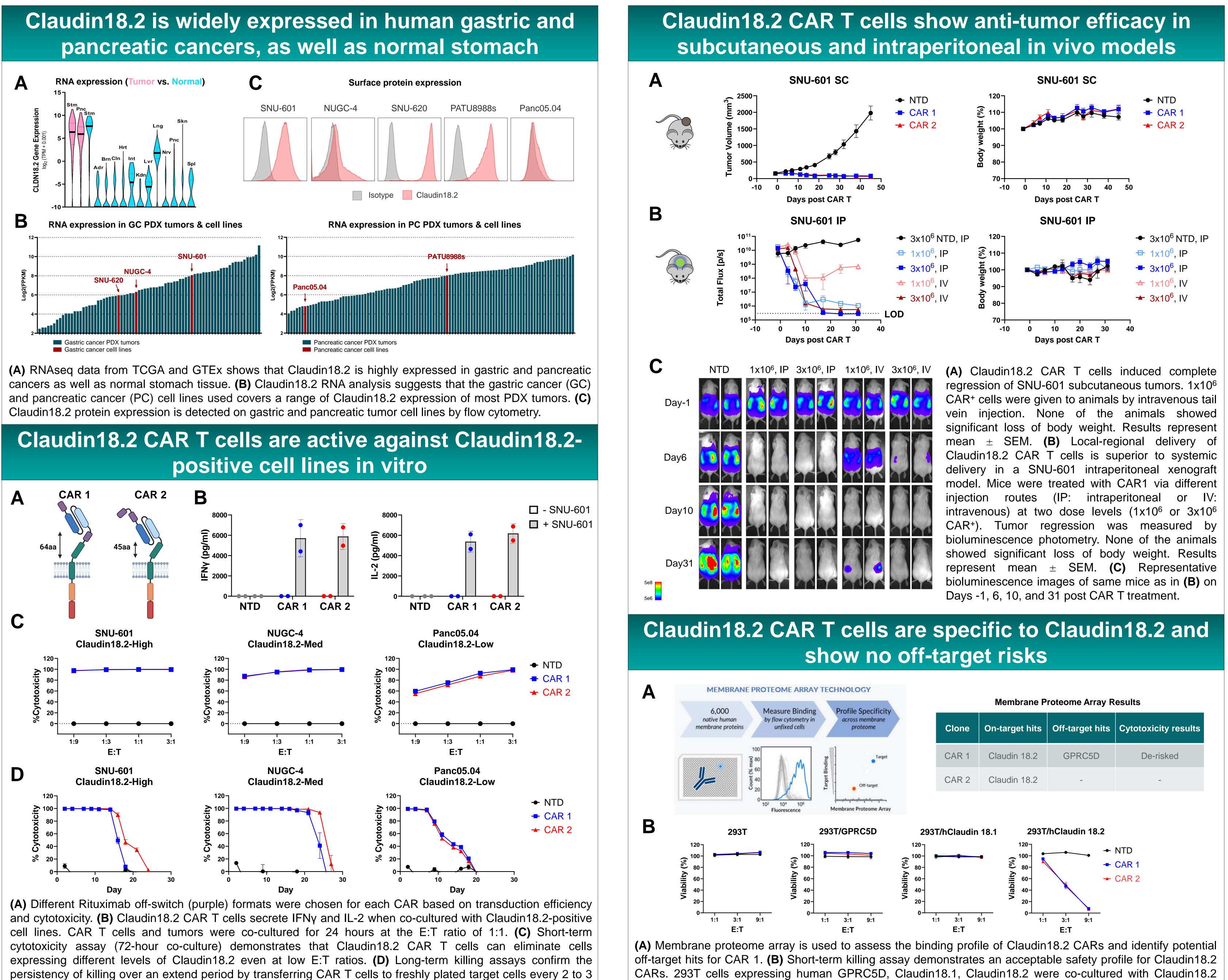
Preclinical development and characterization of allogeneic CAR T cells targeting Claudin18.2-positive tumors

Zhe (Joanne) Li¹, Barbra Sasu¹, Siler Panowski¹

Background: Claudin18.2 (CLDN18.2) has emerged as a promising therapeutic target, with high expression in many types of epithelial cancers including stomach, esophagus, pancreatic, and ovarian cancers. Multiple CLDN18.2-directed monoclonal antibodies and autologous CAR T cell therapies have entered clinical trials and demonstrated promising results. Here, we describe the preclinical benefit with a single treatment and overcome many of the challenges facing autologous CAR T cells. Methods: A panel of allogeneic fully-human scFv-based CAR T cells targeting both human and murine CLDN18.2 were generated. To enable off-the-shelf use, TALEN[®] gene editing was used to knock out (KO) the TRAC locus and CD52 KO was also used to permit CD52-directed lymphodepletion regimens. Binding specificity was studied utilizing cell lines and candidates were also evaluated to provide control over CAR T function.

Results: A subset of CARs showed highly specific binding to CLDN18.2 and evaluation of these CARs in multiple rituximab off-switch formats identified candidates with potent activity in both short-term and repeat stimulation in vitro cytotoxicity assays. Lead allogeneic CAR T cell candidates exhibited cytokine release upon target exposure and intraperitoneal gastric cancer models. Safety evaluation was also performed, including body weight measurement. Taken together, these data support the existence of a therapeutic window and the potential to target CLDN18.2 with allogeneic off-the-shelf CAR T cells.

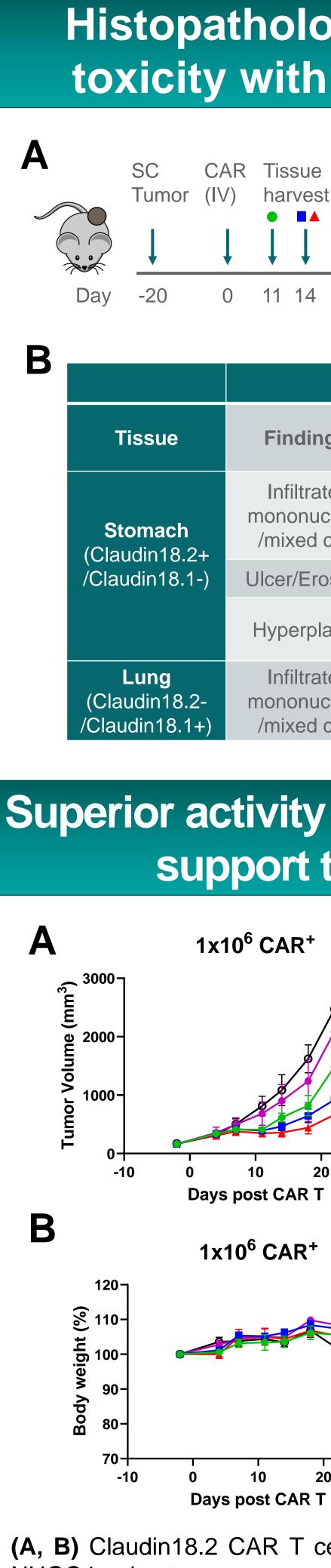


days. Results represent mean ± SEM, n=3 technical replicates. NTD, non-transduced T cells. Experiments were performed three times with CAR T cells from three different donors.

CARs. 293T cells expressing human GPRC5D, Claudin18.1, Claudin18.2 were co-cultured with Claudin18.2 CARs for 72 hours. Results represent mean \pm SEM, n=3 technical replicates. Experiments were performed three times with CAR T cells from two different donors.

¹Allogene Therapeutics, South San Francisco, CA

Membrane Proteome Array Results								
Clone	On-target hits	Off-target hits	Cytotoxicity results					
CAR 1	Claudin 18.2	GPRC5D	De-risked					
CAR 2	Claudin 18.2	-	-					



(A, B) Claudin18.2 CAR T cells induced a dose-dependent anti-tumor responses and body weight loss in a NUGC4 subcutaneous xenograft model. 1x10⁶ or 3x10⁶ CAR⁺ cells were given to animals by intravenous tail vein injection. (A) Differences in efficacy remain consistent at both dose levels. (B) The level of toxicity, measured by loss of body weight, is positively correlated with anti-tumor efficacy. None of the animals showed body weight loss at the 1x10⁶ dose. (C) Claudin18.2 CAR T cells exhibited expansion in the blood. Blood was collected from groups treated with $3x10^6$ CAR⁺ cells. Results represent mean \pm SEM.

- positive cancers.

TALEN® gene-editing is a technology pioneered and controlled by Cellectis. The anti-Claudin18.2 AlloCAR T program, which utilizes Cellectis technology, is licensed exclusively from Cellectis by Allogene and Allogene holds global development and commercial rights.



Abstract #283

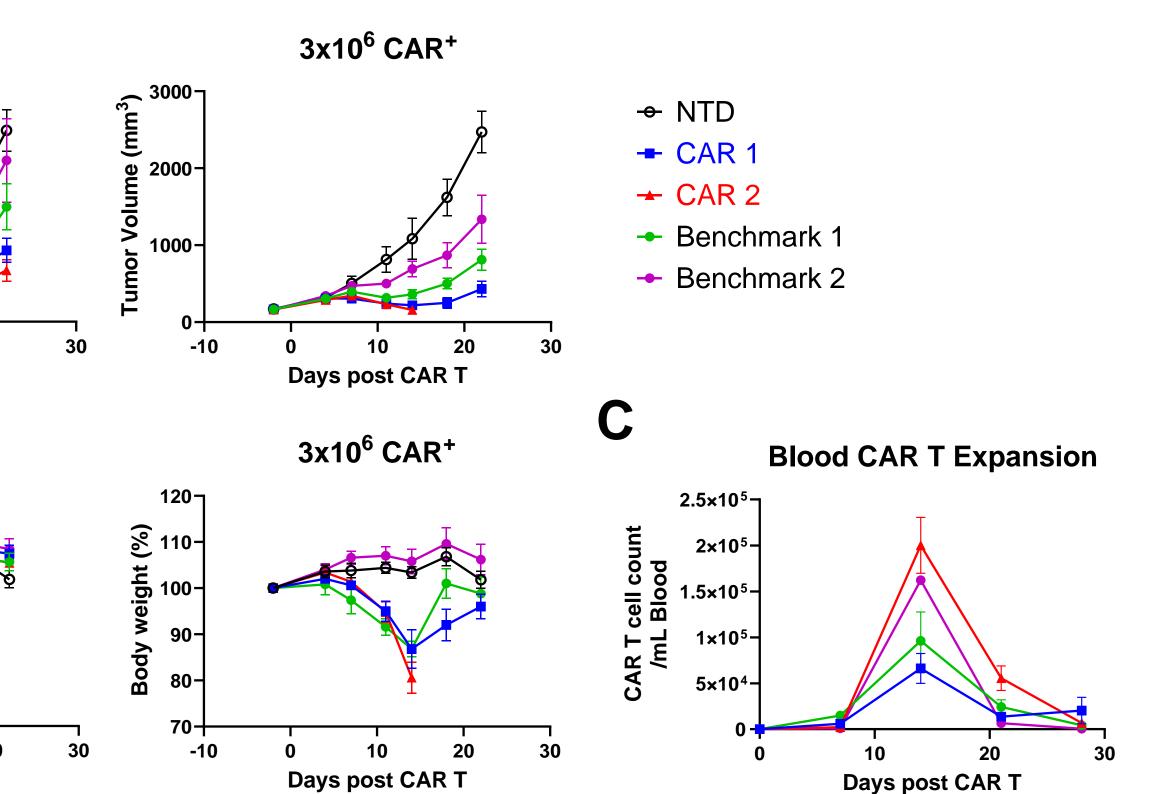
Histopathology analysis identify expected on-target toxicity with Claudin18.2 CAR T cells at high doses

Tissue harvest	NTDCAR 1
40	 CAR 2 Benchmark 1 Benchmark 2

(A) Schematic of in vivo safety evaluation using NUGC4 subcutaneous model. Each Claudin18.2 CAR T treatment group was harvested at different time point due to rapid body weight loss. (B) Histopathology analysis showed infiltration and tissue damage in the stomach, which matches the normal tissue expression of Claudin18.2.

	Treatment					
gs	NTD , 1x10 ⁷ , (n=3)	CAR 1 1x10 ⁷ , (n=3)	CAR 2 1x10 ⁷ , (n=3)	Benchmark 1 1x10 ⁷ , (n=3)	Benchmark 2 1x10 ⁷ , (n=3)	
te, clear cell	0	Moderate (n=2)	Moderate (n=2) Marked (n=1)	Moderate (n=3)	Moderate (n=3)	
osion	0	Mild (n=1)	Moderate (n=1)	0	Marked (n=3)	
asia	0	Mild (n=2)	Minimal (n=1)	Mild (n=1)	Mild (n=1) Moderate (n=2)	
te, clear cell	0	Moderate (n=2) Marked (n=1)	Moderate (n=1) Marked (n=2)	Mild (n=1) Moderate (n=2)	Mild (n=3)	

Superior activity of Claudin18.2 CAR T cells over benchmarks support the existence of a therapeutic window



Conclusions

 Claudin18.2 RNA is expressed in gastric and pancreatic tumors and normal tissue expression is limited to the stomach.

• The efficacy and safety data suggest that allogeneic Claudin18.2 CARs may be an effective and clinically valuable treatment for patients with Claudin18.2-