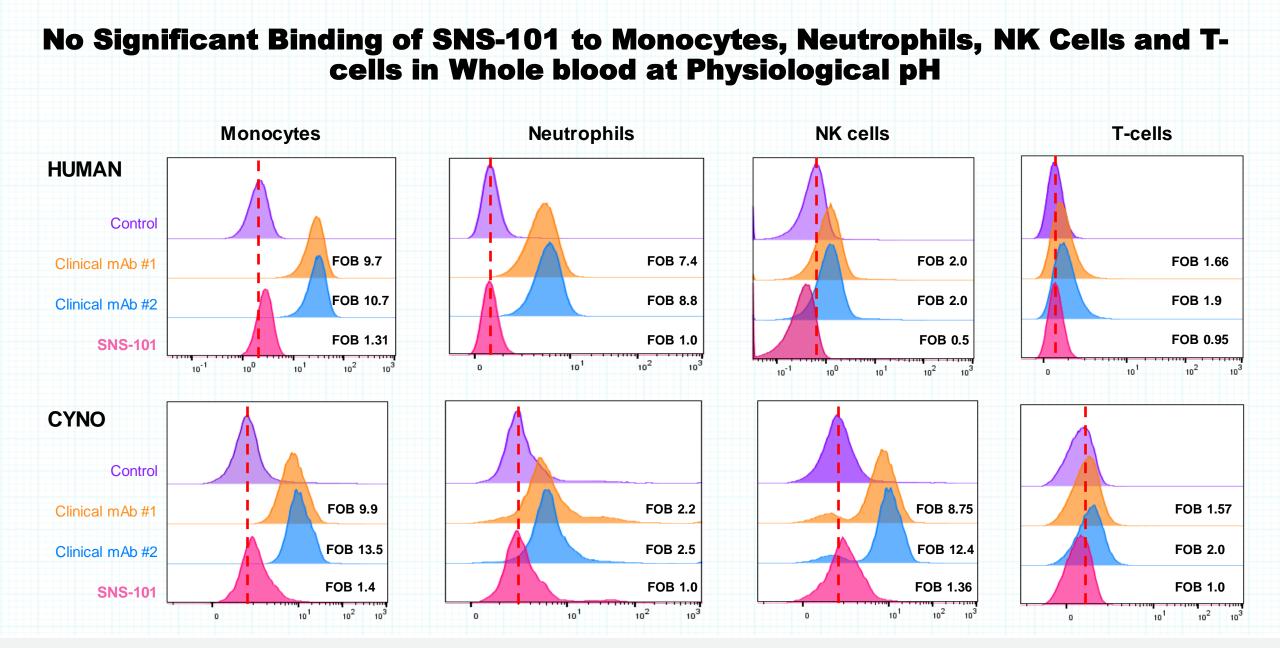




SNS-101 Strongly Inhibits the VISTA:PSGL-1 Interaction And All Other Potential Binding Partners at pH 6.0 in *In Vitro* Assay

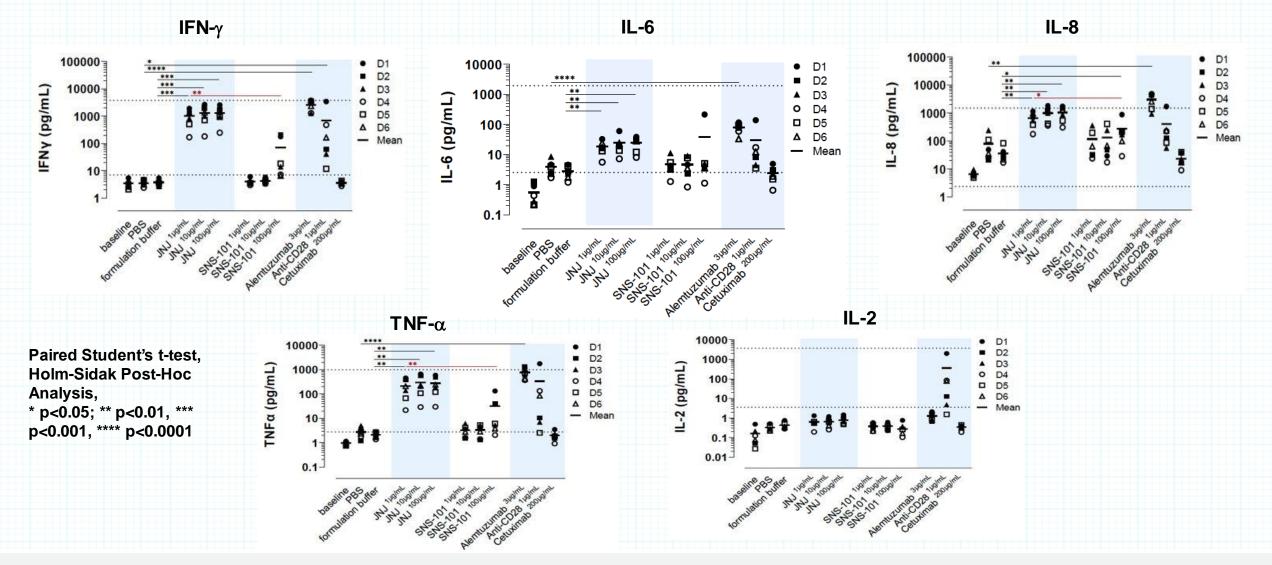






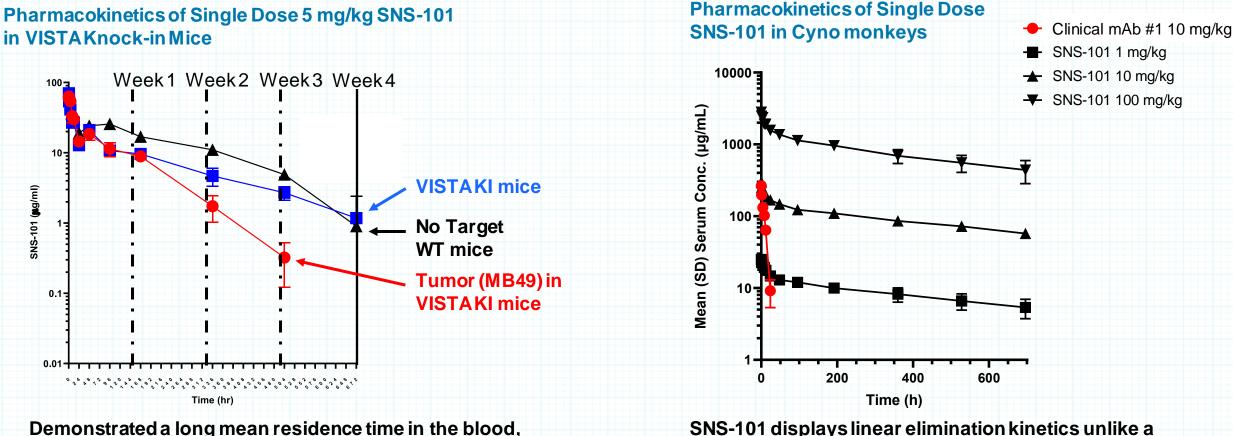


SNS-101 Induced Substantially Lower Cytokine Release in Whole-blood Assay at Neutral pH Compared to pH-independent VISTA Antibody





SNS-101 Has Displayed a Favorable Single-dose PK Profile in Preclinical Studies - *No Significant TMDD in Human VISTA KI Mice or Cyno Monkeys*

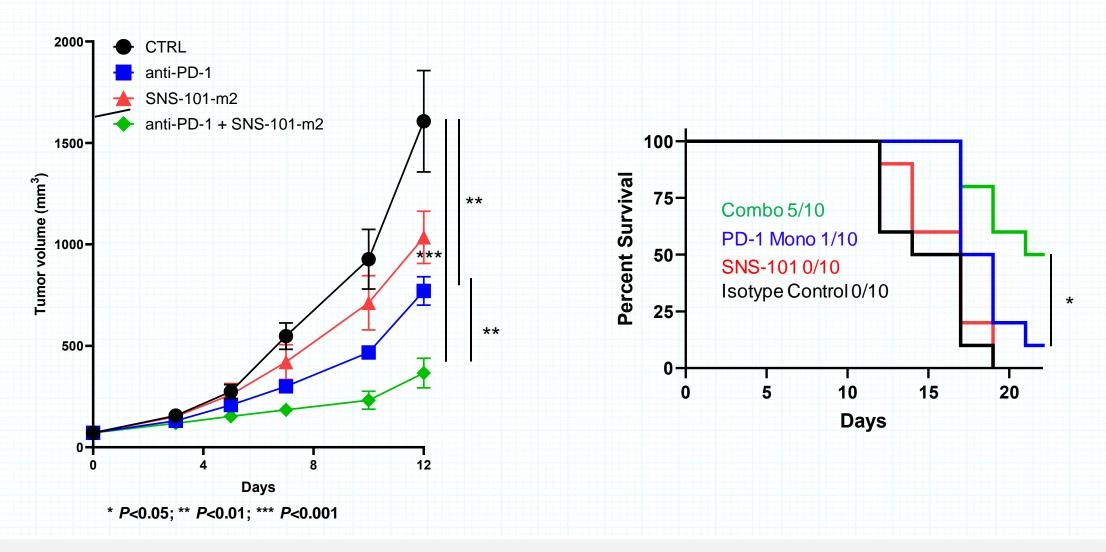


Demonstrated a long mean residence time in the blood, indicating a lack of significant target-mediated drug disposition (TMDD) and clearance in non-malignant tissues

ense

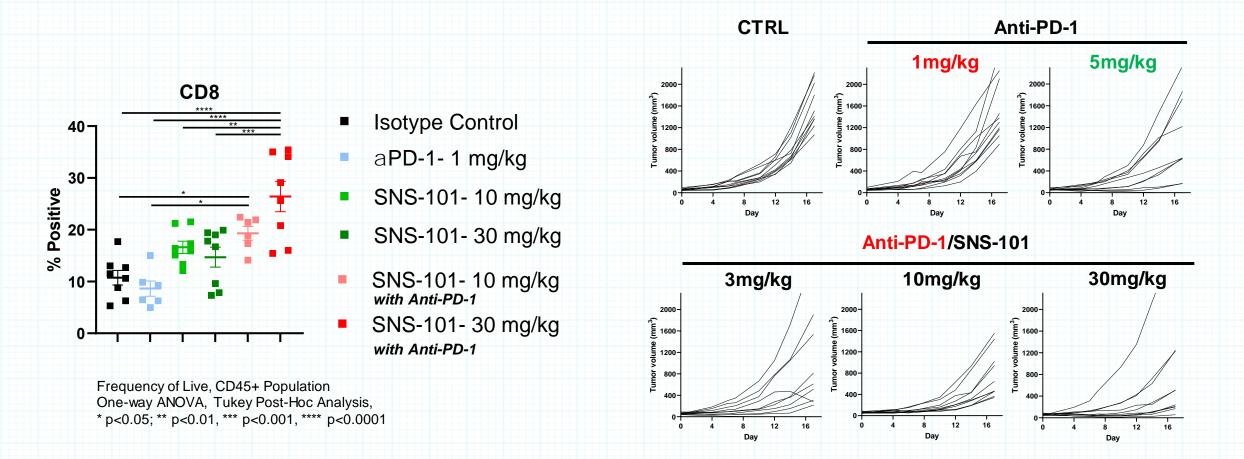
SNS-101 displays linear elimination kinetics unlike a pH-independent anti-VISTA mAb, which demonstrates TMDD and rapid clearance

SNS-101 Demonstrated Strong Combinatorial Activity with Anti-PD-1 in MC38 Model in Human VISTA Knock-in Mice





SNS-101 Demonstrated Increased CD8 T-cells in Combination With Anti-PD-1 in MC38 Tumors *In Vivo*





Robert Pierce, M.D. & Ron Weitzman, M.D. SNS-101 Translational Medicine & Clinical Development Plan

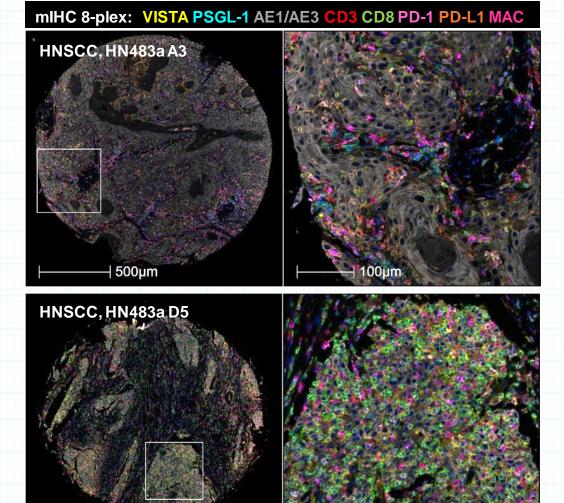


Multiplex IHC Enables Spatial Mapping of VISTA in the TME

Quantification of VISTA and PSGL-1 Expression

 Spatially resolve VISTA⁺ and PSGL-1⁺ within the broader immune-relevant context of the TME

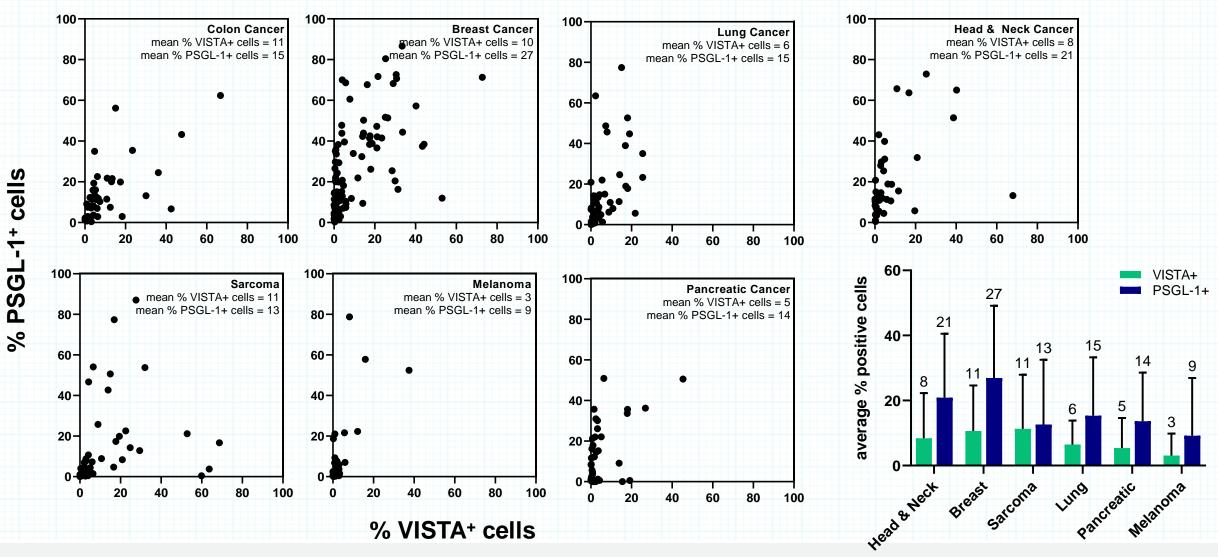
- Spatial resolution of VISTA, PSGL-1 and other candidate immune checkpoints within the TME may reflect critical cell-cell interactions and inhibitory nodes
- VISTA/PSGL-1 proximity may reflect engagement of VISTA checkpoint and high probability of response to SNS-101



500um

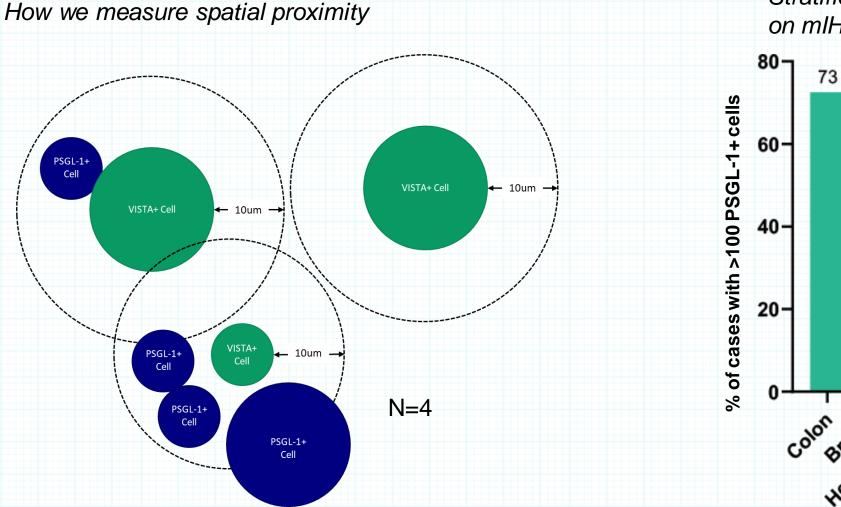


Quantitative Multiplex Immunohistochemistry Enables Stratification of Tumors with High Expression of VISTA and/or PSGL-1

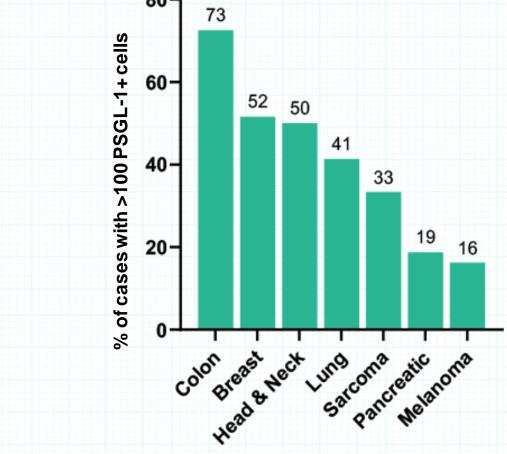




High VISTA/PSGL-1 Proximity Scores in CRC, Breast, HNSCC and Lung



Stratification of tumor types based on mIHC analysis of TMAs





SNS-101: Approach for Phase 2 Cohort Identification to Help Identify and Prioritize High Probability-of-Success Expansion Cohorts

Multiplex immunohistochemistry on patient tumor TMAs using 8-plex assay (VISTA, PSGL-1, PD-L1, PD-1, CD3, CD8, macrophage cocktail (CD68+CD163), tumor marker (pancytokeratin)

- Expression analysis of VISTA, PSGL-1, PD-1, PD-L1 in broader immune context
- Proximity analysis (VISTA/PSGL-1 and PD-L1/PD-1)
- Frequency of VISTA⁺ tumor cells across tumor types
- Distribution of VISTA, PSGL-1 and Proximity Indices across TME phenotypes (e.g. inflamed, immune-excluded, immune-ignored)
- Acquisition of well-curated, clinical samples with ICI treatment history and outcomes
- Is VISTA/VISTA-PSGL-1 proximity upregulated in PD-1 non-responders?

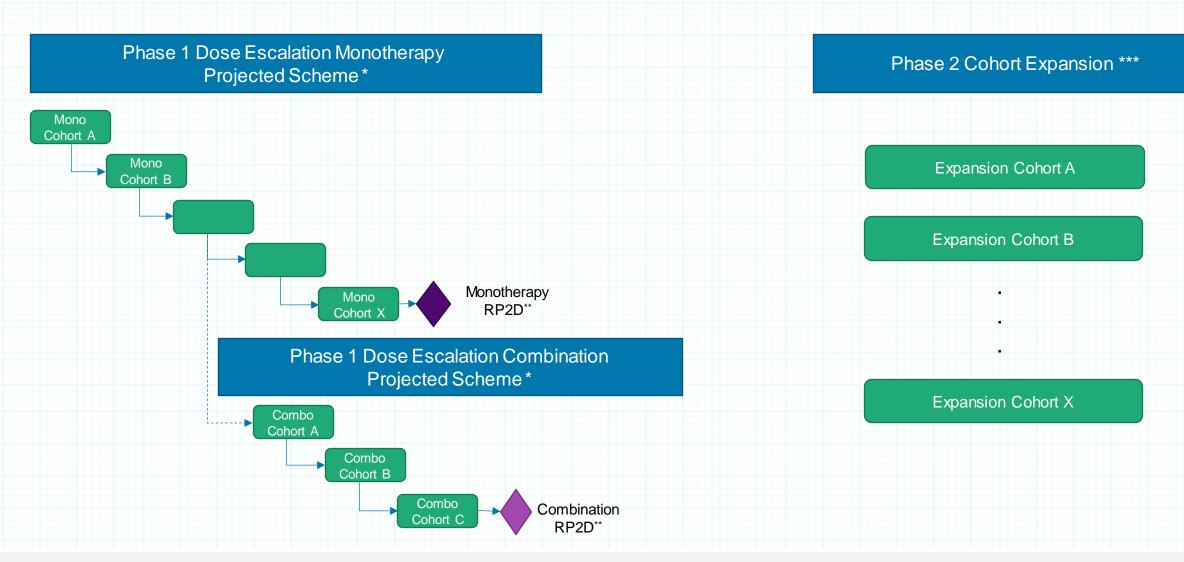
II. Multiplex IHC and scRNAseq on multiple preclinical tumor models

- Does VISTA/PSGL-1 proximity correlate with response to SNS-101?
- Other potential biomarkers of VISTA/PSGL-1 checkpoint engagement?



١.

Preliminary SNS-101 Phase 1/2 Study Schematic





- * Phase 1/2 study design is preliminary and subject to change, including based on feedback from the FDA following submission of IND.
- ** RP2D = Recommended Phase 2 Dose

*** Tumor types, indication and samples size to be determined based on findings from dose-escalation phase and emerging scientific data; cohorts may run concurrently.

SNS-101 Responder Hypothesis based on SNS-101 Preclinical and Translational Data

SNS-101 Preclinical/Translational Strengths	Potential Clinical Direction
 SNS-101 demonstrated strong combinatorial activity with anti-PD-1 in MC38 model in human VISTA-KI mice In-vivo efficacy profile in combination with PD-1 showed increase tumor-infiltrating CD8 T-cells 	 Boosting αPD-1 response in Inflamed/Immunogenic tumor types: <u>Current Plan</u>: checkpoint inhibitor naïve patients with inflamed solid tumors (e.g., HNSCC), who are responsive to αPD-1 treatment, would receive combination treatment of SNS-101 + αPD-1 and be assessed for a boost in anti-tumor activity
 Immunohistochemistry (IHC) staining VISTA/PSGL-1 proximity assay suggests tumors with high VISTA/PSGL-1 proximity signal Include CRC, Breast, NSCLC and HNSCC 	- Inducing a response in inflamed, but typically α PD-1 non-responsive tumors (e.g., MSS colon, HR+ breast) with combination of SNS-101 and α PD-1
 SNS-101 and PD-1 blockade led to enhanced tumor regression in syngeneic 1956 tumors implanted in VISTA-KI mice 	 Bona fide αPD-1 refractory patients (e.g., NSCLC): Does treatment with combination treatment of SNS-101 + αPD-1 overcome resistance?



Fireside Chat with Neil Canavan



Question & Answer Session



VISTA Science Symposium

November 21, 2022

Guest Speaker: Robert Schreiber, Ph.D.

Andrew M. and Jane M. Distinguished Professor of Pathology and Immunology; Professor, Molecular Microbiology; and Director of the Bursky Center for Human Immunology and Immunotherapy Programs at the Washington University School of Medicine. He is also co-leader of the Tumor Immunology Program of Washington University's Siteman Comprehensive Cancer Center, an Associate Director of the Scientific Advisory Board to the Cancer Research Institute and Co-editor-in-Chief of the journal Cancer Immunology Research. Schreiber obtained his PhD in Immunology/Biochemistry at the State University of New York in Buffalo, New York, and received his postdoctoral training at The Scripps Research Institute in La Jolla, California. Sensei IOAB Member.

Sensei Presenters:

John Celebi Chief Executive Officer

Dr. Robert Pierce Chief R&D Officer

Ron Weitzman

Dr. Edward van der Horst

SVP, TMAb Antibody Development

