

Fungal Contamination and Toxigenic *Aspergillus flavus* Strains on Dried Anchovies Sold in North Sumatra

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Abstract: Dried anchovies at Medan traditional markets commonly sold unpacked and stored in open air, as the result, the dried-salted fish was easily spoiled and contaminated by mycotoxigenic fungi. The aims of the study were to enumerate fungal population and to characterize *A. flavus* strains. Each fungal species were isolated from medan anchovies at five traditional markets. The percentage of fungal contamination was determined by a direct plating using dichloran 18% glycerol agar medium. Fungal population was determined using a serial dilutions and pour plate in DG18 medium. Toxigenic *Aspergillus flavus* strains were characterized by the production of yellow pigment on the reverse side of petri dishes containing coconut agar medium 10%. The presence of regulatory (*aflR*) and structural genes (*nor-1*, *ver-1* and *omt-1*) genes determined aflatoxin biosynthesis was determined by amplifying four sets of specific primers (*aflR*, *nor-1*, *ver-1* and *omt-1*). Results showed that medan anchovies were contaminated by storage fungi. *Aspergillus niger* was the most common found followed by *A. flavus* and *A. chevalieri*. Eighty strains of *A. flavus* were isolated from all samples, they are consisted 69 toxigenic and 11 strains were non-toxigenic.

1 INTRODUCTION

Anchovies is dried-salted fish that one of fish products that contain high protein content. Medan anchovies commonly produced traditionally by local fisherman. During processing the fish product was preserved conventionally by adding some salt and dried under the sunlight. Moisture content anchovies between 32 and 34% with salt content between 13 and 17% (Yusra 2017). During storage and high relative humidity the dried-stored anchovies might increase their water activity (a_w) and contaminated by storage fungi.

About 14 of 150 dried-salted fish samples stored at a_w 0.73 to 0.80 were contaminated by *Aspergillus flavus* (Indriati et al. 2017). In addition, *Aspergillus flavus* was the most predominant found on dried salted fish (Youssef et al. 2003) Storage fungi such as *Aspergillus flavus*, *A. candidus*, *A. niger*, *A. wentii*, *A. chevalieri* and *Penicillium islandicum* commonly xerophilic and was able to grow at low a_w (Pitt and Hocking 2009).

Fungal contamination occurred on dried fish at moisture content 12.3 to 32% and genus *Aspergillus* particularly *A. niger*, *A. fumigatus*, *A. ochraceus* and *A. flavus* were the most common found (Sam et

al. (2015). Fungal contamination on food and feedstuffs may cause change taste, colour, reduce nutrition (Hassan et al. 2011). The purpose of recent study was to enumerate fungal contamination and to determine toxigenic *A. flavus* strains on anchovies sold by retailers at traditional markets in North Sumatera, Indonesia.

2 MATERIALS AND METHOD

2.1 Anchovies Sampling

Anchovies as sample were bought from five traditional markets and as much as 100 g anchovies obtained from five retailers at each market. All samples were packed in sterile polyethylene bag and stored in cold until used.

2.2 Determination of the Percentage of Anchovies Contaminated by Fungi

The percentage of anchovies contaminated by each fungal species was determined by direct plating on dichloran 18% glycerol agar (DG18) medium in

petri dish (9 cm diameter). Five replicates were made for each sample.

2.3 Enumeration of Fungal Population

Each fungal population was determined by serial dilution method followed by pour plate in Dichloran 18% Glycerol Agar ((DG18). One hundred gram of each sample were ground using Mill Powder RT-04 no serial number 980923 (Mill Powder Tech. Co LTD, Taiwan) with speed 25 000 rpm for 30 second and 25 g of fine sample in 500 ml flask was dilute by sterilized distilled water until the volume up to 250 ml. The suspension then was homogenized and 1 ml of the aliquot in petri dish (9 cm in diameter) was pour plate in DG18 medium, three replicates was made. All plates then were incubated at 28°C for 5 d. Each fungal population were enumerated, isolated in Potato Dextrosa Agar (PDA) and identified according to Pitt and Hocking (2009) by subculturing in Malt Extract Agar (MEA) and Czapek Yeast Agar (CYA).

2.4 Determination on the Toxicogenicity of *Aspergillus flavus* by Culture Technique

The toxicogenicity of each *A. flavus* strains to produce aflatoxin was determine using coconut agar medium (CAM) according to Lin and Dianese (1976). Each strain of *A. flavus* was culture in CAM for 5 day at 28°C. The presence of yellow pigment at the reverse side of petri dish indicate toxicogenicity.

2.5 Extraction of *A. flavus* Genome

Genome of each *Aspergillus flavus* strain was obtained from two 2 days old mycelial culture on PDA (28°C). As much as 40 g mycelia in 1.5 ml PCR tube containing 600 µl nuclei lysis solution was homogenated using micropestle. Genome extraction kit used was Mini Kit (Promega, Madison, WI, USA). The homogenate then was incubated at 65°C for 15 minute and cooled 5 minute until the temperature 29°C, then, 200 µl protein precipitation was added, vortexed and centrifuged at 13000×g for 3 minute. The supernatant was mixed in 600 µl isopropanol followed by centrifugation 13000×g for 2 minute, the supernatant was discarded and 600 µl ethanol 70% was used to wash the pellet. The mixture was then centrifuged 13000×g for 2 minute. The pellet obtained then rehydrated by 50 µl DNA rehydration solution, vortexed for 5 second and 0.5 µl RNase was add. The DNA mixture was

incubated at 37°C for 15 minute. The DNA concentration was determined using nanophotometer (IMPLEN, Munich, Germany). Electrophoresis DNA was conducted using SCIE-PLAS, Cambridge, England with 1,2% gel agarose and gel was stained with 1 µl ethidium bromide. The photograph used was Gel Doc (Uvitec, Cambridge, Serial) under UV light 303 nm.

2.6 PCR Amplification of Aflatoxin-related Genes

Polymeration chain reaction used was GeneAmp PCR Labcycler Gradient System (Sensoquest, Germany, serial number 1123280105). Specific primer for genes aflatoxin biosynthesis used were *nor-1*, *ver-1*, *omt-1* and *aflR* with basepair 400, 895, 1232 dan 1032 bp respectively. Amplification preparation was conducted as follows: Amplification mix for each primer consisted of 5 µl GoTaq Green Master Mix (Promega), 3 µl nuclease free water, 0,5 µl for each primer F (forward) and R (reverse), then 1 µl DNA template and volume of end DNA Mix reaction on PCR tube was 10 µl. Process of DNA mix in PCR was preincubation step at 94°C for 10 min followed by 35 cycles of denaturation at 94°C for 10 min, annealing at 65°C for 2 min and extension at 72°C for 2 min and final extension at 72°C for 7 min. The PCR product was analyzed by electrophoresis (SCIE-PLAS, Ltd, Cambridge, England) using 1,5 % agarose gel in 1×TAE 40 mM Tris-acetate, 1mM EDTA (pH 8). Gels stained with ethidium bromide 0.1 mg/l, visualized under *Gel Doc* (Uvitec, Cambridge, serial number 13 200263) under UV light (303 nm).

3 RESULTS AND DISCUSSION

3.1 Percentage of Contaminated Anchovies

Eleven species of storage fungi that contaminated anchovies were isolated from 5 traditional market (Table 1).

Table 1: Percentage of anchovies contaminated by fungi at five traditional markets in Medan, North Sumatera

Fungal species	Percentage of anchovies contaminated by fungi				
	Market 1	Market 2	Market 3	Market 4	Market 5
<i>Aspergillus candidus</i>	6.70	0	0	0	0
<i>A. chevalieri</i>	33.30	46.70	0	0	0
<i>A. flavus</i>	33.30	0	63.30	10.00	3.00
<i>A. fumigatus</i>	0	0	0	0	6.60
<i>A. niger</i>	30.0	23.3	23.30	100,00	43.30
<i>A. oryzae</i>	0	0	0	0	10.00
<i>A. tamarii</i>	3.30	46.70	0	10,00	3.00
<i>A. wentii</i>	3.30	0	0	0	0
<i>Penicillium</i> sp. ₁	13.30	0	0	0	10.00
<i>Rhizopus oligosporus</i>	33.30	13.30	3.30	0	10.00
<i>Mycelia sterilia</i>	3.30	0	0	0	0

Table 2: Fungal population (cfu/g) on anchovies contaminated by fungi at five traditional markets on Medan, North Sumatera

Fungal species	Fungal population (cfu/g)				
	Market 1	Market 2	Market 3	Market 4	Market 5
<i>Aspergillus candidus</i>	1×10	0	0	0	0
<i>A. chevalieri</i>	8.3×10	1.33×10 ⁻²	0	0	0.3×10
<i>A. flavus</i>	0.7×10	0	1.13×10 ⁻²	1.7×10	0.3×10
<i>A. fumigatus</i>	0	0	4×10	0.3×10	0.6×10
<i>A. niger</i>	0	0.3×10	3.3×10	3.3×10	0
<i>Cladosporium</i> sp.	0.3×10	0	0	0	0.3×10
<i>Rhizopus oligosporus</i>	0.3×10	0.3×10	0	0	0

CFU = colony forming unit

Table 3: Toxigenicity representative isolates of *A. flavus* strains isolated from anchovies based on cultural in CAM (coconut agar medium) medium and molecular based on the presence of regulatory (*aflR*) gene and structural (*nor-1*, *omt-1*, *ver-1*) genes

Strain	Regulatory gene		Structural genes			The presence of yellow pigment in CAM medium	Toxigenicity
	<i>aflR</i>	<i>nor-1</i>	<i>omt-1</i>	<i>ver-1</i>			
Af1	+	+	-	+	negative	non-toxigen	
Af2	+	+	+	+	positive	toxigen	
Af3	+	+	-	+	negative	non-toxigen	
Af4	+	+	+	+	positive	toxigen	
Af5	+	+	+	+	positive	toxigen	
Af6	+	+	+	+	positive	toxigen	
Af7	+	+	-	+	negative	non-toxigen	
Af8	+	+	+	+	positive	toxigen	
Af9	+	+	+	+	positive	toxigen	

Among the fungal species, the highest percentage anchovies were contaminated by *A. niger* followed by *A. flavus* and *A. chevalieri*, *A. tamarii*, *Penicillium* sp. and *Rhizopus oligosporus*. The presence of *A. niger* was high and found at all markets. Our result in line with Wheeler et al. (1986) who reported that dried salted fish was common infected by *A. niger*. Hassan (2011) also reported that most salted fish was contaminated by *A. niger*

(73.3%) higher than *A. flavus* (66.6%), *Penicillium* spp. (40%) and *Rhizopus* spp. (6.6%).

As shown in Table 1 all anchovies sold by retailers at traditional markets were infected by molds, the contamination might occur processing and fungal population increased during storage. As reported by Azam (2002) conventional drying on fish products susceptible contaminated by microorganisms. In addition, high relative humidity

leads the dried fish absorb water vapor from air and increase moisture content.

3.2 Fungal Population on Anchovies

Fungal population per gram anchovies (colony forming unit or cfu/g) obtained from relailers at five traditional markets are shown in Table 2). Genus of *Aspergillus* was the most dominant (5 species) followed by *Cladosporium* and *Rhizopus oligosporus*.

Table 2 showed that *Aspergillus chevalieri* was the highest population (8.3×10 cfu/g) followed by *A. niger* (3.3×10 cfu/g), *A. flavus* (1.13×10^{-2} cfu/g), and *A. fumigatus* (3.3×10 cfu/g). Among storage fungi *A. chevalieri* is xerophilic and grow at a_w 0.71 to 0.74 (Pitt and Hocking 2009). Previous study by Sam (2015) stated that 23 fungal species were isolated from dried fish and were predominated by *A. flavus* and *A. niger*.

3.3 Toxigenicity of *A. flavus* Strains

Eighty strains of *A. flavus* were successfully isolated. Based on toxigenicity determination using CAM medium, they are consisted of 69 strains are toxigenic (aflatoxin producers) and 11 strains are non-toxigen. Gene analysis determine aflatoxin biosynthesis (Table 3) indicate that regulatory gene (*aflR*) and structural genes (*nor-1*, *omt-1* and *ver-1*) are key enzyme that determine aflatoxin production as previously reported by Erami et al. (2007) that the presence of the for keys genes determine toxigenicity of *Aspergillus flavus*.

Table 3 showed that the presence of regulatory and structural genes on *A. flavus* determined aflatoxin biosynthesis. The toxigenicity determined byn culture technique method in CAM medium and molecular using specific primers showed similar results.

4 CONCLUSION

High fungal contamination on anchovies are potential to spoile and contaminated aflatoxin. Good handling practices was required to reduce fungal growth during storage.

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