



GENOME EDITING IN PERENNIAL RYEGRASS

PROTOPLASTS

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PERENNIAL RYEGRASS

- *Lolium perenne* L. is
 - Important temperate grass used for forage and turf
 - The most widely cultivated forage grass in Europe
- However, it doesn't grow well under

- Drought



- Freezing



Limits its cultivation
in Northern regions

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Concern due to
climate change



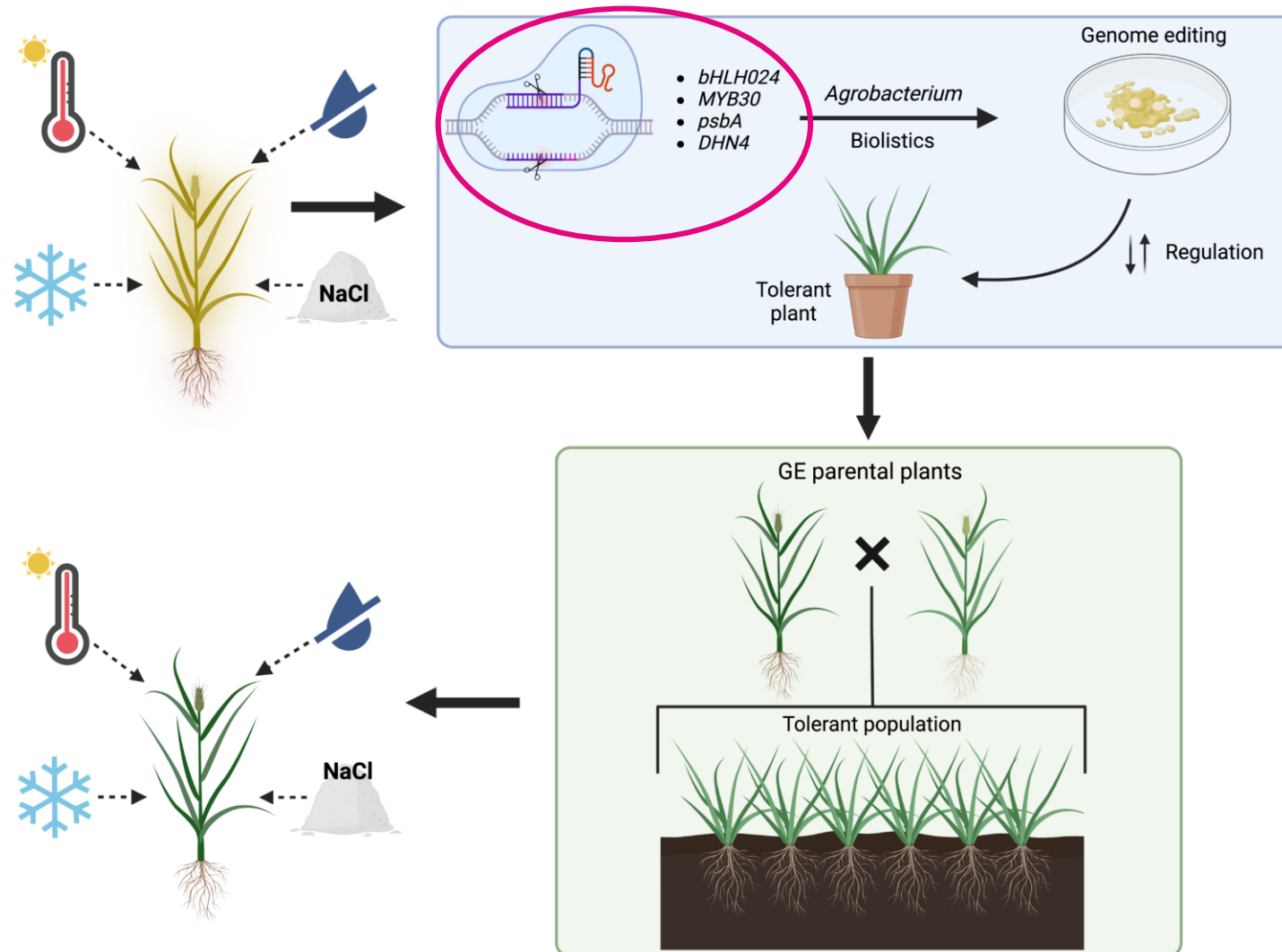
GENETICS AND REPRODUCTION OF *Lolium perenne*

- Wind pollinated
 - Obligate outcrossing species
 - Self-incompatible gametophyte
 - Highly heterogenous genome
 - Selective and traditional breeding
 - Not the best option for generation of abiotic tolerant lines
- 


Genome Editing

Using CRISPR-Cas to create tolerant plants

GENOME EDITING WITH CRISPR-CAS SYSTEMS

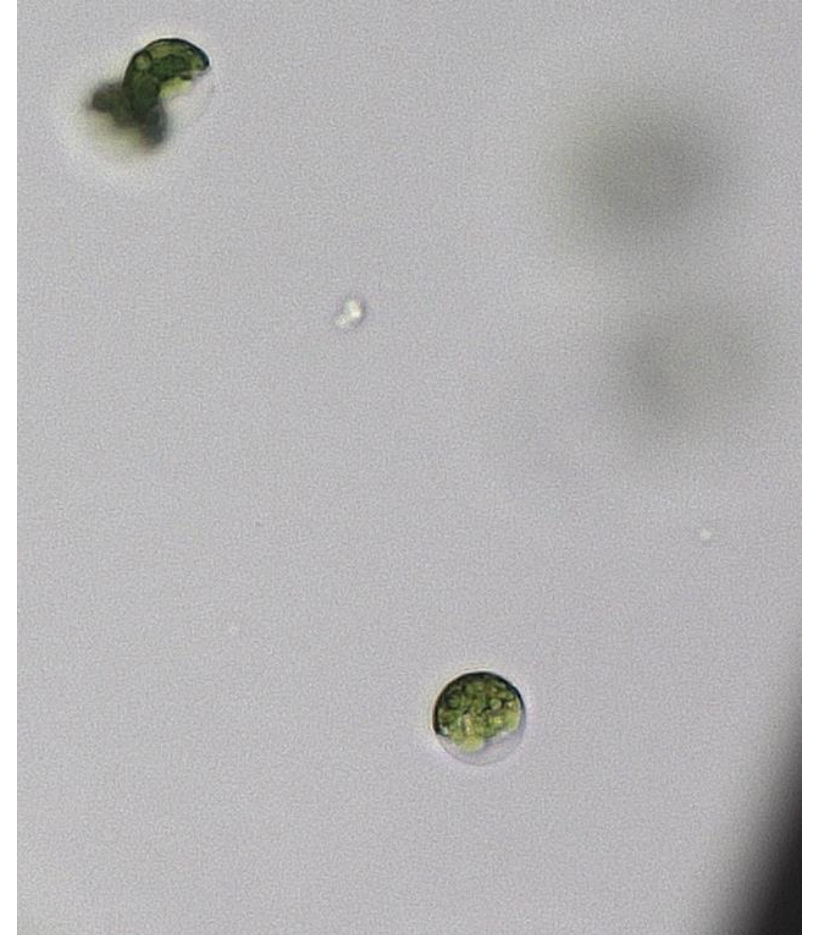


DESIGN AND SELECTION OF gRNAs

- Mostly using bioinformatic tools
 - Provide scores for
 - Specificity
 - Activity/Efficiency
 - The predicted activity is based on
 - The nucleotides of the guide
 - Data sets of previous experiments
 - Bottleneck
 - Predicted efficiency doesn't always correlate to the real activity of the guide
- 
- Test activity of guides **in vivo**
 - **Screen and select** the best performing gRNAs

USING PROTOPLASTS TO SELECT gRNAs

- Protoplasts are plant cells without a cell wall
- Used in multiple cellular, molecular and genetic studies
- Can be **isolated by millions**
 - Perfect for screening experiments
- Used for **selection of gRNAs** in plants
 - Arabidopsis (*Arabidopsis thaliana*)
 - Wheat (*Triticum aestivum*)
 - Rice (*Oryza sativa*)



***Lolium perenne* PROTOPLASTS ISOLATION**

- Different articles describe the isolation of perennial ryegrass protoplasts
- We tested two of the most recent ones
 - “An efficient protocol for perennial ryegrass mesophyll protoplast isolation and transformation, and its application on interaction study between LpNOL and LpNYC1” Yu et al. 2017
 - We did not manage to get the same high number of isolated cells
 - “Genetic Transformation of Protoplasts Isolated from Leaves of *Lolium temulentum* and *Lolium perenne*” Davis et al. 2020
 - We could not get a protoplast suspension with low amount of debris
- Therefore, we decided to establish a protocol based on these two publications

DEVELOPING A PROTOPLAST ISOLATION METHOD

- Highly reproducible and with consistent high yields
 - Around 1×10^6 cells per mL of suspension
- Low amount of debris
- Testing different variables

Mannitol pretreatment (Molar concentration)	Cellulase concentration (w/v)	Enzymatic treatment length	Vacuum infiltration (KPa)
<ul style="list-style-type: none">• 0.2 M• 0.3 M• 0.5 M• 0.6 M	<ul style="list-style-type: none">• 1.5%• 2%• 2.5%• 3%	<ul style="list-style-type: none">• 8 hours• 12 hours• 16 hours• 20 hours	<ul style="list-style-type: none">• 0• 71

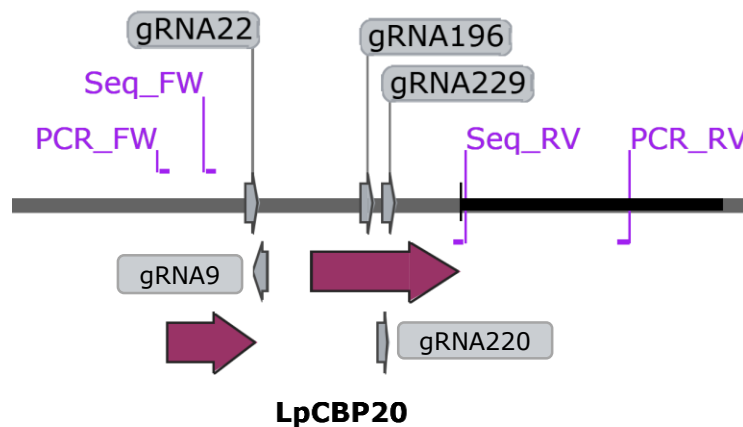
DROUGHT RESISTANCE THROUGH CRISPR-CAS KNOCKOUT

- Targeting gene *LpCBP20*
- Negatively regulates the synthesis of cuticular waxes
- Knocking out *cbp20* in barley produced drought resistant plants (Daszkowska-Golec *et al.* 2020)

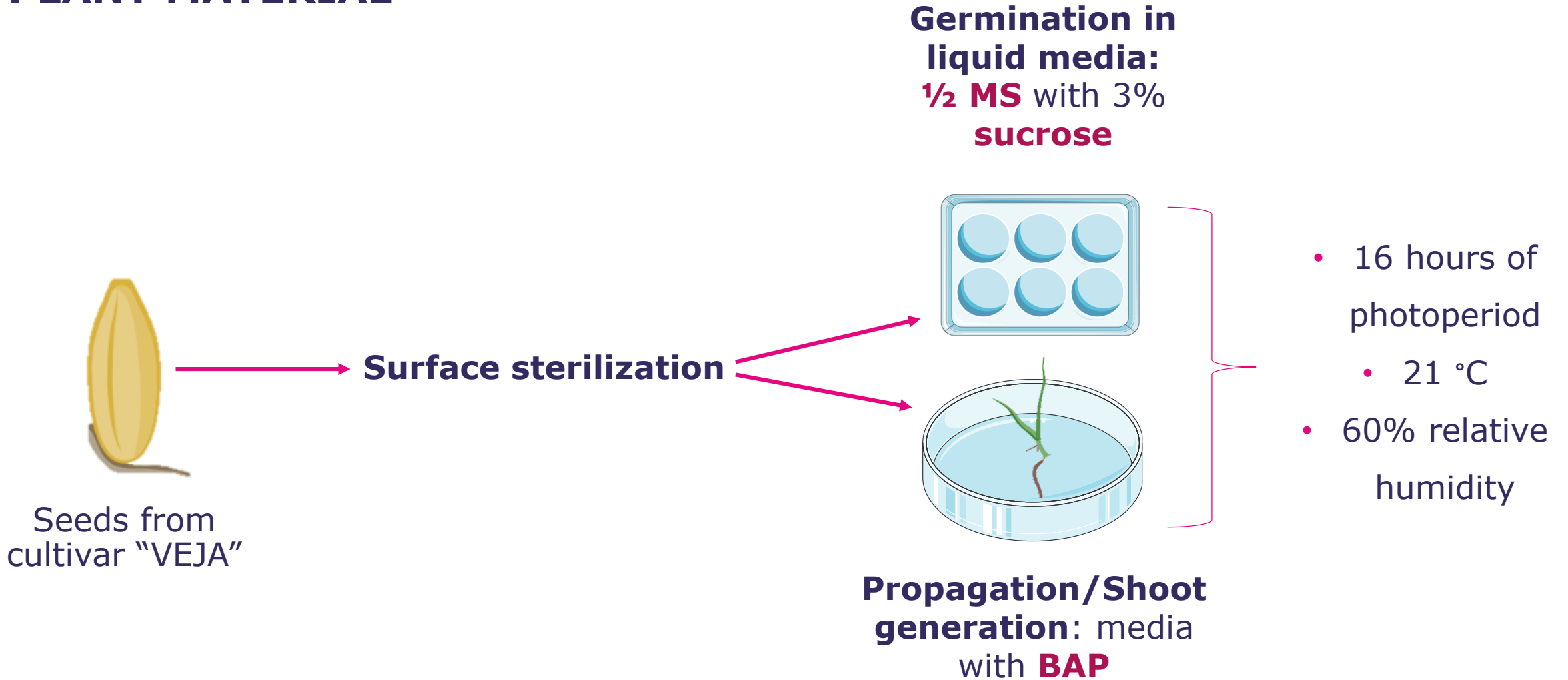


TRANSFORMING PROTOPLASTS WITH DIFFERENT VECTORS

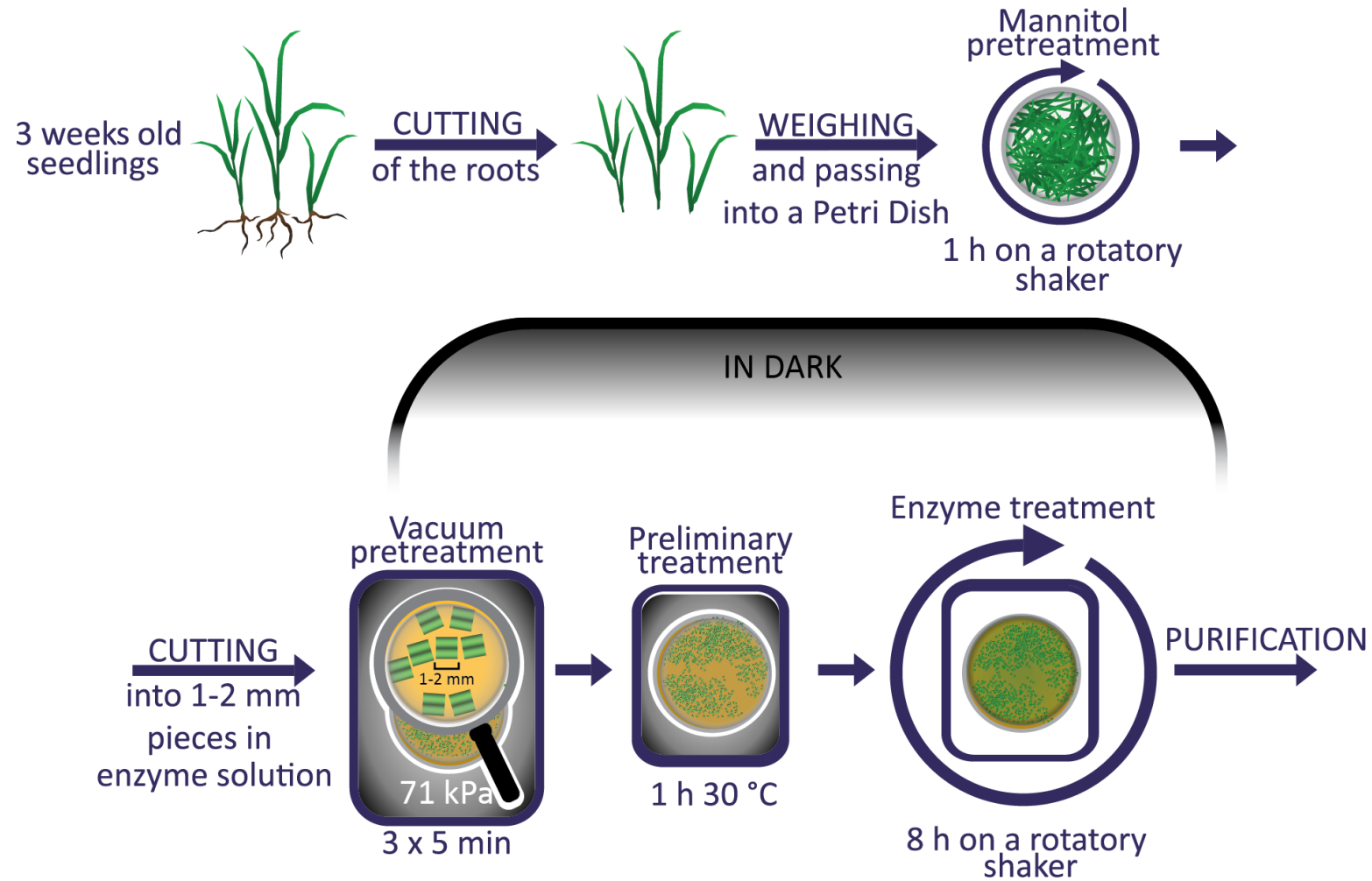
- Using a plasmid with one single gRNA
 - EGFP cassette
- 3 different vectors targeting the second exon of CBP20
 - p196
 - p220
 - p229
- Using a plasmid with 5 different gRNAs
 - ZsGreen cassette
- One vector with
 - 2 gRNAs targeting the first exon
 - 3 gRNAs targeting the second exon



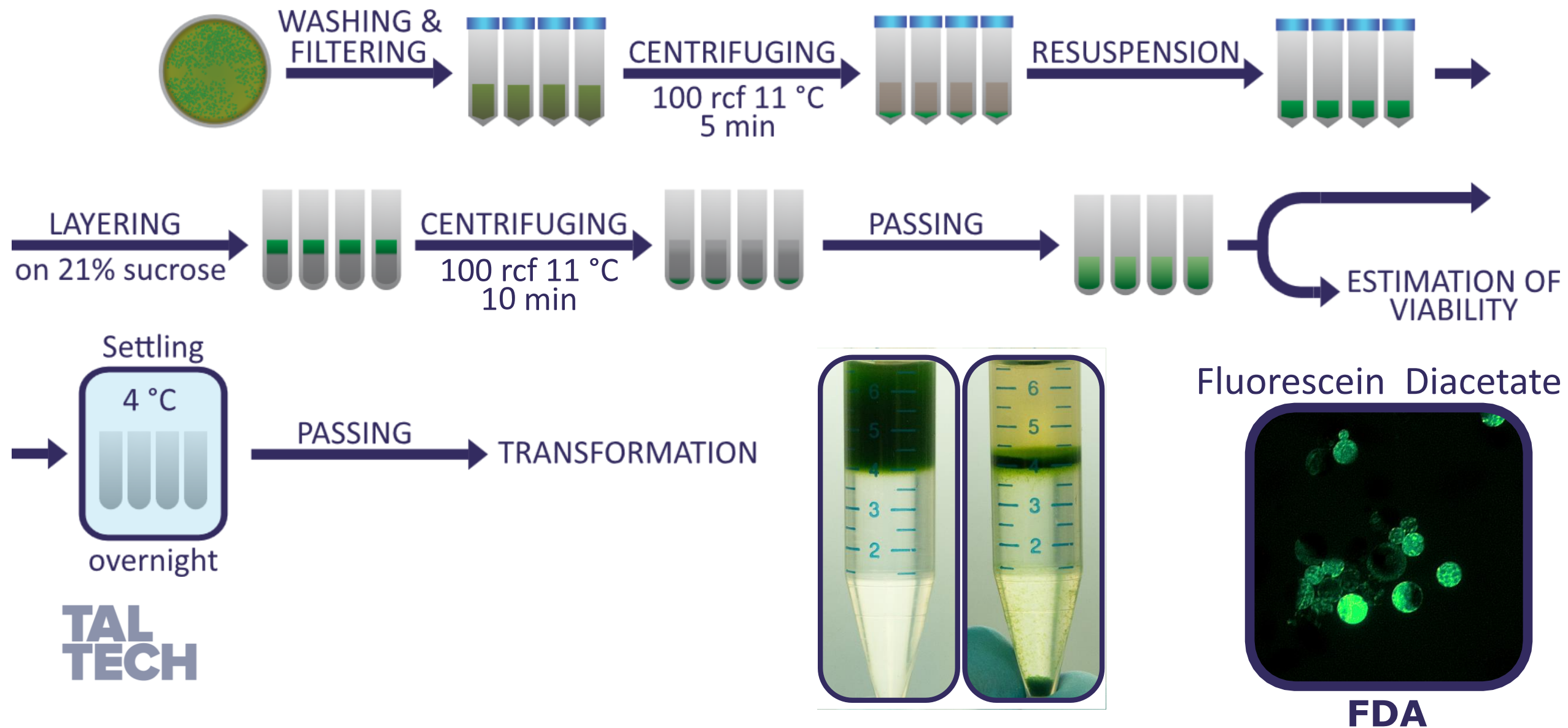
PLANT MATERIAL



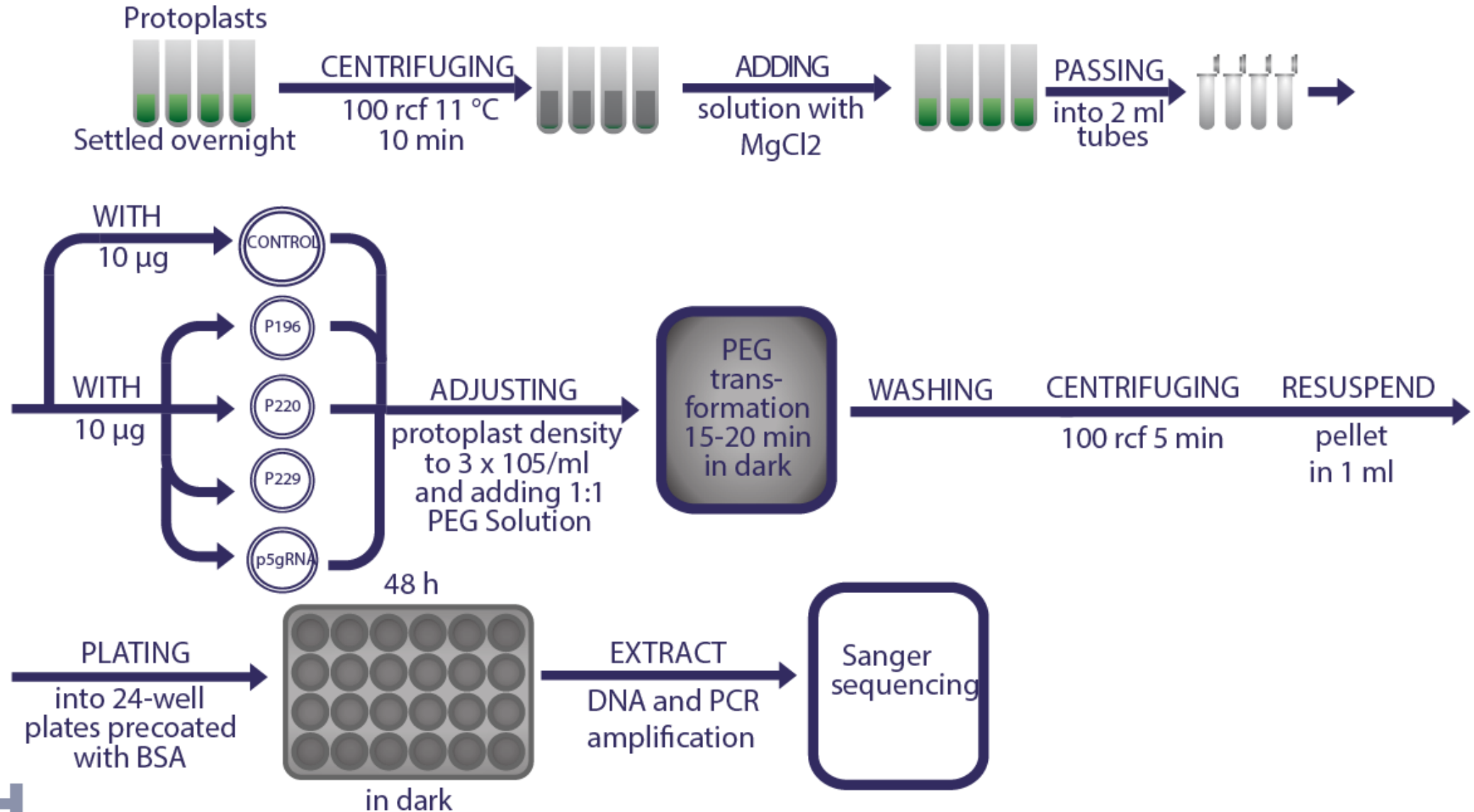
PROTOPLASTS ISOLATION



PROTOPLASTS PURIFICATION

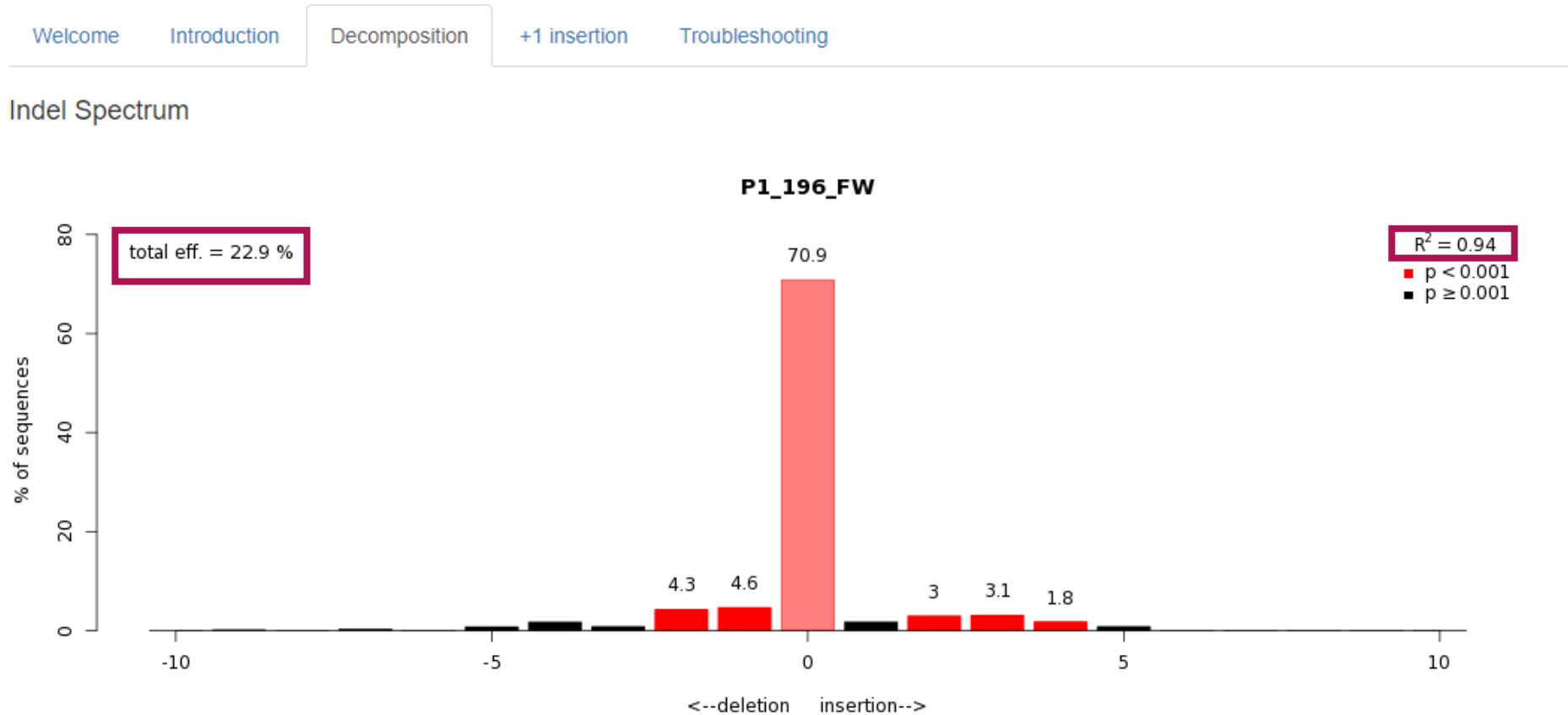


PROTOPLASTS TRANSFORMATION



EDITING EFFICIENCY

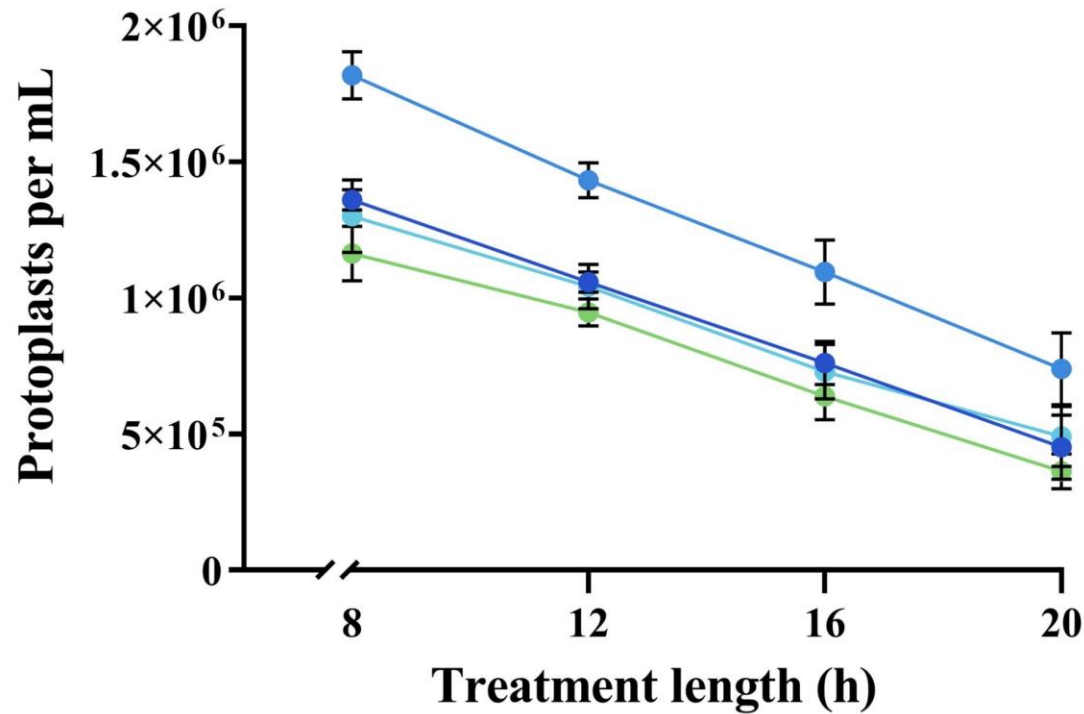
- Using Sanger sequencing trace data
- Analysis by decomposition using TIDE
- Compares non-transformed and transformed sequences
- Provides data showing the frequency of indels present in the samples



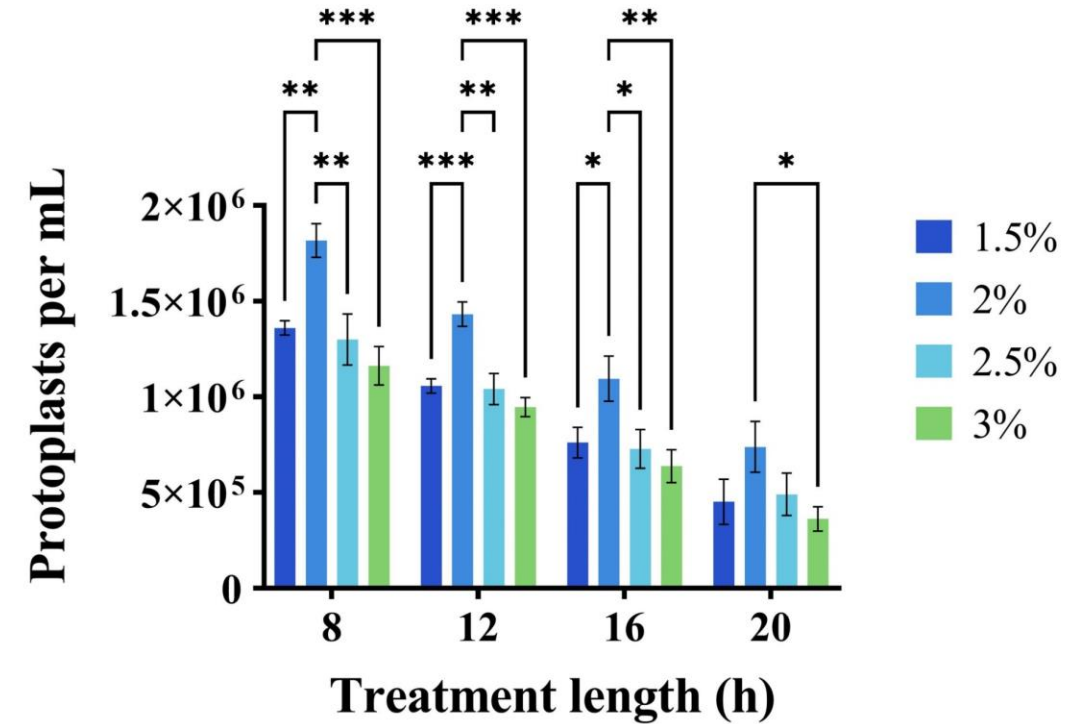
RESULTS

CELLULASE AND ENZYMATIC TREATMENT

a) Viable protoplasts after enzymatic treatment

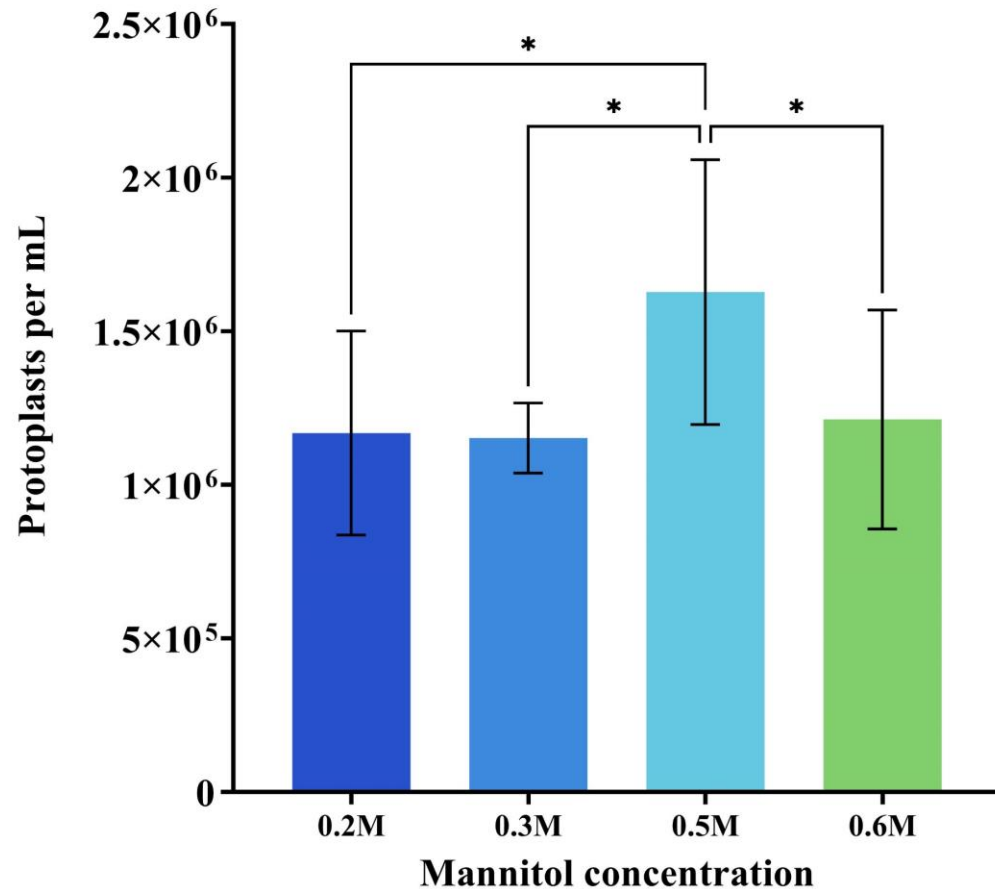


b) Viable protoplasts after enzymatic treatment

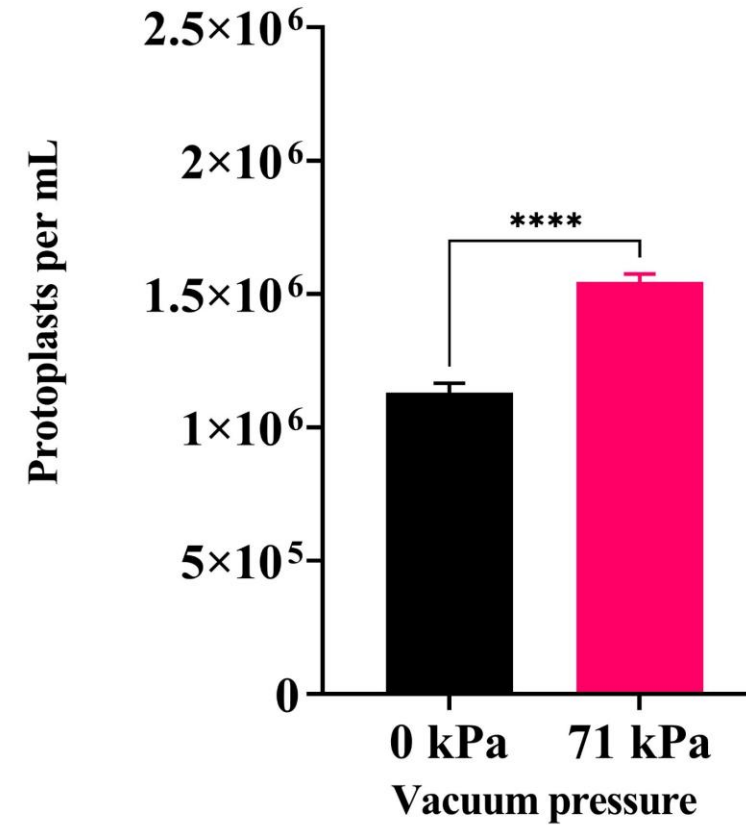


MANNITOL AND VACUUM INFILTRATION

a) Viable protoplasts after mannitol pretreatment

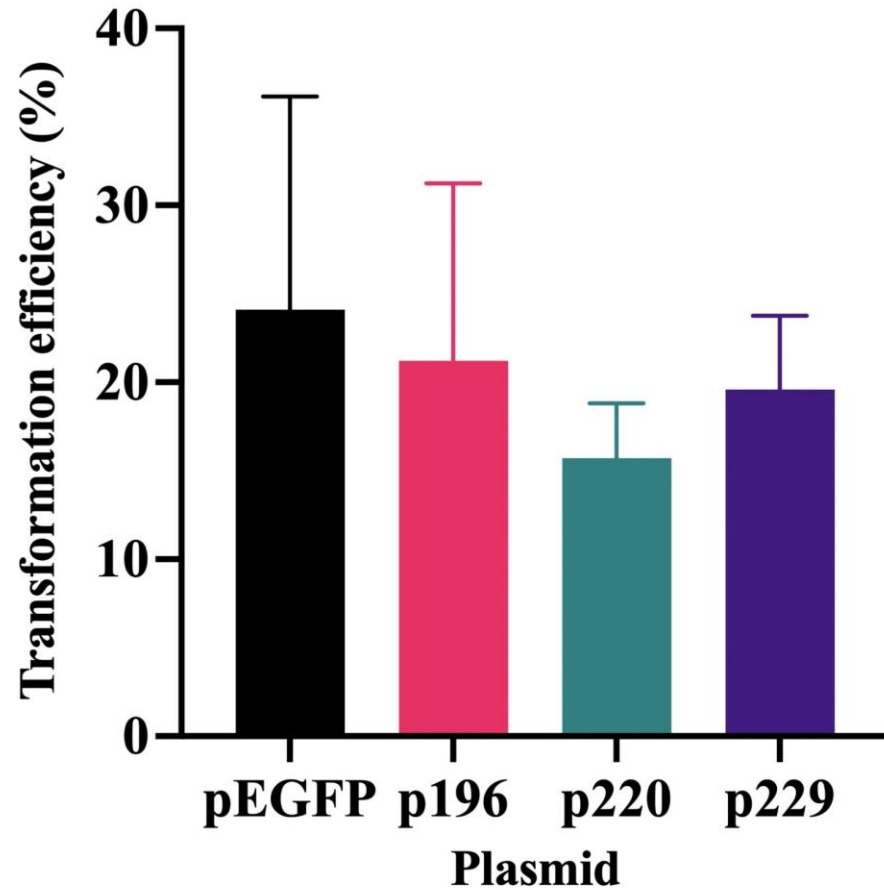


b) Viable protoplasts after vacuum infiltration

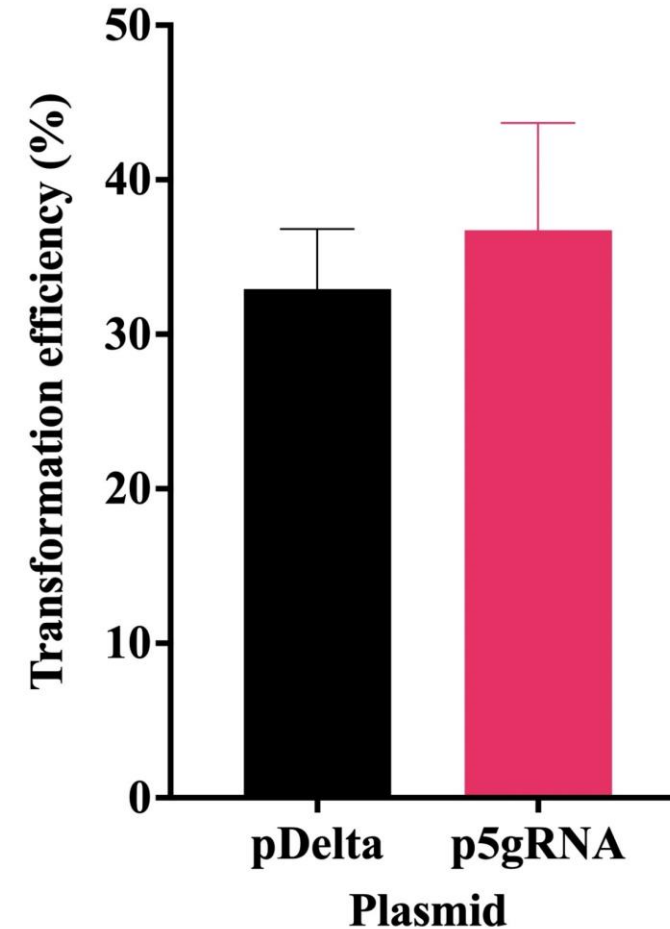


PEG-MEDIATED TRANSFORMATION

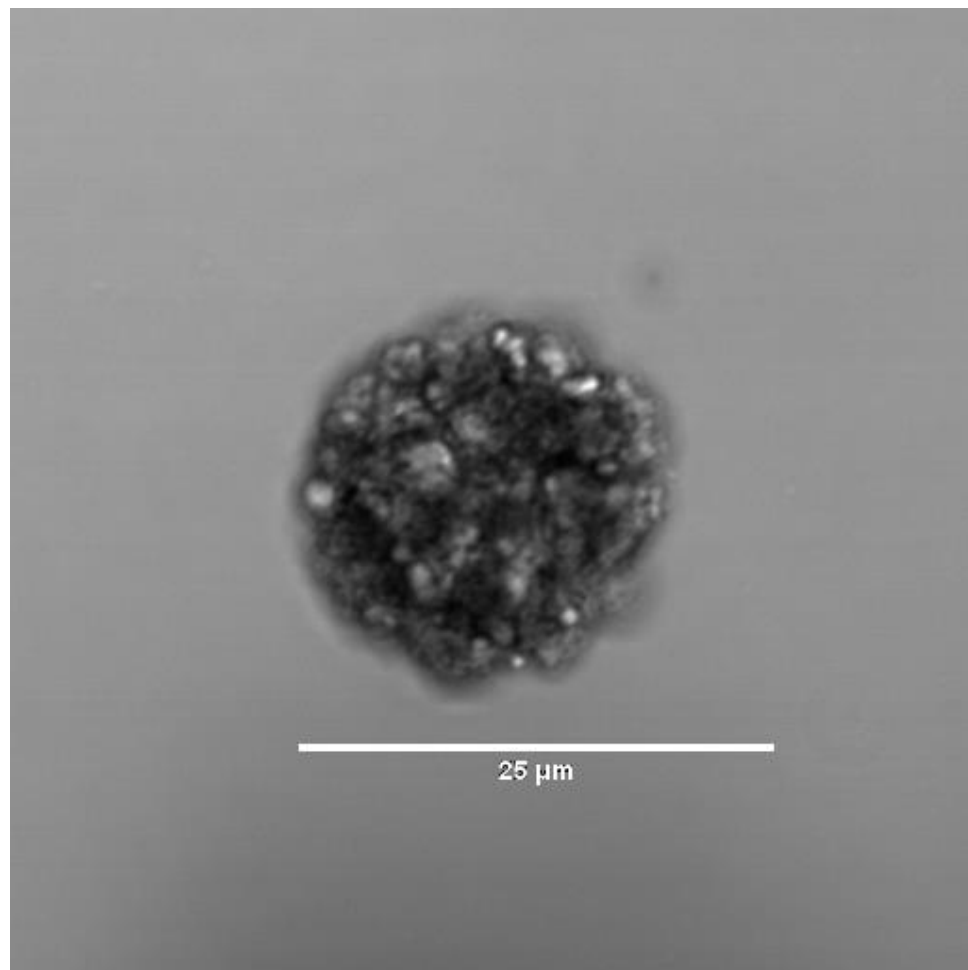
a) Transformation efficiency of pHSE401/EGFP



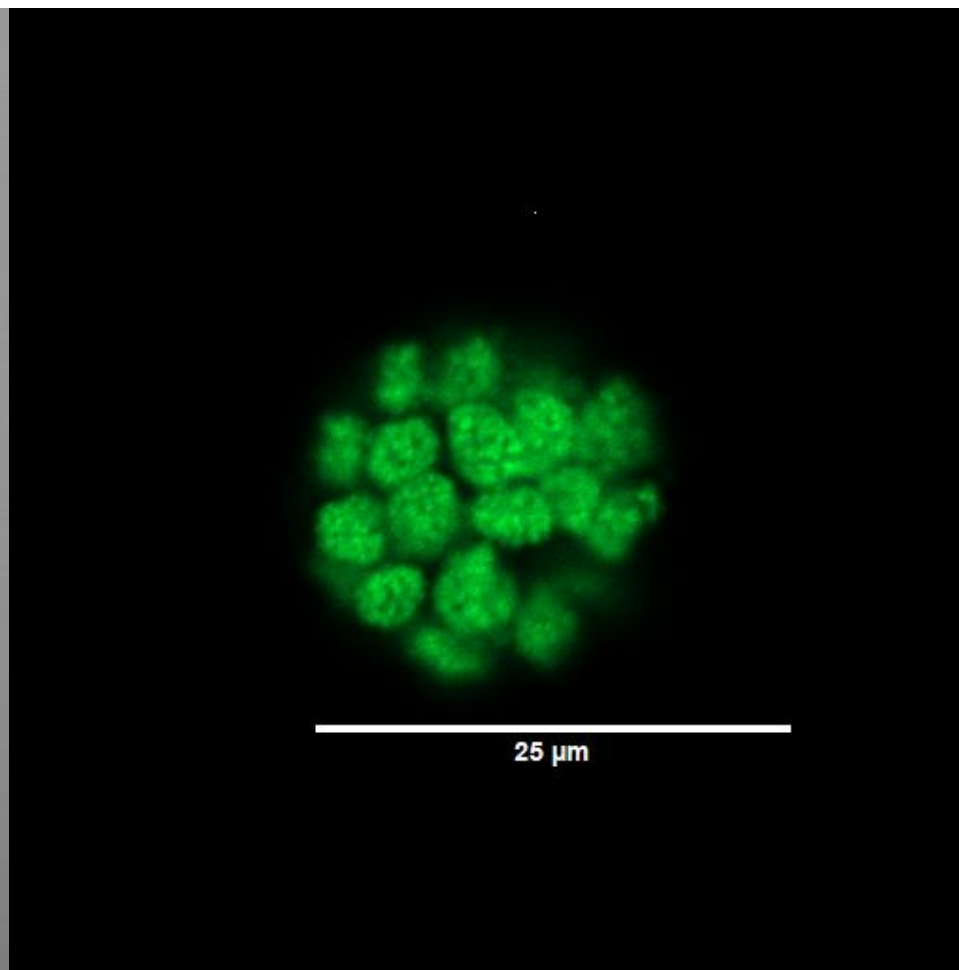
b) Transformation efficiency of the ZsGreen plasmid



ZsGREEN FLUORESCENCE



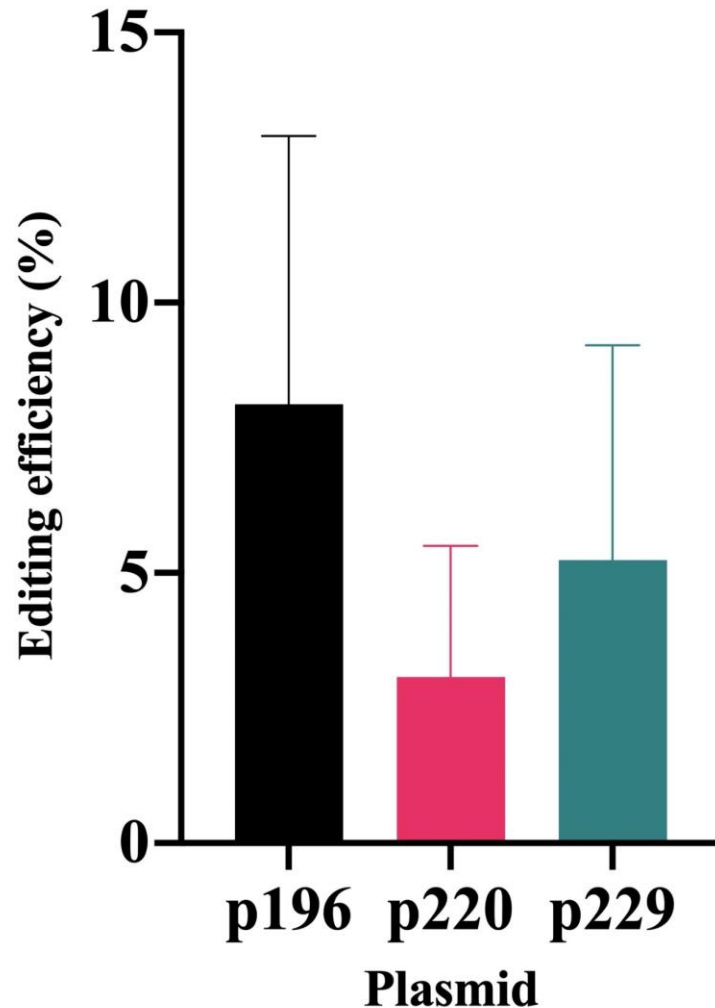
Brightfield



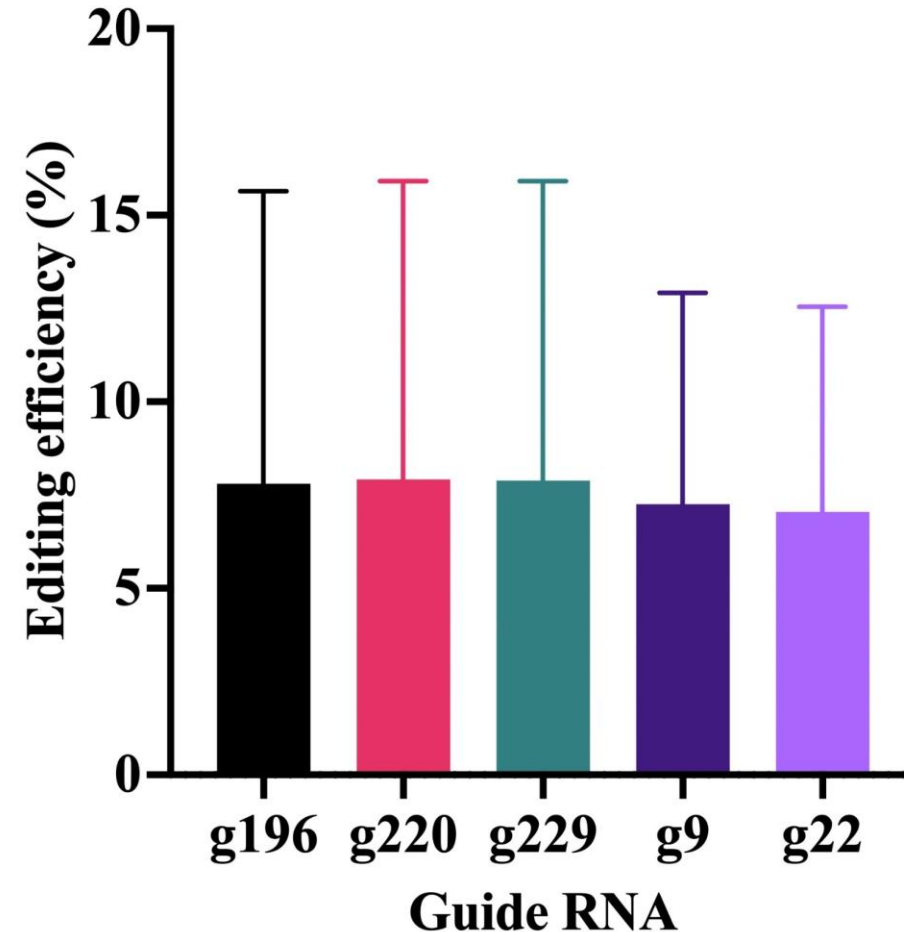
Fluorescence

EFFICIENCY OF THE DIFFERENT VECTORS AND gRNAs

a) Editing efficiency of pHSE401/EGFP



b) Editing efficiency of p5gRNA



FINAL REMARKS

- Successful PEG-mediated transformation
 - More than 20% of the treated cells were transformed (had fluorescence)
- The DNA directly extracted from the transformed cells
 - Suitable for amplicon sequencing
 - Good quality for TIDE analysis
- Editing efficiency was calculated with TIDE for all the gRNAs
 - On average, around 7% of the sequences presented indels

The proposed method can be used to **evaluate** the performance of **guide RNAs in vivo** with the aim of **selecting** the most **suitable material** for later transformation experiments.

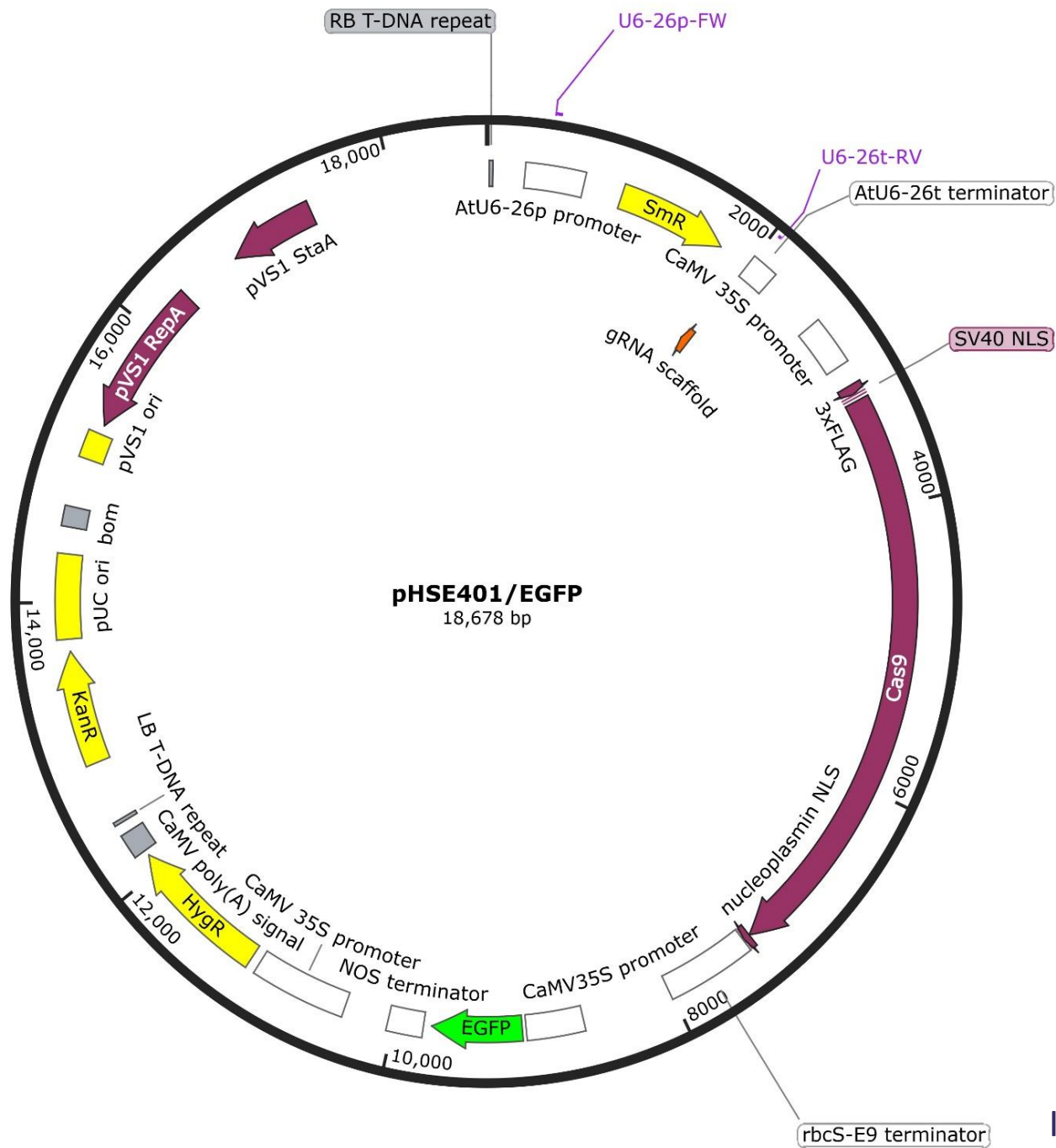
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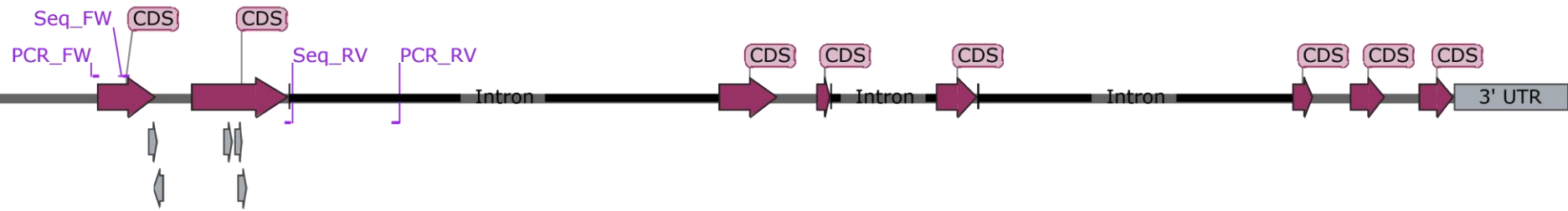


- Cecilia Sarmiento
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- Lenne Nigul
- Signe Nöu

TAL
TECH







LpCBP20