Research Article

Characterization and Application of Enterocin RM6, a Bacteriocin from Enterococcus faecalis

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Received 1 May 2013; Accepted 30 May 2013

Academic Editor: Argyro Bekatorou

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Use of bacteriocins in food preservation has gained great attention in recent years. The goal of this study is to characterize enterocin RM6 from Enterococcus faecalis OSY-RM6 and investigate its efficacy against Listeria monocytogenes in cottage cheese. Enterocin RM6 was purified from E. faecalis culture supernatant using ion exchange column, multiple C 18 silica cartridges, followed by reverse-phase high-performance liquid chromatography. The molecular weight of enterocin RM6 is 7145.0823 as determined by mass spectrometry (MS). Tandem mass spectrometry (MS/MS) analysis revealed that enterocin RM6 is a 70-residue cyclic peptide with a head-to-tail linkage between methionine and tryptophan residues. The peptide sequence of enterocin RM6 was further confirmed by sequencing the structural gene of the peptide. Enterocin RM6 is active against Gram-positive bacteria, including L. monocytogenes, Bacillus cereus, and methicillin-resistant Staphylococcus aureus (MRSA). Enterocin RM6 (final concentration in cottage cheese, 80 AU/mL) caused a 4-log reduction in population of L. monocytogenes inoculated in cottage cheese within 30 min of treatment. Therefore, enterocin RM6 has potential applications as a potent antimicrobial peptide against foodborne pathogens in food.

1. Introduction

Listeria monocytogenes is the causative agent of human listeriosis, a disease with 20–30% fatality [1]. L. monocytogenes is an ubiquitous microorganism that can be found in many raw and processed foods. Soft cheese products have been associated with several listeriosis outbreaks [2–4]. Rudolf and Scherer [5] reported that L. monocytogenes was found in 6.4% of European red smear cheese. Recently, a multistate listeriosis outbreak occurred in the United States due to consumption of contaminated cantaloupe, resulting in 30 deaths across 28 states [6]. Therefore, control of L. monocytogenes remains an important issue for the food industry.

Use of bacteriocins in food preservation has gained a great attention in recent years [7]. The effect of many antilisterial bacteriocins (e.g., nisin, pediocin, and enterocin AS-48) has been investigated in foods including dairy products, meats, and fresh produce [8–10]. In this study, we described a new isolate of Enterococcus faecalis from raw milk, which produces an anti-listerial peptide, enterocin RM6. The bacteriocin was purified by liquid chromatography techniques. The peptide sequence was determined by mass spectrometry and confirmed by structural gene sequencing. In addition, the efficacy of enterocin RM6 against L. monocytogenes in cottage cheese was investigated.

2. Materials and Methods

2.1. Bacterial Strains and Media. Strains were obtained from the culture collection of the food safety laboratory at The Ohio State University (Columbus, OH, USA). The producer strain, E. faecalis OSY-RM6, which was isolated from raw milk [11], was grown in MRS broth (Becton and Dickinson, Sparks, MD, USA). The indicator strain, Lactobacillus
cellobiosus OSU 919, was cultivated in MRS broth. Other bacterial strains used in the test of antimicrobial spectrum are listed in Table 1.

2.2. Strain Identification by 16S rDNA Sequencing. Genomic DNA of strain OSY-RM6 was purified using a DNA isolation kit (DNeasy Blood & Tissue kit; QIAGEN, Valencia, CA, USA). The 16S rDNA sequence was amplified by PCR using two universal primers specific for rDNA gene [12]. PCR amplification was performed using a Taq DNA polymerase kit (QIAGEN) under the following conditions: the reaction mixture was subjected to an initial denaturation at 94°C for 3 min, followed by 35 cycles, including 1 min at 94°C, 1 min at 52°C, and 2 min at 72°C. A final extension was carried out at 72°C for 10 min. The amplified PCR product was purified using a gel extraction kit (QIAquick, QIAGEN), ligated to the pGEM-T Easy vector (Promega, Madison, WI, USA), and introduced into competent E. coli DH5α cells by electroporation. The recombinant plasmid carrying the 16S rDNA fragment was isolated from overnight culture of E. coli using spin column (QIAprep Spin Miniprep kit, QIAGEN). Resultant plasmid DNA was sequenced using a 3730 DNA Analyzer (Applied Biosystems, Foster city, CA, USA) at the Plant-Microbe Genomics Facility at The Ohio State University.

2.3. Purification of Enterocin RM6 from Cultured Broth. An aliquot of 0.5 mL of OSY-RM6 overnight culture was inoculated into a 1-liter flask containing 500 mL MRS broth. The flask was incubated at 30°C for 18 hours without shaking. Cells in the cultured broth were removed by centrifugation at 15,180 x g for 15 min at 4°C (Sorvall RC-5B, DuPont, Wilmington, DE, USA). The supernatant was adjusted to pH 6.5 using 1.0 N NaOH and passed through a cationic exchange column (Macro-Prep High S support; Bio-Rad, Hercules, CA, USA) that was equilibrated with phosphate buffer (50 mM, pH 6.5). Enterocin RM6 was eluted with 1.0 M NaCl in phosphate buffer (50 mM, pH 6.5). The resultant eluate was subjected to solid phase extraction using ten C18 silica cartridges (Sep-Pak; Waters Corporation, Milford, MA, USA) that were head-to-tail connected to increase the binding capacity. Enterocin RM6 was eluted from the cartridges by 70% acetonitrile, and the solvents were removed by lyophilization. The crude extract (CE) of enterocin was obtained by dissolving the freeze-dried powder in high-performance liquid chromatography (HPLC) grade water.

Crude extract of enterocin was further purified by reverse-phase HPLC (Hewlett Packard 1050, Agilent Technologies, Palo Alto, CA, USA). Separation was achieved using a preparative column with 5 μm particle size (250 mm × 10 mm; Alltech Associates, Inc., Deerfield, IL, USA). The mobile phase consisted of (A) a mixture of isopropanol and acetonitrile (2:1, v/v) with 0.4% trifluoroacetic acid (TFA) and (B) HPLC grade water with 0.1% TFA. For each run, aliquots (300 μL) of crude extract were loaded and separated on the column by a linear gradient of 0 to 100% solvent A over 30 min, followed by 100% solvent A for 5 min at a flow rate of 1.5 mL/min. Elution was monitored using a UV-detector at a wavelength of 280 nm. Fractions from each minute were collected automatically using Waters Fraction Collector II (Waters Cooperation, Milford, MA, USA). Fractions with the same retention time from multiple runs were pooled and lyophilized; the resulting powder was dissolved in water. Antimicrobial activity of each faction was determined by microtiter plate bioassay. Active fraction against bacterial indicator strain was stored at 4°C for further analyses.

2.4. Antimicrobial Activity Determination and Inhibition Spectrum. Antimicrobial activity was determined as described by Yousef and Carlstrom [13]. L. cellobiosus OSU 919 was used as the indicator. Briefly, aliquots (10 μL) of overnight indicator bacterium were transferred into 9 mL of MRS soft agar (0.75%) that was held at ~50°C; the mixture was then poured onto a basal MRS agar plate. The tested compound or solution with enterocin RM6 was subjected to serial two-fold dilutions. Aliquots (5 μL) of diluted solution were spotted onto the soft agar layer seeded with indicator bacterium. After incubation at 30°C overnight, inhibitory areas were observed. Antimicrobial activity is expressed in arbitrary unit (AU/mL), which is the reciprocal of the highest dilution factor resulting in a clear inhibitory zone.

Antimicrobial activity of HPLC fractions was examined by microtiter plate bioassay. Briefly, overnight indicator culture was tenfold diluted using MRS medium, and 100 μL of diluted cell suspension was added in a 96-well plate. An equal volume (100 μL) of HPLC fraction was added to each well and incubated at 30°C for 5–8 hours, where sterile distilled water was used as negative control. Optical density at 600 nm (OD600) was measured using a spectrophotometric microplate reader (Vmax Kinetic Microplate Reader, Molecular Devices Corp., Menlo Park, CA, USA). Active fraction showed a lower OD600 value compared to the negative control because of the inhibitory effect against the indicator bacterium.

2.5. ESI-MS and MS/MS Analyses. Accurate molecular weight determination and further peptide sequence investigation of enterocin RM6 were performed on a mass spectrometer (LTQ orbitrap, ThermoFinnigan, West Palm Beach, FL, USA) operated in positive ion mode. Briefly, the sample diluted in the mixture of H2O-MeOH-HAc (50:50:2.5) was infused into the electrospray source at a 6 μL/min flow rate. To achieve the optimal electrospray, spray voltage was set at 2,000 V; source temperature was 175°C. The data were recorded between 400 and 2000 Da, and the resolution was set at 30000 to achieve high mass accuracy determination. The most abundant enterocin RM6 peak was isolated for further MS/MS study. The isolation window was set at 10 Da and the CID fragmentation energy was set to 35%. Data were acquired in continuum mode until well-averaged data were obtained.

2.6. Structural Gene of Enterocin RM6. Genomic DNA from strain OSY-RM6 was used as template for amplifying
3. Results

3.1. Strain Identification. The producer strain OSY-RM6 is a Gram-positive bacterium. Analysis of the 16S rDNA gene (accession number: KF154976) showed a 99% similarity to E. faecalis. The new isolate was designated as E. faecalis OSY-RM6. This strain was positive for acid production (pH 4.8) in Bertani medium.

3.2. Purification of Enterocin RM6. The procedure for preparing enterocin RM6 crude extract recovered ca. 24% total activity in the culture broth (Table 1). The crude extract was further purified by reverse-phase HPLC using a preparative column. The retention time of enterocin RM6 on a C18 column was 31.268 min (Figure 1).

3.3. Molecular Weight and Peptide Sequence Determination by MS and MS/MS. HPLC-purified enterocin RM6 was subjected to electrospray ionization mass spectrometry (ESI-MS) analysis for accurate molecular weight determination.

Table 1: Preparation of crude extract of enterocin RM6.

<table>
<thead>
<tr>
<th>Purification step</th>
<th>Volume (mL)</th>
<th>Arbitrary unit (AU/mL)</th>
<th>Total activity (×10^3 AU)</th>
<th>Recovery rate (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Culture supernatant</td>
<td>500</td>
<td>800</td>
<td>400</td>
<td>100</td>
</tr>
<tr>
<td>Cation exchange</td>
<td>75</td>
<td>1600</td>
<td>120</td>
<td>30</td>
</tr>
<tr>
<td>C18 silica cartridges</td>
<td>60</td>
<td>1600</td>
<td>96</td>
<td>24</td>
</tr>
</tbody>
</table>

*AU/mL: the reciprocal of the highest dilution factor showing a visible inhibitory zone.

The producer strain OSY-RM6 is a Gram-positive bacterium. Two primers (Ent48SF: 5'-GAGGAGTITCATGTTAAGAGGA-3' and Ent48SR: 5'-CATATTGTTAAATTACCAAGCAA-3') were used for PCR [14]. PCR amplification was performed using a Taq DNA polymerase kit (QIAGEN) under the following conditions: the reaction mixture was subjected to an initial denaturation at 95°C for 5 min, followed by 30 cycles, including 1 min at 95°C, 1 min at 52°C, and 1 min at 72°C. A final extension was carried out at 72°C for 10 min. The resultant PCR product was cloned in pGEM-T Easy vector for DNA sequencing.

2.7. Efficacy of Enterocin RM6 against L. monocytogenes in Inoculated Cottage Cheese. Twenty gram of cottage cheese (2% reduced milk fact, Kraft foods) was mixed with 30 mL of peptone water (0.1%), and the mixture was homogenized in a stomacher for 3 minutes. Enterocin crude extract was added to the diluted cheese sample at a final concentration of 80 AU/mL. Peptone water (0.1%) was used as negative control. The mixture was inoculated with L. monocytogenes Scott A at a final concentration of ~10^5 CFU/mL. Inoculated cheese samples were incubated at 35°C for 26 hours. Samples were analyzed at 0.5 h, 4 h, and 26 hours after treatment to examine the inhibitory effect of enterocin. Viable cells after treatment were counted using tryptic soy agar, PALCAM agar, and modified Oxford agar (Becton and Dickinson). Each treatment or control included two independent experiments.

Additional MS/MS analysis was also performed to obtain the amino acid sequence of enterocin RM6. As shown in Figure 2, peaks carrying different charge status ([m/z = 894.14], 1021.73, 1191.85, and 1300.02) were observed for intact enterocin RM6. After deconvolution, the average monoisotopic mass (M + H) of enterocin RM6 was calculated as 1146.08 AU. The most abundant peak ([m/z = 1191.85]) was then isolated and further fragmented by collision-induced dissociation (CID) to obtain the sequential information of this peptide. A series of MS/MS product ions were observed at [m/z = 1419.17], 1438.18, 1467.19, 1486.19, 1519.22, 1542.90, 1576.58, 1614.28, and 1680.98, which led to the identification of

Table 2: Antimicrobial spectrum of enterocin RM6.

<table>
<thead>
<tr>
<th>Strainsa</th>
<th>Mediab</th>
<th>Diameter of inhibitory zone (mm)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gram-positive bacteria</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Pediococcus acidilactici PO2</td>
<td>MRS</td>
<td>17.5</td>
</tr>
<tr>
<td>Pediococcus pentosaceus</td>
<td>MRS</td>
<td>16.8</td>
</tr>
<tr>
<td>Lactobacillus plantarum ATCC 8014</td>
<td>MRS</td>
<td>11.4</td>
</tr>
<tr>
<td>Lactobacillus casei ATCC 7469</td>
<td>MRS</td>
<td>13.4</td>
</tr>
<tr>
<td>Lactobacillus acidophilus ATCC 19992</td>
<td>MRS</td>
<td>10.0</td>
</tr>
<tr>
<td>Pediococcus cerevisiae</td>
<td>MRS</td>
<td>11.1</td>
</tr>
<tr>
<td>Lactobacillus cellobiosus OSU 919</td>
<td>MRS</td>
<td>14.8</td>
</tr>
<tr>
<td>Listeria innocua ATCC33090</td>
<td>TSBYE</td>
<td>7.4</td>
</tr>
<tr>
<td>Enterococcus faecalis ATCC 29212</td>
<td>MRS</td>
<td>8.4</td>
</tr>
<tr>
<td>Listeria monocytogenes Scott A</td>
<td>TSBYE</td>
<td>11.3</td>
</tr>
<tr>
<td>Bacillus cereus ATCC 14579</td>
<td>TSBYE</td>
<td>11.2</td>
</tr>
<tr>
<td>Bacillus cereus ATCC 11778</td>
<td>TSBYE</td>
<td>5.8</td>
</tr>
<tr>
<td>Staphylococcus aureus OSU 6538</td>
<td>NA</td>
<td>6.1</td>
</tr>
<tr>
<td>Staphylococcus aureus (methicillin sensitive)</td>
<td>NA</td>
<td>12.4</td>
</tr>
<tr>
<td>Staphylococcus aureus (methicillin resistant)</td>
<td>NA</td>
<td>10.0</td>
</tr>
<tr>
<td>Gram-negative bacteria</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Yeisina enterocolitica</td>
<td>TSBYE</td>
<td>—</td>
</tr>
<tr>
<td>Salmonella Typhimurium</td>
<td>LB</td>
<td>—</td>
</tr>
<tr>
<td>Escherichia coli O157:H7</td>
<td>LB</td>
<td>—</td>
</tr>
</tbody>
</table>

*Strains obtained from the culture collection of The Ohio State University Food safety laboratory. MRS: lactobacillus MRS broth; TSBYE: tryptic soy broth supplemented with 0.6% yeast extract; NB: nutrient broth; LB: Luria-Bertani medium.
a fragment with 12 amino acids: IVSILTAVGSGG or GGSGVATLISVI (note that L and I can be switched since they have identical masses) (Figure 3). This 12 amino acids sequence partially matches the sequence of a 70-residue cyclic peptide: AS-48 protein, whose complete sequence is MAKEFGIPAAVAGTVLNVVEAGGWVTTVSTLATAVG SGGLSLAAAGRESIKYLIKKEIKKGKRAVIAW, where a tail-to-head linkage is formed between the N-terminal methionine and the C-terminal tryptophan through the dehydration of one water molecule [14, 15]. The theoretical molecular weight of AS-48 protein is 7145.0718 Da, which matches exactly with the observed enterocin RM6 molecular weight (7145.0745 Da). Mass accuracy between theoretical and measured mass of enterocin RM6 is 0.44 ppm. In addition, more product ions were observed in the MS/MS spectrum (Table S1 in Supplementary Material available online at http://dx.doi.org/10.1155/2013/206917) which further supported that enterocin RM6 shares the same sequence with peptide AS-48 [14, 15]. Fragment ions were observed at \( m/z = 575.32, 646.36, 777.40, 963.48, \) and 1034.51, which corresponded to sequences KEFGI, AKEFGI, MAKEFGI, WMMAKEFGI, and AWMAKEFGI, respectively. This observation confirmed that the NH\(_2\) terminus in Met1 was linked with the COOH-terminus in Trp70 to form the head-to-tail cyclized peptide. During CID fragmentation, the cyclized peptide can be broken at different locations to generate different fragmentation patterns. For example, when the linkage was broken between Ile8 and Pro9, internal ions identified as PAAVA (\( m/z = 410.24 \)), PAAVAG (\( m/z = 467.26 \)), PAAVAGT (\( m/z = 568.31 \)), PAAVAGTV (\( m/z = 667.38 \)), and PAAVAGTVLNVVEAGGWVTIV (\( m/z = 1053.59^{2+} \)) were observed. Other fragment ions that were observed included ions that represent amino acid sequences KEFGI, AKEFGI, MAKEFGI, WMMAKEFGI, and AWMAKEFGI, as well as other amino acid sequences between SLAAA-

![Figure 1: High-performance liquid chromatography profile of the crude extract of enterocin RM6. The peak with retention time of 31.27 min (indicated by the arrow) showed antimicrobial activity against Lactobacillus cellobiosus OSU919.](image1)

![Figure 2: Electrospray ionization-mass spectrometry (ESI-MS) analysis of enterocin RM6.](image2)

3.4. Antimicrobial Spectrum of Enterocin. Enterocin RM6 is active against all tested Gram-positive bacteria, but it has no activity against Gram-negative bacteria (Table 2). *Pediococcus acidilactici* PO2 is the most sensitive strain to enterocin RM6. Moreover, this enterocin has strong activity against some important pathogens such as *L. monocytogenes* Scott A, *B. cereus* ATCC 14579, and methicillin-resistant *S. aureus* (MRSA).
3.5. Bactericidal Effect of Enterocin RM6 against Listeria in Cottage Cheese. When enterocin crude extract (final concentration, 80 AU/mL) was added to Listeria-inoculated cottage cheese, a 4-log reduction in pathogen population was observed within 30 min and viable cells were not detected after 26 hrs. In contrast, the population of L. monocytogenes without treatment increased to $10^7$ CFU/mL after 26 hrs at the same incubation conditions (Figure 5).

4. Discussions

Bacteriocins are ribosomally synthesized antimicrobial peptides produced by some Gram-positive bacteria. Many bacteriocins from Enterococcus spp. have been purified and characterized, such as enterocin A, B, P, and AS-48 [16]. In this study, we describe a new bacterial strain isolated from raw milk exhibiting strong antimicrobial activity. The new isolate was identified as E. faecalis by 16S rDNA sequencing and was designated as strain OSY-RM6. MS/MS analyses and structural gene sequencing confirmed that the antimicrobial activity.
activity was attributed to a cyclic peptide with 70 residues, whose chemical structure is the same as enterocin AS-48.

The efficacy of bacteriocins is usually affected by conditions in the food ecosystems such as food composition [7]. The antilisterial efficacy of enterocin RM6 was investigated in cottage cheese. The results indicated that enterocin RM6 has a rapid bactericidal activity against *L. monocytogenes* in cheese products. In all, enterocin RM6 may have a practical application in the food industry to control listerial contamination.

**Acknowledgment**

This research is supported by Kraft Foods, Inc.

**References**


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