

Partial Characterization of Nine Bacteriocins Produced by Lactic Acid Bacteria Isolated from Cold-Smoked Salmon with Activity against *Listeria monocytogenes*

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Nine LAB bacteriocin-producers, isolated from vacuum-packaged cold-smoked salmon (CSS), were phenotypically and genotypically identified as *Lactobacillus curvatus*, *Lactobacillus delbrueckii*, *Lactobacillus fermentum*, *Enterococcus faecium*, and *Pediorococcus acidilactici*. Their bacteriocins were partially characterized. The antimicrobial spectrum was determined against *Listeria monocytogenes*, *E. faecalis*, *E. faecium*, and *Staphylococcus aureus*. The molecular size of bacteriocins ranged from 2.8 to 4.5 kDa. They were inactivated by treatment with proteolytic enzymes but not by lipolytic or glycolytic enzymes. Maximal activity against *L. monocytogenes* ranged between 800 and 10000 AU/mL at pH 6.5. Most of the bacteriocins maintained full activity in a pH range of 2.0 to 8.0 but were partially or completely inactivated at pH 10.0. After heating at 60°C and 100°C, only two bacteriocins from *Lb. curvatus* strains partially lost activity. All bacteriocins showed a narrow spectrum of activity and a high anti-listerial activity, which is characteristic of the class IIa bacteriocins. Isolated bacteriocin-producing LAB could be used successfully in the bio-preservation of CSS and development of new potential bio-preservatives for CSS active against *L. monocytogenes*.

INTRODUCTION

The increasing consumer demand for natural food additives has focused interest on bacteriocins. They are ribosomally synthesized peptides of 30 to less than 60 amino acids, with a narrow to wide antibacterial spectrum against Gram-positive bacteria (Savadogo et al., 2006). The antibacterial compound is heat stable, and a producer strain displays a degree of specific self-protection against its own antibacterial peptide. Bacteriocins of lactic acid bacteria (LAB) are considered biopreservatives, as it is assumed that bacteriocins are degraded by the proteases of the gastrointestinal tract and most of the LAB are considered as GRAS (Generally Recognized as Safe) microorganisms (Holzapfel et al., 1995).

Although by definition all bacteriocins have a protein or peptide component that is essential for their bactericidal function, some have been reported to consist of combinations of different proteins or are composites of proteins together with lipid or carbohydrate moieties (Jimenez-Diaz et al., 1993). Improved protein purification protocols have shown that some bacteriocins previously considered high-molecular-weight protein aggregates may be small peptides that, because of their highly hydrophobic nature, had previously copurified with some other cellular components (Sahl, 1994). Although some Gram-positive bacteria have been shown to synthesize relatively high-molecular-weight, heat-labile bacteriocin-like substances (Vaughan et al., 1994), most of those described to date have been small, heat-stable cationic peptides.

A wide variety of bacterial products of Gram-positive bacteria have been referred to as bacteriocins, and various attempts have been made to classify these agents. Klaenhammer (1993) defined four distinct classes of lactic acid bacterial bacteriocins:

1. *Class I*, lantibiotics, are small (<5 kDa) peptides containing the unusual amino acids lanthionine (Lan), -methyllanthionine (MeLan), dehydroalanine, and dehydrobutyrine;
2. *Class II*, small (<10-kDa), relatively heat-stable, non-lanthionine-containing membrane-active peptides, subdivided into *Listeria*-active peptides with the N-terminal consensus sequence -Tyr-Gly-Asn-Gly-Val-Xaa-Cys- (Class IIa), poration complexes requiring two different peptides for activity (Class IIb), and thiol-activated peptides requiring reduced cysteine residues for activity (Class IIc);
3. *Class III*, large (>30-kDa), heat-labile proteins; and
4. *Class IV*, complex bacteriocins that contain essential lipid or carbohydrate moieties in addition to protein.

In the last decade many bacteriocins from LAB belonging to different groups have been characterized and purified, including nisin, diplococcin, acidophilin, bulgarican, helveticins, lactacins, sacakins, pediocin PA-1, and plantaricins (Nettles and Barefoot, 1993). Of these, nisin produced by *Lactococcus lactis* subsp. *lactis*, has been the most extensively characterized (Moreno et al., 2000). At present, nisin is the only bacteriocin commercially available and marketed because it has been conferred GRAS status by the Food and Drug Administration (FDA, 1988; FAO/WHO, 2006).

Only bacteriocins well characterized could be considered as potential natural food preservatives (De Vuyst and Vandamme, 1994). Therefore, the aim of this study was to isolate and identify potential bacteriocin producing LAB from CSS and to characterize the inhibitory activities of these bacteriocins.

MATERIAL AND METHODS

Fish Source

Fresh-gutted farmed salmon from Norway (*Salmo salar*) were acquired at Matosinhos' Doca (Porto, Portugal). Salmon arrived by lorry (72 h travel) in a chilled container with the temperature controlled between 0°C and 4°C inside polystyrene boxes (two layers of fish between two layers of ice). The fish were transported to the ESB/UCP, under chilled conditions, and submitted to a cold-smoking process (filleted, salted by brining or dry salt method, rinsed, smoked and vacuum-packaged).

Smoking Process

During salting, fillets were placed in a chilled chamber at 5°C. For dry-salting, the salt and sugar added corresponded to one third of the weight of fillet. For brining, 80% of saturated NaCl solution was used (brine/fish: 1/1). Draining was done overnight at 5°C. Smoking was done by two different processes. The first process consisted of drying for 2 h in the smoke chamber but without smoke at less than 30°C, followed by smoking for 6 h at less than 30°C. The second process involved drying for 6 h in the smoke chamber but without smoke at less than 30°C and smoking for 2 h. According to Tomé et al. (2007), these drying/smoking conditions enhance growth of total LAB and lactobacilli during the storage period in these products.

Two batches were processed for each smoking process. Each batch consisted of three salmon and three fillets processed in each smoking process.

The smoked samples were cooled overnight at 5°C. The following day, lug and pin bones and belly flaps were removed and the fillets were then sliced and vacuum packed in a Multivac-Gastrovac (Multivac Sepp Haggenmüller KG, A300/41/42, Germany) 1 mbar/10 s. The permeability of the packs to O₂,

CO₂ and N₂ were 4 mol/m².d.bar., 13 mol/m².d.bar, and 4 mol/m².d.bar, respectively (Vaz-Velho, 2000). Packs were stored for 3 weeks at 5°C and analyzed at the beginning (t₀) and at the end of the storage period (t₁).

Isolation and Phenotypic Characterization of Microorganisms

At the beginning of storage, t₀, and after three weeks, t₁, fillets from each smoking process were cut into small pieces and mixed. Ten grams of this mix were picked randomly and homogenized in 90 mL of sterile ¼-strength Ringer's solution (Lab M, Bury, UK) for 2 min in a Stomacher 400 Lab Blender (Seward Medical, London, UK). Serial decimal dilutions in ¼-strength Ringer's solution were prepared. Three independent samples per batch/smoking process were analyzed at each time interval. Total LAB were enumerated by pour-plating in nitrite actidione polymyxin (NAP) agar, pH 6.7 (Davidson and Cronin, 1973). All colonies were counted as presumptive LAB after 5 days of anaerobic incubation at 25°C. At t₀ and t₁, 10% of colonies overall were picked randomly from NAP plates containing 10–100 colonies. Presumptive LAB were subcultured on NAP agar without selective agents or on All-Purpose Tween (APT, Difco Laboratories, Detroit, Mich., USA) agar, examined for purity and characterized using Gram stain, cytochrome oxidase, and catalase tests. Organisms that were Gram-positive, cytochrome oxidase negative, and catalase negative were stored in APT broth with glycerol (30% v/v) at –80°C until further use.

Anti-listerial Activity of LAB, Nature of the Inhibition and Titre Determination

The antimicrobial activity of 614 presumptive LAB colonies picked from NAP plates, previously isolated from vacuum-packaged CSS was investigated against *L. monocytogenes* 54 (culture collection from ESB/UCP, Porto, Portugal) and *L. innocua* 2030c (Central Public Health Laboratory, Colindale, London, UK), a tetracycline resistant strain, using the spot method described by Tomé et al. (2006). Then, a 48 h-old culture in APT broth at 25°C was made of each LAB culture showing defined inhibition zones, ≥4 mm zone width. Cultures were adjusted to pH 6.5 with NaOH (1N), and cells were collected by centrifugation (7500 x g, 10 min, 4°C) followed by filtration of the supernatant through a 0.22 µm pore size membrane filter (Millipore Co., Bedford, Mass., USA). This solution was designated as a crude filtrate supernatant fluid (CFSF). The nature of the inhibition was assessed by treating the CFSF with the enzymes catalase and trypsin (both from Sigma–Aldrich Chemie GmbH, Steinheim, Germany) separately at a final concentration of 500 IU/mL and 0.1 mg/mL, respectively, for 2 h at 37°C. Cell-free supernatant (CFSF), CFSF treated with catalase and trypsin were spotted against *L. monocytogenes* 54 and *L. innocua* 2030c. Untreated samples were used as positive controls while enzyme solutions were used as

negative controls. The anti-listerial titre of each CFSF was determined against *L. monocytogenes* 54 by the serial twofold dilution assay. It was defined as the reciprocal of the highest dilution showing a distinct inhibition of the target strain and expressed in terms of arbitrary units per milliliter (AU/mL).

Strain Identification

The carbohydrate fermentation pattern of 9 bacteriocin-producing LAB strains was determined by using the API 50 CHL kit (BioMérieux, Mercy-l'Etoile, France). Coccoid strains, were submitted to some characteristic tests for *Enterococcus* spp. for example, the ability to grow at 10°C and 45°C in Tryptone Soy Broth (TSB, Lab M), growth at pH 9.0 and growth in TSB in the presence of 6.5% (w/v) NaCl at 30°C. The Vitek System (BioMérieux) was used for species identification of those strains which were able to grow at 10°C and 45°C and in the presence of 6.5% (w/v) of NaCl, using the Gram-Positive Identification Cards (GPI) according to the manufacturer's instructions.

Identification of all nine strains was genotypically confirmed by using genus/species specific PCR amplification. Each strain was cultured overnight in de Man, Rogosa, and Sharpe (MRS) broth (Merck, Darmstadt, Germany) supplemented with 0.4% DL-threonine (Sigma) at 30°C, and then the DNA was extracted according to Dellaglio et al. (1973).

Strains were further identified using specific-species primers (Table 1). The PCR reactions were performed in a thermal cycler (My Cycler™ Thermal Cycler Firmware, Bio-Rad, Richmond, Calif., USA) in a total volume of 20 µL according Ke et al. (1999), Berthier and Ehrlich (1988), Tilsala-Timisjarvi and Alantossova (1997), Mora et al. (1997), and Chagnand et al. (2001). PCR products were separated by electrophoresis (100 V) in agarose gel (2% agarose gel for *Enterococcus* spp. and *Lb. curvatus*, and 0.8% agarose gel for *Pediococcus* spp. and *Lb. fermentum*) (Qbiogene, Illkirch, France) in 2M Tris-Acetate-0.05M

Table 1: PCR primers used for genus/species specific PCR amplification.

Genus, species	Primer	R/F	Sequence (5'-3')	Reference
<i>Enterococcus</i> spp.	EntF	R	TAC TGA CAA ACC ATT CAT GAT G	(16)
	EntR	F	AAC TTC GTC ACC AAC GCG AAC	
<i>Lb. curvatus</i>	16	R	GCT GGA TCA CCT CCT TTC	(17)
	Lc	F	TTG GTA CTA TTT AAT TCT TAG	
<i>Lb. delbruecki</i>	DelF	R	ACG GAT GGA TGG AGA GCA G	(18)
	DelR	F	GCA AGT TTG TTC TTT CGA ACT C	
<i>P. acidilactici</i>	PacF	R	CGA ACT TCC GTT AAT TGA TTA T	(19)
	PuR	F	ACC TTG CGG TCG TAC TCC	
<i>Lb. fermentum</i>	Ferml	R	GTT GTT CGC ATG AAC AAC GCT TAA	(20)
	Lowlac	F	CGA CGA CCA TGA ACC ACC TGT	

Primers were acquired from Operon Biotechnologies GmbH (Cologne, Germany).

EDTA buffer and then stained with 0.5 µg/mL of ethidium bromide in deionized water. A 50-bp DNA or 100-bp DNA ladder (only for *Lb. curvatus* strains; JulesTM, Qbiogene) was used as molecular size marker. Strains *E. mundtii* PTA-7278 (American Type Culture Collection) and *E. faecium* HKLHS (CCSU); *Lb. curvatus* NCFB 2739T (National Collection of Food Bacteria); *Lb. delbrueckii* subsp. *delbrueckii* ATCC 9649; *P. acidilactici* ATCC 12697; *Lb. fermentum* ATCC 8289 were used as positive controls in specific genus/species PCR amplification reaction for *Enterococcus* strains (ET05, ET12, ET88), *Lb. curvatus* strains (ET06, ET30; ET31); *Lb. delbrueckii* (ET32); *P. acidilactici* (ET34) and *Lb. fermentum* (ET35) identification, respectively. *Lb. brevis* ATCC 14869, and *Lb. plantarum* ATCC 14917^T were used as negative control strains.

Molecular Size of the Bacteriocins

The molecular size of the bacteriocins from LAB ET05, ET06, ET12, ET30, ET31, ET32, ET34, ET35, and ET88 was determined by tricine-SDS-PAGE according the method described by Schagger and Von Jagow (1987). All strains were grown in MRS broth for 20 h at 30°C. The cells were harvested by centrifugation (8000 x g, 10 min, 4°C) and the bacteriocin precipitated from the cell-free supernatant with 70% saturated ammonium sulphate. The precipitate was resuspended in one tenth of the initial volume, in 25 mM ammonium acetate buffer (pH 6.5), desalted by using a 1000 Da cut-off dialysis membrane (Spectrum Inc., Oakland, Calif., USA) and separated by tricine-SDS-PAGE. A low molecular weight marker with sizes ranging from 2.5 to 4.5 kDa (Amersham International, Amersham, UK) was used. The gels were fixed and one half of each was stained with Coomassie Blue R250 (Saarchem, Krugersdorp, South Africa), and the other half remained unstained. The position of the active peptide band in the gel was determined by overlaying an unstained gel with cells of *L. innocua* 2030c (10⁶ CFU/mL) suspended in Brain Heart Infusion (BHI, Oxoid Ltd., Cambridge, UK) agar and incubated at 30°C for 24 h.

Characterization of Crude Filtrate Supernatant Fluid (CFSF)

Sensitivity to Enzymes

To test the sensitivity of the CFSF to the enzymes, samples of 1 mL were treated for 2 h at 37°C with the following filter-sterilized enzyme solutions, at a final concentration of 0.1 mg/mL, (v/v, 1/1): proteinase K in 20 mM Tris - HCl, pH 7.0; trypsin in 40 mM Tris-HCl, pH 8.2; protease E in 20 mM Tris-HCl, pH 7.4; pepsin A in 20 mM buffer Na₂HPO₄, pH 2.8; lysozyme in 20 mM Tris-HCl, pH 7.8, lipase in 0.1 M potassium phosphate, pH 6.0 and α - amylase in 20 mM Tris-HCl, pH 7.0 (with the exception of proteinase K supplied by Boehringer Mannheim GmbH, Germany, all the other enzymes were supplied

by Sigma). After, residual anti-listerial activity against *L. monocytogenes* 54 (ESB/UCP) was determined. Untreated samples were used as positive controls while enzymes solutions were used as negative controls.

Sensitivity to Different pH Values

The sensitivity of the CFSF from isolates ET05, ET06, ET12, ET30, ET31, ET32, ET34, ET35, and ET88 to different pH values was estimated by adjusting each sample to pH 2.0 to 10.0 (2.0, 4.0, 8.0, 10.0) with 1N NaOH or 1N HCl. After incubation at 10°C for 24 h, samples were re-adjusted to pH 6.5, filtered through a 0.22 µm pore size filter (Millipore Co). Serial twofold dilutions of each CFSF were spotted (10 µL) onto fresh APT plates seeded with *L. monocytogenes* 54 and the antimicrobial titres were calculated. Untreated samples were used as control.

Sensitivity to Heat Treatment

The effect of temperature on the CFSF from isolates ET05, ET06, ET12, ET30, ET31, ET32, ET34, ET35, and ET88 was ascertained by heating each of the CFSFs to 60°C for 0 (control), 10, 15, 30, and 60 min, or 100°C for 0 (control), 5, 10, 15, and 20 min. Residual activity was tested against *L. monocytogenes* 54 and titres were determined at each temperature and time interval.

Spectrum of Antimicrobial Activity

The spectrum of activity of the CFSF was tested against a wide range of target strains (Table 2) as well as against the bacteriocin-producing strains and 60 LAB isolated from CSS. These cultures were maintained at 4°C on agar slants. The following culture media and growth conditions used were: TSB supplemented with 0.6% (w/v) Yeast Extract (TSYE) for *Listeria* spp. (30°C), BHI for *S. aureus* and *Escherichia coli* (both at 30°C), TSB for *Salmonella* sp. (30°C), MRS medium for *Enterococcus* spp. (30°C), Columbia Agar containing 5% (v/v) of sheep blood (BioMerieux, for *Campylobacter jejuni* (37°C, microaerophilic atmosphere), and Thiosulfate Citrate Bile Sucrose (TCBS) agar (Merck) for *Vibrio parahaemolyticus* (25°C). The determination of the antimicrobial spectrum was performed quantitatively against *Listeria* spp. and qualitatively against the other microorganisms.

RESULTS

Anti-listerial Activity, Nature of the Inhibition and Titre Determination

From 614 presumptive LAB colonies isolated from NAP plates, 93 showed anti-listerial activity. Of these, two LAB cultures were inhibitory by organic

Table 2: Origin of microorganisms used in determining the antimicrobial spectrum of activity of CFSFs.

Species	Serotype	Origim	Species	Serotype	Source
<i>L. monocytogenes</i> 11994	4b	NCTC	<i>L. innocua</i> 11288	6a	NCTC
<i>L. monocytogenes</i> 4031T	1a	CECT	<i>L. innocua</i> 2030c		PHLS
<i>L. monocytogenes</i> 911	1/2c	CECT	<i>E. faecalis</i>	-	ATCC29212
<i>L. monocytogenes</i> 934	4a	CECT	<i>E. faecalis</i>	-	ESB/UCP
<i>L. monocytogenes</i> 936	1/2b	CECT	<i>E. faecalis</i> E88	-	CCSU
<i>L. monocytogenes</i> 937	3b	CECT	<i>E. faecium</i> HKLHS	-	CCSU
<i>L. monocytogenes</i> 78.39	4c	CIP	<i>S. aureus</i> 8532	-	NCTC
<i>L. monocytogenes</i> 104794	1/2a	CIP	<i>S. aureus</i> 1803	-	NCTC
<i>L. monocytogenes</i> ScottA	4b	PHLS/HAP	<i>S. aureus</i> 25923	-	ATCC
<i>L. monocytogenes</i> 18	1/2a	ESB/UCP	<i>S. aureus</i> 29213	-	ATCC
<i>L. monocytogenes</i> 54	4b	ESB/UCP	<i>C. jejuni</i>	-	Clinical
<i>L. monocytogenes</i> 211	3b	ESB/UCP	<i>E. coli</i> 9001	-	ESB/UCB
<i>L. monocytogenes</i> A92	1/2c or 3c	ESB/UCP	<i>E. coli</i> O157 non pathogenic	-	NCTC
<i>L. monocytogenes</i> B15	4d, 4b or 4e	ESB/UCP	<i>Pseudomonas aeruginosa</i>	-	ESB/UCP
<i>L. monocytogenes</i> B19	4b, 4d or 4e	ESB/UCP	<i>Salmonella enteritidis</i> 05188	-	ESB/UCP
<i>L. monocytogenes</i> G12	4b, 4d or 4e	ESB/UCP	<i>S. typhimurium</i>	-	NCTC
<i>L. monocytogenes</i> L7	4b, 4d or 4e	ESB/UCP	<i>V. parahaemolyticus</i>	-	ESB/UCP
					INIAP/IPIMAR-CRIPN

ATCC: American Type Culture Collection; CECT: Colección Española de Cultivos tipo; CIP: Pasteur Institute Collection; CCSU: Culture Collection Stellenbosch University; Department of Microbiology, South Africa; ESB/UCP: Escola Superior de Biotecnologia/Universidade Católica Portuguesa; INIAP/IPIMAR-CRIPN - Instituto Nacional de Investigação Agrária e das Pescas, Portugal; NCTC: National Collection of Type Cultures, UK; PHLS: Central Public Health Laboratory, London UK.

acid production and nine were bacteriocinogenic. The CFSF of these nine bacteriocin-producing strains were able to inhibit *L. monocytogenes* 54 and *L. innocua* 2030c in an *in vitro* assay performed at 25°C. Anti-listerial activity was not lost in any of the CFSF either after adjusting their pH to 6.5 or by catalase treatment. Anti-listerial activity was lost after incubation with trypsin, indicating the proteolytic nature of the nine inhibitory compounds.

The lowest titres were calculated for ET12 and ET88, 800 AU/mL. ET06, ET31, ET34, and ET35 had titres of 1000 AU/mL. ET30 and ET32 had titres of 2000 AU/mL whilst the highest titre was of 10000 AU/mL, recorded for CFSF ET05.

Strain Identification

All strains considered as LAB and bacteriocin-producers isolated from vacuum-packaged CSS fillets, were Gram-positive, catalase, and oxidase negative. Isolates ET05, ET12, ET34, and ET88 were cocci, whereas ET06, ET30, ET31, ET32, and ET35 were rods. According to the carbohydrate fermentation patterns of the isolates carried out with the API 50 CHL system (results not shown), all the strains could be successfully identified: ET06, ET30, and ET31 as *Lb. curvatus* (% Id: 98.3; 92.7; 93.3, respectively); ET32 as *Lb. delbrueckii* (% Id: 99.6); ET34 as *P. pentosaceus* (% Id: 99.9%); ET35 as *Lb. cellobiosus* (% Id: 99.5%; this species was reclassified recently as *Lb. fermentum*). Strains ET05, ET12, and ET88 were identified as members of the genus *Enterococcus* based on morphology (cocci in pairs), ability to grow in broth at 10°C and at 45°C, at pH 9.0 and in the presence of 6.5% of NaCl, and by PCR with genus specific primers (results not shown). Further identification to species level, *E. faecium*, was based on the Gram-Positive Identification Cards, (GPI) from the Vitek System. Identification obtained with API 50 CHL coincided with those obtained by PCR with specific primers for *Lb. curvatus* (ET06, ET30, ET31, results not shown), *Lb. delbrueckii* (ET32) and *Lb. fermentum* (ET35), whilst isolate ET34 identified as *P. pentosaceus* by API 50 CHL, instead was identified as *P. acidilactici* by PCR (results not shown).

Molecular Size of the Bacteriocins

The molecular size of the nine bacteriocins produced by the LAB strains isolated from vacuum-packaged CSS, ranged from 2.8 to 4.5 kDa. Bacteriocins ET05, ET12, ET32, and ET34 from *E. faecium* strains *Lb. delbrueckii* strain and *P. acidilactici* strain, respectively, are peptides with molecular weights of approximately 3.5 kDa (Fig. 1A and B). Bacteriocins ET88 and ET35 from *E. faecium* and *Lb. fermentum* have molecular weights of approximately 3.7 kDa and 4.0 kDa, respectively (Fig. 1C). Bacteriocins ET30 and ET31 from *Lb. curvatus* strains had molecular weights of 3.1 kDa and

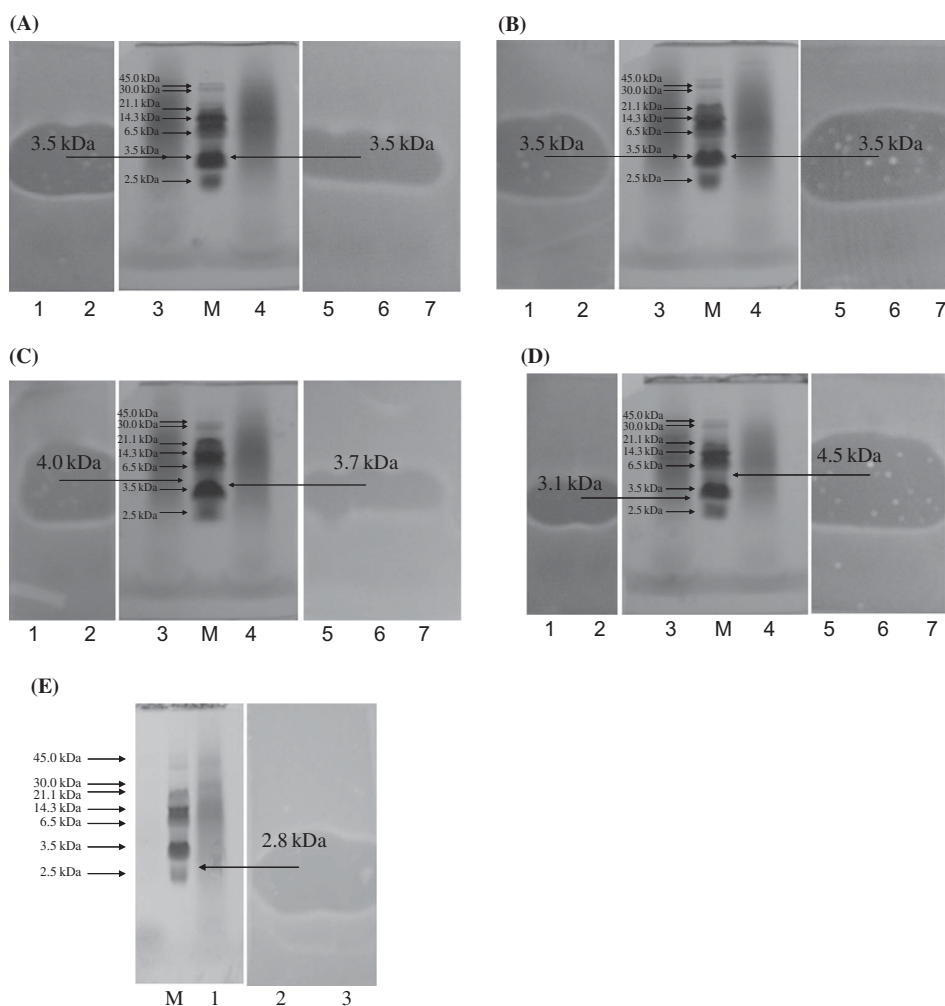


Figure 1: Tricine-SDS-PAGE. (A) Lanes 1-2 and 5-7 = zone of growth inhibition corresponding to the positions of bacteriocins ET05 and ET12, respectively. Lanes 3 and 4 = peptide bands stained with Coomassie Blue R250 of bacteriocins ET05 and ET12, respectively. (B) Lanes 1-2 and 5-7 = zone of growth inhibition corresponding to the positions of bacteriocins ET32 and ET34, respectively. Lanes 3 and 4 = peptide bands stained with Coomassie Blue R250 of bacteriocins ET32 and ET34, respectively. (C) Lanes 1-2 and 5-7 = zone of growth inhibition corresponding to the positions of bacteriocins ET35 and ET88, respectively. Lanes 3 and 4 = peptide bands stained with Coomassie Blue R250 of bacteriocins ET35 and ET88, respectively. (D) Lanes 1-2 and 5-7 = zone of growth inhibition corresponding to the positions of bacteriocins ET30 and ET31, respectively. Lanes 3 and 4 = peptide bands stained with Coomassie Blue R250 of bacteriocins ET30 and ET31, respectively. (E) Lanes 2-3 = zone of growth inhibition corresponding to the positions of bacteriocin ET06. Lane 1 = peptide bands stained with Coomassie Blue R250 of bacteriocin ET06. Lane M = molecular-mass marker (range 2.5 to 45.0 kDa; Amersham). The gel was overlaid with *L. innocua* 2030c (approximately 10^6 CFU mL⁻¹ suspended in BHI agar).

4.5 kDa respectively (Fig. 1D). Finally bacteriocin ET06, also from a *Lb. curvatus* strain, was the smallest peptide with a molecular mass of about 2.8 kDa (Fig. 1E).

Characterization of Crude Filtrate Supernatant Fluid (CFSF)

a. Sensitivity to Enzymes

The anti-bacterial activities of all preparations were not affected by catalase, lipase, α -amylase or lysozyme but were completely inactivated by the proteolytic enzymes protease E, trypsin and proteinase K indicating that anti-bacterial activity was associated with proteinaceous substances. The enzyme pepsin A also inactivated supernatants from *Lb. curvatus* strains.

b. Sensitivity to Different pH Values

The stability of CFSFs in a pH range from 2.0 to 10.0 is shown in Table 3. They differed with regard to their sensitivity to inactivation by changes in pH. Many were stable only in acidic and neutral conditions and were even inactivated at pH 8.0, for example, CFSF from *Lb. fermentum* ET35. Although the nine CFSFs tested were stable at pH 6.5, only CFSF from *E. faecium* ET05 and *Lb. curvatus* ET06 strains remained constant in a wide range of pH from 2.0 to 8.0. All the other bacteriocins were partly inactivated in the extremes of the pH range. CFSF activity of *Lb. curvatus* ET31, *Lb. fermentum* ET35 and *E. faecium* ET88 was completely lost at an alkaline pH while the maximal activity of the others CFSF was reduced to a half.

c. Sensitivity to Heat Treatment

All CFSFs maintained their antimicrobial activity even at 100°C for 20 min. Only CFSFs titres from *Lb. curvatus* ET30 and ET31 lost 20% and 50% of their activities after 10 min and 30 min of heating at 60°C, respectively, and

Table 3: Effect of pH on bacteriocin activity (AU mL⁻¹) produced by the test isolates.

pH	Strains								
	ET05	ET06	ET12	ET30	ET31	ET32	ET34	ET35	ET88
2	10000	1000	800	2000	800	1600	1000	500	400
4	10000	1000	800	2000	800	1600	1000	500	400
6.5 ^a	10000	1000	800	2000	1000	2000	1000	1000	800
8	10000	1000	400	1000	800	1600	500	nd	400
10	1000	800	400	800	nd	1600	500	nd	nd

^aTitre of the CFSF; nd = not detected.

after 10 min and less than 5 min of heating at 100°C, respectively (titre declined from 2000 AU/mL to 1000 AU/mL for the former and from 1000 AU/mL to 800 AU/mL for the last one). Then, both maintained their stability during the remaining time of the heating period.

Spectrum of Antimicrobial Activity

The antagonistic effect of the neutralized and filtered culture supernatants on various Gram-positive bacteria was tested and titres were calculated for each one (Table 4). The inhibition of the various target microorganisms was compared with the inhibitory activity of the nisin-producing *Lc. lactis* subsp. *lactis* ATCC 11454. CFSF from this strain presented a wide range of inhibitory spectrum affecting different nontaxonomically related genera like *Listeria*, *Staphylococcus* and *Enterococcus*. Although CFSF from the other bacteriocin-producing strains inhibited a similar but limited range of target microorganisms, there was considerable variability of sensitivities of *L. monocytogenes* strains inter and intra the nine CFSFs tested. The highest activities (>10000 AU/mL) were obtained from *E. faecium* (ET05), *Lb. curvatus* (ET30) and *Lc. lactis* subsp. *lactis* ATCC 11454 against practically the same *Listeria* strains, i.e. *L. monocytogenes* 11994, *L. monocytogenes* 934, *L. monocytogenes* 936, *L. monocytogenes* Scott A, *L. monocytogenes* 211, *L. monocytogenes* G12 and *L. innocua* 2030c. No inhibitory activity was recorded for CFSF from *Lb. curvatus* ET06 against *L. monocytogenes* 18, *L. monocytogenes* A92 and *L. monocytogenes* L7. No relationship seems to exist among the several serotypes of *Listeria* spp. tested and the level of inhibition obtained. Only CFSFs from *Lb. curvatus* ET06 and from *Lb. fermentum* ET35 were not able to inhibit *E. faecalis* 29212 and *E. faecalis* (ESB/UCP) while partial or complete inhibition was observed for *E. faecalis* 29212, *E. faecalis* (ESB/UCP), *E. faecalis* E88, *E. faecium* HKLHSE with the others CFSFs. On the other hand, just *Lc. lactis* subsp. *lactis* ATCC 11454 showed antimicrobial properties against *S. aureus* strains. None of the CFSFs inhibited any of the Gram-negative strains assessed (results not shown).

The antagonistic activity exhibited by each CFSF against the other bacteriocin-producing strains is recorded in Table 5. Bacteriocin ET31 (*Lb. curvatus*) showed inhibitory activity against the *E. faecium* strains (weak and reversible against *E. faecium* ET12), while bacteriocins ET06 and ET30 from *Lb. curvatus* strains did not show anti-bacterial activities against the same LAB tested. Inhibition by bacteriocins from *E. faecium* was restricted to closely related bacteria, although the antimicrobial compound inhibited *Lc. lactis* subsp. *lactis* ATCC 11454. CFSF from *Lb. delbrueckii* ET32 exhibited activity only against *E. faecium* ET05 while bacteriocin ET35 was inactive against the entire LAB bacteriocin-producing

Table 4: Inhibitory spectrum of bacteriocins against several Gram-positive bacteria.

Microorganisms	ET05	ET06	ET12	ET30	ET31	ET32	ET34	ET35	ET88	<i>L.lactis</i> ATCC11454
<i>L. monocytogenes</i> 11994	20000*	1000	800	40000	1000	8000	2000	400	8000	20000
<i>L. monocytogenes</i> 4031T	8000	800	1000	80000	800	2000	2000	1000	1000	20000
<i>L. monocytogenes</i> 911	8000	100	200	8000	800	1000	1000	400	1000	8000
<i>L. monocytogenes</i> 934	20000	3000	1000	80000	1000	2000	4000	1000	2000	80000
<i>L. monocytogenes</i> 936	80000	8000	1000	80000	8000	8000	8000	1000	8000	100000
<i>L. monocytogenes</i> 937	8000	8000	400	100000	800	4000	8000	400	8000	8000
<i>L. monocytogenes</i> 78.39	1000	200	400	8000	400	800	1000	200	2000	800
<i>L. monocytogenes</i> 104/794	8000	800	800	10000	2000	8000	8000	1000	8000	8000
<i>L. monocytogenes</i> ScottA	20000	4000	8000	80000	1000	8000	10000	2000	8000	40000
<i>L. monocytogenes</i> 18	8000	-	800	2000	4000	1000	4000	800	1000	8000
<i>L. monocytogenes</i> 54	10000	1000	800	2000	1000	2000	1000	1000	800	8000
<i>L. monocytogenes</i> 211	20000	2000	400	100000	800	2000	2000	400	2000	100000
<i>L. monocytogenes</i> A92	2000	-	200	8000	200	800	1000	400	1000	4000
<i>L. monocytogenes</i> B15	8000	2000	200	8000	800	1000	1000	200	2000	8000
<i>L. monocytogenes</i> B19	8000	1000	800	8000	800	1000	8000	200	10000	10000
<i>L. monocytogenes</i> G12	20000	2000	200	100000	400	1000	4000	200	2000	100000
<i>L. monocytogenes</i> L7	6000	-	800	8000	800 ^b	1000	1000	600	1000	8000
<i>L. innocua</i> 11288	4000	100 ^b	400	8000	400	1000	1000	400	800	8000
<i>L. innocua</i> 2030c	8000	200	800	20000	800	8000	2000	800	8000	10000
<i>E. faecalis</i> 29212	+ ^b	-	+ ^b	+	+ ^b	+	+	-	+	+
<i>E. faecalis</i> ESB/UCP	+ ^b	-	+	+	+	+	+	-	+	+
<i>E. faecalis</i> E88	+	+	+	+	+	+	+	+	+	+
<i>E. faecium</i> HKLHS	+	+	+	+	+	+	+	+	+	+
<i>S. aureus</i> 8532	-	-	-	-	-	-	-	-	-	+
<i>S. aureus</i> 1803	-	-	-	-	-	-	-	-	-	+
<i>S. aureus</i> 25923	-	-	-	-	-	-	-	-	-	+
<i>S. aureus</i> 29213	-	-	-	-	-	-	-	-	-	+

*AU mL⁻¹ +:inhibition - : no inhibition ^bbacteriostatic, reversible within 24–48h.

Table 5: Antimicrobial spectrum of activity of CFSFs from LAB against the bacteriocin-producing strains.

Target strains	Crude Filtrate Supernatant Fluid (CFSF)									<i>L.lactis</i> ATCC 11454
	ET05	ET06	ET12	ET30	ET31	ET32	ET34	ET35	ET88	
<i>E. faecium</i> ET05	–	0*	3	0	5	3 ^b	5 ^b	0	4	10
<i>Lb. curvatus</i> ET06	0	–	0	0	0	0	0	0	0	0
<i>E. faecium</i> ET12	11 ^b	0	–	0	5 ^b	0	0	0	2 ^b	6
<i>Lb. curvatus</i> ET30	0	0	0	–	0	0	0	0	0	0
<i>Lb. curvatus</i> ET31	0	0	0	0	–	0	0	0	0	0
<i>Lb. delbrueckii</i> ET32	0	0	0	0	0	–	0	0	0	0
<i>P. acidilactici</i> ET34	0	0	0	0	0	0	–	0	0	0
<i>Lb. fermentum</i> ET35	0	0	0	0	0	0	0	–	0	0
<i>E. faecium</i> ET88	10	0	2 ^b	0	8	0	0	0	–	10
<i>Lc. lactis</i> ATCC11454	8	0	10	0	0	0	0	0	4	–

*inhibition zone (mm); ^b bacteriostatic, reversible within 24–48 h.

bacteria tested. Activity of CFSFs was also screened against 60 LAB isolated from vacuum-packaged CSS; the growth of LAB tested was not inhibited by any of the CFSFs tested.

DISCUSSION

Strain Identification

Our results indicated phenotypic heterogeneity and genetic diversity among the vacuum-packaged CSS bacteriocin-producing isolates, with a good correlation between the phenotypic and genetic identification of the strains. As sugar fermentation patterns are not considered to be a reliable method of distinguishing among *Enterococcus* spp. due to the heterogeneous and atypical profiles displayed by this genus (Pérez et al., 2000), further identification to species level was carried out with the Vitek system, which classified all the enterococcal isolates as *E. faecium* in accordance with the genotypic results at genus level performed by PCR reaction. The fermentation activities recorded with API galleries agreed with those registered with VITEK; however, API 50 CH reactions were not enough to identify microorganisms belonging to *Enterococcus* genus. On the other hand, the biochemical tests carried out with the GPI cards showed that all the *Enterococcus* isolated belonged to the same genus/species *E. faecium*. The genus/species specific PCR identification assay allowed the proper identification of all LAB.

The species identified in our study had been previously isolated from vacuum-packaged CSS or fish products. For instance, González-Rodríguez

et al. (2002), in a study conducted to assess the microbiological quality of vacuum-packaged CSS produced by various processors in Spain, isolated eight species of *Lactobacillus* that included a high percentage of *Lb. curvatus* subsp. *curvatus* as well as *Lb. delbrueckii* subsp. *delbrueckii*, although this last was a minor proportion. Several strains of *Enterococcus* spp. were also isolated. Lyhs et al. (1999) isolated *Lb. curvatus* together with *Leuconostoc mesenteroides* subsp. *mesenteroides*, *Lb. citreum*, *Lb. sakei*, from spoiled vacuum-packaged, cold-smoked rainbow trout. In contrast, the occurrence of *Lb. curvatus* in non-spoiled vacuum-packaged salmon has been reported previously by Truelstrup-Hansen (1995), who identified from 168 LAB, 50% as *Lb. curvatus*.

Few reports have recorded the presence of *Lb. fermentum* in fish. According to Sharpe and Pettipher (1993), this heterofermentative *Lactobacillus* specie has predominated in different kinds of spoiled herring. Also, it has been frequently identified in chickens (Reque et al., 2000), a Balinese sausage (Antara et al., 2002), or boza, a traditional Bulgarian fermented beverage (von Mollendorff et al., 2006).

The importance of *P. acidilactici* strains in the food industry is related to their use as starter cultures in fermented meat and vegetable products (Amézquita and Brashears, 2002). This species, like most other LAB species, is involved in extending the shelf life and improving the hygienic quality of various fermented products, via the production of lactic acid and/or the secretion of anti-bacterial compounds such as bacteriocins (Stiles, 1996). The association of pediococci with proteinaceous foods such as fresh and cured meat, and raw sausages has frequently been reported (Holzapfel et al., 2005), and particularly for *P. acidilactici* in fermented sausages (Parente et al., 2001). *Pediococcus* spp. also have been reported in fresh and marinated fish (Paludan-Muller et al., 2002).

Molecular Size of the Bacteriocins

Although tricine-SDS-PAGE is not an accurate technique to calculate the molecular mass of molecules, it gives valuable information about the presence of either one or two peptides (Moreno et al., 2002). A single inhibition zone was seen for all the bacteriocins assessed by this technique in this study, although the samples precipitated by ammonium sulphate and analyzed by tricine-SDS-PAGE contained more than one protein band.

Bacteriocins ET05 and ET12 from *E. faecium* strains ET05 and ET12, respectively, possessed a molecular mass near 3.5 kDa whereas bacteriocin ET88 from *E. faecium* ET88 has a molecular mass slightly higher. Other enterocins from *E. faecium* strains with close molecular masses have been reported (Moreno et al., 2002) as well as others with higher masses such as enterocin 012 (Jennes et al., 2000). Moreover, strains of *E. faecium* capable of producing two kinds of bacteriocins with different molecular weights have

been described, that is, enterocin A, pediocin-like bacteriocin with a molecular weight of 4.8 kDa (Aymerich et al., 1996), and enterocin B, a small nonlantibiotic bacteriocin not belonging to the pediocin group and with a molecular weight of 5.5 kDa (Nilsen et al., 1998).

All the three bacteriocins isolated from *Lb. curvatus* strains, named ET06, ET30, and ET31, had different molecular masses. While bacteriocins ET30 and ET06 migrated upon SDS-PAGE gel electrophoresis as small peptides of approximately 3.1 kDa and 2.8 kDa (Fig. 1E), respectively, bacteriocin ET31 had a molecular mass slightly higher (4.5 kDa, Fig. 1D). The molecular weight of curvacin A, the first bacteriocin identified and characterized from a strain of *Lb. curvatus*, is around 3.0 – 5.0 kDa (Messens et al., 2002). Like the other bacteriocins assessed here, ET32 from *Lb. delbrueckii* ET32 is a small peptide with an apparent molecular size of approximately 3.5 kDa (as estimated by tricine-SDS-PAGE analysis). Although bacteriocins from *Lb. delbrueckii* have been rarely described, Boris et al. (2001) characterized a bacteriocin UO004 produced by *Lb. delbrueckii* subsp. *lactis* UO004 with a molecular weight near 6.0 kDa by SDS-PAGE analysis. The molecular mass of the pediocin ET34 was approx. 3.5 kDa. Its size is similar to pediocin PD-1 from *P. damnosus* and does not correspond with the molecular mass of the pediocin PA-1 of 4.629 kDa from *P. acidilactici* PAC 1.0 mainly found in fermented sausages and other meat and vegetable fermentations (Jager and Harlander, 1992). Different electrophoretic mobilities of bacteriocins of *Lb. fermentum* have been reported; >1.0 kDa to <5.0 kDa for Fermenticin B produced by *Lb. fermentum* Beijerinck CCRC 14018 (Yan and Lee, 1997), 2.3 kDa – 3.0 kDa (von Mollendorff et al., 2006).

Taking into account their bactericidal activity, proteinaceous nature, heat resistance, and low molecular weight, bacteriocins ET05, ET06, ET12, ET32, ET34, ET35, and ET88 can be classified as small, heat-stable *Listeria*-active peptides possibly belonging to class IIa according to the definition given by Klaenhammer (1993).

Characterization of Crude Filtrate Supernatant Fluid (CFSF)

a. Sensitivity to Enzymes

The sensitivity of the inhibitors to enzymes was tested to gain insight into their chemical structure. Results demonstrated that the active moiety of the entire inhibitory substances was not hydrogen peroxide, a lipid, or a glucan, respectively. The loss of anti-bacterial activity of the CFSFs upon treatment with trypsin, protease E, and proteinase K permits their classification as bacteriocins or/and bacteriocin-like inhibitory substances. The susceptibility of bacteriocins to enzymatic degradation suggests that these peptides will be degraded in the intestinal tract and so will be easily digested without affecting the intestinal flora. From this point of view, *Lb. curvatus* strains and their

bacteriocins could be of great interest as bioprotective cultures, because their bacteriocins will be more quickly digested than the other bacteriocins (they are digested by pepsin A, at pH 2). De Martinis et al. (2003) reported similar bacteriocin sensitivity to these enzymes for the bacteriocins of two strains of *Lb. curvatus* isolated from sausages.

The bacteriocins assessed in this study consisted of pure peptides, as their activities were inactivated by treatment with proteolytic enzymes, while lipolytic and glycolytic enzymes had no effect on activity. All of them shared their sensitivity to treatment with proteolytic enzymes but insensitivity to lipolytic and glycolytic enzymes. According to Piard and Desmazeaud (1992), LAB synthesize many bactericidal agents, some of which are bacteriocins with a proteinaceous active moiety and others are nonprotein agents.

b. Sensitivity to Different pH Values

Bacteriocins differ greatly with regard to their sensitivity to inactivation by changes in pH and temperature. Many are stable only in acid and neutral conditions, and are even inactivated at pH 8.0, for example, lactostrepcins, or pH 10.0, like nisin and pediocin PA-1 (Moreno et al., 2000; Chien-Wei et al., 2004). Most of the bacteriocins maintained full activity over a pH range of 2.0–8.0 and were partially or completely inactivated at pH 10.0. In this case, the loss of the activity was irreversible and could not be regained upon lowering the pH to 6.5. Irreversible inactivation can result from a combination of denaturation and chemical modifications of the molecule. The loss of activity can be related to the solubility of the bacteriocins. In the case of enterocins, the isoelectric point of all known enterocins was around 8.3–10.7, which implies that the solubility increases at pHs below the pI. All the nine inhibitory substances showed maximal activity at pH 6.5. This pH value is very close to the normal pH of CSS fillets (6.0) (Tomé et al., 2006), which is a positive aspect for the addition of these anti-listerial peptides (or their producing bacteria) into this product.

c. Sensitivity to Heat Treatment

Bacteriocins were similar with regard to their sensitivity to inactivation by temperature. Like most of the known bacteriocins, they were mainly heat-tolerant at pH 6.5 (Todorov et al., 1999) after 60 min of treatment at 60°C and 20 min of treatment at 100°C. Hill (1994) and Jennes et al. (2000) also reported the heat stability of enterocins. Only bacteriocins ET30 and ET31 were moderately heat-stable at 60°C and 100°C, thus resembling nisin produced by *Lc. lactis* WNC20 which was inactivated after 15 min at 121°C at pH 7.0 (Noonpakdee et al., 2003) or the pediocin PA-1 showing about 40% activity lost after 15 min of heating at 121°C in the pH range pH 2.5–9.0 (Ray, 1994). The moderate stability of the inhibitory peptide produced by *Lb. curvatus* strains was previously reported (De Martinis et al., 2003).

Spectrum of Antimicrobial Activity

The assessment of the inhibitory spectrum is an important characteristic in order to evaluate the possibility of using the bacteriocin-producing strains or the bacteriocins alone as an additional barrier against spoilage and/or food-borne pathogens in food. Only 2 (22%) out of 9 neutralized culture filtrates displayed activity toward all the Gram-positive microorganisms examined. One characteristic of classical bacteriocins is a narrow spectrum of activity (Tagg et al., 1976). CFSFs of ET05 and ET88 of *E. faecium* demonstrated the widest antimicrobial spectrum of activity. Moreover it was similar to the inhibitory spectrum of the nisin of *Lc. lactis* subsp. *lactis* ATCC 11454, which suppressed the growth of all the *Listeria* spp. as well as *E. faecalis* strains, *E. faecium* and *S. aureus* strains. CFSF from *P. acidilactici* ET34 showed an inhibitory spectrum slightly different from those exhibited by the antimicrobial peptide called pediocin F and pediocin A-1 produced by *P. acidilactici* (Osmanağaoğlu et al., 1998), which is inhibitory to a variety of LAB often encountered in foods, such as *Enterococcus*, as well as spoilage and pathogenic strains i.e. *E. coli* and *S. aureus*. Fimland et al. (2000) showed that the C-terminal disulfide bridge in pediocin-like bacteriocins contributes to widening of the antimicrobial spectrum.

All bacteriocins inhibited strains of *L. monocytogenes*, a pathogen often isolated from a variety of foods. This activity is quite interesting, especially taking into account that many reports showed the occurrence and growth of *L. monocytogenes* in fish. *L. monocytogenes* serovar 4b, which is involved in human infections has been isolated from fresh trout samples (Hangard-Vidaud et al., 1989) as well as smoked salmon. Among the 19 *Listeria* spp. strains tested, 84.2% were inhibited by bacteriocin ET06. The high anti-listerial activity displayed by the nine bacteriocins is characteristic of the class IIa bacteriocins (Klaenhammer, 1993; Ennahar et al., 2000). It is interesting to note the resistance of *L. monocytogenes* 18, *L. monocytogenes* A92 and *L. monocytogenes* L7 to bacteriocin ET06. It has been a common observation by bacteriocin researchers that resistance to bacteriocin action is not only species or strain-specific but also dose-dependent. Strains that have the receptors and relevant characteristics of cytoplasmic membrane for proper attachment and sensitivity to the bacteriocin, are generally inhibited (Ennahar et al., 2000).

Our study confirms that *L. innocua* 2030c resembles *L. monocytogenes* in sensitivity toward LABs ET05, ET06, ET12, ET30, ET31, ET32, ET34, ET35, ET88 bacteriocins and is thus a suitable organism when extensive work with the pathogen is undesirable. On the other hand, the sensitivity differences of each *Listeria* spp. against the nine CFSFs studied is notable. The highest anti-listerial activity was recorded for CFSF ET05, ET30 and nisin from *Lc. lactis* subsp *lactis* ATCC 11454. The dissimilar responses of *Listeria* spp. to bacteriocins did not appear to be determined either by the serological type

of the organisms, nor by the growth phase of *Listeria* strains; all were tested in the stationary phase. As mentioned by Jydegaard et al. (2000), the bacteriocin inactivation of *L. monocytogenes* 412 by nisin and low pediocin concentrations are growth phase dependent, with exponentially growing cells being more susceptible than stationary phase cells. The sensitivity differences of *Listeria* spp., as well other food-borne and spoilage microorganisms toward diverse bacteriocins have been documented previously by several researchers (Moreno et al., 2000; Coventry et al., 1997; Østergaard et al., 1998). As expected, none of the CFSFs inhibited any of the Gram-negative bacteria evaluated. Activity of bacteriocins against Gram-negative bacteria is unusual and has only been reported for a few bacteriocins of LAB (Caridi, 2002; Todorov and Dicks, 2004a; Todorov and Dicks LMT, 2004b).

Well-defined bacteriocins produced by lactobacilli usually have inhibitory activities restricted to closely related species. However, it is worth noting that CFSFs showed limited activity against other bacteriocin-producer strains (Table 5); just CFSF ET05, ET12 and ET88 exhibited antimicrobial properties toward *E. faecium* strains and *Lc. lactis* subsp. *lactis* ATCC11454, although in some cases this inhibitory activity disappeared within 48 h. CFSF from *Lc. lactis* subsp. *lactis* ATCC 11454 such as CFSF ET31 showed limited activity toward bacteria belonging to the genus *Enterococcus* and no activity at all against the *Lactobacillus* strains tested. Contrary to what was expected, most of the bacteriocins did not kill species of bacteria that are known to have the same ecological niche. Similar results were reported for pediocin PD-1 produced by *P. damnosus* as it was not active against other pediococci. In that respect, Coventry et al. (1997) pointed out that the sensitivity of *Listeria* species as well as LAB to bacteriocins may be influenced by the content or type of agar in the growth media. The different inhibition patterns registered for neutralised and filtered culture supernatants of *Lb. curvatus* strains, against the LAB tested, could suggest that the inhibitory compound produced by *Lb. curvatus* ET31 is different from those produced by the other *Lb. curvatus* strains. These results are promising in view of a recent investigation into the use of combinations of LAB bacteriocins or their producing strains in order to broaden the spectrum to a wide variety of pathogens and food spoilage organisms, and avoid the phenomenon of bacteriocin resistance development, which is the main limiting factor for bacteriocin effectiveness. It is generally admitted that each bacteriocin-sensitive bacterial population includes potentially tolerant and/or resistant cells with structural modifications or at least with a high predisposition to such modifications, which would allow them to spontaneously emerge during exposure to the bacteriocin (Hanlin et al., 1993). In particular, modifications in the cytoplasmic membrane composition are often investigated to explain bacteriocin resistance, considering the key role of the membrane in the activity of bacteriocins (Mazzotta and Montville, 1997). On the other hand, no activity of CFSFs was obtained in the screening assay against

60 LAB isolated from vacuum-packaged CSS (results not shown). This specificity could be useful for suppressing growth of *Listeria* in lightly preserved seafood where it could be a serious problem, without interfering with other desirable lactic flora. These results resemble the activity spectrum of the divergicin M35 from *Carnobacterium divergens* M35, which did not inhibit bacteria belonging to the genera *Lactobacillus*, *Lactococcus*, *Enterococcus*, *Pediococcus*, *Propionibacterium*, and *Bifidobacterium* (Tahiri et al., 2004).

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