CHAPTER 11

Sourdough and cereal-based foods: Traditional and innovative products

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Cereal-based foods have been key components of human diets for thousands of years (Alfonzo *et al.* 2013); several historical sources evidence that baking of leavened doughs was a daily practice in several cultures (Corsetti and Settanni 2007). Furthermore, as revealed by the discovery of fossil kernels, humans' utilization of cereals commenced in the Neolithic era (Settanni and Moschetti 2010). Cereals remain a major source of nutrition, particularly in developing and overpopulated countries (Blandino *et al.* 2003). Indeed, the history of several cultures is directly defined by cereals and, consequently, many human populations are identified by the cereals they eat: Chinese are 'rice people', South and Central Americans are 'maize people', North Americans and Mediterranean people are 'wheat people', North Europeans are 'oats and rye people' and Africans are 'millet and sorghum people' (Gifford and Baer-Sinnot 2007).

Millennia BCE, wheat was already one of the most important crops grown in the Mediterranean basin. Following mass migrations, its cultivation underwent a huge expansion, resulting in its production worldwide (Toderi 1989). Among cereals, wheat is critical in the Mediterranean diet: it provides approximately one-third of the daily protein and energy requirements (~2400 kcal) for an adult (Cannella and Piredda 2006). For this reason, wheat surpasses other cereals in terms of the number of hectares dedicated to its cultivation worldwide (Gifford and Baer-Sinnot 2007). However, the world's major sources of energy for humans are rice, wheat and maize (Spiertz and Ewert 2009).

Due to their potential for nutritional enhancement and the fact that their consumption substantially lowers the risk of significant diet-related diseases (Topping 2007), cereals also assume a basic role in the diet of industrialized countries. They are generally consumed after boiling or after fermentation.

In the latter case, cereals are first ground and mixed with water (Salovaara 1998). In this matrix, different microbial groups, including mainly bacteria and yeasts, transform raw materials into final products (Galati *et al.* 2014).

As already mentioned, the process of cereal fermentation has been important for humans since ancient times (Spicher 1999; De Vuyst and Neysens 2005). The acidic product, depending on the proportion of water, may represent a food or a beverage; sometimes it undergoes baking. Either way, fermentation contributes to the microbial stability of the final products (Mensah 1997). Hence, fermented cereal-based foods are complex microbial ecosystems, whose activities confer on the resulting products characteristic features such as palatability, high sensory quality, structure and texture, stability, nutritional and healthful qualities (Corsetti and Settanni 2007) and, when they are in a living form at the moment of consumption (unbaked products), potential probiotic properties (Perricone *et al.* 2014).

Traditional fermented foods prepared from most common types of cereals (such as rice, wheat, maize or sorghum) are well known in many parts of the world. In general, most of these foods are typical to restricted geographical areas, where the cereal substrates (alone or mixed with other cereals or legumes or tubers), in the form of flour, are processed into different products (Blandino *et al.* 2003). Among these foods, bread is common to many societies; it is produced almost everywhere, even in southeast Asian countries that have not been traditional bread consumers (Jenson 1998).

Products derived from cereal flours (e.g. bread, cereal snacks and breakfast cereals) are useful food vehicles to provide micronutrients, but sometimes the amount needed for correct alimentation is quite limited. For this reason, policy and programme responses of several countries, including those located in developing areas, promote food-based strategies, such as food fortification, to prevent micronutrient malnutrition. Food fortification allows delivery of the required nutrients to many populations without requiring radical changes in food consumption (Allen *et al.* 2006).

Bread is mainly produced from wheat, but rye and barley are often used for this purpose. Due to their gluten content, these cereals are toxic to people affected by coeliac sprue (CS), also known as gluten-sensitive enteropathy, an autoimmune disease of the small intestinal mucosa (Silano and De Vincenzi 1999). Although several attempts are being made to decrease the CS-inducing effects of gluten by enzymatic treatment (Caputo *et al.* 2010), to date a strict, life-long gluten-free diet is the only safe and efficient treatment available for this disease (Tack *et al.* 2010). For this reason, the development or enhancement of gluten-free products continues to grow (O'Shea *et al.* 2014). Due to the lower-quality characteristics of these products compared to those made with wheat flour, several studies are in progress to determine the best flour combinations and, for fermented foods, to select the starter strains able to enhance the quality

aspects of gluten-free products. Another strategy to produce products compatible with a CS diet is to hydrolyse the toxic components of flours that contain gluten. From this standpoint, many attempts have been made to apply different microorganisms, mainly lactic acid bacteria (LAB) and microbial proteases (Rizzello *et al.* 2014b), but much more work still needs to be done before the efficacy and effectiveness of the microbial activities are proven to obtain safe foods for CS-affected people.

Cereals used in fermented food production

To date, more than half of the arable land in the world has been planted with cereals, mainly wheat, rice, maize, barley, sorghum and millet (Toderi 1993). Wheat clearly dominates in terms of hectares dedicated to its cultivation. However, thanks to improved plant breeding, rice production per hectare is higher than that for wheat. Other cereal crops are also relevant in several areas throughout the world (Dahlberg 2007).

Thanks to the great adaptability of its several varieties to different climatic conditions, including extreme values of temperature and/or humidity, wheat is cultivated almost everywhere for food and feed production. The species most commonly involved in food production are *Triticum durum* and *Triticum aestivum* (Toderi 1993). *T. durum*, which requires high temperatures, is cultivated in temperate regions, while *T. aestivum*, well suited to lower temperatures, is mainly cultivated in colder areas. In Italy, where cereal-based foods constitute the major part of the daily diet, *T. durum* and *T. aestivum* are traditionally used for pasta and bakery products (bread and leavened baked goods, such us breakfast or recurrent products), respectively. However, in some southern regions, durum wheat flour is used alone or in combination with soft wheat flour for bread production (Settanni *et al.* 2008).

The reason for the great expansion of wheat cultivation is that the other cereals are not as suitable for the production of foods characterized by nutritional value, shelf life and taste as those obtained from wheat flour (Macrae *et al.* 1993; Francis 2000; Cannella *et al.* 2010). In addition to wheat, rye is particularly common for bread production in Scandinavian countries, Germany, Poland and Russia (Bushuk 2001), maize is used in Portugal (Rocha and Malcata 1999) and sorghum in Sudan (Hamad *et al.* 1997), while rice is commonly used for glutenfree bread production (Neumann and Bruemmer 1997; Meroth *et al.* 2004).

The most cultivated cereal in developing countries is maize (Giardini and Vecchiettini 2000). It has been used for human consumption for centuries. The different varieties belonging to the species *Zea mays* are mainly distinguished by colour, basically white or yellow. The latter cultivars are particularly rich in carotenoids (pro-vitamin A; Schober and Bean 2008). The current maize

production could be considered sufficient to meet the caloric needs of nearly 2 billion people (Giardini and Vecchiettini 2000). The peculiar element of maize is starch, which is used in the confectionery industry.

In Asia and Africa, sorghum constitutes an integral part of the basic diet for millions of people (Schober and Bean 2008). Millet is considered to be the oldest cereal, widely cultivated in Asia and Africa and in some areas of eastern Europe (Taylor and Emmambux 2008). Barley is the main food source for a large number of people living in cold or semiarid areas where wheat does not fit (Mosca and Toniolo 2000).

The cereals mentioned are used to produce fermented products throughout the world (Blandino *et al.* 2003; De Vuyst and Neysens 2005; Galati *et al.* 2014). In particular, millet, maize, sorghum, rice and wheat are mainly used in Africa; maize, rice and wheat in America and Australia; wheat, maize, rye, barley and millet in Europe; rice, millet and wheat in Asia.

Technological properties of cereal flours

The choice of raw materials is crucial to obtain bakery products able to satisfy "consumers' needs", and this is particularly important in breadmaking. For this purpose, wheat cultivars have been selected by bread producers since ancient times (Bottega et al. 2010). The technological aptitudes of wheat flour depend on its ability to form gluten, defining of its versatility. Gluten is generated by the interaction between two groups of proteins, gliadins and glutenins, when flour is mixed with water and the mixture is allowed to stand for a while (Cannella et al. 2010). The resulting dough is characterized by a viscoelastic behaviour that is responsible for its extensibility during kneading, leavening and the early stages of cooking, but also confers toughness and elasticity, which allow it to maintain its shape and to develop mass with regularity (Bottega et al. 2010). Although all wheat varieties are able to form gluten, the presence of particular protein subunits (different in structure and relationship gliadin/glutenin, composition and molecular weight) ensures the formation of tenacious nets (MacRitchie 1992; Shewry 2003; Bottega et al. 2010).

Other cereals, such as rye, contain protein subunits similar to those of wheat, but their ability to expand to give a well-developed leavened product is limited (He and Hoseney 1991; Flander *et al.* 2007). Even more critical is the workability of wheat dough enriched with gluten-free flours (Mariotti *et al.* 2006, 2008; Schoenlechner *et al.* 2006). These flours negatively influence the leavening of the dough. Doughs made from durum wheat semolina are characterized by high strength and limited extensibility (Pogna *et al.* 1996). During leavening, therefore, dough tends to have a development by volume lower than that of a good soft wheat flour (Raffo *et al.* 2003).

Cereal microflora

Cereal grains are naturally contaminated by eukaryotic (moulds and yeasts) and procaryotic (bacteria) organisms. The total microbial population and the relative species proportion on wheat grains can be affected by many factors, mainly climatic conditions such as temperature and rainfall, physical damage due to insects or mould attacks and application of insecticides and fungicides. Microorganisms of grains might follow different phases of flour preparation and, since flour does not undergo thermal treatment, it is a source of living and active microorganisms that can be found in the resulting fermented foods (Corsetti and Settanni 2007). The microorganisms that contaminate cereals are generally concentrated in the outer layers of kernels, and they tend to stay in fractions rich in bran during milling. Consequently, flour obtained from milling should theoretically contain a lower bacterial load than caryopses, but the subsequent conditioning phase can increase its microbial content (Berghofer *et al.* 2003).

The levels of living microorganisms present on cereals might range between 10⁴ and 10⁷ colony forming units (cfu) per gram, while they reach cell densities up to 10⁶ cfu/g in the corresponding flours (Stolz 1999). The bacteria, mainly mesophilic, include Gram-negative aerobes (e.g. *Pseudomonas*) and facultative anaerobes (*Enterobacteriaceae*) and Gram-positive species (De Vuyst and Neysens 2005; Minervini *et al.* 2014). Among the latter bacterial groups, LAB, which together with yeasts are relevant for the process of food fermentation, might be found in several spontaneously fermented cereal-based products (Galati *et al.* 2014).

Although several studies have focused on the identification and characterization of LAB in the final fermented products, only a few works have investigated the microbial ecology of raw materials used for cereal-based food production. Some studies are available on the cereals, and the corresponding flours, used in breadmaking. The first document dates to 1987 (Galli and Franzetti 1987), when several samples of Italian wheat flours were analysed for the presence of different microbial groups. Subsequently, Corsetti et al. (1996) isolated and identified LAB and yeasts from common wheat and organic flours. Both studies were performed with a phenotypical/biochemical approach that revealed several lactobacilli (Lactobacillus alimentarius, Lactobacillus plantarum, Lactobacillus rhamnosus, Lactobacillus confusus and Lactobacillus viridescens), even though some species were then reclassified as weissellas (Weissella confusa and Weissella viridescens).

The cultivable LAB populations associated with durum wheat kernels, cultivated in several Italian regions, as well as bran and non-conventional flours (amaranth, chickpea, maize, rice, quinoa and potato) used to produce gluten-free baked goods, were found at levels ranging between 1.00 and 2.16 log cfu/g. The isolates were genetically investigated by applying a polyphasic

strategy consisting of randomly amplified polymorphic DNA-polymerase chain reaction (RAPD-PCR) analysis, partial 16S rRNA gene sequencing, species-specific and multiplex PCRs (Corsetti et al. 2007a). Besides Lactobacillus (Lactobacillus coryniformis, Lactobacillus curvatus and Lactobacillus graminis), the work revealed the presence of other LAB genera: Aerococcus (Aerococcus viridans), Lactococcus (Lactococcus garvieae), Enterococcus (Enterococcus durans, Enterococcus casseliflavus, Enterococcus faecalis, Enterococcus faecium and Enterococcus mundtii) and Pediococcus (Pediococcus pentosaceus). De Vuyst and Neysens (2005) reviewed the presence of several LAB species (Lactobacillus casei, Lb. coryniformis, Lb. curvatus, Lb. plantarum, Lactobacillus salivarius, Lactobacillus brevis, Lactobacillus fermentum, Ent. faecalis, Lactococcus lactis, Pediococcus acidilactici, Pediococcus parvulus, Pe. pentosaceus, Leuconostoc and Weissella) on cereal grains.

More recently, Alfonzo et al. (2013) investigated different wheat (T. durum and T. aestivum) flours used for the production of traditional sourdough breads in Sicily (southern Italy) using culture-dependent phenotypical and genotypical tools, as well as a culture-independent method based on the denaturing gradient gel electrophoresis (DGGE) technique. The last approach was useful to test the technological performance of the dominant strains (which reached concentrations up to 4.75 log cfu/g) and also to detect the species present at undetectable (subdominant) levels and/or as dormant (non-cultivable) flora. Ent. mundtii, Lactobacillus sanfranciscensis, Lb. plantarum, Lactobacillus sakei, Lc. lactis, Leuconostoc mesenteroides, Leuconostoc pseudomesenteroides, Leuconostoc citreum, Pe. pentosaceus, Weissella cibaria and We. confusa were identified. In particular, the most prevalent species detected were We. cibaria, Lb. plantarum, Ln. pseudomesenteroides and Ln. citreum. DGGE analysis confirmed the detection of the genera to which most isolates belonged (Lactobacillus, Enterococcus, Leuconostoc and Weissella), but only two species, Lb. plantarum and Ln. citreum, were clearly identified. The apparent lack of correspondence between culturedependent and culture-independent methods was explained by the fact that when bacteria isolated by plating are not detected with DGGE analysis based on 16S rRNA gene amplification, they are not a major component of the microbial community being investigated (Shinohara et al. 2011). For this reason, a combined approach consisting of both methodologies provides the best strategy for detection of microbial communities within complex food matrices (Carraro et al. 2011).

Regarding eukaryotic organisms, yeasts are also detected both on the cereal surface and in flour samples ranging from a few cells to 10⁴ and 10³ cfu/g, respectively. The species most commonly found are *Candida*, *Cryptococcus*, *Pichia*, *Rhodotorula*, *Saccharomyces*, *Sporobolomyces*, *Torulaspora* and *Trichosporon*. Among fungi (circa 10⁴ cfu/g), *Alternaria*, *Cladosporium*, *Drechslera*, *Fusarium*, *Helminthosporium* and *Ulocladium* (from the field), and *Aspergillus* and *Penicillium* (from storage), are found (De Vuyst and Neysens 2005).

Furthermore, it is also important to consider that the microorganisms performing the fermentation of cereal flours may originate from the equipment used in the milling and/or production process (Berghofer *et al.* 2003).

Cereal fermentation

Since the beginning of human civilization there has been an intimate relationship between the human being and the fermentative activities of microorganisms. These activities have been utilized in the production of fermented foods and beverages, which are defined as those products that have been subordinated to the effect of microorganisms or enzymes determining desirable biochemical changes (Settanni and Moschetti 2014). Fermentation represents the oldest and most economical method of producing and preserving food (Chavan and Kadam 1989; Billings 1998). In fact, fermentation helps in the production of safe and stable foods with a longer shelf life than their raw materials. These foods are more digestible and appealing than unprocessed substrates because they acquire new desired organoleptic characteristics (Settanni and Moschetti 2014). In addition, fermentation provides a natural way to reduce the volume of the material to be transported, to destroy undesirable components, to enhance the nutritive value and to reduce the energy required for cooking (Simango 1997). During cereal fermentation several volatile compounds are formed, which contribute to a complex blend of flavours in the processed products (Chavan and Kadam 1989).

The microorganisms responsible for fermentation may be the microflora indigenously present on the substrate (and this occurred unknowingly for millennia) or they may be added as starter cultures (Harlander 1992). The latter strategy commenced when the microorganisms had been isolated and their activities discovered and studied.

By one biochemical definition, fermentation is an anaerobic process for deriving energy from the oxidation of organic compounds using an endogenous electron acceptor, which is usually an organic compound (Prescott *et al.* 2005). Following this process, the carbohydrates are partially oxidized and several microorganisms produce energy by means of this metabolic pathway. Among the several fermentations employed to produce different foods (Soni and Sandhu 1990), the two main processes that are defining for the transformation of cereal flours are alcoholic and lactic acid fermentations. Alcohol fermentation results in the production of ethanol and yeasts are the predominant organisms involved; lactic acid fermentation is carried out by LAB (Corsetti and Settanni 2007).

In general, natural fermentation of cereals leads to a decrease in the level of carbohydrates as well as some non-digestible poly- and oligosaccharides. This process determines the saccharification of starch and increases the availability of

proteins (Blandino *et al.* 2003). Certain amino acids may be synthesized and the availability of B-group vitamins might be improved. Fermentation also provides optimum pH conditions for the enzymatic degradation of phytate, which is present in cereals in the form of complexes with polyvalent cations such as iron, zinc, calcium, magnesium and proteins. Such a reduction in phytate may increase the amount of soluble iron, zinc and calcium severalfold (Haard *et al.* 1999; De Angelis *et al.* 2003).

Cereal-based fermented products

Based on the type of flour (alone or in combination), the fermenting agent(s) and the technological process(es) applied, different products can be obtained. The fermenting microorganisms for almost all cereal-based fermented products are mainly LAB and a few yeast species. In several cases, their combined action allows the production of foods and beverages with the desired quality characteristics (Galati *et al.* 2014). However, the microbial populations responsible for the fermentation of several niche cereal-based products, especially those produced in countries where this process is not driven by selected starter cultures, remain unknown.

Cereal-based fermented foods are spread all over the world and undoubtedly bread is the main product. Bread is the typical and oldest food of leavened products and it is a symbol of religion. The earliest records date back to the second millennium BCE, when, after a flood of the Nile that covered the grain reserves, the Egyptians realized that grain flour when mixed with water increased in volume over time (Di Giandomenico 2010). During the Second World War, due to the scarcity of cereals, a large number of cereals (maize, rye, oat, barley, rice, sorghum) or other vegetable sources (bean, cassava, soy, potato, chestnut flour etc.) have been used in breadmaking in place of wheat (INSOR 2012).

To date, unlike bread, where biotechnologies are under control during the transformation process carried out at an industrial level, the preparation of many other cereal-based fermented foods is restricted to limited areas and they are mostly produced on a domestic scale (Owczarek *et al.* 2004). However, some of them are particularly important in different countries and have been the object of study for many research groups (Table 11.1).

Bread

The term 'bread' refers to a food of any shape and dimension obtained from a dough prepared with flour and water, with or without salt, fermented naturally or with the addition of yeasts and subsequently subjected to cooking. Breadmaking technology (Figure 11.1) is quite simple (Pagani *et al.* 2006).

The production of leavened bakery products can be summarized as the semisolid mass transformation of the dough, a particular 'emulsion', characterized by

Table 11.1 Traditional fermented cereal-based foods and beverages that have been the object of scientific investigation.

| Products | Cereals | Countries |
|-----------------|-----------------------------------|------------------------------------|
| Amgba | Sorghum, millet | Cameroon, Chad |
| Ang-kak | Rice | China, Indonesia, Thailand |
| Atole | Maize | Guatemala, Mexico |
| Ben saalga | Millet | Burkina Faso |
| Boza | Wheat, millet, maize | Albania, Bulgaria, Romania, Turkey |
| Bread | Wheat, rye, barley, sorghum, rice | Five continents |
| Bushera | Sorghum, millet | Uganda |
| Chicha | Maize | Peru |
| Chikokivana | Maize, millet | Zimbabwe |
| Chongju | Rice | Korea |
| Dolo | Sorghum | Burkina Faso |
| lkikage | Sorghum | Rwanda |
| Jaanr | Millet | Northeastern Himalayas |
| Kachasu | Maize | Zimbabwe |
| Kaffir beer | Maize | South Africa |
| Koko | Maize | Ghana |
| Kunu-Zaki | Millet | Nigeria |
| Lao-chao | Rice | China, Indonesia |
| Mantou | Wheat | China |
| Mirin | Rice | Japan |
| Muramba | Sorghum | Uganda |
| Mutwiwa | Maize | Zimbabwe |
| Ogi | Maize, sorghum, millet | Nigeria, West Africa |
| Pito | Maize, sorghum | Ghana, Nigeria |
| Pozol | Maize | Mexico |
| Puto | Rice | Philippines |
| Sake | Rice | Japan |
| Sourdough bread | Wheat, rye, barley, sorghum, rice | Five continents |
| Takju | Rice, wheat | Korea |
| Tape ketan | Rice | Indonesia |
| Tapuy | Rice | Philippines |
| Tchoukoutou | Sorghum, millet, maize | Benin, Togo |
| Tesguino | Maize | Mexico |
| Togwa | Maize, sorghum, millet | Tanzania |

a continuous phase represented by hydrated gluten that surrounds the starch granules in which are dispersed microbubbles of air, in a 'foam', that is to say in a product in which the continuous phase retains significant volumes of gas (Bottega *et al.* 2010).

The leavening process is of paramount importance during breadmaking. It determines the increase of dough volume, the development of precursors of the aroma compounds and the improvement of the nutritional characteristics of flour.

208

Figure 11.1 General process of bread production.

The volume expansion can be achieved in different ways: the biological approach (thanks to carbon dioxide $[CO_2]$ resulting from yeasts and/or LAB metabolism); the use of chemicals added to the formulation (which exert their action especially during cooking); and the physical approach, with the inclusion of air as a result of intensive mechanical action and typical leavening of some formulations rich in fat (Bottega *et al.* 2010).

Biological leavening assumes a basic importance for the sensory properties of the final product. This process can be carried out by baker's yeast, mainly represented by *Saccharomyces cerevisiae*, and a mixture of yeasts and LAB. In the latter case, the technology applied is that of sourdough, referred to as a mixture of flour and water in which the development of LAB results in the production of lactic acid and acetic acid. LAB developing in the dough may originate as contaminants of flour and/or the bakery environment or might derive from a starter culture containing one or more known species of LAB (De Vuyst and Neysens 2005).

Sourdough

Sourdough is a mixture of cereal flour and water in which a heterogeneous population composed of LAB and yeasts is metabolically active, either by spontaneous fermentation or by fermentation initiated through the addition of a sourdough starter culture, whether or not it involves backslopping (De Vuyst *et al.* 2009).

Based on the technology applied, sourdoughs have been grouped into type I, type II and type III (Böcker *et al.* 1995). Type I sourdoughs are traditionally characterized by continuous, daily refreshment to keep the microorganisms in an active state; fermentation is carried out at room temperature until the pH reaches a final value of around 4.0 (Corsetti and Settanni 2007). Type II sourdoughs are semifluid silo preparations fermented at temperatures higher than 30 °C for long periods (at least 2 days); these sourdoughs are generally added during bread preparation as dough-souring supplements (Böcker *et al.* 1995; Hammes and Gänzle 1998). Type III sourdoughs are dried preparations containing LAB resistant to the drying process (Hammes and Gänzle 1998). Types II and III sourdoughs require the addition of *S. cerevisiae* as a leavening agent. Some authors

(De Vuyst and Neysens 2005) also reported on type 0 dough. This product, although prepared exclusively from baker's yeast and not made with sourdough technology, also contains LAB species as contaminants of the yeast inoculums. These are especially lactobacilli rather than *Pediococcus*, *Lactococcus* and *Leuconostoc* spp. (Jenson 1998), and contribute only to a small degree to the acidification and aroma development of dough because of the short processing time.

The addition of sourdough improves the texture, flavour, nutritional aspects and shelf life (Gänzle *et al.* 2007) of wheat bread produced from baker's yeast due to the synthesis of aroma compounds (Czerny and Schieberle 2002; Hansen and Schieberle 2005), enzymes and antibacterial (Settanni *et al.* 2005) and antifungal (Ryan *et al.* 2008; Poutanen *et al.* 2009) compounds during fermentation. Sourdough fermentation has been demonstrated also to enhance the sensory quality of rye breads (Rizzello *et al.* 2014b).

Besides the technological and nutritional aspects, sourdough fermentation may also provide several health benefits, such as a decrease of the glycemic response of baked goods, enhancement of the content of bioactive compounds and increase of mineral uptake (Gobbetti *et al.* 2013). Compared with sourdough bread, white wheat bread started with baker's yeast alone shows some drawbacks such as low protein digestibility, high carbohydrate content, high glycemic index, low-resistant starch and low level of dietary fibre (Dhinda *et al.* 2011).

Sourdough microorganisms

The proportions of LAB and yeasts in sourdough are not random, but rather respect a defined relationship. The ratio between LAB and yeasts is generally reported to be 100:1 (Gobbetti *et al.* 1994; Ottogalli *et al.* 1996). This could be the result of direct interaction between these microbial groups.

The study of the social activities exhibited by microorganisms is defined as 'sociomicrobiology' (Parsek and Greenberg 2005). The microorganisms were for a long time believed to exist as single cells in a given environment searching for nutrients to multiply. Bacteria were the first microbes to be studied in order to decipher their code of communication. It has been discovered that they are active in performing a census of their population, as well as in investigating the environment for development and in feeling the presence of competitors (Fuqua et al. 1996). Such actions are the result of efficient intercellular communication that is based on the production, release and detection of and reply to small signal molecules, which accumulate and trigger cascade events when a 'quorum' concentration is reached. Hence, the term 'quorum sensing' is used to describe cell-cell communication. Based on this system the bacteria can 'count' one another (Fuqua et al. 1994), acting as a group. Recently, it became apparent that fungi, like bacteria, also use quorum regulation to affect population-level behaviours. Furthermore, considering the extent to which quorum-sensing regulation

controls important processes in many distantly related bacterial genera, it is not surprising that cell density–dependent regulation also appears to be prevalent in diverse fungal species (Hogan 2006).

The studies regarding cell–cell communication in sourdough ecosystems are so far limited to the group of LAB (Di Cagno *et al.* 2007, 2010), but this topic is under study to elucidate the mechanisms of interdomain communication between LAB and yeasts.

Yeasts

Several sourdoughs are reported to host S. cerevisiae. This finding is generally due to introduction through the addition of baker's yeast (Corsetti et al. 2001) or to cross-contamination in bakeries where both conventional and sourdough breads are produced (Valmorri et al. 2010). Yeasts found in sourdoughs belong to different genera (Rossi 1996; Stolz 1999; Gullo et al. 2002). The typical sourdough yeasts are Saccharomyces exiguus, Candida humilis (formerly described as Candida milleri) and Issatchenkia orientalis (Candida krusei; Garofalo et al. 2008; Iacumin et al. 2009). Other yeast species detected in sourdough ecosystems are Pichia anomala (as Hansenula anomala), Saturnispora saitoi (as Pichia saitoi), Torulaspora delbrueckii, Debaryomyces hansenii, Pichia membranifaciens (Corsetti and Settanni 2007) and Candida famata (Mohamed et al. 2007). The extensive variability in the number and type of species found depends on several factors, dough yield (weight of dough/weight of flour × 100), type of cereal used, temperature of fermentation and temperature for sourdough maintenance (Gobbetti et al. 1994). Yeasts via alcoholic fermentation are primarily responsible for the leavening of dough (Corsetti and Settanni 2007).

Lactic acid bacteria

LAB are involved in the process of acidification of sourdough, but heterofermentative species partly contribute to the mass blowing (Gobbetti *et al.* 1995).

Unlike other fermented foods, where LAB responsible for the transformation of raw materials into final products belong to obligate homofermentative and/or facultative heterofermentative species, obligate heterofermentative species play a major role in sourdough (Salovaara 1998), especially when sourdoughs are prepared in a traditional manner (Corsetti et al. 2001, 2003). Typical sourdough LAB mainly belong to the genus Lactobacillus and include all three metabolic groups discussed (Hammes and Vogel 1995). However, other LAB belonging to Leuconostoc, Weissella, Pediococcus, Lactococcus, Enterococcus and Streptococcus genera have been isolated from sourdough (Corsetti and Settanni 2007).

Most of the LAB species commonly detected in sourdough (Figure 11.2) have sourdough as their primary and sole source of isolation, probably because no other ecosystem can support their growth. For instance, except sourdough, no other habitat is known for *Lactobacillus sanfranciscensis* (Hammes *et al.* 2005).

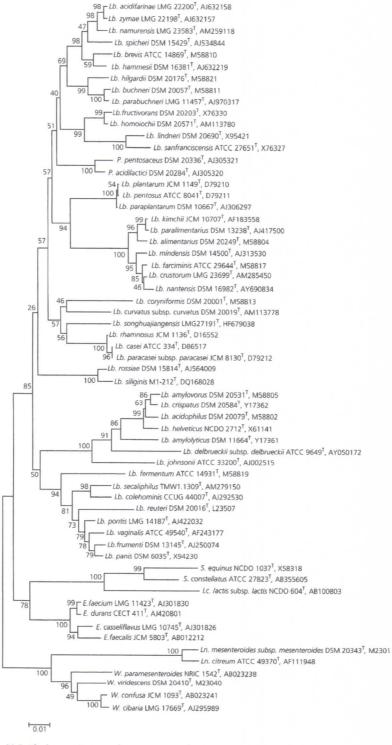


Figure 11.2 Phylogenetic tree of LAB commonly associated with or found in sourdough products based on 16S rRNA gene sequences. Sequence alignment was performed with CLUSTALX (Thompson *et al.* 1997). Sequence and alignment manipulations and calculation of similarity values and nucleotide compositions of sequences were performed with the GeneDoc program version 2.5.000 (K.B. Nicholas and H.B. Nicholas, unpublished data). Positions available for analysis were circa 1150 bp. Phylogenetic and molecular evolutionary analysis was conducted using MEGA version 3.1 (Kumar *et al.* 2004). Bar 0.01 nucleotide substitution per site.

212

Some of the species associated with this environment have been misidentified because of the lack of application of molecular methods, which were unavailable in the past, or have been found in sourdough due to cross-contamination.

Some non-Lactobacillus species isolated from sourdough have only been detected at subdominant levels (Corsetti et al. 2007b). As an example, in mature Italian type I sourdough, E. faecium and P. pentosaceus have been found in the range 10⁴–10⁶ cfu/g, while lactobacilli were about 2–3 orders of magnitude higher (Valmorri et al. 2006). Corsetti et al. (2007b) investigated the role of Ent. faecium and Pe. pentosaceus during sourdough preparation. Strains of both species were followed in dual combination with Lb. sanfranciscensis. During the first steps of sourdough preparation, single inocula of Ent. faecium and Pe. pentosaceus determined a stronger and more rapid acidification of dough than the Lb. sanfrancisensis strain used. Subsequently, the behaviour monitored during the co-fermentation of Ent. faecium/Lb. sanfranciscensis and Pe. pentosaceus/Lb. sanfranciscensis showed that Ent. faecium and Pe. pentosaceus prepare the environment for the establishment of the typical species of mature sourdough, including Lb. sanfranciscensis, by lowering the pH.

Minervini et al. (2014) described the typical LAB population dynamic in sourdough, referred to as 'three-phase evolution', regardless of the type of flour. The three phases indicated are the dominance of LAB species belonging to the genera Enterococcus, Lactococcus and Leuconostoc; the increasingly important presence of sourdough-specific LAB, such as species belonging to the genera Lactobacillus, Pediococcus and Weissella; and the dominance of well-adapted sourdough strains, belonging to obligate heterofermentative species such as Lb. sanfranciscensis, Lb. fermentum and Lactobacillus pontis and to the facultative heterofermentative Lb. plantarum (Gänzle et al. 2007), although the presence of some Leuconostoc spp. is sometimes revealed. This succession of LAB is mainly driven by different tolerance to acidic conditions and to different adaptation mechanisms related to carbohydrate and nitrogen metabolism (Gänzle et al. 2007). The establishment of obligate heterofermentative lactobacilli is essential for the optimal fermentation of traditional sourdough (Salovaara 1998).

Regarding cell–cell communication among lactobacilli in sourdough, Di Cagno et al. (2007) followed, by a proteomic approach, the growth of *Lb. sanfranciscensis* in mono-culture and co-culture with *Lb. plantarum*, *Lb. brevis* or *Lactobacillus rossiae*. When co-cultured, the *Lb. sanfranciscensis* strain, depending on the combination, overexpressed several proteins during the late stationary phase. The induced polypeptides, only in part common to all co-cultures, were identified as stress proteins, energy metabolism–related enzymes and proline dehydrogenase, GTP-binding protein, *S*-adenosyl-methyltransferase and Hpr phosphocarrier protein. Furthermore, two quorum-sensing genes involved, *luxS* and *metF*, were shown to be expressed in the *Lb. sanfranciscensis* strain studied. Later, the same research group studied the effect of pheromone plantaricin A produced by a *Lb. plantarum* strain (DC400) towards other sourdough LAB, and it was found that this

pheromone influenced the growth and survival of the other strains co-cultivated with *Lb. plantarum* DC400 differently (Di Cagno *et al.* 2010).

Starter selection for sourdough production

In the last few years, several research groups have focused their attention on the selection of strains to be used as starter cultures for controlled sourdough fermentation. Analysis of microbial biodiversity and monitoring the dominant microorganisms during fermentation is a key step for the selection of strains able to drive the transformation processes. The starter cultures for fermentation are selected based on their specific technological traits in order to obtain final products with a set of desired characteristics (Figure 11.3).

Generally, LAB are primarily tested for their rapid acidification, whereas yeasts are first tested for their alcoholic fermentation rate. However, depending on the type of bread to be produced, LAB are commonly selected also for other technological performance, such as contribution to the development of the flavour and structure of the dough, reduction of antinutritional factors and, in order to determine microbial stability during fermentation and elongate the

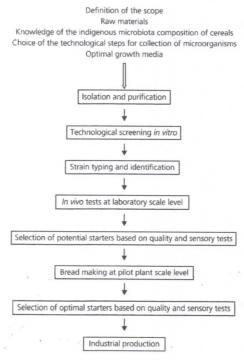


Figure 11.3 Schematic representation of the plan for selection of starter culture(s) to be used for sourdough fermentation.

shelf life of the final products, for their competitiveness with undesired microorganisms based on the production of secondary metabolites, mainly bacteriocins and antifungal compounds (Corsetti and Settanni 2007).

The selection of LAB is performed during fermentation; this allows the collection of dominant strains that have competitive advantages over the native microbiota. The dominant strains will ensure, to a certain extent, the successful transformation of flour into sourdough, thanks to their adaptation to the specific fermentation technology, the environmental conditions and the availability of substrates. Once the dominant isolates are collected, their performance is evaluated in vitro, first using the optimal commercial synthetic media and then using sterilized media prepared from raw material extracts that mimic real conditions, without interaction with indigenous microorganisms (Settanni et al. 2014). Subsequently, the isolates showing interesting properties are identified at strain and species levels. The different strains are tested in vivo using untreated flours, so that their performance is evaluated in the presence of the autochthonous microbiota, and their capacity to dominate the microbial community is monitored (Alfonzo et al. 2013). The final products are generally subjected to sensory and quality evaluations to select the best strains, in individual and/or multiple combinations, able to recreate the traditional aromatic profile, in order to obtain breads with the desired characteristics constant over time (Settanni et al. 2014).

Fortified fermented cereal-based products

Food fortification refers to the addition of micronutrients to processed foods. It represents a valid technology for reducing micronutrient malnutrition when and where existing food supplies and limited access fail to provide adequate levels of the respective nutrients in the diet. In industrialized countries, food fortification has long been used for the successful control of deficiencies of vitamins A and D, several B vitamins (thiamine, riboflavin and niacin), iodine and iron. From the early 1940s onwards, the fortification of cereal products with thiamine, riboflavin and niacin became common practice (Allen et al. 2006). Fortified breads are obtained through the enrichment of flour with nutrients. In fact, the nutritional features of white wheat are quite limited. This is due to the low levels of essential amino acids, such as lysine, and dietary fibre in white flour (Dhinda et al. 2011). For years, the most common constituents added to these kind of breads have been folic acid and iron. However, in the last few years the addition of alternative plant-based protein sources has become usual. It is becoming common practice in white bread production to use dietary fibre and ingredients or by-products rich in fibre (De Angelis et al. 2007, 2009; Rizzello et al. 2012), like a mixture of soy proteins, oat bran and legume flours (Sadowska et al. 2003; Kamaljit et al. 2010; Dhinda et al. 2011; Mohammed et al. 2012; Rizzello et al. 2014a) to enhance its nutritional value.

These products are gaining importance not only because of the increasing number of people on a vegetarian diet, but also due to the high energy requirements for animal protein production, particularly relevant for people who have no access to an animal protein—rich diet. For example, besides bread, several products are produced in different areas, especially Africa and Asia, with cereals in combination with legumes (Blandino *et al.* 2003), thus improving the overall protein quality of the fermented products. As reported earlier, cereals are deficient in lysine, but are rich in cysteine and methionine. Legumes, on the other hand, are rich in lysine but deficient in sulfur-containing amino acids. Thus, by combining cereals with legumes, the overall protein quality is improved (Campbell-Platt 1994). The main non-bread products obtained from the fermentation of mixtures of cereals and other vegetables are listed in Table 11.2.

Several attempts are being made to fortify the common ingredients used in bread production. Since bread is mainly obtained via the fermentation of wheat flour sugars derived from starch involving chemical interactions of the various food components, these interactions can be adjusted to create desirable products only if the chemical and physical processes are well understood (Sivam *et al.* 2010).

The incorporation of legumes in novel, convenient and healthy food products (Schneider 2002; Gòmez *et al.* 2008) represents a valuable strategy for increasing the global consumption of legumes, which is declining (Kohajdová *et al.* 2013) and is below the recommended amount (McCrory *et al.* 2010). For

Table 11.2 Traditional foods and beverages obtained through fermentation of mixtures of cereals and other vegetable sources.

| Products | Cereals and legumes | Countries |
|-----------|-----------------------------|------------------|
| Adai | Cereals and legumes | India |
| Banku | Maize and cassava | Ghana |
| Chee-fan | Wheat and soybean | China |
| Dhokla | Rice, wheat and Bengal gram | India |
| Dosa | Rice and Bengal gram | India |
| Hamanatto | Wheat and soybean | Japan |
| Idli | Rice and black gram | India, Sri Lanka |
| Kanji | Rice and carrots | India |
| Kecap | Wheat and soybean | Indonesia |
| Miso | Rice and soybean | China, Japan |
| Munkoyo | Maize and roots of munkoyo | Africa |
| Shoyu | Wheat and soybean | China, Japan, |
| | | Taiwan |
| Tao-si | Wheat and soybean | Philippines |
| Taotjo | Wheat, rice and soybean | India |
| Tarhana | Wheat and vegetables | Turkey |
| Tauco | Cereals and soybean | Indonesia |
| Vada | Cereals and legumes | India |

this reason, some reports have proposed the addition of legumes to different products such as bread (Dhinda *et al.* 2011; Mohammed *et al.* 2012), biscuits (Eissa *et al.* 2007; Tiwari *et al.* 2011), cakes (Gòmez *et al.* 2008), chapatti (Kadam *et al.* 2012) and crackers (Kohajdová *et al.* 2011).

When legume flours are used for breadmaking, the adjustment of several process parameters is needed to get a high sensory quality that is acceptable to the majority of consumers (Maninder et al. 2007; Kohajdová et al. 2013). One option to improve the sensory and functional quality of breads containing legume flours is represented by the use of sourdough fermentation. With this in mind, Rizzello et al. (2014a) developed a biotechnological protocol for manufacturing a white bread enriched with chickpea, lentil and bean flours through sourdough fermentation. For this purpose, the authors prepared type I sourdough containing legume flours according to traditional protocols routinely used for making typical Italian breads. LAB populations in wheat-legume sourdough included Lb. plantarum, Lb. sanfranciscensis, Ln. mesenteroides, Lb. fermentum, We. cibaria, Lactobacillus pentosus, Lb. coryniformis, Lb. rossiae, Lb. brevis, Lactobacillus parabuchneri and Lactobacillus paraplantarum, most of which are typically associated with mature sourdough. Compared with wheat-legume breads leavened with commercial yeasts, wheat-legume sourdough breads were characterized by higher quality parameters and acceptability by consumers, thus proving the defining role of sourdough technology in improving the characteristics of bakery products obtained with novel formulas.

Gluten-free cereals

Gluten is toxic for people affected by CS. Gluten causes self-perpetuating mucosal inflammation and subsequent loss of absorptive villi and hyperplasia of the crypts. Proteolytic enzymes of the endoluminal tract acting on prolamins of wheat (a-, b-, c-, and x-gliadin), rye (secalin) and barley (hordein) produce proline- and glycine-rich polypeptides that are responsible for the disease (Silano and De Vincenzi 1999). Triticale also contains gluten (Wolter et al. 2014a). The list of proteins that liberate toxic peptides includes the high molecular weight glutenins (Dewar et al. 2006). Both gliadins and glutenins are rich in two amino acids, proline (very resistant to the hydrolysis process) and glutamine. The gluten, when ingested, goes through a process of digestion and the protein is not completely degraded by enzymes with prolyl endopeptidase activities (Catassi and Francavilla 2010). The gluten is degraded into small peptides and their further proteolysis is made difficult by the position and abundance of proline residues (Hausch et al. 2003). Some of these peptides have an immuno-genetic activity for subjects predisposed to develop CS (Catassi and Francavilla 2010). For these reasons, those affected by CS cannot ingest glutencontaining products.

Most of the food products present on the market that characterize the daily diet in many countries are made with gluten-containing cereals. The number of people affected by CS is high, thus the production of baked goods that can be consumed by these subjects is of paramount importance. The main raw materials used for the production of gluten-free leavened baked are rice (Oryza sativa and Oryza glaberrima), maize (Zea mays), sorghum (Sorghum bicolor L. Moench) and millet (Panicum miliacum) and pseudo-cereals such as amaranth (species of genus Amaranthus), quinoa (species of genus Chenopodium) and buckwheat (species of genus Fagopyrum; De Angelis and Di Cagno 2010). Rice is one of the most important cereals used for this purpose because the carbohydrate fraction of the flour consists predominantly of starch present in small granules. Amylopectin, easily digested, is present in higher amounts than amylose. When rice flour is used, it provides softness and stability. Maize finds several uses (e.g. polenta, cornflakes, tortillas, snacks etc.) in the diet of coeliac subjects, but it is scarcely used for the production of gluten-free bread. The use of sorghum flour is recommended in the diet of CS patients, as it is the species phylogenetically most distant from the other cereals. Millet flour does not show a good aptitude for baking and for leavened gluten-free production, therefore it is often used in combination with other flours (Schober et al. 2003). Quinoa and amaranth have been utilized for the manufacture of different bakery products (Lorenz and Coulter 1991; Nsimba et al. 2008) and quinoa flour is also added for the manufacture of enriched gluten-free bakery products (Taylor and Parker 2002).

Most of the gluten-free products present on the market are characterized by a lower quality than conventional products made from gluten-containing flours. Hence, the combination of gluten-free flours and the choice of their ratios is essential for the acceptability of the final products (De Angelis and Di Cagno 2010). Currently, the most utilized gluten-free flours include maize, potato and rice flour and starches, used as base flours due to their bland flavour and neutral effects on baked products. These flours and starches usually tend to be low in nutrition and have very minimal structure-building potential. Chestnut flour presents high potential in the development of gluten-free products (O'Shea et al. 2014).

Gluten-free fermented products

The use of gluten-free flours represents a valid alternative to the complete absence of cereals in the diet of CS patients. However, the poor baking performance of these flours (due to the lack of gluten), the low nutritional quality and the poor sensory characteristics of the resulting products determine that there is an important technological challenge to be faced (Wolter *et al.* 2014a). Furthermore, gluten-free products have a limited microbial shelf life (Gallagher 2009; Hager *et al.* 2011).

In the case of bread, the use of sourdough as a bioprocessing ingredient in gluten-free formulations may provide several positive characteristics in the final product. O'Shea *et al.* (2014) summarized the beneficial role of sourdough in gluten-free bread production as follows:

- Production of peptidase able to detoxify the peptides responsible for CS when using wheat and rye flours.
- Activity against mycotoxins produced from fungi found on maize, sorghum and millet.
- Decrease of pH for degradation of phytic acid.
- Extraction of bioactive compounds from the flour and release of biomolecules that are part of the LAB/yeast metabolism.
- · Elongation of shelf life.
- Enhancement of the flavour profile.
- Production of exopolysaccarides (EPS), proven to be prebiotic, that are useful to produce breads with a softer texture without the addition of hydrocolloid.

All the positive aspects of gluten-free breads observed with the use of sour-dough are due to LAB communities, but their action is strain dependent (Arendt *et al.* 2011).

Among LAB, *Lb. plantarum* is reported to be dominant in several gluten-free sourdoughs produced from amaranth, buckwheat, quinoa, rice and teff flour (Vogelmann *et al.* 2009; Moroni *et al.* 2011a). Strains of this species have been proven to improve the staling rate and crumb hardness of brown rice, buckwheat-based, gluten-free formulations with the addition of sourdough and an inoculum size of 10⁸ cfu/g (Moore *et al.* 2007). However, breads made from different gluten-free flours may exhibit an undesirable aroma (Hager *et al.* 2012), especially when compared to wheat sourdough breads, due to the absence or low intensity of a wheat bread–like note. In general, the characteristic odour attributes of gluten-free breads are pea-like with buckwheat, quinoa and teff flours, cooked potato–like with quinoa and teff flours, vomit-like with sorghum and teff flours and mouldy with buckwheat and quinoa flours (Wolter *et al.* 2014a). The odorants and the resulting undesirable notes cause a negative impact on the aroma quality of gluten-free breads.

Selected LAB strains are able to generate very specific volatile organic compounds in wheat sourdough (Settanni *et al.* 2013). Hence, the use of individual metabolic properties of LAB seems to be a promising approach to increase also the aroma quality of gluten-free breads, although the type of flour influences sourdough fermentation, affecting the availability of carbohydrates as primary fermentation substrates, nitrogen sources and growth factors such as vitamins, minerals and the buffering capacity (Hammes *et al.* 2005).

In order to include starter cultures in gluten-free sourdough, it is important to select the correct strain(s) for a given flour type (Rühmkorf *et al.* 2012a). To study the role of LAB, the effect of *Lb. plantarum* sourdough on gluten-free bread quality was evaluated using a composite recipe (Moore *et al.* 2008; Coda *et al.*

2010). This species has been used by Wolter *et al.* (2014a) to ferment single gluten-free (buckwheat, oat, quinoa, sorghum and teff) flours, in order to investigate the influence of the corresponding sourdoughs added for fermentation in the resulting breads. All sourdoughs decreased dough strength, resulting in softer doughs, reduced the staling rate for buckwheat and teff breads and increased the cell volume in sorghum and teff breads, but they did not prolong the shelf life and did not improve the aroma of breads.

Formation of EPS is a positive characteristic of sourdough LAB, since this feature influences the viscosity of sourdough (Vogel et al. 2002). Homopolysaccharides (HoPS) are generally applied to improve the structural characteristics of baked goods (Corsetti and Settanni 2007). Sourdough lactobacilli have not been found to produce heteropolysaccharides (HePS; Tieking and Gänzle 2005), which are mainly applied in fermented milk products (Laws and Marshall 2001). However, some authors (Galle et al. 2011) have assessed that the utilization of LAB strains producing HePS expands the variety of cultures and the diversity of polysaccharides for applications in gluten-free baking. For this purpose, two LAB, one Lb. casei and one Lactobacillus buchneri, were tested in sorghum sourdough and the resistance to deformation of the sorghum sourdough started with Lb. buchneri was registered at lower levels, due to the presence of HePS. Rühmkorf et al. (2012b) evaluated four EPS-producing LAB strains. One Lb. curvatus provided the best results, in terms of reduced bake loss, higher crumb moisture content and slower rate of staling, when used in a rice/buckwheat bread formulation (Rühmkorf et al. 2012b). Different strains of Lb. sanfranciscensis, Lb. curvatus, Lactobacillus reuteri, Lactobacillus animalis and We. cibaria did not increase the loaf volume of buckwheat, rice and sorghum sourdough breads (Galle et al. 2012; Rühmkorf et al. 2012b). Moroni et al. (2011b) reported that a multiple LAB strain starter culture comprising species Lb. brevis, Lactobacillus paralimentarius, Lb. plantarum and We. cibaria determined a small loaf volume of a buckwheat formulation as a result of a decrease in CO, production due to the sourdough inclusion.

From the works cited it emerged that, although the application of sourdough might not necessarily lead to improved bread quality, the effect of its inclusion in gluten-free breadmaking is strictly dependent on the flour matrix used (Wolter *et al.* 2014b).

Microbial strategies to reduce gluten content

Gluten-free breads are undoubtedly characterized by lower-quality properties than wheat breads. For this reason, several research groups have studied different strategies to reduce the toxic effects of wheat breads for CS patients. Shan *et al.* (2002) proposed oral therapy with a prolyl-endopeptidase produced by *Flavobacterium meningosepticum* that hydrolyses the 33-mer peptide,

reported as one of the most potent peptides involved in triggering CS. This enzyme has also been purified from *Myxococcus xanthus* (Gass *et al.* 2005), *Sphingomonas capsulata* (Shan *et al.* 2004) and *Lactobacillus helveticus* (Chen *et al.* 2003). Stepniak *et al.* (2006) proposed oral supplementation with the prolyl-endopeptidase from *Aspergillus niger* that is stable under gastric conditions. However, some authors found that different LAB can decrease the CS-inducing effects of gluten.

The most significant developments in this field date to the beginning of the 2000s. Di Cagno et al. (2002) demonstrated active hydrolysis of various prolinerich peptides by lactobacilli. Following this finding, sourdoughs were prepared from a mixture of wheat (30%) and gluten-free (oat, buckwheat and millet; 70%) flours inoculated with strains of Lb. alimentarius, Lb. brevis, Lb. sanfranciscensis and Lactobacillus hilgardii. After 24 hours of fermentation, the gliadin fractions of the resulting bread were almost completely hydrolysed and the product was tolerated by CS patients (Di Cagno et al. 2004). The probiotic preparation VSL#3 (VSL Pharmaceuticals, Gaithesburg, MD, USA) containing Streptococcus thermophilus, Lb. plantarum, Lactobacillus acidophilus, Lb. casei, Lactobacillus delbrueckii spp. bulgaricus, Bifidobacterium breve, Bifidobacterium longum and Bifidobacterium infantis was also found to hydrolyse gliadin polipeptides (De Angelis et al. 2006). The same behaviour was registered for Ent. faecalis isolated from fermented wheat doughs (M'hir et al. 2008) and Wieser et al. (2008) confirmed the degradation of gluten proteins during sourdough fermentation in the presence of lactobacilli and enterococci during the selection of glutendegrading LAB. Gerez et al. (2008) reported the functionality of LAB peptidase activities in the hydrolysis of gliadin-like fragments. In that study, none of the LAB strains alone could hydrolyse 57–89 α -gliadin peptide, while the combination of Lb. plantarum and Pe. pentosaceus strains led to the hydrolysis of 57% of the peptide in 8 hours.

Rizzello *et al.* (2007) showed that fermentation by selected sourdough lactobacilli and addition of fungal proteases decreased the residual concentration of gluten of wheat flour below the threshold level indicated by the Codex Alimentarius Commissions of the World Health Organization and the Food and Agricultural Organization for gluten-free foods. Thus, the application of this combined approach based on the activities of sourdough lactobacilli and fungal proteases allowed the production of baked goods made from wheat flour that were not toxic to patients with CS (Greco *et al.* 2011). M'hir *et al.* (2009) used this approach, including a pool of selected enterococci and fungal proteases. However, some limitations to the use of sourdough to reduce the intolerance of CD patients derive from the long fermentation time required for complete hydrolysis of the toxic peptides. Under such conditions, stability and dough resistance are decreased as a result of the disruption of the gluten network (Cabrera-Chávez and Calderón de la Barca 2010). Rizzello *et al.* (2014b) developed a protocol for the manufacture of a traditional wheat flour bread with an

intermediate content of gluten, enhanced digestibility, high bioavailability of free essential amino acids and high protein nutritional quality, but other studies are needed to obtain breads with complete degradation of gluten during flour fermentation and good structural and sensory features.

Thanks to the properties reported here, sourdough LAB cultures are also useful during processing of gluten-free flours because they can eliminate the risks of cross-contamination by gluten (Di Cagno *et al.* 2008).

Conclusion

Cereal-based foods are key components of the diet of several populations. This chapter has analysed the process of fermentation of different cereal flours carried out by different microorganisms, basically LAB and yeasts, whose activities are responsible for the desirable and typical characteristics of the final products. Sourdough technology is applied worldwide to produce breads and provides a useful strategy to solve the main problems related to special bread production. Fortified and/or gluten-free breads are made with mixtures of ingredients and flours that do not generally result in high-quality products. The use of sourdough might improve several quality characteristics, but the effect depends on the active strains. From this standpoint, the selection of an ad hoc starter culture is defining to improve the sensory notes of a given fermented cereal-based product. Several LAB showed the potential to hydrolyse the toxic peptides responsible for CS during long fermentation, but the complete suitability of sourdough for the production of gluten-free breads from wheat or other flours containing gluten is still under study, in order to guarantee the safety of consumers as well as to decrease the time necessary for production.

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