



# **Ammonia Production Using Bacteria and Yeast toward a Sustainable Society**

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Abstract: Ammonia is an important chemical that is widely used in fertilizer applications as well as in the steel, chemical, textile, and pharmaceutical industries, which has attracted attention as a potential fuel. Thus, approaches to achieve sustainable ammonia production have attracted considerable attention. In particular, biological approaches are important for achieving a sustainable society because they can produce ammonia under mild conditions with minimal environmental impact compared with chemical methods. For example, nitrogen fixation by nitrogenase in heterogeneous hosts and ammonia production from food waste using microorganisms have been developed. In addition, crop production using nitrogen-fixing bacteria has been considered as a potential approach to achieving a sustainable ammonia economy. This review describes previous research on biological ammonia production and provides insights into achieving a sustainable society.

Keywords: ammonia; sustainability; bacteria; yeast

## 1. Introduction

In recent years, rapid population growth and industrial development have increased the use of several fossil fuels and have increased the amount of waste and environmental pollutants, leading to destruction of the global environment and causing global warming, ocean acidification, and so on [1]. In alleviating this situation, researchers, governments, companies, and other organizations are greatly interested in improving the global environment, and the Sustainable Development Goals have been set forth [2,3]. Fossil fuels are the most significant energy source in the modern world, which are converted to electrical energy primarily through thermal power generation. Various natural energy sources have been used as alternatives to fossil fuels, including wind, hydro, solar, and geothermal energy [4–6]. In addition to these renewables, biomass as fuel has also received increasing attention. Biomass includes perennial plants, forestry waste, algae, and food waste (municipal solid waste), which are also recognized as carbon-neutral fuels. However, the supply of biomass is greatly affected by weather, time, location, and economics, making it unstable [7–13]. In addition, biodegradable materials could be used as sources of biofuels, and ammonia is expected to be used as a biofuel (Figure 1) [14].

### 1.1. Industrial Uses and the Need for Ammonia

Ammonia is an important compound in a variety of industries [15]. Fixed nitrogen, such as ammonia, is essential for crop growth, and increasing the amount of nitrogen circulating on the planet allows for population growth [16]. The production of ammonia is highly important to sustain life, and about 80% of ammonia is used in the production of fertilizers. Ammonia is also used for refrigerant gas or the synthesis of various chemicals such as plastics, explosives (trinitrotoluene, nitroglycerin, and nitrocellulose), textiles (rayon and nylon), agricultural chemicals, dyes (for cotton, wool, silk, etc.), and so on [17–19]. In recent years, ammonia-fueled batteries have also been devised [18,20–24].



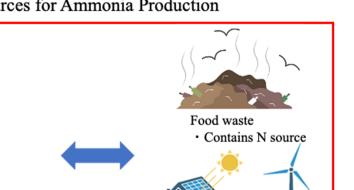
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Solar energy

Green energy

Sustainable

Carbon emission free

Sustainable

Wind power

**Resources for Ammonia Production** 

Figure 1. Sustainable resources of energy for a sustainable society (created with Biorender.com).

The potential application of ammonia as fuel has received increasing attention [8,25–29]. Hydrogen has low transport efficiency because of its low volumetric energy density  $(3 \text{ W h} \cdot \text{L}^{-1})$  and higher calorific value per liquid wight (141.9 MJ/kg) [30]. Therefore, approaches to converting hydrogen to a more transport-efficient substance have been investigated. Ammonia has a flammable range of 16% to 25% (v/v), which can be transported more safely than hydrogen [31,32]. Liquid ammonia contains hydrogen atoms per volume and energy density (MJ  $L^{-1}$ ) that are 1.7 and 1.5 times higher than that of liquid hydrogen, respectively [33,34]. Hydrogen has a low boiling point of -253 °C, and it requires considerable energy to liquefy. By contrast, ammonia has a boiling point of -33.4 °C, and it can be easily liquefied by using general-purpose refrigeration equipment and can be handled easily. Thus, it has been considered for use as a carrier for hydrogen in a hydrogen society [32,35–37]. In addition, the hurdle to the industrial application of ammonia is lower than that of hydrogen because the infrastructure for storage and transportation of ammonia has already been established.

#### 1.2. Chemical Method for Ammonia Production

Fossil fuels

Source limited

Converted into greenhouse gas

Unsustainable

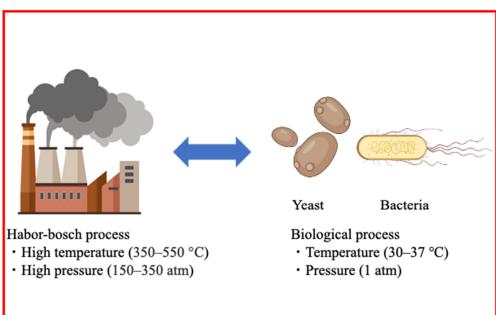
The Haber–Bosch process is a typical example of ammonia nitrogen fixation, and 55% of the world's ammonia is produced by this method [38]. This method requires the cleavage of the triple bond of the nitrogen molecule, which uses a large amount of energy [15,39]. The energy used by the Haber–Bosch process is equivalent to 2–3% of the world's annual fossil fuel use, and it accounts for 1.4% of the global annual carbon dioxide emissions [38,40]. Carbon dioxide is a well-known greenhouse gas (GHG), and its reduction is strongly desired because of environmental issues of growing concern in recent years. Therefore, the Haber–Bosch process has been improved upon in recent years, particularly in catalyst improvements. Moreover, the development of catalysts that allow reactions to proceed under conditions closer to ambient temperature and pressure has been considered. Compared with iron catalysts, Ru-based catalysts enable the fixation of ammonia at lower pressure (90 atm), and they have about 20 times higher catalytic efficiency [41]. However, ruthenium has become increasingly expensive in the last decade, thereby hindering its industrial use [42–44]. The performance of catalysts has been greatly studied and further developed, including the development of several molybdenum-based catalysts that mimic

the active center of nitrogenase (a nitrogen-fixing enzyme) for the synthesis of ammonia at ambient temperature and for pressure in microorganisms [45,46].

At present, most of the hydrogen required by the Haber–Bosch process is obtained by electrolysis. In general, the operation of a water electrolysis unit requires a continuous supply of highly purified water. Furthermore, 9 tons of highly purified water is required to produce 1 ton of hydrogen. Based on these data, 233.6 million tons of water per year is required to produce 1 ton of ammonia using hydrogen obtained from water electrolysis [47]. With the progression of global warming, improving the global environment is necessary [2]. Therefore, attempts are being made to supply ammonia in a sustainable manner using natural energy (green energy) such as wind and solar power generation [48,49]. Ammonia using hydrogen produced from green energy is known as "green ammonia" [50].

In addition to wood biomass, the amount of food waste has continued to grow excessively in recent years, particularly in developed countries [51,52]. Food waste produces considerable amounts of GHGs and environmental pollutants through landfilling and incineration [53]. Thus, many attempts have been made to obtain energy from food waste [13]. In particular, okara is an abundant food waste, and its various uses are being considered [54,55]. Attempts to produce ammonia from food waste by physicochemical methods have also been studied [56–58]. For example, considering that glutamic acid is a common source of nitrogen in food wastes, researchers have targeted glutamic acid contained in sewage sludge and meat and bone meal, and they succeeded in producing ammonia with 35–51% efficiency at 800 °C and 0.5–1.0 g g<sup>-1</sup> carbon content by automated gasification [56]. This reaction is dependent on the catalytic metal ions in the food waste, which may need to be optimized for each feedstock. The addition of LaFeO<sub>3</sub> as a catalyst resulted in an efficiency of 54 vol% [57]. Therefore, this approach could potentially produce 10% of the ammonia used in Europe [56].

Several studies have developed sustainable methods to produce ammonia, most of which include green power generation methods, chemical synthesis methods, and sustainable ammonia production using biological methods [59–61]. Various methods of recovering ammonia from food wastes have also been used, and research is progressing toward practical application [62]. Alternatively, we focus on sustainable ammonia production methods such as biological methods—bacterial and yeast methods—and explore their application potential (Figure 2).



## Ammonia Production Methods

Figure 2. Sustainable ammonia production (created with Biorender.com).

#### 2. Engineered Bacterial Method for Ammonia Production

Microorganisms can degrade and synthesize a wide variety of compounds by applying engineering methods such as directed evolution or genome editing [63–66]. Based on previous reports, pharmaceuticals that cannot be digested by the human body are broken down by microorganisms in the environment [67]. Considering their capability to degrade and synthesize not only natural substances but also man-made substances, microorganism have been used to synthesize a variety of substances (Figure 3) [68,69].

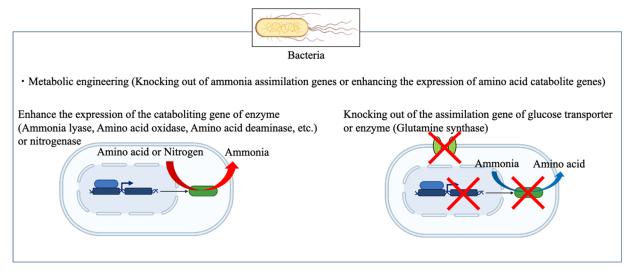


Figure 3. Ammonia production using engineered bacteria (created with Biorender.com).

#### 2.1. Bioengineering of Nitrogen Fixation and Metabolic Engineering for Ammonia Production

Compared with the Haber–Bosch process, the use of nitrogenase is highly sustainable because it can produce ammonia at room temperature and pressure [70]. The amount of nitrogen fixed by forage legumes accounts for  $17.2 \times 10^7$  tons of nitrogen per year [71–73]. Thus, approximately 21% of the nitrogen is produced by nitrogen fixation [74].

Two main species of microorganisms possess nitrogenases. One species is the genus *Rhizobia*, which parasitizes legumes, and the other species is the genera *Azotobacter* and *Klebsiella*, which can grow without parasitizing plants and are the main focus of research [75,76]. Nitrogenase utilizes considerable energy, requiring eight molecules of ATP to produce one molecule of ammonia, and it is easily deactivated by oxygen exposure [77]. Nitrogenase is composed of multiple subunits; the roles of each subunit are interrelated, and the molecular mechanism is complex. Thus, the reaction mechanism remains unclear [78,79].

Attempts to produce ammonia using nitrogenase have been pursued, starting with the construction of a heterologous expression system for the nitrogenase subunits from Klebsiella oxytoca in E. coli [80]. This nitrogenase uses molybdenum as a cofactor, but Azotobacter vinelandii exists as a nitrogen-fixing bacterium with a nitrogenase that uses vanadium and iron. Various attempts have already been made with some success to construct heterologous expression of this nitrogenase gene cluster in *E. coli* and yeast, to elucidate the mechanism by which it is protected from oxygen, and to increase its activity [81–83]. For example, heterologous expression of the nitrogenase complex of A. vinelandii has been successful in E. *coli*, and the minimum number of genes required for nitrogen fixation has been elucidated. A. vinelandii possesses an Fe nitrogenase complex, which consists of 21 subunits, and this nitrogenase complex is actively expressed in only 10 genes in *E. coli* [84]. Furthermore, Paenibacillus sp. is composed of 20 subunits of the Fe-type nitrogenase and functions with a set of nine genes. However, the activity of the heterologously expressed nitrogenase was approximately 10% compared to the nitrogenase of the wild type [85]. Further studies have shown that adding Klebsiella oxytoca NifSU (Fe-S cluster assembly) and Paenibacillus electron transporter genes (*pfoABfldA*) to this minimal gene set improved the expression, yielding 50.1% of the activity of the wild type [85]. A further attempt was made to reconstitute them

by combining fourteen nitrogenase-related genes from *K. oxytoca* into five gene cassettes and cleaving the subunits with a protease from the tobacco etch virus. As a result, this *E. coli* could grow only nitrogen molecules in the air without oxygen [84].

Various theories have been proposed to investigate the mechanism through which nitrogen-fixing bacteria fix nitrogen without inactivating nitrogenase under aerobic conditions; *Pseudomonas stutzeri* A1501 forms a cyst made of a polysaccharide membrane to protect oxygen, thereby protecting nitrogenase from oxygen [86]. *A. vinelandii* also produces polysaccharides to protect nitrogenase from oxygen by localizing the polysaccharides to the plasma membrane [83].

Apart from the aforementioned methods that can directly improve ammonia production capacity, attempts have been made to increase the ammonia supplied to plants by enhancing the ability of nitrogen-fixing bacteria to bind to plants, which has reached practical application [87]. For example, by ingesting *Pseudomonas protegens* Pf-5 X940, 15 nitrogen isotope dilution analyses showed that corn and wheat produced significant amounts of fixed nitrogen by this organism. In addition, the colony in wheat plants was formed on the root surface of Pf-5 X940 expressing GFP [88].

Recently, for industrial applications, various companies such as Ginkgo Bioworks and Pivot Bio, Inc., have been studying the potential use of nitrogen-fixing microbes. Pivot Bio, Inc., performed a study that used nitrogen-fixing bacterium of the genus *Enterobacter* sp. with genetic engineering [89]. These bacteria dis not have enough glutamine because of the low level of expression of the transcription factor GlnR. Thus, when increasing intracellular glutamine, these bacteria synthesized ammonia in the presence of more nitrogenase than the wild type [89]. In addition, corn (*Zea mays*) seeds with these bacteria were pre-infected and provided a steady nutrient supply [89]. In 2019, the average corn N uptake was higher when inoculated with Proven<sup>TM</sup> than that without nitrogen fertilizer application, with yields of 0 kg N ha<sup>-1</sup> (treated with fertilizer) and 10.9 kg N ha<sup>-1</sup> yields (added with bacteria), but this difference was not statistically significant [90].

Many other attempts have been made to use anaerobic microflora to produce ammonia [13,61,91]. A number of methodologies for ammonia recovery have also been proposed, including evaporating the solution after fermentation and increasing the pH [92,93].

#### 2.2. Ammonia Production from Food Waste by Using Bioengineering Methods

Several methods have been reported for producing ammonia from food waste and other biomass using engineered E. coli and Bacillus [54,94–97]. For example, Bacillus subtilis is used to hydrolyze protein biomass with its own protease and to efficiently produce amino acids, and ammonia and alcohol are highly produced from amino acids using metabolic engineering methods [94]. In particular, the authors knocked out the codY gene, a regulator of branched-chain amino acids in the B. subtilis metabolic pathway. Furthermore, the authors knocked out *bkdB*, a lipoamidoacyltransferase, which inhibits the conversion of 2-keto acids to acyl CoA. Then, 2-keto acids are decarboxylated by heterologously expressed 2-keto acid decarboxylase to provide biofuel accumulation. Furthermore, the overexpression of the alpha-ketoisovalerate decarboxylase gene *leuDH* accelerated ammonia production. Consequently, the biofuel production, including ammonia by deamination from media containing amino acid, produced 46.6% of a theoretical yield [94]. Ammonia production from amino acids using *E. coli* has also been attempted [96,97]. Considering the strong ammonia assimilation capacity to keep the cell in homeostasis (about 20% of nitrogen compounds is stored in the cell) of E. coli, attempts were made to knock out genes involved in ammonia assimilation. For example, knockout of the glutamine synthetase *glnA* was predicted to increase the production of ammonia. Furthermore, overexpression of the decarboxylase gene kivD was used to bias the equilibrium reaction between ammonia and amino acid production toward the ammonia-producing side. As a result, the production of ammonia from a medium containing amino acids succeeded with a theoretical yield of 47.8% [97]. However, the abovementioned studies were primarily experiments using amino acid-containing media and not actual food wastes. Therefore, researchers investigated ammonia production from

six different culture media and four different food wastes using *E. coli* to promote the use of actual food wastes [96]. In addition, metabolic profiling showed a correlation between the concentration of substances such as sugars, organic acids and amino acids in the medium, and ammonia production suggested that glucose inhibits the production of ammonia. The M9 yeast extract medium containing sugar at various concentrations showed a negative correlation with ammonia production. Therefore, researchers knocked out the glucose transporter gene *ptsG*, which transports major sugars such as glucose in *E. coli*, deduced the expression of phosphoenolpyruvate- and phosphotransferase-related genes, and succeeded in producing ammonia from the medium containing amino acid and sugar with about 73% yield. Furthermore, ammonia was successfully produced from pretreated soybean residue with a conversion efficiency of about 47% and a high ammonia concentration of approximately 35 mM [96]. In these studies, ammonia was produced in the cells, indicating a trade-off between microbial growth and ammonia production. Moreover, a study using metabolically modified *B. subtilis* for ammonia production reached a plateau at day 6, and it did not increase at day 7 [95]. These results indicated that the produced ammonia was used for cell growth. Therefore, extracellular ammonia production can simultaneously improve production and growth.

#### 3. Engineered Yeast Method for Ammonia Production

In addition to *E. coli*, many attempts have been made to produce ammonia using yeast [98,99]. Numerous attempts have also been made to investigate the heterologous expression of nitrogenases in yeast. For example, the nitrogenase subunit NifB is required for the formation of MoFe clusters at the nitrogenase active center; however, this protein is insoluble in mitochondria of budding yeast [100]. Two combinatorial libraries were constructed for optimization. One library consists of six subclusters including nifUSX and fdxN, and the other library includes 28 NifB genes extracted from the public database and showed different levels of expression based on a factor design [101]. Consequently, NifB genes derived from the archaea Methanocaldococcus infernus or Methanothermobacter *thermautotrophicus* were activated in the mitochondria of yeast [101]. Nitrogenase subunits derived from K. oxytoca within plant mitochondria were also used to attempt expression. In addition, NifF, NifM, NifN, NifS, NifU, NifW, NifX, NifY, and NifZ could be expressed in a soluble form, whereas NifB, NifE, NifH, NifJ, NifK, NifQ, and NifV are insoluble. However, the NifM activity decreased by 10% compared with the previous study because of N-terminal processing after transportation to the mitochondria [102,103]. Furthermore, a NifD protein derived from K. oxytoca contains a mitochondrial targeting peptide (MTP) recognition sequence that undergoes processing, making functional expression difficult in mitochondria, and R98 has been identified as an important residue of this cleavage. As a result, R98P shows to be resistant to processing, and high levels of activity are retained in the mitochondria [104]. Another study found that the Y100Q mutant is resistant to processing within plant and yeast mitochondria [105]. In their search for efficient NifH expression, 32 different NifHs were expressed in tobacco and yeast mitochondria and it was found that NifH from the thermophilic bacterium Hydrogenobacter thermophilus is the most active form [106]. Thus, the expression of nitrogenase subunits from thermophilic bacteria is a potential approach that is worthy of further study because the temperature in mitochondria could reach 50 °C [106]. Genetic diversity can be used to identify the most appropriate Nif protein components from large sequence pools to manipulate eukaryotic nitrogenases [107]. Cloning techniques such as codon optimization of gene synthesis and other synthetic biology tools such as metabolic engineering methods allow for the construction of multi-protein pathways containing the proteins' various sorts of origins. For example, the NifH protein obtained from *H. thermophilus* was found to be soluble in the mitochondria of Saccharomyces cerevisiae and Nicotiana benthamiana, which accumulate at higher levels than A. vinelandii [107]. Therefore, functional nitrogenases in plant cells could be constructed in the future to create practical crops that can fix nitrogen on their own [108]. Recently, researchers have also performed heterologous expression of an active

nitrogenase in the chloroplasts of the cyanobacterium *Synechococcus elongatus* PCC7942, creating an algae that can fix nitrogen using energy from photosynthesis [109].

Metabolically engineered yeast can produce large amounts of ammonia because its of exposure to high concentrations of ammonia, thereby causing growth defects [110]. Yeastbased methods for producing ammonia from food waste have primarily used cell surface engineering (Figure 4) [111]. The nitrogen source of food waste is primarily amino acids derived from proteins; thus, amino acid oxidases, which can efficiently produce ammonia from amino acids, are considered suitable for presentation to the cell surface and are efficient in producing ammonia. Several studies have used yeast cell surface display systems [111]. For example, target proteins are presented on the cell surface by adding a signal for secretion to the N-terminus and a cell wall anchor protein including a glycosylphosphatidylinositol anchor attachment signal sequence to the C-terminus. Approximately 10<sup>5</sup>–10<sup>6</sup> proteins could be presented on the yeast cell surface, and yeast cells can be manipulated as a whole-cell biocatalyst [111]. Furthermore, proteins immobilized on the cell surface can stabilize and exhibit higher enzymatic activity than those in the free state. In addition, the folding machinery of the eukaryote displays a variety of proteins [111]. Thus, given these advantages, the yeast cell surface has been successfully manipulated to produce ethanol from carbohydrates, woody biomass, and large algae with high efficiency [112,113]. Ammonia produced from soybean residues has been used in yeast cell surface engineering. Amino acid catabolic enzymes such as deaminase, transaminase, oxidase, and ammonia lyase are known to produce ammonia from amino acids. Ammonia lyases can be displayed on the yeast cell surface because they do not require any cofactors to make holoenzymes. In a study, glutamine ammonia lyase (YbaS) was displayed on the cell surface, and this yeast successfully produced ammonia from a glutamine solution with high efficiency (83.2%) and high concentration (3.34 g/L) [98]. Moreover, although 0.1% (v/v) ammonia-containing medium inhibited yeast growth, no impairment of the function of this whole-cell catalyst was observed [98,99]. Furthermore, YbaS-displaying yeast successfully produced ammonia with high efficiency from a solution of enzymatically pretreated okara (soybean residues). However, when producing ammonia using YbaS ammonia lyase, only glutamine was used to produce ammonia among the 20 amino acids [114]. L-amino acid oxidase is known to have broad substrate specificity, and it can produce ammonia from a wide variety of amino acids [99]. For example, a L-amino acid oxidase from Aplysia californica is active against L-arginine and L-lysine. In addition, a L-amino acid oxidase obtained from snakes such as Bothrops atrox, Crotalus viridis helleri, and Daboia russelii has been well studied, and it is inactive against amino acids such as L-glutamine and L-aspartic acid [115]. However, it is active against hydrophobic amino acids such as L-leucine and hydrophilic amino acids such as L-histidine and L-lysine. It is also inactive against aromatic amino acids such as L-tyrosine and L-phenylalanine. *Hebeloma cylindrosporum* also contains enzymes with broad specificity, as it has mycelia for the intracellular absorption of ammonia from amino acids contained in soil. In the literature, an enzyme, namely, HcLAAO (L-amino acid oxidase from *H. cylindrosporum*), can produce ammonia from more than 10 known amino acids [116]. Therefore, ammonia can be efficiently produced from food waste using yeast displaying HcLAAO. The display of HcLAAO in budding yeast led to high ammonia production efficiency (about 88%) under mild conditions (30  $^{\circ}$ C) from a pretreated okara solution [99]. In this study, no toxic effects of extracellular ammonia on catalytic activity were observed, which is consistent with previous studies that used glutamine ammonia lyase. Although these attempts to produce ammonia from food waste have been successful, these studies are all laboratory-scale studies. Enzymes have also been developed as tools to produce ammonia from amino acids, with the recent creation of enzymes that can produce ammonia from 13 amino acids [117]. Therefore, industrial-scale attempts for social implementation should be considered in the future.

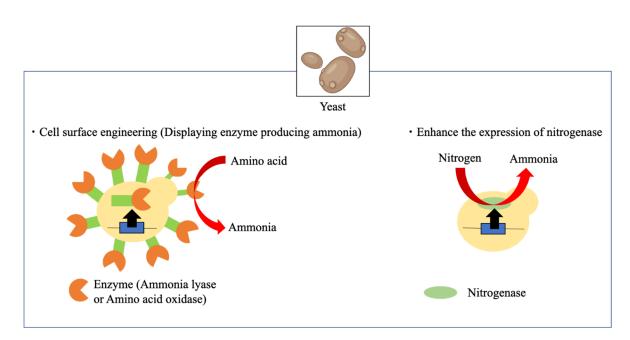


Figure 4. Ammonia production using engineered yeast (created with Biorender.com).

#### 4. Conclusions

This review mainly focused on ammonia production using genetic engineering methods (Table 1). Energy production with low environmental burden is considered essential for the formation of a sustainable society. In addition to studies in this review, for example, anaerobic culture is also one of the methods currently in practical use for energy production from food wastes [118]. These anaerobic cultures usually do not use genetic engineering methods, such that this method has room for optimization by using genetic engineering introduced in this review.

	Metabolic Engineering	Cell Surface Engineering	
Reaction Place	• Inside of the cells	• Outside of the cells	
pH of reaction environment	• Neutral condition (pH 7.2–7.8)	• Depends on the reaction mixture	
Effect on the enzyme	• Nothing	• Stable (fixed on the cell surface)	

**Table 1.** Summary of biological methods for ammonia production.

To achieve a sustainable society, we have to solve some problems other than the development of biological methods such as the storage of food waste [119,120]. In addition, consumption of ammonia is important for creating sustainable environment. For example, ammonia is only 30-50% utilized for crop growth [121]. The remainder eutrophicates the oceans, contaminates drinking water, and, in some areas, changes the composition of vegetation and encourages the propagation of non-native species [122–125]. Ammonia is converted to N<sub>2</sub>O by microorganisms in the environment, which is also a well-known greenhouse gas, and its emission is also a problem [74,126].

Microorganisms such as *E. coli* and yeast contain nitrogen in their bodies, with about 20% of the total amount as proteins [127]. The nitrogen in cells comes from assimilation of amino acids or proteins in a medium. Therefore, the amount of produced nitrogen must be larger than the input nitrogen for effective biofuel production. To solve this problem, cell surface engineered yeast could be repeatedly used for production because the enzyme on the yeast cell surface is stable [128]. Recently, some studies attempted to estimate the metabolism of microorganisms so that it might be possible to calculate the potential of microorganisms to efficiently produce ammonia in the future [129–131].

In this review, we introduced various attempts to improve the Haber–Bosch process, one of the sources of carbon dioxide generation, which uses the largest amount of fossil fuels, in order to realize a sustainable society.

The use of genetically optimized microorganisms to produce ammonia instead of the Haber–Bosch process is a potential and environmentally friendly approach to solve the global ammonia demand problem and to develop a sustainable carbon-free society (Table 2).

	<b>Chemical Method</b>	<b>Bacterial Method</b>	Yeast Method
Resource	<ul><li>Nitrogen in air</li><li>Hydrogen</li></ul>	<ul><li>Nitrogen in air</li><li>Food waste</li></ul>	Food waste
Energy resource	• Fossil fuel (coal, oil, or natural gas)	Not required	Not required
ReactionEnvironment	<ul> <li>High pressure</li> <li>(150–350 atm)</li> <li>High temperature</li> <li>(350–350 °C)</li> </ul>	<ul> <li>Pressure (1 atm)</li> <li>Temperature (37 °C)</li> </ul>	<ul> <li>Pressure (1 atm)</li> <li>Temperature (37 °C)</li> </ul>
Point of improvement for sustainable production	<ul> <li>Alternative energy resource (solar, wind, hydro, etc.)</li> <li>Novel catalyst to react with ambient pressure and temperature</li> </ul>	<ul> <li>Heterologous expression of nitrogenase</li> <li>Metabolic engineering to improve ammonia production from amino acids contained in food wastes</li> </ul>	Cell surface

Table 2. Summary of ammonia production methods.

Therefore, in the future, it may be possible to produce ammonia more efficiently by using it concurrently with such methodologies.

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#### References

- Fawzy, S.; Osman, A.I.; Doran, W.J.; Rooney, D.W. Strategies for mitigation of climate change: A review. *Environ. Chem. Lett.* 2020, 18, 2069–2094. [CrossRef]
- Tortell, P.D. Earth 2020: Science, society, and sustainability in the Anthropocene. Proc. Natl. Acad. Sci. USA 2020, 117, 8683–8691. [CrossRef] [PubMed]
- 3. Capua, I.; Giovannini, E. Coding system to track research progress towards SDGs. Nature 2019, 572, 178. [CrossRef] [PubMed]
- 4. Chu, S.; Majumdar, A. Opportunities and challenges for a sustainable energy future. Nature 2012, 488, 294–303. [CrossRef]
- 5. Chu, S.; Cui, Y.; Liu, N. The path towards sustainable energy. Nat. Mater. 2016, 16, 16–22. [CrossRef]
- 6. Hussain, A.; Arif, S.M.; Aslam, M. Emerging renewable and sustainable energy technologies: State of the art. *Renew. Sustain. Energy Rev.* **2017**, *71*, 12–28. [CrossRef]
- Creutzig, F.; Ravindranath, N.H.; Berndes, G.; Bolwig, S.; Bright, R.; Cherubini, F.; Chum, H.; Corbera, E.; Delucchi, M.; Faaij, A.; et al. Bioenergy and climate change mitigation: An assessment. *Glob. Chang. Biol. Bioenergy* 2015, 7, 916–944. [CrossRef]
- Elishav, O.; Lis, B.M.; Miller, E.M.; Arent, D.J.; Valera-Medina, A.; Dana, A.G.; Shter, G.E.; Grader, G.S. Progress and Prospective of Nitrogen-Based Alternative Fuels. *Chem. Rev.* 2020, 120, 5352–5436. [CrossRef]
- 9. Baliban, R.C.; Elia, J.A.; Floudas, C.A. Biomass and Natural Gas to Liquid Transportation Fuels: Process Synthesis, Global Optimization, and Topology Analysis. *Ind. Eng. Chem. Res.* 2013, 52, 3381–3406. [CrossRef]

- 10. Lynd, L.R.; Larson, E.; Greene, N.; Laser, M.; Sheehan, J.; Dale, B.E.; McLaughlin, S.; Wang, M. The role of biomass in America's energy future: Framing the analysis. *Biofuels Bioprod. Biorefin.* **2009**, *3*, 113–123. [CrossRef]
- Siddiki, S.Y.A.; Mofijur, M.; Kumar, P.S.; Ahmed, S.F.; Inayat, A.; Kusumo, F.; Badruddin, I.A.; Yunus Khan, T.M.; Nghiem, L.D.; Ong, H.C.; et al. Microalgae biomass as a sustainable source for biofuel, biochemical and biobased value-added products: An integrated biorefinery concept. *Fuel* 2022, 307, 121782. [CrossRef]
- Govarthanan, M.; Manikandan, S.; Subbaiya, R.; Krishnan, R.Y.; Srinivasan, S.; Karmegam, N.; Kim, W. Emerging trends and nanotechnology advances for sustainable biogas production from lignocellulosic waste biomass: A critical review. *Fuel* 2022, *312*, 122928. [CrossRef]
- Paritosh, K.; Kushwaha, S.K.; Yadav, M.; Pareek, N.; Chawade, A.; Vivekanand, V. Food Waste to Energy: An Overview of Sustainable Approaches for Food Waste Management and Nutrient Recycling. *BioMed Res. Int.* 2017, 2017, 2370927. [CrossRef]
- 14. Rosenboom, J.-G.; Langer, R.; Traverso, G. Bioplastics for a circular economy. Nat. Rev. Mater. 2022, 7, 117–137. [CrossRef]
- 15. Erisman, J.W.; Sutton, M.A.; Galloway, J.; Klimont, Z.; Winiwarter, W. How a century of ammonia synthesis changed the world. *Nat. Geosci.* **2008**, *1*, 636–639. [CrossRef]
- 16. Smil, V. Population Growth and Nitrogen: An Exploration of a Critical Existential Link. Popul. Dev. Rev. 1991, 17, 569. [CrossRef]
- Fedoruk, M.J.; Bronstein, R.; Kerger, B.D. Ammonia exposure and hazard assessment for selected household cleaning product uses. J. Expo. Sci. Environ. Epidemiol. 2005, 15, 534–544. [CrossRef]
- 18. Park, S.; Jeong, J.; Fujita, K.-I.; Yamamoto, A.; Yoshida, H. Anti-Markovnikov Hydroamination of Alkenes with Aqueous Ammonia by Metal-Loaded Titanium Oxide Photocatalyst. *J. Am. Chem. Soc.* **2020**, *142*, 12708–12714. [CrossRef]
- 19. Zumdahl, S.S. Ammonia. Encyclopedia Britannica. 2022. Available online: https://www.britannica.com/science/ammonia (accessed on 1 January 2022).
- 20. Hu, B.; Huang, S.; Shao, Y.; Chen, J. Thermodynamic analysis of a new ammonia-water power cycle. *Energy Rep.* **2020**, *6*, 567–573. [CrossRef]
- Wakida, T.; Tokuyama, T.; Doi, C.; Lee, M.; Jeong, D.S.; Ishida, S. Mechanical Properties of Polyester/Cotton and Polyester/Rayon Fabrics Treated with Ammonia-Gas. J. Soc. Fiber Sci. Technol. 2004, 60, 34–37. [CrossRef]
- 22. Müller, T.E.; Beller, M. Metal-Initiated Amination of Alkenes and Alkynes. Chem. Rev. 1998, 98, 675–704. [CrossRef] [PubMed]
- 23. Kishimoto, M.; Muroyama, H.; Suzuki, S.; Saito, M.; Koide, T.; Takahashi, Y.; Horiuchi, T.; Yamasaki, H.; Matsumoto, S.; Kuboet, H.; et al. Development of 1 kW-class ammonia-fueled solid oxide fuel cell stack. *Fuel Cells* **2020**, *20*, 80–88. [CrossRef]
- 24. Zhao, Y.; Setzler, B.P.; Wang, J.; Nash, J.; Wang, T.; Xu, B.; Yan, Y. An Efficient Direct Ammonia Fuel Cell for Affordable Carbon-Neutral Transportation. *Joule* 2019, *3*, 2472–2484. [CrossRef]
- Valera-Medina, A.; Amer-Hatem, F.; Azad, A.K.; Dedoussi, I.C.; de Joannon, M.; Fernandes, R.X.; Glarborg, P.; Hashemi, H.; He, X.; Mashruk, S.; et al. Review on Ammonia as a Potential Fuel: From Synthesis to Economics. *Energy Fuels* 2021, 35, 6964–7029. [CrossRef]
- 26. Erdemir, D.; Dincer, I. A perspective on the use of ammonia as a clean fuel: Challenges and solutions. *Int. J. Energy Res.* **2021**, 45, 4827–4834. [CrossRef]
- 27. Herbinet, O.; Bartocci, P.; Dana, A.G. On the use of ammonia as a fuel—A perspective. Fuel Commun. 2022, 11, 100064. [CrossRef]
- 28. Miura, D.; Tezuka, T. A comparative study of ammonia energy systems as a future energy carrier, with particular reference to vehicle use in Japan. *Energy* **2014**, *68*, 428–436. [CrossRef]
- Valera-Medina, A.; Xiao, H.; Owen-Jones, M.; David, W.I.F.; Bowen, P.J. Ammonia for power. Prog. Energy Combust. Sci. 2018, 69, 63–102. [CrossRef]
- Wijayanta, A.T.; Oda, T.; Purnomo, C.W.; Kashiwagi, T.; Aziz, M. Liquid hydrogen, methylcyclohexane, and ammonia as potential hydrogen storage: Comparison review. *Int. J. Hydrogen Energy* 2019, 44, 15026–15044. [CrossRef]
- 31. Moseley, P.T.; Garche, J. *Electrochemical Energy Storage for Renewable Sources and Grid Balancing*; Newnes: Oxford, UK; Elsevier: Amsterdam, The Netherlands, 2014.
- Lan, R.; Irvine, J.T.S.; Tao, S. Ammonia and related chemicals as potential indirect hydrogen storage materials. *Int. J. Hydrog. Energy* 2012, 37, 1482–1494. [CrossRef]
- Jeerh, G.; Zhang, M.; Tao, S. Recent progress in ammonia fuel cells and their potential applications. J. Mater. Chem. A Mater. Energy Sustain. 2021, 9, 727–752. [CrossRef]
- 34. Wang, W.; Herreros, J.M.; Tsolakis, A.; York, A.P. Ammonia as hydrogen carrier for transportation; investigation of the ammonia exhaust gas fuel reforming. *Int. J. Hydrogen Energy* **2013**, *38*, 9907–9917. [CrossRef]
- 35. Palys, M.J.; Daoutidis, P. Using hydrogen and ammonia for renewable energy storage: A geographically comprehensive techno-economic study. *Comput. Chem. Eng.* 2020, 136, 106785. [CrossRef]
- Al-Aboosi, F.Y.; El-Halwagi, M.M.; Moore, M.; Nielsen, R.B. Renewable ammonia as an alternative fuel for the shipping industry. *Curr. Opin. Chem. Eng.* 2021, 31, 100670. [CrossRef]
- 37. Green, L. An ammonia energy vector for the hydrogen economy. Int. J. Hydrogen Energy 1982, 7, 355–359. [CrossRef]
- 38. Schrock, R.R. Reduction of dinitrogen. Proc. Natl. Acad. Sci. USA 2006, 103, 17087. [CrossRef]
- 39. Shilov, A.E. Catalytic reduction of molecular nitrogen in solutions. Russ. Chem. Bull. 2003, 52, 2555–2562. [CrossRef]
- 40. Capdevila-Cortada, M. Electrifying the Haber–Bosch. *Nat. Catal.* **2019**, *2*, 1055. [CrossRef]
- 41. Raróg-Pilecka, W.; Miśkiewicz, E.; Szmigiel, D.; Kowalczyk, Z. Structure sensitivity of ammonia synthesis over promoted ruthenium catalysts supported on graphitised carbon. *J. Catal.* **2005**, 231, 11–19. [CrossRef]

- 42. Over, H. Surface Chemistry of Ruthenium Dioxide in Heterogeneous Catalysis and Electrocatalysis: From Fundamental to Applied Research. *Chem. Rev.* 2012, 112, 3356–3426. [CrossRef]
- Kitano, M.; Inoue, Y.; Yamazaki, Y.; Hayashi, F.; Kanbara, S.; Matsuishi, S.; Yokoyama, T.; Kim, S.W.; Hara, M.; Hosono, H. Ammonia synthesis using a stable electride as an electron donor and reversible hydrogen store. *Nat. Chem.* 2012, *4*, 934–940. [CrossRef] [PubMed]
- 44. Humphreys, J.; Lan, R.; Tao, S. Development and Recent Progress on Ammonia Synthesis Catalysts for Haber–Bosch Process. *Adv. Energy Sustain. Res.* 2020, *2*, 2000043. [CrossRef]
- Ashida, Y.; Arashiba, K.; Tanaka, H.; Egi, A.; Nakajima, K.; Yoshizawa, K.; Nishibayashi, Y. Molybdenum-Catalyzed Ammonia Formation Using Simple Monodentate and Bidentate Phosphines as Auxiliary Ligands. *Inorg. Chem.* 2019, 58, 8927–8932. [CrossRef] [PubMed]
- 46. Garrido-Barros, P.; Derosa, J.; Chalkley, M.J.; Peters, J.C. Tandem electrocatalytic N2 fixation via proton-coupled electron transfer. *Nature* 2022, 609, 71–76. [CrossRef] [PubMed]
- 47. Ghavam, S.; Vahdati, M.; Wilson, I.A.G.; Styring, P. Sustainable Ammonia Production Processes. *Front. Energy Res.* **2021**, *9*, 580808. [CrossRef]
- Wang, M.; Khan, M.A.; Mohsin, I.; Wicks, J.; Ip, A.H.; Sumon, K.Z.; Dinh, C.-T.; Sargent, E.H.; Gates, I.D.; Kibria, G.; et al. Can sustainable ammonia synthesis pathways compete with fossil-fuel based Haber–Bosch processes? *Energy Environ. Sci.* 2021, 14, 2535–2548. [CrossRef]
- 49. Palys, M.J.; Wang, H.; Zhang, Q.; Daoutidis, P. Renewable ammonia for sustainable energy and agriculture: Vision and systems engineering opportunities. *Curr. Opin. Chem. Eng.* **2021**, *31*, 100667. [CrossRef]
- 50. Salmon, N.; Bañares-Alcántara, R. Green ammonia as a spatial energy vector: A review. *Sustain. Energy Fuels* **2021**, *5*, 2814–2839. [CrossRef]
- 51. Melikoglu, M.; Lin, C.S.K.; Webb, C. Analysing global food waste problem: Pinpointing the facts and estimating the energy content. *Cent. Eur. J. Eng.* **2013**, *3*, 157–164. [CrossRef]
- 52. Liu, C.; Hotta, Y.; Santo, A.; Hengesbaugh, M.; Watabe, A.; Totoki, Y.; Allen, D.; Bengtsson, M. Food waste in Japan: Trends, current practices and key challenges. *J. Clean. Prod.* 2016, 133, 557–564. [CrossRef]
- 53. Katami, T.; Yasuhara, A.; Shibamoto, T. Formation of Dioxins from Incineration of Foods Found in Domestic Garbage. *Environ. Sci. Technol.* **2004**, *38*, 1062–1065. [CrossRef]
- 54. Chan, L.Y.; Takahashi, M.; Lim, P.J.; Aoyama, S.; Makino, S.; Ferdinandus, F.; Ng, S.Y.C.; Arai, S.; Fujita, H.; Tan, H.C.; et al. Eurotium Cristatum Fermented Okara as a Potential Food Ingredient to Combat Diabetes. *Sci. Rep.* **2019**, *9*, 17536. [CrossRef]
- 55. Hadj Saadoun, J.; Calani, L.; Cirlini, M.; Bernini, V.; Neviani, E.; Del Rio, D.; Galavernaa, G.; Lazzi, C. Effect of fermentation with single and co-culture of lactic acid bacteria on okara: Evaluation of bioactive compounds and volatile profiles. *Food Funct.* **2021**, *12*, 3033–3043. [CrossRef]
- Gil-Lalaguna, N.; Afailal, Z.; Aznar, M.; Fonts, I. Exploring the sustainable production of ammonia by recycling N and H in biological residues: Evolution of fuel-N during glutamic acid gasification. J. Clean. Prod. 2021, 282, 124417. [CrossRef]
- Wang, P.; Xu, P.; Wang, B.; Shen, C.; Shen, L. Green ammonia production via microalgae steam catalytic gasification process over LaFeO<sub>3</sub> perovskite. *Fuel* 2022, 318, 123322. [CrossRef]
- Wang, P.; Shen, C.; Wang, B.; Xu, P.; Shen, L. Ammonia production from nitrogen-rich biomass gasification: Nitrogen transformation from model amino acids. *Fuel* 2022, 326, 125071. [CrossRef]
- Suryanto, B.H.R.; Du, H.-L.; Wang, D.; Chen, J.; Simonov, A.N.; MacFarlane, D.R. Challenges and prospects in the catalysis of electroreduction of nitrogen to ammonia. *Nat. Catal.* 2019, 2, 290–296. [CrossRef]
- 60. Wang, L.; Xia, M.; Wang, H.; Huang, K.; Qian, C.; Maravelias, C.T.; Ozin, G.A. Greening Ammonia toward the Solar Ammonia Refinery. *Joule* 2018, 2, 1055–1074. [CrossRef]
- 61. Zamri, M.; Hasmady, S.; Akhiar, A.; Ideris, F.; Shamsuddin, A.; Mofijur, M.; Fattah, I.M.R.; Mahlia, T. A comprehensive review on anaerobic digestion of organic fraction of municipal solid waste. *Renew. Sustain. Energy Rev.* **2021**, *137*, 110637. [CrossRef]
- 62. Lin, L.; Yuan, S.; Chen, J.; Xu, Z.; Lu, X. Removal of ammonia nitrogen in wastewater by microwave radiation. *J. Hazard. Mater.* **2009**, *161*, 1063–1068. [CrossRef]
- 63. Yang, D.; Park, S.Y.; Park, Y.S.; Eun, H.; Lee, S.Y. Metabolic Engineering of Escherichia coli for Natural Product Biosynthesis. *Trends Biotechnol.* **2020**, *38*, 745–765. [CrossRef]
- 64. Ko, Y.-S.; Kim, J.W.; Lee, J.A.; Han, T.; Kim, G.B.; Park, J.E.; Lee, S.Y. Tools and strategies of systems metabolic engineering for the development of microbial cell factories for chemical production. *Chem. Soc. Rev.* **2020**, *49*, 4615–4636. [CrossRef]
- 65. Zhang, R.; Li, C.; Wang, J.; Yang, Y.; Yan, Y. Microbial production of small medicinal molecules and biologics: From nature to synthetic pathways. *Biotechnol. Adv.* 2018, *36*, 2219–2231. [CrossRef]
- 66. Montaño López, J.; Duran, L.; Avalos, J.L. Physiological limitations and opportunities in microbial metabolic engineering. *Nat. Rev. Microbiol.* **2022**, *20*, 35–48. [CrossRef]
- 67. Costa, F.; Lago, A.; Rocha, V.; Barros, Ó.; Costa, L.; Vipotnik, Z.; Silva, B.; Tavares, T. A Review on Biological Processes for Pharmaceuticals Wastes Abatement—A Growing Threat to Modern Society. *Environ. Sci. Technol.* **2019**, *53*, 7185–7202. [CrossRef]
- Lopes, M.S.G. Engineering biological systems toward a sustainable bioeconomy. J. Ind. Microbiol. Biotechnol. 2015, 42, 813–838.
   [CrossRef]

- 69. Bornscheuer, U.T.; Huisman, G.W.; Kazlauskas, R.J.; Lutz, S.; Moore, J.C.; Robins, K. Engineering the third wave of biocatalysis. *Nature* 2012, 485, 185–194. [CrossRef]
- Rapson, T.D.; Gregg, C.M.; Allen, R.S.; Ju, H.; Doherty, C.M.; Mulet, X.; Giddey, S.; Wood, C.C. Insights into Nitrogenase Bioelectrocatalysis for Green Ammonia Production. *Chemsuschem* 2020, 13, 4856–4865. [CrossRef]
- 71. Gruber, N.; Galloway, J.N. An Earth-system perspective of the global nitrogen cycle. Nature 2008, 451, 293–296. [CrossRef]
- 72. Field, C.B.; Behrenfeld, M.J.; Randerson, J.T.; Falkowski, P. Primary Production of the Biosphere: Integrating Terrestrial and Oceanic Components. *Science* **1998**, *281*, 237–240. [CrossRef]
- 73. Ishizuka, J. Trends in biological nitrogen fixation research and application. In *Biological Nitrogen Fixation for Sustainable Agriculture: Extended Versions of Papers Presented in the Symposium, Role of Biological Nitrogen Fixation in Sustainable Agriculture at the 13th Congress of Soil Science, Kyoto, Japan, 1990*; Ladha, J.K., George, T., Bohlool, B.B., Eds.; Springer: Dordrecht, The Netherlands, 1992; pp. 197–209.
- Canfield, D.E.; Glazer, A.N.; Falkowski, P.G. The Evolution and Future of Earth's Nitrogen Cycle. *Science* 2010, 330, 192–196. [CrossRef] [PubMed]
- Lindström, K.; Mousavi, S.A. Effectiveness of nitrogen fixation in rhizobia. *Microb. Biotechnol.* 2020, 13, 1314–1335. [CrossRef] [PubMed]
- 76. Philippot, L.; Raaijmakers, J.M.; Lemanceau, P.; van der Putten, W.H. Going back to the roots: The microbial ecology of the rhizosphere. *Nat. Rev. Microbiol.* **2013**, *11*, 789–799. [CrossRef] [PubMed]
- Milton, R.D.; Minteer, S.D. Nitrogenase Bioelectrochemistry for Synthesis Applications. Acc. Chem. Res. 2019, 52, 3351–3360. [CrossRef] [PubMed]
- 78. Einsle, O.; Rees, D.C. Structural Enzymology of Nitrogenase Enzymes. Chem. Rev. 2020, 120, 4969–5004. [CrossRef]
- Cai, R.; Minteer, S.D. Nitrogenase Bioelectrocatalysis: From Understanding Electron-Transfer Mechanisms to Energy Applications. ACS Energy Lett. 2018, 3, 2736–2742. [CrossRef]
- Temme, K.; Zhao, D.; Voigt, C.A. Refactoring the nitrogen fixation gene cluster from *Klebsiella oxytoca*. Proc. Natl. Acad. Sci. USA 2012, 109, 7085–7090. [CrossRef]
- Yang, J.; Xie, X.; Wang, X.; Dixon, R.; Wang, Y.P. Reconstruction and minimal gene requirements for the alternative iron-only nitrogenase in *Escherichia coli. Proc. Natl. Acad. Sci. USA* 2014, 111, E3718–E3725. [CrossRef]
- López-Torrejón, G.; Burén, S.; Veldhuizen, M.; Rubio, L.M. Biosynthesis of cofactor-activatable iron-only nitrogenase in Saccharomyces cerevisiae. *Microb. Biotechnol.* 2021, 14, 1073–1083. [CrossRef]
- Takimoto, R.; Tatemichi, Y.; Aoki, W.; Kosaka, Y.; Minakuchi, H.; Ueda, M.; Kuroda, K. A critical role of an oxygen-responsive gene for aerobic nitrogenase activity in Azotobacter vinelandii and its application to Escherichia coli. *Sci. Rep.* 2022, *12*, 4182. [CrossRef]
- Yang, J.; Xie, X.; Xiang, N.; Tian, Z.-X.; Dixon, R.; Wang, Y.-P. Polyprotein strategy for stoichiometric assembly of nitrogen fixation components for synthetic biology. *Proc. Natl. Acad. Sci. USA* 2018, 115, E8509–E8517. [CrossRef]
- Li, X.-X.; Liu, Q.; Liu, X.-M.; Shi, H.-W.; Chen, S.-F. Using synthetic biology to increase nitrogenase activity. *Microb. Cell Fact.* 2016, 15, 43. [CrossRef]
- Wang, D.; Xu, A.; Elmerich, C.; Ma, L.Z. Biofilm formation enables free-living nitrogen-fixing rhizobacteria to fix nitrogen under aerobic conditions. *ISME J.* 2017, 11, 1602–1613. [CrossRef]
- 87. Klimasmith, I.M.; Kent, A.D. Micromanaging the nitrogen cycle in agroecosystems. *Trends Microbiol.* **2022**, *30*, 1045–1055. [CrossRef]
- Fox, A.R.; Soto, G.; Valverde, C.; Russo, D.; Lagares, A.; Zorreguieta, Á.; Alleva, K.; Pascuan, C.; Frare, R.; Mercado-Blanco, J.; et al. Major cereal crops benefit from biological nitrogen fixation when inoculated with the nitrogen-fixing bacterium *Pseudomonas* protegens Pf-5 X940. Environ. Microbiol. 2016, 18, 3522–3534. [CrossRef]
- 89. Bhatti, M.; Feng, P.C.C.; Pitkin, J. Methods and Compositions for Improving Plant Health. U.S. Patent 8754011, 17 June 2014.
- 90. Davis Pires, D. Pivot Bio Proven Inoculant as a Source of Nitrogen in Corn. Kans. Agric. Exp. Stn. Res. Rep. 2020, 6, 7. [CrossRef]
- 91. Yenigün, O.; Demirel, B. Ammonia inhibition in anaerobic digestion: A review. Process Biochem. 2013, 48, 901–911. [CrossRef]
- 92. Whelan, M.; Everitt, T.; Villa, R. A mass transfer model of ammonia volatilisation from anaerobic digestate. *Waste Manag.* 2010, 30, 1808–1812. [CrossRef]
- 93. Walker, M.; Iyer, K.; Heaven, S.; Banks, C. Ammonia removal in anaerobic digestion by biogas stripping: An evaluation of process alternatives using a first order rate model based on experimental findings. *Chem. Eng. J.* **2011**, *178*, 138–145. [CrossRef]
- 94. Choi, K.-Y.; Wernick, D.G.; Tat, C.A.; Liao, J.C. Consolidated conversion of protein waste into biofuels and ammonia using Bacillus subtilis. *Metab. Eng.* 2014, 23, 53–61. [CrossRef]
- Huo, Y.-X.; Cho, K.M.; Rivera, J.G.L.; Monte, E.; Shen, C.R.; Yan, Y.; Liao, J. Conversion of proteins into biofuels by engineering nitrogen flux. *Nat. Biotechnol.* 2011, 29, 346–351. [CrossRef] [PubMed]
- 96. Tatemichi, Y.; Kuroda, K.; Nakahara, T.; Ueda, M. Efficient ammonia production from food by-products by engineered Escherichia coli. *AMB Express* **2020**, *10*, 150. [CrossRef] [PubMed]
- Mikami, Y.; Yoneda, H.; Tatsukami, Y.; Aoki, W.; Ueda, M. Ammonia production from amino acid-based biomass-like sources by engineered Escherichia coli. AMB Express 2017, 7, 83. [CrossRef] [PubMed]
- 98. Watanabe, Y.; Kuroda, K.; Tatemichi, Y.; Nakahara, T.; Aoki, W.; Ueda, M. Construction of engineered yeast producing ammonia from glutamine and soybean residues (okara). *AMB Express* **2020**, *10*, 70. [CrossRef]

- 99. Watanabe, Y.; Aoki, W.; Ueda, M. Improved ammonia production from soybean residues by cell surface-displayed l-amino acid oxidase on yeast. *Biosci. Biotechnol. Biochem.* 2021, 85, 972–980. [CrossRef] [PubMed]
- Burén, S.; Jiang, X.; López-Torrejón, G.; Echavarri-Erasun, C.; Rubio, L.M. Purification and In Vitro Activity of Mitochondria Targeted Nitrogenase Cofactor Maturase NifB. Front. Plant Sci. 2017, 8, 1567. [CrossRef] [PubMed]
- 101. Burén, S.; Pratt, K.; Jiang, X.; Guo, Y.; Jimenez-Vicente, E.; Echavarri-Erasun, C.; Dean, D.; Saaem, I.; Gordon, B.; Voigtet, C.; et al. Biosynthesis of the nitrogenase active-site cofactor precursor NifB-co in Saccharomyces cerevisiae. *Proc. Natl. Acad. Sci. USA* 2019, 116, 25078–25086. [CrossRef]
- 102. Okada, S.; Gregg, C.M.; Allen, R.S.; Menon, A.; Hussain, D.; Gillespie, V.; Johnston, E.; Byrne, K.; Colgrave, M.L.; Wood, C.C. A Synthetic Biology Workflow Reveals Variation in Processing and Solubility of Nitrogenase Proteins Targeted to Plant Mitochondria, and Differing Tolerance of Targeting Sequences in a Bacterial Nitrogenase Assay. *Front. Plant Sci.* 2020, *11*, 552160. [CrossRef]
- 103. Smanski, M.J.; Bhatia, S.; Zhao, D.; Park, Y.; Woodruff, L.B.A.; Giannoukos, G.; Ciulla, D.; Busby, M.; Calderon, J.; Nicol, R.; et al. Functional optimization of gene clusters by combinatorial design and assembly. *Nat. Biotechnol.* 2014, 32, 1241–1249. [CrossRef]
- 104. Xiang, N.; Guo, C.; Liu, J.; Xu, H.; Dixon, R.; Yang, J.; Wang, Y.-P. Using synthetic biology to overcome barriers to stable expression of nitrogenase in eukaryotic organelles. *Proc. Natl. Acad. Sci. USA* **2020**, *117*, 16537–16545. [CrossRef]
- 105. Allen, R.S.; Gregg, C.M.; Okada, S.; Menon, A.; Hussain, D.; Gillespie, V.; Johnston, E.; Devilla, R.; Warden, A.C.; Taylor, M.; et al. Plant expression of NifD protein variants resistant to mitochondrial degradation. *Proc. Natl. Acad. Sci. USA* 2020, 117, 23165–23173. [CrossRef]
- 106. Chrétien, D.; Bénit, P.; Ha, H.-H.; Keipert, S.; El-Khoury, R.; Chang, Y.-T.; Jastroch, M.; Jacobs, H.T.; Rustin, P.; Rak, M. Mitochondria are physiologically maintained at close to 50 °C. *PLoS Biol.* 2018, *16*, e2003992. [CrossRef]
- 107. Jiang, X.; Payá-Tormo, L.; Coroian, D.; García-Rubio, I.; Castellanos-Rueda, R.; Eseverri, Á.; López-Torrejón, G.; Burén, S.; Rubio, L.M. Exploiting genetic diversity and gene synthesis to identify superior nitrogenase NifH protein variants to engineer N2-fixation in plants. *Commun. Biol.* 2021, 4, 4. [CrossRef]
- 108. Li, Q.; Chen, S. Transfer of nitrogen fixation (nif) genes to non-diazotrophic hosts. ChemBioChem 2020, 21, 1717–1722. [CrossRef]
- 109. Dong, F.; Lee, Y.S.; Gaffney, E.M.; Grattieri, M.; Haddadin, H.; Minteer, S.D.; Chen, H. An engineered, non-diazotrophic cyanobacterium and its application in bioelectrochemical nitrogen fixation. *Cell Rep. Phys. Sci.* **2021**, *2*, 100444. [CrossRef]
- Santos, J.; Sousa, M.J.; Leão, C. Ammonium Is Toxic for Aging Yeast Cells, Inducing Death and Shortening of the Chronological Lifespan. PLoS ONE 2012, 7, e37090. [CrossRef]
- 111. Ueda, M.; Tanaka, A. Cell surface engineering of yeast: Construction of arming yeast with biocatalyst. J. Biosci. Bioeng. 2000, 90, 125–136. [CrossRef]
- Takagi, T.; Yokoi, T.; Shibata, T.; Morisaka, H.; Kuroda, K.; Ueda, M. Engineered yeast whole-cell biocatalyst for direct degradation of alginate from macroalgae and production of non-commercialized useful monosaccharide from alginate. *Appl. Microbiol. Biotechnol.* 2016, 100, 1723–1732. [CrossRef]
- 113. Motone, K.; Takagi, T.; Sasaki, Y.; Kuroda, K.; Ueda, M. Direct ethanol fermentation of the algal storage polysaccharide laminarin with an optimized combination of engineered yeasts. *J. Biotechnol.* **2016**, 231, 129–135. [CrossRef]
- Brown, G.; Singer, A.; Proudfoot, M.; Skarina, T.; Kim, Y.; Chang, C.; Dementieva, I.; Kuznetsova, E.; Gonzalez, C.F.; Joachimiak, A.; et al. Functional and Structural Characterization of Four Glutaminases from Escherichia coli and Bacillus subtilis. *Biochemistry* 2008, 47, 5724–5735. [CrossRef]
- 115. Pollegioni, L.; Motta, P.; Molla, G. L-amino acid oxidase as biocatalyst: A dream too far? *Appl. Microbiol. Biotechnol.* **2013**, *97*, 9323–9341. [CrossRef] [PubMed]
- Bloess, S.; Beuel, T.; Krüger, T.; Sewald, N.; Dierks, T.; von Mollard, G.F. Expression, characterization, and site-specific covalent immobilization of an L-amino acid oxidase from the fungus Hebeloma cylindrosporum. *Appl. Microbiol. Biotechnol.* 2019, 103, 2229–2241. [CrossRef] [PubMed]
- 117. Nakano, S.; Kozuka, K.; Minamino, Y.; Karasuda, H.; Hasebe, F.; Ito, S. Ancestral L-amino acid oxidases for deracemization and stereoinversion of amino acids. *Commun. Chem.* **2020**, *3*, 181. [CrossRef]
- 118. Kastner, V.; Somitsch, W.; Schnitzhofer, W. The anaerobic fermentation of food waste: A comparison of two bioreactor systems. *J. Clean. Prod.* **2012**, *34*, 82–90. [CrossRef]
- 119. Degueurce, A.; Picard, S.; Peu, P.; Trémier, A. Storage of Food Waste: Variations of Physical–Chemical Characteristics and Consequences on Biomethane Potential. *Waste Biomass-Valoriz.* 2020, *11*, 2441–2454. [CrossRef]
- Daly, S.E.; Usack, J.G.; Harroff, L.A.; Booth, J.G.; Keleman, M.P.; Angenent, L.T. A systematic analysis of factors that affect food-waste storage: Toward maximizing lactate accumulation for resource recovery. ACS Sustain. Chem. Eng. 2020, 8, 13934–13944. [CrossRef]
- 121. Dawson, J.C.; Huggins, D.R.; Jones, S.S. Characterizing nitrogen use efficiency in natural and agricultural ecosystems to improve the performance of cereal crops in low-input and organic agricultural systems. *Field Crop. Res.* **2008**, *107*, 89–101. [CrossRef]
- Congreves, K.A.; Otchere, O.; Ferland, D.; Farzadfar, S.; Williams, S.; Arcand, M.M. Nitrogen Use Efficiency Definitions of Today and Tomorrow. *Front. Plant Sci.* 2021, 12, 637108. [CrossRef]
- 123. Galloway, J.N.; Winiwarter, W.; Leip, A.; Leach, A.; Bleeker, A.; Erisman, J.W. Nitrogen footprints: Past, present and future. *Environ. Res. Lett.* **2014**, *9*, 115003. [CrossRef]

- 124. Silliman, B.R.; Bertness, M.D. Shoreline Development Drives Invasion of Phragmites australis and the Loss of Plant Diversity on New England Salt Marshes. *Conserv. Biol.* 2004, *18*, 1424–1434. [CrossRef]
- 125. Hutchins, D.A.; Capone, D.G. The marine nitrogen cycle: New developments and global change. *Nat. Rev. Genet.* 2022, 20, 401–414. [CrossRef] [PubMed]
- 126. Montzka, S.A.; Dlugokencky, E.J.; Butler, J.H. Non-CO<sub>2</sub> greenhouse gases and climate change. *Nature* 2011, 476, 43–50. [CrossRef] [PubMed]
- 127. Schulz, E.; Oslage, H. Composition and nutritive value of single-cell protein (SCP). *Anim. Feed. Sci. Technol.* **1976**, *1*, 9–24. [CrossRef]
- 128. Yamakawa, S.-I.; Yamada, R.; Tanaka, T.; Ogino, C.; Kondo, A. Repeated fermentation from raw starch using Saccharomyces cerevisiae displaying both glucoamylase and α-amylase. *Enzym. Microb. Technol.* **2012**, *50*, 343–347. [CrossRef]
- 129. Zhou, Z.; Tran, P.Q.; Breister, A.M.; Liu, Y.; Kieft, K.; Cowley, E.S.; Karaoz, U.; Anantharaman, K. METABOLIC: High-throughput profiling of microbial genomes for functional traits, metabolism, biogeochemistry, and community-scale functional networks. *Microbiome* **2022**, *10*, 33. [CrossRef]
- 130. Kempes, C.P.; Dutkiewicz, S.; Follows, M.J. Growth, metabolic partitioning, and the size of microorganisms. *Proc. Natl. Acad. Sci.* USA **2012**, *109*, 495–500. [CrossRef]
- Calabrese, S.; Chakrawal, A.; Manzoni, S.; Van Cappellen, P. Energetic scaling in microbial growth. *Proc. Natl. Acad. Sci. USA* 2021, 118, e2107668118. [CrossRef]

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