

# CRISPR Crops— Plant Genome Editing Made Easy

 @KamounLab

 <http://KamounLab.net>



# Today's outline

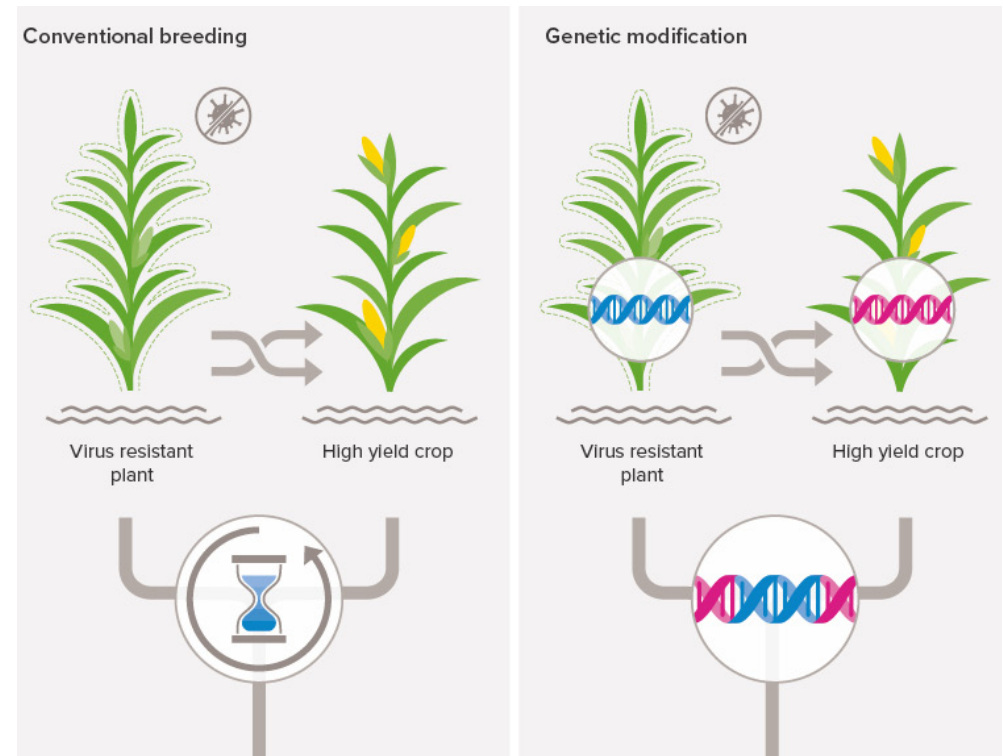
- ➔ What is CRISPR?
- ➔ Applications of CRISPR gene editing
- ➔ CRISPR and conservation biology





# Genetic improvement of plants

- ➔ Hybridization
- ➔ Introgression
- ➔ Mutagenesis
- ➔ Transgenics



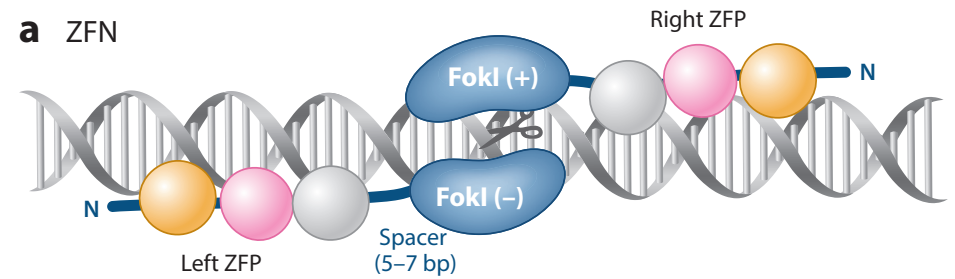
Gene editing – the ultimate in precision is to replace one of a few letters in the genome

# Reverse genetics in plants *circa* ~2012

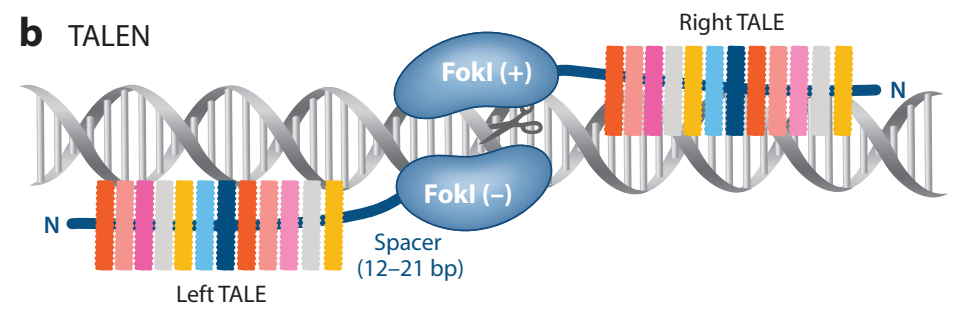
- ➔ Genome sequencing of mutagenized plants
- ➔ TILLING — targeted sequencing of mutant plants
- ➔ T-DNA or transposon mutagenesis
- ➔ Gene knock-down using RNAi (VIGS, artificial miRNA, hairpin constructs)
- ➔ Site-specific nucleases (e.g. ZFNs and TALENs) for gene editing

# Sequence-specific nucleases (SSNs)

- Zinc-Finger Nucleases

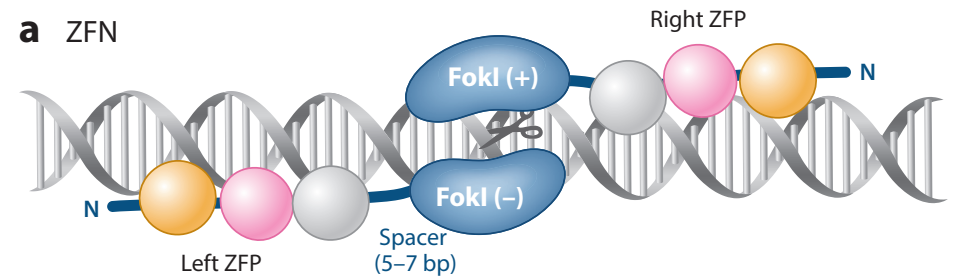


- TAL-Effector Nucleases

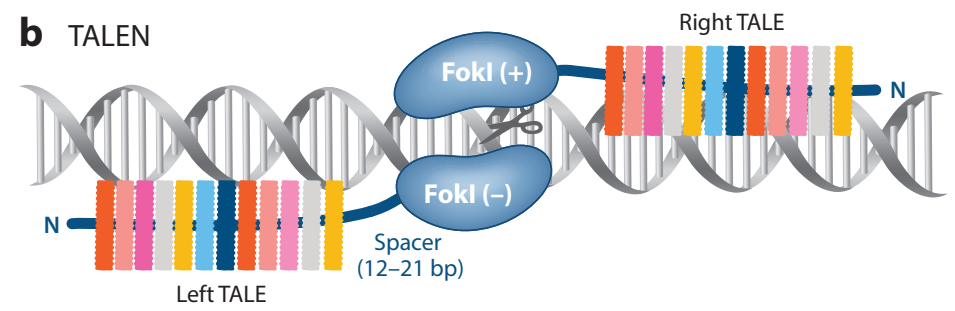


# Sequence-specific nucleases (SSNs)

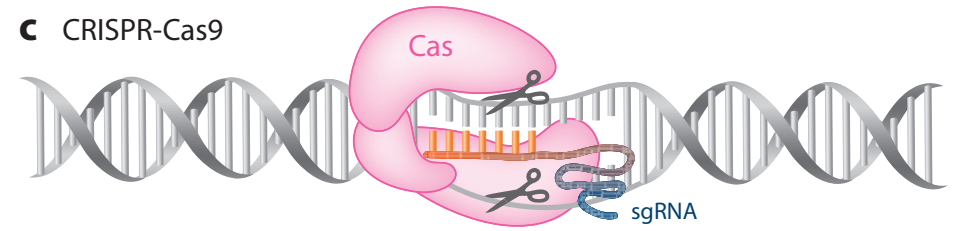
- Zinc-Finger Nucleases



- TAL-Effector Nucleases



- CRISPR/Cas9





Opinion

# This Year's Nobel Prize in Chemistry Honors a Revolution

With Crispr, two scientists turned a curiosity of nature into an invention that will transform the human race.

The CRISPR Craze





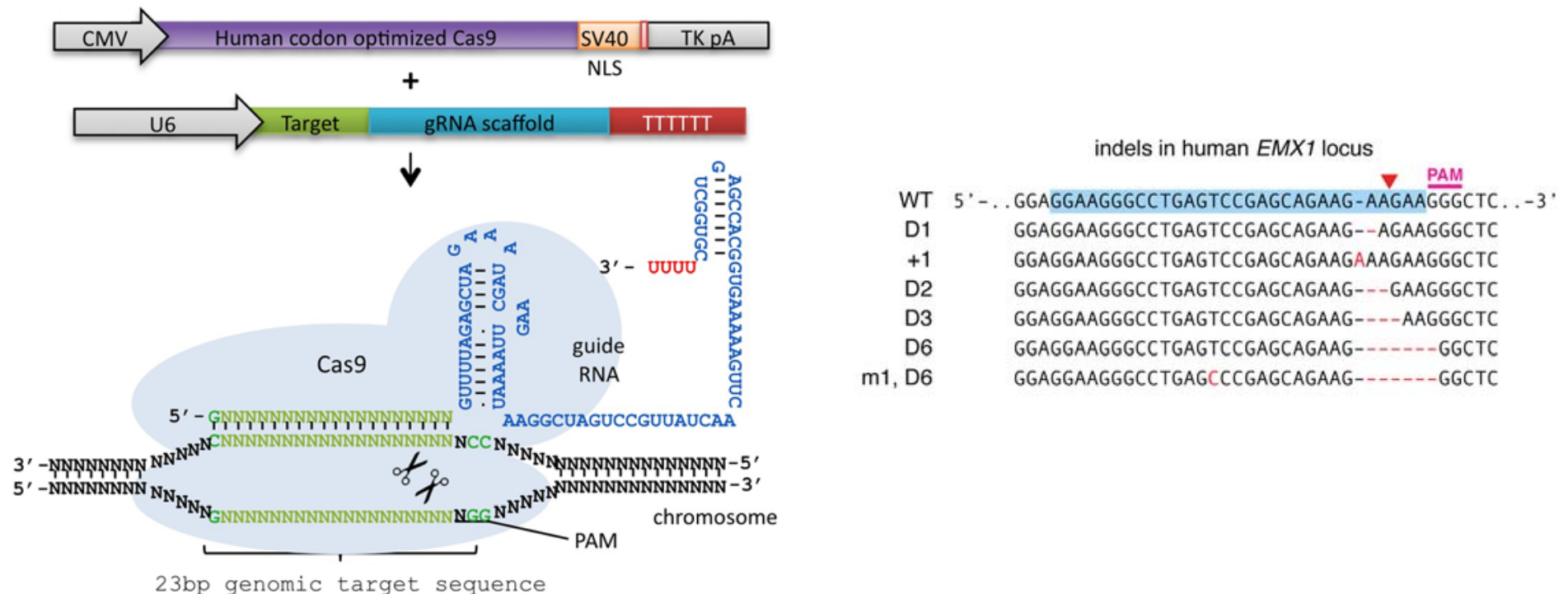
03 January 2013

## RNA-Guided Human Genome Engineering via Cas9

Prashant Mali,<sup>1,5</sup> Luhan Yang,<sup>1,3,5</sup> Kevin M. Esvelt,<sup>2</sup> John Aach,<sup>1</sup> Marc Guell,<sup>1</sup> James E. DiCarlo,<sup>4</sup> Julie E. Norville,<sup>1</sup> George M. Church<sup>1,2\*</sup>

## Multiplex Genome Engineering Using CRISPR/Cas Systems

Le Cong,<sup>1,2\*</sup> F. Ann Ran,<sup>1,4\*</sup> David Cox,<sup>1,3</sup> Shuailiang Lin,<sup>1,5</sup> Robert Barretto,<sup>6</sup> Naomi Habib,<sup>1</sup> Patrick D. Hsu,<sup>1,4</sup> Xuebing Wu,<sup>7</sup> Wenyan Jiang,<sup>8</sup> Luciano Marraffini,<sup>8</sup> Feng Zhang<sup>1†</sup>





**Multiplex and homologous recombination-mediated genome editing in *Arabidopsis* and *Nicotiana benthamiana* using guide RNA and Cas9**

Jian-Feng Li<sup>1,2</sup>, Julie E Norville<sup>2,3</sup>, John Aach<sup>2,3</sup>, Matthew McCormack<sup>1,2</sup>, Dandan Zhang<sup>1,2</sup>, Jenifer Bush<sup>1,2</sup>, George M Church<sup>2,3</sup> & Jen Sheen<sup>1,2</sup>

**Open**  
LETTER TO THE EDITOR

Cell Research (2013) 1:4.  
© 2013 IBCB, SIBS, CAS All rights reserved 1001-0602/13  
www.nature.com/cr



**Efficient genome editing in plants using a CRISPR/Cas system**

Cell Research advance online publication 20 August 2013; doi:10.1038/cr.2013.114

Zhengyan Feng<sup>1, 2, 3, \*</sup>, Botao Zhang<sup>1, \*</sup>, Wona Ding<sup>4</sup>, Xiaod Fengqi Yanfei

**Open**  
LETTER TO

**Target**  
Cell Research

Jin Mi  
Qingp  
Jianm

# The CRISPR Craze

## infects plant scientists

Nucleic Acids Research, 2013, 1–12  
doi:10.1093/nar/gkt1780

**Demonstration of CRISPR/Cas9/sgRNA-mediated targeted gene modification in *Arabidopsis*, tobacco, sorghum and rice**

Wenzhi Jiang<sup>1</sup>, Huanbin Zhou<sup>2</sup>, Honghao Bi<sup>2</sup>, Michael Fromm<sup>3</sup>, Bing Yang<sup>2</sup> and Donald P. Weeks<sup>1,\*</sup>

**Targeted genome modification of crop plants using a CRISPR-Cas system**

Qiwei Shan<sup>1,4</sup>, Yanpeng Wang<sup>1,4</sup>, Jun Li<sup>1,4</sup>, Yi Zhang<sup>1</sup>, Kunling Chen<sup>1</sup>, Zhen Liang<sup>1</sup>, Kang Zhang<sup>1</sup>, Jinxing Liu<sup>1</sup>, Jianzhong Jeff Xi<sup>2</sup>, Jin-Long Qiu<sup>3</sup> & Caixia Gao<sup>1</sup>

**Targeted mutagenesis in the model plant *Nicotiana benthamiana* using Cas9 RNA-guided endonuclease**

Vladimir Nekrasov<sup>1</sup>, Brian Staskawicz<sup>2</sup>, Detlef Weigel<sup>3</sup>, Jonathan D G Jones<sup>1,4</sup> & Sophien Kamoun<sup>1,4</sup>

SCIENCE VOL 341 23 AUGUST 2013

is System  
013

Kabin Xie and Yinong Yang\*

G3: Genes|Genomes|Genetics Early Online, published on October 11, 2013

RNA guided genome editing for target gene mutations in wheat

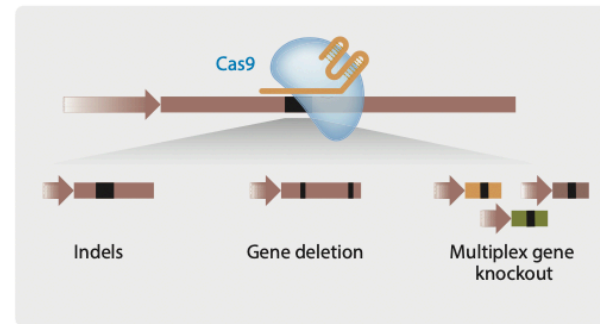
Santosh Kumar Upadhyay, Jitesh Kumar, Anshu Alok, RakeshTuli

# CRISPR/Cas Genome Editing and Precision Plant Breeding in Agriculture

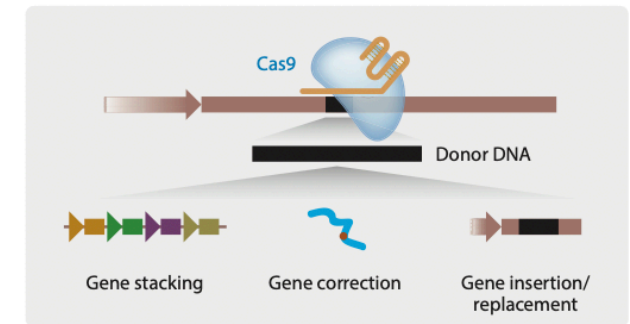
## What can be done with CRISPR?

Kunling Chen,<sup>1,\*</sup> Yanpeng Wang,<sup>1,\*</sup> Rui Zhang,<sup>1</sup> Huawei Zhang,<sup>1</sup> and Caixia Gao<sup>1,2</sup>

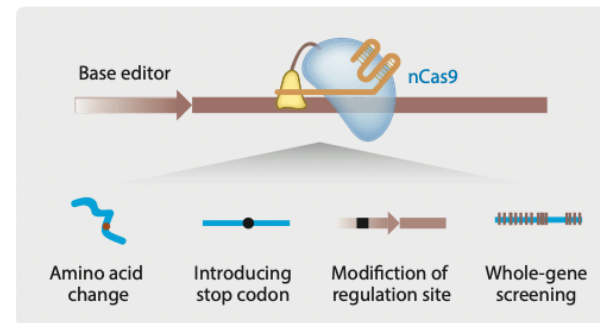
**a** Gene knockout



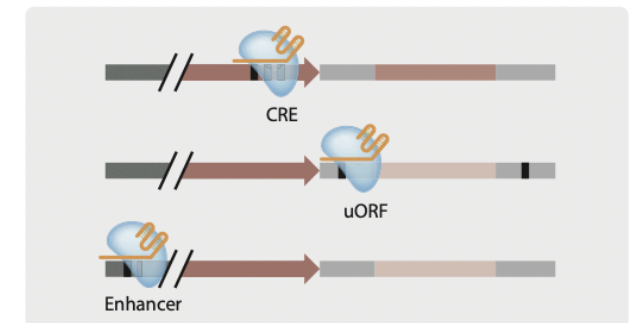
**b** Gene knock-in/replacement



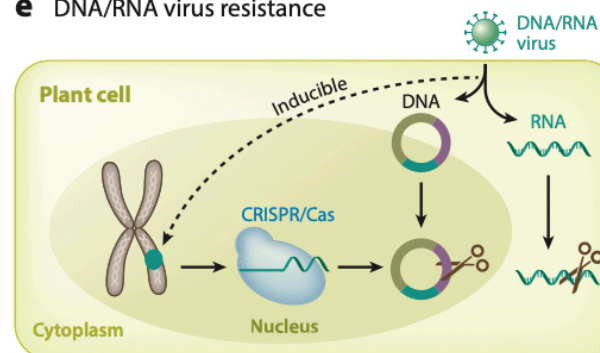
**c** Applications of base editing



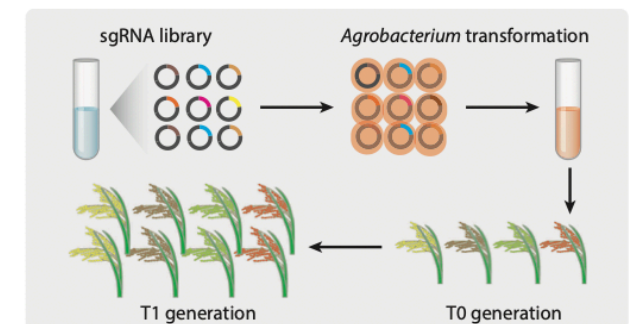
**d** Fine-tuning gene regulation



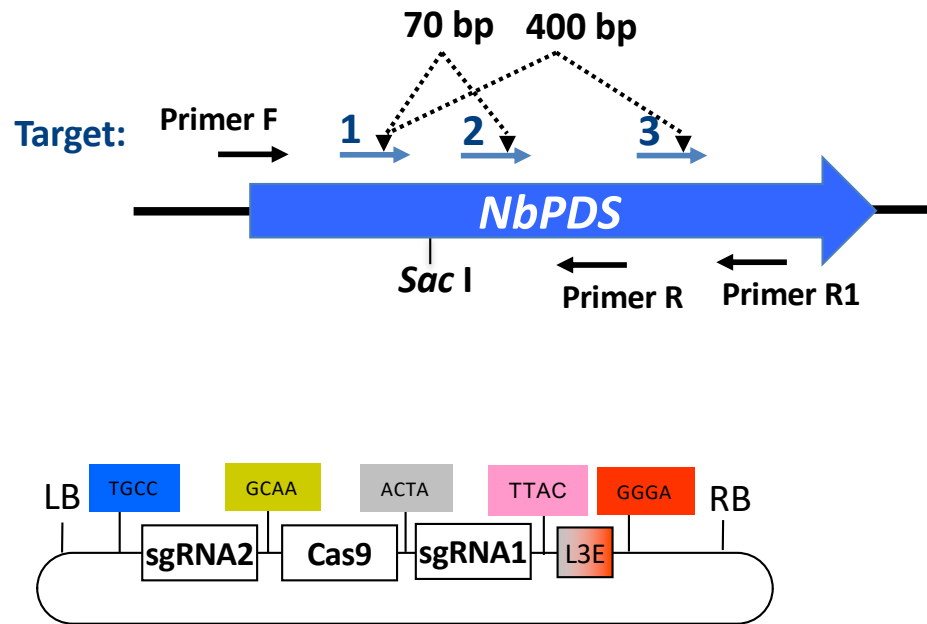
**e** DNA/RNA virus resistance



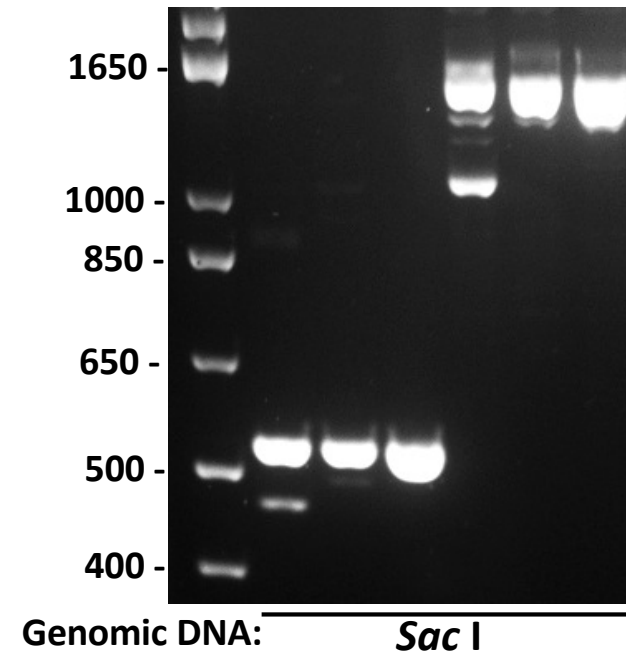
**f** High-throughput mutant library



# CRISPR/Cas9 enables making *small* deletions...

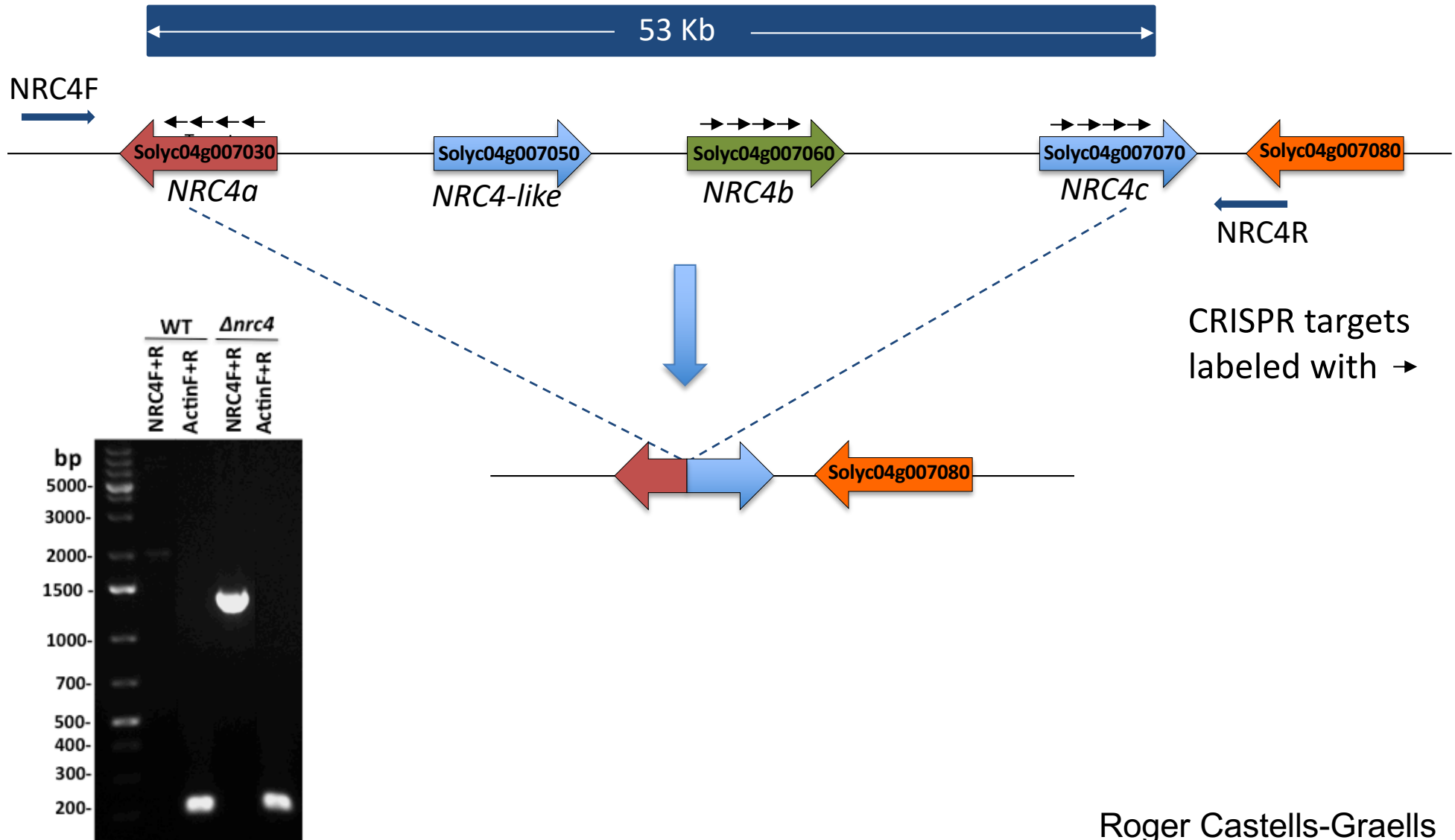


	1	2	3	4	5	6
sgRNA_PDS1	+	+	-	+	+	-
sgRNA_PDS2	+	-	-	-	-	-
sgRNA_PDS3	-	-	-	+	-	-
sgRNA_GFP	-	+	-	-	+	-
Cas9	+	+	+	+	+	+



	Target 1	PAM	Target 2	PAM
PDS	TTCTGCCGTTAATTTGAGAGT	-CCAAGGTAATTCAGCTTATCTTTGGAGCTCGAGGTCTTCGTTGGGAACTGAAA	GTCAAGATGTTTGCTTGCAA	AAGGAATT
m1	TTCTGCCGTTAATTT	-----	-----	CAAAGGAATT -76
m2	TTCTGCCGTTAATTTGAGAGT	-----	-----	CAAAGGAATT -70
m3	TTCTGCCGTTAATTTGAGAGTT	-----	-----	CAAAGGAATT -70 +1 T

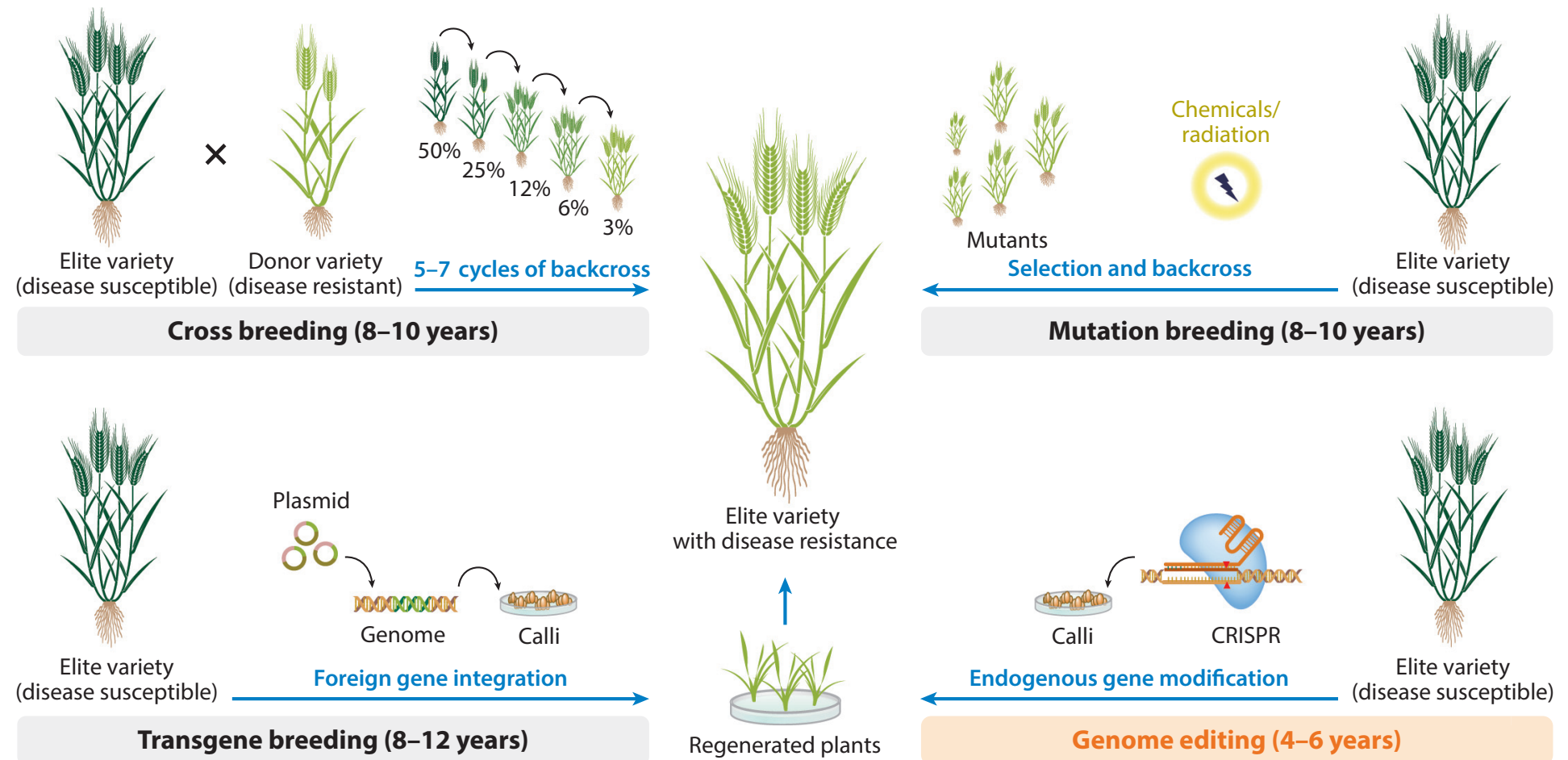
# ...and *B/G* deletions



Roger Castells-Graells  
Vladimir Nekrasov

# CRISPR/Cas Genome Editing and Precision Plant Breeding in Agriculture

Kunling Chen,<sup>1,\*</sup> Yanpeng Wang,<sup>1,\*</sup> Rui Zhang,<sup>1</sup>  
Huawei Zhang,<sup>1</sup> and Caixia Gao<sup>1,2</sup>





# Crop plants with DNA deletions are not GMOs

by [Sophien Kamoun](#) and [Eric Ward](#)

16

MAY 2012

## Crop plants with DNA deletions are not GMOs

by [Sophien Kamoun](#) and [Eric Ward](#)

16

MAY 2012

- ➔ A regulatory not scientific question
- ➔ Two main regulatory frameworks for new crop varieties:
  - ➔ *process-based*
  - ➔ *product-based*





Frank Hochholdinger

@HochholdingerF

Following



The list of countries, which do not regulate targeted mutagenesis by genome editing under GMO legislation grows:

Japan 

USA 

Canada 

Argentina 

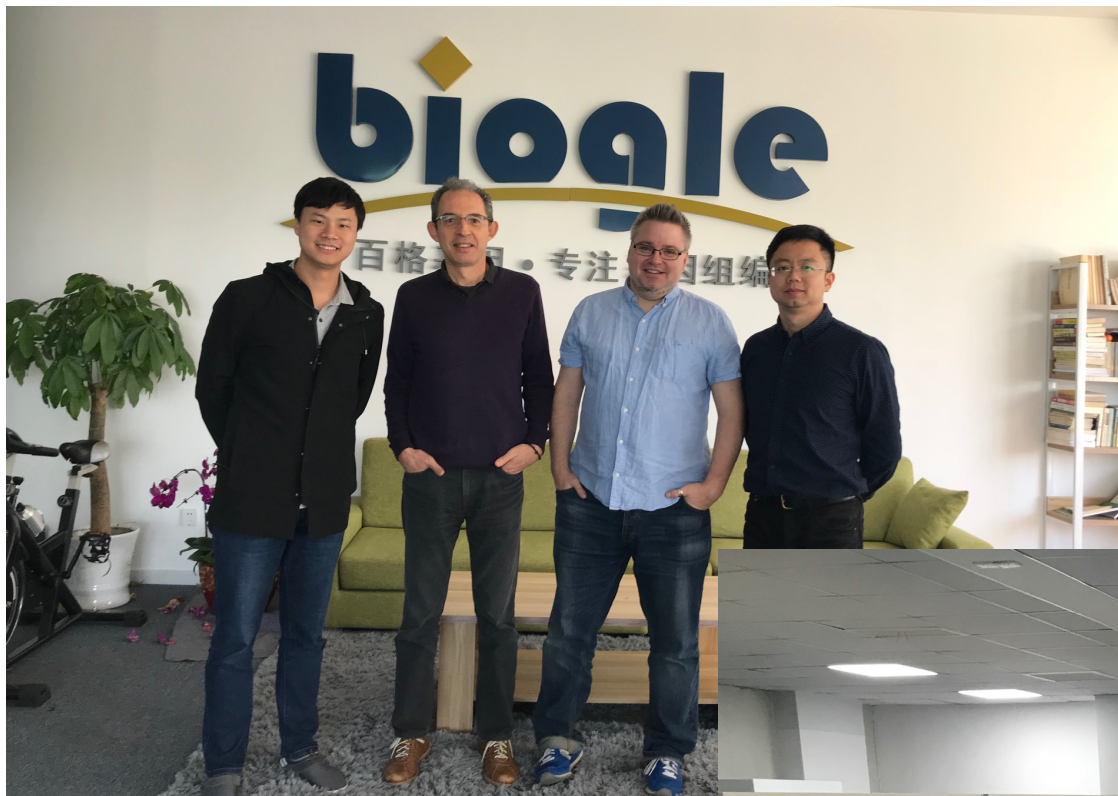
Brazil 

Chile 

Israel 

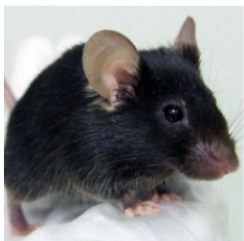
Australia  is likely next.

Only the EU  ignores scientific consensus.





# — 我们专注于基因组编辑 —



## 小鼠基因组编辑

- 单/多基因敲除, 长片段敲除, 条件性敲除
- DNA片段定点敲入、碱基替换
- 服务周期2~4个月
- 提供Founder小鼠或F1代小鼠



## 细胞系基因组编辑

- 常用细胞系 (293T、Hela), 小鼠胚胎干细胞以及iPS干细胞
- 单/多基因敲除, 长片段敲除, DNA片段定点敲入、碱基替换
- 服务周期1~3个月



## 水稻基因组编辑

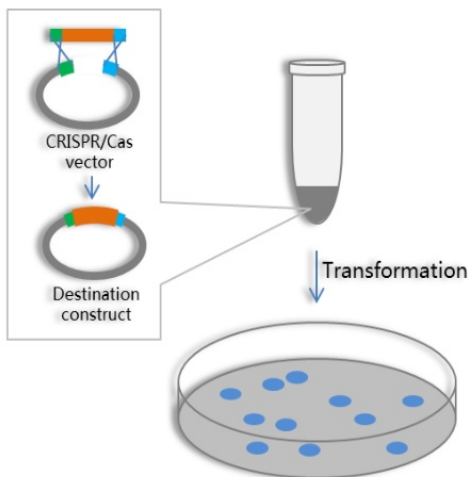
- 粳稻 (如日本晴)、籼稻 (如93-11)
- 单/多基因敲除, 长片段敲除
- 服务周期2~4个月
- 提供至少3株成功突变的T0幼苗



## 大豆基因组编辑

- 常用品种Williams82、Jack
- 单基因敲除
- 服务周期3~9个月
- 提供至少1株成功突变的T0幼苗或3株T1幼苗

## 百格CRISPR/Cas载体构建试剂盒



### 一步载体构建, 直接遗传转化, 数百次的成功实验验证

- 简便 无需酶切, 一步载体构建, 直接用于遗传转化
- 快速 20°C反应30~60分钟即可
- 高效 1000个以上的菌落数, 95%以上的阳性率, 提供您所需的克隆
- 可靠 数百次的植物遗传转化实验, 90%以上基因敲除成功率
- 广泛 涵盖拟南芥、油菜、水稻、小麦和大豆等

# Genome-wide Targeted Mutagenesis in Rice Using the CRISPR/Cas9 System

Yuming Lu<sup>1,3</sup>, Xiao Ye<sup>1</sup>, Renming Guo<sup>1</sup>,  
Jing Huang<sup>1</sup>, Wei Wang<sup>2</sup>, Jiuyou Tang<sup>2</sup>,  
Longtao Tan<sup>4</sup>, Jian-kang Zhu<sup>3</sup>, Chengcai Chu<sup>2</sup>  
and Yangwen Qian<sup>1,\*</sup>

<sup>1</sup>Biogle Genome Editing Center, Changzhou, Jiangsu Province 213125, China

<sup>2</sup>State Key Laboratory of Plant Genomics, National Center for Plant Gene Research (Beijing), Institute of Genetics and Developmental Biology, Chinese Academy of Sciences, Beijing 100101, China

<sup>3</sup>Shanghai Center for Plant Stress Biology and Center for Excellence in Molecular Plant Sciences, Chinese Academy of Sciences, Shanghai 201602, China

<sup>4</sup>Key Laboratory of Agro-ecological Processes in Subtropical Region, Institute of Subtropical Agriculture, Chinese Academy of Sciences, Changsha 410125, China

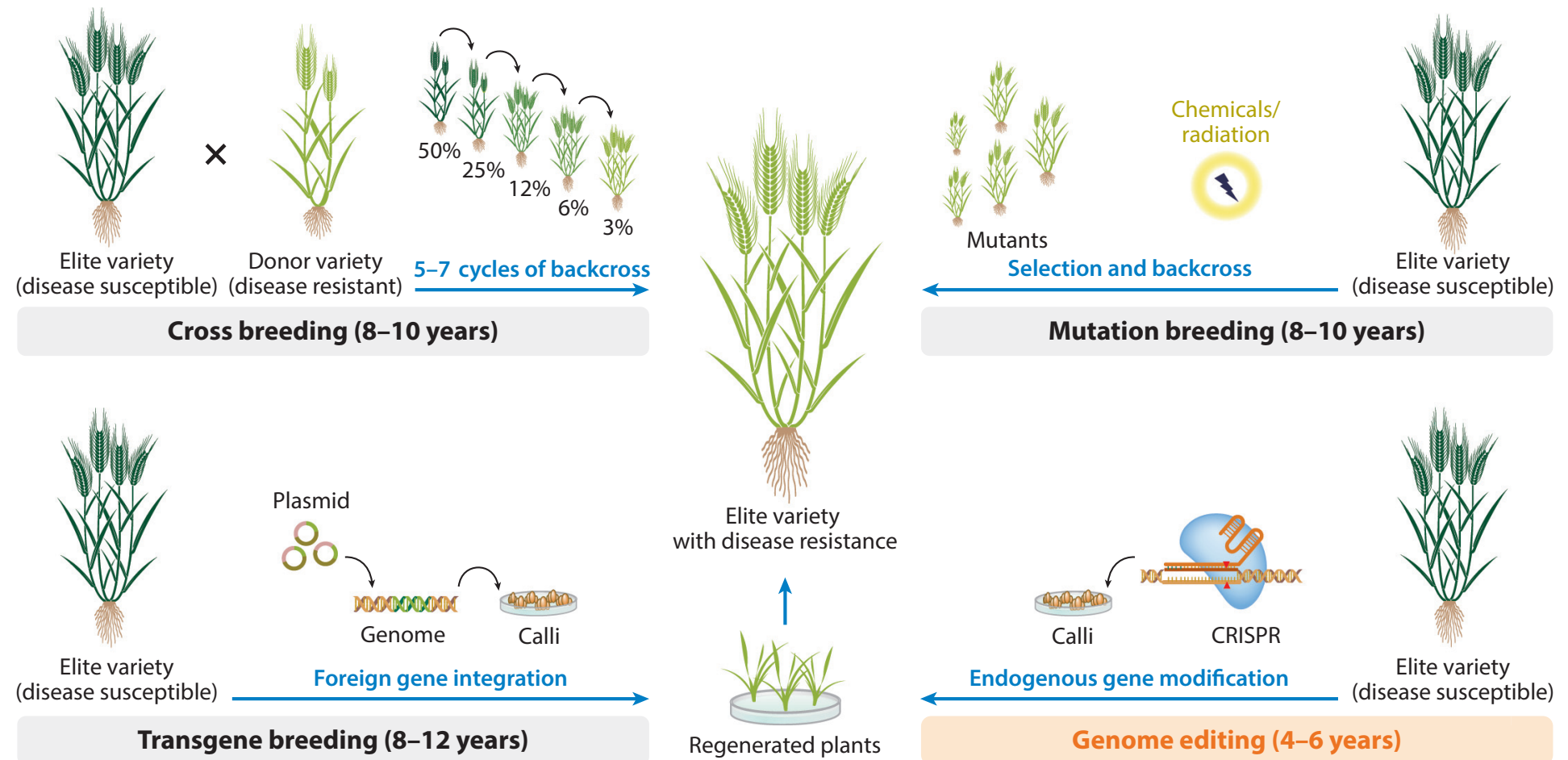
\*Correspondence: Yangwen Qian ([qyw@biogle.cn](mailto:qyw@biogle.cn))

<http://dx.doi.org/10.1016/j.molp.2017.06.007>



# CRISPR/Cas Genome Editing and Precision Plant Breeding in Agriculture

Kunling Chen,<sup>1,\*</sup> Yanpeng Wang,<sup>1,\*</sup> Rui Zhang,<sup>1</sup>  
Huawei Zhang,<sup>1</sup> and Caixia Gao<sup>1,2</sup>



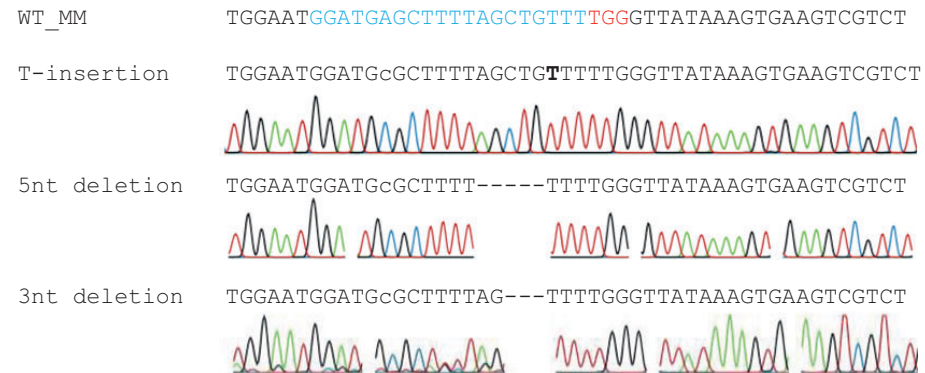


## Using CRISPR/Cas9 genome editing in tomato to create a gibberellin-responsive dominant dwarf DELLA allele

Laurence Tomlinson<sup>1</sup> , Ying Yang<sup>1</sup>, Ryan Emenecker<sup>2</sup>, Matthew Smoker<sup>1</sup>, Jodie Taylor<sup>1</sup>, Sara Perkins<sup>1</sup>, Justine Smith<sup>1</sup>, Dan MacLean<sup>1</sup>, Neil E. Olszewski<sup>2</sup> and Jonathan D. G. Jones<sup>1,\*</sup>

<sup>1</sup>The Sainsbury Laboratory, Norwich Research Park, Norwich, UK

<sup>2</sup>Department of Plant and Microbial Biology, University of Minnesota, St. Paul, MN, USA



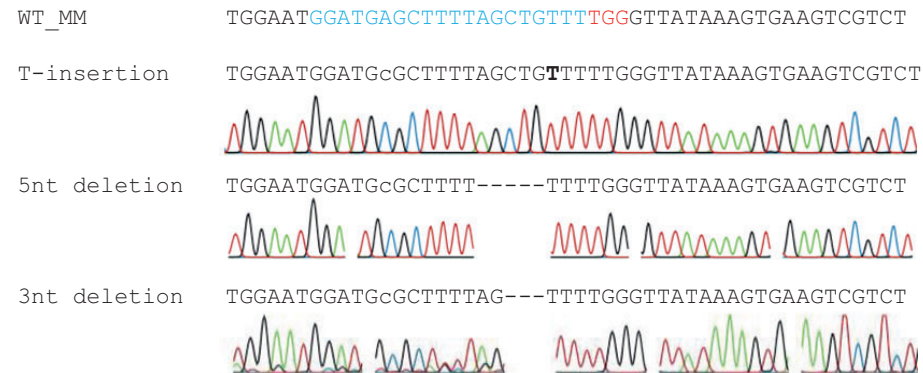
Allelic series!

# Using CRISPR/Cas9 genome editing in tomato to create a gibberellin-responsive dominant dwarf DELLA allele

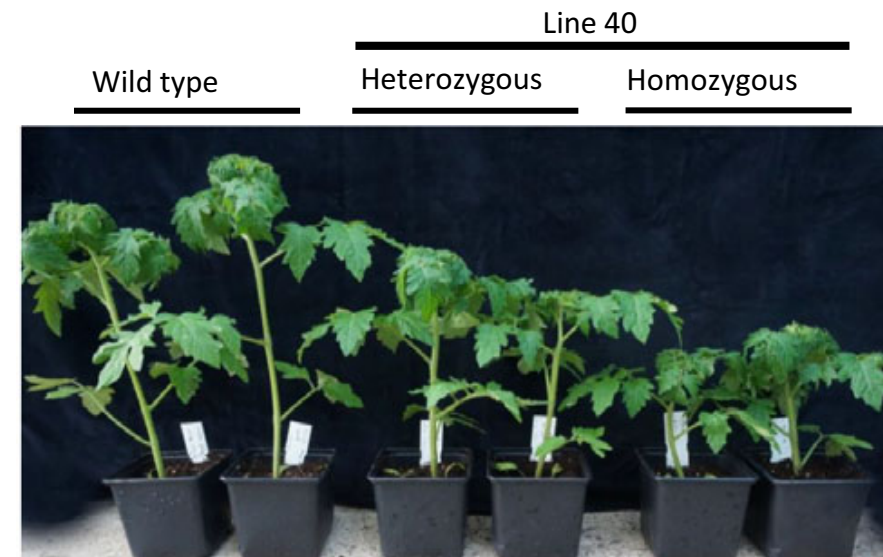
Laurence Tomlinson<sup>1</sup> , Ying Yang<sup>1</sup>, Ryan Emenecker<sup>2</sup>, Matthew Smoker<sup>1</sup>, Jodie Taylor<sup>1</sup>, Sara Perkins<sup>1</sup>, Justine Smith<sup>1</sup>, Dan MacLean<sup>1</sup>, Neil E. Olszewski<sup>2</sup> and Jonathan D. G. Jones<sup>1,\*</sup>

<sup>1</sup>The Sainsbury Laboratory, Norwich Research Park, Norwich, UK

<sup>2</sup>Department of Plant and Microbial Biology, University of Minnesota, St. Paul, MN, USA



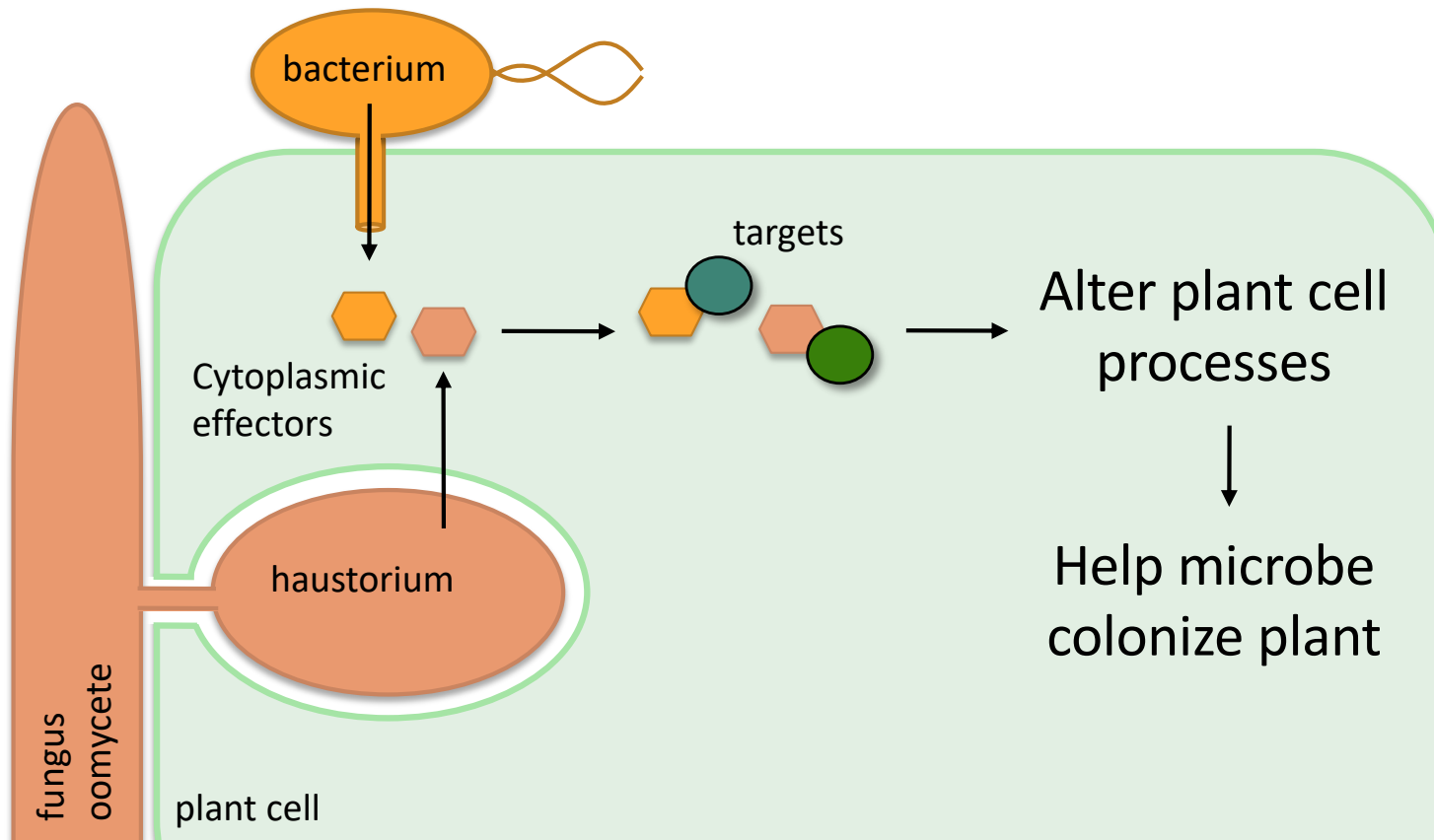
Allelic series!





The plant targets of pathogens *facilitate* infection

Susceptibility S genes!



# CRISPR crops — removing genes for fungal resistance

	Target 1	PAM	PAM	Target 2	
	ACATAGTAAAA <b>GGTGTACCTGTGGTGGAGAC</b>	<u>TGGTGACC</u>	ATCTTTTCTGGTTTAATCGCCCTGCCCTTGTCC	<b>ATTCTTGATTAAC</b>	<b>TTTGTACTCTTTTCAGG</b>
Plant 1	ACATAGTAAAA <b>GGTGTACCTGTGGTGGAGAC</b>	<u>TGGTGACC</u>	ATCTTTTCTGGTTTAATCGCCCTGCCCTTGTCC	<b>ATTCTTGATTAAC</b>	<b>TTTGTACTCTTTTCAGG</b>
Plant 2	ACATAGTAAAA <b>GGTGTACCTGTGGTGGAGAC</b>	<u>TGGTGACC</u>	ATCTTTTCTGGTTTAATCGCCCTGCCCTTGTCC	<b>ATTCTTGATTAAC</b>	<b>TTTGTACTCTTTTCAGG</b>
Plant 8	ACATAGTAAAA <b>GGTGTACCTGTGGTGGAGAC</b>	<u>TGGTGACC</u>	ATCTTTTCTGGTTTAATCGCCCTGCCCTTGTCC	<b>ATTCTTGATTAAC</b>	<b>TTTGTACTCTTTTCAGG</b>
Plant 10	ACATAGTAAAA <b>GGTGTACCTGTGGTGGAGAC</b>	<u>TGGTGACC</u>	ATCTTTTCTGGTTTAATCGCCCTGCCCTTGTCC	<b>ATTCTTGATTAAC</b>	<b>TTTGTACTCTTTTCAGG</b>

susceptible plant

resistant plant



# New Tomato Ideal for Urban Gardens and Even Outer Space Created Through Genetic Editing

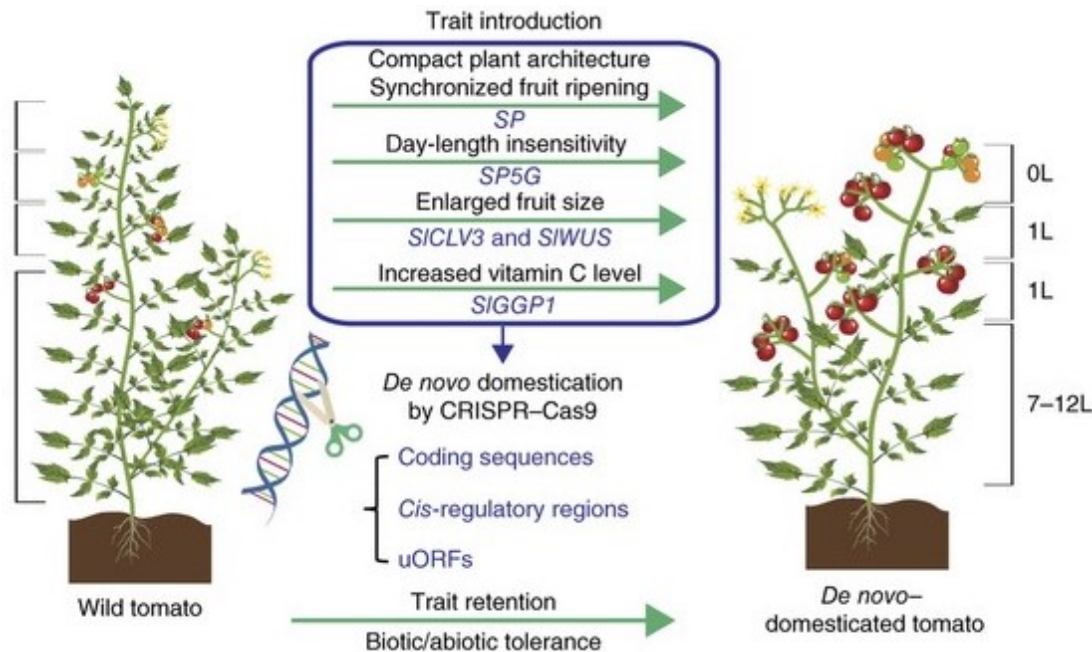
By COLD SPRING HARBOR LABORATORY DECEMBER 31, 2019





# *De novo* domestication of wild tomato using genome editing

Agustin Zsögön<sup>1,7</sup> , Tomáš Čermák<sup>2,6,7</sup>, Emmanuel Rezende Naves<sup>1</sup>, Marcela Morato Notini<sup>3</sup>, Kai H Edel<sup>4</sup>, Stefan Weigl<sup>4</sup>, Luciano Freschi<sup>5</sup>, Daniel F Voytas<sup>2</sup>, Jörg Kudla<sup>4</sup>  & Lázaro Eustáquio Pereira Peres<sup>3</sup> 



PLANT SCIENCES · 05 FEBRUARY 2021

## Insta-crop: CRISPR enables high-speed plant domestication

A lanky species of wild rice turns compact and docile in a jiffy.

# CRISPR and conservation biology

➔ Crop conservation biology needs **a bigger toolbox** to meet unprecedented challenges

➔ *climate change*

➔ *habitat loss*

➔ *invasive pathogens and pests*

➔ *etc.*

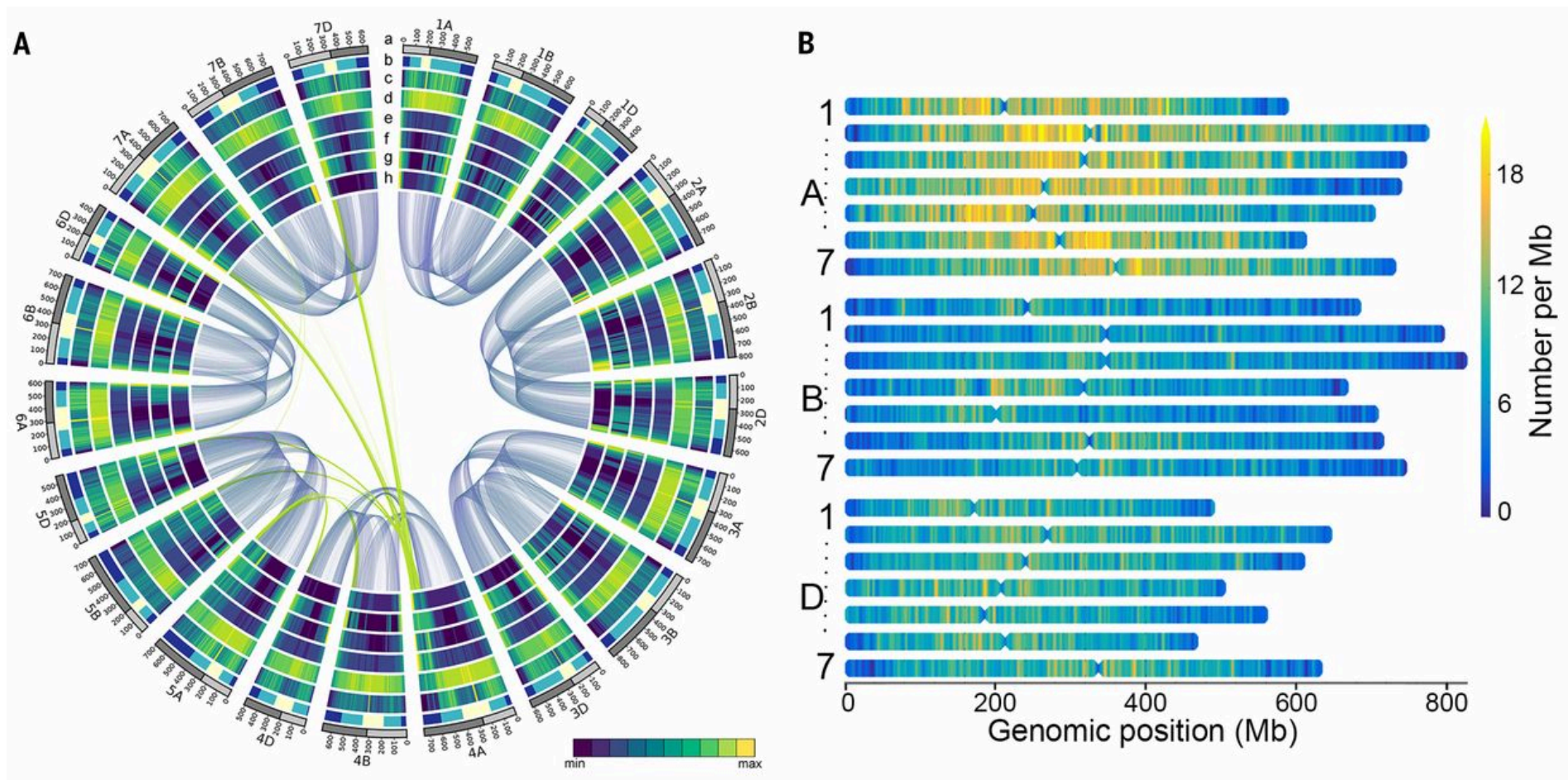
# CRISPR and conservation biology

- ➔ **Conservation genomics**—before we conserve, we need to know what's out there
- ➔ **Facilitated adaptation**—introduce adaptive traits for conservation of endangered crops
- ➔ **Crop de-extinction?**—CRISPR can deliver it

Crop genomics—*just getting started!*

2018

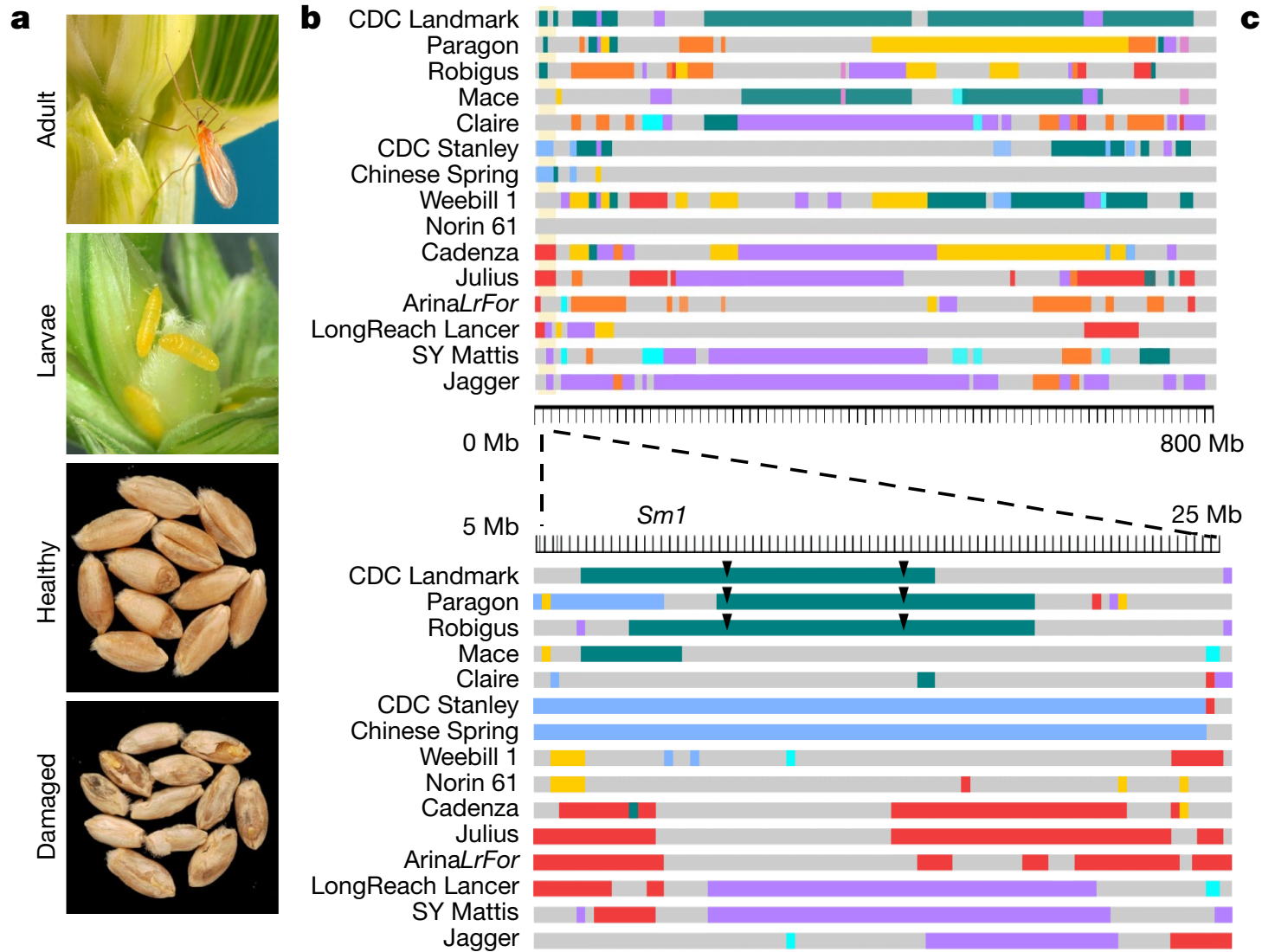
# SLICING THE wheat genome



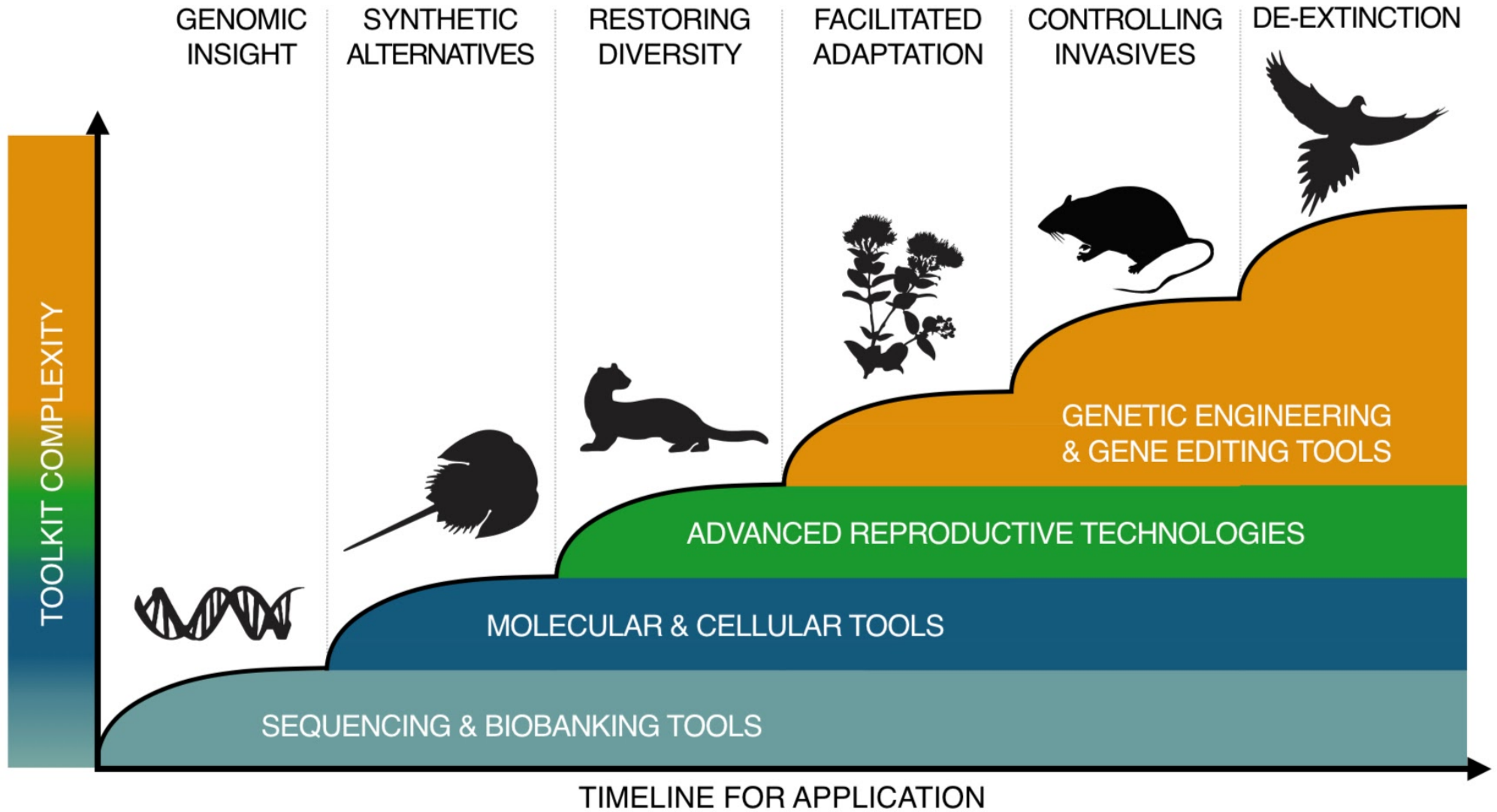


# Multiple wheat genomes reveal global variation in modern breeding

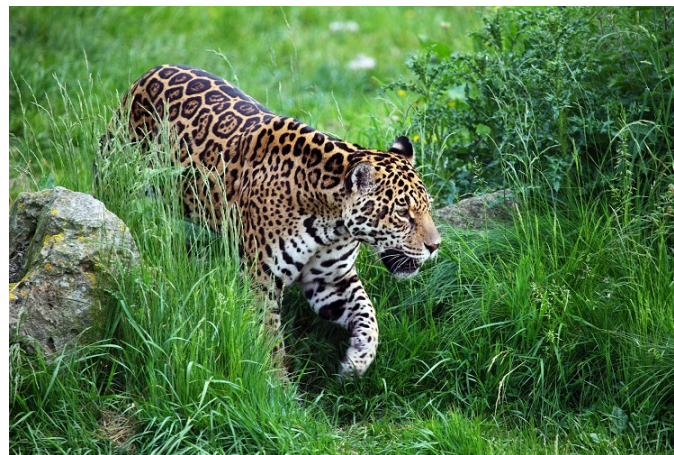
2020



# REVIVE & RESTORE IS BRINGING BIOTECHNOLOGIES TO CONSERVATION









# Plant genome editing is in its infancy...

Kamoun Lab @ TSL



**Sophien Kamoun**

@KamounLab

"I didn't come here to tell you how this is going to end. I came here to tell you how it's going to begin." Neo 'The Matrix'

# Acknowledgements



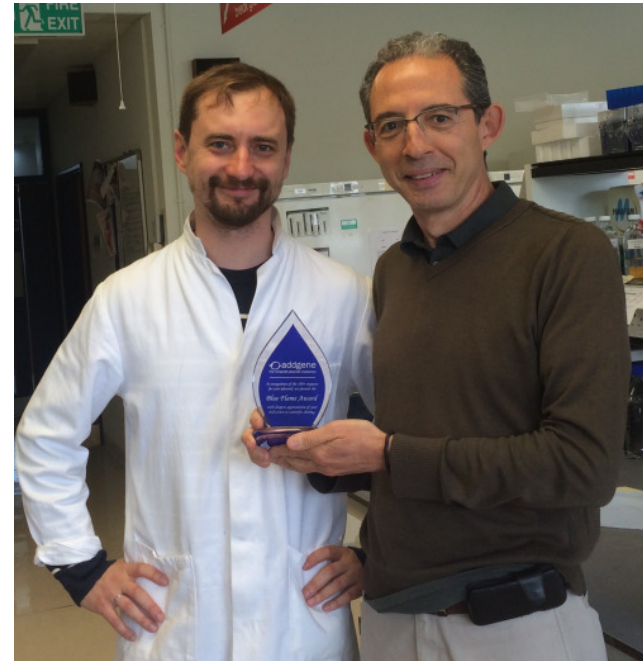
Khaoula Belhaj



Angela Chaparro Garcia



Joe Win



Vladimir Nekrasov



Diana Gomez



Adeline Harant




Thorsten Langner

# Great Resource: Plant Genome Editing Database /via Greg Martin @BTIscience @NSF

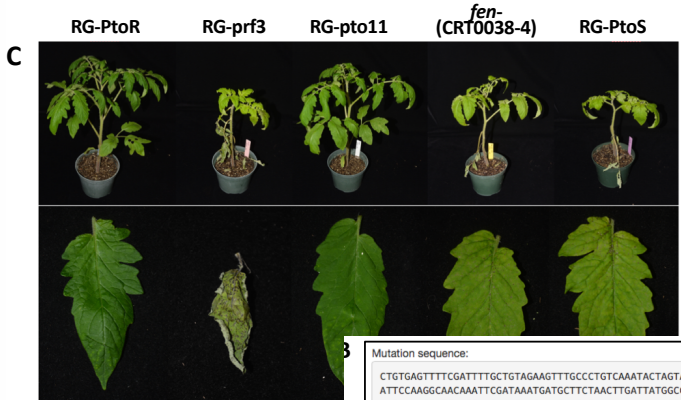
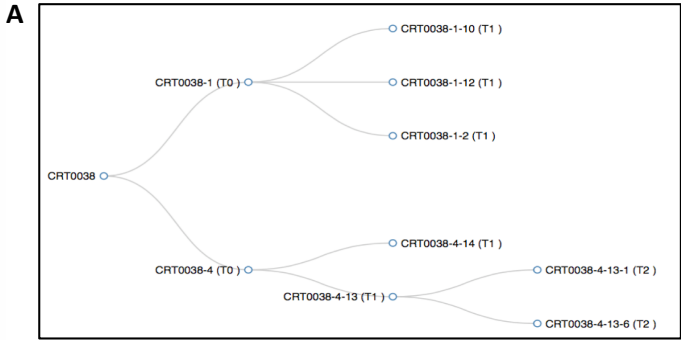


Home



Resource: Plant Genome Editing Database (2018) | Plants and Microbes scoop.it

29/10/2018, 19:13



Mutation sequence:

```
CTGTGAGTTTCGATTTGCTGTAGAAGTTTGCCCTGTCAAATAGTAGATTTCATTATCTATTAAATGGGAAGCAAGT
ATCCAAGGCCAACAAATTCGATAAAATGATGCTCTCAACTGATTTATGGCGTTCCTTTGAAAATTATCGAGTTCCTTTT
TAGATTTGGAGGAAGCAACTAACCACTTTGATGACATTTTTCATTGGAGAGGGGGATTTGGGAAGGTTTACAGGGGT
GTTTTCGTGATGGAACAAAGTCCCTCGAAAAGCATAAACCTGAGTCTCCACAAGGATTGGGAAGTTGGAAGACAGA
AATTGAGATTCCTCATTTTTCAGCCATCCGATCTGGTTTCATTGATGAGGATCTGTGATGAAAAGAAATGAGATGATTC
TAATTTAGACTACATGGAGAATGGGAACCTCAA
```

Mutation alignment:

	596	646	696	746	796	846	896	946
Wildtype	TTGAGGTTCCCATCTCCATGTAGTCATAAAATAGAATCATCTCATTTCCT	TTTCATCACAGAATCCATCAATGAACCAGATGCCGATGGCTGCAAAATG	AGAGAACTCAATTTCTGTTTGAACCTCTCAATACCTTGTGAGGACTCA	CGTTTATGCTTTTTCAGGGCGACCTTGTTCATCAGCAAACACCCCT	GTAACCTCCCAAATCCACCTCTCAATGAAAATTTGTATCAAAAGT	TGTTAGTGTCTCCTCAAATCTACAAAAGGAAGCTGATAAATTTGAAA	GGAATGCCATAA	GAATGCCATAA
MutationRev	TTGAGGTTCCCATCTCCATGTAGTCATAAAATAGAATCATCTCATTTCCT	TTTCATCACAGAATCCATCAATGAACCAGATGCCGATGGCTGCAAAATG	AGAGAACTCAATTTCTGTTTGAACCTCTCAATACCTTGTGAGGACTCA	GGTTTATGCTTTTTCAGGGCGACCTTGTTCATCAGCAAACACCCCT	GTAACCTCCCAAATCCACCTCTCAATGAAAATTTGTATCAAAAGT	TGTTAGTGTCTCCTCAAATCTACAAAAGGAAGCTGATAAATTTGAAA	GGAATGCCATAA	GAATGCCATAA
gRNA							TAACCTCAAGTTAGAAGCATC	
MutationRev							TAACCTCAAGTTAGAAGCATC	



