



UNIVERSITI PUTRA MALAYSIA

***THE FUNCTIONAL STUDY OF ORYZACYSTATIN-1 PROTEASE
INHIBITOR UNDER SALINITY STRESS IN MALAYSIAN RICE
CULTIVAR***

NUR HAZIQAH BINTI MOHAMAD

FBSB 2015 67

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(Professor Dato' Dr. Abu Bakar Salleh)

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LIST OF ABBREVIATION

μL	Microlitre
BLAST	Basic Local Alignment Search Tools
Bp	Base pair
cDNA	Complement Deoxyribonucleic Acid
Cl^+	Calcium ion
dH ₂ O	Distilled water
DNA	Deoxyribonucleic Acid
DNTP	Deoxynucleotide triphosphate
G	Gram
Hr	Hour
MgCl ₂	Magnesium Chloride
mL	Mililitre
MR	Malaysia Rice
mRNA	Messenger Ribonucleic Acid
Na^+	Sodium ion
NaCl	Sadium Hydroxide
OC-1	Oryzacystatin-1
PCR	Polymerase Chain Reaction
qPCR	Real-time PCR
RNA	Ribonucleic Acid
RT-PCR	Reverse Transcriptase Polymerase Chain Reaction
TAE	Tris-Acetate EDTA
T _m	Melting Temperature
UBQ5	Ubiquitin5

ABSTRACT

Phytocystatin is a protein that encodes cysteine proteinase inhibitor. It is well known for its antipathogenic properties and responsible in plant stress tolerance. Proline is an amino acid that acts as osmoprotectant in plant. Previously, both phytocystatin and proline show response in plant abiotic stresses and have become a potential strategy in combating abiotic stress. In this study, we reported the isolation and characterization of Oryzacystatin-1 (*OC-1*) gene from Malaysian cultivar of *Oryza sativa* L., the gene expression profiling by using RT-PCR and the accumulation of proline in stress-induced rice plantlets of MR211, MR220 and MR219. Three Malaysian rice (*Oryza sativa* L.) cultivars, MR 211, MR 220 and MR 219 were tested in pot culture under different salinity concentration (0, 4, 8, 12 dS m⁻¹) and five different duration of salinity treatment (0, 3, 6, 9, 24 hr) for molecular and biochemical response. All the tests were conducted in randomized complete block design with three replicates in a culture room. The result showed that, the *OC-1* gene expression level and proline content were influenced by salinity stress. A cDNA fragment encoding phytocystatin, *OC-1* was also isolated. Sequence analysis shows that *OC-1* contain all common motifs found in phytocystatin such as a single G residue, QVVAG sequence and PW dipeptide motif. The presence of several cystatin superfamily was proven through Protein BLAST confirming the *OC-1* gene isolation earlier. Then, phylogeny studies have shown that the *OC-1* gene is closely related to phytocystatin from *Zea mays* and *Coix lacryma-jobi*.

ABSTRAK

Phytocystatin adalah protein yang mengekod perencat proteinase sisteina. Phytocystatin terkenal dengan sifat antipatogen dan terlibat pada toleransi pokok terhadap tekanan. Prolin merupakan asid amino yang bertindak sebagai pelindung osmotik di dalam pokok. Sebelum ini, kedua-dua phytocystatin dan prolin telah menunjukkan tindak balas terhadap tekanan abiotik di dalam pokok dan telah menjadi satu strategi yang berpotensi dalam memerangi tekanan abiotik. Di dalam kajian ini, kami melaporkan pengasingan dan pencirian *Oryzacystatin-1* gen daripada *Oryza sativa* L. bagi kultivar Malaysia, profil ekspresi gen dengan menggunakan RT-PCR serta pengumpulan prolin didalam pokok padi MR 211, MR 220 dan MR 219 yang telah diberikan tekanan kemasinan. Tiga kultivar padi Malaysia telah diuji didalam bekas kultur dengan menggunakan tahap kemasinan yang berbeza (0, 4, 8, 12 dS m⁻¹) serta lima kadar rawatan yang berbeza (0, 3, 6, 9, 24 jam) bagi tujuan tindak balas molekular dan biokimia. Kesemua ujian tersebut telah dijalankan secara rawak didalam reka bentuk lengkap dengan tiga replikasi didalam bilik kultur. Hasilnya, tingkatan ekspresi *Oryzacystatin-1* gen dan kandungan prolin didalam pokok di pengaruhi oleh tekanan kemasinan. Analisa jujukan menunjukkan *OC-1* mempunyai kesemua motif yang biasa dijumpai pada phytocystatin, seperti G residu, urutan QVVAG dan motif dipeptid PW. Hadirnya beberapa keluarga cystatin yang dibuktikan melalui BLAST protein, mengesahkan gen yang telah di asingkan adalah *OC-1*. Seterusnya, kajian filogeni telah menunjukkan *OC-1* gen berkait rapat dengan phytocystatin daripada *Zea mays* dan *Coix lacryma-jobi*.

CHAPTER 1

INTRODUCTION

Rice (*Oryza sativa* L.) is the third top ranking crop planted in Malaysia that covers about 205,548 hectares area in Peninsular Malaysia (*Buku Perangkaan Pertanian*, 2007). The Malaysian Agriculture and Agro-based Industry Ministry, reported to target Malaysia to achieve a 100% self-sufficiency level (SSL) in paddy production by 2020 (“Self Sufficiency in Rice”, 2015). To fulfill the current and future increasing demands, Malaysia needs to expand its rice area. It is predicted that in 2056, salinity problems would affect 100000 hectares of rice area. The continuous of salinity problem may result in reduction of rice area which would generate food shortage in domestic and global market (Selamat and Ismail, 2009).

Salinity is the biggest problem in rice production at all growth stages where it creates the osmotic effect and ion toxicity (Hakim *et al.*, 2014). High salt concentration in the soil will lower the soil water potential and cause water to move from the plant root to the soil. While, high accumulation of Na^+ and Cl^- inside the cytoplasm of plant will inhibit many enzymatic processes (Jampeetong and Brix, 2009). In order to deal with the unfavorable growth condition, plant develops several mechanisms of biochemical and molecular adaptation by preserving the basic metabolic activities (Basu and Roycoudhury, 2014).

In recent years, several biotechnological approaches mainly in over-expression of phytocystatins gene have been successful studied for plant improvement. For instant, phytocystatin has been used to prevent from insect feeding in transformed plants (Christou *et al.*, 2006; Kiggundu *et al.*, 2010) and used to improve the yield of bio-engineered proteins such as vaccines and metabolic

enzymes (Pillay *et al.*, 2012; Rivard *et al.*, 2006). Cysteine protease (CP) is the main target of phytocystatins by controlling CP activities which regulated by developmental and environmental cues (Quain *et al.*, 2014). Interestingly, studies have suggested that plant cystatin are also involved in responding to abiotic environment stress as they are highly expressed during harsh condition such as cold, drought, oxidant stress, wounds and salinity stress (Benchabane *et al.*, 2010; Bangrak and Chotigeat, 2011; Pereira *et al.*, 2011). However, the information of phytocystatin on Malaysian rice is very limited.

On the other hand, proline accumulation is a biochemical response in many plant species incited by salt stress. Proline as an osmoprotectant will preserve the osmotic stability and prevent damage in plant. However, its role as the plant defence system toward salt stress is controversial which proline are more recognized as a symptom of stress injury instead of being an indicator of stress tolerance (Htwe *et al.*, 2011). Therefore, the clear explanation on molecular adaptation and biochemical mechanism that occur in plant as responds to salt stress are important in developing salt-tolerant plant that are tolerable to the saline environment.

OBJECTIVE

1. To isolate and characterize the Oryzacystatin-1 gene (*OC-1*) from Malaysian rice cultivars.
2. To study the *OC-1* gene expression level in stress-induced rice plantlets of MR211, MR220 and MR219.
3. To analyze the proline content in stress-induced rice plantlets of MR211, MR220 and MR219.

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