

## Electrical Supporting Information

### Design of a self-sufficient hydride-shuttling cascade for concurrent bioproduction of 7,12-dioxolithocholate and L-*tert*-leucine

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# 1. Experimental Section

## 1.1 General information

Cholic acid (3 $\alpha$ ,7 $\alpha$ ,12 $\alpha$ -trihydroxy-5 $\beta$ -cholanic acid; CA), deoxycholic acid (3 $\alpha$ ,12 $\alpha$ -dihydroxy-5 $\beta$ -cholanoic acid; DCA), and chenodeoxycholic acid (3 $\alpha$ ,7 $\alpha$ -dihydroxy-5 $\beta$ -cholanic acid; CDCA) were purchased from Aladdin Reagents Co., Ltd. (Shanghai, China). Trimethylpyruvic acid (TMP) and L-*tert*-leucine were obtained from Adamas Reagent Co., Ltd. (Shanghai, China). Standards of 7-oxo-deoxycholic acid (3 $\alpha$ ,12 $\alpha$ -dihydroxy-7-oxo-5 $\beta$ -cholanic acid; 7-oxo-DCA), 12-oxo-chenodeoxycholic acid (3 $\alpha$ ,7 $\alpha$ -dihydroxy-12-oxo-5 $\beta$ -cholanic acid; 12-oxo-CDCA), and 7,12-dioxo-lithocholic acid (3 $\alpha$ -hydroxy-7,12-dioxo-5 $\beta$ -cholanoic acid; 7,12-dioxo-LCA) were enzymatically synthesized in our previous work.<sup>S1</sup> Unless otherwise specified, all other chemicals and reagents were obtained commercially and were of reagent grade. PrimeSTAR™ HS, Taq DNA polymerase, and restriction endonucleases were all purchased from Takara Biotechnology Co., Ltd. (Dalian, China). The Gibson assembly kit was purchased from Vazyme Biotech Co., Ltd. (Nanjing, China). *E. coli* DH5 $\alpha$  and *E. coli* BL21 (DE3) were used as hosts for gene cloning and protein expression, respectively. Plasmids pET-28a (+), pETDuet-1, and pRSFDuet-1 (Novagen, Shanghai, China) were used for the heterogeneous expression of recombinant protein. The recombinant plasmids pET28a (+)-*Ec*7 $\alpha$ -HSDH, pET28a (+)-*Rr*12 $\alpha$ -HSDH and pET28a (+)-*Es*LeuDH were previously constructed by our laboratory.<sup>S1-S3</sup> The molar yield of 7,12-dioxo-LCA were determined by HPLC analysis using a Shimadzu LC-2010A high performance liquid chromatography equipped with an Hypersil ODS2 column (250 mm  $\times$  4.6 mm, 5  $\mu$ m particle size, Elite) and a UV detector. <sup>1</sup>H NMR and <sup>13</sup>C NMR were measured on a Bruker Ascend spectrometer.

## 1.2 Gene sequences

### Gene sequence of *Rr*12 $\alpha$ -HSDH (GenBank: QDC17258.1)

```
ATGAAACTGCGCGGAAGACCGCCGTCGTCACCGGCGGTGCGGGCGGGATCGGCCGCGCGGTGACCCGCG  
TGTTTCGTCGCGAGGGCGCCCGGTGCTGTTTCGTCGACGTCGACGACGATCGGGGGCGCGCGCTCGAGTCC  
GAGCTGACCGGGGCCGCGGTGAGGCGAAGTTCCTGCAGGCCGACATCTCCCGCGCGAGAGCGCGGACC  
AGATCCGCGACGCCCGCTCGCGGCGTTCGGCGGCATCGACATCTGGTCAACAACGCGCACGCGTCGCGC
```

CAGGCACTGCTGGTTCGAGCACACCCCGGAGATGTTTCGAGCTGTCGTTTCGGCACGGGGTTCTACCCACCGTG  
CACCTCATGCAGGCCTGCTACCCGCAGCTCAAGCAGGCCCGGGTTCCGTCGTCAACTTCGGCTCCGGGTCC  
GCCCTCGACGGCATGCCGACGCAGACGTTCGTACGCGCGGCGAAGGAGGCGATCCGGGCGGTTCAGCCGGG  
TGGCCGCGAACGAATGGGCCGCGACGGCATCCGCGTCAACGTCGTGTGCCCGTTTCGCCGCGACCGAAGGC  
GTGCAGGCCTGGCAGCAGGCGTTCCCCGACCGGGCGGCCCGCGGGCGAAGGTGCCGTTGCAGCGCA  
TCGGCGACCCGGAGACGGACATCGCGCCGGTGGTGGTTCCTCGCCTCCGACGACTCGAAGTACATGACG  
GGCAGACGCTGATGGCCGACGGGGGCAGCATCAAGCTGCGGTGA

**Gene sequence of *Ec7 $\alpha$ -HSDH* (GenBank: BAA01384.1)**

GTGTTTAATTCTGACAACCTGAGACTCGACGGAAAATGCGCCATCATCACAGGTGCGGGTGCAGGTATTGGT  
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AACCATGTTGTAGACGAAATTCAACAACCTGGGTGGTCAGGCATTTGCCTGCCGTTGTGATATACTTCCGAAC  
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CTGGGCCAACCGCAAGATATTGCTAACGCAGCGCTGTTCTTTGCTCGCCTGCTGCGAGCTGGGTAAGCGGA  
CAAATTCTCACCGTCTCCGGTGGTGGGGTACAGGAGCTCAATTA

**Gene sequence of *EsLeuDH* (GenBank: ACB60396.1)**

ATGGTTGAAACAAACGTAGAAAGCACGATTCAGTATTTTCGAAACGATGGCAATGGAAGATTACGAACAAGTC  
GTATTTTGTACGATAAAGTCTCAGGATTAAGGGCATTATCGCGATTCATGATACGACACTCGGACCAGCA  
CTCGGCGGACTCCGTATGTGGAACCTATGCGTCTGACGAGGAAGCATTGATCGACGCGCTTCGTTTGGCAAAA  
GGCATGACGTATAAAAATGCGGCAGCCGGTCTGAACCTTGGCGGCGGGAAAGCGGTCATCATCGGTGATGC  
GAAAACGCAAAAATCAGAAGCTCTGTTCCGTGCATTCGGTCGTTACGTACAGTCGTTAAACGGACGTTACAT  
CACTGCGGAAGACGTCAACACAACAGTCGCCGACATGGATTATATCCACATGGAAACAGATTTTCGTAACCGG  
TGTCAGCCCGGCATTCGGATCAAGCGGCAATCCGTCACCAGTCACGGCTTATGGCGTTTACCGCGGAATGAA  
GGCAGCCGCTAAAGAAGTATATGGCACAGATTCCTCGGAGGAAAAACAGTTGCGATTCAAGGTGTTGGTA

ACGTTGCTTTCAACCTATGCCGTCACTTGCATGAAGAAGGCGCAAAATTGATTGTCACAGACATCAATCAAGA  
 TGCATTACGCCGTGCAGAAGAAGCGTTTGGCGCTCTCGTCGTCGGACCGGATGAAATTTACAGCGTCGATGC  
 CGATATCTTTGCGCCGTGTGCCTTAGGTGCGACATTGAACGATGAGACGATTCCACAACCTGAAAGTGAAAAT  
 CATTGCCGGAGCAGCAAACAACCAACTCAAAGAAGATCGTCACGGAGATATGCTCCAGGAACGCGGTATTTT  
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 ACCATCTGCCGACTTACCGGGCAGCAGAGAAGATGGCAGAAGAACGGATCGCGACAATGGGCAGTGCCCCG  
 CAGCCAGTTCTTACGCCGGGATAAAAACATTTTAGGATCACGCGGTAA

### 1.3 Calculation of E factors

E factors were calculated as  $\text{Weight}_{\text{waste}}/\text{Weight}_{\text{product}}$ . The calculations did not take into account the waste generated for the preparation of the catalysts.

#### (1) 12 $\alpha$ -HSDH/7 $\alpha$ -HSDH/LeuDH system, 100 mM CA, 250 mM TMP, 50 mL (this work).

	Compound	Weight (mg)	Quantity (mg per mg products)
Product	7,12-dioxo-LCA + L- <i>tert</i> -leucine	1600 + 850	
	CA	400	0.16
	Trimethylpyruvic acid	750	0.31
	Cells	1000	0.41
Waste	Ammonium chloride	2650	1.08
	H <sub>2</sub> O	50000	20.4
	1 M HCl	1200	0.49
	Ethanol	39500	16.1
E factor	Including water		38.9
	Excluding water		18.5

#### (2) O<sub>2</sub>/NOX system, 10 mM CA, 75 mL, flow reactor (ref. 43).

	Compound	Weight	Quantity
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		(mg)	(mg per mg product)
Product	12-oxo-CDCA	297	
	CA	3	0.01
	NAD <sup>+</sup>	35.5	0.12
	12 $\alpha$ -HSDH	0.26	0.001
Waste	NOX	37.5	0.13
	MeOH	5925	19.9
	50 mM KPi pH 8.0	640	2.15
	H <sub>2</sub> O	67500	227.3
E factor <sup>a</sup>	Including water		249.6
	Excluding water		22.3

<sup>a</sup> The calculations did not take into account the downstream processes.

### (3) Syringaldazine/MtLac system, 120 mM CA, 410 mL (ref. 44).

	Compound	Weight (mg)	Quantity (mg per mg product)
Product	12-oxo-CDCA	18400	
	CA	1600	0.09
	NAD <sup>+</sup>	136	0.007
	Syringaldazine	21	0.001
	12 $\alpha$ -HSDH	5.33	0.0003
Waste	<i>MtLac</i>	139	0.008
	DMSO	45100	2.45
	100 mM KPi pH 7.2	6700	0.36
	H <sub>2</sub> O	369000	20.1
	Diethyl ether	710000	38.9
E factor	Including water		61.6
	Excluding water		41.5

**(4) Pyruvate/LDH system, 200 mM CA, 100 mL (ref. 36).**

	Compound	Weight (mg)	Quantity (mg per mg product)
Product	12-oxo-CDCA	6800	
	CA	1200	0.18
	Pyruvate	2640	0.39
	NAD <sup>+</sup>	35.5	0.005
Waste	Cells	1000	0.15
	100 mM KPi pH 8.0	1720	0.25
	H <sub>2</sub> O	100000	14.7
	1 M HCl	4800	0.71
	Ethanol	79000	11.6
E factor	Including water		28.0
	Excluding water		13.3

**(5) Acetone/Alcohol Dehydrogenase (ADH) system, 100 mM CA, 10 mL (ref. 42).**

	Compound	Weight (mg)	Quantity (mg per mg product)
Product	12-oxo-CDCA	170	
	CA	230	1.35
	Acetone	1970	11.6
	NADP <sup>+</sup>	0.75	0.004
	12 $\alpha$ -HSDH	16.7	0.1
Waste	Dithiothreitol	1.6	0.01
	Bovine serumalbumin	60	0.35
	100 mM KPi pH 8.0	172	1.01
	H <sub>2</sub> O	10000	58.8
	Ethyl acetate	9000	52.9

E factor	Including water	126.2
	Excluding water	67.4

**(6)  $\alpha$ -Ketoglutarate/Glutamate Dehydrogenase (GluDH) system, 100 mM CA, 70 mL (ref. 39).**

	Compound	Weight (mg)	Quantity (mg per mg products)
Product	12-oxo-CDCA	2464	
Waste	CA	336	0.14
	$\alpha$ -Ketoglutarate	1691	0.69
	NADP <sup>+</sup>	3.5	0.001
	12 $\alpha$ -HSDH	10.8	0.004
	GLuDH	0.58	0.0002
	Dithiothreitol	11	0.004
	Sodium azide	14	0.006
	Ammonium acetate	1348	0.55
	100 mM KPi pH 8.0	1207	0.49
	H <sub>2</sub> O	70000	28.4
	Acetone	276500	112.2
	Ethanolamine	427	0.17
E factor	Including water		142.7
	Excluding water		114.3



## 2. Supporting Tables

**Table S1.** Primers used for constructing recombinant *E. coli* strains.

Primer	Sequence (5'-3')
mcs1-12 $\alpha$ -FP (BamH I )	TCATCACCACAGCCA <u>GGATCC</u> GATGAAACTGCGCGGAAGA
mcs1-12 $\alpha$ -RP (Hind III)	TTAAGCATTATGCGGCCGC <u>AAGCTT</u> TACC GCAGCTTGATGCTGC
mcs1-7 $\alpha$ -FP (BamH I )	TCATCACCACAGCCA <u>GGATCC</u> GATGTTAATTCTGACAACC
mcs1-7 $\alpha$ -RP (Hind III)	TTAAGCATTATGCGGCCGC <u>AAGCTT</u> TTAATTGAGCTCCTGTAC
mcs1-Es-FP (BamH I )	TCATCACCACAGCCA <u>GGATCC</u> GATGGTTGAAACAAACG
Mcs1-Es-RP (Hind III)	TTAAGCATTATGCGGCCGC <u>AAGCTT</u> TTAACCGCGTGATCCT
mcs2-12 $\alpha$ -FP (Nde I )	GTATAAGAAGGAGATATA <u>CATATG</u> ATGAAACTGCGCGGAAGACCGC
mcs2-12 $\alpha$ -RP (Kpn I )	TTTCTTTACCAGACTCGAG <u>GGTACC</u> TACC GCAGCTTGATGCT
mcs2-7 $\alpha$ -FP (Nde I )	GTATAAGAAGGAGATATA <u>CATATG</u> ATGTTAATTCTGACAACCTGAGACTCG
mcs2-7 $\alpha$ -RP (Kpn I )	TTTCTTTACCAGACTCGAG <u>GGTACC</u> TTAATTGAGCTCCTGTA
mcs2-Es-FP(Nde I )	GTATAAGAAGGAGATATA <u>CATATG</u> ATGGTTGAAACAAACGTAGAAGCACG
mcs2-RP(Kpn I )	TTTCTTTACCAGACTCGAG <u>GGTACC</u> TTAACCGCGTGATCCTAAAATGTT
pETDuet-MCS1-FP	ATCGGTGATGTCGGCGATAT
pRSFDuet-MCS1-FP	GTTTTGCGCCATTCGATGGT
Duet-MCS1-RP	GATTATGCGGCCGTGTACAA
Duet-MCS2-FP	TTGTACACGGCCGCATAATC
Duet-MCS2-RP	TGCTAGTTATTGCTCAGCGG

**Table S2.** Strains and plasmids used in this study.

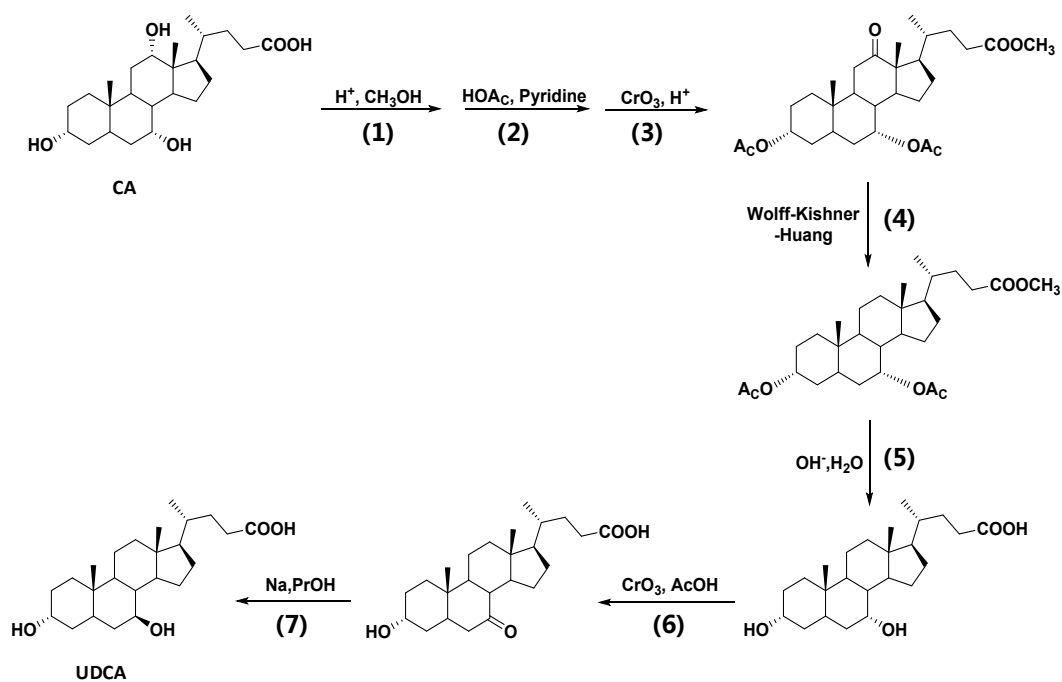
Strain and plasmids	Relevant characteristics	Source
pRSFDuet-1	double T7 promoters, RSF ori, Kan <sup>R</sup>	Novagen
pETDuet-1	double T7 promoters, pBR322 ori, Amp <sup>R</sup>	Novagen
pET28a (+)	T7 promoter, pBR322 ori, Kan <sup>R</sup>	Novagen
pET28a (+)- <i>Rr12α</i> -HSDH	pET28a (+) carrying <i>12α-hsdh</i>	laboratory stock
pET28a (+)- <i>Ec7α</i> -HSDH	pET28a (+) carrying <i>7α-hsdh</i>	laboratory stock
pET28a (+)- <i>EsLeuDh</i>	pET28a (+) carrying <i>leudh</i>	laboratory stock
pRSFDuet- <i>Rr12α</i> -HSDH- <i>EsLeuDh</i>	pRSFDuet carrying <i>12α-hsdh</i> and <i>leudh</i>	this study
pRSFDuet- <i>Rr12α</i> -HSDH	pRSFDuet carrying <i>12α-hsdh</i>	this study
pRSFDuet- <i>Rr12α</i> -HSDH- <i>Ec7α</i> -HSDH	pRSFDuet carrying <i>12α-hsdh</i> and <i>7α-hsdh</i>	this study
pRSFDuet- <i>EsLeuDh</i> - <i>Rr12α</i> -HSDH	pRSFDuet carrying <i>leudh</i> and <i>12α-hsdh</i>	this study
pRSFDuet- <i>EsLeuDh</i>	pRSFDuet carrying <i>leudh</i>	this study
pRSFDuet- <i>EsLeuDh</i> - <i>Ec7α</i> -HSDH	pRSFDuet carrying <i>leudh</i> and <i>7α-hsdh</i>	this study
pETDuet- <i>Ec7α</i> -HSDH	pETDuet carrying <i>7α-hsdh</i>	this study
pETDuet- <i>Ec7α</i> -HSDH- <i>EsLeuDh</i>	pETDuet carrying <i>7α-hsdh</i> and <i>leudh</i>	this study
pETDuet- <i>Rr12α</i> -HSDH	pETDuet carrying <i>12α-hsdh</i>	this study
pETDuet- <i>Rr12α</i> -HSDH- <i>Ec7α</i> -HSDH	pETDuet carrying <i>12α-hsdh</i> and <i>7α-hsdh</i>	this study
pETDuet- <i>EsLeuDh</i>	pETDuet carrying <i>leudh</i>	this study
Strain 1	<i>E. coli</i> BL21(DE3)/(pRSFDuet- <i>Rr12α</i> -HSDH- <i>EsLeuDh</i> and pETDuet- <i>Ec7α</i> -HSDH- <i>EsLeuDh</i> )	this study
Strain 2	<i>E. coli</i> BL21(DE3)/(pRSFDuet- <i>Rr12α</i> -HSDH and pETDuet- <i>Ec7α</i> -HSDH- <i>EsLeuDh</i> )	this study

Strain 3	<i>E. coli</i> BL21(DE3)/(pRSFDuet- <i>EsLeuDH-Rr12<math>\alpha</math>-HSDH</i> and pETDuet- <i>Ec7<math>\alpha</math>-HSDH-EsLeuDH</i> )	this study
Strain 4	<i>E. coli</i> BL21(DE3)/(pRSFDuet- <i>Rr12<math>\alpha</math>-HSDH-EsLeuDH</i> and pETDuet- <i>Ec7<math>\alpha</math>-HSDH</i> )	this study
Strain 5	<i>E. coli</i> BL21(DE3)/(pRSFDuet- <i>EsLeuDH-Rr12<math>\alpha</math>-HSDH</i> and pETDuet- <i>Ec7<math>\alpha</math>-HSDH</i> )	this study
Strain 6	<i>E. coli</i> BL21(DE3)/(pRSFDuet- <i>EsLeuDH</i> and pETDuet- <i>Rr12<math>\alpha</math>-HSDH-Ec7<math>\alpha</math>-HSDH</i> )	this study
Strain 7	<i>E. coli</i> BL21(DE3)/(pRSFDuet- <i>EsLeuDH-Ec7<math>\alpha</math>-HSDH</i> and pETDuet- <i>Rr12<math>\alpha</math>-HSDH</i> )	this study
Strain 8	<i>E. coli</i> BL21(DE3)/(pRSFDuet- <i>Rr12<math>\alpha</math>-HSDH-Ec7<math>\alpha</math>-HSDH</i> and pETDuet- <i>EsLeuDH</i> )	this study

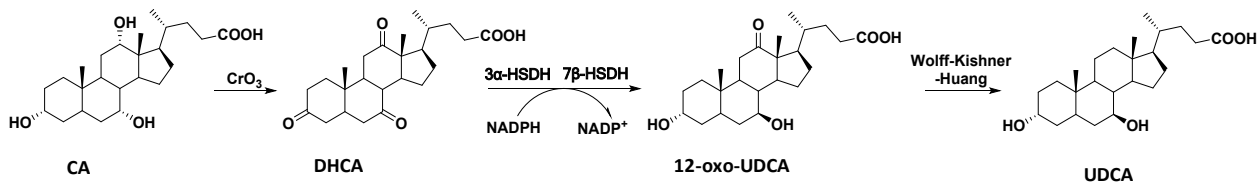
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### 3. Supporting Figures

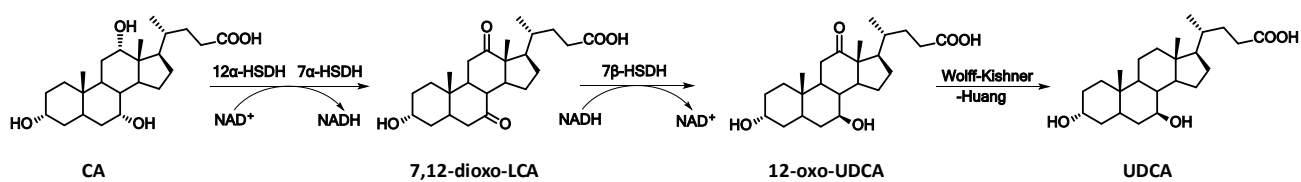
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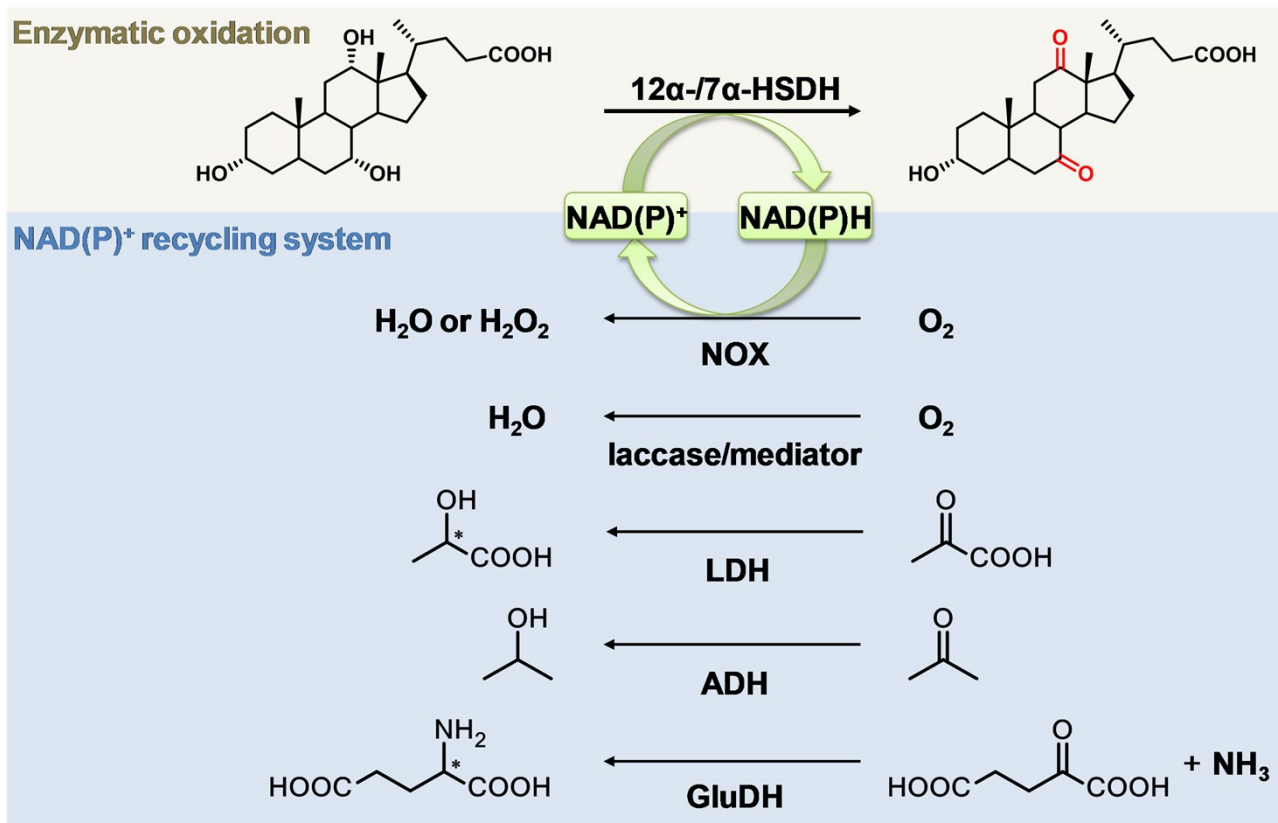
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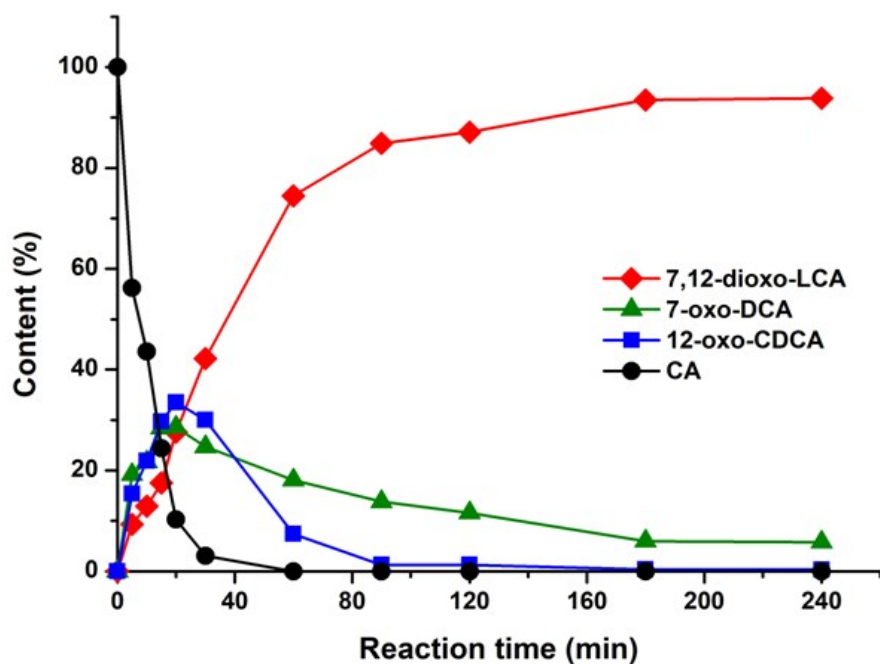
(C)



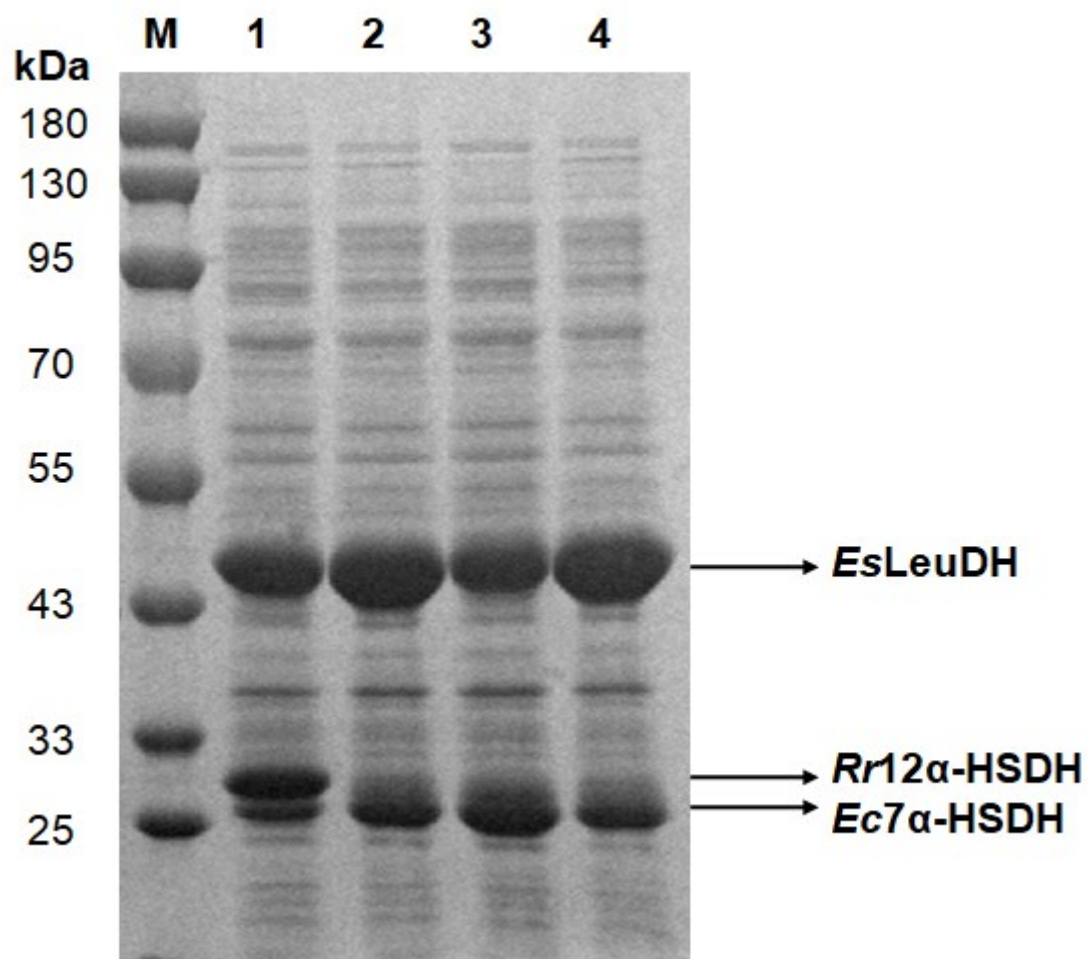
**Figure S1.** Synthesis of UDCA from CA by chemical (A) or chemo-enzymatic methods (B and C).



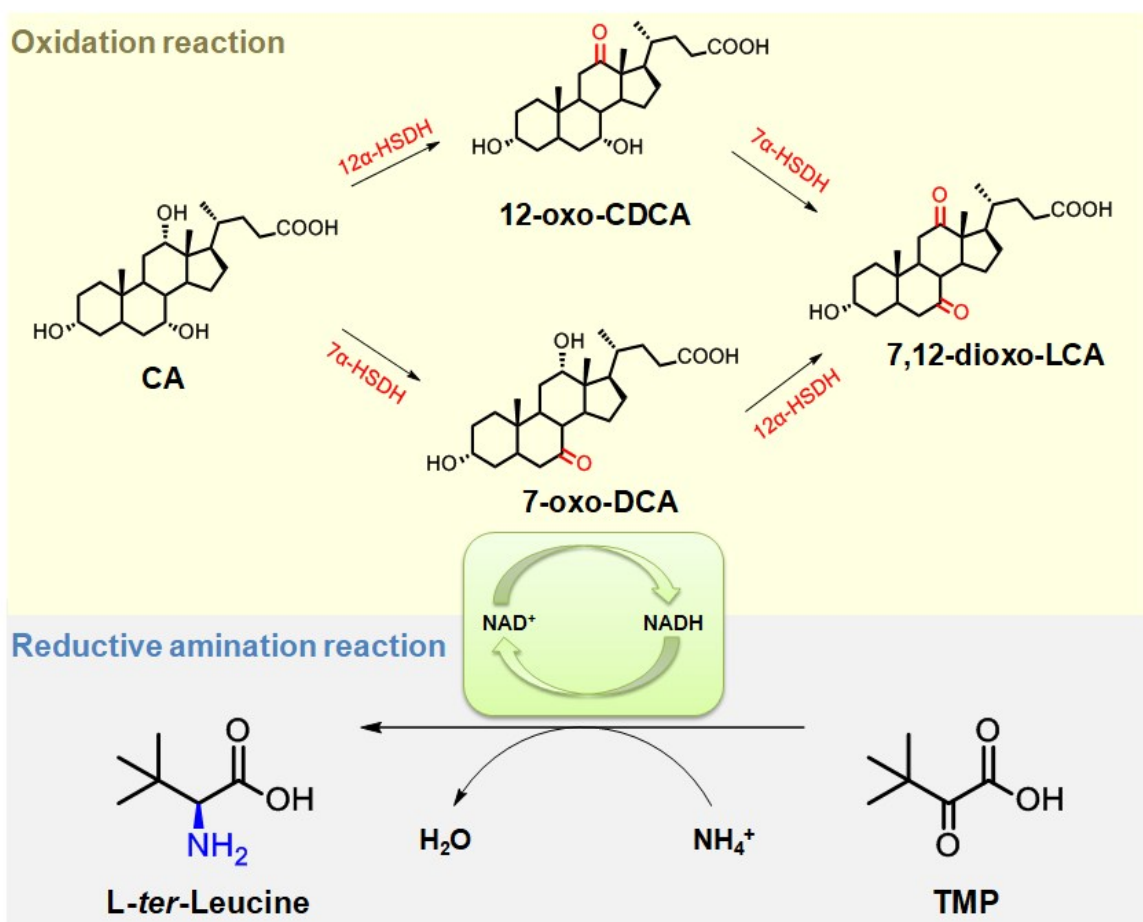
**Figure S2.** Currently available NAD(P)<sup>+</sup>-recycling systems for the biocatalytic oxidation of CA.



**Figure S3.** Reaction progress curves of the self-sufficient hydride-shuttling cascade via *in vitro* biocatalysis. Reaction conditions: 10 mL reaction mixture, containing 25 mM CA, 75 mM TMP, 0.1 mM NAD<sup>+</sup>, lyophilized cell-free extracts of *Rr12α*-HSDH, *Ec7α*-HSDH and *EsLeuD*H (5 U mL<sup>-1</sup> each), and NH<sub>4</sub>Cl/NH<sub>4</sub>OH buffer (1 M, pH 9.5), was incubated at 30°C and 200 rpm, the samples were withdrawn at regular intervals for HPLC analysis.



**Figure S4.** SDS-PAGE analysis of whole-cell proteins expressed in selected *E. coli* strains. Lane M: protein marker (kDa); Lanes 1–4 represent strain 2, strain 4, strain 6 and strain 8, respectively.



**Figure S5.** A detailed synthetic roadmap of the smart self-sufficient hydride-shuttling cascade for concurrently biosynthesis of 7,12-dioxo-LCA and L-tert-leucine.



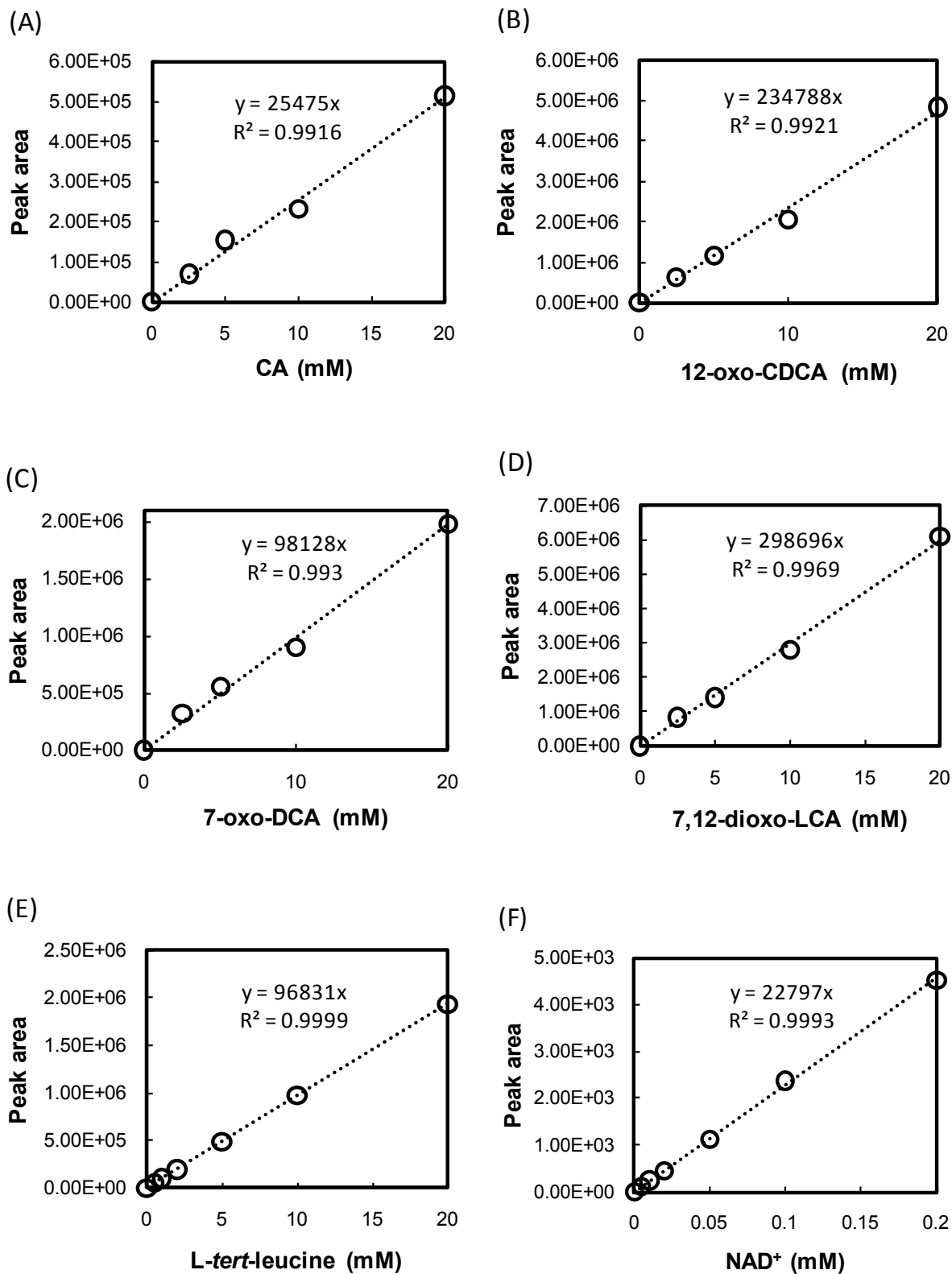
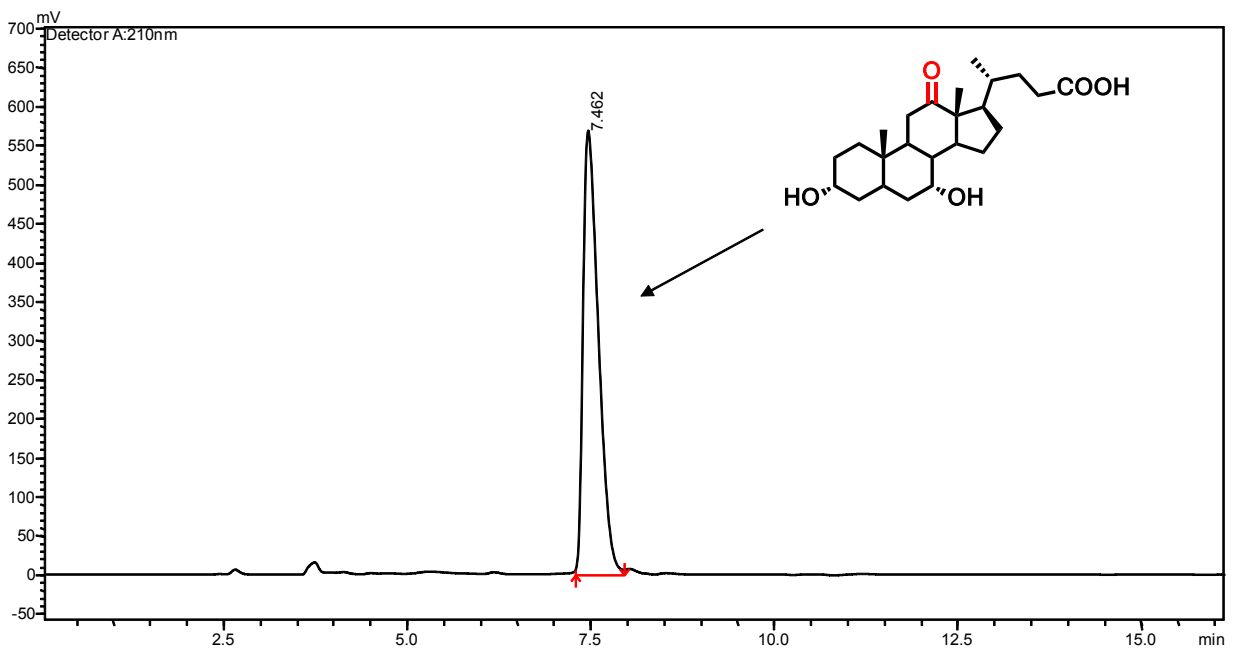
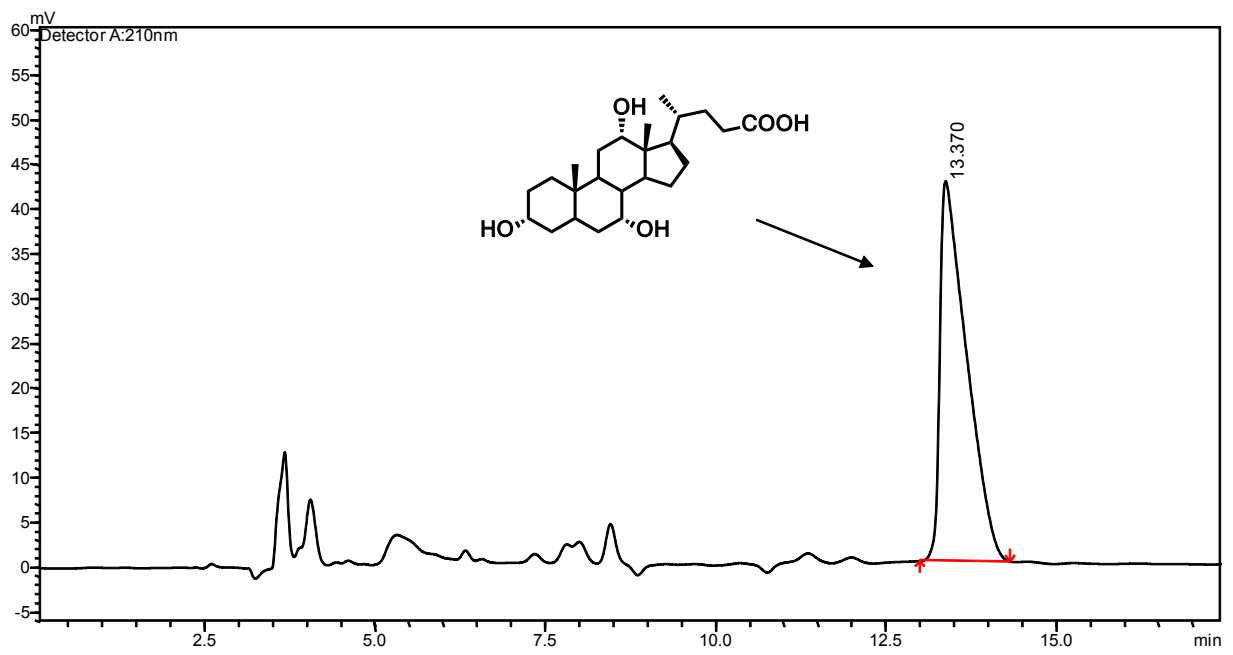
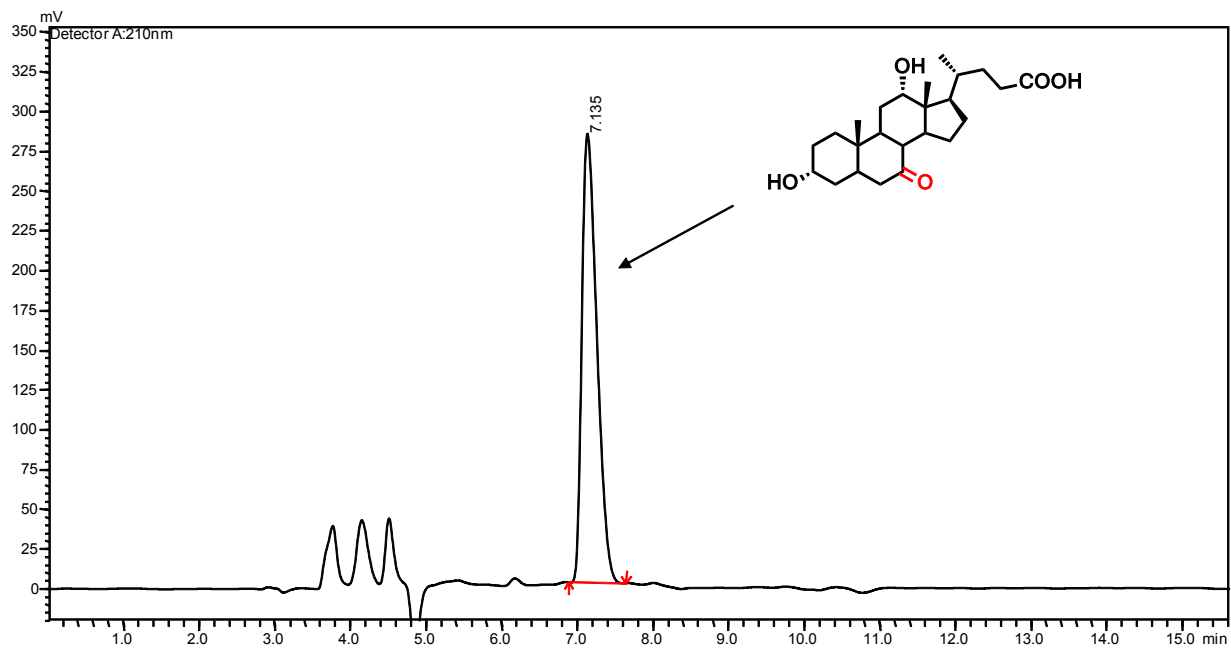


Figure S6. Calibration curves of CA (A), 12-oxo-CDCA (B), 7-oxo-DCA (C), 7,12-dioxo-LCA (D), L-tert-leucine (E), and NAD<sup>+</sup>

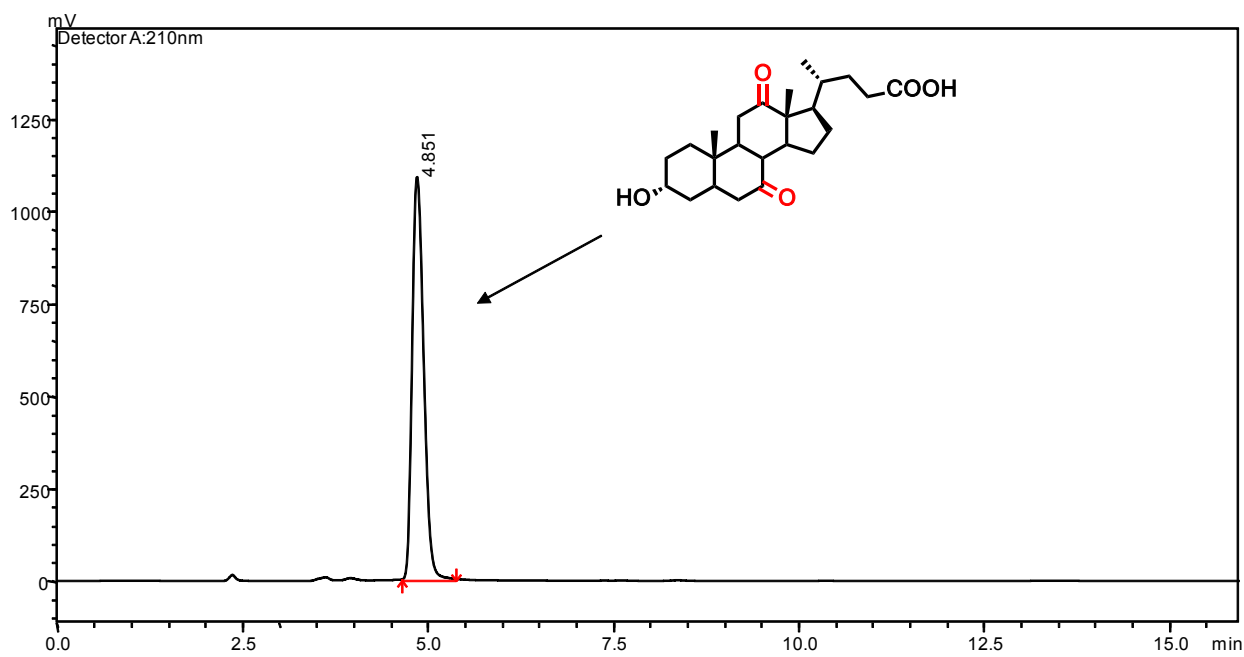
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## 4. HPLC spectra

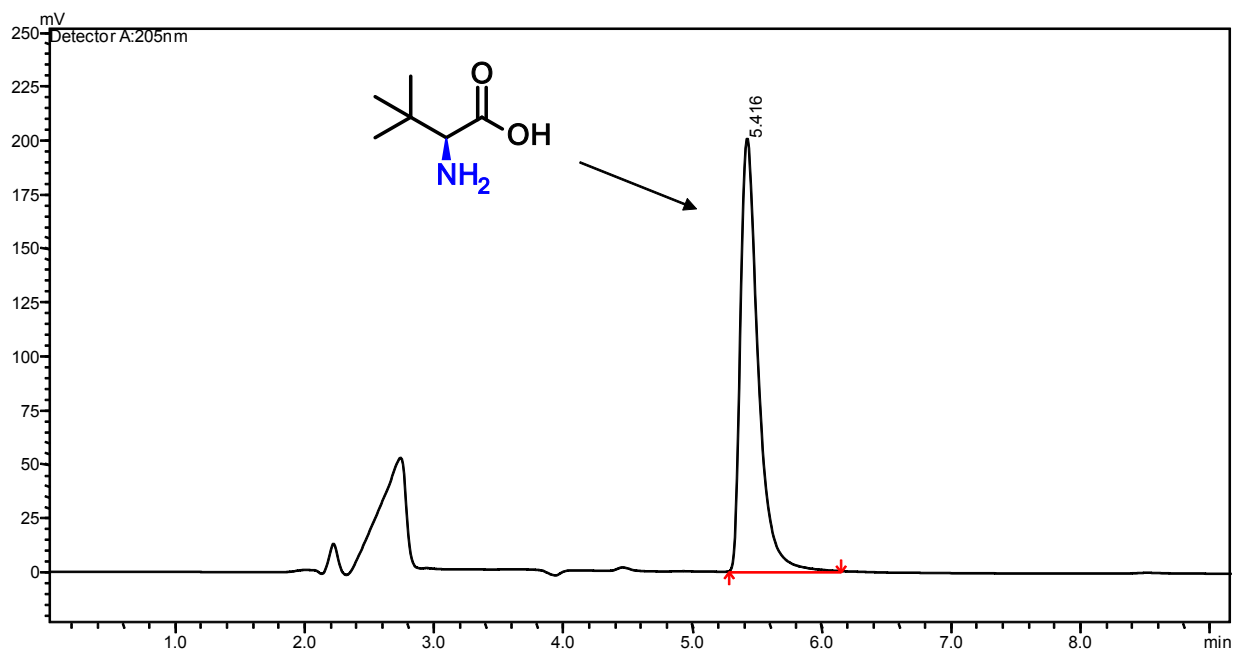




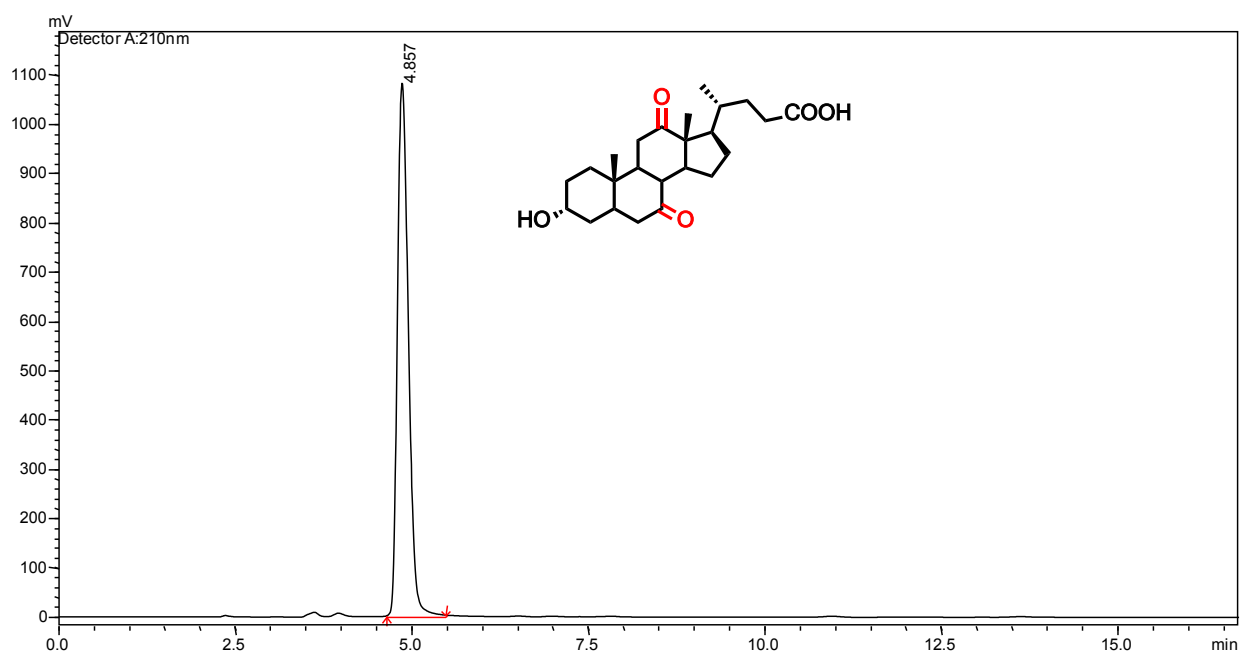
Standard 7-oxo-DCA.



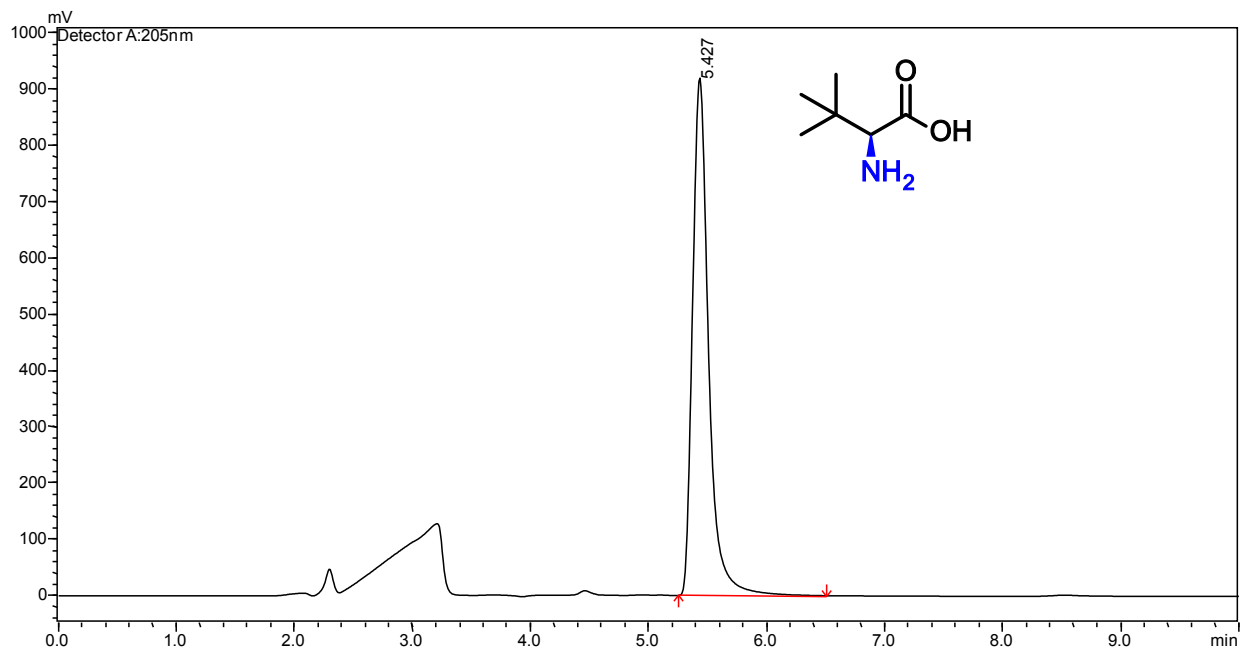
Standard 7,12-dioxo-LCA.



Standard L-*tert*-Leucine.



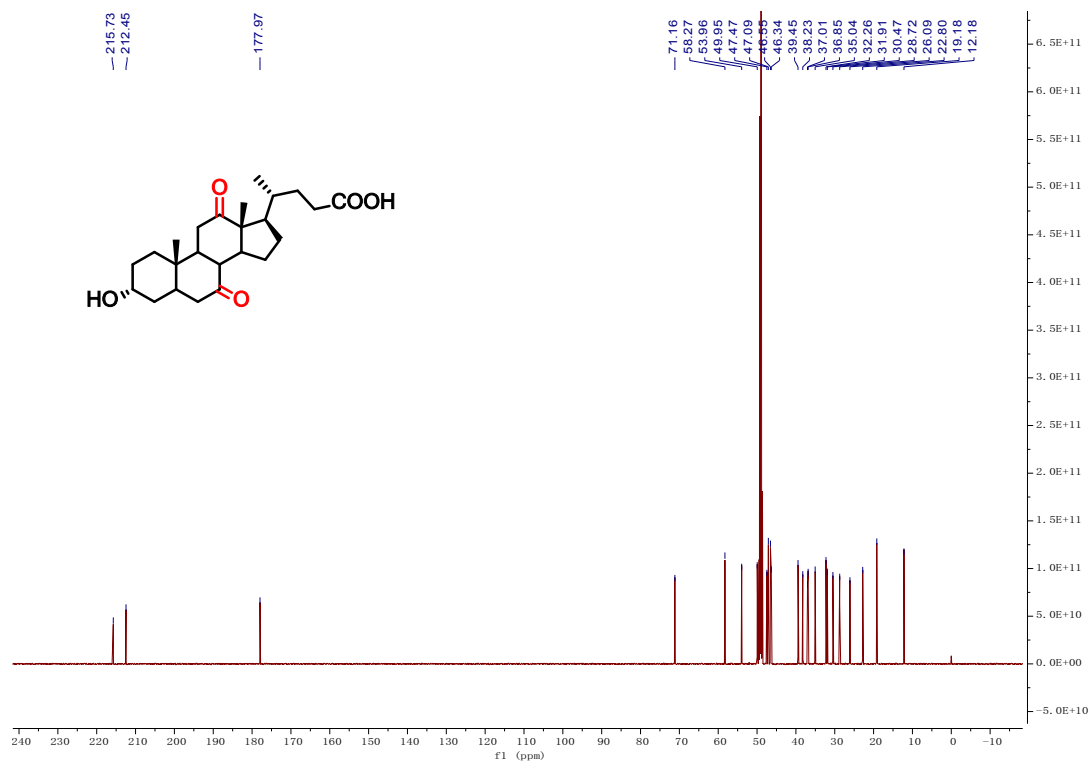
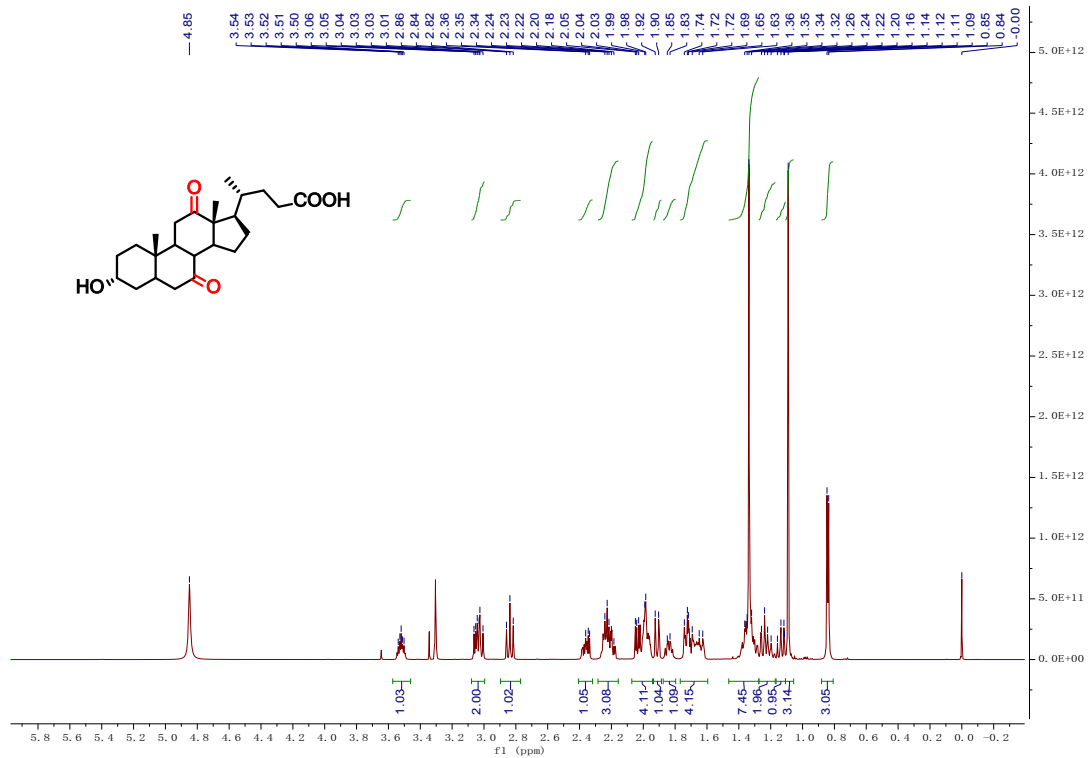
The 7,12-dioxo-LCA prepared by enzymatic cascade reaction.



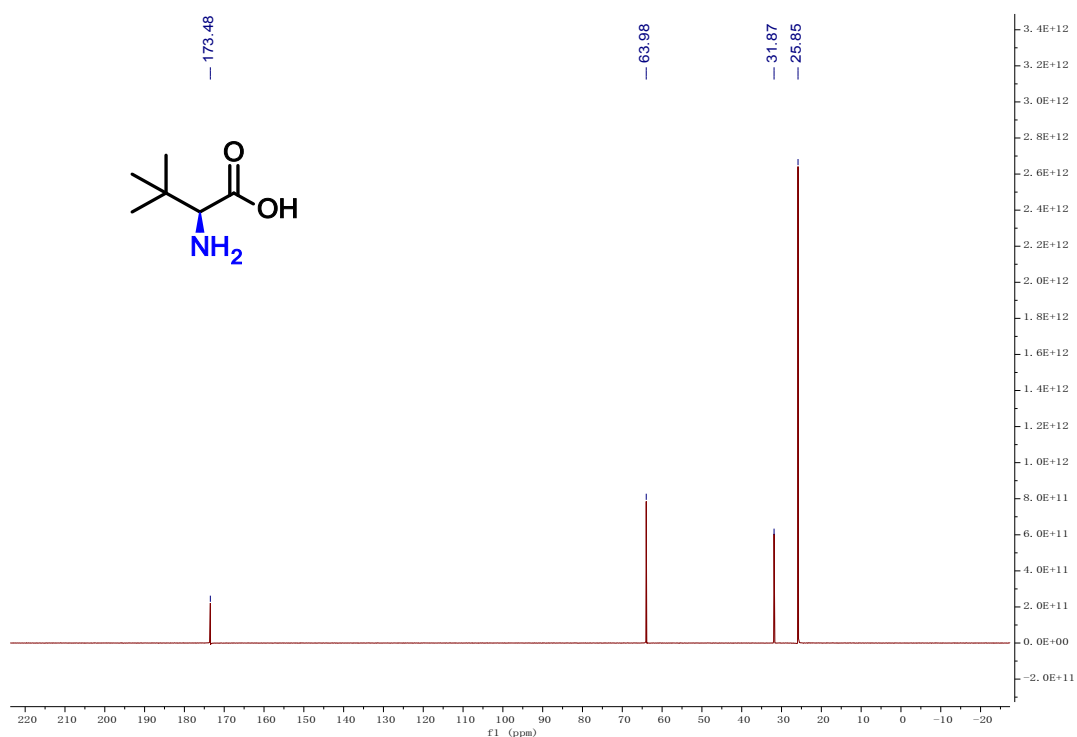
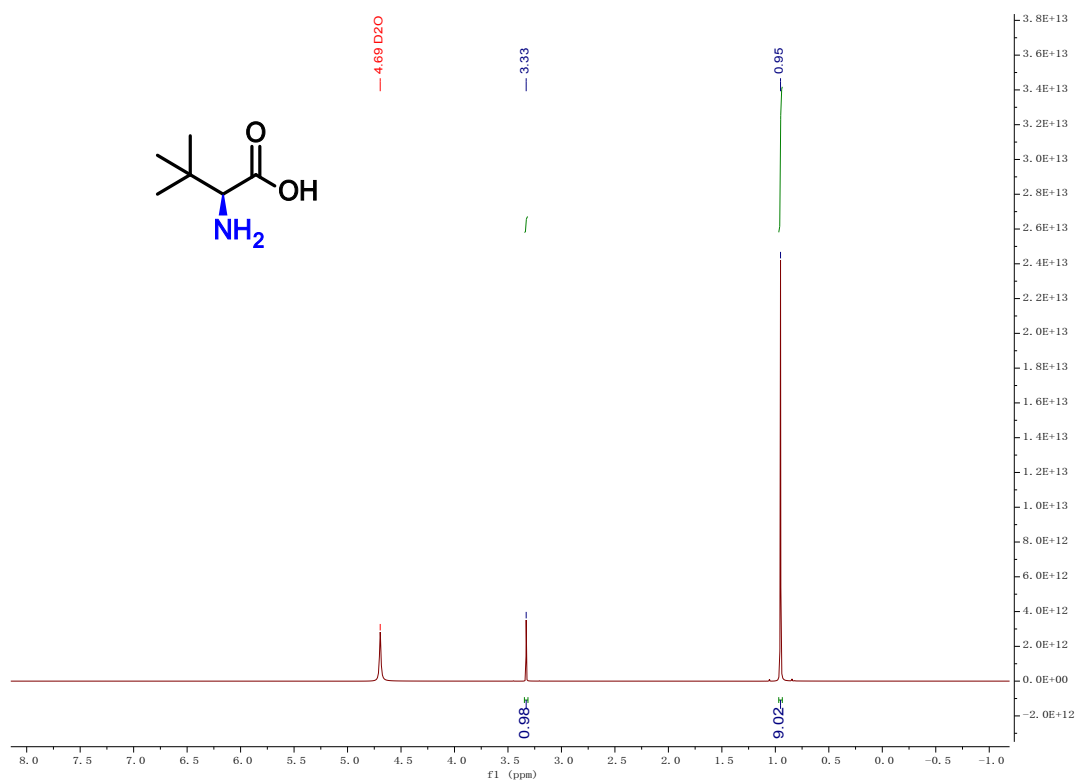
The L-*tert*-Leucine prepared by enzymatic cascade reaction.

## 5. $^1\text{H}$ NMR and $^{13}\text{C}$ NMR spectra

### 5.1 NMR of product 7,12-dioxo-LCA prepared by enzymatic cascade reaction



## 5.2 NMR of product L-*tert*-Leucine prepared by enzymatic cascade reaction



## 6. References

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