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The significance of calcium-sensing receptor in sustaining photosynthesis and ameliorating stress responses in plants

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Calcium ions (Ca^{2+}) regulate plant growth and development during exposure to multiple biotic and abiotic stresses as the second signaling messenger in cells. The extracellular calcium-sensing receptor (CAS) is a specific protein spatially located on the thylakoid membrane. It regulates the intracellular Ca²⁺ responses by sensing changes in extracellular Ca^{2+} concentration, thereby affecting a series of downstream signal transduction processes and making plants more resilient to respond to stresses. Here, we summarized the discovery process, structure, and location of CAS in plants and the effects of Ca²⁺ and CAS on stomatal functionality, photosynthesis, and various environmental adaptations. Under changing environmental conditions and global climate, our study enhances the mechanistic understanding of calcium-sensing receptors in sustaining photosynthesis and mediating abiotic stress responses in plants. A better understanding of the fundamental mechanisms of Ca²⁺ and CAS in regulating stress responses in plants may provide novel mitigation strategies for improving crop yield in a world facing more extreme climate-changed linked weather events with multiple stresses during cultivation.

KEYWORDS

abiotic stress, photosynthesis, calcium, biological functionality, calciumsensing receptor

Introduction

Calcium, a crucial macronutrient for plants, has important physiological functions such as maintaining cell morphology and regulating ion balance and osmotic pressure. Interestingly, it is also a key component for several early signaling pathways involved in plant-environmental stresses interactions (Lambers and Oliveira, 2019; Song et al., 2020; Wu et al., 2020). Most elemental calcium in plant cells exists in ionic forms, which contribute to forming microtubules in the cytoskeleton and maintaining the fluidity of cell membranes (Kong et al., 2020; Liu, 2021). In addition, calcium signals assist in the integration of various signaling pathways to coordinate plants' growth response to multiple environmental signals (Edel and Kudla, 2016; Kudla et al., 2018; Luan and Wang, 2021; Kim et al., 2022). The extracellular calcium-sensing receptor (CAS) is an important protein that specifically exists in plants and has specific regulatory functions. These proteins regulate intracellular Ca²⁺ responses by sensing changes in extracellular Ca²⁺ concentration, thereby affecting a series of downstream signaling activities and fine-tuning plants' responses to environmental perturbations. We summarized the research progress of Ca²⁺ and CAS in regulating photosynthesis and stress responses. Here, we focused on the fundamental role of CAS in regulating growth and signal transduction in response to the external environment.

Structural characteristics of the calcium-sensing receptor

Discovery and location of CAS

The extracellular Ca²⁺ receptor gene was cloned for the first time using the functional gene screening method in Arabidopsis thaliana and named as the calcium-sensing receptor (CAS) (Li et al., 2022). With the aid of fluorescence imaging and proteomics technologies, the CAS was located on the chloroplast thylakoid membrane of Arabidopsis thaliana, spinach, and green algae (Allmer et al., 2006; Naumann et al., 2007; Nomura et al., 2008; Weinl et al., 2008). Interestingly, the CAS was also detected within the purified eyespot of green algae (Trippens et al., 2017). Yamano et al. (2018) fused the CAS-Clover with high-resolution fluorescence imaging to coincide with the eyespot, which also verified the existence of CAS. In addition, high-resolution fluorescence images of CAS were obtained using the sensitive hybrid detector and image deconvolution technology, and its cellular localization on the thylakoid membrane was authenticated and observed with greater clarity.

Structural features of CAS

The carboxyl-terminal of CAS is located within the cell and has two domains (Vainonen et al., 2008); one of which is a noncatalytic thiocyanate domain with high homology, and the other is related to protein interaction which can interact with the 14-3-3 and FHA (forkhead-associated) domain. Under light conditions, the CAS light-regulated kinase STN8 is targeted for phosphorylation at Thr-380 (Vainonen et al., 2008). The amino terminal of CAS is located outside the cell membrane, which may plausibly be the Ca²⁺ binding region, although its conserved sequence has not been found. It was later verified that the binding site of Ca²⁺ in CAS was the amino end rather than the carboxyl end in *Arabidopsis* and *Chlamydomonas* (Wang et al., 2016a).

CAS and stomatal functionality

The pair of guard cells have a fundamental role in regulating gas exchange between plants and the atmosphere, and biotic infection (Melotto et al., 2008; Chen et al., 2012; Sawinski et al., 2013; Sun et al., 2014; Murata et al., 2015; Agurla et al., 2016; Bharath et al., 2021). Generally, Ca²⁺ plays an important role in regulating guard cell turgor and the movements of guard cells. Ca^{2+} ions affect stomatal conductance to water vapor and CO_2 by regulating the pore size. Peng et al. (2022) discovered that an exogenous application of 18 Mm Ca²⁺ could increase the stomatal conductance of Paris polyphylla under high temperatures and strong light conditions. The CO2-induced stomatal closure process is also calcium-dependent and the CPK (calciumdependent protein kinase) has a role in signal transduction and regulating the ion channels of guard cells. In the pentaploid mutant plants with cpk3/5/6/11/23, the stomatal movements were significantly impaired, suggesting that CPK has important functions in facilitating stomatal movement and is plausibly regulated by CO₂ concentration (Schulze et al., 2021). Jakobson et al. (2016) confirmed that there were significant differences in stomatal phenotypes among cas mutants with different alleles and the cas overexpression of Arabidopsis transformants promoted stomatal closure. Li et al. (2017) also demonstrated that the deletion of the CAS gene could disrupt stomatal closure and this phenomenon was attributed to high levels of extracellular Ca^{2+} . It was later verified that the extracellular high Ca^{2+} in CAS gene-deficient mutants could not induce stomatal closure, indicating that CAS was essential for extracellular Ca²⁺-induced stomatal closure in Arabidopsis (Qi et al., 2018).

Interestingly, it was also found that CAS has an important role in elevating intracellular Ca^{2+} concentration when induced by extracellular Ca^{2+} . These studies provided evidence for *Arabidopsis* chloroplasts in regulating extracellular Ca^{2+} -induced intracellular Ca²⁺ concentration increases and subsequent stomatal closure (Nomura et al., 2008; Li et al., 2022). Wang et al. (2016b) found that the chloroplastic regulation of stomatal closure required the reduction of quinone, foliar production of H₂O₂, and the phosphorylation of CAS and LHCII during exposure to unfavorable perturbation. It was confirmed that stomatal movement was associated with NO and the NO acted downstream of the H₂O₂ step in regulating stomatal movement (Bright et al., 2006). Specifically, the stomatal closure induced by H₂O₂ and NO requires the involvement of extracellular Ca²⁺ and the accumulation of H₂O₂ and NO in guard cells (Li et al., 2009; Zhao et al., 2011). Tang et al. (2007) suggested that CAS might regulate the stomatal closure process through the Ca²⁺-CAS-IP signaling pathway. Subsequently, Wang et al. (2012) found that the extracellular Ca2+ induced H2O2 and NO accumulation through CAS in Arabidopsis thaliana, which induced transient changes in cytoplasmic Ca²⁺ concentration and was able to regulate stomatal movement. In addition, Arabidopsis plants expressing PHPLCS-GFP protein were used for PHPLC8-GFP imaging of guard cells, which indicated that H2O2 and NO were involved in the CAS-IP3 signaling pathway (Wang, 2013). Li et al. (2017) found that H₂O₂ might directly or indirectly regulate CAS activity and serve as redox signaling molecules, thereby regulating stomatal closure. Interestingly, the data also revealed that the exogenous NOinduced stomatal closure of CAS guard cells was facilitated by exogenous NO bypassing the damaged CAS. Furthermore, it was indicative that NO has a downstream role relative to CAS in the signal transduction pathway. Recent studies suggested that stomatal movement induced by extracellular Ca2+ was the culmination of multiple interactions among CAS, H₂O₂ and NO signals (Zhang et al., 2018; Li et al., 2022). Li et al. (2017) used multi-disciplinary approaches to examine the responses of different mutants to extracellular Ca2+ in guard cells. It was found that there was a cascade of H₂O₂, ABA, and NO in the CAS signaling pathway. With further studies, it was confirmed that seed germination and stomatal closure of mutant cau1 transformants were excessively sensitive to ABA. In addition to extracellular Ca²⁺, the CAU1-CAS could also respond to ABA through interaction, and ABA and CAUI could also regulate stomatal closure (Ueda and Seki, 2020), although the specific role of CAS in this process required further clarification. Li et al. (2017) found that ABA in CASas mutant plants was able to induce stomatal closure, indicating that the ABAinduced signal transduction pathway was not dependent on CAS; here, CASas refers to the CAS antisense line of Arabidopsis. Based on these studies, the signal transduction cascade is summarized accordingly in a scheme (Figure 1).

CAS and photosynthesis

Photosynthesis is one of the most important biological processes on earth (Liu et al., 2011; Stael et al., 2012; Lambers

and Oliveira, 2019). Calcium is involved in regulating various photosynthetic processes, such as CO₂ fixation and protein phosphorylation in chloroplasts (Rocha and Vothknecht, 2012; Liu, 2020). Hu et al. (2022) found that low calcium levels caused the decline in photosynthesis of Taxus wallichiana varieties, but this negative effect could be reversed by applying supplementary calcium nutrition, which was consistent with previous findings in grape and cucumber (He et al., 2018; Duan et al., 2020). Harnessing the pull-down analysis (network database GeneCAT) combined with proteomics and gene co-expression method, Wang (2013) found that 52% of the CAS co-expression genes in Arabidopsis were related to photosynthesis. Combining photosynthetic indexes, chloroplast structural observation and gene expression profiling, these observations verified that the photosynthetic level and biomass accumulation of Arabidopsis CASas plants were significantly reduced. Specifically, the inhibition of CAS caused the transcription level of photosynthetic electron-related genes to be down-regulated (Wang et al., 2014). Navazio et al. (2020) found that the transcription expression of the CAS gene in Arabidopsis was up-regulated under strong light conditions, and the CAS transcription level under dark and weak light conditions was also significantly up-regulated compared with that under normal light conditions. Petroutsos et al. (2011) confirmed that CAS played a role in maintaining PSII functionality and helping plants to adapt to strong light. Additionally, Li (2021) found three AtCAS homologous genes in some shade plants such as ginseng, and two AtCAS homologous genes in the non-shade plant such as tomato. As CAS is a phosphorylated protein regulated by light intensity; it is plausible that CAS may be closely related to light adaptability. The stomatal movement of terrestrial plants regulates CO2 entry and exit through HTI protein kinase and carbonic anhydrase (Hu et al., 2010), and aquatic plants maintained stable photosynthesis through the CO₂-concentration mechanism (CCM). Studies have indicated that Ca²⁺ and CAS are related to CCM. In response to low CO₂ concentration, the CAS can be transferred to the protein nucleus, and then mediate the chloroplastic retrograde signaling to maintain the expression of HLA3 (high-light activated 3) and LCIA (low-CO2-inducible gene A) (Wang et al., 2016a). Yamano et al. (2018) obtained interesting CAS-Clover high-resolution fluorescence images using a sensitive mixed detector coupled with image deconvolution technology. It was found that CAS moved along the thylakoid membrane in response to low concentration CO₂, and gathered spatially in the protein nucleus during the CCM mechanism. As a chloroplastic thylakoid membrane protein, CAS functionality was examined during de-etiolation and chloroplast development in Arabidopsis thaliana (Huang et al., 2012). The thylakoid-localized CAS protein could assist in the production of cytosolic calcium transients, and the activation of the MPK3/MPK6 signal system. Subsequently, the activated MPK3/MPK6 was involved



in the nucleus-induced phosphorylation of ABI4. The activation of ABI4 at both the transcriptional and posttranslational levels induced the inhibition of LHCB (Gao et al., 2016). These results showed that the cystoid-located CAS was involved in the retrograde signaling pathway from chloroplast to nucleus (Li et al., 2022).

Protein phosphorylation is vital for photosynthesis and adaptation to multiple environmental stresses (Vener, 2007; Fristedt et al., 2010). Vainonen et al. (2008) found that CAS could be phosphorylated by STN8, and the phosphorylation intensity increased with light intensity enhancement. At the same time, a new phosphorylation site Thr-367 was also found in the CAS of green algae (Lemeille et al., 2010). Cutolo et al. (2019) conducted proteomics analysis and phosphorylation determination of CASas, and identified Thr-376, Ser-378, and Thr-380 as the main phosphorylation sites of STN kinase. Protein phosphorylation is also important to regulate photosynthetic electron transport efficiency (Reiland et al., 2011). The cyclic electron transport (CET) process is involved in chloroplast energy regulation and redox metabolism, which is particularly important for photosynthesis. The studies demonstrated that CET has two pathways, PGR5/PGRL1mediated pathway and the NDH-mediated pathway (Johnson et al., 2014; Ma et al., 2021a). The protein interaction between CAS and PGRL1 was confirmed in *Chlamydomonas reinhardtii*, indicating that there is a link between CAS and CET (Terashima et al., 2012; Ma et al., 2021b). Chen et al. (2015) found that Ca²⁺ controlled the biosynthesis of lipids induced by chloroplastic nitrogen starvation by promoting the expression of CAS in the PGRL1-mediated CET pathway.

CAS, stress resilience, and plant defense

When plants are exposed to environmental stresses, they will improve their resilience by self-regulating several *in vivo* processes (e.g. changes in intracellular calcium concentration) to cope with these perturbations (Liu, 2020). Salinity is an important environmental factor, which has a negative impact on plant growth and photosynthesis (Acosta-Motos et al., 2017; Ma et al., 2022). Similarly, Ca²⁺ is also crucial in mediating stress responses (Park et al., 2016). It was verified that Ca²⁺ can enhance the salt tolerance of plants by improving water

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balance, regulating sodium secretion, and enhancing membrane integrity (Ahmed et al., 2021). In addition, Ca²⁺ also enhances resilience to chilling stress. Liu et al. (2013) found that exogenous Ca2+ could mitigate the significant decline in peanut photosynthesis and biomass accumulation during exposures to low night temperatures. It was also confirmed that exogenous Ca2+ ameliorated night chilling-dependent feedback inhibition of photosynthesis by improving sink demand and facilitating nonstructural carbohydrate export from chloroplasts, to restore peanut growth, dry matter production, and leaf photosynthetic capacity (Shi et al., 2020; Song et al., 2020; Wu et al., 2020). Sun et al. (2022) reported that the Ca²⁺ mediated-CAS was crucial for alleviating photoinhibition in peanut growing under conditions of moderate phosphorus deficiency. In addition, Zhao et al. (2015) confirmed that OsCAS could improve drought tolerance in Arabidopsis. When drought and heat stresses were imposed on the creeping bentgrass (Agrostis stolonifera), the expression of CAS was down-regulated, indicating that the calcium sensitivity to drought and heat stress responses was reduced (Xu and Huang, 2018). The molecular mechanism of proline metabolism regulation under drought also revealed a new drought tolerance pathway mediated by CAU1 (Fu et al., 2018). Huang et al. (2012) provided further evidence to support the role of Ca²⁺ and CAS in leaf re-greening and chloroplast development of Arabidopsis. Zheng et al. (2017) screened a chloroplast protein QUA1 by harnessing the large-scale forward genetic method. It is plausible that the CAS-mediated salt and drought tolerance of plants through the calcium signaling-linked QUA1 that influence the stability of CAS. It was further suggested that CAS has a regulatory role in the downstream of QUA1. Interestingly, there are some pieces of evidence to indicate that CAS is a prerequisite for chloroplast Ca²⁺ to induce light response and light-dark transition (Nomura et al., 2012). The expression of cas in Arabidopsis thaliana was significantly reduced under high-temperature treatment. It is plausible that the heat-sensing ability of chloroplasts was partially dependent on CAS during high-temperature exposure (Lenzoni and Knight, 2019).

Plant defense against invasive pathogens is dependent on the dual innate immune system. Meta-analysis indicated that CAS regulated the expression of flg22-induced immune genes through ${}^{1}O_{2}$ -mediated retrograde signaling; flg22 is a peptide derived from bacterial flagellins. These studies revealed that the weakened signals, chloroplasts, and superoxide anions were collectively involved in the innate immune system (Sano et al., 2014; Stael et al., 2015). In the study of plant-pathogen interaction mediated by small molecules during the invasion of *Fusarium graminearum*, Jia et al. (2019) found that the fungal infection or the use of exogenous fusaoctaxin A inhibited the expression of three CAS-like genes and chloroplast genes. Thus, the lowered

susceptibility of wheat to Fusarium A, and the inhibition of photosynthesis provided some evidence that chloroplasts might play a key role in regulating early immune responses (de Torres Zabala et al., 2015). Sclerotinia sclerotiorum is a well-known necrotizing fungus that attacks many crops (Bolton et al., 2006). The CAS is involved in the defense response of Arabidopsis thaliana to S. sclerotiorum, and actively regulates the accumulation of salicylic acid (SA) by promoting the expression of SA biosynthesis-related genes, thereby enhancing the resistance of plants to S. sclerotiorum (Tang et al., 2020). HopAU1 is an immune inducer existing in Pseudomonas syringae pv. actinidiae (Psa) which can interact with NbCAS in tobacco. Thus, silencing the NbCAS by RNAi in N. benthamiana greatly attenuated HopAU1-triggered cell death, suggesting that HopAU1 targets CAS and enhances plant immunity. Further study showed that the overexpression of NbCaS in N. benthamiana significantly improved plant resistance against Sclerotinia sclerotiorum and Phytophthora capsici (Zhang et al., 2022). Taken together, the CAS also serves as a promising resistant-related gene for breeding new disease-resistant varieties.

Future outlook

The CAS has an important role in facilitating Ca²⁺-mediated stomatal closure and is associated with several signaling molecules. As a specific protein located on the chloroplast thylakoid membrane, the CAS functions as the first messenger along the signal transduction pathway in plants. There is some evidence that CAS may participate in various photosynthetic processes such as CO₂ fixation and protein phosphorylation. With further research, we could gain insights into the Ca²⁺induced signaling pathway and the functional characteristics of CAS in plant cells. In the future, it is important to further explore the following: (i) the various relationships between CAS and the other calcium-linked signals; (ii) the potential cross-talk between CAS and light signaling at the chloroplast thylakoid membranes; (iii) the CAS and its specific role in facilitating plant abiotic and biotic stress responses using advanced techniques/tools (proteomics, transcriptomics, and bioinformatics, etc.); (iv) extrapolating the understanding of CAS in model plants to non-model plants such as the legumes and horticultural crops, etc.

Author contributions

YL, RB, CB, XH, and JY are responsible for the general review described in the manuscript. YL, XH, and JY provided further discussion and edited the manuscript. All authors reviewed and approved the final version of the submitted manuscript.

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Conflict of interest

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