# A novel pediocin-like bacteriocin produced by *Lactobacillus plantarum* KMITL-QU 54 isolated from traditional thai fermented meat-rice sausage (Sai-krog Isan)

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#### Abstract

A total of 150 strains of lactic acid bacteria (LAB) were randomly isolated from the 15 samples of traditional thai fermented meat-rice sausage (Sai-krog Isan) sold in Bangkok and screened for bacteriocinproducing strains. Only the LAB strain KMITL-QU 54, which implied to produce bacteriocin and displayed the strongest antilisterial effect as pediocin, was later identified at the DNA level as *Lactobacillus plantarum*. Molecular mass of the purification to homogeneity of the bacteriocin produced by this strain showed that it was 4,689 Da, which was about 60 Da higher than pediocin AcH/PA-1 (4,628 Da) produced by known pediocin producers of *Pediococcus acidilactici* H and *Lb. plantarum* WHE 92. The study of pediocin producing structural gene of this pediocin-like producing strain revealed that *Lb. plantarum* KMITL-QU 54 exhibited 96% and 94% of the structural genes identical to pediocin family of Coagulin (CoaA) produced by *Bacillus coagulans* I<sub>4</sub> and pediocin-like structural sequence genes from KMITL-QU 54 informed that its molecular weight calculation of the pediocin-like structural sequence genes from KMITL-QU 54 informed that its molecular weight was about 4,691 Da and more related to the prior purified bacteriocin (4,689 Da) than coagulin (4,615 Da) and Pediocin AcH/PA-1 (4,628 Da). By these results, hence, we realize that *Lb. plantarum* KMITL-QU 54 might be produced a novel pediocin-like bacteriocin and the strain could be applied as starter for microbiological quality enhancement during this traditional thai fermented meat product fermentation.

#### Introduction

Lactic acid bacteria (LAB) are widely used as starter cultures for reliable and consistent acid production in various fermented foods. The inhibition of other microorganisms may also occur by the formation of various compounds, which produces during fermentation. Among the variety of these inhibitory compounds synthesized by these LAB, bacteriocins have been received much attention in the past decade (Ennahar et al., 1996; Gaenzle et al., 1997; Swetwiwathana, 2005). Thus, an attempt on finding the most potent bacteriocin-producing LAB strains from fermented meat-rice sausage (Sai-krog Isan) and use of these potent bacteriocin producing strains as starter cultures in order to improve the quality and safety during the fermentation of this product had been studied and reported (Swetwiwathana et al., 2007). Only 2 strains of *Lactobacillus plantarum* RS-49 and RS-54 had been reported to produce bacteriocins was different from the known bacteriocin of nisin A and pediocin PA-1 which produced from *Lactococcus lactis* NCDO and *Pediococcus pentosaceus* TISTR 536, respectively. Hence, the further characterization of bacteriocin produced from RS-54 had been studied and reported in this paper. The code of strain had been changed from RS-54 to be KMITL-QU 54 according to the collaborative works between King Mongkut's institute of Technology Ladkrabang (KMITL) – Thailand and Kyushu University (QU) – Japan.

#### Materials and methods

**Bacteriocins purification :** The cell-free supernatant of 1 liters culture incubated at  $30^{\circ}$  C of KMITL-QU 54 was purified by a four step procedures as described by Ennahar *et al.* (1996). The final sample containing the purified bacteriocins was dried by Speed-Vac rotary evaporator (Savant Instruments) and stored at  $-20^{\circ}$ C for molecular mass determination.

*Mass spectrometric* : The molecular masses of purified bacteriocins were determined using a Accu TOF spectrometer, model JMS-T100LC (Agilent Technologies, Germany).

PCR analysis and DNA sequencing of bacteriocin genes from KMITL-OU 54 : The total DNA of KMITL-QU 54 was isolated by using the method described by Anderson and McKay (1983). Pediocin PA-1 primers designed and synthesized (Hokkaido System Science Co. Ltd., Hokkaido, Japan) for PCR amplification are Pedi-1F (5'-GAGTGGGAACTAGAATAAGCGCGTA-3') and Pedi-1R (5)TTACTCTTATTCATAAAATCACCCC -3'). DNA analyses and sequence alignments were carried out using GENETYX-WIN software (GENETYX, Tokyo, Japan). Database searches were performed using BLAST of the National Center for Biotechnology (NCBI, http://www.ncbi.nlm.nih.gov/BLAST/). Alignment analysis of performed ClustalW of the DNA of peptides was using Data Bank Japan (DDBJ, http://www.ddbj.nig.ac.jp/search/clustalw-e.html).

#### **Results and discussion**

*Molecular mass of bacteriocin produced by KMITL-QU 54 :* Bacteriocin produced by *Lb. plantarum* KMITL-QU 54 was showed the molecular mass at about 4,690 d (Fig. 1). This molecular mass of bacteriocin from KMITL-QU 54 revealed about 67 d different from the molecular mass of those pediocins produced by *P. pentosaceus* TISTR 536 (produced pediocin PA-1, molecular mass about 4,623 d – Swetwiwathana, 2005) and *Lb. plantarum* WHE 92, which was reported to produce pediocin AcH (molecular mass 4,623 d; Ennahar *et al.*, 1996).

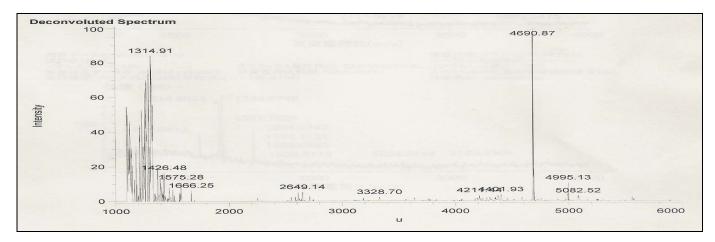


Figure 1. Molecualr mass of purified bacteriocin fraction from Lb. plantarum RS-54.

The results of pediocin producing structural genes from KMITL-QU 54 (Fig. 2) informed that KMITL-QU 54 exhibited 96% of pediocin-like structural genes identical to CoaA (Coagulin) produced by *Bacillus coagulans*  $I_4$  (Le Marrec *et al.*, 2000) and 94% of structural genes identical to pediocin AcH produced by *Pediococcus acidilactici*. The molecular weight calculation of this pediocin-like structural sequence genes from KMITL-QU 54 was later compared to both structural sequence genes from coagulin of *B. coagulans*  $I_4$  and pediocin of *Pediococcus acidilactic* (Fig. 3). It was informed that the pediocin-like structural sequence genes of *Lb. plantarum* KMITL-QU 54 exhibited the related molecular weight to the purified bacteriocin (4,691.1 d). We are, hence, realized that the bacteriocin from KMITL-QU 54 might be a novel pediocin-like bacteriocin. This bacteriocin is under studying the best purification methods for further amino acid sequence step.

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Alignments	入力されたアミノ酸配列
<pre>&gt;gi 11066919 gb AAG28763.1 AF300457_1 CoaA [Bacillus coagulans] Length=62 Score = 124 bits (312), Expect = 7e-27 Identities = 54/56 (96%), Positives = 55/56 (96%), Gaps = 0/56 (0%) Frame = +2 Query 473 LTEKEMANIIGGKYYGNGLYCGKHSCSVDWGKATTCIINNGAMAWATGGHQGTHKC 640 LTEKEMANIIGGKYYGNG+ CGKHSCSVDWGKATTCIINNGAMAWATGGHQGTHKC 62 Sbjct 7 LTEKEMANIIGGKYYGNGVTCGKHSCSVDWGKATTCIINNGAMAWATGGHQGTHKC 62</pre>	KYYGNGLYCGKHSCSVDWGKATTCIINNGAMAWATGGHQGTHKC         分子量         4691.1 Da
<pre>&gt;gi 31790963 ref NF_857602.1  G PapA {Pediococcus acidilactici] gl 548575 ap F29430 FPA1 PEDAC G Bacteriocin pediocin PA-1 precursor (Pediocin ACH) gl 152643 ab AAA955551] G bacteriocin operon-associated ORF gl 152643 ab AAA9337.1  pediocin ACH gl 407297 ab AAC43293.1  G PapA [Pediococcus acidilactici] gl 32654803 ab AAP63155.1  PapA [Pediococcus parvulus] gl 32654803 ab AAP63155.1  PapA [Dediococcus parvulus] gl 32654803 ab AAP63155.1  PapA [Dediococcus acidilactici] gl 51201243 ab AAP63157.1  PapA [Dediococcus acidilactici] gl 51201243 ab AAP63157.1  PapA [Dediococcus acidilactici] gl 51201243 ab AAP63422.1  G PedA [Pediococcus acidilactici] gl 61243126]prf  1918342A G pediocin Length=62</pre>	<u>入力されたアミノ酸配列</u> KYYGNGVTCGKHSCSVDWGKATTCIINNGAMAWATGGHQGTHKC <u>分子量</u>
Score = 122 bits (307), Expect = 3e-26 Identities = 53/56 (94%), Positives = 54/56 (96%), Gaps = 0/56 (0%) Frame = 42 Query 473 LTEREMANIIGGRYYGNGLYCGKHSCSVDWGKATTCIINNGAMAWATGGHGGTHKC 640 LTEREMANIIGGRYYGNG+ CGKHSCSVDWGKATTCIINNGAMAWATGGHGGTHKC 640	4615 Da Coagulin
Sbjet 7 LTEKEMANIIGGKYYGNGVTCGKHSCSVDWGKATTCIINNGAMAWATGGHQGNHKC 62 >g[ 18390350 gb AAC60413.2] prepediocin AcH [Pediococcus acidilactic] Length=62 Score = 119 bits (298), Expect = 3e-25 Identities = 52/56 (92%), Positives = 53/56 (94%), Gaps = 0/56 (0%)	入力されたアミノ酸配列 KYYGNGVTCGKHSCSVDWGKATTCIINNGAMAWATGGHQGNHKC
Juericities     53/36     53/36     (348), Gaps     0/36     (08)       Prame     +2       Query     473     LITEKEMANIIGGKYYGNGLYCGKHSCSVDWGKATTCIINNGAMAWATGGHQGTHKC     640       LITEKEMANIIGGKYYGNG+     CGKHSCSVDWGKATTCIINNGAMAWATGGHQG     KC       Sbjet     7     LITEKEMANIIGGKYYGNGVTCGKHSCSVDWGKATTCIINNGAMAWATGGHQGNDKC     62	分子量 4628 Da Pediocin AcH/PA-1

## Figure 2. Pediocin-like structural genes of *Lb. plantarum* KMITL-QU 54.

**Figure 10.** Comparison of molecular weight from pediocin-like structural sequence genes of *Lb. plantarum* KMITL-QU 54, coagulin and pediocin AcH/PA-1.

#### Conclusions

The study informs that *Lb. plantarum* KMITL-QU 54 might be produced a novel pediocin-like bacteriocin with MW about 4,691 d. Moreover, the strain could be applied as starter for microbiological quality enhancement during this traditional thai fermented meat product fermentation.

#### Acknowledgements

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