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"Industry based on Knowledges"

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### The Proceeding Of

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"Industry Based On Knowledges"

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# **ORAL PRESENTATION**

The 1st Conference Technology on Biosciences and Social Sciences 2016

## **ANIMAL SCIENCE**

### Association Analysis of NRAMP1 Gene Related to Resistance Against Salmonella pullorum Infection in Kampung Chicken

Jumatriatikah Hadrawi<sup>a</sup>\*, Asep Gunawan<sup>b</sup>, Niken Ulupi<sup>b</sup>, Sri Darwati<sup>b</sup> Cece Sumantri<sup>b</sup>

<sup>a</sup>Postgraduate School, Department of Animal Production And Technology, Faculty of Animal Science, Bogor Agricultural University <sup>b</sup>Department of Animal Production And Technology, Faculty of Animal Science, Bogor Agricultural University \*Corresponding author: atikahjumatri@gmail.com

### **Abstract**

Natural Protein Resistance-Associated Makrofag 1 (NRAMP-1) gene plays a role in controlling disease resistance. Kampung Chicken is one of the local Indonesian chickens which have high diversity in term of productivity. Breeding programs to improve disease resistance through molecular selection is one of the efforts that have evolved to increase theproductivity of chicken. The present research was to study the association of apolymorphism of NRAMP-1 with resistance disease in Kampung chicken. The PCR-RFLP method was applied to analyze the association between the polymorphism of NRAMP1 with resistant disease against Salmonella Pullorum. NRAMP-1 gene was genotyped in Kampung chickens using the PCR-RFLP method. The result showed three genotypes were identified of NRAMP-1 gene in Kampung chicken, namely TT, TC, and CC. The NRAMP-1 gene was polymorphic in all native chickens. The chi-square value of the Kampung chicken showed deviate in Hardy-Weinberg equilibrium. The TT and TC genotype revealed higher (P<0.05) mortality bacterium compare to CC genotype. In conclusion, polymorphism in chicken NRAMP-1 gene could be used as acandidate gene to increase resistance to disease in Kampung chicken.

Keyword: Nramp-1 gene, Kampung Chicken, resistance trait, and Salmonella pullorum

### 1. Introduction

Indonesia is one of the genetic diversity producerassociation. center of local chickens in the world[1]. The population of local chicken from 2012 to 2015 according to Dirjen PKH [2]very small increase in the value just of 1%. Kampung chicken is a kind of Indonesian local chicken that does not have special characteristic and spread out in various regions of Indonesia. There is two big problems which become astumblingblockin developing kampung chicken. The first problem is the difficulty to institutions belong to the government with chickens act as carriers. ([3], [4]).

aresearchinstitution and with local chicken

Most of thenative chicken were raised extensively with marginal feed, with environmental hygiene low and implementation of biosecurity. even though kampung chicken is able to thrive despite an increase in low population. Kampung chicken is resistant to several deadly diseases such as a pullorum. The disease is caused by the bacterium Salmonella pullorum, Salmonella pullorum is the cause of pullorum disease get day old chick of local chicken. This attacking young ages under a month with problem can be solved by integrating breeder amortality rate of 20% and 80% and adult

Resistance immune has theenvironment and feeds also controlled by genes. One of the genes controlledis a Natural Resistance-Associated Protein-1 (NRAMP-1) 2.3. Identification of the polymorphism genes. In poultry, a homologue of NRAMP1 gene has been mapped on chromosome 7 which consists of a promoter region, 15 exons, 14introns, and flanking regions 5760bp in length [5], NRAMP1 gene restricts microbial access to essential micronutrients, such as Fe<sup>2+</sup>,  $Co^{2+}$  $Mn^{2+}$ , andZn<sup>2+</sup>,within professional phagosomes. NRAMP1 gene belongs to a large gene family encoding divalent cation localized transporters that are to late endosomes/ lysosomes and are proposed to affect intraphagosomal microbial replication by modulating divalent cation content in this organelle. The many cellular functions that depend on metal ions as cofactors explain the pleiotropic effects of NRAMP1 and its complex role in infectious diseases ([6],[7], ([8],([9],).

The results study NRAMP-1 gene is polymorphic in broiler chickens [10], the native chicken Malaysia [11], and native China ([12],[13]).Until chickens now, variations in NRAMP1gene and their effect on disease have not been well investigated in Indonesian native chickens. Therefore, the objective of the present study was to identify the association of NRAMP1gene polymorphisms with immune traits

#### 2. Material and Methods

### 2.1.The time and place of study

The study was conducted in Juli 2015 until November 2017at the Laboratory of Animal Breeding and Genetics IPB and Laboratory of Medical Microbiology Faculty of Veterinary Science IPB.

### 2.2.Blood Samples

Blood samples were 44 population kampung chicken. DNA was extracted from blood samples at the Laboratory of Animal Breeding and Genetics, Faculty of Animal

affected Science, **Bogor** Agricultural University (Indonesia).

# NRAMP-1gene

The first Blood samples were 44 population kampung chicken. A blood sample was taken from the brachial vein in the wing Identification of the polymorphism NRAMP-1gene against kampung chicken consists of 3 phases: DNA extraction, PCR amplification and **RFLP** (Restriction Fragment LengthPolymorphism). Genomic phenol-chloroform DNA extraction used method[14]and the DNA was dissolved in theelution buffer. The quality of the total genomic extraction was assessed by 1% agarose gel electrophoresis. Polymerase chain reaction (PCR) was carried out using primers specific for a part of exon 11 (421 bp) of NRAMP1 gene (GenBank Accession No. 5'-AY072001): forward caatgagacggtgtctgtgg-3'; reverse 5'cccagaagaaatctccctgc-3'. Amplification was carried out with a GeneAmp® PCR 9700 System (Applied Biosystems, USA). Thermal consisted cvcling conditions predenaturation at 95°C for 5 min, followed by 35 cycles of denaturation at 95°C for 10 s. annealing at 60°C for 20 s, and extension at 72°C for 30 s; the final extension step was at 72°C for 5 min. RFLP method was used to determine the genotype NRAMP-1 gene. PCR result of the NRAMP-1 genefragments was bySacIrestriction enzymes.DNA amplification products and a standard DNA ladder were separated on 1.5% agarose gels in 0.5X TBE buffer.

### 2.4. Clearance Test

Immune traitswere detected in blood samples using the clearance test [15]. This method was used to look at normal bacterial (S.pullorum) population growth that of populations given specific treatment. The treatment impact on bacterial growth was

measured after incubating for 24-48 hat  $35\pm1^{\circ}\text{C}$ . Preparation of bacteria culture begins with the rejuvenation of culture in nutrient medium at a temperature of  $36\pm1^{\circ}\text{C}$  for 18-24 h and a sub-culture on Brain Heart Broth medium at a temperature of  $36\pm1^{\circ}\text{C}$  for 18-24 h.

### 2.5. Data analysis

Data were analyzed with ANOVA using completely randomized design. NRAMP-1 genotype was as treatment biological assays data were as aresponse. A  $\mu$  + Pi + statistical model was used Yij =  $\varepsilon ij$ , where  $Y_{ij}$  is the observation on immune traits,  $\mu$  is the overall mean,  $P_i$  is the effect of polymorphism the single nucleotide genotypes, andeij is the random residual effect[16].

### 3. Result and Discussion

NRAMP-1 gene in chicken located on chromosome 7. The data obtained from GenBank (GenBank accession number: AY072007). The size of NRAMP1 gene was 5760pb. The structure of this gene was begun by a promoter region, exons (15), introns (14) and theend was the flanking region.

NRAMP-1 gene *Genotyping* on exon 11, and with 421 bp PCR product. NRAMP1 gene cutting by *Saclenzyme* restriction. The result showed two alleles (C and T). The genotype were identified of NRAMP-1 gene in Kampung chicken, namely TT, TC and CC where restriction fragments included asingle, uncut fragment of 421 bp (TT genotype), two fragments of 258and163 bp(CC genotype), and three fragments of 421, 258, and 163 bp (TC genotype) (Figure.1).

# Genotypes and allele frequency of NRAMP1 gene inkampung chickens

Allele and genotype frequency values of NRAMP1 genes against kampung chicken presented in Table 1.

The results showed of genotyping frequency of CC genotype dominated, could be interpreted that the gene NRAMP loci SACI-1 polymorphic trait. So for [17] which states that genetic polymorphic in an individual can be seen when there are two alleles for the like gene but different DNA configuration which occupied the same locus on a chromo some. According to Nei and Kumar [18], polimorpysm can be indicated by the presence of two or more alleles in a population and the allele frequency is equal to or below 0.99.

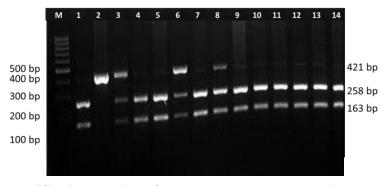


Fig. 1. PCR-RFLP amplification product of NRAMP1 gene at exon 11 that was cut by the SacI.

Table 1. Genotypes and allele frequencies of polymerase values of NRAMP1 genes against kampung chicken

Chicken	N	Allele frequency		genotype frequency		су
Chicken	IN	T	C	TT	TC	CC
Kampung	44	0.24	0.76	0.14 (6)	0.21 (9)	0.66(29)

# Heterozygosity and Hardy-Weinberg equilibrium Genotype Gen NRAMP1

The results of heterozygosity values and NI Hardy-Weinberg equilibrium of NRAMP ka gene in locus SacI presented of Table 2. Genotype and allele frequencies The showed high that the deviate in Hardy-Weinberg to equilibrium.

Hardy-Weinberg equilibrium showed that the value of chi-squared higher than aChi-squared table at the 1% level of confidence, that the results were significantly different. In other, the ratio deviates from higher expectations. The the degree of heterozygosity in a population, the survival of the population will be higher. The degree of heterozygosity is the average percentage of loci heterozygosity of each individual or the percentage of heterozygous average individuals in the population [18].

### Resistance of Kampung Chickens

The resistance ofkampung chickens in this study demonstrated the ability of chicken in ingesting and killing of bacteria. Ingesting

killing ability has shown by clearance test with infection *S. pullorum*. The association of *NRAMP1* genegenotype with immune traits in kampung chickensare presented in Table 3.

The TT and TC genotype revealed higher (P<0.05) mortality bacterium compare to CC genotype. The data showed polymorphism NRAMP-1 gene in kampung chicken associated with immune traits of infection *S. Pullorum*. As a candidate disease resistance gene, NRAMP1 gene has been studied by a number of researchers throughout the world.

This study conducted by Liu et al[10], demonstrated theassociation of an SNP polymorphism in a highly conserved region of NRAMP1 with Salmonella enteritidis vaccine and pathogen challenge response in young chicks, indicating that either NRAMP1 or a gene controls these S.enteritidis response traits. Hu et al [19] report about theassociation of apolymorphism of immune functions in NRAMP1 with same chicken takes effect.

Table2. heterozygosity observations Value, expectations and Hardy-Weinberg equilibrium NRAMP 1 gene

Chicken	N	Но	Не	$x^2$
Kampung	44	0.36	0.21	8.389 <sup>n</sup>

n: significantly different,  $x^2$  (0.05,1) = 3.84

Ho: heterozygosity observation He: heterozygosity expectations  $x^2$ : Hardy-Weinberg equilibrium

Table 3. Association of NRAMP1 gene genotypein kampung chickens resistant to Salmonella pullorum

Genotype	Early concentration	Final concentration	Death rate of
	(CFU/ml)	(CFU/ml)	bacteria (%)
TT	$6.8 \times 10^{10}$	$4.17 \times 10^8$	99,39 <sup>ab</sup>
TC	$6.8 \times 10^{10}$	$1,73 \times 10^8$	99,74 <sup>a</sup>
CC	$6.8 \times 10^{10}$	$7,07 \times 10^8$	98,96 <sup>b</sup>

#### Conclusion

The TT and TC genotype revealed higher (P<0.05) mortality bacterium compare to CC genotype. In chicken, NRAMP-1 gene could be used as acandidate gene to increase resistance to disease in Kampung chicken.

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