




## Review

# Value-Added Products from Ethanol Fermentation—A Review

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**Abstract:** Global demand for renewable and sustainable energy is increasing, and one of the most common biofuels is ethanol. Most ethanol is produced by *Saccharomyces cerevisiae* (yeast) fermentation of either crops rich in sucrose (e.g., sugar cane and sugar beet) or starch-rich crops (e.g., corn and starchy grains). Ethanol produced from these sources is termed a first-generation biofuel. Yeast fermentation can yield a range of additional valuable co-products that accumulate during primary fermentation (e.g., protein concentrates, water soluble metabolites, fusel alcohols, and industrial enzymes). Distillers' solubles is a liquid co-product that can be used in animal feed or as a resource for recovery of valuable materials. In some processes it is preferred that this fraction is modified by a second fermentation with another fermentation organism (e.g., lactic acid bacteria). Such two stage fermentations can produce valuable compounds, such as 1,3-propanediol, organic acids, and bacteriocins. The use of lactic acid bacteria can also lead to the aggregation of stillage proteins and enable protein aggregation into concentrates. Once concentrated, the protein has utility as a high-protein feed ingredient. After separation of protein concentrates the remaining solution is a potential source of several known small molecules. The purpose of this review is to provide policy makers, bioethanol producers, and researchers insight into additional added-value products that can be recovered from ethanol beers. Novel products may be isolated during or after distillation. The ability to isolate and purify these compounds can provide substantial additional revenue for biofuel manufacturers through the development of marketable co-products.

**Keywords:** bioethanol; fermentation; nutraceuticals; organic acids; fusel alcohols; thin stillage; added-value products; 1,3-propanediol; bacteriocins



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## 1. Introduction

The demand for fuel ethanol continues to grow, with global production projected to surpass 140 billion litres/year [1]. Sugarcane and maize continue to be dominant feedstocks for bioethanol production. As such, bioethanol production is projected to consume 25% and 14% of global sugarcane and maize by 2029 [1]. Bioethanol is typically produced using a range of fermentation processes, each specialized to utilize a narrow range of inputs. The inputs are classified based on feedstock types (e.g., sucrose based, starch based, lignocellulosic, and algal), as belonging to a specific generation [2]. The type of feedstock, nutrients, and fermentation conditions affect bioethanol yield, and the nature of co-products [2–4]. Fermentation co-products can also affect bioethanol yield, as some act as both yeast nutrients and antinutrients. While the yield of ethanol is a primary driver for ethanol production, the identification and valorization of fermentation co-products add value to ethanol production.

Yeasts, and particularly *Saccharomyces cerevisiae*, used in ethanol production are most efficient in warm (e.g., 20–30 °C) and acidic (pH between 4.5–6.5) environments [5]. During microbial fermentation, yeasts produce glycerol, as both an osmoprotectant [6–8] and for

metabolic recovery of NAD<sup>+</sup>. Yeast also acidifies their environment to promote growth through proton secretion, secretion of organic acids (discussed below), removal of buffering agents, and dissipation of carbon dioxide [5]. The most common yeast used in alcoholic fermentation, *Saccharomyces cerevisiae*, is preferred for rapid and efficient conversion of sugar solutions to solutions with correspondingly high ethanol concentrations. The latter solutions are readily distilled and dehydrated to yield products suitable for blending with gasoline. Dehydration processes to produce anhydrous ethanol include heterogeneous azeotropic distillation using solvents (e.g., benzene), extractive distillation with solvents and entrainers (e.g., salts), adsorption using molecular sieves, and pervaporation membranes (e.g., zeolite, silica, etc.) [9]. During fermentation with yeast, value-added co-products (e.g.,  $\alpha$ -glycerylphosphorylcholine) can accumulate along with ethanol, while fusel alcohols can be recovered during distillation and are often added to ethanol used in fuel applications. Common lower value co-products can also accumulate (e.g., acetic acid, succinic acid, and glycerol) that are more difficult to valorize, and nuisance coproducts can accumulate (methanol, hydrogen sulfide, and methyl mercaptan). A portion of the volatile co-products are volatile, toxic, and/or contribute to odors that can be co-distilled with the ethanol product. Multi-stage distillation and other purification technologies are required to remove these compounds. A group of alcohols and aldehydes (e.g., aldehydes, butanols, propanols, etc.) are naturally synthesized from amino acids through the Ehrlich pathway [10] and simple sugars during fermentation [11,12]. In addition to alcohols, organic acids can also accumulate (e.g., acetic acid, succinic acid, and lactic acid) [3,13–15] because of yeast metabolic processes and even metabolism by adventitious bacteria present during ethanolic fermentation. Secondary fermentation (two-stage fermentation) of stillage can affect the contents of these compounds.

Upon the completion of ethanolic fermentation, the distillers' grain waste by-product can be further upgraded and enriched (e.g., secondary fermentation) to produce additional added-value compounds (e.g., 1,3-propanediol), as well as a highly concentrated protein that can be utilized as feed for domestic animals [14,15]. Altogether, the production of these co-products and others can add considerable value to ethanol coproducts (Table 1). Therefore, to maximize the utilization of grain crops, it is beneficial to identify valuable co-products produced during the fermentation process. This review will examine several valuable co-products that accumulate during alcoholic fermentation ( $\alpha$ -glycerylphosphorylcholine, and fusel alcohols), followed by those produced during secondary fermentation (e.g., organic acids) of the distillers' grain co-products (whole and thin stillage), which are primarily from first generation ethanol production. The recovery and purification of these compounds can further add value and provide opportunities for increased utilization of grain crops and provide ethanol producers access to new markets.

**Table 1.** Market size of coproduct solutes from alcoholic fermentation [16–22].

Co-Products	Market Size (US Dollars)	Project Compound Annual Growth Rate (CAGR)
Ethanol	89.1 billion (2019)	4.8% by 2027
Acetic acid	8.92 billion (2019)	5.2% by 2027
Succinic acid	181.6 million (2019)	9.2% by 2022
Lactic acid	2.7 billion (2020)	8.0% by 2028
Glycerol	2.6 billion (2019)	4.0% by 2027
Nootropics	2.42 billion (2020)	12.7% by 2028
Dried distillers' grains with solubles	112.5 million (2020)	5.2% by 2026

## 2. Thin Stillage and Distillers' Grains

Following ethanolic fermentation and distillation processes, the by-product stillage contains much of the protein, oil, fiber, and non-starch carbohydrate that were not available to the yeast during fermentation. A common process for using these components starts with the separation of whole stillage into a liquid portion with suspended solids (thin

stillage) and wet solids (distillers' wet grains), using centrifugation, vibratory separation, or a press [23,24]. These by-products can then be further processed (i.e., drying and fractionation) or extracted into specific components.

Without any further processing, all or a portion of thin stillage can be returned or backset to the next fermentation. This practice replaces some of the water required to soak incoming feedstocks intended for fermentation [25]. Protein is a major nutrient remaining in the stillage, and many methods have been developed to recover stillage proteins. Grain thin stillage contains approximately 37% protein (*w/w* dry basis) [26], and research has been conducted to develop stillage protein concentrates. Physical clarification techniques using additives [27,28], gas flotation [29,30], centrifugation [31], and filtration [32,33] have been tested as approaches to produce protein-enriched solids from thin stillage. Another approach for improving stillage protein quality and concentration is through a two-stage fermentation strategy [15,34], where it is also possible to upgrade glycerol to higher value compounds. Where thin stillage is not suited for use as a feed, it may be used as a nitrogen-rich fertilizer for crops [35]. Another strategy is to pair thin stillage valorization with a protein extraction process. Protein extraction from oilseed meal requires the use of large volumes of solution to dissolve proteins before precipitation. Thin stillage has some dissolved protein, but it can be used as a solution for protein extraction from oilseed meal [36]. An economical use of thin stillage that avoids the need for evaporation while providing the benefit of the stillage as a nutrient solution involves simply providing stillage in the water for cattle. In this way, the stillage becomes a nutrient-rich water source [37].

Once thin stillage is separated by dewatering distillers' wet grains [38,39], the thin stillage can then be dried to a concentrated syrup called distillers' solubles (DS), which is useful as an animal feed component. Remaining solids or distillers' wet grains (DWG) can be dried to produce distillers' dried grains (DDG) for storage and shipping. An alternative practice is to add DS to the grains as they dry to produce dried distillers' grains with solubles (DDGS), a product that is commonly used with cattle feed [40]. Compared to wet feed products, DDG and DDGS have extended shelf-life and are more easily shipped. The sale of fermentation by-products for use as cattle feed generally provides 10–20% of the total revenue of ethanol production facilities [41] while avoiding revenue losses that would be incurred if co-product disposal was necessary. Fractionation of the DDGS can concentrate protein and generate fractions with high fiber contents to produce additional protein and fiber products [42]. In some rations, the higher fiber content of DDGS is undesirable. Producing a higher protein- and fat-content feed ingredient can improve the value of this product stream. In addition to the use of stillage products in animal feed, a portion DDGS proteins can be more readily solubilized and extracted for a wide variety of industrial uses (e.g., biopolymer production) [43].

DDG, produced during first-generation biofuels processes, can be used as a substrate for a second fermentation after pre-treatment that converts unhydrolyzed and unprocessed cellulose into fermentable sugars [44]. Pre-treatment conditions are like those employed in second-generation biofuel processes and can include physical (e.g., milling), chemical (e.g., alkaline treatments), physicochemical (e.g., steam or CO<sub>2</sub> explosion), or biological processes (e.g., enzymatic hydrolysis) [2]. Nonetheless, variability of DDG composition could affect uniform enzymatic digestibility, and fermentability. Modification of the overall process design to accommodate new processing steps, or variable input materials would affect the economics of biofuel production [44].

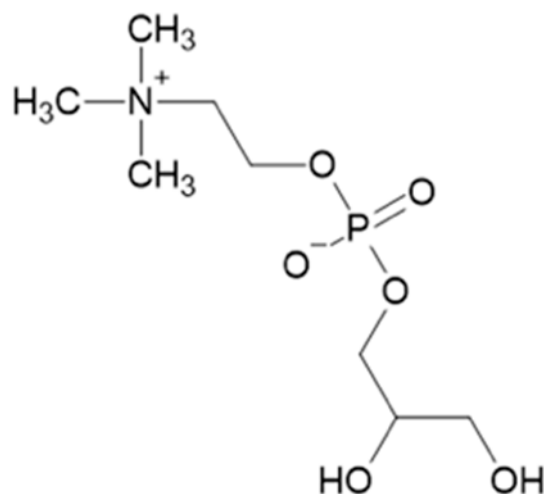
Alternatively, separation of fiber and germ, prior to fermentation, can enhance the value of non-starch/sugar nutrients in grain [45]. This can be accomplished through a combination of processes, including soaking, grinding, enzymatic hydrolysis, and sieving [45]. Wet milling ethanol production facilities can also implement front-end fractionation using conventional hydrocyclone systems [41], to further separate starch, protein, and fibre [46], prior to alcoholic fermentation.

Finally, oil is another product of bioethanol production that is often poorly utilized. For corn, methods of oil extraction from thin stillage have been patented [47], and other

oil extraction methods from corn DDGS have been developed [48]. However, due to the lower oil content of wheat and other grains, oil recovery techniques have remained largely undeveloped. In addition to fractionating fermentation products, other high-value compounds can also be extracted during the initial and two-stage fermentation processes.

### 3. $\alpha$ -Glycerylphosphorylcholine

$\alpha$ -Glycerylphosphorylcholine ( $\alpha$ -GPC) (Figure 1) is a yeast metabolite and precursor involved in the synthesis of acetylcholine and membrane phospholipids [49]. This compound has garnered interest in the natural products, medical, pharmaceutical, food, athletic performance, and cosmetic industries [50,51]. As  $\alpha$ -GPC is a source of choline, it has broad potential applications in foods related to health and performance. It is commonly marketed as a nootropic, due to its ability improve cognitive recovery and neurological function in healthy individuals, maintain neurological function after brain injury or mitigate deterioration of those affected by brain disease [49,52]. For example,  $\alpha$ -GPC has been used to improve the learning and memory abilities in stroke patients [53] and is also being investigated as a nootropic in treating psychiatric and neurological conditions (e.g., Alzheimer's, dementia, schizophrenia, etc.) [53–57]. Furthermore, relating to its applicability as a nootropic supplement,  $\alpha$ -GPC has been demonstrated to improve muscle strength [58]. With Alzheimer's disease predicted to increase three-fold, to affect 131.5 million people, by 2050 [58], the demand for  $\alpha$ -GPC could increase correspondingly, in addition to its use for muscle therapy. Collectively, the global market projections for nootropic supplements are predicted to surpass \$10 billion by 2025 [59], making this nootropic an attractive value-added product if it can be recovered efficiently.

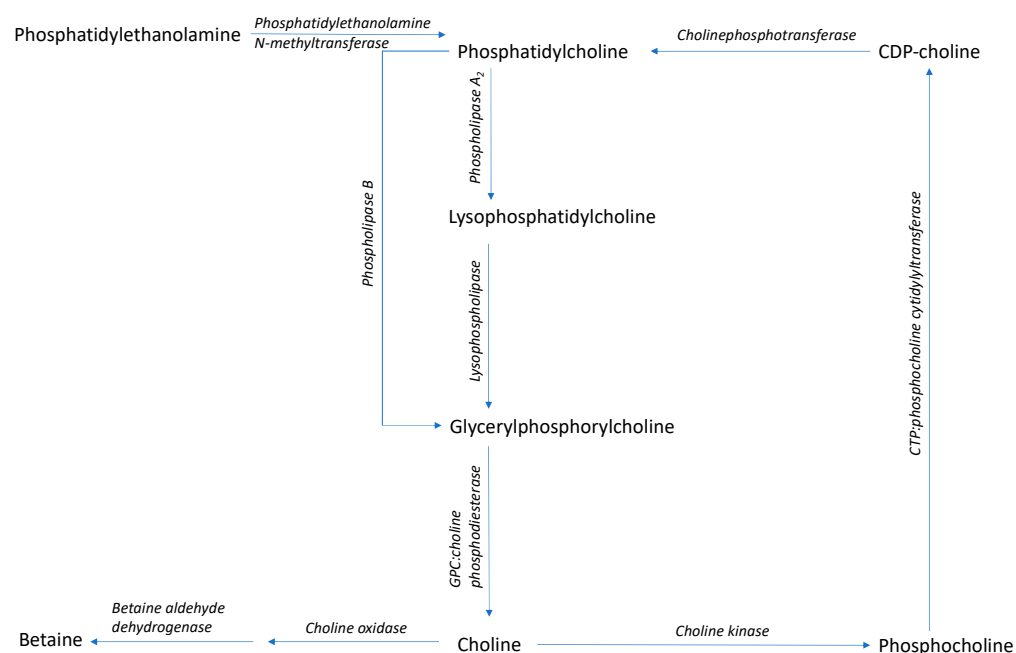


**Figure 1.** Chemical structure of  $\alpha$ -glycerylphosphorylcholine.

#### *Synthesis of $\alpha$ -Glycerylphosphorylcholine*

$\alpha$ -GPC can be produced chemically or with the use of enzymes. Chemical methods typically involve either hydrolysis of phosphatidylcholine (PC) or condensation of glycerol derivatives with phosphocholine donors using basic catalysts [60–62]. The chemical processes can produce toxic fumes and require the use of strong acids and harmful or undesirable solvents [62,63]. The use of toxic substrates can produce  $\alpha$ -GPC that is not safe for use in food and, thus, not marketable. Alternatively,  $\alpha$ -GPC has been produced by enzymatic hydrolysis of PC in aqueous media [52,57,64–66], employing phospholipases (Figure 2) [64]. Enzymatic production of  $\alpha$ -GPC is advantageous, as the amounts of chemical reagents can be reduced, thereby making a comparably inexpensive product that is suited for use in food and cosmetic products [64]. However, enzymatic production of  $\alpha$ -GPC can also be difficult due to the limited solubility of PC in aqueous phases and long

reaction times (e.g., low activity) for phospholipases [67–69]. Surfactants can be used to improve  $\alpha$ -GPC production, while being environmentally friendly [69].



**Figure 2.** Synthesis and metabolism pathway of  $\alpha$ -glycerylphosphorylcholine (Adapted from Li and Vance 2008 [70]; Gallazzini and Burg 2009 [71]). The blue area represents processes of the Kennedy Pathway, and the green area represents the synthesis of phosphatidylcholine via methylation of phosphatidylethanolamine.

Fermentation of ginger feedstocks using *Schizosaccharomyces pombe* yeast [72] has also resulted in the production of a nootropic compound (e.g., 6-paradol) [72]; however, these fermentations are often slower and produce by-products [73,74] that can make purification costly. Nonetheless, the ability to isolate and purify these nootropic coproducts can add significant revenue for bioethanol producers, by providing ethanol producers with new products that meet the needs of other markets. For example, nootropics (e.g.,  $\alpha$ -GPC) had a value of 7.21 billion USD in 2020, and they are expected to grow with a CAGR of 80% until 2028 [75]. Accumulation of  $\alpha$ -GPC in the mash has been observed during alcoholic fermentation of cereal crops, such as wheat, barley, and oat, although the concentration produced can differ among cultivars [3,13]. For example, fermentation of 28 barley and 12 oat cultivars resulted in the accumulation of between 0.84 g/L to 1.81 g/L for barley and 0.62 g/L to 0.88 g/L for oat, depending on cultivar [3]. Meanwhile, fermentation of wheat resulted in  $\alpha$ -GPC accumulation of approximately 1.68 g/L [13]. Oyeneye et al. also found that treatment of the grain with phospholipase A1, an enzyme that readily hydrolyses phosphatidyl choline and lysophosphatidyl choline, produced a beer with higher  $\alpha$ -GPC than other treatments. This treatment effect strongly suggests that  $\alpha$ -GPC might accumulate as a result of phosphatidyl choline hydrolysis. Pre-treatment of the feedstock (e.g., soaking, germination, incubation temperature, etc.) can also influence  $\alpha$ -GPC yield (unpublished data). Therefore, alcoholic fermentation of cereal feedstocks can be highly advantageous due to the inexpensive processes involved and because  $\alpha$ -GPC can be concurrently produced with ethanol during fermentation.  $\alpha$ -GPC is a naturally produced endogenous choline derivative; however, it is rarely found at high concentrations in nature. Therefore, there is great potential in developing alternative, inexpensive, and sustainable means for commercial production to supply this compound.



#### 4. Fusel Alcohols

A series of primary alcohols are found as natural co-products, generated during the fermentation process. These compounds are largely produced by metabolic processes, called the Ehrlich pathway in yeast, that recover nitrogen required for growth and metabolism. Enzymes of the pathway catalyze the transfer of amines between amino acids and ketones and the decarboxylation of  $\alpha$ -keto acids to produce aldehydes, and they reduce aldehydes to form fusel alcohols (Figure 3). Products from the Ehrlich pathway can be influenced by the presence of certain amino acid intermediates (Table 2).

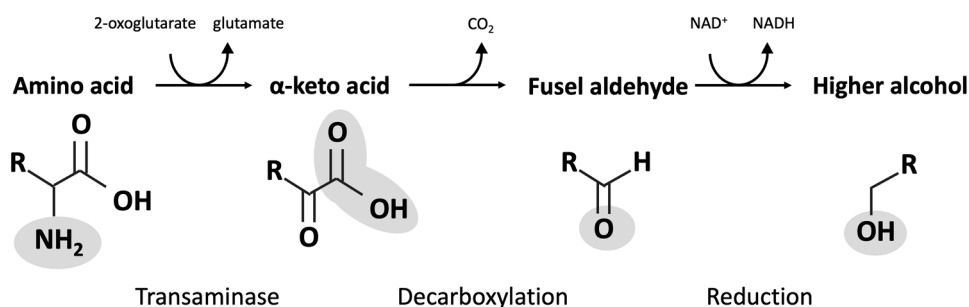


Figure 3. Ehrlich pathway conversion of amino acids into fusel alcohols.

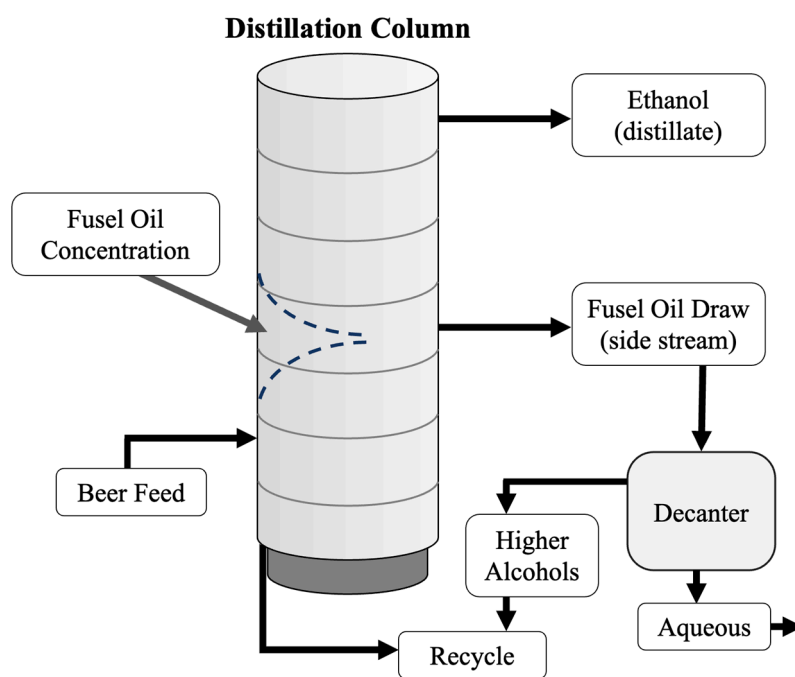
Table 2. Amino acids intermediates and products of the Ehrlich pathway. Reconstructed from Hazelwood et al., (2008) [10].

Amino Acid	$\alpha$ -Keto Acid	Fusel Aldehyde	Fusel Alcohol
Isoleucine	$\alpha$ -Ketomethylvalerate	Methylvaleraldehyde	Active amyl alcohol
Leucine	$\alpha$ -Ketoisocaproate	Isoamylaldehyde	Isoamyl alcohol
Methionine	$\alpha$ -Keto- $\gamma$ -	Methional	Methionol
Phenylalanine	(methylthio)butyrate	Phenylethanal	Phenylethanol
Threonine	Phenylpyruvate	Propanal	Propanol
	2-Ketobutyrate		
Tryptophan	3-Indole pyruvate	3-Indole acetaldehyde	Tryptophol
Tyrosine	<i>p</i> -Hydroxyphenylpyruvate	<i>p</i> -Hydroxyphenylacetaldehyde	<i>p</i> -Hydroxyphenylethanol or tyrosol
Valine	$\alpha$ -Ketoisovalerate	Isobutanal or isovaleraldehyde	Isobutanol

Fusel alcohols are somewhat volatile, and as such, distillation enables their separation from fermented mash. These are a mixture of primarily alcohols, including active amyl alcohol (2-methyl-1-butanol), isoamyl alcohol, isobutyl alcohol, and, in lesser amounts, n-amyl alcohol, n-butyl alcohol, and methionol. Less volatile alcohols are also present in the mash and are poorly extracted by distillation, including phenethyl alcohol and tyrosol. Some of these alcohols are aromatic and are associated with strong tastes and pungent odours. Used sparingly, fusel alcohols and esters can contribute positively to foods and beverage flavours, but at higher concentrations, they are associated with unpleasant flavours and “hangover” symptoms [76]. These compounds can be detrimental to yeast growth; therefore, their removal is essential in maintaining efficient fermentation [10,76–78]. When these materials are extracted from backset or continuous distillation processes, the fermentation efficiency is increased, and a valuable co-product can be isolated.

During ethanol distillation, fusel alcohols concentrate in the distillation column as they are less volatile than the ethanol water azeotrope. The removal of these compounds is not required for fuel production but is essential for ethanol destined for food or pharmaceutical applications. If these higher alcohols are not removed, they can reach their limit of solubility and increase vapour pressures in the column. In turn, the increase in vapour pressure can cause boiling and flooding and thus interfere with distillation. In bioethanol plants with continuous distillation processes, the fusel alcohols are distributed in the distillation or rectifier column, where the product at the top of the column is purified alcohol (~95% by

wt.), and at the bottom is mostly water. Fusel alcohols are most volatile at lower ethanol concentrations and tend to collect in the intermediate region where ethanol concentration is approximately 45% (v/v) [76]. In continuous distillation columns, fusel alcohol removal is accomplished by drawing a solution of water, ethanol, and fusel alcohols from the center of the rectifier (Figure 4).



**Figure 4.** Distillation column with a fusel alcohol draw near the middle of the column. The fusel alcohols can be removed from the distillation process and decanted, reusing the higher alcohols and removing the aqueous phase. Without a fusel alcohol draw, the fusel alcohols will fall in the column and form a two-phase system. This can interfere with the distillation efficiency.

In batch distillation, fusel alcohols form azeotropes with water that can accumulate at the bottom of the column to become concentrated late distilling fractions called “tails” that evaporate when the column temperature rises at the end of a distillation.

The accumulation of fusel alcohol in commercial fermentations can vary from 1–11 mL/L of ethanol produced [79]. With global biofuel production capacity increasing, the capacity for the production of substantial amounts of fusel alcohol is possible. For example, the US Energy Information Administration announced that the total US biofuels plant production capacity reached 21 billion gallons per year as of January 2021 [80]. If fusel oils were separated from fuel ethanol, hundreds of millions of litres of fusel alcohols could be recovered annually. Ultimately, it is beneficial to investigate additional uses for these compounds to promote sustainability and discover new market values for these co-products. Typically, fusel alcohols are blended with ethanol to make a product that is suited for fuel applications. Separated fusel oils can also be used as an energy source, but this is not an ideal fuel, as negative environmental impacts can outweigh the value of energy recovered when combusting these materials [81]. Despite some of the drawbacks in fermentations that accumulate fusel alcohols, they have utility as gasoline and diesel additives that improve fuel properties and combustion [5,10,76–78,82]. In particular, fusel alcohol gasoline blends do not exhibit any phase separation, and engines operating on these blends can achieve higher compression ratios and performance than possible with gasoline alone.

Fusel alcohols can also be purified and esterified to yield a range of valuable esters. Isoamyl alcohol is the primary component in fusel alcohols and can be used in the production of organic esters for industrial purposes. For example, isoamyl acetate is formed by esterification of isoamyl alcohol with acetic acid, and it has applications in commercial

products such as lacquers and as a food flavouring additive, with a banana-, caramel-, or pear-like flavour profile [76–78]. As such, the recovery of isoamyl alcohol can add significant value to the bioethanol producer, as this compound has three-times the market value of fuel ethanol [81]. Unfortunately, isolating and purifying isoamyl alcohol from fusel alcohols is challenging, due to the formation of heterogeneous azeotropes of several alcohols with water during distillation. Multiple decanters have been used to collect fusel alcohol draws at alternating positions in the distillation column, to separate and purify fusel alcohols, although these methods typically do not afford products that are sufficiently pure for commercial application [76,79,81]. Esterification of fusel alcohols, a process called reactive distillation, has also been proposed and demonstrated successfully at a bench-scale to simultaneously obtain highly enriched alcohol esters [76,77]. Another method to aid in fusel alcohol separation is the addition of water to promote phase separation of higher alcohols from the aqueous phase (containing methanol, ethanol, propanol, and isopropanol). The organic phase can then be further distilled to enrich isoamyl alcohol [76,79].

Enrichment and purification of individual fusel alcohols and their esters can produce value-added compounds, suitable for inclusion in flavours and fragrances. The accumulation of these alcohols depends greatly on yeast genetics [76–78] and fermentation medium components (e.g., sugar source) [76,77]. For example, the production of isopropanol can be manipulated through the addition of exogenous acetone to the mash [76,77]. Yeast genetics determines the production of volatile compounds, such as fusel alcohols, that contribute to the product flavours and aromas [77,83,84]. Identifying yeasts that appropriately influence the flavour profiles of the final product is of great importance for the distillation of materials such as beers, ciders, and spirits. Alteration of fusel compounds is possible, although the complete suppression of fusel alcohol formation by selection of yeast is not, due to the biosynthesis of the  $\alpha$ -keto acid present in the Ehrlich pathway (Figure 3) [83,85,86]. Overall, modifications can influence the production of alcohol by-products, and they can add considerable value to the bioethanol producer if these compounds can be further isolated and purified.

## 5. Two-Stage Fermentation of Thin Stillage

In addition to physical processing, stillage can be fermented by a range of organisms. The biorefinery of thin stillage has been accomplished via fermentation with a consortium of LAB (e.g., *Lactobacilli*) selectively recovered from a stillage storage tank (Figure 5) [14,15]. Fermentation was effective in modifying the stillage, allowing the efficient separation of a protein-rich fraction. The consortium organisms belong to species that are classified as Generally Recognized as Safe (GRAS) and are routinely used in the food industry. Clarification of thin stillage via LAB fermentation may result from a combination of gas production that leads to anoxic gas flotation [87], production of exopolysaccharides that cause particle aggregation [14], and/or allowing sufficient time for settling/floatation of aggregated particles or separation of the particles from the solution with decanting and desludging centrifuges. The resulting protein-rich slurry contains much of the original thin stillage protein, as well as proteins produced by the LAB. Drying the slurry creates a concentrate of up to 60% protein [88]. The bacteria-rich protein concentrate can then serve as a probiotic animal feed supplement [89]. Furthermore, LAB are capable of utilizing and thriving on complex carbohydrates as their carbon source. Two-stage fermentation of wheat-based thin stillage using LAB produced succinic acid (>2.0 g/L), lactic acid (>4.5 g/L), and acetic acid (>4.5 g/L) within 72 h at 37 °C [15].





**Figure 5.** Wheat thin stillage before (left) and after (right) 48 h of secondary fermentation by lactic acid bacteria. The initially homogenous solution separates into an upper layer composed of, mostly, clear liquid and a protein rich slurry in the bottom layer.

### 5.1. Organic Acids

Many of the organic acids (e.g., acetic acid, succinic acid, and lactic acid) produced by yeast are acids that typically arise from glycolysis (e.g., acetic acid and lactic acid) and the citric acid cycle (e.g., succinic acid) [90]. These compounds are typically produced commercially via fermentation microbial processes [90–92] for use in the food, beverage, and manufacturing industries. The efficiency of production of these organic acids is dependent on the microorganisms, feedstock, fermenter productivity, and development of efficient recovery processes [93–95].

Due to its pH tolerance and simple nutrient requirements, *Saccharomyces cerevisiae* has been investigated and genetically modified for organic acid production [96]. Organic acids are not traditionally produced by *Saccharomyces cerevisiae* fermentation. The production of these organic acids could produce considerable additional value (Table 1), but a portion of the organic acids (e.g., acetic acid and lactic acid) seen in commercial ethanol beers are typically attributable to nuisance organisms present during bioethanol production. At modest concentrations, these organic acids can inhibit fermentation [5,97,98]. For example, acetic acid present in commercial *Saccharomyces cerevisiae* fermentations is undesirable, with concentrations > 0.4 g/L signifying bacterial contamination and concentrations > 0.6 g/L leading to impaired fermentations [95,99]. Thus, during ethanolic fermentation, it is often preferred to minimize production of these compounds. However, after the completion of primary ethanolic fermentation, the resulting thin stillage by-product can be used as culture media for lactic acid bacteria (LAB) fermentation (two-stage fermentation; discussed below) for producing additional succinic acid, lactic acid, and acetic acid [14,15]. For example, fermentation (72 h at 37 °C) on wheat-based thin stillage with an initial glycerol content of 10 g/L using *Lactobacilli* resulted in the accumulation of succinic acid (>2.0 g/L), lactic acid (>4.5 g/L), and acetic acid (>4.5 g/L) in the stillage medium [15].

### 5.2. Conversion of Glycerol to 1,3-Propanediol

In addition to the production of organic acids, during fermentation, glycerol is also produced by yeast to protect cells against lysis and regenerate  $\text{NAD}^+$  needed for glycoly-

sis [6–8]. Interestingly, glycerol in the fermentation mash can be further upgraded through its conversion to 1,3-propanediol using *Lactobacilli* [14,15]. This compound is useful for the production of textiles, carpets, adhesives, moldings, etc. [100,101]. The conversion of glycerol to 1,3-propanediol is catalyzed via a two-step reaction. First, glycerol is converted to 3-hydroxypropionaldehyde by glycerol dehydratase, and subsequently, it is converted to 1,3-propanediol by 1,3-propanediol oxidoreductase [101,102]. Glycerol dehydratase and 1,3-propanediol oxidoreductase activity require cobalamin (aka vitamin B<sub>12</sub>) as a cofactor for catalysis [101,102]. Furthermore, cobalamin is an essential nutrient for DNA synthesis and cellular energy production. Some lactic acid bacteria (e.g., *Lactobacillus reuteri*) can produce this essential nutrient [103–106]. A two-stage fermentation of wheat-based thin stillage with a consortium of *Lactobacilli* converted most glycerol (10 g/L) present to 1,3-propanediol (6.1 g/L) [14,15] within 72 h. Genetic sequencing of members of the consortium identified *Lactobacilli* gene sequences that encoded for proteins that likely produce cobalamin [15]. The inoculum size for this study was 0.01% (v/v), and thus, it could be easily implemented at other bioethanol facilities, to facilitate the production of these valuable compounds and vitamins.

### 5.3. Bacteriocins

Bacteriocins are another potential value-added product that could be purified after a second two-stage fermentation of ethanol stillage with *Lactobacillus*. Bacteriocins are antimicrobial proteins produced by most bacteria [107,108]. This broad class of compounds inhibits the growth of competing bacteria. Typically, the bacteria that produce bacteriocins simultaneously produce immunity proteins [109]. Some bacteriocins exhibit broad-spectrum antagonistic effects. For example, bacteriocins produced by Gram-negative bacteria typically affect closely related species, whereas Gram-positive bacteria can produce bacteriocins that exhibit a broader spectrum of activity [110]. These antimicrobial compounds might have utility as natural food preservatives or for pharmaceutical applications [110,111]. Bacteriocins derived from Generally Recognized as Safe (GRAS) organisms would have greater potential for such applications (e.g., *Lactobacilli*). Genes encoding known bacteriocin proteins were present in a consortium of *Lactobacilli* that was capable of two-stage fermentation of wheat-based thin stillage [15]. It is not known if the identification of bacteriocins was complete, as most bacteriocins and their sequences remain unidentified, and only a few have been investigated for their utility in foods as antagonistic compounds [112]. Furthermore, growth media composition [113], fermentation temperature [114], and pH [115] can also affect bacteriocin production and yield.

Currently, there are five recognized classes of bacteriocins that are segregated primarily on their molecular size and properties [34]. Unlike traditional antibiotics, which typically act as enzyme inhibitors [116], bacteriocins elicit adverse effects by inhibiting bacterial cell growth, by disrupting essential functions (e.g., translation and transcription) [117], and by targeting the cell surface and altering membrane permeability [110,117–121] (e.g., formation of membrane channels). Bacteriocin-producing organisms simultaneously express bacteriocin-immunity proteins that protect producing organisms from their own toxins [122,123]. Bacteria can acquire or lose immunity against specific bacteriocins through horizontal gene transfer [124–130].

Although the applications for bacteriocins as a food preservative and for pharmaceutical use are promising, the purification of bacteriocins can be difficult. This is likely due to their complex molecular structure, physicochemical properties, and heterogeneity [110]; therefore, specific purification processes might be required for individual bacteriocins [110]. These difficulties and current separation and approaches to bacteriocins purification are further reviewed in Tse and Reaney (2020) [34].

## 6. Spent Yeast

Spent yeast is another by-product from the brewing and biofuels industry. The spent yeast is typically removed at the end of the fermentation, although a small amount

can be retained for subsequent fermentation batches [131]. Due to its protein content, discarded yeast is typically used as inexpensive animal feed materials [131]. However, spent yeast also contains valuable nutrients, such as vitamins (e.g., vitamins B<sub>1</sub>, B<sub>2</sub>, B<sub>3</sub>, B<sub>6</sub>, B<sub>9</sub>, and B<sub>12</sub>) [132–134], minerals (e.g., Na, K, Ca, Mg, Fe, Mn, Zn, Cu, Se, Cr, and Mo) [134,135], proteins (e.g., mannoproteins and hydrolysates) [135], carbohydrates (e.g.,  $\beta$ -glucans) [135], antioxidants (e.g., glutathione) [136], and phenolic compounds (e.g., gallic acid and ( $\pm$ )-catechin) [132]. Retrieval of these compounds can have added value in animal feeds, nutritional supplements, and functional foods (e.g., flavor enhancers) and non-food additives (e.g., cosmetics) [131,137]. Hydrolysis of spent yeast can yield a complex mixture of oligopeptides, peptides, and free amino acids, also known as hydrolysates [137]. Total solids recovery of hydrolysates, proteins, and  $\alpha$ -amino nitrogen content in dried spent yeast was reported to be 50%, 55.9%, and 4.8%, respectively [137]. However, the composition of the spent yeast (e.g., the presence of contaminating bacteria) during fermentation can have direct implications on the yeast extract and concentration of nutrients in these fractions [138]. Furthermore, as yeast are living cells with vigorous metabolism, they have a high ratio of RNA to other nutrients. Consumption of large amounts of nucleic acids can impart detrimental health effects to foods. Consumption of nucleic acid-rich materials can lead to uric acid accumulating in tissues and the consequent symptoms of gout [131]. Therefore, processing of spent yeast should include RNA degradation processes, prior to its application in food supplements.

## 7. Enzymes and Pharmaceuticals Products Produced via Microbial Fermentation

Hydrolytic enzymes have a market value close to a \$1 billion/year, and they play an important role in many industries, including in the preparation of pharmaceuticals, cosmetics, medicines, nutritional supplements, chemicals, beverages, and foods (Table 3) [139]. Although hydrolytic enzymes and pharmaceutical products may be recovered from microbial fermentation, other methods are more effective, so they are not described in detail here. Solid-state and submerged fermentations have been employed for enzyme production [140–146], with the former method demonstrating advantages for certain enzymes' production (e.g., invertase, pectinases, and tannases from *Aspergillus* sp.), rather than liquid-based fermentation systems [145,147]. Solid-state fermentation can also be more productive than liquid systems while being simpler technically and requiring lower capital investment, lower energy input, lower water requirement, better product recovery, and a lack of foam build up when compared to submerged fermentations [148]. Improved enzyme production has also been demonstrated using co-cultivation methods [145]. Nonetheless, both submerged and solid-state fermentation technologies are being investigated and improved upon to increase industrial enzymes production.

**Table 3.** Examples of industrial enzymes and the microbial producers. Reconstructed from Ventura-Sobreville et al., 2015 [139]; de Souza Vandenberghe et al., 2016 [149]; Singh et al., 2019 [150], and citations therein.

Enzyme Type	Enzyme	Application	Microbial Producer
Oxidoreductases	Glucose oxidase	Food industry [149,150]	<i>Aspergillus</i> sp. [151]
	Lactases	Food industry [149,150]	Bacteria ( <i>Lactobacillus</i> sp.), fungi ( <i>Aspergillus</i> sp.), yeast [152–154]
Transferases	Glycosyltransferases Fructosyltransferases	Food, cosmetics, pharmaceutical industries [149,150]	Bacteria (e.g., <i>Bacillus</i> sp., <i>Klebsiella</i> sp., <i>Geobacillus</i> sp., <i>Thermoanaerobacter</i> sp.) [155–157]
	Transglutaminase	Food industry [149,150]	<i>Bacillus</i> sp., <i>Streptomyces</i> sp. [158,159]
Hydrolases	Amylases	Food, brewing, and bioethanol industries [149,150]	<i>Bacillus</i> sp., <i>Lactobacillus</i> sp., <i>Aspergillus</i> sp., <i>Mucor</i> sp., <i>Saccharomyces</i> sp. [148]
	Cellulases and hemicellulases	Food and biofuel industries [149,150]	<i>Aspergillus</i> sp., <i>Trichoderma</i> sp., <i>Bacillus</i> sp., <i>Cellulomonas</i> sp., <i>Clostridium</i> sp. [149,160]
	Chitinases	Pharmaceutical industry [150]	<i>Bacillus</i> sp., <i>Streptomyces</i> sp., <i>Talaromyces</i> sp., <i>Trichoderma</i> sp., <i>Nocardia</i> sp. [161]
	Invertase	Food industry [149,150]	<i>Aspergillus</i> sp., <i>Saccharomyces</i> sp. [162]
	Lipases	Food industry [149,150]	<i>Aspergillus</i> sp., <i>Bacillus</i> sp., <i>Rhizopus</i> sp., <i>Trichosporon</i> sp., <i>Lactobacillus</i> sp., <i>Penicillium</i> sp., <i>Pseudomonas</i> sp. [149]
	Mannanases	Food, animal feed, and biorefinery industries [163]	<i>Bacillus</i> sp., <i>Trichoderma</i> sp., <i>Aspergillus</i> sp., <i>Thermomyces</i> sp., <i>Rhizopus</i> sp. [149]
	Pectinases	Food and animal feed industry [149,150]	<i>Aspergillus</i> sp., <i>Rhizopus</i> sp., <i>Penicillium</i> sp. [149]
	Phytases	Food, animal feed, and bioethanol industries [149,150]	<i>Aspergillus</i> sp., <i>Lactobacillus</i> sp., <i>Saccharomyces</i> sp., <i>Bacillus</i> sp., <i>Candida</i> sp., <i>Pseudomonas</i> sp. [149]
	Proteases	Food industry [149,150]	<i>Bacillus</i> sp., <i>Aspergillus</i> sp., <i>Pseudomonas</i> sp., <i>Synergistes</i> sp., <i>Rhizopus</i> sp. [149]
	Xylanases	Food, pulp, and ethanol industries [149,150]	<i>Aspergillus</i> sp., <i>Rhizomucor</i> sp., <i>Bacillus</i> sp. [149]
Peroxidase	Peroxidases	Pharmaceutical industry [150]	<i>Bacillus</i> sp., <i>Ensifer</i> sp. [164,165]
Acylase	Penicillin acylase	Pharmaceutical industry [150]	<i>Bacillus</i> sp., <i>Escherichia</i> sp. [166–168]

## 8. Conclusions

In conclusion, microbial fermentation is often utilized to produce a wide variety of valuable compounds used in several industries. First-generation biofuel producers utilizing *Saccharomyces cerevisiae* for grain fermentation can potentially generate substantial revenue through the production, isolation, and purification of numerous added-value

co-products. These products can be co-produced during initial ethanolic fermentation (e.g., ethanol and  $\alpha$ -GPC), distillation (e.g., fusel alcohols), and two-stage fermentation processes (e.g., 1,3-propanediol, organic acids, essential nutrients, and high protein domestic feed). The application of two-stage fermentation using LAB opens the possibility of not only upgrading and enriching thin stillage products, but it is also safe for feed production, due to the GRAS status of known Lactobacilli. Fortunately, these technologies have demonstrated success in producing value-added products, and they can be rapidly implemented into existing facilities, and downstream LAB fermentation utilizes minimal inoculant (0.01% *v/v*). However, production yields of these products can vary depending on the composition of the feedstock, the fermentation medium, microorganisms present, and fermentation conditions. In addition, complications can arise in the development of isolation and purification for these compounds. Nonetheless, the co-production of these added-value compounds will not only increase the value and utilization of grain crops, but it can also provide ethanol producers with significant additional revenue and new market entries for these fermentation co-products.

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