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CHINESE ACADEMY OF SCIENCES

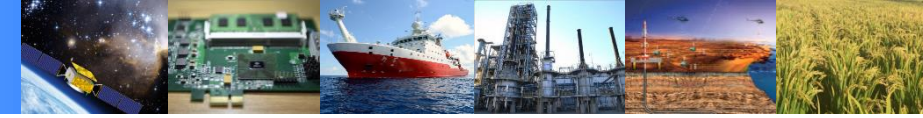
# State of Genetically Modified Pigs for Agriculture in China

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2021-08-31

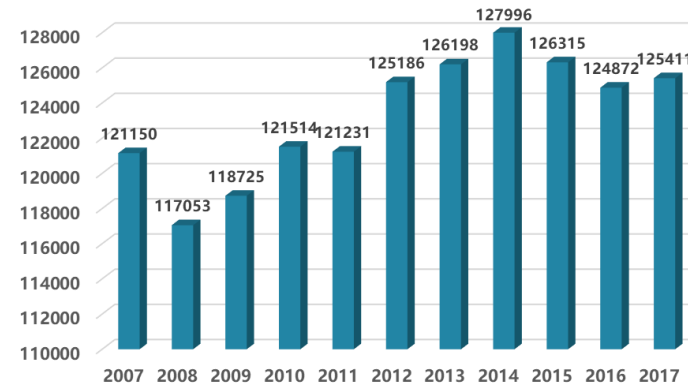




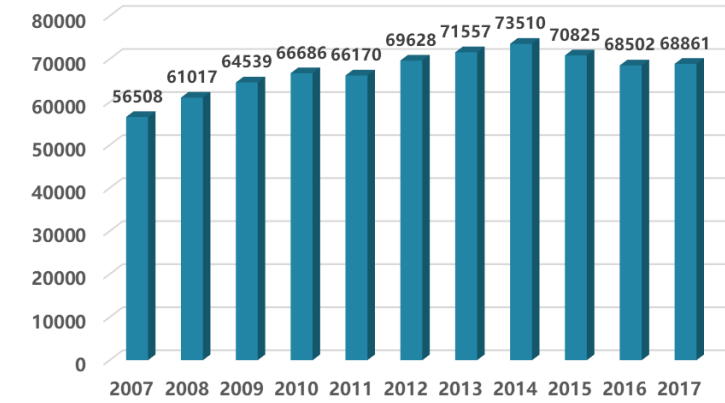
**Pork is the main meat food in China (65%)**

**China consumes more than 50% pork of the World**

**World: about 1.4 billion pigs; China: about 0.7 billion pigs**



The number of slaughtered pig in the world



The number of slaughtered pig in China

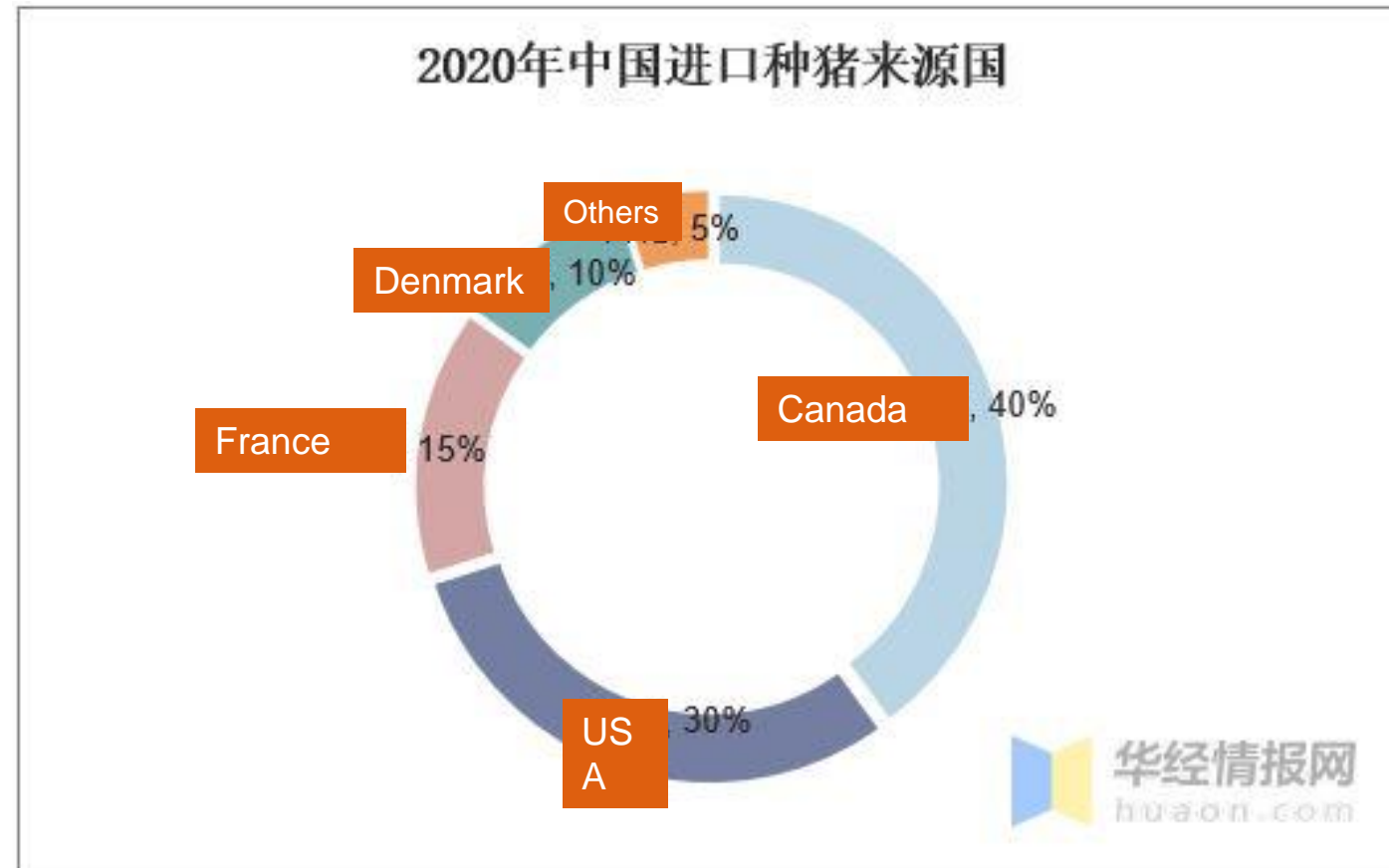


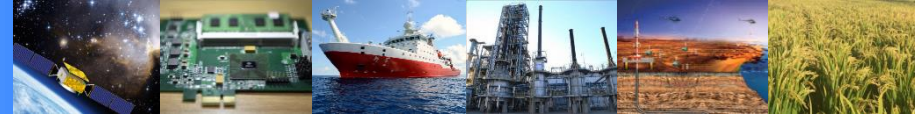


China has no native breeds of pigs with high productivity



Import 20 thousand elite boars from other countries each year





The Chinese government attaches importance to generation of genetically modified pig strains with favorable traits for agriculture

1. Setting up National special funding for transgenic livestock
2. Setting up big pig facilities for large scale characterization of GM pigs
3. More than 15 research teams working on GM pigs



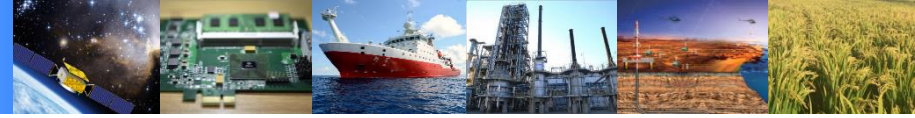




# The genetical modification of pigs can be made to:

- ✓ Promote growth: GH, IGF
- ✓ Improve meat quality: FAT1, FAT2, MSTN
- ✓ Reduce environmental emissions: NSP-degrading and phytate-degrading enzymes
- ✓ Gain resistance to pathogens or other environmental stress





# 1. Genetical Modification of pigs for promoting growth



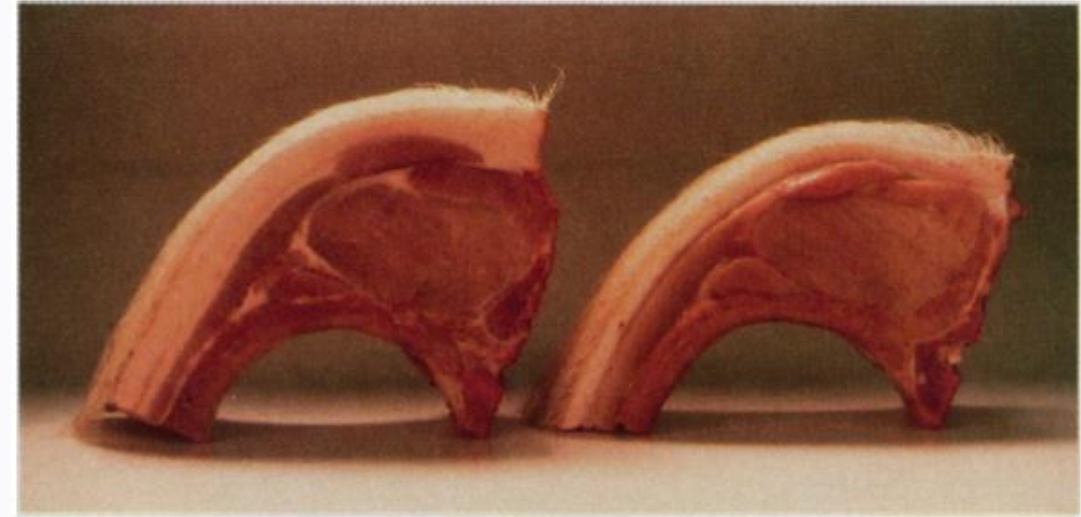


# Over expression of growth hormone

Significant improvements in both daily weight gain and feed efficiency

Group	Average daily weight gain (g)	Feed efficiency (kg feed/kg gain)
<b>Founder animals*</b>		
Control	743 ± 32 (6)	3.12 ± 0.15
Transgenic	690 ± 65 (6)	2.62 ± 0.12
	<i>P</i> = 0.480	<i>P</i> = 0.026
<b>37-06 G2 progeny†</b>		
Control	760 ± 24 (8)	2.99 ± 0.12 (8)
Transgenic	874 ± 30 (5)	2.46 ± 0.16 (5)
	<i>P</i> = 0.016	<i>P</i> = 0.026
<b>37-06 G3 progeny‡</b>		
Control	867 ± 21 (15)	ND
Transgenic	933 ± 31 (8)	
	<i>P</i> = 0.098	
<b>31-04 G2 progeny§</b>		
Control	869 ± 44 (7)	ND
Transgenic	988 ± 62 (7)	
	<i>P</i> = 0.15	
<b>Combined progeny</b>		
Control	815 ± 17 (30)	ND
Transgenic	905 ± 21 (20)	
	<i>P</i> = 0.001	

a marked reduction in subcutaneous fat



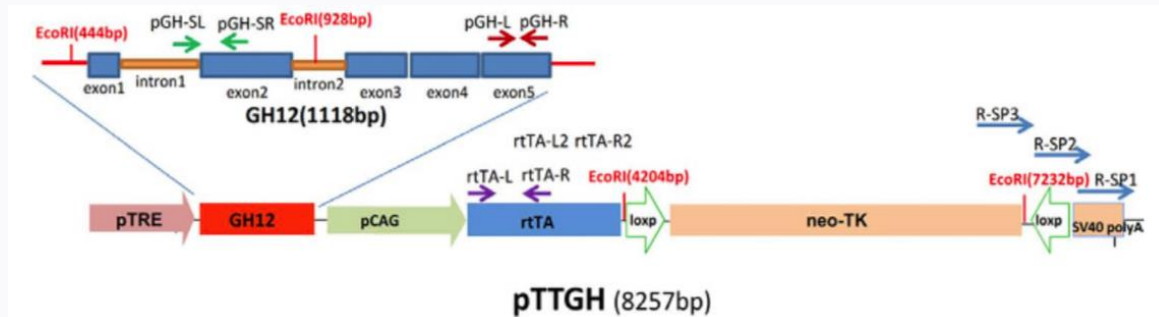
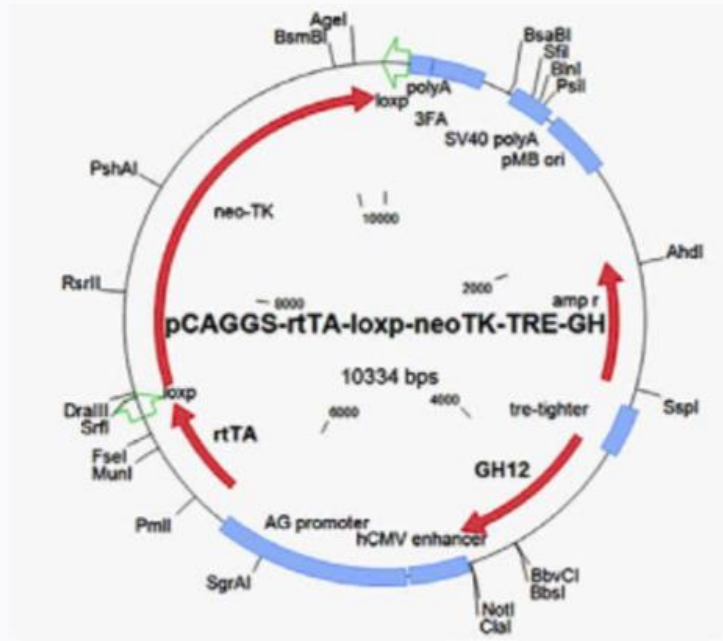
pathological changes

Diagnosis*	Number of animals	
	Transgenic	Control
Gastric ulcers	5/5	0/3
Synovitis	4/5	0/3
Cardiac myocyte nuclear hypertrophy	4/5	0/3
Dermatitis	4/5	1/3
Nephritis	3/5	0/3
Pneumonia	3/5	1/3

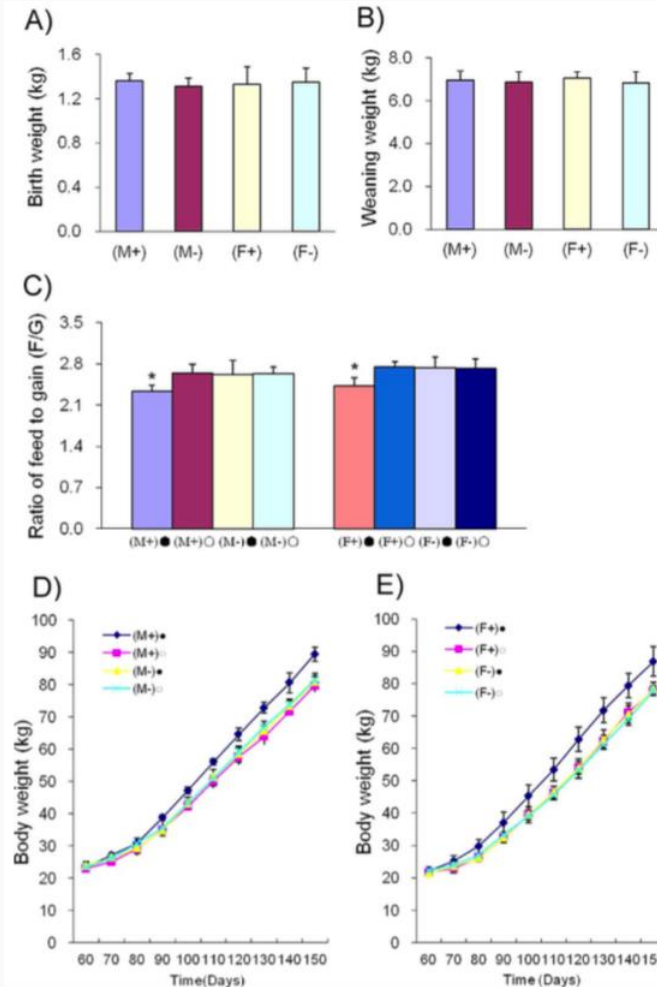


# The transgenic pig with controllable expression of growth hormone

the pCAGGS-rtTA-TRE-GH12 (pTTGH) vector



Growth rates and feed efficiencies







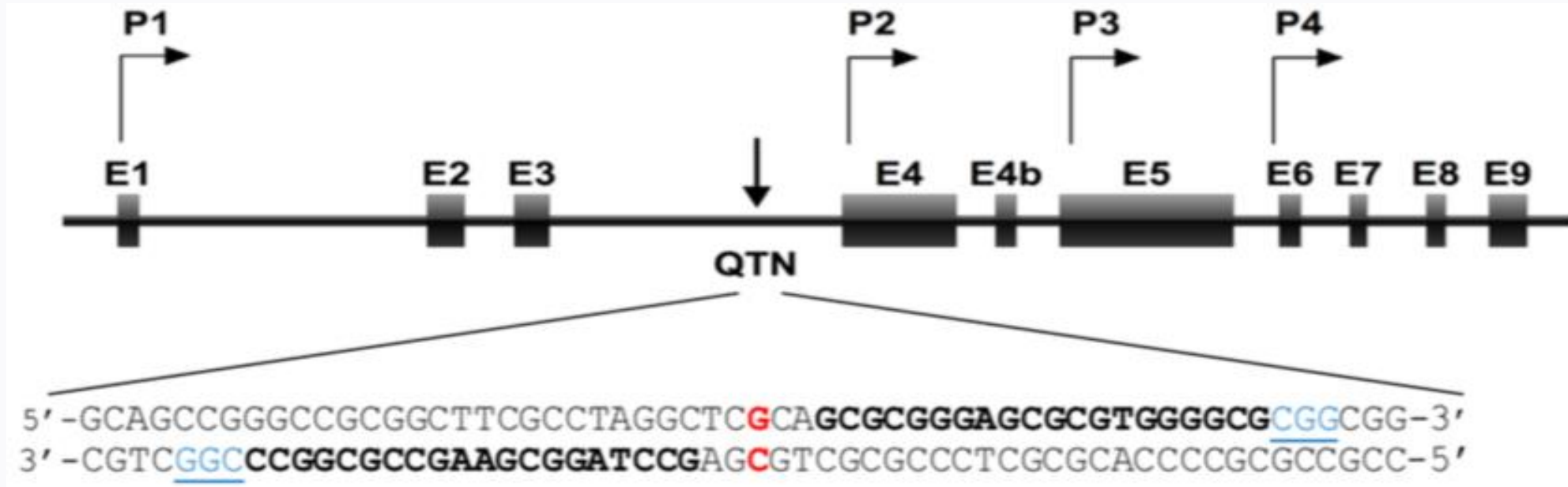
## Editing porcine *IGF2* regulatory element to improve meat production in Chinese Bama pigs

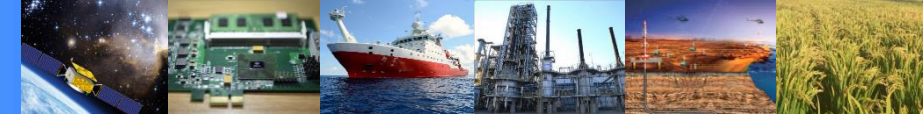
*IGF2*, an important growth factor which affects skeletal muscle and fat deposition

*IGF2*-intron 3–nucleotide 3072:

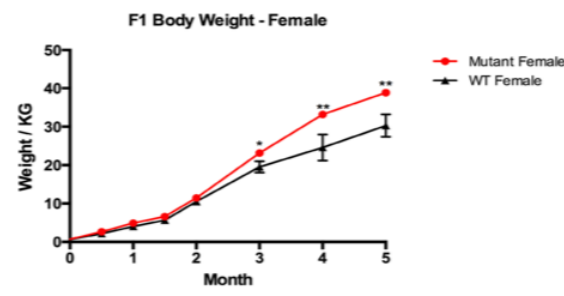
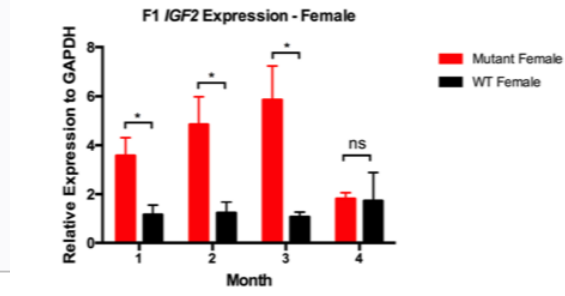
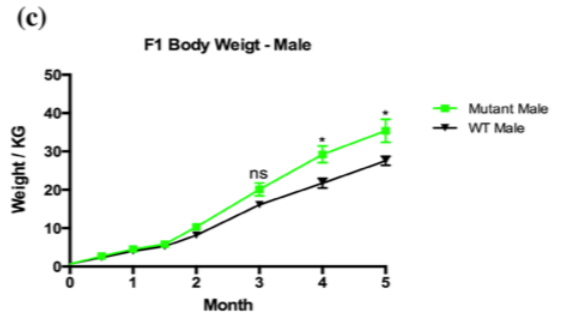
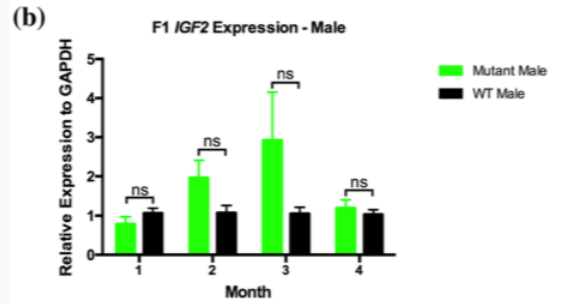
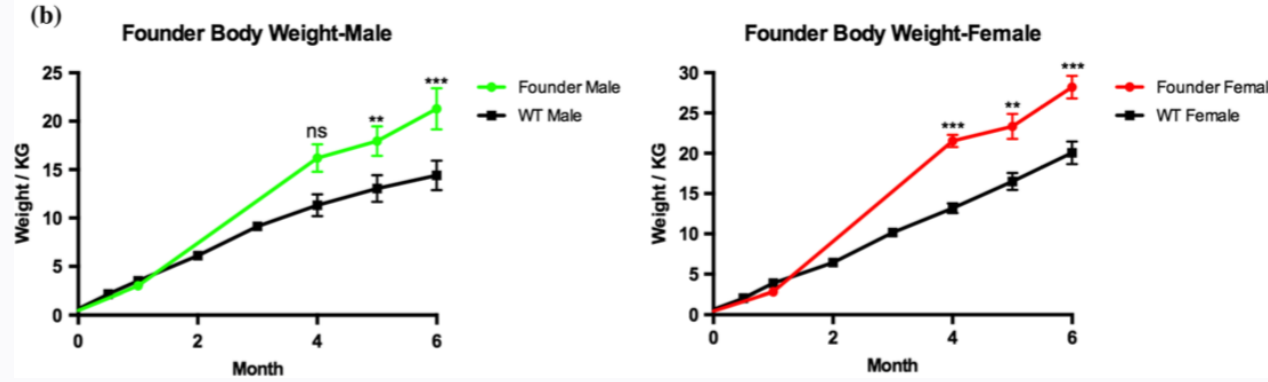
5'-GCTCGC-3', the G allele, recognized by repressor ZBED6, negatively regulates *IGF2* expression

### Experiment Design

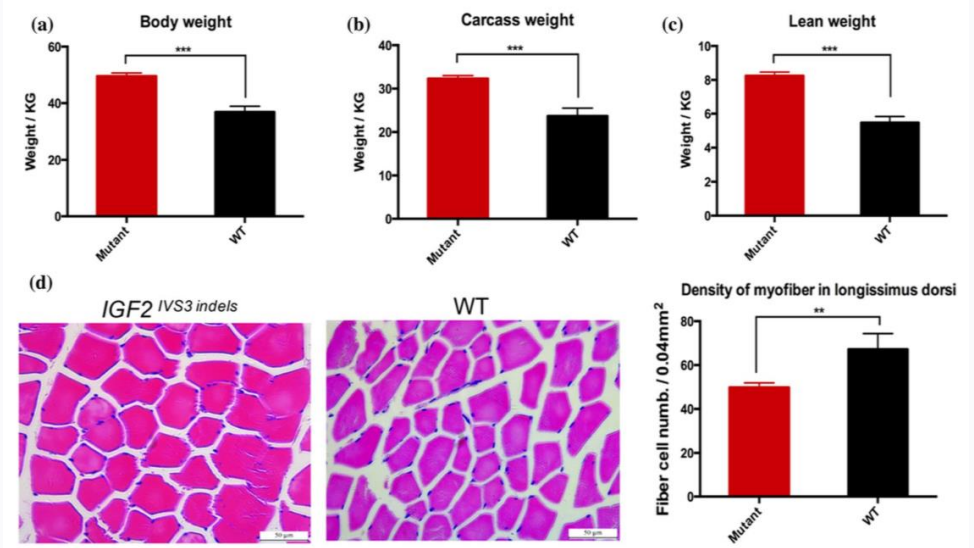
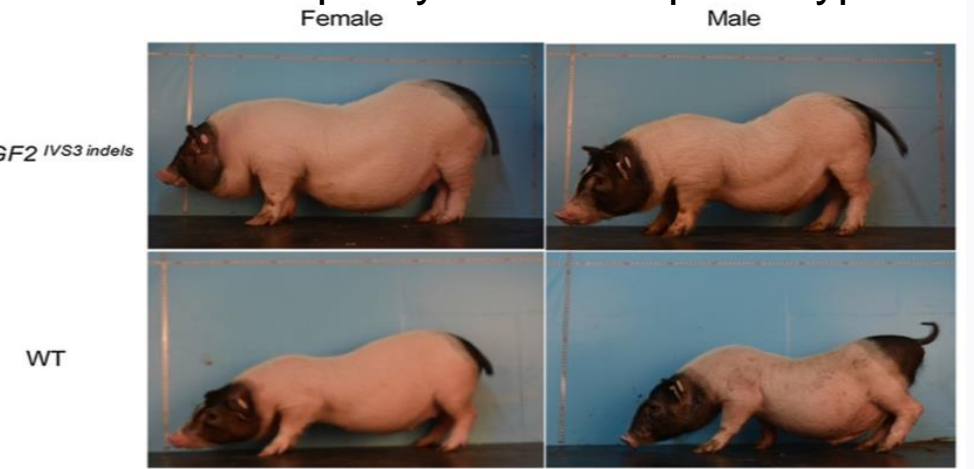




## Growth performance of founder and F1 gene-edited pigs



## Meat quality and other phenotypes





## 2. Improvement of the pig meat quality



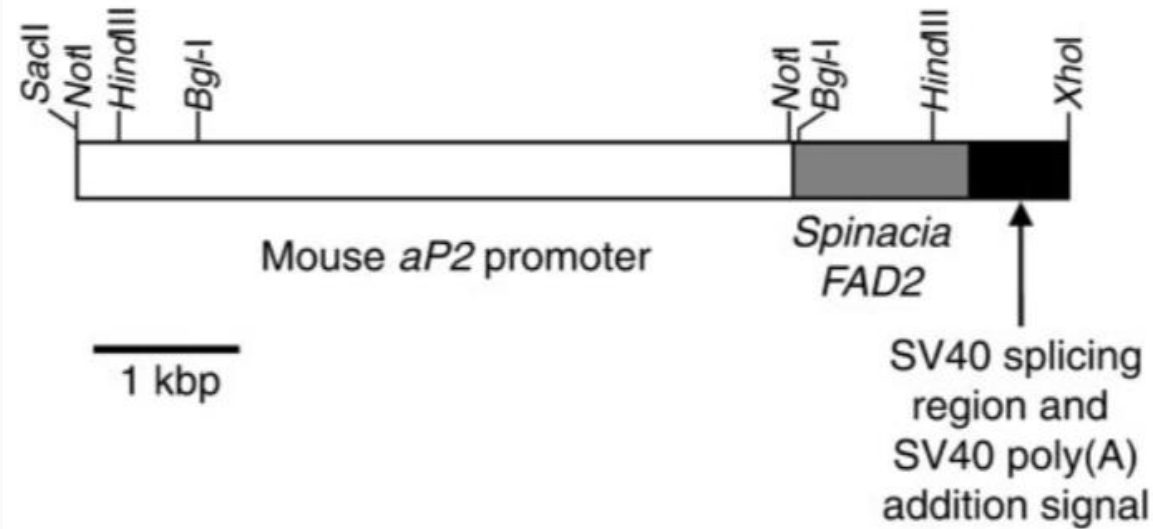


## Expression of FAD2 in transgenic pigs promotes synthesis of polyunsaturated fatty acids

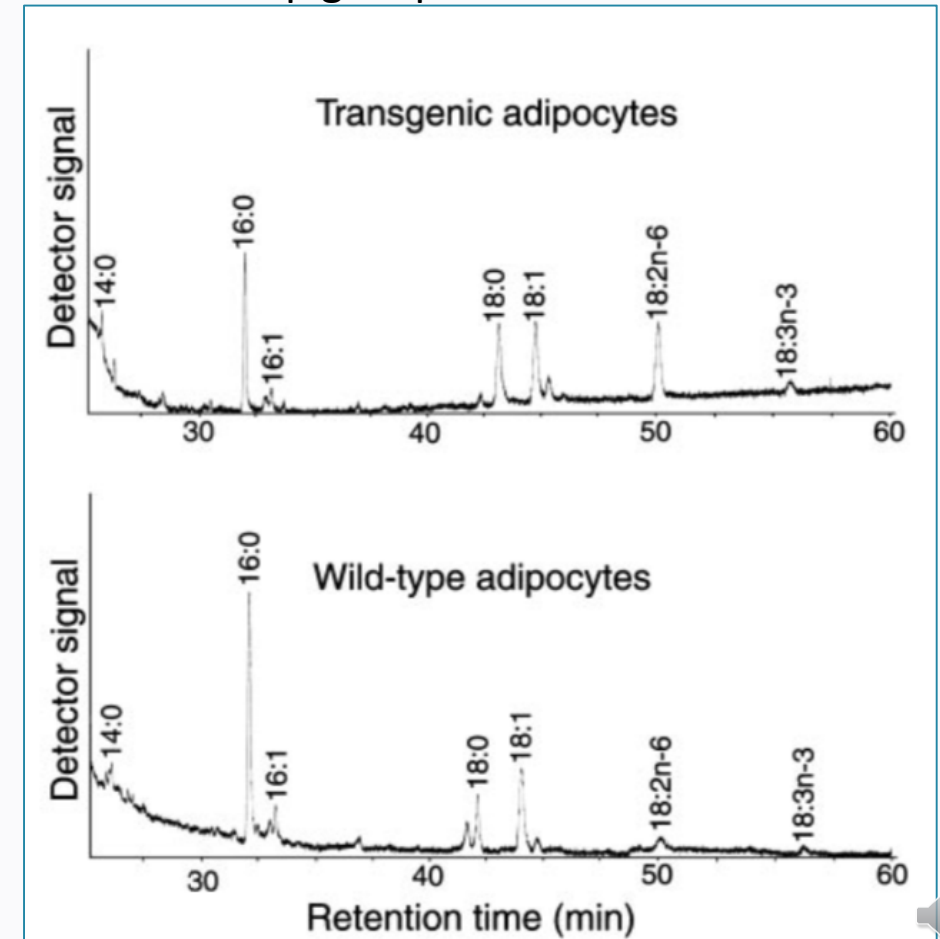
Mammals lack the desaturases required for synthesis of polyunsaturated fatty acids

FAD2: a delta12 fatty acid desaturase from spinach

The coding region of cDNA for a 12 desaturase (FAD2) from *Arabidopsis thaliana*



fatty acid composition of accumulated lipids in pig adipose cells.



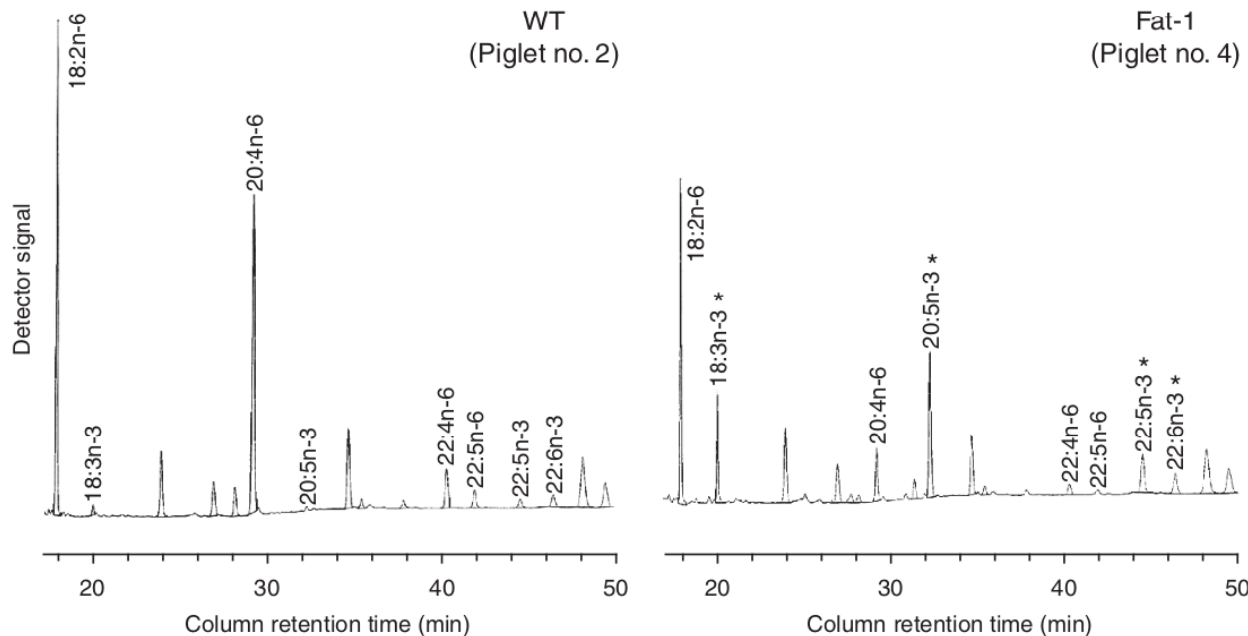




## The delta-15 desaturase (*fat1*) gene pigs synthesize n-3 PUFAs from n-6 PUFAs

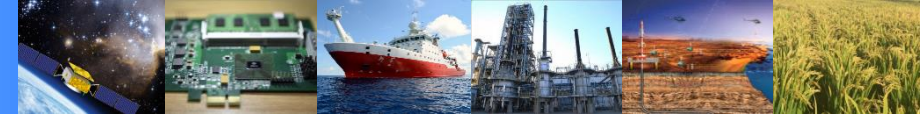
Transgenic pigs with *hfat-1*, humanized *Caenorhabditis elegans* gene, encoding an n-3 fatty acid desaturase

the polyunsaturated fatty acid profiles of total lipids



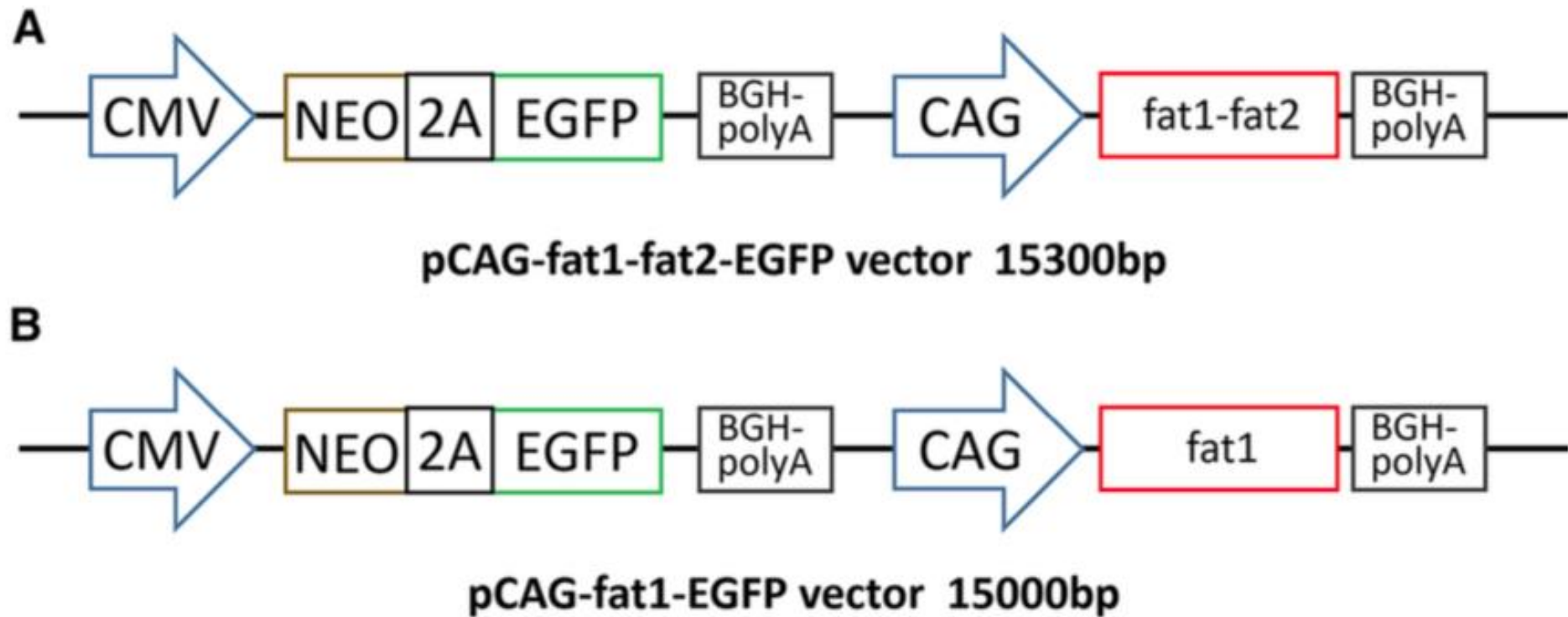
**Table 1** *n-3* and *n-6* fatty acids concentration and *n-6/n-3* ratios in tail samples from *hfat-1* transgenic and wild-type piglets

Fatty acids in tails <sup>a</sup>	Transgenic piglets ( <i>n</i> = 8)	Wild-type piglets ( <i>n</i> = 8)
ALA (18:3 <i>n-3</i> , %)	0.94 ± 0.10	0.63 ± 0.04
EPA (20:5 <i>n-3</i> , %)	4.21 ± 0.60	0.26 ± 0.07
DPA (22:5 <i>n-3</i> , %)	1.69 ± 0.19	0.35 ± 0.05
DHA (22:6 <i>n-3</i> , %)	1.75 ± 0.23	0.95 ± 0.21
Total <i>n-3</i> FA (%)	8.59 ± 0.84	2.18 ± 0.25
Total <i>n-6</i> FA (%)	14.28 ± 1.31	18.46 ± 1.41
<i>n-6/n-3</i> ratio	1.69 ± 0.30	8.52 ± 0.62



## Co-expression of fat1 and FAD2 in transgenic pigs

Endogenously produced n-6 PUFAs was used as substrates to synthesize n-3 PUFAs

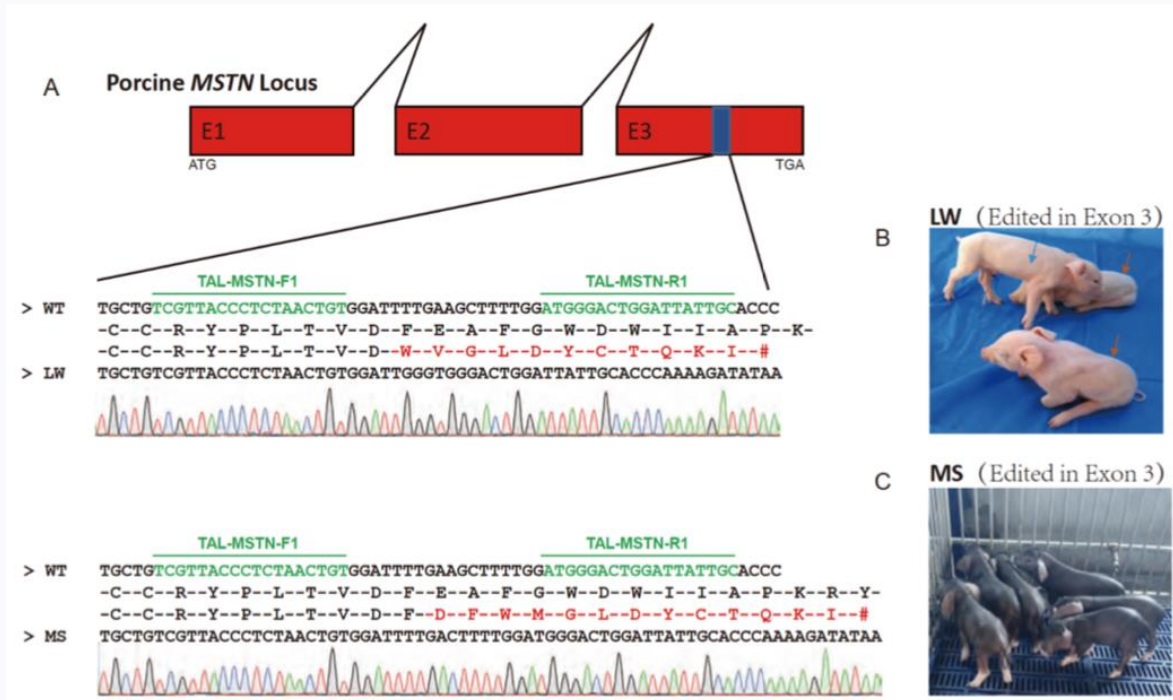




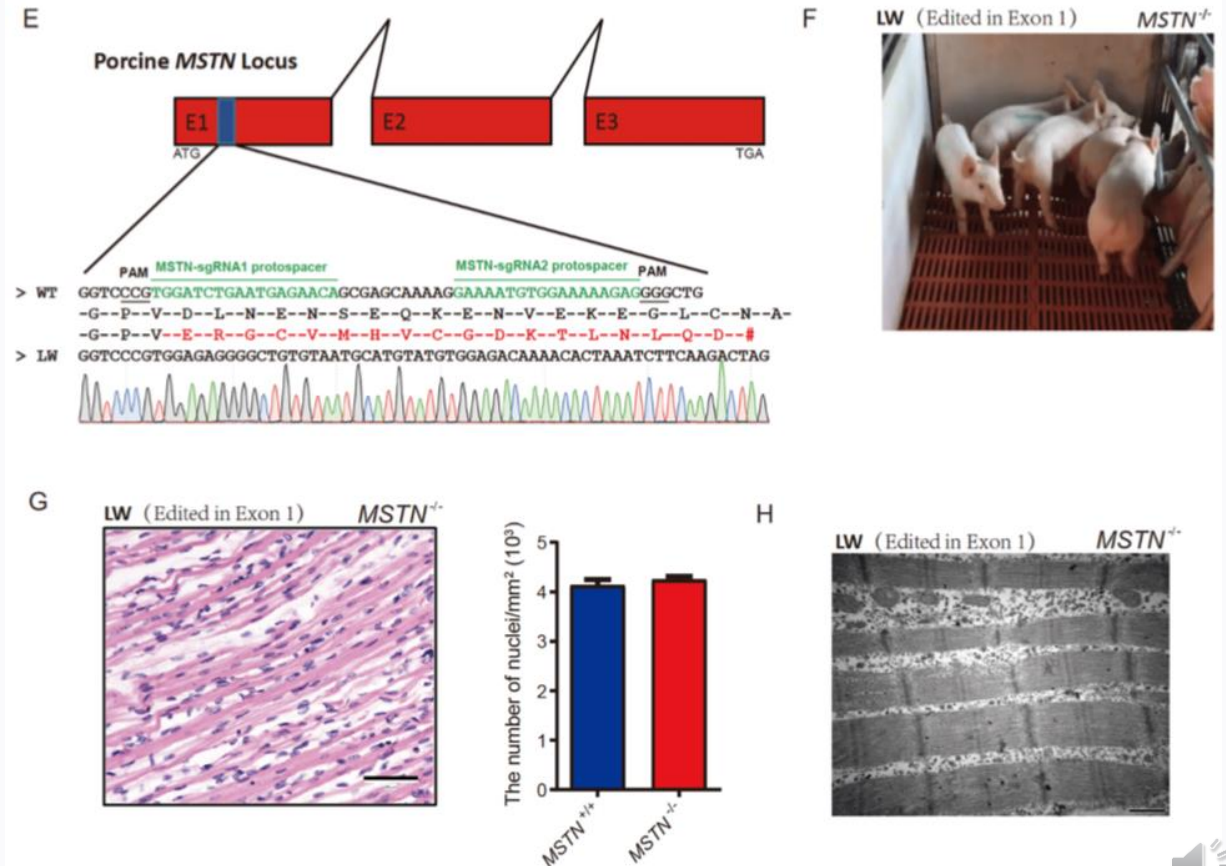
# MSTN-edited pigs to overcome lameness and sustainably improve nutritional meat production

Genes related to muscle mass: *myostatin (MSTN)*, negatively regulates skeletal muscle cell proliferation

hindlimb weakness



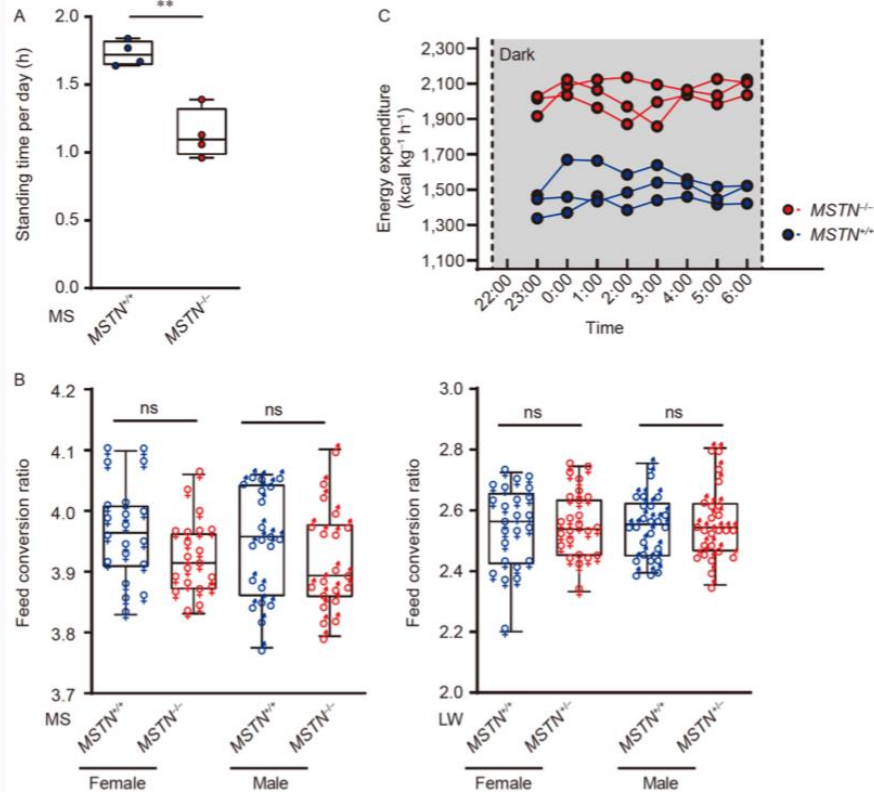
Alternative edit-site-based solution avoids ER stress and overcomes the hindlimb weakness



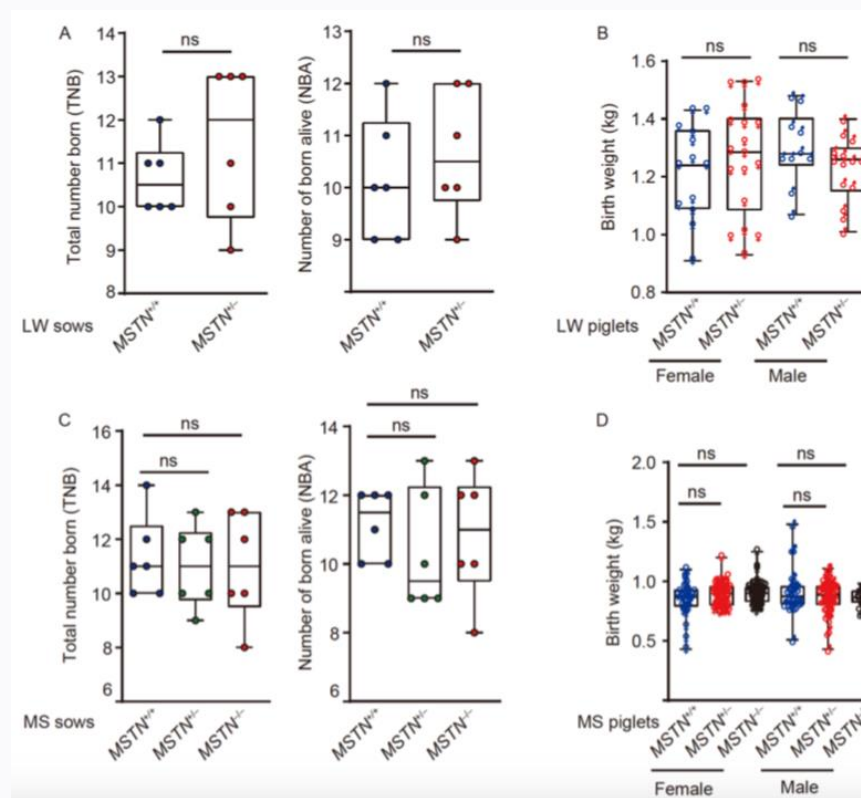




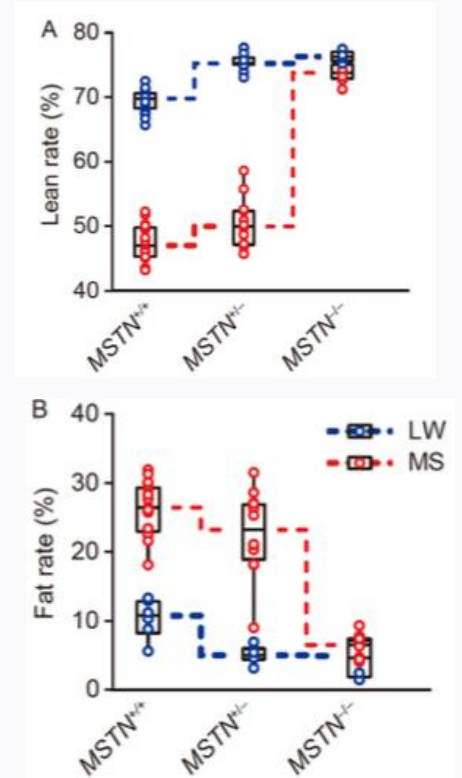
## Similar feed-conversion ratio



## No dystocia or significant effects on the maternal reproductive traits



## different trends of change for lean and fat rates







## 3. Reduction of environmental impact

Only 1/3 of feed nitrogen and phosphorus were utilized from feedstuff diets in pig production. Inefficient feed digestion can cause serious nutrient emissions to the environment.

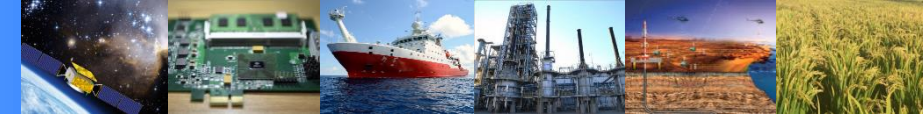
- Phytates, negatively charged saturated cyclic acids, bind to positively charged molecules in the diet such as minerals and protein, thereby reducing nutrient digestibility and increasing discharge of the unabsorbed nutrients to the environment.
- Pigs are inherently incapable of digesting Non-starch polysaccharides that are primarily present in plant cell walls.



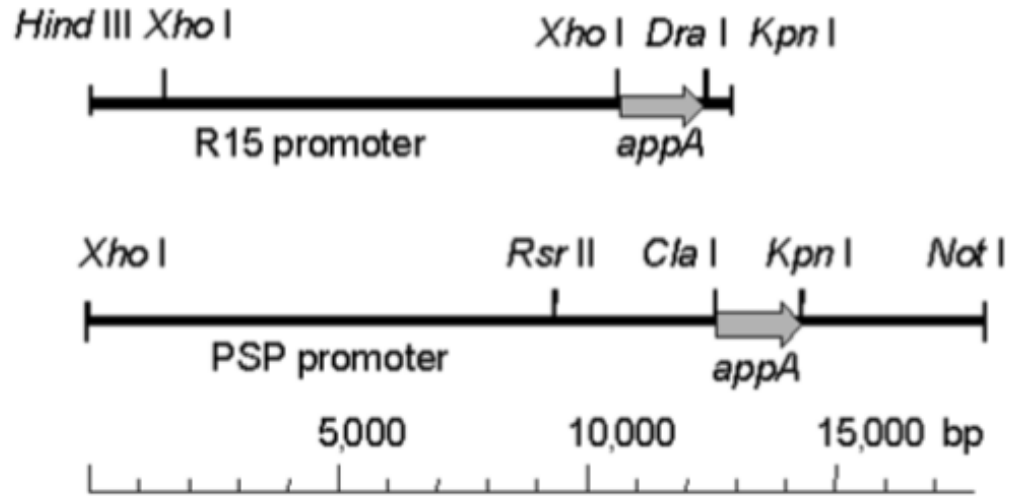
# Salivary gland-specific expression of phytase

Phytate phosphorus passes undigested, most important manure pollutant

Phytase: allows the pigs to digest the phosphorus in phytate

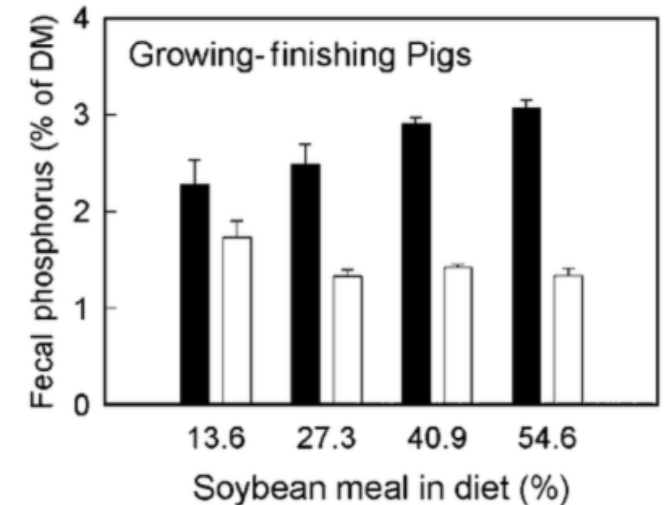
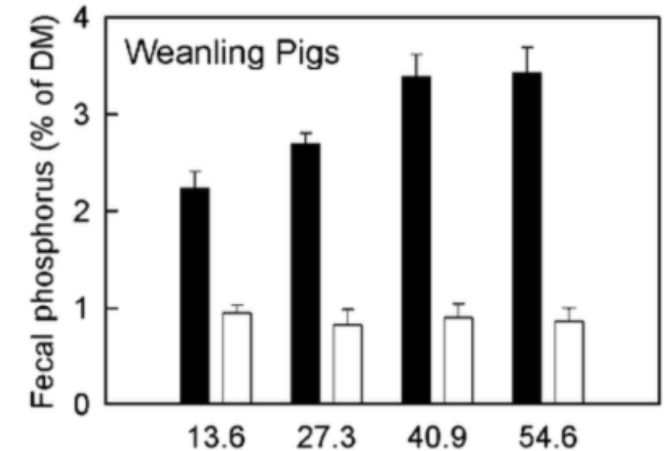


Phytase transgene constructs



◆ salivary gland-specific expression promoter: PSP

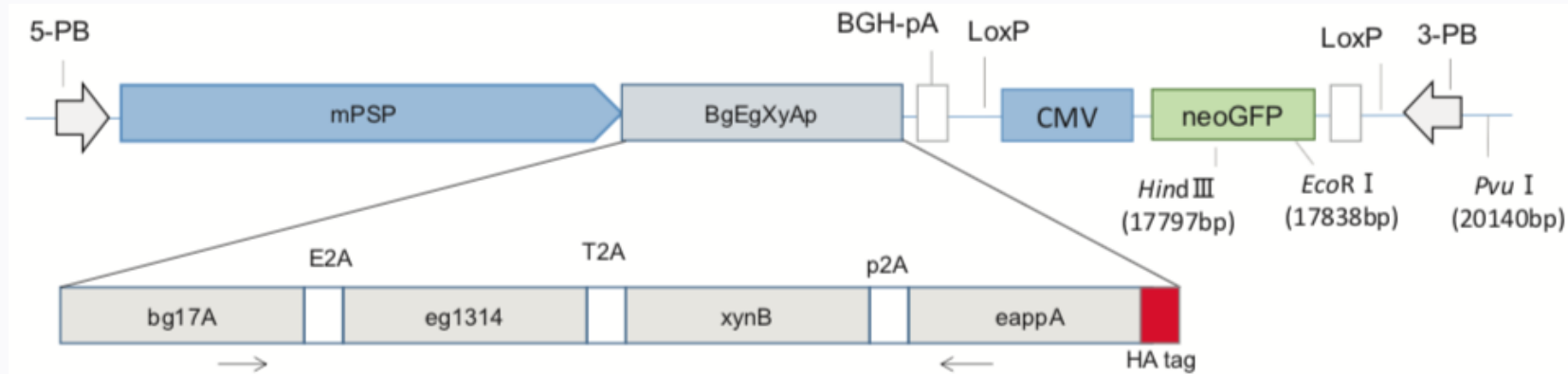
Total phosphorus content of fecal matter





## Expression of three microbial enzymes, b-glucanase, xylanase, and phytase in the salivary glands

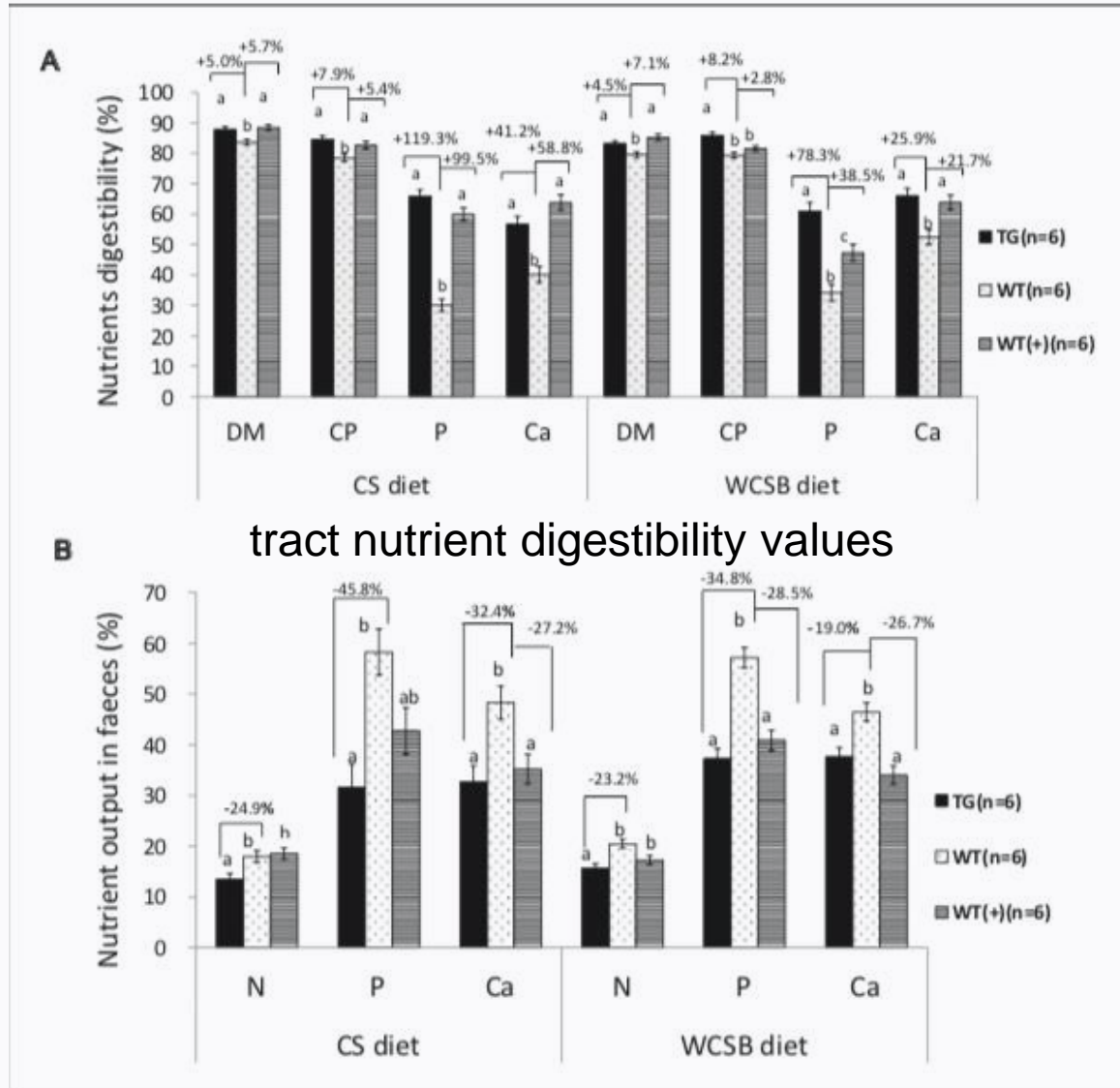
### Experiment Design



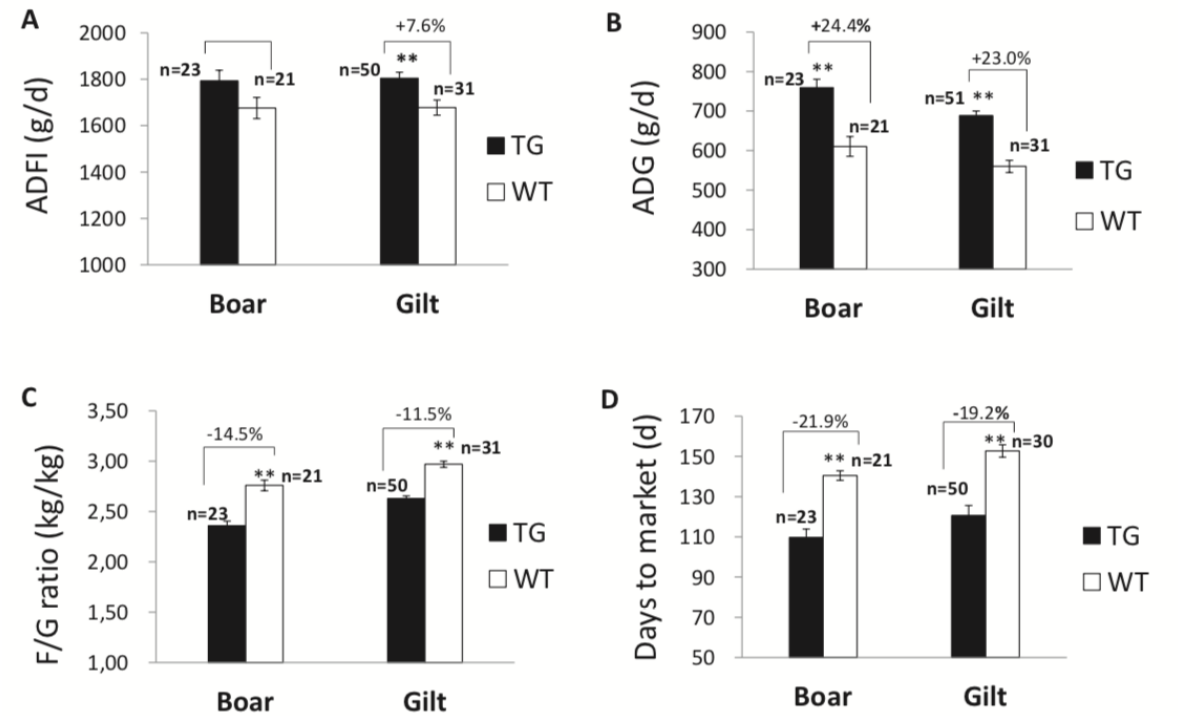
- ◆ salivary gland-specific expression promoter: mPSP
- ◆ **NSP-degrading enzymes**: two  $\beta$ -glucanases genes (bg17A and eg1314), a xylanase gene (xynB)
- ◆ **Phytate-degrading enzyme**: phytase gene (eappA)



## fecal nutrient output



## Growth performance







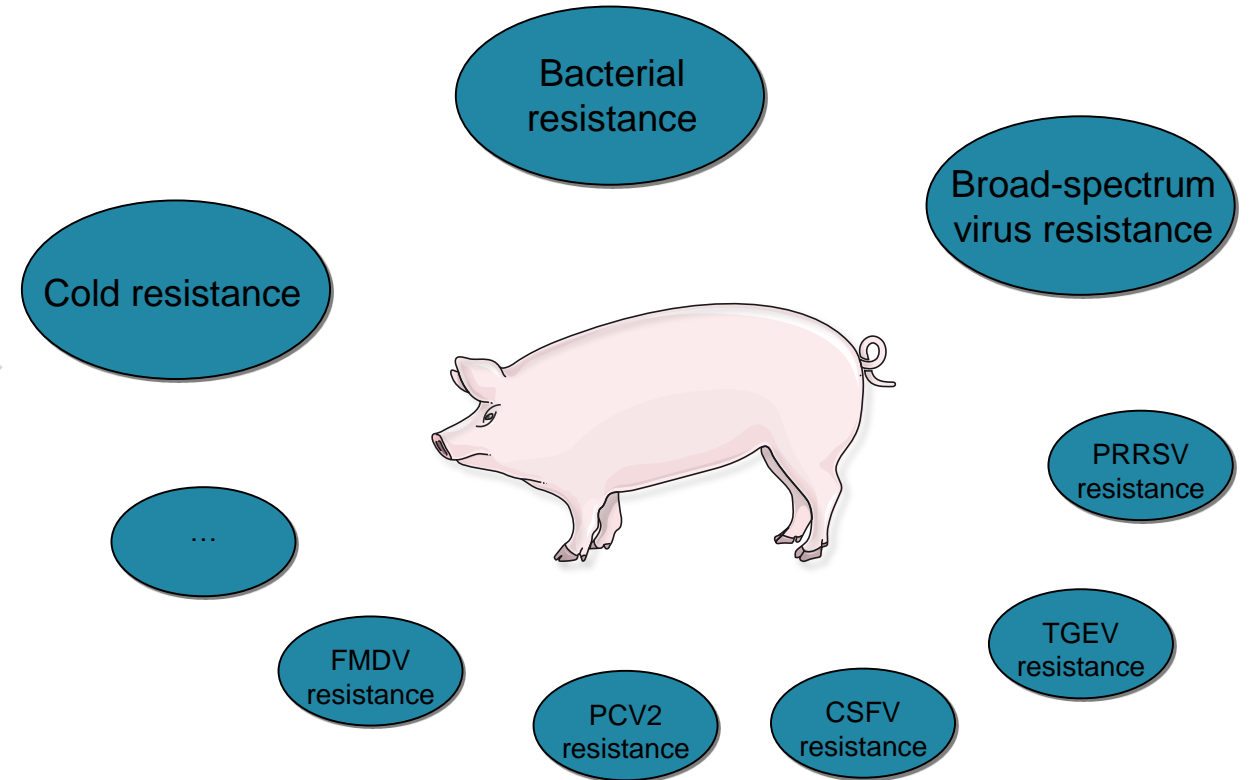
# 4. Stress resistance pigs

Overexpressing endogenous resistance genes

Introducing exogenous resistance genes

Editing pathogens targeting receptor genes

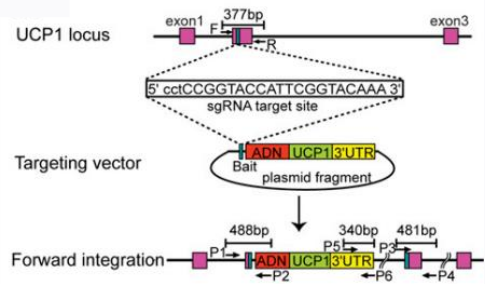
Introducing synthetical virus-killing genes



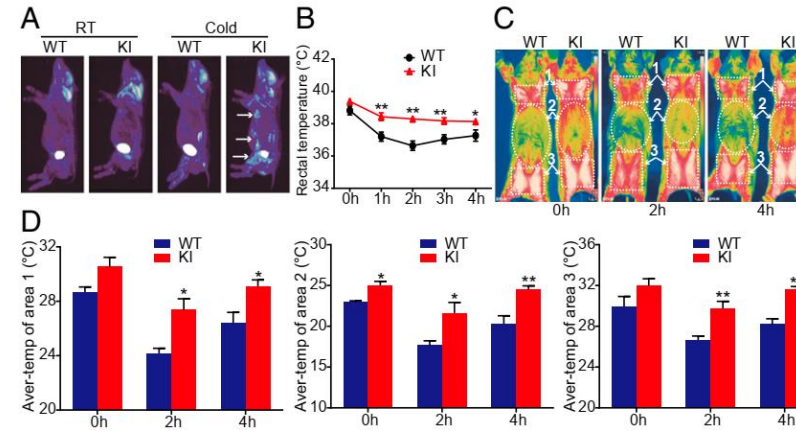
# Generation of adiponectin-UCP1 KI transgenic pigs: Cold resistance transgene pigs

Pigs lack a functional UCP1 gene, resulting in poor thermoregulation and susceptibility to cold.

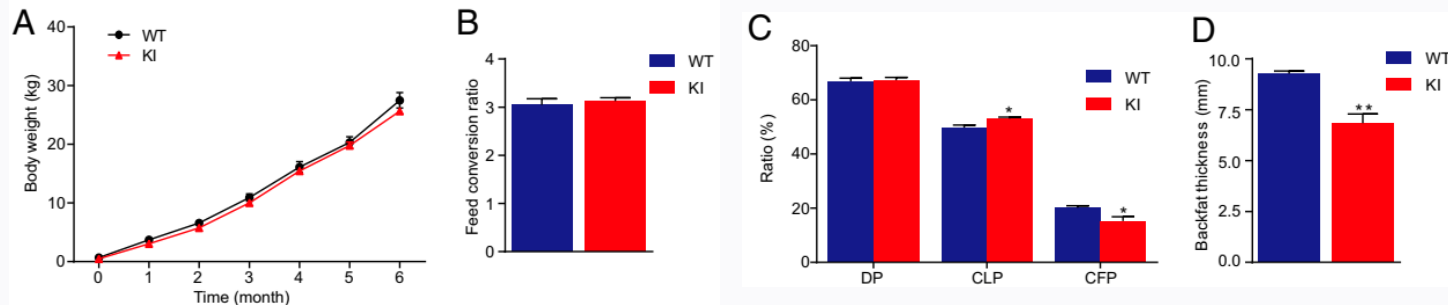
## Generation of adiponectin-UCP1 KI transgenic pigs



## UCP1 KI pigs exhibited improved thermoregulation



## Decreased Fat Deposition and Increased Adipose Lipolysis in UCP1 KI Pigs





# Anti-bacterial transgenic pigs

Lysozyme has Broad-spectrum antibacterial activities

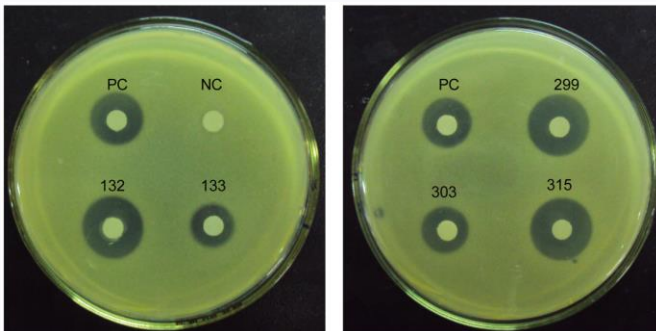


- Bacillus subtilis*
- Bacillus cereus*,
- Staphylococcus aureus*,
- Escherichia coli*,
- Klebsiella pneumoniae*,
- Pseudomonas aeruginosa*,
- Streptococcus agalactiae*,
- Salmonella typhimurium*,
- ...

Human lysozyme gene transgenic pigs



Transgenic pigs milk can inhibit growth of *E.coli*



Transgenic pigs milk can inhibit the growth of *E.coli* in the duodenum of sucking pigs

Item	Non-transgenic	Transgenic	p-value
<i>Escherichia coli</i>			
Duodenum	7.62±0.24	6.56±0.17	<0.001
Jejunum	7.00±0.39	6.85±0.66	0.707
Ileum	7.70±0.39	7.58±0.43	0.691
Colon	7.09±0.35	6.77±0.34	0.236

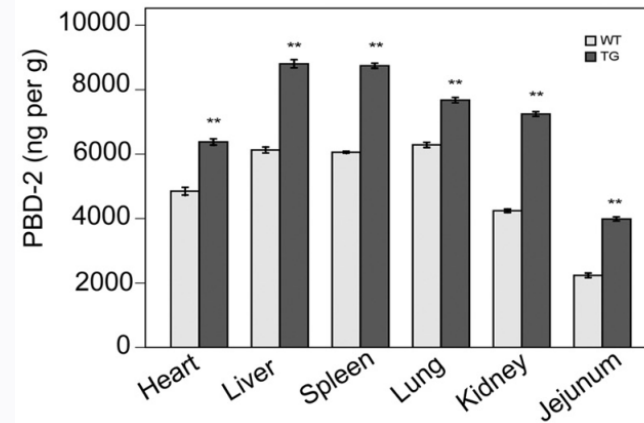


# Anti-bacterial transgenic pigs

Porcine beta-defensin 2 (PBD-2) can against *Actinobacillus pleuropneumoniae*

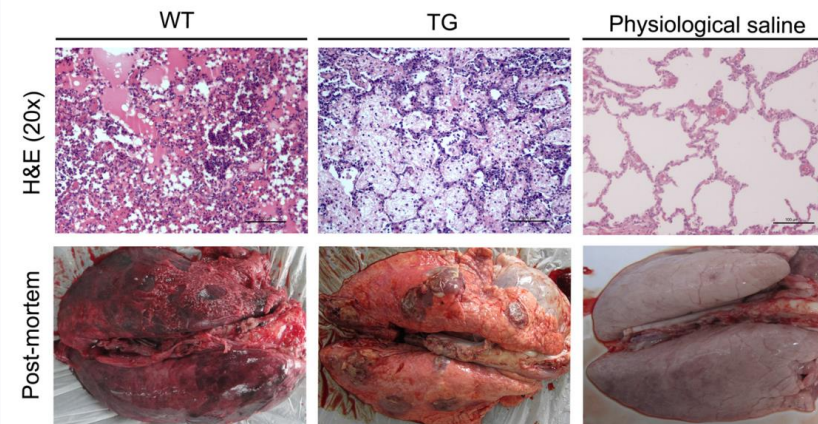
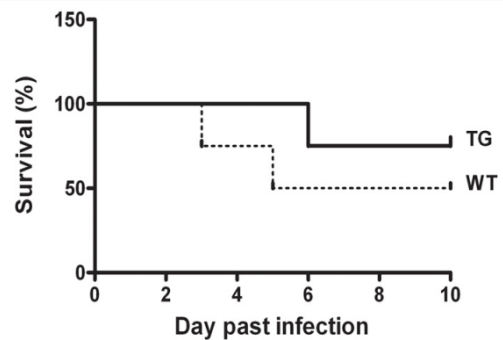
*Actinobacillus pleuropneumoniae* is an important respiratory pathogen causing porcine contagious pleuropneumonia

Construct of transgenic pigs overexpressing PBD-2



PBD-2 gene is driven by the CAG promoter

Overexpression of porcine beta-defensin 2 (PBD-2) enhanced resistance to cohabitation infection by *A. pleuropneumoniae*.

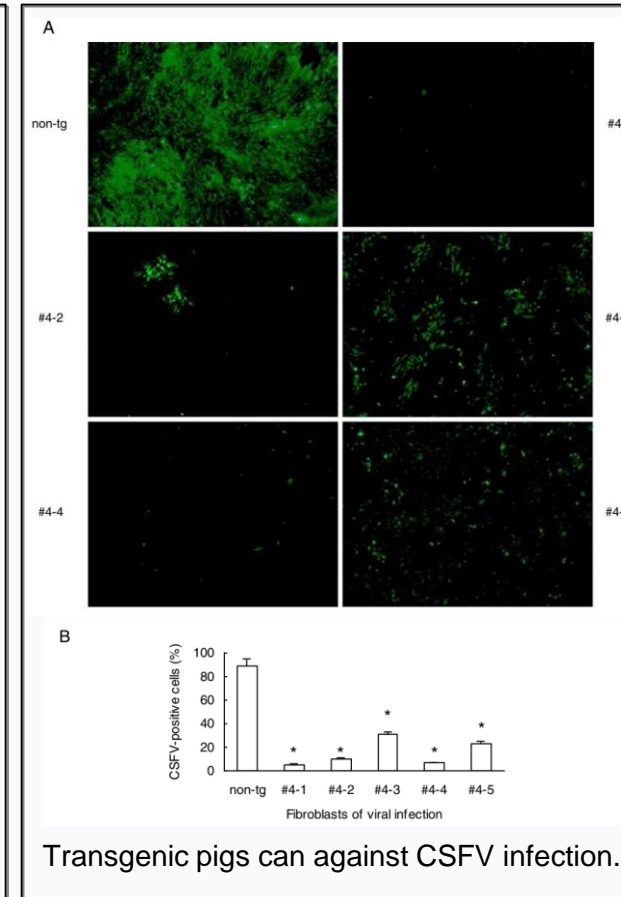
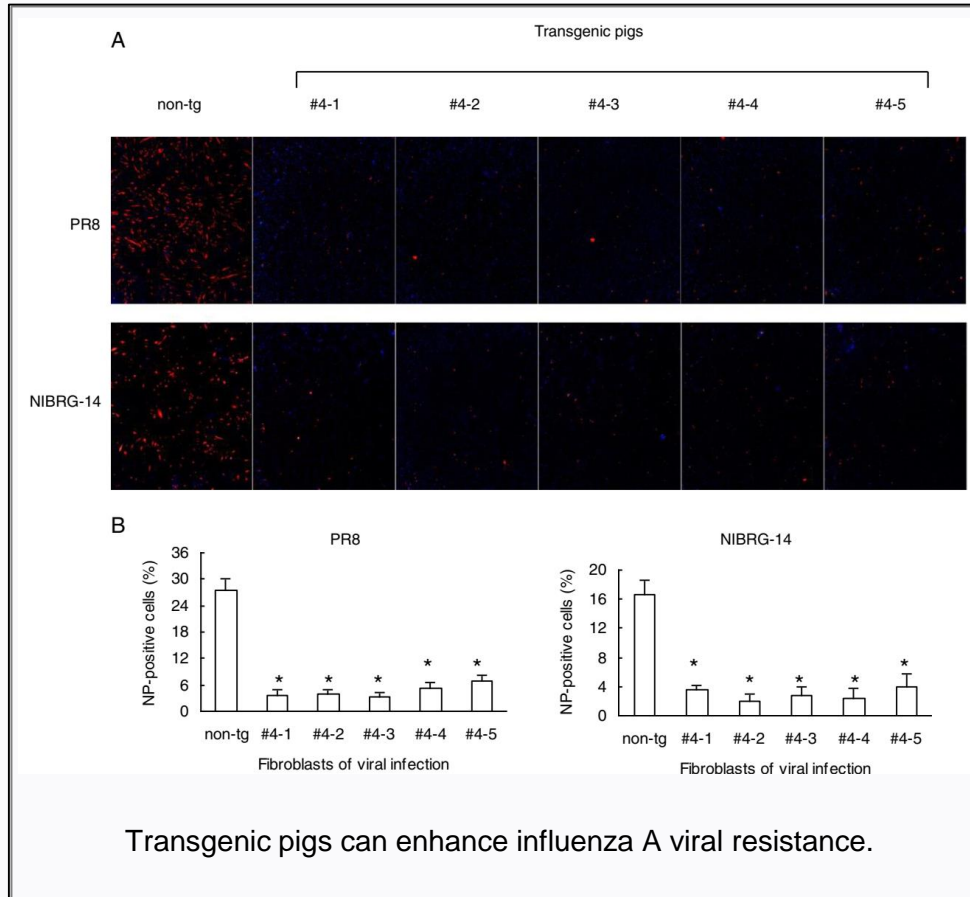


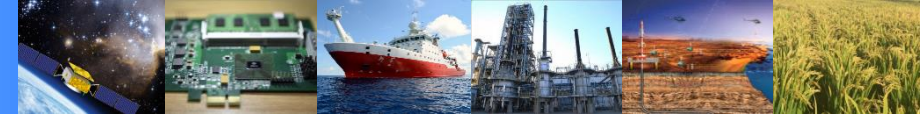




# Broad-spectrum antiviral transgenic pigs

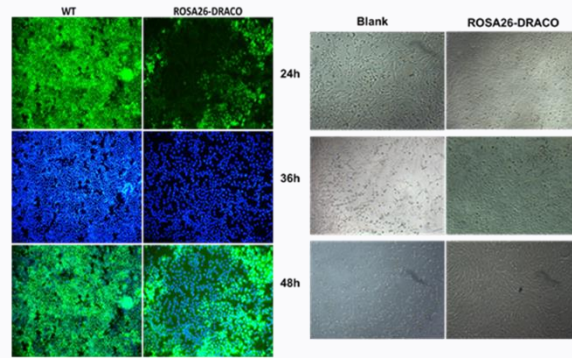
1. The transgenic pigs over-expressing myxovirus resistance gene have broad-spectrum antiviral activities





# Broad-spectrum antiviral transgenic pigs

## 2. The transgenic pigs express DRACO gene have broad-spectrum antiviral activities



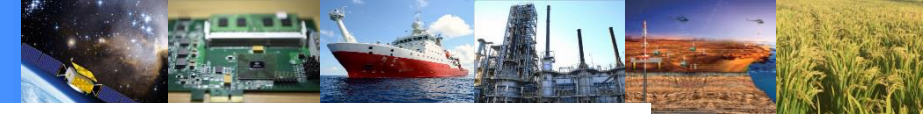
Transgenic pigs can enhance CSFV and PRV resistance

## 3. Gene editing pigs enhance NLRP3 expression have broad-spectrum antiviral activities



Transgenic pigs can enhance NLRP3 expression during the stimulation

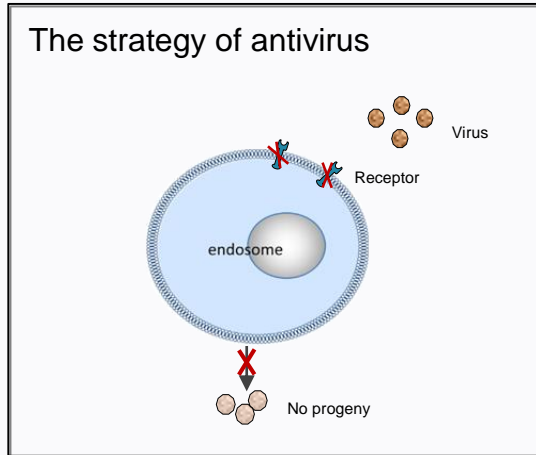




# Specific viral resistance transgenic pigs

## 1. Knockout virus entry receptors CD163 and porcine aminopeptidase N (pAPN) Gene

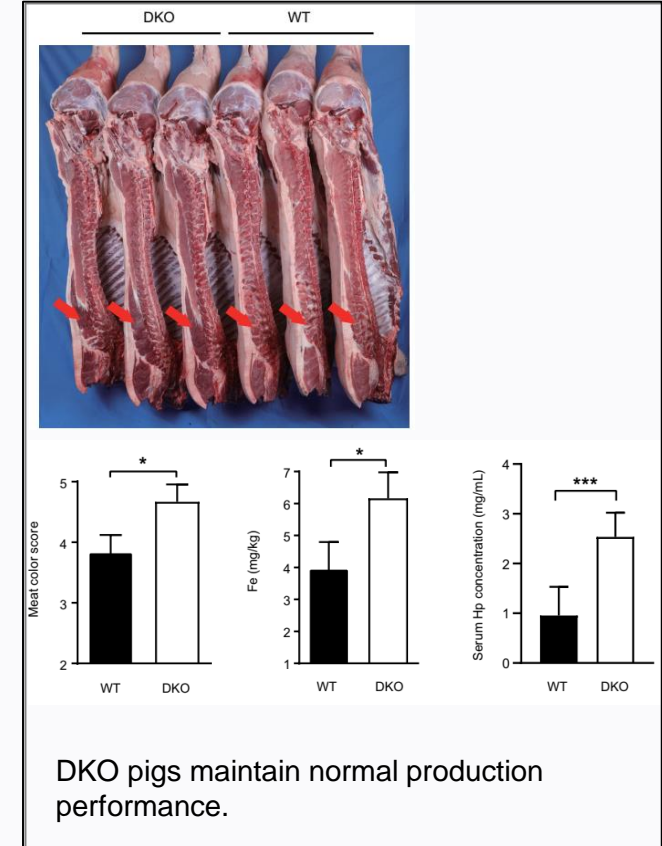
pigs are resistant to PRRSV and porcine transmissible gastroenteritis virus (TGEV) and decreased susceptibility to deltacoronavirus (PDCoV)

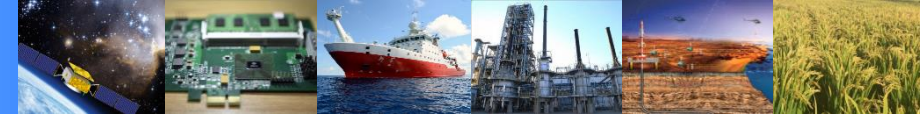


CD163&pAPN DKO pigs

	WT (Infected)	DKO (Infected)
TGEV (3 dpi)		
TGEV (14 dpi)		
PDCoV		

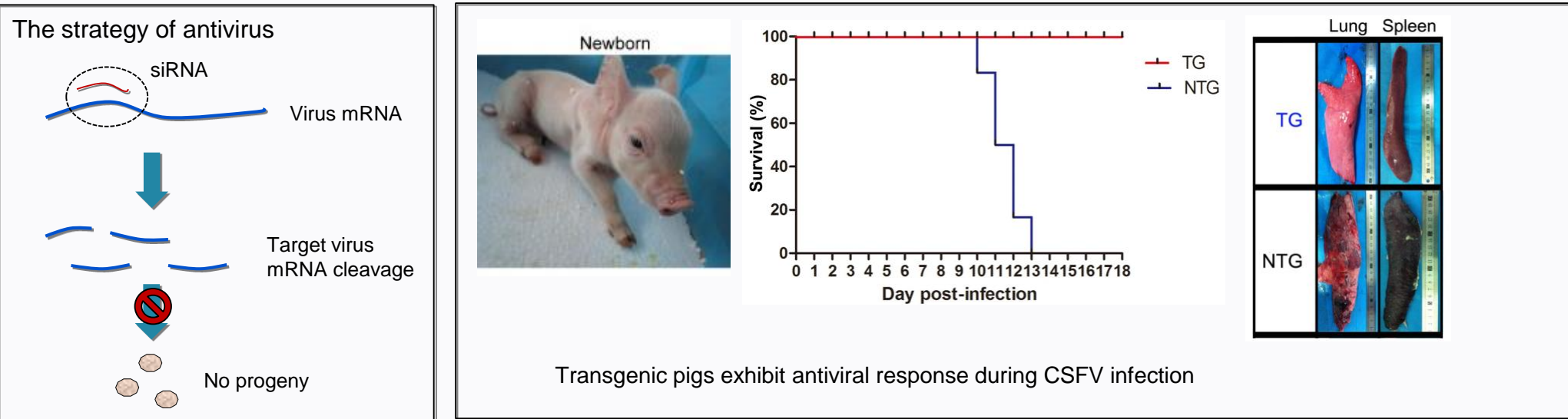
CD163 and pAPN DKO pigs are completely resistant to genotype 2 PRRSV and TGEV, and also decreased susceptibility to PDCoV.





# Specific viral resistance transgenic pigs

## 2. Constructing CSFV resistance pigs through knockin the shRNAs of targeting virus RNA



### Others anti-viral transgenic pigs through introducing RNA interference

Type of virus	Journal	Organization
Foot-and-mouth disease virus (FMDV)	eLife, 2015	Shihezi University
Porcine circovirus type 2 (PCV2)	Chinese Journal of Veterinary Science, 2017	South China Agricultural University





# Thanks

