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# IDENTIFICATION OF VOLATILE COMPOUNDS PRODUCED BY Brevibacterium linens THAT INHIBIT MOLDS

ΒY

SAMUEL E. BEATTIE

A thesis submitted in partial fulfillment of the requirements for the degree Master of Science Major in Dairy Science South Dakota State University 1985

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## IDENTIFICATION OF VOLATILE COMPOUNDS PRODUCED BY Brevibacterium linens THAT INHIBIT MOLDS

This thesis is approved as a creditable and independent investigation by a candidate for the degree, Master of Science, and is acceptable for meeting the thesis requirements for this degree. Acceptance of this thesis does not imply that the conclusions reached by the candidate are necessarily the conclusions of the major department.

> George S. Torrey Thesis Advisor

Date

John G. Parsons Head, Dairy Science Dept. Date

Knowing others is wisdom; Knowing the self is enlightenment. Mastering others requires force; Mastering the self needs strength.

He who knowns he has enough is rich. Perseverance is a sign of will power. He who stays where he is endures. To die but not to perish is to be eternally present.

Lao Tsu, 600 B. C.

#### ACKNOWLEDGEMENTS

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SEB

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#### INTRODUCTION

Mold contamination of processed food and its raw ingredients is a major concern in agriculture. Although at one time mold contamination of cheese and other foods was considered to be of little consequence, discovery of carcinogenic and/or teratogenic toxins (73) produced by molds commonly associated with food has changed this innocuous and frequent inconvenience into a serious health consideration.

The presence of mold on or in cheese is a common occurence, indeed, many cheeses depend upon molds to provide characteristic cheese flavor and texture. Cheeses such as blue, Roquefort, Camembert, and Brie are ripened under controlled conditions by species of <u>Penicillium</u>.

While certain molds are desirable, adventitious molds are undesirable contaminants in manufacture of most cheeses. Long ripening periods at controlled temperatures and humidity favor the growth of adventitious molds on Cheddar and other hard cheeses. Control of molds in ripening rooms and throughout cheese manufacture is important. One mold colony can give rise to thousands of airborne spores which can contaminate many kilograms of cheese. Some of the molds isolated from cheese produce mycotoxins in the cheese (10, 11, 48); therefore, control of mold growth is of paramount importance for the cheese industry.

Surface-ripened cheeses are ripened by a smear of

bacteria and yeasts growing on the surface of the cheese. <u>Brevibacterium linens</u>, the predominant microorganism in the smear of surface-ripened cheeses, is responsible for the typical aroma and flavor of Limburger, Trappist, Brick, and similar cheeses (1, 5, 8, 41, 42, 43, 60, 84). The aroma of these cheeses is a well-known characteristic and is due to the proteolytic metabolism of microorganisms in the surface smear (1, 2, 13, 21, 22, 23, 24, 26, 81, 84, 85).

An interesting aspect of surface-ripened cheeses noted by Grecz et al. (29) is resistance to spoilage by bacteria or molds. These workers found that extracts of surface-ripened cheeses and of <u>B</u>. <u>linens</u> produced similar antimicrobial activity (30, 31). Grecz et al. did not examine the aroma of surface-ripened cheeses for antimicrobial activity. The aromas of surface-ripened cheeses contain a variety of compounds, several of these compounds are inhibitory to mold growth in environments other than cheese.

The purpose of research presented in this thesis was to evaluate volatile compounds produced by <u>B</u>. <u>linens</u> for antimycotic activity. A method was developed to quantify inhibition of mold growth and spore germination in the presence of <u>B</u>. <u>linens</u>. The identity of the inhibitory compound was determined. Furthermore, the concentration of the inhibitory compound necessary to inhibit spore germination was determined.

This thesis is arranged in an alternative style. The comprehensive literature review is followed by two journal articles which describe the majority of materials and methods. Results and discussion are also included in the journal articles. Materials and methods or results not presented in an article are placed in appendices as are statistical information and raw data. The conclusions derived from the project are included as a separate section.

#### LITERATURE REVIEW

The microorganism most responsible for ripening of surface-ripened cheeses is <u>Brevibacterium linens;</u> this microorganism has a significant role in development of flavor, aroma, and texture in Limburger, Trappist, and Leiderkranz cheeses. Knowledge of the microbial succession occurring on the surface of this type of cheese and how the ecology of the cheese relates to organoleptic properties of the cheese is necessary to understand the antimycotic phenomena exhibited by these cheeses. The presence of sulfur-containing compounds in the aroma of cheese and production of these compounds by the surface microflora provide insight to the antimycotic activity. Furthermore, examination of cheese spoilage and methods to prevent mold spoilage of cheese is desirable to understand the need for new antimycotic agents. Finally, this literature review would not be complete without an overview of surface-ripened cheese manufacture.

#### Brevibacterium linens

<u>Bacterium linens</u> was first described by Wolff (as cited in 9, 21) as a short gram-positive rod that produced a red-brown pigment and was responsible in part for the slime on surface-ripened cheeses. Steinfatt (as cited in 21) provided a more detailed description of the bacterium as a gram-positive, aerobic, nonsporeforming, nonmotile rod.

The bacterium liquefied gelatin and produced an alkaline reaction in milk after 10 days of incubation.

Breed (as cited in 70) in 1963 proposed changing the name of <u>Bacterium linens</u> to <u>Brevibacterium linens</u> to define more clearly the genus <u>Bacterium</u>. The seventh edition of <u>Bergey's Manual of Determinative Bacteriology</u> placed the genus <u>Brevibacterium</u> in the family <u>Brevibacteriaceae</u> and classified <u>B</u>. <u>linens</u> as the type species for the genus (9). <u>B</u>. <u>linens</u> was described as a gram-positive, short, unbranched, gram-positive rod that gave a positive catalase test, liquefied gelatin, and was salt tolerant.

The coryneforme morphology exhibited by <u>B</u>. <u>linens</u> was not described in the seventh edition of <u>Bergey's Manual</u> of <u>Determinative Bacteriology</u>; however, Schefferle (as cited in 21) had observed snapping cell division and palisade cell arrangements, typical of corynebacteria, in cultures of <u>B</u>. <u>linens</u>. Mulder and Antheunisse (58) examined 150 strains of <u>Arthrobacter</u>, a genus of bacteria with coryneforme morphology, isolated from soil, activated sludge and dairy products and compared these isolates with <u>B</u>. <u>linens</u> and bacteria of other genera. These workers found <u>Arthrobacter</u> strains isolated from soil and activated sludge quite similar to each other yet different from strains isolated from dairy products. Furthermore, some of the strains isolated from dairy products were probably identical to <u>B</u>. <u>linens</u>. <u>B</u>. <u>linens</u> and these dairy strains were similar in nutritional,

morphological, and gram-staining characteristics. The pleomorphic tendencies of isolates of <u>Arthrobacter</u> and <u>B</u>. <u>linens</u> were also noted. Pleomorphism was most pronounced as the cultures aged or when incubation was in minimal medium. <u>Arthrobacter globiformis</u> was found to be very similar to <u>B</u>. <u>linens</u>.

Mulder et al. (59) further compared <u>B</u>. <u>linens</u> and <u>Arthrobacter</u> strains isolated from cheese. Two types of coryneforme bacteria were isolated from cheese; one forming grey-white colonies, the other forming orange colonies. <u>B</u>. <u>linens</u> and the orange colony-forming bacteria were similar morphologically and nutritionally. Strains of both types were able to grow within 31 days at 30 C in media containing up to 15% sodium chloride. Both types, in agreement with studies on Limburger and similar cheeses (1, 2, 13, 21, 22, 23, 24, 26, 81, 84, 85), were proteolytic. Because of the similarities between the orange-colony forming bacteria and <u>B</u>. <u>linens</u>, Mulder et al. (59) concluded that the pigmented bacteria should not be placed in the genus <u>Arthrobacter</u>.

The similarities among strains of <u>Arthrobacter</u>, <u>B</u>. <u>linens</u>, and other coryneforme bacteria have been examined on the basis of 57 characteristics (16). <u>A</u>. <u>globiformis</u> and <u>B</u>. <u>linens</u> were found to be very similar; 84% of the characteristics were properties of both microorganisms. This prompted daSilva and Holt (16) to propose that the name of <u>B</u>. <u>linens</u> be changed to "<u>A</u>. <u>linens</u> (Wolff) daSilva and Holt"

and that other species in the genus <u>Brevibacterium</u> be examined for similarities to other genera. The eighth edition of <u>Bergey's Manual of Determinative Bacteriology</u> lists all species of brevibacteria as <u>species incertae sedis</u> (70), indicating <u>Brevibacterium</u> may not be a discrete genus and individual species may belong in other genera.

### Microbial ecology of surface-ripened cheese

The observed softening of surface-ripened cheese during ripening because of growth of <u>B</u>. <u>linens</u>, indicates that the organism produces proteolytic enzymes. Albert et al. (5) found that pure cultures of <u>B</u>. <u>linens</u> were proteolytic in skim milk. An increase in soluble nitrogen in cultures was noted for a period of up to 81 days after inoculation. These workers also found an alkaline reaction in litmus milk with slight digestion of the medium observed 10 days post-inoculation and complete digestion of the medium in several weeks. The cultures were not fermentative.

The presence of yeasts on the surface of young surface-ripened cheeses was found to be important in the normal development of the cheese (36, 39, 40, 42, 67, 81, 84). Kelly (39), using microscope slide pressings from the surface of Limburger cheese, followed the succession of microorganisms over time. Initially, the dominant group of microorganisms occurring on the surface of the cheese was yeasts. Coincident with lysis and subsequent disappearance

of yeasts was appearance of short gram-positive rods. Kelly and Marquardt (40) found that the surface pH of the cheese increased from 4.9 to 5.8 in the eight days that yeasts were present. <u>B</u>. <u>linens</u> was found to be incapable of growth at pH lower than 5.8, and was not able to grow on fresh cheese curd having a low pH. These authors concluded that yeasts were able to metabolize lactic acid and proteins, raising the pH of the cheese to 5.8 and above, thereby providing conditions suitable for growth of B. linens.

While increasing the pH of the cheese surface aided growth of <u>B</u>. <u>linens</u>, yeasts also provided necessary growth cofactors (21, 36, 67, 84). Purko et al. (67) found that pure cultures of yeasts isolated from Limburger cheese were able to produce pantothenic acid, riboflavin, biotin and niacin. These yeasts were able to increase the growth of <u>B</u>. <u>linens</u> in a vitamin-free medium. Yeasts isolated from surface-ripened cheeses were identified as species of <u>Mycoderma</u> and <u>Torula</u> (21, 81, 84). Later work by Cone and coworkers (2, 81, 85) identified the major proteolytic yeasts occurring on surface-ripened cheese as species of <u>Trichosporon</u>.

Proteolytic activity of yeasts isolated from Limburger and Brick cheese appeared to be of minor nature with the major portion of proteolysis having occurred in the cheese after the yeasts had been replaced by bacteria or in 8-10 days (21, 39, 42). Amounts of free amino acids increased as the proteolytic bacteria digested the cheese. Methionine,

lysine, and glutamic acid were liberated in the greatest quantities from the proteins. The largest increase in free amino acids was found between 9 and 14 days post salting of the cheese (21).

Yeasts associated with Trappist cheese produced endocellular proteinases, polypeptidases, and peptidases and were important in increasing free amino acids in the cheese (2, 81, 85). Cheese ripened with yeast and B. linens had a higher free amino acid content through 35 days of ripening than cheese ripened with B. linens alone. An extracellular aminopeptidase from B. linens was isolated and characterized by Foissy (22, 23, 24). The pH optimum for the enzyme was 9.6, somewhat higher than the pH at the surface of the cheese during ripening (23). Torgeson and Sorghaug (82) found five enzymes that were able to hydrolyze a variety of dipeptides. An additional enzyme was found that was very active in peptide hydrolysis. The authors conclude that this enzyme was possibly a proteinase rather than a pepti-Lysis of cells during starvation or for other reasons dase. would release these enzymes into the cheese furthering proteolysis. An extracellular proteinase identified by Freidman et al. (26) was specific for casein, especially  $\beta$ -casein, and had a pH optimum between 7.2 and 7.3. Release of proteolytic enzymes by B. linens progressively degrades cheese proteins, increasing soluble nitrogen with age as peptides are formed (21). Textural and flavor changes occur with

enzymatic degradation of casein.

Importance of sulfur-containing compounds in aroma of cheeses

The aroma and flavor of a cheese is dependent upon many factors, most important being the components of the cheese and the microbial growth on or in the cheese (1, 60, 69, 88). Compounds important in cheese aroma are produced as by-products of microbial growth or are produced by chemical reactions facilitated by conditions created by bacterial growth. The protein and fat portions of cheese are substrates which microorganisms use readily to produce important and characteristic aroma compounds. For example, the aroma of blue cheese varieties is derived from the presence of methylketones (1, 62). Methylketones are catabolic endproducts of lipolytic molds associated with blue cheeses (1, 62).

Enzymatic degradation of cheese proteins by microbial enzymes yield compounds which have been found to be important in the aroma of surface-ripened and Cheddar cheeses (32, 33, 44, 49). Early work on the aroma of Cheddar cheese detected sulfur-containing compounds. Removal of these sulfur compounds by chemical traps altered the aroma of the cheese (47, 50, 52, 53, 86). Dateo et al. (17) devised a gas entrainment system for volatile sulfur compounds produced by cooking cabbage. This entrainment system found use in studying cheese aromas. The entrainment system

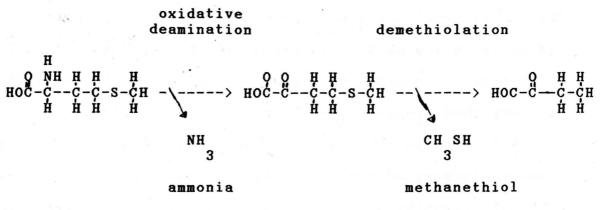
was composed aqueous solutions of lead acetate, mercuric cyanide, and mercuric chloride (Table 1). Walker (86) used cadmium sulfate in place of lead acetate to remove hydrogen sulfide and thiols in the presence of moisture. These trapping compounds coupled with aroma or taste panels were useful in determining which sulfur compounds were the most important constituents of Cheddar cheese aroma. The aroma was most altered after passing through trapping compounds that removed thiols and hydrogen sulfide (47, 50, 52, 53, 86).

Methanethiol, the most important constituent in the aroma of Cheddar cheese (52, 53), and other volatile sulfur compounds are derived from sulfur-containing amino acids present in cheese proteins (38, 43, 52, 53). Formation of methanethiol from methionine by adventitious and starter bacteria is not a major source of methanethiol in Cheddar The starter cultures employed in Cheddar cheese cheese. manufacture do not have this capablity and conditions in the cheese during aging do not favor growth of adventitious microorganisms (44, 74, 75). Law and Sharpe (44) hypothesized that the reducing environment produced by the starter culture favors nonenzymatic production of methanethiol from low molecular weight precursors produced by the starter culture. Furthermore, cheese made with gluconic acid lactone did not contain methanethiol and had a more positive redox potential (44, 50, 51).

Trap	Concentration of trapping agent	Compounds trapped
Mercuric chloride	3% (W/V)	Thiols, sulfides, disulfides
Mercuric cyanide	4% (W/V)	Thiols, hydrogen sulfide
Cadmium sulfide	5% (W/V)	Hydrogen sulfide

Table 1. Chemical traps employed to remove sulfur-containing compounds form atmospheres in growth chambers.

Methionine is necessary for production of methanethiol by pseudomonads and corynebacteria associated with cheeses (1, 33, 38, 44, 72, 74). Segal and Starkey (72) proposed that methionine is degraded to methanethiol by the following method:



methionine

🎗 -ketomethionine

*a*-ketobutryic acid.

Deamination and demethiolation are enzyme-mediated reactions. Some bacteria oxidize methanethiol to produce methyl disulfide and methyl sulfide which are also important in cheese aroma (38, 72). Methanethiol production from me-35thionine labelled with S demonstrated that the amino acid was demethiolated by the bacterium associated with Trappist cheese, i. e. <u>B</u>. <u>linens</u> (38). Grill et al. (33) used an entrainment similar to that of Dateo et al. (17) and trapped sulfur-containing compounds derived from the degradation of isotopically marked methionine. In traps specific for hydrogen sulfide little radioactive material was found when compared to traps specific for methanethiol. Grill et al. (33) hypothesized that sulfur-containing compounds other than methionine were precursors to hydrogen sulfide. Kadota and Ishida (38) reviewed several precursors to hydrogen sulfide production by bacteria. Cysteine and cystine, both present in casein and other proteins of milk and cheeses, are common precursors of hydrogen sulfide. Cuer et al. (15) further demonstrated that hydrogen sulfide was produced by <u>B. linens</u> when incubated in media enriched with cysteine.

Small concentrations of methanethiol, hydrogen sulfide, and methyl disulfide are found in a variety of food products. Although these compounds may be present in small quantities their presence is often quite evident. The odor thresholds of these sulfur compounds are low: methanethiol, 0.02 ppb; hydrogen sulfide, 0.18 ppb; dimethyl sulfide, 0.33 ppb (6, 61). Jansen found that the taste threshold for these compounds in beer ranged from 1.0 ppm for methanethiol to 10.0 ppm for methyl disulfide (37). Taste and odor thresholds are different; odor thresholds are approximately 10,000 times lower than taste thresholds (6).

More recent studies on the aroma of surface-ripened cheeses have relied upon more sensitive and accurate separation techniques using mass spectrometry to identify compounds (19, 20, 64). Vacuum distillation of extracts from several surface-ripened cheeses yielded over sixty neutral compounds (19) and in further work on Pont l'Eveque, a surface-ripened cheese, 123 compounds were identified (20). 414136

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Methanethiol, methyl disulfide, and thioesters were identified (20). This was the first evidence of the production of thioesters by bacteria associated with cheeses. S-methylthioacetate, thought to be formed by enzymatic acylation of methanethiol, has a low odor threshold, 5 ppb, and an odor similar to that of cooking cauliflower (1, 64). Thioesters are important in off flavor development in beers and other fermented beverages (37, 54, 55), and may be significant constituents in the aroma of surface-ripened cheese (19, 20, 64).

#### Spoilage and preservation of cheese

Manufacturing milk into cheese is an efficient method of concentrating milk constituents and preserving them from microbial spoilage. Many microorganisms, pathogenic and nonpathogenic, find milk a source of readily available nutrients, thus preventing long term storage of milk. Cheese has several characteristics which make it less susceptible to spoilage by microorganisms; (a) low pH because of acid production by starter cultures (41, 60); (b) salt concentration (1 to 2%) inhibitory to growth of spoilage organisms; (c) reduced oxidation/reduction potential making growth difficult for aerobic microorganisms; (d) inhibitory compounds produced by starter cultures (10, 35).

Cheese will, however, spoil if proper precautions

are not taken during manufacture and storage. Spoilage of cheese may vary from a slight flavor defect that is detectable only by experienced cheese graders to gross defects such as holes in the cheese or bloated packages caused by gas-producing microorganisms (13, 41, 60, 69, 88). Preventing spoilage of cheese by adventitious microorganisms is best accomplished through maintaining sanitary conditions during cheese manufacturing and using an active starter culture.

Perhaps the major type of spoilage found with cheese is surface mold growth. Species of <u>Penicillium</u> and <u>Asper-</u> <u>gillus</u> are the most common molds found on Cheddar and Swiss cheese (10, 11). Unlike many bacteria, molds can grow at low pH and temperature causing serious problems during aging and storage of cheese. The amount of cheese discarded or downgraded because of mold growth has not been estimated.

Mold contamination of foods is aesthetically unpleasant, and, with the discovery of mycotoxin production by molds commonly associated with foods, mold contamination has been recognized as a serious health hazard. Mycotoxins are produced by a variety of molds, including species of <u>Asper-</u> <u>gillus</u>, <u>Penicillium</u>, <u>Alternaria</u>, and <u>Fusarium</u> (34, 79); all of these molds have been isolated from cheeses (10, 11). Perhaps the most economically important mycotoxin-producing molds are species of <u>Aspergillus</u> especially <u>A</u>. <u>parasiticus</u> and <u>A</u>. <u>flayus</u> which produce aflatoxins (10, 11, 34, 73).

Aflatoxins are among the most carcinogenic and toxic compounds yet identified (73). Somewhat less toxic are toxins produced by species of <u>Penicillium</u>; these toxins include penicillic acid, ochratoxin A, and patulin (66, 79).

The production of mycotoxins by molds depends upon the substrate on which the organism is growing (34, 66). Molds isolated from cheeses are able to produce a host of toxins including aflatoxins, patulin, and penicillic acid (10, 11). Aflatoxins can be produced by <u>A</u>. <u>flavus</u> and <u>A</u>. <u>parasiticus</u> growing on cheese, but more importantly, aflatoxins can diffuse into the cheese to a depth where simply removing the visible mold may not remove all of the mycotoxin (25, 48, 76). This feature of mold contamination has serious health considerations.

Cheese is a unique food because of the long aging time many cheeses require before use. Cheeses made from unpasteurized milk are required by law to have at least 60 days aging to reduce health risks inherent in consuming raw milk. The long aging period necessitates use of antimycotic agents to reduce surface mold growth.

The most commonly used antimycotic agents for cheese preservation have been salts of sorbic acid (2, 4-hexadienoic acid) (68, 77). Potassium sorbate is used because sorbic acid, a polyunsaturated organic acid, is sparingly soluble in water. At 20 C, only 0.16 g sorbic acid will dissolve in 100 ml water, whereas 139 g potassium sorbate

will dissolve in 100 ml water (12). Organic acids, e. g. sorbic, propionic, and lactic acids, are most effective as antimicrobial agents in the undissociated form. Sorbic acid is most effective in low acid foods (12, 77). Sofos and Busta (77) state that sorbates are effective against molds at levels of 0.05-0.3%. Both potassium sorbate and sorbic acid are generally recognized as safe (GRAS) by the Food and Drug Administration (FDA) (68).

Application of sorbic acid to cheese may be by several methods including spraying or dipping, dusting the cheese surface with sorbic acid powder, or impregnating the compound into the packaging material (77). Sorbic acid has a strong odor and flavor that may impart an off-flavor to cheese. Moreover, sorbic acid can be metabolized by species of <u>Penicillium</u> to yield a compound that will impart a hydrocarbon off-flavor to the cheese.

Another organic acid which has been used as a cheese preservative is propionic acid (12, 18). As with other organic acids, propionic acid and salts of propionic acid are most effective against molds in the undissociated form, therefore, propionic acid is most effective in foods of low pH (18). By law, the upper limit for direct addition of propionic acid to cheese and cheese products is 0.3% (18); propionate may also be added to the packaging material. Propionic acid and its salts are GRAS. The compound has a strong odor and flavor and is an important component in the

aroma of Swiss cheese (12, 18).

In 1982 the FDA approved limited use of natamycin (previously known as pimaricin), an antibiotic produced by <u>Streptomyces natalensis</u>, as an antimycotic on cheese cuts and slices (28, 68) The use of natamycin on Italian and natural cheeses, including Swiss, Cheddar, and blue was approved in 1984 (28). The antibiotic was proven to be effective against mold growth at concentrations as low as 50 ppm, however, natamycin does not affect bacteria (28, 68). FDA regulations allow spraying or dipping cheese in aqueous solutions containing 200-300 ppm natamycin (14). Natamycin may be incorporated into the packaging material either as a constituent of packaging film (0.1% natamycin) or as an edible coating material containing 0.2% natamycin.

### Antimycotic agents produced by bacteria

Certain volatile sulfur compounds are known antimycotic agents (3, 27, 45, 46, 56, 57, 63, 65, 80). The ecology of various environments, including cheeses, is influenced by the presence of bacteria producing antimycotic agents during the decomposition of a substrate. Microorganisms may also produce nonvolatile antimicrobial substances such as antibiotics and toxins which aid the microorganism in competing for available nutrients.

The antimicrobial activity of surface-ripened cheese was first noted by Grecz et al. (29). In a series of experiments these workers demonstrated that growth of <u>Clostridium</u>

botulinum and species of Penicillium was inhibited when these organisms were inoculated onto sections of surfaceripened cheese. Inhibition of molds occurred on cheese at least eight weeks old that had been heated at 121 C for 15 min. Cheeses aged less than eight weeks were less inhibitory to growth of molds and <u>C</u>. <u>botulinum</u> (29). Extracts of the cheese were further examined for antimicrobial activity. Growth of C. botulinum, Staphylococcus aureus, and <u>Bacillus</u> cereus was inhibited by extracts of Leiderkranz cheese (30, 31). Finally, cultures of microorganisms isolated from surface-ripened cheese were examined for antimicrobial activity against B. cereus and C. botulinum. Antimicrobial activity was produced by mixed cultures only when cultures were aerated, and pure cultures of B. linens were found to exhibit antimicrobial activity (30). Because of the similarities in characteristics, the antimicrobial agent isolated from cheese and the antimicrobial agent isolated from <u>B. linens</u> were thought to be related antibiotics (31).

Weckbach and Marth demonstrated a reduction in toxin production by <u>A</u>. <u>parasiticus</u> when grown in a competetive environment with <u>B</u>. <u>linens</u> (87). Culturing <u>A</u>. <u>parasiticus</u> in the presence of <u>B</u>. <u>linens</u> reduced growth of mycelium, although the bacterium could no longer be isolated from the culture after three days of contact with the mold. The authors concluded that aflatoxins were toxic to <u>B</u>. <u>linens</u>.

Production of an antibiotic by brevibacteria associated with environments other than cheese has been demonstrated (4, 71). Brevibacteria isolated from human skin were able to produce an antibiotic effective against §. <u>aureus</u> (4). Ryall et al. (71), using synthetic skin medium, showed that brevibacteria were able to inhibit growth of fungi associated with <u>tinea pedis</u> (athlete's foot). These workers hypothesized that methanethiol produced by the bacteria was responsible for the observed inhibition. Proteolysis of proteins in the synthetic skin by enzymes produced by the bacteria released methionine which was then degraded to methanethiol.

Methanethiol and other sulfur compounds are produced during ripening of Cheddar and surface-ripened cheeses. These sulfur compounds may be found during degradation of substances other than cheese. Cabbage, kale, and Brussels sprouts (crucifers) release methanethiol, methyl disulfide, and methyl sulfide during microbial degradation in soil (45). The presence of crucifer stem and leaf tissue in soils reduced the amount of pea root rot caused by <u>Aphano=</u> <u>myces euteiches</u>. Further studies using pure sulfur-containing compounds, including methanethiol, methyl disulfide, and methyl sulfide, replicated inhibition of <u>A</u>. <u>euteiches</u> zoospore germination, hyphal growth, and zoospore motility (46, 63). Methanethiol when added to the headspace of growth jars completely inhibited the growth of the fungi at

a level of 10 to 30 ppm methanethiol.

Sulfur is an important part of many fungicides and exhibits fungicidal activity in the elemental form (3, 95). The reactivity of the sulfhydryl group is important for the biological activity of thiols and some sulfhydryl-containing fungicides, although the exact mode or site of toxicity is unknown. Several possible reactions leading to thiol toxicity in biological systems were outlined by Torgeson (83):

- Interactions with disulfide bonds in enzymes, thereby preventing the function of the enzyme. Thiols could also covalently bond with sulfhydryl groups in enzymes which rely upon a sulfhydryl group to function, e. g. acetyl Coenzyme A.
- 2. Chelating metal ions in cytochrome and other metallo-enzymes necessary for respiration. Hydrogen sulfide and short chain thiols are toxic to humans, part of this toxicity is due to the chelation of the metal ion in cytochrome oxidase (78).
- 3. Production of free radicals in the cell.

The presence of sulfur-containing compounds, such as methanethiol and hydrogen sulfide, in the aroma of surfaceripened cheese is well documented (1, 8, 13, 19, 20, 32, 33, 38, 44, 47, 49, 50, 51, 52, 60, 75, 86). While these compounds are toxic to humans in high concentrations, surface-ripened cheeses do not contain concentrations sufficient to be harmful to humans. The resistance to molding exhibited by surface-ripened cheese may be in part because of toxic concentrations of sulfur-containing compounds produced by microorganisms, especially <u>B</u>. <u>linens</u>, responsible for ripening of this type of cheese.

#### Surface-ripened cheese production

Production of surface-ripened cheese is similar to that of other cheeses up to the point of whey draining and matting of the cheese curd (41, 60). Unlike Cheddar and other hard cheeses, most surface-ripened cheeses are salted after pressing by placing in a brine maintained at 22% sodium chloride. This method of salting also inhibits growth of undesirable microorganisms. Halophilic yeasts, present in the brine, are inadvertently inoculated onto blocks during handling and brining. These yeasts are important in providing conditions suitable for bacterial growth.

Typically, surface-ripened cheeses are produced in small-sized blocks, not larger than five pounds and as small as four ounces (41, 60). Limburger and other surfaceripened cheeses rely upon microorganisms growing on the cheese surface for flavor development. Small blocks with an actively growing surface microflora are desirable so that enzymes produced by the surface microflora can diffuse throughout the cheese block. In contrast to surface-ripened cheeses, hard cheeses develop flavor uniformly throughout,

therefore, large blocks of cheese can be manufactured.

Surface-ripened cheeses are ripened in closely controlled conditions of temperature and high relative humidio ty; 15.6 C and 95% relative humidity are typically used (41, 60). The cheese blocks are placed on wooden shelves inoculated with <u>B</u>. <u>linens</u>. Turning the blocks and brushing them with water or saline solution create a uniform microbial smear on the surface of the cheese. The first stage of ripening may last up to two weeks depending upon the amount of flavor desired in the finished product. After the initial ripening period, the surface-ripened block is wrapped in parchment paper and curing continues for up to eight weeks before the cheese is marketed.

The finished product should be soft bodied with a uniform smear on the outside and a nutty, smooth taste. Overripened cheese will have an aroma and flavor reminiscent of ammonia because of extensive proteolysis, which also yields an overly soft cheese (41).

#### Summary

Surface-ripened cheeses are those cheeses ripened by a microbial smear growing on the cheese surface. Limburger, Trappist, Leiderkranz, and Brick are examples of this type of cheese. Halophilic yeasts and bacteria are the dominant microorganisms found in the smear. Yeasts appear first upon the cheese and are responsible for creating conditions suitable for bacterial growth. <u>Brevibacterium linens</u> becomes

the dominant bacteria in the smear and is responsible for development of the typical aroma and texture of surfaceripened cheeses. This bacterium may also be responsible for the resistance to molding exhibited by these cheeses.

Aromas of surface-ripened cheeses are complex mixtures. Catabolism of proteins by <u>B</u>. <u>linens</u> is an important source of many of aroma compounds. Sulfur-containing compounds, such as methanethiol, hydrogen sulfide, and thiolesters, are especially important components of surfaceripened cheese aroma. These compounds are known to possess fungicidal activity, and may play a role in antimycotic activity exhibited by surface-ripened cheeses.

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Volatile compounds produced by <u>Brevibacterium linens</u> inhibit spore germination and growth of several food-borne molds.

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Running head:

Volatile compounds from <u>B</u>. <u>linens</u> inhibit mold growth.

Volatile compounds produced by <u>Brevibacterium linens</u> inhibit spore germination and growth of several food-borne molds.

## Abstract

Volatile compounds produced by <u>Brevibacterium linens ATCC</u> 8377 and 9172 inhibited germination of spores and mycelial growth of <u>Aspergillus parasiticus</u> NRRL 2999, <u>A. clavatus</u> NRRL 6, <u>Penicillium camemberti</u> NRRL 973, and <u>P. expansum</u> NRRL 877. Molds and bacteria were grown on separate agar surfaces in the same enclosed atmosphere using growth chambers constructed of flexible film. A defined medium was used to determine the influence of methionine upon the production of antimycotic, volatile compounds by <u>B. linens</u>. Medium lacking methionine did not inhibit spore germination or hyphal extension.

The presence of <u>Brevibacterium linens</u> in the bacterial smear of surface-ripened cheeses has been correlated with the pungent aroma associated with these cheeses (7, 11, 16). Methanethiol, methyl disulfide, hydrogen sulfide and most recently methylthioacetate have been detected in the aroma of surface-ripened cheeses (7, 8, 17). During ripening of surface-ripened cheeses, amino acids from milk proteins are metabolized by the microflora of the surface smear to form these sulfur-containing compounds. Especially of note is the production of methanethiol (5, 11) and subsequent formation of thioesters (4, 5) by strains of B.

## linens.

Volatile compounds produced by bacteria have been shown to inhibit the growth of molds (9, 10, 15) and mycotoxin production (2). Ryall et al. (19) demonstrated inhibition of dermatophytic fungi by brevibacteria grown upon a proteinaceous medium, composed of synthetic skin and postulated that methanethiol was the responsible antifungal agent.

<u>B. linens</u> growing upon a cheese generates volatile compounds, including sulfur-containing compounds (7, 8, 16, 17). This study was undertaken to evaluate the ability of volatile compounds produced by <u>B</u>. <u>linens</u> to inhibit spore germination and hyphal growth of molds. The influence of methionine in the bacterial growth medium was also examined with regard to germination and hyphal growth.

### MATERIALS AND METHODS

Media and organisms. Cultures of <u>Brevibacterium linens</u> ATCC 8377 and 9172 were obtained from the American Type Culture Collection, Rockville, MD. Stock cultures were maintained on trypticase yeast extract agar (TYE) which contained (per liter): trypticase (BBL Microbiology Systems, Cockeysville, MD), 10 g; yeast extract, 5 g; agar, 15 g (both Difco Laboratories, Detroit, MI). <u>B</u>. <u>linens</u> was subcultured bimonthly onto fresh TYE slants and stored at 4 C.

Aspergillus parasiticus NRRL 2999, Penicillium camem-

<u>berti</u> NRRL 877 and <u>P. expansum</u> NRRL 877 were obtained from the Northern Regional Research Center, Peoria, IL. <u>A. cla-</u> <u>vatus</u> NRRL 6 was obtained from the Plant Science Department, South Dakota State University. All molds were maintained on malt agar (Difco) supplemented with 3 g yeast extract (MYE) per liter. Molds were subcultured bimonthly onto fresh MYE slants and stored at 25 C.

A defined medium (DMM), which was also used to culture <u>B</u>. <u>linens</u>, contained buffered salts solution described by Meynell and Meynell (14); amino acids (125 ug/ml), alanine, aspartic acid, glutamic acid, lysine, tryptophan, and methionine (all Sigma Chemical Co., St. Louis, MO); and vitamins (per liter); biotin, 5 ug, folic acid, 50 ug, and thiamine, 1.0 mg (all Sigma). Methionine was deleted from the medium (DMA) for some experiments. When required, media were solidified with Noble agar (15 g/l). The pH of each defined medium was adjusted to 7.0.

Spore germination studies. An adaptation of the slide-germination method of the American Phytopathological Society (1) was used to evaluate the influence of volatile compounds produced by strains of <u>B</u>. <u>linens</u> on germination of mold spores. Suspensions of spores were prepared by rinsing sporulated cultures with cold, sterile distilled water. The resulting spore suspension was passed through cheesecloth to remove mycelial fragments and diluted to contain 5000 to 10,000 spores/ml. Slide cultures were prepared by placing a

drop of molten MYE onto a sterile microscope slide in the cover of an inverted petri plate and transferring 0.01 ml of the diluted spore suspension, about 100 spores, onto the solidified agar.

Bacterial inocula were prepared in a chemically defined medium lacking an organic sulfur source (DMA). Cultures were incubated aerobically at 30 °C for 12-18 h. Petri plates containing TYE, DMM, or DMA agar were inoculated by spreading 0.5 ml of the broth culture of <u>B</u>. <u>linens</u> on each agar surface. Inoculated plates were incubated for 24 h at o 30 °C before preparation of growth chambers.

Growth chambers were prepared by combining covers of inverted petri plates containing freshly-prepared slide cultures of spores with the bottoms of petri plates that contained confluent, 24 h surface growth of <u>B</u>. <u>linens</u> on the desired agar medium. Combined petri plates were placed in flexible pouches composed of polyester/polyethylene laminate (Dazey Corp., Industrial Airport, KS), and the pouches were heat-sealed to form growth chambers (Fig. 1). Growth chamo bers were incubated at 25 C.

Inhibition of spore germination by volatile sulfur compounds produced by <u>B</u>. <u>linens</u> was evaluated using trapping agents to remove specific compounds from the atmospheres of growth chambers. Aqueous solutions of mercuric chloride, 3%(w/v), mercuric cyanide, 4% (w/v), and cadmium sulfate 5%(w/v in 0.2 N HCl) were used as trapping agents (3, 6, 13, 20).

Two leaves of 55 mm filter paper (Whatman No. 1), containing a single trapping solution were used as traps. Traps were placed under the slide culture and only one trapping agent was used in each growth chamber. Distilled water replaced trapping solutions in growth chambers containing no traps and control growth chambers.

Growth chambers were incubated at 25 C. Control chambers without bacteria and containing slide cultures were opened for evaluation after 24 h. Growth chambers containing <u>B. linens</u> along with slide cultures were opened after 48 h incubation. Spore germination was determined by means of a microscope, a spore was considered germinated only if a germ tube was apparent. Germination of spores in growth chambers was recorded as a percentage of the average number of spores germinating in control growth chambers. Hyphal extension studies. Slide cultures of spores were prepared described for spore germination studies. Mold cultures were incubated for 12-18 h in the absence of bacteria to allow spore germination. As in germination studies, slide cultures were then placed in inverted petri plates containing confluent surface growth of 24-h cultures of B. linens on TYE, DMM, or DMA agar. These combined plates were sealed into growth chambers and incubated at 25 C. Control chambers contained only slide cultures. Half of the growth chambers containing molds and bacteria were opened after 12 h incubation (Treatment 1), hyphal measure-

ments were made, and slide cultures and bacteria were resealed into chambers. This procedure was repeated at 12-h intervals for the duration of the experiment. The remaining growth chambers were incubated for 24 h (Treatment 2), chambers were opened, bacteria were removed, and hyphal measurements were made at 12-h intervals to determine if growth resumed. Hyphal extension measurements were made by means of a microscope fitted with a calibrated ocular micrometer. Measurements were made at a magnification of 100x and the largest colony diameter was measured. Measurements were made on five randomly selected propagules per chamber, six chambers were used each time, treatment, and medium for a total of 30 measurements.

Statistical interpretation. Spore germination percentages 0.5 were transformed using arcsin(%germination ) and then analysed by analysis of variance, each treatment was replicated eight times. Duncan's multiple range test was used to determine differences between treatments. Analysis of covariance was used to interpret hyphal extension data. Significance in all analysis was considered to be at P <

#### RESULTS AND DISCUSSION

Preliminary studies showed that the trapping agents.

used in this study did not reduce spore germination when  $\underline{B}$ . <u>linens</u> was absent. Germination of mold spores was inhibited when spores were incubated in the same atmosphere as B. linens ATCC 9172 and 8377 (Table 1). A pungent thiol-like aroma was apparent upon opening growth chambers containing only bacteria and slide culture (No Trap, Table 1) when TYE was used as growth medium for <u>B</u>. <u>linens</u>. Spores of all species of molds tested did not germinate in trapless growth chambers. Trapping agents included in growth chambers increased spore germination and reduced the pungent aroma compared to trapless chambers. Cadmium sulfate traps were superior to traps containing mercuric chloride or mercuric cyanide for improving the percentage of spores germinating compared to trapless chambers. When mercuric cyanide traps were used in growth chambers, the percentage of spores of <u>A</u>. parasiticus germinating was increased; however, an increase in the percentage of spores germinating was not noted for <u>A</u>. <u>clavatus</u>, <u>P. expansum</u>, or <u>P. camemberti</u>. The percentage of spores of <u>P. camemberti</u> germinating increased in growth chambers containing B. linens and mercuric chloride traps. Spore germination was also improved when P. expansum and B. linens ATCC 8377 were included in growth chambers containing mercuric chloride traps. Mercuric chloride traps did not increase the percentage of spores geminating for the molds used in the study. Other researchers (6, 11, 12, 13, 20) have used mercuric cyanide to remove hydrogen sulfide and

thiols, and, mercuric chloride to remove sulfides, disulfides, and thiols from atmospheres of several environments. Cadmium sulfate was used by Walker (20) to remove hydrogen sulfide from cheese aroma. As used in this study, the reaction of the trapping agents may have released volatiles which were inhibitory to spore germination. For example, reaction of thiols with mercuric cyanide forms hydrogen cyanide (3). These reaction products may cause the differential responce of mold spores to the presence of mercuric chloride or mercuric cyanide and B. linens.

A defined medium lacking organic sulfur, DMA, was used to determine the influence of methionine upon production of antimycotic compounds by <u>B</u>. <u>linens</u> (Table 2). Germination of mold spores was inhibited when methionine was included in the bacterial growth medium (DMM); however, when growth chambers contained bacteria grown upon DMA, mold spore germination was not inhibited. Comparison of spore germination in chambers containing <u>B</u>. <u>linens</u> grown upon DMA with spore germination in chambers containing <u>B</u>. <u>linens</u> grown upon DMM indicates that methionine was necessary for inhibition.

Extension of mold hyphae was also inhibited by volatile compounds produced by <u>B</u>. <u>linens</u> grown upon TYE or DMM (Fig. 2-5). Hyphal growth in the presence of <u>B</u>. <u>linens</u> grown upon DMA (data not shown) was similar to hyphal growth in control chambers. The bacteria were removed from growth

chambers (Treatment 2, Fig. 2-5) after 24 h to determine if inhibition of hyphal growth was permanent. When treatment two was used, inhibition of hyphal growth was apparent for all molds except P. <u>expansum</u> grown in the presence of B. <u>linens</u> grown upon DMM (Fig. 5). For all other molds, B. <u>linens</u> ATCC 9172 tended to restrict growth of hyphae more than B. <u>linens</u> ATCC 8377 whern the bacteria were grown on DMM. <u>B. linens</u> grown upon TYE inhibited hyphal extension regardless of the treatment or mold. Furthermore, when TYE was used as the bacterial growth medium hyphal lengths were shorter compared to those in growth chambers with <u>B. linens</u> grown upon DMM, albeit inhibition was observed in chambers containing either medium.

Morphological changes of mold hyphae occurred when incubated in the presence of <u>B</u>. <u>linens</u> grown upon TYE or DMM, (Fig. 5). Hyphae became swollen and highly vacuolated after 48 h of incubation in the presence of <u>B</u>. <u>linens</u> (Fig. 5a). In some cases, hyphae appeared to rupture with loss of cytoplasm (Fig. 5b). Hyphae in growth chambers with <u>B</u>. <u>linens</u> grown upon DMA did not exhibit morphological changes (Fig. 5c). Mold spores also exhibited morphological changes when incubated in the presence of <u>B</u>. <u>linens</u> grown upon DMM and TYE. Spores became smaller and appeared to become desicated. Pezet and Pont (18) observed morphogenetic changes in <u>Phomopsis vitacola</u> when the fungi produced methanethiol from a medium containing methionine. Methanethiol is also known to cause alterations in the life cycle of <u>Aphanomyces euteiches</u> (12). Methanethiol is known to be produced by <u>B</u>. <u>linens</u> during ripening of surface-ripened cheeses (7, 8, 18).

The pungent aroma noted upon opening growth chambers containing <u>B</u>. <u>linens</u> grown upon TYE or DMM was garlic-like and reminiscent of organic sulfur compounds. Methanethiol, methyl disulfide, and other sulfur-containing compounds are produced by B. linens during aging of surface-ripened cheeses (7, 8, 17). When growth chambers containing <u>B</u>. linens grown upon DMM or DMA were opened only DMM chambers had a pungent aroma. Since the presence of methionine was the only difference between these media it is concluded that B. linens produced a volatile metabolite from methionine that was capable of inhibiting mold growth and spore ger-The ability of B. linens to produce methanethiol mination. from methionine is well-documented (4, 5, 11) and we suggest that this is the compound responsible for inhibition. Furthermore, we suggest that production of volatile compounds such as methanethiol during aging of surface-ripened cheeses is important in suppressing mold spoilage.

# Acknowledgements

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	Strain of Number of s	Number of spores	pores Trapping Agent No Trap			
Mold	B. linens	in control <sup>a</sup>	HgC12	Hg(CN) <sub>2</sub>	; CdSO4	
A. parasiticus	9172	56 ± 1.6	0c	92.9 ± 3.7 <sup>b</sup>	$40.3 \pm 6.8^{d}$	0 <sup>c</sup>
	8377	77 ± 2.7	0c	83.1 ± 3.5	90.3 ± 4.9	0c
A. clavatus	9172	89 ± 5.4	0	0	91.8 ± 6.3 <sup>C</sup>	0
	8377	75 ± 2.8	3.3 ± 1.1	$2.0 \pm 0.7$	87.2 ± 3.6 <sup>C</sup>	2.7 ± 0.8
P. expansum	9172	109 ± 7.3	0	0	94.1 ± 9.2 <sup>c</sup>	0
	8377	88 ± 3.5	94.5 ± 3.1 <sup>C</sup>	0	95.6 ± 4.0 <sup>C</sup>	0
P. camemberti	9172	74 ± 2.8	83.7 ± 3.6 <sup>C</sup>	0	89.2 ± 3.5 <sup>c</sup>	0
	8377	95 ± 4.8	81.7 ± 4.2 <sup>d</sup>	0	$67.9 \pm 6.2^{\circ}$	0

TABLE 1. Percentage of mold spores germinating in the presence of <u>Brevibacterium</u> <u>linens</u> ATCC 8377 or 9172 grown upon TYE in the presence of trapping agents.

<sup>a</sup>Average number ± standard error of spores germinating in eight chambers not containing bacteria.

<sup>b</sup>Mean ± standard error of percentage of spores germinating in eight replicates.

 $c,d_{Values}$  with different superscripts within a row are different (P<.05).

TABLE 2. Percentage of mold spores germinating in the presence of <u>Brevibacterium linens</u> ATCC 8377 and 9172 grown upon DMA and DMM media.

	Strain of	Medi	Spores germinating, % Medium		
Mold	B. linens	DMM	DMA		
A. parasiticus	9172	0	89.4 ± 4.0 <sup>a</sup> ,b		
	8377	0	99.4 ± 1.3 <sup>b</sup>		
A. clavatus	9172	0	$97.0 \pm 5.7^{b}$		
ананананананананананананананананананан	8377	6.9 ± 1.4	94.2 ± 1.5 <sup>b</sup>		
P. expansum	9172	0	82.7 ± 7.2		
	8377	0	85.7 ± 4.3		
P. camemberti	9172	29.2 ± 3.1	90.3 $\pm$ 1.1 <sup>b</sup>		
· · · · · · · · · · · · · · · · · · ·	8377	0	90.2 ± 2.8 <sup>b</sup>		

Control = 100% germination.

<sup>a</sup>Mean ± standard error.

<sup>b</sup>Not significantly different from control (P<.05).

Figure 1. Hyphal extension of <u>Aspergillus parasiticus</u> NRRL 2999 in the presence of <u>Brevibacterium linens</u> ATCC 8377 and 9172 grown upon DMM (A) or TYE (B). Treatment 1 (+ = 8377, Δ = 9172) - bacteria remained in growth chambers for the entire experiment. Treatment 2 (□ = 8377, # = 9172) - bacteria removed from growth chambers after 24 h incubation. Control (X) growth chambers did not contain bacteria.

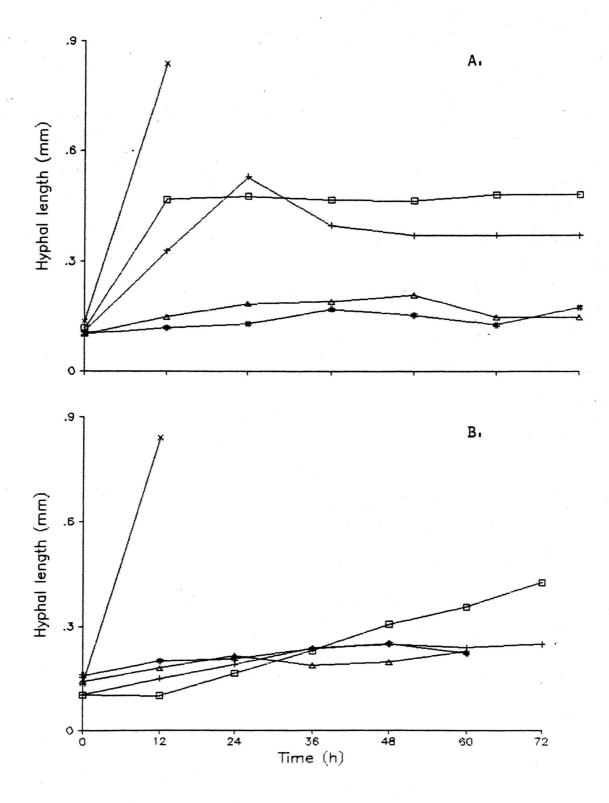


Figure 2. Hyphal extension of <u>Aspergillus clavatus</u> NRRL 6 in the presence of <u>Brevibacterium linens</u> ATCC 8377 and 9172 grown upon DMM (A) and TYE (B). Treatment 1 (+ = 8377, △ = 9172) - bacteria remained in growth chambers for the entire experiment. Treatment 2 (□ = 8377, # = 9172) - bacteria removed from growth chambers after 24 h incubation. Control (X) growth chambers did not contain bacteria.

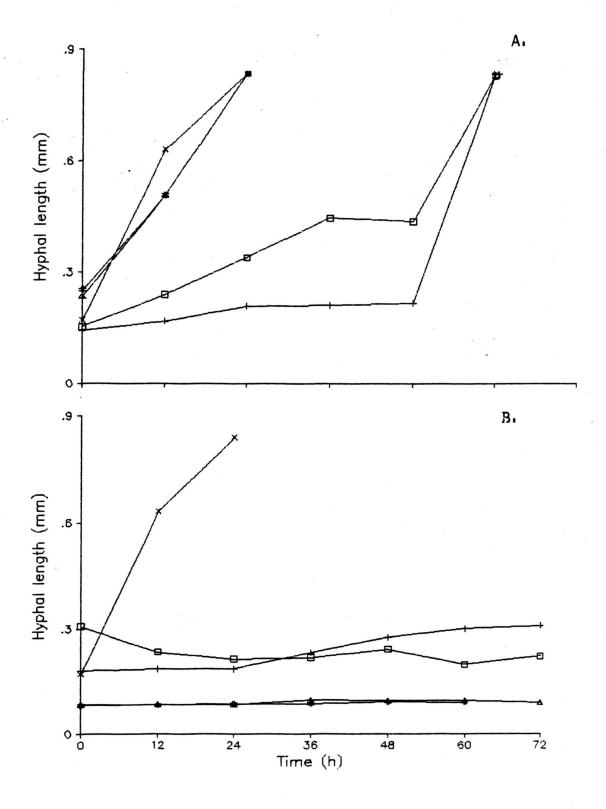


Figure 3. Hyphal extension of <u>Penicillium camemberti</u> NRLL 973 in the presence of <u>Brevibacterium linens</u> ATCC 8377 or 9172 grown upon DMM (A) or TYE (B). Treatment 1 (+ = 8377, △ = 9172) - bacteria remained in growth chambers for the entire experiment. Treatment 2 (□ = 8377, # = 9172) - bacteria removed form growth chambers after 24 h incubation. Control (X) growth chambers did not contain bacteria.

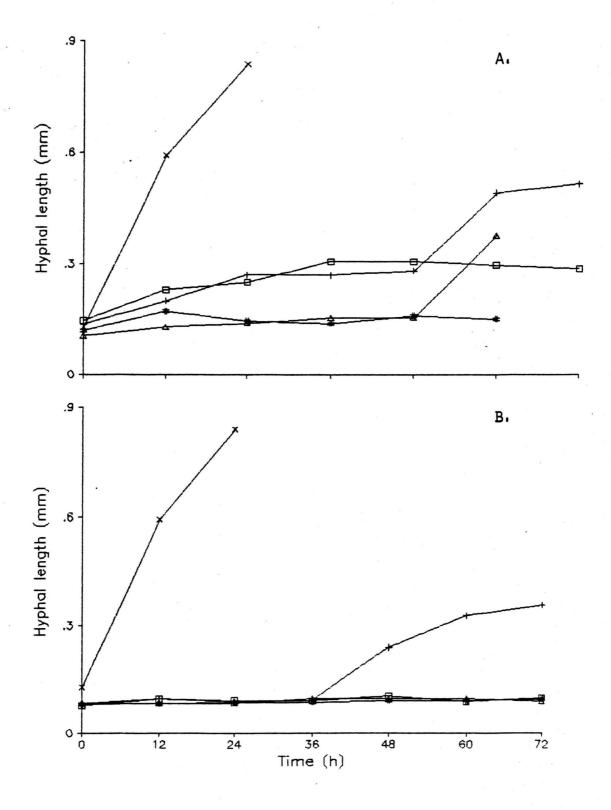


Figure 4. Hyphal extension of <u>Penicillium expansum</u> NRRL 877 in the presence of <u>Brevibacterium linens</u> ATCC 8377 and 9172 grown upon DMM (A) or TYE (B). Treatment 1 (+ = 8377, △ = 9172) - bacteria remained in growth chambers for the entire experiment. Treatment 2 (□ = 8377, # = 9172) - bacteria removed from growth chambers after 24 h incubation. Control (X) growth chambers did not contain bacteria.

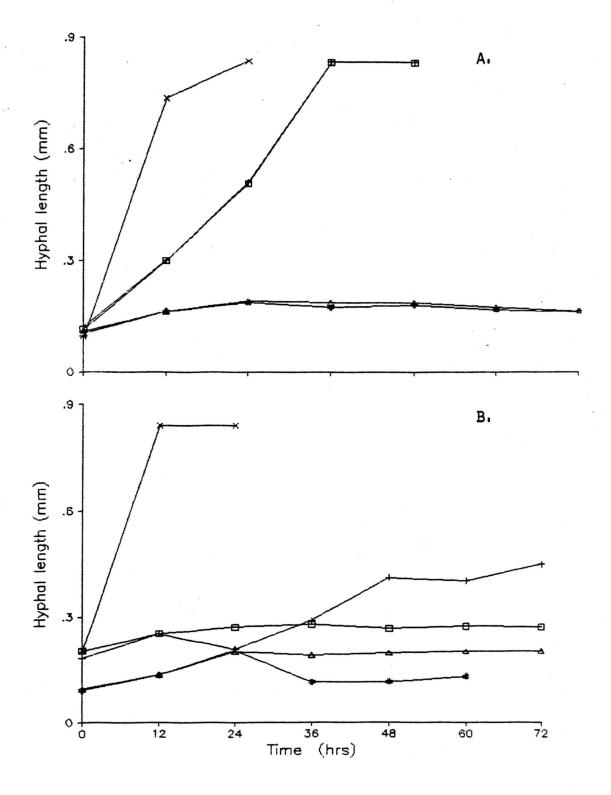
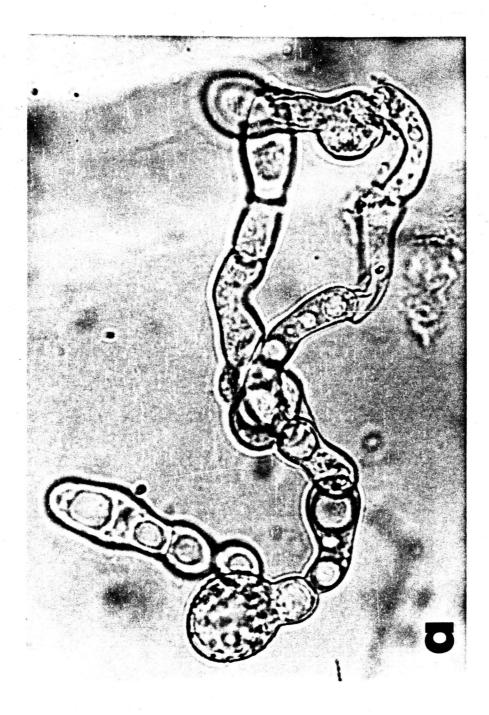
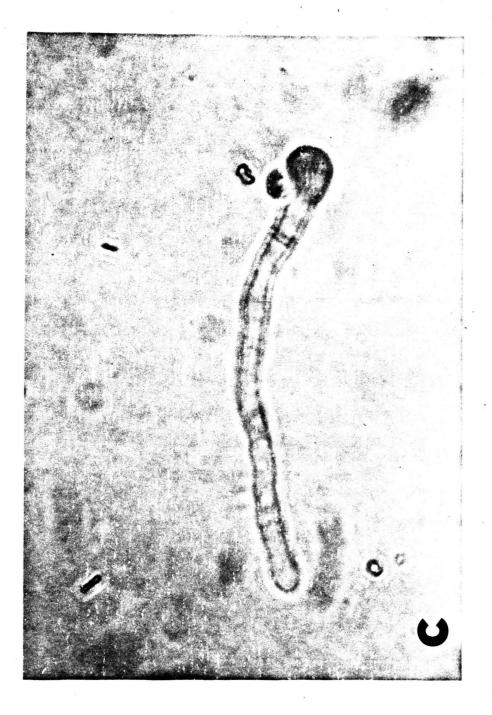


Figure 5.

Micrographs showing morphological changes in hyphae of molds caused by volatile compounds produced by Brevibacterium linens ATCC 8377 grown upon DMM (400 x). A - Aspergillus parasiticus, B - Penicillium camemberti,

C - Aspergillus parasiticus control.





Fungal toxicity of volatile compounds produced by <u>Brevibacterium linens</u> caused by methanethiol

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## Abstract

Methanethiol production was studied using cultures of <u>Brevibacterium linens</u> ATCC 8377 grown upon three different media. <u>B</u>. <u>linens</u> did not produce methanethiol when grown on a medium lacking methionine, but did produce methanethiol and methyl disulfide when grown on a defined medium containing methionine and an undefined medium containing trypticase. Methanethiol concentrations produced by <u>B</u>. <u>linens</u> are capable of inhibiting germination of spores of <u>Penicil-</u> <u>lium expansum</u> NRRL 877 (toxic concentrations being above 0.33 ppm methanethiol.

The characteristic aroma and flavor of surface-ripened cheeses result in part from the proteolytic activity of microorganisms found in the surface smear of these cheeses. <u>Brevibacterium linens</u>, the dominant bacterium of the smear, produces proteolytic enzymes and is responsible for the pungent aroma associated with these cheeses (Boyavel and Desmazeaud, 1983; Parliment et al., 1983). Grill et al. (1967) found that the surface smear of Trappist cheese was necessary for production of methanethiol from methionine. Methanethiol, hydrogen sulfide, and methylthioacetate are important sulfur-containing compounds in the aroma of surface-ripened cheeses (Dumont et al., 1974, 1976; Parliment

### et al., 1982).

A notable aspect of Limburger, Trappist, and other surface-ripened cheeses is their inability to support mold growth (Grecz et al., 1959; Shih and Marth, 1972). Grecz et al. (1961, 1962) investigated the possiblity that an antibiotic, produced by <u>B</u>. <u>linens</u>, was responsible for the resistance to molding. More recent work conducted with brevibacteria from human skin determined these bacteria produced an antibiotic active against gram-positive and -negative bacteria (Ryall et al., 1981; Al-Admany and Noble, 1981). Ryall et al. (1981) postulated that methanethiol produced by brevibacteria may inhibit the growth of dermatophytic fungi. Whether a constituent of surface-ripened cheese aroma is responsible for the resistance to molding exhibited by surface-ripened cheeses has not been examined.

Volatile compounds produced by <u>B</u>. <u>linens</u> have been demonstrated to inhibit irreversibly spore germination and mycelial growth of several mold species commonly reported on spoiled cheese (Beattie and Torrey, 1985), and inhibition of mold growth depended upon the presence of methionine in bacterial growth media. We report herein on the toxicity of methanethiol on spores of <u>Penicillium expansum</u>, the rate of production of methanethiol by <u>B</u>. <u>linens</u>, and use of polyester/polyethylene laminate pouches as controlled atmosphere growth chambers.

### MATERIALS AND METHODS

Organisms and culture conditions. <u>Brevibacterium linens</u> ATCC 8377 was obtained from the American Type Culture Collection, Rockville, MD, and <u>Penicillium expansum</u> NRRL 973 was obtained from the Northern Regional Research Center, Peoria, IL. Malt agar (Difco Laboratories, Detroit, MI) supplemented with 3 g/L yeast extract (MYE) was used for growth and maintenance of <u>P. expansum</u>. A medium (TYE) containing trypticase (BBL Microbiology Systems, Cockeysville, MD), 10 g/L; yeast extract, 5 g/L; and agar, 15 g/L (both Difco) was used for growth and maintenance of <u>B</u>. <u>linens</u>.

A defined medium (DMM), which was also used for bacterial growth, contained buffered mineral salts solution (Meynell and Meynell, 1970); amino acids (0.125 ug/mL each), alanine, aspartic acid, glutamic acid, lysine, methionine and tryptophan (all Sigma Chemical Co., St. Louis, MO); and vitamins (per L); biotin, 5 ug, folic acid, 50 ug, and thiamine, 1.0 mg (all Sigma). Methionine could be deleted from the medium (DMA) without affecting bacterial growth. When required, media were solidified with 15 g/L Noble agar (Difco). The pH of each media was 7.0.

Bacterial inocula were prepared in DMA broth; these o cultures were incubated aerobically at 30 C for 12-18 h. Petri plates containing TYE, DMM, or DMA agar were inoculated by spreading 0.5 mL of the broth culture of <u>B</u>. <u>linens</u> on the agar surface. Inoculated plates were placed in flexible pouches (16.2 cm x 24.4 cm), composed of polyester (0.5 mil)/polyethylene (1.5 mil) laminate (Dazey Corp., Industrial Airport, KS). A teflon/rubber septum coated with silicon high vacuum grease was sealed into a pocket on the interior wall of each pouch. Pouches were closed by means of a heat-sealer (Dazey Corp.) to form growth chambers. The positions of mold and bacterial cultures within growth chambers are illustrated in Figure 1. Samples of the atmospheres of growth chambers were taken through the septum by means of a gas-tight syringe. Growth chambers were incu-

Analyses of growth chamber atmospheres. Atmospheres samples were directly injected into a Varian 3700 gas-liquid chromatograph (Varian Instruments, Palo Alto, CA) equipped with a flame ionization detector, on-column injection, and a glass column (1.83 m x 0.63 cm o.d.) packed with 10% DC-200 on Chromosorb W(AW) as described by Leppanen et al. (1980). The detector and injection temperatures were, respectively, 200 and 180°C, and the oven was programmed (without delay) from 40 to 110°C at 8°C/min and equipped with flame ionization detector. Nitrogen was used as carrier gas at a flow rate of 30 mL/min. To determine the tentative identity of compounds occurring in atmospheres of growth chambers, retention times for peaks occurring in sample chromatograms were compared with retention times of peaks in chromatograms of standards. Methanethiol (Eastman Kodak, Rochester, NY), methyl disulfide (Aldrich Chemical Co., Milwaukee, WI), and methyl thioacetate (Alfa Products, Danvers, MA) were used as standards. Retention times of standards were .71 min, 3.36 min, and 2.27 min, respectively.

Stock mixtures of methanethiol in nitrogen gas were made in empty sealed pouches. Evacuated pouches were reinflated with nitrogen gas, and then injected, via a gas-tight syringe, with a known weight of liquid methanethiol. Dilutions were made at room temperature from this stock mixture into pouches containing nitrogen. Concentrations of methanethiol in growth chambers were determined from a standard curve correlating peak height with concentration. Spore germination assay. An adaptation of the slidegermination method of the American Phytopathological Society (1943) was used to evaluate the influence of methanethiol on germination of spores of P. expansum. A drop of MYE on a sterile microscope slide was inoculated with 50 to 100 mold spores; this slide culture was placed into an inverted petri Moistened filter papers were added to the petri plate. plate to provide adequate humidity. The slide culture and petri plate were then sealed into a growth chamber. Atmospheres in sealed growth chambers were removed by means of a gas-tight syringe. A known volume of air was injected into the collapsed growth chambers using a gas-tight syringe.

Methanethiol from stock mixtures was then injected into growth chambers to yield concentrations of 0.16, 0.33, 0.82, 1.6, and 3.3 ppm. Growth chambers with no methanethiol were used as controls.

Growth chambers were incubated at 25 C. Chambers containing 0, 0.16, and 0.33 ppm methanethiol were opened after 24 h incubation because outgrowth of spores was observed. Germinated spores were counted by means of a microscope. Remaining chambers were opened and examined after 48 h incubation.

### **RESULTS AND DISCUSSION**

We have observed previously (Beattie and Torrey, 1985) that mold spores representing four species did not germinate when incubated in growth chambers containing strains of <u>B</u>. <u>linens</u> grown upon TYE or a defined medium that contained methionine (DMM). Spores did germinate in chambers containing <u>B</u>. <u>linens</u> grown upon medium from which methionine was absent (DMA). These observations indicate that the mold is inhibited by a metabolite of methionine. Comparison of odors from growth chambers in which <u>B</u>. <u>linens</u> was incubated on DMM or DMA media indicated presence of a volatile sulfur compound in atmospheres over DMM but not over DMA.

Periodic gas chromatographic analyses were done on samples from atmospheres in growth chambers containing <u>B</u>. <u>linens</u> growing upon TYE, DMM, and DMA media. Methanethiol was not detected in atmospheres when <u>B</u>. <u>linens</u> was grown on DMA, a medium which lacked methionine. However, when samples from chambers containing <u>B</u>. <u>linens</u> grown on media including protein (TYE) or methionine (DMM) were chromatographed (Figure 1), methanethiol and methyl disulfide were detected but not methylthioacetate. Methyl disulfide, probably from oxidation of methanethiol (Parliment et al., 1982), was detected only in trace amounts at incubation times of 84 h and longer. Since these incubations were longer than the 48 h necessary to develop fungal toxicity in growth chambers (Table 1), and since methyl disulfide is less toxic to fungi than methanethiol (Owens, 1969; Lewis and Papavizas, 1971), these traces of methyl disulfide were considered unlikely to influence spore germination.

Concentrations of methanethiol in growth chambers were determined at intervals during incubation (Figure 2). Data from growth studies in this laboratory demonstrated that toxic concentrations of methanethiol occur in growth chambers after 24 to 36 h incubation on DMM and TYE media. Examination of Figure 2 reveals that methanethiol concentrations on both media during this period ranged from about 0.2 to 1.2 ppm.

Spores of <u>P</u>. <u>expansum</u> were used to correlate concentration of methanethiol with toxicity. A series of growth chambers containing slide cultures of spores were injected with varying concentrations of methanethiol, and the chambers then incubated. Spore germination was inhibited permanently by the presence of 0.82 but not 0.33 or 0.16 ppm methanethiol (Table 1).

Cuer et al. (1979) have reported methanethiol, hydrogen sulfide, methyl disulfide, and methylthioacetate in volatiles from strains of <u>B</u>. <u>linens</u> cultured on a medium containing protein. Methanethiol produced during decay of cruciferous plants inhibited fungal growth and zoospore germination (Lewis and Papavizas, 1970, 1971). Owens (1969) has suggested that reaction of methanethiol with sulfhydryl groups may cause inactivation of enzymes necessary for germination. We report herein the capability of methanethiol in concentrations less than 1 ppm to inhibit germination of spores of P. expansum. B. linens grown on DMM or TYE produced methanethiol in concentrations similar to those found toxic to the fungi. Production of methanethiol by B. linens during the manufacture of surface-ripened cheeses serves to discourage mold spoilage.

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This research was supported by funds from the South Dakota State University Graduate Research Fund, and the Dairy Research Foundation of the National Dairy Council. Portions of this research were presented at the annual meeting of the American Society for Microbiology, Las Vegas, NV, March 3-7, 1985. Figure 1. Diagram of flexible laminate growth chamber. A slide culture, B - growth chamber, C - bacterial culture, D - septum sealed into growth chamber wall to allow sampling of growth chamber atmosphere.



Figure 2. Chromatogram of atmosphere from growth chamber containing <u>Brevibacterium linens</u> ATCC 8377 grown upon DMM medium for 84 h. Attenuation 32x at 10<sup>-12</sup> sensitivity. A. Methanethiol. B. Methyl disulfide.

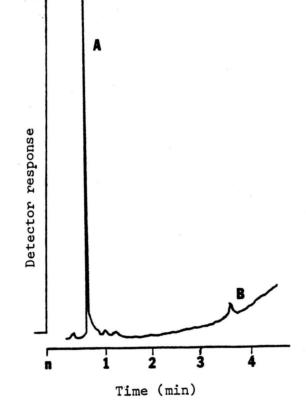


Figure 3. Methanethiol production by <u>Brevibacterium linens</u> ATCC 8377 grown upon DMM (•) and TYE (•) media, average of three replications.

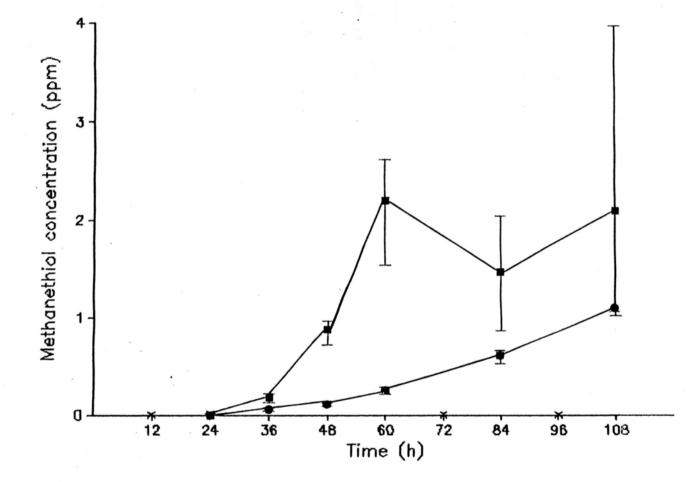


Table 1. Effect of methanethiol concentration on germination of spores of <u>Penicillium expansum</u> NRRL 973.

Methanethiol concentration	Number of spores germinated
0.0 <sup>a</sup>	107 <u>+</u> 5 <sup>b</sup>
0.16	107 <u>+</u> 6
0.33	86 <u>+</u> 46
0.82	0
1.6	0
3.3	0

<sup>a</sup>Control - no methanethiol added.

<sup>b</sup>Mean of three replicates <u>+</u> standard deviation

### SUMMARY OF CONCLUSIONS

Volatile compounds produced by <u>Brevibacterium linens</u> inhibited spore germination and hyphal growth of several molds. Inhibition of molds inside growth chambers in which molds and bacteria were grown on separate media was not observed when methionine was excluded from bacterial media. Incorporation of methionine in bacterial media caused inhibition of mold growth and resulted in production of methanethiol by <u>B</u>. <u>linens</u>. Methanethiol, in concentrations similar to those produced by <u>B</u>. <u>linens</u> in growth chambers was shown to inhibit germination of mold spores. Production of methanethiol in surface-ripened cheese may be responsible, in part, for the resistance to molding these cheese exhibit.

# Appendix A

# Development of defined medium

The ability of a medium to support bacterial growth can be determined by measuring an increase in cell concentration after a period of growth. The increase in biomass over a period of time can be described mathematically as dx/dt = ux (2, 9); where dx/dt is change in cell concentration, dx, over time interval, dt. x is the actual cell concentration at any given time, and, u is the specific -1 growth rate (h ). This formula is valid only during the exponential phase of growth, and for a given microorganism in a given medium under constant conditions, u is a constant and the slope of the exponential portion of a growth curve.

In dilute cultures, cell concentration and medium turbidity are directly related (2), therefore, the amount of light absorbed or transmitted through a liquid culture can be related to bacterial growth and actual cell concentration. The slope of the growth curve during exponential growth in a dilute culture is an indication of the ability of a medium to support bacterial growth.

Growth of <u>Brevibacterium linens</u> upon a complex medium such as cheese yields a variety of compounds, most notable are sulfur compounds which are important components of cheese aroma (1, 4). A defined medium was developed to determine the influence of organic sulfur upon the production of volatile sulfur compounds by <u>B</u>. <u>linens</u>. Growth curves were done to determine the influence of an added organic sulfur source upon the growth of <u>B</u>. <u>linens</u>.

# Materials and Methods

The media used for growth curves are given in Figure Al. Stock cultures of <u>B</u>. <u>linens</u> ATCC 8377 and 9172, obtained from the American Type Culture Collection, were streaked onto a medium (TRP) containing; trypticase, 1% (BBL Microbiology Systems, Cockeysville, Md), and agar, 1.5% (Difco Laboratories, Detroit, MI). Growth from an isolated colony on TRP was used to inoculate 25 ml of test medium. Flasks of inoculated test medium were placed upon an incubatorshaker (New Brunswick Scientific, Edison, NJ) and incubated at 30°C for 24 h with agitation (200 rpm). If a test medium did not become turbid after 24 h incubation it was discarded. Turbid cultures were gram stained to insure purity. These cultures were used as inocula for growth curves.

Growth curves were initiated by inoculating 50 ml of the test medium in a 300 ml side-arm flask. A 3% inoculum was used for all media except those containing 5 or 6 amino acids, these were inoculated at 10%. A spectrophotometer (Spectronic 20, Bausch and Lomb, Rochester, NY) was used to determine initial (time = 0) absorbance of media. Cuvettes containing uninoculated media were used as blanks. Inoculated side-arm flasks were incubated as described earlier and absorbance readings were made periodically.

## **Results and Discussion**

The goal of this research was to develop a chemically defined medium that would support growth of B. linens in the absence of an organic sulfur source. This medium, as developed or supplemented with methionine, was used to evaluate the production of antimycotic volatiles by B. linens. The medium selected contained five amino acids, three vitamins, and basal salts (DMA, Figure Al). When an organic sulfur source, methionine, (DMM) was added, growth of B. linens was not decreased (Figure A2). Specific growth rate constants of <u>B</u>. <u>linens</u> grown upon DMA and DMM were quite similar (Table Al). While growth of <u>B</u>. <u>linens</u> on DMA or DMM, as evidenced by smaller values for u (Table Al), was slower than on other media, the cell morphology remained constant. There is a tendency for B. linens and other corynebacteria to to become pleomorphic when media is nutritionally inadequate (10, 11).

Figure Al. Media used in development of a defined medium for the growth of <u>B</u>. <u>linens</u> ATCC 8377 and 9172.

Trypticase yeast extract (TYE): trypticase, 0.5%; yeast extract, 0.5%.

1% Trypticase (TRP): trypticase, 1%.

20 amino acids (20 AA):

Amino acids (125 ug/ml): cystine, tryptophan, aspartic acid, tyrosine, glutamic acid, arginine, proline, asparagine, threonine, isoleucine, methionine, phenylalanine, serine, glycine, alanine, valine, lysine, leucine, histidine, cysteine.

Vitamins: biotin, 0.005 ug/ml; folic acid, 0.05 ug/ml; thiamine, 1.0 ug/ml.

Basal salts solution (trace quantities): NaCl, KCl,

NH Cl, MgCl, KH PO, Na PO, FeCl in Tris 4 2 2 4 2 4 3 buffer, pH 7.0 (8).

ll amino acids (ll AA):

Amino acids (125 ug/ml): tyrosine, lysine, tryptophan, methionine, cysteine, proline, alanine, threonine, glutamic acid, aspartic acid, isoleucine. Vitamins: same as 20 AA medium.

Basal salts solution: same as 20 AA medium.

Continues,

Figure 1A. (continued) a 6 amino acids (DMA , DMM):

Amino acids (125 ug/ml): lysine, alanine, glutamic

acid, aspartic acid, tryptophan, methionine.

Vitamins and basal salts solution same as 20 AA medium.

Media brought to neutrality (pH 7.0) by addition of 0.1 N NaOH or HCl.

#### a

Methionine not added to DMA medium.

ledium		Growth constant, u* (h-)
TRP .	8377	. 134
	9172	.127
TYE	8377	.115
	9172	.168
20 AA	8377	.113
	9172	.157
11 AA	8377	.153
	9172	.154
DMM	8377	.041
	9172	.043
DMA	8377	.039
	9172	.038

Table Al. Specific growth rate constants of <u>B</u>. <u>linens</u> ATCC 8377 and 9172 grown at 30 oC upon various media.

\*

Growth constant derived from regression equation of the exponential portion of growth curve, f(x) = ux + b (2, 9).

Figure A2. Growth of <u>Brevibacterium</u> <u>linens</u> ATCC 9172 in six different media at  $30^{\circ}C$ .  $\Box$  - TRP,  $\blacksquare$  - TYE,  $\nabla$  - 20 aa,  $\bullet$  - 11 aa,  $\blacktriangle$  - DMA, O - DMM.

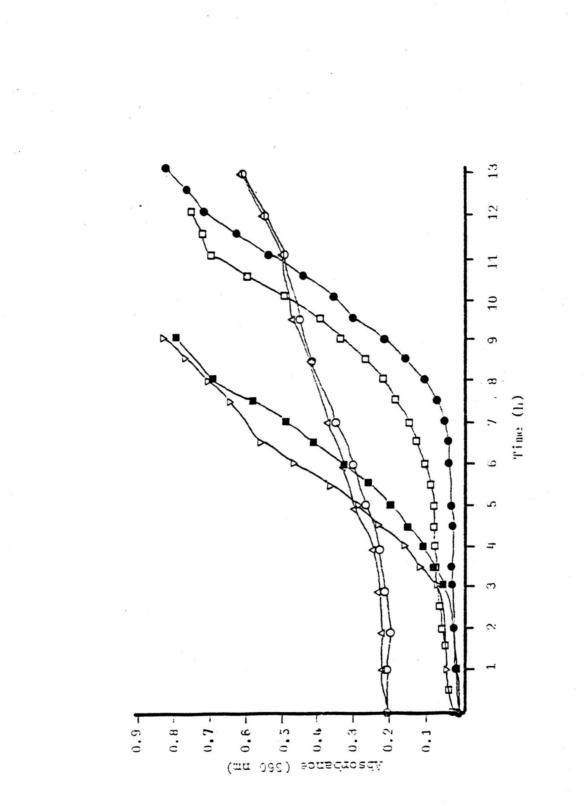
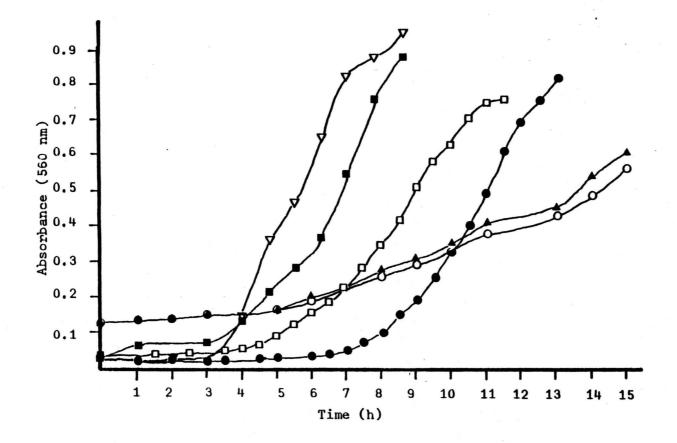


Figure A2. Growth of <u>Brevibacterium</u> <u>linens</u> ATCC 8377 in six different media at  $30^{\circ}C$ .  $\Box$  - TRP,  $\blacksquare$  - TYE,  $\nabla$  - 20 aa,  $\bullet$  - 11 aa,  $\Delta$  - DMA, O - DMM.



# Appendix B

Chemical methods: quantitation of volatile fatty acids, and efficiency of chemical traps for certain sulfur compounds Gas chromatography was done to determine if volatile fatty acids (VFA) were present in atmospheres of growth chambers and in broth cultures of <u>Brevibacterium linens</u>. Chemical trapping agents were used in growth chambers to remove specific sulfur-containing compounds (Table B1) from the atmospheres of growth chambers. The relative efficiency of these trapping agents in removing the specific sulfur-containing compounds was also determined by gas chromatography.

### Materials and methods

The chromatograph used was a Varian 1400 equipped with a flame-ionization detector and a stainless steel column (1.83 m x 0.32 cm) packed with 10% SP-1200, 1% H PO 3 4 on Chromosorb W(AW) 80/100 mesh (Supelco, Inc., Bellefonte, PA). The gas chromatograph was operated isothermally at 0 140 C.

Samples of growth chamber atmospheres and broth culture media were analysed for VFA. Broth cultures of <u>B</u>. <u>linens</u> were incubated for 24 h on an incubator/shaker and o for 24 h with agitation (200 rpm). Following incubation, cultures were acidified with 25% metaphosphoric acid (2 ml HPO /10 ml media) and centrifuged at 3 10,000 x g for 10 min to remove bacterial cells from culture media (12). A portion of the supernatant was injected into the chromatograph. Growth chamber atmospheres were sampled by means of a gas-tight syringe and the samples were injected directly into the chromatograph.

Retention times of peaks occurring in sample chromatograms were compared with retention times of peaks occurring in chromatograms of standards. The VFA standard solution contained the acids: acetic, 3.15 mg/ml; propionic, 0.993 mg/ml; butyric; 0.958 mg/ml; valeric, 0.47 mg/ml; and isovaleric, 0.431 mg/ml. Standards were run at chromatograph -11 settings of 10 sensitivity and 128x attenuation, samples -12 were run at 10 sensitivity and 32x attenuation.

Atmospheres of growth chambers containing <u>B</u>. <u>linens</u> and trapping agents were analysed by means of gas chromatography. A Varian 3700 gas chromatograph equipped with a flame-ionization detector and glass column (1.83 m x 0.63 cm) packed as described by Leppanen et al. (5) was used. The chromatograph was temperature programmed from 40 to 110 o o C at 8 C/min. Samples of atmospheres were injected directly into the gas chromatograph.

### Results and Discussion

Volatile fatty acids were not detected in the atmospheres of growth chambers, very small amounts of acetic, propionic, and isovaleric acids were found in supernatents of broth cultures. Hosono et al. (as cited in 1) found that <u>B</u>. <u>linens</u> produces organic acids when grown upon casein. Further work by Hosono et al. (as cited in 1) determined that acetic and isovaleric acids originate from alanine and leucine, respectively. Since no VFA were detected in samples of atmospheres of growth chambers, it is reasonable to conclude that these compounds do not play a role in the observed fungitoxicity of volatile compounds produced by <u>B</u>. <u>linens</u>. Volatile fatty acids found in broth cultures most likely remain in the salt form until acidification of the growth medium by metaphosporic acid.

The trapping agents used in this study (Figure B1) have been used by others (3, 4, 6, 7, 15) to study the effect of removing specific sulfur-containing compounds from cheese aroma and from atmospheres above decomposing cruciferous plants (6). Manning and Price (7) determined that traps composed of mercuric cyanide and mercuric chloride were up to 99% effective in trapping the sulfur compounds for which they were intended. Cadmium sulfate was used by Walker (15) to remove hydrogen sulfide in the presence of thiols and water vapor.

Gas chromatography of atmospheres of growth chambers containing trapping compounds and <u>B</u>. <u>linens</u> revealed that mercuric chloride and mercuric cyanide were very effective in trapping methanethiol. After incubation of <u>B</u>. <u>linens</u> in the presence of these trapping agents for 48 h, methanethiol was not detected in samples from growth chamber atmospheres. The aroma emitted from growth chambers was also changed, the putrid, sulfury aroma apparent in chambers containing only <u>B</u>. <u>linens</u> was completely eliminated by including traps of

mercuric cyanide or mercuric chloride in growth chambers. Chambers containing traps of cadmium sulfate were less effective in removing the typical aroma, and, gas chromatography of atmospheres revealed small concentrations of methanethiol, about 0.16 ppm. Table Bl. Chemical traps employed to remove sulfur-containing compounds from atmospheres in growth chambers.

 Trap	Concentration of trapping agent	Compounds trapped
Mercuric chloride	3% (w/v)	Thiols, sulfides, disulfides
Mercuric cyanide	4% (w/v)	Thiols, hydrogen sulfide
Cadmium sulfate	5% (w/v)	Hydrogen sulfide

Appendix C

Statistics and summary of raw data

Hyphal extension data for all molds with both strains of Brevibacterium linens have been summarized and mean values with standard deviations are given in Tables 1 to 8. Each value in these tables is the average from thirty measurements, five measurements taken from six slide cultures for each treatment. There were three treatments for each set of mold and bacterium, however, only treatments one and two are given in the following tables. Treatment three was a control containing no bacteria. Hyphal extension of molds in treatment three chambers proceeded at the same rate as hyphal extension in chambers containing B. inens grown upon DMA. Treatment one growth chambers contained B. linens and mold hyphae for the entire experiment. Treatment two growth chambers contained <u>B. linens</u> for only 24 hours after this period of incubation the slide culture was removed from the growth chamber and sealed into a chamber without the bacterium.

The remaining tables are summararies of statistics from Statistical Analysis System (SAS, 13) output. Hyphal extension data was evaluated using general linear model (GLM) procedure and deleting treatment three and DMA data. Spore germination data was evaluated using analysis of variance (ANOVA), differences between means of treatments of spore germination data were determined by Duncans multiplerange test with significance set at 0.05 level (13, 14).

						Medi	um					
		DM	A	-		DMM				TYE	- 14	
ſime (h)	Trt l	Sd	Trt 2	Sd	Trt l	Sd	Trt 2	Sd	Trt 1	sd	Trt 2	Sd
0	a .09	.03	.10	.03	. 10	.03	. 10	.03	. 14	.06	.16	. 05
12	. 84		. 84		ь		.12	.04	ь		. 20	.07
24	. 84		. 84		.19	.06	.13	.04	. 22	.10	. 21	. 09
36					. 19	.06	. 17	.06	. 19	.09	. 23	. 12
48					. 21	.07	.16	.05	. 20	.09	. 25	.13
60					.15	.05	.13	.05	. 23	.10	. 22	. 10
72					.15	.06	.18	.06	. 22	.06	. 25	. 11

Table Cl.	Aspergillus parasiticus	hyphal	extension	in	the	presence	of	₿.	linens	ATCC	8377	
	DMM. DMA. or TYE.							_				

a

						Med	ium					
		DM	A			DMM				TYE		
Time (h)	Trt 1	Sd	Trt 2	Sd	Trt l	Sd	Trt 2	Sd	Trt 1	Sd	Trt 2	Sd
0	a .14	. 04	. 12	. 04	.11	.05	. 12	.03	. 10	.03	.10	. 03
12	ь		. 82	.08		ь	. 47	.15	ь		.10	.02
24	. 84		. 84		.53	.16	.48	.13	.19	.10	.16	.08
36	. 84		. 84		.40	.15	. 47	.15	. 24	.14	.23	.12
48					. 37	.15	. 47	. 14	. 25	.15	. 31	.14
60					. 37	.16	.48	.15	. 24	.13	. 35	.14
72					. 37	.15	.48	.16	. 25	.13	. 42	. 25

Table C2. <u>Aspergillus parasiticus</u> hyphal extension in the presence of <u>B</u>. <u>linens</u> ATCC 9172 grown upon DMM, DMA, or TYE.

a

						Medi	um					
		DM	٨			DMM				TYB		
ſime (h)	Trt l	Sd	Trt 2	Sd	Trt l	Sd	Trt 2	Sd	Trt 1	Sd	Trt 2	Sd
0	.lla	.03	. 10	.03	. 12	.04	. 12	. 05	.18	.07	. 20	.06
12	ь		.53	.13	1	,	. 30	. 07		,	. 25	.08
24	. 84		. 84		.51	.10	.51	. 11	. 21	.08	. 27	. 07
36	. 84		. 84		. 84		. 84		.29	. 05	. 28	. 07
48					. 84		. 84		.41	.03	. 27	. 07
60									.40	.05	. 27	.06
72									. 45	.03	. 27	.08

Table C3. <u>Aspergillus clavatus</u> hyphal extension in the presence of <u>B</u>. <u>linens</u> ATCC 9172 grown upon DMM, DMA, or TYE.

a

					Medi	um					
		DMA			DMM				TYE		
Time (h)	Trt l S	d Trt 2	Sd	Trt l	Sd	Trt 2	Sd	Trt l	Sd	Trt 2	Sd
0	.14a .0	4.12	.04	.10	.03	. 11	. 02	.09	. 04	.09	. 03
12	b	. 84		1	•	.16	.04	ь		.13	.04
24	.84	84		.19	.05	.19	.05	. 20	.07	. 20	.09
36	.84	84		. 19	.03	.18	.04	. 19	.06	. 11	.05
48				.19	.04	.18	.04	. 20	.06	. 12	.05
60				.18	.04	. 17	.04	. 20	.04	.13	. 05
72				. 17	.04	.16	.04				

Table C4. <u>Aspergillus clavatus</u> hyphal extension in the presence of <u>B</u>. <u>linens</u> ATCC 8377 grown upon DMM, DMA, or TYE.

a

						Medi	um					~
		DM	A			DMM				TYE		
Time (h)	Trt l	Sd	Trt 2	Sd	Trt l	Sd	Trt 2	sd	Trt 1	Sd	Trt 2	Sd
0	.lla	.04	.13	.04	. 11	.03	.12	.05	.08	. 02	.08	.03
12	ь		. 50	.14	ь	,	. 17	. 09	ь	,	.08	. 02
24	. 84		. 84		. 14	.05	.15	.07	.08	.02	.09	.03
36	. 84		. 84		.16	.06	.14	.06	.10	. 02	.09	.02
48					.16	.05	.16	.16	.10	.02	.09	. 02
60					. 38	. 33	. 15	.06	.10	.02	.09	. 02
72					.42	. 36	. 15	.05	.09	.02	. 10	.03

Table C5. <u>Penicillium camemberti</u> hyphal extension in the presence of <u>B</u>. <u>linens</u> ATCC 8377 grown upon DMM, DMA, or TYB.

Mean and standard deviation (Sd) of 30 measurements, in millimeters, taken from six microscope slide cultures, five measurements/slide culture.

a

						Medi	um					
		DM	A			DMM				TYE		
lime (h)	Trt l	Sd	Trt 2	Sd	Trt l	Sd	Trt 2	Sd	Trt 1	Sd	Trt 2	sd
0	. 14a	.03	. 16	.05	. 14	.03	.15	. 04	.08	. 02	. 08	. 02
12	ь		. 83	.08	-1	•	.23	.07	ь		.10	. 02
24	. 84		. 84		. 27	.07	.25	.05	.09	.02	.09	. 02
36	. 84		. 84		. 27	. 05	. 31	.06	.09	.03	.09	. 02
48					. 28	.05	. 31	.06	. 24	.12	.10	. 02
60					. 49	. 21	. 30	.04	. 33	. 32	.09	. 02
72					. 52	. 20	. 29	.03	. 36	.36	. 10	.02

Table C6. <u>Penicillium camemberti</u> hyphal extension in the presence of <u>B</u>. <u>linens</u> ATCC 9172 grown upon DMM, DMA, or TYE.

a

			Medi	um		
	D	MA	DMM		TYE	
Time (h)	Trtl Sd	Trt 2 Sd	Trtl Sd	Trt 2 Sd	Trt l Sd	Trt 2 Sd
0	.18a .05	.20 .20	.24 .07	.25 .08	.08 .03	.08 .0
12	ь	.83 .08	ь	.51 .13	ь	.08 .0
24	. 84	.84	.84	.84	.08 .03	.09 .0
36	.84	.84	.84	.84	.10 .03	.09 .0
48					.10 .03	.09 .0
60					.10 .02	.09 .0
72					.09 .02	.10 .0

Table C7. <u>Penicillium expansum</u> hyphal extension in the presence of <u>B</u>. <u>linens</u> ATCC 8377 grown upon DMM, DMA, or TYB.

a

					Medi	um					
		DMA			DMM				TYE		
ſime (h)	Trt 1	Sd Trt 2	Sd	Trt l	Sd	Trt 2	Sd	Trt l	Sd	Trt 2	Sd
0	.17a .0	.17	.05	. 14	. 05	. 15	.06	. 18	. 10	.31	. 16
12	ь	. 57	.12	b	,	. 24	.06	ь		. 23	.16
24	. 84	84		. 21	.07	. 34	. 23	.19	. 10	. 21	. 15
36	. 84	84		. 21	.06	.45	. 29	. 23	.14	. 22	. 15
48				. 22	.05	. 44	. 30	. 27	. 20	. 24	. 15
60				. 84		. 84		. 30	. 26	. 20	. 15
72				. 84		. 84		. 31	. 25	. 22	. 15

Table C8. <u>Penicillium expansum</u> hyphal extension in the presence of <u>B</u>. <u>linens</u> ATCC 9172 grown upon DMM, DMA, or TYB.

a

Mold	Strain of <u>B. linens</u>	Standard error
aspergillus parasiticus	8377	. 12
	9172	.15
Aspergillus clavatus	8377	.05
	9172	.13
<u>Penicillium camemberti</u>	8377	.16
	9172	.13
<u>Penicillium expansum</u>	8377	.20
	9172	.09

Table C9. Standard error terms from analysis of variance of hyphal extension data.

Root mean square error - standard deviation of hyphae lengths over entire experiment, including all treatments, media, and times, and not including treatment three or DMA.

×

variance of mold spore gen of trapping agents and <u>B</u> . <u>linens</u> grown upon defined	rmination perce <u>linens</u> and in	entages in t	
Mold	<u>B</u> . <u>linens</u> strain	Stand	ard error
· · · ·		TAa	DMP
Aspergillus parasiticus	8377	8.90	11.30
	9172	7.82	10.69
Aspergillus clavatus	8377	6.86	20.24
	9172	10.96	11.74
<u>Penicillium camemberti</u>	8377	11.38	11.70
	9172	7.46	13.92
<u>Penicillium expansum</u>	8377	8.08	9.89
	9172	14.36	10.15

a

Root mean square errors from experiments using trapping agents.

bRoot mean square errors from experiments using defined media.

Table Cl0. Standard error terms of analysis of

95

Source of variation	Degrees of freedom	Mean squarea							
	Treedom	Penicillium camemberti	Penicillium expansum	Aspergillus parasiticus	Aspergillus clavatus				
Treatment	1	3.64*	605.51	997.24	19.29*				
Medium	1	6432.27	17809.17	14723.20	20.49*				
Time	5	20.56	417.45	354.93	558.17				
Slide(treat x time)	12	233.81	279.33	164.64	1102.29				
Model	20	737.77	1848.78	1688.72	1484.38				
Error	ь	28.03	29.92	41.46	45.76				
Total	ь								

Table Cll.	Analysis of variance mean squares for hyphal extension data using B. linens	
	DMA and treatment three deleted from analysis.	

Sum of squares/degrees of freedom (df).

P not significant at .01.

8

\*

b - Degrees of freedom: 669, P. <u>camemberti</u>; 609, P. <u>expansum</u>; 609, A. <u>parasiticus</u>; 639, A. <u>clavatus</u>. Total degrees of freedom = error df + model df

Source of variation	Degrees of freedom	Mean squarea			
	Treedom	Penicillium camemberti	Penicillium expansum	Aspergillus parasiticus	Aspergillus clayatus
Treatment	1	164.37	* 347.44	615.40	* 2.44
Medium	1	1578.79	12339.88	476.77	48.34
Treatment x medium	1	88.12	329.12*	268.97	109.38
Time	5	9.23*	1605.34	180.85	18.39
Slide(treat x time)	12	87.69	69.90*	906.20	18.73
Model	20	187.33	1675.40	978.61	82.56
Error	ь	12.59	70.35	23.96	4.07
Total	ь				

Table Cl2. Analysis of variance mean squares for hyphal extension data using <u>B</u>. <u>linens</u> ATCC 8377, DMA and treatment three deleted from analysis.

Sum of squares/degrees of freedom (df).

\*

a

P not significant at .01.

b - Degrees of freedom: 639, P. <u>camemberti</u>; 402, P. <u>expansum</u>; 509, <u>A. parasiticus</u>; 424, <u>A. clavatus</u>. Total degrees of freedom = error df + model df

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