

# Lytic Activity of LysH5 Endolysin Secreted by *Lactococcus lactis* Using the Secretion Signal Sequence of Bacteriocin Lcn972

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**Bacteriophage endolysins have an interesting potential as antimicrobials. The endolysin LysH5, encoded by *Staphylococcus aureus* phage vB\_SauS-phi-IPLA88, was expressed and secreted in *Lactococcus lactis* using the signal peptide of bacteriocin lactococcin 972 and lactococcal constitutive and inducible promoters. Up to 80 U/mg of extracellular active endolysin was detected in culture supernatants, but most of the protein (up to 323 U/mg) remained in the cell extracts.**

*Staphylococcus aureus* is a cause of serious concern in clinical settings especially due to the emergence of methicillin-resistant *S. aureus* (MRSA) and vancomycin-resistant *S. aureus* (VRSA) (13). Moreover, *S. aureus* is a threat to food safety (3). In the dairy environment, this pathogen is recognized as a frequent cause of subclinical intramammary infections in dairy cows (28). Bacteriophage endolysins mediate lysis of the host bacteria at the end of the lytic cycle to release phage progeny (39). They have a huge potential as novel therapeutic antibacterial agents (7), as biopreservatives in foods (27), in pathogen detection (38), and as disinfectants of industrial facilities (32).

*Lactococcus lactis*, a GRAS (generally regarded as safe) microorganism with a long history of safe use in food fermentations, has been proven as a suitable host for expression and purification of heterologous proteins with food safety applications such as bacteriocins (1, 23) and bacteriophage lytic enzymes against *Listeria monocytogenes* (8, 36) and *Clostridium difficile* (24). To facilitate secretion, proteins have been fused to bacteriocin signals (12) or to the signal peptide (SP) of Usp45 (SP<sub>usp45</sub>), the major Sec-dependent protein secreted by *L. lactis* (1). The capacity of this microorganism for protein secretion can be exploited to improve large-scale production processes and downstream purification steps (19).

LysH5 endolysin from *S. aureus* bacteriophage vB\_SauS-phi-IPLA88 has been characterized. LysH5 is a 53.7-kDa protein, encoded by a 1,446-bp gene (10), which lysed a wide range of staphylococci and also inhibits *S. aureus* growth in milk (11, 27).

The aim of this work was cloning and expression of the LysH5-encoding gene in an *L. lactis* strain under the control of lactococcal inducible and constitutive promoters to facilitate the potential application of this endolysin as a food preservative. For secretion, the signal peptide (SP<sub>Lcn972</sub>) of the bacteriocin lactococcin 972 (Lcn972), which shows a Sec-dependent processing signal (22), has been tagged with LysH5 endolysin.

**Heterologous expression of the endolysin LysH5 in *L. lactis*.** Prior to the construction of an *L. lactis* secretion system for the endolysin LysH5, we assessed the ability of this bacterium to express active LysH5 protein. Phage phiIPLA88 DNA (9) was used to amplify the LysH5-encoding gene with primers AMI-1 and AMI-2 (see Table S1 in the supplemental material). PCR amplifications were carried out using the PureTaq Ready-To-Go PCR bead kit (GE Healthcare, United Kingdom), and PCR fragments were purified using the GenElute PCR cleanup kit (Sigma, St. Louis, MO). The resulting PCR product was cloned into the plasmid

pNZ8020 (5) using the restriction sites PstI and BamHI. The recombinant plasmid pNZ8020-LysH5, bearing the LysH5-encoding gene under the control of the P<sub>nisA</sub> promoter, was transformed into *L. lactis* NZ9000 (15). This recombinant plasmid and those constructed thereof were confirmed by DNA sequencing. Aliquots of cultures *L. lactis* NZ9000(pNZ8020) and *L. lactis* NZ9000(pNZ8020-LysH5) at an optical density at 600 nm (OD<sub>600</sub>) of 0.4 were induced for 4 h at 30°C with nisin A (10 ng/ml). Cells from 10-ml cultures were resuspended in 50 mM sodium phosphate buffer, pH 7, and lysed using glass beads in a FastPrep system (ThermoSavant-Bio 101/Q-Biogen, Holbrook, NY). Lysates were centrifuged at 10,000 × g for 10 min at 4°C and tested for lytic activity by turbidity reduction assay against *S. aureus* Sa9 cell suspensions (27). A specific activity of 166.85 ± 34.8 U/mg (Table 1) was observed, indicating that LysH5 is synthesized in *L. lactis* as an active protein. No lysis of *S. aureus* Sa9 was detected in extracts of the control strain *L. lactis* NZ9000(pNZ8020).

**Lcn972 signal peptide (SP<sub>Lcn972</sub>) leads to LysH5 secretion by *L. lactis*.** To obtain a recombinant strain able to release active LysH5 into the culture supernatants, a fusion between SP<sub>Lcn972</sub> and the complete LysH5 was designed. Plasmid pBL1 (31) was used as the template to amplify an 81-bp fragment containing SP<sub>Lcn972</sub> using the primers Lcn-AMI and Lcn-AMI-3 (see Table S1 in the supplemental material). A PCR fragment containing the LysH5-encoding gene was obtained after amplification of phage phiIPLA88 DNA with primers AMI-2 and AMI-7 (see Table S1). The two PCR fragments were merged by a PCR with primers AMI-2 and Lcn-AMI (see Table S1). The resulting PCR product (1,566 bp) was cleaved with BamHI and PstI and cloned into plasmid pNZ8020 to generate the recombinant plasmid pNZ8020-SP<sub>Lcn972</sub>-LysH5. Transformants containing this plasmid were induced with nisin A as indicated above. Expression and secretion of LysH5 by *L. lactis* NZ9000 under the control of the nisin A promoter (P<sub>nisA</sub>) were tested in both cell extracts and culture super-

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TABLE 1 Endolysin LysH5 production in supernatants and cell extracts from recombinant strains

Lactococcal strain	Lytic activity (sp act, U/mg)	
	Supernatant	Cell extract
<i>L. lactis</i> NZ9000(pNZ8020)	0	0
<i>L. lactis</i> NZ9000(pIL252)	0	0
<i>L. lactis</i> NZ9000(pMG36c)	0	0
<i>L. lactis</i> NZ9000(pNZ8020-LysH5)	0	166.85 ± 34.8
<i>L. lactis</i> NZ9000(pNZ8020-SP <sub>Lcn972</sub> -LysH5)	40 ± 7	116.36 ± 9.6
<i>L. lactis</i> NZ9000(pNZ8020-SP <sub>Lcn972</sub> -LysH5opt)	80 ± 3	323.63 ± 17.1
<i>L. lactis</i> NZ9000(pIL252-P <sub>llmg0169</sub> -SP <sub>Lcn972</sub> -LysH5opt)	0.01 ± 0.005	3.63 ± 0.5
<i>L. lactis</i> NZ9000(pMG36c-P <sub>32</sub> -SP <sub>Lcn972</sub> -LysH5opt)	0.01 ± 0.003	3.34 ± 1.1

nanatants by immunoblotting and lytic activity assays. Rabbit polyclonal antibodies were raised against purified LysH5 according to a previous protocol (30). Cell extracts and supernatants from induced cultures were subjected to Western blotting and immunological detection (chemiluminescent Western blotting kit; Roche) using anti-LysH5 (1:4,000 dilution). As shown in Fig. 1A (lane 2), a single protein band of about 53 kDa reacted with the antibody in the cell extract from *L. lactis* NZ9000(pNZ8020-SP<sub>Lcn972</sub>-LysH5). A weak band was also detected in the corresponding supernatants. These results agree well with the predicted mass of LysH5. The functional expression of LysH5 was determined and quantified by using *S. aureus* Sa9 cell suspensions in a turbidity reduction assay (Table 1). The cell extract from *L. lactis* NZ9000(pNZ8020-SP<sub>Lcn972</sub>-LysH5) showed a significantly stronger activity than did the supernatant. Although several examples have been published supporting the idea that the best production yields in *L. lactis* are obtained when proteins are secreted to the culture medium (19), LysH5 production did not follow this pattern. It is well known that protein size, the nature of the signal peptides (SPs), and the presence of propeptides are factors that may interfere with protein secretion (4). Moreover, secretion efficiency might be also influenced by the proper combination between the mature protein and the SP used to drive secretion (20).

Regarding the low specific activity of LysH5 in the supernatants of *L. lactis* cultures, this could be due, at least partially, to the activation of housekeeping proteases by the stress response to overexpression of secreted proteins (2). In fact, the inactivation of

HtrA, the unique housekeeping protease in the *L. lactis* cell surface, results in higher heterologous protein yields (26). The low specific activity of the extracellular LysH5 may also be ascribed to inactivation by low pH as previously observed (27).

In order to improve these results, we proceeded to the codon optimization of the SP<sub>Lcn972</sub>-LysH5-encoding gene based on the *Lactococcus lactis* subsp. *cremoris* MG1363 codon usage (Genescript) (17). The resulting DNA fragment was cloned into BamHI/PstI restriction sites from pNZ8020 to generate pNZ8020-SP<sub>Lcn972</sub>-LysH5opt. Expression of the optimized version of the LysH5 gene under the control of the P<sub>nisA</sub> promoter was evaluated (Table 1; Fig. 1, lanes 1). The codon optimization of the LysH5-encoding gene resulted in higher levels of endolysin activity in both cell extracts and supernatants from *L. lactis* NZ9000(pNZ8020-SP<sub>Lcn972</sub>-LysH5opt). Increases of 2.7- and 2-fold in the cell extract and supernatant activities, respectively, were obtained over the wild-type version of the LysH5 gene (Table 1; Fig. 1, lanes 2).

**Induction of LysH5 production by coculturing with the nisin Z producer strain *L. lactis* IPLA 729.** In regard to the industrial production of LysH5, nisin addition remains costly. Therefore, the activity of LysH5 produced by cultures of *L. lactis* NZ9000 carrying pNZ8020 derivatives in the presence of the nisin Z producer strain *L. lactis* IPLA 729 (29) was assessed as an alternative approach. Initially, we determined the survival of the *L. lactis* NZ9000 strains in the presence of *L. lactis* IPLA 729. We defined the optimal coculture conditions as an inoculum of 0.01% (vol/vol) *L. lactis* IPLA 729 ( $3 \times 10^6$  CFU/ml) and 3% (vol/vol) *L. lactis* NZ9000(pNZ8020-LysH5) or *L. lactis* NZ9000(pNZ8020-SP<sub>Lcn972</sub>-LysH5opt) ( $3 \times 10^7$  CFU/ml) (data not shown). Then, *L. lactis* IPLA 729 and *L. lactis* NZ9000(pNZ8020) derivatives were grown to stationary phase, centrifuged, and resuspended at the same cell density. Cocultures (20 ml) of *L. lactis* NZ9000(pNZ8020) derivatives (3%, vol/vol) and *L. lactis* IPLA 729 (*nisZ*<sup>+</sup>) (0.01%, vol/vol) were incubated for 4 and 6 h at 30°C. To determine the viable counts of each strain, culture dilutions were plated onto GM17 and GM17 containing chloramphenicol (10 µg/ml). LysH5 activity was tested in both cell extracts and supernatants of cocultures. LysH5 activity was detected in the cell extract of *L. lactis* NZ9000(pNZ8020-LysH5), the highest specific activity being  $17.74 \pm 0.85$  U/mg after 4 h of incubation (Fig. 2) and matching the highest nisin Z concentration in the supernatants (400 AU/ml). This result suggests that the bacteriocin secreted by the nisin Z-producing strain induced production of the endolysin. A similar result was obtained when cocultures of *L. lactis* NZ9000(pNZ8020-SP<sub>Lcn972</sub>-LysH5opt) and *L. lactis* IPLA 729 were used (data not shown). However, no LysH5 activity was detected in the supernatant from these cocultures, although the presence of LysH5 was detected by Western

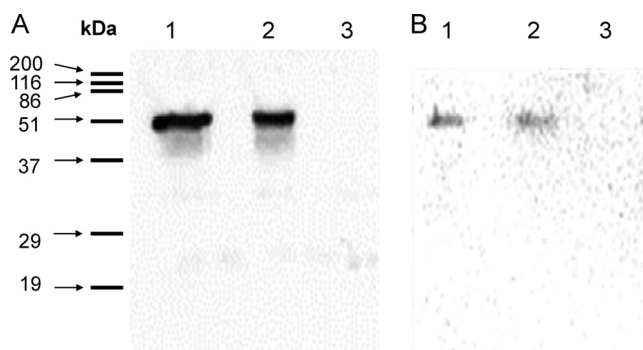


FIG 1 Detection of heterologously synthesized LysH5 by immunoblotting in cell extracts (A) and in supernatants of cultures (B) of *L. lactis* NZ9000(pNZ8020-SP<sub>Lcn972</sub>-LysH5opt) (lanes 1), *L. lactis* NZ9000(pNZ8020-SP<sub>Lcn972</sub>-LysH5) (lanes 2), and *L. lactis* NZ9000(pNZ8020) (lanes 3). Protein molecular mass markers with sizes in kilodaltons are indicated in the left margin.

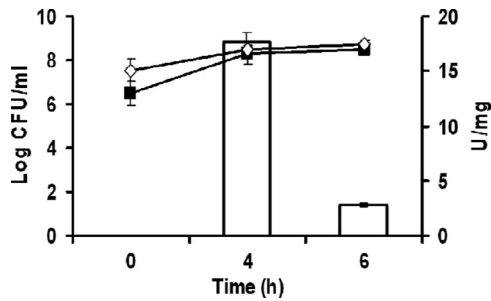


FIG 2 Coculture of *L. lactis* NZ9000(pNZ8020) derivative with *L. lactis* IPLA 729 (nisin Z producer). Survival of both strains, *L. lactis* NZ9000(pNZ8020-LysH5) (◇) and *L. lactis* IPLA 729 (■), was monitored as viable counts at different incubation times. Bars indicate the endolysin LysH5 activity expressed by *L. lactis* NZ9000(pNZ8020-LysH5) in coculture with *L. lactis* IPLA 729.

blot analysis (data not shown). This could indicate that secreted nisin Z was not enough to induce LysH5 in *L. lactis*(pNZ8020-Lcn-LysH5opt). Therefore, further coculturing conditions and alternative nisin Z producers should be tested.

**LysH5 expression under different *L. lactis* promoters.** For a constitutive secretion of LysH5, the plasmid pGM36c (37) containing the P<sub>32</sub> promoter was used. Primers Lcn-Lys-1 and Lcn-Lys-2 (see Table S1 in the supplemental material) were used to amplify a 1,573-bp fragment from plasmid pNZ8020-SP<sub>Lcn972</sub>-LysH5opt. The PCR-generated fragment was cloned into SacI/PstI restriction sites of plasmid pMG36c to generate the recombinant plasmid pMG36c-P<sub>32</sub>-SP<sub>Lcn972</sub>-LysH5opt. The heterologous production of LysH5 was determined in both supernatants and cell extracts. The results showed a very low production of LysH5 in both the cell extracts and the supernatants (Table 1). To test a presumably more efficient promoter, the inducible P<sub>llmg0169</sub> promoter of the *L. lactis* MG1363 *llmg0169* gene was amplified from plasmid pAB0169 (21) using primers pAB0169BamHI and pAB2164SmaI (see Table S1). The PCR-generated fragment (241 bp) was cloned into the BamHI/SmaI restriction sites of plasmid pIL252 (33). Then, the gene encoding SP<sub>Lcn972</sub>-LysH5opt fusion protein was cloned into BamHI/PstI restriction sites from plasmid pIL252-P<sub>llmg0169</sub>, resulting in the plasmid pIL252-P<sub>llmg0169</sub>-SP<sub>Lcn972</sub>-LysH5opt. With the purpose of increasing the transcription under the P<sub>llmg0169</sub> promoter, transformed cultures (OD<sub>600</sub> 0.4) were induced with bacitracin (1 μg/ml) for 4 h at 30°C. As occurred when the P<sub>32</sub> promoter was used, *L. lactis*(pIL252-P<sub>llmg0169</sub>-SP<sub>Lcn972</sub>-LysH5opt) extracts also showed low endolysin activity, whereas only scant lytic activity was observed in the supernatants (Table 1). The lower production of LysH5 by these derivatives than by the pNZ8020-based plasmids may be due to copy number differences but, more likely, is caused by the intrinsic differences among promoters used to drive gene expression (6, 14). Thus, the highest endolysin activity levels were obtained under the transcriptional control of P<sub>nisA</sub>, the promoter of the NICE system, which has been extensively used to produce proteins in *L. lactis* (16, 19, 25). In contrast, LysH5 activity was dramatically reduced in the constructs carrying the constitutive P<sub>32</sub> promoter, which was previously used to successfully drive the cytoplasmic production of the *Listeria* endolysins Ply118 and Ply511 in *L. lactis* (8).

Production of heterologous proteins is determined by a number of host genes. Different *L. lactis* mutants with higher lyso-

staphin and endolysin Ply511 activities were obtained by insertions in four different genes related to cell wall biosynthesis (35). In addition, other tools to enhance protein secretion have been developed, such as the fusion of a short synthetic propeptide between the SP and the mature moiety (18). Recently, the overexpression of the lactococcal CsiA protein, which inhibits the last stages of cell wall biosynthesis, promoted the efficient release of the heterologous intracellular *Listeria* endolysin LM4 in its active form without affecting cell viability (34). In conclusion, the synthesis of active LysH5 in *L. lactis* has been achieved, supporting the use of this GRAS host for its production as a food additive or for therapeutic purposes. However, further optimization to enhance secretion that facilitates downstream purification is needed for cost-effective LysH5 large-scale production.

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