องค์ประกอบทางเคมีของผลพิลังกาสา Ardisia colorata Roxb. และฤทธิ์ทางชีวภาพ

นายพิชยา ประเสริฐแสง

สถาบนวิทยบริการ

วิทยานิพนธ์นี้เป็นส่วนหนึ่งของการศึกษาตามหลักสูตรปริญญาวิทยาศาสตรมหาบัณฑิต สาขาวิชาเคมี ภาควิชาเคมี คณะวิทยาศาสตร์ จุฬาลงกรณ์มหาวิทยาลัย ปีการศึกษา 2545 ISBN 974-17-1965-5 ลิขสิทธิ์ของจุฬาลงกรณ์มหาวิทยาลัย CHEMICAL CONSTITUENTS OF Ardisia colorata Roxb. FRUIT AND THEIR BIOLOGICAL ACTIVITIES

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สถาบนวทยบรการ

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พิชยา ประเสริฐแสง : องค์ประกอบทางเคมีของผลพิลังกาสา *Ardisia colorata* Roxb. และฤทธิ์ ทางชีวภาพ (Chemical constituents of *Ardisia colorata* Roxb. fruit and their biological activities) อ. ที่ปรึกษา: ผศ. คร.วรินทร ชวศิริ, 110 หน้า. ISBN 974-17-1965-5.

ผลการสึกษาฤทธิ์ทางชีวภาพเบื้องค้นชี้ให้เห็นว่าสิ่งสกัดไดลออโรมีเทนของผลพิลังกาสา แสดงกวามเป็นพิษต่อไรสีน้ำตาล Artemia salina ในระดับสูง และแสดงฤทธิ์ด้านอนุมูลอิสระ DPPH จึงสึกษาองก์ประกอบทางเกมีของสิ่งสกัดดังกล่าว พบสารบริสุทธิ์ 3 ชนิด ได้แก่ 2,5dihydroxy-3-undecyl-1,4-benzoquinone (embelin), stigmasterol, octadeca-9,12-dienoic acid (linoleic acid) และของผสม 4 ชนิด ได้แก่ ของผสมแอลกอฮอล์โซ่ตรง, ของผสมของ stigmasteryl-3-O-palmitate และ β-sitosteryl-3-O-palmitate, ของผสมซึ่งมี α-amyrin เป็นส่วน ประกอบ และของผสมของ 1,3-dihydroxy-5-(heptadec-8,11-dienyl)benzene และ 1,3dihydroxy-5-(pentadec-8-enyl)benzene พบว่า embelin ซึ่งเป็นองก์ประกอบหลัก แสดงกาวม เป็นพิษต่อไรสีน้ำตาลและแสดงฤทธิ์ด้านอนุมูลอิสระ DPPH สูงสุด ในขณะที่แสดงฤทธิ์ด้านการ กินของแมลงในระดับปานกลาง แสดงฤทธิ์ยับยั้งจุลชีพ Bacillus subtilis และฤทธิ์การยับยั้ง เอนไซม์ p-hydroxypyruvate dioxygenase พบว่าของผสมของ 1,3-dihydroxy-5-(heptadec-8,11-dienyl)benzene และ 1,3-dihydroxy-5-(pentadec-8-enyl)benzene แสดงฤทธิ์ยับยั้งการ เจริญของเชื้อราโรกพืช Alternaria sp. โดยอาศัยวิธี bioautographic และแสดงฤทธิ์ฆ่าหนอน กระทู้ผัก Spodoptera litura สารทั้งสองชนิดแสดงฤทธิ์ยับยั้งเซลล์มะเร็งเด้านมและเซลล์มะเร็ง ปอดได้ในระดับสูง

ภาควิชาเคมี	ลายมือชื่อนิสิต
สาขาวิชาเคมี	ลายมือชื่ออาจารย์ที่ปรึกษา
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The preliminarily biological activity results pointed out that the dichloromethane crude extract of Ardisia colorata Roxb. fruits exhibited high cytotoxic activity against brine shrimp Artemia salina as well as scavenged the DPPH radical. The chemical constituents investigation disclosed three pure compounds: 2,5-dihydroxy-3-undecyl-1,4-benzoquinone (embelin), stigmasterol and octadeca-9,12-dieonoic acid (linoleic acid), and four mixtures: a mixture of long chain alcohols, a mixture of stigmasteryl-3-Opalmitate and β -sitosteryl-3-O-palmitate, a mixture containing α -amyrin, and a mixture of 1,3-dihydroxy-5-(heptadec-8,11-dienyl)benzene and 1,3-dihydroxy-5-(pentadec-8enyl)benzene. Among isolated substances, embelin, the major component, exhibited the highest cytotoxic activity against brine shrimp, the highest scavenging effect on DPPH radical, moderate insect antifeedant activity, and gave positive tests for antimicrobial activity against Bacillus subtilis, enzymatic inhibitor on p-hydroxypyruvate dioxygenase. A mixture of 1,3-dihydroxy-5-(heptadec-8,11-dienyl)benzene and 1,3-dihydroxy-5-(pentadec-8-enyl)benzene displayed the growth inhibition effect on phytophatogenic fungi, Alternaria sp., by bioautographic method as well as showed insecticidal activity against Spodoptera litura. These two substances displayed strongly cytotoxic activity against the breast cancer cell and the small cell lung cancer.

จุฬาลงกรณ์มหาวิทยาลัย

DepartmentChemistry	Student's signature
Field of studyChemistry	Advisor's signature
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BHA butyrated hydroxyanisole °C degree of celcius dichloromethane, methylene chloride CH_2Cl_2 CHCl₃ chloroform cm^{-1} unit of wavelength d doublet (NMR) DPPH 2,2-diphenyl-1-picryhydracyl ED₅₀ 50% effective dose EtOAc ethyl acetate DMSO dimethylsulfoxide gram (s) g gas chromatography GC J coupling constant 50% inhibitory concentration IC₅₀ kilogram (s) kg weight wt NMR nuclear magnetic resonance IR infrared L liter (s) 50% lethality concentration LC_{50} LD_{50} 50% lethality dose multiplet (NMR) m milligram (s) mg mL milliliter (s) melting point m.p. mass to charge ratio m/z MW molecular weight M^+ molecular ion

List of Abbreviations

broad

br

nm nanometer

ppm part per million

s singlet (NMR)

List of abbreviations (continued)

t	triplet (NMR)
TLC	thin layer chromatography
δ	chemical shift
μg	microgram (s)
λ	wavelenght
MeOH	methanol
MS	mass spectrometry
R _f	retardation factor



CHAPTER I

INTRODUCTION

Thailand is a tropical country located in suitable landscape. Therefore, a variety of plants species is widely spread. A lot of species exhibit significance for medically useful drugs. In early age, the use of medicinal plants for medical treatments was not complicated and mainly depended on ancient belief. They were normally used in both a single form and multiple forms, which have been existed by the passage of knowledge down the generations within each ethic group.

Although several plant species have been distributed in Thailand, only small proportion has been thoroughly phytochemical and pharmacological investigation. When one considers that a single plant may contain up to thousands of constituents, the possibilities of making new discoveries become evident. Fransworth claims that 119 characterized drugs are still obtained commercially from higher plants and that of 74% were found from ethnobotanical information.¹

At the present time, the research and development of new drugs for treatments are still main problems of medicinal development. In this research, the searching for biologically active compounds from a medicinal plant, *Ardisia colorata* Roxb., has been focused.

About half of useful drugs nowadays are derived from natural sources. Therefore, the natural products have served as an important source of drugs since ancient times. Rapid identification of bioactive compounds from natural product mixtures still remained as a critical factor since the development procedures for searching biologically active compounds from natural source such as plants, microorganisms, marine organisms *etc.* required a long and tedious process. For this reason, it is necessary to be of methods available which eliminate unnecessary separation procedures. Chemical screening is thus performed to allow localization and targeted isolation of new or useful types of constituents with potential activities.

Screening program for biologically active natural products requires the right bioassays. Bioassays are also essential for detecting the required effects throughout activity-guided fractionation. They must be simple, inexpensive and rapid in order to cover with large number of samples.

Since most active plant principles are toxic at elevated doses, a possible approach to develop an effective general bioassay might be simply to screen for substances that are toxic to zoologic systems. Desiring a rapid, inexpensive, in-house bioassay for screening and fractionation monitoring of physiologically active plant extracts. Brine shrimp lethality cytotoxic test has been used as a bioassay for a variety of toxic substances. This method has also been applied to plant extracts in order to facilitate the isolation of biologically active compounds.

Brine shrimp *Artemia salina* Linn. as shown below is a tiny crustacean. It can response to biologically active compound comparable with mammalian system. It has also DNA-dependent RNA polymerase enzyme and ouabaine sensitive Na⁺ and K⁺ dependent ATPase like in mammalian system.²

Therefore, the results from brine shrimp assay can be used as a primary toxic representative for other mammal species.



Brine shrimp

The searching for cytotoxic compound such as anticancer, antimicrobial, insecticide *etc*. has been frequently used brine shrimp as general tools by microwell method.^{3,4} The eggs of brine shrimp are readily available at low cost in pet shops as food for tropical fish, and they remain viable for years in the dry state. Upon being placed in as brine solution, the eggs hatch within 24 hours, providing large number of larvae (nauplii). This method is convenient since it requires a little material, rapid,

inexpensive and reliable. For this reason, this brine shrimp assay is selected as one of primary screening methods of this study.

The second screening method involving to this research is an antioxidant activity. Oxidative processes are well known to be major causes of degradation of food and materials. More recently, free radicals in particular oxygen reactive species, have been recognized to be involved in many diseases, including cancer and arteriosclecrosis. Aging also may be the sum of the free radical reactions which occur continuously throughout cells and tissues. Plant contains several widespread phenolic compounds with well established antioxidant activity. Typical examples are common flavonoids such as quercetin or rutin, phenolic acid such as gallic or caffeic acids and tannins. The structures of each compound are shown below.



The screening strategy for searching new antioxidant compounds from crude plant extract based on simple and rapid TLC autographic assays. In 1994, Takao *et al.*⁵ used 2,2-diphenyl-1-picryhydrazyl (DPPH) radical for screening antioxidants in marine bacteria as a TLC spray reagent. It was also proved to be well suited for the detection of antioxidants in crude plant extracts.

Scavenging effect towards DPPH radical consisted of two assays for qualitative and quantitative analysis.⁶ The assay for qualitative analysis using TLC autographic method, and spectrophotometric assay for quantitative analysis respectively. Both of them use DPPH radical as a radical source. The different color between radical form (purple) and non-radical form (colorless) of DPPH was used for regarding. The structures of DPPH radical and non-radical forms are illustrated as shown below.



From primarily biological screening of Thai medicinal plants using two screening methods described above, under co-operation between Department of Chemistry and Department of Biology, Faculty of Science, Chulalongkorn University, it was found that the dichloromethane extract of dried fruits of *Ardisia colorata* Roxb. (Pi-Lang-Ka-Sa) belonging to Myrsinaceae family showed attractive results for both screening methods.

1.1 Botanical Aspects and Distribution

Ardisia colorata Roxb. Shrub or small tree to 5-10 m. The bark is brown, quite smooth and very thin. The oblong-lanceolate with point or slightly tapering tip and blunt or slightly pointed base leaf about 13-28 cm. long and 3-8 cm. wide. These plants have no teeth or gland on margin. Mature leaves without hairs but with scattered tiny rusty-brown scale and dark gland dots below. It has 15-20 pairs of side veins with many shorter intermediate ones. Stalks 0.6-1.5 cm the young twigs pale cream, densely scaly. The flower is pale pink about 0.25-0.3 cm in branched pyramidal clusters (panicles) at end of twigs, 10-18 cm. Individual stalks 0.2-0.4 cm \pm scaly. Calyx \pm 0.1 cm spilt 2/3 into spreading lobes, not overlapping, black-dotted. The corolla deeply spilt with blunt lobes, no gland dots. Slender and projecting before petals open. The fresh fruit is pale yellow, when old is black.⁷ The pictures of flower, leaf and fruit of *Ardisia colorata* Roxb. are displayed below.



According to Thai herbal ancient believes, the leaf was used to treat for the liver diseases. The fruits showed antidiarrheal and antipyratic activity and the roots were used for treatment of gonorrheal diseases. The isolated mixtures from fruits can inhibit the growth of *Salmanella* spp. and *Shigella* spp. which cause of diarrhea diseases.⁸

In Thailand, the plants belonging to Myrsinaceae family are consisted of 6 species⁷ i.e. *Ardisia colorata* Roxb. (Pi-Lang-Ka-Sa), *Ardisia nervosa* Kurz. (Gee-Par-Taek), *Ardisia viren* Kurz.(Tar-Nok), *Rapanea yunnanensis* Mez. (Lang-Ka-Tae), *Maesa ramentacea* A.Dc. (Kra-Duk-Kai) and *Maesa montana* A.Dc. (Hat-Sa-Kun-Kreua). In this research, the fruits of *Ardisia colorata* Roxb. were selected to examine for their chemical constituents.

1.2 Literature Reviews for Ardisia colorata Roxb.

In 1986, O. Luanratana *et al.* reported the orange pigment, rapanone which was obtained from petroleum ether, chloroform and ethanol extracts of the barks of *A.colorata* Roxb. It exhibited antileprotic activity on mice which infected with *Mycobacterium leprae*.⁹



Furthermore, O. Luanratana *et al.* addressed the microbial growth inhibition results of the ethanolic extract from the barks. The ethyl acetate fraction exhibited good activity to inhibit the growth of *Aeromonas hydrophila*, *Vibrio cholerae* and *Shigella dysenteriae*, whereas the butanolic fraction revealed antigonorrhoeal activity against *Neisseria gonorrhoea*.¹⁰

1.3 Chemical Constituents Studies on Ardisia Genus

Literature surveys of chemical constituents of the plants belonging to *Ardisia* genus are presented in Table 1.1. The structures of some isolated compounds are shown below.

Scientific Name	Compound	Ref.
Ardisia japonica Thunb.	Norbergenin (1), Bergenin (2)	11
	and tri-O-methylnorbergenin (3)	
	Triterpenoid saponin (4),	
	Triterpenoid glycoside,	
Ardisia iwahigensis Elmer.	Ardisenone (5)	12
Ardisia cornudentata Mez.	Ardisianone (6),	13
	Cornudentanone (7)	
Ardisia solanacea Roxb.	Bauerenol acetate (8)	14
2	α -Amyrin and β -Amyrin	
Ardisia mamillata Hance.	Ardisimamillosides A-D (9-12)	15
Ardisia crenata	Ardisicrenoside A-F (13-18)	16
Ardisia sieboldii	Ardisiaquinone A-F (19-24)	17

Table 1.1Compounds found in Ardisia genus









1.4 The Goal of This Research

The attractive preliminary results for bioassay of *Ardisia colorata* Roxb. crude extract call for intensive investigation. Therefore, the goal of this research can be summarized as:

- To extract and isolate the chemical constituents from the dried fruit of A. colorata Roxb.
- 2. To elucidate the structures of isolated compounds.
- 3. To explore the biological activities of isolated compounds.



CHAPTER II

EXPERIMENTAL

2.1 Plant Materials

The dried fruits of *Ardisia colorata* Roxb. (Pi-Lang-Ka-Sa) were purchased from Vetchapong, Bangkok, Thailand, in June 2001. The plant sample has been compared with voucher specimen deposited in the Herbarium of the Princess Sirindhon, Thailand (BK 14133).

2.2 Instruments and Equipment

Melting points were determined with a Fisher-Johns melting point apparatus and are uncorrected. Thin layer chromatography (TLC) was performed on an aluminium sheet precoated with silica gel (Merck's Kieselgel 60 PF₂₅₄). Column chromatography was performed on silica gel (Merck's Kieselgel 60G) and flash chromatography was proceeded on silica gel (40 μ m average particle diameter). Chromatotron was performed on Harrison research, Model 7924 T, Serial No. W34 Patented, made in U.S.A.

The Fourier Transform-Infrared Spectra (FT-IR) were recorded on NICOLET IMPACT 410 FT-IR spectrometer. The ¹H and ¹³C NMR spectra (both 1D and 2D) were performed in deuterated chloroform or dimethylsulfoxide with tetramethylsilane as an internal reference on Bruker Fourier Transform Nuclear Magnetic Resonance Spectrometer, model AC-F200 and a Joel, model JNM-A500 and a Bruker Avance 300 FT-NMR spectrometer. Mass spectrometry (MS) analysis was conducted on Fisson Instrument Model Trio 2000. GC analysis was carried out on a Shimadzu GC GC-9A instrument equipped with flame ionization detector with N₂ as a carrier gas. The GC-MS analysis was performed on GC model star 3400Cx and MS model saturn 4D from Varian.

2.3 Chemical Reagents

All solvents used in this research were purified by standard methodology except for those which were reagent grades. The reagents utilized for synthesizing all derivatives were purchased from Fluka Chemical Company or otherwise stated and were used without further purification.

2.4 Chemical Tests

- Liebermann-Berchard's Test

The sample was dissolved in dry chloroform 0.5 mL, then slowly added 2-3 drops of acetic anhydride followed by one drop of concentrated sulfuric acid. Development of the color suggests the presence of steroids or triterpenoids. If the solution was dark blue or greenish blue, the sample may contain steroidal nucleus, whereas if the solution turned reddish, the sample should be triterpenoidal compound.

2.5 Extraction of Ardisia colorata Roxb.

The dried and powdered fruits of *A. colorata* 5 kg were extracted by soaking with dichloromethane for two days at room temperature. The same process was repeated twice. During the evaporation of dichloromethane from the extract, the orange plate was precipitated. This orange solid was recrystallized from methanol twice to give Compound <u>1</u>. The remained portion was completely evaporated to furnish a dark brownish crude as Fraction I. The marc was similarly extracted with ethyl acetate and methanol to yield Fractions II and III as dark brownish crudes. The extraction process is summarized as shown in Scheme 2.1.



Scheme 2.1 Extraction procedure for the dried fruits of *A. colorata* Roxb.

2.6 Chemical Reactions

2.6.1 Hydrolysis of Mixture <u>3</u>¹⁸

Mixture $\underline{3}\ 2.0$ g was dissolved with 10% ethanolic KOH 40 mL in the 100 mL round bottom flask. The solution was stirred under refluxing on a water bath for 4 hours. The reaction was monitored until the reaction was completed by thin layer chromatography compared with starting material. After the reaction was completed, the solvent was evaporated and the residue was extracted with diethyl ether 150 mL for three times. The combined diethyl ether layer was dried over anhydrous sodium sulfate. Evaporation the solvent furnished a hydrolyzed product, assigned as Mixture $\underline{3A}$.

The remained aqueous layer after extraction with diethyl ether was acidified with dilute hydrochloric acid to yield the acid part designated as Mixture <u>3B</u> which was further analyzed as methyl ester.

2.6.2 Methylation of Mixture <u>3B</u>

Mixture <u>3B</u> was dissolved in a small amount of $CHCl_3$ and ethyl acetate. Then, two drops of trimethylsilyl diazomethane (TMSCHN₂) were added under nitrogen gas flow condition to give the permanent yellow solution. The solution was left at room temperature overnight. After the reaction was completed, the product was further analyzed by GC-MS analysis.

2.7 Bioassay Experiments

2.7.1 Brine Shrimp Cytotoxic Lethality Test⁴

This bioassay is a procedure for general toxicity screening. The advantage of this assay is rapid, reliable, convenient and inexpensive. It requires small quantities of material and is able to identify a broad spectrum of activities. Thus, it is essential as a preliminary testing for the search of bioactive compounds. There are several techniques for this assay. A microwell method has been used for this experiment and was described as follows:

2.7.1.1 Sample Preparation

The samples were prepared by dissolving 4 mg of tested compound (either crude extract or pure compound) in 80 μ L of dimethyl sulfoxide (DMSO) then added artificial sea water (dissolving 38.5 g of NaCl in 1 L of distilled water and filtered by cellophane paper) to the solution to make 4000 μ L and allowed to shake-well to afford solution A (1000 ppm). Serial dilution of this stock solution was made to obtain solution B (100 ppm) and solution C (10 ppm), respectively. The control solution was prepared by using only DMSO and artificial sea water.

2.7.1.2 Hatching Brine Shrimp

Brine shrimp eggs *Artemia salina* obtained locally were hatched in artificial sea water in gloomy plastic box. The box was divided into unequal two sections linked with four 2 mm diameter holes. The eggs were scattering into the larger section which was darkened with aluminum foil while the smaller section was

illuminated with the 20 watt lamp. The box was kept at 22-29 °C. After 24 hours, nauplii were collected by micropipette from the smaller section.

2.7.1.3 Bioassay

Transferred five nauplii in 100 μ L of artificial sea water into each well of 24-well microplates by micropipette. Each concentration was performed for six replications. The covered plates were kept under the same conditions as hatching. The number of dead nauplii in each well was counted under binocular microscope after at 6 hours (for acute toxicity) and 24 hours (for chronic toxicity).

2.7.1.4 LC₅₀ Determination

LC₅₀ values were calculated by probit analysis program.¹⁹

2.7.2 Scavenging Effects on DPPH Radicals⁶

2,2-Diphenyl-1-picryhydrazyl (DPPH) radical is a stable radical with a purple color (λ_{max} 517 nm). Upon reduction by a scavenger, the extensive conjugation is disrupted and the compound turns yellow.

2.7.2.1 TLC Autographic Assay

After developing and drying, TLC plates were sprayed with a 0.2% DPPH in methanolic solution. The plates were examined 5 minutes after spraying. Active compounds appeared as yellow spots against purple background.

2.7.2.2 Spectrophotometric Assay

Samples of various concentrations (0.5 mL) were added to a 1 mL methanolic solution of DPPH radical (final concentration of DPPH was 0.2 mM). The mixture was shaken vigorously and then left for 30 minutes. The absorbance of the resulting solution was measured at 517 nm with a spectrophotometer. All tests and analyses were run in three replicates and averaged. Calculate the percentage of radical scavenging by the following equation.

The percentage of radical scavenging = $(1 - A_{sample}/A_{control}) \times 100$

 $A_{sample} = Absorbance of sample solution with DPPH$ $A_{control} = Absorbance of only DPPH and used solvent$

2.7.3 Antifungal Activity by Bioautographic Method²⁰

Direct bioautographic detection of the TLC plate is applicable to microorganisms that can grow directly on the TLC plate and suitable precautions are required. Each isolated substance was spotted on silica gel TLC plate and developed in suitable solvent system. This plate was allowed to air-dry for complete removal of solvents. The TLC plate was sprayed with a spore suspension of *Fusarium oxysporum* and *Alternaria* sp., and incubated at room temperature for 3 days. After incubation, the TLC was stained with 1% (V/V) lacto-phenol in 5% (V/V) acetic acid for 5 minutes and then destained with 5% (V/V) acetic acid for 10 minutes. The active components appeared as clear zone against blue background.

2.7.4 Insecticidal Activity by Vial Test Contact Toxicity²¹

This assay is dry film method or vial test to provide the potential substance on insecticidal activity, so it is a chance of finding substance for the phytophagous control. A polyphagous insect, the common cutworm, *Spodoptera litura* was used as a model for meanwhile investigation.

2.7.4.1 Sample preparation

The samples were prepared by dissolving 4 mg of test substances in 4 mL of acetone to provide 1000 ppm solution. Serial dilution of this stock solution was made to obtain 500, 100, 50, and 10 ppm, respectively. Then 1 mL of each concentration solution was poured into the glass vial (3 replication). Following, each vial that contained 1 mL of sample was evaporated the acetone, and the treated vial was placed in an open space for a few minutes to ensure complete removal of acetone. The test substances have already coated on the wall and bottom of vial.

2.7.4.2 Bioassay

Common cutworms *Spodoptera litura* were reared on an artificial diet in a controlled environment. The 15 of first instar larvae were placed in the vial to free movement for 6 hrs, then the larvae were transferred into the new vial containing artificial diet and were kept at 25°C for 5 days.

After the fifth day, the died cutworms were counted and converted to percentage of died larvae of *S.litura*. Finally, the LD_{50} in ppm was calculated by probit analysis program.

CHAPTER III

RESULTS AND DISCUSSION

Stemmed from the preliminary results that the dichloromethane extract from *Ardisia colorata* Roxb. fruit exhibited high activity for brine shrimp cytotoxicity as well as showed positive results for radical scavenger properties toward DPPH, this extract was rationalized to further investigate for chemical constituents and their biological activities.

3.1 Results of Extraction General Extraction

The crush dried fruits of *Ardisia colorata* Roxb. were extracted according to the procedure described in Chapter II. Each crude extract was subjected to preliminarily biological screening test followed the procedure described in Chapter II. The results of extraction are exhibited as shown in Scheme 3.1. Weight of each crude extract and percentage yield (w/w of dried plant material) are summarized as shown in Table 3.1.

From the extraction result (Scheme 3.1 and Table 3.1), it was signified that the orange plate solid (Compound <u>1</u>) 9.33 g or 0.19 % (w/w of dried fruit material) was deposited form CH_2Cl_2 extract. The extraction procedure was proceeded by increasing solvent polarity. Dichloromethane extract gave the best yield 114.20 g or 2.28 %(w/w). All crude extracts including Compound <u>1</u> were further screened for the biological activity using cytotoxicity test against brine shrimp and radical scavenging effect on DPPH radical.



Scheme 3.1 The results of extraction of dried fruits of *Ardisia colorata* Roxb.

Table 3.1Weight and percentage yield of each crude extract from the extraction
procedure of dried fruits of *Ardisia colorata* Roxb.

Crude extract	Weight (g)	Percentage yield*
CH ₂ Cl ₂ (Fraction I)	114.20	2.28
EtOAc (Fraction II)	75.40	1.51
MeOH (Fraction III)	95.60	1.91
Orange plate (Compound <u>1</u>)	9.33	0.19

* The percentage yield was calculated based on dried fruit materials (5 kg)

3.2 Preliminary Study on Biological Activity Test

3.2.1 Brine Shrimp Cytotoxic Lethality Test

This bioassay was performed following the methodology described in Chapter II. For preliminary study, each crude extract as referred in Scheme 3.1 was subjected for this assay. The results are displayed in Figure 3.1.



Figure 3.1LC50 values from brine shrimp cytotoxic lethality test for crude
extracts of Ardisia colorata Roxb.

Among crude extracts, the dichloromethane extract displayed high toxicity level with LC_{50} value of 13 ppm. Other crude extracts showed moderate toxicity against brine shrimp. This obtained result implied that the dichloromethane extract should contain substances responsible for cytotoxic activity.

3.2.2 Scavenging Effect on DPPH Radical TLC Autographic Assay

Antioxidant assay was accomplished by the protocol described in Chapter II. The chromatograms of all crude extracts before and after spraying with DPPH radical reagent are demonstrated in Figure 3.2.



Figure 3.2TLC autographic assay for DPPH radical scavenger assay

(A) TLC chromatogram before spraying with DPPH reagent

(B) TLC chromatogram after spraying with DPPH reagent

D : dichloromethane, E : ethyl acetate and M : methanol extracts

Figure 3.2A shows the TLC of all crude extracts which was developed in 30% ethyl acetate: hexane solvent system before spraying with DPPH. After sprayed with DPPH reagent, the active components were visualized as yellow spot against purple background (Figure 3.2B). It was found that the dichloromethane crude extract revealed at least 3 components active towards DPPH radical reagent.

From these preliminary results for biological activities of both brine shrimp cytotoxic lethality test and TLC autographic assay for DPPH radical scavenger, the most tendency potent extract was dichloromethane. The other crude extracts gave medium activity. Therefore, the dichloromethane extract was selected for further examination.
3.3 Chemical Constituents of Dichloromethane Extract (Fraction I)

During the evaporation of solvent from dichloromathane extract, it was noticed that some orange plate deposited. These solids were filtered and designated as Compound <u>1</u>. The remaining dichloromethane extract was further concentrated to furnish sticky dark brown material (Fraction I), which was further examined its constituents by separation with column chromatography.

3.3.1 Structural Elucidation of Compound <u>1</u>

The orange plate solid deposited upon the solvent evaporation of the dichloromethane extract was recrystallized with methanol twice to afford Compound <u>1</u> as orange plate crystal, 9.33 g (0.19% w/w of dried fruit material), m.p. 140-142 °C. This compound was soluble in dichloromethane and chloroform.

The IR spectrum of this compound (Figure 3.3) gave the absorption peaks of hydroxyl group at 3308 cm⁻¹, carbonyl group at 1613 cm⁻¹ and C-O bond at 1332 cm⁻¹. Other signals were tentatively assigned as shown in Table 3.2.

The mass spectrum (Figure 3.4) showed the molecular ion peak, M^+ , at m/z 294 (Calcd. for $C_{17}H_{26}O_4$: MW 294.18) and other important fragmentation ion peaks at m/z (% relative intensity) 155(40), 154(100), 153(28), 142(25) and 125(20). According to literature search, the fragmentation pattern of this compound²² was proposed as shown in Scheme 3.2.

The ¹H NMR spectrum (CDCl₃) of this compound (Figure 3.5) displayed the proton signals as follows: protons of hydroxyl group (2-OH, 5-OH) at δ 7.67 ppm (br, 2H), aromatic proton (H-6) at δ 5.98 ppm (s, 1H), methylene group (H-1') at δ 2.43 ppm (t, 2H, J = 7.05 Hz), methylene group (H-2') at δ 1.48 ppm (t, 2H, J = 6.89 Hz), methylene group (H-3'-10') at δ 1.24 ppm (s, 16H) and terminal methyl group (H-11') at δ 0.86 ppm (t, 3H, J = 6.20 Hz). The ¹H NMR chemical shift assignment of Compound <u>1</u> is shown in Table 3.3 compared with those of embelin from previous report.²³

The ¹³C NMR spectrum (CDCl₃) of this compound (Figure 3.6) displayed the important carbon signals at 161.1 (C-1, C-2, C-4 and C-5), 116.1 (C-6), 102.2 (C-3) and 22.5 (C-1'). Others signals were detected and compared with those of embelin from previous report²³ as presented in Table 3.4.



Figure 3.3The IR spectrum of Compound 1

Wavenumber(cm ⁻¹)	Intensity	Tentative assignments
3308	Strong	O-H stretching vibration of hydroxyl group
2919, 2847	Medium	C-H stretching vibration of CH ₂ and CH ₃
1613	Strong	C=O stretching vibration of benzoquinone
1460	Weak	C-H bending vibration of CH ₂
1332	Strong	C-H bending vibration of CH ₃
1193	Strong	C-O stretching vibration
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Figure 3.4 The mass spectrum of Compound <u>1</u>



Scheme 3.2 The possible mass fragmentation pattern of Compound $\underline{1}^{22}$



Figure 3.5 The ¹H NMR spectrum of Compound $\underline{1}$

Table 3.3	The ¹ H NMR chemical shift assignment of Compound <u>1</u> compared
	with embelin ²³

Position	Chemical shift (ppm)			
	Compound <u>1</u>	Embelin		
-OH	7.67 (br, 2H)	7.72 (br, 2H)		
H-6	5.98 (s, 1H)	6.01 (s, 1H)		
H-1'	2.43 (t, 2H, <i>J</i> = 7.05 Hz)	2.45 (t, 2H, $J = 7.5$ Hz)		
H-2′	1.48 (t, 2H, <i>J</i> = 6.89 Hz)	1.47 (m, 2H)		
H-3'-H-10'	1.24 (overlapping, 16H)	1.22 (overlapping, 16H)		
H-11′	0.86 (t, 3H, J = 6.20 Hz)	0.88 (t, 3H, $J = 7.0$ Hz)		



Figure 3.6 The ¹³C NMR spectrum of Compound $\underline{1}$

Table 3.4	The	^{13}C	NMR	chemical	shift	assignments	of	Compound	<u>1</u>	and
	embe	elin ²³								

	Position	Chemical shift (ppm)			
		Compound <u>1</u>	Embelin		
	1,2,4,5	161.1	170.0		
	3	102.2	103.5		
	6	116.1	117.5		
ר ב	11	22.5	22.3		
ġ.	2'-10'	22.6 - 31.9	27.8 - 29.3		
	11′	14.1	14.0		

In distinction to all spectroscopic evidences and physical properties compared with those reported in literature, Compound $\underline{1}$ was confident to conclude its structure as 2,5-dihydroxy-3-undecyl-1,4-benzoquinone (embelin).



Compound 1: 2,5-dihydroxy-3-undecyl-1,4-benzoquinone

According to previous reports concerning with chemical constituents determination of *Aridisia colorata* Roxb. Luanratana *et al.* claimed that the barks and fruits of this plant contained a dihydroxybenzoquinone, 2,5-dihydroxy-3-tridecyl-1,4-benzoquinone or rapanone, as the major component.⁹ Nevertheless, endorsing with various spectroscopic information this present examination manifestly demonstrated that the major content was not rapanone, but was 2,5-dihydroxy-3-undecyl-1,4-benzoquinone or embelin. The difference of these two structures: rapanone and embelin was the number of carbon atoms containing in saturated aliphatic side chain The methylene group on the side chain of embelin is less than that of rapanone by two methylene groups, 11 and 13 atoms, respectively.

As aforementioned discussion, the mass spectrum and the 13 C NMR spectrum of Compound <u>1</u> that revealed the molecular ion peak at m/z 294 and the 11 carbon signals of saturated side chain this molecule, respectively. These informative data strongly supported the occurrence of embelin as a major constituent in this particular plant.

3.4 Solvent Variation in Embelin Extraction Procedure

The search for appropriate solvents for embelin extraction procedure was attempted. In section 3.3, embelin was deposited upon the evaporation of solvent after soaking dry fruit material with dichloromethane. Consequently, other solvents (1000 mL) including diethyl ether, ethyl acetate and acetone were employed to examine whether it could be used in place of dichloromethane. The general protocol was as follows: 300 g of dry fruit materials was extracted with selected solvent under refluxing for 2 hours. The acquired quantity of embelin from extraction with each solvent is summarized as shown in Table 3.5.

 Table 3.5
 Weight and percentage yield of embelin from the extraction with selected solvents

Solvent	Weight (g) of embelin	Percentage yield
Diethyl ether	0.030	0.010
Dichloromethane	0.048	0.016
Ethyl acetate	0.015	0.005
Acetone		-

The extraction result showed that embelin could be extracted with non-polar solvents. The extraction of *Ardisia colorata* Roxb. fruits with dichloromethane gave the best percentage yield among all selected solvents.

3.5 Separation of Dichloromethane Extract (Fraction I)

Fraction I, the dichloromethane extract was obtained as a dark brown material 114 g (2.28% w/w of dried fruit materials). From thin-layer chromatography, it was demonstrated that at least six components (solvent system: 30% ethyl acetate in hexane) were present in this fraction. A portion of crude (100 g) was chromatographed on silica gel column chromatography to separate Fraction I into small fractions according to their polarity, using gradient elution started from *n*-hexane followed by a mixture of *n*-hexane:ethyl acetate, ethyl acetate and ethyl acetate:methanol, respectively. The eluted solution was collected approximately 500 mL. Monitoring each fraction by thin-layer chromatography technique, the fractions

that demonstrated similar features were combined. The results of separation Fraction I is presented as shown in Table 3.6.

Fraction	Solvent system	Remarks	Weight (g)
IA	Hexane,	Yellow oil	6.81
	20%EtOAc:Hexane	11/2_	
IB	30%EtOAc:Hexane	Sticky brown material	17.01
IC	50%EtOAc:Hexane	Dark brown semisolid	5.22
ID	75%EtOAc:Hexane	Dark brown semisolid	4.45
IE	EtOAc	Dark brown semisolid	11.38
IF	30% MeOH:EtOAc	Dark brown semisolid	19.68
IG	50%MeOH:EtOAc	Sticky dark brown material	12.54

Table 3.6The separation of dichloromethane extract by silica gel column
chromatography

3.5.1 Separation of Fraction IA

Thin-layer chromatography of Fraction IA using 30% ethyl acetate-hexane as a developing solvent demonstrated that there were at least 2 components in Fraction IA. This fraction was further subjected to silica gel column chromatography. Hexane, a mixture of ethyl acetate and hexane, and ethyl acetate were used as eluents. The eluted solution was collected about 50 mL for each fraction and monitored by TLC. The results of the separation of Fraction IA are presented in Table 3.7.

Eluents	Fraction no.	Remarks	Weight (g)
Hexane	1-5	White solid in yellow oil, Substance 2	0.98
3%EtOAc:Hexane	6-12	Yellow liquid, Mixture <u>3</u>	3.24
10%EtOAc:Hexane	13-21	Sticky brown material	1.14
30%EtOAc:Hexane	22-30	Sticky brown material	1.05

From the results of separation, Fraction no. 1-5 was purified by recrystallization with hot ethyl acetate several times to afford bright white plates (21.5 mg) designated as MIxture $\underline{2}$. In addition, according to TLC monitoring of Fraction no. 6-12, one spot with tail was clearly detected. This fraction was then purified by flash column chromatography. The yellow semisolid (2.5 g) designated as Mixture $\underline{3}$ was received.

3.5.1.1 Structural Elucidation of Mixture 2

Mixture <u>2</u> (21.5 mg, 0.0245 % w/w of dichloromethane crude extract) had R_f 0.80 (40% EtOAc:Hexane), m.p. 51-53 °C soluble in dichloromethane, chloroform and methanol; slightly soluble in hexane and ethyl acetate. The IR spectrum of this mixture (Figure 3.7) displayed the characteristic absorption peak of hydroxyl group at 3446 cm⁻¹. Other signals were tentatively assigned as shown in Table 3.8.

The ¹H NMR spectrum of Mixture <u>2</u> (Figure 3.8) showed signals of terminal methyl proton at δ 0.86 ppm, methylene proton at δ 1.24 ppm and methylene protons which directly connected to oxygen atom at δ 1.55 ppm.

Table 3.8 The IR absorption band assignments of Mixture	e <u>í</u>
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Wave number (cm ⁻¹)	Intensity	Tentative assignment
3446	Broad	O-H stretching vibration of hydroxyl group
2909,2842	Strong	C-H stretching vibration of CH ₂ and CH ₃
1465	Medium	C-H bending vibration of CH ₂

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Figure 3.7The IR spectrum of Mixture 2



Figure 3.8 The ¹H NMR spectrum of Mixture $\underline{2}$

$CH_3-(CH_2)_n-OH$

Mixture <u>2</u>

3.5.1.2 Structural Elucidation of Mixture 3

Mixture <u>3</u>, the yellow semisolid (2.5 g, 2.85% w/w of crude extract) had R_f 0.62 (10% EtAOc:Hexane). It was soluble in ethyl acetate, dichloromethane, chloroform and methanol but slightly soluble in hexane. This mixture showed greenblue solution with Liebermann Burchard's reagent expressing that this mixture contained a steroidal nuclei.

The IR spectrum of this mixture (Figure 3.9) showed a major characteristic absorption band of carbonyl group belonging to an ester at 1743 cm⁻¹. Other signals were tentatively assigned as shown in Table 3.9.

Table 3.9The IR absorption band assignments of Mixture 3

Wave number (cm ⁻¹)	Intensity Tentative assignment		
2921, 2846	Medium C-H stretching vibration of CH ₂ and Cl		
1743	Strong C=O stretching vibration of ester		
1461	Medium C-H bending vibration of CH ₂		
1376	Weak C-H bending vibration of CH ₃		
1160	Strong	C-O stretching vibration	
723	Weak	CH_2 rocking vibration	
	59.9		



Figure 3.9 The IR spectrum of Mixture <u>3</u>

From the physical properties and all spectroscopic data, this mixture could be concluded as a steroidal ester. Therefore, the hydrolysis reaction was set up for further identification of this mixture.

Hydrolysis of Mixture 3

A solution of 10% ethanolic KOH (40 mL) was added to Mixture <u>3</u> (2.0 g) and the solution was stirred under refluxing on a water bath for 4 hours. Evaporation of ethanol gave a solid, which was further extracted with diethyl ether 150 mL three times. The combined diethyl ether was dried over anhydrous sodium sulfate. Evaporation of the solvent furnished a solid with yellow oil. After recrystallization with a mixture of hexane-ethyl acetate several times, white amorphous solid designated as Mixture <u>3A</u> (18.4 mg), m.p. 146-147 °C was received.

Study on Mixture <u>3A</u>

Mixture <u>3A</u> was thought to be a mixture of steroids which were generally detected as constituents in plants. Thus, this mixture was *co*-TLC with a mixture of authentic steroids, namely campesterol, stigmasterol and β -sitosterol, previously isolated and characterized from *Rhizophora apicalata* Bl.¹⁸ The same R_f values on TLC plates suggested that Mixture <u>3A</u> be steroidal compounds. Further study was

carried on utilizing GC compared with authentic steroids, namely cholesterol, campesterol, stigmasterol and β -sitosterol. The results of GC analysis (Figure 3.10) of this mixture showed the retention times at 28.16 and 31.91 min, respectively corresponding to those of stigmasterol and β -sitosterol, respectively. The composition of steroids in this mixture is presented in Table 3.10.



Figure 3.10 The chromatogram of Mixture <u>3A</u> from GC analysis

Table 3.10The composition of steroids in Mixture <u>3A</u>

Name	Retention time (min)	% Composition
Stigmasterol	28.16	71
β-sitosterol	31.91	28

Based upon the color test result, physical properties and all spectroscopic data including GC analysis, it could be concluded that Mixture <u>3A</u> is a mixture of stigmasterol and β -sitosterol.



Study on Mixture <u>3B</u>

Mixture <u>3B</u> is an acid part from the hydrolysis of Mixture <u>3</u>. This mixture was purified with column chromatography using gradient elution of *n*-hexane and ethyl acetate as solvent system. Methylation of this mixture by tetramethylsilyl diazomethane followed the method described in section 2.6.3 led to the formation of methyl ester of Mixture <u>3B</u>. The methyl ester derivative was further examined by GC-MS analysis without further purification. The gas chromatography was carried out on DB-35 column. The chromatogram of this derivative is exhibited as shown in Figure 3.11.

The component with retention time at 11.99 min was found to be a major component in this mixture. The mass spectrum of this component gave the molecular ion peak, M^+ , at m/z 270. Compared with library data (NIST database), it was found that this major composition was corresponding to hexadecanoic acid methyl ester (Calcd. for C₁₇H₃₄O₂: MW 270.45). The mass spectrum of this component and library data of hexadecanoic acid (palmitic acid) methyl ester is shown in Figure 3.12.



Figure 3.11 The chromatogram of Mixture <u>3B</u> methyl ester



Figure 3.12The mass spectrum of the major component ($R_t = 11.99$ min) of
Mixture <u>3B</u>

With the aids of the chemical reaction together with spectroscopic data and GC-MS analysis, it could be obviously concluded that Mixture <u>3</u> was stigmasteryl-3-*O*-palmitate and β -sitosteryl-3-*O*-palmitate.



Stigmasteryl-3-O-palmitate



β-Sitosteryl-3-*O*-palmitate

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3.5.2 Separation of Fraction IB

The TLC of Fraction IB in 30% ethyl avetate:hexane revealed a spot identical with the authentic sample of α -amyrin or β -amyrin. Thus, Fraction IB was further separated with silica gel column, and the result of the separation of Fraction IB is presented in Table 3.11. The gradient elution of hexane and ethyl acetate was used. Each fraction was monitored by thin-layer chromatography. According to the TLC results, the fraction which exhibited the same spot as authentic sample was Fraction IB.2. Therefore, Fraction IB.2 was further purified by silica gel column. The obtained solid was further recrystallized with hot ethanol several times to yield white needle crystal (59.6 mg) specified as Mixture <u>4</u>.

Eluents	Fraction	Remarks	Weight (g)
10% EtOAc:hexane	IB.1	Yellow oil	0.52
15% EtOAc:hexane	IB.2	White crystal in yellow solution;	
		Mixture <u>4</u>	0.95
30% EtOAc:hexane	IB.3	Sticky dark yellow material	1.78
60% EtOAc:hexane	IB.4	Sticky brown material	2.05

Table 3.11	The results of the separation of Fraction IB

3.5.2.1 Structural Elucidation of Mixture 4

Mixture <u>4</u>, white needle crystal 59.6 mg (0.0679% w/w of dichloromethane crude extract) R_f 0.62 (40% EtOAc:hexane) melted at 189-190 °C. This compound was soluble in chloroform, dichloromethane and ethyl acetate. From the Liebermann-Burchad's test, Mixture <u>4</u> provided a red solution, suggesting the presence of triterpenoidal nuclei.

This mixture was subjected to GC-MS analysis to explore the mixture components using DB-35 column. The chromatogram of Mixture $\underline{4}$ revealed that there were 3 components present as shown in Figure 3.13.

The results of GC-MS analysis of this mixture revealed the retention times at 17.96, 19.56 and 20.66 min, respectively. The major component was component 3 with retention time at 20.66 min. The mass spectrum of each component is exhibited as shown in Figures 3.14 - 3.16.



Figure 3.13 The chromatogram of Mixture <u>4</u> from GC analysis



Figure 3.14 The mass spectrum of component 1 (Rt 17.96 min) of Mixture $\underline{4}$



Figure 3.15 The mass spectrum of component 2 (Rt 19.56 min) of Mixture 4



Figure 3.16 The mass spectrum of component 3 (Rt 20.66 min) of Mixture 4

Table 3.12	The composition of Mixture $\underline{4}$	

Component	Retention time (min)	Mass	% Composition
1	17.96	346	17
2	19.56	426	21
3	21.73	460	62

From the comparison of the mass spectrum of each component with NIST data library, it was found that the first component is possibly 2(1H)Napthalenone. The second component with m/z at 426 was equivalent to $C_{30}H_{50}O$ (MW 426.72). The third component which showed m/z at 460, was perhaps an impurity that would not be considered in the meanwhile.

The spectroscopic study including IR and ¹H NMR for this mixture was performed as shown in Figures 3.17-3.18.

The IR spectrum of this mixture (Figure 3.17) represented the characteristic absorption band of hydroxyl group at 3248 cm⁻¹, C-H stretching at 2940 and 2856 cm⁻¹ and C-O stretching at 1037 cm⁻¹. The ¹H NMR spectrum (Figure 3.18) exhibited the methylene proton signal at 0.72-2.16 ppm, carbinol proton (H-3) at 3.21 ppm and olefinic proton (H-12) at 5.16 ppm. The mass spectrum of this mixture was compared with that of α -amyrin from NIST data library as shown in Figure 3.19 and the possible fragmentation¹⁸ is shown in Scheme 3.3.

Considering between α -amyrin and β -amyrin, these two triterpenoids gave closely characteristic peaks on mass spectrum. Therefore, the structure of second component from Mixture <u>4</u> was ensured by the comparison their m.p. and the ¹³C NMR signals C-12 and C-13. The comparison data of Mixture <u>4</u>, α -amyrin²⁴ and β -amyrin²⁵ are presented as shown in Table 3.13.

Triterpenoids	m.p. (°C)	¹³ C NMR chemical shift (ppm)	
		C-12	C-13
α-amyrin	186.0	124.3	139.3
β-amyrin	197.0-197.5	121.7	145.2
Mixture <u>4</u>	189.0-190.0	124.6	138.9

Table 3.13 The comparison data of α -amyrin and β -amyrin and Mixture <u>4</u>



Figure 3.17The IR spectrum of Mixture $\underline{4}$



Figure 3.18 The ¹H NMR spectrum of Mixture $\underline{4}$



Figure 3.19 The mass spectrum of Mixture $\underline{4}$ (A) and α -amyrin (B)





From all spectroscopic evidences, physical properties and GC-MS analysis compared with literatures, it could be obviously concluded that the second component of Mixture $\underline{4}$ is α -amyrin.



The second component of Mixture $\underline{4}$; α -amyrin

3.5.3 Separation of Fractions IC and ID

According to the monitoring of Fractions IC and ID with TLC (solvent system: 30% ethyl acetate:hexane), these two fractions displayed similar pattern. Hence, Fractions IC and ID were combined and renamed as Fraction ICD. Fraction ICD was further separated with silica gel column using gradient elution system of hexane, a mixture of hexane:ethyl acetate, ethyl acetate and a mixture of ethyl acetate:methanol. Each fraction was collected approximately 50 mL, and was monitored by thin-layer chromatography technique. The fractions demonstrated similar pattern on TLC plates were combined. The results of the separation of Fraction ICD are presented as shown in Table 3.14.

1 able 5.14	The results of the separation of Fraction ICD

Eluents	Fraction No.	Remarks	Weight (g)
Hexane	1-3 (ICD1)	Yellow oil	0.06
5%-10% EtOAc:hexane	4-10 (ICD2)	White solid in brown oil	1.08
20%-50% EtOAc:hexane	11-13 (ICD3)	Sticky brown material	1.54
60%-70% EtOAc:hexane	14-17 (ICD4)	Red-brown oil	1.02
70% EtOAc:hexane-EtOAc	18-25 (ICD5)	Dark brown semisolid	1.57
20% MeOH:EtOAc	26-35 (ICD6)	Dark brown semisolid	2.41

Fraction 4-10 coded as ICD2, was the white solid deposited in brown oil when it was stand at room temperature for a moment. This fraction was further purified by separation with silica gel column. Similarly, Fraction 14-17 coded as ICD4 were redbrown oil that showed two spots on TLC.

Separation of Fraction ICD2

Fraction ICD2 was chromatographed on silica gel column, using gradient elution of hexane and ethyl acetate. Each fraction was monitored with thin-layer chromatography, fraction which showed similar pattern was combined. Fraction 7-8 (see also Table 3.15) obliged white amorphous in solution. After filtration, the white amorphous was recrystallized by ethyl acetate for several times to yield off-white needle crystals (36.9 mg) designated as Compound 5.

Eluents	Fraction No.	Remarks	Weight (g)
Hexane	1-2	Yellow oil	0.050
5% EtOAc:hexane	3-6	White amorphous in yellow oil	0.102
10% EtOAc:hexane	7-8	White amorphous in yellow oil;	0.324
		Compound <u>5</u>	
20% EtOAc:hexane	9-12	Sticky brown material	0.420

Table 3.15The results of the separation of Fraction ICD2

3.5.3.1 Structural Elucidation of Compound 5

The off-white needle crystal (36.9 mg, 0.0421 % w/w of dichloromethane crude extract) displayed a single spot on TLC at R_f 0.47 in 40% ethyl acetate:hexane. It was melted at 139-141 °C. This compound was soluble in chloroform, dichloromethane and ethyl acetate. Compound <u>5</u> showed a positive result with Liebermann-Burchard's reagent: it gave a blue-green solution. Therefore, this compound was presumed to contain a steroidal nuclei. According to the spectroscopic study, the IR spectrum (Figure 3.20) of this compound gave the absorption peaks of hydroxyl group at 3453 cm⁻¹, C=C stretching vibration at 1637 cm⁻¹. Other signals were tentatively assigned as shown in Table 3.16.

GC-MS analysis was selected to study the composition of this compound, using DB-35 column. Figure 3.21 presents the chromatogram of Compound <u>5</u>. The major component with retention time at 6.56 min gave the molecular ion peak at m/z412 (Figure 3.22). Compared with NIST data library, it was found that this compound was compatible with stigmasterol (Calcd for C₂₉H₄₈O: MW 412.69). The mass spectrum of this compound and library data of stigmasterol are shown in Figure 3.23. The possible mass fragmentation of this compound was proposed as shown in Scheme 3.4.



Figure 3.20 The IR spectrum of Compound <u>5</u>

Table 3.16The IR absorption band assignments of Compound 5

Wavenumber (cm ⁻¹)	Intensity	Tentative assignments		
3435	Strong	O-H stretching vibration of hydroxyl group		
2937, 2867	Medium	C-H stretching vibration of CH ₂ and CH ₃		
1637	Medium	C=C stretching vibration		
1458	Weak	C-H asymmetric bending of CH ₂ and CH ₃		
1376	Weak	C-H symmetric bending of CH ₃		
1038	Weak	C-O stretching vibration		
- M 101	1199	791 I I I I I I I I I I I I I I I I I I I		



Figure 3.21 The chromatogram of Compound <u>5</u>



Figure 3.22 The mass spectrum of Compound <u>5</u>



Figure 3.23 The mass spectrum of Compound (A) and stigmasterol (B)



Scheme 3.4 The possible mass fragmentation pattern of Compound <u>5</u>

As a result of its physical properties, GC-MS analysis and spectroscopic data, it could noticably be concluded that this compound was stigmasterol.



Compound 5: stigmasterol

Separation of Fraction ICD4

Fraction ICD4 containing red-brown oil was further separated by silica gel column. The gradient elution system of hexane, ethyl acetate and methanol was used.

In addition, from the DPPH screening for radical scavenging by TLC autographic assay, this fraction had at least 2 components that showed scavenging for DPPH assay. The results of separation are summarized as shown in Table 3.17.

Table 3.17The results of the separation of Fraction ICD4

Eluents	Fraction No.	Remarks	Weight (g)
20% EtOAc:hexane	ICD4.1	Yellow oil	0.025
50% EtOAc:hexane	ICD4.2	White solid, Compound <u>5</u>	0.084
70% EtOAc:hexane	ICD4.3	Red oil	0.898
100 % EtOAc	ICD4.4	Brown material	0.091

Fraction ICD4.2 gave a white solid which was further purified by recrystallization from hot ethyl acetate for several times to yield the off-white needle crystal. This crystal was melted at 140 °C and soluble in chloroform and dichloromethane. According to the spectroscopic study of this crystal and its physical properties, it was believed that Compound 5 was stigmasterol.

Separation of Fraction ICD4.3

According to the result of separation Fraction ICD, Fraction ICD4.3 was obtained as red oil material. The TLC of this fraction in 10% methanol in chloroform revealed 2 spots. This fraction was then rechromatographed with flash column, using gradient elution of chloroform and methanol. After separation, it still contained impurities. Therefore this portion was reseparated by chromatotron technique using gradient elution of 100% chloroform to 10 % methanol in chloroform and operated under ultraviolet light wavelength 254 nm condition. The separation of Fraction ICD4.3 by flash column chromatograph and chromatotron techniques yielded two compounds. The result of the separation of Fraction ICD4.3 is presented in Table 3.18. The first is brown oil 808.5 mg, designated as Mixture <u>6</u>, and the second is pale-yellow oil 19.2 mg, specified as Compound <u>7</u>. Both of them gave a positive result for TLC autographic assay for scavenging effect on DPPH radical.

Table 3.18The results of the separation of Fraction ICD4.3

Eluents	Fraction	Remarks	Weight (g)
CHCl ₃	1-15	Brown oil; Mixture <u>6</u>	0.808
10% MeOH/CHCl ₃	16-23	Pale-yellow oil; Compound <u>7</u>	0.019

3.5.3.2 Structural Elucidation of Mixture <u>6</u>

The brown oil (808.5 mg, 0.9217 % w/w of dichloromethane crude extract) revealed only one spot at R_f 0.49 in 10% methanol in chloroform. This mixture was soluble in ethyl acetate, dichloromethane and chloroform.

The IR spectrum (Figure 3.24) of this mixture exhibited absorption peaks of hydroxyl group at 3350 cm⁻¹, double bond of aromatic ring signal at 1598 cm⁻¹ and 1466 cm⁻¹. Other signals were tentatively assigned as shown in Table 3.19.



Figure 3.24 The IR spectrum of Mixture <u>6</u>

Table 3.19	The IR absorption	a band assignments	of Mixture 6
		0	

Intensity	Tentative assignments	
Strong	O-H stretching vibration of hydroxyl group	
Strong	C-H stretching vibration of CH ₂ and CH ₃	
Strong	C-C stretching vibration of aromatic	
Strong	=CH bending vibration in plane	
Strong	C-H bending in plane vibration of aromatic	
Medium	C-H bending out of plane vibration of aromatic	
	Intensity Strong Strong Strong Strong Strong Medium	

The ¹H NMR spectrum (CDCl₃) of this mixture (Figure 3.26) demonstrated the signal of aromatic protons at δ 6.25 ppm (m, 2H) and δ 6.16 ppm (m, 1H), olefinic proton at δ 5.36 ppm (m, 3H), bisallylic methylene proton at δ 2.76 ppm (t, 1H, *J*=5.78 Hz), benzyl methylene protons at δ 2.42 ppm (t, 2H, *J*=7.36 Hz), allylic methylene protons at δ 2.02 ppm (br, 4H), homobenzyl methylene protons at δ 1.50 ppm (br, 2H), methylene protons at δ 1.27 ppm (overlappig, 16H) and terminal methyl protons at δ 0.87 ppm (t, 3H, *J*=6.28 Hz).

The ¹³C NMR spectrum (CDCl₃) of this mixture (Figures 3.27-28) gave the signals for tetra-substituted benzene at δ 156.2, 146.4, 108.2 and 100.3 ppm and six olefinic carbon signals at δ 130.3, 130.2, 129.9, 129.8, 128.0 and 127.9 ppm.

According to the infrared, ¹H and ¹³C NMR spectra, it was found that this mixture may contain resorcinol ring and unsaturated side chain. Thoroughly, this unsaturated side chain is consisted of 15 and 17 carbon atoms, respectively. This observation was confirmed by MS analysis (Figure 3.25). To illustrate this, the mass spectrum showed two molecular ion peaks (M^+) at m/z 344 (calcd. for C₂₃H₃₆O₂: MW 344.27) designated as Mixture <u>6A</u> and m/z 318 (calcd. for C₂₁H₃₄O₂: MW 318.26) designated as Mixture <u>6B</u>. Its also showed a base peak at m/z 124 corresponding to a benzylic fragmentation.

The unsaturated 17 carbon atom side chain of Mixture <u>6A</u> contained two skipped double bonds. This was endorsed by the bisallylic methylene proton signal at δ 2.76 ppm and olefinic carbon signals at δ 127.9, 128.0, 130.2, and 130.3 ppm. While Mixture <u>6B</u> had one double bond on 15 carbon atom side chain that showed two olefinic carbon signals at δ 129.8 and 129.9 ppm.



Figure 3.25 The mass spectrum of Mixture <u>6</u>





Figure 3.27 The 13 C NMR spectrum of Mixture <u>6</u>



Figure 3.28 The ¹³C NMR spectrum of Mixture <u>6</u> (expanded)

The double bond was unequivocally determined to be Z stereochemistry from the comparison of the ¹³C NMR data with previous reports²⁶, the known 3-(heptadec-8'-Z-enyl)-1,2-dimethoxybenzene and its 8'-E isomer. The comparison of ¹³C NMR data of Mixture <u>6</u> with both Z- and E-isomers are summarized as shown in Table 3.20.

Table 3.20	The ¹³ C NMR	data of Mixture 6 compared	with their references ²⁶
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Position	Chemical shift (ppm)				
	Mixture <u>6</u>	Z-isomer	<i>E</i> -isomer		
Allylic carbon	27.21	27.18	32.58		
Olefinic carbon	129.91, 128.02	129.82, 129.87	130.28,130.34		

According to Table 3.20, it was found that the carbon signals of Mixture <u>6</u> were coincident with those of 8'-Z-isomer, but not with those of 8'-E-isomer. Thus, the stereochemistry of double bond in unsaturated side chain of Mixture <u>6</u> was believed to be Z-form. The assignments of proton and carbon signals of Mixtures <u>6A</u> and <u>6B</u> are summarized as shown in Figure 3.29 and Table 3.21.





Figure 3.29 The ¹H NMR and ¹³C NMR assignments of Mixture $\underline{6}$
Position	n Mixture <u>6A</u>		Position	Mixture <u>6B</u>		
	¹ H	¹³ C		¹ H	¹³ C	
1	-	156.2	1	-	156.2	
2	6.16 (m, 1H)	100.3	2	6.16 (m,1H)	100.3	
3	-	156.2	3	-	156.2	
4,6	6.25 (m, 2H)	108.2	4,6	6.25 (m, 2H)	108.2	
5	-	146.4	5	-	146.4	
1′	2.42 (t, 2H, <i>J</i> =7.36 Hz)	35.9	-1'	2.42 (t, 2H, <i>J</i> =7.36 Hz)	35.9	
2'	1.50 (br, 2H)	31.8	2'	1.50 (br, 2H)	31.8	
3'		31.5	3'		31.5	
4′		31.1	4′		31.1	
5'	1.27 (overlapping)	29.7	5'	1.27 (overlapping)	29.7	
6′		29.3	6'		25.3	
7′,13′	2.02 (br, 4H)	27.2	7′,10′	2.02 (br, 4H)	27.2	
8′	5.36 (m, 3H)	130.3	8'	5.36 (m, 3H)	129.9	
9′	5.36 (m, 3H)	130.2	9'	5.36 (m, 3H)	128.0	
10′	2.76 (t, 1H, <i>J</i> =5.78 Hz)	25.6	11′		27.2	
11′	5.36 (m, 3H)	128.0	12'	1.27 (overlapping)	25.6	
12′	5.36 (m, 3H)	127.9	14′		22.7	
14′		25.7	15'	0.87 (t, 3H, <i>J</i> =6.28 Hz)	14.1	
15′	1.27 (overlapping)	25.6				
16′		22.7				
17′	0.87 (t, 3H, <i>J</i> =6.28 Hz)	14.1	וכחו			
$\begin{array}{c} 0H \\ 2 \\ HO \\ 6 \\ 1' \\ HO \\ 6 \\ 1' \\ 1' \\ 1' \\ 1' \\ 1' \\ 1' \\ 1' $						

Table 3.21The 1 H and 13 C NMR chemical shift assignments of Mixture 6

According to all spectroscopic data, the physical properties and previous study²⁸ of this mixture. It was believed that this mixture was composed of two components as Mixture <u>6A</u> or 1,3-dihydroxy-5-(heptadec-8,11-dienyl)benzene and Mixture <u>6B</u> or 1,3-dihydroxy-5-(pentadec-8-enyl)benzene.



1,3-dihydroxy-5-(heptadec-8,11-dienyl)benzene; Mixture 6A



1,3-dihydroxy-5-(pentadec-8-enyl)benzene; Mixture 6B

Mixture 6

3.5.3.3 Structural Elucidation of Compound 7

Compound <u>7</u>, the pale-yellow oil 19.2 mg (0.0219 % w/w of dichloromethane crude extract) showed a single spot at R_f 0.65 in 10% methanol in chloroform was obtained from re-separation of Fraction ICD4.3. This compound was soluble in chloroform and methanol.

The IR spectrum of this compound (Figure 3.30), exhibited the absorption peaks of hydroxyl group of carboxylic acid in region of 3100-3300 cm⁻¹, carbonyl group at 1712 cm⁻¹. Other signals were tentatively assigned in Table 3.22.



Figure 3.30 The IR spectrum of Compound <u>7</u>

Table 3.22The IR absorption band assignments of Compound 7

Wavenumber (cm ⁻¹)	Intensity	Tentative assignments
3100-3300	Broad	O-H stretching vibration of carboxylic acid
2921, 2856	Strong	C-H stretching vibration of CH ₂ and CH ₃
1712	Strong	C=O stretching vibration of carboxylic acid
1456	Medium	C-H bending vibration of CH ₂ and CH ₃

As an informative result of mass spectrometry (Figure 3.31), this compound showed the molecular ion peak at m/z 280 (Calcd. for $C_{18}H_{32}O_2$: MW 280.45) corresponding to unsaturated long chain fatty acid, linoleic acid. Furthermore, the relatively intense fragment ions at m/z (relative intensity) 60(22), 67(100), 81(87), 95 (83) and 109(43) are detected. The fragment ion at m/z 60 is formed by McLafferty (H)rearrangement. The possible mass fragmentation pattern is proposed as shown in Scheme 3.5.

The ¹H NMR spectrum of this compound (Figure 3.32) exhibited the chemical shift of proton signals of the olefinic signal at 5.32 ppm (m, 4H), the bisallylic signal at 2.76 ppm (t, 2H, J=5.83 Hz), the α -proton (H-2) at 2.33 ppm (t, 2H, J=7.53 Hz), the allylic signal at 2.04 ppm (br, 4H), the β -proton (H-3) at 1.61 ppm (br, 2H), the methylene signal is overlapped at 1.29 ppm (14H), and the terminal methyl proton at 0.87 ppm (t, 3H, J=6.97 Hz).

According to the ¹³C NMR of this compound (Figure 3.33), it displayed 18 carbon signals. The important signals are as follows: the carbonyl signal (C-1) at 179.4 ppm and four olefinic carbons at 127.9, 128.0, 130.0, and 130.2 ppm, respectively. The other carbon signals are assigned as presented in Table 3.23 (included with the proton signals). The structure of octadeca-9,12-dienoic acid or linoleic acid, is shown below.



Octadeca-9,12-dienoic acid or linoleic acid



Figure 3.31The mass spectrum of Compound 7



Scheme 3.5 The possible mass fragmentation pattern of Compound <u>7</u>



Figure 3.32 The ¹H NMR spectrum of Compound <u>7</u>



Figure 3.33 The ¹³C NMR spectrum of Compound <u>7</u>

acid²⁸

	Chemical shift (ppm)						
Position	Compound	7	linoleic acid				
	¹ H	¹³ C	¹ H	¹³ C			
1	-	179.4	-	177.0			
2	2.33 (t, 2H, <i>J</i> =5.83) 33.9 2.37 (t, 2H, <i>J</i> =7.1)		35.8				
3	1.61 (m, 2H)	24.7	1.67 (m, 2H)	25.1			
4	*	29.8	**	29.7			
5	*	29.3	**	30.1			
6	*	29.1	**	30.4			
7	*	29.0	**	30.3			
8	2.04 (m, 2H)	27.2	2.10 (m, 2H)	27.4			
9	5.32 (m, 2H)	130.2	5.32 (m, 2H)	131.0			
10		129.0		128.4			
11	2.76 (t, 2H, <i>J</i> =7.53)	25.6	2.80 (m, 2H)	25.2			
12	5.32 (m, 2H)	127.9	5.32 (m, 2H)	128.0			
13		130.0		130.6			
14	2.04 (m, 2H)	27.2	2.10 (m, 2H)	27.4			
15	*	29.0	**	30.0			
16	161 1 * 1 d 61	31.5		32.6			
17	*	22.6	**	23.2			
18	0.87 (t, 3H, <i>J</i> =6.97)	14.1	0.91 (t, 3H, <i>J</i> =7.4)	14			

Note: The methylene signals of Compound <u>7</u> were resonated at 1.29 ppm

(overlapping, 14H), and those of linoleic acid were reported at 1.35 ppm (br, 14H).

Additionally, the structure of Compound $\underline{7}$ was deduced from the analysis of the two-dimensional ¹H correlation spectroscopy (COSY), and heteronuclear multiple bond connectivity (HMBC) spectra. The COSY spectrum of Compound $\underline{7}$ (Figure 3.35) reasonably exposed correlations between H-2 (δ 2.33) and H-3 (δ 1.61), between allylic protons H-8, H-14 (δ 2.04) and olefinic protons H-9, H-13 (δ 5.32), and between bisallylic proton H-11 (δ 2.76) and olefinic protons H-10, H-12 (δ 5.32), respectively.

According to the HMBC spectrum (Figure 3.36), it was clearly manifested the correlations between C-1 (δ 179.4) and H-2, H-3, between C-8 (δ 27.2) and H-10, between C-14 (δ 27.2) and H-12, and C-11 (δ 25.6) and H-9, H-13, respectively. The selected COSY and HMBC correlations of Compound <u>7</u> are demonstrated as depicted in Figure 3.34.



Figure 3.34 The selected COSY and HMBC correlations of Compound 7

Additional proof was carried out by allowing Compound $\underline{7}$ and authentic linoleic acid react with trimethylsilyl diazomethane (TMSCHN₂) followed the procedure described in Section 2.6.3. The methylation reaction of Compound $\underline{7}$ yielded the methyl ester derivative. The methyl esters of both Compound $\underline{7}$ and authentic linoleic acid were further subjected to GC analysis to confirm that this compound is octadeca-9,12-dienoic acid or linoleic acid. The chromatograms and retention time of both compounds were presented as shown in Figure 3.37.



Figure 3.35 The COSY spectrum of Compound <u>7</u>



Figure 3.36 The HMBC spectrum of Compound <u>7</u>



Figure 3.37 The chromatograms of methyl ester derivatives of authentic linoleic acid (A) and Compound <u>7</u> (B)

The GC analysis of methyl ester derivative of this compound revealed the retention time at 7.48 min, which was corresponded to that of authentic linoleic acid.

Therefore, as a result of physical properties, spectroscopic evidences, chemical reaction, and GC-analysis, Compound <u>7</u> was confident to conclude its structure as octadeca-9,12-dienoic acid (linoleic acid).



Furthermore, according to the monitoring the EtOAc and MeOH crude extracts from *Ardisia colorata* Roxb. fruits by thin layer chromatography, these crude extracts exhibited almost similar spots as detected in CH_2Cl_2 crude extract. Moreover, these two crudes showed insignificant effect to both primary screening methods (brine shrimp cytotoxicity and scavenging effect towards DPPH radical). Therefore, there is rationalized not to separate and further explore these two fractions.

3.6 Study on Biological Activities of Isolated Substances from CH₂Cl₂ Extract of *Ardisia colorata* Fruit

As discussed above, the separation of dichloromethane extract from *Ardisia colorata* Roxb. fruits led to the isolation of seven substances. The isolated substances were further studied to search for biological activity. The cytotoxicity test, radical scavenger effect on DPPH radical, antifungal activity and insecticidal activity were selected for exploration followed the protocols described in Chapter II.

3.6.1 Brine Shrimp Lethality Cytotoxic Test

Seven isolated substances were tested for cytotoxic activity against brine shrimp, the LC_{50} values determined by probit analysis program are exhibited as shown in Table 3.24.

Substance	LC ₅₀ (1	Toxicity	
	Acute	Chronic	
1 1 1	5.76	1.72	High
2	UK-9110	1486.31	No
3 3 9	กรถ์บ	650.32	Low
4	1 9 9 99 94	698.55	Low
5	-	709.22	Low
6	230.44	32.36	Moderate
7	443.96	4.99	High

	Table 3.24	Cytotoxicity	activity	against A	rtemia	salina	(brine	shrimp)
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Note: 0-10 μg/mL; High toxicity, 10-100 μg/mL; Moderate toxicity 100-1000 μg/mL; Low toxicity, >1000 μg/mL; No toxicity

Among all isolated substances, Compounds <u>1</u> and <u>7</u> showed high cytotoxicity against brine shrimp at LC₅₀ 1.72 and 4.99 μ g/mL, respectively. The Mixture <u>6</u> showed moderate cytotoxicity at LC₅₀ 32.36 μ g/mL. The other substances were inactive or gave low toxicity against brine shrimp.

From the result above, the cytotoxic activity against brine shrimp of Compounds <u>1</u> and <u>7</u> were found to comparatively be higher than that of dichloromethane extract (LC₅₀ 13 μ g/mL), whereas the other substances were lower. This attained result could clearly convince that the active components responsible for cytotoxic activity from dichloromethane extract should be Compound <u>1</u> or embelin and Compound <u>7</u> or linoleic acid.

3.6.2 Scavenging Effect on DPPH Radical

The DPPH radical scavenging activity by TLC autographic method of isolated substances revealed that Compounds <u>1</u> and <u>7</u> and Mixture <u>6</u> showed significant activity against DPPH radical.

The quantitative analysis for determination IC_{50} value of those substances were then performed using spectrophotometric method as described in Chapter II. The result of radical scavenging effect on DPPH of each compound is presented as shown in Table 3.25. The curve between the percentage of radical scavenging and the concentration of each sample was plotted. Figure 3.38 as an instance shows the curve of radical scavenging effect for Compound 1.

Sample	Concentration (µg/mL)	% radical scavenging		
	1000	74.66		
	500	73.39		
Compound 1	250	66.17		
	125	57.76		
	62.5	41.26		
	1000	62.81		
Mixture <u>6</u>	500	51.84		
	250	48.83		
	125	31.49		
	62.5	24.88		
	500	26.06		
Compound <u>7</u>	250	16.80		
	125	13.54		
	62.5	7.40		
	1000	68.38		
	500	81.67		
BHA	250	79.77		
	125	75.26		
	62.5	66.39		

Table 3.25Radical scavenging effect on DPPH radical



Figure 3.38 DPPH radical scavenging effect of Compound <u>1</u>

The IC_{50} values of each substance were calculated from the logarithmic equation of each curve and were tabulated in Table 3.26.

Table 3.26IC₅₀ values of radical scavenging effect on DPPH radical

Substance	IC ₅₀ (µg/mL)
1	86.34
<u>6</u>	386.03
7	9505.40
BHA [*]	16.88

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Butylated hydroxyanisole (BHA)

From the result of radical scavenging effect on DPPH radical assay, it was found that Compound <u>1</u> (embelin) showed the highest activity at IC₅₀ 86 ppm among all tested substances. From the comparison with BHA (butylated hydroxyanisole), a commercial antioxidant, the activity of Compound <u>1</u> was lower than that of BHA.

As the comparison with preliminary result of dichloromethane crude extract, it was detected that the active component which scavenged the DPPH radical on TLC (Section 3.2.2) should be Compound <u>1</u> or Compound <u>7</u> or Mixture <u>6</u>. Therefore, this inference was ensured by spectrophotometric method. The IC_{50} values of each substance pointed out that Compound <u>1</u> or embelin gave the lowest IC_{50} among those substances tested, although it gave higher value than that of BHA. By this reason, it could obviously conclude that embelin was the active antioxidant agent from dichloromethane crude extract from *A. colorata* fruits.

3.6.3 Preliminary Antifungal Activity on Fusarium oxysporum and Alternaria sp. by Bioautographic Assay

Seven isolated substances from the dichloromethane extract from the fruits of *Ardisia colorata* Roxb. were investigated for antifungal activity. Two phytophatogenic fungi *Fusarium oxysporum* and *Alternaria* sp. were employed. The TLC plate containing spot of each substance was sprayed with spore suspension of those two mentioned fungi and incubated in a water agar for 3 days. The inhibition was observed by clear zone of active component against blue background.

When the TLC plate was sprayed with spore suspension of *Fusarium oxysporum*, all of isolated substances did not show any inhibition effect. That was noticed by no clear zone on the TLC plate as displayed in Figure 3.39.



Figure 3.39Bioautographic pattern of isolated substances againstFusarium oxysporum 43-68

Furthermore, when spore suspension of *Alternaria* sp. was sprayed on TLC plate, the spot of Mixture <u>6</u> displayed a clear zone which indicated the inhibition zone. The TLC plates are shown in Figure 3.40



Figure 3.40 Bioautographic pattern of isolated substances against *Alternaria* sp.43-89

According to preliminary result of antifungal activity, Mixture <u>6</u> (a mixture of alkenylresorcinol) exhibited inhibition zone on TLC plate when sprayed with spore suspension of *Alternaria* sp., while the other substances did not show inhibition zone. Therefore, it could be elementary proposed that the resorcinol moiety of Mixture <u>6</u> displayed as important portion responsible for this activity.

3.6.4 Insect Contact Toxicity Test

The contact toxicity against the larvae of *S. litura* by the vial test method was performed followed the procedure that described in Section 2.7.4. Two potential substances, Compound <u>1</u> and Mixture <u>6</u> containing phenolic lipid derivatives were investigated. The dead larvae amount after 5 days is reported as shown in Table 3.27.

The correlation curve between the concentration and the percentage of died larvae were plotted, and the results are exhibited in Figure 3.41. The LD_{50} of each sample was calculated by probit analysis program, and the LD_{50} results are demonstrated in Table 3.28.

Concentration	Compo	ound <u>1</u>	Mixt	ure <u>6</u>
(µg/mL)	Died larvae	ed larvae Percentage		Percentage
1000	24	53	-	-
500	23	51	45	100
100	15	33	43	96
50	11	24	26	58
10	14	31	15	33

 Table 3.27
 The died larvae (S. litura) after treating with Compound 1 and Mixture 6



Figure 3.41 Insecticidal activity (contact toxicity vial test) against S. litura

Table 3.28The LD50 of Compound 1 and Mixture 6 against S. litura by contact
toxicity vial test

Substance	LD ₅₀ (µg/mL)
Compound <u>1</u>	746.73
Mixture <u>6</u>	33.6

The LD₅₀ of Mixture <u>6</u> is significantly less than that of Compound <u>1</u> about 80 times. Mixture <u>6</u> displayed higher insect toxicity than Compound <u>1</u>. Considering the structure of both substances, it could be uncomplicatedly recommended that the resorcinol moiety or the unsaturated aliphatic side chain may be the active part of Mixture <u>6</u>.

3.6.5 Anticancer Activity

As aforementioned, phenolic lipid derivatives (Compound <u>1</u> and Mixture <u>6</u>) are known to possessed cytotoxic activity against brine shrimp, the anticancer activity was selected for further investigation. The anticancer activity against human mouth carcinoma (KB), breast cancer (BC), and small cell lung cancer (NCI-H187) was performed by Bioassay Laboratory, National Center for Genetic Engineering and Biotechnology using microculture tatrazolium assay. This assay utilized the maximum concentration at 20 μ g/mL. The cytotoxicity results are deduced as shown in Table 3.29.

Table 3.29The result	s of	anticancer	activity
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	Cancer Cell							
Substances	КВ		BC		NCI-H187			
	$IC_{50}\mu g/mL$	Acitvity	$IC_{50}\mu g/mL$	Activity	$IC_{50}\mu g/mL$	Activity		
Compound <u>1</u>	F	Inactive	2.03	Strong	0.30	Strong		
Mixture <u>6</u>		Inactive	1.95	Strong	3.30	Strong		

Note: >20 µg/mL; Inactive, 10-20 µg/mL; Weak, 5-10 µg/mL; Moderate,

<5 µg/mL; Strong

As the results presented in Table 3.29, it was found that both Compound $\underline{1}$ and Mixture <u>6</u> demonstrated potent cytotoxic against the breast cancer (BC) and the small cell lung cancer (NCI-H187), whereas they were inactive for the human mouth carcinoma (KB). The presence of *o*-hydroxybenzoquinone nucleus of Compound <u>1</u> and resorcinol moiety of Mixture <u>6</u> might be important portion for expressing this activity.

The antifungal and insecticidal activities result indicated that Mixture $\underline{6}$ or a mixture of alkenylresorcinol showed positive result for both assays. The other biological activity of resorcinolic lipid derivative was additionally literated from previous reports. In 1999 Kozubek and Tyman reviewed the biological activity of resorcinolic lipid.²⁹ Antibacterial activity is one of activities that have been reported. Recent experiment indicated that the extracts from Gingko biloba fruit, Ardisia *japonica* plant and seed cover of *Myristica fragrans* or cashew nut shell liquid (CNSL) which contained resorcinolic lipid derivative as a major component exhibited high activity toward phatogenic Gram-positive bacteria, Mycobacterium smegmatis and Mycobacterium tuberculosis as well as phytophatogenic bacteria. Similarly to their antibacterial activity, resorcinolic lipid displayed antifungal activity that inhibited the growth of Trichophyton mentagrophytes and Saccharomyces cerevisiae. The resorcinolic lipids are nontoxic to higher animals, it was found by the toleration by rats with an oral intake of 5 g/kg. This result was accordingly led to other applications such as the treatment of mouth and gingival infections, antifungal fluids and also in hair restoration lotion preparations. Furthermore, the biological activity of this substance as the growth regulators and in host-parasite relationship, effect of resorcinolic lipids on nucleic acid, interaction with proteins and effects on enzymatic activity, contact dermatitis, interaction of resorcinolic lipids with phospholipids bilayer biological membranes and the modulators of lipid oxidation have been addressed.

Moreover, as a collaboration work with Dr. Masanori Morimoto from Department of Agricultural Chemistry, Kinki University, Nakamachi, Nara, Japan, in some biological activity tests on Compound <u>1</u> or embelin, a major component of *Ardisia colorata* Roxb. fruits were conducted. The bioassay studied at Kinki University included antifeedant activity, mitochondria respiration inhibitor, antimicrobial activity, and the inhibitory enzymatic activity.

The insect antifeedant activity of Compound <u>1</u> using leaf disk bioassay³⁰ against *Spodoptera litura* gave the ED₅₀ of 0.585 mg/disk (moderate activity).

For the antimicrobial activity against *Bacillus subtilis*, the gram-positive bacteria that normally considered to be non-pathogenic, Compound <u>1</u> showed complete inhibition at 31.5 ppm.

In the case of the inhibitory enzymatic activity on p-hydroxypyruvate dioxygenase or HPPD enzyme, the inhibition of HPPD enzyme will prevent the

accumulation of toxic fumarylacetoacetate and succinylacetone in the liver, Compound <u>1</u> exhibited IC₅₀ value of 24.5 μ g/mg protein.

For the mitochondria respiration inhibitor test, mitochondria are essential to the creation of ATP. In general, the mitochondria is an important organelle in respiration system. In this experiment, Compound <u>1</u> displayed as electron transport inhibitory on succinic acid to H₂O was tested. Compound <u>1</u> demonstrated the weak mitochondria inhibition, and showed IC_{50} at 3.56×10^{-4} M.

The biological activity result of isolated substances indicated that the active components of the dichloromethane extract from *A. colorata* fruit are Compound <u>1</u> or embelin and Mixture <u>6</u>. These two substances could be classified as phenolic lipid type derivative. Compound <u>1</u> (embelin) exhibited the highly cytotoxic activity against brine shrimp at LC₅₀ 1.72 µg/mL, and the extremely antioxidant against DPPH radical. Mixture <u>6</u> revealed the moderately cytotoxic activity against brine shrimp at LC₅₀ 32.36 µg/mL, but it showed high activity on the growth inhibitory effect of fungi *Alternaria* sp. and the insecticidal activity against *S. litura*. Furthermore, Compound <u>1</u> and Mixture <u>6</u> exhibited strongly cytotoxic activity against the breast cancer cell at IC₅₀ 2.03 µg/mL and 1.95 µg/mL, and also showed powerfully cytotoxic activity against the small cell lung cancer at IC₅₀ 0.30 µg/mL and 3.30 µg/mL, respectively.

Compound $\underline{7}$ or linoleic acid belonged to the unsaturated long chain fatty acid demonstrated the highly cytotoxic activity against brine shrimp at LC₅₀ 4.99 µg/mL. However, no other activity was observed. The other substances, Mixtures $\underline{3}$ and $\underline{4}$ and Compound $\underline{5}$ that categorized in triterpenoid and steroid type illustrated low cytotoxicity against brine shrimp.

CHAPTER IV

CONCLUSION

During the course of this research, with the aim to study chemical constituents and their bioactivities from Thai medicinal plants, it was found that the dichloromethane extract of *Ardisia colorata* Roxb. fruits, belonging to Mysinaceae family, showed highly cytotoxic activity against *Artemia salina* (brine shrimp) and exhibited significant radical scavenging effect on DPPH radical. After fractionation and purification, three pure compounds and four mixtures were obtained. All isolated substances were further elucidated by means of their physical properties, chemical reactions and spectroscopic evidences. The structures of all isolated substances are summarized as shown in Table 4.1.

Table 4.1Isolated substances of dichloromethane extract from *Ardisia colorata* Roxb. fruits

Substance	Weight	% yield*	Structure	Remarks
Compound <u>1</u>: 2,5-dihydroxy-3-undecyl-1,4-benzoquinone (embelin)	9.33 g	0.19	но	Orange plate solid
Mixture <u>2</u> :			CH ₃ -(CH ₂) _n -OH	Bright white plate
Mixture of long chain alcohols	21.5 mg	0.0245		solid
Mixture <u>3</u> : Stigmasteryl-3- <i>O</i> -palmitate	2.5 g	2.85		Yellow semisolid
and				
β-Sitosteryl-3- <i>O</i> -plamitate	สถา			
จพ	าลง	กรณ์	มหาวิทยาลย	

Table 4.1	(Continued)
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Substance	Weight	% yield*	Structure	Remarks
Mixture <u>4</u>: α-amyrin	56.9 mg	0.0679	$ \underbrace{ H } (f) = f) $	White needle crystal
Compound <u>5</u> : Stigmasterol	36.9 mg	0.0421	+ 0	Off-white needle crystal
	ล์ถา	บันวิเ	ายบริการ	

Table 4.1 (Continued)

Substance	Weight	% yield*	Structure	Remarks
Mixture <u>6</u> 1,3-dihydroxy-5-(heptadec-8,11- dienyl)benzene and	808.5 mg	0.9217	HO	Brown oil
1,3-dihydroxy-5-(pentadec-8-enyl) benzene			HO CHI	
Compound <u>7</u>: Octadeca-9,12-dienoic acid	19.20 mg	0.0219	HO	Pale-yellow oil

*The percentage yield of all isolated substances was calculated based on crude CH_2Cl_2 extract (114.20 g), while Compound <u>1</u> was calculated based on dried fruit materials (5 kg).

All isolated substances were further bioassayed. The bioactivity assay that selected for determination in this research are as follows; cytotoxicity test against brine shrimp, scavenging effect on DPPH radical, antifungal (plant phatogenic) activity, insecticidal activity (contact toxicity, vial test against *S. litura*) and anticancer activity. Compound <u>1</u> (embelin), the orange dihydroxybenzoquinone pigment obtained as a major component, exhibited high cytotoxic activity against brine shrimp, the breast cancer and the small cell lung cancer as well as the highest scavenging activity toward DPPH radical, and moderate toxicity against the common cutworm *S. litura*. While this compound was subjected to antifungal activity, it displayed inactive for growth inhibitor of *Fusarium oxysporum* and *Alternaria* sp.

Previous report on the chemical constituent determination of *A. colorata* fruit resulted that the major orange pigment was rapanone. Nevertheless, in this course of research endorsed by various apectroscopic evidences, the major orange pigment was obviously identified as embelin.

Concerning with bioactivities of other substances, Mixture <u>6</u> or a mixture of alkenylresorcinol exhibited moderate toxicity and strong cytotoxic activity against the breast cancer and the small cell lung cancer. It also showed significant activity on antifungal against *Alternaria* sp. Compound <u>7</u> or linoleic acid displayed high toxicity against brine shrimp while it showed inactive for other assays tested. The surplus substances revealed insignificant bioactivities result.

The occurrence of linoleic acid clearly showed the possible biosynthesis pathway for Mixture $6A^{30}$ or 1,3-dihydroxy-5-(heptadec-8,11-dienyl)benzene. The unsaturated side chain moiety of Mixture 6A which had non-conjugated two double bonds should be arose by the decarboxylation of linoleic acid and then attached to position 3 of resorcinol moiety.

The antifungal activity and insecticidal activity of chemical constituents from the dichloromethane extract from *A. colorata* fruit was addressed for the first time in chemical literature.

In conclusion, it could be clearly seen that the fruits of *Ardisia colorata* Roxb. could be used as a good source of embelin, a bioactive benzoquinone pigment. Moreover, this plant produced resorcinol substance in good yield.

Proposal for The Future Work

The incidence of embelin as a major constituent, one of an active ingredient and the outcome from this research opened many possibilities to carry on for further exploration. For example, the utilization of embelin as a cytotoxic agent against various cell lines. The possible future work on structure activity relationship of embelin and other substances may provide an opportunity to understand what parts of the molecule have an influence for those interest activity. For instance, the most importantly active part of embelin was o-hydroxybenzoquinone moiety, the exploration of structure activity relationship of this compound should be studied on the variation of number of carbon atom and degree of unsaturation in side chain and substituted group on *o*-hydroxybenzoquinone moiety. Another aspect that would make this research fulfill is the chemical constituents and biological activity investigations of ethyl acetate and methanol crude extracts from the dried fruits, and other parts of Ardisia colorata Roxb. This would provide informative data for the chemotaxonomy of this plant. This present investigation is one of excellent examples to endorse the concept of the necessity on studying chemical constituents in conjunction with biological activity which will certainly be a gateway for disclosing lead bioactive compounds for specific proposes.

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APPENDIX

Probit analysis (brine shrimp) of dichloromethane extract from A.colorata fruit

* * * * * * * * * * * * PROBIT ANALYSIS * * * * * * * * * * * * *

Parameter estimates converged after 19 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .03594 .01687 2.12985

Intercept Standard Error Intercept/S.E.

-.47802 .19253 -2.48284

Pearson Goodness-of-Fit Chi Square = .114 DF = 2 P = .945

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of Observed Expected VAR00001 Subjects Responses Responses Residual Prob

| 1000.00 | 30.0 | 30.0 | 30.000 | .000 | 1.00000 |
|---------|------|------|--------|---------|---------|
| 100.00 | 30.0 | 30.0 | 29.973 | .027 | .99908 |
| 10.00 | 30.0 | 13.0 | 13.584 | 584 | .45279 |
| .00 | 30.0 | 10.0 | 9.490 | .510 .: | 31632 |

Confidence Limits for Effective VAR00001

| | 9 | 5% Confidence | ce Limits | |
|------|-----------|-------------------------|-----------|--|
| Prob | VAR00001 | Lower | Upper | |
| .01 | -51.42761 | -714.02422 | -23.18061 | |
| .02 | -43.84287 | -619.02806 | -19.11494 | |
| .03 | -39.03059 | -558.77621 | -16.51524 | |
| .04 | -35.41050 | -513.46537 | -14.54530 | |
| .05 | -32.46584 | -476.62021 | -12.93120 | |
| .06 | -29.95947 | -445.26953 | -11.54700 | |
| .07 | -27.76187 | -417.79070 | -10.32375 | |
| .08 | -25.79418 | -393.19586 | -9.21930 | |
| .09 | -24.00465 | -370.83684 | -8.20584 | |
| .10 | -22.35738 | -350.26433 | -7.26395 | |
| .15 | -15.53726 | -265.22331 | -3.22959 | |
| .20 | -10.11684 | -197.91892 | .26030 | |
| .25 | -5.46660 | -140.65421 | 3.73078 | |
| .30 | -1.29055 | -90.23857 | 7.85723 | |
| .35 | 2.57919 | -46.41435 | 14.57437 | |
| .40 | 6.25119 | -14.95507 | 31.07391 | |
| .45 | 9.80390 | -1. <mark>4</mark> 4194 | 63.96149 | |
| .50 | 13.30028 | 4.18560 | 103.99899 | |
| .55 | 16.79666 | 7.67931 | 146.17032 | |
| .60 | 20.34936 | 10.45487 | 189.79546 | |
| .65 | 24.02137 | 12.9 <mark>5</mark> 749 | 235.25166 | |
| .70 | 27.89111 | 15.3 <mark>8708</mark> | 283.36344 | |
| .75 | 32.06716 | 17.87396 | 335.41864 | |
| .80 | 36.71740 | 20.54494 | 393.48285 | |
| .85 | 42.13782 | 23.57827 | 461.24381 | |
| .90 | 48.95794 | 27.32006 | 546.57740 | |
| .91 | 50.60521 | 28.21516 | 567.19670 | |
| .92 | 52.39474 | 29.18459 | 589.59974 | |
| .93 | 54.36243 | 30.24735 | 614.23627 | |
| .94 | 56.56002 | 31.43080 | 641.75490 | |
| .95 | 59.06639 | 32.77661 | 673.14397 | |
| .96 | 62.01106 | 34.35315 | 710.02669 | |
| .97 | 65.63115 | 36.28550 | 755.37512 | |
| .98 | 70.44342 | 38.84606 | 815.66612 | |
| .99 | 78.02817 | 42.86708 | 910.70693 | |
| | | | | |
| | | | | |
| | | | | |

Probit analysis (brine shrimp) of ethyl acetate extract from A.colorata fruit

Parameter estimates converged after 12 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .00689 .00299 2.30258

Intercept Standard Error Intercept/S.E.

-.42568 .17554 -2.42494

Pearson Goodness-of-Fit Chi Square = .370 DF = 2 P = .831

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of Observed Expected VAR00001 Subjects Responses Responses Residual Prob

| 1000.00 | 30.0 | 30.0 | 30.000 | 1.4954E-09 | 1.00000 |
|---------|------|------|--------|-------------|---------|
| 100.00 | 30.0 | 18.0 | 18.119 | 119 .60 |)397 |
| 10.00 | 30.0 | 12.0 | 10.819 | 1.181 .36 | 064 |
| .00 | 30.0 | 9.0 | 10.055 | -1.055 .335 | 17 |

Confidence Limits for Effective VAR00001

| Prob VAR00001 Lower Upper .01 -275.73480 -2070.13544 -128.97750 .02 -236.18814 -1804.72630 -107.21648 .03 -211.09704 -1636.39808 -93.34444 .04 -192.22199 -1509.81681 -82.86355 .05 -176.86860 -1406.88942 -74.30144 .06 -163.80044 -1319.31401 -66.98179 .07 -152.34223 -1242.55670 -60.53468 .08 -142.08277 -1173.85725 -54.73451 .09 -132.75219 -1111.40451 -49.43282 .00 -124.16339 -1053.94295 -44.52631 .15 -88.60343 -816.41558 -23.83323 .20 -60.3151 -628.38063 -6.64275 .25 -36.09530 -468.17573 9.21749 .30 -14.32145 -326.28751 25.4121 .35 5.85527 -199.13301 44.80099 .40 25.00100 -90.32605 | | ç | 5% Confidenc | e Limits | |
|---|------|------------|-------------------------|------------|--|
| $\begin{array}{rrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrrr$ | Prob | VAR00001 | Lower | Upper | |
| $\begin{array}{cccccccccccccccccccccccccccccccccccc$ | .01 | -275.73480 | -2070.13544 | -128.97750 | |
| 03 -211.09704 -1636.39808 -93.34444 04 -192.22199 -1509.81681 -82.86355 05 -176.86860 -1406.88942 -74.30144 06 -163.80044 -1319.31401 -66.98179 07 -152.34223 -1242.55670 -60.53468 08 -142.08277 -1173.85725 -54.73451 09 -32.75219 -1111.40451 -49.43282 10 -124.16339 -1053.94295 -44.52631 15 -88.60343 -816.41558 -23.83323 20 -60.34151 -628.38063 -6.64275 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 517 79.9478 41.40057 342.56267 60 98.50849 56.71072 461.76345 <t< td=""><td>02</td><td>-236 18814</td><td>-1804 72630</td><td>-107 21648</td><td></td></t<> | 02 | -236 18814 | -1804 72630 | -107 21648 | |
| 04 -192.22199 -1509.81681 -82.86355 05 -176.86860 -1406.88942 -74.30144 06 -163.80044 -1319.31401 -66.98179 07 -152.34223 -1242.55670 -60.53468 08 -142.08277 -1173.85725 -54.73451 09 -132.75219 -1111.40451 -49.43282 10 -124.16339 -1053.94295 -44.52631 15 -88.60343 -816.41558 -23.83323 20 -60.34151 -628.38063 -6.64275 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.08751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 55 79.98478 41.40057 342.56267 60 98.50849 56.71072 461.76345 65 117.65423 70.17032 587.31157 | 03 | -211 09704 | -1636 39808 | -93 34444 | |
| 176.176.86860 -1406.88942 -74.30144 06 -163.80044 -1319.31401 -66.98179 07 -152.34223 -1242.55670 -60.53468 08 -142.08277 -1173.85725 -54.73451 09 -132.75219 -1111.40451 -49.43282 10 -124.16339 -105.94295 -44.52631 15 -88.60343 -816.41558 -23.83323 20 -60.34151 -628.38063 -6.64275 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 .40 25.00100 -90.32605 75.02174 .45 43.52472 -16.02898 135.23560 .50 61.75475 20.80164 230.78329 .55 79.98478 41.40057 342.56267 .60 98.50849 56.71072 461.76345 .65 117.65423 70.17032 587.33157 .70 137.83094 83.13328 720.88289 .75 | .00 | -192 22199 | -1509 81681 | -82 86355 | |
| 106 -163.80044 -1319.31401 -66.98179 07 -152.34223 -1242.55670 -60.53468 08 -142.08277 -1173.85725 -54.73451 09 -132.75219 -1111.40451 -49.43282 10 -124.16339 -1053.94295 -44.52631 15 -88.60343 -816.41558 -23.83323 20 -60.34151 -628.38063 -6.64275 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 .45 43.52472 -16.02898 135.23560 .50 61.75475 20.80164 230.78329 .55 79.98478 41.40057 342.56267 .60 98.50849 56.71072 461.76345 .65 117.65423 70.17032 587.33157 .70 137.83094 83.13328 720.82829 .75 159.60479 96.38341 865.74469 . | 05 | -176 86860 | -1406 88942 | -74 30144 | |
| $\begin{array}{cccccccccccccccccccccccccccccccccccc$ | .06 | -163.80044 | -1319.31401 | -66.98179 | |
| $\begin{array}{cccccccccccccccccccccccccccccccccccc$ | .07 | -152.34223 | -1242.55670 | -60.53468 | |
| 09 -132.75219 -1111.40451 -49.43282 10 -124.16339 -1053.94295 -44.52631 15 -88.60343 -816.41558 -23.83323 20 -60.34151 -628.38063 -6.64275 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 55 79.98478 41.40057 342.56267 60 98.50849 56.71072 461.76345 65 117.65423 70.17032 587.33157 70 137.83094 83.13328 720.88289 75 159.60479 96.38341 865.74469 80 183.85101 110.62935 1027.56390 85 212.11293 126.83892 1216.57976 90 247.67288 146.87823 1454.76089 91 256.26169 151.67826 1574.88236 93 275.85173 162.58362 1643.67762 94 287.30994 168.93886 1720.52680 95 300.37809 176.16952 1808.19120 96 315.73149 184.64417 1911.20605 97 334.60654 195.03717 2037.87521 98 359.69763 208.81729 2206.2 | .08 | -142.08277 | -1173.85725 | -54.73451 | |
| 10 -124.16339 -1053.94295 -44.52631 15 -88.60343 -816.41558 -23.83323 20 -60.34151 -628.38063 -6.64275 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 55 79.98478 41.40057 342.56267 .60 98.50849 56.71072 461.76345 .65 117.65423 70.17032 587.33157 .70 137.83094 83.13328 720.88289 .75 159.60479 96.38341 865.74469 .80 183.85101 110.62935 1027.56390 .85 212.11293 126.83892 1216.57976 .90 247.67288 146.87823 1454.76089 .91 256.59226 156.87926 1574.88236 .93 275.85173 162.58362 1643.67762 .94 287.30994 168.93886 1720.52680 .95 300.37809 176.16952 1808.19120 .96 315.73149 184.64417 1911.20605 .97 334.60654 195.03717 2037.87521 .98 359.69763 208.81729 2206.29535 .99 399.24430 230.47297 2471.80983 | .09 | -132.75219 | -1111.40451 | -49.43282 | |
| 15 -88.60343 -816.41558 -23.83323 20 -60.34151 -628.38063 -6.64275 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 55 79.98478 41.40057 342.56267 60 98.50849 56.71072 461.76345 65 117.65423 70.17032 587.33157 70 137.83094 83.13328 720.88289 75 159.60479 96.38341 865.74469 80 183.85101 110.62935 1027.56390 85 212.11293 126.83892 1216.57976 90 247.67288 146.87823 1454.76089 91 256.26169 151.67826 1574.88236 92 265.59226 156.87926 1574.88236 93 275.85173 162.58362 1643.67762 94 287.30994 168.93886 1720.52680 95 300.37809 176.16952 1808.19120 96 315.73149 184.64417 1911.20605 97 334.60654 195.03717 2037.87521 98 359.69763 208.81729 2206.29535 99 399.24430 230.47297 2471.80983 | .10 | -124.16339 | -1053.94295 | -44.52631 | |
| 20 -60.34151 -628.38063 -6.64275 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 55 79.98478 41.40057 342.56267 60 98.50849 56.71072 461.76345 65 117.65423 70.17032 587.33157 70 137.83094 83.13328 720.88289 75 159.60479 96.38341 865.74469 80 183.85101 110.62935 1027.56390 85 212.11293 126.83892 1216.57976 90 247.67288 146.87823 1454.76089 91 256.26169 151.67826 1572.32893 92 265.59226 156.87926 1574.88236 93 275.85173 162.58362 1643.67762 94 287.30994 168.93886 1720.52680 95 300.37809 176.16952 1808.19120 96 315.73149 184.64417 1911.20605 97 334.60654 195.03717 2037.87521 98 359.69763 208.81729 2206.29535 99 399.24430 230.47297 2471.80983 | .15 | -88.60343 | -816.41558 | -23.83323 | |
| 25 -36.09530 -468.17573 9.21749 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 55 79.98478 41.40057 342.56267 60 98.50849 56.71072 461.76345 65 117.65423 70.17032 587.33157 70 137.83094 83.13328 720.88289 75 159.60479 96.38341 865.74469 80 183.85101 110.62935 1027.56390 85 212.11293 126.83892 1216.57976 90 247.67288 146.87823 1454.76089 91 256.26169 151.67826 1512.32893 92 265.59226 156.87926 1574.88236 93 275.85173 162.58362 1643.67762 94 287.30994 168.93886 1720.52680 95 300.37809 176.16952 1808.19120 96 315.73149 184.64417 1911.20605 97 334.60654 195.03717 2037.87521 98 359.69763 208.81729 2206.29535 99 399.24430 230.47297 2471.80983 | .20 | -60.34151 | -628.38063 | -6.64275 | |
| 30 -14.32145 -326.28751 25.44121 35 5.85527 -199.13301 44.80099 40 25.00100 -90.32605 75.02174 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 55 79.98478 41.40057 342.56267 60 98.50849 56.71072 461.76345 65 117.65423 70.17032 587.33157 70 137.83094 83.13328 720.88289 75 159.60479 96.38341 865.74469 80 183.85101 110.62935 1027.56390 85 212.11293 126.83892 1216.57976 90 247.67288 146.87823 1454.76089 91 256.26169 151.67826 1574.88236 93 275.85173 162.58362 1643.67762 94 287.30994 168.93886 1720.52680 95 300.37809 176.16952 1808.19120 96 315.73149 184.64417 1911.20605 97 334.60654 195.03717 2037.87521 98 359.69763 208.81729 2206.29535 99 399.24430 230.47297 2471.80983 | .25 | -36.09530 | -468,17573 | 9.21749 | |
| .355.85527.199.1330144.80099.4025.00100-90.3260575.02174.4543.52472-16.02898135.23560.5061.7547520.80164230.78329.5579.9847841.40057342.56267.6098.5084956.71072461.76345.65117.6542370.17032587.33157.70137.8309483.13328720.88289.75159.6047996.38341865.74469.80183.85101110.629351027.56390.85212.11293126.838921216.57976.90247.67288146.878231454.76089.91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .30 | -14.32145 | -326.28751 | 25.44121 | |
| .4025.00100-90.3260575.02174.4543.52472-16.02898135.23560.5061.7547520.80164230.78329.5579.9847841.40057342.56267.6098.5084956.71072461.76345.65117.6542370.17032587.33157.70137.8309483.13328720.88289.75159.6047996.38341865.74469.80183.85101110.629351027.56390.85212.11293126.838921216.57976.90247.67288146.878231454.76089.91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.93861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .35 | 5.85527 | -199.13301 | 44.80099 | |
| 45 43.52472 -16.02898 135.23560 50 61.75475 20.80164 230.78329 55 79.98478 41.40057 342.56267 60 98.50849 56.71072 461.76345 65 117.65423 70.17032 587.33157 70 137.83094 83.13328 720.88289 75 159.60479 96.38341 865.74469 80 183.85101 110.62935 1027.56390 85 212.11293 126.83892 1216.57976 90 247.67288 146.87823 1454.76089 91 256.26169 151.67826 1512.32893 92 265.59226 156.87926 1574.88236 93 275.85173 162.58362 1643.67762 94 287.30994 168.93886 1720.52680 95 300.37809 176.16952 1808.19120 96 315.73149 184.64417 1911.20605 97 334.60654 195.03717 2037.87521 98 359.69763 208.81729 2206.29535 99 399.24430 230.47297 2471.80983 | .40 | 25.00100 | -90.32605 | 75.02174 | |
| 50 61.75475 20.80164 230.78329 $.55$ 79.98478 41.40057 342.56267 $.60$ 98.50849 56.71072 461.76345 $.65$ 117.65423 70.17032 587.33157 $.70$ 137.83094 83.13328 720.88289 $.75$ 159.60479 96.38341 865.74469 $.80$ 183.85101 110.62935 1027.56390 $.85$ 212.11293 126.83892 1216.57976 $.90$ 247.67288 146.87823 1454.76089 $.91$ 256.26169 151.67826 1512.32893 $.92$ 265.59226 156.87926 1574.88236 $.93$ 275.85173 162.58362 1643.67762 $.94$ 287.30994 168.93886 1720.52680 $.95$ 300.37809 176.16952 1808.19120 $.96$ 315.73149 184.64417 1911.20605 $.97$ 334.60654 195.03717 2037.87521 $.98$ 359.69763 208.81729 2206.29535 $.99$ 399.24430 230.47297 2471.80983 | .45 | 43.52472 | -16.02898 | 135.23560 | |
| .55 79.98478 41.40057 342.56267 $.60$ 98.50849 56.71072 461.76345 $.65$ 117.65423 70.17032 587.33157 $.70$ 137.83094 83.13328 720.88289 $.75$ 159.60479 96.38341 865.74469 $.80$ 183.85101 110.62935 1027.56390 $.85$ 212.11293 126.83892 1216.57976 $.90$ 247.67288 146.87823 1454.76089 $.91$ 256.26169 151.67826 1512.32893 $.92$ 265.59226 156.87926 1574.88236 $.93$ 275.85173 162.58362 1643.67762 $.94$ 287.30994 168.93886 1720.52680 $.95$ 300.37809 176.16952 1808.19120 $.96$ 315.73149 184.64417 1911.20605 $.97$ 334.60654 195.03717 2037.87521 $.98$ 359.69763 208.81729 2206.29535 $.99$ 399.24430 230.47297 2471.80983 | .50 | 61.75475 | 20.80164 | 230.78329 | |
| .60 98.50849 56.71072 461.76345 .65 117.65423 70.17032 587.33157 .70 137.83094 83.13328 720.88289 .75 159.60479 96.38341 865.74469 .80 183.85101 110.62935 1027.56390 .85 212.11293 126.83892 1216.57976 .90 247.67288 146.87823 1454.76089 .91 256.26169 151.67826 1512.32893 .92 265.59226 156.87926 1574.88236 .93 275.85173 162.58362 1643.67762 .94 287.30994 168.93886 1720.52680 .95 300.37809 176.16952 1808.19120 .96 315.73149 184.64417 1911.20605 .97 334.60654 195.03717 2037.87521 .98 359.69763 208.81729 2206.29535 .99 399.24430 230.47297 2471.80983 | .55 | 79.98478 | 41.40057 | 342.56267 | |
| .65117.6542370.17032587.33157.70137.8309483.13328720.88289.75159.6047996.38341865.74469.80183.85101110.629351027.56390.85212.11293126.838921216.57976.90247.67288146.878231454.76089.91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .60 | 98.50849 | 56.71072 | 461.76345 | |
| .70137.8309483.13328720.88289.75159.6047996.38341865.74469.80183.85101110.629351027.56390.85212.11293126.838921216.57976.90247.67288146.878231454.76089.91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .65 | 117.65423 | 70.17032 | 587.33157 | |
| .75159.6047996.38341865.74469.80183.85101110.629351027.56390.85212.11293126.838921216.57976.90247.67288146.878231454.76089.91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .70 | 137.83094 | 83.1 <mark>3</mark> 328 | 720.88289 | |
| .80183.85101110.629351027.56390.85212.11293126.838921216.57976.90247.67288146.878231454.76089.91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .75 | 159.60479 | 96.38341 | 865.74469 | |
| .85212.11293126.838921216.57976.90247.67288146.878231454.76089.91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .80 | 183.85101 | 110.62935 | 1027.56390 | |
| .90247.67288146.878231454.76089.91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .85 | 212.11293 | 126.83892 | 1216.57976 | |
| .91256.26169151.678261512.32893.92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .90 | 247.67288 | 146.87823 | 1454.76089 | |
| .92265.59226156.879261574.88236.93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .91 | 256.26169 | 151.67826 | 1512.32893 | |
| .93275.85173162.583621643.67762.94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .92 | 265.59226 | 156.87926 | 1574.88236 | |
| .94287.30994168.938861720.52680.95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .93 | 275.85173 | 162.58362 | 1643.67762 | |
| .95300.37809176.169521808.19120.96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .94 | 287.30994 | 168.93886 | 1720.52680 | |
| .96315.73149184.644171911.20605.97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .95 | 300.37809 | 176.16952 | 1808.19120 | |
| .97334.60654195.037172037.87521.98359.69763208.817292206.29535.99399.24430230.472972471.80983 | .96 | 315.73149 | 184.64417 | 1911.20605 | |
| .98 359.69763 208.81729 2206.29535
.99 399.24430 230.47297 2471.80983 | .97 | 334.60654 | 195.03717 | 2037.87521 | |
| .99 399.24430 230.47297 2471.80983 | .98 | 359.69763 | 208.81729 | 2206.29535 | |
| | .99 | 399.24430 | 230.47297 | 2471.80983 | |
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Probit analysis (brine shrimp) of methanol extract from A.colorata fruit

* * * * * * * * * * * * PROBIT ANALYSIS * * * * * * * * * * * * *

Parameter estimates converged after 18 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .00459 .00153 3.01209

Intercept Standard Error Intercept/S.E.

-1.60776 .21486 -7.48277

Pearson Goodness-of-Fit Chi Square = .345 DF = 2 P = .842

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of Observed Expected VAR00001 Subjects Responses Responses Residual Prob 1000.00 29.958 30.0 30.0 .042 .99859 3.0 100.00 30.0 3.762 -.762 .12541 10.00 30.0 2.0 1.775 .225 .05917 .00 30.0 2.0 1.618 .382 .05394

Confidence Limits for Effective VAR00001

| | ç | 5% Confidence | e Limits | |
|------|------------|--------------------------|------------|--|
| Prob | VAR00001 | Lower | Upper | |
| .01 | -156.41324 | -584.14924 | -47.69094 | |
| .02 | -97.07766 | -421.81899 | -4.20030 | |
| .03 | -59.43114 | -321.69132 | 26.25878 | |
| .04 | -31.11113 | -248.98521 | 51.78800 | |
| .05 | -8.07498 | -192.43352 | 75.14318 | |
| .06 | 11.53240 | -146.906 <mark>87</mark> | 97.62982 | |
| .07 | 28.72423 | -109.5 <mark>7749</mark> | 119.93478 | |
| .08 | 44.11747 | -78.63855 | 142.39129 | |
| .09 | 58.11700 | -52.78684 | 165.10061 | |
| .10 | 71.00360 | - <mark>31.00374</mark> | 188.01801 | |
| .15 | 124.35754 | 40.63195 | 301.45434 | |
| .20 | 166.76156 | 82.83411 | 406.34156 | |
| .25 | 203.14043 | 113.59304 | 501.77222 | |
| .30 | 235.80979 | 138.78415 | 589.90325 | |
| .35 | 266.08281 | 160.83745 | 672.85981 | |
| .40 | 294.80895 | 180.98608 | 752.35527 | |
| .45 | 322.60182 | 199.96427 | 829.78389 | |
| .50 | 349.95405 | 218.27318 | 906.35330 | |
| .55 | 377.30628 | 236.30260 | 983.20219 | |
| .60 | 405.09915 | 254.39912 | 1061.51248 | |
| .65 | 433.82529 | 272.9 <mark>1</mark> 613 | 1142.63955 | |
| .70 | 464.09832 | 292. <mark>265</mark> 99 | 1228.29956 | |
| .75 | 496.76767 | 312.99607 | 1320.89163 | |
| .80 | 533.14654 | 335.93213 | 1424.14516 | |
| .85 | 575.55057 | 362.51119 | 1544.65547 | |
| .90 | 628.90451 | 395.76919 | 1696.46949 | |
| .91 | 641.79110 | 403.77677 | 1733.16241 | |
| .92 | 655.79063 | 412.46626 | 1773.03395 | |
| .93 | 671.18387 | 422.00997 | 1816.88568 | |
| .94 | 688.37571 | 432.65636 | 1865.87365 | |
| .95 | 707.98308 | 444.78386 | 1921.75944 | |
| .96 | 731.01923 | 459.01376 | 1987.43640 | |
| .97 | 759.33924 | 476.48316 | 2068.20232 | |
| .98 | 796.98577 | 499.66888 | 2175.60336 | |
| .99 | 856.32135 | 536.14039 | 2344.95275 | |
| | | | | |
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Probit analysis (brine shrimp) of Compound 1

>Warning # 13527>Parameter estimates did not converge in maximum number of iterations.

Number of iterations = 20Optimal solution not found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .49095 .89505 .54852

Intercept Standard Error Intercept/S.E.

-.84316 .26094 -3.23122

Pearson Goodness-of-Fit Chi Square = .001 DF = 2 P = 1.000

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of Observed Expected VAR00001 Subjects Responses Responses Residual Prob

| 1000.00 | 30.0 | 30.0 | 30.000 | .000 1.00000 |
|---------|------|------|--------|--------------|
| 100.00 | 30.0 | 30.0 | 30.000 | .000 1.00000 |
| 10.00 | 30.0 | 30.0 | 29.999 | .001 .99998 |
| .00 | 30.0 | 6.0 | 5.987 | .013 .19957 |

จุฬาลงกรณมหาวทยาลย

* * * * * * * * * * * P R O B I T A N A L Y S I S * * * * * * * * * * * * *

| | 95% | 6 Confidence | Limits | | |
|------|----------|-----------------|--------|--|--|
| Prob | VAR00001 | Lower | Upper | | |
| | | | | | |
| .01 | -3.02105 | | | | |
| .02 | -2.46580 | | | | |
| .03 | -2.11352 | | | | |
| .04 | -1.84851 | | | | |
| .05 | -1.63294 | | | | |
| .06 | -1.44946 | | | | |
| .07 | -1.28858 | | | | |
| .08 | -1.14454 | | | | |
| .09 | -1.01354 | | | | |
| .10 | 89295 | | | | |
| .15 | 39367 | / | | | |
| .20 | .00313 | | | | |
| .25 | .34355 | | | | |
| .30 | .64927 | • • • • • | | | |
| .35 | .93255 | | | | |
| .40 | 1.20136 | •/ // // 8 | | | |
| .45 | 1.46144 | . // / .~ | | | |
| .50 | 1.71740 | | | | |
| .55 | 1.97335 | · / / / · · · · | | | |
| .60 | 2.23343 | | | | |
| .65 | 2.50224 | | | | |
| .70 | 2.78553 | | | | |
| .75 | 3.09124 | · | | | |
| .80 | 3.43166 | | | | |
| .85 | 3.82847 | | | | |
| .90 | 4.32774 | | | | |
| .91 | 4.44833 | | | | |
| .92 | 4.57933 | | | | |
| .93 | 4.72338 | | | | |
| .94 | 4.88425 | | | | |
| .95 | 5.06774 | | | | |
| .96 | 5.28330 | 79 19 14 | | | |
| .97 | 5.54831 | ·IU M | | | |
| .98 | 5.90060 | | | | |
| .99 | 6.45584 | | | | |
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Probit analysis (brine shrimp) of Mixture 2

Parameter estimates converged after 12 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .00070 .00029 2.38385

Intercept Standard Error Intercept/S.E.

-1.03937 .16634 -6.24859

Pearson Goodness-of-Fit Chi Square = .118 DF = 2 P = .943

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of Observed Expected VAR00001 Subjects Responses Responses Residual Prob 1000.00 30.0 11.0 11.007 -.007 .36690 100.00 30.0 5.0 4.985 .015 .16616 10.00 30.0 5.0 4.528 .472 .15095 .00 4.0 4.479 -.479 .14932 30.0

| | C | 95% Confidence | e Limits | |
|------|-------------|---------------------------|---------------------------|--|
| Prob | VAR00001 | Lower | Upper | |
| .01 | -1840.38413 | -11983.85760 | -816.37638 | |
| .02 | -1450.56560 | -9799.55977 | -594.27935 | |
| .03 | -1203.23824 | -8415.78934 | -451.26940 | |
| .04 | -1017.18351 | -7376.53566 | -3 <mark>41.98</mark> 567 | |
| .05 | -865.84233 | -6532.76338 | -251.51137 | |
| .06 | -737.02722 | -5816.15 <mark>712</mark> | -172.92704 | |
| .07 | -624.08157 | -5189. <mark>48319</mark> | -102.37440 | |
| .08 | -522.95221 | -4630.16026 | -37.41410 | |
| .09 | -430.97910 | -4123.47557 | 23.66185 | |
| .10 | -346.31771 | -3659.35826 | 82.16913 | |
| .15 | 4.20309 | -1788.46557 | 375.08187 | |
| .20 | 282.78591 | -515.79263 | 822.13289 | |
| .25 | 521.78516 | 100.82163 | 1680.88866 | |
| .30 | 736.41387 | 365.96120 | 2740.67699 | |
| .35 | 935.29937 | 533.93725 | 3800.44407 | |
| .40 | 1124.02225 | 668.33479 | 4831.05472 | |
| .45 | 1306.61382 | 787.57282 | 5838.97558 | |
| .50 | 1486.31047 | 899.23264 | 6836.60398 | |
| .55 | 1666.00712 | 1007.45678 | 7837.66806 | |
| .60 | 1848.59869 | 11 <mark>1</mark> 5.12917 | 8857.15456 | |
| .65 | 2037.32158 | 1224. <mark>75</mark> 578 | 9912.53614 | |
| .70 | 2236.20707 | 1338.99718 | 11026.03788 | |
| .75 | 2450.83578 | 1461.21506 | 12228.74789 | |
| .80 | 2689.83503 | 1596.36 <mark>507</mark> | 13568.96792 | |
| .85 | 2968.41785 | 1752.98857 | 15132.06838 | |
| .90 | 3318.93865 | 1949.06727 | 17099.79511 | |
| .91 | 3403.60005 | 1996.29918 | 17575.18779 | |
| .92 | 3495.57315 | 2047.56353 | 18091.68407 | |
| .93 | 3596.70251 | 2103.88004 | 18659.65080 | |
| .94 | 3709.64816 | 2166.71889 | 19294.03852 | |
| .95 | 3838.46327 | 2238.31968 | 20017.62831 | |
| .96 | 3989.80445 | 2322.35964 | 20867.83493 | |
| .97 | 4175.85919 | 2425.56971 | 21913.16227 | |
| .98 | 4423.18654 | 2562.61382 | 23302.89855 | |
| .99 | 4813.00507 | 2778.31773 | 25493.58949 | |
| | | | | |
| | | | | |
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Probit analysis (brine shrimp) of Mixture 3

Parameter estimates converged after 10 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .00138 .00031 4.42221

Intercept Standard Error Intercept/S.E.

-1.46642 .20124 -7.28677

Pearson Goodness-of-Fit Chi Square = .559 DF = 2 P = .756

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of
VAR00001Observed
SubjectsExpected
ResponsesResidualProb1000.0030.014.013.917.083.46388100.0030.02.02.759-.759.0919510.0030.03.02.195.805.07316

| 00 | 20.0 | 2.0 | 2 1 2 0 | 120 | 07107 |
|-----|------|-----|---------|-----|--------|
| .00 | 30.0 | 2.0 | 2.138 | 138 | .0/12/ |

| | ç | 5% Confidence | e Limits | |
|------|------------|---------------------------|------------|--|
| Prob | VAR00001 | Lower | Upper | |
| .01 | -625.05993 | -1517.23675 | -253.03731 | |
| .02 | -426.91570 | -1170.75000 | -106.35727 | |
| .03 | -301.19953 | -953.08202 | -11.12673 | |
| .04 | -206.62816 | -790.98050 | 62.15311 | |
| .05 | -129.70164 | -660.53497 | 123.17216 | |
| .06 | -64.22510 | -550.79749 | 176.40112 | |
| .07 | -6.81497 | -455.805 <mark>66</mark> | 224.29897 | |
| .08 | 44.58895 | -371.94205 | 268.37602 | |
| .09 | 91.33875 | -296.84236 | 309.63325 | |
| .10 | 134.37203 | -228.87389 | 348.77151 | |
| .15 | 312.54128 | 36.57611 | 526.77200 | |
| .20 | 454.14455 | 222.74675 | 693.04174 | |
| .25 | 575.62755 | 362.25991 | 855.89106 | |
| .30 | 684.72304 | 473.50914 | 1016.17268 | |
| .35 | 785.81627 | 567.56933 | 1173.72628 | |
| .40 | 881.74386 | 651.04218 | 1329.01032 | |
| .45 | 974.55491 | 727.97626 | 1483.07629 | |
| .50 | 1065.89448 | 801.03618 | 1637.35400 | |
| .55 | 1157.23405 | 872.16226 | 1793.56556 | |
| .60 | 1250.04510 | 942.95608 | 1953.77178 | |
| .65 | 1345.97269 | 1014. <mark>9</mark> 4349 | 2120.54126 | |
| .70 | 1447.06592 | 1089.81417 | 2297.28437 | |
| .75 | 1556.16141 | 1169.73547 | 2488.89393 | |
| .80 | 1677.64441 | 1257.91315 | 2703.07872 | |
| .85 | 1819.24768 | 1359.87035 | 2953.56191 | |
| .90 | 1997.41693 | 1487.22243 | 3269.66032 | |
| .91 | 2040.45021 | 1517.85860 | 3346.13088 | |
| .92 | 2087.20001 | 1551.09456 | 3429.25184 | |
| .93 | 2138.60393 | 1587.58819 | 3520.69887 | |
| .94 | 2196.01406 | 1628.28789 | 3622.88885 | |
| .95 | 2261.49060 | 1674.63819 | 3739.50499 | |
| .96 | 2338.41712 | 1729.01059 | 3876.59716 | |
| .97 | 2432.98849 | 1795.74528 | 4045.24384 | |
| .98 | 2558.70466 | 1884.29591 | 4269.59173 | |
| .99 | 2756.84889 | 2023.55350 | 4623.50093 | |
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Probit analysis (brine shrimp) of Compound 5

Parameter estimates converged after 10 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .00113 .00029 3.95835

Intercept Standard Error Intercept/S.E.

-.80130 .15271 -5.24718

Pearson Goodness-of-Fit Chi Square = .487 DF = 2 P = .784

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of Observed Expected VAR00001 Subjects Responses Responses Residual Prob 1000.00 30.0 19.0 18.862 .138 .62875 30.0 100.00 6.0 7.369 -1.369 .24563 10.00 30.0 7.0 6.443 .557 .21476 .00 7.0 6.344 30.0 .656 .21148

| | 9 | 5% Confidence | e Limits |
|------|-------------|--------------------------|------------|
| Prob | VAR00001 | Lower | Upper |
| 01 | 1240 70422 | 2004 04544 | 772 02052 |
| .01 | -1349.79433 | | -//2.02852 |
| .02 | | -2532.75401 | |
| .03 | -955.44087 | -2233.31279 | -499.88769 |
| .04 | -840.28462 | | -419.40503 |
| .05 | -/46.61390 | -1826.45825 | -353.42648 |
| .00 | | -10/1.82101 | -290.81210 |
| .07 | -596.97911 | -1536.65976 | -246.74701 |
| .08 | -534.38636 | -1416.04800 | -201.51068 |
| .09 | -4//.460/6 | -1306.75874 | -159.96773 |
| .10 | -425.06065 | -1206.56046 | -121.32472 |
| .15 | -208.11019 | -797.67528 | 44.63014 |
| .20 | -35.68485 | -484.5/464 | 188.39415 |
| .25 | 112.24074 | -232.43855 | 328.20725 |
| .30 | 245.08250 | -28.16119 | 475.91236 |
| .35 | 368.18018 | 135.70813 | 638.20727 |
| .40 | 484.98784 | 268.19373 | 815.21962 |
| .45 | 598.00061 | 379.51553 | 1003.34068 |
| .50 | 709.22160 | 478.01922 | 1199.53229 |
| .55 | 820.44259 | 569.41313 | 1402.83368 |
| .60 | 933.45535 | 657.56096 | 1614.12870 |
| .65 | 1050.26302 | 745. <mark>37</mark> 278 | 1835.81484 |
| .70 | 1173.36070 | 835 <mark>.47367</mark> | 2071.87817 |
| .75 | 1306.20245 | 930.78662 | 2328.54770 |
| .80 | 1454.12804 | 1035.30504 | 2615.97846 |
| .85 | 1626.55339 | 1155.65469 | 2952.49346 |
| .90 | 1843.50384 | 1305.55598 | 3377.43221 |
| .91 | 1895.90395 | 1341.57338 | 3480.25611 |
| .92 | 1952.82955 | 1380.63394 | 3592.02776 |
| .93 | 2015.42230 | 1423.50982 | 3714.99997 |
| .94 | 2085.32860 | 1471.31394 | 3852.42219 |
| .95 | 2165.05710 | 1525.74130 | 4009.24643 |
| .96 | 2258.72782 | 1589.57444 | 4193.60675 |
| .97 | 2373.88407 | 1667.90571 | 4420.39778 |
| .98 | 2526.96424 | 1771.82728 | 4722.08316 |

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Probit analysis (brine shrimp) of Mixture 6

Parameter estimates converged after 13 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .00805 .00306 2.63309

Intercept Standard Error Intercept/S.E.

-.26040 .17233 -1.51099

Pearson Goodness-of-Fit Chi Square = 1.144 DF = 2 P = .565

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of Observed Expected VAR00001 Subjects Responses Responses Residual Prob 1000.00 30.0 30.0 30.000 1.0303E-13 1.00000 100.00 30.0 21.0 21.207 -.207 .70689 10.00 30.0 15.0 12.858 2.142 .42861

.00 30.0 10.0 11.918 -1.918 .39728

| | ç | 5% Confidenc | e Limits | |
|------|------------|---------------------------|------------|--|
| Prob | VAR00001 | Lower | Upper | |
| .01 | -256.72814 | -1108.52756 | -129.77412 | |
| .02 | -222.85329 | -976.33625 | -110.02892 | |
| .03 | -201.36079 | -892.51302 | -97.45332 | |
| .04 | -185.19281 | -829.48842 | -87.96082 | |
| .05 | -172.04141 | -778.24821 | -80.21398 | |
| .06 | -160.84750 | -734.65 <mark>635</mark> | -73.59858 | |
| .07 | -151.03263 | -696.45 <mark>41</mark> 1 | -67.77888 | |
| .08 | -142.24459 | -662.26636 | -62.55023 | |
| .09 | -134.25222 | -631.19081 | -57.77815 | |
| .10 | -126.89523 | -602.60194 | -53.36921 | |
| .15 | -96.43531 | -484.45730 | -34.89412 | |
| .20 | -72.22673 | -390.95220 | -19.81818 | |
| .25 | -51.45793 | -311.23268 | -6.38484 | |
| .30 | -32.80690 | -240.35955 | 6.39617 | |
| .35 | -15.52395 | -175.84143 | 19.39609 | |
| .40 | .87589 | -116.72356 | 33.83525 | |
| .45 | 16.74292 | -63.79820 | 52.07717 | |
| .50 | 32.35838 | -20.37055 | 78.68848 | |
| .55 | 47.97384 | 10.24693 | 118.10995 | |
| .60 | 63.84087 | 3 <mark>0.7444</mark> 5 | 168.77971 | |
| .65 | 80.24071 | 4 <mark>6.2345</mark> 1 | 226.84668 | |
| .70 | 97.52366 | 59.7 <mark>4</mark> 947 | 290.84976 | |
| .75 | 116.17469 | 72.80932 | 361.44406 | |
| .80 | 136.94349 | 86.40817 | 440.99806 | |
| .85 | 161.15207 | 101.59042 | 534.39685 | |
| .90 | 191.61198 | 120.13902 | 652.46798 | |
| .91 | 198.96898 | 124.56014 | 681.04467 | |
| .92 | 206.96135 | 129.34381 | 712.10864 | |
| .93 | 215.74939 | 134.58353 | 746.28531 | |
| .94 | 225.56426 | 140.41389 | 784.47689 | |
| .95 | 236.75817 | 147.03966 | 828.05837 | |
| .96 | 249.90957 | 154.79674 | 879.28835 | |
| .97 | 266.07754 | 164.29957 | 942.30262 | |
| .98 | 287.57005 | 176.88603 | 1026.11500 | |
| .99 | 321.44490 | 196.64372 | 1158.29380 | |
| | | | | |
| | | | | |
| | | | | |

Probit analysis (brine shrimp) of Compound 7

Parameter estimates converged after 13 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .11353 .03565 3.18500

Intercept Standard Error Intercept/S.E.

-.16789 .23000 -.72997

Pearson Goodness-of-Fit Chi Square = 3.334E-10 DF = 2 P = 1.000

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number of Observed Expected VAR00001 Subjects Responses Responses Residual Prob 1000.00 30.0 30.0 30.000 .000 1.00000 100.00 30.0 30.0 30.000 .000 1.00000 10.00 30.0 25.0 25.000 .000 .83333 .00 13.0 13.000 30.0 .000 .43333

| | 9 | 5% Confiden | ce Limits | |
|------|-----------|-------------------------|-------------------------|--|
| Prob | VAR00001 | Lower | Upper | |
| .01 | -19.01191 | -56.41258 | -9.86288 | |
| .02 | -16.61083 | -50.20620 | -8.34007 | |
| .03 | -15.08742 | -46.27304 | -7.36931 | |
| .04 | -13.94142 | -43.31725 | -6. <mark>6360</mark> 7 | |
| .05 | -13.00924 | -40.91521 | -6.03737 | |
| .06 | -12.21580 | -38.872 <mark>56</mark> | -5.52592 | |
| .07 | -11.52012 | -37.08317 | -5.07585 | |
| .08 | -10.89721 | -35. <mark>48244</mark> | -4.67143 | |
| .09 | -10.33071 | -34.02797 | -4.30228 | |
| .10 | -9.80924 | -32.69039 | -3.96123 | |
| .15 | -7.65021 | -27.16843 | -2.53317 | |
| .20 | -5.93428 | -22.80527 | -1.37268 | |
| .25 | -4.46217 | -19.08970 | 34946 | |
| .30 | -3.14017 | -15.78571 | .60214 | |
| .35 | -1.91514 | -12.76555 | 1.52542 | |
| .40 | 75270 | -9.95554 | 2.45735 | |
| .45 | .37197 | -7.31598 | 3.43815 | |
| .50 | 1.47881 | - <mark>4.835</mark> 15 | 4.52029 | |
| .55 | 2.58565 | -2.52963 | 5.77773 | |
| .60 | 3.71032 | <mark>44038</mark> | 7.30885 | |
| .65 | 4.87275 | 1.39 <mark>30</mark> 2 | 9.21738 | |
| .70 | 6.09779 | 2.97776 | 11.57607 | |
| .75 | 7.41979 | 4.38324 | 14.42619 | |
| .80 | 8.89190 | 5.71215 | 17.83607 | |
| .85 | 10.60783 | 7.08233 | 21.98954 | |
| .90 | 12.76685 | 8.66055 | 27.36132 | |
| .91 | 13.28832 | 9.02685 | 28.67367 | |
| .92 | 13.85483 | 9.42006 | 30.10407 | |
| .93 | 14.47774 | 9.84756 | 31.68172 | |
| .94 | 15.17342 | 10.31991 | 33.44883 | |
| .95 | 15.96686 | 10.85310 | 35.46974 | |
| .96 | 16.89904 | 11.47330 | 37.85028 | |
| .97 | 18.04504 | 12.22829 | 40.78432 | |
| .98 | 19.56845 | 13.22195 | 44.69459 | |
| .99 | 21.96953 | 14.77118 | 50.87456 | |
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Probit analysis (insecticidal activity) of Compound 1

Parameter estimates converged after 8 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .00071 .00023 3.16759

Intercept Standard Error Intercept/S.E.

-.53391 .11637 -4.58810

Pearson Goodness-of-Fit Chi Square = 2.422 DF = 3 P = .490

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

Number ofObservedExpectedVAR00001SubjectsResponsesResponsesResidual1000.0045.024.025.7331.73357195

| 1000.00 | 45.0 | 24.0 | 25.733 | -1./33 | .5/185 |
|---------|------|------|--------|--------|--------|
| 500.00 | 45.0 | 23.0 | 19.349 | 3.651 | .42999 |
| 100.00 | 45.0 | 15.0 | 14.485 | .515 | .32189 |
| 50.00 | 45.0 | 11.0 | 13.913 | -2.913 | .30919 |
| 10.00 | 45.0 | 14.0 | 13.463 | .537 | .29918 |

| | 9 | 5% Confidence | e Limits | |
|------|-------------|-------------------------|-------------|--|
| Prob | VAR00001 | Lower | Upper | |
| .01 | -2506.93112 | -7155.18332 | -1400.23227 | |
| .02 | -2125.67060 | -6157.51347 | -1162.30375 | |
| .03 | -1883.77302 | -5524.94477 | -1010.92490 | |
| .04 | -1701.80290 | -5049.38638 | -896.75009 | |
| .05 | -1553.78424 | -4662.80092 | -803.63308 | |
| .06 | -1427.79712 | -4333.97214 | -724.15972 | |
| .07 | -1317.33105 | -4045.85389 | -654.27686 | |
| .08 | -1218.42187 | -3788.07006 | -591.51341 | |
| .09 | -1128.46793 | -3553.81397 | -534.24456 | |
| .10 | -1045.66517 | -3338.36877 | -481.34058 | |
| .15 | -702.83965 | -2449.16968 | -259.50237 | |
| .20 | -430.37279 | -1748.29830 | -77.35653 | |
| .25 | -196.62048 | -1156.51441 | 88.40974 | |
| .30 | 13.29631 | -643.57006 | 255.76885 | |
| .35 | 207.81551 | -209.79494 | 452.39627 | |
| .40 | 392.39520 | 116.87816 | 723.91352 | |
| .45 | 570.97819 | 335.08711 | 1084.46067 | |
| .50 | 746.72981 | 494.06568 | 1495.06223 | |
| .55 | 922.48143 | 629.19020 | 1929.51785 | |
| .60 | 1101.06442 | 75 <mark>5.51817</mark> | 2381.94597 | |
| .65 | 1285.64411 | 880.24325 | 2855.41125 | |
| .70 | 1480.16331 | 1008.13660 | 3357.92044 | |
| .75 | 1690.08010 | 1143.74837 | 3902.61212 | |
| .80 | 1923.83240 | 1292.95612 | 4510.95453 | |
| .85 | 2196.29926 | 1465.37682 | 5221.55106 | |
| .90 | 2539.12479 | 1680.89581 | 6117.06937 | |
| .91 | 2621.92754 | 1732.78362 | 6333.53073 | |
| .92 | 2711.88149 | 1789.09499 | 6568.74431 | |
| .93 | 2810.79067 | 1850.95051 | 6827.43607 | |
| .94 | 2921.25673 | 1919.96567 | 7116.42202 | |
| .95 | 3047.24386 | 1998.60121 | 7446.08861 | |
| .96 | 3195.26252 | 2090.89748 | 7833.49481 | |
| .97 | 3377.23264 | 2204.25014 | 8309.87536 | |
| .98 | 3619.13021 | 2354.77196 | 8943.30108 | |
| .99 | 4000.39073 | 2591.72193 | 9941.94948 | |
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Probit analysis (insecticidal activity) of Mixture 6

Parameter estimates converged after 14 iterations. Optimal solution found.

Parameter Estimates (PROBIT model: (PROBIT(p)) = Intercept + BX):

Regression Coeff. Standard Error Coeff./S.E.

VAR00001 .02252 .00385 5.84637

Intercept Standard Error Intercept/S.E.

-.75734 .20607 -3.67518

Pearson Goodness-of-Fit Chi Square = 1.515 DF = 2 P = .469

Since Goodness-of-Fit Chi square is NOT significant, no heterogeneity factor is used in the calculation of confidence limits.

Observed and Expected Frequencies

| Nun | nber of | Observed | Expecte | d | | |
|----------|----------|----------|-----------|---------|----------|------|
| VAR00001 | Subjects | Respor | nses Resp | onses F | Residual | Prob |
| 500.00 | 45.0 | 45.0 | 45.000 | | 1 00000 | |
| 500.00 | 45.0 | 45.0 | 45.000 | .000 | 1.00000 | |
| 100.00 | 45.0 | 43.0 | 41.964 | 1.036 | .93254 | |
| 50.00 | 45.0 | 26.0 | 28.974 | -2.974 | .64387 | |
| 10.00 | 45.0 | 15.0 | 13.380 | 1.620 | .29733 | |

| | ç | 5% Confidence | e Limits | |
|------|-----------|--------------------------|-----------|--|
| Prob | VAR00001 | Lower | Upper | |
| .01 | -69.66227 | -127.88879 | -39.91774 | |
| .02 | -57.55919 | -109.85618 | -30.67907 | |
| .03 | -49.88017 | -98.43710 | -24.79540 | |
| .04 | -44.10354 | -89.86124 | -20.35508 | |
| .05 | -39.40470 | -82.89624 | -16.73242 | |
| .06 | -35.40525 | -76.976 <mark>82</mark> | -13.64007 | |
| .07 | -31.89851 | -71.794 <mark>3</mark> 2 | -10.92102 | |
| .08 | -28.75865 | -67.16086 | -8.47958 | |
| .09 | -25.90307 | -62.95321 | -6.25290 | |
| .10 | -23.27450 | -59.08593 | -4.19736 | |
| .15 | -12.39154 | -43.14836 | 4.38712 | |
| .20 | -3.74210 | -30.59700 | 11.32510 | |
| .25 | 3.67835 | -19.94873 | 17.39696 | |
| .30 | 10.34214 | -10.52004 | 22.98346 | |
| .35 | 16.51714 | - <mark>1.9398</mark> 1 | 28.31703 | |
| .40 | 22.37660 | 6.01248 | 33.56759 | |
| .45 | 28.04571 | 13.47402 | 38.87995 | |
| .50 | 33.62493 | 20.53315 | 44.39220 | |
| .55 | 39.20415 | 27.25301 | 50.24372 | |
| .60 | 44.87325 | 33.69391 | 56.57673 | |
| .65 | 50.73272 | 39.93 <mark>57</mark> 3 | 63.53776 | |
| .70 | 56.90771 | 46.09708 | 71.29021 | |
| .75 | 63.57151 | 52.35117 | 80.05131 | |
| .80 | 70.99195 | 58.95204 | 90.17057 | |
| .85 | 79.64139 | 66.30914 | 102.30281 | |
| .90 | 90.52436 | 75.23234 | 117.90166 | |
| .91 | 93.15293 | 77.34862 | 121.70819 | |
| .92 | 96.00851 | 79.63437 | 125.85679 | |
| .93 | 99.14837 | 82.13352 | 130.43253 | |
| .94 | 102.65511 | 84.90935 | 135.55825 | |
| .95 | 106.65456 | 88.05810 | 141.42127 | |
| .96 | 111.35340 | 91.73759 | 148.32944 | |
| .97 | 117.13002 | 96.23646 | 156.84675 | |
| .98 | 124.80904 | 102.18298 | 168.20299 | |
| .99 | 136.91213 | 111.49578 | 186.16146 | |
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VITA

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