

## 제주특산 흑오미자 재배 및 대량증식기술과 특수성분 및 천연음료개발

Development of natural beverage and a special components and, methods of mass production and cultivation technique of Schisandra nigra as special-product of Cheju

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Development of natural beverage and a special components and, methods of mass production and cultivation technique of *Schisandra nigra* as special-product of Cheju

66

3

1999. 10.

가 가

가

가

가

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1)

2)

, 3) DNA

, 4)

, 5)

, 6) 가

가

가

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1.

## 2. RAPD

## RAPD

genetic marker

, DNA, ,  
DNA yield  $\gamma$  PCR,  
genomic DNA .

, PCR = template DNA, Mg<sup>2+</sup>, *Taq* DNA polymerase, dNTP annealing cycle band pattern 가 RAPD

, 가 Callus

가 가

, marker marker  
, marker cloning kit 가

3.

,  
,  
, , , citral  
, , , total phenol  
phenolic acid , ( )  
, lignan schizandrin,

4.

,  
,  
,  
( , ) ,

## A.

1.

, 가 가

600 1,350m

가 35.

, , 2  
, 400m<sup>2</sup> 3 39 . 3,000 3,094 mm  
, 8.8 10.5 0.6  
1,300m - 7 가  
가 . 50  
5 6  
가 , 1:6.3 .  
19.2 19.8 106.2 ,  
9.3 11.1 가 0.99g, 722 795  
, IBA 500ppm  
80% 가 , 63 92%  
WPM BAP GA<sub>3</sub> 1.0  
mg/ , BAP 3.0mg/ 가 3

가 200

, 가

가 ,

## 2. RAPD

genomic DNA

Roggers Bendish (1988), Dellaporta (1983)

genomic DNA ,

DNA

가

가 . RAPD southern hybridization

marker OPA - 17 primer 770 bp, OPB - 03 primer

749 bp, OPB - 9 609 bp,

OPF - 16 1,100 bp, OPA - 17 580 bp

marker ,

primer . primer , ,

genetic marker

marker ,

3.

sucrose	1.13%	0.02%	,	
fructose	glucose	3.71%, 2.51%	1.83%, 1.05%	
,	malic acid	47,684 ppm, 38,691 ppm	†	
,	citric acid	11,939 ppm, 3,330 ppm	.	
malonic acid	maleic acid	2.8 ppm, 2.3 ppm	0.3	
ppm, trace				
722 mg %			7,577	
mg %	,	202 mg %, 5,596 mg %		
lysine		300 mg %	288	
mg %	.			
glutamic acid, aspartic acid, threonine, valine		Leucine	.	
	1.275%,		1.560%	
,	†			
chlorogenic acid	1,276 ppm, 802 ppm	,	gentisic acid	
69 ppm,	136 ppm			
ferulic acid	187 ppm,	23 ppm		
.	cinnamic acid, caffeic acid, coumaric			
acid				
145 mg/g		160 mg/g		
.	,			

linoleic acid 21,765 ppm, 45,584 ppm      가

palmitic acid 3,974 ppm, 10,872 ppm      가

capric acid .

,

,

-ylangene, , -elemene, -himachalene,

-selinene widdrene ,

caryophyllene, calarene, cubebene, acoradiene -himachalene

가

86.54mg% , 138.75mg%

2

peonidin-3-monoglucoside peonidin-3,5-diglucoside ,

delphinidin-3-monoglucoside malvidin-3-monoglucoside

schizandrin 0.11%/100g( ) ,

0.13%/100g( ) .

4.

가

, ,

가

가      가      ( )

( , )

가

가

가

, , , ,

가

가

가

80

4

60

6

가 5 6 ,  
가 . pH

4 7 , , , , ,

21

20 ( 5 )

(isoascorbic acid) 20 가

가

**B.**

가

가가

RAPD

genetic marker

marker

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가

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가

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가

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가

가

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RAPD

marker ( 25(4) 1998.12.)  
( 12(1) 1998.12.)  
( 12(1) 1998.12.)  
( 1999. 6.)

Sex specific marker of *Schisandra nigra* (1999. 10.)

· ( 1999.12.)  
· ( 2000. 3.)

3

(*Schisandra nigra* Max.),

( 1998. 2.)

DNA marker (2000, 8. )

♂ ( 1999. 2.)

## SUMMARY

1. *Schisandra nigra* is a special use species used as a medical and edible fruit. In addition, it is an endemic species which has a unique habitat at the altitude of 600~1,400m in Cheju island. At present, *Schisandra nigra* is confronted with a crisis due to a sudden increase of demand and imprudent destruction of a rare species. Thus, this study was conducted to find 1)multiplication method and genetic variation, 2)ecophysiological characteristics of the natural habitat of *Schisandra nigra* and 3)detection and preservation of genetic resources and upbringing for a superior description.

1.1. *S. nigra* scatteringly grow at 750~1,200m in the western part, at 600~1,100m in the northern part and at 650~1,400m in the southern part. The geographical distribution of *S. nigra* is the secondary forest of broad leaved trees that are mainly composed of *Piuus densflora*, *Quercus serrata*, *Styrax japonica*, *Lindera erythrocarpa* and *Carpinus laxflora*. The density of a main distribution sector of *S. nigra* appears in the range of 3~39 individuals in each 20×20m plot.

1.2. It was investigated that morphological characteristics on xylem fibers, vessel elements, stomata, and a leaf character of *S. nigra*, *S. chinensis*, *Kadsura japonica*. The results of morphological characteristics were significant at 5% levels within the species and within the individuals as a male, a female and monocious.

1.3. Blooming and fruition of the natural habitat of *S. nigra* were significant

by a region. The most superior region was Youngsil in the western part and Kwaneumsa in the northern part. However, Sanghyo in the southern part was lower than the other regions in the survival rates of a female flower and a quantity of a attached fruit. Fruit characteristics of selected *S. nigra* individuals were averaging at 1.1cm in length, and 741 per liter and 10.3 per cluster in the number of fruits.

1.4. In seed production, the seed germination of the cold moist stratification treatment and the storage in the open ground treatment was the highest in the secondary year(over 93%). In the seed storage and storage pretreatment, the seed current-year germination rate was the highest in the treatment stored at the cold and moist place for 4 mon after immersed in GA<sub>3</sub> 1,000ppm solution during 12 hours.

In cutting reproduction, IBA 500ppm treatment(rooting rate over 80%) showed the higher rate than the other treatments, and the rooting rate by the cutting period was the highest in July(juvenile cutting).

In a grafting reproduction of selected individuals, it indicated a slight difference among individuals, but percents of the graft union was averaging at over 83%.

To propagate the selected individuals by tissue culture, the winter buds were used as a material. Because they were grown in the field, the materials were contaminated after the culture in spite of the surface sterilization. But from the two-years-old twigs showed good growth. WPM medium was better than MS and GA<sub>3</sub> was very effective to differentiate shoots from the buds.

1.5. The distribution and frequency of allele to 4 isozymes among 3 populations of *Schisandra nigra* natural habitat showed a polymorphism in *Idh*, *M dh-2*, *Mnr*, and *Pgi-2* among 6 loci, except *M dh-1*, *Pgi-1*. All allele(*Idh*, *M dh-2*, *Pgi-1*, *M dh-1*, and *Mnr*) except *Pgi-2*, also showed the highest main allele.

2. RAPD (Random Amplified Polymorphic DNA) analysis and southern hybridization were conducted to *S. nigra* plants in order to select the specific markers for monoecious and dioecious individuals. RAPD results using one hundred eighty random 10-mer primers revealed that *S. nigra* had a different banding pattern from *S. chinensis* and *Kadsura japonica*. When DNA isolated from leaves of monoecious and dioecious plants were used as PCR template, only six primers, OPA-17, OPA-19, OPB-03, OPB-09, OPB-16 and OPF-16, showed polymorphic band patterns. No variation in banding profiles within male or female individuals was observed when these six primers were used whereas three monoecious plants(No 6, No 41, No 51) showed different banding patterns one another.

2.1 One primer, OPA-17, yielded DNA fragment of 770 bp, which was detected in male plants but not in any of the female plants tested. By contrast, a 749 bp DNA fragment was amplified in female plants by primer OPB-03, which did not amplified with male DNA samples. We also found three DNA fragments, 609 bp, 1115 bp and 580 bp, which were detected in monoecious but not in male and female plants with primers OPB-09, OPF-16, OPB-03, respectively.

2.2. To verify the specificity of isolated RAPD markers, southern blot hybridization was performed with male, female and monoecious plants DNA. Potential sex-specific RAPD products were isolated from agarose gel and cloned into the pGEM-T Easy vector. Cloned fragments were digested with *Eco*RI and used as probes in southern hybridization analysis of genomic DNA.

2.3. DNAs isolated from male and female plants were separately digested with *Eco*RI, *Bam*HI and *Hind*III. When the male and female DNAs were allowed to hybridize with these probes, the 770 bp male probe yielded some bands specific to male plants whereas the 749 bp female probe resulted in single band only for DNA samples from female plants. The 609 bp of monoecious probe hybridized specifically with DNA from three monoecious plants (6, 41, 51), whereas 1,115 bp probe hybridized specifically with DNA from two monoecious plants (41, 51). The 530 bp of monoecious probe hybridized specifically with only one monoecious plant DNA (51). Three monoecious-specific probes did not hybridized with male and female DNA.

2.4. Five RAPD markers found to be sex-specific were sequenced and the homology searches of the data banks were carried out. The sequences did not include long open reading frames and they exhibited no significant similarity to previous reported sequences. Five set of PCR primers were synthesized based on the nucleotide sequences of the cloned fragment, named male 1, female 1, mono 2-1, mono 2-2 and mono 3 and used for the identification of dioecious and monoecious plants by PCR analysis. These primer sets

amplified 460 bp, 436 bp, 411 bp, 684 bp and 207 bp of DNA fragments, respectively. With the designated male and female primer sets, we could identify the sex of dioecious plants. The primer mono 2-1 amplified expected size for all three monoecious plants, whereas mono 2-2 gave a PCR product with two monoecious plants, No. 41 and 51. However, mono 3 primer produced expected 207 bp fragment with only one monoecious plant of No. 51. All monoecious specific primers did not amplify any bands for male or female plants.

2.5. With these primer sets we could easily and rapidly distinguish male, female or monoecious plant within the *S. nigra* population. These results indicate that the five primer sets could be used as genetic markers for the early discrimination of dioecious and monoecious individuals of *S. nigra*.

3. The results from the determination of content of components and biological active substances in *S. chinensis* and *S. nigra* are as follows.

3.1. The contents of free sugar in *S. chinensis* and *S. nigra* was examined. In the *S. chinensis*, content of sucrose was about 1.13%, and there only a few amount of glucose and fructose. But in the *S. nigra*, content of fructose and glucose was 3.71% and 2.51%, respectively, and only a few amount of sucrose.

3.2. In case of organic acid, *S. chinensis* and *S. nigra* contained the higher percentage of malic acid and citric acid than the others. The contents of

malic acid and citric acid in *S. chinensis* was 47,684 and 11,939 ppm/dry weight 100g, and in *S. nigra* was 38,691 and 3,330 ppm/dry weight 100g, respectively.

3.3. Contents of amino acid in *S. chinensis* was higher than that of *S. nigra*. But, content of lysine and arginine in *S. nigra* was higher than that of *S. chinensis*.

3.4. The contents of total phenolic compounds in *S. chinensis* and *S. nigra* was 1.275% and 1.560%, respectively. The predominating phenolic acid was cinnamic acid, gentisic acid, coumaric acid, chlorogenic acid and ferulic acid.

3.5. Contents of crude lipids in *S. chinensis* and *S. nigra* was 145.5mg/g and 160.5mg/g, respectively. Most of fatty acid in lipids was oleic acid, linoleic acid and linolenic acid as a unsaturated fatty acid, and palmitic acid as a saturated fatty acid.

3.6. In case of essential oils, The predominating components in *S. chinensis* was -ylangene, , -elemene, -himachalene, -selinene and widdrene, *S. nigra* was caryophyllene, calarene, cubebene, acoradiene and -himachalene.

3.7. Contents of total anthocyan pigment in *S. chinensis* and *S. nigra* was 86.54 and 138.75mg%, respectively. Antocyanin, as a aglycone of anthocyan, was supposed to peonidin-3-glycoside in *S. chinensis* and supposed to malvidin-3-glycoside, petunidin-3-glycoside, delphinidin-3- glycoside in *S. nigra*.

3.8. Schizandrin, as a biological active substance, was detected in the extract of *S. chinensis*. Schizandrin-like substance was detected in the extract of *S. nigra*.

4.1. The rate of extract yield from *S. nigra* was best when extracted for 4 hrs. at 80 °C. And we found the possibility that extract from dried stems of *S. nigra* could be used as material for beverage.

4.2. Among Hunter values, both a and b value of the extract from *S. nigra* was much higher than that of *S. chinensis*. As red color was deeper, the extract from *S. nigra* was found to be good in desired process of material to manufacture deep-colored beverage.

4.3. L value, one of Hunter value, and absorbance at 520 nm of the extract from *S. nigra* was low for fresh material, but high for dried material.

4.4. The pH of the extract from *S. nigra* was 0.1-0.2 higher than that of *S. chinensis*. Although the pH of the extract from *S. nigra* was somewhat low when extracted by water, the pH range was to be able to maintain the stability of color the range like *S. chinensis*.

4.5. The sugar content of the extract of *S. nigra* was lower than that of *S. chinensis*, but the difference was insignificant.

4.6. Hunter value a and b of the extract of *S. nigra* extracted for 6hrs. at 60 °C by water was six times higher and 1.5 times higher than that extracted for 24hrs. at room temperature by water. The color of the extract did not

vary when stored in cold for 7 weeks. Hunter value L and absorbance at 520nm was measured by storing at low temperature to investigate the lightness of the extract. Until 4 weeks we found the turbidity increased as L value decreased and absorbance increased. But it seemed to be temporary and little worth consideration because L value and absorbance increased after 5 weeks.

4.7. Total polyphenol of the extract of *S. nigra* decreased during storing period for all samples. Therefore, long-term storage could be the primary factors for browning and altered taste.

4.8. pH of the extract of *S. nigra* decreased but acidity increased during storing period. Therefore, the taste and red color did not change, keeping stability.

4.9. Sugar content of the extract of *S. nigra* did not change at all samples during storing period. For this reason, the quality did not deteriorate easily during storage.

4.10. Beverage products extracted by water, 3% ethanol and aqueous solution from *S. nigra* were stored in cold for 14 weeks and 21 weeks, respectively. Red color was the best for 3% ethanol aqueous solution product at 80<sup>o</sup>, the decrease of which was insignificant during storage. Red color of the extract from dried *S. chinensis* was 1/4 - 1/5 as much as *S. nigra*, which decreased significantly during storage. When we investigated the stability of red color by adding erythobic acid as antioxidant, it seemed to be effective in spite of a small quantity.

4.11. Lightness of the extract of *S. nigra* was higher than that of *S. chinensis*. Lightness was the highest in the extract of dried *S. chinensis*. These results was considered not to be due to turbidity difference but to inverse proportion to the amount of red color. We confirmed the results by measuring absorbance at 520 nm.

4.12. The content of total polyphenol as a barometer of oxidized browning was measured during storage. Total polyphenol content of *S. nigra* and *S. chinensis* decreased a little until 14 weeks, but much after 21weeks. On the contrary, total polyphenol content of dried *S. chinensis* decreased very much after 14 weeks, which indicated the possibility to be browned. Total polyphenol content of product added by antioxidant decreased a liitle until 20 weeks.

4.13. There was no difference in sugar content between *S.nigra* and *S. chinensis*. As sugar content of dried *S. chinensis* was less half than that of *S. nigra*, it seemed that dried *S. chinensis* was not proper to raw material for beverage. Sugar content of extracted product increased during storage, which was considered to be due to altered composition of sugar.

4.14. The acidity of product, which have effects on sour and pH, was the highest for dried *S. chinensis*, in order of *S. chinensis* and *S. nigra*. The acidity of *S. nigra* was 1/3 - 1/4 as much as *S. chinensis*. The acidity decreased alittle during storage. Antioxidant had no effect on acidity.

4.15. pH of the extract product of *S. nigra* ranged from 3.0 to 3.5, and that of *S. chinensis* and dried *S. chinensis* ranged from 2.5 to 2.9. pH had no effect on commercial quality of the product. Adding antioxidant was effective to keep pH from decrease.

4.16. The quality of product, adjusting similar red color and having 6% of sugar content by adding sugar, was investigated. Red color of *S. nigra* product was 6 times higher than that of dried *S. chinensis*, thus less sour and fairly sweet.

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3.	.....	66
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2)	.....	69
4.	.....	70

1)		70
(1)		70
(2)		71
(3) Wright F		74
(4) Nei		75
5.		76
1)		76
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, 20m × 20m (400m<sup>2</sup>)

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, (aspect), (relative location), (local topography), (slope), (soil depth), (soil moisture)

### Global Positioning System

, , 凹面 (concave), 凸面 (convex), (smooth)

(60cm ), (31 60cm ), (30cm )

20

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1995 10 1996 12 , 1  
(8m ), (2 8m ), (2m )

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, DBH,

(1)

(X<sub>ij</sub>)

vegetational data matrix

$$X_{ij} = (d_{ij} + D_{ij}) / 2 \quad (1)$$

X<sub>ij</sub> : j                            i

d<sub>ij</sub> :

D<sub>ij</sub> :

vegetational data matrix      ¶      cluster

Ludwig      Reynolds ¶      statistical ecology basic

program . percent dissimilarity (PD)  
. cluster

Curtis Mckintosh

(importance value, IV) (2) . . .  
 (mean importance value, MIV) (6)

$$IV = ( RD + RC + RF ) / 3 \quad \text{---} \quad (2)$$

IV :

RD

RC :

RF

$$RD = \frac{\Delta P}{P} \times 100(\%) \quad (3)$$

$$RC = \frac{\text{_____}}{\text{_____}} \times 100(\%) \quad (4)$$

$$RF = \text{_____} \times 100(\%) \text{ _____} \quad (5)$$

$$((\text{IV} \times 3) + (\text{IV} \times 2) + (\text{IV} \times 1)) / 6 = \dots \quad (6)$$

Shannon (H')

(Maximum H')

$$H'_{\max} = \log S(S) \quad )$$

$$(J') \quad J' = H' / H' \max \quad , \quad 1 - J'$$

$$H' = - p_i \log p_i \quad (7)$$

( ,  $p_i$  )

Brower Zar Morista's index

$$\text{Morista index} = n \frac{X^2 - N}{N(N-1)} \frac{1}{X} \quad (8)$$

( , n , N , X )

†

Ludwig Reynolds

Pearson Spearman

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F.A.A. (Formalin 5 : Acetic acid

5 : 70% Ethyl alchhol 90 2 3

Acete- carmin	10	70%	ethyl
alcohol	glycerin		

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2 3cm

Schurze 5 7  
 safranin 10 70% ethyl alcohol glycerin  
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 starch gel mold  
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 gel 0.5cm  
 gel gel  
 . gel

### 1.1. Gel tray Buffer

System	Gel Buffer	Tray Buffer
A Formulation	Tris citrate(pH 8.3) Trizma base ----- 62.0g Citric acid ----- 14.6g Distilled water ----- 10.0 Dissolve the chemicals and check the pH. Store at room temperature	Lithium borate(pH 8.3) Lithium hydroxide--- 12.0g Boric acid----- 118.9g Distilled water----- 10.0 Dissolve the chemicals and check the pH. Store at room temperature
	To use, add 75ml of the lithium borate buffer to 765ml of the tris citrate gel buffer to make the required 750ml	Use as it is
C Formulation	Tris citrate(pH 6.2) Trizma base 162g Citric acid 108.9g Distilled water 3.0 Dissolve the chemicals and titrate to pH 6.2 with 4N NaOH.	Same as gel buffer Same as buffer
	Store in the refrigerater To use, mix 16ml of the buffer with 734ml of distilled water.	Mix 250ml of the buffer with 750ml distilled water.

1.2.

System	Stain buffer system used	Tray buffer system used	Stain components
IDH (Isocitric dehydrogenase)	75ml 0.2M Tris HCl	C	DL- Isocitric acid 200mg NADP 10mg MTT 5mg 1% MgCl <sub>2</sub> Solution 1mg
MDH (Malic dehydrogenase)	75ml 0.05M Tris HCl pH 8.0	C	Malic acid solution 5ml (134.1g DL-malic acid, 80g NaOH, adjust to pH 7.0 with about nase of 4N NaOH NAD 1ml NBT 1ml PMS 0.5ml
MNR (Menadione reductase)	75ml 0.05M Tris HCl pH 7.0	A	NADH 25mg Menadione 20mg NBT 1mg
PGI (Phosphogluco se isomerase)	75ml 0.05M Tris HCl pH 8.0	A	D-fructose- 6 phosphate 25mg 1% MgCl <sub>2</sub> Solution 1ml NADP 1ml NBT 1ml PMS 0.5ml G6PDH 20 units

aconitase (ACO, E.C. 4.2.1.3), glutamate dehydrogenase (GDH, E.C.14..3), glutamate-oxaloacetate transaminase (GOT, E.C.2.6.1.1), isocitric dehydrogenase(IDH, E.C. 1.1.1.42), leucine aminopeptidase(LAP, E.C.3.4.11.1), menadion reductase(MNR, E.C. 1.6.99.2), malate dehydrogenase(MDH, E.C. 1.1.1.37), 6-phosphogluconate dehydro-genase(6PG, E.C. 1.1.1.44), phosphoglucose isomerase(PGI, E.C.5.3.1.9),

phosphoglucomutase(PGM, E.C. 2.7.5.1), shikimate - dehydrogenase(SKDH, E.C. 1.1.1.25) 11 가 GDH, GOT , ACO, LAP, 6PG, PGM 가

IDH, MDH, MNR, PGI 4 , 6 (*Idh, M dh-1, M dh-2, M nr, Pgi-1, Pgi-2*)  
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(A/L), (P),  
(H<sub>o</sub>) (H<sub>e</sub>), ,  
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Fisher exact-test , X<sup>2</sup>-test  
exact-test Hardy - Weinberg

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UPGMA PC  
BIOSYS - 1 program

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IBA, NAA, IAA

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4% 1

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, 1:1:1(v/v) , IBA 500 mg/

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가 黑五味子

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VPER(vermiculite + perlite), VPEA(vermiculite +  
peatmoss), VPPL(vermiculite+peatmoss+perlite) 1:1:1(v/v)  
IBA 500mg/

(4)

. 1,  
2, 3, 4, 5, 6, 7, 8, 9, 10, 13, 15, 16, 20 14 7 12  
. , , 1:1:1(v/v)  
, IBA 500mg/

3)

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Tween 20 2%

NaClO                  20                  , 0.2%      HgCl<sub>2</sub>                  10  
4                  .                  95%      EtOH  
.                  .                  †                  4†

BAP      TDZ                  , BAP      GA<sub>3</sub>

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2	870 890	5 10	SW			
1	770 790	5 20	E			
1	880 910	10 30	NE NW			
1	780 800	5 30	SE			

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1.4.

pH 4.36 5.28

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(C.E.C.)

$16.7 \pm 3.83$  (10.21 22.74) me/ 100mg

11.34

me/ 100mg

1.4.

	(% )	(% )	pH	(m e/ 100g)					P <sub>2</sub> O <sub>5</sub>
				K <sup>+</sup>	Na <sup>+</sup>	Ca <sup>++</sup>	Mg <sup>++</sup>	C.E.C	
1	22.51	0.61	4.93	0.20	0.16	0.16	0.34	15.74	3.05
2	23.58	1.00	4.37	0.38	0.24	0.40	0.90	12.15	1.30
3	21.27	0.85	5.14	0.26	0.19	0.40	0.50	12.57	0.30
4	25.08	1.33	4.97	0.46	0.33	2.30	1.50	10.21	1.80
5	27.21	0.69	4.92	0.16	0.24	0.30	0.30	13.32	2.02
6	24.86	0.53	5.02	0.35	0.25	2.33	1.48	17.61	2.46
7	24.18	0.96	4.79	0.47	1.43	6.14	0.31	22.19	4.07
8	23.82	0.99	5.28	0.31	0.83	1.40	0.80	16.07	0.30
9	24.51	0.92	4.81	0.42	1.23	3.25	0.56	19.82	3.12
10	22.48	0.61	4.53	0.38	0.33	2.52	0.85	18.22	2.07
11	23.51	1.27	4.90	0.44	0.98	1.70	1.00	19.21	2.40
12	19.27	1.14	4.90	0.35	1.32	0.80	0.60	20.23	1.70
13	15.45	0.68	5.04	0.38	1.43	7.78	3.68	22.74	2.21
14	21.83	0.81	4.58	0.37	0.19	1.00	0.88	16.74	1.88

(2)

, , , ,

2

(400m<sup>2</sup>)

3 39

( 1.5.).

1.5.

		plot (No./400m <sup>2</sup> )	
.	-	14	
.	-	1	13 13
a.	-	3	39 26 49
b.	-	3	32 16 48
c.	-	3	12 6 20
a.	-	2	6 4 - 7
b.	-	2	3 4 - 2

Spearman

負

, , 가

, , , 가

가 5

2	106.2	가	,			9.3
11.1	가	.	2	0.99g	가	
			2	795		1
2	722	727			( 1.6.).	

1.6.

	( )	(m)	(cm)	( )	( )	(g)	(cm)	( )
1	50	770- 890	2.32	70.2	10.0	0.83	1.12	740
2	27	850- 890	2.13	64.7	10.4	0.82	1.12	735
2	6	850- 870	2.55	70.3	9.8	0.75	1.09	795
	5	850- 860	2.76	-	-	-	-	-
1	5	770- 790	2.34	63.8	11.1	0.99	1.19	722
2	5	770- 780	2.70	106.2	9.3	0.84	1.11	727
	2	850- 870	2.75	-	-	-	-	-

3)

가 ( 1.1.).

가

가

( 1.2). 가

, 가

가

,

,

가

( 1.7).



1.1.

A :                  B :                  C:



1.2.

A :                  , B :                  , C :

1.7.

	(m)	(cm)	( )	( )	(g)	(cm)
1- 1	880	2.6	16	11.2	0.62	1.03
" 2	890	1.8	15	11.2	0.65	1.02
" 3	890	3.0	23	9.2	0.78	1.10
" 4	880	1.5	20	10.2	0.74	1.10
" 13	890	2.9	63	9.0	0.53	0.92
2- 5	850	1.5	102	11.8	0.69	0.98
" 6	850	2.5	135	18.6	0.86	1.10
" 7	850	1.2	4	10.0	0.60	1.03
" 8	850	2.5	66	9.6	0.84	1.14
" 9	850	1.2	11	8.8	0.56	0.94
" 10	850	2.6	6	10.0	0.59	0.95
" 11	850	2.0	28	12.0	0.62	0.97
" 12	850	1.5	9	8.8	0.67	1.01
" 14	860	1.6	4	8.0	0.72	1.11
" 15	850	1.5	5	9.6	0.56	0.99
" 16	850	1.5	16	9.6	0.54	0.99
" 17	850	2.0	15	11.8	0.61	1.01
" 18	850	2.0	80	14.2	0.62	1.03
" 19	850	2.2	8	9.0	0.73	1.11
" 20	860	1.5	52	8.4	0.49	0.94
" 21	850	1.9	16	11.4	0.67	1.08
" 22	850	2.1	6	11.2	0.57	0.99
" 23	850	1.7	28	11.4	0.65	1.04
" 24	850	1.8	10	12.6	0.74	1.06
" 25	850	1.5	47	9.4	0.57	0.99
" 26	850	2.3	16	12.8	0.60	0.93
" 27	860	1.3	5	8.8	0.50	0.83
" 28	790	2.4	62	12.0	0.67	1.00
" 29	790	2.3	17	10.8	0.71	1.00
" 30	780	1.8	18	8.6	0.66	1.07

2)

(1)

가 가 ,  
가  
가 가 .  
가 2 ,  
가

Table 1.8 1.9 , , , ,

1.8.

(m)				
600	$5.11 \pm 0.70$	$3.34 \pm 0.48$	$1.91 \pm 0.56$	$0.12 \pm 0.02$
900	$5.16 \pm 1.01$	$3.23 \pm 0.60$	$1.51 \pm 0.53$	$0.12 \pm 0.02$
1100	$5.23 \pm 0.71$	$3.37 \pm 0.53$	$1.81 \pm 0.45$	$0.12 \pm 0.01$
1300	$5.30 \pm 0.85$	$3.29 \pm 0.62$	$1.78 \pm 0.54$	$0.11 \pm 0.01$
Total	$5.20 \pm 0.83$	$3.31 \pm 0.55$	$1.75 \pm 0.54$	$0.12 \pm 0.01$

600, 900, 1,100, 1,300 m ,

가

900 m

가

가

1.9.

	$5.17 \pm 0.76$	$3.31 \pm 0.53$	$1.82 \pm 0.36$	$0.12 \pm 0.01$
	$5.18 \pm 0.76$	$3.30 \pm 0.52$	$1.78 \pm 0.44$	$0.11 \pm 0.01$
	$5.20 \pm 0.87$	$3.29 \pm 0.56$	$1.66 \pm 0.51$	$0.12 \pm 0.01$
	$5.18 \pm 0.80$	$3.34 \pm 0.54$	$1.75 \pm 0.45$	$0.11 \pm 0.01$

가

가

가

2)

Perigenous

가 79.90 μm

가

75.20  $\mu\text{m}$

92.50 110.75  $\mu\text{m}$

,

가

가 694.38/ $\text{mm}^2$

가

395.33 ,

, 4 가 , ,  
,

가 .

1.10.

	( $\mu\text{m}$ )	( $\mu\text{m}$ )	/	( $\text{mm}^2$ )
<i>S. chinensis</i>	$79.90 \pm 0.351$	$44.20 \pm 0.264$	1.939	$39.533 \pm 4.048$
<i>S. nigra</i>	$97.50 \pm 0.543$	$63.50 \pm 0.812$	1.535	$31.400 \pm 2.472$
<i>S. nigra</i> 41	$105.88 \pm 0.960$	$58.75 \pm 0.244$	1.857	$20.435 \pm 2.635$
<i>S. nigra</i> 6	$110.75 \pm 1.325$	$64.25 \pm 0.504$	1.724	$23.115 \pm 2.207$
<i>S. nigra</i>	$92.50 \pm 0.598$	$57.75 \pm 0.717$	1.602	$25.714 \pm 2.358$
<i>Kadsura japonica</i>	$76.20 \pm 0.268$	$42.40 \pm 0.504$	1.797	$69.438 \pm 4.527$

3)

4 , 가  
.

가  
.

752  $\mu\text{m}$  가 , 가 820  $\mu\text{m}$  , 가 862 947  $\mu\text{m}$   
가

10% 가 .

가 가 , ,

3

,

,

가

가

1.11.

	( $\mu\text{m}$ )	( $\mu\text{m}$ )	( $\mu\text{m}$ )	( $\mu\text{m}$ )
<i>S. chinensis</i>	510.80 $\pm$ 8.685	64.84 $\pm$ 1.282	75.200 $\pm$ 12.092	17.20 $\pm$ 0.220
<i>S. nigra</i>	702.67 $\pm$ 14.431	64.90 $\pm$ 1.474	90.333 $\pm$ 18.023	21.97 $\pm$ 0.381
<i>S. nigra</i> 41	690.00 $\pm$ 13.646	71.92 $\pm$ 1.393	86.267 $\pm$ 16.671	21.50 $\pm$ 0.163
<i>S. nigra</i> 6	732.00 $\pm$ 12.629	146.66 $\pm$ 3.060	94.700 $\pm$ 21.011	22.47 $\pm$ 0.334
<i>S. nigra</i>	574.33 $\pm$ 11.789	91.13 $\pm$ 2.310	68.033 $\pm$ 19.907	21.87 $\pm$ 0.307
<i>Kadsura japonica</i>	680.00 $\pm$ 12.628	76.47 $\pm$ 2.261	82.000 $\pm$ 17.928	20.60 $\pm$ 0.178

3

2)

(1)

가

,      가      ,      (pollen   sac)      ,

, 가 , stigma가 ,

( 1.3). 가

3

3

가 , 3 가 9 가

6 3

( )

가 15      가 , 15  
( 1.4 ).  
2 3 ,

10 ( 1.5 ).

1: 6.3

가

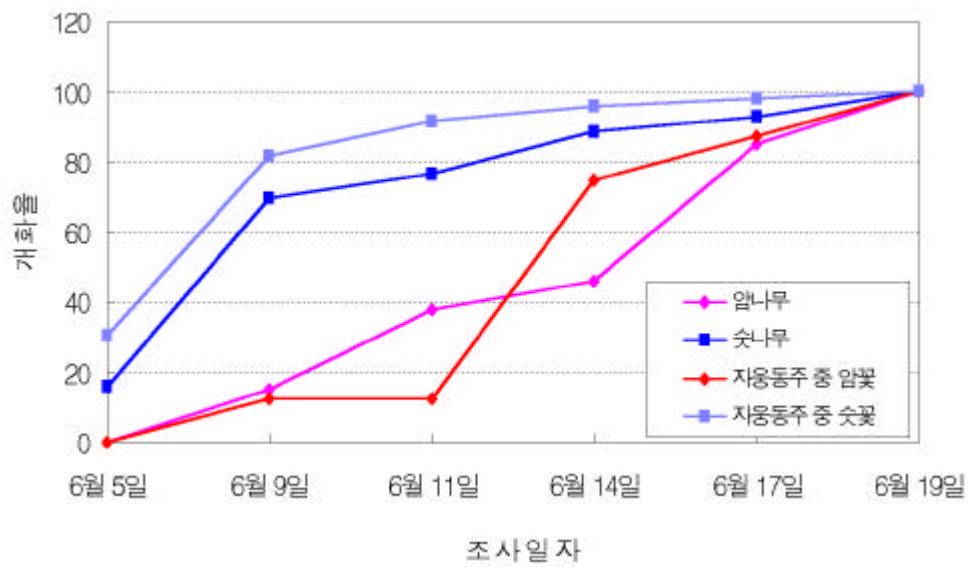
, 가

1.12.

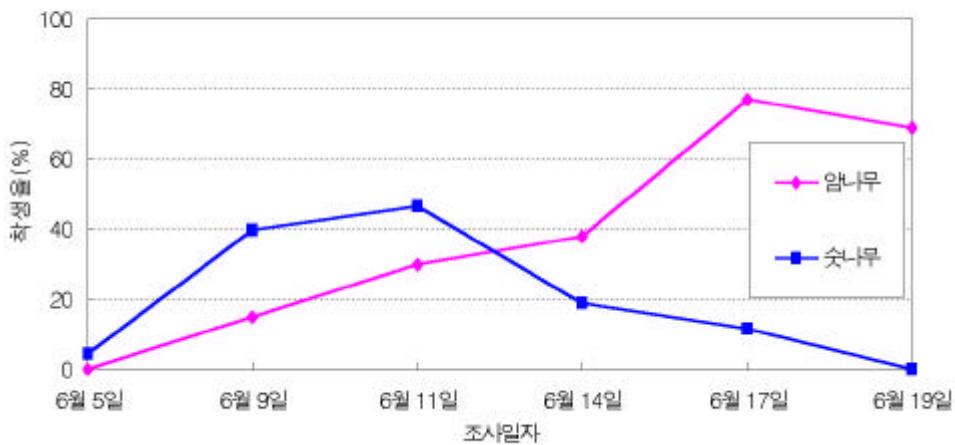
		6 5	6 9	6 14	6 17
		-	$5.35 \pm 1.44$	$5.49 \pm 1.52$	$5.59 \pm 1.56$
		-	$3.00 \pm 0.48$	$3.02 \pm 0.49$	$3.06 \pm 0.62$
(cm)		$7.04 \pm 0.43$	$7.08 \pm 0.57$	$7.21 \pm 0.64$	$7.52 \pm 0.49$
		$3.35 \pm 0.62$	$3.50 \pm 0.81$	$3.76 \pm 1.75$	-



1.3. (A: , B: , C: )



1.4.



1.5.

(2)

5 , ,  
21.2 , 14.4  
68.0% .  
905g ( 1.13).

1.13.

	(m)	(cm)	( )	( )	(%)	(g)
1	850	2.5	20.8	12.8	62	710
2	850	2.5	24.5	18.6	76	2,159
3	890	2.9	20.1	14.2	71	744
4	850	2.0	20.8	14.2	68	704
5	850	2.0	19.6	12.0	61	208
		2.4	21.2	14.4	68	905

4 4

, , , , 19.22 19.82  
가 , 1.8 10.5 가  
. . 10.5 54%  
( 1.14).

1.14.

	(m)	( )	( )	(%)
	850-890	19.32	8.9	46
	770-790	19.51	10.5	54
	550-800	19.22	1.8	9
	650-750	19.82	6.5	33
	550-750	19.50	6.9	36

4.

1)

(1)

$$6 \quad \text{가} \quad M dh-1 \quad Pg i-1$$

$$2 \quad \text{가} \quad 4$$

$$(1.15). Pg i-2 \quad \text{가}$$

$$\text{가} \quad \text{가}, Pg i-2$$

$$3 \quad \text{가} \quad 2$$

$$, b\text{가}$$

$$a\text{가}$$

$$Idh-1 \quad a\text{가} \quad 0.033$$

$$, M dh-2 \quad 3$$

$$\text{가} \quad , \quad 1$$

Population				Population			
Locus	1	2	3	Locus	1	2	3
<i>Idh</i>							
(N)	30	30	30	(N)	30	30	30
a	.033	.000	.000	a	1.000	1.000	1.000
b	.967	1.000	1.000				
<i>M dh- 1</i>							
(N)	30	30	30	(N)	30	30	30
a	.083	.000	.133	a	.783	.833	.983
b	.883	1.000	.850	b	.217	.167	.017
c	.033	.000	.017				
<i>M dh- 2</i>							
(N)	30	30	30	(N)	30	30	30
a	.083	.000	.133	a	.783	.833	.983
b	.883	1.000	.850	b	.217	.167	.017
c	.033	.000	.017				
<i>Pgi- 1</i>							
(N)	30	30	30	(N)	30	30	30
a	1.000	1.000	1.000	a	.250	.283	.617
				b	.683	.667	.383
				c	.067	.050	.000
<i>Pgi- 2</i>							
(N)	30	30	30	(N)	30	30	30
a	.083	.000	.133	a	.783	.833	.983
b	.883	1.000	.850	b	.217	.167	.017
c	.033	.000	.017				

N: , 1: Youngsil( ) 2: Sanghyo( ); 3: Kwaneumsa( )

(2)

1. 26

1.7 , 95% 99%

38.9% 50.0%,

0.141 0.147

, 가

가

,

96

( $A/L = 1.68$ ;  $P = 45.1\%$ ;  $H_e = 0.143$ : Hamrick et al., 1992)

,

가 (endemic:  $A/L = 1.48$ ;  $P = 26.3\%$ ;  $H_e = 0.056$ )

( $P = 67.7\%$ ,  $A/L = 2.15$ ,  $H_e = 0.251$ )

( $A/L = 1.90$ ;  $P = 60.3\%$ ;  $H_e = 0.208$ )

가

1.16.

Population	$A/L$	$P_{95}$	$P_{98}$	$H_o$	$H_e$
Youngsil	2.0	50.0	66.7	.172	.183
	( .4)			( .077)	( .080)
Sanghyo	1.5	33.3	33.3	.111	.127
	( .3)			( .070)	( .084)
	1.7	33.3	50.0	.139	.130
( .3)				( .087)	( .082)
Mean	1.7	38.9	50.0	.141	.147

Note:  $A/L$ , number of alleles per locus;  $P_{95}$ , percentage of polymorphic loci at 95% level;  $P_{99}$ , percentage of polymorphic loci at 99% level;  $H_o$ , observed heterozygosity;  $H_e$ , Nei's unbiased expected heterozygosity

가

가

가

,

가

가

가

,

가

가

(鳥)

,

,

가

가

子

가

가

가

가

(3) Wright  $F$

Wright  $F$        $F_{IS}$        $F_{IT}$       0      가

Hardy - Weinberg

( ).

Hardy - Weinberg

,

Hardy - Weinberg

가

가

Hardy - Weinberg

가

,

가

가

1.17.

Wright  $F$

Locus	$F_{IS}$	$F_{IT}$	$F_{ST}$
<i>Idh</i>	-.034	-.011	.022
<i>Mdh-2</i>	.009	.054	.045
<i>Mnr</i>	-.231	-.154	.063
<i>Pgi-2</i>	.150	.227	.091
Mean	.025	.097	.074

가 7.4% ( $F_{ST} = 0.074$ ),  
92.6%

가 ( $G_{ST} = 0.124$ ) ( $G_{ST} = 0.141$ )

가  
가

pattern

가

#### (4) Nei

Nei	(D)	(I)	$F_{ST}$	가
	0.018	0.983		

1.18.

Nei

Population	1	2	3
Young sil	*****	.000	.028
Sanghyo	1.000	*****	.025
Kwaneumsa	.972	.976	*****

UPGMA

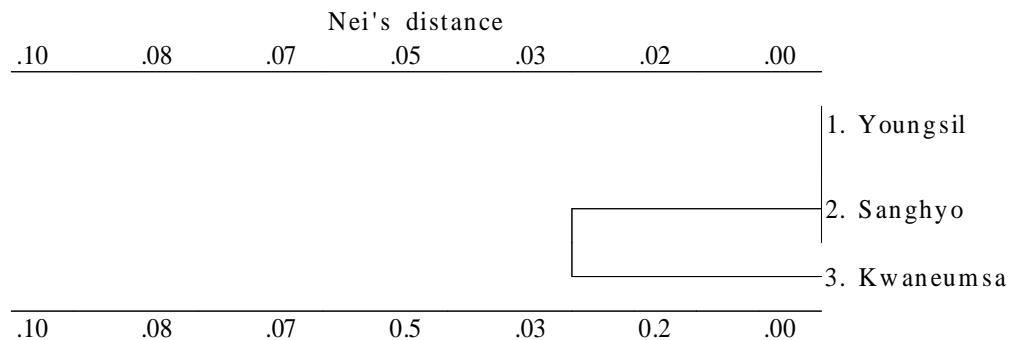
1.6

Pgi-2

가

가

가



Cophenetic correlation = .994

1.6. Nei

5.

1)

가	.	1995	9	20	.	1995	10	5
<b>果肉</b>								
.		4		,	4	,	,	,
4			5			300		
2				1.19				

1.19.

	( )		
		(%)	2 (%)
	300	0	61
	300	9	95
	300	8	93
	300	0	38

가

8 9%

,

가

. 2

가 93 95%

38 61%

가

가

, H<sub>2</sub>O<sub>2</sub> 10%, GA<sub>3</sub> 1,000ppm, BAP 1,000ppm,

TWEEN Alcohol 30%

12

, , ,

25

, 4

,

4 7 , 25 7

5

× 4

,

60

GA<sub>3</sub> 1,000ppm 가

15 29% 가

,

가 가

GA<sub>3</sub>

가 가

29%

,

9%

가

3

가

Hartman Kester

가

,

가

2

가

가

4가

5

95% 93%

( 1.20).

1.20.

		( )	( )	(%)
4	H <sub>2</sub> O <sub>2</sub>	150	20	13
	GA <sub>3</sub>	150	44	29
	TWEEN	150	28	19
	Alcohol	150	0	0
	BAP	150	0	0
2 + 3	H <sub>2</sub> O <sub>2</sub>	150	16	4
	GA <sub>3</sub>	150	38	14
	TWEEN	150	32	10
	Alcohol	150	11	4
	BAP	150	5	2
4	H <sub>2</sub> O <sub>2</sub>	150	16	3
	GA <sub>3</sub>	150	22	7
	TWEEN	150	15	5
	Alcohol	150	7	3
	BAP	150	0	0
4	H <sub>2</sub> O <sub>2</sub>	150	14	4
	GA <sub>3</sub>	150	26	10
	TWEEN	150	18	6
	Alcohol	150	5	2
	BAP	150	0	0
24		150	33	14
4		150	0	0

2)

(1)

IBA 69%, NAA 55%, IAA  
 52%, 52% IBA 가 가 NAA 50  
 mg/ 가 30% 가 .  
 IBA 가 6.6, 6.1 cm 가  
 , 가 47, IAA 가 5.1  
 cm 가 .

1.21.

(mg/ )	( )	( )	(%)	( )	(cm)
IBA 50	60	34	57	5.7	6.4
100	60	46	77	6.6	6.2
500	60	48	80	6.4	6.1
1000	60	37	62	7.6	5.8
Mean	60	41	69	6.6	6.1
NAA 50	60	18	30	4.8	5.2
100	60	38	63	5.4	5.3
500	60	46	77	5.9	6.2
1000	60	29	48	6.1	5.6
Mean	60	33	55	5.6	5.6
IAA 50	60	24	40	4.8	5.4
100	60	29	48	4.6	5.9
500	60	38	63	4.9	4.9
1000	60	34	57	4.3	4.2
Mean	60	31	62	4.7	5.1
	60	31	52	4.7	5.8

7 10 IBA 500 mg/  
 가 , 8 24 가  
 48% 가 .  
 7 10 가 73% 가 8 24  
 가 63% .  
 8

3

가 7 IBA 500  
가 가 .

Table 1.22. IBA

	3	20	6	10	8	1	8	24
IBA 50ppm	57		63		68		55	
	100	77		78		78		77
	500	80		83		78		78
	1000	62		80		58		57
		52		60		57		48
		66		73		67		63

IBA 500mg/  
가

가

(2)

63 88% 가

5.9      9.6    ,                5.8      7.3 cm,            6.1      9.0 cm  
 ՚

가 , Kim Nam 5, 10, 20

가 85.7%, 81.7%, 62.4%

Yim 1, 2, 10, 20 84%, 43%,  
 28%, 15%  
 15 63% 7

1.23.

---

	( )	( )	(%)	( )	(cm)	(cm)	(mm)
1	60	53	88	9.6	7.3	9.0	1.2
5	60	49	82	8.1	6.9	7.5	1.3
10	60	45	75	7.3	6.2	7.4	1.2
15	60	38	63	5.9	5.8	6.1	1.2
		60	77	7.7	6.6	7.5	1.2
F -			15.90**	12.78**	9.30**	9.42**	0.97 <sup>NS</sup>

\*\* : significant at 1% level

NS : Non significant at 1 or 5% level

(3)

60%, 67%,  
 50% 7  
 . , ,  
 5.0 7.4 , 5.7 6.7cm, 5.7 7.0cm 1.0  
 2.1mm 7

## 1. 24.

	( )	(%)	( )	(cm)	(cm)	(mm)
	60	36	60	7.1	6.4	6.8
	60	40	67	7.4	6.7	7.0
	60	30	50	5.0	5.7	5.7
	60	35	59	6.5	6.3	6.5
F -			4.618**	89.36**	13.80**	27.40**
						6.89**

\*\* : significant at 1% level

NS : Non significant at 1 or 5% level

## Kummerow

가

가

, ,

Farrar and Grace *Pinus monticola*

가 , 가

가

가

Table 21

VPPL	가 68% 가	가	가	가
38%	가		가	.
	,	,		.
6.0	6.5cm,	6.4	6.7cm	6.2 7.0 ,
				2.0 2.1 mm

	( )	( )	(%)	( )	(cm)
VPER	60	23	38	6.2	6.0
VPEA	60	38	63	6.6	6.5
VPPL	60	39	65	6.3	6.2
	60	41	68	7.0	6.3
	60	35	59	6.5	6.3
F -			17.31 <sup>**</sup>	0.868 <sup>NS</sup>	0.826 <sup>NS</sup>

\*VPER:vermiculite+perlite 1:1(v/v), VPEA:vermiculite+peatmoss 1:1 (v/v),

VPPL:vermiculite+peatmoss+perlite 1:1(v/v), \*\* : significant at 1% level, NS :

Non significant at 1 or 5% level

가

,

가

가

, , , ,

,

, , ,

,

(4)

40 85%

2, 9, 25 가 80% 가 , 10

가 40%

가

, 5.0 7.6 ,

5.4 7.4cm

가

Table 1.26.

	( )	( )	(%)	( )	(cm)
1	20	15	75	6.4	7.4
2	20	17	85	5.8	6.8
3	20	11	55	6.2	6.9
4	20	13	65	7.6	6.6
5	20	16	80	5.2	5.4
6	20	9	45	6.4	6.6
7	20	12	60	5.2	6.3
8	20	14	70	5.0	5.8
9	20	17	85	7.3	6.8
10	20	8	40	6.6	6.4
13	20	13	65	6.2	7.6
15	20	17	85	5.8	7.8
16	20	15	75	6.2	6.4
20	20	10	50	5.6	6.8
	20	13	67	6.1	6.7

3)

30

63 92%

( 1.27).

83%

†

1.27.

	( )	( )	(% )		( )	( )	(% )
1	24	20	83	30	24	15	63
2	24	17	71	31	24	18	75
3	24	22	92	32	26	17	65
4	24	21	88	33	18	13	72
5	24	20	83	34	25	19	76
6	24	22	92	35	30	18	60
7	24	21	88	36	20	10	50
8	24	18	75	37	22	12	55
9	24	20	83	38	26	15	58
10	24	18	75	39	24	20	83
12	24	23	72	40	24	18	75
13	24	22	92	41	24	18	75
14	24	21	88	42	22	14	64
15	24	28	75	43	22	17	77
19	24	16	73	46	24	14	70
20	24	20	83	47	26	20	83
22	24	16	73	48	28	11	42
28	24	17	72	49	24	18	64
30	24	15	63	50	26	17	71
					983	702	71

4)

. 6 41 , 8 , 14 , 22 27

(1)

가 11 가

가

Tween 20 2% NaClO  
20 , 0.2% HgCl<sub>2</sub> 10 4  
95% EtOH .  
가 4가

1.28.

		(%)		(%)
6	291	47.8	356	23.9
8	80	51.2	72	19.4
14	61	47.5	59	35.5
22	174	54.0	160	25
27	40	67.5	56	21.4
41	39	56.4	24	20.8
	114.2	54.1	121.2	24.3

가

가

(54.1% )

24.4%

가

1.29.

( 6 )

(mg/ )		
WPM 가	4.4 ± 1.0	3.9 ± 1.2
	2.9 ± 0.9	2.7 ± 1.0
	3.2 ± 1.5	3.8 ± 1.4
	3.2 ± 0.7	2.9 ± 0.9
MS 가	3.5 ± 1.5	3.8 ± 1.3
	2.8 ± 1.1	1.9 ± 0.8
	2.8 ± 1.1	2.7 ± 1.1
	2.7 ± 0.8	2.2 ± 0.8

2 4

,

가

1.30.

( 6 )

(mg/ )		
WPM 가	76.2 ± 21.1	74.6 ± 22.9
	100.5 ± 37.6	80.9 ± 26.6
	80.4 ± 36.0	97.7 ± 20.7
	120.6 ± 57.4	91.9 ± 31.1
MS 가	75.3 ± 38.1	82.2 ± 27.7
	98.3 ± 39.5	82.0 ± 31.2
	88.0 ± 32.9	87.1 ± 40.4
	126.0 ± 45.7	82.4 ± 37.4

BAP 1.0mg/ BAP GA<sub>3</sub> 가  
WPM MS , 가

### 1.31. (%)

(mg/ )			
WPM	가	0	0
	BAP 1.0	9.1	0
	GA <sub>3</sub> 1.0	55.6	19.4
	BAP 1.0+GA <sub>3</sub> 1.0	23.1	4.0
MS	가	0	0
	BAP 1.0	0	0
	GA <sub>3</sub> 1.0	35.7	14.3
	BAP 1.0+GA <sub>3</sub> 1.0	11.8	0

GA<sub>3</sub> BAP가 가

가

GA<sub>3</sub> 가 BAP 가

WPM

가 . 가 GA<sub>3</sub>  
가

1.32.

,

BAP 0.5	264.8 ± 165.3	1.1 ± 1.1	10.6 ± 5.9
BAP 1.0	243.5 ± 113.3	2.1 ± 1.9	8.0 ± 3.6
BAP 2.0	277.8 ± 154.6	3.0 ± 2.5	7.3 ± 3.0
BAP 3.0	267.6 ± 59.4	2.3 ± 2.3	7.5 ± 3.8
BAP 5.0	315.6 ± 140.9	1.9 ± 1.7	9.4 ± 5.5
TDZ 0.05	349.7 ± 185.2	2.8 ± 2.9	7.4 ± 3.3
TDZ 0.1	302.2 ± 146.1	2.0 ± 1.8	7.9 ± 3.9
TDZ 1.0	296.3 ± 169.4	2.8 ± 2.5	7.4 ± 3.9

BAP GA<sub>3</sub>

BAP GA<sub>3</sub>

GA<sub>3</sub>

가 BAP

,

GA<sub>3</sub> 3.0mg/ BAP 2.0mg/ 가 가 가

( 1.32.).

1.33.

BAP 1.0 + GA <sub>3</sub> 1.0	218.9 ± 89.1
BAP 1.0 + GA <sub>3</sub> 2.0	159.5 ± 72.3
BAP 1.0 + GA <sub>3</sub> 3.0	217.5 ± 74.7
BAP 2.0 + GA <sub>3</sub> 1.0	194.3 ± 107.5
BAP 2.0 + GA <sub>3</sub> 2.0	251.8 ± 126.2
BAP 2.0 + GA <sub>3</sub> 3.0	153.9 ± 77.8
BAP 3.0 + GA <sub>3</sub> 1.0	169.3 ± 68.6
BAP 3.0 + GA <sub>3</sub> 2.0	258.3 ± 92.1
BAP 3.0 + GA <sub>3</sub> 3.0	172.3 ± 123.4

,  
, ( 1.7).

5)

,  
 $18\text{m}^2$  1 ,  $36\text{m}^2$  1 ,  
 $170\text{m}^2$   
25 86 ( 1.8).  
1997 1999 4 ♂ ( 1.34).  
26.4% , 327 , 248  
, 6.3:1 .

1.34.

( )	( )	(% )		
125	33	26.4	327	248



1.7. (A : , B : , C : )



1.8. Pergola

## 4

, 가 가

600 1,350m ,  
 가 35. , 2 , 400m<sup>2</sup>  
 3 39 . 3,000 3,094 mm ,  
 8.8 10.5 0.6  
 1,300m - 7 가  
 가 . 50 . 5  
 6 . 가  
 , 1:6.3 . 19.2 19.8  
 . 106.2 , 9.3  
 11.1 가 . 0.99g, 722 795 .  
 , IBA 500ppm 80% 가  
 , 63 92% .  
 WPM BAP GA<sub>3</sub> 1.0 mg/  
 , BAP 3.0mg/ 가 3

,  
 ,  
 , 가 200

, 가

가 ,

### 3 RAPD

1

RAPD

genetic marker

DNA

DNA yield 가 PCR

genomic DNA

, PCR template DNA, Mg<sup>2+</sup>, Taq DNA polymerase, dNTP  
annealing cycle band pattern 가 RAPD

가

Callus

가 가

, marker marker  
, marker cloning kit 가

2

1.

*Schisandra* 19

3 (6 , 41 , 51 ), 8 (46, 45 , 47 , 48 ,

49 , 50 , 52 , 53 ), 7 (29 , 32 , 33 , 42 , 44 , 54 , 55 )

1

*Kadsura japonica*

## 2. Genomic DNA

500 mg  
10  $\gamma$  isolation buffer [10% polyethylene glycol 6000, 0.35 M sorbitol, 0.1 M Tris (pH 8.0), 0.5% spermidine, 0.5% spermine, 0.5% - mercaptoethanol], (4 , 15,000 rpm, 10 )  
5 lysis buffer [0.35 M sorbitol, 0.1 M Tris (pH 8.0), 0.5% spermidine, 0.5% spermine, 0.5% - mercapto - ethanol] 1/10  
10% sarcosine  $\gamma$ , 10 , 2 CTAB  
[2% cetyltrimethylammonium bromide (CTAB), 0.1 M Tris (pH 9.5), 20 mM EDTA, 1.4 M NaCl, 0.5% - mercaptoethanol], 65 , 10  
chloroform (chloroform:isoamylalchol = 24:1)  $\gamma$   
(25 , 15,000 rpm, 10 ),  
isopropanol . DNA pipet  
 $\gamma$  99.9% ethanol tube ,  
 $\gamma$  70 80% ethanol 1 M $\ell$   $\gamma$  ,  
, 200  $\mu\ell$  TE buffer [10 mM Tris (pH 8.0), 1 mM EDTA], phenol:chloroform (1:1)  
(25 , 15,000 rpm, 10 )  
isopropanol , 99.9% ethanol 1 M $\ell$   $\gamma$   
(4 , 15,000 rpm, 10 ) 70%  
ethanol (4 , 15,000 rpm, 10 )  
DNA TE buffer [10 mM Tris (pH 8.0), 1 mM EDTA]

EDTA 1% PCR buffer 10 mM Mg<sup>2+</sup>

DNA Hoefer DNA fluorometer (TKO-100)

### 3. Random primers

PCR Primer Operon Technologies, Inc. random primer  
primer OPA kit (20 primers), OPB kit (20 primers), OPC kit (20 primers),  
OPD kit (20 primers), OPE kit (20 primers), OPF kit (20 primers), OPG kit  
(20 primers) , OPH kit (20 primers), OPI kit (20 primers) 180 primer

### 4. PCR

#### A. RAPD (Random Amplified Polymorphic DNA)

PCR 20 ng template DNA, 25 pmole primer (decanucleotide),  
200 μM deoxyribonucleic acid, 1.5 mM MgCl<sub>2</sub>, 1/10 10 × PCR buffer  
[100 mM Tris-HCl, 25 mM MgCl<sub>2</sub>, 500 mM KCl (pH 8.3)], 1.25 unit Taq  
DNA polymerase (Takara) 25 μl PCR  
Perkin Elmer GeneAmp PCR system 2400 94 5  
denaturation, 94 30 denaturation, 37 30  
annealing, 72 30 extension 45 . 72  
5 PCR 1% agarose gel  
ethidium bromide

marker PCR  
PCR 94 5

denaturation , 94 30 denaturation, 55 30  
annealing, 72 30 extension 30 . 72  
5 . PCR 1.2% agarose gel  
ethidium bromide

5. Agarose gel DNA

RAPD DNA , marker  
band agarose gel GENE CLEAN kit  
(Bio101) . gel 3 NaI  
, 45 55 5 . 5  $\mu\text{l}$  glassmilk  
, 5 (15,000 rpm, 5 )  
200  $\mu\text{l}$  New wash (New concentrate 14 M $\text{M}$ , H<sub>2</sub>O 280 M $\text{M}$ , 99.9%  
ethyl alcohol 310 M $\text{M}$ ) 3 . TE buffer 4  
5 55 3 (15,000 rpm, 3 )  
. DNA sample Hoefer DNA fluorometer (T KO- 100)

6. marker cloning

, marker band  
, pGEM-T Easy Vector System (Promega ) cloning  
. insert vector 3:1 ↗ . DNA  
1  $\mu\text{l}$  T4 DNA ligase 10 x buffer, pGEM-T easy vector (50 ng, 2-1) 1  
 $\mu\text{l}$ , T4 DNA ligase (3 Weiss units/ $\mu\text{l}$ ) , 10  $\mu\text{l}$   
. 4 16 24  
ligation 50  $\mu\text{l}$  competent cell

, 20 . 42 45 50  
 , . 950  $\mu\text{l}$  SOC medium [0.5% (w/v)  
 yeast extract, 2% (w/v) tryptone, 10 mM NaCl, 2.5 mM KCl, 10 mM MgCl<sub>2</sub>,  
 20 mM MgSO<sub>4</sub>, 20 mM glucose]  
 , 37 1  
 30 . ampicillin (50  $\mu\text{g}/\text{Ml}$ ), 20  $\mu\text{l}$  X-gal (50  
 mg/Ml in dimethylformamide), 100  $\mu\text{l}$  IPTG (100 mM) ↗ LB (tryptone  
 10 g, yeast extract 5 g, NaCl 5 g, 1 M NaOH 1 Ml/ ) 37  
 16 24 . colony  
 plasmid DNA *EcoR*  
 marker

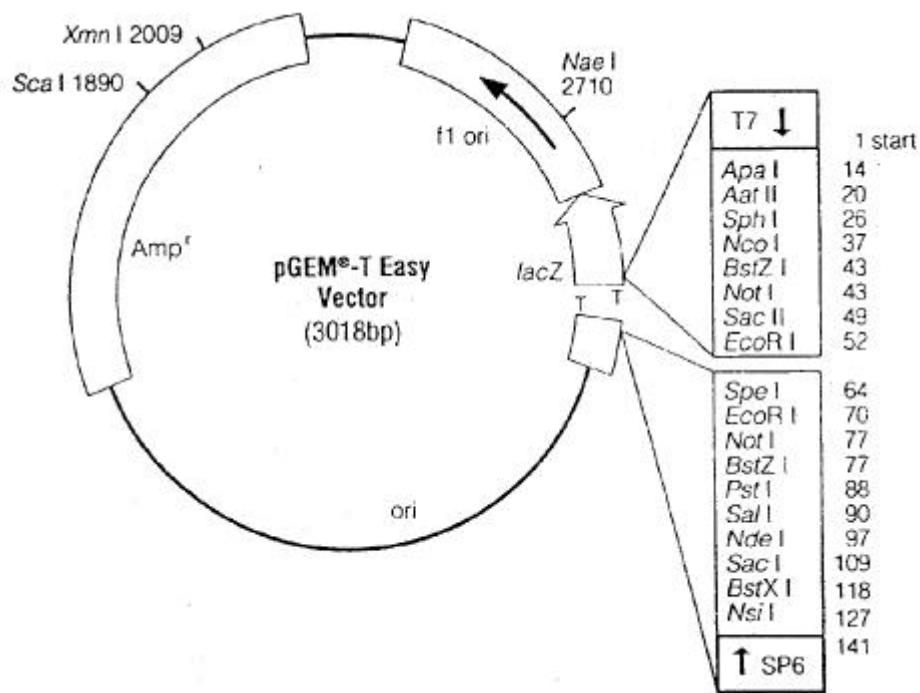
#### 7. callus

70% 30 2% sodium  
 hypochlorite 15  
 . (NAA, BA) MS,  
 WPM, GD callus .

#### 8. Plasmid DNA

↗. Cloning plasmid DNA  
 Cloning colony ampicillin (50  $\mu\text{g}/\mu\text{l}$ ) ↗ LB 37  
 saturation , Wizard<sup>TM</sup> Minipreps DNA Purification  
 System (Promega) plasmid DNA . 1.5 Ml  
 eppendorff tube 15,000 rpm, 1 cell  
 resuspension solution (50 mM Tris-HCl, pH 7.5, 10 mM EDTA, 100  $\mu\text{g}/\text{Ml}$ )  
 , 200  $\mu\text{l}$  cell lysis solution

200  $\mu\ell$  neutralization solution  
 15,000 rpm 5 1 M $\ell$   
 Wizard<sup>TM</sup> Minipreps DNA



### 2.1. pGEM-T easy vector

pGEM-T vector pGEM-5zf(+) DNA *Eco*R ,  
 thymidine . 3'-T overhang PCR ligation  
 ↗ . purification resin , resin/DNA  
 Wizard<sup>TM</sup> minicolumn elution DNA . 2 M $\ell$   
 column wash solution , 4 , 15,000 rpm 2  
 30 . 60 20  $\mu\ell$  DNA

4 , 15,000 rpm, 2 DNA .  
 . Sequencing plasmid  
 Cloning colony ampicillin (50  $\mu$ g/ $\mu$ l) 2 M $\ell$  LB  
 37 saturation 500 M $\ell$  LB  
 overnight . centrifuge tube 5 ,  
 4 , 7,000 rpm, 5 18M $\ell$  TEG (50 mM glucose, 10 mM EDTA, 25 mM Tris-HCl, pH 8.0)  
 (lysozyme 10 mg / TEG M $\ell$ ) 2 M $\ell$  2 , 10  
 . 40 M $\ell$  (0.2 N NaOH, 1% SDS)  
 , 10 . 30 M $\ell$  3 M potassium acetate (pH 5.5) 15 , 4 , 8,000 rpm, 5  
 . (cheese cloth) tube  
 60 M $\ell$  2 isopropanol 2 , deep freezer (-70 ) 30  
 . 4 , 12,000 rpm, 15 10  
 4.3 M $\ell$  2 , 4.8 g  
 CsCl, 0.5 M $\ell$  EtBr (10 mg/M $\ell$ ) , 4 , 12,000 rpm  
 15 . 18 gage  
 tube , (Beckman) , 20 ,  
 50,000 rpm, 17 21 gage  
 tube , 18 gage plasmid  
 DNA tube , 20 SSC isopropanol  
 EtBr , TE  
 buffer [10 mM Tris (pH 8.0), 1 mM EDTA]  
 DNA DNA sequencing

9.

Boehringer Mannheim . DNA TE  
buffer 37 1 3 . 65  
80 10 , 1/6  
(50% glycerol, 0.1 M EDTA, 0.025% bromophenol blue, 0.025% xylene cyanol)

10. Southern hybridization

, marker  
southern hybridization . probe recombinant plasmid  
DNA *Eco*R Gene clean kit agarose gel  
DNA DIG DNA labelling and detection kit  
(Boehringer Mannheim Co.) labeling probe .  
probe DNA 15  $\mu\ell$  95 10  
3 . 2  $\mu\ell$  hexanucleotide mixture, 2  $\mu\ell$  dNTP labelling  
mixture, 1  $\mu\ell$  Klenow enzyme (100 U/ $\mu\ell$ ) 7 37 60  
. 2  $\mu\ell$  0.2 M EDTA (pH 8.0), 2.5  $\mu\ell$  4 M LiCl, 75  $\mu\ell$  7  
ethyl alcohol 7 -70 30 .  
(4 , 15,000 rpm, 15 ) 50  $\mu\ell$  70% ethyl alcohol  
DNA . DNA 50  $\mu\ell$   
TE buffer probe .  
agarose gel gel denaturation solution (1.5  
M NaCl, 0.5 N NaOH) 45 . Gel  
, neutralization solution [1 M Tris (pH 8.0), 1.5 M NaCl]  
45 . Nylene membrane (Boehringer

Mannheim)  $\gamma$  10 $\times$  SSC (1.5 M NaCl, 0.15 M Na-Citrate,  
 pH 7.0) 10 . Gel nylone membrane  
 , 2 $\times$  SSC (0.3 M NaCl, 30 mM Na-Citrate) paper  
 . 10 cm paper towel 500  
 g 8 . DNA transfer . 5 $\times$  SSC  
 (0.75 M NaCl, 75 mM Na-Citrate) membrane 5  
 , 30 . 80 1 2  
 .  
 DNA membrane 20 M $\ell$  prehybridization  
 [5 $\times$  SSC, 0.02% (w/v) SDS, 0.1% (w/v) N-lauroylsarcosine, 1% blocking  
 reagent] . 68 30  
 probe DNA (5-25 ng/M $\ell$ ) 68 16  
 . Nylon-membrane Solution (2 $\times$  SSC,  
 0.1% SDS) 5 , 68 Solution (0.1 $\times$  SSC,  
 0.1% SDS) 15 . Washing buffer (0.1 M maleic acid,  
 0.15 M NaCl, pH 7.5, 0.3% Tween 20) 1 5 blocking  
 solution (0.1 M maleic acid, 0.15 M NaCl, 1/10 10X blocking solution)  
 30 . 20 M $\ell$  blocking solution antibody solution 75  
 mU/buffer . 100 M $\ell$  washing buffer 15 2  
 , 20 M $\ell$  detection buffer (0.1 M Tris-HCl, 0.1 M NaCl, 50 mM MgCl<sub>2</sub>,  
 pH 9.5) 2 5 . Membrane hybridization bag 20  
 CSPD ready-to-use solution ,  $\gamma$   
 5 . ,  $\gamma$   
 37 5 15 , X-ray film .

11. Primer

4  $\mu$ g DNA 18  $\mu$ l , 2  $\mu$ l 2M NaOH, 2 mM EDTA  
, 5 . 75  $\mu$ l 99.9% ethylalcohol  
, -70 10 . 4 , 14,000 rpm 10  
20  $\mu$ l が 70% ethylalcohol 1 .  
18  $\mu$ l . ALF-Express  
automatic sequencer (Pharmacia Biotech) sequencing .  
sequence homology http://www.nebi.  
nlm.nih.gov/BLAST/ Basic Local Alignment Search Tool (BLAST) program  
, marker Primer  
Primer3 Input (http://www.genome.wi.mit.edu) site , primer  
가 , ( ) primer

### 3

#### 1. genomic DNA

CTAB (cetyltrimethylammonium bromide) spermine- spermidine

genomic DNA 1 g 1.6 mg

genomic DNA . CTAB

(Rogers and Bendish, 1988) Dellaporta genomic DNA

, , DNA

genomic DNA 가

#### 2. PCR

RAPD marker PCR 가

annealing , 36 55 가

(Williams , 1990; Hu and Quiros, 1991; Demeke , 1992; Kim , 1997).

, , 90% RAPD marker

annealing 36 (Koller , 1993; Adams , 1993) , 37 (Bellamy , 1996; Lee , 1995) denaturation 94 ,

extension 72 , annealing 37 , 42 , 50

PCR , 37 가 band

pattern .

cycle 37 annealing , 30cycle,

35cycle, 40cycle, 45cycle , 45cycle

PCR DNA band pattern

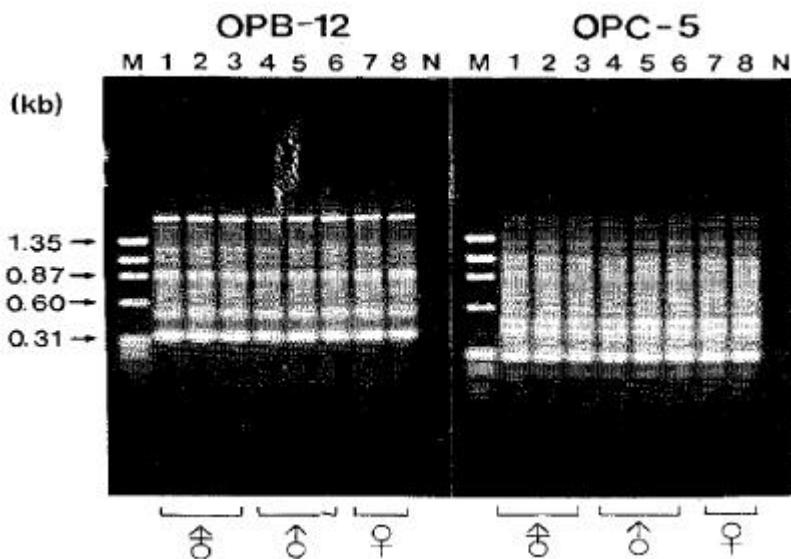
10 ng, 15 ng, 20 ng, 30 ng, 50 ng, 100 ng DNA が PCR  
, 10 ng, 15 ng band が , 100 ng  
band smear band pattern . (Adams , 1993,  
(Koller , 1993), (Lee , 1995) 30 ng DNA  
band pattern .  
PCR annealing 37 , 45cycles , DNA  
30 ng/25 μℓ , 3

3. ,  
primer 10-mer OPA kit (20 primers), OPB kit  
(20 primers), OPC kit (20 primers), OPD kit (20 primers), OPE kit (20  
primers), OPF kit (20 primers), OPG (20 primers), OPH kit (20 primers),  
OPI kit (20 primers) 180 primer .  
2 1 RAPD  
2-2 , 180 primer 9 が primer  
, 180 primer , , 3  
band pattern . OPD-15 primer  
800 bp, 2,500 bp, 1,900 bp  
band が . 9 primer が  
band が 128 , 1 10 band pattern  
band pattern 150-3,000 bp .  
,

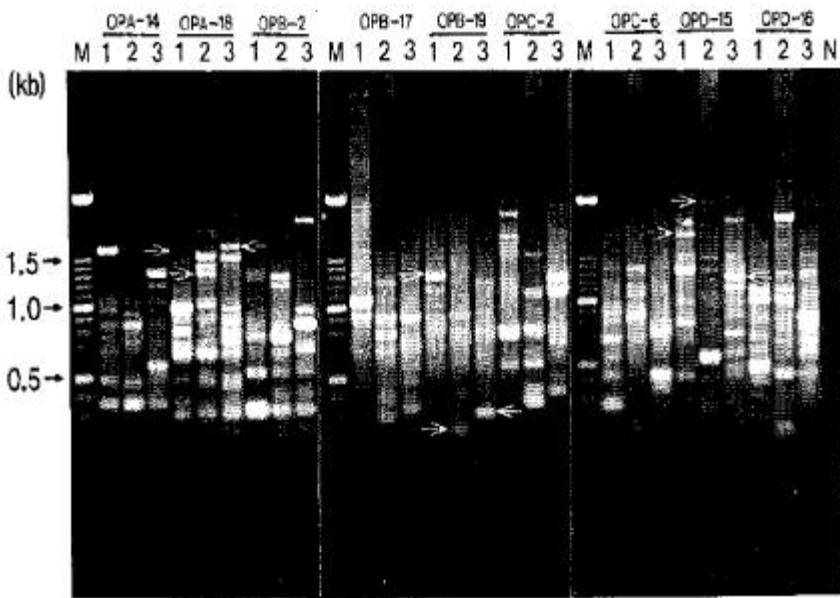
, , ,  
가  
가 .

4. marker

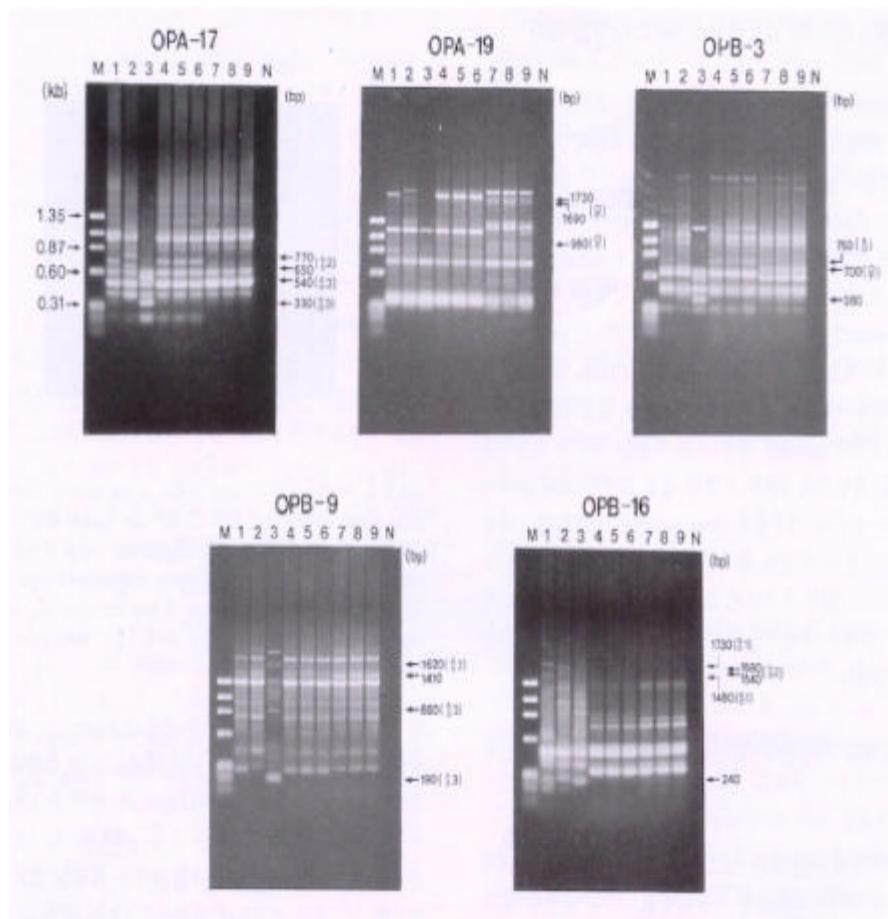
, , , 3 180 primer  
RAPD , 174 primer , ,  
primer band pattern ( 2.3).  
6 random primer ( 2.1) polymorphic band pattern  
band 가 ( 2.4, 2.5).



2.2. Primer OPB-12 OPC-5 PCR . Lane M: size marker, *Hae* X174 DNA; lane 1-3: 6, 41, 51; lane 4-6: 46, 52, 53; lane 7-8: 42, 54; N:



**2.3. Primer PCR .** Lane M: size marker 100 bp ladder; lane 1: ; lane 2: ; lane 3: ; lane N: ; band .



#### 2.4. Primer OPA - 17, OPA - 19, OPB - 03, OPB - 9, OPB - 16

PCR . Lane M: *Hae*

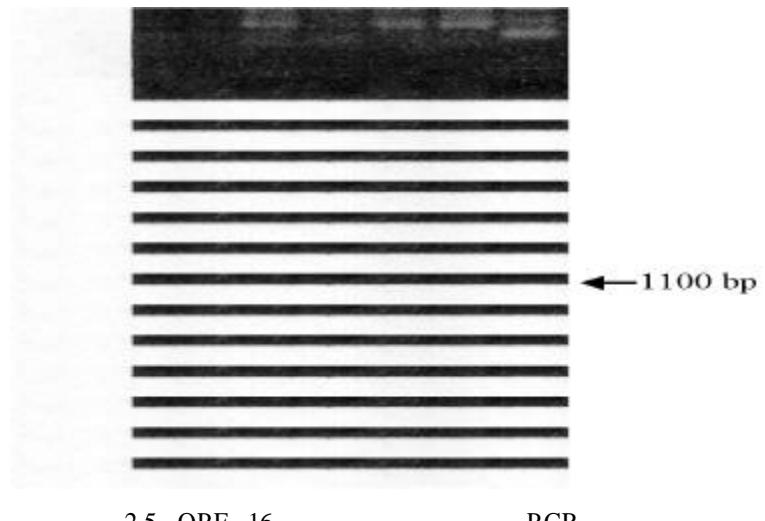
X174 DNA; lane 1-3:

6, 41, 51; lane 4-6:

46, 52, 53; lane 7-8:

42,

52, 54; lane N:



2.5. OPF- 16

PCR

Lane M: DNA      *EcoR*      *Hind*      DNA; lane 1:  
 6; lane 2:                  41; lane 3:                  51; lane 4:  
 42; lane 5:                  46

#### 2.1. RAPD primer

Primer number	Nucleotide sequence (5' to 3')	GC content(%)
OPA - 17	G A C C G C T T G T	60
OPA - 19	C A A A C G T C G G	60
OPB - 03	C A T C C C C C T G	70
OPB - 09	T G G G G G A C T C	70
OPB - 16	T T T G C C C G G A	60
OPF - 16	G G A G T A C T G G	60

marker OPA - 17 770 bp, OPB - 03 1360 bp,  
marker OPB - 03 980 bp, OPA - 19 1,300 bp,  
6 marker OPA - 17 1,340 bp, 41 marker  
OPB - 09 609 bp, OPF - 16 1,115 bp, 51 marker OPA - 17  
580 bp ( 2.4, 2.5).

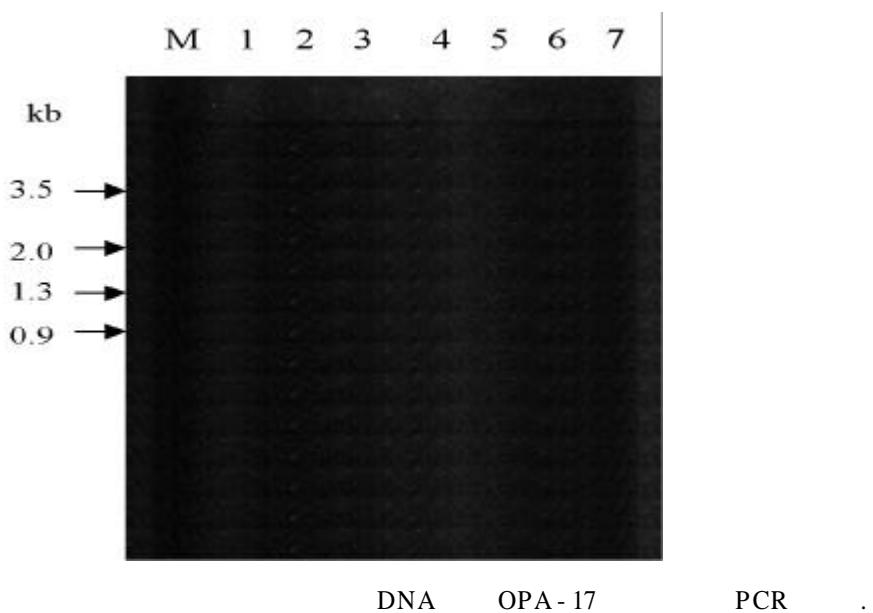
6 primer 87 가 band 가 ,  
primer 14.5 band 가 . band 200-2,000 bp  
. 가 ,  
band pattern

6 OPA - 17 primer 680 bp band  
, 41 OPA - 19 primer  
2 band pattern (1,200 bp, 1,350 bp) . 51  
6 , 41 , band

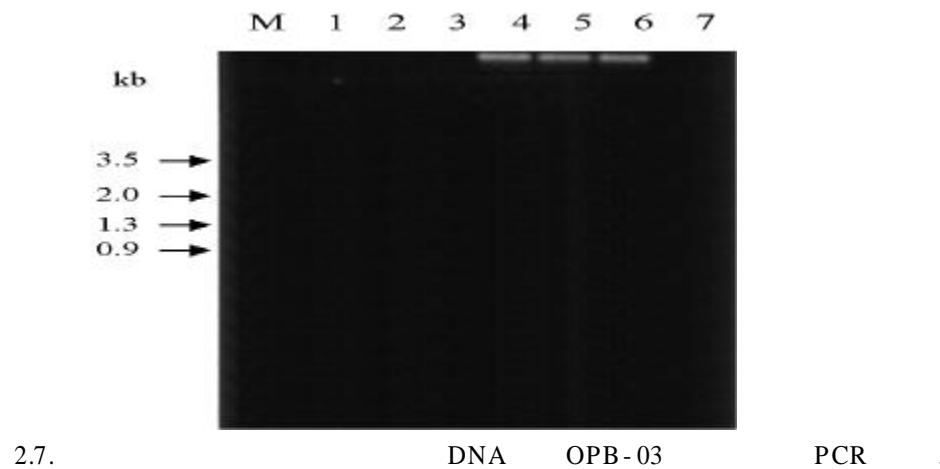
가 .  
phylogenetic relation cluster 2.3  
180 primer 174 primer monomorphic band  
pattern 가 .

##### 5. marker cloning

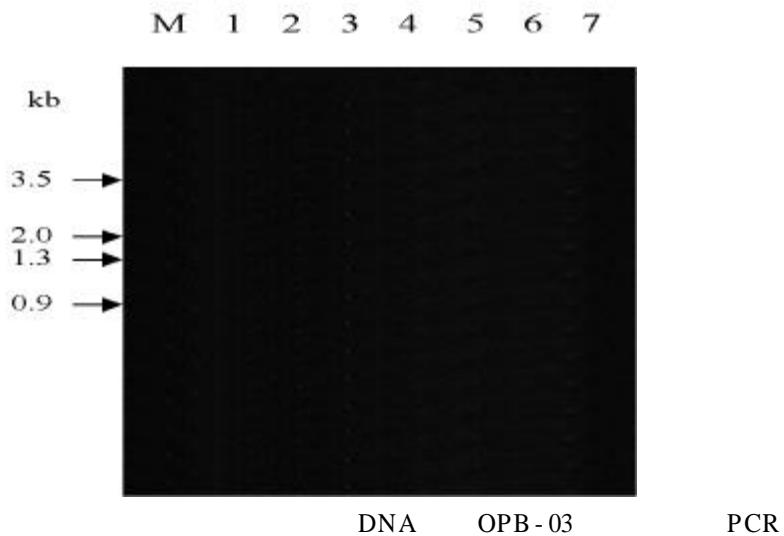
RAPD genetic marker 가 band GENECLEAN kit  
pGEM-T Easy vector system cloning  
marker OPA - 17 789 bp ( 2.6), OPB - 03 1,360  
bp ( 2.7), marker OPB - 03 749 bp ( 2.8),



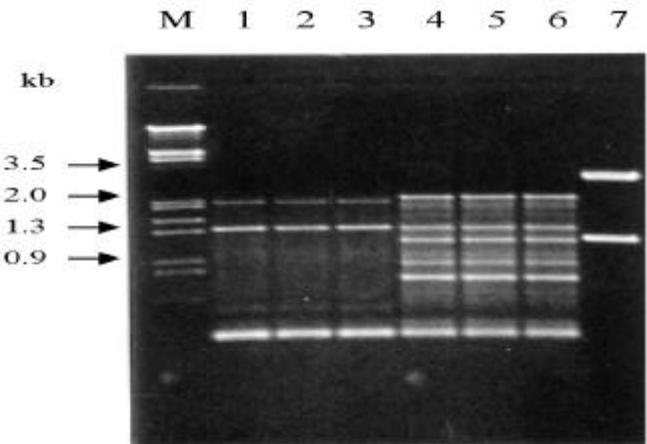
2.6. DNA OPA - 17 PCR  
 Lane M, DNA *Hind* - *EcoR* DNA; lane 1-3:  
 42, 52, 53; lane 4-6: 46, 54, 55. lane 7: *EcoR*  
 plasmid DNA



Lane M:    DNA    *Hind* - *EcoR*                          DNA ; lane 1-3,  
 42, 52, 53; lane 4-6,                  46, 54, 55; lane 7, *EcoR*  
 plasmid DNA.

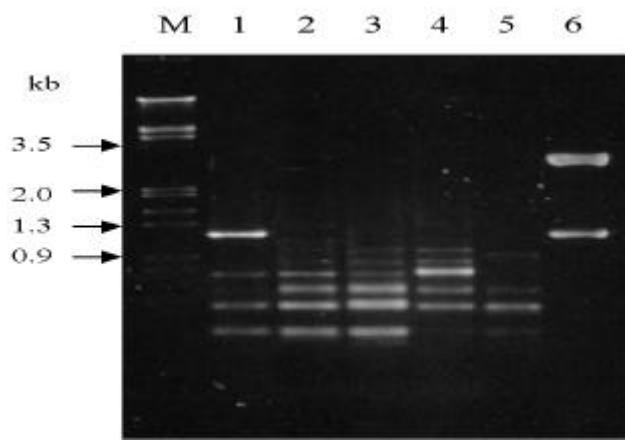


Lane M:    DNA    *Hind* - *EcoR*                          DNA ; lane 1-3,  
 42, 52, 53; lane 4-6:                  46, 54, 55; lane 7: *EcoR*  
 plasmid DNA



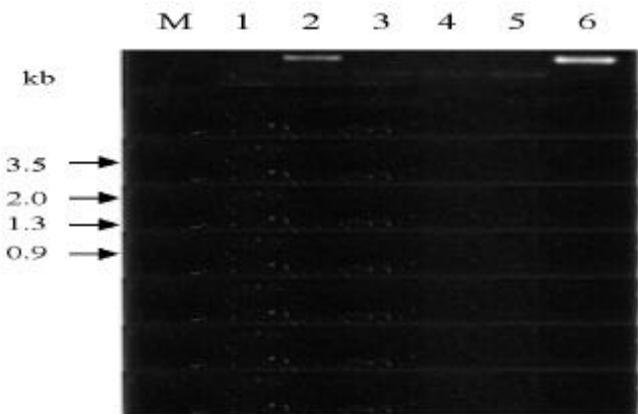
2.9. DNA OPA - 19 PCR

Lane M: DNA *Hind* -*EcoR* DNA; lane 1-3:  
 42, 52, 53; lane 4-6: 46, 54, 55; lane 7: *EcoR*  
 plasmid DNA



2.10. 6 DNA OPA - 17 PCR

Lane M: DNA *Hind*-*EcoR* DNA; lane 1-3:  
 6, 41, 51; lane 4, 46; lane 5: 42; lane 6: *EcoR*  
 plasmid DNA

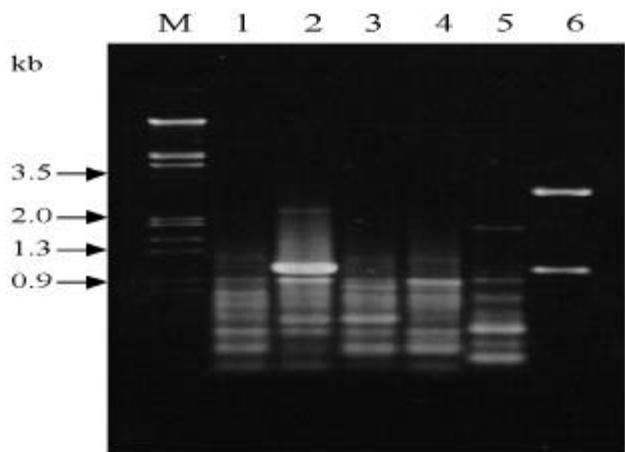


2.11. 41 DNA OPB - 09 PCR .

Lane M: DNA *Hind* - *EcoR* DNA; lane 1-3:

6, 41, 51; lane 4: 46; lane 5: 42; lane 6: *EcoR*

plasmid DNA

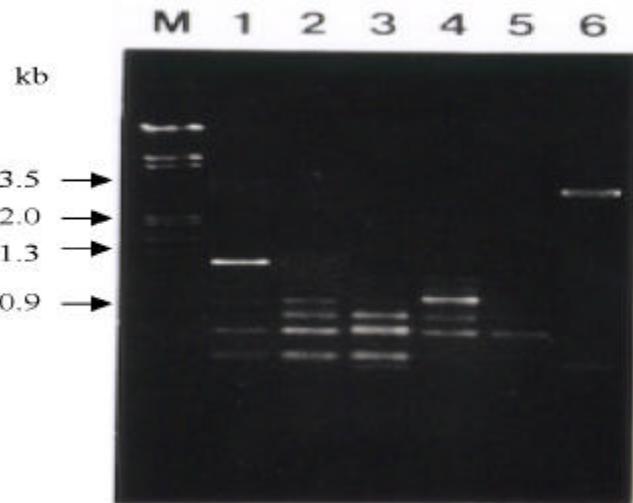


2.12. 41 DNA OPF - 16 PCR .

Lane M: DNA *Hind* - *EcoR* DNA; lane 1-3:

6, 41, 51; lane 4: 46; lane 5: 42; lane 6: *EcoR*

plasmid DNA



2.13. 51 DNA OPA - 17 PCR .

Lane M: DNA *Hind* - *EcoR* DNA; lane 1-3:

6, 41, 51; lane 4: 46; lane 5: 42; lane 6: *EcoR*

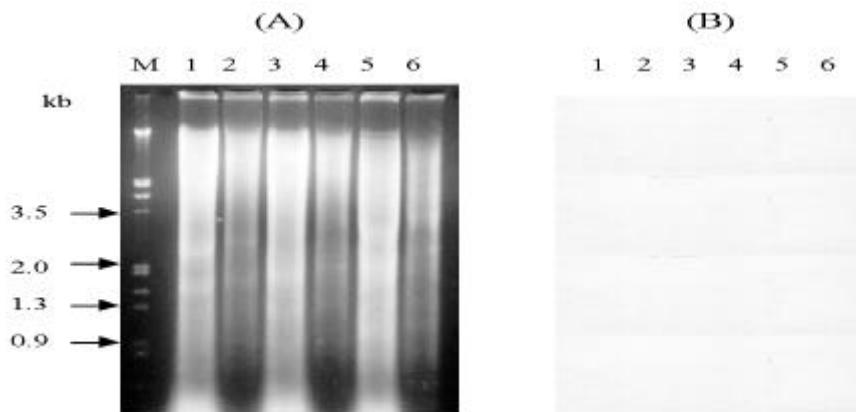
plasmid DNA

6. marker

RAPD marker ,  
homology southern hybridization .  
marker† cloning plasmid , *EcoR* probe

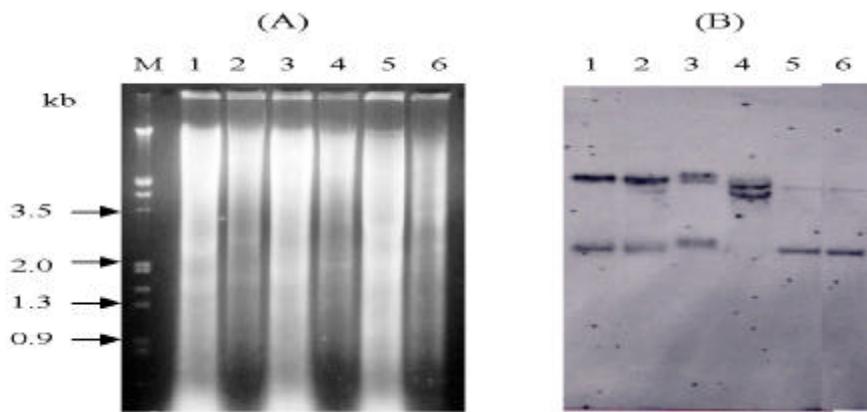
marker OPA - 17 789 bp DNA  
hybridization , 2.14 band†  
genetic marker † , †  
marker † . OPB - 03 1,360 bp  
DNA probe 2.15  
marker † .

marker OPB - 03 749 bp probe  
 , , genetic marker ♂  
 ( 2.16).  
 6 marker OPA - 17 1,340 bp  
 hybridization , , , marker  
 ♂ (data not shown). 41 RAPD marker  
 OPB - 09 600 bp  
 genetic marker ♂ ( 2.17). OPF - 16 1,100 bp DNA probe 41  
 51 ( 2.18). 51 genetic marker  
 OPA - 17 580 bp probe hybridization  
 , 2.19 51 .

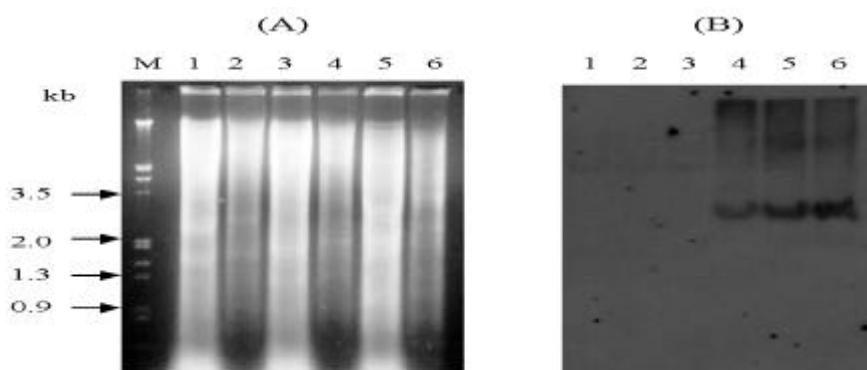


#### 2.14. OPA - 17 PCR

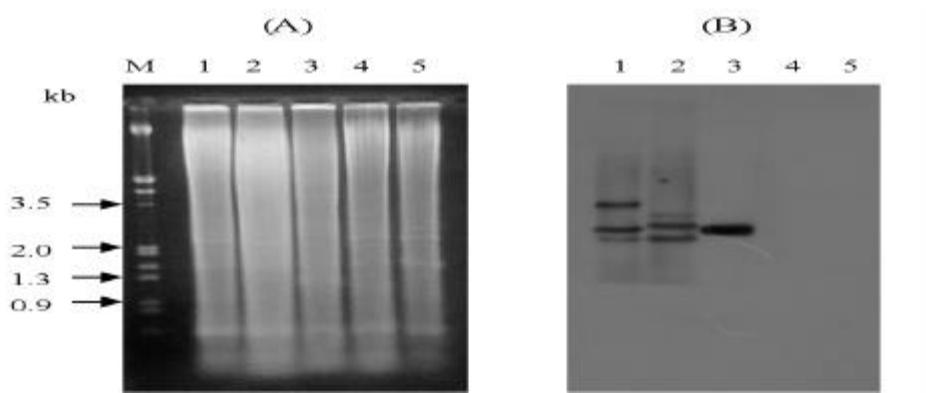
marker probe southern hybridization.  
 Lane M: DNA *Hind* - *EcoR* DNA; lane 1-3:  
 46, 52, 53; lane 4-6: 42, 54, 55; (A) DNA  
*Hind* , 0.8% agarose ; (B) southern  
 hybridization X-ray film



2.15. OPB-03 PCR  
 marker probe southern hybridization.  
 Lane M: DNA *Hind* -*EcoR* DNA; lane 1-3:  
 46, 52, 53; lane 4-6: 42, 54, 55; (A) DNA  
*Hind*, 0.8% agarose gel ; (B) southern  
 hybridization X-ray film .



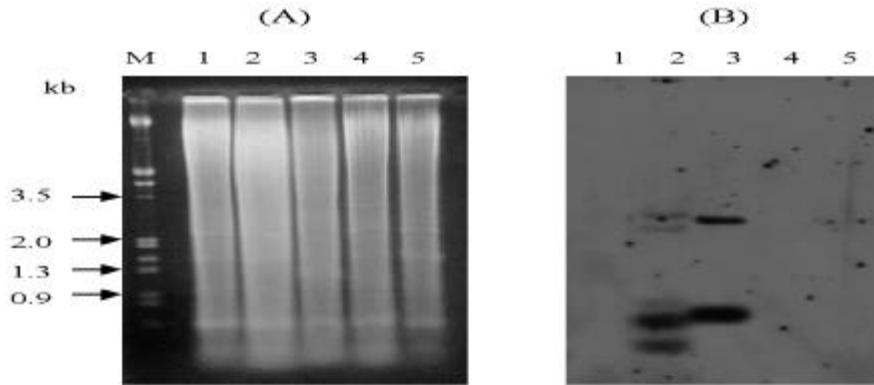
2.16. OPB-03 PCR  
 marker probe southern hybridization.  
 Lane M: DNA *Hind* -*EcoR* DNA; lane 1-3:  
 46, 52, 53; lane 4-6: 42, 54, 55; (A) DNA  
*Hind*, 0.8% agarose gel ; (B) southern  
 hybridization X-ray film



### 2.17. OPB - 09                          PCR                          41

RAPD marker    probe                          southern hybridization

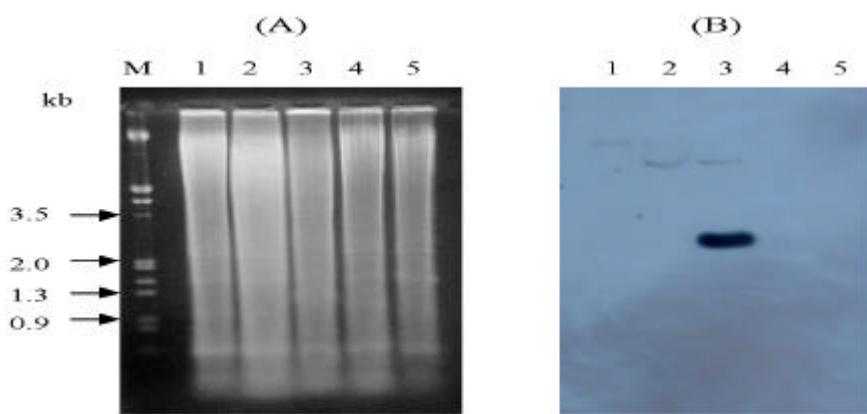
Lane M:        DNA    *Hind* -*EcoR*                          DNA; lane 1:        6;  
 lane 2:        41; lane 3:        51; lane 4:        46; lane 5:        42;  
 (A)        DNA    *EcoR*                          , 0.8% agarose gel  
 ; (B)        southern hybridization                          X-ray film



### 2.18 OPF - 16                          PCR                          41

RAPD marker    probe                          southern hybridization

Lane M:        DNA    *Hind* -*EcoR*                          DNA; lane 1:        6;  
 lane 2:        41; lane 3:        51; lane 4:        46; lane 5:        42;  
 (A)        DNA    *EcoR*                          , 0.8% agarose gel  
 ; (B)        southern hybridization                          X-ray film



## 2.19. OPA - 17

PCR

### southern hybridization.

41

Lane M: DNA *Hind*-*EcoR* DNA; lane 1: 6;  
 lane 2: 41; lane 3: 51; lane 4: 46; lane 5: 42;  
 (A) DNA *EcoR*, 0.8% agarose gel  
 ; (B) southern hybridization X-ray film

7. kit 가

1) marker

## Southern hybridization

DNA

2.20

789 bp

open reading frame 가

## homology

749 bp

homol

DNA

가

ORF

homology

(2.21).

10	20	30	40	50	60	70	80	
5'	GACCGCTTCT	TGTTGGAACT	CTTTGGAATA	ACCTAAGTC	GTGAGGGTAG	ACTACTCTTG	GCATTTACAT	CAGCATGTGT
90	100	110	120	130	140	150	160	
GTATAAGTTT	<u>CA</u> <u>GGAAATTG</u> A	<u>G</u> <u>TCACACT</u> A	<u>T</u> <u>CCATCAATA</u>	TCACATGTAA	ATTCATTCTAT	TATTTCTATT	TATGITCTCT	
170	180	190	200	210	220	230	240	
TATAATTATA	TTAACGTATC	TACTGAAAAA	ATCTCTATAA	AATATTTGTA	AGTATCCAAT	ATTTGTTGAT	TGTATTGTGT	
250	260	270	280	290	300	310	320	
TGAAATCAAA	ATACTAAAAAA	ATTGTGTTAT	CTAGTCACG	TGAGTCACAT	TTCAGGTCG	TGATCAAAAT	AGGCACGTAA	
330	340	350	360	370	380	390	400	
TGTCATCTAT	TTCTAGTTTT	GAGTCITACA	TAATATATTA	GAGTATATTG	GATTCAATTG	GCATGTTGGT	TTTTGGAATT	
410	420	430	440	450	460	470	480	
TTATCTCAT	TTTAAAATT	TGAAAATTG	GTGCAAGGCA	GTCGACCACT	CACTTGAA	TGGTAGAAC	GTCATGGTGA	
490	500	510	520	530	540	550	560	
GCTTCGGTTT	CCTAACATTA	AAAATAGGT	GTTATGTTGG	TTTGGTCATT	<u>TG</u> <u>GA</u> <u>ACTCAA</u>	<u>GAGACCAAGT</u>	<u>GACCC</u> TTGCT	
570	580	590	600	610	620	630	640	
CCCCAACATT	GGTAGGITAC	CTTCCTTGCT	TCTTGTGTTTC	ACTTCCTTTC	GCTCTTTCA	AGTTTCGIG	CTAGGGTTTC	
650	660	670	680	690	700	710	720	
TATTGATT	CACTATTTCT	ACCCTAGATT	GTTTTTCATA	GCTCTCTAAC	CITCTAAGAA	GTCAAAGTCC	TCTTCGGGTA	
730	740	750	760	770	780			
CCTATGATAC	ATGCAAGTT	GTGCCCCTC	GTGTCACAGCT	AGGCACAAAT	AATATGATCA	CAAGGGTCA	3'	

## 2.20. marker .

10	20	30	40	50	60	70	80	
5'	GGACTGCAGA	CATCGATTAC	ATAACGAAAA	TCACCATAGA	AACCTCTCTT	TCITATAGGC	GTACAAAGAG	GGTCCCAG
90	100	110	120	130	140	150	160	
ATCAGGATAT	AGGITGAGAA	CCACTACGAA	GGACTACCGG	CAAAAGAGTA	AATCATCGAT	CGACTCAATT	CTTTCTTATC	
170	180	190	200	210	220	230	240	
TTGGTCTAGA	AAGAGAGGCC	CTGAAGTAAC	TGGAATAGT	GCAGAAAAAA	GCTTTACCAT	CAAGCCGCTT	GCTTTGTCT	
250	260	270	280	290	300	310	320	
CTGCCCTCTT	TTGTGAGAAAT	TTATCGGTAC	CGTACACACC	CTGAAGGGG	GTGGAG <u>C</u> <u>T</u>	<u>G</u> <u>TG</u> <u>CC</u> <u>CT</u> <u>AT</u>	<u>T</u> <u>CT</u> <u>TC</u> <u>ACT</u> <u>G</u> A	
330	340	350	360	370	380	390	400	
CCACAGTGG	CTGACTTTGC	TTTGTATGAT	TTTCCTTGC	CCAAGACTGA	CATTCGTTTC	GTTTGAGTG	ATGAACTGCT	
410	420	430	440	450	460	470	480	
GTTGTTAAC	TGTGTAAC	AAAAAATAAA	AGTAGATGTT	TGATGTTAGA	GAAGTCCTT	CTATGATTCC	TTACCGAAC	
490	500	510	520	530	540	550	560	
GTATGCC	GAGITGTTTT	CCTTCACAA	GACAAGGGG	TGTGCTCCCA	TCCCTCCGG	GATCATAGGA	AAGATGGAGT	
570	580	590	600	610	620	630	640	
GATTCTCCAT	CTGGCTCTA	CCTATTTAAG	AGAGTCGATA	TTTCACTTTA	AAGACAGCT	GTGCCCTACC	TACCAAGATA	
660	660	670	680	690	700	710	720	
AATACAGTCT	TACCTGTCCT	AGCTTGATAG	AAACCTAGCT	ACAAAAGCGA	CITCCCCCTG	TTGAAAATAC	CAG <u>TG</u> <u>GAG</u> <u>G</u>	
730	740							
<u>AGTACCTGGT</u>	<u>G</u> <u>T</u> <u>G</u> <u>C</u> <u>T</u> <u>A</u> <u>C</u>	CTGCAGTCC	3'					

## 2.21. marker .

41 RAPD marker RAPD marker OPB - 09 609  
 bp 2.22, OPF - 16 1,115 bp 2.23  
 . Southern hybridization , 偈 609 bp  
 frame stop codon ORF偈  
 . Homology  
 retroviral protease reverse transcriptase  
 retroviral molecular fossil .  
 Southern hybridization 3 51  
 580 bp 2.24 .

	10	20	30	40	50	60	70	80
5'	TCATCCCCCT	GCCCTTTCA	AGCCITCITC	TTCTCACATA	AGATACACAT	GCCITAGACA	ATTTTGGAA	GGCAAAGCC
	90	100	110	120	130	140	150	160
TTAGGCAAGT	TTGGGAGGTT	TTCITCAGTT	GGCACGAAAA	CGGACCCGAA	CGGCAGGATT	TTTGAAACAT	GAGCITTTCA	
	170	180	190	200	210	220	230	240
TGAATTTITC	AGATTOGCCCT	TCGGAATGG	GTGGIT <u>GCC</u>	<u>ATTTTGATGC</u>	<u>TCCATT</u> ATC	GAACGGATCA	TTTCAATT	
	250	260	270	280	290	300	310	320
TTTGGGIGGGT	CCCTTTAGGA	TTCATCAAGA	ATAGTCCTCT	TTTACGAAGA	ACGAACCGAC	CGACGGGTT	TTTCAAACITG	
	330	340	350	360	370	380	390	400
CATCTTTITC	TGAATTTTT	GAACCATGG	ACTCCATGAG	TGATTGGATC	ATTTTGAAAT	TTTCTACCG	TCGCTGCCT	
	410	420	430	440	450	460	470	480
TATGCTTAAA	GFGCTTCTCA	TGCCCTAGCC	AATTTAGAG	GCTTTGACC	ATGGACATGG	AGTGCTCAT	GACTAAGCAA	
	490	500	510	520	530	540	550	560
ATTTTGGCC	CCAACAATGG	GATTCAACAA	CITAAAAAAC	TTCTCCAAA	TTCTTTAAG	ATGAAAGATT	TAGGACCAT	
	570	580	590	600				
AACITTATTTC	TTGGACITAG	AGCTATAT <u>AT</u>	<u>TGATACACCA</u>	<u>GGGGATGA</u>	3'			

2.22. marker .

	10	20	30	40	50	60	70	80
5'	TGGACTGCAG	ACTGGCCTAA	TTCGCCATA	ATATAGTAGT	CAACATTGTA	TGATGCAGCA	ACAAGGCCA	CTTGAGAOct
	90	100	110	120	130	140	150	160
TGAATCGTAA	GCCCCATGAA	GGTGTGAAG	GGCCTGTTCT	TGGTATAGCA	GTTAGCTCG	AAGTTCATCA	AAGGITAACG	
	170	180	190	200	210	220	230	240
ACGAATTATT	GTTGGTGACC	GTCCTGATAA	AGGATTCATCA	TTCTTCAGGG	AGTCCCTGCCA	GTGCATAGAT	CACCATGTCT	
	250	260	270	280	290	300	310	320
TTAGAAGAAA	GG AGAGTT	TATGGCTCT	AGGAATTTC	CA <u>ATGGACCG</u>	<u>AAGATAACG</u> IT	<u>CC</u> CATGGGTT	GGATCCCTTT	
	330	340	350	360	370	380	390	400
TTCAAAGATT	GAAATTTGAG	TTGAGATGTA	ACTCACGACG	AGTAGGCTGG	TCTAAAAATC	TTTTTTACAA	GGCTAACCAA	
	410	420	430	440	450	460	470	480
AATATCACGA	CAGGGTGICA	GTTCATGAAC	CTCTTGAAGA	ATCTCTTTG	TAAGAGITGC	ATTGATCCAT	GATAAGGAGC	
	490	500	510	520	530	540	550	560
TTTGGTCTGT	CCGAACCCAA	GGCGAATATG	CAGGGTTTG	TATGGCGTTG	CCATTGGCAT	AACAAATAAA	CTGAGGAGGA	
	570	580	590	600	610	620	630	640
CTGGGAAACG	TTCCGTGAC	AAAGCCCAGA	AAAGAATTTC	AGATTATAAT	GTAGGAATTG	GGATTCAAA	AGAAGATAAT	
	650	660	670	680	690	700	710	720
TGGTAGAGTC	AAGTTTAAGG	GAAACAAGAT	GGGTGATGTT	TGGTGTAACT	ACCATAGGAT	TOGGGGTCTC	TTCCGTGCT	
	730	740	750	760	770	780	790	800
GAAGAACCGA	GCATGAGGGA	TTGATGAAAG	AGAACCCATA	GTGTTGTCAA	ACCACCTTGG	AATACTACAA	TCAGAGCCTC	
	810	820	830	840	850	860	870	880
CATAGCTTTC	CGTTTCTATC	TATCATAAACG	GGAAAGGGAC	CAAAAAATCC	CTCAAAGTGT	GGAACAAGGC	TAACCTCGT	
	890	900	910	920	930	940	950	960
AAGAAGATGT	TCGATGCAAG	TGCTGTGAA	TGGAGAGAAA	AGATAGGACA	GAGAAGGAGA	AAGAGTGAAG	TTCTGGTCTT	
	970	980	990	1000	1010	1020	1030	1040
TCTACTAGGT	<u>CCTCCCTTCG</u>	<u>AGCCTTCAGA</u>	<u>TCAACTCGTT</u>	<u>GGGAACGAA</u>	<u>TGTTCTGGTT</u>	<u>GTATTGAGC</u>	<u>CCTCTACCGC</u>	
1	1050	1060	1070	1080	1090	1100	1110	1115
TOGCCCTTGCC	TCGAAGCCCG	TGGAAGGTTCA	AAAGCAATAA	ACTCAGGCCCT	CITAACCITC	TCCATCTCCA	GTCCA	

2.23.

6 41

.

	10	20	30	40	50	60	70	80
5'	TGACCGCTTG	TGGTAAGAGA	GATCCCTCCG	AAGCCCTTT	CGCTTCGCTC	AAGGGCGGAA	GCTCAGGAGG	ACATGCCCG
	90	100	110	120	130	140	150	160
TCCTTTGAA	TGACTTCTGG	TTTTCGCAA	GGCCCTGTATT	TCTTTGAA	ACCATATCAT	TTTGTATTTG	AGGGTGAAGC	
	170	180	190	200	210	220	230	240
CAACAGACTC	CGCAGTTTT	TGCATTGCTA	TGTTCTT <u>ACC</u>	<u>ACTCGTTCCC</u>	<u>AGTGTAG</u> CGG	ACCGCATGCC	GGGITCTTTG	
	250	260	270	280	290	300	310	320
ACCAACTGCA	AACCTGTCA	CAAAAGGGCG	CGGGCCAATC	CCAGAGCGTG	ATATTGTGGA	TGTGTGCGGA	CGGGATCCGT	
	330	340	350	360	370	380	390	400
GTITCCAATT	TTGCTTTCTT	CATTGAGCGA	GCAGATGCAA	TITGCCGCAA	TGTTCCGTC	TOGTG <u>CAACA</u>	<u>CCGACCAGAT</u>	
	410	420	430	440	450	460	470	480
<u>CAAGA</u> GACAG	CTCATATTCA	CTTGTGTTCA	TGATGACGAG	ACGATTTTCG	GGCGTCATGT	ATTCTGTGCC	AAAGGGGCCG	
	490	500	510	520	530			
CGATGCATCT	TTGCAACTGC	GTCTGCTTCT	TCGGFCCTA	CAAGCGGTC	3'			

2.24.

51

marker

2) primer

DNA

PCR primer                    primer  $\gamma$  , , ,  
( 2.3).

primer set                    PCR

460 bp,                  436 bp      PCR

primer                        southern hybridization

5 ,                  5            genomic DNA                    PCR

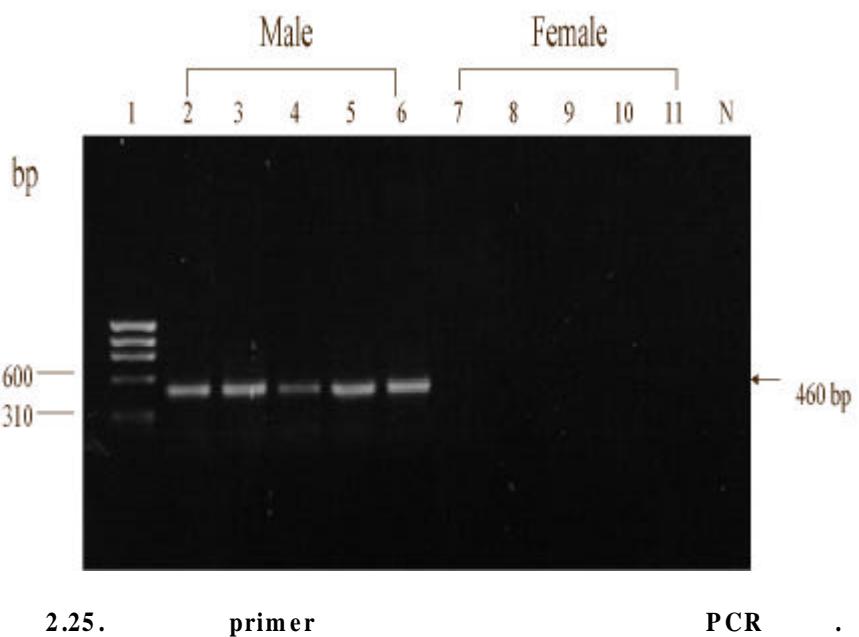
primer                        460 bp      PCR

( 2.25).                   $\gamma$                     primer  
436 bp      DNA                  ( 2.26).                    primer

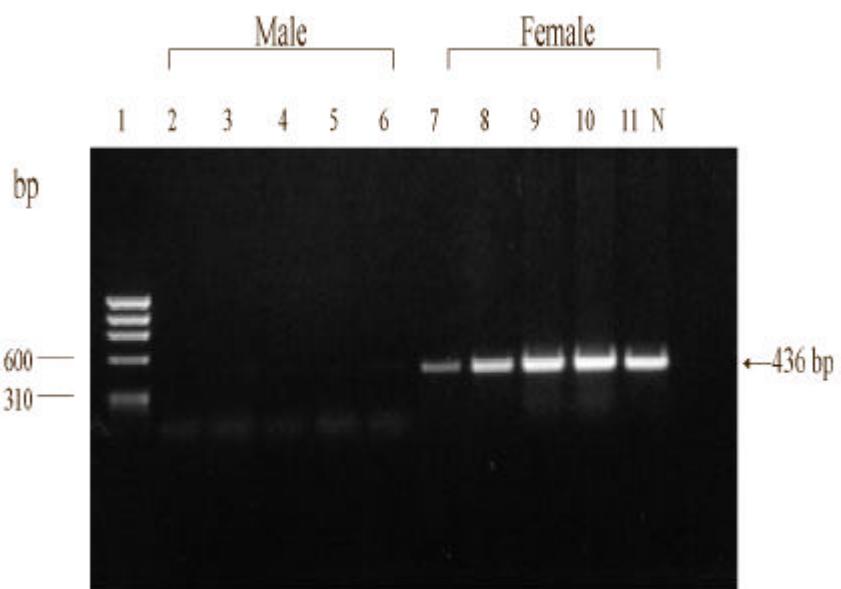
PCR marker

### 2.3. primer

	primer	primer	PCR product
male 1	F : 5'-GGAATTGAGITCACCTCATC-3' R : 5'-CACTTGGTCTCTTGAGTTCC-3'		460 bp
female 1	F : 5'-CTTGTGCGCTATTCTTCAC-3' R : 5'-AACACCAGCTACTCTCTCCA-3'		436 bp
41- 1	mono 2- 1	F : 5'-GCCATTTGATGCTCCATT -3' R : 5'-GTAGGGGGACCACATAGTT -3'	411 bp
41- 2	mono 2- 2	F : 5'-ATGGACCGAGATAACGTT -3' R : 5'-TCCGAGCTTCCCTCCTGGAT -3'	684 bp
51	3	F : 5'-ACCACTCGITCCCAGTTGAG -3' R : 5'-TCTTGATCTGGFCGCTGTTG -3'	207 bp



**2.25.** primer PCR



**2.26. Female primer** PCR .

## 2.27. Male 1 primer    Female 1 primer

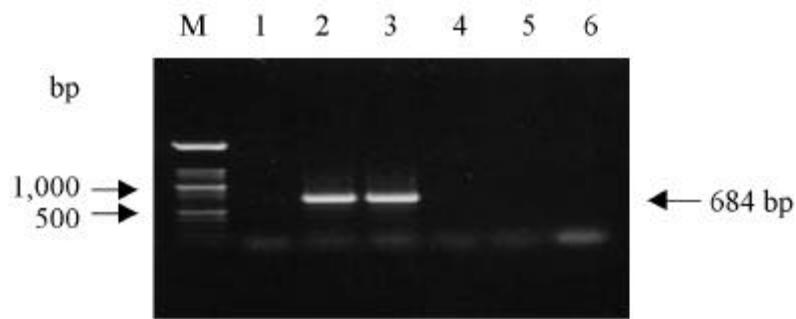
**PCR** . Lane M: size marker, *Hae* VI digest of X174 DNA; lane 1-3: 6, 41, 51 ; lane N,

41 RAPD marker 6, 41, 51  
 609 bp primer (mono2- 1)  
 PCR 2.28 684 bp  
 band ↗ . Southern hybridization ,  
 41, 51 primer PCR  
 , 884 bp band ↗ 41, 51  
 ( 2.29).



### 2.28. Mono 2-1 primer PCR .

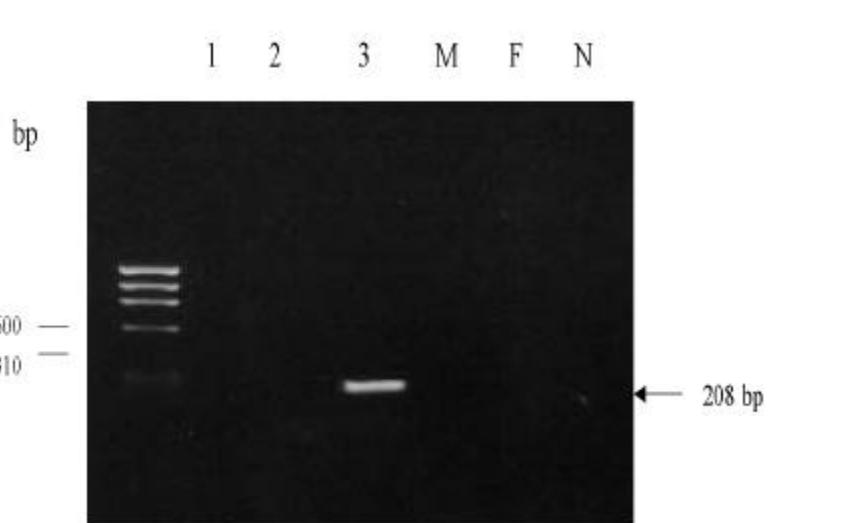
Lane M: size marker, 100 bp ladder; lane 1-3: 6, 41, 51 ;  
 lane 4-5: 46, 42; lane 6:



### 2.29. Mono2-2 primer PCR .

Lane M: size marker, 100 bp ladder; lane 1-3: 6, 41, 51 ;  
 lane 4-5: 46, 42; lane 6:

3                    51                    primer                    PCR                    ,  
 ,                    band  $\gamma$ †                    , 51  
 207 bp    band  $\gamma$ †                    (    2.30)                    51                    genetic marker  
 $\gamma$ †                    .



### 2.30. Mono3 primer                    PCR

Lane 1-3:                    6, 41, 51 ; lane M:                    46 ;  
 F:                    42 ; N:

RAPD                    southern hybridization  
 genetic marker                    OPA - 17 primer                    770 bp, OPB - 03  
 primer                    749 bp,                    OPB - 9  
 600 bp, OPF - 16    1,100 bp,                    51                    OPA - 17    580  
 bp                    .                    marker                    ,  
 primer                    .                    marker                    primer                    male 1  
 primer,                    marker                    female 1 primer

,	primer	mono2- 1, mono2- 2, mono3
.	primer	PCR
male 1 primer	460 bp	band ↗ , female 1 primer
436 bp band		marker
mono 2- 1 primer		411
bp band		mono2- 2 primer
41 51		684 bp band ↗ .
51 marker		mono3 primer
PCR , ,		
51 207 bp band ↗ .		
primer set		
	, mono2- 1 primer	PCR
1		1.5

genomic DNA

Roggers Bendish (1988), Dellaporta (1983)

genomic DNA ,

DNA .

가

,

가 RAPD southern hybridization

marker OPA - 17 primer 770 bp, OPB - 03 primer

749 bp, OPB - 9 609 bp,

OPF - 16 1,100 bp, 51 OPA - 17 580 bp

marker ,

marker primer , ,

genetic marker

marker ,

,

,

4

1

“

anthocyanin ”(1982, ) anthocyanin ,  
pH, , Ascorbic acid,  
γ†, “ ”(1986, )  
Ascorbic acid, , ,  
anthocynin, tannin γ†, “ Alloxan 負荷家鬼  
”(1987, ) MeOH & ether  
mouse , γ†, “  
”(1989, ), “ , , ,  
, , ”(1990, ), “  
”(1990, )  
rats alcohol , , microsomal protein , glycogen  
, glucose-6-phosphate dehydrogenase γ†, “  
”(1992, ) γ† ,  
  
,  
, Schizandrin ,  
Kochetkov, N.K. (1961), Chen, Y.-Y. (1978) ,  
absolute configuration Chizhov, O.S. (1961), Ikeya, Y.  
(1978, 1979, 1982), Liu, C.-S. (1978) , Ben  
David Y. (1978), Ghera, E. (1978), Warshawsky, A.M. (1990) .  
Gomisin A , Taguchi, H. (1975),  
Chen, Y.-Y. (1976) , Taguchi, H. (1977),  
Ikeya, Y. (1979) , Nagai, H. (1989), Mizoguchi,

Y. (18991) . Gomisin D , Ikeya, Y.  
 (1979) , Lian-niang (1985) . Gomisin F  
 , Ikeya, Y. (1979) ,

Gomisin K1 , Ikeya, Y. (1978,  
 1979, 1980) , Suekawa, M. (1987) .

Gomisin N O , absolute configuration  
 Ikeya, Y. (1972, 1973, 1979, 1982) . Gomisin P ,  
 Taguchi, K. (1975), Liu, C.-S. (1978), Ikeya, Y. (1979)  
 , Ikeya, Y. (1980, 1981, 1982, 1983) ,  
 Hikino, H. (1984) . Gomisin Q ,  
 Ikeya, Y. (1979, 1990) ,

Liu, C.-S. (1982) .

Sesquicarene , Ohta, Y.  
 (1968), Bohlmann (1979) , Garbers, C.F.  
 (1975), Kitatani, K. (1976), Uyehara (1985), Johnson, C.R. (1987) .

Bisabolene , Bohlmann  
 (1974), Zdero, C. (1988) , Vig, O.P. (1979)

Chamigrenal Chamigrene , Ohta, Y.  
 (1968), Ito, S. (1967) , Tanaka, A. (1967),  
 Ireland, R.E. (1984), Martin, J.D. (1986) .

3-Copaene(=Ylangene) , Motl, O.  
 (1962), Irie, T. (1964), Ohta, Y. (1969) , Corey,  
 E.J. (1973), Wenkert, E. (1992) .

, Nigranoic acid Kikuchi, M  
 (1972), Schizandronic acid Takahashi, K. (1975, 1976), Schizandrol  
 Takahashi, K. (1976), Ayer, W.A. (1982)

, 2-Tridecanone                    Allen, R.R. (1965), Regnier, F.E. (1968)

, Majek, M. (1974), McDaniel, C.A. (1987)

가, , .

가 日本 中國

가 .

anthocyan , 五味

, , , tannin,

, alcohol

alloxan 負荷家鬼 , 가

phenol ,

**lignan** gomisin, schizandrin,

sesquicarene, bisabolene, chamigrene, ylangene



## 1.

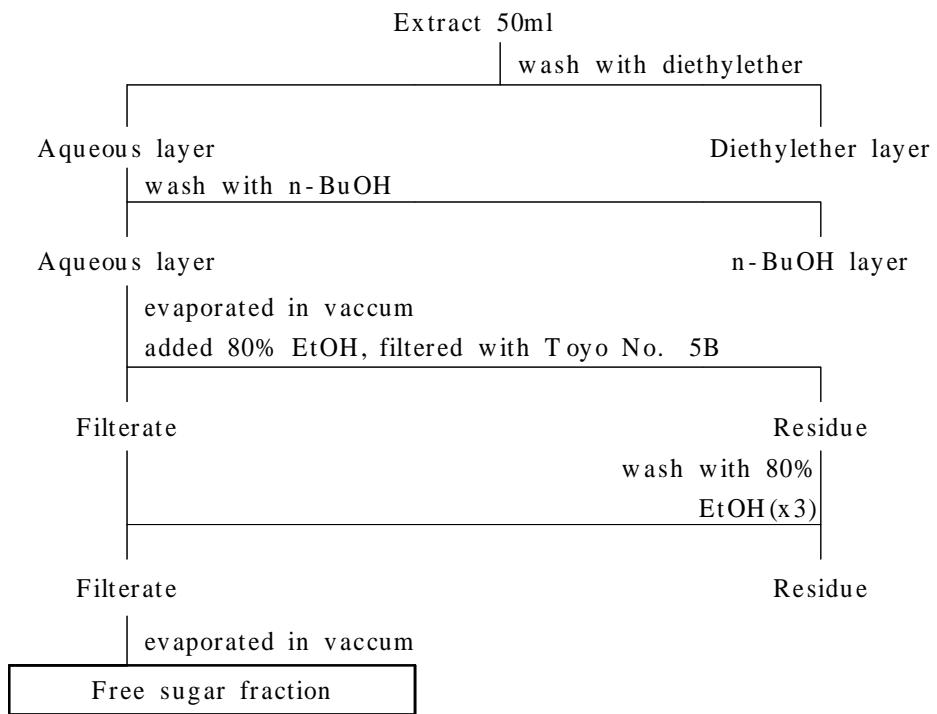
,

## 2.

가†.

- 1) , 10g .  
 2) 80% EtOH 50ml homogenizer(Nissei , Ace homogenizer  
 AM-11, Japan) , 80 2 , Whatman No. 2  
 , 45 50ml  
 가 50ml가 EtOH  
 . . 4-1.

- 3) C<sub>18</sub> cartridge , C<sub>18</sub>  
 cartridge MeOH ,  
 C<sub>18</sub> cartridge  
 가 4ml , 1ml membrane filter(Millipore,  
 0.45μm, Waters ) 3.1. ( HPLC)



. 3.1. , .

### 3.1. HPLC

Instrument	Waters Associate
Column	Carbohydrate analysis(4mm x 30cm)
Solvent	Acetonitrile / Water (80:20, v/v)
Detector	Differential Refractometer(RI)
Flow rate	1.5ml/min
Injection Volume	20μl

#### 1) HPLC

7) 10g 20ml homogenizer(Nissei , Ace homogenizer  
AM-11, Japan) , ,

3 . . ehter 2  
, HPLC 40ml  
) C<sub>18</sub> cartridge , Membrane  
filter (0.45μm)

### 3.2. HPLC

Instrument	Waters Associate
Column	RSpak KC-811(4mm x 30cm)
Column Temp.	40
Solvent	0.1% H <sub>3</sub> PO <sub>4</sub> in water
Detector	Differential Refractometer(RI)
Flow rate	1.0 ml/min
Injection Volume	20μl

2) 가 ( GC)

가) 1) HPLC 가) , 10ml

) ,  
BF<sub>3</sub>/MeOH CHCl<sub>3</sub> 2ml 가 60 25  
ammonium sulfate 4ml 가 , CHCl<sub>3</sub> GC

) Hewlett packard 5890 series , Ultra 2 (Crosslinked  
MethylSiliconGum, 25m x 0.32mm x 0.52 μm film thickness,  
Hewlett Packard Co.), FID, 250 , 270 ,  
70 1 5 210

1)

7) 10g 100ml Homogenizer (Nissei , Ace  
 homogenizer AM-11, Japan) , 80 40

) 25% TCA (Trichloroacetic acid) 7) , 1  
 , (5,000xg)

) ethylether 7) , TCA, ,  
 50m1 7) 0.2M Na-citrate  
 (pH 2.2)

2) : 7)

7) 1g 6N HCl 10ml 110 24 7)  
 ,  
 ) 0.2N sodium citrate buffer(pH2.2) 10ml  
 3.3.

### 3.3.

Instrument	LKB 4150 ALPHA
Ion exchange resin	Ultrapac 11 cation exchange resin
Column Temp.	60
Buffer flow rate	45 ml/h
Ninhydrin flow rate	35 ml/h
Buffer pressure	22 bar
Ninhydrin pressure	16 bar
Reaction bath Temp.	130
pH range	3.2 10.0
Injection volume	40 $\mu\ell$

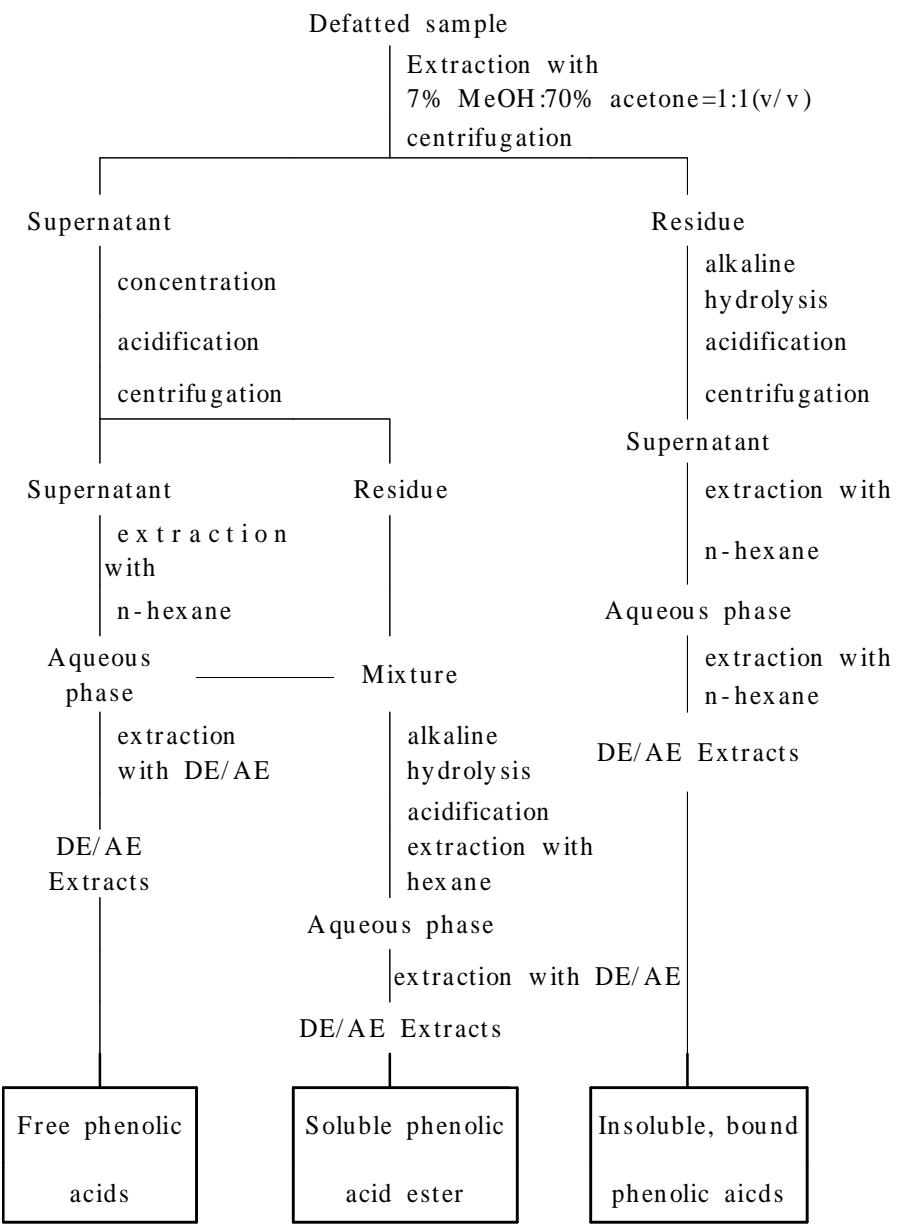
### . Phenol

- 1) Total phenol
- 7) Phenol phosphomolybdate  
 Folin - Denis

) Folin-Denis reagent : Sodium tungstate 25g, phosphomolybdate 5g,  
phosphoric acid 12.5ml 188ml 2  
, 250ml .  
)  
(1) 10g 95% ethanol 40ml ,  
homogenizer (Nissei , Ace homogenizer AM-11, Japan) 3  
Whatman No. 3 (5 ).  
(2) 100ml mass flask 80ml, 1g ,  
Folin-Denis 2ml 5ml 250ml  
100ml .  
(3) 1 , 660nm 3  
blank test 0.1% 3.0% tannic  
acid spectrophotometer (Schimadzu, spectrophotometer UV-120-02,  
Japan) , standard curve total phenol  
tannic acid .  
2) , ester  
250ml  
(1) Krygier  
(2) , Soxhlet diethylether 12  
30g 250ml 70%  
EtOH : 70% acetone = 1 : 1(v/v) 250ml , sonicator 5  
3,000 rpm 30 .  
(3) 5 , 1,250ml 100ml ,  
100ml 250ml 50ml 250ml ,  
(4)

)  
(1) 6N HCl pH 2 ,  
4,000rpm 30 .  
(2)  
250ml hexane 5 .  
(3) diethylether : ethylacetate = 1 : 1(v/v, DE/AE)  
6 .  
(4) DE/AE anhydrous sodium sulfate  
30 , MeOH  
10ml , 偈 .

) Ester  
(1) Sodium sulfate  
,  
(2) 偈 4N NaOH 250ml 偈 (偈 ) 4 .  
偈 )  
(3) 6N HCl偈 pH 2 ,  
hexane ) (3), (4)  
)  
(1) {偈} (4) 4N NaOH 250ml偈 ,偈  
4偈 .  
(2) , 6N HCl偈 pH 2 ,  
hexane DE/AE ,  
MeOH , 10ml  
偈 . 4-2.



. 3.2. , .

) ( TLC)  
(1) 10ml , Ester  
TLC .  
(2) Plate(silica gel F<sub>254</sub>, Merck , 0.25mm) (1mg/ml )  
, Ester  
spotting toluene:ethylacetate:formic acid(5:4:1, v/v)  
ferric chloride 5% potassium ferricyanide 0.5%  
1:1 ,  
HPLC GC .

) Phenol HPLC  
10ml , Ester  
1ml membrane filter(Millipore, 0.45μm, Waters ) 3.4.  
(1,000ppm) HPLC

### 3.4. HPLC

Instrument	Waters Associate			
Column	μ Bondapak C <sub>18</sub> (4mm x 30cm)			
Solvent system	Solvent A ; 100% CH <sub>3</sub> CN Solvent B ; Water : CH <sub>3</sub> CN : CH <sub>3</sub> COOH = 86 : 8 : 4(v/v)			
Elution programming	0 - 10 min	solvent A 1 %	solvent B 99 %	
	10 - 50 min	solvent A 60 %	solvent B 40 %	
	50 - 55 min	solvent A 60 %	solvent B 40 %	
	55 - 60 min	solvent A 1 %	solvent B 99 %	
Detector	UV 280 nm			
Flow rate	1.5 ml/min			

) Phenol GC  
(1)  
( 2g ) 2ml Vial γ

, 2가

(2) N,O-bis(trimethylsilyl)acetamide

(가) N,O-bis(trimethylsilyl)acetamide 0.1ml 가 Vial

70 7 가

( ) MeOH 가 2ml , GC

(3) 14% BF<sub>3</sub>-MeOH

(가) 14% BF<sub>3</sub>-MeOH 1ml 가 Vial 2

, 100ml ,

( ) NaCl 20ml ethylether 10ml

( ) Ethylether 가 ethylether 2ml

GC

(4) GC 3.5.

3.5.

GC

Instrument	Hewlett packard 5890 series
Column	Ultra 1(Crosslinked MethylSiliconGum, 25m x 0.32mm x 0.52 μm film thickness, Hewlett Packard Co.)
Detector	FID
Carrier gas	N <sub>2</sub>
Detector & Injector Temp.	300
Oven Temp.	60 (1 ), 10 , 300 (15 )

1)

가) 20g chloroform : MeOH : water = 2 : 1 : 1(v/v)

Bligh Dyer

) , 20 (1,000ml homogenizer(Nissei ,

Ace homogenizer AM-11, Japan) ,

48 , chloroform

24 , chloroform

3 .

) Chloroform oven

24 , が

2)

가) Rouser silicic acid ( SACC)

) , silicic acid(100~200 mesh, Sigma , U.S.A)

MeOH 110 oven

24 .

) silicic acid 30g chloroform slurry が

column chloroform ,

column 2 .(column size : 1.7 x 40 cm)

) chloroform column , chloroform

( ), acetone( ) methanol( ) <sup>27)</sup>

) ,

3.3.

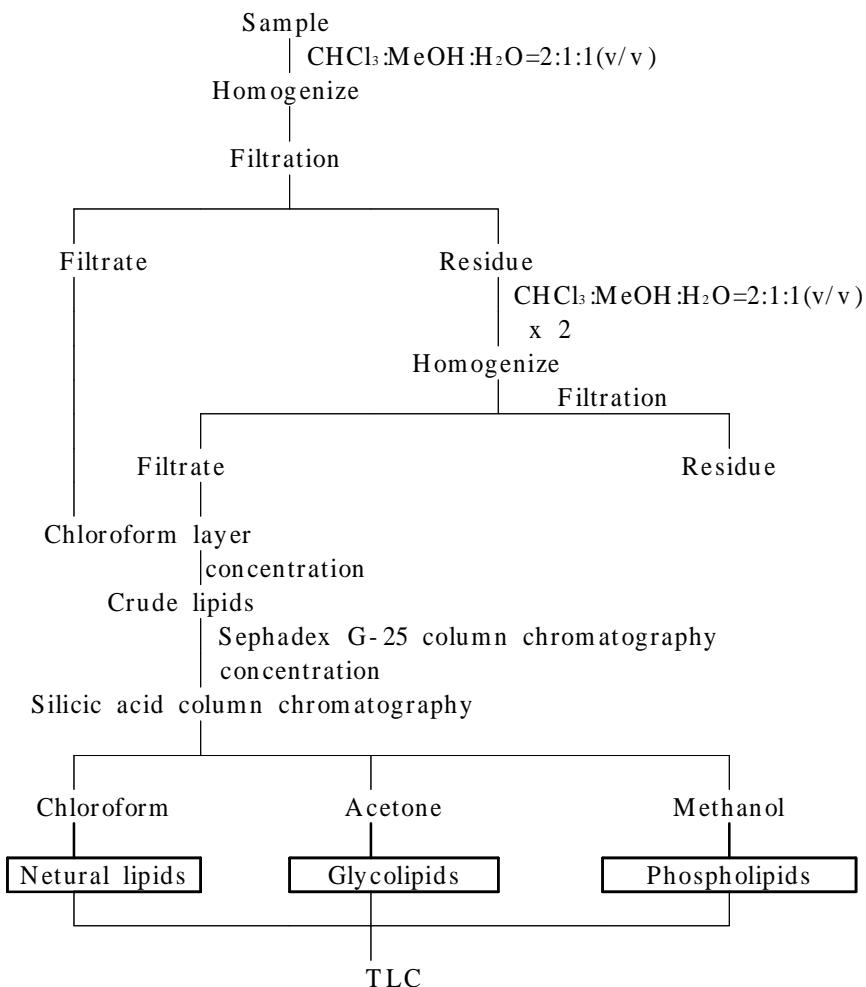
3) GC

가)

GC(Hewlett Packard 5890 series II)

GC 3.6. .

)



3.3. . ,

## 3.6.

## GC

Instrument	Hewlett packard 5890 series
Column	15% DEGS, glass, 3mm x 1m,
Detector	FID
Carrier gas	N <sub>2</sub>
Detector & Injector Temp.	250
Oven Temp.	80 (2), 5, 180 (38)

(1) ethylether 10ml 2ml  
 가 0.5N NaOH in MeOH 8ml 가  
 100 . 가 ( 5 ) , 5ml 14%  
 BF<sub>3</sub>-MeOH 가 2 .  
 (2) NaCl 20ml flask 가 ,  
 , 20ml petroleum ehter(b.p 30 60 ) 가 1  
 , ether 50ml  
 (whatman No. 2) , 가 .  
 (3) petroleum ehter 1ml (2 ) GC  
 . (500 ppm )

## 3.

## 가.

50g ethylether 50ml 3  
 가 , ethylether 50ml

## . GC/MS

1) GC/MS (Hewlett packard Co., 5970 series Mass Selective Detector)

2) : column Ultra 2(Cross-Linked 5% Phenyl Methyl Silicone, 25m  
x 0.32mm x 0.52 μm film), FID, 300 ,  
1 μl , Oven 60 1 ,  
7 100 100 1  
3 160 , 10 10  
280 10 .

#### 4.

7).

1)  
7) Hayashi , 100g 500ml 1%  
HCl in MeOH homogenizer ,  
, Toyo No. 2 G<sub>3</sub> glass filter  
, 3 .  
) 1% HCl in MeOH 100ml

2) : 10g 0.01% HCl in MeOH  
0.01% HCl in EtOH 20ml 7 Sonicator 1  
3 , . , 440nm E<sub>440</sub>/E<sub>MAX</sub>

#### . Anthocyan

1) 0.1N NaOH 0.1N HCl  
7  
2) Anthocyanin

1) Aluminum chloride : AlCl<sub>3</sub> 5g ethanol 100ml  
 2) Ammonium molybdate : ammonium molybdate 2N  
 HCl pH 2  
 3) Lead acetate : 1g lead acetate 75% ethanol 150ml

3)

Harborne Spectrophotometer  
 , 5% AlCl<sub>3</sub> ethanol 3 1

4) Anthocyanin

Fuleki Francis Philip

TAN in mgs per 100g = OD X DV X 100/SV X TEV/SW X 10/E<sup>1cm 1%</sup>  
 OD = , DV = , SV = , TEV = , SW =

$$E^{1cm\ 1\%} = \frac{\text{molar absorbance} \times 10}{\text{molecular weight}}$$

1) Amberite IRC-50

1)

(Amberite IRC-50)

5% HCl 500ml H<sup>+</sup> , 500ml  
 0.1N NaOH 500ml 500ml  
 5% HCl 500ml  
 H<sup>+</sup> , 500ml 0.1N NaOH 500ml ,  
 500ml

)
 (1) charge( , 100g  
 MeOH 100ml 40ml charge)  
 (800ml, ) 0.3% HCl in EtOH  
 (500ml, ) .  
 (2) , (1)  
 , 3 .  
 (3) Whatman No. 1  
 ( PC) , 偈

3.7.

3.7. PC

a	- -	15:82:3	
b	n - BuOH - -	4:1:5	
c	- -	5:2:3	
d	- -	30:10:3	
e	-	1:99	
f	EtOAc - -	80:15:15	
g	n - BuOH - pyridine -	6:3:1	
h	n - BuOH - pyridine -	6:3:3	
i	Benzene - -	2:2:1	
j	Benzene - MeOH -	90:16:8	

) PC

(1) Alkaline potassium permanganate : 2% Na<sub>2</sub>CO<sub>3</sub> 1% KMnO<sub>4</sub>

100 , ,

, ,

(2) AgNO<sub>3</sub> 0.1ml acetone 20ml偈 AgNO<sub>3</sub>

偈 paper

0.5M-NaOH in EtOH

(1), (5-10).

2) Aglycone

가)

(1) aglycone

4ml 50ml 1N HCl 20ml 가 30 가

(2) Aglycone

가 Amberite IRC-50

300ml , aglycone 0.3% HCl in EtOH 200ml

, 4.7. PC (Whatman No. 1)

aglycone

) Aglycone

(1) Whatman No. 3MM PC , aglycone

가 3.8.

(2) , aglycone Whatman No. 3MM

(20x20cm) banding / / (15:82:3) ,

aglycone 1% HCl in MeOH 가

, ,

## 3.8. Aglycone

PC

A	n - BuOH/ /	100:25:60	
B	n - BuOH/ /	4:1:5	
C	n - BuOH/ 2N HCl	1:1	
D	/	97:3	
E	/ /	5:2:3	
F	/ /	15:82:3	
G	/ /	30:10:3	
H	/	2:98	
I	/ /	5:1:5	
J	/	99:1	
K	EtOAc/ /	80:15:15	
L	n - BuOH/ /	7:2:5	
M	Acetone/ 10%	1:1	
N	Benzene/ /	2:2:1	
O	Benzene/ MeOH/	90:16:8	
P	phenol/	4:1	
Q	isoBuOH/ /	8:2:3	
R	n - BuOH/ Pyridine/	6:3:1	
S	Isoamylalcohol/ /	21:5:4	
T	Isopropylalcohol/ 10%	1:1	

1)

PC , aglycone spectrophotometer  
 (Hewlett packadr, HP 8453E UV-visible spectroscopy system)

2) HPLC

Whatman No. 3MM PC , aglycone

HPLC      3.9.

3.9.      aglycone      HPLC

Instrument	Waters Associate
Column	$\mu$ Bondapak C <sub>18</sub> (4mm x 30cm)
Solvent	/ / MeOH(10:73:17)
Detector	546nm
Flow rate	1.0 ml/min
Injection Volume	10 $\mu$ l

5.      : Schizandrin

71.

1) , 150g Soxhlet ether 8  
, MeOH 8

2) Ether

ether (Ether )

3) MeOH 200ml , EtOAc 200ml 3  
, MeOH 40ml , celite 16g

celite

4) Celite 30g column 3) Celite column  
charge n-hexane 120ml, CHCl<sub>3</sub> 80ml, MeOH 120ml

5) n-hexane , ether

1)

, n-hexane ether , Silica

gel(70~250 mesh, merck ) 200g      hexane      slurry ,  
 column      3.10.

3.10.

No.			No.		
A	Hexane	500ml	G	Hexane-EtOAC(50:50)	500ml
B	Hexane-EtOAC(95:5)	"	H	100% EtOAC	"
C	Hexane-EtOAC(90:10)	"	I	EtOAC-MeOH(95:5)	"
D	Hexane-EtOAC(85:15)	"	J	EtOAC-MeOH(90:10)	"
E	Hexane-EtOAC(80:20)	"	K	EtOAc-MeOH(80:20)	"
F	Hexane-EtOAC(70:30)	"	L	100% MeOH	"

2)

†)

) Silica gel(70~250 mesh, merck ) 30g      n-hexane      slurry  
 , column      4.11.

3.11.

No.			No.		
†	n-hexane	200ml		Benzene-Acetone (50:50)	200ml
	n-hexane-Benzene (1:1)	"		Benzene-Acetone (40:60)	"
	Benzene	"		Benzene-Acetone (30:70)	"
	Benzene-Acetone (90:10)	"		Benzene-Acetone (10:90)	"
	Benzene-Acetone (80:20)	"		Acetone	"
	Benzene-Acetone (70:30)	"		Acetone-MeOH (50:50)	"
	Benzene-Acetone (60:40)	"		MeOH	"

. Sephadex LH- 20

1) Sephadex LH- 20(25~100 $\mu$ , Sigma )  
2) MeOH- CHCl<sub>3</sub>(1:2, v/v)  
( 400ml ),  
charge , 80ml  
, 10 .

, Sephadex LH- 20  
schizandrin(Wako ) TLC(silica gel  
60F<sub>254</sub>, merck , 10x10cm) spotting TLC  
, benzene- acetone (9:1, v/v)  
n-hexane- acetone(7:3, v/v)

1) Sephadex LH- 20 , TLC  
2) TLC plate(20x10cm) banding n-hexane- acetone(7:3, v/v)  
, Rf  
MeOH  
3) 2) TLC 2) TLC ,  
benzene- acetone(9:1, v/v)

. HPLC

TLC membrane filter(Millipore, 0.45 $\mu$ m, Waters )  
3.12. HPLC

## 3.12.

## HPLC

Instrument	Waters Associate
Column	$\mu$ Bondapak C <sub>18</sub> (4mm x 30cm)
Solvent	acetonitrile/ anf(50:50)
Detector	UV 254nm
Flow rate	1.0 ml/min
Injection Volume	10 $\mu$ l

## 3

1.

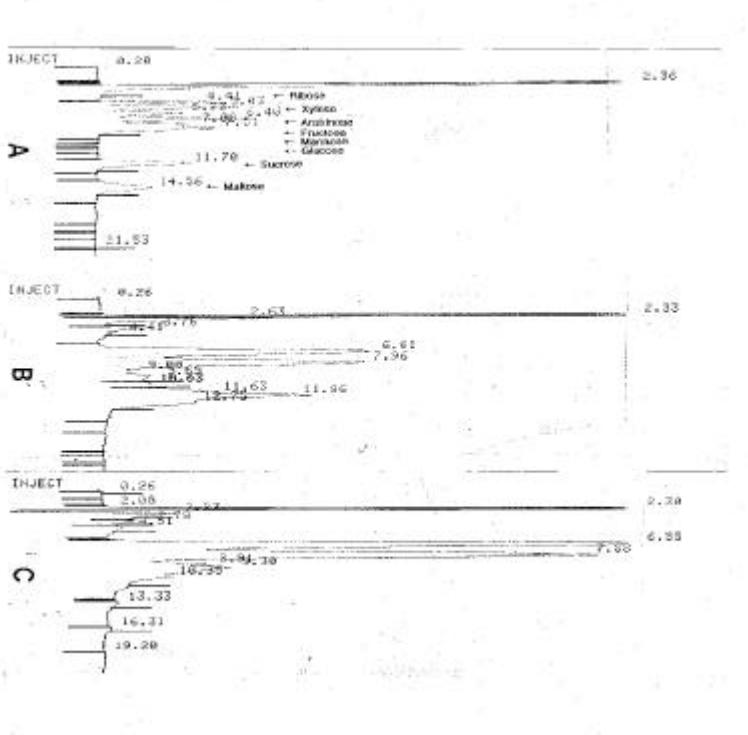
가.

1) . .  
HPLC 3.13. 3.4.

3.13.

( : % )

	Ribose	Arabinose	Fructose	Glucose	Sucrose
	0.04	-	1.83	1.05	1.13
	0.08	0.01	3.71	2.51	0.02



3.4. , HPLC

A : , B : , C :

2) 3.13.

, sucrose  
, fructose glucose  
. sucrose  
, sucrose  
가 가

1) HPLC

가) 9 (2%) HPLC 3.14.

3.5.

3.14. HPLC

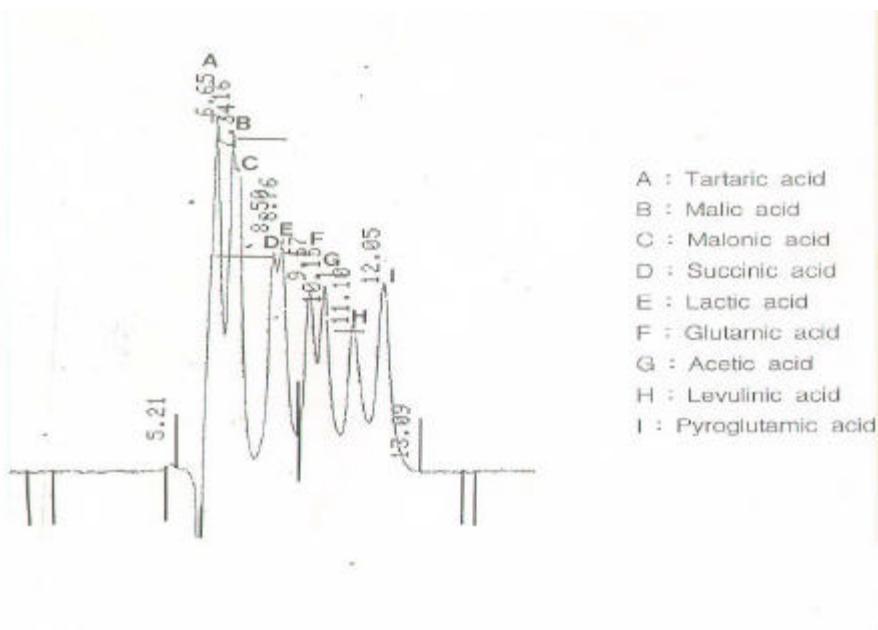
	( )		( )
Glutamic acid	9.67	Tartaric acid	6.65
Malonic acid	7.34	Succinic acid	8.50
Lactic acid	8.76	Acetic acid	10.15
Levulinic acid	11.10	Malic acid	7.16
Pyroglutamic acid	12.05	Maleic acid	6.70

) HPLC 3.15.

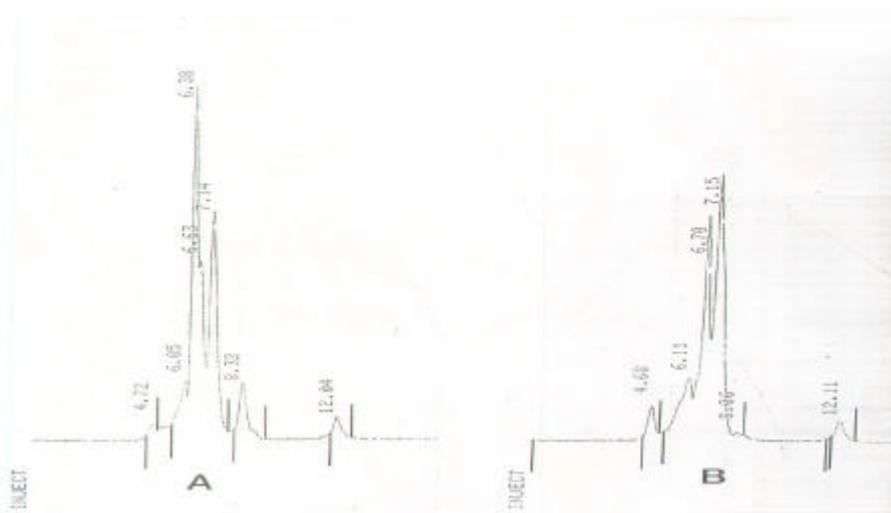
3.6.

3.15. HPLC ,

Citric acid	17.40 %	.
Tartaric acid	4.68 %	
Maleic acid		4.94 %
Malic acid		14.20 %
Pyroglutamic acid	1.02 %	1.17 %



3.5. HPLC



3.6. HPLC

A : , B :

) citric acid 가 가 , tartaric  
 acid pyroglutamic acid 가 . malic acid 가 가  
 , maleic acid pyroglutamic acid 가  
 3.15. 가  
 . , 3.6. retention  
 time( Rt) 4.72, 6.05 8.32 peak 가 Rt 8.32  
 succinic acid 3.41% . Rt 4.68,  
 6.11 8.06 peak 가 , Rt 4.72 Rt  
 4.68 , Rt 6.05 Rt 6.11  
 . 6

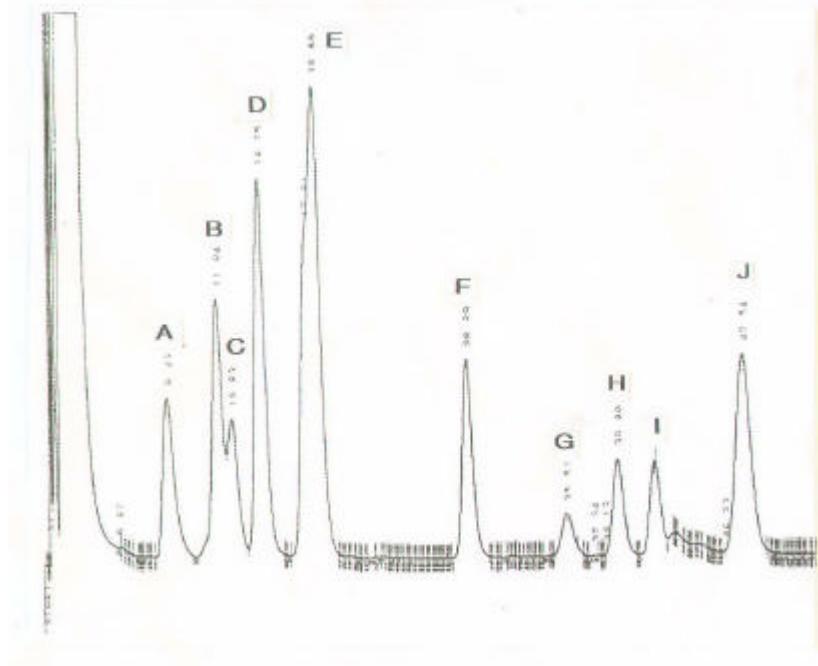
) HPLC RI  
 가  
 , External  
 GC

## 2) GC

가) 10 (1,000ppm ) GC

3.7.

) Oxalic acid methylester Retention Time( Rt) 8.43, malonic acid  
 methylester Rt 11.84, fumaric acid methylester Rt 12.83, succinic acid  
 methylester Rt 14.75, maleic acid methylester Rt 18.60, glutaric acid  
 methylester Rt 28.88, malic acid methylester Rt 35.51, tartaric acid  
 methylester 39.00, citric acid methylester Rt 41.65, pyroglutamic acid  
 methylester Rt 47.54 peak 가 , malonic acid methylester  
 fumaric acid methylester 가 .



3.7.

GC

A : oxalic acid methylester, B : malonic acid methylester, C : fumaric acid methylester, D : succinic acid methylester, E : maleic acid methylester, F : glutaric acid methylester, G : malic acid methylester, H : tartaric acid methylester, I : citric acid methylester, J : pyroglutamic acid methylester.

)

GC

3.16.

3.8.

3.16. GC

Malonic acid	0.3 ppm	2.8 ppm
Sunnic acid	247.2 ppm	21.6 ppm
Maleic acid	-	2.3 ppm
Malic acid	47,684.4 ppm	38,691.1 ppm
Pyroglutamic acid	1,241 ppm	101 ppm
Glutaric acid	74.3 ppm	-
Citric acid	11,939.1 ppm	3,330.5 ppm

) 3.16.

HPLC

, malonic acid maleic acid

가

가

1).

3.9.

,

3.17.

2) lysine

. Lysine

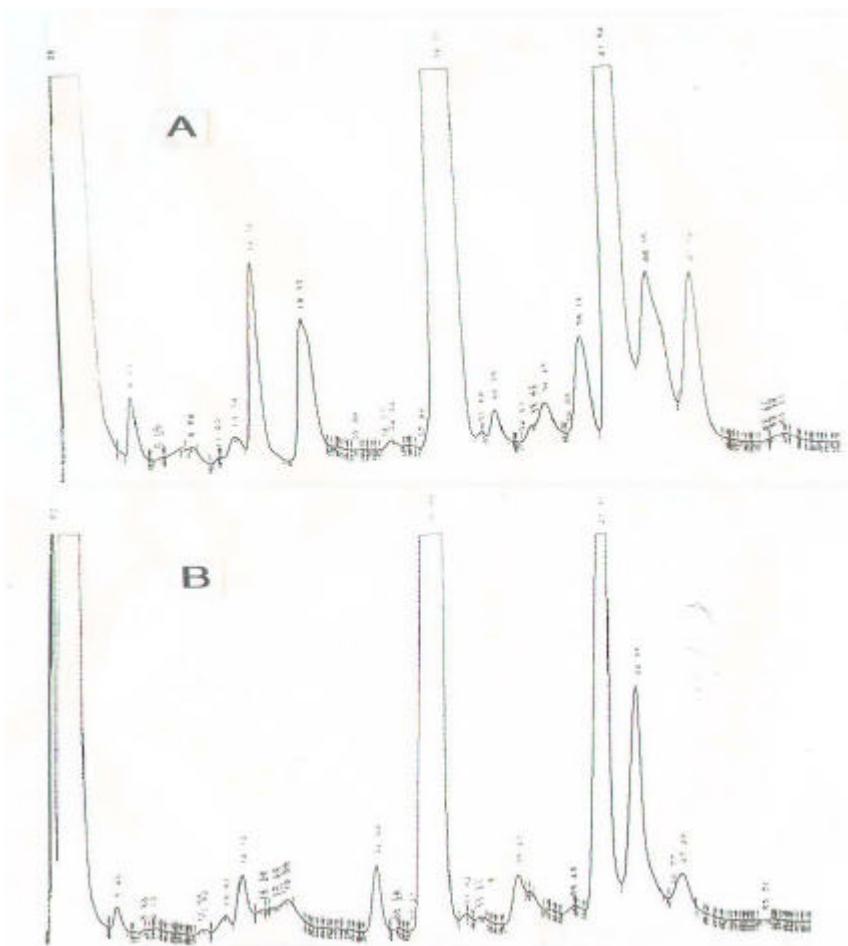
300mg %

가

ly sine 350mg%

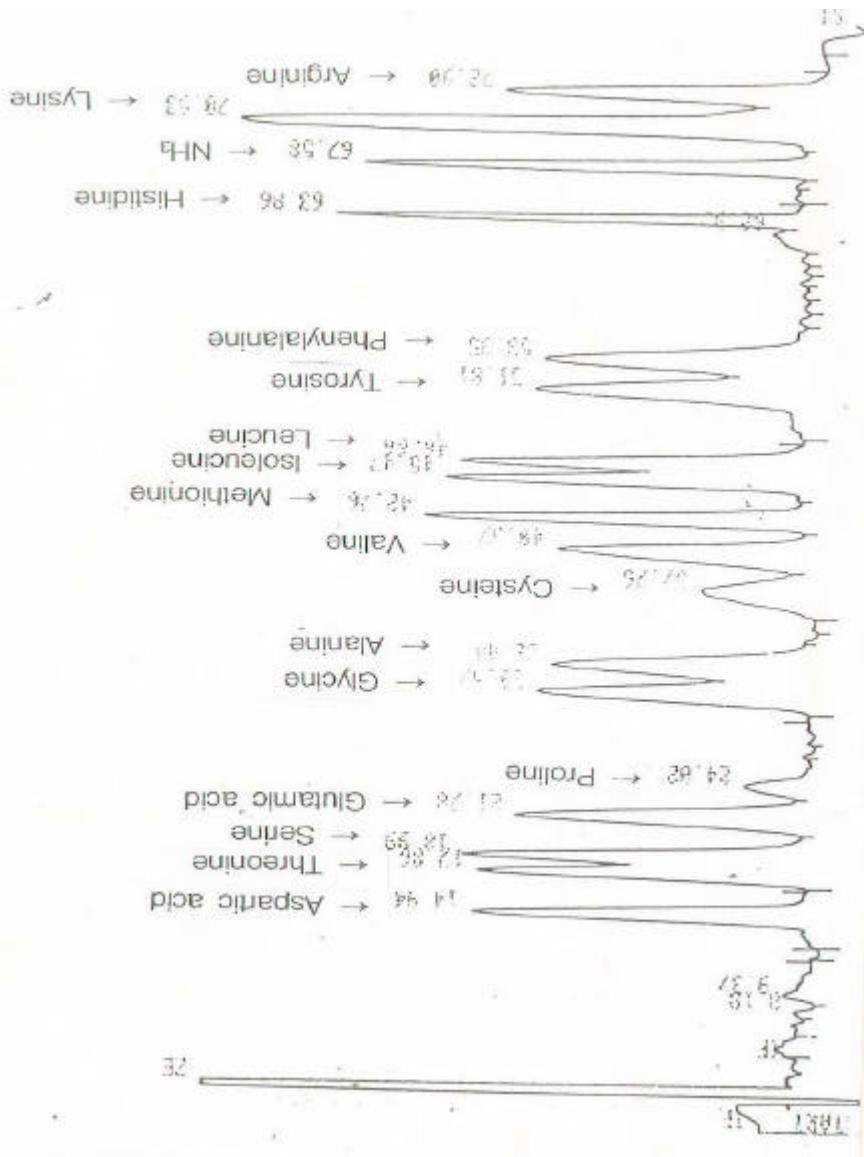
가

가 가



3.8. , GC

A : , B :



3.9.

3.17. ,

	(mg % )		(mg % )	
Aspartic acid	52.23	422.76	11.80	366.97
Threonine	154.73	511.36	31.41	427.55
Serine	0.00	600.01	0.00	491.94
Glutamic acid	97.37	721.95	11.40	677.03
Proline	23.42	427.13	5.42	239.04
Glycine	14.70	502.92	4.54	379.14
Alanine	50.19	348.16	7.14	273.84
Cysteine	0.00	211.75	12.79	15.93
Valine	34.48	638.41	16.10	426.77
Methionine	21.58	148.35	0.00	35.93
Isoleucine	24.26	469.02	0.00	358.68
Leucine	31.10	653.56	0.00	447.41
Tyrosine	12.07	316.80	13.69	139.95
Phenylalanine	12.75	390.08	0.00	214.07
Histidine	120.30	518.55	40.40	371.96
Lysine	22.37	288.74	47.72	300.17
Arginine	50.37	408.26	0.00	430.43
Total	721.94	7,577.79	202.42	5,596.84

1)

Folin-Denis

, 1.275%,

tannic acid

1.560%

2)

TLC

가) 3.2.

Ester TLC 3.18.  
3.10.

3.18. TLC

		CH	EP	CAT	TAN	PHC	CAF	GEFE	CI
	Ester	O	X	X	X	O	O	O	X
		O	X	X	X	X	X	X	X
		O	X	X	X	O	O	O	X
	Ester	O	X	X	X	O	X	O	X
		O	X	X	X	X	X	X	X
		O	X	X	X	O	O	O	X

\* CH:chlorogenic acid, EP:epicatechin, CAT:catechin, TAN:tannic acid,  
PHC:phloroglucinol & coumaric acid, CAF:caffeic acid, GFFE:gentisic acid  
& Ferulic acid, CI:cinnamic acid, O: 가 , X: 가 .

) 3.18. 3.10. TLC

GC HPLC

3) GC HPLC

가) HPLC 3.11. 3.19.

GC 3.12. 3.20.

, GC ,

가 HPLC , , HPLC

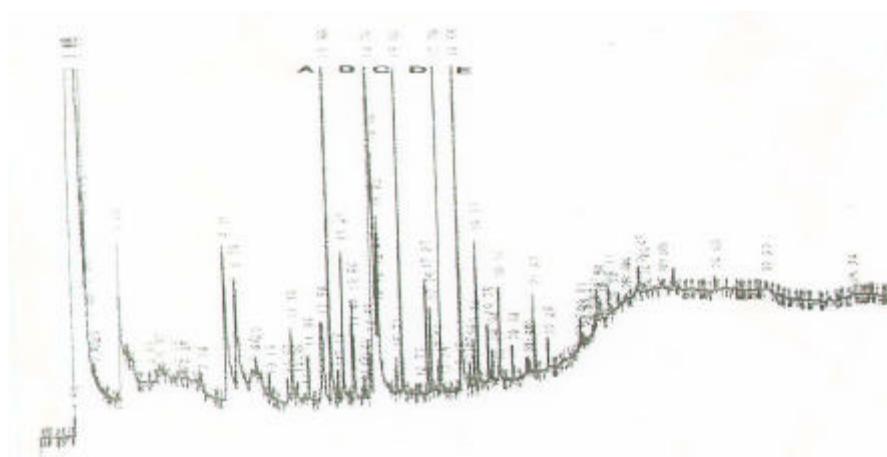
(gradient method) ,

Rt 가 GC



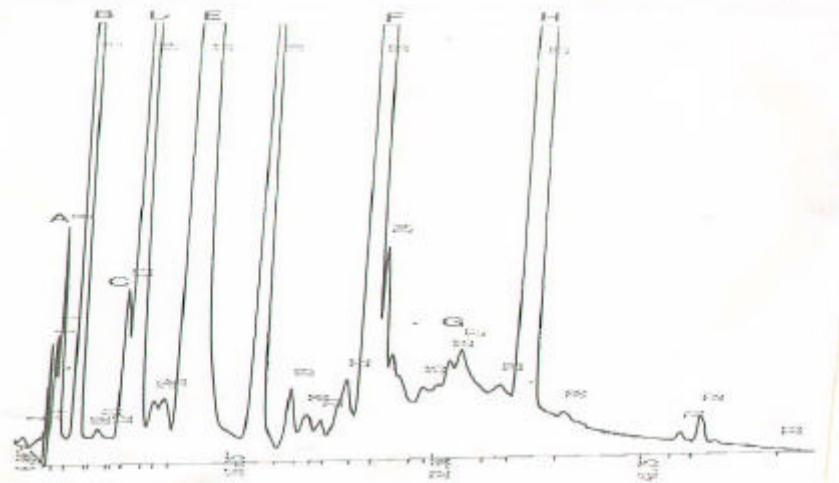
3.10. , TLC

A : , B : , C :  
 , M : , D : ,  
 E : , F :



4.11. HPLC

A : phloroglucinol, B : coumalicacid and gentisic acid, C : chlorogenic acid, D : catechin, E : epicatechin, F : ferulic acid, G : tannic acid, H : cinnamic acid.



3.12. GC (BF<sub>3</sub>/MeOH )

A : cinnamic acid, B : gentisic acid, C : coumalic acid, D : chlorogenic acid,  
E : ferulic acid.

3.19. HPLC

	Rt		Rt
Phloroglucinol	3.215	Catechin	8.958
Coumalic acid	4.588	Epicatechin	13.175
Gentisic acid	4.588	Ferulic acid	27.001
Caffeic acid	6.408	Tannic acid	34.528
Chlorogenic acid	8.161	Cinnamic acid	39.555

3.20. GC

	Cinnamic acid	Caffeic acid	Gentisic acid	Coumalic acid	Chlorogenic acid	Ferulic acid
Rt	12.90	14.42	14.76	15.98	17.70	18.55

\* Caffeic acid N,O-bis(trimethyl silyl)acetamide

, 14% BF<sub>3</sub>-MeOH

)

GC HPLC

3.21.

) 3.21.

cinnamic acid

, caffeic acid

가

. Gentisic acid

, coumaric acid

가

chlorogenic acid ferulic acid

) 3.21.

3.18.

, epicatechin, catechin,

phloroglucinol

tannic acid

가

,

chlorogenic acid, gentisic acid, coumaric acid, ferulic acid

cinnamic acid

TLC

3.21.

( : ppm)

	CI	CAF	GE	CO	CH	FE
Ester	22.71	46.73	28.14	trace	234.08	25.20
	trace	-	17.20	trace	305.15	15.172
	33.00	trace	24.37	91.58	637.67	147.34
Ester	1.84	-	96.76	58.44	167.71	9.36
	trace	-	22.36	trace	357.97	4.06
	8.44	trace	18.53	46.70	302.04	10.77

\* CI : cinnamic acid, CAF : caffeic acid, GE : gentisic acid, CO : coumaric acid, CH : chlorogenic acid, FE : ferulic acid

1)

7) Bligh Dyer silicic acid

3.22.

3.22.

(unit : mg/g. dry weight)

	110.5	32.0	3.0	145.5
	120.7	35.8	4.0	160.5

)

10%

7)

GC

2)

GC

7)

17 (500 ppm )

GC

3.23.

3.13.

3.23.

GC

	Rt (min)		Rt (min)
Caproic acid(C6:0)	2.08	Stearic acid(C18:0)	24.15
Caprylic acid(C8:0)	5.12	Linoleic acid(C18:2)	26.27
Capric acid(C10:0)	9.20	Linolenic acid(C18:3)	28.66
Lauric acid(C12:0)	13.31	Arachidic acid(C20:0)	29.33
Myristic acid(C14:0)	17.15	Eicosenoic acid(C20:1)	30.31
Myristoleic acid(C14:1)	18.08	Arachidonic acid(C20:4)	36.76
Palmitic acid(C16:0)	20.70	Behenic acid(C22:0)	38.80
Palmitoleic acid(C16:1)	21.36	Erucic acid(C22:1)	40.34
Oleic acid(C18:1)	24.83		

) , , ,

GC

3.24.

3.14.

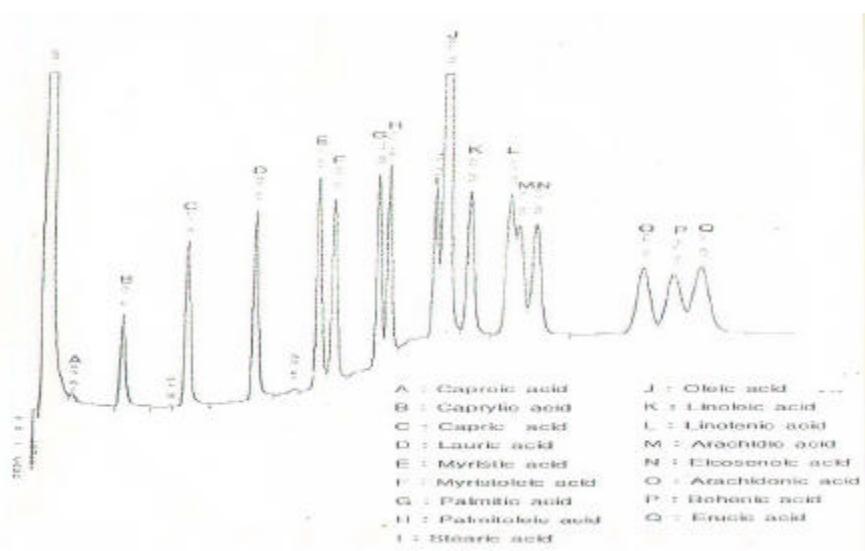
3.24. ,

( : ppm )

	A	B	C	D	E	F
Caprylic acid	50	14	3	169	25	2
Capric acid	1,795	14	9	2,178	29	5
Lauric acid	112	4	0.5	358	3	9
Myristic acid	1	0.5	0.5	8	10	16
Myristoleic acid	7	trace	14	3	6	trace
Palmitic acid	1,942	1,990	42	9,888	984	trace
Palmitoleic acid	trace	5	trace	trace	7	trace
Oleic acid	731	34	9	2,687	139	5
Linoleic acid	12,732	8,479	554	29,364	13,368	2,852
Linolenic acid	49	106	57	322	248	16
Eicosenoic acid	60	6	trace	131	3	2
Total	17,479	1,0652	689	45,108	14,822	2,907

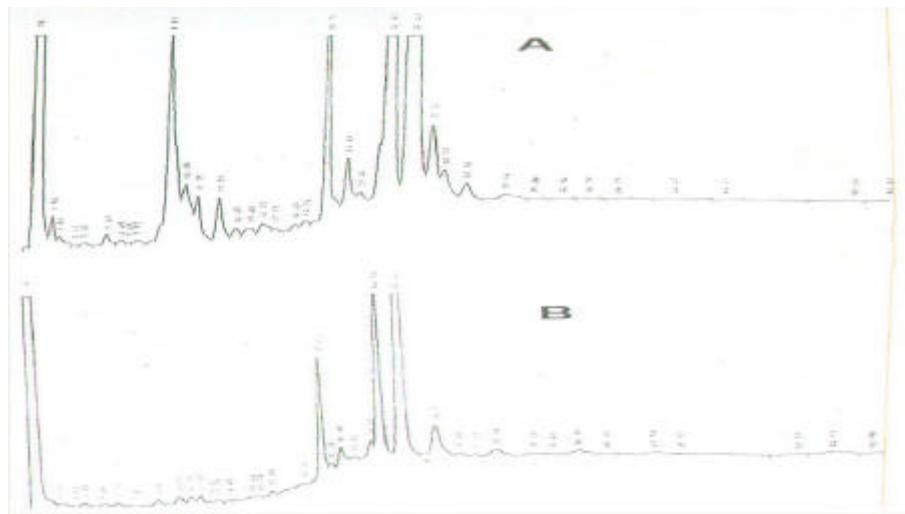
A: , B: , C:

D: , E: , F:



3.13.

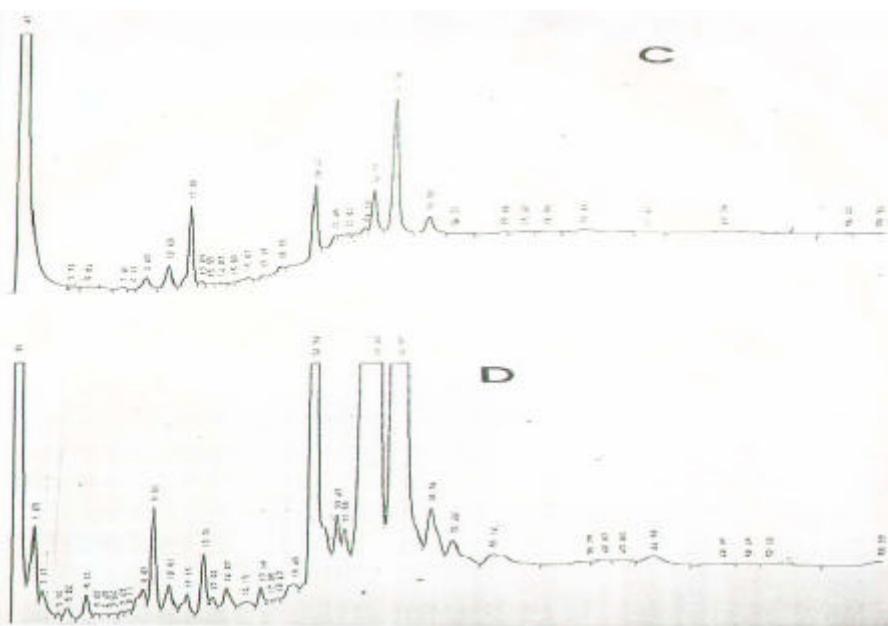
GC



3.14. ,

GC

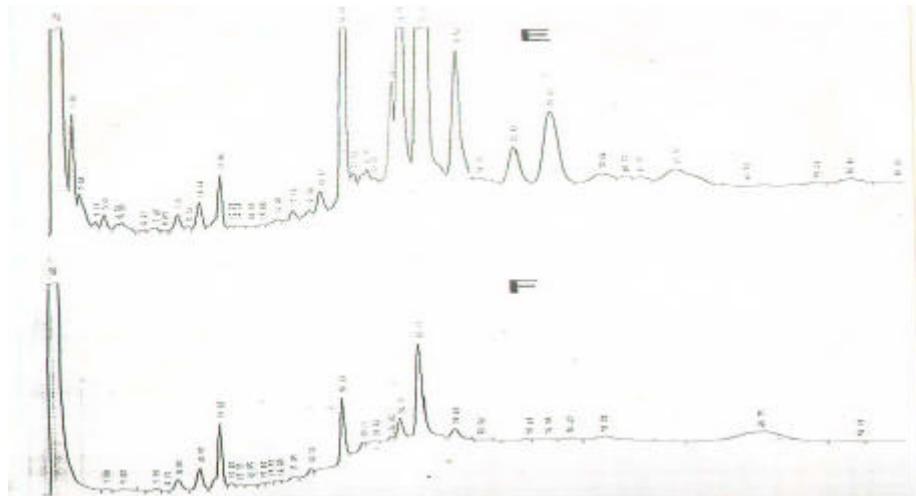
A : , B :



3.14. ,

GC

C : , D :



3.14. ,

GC

E : , F : .

) GC

oleic acid linoleic acid, linolenic acid

가 2

2.

가.

ethylether , , GC/MS

3.15. , GC peak MS

3.25. 3.16. 3.26. 3.17.

. terpene monoterpane sesquiterpene

90%

가

, , 3.15.

, ,

(R.T. 20 )

. 3.25. 3.26. -ylangene, ,

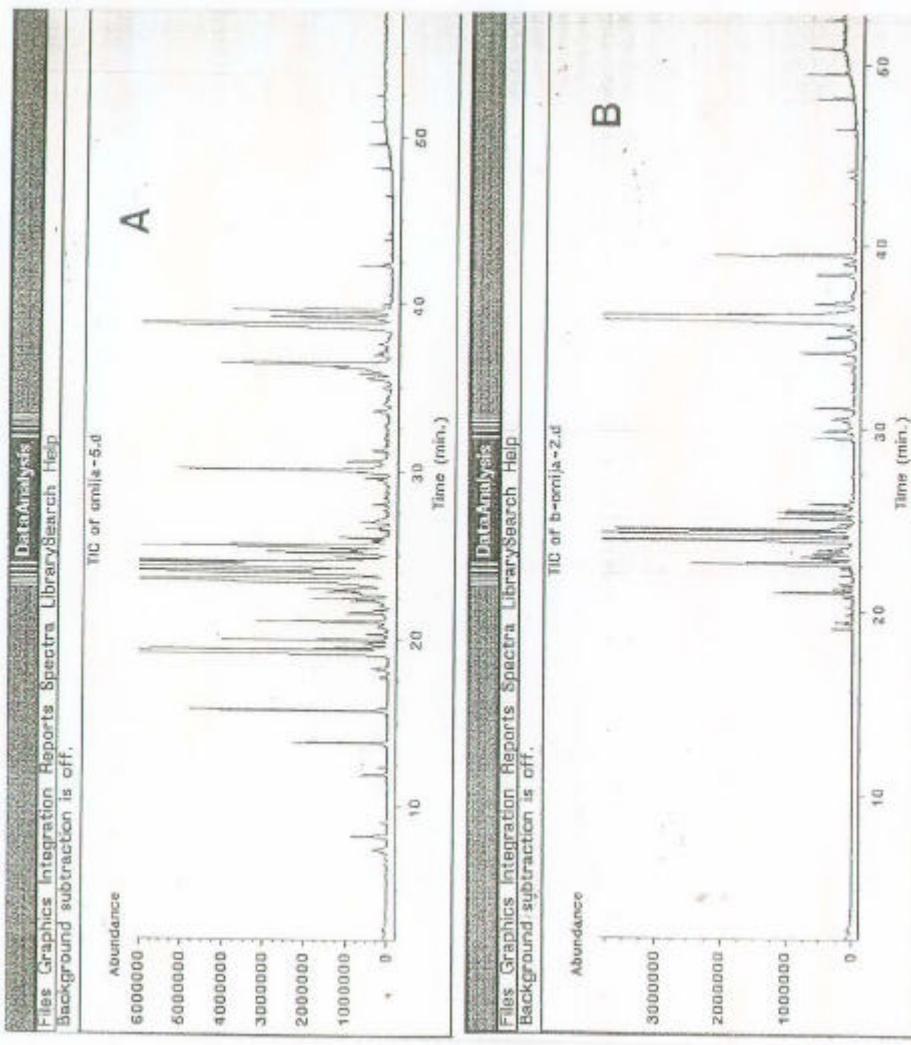
-elemene, -himachalene, -selinene widdrene

, caryophyllene, calarene, cubebene, acoradiene

-himachalene

가

가



3.15.

GC

A : , B :

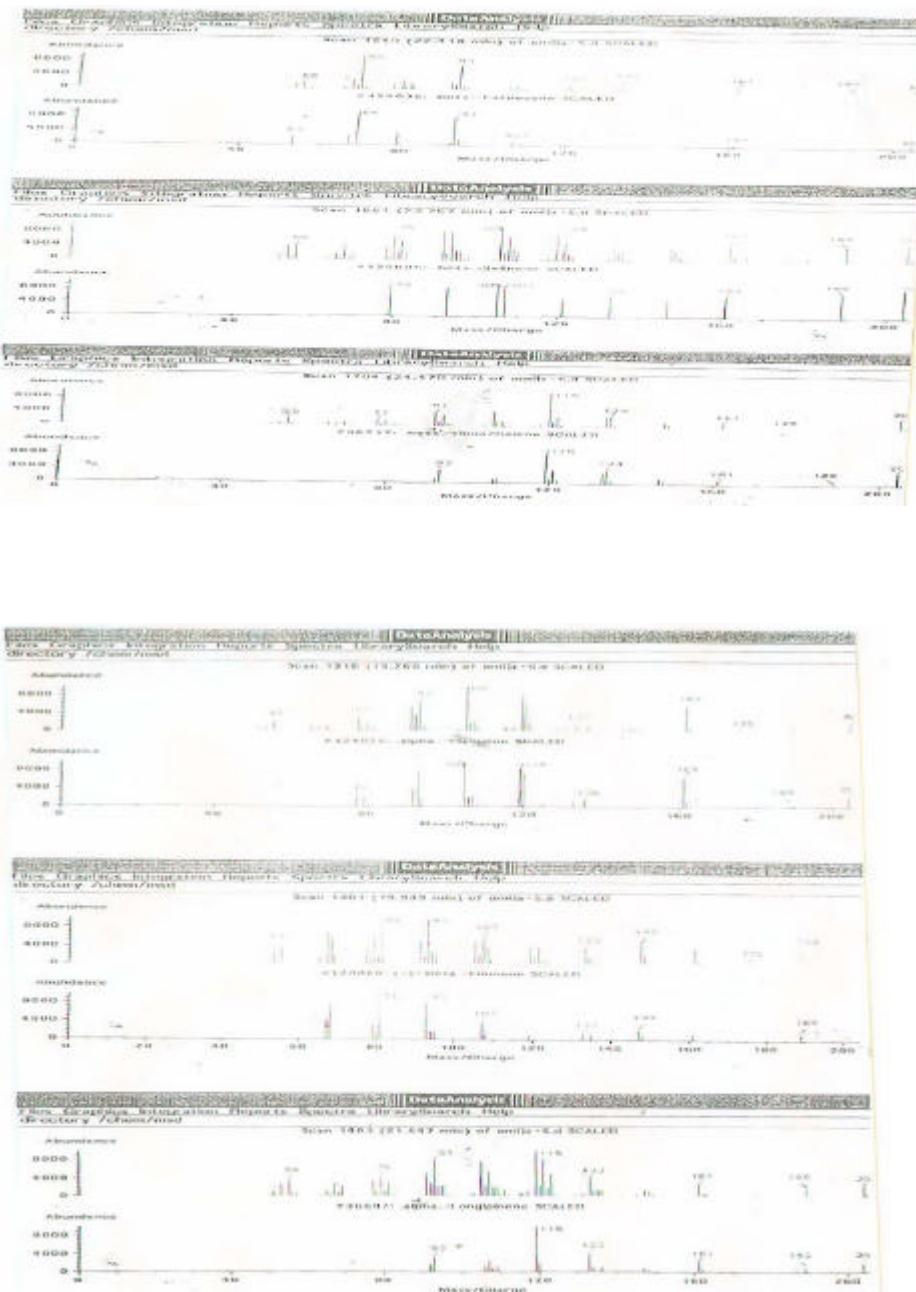
3.25.

GC/MS

Rt	Compounds	% Sg
7.437	Benzene, 1-methyl-4-(1-methylethyl)-	0.42
8.211	-Terpinene	0.69
11.857	3-cyclohexen-1-ol, 4-methyl-1-(1-methylethyl)-	0.34
12.315	1--Terpineol	0.11
13.758	Methyl Thymylether	1.07
17.749	-Terpinolene	0.12
18.263	5H-Inden-5-one ,octahydro-,trans-	0.30
19.296	-Ylangene	13.19
19.473	-Guaiene	0.78
19.707	1H-Pyrrole, 1-butyl-	0.40
19.944	(-)-Elemene	1.81
20.305	1,2-Naphthalenediol, 2-ethyl-1,2,3,4-tetrahydro-1-methyl-.ci	0.26
20.738	Tricyclo[2.2.1.0 <sup>2,6</sup> ]heptane,1,7-dimethyl-7-(4-methyl-3-pente	0.11
21.052	-Elemene	1.88
21.554	-Longipinene	0.58
21.740	-Terpinene	0.13
22.045	Longipinene	0.14
22.274	(+)-Aromadendrene	0.55
22.415	-Farnesene	0.93
22.713	-Cadinene	0.54
22.828	Acoradiene	0.86
22.959	(-)-Acoradiene	0.19
23.478	-Selinene	7.23
23.768	-Selinene	0.34
24.176	Naphthalene,1,2,3,4,4a,5,5,8a-octahydro-7-methyl-4-methylene	0.82

## 3.25.

Rt	Compounds	% Sg
24.469	- Himachalene	10.05
24.590	Widdrene	5.53
25.173	- Cadinene	1.52
25.286	6,10,11,11-T etramethyl-Tricyclo[6,3,0,1E2,3] Undec- 1(7)ene	0.57
25.761	- Muurolene	0.16
25.835	- Humulene	0.19
25.978	Quinoline, 2,6-dimethyl-	0.52
26.444	2- (5-Methyl- 3-Isoazoyl)-4-Acetyl-5 -Methylpyrrole	0.12
26.638	Nerolidol	0.28
27.943	2- (1',1'- Dideutero- Allyl)Aniline	0.23
29.910	Bicyclo[4,4,0]Dec- 1-En,2- Isopropyl- 5-Methyl- 9-Methylene-	0.41
30.110	Cycloheptene,5- ethylidene- 1-methyl-	4.22
30.587	T -Muurolol	0.75
33.591	(+)-Oxo- - Ylangene	0.40
35.010	Azocene, 2-methoxy- 8-methyl-	0.15
35.552	2,3,4- Trimethylbenzaldehyde	0.58
35.844	2-Vinyladamantane	0.63
36.112	(E)- And(z)- 7-Methyl- 5,7-Octadien- 1-Ol	1.27
36.382	Cycloisolongifolene	4.54
36.877	2,2-Diisobutenyl-3- cyclohepten- 1-one	0.32
37.158	2-Naphthalenecarboxylicacid,8- ethenyl- 3,4,4a,5,6,7,8,8a-oct	0.22
37.584	4(1H)- Quinolinone, 3-methoxy- 1-methyl-	0.29
38.709	Benzenemethanol,ar,ar, - trimethyl-	8.15
39.149	- 4-Carene	2.05
39.562	Phenol, - (2-methylallyl)-	2.67
42.262	4- (2'- ethyl- 5'- phenyl- pyrro- 3'- yl)pyridine	0.25
46.429	Tetracosamethylcyclododecasiloxane	0.05
50.854	1,1,3,3,5,5,7,7,9,9,11,11,13,13,15,15-Hexadecamethyl- octasilo	0.10



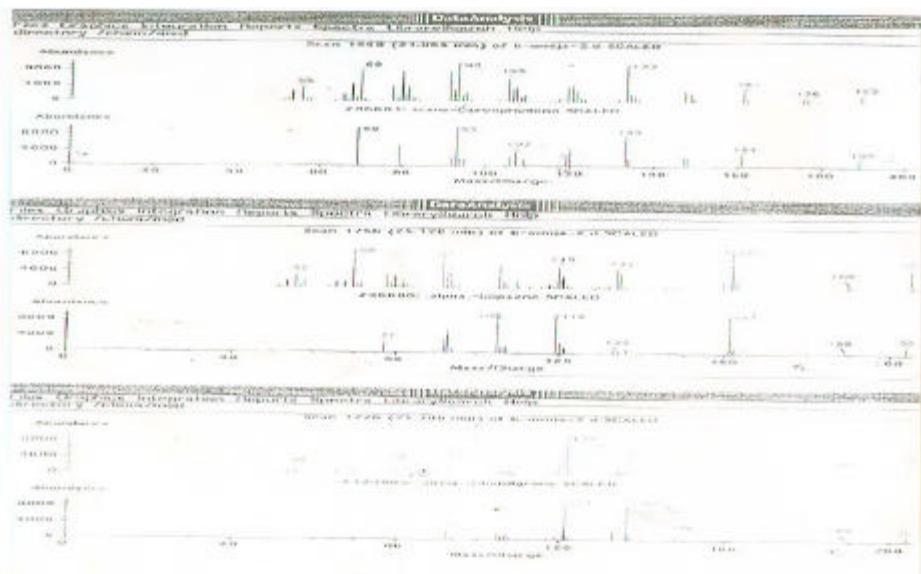
316.

MS

3.26.

G/MS

Rt	Compounds	% Sg
19.442	Selina-3,7(11)- diene	0.36
21.073	trans - Caryophyllene	1.65
21.255	1H-3a,7-Mrthan oazulene, 2,3,4,7,8,8a-hexahydro-3,6,8,8-tetram	0.51
21.542	- Longipinene	0.31
22.028	Tricyclo[3.3.0.0E4,6]octan-3-one,6-methyl-	0.30
22.661	Calarene	3.58
22.767	- Bisabolene	0.49
23.031	- Cubebene	1.49
23.247	- Chamigrene	0.96
23.385	Longipinene	1.56
23.955	(-)- Acoradiene	11.52
24.337	- Himachalene	10.19
24.519	1,2,2-trimethyl-1-( -tolyl)- cyclopentane	5.74
25.129	- Copaene	1.74
25.390	- Chamigrene	1.67
25.911	N-(-tolyl)-acrylic acid amide	1.14
29.516	- Selinene	0.96
29.893	(+)-Isobicyclogermacrene	1.05
31.143	Caryophylla-2(12),6-dien-5-one	1.09
33.588	Cyclopentene,1,3-dimethyl-2-(1-methylethyl)-	0.52
35.024	Aristolone	1.08
36.152	2-Naphthalenecarboxylic acid, 8-ethenyl- 3,4,4a,5,6,7,8,8a-oct	32.88
36.267	Benzene,1,4-dimethyl-2,5-bis(1-methylethyl)-	3.68
36.854	Tricyclo[3.2.1,0E2,8]Octan-7-one, 6-methyl-6-(2-methyl-2-prop	1.46
38.457	1,2,3,3a,5,6,6a,7-Octahydro-1,3a,6-trimethyl- 4H-cyclopent[d]i	1.40
39.496	2H-1-Benzopyran, 3,4-dihydro-2-methyl-	4.35
46.430	Benzeneacetic acid, ,3,4-tris[(trimethylsilyl)oxy]-t	0.20
50.857	Tetracosamethylcyclododecasiloxane	0.40
52.420	1,1,3,3,5,5,7,7,9,9,11,11,13,13,15,15-hexadecamethyl-octasilo	0.28



3.17.

MS

3.

### ▷. Anthocyan

1)

▷), , ▷ anthocyan

3.27.

3.27. ,

▷

		0.1N NaOH	2N HCl	AlCl <sub>3</sub>
	Ammonium Molybdate	Vanillin	Lead acetate	FeCl <sub>3</sub> - K <sub>3</sub> Fe(CN) <sub>6</sub>

) , 0.1N NaOH 가 ,  
, 2N 가

, , , ,  
,

) Vanillin ,

4-27.

) anthocyanin 3.18.

B OH 가 cyanidin, delphinidin, petunidin  
aluminium chloride , ammonium molybdate , lead acetate 가  
, OH 가 pelagonidin,  
peonidin, malvidin 가

anthocyanin . 3.27.

pelagonidin, peonidin, malvidin ,  
cyanidin, delphinidin, petunidin .

2)

가) , 0.01% HCl in MeOH 0.01% HCl in EtOH  
5% 3 가  
3.28.

3.28. ,

	(nm)		가 (nm)
	0.01% HCl in MeOH	0.01% HCl in EtOH	
	533	541	0
	540	554	15

) 3.28.

533, 541nm

가

peonidin

540, 554nm

가

15nm

540, 554nm malvidin

가 ,

cyanidin,

petunidin, delphinidin

가

delphinidin, petunidin

가

,  $E_{440}/E_{MAX}$

25.4,

28.1 3-glycoside

peonidin-3-

glycoside

malvidin-3-glycoside,

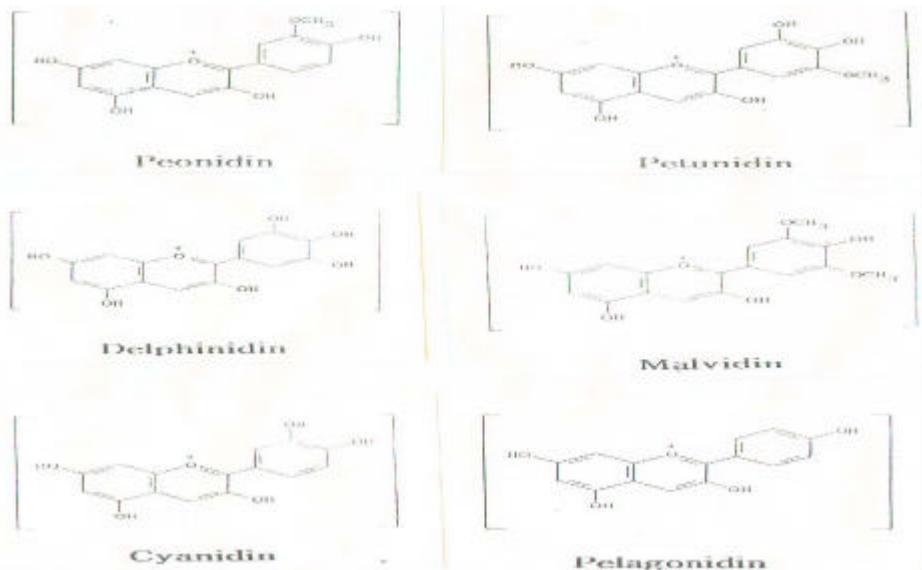
petunidin-3-glycoside, delphinidin-3-glycoside

3가

,

3가

가



### 3.18. Anthocyanin

3)

Fuleki Francis Philip

, peonidin-3-glucoside

86.54m g %

138.75mg%

2

1) Amberite IRC- 50

가)

(Amberite IRC-50)

)

## charge

0.3% HCl in EtOH

가

가

0.3% HCl in EtOH

),

, 가 , , , ,

2)

가)

0.3% HCl in EtOH

, n - BuOH/

pyridine/ (6:3:1, v/v)

가 가

) n-BuOH/pyridine/ (6:3:1, v/v)

Whatman No. 1

0.3% HCl in EtOH

## spotting

,

가

, 0.3% HCl in EtOH

가

1)

7) 67 ( 3.18.) aglycone(anthocyanin)

anthocyanin

anthocyanin

) Anthocyanin

가

가

, Amberite IRC-50

anthocyanin

HPLC

,

glucose

,

glucose

, 7), , , ,

2) PC

7) , , , , 7), , , ,

anthocyanin Whatman No. 1 spotting 3.8.

n-BuOH/ / (4:1:5, v/v, , BAW), / / (82:15:3, v/v,  
WHA) 4-29.

) , 2 , 2 anthocyanin

, , ,

anthocyanin . , anthocyanin

anthocyanin ,

anthocyanin delphinidin, malvidin, petunidin

, anthocyanin peonidin, cyanidin

3.29. anthocyanin PC .

							가
BAW	Rf 0.140	Rf 0.140	Rf 0.133	Rf 0.273	Rf 0.193	Rf 0.140	-
	Rf 0.233	Rf 0.273			Rf 0.433	Rf 0.273	
WHA	Rf 0.066	Rf 0.120	Rf 0.080	Rf 0.066	Rf 0.133	Rf 0.147	Rf 0.167
	Rf 0.266	Rf 0.273	Rf 0.253	Rf 0.233	Rf 0.353	Rf 0.273	Rf 0.320
			Rf 0.413	Rf 0.366			Rf 0.507

) anthocyanin BAW 2

, WHA

, 3.29. WHA 가

WHA .

) Anthocyanin

3.29. , 2 anthocyanin

Whatman No. 3MM (20 X 20cm)

anthocyanin banding WHA

, Rf 0.080 band, Rf 0.266 band 가

, Rf 0.060 , Rf 0.130 band 가

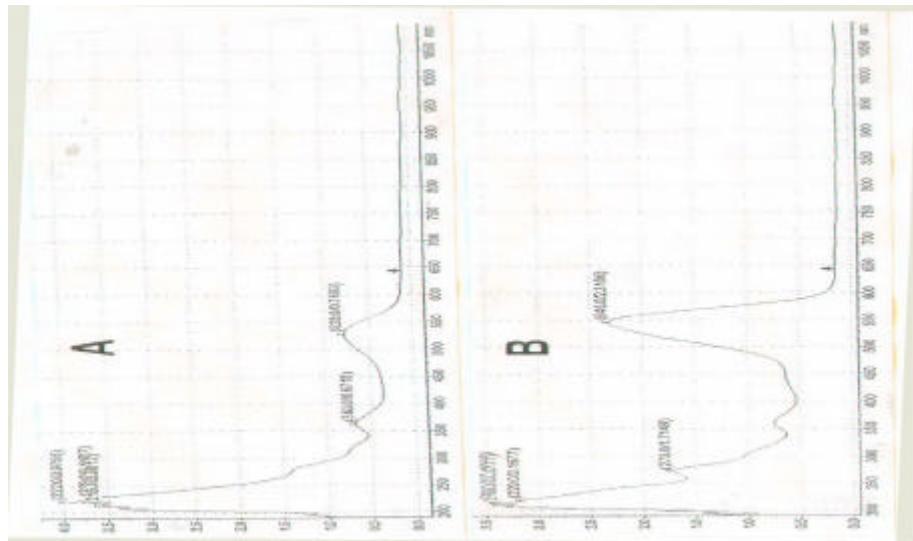
. Rf 1% HCl in MeOH .

1)

가) Rf 0.080( - 0.080), Rf 0.266( - 0.266),

Rf 0.060( - 0.060), Rf 0.130( - 0.130)  
, - 0.080 535nm - 0.266 528nm ( 3.19.) - 0.060

546nm ( 3.19.) - 0.130 538nm .



3.19. - 0.266(A) - 0.060(B)

) - 0.080 cyanidin peonidin  
 , - 0.266 peonidin , - 0.060 delphinidin , - 0.130 malvidin

2) PC  
 2 glucose  
 peonidin - 3 - monoglucoside peonidin - 3,5 - diglucoside ,  
 PC 2 glucose  
 delphinidin - 3 - monoglucoside malvidin - 3 - monoglucoside

### 3) HPLC

†) 4 4-9. HPLC , - 0.080  
 Rt 4.032, Rt 13.042 , - 0.266 Rt 4.432, Rt 12.037 , - 0.060 Rt  
 3.202, Rt 21.338 , - 0.130 Rt 4.673, Rt 24.093 peak † ,  
 . , - 0.080 Rt

13.042                  peak      Rt 13.095                  ,      - 0.266      Rt 12.037  
                         peak      Rt 12.282                  ,      - 0.060      Rt 21.338                  peak  
                         Rt 20.995                  ,      - 0.130      Rt 4.673                  peak      Rt 4.410  
                         가                  ,                  가

#### 4. : Schiznadrin

가.

	ether	MeOH	
)	ether	(ether )	MeOH
	, ethylacetate		celite
large	n-hexane, chloroform, methanol		.
	, ether , n-hexane , chloroform ,		,
	schizandrin	benzene/ acetone (9:1, v/v)	
/acetone(7:3, v/v)		TLC	
	, ether	n - hexane	
	chloroform	†,	
	†		ether

1)

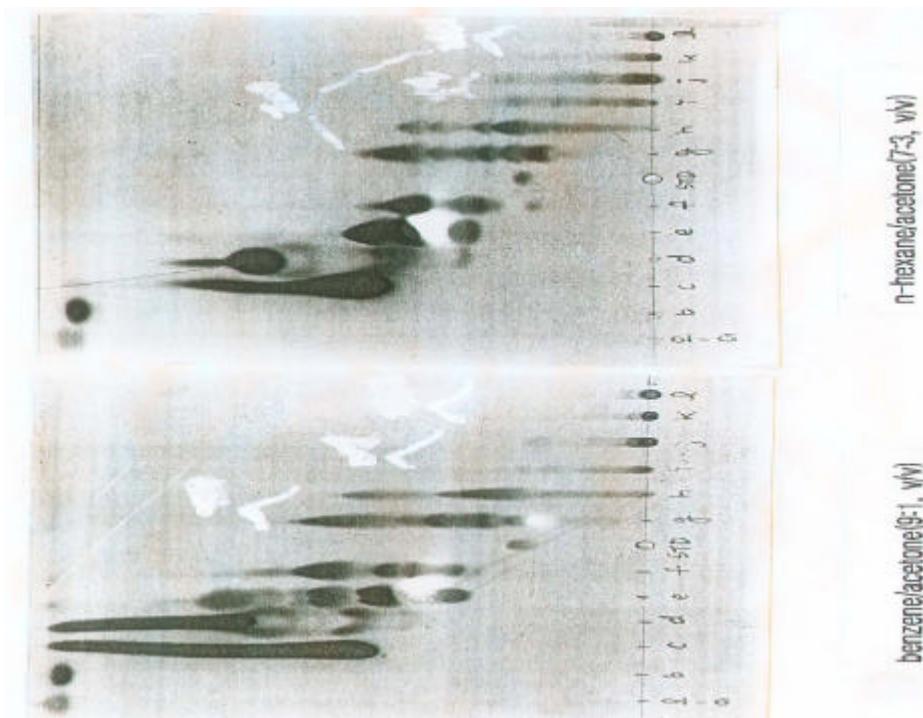
가) Silica gel	n-hexane	charge
3.10.	.	12
schizandrin	TLC	(
3.21.), G, H		가)
, I, J	.	G, H, I, J

) Silica gel n-hexane charge  
 3.11. 14  
 schizandrin TLC ( 3.22.,  
 3.23.), , γ†  
 , Sephadex

LH-20

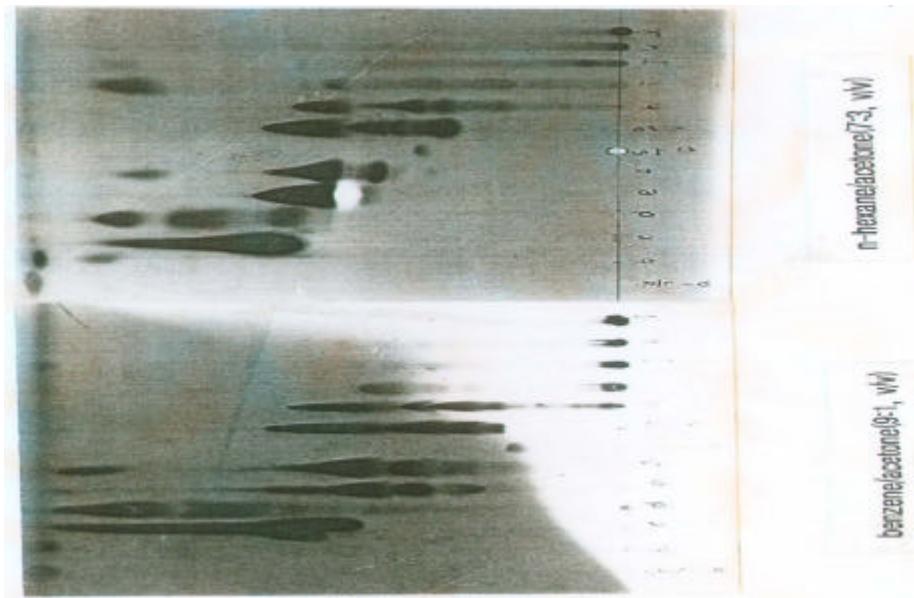
2) Sephadex LH-20

γ†)  
Sephadex LH-20



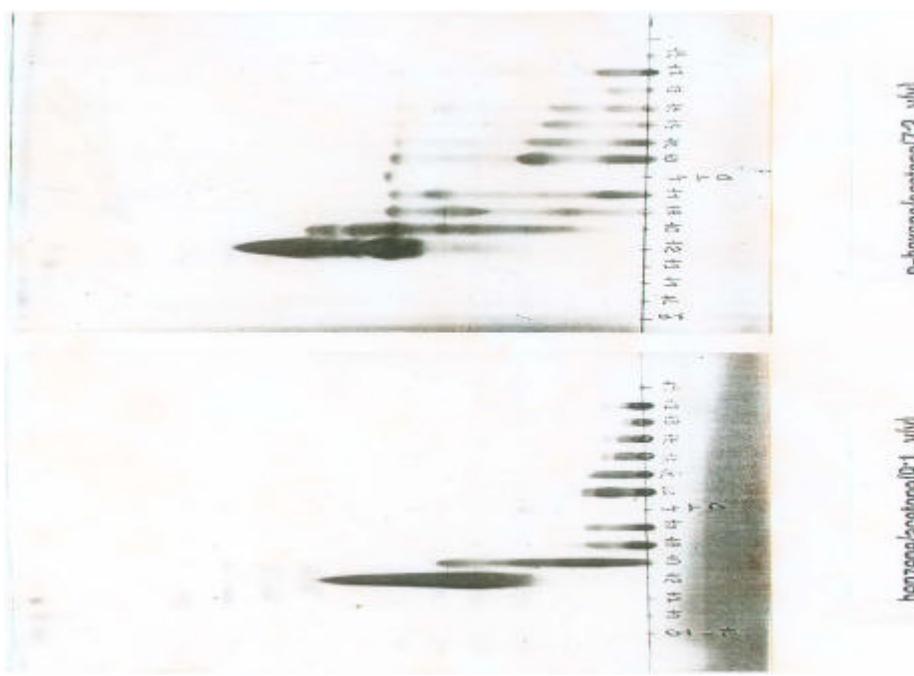
3.20.

TLC



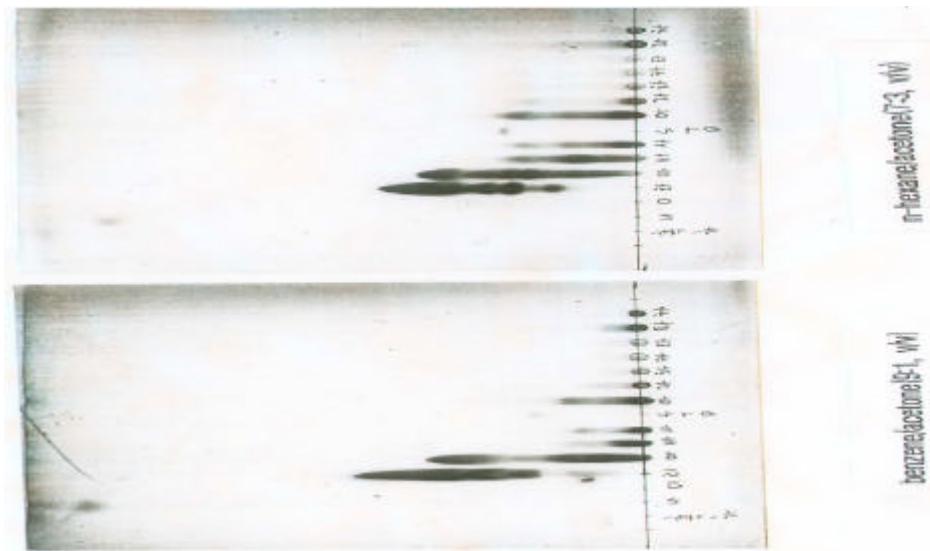
3.21.

TLC



3.22.

TLC



3.23.

TLC

) Sephadex LH-20      chloroform/MeOH(3:1, v/v)      slurry

80ml      10      .      10  
 schizandrin      TLC      , 3, 4, 5      (  
 0.6  1.0)      가      .      ,  
 가

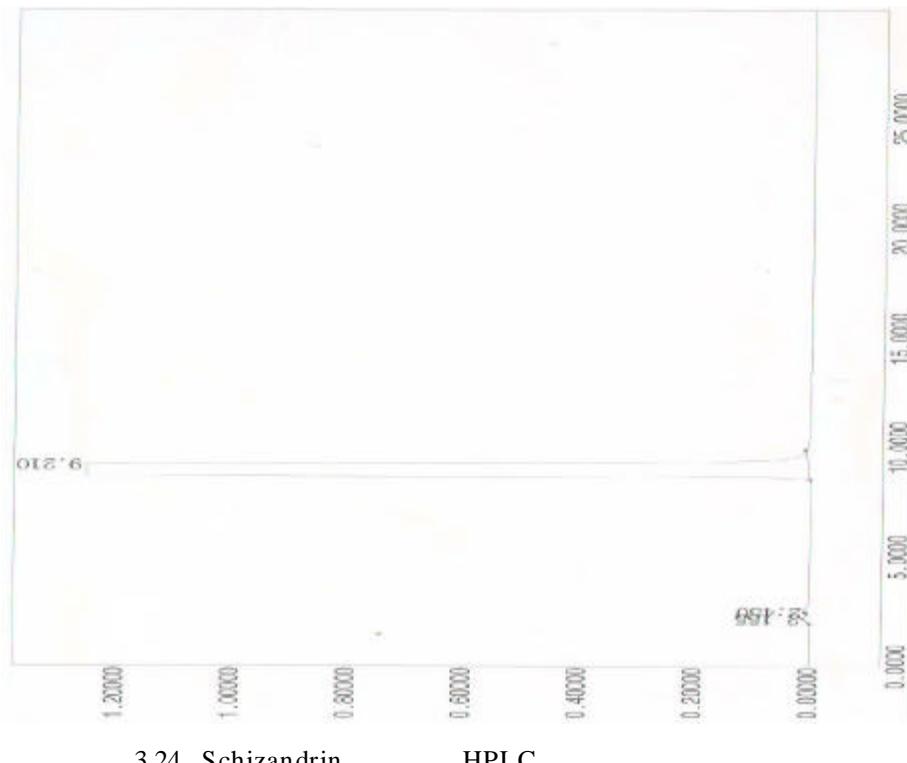
3)

가) Sephadex LH-20      가      3,  
 4,  5      ,      TLC      banding  
 n-hexane/ acetone(7:3, v/v)      schizandrin  
 schizandrin      Rf      MeOH      4      ,  
 ,

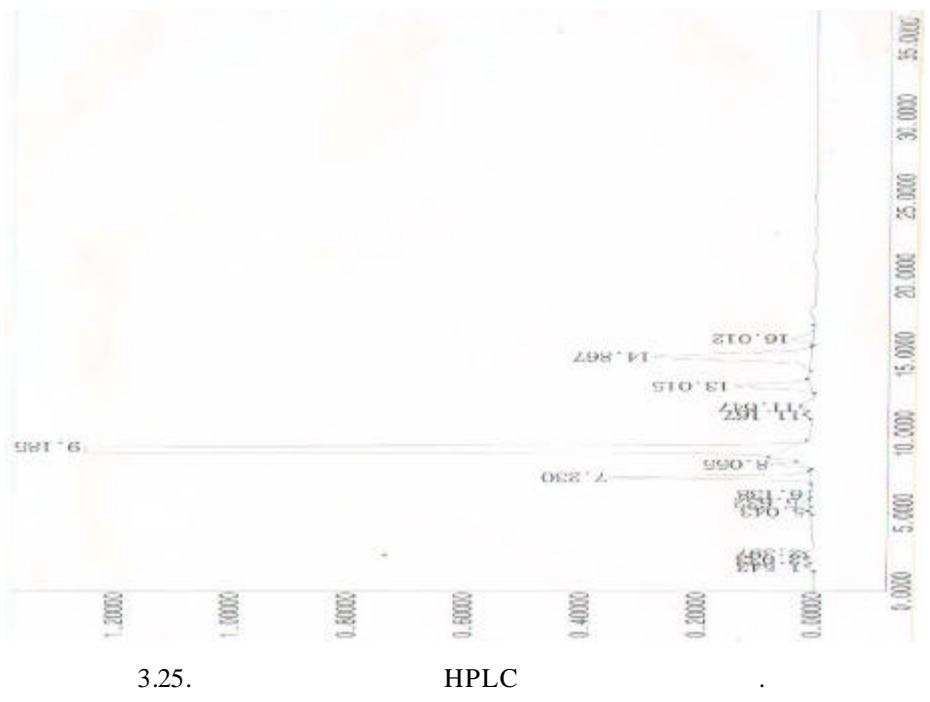
) TLC benzene/ acetone (9:1,  
v/v) 7)

### . HPLC

1) TLC membrane filter(Millipore, 0.45μm, Waters  
) 3.12 HPLC , schizandrin  
3.24, 3.25, 3.26 .  
2) 3.24. schizandrin Rt 9.210 ,  
3.25. Rt 9.185 peak 7) ,  
Rt 7.230, Rt 13.015, Rt 14.867 peak 7)  
3.26. Rt 9.155 peak 7) , Rt  
13.210, Rt 13.715, Rt 14.905, Rt 15.880, Rt 17.982 peak 7)

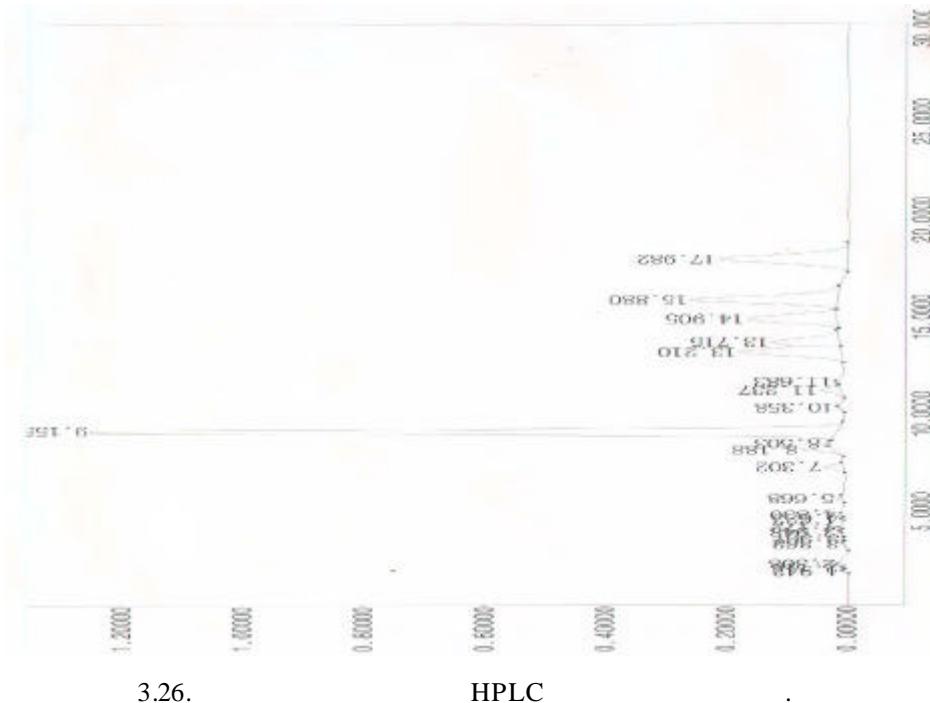


3.24. Schizandrin HPLC



3.25.

HPLC



3.26.

HPLC

3) Rt 9.185 Rt 9.155

가 . Schizandrin

5

schizandrin 가

schizandrin

schizandrin 가

4) LC/MS

sucrose            1.13%     0.02%                                  ,
   
 fructose     glucose                3.71%, 2.51%     1.83%, 1.05%

,
 malic acid                47,684 ppm, 38,691 ppm     が
   
 ,
 citric acid                11,939 ppm, 3,330 ppm     .
   
 malonic acid     maleic acid                2.8 ppm, 2.3 ppm     0.3

ppm, trace    .
   
 mg %                        722 mg%                                      7,577

mg %                        202 mg%, 5,596 mg%
   
 lysine                        300 mg%                                    288

mg %

glutamic acid, aspartic acid, threonine, valine                Leucine                                  .
   
 1.275%,                        1.560%
   
 ,
 が

chlorogenic acid            1,276 ppm, 802 ppm            , gentisic acid
   
 69 ppm,                        136 ppm

ferulic acid                187 ppm,                        23 ppm
   
 .
 cinnamic acid, caffeic acid, coumaric

acid                                  .
   
 145 mg/g                        160 mg/g

,
 linoleic acid が 21,765 ppm, 45,584 ppm     が
   
 palmitic acid が 3,974 ppm, 10,872 ppm     が

capric acid

,  
,

- ylangene, , - elemene, - himachalene,  
- selinene widdrene ,  
caryophyllene, calarene, cubebene, acoradiene - himachalene  
γ  
86.54mg % , 138.75mg %  
2  
peonidin-3-monoglucoside peonidin-3,5-diglucoside ,  
delphinidin-3-monoglucoside malvidin-3-monoglucoside  
  
schizandrin 0.11%/ 100g( ) ,  
0.13%/ 100g( ) .

5

1

, 가

,

가

,

가

가

가가

package

2.5

,

가

가

, 가

가

가 가

가

,

가

가

.

,

,

가

가

, soft drink, extract

( )

,

가

,

가 가가 , 가

가 가

가 route 가 .

가 .

가 가 가

UR , 가

가 가 route 가

## Alloxan 가

茶 가

荟(

가

( ), ( )

가

가

2

1.

가. : 1996 1998 10 11  
4 ,

: 1996 11

4 ( $\pm 1$ )

: 1997 10

- 18 4

: 1997 1999 1

1996 1998 가

4

: 1997 1

0-4

: 80 4 40 7

25%

: 96

2.

†.

, , A.O.A.C. 105  
, Soxhlet , 600  
, (Auto kjeldhal system, Buchi,  
Switzerland) (Fibertec system M  
1017, Tecator, Switzerland) 100  
(† )

4† 2† 2†  
4† .  
1) : , 20% ethanol , 25% ethanol  
2) : 80 40  
3) : 80 3, 4, 5 40 7  
† 4  
20% ethanol 5 80 3  
, 4 , 5 1 1  
3  
45 50 30 ( )  
40 7  
3

4

2†

2†

3, 4, 5, 7

3 (Tokyo Denshoku Co. SP-80,  
Japan) Hunter L ( ), a ( ), b ( )  
spectrophotometer (UV/Visible Spectrophotometer,  
Pye Unicam, U.K.) 520nm

5ml 100ml 5ml Folin  
2ml 100ml 660nm  
catechol

, , pH meter (Orion 720A, U. S. A)  
pH 0.1N NaOH (pH 8.3 )  
(ATAGO, Japan)

27 ( 25% ) , ,  
, , pH, ,  
27 ( 4 ± 1 )

, , 3%  
60 80 4

isoascorbic acid(=erythorbic acid)

0.025%

가

가

가

140 150Mℓ Al-pack

80 5 6

17

(4±1 )

, ,

,

7

..

.

( )

, ,

3%

60 4

5%가 4

가

가

Al-pack

, ,

(4±1 )

가

.

, , ,

1.

, ,

4.1.

가† 86.3%

, , 가

가

가 50%

, ,

,

가

,

가 ,

4.1.

						(가)
( )	86.30	1.05 (7.66)	2.07 (15.11)	3.21 (23.43)	0.68 (4.96)	6.69 (48.83)
( )	19.17	6.23 (7.71)	16.04 (19.84)	15.22 (18.83)	3.85 (4.76)	39.49 (45.17)
( )	22.12	5.58 (7.16)	13.93 (17.89)	15.21 (19.53)	3.90 (5.01)	39.26 (50.41)
( )	10.41	0.84 (0.94)	1.57 (1.75)	44.11 (49.24)	2.60 (2.90)	40.47 (45.17)

( ) (%)

2.

가

4.2. .

4.2. ( % )

( % )	80						40	
	3		4		5		7	
	0	20	0	20	0	20	0	20
( )	9.32	9.68	9.52	10.08	9.40	9.16	8.31	8.40
( )	45.63	39.25	47.04	39.12	47.48	45.45	38.44	35.02
( )	39.78	38.56	40.79	38.74	48.61	41.53	38.39	37.63
( )	12.02	12.13	13.99	15.07	14.31	13.19	9.79	9.45

20% 80

(3 , 4 , 5 ) 40 (7 ) 80

3

80

4 가

3

5

4

20%

5

가

가

10%

80

4

4

8 12% 가

가

1 3%

가

가

(3

5%)

, 80

4

80

4

20%

가 ( , )

가

가 가

가

45%

90 3

가

가

3.

1)

(a )

가

(

)

가

a , b , L

a 4.3.

가

(80 )

(40 )

, 80 4

a , 40 7

20%

a 가

80 5 40 7

a

a

a

4.3.

(a )

a ( %)	80						40	
	3		4		5		7	
	0	20	0	20	0	20	0	20
( )	47.57	67.62	65.97	70.05	60.81	68.62	60.07	63.44
( )	26.75	27.53	28.63	27.65	27.97	29.34	32.48	35.70
( )	13.38	18.36	12.58	19.14	17.39	20.60	16.55	22.16
( )	3.44	11.02	5.52	10.65	7.18	12.06	0.22	2.53

(80 )

(40

)

a

20%

a 2 ,

a

, a 가

2)

(b )

가

b

4.4.

4.4.

(b )

(% ( ))	80						40	
	3		4		5		7	
	0	20	0	20	0	20	0	20
( )	30.26	56.64	58.92	41.04	68.41	56.14	34.68	50.87
( )	32.33	32.69	31.83	32.90	31.87	48.04	22.46	19.02
( )	21.91	32.23	23.68	36.00	18.14	40.48	14.30	19.80
( )	40.91	55.08	47.59	55.90	53.40	59.62	35.19	54.24

b 가 (80 , )

80 20%

40 7

a

가

b

a

b

5

b

20%

b

b

20%

b

b

4.

1)

(L )

가

1

45.

4.5.

(L )

(% )	80						40	
	3		4		5		7	
	0	20	0	20	0	20	0	20
( )	29.59	17.08	18.40	8.03	19.71	16.79	14.35	23.32
( )	49.54	39.49	50.51	47.95	45.81	30.10	66.02	64.68
( )	71.58	65.08	68.19	45.13	63.85	50.35	76.92	69.66
( )	79.64	60.61	75.17	61.82	72.85	59.37	84.00	81.08

L ( ) 80

4

20%

L ( )

L

L ( )

20%

40

7

L

L 가

가

L

alcohol

L

2)

(OD )

4가

( )

520nm

4.6.

4.6.

(OD)

	80						40	
( % )	3		4		5		7	
	0	20	0	20	0	20	0	20
	( )	1.71	4.54	4.52	7.52	3.90	4.60	4.52
( )	1.26	1.19	1.32	1.34	1.43	1.81	0.93	0.94
( )	0.52	0.72	0.55	1.04	0.68	0.90	0.46	0.54
( )	0.26	0.52	0.36	0.57	0.43	0.66	0.19	0.25

OD 80 20% 4

3 7

20% OD 20%

가 . 20%

가

OD 가

3

2 가 ,

2 가

5.

1) pH

pH 4.7 .

pH 3.39 3.62

20%

pH

pH 40 80

pH 가

20%

pH

pH

4.7.

pH

( % )	80						40	
	3		4		5		7	
	0	20	0	20	0	20	0	20
( )	3.42	3.62	3.41	3.48	3.39	3.57	3.45	3.57
( )	3.32	3.49	3.26	3.44	3.13	3.46	3.26	3.41
( )	3.02	3.21	3.02	3.22	2.98	3.24	3.00	3.24
( )	5.36	5.63	5.32	5.54	5.29	5.56	5.49	5.55

pH

가 가

pH

pH

가

pH 4.0

pH 4.5

, pH 5.0

pH 3.6

pH

pH 5.3

2)

4.8.

4.8.

(%)

(% ( ))	80						40	
	3		4		5		7	
	0	20	0	20	0	20	0	20
( )	2.2	2.6	2.6	2.6	2.4	2.0	1.4	2.2
( )	2.8	2.0	3.0	2.2	2.8	2.2	2.2	2.0
( )	1.6	2.0	1.8	2.0	2.2	2.0	1.6	2.0
( )	0.8	0.8	1.0	1.0	1.0	1.0	0.6	0.4

80

4

4

20%

가

가 가 , ,  
가 ,

2 3%

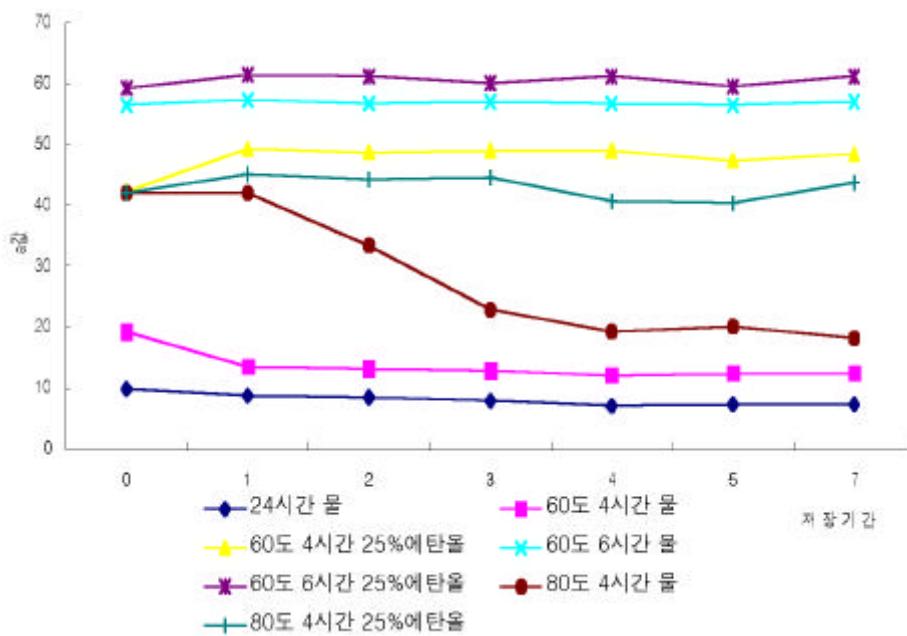
50%

3%

6.

1)

(a )



4.1.

(a )

가

, ,

Hunter a ( )

4.1.

1

80

4

60 6

25%

60 4

80

4

25%

(17 19 ) 24

가

60 6 (25%) 6

(a )

7

가

80 4

a

a

80 4

a

가

가

,

2)

(b )

가

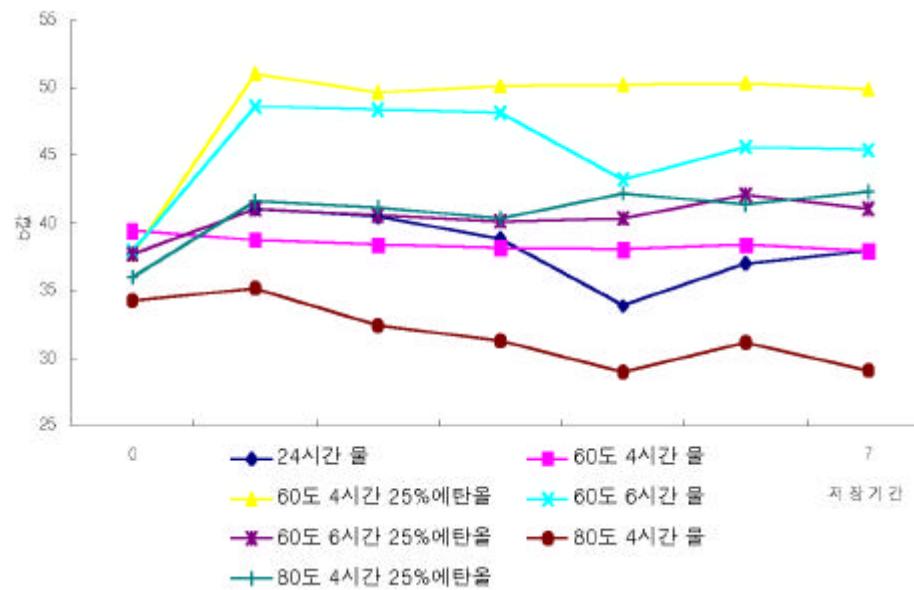
가

가

Hunter

4.2.

b ( ) 60 4  
 60 4 25% 가 가  
 60 6 가 가 80 4  
 가 .  
 24  
 1 가 .  
 가 .

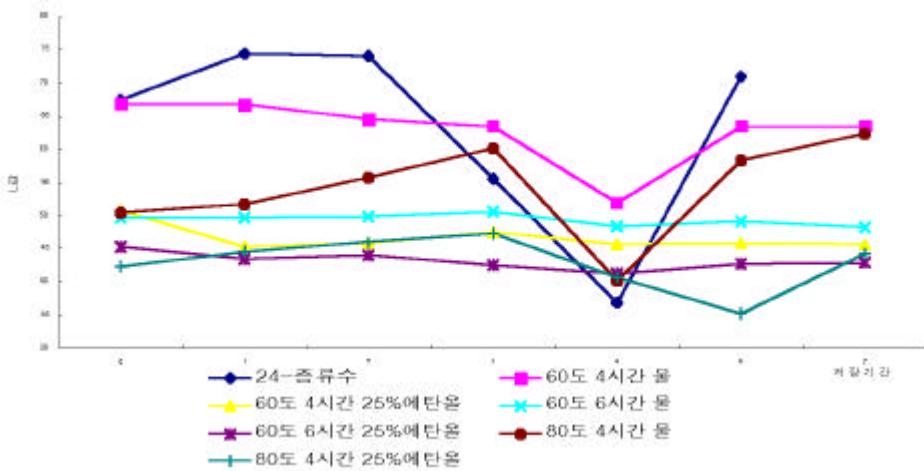
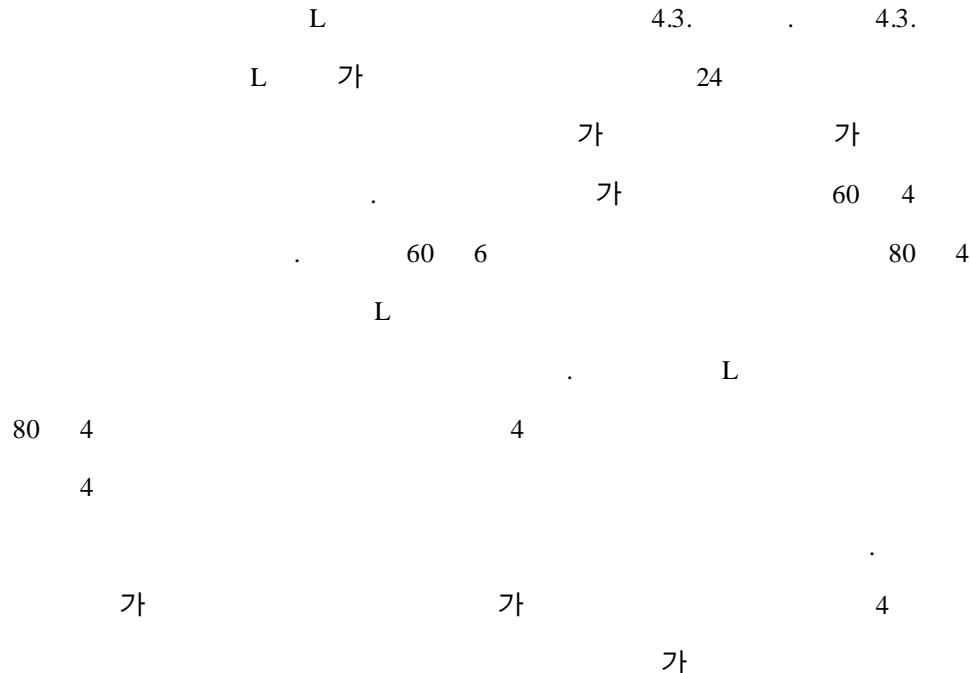


4.2. (b )

가 가 가  
 . 80 4 , 60 4 , 60 6  
 4  
 a (L ) 가 .

3)

(L )



4.3.

(L )

4)

520 nm  
4.4.

(OD) 60 6 25%

가 가 , 60 6

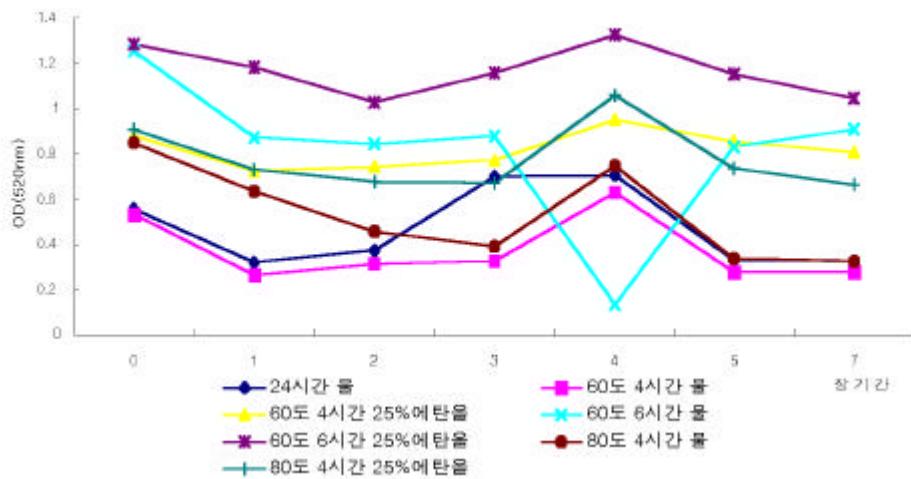
24 60 4

2가

4

가

3



4.4.

(OD)

5)

(total polyphenol)

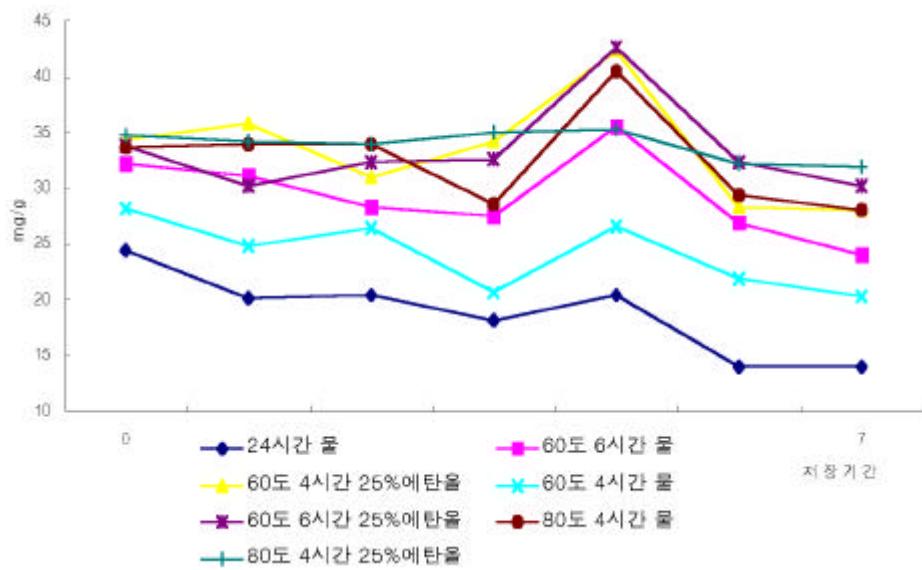
(polyphenol)

2

가

가

4.5.



4.5.

24

60 4

25%

4

4

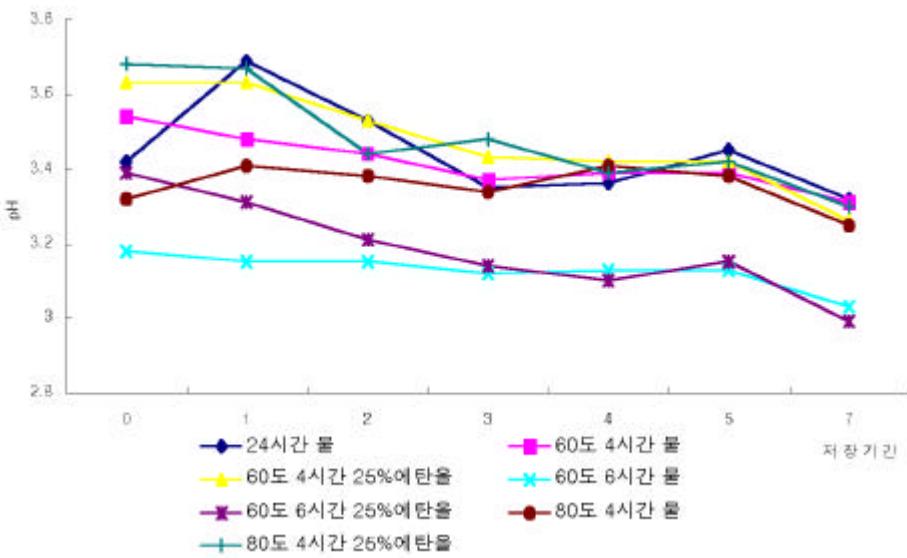
가

가 . 25%

가

가 . 가

6) pH



4.6.

pH

pH

pH

4.6.

pH 3.68 2.99

6

0 6 25%

pH 3.6 3.3

24

pH

pH

pH

pH

pH

pH 4.0

7)

가

가

0.1N-NaOH

가

4.7.

가

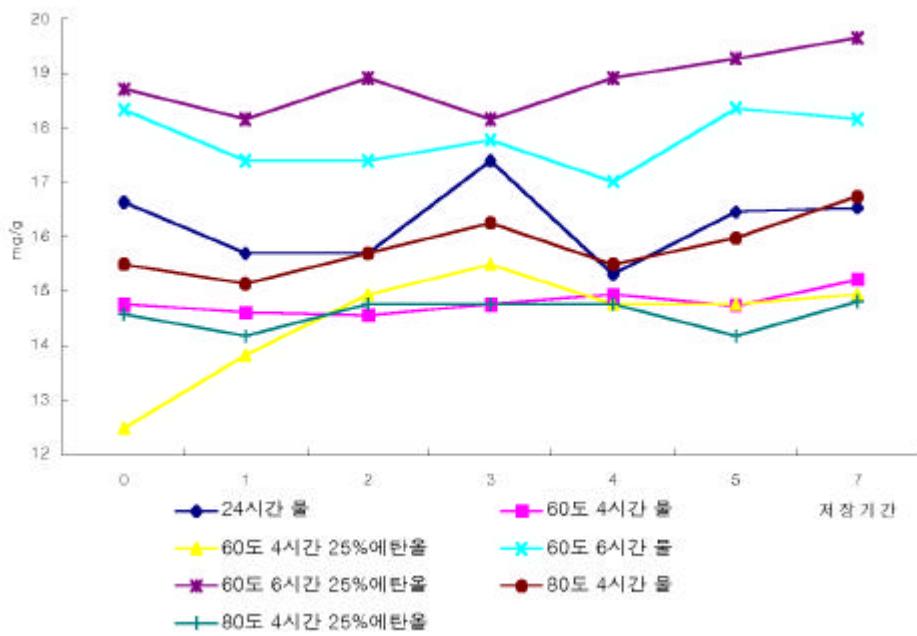
60 6 25%

가 가

24

80 4

pH



4.7.

가

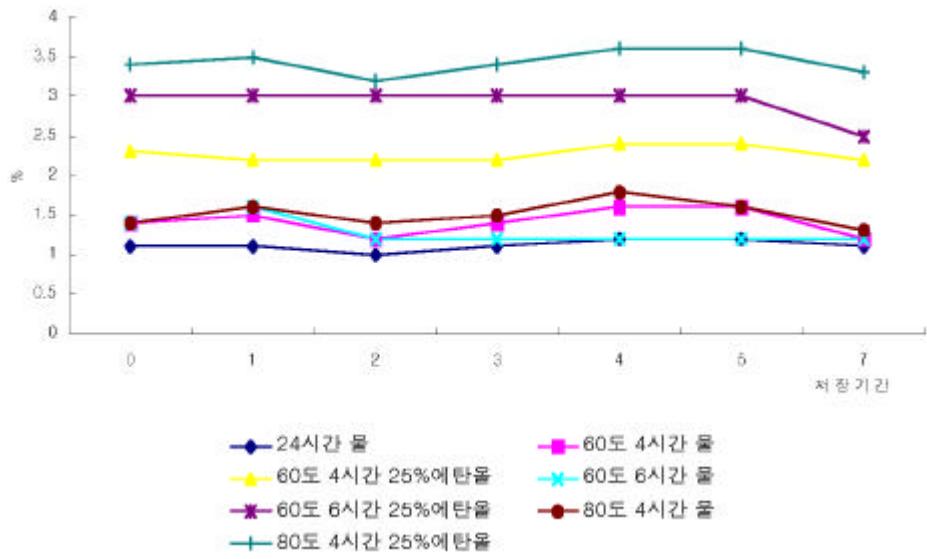
가

8)

4.8.

가

가



4.8.

80      4      25%	60      6
가	
. 25%	
2.2      3.6	1.0      1.8
가 2  3%	

가

7.

1)

(a )

가

3

a

4.9.

4.9.

a

	1	2	3	4	5	6	7	8	9	10	13	14	20	21
80	52.6	51.6	52.0	51.4	51.8	51.6	53.1	51.8	50.9	51.2	51.8	51.5		
	49.3	48.2	49.0	48.5	49.3	48.4	50.2	48.7	48.8	48.6	48.8	48.3		
60	46.1	45.4	45.7	45.6	46.3	45.6	48.4	45.7	46.2	45.8	46.5	46.4	45.6	46.8
	42.0	40.5	40.9	40.8	41.3	40.9	42.9	40.3	41.3	41.0	41.8	41.5	40.9	41.8
80	40.2	39.8	39.4	39.3	39.3	39.6	40.2	39.4	40.1	39.4	39.2	39.2	38.9	40.9
	36.3	35.8	35.4	34.9	36.3	35.8	37.0	35.7	35.7	35.1	34.3	35.0	34.9	33.4
60	41.8	41.4	41.3	41.2	41.8	41.8	42.5	41.6	42.0	41.6	41.7	41.2	42.9	42.8
	43.5	43.1	43.2	42.7	43.2	43.4	44.4	43.5	43.5	43.1	43.4	43.2	41.0	45.2
60	11.8	11.8	11.9	11.9	7.0	8.2	8.7	7.8	8.2	7.9	8.3	8.2	6.1	5.9
	12.2	12.0	12.1	12.0	6.8	7.8	8.3	7.8	7.9	7.8	8.0	7.6	5.7	5.5

, ,

,

60      80

1%

3%

a

가

가

가

가

가

1/4 1/5

가

4

80 가 60

가

,

가

60 가

60

3%

가

.

21

a ( )

가

.

60

3%

20 (5 )

가

2)

가

(a )

,

erythorbic acid

(iso ascorbic acid) 0.025%

가

가

ion

,

가

가

a

4.10.

.

4.10.

가

a

	1	2	3	4	5	6	7	8	9	10	13	14	20
80	50.9	50.8	51.2	51.7	51.8	50.9	51.0	52.7	53.9	50.8	51.2	50.1	
	47.2	47.4	47.4	48.0	48.0	46.6	46.2	47.7	48.4	45.9	46.1	45.8	
	44.9	44.3	45.0	43.8	44.5	43.8	43.0	45.0	45.3	43.5	44.1	42.7	38.8
60	38.3	37.3	37.6	37.8	37.8	36.9	36.7	38.2	38.0	36.3	36.2	35.4	35.4
	36.2	36.2	36.4	36.5	36.6	35.5	35.0	36.6	37.2	35.2	32.6	34.5	35.2
	32.3	32.8	32.6	32.8	32.8	31.0	31.7	32.0	31.8	32.0	30.8	30.8	27.5
80	40.2	38.0	38.3	38.9	38.6	38.5	37.8	37.4	39.4	38.2	32.9	37.5	37.8
	42.6	40.5	40.4	40.6	39.2	40.5	39.9	40.0	41.2	39.5	39.4	39.0	39.4
	11.8	10.0	9.8	10.5	6.8	6.6	5.8	6.8	5.8	5.8	6.2	5.7	6.4
60	11.4	10.1	10.0	10.0	6.2	6.5	5.9	6.2	6.0	5.9	5.8	5.6	5.6

가

a

가

a

2 5

a

a

가

a

,

20 ( 5 )

pH

,

3)

(b )

b

4.11.

b 80

가

60

가

3%

b (a )

b 3 4

가

b 10

가

가

가

b 가

b

b a L

가

가

4.11.

b

	1	2	3	4	5	6	7	8	9	10	13	14	20	21
80	40.7	41.2	41.3	41.9	41.7	41.8	42.8	42.4	41.6	42.6	43.0	44.2		
	37.7	38.0	38.2	38.8	38.8	38.4	38.6	38.9	39.2	39.3	39.3	39.7		
60	30.6	30.7	30.8	31.2	31.1	31.0	31.3	31.4	31.6	31.5	31.1	31.5	31.4	33.0
	29.8	29.7	29.9	30.3	30.3	30.3	30.4	29.3	30.7	30.7	30.5	30.8	30.8	32.2
80	40.7	41.0	41.7	42.3	42.9	42.7	43.4	44.0	43.0	44.4	47.0	47.3	48.2	52.4
	30.6	30.6	30.6	30.8	31.6	31.1	31.6	31.9	32.5	32.6	32.0	33.8	34.8	39.3
60	37.8	38.8	39.6	39.4	39.5	39.4	39.5	39.9	40.7	40.3	40.5	41.5	46.5	51.9
	42.2	43.4	43.4	44.0	44.1	44.5	44.1	44.0	44.7	44.9	44.9	45.1	45.0	56.5
60	10.3	10.9	10.9	11.2	6.7	7.6	7.5	7.5	9.4	7.7	8.0	8.1	6.7	6.8
	9.9	10.4	10.4	10.7	6.4	7.2	7.4	7.3	9.5	7.6	7.6	6.7	6.4	6.3

4)

가

(b )

, erythorbic

acid(iso ascorbic acid)

가

,

b

4.12.

가

가

,

가

a

erythorbic acid

가 a, b

가

4.12.

가

b

	1	2	3	4	5	6	7	8	9	10	13	14	20
80	39.3	38.8	40.4	41.0	42.0	40.2	42.1	42.6	42.4	42.1	42.2	41.9	
	37.0	37.2	37.8	38.5	38.9	36.2	36.8	37.2	38.3	38.5	37.6		
	30.4	29.5	28.5	29.4	29.8	29.1	29.5	29.8	29.3	29.2	28.9	28.8	30.1
60	30.3	30.0	30.4	30.6	30.7	30.4	31.0	31.1	30.3	30.7	30.1	30.4	30.0
	38.1	34.6	36.0	35.8	35.8	36.9	39.0	39.6	37.1	36.4	33.4	38.2	37.8
80	29.0	27.9	28.6	28.9	28.4	29.3	30.3	30.2	31.0	28.6	30.7	30.0	29.0
	30.0	31.0	32.2	32.7	32.2	32.0	32.7	32.5	32.4	32.5	32.2	31.6	31.4
60	35.0	36.5	37.0	36.6	36.2	36.3	36.6	36.2	35.8	35.4	34.6	35.7	34.6
	9.6	9.4	9.8	10.2	7.2	6.4	7.2	7.4	7.3	6.9	7.4	7.1	7.6
60	9.4	9.7	9.6	10.1	6.9	6.5	6.9	7.3	7.0	7.2	7.5	7.6	7.6

5)

(L )

L ( )

2가

가

L

L

a ( )

L 80 4

, 60 4 , 80 60

60 4 L

a

3%

L

가

가

가

2가 가 가

가가 4.9 a 가

1

a

( 4.13).

4.13.

L

6)

가

(L )

가

L

4.14.

L ( )

가

, ,

L

L

3.10.

a

a 가 L

, b

L

가

L 가

가

가 가

4.14.

가

L

	1	2	3	4	5	6	7	8	9	10	13	14	20
80	45.2	45.0	44.5	45.1	43.9	41.4	43.2	42.8	40.2	43.2	41.6	40.4	
	47.9	47.2	47.0	47.5	46.9	45.1	46.9	46.2	44.1	46.4	45.8	46.1	
60	51.5	53.2	52.1	52.8	52.4	53.3	53.5	51.0	50.7	52.7	51.3	52.5	51.6
	54.6	55.2	56.0	56.9	55.4	55.8	56.4	53.7	53.8	55.8	55.4	55.9	55.4
80	54.6	56.8	55.1	55.6	54.9	52.8	53.4	51.0	53.6	53.9	53.4	50.5	52.4
	62.9	64.5	63.6	63.6	63.4	61.5	60.6	60.9	60.9	61.3	58.3	61.5	71.9
60	60.8	62.6	61.1	61.1	60.5	59.8	59.9	58.1	57.9	59.9	57.4	59.6	59.8
	60.5	61.9	61.1	62.0	61.3	60.9	62.8	62.0	60.8	62.4	61.8	60.9	62.6
60	68.9	66.8	66.9	70.9	73.7	75.2	78.0	79.1	78.4	79.1	76.6	79.9	79.3
	79.6	79.1	79.5	79.9	80.5	80.7	85.4	84.8	84.4	85.9	84.9	84.6	85.9

7)

(OD )

520nm

4.15.

가

60

가 가

80

80

60

a

가

520nm

4.15.

	1	2	3	4	5	6	7	8	9	10	13	14	20	21
80	1.62	1.66	1.71	1.67	1.67	1.61	1.70	1.67	1.60	1.57	1.58	1.66	1.65	
	1.56	1.49	1.54	1.52	1.34	1.43	1.39	1.55	1.43	1.45	1.45	1.48	1.48	
	1.32	1.30	1.33	1.31	1.15	1.22	1.26	1.30	1.30	1.21	1.18	1.28	1.16	1.23
60	1.13	1.12	1.12	1.09	1.13	1.07	1.10	1.16	1.06	1.08	1.11	1.09	1.03	1.05
	1.69	1.70	1.75	1.70	2.02	1.67	1.76	1.79	1.56	1.69	1.72	1.52	1.50	1.53
80	1.25	1.26	1.26	1.28	1.09	1.18	1.23	1.30	1.26	1.21	1.30	1.16	1.29	1.18
	1.77	1.62	1.66	1.70	1.49	1.54	1.62	1.63	1.62	1.70	1.62	1.55	1.52	1.54
60	1.82	1.74	1.90	1.87	1.86	1.75	1.86	1.87	1.83	1.91	1.78	1.66	1.63	1.66
	0.48	0.44	0.46	0.44	0.21	0.24	0.24	0.24	0.22	0.24	0.24	0.24	0.17	0.18
60	0.38	0.37	0.38	0.24	0.18	0.19	0.20	0.20	0.19	0.19	0.20	0.23	0.14	0.14

a

가

20 , 21

가

8)

가

(OD )

가

4.16.

4.16.

가

	1	2	3	4	5	6	7	8	9	10	13	14	20
80	1.66	1.51	1.51	1.54	1.58	1.52	1.58	1.56	1.55	1.56	1.58	1.59	
	1.36	1.40	1.39	1.39	1.46	1.42	1.40	1.36	1.40	1.33	1.40	1.40	
	1.25	1.20	1.22	1.24	1.28	1.21	1.24	1.16	1.18	1.12	1.19	1.18	1.29
60	1.03	1.07	1.02	1.00	1.06	1.02	1.02	1.06	1.02	0.98	1.03	0.94	1.05
	1.62	1.52	1.46	1.49	1.52	1.45	1.53	1.48	1.60	1.38	1.32	1.27	1.27
	1.16	1.16	1.10	1.10	1.15	1.09	1.18	1.12	1.13	1.07	1.08	0.97	1.00
80	1.62	1.49	1.46	1.49	1.51	1.43	1.49	1.38	1.40	1.34	1.37	1.32	1.28
	1.74	1.66	1.64	1.61	1.65	1.57	1.69	1.49	1.55	1.38	1.42	1.33	1.32
	0.43	0.44	0.42	0.41	0.23	0.23	0.22	0.23	0.20	0.20	0.22	0.20	0.18
60	0.34	0.34	0.32	0.32	0.18	0.22	0.17	0.17	0.17	0.16	0.16	0.17	0.17

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9)  
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가  
(polyphenol)  
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가 가	가 가	.	polyphenol
가		가	
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polyphenol	14		
		21	
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polyphenol	가		가
		가	
		.	

4.17.

(mg/ 100Mℓ)

	1	2	3	4	5	6	7	8	9	10	13	14	20	21
80	72.8	73.5	74.2	73.5	67.0	67.6	72.5	63.8	70.4	68.2	67.8	67.1	64.9	
80	72.8	72.1	73.5	72.1	60.0	71.1	71.1	67.1	68.5	66.7	66.7	68.5		
60	67.3	68.0	69.3	67.3	65.0	64.8	66.7	60.5	60.2	64.8	62.3	63.4	61.2	48.4
80	67.3	66.6	68.0	67.3	64.0	59.1	66.2	65.1	64.8	64.8	64.4	58.0	57.7	48.4
	57.7	56.6	59.8	58.7	55.2	55.3	56.4	54.6	56.3	56.9	54.4	54.9	54.9	43.6
80	57.1	55.5	57.7	57.1	50.4	51.8	55.8	52.4	54.7	55.0	49.5	53.6	49.5	45.3
60	57.1	56.6	58.2	57.1	54.4	53.8	56.8	52.7	54.6	55.5	53.8	53.0	55.5	43.6
60	56.0	56.0	58.2	57.1	53.6	54.6	56.8	52.4	53.8	55.3	52.7	53.5	52.7	43.6
60	66.0	67.6	67.6	68.0	59.6	61.8	61.2	59.0	58.4	58.1	59.1	56.8	55.2	43.2
	64.0	68.4	66.0	64.4	58.4	56.1	60.0	59.6	58.1	57.8	57.8	57.1	56.4	41.2

10)

가

erythorbic acid      가

4.18.

가      가

1

가

가  
가

20

가

14

가

가가      가

4.18.

가

pholypheol                  (mg/ 100Mℓ)

	1	2	3	4	5	6	7	8	9	10	13	14	20
80	74.9	72.8	73.9	70.0	72.4	73.4	71.6	69.4	70.7	70.5	72.6	69.8	
	73.5	72.1	72.5	70.7	72.1	73.3	71.3	67.0	70.4	69.5	72.6	68.6	
	71.4	68.7	69.4	67.3	65.3	71.6	75.9	66.6	66.2	67.3	68.0	61.9	64.8
60	70.0	68.7	69.4	68.7	69.4	72.0	72.3	64.3	66.2	59.8	64.4	60.9	64.1
	59.3	58.7	58.7	57.1	55.7	61.4	61.5	55.5	58.1	55.4	57.5	55.3	54.4
80	58.7	56.6	57.6	55.4	53.3	60.9	61.1	53.2	55.7	54.3	55.3	53.6	51.8
	58.7	55.7	58.2	55.5	54.1	60.8	59.6	56.0	56.4	54.4	56.9	54.7	54.7
60	58.2	58.2	57.6	55.5	54.3	62.6	60.7	56.0	55.5	53.5	56.3	55.0	54.1
	69.2	67.6	69.6	67.8	67.8	67.9	69.2	66.8	63.3	65.5	61.5	60.1	59.8
60	67.6	66.8	66.0	67.2	66.6	66.6	68.4	65.1	60.6	64.5	60.5	59.9	60.1

11)

4.19.

가

4.19.

(%)

	1	2	3	4	5	6	7	8	9	10	13	14	20	21
80	0.8	0.8	1.0	0.9	1.0	1.1	1.1	1.4	1.0	1.1	1.4	1.2		
	0.8	0.6	0.6	0.9	1.0	1.0	1.0	1.2	1.0	1.0	1.2	1.0		
	1.0	0.8	1.0	1.1	1.2	1.2	1.2	1.4	1.4	1.4	1.6	1.2	1.8	1.8
60	0.8	0.6	0.8	0.9	1.0	1.0	1.1	1.2	1.1	1.1	1.4	1.2	1.4	1.4
	1.6	1.6	1.8	1.7	1.9	1.9	1.9	1.9	1.9	2.1	2.1	2.1	2.1	1.9
	1.1	1.0	1.1	1.4	1.3	1.6	1.6	1.6	1.6	1.6	1.6	1.6	1.7	1.6
80	1.0	1.4	1.6	1.6	1.6	1.6	1.7	1.8	1.8	1.8	1.6	1.8	1.8	1.9
	1.4	1.4	1.6	1.6	1.6	1.6	1.8	1.8	1.8	1.8	1.8	1.8	1.8	1.9
	0.6	0.6	0.7	0.7	0.4	0.4	0.5	0.6	0.4	0.5	0.6	0.5	0.5	0.4
60	0.5	0.5	0.6	0.7	0.3	0.3	0.4	0.4	0.4	0.4	0.5	0.4	0.4	0.4

가

가

가

가

가 가

12) 가

가

4.20.

4.20.

가

(%)

		1	2	3	4	5	6	7	8	9	10	13	14	20
80		0.8	0.8	0.8	1.1	1.2	1.0	1.0	1.2	1.0	1.2	1.2	1.2	1.2
		0.8	0.8	1.0	1.0	1.0	0.8	1.0	1.2	1.0	1.0	1.0	1.0	1.4
		1.0	1.0	1.4	1.1	1.3	1.0	1.4	1.4	1.4	1.6	1.8	1.8	2.0
60		0.8	0.8	1.2	1.0	1.2	1.0	1.0	1.2	1.0	1.2	1.4	1.4	1.4
		1.8	1.6	1.8	1.7	1.8	1.8	1.9	1.9	1.9	1.9	2.1	2.1	2.1
80		1.6	1.4	1.4	1.5	1.4	1.2	1.6	1.6	1.6	1.6	1.6	1.6	1.4
		1.6	1.4	1.6	1.6	1.8	1.6	1.8	1.9	1.8	1.9	1.9	1.9	1.9
60		1.6	1.4	1.6	1.6	1.8	1.6	1.8	1.9	1.6	1.9	1.9	1.9	1.9
		0.8	0.7	0.8	0.8	0.5	0.4	0.5	0.5	0.4	0.5	0.6	0.6	0.4
60		0.6	0.6	0.6	0.8	0.4	0.4	0.4	0.5	0.4	0.4	0.5	0.4	0.5

가

가

가 . . . . .  
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가 . . . . .  
가 . . . . .  
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가 . . . . .

13)

가 . . . . .  
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4.21. . . . .  
가 . . . . .

가 . . . . .  
1/3 1/6 . . . . .  
3% . . . . .  
가 . . . . .

가 . . . . .  
20 . . . . .

6 . . . . .

가 . . . . .

4.21.

(mg/g)

	1	2	3	4	5	6	7	8	9	10	13	14	20
80	42.8	43.5	43.7	43.7	42.0	41.3	38.8	41.3	40.1	38.0	39.3	41.6	
	42.2	43.5	42.8	43.7	40.9	40.4	37.7	39.4	41.1	39.1	40.5	40.6	
	23.3	22.9	21.4	23.6	23.3	22.7	22.7	22.1	23.4	21.3	21.3	22.3	
60	23.6	23.1	24.0	24.2	23.8	23.0	23.5	21.8	23.3	23.4	22.5	23.9	
	146.1	131.4	130.6	153.8	140.2	140.2	140.0	138.2	139.5	138.0	136.6	136.3	130.2
	148.7	143.0	143.4	157.7	144.2	141.5	140.9	125.8	143.8	133.2	137.8	138.9	134.9
60	122.6	138.2	137.4	142.6	125.4	123.0	128.8	129.1	129.1	127.1	142.6	129.8	120.9
	115.8	118.6	109.6	122.7	113.6	109.0	112.9	115.5	110.4	110.0	1103. 7	109.5	107.9
	151.5	150.6	151.8	145.7	145.6	123.5	123.2	121.8	107.0	116.2	123.6	123.6	113.5
60	147.3	143.8	150.1	149.6	151.5	123.5	126.8	114.9	119.4	120.9	123.6	125.6	116.0

14)

가

가

4.22.

가 가

가 가

,

5

20

가

가

pH

가가

4.22.

가

(mg/g)

	1	2	3	4	5	6	7	8	9	10	13	14	20
80	45.7	44.7	43.9	44.8	43.4	42.3	43.4	42.4	42.7	45.8	44.2	43.4	44.5
	45.9	39.4	43.4	45.9	44.2	42.5	43.4	45.7	46.5	41.1	42.4	42.4	44.2
	24.2	22.9	25.2	28.7	24.4	24.1	23.0	23.1	25.1	22.9	24.1	25.6	
60	24.5	25.9	27.8	27.9	25.2	25.8	21.9	22.3	27.0	26.1	24.1	24.7	
	148.5	133.7	145.4	149.4	138.6	141.3	137.8	137.7	141.2	138.6	125.0	139.5	135.3
	146.7	141.4	147.6	159.2	145.0	143.4	144.8	139.5	139.0	138.5	115.0	143.0	141.8
60	139.8	131.1	140.8	139.8	127.4	131.0	135.0	130.6	131.1	125.7	116.9	133.0	141.8
	148.2	114.1	120.2	119.0	116.8	116.6	113.3	103.0	116.6	109.5	109.9	103.2	109.8
	148.3	136.0	150.4	153.7	124.3	125.5	121.7	121.8	123.7	123.2	114.5	127.6	128.9
60	155.4	143.2	151.6	149.8	123.3	126.6	122.5	117.9	119.2	123.6	115.2	124.6	125.6

15)

pH

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pH

4.23

pH

4.23.

pH

16) 가 pH

(iso ascorbic acid)

가 가 pH

4.24.

4.24. , 가 pH

가 pH 2.80 3.20 2.82

3.41 pH 2.53 2.88 2.66 2.80

pH 2.69 2.97 가 pH

가

pH가

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4

7

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(isoascorbic acid)

20

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