

# Yeast derived from lignocellulosic biomass as a sustainable feed resource for use in aquaculture

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## Abstract

The global expansion in aquaculture production implies an emerging need of suitable and sustainable protein sources. Currently, the fish feed industry is dependent on high-quality protein sources of marine and plant origin. Yeast derived from processing of low-value and non-food lignocellulosic biomass is a potential sustainable source of protein in fish diets. Following enzymatic hydrolysis, the hexose and pentose sugars of lignocellulosic substrates and supplementary nutrients can be converted into protein-rich yeast biomass by fermentation. Studies have shown that yeasts such as *Saccharomyces cerevisiae*, *Candida utilis* and *Kluyveromyces marxianus* have favourable amino acid composition and excellent properties as protein sources in diets for fish, including carnivorous species such as Atlantic salmon and rainbow trout. Suitable downstream processing of the biomass to disrupt cell walls is required to secure high nutrient digestibility. A number of studies have shown various immunological and health benefits from feeding fish low levels of yeast and yeast-derived cell wall fractions. This review summarises current literature on the potential of yeast from lignocellulosic biomass as an alternative protein source for the aquaculture industry. It is concluded that further research and development within yeast production can be important to secure the future sustainability and economic viability of intensive aquaculture.

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**Keywords:** yeast; lignocellulosic biomass; wood; nutritional value; fish health; aquaculture

## INTRODUCTION

Fish protein now contributes 17% of the global human population's intake of animal protein, and aquaculture accounts for a rapidly increasing share.<sup>1</sup> The increased feed demand of the rapidly expanding aquaculture sector has led to a concern that available feed resources will limit further growth. Fishmeal and fish oil were historically the major sources of protein and lipid in the intensive farming of carnivorous fish, and salmon farming has been criticised for reducing the amount of fish protein and lipids for human consumption.<sup>2,3</sup> Currently, there is insufficient fish oil and fishmeal on the world market to meet demand.

The recent development of salmon farming has shown reduced dependence on marine ingredients in the feed.<sup>4,5</sup> Following the rapid expansion in aquaculture and limited resources of fishmeal, plant ingredients have become the main protein sources in diets for most aquaculture species.<sup>6,7</sup> Thus, Norwegian salmon feed contained 66% plant ingredients in 2012.<sup>5</sup> Currently, the preferable plant protein sources in diets for carnivorous fish are refined and expensive products such as soybean protein concentrate. These plant ingredients are directly applicable for human consumption, and the use of human food as feed ingredients is doubtful, for ethical as well as economic reasons. Sustainability of food production should aim at maximised nutritional output for human consumption and minimised input of resources, with the lowest possible impact on the environment.<sup>5</sup> The change to plant ingredients implies that aquaculture uses large land areas and water resources. As arable land and clean water resources are already limited, this means that the sustainability of substituting marine feed

ingredients with high quality plant products in fish feed is questionable.<sup>5</sup> Farmed fish species such as the Atlantic salmon (*Salmo salar*) are, however, more efficient than terrestrial farmed animals in retaining protein and energy.<sup>4,5</sup>

Microbial products, particularly yeast, are potential sustainable ingredients in aquafeeds due to the ability to convert low-value non-food biomass from forestry and agricultural industry into high-value feed with limited dependence on arable land, water and changing climatic conditions.<sup>8</sup> Under-utilised wood and co-products from agriculture and forestry can provide resources for production of feed ingredients from lignocellulosic biomass. Microbial conversion of the biomass to liquid biofuel has been an attractive research area in recent years.<sup>9,10</sup> Use of lignocellulosic biomass for production of high-value lipids and chemicals has also received considerable attention,<sup>11–13</sup> while there has been less interest in yeast biomass as an alternative protein source. Yeast cells are able to synthesise all precursors required for macromolecular constituents such as protein and nucleic acids from sugars and essential nutrients like nitrogen, inorganic phosphate and sulfate, and additional minerals and vitamins. The composition of yeast is dependent on strain, growth media and growing

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conditions; the content of crude protein varies between 40% and 55% of the dry matter including contents of nucleic acids.<sup>14–16</sup> The potential of different products from *Saccharomyces cerevisiae* for use in fish feed has been reviewed by Ferreira *et al.*<sup>17</sup>

The conversion of cellulose and hemicellulose to protein-rich biomass and ethanol by yeast fermentation is well established. *Torula* yeast, or *Candida utilis*, has been commercially available for more than 70 years as a nutritional supplement in animal feed.<sup>18</sup> During World War II, the Germans cultured *Torula* yeast as a protein source based on sulfite waste liquor from pulp and paper manufacture and wood sugar obtained by acid hydrolysis of wood.<sup>19</sup> Scientific progress and advances in technology have improved industrial yeast production from a wide range of lignocellulosic sources, including wastes or by-products from forestry and agriculture.<sup>20,21</sup> Recent studies with Atlantic salmon have documented excellent nutritional properties of *C. utilis* and *Kluyveromyces marxianus* yeasts grown on lignocellulosic biomass.<sup>8</sup> Both of these yeast species can metabolise a wide range of monosaccharides obtained from lignocellulosic biomass.<sup>22–24</sup> This is promising for future exploitation and economical industrial-scale production of yeast products based on lignocellulosic biomass as sustainable protein sources. Like *S. cerevisiae*, strains of *C. utilis* and *K. marxianus* have obtained the generally-regarded-as-safe (GRAS) status assigned by the US Food and Drug Administration (FDA) to substances not known to be hazardous to health and which are approved for use in foods.

Recent research on effects of yeast products in fish diets has focused mainly on the role as immunostimulants, gut health promoters and intestinal microbiota modulators at low inclusion rates. The literature pertaining to yeast products as a major nutrient source for different fish species is scarce, despite positive outlooks on future needs and prospects on new production technology development. The objective of this review is to summarise current knowledge on the potential of yeast products produced from lignocellulosic substrates as sustainable protein sources in aquaculture.

## YEAST PRODUCTION FROM LIGNOCELLULOSIC BIOMASS

In the late 20th century, the use of first-generation feedstock dominated the development of biofuel, but concerns exist about the impact this may have on biodiversity, water and land use, and competition with human food. So there is an increasing interest in second-generation lignocellulosic biomass such as by-products from the agricultural and forestry sectors, as this represents an abundant, natural, renewable and cheap resource for biorefinery. Lignocellulosic biomass is a heterogeneous complex of the carbohydrate polymers cellulose and hemicellulose, and the aromatic lignin. Cellulose is generally the largest fraction in woody biomass, representing about 35–50% of the biomass weight, while hemicellulose represents about 20–30%.<sup>25</sup> Woody biomass has higher lignin content of about 20–30% and is structurally stronger and denser than agricultural biomass.<sup>25</sup> Woody biomass generally has a lower content of pentoses than agricultural biomass, but hard wood has a higher content of pentoses than soft wood species.<sup>26</sup>

Processing of lignocellulosic biomass for yeast production requires four major steps: (1) pre-treatment, (2) enzymatic hydrolysis, (3) fermentation and (4) downstream processing. Figure 1 shows the main steps in using woody biomass in the production of yeast. Pre-treatment of lignocellulosic biomass is essential to break the hemicellulose–lignin complex, separate out the lignin, and

disrupt the crystalline structures of cellulose to make cellulose and hemicellulose more accessible before enzymatic hydrolysis to monosaccharides. The pre-treatment of the feedstock includes chemical, physical, thermal and biological methods.<sup>10</sup> Pre-treatment and saccharification of woody biomass, especially from softwood species, differ from non-woody materials because it is more resistant to microbial and enzymatic actions and requires more energy. Pre-treatment processing conditions must therefore be adapted to the properties of each different source of lignocellulosic biomass to get maximum yield of sugars for yeast production.<sup>21,27</sup>

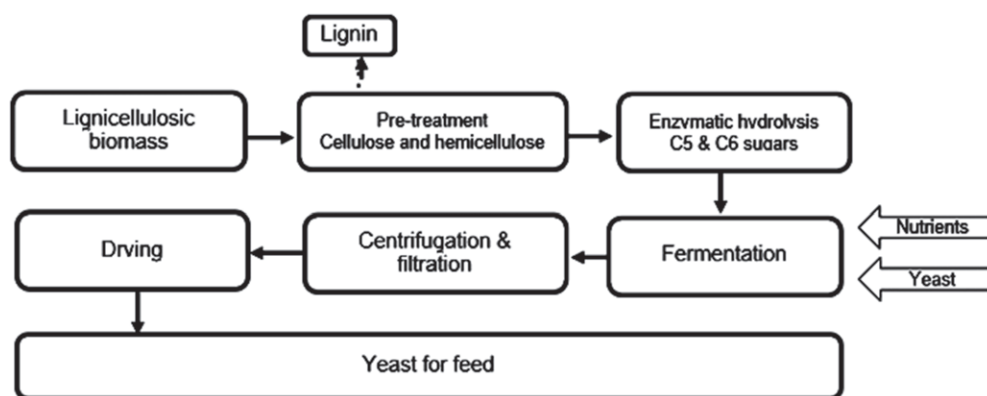
Efficient conversion of crystalline polysaccharides to fermentable sugars is crucial for an economical and sustainable yeast production from lignocellulosic biomass. Maximum hydrolysis to monomeric sugars requires appropriate delignification and a cocktail of biomass-degrading enzymes.<sup>10,28</sup> Recent findings on enzymes that catalyse cleavage of glycosidic bonds in recalcitrant polysaccharides show the potential for increased efficiency of enzymatic conversion.<sup>29,30</sup> These enzymes increase the efficiency of hydrolytic enzymes by acting on the surface of the insoluble substrate.<sup>30</sup> Enzymatic hydrolysis may be carried out as separate hydrolysis and fermentation or simultaneous saccharification and fermentation.<sup>13,28</sup>

The sugars from lignocellulosic biomass can be converted into yeast biomass under aerobic conditions by fermentation, using suitable yeast strains and supplementary essential nutrients like nitrogen, inorganic phosphate and sulfate. *S. cerevisiae* strains are highly efficient in metabolising hexose sugars, and is predominant in ethanol production using cane and beet molasses as substrate. Wild-type strains of *S. cerevisiae* cannot ferment pentoses, but population genetics has been applied to develop non-recombinant *S. cerevisiae* strains that can grow on xylose.<sup>31</sup> Genetic engineering is also used to generate strains of *S. cerevisiae* that can grow on lignocellulosic feedstocks.<sup>31–33</sup> Fermenting mixed carbon components derived from lignocellulosic biomass is a challenge to cost-efficient production of yeast protein. However, recently, Wei *et al.*<sup>34</sup> were able to integrate the fermentation pathways of hexose and pentose sugars and an acetic acid reduction pathway into one *S. cerevisiae* strain using synthetic biology and metabolic engineering.

Natural yeast species that can co-ferment hexose and pentose sugars include *C. utilis* and *K. marxianus*.<sup>22,35</sup> Both *C. utilis* and *K. marxianus* can be grown on a broad substrate spectrum and are able to utilise pentoses such as xylose, the second most abundant carbohydrate in lignocellulosic biomass.<sup>23,24</sup> *C. utilis* is strictly aerobic and does not produce ethanol, whereas strains of *K. marxianus* can produce both ethanol and yeast biomass.<sup>22,29</sup> Yeast biomass may also be obtained as a by-product from production of biofuel or lipids.<sup>10,13,36</sup> After fermentation, the harvested yeast biomass is subjected to downstream processing like washing, cell disruption and drying.

## CHEMICAL COMPOSITION OF YEAST

The chemical composition of whole cell yeast is dependent on strain, growth media and growing conditions, and downstream processing after fermentation.<sup>14</sup> Reported proximate composition of *S. cerevisiae*, *C. utilis* and *K. marxianus* yeasts is shown in Table 1. The average crude protein contents were similar for the three yeast species, but limited comparable data were found for *K. marxianus*. The reported contents of lipids and fibre in these yeast strains were generally low, but variable, possibly due to different analytical methods. Fatty acid composition may depend



**Figure 1.** Flow chart of yeast production from lignocellulosic biomass involving four major steps: (1) pre-treatment of the biomass to remove lignin and to make cellulose and hemicellulose more accessible to hydrolysis; (2) enzymatic hydrolysis to convert cellulose and hemicellulose into C6 and C5 sugars; (3) fermentation of sugars, nitrogen, phosphate, and other nutrients; and (4) downstream processing into dry yeast product for use as a protein source in fish feed.

on growth stage and culture conditions,<sup>14,37</sup> but characteristically these yeast strains may contain mainly saturated fatty acids.<sup>38</sup> The nucleic acid contents in foods depend mainly on cell density,<sup>39</sup> and contents in yeasts are much higher than in plant and animal foods. The studied yeast species showed large variation in nucleic acid content among the different studies. This could be due to differences in growth rate, as rapidly dividing cells typically result in a higher level of nucleic acids than slower growth rates,<sup>35,40</sup> or different analytical methods.<sup>41</sup> Yeast is also a source of minerals such as phosphorus, calcium, sodium, zinc, iron, copper, manganese, and selenium.<sup>42,43</sup> Furthermore, yeasts such as *S. cerevisiae* and *C. utilis* are a moderately rich source of vitamins, predominantly B-vitamins, i.e. riboflavin, pyridoxine, niacin and pantothenic acid.<sup>42,44</sup>

The amino acid profile of yeasts compared with fish meal and soybean meal is shown in Table 2. The yeasts and fishmeal revealed similar contents of most indispensable amino acids on a crude protein basis, despite higher contents of non-protein nitrogen in the form of nucleic acids in yeast, but the content of methionine was lower in yeast than in fishmeal. Yeast protein is characteristically low in the sulfur-containing amino acids methionine and cysteine.<sup>8,45</sup> In general, the amino acid profile of the yeasts compiled in Table 2 agree well with reported ranges for yeast protein.<sup>35</sup> The data indicate that *S. cerevisiae* may have higher contents of methionine compared to the other yeast species, but slightly lower content of lysine. Except for the low methionine contents, yeast protein has a favourable amino acid composition compared with the fish requirements.<sup>46</sup> In agreement with previous studies,<sup>38</sup> the yeasts thus have potential as a protein source in a mixed diet for fish.

## DIGESTIBILITY OF YEAST IN FISH

Digestibility is an important characteristic of yeast as an economically competitive protein source for aquaculture. Yeast products have a protein digestibility comparable to conventional protein sources in diets for different fish species. Studies with sea bass (*Dicentrarchus labrax*) showed that the apparent protein digestibility of *S. cerevisiae* was high (88.3%), but lower than that of high quality fishmeal (92.8%).<sup>47</sup> Increasing replacement of fishmeal with *S. cerevisiae* in the latter study also resulted in slightly decreasing apparent digestibility of dry matter and energy. Replacing fishmeal with up to 380 g kg<sup>-1</sup> dried *S. cerevisiae* yeast in

diets for pacu (*Piaractus mesopotamicus*, Holmberg 1887) had no significant effect on protein digestibility, whereas lipid digestibility was significantly increased.<sup>48</sup> In Gilthead sea bream (*Sparus aurata*), increased protein digestibility has been obtained when replacing fishmeal with 100 g kg<sup>-1</sup> and 200 g kg<sup>-1</sup> *S. cerevisiae* in the diets.<sup>49</sup> In the redclaw crayfish (*Cherax quadricarinatus*), the apparent protein digestibility of dried brewers yeast was of 92.6%, compared with 89.8% for fishmeal.<sup>50</sup> In contrast to the latter studies, rather poor protein and amino acid digestibility of intact and dried *S. cerevisiae* have been shown in salmonid fish species such as rainbow trout, *Oncorhynchus mykiss*,<sup>43,51</sup> Atlantic salmon,<sup>8</sup> and Arctic char (*Salvelinus alpinus*).<sup>52</sup> Lower energy and amino acid digestibility for intact than for extracted *S. cerevisiae* has been reported for Arctic char, while there was no significant difference in Eurasian perch (*Perca fluviatilis*).<sup>52</sup> This indicates differences in digestion efficiency among species, possibly due to different gastro-intestinal enzyme activity.<sup>53</sup>

High *in vitro* digestibility of *C. utilis* cells cultured in *Eucalyptus globulus* hydrolysates has been observed.<sup>35</sup> Results from studies with up to 347 g kg<sup>-1</sup> *C. utilis* in diets for tilapia fry indicated that the apparent protein digestibility was similar to that of fishmeal.<sup>54</sup> Protein digestibility in rainbow trout was increased when freeze-dried *C. utilis* biomass was fed at levels up to 200 g kg<sup>-1</sup>, corresponding to 35% of total dietary crude protein replacing fishmeal protein.<sup>55</sup> Very few studies have compared the digestibility of different yeasts in the same experiment. In studies with Atlantic salmon, replacement of 40% of low temperature (LT) fishmeal with *C. utilis* or *K. marxianus* yeast had no significant effect on digestibility of crude protein and energy, whereas substitution with *S. cerevisiae* significantly reduced protein and energy digestibility.<sup>8</sup> Digestibilities of individual amino acids were also unaffected by dietary inclusion of *C. utilis* and *K. marxianus*, except for a reduced digestibility of methionine and cysteine in the *K. marxianus* diet. Conversely, the diet containing *S. cerevisiae* revealed significantly lower amino acid digestibility than the fishmeal control except for glycine and cysteine.<sup>8</sup> The latter study<sup>8</sup> indicated considerable differences in digestibility among different whole cell yeast products.

## Effects of processing of yeast on digestibility

The harvested yeast biomass should be subjected to cost-efficient downstream processing to preserve valuable nutrients and bioactive components, and to promote as high digestibility as

**Table 1.** Concentration (g kg<sup>-1</sup> DM) of crude protein (N × 6.25), nucleic acids, lipids, fibre, carbohydrates, and ash of yeast

Organism	Crude protein	Nucleic acids	Lipids	Fibre	Carbohydrates	Ash	Reference
<i>Saccharomyces cerevisiae</i>	445	–	–	6.3 <sup>¶</sup>	–	64	43
	485	75 <sup>*</sup>	34 <sup>§</sup>	122 <sup>**</sup>	329 <sup>¶¶</sup>	83	61
	539	–	06 <sup>‡</sup>	23 <sup>††</sup>	50 <sup>***</sup>	68	116
	473	48 <sup>*</sup>	–	–	77 <sup>***</sup>	–	117
	475	60 <sup>†</sup>	21 <sup>‡</sup>	–	11.4 <sup>***</sup>	66	8
	456	–	80 <sup>§</sup>	–	232 <sup>¶¶</sup>	103	42
	466	–	10 <sup>§</sup>	–	–	63	64
	396	9.0 <sup>*</sup>	5	314 <sup>**</sup>	–	45	37
<i>Candida utilis</i>	501	–	–	–	–	–	55
	462	11.5 <sup>†</sup>	24 <sup>§</sup>	12 <sup>¶</sup>	–	95	44
	482	–	16 <sup>§</sup>	6.5 <sup>¶</sup>	–	–	54
	598	99 <sup>†</sup>	3.2 <sup>‡</sup>	–	40 <sup>***</sup>	58	8
<i>Kluyveromyces marxianus</i>	544	109 <sup>†</sup>	9 <sup>‡</sup>	–	9 <sup>***</sup>	81	8

\*RNA; †Total nucleic acids; ‡HCl-EE; §Ether extract (EE); ¶Crude fibre; \*\*Total fibre; ††Acid detergent fibre (ADF); §§Neutral detergent fibre (NDF); ¶¶Total carbohydrates; \*\*\*Starch.

**Table 2.** Average amino acid composition (g 16 g N<sup>-1</sup> ± standard deviation of non-hydrated amino acids) of yeast compared with fish meal and soybean meal

Amino acid	Fishmeal <sup>46</sup>	Soybean meal <sup>46</sup>	<i>Saccharomyces cerevisiae</i> <sup>8,37,43,61,64,116</sup>	SD	<i>Candida utilis</i> <sup>8,44,54,55,118</sup>	SD	<i>Kluyveromyces marxianus</i> <sup>8,119,120</sup>	SD
<i>Indispensable amino acids</i>								
Arginine	5.74	7.38	4.68	0.60	5.20	0.71	4.34	0.20
Histidine	2.36	2.67	2.47	0.87	1.97	0.28	1.80	0.18
Isoleucine	4.53	4.94	4.43	0.76	4.29	0.34	4.20	0.14
Leucine	7.06	7.80	6.73	1.16	6.19	0.61	6.81	1.37
Lysine	8.18	5.53	6.95	0.70	7.71	1.16	7.39	0.65
Methionine	2.87	1.41	1.81	0.49	1.08	0.23	1.33	0.25
Phenylalanine	3.84	5.26	4.18	0.72	3.64	0.51	3.96	0.25
Threonine	4.00	4.03	4.71	0.78	4.71	0.23	5.11	0.33
Tryptophan	1.05	1.41	1.08	0.15	1.17	0.12	0.98	
Valine	4.87	5.51	5.24	0.82	5.08	0.56	5.11	0.62
<i>Dispensable amino acids</i>								
Alanine	–	–	6.13	1.17	5.75	0.74	8.49	1.39
Aspartic acid	–	–	9.51	1.48	8.30	1.59	10.59	1.11
Cysteine	1.31	1.53	1.23	0.65	0.81	0.21	0.58	0.17
Glutamic acid	–	–	13.01	1.02	10.43	1.37	13.63	1.92
Glycine	–	–	4.32	0.77	4.15	0.27	4.47	0.45
Proline	–	–	3.69	0.71	3.85	0.83	3.99	0.66
Serine	–	–	4.48	0.94	4.07	0.74	5.34	0.34
Tyrosine	3.08	3.21	3.69	1.01	3.07	0.50	3.18	0.28

possible. There are few data on effects of processing on digestibility, and whether different yeasts require different processing methods. It has been suggested that the thick and rigid cell walls are a major problem inhibiting industrial production and utilisation of dietary yeast protein.<sup>37,56–58</sup> A poorly digestible cell wall may limit enzymatic access to cellular contents, depending on yeast cell characteristics as well as growth substrate and processing conditions. Rumsey *et al.*<sup>51</sup> showed that intact brewers yeast had lower protein and energy digestibility in rainbow trout than disrupted yeast cells, yeast extract and yeast protein isolate.

Cell walls represent 26–32% of the cell dry weight and contain varying proportions of mannan-oligosaccharides,  $\beta$ -glucan, chitin and nucleic acids depending on species and strain.<sup>59,60</sup> Various chemical, enzymatic, physical or mechanical methods can be used to rupture the yeast cell walls (reviewed by Nasser *et al.*<sup>23</sup>). Digestibility of yeast nutrients can be increased by mechanical rupturing of cell walls<sup>61</sup> or enzymatic hydrolysis.<sup>58</sup> Enzymatic pre-treatment followed by high-pressure mechanical homogenisation have been shown to be efficient for *C. utilis*.<sup>62</sup> Yeast extract with higher protein content than in whole or hydrolysed yeast



is obtained by removal of cell wall material; this represents the water-soluble cell contents. Increasing inclusion of yeast extract as a replacement for fishmeal in diets for shrimp (*Litopenaeus vannamei*) increased the apparent digestibility of protein,<sup>63</sup> most likely due to the combined effect of removal of cell walls and increased proportion of water-soluble low molecular weight proteins. In Arctic char, Langeland *et al.*<sup>52</sup> showed that the absence of intact cell walls had a positive effect on digestibility of *S. cerevisiae* protein. This confirms the finding that a phosphorylated *S. cerevisiae* protein concentrate, produced by rupturing of cell walls and centrifugation to remove debris, produced much higher protein digestibility in rats than whole yeast cells.<sup>37</sup> However, the cell wall fraction is rich in bioactive and immunostimulating compounds like  $\beta$ -glucan and mannan oligosaccharides. Hence, it seems likely that the whole yeast biomass, after rupturing of cell walls, may be the most attractive feed ingredient, by combining the properties as a source of nutrients and bioactive components. Extrusion of feed may cause partial disruption of yeast cell walls and potentially increase protein and amino acid digestibility.<sup>64</sup> Further research is warranted to determine optimal extrusion conditions for fish feed with contents of different yeast whole cell products.

## EFFECT OF DIETARY YEAST ON GROWTH PERFORMANCE, NITROGEN UTILISATION AND CARCASS COMPOSITION

Growth responses and protein utilisation in fish fed yeasts may depend on a number of factors, including yeast substrate, yeast strain, and post-fermentation processing, as well as fish species and diet formulation. As an easily available natural industrial product, strains of *S. cerevisiae* have been used as a protein source in a number of studies with different fish species. Several growth performance studies have shown that intact *S. cerevisiae* can be used to partly replace fishmeal protein in diets for fish species such as sea bass,<sup>47</sup> the omnivore pacu, *Piaractus mesopotamicus*;<sup>48</sup> Nile tilapia, *Oreochromis niloticus*;<sup>65</sup> Atlantic salmon;<sup>8</sup> Thai Panga, *Pangasianodon hypophthalmus*  $\times$  *Pangasius bocourti*;<sup>66</sup> and Arctic char.<sup>64</sup> A majority of these studies have been carried out with isonitrogenous and isoenergetic diets, with or without balancing amino acid composition with synthetic amino acids. Some studies have shown that partial substitution of fishmeal with *S. cerevisiae* increased growth rate and nitrogen retention.<sup>47,48</sup> Moderate levels of 250 g kg<sup>-1</sup> brewers yeast (*S. cerevisiae*) in diets for rainbow trout, replacing casein, improved growth performance, whereas the very high levels of 500 g kg<sup>-1</sup> and 750 g kg<sup>-1</sup> reduced feed intake and growth.<sup>67</sup> Partially replacing fishmeal with 345 g kg<sup>-1</sup> spray-dried and inactivated *S. cerevisiae* significantly reduced specific growth rate and feed conversion ratio, but did not affect retention of digested N in Atlantic salmon.<sup>8</sup> Conversely, replacing fishmeal protein with 289 g kg<sup>-1</sup> intact *S. cerevisiae* in extruded diets for Arctic char had no negative effects on growth performance.<sup>64</sup> The adverse effect of *S. cerevisiae* on growth and feed conversion ratio in the study by Øverland *et al.*<sup>8</sup> might be associated with poor digestibility of protein, amino acids and energy, possibly indicating an unfavourable yeast post-fermentation procedure.

Unlike *S. cerevisiae*, there are few studies with *C. utilis* and *K. marxianus* in diets for farmed fish. *C. utilis* is recognised for rapid growth to high cell density based on numerous inexpensive substrates including lignocellulosic biomass and wastes.<sup>18</sup> It is

commonly used as a food additive, partly due to umami flavour, high palatability, and documented safety. In studies with rainbow trout, *C. utilis* biomass grown on peat could partly replace fishmeal protein without significant reduction of growth performance.<sup>55</sup> Studies with Atlantic salmon showed that *C. utilis* grown on a lignocellulosic sugar substrate could be fed at 283 g kg<sup>-1</sup> of the diet, replacing 40% of the fishmeal protein in isonitrogenous diets, without adverse effects on growth, feed intake or feed conversion ratio.<sup>8</sup> Nitrogen retention was significantly enhanced in salmon fed *C. utilis* as compared with the fishmeal control diet, despite lowered level of methionine (the likely limiting essential amino acid for protein synthesis), higher level of non-protein nitrogen, and unaffected digestibility of nitrogen and amino acids. In previous studies with tilapia fry, up to 350 g kg<sup>-1</sup> *C. utilis* was fed without adverse effects on growth and feed conversion ratio.<sup>54</sup> Recent studies with shrimp (*Litopenaeus vannamei*) showed that growth and survival rate were similar when feeding diets containing varying proportions of *C. utilis* and fishmeal.<sup>68</sup> They showed that by applying a nitrogen stable isotope technique (<sup>15</sup>N), the relative body growth incorporation of dietary nitrogen from *C. utilis* consistently increased with increasing proportion in the diets. Previous studies have indicated that *K. marxianus* yeast is a promising protein source for aquaculture.<sup>57</sup> The results obtained by Øverland *et al.*<sup>8</sup> support this by documenting that a diet containing 302 g kg<sup>-1</sup> *K. marxianus*, replacing 40% of the crude protein from fishmeal, promoted similar growth, feed conversion ratio, and nitrogen retention in Atlantic salmon as the fishmeal-based control diet. No other study apparently has examined the suitability of inactive *K. marxianus* as a major protein source in aquaculture.

Recovery of microbial biomass as a by-product from fuel ethanol biorefineries is a new concept to produce yeast protein sources. In studies with carp (*Cyprinus carpio*), significantly improved growth and feed conversion ratio was obtained when dried yeast from distillers grains and solubles, containing 340 g kg<sup>-1</sup> protein on a DM basis, replaced 15% or 20% of a high quality fishmeal.<sup>36</sup> A higher level of this yeast (460 g kg<sup>-1</sup>, replacing 50% of fishmeal protein) resulted in similar growth and feed conversion as the fishmeal control diet. Grain distillers dried yeast containing 520 g kg<sup>-1</sup> crude protein in the dry matter has been evaluated as a protein source for rainbow trout,<sup>69</sup> and the results showed no significant differences in growth performance when 25% and 37.5% of the fishmeal were replaced with this yeast product on a digestible protein basis. Further fishmeal replacement resulted in linear reductions in growth and poorer feed conversion ratio, although the diets were balanced for digestible lysine, methionine and threonine. The reduction in growth performance at high inclusion levels was not associated with reduced feed intake.<sup>69</sup>

In addition to amino acids and protein, yeast cells synthesise nucleic acids as substantial N-containing components. In fast-growing yeast cells, about 10–15% of the total nitrogen is in the form of nucleic acids.<sup>40</sup> In contrast to indispensable amino acids, nucleotides are endogenously synthesised and not considered essential nutrients, but dietary nucleic acids may be partially salvaged and used by the animals, thus influencing growth performance and nitrogen balance.<sup>70</sup> Rumsey *et al.*<sup>71</sup> showed that feeding rainbow trout high levels of nucleic acids, corresponding to up to 500 g kg<sup>-1</sup> dietary *S. cerevisiae*, had no effect on feed intake, but increased growth and nitrogen retention. This may be consistent with the results obtained by Øverland *et al.*<sup>8</sup> where nitrogen retention was increased by feeding *C. utilis* containing 93 g kg<sup>-1</sup>

nucleic acids as a substitute for fishmeal for Atlantic salmon, thus reducing the proportion of amino acid nitrogen in total dietary nitrogen. Similarly, replacing fishmeal with *K. marxianus*, containing 102 g kg<sup>-1</sup> nucleic acids, in salmon diets had no negative effect on nitrogen retention.<sup>8</sup> This indicates that the nucleic acids in yeast may be directly incorporated in the body or spare non-essential amino acid nitrogen through endogenous utilisation. In contrast, no clear nitrogen sparing effect of yeast or dietary nucleic acids have been found in turbot (*Psetta maxima*) or rainbow trout.<sup>72</sup> The digestibility of nucleic acid nitrogen seems to be high in many species,<sup>41</sup> and a variable proportion is excreted in the urine.<sup>73</sup> The urolytic pathway in Atlantic salmon seems to be able to control high dietary levels of nucleic acids.<sup>74</sup> This may partially explain the positive effect of dietary inactive yeast on growth and nitrogen utilisation in studies with different fish species.

### Carcass composition

Fillet characteristics of fish fed yeast products may depend on fish growth and body condition.<sup>54,75</sup> However, studies with different fish species have shown no or only minor effects of replacing fishmeal by different yeast products on whole body or muscle proximate composition.<sup>36,65,66,69</sup> Carcass composition was not significantly affected by protein source when *C. utilis* replaced fishmeal in diets for rainbow trout,<sup>55</sup> or fishmeal and blood meal in diets for fingerling grey mullet (*Mugil cephalus*).<sup>76</sup> Nor were there any clear effects on carcass composition when using high levels of *C. utilis* in diets for tilapia (*Oreochromis mossambicus* Peters).<sup>54</sup>

The fatty acid composition of fish muscle reflects the dietary fatty acid profile. The fatty acid profile of fish muscle may thus be slightly modified by using yeast as a lipid source. However, use of diets with a low proportion of dietary lipids from yeast products may have no or minor effects on fillet fatty acid profile. This has been clearly shown in studies with live *K. marxianus* cells as a substitute for fishmeal in diets for tilapia (*Oreochromis niloticus* Linnaeus).<sup>77</sup> Likewise, fillet colour measurements in Thai Panga have revealed no effects of dietary substitution of fishmeal with *S. cerevisiae*.<sup>66</sup> In general, the knowledge from using yeast biomass in nutritionally balanced fish diets indicates no major effects on characteristics of fish as human food. Assuming use of GRAS yeast strains, the evaluation as fish feed ingredients should be mainly based on effects on fish growth performance and health.

## EFFECT OF DIETARY YEAST AND YEAST PRODUCTS ON FISH HEALTH

Yeast is commonly used in aquaculture as a growth promoter and immunostimulant in functional feeds. Yeasts contain various bioactive components such as  $\beta$ -glucans, mannose polymers covalently linked to proteins (manno proteins), minor amounts of chitin, and nucleic acids. Positive health effects have been documented when adding yeast to fish diets at low levels (10–40 g kg<sup>-1</sup>), especially with whole *S. cerevisiae* yeast or yeast products, but few studies evaluate the health beneficial effects of yeast when used as a protein source. Low levels of *S. cerevisiae* yeast have shown to enhance growth performance, immune responses, and/or protection against bacterial infection, and to increase disease resistance in several fish species, such as salmonids,<sup>58,78</sup> hybrid striped bass,<sup>79,80</sup> gilthead seabream,<sup>81</sup> seabream,<sup>82</sup> hybrid tilapia (*Oreochromis niloticus*),<sup>83</sup> common carp,<sup>60</sup> Indian common carp (*Labeo rohita*),<sup>84</sup> channel catfish (*Ictalurus punctatus*),<sup>85</sup> Japanese seabass (*Lateolabrax japonicus*),<sup>86</sup>

and Nile tilapia.<sup>87</sup> Yeasts such as *C. utilis* have also been shown to modulate immune responses in rainbow trout,<sup>88,89</sup> and *K. marxianus* has shown probiotic properties in an *in vitro* colonic model system.<sup>90</sup> Feeding live bakers yeast (10 g kg<sup>-1</sup>) improved growth performance and feed utilisation, and resistance to *Aeromonas hydrophila* in Nile tilapia<sup>91</sup> and enhanced intestinal colonisation or accelerated the maturation of the digestive system of early feeding fry.<sup>92</sup> When used in moderate levels of 200 g kg<sup>-1</sup> in diets for Atlantic salmon, *C. utilis* and *K. marxianus* affected the bacterial composition in the distal intestine, particularly by reducing the relative amount of *Firmicutes* bacteria.<sup>93</sup> Inclusion of 200 g kg<sup>-1</sup> inactive *C. utilis* whole yeast cells, and also *K. marxianus*, counter-acted soybean meal induced enteritis (SBMIE) in the distal intestine of Atlantic salmon,<sup>93</sup> while *S. cerevisiae* did not protect against SBMIE. Adding 10–20 g kg<sup>-1</sup> dietary yeast cell wall extracted from *S. cerevisiae* improved growth performance and intestinal mucus development of Japanese seabass fed a diet with 210 g kg<sup>-1</sup> SBM.<sup>86</sup> Similarly, adding 2 g kg<sup>-1</sup> bakers yeast cell wall fraction rich in MOS to a diet with moderate (140 g kg<sup>-1</sup>) inclusion of SBM eliminated SBMIE in the distal intestine of Atlantic salmon.<sup>78</sup> The protective mechanisms of yeast against SBMIE could include a beneficial effect on gastrointestinal morphology, reduced inflammatory responses, and stimulating wound healing processes and modulation of gastrointestinal microbiota.

Enhanced growth performance and reduced mortality after pathogen challenge and increased disease resistance are attributed to several immunostimulatory properties of yeast, including stimulation of both humoral and cellular immune responses. These responses include blood neutrophil oxidative radical and extracellular superoxide anion production of head kidney macrophages,<sup>79</sup> enhanced serum leucocytes and lysozyme activities and complement activity,<sup>58,83</sup> enhanced cellular immune responses of the innate immune system as measured by phagocytic index, respiratory burst activity,<sup>83,84</sup> number of erythrocytes and lymphocytes,<sup>84</sup> and increased antioxidant status by increasing catalase, superoxide dismutase, and total antioxidant activity in serum.<sup>94</sup> Enhanced growth performance and health in fish fed yeast might also be attributed to improved digestive enzyme activity,<sup>95</sup> and by supplying digestive enzymes that aid in the digestion of complex carbohydrates,<sup>84</sup> or by enhancing digestive capacity by improving gastrointestinal morphology and thereby increasing the absorptive surface.<sup>8,58,95,96</sup> Yeast also provides dietary nucleotides that enhance immune responses in several fish species, including Atlantic salmon,<sup>97</sup> hybrid striped bass,<sup>80</sup> and grouper (*Epinephelus malabaricus*),<sup>98</sup> enhances intestinal epithelial cells in Atlantic salmon,<sup>97</sup> and increases weight of the gastrointestinal tract in Atlantic salmon.<sup>8</sup> Modulation of intestinal microbiota may also be a contributing factor as reported for several fish species, including salmonids,<sup>58</sup> common carp,<sup>60</sup> hybrid striped bass,<sup>80,99</sup> hybrid tilapia,<sup>83</sup> juvenile beluga sturgeon (*Huso huso*),<sup>100</sup> and Nile tilapia.<sup>87</sup> Modulation of intestinal microbiota and protection against bacterial infection might be associated with the mannan oligosaccharides in the yeast cell walls, which prevent colonisation of pathogens and thereby eliminating them from the intestinal tract, or by increasing the gastrointestinal tract mucus secretion and by that improving the gastrointestinal barrier function and protection.<sup>101</sup>

The effect of yeast on immunity and survival rate of fish is inconsistent, however. This could be attributed to factors such as type of yeast products,<sup>93</sup> concentration,<sup>83,86</sup> feeding duration of the yeast products,<sup>85</sup> and fish age, size and species. The fermentation conditions or downstream processing conditions during manufacture

may also affect the nucleic acid levels<sup>71</sup> and the cell wall properties of the yeast and thus the health promoting effects. Fish fed diets supplemented with nucleotides have shown enhanced resistance to viral, bacterial and parasitic infection, thus indicating that nucleotides are conditionally or semi-essential nutrients for fish.<sup>99</sup> Grinding yeast cells or extracting  $\beta$ -(1,3)-glucan from yeast also have implications for the health promoting properties.<sup>60,102</sup>

## CONCLUSIONS AND FUTURE PERSPECTIVES

The human world population is estimated to increase from 6.9 billion in 2010 to 8.2 billion by 2030, and probably to more than 9 billion by 2050.<sup>1</sup> The food supply will have to increase and become more efficient to grow given the availability of natural resources and existing technology.<sup>103</sup> The total supply of seafood has been growing more rapidly than the global population.<sup>1</sup> Almost 90% of the capture fisheries are now fully exploited or over-exploited. Aquaculture has grown to almost half of total fish supply, and is expected to continue to grow at a rapid rate in the coming decades. The rapid growth of aquaculture production has been driven by productivity growth, technological progress, globalising trade, and favourable economics of large-scale intensive farming.<sup>104</sup>

Increasing production of carnivorous species like salmonids has greatly enhanced the demand of commercial feed, and further expansion will depend on a sufficient supply of high-quality sustainable feed resources. The increased production has been made possible through diet modifications, mainly replacing fishmeal and oil with plant ingredients derived from agricultural commodity crops like soybean and rapeseed. Conceivably, future salmon farming may be a net producer of fish protein and oil.<sup>5,105–107</sup> Challenges associated with a further increase in plant protein sources include contents of anti-nutrients,<sup>108–110</sup> potential contents of mycotoxins,<sup>111</sup> and insecticides.<sup>112</sup> The increasing use of genetically modified plants<sup>113</sup> is subject to regulatory constraints and may limit acceptability in some markets and among groups of consumers. The sustainability of substituting marine feed ingredients with plant products in fish feed is questionable, considering the need of water and arable land, use of scarce global phosphorus fertilisers, use of pesticides, and environmental human health implications.<sup>114</sup> Although these concerns are not specific to plant products used in fish feed, it is urgent to identify non-conventional sources of protein that can be converted into high-quality feed ingredients for aquaculture. Especially the further expansion in the production of carnivorous species like salmon may require economically competitive and sustainable feed ingredients produced from novel non-food resources.

Efforts towards sustainability in the aquaculture industry may benefit from combining the approaches of different research disciplines. Fast-growing yeasts, produced from non-edible feedstocks and categorised as GRAS microorganisms, may be attractive candidates for industrial upscaling to important feed resources. A number of studies with different yeast products and fish species have shown that yeast products produced from lignocellulosic substrates can be used as attractive protein sources with the additional benefit of improving immune function and intestinal health of the fish. Yeast production from lignocellulosic biomass offers the advantage of making food production less dependent on arable land and climate, and relieving pressure on resources for direct human food production. No single cost-efficient technology may meet all the challenges of large-scale industrialisation,<sup>20,115</sup> and yeast production from lignocellulosic biomass is less efficient

compared to feedstocks such as sugar or starch. Currently, the process cost of yeast protein production from lignocellulosic biomass may be too high and there is a need to develop efficient processes for economic utilisation. Production of multiple co-products such as lignin, biochemicals and biofuel in biorefineries can contribute to reduce the cost as the technology matures. In the future, this may open diverse options that reach beyond current use of land areas for production of food and feed. Continued research and development in yeast production from lignocellulosic substrates can be an important contribution to securing the sustainability and economic viability of future aquaculture.

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