ISOLATION OF LACTIC ACID BACTERIA FROM MALAYSIAN NON-BROILER CHICKEN (GALLUS GALLUS) INTESTINE WITH POTENTIAL PROBIOTIC FOR BROILER FEEDING

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ABSTRACT: Probiotic supplement can function as substitute for antibiotics especially in the broiler chicken feeding which can form an integral part of organic farming. Broiler forms one of an important protein source in South East Asia. Lactic acid bacteria (LAB) are important inhabitants of animal intestine and are useful source of probiotic microorganisms. Non-broiler chicken could be an ideal source of probiotic microorganisms that can be utilized for large scale broiler feeding. Our studies have successfully identified, through morphological and biochemical tests, 11 LAB isolates from gastrointestinal tract of local non-broiler chicken (Gallus gallus). These isolates have the ability to utilize lactose as part of their metabolism process and all showed negative reactions on catalase test. Out of the eleven (11) isolates, three (3) isolates were Gram-positive cocci and remaining isolates were of Gram-positive bacilli. Three isolates (E4, E11 and E17) showed at least 10 mm inhibitory effects on disc diffusion test against pathogenic bacteria Salmonella typhimurium. The partial 16S rRNA gene sequencing showed that one isolate (E17) has 89% similarity with Lactobacillus rhamnosus. These LAB strains isolated from Malaysian domestic non-broiler chicken gastrointestinal tract can potentially be used as a component for probiotics formulation in poultry feeding.

ABSTRAK: Makanan tambahan probiotik boleh berfungsi sebagai pengganti antibiotik terutamanya dalam pemakanan ayam pedaging yang akan membentuk bahagian kamiran dalam penternakan organik. Ayam pedaging merupakan sumber protein penting di Asia Tenggara. Bakteria asid laktik (lactic acid bacteria (LAB)) merupakan penghuni penting dalam usus haiwan dan merupakan sumber penting dalam mikroorganisma probiotik. Ayam bukan pedaging sesuai dijadikan sumber mikroorganisma probiotik agar dapat digunakan sebagai pemakanan ayam pedaging secara besar-besaran. Kajian telah berjaya mengenal pasti, melalui kaedah morfologi dan uji kaji biokimia, 11 LAB diasingkan daripada salur gastrousus ayam bukan pedaging (Gallus gallus). Pencilan ini berupaya menggunakan laktosa sebagai sebahagian daripada proses metabolisma dan semuanya menunjukkan kesan negatif terhadap ujian katalase. Daripada sebelas (11) pencilan, tiga (3) pencilan adalah kokus gram-positif dan pencilan yang lainnya adalah basilus grampositif. Tiga pencilan (E4, E11 dan E17) menunjukkan sekurang-kurangnya 10 mm kesan rencatan terhadap ujian resapan cakera terhadap bakteria patogen Salmonella typhimurium. Sebahagian daripada urutan gen 16S rRHA menunjukkan bahawa satu pencilan (E17) mempunyai 89% persamaan dengan Lactobacillus rhamnosus. Strain LAB ini diasingkan daripada salur gastrousus ayam tempatan bukan pedaging, sesuai digunakan sebagai komponen untuk rumusan probiotik dalam pemakanan ayam-itik.

KEYWORDS: lactic acid bacteria; probiotic; non-broiler chicken; Lactobaccilus rhamnosus

1. INTRODUCTION

The variable use of probiotics as a daily supplement has become a popular routine in commercial poultry feeding, particularly following antibiotic treatment. Lactic acid bacteria or LABs are the most common microbes employed as probiotics and these strains are usually of Lactobacillus sp, Bifidobacterium sp and Enterococcus sp [1]. Two principal kinds of probiotic bacteria, the members of the genera Lactobacillus and Bifidobacterium, have been studied in detail [2]. These are Gram-positive lactic acidproducing bacteria that constitute a major part of the normal intestinal microflora in animals and humans. These friendly bacteria play key role in enhancing the resistance to colonization by potentially pathogenic organisms in the gut. Bacterial food-borne diseases in chicken are the major problem affecting the productivity in poultry industry. Commonly, the control of this disease is based on the administration of antibiotics. Meanwhile, the use of antibiotic as growth promoter is so extensive in poultry and pig industries [3,4]. However, such treatment could develop resistant bacteria [5,6] and result in antibiotic residues leftover in chicken which is a potential hazard to public health. Furthermore, the normal microflora which is beneficial to chicken may also be killed or inhibited due to the administration of antibiotics [7] and a new approach of using probiotic is gaining acceptance [8]. In the recent years, there are great interests in the use of probiotic in poultry feeding to improve disease resistance and growth performance. Probiotic bacteria competed with the pathogen for nutrient, stimulated the immune system [7] and showed good colonizing and antagonistic properties against the pathogens [9].

However, the proper probiotic supplements are relatively expensive, and different studies have used different plethora of LAB species and strains. In some countries, successful probiotic supplement could be subjected to trade secret and consequently, commercially available probiotics formulation may not contain the most appropriate strains or doses. As a result, the extensive use of probiotic is not quantitative and economically ineffective at the larger scale. Due to the LAB diversity and poorly understood interactions between the microbes and intestine, determining the correct types and quantity of probiotic for broiler feeding. Malaysian non-broiler chicken could be a good source of probiotic for broiler feeding. Malaysian non-broiler chicken referred as '*ayam kampung*' is usually fed with household foods, and these are usually free of antibiotic. In this work, LAB strains from the intestinal tissue of non-broiler chickens were screened and characterized. It is hope that the isolated LAB can potentially be used as supplement to the young or growing broiler.

2. MATERIALS AND METHODS

A selected non-broiler chicken was slaughtered and it intestinal tissues was dissected and collected. The intestinal tissue was cut and suspended into 5 ml phosphate buffered saline PBS buffer and briefly vortexed. The tissue was grinded until homogenised and the solution kept as a stock. This was serially diluted in PBS and each dilution was plated on MRS agar followed by incubation at 37 °C overnight, anaerobically. MRS broth and media contained 1.0 % peptone, 0.8% meat extracxt, 0.4 yeast extract, 2.0% glucoase, 0.5 % sodium acetate, 0.1 % Tween 80, 0.2 % dipotassium hydrogen phosphate, 0.2 % triammonium citrate, 0.02 % magnesium sulfate and 0.005 % manganese sulfate. Single colonies were sub-cultured and kept in 80 %v/v glycerol and stored at -80°C for future identification processes. Gram staining, lactose utilization test, catalase tests were carried out on the selected colonies. In lactose utilization test, lactose and few drops of bromocresol purple was added into nutrient agar (NA).

Genomic DNAs were extracted from 11 isolated lactic acid bacteria strain. DNA was extracted using method as outlined in the protocol on DNeasy® Blood and Tissue Extraction Kit (QIAGEN). The 16S ribosomal gene was amplified by PCR using the universal primers pA forward (5'-AGAGTTTGATCCTGGCTCAG-3') and pE reverse (5'-CCGTCAATTCCTTTGAGTTT-3'). The PCR products were then loaded in 1.0 % agarose gel for electrophoresis and visualized on under UV transilluminator.

For detection of antimicrobial activity, a disc diffusion method was used. Sterile filter discs (12 mm) were dipped into the culture broth (MRS broth) of lactic acid bacteria, incubated for 42 hours and placed on solidified nutrient agar NA (OXOID) which was seeded with 12 to 14 hours cultures of *Salmonella typhimurium* (pathogenic bacteria). The plates were stored at 4 °C for 3 to 4 hours to permit diffusion, and finally incubated at 37 °C for 14 to 16 hours. Disc dipped in uninoculated MRS broth was served as control. Zones of inhibition observed were then measured [11, 12]. Disk impregnated in 0.4 mg/ml Streptomycin was taken as control.

3. RESULTS AND DISCUSSION

All isolates appeared positive in lactose utilisation test. These bacteria were able to ferment lactose to produce lactic acid that lowers the pH of the media that, in turn, changed the purple indicator dye to yellow indicative of fermentation activities (Fig. 1a). Gram reaction and morphology studies showed that out of eleven (11) isolates, three (3) isolates were Gram-positive bacilli and eight (8) isolates were Gram-positive cocci (Table 1 and Fig. 1b). In the antimicrobial test, all eleven (11) isolates produced antimicrobial compound that were active against pathogenic bacteria *Salmonella typhimurium* strains. This was indicated by the formation of clear inhibitory zones near the diffusion spots (Fig. 2). As also shown in Table 1, sample E4, E11 and E17 produced more that 10 mm (12-15 mm) diameter zones of inhibition; while all other samples produced less than 10 mm zone of inhibitions. Several factors could cause these inhibitions. Other than the production of organic acid, the antagonistic activity of LAB towards pathogens was also due to the production of bactericidal substances like bacteriocins [1].

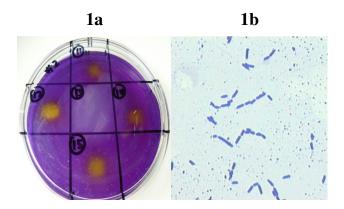


Fig. 1: a) Lactic acid bacteria changed the agar from purple to yellow indicative for lactic acid fermentation and formation of lactate;b) Morphology and gram staining of isolate 17 showing gram positive coccobacilli (oil immersion, 1000x magnification).

Isolates	Gram Reaction	Shape	Catalase	Lactose	Zone of inhibition*
E2	+	Cocci	-	+	+
E4	+	Cocci	-	+	++
E10	+	Bacilli	-	+	+
E11	+	Cocci	-	+	++
E12	+	Cocci	-	+	+
E14	+	Cocci	-	+	+
E15	+	Cocci	-	+	+
E17	+	Bacilli	-	+	++
E18	+	Cocci	-	+	+
E19	+	Cocci	-	+	+
E24	+	Bacilli	-	+	+

Table 1: Biochemical and morphological test for isolates from chicken intestine.

*Using Salmonella thyphimurium as indicator

+ < less than 10 mm

++> more than 10 mm

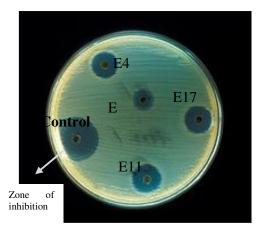


Fig. 2: Lactic acid bacteria against *Salmonella typhimurium* strains in disk diffusion method. Zone of inhibitions form surrounding the disk impregnated with sample LAB cultures E4, E11, E24 and E17.

As shown in Table 2, the result ribosomal rRNA sequence from one of the sample (isolate E17) produced 1.5 kb fragment (Fig. 3). The sequence indicated 89 % similarity to *Lactobacillus rhamnosus* GG (ATCC 53103) (Table 2). The other closely similar sequences were *Lactobacillus plantarum* followed by other common LAB, *Lactobacillus casei* and others. *Lactobacillus rhamnosus* GG was isolated from healthy human intestinal tract. This strain has however, established clinical health effects to human that prevents <u>urogenital tract</u> infection from pathogens, human diarrhea [13-15] and respiratory infections risk in children

[16]. It was shown that a strong antimicrobial activity of *Lactobacillus rhamnosus* GG against *Salmonella typhimurium* is due to the accumulation of lactic acid produced [17]. The usefulness of *Lactobacillus rhamnosus* strain in formulating probiotic feeding was previously suggested [1].

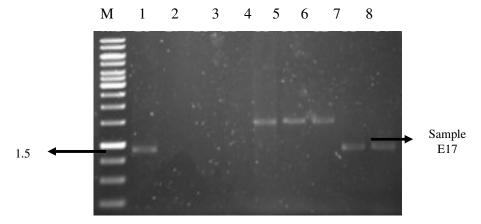


Fig. 3: The polymerase chain reaction amplification of ribosomal RNA gene using rRNA universal primer . Lane M in 1 kb ladder (Fermentas). Lane number 1 to 9 correspond to genomic sample from the 9 of the isolates, of which lane 9 is for isolate E17.

Strain from NCBI databse	Reference	Score	E-value
Lactobacillus rhamnosus GG	NC 013198.1	477	1e-133
Lactobacillus plantarum	NZ ACGZ01000098.1	473	2e-132
Lactobacillus plantarum JDM1	NC 012984.1	473	2e-132
Lactobacillus plantarum wcfS1	NC 004567.1	472	6e-132
Lactobacillus rhamnosus LMS2-1	NZ ACIZ01000148.1	472	6e-132
Lactobacillus rhamnosus HN001	NZ ABWJO1000068.1	472	6e-132
Lactobacillus rhamnosus Lc705	NC 013199.1	472	6e-132
Lactobacillus casei BL23	NC 010999.1	472	6e-132
Lactobacillus casei ATCC334	NC 008526.1	472	6e-132
Lactobacillus salivarius ATCC 11741	NC 007929	459	5e-128
Lactococcus lactis KF 147	NC 013656.1	361	1e-98

Table 2: The NCBI BLAST hit list for rRNA sequence from Sample E17 isolate.

4. CONCLUSION

In this work, we have successfully isolated lactic acid bacteria (LAB) strain from the gastrointestinal tract of Malaysian domestic non-broiler chicken. This isolated bacteria demonstrated common probiotic properties. Out of 25 isolates, 11 isolates have the ability to utilize lactose as part of their metabolism process, all showed negative reactions towards

catalase test and active against pathogenic bacteria *Salmonella typhimurium*. Three out of the 11 isolates were Gram-positive cocci and remaining 8 isolates were Gram-positive bacilli. Based on ribosomal RNA sequencing, one of the isolated bacilli have 89% similarity to *Lactobacillus rhamnosus*. While this strain was established it uses as probiotic in yogurt and dairy product, it was also shown to be pathogenic in some other circumstances. From this study, the isolated strains could potentially be used as probiotics in poultry feeding formulation.

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