

# Transgenic Microalgae – Problems and Perspectives



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# Transgenic Microalgae – Problems and Perspectives

- Rationale – why do we want to genetically manipulate microalgae?

- Strategy for microalgal transfection (transformation)

- Overview to transformed microalgae

Diatoms

*Thalassiosira pseudonana*

*Phaeodactylum tricornutum*

Rhodophytes (red algae)

*Cyanidioschyzon merolae*

Chlorophytes

*Chlorella*

*Haematococcus pluvialis*

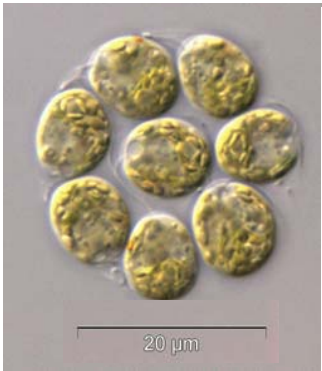
*Dunaliella salina*

*Chlamydomonas reinhardtii* (*Volvox carteri*)

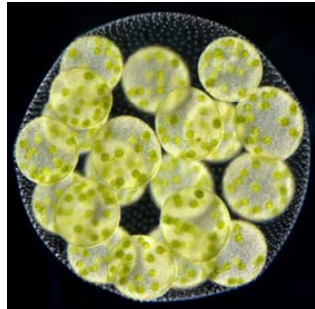
- Strategies for targeted knock-out / knock-down in *Chlamydomonas reinhardtii*  
knock-out libraries  
(inducible) amiRNAs
- Overcoming problems in gene expression by targeted chromatin remodeling
- Summary & Outlook

# Why transform microalgae?

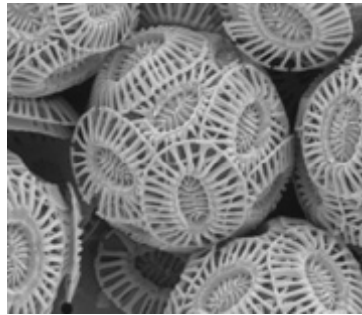
More and more microalgal genome sequences are completed:



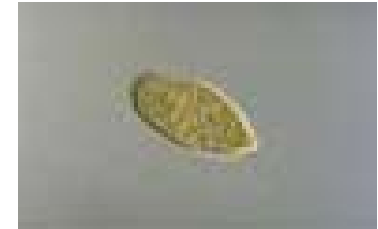
*Chlamydomonas reinhardtii*  
(Chlorophytes)



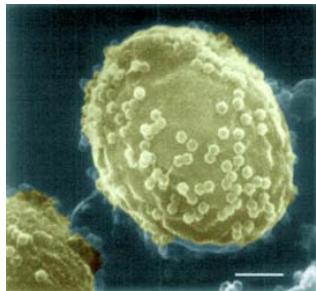
*Volvox carteri*  
(Chlorophytes)



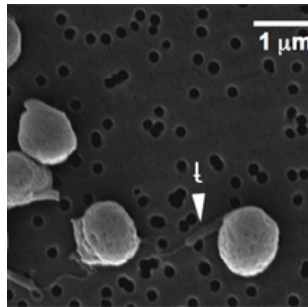
*Emiliana huxleyi*  
(Coccolithophores)



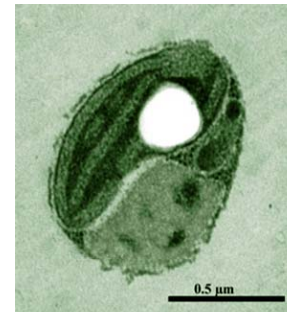
*Cyanidioschyzon merolae*  
(Rhodophytes)



*Chlorella variabilis*  
(Chlorophytes)



*Micromonas pusilla*  
(Prasinophytes)



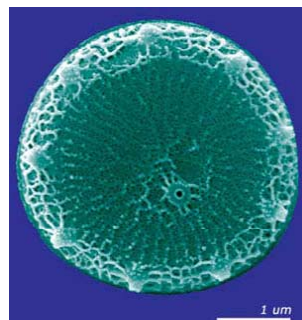
*Ostreococcus lucimarinus / tauri*  
(Prasinophytes)



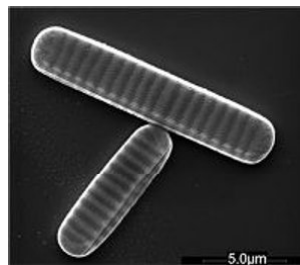
*Guillardia theta* & *Bigelowiella natans*  
(Cryptomonads)



*Phaeodactylum tricornutum*  
(Diatoms)



*Thalassiosira pseudonana*  
(Diatoms)



*Fragilariopsis cylindrus*  
(Diatoms)



*Aureococcus anophagefferens*  
(Pelagophytes)

Source: mainly JGI

# Why transform microalgae?

## Basic research

- Photosynthesis (*Chlamydomonas reinhardtii*)
- Cilia/flagella (*Chlamydomonas reinhardtii*)
- Development of multicellularity (*Volvox carteri* / *Chlamydomonas reinhardtii*)
- Primary and secondary endosymbiosis (Rhodophytes, Diatoms, Cryptophytes)
- Extremophiles (*Cyanidioschyzon merolae*, *Chlamydomonas nivalis/acidophila*)

## Ecological importance

- ~50% of annual carbon fixation by microalgae
- Algal blooms (*Aureococcus anophagefferens*)

## High-end commercial products

- Carotenoids (*Haematococcus*, *Dunaliella*)
- Feed stock for aquaculture (*Nannochloropsis*)
- Food supplementals (*Chlorella*)
- Hydrogen (*Chlamydomonas reinhardtii*)
- Biodiesel (*Botryococcus braunii*)

# Why transform microalgae?

## Metabolic engineering

### Reduce expression of algal genes

knock-out; knock-down by expressing antisense, inverted repeats, amiRNAs  
→ eliminate pathways competing with that leading to desired product

### Express foreign genes

→ generate strains with new biosynthetic properties

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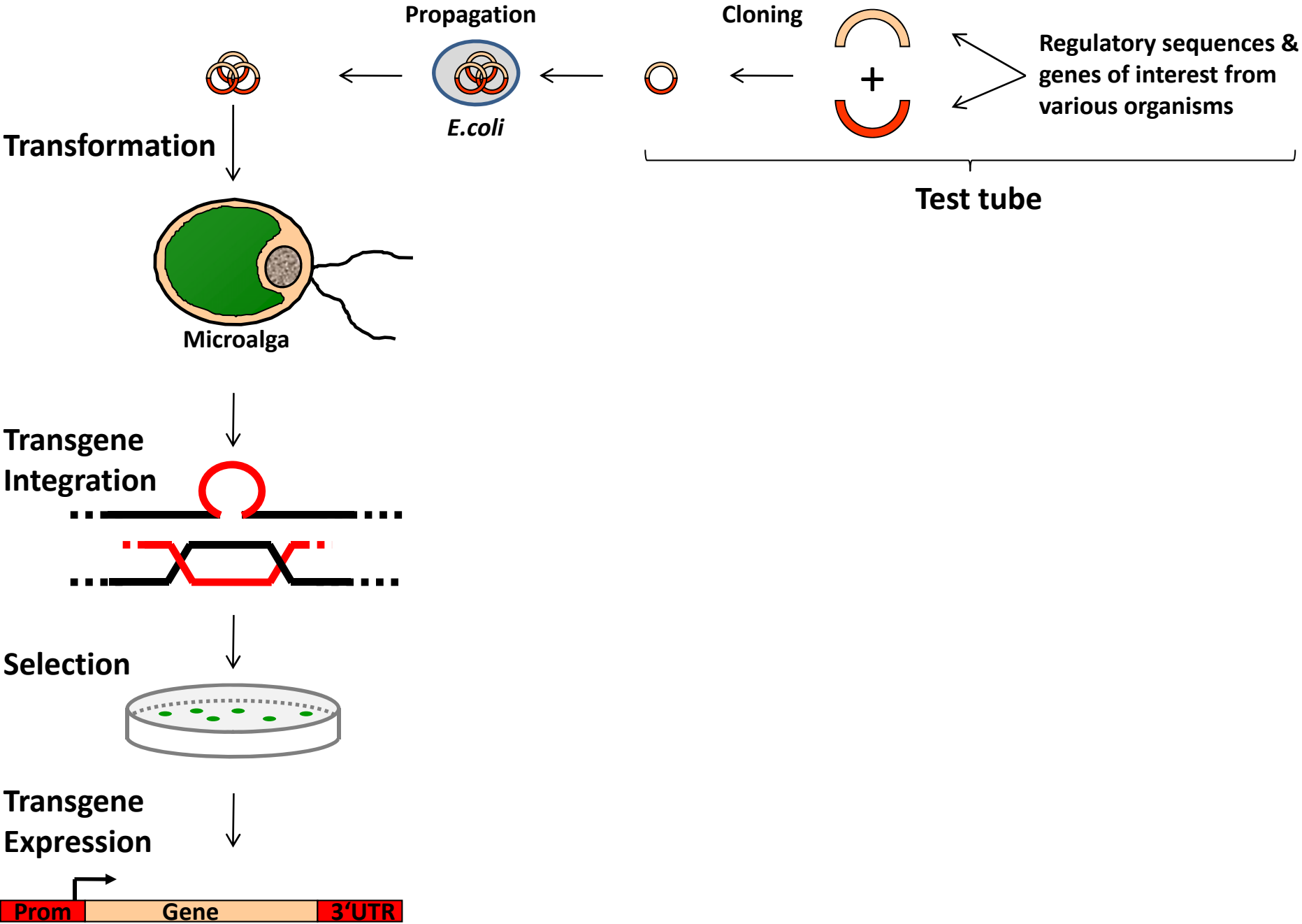
*Haematococcus pluviales*

*Dunaliella salina*

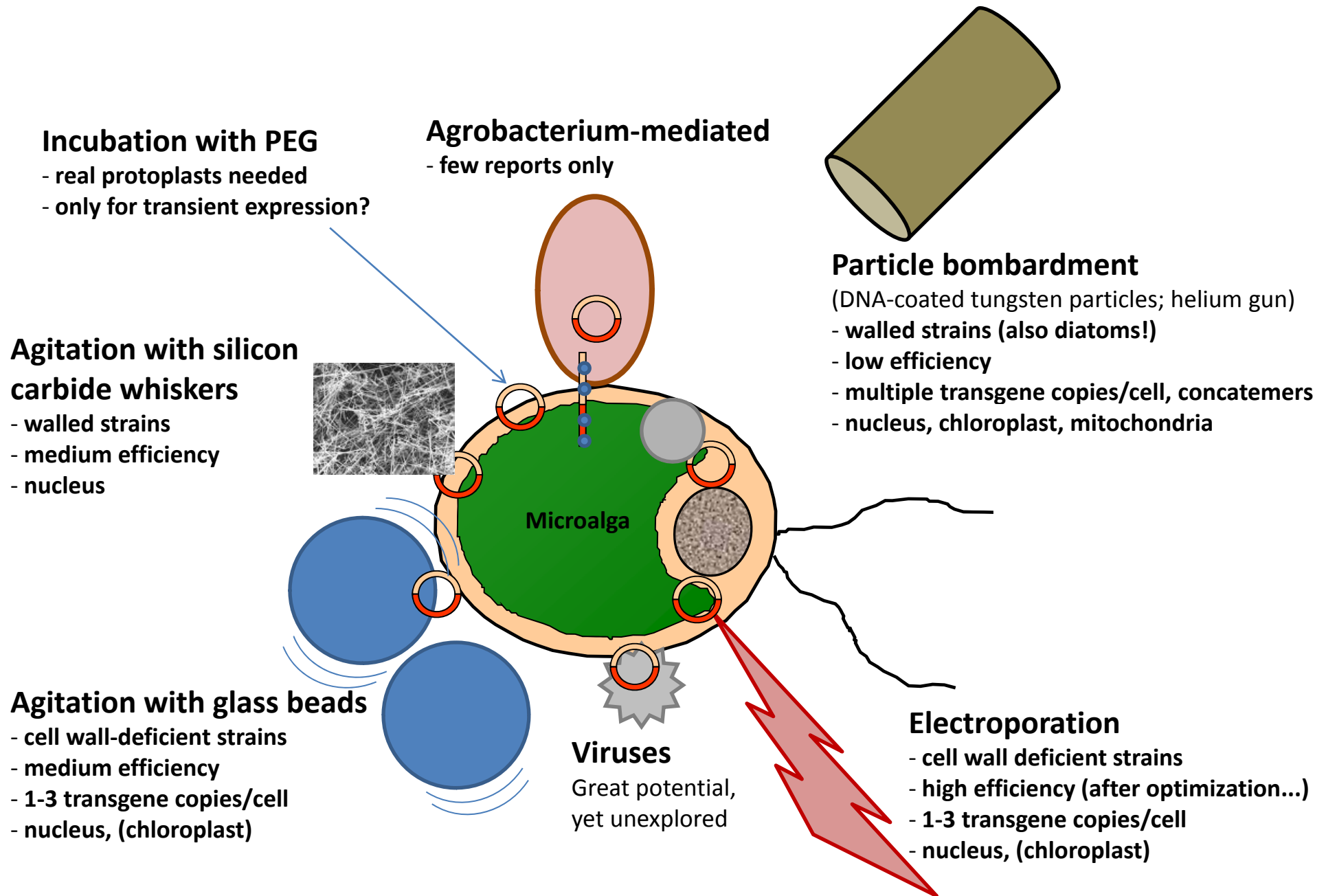
*Chlamydomonas reinhardtii* (*Volvox carteri*)

- Strategies for targeted knock-out / knock-down in *Chlamydomonas reinhardtii*
  - knock-out libraries
  - (inducible) amiRNAs
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# Strategie for microalgal transformation

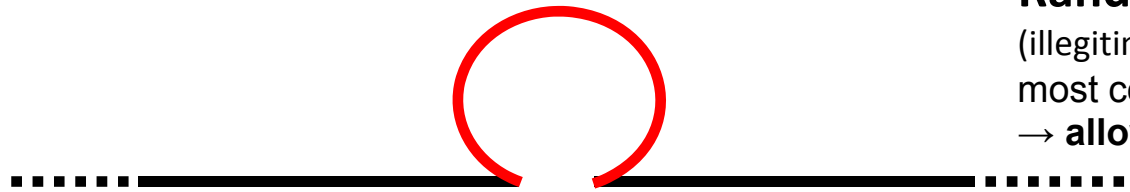


# Transformation techniques for microalgae





# Transgene Integration



## Random integration

(illegitimate recombination, non-homologous recombination)  
most common in algal nuclei  
→ **allows generation of knock-out libraries**



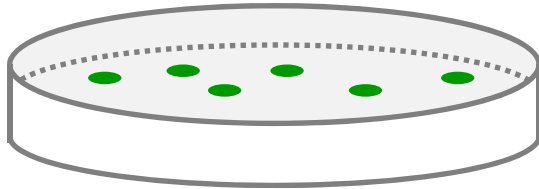
## Homologous recombination

usually rare in nuclei, most common in organelles  
→ **allows targeted knock-out / gene manipulation**



## Transient, no stable integration

# Selection



## Complementation of auxotrophic strains (recessive selectable markers)

- Generate strains by random mutagenesis (e.g. UV) that cannot synthesize essential metabolite
  - Grow cells by exogenously supplementing essential metabolite
  - Transform cells with wild-type gene
  - Select for transformants by plating cells on medium lacking essential metabolite
- (e.g.: defects in nitrate reductase, grow cells on ammonium, transform with *NR* gene, select on nitrate  
defect in argininosuccinate lyase, grow with arginine, transform with *ARG7* gene select on medium lacking arginine)

## Introducing resistance gene to antibiotic or herbicide (dominant selectable markers)

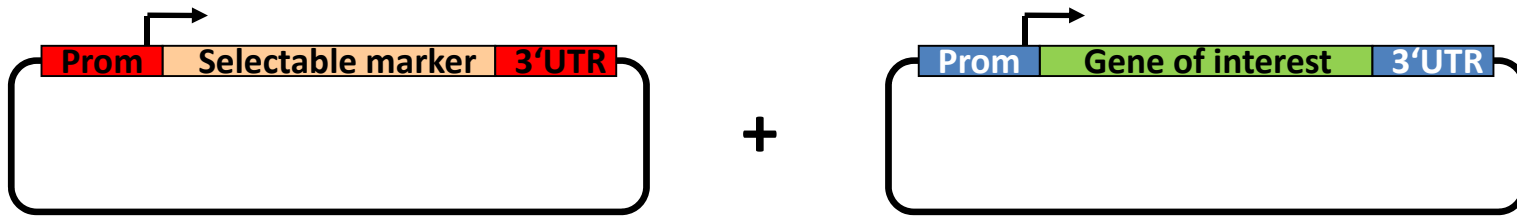
- Transform cells with resistance gene (usually under control of homologous promoters)
- Select for transformants by plating cells on medium containing antibiotic / herbicide

(e.g.:

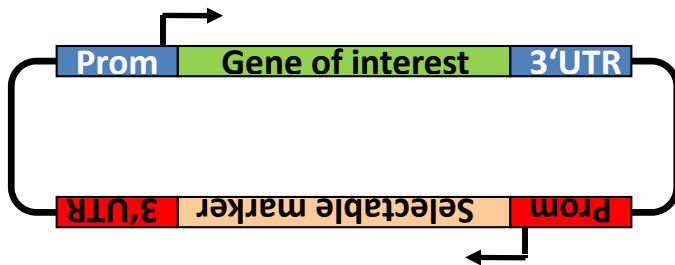
<i>ble</i>	resistance to	phleomycin antibiotics (zeocin)
<i>aadA</i>		spectinomycin, streptomycin
<i>aphVIII</i>		paromomycin
<i>aph7''</i>		hygromycin B
<i>nat</i>		nourseothricin
<i>nptII</i>		kanamycin
<i>bar</i>		phosphinothricin (herbicide)

# Transgene Expression

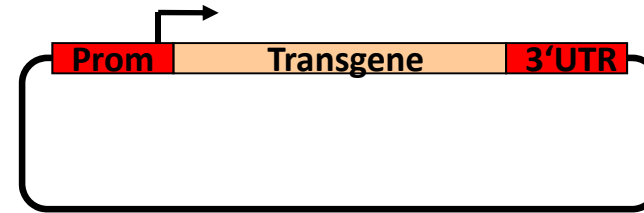
**Co-transformation** with 2 separate plasmids for selectable marker and gene of interest  
(20-50% of transformants contain construct with gene of interest)



**Co-transformation** with construct for selectable marker and gene of interest on one plasmid  
(~80% of transformants contain construct with gene of interest)



# Transgene Expression



## Promoter

- Should be homologous promoter (from the algal species to be manipulated)
- Strong constitutive promoter (nuclear: *RBCS*, *FCP*; chloroplast: *psbA*)
- Inducible promoter (nuclear: nitrate reductase, carbonic anhydrase, cytochrome  $c_6$ )

## Transgene

- Consider codon usage!  
(e.g. *Chlamydomonas* nuclear genes: G/C-rich, bias towards G/C in 3rd position  
chloroplast genes: A/T-rich)
- Introns (if genes in target alga are intron-rich, like in *Chlamydomonas reinhardtii*)

## 3'UTR

- Should be homologous (from the algal species to be manipulated)
- Ensure presence of polyadenylation signal (transcript stability!)

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- Diatoms

- Thalassiosira pseudonana*

- Phaeodactylum tricornutum*

- Rhodophytes (red algae)

- Cyanidioschyzon merolae*

- Chlorophytes

- Chlorella*

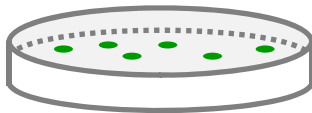
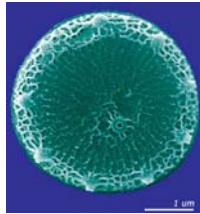
- Haematococcus pluvialis*

- Dunaliella salina*

- Chlamydomonas reinhardtii* (*Volvox carteri*)

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# *Thalassiosira pseudonana*



## Transformation

Particle bombardment

## Transgene Integration

Stable, random integration

## Selection

*nat* gene (nourseothricin resistance)

## Transgene Expression

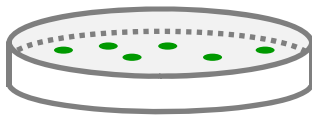
*LHCF9* promoter (constitutive)

*NR* promoter (inducible)

# *Phaeodactylum tricornutum*

Second best established algal genetic system  
**But...**

- Diploid in vegetative phase
- Particle bombardment: only low frequency
- No homologous recombination



## Transformation

Particle bombardment

## Transgene Integration

Stable integration

## Selection

*ble* gene (resistance to phleomycin antibiotics)

*nat* gene (nourseothricin resistance)

*nptII* gene (kanamycin resistance)

*sat-1* gene (streptothricin resistance)

*cat* (chloramphenicol resistance)

## Transgene Expression

### Promoters

Fucoxanthin chl-binding protein (*fcpA*)

### Transgenes

*GFP*

Luciferase

*uidA*

*Human Glut1*

*Chlorella HUP1*

*etc....*

**Inverted Repeat: RNAi!**

Apt, K. E. et al. (1996) Mol. Gen. Genet. 252:572–579.

Zaslavskaja, L. A. et al. (2000) J. Phycol. 36:379–386.

Zaslavskaja, L. A. et al. (2001) Science 292:2073–2075.

Sakaguchi, T., Nakajima, K., & Matsuda, Y. (2011) *Plant Physiol.*, in press.

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*Haematococcus pluvialis*

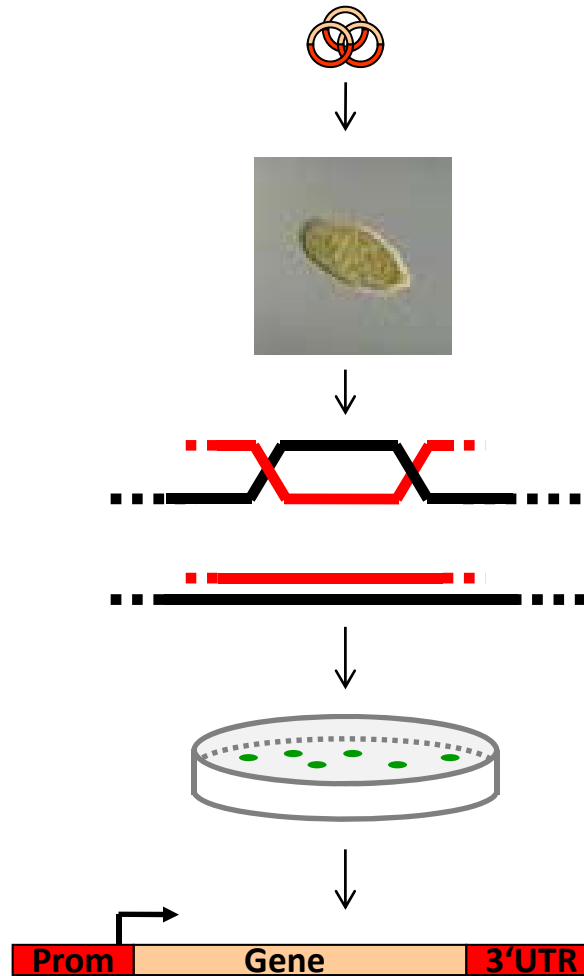
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# Cyanidioschyzon merolae



## Transformation

Electroporation, PEG

## Transgene Integration

Homologous recombination

Transient

## Selection

GFP fluorescence

*URA3*, Complementation of uracil-requiring mutants  
(selected initially for spontaneous mutants resistant to 5-fluoroorotic acid)

## Transgene Expression

### Promoters

*URA3* promoter and  
Catalase promoter  
*βTUB* promoter

### Transgenes

*URA3* wild-type gene  
antisense against catalase  
*GFP*  
HA-tagged  $\beta$ -tubulin

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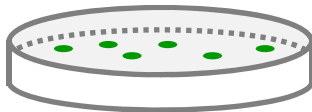
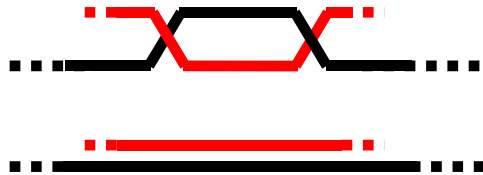
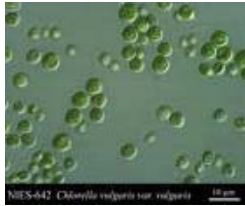
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# **Chlorella** (*C. ellipsoidea*, *C. saccharophila*, *C. sorokiniana*, *C. vulgaris*, *C. kessleri*)



## **Transformation**

Particle bombardement, PEG, electroporation

## **Transgene Integration**

Homologous recombination

Transient

## **Selection**

none (...)

Complementation of *NR* gene mutant with wild-type *NR* gene

*hpt* (hygromycin resistance)

*nptII* (kanamycin resistance)

## **Transgene Expression**

### **Promoters**

plant CaMV-35S promoter

*Chlorella* virus promoter

*Chlamydomonas* *RBCS2*

### **Transgenes**

*Renilla* luciferase

*E. coli* *GUS*

human growth hormone

neutrophil peptide-1

flounder fish growth hormone

mosquito ovary peptide hormone

Jarvis, E. E. & Brown, L. M. (1991) *Curr. Genet.* 19:317–21.

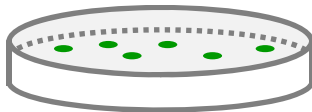
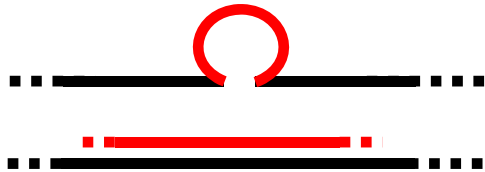
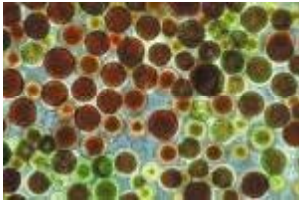
Dawson, H. N. et al. (1997) *Curr. Microbiol.* 35:356–62.

Hawkins, R. L. & Nakamura, M. (1999) *Curr. Microbiol.* 38:335–41.

Borovsky, D. (2003) *J. Exp. Biol.* 206:3869–75.

Kim, D.-H. et al. (2002) *Mar. Biotechnol.* 4:63–73.

# Haematococcus pluvialis



## Transformation

Particle bombardment, electroporation

## Transgene Integration

Stable integration (concatemers?)

Transient

## Selection

none (...)

Modified phytoene desaturase gene (norflurazon resistance)

## Transgene Expression

### Promoters

Viral SV40 promoter

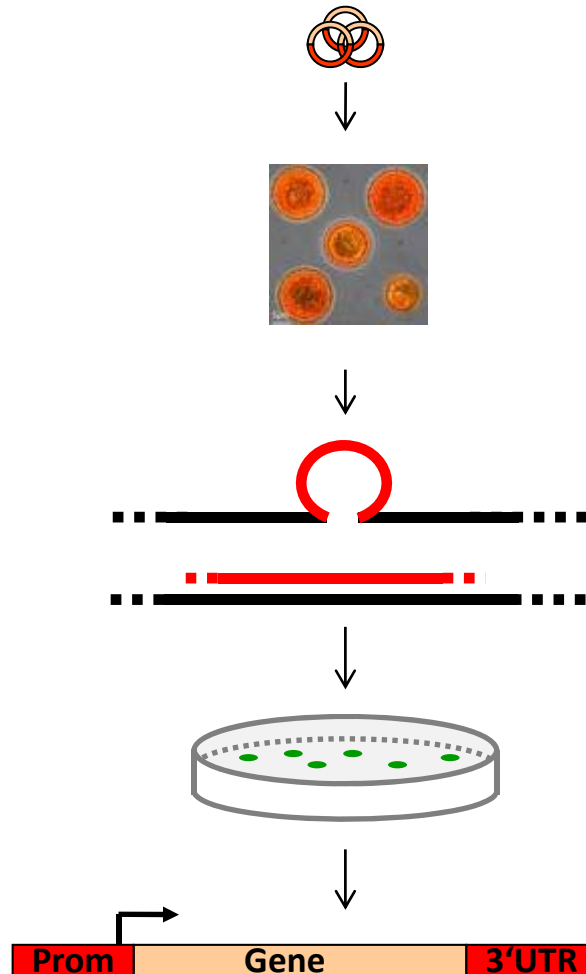
Phytoene desaturase promoter

### Transgenes

*E. coli lacZ*

and modified gene

# *Dunaliella salina*



## Transformation

Particle bombardment, electroporation, glass bead agitation

## Transgene Integration

Stable integration

Transient

## Selection

none (...)

*bar* gene (resistance to herbicide phosphinothricin)

*ble* gene (resistance to phleomycin antibiotics)

(Transient) complementation of NR mutant

## Transgene Expression

### Promoters

Maize ubiquitin promoters

CaMV-35S promoter

Chlamydomonas *RBCS2* promoter

Actin promoter

Carbonic anhydrase (salt-induced)

Nitrate reductase (NO<sub>3</sub>-induced)

### Transgenes

*EGFP*

*GUS*

Nitrate reductase

Geng DG et al. (2002) High Tech. Lett. 12:35–39.

Sun Y et al. (2005) Mol. Biotechnol. 30:185–192.

Tan CP et al. (2005) J. Microbiol. 43:361–365.

Geng DG et al. (2004) Acta Bot. Sin. 46:342–346.

Jiang GZ et al. (2005) Acta Gen. Sin. 32:424–433.

Li, J. et al. (2007) Gene 403, 132–142.

Li J et al. (2008) J. Appl. Phycol. 20:137–145.

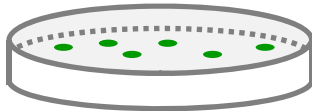
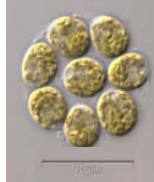
Li, J. et al. (2010) Mol. Biol. Rep. 37, 1143–1154.

# *Chlamydomonas reinhardtii*

Best established algal system for cp transformation

But...

- High level expression only in *psbA* mutant



## Transformation

Particle bombardment (agitation with glass beads)

## Transgene Integration

Homologous recombination

## Selection

Complementation of cp gene mutants (e.g. *atpB* in FUD50)  
*aadA* gene (spectinomycin, streptomycin resistance)  
*aphA-6* (kanamycin resistance)

## Transgene Expression

### Promoters

*psbA*

*atpA*

*rbcL*

*psbC*

*etc.*

### Transgenes

*GFP*

*Luciferase*

*uidA*

HSV8-scFv (single-chain antibody)

human therapeutic proteins

*etc.*

Boynton, J. E. et al. (1988) *Science* 240:1534–1538.

Goldschmidt-Clermont, M. (1991) *NAR* 19, 4083-4089.

Bateman, J. M. & Purton, S. (2000) *Mol. Gen. Gen.* 263:404–410.

Mayfield S.P et al. (2003) *PNAS* 100:438–442.

Mayfield, S. & Schultz, J. (2004) *Plant J.* 37:449–458.

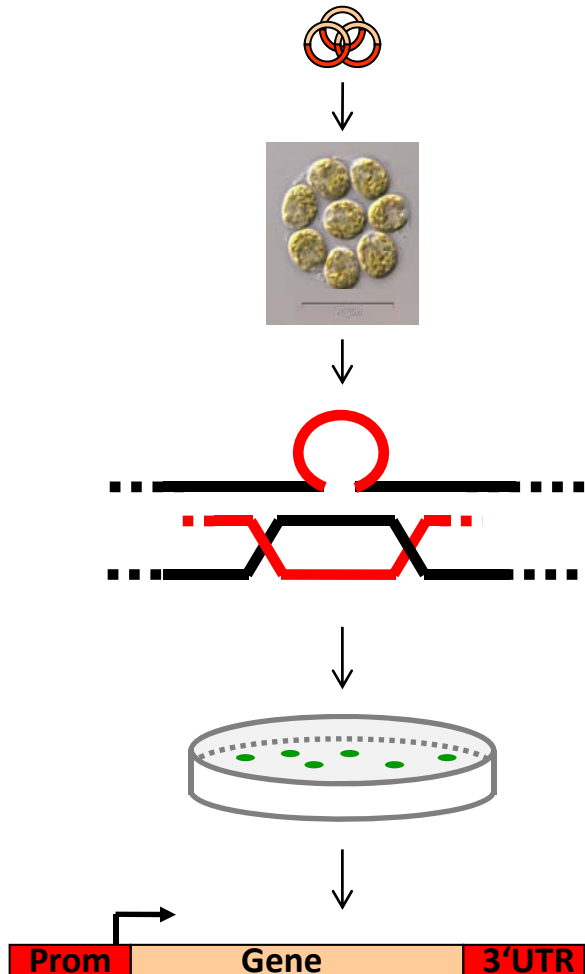
Rasala, B. A. et al. (2010) *Plant Biotechnol. J.* 8:1–15.

# *Chlamydomonas reinhardtii*

Best established algal system for nuclear transformation

But...

- No efficient homologous recombination
- No reliable high-level nuclear transgene expression



## Transformation

Particle bombardment, electroporation, agitation with glass beads or silicon carbide whiskers, agrobacterium

## Transgene Integration

Stable integration

Homologous recombination (ssDNA)

## Selection

Complementation of auxotrophic strains (*arg7*, *nit1*, *atpC*, ...)

*ble* gene (resistance to phleomycin antibiotics)

*aphVIII* gene (resistance to paromomycin, kanamycin and neomycin)

*aph7''* gene (resistance to hygromycin B)

*aadA* gene (spectinomycin, streptomycin resistance)

## Transgene Expression

### Promoters

***HSP70A-RBCS2***

*PSAD*

$\beta_2$ *TUB*

*NIT1* (inducible)

*CYC6* (inducible)

etc...

### Transgenes

GFP

Luciferase (*Gaussia*, *Renilla*)

*Chlorella HUP1*

antisense, Inverted Repeat

amiRNA!

etc...

- Debuchy, R., et al. (1989) *Embo J* 8, 2803-2809.  
Fernandez, E. et al. (1989) *PNAS* 86, 6449-6453.  
Kindle, K. L. (1990) *PNAS* 87, 1228-1232.  
Stevens, D. R. et al. (1996) *Mol Gen Genet* 251, 23-30.  
Fuhrmann, M. et al. (1999) *Plant J* 19, 353-361.  
Schroda, M. et al. (2000) *Plant J* 21, 121-131.  
Fuhrmann, M. et al. (2001) *J Cell Sci* 114, 3857-3863.  
Sizova, I., et al. (2001) *Gene* 277, 221-229.  
Molnar, A. et al. (2009) *Plant J* 58, 165-174.  
Schmollinger, S. et al. (2010) *Curr Genet* 56, 383-389.

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- Strategy for microalgal transfection (transformation)
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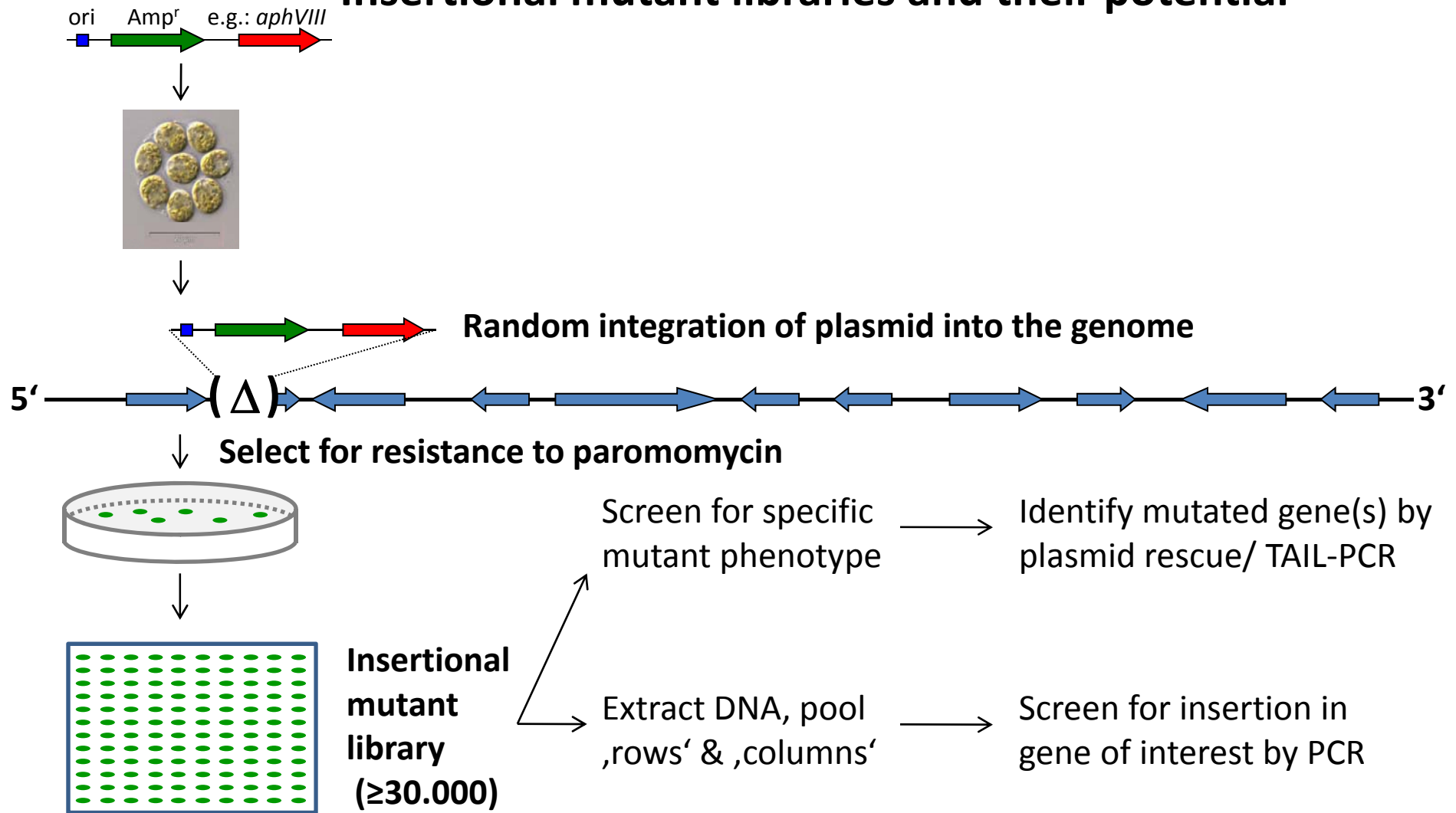
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# Insertional mutant libraries and their potential



## Challenges

- High transformation efficiency needed
- Screen for phenotype: vegetative cells need to be haploid
- Maintenance of thousands of transformants

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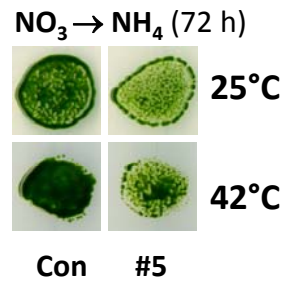
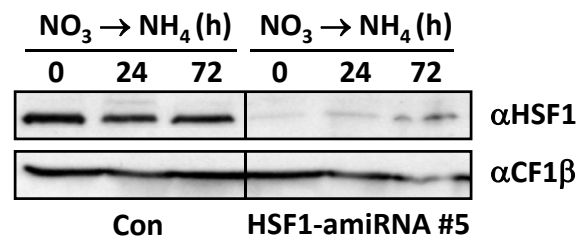
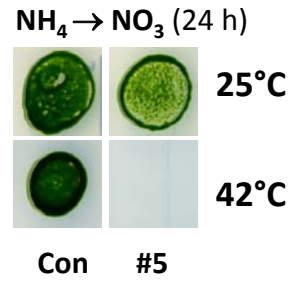
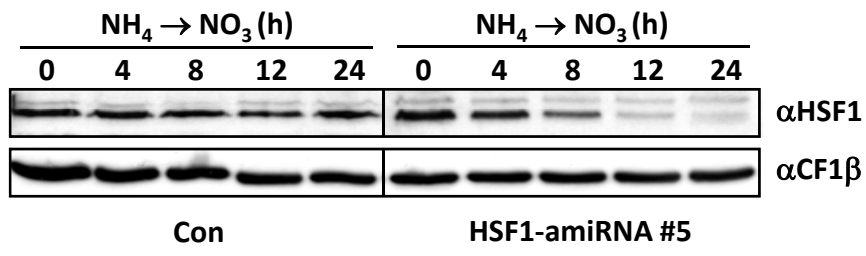
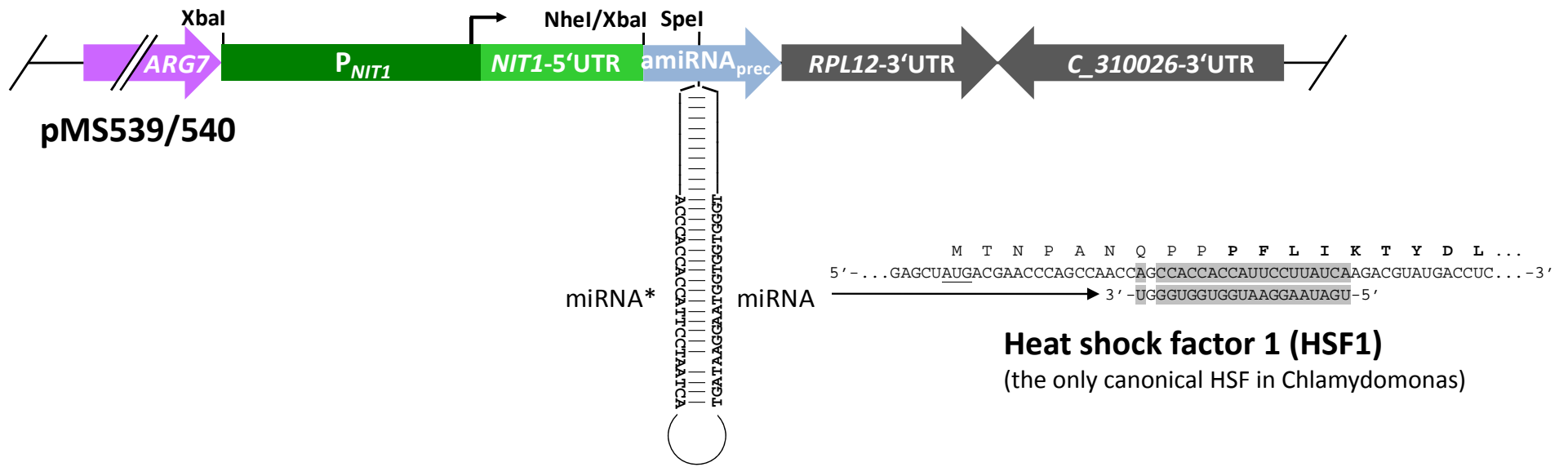
*Haematococcus pluvialis*

*Dunaliella salina*

*Chlamydomonas reinhardtii* (*Volvox carteri*)

- Strategies for targeted knock-out / knock-down in *Chlamydomonas reinhardtii*
  - knock-out libraries
  - (inducible) amiRNAs
- Overcoming problems in gene expression by targeted chromatin remodeling
- Summary & Outlook

# Establishment of a conditional amiRNA system for *Chlamydomonas* using heat shock factor 1 (HSF1) as target



# Transgenic Microalgae – Problems and Perspectives

- Rationale – why do we want to genetically manipulate microalgae?
- Strategy for microalgal transfection (transformation)
- Overview to transformed microalgae

Diatoms

*Thalassiosira pseudonana*

*Phaeodactylum tricornutum*

Rhodophytes (red algae)

*Cyanidioschyzon merolae*

Chlorophytes

*Chlorella*

*Haematococcus pluviales*

*Dunaliella salina*

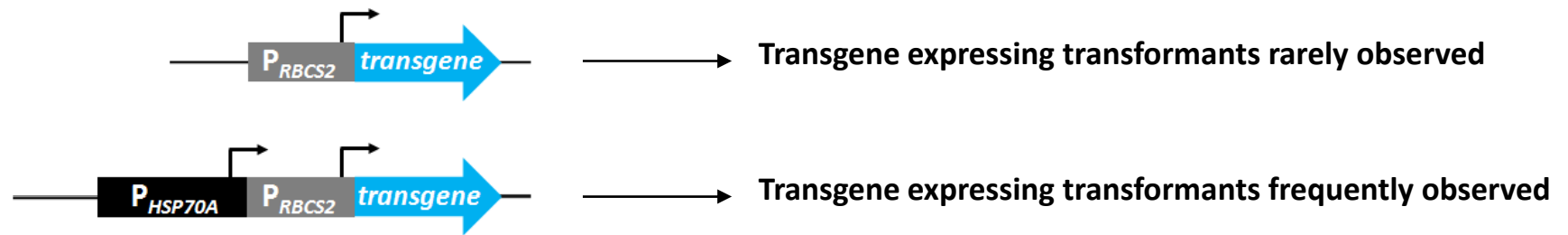
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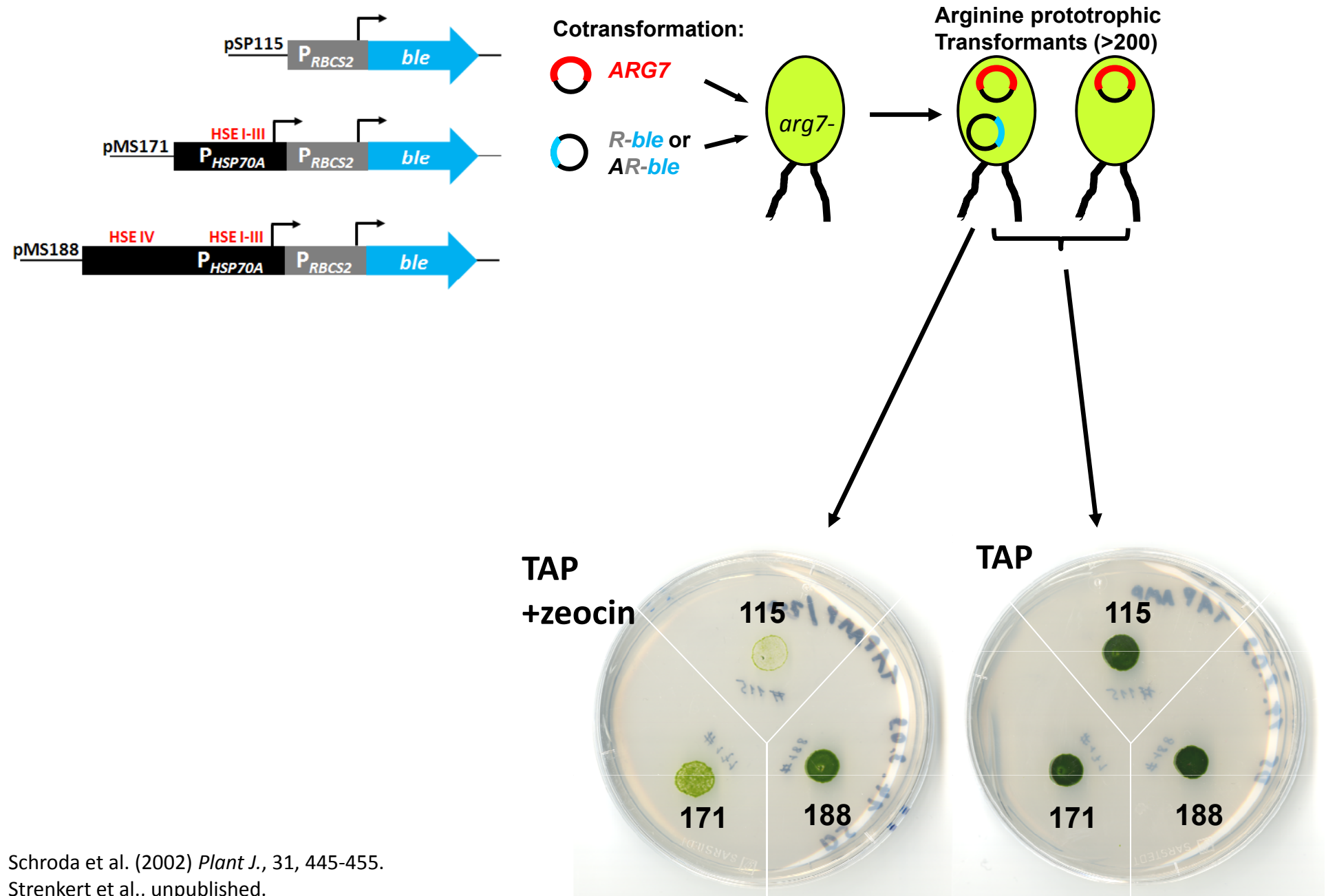
## The *HSP70A* promoter improves transgene expression



# The *HSP70A* promoter improves transgene expression

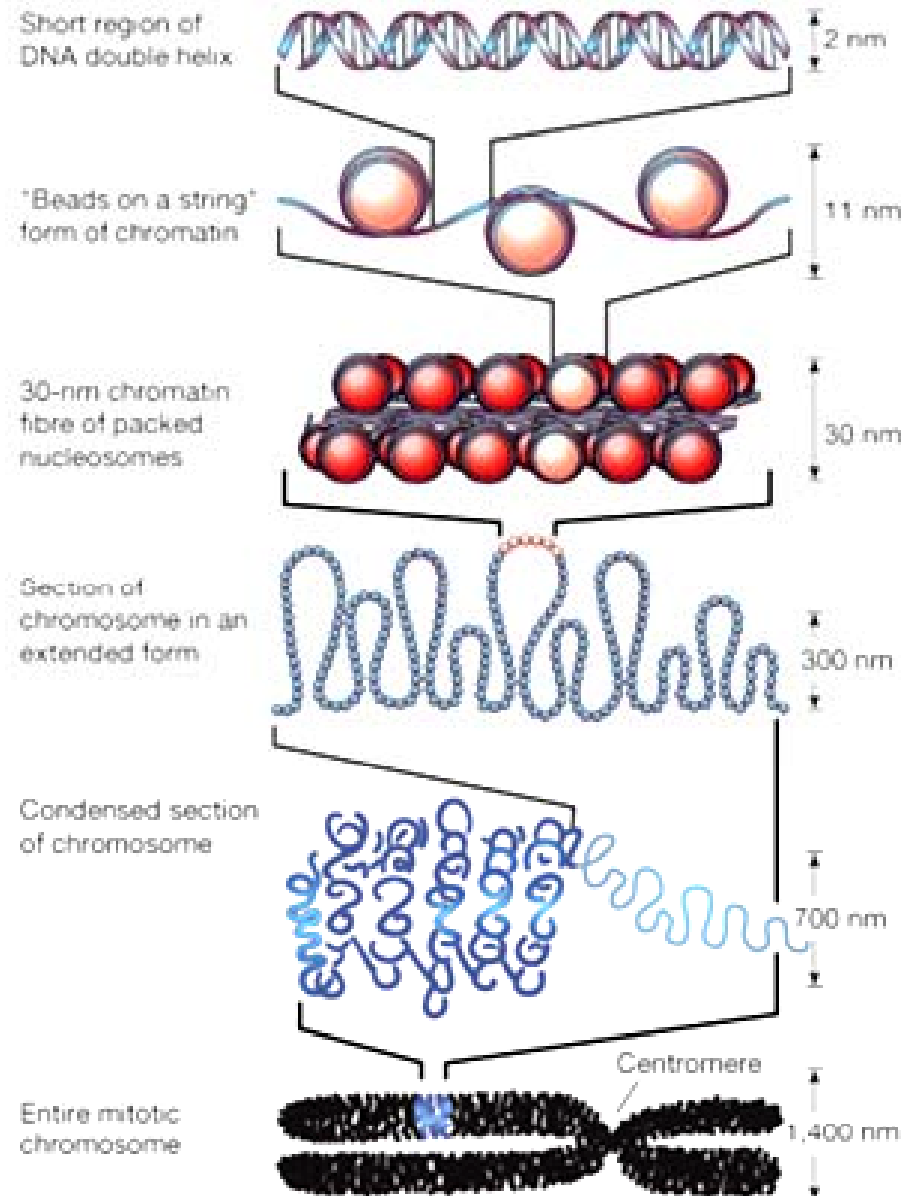


# The *HSP70A* promoter counteracts silencing of downstream promoters



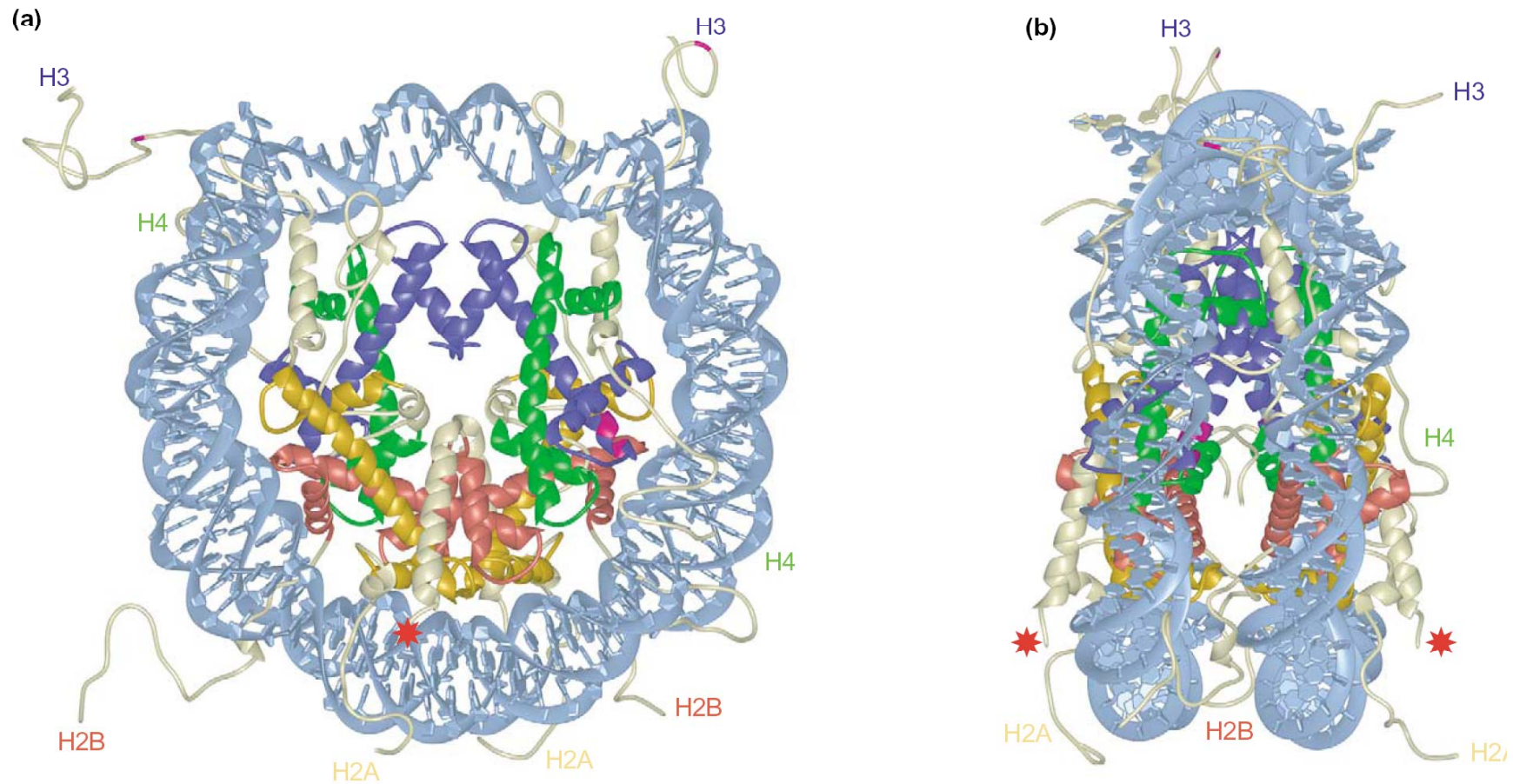
Schroda et al. (2002) *Plant J.*, 31, 445-455.  
Strenkert et al., unpublished.

# DNA is organized into chromatin



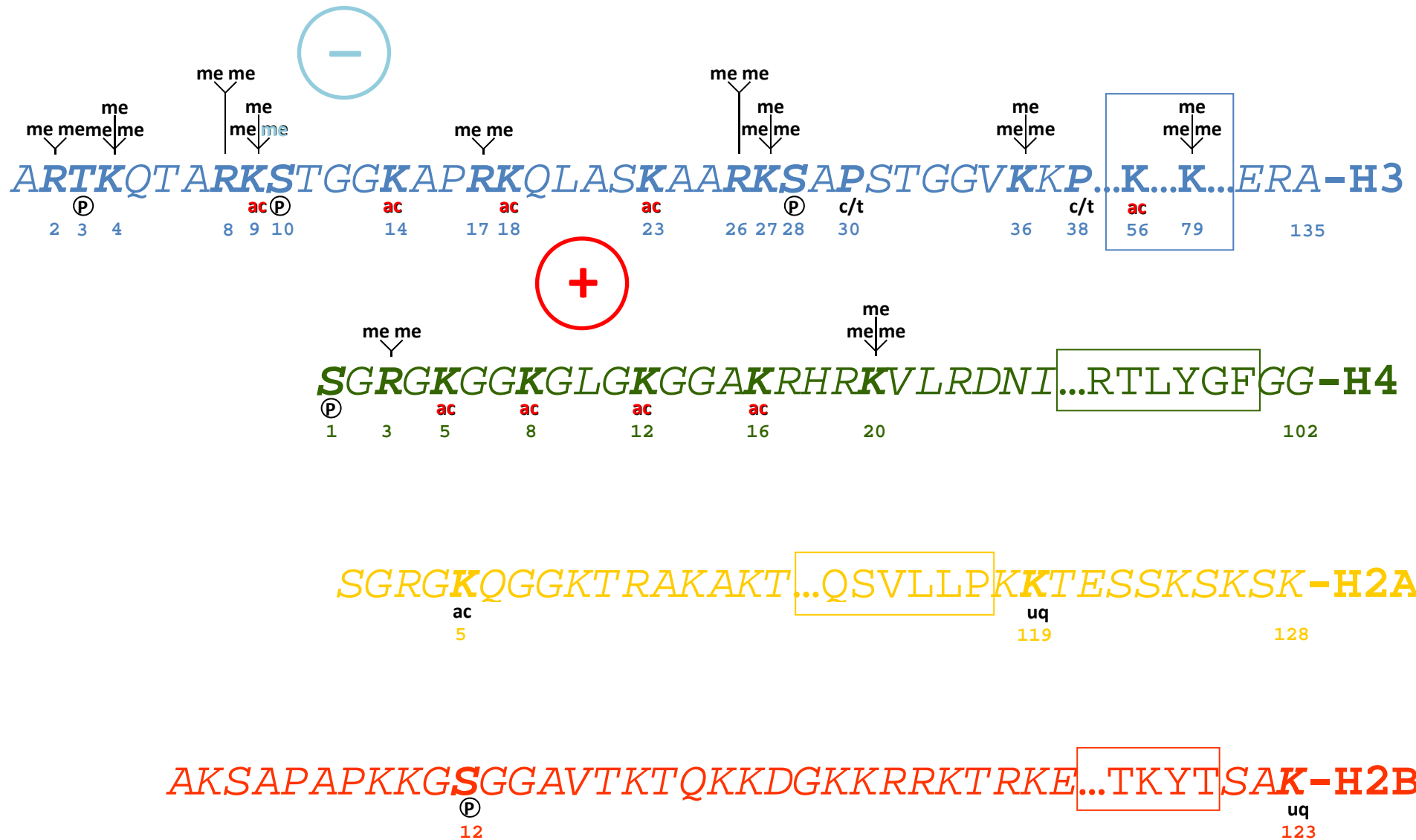


# Structure of a nucleosome

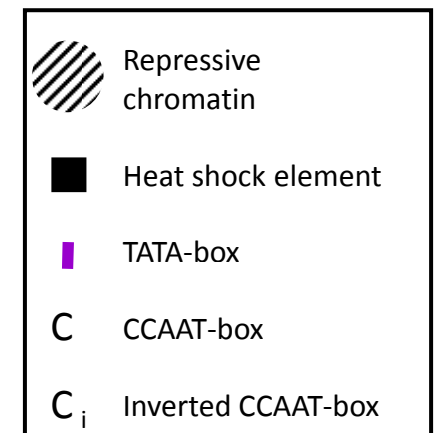
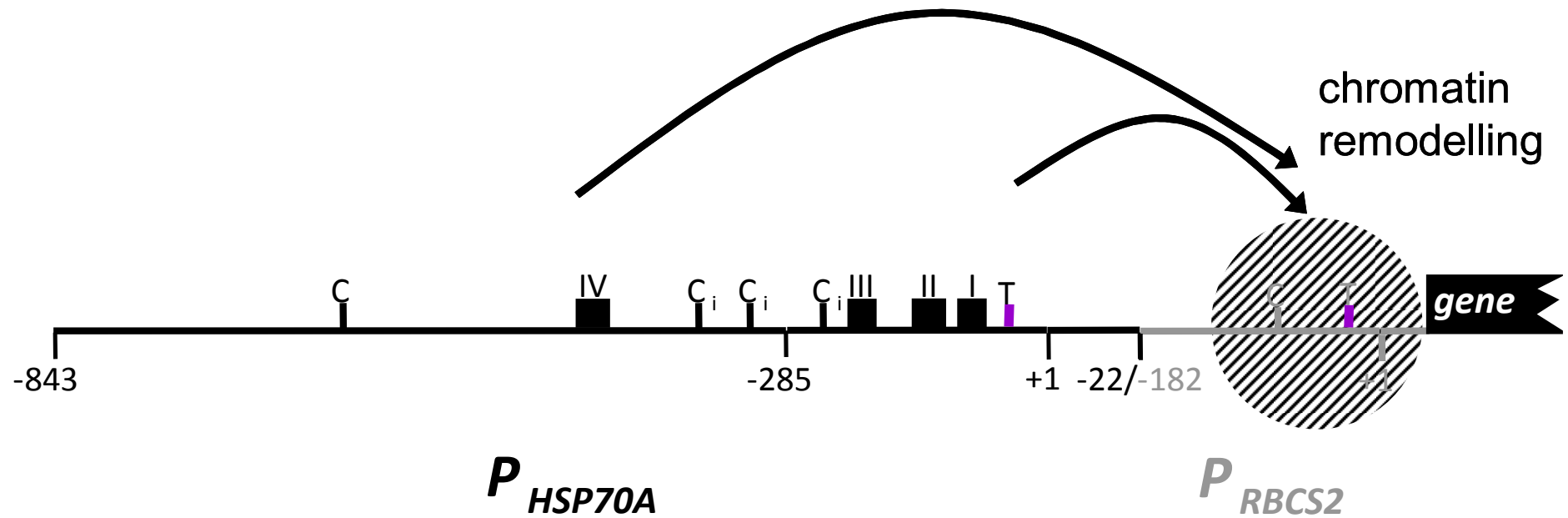


- 147 bp DNA wrapped around octamer consisting of dimers of histones **H2A**, **H2B**, **H3** and **H4**

# The Histone code



# Model explaining how the *HSP70A* promoter may activate transgenes



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# Summary & Outlook

## Chloroplast

- Significant transgene expression only in *Chlamydomonas psbA* mutants
- needs to be established in other algal systems (started in *Euglena gracilis* & *Porphyridium sp.*)

## Nucleus

### Insertional mutant libraries

- Only established for *Chlamydomonas*
- higher transformation efficiencies needed for other systems

### Homologous recombination

- encouraging results with ssDNA in *Chlamydomonas*
- perhaps easy in *C. merolae* and *Chlorella*?!  
→ Much more research needed

### RNA silencing

- Well established for *Chlamydomonas* (antisense, IR, amiRNA)
- Encouraging: first reports on successful IR in *P. tricornutum*; antisense in *C. merolae*  
→ more research needed; micro-RNAs present in other microalgae?

### Transgene expression

- Transient expression appears to work well in *C. merolae*, *Chlorella*, *D. salina* (*N. occulata*?!)
- Expression of stable transgenes appears fine in *P. tricornutum*, bad in *Chlamydomonas*  
→ Solvable when epigenetic gene silencing mechanisms understood?