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วิทยานิพนธ์นี้เป็นส่วนหนึ่งของการศึกษาตามหลักสูตรปริญญาวิทยาศาสตรมหาบัณฑิต สาขาวิชาเทคโนโลยีชีวภาพ มหาวิทยาลัยเทคโนโลยีสุรนารี ปีการศึกษา 2552

SELECTION OF MONOCLONAL ANTIBODIES SPECIFIC TO AFLATOXIN FROM PHAGE DISPLAY ANTIBODY LIBRARY

KUNTALEE RANGNOI

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SELECTION OF MONOCLONAL ANTIBODIES SPECIFIC TO AFLATOXIN FROM PHAGE DISPLAY ANTIBODY LIBRARY

Suranaree University of Technology has approved this thesis submitted in partial fulfillment of the requirements for a Master's Degree.

Thesis Examining Committee

(Assoc. Prof. Dr. Neung Teaumroong)

Chairperson

(Assoc. Prof. Dr. Montarop Yamabhai)

Member (Thesis Advisor)

(Dr. Amara Chinaphuti)

Member

(Prof. Dr. Sukit Limpijumnong)

Vice Rector for Academic Affairs

Dean of Institute of Agricultural Technology

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อะฟลาท๊อกซินเป็นสารพิษที่เกิดจากเชื้อรา ซึ่งพบปนเปื้อนในอาหารคนและสัตว์ รวมทั้ง ้ผลิตผลทางการเกษตรในหลายพื้นที่ วิธีการตรวจหาอะฟลาท๊อกซินที่ใช้กันอยู่ในปัจจุบัน ได้แก่ วิธี ้โครมาโตรกราฟี แบบแผ่นบาง และแบบของเหลวประสิทธิภาพสูง หรือวิธีทางอิมมูน ซึ่งเป็นวิธีการ ้ที่มีขั้นตอนที่สะควก รวคเร็ว และมีความกุ้มทุนมากที่สุด โดยวิธีการนี้ต้องใช้แอนติบอคีที่จำเพาะ ต่ออะฟลาท๊อกซินเป็นสารสำหรับตรวจสอบ เทคโนโลยีการแสดงโปรตีนบนผิวเฟจ นับเป็น ้ทางเลือกหนึ่งที่มีประสิทธิภาพสูงในการผลิตแอนติบอดี งานวิจัยนี้ได้ทำการคัดเลือกโมโนโคนอล แอนติบอดีที่จำเพาะต่ออะฟลาท๊อกซินบี 1 จากกลังเฟจซึ่งแสดงแอนติบอดีแบบเส้นเดี่ยว เอสซีเอฟวี บนผิวเฟจ โดยได้ทำการกัดเลือกจำนวน 3 รอบ แล้วนำเฟจจำนวน 96 โกลนที่กัดหามาได้ไปผลิต เป็นโมเลกุลแอนติบอดีเส้นเดี่ยวที่ไม่ติดอยู่บนผิวเฟจ จากแบกทีเรียสายพันธุ์ อีโคไล เอชบี 2151 ้จากนั้นนำแอนติบอคีเหล่านี้ไปทคสอบความสามารถในการจับกับอะฟลาท๊อกซิน โคยใช้วิธีการ ้อี่ไลซ่าแบบแข่งขัน จากนั้นจึงได้นำโคลนที่แสดงความสามารถในการจับอะฟลาท๊อกซินอย่าง ้ จำเพาะเจาะจงมาศึกษาลำคับเบสของคีเอนเอ โคยการวิเคราะห์ค้วยเครื่องวิเคราะห์ลำคับเบส อัตโนมัติ และนำแอนติบอดีที่ได้นำมาทำการแยกให้บริสุทธิ์ แล้วทำการศึกษาโครงสร้างโคยใช้วิธี เอสคีเอสเพจ และวิธีเวสเทิร์น บลอต รวมทั้งทำการวิเคราะห์คุณสมบัติในการจับจำเพาะต่ออะฟ ลาท๊อกซินโดยใช้วิธีการอีไลซ่า แบบแข่งขัน นอกจากนั้นแล้ว ได้นำแอนติบอดีที่คัดเลือกแล้วนี้ไป ้สร้างให้เป็นโมเลกุล เอสซีเอฟวีที่เชื่อมต่อกับ เอนไซม์อาลคาไล ฟอสฟาเตส (scFv-AP) เพื่อให้ ้สะควกในการตรวจจับค้วยวิธี อีไลซ่า แอนติบอคีเชื่อมต่อที่ไค้สร้างขึ้นนี้ มีความสามารถในการจับ กับอะฟลาท๊อกซินดีกว่าเมื่อเปรียบเทียบกับ โมเลกุลแอนติบอดีเส้นเดี่ยวประมาณ 3-4 เท่า โดย แอนติบอดีนี้สามารถใช้ในการตรวจจับอะฟลาท๊อกซินบี 1 ใด้ต่ำสุดประมาณ 0.006 ถึง 0.03 ้ไมโครกรัมต่อมิลลิลิตร (6-10 ส่วนในล้านส่วน) และพบว่าค่าความสามารถในการยับยั้งการจับกึ่ง หนึ่งอยู่ในช่วงระหว่าง 0.035-0.02 ไมโครกรัมต่อมิลลิลิตร เมื่อนำแอนติบอคีที่เชื่อมต่อกับเอนไซม์ อาลคาไล ฟอสฟาเตส นี้ไปทดสอบการจับกับอะฟลาท๊อกซินชนิดอื่นๆ ได้แก่ อะฟลาท๊อกซิน บี 2, จี 1, จี 2 และ เอ็ม1 พบว่าร้อยละความสามารถในจับกับจี 1 มีมากที่สุด ผลจากการศึกษานี้แสดงว่า ้สามารถใช้เทคโนโลยีการแสดงโปรตีนบนผิวเฟจในการคัดเลือกแอนติบอดีชนิดเส้นเดี่ยวที่มี ความจำเพาะต่ออะฟลาท๊อกซินบี 1 ได้ และยังสามารถทำการตัดต่อยืนเพื่อสร้างเป็นแอนติบอดีที่ เชื่อมต่อกับเอนไซม์อาลกาไล ฟอสฟาเตส ซึ่งสามารถนำไปใช้ประโยชน์ในการพัฒนาเป็นชุด

ตรวจสอบ สำหรับตรวจวิเคราะห์สารพิษอะฟลาท็อกซินที่ปนเปื้อนในผลิตผลทางการเกษตรได้ ต่อไป

สาขาวิชาเทคโนโลยีชีวภาพ ปีการศึกษา 2552

ลายมือชื่อนักศึกษา
ลายมือชื่ออาจารย์ที่ปรึกษา
ลายมือชื่ออาจารย์ที่ปรึกษาร่วม

KUNTALEE RANGNOI : SELECTION OF MONOCLONAL ANTIBODIES SPECIFIC TO AFLATOXIN FROM PHAGE DISPLAY ANTIBODY LIBRARY. THESIS ADVISOR : ASSOC. PROF. MONTAROP YAMABHAI, Ph.D., 83 PP.

PHAGE DISPLAY/SCFV/ ANTIBODY/ NAÏVE /BIO-PANNING/MONOCLONAL /RECOMBINANT /AFLATOXIN/ALKALINE PHOSPHATASE FUSION

Aflatoxins are one of the major mycotoxins that contaminate several agricultural products and human food in many areas of the world. Several methods for aflatoxin determination have been developed, including thin-layer chromatography (TLC), high performance liquid chromatography (HPLC), and immunological method. Among these, the most cost-effective method is immunological analysis that uses monoclonal antibodies as detection reagents. Phage display technology is a powerful alternative method for the production of monoclonal antibody. A specific antibody to aflatoxin B_1 (AFB₁), the most toxic and prevalence isoform, could be isolated by bio-panning from Phage-displayed human single-chain-variable-fragment (scFv) antibody libraries. After three rounds of affinity selection, ninety-six positive clones were induced for the production of soluble scFv antibody fragments by using Escherichia coli non-suppressor strain (HB2151), and tested for their ability to bind soluble AFB₁ by competitive enzyme-linked immunosorbent assay (ELISA). Amino acid sequence analysis of the selected clones was performed using information obtained from automated DNA sequencing. ScFv antibodies that could be inhibited by soluble AFB₁ were purified to analyze their structure by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) and western blot analysis, and confirm their specific binding to AFB₁ by competitive ELISA. In addition, selected scFv clones were engineered to create scFv- alkaline phosphatase (scFv-AP) fusions and used as

reagents for one-step detection in ELISA format. The scFv-AP fusions showed 3-4 fold improved binding property when compared to soluble scFv form. The IC₅₀ of scFv-AP by AFB₁ varied between 0.035-0.02 μ g/ml and the limit of detection was approximately 0.006-0.03 μ g/ml (6-30 ppb). The selected antibodies were specific to AFB₁ and could cross-react with AFG₁, but not to other aflatoxins, namely AFB₂, AFG₁, AFG₂, AFM₁. These results indicated that phage display technology could be used to obtain a specific antibody against aflatoxin, and that the scFv-AP fusion was an efficient detection reagent that could be further developed to generate a cost-effective diagnostic kit or biosensor for the detection of aflatoxin contamination in agricultural commodities and products in the future.

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LIST OF ABBREVIATIONS

°C	degree Celsius
μg	microgram
μl	microlitre
bp	base pair
DNA	deoxyribonucleic acid
et al.	Et alia (and other)
g	gram
h	hour
Kda	Kilo daltal
1	litre
mg	milligram
min	minute
ml	milliliter
mM	millimolar
ng	nanogram
pmol	picomol
rpm	revolution per minute
UV	ultraviolet
v/v	volume per volume
w/v	weight per volume

CHAPTER I

INTRODUCTION

1.1 Significant of this study

Aflatoxin is a potent mycotoxin produced by certain strains of Aspergillus flavus and A. parasiticus (Woychik et al., 1984). Among 18 different types of aflatoxins identified, major members are aflatoxin B_1 , B_2 , G_1 , and G_2 . Aflatoxin B_1 (AFB₁) is normally predominant in amount in cultures as well as in food products. These toxins especially AFB₁ are highly carcinogenic agents consistently found as contaminants in human food supplies and animal feeds in many areas of the world (Taylor et al., 1997). As a consequence, there is a need for routine screening of products destined for animal and human consumption. The methods of analysis include thin layer chromatography (TLC), gas liquid chromatography (LC), high-pressure and liquid chromatography (HPLC), which are laborious and require expensive instrumentation and clean-up of the samples (Devi et al., 1999). Immunological methods are cost-effective and adaptable to the situation of developing countries. Commercial test kits for detection of aflatoxin have been developed, which used polyclonal or monoclonal antibody as detection reagents. These techniques need the immunization of animals to obtain specific antibodies, which are costly and timeconsuming. Thus, an alternative method for large scale production of specific monoclonal antibody at lower cost in the form of a recombinant proteins obtain from phage display technology is attractive. Phage display antibody technology is also beneficial for the production of convenient test kits that can be distributed to a wide range of customers and user at lower cost, resulting in a better health of the populations in the future (Yuan et al., 1997). Also, the monoclonal antibodies have proven to be useful analytical tools for quantifying the aflatoxins.

Several *in vitro* methods have been developed for the production of antibodies an alternatives to *in vivo* animal-based methods (Bradbury et al., 2003). The most commonly used technology among the various *in vitro* strategies is phage display (Hust and Dubel, 2004) which is, recognized as a powerful tool for selecting recombinant antibody fragments with specific binding properties from a vast number of variants against a wide range of target molecules, such as proteins, glycoproteins, oligosacharides, nucleic acid toxins or low molecular weight compounds – haptens (Willats, 2002; Yau et al., 2003). Phage display, first described by Smith (1985), refers to the display of functional foreign peptides, proteins or antibody fragments on the surface of a bacteriophage (Clackson et al., 1991).

Filamentous bacteriophages have been used to display single-chain variable fragment (scFv) and Fab antibody fragments by fusion to the phage minor coat protein pIII (Hoogenboom et al., 1991; McCafferty et al., 1990). The antibodies phage display library can be used to select antibodies against any antigen, including foreign antigen, nonimmunogenic antigens and toxic antigens. Display of large and diverse antibody fragment gene repertoires on phages permits the direct selection of monoclonal antibodies using antigen-affinity chromatography without the need for immunization (Marks et al., 1991).

The procedure for selecting phage is simple and inexpensive. Using recombinant DNA technology and microbiological methods, peptides or proteins are fused to one of the phage capsid proteins by genetic engineering into appropriate vector (plasmid or phagemid). The peptides or proteins that displayed on the phage coat proteins are accessible to various molecular interactions. Affinity selection of antibodies is accomplished by exposing the phage library to immobilized antigen molecules (binding clones are captured and non-binding clones are washed away). The captured phage particles are eluted from antigen, amplified by infecting *Escherichia coli* host cells and used in a

subsequent round of affinity selection. After the final round of affinity selection phage particles are amplified in order to prepare and characterize their displayed antibodies individually (Petrenko and Vodyanoy, 2003). Finally, the monoclonal phage population with the desired binding specificities can be isolated (Willats, 2002).

Advantages of phage-display antibody technology include (i) selection of antibodies from a phage library is unbiased by the immunogenicity and less dependent on quantitative abundance of the target ligands (Schier et al., 1996), (ii) antibodies are produced in DNAencoded plasmids that are readily cloned and modified, (iii) antibodies can be produced in large quantities from *Escherichia coli* without the use of animals, (iv) the genes of the antibody can be further engineered to obtain improved antibody property such as sensitivity.

In this thesis, scFv monoclonal antibody which is specific to aflatoxin was produced by using phage display technique. The phage displayed antibody fragment was induced to produce soluble scFv fragment against aflatoxin. In addition, a fusion of scFv to bacterial alkaline phosphatase (AP) was generated. This construct was used as convenient one step detection reagents for Enzyme-linked Immunosorbent Assay (ELISA). This scFv-AP can be developed for the detection of aflatoxins in agricultural products and food products in the future.

1.2 Research objectives

- 1. To select phage displayed scFv antibodies which are specific to aflatoxin by biopanning.
- 2. To produce scFv fragments that can interact specifically with aflatoxin.
- 3. To create scFv-AP fusion for one-step detection of aflatoxin in ELISA format.

1.3 Scope and limitation of the study

In this study, the specific antibodies to aflatoxin B_1 were selected from human scFv antibody libraries by using *in vitro* panning process. Antibody was purified using Ni²⁺-NTA resin and was analysed by SDS-PAGE and western blot. Amino acid sequences of each scFv antibody were determined by DNA sequence analysis. Selected clones were engineered to create scFv-alkaline phosphatase fusions (scFv-AP) and used as convenient one-step detection probes for competitive ELISA. The sensitivity and cross-reactivity to unconjugated aflatoxin B_1 , B_2 , G_1 , G_2 and M_1 was determined by competitive ELISA.

CHAPTER II

LITERATURE REVIEWS

2.1 Aflatoxins

Aflatoxins molecules were first discovered in the early 1960 when 100,000 turkeys died in England after consuming contaminated brazillian peanut meal (Hartley et al., 1963). These toxins are highly carcinogenic secondary metabolites of *Aspergillus flavus* and *A. parasiticus* which produced during fungal infection of a susceptible crop in the field or after harvest. The productions of aflatoxins occur at high temperatures and humidity, therefore their contamination are most common in tropical and subtropical countries. Contamination can occur in the field, during harvest, transportation and storage under optimal conditions for fungal growth (Bedard and Massey, 2006). Conditions that contribute to fungal growth and the production of aflatoxins are hot and humid climate, kernel moisture, favorable substrate characteristics, and factors that decrease the host plant's immunity (insect damage, poor fertilization, and drought).

There are four principle types of aflatoxin: B_1 , B_2 , G_1 and G_2 plus two additional metabolic products, M_1 and M_2 which are named for their respective innate fluorescent properties and mobility in chromatogram (Pitt, 2000). Whereas the B designation of aflatoxins B_1 and B_2 resulted from the exhibition of blue fluorescence under UV-light, while the G designation refers to the yellow-green fluorescence of the relevant structures under UV-light. Aflatoxin M_1 and M_2 are major metabolites of aflatoxin B_1 and B_2 respectively. The aflatoxins M_1 and M_2 were first isolated from milk of lactating animals fed with aflatoxin. AFB₁ is the most toxic and most prevalent compound, followed by G_1 , B_2 and G_2 with decreasing toxicity (Busby and Wogan, 1984). Aflatoxins have been found in many foods and animal feeds. These toxins dissolve in various polar organic solvents including methanol, chloroform, dimethyl sulfoxide (DMSO), aqueous acetone and aqueous hexane-acetone-water azeotrope that developed for extraction procedures of natural products. They are heat stable, even at temperatures in excess of 100°C and are difficult to destroy. These toxins have closely similar structures and form a unique group of highly oxygenated, naturally occurring heterocyclic compounds. Their molecular formulas as established from elementary analyses and mass spectrometric determinations are B_1 : $C_{17} H_{12} O_6$, B_2 : $C_{17} H_{14} O_6$, G_1 : $C_{17} H_{12} O_7$ and G_2 : C_{17} $H_{14} O_7$. Aflatoxins B_2 and G_2 were established as the dihydroxy derivatives of B_1 and G_1 , respectively. Whereas, aflatoxin M_1 is 4-hydroxy aflatoxin B_1 and aflatoxin M_2 is 4dihydroxy aflatoxin B_2 . The chemical structures of these aflatoxins are shown in Figure 1. Some physical properties of the compounds are summarized in Table 1.



Figure 1. Chemical structures of major aflatoxins namely aflatoxin B₁, G₁ and M₁ with the double bonds in 8-9 positions and aflatoxins B₂, G₂ and M₂ without the double bond. Other methabolites are AFM₁ and AFM₂ have been found in milk and urine of lactating animals. AFM₁ and AFM₂ are produced from AFB₁ and AFB₂, respectively, by hydroxylation in lactating

 Aflatoxin	Molecular formula	Molecular weight	Melting point (°C)	
 B1	$C_{17}H_{12}O_6$	312	268-269	
B2	$C_{17}H_{14}O_6$	314	286-289	
G1	$C_{17}H_{12}O_7$	328	244-246	
G2	$C_{17}H_{14}O_6$	330	237-240	
M1	$C_{17}H_{12}O_7$	328	299	
M2	$C_{17}H_{14}O_6$	330	293	

Table 1. Chemical and physical properties of aflatoxins. Adapted from Wogen (1966).

2.1.1 Effects of aflatoxin contamination

Agricultural products are often contaminated with fungi that can produce toxic metabolites referred as mycotoxins. Among these, aflatoxins have assumed economic importance because of their influence on the health of human beings and livestock and on the marketability of agricultural products. The tolerance levels currently set by the regulatory bodies worldwide are typically 0.05 ppb for AFM₁ in milk, 10 ppb for AFB₁ and 20 ppb for total aflatoxins in food intended for human consumption and 20-300 ppb for total aflatoxins in animal feeds (Lee et al., 2004). Food and Drug Administration (FDA) action levels for aflatoxin present in human food is 20 ppb with the exception of milk is 0.5 ppb. The action level for most feed is 20 ppb.

Aflatoxins are both acutely and chronically toxic to animals, including man, causing acute liver damage, liver cirrhosis, induction of tumours and teratogenic effects (Stoloff, 1977). In humans, numerous studies have linked the incidence of primary hepatocellular carcinoma with the intake of aflatoxins, leading to the classification of AFB₁ as class 1 human carcinogen by the IARC (IARC, 1993). Toxic and especially carcinogenic effects of aflatoxins have been reported in several different animals, but susceptibility to these toxins

varies greatly with sex, age, species and strain within a species (Busby and Wogan, 1984; CAST, 2003). Numerous animal studies have shown that the liver is the main target organ and therefore the main symptoms of aflatoxin exposure in domestic and laboratory animals are hepatic injuries. Aflatoxin is hepatotoxic and its acute and chronic effects in swine are largely attributable to liver damage (Armbrecht, 1978). In cattle, the primary symptom is reduced weight gain as well as liver and kidney damage. Milk production is reduced in cow (Keyl, 1978). Lower aflatoxin doses may lead to milder hepatic injuries and reduced growth rate, especially in young animals (Pier, 1992). Domesticated animals are variably susceptible to aflatoxins; some oral LD_{50} values (lethal dose for 50% of animals) are given for aflatoxin B₁ in several animal species in Table 2.

Animal	sex	Age/size	LD ₅₀ (mg/kg)
Duckling	М	1 d	0.37
Rat	M-F	1 d	1.0
Rat	М	21 d	5.5
Rat	F	21 d	7.4
Dog	M-F	Adult	0.5
Hamster	М	30 d	10.2

Table 2. Oral LD₅₀'s of Aflatoxin B₁. Adapted from Robens and Richard (1992).

As for the mechanism of AFB_1 induced mutagenicity and carcinogenicity, AFB_1 epoxide adducts to DNA, preferably guanine nucleotides, causing point mutations mainly G-C to T-A (94%) or A-T (6%) (Bailey et al., 1996). Depending on the location this mutation, activation of proto-oncogenes or silencing of suppressor genes will cause initiation of the cancer process. Mutations in the *p53* tumor suppressor gene at the hot spot of codon 249, are discussed in the context of AFB₁, but relations between this mutation, AFB₁ and hepatitis B virus infection are not entirely clear (Smela et al., 2001).

It is clear that exposure to aflatoxins is hazardous to human and animal health. For that reason, most countries have regulations governing the allowable concentrations of aflatoxin in food and feed (Van Egmond, 1989).

2.1.2 Recent methods of analysis for aflatoxins

Aflatoxin contamination is widespread in staple crops like peanut, maize, sorghum, pearl millet, chillies, pistachio, cassava etc., and compromises the safety of food and feed supplies (Waliyar et al., 2009). It is important to be able to detect and quantify aflatoxins in commodities to protect human and animal health. Food and feed industries should monitor their products routinely to ensure that aflatoxins levels are below the regulatory limits. Sensitive, accurate, and precise methods of analysis are needed for any monitoring programme to be effective. Many analytical methods including thin-layer chromatography (TLC), high performance liquid chromatography (HPLC), gas-liquid chromatography (GLC), immunoassay and combined methods have been employed to detect AFB₁. TLC, GC and HPLC methods for determining aflatoxins in food are widely accepted as official methods. However, these methods are laborious and time consuming. Often these techniques require knowledge and experience chromatographic techniques to solve separation and interference problems. Immunoassays are preferred over analytical methods because of their simplicity and cost-effectiveness. However, commercial kits based on immunological methods are expensive, and may be difficult to import them. These methods have a unique ability to routinely handle a large number of samples and do not require time-consuming procedures and sophisticated equipment, many imunoassays for aflatoxin have been developed. The described immune assays have integrated either polyclonal or monoclonal antibodies. Antibody-based techniques, such as ELISA, are widely used for routine screening of possible mycotoxin contamination. Diagnostic assays usually are based on competitive ELISA formats in which the binding of the anti-hapten antibodies to a labeled tracer molecule is inhibited by free antigen from the sample. The decrease of labeled tracer bound to the antibody is then determined (Lauer et al., 2005).

In the present day, DOA-Afatoxins ELISA test kit was approved for determination of aflatoxins at the level of 0.4 ppb by using polyclonal for detection agents (according to the manufacture leaflet, Department of Agricultural, Thailand). However, these kits require immunization of animals to produce polyclonal antibody. The alternative method to produce antibodies at lower cost by using recombinant proteins obtains from phage display technology. There have been previous reports on the isolation single chain variable fragments (scFv) specific to aflatoxin B_1 from phage display antibody library (Daly et al., 2002; Dunne et al., 2005; Moghaddam et al., 2001; Yang et al., 2009).

2.2 Phage display technique

Phage display was first described by Goerge P. Smith in 1985 (Smith, 1985). This technique refers to the display of functional foreign peptides, proteins or antibody fragments on the surface of a filamentous bacteriophage. A gene of interest is fused to a gene encoding a phage coat protein, resulting in phage particles that display the encoded protein and contain its gene, providing a direct link between phenotype and genotype (Figure 2). This connection between genotype and phenotype enables large libraries of proteins to be screened and amplified in a process called *in vitro* selection (bio-panning).

Two key discoveries were essential for the development of antibody phage display technology. First, the demonstration that foreign DNA inserted into filamentous phage gene III (g3) is expressed as a fusion protein and displayed on the surface of the phage (Smith, 1985). Second, the successful expression of functional antibody fragments in the periplasmic space of *E. coli* (Better et al., 1988; Skerra and Plu[°]ckthun, 1988)

The advantages of phage display are easy to manipulate, rapid, inexpensive and appropriate for large scale production. Many recombinant anibodies with specific for various antigens such as human-self antigen, non-immunogenic antigens and small haptens were produced by phage display technology (Yau et al., 2003). This technique can also be used to study protein-ligand interactions (Cesareni, 1992), receptor and antibody-binding sites (Griffiths, 1993; Winter et al., 1994), and to improve or modify the affinity of proteins for their binding partners (Burton, 1995; Neri et al., 1995).

Large repertoires of antibodies (from 10^9 - to 10^{12}) were constructed and displayed on the surface of filamentous phages, thus exceeding the natural repertoire by several magnitudes (Hoogenboom et al., 1998). These libraries can then be easily used to select and purify specific phage particles bearing sequences with desired binding specificities from the non-binding variants.



Figure 2. Diagram of filamentous phage displaying scFv antibody fragments (Byrne et al., 2009). The phage consists of circular ssDNA surrounded by a coat protein. g8p (pVIII) is the major coat protein g3p, at the tip of the phage, is one of the minor coat proteins. The genes encoding variable domain of the scFv are fused to gene III in the genome of the filamentous phage. The scFv is displayed as a fusion to pIII protein at the top of the phage. The phage retains its ability to infect bacteria since the scFv is not fused to all pIII protein molecules. Four g3p molecules are illustrated in the figure, three of which display scFv molecules.

2.2.1 Biology of the filamentous bacteriophage

2.2.1.1 Structure

Filamentous phage have been identified which are able to infect a variety of gram negative bacteria. The best characterized of these nonlytic bacteriophage (M13, fl and fd) infect *E. coli* containing the F conjugative plasmid. The genomes of these three bacteriophage have been completely sequenced and are 98% homologous (Beck and Zink, 1981; Hill and Petersen, 1982; Van Wezenbeek et al., 1980). M13, Fl and fd are

collectively refered to as the Ff phage because of their dependence on the f plasmid for infection and their similarity. Ff phage has a single strand, covalently closed DNA genome which is enclosed in a long cylinder of approximately 7 nm wide by 900 to 2000 nm in length. Each genome encodes 11 genes, the products of which are listed in Table 3. The genes are grouped in the genome according to their functions in the life cycle of the phage. There are three groups of genes. One group (gene II, V, and X) encodes proteins required for DNA replication of the phage genome (Rapoza and Webster, 1995). The second group (gene III, VI, VII, VIII and IX) encodes the proteins that make up the capsid proteins for phage particle (Endemann and Model, 1995; Rapoza and Webster, 1993; Simons et al., 1981). Among these, pVIII is the major coat protein (with about 2700 copies per phage), while pIII, pVI, pVII and pIX are minor coat proteins (each with five copies per phage). Protein pIII, which is commonly fused with recombinant proteins (Smith, 1985), is presented as approximately 3-5 molecules. The last group (gene I, IV, and XI) encodes three proteins that are involved in the membrane associated with assembly of the bacteriophage. Gene products pI, pIV, and pXI are required for phage assembly (Horabin and Webster, 1988; Rapoza and Webster, 1995), but they are not part of the phage particle.

Gene	Function	Amino acid
II	DNA replication	410
Х	DNA replication	111
V	Binding of ssDNA	87
VIII	Major capsid protein	50
III	Minor capsid protein	406
VI	Minor capsid protein	112
VII	Minor capsid protein	33
IX	Minor capsid protein	32
Ι	Assembly	348
IV	Assembly	405
XI	Assembly	108

Table 3. Genes and gene products of fl bacteriophage (Webster, 1996).



A

Figure 3. (A) Electron microscopic of filamentous bacteriophage (http://www.mardre.com /homepage/mic/tem/samples/bio/virus/m13a.htm). (B) Filamentous phage fd architecture (Gao et al., 1999). Its single-stranded DNA genome is encased in a cylindrical protein coat. The long site of the cylinder consists of about 2,800 molecules of the major coat protein (pVIII). On one end, the particle is capped by about five copies of the proteins pVII and pIX. During the phage assembly process, this end emerges first from the bacterium. The incorporation of the minor coat proteins pIII and pVI on the other end of the particle terminates the assembly process. Protien III is the first protein to interact with the host cell during infection.

2.2.1.2 Life cycle of phage

The general stages to a phage life cycle are infection, replication of the viral genome, assembly of new viral particles and then release of the progeny particles from the host. Infection is a multistep process requiring interactions with the F conjugative pilus and the bacterial TolQ, R and A cytoplasmic membrane proteins. The infection begins when pIII attaches to the F pilus of *E.coli*. The phage genome is then transferred to the cytoplasm of the bacterial cell where resident proteins convert the single stranded DNA genome to a double stranded replicative form ("RF") (Webster, 1996). This RF undergoes rolling circle replication to make ssDNA, and serves as a template for expression of phage capsid proteins, and other phage proteins involved in the assembly process. Once the phage DNA has been fully coated with pVIII, the secretion terminates by adding the pIII and pVI cap (Marvin, 1998), and the new phage detaches from the bacterial surface. The bacterial host can continue to grow and divide, allowing this process to continue indentifinity. Filamentous bacteriophage does not produce lytic infection in their host.



Figure 4. Life cycle of filamentous phage. Phage particle infected to bacterial host cell by binding of pIII to the tip of the F pilus and entry of the single stranded DNA (ssDNA) into the cytoplasm. The ssDNA is converted by host enzymes to a double stranded replicative form (RF). The RF undergoes rolling circle replication to make ssDNA and also serves as a template for expression of the phage proteins gIIIp and gVIIIp. Phage progeny are assembled by packaging of ssDNA into protein coats and extruded through the bacterial membrane into the medium.

2.2.1.3 Phage vector

Phagemids are the most popular vector for display which is hybrids of phage and plasmid vectors. These vectors contain the origins of replications for both M13 phage and *E. coli*, gene coding part of pIII protein with the appropriate cloning site, and an antibiotic-resistance gene (Azzazy and Highsmith, 2002). Moreover, phagemids often contain an amber stop codon, to allow host specific expression of pIII fusion protein or soluble fusion partner, respectively (Hoogenboom et al., 1991), and a gene encoding one coat protein that will be fused to the foreign DNA that is to be expressed (Barbas et al., 1991; Kang et al., 1991).

If a phagemid vector is used, phage particles will not be released from the *E. coli* cells until they are infected with helper phage, which enables packaging of the phage DNA and assembly of the mature virions with the relevant protein fragment as part of their outer coat on either the minor (pIII) or major (pVIII) coat protein. The incorporation of many different DNA fragments into the pIII or pVIII genes generates a library from which members of interest can be isolated.



Figure 5. Schematic picture of a phagemid vector used to display antibody fragment on the surface of filamentous phage through pIII coat protein (Yau et al., 2003). Gene pIII is represented as a black box, the antibody fragment DNA insert as a grey box. The phagemid vector replicated in *E. coli*, but co-infection with a helper phage is required for the production of phage particles.
2.3 Phage display antibody library

Phage display has been used for isolation recombinant antibodies with a range of unique specificities and produces many high affinity antibodies without immunization. The surface of filamentous bacteriophage can be used displayed to antibody fragments in different formats, including Fab fragment (Cabilly, 1989), Fv fragment (Skerra and Plu[°]ckthun, 1988), scFv fragment (McCafferty et al., 1990), and diabody fragments(Holliger et al., 1993).

In general, two kinds of library can be used: naïve and immune. The naïve libraries are derived from natural unimmunized human rearranged V genes (Sblattero and Bradbury, 2000; Sheets et al., 1998), synthetic human V genes (Knappik et al., 2000) or shuffled V genes (Rader et al., 1998). Immune libraries are created from V genes from immunized humans (Dubrovskaya et al., 2007) or mice (Orum et al., 1993). Antibodies selected from immunized libraries tend to have much higher affinities for the antigen used for immunization when compare to a naïve library of equivalent size.

Key advantages of naïve library or single-pot repertoires include: (i) isolation of human antibodies to self, nonimmunogenic or toxic antigens; (ii) a single library can be used for all antigens; (iii) short time needed for antibody generation (2—4 rounds of selection in two weeks); and (iv) direct isolation of high affinity antibodies when very large repertoires are used (de Haard et al., 1999; Vaughan et al., 1996). Disadvantages of naïve libraries are: (i) low affinity of antibodies isolated from small sized libraries; (ii) the time needed to construct large libraries, and (iii) content and quality of the library are unequal.

Advantages and disadvantages of immune libraries include: (i) the long time requirement for animal immunisation, (ii) easier preparation than naive libraries (only medium sized repertoires, 10^{6} – 10^{7} of different phage particles are desired), (iii) the unpredictability of the immune response of the animal to an antigen of interest, (iv) the lack of immune response to some antigens, (v) the restrictions in generating human antibodies,

(vi) construction of a new library for each new antigen which increases the total time of the procedure by 1–3 month.

Thus, Phage display antibody has many advantages for production of antibody over conventional methods. Researcher can use appropriate kind of library, depending on their research objectives.

2.3.1 Phage display antibody expression formats

The antibody molecule has four polypeptide chains, two heavy (H) chains with molecular weights of 50 kDa and two light (L) chains (25 kDa molecular weight) linked by disulphide bonds. The chains have both constant (C) and variable (V) regions. The antigenbinding site of an intact antibody consists of a paired of variable domains VH and VL.

Antibodies have been displayed as functional binding molecules in different formats of antibody fragments. Effective display formats of antibodies are single-chain fragment variable (scFv) (Clackson et al., 1991; Marks et al., 1991), Fab (fragment antibody binding) fragments (Kang et al., 1991), Fv's (fragment variable) with an engineered intermolecular disulphide bound to stabilize the V_{H} . V_{L} pair (dsFv's) (Brinkmann et al., 1995), and diabody fragments (Holliger et al., 1993; McGuinness et al., 1996).

ScFv are relatively small (26-27 kDa), generally quite stable, and are encoded by a single gene, which simplifies genetic manipulations. The most common polypeptide linker to fuse the V_H and V_L chains together into a scFv antibody fragment is $(Gly_4Ser)_3$. The two variable domains can be connected either as V_H -linker- V_L or V_L -linker- V_H .

The other commonly used recombinant antibody fragments are Fab. Fab consist of two polypeptide chains, one containing the light chain variable and constant domains, V_{L} . $C_{\kappa \text{ or } \lambda}$, the other a truncated heavy chain containing the variable domain and one constant domain, V_{H} . C_{H1}. The illustrating picture of antibody fragment is shown in Figure 5, and the characteristics of different antibody fragments are presented in Table 4.



Figure 6. A cartoon representation of Immunoglobulin G (IgG) and antibody fragments. (Fj llman and Hall, 2005) Heavy chains are light yellow, whereas light chains are dark orange. Striped domains represent the variable domains ontaining complement determining regions (CDRs) in blue. The antigen binding fragment (Fab) is a large and stable fragment, whereas the variable fragment (Fv) is an unstable association of the two original variable domains. The single chain variable fragment (scFv) is recombinantly expressed with a linker to make it a small and stable antibody fragment.

Antibody fragment	Size (kDa)	Paratopes (valency)	Structure		
scFv	25-30	1	$V_{\rm H}$ and $V_{\rm L}$ domains are linked by a linker peptide. Changing the linker length can lead to the formation of diabody (60 kDa), triabody (90kDa), or tetrabody (120 kDa)		
Fv	25	1	$V_{\rm H}$ and $V_{\rm L}$ with no linker between the V domain		
Fab	50	1	Composed two chains of $V_{\rm H}\mathchar`-\mbox{C}_{\rm H}\mbox{1}$ and $V_{\rm L}\mbox{-}\mbox{C}_{\rm L}$		
F(ab) ₂	100	2	Two Fab molecules		
IgG	150	2	Parent antibody molecule consists of two heavy chain (V _H -C _H 1-C _H 2-C _H 3) and two light chain (V _L -C _L)		

Table 4. Characteristics of antibody fragments. Adapted from Azzazy and Highsmith

(2002).

2.3.2 Application of phage display antibody

The most successful applications of phage display antibody include (i) the de novo isolation of high affinity human antibodies from non-immune and synthetics libraries (Knappik et al., 2000); (ii) isolation of genes encoding antigen-specific antibodies from immunized animals and from hybridomas; (iii) the discovery of antibodies with unique properties from non-immune (Huie et al., 2001) and immune libraries from animal or human donors (Kramer et al., 2003). Antibody libraries displayed on phage are frequently used in the pharmaceutical industry for isolation of therapeutic antibody leads, for development into primarily anti-cancer or anti-inflammatory antibody drugs.

Phage display is particularly useful in cases where monoclonal antibodies could not be obtained by classical hybridoma technique such as antibodies against nonimmunogenic or toxic antigens such as Aflatoxin, which is hapten molecule (Daly et al., 2002; Moghaddam et al., 2001). Phage antibody technology can also be used to clone and rescue monoclonal antibodies from genetically unstable hybridomas. Phage antibody genes can be easily sequenced, mutated, and engineered to improve antigen binding (Azzazy and Highsmith, 2002). Finally, soluble recombinant antibodies can be produced quickly and economically and can be used as *in vitro* diagnostic reagents.

2.3.3 Selection of monoclonal antibody by in vitro bio-panning

The specifically binding clones are selected from phage display antibody libraries by using bio-panning method. This technique is used to enrich phage particles from phagedisplay libraries. The phage display antibodies are incubated with immobilized target or interested antigen. The unbound phages are removed by washing with washing buffer such as Phosphate Buffer Saline buffer (PBS) or PBS containing Tween. Phage antibodies bound to target can be eluted in different ways namally; acidic solution such as glycine-HCl buffer (Kang et al., 1991; Parmley and Smith, 1988), basic solutions like triethylamine (Marks et al., 1991), by enzymatic cleavage of protease site engineered between the antibody and gene III (Ward et al., 1996) or by competition with excess antigen (Clackson et al., 1991) or antibodies to the antigen (Meulemans et al., 1994). After elution, the specifically bound phages are amplified in *E.coli* cells for further rounds of bio-panning. Ideally, only one round of selection should be required, non-specific binding limits the enrichment that can be achieved per selection round. In most cases, several rounds (approximately 2-4 rounds) of selection and amplification are needed to select the best binders from the phage display library. Figure 7 show illustration diagram of in vitro biopanning method.



Figure 7. Overview of bio-panning method. The target or antigens of interest are immobilized on to immuno tube or ELISA plate. The phage display antibodies are leaved to bind to target protein. Unbound phages are washed away. After that, the bound phages are eluted appropriate methods as described in the text. The specific phages are amplified in *E.coli*. and are enriched in each round of selection. The binder phages are screened in ELISA technique.

2.3.4 Production of soluble scFv fragments

If the phagemid contain amber stop codon after the scFv gene, the soluble scFv can be produce by infecting *E. coli* strain HB2151 is nonsuppressor cells as they recognize an amber stop codon, engineered between the scFv gene and gIII in the phagemid. Thus only express the scFv will be expressed without the g3p protein. Phagemid that is used in this thesis also designed to introduce polyhistidine tag fused to the expressed scFv, thereby permitting rapid and simple protein purification by immobilized metal affinity chromatography. Depending on the isolated clone, soluble antibodies may be present in the culture supernatant, the bacterial periplasm, and inside the bacterial cells. All three fractions must be isolated and analyzed by western blot, using a commercially available conjugated antihistidine (His) tag antibody, to determine the location of the soluble antibodies. Soluble scFv are relatively simple to isolate, can be economically produced in bacteria in very large quantities (Kipriyanov et al., 1997; Mavrangelos et al., 2001), and do not entail complex refolding procedures (Sánchez et al., 1999).

CHAPTER III

MATERIALS AND METHODS

3.1 Materials

3.1.1 Phage

- Griffin.1, TomlinsonI and TomlinsonJ scFv libraries from the MRC, Cambridge Centre for Protein Engineering, Cambridge England were kindly provided by Prof. Greg Winter and Dr. Ian Tomlinson, respectively.
- KM13K07 helper phage NEBa (Vieira and Messing, 1987) and Propagated as described in the MRC phage display protocols.

3.1.2 Antigen

Aflatoxins conjugated with BSA and soluble Aflatoxin B₁, B₂, G₁, G₂ were prepared from *Aspergillus flavus* (Sigma, Germany).

Bovine albumin serum BSA was natural purified protein obtained from Fluka, USA.

3.1.3 Bacteria

E. coli TG1: Suppressor strain was used for cloning and for amplification of phage between rounds of panning (Supplied by MRC, Cambridge, UK).

E. coli HB2151: Non-Suppressor strain was used for inducing scFv fragment.

3.1.4 Antibodies

Anti-M13 Horseradish peroxidase (HRP): (Amersham-Pharmacia Biotech, Sweden)

Anti-Protein L peroxidase HRP (Sigma, Germany)

3.1.5 Instruments

Autoc	lave:	Hiclave HA-3000MIV, Hirayama, Japan				
Balan	ce:	Precisa 205A, Precisa Instruments, Switzerland				
		Precisa 3000C, Precisa Instruments, Switzerland				
Centrifuge machine:		Sorvall RC5C plus, Kendro laboratory Products, USA				
		Eppendrof centrifuge 5810 R, Eppendrof, USA				
Deep	freezer -70 °C:	Heto, Ultra Freeze, Denmark.				
ELISA	A reader:	Sunrise, TECAN, Austria				
Electr	oporator :	Eppendrof 2510, Eppendrof, USA				
Freezo	er -20 °C:	Heto, HLLF 370, Denmark.				
		MyBio LFT420, DAIREI, Denmark				
Gel dryer:		Drygel sr. SLAB GEL Dryer model SE1160, Hoefer Scientific				
		Instruments, USA				
Gel el	ectrophoresis	Mini Protean [®] 3 cell, BioRad, USA				
appar	atus:					
Heat	Box:	HB1, Wealtee Corp., USA				
Incub	ator shaker:	C24 Incubator shaker, New Brunswick Scientific, USA				
Incub	ator:	Memmert, BE 500, WTB Binder BD115, Shel-Lab 2020 Low				
		Temperature Incubator, Sheidon, USA				
Laminar hood		Holten LaminAir HBB 2448, Denmark.				
		BH2000 Series ClassII Biological Safety Cabinets,				
		BHA120 & BHA180, Clyde-Apac,				
Memb	orane transfer	Semi Phor, Hoefer Scientific instruments, USA				
Micro	centrifuge:	Iini spin plus, Eppendrof, USA				
		Eppendorf 54154, Eppendorf, Germany				

pH meter:	Ultra Basic pH meter, Denver Instruments, Germany				
Rotator:	Certomat TCC, B. Braun Biotech International, Germany				
Sequencing machine:	ABI prism model 310 Genetic Analyzer, Applied				
	Biosystems, USA				
Shaker:	Innova 2300 platform shaker, New Brunswick Scientific, UK				
	Certomat TC2, B. Braun Biotech International, Germany				
Sonicator:	Waken GE100 Ultrasonic processor, Japan				
Spectrophotometer:	Ultrospec 2000, Pharmacia biotech, UK				
Stirrer:	Variomag Electronicrührer Poly 15, Germany				
	Magnetic stirrer MSH300, USA				
Thermomixer:	Thermomixer compact, Eppendrof, USA				
Vortex:	Vortex-Genie 2 G506, Scientific Industries, USA				

3.2 Methods

3.2.1 Selection of scFv fragment from naive semi-synthetic scFv antibody library

3.2.1.1 Amplification of phage libraries for panning

The Tomlinson I and J library are contributed by MRC (Cambridge, UK). These are naive semi-synthetic scFv antibody library that modify at CDR3 region. The 500 μ l libraries stock (Tominson I and J) were added into 50 ml pre-warmed 2XYT (16g/L tryptone, 10g/L yeast extract and 86mM NaCl) containing 100 μ g/ml ampicillin and 1% (w/v) glucose and grown shaking at 37°C OD₆₀₀ 0.4 (approximately 2 hours). The 1×10¹¹ KM13 helper phage was added into 50 ml cultures and incubated at 37°C for 30 minutes. Then, the cultures were centrifuged at 4,000 rpm for 10 minutes and resuspended in 100 ml of 2XYT containing 100 μ g/ml ampicillin, 50 μ l/ml kanamycin and 0.1% (w/v) glucose. The cultures were incubated shaking at 30°C for overnight. The next day, the overnight

cultures were spun at 4,000 rpm for 30 minutes. The 20 ml PEG/Nacl (20% Polyethylene glycol 6000, 2.5 M NaCl) was added into 80 ml supernatant and left for 1 hour on ice and spun again at 4,000 rpm for 30 minutes. The pellet was resuspended with 4 ml Phosphate Buffer Saline (PBS) with 15% glycerol and stored at -70°C. These libraries were used for the selection of the specific scFv clones to aflatoxin.

3.2.1.2 Selection of phage antibodies (Bio-panning)

The selection was done on Nunc Maxisorp immuno tube (Nunc, Denmark). Aflatoxin B₁ conjugated with Bovine serum albumin (BSA) (Sigma, Germany) in 100 µl phosphate buffered saline (PBS) was immobilized on immune tube and incubated for overnight (16-18 hours) at 4°C. This experiment was done 3 round of selection. Amount of targets were used for selection in each round 50 µg, 15 µg and 5 µg respectively. The immune tube were washed three times with PBS and blocked with PBS supplemented with 4% (w/v) skimmed milk (4% MPBS) for 2 hours at room temperature. After, removal of the blocking solution, the tube was washed three times with PBS. The libraries were pre-incubated in the 4% (w/v) MPBS and 1% (w/v) BSA for 30 minute at room temperature. After incubation for 30 minute at room temperature, the libraries were added to immune tube and incubated at room temperature for 2 hours. Unspecific phages were washed away three times with PBS supplemented with 0.05% (v/v) Tween 20 (PBST) and three times with PBS. Bound phages were eluted with 500 µl of 1µg/ml trypsin buffer (50 µl trypsin in 450 µl PBS) incubating the immune tube at room temperature for 20 minutes with rotating.

The eluted phages were recovered by infection of 350 µl of exponentially growing *E. coli* TG1 at 37°C for 30 minute. Then, phages infected were performed 10-fold-serial dilution (10^2-10^6) and spread each dilution on TYE agar plates (10g/L tryptone, 5g/L yeast extract, 137mM NaCl and 15g/L Bacto-agar) supplemented with 100 µg/ml ampicillin and 1% (w/v) glucose. The agar plates were incubated at 37°C for overnight.

3.2.1.3 Further rounds of selection

1 ml 2XYT media were added to the agar plates that have a lawn of infected bacteria and loosen the cells with glass spreader. The scraped cells were kept as 15% (v/v) glycerol stock at -70°C. The 10 μ l of scraped bacteria were added into 10 ml 2XYT supplemented with 100 μ g/ml ampicillin and 1% w/v) glucose and incubated at 37°C with shaking until OD₆₀₀ was 0.4 (approximately 2 hours). After this procedure, the phages were rescued by 5×10¹⁰ helper phage KM13 (0.65 μ l) and incubating 37°C without shaking for 30 minutes. After centrifugation at 4,000 rpm, 4°c for 15 minutes, supernatant was removed and resuspend the pelleted bacteria with 5 ml of 2XYT containing 100 μ l/ml ampicillin, 50 μ l/ml kanamycin and 0.1% (w/v) glucose prior to incubated at 30°c with shaking for overnight. Next day, the overnight culture was centrifuged at 4,000 rpm, 4°C for 15 minutes and kept the supernatant. After, 1ml PEG/NaCl was added with 4 ml of the supernatant was removed and the pellet was resuspened in 100 μ l PBS for the next round of selection. Three rounds of selections were performed.

3.2.1.4 Polyclonal phage ELISA

The Immuno 96 MicroWellTM plate (Nunc, Denmark) was immobilized with 5 μ g of AflatoxinB1-BSA and 1% (w/v) Bovine serum albumin BSA in 100 μ l PBS buffer as control. After incubating plate at 4°C for overnight, the plate was washed 3 times with PBS. The wells were blocked with 4% (w/v) skimmed milk for 2 hours at room temperature. Then, wells were washed 3 times with PBS and added 50 μ l precipitated phage from the end round of selection in 50 μ l 4% (w/v) MPBS. The plate was incubated at room temperature for 2 hours after that, phage solution was discarded. Unspecific phages were washed away three times with PBST, followed by three times with PBS and incubated for 1 hour with a 1:5,000 dilution of a mouse anti-M13 phage-horseradish peroxidase (HRP) conjugate (Amersham-Pharmacia Biotech, Sweden) in 2% (w/v) MPBS. The plate was

washed again as described earlier. The 100 μ l 3,3', 5,5'- Tetramethybenzidine (TMB; Sigma, Germany) substrate was added. The reaction was developed to be a blue color at 37°C for 30 minutes and was then stopped using 50 μ l/well of 10% (v/v) HCl. The absorbance was read at 450 nm, using a Sunrise absorbance reader (TECAN, Austria).

3.2.1.5 Preparation of monoclonal phage

In the 1st and 3rd selection round, individual clone was randomly picked out from the TYE plate to different well (100 μ l of 2XYT with 100 μ g/ml ampicillin and 1% (w/v) glucose was added) of immuno plate (Nunc, Denmark). After culturing, small inocula (5 μ l) from each well was transferred to a second ELISA plate containing 200 μ l of 2XYT with 100 μ g/ml ampicillin and 1% (w/v) glucose per cell. The original ELISA plate or master plate was added with glycerol to a final concentration of 20 % and kept at -20°C. The transferred plate was shaken at 37°C for 2 hours, and phages were rescued in *E.coli* TG1 cells by adding 10¹⁰ helper phage to each well, the plate was shaken at 37°C for 1 hour before spinning the plate at 4,000 rpm for 10 minutes. The supernatant was thrown away and the pellet was resuspended in 200 μ l of 2XYT containing 100 μ g/ml ampicillin and 50 μ g/ml kanamycin, and cultured at 30°C 250 rpm for overnight. The overnight culture was spun at 4,000 rpm for 10 minutes and 50 μ l of the supernatant containing phage was used in monoclonal phage ELISA.

3.2.1.6 Confirmation of binding activity of monoclonal phage clones by ELISA

Immuno 96 WellsTM Plate was coated and blocked as in polyclonal phage ELISA. After incubation for 2 hours at room temperature, the wells were washed 3 times with PBS. Then 50 μ l of phage supernatant and 50 μ l 4% (w/v) MPBS were added into each well of the plate. Unspecific phages were washed away 3 times with PBST and 3 times with PBS. After stand on the bench 2 hours, the plate was washed 3 times with PBST and followed 3 times with PBS. Subsequently, 100 μ l HRP-anti-M13 (1:5000 dilution in 2% (w/v) MPBS) was added to each well. After incubation for an additional 1 hour at room temperature, the wells were washed again and 100 μ l substrate solutions (TMB, SIGMA) was added to each well and incubated at 37°C for 30 minutes. The reaction was stopped with 10% (v/v) HCl. The yellow color was developed after adding HCl. Absorbance was measured at 450 nm.

3.2.2 Expression and purification of scFv antibody

3.2.2.1 Induction of soluble antibody fragment

In total, 35 positive clones from Tomlinson J and 61 clones from Tomlinson I library that shown high activities were infected into a non-suppressor strain HB2151 which reads the amber codon as a stop signal, thereby solubly expressing the antibody fragment's. The 5 µl of phage particles from master glycerol stock were infected to 150 µl exponentially growing E.coli HB2151 bacterial at an OD₆₀₀ of 0.4, and incubated at 37°C for 30 minutes. After that, the infected bacteria were streak on TYE agar plates containing 100 µg/ ml ampicillin and 1% (w/v) glucose. The plates were incubated at 37°C for overnight. The next day, single colonies were picked with pipette tips and used to inoculate 100 µl of 2xTY, containing 100 µg/ ml ampicillin and 1% (w/v) glucose in 96 immuno plate. The culture was incubated with shaking at 37°C for overnight. The following day, the 8 µl cultured were transferred to 96 Deep Well immune plate containing 800 µl 2XYT, 100 μ g/ ml ampicillin and 0.1% (w/v) glucose and incubated shaking at 37°C until OD₆₀₀ reached 0.9. The master plate was added glycerol to a final concentration of 15% and stored at -70°C. The 100 μl 2XYT, 100 μg/ ml ampicillin and 9 mM isopropyl-β-dthiogalactopyranoside (IPTG), achieving a final concentration of 1mM were added into 96 deep well plate. The incubation was continued at 30°C for overnight. The secreted antibody could be found in the supernatant.

3.2.2.2 Small scale production of soluble scFv antibody

A single colony of HB2151 *E.coli* cells containing the pIT2 bearing the 3I-F6 encoding gene was inoculated into 5 ml 2XYT broth containing 100µg/ml Ampicillin, 1% (v/v) glucose and was grown overnight at 37°C, with shaking at 250 rpm. The next day, overnight culture was inoculated into 15 ml 2XYT medium containing 100µg/ml Ampicillin, 0.1% (v/v) glucose at 37°C for 3 hours with shaking at 220 rpm. After 3 hr, the culture was induced with 1mM IPTG and incubated overnight at 30°C with shaking at 220 rpm. The culture was centrifuged at 4,000 rpm for 20 min at 4°C and the supernatant was discarded. The pellet was resuspended in 1ml of lysis buffer (20mM Tris; pH 8.0, 150mM NaCl and 1mM PMSF). The cells were sonicated twice on ice for 30sec at 50% output. They were then centrifuged in microcentrifuged tube at 14,000 rpm at 4°C for 10 minutes. The supernatant is cell lysate fraction and was used to purify scFv fragment.

3.2.2.3 Small scale purification by using Ni-NTA resin

ScFv antibody containing a hexahistidine tag was purified using Ni⁺⁺ ions immobilized on resin by covalent linkage to nitrilotriacetic acid (NTA) (Novagen). The high affinity interaction between the histidine residues and the Ni⁺⁺ ions allow the specific purification of histidine tagged recombinant proteins. The antibody fragments were eluted with imidazole, which competes for binding to the Ni⁺⁺ ions. Before use of the Ni⁺⁺ ions, the resin was equilibrated with lysis buffer (20mM Tris; pH 8.0, 150mM NaCl, 1mM PMSF). Thus, the same pH and buffer composition with that of the Ni⁺⁺ ions resin ensuring that the sample would bind properly. The cell lysate fraction was incubated in a microcentrifuge tube with 200 μ l Ni-NTA resin for 2 hour to allow the binding of hexahistidine tag to Ni-NTA resin. The solution was centrifuged at 2000 rpm for 5 min. The flow-through from the wash was collected for further analysis. The washing buffer 1 (20mM Tris; pH 8.0, 150mM NaCl, 5mM imidazole) was added into the microcentrifuge tube and spun again. The Ni-NTA matrix was washed with washing buffer 2 (20mM Tris; pH 8.0, 150mM NaCl, 20mM imidazole). After that, the elution buffer (20mM Tris; pH 8.0, 150mM NaCl, 250mM imidazole) was added into the tube and they were centrifuged again. The scFv fragment should be presented in the eluted fraction. To monitor antibody purity, SDS-PAGE electrophoresis was used. To determine the efficiency of purification, the flow through fraction, wash fraction and elution fraction were compared by SDS polyacrylamide gel and western blot analysis. ELISA was used to detect the antibody binding.

3.2.2.4 Large scale production of soluble scFv antibody

A single colony of HB2151 *E.coli* cells containing pIT2 bearing the 31-F6 encoding gene was inoculated into 5 ml 2XYT broth, containing 100μ g/ml Ampicillin, 1% (w/v) glucose and was grown overnight at 37°C, with shaking at 250 rpm. The next day, an overnight culture was inoculated into 500 ml Terrific broth, containing 100μ g/ml ampicillin and 0.1% (w/v) glucose at 37°C for 4 hours (OD₆₀₀ is 0.9) with shaking at 220 rpm. And then, the culture was induced with 1mM IPTG and incubated overnight at 30°C at 220 rpm. The culture was poured into 250 ml centrifuge bottles and centrifuged at 10,000 rpm for 20 min at 4°C and the supernatant was discarded. The pellet was resuspended in 30 ml of sonication buffer (1xPBS, 0.5M NaCl and 20mM Imidazole). The resuspended solution was aliquoted in 1 ml fractions into 2 ml centriguation tubes. The cells were sonicated on ice for 45second min at 40% output for three times. The cells were centrifuged in microcentrifuged tube at 14,000 rpm at 4°C for 10 minutes. The cell lysate fractions were passed through 0.2 μ m filter for sterisation.

3.2.2.5 Purification scFv fragment by immobilized metal affinity chromatography (IMAC) using Ni-NTA

The column was prepared by loading 4 ml of Ni-NTA resin (Novagen, USA) and left to form a 2 ml packed resin. The column was equilibrated with running buffer (1xPBS, 0.5M NaCl, 20mM Imidazole and 1% (v/v) Tween). The filtered supernatant was transferred to prepared IMAC columns and passed through the column twice and the flow through was collected. The column was washed once with 30 ml running buffer to remove any non-specifically proteins. The scFv was eluted with 20 ml of 100mM sodium acetate, pH 4.4 and collected in 1.5 ml microcentrifuge tubes (final volume 500µl in each tube) containing 50µl 10x PBS (filtered through a 0.2 µm filter) and 50 µl 100 mM NaOH. The eluted antibody was buffer exchange into PBS and concentrated by ultrafiltration (Vivaspin, Sartorius) and then the scFv solution was dispensed in 20 µl volumes and stored at -20°C until required. SDS-PAGE electrophoresis was used to monitor antibody purify. The efficiency of purification, the flow through fraction, wash fraction and elution fraction were compared by SDS polyacrylamide gel and western blot analysis. ELISA was used to detect the antibody binding.

3.2.2.6 ScFv antibody ELISA

Immuno 96 WellsTM Plate was performed by coating microtitre plates with 5 μ g of AflatoxinB₁-BSA in a volume of 100 μ l and with 1% (w/v) BSA as control. The ELISA plate were washed 3 times with PBS and blocked with PBS supplemented with 3% (w/v) BSA. After incubation for 2 hours at room temperature, the wells were washed three 3 times with PBS. Then 50 μ l of scFv antibody and 50 μ l 3% (w/v) BSA were added into each well of the plate. After stand on the bench 2 hours, the plate was washed 3 times with PBST and followed 3 times with PBS. Subsequently, 100 μ l Protein L peroxidase (1:5000 dilutions in 3% (w/v) BSA) was added into each well. After incubation for an additional 1 hour at room temperature, the wells were washed again and 100 μ l substrate solutions (TMB, SIGMA) was added to each well and incubated at 37°C for 30 minutes. The reaction was stopped with 10% (v/v) HCl. The yellow color was developed after adding HCl. Absorbance was measured at 450 nm.

3.2.2.7 Sequence analysis

Plasmid DNA from three different clones were extracted and purified from overnight culture by using a commercially plasmid preparation kit (Mini Preps: Qiagen, Germany) and were completely sequenced by Macrogen (Seoul, Korea) using primers: For VH use link seq new CGA CCC GCC ACC GCC GCT G and For Vκ use pHEN seq CTA TGC GGC CCC ATT CA. Immunogenic analysis of variable regions was done using IgBLAST and V-Base programme.

3.2.2.8 SDS polyacrylamide gel

SDS polyacrylamide gel was used to determine the efficiency of purification, the flow through fraction, wash fraction and elution fraction were compared. A SDS polyacrylamide gel, consisting of 12 % running gel and 5 % stacking gel in SDS-PAGE electrophoresis buffer. The samples were boiled for 5 min before they were loaded onto the gel. The proteins were detected with Coomassie Brilliant Blue (CBB) staining solution for 15 minutes, followed by destaining with destaining solution (40% (v/v) methanol, 10% (v/v) glacial acetic acid). The gel was then dried using the Gel Drying System (Drygel sr. SLAB GEL Dryer model SE1160, Hoefer Scientific instruments, USA).

3.2.2.9 Western Blot analysis

For western blot analysis, samples will be run on SDS-PAGE gel and the scFv were blotted on to nitrocellulose membrane or PVDF membrane. After run SDS-PAGE, The membrane and filter papers were cut to the same size of gel. All of them were incubated in transfer buffer (25 mM Tris, 192 mM glycine, 10% (v/v) methanol) for 30 minutes at room temperature. The proteins on the SDS polyacrylamide gel were transferred to membrane using a membrane transfer machine (Semi Phor, Hoefer Scientific Instruments, USA). And, membrane was blocked with 3% (w/v) BSA in PBS for 1 hour at room temperature. Then, the membrane was washed 3 times with PBST following 3 times with PBS by rotating for 10 minutes. After that, the membrane was incubated with Protein L peroxidase (a dilution of 1:5000 in 3% (w/v) BSA) for 1 hour. Membrane was washed the same as previous step. The 3,3',5-5'-Tetramethylbenzidine (TMB) substrate (Sigma, Germany) was used according to the manufacturers protocol for developing the blue color on the membrane. Stop reaction was done by adding 1M EDTA.

3.2.2.10 Competition ELISA for anti Aflatoxin B₁

For the inhibition assay, $5\mu g$ of AFB₁-BSA was coated on an ELISA plate. The antibodies, at dilutions previously determined by an antibody titre, were pre-incubated with varying concentrations of soluble aflatoxin-B1 from 5 μg to 0.019 μg . After incubation at 37 °C for 30 minutes, the mixture was then transferred to the previously coated and blocked microtitre plate and incubated for 1 hour. The unbound antibodies were washed away 3 times with PBS containing 0.05% (v/v) Tween 20 and 3 times with PBS. The secondary antibody (Protein L peroxidase) was added to wells and incubated for 1 hour at room temperature. Following this, the substrate solution was added and incubated at 37 °C for 30 minutes, 10% (v/v) HCl was used to stop the reaction. The reaction was then measured by reading the absorbance at 450 nm using the Tecan plate reader.

3.2.3 Cloning of scFv fragment into vector containing alkaline phosphatase gene

The anti-AFB₁ selected from Tomlinson I library (TomI-F6) and YAMO 1 library (YM1-C3) were engineered to create scFv-AP fusion.

3.2.3.1 Cloning of scFv fragment into pKP300AIII

pKP300 Δ III was generated from Brian, Kay Lab (UIC, USA). It is expressions vector that containing alkaline phosphatase gene. The scFv fragments DNA were inserted into pKP300 Δ III between *NcoI* and *Not*I sites. The DNA of scFv fragments and pKP300 Δ III vector was double digested with *NcoI* (10U/µl, NEB, USA) and *Not*I (10U/µl, NEB, USA) enzymes, to generate compatible sticky ends. The digestion reactions of scFv fragments and pKP300 Δ III vector were performed separately, each in a total volume of 30

µl. The reaction mixtures consisted of 1 µg of insert DNA, 0.5 µg of vector DNA, 1x NEB buffer 3, 1 µg/ml BSA, 5U of Ncol (10U/µl, NEB, USA) and 5U of NotI (10U/µl, NEB, USA). The reactions were incubated at 37°C for 3 hours. The reactions were heat inactivated at 65°C for 20 minutes. After heat inactivate enzyme, the NcoI/NotI digested vector was dephosphorylated by adding 0.5 µl of CIP enzyme (10U/µl, NEB, USA) and incubated at 37°C for 1 hour. The inserts and vectors were separated from stuffer fragments by gel electrophoresis followed by Wizard clean up kit (Promega, USA). The scFv DNA was ligated into pKP300ΔIII vectors at a 3:1 ratio. The total volume of ligation reaction was 10 μl, which consisted of 24.6 μg of inserted DNA, 12.1 μg of pKP300ΔIII vector, 1x T4 DNA ligase buffer and 1 µl of T4 DNA ligase (400U/µl, NEB, USA). The ligation reaction was incubated at 25°C for 2 hours and heat inactivate enzyme at 65°C for 10 minutes. Then, the ligation reaction was transformed into 100 µl of E.coli TG1 cells by electroporation. The electroporation was performed at 1.8 kV, 18 μ F, 200 Ω and τ approximately 4 msec using an electroporation machine (Eppendrof 2510, Eppendrof, USA). The cuvette was added immediately with 900 µl of 2XYT medium room temperature. The 1 ml of combined transformed cells was incubated at 37°C for 1 hour. After that the transformed cells were spread on LB plates, containing 100 µg/ml ampicillin and incubated for overnight at 30°C. Five of individual clones were picked and analyzed by double digestion with Ncol/Notl and ELISA format.



Figure 8. Map of pKP300ΔIII (Krittika, personal contact). The arrow shows the position of PhoA promoter, alkaline phosphatase and ampicillin resistance genes.

3.2.3.2 Digestion to check positive clones

Five individual colonies were picked with sterilized toothpicks into 5 ml of low phos media (30 mM (NH₄)₂SO₄, 2.4 mM (CH₂COONa)₂.2H₂O, 14 mM KCl, 5.36g/L yeast extract, 5.36g/L Hy-case SF casein hydrolysate, 7mM MgSO₄, 14mM glucose, pH 7.3 adjust by KOH). After incubation for 18 hour, the samples were centrifuged at 4000 rpm for 10 minutes. The supernatants were tested the activity specifically to aflatoxin on ELISA format. The DNA of each clones were extracted DNA by using Qiagen mini prep (Qiagen, Germany). The digestion reactions of DNA were performed, in a total volume of 10 μ l. The reaction mixtures consisted of 1 μ l of DNA, 0.5 μ g of vector DNA, 1x NEB buffer 3, 1 μ g/ml BSA, 1U of *NcoI* (10U/ μ l, NEB, USA) and 1U of *NotI* (10U/ μ l, NEB, USA). The reactions were incubated at 37°C for 1 hour and were run on 2% (w/v) agarose gel in TAE

buffer at 100 volts for 40 minutes. The positive clones were shown 2 bands in approximately size 5kb and 800 bp.

3.2.3.3 ScFv-AP ELISA

Immuno 96 WellsTM Plate was performed by coating microtitre plates with 5 μ g of AflatoxinB₁-BSA in a volume of 100 μ l and with 1% (w/v) BSA as control. The ELISA plates were washed three times with Tris-buffered saline (TBS; 25mM Tris-HCl, pH 7.5, 140 mM Nacl, 3mM KCl) supplemented with 0.05% (v/v) Tween 20 (TBST) and blocked with TBS supplemented with 3% (w/v) BSA. After incubation for 1 hour at room temperature, the wells were washed 5 times with TBST. Then 50 μ l of scFv-AP supernatant and 50 μ l 3% (w/v) BSA were added into each well of the plate. After standing on the bench for 2 hours, the plate was washed 5 times with TBST. Subsequently, 100 μ l p-Nitrophenyl phosphate disodium salt hexahydrate (PNPP; SIGMA, USA) was added into each well. After incubation for 20 minute at room temperature, an absorbance was measured at 405 nm.

3.2.3.4 Expression and purification scFv-AP by using Ni-NTA column

Individual colony was picked with sterilized toothpicks into 10 ml of low phos media. After incubation for 20 hour, the samples were centrifuged at 4000 rpm for 10 minutes. The pellet was resuspended in 1 ml of 1X BugBuster lysis buffer (Novagen, USA) and incubated on ice for 20 minutes. The solution was centrifuged at 14,000 rpm for 10 min. The supernatant was kept to run SDS-PAGE. The Ni-NTA column was equilibrated with 600 µl of lyses buffer and spun at 2,000 rpm for 2 minutes. The cell lysate fraction was added into column and spun at the same speed again. The flow through was collected for SDS-PAGE. The washing buffer (50mM NaH₂PO₄, 300mM NaCl, 20mM imidazole; pH 8.0) was added to column and spun. The column was washed anther two times. After that, 200 µl the elution buffer (50mM NaH₂PO₄, 300mM NaCl, 250mM imidazole) was added into the column and they were centrifuged again. The scFv fragment should be presented in

the eluted fraction. To monitor antibody purity, SDS-PAGE electrophoresis was used. To determine the efficiency of purification, the flow through fraction, wash fraction and elution fraction were compared by SDS polyacrylamide gel.

3.2.3.5 Extraction soluble Aflatoxin from peanut

The 20 grams of peanut were blended with mortar. After that, 100 ml of 70% (v/v) methanol were added into sample. The sample was shaking at 180 rpm for 30 min. Then, the sample was passed though filter papers. Sample was diluted with 5% (v/v) methanol in TBST before using in competitive ELISA analysis.

3.2.3.6 Cross reactivity scFv-AP fusion in competitive ELISA

ScFv-AP antibody as assayed against a range of soluble aflatoxins B_1 , B_2 , G_1 and G_2 . Stock solutions of aflatoxin were prepared in 100% methanol and diluted using TBST containing 5% (v/v) methanol. The assays were immobilized and blocked as described in 3.2.3.3. After washing 3 times with TBST, the dilution antibodies were pre-incubated with varying concentrations of soluble aflatoxin from 5 µg to 0.019 µg and 1:2 dilution of peanut's extraction. After incubation at 37 °C for 30 minutes, the mixture was then transferred to the previously coated and blocked microtitre plate and incubated for 1 hour. The unbound antibodies were washed away 5 times with TBST. Following this, the PNPP substrate solution was added and incubated at room temperature for 30 minutes. The reaction was then measured by reading the absorbance at 405 nm using the Tecan plate reader.



Figure 9. Schematic diagrams of competitive ELISA to test the binding of scFv-AP to soluble AFB₁. The AFB₁-BSA was immobilized on ELISA plate. The mixtures of scFv-AP and soluble aflatoxin were added to plate. The binding of soluble scFv-AP to AFB₁-BSA, hence the ELISA, depend on the amount of soluble AFB₁ which is a competitor. In the presence of high amount of soluble AFB₁, low scFv-AP can bind to AFB₁-BSA (low signal). On the contrary, if the amount of soluble AFB₁ is low, ELISA signal will be high.

CHAPTER IV

RESULTS AND DISCUSSIONS

4.1 Polyclonal phage ELISA

The human naïve scFv Tomlinson I and J libraries were used for selection of scFv specific to aflatoxin. Three rounds of panning were performed. After scraped cell in each rounds of selection, the polyclonal phages that bind to aflatoxin B_1 were analyzed by polyclonal phage ELISA to monitor the enrichment of the phage population of the selection process using mouse anti-M13 antibodies conjugated to HRP for detection. Polyclonal phages selected from Tomlinson I library showed positive signal in every rounds of selection (Figure 10A). The results indicated that the phage display antibodies specific to aflatoxin B_1 conjugated BSA were enriched in each round of selection (Figure 10) whereas, screening of polyclonal phage antibodies from Tomlinson J library did not show any enrichment after second round of selection (Figure 10B). Yang et al. reported the signals of polyclonal phage antibodies against AFB₁ increased after each round of selection which similar as this result.

44

A

B



Figure 10. Polyclonal phages ELISA of (A) Tomlinson I library, (B) Tomlinson J library specific to AflatoxinB₁-BSA (AFB₁-BSA) were detected with HRP conjugated mouse anti-M13 antibodies. After each round of selection, polyclonal phages were prepared and analyzed for binding to AFB₁-BSA. 1% BSA was used as negative control in this experiment. All analyses were performed in duplicate.

4.2 Selection of scFv fragments specific to aflatoxin by using biopanning

The Tomlinson I and J libraries were grown to use for selecting scFv specific to Aflatoxin. Three rounds of selection were performed. Titration of eluted phage from Tomlinson I and J library showed that there was significant enrichment in their number between the first and second rounds of panning, but not in the third round (Table 5).

The results in Table 5 showed that the phages were amplified the same antibody formats after 3rd round of selection and antibody can bind to AFB₁-BSA more than soluble AFB₁. This problem can be solved by;

- (i) decreasing the coating concentration of antigen in each round (Strachan et al., 2002) or increasing more times of washing.
- (ii) The free aflatoxin may be used to elute the aflatoxin specific antibodies
- (iii) The phage antibodies are pre-incubated with the carrier protein prior to incubation with the aflatoxin-protein conjugate. Subtractive panning should remove any antibody binding to the carrier protein, but not those antibodies binding to the bridging group linking the protein and the afaltoxin (Sheedy et al., 2007).

Nonetheless, such panning optimization strategies have led to the isolation of antibodies with higher specificity and lower levels of cross-reactivity (Chames et al., 1998).

However, with naïve libraries, the isolation of antibodies binding strictly to the free hapten such as aflatoxin is difficult, and only few publications have addressed the problems associated with the selection of antibodies that only recognize the hapten when the latter is conjugated with its carrier protein (Moghaddam et al., 2001). This study can use standard bio-panning method to isolate antibody that bind to soluble aflatoxin B₁, when using BSA-conjugated toxin as a target, as demonstrated by competitive ELISA.

scFv library source	Tomlinson I	Tomlinson J
Number of clone after 1 st panning	3.2×10 ³	4×10 ²
Number of clone after 2 nd panning	1×10 ⁵	1.4×10 ⁵
Number of clone after 3 rd panning	2.2×10 ⁵	2×10 ⁶
Number of positive phage clones after 1 st round	6/96	8/96
Number of positive phage clones after 3 rd round	71/96	32/96
Number of soluble scFv expessed	19/62	2/34
Number of scFv fragments that bound soluble Aflatoxin B_1	3/8	NA
Number of scFv producing different clones	1/3	NA

Table 5. Overview of selection of antibodies against Aflatoxin B₁-BSA.

* The number of positive clones / the number of screened clones. NA is an abbreviation for not applicable.

4.2.1 Monoclonal phage ELISA in 1st round of selection.

The phage clones specific to aflatoxin B_1 - BSA from 1st round were screened for the positives clones by ELISA method using mouse anti-M13 antibodies conjugated to HRP for detection. The phage clones were scored as positive by ELISA when the absorbance value on AFB₁-BSA coated wells was at least two times higher than binding of phage to 1% (w/v) BSA coated-wells. Six positives individual clones from Tomlinson I (Figure 11A) and eight clones from Tomlinson J (Figure 11B) from 1st round of panning showed positive signal from 96 picked clones from each library.



Figure 11. Binding of specific phage to AFB₁-BSA (Aflatoxin B₁ conjugated with BSA) from 1st round of panning. Clones were selected from (A) Tomlinson I antibody library and (B) Tomlinson J antibody library. Signal developed using TMB substrate and monitored by optical density at 450 nm. Six positive clones (B12, D6, D10, D11, F9 and F11) from Tomlinson I and eight clones (C1, C2, A3, E4, F5, H9, F10 and G10) from Tomlinson J library showed specific binding to AFB1-BSA and not to 1% (v/v) BSA (negative control).

4.2.2 Monoclonal phage ELISA 3rd round of selection.

After 3^{rd} round of selection, individual 96 clones were randomly picked and their binding were tested by phage ELISA using mouse anti-M13 antibodies conjugated to HRP for detection. Seventy-one positive clones from Tomlinson I (Figure 12A) and thirty-two clones from Tomlinson J (Figure 12B) library showed specific binding to AFB1-BSA. The positive phage clones shown absorbance two times higher than the phage binding to 1% (w/v) BSA. These positive phage clones were enriched from 1st round of selection.



Figure 12. Binding of specific phage to AFB₁-BSA (Aflatoxin B₁ conjugated with BSA) from 3rd round of panning. Clones were selected from (A) Tomlinson I antibody library and (B) Tomlinson J antibody library. Data is expressed as absorbance at 450 nm. Seventy-one positive clones from Tomlinson I and thirty-two clones from Tomlinson J library showed specific binding to AFB1-BSA and not to 1% (v/v) BSA (negative control).

4.3 Production of soluble antibody fragments

From total, 62 positive clones from Tomlinson I and 34 clones from Tomlinson J library were brought forward to be assessed in induction of soluble antibody formats. The 96 selected clones were infected into a non-suppressor *E.coli* strain HB2151 (non-supressor strain) which reads the amber codon (TAG) as a stop signal, thereby solubly expressing the antibody fragment's. The scFv-gene III fusion is under the control of the LacZ promoter. Expression was induced by the addition of IPTG. The scFv fragments are then secreted in the supernatant or cell lysate.

ScFv fragments specific to aflatoxin B_1 were determined by ELISA. Bound scFv were detected using an anti-protein L-HRP labeled antibody. The substrate used to detect bound secondary antibody was TMB (Fluka) and the reaction was stopped with 10% (v/v) HCl. The reaction was measured by reading the absorbance at 450 nm with Tecan plate reader.

Nineteen clones were shown to secrete scFv antibody specifically binding to AFB₁-BSA detected in cell culture supernatants by ELISA (Figure 13). From these, eight clones which showed the highest signal were brought forward for competitive studies.



Figure 13. Binding of scFv antibody fragments to AFB₁-BSA. Sixty-one positive clones from Tomlinson I and thirty-five clones from Tomlinson J library were induced and scFv fragments binding to AFB₁-BSA in ELISA was analysed. The data is expressed as absorbance at 450 nm. Nineteen clones were shown to secrete scFv antibody specifically binding to AFB1-BSA and not to 1% (v/v) BSA (negative control).

4.4 Competitive ELISA of scFv fragments

The eight positive clones that shown highest absorbance from the earlier experiment were tested the binding to soluble AFB₁ using competitive ELISA. The positive result was determined if the absorbance decreased when the concentrations of soluble AFB₁ increased. TomI-F6, TomI-H2 and TomI-E9 were inhibited from binding immobilized aflatoxin by free toxin, indicating specific binding to soluble AFB₁ (Figure 14).



Figure 14. Binding of scFv fragments to AFB₁-BSA in an inhibition assay. The three clones exhibiting specific binding for free aflatoxin are TomI-H2, TomI-F6 and TomI-E9. This can be seen from the decreasing absorbance shown when higher concentrations of aflatoxin B₁ are present.

4.5 Sequence analysis of selected clones

Three clones (TomI-F6, TomI-H2 and TomI-E9) that can be inhibited by soluble aflatoxin B_1 were selected for sequence analysis. After sequencing, the origin of the V gene and complimentarity determining region 3 (CDR3) lengths were determined using V BASE immunoglobulin V gene database (Tomlinson et al., 1996) and IgBLAST program. The analysis showed that the three clones of phages displaying anti-aflatoxin B_1 , TomI-F6, TomI-H2 and TomI-E9 were identical. The closest germline sequence for anti-aflatoxin scFv antibody could be identified by comparison to database. The deduced amino acid sequences of V_H and V_L are illustrated in Figure 15. According to the amino acid sequences, the V_H was classified as human heavy chain family V_H3 derived germline V3-

23*05 while V_L fell into human kappa light chain family Vk1 derived germline V1D-39*01. The numbers of different amino acids from germ line of V_H and V_L of phage clone were 7 and 2 amino acids, respectively. The CDR3 of the binders had lengths 9 amino acids for the VH genes, and for the VL genes as shown in Table 6. TomI-A8, TomI-A11, YM1-C5 and YM-D2 that the clones not bind to soluble aflatoxin were identified sequences (Table 6). The CDRH3 is critical importance in the interaction of an antibody with its respective antigen. This region, along with the CDRL3, is considered to make the most significant contributions to affinity and specificity.

(A) V_H

______ FWR 1 _____ CDR H1 ____ FWR 2 _____ CDR H2 EVQLLESGGGLVQPGGSLRLSCAASGFTFS **SYAMSW** VRQAPGKGLEWVS <u>SISNAGTY</u>TYY<u>V</u>DSVKGR

______FWR 2 ______ CDR H3 _____ FWR 4 _____ FTISRDNSKNTLYLQMNSLRAEDTAVYYC AKNYTTFDY WGQGTLVTVS

(B) V_L

_____ FWR 1 _____ CDR V1 _____ FWR 2 _____ CDR V2 _____ FWR 3 _____ DIQMTQSPSSLSASVGDRVTITC **RASQSISSYLNW** YQQKPGKAPKLLIY <u>S</u>ASSLQSGV PSRFSGSGSGTD

FTLTISSLQPEDFATY YC QQGANYPNT FGQGTK VEIK

Figure 15. Amino acids sequence alignment of scFv antibody fragments (TomI-F6) selected against AFB₁.The heavy (A) and light chain (B) complementarity determining regions (CDRs) were defined according to IgBLAST program and are shown in bold letters. The underline letters has shown amino acids difference from germ line.

Table 6. Amino acids sequence of selected clones.

Clone	FR1	CDR1	FR2	CDR2	FR3	CDR3	Germline	Amino acid different from germline	Family
VH									
TomI-F6	EVQLLESGGGLVQPGGSL RLSCAASGFTFS	SYAMSW	VRQAPGKGLE WVS	SISNAGTYTY YVDSVKGR	FTISRDNSKNTLYLQMNSLR AEDTAVYYC	AKNYTTFDY	IGHV3-23*05	7	VH3
TomI-A8	EVQLVESGGGLVQPGGSL RLSCAASGFTFS	SYAMSW	VRQAPGKGLE WVS	SIAGSGNYTS YADSVKGR	FTISRDNSKNTLYLQMNSLR AEDTAVYYC	AEASGAFDY	IGHV3-23*04	6	VH3
TomI-A11	EVQLVESGGGLVQPGS*T RLSCAASGFTFS	SYAMRW	VH*APGKGLE WVS	AISGYGSYTT YADSVKGR	FTISRDNSKNTLYLQMNSLR AEDTAVYYC	AKGGYSFDY	IGHV3-23*04	11	VH3
YM1-C3	QVQLVQSGAEVKKPASVK VSCKAS	GYTFTSYA	MHWVRQAXGQ RLEWMGW	INAGNGNT	KYSQKFQGRVTITRDTSAST AYMELSSLRSEDTAVYYCAR	ADDYGSGSYG FDY	IGHV1-3*01	1	VH1
YM1-C5	EVQLVESGGGLVQPGGSL RLSCAASGFTFS	SYAMRW	VRQAPGKGLE WVS	AISGSGGSTY YADSVKGR	FTISRDDSKNTAYLQMNSLK TEDTAVYYC	SRVGLWGPRY YYYYGMDVW	IGHV3-23*04	4	VH3
YM1-D2	QMQLVQSGAEVKKTGSSM KVSCRVSGYTFT	NRYLHW	VRQAPGQALE WMG	WITPFNGNTN YAQKFQDR	VTITRDRSMSTAYLELSSLR SEDTAMYYCAS	GGPLDY	IGHV1-45*02	5	VH1
VL									
TomI-F6	DIQMTQSPSSLSASVGDR VTITC	RASQSISSYL NW	YQQKPGKAPK LLIY	ASSLQSGV	PSRFSGSGSGTDFTLTISSL QPEDFATYYC	QQGANYPNT	IGKV1D-39*01	2	VK1
TomI-A8	DIQMTQSPSSLSASVGDR VTITC	RASQSISSYL NW	YQQKPGKAPK LLIY	NASALQSGV	PSRFSGSGSGTDFTLTISSL QPEDFATYYC	QQNAYSPDT	IGKV1D-39*01	2	VK1
TomI-A11	DIQMTQSPSSLSASVGDR VTITC	RASQSISSYL NW	YQQKPGKAPK LLIY	ASNLQSGV	PSRFSGSGSGTDFTLTISSL QPEDFATYYC	QQNADSPAT	IGKV1D-39*01	2	VK1
YM1-C3	DTVMTQSPSSLSASVGDR VTITC	RASQSISSYL NW	YQQKPGKAPR LLIYA	ASSLQSGV	PSRFSGNGSGTDFTLTISSL QPEDFATYYC	QQSYSTPYA	IGKV1D-39*01	4	VK1
YM1-C5	SYVLTQPPSVSLTLGQTA RITCE	GNSIGRKHVH W	YQQKPGQAPV LVYR	DIQRPSGI	PERFSGSNSGNTATLTITGV QVGDDSDYYC	QVWDRDSRTI V	IGLV3-9*01	16	VL3
YM1-D2	QSALTQPASASGSPGQSV TISCT	GTGSDVGGYN YVSW	YQQHPGKAPK LMIYE	VSKRHSGV	PDRFSGSKSGNTASLTVSGL QAEDEADYYC	SSYAGSNNLV	IGLV2-8*01	3	VL2
4.6 Purification of soluble scFv antibodies

The pIT2 vector of the Tomlinson's library encodes a carboy-terminal hexahistidine tag, which allowed for rapid purification of the recombinant antibodies using IMAC. Clone TomI-F6 that showed to be inhibited by soluble aflatoxin was selected to purify. For small scale production, induction was performed at 30°C for 20 hours, using 1mM IPTG in a total culture volumn of 15 ml. The antibody fragment was purified from cell lysate by using Ni-NTA resin, and analyzed by SDS-PAGE and Western Blot (Figure 16). All factors affecting purification, including the quantity of resin and concentration of imidazole in both the wash and elution buffers. Imidazole were incorporated in the wash buffer, reduced the non-specific binding of contaminant proteins to the resin. scFv fragments were eluted with 250mM imidazole. The results indicated that the purity of antibodies greatly increased following elution with imidazole in comparison to the crude cell lysate.



Figure 16. SDS-PAGE (A) and western blot analysis (B) of purified scFv antibody in small scale production (anti-Aflatoxin clone TomI-F6) by IMAC. SDS-PAGE stained with CBB. Western blot probed with anti-protein L peroxidase labeled antibody. Color was developed by using TMB substrate. Lane (M): Protein Molecular weight Markers (Sigma Colour burst prestain); Lane (L): cell lysate fraction; Lane (F): Flow through fraction; Lane (W1-3): Wash fraction; Lane (E1-3): Elution fraction containing purified scFv antibody.

For the large scale production of soluble scFv antibody, the culture volumn were increased to 500 ml. For high levels of expression, antibody was grown in Terrific broth, which contains high concentrations of tryptone and yeast extract also includes glycerol as source of carbon and potassium phosphate salts as a buffering agent to prevent cell death at decreased pH. The results of SDS-PAGE and Western Blot analysis (Figure 17) showed that the scFv-hexahistidine fusion antibodies with a size of about 30 kDa were successfully purified from lysed cells using Ni-NTA resin. However, the contaminated protein appeared to co-purifying along with desired scFv antibodies. The contaminants removal approach will be necessary in further purification step. The western blot result showed that only monomer scFv was obtained and that it was functionally active. The quality of product seems to be lower than small scale production because sometimes inclusion body format can occur during production scFv, which results in poor expressivity (Brichta et al., 2005). This problem can be solved by cloning this scFv to pKP300 Δ III (expression vector) encode an carboxy-terminal (His)₆ tag, which allowed for the rapid purification of recombinant antibodies using immobilized metal ion affinity chromatography (IMAC).



Figure 17. SDS-PAGE (A) and Western blot analysis (B) of purified scFv antibody in largescale production (anti-Aflatoxin clone TomI-F6) by IMAC. SDS-PAGE stained with CBB. Western blot probed with anti-protein L peroxidase labeled antibody. Color was developed by using TMB substrate. Lane (M): Protein Molecular weight Markers (Sigma Colour burst prestain); Lane (L): cell lysate fraction; Lane (F): Flow through fraction; Lane (W): Wash fraction; Lane (E1-2): Elution fraction containing purified scFv antibody.

4.7 Cloning of scFv fragments into pKP300∆III vectors

The scFv fragments against aflatoxin B_1 that selected from compact human naïve library (Pansri et al., 2009) and a semi-synthetic library (Tomlinson; TomI-F6) as engineered to create scFv-alkaline phosphatase fusions (scFv-AP) and used as convenient one-step detection probes for competitive ELISA.

The resulting scFv fragment (TomI-F6) and pKP300 Δ III vector were digested with restriction enzymes *NCoI* and *Not I*. The digested products were assessed on agarose gel and the resulting gel images are shown in Figure 18 with bands visible at approximately 5,000 bp representing vector and 750-800 bp representing insert.



Figure 18. Agarose gel analysis of *NCoI* and *NotI* digestion products. M, 1 kb DNA ladder and 100 bp ladder. A vector (pKP300ΔIII) and insert (scFv-F6) band can be seen at approximately 5,000 bp and 800 bp, respectively. The concentration of DNA of vector and insert were 12.1 µg/ml and 12.3 µg/ml, respectively.

Full length scFvs had been prepared; they were ligated into pKP300ΔIII vector at *NCoI* and *NotI* sites. The ligation reactions were transformed to *E.coli* TG1 by electroporation. After incubation for 16 hour, the 5 individually colonies were randomly picked and grown in Low phos media at 30°C for 20 hour. The scFv-AP was expressed and secrete into culture media. Cell pellets were used to extracted plasmid by using Plasmis mini Prep (Qiagen, Germany). Plasmids were digested with *NCoI* and *Not I* for checking the scFv were clone into expression vector. The result showed that all of 5 clones were successfully cloned into vector (Figure 19).



Figure 19. Agarose gel analysis of NCoI and NotI digestion products to check positive clones. Lane 1, 1 kb DNA ladder; lane 2, 4, 6, 9, 11, undigested of clone 1-5; lane 3, 5, 7, 10, 12, clone 1-5 were digested with enzymes; lane 8, 100 bp DNA ladder. The digestion products were seen in two sites indicated by arrow.

4.8 ScFv-AP ELISA

The five scFv-AP clones were tested the specific to bind AFB₁-BSA by ELISA using PNPP as substrate. The yellow color was developed and measure absorbance at 405 nm. Clone 1 that showed the highest signal bound to AFB₁-BSA (Figure 20) was assessed to determine the optimal antibody dilution to the further step in competitive ELISA. The result showed that an antibody dilution of 1/10 (TomI-F6) and 1/20 (YM1-C3) were appropriated for applies in a competitive ELISA to obtain the half maximal inhibitory concentration (IC₅₀) of soluble AFB₁ and % cross-reactivity (Figure 21).



Figure 20. The binding scFv-F6-AP specific to AFB₁-BSA. PNPP was used to be substrate for AP. Absorbance signals were presented with standard deviation of triplicates.



Figure 21. ELISA for determination of scFv-AP antibody dilution for use in a competitive ELISA. Antibody dilutions from 1:2 to 1:256 were assayed. Absorbance signals were presented with standard deviation of triplicates.

4.9 scFv-AP competitive ELISA

Various 1:3 (B: YM1-C3) and 1:2 dilutions (A: TomI-F6) of soluble AFB₁ from 5.0 μ g/ml-0.762 ng/ml in 5% methanol in TBST were incubated with two scFv-AP (TomI-F6 and YM1-C3) at 37°C for 30 minutes before adding into wells of Immuno 96 MicroWellTM plates, which coated with 5 μ g AFB₁-BSA. The plated were washed five times with TBST after 1 hour of incubation. PNPP were used as substrate for AP. The IC₅₀ values for the TomI-F6-AP and YM1-C3-AP were found to be 0.2 μ g/ml and 0.035 μ g/ml, respectively. This result showed that the YM1-C3 demonstrated approximately 10 fold increase in assay sensitivity when compared to TomI-F6. ELISA inhibition curves for both antibodies are shown in Figure 22. The limit of detection (LOD) is 0.006-0.03 μ g/ml. This is the smallest concentration of the analyses that produces a signal. This value was determined by selecting

the mean normalized absorbance minus three standard deviations for the negative standard (no AFB₁). Moghaddam et al., and Yang et al. reported that the IC_{50} of the selected clones were about 400, 0.4 ng/ml, respectively. The previous study showed more sensitivity than this study. In the future study, the sensitivity of my antibodies can be further improved by various methods of affinity maturation.



Figure 22. Competitive ELISA of scFv-AP antibody for detection of different concentration of AFB₁. The average absorbance at a wavelength of 405 nm and S.D are shown. Data was normalized by dividing the mean absorbance obtained at each AFB₁ concentration (A) by the absorbance value determined in the presence of zero toxins (A₀). Normalized absorbance values (A/A₀) were plotted against the logarithm of AFB₁ concentration. All analyses were performed in triplicate. The limit of detection of TomI-F6 and YM1-C3 were found to be 0.003 and 0.02 µg/ml, respectively.

4.10 Comparison sensitivity of scFv fragment and scFv-AP

The results of competitive ELISA employing the scFv YM1-C3 antibody fragments were compared to scFv-AP YM1-C3 in terms of assay sensitivity. Standard of each potential, ranging from 2.3 ng/ml to 5,000 ng/ml were prepared in TBST containing 5% (v/v) methanol and mixed with an equal volume of antibody at the optimal concentration. The IC₅₀ values for the scFv YM1-C3 and YM1-C3-AP were found to be 0.13 μ g/ml and 0.035 μ g/ml, respectively. The scFv-AP fusions showed improved binding affinity of approximately 3-20 folds better than the soluble form and phage-displayed scFv formats, respectively (Pansri et al., 2009).



Figure 23. Competitive ELISA for the detection of AFB₁ using the scFv antibody fragment and scFv-AP fusion antibody. The average absorbance at a wavelength of 405 nm and S.D are shown. Data was normalized by dividing the mean absorbance obtained at each AFB₁ concentration (A) by the absorbance value determined in the presence of zero toxins (A₀). Normalized absorbance values (A/A₀) were plotted against the logarithm of AFB₁ concentration. All analyses were performed in triplicate. The IC₅₀ values for the scFv YM1-C3 and YM1-C3-AP were found to be 0.13 µg/ml and 0.035 µg/ml.

4.11 Cross-reactivity

The anti-aflatoxin antibodies were demonstrated for cross-reactivity with three structurally related aflatoxins; aflatoxin B₂, G₁, G₂ and M₁ using competitive ELISA. Standard of each potential cross-reactant, ranging from 2.3 ng/ml to 5,000 ng/ml were prepared in TBST containing 5% (v/v) methanol and mixed with an equal volume of antibody at the optimal concentration. The cross-reactivity potential was approximated at the IC₅₀. IC₅₀ value is the analyze concentration that results in 50% inhibition. The percent inhibition was determined according to the equation: % inhibition = [(OD sample/ OD control)] × 100 (Figure 24). The percentage cross-reactivity determined at IC₅₀ (%CR₅₀) was expressed as 50% inhibitory concentration of aflatoxin B₁ divided by the 50% inhibitory concentration of other aflatoxin and multiplied by 100.



Figure 24. Percent inhibition of aflatoxin antibody (A) TomI-F6 and (B) YM1-C3 was calculated from ELISA value of competitive ELISA. Results were plotted as the mean percent inhibition against the logarithm of soluble aflatoxins concentrations. Data were presented with standard deviation of triplicates.



Figure 24. (Continued)

Table 7 shows the results of the studies, which the scFv-AP antibody showed high degree of cross-reactivity, especially to $AFG_1 > AFB_2 > AFG_2 > M_1$ (in decreasing order). Anti-aflatoxin scFv-AP TomI-F6 and YM1-C3 showed the highest level cross-reactivity with aflatoxin G₁ were 68.75% and 70%, respectively. Moghaddam (2001) found that their antibody was able to bind as effectively to AFG₁ as AFB₁. Yang (2009) reported their antibody showed cross-reactivity of 12% with AFB₂, 42% with AFG₁ and 9% with AFG₂. This is similarity in this study. The YM1-C3 showed low percentage cross-reactivity with aflatoxin M_1 (0.88%). This can be shown by the fact that hydroxylation C9 residue in aflatoxin M₁ results in a reduction in binding affinity.

Table 7. Percentage cross-reactivity value of the TomI-F6 and YM1-C3 scFv-AP antibody to various potential cross-reactants. The cross-reactivity potential was proximated at the IC₅₀ value, which was estimated at 50% A/A₀.

	TomI-F6		YM1-C3	
Aflatoxin compound	IC ₅₀ (ug/ml)	CR ₅₀ (%)	IC ₅₀ (ug/ml)	CR ₅₀ (%)
B_1	0.22	100	0.035	100
B_2	1.3	16.92	0.13	26.92
G_1	0.32	68.75	0.05	70
G_2	1.6	13.75	0.12	29.17
M_1	1.7	12.94	4	0.88

4.12 Purification of scFv-AP

The pKP300ΔIII vector expresses scFv with FLAG tag at N terminal and enzyme alkaline phosphatase and hexahistidine tag fusion at C terminal, which allowed for rapid purification of the recombinant antibodies using IMAC. Clone scFv-AP TomI-F6 was cultured in a total culture volumn of 10 ml. Induction was performed at 30°C for 20 hours. The scFv-AP fusion was purified from cell lysate by using Ni-NTA resin, and analyzed by SDS-PAGE (Figure 25). The result showed that the scFv-AP fusion antibodies with a size of about approximately 70 kDa were successfully purified from lysed cells using Ni-NTA resin.



Figure 25. SDS-PAGE purified scFv-AP (TomI-F6) antibody. SDS-PAGE stained with CBB. Lane (M): Protein Molecular weight Markers (Fermentous unstain); Lane (L): cell lysate fraction; Lane (FT): Flow through fraction; Lane (W1-3): Wash fraction; Lane (E1-3): Elution fraction containing purified scFv-AP antibody.

4.13 ScFv-AP binds to AFB₁ in peanut samples

To assess the assay method in practice, peanuts were taken from farmers' fields that were presumed to be at high risk of aflatoxin contamination. These were extracted and assayed by using competitive ELISA. The peanut extract was diluted at 0, 1:16 and 1:256 and was incubated with scFv-AP TomI-F6 before adding to ELISA coated well with AFB₁-BSA. The scFv-AP can bind to AFB₁-BSA (high signal), when the dilution of peanut extract was increased (the concentration of AFB₁ contaminated decreased). The result (Figure 26) showed that this scFv-AP can be used to detected aflatoxin in agricultural product.



Figure 26. Inhibition of peanuts extract by competitive ELISA. Soluble scFv-AP was incubated with (A) 5% methanol in TBST, (B) 1:256 dilutions of peanut extract, (C) 1:16 dilutions of peanut extract and (D) undiluted peanut extract before transferring to AFB₁-BSA that coated microtitre wells. Absorbance signals were presented with standard deviation of duplicates.

CHAPTER V

CONCLUSION

In summary, scFv antibody specific to aflatoxin B₁ can be selected from naïve semisynthetic scFv phage display antibody libraries by standard bio-panning. Polyclonal phages showed positive signal in each round of panning. The positive clones from 1st and 3rd round of selection were determined for specific binding to aflatoxin B_1 by competitive ELISA. Three clones from eight clones, that produced high affinity antibody specific to aflatoxin B_{1} . were selected. All selected clones showed the same nucleotide sequence after automated DNA sequence analysis. DNA sequence of variable regions using IgBLAST and VBASE program analysis revealed that the antibody contained IGHV3 of human V_H region and IGKV1 of human VL region. One of soluble scFv antibody clone that shown high specific to aflatoxin B_1 was expressed and purified by using $Ni^{2+}\mbox{-}NTA$ affinity column chromatography. This scFv antibody and a compact human naïve phage display library (YAMO1) were engineered to create scFv-alkaline phosphatase fusion (scFv-AP) and can be used as convenient one-step detection probe for competitive ELISA. The IC₅₀ of scFv-AP by AFB₁ varied between 0.035-0.02 μ g/ml and the limit of detection was approximately $0.006-0.03 \mu g/ml$ (6-30 ppb). The selected antibodies were specific to AFB₁ and could cross-react with AFG₁, but not to other aflatoxins. These antibodies can be used in therapeutic study in the future.

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antibody to the mycotoxin zearalenone. Applied and Environmental Microbiology 63, 263.

BIOGRAPHY

Miss Kuntalee Rangnoi was born on November 14, 1983 in Phetchaburi, Thailand. She graduated with the Bachelor Degree of Animal Production Technology, Suranaree University of Technology, Thailand in 2005. She had opportunity to study Master Degree enrollment in School of Biotechnology, Institute of Agricultural Technology, Suranaree University of Technology (2005-2009). During her study, she has experience on her research for 6 months at Applied Biochemistry Group, school of Biotechnology and Biomedical Diagnostics Institute, Dublin City University, Dublin, Ireland. She had presented research work in Agricultural Biotechnology for Better Living and a Clean Environment (ABIG 2009), September 22-25, 2009, Queen Sirikit National Convention Center, Bangkok, Thailand (Poster presentation; in "Selection of scFv antibody fragment specific to Aflatoxin by using phage display technology"). She conducted to research in the topic of selection of monoclonal antibodies specific to aflatoxin from phage display antibody library.