**Review** article

## Germination: a means to improve the functionality of oat

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The biochemical and physiological reactions of germination have long been utilised to produce barley malt for brewing and other purposes. Also some oat malt has been produced as raw-material of ale and stout production. The main goals of malting have been the degradation of grain storage components to soften the kernel structure, synthesis of amylolytic enzymes and production of nutrients for brewing yeast. Also flavour and colour attributes have been important. During the recent years interest has arisen also in the secondary metabolites produced during germination, which can have valuable health promoting properties and act as bioactive or functional compounds in foods. By using a tailored germination/malting process a desired combination of valuable properties may be obtained in germinating grains or seeds. All this requires knowledge and know-how of the germination process and the biochemistry behind it. This paper reviews the scientific knowledge about germination/malting of oat with special emphasis on changes in grain characteristics.

Key words: oat, germination, processing, bioactivity

#### Introduction

Germination of cereals has been used for centuries to soften kernel structure, to increase nutrient content and availability, to decrease the content of antinutritive compounds, and to add new flavours without knowing the biochemistry behind these phenomena. Barley malting is the most widely known controlled germination process, used to produce malt for brewing purposes and food applications. Germination of a grain or seed is a chain of events that commences when viable, dry seeds imbide water, and terminates with the elongation of the embryonic axis. Upon imbibition, the quiescent seed rapidly resumes metabolic activity including respiration, enzyme and organelle activity, and RNA and protein synthesis (Bewley and Black 1994). Enzymes are synthesized to degrade storage macromolecules. These reactions lead to structural modification and development of new compounds, many of which have high bioactivity and can increase the nutritional

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value and stability of the grains. Furthermore, many of the developed compounds are flavour precursors participating in the formation of palatable malt flavour.

A traditional malting process can be divided in three main steps: steeping, germination and kilning (drying). During steeping the moisture content of the kernel is increased, normally to 43–45% to initiate germination. During germination (typically 4–7 days) enzyme synthesis and kernel modification occur under strictly controlled conditions (temperature, moisture, aeration). The aims of the kilning step (approximately 20– 24 h) are to dry the kernel to a moisture content of 3–5% to ensure microbiological stability of the product, to stop or retard the biochemical reactions, and to produce aroma and flavour compounds (Kunze 1999).

Oat malt is a minor product when compared to barley malt, but it is continuously produced e.g. for ale and stout brewing and for special food ingredients. Native oat is known as a health-promoting cereal with high content of soluble dietary fiber, beneficial fatty acid content and a large selection of vitamins, minerals, sterols and antioxidants. The studies of germinated/malted oat are few but cover a wide range of topics: influence of processing on the main enzyme activities, especially amylolytic, proteolytic and lipolytic activities, modification of amino acid content, fat content and composition, fiber degradation, the amount of bioactive and antinutritive compounds, and flavour formation. These topics will be discussed in more detail in the following chapters.

## Modification of oat macromolecules and the development of related enzyme activities during germination

New groups of healthy and functional foods or food ingredients have been developed during

recent years including foods with slowly digestible starch, functional oligosaccharides and bioactive peptides (Crittenden and Playne 1996, Hilliam 2000, Quinquenet 2000, Peptide enriched sport drinks for enhanced recovery 2001, Company listing by category 2003). Grain storage compounds, or their degradation products could also serve as raw-materials for these novel foods or ingredients. Especially germination is a process where modified compounds and enzymes that degrade storage macromolecule are naturally formed. With the basic knowledge of hydrolysis of macromolecules, controlled utilization of amylolytic, glucanolytic and proteolytic activities of germinating grains, a range of cereal based, novel food ingredients could be achievable.

#### Starch and other polysaccharides

Starch is the main storage compound in oat grains constituting 60% of the dry weight (Wood et al. 1991). The starch molecules amylopectin and amylose are arranged in compound granules within the endosperm cells (Bechtel and Pomeranz 1981). During germination, starch degradation is very limited, despite the fact that starch degrading enzymes are synthesized. According to Manners and Bathgate (1969) malting of oat caused a very limited decrease in the molecular size of amylose and a small, but significant reduction in the exterior chain length of amylopectin. Sutcliffe and Baset (1973) demonstrated a close correlation between the rate of increase in amylase activity and the rate of starch degradation. The comparison of germinated oat starch to other cereal starches showed that oat starch granules were most resistant to  $\alpha$ -amylolysis with little increase in damaged starch and no erosion of granule surface (Lineback and Ponpipom 1977). However, release of small granules from the compound starch granules was observed. Salmenkallio-Marttila et al. (2004) found in their detailed microstructure studies that after 3 days of germination the starch granules remained intact but the protein network surrounding the

starch granules was more dense and tightly bound than in native grains.

Hughes et al. (1997) malted different cereals by using a typical pilsner malt type malting programme and found that  $\alpha$ - and  $\beta$ -amylase activities were much lower in oat than in barley, wheat or rye. Peterson (1998) reported that  $\alpha$ -amylase activity reached levels almost as high as in some barley samples but the diastatic power, characterizing the combined effects of  $\beta$ -amylase and  $\alpha$ -amylase, was much lower than in barley. Similar results were obtained already much earlier in the studies of Pomeranz and Shands (1974) who found that the levels of  $\alpha$ -amylase in germinated oat could be increased to the same level as in barley by addition of gibberellic acid during germination. The diastatic power, however, was lower in germinated oat than in germinated barley regardless of gibberellic acid treatment.

The main cell wall component (2-7% of the total dry weight) in the oat endosperm is a mixed linked  $(1\rightarrow 3), (1\rightarrow 4)$ -  $\beta$ -D-glucan. Oat  $\beta$ -glucan has a higher molecular weight than barley or rye  $\beta$ -glucan (Beer et al. 1997). Nutritionally,  $\beta$ -glucan is a dietary fiber not degraded by digestive enzymes. β-Glucan is easily extractable, forming viscous solutions and is known to decrease the levels of serum cholesterol and balance the glucose and insulin increase after meals (Ripsin et al. 1992, Welch 1995, Kahlon and Chow 1997). According to Hughes et al. (1997)  $\beta$ -glucanase activity of oat was comparable to that of wheat but much lower than that of barley and rye. Despite this considerably low  $\beta$ -glucanase activity, the  $\beta$ -glucan content decreased significantly during malting from 3.5% to 0.6%. Peterson (1998) also found that  $\beta$ -glucans were almost totally degraded during a 6-day germination. However, the content and properties of  $\beta$ glucans are important also in malted oat products if the cholesterol lowering and other functional effects of fiber are to be maintained. Especially, the ability of oat products to develop highly viscous matrices is known to depend on the molecular weight and viscosity of  $\beta$ -glucans, and furthermore, this ability is known to be influenced by processing (Beer et al. 1997, Zhang et al. 1998, Gallaher et al. 1999). Therefore, Wilhelmson et al. (2001) developed a short malting process for oat aiming to retain high  $\beta$ -glucan content also in malted oat. Up to 70% of the  $\beta$ -glucans could be restored, and the average molecular weights ( $M_w$  and  $M_n$ ) of  $\beta$ -glucan molecules changed only slightly during malting. The major changes in  $\beta$ -glucan reserves during this short malting were observed in the crushed cell layer beneath the scutellum (Salmenkallio-Marttila et al. 2004).

## Amino acid composition and protein degrading enzymes

Oat contains 15-20% proteins (Robbins et al. 1971, McMullen 1991). The nutritional value of oat proteins is higher than that of other cereal proteins due to lower content of prolamine protein (Peterson 1976, Hamad and Fields 1979). The total protein content of oat increases slightly during germination due to more intensive degradation of other kernel components for respiration (Dalby and Tsai 1976). Dalby and Tsai (1976) and Wu (1983) observed an increase in albumin content (rich in essential amino acids, lysine and tryptophane) during oat germination, and a subsequent decrease in globulin and prolamine contents (poor in lysine) leading to increased nutritional value of germinated oat. Hamad and Fields (1979) also found an increase in lysine content during germination, but the calculations did not show any significant increase in the nutritional value of oat when compared to casein. Shutov and Vaintraub (1987) have proposed that globulins undergo changes in their solubility at the beginning of germination, making them more susceptible for seed endoproteinases. These endoproteinases are also synthesized at the beginning of germination.

Sutcliffe and Baset (1973) showed an increase in casein hydrolysing activity in oat during germination. The rate of activity increased during the first two days of germination. The results of Mikola and Jones (2000a) showed that a wide range of oat endoproteinases were able

to hydrolyse the insoluble oat storage proteins. The serine and metalloproteinases hydrolysed oat proteins in dissolved form, and when oat proteins were in a bound form cysteine endoproteinase was the main degrading activity. Mikola and Jones (2000b) and Mikola et al. (2001) studied the hydrolysis on oat globulins and oat avenins in detail. In the beginning of germination, oat globulins were hydrolysed into intermediate molecular weight peptides. These peptides did not accumulate in the grain but they disappeared at the same rate as intact globulins (Mikola and Jones 2000b). Cysteine proteinases are assumed to be responsible for the initiation of oat globulin hydrolysis. The degradation of oat prolamins, avenins, started in the beginning of germination with hydrolysis of  $\alpha$ -avenins followed with hydrolysis of  $\beta$ -avenins after an 8-day germination (Mikola et al. 2001). Additionally, cysteine proteinases are regarded responsible for the initiation of avenin hydrolysis.

#### Lipids and lipid degradation

Oat contains more lipids than other cereals, normally 5-9% (Sahasrabudhe 1979, Youngs 1986, Saastamoinen et al. 1989). In oat, lipids are located throughout the kernel, while lipids in other cereals are concentrated in the embryo (Lockhart and Hurt 1986). Most of the oat fatty acids are unsaturated. The oleic acid portion of oat lipids is exceptionally high among cereal lipids (Youngs 1986, Saastamoinen et al. 1989). Furthermore, oat is exceptional among cereals in that it contains a significant lipid hydrolysing (lipase) activity already in native, quiescent grains (Urquhart et al. 1983, 1984, Liukkonen et al. 1993, Ekstrand et al. 1993). Still, lipids of dry, intact oat grains can stay stable for years when stored at proper conditions (Welch 1977). On the other hand, Ekstrand et al. (1993) reported a marked increase in free fatty acids during storage of whole grains already after 16 weeks at 30°C indicating the susceptibility of oat lipids to hydrolysis. Lehtinen et al. 2002 observed that routinely applied heat treatment used to inactivate oat lipase was insufficient to increase the storage stability of oat products and instead, could even accelerate the oxidative deterioration of oat lipids. All together, even though native oat contains less oxidative lipoxygenase enzyme than many other grains, like barley, wheat or maize (von Ceumern and Hartfield 1984), the stability of oat lipids and the preventation of off-flavour formation are great challenges to the oat processing industry due to the high lipid content, high native lipolytic activity of oat and, according to Lehtinen et al. (2002) the susceptability of some oat lipid fractions to normally used heat treatments.

Urquhart et al. (1983) did not observe any marked increase in oat lipase activity during a 4-day germination. However, the location of lipase partly changed during germination from the bran-endosperm fraction to the embryo. Ekstrand et al. (1992) noticed an increase in lipase activity at neutral and alkaline pH while the activity at acidic pH remained relatively constant. They suggested that the lipase activity in alkaline pH was related to the metabolic processes of the embryo during germination. Peterson (1998) studied the degradation of lipids during malting in 39 hull-less oat varieties (46 samples) and in 10 hulled varieties. The hull seemed to have influence on lipid reactions during germination. The total lipid content of hull-less varieties was higher than in hulled varieties, 5.7-10.2% and 4.4–9.6%, respectively. The proportion of free fatty acids in native, hull-less samples was higher than in hulled samples. During germination, the decrease in the total lipid content of hull-less samples was 15% on average and in hulled samples approximately 5%. In fact, in half of the hulled samples, the total lipid content increased during malting. On the other hand, the formation of free fatty acids during germination was more intense in hulled samples than in hull-less ones indicating the occurrence of both formation and degradation of lipids during germination.

Peterson (1999) studied further the relationship between lipase activity and the lipid metabolism during germination. He found that the

concentration of free fatty acids increased remarkably during a 6-day malting, but the total lipase activity decreased 40%. He concluded that there was no correlation between total lipase activity and the degradation of triglycerides and suggested the occurrence of oil body bound lipase which is responsible for the degradation of storage lipids.

Outinen (1999) studied the influence of malting conditions on the lipid content and lipolytic activity on one hulled (originally high in lipolytic activity) and on one hull-less (originally moderate lipolytic activity) variety, and reported quite different results from those of Peterson (1998). In general, no lipid degradation or significant formation of free fatty acids was seen during germination. On the contrary, in most of the malted samples the lipid content increased. Lipolytic activity of hulled oat with high original activity remained stable or decreased a little in all used malting conditions. On the other hand, in the hull-less sample with moderate original lipolytic activity this activity was further increased under some malting conditions. No increase in lipid oxidising activity during malting of oat was observed in studies of Outinen (1999). It can be concluded that the research results on lipolysis in oat germination are to some extent contradictionary and further research is needed in this area.

# Bioactive compounds of germinated oat

In addition to macrocomponents of which especially  $\beta$ -glucan has interesting positive health effects, oat contains microcomponents of which many are also physiologically active. Phytates are Ca-, K- and Mg-salts of myo-inositolhexaphosphate, which serves as a phosphate reserve in the kernel (Hall and Hodges 1966, Fredlund et al. 1997, greiner and Alminger 1999). Furthermore, oat contains various interesting bioactive compounds that are products of secondary metabolism. These compounds include vitamins, sterols and phenolic compounds. They can affect the nutritional and functional properties and stability of oat products. By germination, the concentration and availability of these compounds can be adjusted.

#### Phytate content and phytase activity

Phytates are negatively charged molecules and can form complexes or otherwise bind metal ions and proteins, and thereby influence the absorption of trace elements and activity of proteins in the gastrointestinal tract (Zhou and Erdman 1995). Phytic acid can also prevent absorption of calcium (Sandberg et al. 1993, Saha et al. 1994). Even though oat has a high mineral content, the absorption of iron and zinc from a breakfast containing oat porridge is low (Sandström et al. 1987, Rossander-Hultén et al. 1990). In a multifunctional diet, which contains high amounts of mineral substances, the influence of phytate is insignificant. However, in diets rich in cereal based foods, the influence of phytate on mineral balance can be remarkable. This can be the case with vegetarians and people of the third world (Larsson et al. 1996). On the other hand, phytic acid has also been reported to have health promoting effects. Phytic acid and its degradation products are believed to contribute to prevention of colon cancer and cardiovascular diseases (Graf and Eaton 1993, Shamsuddin 1995).

Oat phytate is known to be difficult to degrade (Larsson and Sandberg 1992, Marklinder et al. 1995). Fretzdorff (1992) noticed that the degradation of phytate during germination was significantly lower in oat than in other cereals. On the other hand, Marklinder et al. (1995) reported that malted barley flour and malted oat flour had the same capacity to degrade oat phytate. The phytate content could be further decreased when malted oat or malted oat flour was incubated in water, indicating an increase in phytase activity during germination (Fretz-

dorff 1992, Larsson and Sandberg 1992). The optimal condition for incubation was 17 h at 37–40°C (Larsson and Sandberg 1992, 1995). The results from germination studies were reflected in the clinical data: Larsson et al. (1996) reported that the use of malted oat in preparation of breakfast porridge significantly increased the uptake of iron and zinc in human subjects. According to Oksman-Caldentey et al. (2001) the phytate content could be decreased 15–35% during even a short 3 day germination process.

Rather recently oat phytase was isolated and purified for the first time (Greiner and Larsson-Alminger 1999). The enzyme is a 67 kDA monomeric protein with pH optimum at 5.0 and it is strongly inhibited by phosphatate resulting from phytate degradation. Greiner and Larsson-Alminger (1999) suggested that inhibition will not occur in a growing plant since the seedling can probably remove the released phosphate actively or passively.

#### **Sterols**

Sterols are minor lipids with significant biological functions. White and Armstrong (1986) reported an improved stability of soybean oil after addition of purified oat sterols. In plants sterols are present as free sterol alcohols, steryl esters and steryl glycosides (Dyas and Goad 1993). The major sterol compound in oat is  $\beta$ -sitosterol (Lásztity 1998, Määttä et al. 1999). Oksman-Caldentey et al. (2001) measured up to 20% increase in sterols during germination of oat. Sitostanol was found to be synthesized during germination. Oat sterols were shown to be heat-stable when germinated oat was dried at different temperatures.

#### Vitamins

Oat contains mainly lipid soluble tocols (vitamin E) and water soluble vitamin B. The main tocols in oat are  $\alpha$ -tocopherols and  $\alpha$ -tocotrienols (Hammond 1983). Oat and oat products contain high amounts of thiamin and pantothenic acid (Lockhart and Hurt 1986) and are especially rich in biotin compared to other cereals (Lásztity 1998). Germination has been used to produce a vitamin-rich oat drink (DE 3741991). Oksman-Caldentey et al. (2001) reported a 20% increase in the content of several oat B vitamins like biotin, niacin, pantothenic acid and B6 vitamin during germination.

#### Phenolic compounds

Phenolic compounds act as a part of the defence mechanisms of plants, protecting plants against pathogens, pests and other stress conditions (drought, UV-light, etc). In general, they have antimicrobial, antioxidative and anti-carcinogenic properties (Pratt 1992). Oat contains quinones of benzoic acid cinnamic acids, flavones, flavonols, flavanones, anthocyanidins, aminophenols and precursors of these compounds (Collins 1986). Totally native oat flour contains 87 mg/kg fenolic acids, the main component being ferulic acid (Sosulski et al. 1982). Oksman-Caldentey et al. (2001) reported that the total phenolic content of oat increased 3- to 4-fold during a short germination process. This could also reflect to the better extractability of phenolic compounds from kernel structures after germination. After germination most of the phenolics were bound (58%), in ester form (25%), or in glucosidic form (15%). Only 2-3% of phenolic acids were free. The major free acids were caffeic acid, syringic acid, ferulic acid and sinapic acid.

Avenanthramides are special oat phenolics, which are regarded as central compounds of the defence mechanism (Collins 1989, Dimberg et al. 1993). The avenantramides occur as constitutive components of oat grains. In oat leaves, avenanthramides are not detected until the leaves are inoculated with an incompatible race of the oat crown rust pathogen, *Puccinia coronata* f. sp. *avenae* (Miyagawa et al. 1995) or other fungal elicitor, such as victorin (Miyagawa et al. 1996). Germination increased the amount of avenanthramides significantly (Kaukovirta-Norja et

al. 2001, Oksman-Caldentey et al. 2001). An especially high increase in avenanthramide content was observed in a hull-less variety (up to 35-fold increase) indicating the special role of avenanthramides in plant protection. Bryngelsson et al. (2003) also observed a significant increase in concentration of avenantramides during steeping and germination of oat.

Phenolic compounds are known to be bitter tasting and therefore may not be advantageous for cereal products. However, different phenolics have different flavour characteristics. According to Dimberg et al. (1996) and Molteberg et al. (1996) avenanthramides and caffeic acid were significantly correlated with low rancidity, a lack of bitterness, and freshness, whereas the opposite was found for most of the simple phenolic compounds.

Phytoestrogens are one interesting group of plant phenolic compounds with reported antitumoral (Adlercreutz et al. 1995) and antioxidative activities (Fleury et al. 1992). However, germination has not been shown to influence oat phytoestrogens, e.g. lignan content or composition (Oksman-Caldentey et al. 2001).

#### Total antioxidant activity

Special antioxidative properties of oat have been known for a long time. The antioxidants are concentrated mainly in the outer layers of the oat kernel (Peterson 2001). Phenolic compounds and tocols are mainly regarded responsible for oat antioxidant activity (Hammond 1983, Forssell et al. 1990). Oat sterols have also been reported to have antioxidative properties (White and Armstrong 1986, Kahlon 1989). Forssell et al. (1992) reported that oat lecithin is a better antioxidant than soy or turnip rape lecithin. Emmons et al. (1999) found that the total phenolic content of oat was significantly correlated with antioxidant activity, measured by  $\beta$ -carotene bleaching. Also other methods measuring total antioxidant activity gave similar results.

The total antioxidativity of oat, measured as DPPH (diphenylpicrylhydratcyl) radical scav-

enging activity, increased during germination (Kaukovirta-Norja et al. 2001, Oksman-Caldentey et al. 2001). The total phenolic content of germinated oat correlated well with total antioxidativity, indicating that a significant part of the increased antioxidativity was due to phenolic compounds. The antioxidativity was significantly higher in the hull-less oat variety than in the hulled variety (Oksman-Caldentey et al. 2001).

# Influence of germination on oat flavour

Oat is perceived as a tasty cereal with palatable, nutty-like flavour (Heydanek and McGorrin 1986). Oat flavour is mainly formed during processing. The germination process is well known to intensify both colour and flavour of grain products. The kilning (drying) step of a germination/malting process is especially important in the formation of different kind of flavours in germinated grains. The Maillard reaction is one of the major reactions influencing the flavour and colour of a germinated, dried products (Fayle and Gerrard 2002). For germinated oat, the most salient sensory attributes were roasted odour and flavour, sweet taste, intense odour, intense aftertaste and hard, crisp, brittle texture (Heiniö et al. 2001). Relatively high drying temperatures (>  $85^{\circ}$ C) were necessary to produce these sensory attributes. Sensory and instrumental profile analyses of selected volatile compounds showed that roasted, sweet and nutty attributes were related to dimethyl sulphides and isobutanol. Furthermore, phenolic compounds seemed to influence oat flavour significantly. Although free fatty acids are generally supposed to affect the sensory quality, they had a negligible effect in the study of Heiniö et al. (2001).

## Microbiological quality and shelflife of germinated oat

When oat germination or malting are used to produce novel food ingredients, a controlled process and the stability and shelf-life of the products are essential. Microbial aspects have to be taken into account in wet grain processing.

#### Microbiological quality of germinated oat

The microbial flora characteristic of oat and oat products develops in the field before harvest, under storage, and during oat processing. Moulds, normally divided into field and storage fungi, are typical spoilage microbes of cereals. Especially Aspergillus and Penicillium storage fungi are known to be responsible for the deterioration of dry oat products like flakes and muesli (Weidenbörner and Kunz 1994, 1995, Nogueira and Cavalcanti 1996, Nogueira et al. 1996). Toxin forming Fusarium species are typical field fungi of oat products. According to Hietaniemi and Kumpulainen (1991, 1993), oat seemed to be more susceptible to Fusarium mycotoxins than rye, wheat or barley. However, Fusarium toxin formation is greatly dependent on the Fusarium species and climate conditions (Gareis et al. 1989).

Germination conditions favour microbial growth. Changes in the composition of microflora during barley malting and the effects of microbes on malt and beer quality have been studied extensively (Petters et al. 1988, Flannigan 1996, Noots et al. 1998). It is well known that microbes in and on the outer layers of kernels greatly influence the quality of the final product. Aerobic microbes compete with grain tissue for dissolved oxygen during the steeping of the kernels, and uncontrolled multiplication of microbes may lead to poor germination (Doran and Briggs 1993). Wilhelmson et al. (2001) studied the changes in microflora during germination of hulled and hull-less oat varieties at dif-

ferent germination temperatures. Results showed clearly that elevated germination temperatures led to an increase in Fusarium, aerobic heterotrophic bacteria, Pseudomonas spp., lactic acid bacteria, and aerobic spore-forming bacteria. The hull-less variety was more susceptible to Fusarium contamination at high temperatures. However, at moderate 15°C temperature only one oat sample contained low amounts of the Fusarium toxin deoxynivalenol indicating the importance of germination temperature for control of microflora and toxin formation. Bacillus cereus or Clostridium species were not detected in any of the samples. This was beneficial because heatresistant spores of Bacillus species cause spoilage of bread by rope formation and may also constitute a health risk (Kramer and Gilbert 1989, Thomson et al. 1993).

Oksman-Caldentey et al. (2001) used lactic acid starter cultures to control microflora during germination. The use of starter cultures significantly decreased the total bacterial content and content of *Pseudomonas* spp. of malted oat. Furthermore, starter cultures inhibited the growth of *Fusarium* spp. during oat germination.

#### Stability of oat lipids after germination

The germination process itself does not, in general, deteriorate oat lipids. However, during germination the amount of reducing sugars increases, resulting in perceived sweetness, and the amount of free fatty acids may rise, increasing the risk of rancidity (Peterson 1998). Heiniö et al. (2002) studied the influence of germination on lipid and flavour stability of oat products (oat groats). A possible relationship between a decrease in lipolytic activity during oat germination (Outinen 1999, Heiniö et al. 2001) on storage stability was investigated. In accordance with the lower lipolytic activity of germinated oat, the accumulation of free fatty acids was slower in germinated oat when compared to native oat during the first two months of storage (Heiniö et al. 2002). After that, the formation of free fatty acids was similar or faster in germinated oat than

in native oat. However, the degree of unsaturation characterizing oxidation of lipids did not change in germinated oat samples but was clearly reduced in native oat samples. Furthermore, the concentration of volatile aldehydes and ketones regarded as secondary oxidation products of unsaturated fatty acids was lower in germinated oat samples during the whole 12 month storage period when compared to native oat samples. The formation of volatile compounds related to lipid oxidation was closely correlated with the development of the undesired sensory attributes. Therefore, the sensory stability of germinated oat groats was better than that of oat groats made from native oat.

### Utilization of germinated oats

Germination process can be regarded as a pretreatment of cereal raw material with aims to modify the structure and composition of kernel for further processing. The changes in structure may influence the processability of kernels e.g. in fractionation or extraction processes, and furthermore, influence the absorption of nutrients or the bioavailability of functional components in humans and aminals. In addition, the softer kernel structure of germinated oat opens new possibilities to use whole kernels in human and animal nutrition.

Oat malt is commercially available from several companies. So far, it has been mainly used in breweries as a minor raw-material in ale and stout production to tune beer flavor. A recent study (Taylor et al. 1998) showed that malted oat provided a pronounced toasted, biscuity aroma and palate, combined with a creamy and relatively intense mouthfeel in beer. These flavour effects were apparent at less that 10% replacement of barley malt with malted oat.

Some other application have also been developed for malted oat in bakery, biscuit, cereal bar and confectionary industries (Oat cuisine 1989). Also the production of a vitamin-rich oat drink (DE 3741991) and snack foods (PCT/FI01/ 00865) based on malted oat has been suggested. However, only a minor part of the great potential of germinated/malted oat has currently been utilised. Bread and bisquits as well as breakfast cereals are natural end-products for the use of oat malt. Textural and flavour changes induced during germination and finalised in the subsequent heat treatments could also be the basis of new product concepts, providing the modern consumer with healthy, whole grain foods in a palatable form.

As the present review shows, the basic biochemical reactions in oat kernel during germination are quite well known and documented. The tools to develop tailored germination processes and to create novel, functional oat compounds and fractions are therefore available. The main challenges for the development of new ingredients and products from germinated oat is to find the innovative ways to combine different prosessing methods i.e. drying, fractionation, extraction, extrusion with germination and to formulate the final product concepts that attract consumers of different origin. In addition, the stability and sufficient shelf-life are important success criteria of all oat products and should therefore always be addressed.

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## SELOSTUS

#### Idättämällä voi muokata kaurasta uudenlaisia elintarvikeraaka-aineita

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Kaura on terveellinen ja monipuolinen elintarvikeraaka-aine, ja se on tärkeä viljelykasvi suomalaisessa maataloudessa. Kauraa käytetään varsin perinteisesti, ja niinpä kauran menekin ja erityisesti jalostettujen kauratuotteiden viennin lisäämiseksi tarvitaan uudenlaisia prosessointikeinoja ja tuoteideoita. Idätys on vanha keino muokata viljojen siementen koostumusta ja pehmentää niiden rakennetta. Idättämällä on pilkottu huonosti sulavia siementen komponentteja sekä parannettu siementen makua ja ravintoarvoa. Idätyksen tärkein sovellus on ohran mallastaminen oluen raaka-aineeksi. Viime aikaisen terveysvaikutteisten elintarvikekomponentien tutkimuksen myötä on kiinnostuttu myös itäviin siemeniin muodostuvista bioaktiivisista yhdisteistä ja itävän siemenen tuottamien entsyymien hyödyntämisestä muokattujen elintarvikeraaka-aineiden valmistuksessa. Bioaktiivisilla yhdisteillä on mm. antioksidatiivisia ja antimikrobisia ominaisuuksia. Hallitulla idätystysprosessilla on mahdollista saada haluttuja yhdisteitä ja ominaisuuksia idätettyihin viljoihin. Tämä kuitenkin edellyttää, että sekä idätysprosessi että itämisen aikaiset biokemialliset tapahtumat tunnetaan hyvin. Kauran idätystä on tutkittu jonkin verran, ja tämä katsaus osoittaa, että idätys on potentiaalinen keino muokata kauraa maistuvien, terveellisten ja laadultaan korkealuokkaisten kaurajalosteiden raaka-aineeksi.