

Anabaena: A New Chassis for Space Exploration

Abstract

The next era of human space exploration demands sustainable biological systems capable of supporting long-duration missions to the Moon, Mars, and beyond. Among candidate organisms for biological in situ resource utilization (bio-ISRU), the filamentous cyanobacterium *Anabaena* has emerged as a particularly promising chassis. This nitrogen-fixing, photoautotrophic microorganism combines exceptional tolerance to extreme environmental stressors with the unique capacity to differentiate specialized heterocyst cells, creating a natural compartmentalization system ideal for metabolic engineering. Recent advances have demonstrated that *Anabaena* species—particularly *Anabaena* sp. PCC 7938 and PCC 7120—can thrive under Mars-like atmospheric conditions, utilize regolith simulants as nutrient sources, and serve as foundational elements in closed-loop life support systems. This review critically examines the biological features that position *Anabaena* as a next-generation space biotechnology platform, analyzes current limitations in genetic manipulation and productivity optimization, and highlights critical knowledge gaps that must be addressed before operational deployment. We argue that the convergence of synthetic biology tools, systems biology understanding, and space-adapted cultivation hardware positions *Anabaena* as the cornerstone organism for establishing self-sustaining human presence beyond Earth.

Introduction: The Imperative for Biological Life Support Systems

The transition from short-duration exploration missions to permanent human settlement of extraterrestrial environments represents one of humanity's most ambitious endeavors. Current mission architectures for crewed Mars exploration, planned by NASA, ESA, and emerging space agencies for the 2030s and beyond, face a fundamental constraint: the prohibitive cost of transporting consumables—oxygen, water, nutrients, and materials—from Earth[1]. Mass-to-orbit costs, even with next-generation launch systems, render long-term resupply economically unsustainable. The solution lies in biological in situ resource utilization (bio-ISRU), wherein engineered microorganisms convert local resources into life-support consumables[2].

Among photosynthetic microorganisms, cyanobacteria have long been identified as prime candidates for space biotechnology applications[3]. These ancient prokaryotes dominated Earth's early biosphere, precipitating the Great Oxidation Event approximately 2.4 billion years ago through oxygenic photosynthesis. Modern cyanobacteria retain the metabolic versatility and stress tolerance that enabled their ancestors to thrive in harsh primordial environments—traits directly relevant to space applications. However, not all cyanobacteria are equally suited for extraterrestrial deployment.

The genus *Anabaena* has recently emerged as the leading candidate chassis for Mars-focused biotechnology development[4][5]. Unlike unicellular cyanobacteria such as

Synechocystis sp. PCC 6803 or *Synechococcus elongatus*, filamentous *Anabaena* species possess a critical advantage: the capacity for cellular differentiation into heterocysts, specialized nitrogen-fixing cells that create microoxic compartments to protect the oxygen-sensitive nitrogenase enzyme complex[6]. This natural compartmentalization obviates the need for complex bioprocessing infrastructure to separate aerobic and anaerobic metabolism. Furthermore, recent strain selection efforts have identified *Anabaena* sp. PCC 7938 as uniquely capable of utilizing Martian atmospheric gases at reduced pressures and extracting nutrients from regolith simulants[7][8]. These capabilities position *Anabaena* not merely as a research model but as an operational organism ready for engineering toward deployment.

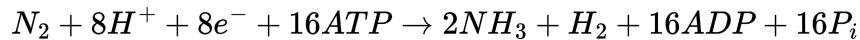
This review synthesizes recent advances in *Anabaena* biology relevant to space applications, critically evaluates current genetic engineering capabilities, examines performance under space-relevant environmental conditions, and identifies the most pressing research gaps. We contend that while significant challenges remain—particularly in achieving industrial-scale productivity and completing genome-to-phenome understanding—*Anabaena* represents the most viable photosynthetic chassis for establishing sustainable extraterrestrial human presence.

Biological Features: Why *Anabaena* Stands Apart

Heterocyst Differentiation and Nitrogen Fixation

The defining feature of *Anabaena* is its capacity for cellular differentiation into heterocysts, a developmental program unparalleled among cyanobacteria[9]. Under nitrogen-limited conditions, approximately 5-10% of vegetative cells in a filament differentiate into heterocysts, forming a quasi-regular spatial pattern with one heterocyst per 10-15 vegetative cells[10]. This remarkable patterning system operates through an elegant molecular mechanism involving two opposing gradients: the activator HetR, a serine-type protease and transcription factor that promotes heterocyst fate, and inhibitors PatS and HetN that suppress differentiation in adjacent cells[11][12].

The heterocyst developmental program involves profound morphological and metabolic remodeling. Differentiating cells deposit a multilayered envelope consisting of an inner glycolipid layer and an outer polysaccharide layer, creating a barrier to oxygen diffusion that reduces internal O₂ concentrations by 100-1000 fold[13]. Photosystem II (PSII) is selectively degraded, eliminating oxygen-producing photosynthesis while retaining photosystem I for ATP generation via cyclic photophosphorylation[14]. The nitrogenase enzyme complex—comprising the iron protein (NifH) and molybdenum-iron protein (NifDK)—is then expressed exclusively in these microoxic cells, enabling biological nitrogen fixation according to the reaction:



Fixed nitrogen is exported as glutamine to adjacent vegetative cells via intercellular molecular exchange through septal junctions, while vegetative cells reciprocally provide heterocysts with reducing equivalents and carbon skeletons from photosynthetically fixed CO₂[15].

For space applications, this natural compartmentalization solves a critical bioprocessing challenge. Oxygen production and nitrogen fixation—both essential for life support—are inherently incompatible in biochemical terms due to the extreme oxygen sensitivity of

nitrogenase (inactivated within seconds at atmospheric O₂ levels). Engineering oxygen tolerance into nitrogenase has proven extraordinarily difficult despite decades of effort[16]. *Anabaena*'s heterocyst system provides an elegant, evolution-tested solution that requires no external energy for maintaining spatial separation.

Genomic Organization and Nitrogen Fixation Gene Rearrangement

The *Anabaena* genome exhibits sophisticated regulatory architecture optimized for coordinating multicellular development. The model strain *Anabaena* sp. PCC 7120 possesses a 7.2 Mb circular chromosome encoding approximately 6,500 genes, along with multiple plasmids[17]. Notably, nitrogen fixation genes undergo developmentally-programmed genomic rearrangements during heterocyst differentiation—a phenomenon rare among prokaryotes[18].

Specifically, an 11 kb DNA element interrupting the *nifD* gene in vegetative cells is excised via site-specific recombination at 11 bp direct repeats flanking the element, creating a functional *nifHDK* operon exclusively in heterocysts[19]. This genetic excision serves multiple functions: it prevents premature nitrogenase expression in vegetative cells (where oxygen would inactivate the enzyme), ensures irreversibility of differentiation, and provides a stable genetic marker distinguishing cell types. Understanding this recombination mechanism has proven crucial for developing heterocyst-specific expression systems in synthetic biology applications[20].

The genome also encodes an extensive repertoire of regulatory proteins governing heterocyst development. The master regulator HetR controls expression of over 100 genes during differentiation, with its regulon encompassing genes for envelope deposition (*hep* genes), nitrogen metabolism (*nir* and *amt* genes), pattern formation (*pat* genes), and metabolic remodeling[21]. The transcription factor NtcA, which responds to cellular nitrogen status, acts upstream of HetR to initiate the developmental cascade when combined nitrogen becomes limiting[22]. This hierarchical regulatory architecture provides multiple intervention points for synthetic biology engineering.

Stress Tolerance and Extremophile Characteristics

While *Anabaena* species are not extremophiles in the classical sense—they do not require extreme conditions for growth—many strains exhibit remarkable tolerance to environmental stressors relevant to space applications. *Anabaena cylindrica* forms akinetes, thick-walled resting cells that confer resistance to prolonged desiccation, enabling survival for years in dried states[23]. These akinetes also show increased tolerance to ionizing radiation, withstanding cumulative doses exceeding 240 mGy during 548 days of exposure in low Earth orbit[24].

UV radiation tolerance varies substantially among strains. While space-exposed *Anabaena cylindrica* biofilms experienced surface bleaching and carotenoid degradation under unfiltered UV, subsurface layers retained viability, suggesting that biofilm architecture provides inherent protection against radiation damage[25]. Desert-dwelling *Chroococcidiopsis* (a related cyanobacterium) shows superior UV tolerance through over-expression of DNA repair genes and synthesis of UV-absorbing pigments[26], traits that could potentially be engineered into *Anabaena* chassis.

Temperature tolerance also merits consideration. *Anabaena* sp. PCC 7938 grows optimally at 25-30°C but retains viability at 15-35°C[27]. Mars surface temperatures ranging from

-125°C to +20°C present challenges, but cultivation hardware would provide thermal control. More relevant is tolerance to rapid temperature fluctuations, an underexplored area requiring systematic characterization.

Critically, recent studies demonstrate that *Anabaena* sp. PCC 7938 grows vigorously under low-pressure, Mars-relevant atmospheric conditions (96% N₂, 4% CO₂ at 100 hPa total pressure), with growth rates comparable to standard laboratory conditions[28]. This finding was unexpected, as most organisms experience severe stress at reduced pressures. The mechanism underlying this pressure tolerance remains unclear but may involve altered gas solubility kinetics favoring more efficient CO₂ and N₂ uptake.

Genetic Engineering Tools: Current Capabilities and Limitations

Transformation Methods and Vectors

Genetic manipulation in *Anabaena* has historically lagged behind model organisms like *Escherichia coli* or *Synechocystis*, but recent years have witnessed substantial progress. Natural transformation (uptake of naked DNA from the environment) occurs in *Anabaena* but with relatively low efficiency (<10⁻⁶ transformants per cell), necessitating optimized protocols[29]. Conjugation from *E. coli* donor strains provides a more reliable alternative, achieving transformation frequencies of 10⁻³ to 10⁻⁴ using self-transmissible or mobilizable plasmids[30].

Several vector systems are now available for *Anabaena*. Broad-host-range plasmids based on RSF1010 and RK2 replicons can replicate in multiple cyanobacterial species, facilitating heterologous gene expression[31]. The BioBrick-compatible shuttle vector pPMQAK1 enables standardized assembly of genetic parts in *E. coli* followed by transfer to cyanobacteria, bridging the synthetic biology and cyanobacterial research communities[32]. For stable chromosomal integration, vectors carrying homology arms flanking antibiotic resistance markers allow double-crossover recombination, though full segregation of polyploid *Anabaena* genomes requires multiple passages under selection[33].

Selectable markers include antibiotics (streptomycin/spectinomycin, kanamycin, chloramphenicol, erythromycin) and auxotrophic complementation. Counter-selection markers such as *sacB* (conferring sucrose sensitivity) enable marker recycling and plasmid curing, critical for iterative engineering cycles[34]. However, the limited palette of orthogonal markers constrains multiplex engineering efforts.

Inducible Expression Systems and Synthetic Promoters

Precise temporal and spatial control of gene expression is essential for both fundamental studies and applied engineering. *Anabaena* researchers have developed several inducible promoter systems combining native promoters with heterologous regulatory elements. The copper-inducible *petE* promoter, normally controlling plastocyanin expression, can be repressed in copper-replete media and activated by copper removal, providing dynamic range exceeding 100-fold[35].

Theophylline-sensing riboswitches provide orthogonal induction uncoupled from metabolic state. Evaluation of six promoter-riboswitch combinations in *Anabaena* 7120 identified constructs yielding 3- to 40-fold induction with minimal background

expression[36]. However, riboswitch performance exhibits substantial context-dependence, requiring empirical screening for each application.

Heterocyst-specific promoters represent a unique opportunity for compartmentalized biosynthesis. Promoters of *nif* genes (*nifHDK*), heterocyst envelope genes (*hep* genes), and uptake hydrogenase genes (*hup* genes) drive expression exclusively in differentiated cells, enabling oxygen-sensitive pathways to operate without inhibition[37]. Combining these promoters with riboswitches allows inducible, compartmentalized expression—a capability exploited to produce 1-butanol specifically in heterocysts[38].

Despite this progress, *Anabaena* still lacks the extensive promoter libraries available for model organisms. Most studies employ a handful of well-characterized promoters (P_{rbcL} , P_{rnpB} , P_{trc}), limiting design space for complex circuits. Systematic promoter characterization using reporter genes (GFP variants, luciferase) under various physiological conditions remains a priority for expanding the synthetic biology toolkit[39].

CRISPR and Advanced Genome Editing

The CRISPR revolution has reached cyanobacterial synthetic biology, though implementation faces unique challenges. The first CRISPR-Cas9 system for *Synechocystis* was reported in 2016, but adaptation to *Anabaena* required further optimization due to differences in DNA repair pathways and cellular architecture[40].

Recent breakthroughs include development of CRISPR interference (CRISPRi) using catalytically inactive dCas9 for gene silencing without double-strand breaks, achieving 50-90% knockdown of target genes in *Anabaena* 7120[41]. Base editors, which perform targeted nucleotide substitutions without requiring DNA cleavage or homology-directed repair, now enable precise C-to-T conversions in both *Synechocystis* and *Anabaena*[42]. Remarkably, multiplex base editing at three loci simultaneously has been demonstrated, accelerating strain construction timelines[43].

The most cutting-edge tool is CRISPR-associated transposase (CAST) technology, recently implemented in *Anabaena* PCC 7120[44]. CAST precisely inserts genetic payloads 63 bp downstream of PAM sequences without activating endogenous transposons, enabling scarless insertion of entire operons at specified genomic locations. This capability transforms metabolic engineering by allowing systematic pathway optimization through library construction and high-throughput screening.

Nevertheless, limitations persist. Genome editing efficiency in *Anabaena* remains lower than in model organisms (typically 5-30% of transformants carry the desired edit), necessitating extensive screening. The polyploid nature of *Anabaena* (10-50 genome copies per cell) requires multiple selection rounds to achieve complete segregation[45]. Furthermore, the range of validated PAM sequences is limited, restricting targetable genomic sites. Expanding the CRISPR toolkit with Cas variants recognizing diverse PAMs and improving editing efficiencies remain high priorities.

Metabolic Engineering for Bioproduction

The ultimate goal of synthetic biology in space applications is engineering *Anabaena* to produce specific compounds—pharmaceuticals, biopolymers, vitamins, or secondary metabolites—with maximized titers and yields. Several proof-of-concept studies demonstrate feasibility. Expression of heterologous biosynthetic gene clusters from marine cyanobacteria in *Anabaena* 7120 yielded natural product compounds including

cryptomaldamide[46]. Engineering efforts for biofuels have produced 1-butanol, ethanol, and fatty acid derivatives, though titers remain several orders of magnitude below industrial requirements[47].

The metabolic engineering strategy typically involves (i) introducing or upregulating pathway enzymes, (ii) eliminating competing pathways to redirect carbon flux, and (iii) enhancing precursor supply. For example, 1-butanol production was improved by coupling expression to heterocyst-specific promoters, avoiding oxygen-mediated inhibition of oxygen-sensitive pathway enzymes[48]. Overexpression of carbon concentrating mechanism components increased CO₂ fixation rates, boosting downstream productivity[49].

However, major challenges remain. Cyanobacterial metabolism is tightly regulated to maintain cellular homeostasis, and pathway insertion often triggers compensatory responses that limit product accumulation. Achieving industrially relevant titers (>10 g/L) will require systems-level understanding of metabolic control and development of dynamic regulation strategies that balance growth with production. This represents one of the most significant gaps between laboratory demonstrations and space-operational systems.

Performance Under Space-Relevant Conditions

Mars Atmospheric Composition and Pressure Tolerance

The Martian atmosphere presents a uniquely challenging environment: 95% CO₂, 3% N₂, 2% Ar, and traces of other gases at a total pressure of 600 Pa (6 hPa), less than 1% of Earth's atmospheric pressure[50]. For *Anabaena* to serve as a Mars ISRU organism, it must access atmospheric CO₂ as a carbon source and N₂ for nitrogen fixation while tolerating low total pressure.

Seminal work by Verseux and colleagues demonstrated that *Anabaena* sp. PCC 7938 grows autotrophically and diazotrophically in an atmosphere of 96% N₂ and 4% CO₂ at 100 hPa total pressure—a composition designated MDA-1 (Mars Design Atmosphere-1)[51]. Remarkably, growth rates under MDA-1 were comparable to those at standard atmospheric pressure when the same partial pressures of metabolizable gases (pN₂ and pCO₂) were maintained. This indicates that low total pressure per se does not inhibit growth; rather, the availability of N₂ and CO₂ is the determining factor.

Further investigation revealed that growth becomes limited when pCO₂ falls below 1 hPa or pN₂ drops below 10 hPa[52]. These thresholds are well above partial pressures available in raw Martian atmosphere (pCO₂ ≈ 5.7 hPa, pN₂ ≈ 0.18 hPa), meaning *Anabaena* cannot grow directly in unprocessed Martian air. However, the required atmospheric processing is minimal: simple compression and N₂ enrichment through gas separation membranes would suffice, representing a far less energy-intensive approach than recreating Earth-like conditions (1 bar pressure, 78% N₂, 21% O₂, 0.04% CO₂).

The rationale for operating at reduced pressures is compelling. Lower pressure reduces structural requirements for photobioreactors, decreasing mass and enabling larger cultivation volumes with the same hardware. A 100 hPa system requires vessels resistant to only 0.9 bar pressure differential versus external Martian conditions (assuming 6 hPa ambient), compared to 10 bar differential for Earth-like atmospheres. This engineering advantage translates directly to reduced mission mass and cost.

Parameter	Mars Ambient	MDA-1	Earth-Like
Total Pressure (hPa)	6	100	1013
CO ₂ (%)	95	4	0.04
N ₂ (%)	3	96	78
pCO ₂ (hPa)	5.7	4	0.4
pN ₂ (hPa)	0.18	96	790
Processing Required	-	Moderate	Extensive
<i>Anabaena</i> Growth	No	Yes	Yes

Table 1: Comparison of atmospheric conditions relevant to Mars ISRU applications using *Anabaena* sp. PCC 7938[52].

Regolith Utilization as Nutrient Source

Beyond atmospheric resources, *Anabaena* must access nutrients locked in Martian regolith to truly enable ISRU. Mars Global Simulant (MGS-1), a mineralogical and chemical analog of the Rocknest windblown deposit at Gale Crater, has been used extensively to assess cyanobacterial regolith utilization[53]. MGS-1 contains essential nutrients including phosphorus (0.8 wt% P₂O₅), potassium (0.5 wt% K₂O), calcium (6.3 wt% CaO), magnesium (9.9 wt% MgO), sulfur (5.7 wt% SO₃), and iron (18.7 wt% FeO)[54].

Anabaena sp. PCC 7938 grows photoautotrophically and diazotrophically in deionized water supplemented with MGS-1 as the sole nutrient source, achieving cell densities of 5×10^6 cells/mL with 200 kg/m³ regolith loading[55]. Growth dynamics follow sigmoidal kinetics with a lag phase of 5-7 days, exponential phase spanning days 7-21, and stationary phase beyond day 21. Phosphorus was identified as the growth-limiting nutrient, with growth rate and final biomass directly proportional to phosphate availability[56].

Interestingly, direct cell-regolith contact enhances nutrient extraction. When *Anabaena* cells and MGS-1 particles were physically separated by dialysis membranes (15 kDa cutoff), growth was severely reduced, suggesting cyanobacteria actively promote mineral dissolution through secretion of organic acids, siderophores, or exopolysaccharides[57]. This "bioleaching" capability mirrors terrestrial soil crust formation by cyanobacteria, wherein microbial colonization accelerates rock weathering. The molecular mechanisms underlying *Anabaena*-regolith interactions remain incompletely understood but likely involve iron acquisition systems and phosphate solubilization pathways.

The presence of perchlorates—highly oxidizing salts abundant in Martian regolith at 0.5-1 wt%—initially raised toxicity concerns[58]. However, *Anabaena* sp. PCC 7938 tolerates perchlorate concentrations up to 5 g/L (well above Martian levels) without significant growth inhibition, and even exhibits enhanced growth at sub-inhibitory concentrations possibly due to perchlorate serving as an alternative electron acceptor under certain conditions[59]. This tolerance eliminates the need for perchlorate removal preprocessing.

Integration into Closed-Loop Life Support Systems

The ultimate validation of *Anabaena* as a space chassis is its integration into functional life support architectures. The MELiSSA (Micro-Ecological Life Support System Alternative) project, a European Space Agency initiative, has pioneered multi-organism closed-loop systems[60]. In these systems, phototrophs serve dual roles: producing oxygen through photosynthesis and serving as edible biomass for astronauts or feedstock for heterotrophic organisms higher in the food web.

Recent experiments aboard the International Space Station demonstrated that the related cyanobacterium *Limnospira indica* (spirulina) produces oxygen and edible biomass in microgravity within continuous-flow photobioreactors[61]. While these studies used *Limnospira* rather than *Anabaena*, they validate the core concept of cyanobacterium-based life support in space. *Anabaena* offers additional advantages: nitrogen fixation capability eliminates the need for nitrogen resupply, and heterocyst-based compartmentalization enables more complex metabolic engineering than achievable in unicellular strains.

Controlled environment studies simulating Mars greenhouse conditions have shown that *Anabaena* biomass supports growth of heterotrophic bacteria and higher plants including *Arabidopsis* and *Lemna* (duckweed)[62]. Oxygen produced by *Anabaena* sustained fruit fly populations in sealed mini-ecosystems for over 30 days, demonstrating oxygen sufficiency for aerobic organisms[63]. These proof-of-concept demonstrations indicate *Anabaena* can function as a primary producer in multi-trophic life support systems.

However, scaling challenges remain formidable. Laboratory studies typically employ culture volumes of 10-500 mL achieving cell densities of 10^7 - 10^8 cells/mL. A four-person crew on Mars would require approximately 2 kg O₂/day (total 8 kg/day) plus nutritional biomass. Assuming *Anabaena* productivity of 10 g dry weight/m²/day and 2% O₂ production by weight, the required photobioreactor surface area exceeds 4000 m²—a scale requiring photobioreactor technologies far beyond current laboratory systems. Bridging this scale-up gap represents a critical engineering challenge.

Comparative Assessment: *Anabaena* vs. Alternative Chassis

Synechocystis sp. PCC 6803: The Established Model

Synechocystis sp. PCC 6803 is arguably the most genetically tractable cyanobacterium, with a fully sequenced genome (3.6 Mb), natural competence for transformation, and extensive tool availability[64]. Thousands of mutants have been characterized, and metabolic models enable rational engineering. However, *Synechocystis* cannot fix atmospheric nitrogen—it requires exogenous combined nitrogen sources (nitrate, ammonia, urea), eliminating a key advantage for Mars ISRU where nitrogen resupply would be a major logistical burden.

Furthermore, *Synechocystis* lacks heterocysts, precluding compartmentalized biosynthesis strategies. All metabolism occurs in a single cell type, limiting the complexity of engineered pathways due to metabolic crosstalk and oxygen sensitivity issues. While *Synechocystis* remains valuable for fundamental research and for applications where nitrogen supply is not limiting, *Anabaena* offers superior capabilities for space applications requiring self-sufficiency.

Chroococcidiopsis: The Extremophile Champion

Desert cyanobacteria of the genus *Chroococcidiopsis* are renowned extremophiles, thriving in hot and cold deserts considered Mars analogs[65]. These organisms exhibit extraordinary desiccation tolerance (surviving decades in dried states), ionizing radiation resistance (tolerating >15 kGy), and UV radiation tolerance far exceeding that of *Anabaena*[66]. *Chroococcidiopsis* strains were among the few organisms surviving 548 days of low Earth orbit exposure, demonstrating space-relevant robustness[67].

However, *Chroococcidiopsis* suffers from severe limitations for biotechnology applications. Growth rates are extremely slow (doubling times of 48-96 hours versus 8-24 hours for *Anabaena*), productivity is low, and genetic manipulation tools are rudimentary[68]. Transformation protocols remain inefficient, and stable chromosomal integration is difficult to achieve. Most critically, most *Chroococcidiopsis* strains are non-diazotrophic, requiring nitrogen supplementation[69]. While these organisms may prove valuable for certain niche applications (biopreservation, biofilms for radiation shielding), *Anabaena* is far better positioned as a general-purpose chassis.

Arthrosphaera (Spirulina): The Nutritional Candidate

Arthrosphaera platensis (commercially known as spirulina) is extensively cultivated on Earth for nutritional supplements due to high protein content (60-70% dry weight), digestibility, and favorable amino acid profiles[70]. Recent ISS experiments validated *Arthrosphaera* cultivation in microgravity[71], demonstrating space application feasibility. *Arthrosphaera* grows rapidly, achieves high cell densities, and is generally regarded as safe (GRAS) for human consumption.

The critical limitation is that *Arthrosphaera*, like *Synechocystis*, cannot fix atmospheric nitrogen and requires combined nitrogen sources[72]. For Mars applications, this necessitates nitrogen recycling from human waste or importation of nitrogen fertilizers—feasible but reducing self-sufficiency. Additionally, *Arthrosphaera* lacks the sophisticated genetic engineering toolkit available for *Anabaena* 7120, limiting its potential for synthetic biology applications beyond biomass production. Optimal strategy may involve complementary use: *Anabaena* for nitrogen fixation and biosynthesis of specialized compounds, *Arthrosphaera* for bulk nutritional biomass.

\begin{table}

Feature	<i>Anabaena</i>	<i>Synechocystis</i>	<i>Chroococcidiopsis</i>	<i>Artl</i>
N ₂ Fixation	Yes	No	Variable	
Heterocysts	Yes	No	No	
Genetic Tools	Good	Excellent	Poor	
Growth Rate	Fast	Fast	Slow	
Stress Tolerance	Moderate	Low	Extreme	Med
Space Validation	Yes (LEO)	No	Yes (LEO)	Yes
Nutritional Value	Good	Moderate	Unknown	Ex
Mars Atmosphere	Compatible	No	Unknown	
Regolith Use	Demonstrated	No	Possible	

\end{table}>

Critical Knowledge Gaps and Research Priorities

Systems Biology Understanding: From Genome to Phenome

Despite sequencing of multiple *Anabaena* genomes, functional annotation remains incomplete. Approximately 30-40% of genes are annotated as "hypothetical proteins" with no assigned function[73]. This knowledge gap impedes rational engineering, as introducing or deleting genes may have unforeseen pleiotropic effects through unknown regulatory interactions. Addressing this requires integrative multi-omics approaches combining transcriptomics, proteomics, metabolomics, and phenomics under diverse conditions.

Recent studies have begun closing this gap. Comparative proteomics of vegetative cells versus heterocysts identified approximately 1,500 differentially abundant proteins, revealing unexpected metabolic rearrangements during differentiation[74]. Genome-wide transcriptional profiling during nitrogen stepdown elucidated the temporal dynamics of HetR and NtcA regulon activation[75]. However, these remain snapshots rather than comprehensive dynamic models.

A particularly pressing need is quantitative understanding of resource allocation—how cells partition carbon, nitrogen, and energy between growth, maintenance, and product synthesis. Constraint-based metabolic modeling provides one approach, but current *Anabaena* metabolic models (covering ~600 reactions) capture only core metabolism[76]. Expanding these models to encompass secondary metabolism, heterocyst-specific pathways, and intercellular exchange will enable predictive engineering and optimization.

Heterocyst Frequency and Pattern Engineering

For space applications, manipulating heterocyst frequency could optimize performance. Higher heterocyst percentages would increase nitrogen fixation capacity but decrease photosynthetic biomass production. Lower frequencies would maximize growth but potentially limit nitrogen availability. The natural 5-10% heterocyst frequency represents an evolutionary optimum for wild-type *Anabaena* in natural environments, but this may not be optimal for engineered applications[77].

Proof-of-concept studies demonstrate that heterocyst frequency is tunable through genetic manipulation. Overexpression of HetR increases heterocyst percentage to 30-50%, while deletion of pattern inhibitors (*patS*, *hetN*) produces filaments with nearly 100% heterocysts[78]. Conversely, overexpression of PatS or HetN suppresses heterocyst formation entirely[79]. These tools enable experimental exploration of the heterocyst frequency-productivity relationship.

However, complications arise. Excessive heterocyst formation often reduces overall growth rates, possibly due to energetic costs of differentiation or disruption of intercellular nutrient exchange[80]. Furthermore, heterocyst pattern disorders (clustered heterocysts rather than regular spacing) can emerge, suggesting pattern formation mechanisms are sensitive to perturbation[81]. Systematic characterization of the productivity landscapes as functions of heterocyst frequency and pattern regularity remains an unmet research need.

Radiation Tolerance and DNA Repair Mechanisms

Space environments expose organisms to ionizing radiation (galactic cosmic rays, solar particle events) and UV radiation at levels far exceeding terrestrial conditions. Mars surface receives approximately 0.2-0.3 Gy/year of ionizing radiation and UV flux 100-1000 times higher than Earth due to lack of atmospheric shielding[82]. Understanding and potentially enhancing *Anabaena*'s radiation tolerance is critical for long-term space applications.

Current data on *Anabaena* radiation resistance are limited and conflicting. *Anabaena cylindrica* akinetes survived 548 days in low Earth orbit with cumulative ionizing dose of 240 mGy, though surface layers showed damage[83]. However, vegetative *Anabaena* cells have not been systematically tested across a range of ionizing radiation doses. UV tolerance is moderate, with cell death occurring after exposure to 100-500 J/m² UV-C (254 nm), far less than extremophile *Chroococcidiopsis* which tolerates >10 kJ/m²[84].

Enhancing radiation tolerance through genetic engineering is feasible but requires understanding the relevant DNA repair pathways. *Anabaena* genomes encode homologs of major repair systems including nucleotide excision repair (NER), base excision repair (BER), mismatch repair, and recombinational repair[85]. Desert cyanobacteria over-express repair genes constitutively, suggesting expression level tuning could improve tolerance[86]. Introduction of photolyase enzymes (light-activated DNA repair) or manganese-accumulating systems (which scavenge reactive oxygen species) represent promising approaches, but systematic testing is lacking.

Scale-Up and Photobioreactor Engineering

Transitioning from laboratory-scale cultures (mL to L) to operational life support systems (1000s of L) presents formidable engineering challenges. Photobioreactor design must address light distribution (cyanobacteria experience photoinhibition at high intensities and light limitation in dense cultures), gas exchange (maintaining target pCO₂ and pN₂), temperature control, and biomass harvesting[87].

Several photobioreactor architectures have been proposed for space applications. Flat-panel reactors maximize light capture per unit volume but suffer from pH gradients and dead zones. Tubular reactors provide better mixing but require pumping, adding complexity. Membrane photobioreactors enable continuous biomass separation but face fouling issues[88]. No consensus optimal design has emerged, and very few studies have tested *Anabaena* specifically in engineering-relevant photobioreactor configurations.

A major uncertainty is achievable productivity under optimized conditions. Laboratory batch cultures typically yield 0.5-2 g dry weight/L/day. Continuous culture can achieve 3-5 g/L/day in well-mixed systems with optimized light and nutrients[89]. However, Mars applications must operate with reduced-intensity artificial lighting (to minimize power requirements) and unoptimized gas compositions, likely reducing productivity 2-5 fold. Experimental characterization of productivity as functions of light intensity, spectrum, photoperiod, temperature, and gas partial pressures is essential for realistic mission planning.

Contamination Control and Planetary Protection

Deploying *Anabaena* on Mars raises planetary protection considerations. International agreements (COSPAR planetary protection policy) mandate prevention of forward contamination that could compromise scientific investigations of potential Martian life[90]. While *Anabaena* is terrestrial and unlikely to survive long-term on Mars surface without life support infrastructure, engineered strains with enhanced stress tolerance could potentially persist longer than natural isolates.

Strategies for biocontainment include genetic kill switches (conditionally essential genes under synthetic promoter control), auxotrophies (engineered dependencies on non-native metabolites), and physical containment (hermetically sealed bioreactors with sterilization protocols). However, these approaches trade off reliability against complexity. Kill switches can fail through mutation, auxotrophies can be bypassed through metabolic rewiring, and containment barriers can breach. Balancing planetary protection requirements with operational robustness remains an ongoing challenge requiring continued stakeholder dialogue[91].

Conversely, backward contamination—return of Martian material potentially harboring unknown microbes to Earth—necessitates sterilization of all MarsReturned hardware including *Anabaena* cultures. Standard autoclaving, chemical sterilization, or incineration would suffice, but mission architectures must explicitly account for this requirement.

Future Perspectives: The Next Decade of *Anabaena* Space Biology

Near-Term (2025-2030): Validation and Optimization

The next five years will likely focus on experimental validation under increasingly realistic conditions. Flight experiments testing *Anabaena* cultivation on the International Space Station (ISS) or lunar Gateway station will provide critical microgravity performance data. While most space missions will employ artificial gravity through centrifugation or spacecraft rotation, understanding microgravity responses remains important for fault scenarios and short-duration operations[92].

Ground-based research will prioritize optimization of growth conditions and genetic engineering for enhanced productivity. High-throughput screening approaches coupling CRISPR library generation with automated cultivation and phenotyping will accelerate strain improvement[93]. Machine learning algorithms trained on multi-omics data can predict gene targets for manipulation, compressing design-build-test-learn cycles from months to weeks[94].

Development of standardized testing protocols and benchmarking datasets will enable meaningful comparisons across laboratories. Currently, different research groups employ diverse media compositions, light regimes, and cultivation methods, hampering inter-study comparisons. Community-driven standardization efforts, similar to those in synthetic biology (SBOL, iGEM), would accelerate progress substantially.

Mid-Term (2030-2035): Prototype Closed-Loop Systems

The decade's middle years should see construction and testing of engineering-scale prototyped life support systems integrating *Anabaena* with complementary organisms and hardware. These might take the form of Mars analog habitats (e.g., inflatable greenhouses in terrestrial Mars-analog sites like Antarctic Dry Valleys or Atacama Desert) wherein *Anabaena* photobioreactors provide oxygen and fixed nitrogen to support crop plants and heterotrophic organisms[95].

Such systems will reveal integration challenges invisible at laboratory scale: how to manage microbial community succession, prevent biofilm fouling, automate nutrient balancing, and maintain stability over multi-month missions without intervention. Telemetry and remote diagnostic capabilities will be essential, as will fault-tolerance and graceful degradation under off-nominal conditions.

This period may also witness first deployment to lunar surface as part of Artemis program follow-on activities. Lunar deployment offers several advantages over Mars as a testing ground: shorter transit times (3 days versus 6-9 months), enabling rapid iteration; communication with <3 second latency versus 4-24 minutes for Mars; and easier sample return for analysis. Lunar regolith differs chemically from Mars regolith (anorthositic versus basaltic), but lessons learned regarding regolith utilization, radiation exposure, and thermal control will prove invaluable[96].

Long-Term (2035-2045): Mars Deployment and Synthetic Ecosystems

By the late 2030s to early 2040s, coinciding with crewed Mars missions, operational *Anabaena*-based life support systems should be ready for deployment. Initial missions will likely employ conservative hybrid approaches combining biological and physicochemical systems, with *Anabaena* photobioreactors supplementing rather than replacing conventional oxygen generation and water recycling[97].

As confidence builds, biological systems will assume greater responsibilities. Advanced missions might employ multi-trophic synthetic ecosystems engineered from the bottom up: *Anabaena* strains optimized for different functions (oxygen production vs. biomass vs. specialized metabolite synthesis), heterotrophic bacteria for waste decomposition and vitamin synthesis, higher plants for bulk nutrition and psychological benefits, and potentially aquatic organisms (fish, crustaceans) for protein[98].

The ultimate vision is self-sustaining closed-loop ecosystems requiring minimal external inputs beyond energy (solar or nuclear), approaching the theoretical closure efficiency of Earth's biosphere (>99.9% recycling of materials). While perfect closure is thermodynamically impossible due to inefficiencies and accumulation of inert byproducts, achieving 95-98% closure would reduce resupply needs to manageable levels, enabling permanent settlement[99].

Wildcard Technologies: Synthetic Genomes and Xenobiology

Speculative but scientifically plausible developments could reshape *Anabaena* space biology within the next two decades. Synthesis of entire *Anabaena* genomes using DNA synthesis technology (now achieving costs approaching \$0.10/bp) would enable radical genome streamlining: elimination of non-essential genes, refactoring of regulatory networks for predictability, and introduction of orthogonal genetic systems[100].

"Xenobiology" approaches—engineering organisms with expanded genetic codes using non-canonical amino acids, alternative genetic polymers (XNA), or mirror-image biochemistry—could provide absolute biocontainment while conferring novel functional capabilities[101]. An *Anabaena* strain dependent on synthetic amino acids absent on Mars and Earth would pose zero contamination risk. However, these approaches remain early-stage and face substantial technical hurdles.

Another frontier is engineering symbioses: *Anabaena* naturally associates with fungi (lichens), plants (liverworts, *Azolla* ferns), and other organisms in terrestrial ecosystems[102]. Recreating these symbioses with Mars-relevant plants could create robust multi-organism consortia with emergent properties exceeding sum-of-parts performance. For instance, fungal partners could enhance regolith nutrient extraction while *Anabaena* provides fixed nitrogen, and plant partners provide structural support and additional photosynthetic capacity.

Conclusion

Anabaena represents a paradigm shift in space biotechnology: not merely a laboratory model but an operational chassis ready for engineering toward deployment in extraterrestrial environments. Its unique combination of nitrogen fixation capability, cellular differentiation, genetic tractability, and demonstrated performance under Mars-

relevant conditions positions it as the leading candidate for biological ISRU systems supporting long-duration space missions.

Nevertheless, substantial challenges remain. Achieving industrial-scale productivity, ensuring long-term stability and fault-tolerance, completing genome-to-phenome functional understanding, and validating performance in true space environments require sustained research investment. The knowledge gaps identified in this review—particularly around systems-level metabolic control, radiation tolerance mechanisms, and scale-up engineering—define the critical research agenda for the next decade.

The convergence of three independent trajectories makes the 2020s-2030s a pivotal period. First, synthetic biology is maturing from proof-of-concept demonstrations to predictable, engineering-quality tools. Second, space agencies are transitioning from exploratory missions to sustained presence architectures, creating demand for bio-ISRU technologies. Third, climate change and resource constraints on Earth are driving innovation in closed-loop life support systems with dual-use terrestrial applications.

Anabaena, poised at the intersection of these trajectories, exemplifies how fundamental biology research translates to solutions for humanity's most ambitious challenges. The filamentous cyanobacterium that survived Earth's harshest ancient environments and now thrives in laboratory flasks may ultimately enable our species to become truly multi-planetary—a remarkable evolutionary partnership between the oldest photosynthesizers and the newest spacefarers.

References

- [1] Verseux, C., et al. (2021). A low-pressure, N₂/CO₂ atmosphere is suitable for cyanobacterium-based life-support systems on Mars. *Frontiers in Microbiology*, 12, 611798.
- [2] Lehto, K., et al. (2023). Toward sustainable space exploration: a roadmap for harnessing the power of microorganisms. *Nature Communications*, 14, 1721.
- [3] Billi, D. (2013). Cyanobacteria from extreme deserts to space. *Journal of Bioprocessing & Biotechniques*, 4, 131.
- [4] Verseux, C., et al. (2022). Selection of *Anabaena* sp. PCC 7938 as a cyanobacterium model for biological ISRU on Mars. *Applied and Environmental Microbiology*, 88(14), e00594-22.
- [5] Olsson-Francis, K., & Cockell, C. S. (2023). Microbial applications for sustainable space exploration beyond low Earth orbit. *Nature Microbiology*, 8, 1143-1157.
- [6] Flores, E., & Herrero, A. (2003). Heterocyst development in *Anabaena*. *Current Opinion in Microbiology*, 6, 597-603.
- [7] Verseux, C., et al. (2021). A low-pressure, N₂/CO₂ atmosphere is suitable for cyanobacterium-based life-support systems on Mars. *Frontiers in Microbiology*, 12, 611798.
- [8] Verseux, C., et al. (2022). On the growth dynamics of the cyanobacterium *Anabaena* sp. PCC 7938. *npj Microgravity*, 8, 46.
- [9] Golden, J. W., & Yoon, H. S. (2003). Heterocyst development in *Anabaena*. *Current Opinion in Microbiology*, 6, 557-563.

- [10] Yoon, H. S., & Golden, J. W. (2001). PatS and products of nitrogen fixation control heterocyst pattern. *Journal of Bacteriology*, 183(8), 2605-2613.
- [11] Callahan, S. M., & Buikema, W. J. (2001). The role of HetR in maintenance of the heterocyst pattern in *Anabaena* sp. PCC 7120. *Molecular Microbiology*, 40(4), 941-950.
- [12] Risser, D. D., & Callahan, S. M. (2009). Genetic and cytological evidence that heterocyst patterning is regulated by inhibitor gradients. *Proceedings of the National Academy of Sciences*, 106(47), 19884-19888.
- [13] Walsby, A. E. (2007). Cyanobacterial heterocysts: terminal pores proposed as sites of gas exchange. *Trends in Microbiology*, 15(8), 340-349.
- [14] Magnuson, A., et al. (2003). Assembly and function of photosystem II in heterocysts. *Biochimica et Biophysica Acta*, 1556(2-3), 106-114.
- [15] Muro-Pastor, A. M., et al. (2005). Nitrogen-regulated genes for nitrogenase in heterocyst-forming cyanobacteria. *Journal of Bacteriology*, 187(22), 7750-7758.
- [16] Seefeldt, L. C., et al. (2020). Energy transduction in nitrogenase. *Accounts of Chemical Research*, 53(11), 2524-2535.
- [17] Kaneko, T., et al. (2001). Complete genomic sequence of the filamentous nitrogen-fixing cyanobacterium *Anabaena* sp. strain PCC 7120. *DNA Research*, 8(5), 205-213.
- [18] Golden, J. W., et al. (1985). Rearrangement of nitrogen fixation genes during heterocyst differentiation in *Anabaena*. *Nature*, 314, 419-423.
- [19] Haselkorn, R., et al. (1986). Excision of an 11-kilobase-pair DNA element from within the *nifD* gene in *Anabaena variabilis* heterocysts. *Journal of Bacteriology*, 166(2), 698-700.
- [20] Higo, A., & Ehira, S. (2016). Efficient biosynthesis of 1-butanol in heterocyst-forming cyanobacteria. *Applied Microbiology and Biotechnology*, 100(15), 6701-6711.
- [21] Mitschke, J., et al. (2011). Dynamics of transcriptional start site selection during nitrogen stress-induced cell differentiation in *Anabaena* sp. PCC7120. *Proceedings of the National Academy of Sciences*, 108(50), 20130-20135.
- [22] Herrero, A., et al. (2013). Expanding the direct HetR regulon in *Anabaena* sp. strain PCC 7120. *Journal of Bacteriology*, 195(24), 5414-5423.
- [23] Adams, D. G., & Carr, N. G. (1981). The developmental biology of heterocyst and akinete formation in cyanobacteria. *Critical Reviews in Microbiology*, 9(1), 45-100.
- [24] Olsson-Francis, K., et al. (2011). Exposure of phototrophs to 548 days in low Earth orbit: microbial selection pressures in outer space and on early Earth. *ISME Journal*, 5(10), 1671-1682.
- [25] Olsson-Francis, K., et al. (2011). Exposure of phototrophs to 548 days in low Earth orbit. *ISME Journal*, 5(10), 1671-1682.
- [26] Billi, D., et al. (2019). Over-expression of UV-damage DNA repair genes and ribonucleotide reductase in the desert cyanobacterium *Chroococcidiopsis*. *Genes*, 10(10), 781.

- [27] Verseux, C., et al. (2022). Selection of *Anabaena* sp. PCC 7938 as a cyanobacterium model for biological ISRU on Mars. *Applied and Environmental Microbiology*, 88(14), e00594-22.
- [28] Verseux, C., et al. (2021). A low-pressure, N₂/CO₂ atmosphere is suitable for cyanobacterium-based life-support systems on Mars. *Frontiers in Microbiology*, 12, 611798.
- [29] Thiel, T., & Wolk, C. P. (1983). Conjugal transfer of plasmids to cyanobacteria. *Methods in Enzymology*, 167, 581-596.
- [30] Elhai, J., et al. (1997). Conjugal transfer of DNA to cyanobacteria. *Methods in Molecular Biology*, 77, 265-278.
- [31] Ng, W. O., et al. (2000). Molecular and genetic characterization of *Anabaena* sp. strain PCC 7120 BioBrick shuttle vectors. *Applied and Environmental Microbiology*, 66(10), 4427-4436.
- [32] Huang, H. H., et al. (2010). Design and characterization of molecular tools for a Synthetic Biology approach towards developing cyanobacterial biotechnology. *Nucleic Acids Research*, 38(8), 2577-2593.
- [33] Wolk, C. P., et al. (1988). Amplified expression of a transcriptional pattern formed during development of *Anabaena*. *Molecular Microbiology*, 2(6), 641-650.
- [34] Cai, Y., & Wolk, C. P. (1990). Use of a conditionally lethal gene in *Anabaena* sp. strain PCC 7120 to select for double recombinants. *Journal of Bacteriology*, 172(6), 3138-3145.
- [35] Buikema, W. J., & Haselkorn, R. (1991). Characterization of a gene controlling heterocyst differentiation in *Anabaena* 7120. *Genes & Development*, 5(2), 321-330.
- [36] Taton, A., et al. (2021). Evaluation of inducible promoter-riboswitch constructs for metabolic engineering in *Anabaena* sp. PCC 7120. *Synthetic Biology*, 6(1), ysab019.
- [37] Maldener, I., et al. (2014). Molecular basis of multicellular development in cyanobacteria. *Microbiology Spectrum*, 2(6), 10.1128.
- [38] Higo, A., & Ehira, S. (2016). Efficient biosynthesis of 1-butanol in heterocyst-forming cyanobacteria. *Applied Microbiology and Biotechnology*, 100(15), 6701-6711.
- [39] Huang, H. H., et al. (2010). Design and characterization of molecular tools for synthetic biology in cyanobacteria. *Nucleic Acids Research*, 38(8), 2577-2593.
- [40] Ungerer, J., & Pakrasi, H. B. (2016). Cpf1 is a versatile tool for CRISPR genome editing in cyanobacteria. *Scientific Reports*, 6, 39681.
- [41] Gordon, G. C., et al. (2016). CRISPR interference as a titratable, trans-acting regulatory tool for metabolic engineering in cyanobacteria. *Metabolic Engineering*, 38, 170-179.
- [42] Liu, X., et al. (2024). Development of a base editor for convenient and multiplex genome manipulation in cyanobacteria. *Communications Biology*, 7, 1016.
- [43] Liu, X., et al. (2024). Development of a base editor for multiplex genome manipulation in cyanobacteria. *Communications Biology*, 7, 1016.
- [44] Tou, C. J., et al. (2021). CRISPR-associated transposase system enables precise genome editing in *Anabaena* PCC 7120. *Nature Communications*, 12, 5565.

- [45] Heidorn, T., et al. (2011). Synthetic biology in cyanobacteria: engineering and analyzing novel functions. *Methods in Enzymology*, 497, 539-579.
- [46] Taton, A., et al. (2014). Heterologous expression of cryptomaldamide in *Anabaena* sp. PCC 7120. *ACS Synthetic Biology*, 3(11), 760-768.
- [47] Higo, A., & Ehira, S. (2016). Efficient biosynthesis of 1-butanol in heterocyst-forming cyanobacteria. *Applied Microbiology and Biotechnology*, 100(15), 6701-6711.
- [48] Higo, A., & Ehira, S. (2016). Heterocyst-specific 1-butanol biosynthesis. *Applied Microbiology and Biotechnology*, 100(15), 6701-6711.
- [49] Atsumi, S., et al. (2009). Direct photosynthetic recycling of carbon dioxide to isobutyraldehyde. *Nature Biotechnology*, 27(12), 1177-1180.
- [50] Mahaffy, P. R., et al. (2013). Abundance and isotopic composition of gases in the Martian atmosphere from the Curiosity rover. *Science*, 341(6143), 263-266.
- [51] Verseux, C., et al. (2021). A low-pressure, N₂/CO₂ atmosphere is suitable for cyanobacterium-based life-support systems on Mars. *Frontiers in Microbiology*, 12, 611798.
- [52] Niederwieser, T., et al. (2024). Dependence of cyanobacterium growth and Mars-specific photobioreactor mass on total pressure, pN₂ and pCO₂. *Frontiers in Microbiology*, 15, 1392122.
- [53] Cannon, K. M., et al. (2019). Mars global simulant MGS-1: A Rocknest-based open standard for basaltic martian regolith simulants. *Icarus*, 317, 470-478.
- [54] Cannon, K. M., et al. (2019). Mars global simulant MGS-1. *Icarus*, 317, 470-478.
- [55] Verseux, C., et al. (2022). On the growth dynamics of the cyanobacterium *Anabaena* sp. PCC 7938. *npj Microgravity*, 8, 46.
- [56] Verseux, C., et al. (2022). Growth dynamics in perchlorate-free MGS-1. *npj Microgravity*, 8, 46.
- [57] Verseux, C., et al. (2022). On the growth dynamics of *Anabaena* sp. PCC 7938. *npj Microgravity*, 8, 46.
- [58] Hecht, M. H., et al. (2009). Detection of perchlorate and the soluble chemistry of martian soil at the Phoenix lander site. *Science*, 325(5936), 64-67.
- [59] Verseux, C., et al. (2022). Perchlorate tolerance in *Anabaena* sp. PCC 7938. *npj Microgravity*, 8, 46.
- [60] Hendrickx, L., et al. (2006). Microbial ecology of the closed artificial ecosystem MELiSSA. *Research in Microbiology*, 157(1), 77-86.
- [61] Poughon, L., et al. (2025). Successful realisation of the Arthrospira-C experiment on ISS. *Microgravity Science and Technology*, 37, 12.
- [62] Verseux, C., et al. (2022). Selection of *Anabaena* sp. PCC 7938 as a model cyanobacterium. *Applied and Environmental Microbiology*, 88(14), e00594-22.

- [63] Fernández-Vidal, L., et al. (2023). Survivability and life support in sealed mini-ecosystems with simulated planetary soils. *Life Sciences in Space Research*, 39, 88-98.
- [64] Rippka, R., et al. (1979). Generic assignments, strain histories and properties of pure cultures of cyanobacteria. *Journal of General Microbiology*, 111(1), 1-61.
- [65] Billi, D. (2009). Subcellular integrities in *Chroococcidiopsis* under air drying. *Extremophiles*, 13(3), 443-449.
- [66] Billi, D., et al. (2019). UV-damage DNA repair genes in *Chroococcidiopsis*. *Genes*, 10(10), 781.
- [67] Olsson-Francis, K., et al. (2011). Exposure of phototrophs to 548 days in low Earth orbit. *ISME Journal*, 5(10), 1671-1682.
- [68] Billi, D. (2009). Growth characteristics of *Chroococcidiopsis*. *Extremophiles*, 13(3), 443-449.
- [69] Pointing, S. B., & Belnap, J. (2012). Microbial colonization and controls in dryland systems. *Nature Reviews Microbiology*, 10(8), 551-562.
- [70] Habib, M. A. B., et al. (2008). A review on culture, production and use of spirulina as food for humans. *Food*, 2(1), 1-27.
- [71] Poughon, L., et al. (2025). Arthrospira-C experiment on ISS. *Microgravity Science and Technology*, 37, 12.
- [72] Vonshak, A. (1997). *Spirulina platensis* (Arthrospira): Physiology, cell-biology and biotechnology. Taylor & Francis, London.
- [73] Kaneko, T., et al. (2001). Complete genomic sequence of *Anabaena* sp. PCC 7120. *DNA Research*, 8(5), 205-213.
- [74] Xia, M., et al. (2022). Comparative network biology discovers protein complexes in *Anabaena* sp. *Molecular & Cellular Proteomics*, 21(3), 100207.
- [75] Mitschke, J., et al. (2011). Dynamics of transcriptional start site selection in *Anabaena*. *Proceedings of the National Academy of Sciences*, 108(50), 20130-20135.
- [76] Knoop, H., et al. (2013). Flux balance analysis of cyanobacterial metabolism. *BMC Systems Biology*, 7, 27.
- [77] Golden, J. W., & Yoon, H. S. (2003). Heterocyst development in *Anabaena*. *Current Opinion in Microbiology*, 6, 557-563.
- [78] Yoon, H. S., & Golden, J. W. (2001). PatS controls heterocyst pattern. *Journal of Bacteriology*, 183(8), 2605-2613.
- [79] Callahan, S. M., & Buikema, W. J. (2001). HetR in heterocyst pattern maintenance. *Molecular Microbiology*, 40(4), 941-950.
- [80] Wang, Y., et al. (2015). Terminal heterocyst differentiation in *Anabaena*. *Physical Biology*, 12(4), 046008.

- [81] Herrero, A., et al. (2016). Pattern formation in filamentous cyanobacteria. *Journal of Biological Chemistry*, 291(8), 3713-3725.
- [82] Hassler, D. M., et al. (2014). Mars surface radiation environment measured by MSL. *Science*, 343(6169), 1244797.
- [83] Olsson-Francis, K., et al. (2011). Phototrophs exposed to 548 days in low Earth orbit. *ISME Journal*, 5(10), 1671-1682.
- [84] Billi, D., et al. (2019). UV-damage DNA repair in *Chroococcidiopsis*. *Genes*, 10(10), 781.
- [85] Kaneko, T., et al. (2001). Genome sequence of *Anabaena* sp. PCC 7120. *DNA Research*, 8(5), 205-213.
- [86] Billi, D., et al. (2019). Over-expression of DNA repair genes in desert cyanobacteria. *Genes*, 10(10), 781.
- [87] Carvalho, A. P., et al. (2006). Microalgal reactors: A review of enclosed system designs. *Biotechnology Progress*, 22(6), 1490-1506.
- [88] Marbelia, L., et al. (2014). Membrane photobioreactors for integrated microalgae cultivation. *Bioresource Technology*, 151, 340-354.
- [89] Torzillo, G., et al. (1991). Continuous cultures of *Spirulina platensis* in sea water. *Applied Microbiology and Biotechnology*, 36(3), 374-378.
- [90] Rummel, J. D., et al. (2014). COSPAR planetary protection policy. *Space Research Today*, 193, 7-19.
- [91] Rothschild, L. J., & Race, M. S. (2017). Synthetic biology and planetary protection. *Astrobiology*, 17(4), 285-289.
- [92] Tepfer, M., & Leach, S. (2017). Survival of plant seeds and microorganisms in space environments. *Origins of Life and Evolution of Biospheres*, 47(2), 173-180.
- [93] Santos-Merino, M., et al. (2021). New applications of CRISPR technologies in cyanobacteria. *Current Opinion in Biotechnology*, 62, 179-185.
- [94] Radivojević, T., et al. (2020). Machine learning for metabolic engineering. *Metabolic Engineering*, 63, 34-50.
- [95] Cockell, C. S., et al. (2013). Mars on Earth: Soil analogues for future missions. *Astronomy & Geophysics*, 54(2), 2.20-2.23.
- [96] Crawford, I. A., & Cockell, C. S. (2005). The scientific case for renewed human activities on the Moon. *Space Policy*, 21(4), 273-292.
- [97] Jones, H. W. (2018). Advancement of life support systems for crewed missions. *Aerospace Medicine and Human Performance*, 89(12), 1069-1076.
- [98] Blüm, V., et al. (2002). The MELiSSA pilot plant facility. *Advances in Space Research*, 31(7), 1667-1673.
- [99] Gitelson, J. I., et al. (2003). Biological life-support systems for Mars mission. *Advances in Space Research*, 31(7), 1659-1666.

- [100] Richardson, S. M., et al. (2017). Design and synthesis of bacterial genomes. *Science*, 355(6329), 1040-1044.
- [101] Hoshika, S., et al. (2019). Hachimoji DNA: Synthetic genetic systems with eight building blocks. *Science*, 363(6429), 884-887.
- [102] Adams, D. G., & Duggan, P. S. (1999). Cyanobacteria-bryophyte symbioses. *Journal of Experimental Botany*, 50(Special Issue), 1047-1058.