# ANTIMICROBIAL RESISTANCE DETECTION OF CITROBACTERFREUNDII ISOLATED FROM RECTAL SWAB IN EAST JAVA

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Submission date: 31-Dec-2019 10:54AM (UTC+0800)

**Submission ID:** 1238821370

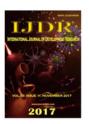
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## ANTIMICROBIAL RESISTANCE DETECTION OF CITROBACTERFREUNDII ISOLATED FROM RECTAL SWAB IN EAST JAVA

Vol. 07, Issue, 11, pp.17119-17122, November, 2017

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### ARTICLE INFO

### Article History:

Received 05th August 2017 Received in revised form 29th September, 2017 Accepted 18th October, 2017 Published online 30th November, 2017

### Key Words:

Antimicrobial resistance, Citrobacterfreundii, Rectal swab, bla<sub>TEM</sub>gene, ESBL.

### ABSTRACT

The aim of this study was to detect  $bla_{\text{TEM}}$ gene which the agent of antibiotic resistance in Citrobacterfreundii isolated from rectal swab in East Java. A total of 275 rectal swab samples were collected from 12 district of East Java. Then, samples were isolated with selective media and biochemical tests. Antibiotic Resistance test was using the disc diffusion against amoxycillin, amoxycillin-clavulanic acid, ampicillin-sulbactam, ceftoaxime, ceftazidime, sulfametaxole-trimethoprim, and tetracycline and then detection of bla  $_{\text{TEM}}$  gene used PCR. The number of Citrobacterfreundii was 7 (2,55 %). Antimicrobial resistance showed 2 broilers and 1 dairy cattle from Surabaya resistant to ampicillin-sulbactam and tetracycline, 4 milkfish from Sidoarjo resistant to amoxicillin and ampicillin-sulbactam. All seven samples were positive  $bla_{\text{TEM}}$  gene with 445 bp size used PCR.

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Citation: Mustofa Helmi Effendi, Intan Permatasari Hermawan, Wiwiek Tyasningsih, Suwarno, Intan Galuh Bintari, Risi Cicilia and Rinda Dewi Safitri. 2017. "Antimicrobial resistance detection of citrobacterfreundii isolated from rectal swab in east java", International Journal of Development Research, 7, (11), 17119-17122.

### INTRODUCTION

Citrobacter freundii is a group of Enterobacteriaceae bacteria that can be isolated from stool specimens or intestinal tract of humans and animals (Kao et al., 2010). Farm products became the biggest source of fecal contamination that can cause human pathogens. Fecal contamination of water or feces usually present the environment or migrate through the water. Still little is known about the effects of antibiotic resistance in animal faeces. Antibiotic resistance in the particles containing the gene resistant genetically and can spread through animal feces in the soil and aquatic ecosystems (Adelowo and Fagade, 2010). Excessive use of antibiotics as well as non-compliance with medication causing downtime

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residues in poultry meat, eggs and milk or other dairy products (Maynard et al., 2003). National Institutes of Health reported that the death rate in the United States reached 90,000 people per year due to antibiotic resistance. Antimicrobial Resistance Bank CDC in 2013 reported That Citrobacterfreundii resistant to the antibiotic Amoxicillin, Ampicillin, Cefazolime and Cefoxitin (Guilfoile et al., 2007). Antibiotic resistance to betalactam antibiotics caused by the production of beta-lactamase, an enzyme that can hydrolyze beta-lactam ring so that the bacterial cell wall synthesis process is not disrupted. One of the enzymes produced by bacteria is the extended-spectrum beta-lactamase, or ESBL which is the result of mutations of beta-lactamase. Most ESBL derived from TEM-type enzyme encoded by the gene blatem. Blatem genes are genes that cause 11 tibiotic resistance in the plasmid of the most frequently detected in clinical populations of gram-negative microorganisms (Wilopo et al., 2015). Based on this background it is necessary to detect the gene encoding blaTEM as antimicrobial resistance in bacteria Citrobacterfreundii isolated from rectal swab of animal that endanger public health.

### **MATERIALS AND METHODS**

### Samples

A total of 275 rectal swab samples were collected from 12 district. Rectal swab samples were taken aseptically. The sampling technique is purposive sampling. Rectal swab taken with a sterile cotton swab that has been prepared and then inserted into the sample pot containing physiological saline and has been labelled the previous location. Rectal swabs taken directly from the rectum or spontaneous defecation (rectal toucher). Taken stool samples must be free of urine to prevent bacterial contamination of urine. All samples were stored in the cool box or directly checked exceed 30-40 minutes.

**Table 1.Samples** 

No	District	Sample	Number of sample	
	Surabaya	Catfish, Tilapia fish, Parrot fish, Shark	33	
1		catfish, Shrimp, Dairy cattle, Broiler		
2	Sidoarjo	Tilapia fish, Milkfish, Catfish, Shark	31	
		catfish, Beef cattle		
3	Gresik	Tilapia fish, 13 mp, Milkfish	8	
4	Bojonegoro	Tilapia fish, Catfish, Gurame fish	12	
5	Blitar	Tilapia fish, Catfish, Gurame fish	2	
6	Probolinggo	Tilapia fish, Milkfish, Shrimp, Grouper	11	
		fish, Mullet fish		
7	Mojokerto	Tilapia fish, Catfish, Gurame fish, Shark	22	
		catfish		
8	Malang	111 pia fish, Milkfish, Shrimp	8	
9	Nongkojajar	Dairy cattle	39	
10	Grati	Dairy cattle	48	
11	Senduro	Dairy cattle	30	
12	Batu	Dairy cattle	31	
	Total sample		275	

Table 2. Results of resistance phenotype

No	Code	Sample	Location	Resistance phenotype
1	AW-1	Broiler	Wonokromo, Surabaya	SAM, TET
2	AK-5	Broiler	Keputran, Surabaya	SAM, TET
3	SB-3	Dairy cattle	BendulMerisi, Surabaya	SAM, TET
4	SPT.K-1	Beef cattle	Krian, Sidoarjo	AML, SAM
5	IB.S-1	Milkfish	Sidoarjo	AML, SAM
6	IB.S-2	Milkfish	Sidoarjo	AML, SAM
7	IB.S-4	Milkfish	Sidoarjo	AML, SAM

AML= amoxicillin, SAM= ampicillin-sulbactam, TET= Tetracycline

### Isolation of Citrobacterfreundii

All samples were diluted and the 10 olated on Eosin Methyline Blue Agar and Mac Conkey's Agar. The inoculated plates were then incubated at 37°C for 24 hours and then observed macroscopically. Characteristics of bacterial colonies Citrobacterfreundii on EMBA appeared diameter of 2-4 mm, smooth, low, convex, moist, metallic green, on MCA appear pink colonies (Holt et al., 1994). Furthemore, all bacterial isolated and identified by magnification microscopic examination with negative Gram staining. Negative gram stain showed short stems (bacilli) and then all bacterial isolated and identified by biochemical test (Lay, 1994). Identification test using Triple Sugar Iron Agar, glucose, lactose, sucrose,

mannitol, maltose, Simon's Indol Motility, MR- Vogen Prekauer (VP), Simon's Citrate Agar and Urea Agar.

### Antibiotic Susceptibility Testing

The antimicrobial sensitivity phenotypes of bacteria lisolates were determined using a Kirby-Bauer disk diffusion assay according to the standards and interpretive criteria described by the Clinical and Laboratory Standards Institute. And used stand art Mac Farland number 0.5 so that the concentration of bacteria to be equivalent to 3.0 x 108 CFU / ml (Mcfarland, 1907). Inspection began with a cotton stick sterile dipped into the bacterial suspension then pressed on the walls of the tube until the cotton is not too wet, then applied to the surface of the Muller Hinton Agar until blended, wait for about 15 minutes, then put the disc containing antibiotics on it and incubated at 37°C for 19-24 hours. Antibiotic resistance tests against Amoxycillin 20 mg, Amoxicillin-clavulanate 20/10 mg, ampicillin-sulbactam 10μg / 10μg, Cefotaxime 30 g, Ceftazidime 30 g, Sulfametaxole-Trimethoprim 23.75 1,25ug, Tetracycline 30 ug. Observations of each antibiotic susceptibility test results based on the diameter of that zone was measured using calipers (Ahmed and Shimamoto, 2011).

### Detection of blaTEM Using PCR

### **DNA** extraction

DNA bacteria extraction were grown on agar overnight at a mperature of 36°C and then taken colonies using 1µl sterile plastic loop and transferred into 184 mL of sterile distilled water and added to 5 µl lysozyme (5 mg/ml) followed by 1 cubation at 56°C for 30 minutes.

Furthermore, the DNA was diluted to 100µl with buffer kit (QIAamp DNA mini kit 50). A total of 1µl DNA solution used for PCR amplification (Moenstein et al., 2007).

### Amplification

Primers used for detection of gene bla TEM shown in Table 3. In the process annel (attachment) used 1µl DNA solution Hot Star Taq Qiagen Master Mix (Qiagen Nr. 203 445) and 10 pmol primer specific gress reach a final volume of 25µl. Amplification followed denaturation process at 95°C for 15 min, 30 cycles of denaturation at 94°C for 30 seconds, anneling at 50°C for 30 seconds, extension at 72°C for 20 minutes (Moenstein et a 6 ., 2007). The next step was reactingthe PCR products by electrophoresis in 1% (w/v) agarose gel, then stained with ethidium-bromide and visualized using UV light. primersand DNA sequence was shownin Table 2. (Ahmedand Shimamoto, 2011). In genedetectionbla TEM positive results when there are 445 base paresized DNA band (Moenstein etal., 2009).

### RESULTS

### Isolation of Citrobacterfreundii

The results of this study from the isolation and identification of bacteria from a total of 275 samples found positive *Citrobacterfreundii* were seven samples (2,55%) that the samples of rectal swab of Boriler from Wonokromo Surabaya (AW-1), Broiler from Keputran Surabaya (AK-5), Dairy Cattle from Bendul Merisi Surabaya (SB-3), Beef cattle from KrianSidoarjo (SPT.K-1), Milkfishes from Sidoarjo (IBS-1), (IBS-2), (IBS-4).

Table 3. Primer used for PCR (Moenstein et al., 2007)

Primer	Sekunes (5 to 3)	Gene
TEM-F	TCGCCGCATACACTATTCTCAGAATGA	Bla <sub>TEM</sub>
TEM-R	ACGCTCACCGGCTCCAGATTTAT	Bla <sub>TEM</sub>

### Antibiotic Susceptibility Testing

Interpretation of the results obtained diameter of antibiotic zone that Broiler from Keputran (AK-5) and Wonokromo (AW-1) and Dairy cattle (SB-3) are resistant to ampicillin-sulbactam and tetracycline, but Beef cattle from Krian (SK-1), Milkfishes from Sidoarjo (IBS-1), (IBS-2), (IBS-4) are resistant to Amoxycillin and Ampicillin-Sulbactam. But on the other antibiotics are still visible intermediete and sensitive.

### Detection of blatem Using PCR

PCR result identified of bla<sub>TEM</sub> in *Citrobacterfreundiis*howed on seven samples of broiler from Keputran (AK-5), broilers from Wonokromo (AW-1), Dairy cattle from BendulMerisi (SB-3), Beef cattle from Krian (SK-1), Milkfishes from Sidoarjo (IBS-1), (IBS-2), (IBS-4). The results showed the band in 445 basepare (bp).

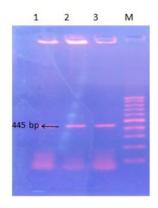


Figure 1. PCR assay to detect bla TEM Lane 1= AK-5; Lane 2= AW-1; Lane 3= SB-3; M= Marker

### DISCUSSION

The increasing prevalence of ESBL-producing Enterobacteriaceae very harmful to people's health, because the transmission can be derived from food products of animal origin or can be derived from animal faeces. In this study found 7 (2,55%) samples positive *Citrobacterfreundii*from the total 275 samples of rectal swab. Seven samples were broiler from Keputran (AK-5), broiler from Wonokromo (AW-1), Dairy cattle from BendulMerisi (SB-3), Beef cattle from Krian (SPT.K-1), Milkfishes from Sidoarjo (IBS-1), (IBS-2), (IBS-4).

Three samples were AK-5, AW-1, SB-3 found to be resistant to the antibiotic ampicillin-sulbactam, and tetracycline. And SPT.K-1, IBS-1, IBS-2, IBS-4 found to be resistant to the amoxicillin and ampicillin-sulbactam (Table 4). Then followed the PCR test to detect gene bla<sub>TEM</sub> as agents ESBL and found positive for the band DNA of seven samples in 445 basepare (Figure 1).

In research done by Ahmed and Shimamoto (2011) That the gene coding for antimicrobial resistance blaTEM as found as many as 23 isolates (20.5%) of 34 isolates of gram-negative bacteria in the case of mastitis in Egypt and in Citrobacterfreundii found blaTEMgenes as much as 3 isolates. Results of research conducted by Kao et al. (2010) found a gene in Citrobacterfreundii showed blaTEM much as 2 isolates of 30 isolates sampled. Extended Spectrum Beta Lactamase (ESBL) derived from TEM-type enzyme encoded by the gene blatem. The Blatem genes are genes that cause atibiotic resistance in the plasmid of the most frequently detected in clinical populations 9 gram-negative microorganisms (Wilopo et al., 2015). Extended Spectrum Beta Lactamase (ESBL) have the ability to transfer genes from one organism into another organism, so that the spread of resistance very easily occur between strains and even between species (Wilopo et al., 2015). Plasmids are also responsible for coding genes that carry resistance genes to other drug classes (eg, classes (aminoglycoside). This makes selection of antibiotics against ESBL-producing organisms are very limited (Jacob and Munoz, 2005). Members of the family Enterobacteriaceae including Citrobacterfreundii expressing plasmid-encoded βlactamase. Generally ESBL genes derived from TEM-1, TEM-2 or SHV-1 experienced mutase and change the configuration of amino acids around the active site of beta-lactamase. This situation makes the spectrum beta-lactam antibiotics susceptible to hydrolysis by this enzyme (Bush and Jacoby, 2010).

### Conclusion

The conclusion of this study shows that seven samples positive *Citrobacterfreundii* found in broiler chickens, dairy cattle, beef cattle and milkfishes in East Java resistance to antibiotics. And *Citrobacterfreundii* showed as ESBL agent to detect bla<sub>TEM</sub> gene using PCR with 445 basepare size.

### Acknowledgement

Supported by Ministry of Research, Technology and Higher Education Indonesia on research of Penelitian Terapan Unggulan Terpadu fiscal year 2017.

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