

LEGENDS TO SUPPLEMENTARY FIGURES

Fig. S1. Amplification of cDNAs by the primers designed on the basis of the EST information. The cDNA fragments for CURS (A) and DCS (B) were amplified.

Fig. S2. Multiple alignment of the amino acid sequences of plant type III PKSs. The red box indicates the three amino acids forming the Cys-His-Asn catalytic triad that is essential for starter substrate loading and malonyl-CoA condensation. DCS, diketide-CoA synthase; CURS, curcumin synthase; MsCHS, a chalcone synthase from *Medicago sativa* (DNA

accession no. AAA02824); RhBAS, a benzalacetone synthase from *Rheum palmatum* (AAK82824); WtPKS1, a polyketide synthase from *Wachendorfia thyrsiflora* (AAY727928).

Fig. S3. Expression of *DCS* (A) and *CURS* (B) in turmeric. The relative expression was quantified by qPCR and normalized to 18S rRNA.

Fig. S4. SDS-PAGE of purified DCS and CURS by Ni-NTA column. Lane 1, marker; lane 2, purified DCS and CURS.

Fig. S5. Analysis of the compound derived by hydrolysis of feruloyl-diketide-CoA and *trans*-5-(4-hydroxy-3-methoxyphenyl)-3-oxopent-4-enoic acid methyl ester. The alkaline hydrolysis of feruloyl-diketide-CoA [3b] (A) and *trans*-5-(4-hydroxy-3-methoxyphenyl)-3-oxopent-4-enoic acid methyl ester (B) resulted in the formation of the same compound, *trans*-5-(4-hydroxy-3-methoxyphenyl)-3-oxopent-4-enoic acid [3c]. *Trans*-5-(4-Hydroxy-3-methoxyphenyl)-3-oxopent-4-enoic acid [3c] is readily decarboxylated to yield dehydrozingerone [3d]. A methyl ester synthesized from *trans*-5-(4-hydroxy-3-methoxyphenyl)-3-oxopent-4-enoic acid, as a reference, was hydrolyzed by 1 M KOH just before LC-ESIMS analysis. The UV (C), MS (E), and MS/MS (G) spectra of the compound derived by hydrolysis of feruloyl-diketide-CoA are shown, together with the UV (D), MS (F), and MS/MS (H) spectra, of the hydrolysis product of the control methyl ester.

Fig. S6. HPLC analysis of the products after co-incubation of DCS and CURS in the presence of feruloyl-CoA [3a], *p*-coumaroyl-CoA [2a], and malonyl-CoA (A) and a similar HPLC analysis of an ethyl acetate extract of the rhizome of turmeric (B). Curcumin [3e], demethoxycurcumin [4e], and bisdemethoxycurcumin [2e] were detected in both analyses, although the abundance ratios of curcumin [3e] to demethoxycurcumin [4e] and bisdemethoxycurcumin [2e] were different.

Supplementary Table 1. Primers used for quantitative real time PCR

Gene	Primer sequence (5' to 3')
<i>DCS</i> (forward)	CAACAGCACGCCAGTCGA
<i>DCS</i> (reverse)	GTGCTGTTCATCCTGGACGAG
<i>CURS</i> (forward)	TCAGCTCATCCATCACGAAGTACAC
<i>CURS</i> (reverse)	CATCATTGACGCCATCGAACGC
18S rRNA (forward)	CCTTCCTCTAAATGATAAGGTTCAATGG
18S rRNA (reverse)	GATTGAATGGTCCGGTGAAAGTGT

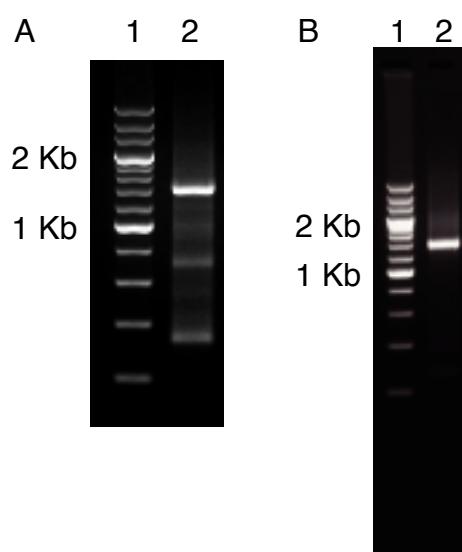


Fig. S1. Katsuyama et al.

		DCS	CURS	MsCHS	RhBAS	OscUS	WtPKS1
1	-----	MEANGYRITHS	-ADGPATILAIGTANPTNVVDQNAYPDFYFRVTNSEY				
1	-----	MANLHALRREQR	-AQGPATIMAIGTA TPPNLYEQSTFPDFYFRVTNSDD				
1	-----	MVSVSEIRKAQR	-AEGPATILAIGTANPANCVEQSTYPDFYFKITNSEH				
1	-----	MATEEMKK	-LATVMAIGTANPPNCYYQADFPDFYFRVTNSDH				
1	MAPTTTMGSALYPLGEMRRSQR	-ADGLAAVLAILGTANPPNCVTQEELPDFYFRVTNSDH					
1	-----	MASTEGLQAYRNNMAEGPATIMAIGTANPPNVVDASTFPDYYWRVTNSEH					
48	LQ	-ELKAKFRRICEKAAIRKRHLYLTEEILRENPSLLAPMAPSF	DARQAIIVVEAVPKLAK				
49	KQ	-ELKKKFRRMCEKT	MVKKRYLHLTEEILKERPKLCSYKEASFDDRDQDIVVE	EIPRLAK			
49	KT	-ELKEKFQRMCDKSMIKRRYMYL	TTEEILKENPNVCEYMAPSLDARQDMVVVEVPRLGK				
42	LI	-NLKQKFKRLCENSRIEKRYLHV	TEEILKENPNIAAYEATSLNVRHKMQVKGVAELEGK				
59	LT	-ALKDKFKRICQEMGVQRRYLIHTEEMLSAHPEF	VDRDAPSLDARLDIAADAVPELAA				
51	LSPEYRV	KLKRICERSSIRKRHLVLTEQLKENPT	LTTYVDASYDERQSIVLDAPVKLAC				
107	EAAEKA	IKIEWGRPKSDITHLVFC	SAGIDMPGSDLQLLKLLGLPPSVNRVMLYNVGCHAG				
108	EAAEKA	IKIEWGRPKSEITHLVFC	SISGIDMPGADYRLATLLGLPLTVNRLMIYSQACHMG				
108	EAAVKA	IKIEWGQPKSKITHLIVCT	TSGVDMPGADYQLTKLLGLRPYVKRYMMYQQGCCFAG				
101	EAAALK	AIKEWGQPKSKITHLIVC	LAGVDMPGADYQLTKLLDLPSPVKRFMFYHLGCYAG				
118	EAAKK	AIKEWGRPAADITHLVVT	TNSGAHVPGVDFRLVPLLGLRPSVRRTMLHLNGCFAG				
111	EAAA	AKIKEWGRPKTDITHMV	VCTGAGVDVPGVDYKMMNLLGLPPTVNRVMLYNVGCHAS				
167	GTA	LRAKDLAENNRGARVLAV	CSEVTVLSYRGPHPAHIESLFVQALFGDGAAALVVGSD				
168	AAML	RIAKDLAENNRGARVLVV	ACEITVLSFRGPNEGDFEALAGQAGFDGAGAVVVGAD				
168	GT	VLRAKDLAENNKGARVL	VVCSEVTAVTFRGPSDTHLDLSVQALFGDGAAALIVGSD				
161	GT	VLRAKDIENNGARVL	IVCSEM TTTCFRGPSETHLDMSIGQAI LGDAAVIVGAD				
178	CA	ALRLAKDLAENSRGARVL	VVAE LTLMYFTGPDEGCFR LLVQGLFGDAAVIVGAD				
171	GT	VLRIAKDLAENNKGARVL	VVSSEVSVMF FRGPAEGDVEILLQALFGDGSAAIIVGAD				
227	PVD	GVERPIFEIASASQVML	PESAEAVGGHLREIGLTFHLKSQLPSIIASNIEQSLTTAC				
228	PLE	GIKEKPIYEIAAMQETVA	EISQGAVGGHLRAFGWTFYFLNQLPAIIADNLGRSLERAL				
228	PV	PEIEKPIFEMVWT	TAQTIAPDSEGAI DGH LREAGLTFHLLKDVP GIVSKNITKALVEAF				
221	PDL	TVERPIFELVSTAQTIV	PESHGAI EGH LLESGLSFHLYKTVPTLISNNIKTCLSDAF				
238	-ADD	VERPLFEIVSAQTII	IPESDH ALNMRF TERR LDGV LGRQV PGLIGDNVERCLLDMF				
231	PIE	GVEKPIFQIFSASQM	TLP EGEHLVAGHLRELGLTFHLKPQLPNTVSSNIHKPLKKA F				
287	SPL	GLLSDWNQLFWAVHPGG	RAILDQVEARLGLEKDR LAATRHVLSEYGNMQSATVL				
288	APL	GVREWNDVFWVAHPGN	WAI IDAI EAKLQLSPDKLSTARHVTEYGNMQSATVY				
288	EPL	GISD YNSIFWIAHPGG	PAILDQVEQKLALKPEKMNATREV LSEYGNMSSACVL				
281	TPL	NISD WNSLFWIAHPGG	PAILDQV TAKV GLEKEKLKVTRQVLKDYGNMSSATVF				
297	GPL	LLGGDGGGGWNDLFWA	HVGPGS STIMD QVDA ALGLEPGKLA ASRRV LSDYGNMMSGATVI				
291	EPL	NITD WNSIFWIVHPGG	RAILDQVQE KIGLEENKLDV SRYVLAENG NMM SASVF				
343	FIL	DEM RNRSAAE GHATT	GEGLDWGVLLGF GPG GLS IETVVLHSCRLN	-			
344	FVM	DEL RKR SAVE GR	STTGDGLQWGVL LGFGP GLS IETVVL RSMPL	--			
344	FIL	DEM RKKSTQNGLK	T GEGLEWGVL FGFGP GLT IETVVL RSV A I	--			
337	FIM	DEM RKKSL ENQ	ATT GEGLEWGVL FGFGP GIT VETVVL RSV PVI S				
357	FAL	DEL RRQRKE	AAA AGEWPELGV MMAFGPGMTVDAM LLHATSHVN				
347	FIM	DEM RKR SAAQGC	STT GEGHEWGVL FGFGP GLS IETVVL HSVPLSI				

Fig. S2. Katsuyama et al.

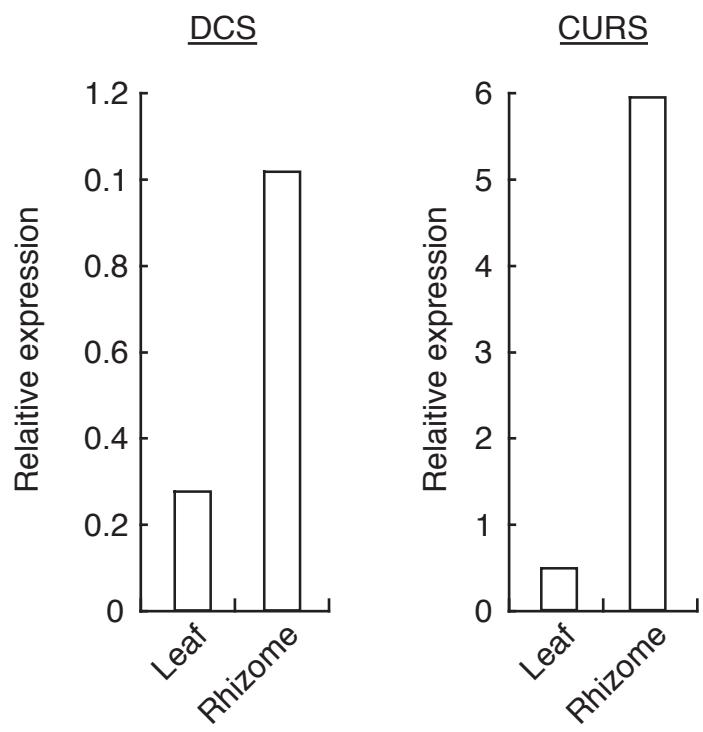


Fig. S3. Katsuyama et al.

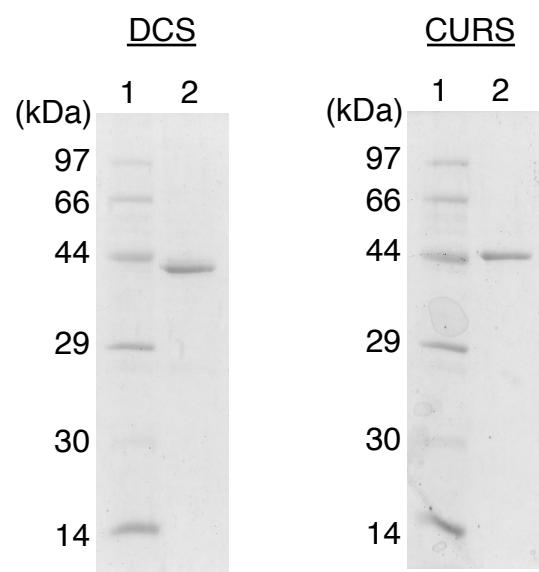


Fig. S4 Katsuyama et al.

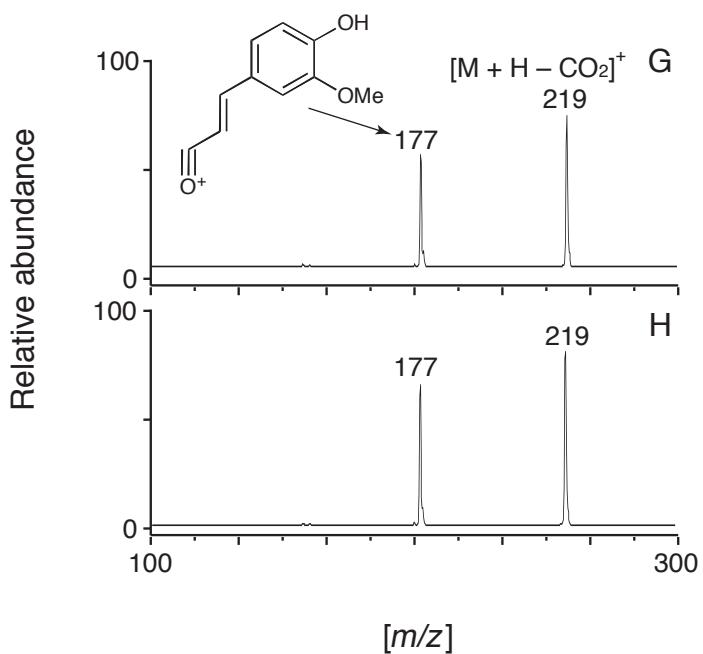
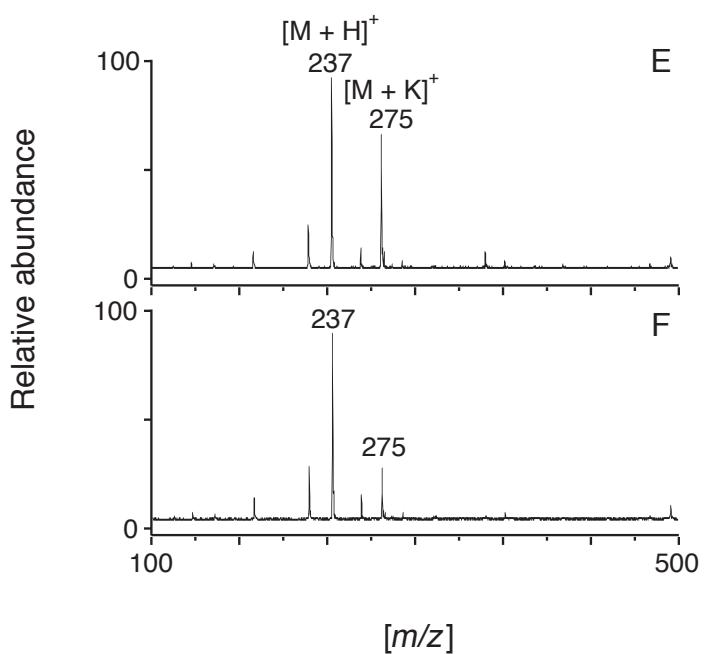
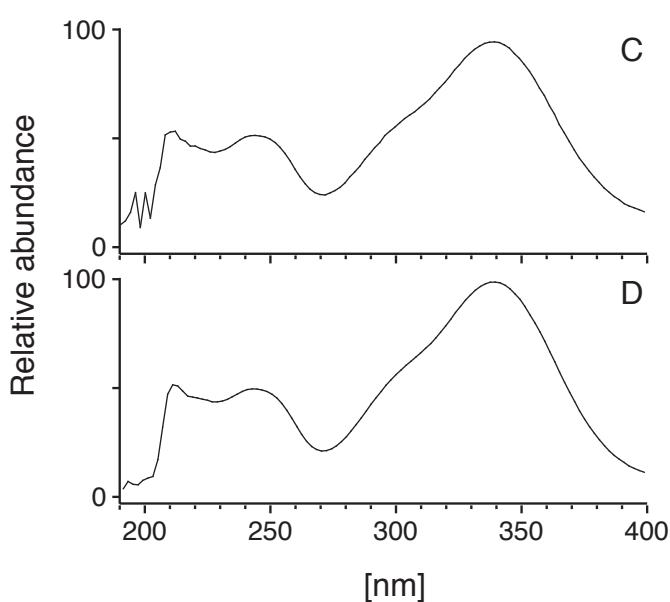
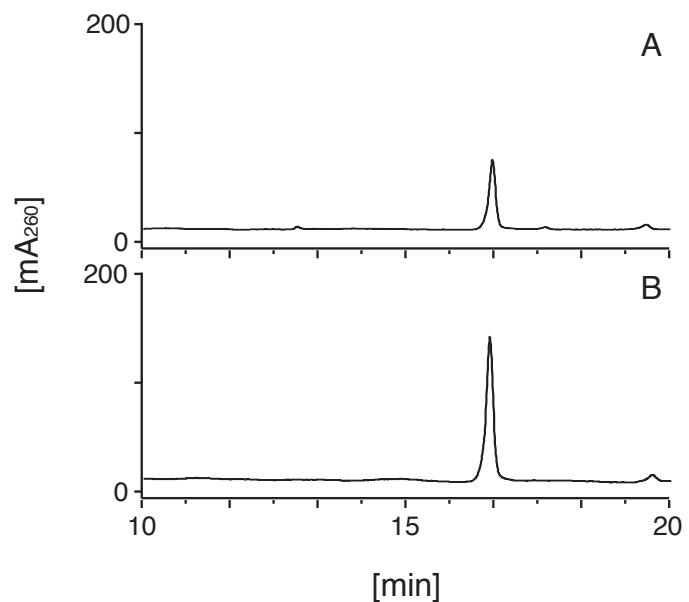


Fig. S5. Katsuyama et al.

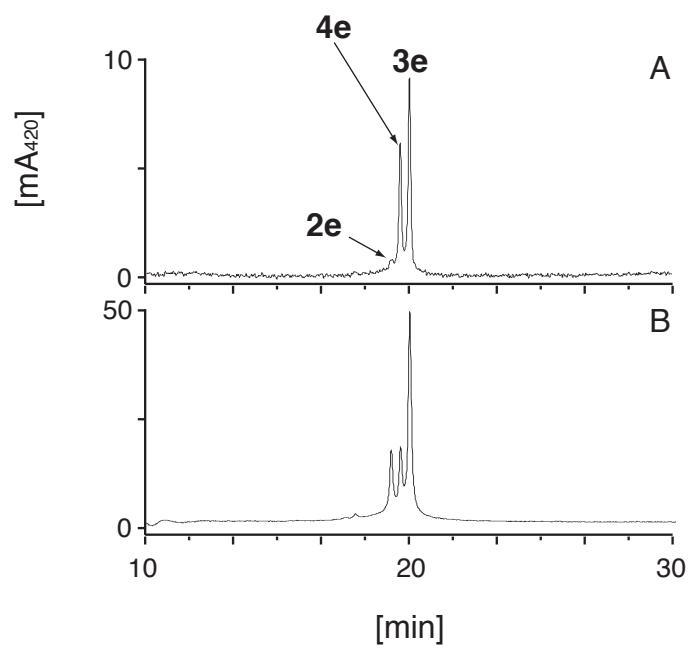


Fig. S6. Katsuyama et al.