

Table S1. Name of each primer obtained, size of the insert ligated in each plasmid, and sequences of the primers used to amplify each insert.

Plasmid	Size of the insert (bp)	Sequence of primers (5'--3') used in amplification of inserts*
pJL-RNAi-adrA	382	Fw: AGACTCCATGGTTGCCGTAAATCCGACT
		Rv: AGACTCCATGGCGTCCACAAAGATCAAGG
pJL-RNAi-adrC	412	Fw: AGACTCCATGGACCAGCTAGAACTTACTC
		Rv: AGACTCCATGGTCGATTAGAGCAGTGGAT
pJL-RNAi-adrD	324	Fw: AGACTCCATGGCACCAAGATAACGAAGCAG
		Rv: AGACTCCATGGTTGTGCATCGCAGACGCT
pJL-RNAi-adrE	287	Fw: AGACTCCATGGCTTGGTTGTGGATGTGG
		Rv: AGACTCCATGGACCGCTCTCTAGGCTGAC
pJL-RNAi-adrF	401	Fw: AGACTCCATGGATCACCGTAGCGCCTCT
		Rv: AGACTCCATGGACCAACTCGTAGCTGCCT
pJL-RNAi-adrG	329	Fw: AGACTCCATGGTTGCCTTATAATCACATT
		Rv: AGACTCCATGGCCATCGCGAGGAATCCGG
pJL-RNAi-adrH	325	Fw: AGACTCCATGGTAGTTCACGCAGATGCC
		Rv: AGACTCCATGGTCCACTGCTGCTGAGGCC
pJL-RNAi-adrI	374	Fw: AGACTCCATGGATCATAGAGATGCCTTGG
		Rv: AGACTCCATGGCCCAACCTCTCGCGCAA
pJL-RNAi-adrJ	394	Fw: AGACTCCATGGGCCCAACTATCTATACA
		Rv: AGACTCCATGGCTCTTCTTCCCAGTGA
pJL-RNAi-adrK	371	Fw: AGACTCCATGGTCCTTGAGCTTGGCCTG
		Rv: AGACTCCATGGTTGTCGACTTTAGCTT

\* *Nco*I sites are highlighted in red.

Table S2. Primers used in qRT-PCR experiments

Target gene	Name of primer	Sequence 5'---3'	Amplicon size (pb)
<i>adrA</i>	adrA-qPCR fw	CGACTGGCTACGTCTCACTG	106
	adrA-qPCR rv	TAAAAAACCTAGCCCGATTG	
<i>adrC</i>	adrC-qPCR fw	TGTGTCAGATCCTCGTTGAA	85
	adrC-qPCR rv	CTGCCGTAAACCGAAAAGTA	
<i>adrD</i>	adrD-qPCR fw	GGCTCGGACGACTATACTGA	84
	adrD-qPCR rv	AGTACAGAACGCCTGGAGTG	
<i>adrE</i>	adrE-qPCR fw	CTCGTATTCCAGCTGCTCCT	94
	adrE-qPCR rv	CCAAAAGTCTCACGAAGCAA	
<i>adrF</i>	adrF-qPCR fw	CCATGAGACGACACCAAGGC	112
	adrF-qPCR rv	ATCAACGCTGGAATTCTGGT	
<i>adrG</i>	adrG-qPCR fw	GAGTCATCGAGCTCCTGTC	98
	adrG-qPCR rv	ACGAAAAATGACAGGCCAAG	
<i>adrH</i>	adrH-qPCR fw	GACACCCAATATCGGACAAG	93
	adrH-qPCR rv	AAGGCATCTCGTGAACCTAC	
<i>adrI</i>	adrI-qPCR fw	ACGTCGCGAAAAGACAAGAT	97
	adrI-qPCR rv	TCGCGGTTGGGTAGATAAAG	
<i>adrJ</i>	adrJ-qPCR fw	TGCGTGGACTGCTCTACTTC	82
	adrJ-qPCR rv	GAGTTCTGTGAGCGGGTCTA	
<i>adrK</i>	adrK-qPCR fw	TGACCCAAAGAGCCTCTACA	88
	adrK-qPCR rv	CGCTGGAATCTGCTGTATT	

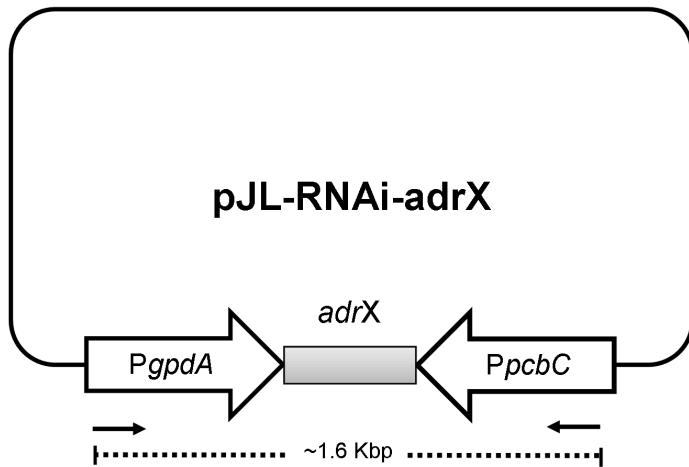
**A****B**

Figure S1. A) Example of the plasmid constructed for each gene from the *adr* cluster. *Pgpda*: promoter from the *gpd* gene from *A. nidulans*. *Ppcbc*: promoter from the *pcbc* gene from *P. chrysogenum*. Both promoters are oriented in opposite directions, generating double-stranded RNA molecules (dsRNAs) from the DNA fragment inserted, and thus triggering the fungal RNA-silencing machinery. The grey box between the promoters represents a fragment from a given *adr* gene (*adrX*, where X identifies the specific gene; see Supplementary Table S1). The small black arrows represent primers ConfRNAiFW (5'- GCATGCCATTAAACCTAGG-3') and ConfRNAiRV (5'- ACGGTGGCTGAAGATTC-3'). These primers were used to confirm the integration of the full silencing cassette (see panel B). The expected size of the amplicon containing the full silencing cassette (around 1.6 Kbp) is indicated. B) PCR assay demonstrating integration of the full silencing cassette in each transformant obtained. Amplicons obtained were subjected to electrophoresis in agarose gels. Lane WT: wild-type strain *P. roqueforti* CECT 2905; lane E: *P. roqueforti* CECT 2905 containing empty pJL43-RNAi vector; lane S: Standard GeneRuler 1 kb DNA Ladder (Fermentas). Relevant sizes expressed in kb are shown at left. Please note that as expected, length of amplicon obtained from *P. roqueforti* CECT 2905 containing empty vector is shorter (expected size 1.2 Kbp).