

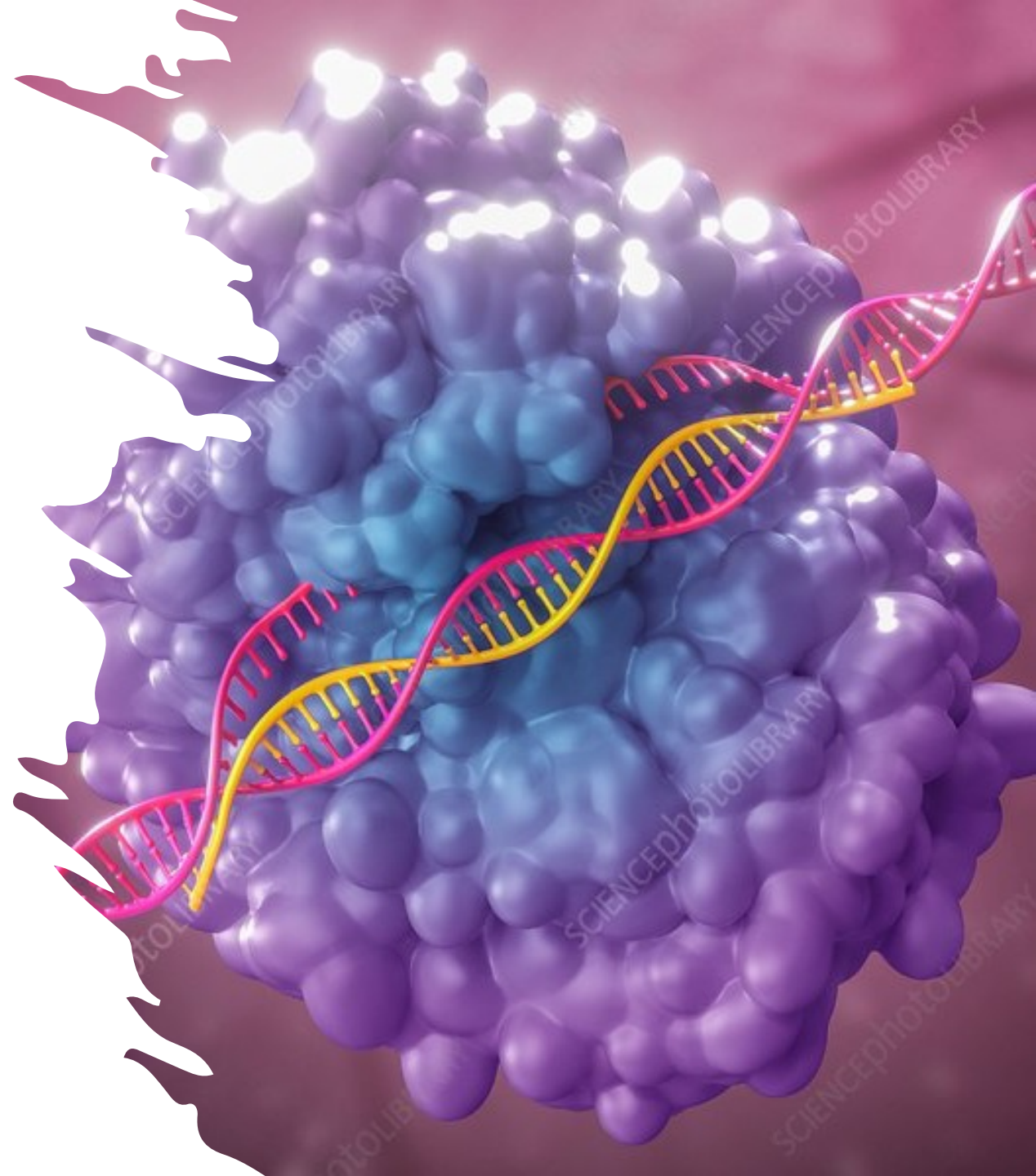


Presentation And Discussion Of The Paper:

Engineering herbicide-resistant watermelon variety through CRISPR/Cas9-mediated base-editing

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29th Jun, 2021



Outline

- The paper
- Aim of the study
- The CRISPR/cas9 system
- CRISPR/cas9 Base Editing
- CRISPR/cas9 delivery into plants
- Methods
- Results
- Discussion
- Conclusions



The paper


Plant Cell Reports

<https://doi.org/10.1007/s00299-018-2299-0>

FOCUS ARTICLE



Engineering herbicide-resistant watermelon variety through CRISPR/Cas9-mediated base-editing

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Keywords Base-editing · Herbicide-resistant watermelon · Transgene-free



Aim of the study

Utilizing CRISPR/ Cas9 base editing to produce herbicide-resistant-watermelon (broadleaved weed)

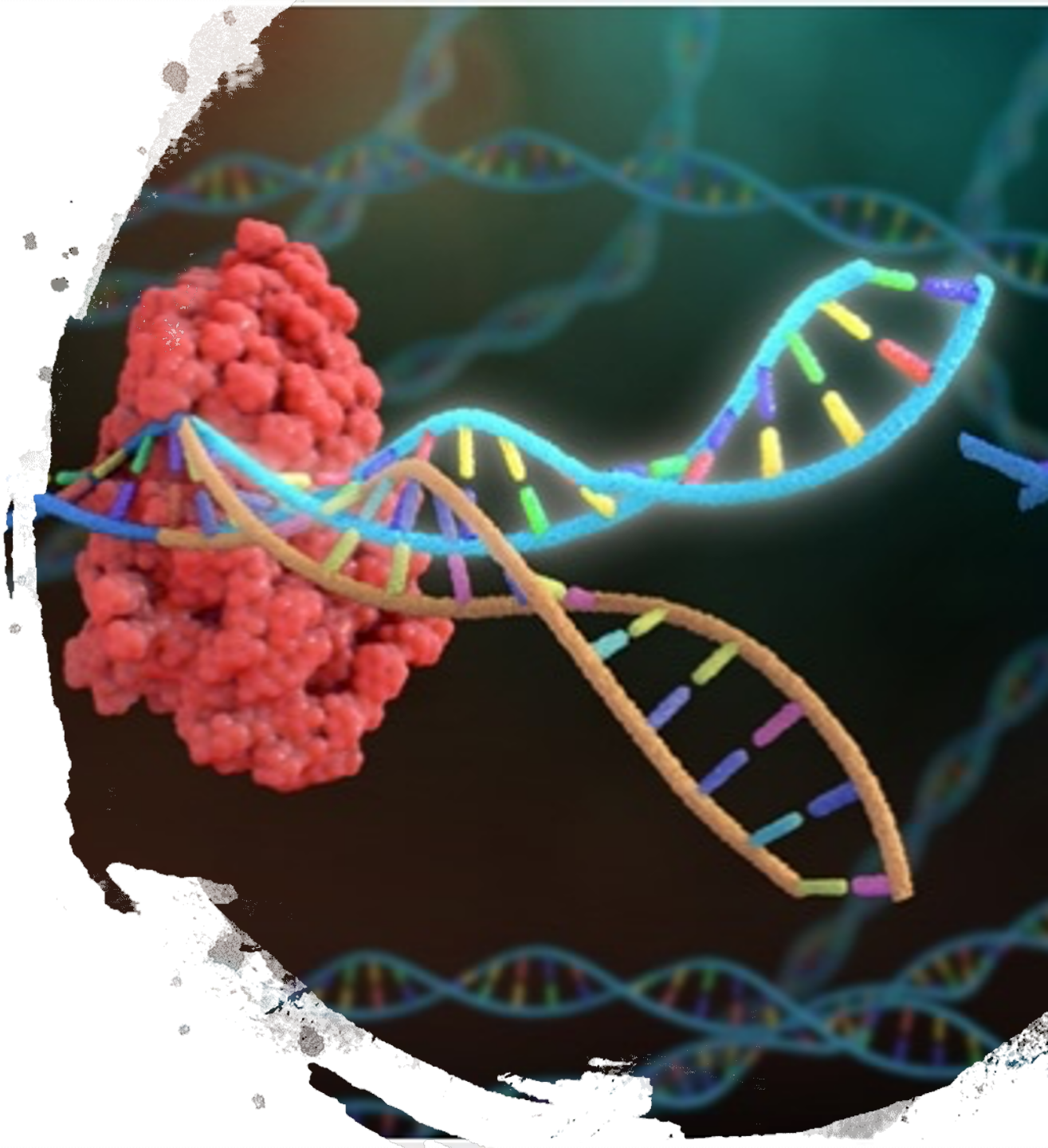
Broadleaved weed



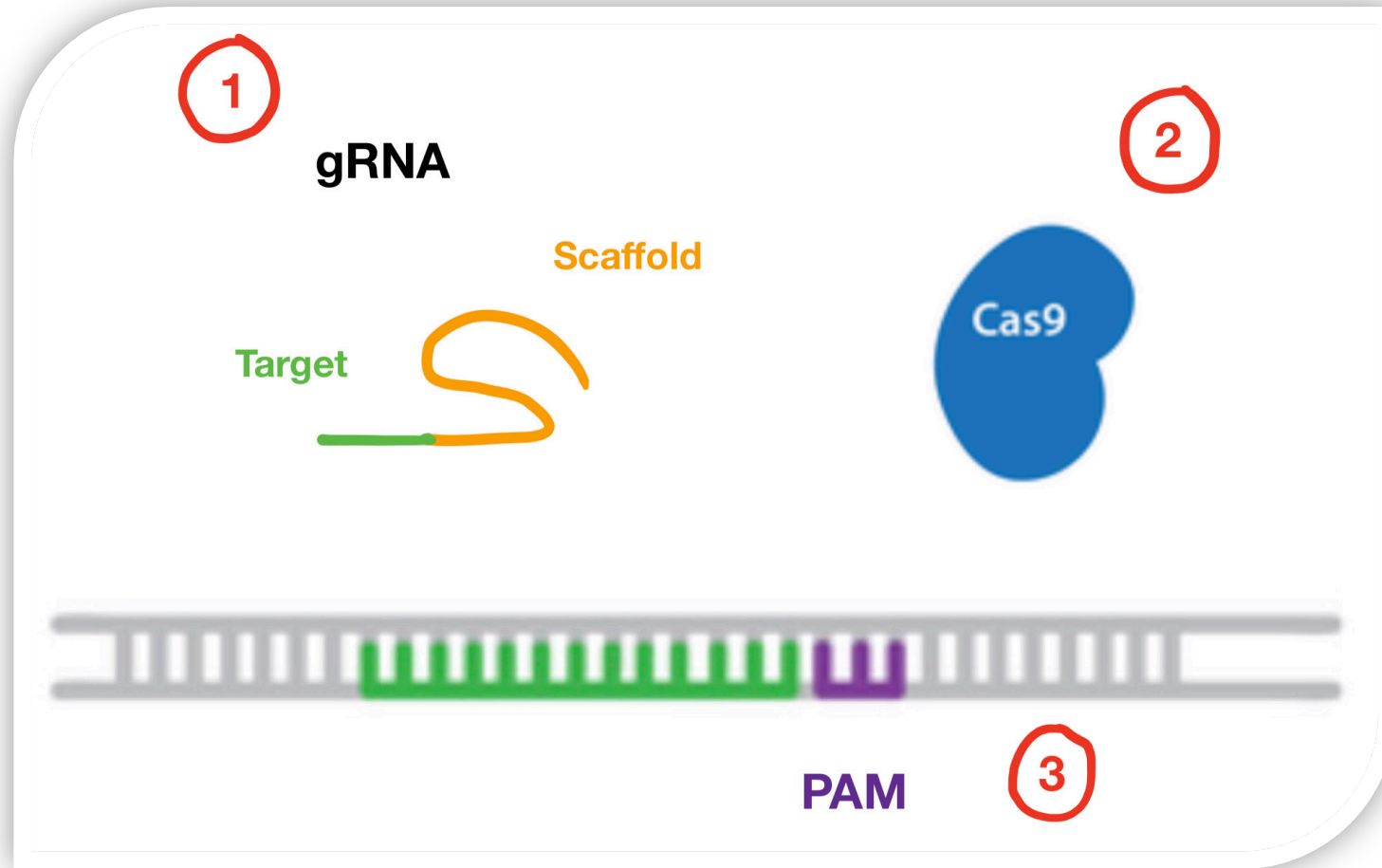
Previous studies



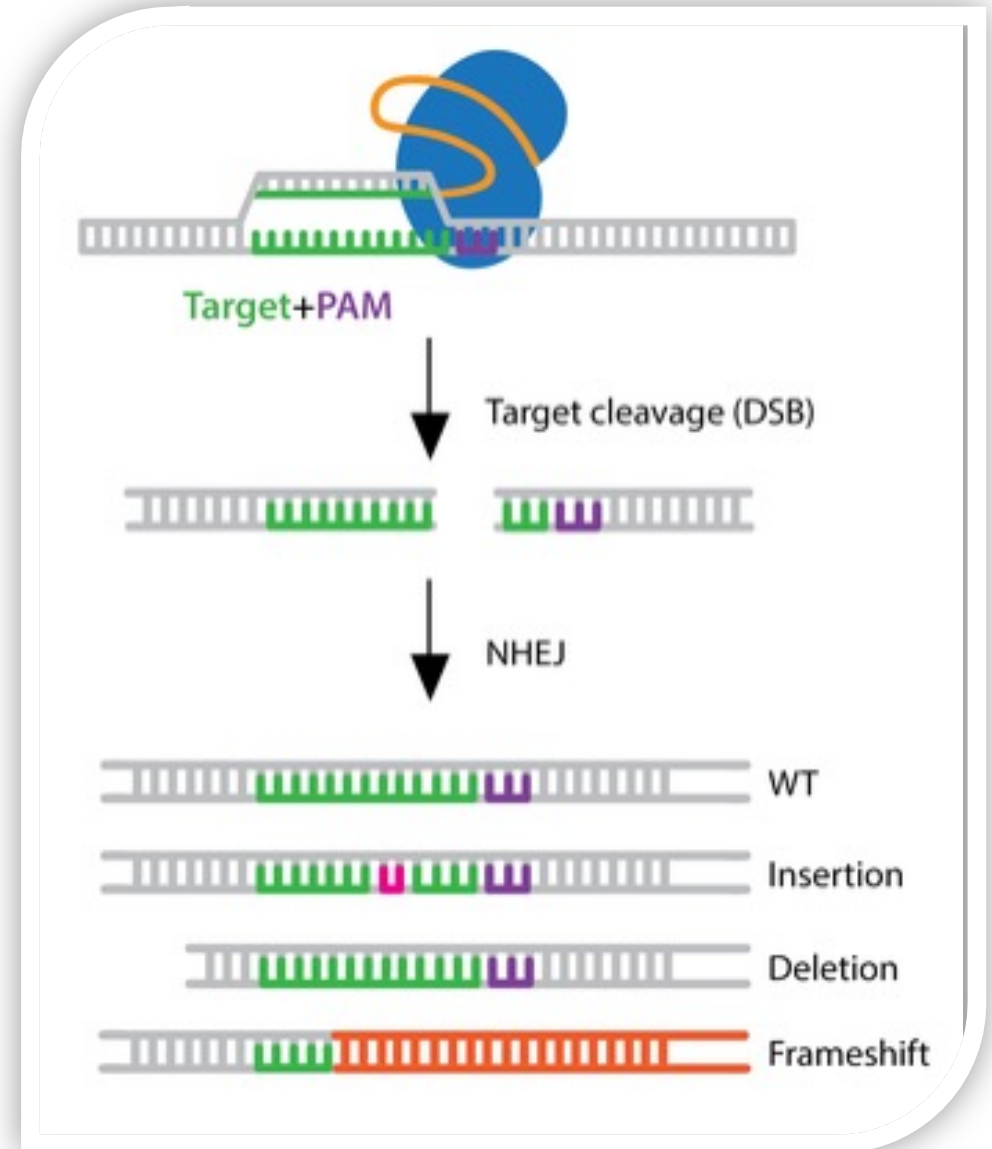
CRISPR/cas9 system



CRISPR/cas9 System Components



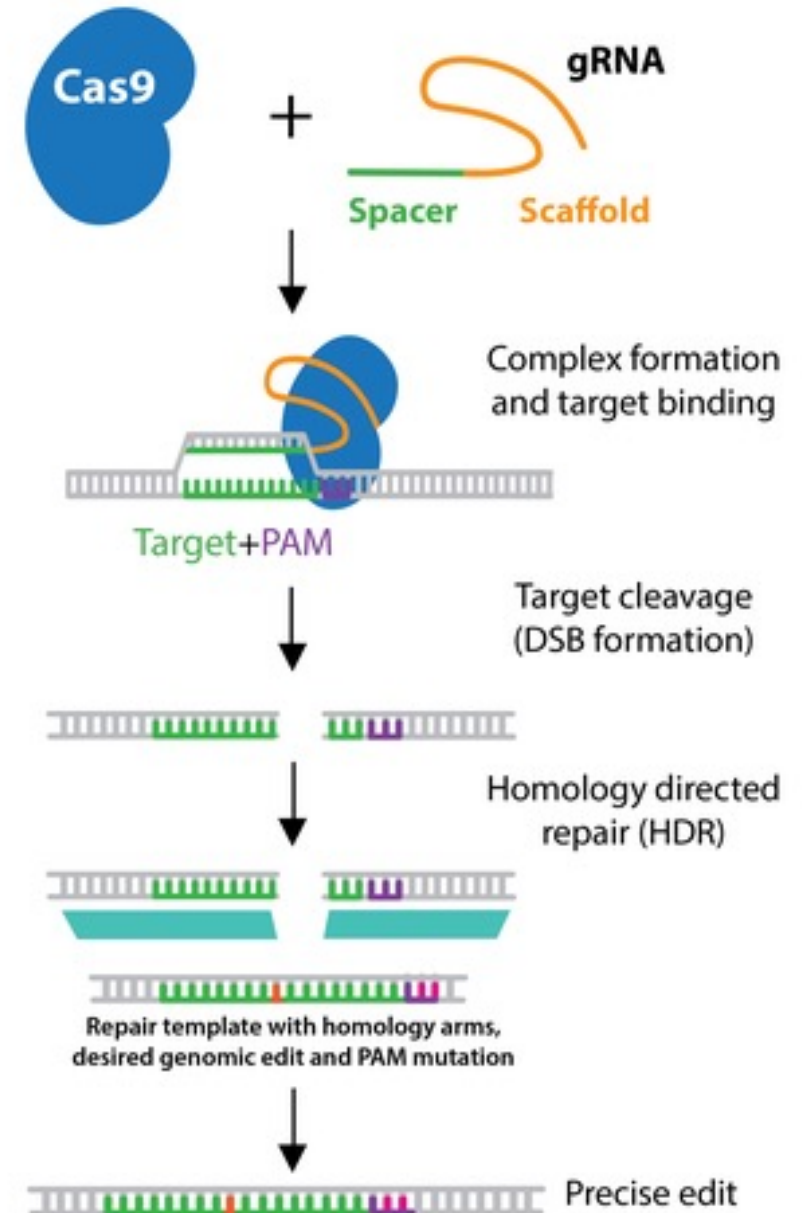
Non-Homologous End Joining



Homology Directed Repair

Limitations

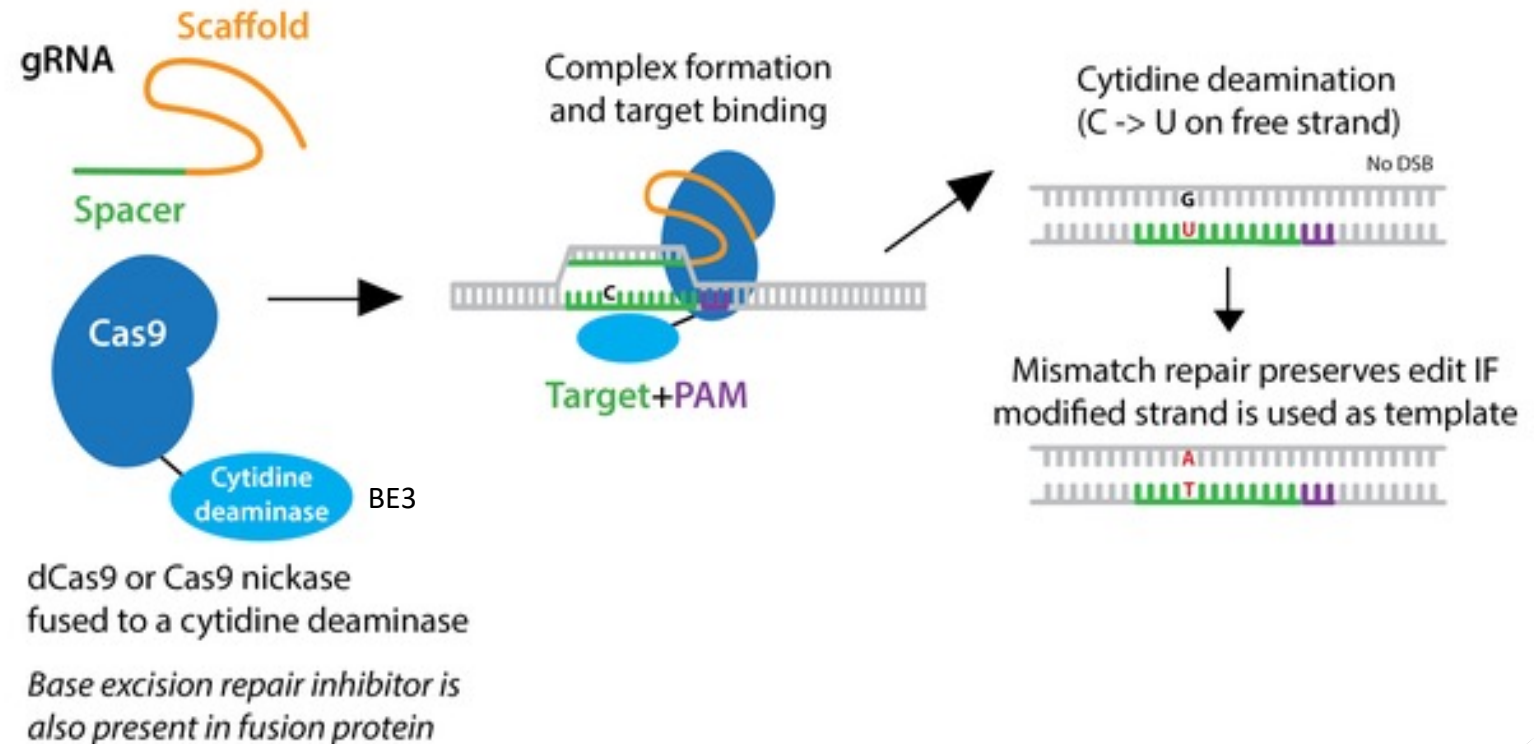
Difficulties delivering sufficient template DNA when repairing the double strand break in homology directed repair



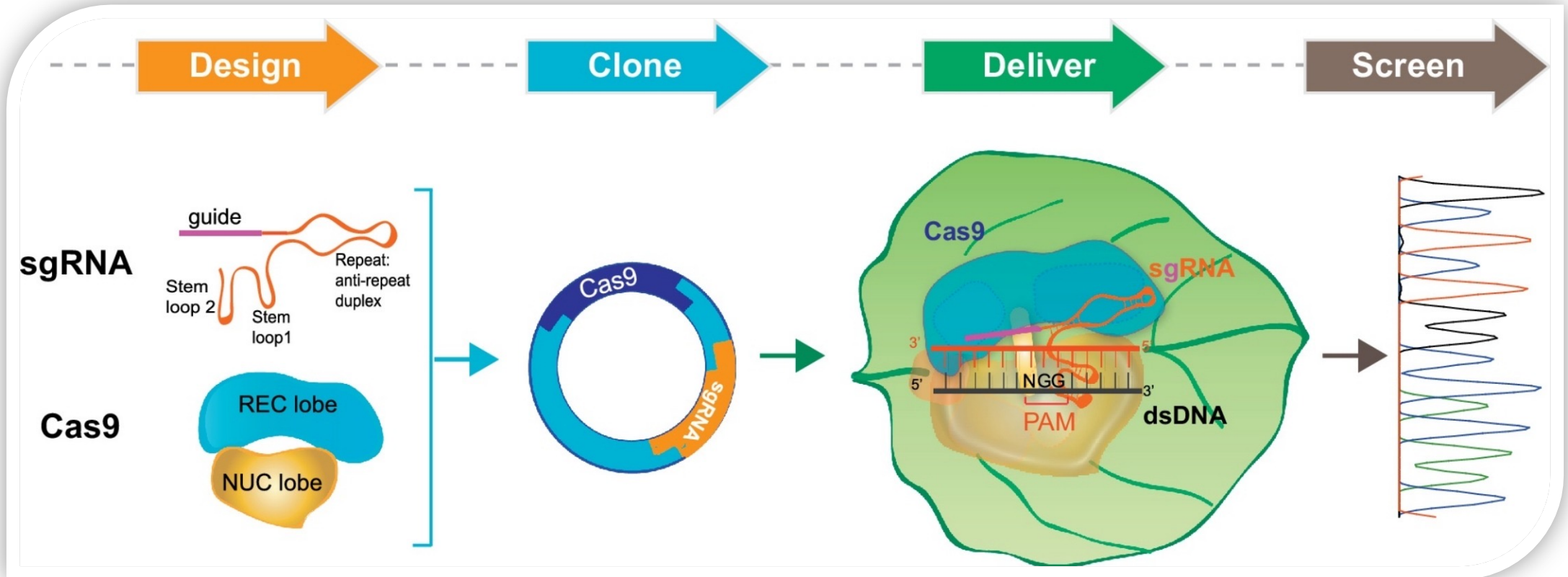
CRISPR/cas9 Base Editing

Cytosine
Base
Editors
(CBEs)
C to T

adenine
Base
Editors
(ABEs)
A to G



CRISPR/cas9 delivery into plants



Methods





The Target Gene

Watermelon Acetolactate
Synthase gene
CIALS



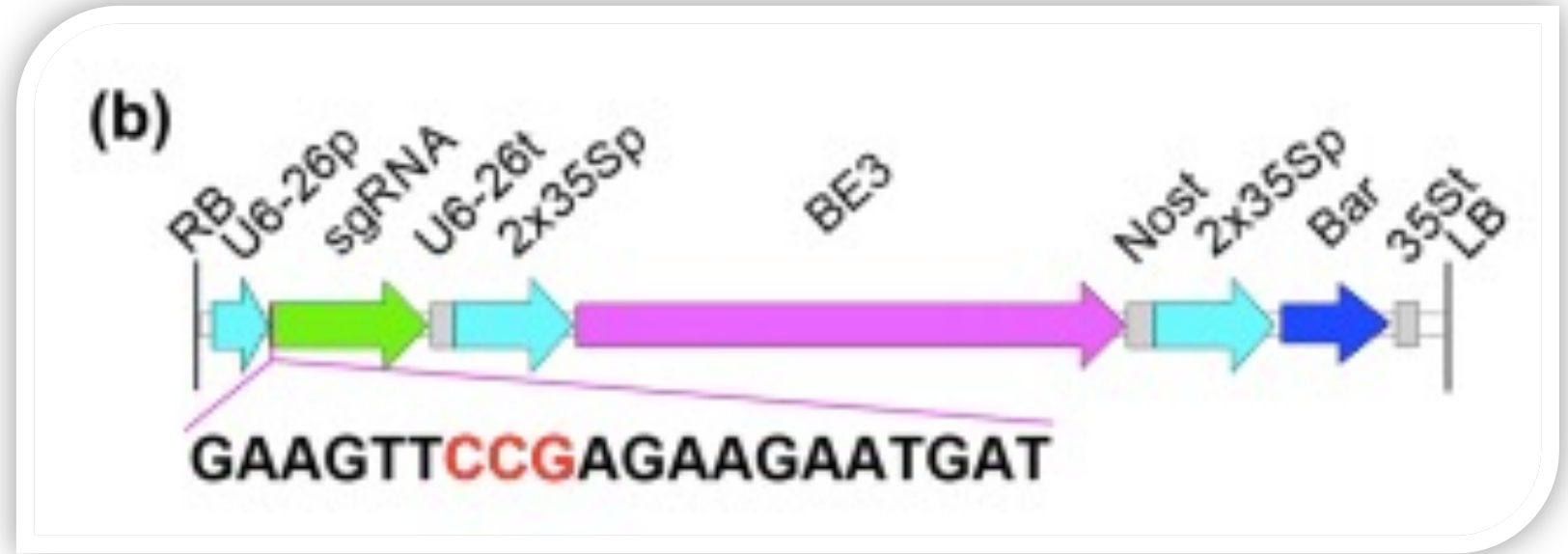
Point mutation
C to T In Pro190 (CCG)

Herbicide resistance



Transformation vector

pBSE901



(a)

3 4 5 6 7 8 9

At CAAGTCCCTCGTCGTATGATTGG

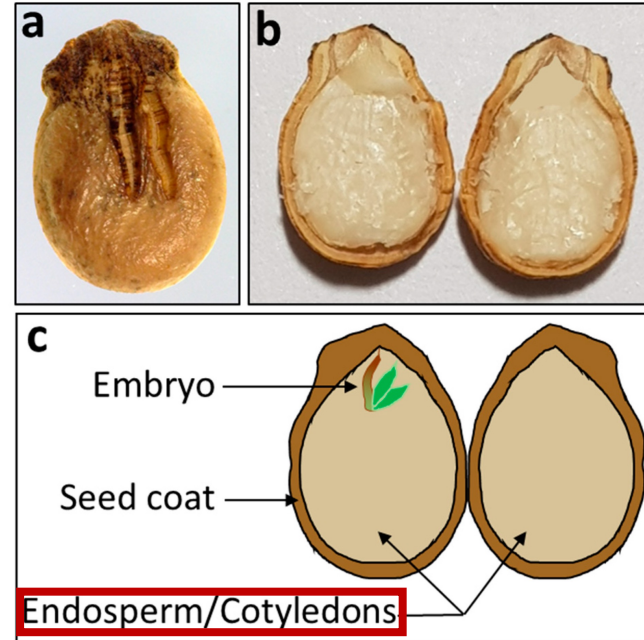
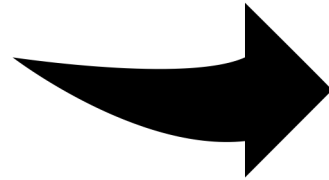
Cl CAAGTCCGAGAAGAATGATCGG

Q V P R R M PAM

Agrobacterium transformation



EHA105



**Watermelon
ZG94**





Results

PCR



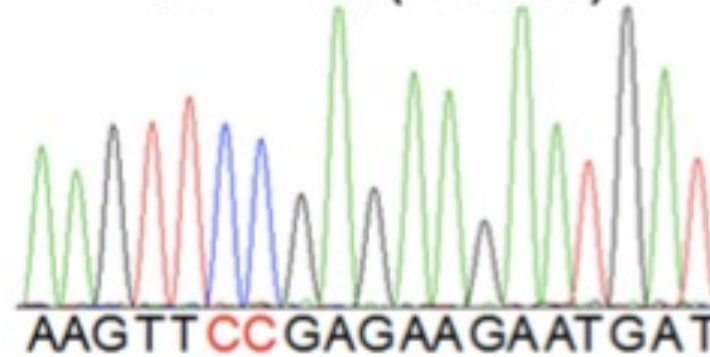
all candidate plants contained genes of *BE3* and *gRNA*

Sanger Sequencing (T0)

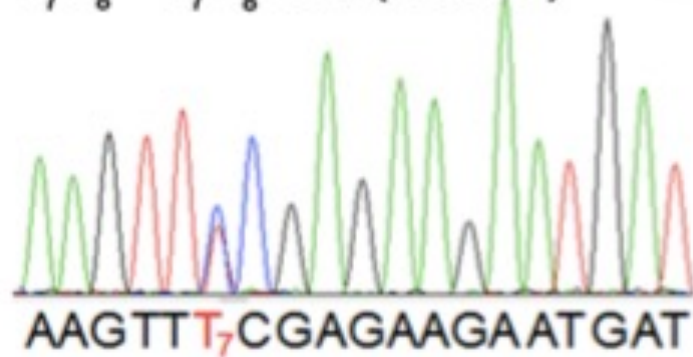
P190S

P190L

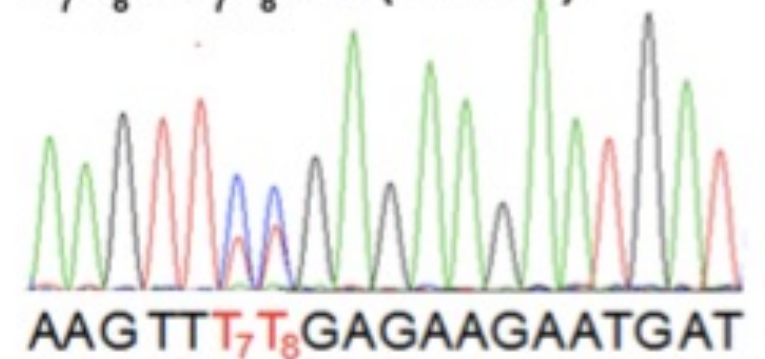
Wild Type 77%(154/199)



$C_7C_8 \rightarrow T_7C_8$ 17%(34/199)



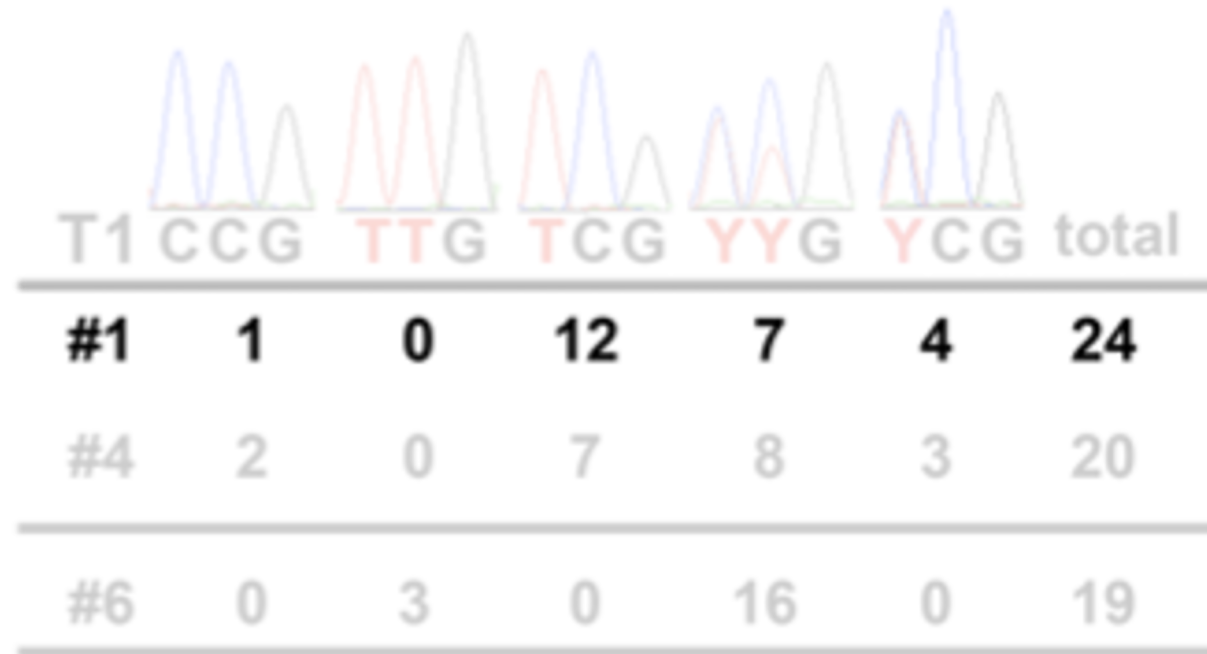
$C_7C_8 \rightarrow T_7T_8$ 6%(11/199)



Inheritance

T0: **CCG** to **TCG**

Wild	4% (1/24)
Homozygous	50% (12/24)
Hetrozygous	46% (11/24)



Inheritance

T0: **CCG** to **TCG**

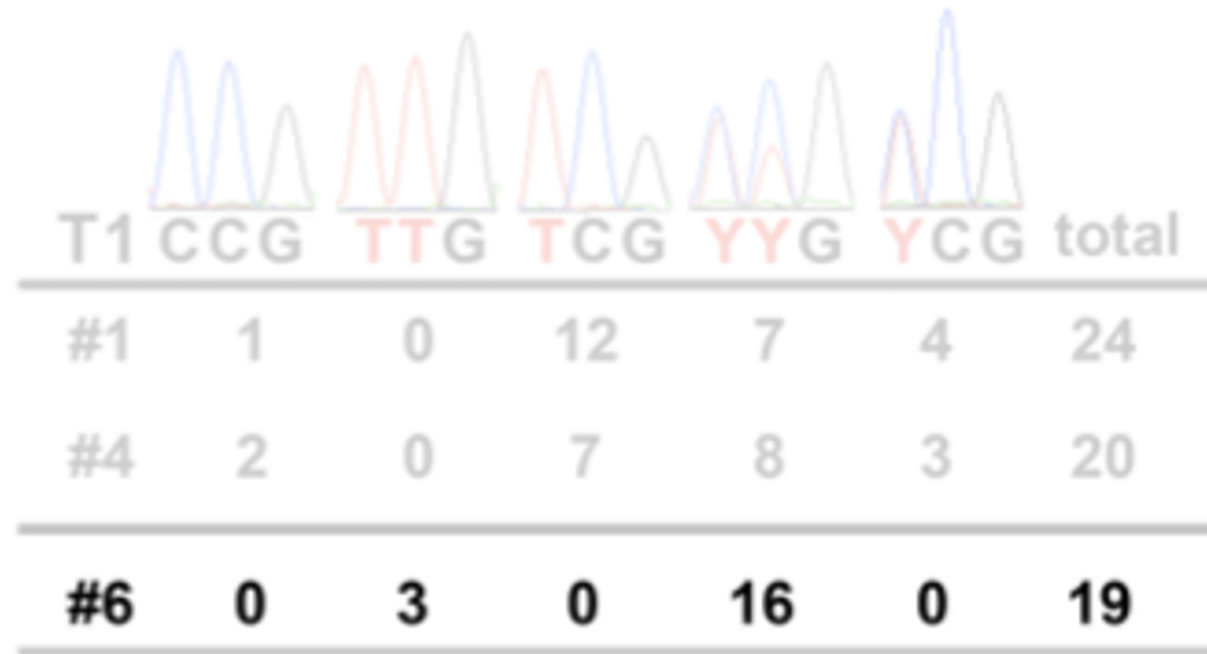
Wild	10% (2/20)
Homozygous	35% (7/20)
Hetrozygous	55% (11/20)



Inheritance

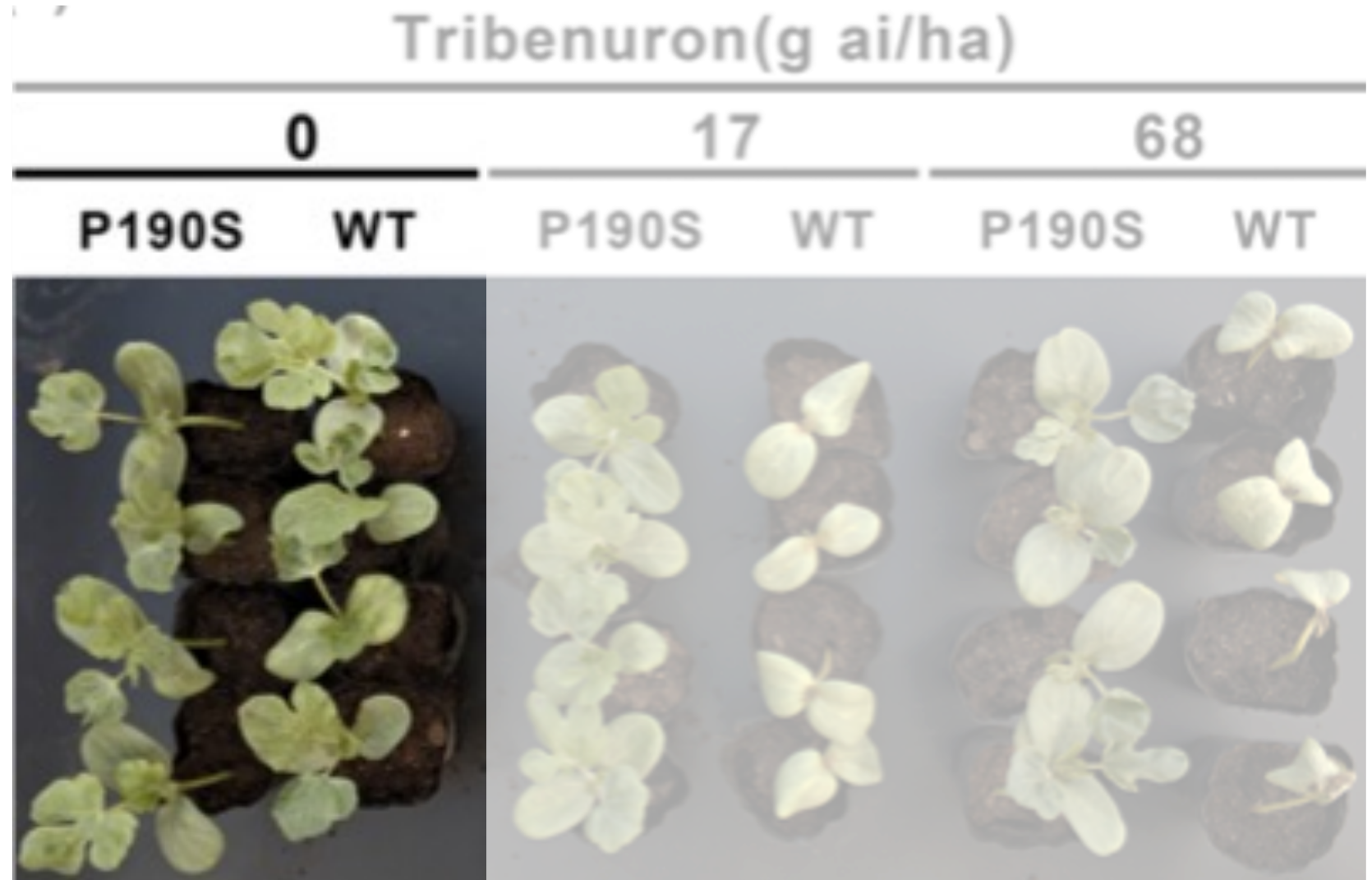
T0: **CCG** to **TTG**

Wild	null
Homozygous	16% (3/19)
Hetrozygous	84% (16/19)



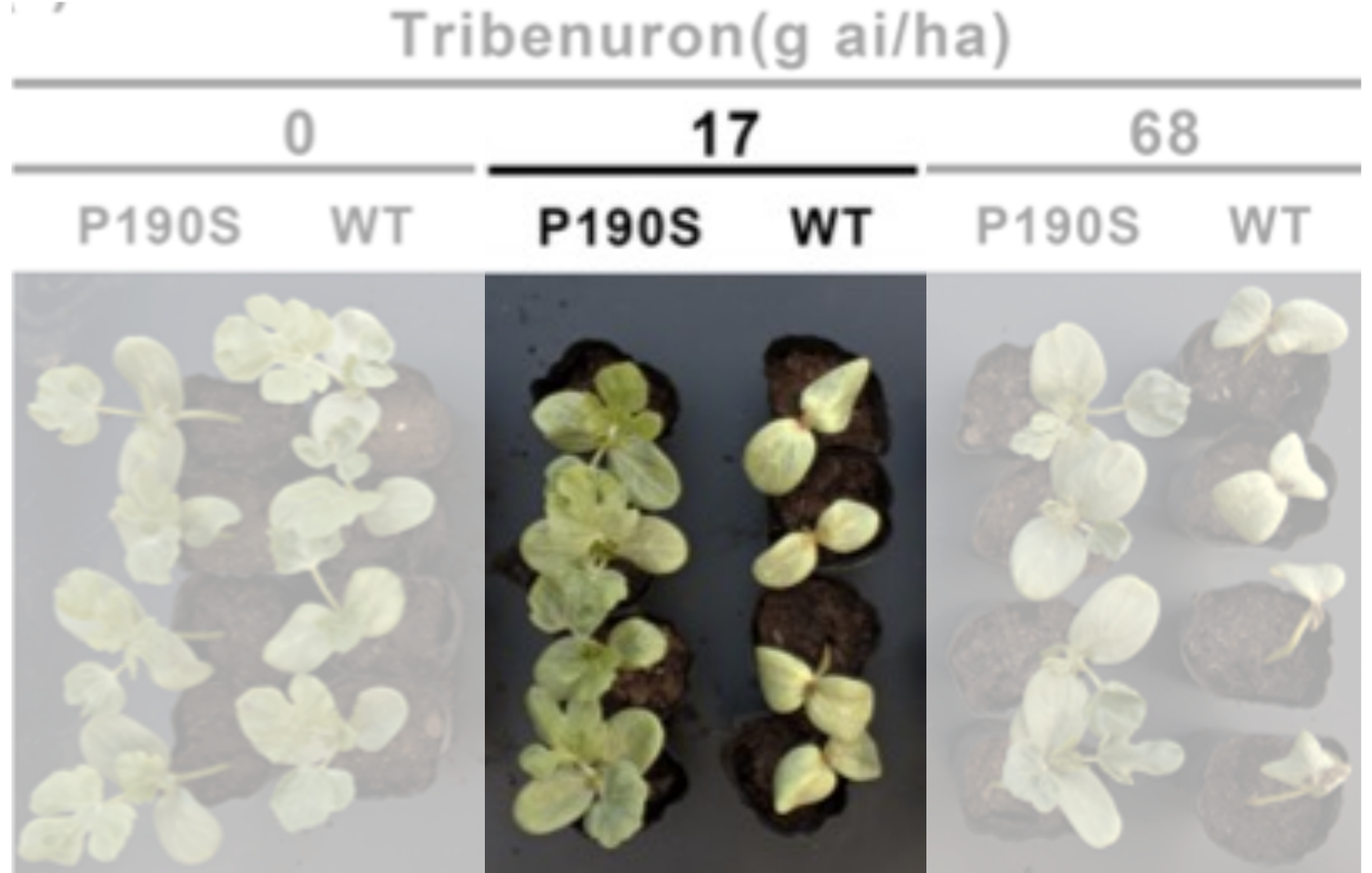
Herbicide Resistance

14 days after
herbicide
treatment



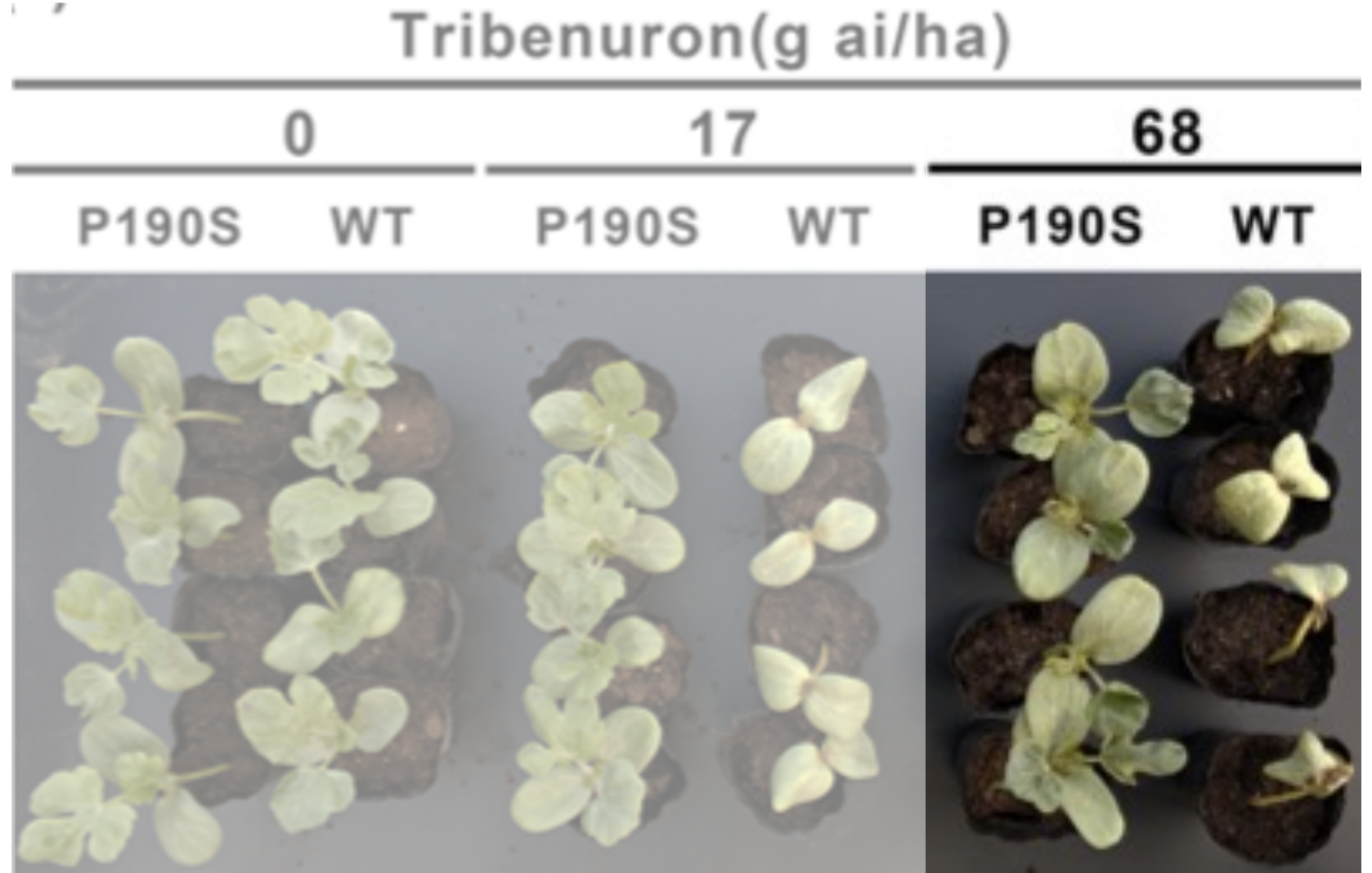
Herbicide Resistance

14 days after
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Herbicide Resistance

14 days after
herbicide
treatment



Discussion

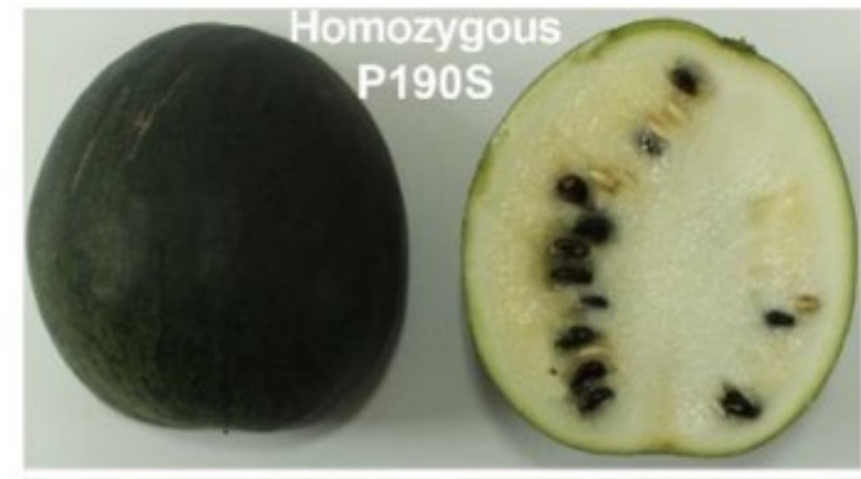


**What do you think the problems with
this technique would be ?**

Properties of the base edited watermelon



Fruit size and colour did not change



Properties of the base edited watermelon

	^a Seeds Number	^b Seeds Length(cm)	^c Seeds Weight(g)
Wild type	205±7	10.37±0.12	2.31±0.04
Homozygous	213±4.5	10.16±0.11	2.33±0.03
Heterozygous	209±4.7	10.33±0.15	2.30±0.02

The seed number per fruit, seed length and weight per 10 seeds
Did not change

Wild Type

Homozygous
P190S

Heterozygous
P190S



The seed size did not change

Off-Target potential



Whole genome search

5 locations with mismatches

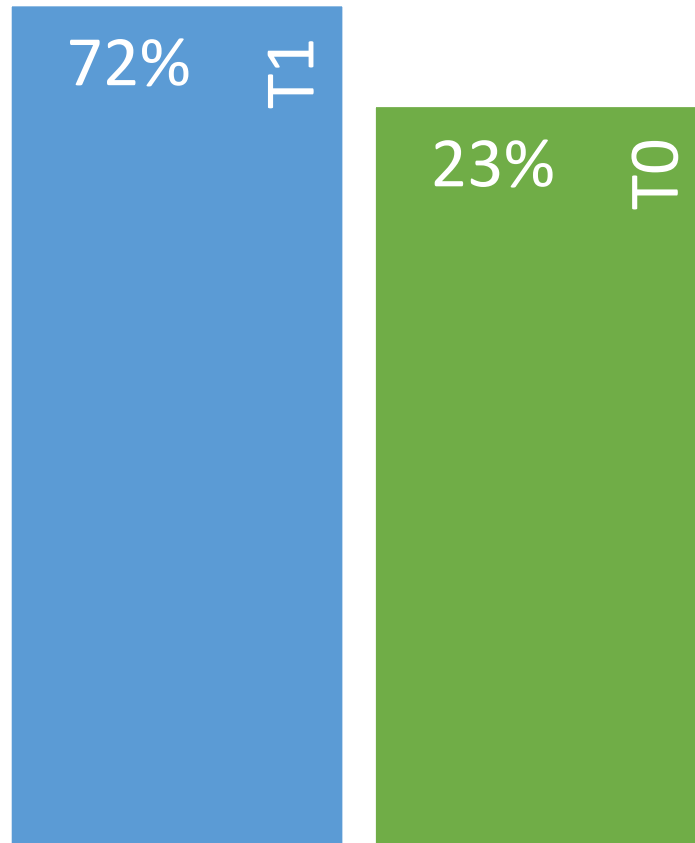
PCR + Sanger sequencing

No off-target edits found or indels




Conclusions

System efficiency



Implementation

Transgene-free base-edited herbicide-resistant watermelon plants are genetically identical to those bred via traditional mutagenesis



No extra regulations should be applied



Ready for immediate field application



**Thank you
for
listening**