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The Role of Thiamine in Plants and Current Perspectives in Crop Improvement

Atiqah Subki, Aisamuddin Ardi Zainal Abidin and Zetty Norhana Balia Yusof

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Abstract

Current research is focusing on selecting potential genes that can alleviate stress and produce disease-tolerant crop variety. The novel paradigm is to investigate the potential of thiamine as a crop protection molecule in plants. Thiamine or vitamin B_1 is important for primary metabolism for all living organisms. The active form, thiamine pyrophosphate (TPP), is a cofactor for the enzymes involved in the synthesis of amino acids, tricarboxylic acid cycle and pentose phosphate pathway. Recently, thiamine is shown to have a role in the processes underlying protection of plants against biotic and abiotic stresses. The aim of this chapter is to review the role of thiamine in plant growth and disease protection and also to highlight that TPP and its intermediates are involved in management of stress. The perspectives on its potential for manipulating the biosynthesis pathway in crop improvement will also be discussed.

Keywords: thiamine, vitamin B1, plant protection, stress, crops

1. Introduction

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Thiamine also known as vitamin B_1 was the first vitamin type B identified [1]. Free thiamine, thiamine monophosphate (TMP) and thiamine pyrophosphate (TPP) are the three most predominant forms of B_1 that exist in the cells [2]. Vitamin B_1 is a colourless, water-soluble vitamin made solely by plants and microorganisms and act as essential micronutrient in the human diet [3].

Thiamine occurrence in plants is widely distributed across organs, namely, leaves, flowers, fruits, seeds, roots, tubers and bulb [4]. Studies in *Arabidopsis* plant showed that the most abundant vitamer is TPP followed by TMP and thiamine, respectively [5]. The concentration

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of thiamine vitamer can be increased by supplementation of hydroxyethyl-thiazole (HET) and hydroxy-methylpyrimidine (HMP) [3]. The highest concentration of vitamin B in plants can be secured up to μ g/g relatively [6]. Diverse B₁ sources include yeast, cereal grains, beans, nut and meat [7].

2. Role of thiamine

In plants, thiamine is known to have its role as a cofactor for important metabolic activities [8]. Thiamine is known to be an essential regulator that plays an important role in plant's primary regulatory system [9]. Living organisms require the active form of thiamine which is known as thiamine pyrophosphate (TPP) in order to play the role as an important cofactor. TPP is a crucial component required in many metabolic activities such as acetyl-CoA biosynthesis, amino acid biosynthesis, Krebs cycle and Calvin cycle [10].

3. Role of thiamine in plant protection

In plants, thiamine plays a role as a response molecule towards abiotic and biotic stresses, and data from the literature suggest that boosting thiamine content could increase resistance to stresses [11]. Biotic stress is usually involve in the damage of plants caused by living organisms, while abiotic stress is due to environmental factors which cause a series of morphological, physiological, biochemical and molecular changes to plants that will affect the plants' growth, development and productivity [12].

Previous study that the effect of the infection of *Ganoderma boninense*, a pathogenic fungus, to the expression of *ThiC* gene in oil palm suggests that thiamine may play an important role in dealing with biotic stress [13]. Comprehensive studies on the effect of abiotic stresses on the regulation of thiamine in oil palm were also done where various types of stresses, namely, oxidative, salinity and osmotic stresses, have been induced in oil palm where an increase in gene expression and also total thiamine content was observed post-stress inductions [14–17]. A study by Kamarudin et al. explored the application of an endophytic fungus, *Hendersonia toruloidea*, in elevating the expression of thiamine and its intermediates in the plant [18, 19]. It was clear that a fungal endophyte could also boost thiamine content in oil palm. Current work is providing data on thiamine accumulation in oil palm seedlings upon application of beneficial endophytic bacteria, namely, *Pseudomonas aeruginosa* and *Burkholderia cepacia*.

Besides that, a study on the impact of *ThiC* promoter as well as its riboswitch on thiamine regulation in *Arabidopsis* sp. showed that the transcript of *ThiC* gene is highest at the end of the light period and lowest at the end of dark period [20]. Other than that, the responses of thiamine biosynthesis genes under several types of abiotic stresses such as salt and osmotic stress in *Arabidopsis* were examined, and it was found that these conditions have caused the

upregulation of the expression of the genes, thus eventually causing significant changes in thiamine level [5].

A study by Croft et al. revealed the declination of *ThiC* gene expression upon exogenous application of thiamine, which suggests a feedback regulation system in thiamine biosynthesis of green alga, *Chlamydomonas reinhardtii* [21]. On the other hand, Mcrose et al. proved that the relative gene expression of prasinophyte algae, *Emiliania huxleyi*, was significantly increased when thiamine supply was exhausted [22]. It has been demonstrated in *Cassava* sp. plant that the application of exogenous thiamine led in the formation of splicing variants of *ThiC* gene suggesting the presence of TPP riboswitch [23].

Thiamine possesses an antioxidant capacity as it has O_2^-/OH^- scavenger properties [24]. Vitamin B₁ is responsible for the recycling of vitamin C through the synthesis of nicotinamide adenine dinucleotide phosphate (NADPH) [4]. The antioxidant properties of thiamine were seen in a study on *Arabidopsis* sp. where paraquat-treated plant caused reduction in protein carbonyls and dichlorofluorescein diacetate (indicator of oxidative stress) when thiamine was applied [25]. Thiamine pyrophosphate indirectly acts as antioxidant by supplying NADH and NADPH to tackle oxidative stress [25]. However, out of all the studies conducted, scientists still find difficulties to unravel the cellular mechanism of B₁ as an antioxidant either through indirect effect of cofactor or as direct effect as antioxidant [4].

Recently, it has been reported that thiamine formed an indirect role in enhancing anti-oxidative capacity in plants, which is important in defence responses [26]. In addition, systemic acquired resistance (SAR) in *Oryza sativa, Arabidopsis thaliana, Nicotiana* sp. and *Cucumis sativus* was shown to be induced when thiamine was applied to these plants [27].

4. Thiamine biosynthesis is regulated by TPP riboswitch

In the past years, we recognise DNA as the main key on every single reaction that occurs in the cellular environment. The paradigm has been shifted to RNA nowadays. Since RNA sequences can carry out diverse tasks and are amenable to engineering both *in vitro* and *in vivo*, they are particularly attractive for controlling cell behaviour [28].

5. Thiamine pyrophosphate: the dominant class of riboswitch

Riboswitch is a natural RNA sensor that allows the direct binding of small metabolites, thus regulating the expression of various metabolic genes without the needs of protein cofactor [29, 30]. Without the protein involvement, regulation of gene expression can still occur due to the direct metabolite binding at riboswitch sequence [31]. RNA can specifically recognise and bind other molecules, including low molecular weight metabolites [32]. This includes nucleobases, cofactors, amino acids, second messenger and metal ion [33]. The metabolites are usually small, non-toxic molecule which exhibits a good cell permeability [34].

To date, there are about 15 riboswitch classes reported as shown in **Table 1**, and more of it is still unknown [35]. Among all classes of riboswitches, TPP riboswitches are the most ubiquitous in three life domains [36]. Thiamine pyrophosphate (TPP) is the most abundant riboswitch and is known to be present even in eukaryotes [37]. It has an intermediate level of sequence conservation [38]. So in many organisms (prokaryotes, algae, plants and fungi), riboswitch has been found to play the role of regulating thiamine biosynthesis [39].

In all plant taxa, the TPP riboswitch is present in the *ThiC* gene, and some of the TPP riboswitches that are lost during the gymnosperm evolution are present in the *Thi1* gene of ancient plants [40]. Studies by Cheah and co-worker testified that *Thi4* and N-myristoyltransferase (*NMT*) genes in *Neurospora crassa* are controlled by TPP riboswitch by splicing mechanism of an intron located in the 5' untranslated region (UTR) [39].

From the perspective of evolution, the presence of TPP riboswitch in ancient plant taxa suggests that this mechanism is active 400 million years ago, in early emergence of vascular plants [40]. The ancient plant taxa including ancient land plants consist of supplementary TPP riboswitch which ought to be found in the *Thi1* gene and no longer found, suggesting that during gymnosperm evolution, this sequence might be lost from this family gene [40]. Apart from that, the alternative splicing of 3' UTR gene also found in lycophytes, which are an ancient vascular plant family that existed around 150–200 million years before angiosperm (i.e. *Arabidopsis* and rice) [40]. **Table 2** shows the list of the discovered TPP riboswitches in various organisms.

Generally, riboswitches in bacteria can be found on the upstream 5' region of the non-coding region of mRNA, while in plant and fungi, this regulatory element resides at the 3' end of the untranslated region of a gene [20, 40, 41]. Although the location of TPP riboswitch in prokaryotes and eukaryotes might differ, its structure reveals a high similarity. This difference in location suggests a unique mode of action for the plant riboswitch [40].

Туре	Riboswitches class	Gene	Reference
Amino acid derivatives	Purine	ydhL	[48]
	Lysine	Asd	[41]
	Glycine		[49]
Carbohydrates	Glucosamine-6-phosphate	glmS	[50]
Enzyme cofactor	Flavin mononucleotide	ThiC	[29]
	Thiamine pyrophosphate	BtuB	[51]
	Cobalamin (B_{12})	S-box	[52]
	Tetrahydrofolate (THF)		[53]
	S-adenosyl methionine		
	S-adenosyl homocysteine		
Nucleotide precursor	Adenine, guanine	pbuE	[54]
	c-di-GMP	tfoX	[55]
	pre-queuosine (preQ1)	ykv	[56]

Table 1. Riboswitch classes reported across all kingdom of life.

Gene	Location	Organism	Reference
ThiC	3' UTR	Arabidopsis thaliana	[41]
	5′ UTR	Chlamydomonas reinhardtii	[21]
	3' UTR	Oryza sativa	[49]
		Poa secunda	[49]
		Solanum lycopersicon	[26]
		Thalassiosira pseudonana	[37]
		Phaeodactylum tricornutum	[37]
		Alishewanella sp.	[36]
		Flowering plant	[58]
Thi4	5′ UTR	Neurospora crassa	[39]
		Volvox carteri	[21]
		Fusarium oxysporum	[41]
ThiA		Aspergillus oryzae	[52]
ThiM	5′ UTR	Escherichia coli	[49]
Thi-box		Bacillus subtilis	[29]
		Rhizobium sp.	[60]
Thi1	3' UTR	Ancient plant (bryophytes, lycophytes)	[40]
ThiR	5′ UTR	Haloferax volcanii	[61]

Table 2. The list of RNA regulatory element involved in thiamine biosynthesis pathway, TPP riboswitch, in various organisms.

The biosynthesis of thiamine is uncommon from other vitamins. This is because previous study by Guan et al. revealed that the energy cost of thiamine synthesis is higher as compared to other vitamin cofactors [42]. Therefore, the location of riboswitch at the initial pathway strongly suggests that a novel riboswitch regulates the regulation of thiamine.

6. Thiamine biofortification in plants

As previously mentioned, thiamine has shown to act as cofactor and activator for plant stress and disease resistance. Furthermore, supplementation and accumulation of thiamine in plants showed no evidence of toxicity towards the plants as supported by the feeding studies [3]. However, a review by Goyer in 2010 suggested that thiamine production will be regulated in order to perfectly match the production to the demand of the cofactor. The study also stated that thiamine biosynthesis is regulated via (1) riboswitch-dependent gene regulation and (2) tissue specificity, stress dependence and post-translational regulation. Tissue-specific transcription factors have been found in THI1 gene, and the regulation has been widely studied [43] at the promoter level. The promoter activity in the roots is not due to light regulation but rather to promoter tissue specificity. On the other hand, stress dependence can be seen in maize seedlings where under osmotic and oxidative stresses, TPK enzyme activity increased [44] but exhibited a decrease under normal condition [45]. Furthermore, post-translational regulation or feedback inhibition has been identified in TH1 where excess of HMP-PP and ATP has shown to inhibit TH1 activity.

Gene transcript/ enzymes/thiamine derivatives	Stress	Outcomes	References	
THIC	Oxidative, osmotic, temperature (cold), biotic (colonisation by endophyte)	Increase in expression	[15, 18, 25, 62, 63]	
	Exogenous thiamine	Decrease in expression	[64]	
THI4	Light, oxidative, biotic (colonisation by endophyte)	Increase in expression	[18, 25, 65, 66]	
	Dark	Decrease in expression		
TH1	Oxidative, biotic (colonisation by endophyte)	Increase in expression	[18, 25, 45]	
ТРК	Osmotic, salinity, oxidative, biotic (colonisation by endophyte)	Enzyme activity increase	[18, 44, 45]	
Total thiamine	Osmotic, salinity, oxidative, biotic (colonisation by endophyte)	Increase in concentration	[18, 25, 45]	
	Exogenous thiamine	Decrease in expression	[64]	

Table 3. Effects of stress towards thiamine biosynthesis in plants.

Total thiamine content in wild-type plants is mainly composed of thiamine, thiamine monophosphate (ThMP) and thiamine diphosphate (ThDP) [27]. Overexpression of THIC and THI4 simultaneously has shown to increased thiamine levels up to sixfold and ThDP levels twofold compared to single overexpression of either THIC or THI4 which showed no elevation of total thiamine content [11]. This shows the relationship between thiamine biosynthesis genes and thiamine production. Elevation of thiamine content and also the thiamine biosynthesis gene transcripts in plants have been demonstrated quite extensively via the application of biotic and abiotic stresses. Utilisation of these stresses may aid in the fortification of thiamine in crops. **Table 3** shows the studies done in understanding the effects of the application of stress towards thiamine production in plants.

Apart from that, higher possibilities of thiamine fortification in plants could be achieved via genetic manipulation. Genetic engineering via mutation of riboswitch coding sequence in plant model organism, *Arabidopsis*, has produce an organism with deficiency in TPP riboswitch activity and enhanced accumulation of total thiamine esters [20]. However, due to increasing TPP concentrations, this condition has led to an increase of metabolic flux into the TCA cycle and pentose phosphate pathway which causes a significant increase in the organism respiratory rate, hence more CO₂ production [20]. Genetic manipulation in *Arabidopsis* and rice by overexpression of THIC and THI4 has shown to increase thiamine levels up to sixfold and ThDP levels twofold in *Arabidopsis* and increased total thiamine level up by fivefold in *Oryza sativa* [11, 46]. Furthermore, genetic manipulation of TPK via promoter enhancement in *Arabidopsis* has led to an increased expression of TPK up to 30-fold and transketolase enzyme activity by 2.5-fold [47]. The mutant plant also resulted in chlorotic and slow-growth characteristics. However, levels of total thiamine of mutant plants were significantly lower compared to control.

7. Conclusion

Overall, based on the extensive studies done, thiamine fortification in plants could be achieve via both abiotic and biotic stress and genetic engineering [20, 24, 25, 45]. Manipulation by the knowledge available on the riboswitch associated with THIC could likely be an effective strategy to manipulate thiamine levels in plants, especially in terms of biofortification. However, it is well agreed that the process on enhancing thiamine levels in plants is not as straightforward and as easy as it seems. Further understanding of the two key precursors (HMP and HET) will be required as this will lead to the accumulation of thiamine, with hopefully least side effects. These two intermediates have been shown to be not toxic to plants, and plant tolerance towards stress is expected to increase when the levels of these two intermediates are enhanced. However, the modification of this will still come with its own challenges since it involves highly complex enzymes which are regulated very tightly and there have not been much studies on the understanding of the mechanisms just yet.

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Conflict of interest

The authors declare that there is no conflict of interest.

Author details

Atiqah Subki, Aisamuddin Ardi Zainal Abidin and Zetty Norhana Balia Yusof*

*Address all correspondence to: zettynorhana@upm.edu.my

Department of Biochemistry, Faculty of Biotechnology and Biomolecular Sciences, Universiti Putra Malaysia, Serdang, Selangor, Malaysia

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