

Supplementary Material



**Figure S1.** In silico (Genevestigator) deduced differential expressions of up-and downregulated cytokinin-Oglucosyltransferase (CGT) genes of rice in response to infections with *Rhizoctonia solani*, *Magnaporthe oryzae*, *Xanthomonas oryzae* pv. *oryzae*, and *Xanthomonas oryzae* pv. *oryzicola*. Downregulation is indicated by green colour and upregulation by red colour.

**Table S1.** Primers used in PCR assay to assess the expressions of CGT genes in on challenge inoculation with *M. oryzae*, *Xoo* and *R. solani*.

Gene	Primers	Sequence (5'-3')
<i>LOC_Os04g25440</i>	CGT-F	GGACGCACACGTAACAGTAA
	CGT -R	GTCAAGTCATGGAGGCAAGT
<i>LOC_Os04g25800</i>	CGT -F	TGCACAAGAGTGGAAGAGAA
	CGT -R	CTCACCATTCCACCCTGCTAAT
<i>LOC_Os04g25490</i>	CGT -F	TGGACACGAAGATGGATTGG
	CGT-R	CGTCACGTCGATGAAGAAGT
<i>LOC_Os04g37820</i>	CGT -F	GACGCATACAGAGACATGGG

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	CGT-R	GTTGACGAAGGTGATGTGGA
<i>LOC_Os07g30620.1</i>	CGT -F	CTAATCAGGCATATCTTCCCCAC
	CGT -R	TCACCCAAGTCGTTGTAGAAAG
<i>LOC_Os07g30610.1</i>	CGT-F	GTATCAACACCTTCCACCCC
	CGT-R	GATCTTGTACACGCCTAGAGG

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