Starch Overproduction by Means of Algae

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Abstract This chapter provides an overview of the state of knowledge of starch production as an ultimate energy reserve in algae. It includes a survey of recent discoveries on controls that direct the metabolism of algal cells towards starch hyper-accumulation with the aim of providing starch-enriched biomass for the production of bioethanol as a biofuel of the future. We also outline basic research from the 1960s, from which the recent starch research stems, although the use of algal starch for biofuel production was not considered at that time. The principles of, and basic approaches to a directed synthesis of starch are described in both laboratory experiments and large scale-up outdoor photobioreactors.

Keywords Algae **·** Bioethanol **·** Cell cycle **·** Carbon dioxide **·** Inhibitors **·** Light intensity **·** Nutrient limitation **·** Starch **·** Temperature

List of Abbreviations

CHX Cycloheximide DW dry weight FdUrd 5-fluorodeoxyuridine PBR photobioreactor

1 Introduction

Algae represent a vast variety of photosynthetic species growing photoautotrophically using light as a source of energy for photosynthesis and fixing inorganic carbon, from which a reserve material such as starch and/or lipid is synthesized

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and then serve as an energy and carbon source for future cellular growth and reproduction.

Despite the pioneering papers from the 1960–1980s, not much attention was paid to starch research for more than twenty years. The main motivation for the recent boom in starch research is that algae can be a high-yielding source of energy-rich reserve components, which can be used for the production of biofuels (third generation biofuels) without compromising food supply chains and be a renewable alternative to fossil fuels (Nigam and Singh [2010](#page-21-0)). In general, the algae from different taxons can produce either lipids (*Nannochloropsis, Trachydiscus* and other members of Eustigmaceae) or starch (most chlorococcal and volvocean algae) as their energy and carbon reserves. In general, starch-producing algae can also accumulate small amounts of lipid under standard conditions, but only some of them, under specific growth conditions, can be induced to overproduce lipids instead of starch (Zhukova et al. [1969](#page-23-0)).

In this review, we will focus exclusively on those algal species that accumulate polysaccharides (starch, cellulose), which can be used as a feedstock for the production of bioethanol.

In these species, starch is the final energy storage molecule formed in the photosynthetic fixation of carbon dioxide. Located in chloroplasts, it is utilized to provide energy for the cell's vital metabolic processes, and is independent of varying external energy and carbon supplies.

Many microalgae grow rapidly and are considerably more productive than land plants and macroalgae (seaweed) (Chisti [2007](#page-19-0)). There are several main groups of microalgae, which differ primarily in pigment composition, biochemical constituents, ultrastructure, and life cycle. In this chapter, we will pay attention to those representatives of the green algae (Chlorophyta) that may be a potential source of starch for future biotechnological applications in the production of bioethanol.

The green algae are a large group from which the Embryophytes (higher plants) emerged. The group, including both green algae and embryophytes, is monophyletic (and often just known as the Kingdom Plantae). There are about 6,000 species of green algae; many live most of their lives as single cells, while others form colonies or long filaments.

Almost all forms have chloroplasts. These contain chlorophylls *a* and *b*, giving them a bright green color (as well as the accessory pigments β-carotene and xanthophylls), and have stacked thylakoids (van den Hoek et al. [1995\)](#page-22-0). All green algae have mitochondria with flat cristae. The storage product for members of this group is true starch, found inside the chloroplasts and consisting of amylose and amylopectin. The starch (seen as whitish granules under the electron microcopy) can often be observed surrounding the pyrenoid, a distinct spherical structure embedded in the chloroplast. There may be more than one pyrenoid, the pyrenoid may not always be present (e.g., *Ankistrodesmus* and *Tetraedron*), or the pyrenoid may be lacking completely. In most representative taxa, the cells are surrounded by a cellulosic wall. Some taxa may also have sporopollenin deposited on the wall. This gives added strength and is thought to help prevent desiccation.

Some microalgae (*Chlorella*, *Scenedesmus, Chlamydomonas*, *Tetraselmis* etc.) are known to contain a large amount of starch and cellulose, providing the raw materials for bioethanol production (Matsumoto et al. [2003](#page-21-1)). Many of the properties of starches, which determine their suitability for particular end-uses, depend on their amylose/amylopectin ratios. These properties include gelatinisation characteristics, solubility, and the formation of resistant starch. The structure of algal starch (*Chlorella*) resembles that of cereals (ca. 34% of amylose content) and its gelatinisation temperature (ca. 65° C), as determined by viscosity measurements, also suggests a structural similarity (Maršálková et al. [2010\)](#page-21-2).

2 Starch Synthesis During the Cell Cycle

As early as 1967, Duynstee and Schmidt [\(1967](#page-20-0)) monitored the net content of starch within the cell cycle in synchronous cultures of a high-temperature strain of *Chlorella pyrenoidosa*. They found that in light, starch accumulates continuously nearly to the end of the cell cycle, followed by cessation immediately prior to and during the period of nuclear division and cytokinesis (Fig. [1\)](#page-2-0).

These results were confirmed in synchronized cultures of the chlorococcal alga *Scenedesmus quadricauda* (Šetlík et al. [1972](#page-22-1)) where the starch content, promoted mainly by light energy input, was also dependent on the stage of cell development. Starch content rose in the early growing phase and decreased thereafter, even under continuous light (Hirokawa et al. [1982\)](#page-20-1). Net starch accumulation can be an indicator of the balance between material and energy supply (provided by photosynthesis) and their use in synthetic processes. During nuclear and cellular divisions, starch is dramatically degraded, as indicated by cytological (Murakami et al. [1963](#page-21-3)) and biochemical (Baker and Schmidt [1964a](#page-19-1); Baker and Schmidt [1964b](#page-19-2); Curnutt and Schmidt [1964](#page-20-2); Herrmann and Schmidt [1965;](#page-20-3) Johnson and Schmidt [1966](#page-21-4); Cole et al.

Fig. 2 Electron microscopic photographs of daughter (**a**) and mother (**b**) cells of *Chlorella vulgaris* grown in complete mineral medium, *N*, nucleus; *S*, starch granules. Bars: 2 μm. (modified according to Branyiková et al. [2011](#page-19-3))

[1998;](#page-20-4) Schmidt [1966](#page-21-4)) changes and increased activity of starch degrading enzymes (Levi and Gibbs [1984;](#page-21-5) Wanka et al. [1970](#page-22-2)). Starch degradation products fulfill demands for both carbon building blocks for organic synthesis and increased energy demands during nuclear and cellular division. Because the photosynthetic rate was approaching its lowest level of the cell cycle during this stage of cellular development (Sorokin [1957](#page-22-3)), accumulated carbon and energy reserve starch was mobilized and utilized during this stage of cellular development. The net content of starch is comprised of starch that was synthesized minus that which was utilized. It is apparent therefore that net accumulation data underestimates the value of actual starch synthesis during the cell cycle. Accumulation of starch during the cell cycle and its phases was also studied in synchronized batch cultures of *Chlorella vulgaris* (Fig. [2\)](#page-3-0). Culture proliferation and changes in starch content were monitored under continuous illumination. The relative starch content reached a maximum of 46% of DW just prior to cell division, which commenced at about the $10th$ hour of growth in light (Brányiková et al. [2011\)](#page-19-3).

During cell division, the relative starch content decreased, even in cells grown in light, to 13% of DW. If the cells were transferred to the dark, they exhausted the stored starch nearly completely (4% of DW). Hence, significant cellular energy and material reserves are maintained to ensure that at any point in development, vital processes can run to completion even if the cells are placed in the dark.

2.1 Starch Production as Affected by Light Intensity

As the ultimate storage form of photosynthetically fixed carbon, starch plays a central role in algal metabolism. Chloroplasts convert carbon dioxide into glucose, which, with the action of starch synthases and branching enzymes, is converted to a mixture of linear and branched glucans (amylose and amylopectin) that determine the properties of the starch. The extent of starch synthesis and degradation in the chloroplast is tightly regulated by the available light (Sundberg and Nilshammar-Holmvall [1975;](#page-22-4) Brányiková et al. [2011](#page-19-3)) and by a number of environmental factors.

Synchronized cultures of *Scenedesmus quadricauda* were grown at two different light intensities and thus differing in specific growth rate. Populations characterized by low growth rates accumulated starch to somewhat more than one third of the value (per cell) obtained in cells with a high specific growth rate. Hence even in lightlimited growth, significant cellular energy and material reserves are maintained. As described above for *Chlorella*, this is, perhaps, to ensure that at any point in development, vital processes can run to completion even if the cells are placed in the dark. Starch reserves were consumed nearly completely if cultures were transferred to the dark (Šetlík et al. [1972\)](#page-22-1).

In synchronized cultures of *Chlorella vulgaris* grown under continuous illumination, the relative content of starch was substantially reduced (maximum 26% of DW) in response to an increase in biomass concentration and corresponding decreased mean light intensity (Sundberg and Nilshammar-Holmvall [1975;](#page-22-4) Brányiková et al. [2011](#page-19-3)). The important role of light intensity on starch content was confirmed in experiments with asynchronous cultures grown at different starting biomass concentrations and consequently, at different mean light intensities, which were continuously decreasing with increasing biomass (Fig. [3\)](#page-5-0).

The higher the mean light intensity, the higher was the starch content. A decrease in mean light intensity below a certain level $(275 \text{ }\mu\text{mol}/(\text{m}^2 \text{ s}^1))$ caused a gradual decline in the rate of starch synthesis, leading finally, to no net increase in starch content or even to its decrease in the case of very low mean light intensities (Fig. [3](#page-5-0)) (Zhang et al. [2002;](#page-23-1) Eriksen et al. [2007](#page-20-5); Thu et al. [2009](#page-22-5)).

Beside light quantity, its quality can also play a role in starch synthesis and its degradation. Kamiya and Kowallik ([1986](#page-21-6)) studied starch synthesis and degradation in *Chlorella vulgaris* under autotrophic conditions in blue and red light and found that in blue light, increased starch degradation occurred, causing a decrease in total starch content when compared with cells grown in red light (Miyachi et al. [1978;](#page-21-7) Kamiya and Kowallik [1986\)](#page-21-6).

2.2 Starch Production as Affected by Temperature

Semenenko and his collaborators published a series of papers in which they described and analyzed restrictive effects of high temperatures on the reproductive and synthetic processes in different strains and species of *Chlorella* (Semenenko et al. [1967](#page-22-6), [1969;](#page-22-7) Semenenko and Zvereva [1972a](#page-21-8), [b\)](#page-22-8). Unfortunately, their work was published in Russian more than 40 years ago and was neglected by researchers both then and now. They had, however, discovered the basic principles and rules, which control the production of energy reserves in algae, information, which only recently has been rediscovered.

The authors found that for the high-temperature strain of *Chlorella* sp. K (growth optimum at $36-38$ °C) the effective restrictive temperature was $43-44$ °C. With a mesophilic strain of *C. pyrenoidosa* (growth optimum 26–28°C) the restrictive **Fig. 3** The effect of mean light intensity (**c**) on changes in biomass concentration (DW in g/L) (**^a**), starch content in g/L (**b**) in asynchronous cultures of *Chlorella vulgaris*. The cultures were grown in a laboratory photobioreactor continuously illuminated by a constant incident light intensity of 780 μmol/(m² s¹) and at various biomass concentrations (numerals on curves indicate initial dry weight in g/L). *Dashed horizontal lines* indicate the biomass concentration (panel **a**) and the mean light irradiance (panel **c**) under which the net starch content begins to decrease. (modified according to Brányiková et al. [201](#page-19-3)1)

temperature for cellular division is 35–36°C. At the time when cells ceased dividing, giant cells with a diameter up to 5–6 times that of the daughter cells formed a significant part of the population. The higher the average irradiation received by the cells, the greater were the number of giant cells ultimately obtained, and also the faster was attainment of the final stage. Very large quantities of starch were found in the giant cells of *Chlorella* sp. K, while extensive lipid inclusions appeared in *C. pyrenoidosa*. To explain the behavior of cell populations at the restrictive temperature, Semenenko et al. assumed that the extreme temperatures effectively blocked reproductive events while other synthetic processes were mainly unaffected or were even stimulated by the high temperature, as was the case with photosynthesis. However, with reproduction processes inhibited, the demand for cell construction material gradually decreased, photosynthates accumulated in the later phases and by a feedback type of inhibition, they finally stopped photosynthesis.

The effect of temperature on $CO₂$ fixation in photosynthesis and on starch conversion into sucrose was studied in a series of papers by Nakamura and Miyachi (Nakamura and Miyachi [1982a,](#page-21-9) [b,](#page-21-10) [1980](#page-21-11); Nakamura [1983](#page-21-12); Nakamura and Imamura [1983\)](#page-21-13). They found that in *Chlorella vulgaris*, a temperature higher than 30°C caused acceleration of the conversion of starch to sucrose. This temperature-induced degradation of starch also occurred in the dark, proving that photosynthetic electron transport is not involved (Nakamura and Miyachi [1982a;](#page-21-9) Nakamura and Miyachi [b\)](#page-21-10).

3 Treatments for Overproduction of Starch

Algal cells grown in light synthesize starch in chloroplasts, in the form of starch bodies (see, chapter *Starch synthesis during the cell cycle*). However, these starch reserves only represent the net production of starch, comprising the difference between gross synthesis of starch and that used to drive many cellular processes and events, particularly for the highest consumers of energy and carbon such as DNA replication, nuclear division and cytokinesis.

The final cellular content of starch is determined by two main factors: (1) the mean light intensity, which determines the rate of starch synthesis and its final content, and (2) processes (mainly cell division) consuming starch for their energy demands. Therefore to attain a maximum yield of starch, processes that utilize starch as energy and carbon source must be stopped or minimized, while conditions enabling production of starch should be optimized (Brányiková et al. [2011\)](#page-19-3).

3.1 Applying Specific Inhibitors

3.1.1 Inhibition of Nuclear DNA Replication

In the chlorococcal alga *Scenedesmus quadricauda*, specific inhibition of nuclear DNA synthesis by 5-fluorodeoxyuridine (FdUrd) was used to study cell cycle regulatory processes (Zachleder [1994;](#page-22-9) Zachleder [1995](#page-22-10)). FdUrd interrupts the

Fig. 4 Microphotographs of the mixture of daughter coenobia (*A*), dividing mother cells (*M*) in a control culture of *Scenedesmus quadricauda* (after 14 h of light), and "giant" cells (*F*) from an FdUrd-treated culture (36 h of growth in the presence of FdUrd 25 mg/L). Scale bar=10 μm. (modified according to Zachleder [1995\)](#page-22-10)

deoxyribonucleotide metabolic pathway just before the last step by inhibiting the enzyme thymidylate synthase (EC 2.1.1.45), which catalyzes the reductive methylation of 2′-deoxyuridylate to form deoxythymidylate (Cisneros et al. [1993](#page-19-4); Follmann [1983;](#page-20-6) Bachmann et al. [1983\)](#page-19-5). As a consequence, subsequent events such as nuclear division and cytokinesis were also blocked (Zachleder et al. [1996\)](#page-22-11). If FdUrd was added to a synchronized culture of *Scenedesmus quadricauda* at the beginning of the cell cycle, the cells remained uninuclear with no nuclear DNA replication, and no nuclear or cellular division occurred during experiments (Zachleder [1995](#page-22-10)). The growth of cells and processes involved in their growth, like RNA and protein accumulation, were not affected for an interval corresponding to at least the duration of two or three cell cycles, attaining six doublings in cell volume by the end of this period. While FdUrd effectively blocked DNA replication in *Scenedesmus quadricauda*, growth processes were not affected by the presence of the inhibitor. Because activity of the chloroplast was not affected, cells grew to a giant size, producing large numbers of starch grains in their chloroplasts (Fig. [4](#page-7-0)).

3.1.2 Inhibition of Cytoplasmic Proteosynthesis

Cycloheximide, an antibiotic inhibiting eukaryotic cytoplasmic protein synthesis, prevented cultures of *Scenedesmus quadricauda* (Zachleder et al. [2002](#page-22-12)) or *Chlorella vulgaris* (Brányiková et al. [2011](#page-19-3)) from undergoing nuclear division and consequently no cell division occurred. Starch, however, was intensively synthesized and in the case of *Chlorella vulgaris,* attained (within 4–5 h) a level of at least 60% of DW $(Figs. 5 and 6).$ $(Figs. 5 and 6).$ $(Figs. 5 and 6).$ $(Figs. 5 and 6).$ $(Figs. 5 and 6).$

After 5 h of growth in the presence of cycloheximide, the rate of starch synthesis equilibrated with biomass growth, and a high relative content of starch remained constant (Fig. [5](#page-8-0)). This shows that starch synthesis in the chloroplast is independent of cytoplasmic protein synthesis as well as DNA replication and other cell reproductive processes.

Fig. 6 Electron microscopic photographs of cells of *Chlorella vulgaris* grown in the presence of cycloheximide (1 mg/L). *S*, starch granules. Bar: 5 μm. (modified according to Brányiková et al. [2011](#page-19-3))

3.2 Nutrient Limitation

While treatments with inhibitors are useful and important from the viewpoint of basic research, their industrial utilization is problematic for reasons of economics and possible environmental contamination. Suppression of cell cycle events requiring energy and carbon from starch reserves can however, be attained by limitation of macroelements in mineral media used for culturing autotrophically grown algae. An increase of starch content in algal cells upon limitation of some elements (nitrogen, sulfur, or phosphorus) was described a long time before the utilization of algal starch for production of biofuels was considered (Klein [1987;](#page-21-14) Ball et al. [1990](#page-19-6); Zachleder et al. [1988;](#page-22-13) Šetlík et al. [1988](#page-22-14); Ballin et al. [1988\)](#page-19-7). Although in contrast to the use of specific inhibitors, the exact mechanism of element limitation-induced starch synthesis is neither well known nor specific, their application for scale up culturing is advantageous both from economic (saving fertilizers) and environmental points of view. Nevertheless, the principles of this approach are similar to that of inhibitors;

the absence of different elements caused a cessation of processes consuming starch while starch production was unaffected.

3.2.1 Effect of Nitrogen Starvation

When microalgae are starved for nitrogen, the main element of protein and nucleic acids, cell division ceases and photosynthetic carbon partitioning is switched towards starch synthesis. In *Chlamydomonas reinhardtii* cultivated in 1.0 mM KNO₃, protein and chlorophyll *a* content decreased when nitrate was exhausted, while carbohydrate content increased nearly six-fold (Klein [1987](#page-21-14)). Similarly in *Scenedesmus quadricauda,* macromolecular syntheses of RNA, protein and DNA were arrested when a nitrogen source was lacking (Fig. [7](#page-10-0)). This diverted the photosynthetic carbon flow into starch without it being consumed as a carbon and energy source (Ballin et al. [1988\)](#page-19-7).

More than 20 years later, the effect of nitrogen starvation was verified in a production strain of *Chlorella vulgaris* p12 that accumulated starch from 35 to 58% of dry weight (DW) in nitrogen-limited medium (Brányiková et al. [2011](#page-19-3); Behrens et al. [1989;](#page-19-8) Dragone et al. [2011\)](#page-20-7).

Starch accumulation has been studied mostly in freshwater microalgal strains including *Chlamydomonas reinhardtii* (Zhang et al. [2002;](#page-23-1) Eriksen et al. [2007](#page-20-5); Thu et al. [2009\)](#page-22-5), *Chlorella vulgaris* (Brányiková et al. [2011](#page-19-3); Behrens et al. [1989;](#page-19-8) Dragone et al. [2011](#page-20-7)) and *Scenedesmus obliquus* (Rodjaroen et al. [2007\)](#page-21-15). Less attention, however, has been paid to marine microalgae other than *Dunaliella tertiolecta* (Ike et al. [1997](#page-20-8)), and *Chlamydomonas perigranulata* (Hon-Nami [2006\)](#page-20-9).

The marine green microalga, *Tetraselmis subcordiformis*, accumulates starch autotrophically (Zheng et al. [2011\)](#page-23-2) or mixotrophically (Ji et al. [2011\)](#page-20-10). The maximum starch content was 35% DW and starch productivity was 0.26 g L^{-1} d⁻¹. Using sulfur starvation to direct starch overproduction, *T. subcordiformis* could achieve a starch productivity of 0.62 g L⁻¹ d⁻¹ and a starch content of 62.1% DW, thus endowing it with great potential for starch production (Yao et al. [2012](#page-22-15)).

3.2.2 Effect of Phosphorus Starvation

Phosphorus is indispensable for the basic processes of energy flow in cells and is one of the essential macromolecules. In *Scenedesmus quadricauda,* the first process affected by the absence of phosphorus was RNA synthesis, which leveled off relatively early in the cell cycle, *i.e.* around the 5th hour. Protein synthesis showed the first signs of restriction much later and ceased at the time when protoplast fissions were taking place and the release of daughter cells started (Zachleder et al. [1988](#page-22-13)). Accumulation of starch, however, indicated that the rate of photosynthesis was nearly normal during the entire cycle in which RNA synthesis stopped as well as in the next cycle, in which no macromolecular syntheses took place (Fig. [8](#page-11-0)). There were sufficient cellular phosphate reserves in diverse forms at the stage when RNA synthesis leveled off (Voříšek and Zachleder [1984\)](#page-22-16).

Fig. 7 Variation in RNA (**a**), protein (**b**), DNA (**c**), starch (**d**) content during the cell cycle in control (*open symbols*) and nitrogen-starved (*closed symbols*) synchronous populations of *Scenedesmus quadricauda. Arrows* indicate the beginning of the dark period in control cultures. The nitrogenstarved culture was kept in continuous light. (modified according to Ballin et al. [1988\)](#page-19-7)

Daughter cells generated under phosphate deficiency and exposed to light in a phosphate-free medium were unable to develop further. The synthesis of nucleic acids was completely suppressed. Protein synthesis was also severely restricted, and only a slight increase in the amount of protein was observed (Fig. [8\)](#page-11-0). The photosynthetic apparatus continued to work at a moderate but sustained rate and starch accumulated, characteristic of a phosphorus-fed cell.

In complete medium, the phosphorus-starved daughter cells resumed macromolecular syntheses with a lag of about 5 h. Due to the large amount of starch accumulated in these cells, they were able to progress through an entire cell cycle in the dark after being supplied with phosphorus (Zachleder et al. [1988\)](#page-22-13).

Phosphorus limitation in *Chlorella vulgaris* caused cell division to be blocked and was severe enough to reduce biomass synthesis from the very beginning of the experiment and to stop it completely after 12 h in the absence of phosphorus in the medium, or after 20 h of growth in medium with a five fold decreased concentration of phosphate. Thereafter, starch began to degrade, and its concentration (in g/L) and relative content (% of DW) decreased (Fig. [9](#page-12-0), Brányiková et al. [2011](#page-19-3)).

3.2.3 Effect of Sulfur Starvation

Sulfur, which is necessary for the metabolism of proteins, lipids, and molecules associated with electron transport chains, is another essential element that regulates starch accumulation in microalgae.

If synchronized cultures of *Scenedesmus quadricauda* were incubated under photosynthesizing conditions in a sulfur-free medium, inhibition of RNA synthesis occurred close to the end of the first cell cycle and protein synthesis ceased two hours later (practically at the time of protoplast fission). If the daughter cells derived from the starved populations were kept in sulfur-free medium, macromolecular syntheses were dramatically restricted (Fig. [10\)](#page-12-1).

Only photosynthesis continued to produce starch at a similar rate to that of normally grown cells. Thus, a very large amount of starch accumulated (Figs. [10](#page-12-1) and [11\)](#page-12-2).

Supported by these reserves, when starved cells were refed with sulfur, they progressed through an entire cell cycle in the dark and divided into eight daughter cells. In sulfur-supplied cells, both in the dark and in light, RNA protein and DNA synthesis started without any delay in a similar way to the control culture. If competition for sulfur reserves occurred between growth and division processes, the former were preferred in the light and the latter in the dark (Šetlík et al. [1988](#page-22-14)).

Fig. 9 Effect of nitrogen, phosphorus, or sulfur limitation on changes to relative starch content (% of DW) in asynchronous cultures of *Chlorella vulgaris*. The cultures were grown in a laboratory photobioreactor with continuous illumination at an incident light intensity of 780 μ mol/(m² s¹) in a complete mineral medium (*diamonds, dotted line*) or in nitrogen (*diamonds, solid line*), phosphorus (*triangles*), or sulfur (*squares*) limited media. (modified according to Brányiková et al. [2011](#page-19-3))

60

40

Fig. 11 Electron microscopic photographs of cells of *Chlorella vulgaris* grown in sulfur-limited medium. *S*, starch granules. Bar: 5 μm. (modified according to Brányiková et al. [201](#page-19-3)1)

In *Chlamydomonas reinhardtii,* deprivation of sulfur caused changes in cellular metabolism resulting in cell expansion, regulated protein degradation and a significant increase (nearly ten times) in the starch/protein ratio (Ball et al. [1990](#page-19-6); Zhang et al. [2002](#page-23-1)). Similarly, the alga *Dunaliella salina* altered the partitioning of photosynthate between starch and protein and, under sulfur-deprived conditions, induced a ten-fold increase in the starch/protein ratio (Cao et al. [2001](#page-19-9)).

Similarly, it was recently demonstrated that a sulfur-limited culture of *Chlorella vulgaris* maintained a high starch content (60%) and a starch productivity of 0.8 g $l⁻¹$ d⁻¹ (Brányiková et al. [2011\)](#page-19-3), levels significantly higher than those calculated under nitrogen limiting conditions (0.119–0.55 g l^{-1} d⁻¹) (Dragone et al. [2011;](#page-20-7) Fernandes et al. [2010](#page-20-11); Behrens et al. [1989\)](#page-19-8). This suggests that compared to nitrogen and phosphorus, sulfur limitation is more effective in enhancing starch productivity. For practical and industrial applications, it was also important that, in the case of sulfur limitation, the high level of starch was maintained for more than 24 h, whereas limitation of either nitrogen or phosphorus caused a decay in starch content and cell death within a few hours (Fig. [9,](#page-12-0) Brányiková et al. [2011\)](#page-19-3).

Comparing cultures grown in complete medium and sulfur-starved cultures under conditions simulating alternating days and nights, degradation of starch was substantially lower in sulfur-starved cultures than in cultures grown in complete medium, while net synthesis during the light period was substantially higher (Fig. [12](#page-14-0)).

The effect of sulfur limitation was apparently due to inhibition of processes using energy and carbon derived from starch reserves (Brányiková et al. [2011\)](#page-19-3) as is the case for other treatments.

3.3 Utilization of Waste CO₂

To attain high productivity, microalgae require $CO₂$ as a photosynthetic carbon source at the same time the cost of CO_2 represents one of the major cost factors for algal biomass. Microalgae can tolerate and utilize substantial levels of CO_2 ; thus, photosynthetic carbon dioxide can be derived from cheap sources such as combustion of organic waste, fermentation processes or other sources (Mann et al. 2009 ; Douskova et al. 2009 ; Doucha et al. 2005). Some cheap sources of $CO₂$ such as flue gas from incineration of domestic waste (Douskova et al. [2009](#page-20-12)), biogas produced from distillery stillage (Doušková et al. [2010;](#page-20-14) Douskova et al. [2009;](#page-20-12) Kaštánek et al. [2010\)](#page-21-17) or flue gas from cogeneration unit combusting biogas from anaerobic digestion of swine manure (Doušková et al. [2010;](#page-20-14) Douskova et al. [2009;](#page-20-12) Kaštánek et al. [2010](#page-21-17)) have been used successfully.

This provides an additional benefit of algal biomass production because carbon dioxide from different sources including waste incinerators, power stations, limekilns, cogeneration units, etc. *in situ* could be bioremeditated (Doušková et al. [2010;](#page-20-14) Douskova et al. [2009](#page-20-12); Kaštánek et al. [2010;](#page-21-17) Nigam and Singh [2010\)](#page-21-0).

Fig. 12 Effect of alternating light and dark periods on changes in biomass concentration (DW in g/L) (**a**), starch content (g/L) (**b**) and relative starch content (% of DW) (**c**) in untreated (*empty symbols*) and in sulfur limited (*full symbols*) cultures of *Chlorella*. (modified according to Brányiková et al. [2011](#page-19-3))

3.4 Scale Up Culturing

To verify laboratory findings, experimentation using large-scale production units is the only way to prove real outcomes of any proposed approach. Unfortunately, to date, with the exception of the paper by Brányiková et al. [\(2011\)](#page-19-3), large-scale production studies giving relevant data on the production of lipid-enriched algal biomass have not been carried out.

Fig. 13 Changes in biomass concentration (DW in g/L), starch content (g/L) (**a**), and relative starch content (% of DW) (**b**) in cultures of *Chlorella vulgaris*. The cultures were grown in an outdoor scale-up thin-layer photobioreactor in complete mineral medium for 96 h and then in sulfurlimited medium (-S). *Dark periods* (nights) are marked by black stripes and separated by vertical solid lines. Linear regression curves of the 1st order (*solid lines*) were fitted to the experimental data separately for the periods of culture grown in complete medium and sulfur-limited cultivation. (after Brányiková et al. [2011](#page-19-3))

An increase in starch production in sulfur-limited culture, up to a maximum of 50% of algal biomass (DW) was demonstrated under field conditions using the outdoor scale up, thin-layer solar photobioreactor (Fig. [13](#page-15-0)).

Despite the relatively unfavorable climatic conditions of Trebon (Czech Republic), a total yield of starch, calculated per ha over a season of 150 days, was 7 t (Brányiková et al. [2011](#page-19-3)). In optimum locations for photoautotrophic production, such as Greece, with a season lasting approximately 250 days, the overall harvest might be increased by a factor of 5.

Based on the findings described above cheap starch-enriched biomass could be produced from highly productive *Chlorella* cultures if grown in suitable outdoor photobioreactors in which photosynthetic carbon dioxide is derived from combustion of organic waste, fermentation processes or other sources (Mann et al. [2009;](#page-21-16) Douskova et al. [2009](#page-20-12); Doucha et al. [2005](#page-20-13)). Furthermore, after starch extraction, the remaining cell residue, comprising mostly protein, can be used as an animal feed supplement to further decrease the cost of starch production. This characteristic enhances the ecological and economic impact of the proposed technology because of its potential to bioremediate carbon dioxide emissions from different CO₂ sources including waste incinerators, power stations, limekilns, cogeneration units, etc. *in situ*.

4 Bioethanol

4.1 Principles of Bioethanol Production from Algae

Even though some algae can act as mini factories for the production of ethanol during dark fermentation, and attempts have been made to create genetic engineered microalgae for the direct production of ethanol (Ueda et al. [1996](#page-22-17); Hirano et al. [1997](#page-20-15)), these approaches are still not applicable in large-scale biotechnology. The general principle of bioethanol production from microalgal biomass is similar to its production from higher plants. Algal polysaccharides (starch, eventually cellulose and hemicellulosis) are enzymatically hydrolyzed to fermentable sugars, which are subsequently fermented to ethanol by a suitable ethanol producer (yeast, bacteria) (Kelsall and Lyons [1999\)](#page-21-18). Ethanol is then concentrated by distillation or membrane filtration and dehydrated. Despite the fact that the overall procedure is quite simple, and for different raw materials, well managed even in large scale, each step poses specific difficulties in the case of microalgal biomass. The microalgae store starch inside the cell, which is why the cells must be disintegrated prior to hydrolysis in order to liberate starch for the action of enzymes. Disintegration can be by mechanical means (e.g., ultrasonic, explosive disintegration, mechanical shear, etc.) or by dissolution of cell walls using enzymes (Percival Zhang et al. [2006](#page-21-19)). The starch-rich biomass is then processed using technologies similar to other starch-based feedstocks, involving processes of (1) gelatinization (heating in water) which improves the availability of starch for amylase hydrolysis (2) liquefaction i.e. breaking of starch via α-amylases into soluble, shortchain dextrins and oligosaccharides resulting in significant reduction of viscosity, (3) saccharification of the starch-hydrolysate to glucose syrup catalyzed by glucoamylase and (4) fermentation of simple sugars to ethanol by a suitable strain of yeast, (e.g. *Saccharomyces cerevisiae,* for review, see Singh et al. [2010\)](#page-22-18) or bacteria, e.g. *Zymomonas mobilis* (Matsumoto et al. [2003](#page-21-1); Rubin [2008\)](#page-21-20). Using special enzymes and yeast strains, the whole process can be designed so that no gelatinisation is necessary and liquefaction, saccharification and fermentation are carried out simultaneously (Doran-Peterson et al. [2009](#page-20-16)).

Finally, the ethanol is separated from the fermentation broth by distillation or membrane filtration and the concentrated ethanol (95%) can be directly used in special types of engines as so called E100, or dehydrated and blended with gasoline (Demirbas [2001](#page-20-17); Nigam and Singh [2010](#page-21-0); Brennan and Owende [2010](#page-19-10)). The solid residue from the process can be used as animal feed (McKendry [2002](#page-21-21)), agricultural fertilizer or as a feedstock for biogas production (for more details, see review John et al. [2011](#page-20-18)).

The recent extensive research effort to develop algal biomass with a high starch/ cellulose content is aimed at providing a product that could serve as a substrate for ethanol production. As mentioned in the preceding chapters there are algal species that can be directed to produce large amounts $(50\%$ of the dry weight) of starch, suitable for ethanol production after hydrolysis (Brányiková et al. [2011;](#page-19-3) Yao et al. [2012\)](#page-22-15).

4.2 Approaches to Cheapen Bioethanol Production

4.2.1 Carbon Dioxide Supply from Flue Gases

Since biomass accumulation by feedstock crops can utilize carbon dioxide from flue gases, their growth for bioethanol production can reduce green house gas levels as well as the price of bioethanol production (Borowitzka [2008](#page-19-11); Douskova et al. [2009\)](#page-20-12).

4.2.2 Hydrolysis of Algal Biomass for Yeast Nutrition

In order to achieve the highest possible fermentation yield (theoretically 0.5L g of ethanol/L g of glucose) appropriate nutrition for yeast has to be ensured (source of N, P, S etc.). The cost of nutrients can be cut by adding proteases during polysaccharide hydrolysis, which liberates free amino acids (a source of nitrogen for yeast) from the protein fraction of the algal biomass. The processes of enzymatic hydrolysis and fermentation can also be carried out simultaneously (Doran-Peterson et al. [2009\)](#page-20-16).

4.2.3 Production of Ethanol from Residual Algal Biomass

The residual biomass obtained e.g. after oil extraction can also be used as a substrate for ethanol production, as was shown when *Chlorococcum* sp. was used as a substrate for bioethanol production through fermentation by *Saccharomyces bayanus* (Harun et al. [2010](#page-20-19); Harun et al. [2011\)](#page-20-20)*.* The lipid-extracted microalgal debris was used for fermentation and the yield of bioethanol was about 3.8 g/L from 10 g/L of the substrate. Enzymatic hydrolysis of *Chlorococcum* sp. by cellulase obtained from *Trichoderma reesei* was examined. Hydrolysis was conducted under varying conditions of temperature, pH and substrate concentration, with constant enzyme dosage. The highest glucose yield of 64.2% (w/w) was obtained at a temperature of 40 °C. pH 4.8, and a substrate concentration of 10 g/L of microalgal biomass (Ueno et al. [1998\)](#page-22-19). Overall, the enzymatic hydrolysis process proved to be an effective mechanism to enhance the saccharification process of microalgal biomass.

4.2.4 Hydrolysis of Cellulose from Cell Walls

Besides starch, green algae can accumulate cellulose as the cell wall carbohydrate, which can also be used for ethanol production. The cellulosic biomass from other plant sources that from algae can be enzymatically or chemically hydrolyzed to simple sugars, which can then be fermented to ethanol (Kelsall and Lyons [1999;](#page-21-18) Doran-Peterson et al. [2009](#page-20-16)).

The advantage of microalgal cellulose is that it is not linked to lignin (unlike cellulose in wood, straw, etc.), which makes hydrolysis easier comparing to the lignocellulosic materials.

4.2.5 Dark Fermentation—Direct Production of Ethanol

Some algae can even serve as a self-biorefinery for ethanol production during anaerobic dark conditions by utilizing their photosynthates. Microalgae fix CO_2 during photosynthesis and accumulate starch in their cells. Some microalgae can also grow under dark conditions in the presence of organic nutrients such as sugars and thereby accumulate starch (heterotrophic nutrition) (Chen et al. [2009](#page-19-12)). If dark and anaerobic conditions are established, the oxidative degradation of starch becomes incomplete and, depending on the type of the microalga, hydrogen gas, carbon dioxide, ethanol, lactic acid, formic acid, acetic acid and other products can be produced in varying proportions (John et al. [2011](#page-20-18)). Algal cells contain large amounts of polysaccharides composed of glucose, which can be catabolized rapidly under dark and anaerobic conditions to ethanol. Conversion from intracellular starch to ethanol under dark and anaerobic conditions was observed in several algal strains (Hirano et al. [1997\)](#page-20-15). However, the levels of conversion to ethanol were significantly different from each other. Relatively high conversion rates of $30-40\%$ (vs. a theoretical yield of 0.56 g of ethanol/L g of starch) were observed in two strains, *Chlamydomonas reinhardtii*

(UTEX2247) and Sak-1. Importantly, there was no need for nitrogen flushing due to the complete utilization of oxygen and there was no need for agitation for ethanol production. The optimal pH for ethanol production from *Chlamydomonas* was 7–8 and a temperature of 25–30°C. Ethanol was also produced via dark fermentation of cellular starch from *Chlorococum littorale*, where an increase in the incubation temperature affected the mode of cellular starch decomposition and brought about an increase in ethanol productivity. If methyl viologen was added to the reaction, hydrogen formation drastically decreased while ethanol productivity increased (Ueno et al. [1998](#page-22-19)). Exploiting microalgal strains to accumulate starch/cellulose and directly utilize their enzymatic or anaerobic digestion systems to produce ethanol can provide a cost-effective bioethanol production process. (For detail review of these alternative ways of ethanol production, see John et al. [2011\)](#page-20-18).

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