

# **A role in 15-deacetylcalonecetrin acetylation in the non-enzymatic cyclization of an earlier bicyclic intermediate in *Fusarium trichothecene* biosynthesis**

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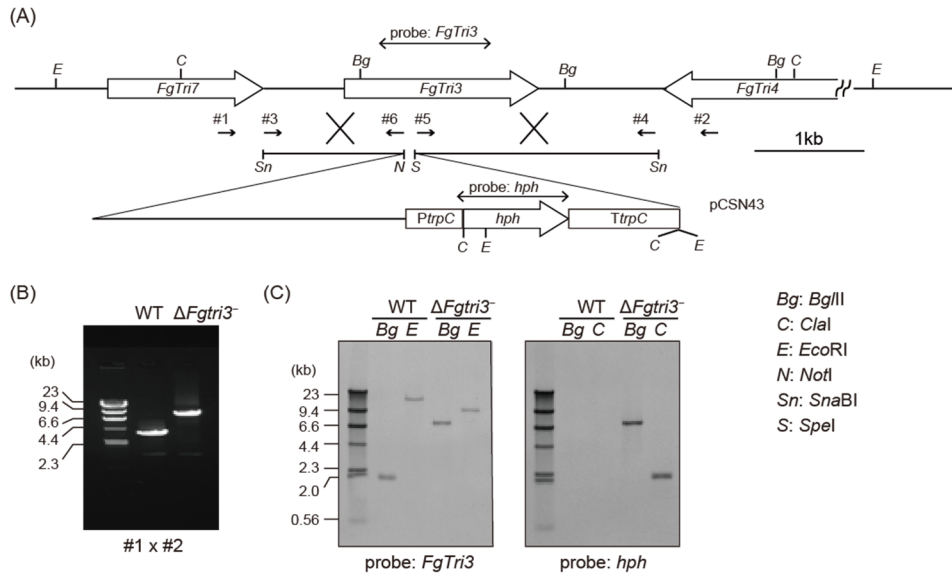
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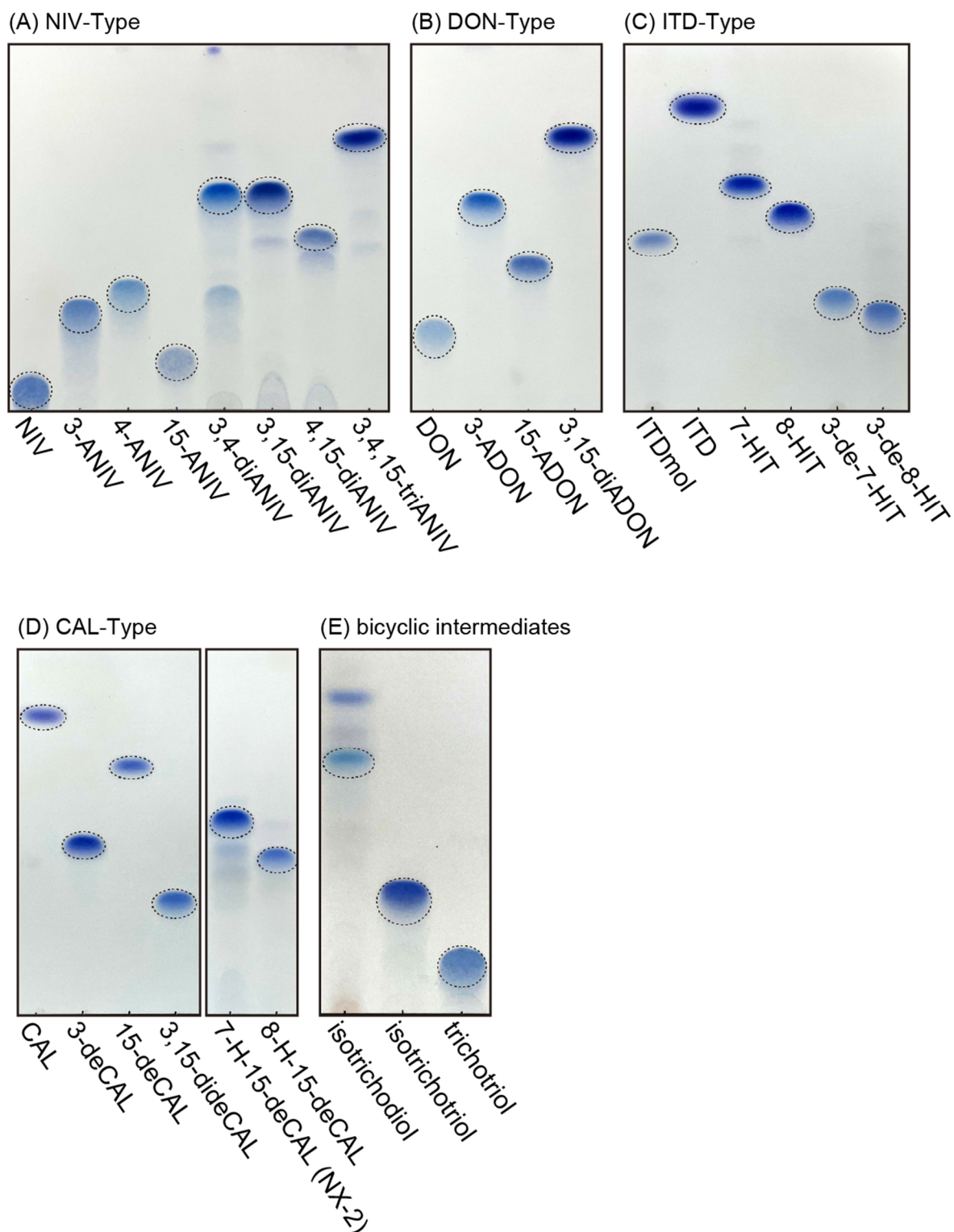
**Table S1.** Primers used for generation of the  $\Delta Fgtri3$  (FGD3) and  $\Delta Fgtri5\Delta Fgtri11$  (FGD5/11) mutants

No.	Primer name	Primer sequence	Comments
#1	MFTri3-LAU	5'-TATCTTTACCACAACAGCCGATTGCCCC-3'	Primer for screening of the <i>FgTri3</i> gene disruption mutant (used in combination with #2)
#2	MFTri3-LAD	5'-GCACTCTCTGATTGGAGGAGAACACTTG-3'	Primer for screening of the <i>FgTri3</i> gene disruption mutant (used in combination with #1)
#3	MFTri3U-Sna	5'-GT <u>ACGTAG</u> TCATTGTCTCCATTATAACAAGGC-3'	Inward primer containing a <i>Sna</i> BI site (underlined) used for the construction of the <i>FgTri3</i> gene disruption vector (used in combination with #4)
#4	MFTri3D-Sna	5'-TTT <u>ACGTAT</u> TGCAGGAATGGAAGTTCATTGTAG-3'	Inward primer containing a <i>Sna</i> BI site (underlined) used for the construction of the <i>FgTri3</i> gene disruption vector (used in combination with #3)
#5	MFTri3U-inv/Spe	5'-A <u>ACTAGT</u> ATGGGGCCAAGAGATCAAGAACCTG-3'	Outward primer containing a <i>Spe</i> I site (underlined) used for the construction of the <i>FgTri3</i> gene disruption vector (used in combination with #6)
#6	MFTri3D-inv/Not	5'-A <u>GCGGCCGCT</u> TGAGGGATTGGCGTCGAGTCTG-3'	Outward primer containing a <i>Not</i> I site (underlined) used for the construction of the <i>FgTri3</i> gene disruption vector (used in combination with #5)
#7	MFTri11-LAU2	5'-TCAAGGCTCTCCCTGGGATG-3'	Primer for PCR confirmation of <i>FgTri11</i> gene disruption (used in combination with #8)
#8	MFTri11-LAD	5'- ATGAGTCGATCAATTTAGCGGCTGCGAG-3'	Primer for PCR confirmation of <i>FgTri11</i> gene disruption (used in combination with #7)





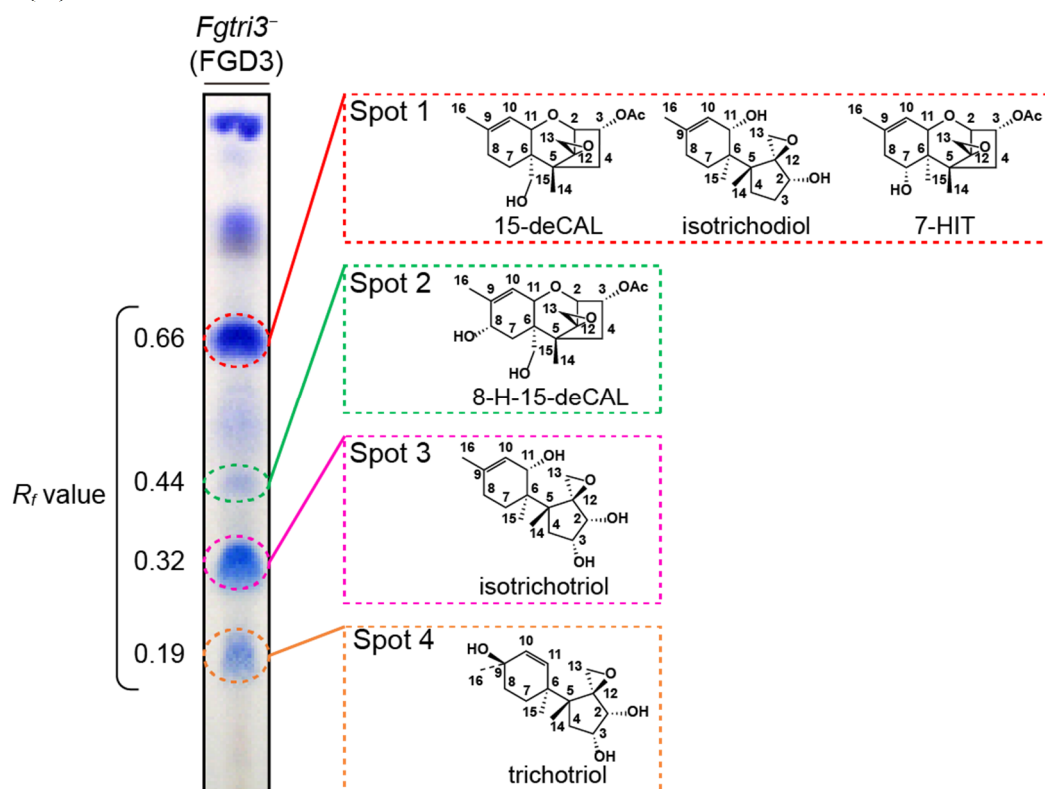
**Supplementary Figure S1.** Generation of a  $\Delta Fgtri3$  mutant used in this study. (A) The gene disruption vector p $\Delta FgTri3$ -hph was constructed by replacing a portion of an *FgTri3* coding region with pCSN43 [1] that contains the *hph* (hygromycin B phosphotransferase gene) cassette by the inverse polymerase chain reaction (IPCR) method [2] as follows: (1) *FgTri3* and its flanking regions were amplified by long PCR with inward primers #3 and #4 (Table S1), both of which had a *Sna*BI recognition site (Table S1); (2) the amplified product was self-ligated after digestion with *Sna*BI; (3) the ligated flanking regions were amplified by IPCR with the outward primers #5 and #6 (Table S1) that had *Spe*I and *Not*I sites, respectively (Table S1); and (4) the IPCR product was cloned between *Spe*I and *Not*I sites downstream of the *trpC* terminator. After transformation, the colonies were selected with 300  $\mu$ g/ml of hygromycin B. (B) PCR analysis of genomic DNA isolated from the wild-type (WT) and  $\Delta Fgtri3$  mutant strains. Two primers #1 and #2 that amplify the region encompassing *FgTri3* were used. An amplicon from the disruption mutant is 4.4 kb larger than the one from the WT strain. (C) Southern blot analysis of genomic DNA isolated from the WT and  $\Delta Fgtri3$  mutant strains. The probes were synthesized with a PCR DIG probe synthesis kit [3] and detected with a DIG Luminescent Detection kit (Roche Diagnostics GmbH, Mannheim); standard hybridization, washing, and detection conditions provided by the manufacture were used for detection. Either blot showed expected banding patterns with probes that correspond to *FgTri3* and *hph*.



**Supplementary Figure S2.** Trichothecene standards on TLC. (A) NIV-type trichothecenes. NIV was produced by the deacetylation of 4-acetylivalenol (4-ANIV). 4-ANIV and 4,15-diacetylivalenol (4,15-diANIV) were produced through the incubation of MAFF 111233 WT culture. 3-Acetylivalenol (3-ANIV) was produced through the incubation of the culture of

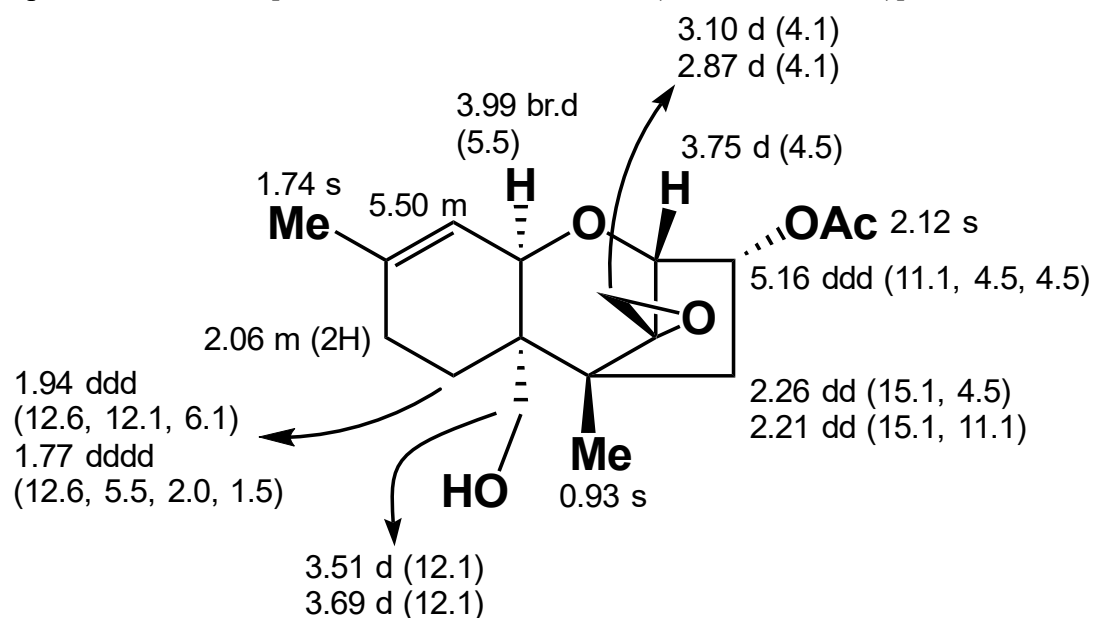
NBRC 113183, an *Fgtri13*\_o/e strain of *F. graminearum* MAFF 101551 [4]. 15-Acetylnivalenol (15-ANIV) and 3,15-diacetylnivalenol (3,15-diANIV) were produced through the incubation of the culture of NBRC 113182, an *Fgtri7*<sup>-</sup> #1 s1\* strain of *F. graminearum* MAFF 111233 [4]. 3,4,15-Triacetylnivalenol (3,4,15-triANIV) was produced through the incubation of the culture of NBRC 114123, an *FgTri8* disruptant of *F. graminearum* MAFF 111233 [5]. 3,4-Diacetylnivalenol (3,4-diANIV) was obtained by 3-*O*-acetylation of 4-ANIV by Tri101p. (B) DON-type trichothecenes. 3,15-diADON was produced through the incubation of the culture of YN\_149. (C) ITD-type trichothecenes. 3-Deacetyl-7-hydroxyisotrichodermin (3-de-7-HIT) and 3-deacetyl-8-hydroxyisotrichodermin (3-de-8-HIT) were obtained through the alkaline deacetylation of 7-HIT and 8-HIT, respectively. (D) CAL-type trichothecenes. 3,15-dideCAL is the abbreviation for 3,15-dideacetylcalonectrin. 7-H-15-deCAL was obtained through the incubation of the culture of *F. graminearum*  $\Delta Fgtri1/FgTri1_{NX}$  chemotype [6]. Another name for 7-H-15-deCAL is NX-2. 8-H-15-deCAL was obtained through the incubation of FGD3 culture, and its structure was confirmed by NMR analysis (Supplementary Figure S3, spot 3). (E) Bicyclic trichothecene intermediates. Isotrichodiol, isotrichotriol and trichotriol were produced through the incubation of FGD3 culture, and their structures were confirmed by NMR analysis (Supplementary Figure S3, spots 1, 3, and 4).

(A)



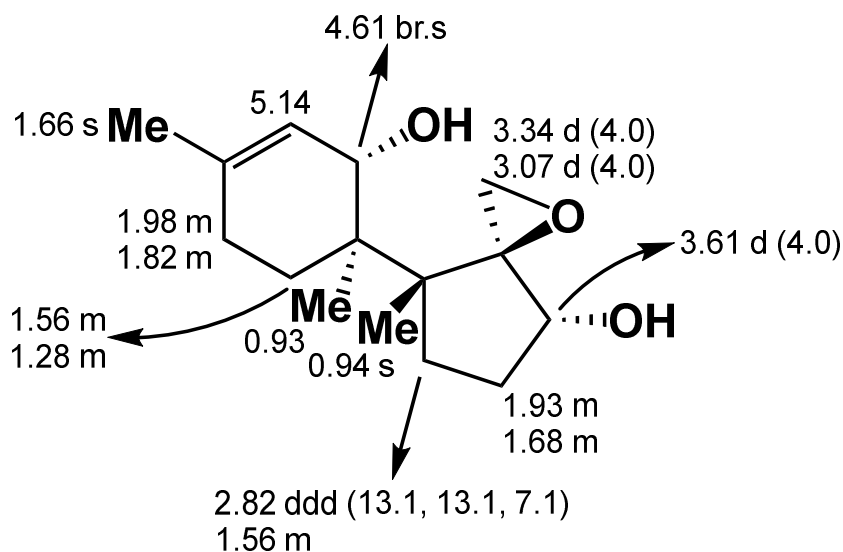
(B)

Spot 1 : 15-deCAL [ $^1\text{H}$  NMR Data of 15-deCAL (600 MHz,  $\text{CDCl}_3$ )]



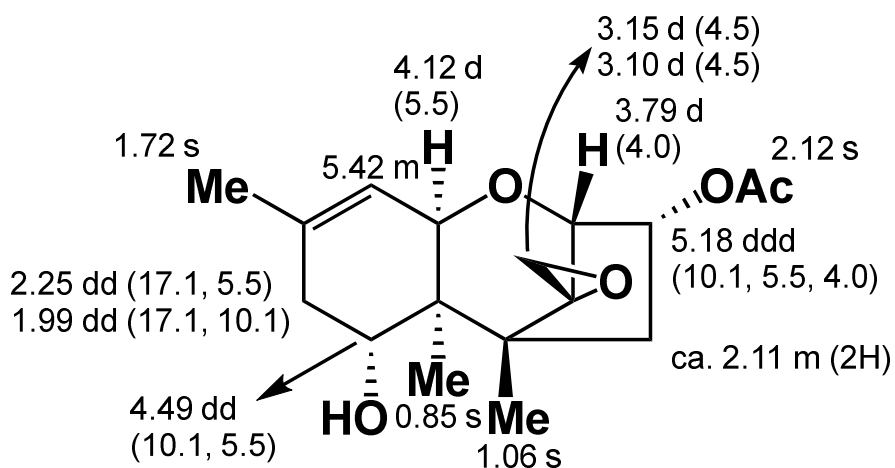
$^{13}\text{C}$  NMR Data of 15-deCAL (151 MHz,  $\text{CDCl}_3$ ) 78.1 (C-2), 71.4 (C-3), 39.3 (C-4), 45.3 (C-5), 44.2 (C-6), 20.8 (C-7), 28.4 (C-8), 140.5 (C-9), 119.3 (C-10), 68.3 (C-11), 65.2 (C-12), 48.6 (C-13), 12.5 (C-14), 62.8 (C-15), 23.3 (C-16), 21.0, and 170.5 (3-OAc) ppm.

Spot 1 : isotrichodiol [ $^1\text{H}$  NMR Data of isotrichodiol (600 MHz,  $\text{CDCl}_3$ )]



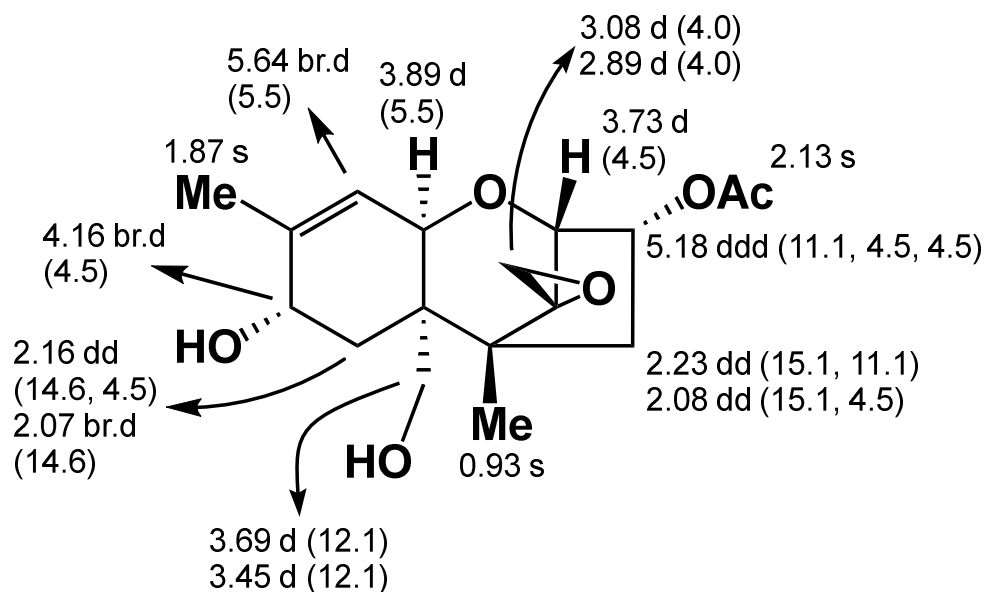
$^{13}\text{C}$  NMR Data of isotrichodiol (151 MHz,  $\text{CDCl}_3$ ) 81.7 (C-2), 30.3 (C-3), 37.1 (C-4), 46.8 (C-5), 41.0 (C-6), 29.2 (C-7), 27.8 (C-8), 130.5 (C-9), 125.5 (C-10), 72.0 (C-11), 69.3 (C-12), 49.5 (C-13), 20.2 (C-14), 13.0 (C-15), and 22.3 (C-16) ppm.

Spot 1 : 7-HIT [ $^1\text{H}$  NMR Data of 7-HIT (600 MHz,  $\text{CDCl}_3$ )]



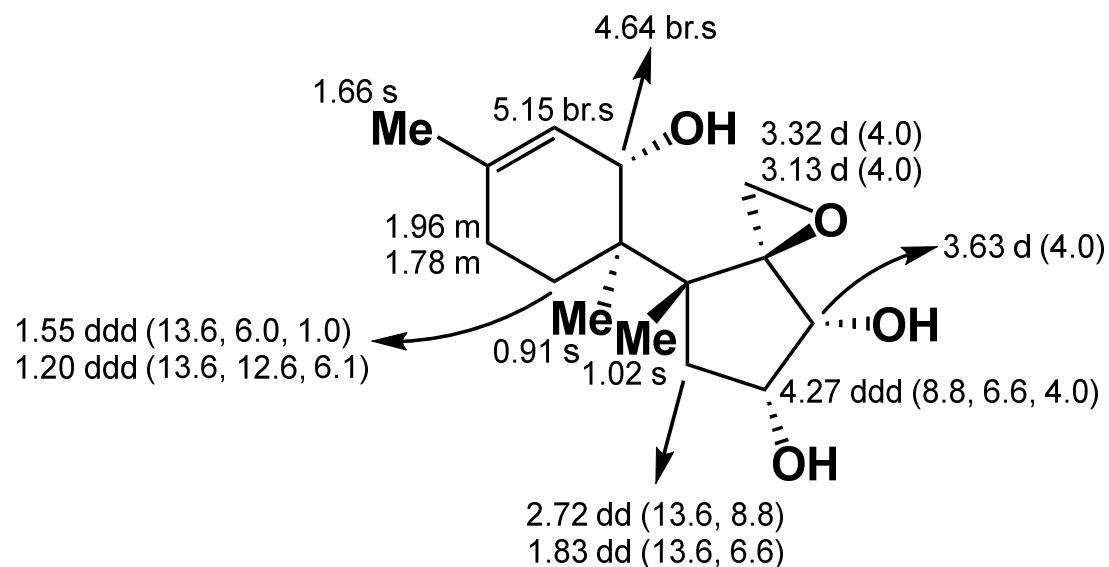
$^{13}\text{C}$  NMR Data of 7-HIT (151 MHz,  $\text{CDCl}_3$ ) 79.0 (C-2), 71.4 (C-3), 41.5 (C-4), 46.2 (C-5), 44.7 (C-6), 69.1 (C-7), 39.6 (C-8), 137.9 (C-9), 119.5 (C-10), 74.7 (C-11), 65.0 (C-12), 47.7 (C-13), 15.1 (C-14), 10.3 (C-15), and 22.5 (C-16) ppm.

Spot 2 : 8-H-15-deCAL [ $^1\text{H}$  NMR Data of 8-H-15-deCAL (600 MHz,  $\text{CDCl}_3$ )]



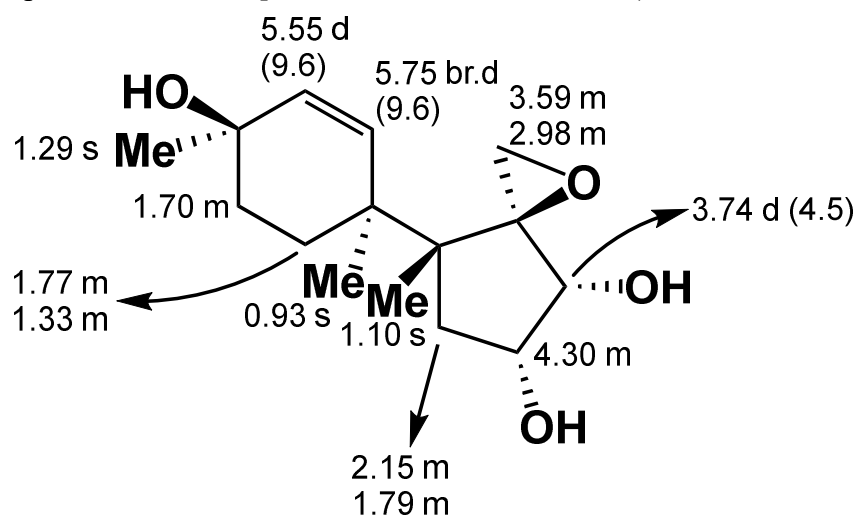
$^{13}\text{C}$  NMR Data of 8-H 15-deCAL (151 MHz,  $\text{CDCl}_3$ ) 77.9 (C-2), 71.3 (C-3), 39.5 (C-4), 45.7 (C-5), 44.0 (C-6), 28.3 (C-7), 67.3 (C-8), 138.8 (C-9), 122.8 (C-10), 68.6 (C-11), 67.0 (C-12), 48.5 (C-13), 12.6 (C-14), 62.7 (C-15), 20.6 (C-16), 21.1, and 170.7 (3-OAc) ppm.

Spot 3 : isotrichotriol [ $^1\text{H}$  NMR Data of isotrichotriol (600 MHz,  $\text{CDCl}_3$ )]



$^{13}\text{C}$  NMR Data of isotrichotriol (151 MHz,  $\text{CDCl}_3$ ) 79.6 (C-2), 70.0 (C-3), 43.5 (C-4), 45.2 (C-5), 41.2 (C-6), 29.1 (C-7), 27.6 (C-8), 135.3 (C-9), 125.3 (C-10), 71.5 (C-11), 67.5 (C-12), 49.3 (C-13), 22.0 (C-14), 13.1 (C-15), and 22.5 (C-16) ppm.

Spot 4 : trichotriol [ $^1\text{H}$  NMR Data of trichotriol (600 MHz,  $\text{CDCl}_3$ )]

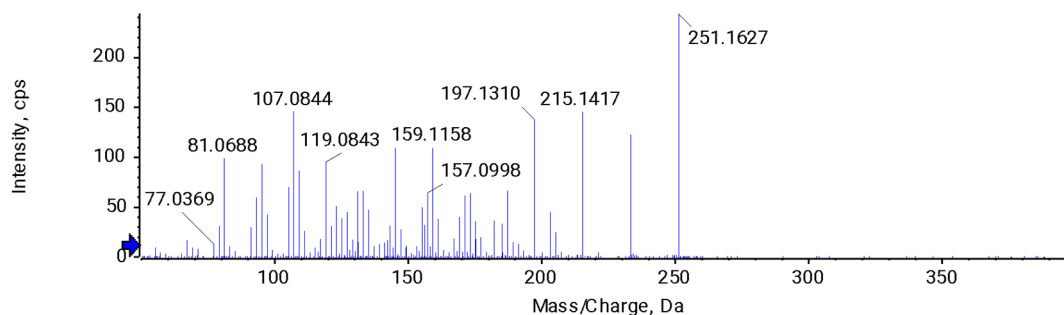


$^{13}\text{C}$  NMR Data of trichotriol (151 MHz,  $\text{CDCl}_3$ ) 77.5 (C-2), 69.0 (C-3), 41.5 (C-4), 43.9 (C-5), 39.5 (C-6), 27.0 (C-7), 34.0 (C-8), 66.5 (C-9), 131.3 (C-10), 136.3 (C-11), 69.2 (C-12), 49.9 (C-13), 20.4 (C-14), 21.3 (C-15), and 30.7 (C-16) ppm.

**Supplementary Figure S3.** Bicyclic precursors and tricyclic trichothecene intermediates of FGD3. (A) TLC analysis of trichothecene metabolites. Spot 1 contained 15-deCAL, isotrichodiol, and 7-HIT. Spots 2, 3, and 4 contained 8-H-15-deCAL, isotrichotriol, and trichotriol, respectively. (B) The compounds in spots 1–4 were confirmed by NMR analyses.

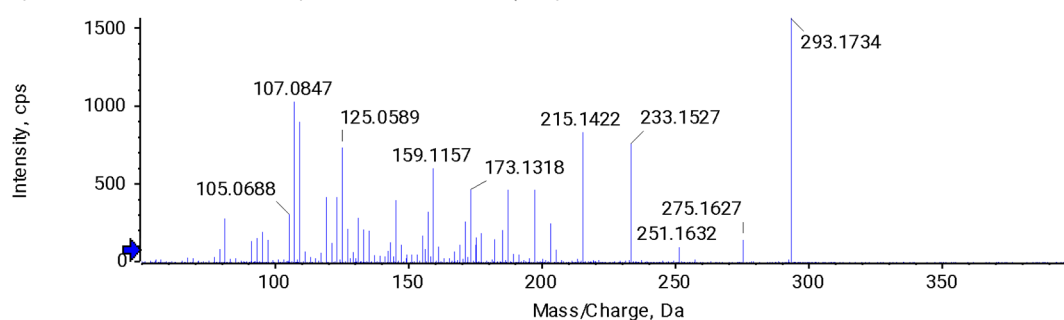
(A) ITDmol  $[M+H]^+$   $m/z$ : 251.164, RT: 3.88 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET12.wiff (samp...from 3.835 to 3.972 min Precursor: 251.2 Da, CE: 25.0)



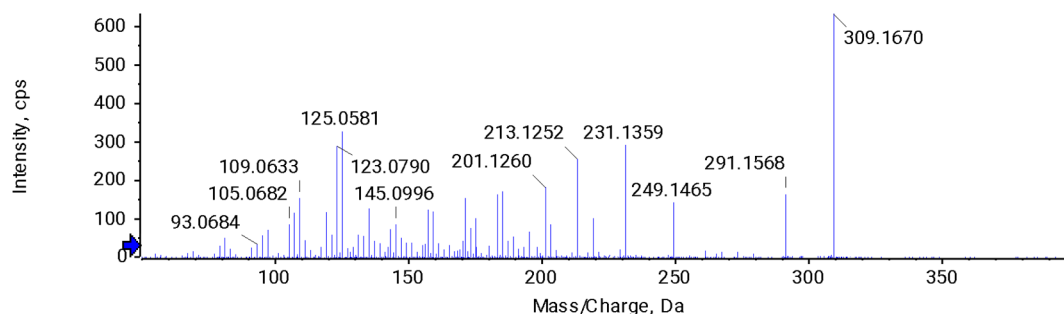
(B) ITD  $[M+H]^+$   $m/z$ : 293.175, RT: 4.95 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET13.wiff (samp...from 4.881 to 5.018 min Precursor: 293.2 Da, CE: 25.0)



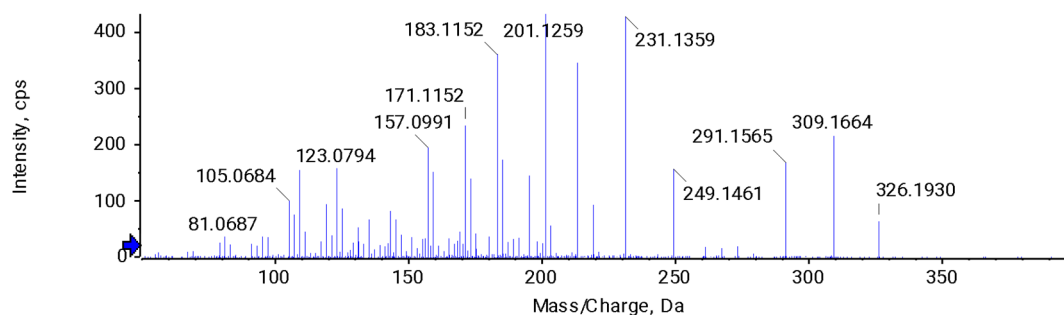
(C-1) 7-HIT  $[M+H]^+$   $m/z$ : 309.170, RT: 3.84 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET15.wiff (samp...from 3.787 to 3.899 min Precursor: 309.2 Da, CE: 25.0)



(C-2) 7-HIT  $[M+NH_4]^+$   $m/z$ : 326.196, RT: 3.84 min

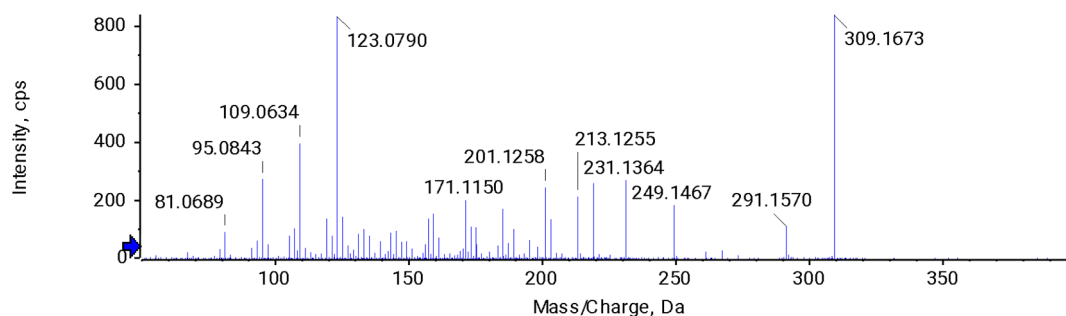
Spectrum from 231224\_KY\_TOFposi\_4\_DataSET15.wiff (samp...from 3.788 to 3.908 min Precursor: 326.2 Da, CE: 25.0)





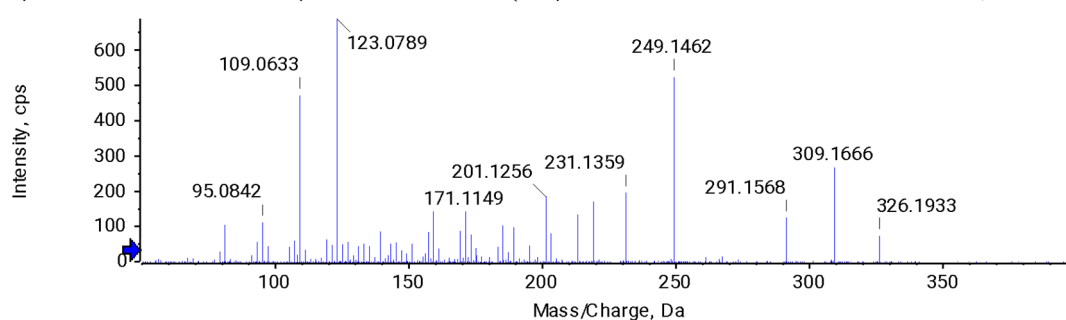
(D-1) 8-HIT  $[M+H]^+$   $m/z$ : 309.170, RT: 3.45 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET16.wiff (samp...from 3.394 to 3.505 min Precursor: 309.2 Da, CE: 25.0



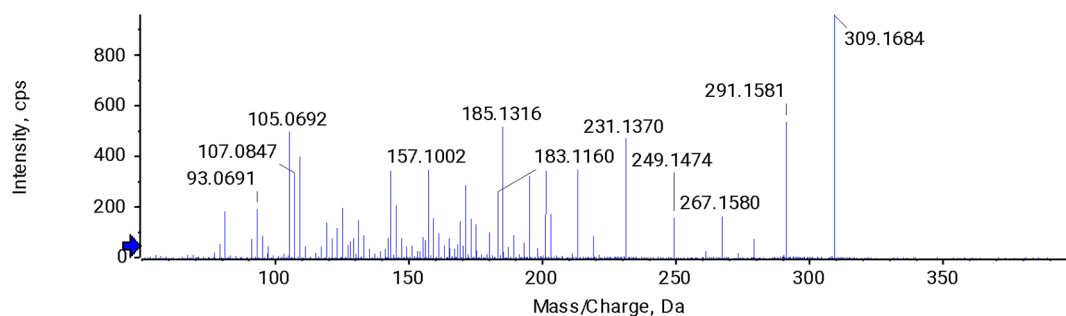
(D-2) 8-HIT  $[M+NH_4]^+$   $m/z$ : 326.196, RT: 3.44 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET16.wiff (samp...from 3.403 to 3.506 min Precursor: 326.2 Da, CE: 25.0



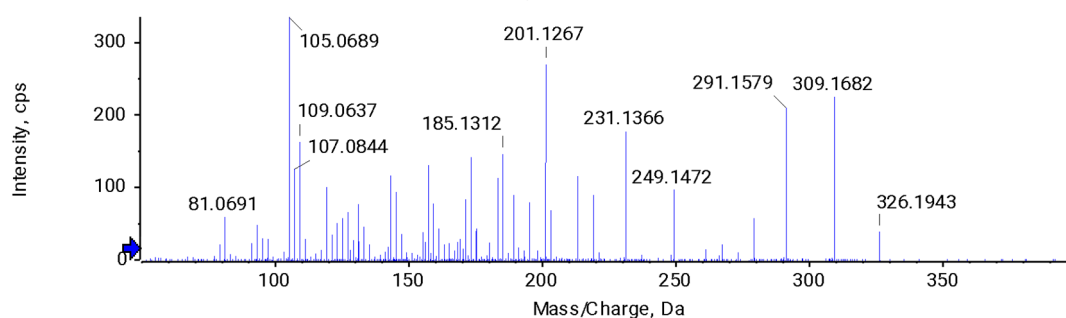
(E-1) 15-deCAL  $[M+H]^+$   $m/z$ : 309.170, RT: 3.55 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET112.wiff (sam...from 3.497 to 3.616 min Precursor: 309.2 Da, CE: 25.0



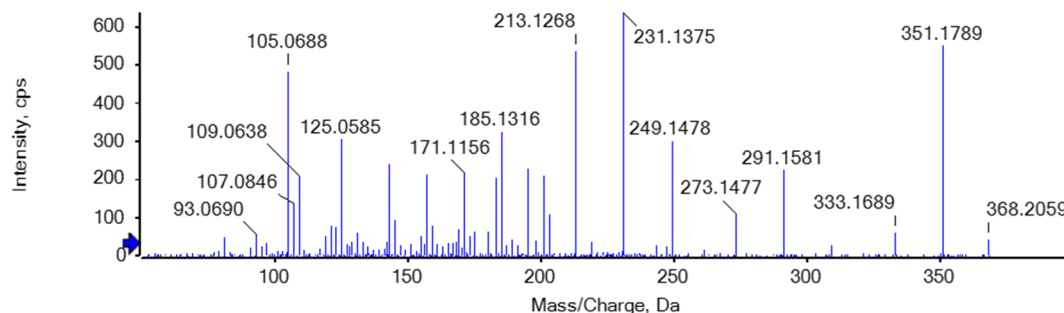
(E-2) 15-deCAL  $[M+NH_4]^+$   $m/z$ : 326.196, RT: 3.54 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET112.wiff (sam...from 3.497 to 3.626 min Precursor: 326.2 Da, CE: 25.0



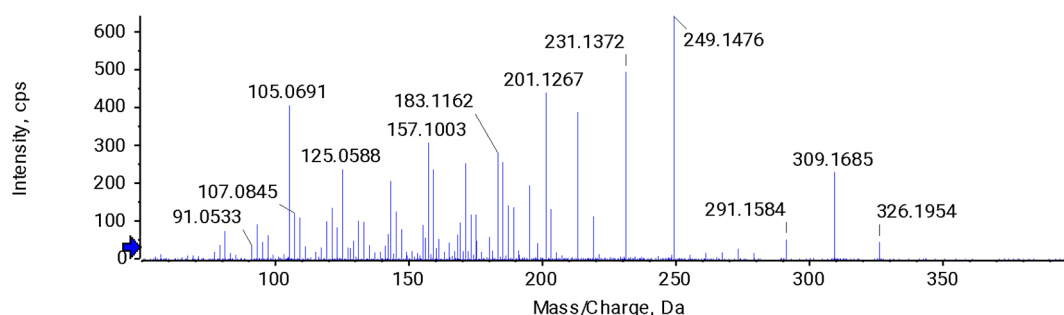
(F) CAL  $[M+NH_4]^+$   $m/z$ : 368.207, RT: 4.41 min

Spectrum from 231224\_KY\_TOFposi\_3\_DataSET17.wiff (samp...from 4.363 to 4.466 min Precursor: 368.2 Da, CE: 25.0)



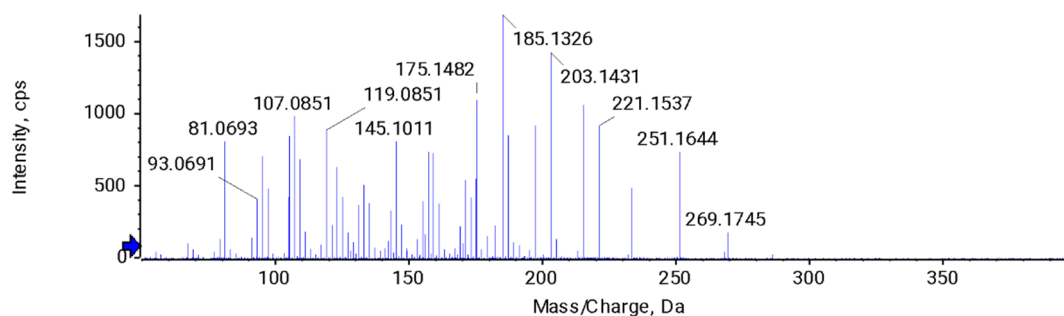
(G) 3-deCAL  $[M+NH_4]^+$   $m/z$ : 326.196, RT: 3.58 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET111.wiff (sam...from 3.523 to 3.643 min Precursor: 326.2 Da, CE: 25.0)



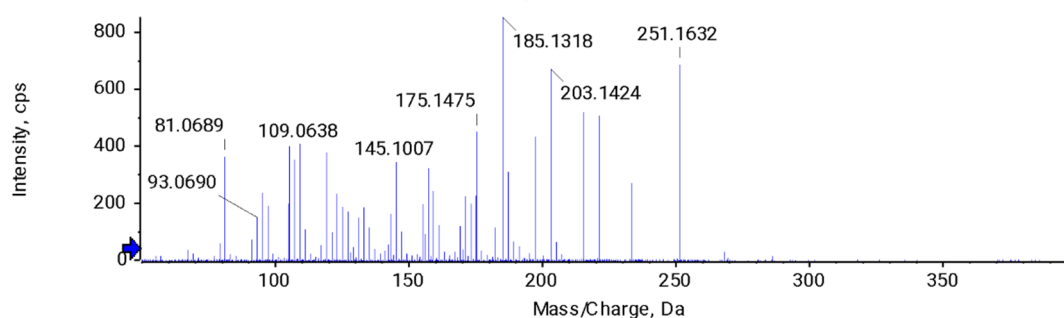
(H) isotrichotriol  $[M+NH_4]^+$   $m/z$ : 286.201, RT: 3.45 min

Spectrum from 230614\_KY\_TOFposi\_3\_DataSET117.wiff (sam...from 3.380 to 3.563 min Precursor: 286.2 Da, CE: 25.0)



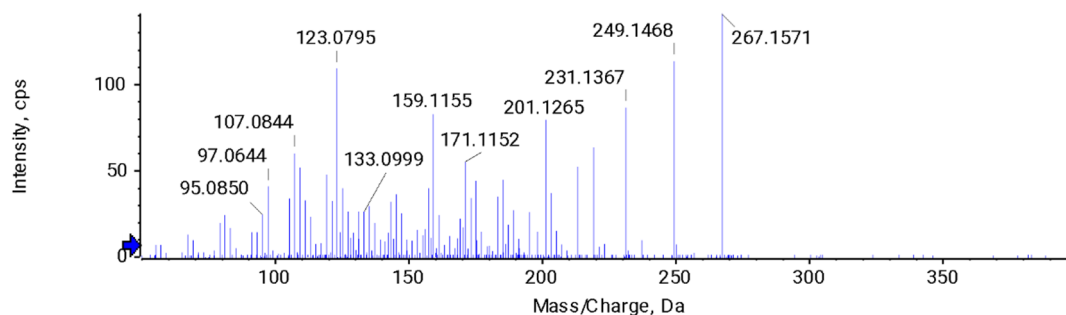
(I) trichotriol  $[M+NH_4]^+$   $m/z$ : 286.201, RT: 3.04 min

Spectrum from 230614\_KY\_TOFposi\_3\_DataSET120.wiff (sam...from 2.979 to 3.149 min Precursor: 286.2 Da, CE: 25.0)



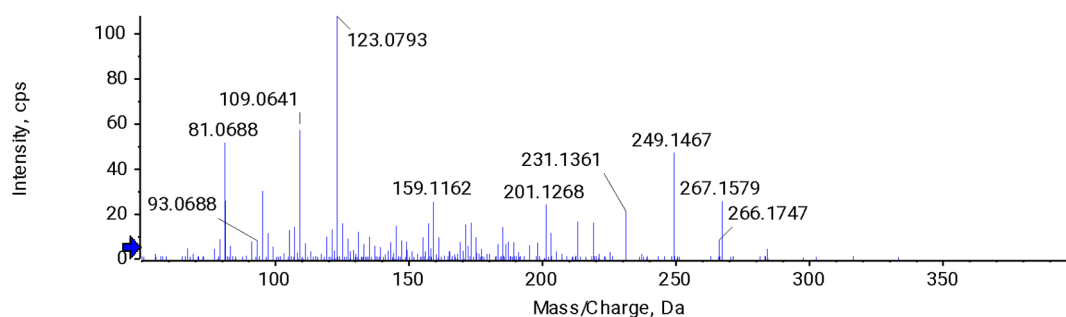
(J) 3-deacetyl-7-hydroxyisotrichodermin (3-de-7-HIT)  $[M+H]^+$   $m/z$ : 267.159, RT: 2.80 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET18.wiff (samp...from 2.745 to 2.865 min Precursor: 267.2 Da, CE: 25.0)



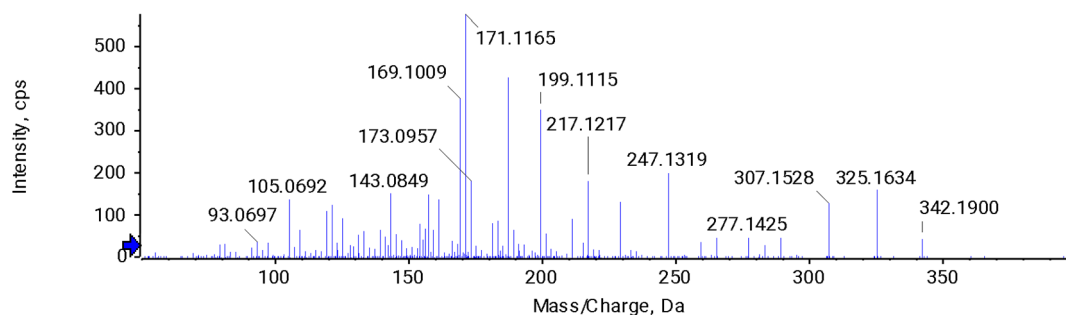
(K) 3-deacetyl-8-hydroxyisotrichodermin (3-de-8-HIT)  $[M+NH_4]^+$   $m/z$ : 284.186, RT: 2.48 min

Spectrum from 231224\_KY\_TOFposi\_4\_DataSET19.wiff (samp...from 2.412 to 2.541 min Precursor: 284.2 Da, CE: 25.0)



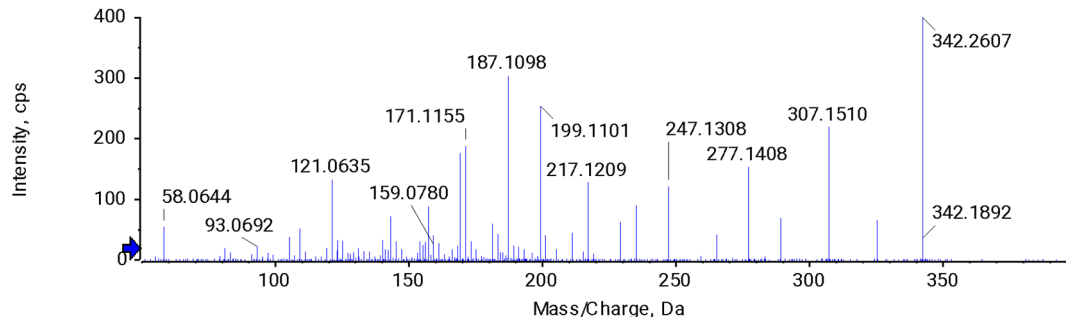
(L) 7-H-15-deCAL (NX-2)  $[M+NH_4]^+$   $m/z$ : 342.191, RT: 3.15 min

Spectrum from 230812\_KY\_TOFposi\_2\_DataSET16.wiff (samp...from 3.118 to 3.222 min Precursor: 342.2 Da, CE: 25.0)



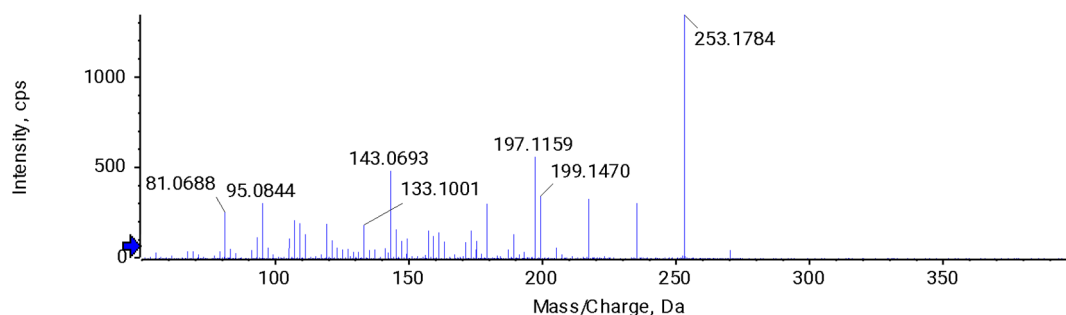
(M) 8-H-15-deCAL  $[M+NH_4]^+$   $m/z$ : 342.191, RT: 3.10 min

Spectrum from 231224\_KY\_TOFposi\_5\_DataSET19.wiff (samp...from 3.035 to 3.155 min Precursor: 342.2 Da, CE: 25.0)



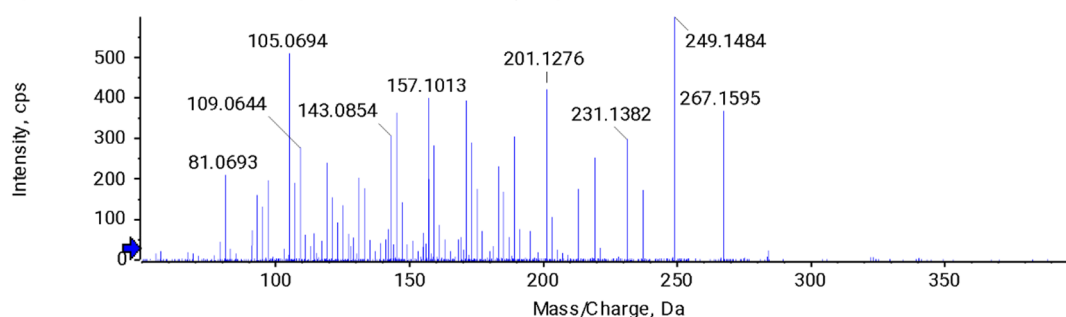
(N) isotrichodiol  $[M+NH_4]^+$   $m/z$ : 270.206, RT: 4.11 min

Spectrum from 231224\_KY\_TOFposi\_5\_DataSET15.wiff (samp...from 4.043 to 4.189 min Precursor: 270.2 Da, CE: 25.0)



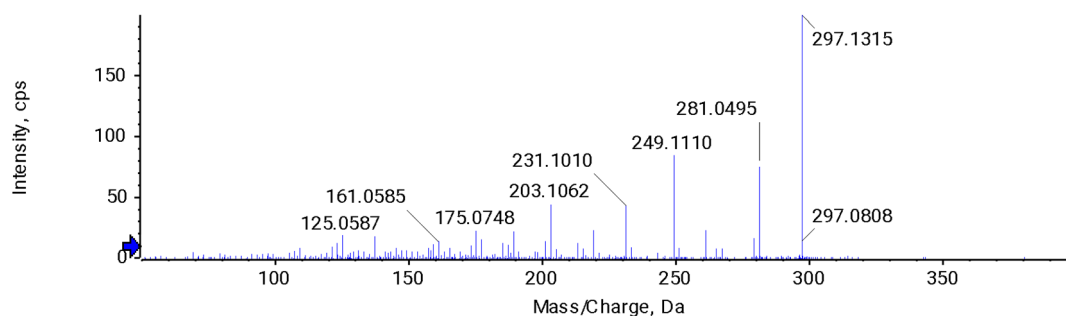
(O) 3,15-dideacetylcalonecitrin (3,15-dideCAL)  $[M+NH_4]^+$   $m/z$ : 284.186, RT: 2.77 min

Spectrum from 230626\_KY\_TOFposi\_1\_DataSET12.wiff (samp...from 2.713 to 2.822 min Precursor: 284.2 Da, CE: 25.0)



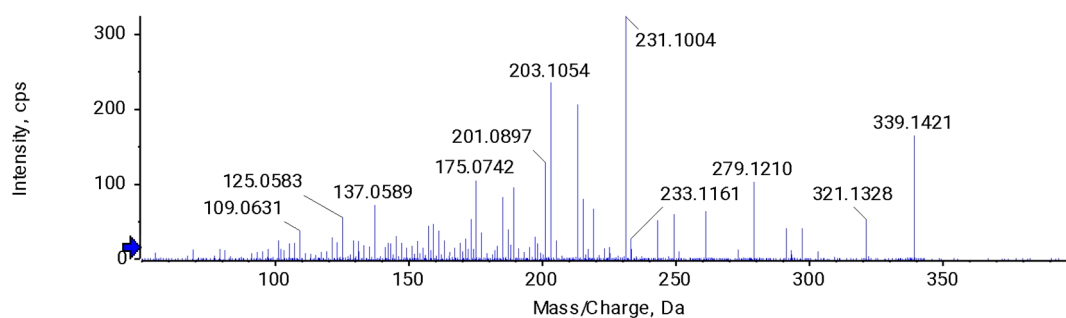
(P) DON  $[M+NH_4]^+$   $m/z$ : 314.160, RT: 2.29 min

Spectrum from 231107\_KY\_TOFposi\_2\_DataSET12.wiff (samp...from 2.258 to 2.389 min Precursor: 314.2 Da, CE: 25.0)



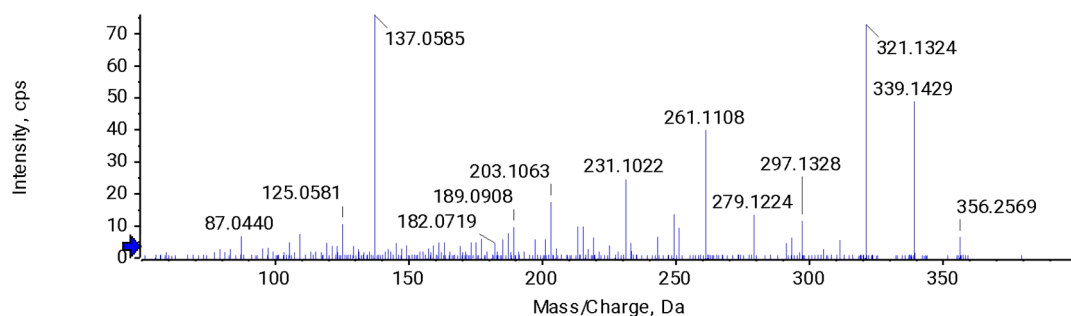
(Q) 3-ADON  $[M+H]^+$   $m/z$ : 339.144, RT: 3.10 min

Spectrum from 231107\_KY\_TOFposi\_2\_DataSET13.wiff (samp...from 3.068 to 3.205 min Precursor: 339.1 Da, CE: 25.0)



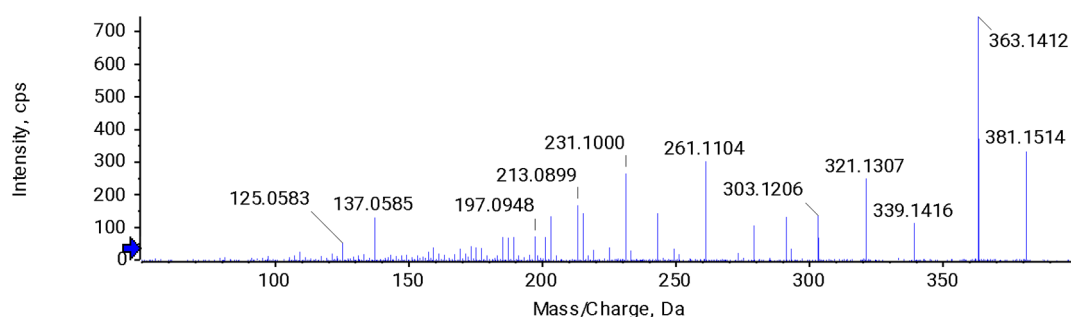
**(R) 15-ADON [M+NH<sub>4</sub>]<sup>+</sup> *m/z*: 356.170, RT: 3.05 min**

Spectrum from 231107\_KY\_TOFposi\_2\_DataSET14.wiff (samp...from 2.994 to 3.131 min Precursor: 356.2 Da, CE: 25.0)

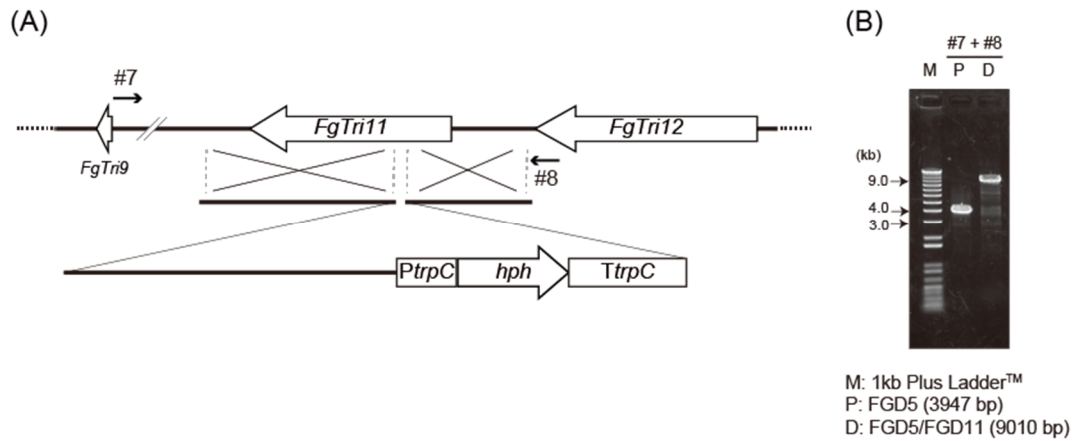


**(S) 3,15-diADON [M+NH<sub>4</sub>]<sup>+</sup> *m/z*: 398.181, RT: 3.84 min**

Spectrum from 231224\_KY\_TOFposi\_3\_DataSET15.wiff (samp...from 3.788 to 3.925 min Precursor: 398.2 Da, CE: 25.0)

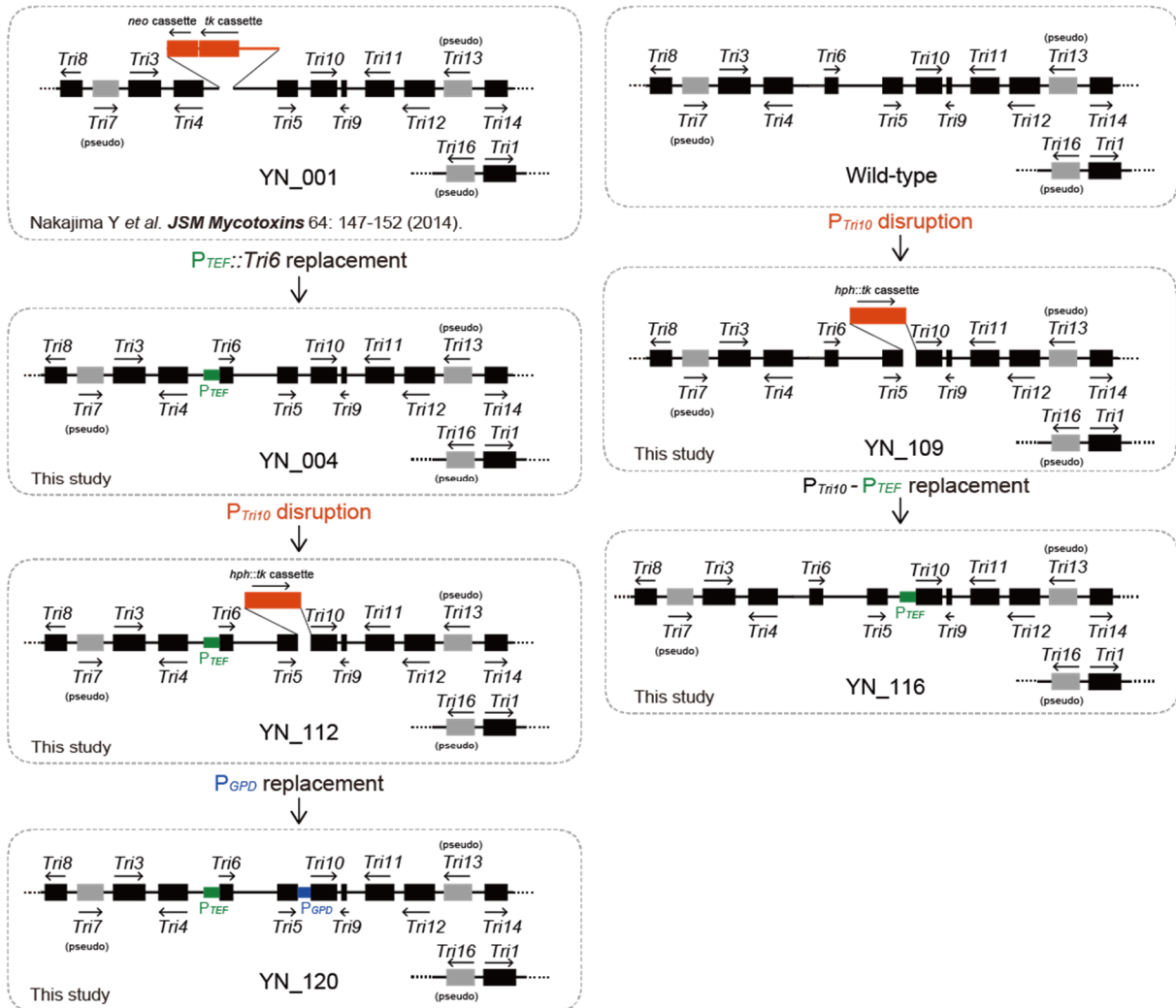


**Supplementary Figure S4.** MS/MS spectra of 16 tricyclic and 3 bicyclic trichothecenes and shunts. The purified samples were applied to LC-MS, and their MS/MS spectra were obtained in the positive ion mode, using the TOF-MS methods.

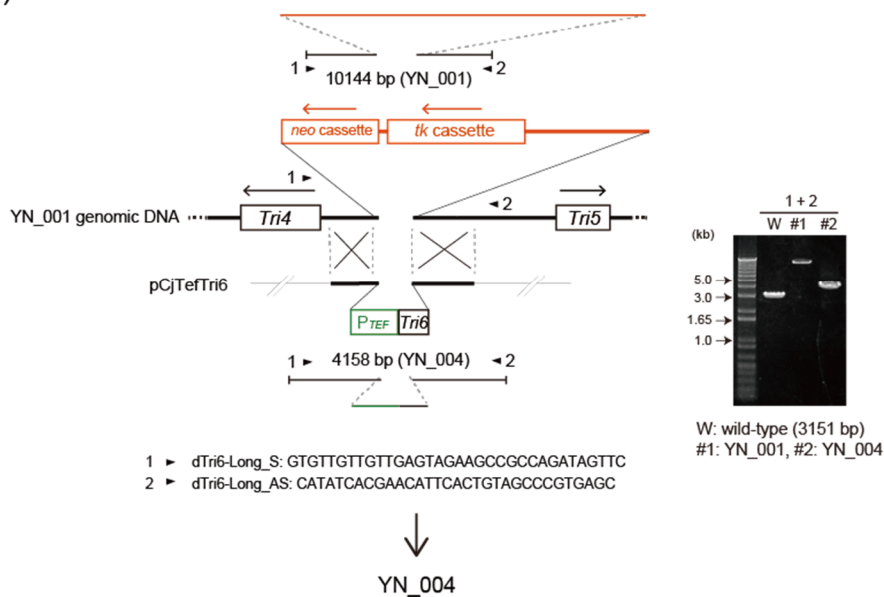


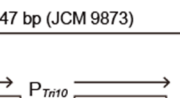
**Supplementary Figure S5.** Generation of a  $\Delta Fgtri5\Delta Fgtri11$  (FGD5/11) disruption mutant used in this study. (A) The *FgTri11* gene disruption vector [7] that contains the *hph* (hygromycin B phosphotransferase gene) cassette was transformed into FGD5 mutant strain [8]. After transformation, colonies were selected with 300  $\mu\text{g/ml}$  of hygromycin B. (B) PCR confirmation of genomic DNA isolated from the FGD5 and FGD5/FGD11 mutant strains. Primers #7 and #8 were used to amplify the region encompassing *FgTri11*. An amplicon from the double disruption mutant is 5.0 kb larger than that from its parent FGD5.

(A)

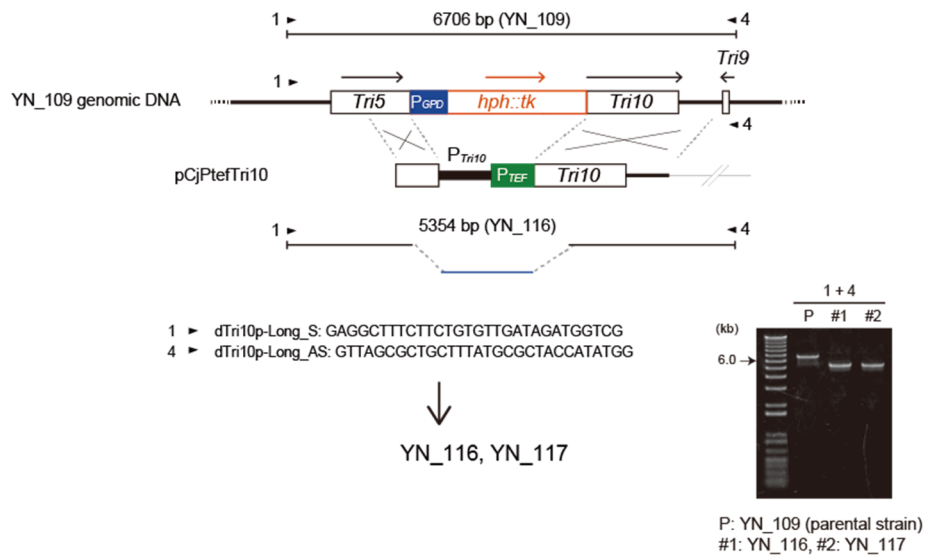


(B)

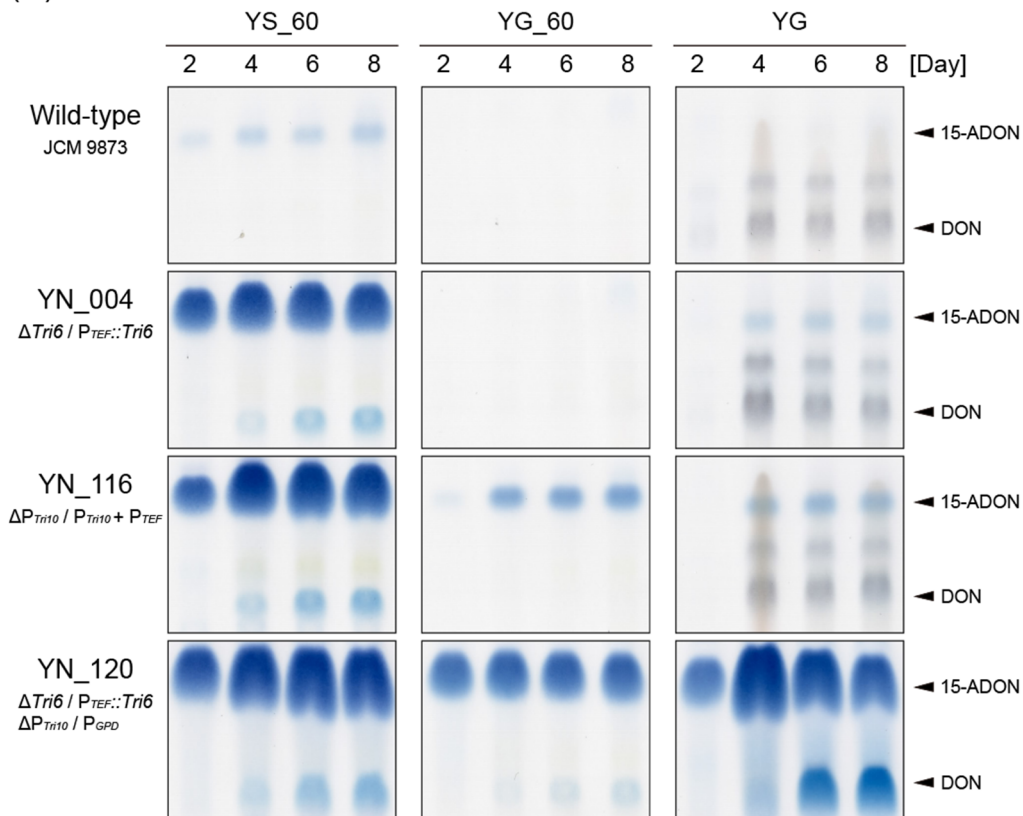








(C)



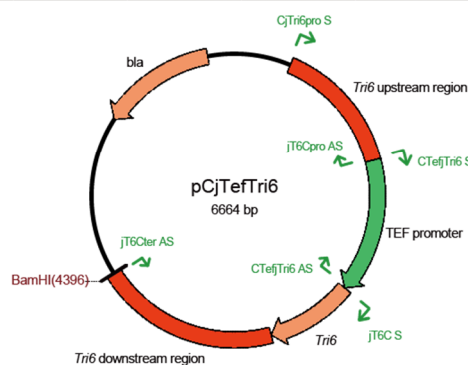
(D)

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201	CCGCACAGAT	GCGTAAAGAG	AAATACCGC	ATCAGGCGCC	ATTCCGCATT	CAGGCTGCGC	AACTGTTGGG	AAGGGCGATC	GGTGCGGGCC	TCTTCGCTAT
301	TACGCCACST	GGCGAAGGG	GGATGTGCTG	CAAGGCGATT	ANGTTGGTA	ACGCGAGGT	TTTCCGATC	ACGACGTTGT	AAACGACGG	CCAGTGAATT

CjTri6pro\_5

		CjTri6pro_S	
		Tri6 upstream region	
401		CGAGCTCGGT ACCCAGGTAC CTGTCTATC GCGTCTCTAG AGTGCCTTGC ATGCGTTGTG GCGGTAAAGC CTCACAACTC TGAAGTTGTC CTCAGTATCG	
		Tri6 upstream region	
501		CCGTGTGAGA TAGGCTTCGC AAGAGTGTCC GGTCCGAAAC TCACCAATCA ACTCAGCAGG ATGAACAAGG GGTCTGAAGG GCCTGSCAGG CCTGACAGGA	
		Tri6 upstream region	
601		GTGATAAAT GTGAGAGAG ATATGCCAT ACAACCTGT AMCTTGTGAA ACSSGSCATG GAATCCCATG GCAAGTTATG GSGTCAGCAG CAACTGAAAT	
		Tri6 upstream region	
701		GCTACAGGT CAGAAGTGC ATCTTTTAC CGGCGSCTTA TCCGAAGTTG CTGCCATCA GATACAGACA TGATGAGAGA GTGATACGAC TCGCGCGAAG	
		Tri6 upstream region	
801		AATAAGATC ATCAGTSCGC CGCAATGTTA AAACTGATG TCGCGAAGCA ACATTAGCT TTGAGGAGCAT GCCAGGGTCT TGCTCGAAA TATCTTTGTC	
		Tri6 upstream region	
901		TACCGAGCC CATGATGTT CAAAGTATG TACATGATG GTCTTGACA GAGACAGCC TCGAGTGTTA TCGAGACTGT CAGGCTGCA GTAAGTTGCC	
		Tri6 upstream region	
1001		ACAGACTGA ATCGATTATC ATTGACGTT CGGAAGCGCT CTGTTAGGAA TCTTTCTAGA CCAACTAC CACTTTGACA TCTGATACT AACACTAGTA	
		Tri6 upstream region	
1101		GCACATAGT AAACCTTCA CTGCGCGCG ATCAACTGT AAACAGGTAC CGGCGAGCG GTCTGGGATA AGATACCTT TTAACTGCC GTAGCAACT	
		Tri6 upstream region	
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		TEF promoter	
		CjTefjTri6_S	
		Tri6 upstream region	
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		jT6Cpro_AS	
		CjTefjTri6_S	
		TEF promoter	
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		TEF promoter	
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		TEF promoter	
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1901		ACCGTAGAG CGATAGCAG TAAAGAAAG CTAACAATA AAAATTCTT GCCCTAAGC CATGAAGAG AGATGGGTG GAGCAGACC AAGGAAGAG	
		TEF promoter	
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		TEF promoter	
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		TEF promoter	
2201		CCACAAAT TTTCTTCCCT TCCTCTCCT GTCCGCTCA GTACGTATAT CTCCCTTCC CTGCTTCTC TCCTCATCC TTTTTCATC CATCTCTGC	
		TEF promoter	
		Tri6	
		jT6C_S	
		M I	
2301		TAACTTCT GCTCAGACC TCTACGATT ACTAGCGTA GTATCTGAG ACTTCTCCCT TTTATATCC ACAAACATA ACACACCTT CACCATATT	
		CjTefjTri6_AS	
		jT6C_S	
		Tri6	
2401		Y H E A E S H Y E S W S A L P L F D R V A S P D P A K D F V P D L N TACATGAGG CGGAATCTCA CTACGAATCT TGAAGCGCT TCGCCCTCT TGAATGAGT GCGTCTCCG ATCTCGCAA GGAATTTGT CCAATCTAA	
		CjTefjTri6_AS	
		Tri6	
2501		N D Y E S P T F E I D L L S E T Y D F D N F P T Y S L P T V D S T K ACGACTTGA ATACAGACA TTGAAATAG ATCTTCTCT AGAATCTTAT GACTTTGACA ACTTCCACC ATACTCTCTA CCAAGGTGG ATTCAGCAA	
		Tri6	
2601		K T L Y S E E P L V C F D F D F A N P A I E N Y I T T S S G L L D A GACTTTGAT TCGAAGAAC CACTTGTGT CTTGACTTT GACTTCGCA ACCCGCTAT CGAAATATAT ATACACAT CTTGAGACT GTTGAGCGA	
		Tri6	
2701		V P S Q L I A L P T F T R P S K C P F P S C K S A T V F E S G R D F GTCCAGGCC AGCTTATGC CTTTCCACC TTACAGAGC CAAGCAATG CCAATCCCT AGTTGCAAGT CCGCCACAGT CTTTGAAGC GAGCGGACT	
		Tri6	
2801		F R R H Y R Q H F K R F F C R Y S E C P Q S A Q D L Q E V G T K G F TTAGCGGCA TTACGECOA CACTTCAAG GCTTTTCTG TCGCTACTCA GAATGCCCTC AGTCAGTCA AGACTGCAA GAGTCCGCA CCAAGGCTT	
		Tri6	
2901		F A T R K D R A R H E S K H K P T V R C P W Q D K E G Q Q C L R V F TGCGATGCG AAGGAGCTG CTGCGATGA GTCTAAGCAC AAACCAAGC TGCGGTGCC TTGCGAGAC AAGGAGGAC AACATGTCT GAGGCTTTT	
		Tri6	
3001		S R V D N M R D H Y R R I H K C * AGCAGGTGG ATAACTGCG AGATCACTAT AGGCGGATC ATAGTGTG ACAGGAGATC GGTGTGAAA ACTGAGATA GTTACTATA AAAGCAACA	
		Tri6 downstream region	
3101		TTTGGAAAA TCAATAACA TATCCGATA AATATCCAT TAGACTTTT GAGTTTATA AATGATGAG AGTTTGGCG CGATCGATG TGTTCAGC	
		Tri6 downstream region	
3201		ACCAATCAT AATATACTC ATAGTAGCG AGACCTGCA GCTCATGTT ACCAGCCCTG TTTTATTGC ACAATTAGT ACATCTATT CTGACAGCA	
		Tri6 downstream region	
3301		CTACAGGCT TTCAACTT ACATATTGC TCTAGCGTG GCTGTGCTA AGCAGATCA GCGCGTAT TGTATTGCT GCGTGTCTT GCGCCGATC	
		Tri6 downstream region	
3401		GTCCAGAAC AATCTGGCA CTGCAAGTA TTTGCGGCG CTGGTAAAT TCTTTAGTA GTCTGCTCA TTTGGCAGT GCAATAATA TATTTACC	
		Tri6 downstream region	
3501		GCGTTGCT GCGCATGTA CCAAAAGC TTCAATATA TCTGCGCAT ATCGTTACT CGGTTCCAG TTTACAGCA TACGCTCTC GTATCAATG	

3601	CCCAACAAA	CATACGTAT	TGGTACTAT	GCCGGTTTC	CCCGAACAT	TGCTGTCCG	CAAAAGGATG	CATCGGACT	CTAAAGATTG	CGATCTGTG
Tri6 downstream region										
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Tri6 downstream region										
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Tri6 downstream region										
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Tri6 downstream region										
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Tri6 downstream region										
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Tri6 downstream region										
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Tri6 downstream region										
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jT6Cter_AS										
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jT6Cter_AS										
4501	GCCGGAAGCA	TAAAGTGTAA	AGCTTGGGT	GCCTAATGAG	TGAGCTAATC	CACATTAATT	CGTTTGGCT	CAGTGCCTGC	TTTCCAGTGC	GGAAACCTGT
4601	CGTGCCAGCT	GCATTAATGA	ATCGGCCAAC	GCGCGGGGAG	AGCGGTTTGG	CGATTGGGG	GCCTTCCGC	TTCTCTGCTC	ACTGACTCGC	TGCGCTCGGT
4701	CGTTGGGCTG	CGCGAGCGG	TATCAGCTCA	CTCAAGGCG	GTAAACGGT	TATCCACAGA	ATCAGGGGAT	AACGACAGAA	AGACATGTTG	AGCAAAAGGC
4801	CAGCAAAAG	CCAGAACCG	TAAAAGGCC	CGCTTGGTG	CGTTTTTCCA	TAGGCTCCGC	CCCCCTGAGC	AGCATCACAA	AAATCGAGCG	TCAAGTCAGA
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5001	GTCCGCTTT	CTCCCTTGGG	GAGCGTGGC	GCTTTCTCAA	TGCTCAGCT	GTAGGTATCT	CAGTTGGGTG	TAGGTGTTTC	GCTCCAGCT	GGGCTGTGTG
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bla										
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bla										
5801	CCGAGCGCAG	AAGTGGTCT	GCAACTTAT	CCGCTCCAT	CCAGTCTATT	AATTGTGCTC	GGGAAGCTAG	AGTAAGTAGT	TGCGCAGTTA	ATAGTTTGGG
bla										
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bla										
6001	TCCCCATGT	TGTGCAAAA	AGCGGTTAGC	TCCTTGGCTC	CTCCGATCGT	TGTGAGAGT	AGTTGGGCGG	CAGTGTATC	ACTCATGTT	ATGGCAGCAC
bla										
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bla										
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bla										
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bla										
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bla										
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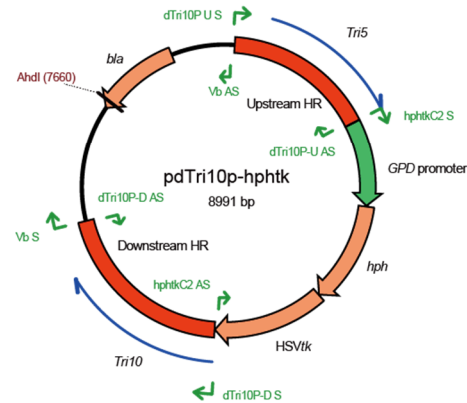
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Upstream HR										
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Tri5										
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Tri5										
401	ATTTTCTCAA	CAGTAGCGTG	CGCTTCTCTG	AGTATATTGG	ATACCGAGAC	AGCAATTACA	CCCGAGAGGA	GCGCATCGAG	AATTTCAGCT	ATGCTTACAA
Upstream HR										

		Tri5	
		Upstream HR	
501	CAGGCTGCG	CACCACTTGG	CTCAGCCTCG
		Tri5	
		Upstream HR	
601	GTATACAGCT	GGGCAAGGT	GTCCAAGAG
		Tri5	
		Upstream HR	
701	CTGCCATGT	GACTATTTT	GAAGCCTTC
		Tri5	
		Upstream HR	
801	TTTTGGACT	TTCTGCTCAT	TGAACCTTAT
		Tri5	
		Upstream HR	
901	CACACAGTTT	TGAGGATGT	TGATTGAGC
		Tri5	
		Upstream HR	
1001	TCATTGTGT	GGGCTCTCT	TATGGCCAA
		Tri5	
		Upstream HR	
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		Tri5	
		Upstream HR	
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		Tri5	
		Upstream HR	
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		Tri5	
		Upstream HR	
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		Tri5	
		Upstream HR	
1501	GGGCCAGGA	CGATGTGAAG	GAGGCTCAGA
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		Upstream HR	
1601	ACGACAGCAG	AGAGAGGGG	TGAGTAATAA
		Tri5	
		Upstream HR	
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		Tri5	
		Upstream HR	
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		Tri5	
		Upstream HR	
1901	CACGAGATAG	TACCTTCTCC	GAAGTAGSTA
		Tri5	
		Upstream HR	
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		Tri5	
		Upstream HR	
2101	TCGTCACTG	ACCTGCTGAG	GTCCCTCAGT
		Tri5	
		Upstream HR	
2201	TCACACATT	GTTCATAT	TTTCTGCTC
		Tri5	
		Upstream HR	
2301	ACCTTTATTT	CCCTTAAGTA	AGTACTTTGC
		Tri5	
		Upstream HR	
2401	GCAGCTGAC	TAAACCTAC	CCCGCTTGA
		Tri5	
		Upstream HR	
2501	TCCTGAGCT	GATGAGCTC	TGCGAGGCG
		Tri5	
		Upstream HR	
2601	TGGTTCTAC	AAAGATGTT	ATGTTATCG
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		Upstream HR	
2701	TATTCATCT	CCCGCGTGC	ACAGGTGTC
		Tri5	
		Upstream HR	
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		Tri5	
		Upstream HR	
3801	CGGGGACGCG	CGGTGTGTAA	TGACAGAGCG
		Tri5	
		Upstream HR	



3901	GGAGCTCAGC	ATGCCCCGCG	CCCGGCCCTC	ACCTCTGATC	TGGACCGCCA	TCCGATCGCC	GCCTCTCTGT	GCTACCCGGC	CGCGCGATAC	CTTATGGSCA
HSVtk										
4001	GCATGACCCC	CGAGGCCGTG	CTGGCGTTCG	TGGCCCTCAT	CCCCGCCGAC	TTGCCCGSCA	CCAAATCGTG	GCTTGGGGCC	CTTCCGGAGG	ACAGACACAT
HSVtk										
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HSVtk										
4201	TATCTGACGT	CGCGCGGGTC	GTGGCGGGAG	GACTGGGGAC	AGCTTTCCGG	GACGGCCGTG	CCGCCCCAGG	GTGCCGAGCC	CCAGAGCAAC	GCGGGCCAC
HSVtk										
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HSVtk										
4401	CTTGGCCAAA	CGCTCCGTTT	CCATGACAGT	CTTTATCTCG	GATTACGACC	AATCGCCCGC	CGGCTGCCGG	GACGCCCTGC	TGCAACTTAC	CTCCGGGATG
dTri10P-D_5										
Downstream HR										
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hphtkC2_AS										
Downstream HR										
Tri10										
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Tri10										
Downstream HR										
4701	AGAGGAGATG	CGTGTGACT	ATACTGACCT	CTGGCGGCC	TACGTACTAT	GCACATTGT	GCATGTGCT	CCTCTATAAA	GAATCGCTTT	CAAGCCCTTG
Tri10										
Downstream HR										
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Tri10										
Downstream HR										
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Tri10										
Downstream HR										
5001	GATGTGATG	CTCGATATCC	GATCTACGAT	TATCGTTGGT	CACTAACAAA	TTAAATAGT	CTTGGACCTT	AAGCAGGGGA	GATTGGCGGG	TTCACTTCCA
Tri10										
Downstream HR										
5101	TGCGGCCAAC	ATACTCATTC	CTGTCTTGST	TGAGGGATGG	TCCACAGCTT	TGCAATCAGG	CCCCCGAGCC	ACCTCATAT	GGTGGAGACT	GGATGATCA
Tri10										
Downstream HR										
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Tri10										
Downstream HR										
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Tri10										
Downstream HR										
5401	GACCATTTTG	ACTATTCTCG	AGTGGGGTAA	GCTGGATCGT	TGGAGGGGAC	AGGAGCAAGA	ACATAATCGC	TTGAGCCTAA	AGAGCGCTCG	TAGGGCGGCG
Tri10										
Downstream HR										
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Tri10										
Downstream HR										
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Tri10										
Downstream HR										
5701	TGAAGCTGTC	ACGAGCGTTA	CTTGGCTCT	AGCTGTCACA	GGATGATGGG	CCTCAGAAG	TCATAAGGAC	TTTTTCAGAA	ATACTCTGAG	GTGCTATGAG
Tri10										
Downstream HR										
5801	GGCAGCTTCA	GCCTCTTAAA	AAAGATGAGC	GGACCTCTTC	AGGCTCTGGA	AGAGCGTTGG	AAGAGAGAGG	AGATAGATAC	AGAGTCTCCA	ATGAGATGGG
Tri10										
Downstream HR										
5901	AGAGCTTGAC	GGATCACCAT	GGGCTTCCAG	TGCTACTTTG	GTAGGGATGG	TATCGGCGCA	GGAGCTACGG	ATACCGATAC	CCGAAGTCTT	TCTTCTGCA
Tri10										
Downstream HR										
6001	CAGCTACCGG	ATACATGTAA	GTTGATGCT	CACCTACAGA	TGATGACTGC	AGGAGGTTT	TATTAGATGT	TATGGSTTCG	ACTCATTTTG	GCSTTTAAAC
Tri10										
Downstream HR										
6101	AGGAGTTGGC	ATGATGCATG	ATTTAATATG	AATGAGAGGG	CGGAAACGTT	TTCCCATGTA	ACTTGATGTA	TGGACCTTCA	ATATATATTT	CTAGTATACT
Tri10										
Downstream HR										
6201	TTTTTAACGG	TGAGCCGAT	CTTCTCCATC	TATCCAAACA	ATCGTGTGTA	AAGCAGTCTG	GAACCATGGT	GGGCGAATCA	GTGCTGTGTC	CCTTGATTGT
Vb_5										
Downstream HR										
6301	CTCGACATGG	CTTGACATTG	TTTAGCCCGT	ACGCAAAAGC	ATATTGATCA	TAGTGAAAGG	GTGTTGATTT	CAAGCGAGGG	GATTTGTGGA	AGTTCCGGGGT
dTri10P-D_AS										
Vb_5										
6401	ACCGAGCTCG	AATTGCTAAT	CATGGTCATA	CGTGTTCCT	GTGTGAAATT	GTATCCGCT	CACAATTCCA	CACAACATAC	GAGCCGGAGG	CATAAAGTGT
dTri10P-D_AS										
6501	AAAGCTCGGG	GTGCTTAATG	AGTGAGCTAA	CTCAATAA	TTGGGTTGGG	CTCACTGCCC	GCCTTCCAGT	CGGGAAACCT	GTGCTGCCAG	CTGCATTAA
6601	GAATCGGCCA	ACGGCGGGGG	AGAGGCGGTT	TGGCTATTGG	CGCGCTCTCC	GCCTCTCGCG	TCACTGACTC	GCCTGCCGTC	GTGCTTCGGG	TGCGGCGAGC
6701	GGTATCAGCT	CACCTAAAGG	CGGTAATACG	GTATCCACAA	GAATCAGGGG	ATAACGACAG	AAAGAACATG	TGAGCAAAAG	GCCAGCAAAA	GGCCAGGAAC
6801	CGTAAAGAGG	CGCGGTTGCT	GGCGTTTTTC	CATAGGCTCC	GGCCCCCTGA	CGAGCATCAC	AAAAATCGAC	GCCTAAGTCA	GAGGTGGGGA	AAACCGACAG
6901	GACTATAAG	ATACAGGGCG	TTTCCCGCTG	GAGGCTCCCT	CGTGGGCTCT	CTGTTCGCGA	CCCTGGCGCT	TACCGGATAC	CTGCTCGGCT	TTCTCCCTTC
7001	GGGAGGCGTG	GGCGTTCTCT	ATAGCTCAGG	CTGTAGGAT	CTCAGTTGCG	TGTAGGCTGT	TGCGTCCAGG	CTGGGCTGTG	TGCACGAACC	CCCCGTTGAG
7101	CCGACCGCGT	GGCGCTTTAT	CGGTAACTAT	CGTCTTGAGT	CCAAACCCGG	AAGACAGGAC	TTATCGCCAC	TGGCAGCAGC	CACCTGGTAA	AGGATTAAGA
7201	GAGCGAGGTA	TGTAGGCGGT	GCTACAGAGT	TCTTGAAGTG	GTGGGCTAAC	TACGGCTACA	CTAGAGGAGC	AGTATTGGT	ATCTGCGCTC	TGCTGAAGCC
7301	AGTTACTCTC	GGAAAAAGAG	TTGGTAGCTC	TTGATCCGGC	AAACAACCA	CCGCTGGTAG	CGGTGGTTTT	TTTGTTTGCA	AGCAGAGAT	TACGCCGAGA

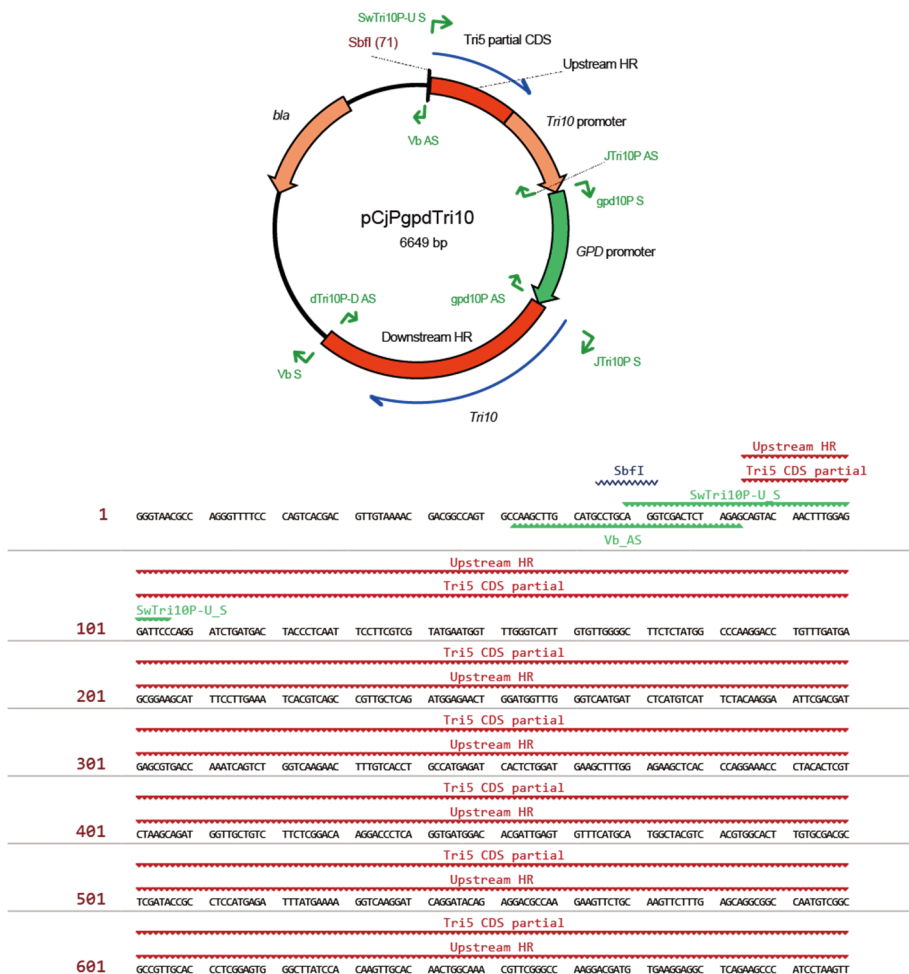
7401	AAAAAGAT	CTCAAGAAGA	TCCTTTGATC	TTTCTACGG	GGTCTGACGC	TCAGTGGAAC	GAAACTCAC	GTAAAGGAT	TTTGTCTATG	AGATTATCAA
7501	AAAGGATCTT	CACCTAGATC	CTTTTAATT	AAAAATGAG	TTTTAATCA	ATCTAAGATA	TATATGAGTA	AACTTGGTCT	GACAGTTACC	AATGCTTAAT
						AhdI				bla
7601	CAGTGAAGCA	CCTATCTCAG	CGATCTGTCT	ATTTCGTTC	TCCATAGTTG	CCTGACTCCC	CGTGTGTAG	ATAACTACGA	TACGGGAGGG	CTTACCATCT
						bla				
7701	GGCCCCAGTG	CTGCAATGAT	ACCGCGAGAC	CCACGCTCAC	CGGCTCCAGA	TTTATCAGCA	ATAAACACGC	CAGCCGGAAG	GGCCGAGCGC	AGAAGTGTCT
						bla				
7801	CTGCACTTTT	ATCCGCTCTC	ATCCAGTCTA	TTAATTGTTG	CCGGGAAGCT	AGAGTAAGTA	GTTCGCGAGT	TAATAGTTTG	CGCAACGTTG	TTCGCAATTG
						bla				
7901	TACAGGCATC	GTGGTGTAC	GCTCGTCTT	TGGTATGGCT	TCATTCACT	CCGGTTCCCA	ACGATCAAGG	CGAGTTACAT	GATCCCCCAT	GTGTGTCAAA
						bla				
8001	AAAGCGTTA	GCTCCTTCGG	TCCTCCGATC	GTGTGCAGAA	GTAAATTGGC	CGCAGTGTTA	TCATCATGG	TTATGGCAGC	ACTGCATAAT	TCTCTTACTG
						bla				
8101	TCATGCCATC	CGTAAGATGC	TTTTCTGTGA	CTGGTGAAGTA	CTCAACCAAG	TCATTCTGAG	AATAGTGTAT	GGGGGACCG	AGTTGCTCTT	GGCCGGGCTC
						bla				
8201	AATACGGGAT	AATACCGCG	CACATAGCAG	AACTTTAAAA	GTCTCATCA	TTGGAAGACG	TTCTTGGGG	CGAAACTCT	CAAGGATCTT	ACCGCTGTTG
						bla				
8301	AGATCCAGTT	CGATGTAACC	CACCTGTGCA	CCCACTGAT	CTTCAGCATC	TTTACTTTT	ACCAGGTTT	CTGGGTGAGC	AAAAACAGGA	AGGCAAAATG
						bla				
8401	CCGCAAAAA	GGGAATAAGG	CGGACACGGA	AATGTTGAAT	ACTCATACTC	TTCTTTTTC	AATATTATTG	AAGCATTAT	CAGGGTTATT	GTCTCATGAG
						bla				
8501	CGGATACATA	TTTGAATGTA	TTTAGAAAA	TAAACMAATA	GGGGTCCGC	GCACATTCC	CCGAAAGTG	CCACTGACG	TCTAAGAAAC	CATTATTATC
8601	ATGACATTAA	CTATAAAAA	TAGGCTGATC	ACGAGGCCCT	TTGCTCTCG	GGTTTCGGT	GATGACGGTG	AAACCTCTG	ACACATGACG	CTCCCGGAGA
8701	CGGTCACAGC	TTGTCTGTAA	CGCGATGCCG	GGAGCAGACA	AGCCCGTCAG	GGCGGCTCAG	CGGGTGTTCG	CGGGTGTTCG	GGCTGGCTTA	ACTATGCGGC
8801	ATCAGACGAG	ATTGTAAGTA	GAGTGCACCA	TATGCGGTGT	GAAATACCGC	ACAGATGCGT	AAGGAGAAAA	TACCGCATCA	GGCGCCATTC	GCCATTCAGG
8901	CTCGCAACT	GTGGGAGG	GCATCGGTG	CGGCTCTT	CGCTATTACG	CCAGCTGGCG	AAAGGGGAT	GTGCTGCAAG	GCATTAAAT	T



1	GGGTAGGCC	AGGGTTTTCC	CAGTACGAC	GTGTAAAC	GACGGCCAGT	GCACAGCTTG	CATGCTGCA	GGTCACTCT	AGAGCAGTAC	AACCTTGGAG
						SwTri10P-U.S				
						Vb_AS				
						Tri5 partial CDS				
						Upstream HR				
101	GATTCACAG	ATCTGATGAC	TACCTCAAT	TCCTTCGTG	TATGATGAT	TTGGGTCTT	GTGTTGGGG	TTCTCTATG	CCCAAGGACC	TGTTTGATGA
						Upstream HR				
201	GGCGAGCAT	TTCTTGAAA	TCACGTGAC	CGTTGCTCAG	ATGAGAACT	GGATGTTTG	GGTCAATGAT	CTCATGTCAT	TCTACAGGA	ATTGAGCAT
						Upstream HR				
301	GAGCGTACC	AATCAGTCT	GGTCAAGAC	TTTGTCACT	GCCATGAGAT	CACCTGGAT	GAGCTTTGG	AGAGGCTGAC	CCAGGAAACC	CTACACTGCT
						Upstream HR				
401	CTAAGCAGAT	GGTTGCTGTC	TTCTCGACA	AGGACCTCTA	GGTGTGAGC	ACGATTGAT	GTTCATGCA	TGGCTAGCTC	ACGTGGCACT	TGTGGGAGCG
						Upstream HR				
501	TCGATACCG	CTCCATGAGA	TTTATGAAA	GGTCAAGGAT	CAGGATACAG	AGGACGCCAA	GAAGTTCTGC	AACTTCTTTG	AGCAGCGCGC	CAATGTCGGC
						Upstream HR				
601	GCCCTTGAC	CCTCGGAGTG	GGCTTATCCA	CAAGTTGAC	AACTGGCAAA	CGTTCCGGCC	AAGGAGCATG	TGAAGGAGCG	TCAGAGGCC	ATCTAAAGT
						Tri10 promoter				
						Upstream HR				
						Tri5 partial CDS				
701	CAATTGACCT	AGTGGATAA	CCGAGGCGA	GTTTGGAGT	ATGTTTGGG	GGTACGGATA	CTGCTTTGGA	GAATGTTGCT	CTGTTAAT	GATTACAAAT
						Tri10 promoter				
801	AGTTCGGTG	TGTTTGTGA	GAATGACAG	TTGAGCAAG	ATAATTACTT	CGGATAGGC	AGTTGAAGCT	GAATGCTCTG	ACGTAACTGT	AGCCTGTAA
						Tri10 promoter				
901	CATTTCACG	TTGAGTGCAG	GCCTTTGGCT	AACCAAGTCT	GTACACCGCT	GGTGGGACCA	GGGCTACCCC	CAACCTGCA	ACTGATCTG	CAGCTGACG
						Tri10 promoter				
1001	TGGCAGAGTG	GTAGACTGGC	GCTACGAAT	GTAGTGGAT	GGCGGAATCT	TGTATCCGCT	CGGAGTTTGG	AGGGAGCTGG	CACATTCTCT	AGACCCGCGA
						Tri10 promoter				
1101	ATTGATCTTC	AAAGCGCTTG	CGTTTGTGTC	AGATCAGTGA	CCATACCTT	GCTTTTGGC	ACCACCCAAA	CGTCCACTGA	ACGAGGCGTA	CAGAAACAC

1201	ACAGATAGG GTTTGATGCC TGCTTGAGCA CTATGAGGGA CACGACACT CTGTAAACT CTATCCTTGC ATTATATTGT AACATCGTTT AACTTCTCCA
	Tri10 promoter gpd10P_S GPD promoter
1301	CGCCCATCC TTCCCGTCTT TCCCCCACT TCAATTGTAT GCCAACCAAC AATCATCAA TTATCATTAT TATTGTGTT AGTCATCCCG GTGACTCTTT
	Tri10 promoter JTri10P_AS GPD promoter
1401	CTGGCATGCG GAGAGACGGA CGAGACGAGA GAGAAGGGCT GAGTAATAAG CGCCACTGCG CCAGACAGCT CTGGCGGCTC TGAGGTGAG TGAGTGATTA
1501	TTAATCCGGG ACCCGCCGCC CCTCCGCCCC GAAGTGGAAA GGTGTGTGTG CCCTCGTTTG ACCAAGATC TATTGCATCA TCGGAGATA TGGAGCTTCA
1601	TGGAATCACC GGCATTAAGC GAGGAGAAAT GTGAAGCGAG GGTGTATAG CCCTCGGCGA AATAGCATGC CATTACCTA GGTACAGAG TCCAATTGCT
1701	TCGATCTGG TAAAGATTCC ACAGATAGT ACCTTCTCCG AAGTAGTAG AGCAGATACC CGGCGGTAA GCTCCTAAT TGGCCATCC GGCATCTGTA
1801	GGCGSTCAA ATATCGTGCC TCTCTGCTT TGCCCGGTGT ATGAACCGG AAGGCCGCT CAGGAGCTGG CCAGCGGCG AGACCGGAA CACAAGCTGG
1901	CAGTCAGCCC ATCCGGTGCT CTGCATCGA CCTCTGAGC TCCCTGAGTC CCGTAGGAG AGCTTTGCCC GGTCTGCGC CCCGTTGTGT CGGCGGGTT
2001	GACAAGTGG TTGGCTAGT CCAACATTG TTGCATATT TTCTGCTCT CCCACAGC TGCTCTTTTC TTTTCTCTT CTTTCCCAT CTTAGATATA
2101	TTATCTTCC CATCCAGAA CTTTATTTC CCCTAAGTAA GTACTTGTCT ACATCCATAC TGCATCCTTC CCATCCCTTA TTCTTTGAA CTTTCAGT
	GPD promoter Downstream HR Tri10 JTri10P_S gpd10P_AS
2201	CGAGCTTCC CACTTCATG CAGCTTGACT AACAGTACC CCGCTTGAGA TGATTTCCT AAGCGTAGA CAGTCCGAG AGACGAGCCT GTTGATGTAC
2301	TACTTAGAG TCGTGTTCCT TCTGATGTC ATCAACCAA ACAACAATTG TCTGGGAAG AGAGAGTGG TGTGTACTAT ACTGACTCT GCGCGGCTTA
2401	CGTACTATGC CACATTGTGC ATGTGCGTCC TCTATAAGA ATGCTTTTCA AGCCCTTGCA GATCTGAGA GCGATGGTA TGGAGAGAG AGAAGACATA
2501	CTACTACAT CTTGCACTCC AGAGTCTCA GAAGCTGCTG GGTGGGCTG ACAAGACATT TGGATCACA AGGCTGAAG GTACCGTGT TGGCTTGTCT
2601	TGACTGTAC AGCTTATCAG TTTTGAGTA AGACGAATCC ACCATTGTT CGATGCTGGA TGTGATGCT CGATATCCGA TCTAGATTA TCGTTGTCA
2701	CTAACAAAT AAAATAGTCT TCGACCTAA GCGAGGAGGA TTGGCGCGTT CACCTCCATG CGGCCAAT ACTCATTCT GTCTTGGTTG AGGATGGTC
2801	CGAGCTTTG GATCAGGCC CCCAGCCAC CTCATATGG TCGAGCTGG ATGAATACA CTTCGGCTG ACTGAATC AACTCTCTT GAGCTTCGA
2901	TAGCTCGAG CTTTGAGATT CCTGTCAAC TCACTCGCG CAGTCGGCAT CCGTCTTGC ATATCTATTG GCCCATCAG ACCATTGAA GATTACGGC
3001	ATCTCTCGA CCAGCCAGGC CTTATACAG TGGACGAGT GCTGGGGTG AGGAATTGA CCATGTTGAC TATTCTGAA GTGGTAAGC TGGATCGTTG
3101	GAGCGACAG GAGCAAGAAC ATATCGCTT GAGCTAAG ACCTCTGCTA GCGCGCCAT GATGATTGAG GATATGTTG CAGAGAGCT ACAAGGCTA
3201	CCGACAGAG AGAGCTTCC AGACTCATC ACTCAGATT ACGCCGCTC TATCATGAG TATCTGCATA CAGTAGTTC CGGACTCAI CCCAACCCTT
3301	CAGAGTTCA GATAGTGTG GCGGAGGCG TTCAATTGTT GAGAGGCTC CCAATCTTG AAGCTGTAC GAGCGTACT TGGCTCTAG CTGTACAGG
3401	ATGATGACC TCAGAAAGTC ATAAGGACTT TTTCAGAAAT ACTCTGAGT CGTATGAGGC GACATTGAG TCCTTAAAA AGTATGAGCG AACTCTTCA
3501	GTCTTGGAG ACCTCTGGAA GAGAGAGAG ATAGATACAG AGTCTCCAAT GAGATGGAA GACTTGAGCG ATACCATGG GCTTCCAGT CTACTTGTG
3601	AGGATGGTA TCGGCGCAGA GACTACGAT ACCGATACC GAAGTCTTC TTCTGCATCA GCTACCGAT ACATGTAAG TTGATGCTA CTTAGAGATG
3701	ATGATGAGC GGAAGTTTTA TTAGATGTA TGGGTTGAG TCATTTTGGC GTTTAAACAG GATTTGGCAT GATGATGAT TTAATGGAA TAGAAGAGCG
3801	GAAAAGTTT CCATGTATC TTGATGATG GACTTCAAT ATATATTCT AGTATACTTT TTTAAGCGTG AACCGATCT TCTCATCTA TCCAACCAAT
3901	CGTCTGAAA GCACTCTGA ACCATGTTGG GCGAATCAGT GCTGTGTCCC TTGATGTCT CGACATGGCT TGACATTGT TABCCGCTAC GACAACGAT
4001	ATTGATACA GTGAAGAGT GTTGATTCA AGCGAGGGA TTTGTGAG TTCCGGGTAC CGAGCTCGAA TTGTAATCA TGGTCATAG TGTTCCTGT
	Downstream HR Vb_S dTri10P-D_AS

4101	GTGAATTGT	TATCGCTCA	CAATCCACA	CAACATACGA	GCCGGANGCA	TAAAGTGTA	AGCTGGGGT	GCTTAATGAG	TGAGCTAACT	CACATTAACT
4201	GGTTTGGCT	CACCTGCCG	TTTCCAGTCG	GAAACCTGT	CTGTCAGCT	GCAATTAATG	ATCGGCCAAC	GCGCGGGGAG	AGCGGGTTTG	CGTATTGGCC
4301	GCCTTCCGC	TTCTCGCTC	ACTGACTCG	TGCGCTCGGT	CGTTCGCGTG	CGCGAGCGG	TATCAGCTCA	CTCAAGGCG	GTAATACGCT	TATCCACAGA
4401	ATCAGGGGAT	AACGACGAA	AGAATCTGTG	AGCAAAAGGC	CAGCAAAAGG	CCAGGAACCG	TAAAAAGGCC	GCCTTCTGCG	CGTTTTTCCA	TAGGCTCCGC
4501	CCCCCTGACG	AGCATCACAA	AAATCGACGC	TCAAGTCAGA	GGTGGCGAAA	CCGACAGAGA	CTATAAGAT	ACCAAGCGTT	TCCCCCTGGA	AGCTCCCTCG
4601	TGCGCTCTCC	TGTTCCGACC	CTGCGCTTAA	CCGATACCTT	GTCCGCTCTT	CTCCCTTCGG	GAAGCGTGGC	GCCTTCTCAT	AGCTACGCT	GTAAGTATCT
4701	CAGTTCGGTG	TAGGTCGTTT	GTCTCAAGCT	GCGCTGTGTG	CAGCAACCCC	CGTTCAGCC	CGACGCTGCG	GCCTTATCCG	GTAAGTATCG	TCTTGAGTCC
4801	AACCGGTAA	GACACGACTT	ATCGCACTG	GCAGCAGCCA	CTGGTAAACG	GATTACAGA	GCAGGATATG	TAGGCGGTGC	TACAGAGTTC	TTGAAGTGGT
4901	GGCTAACTA	CGGCTACACT	AGAAGACAG	TATTTGGTAT	CTGCGCTCTG	CTGAAGCAG	TTACCTTCGG	AAAAAGATT	GSTAGCTCTT	GATCCGCGAA
5001	ACAAACACC	GCTGTGAGCG	GTGTTTTTTT	TGTTTGCAG	CAGCAGATTA	CGCGCAGAAA	AAAGGATCT	CAGAAGATC	CTTTGATCTT	TTCTACGGGG
5101	TCTGACGCTC	AGTGGAAACG	AAACTCACGT	TAAAGGATT	TGGTCATGAG	ATTATCAAAA	AGGATCTTCA	CCTAGATCCT	TTTAAATTA	AAATGAAGTT
5201	TTAATCAAT	CTAAGTATA	TATGAGTAAA	CTTGGTCTGA	CAGTTACCAA	TGCTTAATCA	GTGAGGCACC	TATCTCAGCG	ATCTGTCTAT	TTGTTTATC
b1a										
5301	CATAGTTGCC	TGACTCCCCG	TCGTGTAGAT	AACTACGATA	CGGGAGGGCT	TACCACTGG	CCCCAGTCT	GCAATGATAC	CGCGAGACCC	ACGCTCACCG
b1a										
5401	GCCTCAGATT	TATCAGCAAT	AAACAGCCA	GCCGGAAGGG	CCGAGCGCAG	AAGTGGTCT	GCAACTTTAT	CCGCTCCAT	CCAGTCTATT	AATTGTTGCC
b1a										
5501	GGGAGCTAG	AGTAAGTAGT	TCGCCAGTTA	ATAGTTTGGC	CAACGTTGTT	GCCATTGCTA	CAGGATCGT	GGTGTACGC	TCGCTGTTTG	GTATGGCTTC
b1a										
5601	ATTACGCTCC	GGTTCCTAAC	GATCAAGGCG	AGTTACATGA	TCCCCCATGT	TGTGCAAAA	AGCGGTTAGC	TCCTCGGTC	CTCCGATCGT	TGTGAGAGT
b1a										
5701	AAGTTGGCCG	CAGTGTATC	ACTCATGTT	ATGGCAGCAC	TGCATAATTC	TCTTACTGTC	ATGCCATCCG	TAGATGCTT	TTCGTGACT	GGTGAAGTACT
b1a										
5801	CAACGAGTC	ATTCTGAGAA	TAGTGTATGC	GGGACCGAG	TTGCTCTTGC	CGGCGTCAA	TACGGGATA	TACCGGCCA	CATAGCAGAA	CTTTAAAGT
b1a										
5901	GCATCATTT	GGAAACGTT	CTTCGGGGCG	AAACTCTCA	AGGATCTTAC	CGCTGTTGAG	ATCCAGTTCG	ATGTAACCCA	CTCGTGCACC	CAACTGATCT
b1a										
6001	TCAGCATCTT	TTACTTTTAC	CAGCGTTTCT	GGGTGAGCAA	AAACAGGAAG	GCAAAATGCC	GCAAAAAGG	GAATAAGGC	GACACGAGAA	TGTTGAATAC
b1a										
6101	TCATCATCTT	CTTTTTTCAA	TATTATTGAA	GCATTATCA	GGGTATTGTT	TCATGAGCG	GATACATATT	TGAATGTATT	TAGAAAAATA	AACAATAGG
b1a										
6201	GGTTCGCGC	ACATTTCCCG	GAAAGTGCC	ACCTGAGCTC	TAAGAAACCA	TTATTATCAT	GACATTAAAC	TATAAAATA	GGCGTATCAC	GAGGCCCTTT
6301	CGTCTCGCG	GTTCGGTGA	TGACGCTGAA	AACCTCTGAC	ACATGACGCT	CCCCGAGACG	GTCACAGCTT	GTCTGTAGC	GGATGCCGGG	AGCAGACAG
6401	CCCGTCAGGG	CGGTCAGCG	GGTGTGGCG	GGTGTGGGG	CTGGCTTAAC	TATCGGCAT	CAGACAGAT	TGTACTGAGA	GTGACCATTA	TGCGGTGTGA
6501	AATACGCGAC	AGATCGGTAA	GGAGAAATA	CCGATCAGG	GCGCATTCGC	CATTACGGCT	GCACAACGT	TGGGAAGGCG	GATCGGTGCG	GGCCTCTCTG
6601	CTATTACGCC	AGCTGGCGAA	AGGGGAGTGT	GCTGCAAGGC	GATTAAAGT					



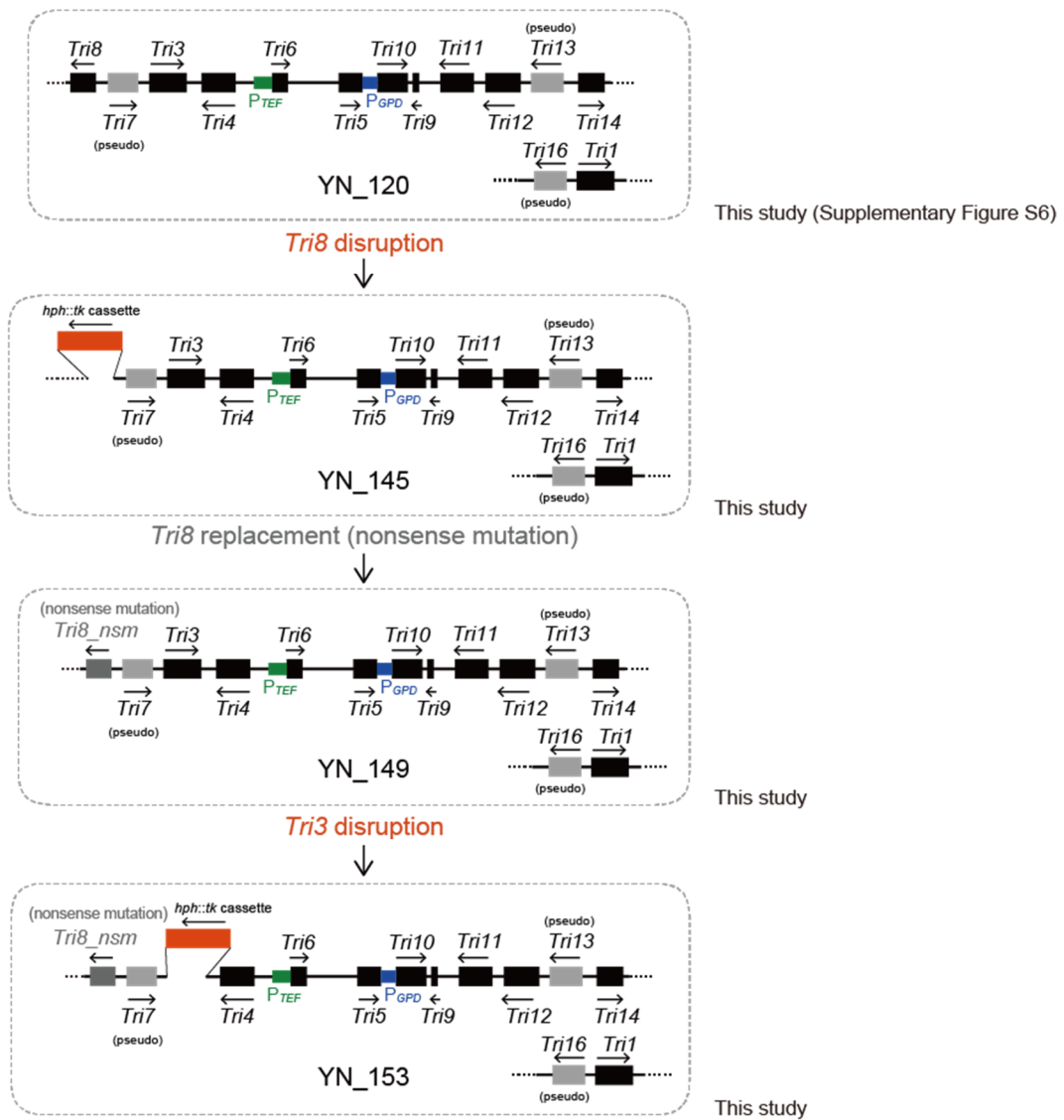


	Tri5 CDS partial									
	Upstream HR					Tri10 promoter				
701	CAATTGAGCT	AGTGGATGAA	CCGAGGCCA	GTTTGGAGT	ATGTTTTCG	GGTACGGATA	CTCGTTTGA	GAATGGTGGT	CTGTATAAT	GATTACAAAT
801	Tri10 promoter									
	AGTTCGGTCG	TGTTTTGTTA	GAATGAACAG	TTGAACAAGG	ATAATTACTT	CGGAATAGGC	AGTTGAACCT	GAATGCTGT	ACGTAACTCT	AGCCTGTAAAC
901	Tri10 promoter									
	CATTTCCCACT	TTGAGTGCAG	GCTTTTGGCT	AACCAAGTCT	GTACACCCGT	CGGTGCGACA	GGGCTACCCC	CAACCTCGCA	ACTGCATCTG	CAGCTGCAAC
1001	Tri10 promoter									
	TGGCAGACTG	GTAGACTGGC	GCTACGAATC	GTAGTGCAT	GCGGGAATCT	TGTACCCGCT	CGGAGGTTGG	AGGGAGCTGG	CACATTCTCT	AGACCCGCGA
1101	Tri10 promoter									
	ATTGATCTTC	AAAGCGCTTG	CGTTTTGTCC	AGATCAGTGA	CCATACCTT	GCTTTTTCGC	ACCACCCAAA	CGTCCACTGA	ACGAGGCGTA	CAGAAACAC
1201	Tri10 promoter									
	ACAAGATAG	GTTTGATGCC	TGCTTGAGCA	CTATGAGGGA	CACGACACTT	CTGTAAACT	CTATCCTTGC	ATTATATTGT	AACATCGTTT	AACTCTCTCA
	tef10P_S									
	TEF promoter									
1301	Tri10 promoter									
	CGCCCATCC	TTCCCGTCTT	TCCCCCAACT	TCAATTGTAT	GCCAAACAC	AATCATCAA	TTATCATTAT	TATTGTGCTT	AGTCATCTAG	CGACTTAGAG
	JTri10P_AS									
	TEF promoter									
1401	tef10P_S									
	TGCACGCTCA	AAAGATAACC	CTAACTCTG	ATTGTATAG	ATTGTATAT	GGCATATAGT	ACGGCTGAG	CATAAAAAA	AAACAGAT	TATTATAGTT
1501	TEF promoter									
	TATTATACG	AGACAGCAGA	ATACCCGCC	AAGTTAAGCC	TTTGTGCTGA	TCATGCTCTC	GAACGGGCCA	AGTTCCGGAA	AAGCAAAGGA	GCSTTTAGTG
1601	TEF promoter									
	AGGGCAATT	TGACTCACCT	CCAGGCAAC	AGATGAGGGG	GGCAAAAAGA	AAGAAATTTT	CGTGAATCAA	TATGGATTCC	GAGCATCAIT	TTCTTGCST
1701	TEF promoter									
	CTATCTGCT	ACGTATGTTG	ATCTTGACGC	TGTGATCAA	GCAACGCCAC	TCGCTCGCTC	CATCGCAGGC	TGGTCGCGA	CAAAITAAAA	GGCGGCAAC
1801	TEF promoter									
	TGCTACAGCC	CGGGGGTGT	CCGCTGCAA	GTACAGAGTG	ATAAAGCGG	CCATGCGACC	ATCAACGCGT	TGATGCCAG	CTTTTTCGAT	CCGAGATCC
1901	TEF promoter									
	ACGCTAGAGG	CGATAGCAAG	TAAAGAAAAG	CTAAACAAA	AAAAATTCT	GCCCCAAGC	CATGAAAAGG	AGATGGGGTG	GAGCAGAAC	AAGGAAAGAG
2001	TEF promoter									
	TCGCGCTGGG	CTGCCGTTC	GGAGGTGTT	GTAAAGGCTC	GACGCCCAAG	GTGGAGTCT	AGGAGAGAA	TTTGCAATCG	GAGTGGGGCG	GTTTACCCCT
2101	TEF promoter									
	CCATATCCA	TGACAGATAT	CTACAGCCA	AGGGTTGAG	CCCGCCGCT	TAGTGTGCTG	CTCGGCTTG	CCCTCCATA	AAGGATTTC	CCTCCCCCTC
2201	TEF promoter									
	CCACAAATT	TTCTTTCCCT	TCCTCTCCT	GTCCGCTTCA	GTACGATAT	CTCCCTTCC	CTCGCTCTC	TCCTCCATCC	TTCTTTCATC	CATCTCTGC
	Downstream HR									
	Tri10									
	JTri10_S									
2301	TEF promoter									
	TACTTCTCT	GTACAGACC	TCTACGATT	ACTAGCGTA	GTATCTGAGC	ACTTCTCCCT	TTTATATCC	ACAAACATA	ACACACCTT	CACCATGAT
	tef10P_AS									
2401	Tri10									
	JTri10_S									
	Downstream HR									
	TTCCCAAGC	CTAGACAAGT	CCGAGAGAGC	AGCTGTTGA	TGTACTACCT	AGAGCTGGTG	TTTCTCTGCG	AATGCTCAA	CCCAACAC	AATTGTCTGG
	tef10P_AS									
2501	Tri10									
	Downstream HR									
	GAAGAGAGA	GTGGCTGTTG	ACTATACTGA	CCTTGCAGG	GCCTACGTAC	TATGACCAT	TGTGCAATG	GCCTCTAT	AAAGATCGC	TTTCAAGCCC
2601	Tri10									
	Downstream HR									
	TTGCAGTCT	GAACAGCGA	TGGTATGAA	GAGAGAGAG	ACATACTACT	ACATTCTTGC	ACTCAGGAG	TCTCAGAGC	TGCTGGGTGG	GCTCGACAG
2701	Tri10									
	Downstream HR									
	ACATTTGCA	TCACAGGCT	GAAGGTACC	GTGCTGCCC	TTGCTTGAT	GCTACAGCTT	ATCAGTTTGG	AGGTAGAGC	AATCCACAT	TGTTGCGATG
2801	Tri10									
	Downstream HR									
	CTGAGTGG	ATGCTGATA	TCCGATCTAC	GATTATGTT	GGTCACTAGC	AAATTAAAT	AGTCTTGGCA	CCTAGCAGG	GGAGATTGGC	GGTTTCACT
2901	Tri10									
	Downstream HR									
	CCATGCGGC	AACTACTCA	TTCTGTCTT	GTTTGAGGA	TGGTCCACAG	CTTTGCAATC	AGGCCCCCA	GGCACTCCA	TATGTTGCGA	GCTGGATGA
3001	Tri10									
	Downstream HR									
	TCACACTCTG	GCTCGACTGA	AGATCAAAAC	TCTTTGAGCT	TCGAATAGCT	CGAGCTTTG	AGATTCCTGT	CAAACTCACT	CGCCGAGTC	GGCATCCTGT
3101	Tri10									
	Downstream HR									
	CTGCTATATC	TATTGGCCA	TCAGACCAT	TTGAAGATTA	CGGCACTCTC	CTGACCAAGC	CAGGCTTAT	ACAGCTGAGC	GAGTGTGG	GTTGCGAGAA
3201	Tri10									
	Downstream HR									
	TTGACCATG	TTGACTATC	TGAGTGGG	TAGCTGGAT	CGTTGGAGC	GACAGAGCA	AGAACATAAT	CGCTTGAGCC	TAAAGAGCT	CGTAGGCGC
3301	Tri10									
	Downstream HR									
	GCATATGTA	TGAGGATAT	GTGTGAGC	GAGCTACAAA	GCTACCGAC	AGACGAGCG	CTTCAGAGC	TCATCACTCA	GATTAGGCC	GCCTCTATCA
3401	Tri10									
	Downstream HR									
	TGACGTATCT	GCATACAGTA	GTTTCCGAGC	TCAATCCCAA	CCTTTCAGAG	GTTCAAGATA	GTGTGCGCG	GACGCTTCAA	TTGTGGAGA	GCTCCCAA
3501	Tri10									
	Downstream HR									
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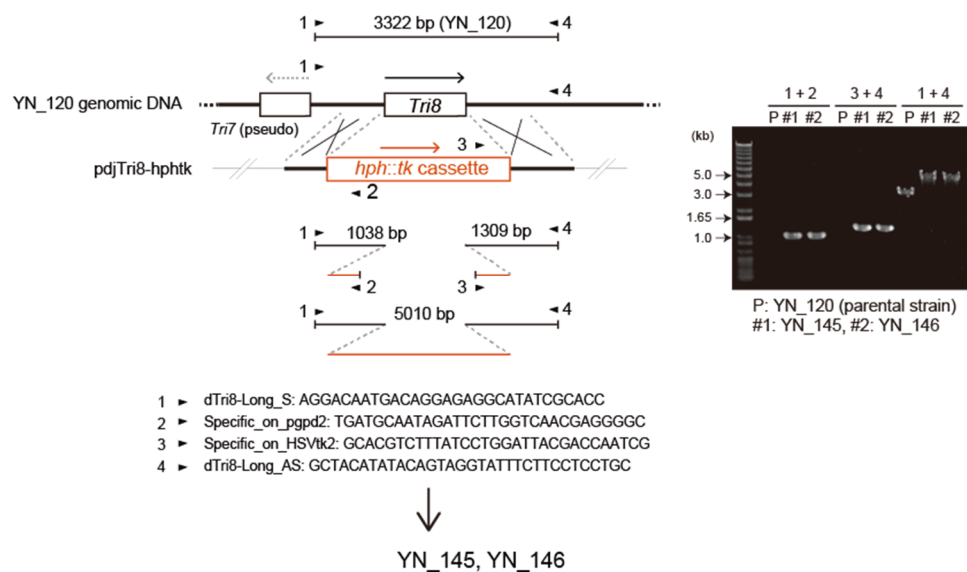


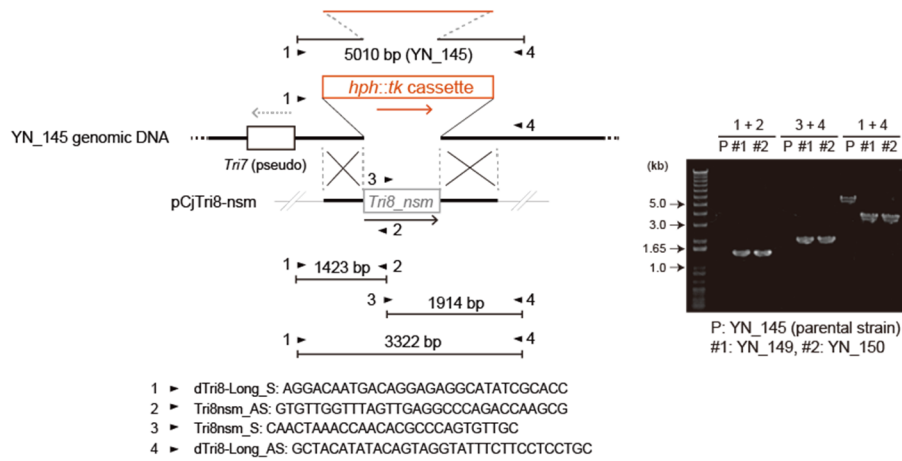
multiple genetic manipulations for the overproduction of 15-ADON. The previously described strain  $\Delta Tri6$  tk (YN\_001) contains a hygromycin B phosphotransferase gene translationally fused to a herpes simplex virus thymidine kinase gene (*hph::tk*) [9, 10]. This strain was used to generate a marker-free *Tri6*-overexpressor strain YN\_004 ( $\Delta Tri6$  /  $P_{TEF}::Tri6$ ) by conditional negative selection against the *hph::tk* cassette with 2'-deoxy-5-fluorouridine (5-FdU). In strain YN\_004, a strong *Aspergillus nidulans* *TEF1 $\alpha$*  promoter fragment [9] is inserted upstream of the *Tri6* coding region. The promoter disruptants YN\_112 ( $\Delta Tri6$  /  $P_{TEF}::Tri6$ ,  $\Delta P_{Tri10}$  /  $P_{GPD}::hph::tk$ ) and YN\_109 ( $\Delta P_{Tri10}$  /  $P_{GPD}::hph::tk$ ) were obtained from YN\_004 and the WT strains, respectively, by positive selection against *hph::tk* with hygromycin B. Marker-free transformants YN\_120 ( $\Delta Tri6$  /  $P_{TEF}::Tri6$ ,  $\Delta P_{Tri10}$  /  $P_{GPD}$ ) and YN\_116 ( $\Delta P_{Tri10}$  /  $P_{Tri10}$  +  $P_{TEF}$ ) were obtained from YN\_112 and YN\_109, respectively, by the negative selection. (B) Generation of strains YN\_120 and YN\_116. YN\_004 was created from YN\_001 following transformation with pCjTefTri6, where 5-FdU resistant transformants with double crossover homologous recombination were negatively selected against *hph::tk*. YN\_112 and YN\_109 were created from YN\_004 and the WT strains, respectively, by positive selection with hygromycin B following transformation with pdTri10p-hphtk. YN\_120 and YN\_116 were created by negative selection with 5-FdU, respectively, following transformation of YN\_112 with pCjPgpdTri10 and YN\_109 with pCjPtefTri10. (C) Production of trichothecenes by the transgenic strains. The WT (JCM 9873), YN\_004, YN\_116, and YN\_120 strains were cultured on 100 mL of YS\_60 (0.1% [w/v] Bacto™ yeast extract, 6% [w/v] sucrose) and YG\_60 (0.1% [w/v] Bacto™ yeast extract, 6% [w/v] glucose) medium in a 300 mL-Erlenmeyer flask with gyratory shaking at 25°C. In contrast to the WT and *Tri6*-overexpressor YN\_004 strains, YN\_116 produced 15-ADON on YG\_60 medium that does not contain sucrose. The 15-ADON production was further enhanced in the culture of YN\_120, in which both *Tri6* and *Tri10* were constitutively overexpressed. YN\_120 also produced 15-ADON on YG medium (0.5% [w/v] Bacto™ yeast extract, 2% [w/v] glucose) that contains a lower amount of glucose and a higher amount of yeast extract. (D) Sequences and maps of disruption and replacement vectors.

(A)

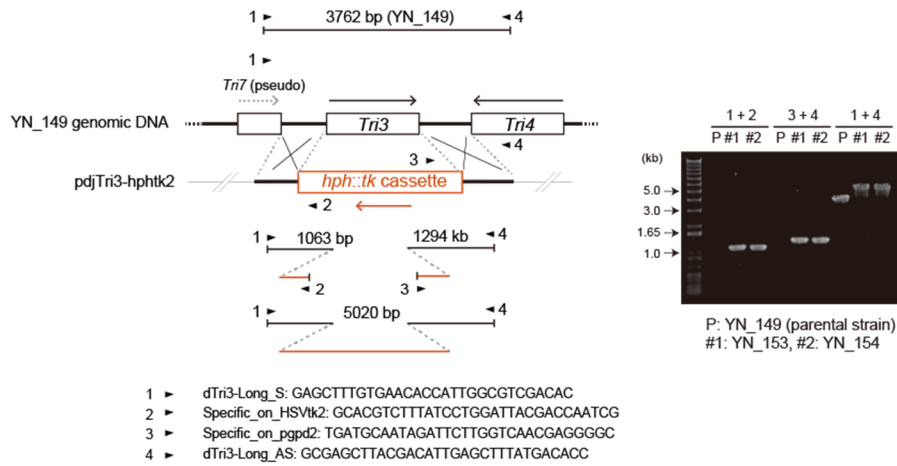


(B)



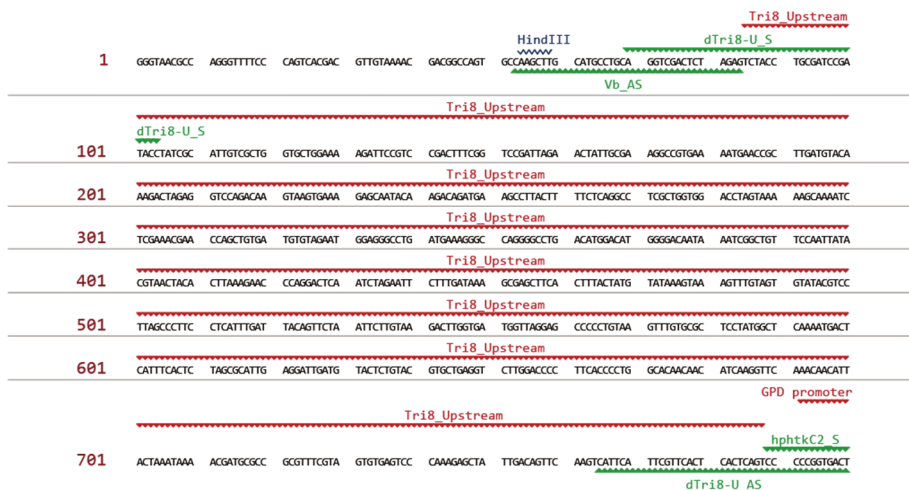


YN149, YN150



YN\_153, YN\_154

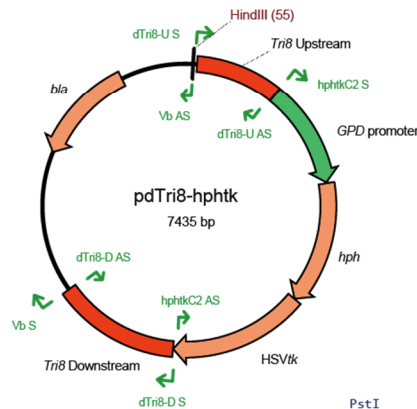
(C)





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	dTri8-U_AS										
901		GPD promoter									
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1001		GPD promoter									
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1201		GPD promoter									
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1501		GPD promoter									
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1601		GPD promoter					hph				
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1701		hph									
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1901		hph									
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2001		hph									
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2101		hph									
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2401		hph									
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3701		HSVtk					dTri8-D_S				
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										hphTkC2_AS	
3801		Tri8 Downstream									
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		hphTkC2_AS									
3901		Tri8 Downstream									
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4001		Tri8 Downstream									
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4101		Tri8 Downstream									
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4201		Tri8 Downstream									
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4301		Tri8 Downstream									
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4401		Tri8 Downstream									
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4501		Tri8 Downstream									
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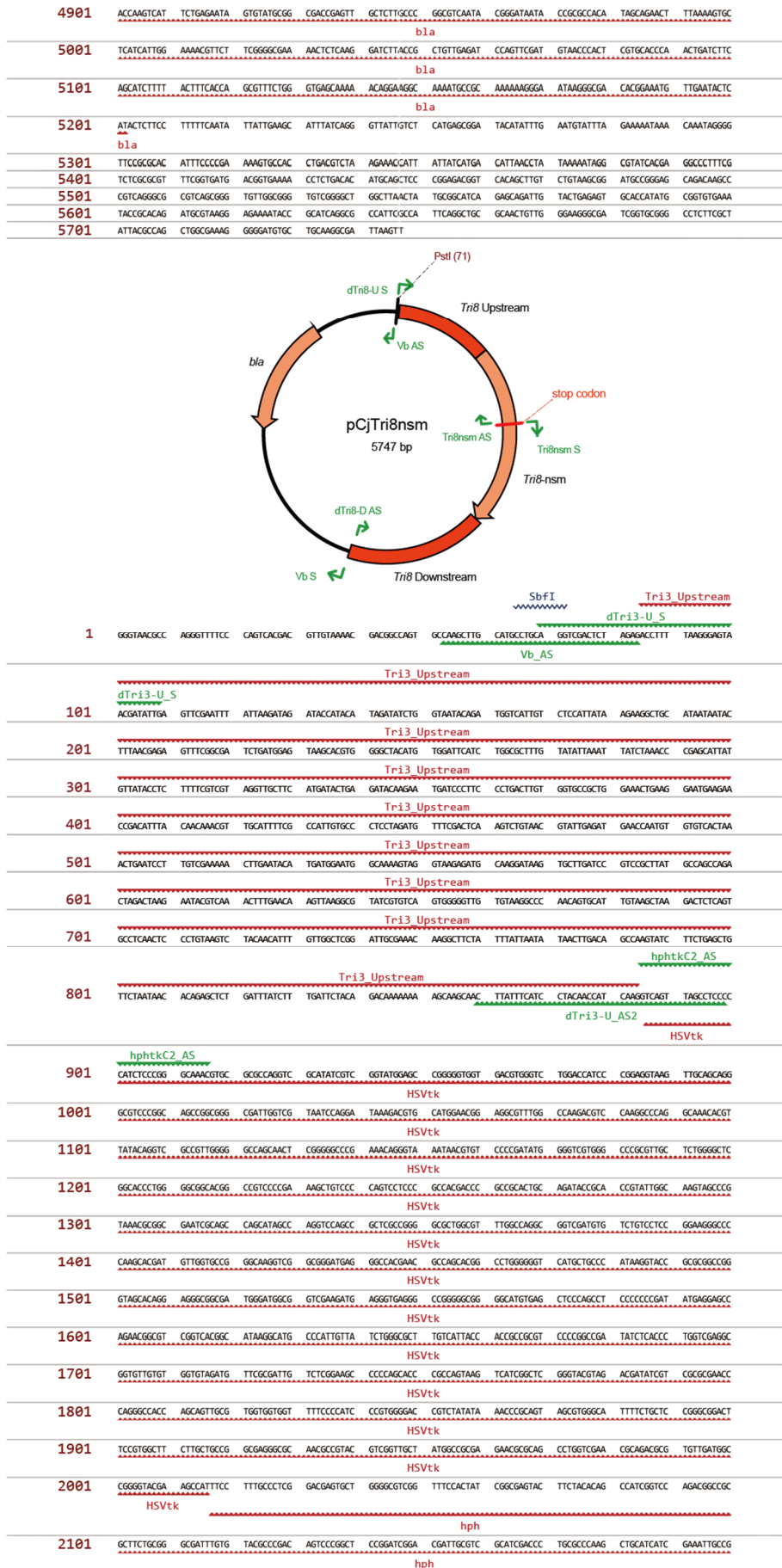
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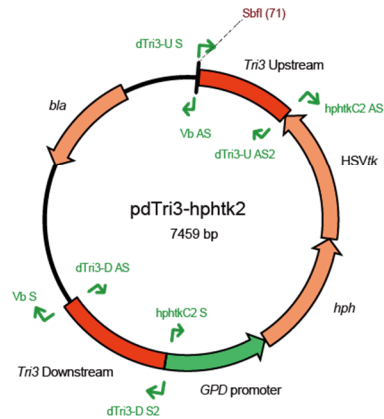






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	Tri3 Downstream
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4501	ACATAGTGG CCTTCAGAG TAACTACAT GTCCAGACA GTTGATGTAT GTAAATATA CTTAGCTTA TTCTAGTAA AGCATATTA TTAGTAATGC
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4601	AGAGAGACT TCCACAGGA GTTAGTTTGA TAGAATGTC ACAAGTATC GGTCTGTAT TCGTCCACA ATACATTGCC TGAGTGAAT AGTGTATAA
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4701	TTGACGAGC TCTAATAGG GTACAATTA TTTTACGTC ASATGTGCT ACTATGACT TCCATCTTG CATACAAAA ATGAGTGA AAGAGAAC
	Vb_S
4801	AGATTCGAT GCCGATACG TGAGTTATCC AGTAAGTAT TGAGACAGT TGTGATGC GCCGGGTAC CGAGCTCGAA TTCGTAATCA TGSTCATAG
	dTri3-D_A5
4901	TGTTCTCTG GTGAATTGT TATCCGCTCA CAATTCCACA CAACATAGCA GCGGAGACA TAAAGTTGAA AGCTGGGGT GCCTAATGAG TGAGCTAAT
5001	CACATTAAAT GGTGTGCGCT CACTGCGCG TTTCCAGTCG GGAACCTGT GTGCCAGCT GCATTAAATGA ATCGGCCAAC GCGCGGGAG AGGCGTTTG
5101	CGATTGGGC GCTCTCCGC TTCTCTGCT ACTGACTCGC TGCGCTCGT CGTTCGGTG CCGGAGCGG TATCAGCTCA CTCGAAGCG GTAATACGT
5201	TATCCACAGA ATCAGGGAT AACGAGAA AGAATGTG ASCAAAAGC CAGAAAGG CAGGAAACG TAAAAGGCC GCGTCTGTC GGTTTTCCA
5301	TAGGCTCCG CCCCCTGAGC AGCATCACA AAATCGAGC TCAAGTCAGA GGTGGCGAAA CCGCAGAGGA CTATAAGAT ACCAGCGTT TCCCCCTGA
5401	AGCTCCCTG TGCGCTCTCC TGTTCCGACC CTGCGCTTA CCGGATACCT GTCCGCTTT CTCCCTTCGG GAAGGTGGC GCTTTCTCAT AGCTCACGCT
5501	GTAGGTATCT CAGTTCGGTG TAGTGTGTC GCTCCAAGCT GGGCTGTGT CACGAACCC CGTTTCAGCC CGAGCGCTG GCTTATCCG GTAACATCG
5601	TCTTGATCC AACCGGTAA GACAGACTT ATCGGCACTG GCAGCAGCA CTGTAACAG GATTAGCAGA GCGAGGTATG TAGGCGTGC TACAGATTCT
5701	TTGAGTGGT GGCCTAACTA CCGCTACACT AGAAGGAGC TATTGTGAT CTGCGCTCTG CTGAAGCCAG TTACTCTCG AAAAAGATT GGTAGCTCT
5801	GATCCGCAA ACAACACCC GCTGGTAGCG GTGGTTTTT TGTGTGCAAG CAGCAGATTA CCGCAGAAA AAAAGSATCT CAAGAGATC CTTTGTACTT
5901	TTTACGGGG CTGACGCTC AGTGAGCA AACTCAGCT TAAGGATTG TGTCTAGAG ATTATCAAA AGATCTTCA CCTAGATCT TTTAAATTA
6001	AAATGAGTT TTAATCAAT CTAAGTATA TATGATGAA CTTGCTGTA CAGTTACAA TGCTTAATCA GTGAGGCACT TATCTCAGCG ATCTGTCTAT
	bla
6101	TTGCTTATC CATAGTGGC TGACTCCCG TCGTGTAGT AACTACGATA CCGGAGGCT TACCATCTGG CCCCAGTGT GCAATGATAC CCGAGAGCC
	bla
6201	ACGCTCACC GCTCCAGATT TATCAGCAAT AAACAGCCA GCCGGAAGG CCGAGCGAG AAGTGGTCT GCAACTTAT CCGCTCCAT CCAGTCTATT
	bla
6301	AATTGTTGC GGGAGCTAG AGTAAGTAG TCGCCAGTTA ATAGTTTGG CACGTTGTT GCCATTGCTA CAGGATCTGT GGTGTACGC TCGTCTTTG
	bla
6401	GTATGCTTC ATTACGCTC GGTTCGCAAC GATCAGGCG AGTTACATGA TCCCCATGT TGTGCAAAA AGCGGTTAG TCTTCTGCTC CTCCGATCT
	bla
6501	TGTCAAGT AAGTTGCCG CAGTGTATC ACTCATGTT ATGGCAGCAC TGCAATTC TCTTACTGC ATGCCATCG TAAGATGCT TCTGTGACT
	bla

6601	GGTGAGTACT	CAACCAAGTC	ATTCTGAGAA	TAGTGTATGC	GGCGACCGAG	TTGCTCTTGC	CCGGCGTCAA	TACGGGATAA	TACCGCGCCA	CATAGCAGAA
	bla									
6701	CTTTAAAGT	GCTCATCATT	GGAAACGTT	CTTCGGGGCG	AAAACCTCA	AGGATCTTAC	CGCTGTTGAG	ATCCAGTTCC	ATGTAACCCA	CTCGTGCACC
	bla									
6801	CACTGATCT	TCAGCATCTT	TTACTTTTAC	CAGCGTTTCT	GGGTGASCAA	AACAGGAGAG	GCAAAATGCC	GCAAAAGAG	GAATAAGGCG	GACACGGAAA
	bla									
6901	TGTTGAATAC	TCATACCTTT	CTTTTTCAC	TATTATTGAA	GCATTATCA	GGGTTATTGT	CTCATGAGCG	GATACATATT	TGAATGTATT	TAGAAAAATA
	bla									
7001	AACAATAGG	GGTCCGCGC	ACATTCCCC	GAAAAGTGGC	ACCTGACGTC	TAAGAAACCA	TTATTATCAT	GACATTAAAC	TATAAAAAATA	GGCGTATCAC
7101	GAGGCCCTTT	CGTCTCGCGC	GTTTCGGTGA	TGACGGTGA	AACCTCTGAC	ACATGCAGCT	CCCGGAGACG	GTCACAGCTT	GTCGTGTAAGC	GGATGCCGGG
7201	AGCAGACAG	CCCGTCAGGG	CGCGTCAGCG	GGTGTGGCG	GGTGTGCGGG	CTGGCTTAAC	TATGCGGCAT	CAGAGCAGAT	TGTACTGAGA	GTGCACCATATA
7301	TGCGGTGTGA	AATACCGCAC	AGATGCGTAA	GGAGAAAAATA	CCGCATCAGG	CGCCATTCCG	CATTGAGGCT	GCGCAACTGT	TGGGAAGGGC	GATCGGTGCG
7401	GGCTCTCTCG	CTATTACGCC	AGCTGGCGAA	AGGGGGATGT	GCTGCANGGC	GATTAAAGTT				



**Supplementary Figure S7.** Transgenic *F. graminearum* strains used for the investigation of the effect of *Tri6* and *Tri10* overexpression on 15-deCAL accumulation. (A) Pedigree of transformants with multiple genetic manipulations for the generation of a *Tri3*-disrupted strain, YN\_153, with overexpression of *Tri6* and *Tri10*. The gene disruptants, YN\_145 ( $\Delta Tri6$  /  $P_{TEF}::Tri6$ ,  $\Delta P_{Tri10}$  /  $P_{GPD}$ ,  $\Delta Tri8$  /  $P_{GPD}::hph::tk$ ) and YN\_153 ( $\Delta Tri6$  /  $P_{TEF}::Tri6$ ,  $\Delta P_{Tri10}$  /  $P_{GPD}$ ,  $\Delta Tri8$  /  $Tri8_{nsm}$ ,  $\Delta Tri3$  /  $P_{GPD}::hph::tk$ ), were obtained from the marker-free transformants YN\_120 (Supplementary Figure S6) and YN\_149 ( $\Delta Tri6$  /  $P_{TEF}::Tri6$ ,  $\Delta P_{Tri10}$  /  $P_{GPD}$ ,  $\Delta Tri8$  /  $Tri8_{nsm}$ ), respectively, by positive selection. YN\_149 was created from YN\_145 by negative selection. (B) Generation of strain YN\_153. YN\_145 was created from YN\_120 by positive selection with hygromycin B following transformation with pdjTri8-hphtk. YN\_149 was created from YN\_145 by negative selection with 5-FdU following transformation with pCjTri8-nsm. YN153 was created from YN\_149 by positive selection with hygromycin B after transformation with pdjTri3-hphtk2. (C) Sequences and maps of disruption and replacement vectors.

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