

Supplemental Table S1. Inhibition of mycelium of three plant pathogenic fungi by isoxanthohumol.

Compound	Concentration (μ g/mL)	Inhibition rate ^a (%) \pm SD		
		<i>Sclerotinia sclerotiorum</i>	<i>Botrytis cinerea</i>	<i>Fusarium graminearum</i>
Isoxanthohumol	50	84.46 \pm 0.30	85.90 \pm 0.24	63.88 \pm 0.23
	25	79.49 \pm 0.05	78.45 \pm 0.32	60.43 \pm 0.82
	10	33.28 \pm 0.64	61.03 \pm 0.43	45.11 \pm 0.23
	5	11.15 \pm 1.82	57.00 \pm 0.60	27.08 \pm 1.24
	2.5	5.73 \pm 0.14	39.03 \pm 0.93	0.00 \pm 0.00
	EC₅₀	14.52	4.32	16.50

^a Repeat each treatment three times. Data are displayed as mean \pm SD.

Supplemental Table S2. Key DEGs involved in Carbon metabolism and TCA cycle and their primer sequences.

Gene alias	Control FPKM	Treat _FPKM	log ₂ Fold Change	Annotation	Primer sequences
BCIN_07g05430	0.273	37.186	6.58	Carbon metabolism	F:CGAGAACGGTGTGAGTTAGA TG R:CCTTCAGGTTCAGGGAAGA TT
BCIN_15g02910	0.790	4.343	2.46	Carbon metabolism	F:CTGTATCCACCCAAGCCTATA C R:TTTCTCTCGTCGTAGCATTC F:CAATCACTGACGAACCCTCT AA
Bcboa1	1.436	5.503	1.94	Carbon metabolism	R:AACCAGTCTCCATGCTTTTC F:ATGCCACCAAGACCATCAC R:CATCTCCGTCGCGTCAATAA F:CTCGTCTGTATGGCCGTATT R:CAAACCGACGAAGCACATTA C
Bcpio2	0.393	1.596	1.81	Carbon metabolism	
BCIN_12g00360	137.170	51.030	-1.33	Carbon metabolism	
BCIN_02g02750	316.836	529.293	0.82	Citrate cycle (TCA cycle)	F:GGATTTGGCGAGAACAAAGGA R:GTGAGTGGTGTGAGCAGAAA

BCIN_05g04430	176.703	288.606	0.79	Citrate cycle (TCA cycle)	F:TTGAATCCAGCTACCGATGC R:GCACTCTTCGATGGAAA TA
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