Review

Food-Borne Fungi Kingdom

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Food-borne fungi may be divided into two categories, beneficial and detrimental. This has provided endless fascination and intrigue to those who have studied fungi, especially foodborne fungi. However, fungi are among the most versatile and diverse organisms in their morphology, life cycle, and ecology and the situation has made it difficult to understand fungal biology from the perspective of the broader fields of evolution, ecology, genetics, and population biology. Nowadays the population biology has come of age during the molecular revolution and this has influenced the choice of approaches and tools for foodborne fungi.

Characterization of food-borne fungi

Fungi are eukaryotes feed by adsorbing organic compounds from their environment and are generally composed of hyphae with cell walls. The non-motile organisms are recognized to reproduce by spores with one or several cells.

They are generally grouped in 4 division as follows due to similarity of structure, ribosomal DNA, cell wall composition, and lysine synthesis pathway. Of these, Deuteromycetes, unlike the other groups, has been well characterized on the basis of asexual reproduction (1, 2).

- 1) Zygomycetes
- 2) Ascomycetes
- 3) Deuteromycetes
- 4) Basidiomycetes.

Deuteromycetes has been further classified in 23 genera including Acremoniu, Aspergillus, Botrti, Fusarium, Penicillium, Phoma, Paecilomyces, Scopulariopsis, Trichothecium, Trichoderma, and Wallemia (3).

Aspergillus

Aspergillus species are mainly described and keyed out in many literatures, regardless of beneficial or detrimental. Aspergilli have been classified into 16 species, as shown in Table 1 (3).

Colonies usually growing rapidly, white, yellow, yellow-brown, brown to black or shades of green, mostly consisting of a dense felt of erect conidiophores (Fig. 1).

The distinction of this species from aflatoxigenic A. flavus is often difficult because of the many intergrading strains. Beneficial A .oryzae isolates differ from A. flavus by lighter green colonies and larger, less ornamented conidia, which are often not uniform in size.

Several species have attracted attention as human and animal pathogens or because of their

${\bf Table \ 1.} \quad {\bf Keys \ to \ the \ genera \ of} \ {\it Aspergillus \ species.}$

MEA, malt extract agar; Czapek, Czapek Dox agar medium. Handbook of Microbiological Media (Atlas, R.M. ed.) CRC Press (1993).

1a. Colonies white, black or in yellow, brown or grey colours	
2a. Conidial heads white, often wet	candidus
3a. Conidial heads dark brown to black	niger
4a. Conidial heads columnar, often cinnamon-brown to pinkish-brown	terreus
 5a. Conidial heads olive to light brown; stipe brown. Hülle cells often produced	ustus
6a. Conidial heads pure yellow, conidia smooth to finely roughened	ochraceus
 7a. Conidia conspicuously ornamented with warts and tubercles, outer and inner wall can be distinguished	tamarii wentii
8a. Conidiophores typically brown, Hülle cells and <i>Emericella</i> teleomorph mostly present	nidulans
9a. Colonies on Czapek or MEA mostly restricted (colony diameter usually less than 1.5 cm within one week)	
 10a. Colonies variably coloured, conidial heads biseriate, sometimes Hülle cells present	
Yellow Eurotium teleomorph produced in old cultures or on low water activity media	
12a. Conidial heads yellow-green to dark yellow green	
Conidial heads predominantly uniseriate, conidia dark yellow green, conspicuously echinulate	parasiticus
14a. Conidia minutely echinulate, yellow green	
15a. Conidial heads columnar, vesicles broadly clavate, conidia rough to echinulate	fumigatus clavatus

ability to produce toxic metabolites. Others are important for their role in fermentation of oriental food products or industrial application in the production of organic acids or enzymes. The classification is mainly based on morphological characters (4, 5)

Fungal metabolites such as organic acids and enzymes are frequently employed in dairy, bakery and fermented food products, as well as in preservation. The fungi could also provide a direct source of protein. The beneficial forms may be purposely introduced or naturally present. while the detrimental ones appear as unwanted natural contaminants. In the last decades fungal contamination during processing and storage has attracted much attention, particularly because of the production of toxic metabolites. Although toxic fungi are ubiquitous and belong to the common contamination flora, their recognition is hampered by incomplete and often confusing literature.

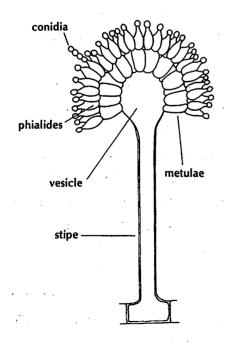


Fig.1. Typical morphological structure in Aspergillus

Aflatoxigenic Aspergilli

The profiles of secondary metabolites, including mycotoxins are of great use in systematics (4). A. flavus and A. parasiticus are well known for aflatoxigenic species (5). The presence of fungal growth on or in food products or in a Petri dish should be first confirmed directly with the naked eye and subsequently with a dissecting microscope. Fungal tissue (mycelium, fruit bodies, or sporulating structures) can be taken and subjected to examination and identification.

Useful roles of Aspergilli in food processing

(a) Production of organic acids

Citric acid is the most important organic acid produced by fermentation (5). Its annual production is estimated at 400,000 tons which are made mainly with A. niger (6). Citric acid is extensively used in the food industry as an acidualnt and flavoring substance.

Gluconic acid (50,000 tons/year) and its deltalactone are applied in foods and in the medical field.

Itaconic acid can be made with *A. terreus* using molasses or wood hydrolysates, and finds application in the chemical industry.

(b) Detoxification of mycotoxins

Aflatoxin B1 in some kinds of nut meals could be fully degraded during 7 days solid-substrate fermentation with a selected *Aspergillus* (7). Although the processes may be time consuming, biological detoxification of mycotoxins has the advantage over chemical and physical processes.

(c) Production of industrial enzymes

Within the range of enzymes produced by fermentation, the majority is hydrolytic enzymes,

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especially proteases. Proteases obtained from Aspergillus oryzae are applied in detergents, and in food processing (cheese ripening, bread making, and tenderization of meat).

Lipases synthesized by Aspergillus sp. are similar to pancreatic lipase in that they do not attack the 2-position in the triglyceride and show low specificity for the type of fatty acid whereas Mucor miechei lipase is active on short chain triglycerides with no marked positions specificity.

Starch-hydrolysing enzymes (alpha-amylases and glucoamylases) from *A. oryzae* and *A. niger* are applied in brewing and bread making.

Other hydrolases such as cellulases by A. niger, and pectinases by Aspergillus sp.are applied to improve digestibility of fibrous foods, and filterability of fruit juices and beer.

Fungi tend to secrete transeliminase type of pectinases that are specific for pectin (polymethylgalacturonate lyases).

(d) Transformation by protoplast fusion technique

Although any gene transfer technique can be used to construct high yielding strains, many fungal groups are said to lack a suitable system (7). For this organisms protplast fusion may hold the answer. If the cell wall is removed from fungi using enzymes in an isotonic medium, spherical proplasts result. In the fused product, if the two target chromosomes are largely homologous, recombination will occur.

Protoplast fusion offers the unique opportunity of bringing together two complete genomes and a very high frequency of recombinants can be obtained. Recently, one particularly successful host-vector system concentrated on employing both nitrate rductase (niaD) - deletion mutant: A. oryzae as a host and a potential chimera plasmid

that directed highly effective promoter gene as a vector (8).

A homologous transformation system was developed which is applicable to analysis of the amylase gene promotor function in *A. oryzae* (9). The unique host-vector recombinant system may provide much of the impetus for recent developments in eukaryotic protein secretion and thus enable to the products to be made at such high levels that their recovery will become commercially feasible (10).

Finally, this unique system allows us to manufacture genetically modified organisms that might be accepted as more suitable food materials than those yielded by using bacterial host and plasmids bearing antibiotic-resistant genes, since they might cause undesirable state of systematic immunological responsiveness after their intake and digestion.

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