Gao Delivery Team Summary

<u>Goals</u>

Detect and quantify genome editing in lung airway epithelia following AAV delivery

Examine non-target tissues for evidence of editing

<u>Strategy</u>

Activation of fluorescent tdTomato following CRISPR/Cas9 editing

Delivery reagents

Study Arm	Vector	Cargo	Dosage	Solvent	Volume	Delivery Method
1 - Experimental	ssAAV5	B guide + SaCas9	1.7E11 vg	Saline	40 µl	Intratracheal intubation
2 - Experimental	ssAAV5 x 2	 SpCas9 A & B guides + GFP 	1) 1.7E11 vg 2) 1.7E11 vg	Saline	40 µl	Intratracheal intubation
3 - Positive control	ssAAV5	Cre recombinase	1.7E11 vg	Saline	40 µl	Intratracheal intubation
4 - Negative control	-	saline	-	Saline	40 µl	Intratracheal intubation

Test mice

Study Arm	Mouse genotype	Age	Incubation time	Number
1 - Experimental	SaSp Ai9/SaSp Ai9	8 weeks	4 weeks	7M 7F
2 - Experimental	SaSp Ai9/SaSp Ai9 and Ai9/Ai9	8 weeks	4 weeks	7M 9F
3 - Positive control	SaSp Ai9/SaSp Ai9	8 weeks	4 weeks	2M 2F
4 - Negative control	SaSp Ai9/SaSp Ai9	8 weeks	4 weeks	3M 3F

Reporter mouse model: conventional Ai9



= SpCas9 guide target
= SaCas9 guide target

Reporter mouse model: single-guide SaSp Ai9



Animal Health data



H&E stains



Male

Study Arm 2 Experimental AAV5-SpCas9 & AAV5-Guides A&B-GFP 2857

Female

Study Arm 2 Experimental AAV5-SpCas9 & AAV5-Guides A&B-GFP 2875

Animal health summary

No anomalies were observed with respect to body weight, liver weight or spleen weight in any mouse, regardless of the treatment

Microscopic examination of the lung (target tissue), liver and spleen revealed normal histologies, with no evidence of inflammation or toxicity

Following dissection, organs and removal of small samples for molecular analysis, organs were fixed overnight in 4% paraformaldehyde, then saturated overnight in 30% sucrose. Samples were frozen in OCT and stored at -80 C

Lung and other organ samples were sectioned at 14 µm and imaged on an AxioScan.X1 slide scanner with a 20X objective

Each section was imaged for tdTomato, indicating a successful editing event, and DAPI to highlight nuclei and identify airways

Some sections were also imaged for GFP, to mark successful AAV delivery (not shown)

<u>Males</u>

Target tissue Lung airway epithelia

tdTomato

DAPI

Merged

Study Arm 1 Experimental AAV5-guideB-SaCas9

100 µm

100 µm

100 µm

2857







<u>100 µm</u>

Study Arm 3 Positive Cont. AAV5-Cre

<u>100 µm</u> 2845

Study Arm 4 Negative Cont. Saline



2868

<u>Females</u>

Study Arm 1 Experimental AAV5-guideB-SaCas9

<u>100 µm</u>

Target tissue Lung airway epithelia

tdTomato

DAPI

Merged

3304

Study Arm 2 Experimental AAV5-SpCas9 + AAV5-guidesA&B-GFP



2847

Study Arm 3 Positive Cont. AAV5-Cre

Study Arm 4 Negative Cont. Saline <u>100 µm</u>

<u>100 µm</u>

100 µm



<u>100 µm</u>

Study Arm 1 - Experimental: AAV5 - guideB-SaCas9

Few to no red cells observed localized to airway epithelia, indicating no significant editing of target. Five out of 14 (1 M, 4 F) showed sporadic red cells (>5 per section) not localized to airways.

<u>Study Arm 2 - Experimental: AAV5-SpCas9 + AAV5-guidesA&B-GFP</u>

In 7 animals (1 M, 6 F), significant number of red cells in airway epithelia, indicating successful editing. In remaining animals, few to no red cells observed.

Study Arm 3 - Positive control: AAV5-Cre

Significant red cell accumulation observed, as expected.

Study Arm 4 - Negative control: saline

No red cells detected, as expected.

Quantification of editing in target tissue - lung airway epithelia



Therefore, 26.3% editing efficiency, in a 34960 μ m² airway

Quantification results

Mouse	Small airways (n=3) %	Large airways (n=3) %
2857 M	21.5 ± 4.6	17.6 ± 6.8
2867 F	17.6 ± 2.5	14.5 ± 1.0
2874 F*	22.1 ± 3.7	19.3 ± 5.7
2875 F*	25.3 ± 3.7	20.6 ± 5.8
3305 F	21.2 ± 3.1	11.6 ± 3.2
3307 F	17.3 ± 3.7	13.9 ± 1.3
3309 F	18.2 ± 1.1	14.6 ± 2.5

* Conventional Ai9, not SaSp Ai9

Molecular analysis: PCR to detect AAV genomes

- PCR was performed to detect the presence of vector genomes in genomic DNA isolated from lung samples
- Primers were specific for SaCas9, SpCas9 or GFP transgenes
- Presence of AAV genomes serves as confirmation of successful intratracheal infusion

<u>Result:</u> presence of AAV genomes correlated very well with detection of tdTomato+ cells

		Study Arı Negative Saline	m 4 Cont.		Study Positiv AAV5-	Arm 3 ve Cont. Cre	St E> A/	udy Arr perime AV5-gui	n 1 :ntal deB-S	aCas9				Study Exper AAV5 AAV5	Arm 2 iment -SpCa: -guide	2 :al s9 + esA&B-C	iFP						blank		
primers detect: SaCass	- Ə						*	* *			- +	+	+	+ +	+	+ -				-				+ airway edi detected by red cells	iting
SpCas	9										-	-	-	-		-								* Editing detected by re cells, but not in airways	d n
GFI											-	-													
Contro	mouse 82	2846 2850	2863	2864	2845 2847	2852 2864	2868	2848 2866	2853	2854	2867 2867	3305	3307	3309 2857	2874	2875 2849	2855	2856 2869	3301	3302	3303	3308			

Analysis of genome editing in non-target tissues

- From every animal in all four study arms, a full panel of non-target organs was dissected and prepared for imaging
 - Brain, Eye, Heart, Trachea, Liver, Kidney, Pancreas, Spleen, Stomach, Small Intestine (Duodenum, Jejunum, Ileum), Large Intestine, Muscle (Gastrocnemius, Soleus, TA, EDL and Diaphragm), White Adipose (Subcutaneous and Perigonadal), Brown Adipose, Testes or Ovary, Epididymis or Uterus
- For those animals showing significant editing in the target tissue, the non-target organs were sectioned and imaged as above
 - Three non-consecutive sections from each organ were imaged
- Images below are from one male and one female mouse, each of which showed significant editing in the target tissue

<u>Results:</u> Very rare red cells were observed in some non-target tissues.

Study Arm 2 Experimental AAV5-SpCas9





Study Arm 2 Experimental AAV5-SpCas9





Study Arm 2 Experimental AAV5-SpCas9

3305 F

Trachea



Study Arm 2 Experimental AAV5-SpCas9



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Kidney



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Pancreas



Study Arm 2 Experimental AAV5-SpCas9



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Stomach



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Small Intestine (Duodenum)



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Small Intestine (Jejunum)



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Small Intestine (Ileum)



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Large Intestine



Study Arm 2
Experimental
AAV5-SpCas9

3305 F

Muscle (Gastrocnemius)



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Muscle (Soleus)



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Muscle (TA)



Study Arm 2 Experimental AAV5-SpCas9



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Muscle (Diaphragm)



Study Arm 2 Experimental AAV5-SpCas9

3305 F

Brown Adipose Tissue



Study Arm 2 Experimental AAV5-SpCas9

3305 F

White Adipose (Perigonadal)



Study Arm 2 Experimental AAV5-SpCas9

3305 F

White Adipose (Subcutaneous)



Study Arm 2 Experimental AAV5-SpCas9





Study Arm 2 Experimental AAV5-SpCas9



Study Arm 2 Experimental AAV5-SpCas9

2857 M





Study Arm 2 Experimental AAV5-SpCas9

2857 M

Epididymis



Zoomed images of red cells in non-target tissues

Gastroc



3305 F

Trachea



Liver



Testes



<u>Conclusion:</u> Extremely rare red cells are detectable in non-target tissues