

Traditional Algal Transformation Techniques

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Microalgae and cyanobacteria are photosynthetic microbes that can be grown with the simple inputs of water, carbon dioxide, (sun)light, and trace elements.

microalgae

transformation

cyanobacteria

DNA

1. Introduction

Microalgae and cyanobacteria are interesting study organisms capable of photosynthetic growth on carbon dioxide (CO₂) as a sole carbon source. These organisms naturally synthesize different metabolites such as pigments, oils and lipids, sterols, starches, polysaccharides, and halogenated compounds, and algae are also amenable to contained, scalable growth in photobioreactors driven by (sun)light energy [\[1\]\[2\]](#). Genetic engineering of algae and cyanobacteria in the era of synthetic biology holds the promise of tailored production of novel and customized metabolites using sustainable waste CO₂ as a feedstock. However, algae are a diverse and polyphyletic group of organisms which do not share close evolutionary relatedness and exhibit incredible variability in their genomes [\[3\]](#). These features have hindered intensive molecular tool development except in a handful of model species, and their broader application to biotechnology has been slower compared to other hosts such as bacteria, yeast, plant, and mammalian cells [\[2\]\[4\]](#).

The genomic diversity of algae necessitates customized molecular tools that work with genetic architecture of a specific host. Once these tools are produced by piecemeal cloning or complete DNA synthesis, the reliable introduction of foreign genetic material into cells is an essential prerequisite to biotechnological concepts [\[5\]\[6\]\[7\]](#). DNA delivery methods result in stable chromosomal integration or episomal/plasmid extrachromosomal replication of foreign transgene expression elements [\[7\]\[8\]\[9\]](#). Methodologies of DNA delivery and cell membrane/wall permeabilization vary according to the host organism and target cellular compartment (organellar or nuclear). For example, some cyanobacteria will natively take up DNA from their environment without the need for manipulation, while many eukaryotic algae maintain cell walls that necessitate more aggressive delivery methods [\[10\]\[11\]](#).

Foreign DNA delivery can be achieved by mechanical agitation, surfactant permeabilization, electroporation, particle bombardment, and bacterial DNA transfer (conjugation or *Agrobacterium tumefaciens*-mediated) [\[9\]\[12\]\[13\]\[14\]\[15\]](#). In addition to these classical methods, emerging techniques used in other cell systems that have not been widely applied to algae may potentially improve DNA delivery ([Figure 1](#)). Strategies based on cell penetrating peptides (CPP), polymers, metal-organic frameworks, nanoparticles, and liposomes have all been demonstrated in

non-algal hosts [16][17][18]. This review examines recent progress in gene delivery methodologies and discusses their technical aspects, advantages, limitations, and potential in the context of algal biotechnology.

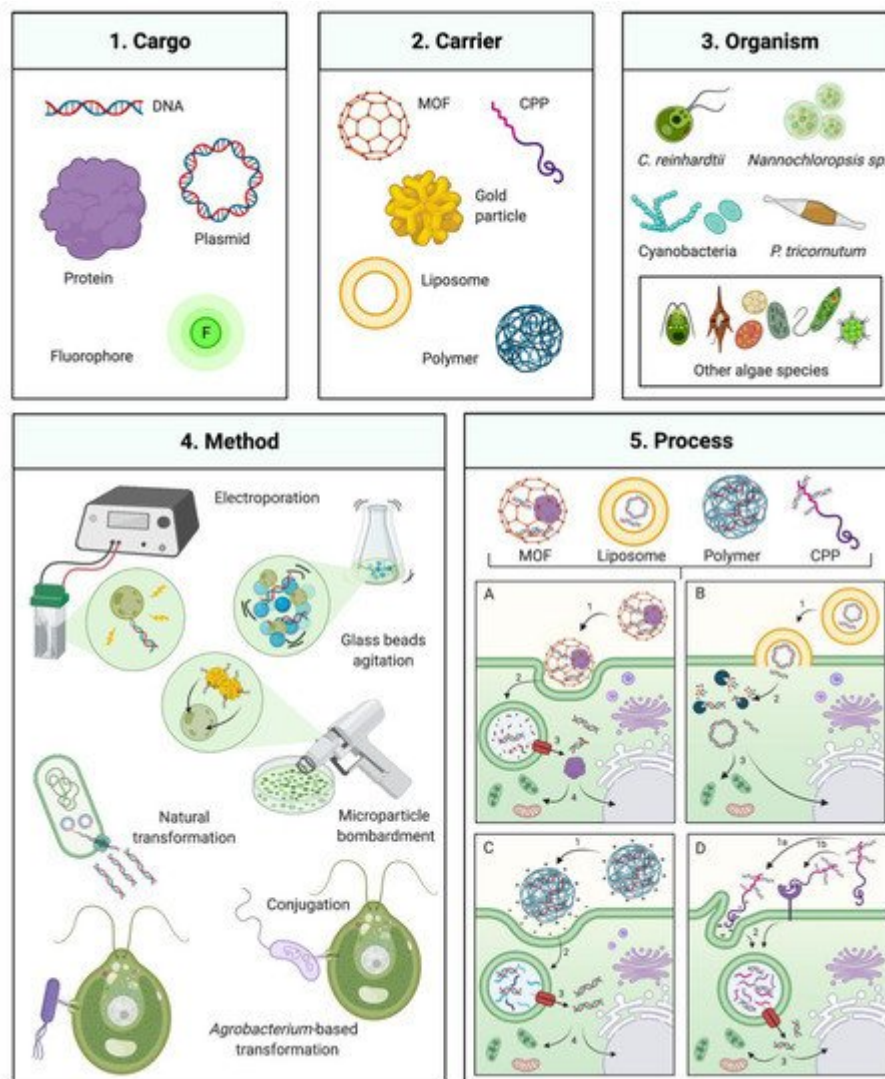


Figure 1. Gene delivery technologies for microalgae engineering. Overview of several available transformation technologies which have been or could be applied to algal engineering. (1) and (2) Several carriers that can mediate transformation of a DNA, protein, or chemical cargo (left panel) MOF: metal-organic framework, CPP: Cell-penetrating peptide. (3) Algal species which transformation has been commonly employed are depicted on the upper right: *C. reinhardtii*, *Nannochloropsis sp.*, and *P. tricornutum*, including various cyanobacteria and other algal species as described in the manuscript. (4) Classical transformation methods are presented in the bottom left panel. (5) Process of transformation mediated by different DNA carriers and their interaction with the cell for delivery of the respective cargo. (5A) MOF-DNA, (5A1) recognition and cell uptake by endocytosis, (5A2) internalization and fusion with phagolysosome, (5A3) phagolysosome escape, (5A4) transport to nucleus, chloroplast or mitochondria; (5B) Liposome-DNA, (5B1) liposome integration with cell membrane, (5B2) cargo (DNA or protein) exposed in cytoplasm and potential enzymatic degradation, (5B3) intact cargo transport to nucleus, mitochondria or chloroplast; (5C) Polymer-DNA, (5C1) cell recognition of polymer nanoparticle, charge interaction and cell uptake by endocytosis, (5C2) internalization and fusion with phagolysosome, (5C3) phagolysosome escape, (5C4) transport to nucleus, chloroplast or mitochondria; (5D) CPP, (5D1) recognition and cell uptake by endocytosis, (5D2) internalization and fusion with phagolysosome, (5D3) phagolysosome escape, (5D4) transport to nucleus, chloroplast or mitochondria.

transport to nucleus, chloroplast or mitochondria; (5D) CPP-DNA, 1 cell recognition of CPP, by charge interaction (5D1a) or receptor recognition (5D1b) and cell uptake by endocytosis, (5D2) internalization and fusion with phagolysosome, (5D3) phagolysosome escape, (5D4) Transport to nucleus, chloroplast or mitochondria. Created with BioRender.com.

2. Traditional Algal Transformation Techniques

2.1. Agitation of Cells in the Presence of DNA and Non-Ionic Surfactants

The agitation of algal cells in the presence of glass beads, polyethylene glycol (PEG), and foreign DNA has become a standard protocol for the delivery of foreign DNA into the model green microalgae *C. reinhardtii* and other species as *Cyanidioschyzon merolae*, *Chlorella vulgaris*, and *Dunaliella salina* [14][19][20][21]. This protocol has been adopted for nuclear and chloroplast transformation and can be applied to cell wall deficient cell lines or those with a cell wall when chemo/enzymatic removal is performed before transformation (as with *Chlorella* strains) [13][14][15][22][23][24]. Agitation-based transformation is achieved by agitation (vortexing) of cells in the presence of a physical agitator (glass beads), a non-ionic surfactant (i.e., polyethylene glycol (PEG)), and DNA [14][25]. Transformation efficiency depends on many factors such as cell size, presence or absence of a cell wall, duration of agitation, velocity, the concentration of surfactant, and the use of linear or circular DNA [26]. The frequency of DNA integration into the *C. reinhardtii* genome with glass bead agitation has been reported at $\sim 10^3$ transformants μg of DNA⁻¹ with $\sim 10^8$ cells mL⁻¹ starting cell concentration [14]. Reports have shown that numerous other non-reactive materials, such as silicon carbide whiskers can also serve this purpose [27]. Agitation-based transformation protocols are advantageous as they do not require specialized equipment, and are inexpensive and relatively fast [14][15].

A significant limitation of the technique is the requirement of cell wall removal, which can be achieved with the autolysin protein of *C. reinhardtii* for itself, or other enzyme cocktails (e.g., cellulase, macerage, pectinase, and hemicellulose) for other hosts [14][19][28][29][30][31][32]. Reports have shown that cell-wall free protoplasts of *Chlorella ellipsoidea* and the naturally cell wall deficient *C. merolae* have been transformed using DNA and PEG without mechanical agitation, suggesting that surfactant mediated permeabilization is sufficient to enable DNA uptake [33]. However, the main drawback of agitation-based transformation techniques is the low transformation efficiency [34]. This has led to development of other more efficient strategies.

2.2. Electroporation

DNA transformation by electroporation is as an alternative method in numerous algae and pioneered in *C. reinhardtii* [34]. This technique uses electrodes to generate voltage differential across the cell membrane, temporarily disturbing the phospholipid bilayer, allowing molecules to pass into the cell [35][36]. Transformation efficiencies from electroporation vary according to factors such as field-strength, pulse length, ionic strength of medium composition, temperature, cell membrane characteristics, the species used, its molecular toolkit, and the presence of cell-wall [36][37]. Electroporation can increase transformation efficiency up to 100-fold over agitation and

is not affected by the cell wall. Fewer sequence deletions have been observed at the genomic insertion sites than those seen from agitation transformation [34][37][38]. Electroporation has been shown to enable the transformation of cell-wall containing algae and cyanobacteria such as *Monoraphidium neglectum*, *Nannochloropsis* sp., *Phaeodactylum tricornutum*, *C. reinhardtii*, *Anabaena*, and *Nostoc punctiforme* [34][35][39][40][41][42][43][44]. Electroporation is a useful technique for delivering DNA, RNA, proteins, nucleotides, and dyes into cells [34][35][45]. Electroporation is a rapid protocol and requires equipment which is often present in microbial laboratories to transform bacterial cells [37][46].

Electroporation efficiencies were increased by considering the specific mitotic phase of *C. reinhardtii* growth [47]. Synchronization of *C. reinhardtii* cultures has been shown to increase transformation efficiency after three days. *C. reinhardtii* culture synchronization using 28 °C during the light phase and 18 °C in the dark phase was also shown to enable increased homologous recombination efficiencies specifically at 12 hours after illumination initiation [48]. Digital microfluidics (DME) have also been used to optimize electroporation protocols, using a fluid mixture of cell/DNA droplets encapsulated in biocompatible oil, and electric pulses applied from an array of microelectrodes to the droplets [49][50]. DME showed an efficiency of 2.5×10^4 of *C. reinhardtii* mutants per μg of DNA without cell wall removal with an initial concentration of 2×10^6 cells mL^{-1} algal cells, and 1 μg DNA [49]. A square-electric-pulse electroporation technique has also been reported with efficiencies of 6×10^3 *C. reinhardtii* transformants per μg of DNA with 1.5×10^7 cells mL^{-1} initial cell density 0.1 μg of DNA [47]. However, this technique has not been broadly adopted, as it requires specialized equipment. It was described in *N. limnetica*, which contains a rigid cell wall, that treatment with 10 mM lithium acetate and 3 mM dithiothreitol (DTT) before electroporation improved transformation efficiency by increasing cell wall permeability, resulting in 1.1×10^7 transformants per μg of DNA with an initial concentration of 3.3×10^9 cells mL^{-1} and 4 μg of DNA [40].

2.3. Microparticle Bombardment

Microparticle bombardment is one of the most versatile gene delivery methods due to its ability to transform the nucleus, mitochondria, or chloroplast genomes and even transform cells containing cell walls [13][51][52][53]. Microparticle bombardment is based on accelerated non-reactive metal (gold or tungsten) micro-projectiles coated with DNA being shot at and colliding with target cells [1][51]. The impact of these particles allows them to penetrate the cells and deliver foreign DNA. Transformation efficiency depends on the starting cell density, target organelle, selection efficiency, the number of DNA-coated particles, the DNA concentration on each particle, the kinetic energy of the particles, temperature, and ability of the cell to regenerate after particle damage [54].

The most common use for this technique is the transformation of chloroplast genomes, as it enables penetration of DNA through multiple membrane layers, but it has also recently been used to transform circular plasmids into the nucleus of the red alga *Porphyridium purpureum* [23][51][55]. Unlike agitation, but similar to electroporation, cell-wall presence does not affect transformation efficiency; however, cell viability can be disturbed if many micro-projectiles are used [51][56]. The efficiency of nuclear transformation in *C. reinhardtii* has been as low as 15 transformants per μg of DNA, with $\sim 10^7$ cells mL^{-1} starting cell concentration and 0.8 μg of DNA [13]. Efficiencies reported in *D. salina*, *Volvox carteri* and *P. tricornutum* are 2.5×10^{-5} , 1.7×10^2 , and 1.0×10^2 transformants per μg of DNA (0.1–0.7 μg

of DNA), respectively, with 10^5 – 10^7 cells mL⁻¹ as starting cell density [57][58][59]. However, it was reported that in *C. reinhardtii*, transformation efficiency could be improved when smaller particles are used (0.6 µm vs. 1.0 µm) [60].

Microparticle bombardment has also been described for the delivery of 24–68 kDa proteins, a technique called proteolistics. This approach is a simple physical deposition of target protein onto delivery substrate which is then used as other microprojectiles [61]. The application was shown to deliver Cas9–gRNA ribonucleoprotein (RNP) into *P. tricornutum*, where knock out of the *PtUMPS* and *PtAPT* genes led to 5-fluoroorotic acid (5-FOA) and 2-fluoroadenine (2-FA) resistance at an efficiency of $\sim 10^{-6}$ per 10^6 cells mL⁻¹ [62]. This technique was also found to be suitable for multiplexing mutations [62] and could become a powerful strategy for generating targeted knockouts in this and other diatoms/algae.

One disadvantage of bombardment is the requirement of gene-gun infrastructure as well as reagents and equipment. Transformation parameters must also be optimized for every alga and target cell compartment/genome [9][63]. One thing that must be considered is the potential post-transcriptional gene silencing (PTGS) due to the relatively high copy number of transgenes integrated into the genome [53][64].

3.2. Bacterial Conjugation

Conjugation is based on the ability of bacteria to share genetic information by exchanging plasmids through pili. This natural process can be employed using either a double or triple vector approach combining shuttle, conjugative, and helper plasmids. The shuttle and helper plasmids mediate the transfer of a conjugative plasmid between *E. coli* and the target host while the helper plasmid aids transformation by preventing vector degradation by endogenous restriction systems; it codes for DNA methylases (AvaI, AvaW, and AvaIII) that prevent target host restriction enzyme recognition [65][66]. The helper vector contains an oriT-region, *bom*-site, and *mob* genes that encode a nickase and enable conjugation [67][68]. Bacterial conjugation has been used commonly in cyanobacterial species such as *Synechococcus*, *Prochlorococcus*, *N. punctiforme*, *Anabaena*, and *Synechocystis* sp. and recently in eukaryotic algae such as *P. tricornutum*, *Thalassiosira pseudonana*, *Acutodesmus obliquus*, and *Neochloris oleoabundans* [66][68][69][70][71][72][73][74][75]. The effectiveness of transformation by conjugation is based on the capability of the algal recipient strain to integrate and maintain the vector, either in the chromosome, in endogenous plasmids, or as an episomal plasmid [65][66][74][75].

Episomes, circular plasmids which self-replicate and do not integrate into the chromosome, can be efficient and straightforward vectors to enable transgene delivery of desired genetic elements between bacteria, cyanobacteria, and eukaryotes. Episomes avoid insertions and knock-out of non-targets and replicate independently from chromosomes [76][77][78]. Yeast centromere sequences (CEN/ARS) were found to act as autosomal replicating elements that allow stable maintenance of the circular episomal plasmid in diatoms [66][79]. Factors such as the growth phase of both target and donor organisms are crucial factors for proper conjugation [80]. Episomal vectors to transform microalgae via conjugation provide an efficient means of multi-gene pathway transfer due to stable self-replication of episomal vector and minimal possibility of positional or epigenetic effects [66]. Conjugation has the advantage over electroporation, microparticle bombardment, and glass-beads agitation to deliver larger DNA

fragments [45]. The use of the conjugation-based method in *P. tricornutum*, *N. oceanica*, and *Neochloris oleoabundans* has been demonstrated to generate mutants with higher transformation efficiency compared to microparticle bombardment transformation: 500–1000 transformants/10⁸ cells/200 ng DNA (conjugation) compared to 5–25 transformants/10⁸ cells/2.5 µg DNA (bombardment) in *P. tricornutum* [74][81][82][83].

3.3. Agrobacterium-Mediated Transformation

The use of *A. tumefaciens* to transform plant cells relies on the natural infection process of the bacterium, which causes crown gall tumors on various plant species [84]. Natural tumor formation is a consequence of replicating a single-stranded copy of the transferred bacterial tumor-inducing (Ti) plasmid. The Ti plasmid and *A. tumefaciens* have been exploited since 1988 when it was found to allow transfer and permit stable integration of DNA fragments into a target higher plant genomes and has even been shown to transform mammalian cells [85][86][87]. The technique has the reported advantage of low gene rearrangements and foreign transcript silencing in plant cell lines [88][89].

A. tumefaciens-based transformation of algae has been reported in several species, although its use has not been widely adopted. Reports exist of transformation by *Agrobacterium* method in *C. reinhardtii*, *H. lacustris*, *Chlorella* sp., *Dunaliella bardawil*, *Symbiodinium* sp., *Nannochloropsis* sp. and *Parachlorella kessleri* [85][90][91][92][93][94]. However, questions remain about how the bacterium, which evolved to infect plant cells, can infect evolutionarily distant algae and whether the infection requires specific recognition machinery on the target, or whether it is target independent. The standard protocol for *Agrobacterium*-mediated transformation is to mix target cell cultures with *Agrobacterium* containing a transgene cassette of interest; the mixture is exposed with the virulence agent acetosyringone, which signals the bacterium to infect wounded plant tissue. After transformation of the target cell by the bacterium, the selection is made using an antibiotic that can be selective against the bacterium while selecting the transformed cells. Variable transformation efficiencies have been reported depending on the protocol followed [90][95][96][97]. Factors such as temperature, pH, and time of virulence gene induction have been reported to have a substantial effect on transformation efficiency [98]. This method may enable integrating multigene pathways into host genomes as the capacity of the Ti plasmid has been reported as large as 150 kbp. If stably integrated into the genome, this would vastly outperform current methods of gene delivery that exhibit random nuclease digest *en route* to the nucleus, especially in green algae [99][100][101]. However, a recent report indicated that this method is no better than electroporation for stable integration of a plasmid containing a luciferase expression cassette and a selectable marker cassette into *Parachlorella kessleri* [90]. Further investigation is required to determine if this standard plant transformation protocol can be robustly applied to enhancing multi-gene expression cassette delivery to algal hosts.

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