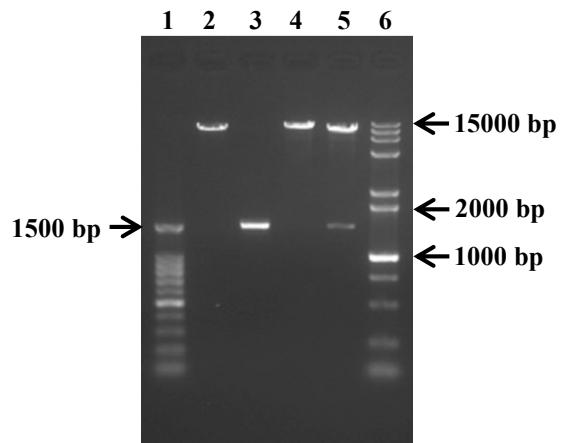


# Supplementary Materials: The Role of Sugarcane Catalase Gene *ScCAT2* in the Defense Response to Pathogen Challenge and Adversity Stress

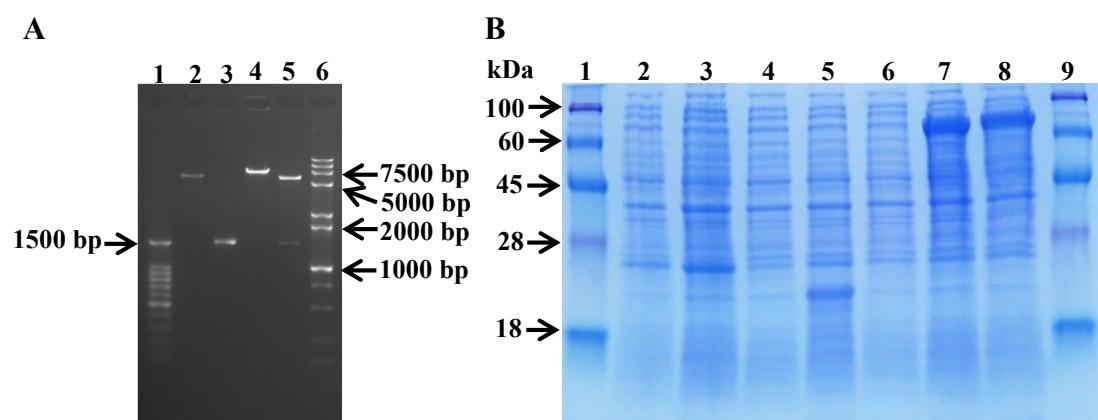
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**Table S1.** Primers used in this study.

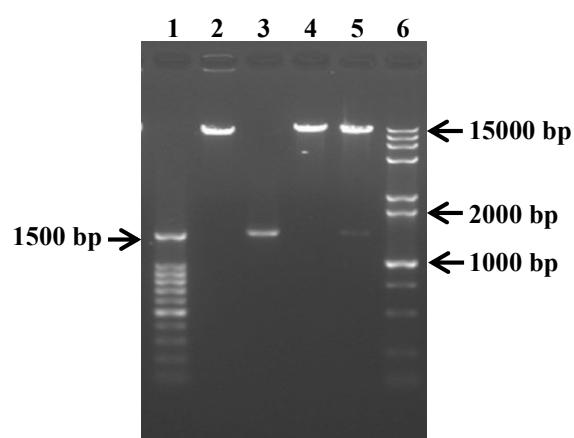
Primer	Sequence	Strategy
<i>ScCAT2</i> -cDNAF	ACACCTCCACTCTCCAACACT	RT-PCR
<i>ScCAT2</i> -cDNAR	CAATTGCCATCACTCACAT	RT-PCR
<i>ScCAT2</i> -QF	GGAGTGGAAAGCTGTTCGTGC	qRT-PCR
<i>ScCAT2</i> -QR	CGCCGTCGTAGTGGTTGTTA	qRT-PCR
<i>GAPDH</i> -QF	CACGGCCACTGGAAGCA	qRT-PCR
<i>GAPDH</i> -QR	TCCTCAGGGTTCCTGATGCC	qRT-PCR
<i>ScCAT2</i> -SublocF	TGCTCTAGAATGGATCCCACCAAGTTC	Subcellular localization vector construction
<i>ScCAT2</i> -SublocR	GGACTAGTCATGTTGGCTTCATGTTGAG	Subcellular localization vector construction
<i>ScCAT2</i> -32aF	CGGAATTCATGGATCCCACCAAGTT	Prokaryotic expression vector construction
<i>ScCAT2</i> -32aR	CCCTCGAGTCACATGTTGGCTTCATG	Prokaryotic expression vector construction
<i>ScCAT2</i> -1301F	TGCTCTAGAATGGATCCCACCAAGTTC	Overexpression vector construction
<i>ScCAT2</i> -1301R	GGACTAGTCACATGTTGGCTTCATGTT	Overexpression vector construction
<i>NtHSR201</i> -F	CAGCAGTCCTTGGCGTTGTC	qRT-PCR
<i>NtHSR201</i> -R	GCTCAGTTAGCCGCAGTTGTG	qRT-PCR
<i>NtHSR203</i> -F	TGGCTCAACGATTACGCA	qRT-PCR
<i>NtHSR203</i> -R	GCACGAAACCTGGATGG	qRT-PCR
<i>NtHSR51</i> -F	TTGGGCAGAACATAGATGGTA	qRT-PCR
<i>NtHSR51</i> -R	TTTGGTGAAAGTCTTGGCTC	qRT-PCR
<i>NtNPR1</i> -F	GGCGAGGAGTCCGTTCTTAA	qRT-PCR
<i>NtNPR1</i> -R	TCAACCAGGAATGCCACAGC	qRT-PCR
<i>NtPR-1a/c</i> -F	AACCTTGACCTGGGACGAC	qRT-PCR
<i>NtPR-1a/c</i> -R	GCACATCCAACACGAACCGA	qRT-PCR
<i>NtPR2</i> -F	TGATGCCCTTGGATTCTATG	qRT-PCR
<i>NtPR2</i> -R	AGTCCTGCCCGCTTT	qRT-PCR
<i>NtPR3</i> -F	CAGGAGGGTATTGCTTGTAGG	qRT-PCR
<i>NtPR3</i> -R	CGTGGGAAGATGGCTTGTGTC	qRT-PCR
<i>NtEFE26</i> -F	CGGACGCTGGTGGCATAAT	qRT-PCR
<i>NtEFE26</i> -R	CAACAAGAGCTGGTGTGGATA	qRT-PCR
<i>NtAccdeaminase</i> -F	TCTGAGGTTACTGATTGGATTGG	qRT-PCR
<i>NtAccdeaminase</i> -R	TGGACATGGTGGATAGTTGCT	qRT-PCR
<i>NtEF1-α</i> -F	TGCTGCTGTAACAAGATGGATGC	qRT-PCR
<i>NtEF1-α</i> -R	GAGATGGGGACAAAGGGGATT	qRT-PCR



**Figure S1.** Enzyme digestion of the subcellular localization expression vector pCAMBIA 2300-*ScCAT2-GFP*. 1, 100 bp ladder marker; 2, 35S::GFP/*Xba*I; 3, *ScCAT2* ORF PCR product; 4, 35S::*ScCAT2*::GFP/*Xba*I; 5, 35S::*ScCAT2*::GFP/*Xba*I + *Spe*I; 6, 15000 + 2000 bp DNA marker.



**Figure S2.** Enzyme digestion of prokaryotic expression vector pET-32a-*ScCAT2* (A) and corresponding protein expression in *Escherichia coli* Rosetta strain cells (B). (A) 1, 100 bp ladder marker; 2, pET-32a/*Eco*RI; 3, *ScCAT2* ORF PCR product; 4, pET-32a-*ScCAT2*/*Eco*RI; 5, pET-32a-*ScCAT2*/*Eco*RI + *Xho*I; 6, 15000 + 2000 bp DNA marker. (B) 1, Protein marker; 2, Rosetta strain cells without induction; 3, Rosetta strain cells induction for 2 h; 4, Rosetta + pET-32a strains without induction; 5, Rosetta + pET-32a strains induction for 2 h; 6, Rosetta + pET-32a-*ScCAT2* strains without induction; 7 and 8, Rosetta + pET-32a-*ScCAT2* strains induction for 1 h and 2 h, respectively.



**Figure S3.** Enzyme digestion of the transient overexpression vector pCAMBIA 1301-*ScCAT2*. 1, 100 bp ladder marker; 2, pCAMBIA 1301/*Xba*I; 3, *ScCAT2* ORF PCR product; 4, pCAMBIA 1301-*ScCAT2*/*Xba*I; 5, pCAMBIA 1301-*ScCAT2*/*Xba*I + *Spe*I; 6, 15000 + 2000 bp DNA marker.