



UNIVERSITI PUTRA MALAYSIA

***PHENOTYPIC AND MOLECULAR VARIATION AMONG SELECTED
Curcuma alismatifolia GAGNEP. MUTANTS DERIVED FROM ACUTE
AND
CHRONIC GAMMA IRRADIATION***

SIMA TAHERI

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By

SIMA TAHERI

**Thesis Submitted to the School of Graduate Studies, Universiti Putra
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Philosophy**

November 2014

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Dedication

This thesis is dedicated to

My beloved father and mother

Whose love and blessings have been sources

Of Inspiration to me

And my

Beloved sisters Sara and Sepideh

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Abstract of thesis presented to the Senate of Universiti Putra Malaysia in
Fulfillment of the requirement for the degree of Doctor of Philosophy

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November 2014

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Faculty: Agriculture

Studies were conducted to study phenotypic and molecular variation among selected *Curcuma alismatifolia* mutants through acute and chronic gamma irradiation and to develop new variants of *C. alismatifolia*. In acute gamma irradiation, rhizomes in the sprouting bud stage were irradiated at eight different doses of 0, 10, 20, 25, 35, 40, 60, and 100 Gy. Radiation sensitivity tests revealed that the LD₅₀ of the varieties were achieved at 21 Gy for 'Chiang Mai Red', 23 Gy for 'Sweet Pink', 25 Gy for 'Kimono Pink', and 28 Gy for 'Doi Tung 554'.

From the analysis of variance (ANOVA), significant variations were observed for vegetative traits, flowering development, and rhizome characteristics among the four cultivars of *C. alismatifolia* and dose levels as well as the dose × variety interaction. In first generation (M₁V₁), as many as 52 chlorophyll mutants were observed, resulting in a mean mutation frequency of 38.5 % in which 13.1 and 19.3% were produced by 10 and 20 Gy doses, respectively. Individual mutant DT20-1 with two-tone purple bract color was the most attractive mutant generated.

In SSR analysis, the obtained results indicated the high efficiency of polymorphic SSR loci for genetic studies among treated and non-treated *C. alismatifolia* individual plants. 20 Gy acutely irradiated individuals showed a higher mean percentage of polymorphic loci (62.5%) than the 10 Gy (59.38%) and non-treated (22%) ones. By HRM analysis, all studied irradiated individual plants produced melting curves with different melting temperature (T_m) from their control individual plants.

In chronic gamma irradiation, plants that were exposed to radiation at doses of 14.6, 33, and 87.4 Gy showed significant decreases in the vegetative traits as compared to the controls. Interestingly, low doses of gamma irradiation stimulated the vegetative growth and promoted earlier flowering of mutants compared to controls. However, higher dose rates decreased the flowering capacity and reduced the quality of the rhizomes of *C. alismatifolia* mutants. In SSR analysis, the overall genetic variability for the varieties studied showed that chronic gamma irradiation particularly at higher dose rates (14.6, 33, and 87.4 Gy) were able to induce more genetic variations to genome of studied *C. alismatifolia* individual plants.

In order to study the morphological and genetic stability of mutants, for both acute and chronic gamma irradiations, studies were continued into the second generation (M_1V_2). In acute gamma irradiation chlorophyll mutation frequency decreased significantly at 10 and 20 Gy irradiated plants in M_1V_2 . Individual DT20-1 with two tone purple color at first generation produced solid mutant with deeper purple bracts in M_1V_2 . In order to gain distinctness, uniformity and stability (DUS) of mutants, the rhizomes of all individual plants were grown in M_1V_3 generation and observations were continued. Nine individual plants with bract color variations and dwarfism were selected for further studies in M_1V_4 generation. DT20-1 mutant that maintained the deep purple color in M_1V_3 was considered a desirable potential mutant. SSR analysis in the second generation showed that the percentage of polymorphic loci decreased from 59.3% to 40.6% for 10 Gy individual plants. Similarly, at 20 Gy, percentage of polymorphic loci decreased from 62.5% to 50.0 %.

In chronic gamma irradiation, none of the flower shape and color variations observed in first generation were transferred to the next generation. The effects induced by chronic gamma radiation on these plants are more likely to be physiological changes at sublethal dosages rather than irreversible genetic effects. In SSR analysis the average number of alleles, mean Shannon's information index, and the mean PIC values decreased in second generation as compared to first generation. The percentage of polymorphic loci decreased from 59.38% in M_1V_1 to 37.5% in M_1V_2 .

In conclusion, the above studies demonstrated that the mutation frequencies induced by acute gamma irradiation were higher than that of chronic gamma irradiation. In general, new variants of *C. alismatifolia* varieties developed in this study have the potential to be introduced to the Malaysian flower industry.

Abstrak tesis yang dikemukakan kepada Senate Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Doktor Falsafah

VARIASI FENOTIP DAN MOLEKUL ANTARA MUTAN *Curcuma alismatifolia* GAGNEP. TERPILIH HASIL DARIPADA SINARAN GAMMA AKUT DAN KRONIK

Oleh

SIMA TAHERI

November 2014

Pengerusi: Thohirah Lee Abdullah, PhD
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Kajian telah dijalankan untuk menilai variasi fenotip dan molekul antara mutan *Curcuma alismatifolia* terpilih melalui sinaran gamma akut dan kronik serta memperkenalkan varian *C. alismatifolia* yang baru. Dalam sinaran gamma akut, rizom dalam peringkat mata tunas bercambah telah diiradiasi dengan lapan dos berlainan (0, 10, 20, 25, 35, 40, 60, dan 100 Gy). Ujian kesensitifan radiasi telah menunjukkan bahawa LD₅₀ untuk varieti-varieti adalah seperti berikut: 'Chiang Mai Red': 21 Gy, 'Sweet Pink': 23 Gy, 'Kimono Pink': 25 Gy, 'Doi Tung 554': 28 Gy.

Dari analisis varians, variasi yang nyata telah diperhatikan untuk ciri-ciri vegetatif, perkembangan bunga dan ciri-ciri rizom antara keempat-empat kultivar *C. alismatifolia*, paras dos dan juga interaksi di antara dos × varieti. Dalam generasi pertama (M₁V₁), sebanyak 52 mutan klorofil telah diperhatikan, mencatatkan kekerapan min mutasi sebanyak 38.5% di mana 13.1 dan 19.3% adalah dihasilkan oleh dos 10 dan 20 Gy masing-masing. Mutan DT20-1 dengan daun pelindung yang mempunyai dua ton warna, adalah mutan yang paling menarik.

Dalam analisis SSR, keputusan yang diperoleh menunjukkan bahawa alokus SSR yang polimorfik adalah sangat berkesan dalam kajian genetic antara individu *C. alismatifolia* yang dirawat dan tidak dirawat. Individu yang diiradiasi dengan 20 Gy secara akut mencatatkan peratus min lokus polimorfik yang lebih tinggi (62.5%) berbanding dengan 10 Gy (59.38%) dan yang tidak dirawat (22%). Kesemua individu yang dirawat menghasilkan lengkung peleburan dengan suhu peleburan (T_m) yang berlainan berbanding dengan individu kawalan melalui analisis HRM.

Dalam sinaran gamma kronik, tumbuhan yang didedahkan kepada sinaran pada dos 14.6, 33, dan 87.4 Gy menunjukkan pengurangan yang nyata dalam ciri-ciri vegetative berbanding dengan kawalan. Yang menariknya, sinaran gamma padaparas dos yang rendah mampu merangsang pertumbuhan vegetative dan mengalakkan pembungaan awal tumbuhan mutan berbanding dengan tumbuhan kawalan. Namun begitu, kadar dos yang tinggi mengurangkan kapasiti berbunga dan kualiti rizom mutan *C. alismatifolia*. Dalam analisis SSR, variasi genetic keseluruhan untuk varieti-varieti yang dikaji menunjukkan bahawa sinaran gamma kronik, terutamanya

pada kadar dos yang tinggi (14.6, 33, dan 87.4 Gy) berkeupayaan untuk mengaruhkan lebih banyak variasi genetic pada genom individu *C. alismatifolia* yang dikaji.

Untuk mengkaji kestabilan morfologi dan genetic mutan-mutan yang dihasilkan daripada sinaran gamma akut dan kronik, pemerhatian serta kajian telah diteruskan sehingga generasi kedua (M_1V_2). Dalam sinaran gamma kronik, kekerapan mutasi klorofil telah berkurang dengan nyata pada individu M_1V_2 yang disinari dengan 10 dan 20 Gy. Individu DT20-1 dengan dua ton warna ungu pada generasi pertama menghasilkan mutan yang stabil dengan warna ungu yang lebih gelap pada daun pelindungnya dalam M_1V_2 . Untuk memperoleh mutan yang mempunyai kelainanan, keseragaman dan kestabilan (DUS), rizom-rizom dari kesemua individu telah ditanam dalam generasi M_1V_3 dan pemerhatian telah diteruskan. Sembilan individu dengan variasi warna daun pelindung dan kekerdilan telah dipilih untuk kajian selanjutnya dalam generasi M_1V_4 . Mutan DT20-1 yang mengekalkan warna ungu tua dalam M_1V_3 telah dianggap sebagai mutan yang berpotensi. Analisis SSR dalam generasi kedua menunjukkan bahawa peratusan lokus polimorfik telah berkurangan daripada 59.3 % kepada 40.6 % untuk individu yang diiradiasi dengan 10 Gy. Trend yang sama telah ditunjukkan untuk individu yang diiradiasi dengan 20 Gy, di mana peratusan lokus polimorfik berkurangan daripada 62.5 % kepada 50.0 %.

Dalam sinaran gamma kronik, tidakada variasi bentuk dan warna bunga yang diperhatikan dalam generasi pertama dipindahkan ke generasi seterusnya. Kesan yang diaruhkan oleh sinaran gamma kronik pada tumbuhan ini berkemungkinan besar merupakan perubahan fisiologi pada dos sublethal berbanding dengan kesan genetic tidak berbalik. Dalam analisis SSR purata nombor alel, min indeks maklumat Shannon dan min nilai PIC berkurangan pada generasi kedua berbanding dengan generasi pertama. Peratus lokus polimorfik berkurangan daripada 59.38% dalam M_1V_1 kepada 37.5% dalam M_1V_2 .

Kesimpulannya, kajian ini telah menunjukkan bahawa kekerapan mutasi yang diaruhkan oleh sinaran gamma akut adalah lebih tinggi daripada sinaran gamma kronik. Secara umumnya, varian baru *C. alismatifolia* yang dihasilkan daripada kajian ini mempunyai potensi untuk diperkenalkan ke dalam industry bunga di Malaysia.

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I certify that a Thesis Examination Committee has met on 3 November 2014 to conduct the final examination of Sima Taheri on her thesis entitled "Phenotypic and Molecular Variation among Selected *Curcuma alismatifolia* Gagnep. Mutants Derived from Acute and Chronic Gamma Irradiation" in accordance with the Universities and University Colleges Act 1971 and the Constitution of the Universiti Putra Malaysia [P.U.(A) 106] 15 March 1998. The Committee recommends that the student be awarded the Doctor of Philosophy.

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TABLE OF CONTENTS

	Page
ABSTRACT	i
ABSTRAK	iii
ACKNOWLEDGEMENTS	v
APPROVAL	vi
DECLARATION	viii
LIST OF TABLES	xvii
LIST OF FIGURES	xx
LIST OF ABBREVIATIONS	xxvi
CHAPTER	
1 INTRODUCTION	1
1.1 Floriculture industry in Malaysia	1
1.2 Overview of ornamental gingers	1
1.3 Status of <i>Curcuma alismatifolia</i>	2
1.4 The role of induced mutation in producing new cultivars of ornamental plants	2
1.5 The application of biotechnology in mutation programs	3
1.6 Research Objectives	4
2 LITERATURE REVIEW	5
2.1 Taxonomy of <i>C. alismatifolia</i>	5
2.2 Botanical characteristics of <i>C. alismatifolia</i>	5
2.3 Distribution and habitat of <i>C. alismatifolia</i>	6
2.4 Climatic condition of <i>C. alismatifolia</i>	6
2.5 Vegetative propagation of <i>C. alismatifolia</i>	6
2.6 Chromosome number of <i>C. alismatifolia</i>	7
2.7 Mutation	8
2.7.1 Induction of mutations	8
2.7.2 Mutation breeding of ornamental plants	10
2.7.3 Types of gamma irradiation	10
2.7.4 Radiation sensitivity	12
2.7.5 Lethal dose (LD ₅₀)	12
2.8 Using biotechnology techniques in mutation breeding programs	13
2.8.1 Application of molecular markers in mutation breeding programs	14
2.8.1.1 Using simple sequence repeats (SSR) for assessment of genetic variation	15
2.8.1.2 Using inter simple sequence repeats (ISSR) for assessment of genetic variation	16
2.8.2 Mutation scanning by high resolution melting technique (HRM)	16

3	RADIATION SENSITIVITY AND MORPHOLOGICAL STUDIES OF <i>CURCUMA ALISMATIFOLIA</i> CULTIVARS IN SELECTED DOSES OF ACUTE GAMMA RAYS IN M₁V₁, M₁V₂, and M₁V₃ GENERATIONS	20
3.1	Introduction	20
3.2	Material and methods	22
3.2.1	Plant materials	22
3.2.2	Planting media and preparation of rhizomes for acute gamma irradiation	23
3.2.3	Mutation induction by acute gamma rays	23
3.2.4	Radiation sensitivity test and determination of 50 % lethal dose (LD ₅₀)	24
3.2.5	Induction of mutation with selected doses of gamma radiation	24
3.2.6	Post irradiation care	24
3.2.7	Vegetative growth characteristics	24
3.2.7.1	Number of shoots	24
3.2.7.2	Leaf length	25
3.2.7.3	Leaf width	25
3.2.7.4	Number of leaves	25
3.2.7.5	Plant height	25
3.2.8	Flower development characteristics	25
3.2.8.1	Number of days to visible bud	25
3.2.8.2	Inflorescence length	25
3.2.8.3	Number of days to anthesis	25
3.2.8.4	Number of true flowers	25
3.2.8.5	Number of pink bracts	26
3.2.8.6	Number of days to senescence	26
3.2.9	Rhizome characteristics	26
3.2.9.1	Number of new rhizomes	26
3.2.9.2	Rhizome size	26
3.2.9.3	Storage root number	26
3.2.10	Chlorophyll mutation and leaf shape variation in M ₁ V ₁ generation	26
3.2.11	Flower shape mutations	27
3.2.12	Flower color mutations	27
3.2.13	Mutation frequency	27
3.2.14	Preparation of rhizomes for studies in M ₁ V ₂ generation	27
3.2.15	Data collection in M ₁ V ₂ generation	28
3.2.16	Morphological studies in M ₁ V ₃ generation	28
3.2.17	Experimental layout and statistical analysis	28
3.3	Results and discussions	29
3.3.1	Gamma irradiation and radiation sensitivity test	29
3.3.2	Analysis of variance (ANOVA) for morphological characteristics of <i>C. alismatifolia</i> in M ₁ V ₁ generation	32
3.3.3	Effect of acute gamma irradiation on vegetative growth characteristics of <i>C. alismatifolia</i> in M ₁ V ₁ generation	32
3.3.3.1	Number of shoots	33

3.3.3.2	Number of leaves	34
3.3.3.3	Leaf length	34
3.3.3.4	Leaf width	35
3.3.3.5	Plant height	36
3.3.4	Effect of acute gamma irradiation on flower development characteristics of <i>C. alismatifolia</i> in M ₁ V ₁ generation	37
3.3.4.1	Number of days to visible bud	38
3.3.4.2	Inflorescence length	39
3.3.4.3	Number of days to anthesis	40
3.3.4.4	Number of true flowers	41
3.3.4.5	Number of pink bracts	41
3.3.4.6	Days to senescence	42
3.3.5	Effect of acute gamma irradiation on rhizome characteristics of <i>C. alismatifolia</i> in M ₁ V ₁ generation	43
3.3.5.1	Number of new rhizomes	44
3.3.5.2	Rhizome size	45
3.3.5.3	Storage root number	46
3.3.6	Cluster analysis of <i>C. alismatifolia</i> individual plants for morphological characteristics in M ₁ V ₁ generation	47
3.3.7	Principal component analysis (PCA) for morphological traits in M ₁ V ₁ generation	48
3.3.8	Observed mutations and mutation frequencies in M ₁ V ₁ generation	51
3.3.8.1	Chlorophyll mutations	51
3.3.8.2	Morphological mutations	52
3.3.8.3	Flower color and shape mutations	53
3.3.9	Analysis of variance (ANOVA) for morphological traits of <i>C. alismatifolia</i> in M ₁ V ₂ generation	55
3.3.10	Effect of acute gamma rays on vegetative growth characteristics of <i>C. alismatifolia</i> in M ₁ V ₂ generation	55
3.3.10.1	Number of shoots	56
3.3.10.2	Number of leaves	56
3.3.10.3	Leaf length	56
3.3.10.4	Leaf width	57
3.3.10.5	Plant height	57
3.3.11	Effect of acute gamma irradiation on flower development characteristics in M ₁ V ₂ generation	57
3.3.12	Mutation frequencies in M ₁ V ₂	58
3.3.13	Correlation between morphological data in M ₁ V ₁ and M ₁ V ₂ generations	62
3.3.14	Morphological studies in M ₁ V ₃ generation	62
3.3.14.1	Performance of selected mutants in M ₁ V ₃ generation	62
3.4	Conclusion	64

4	ASSESSMENT OF GENETIC VARIATION OF ACUTELY IRRADIATED <i>CURCUMA ALISMATIFOLIA</i> CULTIVARS USING MOLECULAR MARKERS AND HIGH RESOLUTION MELTING TECHNIQUE	65
4.1	Introduction	65
4.2	Materials and methods	66
4.2.1	Plant materials	66
4.2.2	Planting media and preparation of rhizomes for acute gamma irradiation	67
4.2.3	Gamma irradiation of rhizomes of <i>C. alismatifolia</i>	67
4.2.4	SSR primer screening	67
4.2.4.1	DNA extraction	67
4.2.4.2	DNA purity and quantification	67
4.2.4.3	PCR amplification	67
4.2.4.4	PCR product visualization	68
4.2.5	Homogeneity test of the non-treated individual plants for SSR	68
4.2.6	SSR analysis	68
4.2.7	ISSR primer screening	69
4.2.7.1	PCR amplification	69
4.2.7.2	PCR product visualization	69
4.2.8	Data scoring and analysis for SSR and ISSR markers	69
4.2.9	High resolution melting (HRM) analysis	70
4.2.9.1	SSR primers	70
4.2.9.2	PCR amplification and HRM analysis	70
4.2.9.3	DNA sequencing	71
4.2.10	Molecular studies in M_1V_2 generation	71
4.2.10.1	DNA extraction	71
4.2.10.2	SSR analysis	71
4.3	Results and discussions	72
4.3.1	DNA extraction and qualification	72
4.3.2	SSR primer screening	72
4.3.3	Homogeneity test of the non-treated individual plants	74
4.3.4	Molecular studies in M_1V_1 generation	74
4.3.4.1	SSR polymorphism	74
4.3.4.2	Genetic relationship among treated and non-treated individual plants of <i>C. alismatifolia</i> using SSR molecular markers in M_1V_1 generation	78
4.3.4.3	Relationship among treated and non-treated individual plants of <i>C. alismatifolia</i> based on principle component analysis (PCA) using SSR molecular markers in M_1V_1 generation	79
4.3.4.4	ISSR polymorphism in M_1V_1 generation	81
4.3.4.5	Genetic relationship among treated and non-treated individual plants of <i>C. alismatifolia</i> using ISSR molecular markers in M_1V_1 generation	84
4.3.4.6	Comparison among SSR and ISSR markers in	88

	M ₁ V ₁ generation	
4.3.4.7	DNA polymorphism detection through HRM analysis in M ₁ V ₁ generation	89
4.3.5	Molecular studies in M ₁ V ₂ generation	93
4.3.5.1	SSR polymorphism and genetic variability analysis among irradiated <i>C. alismatifolia</i> individual plants in M ₁ V ₂ generation	93
4.3.5.2	Genetic relationship among treated and non-treated individual plants of <i>C. alismatifolia</i> using SSR molecular markers in M ₁ V ₂ generation	96
4.3.6	Correlations between molecular and morphological dissimilarity matrices in M ₁ V ₁ and M ₁ V ₂	98
4.4	Conclusion	98
5	MORPHOLOGICAL STUDIES OF IRRADIATED <i>CURCUMA ALISMATIFOLIA</i> CULTIVARS USING CHRONIC GAMMA IRRADIATION IN M₁V₁ AND M₁V₂ GENERATIONS	100
5.1	Introduction	100
5.2	Materials and methods	101
5.2.1	Plant materials	101
5.2.2	Planting and preparation of the rhizomes for chronic gamma irradiation	102
5.2.3	Mutation induction of <i>C. alismatifolia</i> by chronic gamma rays	102
5.2.4	Plant maintenance in gamma greenhouse (GGH) during gamma irradiation	104
5.2.5	Vegetative growth Characteristics	104
5.2.6	Flower development characteristics	104
5.2.7	Rhizome characteristics in selected doses	104
5.2.8	Chlorophyll mutation and leaf shape variation	104
5.2.9	Flower shape mutation	104
5.2.10	Flower color mutation	105
5.2.11	Mutation frequency	105
5.2.12	Morphological studies in M ₁ V ₂ generation	105
	5.2.12.1 Preparation of rhizomes for studies in M ₁ V ₂ generation	105
	5.2.12.2 Data collection in M ₁ V ₂ generation	105
5.2.13	Statistical analysis	105
5.3	Results and discussion	106
5.3.1	Radiation sensitivity and survival percentage of <i>C. alismatifolia</i> irradiated by chronic gamma irradiation	106
5.3.2	Effect of chronic gamma irradiation on vegetative growth characteristics in M ₁ V ₁ generation	106
	5.3.2.1 Number of shoots	108
	5.3.2.2 Number of leaves	108
	5.3.2.3 Leaf length	109
	5.3.2.4 Leaf width	110
	5.3.2.5 Plant height	110

5.3.3	Effect of chronic gamma irradiation on flower development characteristics in M_1V_1 generation	112
5.3.3.1	Number of days to invisible bud appearance	113
5.3.3.2	Inflorescence length	114
5.3.3.3	Number of days to anthesis	116
5.3.3.4	Number of true flowers	116
5.3.3.5	Number of pink bracts	117
5.3.3.6	Number of days to inflorescence senescence	118
5.3.4	Effect of chronic gamma irradiation on rhizome characteristics in M_1V_1 generation	119
5.3.4.1	Number of new rhizomes	119
5.3.4.2	Rhizome size	120
5.3.4.3	Number of storage roots	121
5.3.5	Cluster analysis of <i>C. alismatifolia</i> individual plants for 14 morphological traits in M_1V_1 generation	122
5.3.6	Principal component analysis of <i>C. alismatifolia</i> for morphological traits in M_1V_1 generation	124
5.3.7	Analysis of variance (ANOVA) for morphological traits of <i>C. alismatifolia</i> in M_1V_2 generation	125
5.3.7.1	Effect of chronic gamma irradiation on vegetative growth characteristics in M_1V_1 generation	125
5.3.8	Effect of chronic gamma irradiation on flower development characteristics in M_1V_2 generation	127
5.3.9	Mutations observed in the M_1V_1 and M_1V_2 generations	128
5.4	Conclusion	132
6	ASSESSMENT OF DNA VARIATIONS OF CHRONICALLY IRRADIATED <i>CURCUMA ALISMATIFOLIA</i> CULTIVARS USING SIMPLE SEQUENCE REPEAT MARKERS	134
6.1	Introduction	134
6.2	Materials and methods	135
6.2.1	Plant materials	135
6.2.2	Planting media and preparation of rhizomes for chronic gamma irradiation	135
6.2.3	Mutation induction of <i>C. alismatifolia</i> by chronic gamma rays	135
6.2.4	SSR analysis	135
6.2.4.1	DNA extraction	135
6.2.4.2	DNA purity and quantification	135
6.2.4.3	PCR amplification	135
6.2.4.4	PCR product visualization	136
6.2.4.5	Data scoring and analysis	136
6.2.5	Molecular studies in M_1V_2 generation	136
6.2.5.1	DNA extraction	136
6.2.5.2	SSR analysis	137
6.2.5.3	PCR product visualization to detect DNA polymorphism	137
6.3	Results and discussions	137
6.3.1	DNA extraction and qualification	137

6.3.2	Microsatellite polymorphism and genetic variability analysis among irradiated <i>C. alismatifolia</i> individual plants in M_1V_1 generation	137
6.3.3	Genetic relationship among treated and non-treated individual plants of <i>C. alismatifolia</i> using SSR molecular markers in M_1V_1 generation	142
6.3.4	Relationship among treated and non-treated individual plants of <i>C. alismatifolia</i> based on principle component analysis (PCA) using SSR molecular markers in M_1V_1	114
6.3.5	SSR polymorphism and genetic variability analysis among irradiated <i>C. alismatifolia</i> individual plants in M_1V_2 generation	145
6.3.6	Correlation between molecular and morphological dissimilarity matrices in M_1V_1 and M_1V_2 generations	147
6.3.7	Comparison between acute and chronic gamma irradiation	148
6.4	Conclusion	148
7	CONCLUSION AND RECOMMENDATIONS FOR FUTURE STUDIES	149
	REFERENCES	152
	APPENDICES	177
	BIODATA OF STUDENT	196
	LIST OF PUBLICATIONS	197

LIST OF TABLES

Table		Page
2.1	Examples of mutation breeding program and the application of molecular markers on selected crops	9
3.1	Distinctive qualitative features of <i>C. alismatifolia</i> cultivars and hybrid used in this study. The leaf and flower color were identified by RHS color chart	22
3.2	The number of irradiated and mortal rhizomes of <i>C. alismatifolia</i> cultivars after acute irradiation with different doses of gamma rays	29
3.3	Effect of acute gamma rays on vegetative growth characteristics of <i>C. alismatifolia</i> in M_1V_1 generation	32
3.4	Effect of acute gamma rays on flower development characteristics of <i>C. alismatifolia</i> in M_1V_1 generation	38
3.5	Effect of acute gamma rays on rhizome characteristics of <i>C. alismatifolia</i> in M_1V_1 generation	44
3.6	Cluster means for 14 traits estimated in 44 individuals of <i>C. alismatifolia</i> in M_1V_1 generation	48
3.7	Eigenvectors, Eigenvalues, and proportions of variability for three principle components among 14 morphological traits of 44 <i>C. alismatifolia</i> in M_1V_1 generation	50
3.8	Different types of observed mutations in four studied cultivars of <i>C. alismatifolia</i> in the M_1V_1 generation	52
3.9	Effect of acute gamma rays on vegetative growth characteristics of <i>C. alismatifolia</i> in M_1V_2 generation	56
3.10	Effect of acute gamma rays on flower development characteristics of <i>C. alismatifolia</i> in M_1V_2 generation	58
3.11	Mutation frequency of Chlorophyll mutants, morphological mutants and flower color and shape mutants of <i>C. alismatifolia</i> individual plants in M_1V_1 and M_1V_2 generations	59
3.12	Performance of selected mutants derived from gamma irradiation in M_1V_3 generation for vegetative growth characteristics	64
4.1	SSR polymorphic primer sequences used for analysis of 44 individuals of <i>C. alismatifolia</i> and their respective annealing temperature (Ta)	73

4.2	ISSR polymorphic primer sequences used for analysis of 44 individuals of <i>C. alismatifolia</i> and their respective annealing temperature (Ta)	73
4.3	Characterization of SSR loci in M ₁ V ₁ generation of <i>C. alismatifolia</i> individual plants irradiated at 0, 10, and 20 Gy	76
4.4	Genetic variation estimates for <i>C. alismatifolia</i> individual plants at 0, 10, and 20 Gy using SSR markers	77
4.5	The principal component of SSR marker data on treated and non-treated individual plants of <i>C. alismatifolia</i>	80
4.6	Characterization of ISSR loci in M ₁ V ₁ generation of <i>C. alismatifolia</i> individual plants irradiated at 0, 10, and 20 Gy	82
4.7	Genetic variation estimates for <i>C. alismatifolia</i> individual plants at 0, 10, and 20 Gy using ISSR markers	84
4.8	Characterization of SSR loci in M ₁ V ₂ generation of <i>C. alismatifolia</i> individual plants irradiated at 0, 10, and 20 Gy	94
4.9	Comparison between SSR loci characteristics in M ₁ V ₁ and M ₁ V ₂ generations	95
4.10	Percentage of polymorphic loci (Ppl%) of varieties of <i>C. alismatifolia</i> in M ₁ V ₁ and M ₁ V ₂ generations	95
5.1	The selected dose rates, ring numbers, and accumulative dose of chronic gamma rays	102
5.2	Effect of chronic gamma rays on vegetative growth characteristics of <i>C. alismatifolia</i> in M ₁ V ₁ generation	107
5.3	Effect of chronic gamma rays on flower development characteristics of <i>C. alismatifolia</i> in M ₁ V ₁ generation	113
5.4	Effect of chronic gamma rays on rhizome characteristics of <i>C. alismatifolia</i> in M ₁ V ₁ generation	119
5.5	Cluster means for 14 characters estimated in 104 individuals of <i>C. alismatifolia</i>	124
5.6	Eigenvectors, Eigenvalues, and proportions of variability for four principle components among 14 characters for 104 <i>C. alismatifolia</i>	125
5.7	Effect of chronic gamma rays on vegetative growth characteristics of <i>C. alismatifolia</i> in M ₁ V ₂ generation	126

5.8	Effect of chronic gamma rays on flower development characteristics of <i>C. alismatifolia</i> in M_1V_2 generation	128
5.9	Mutation frequency of observed mutations induced by chronic gamma irradiation in M_1V_1 generation	132
6.1	Characterization of SSR loci in M_1V_1 generation of 104 chronically irradiated <i>C. alismatifolia</i> individual plants	138
6.2	Genetic variation estimates for <i>C. alismatifolia</i> individual plants at six doses using SSR markers in M_1V_1 generation	141
6.3	The principal component of SSR marker data on treated and non-treated individual plants of <i>C. alismatifolia</i>	144
6.4	Characterization of SSR loci in M_1V_2 generation of 104 chronically irradiated <i>C. alismatifolia</i> individual plants	146
6.5	Comparison between SSR loci characteristics in M_1V_1 and M_1V_2 generations	146
6.6	Percentage of polymorphic loci (Pp1%) of varieties of <i>C. alismatifolia</i> in M_1V_1 and M_1V_2 generations	147

LIST OF FIGURES

Figure		Page
2.1	Development stages of <i>C. alismatifolia</i> . (A) Planting organ is a rhizome with storage roots (ball shape). (B) First shoot development. (C) Flowering. (D) Fully open inflorescence. (E) Inflorescence senescence. (F) Plants at harvest stage – New (light brown) and old (dark brown) storage organs and rhizomes	7
3.1	<i>C. alismatifolia</i> cultivars and one hybrid. (A) ‘Chiang Mai Red’; (B) ‘Doi Tung 554’ (hybrid); (C) ‘Sweet Pink’; (D) ‘Kimono Pink’	22
3.2	Rhizome preparation and gamma irradiation of rhizomes of <i>C. alismatifolia</i> . (A) Rhizomes before irradiation; (B) Planting rhizomes in media; (C) Sprouted rhizomes; (D) preparation of rhizomes for gamma irradiation; (E) Gamma cell used for acute gamma irradiation	23
3.3	PoloPlus plot of linear scale of dose versus mortality percent, (A) ‘Doi Tung 554’, (B) ‘Chiang Mai Red’, (C) ‘Sweet Pink’, (D) ‘Kimono Pink’	31
3.4	Effects of acute gamma irradiation on number of shoots of <i>C. alismatifolia</i> at 0, 10, and 20 Gy doses in M_1V_1 generation. Bars from left to right at each dose: ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, ‘Kimono Pink’	33
3.5	Effect of acute gamma irradiation on number of leaves of <i>C. alismatifolia</i> at 0, 10, and 20 Gy doses in M_1V_1 generation. Bars from left to right at each dose: ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, ‘Kimono Pink’	34
3.6	Effect of acute gamma irradiation on leaf length of <i>C. alismatifolia</i> varieties at 0, 10, and 20 Gy doses in M_1V_1 generation. Bars from left to right at each dose: ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, ‘Kimono Pink’	35
3.7	Effect of acute gamma irradiation on leaf width of <i>C. alismatifolia</i> cultivars at 0, 10, and 20 Gy doses in M_1V_1 generation. Bars from left to right at each dose: ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, ‘Kimono Pink’	36
3.8	Effects of acute gamma irradiation on plant height of <i>C. alismatifolia</i> cultivars at 0, 10, and 20 Gy doses in M_1V_1 generation. Bars from left to right at each dose: ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, ‘Kimono Pink’	36
3.9	Effect of gamma irradiation on plant height of ‘Doi Tung 554’	37

3.10	Effects of acute gamma irradiation on number of days to visible bud appearance of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	39
3.11	Effects of acute gamma irradiation on inflorescence length of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	40
3.12	Effects of acute gamma irradiation on number of days to anthesis of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	40
3.13	Effects of acute gamma irradiation on number of true flowers of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	41
3.14	Effects of acute gamma irradiation on number of pink bracts of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	42
3.15	Effects of acute gamma irradiation on number of days to senescence of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	43
3.16	Effects of acute gamma irradiation on number of new rhizomes of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	45
3.17	Effects of acute gamma irradiation on rhizome size of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	45
3.18	Effects of acute gamma irradiation on number of storage roots of <i>C. alismatifolia</i> in M ₁ V ₁ generation. Bars from left to right: ‘Chiang Mai Red’, ‘Doi Tung554’, ‘Sweet Pink’, ‘Kimono Pink’	46
3.19	Effect of acute gamma irradiation on rhizome size and number of storage roots of at 0, 10, and 20 Gy doses in M ₁ V ₁ generation. (A): ‘Doi Tung554’, (B): ‘Sweet Pink’	47
3.20	Dendrogram representing the morphological variation among 44 treated and non-treated individuals of <i>C. alismatifolia</i> across 14 variables in M ₁ V ₁ generation. CMR: ‘Chiang Mai Red’; DT: ‘Doi Tung554’; SP: ‘Sweet Pink’; KP: ‘Kimono Pink’. SP20-1= 20 Gy, individual no. 1	49
3.21	Three dimensional graph of principal component analysis (PCA) for 14 morphological variables indicating relationships among	51

	treated and non-treated of four varieties of <i>C. alismatifolia</i> in M_1V_1 generation	
3.22	Chlorophyll and morphological mutations observed in 10 and 20 Gy irradiated <i>C. alismatifolia</i> mutants. DT: ‘Doi Tung 554’; SP: ‘Sweet Pink’	53
3.23	Effect of acute gamma rays on flower shape and color of <i>C. alismatifolia</i> in M_1V_1 ; (A) non-treated inflorescence of ‘Chiang Mai Red’, (B) Inflorescence without true flowers cultivar CMR (10 Gy), (C) twin inflorescence within one stalk cultivar CMR (10 Gy), (D) absence of upper bracts of CMR (10 Gy), (E) non-treated true flower with one flag petal, (F) Two- flag petals true flowers (10 Gy), (G) non-treated inflorescence of ‘Doi Tung’, (H), (I) two tone-purple color of bracts of ‘Doi Tung’ (20 Gy), (J) Marbled pattern of bracts of ‘Doi Tung’ (20 Gy), (K) Double flower stalk, hybrid ‘Doi Tung’ (20 Gy)	54
3.24	Effect of gamma irradiation on plant height and flowering of <i>C. alismatifolia</i> individuals in M_1V_2 (A) Dwarfism in ‘Kimono Pink’ cultivar, (B) Flowering of 10 Gy ‘Chiang Mai Red’ individuals in M_1V_1 and M_1V_2	61
3.25	Variation in bract color of 20 Gy irradiated hybrid ‘Doi Tung 554’ (DT20-1) compared to control in M_1V_1 and M_1V_2 generations	62
3.26	Bract color mutant of DT20-1 in M_1V_3 generation	63
4.1	Homogeneity of four <i>C. alismatifolia</i> cultivars/hybrid before irradiation. (A) SSR marker, (B) ISSR marker. DT: ‘Doi Tung’; CMR: ‘Chiang Mai Red’; SP: ‘Sweet Pink’; KP: ‘Kimono Pink’	74
4.2	Dendrogram based on Dice’s similarity coefficient indicating relationships among treated (10 and 20 Gy) of <i>C. alismatifolia</i> compared to their non-treated plants (0 Gy)	79
4.3	Three-dimensional plot of principal components of SSR marker data among treated and non-treated individual plants of <i>C. alismatifolia</i>	81
4.4	Dendrogram based on Jaccard’s similarity coefficient indicating relationships among treated (10 and 20 Gy) of <i>C. alismatifolia</i> compared to their non-treated plants (0 Gy) using ISSR markers	86
4.5	Melting curves of treated and non-treated individuals of <i>C. alismatifolia</i> for primer ‘clon01’. Plot A represents melting curves, SNPs, and different melting temperature for control, 10, and 20 Gy Sweet pink, plot B represents normalized data for 10 and 20 Gy irradiated individuals of ‘Sweet Pink’ cultivar	91
4.6	Clustering of treated and non-treated individuals of <i>C.</i>	92

alismatifolia for clon01. The individuals inside the circles are 10 and 20 Gy mutants. A) ‘Doi Tung554’, B) ‘Chiang Mai Red’, C) ‘Kimono Pink’, D) ‘Sweet Pink’

- 4.7 Alignment of SP10-1 with non-treated ‘Sweet Pink’ (primer ‘clon01’). The sequence shown represents deletions (Adenine and Thymine), Insertions (Guanine), and substitutions (Tymine with Adenine) occurred in 10 Gy ‘Sweet Pink’ 93
- 4.8 Dendrogram based on Dice’s similarity coefficient indicating relationships among treated (10 and 20 Gy) of *C. alismatifolia* compared to their non-treated plants (0 Gy) using SSR markers in M_1V_2 generation 97
- 5.1 (A) Gamma greenhouse (GGH) in Nuclear Malaysia, (B) Gamma rays source (^{137}Cs), (C) The position of rings to gamma source, (D) arrangement of plants in GGH 103
- 5.2 Effect of chronic gamma irradiation on number of new shoots of *C. alismatifolia* cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’ 108
- 5.3 Effect of chronic gamma irradiation on leaf number of *C. alismatifolia* cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’ 109
- 5.4 Effect of chronic gamma irradiation on leaf length of *C. alismatifolia* cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’ 109
- 5.5 Effect of chronic gamma irradiation on leaf width of *C. alismatifolia* cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’ 110
- 5.6 Effect of chronic gamma irradiation on plant height of *C. alismatifolia* cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’ 111
- 5.7 Effect of chronic gamma irradiation on number of days to visible bud appearance of *C. alismatifolia* cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’ 114
- 5.8 Effect of chronic gamma irradiation on inflorescence length of *C. alismatifolia* cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and 115

	‘Kimono Pink’	
5.9	The effect of different doses of chronic gamma irradiation on inflorescence size of <i>C. alismatifolia</i> hybrid ‘Doi Tung 554’	115
5.10	Effect of chronic gamma irradiation on number of days to anthesis of <i>C. alismatifolia</i> cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’	116
5.11	Effect of chronic gamma irradiation on number of true flowers of <i>C. alismatifolia</i> cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’	117
5.12	Effect of chronic gamma irradiation on number of pink bracts of <i>C. alismatifolia</i> cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’	118
5.13	Effect of chronic gamma irradiation on number of days to senescence of <i>C. alismatifolia</i> cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung’, ‘Sweet Pink’, and ‘Kimono Pink’	118
5.14	Effect of chronic gamma irradiation on number of new rhizomes of <i>C. alismatifolia</i> cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’	120
5.15	Effect of chronic gamma irradiation on rhizome size of <i>C. alismatifolia</i> cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’	121
5.16	Effect of chronic gamma irradiation on number of storage roots of <i>C. alismatifolia</i> cultivars in M_1V_1 generation. From left to right at each dose, ‘Chiang Mai Red’, ‘Doi Tung 554’, ‘Sweet Pink’, and ‘Kimono Pink’	122
5.17	Dendrogram representing the morphological variation among 104 treated and non-treated individuals of <i>C. alismatifolia</i> across 14 variables. CMR: ‘Chiang Mai Red’; DT: ‘Doi Tung 554’; SP: ‘Sweet Pink’; Kp: ‘Kimono Pink’. e.g. CMR11-1= “11” number is the R11 (6.25 Gy) and “1” number is individual number	123
5.18	Chlorophyll mutation and morphological mutations in chronic gamma irradiated <i>C. alismatifolia</i> individual plants in (A) M_1V_1 generation and (B) M_1V_2 generation	129
5.19	Induced flower color and shape variations by chronic gamma	131

irradiation

- 6.1 Dendrogram based on Dice's similarity coefficient indicating relationships among treated *C. alismatifolia* compared to their non-treated plants (0 Gy) in M_1V_1 generation 143
- 6.2 Three dimensional graph of principal component analysis (PCA) indicating relationships among treated and non-treated of 104 individuals of *C. alismatifolia* 145



LIST OF ABBREVIATIONS

°C	degree Celsius
µl	microliter
µM	micromole
¹³⁷ Cs	caesium-137
⁶⁰ Co	Cobalt-60
AFLP	amplified fragment length polymorphism
ANOVA	analysis of variance
C.V.	coefficient of variation
CE	capillary electrophoresis
CMR	Chiang Mai Red
CRD	Complete randomized design
CTAB	Cetyltrimethylammonium bromide
DNA	deoxyribonucleic acid
DT	Doi Tung 554
DUS	distinctness, uniformity and stability
dNTP	2'-deoxynucleotide 5'-triphosphate
EDTA	Ethylenediaminetetraacetic acid
EMS	ethyl-methane-sulphonate
ENH	ethyl-nitrosourea
FAO	Food and agriculture organization
G1	Gap 1
G2	Gap 2
GGH	Gamma greenhouse
Gy	Gray
H	Nei's (1973) gene diversity
h	hours
He	Expected heterozygosity
H _o	observed heterozygosity
HRM	high resolution melting
I	Shannon's information index
IAEA	International Atomic Energy Agency
IR	ionizing radiations
ISSR	inter-simple sequence repeat
kR	Kilo radian
KP	Kimono Pink
LD50	50 % lethal dose
LSD	Least significant difference
M	Mitosis
M ₁ V ₁	First vegetative generation of irradiated populations
M ₁ V ₂	Second vegetative generation of irradiated populations
M ₁ V ₃	Third vegetative generation of irradiated populations
MAB	Mutation-assisted breeding
MNH	methyl-nitroso-urea
mg	miligram
min	Minutes
mM	milimolar
n _a	Observed number of alleles
n _e	Effective number of alleles

NTSYS	numerical taxonomy multivariate analysis system
PCA	principal component analysis
PCR	polymerase chain reaction
PIC	polymorphic information content
PVP	polyvinyl pyrrolidone
RAPD	random amplified polymorphic DNA
RCBD	randomized complete block design
RFLP	restriction fragment length polymorphism
ROS	reactive oxygen species
S	Synthesis
SA	sodium azide
SAS	statistical analysis system
SNP	single nucleotide polymorphism
SSR	simple sequence repeats
SP	Sweet Pink
TBE	Tris/Borate/EDTA
TE	Tris EDTA buffer
UPGMA	Unweighted pair group method using arithmetic averages
UV	ultraviolet

CHAPTER 1

INTRODUCTION

1.1 Floriculture industry in Malaysia

The Malaysian cut flower is a relatively new industry as compared to other agricultural business but have grown from its rather low position into a commercial yield in the mid-eighties. Efficient increase in the flower production and export in response to local and foreign demands are witnesses of such tremendous growth in the last decade. It is expected this developing trend will continue in the future along with growing affluence of the local population and that of the developed countries as well as improved market opportunities. In the Tenth Malaysia Plan (2011-2015), cut flowers with qualified potential to meet the growing local and foreign demands and to generate higher income for producers are preferred. In 2006, exports from Malaysia for temperate flowers, orchids, foliage and dried flowers were RM100 million a figure that only represented about 33 % of the world market (Ahmad *et al.*, 2007). This scenario showed that the Malaysian floriculture industry still has the potential for expansion to bring more revenue to the country as well as the industry.

In recent years, the Malaysian floriculture industry has shown a remarkable growth promoted by urbanization and landscaping activities and has become a dynamic industry in the agricultural export sector. Thus, as a contribution of the Tenth Malaysian Plan, which emphasized more on the agriculture sector, the introduction of a new flowering crop is vital as it has vast potential in contributing to Malaysia's economic development. In the quest for new variety of ornamental plants to further enhance the lucrative floriculture industry, ornamental Zingibers, which have long been known of their showy inflorescences, present the highest potential (Lee, 2007).

1.2 Overview of ornamental gingers

The Zingiberaceae or ginger family is a family of flowering plants comprising about 90 genera and approximately 1400 species which continues to grow as more new species are being identified (Chapman, 1995). The genera of flowering gingers include *Alpinia*, *Curcuma*, *Etingera*, *Globba*, *Hedychium*, *Kaempferia* and *Zingiber*. Ornamental gingers with spectacular and brightly colored flowers are amazingly versatile group of plants with a wide range of shapes, size, and colors. These gingers are cultivated in many parts of the tropics and subtropics as landscape plants, pot flowers or cut flowers. The genus *Curcuma* is considered to have originated in the Indo-Malayan Region (Purseglove, 1972), has a widespread occurrence in the tropics of Asia to Africa and Australia. *Curcuma* is the genera in the Zingiberaceae family that have recently gained much attraction.

1.3 Status of *C. alismatifolia*

C. alismatifolia, native to northern Thailand was introduced to the world flower market in the early 1990s. The basic physiology of flowering for *C. alismatifolia* has been intensively studied (Fukai and Udomdee, 2005; Azuma and Takano, 1994; Hajiladi *et al.*, 1997a, Kuehny *et al.*, 2002) and year-round cut flower production has been established. Presently, the species is grown predominantly in Thailand and China where they are the native (indigenous) to the South East Asia (Paz, 2003). Although successful breeding programs of *Curcuma* such as production of inter-specific hybrids have been achieved in Thailand (Wongpiyasatid *et al.*, 2009) only a few have been introduced to the world flower market because of the lack of commercial bodies to lead these flowers onto the world market. Similar growing conditions here in Malaysia offer a great potential and this opportunity is presently unexploited.

This plant has a high potential to enhance the floriculture industry in Malaysia as it is becoming a favorite in the international market – as cut flower production, potted plant as well as for exterior landscapes. It is also becoming a highly demanded plant in Japan, Holland, USA, Taiwan and Israel where it is grown commercially as a cut flower.

Curcuma alismatifolia have colorful, long-lasting inflorescence, with a 180-200 days production cycle and with few pest problems. They are herbaceous perennials with short fleshy rhizomes and tuberous roots, often with a dormancy period (Khuankaew *et al.*, 2010). In Thailand, rhizomes of *Curcuma* in commercial production are harvested during the three months dormant period of storage and distribution (Lee, 2007). Other characteristics that make them attractive to the floricultural industry are ease of production, unique foliage, and numerous flowering stems per pot. In this study, three cultivars of *C. alismatifolia*, ‘Chiang Mai Red’, ‘Sweet Pink’, ‘Kimono Pink’, and one hybrid namely ‘Doi Tung 554’ were chosen due to their free flowering habits and attractive flower colour and shape.

1.4 The role of induced mutation in producing new cultivars of ornamental plants

Mutation breeding is one of the methods for generating genetic variation and obtaining new cultivars of ornamental plants during the past decades (Ahloowalia and Maluszynski, 2001; Broertjes *et al.*, 1976; Zalewska and Jerzy, 1997). Induced mutations have been utilized for the past 70 years to produce mutant cultivars by changing the plant characteristic for a significant increase in plant production among both seed and vegetatively propagated crops. The most common mutagens used are: chemical (alkylating agents)-ethyl-nitrosourea (ENH), methyl-nitroso-urea (MNH), sodium azide (SA), ethyl-methane-sulphonate (EMS), and physical agents–X-rays, gamma rays, fast neutrons, ultra violet, and laser (Jain, 2002). Among these mutagens, gamma rays and EMS are widely used for mutation induction. The advantages of physical mutagens are accurate dosimetry and reasonable reproducibility, as well as high and uniform penetration of multicellular system particularly by gamma rays. Extensive work has been carried out for the last 30 years for improvement of different ornamental crops. Gamma-rays have been most successfully used and 76 new mutant varieties with

changed flower color/shape, and chlorophyll variation in leaves have been developed and released in different ornamentals (Datta, 2009).

1.5 The application of biotechnology in mutation programs

The estimation of genetic variation on the basis of morphological traits alone, which are the product of gene and environmental interactions, does not determine the actual level of genetic variation among studied individuals (Sigrist *et al.*, 2011). In order to reveal the induced variation in genomic DNA of mutants, appropriate molecular markers are required. Among different classes of molecular markers, Simple sequence repeats (SSR) are useful for a variety of applications in plant genetics and breeding. SSR also known as microsatellites are a class of molecular markers based on tandem repeats of short (1-6 bp) DNA sequences which are ubiquitously distributed throughout eukaryotic genomes. These repeat sequences with codominant inheritance are found to be abundant in plant genomes and are frequently highly polymorphic, even among closely related cultivars. Variation could be due to insertion or deletion of nucleotides. In addition, SSR markers have also been shown to have transferability across different species and plants, a fact that increase their value in plant genetic studies. Inter-simple sequence repeat (ISSR) involves amplification of DNA segments present at an amplification distance in between two identical SSR repeat regions oriented in opposite directions (Masumbuko and Bryngelsson, 2006). Based on the presence of the SSR throughout genome, ISSR analysis involves polymerase chain reaction (PCR) amplification of genomic DNA using a single primer that targets the repeat, which is anchored at 3' or 5' end. Unlike SSR markers, ISSR does not require sequence information or prior genetic studies for primer synthesis (Joshi *et al.*, 2000). Besides ISSRs have been demonstrated to provide highly reproducible results and generate abundant polymorphism in a number of crop species. Therefore it provides an efficient tool for assessment of DNA variation induced by gamma irradiation.

SSRs and ISSRs have been highly popular genetic markers for last two decades because of the above mentioned benefits. However, the traditional protocols used for SSR, employ loci-specific primers to PCR amplify the DNA fragment containing nucleotide repeats, and separate the PCR products using laborious electrophoresis. Nevertheless, high resolution melt (HRM) analysis is a technique that measures the disassociation of double-stranded DNA at high temperature resolution, and permits the analysis of genetic variations (SNPs, mutations, methylation) in PCR amplicons (Gundry *et al.*, 2003). This technique allows mutation scanning without the need for costly labeled probes, as it uses high fidelity heteroduplex- detecting double-stranded DNA binding dyes, such as EvaGreen which exhibit equal binding affinities for GC-rich and AT-rich regions and no sequence preference (Lochlainn *et al.*, 2011).

However, due to the lack of new local varieties, most of the varieties planted in Malaysia were imported from Thailand. In addition, unlike other well-established bulbous floral crops, studies on *C. alismatifolia* for induced mutation have not been fully investigated. The development of new varieties of *Curcuma* is expected to reduce the grower's dependence on Thailand's cultivars in the long term.

The present study is aimed to use acute and chronic gamma rays to induce morphological and DNA variations in *C. alismatifolia* and to develop new variants of *C. alismatifolia*, as well as to compare the effectiveness of acute and chronic gamma irradiation in inducing desired morphological and DNA variation.

1.6 Research objectives

This study was designed to achieve the following general objectives:

1. To induce morphological and DNA variations in *C. alismatifolia* cultivars through acute and chronic gamma irradiation.
2. To develop new variants of *C. alismatifolia* using acute and chronic gamma irradiation.
3. To compare the effectiveness of acute and chronic gamma irradiation in induction of desired morphological and DNA variations in *C. alismatifolia*.

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