

Supplementary Materials: The CRISPR/Cas9 Minipig—A Transgenic Minipig to Produce Specific Mutations in Designated Tissues

Martin Fogtmann Berthelsen, Maria Riedel, Huiqiang Cai, Søren H. Skaarup, Aage K.O. Alstrup, Frederik Dagnæs-Hansen, Yonglun Luo, Uffe B. Jensen, Henrik Hager, Ying Liu, Henrik Callesen, Mikkel H. Vendelbo, Jannik E. Jakobsen and Martin Kristian Thomsen

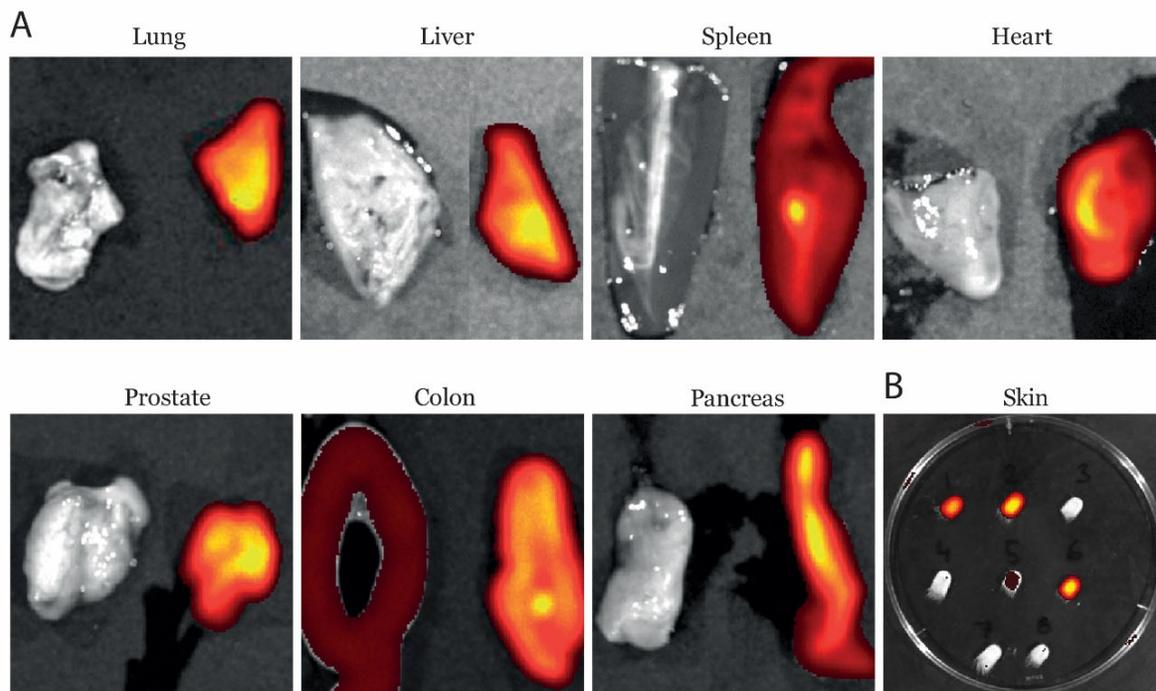


Figure S1. Transgenic expression. (A) IVIS scanning of tissue biopsies from a euthanized Cas9 pig verified transgene RFP expression in major organs. As a negative control tissue biopsies from a Danish Landrace pig were used. (B) IVIS scan of ear biopsies from a F1 litter.

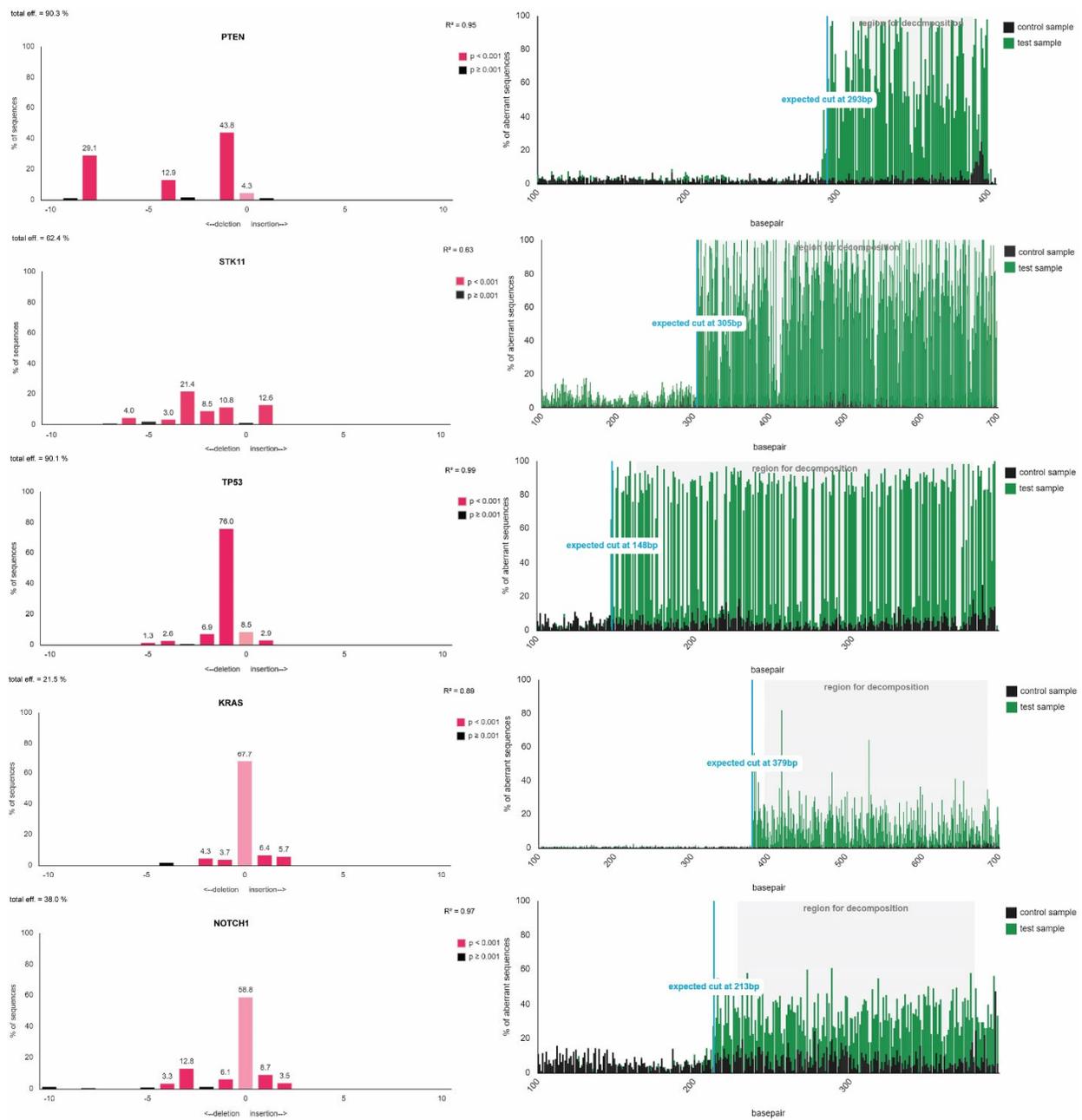


Figure S2. Validation of sgRNA in porcine fibroblast. Porcine fibroblasts transfected with the specified sgRNAs to determine guide efficiency. Mutations efficiency was assessed by TIDE analysis: Tracking of Indels by Decomposition software.

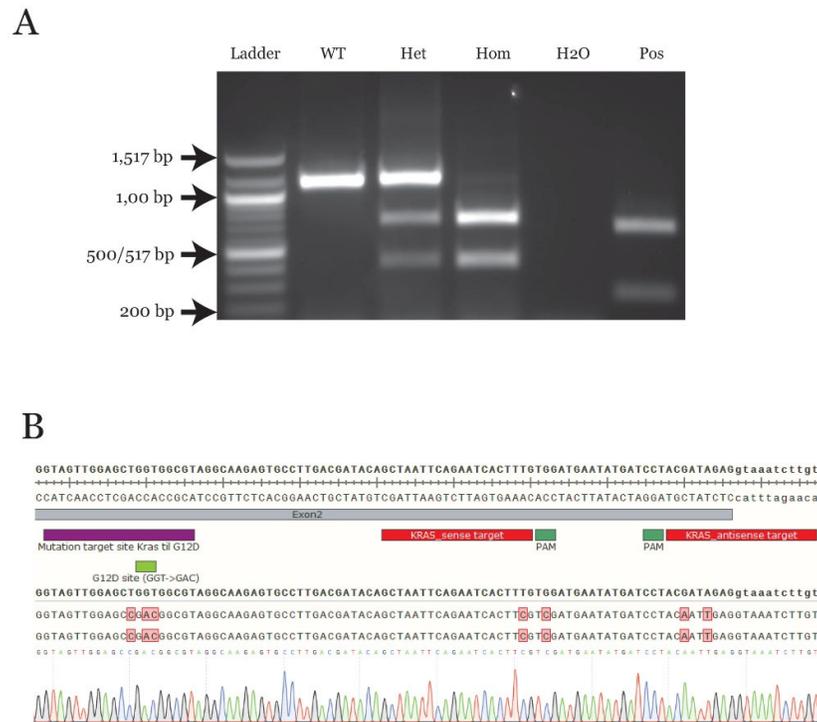


Figure S3. Introduction of KRAS^{G12D} point mutation by HDR. **(A)** The introduction of the KRAS^{G12D} point mutation by CRISPR HDR was assessed in single clones. Fibroblast from Cas9 transgene pic was co-transfected with sgRNA for KRAS and a KRAS^{G12D} repair template. Introduction of KRAS^{G12D} generated a restriction site and clones were analysed by PCR followed by an enzymatic digestion for the present of the new restriction site; KRAS^{G12D} mutated clones revealed two bands and non-mutated clones had a single band. Heterozygotes revealed three bands, one corresponding to the WT band and two to the cleaved mutated allele. Positive control was the KRAS^{G12D} template. **(B)** Sanger sequencing verified that the DNA sequence of the KRAS gene was identical to the sequence of the KRAS^{G12D} HDR template. Western Blot images can be found in Figure S7.

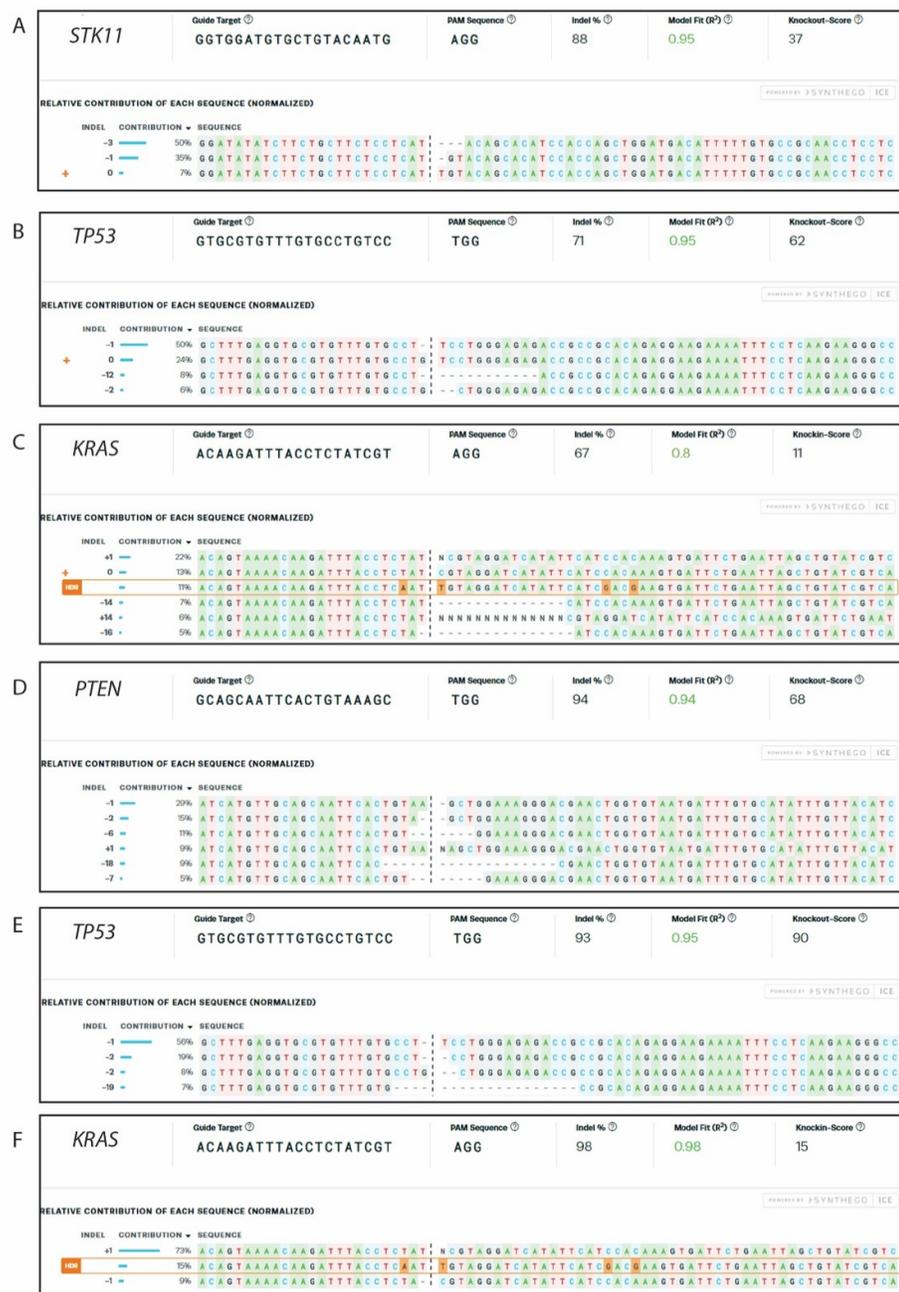


Figure S4. ICE analysis for AAV induces mutations in vitro. (A) *STK11* mutations in SKT_AAV2H22 transduced Cas9 fibroblasts. (B) *TP53* mutations in SKT_AAV2H22 transduced Cas9 fibroblasts. (C) *KRAS* mutations and *KRAS*^{G12D} repair in SKT_AAV2H22 transduced Cas9 fibroblasts. (D) *PTEN* mutations in PTK_AAV9 transduced Cas9 fibroblasts. (E) *TP53* mutations in PTK_AAV9 transduced Cas9 fibroblasts. (F) *KRAS* mutations and *KRAS*^{G12D} repair in PTK_AAV9 transduced Cas9 fibroblasts.

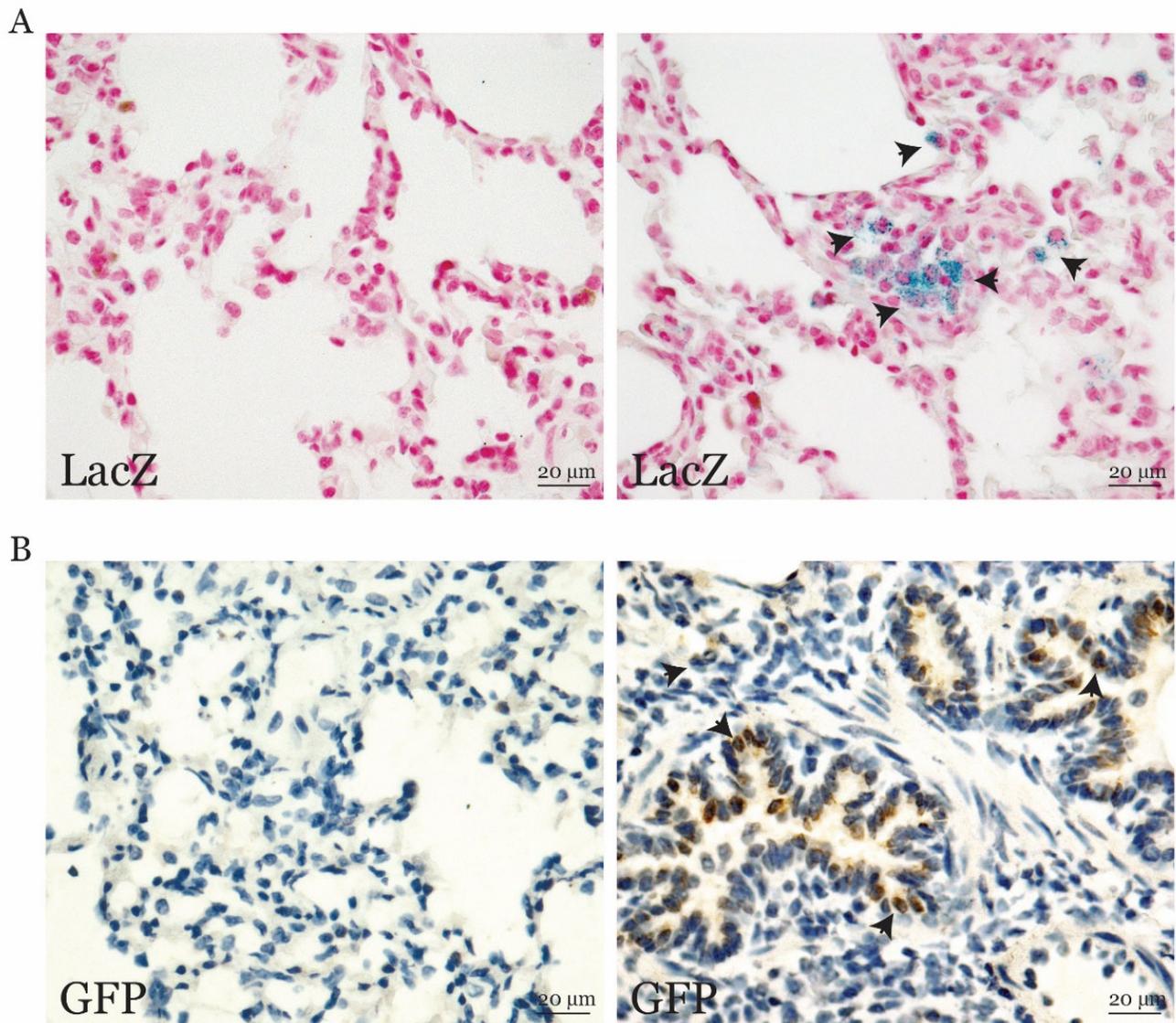


Figure S5. AAV targeting of pig lung epithelium. (A) β -galactosidase for LacZ in LacZ_AAV2H22 treated (right) and non-treated (left) porcine lungs. (B) Immunohistochemically staining for GFP in GFP_AAV9 treated (right) and non-treated (left) porcine lungs. Arrowheads mark positive cells.

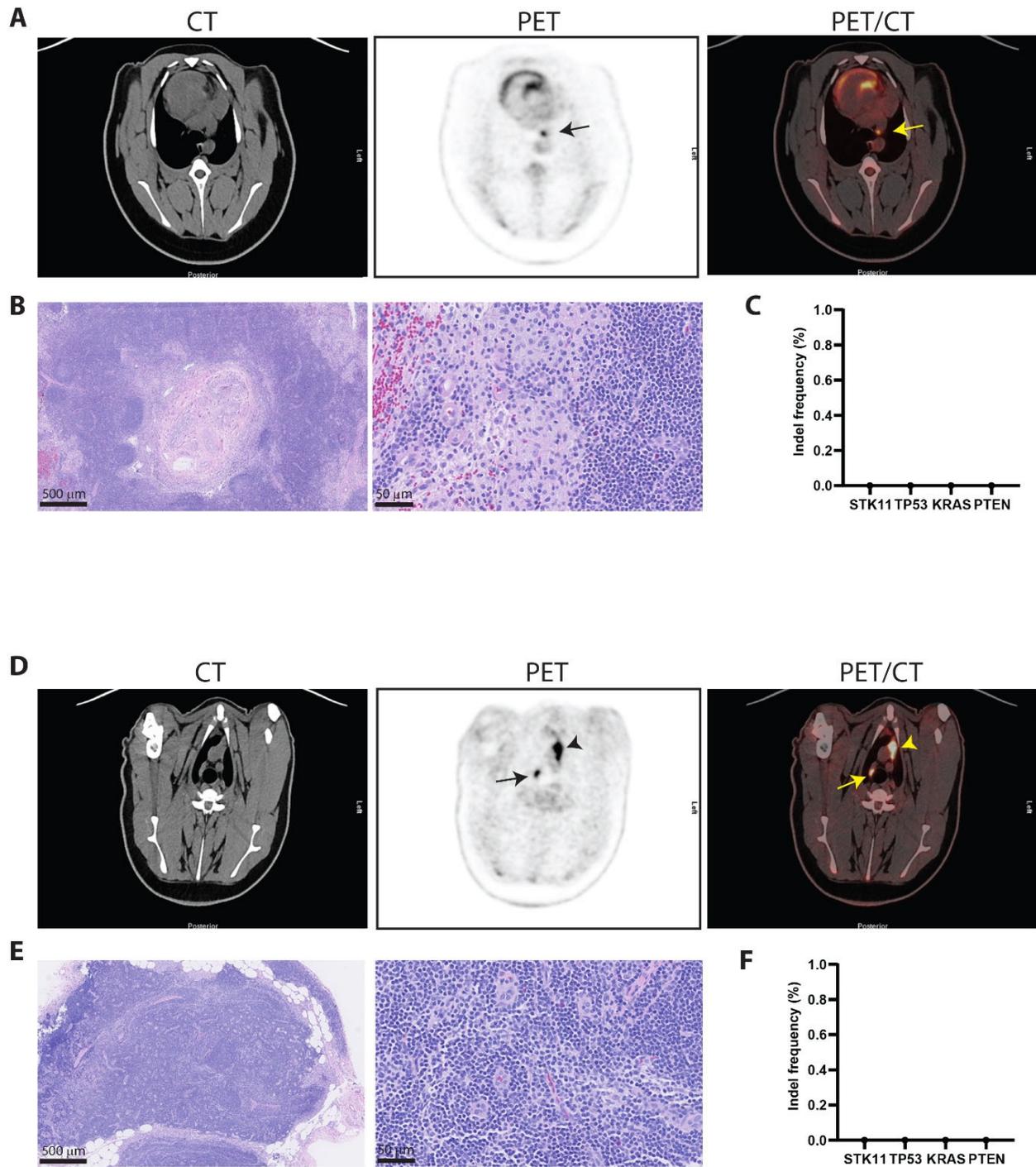
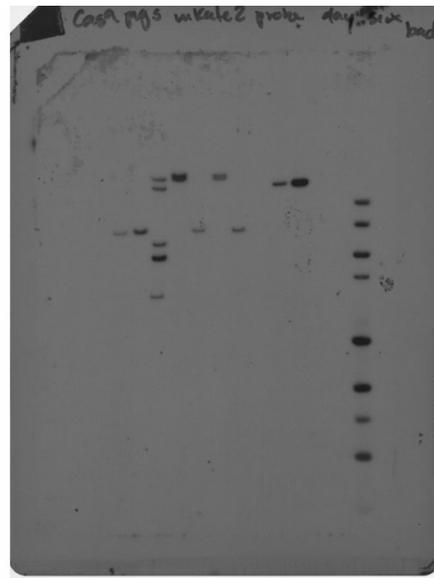
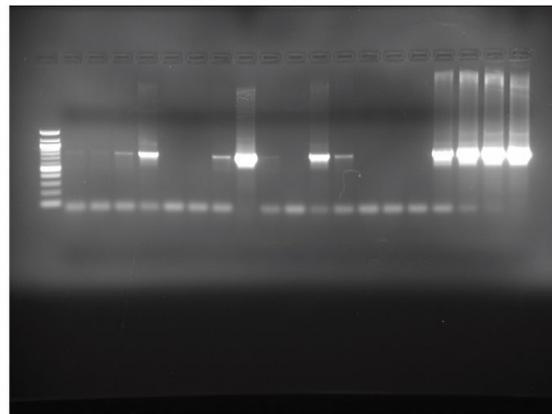


Figure S6. Activated lymph nodes. Activated lymph nodes were detected by increased ^{18}F -FDG signal by PET/CT scans. (A,D) PET/CT scans from two Cas9 mini pigs 18 months post AAV transduction. Arrows mark the activated lymph nodes and arrowhead mark thymus. Representative images are shown. (B,E) Histological sections from activated lymph nodes stained with H&E. (C,F) Indel analysis for CRISPR induced mutations in the four target genes. No Indels were detected by the Sanger sequencing.

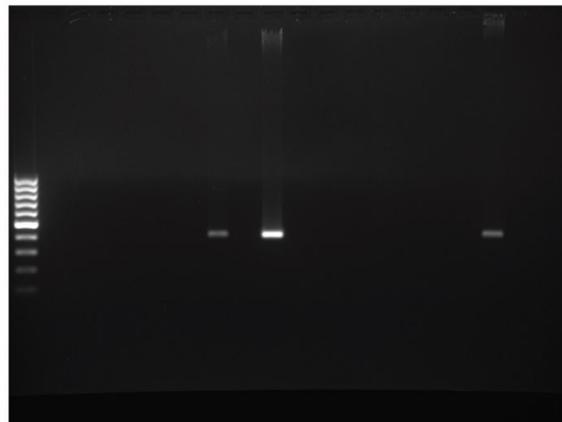
SB - Figure 1



Viral DNA - Figure 4



Recombination
Figure 4



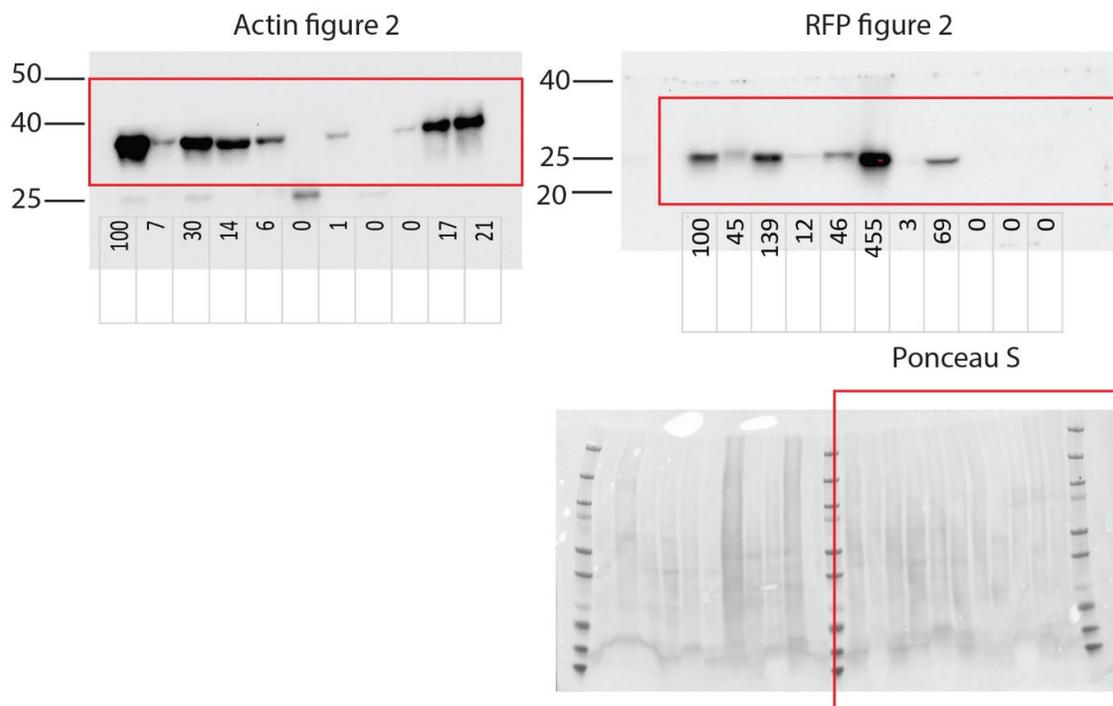
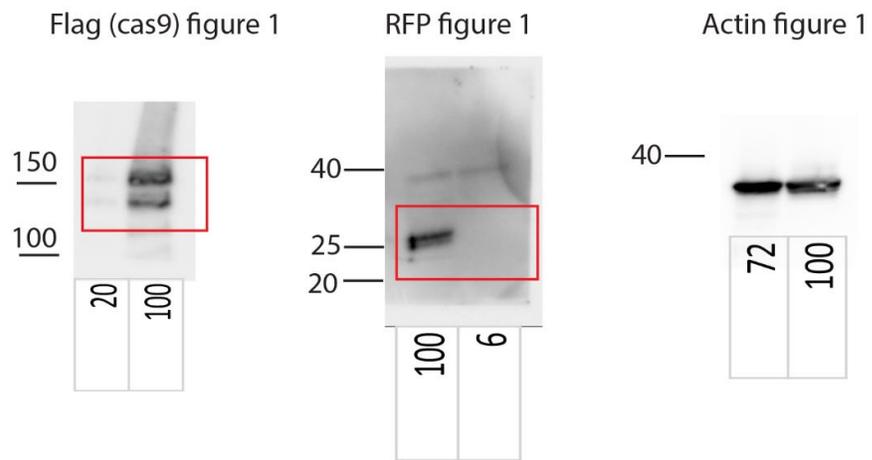
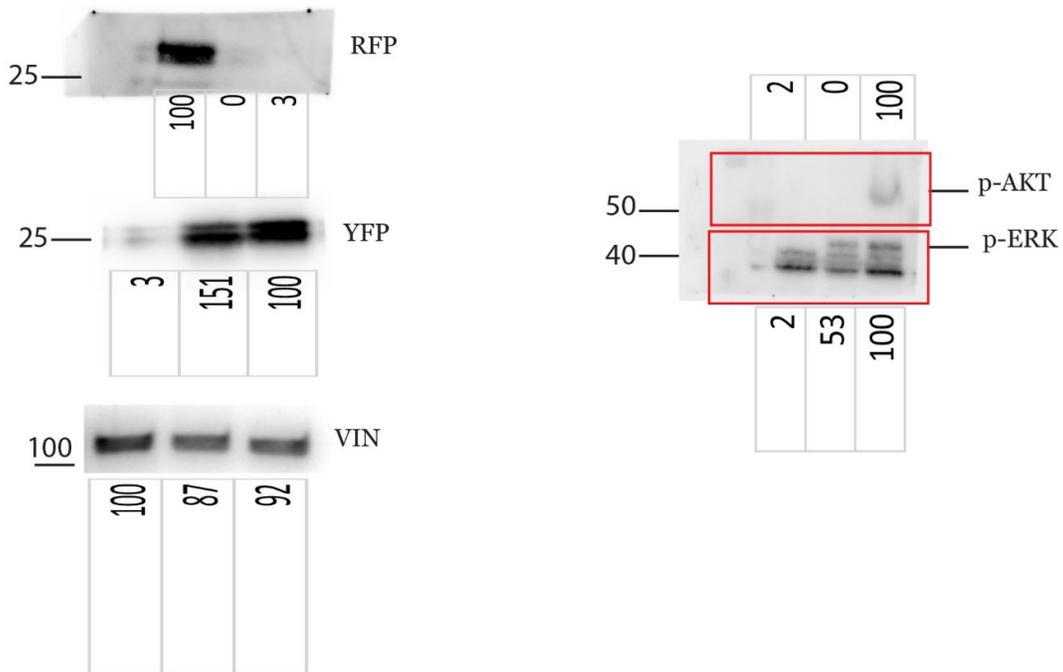
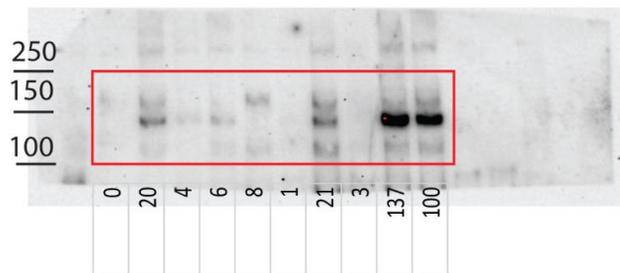


Figure 3



Flag (cas9) figure 5



Actin figure 5

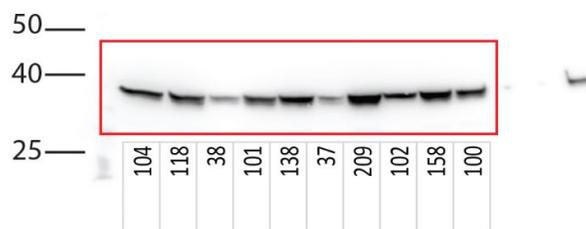


Figure S7. Uncropped Western Blot and Southern Blot images.

Table S1. Primers.

Primer:	Sequence:
KRAS_sgRNA_F	caccGACAAGATTTACCTCTATCGT
KRAS_sgRNA_R	aaacACGATAGAGGTAATCTTGTC
KRAS_seq_F	AGTGAAGTCATGGCCCACTC
KRASG12D_seq_F	AGGCCTGCTGAAAATGACTGA
KRAS_seq_R	AAACACCAAAACCCCATACG
TP53_sgRNA_F	caccGTGCGTGTTTGTGCCTGTCC
TP53_sgRNA_R	aaacGGACAGGCACAAACACGCAC
TP53_seq_F	GGCTTCTTGATCAGCTGGAG
TP53_seq_R	TCGCCATCCAGTGGCTTCTTC
STK11_sgRNA_F	caccGGTGGATGTGCTGTACAATG
STK11_sgRNA_R	aaacCATTGTACAGCACATCCACC
STK11_seq_F	ATTCTTTGGGGCTGCTCTCC
STK11_seq_R	AGCCATAGAGGGGGCAACTA
PTEN_sgRNA_F	caccGCAGCAATTCCTGTAAAGC
PTEN_sgRNA_R	aaacGCTTTACAGTGAATTGCTGC
PTEN_seq_F	TGGCCTCCCTATCTAATGG
PTEN_seq_R	CTCTGGTCCTTACTTCCCAT
NOTCH1_sgRNA_F	caccGCGTAGTCCACCACCCGCCG
NOTCH1_sgRNA_R	aaacCGGCGGTGTGGTGGACTACGC
NOTCH1_seq_F	GGGTACAGTCCCGTGGTGTT
NOTCH1_seq_R	GGCCCTGGGTGTCCTTAC
Activation_primary_F	CAGCCATTGCCTTTTATGGT
Activation_primary_R	TGTCGCCCTCGAACTTAC
Activation_nested_F	GCTGGTTGTTGTGCTGTCTC
Activation_nested_R	AAGTCGTGCTGCTTCATGTG
AAV_presence_F	ATCAGCAAGGAGATGATCGC
AAV_presence_R	TTAGCTGTATCGTCAAGGCACTC
AAV_titration_F	GGAACCCCTAGTGATGGAGTT
AAV_titration_R	CGGCCTCAGTGAGCGA