## **Supplementary Information**

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Table S1. Strains, Kits, Chemical Reagents and other Materials used in this study

, , ,		asca in this study
Material	Purchased From	Cart No.
Trelief 5α	Tsingke	TSC-C01
BL21(DE3)	Biosharp Life Science	BL1287A
V.natriegens ATCC14048	Mingzhou Biology	BMZ146848
Phanta Max Super-Fidelity DNA Polymerase	Vazyme	P505
2 × Rapid Taq Master Mix	Vazyme	P201
FastPure Gel DNA Extraction Mini Kit	Vazyme	DC301
RapidLyse Plasmid Mini Kit	Vazyme	DC211
ClonExpress II One Step Cloning Kit	Vazyme	C112
Ultra GelRed (10,000 ×)	Vazyme	GR501
100 bp DNA Ladder	Vazyme	MD104
DL5000 DNA Marker	Vazyme	MD102
SanPrep Column Plasmid Mini-Preps Kit	Sangon Biotech	B518191
50X TAE Buffer	Sangon Biotech	B548101
IPTG	Sangon Biotech	A100487
Carbenicillin disodium	Sangon Biotech	A429319
Kanamycin sulfate	Sangon Biotech	A430277
Chloramphenicol	Sangon Biotech	A429048
Magnesium chloride hexahydrate	Sangon Biotech	A601336
Potassium acetate	Sangon Biotech	A601169
Magnesium sulfate anhydrous	Sangon Biotech	A601988
Sucrose	Sangon Biotech	A502792
1 kb DNA Ladder (Dye Plus)	TaKaRa	3426A
QuickCut <sup>TM</sup> Dpn I	TaKaRa	1609
QuickCut <sup>TM</sup> Xho I	TaKaRa	1635
QuickCut <sup>TM</sup> Kpn I	TaKaRa	1618
QuickCut <sup>TM</sup> Pst I	TaKaRa	1624
Seamless Cloning Kit	Beyotime	D7010
Manganese Chloride Tetrahydrate	Aladdin	M109463

TE buffer	Aladdin	T301525
Dimethyl sulfoxide	Aladdin	D103272
PIPES Buffered Solution	Aladdin	P301883
D-Sorbitol solution	Aladdin	S104833
Naringenin	Aladdin	N164488
L-Tyrosine	Aladdin	T103976
Naringenin chalcone	Aladdin	N414406
Agar	BIOFROXX	1182
Yeast Extract	OXOID	LP0021
Peptone	OXOID	LP0042B
Sodium Chloride	Sangon Biotech	A100241
Agarose	BIOFROXX	1110
Pure Water	Wahaha	/
Gene Pulser Cuvette 0.1cm	Biorad	1652089
2216E Liquid Medium	Hopebio	HB0132-1
Brain Heart Infusion Broth (BHI)	Hopebio	HB8297-5
Ethanol, anhydrate	Hushi	10009218
Glycerol	Hushi	10010618

Table S2. Genes' Name and Sequences

Gene or Component	Sequence (5'-3')
4CL	ATGGCGCCACAAGAACAAGCAGTTTCTCAGGTGAT
4CL	GGAGAAACAGAGCAGTTTCTCAGGTGAT
	ATTTTCCGATCAAAGTTACCGGATATTTACATCCCGA
	ACCACCTATCTCCACGACTACACCTACACCAAAACA
	TCTCCGAATTCGCCACTAAGCCTTGCCTAATCAACG
	GACCAACCGGCCACGTGTACACTTACTCCGACGTC
	CACGTCATCTCCCGCCAAATCGCCGCCAATTTTCAC
	AAACTCGGCGTTAACCAAAACGACGTCGTCATGCT
	CCTCCTCCCAAACTGTCCCGAATTCGTCCTCTTT
	CCTCGCCGCCTCCTTCCGCGGCGCAACCGCCACCG
	CCGCAAACCCTTTCTTCACTCCGGCGGAGATAGCTA
	AACAAGCCAAAGCCTCCAACACCAAACTCATAATC
	ACCGAAGCTCGTTACGTCGACAAAATCAAACCACT
	TCAAAACGACGACGAGTAGTCATCGTCTGCATCG
	ACGACAACGAATCCGTGCCAATCCCTGAAGGCTGC
	CTCCGCTTCACCGAGTTGACTCAGTCGACAACCGA
	GGCATCAGAAGTCATCGACTCGGTGGAGATTTCAC
	CGGACGACGTGGTGGCACTACCTTACTCCTCTGGC
	ACGACGGGATTACCAAAAGGAGTGATGCTGACTCA
	CAAGGGACTAGTCACGAGCGTTGCTCAGCAAGTCG
	ACGGCGAGAACCCGAATCTTTATTTCCACAGCGATG
	ACGTCATACTCTGTGTTTTGCCCATGTTTCATATCTA
	CGCTTTGAACTCGATCATGTTGTGTGGTCTTAGAGT
	TGGTGCGGCGATTCTGATAATGCCGAAGTTTGAGAT
	CAATCTGCTATTGGAGCTGATCCAGAGGTGTAAAGT
	GACGGTGGCTCCGATGGTTCCGCCGATTGTGTTGGC
	CATTGCGAAGTCTTCGGAGACGGAGAAGTATGATTT
	GAGCTCGATAAGAGTGGTGAAATCTGGTGCTC
	CTCTTGGTAAAGAACTTGAAGATGCCGTTAATGCCA
	AGTTTCCTAATGCCAAACTCGGTCAGGGATACGGA
	ATGACGGAAGCAGGTCCAGTGCTCGCAATGTCGTT
	AGGTTTTGCAAAGGAACCTTTTCCGGTTAAGTCAG
	GAGCTTGTGGTACTGTTGTAAGAAATGCTGAGATGA
	AAATAGTTGATCCAGACACCGGAGATTCTCTTTCGA
	GGAATCAACCCGGTGAGATTTGTATTCGTGGTCACC
	AGATCATGAAAGGTTACCTCAACAATCCGGCAGCTA
	CAGCAGAGACCATTGATAAAGACGGTTGGCTTCATA
	CTGGAGATATTGGATTGATCGATGACGATGACGAGC
	TTTTCATCGTTGATCGATTGAAAGAACTTATCAAGT
	ATAAAGGTTTTCAGGTAGCTCCGGCTGAGCTAGAG
	GCTTTGCTCATCGGTCATCCTGACATTACTGATGTTG
	CTGTTGTCGCAATGAAAGAAGAAGCAGCTGGTGAA
	5.5.1.51.5.5.2.1.1.5.1.1.5.1.1.1.5.1.1.1.0.1.1.0.1.1.0.1.1.0.1.1.0.1.1.0.1.1.0.1.1.0.1.1.0.1.1.0.1.1.0.1.1.0.1

GTTCCTGTTGCATTTGTGGTGAAATCGAAGGATTCG GAGTTATCAGAAGATGATGTGAAGCAATTCGTGTCG AAACAGGTTGTGTTTTACAAGAGAATCAACAAAGT GTTCTTCACTGAATCCATTCCTAAAGCTCCATCAGG GAAGATATTGAGGAAAGATCTGAGGGCAAAACTAG CAAATGGATTGTGA **TAL** ATGCTACAACCATCTGCCCAAGGAAGCTACAGCAA GGCCGTGCTGTCGACATTTCGATCCCTGGATGATCT CACCTGTGATGGTTCCAAAATCATTCTCGATGGCCA ATCGCTCAATGTGAGCTCAGTGGTCGCCGCAGCATG CCATCAAGTGCCAGCGTCAATCAGCAAGGACCCGC AGCTTCATGCGCGGCTGCGAGAAAGCGTCGAGCTG CTTGCCCGCAAGCTGGCTGAAGGCGAAATCGCCTA TGGGGTCAACACCGGGTTTGGTGGCAGCGCTGACA CCCGCACCGATGACTATTCCACCCTGCAGCAGGCA CTGGTCCAGCATCTGGCGTCCGGTGTGTTGCTGCCC ACCGACCGCAGCAGTAGTCGACCATCGTCGCGGTA CCCGCATGGTTTGAGAAGCCATAGCATGCCCACAG CCGTGGTCAAGGCTGCCATGCTCGTCCGATGTAACT CACTCTTGCGAGGGCATTCCGCCGTCCGCCCGAA GTCATTGAGCACATCCTGGCTTTCCTACGCAGTGGT CTAATCCCGGTGGTGCCTGTTCGGGGGAGCATCTCG GCATCCGGCGACCTGTCCCCGCTCTCGTATATCGCC AACGCTCTCGAGGGGAACCCCGACATTGCTATCCA GAATGAGGCCACTGGGGACGTCATCCGCGCCGACG AGGCGTTGCAGCAGTTGGGCCTTGCGCCTCTCAGG TTCGGGCCCAAGGAGGCCTGGGCCTGCTCAACGG CACGGCCTTTAGCGCTGGCGCCGCCAGCTTGGTCC TCTTCGAAGCCAACCAGCTCGTGCTCCTTTCCCAGG TACTGACAGCGATGGGAACGGAAGCGCTGGCGGGC TCGACCGGGAACTACCATCCCTTCATTGCGGGCGTC CGTCCCCACCGCGGCCAAATCGAGGCGGCCGGAAA CATCTTCCACTTACTGCGGGACTCGCGCATGGCCAC CAGCCCTGGCGCCGATGGCGCAAGCCACAGTAGTA GCTTGGCGCAGGACCGGTATGCGCTTCGCACCGCA TCCCAATGGATAGGCCCACAGATCGAAGACATGTCT CTTGCTTCGGAACAGGTCCACTGCGAATTGAACTC CACCACAGACAATCCCTTGCTGGATCCGGGCTCGG GCCACATGCACCATGGAGGGAACTTCCAGGCCACT TCGATTACCAGTGCCATGGAGAAGACCATGAGCGC GATGCAGATGCTTGGCCGCATGATCTTTTCCCAGTG TACCGAGCTCATCAACCCGGCCTTGAACAATGGGC TGCCACCAAACCTCTCCTTCGACGACCCCAGTCTGT CGTTCACCATGAAAGGAATCGACATCAACATGTCG

GCATACATGGCCGAATTAAGCTACCTGAATCACCAC GTCAGCAACCACGTCCAGTCCGCCGAGATGCACAA CCAGGGCCTCAACTCCCTCGCCCTGGTCGCCAGCC GCTACGCGGCCGAGACGGTCGAGGTCTTGTCCTTG ATGGCCGCGCATATCTCTATGCACTGTGTCAGGCG CTCGATCTCCGGGCCTGCCACCTGGAGTTCCTCCGC GACGCTCGCAGCACGGTCGATAGCCTCACCGCCGA GTTATGCCTGTCATTCTCCCCCCTGTTGTCCGAGTCC GACCAGCGTCAGATCCAGGATTCCACATGGGAGCA GCTCCTGCACCACTGGAACCGAAGCAGCACCAGCG ACCTCCACGACCGCAGTCGCAATGCGGCGAGTCAC ACGATGGGTGCCCTCGTCGAGCTGCTGCCGATGCA GCTGAACGAAGCGACTGCTCTGCCCACTCGCATCG CCCCCAGCAGCAGTGGCTCGACGAGGTGTCTGCC ACGTTGGCTGGCAGCTACGATGCAACCCGAGCCAA GTTCCAATCCAATCCGACCACCCCCTTCTATCTCTG CAGTGCATCCCGCAGGATGTACGAATTCGTGCGCA AGGGGCTGGGGGTGCCGCTTCACCGCGGGATTGTT GACCACCCGACCTATCCTTCTGTAACCGAGGAGCA CGCTGGCAAGGAGCTGATCGGGTCGCAGGTGAGCA AGATCTACATGGCATTGCGTGGAGGTGCGTTCCGTG ACGTGCTACAAGATTGTTGGTATAGTTCGGATTCTG TGTGGGGTTTTGCTGGCGAGGAACTGGTTTTGGCTT **CCAAGTTGTGA** 

## CHI (codon optimized)

ATGGCGGATTTTAAATTTGAACCAATGCGTAGTCTC ATATATGTTGACTGCGTCAGCGAAGACTATCGACCT AAACTCCAACGCTGGATTTACAAAGTGCATATTCCT GATTCGATCTCTCAGTTCGAGCCTTATGTGACAAAA TACGCGTTTTATCCCTCTTTCCCAATCCCACCCCAAG GAGATAGGTTTGGTTACGCCCGTATGCAACTTACAG AACACCACTGGTTAGTCTCTGACCTTGATCCACGAT TAGAAATCAAAGCGATTGCCGAAACTTTCCCCATGG ACGTACTAGTGTGGCAAGGGCAAATACCTGCCGCA GCCCACACGGATGCTCAGATCGATAGTGATGGAGAT GCGGGCAATGCCGCGCGAAAATCCAATAATGCTGA GGGAAATCCATTTATTTTCGCGTTTCTTCCGATGTGG TGGGAAAAGGATTTGAAGGGCAAAGGTAGAACAAT AGAAGATGGCGCAAACTATCGGTTTAACATGACAAT TGGCTTTCCAGAGGGGGTGGATAAGGCAGAAGGTG AGAAATGGCTGTTCGAAAAAGTGGTACCGATCTTA CAGGCAGCTCCGGAATGTACTCGTGTGTTGGCGAG CGCGGTGAAAAAAGACATTAATGGATGCGTGATGG ATTGGGTTCTCGAAATATGGTTTGAAAACCAATCGG GCTGGTACAAGGTTATGGTTGATGATATGAAAGCAC

	TAGAAAAACCCTCATGGGCTCAGCAGGACGCATTC
	CCCTTTCTGAAGCCCTATCATAATGTGTGCTCGGCT
	GCCGTTGCAGATTATACCCCATCTAACAACTTAGCT
CIIC ( 1	AACTACCGGGGGTACATCACCATGCGT
CHS (codon	ATGGTACGGAAGCGTTACGTACATTTAACCGAAGAA
optimized)	TTCATCCAAAGAAACCCTAATATATGTGATAACACTT
	CTCCGTCTTTGGATGTGCGACAGGACTTGCTAGTAG
	TGGAAGTTCCCAAACTGGGGCAAGAAGCGGCTACT
	AAAGCTATCAAAGAGTGGGGCCAACCTAAGAGCAA
	CATAACACCTGATTTTTTGTACCAACTCTTGTGT
	GGAGATGCCGGGGTCTGATTATCGACTCGCTAACCT
	CCTGGGTCTAAACCCCTACGTAAAACGCTATATGAT
	GTATCAGCAGGGGTGCTACGCTGGCGGGATGGTTCT
	TAGACTTGCTAAAGACTTGGCCGAGAACACCAAGG
	GAGCCCGTGTCTTAGTCGTATGCTCTGAAATTACTG
	CCATAGCATTTCACGGTCCAAACGAAAACACTTCG
	GTTGATTACCTAGTCGGTCAGGCAATTTTTGGTGAT
	GGAGCAGGTGCAGTCATAGTGGGCAGCGATCCAGA
	TCTCAAGAGAACTGCCGCTGTTTGAAATGGTATC
	GGCAACGCAGACCTTCGTGCCCGACTCAGTGGGCG
	CGATCGGGGGTAGGTTGTCAGAGGCTGGGCTTATGT
	TCTACCTTGGTAAGACTGTTCCCGGTTTGATAAGCG
	AAAATATCGAGAAAGCTTTGATTCAAGCATTTAGTC
	CCTTAGGGATAACTGATTGGAACTCAATCTTTTGGA
	TTGTACACCCTGGCGGTCCTGCTATTCTTGATCAGG
	TTGAACTTAAACTCGGCTTAGAAAAGGACAAGATG
	CGAGCTAGCCGCCACGTGTTGAGCGAATATGGCAAT
	ATGTCCCATGTATCCGTCTTATTCATTATCGATGAGAT
	GAGGAAGAAATCAGTGACGGATGGAGCCGCGACTA
	CGGGTGAAGGACTGGATTGGGGTGTGCTTTTTGGC
	TTTGGTCCCGGCCTAACCATCGAGACGATTGTTCTT
	CATAGTATACCCACGACCCTGCCAGTTTCTTCCACT
	ATTGGCAGTCAGGTC
OMT3	ATGAAGAATTCATCAACGGATGAAGATTTATCCATAT
	TCGCGATGCAAGTGGCTACTTCTTCCATAGTCCCCA
	GAGCTCTGAAAGCTGTCATAGAACTAGACCTGCTG
	GAGATGATGAAGAAGGCCGGCCGTCCCCTTTCCAC
	GTCAGAAATGGCCGCCCAGATTCAGGCCACCAACC
	CGGAGGCCGCCCTCATGATTGACAGGATCCTTCGA
	GTTCTCATCGCCAGCAACATTCTAGAATGCACCACT
	GCTGCCTCTCACGGTGGCGCTGAGCGGCTTTATTCT
	TTGGCTCCGGTTTGCAAGTTCTTCACCAAGAATGAT
	GATGGTGTTTCTTGGGCTCCATTGTTTCTCATGATCC
	AAGACAGAGTCTTCACAGAAGCCTGGGATCATGTA
	THIS TOTAL TOTAL TOTAL AND THE

AAGGATGCAATAGTTGAAGGAGGAATCCCGTTCAA TATAGCCCATGGAATGAGTGGGTTCGAATACCCGGC AACCGACCCGAGATATAACAAGATTTTCAACCAAG CCATGTCTGATGAATCCACCATGTTTATGCATAAAAT TCTTGAATTATATGATGGATTTGACGGTTTGAAATCT GTTGTGGATGTTGGTGGAATTGGAGCTTCACTA AAGATGATTATAACCAAGTATCCATCTATTCAGGCCA TCAATTTCGATTTGCCCCATGTCATCCAAAATGCTCC ATCTCATCCTGGGTTGGAGCACAGAAGTGGAGACA TGTTTGTTAGTGTGCCTACAGGAGATGCCATTTTGT TGAAGTGGATTATCCATAATTGGAGCGATGGGCATT GCTTAAAACTCCTGAAAAACTGCTACGAAGCACTT CCAGAAAAGGGAAAAGTGATAATCGCAGATCGGAT TCTTCCGGAAACAGAGAATTATAAGGAGGCATCGG CGTCGGTAGACTTGGCCGGCGACGCGCTCATGTTA ACGTTGTTCACCGGTGGGAAGGAGAGGGCAGAAG CAGAATTTCAAGCTTTGGCAAAAGCATCTGGTTTCA AACATTTCCGTAAAGTTTGTTGTGCTTTCAGCACTT **GGATCATGGAACTCTACAAATAA** 

T7 RNA polymerase

ATGAACACTATCAACATTGCGAAGAACGATTTCTCT GACATCGAACTCGCGGCAATCCCGTTCAACACCCT GGCAGATCACTACGGTGAACGTCTGGCCCGTGAAC AGCTGGCTCTGGAACACGAAAGCTATGAAATGGGT GAGGCGCGTTTCCGTAAGATGTTCGAACGTCAGTTA AAAGCGGGTGAAGTTGCAGACAACGCCGCGGCGA AGCCGCTGATCACCACCCTGCTGCCGAAAATGATC GCTCGCATCAACGATTGGTTCGAAGAAGTGAAAGC GAAACGTGGCAAACGTCCGACTGCCTTCCAGTTCC TGCAAGAAATTAAACCGGAGGCAGTGGCGTATATC ACTATCAAAACTACCCTGGCGTGCCTGACCTCTGCG GATAACACGACCGTTCAGGCGGTAGCGTCCGCAAT CGGTCGTGCGATCGAAGATGAAGCCCGCTTCGGCC GTATCCGTGACCTTGAAGCAAAACACTTTAAAAAA AACGTTGAAGAACAGCTGAACAAACGCGTTGGTCA CGTATACAAAAAGGCGTTTATGCAGGTTGTGGAAG CTGATATGCTGTCCAAAGGCCTGCTGGGCGGCGAA GCGTGGAGCTCTTGGCATAAAGAAGATTCTATTCAC GTGGGCGTTCGCTGCATTGAAATGCTGATCGAATCG ACTGGTATGGTTTCTCTGCACCGTCAGAACGCTGGT GTTGTTGGCCAGGATTCTGAAACTATCGAACTGGCC CCGGAATATGCGGAAGCGATCGCTACTCGCGCAGG TGCGCTGGCCGGTATTTCCCCGATGTTCCAGCCGTG CGTTGTACCGCCTAAACCGTGGACCGGCATCACCG GCGGTGGTTATTGGGCTAACGGTCGCCGTCCGCTGG CACTCGTTCGTACCCACAGCAAAAAGGCACTGATG CGTTACGAAGATGTGTACATGCCGGAGGTTTACAAA GCCATCAACATTGCTCAGAATACCGCCTGGAAGATT AATAAAAAAGTGCTCGCGGTGGCGAACGTTATTAC CAAGTGGAAACACTGTCCGGTGGAAGATATCCCGG CTATCGAGCGCGAGGAACTGCCGATGAAACCGGAA GATATCGATATGAACCCGGAAGCTCTGACCGCGTGG AAACGTGCTGCTGCCGTATATCGCAAAGACAA GGCGCGTAAGTCTCGCCGTATCTCTCTGGAGTTTAT GCTGGAACAGGCGAACAAATTCGCCAACCACAAA GCCATCTGGTTCCCGTACAACATGGATTGGCGTGGC CGTGTCTATGCAGTTTCTATGTTCAACCCGCAAGGG AATGATATGACCAAAGGCCTGCTGACTCTGGCAAA AGGTAAGCCGATCGGCAAAGAAGGTTATTACTGGC TGAAAATCCATGGCGCGAATTGCGCGGGTGTAGATA AGGTACCGTTCCCGGAACGTATTAAGTTCATTGAAG AGAACCACGAGAATATCATGGCATGCGCGAAAAGC CCGCTGGAAAACACTTGGTGGGCTGAGCAGGACTC CCCGTTCTGCTTCCTGGCGTTCTGCTTCGAGTACGC AGGTGTTCAACATCATGGCCTGAGCTACAACTGCTC CCTGCCTCTGGCATTCGACGGTTCTTGCAGCGGTAT CCAGCACTTTTCCGCTATGTTACGTGATGAAGTAGG TGGCCGTGCTGTCAATCTGCTGCCGTCCGAAACCGT TCAAGATATCTATGGTATCGTAGCGAAAAAGGTTAA CGAGATTCTGCAAGCAGATGCTATCAACGGTACCGA CAACGAGGTAGTTACCGTGACCGATGAAAACACCG GTGAAATTTCCGAAAAAGTAAAACTGGGGACCAAG GCACTGGCCAGTGGCTGGCCTACGGCGTGAC TCGCAGCGTCACCAAACGTAGCGTGATGACGCTCG CTTATGGCAGCAAAGAGTTTGGCTTTCGCCAGCAA GTTCTGGAGGATACTATCCAGCCAGCCATTGACTCG GGCAAGGGTCTGATGTTTACCCAGCCGAACCAGGC CGCTGGATATATGGCAAAACTGATCTGGGAATCTGT AAGCGTCACCGTTGTGGCAGCGGTTGAAGCTATGA ATTGGTTGAAAAGCGCCGCGAAACTGCTGGCAGCG GAAGTGAAAGATAAAAAAACTGGCGAAATCCTGCG TAAACGCTGCGCGGTTCACTGGGTCACTCCGGATG GCTTCCCAGTTTGGCAGGAATACAAAAAACCTATCC AGACCCGCCTGAATCTGATGTTCCTGGGCCAATTCC GTCTGCAACCGACCATCAACACCAACAAAGACAGC GAAATTGACGCCCACAAACAGGAGTCCGGCATTGC CCCTAACTTCGTTCACTCTCAGGACGGCTCTCATCT GCGCAAAACTGTCGTTTGGGCTCATGAAAAGTACG GCATCGAATCCTTTGCGCTGATCCACGACTCCTTCG

	GTACCATCCCGGCGACGCGGCCAACCTGTTCAAA
	GCTGTTCGTGAAACTATGGTAGACACGTACGAAAG
	CTGCGACGTTCTGGCAGACTTCTACGATCAGTTCGC
	TGACCAGCTGCACGAAAGCCAGCTGGACAAGATGC
	CAGCTCTGCCGGCTAAAGGTAATCTGAACCTGCGC
	GACATTCTGGAATCTGACTTCGCTTTCGCCTGA
TtgR	ATGGTTAGACGAACAAAAGAGGAGGCCCAGGAGA
	CCCGCGCGCAAATTATAGAAGCTGCGGAAAAGGCA
	TTTTACAAACGTGGCGTTGCGCGAACCACCCTTGC
	GGACATCGCCGAGCTGGCCGGAGTCACCCGAGGTG
	CTATCTACTGGCACTTCTCGAACAAAGCTGAGTTAG
	TGCAAGCGCTGCTGGACTCATTACACGAGACTCAT
	GACCACCTGGCAAGGGCTTCTGAATCTGAAGATGA
	GCTCGACCCGTTCGGCTGTATGCGCAAATTACTATT
	ACAGGTCTTCAACGAACTTGTACTGGACGCCCGTA
	CACGACGTATTAACGAAATTCTCCATCATAAGTGCG
	AATTTACCGACGATATGTGCGAAATTCGTCAACAAC
	GACAGGGTGCTGTTGGACTGTCACGAAGGTGTT
	GCGTTGGCATTGGCTAATGCTGTTCGTCGGGAGCAA
	CTGCCGACTGATCTTGATATCGAACGTGCTGCCGTG
	GCGATTTTTGCTTACGTCGACGGTCTTATCGGTCGG
	TGGCTACTTCTGCCCGATTCGTTTGACTTACTGCGC
	GATGTGGAAAAATGGGTTGACACAGGTTTGGATAT
CER	GCTTCGGCTGTCCCCGGCCCTCCGAAAA
GFP	ATGCGTAAAGGAGAAGAACTTTTCACTGGAGTTGT
	CCCAATTCTTGTTGAATTAGATGGTGATGTTAATGGG
	CACAAATTTTCTGTCAGTGGAGAGGGTGAAGGTGA
	TGCAACATACGGAAAACTTACCCTTAAATTTATTTG
	CACTACTGGAAAACTACCTGTTCCATGGCCAACACT
	TGTCACTACTTTCGGTTATGGTGTTCAATGCTTTGCG
	AGATACCCAGATCATATGAAACAGCATGACTTTTTC
	AAGAGTGCCATGCCCGAAGGTTATGTACAGGAAAG
	AACTATATTTTCAAAGATGACGGGAACTACAAGAC
	ACGTGCTGAAGTCAAGTTTGAAGGTGATACCCTTG
	TTAATAGAATCGAGTTAAAAGGTATTGATTTTAAAG
	AAGATGGAAACATTCTTGGACACAAATTGGAATAC
	AACTATAACTCACACAATGTATACATCATGGCAGAC
	AAACAAAAGAATGGAATCAAAGTTAACTTCAAAAT
	TAGACACAACATTGAAGATGGAAGCGTTCAACTAG
	CAGACCATTATCAACAAAATACTCCAATTGGCGATG
	GCCCTGTCCTTTTACCAGACAACCATTACCTGTCCA
	CACAATCTGCCCTTTCGAAAGATCCCAACGAAAAG
	AGAGATCACATGGTCCTTCTTGAGTTTGTAACAGCT
	GCTGGGATTACACATGGCATGGATGAACTATACAAA

	TAA
Promoter J23102	TTGACAGCTAGCTCAGTCCTAGGTACTGTGCTAGC
Promoter J23106	TTTACGGCTAGCTCAGTCCTAGGTATAGTGCTAGC
Promoter J23107	TTTACGGCTAGCTCAGCCCTAGGTATTATGCTAGC
Promoter ttg2	CAGCAGTATTTACAAACAACCATGAATGTAAGTATA
	ATCC
Promoter ttg1	CAGCAGTATTTACAAACAACCATGAATGTAAGTATA
	TTCC
RBS B003m	AGAGTCACAGGACTACTA
RBS B0032m	AGAGTCACACAGGAAAGTACTA
RBS for V.natriegens	GGCAGCAGCCAAAGGAGATATA
1	
RBS for V.natriegens	AGAGTCACACAGGAAAGTACTA
2	
Plac and LacO	GAGGATCGAGATCGAAATTAATGTGAGTTAGCTCAC
	TCATTAGGCACCCCAGGCTTTACACTTTATGCTTCC
	GGCTCGTATGTTGTGTGGAATTGTGAGCGGATAACA
	ATTTC

**Table S3. Primers' Name and Sequences** 

D.	T	G (5: 2:)
Primer	Primer's Usage	Sequence (5'-3')
TAL-f-	insert the TAL gene into	ATCGCTGACGTCGGTACCCTCGAGG
XhoI	the pETDuet-4CL	GCAGCAGCCAAAGGAG
	plasmid	
TAL-r-	insert the TAL geneinto	GCAGCGGTTTCTTTACCAGACTCGA
XhoI	the pETDuet-4CL	GTCACAACTTGGAAGCCAAAACCA
	plasmid	GTTCC
pETDuet-	Linearize the vector	TCGATCTCGATCCTCTACGCC
f-Plac	pETDuet-4CL-TAL	
pETDuet-	Linearize the vector	TATTGTACACGGCCGCATAATC
r-	pETDuet-4CL-TAL	
MCS1del		
pETDuet-	Linearize the vector	CAGCACATGGACTCGTCTACTA
r-linear	pETDuet-4CL-TAL	
T7pol-r-	Linearize the vector	TTTCGATCTCGATCCTCTAGAGACCC
Plac-new	pMB1-T7pol	TGATATTGACGGC
T7pol-f-	Linearize the vector	GTGAGCGGATAACAATTTCATGAAC
Plac	pMB1-T7pol	ACTATCAACATTGCGAAGAAC
cap-plac-	Anneal to form short	GAGGATCGAGATCGAAATTAATGTG
up	pieces	AGTTAGCTCACTCATTAGGCACCCC
1		AGGCTTTACACTTTATGCTTCCGGCT
		CGTATGTTGTGGGAATTGTGAGCG
		GATAACAATTTC
cap-plac-	Anneal to form short	GAAATTGTTATCCGCTCACAATTCCA
bottom	pieces	CACAACATACGAGCCGGAAGCATAA
	_	AGTGTAAAGCCTGGGGTGCCTAATG
		AGTGAGCTAACTCACATTAATTTCG
		ATCTCGATCCTC
pMB1-	Linearize the vector	AGAGGATCGAGATCGAAATTAATGT
Plac-	pMB1-plac-T7pol	GAGTTAG
T7pol-f		
pMB1-	Linearize the vector	TGCGGCCGTGTACAATACAGGAAAC
Plac-	pMB1-plac-T7pol	AGCTATGACCATGC
T7pol-r		
pETDuet-	Linearize the vector	TATTGTACACGGCCGCATAATC
r-	pETDuet-4CL-TAL	
MCS1del	1	
4CL-TAL-	Linearize the vector	GTAGACGAGTCCATGTGCTG
r	pETDuet-4CL-TAL	
pACYCD	Linearize the vector	CGGCCGATATCCAATTGAGATCTG
uet-I-S-	pACYCDuet-OMT3	
MCS1-r	F 52 55 51115	
pACYCD	Linearize the vector	GTCGGTACCCTCGAGTCTGGTAAAG
PACT CD	Linearize the vector	GICGGIACCCICGAGICIGGIAAAG

uet-I-S-	nACVCDuet OMT2	
MCS1-f	pACYCDuet-OMT3	
	ingest the CUI canainte	CTCAATTGGATATCGGCCGATGGCG
singlet- CHI-f	insert the CHI gene into	GATTTAAATTTGAACCAATGCG
CHI-I	the pACYCDuet-OMT3 plasmid	GALLITAAALITGAACCAAIGCG
ain alat	<b>-</b>	TACCAGACTCGAGGGTACCGACTTA
singlet- CHI-r	insert the CHI gene into	
СП-г	the pACYCDuet-OMT3	ACGCATGGTGATGTACCCC
-11-4	plasmid	
singlet-	insert the CHS gene into	CAGCGGTTTCTTTACCAGACTCGAG
CHS-r	the pACYCDuet-OMT3-	TTAGACCTGACTGCCAATAGTG
. 1 .	CHI plasmid	
singlet-	insert the CHS gene into	CATGCGTTAAGTCGGTACCCTCGAG
CHS-f	the pACYCDuet-OMT3-	AGAGTCACACAGGAAAGTAC
O) (T)	CHI plasmid	LATTICCA COTTOCA CA
OMT3-f-	insert the OMT3 gene	AATTCGAGCTCGGCGCCCTGCAGA
PstI	into the pACYCDuet-1	TGAAGAATTCATCAACGGATGAAG
0) (77)	plasmid	
OMT3-r-	insert the OMT3 gene	GCCGCAAGCTTGTCGACCTGCAGTT
PstI	into the pACYCDuet-1	ATTTGTAGAGTTCCATGATCCAAGTG
	plasmid	
pACYC17	Linearize the vector	GACAGTTTTATTGTTCATG
7-r-MCS2	pACYC177	
pACYC17	Linearize the vector	CATAAACAGTAATACAAGGG
7-f-MCS2	pACYC177	
Pttg2-	Anneal to form short	TGAACAATAAAACTGTCCAGCAGTA
J23106-	pieces	TTTACAAACAACCATGAATGTAAGT
linker-up		ATAATCCTGTGAAATTCCACAAGAG
		TCACACAGGAAAGTACTACATAAAC
		AGTAATACAAGGGG
Pttg2-	Anneal to form short	CCCCTTGTATTACTGTTATGTAGTA
J23106-	pieces	CTTTCCTGTGTGACTCTTGTGGAATT
linker-		TCACAGGATTATACTTACATTCATGG
bottom		TTGTTTGTAAATACTGCTGGACAGTT
		TTATTGTTCA
pACYC17	Linearize the vector	TTCAACAGATCGGGAAGGG
7-Ttgr-	pACYC177-Pttg2	
MCS3-f		
pACYC17	Linearize the vector	CTTGTCGGTAAAGATGCGG
7-Ttgr-	pACYC177-Pttg2	
MCS3-r		
J23106-	insert the J23106-TtgR	CTTCCCGATCTGTTGAATTATTTTCG
rbs-ttgr-	gene into the	GAGGGCCGG
linker-r	pACYC177-Pttg2-GFP	
	plasmid	

J23106-	insert the J23106-TtgR	GCATCTTTACCGACAAGTTTACGGC
rbs-ttgr-	gene into the	TAGCTCAGTCCT
linker-f	pACYC177-Pttg2-GFP	INGCICAGICCI
IIIIKCI-I	plasmid	
cTtgr pro-	Linearize the vector	GATATACCATGGGCAGC
f	pACYC177-Pttg2-GFP-	G/II/II/ICC/II/GGGC/IGC
1	J23106-TtgR	
cTtgr pro-	Linearize the vector	CTTGTCGGTAAAGATGC
r	pACYC177-Pttg2-GFP-	CITGICGG III MIGNIGO
1	J23106-TtgR	
J23102-	Anneal to form short	GGGTAAAAGCTAACCGCATCTTTAC
rbs-linker-	pieces	CGACAAGTTGACAGCTAGCTCAGTC
up	pieces	CTAGGTACTGTGCTAGCTGTGAAAT
		TCCACAAGAGTCACACAGGACTACT
		ATAAGGAGATATACCATGGGC
J23102-	Anneal to form short	GCCCATGGTATATCTCCTTATAGTAG
rbs-linker-	pieces	TCCTGTGTGACTCTTGTGGAATTTC
bottom	F	ACAGCTAGCACAGTACCTAGGACTG
		AGCTAGCTGTCAACTTGTCGGTAAA
		GATGCGGTTAGCTTTTACCC
J23107-	Anneal to form short	GGGTAAAAGCTAACCGCATCTTTAC
rbs-linker-	pieces	CGACAAGTTTACGGCTAGCTCAGCC
up	1	CTAGGTATTATGCTAGCTGTGAAATT
1		CCACAAGAGTCACACAGGACTACTA
		TAAGGAGATATACCATGGGC
J23107-	Anneal to form short	GCCCATGGTATATCTCCTTATAGTAG
rbs-linker-	pieces	TCCTGTGTGACTCTTGTGGAATTTC
bottom	-	ACAGCTAGCATAATACCTAGGGCTG
		AGCTAGCCGTAAACTTGTCGGTAAA
		GATGCGGTTAGCTTTTACCC
cPttg-	Linearize the vector	GTGTATGCGTAAAGGAGAAG
linear-f	pACYC177-Pttg2-GFP-	
	J23106-TtgR	
cPttg-	Linearize the vector	CTGCTGGACAGTTTTATTG
linear-r	pACYC177-Pttg2-GFP-	
	J23106-TtgR	
petduet-t7-	Colony PCR for	CGACGATGGTCGACTACTG
4cl-tal-	pETDuet-TAL-4CL	
colper-r		
petduet-t7-	Colony PCR for	GAACCAGGCCGCTGGATATATG
4cl-tal-	pETDuet-TAL-4CL	
colper-f		
L66A-	single point mutation	GCTGGACTCAGCACACGAGACTCAT
TtgR-f		GACCACCTGGCAAGG

L66A-	single point mutation	GTCTCGTGTGCTGAGTCCAGCAGCG
	single point mutation	CTTGCACTAACTCAG
TtgR-r	1	
G176R-	single point mutation	CGGTCTTATCCGCCGGTGGCTACTTC
TtgR-f		TGCCCGATTCGTTTG
G176R-	single point mutation	CCACCGGCGGATAAGACCGTCGACG
TtgR-r		TAAGCAAAAATCGCCAC
N110A-	single point mutation	CGACGTATTGCCGAAATTCTCCATCA
TtgR-f		TAAGTGCGAATTTACCGACG
N110A-	single point mutation	CGACGTATTGCCGAAATTCTCCATCA
TtgR-r		TAAGTGCGAATTTACCGACG
177colper-	Colony PCR for	CAAAGCCACGTTGTGTCTC
f	pACYC177-J23106-	
	TtgR-Pttg2-GFP	
177colpcr-	Colony PCR for	CCGTCGTCAATAAACCGAACC
r	pACYC177-J23106-	
	TtgR-Pttg2-GFP	
chi-chs-	Colony PCR for	CTGCGCTAGTAGACGAGTCCATGTG
colper-f	pACYCDuet-OMT-CHI-	
	CHS	
chi-chs-	Colony PCR for	GATATACCATGGGCAGCAGCCATC
colper-r	pACYCDuet-OMT-CHI-	
	CHS	
CHS-	single point mutation	GGCAATTGTAGGTGATGGAGCAGGT
F153V-f		GCAGTCATAG
CHS-	single point mutation	CCATCACCTACAATTGCCTGACCGA
F153V-r		CTAGGTAATCAACCG
CHS-	single point mutation	GATTATCGACTCATGAACCTCCTGG
A82M-f	<i>C</i> 1	GTCTAAACCCCTACG
CHS-	single point mutation	GAGGTTCATGAGTCGATAATCAGAC
A82M-r		CCCGGCATCTCCAC
mutantCH	single point mutation	CAGGTTAATGAATTTGCTCGGTCTG
S-A82M-f	0 r management	AACCCTTACGTG
mutantCH	single point mutation	GCAAATTCATTAACCTGTAATCAGA
S-A82M-r	single point materion	ACCGGCATTTCC
5 7102111		11000001111100

Table S4. Plasmids' Components and Function

Plasmid	Components	Function
pETDuet-T7RNAP-4CL-	Plac-T7RNAP, PT7-4CL-	To express T7 RNAP and
TAL	TAL	4CL, TAL for
		biosynthesis of
		flavonoids.
pACYCDuet-OMT3-	PT7-OMT3, PT7-CHI-	To express OMT3, CHI
CHI-CHS	CHS	and CHS for biosynthesis
		for flavonoids.
pACYC177-Pttg2-GFP-	PJ23106-TtgR, Pttg2-GFP	To building biosensors for
J23106-TtgR		product detection
pACYC177-Pttg2-GFP-	PJ23102-TtgR, Pttg2-GFP	To building biosensors for
J23102-TtgR		product detection
pACYC177-Pttg2-GFP-	PJ23107-TtgR, Pttg2-GFP	To building biosensors for
J23107-TtgR		product detection
pACYC177-Pttg1-GFP-	PJ23106-TtgR, Pttg1-GFP	To building biosensors for
J23106-TtgR		product detection
pACYC177-Pttg1-GFP-	PJ23107-TtgR, Pttg1-GFP	To building biosensors for
J23107-TtgR		product detection
pETDuet-Plac-4CL-	Plac-4CL, Plac-	To express 4CL, CHS and
CHSmutants	CHSmutants-CHI	CHI for detect the effect
		of directed evolution

## S5 Protocols for Molecular Cloning and Bacteria Culture

**PCR:** We use Vazyme's Phanta Max kit for amplifying the genes or fragments. The protocol was followed and the PCR product was purified using Vazyme's FastPure Kit.

**Restriction Enzyme Digestion:** For digesting plasmids or fragments, we add 4μg plasmids or fragments, 3μL FastDigest Buffer in a 30μL system. The reaction liquid was then incubated under 37°C for 30min then heated under 85°C for 20min.

**Point mutation:** To mutate certain site in a gene in cloning vector (pUC19 or pUC57), we used primers with point mutation to linearize the plasmid and introduced the mutation to it. The linearized plasmid contains 15-20bp of homologous sequences (which contains the mutated site). It is recircled using Vazyme's CE II kit.

**Homologous Recombination:** We used Vazyme's CE II kit for single fragment insertion and Beyotime's Seamless Clone kit for multi-fragments insertion. The reaction liquid was mixed by the instruction then incubated under 37°C for 30min.

Plasmid transformation for *E. Coli*: Unfreeze Trelief 5α Competent Cell on ice for about 15min. Then add no more than 10μL recombination product or 1μg plasmid to 100μL cell. Incubated on ice for 30min, then heat shock under 42°C for 60s. Then after recover on ice for 3min, add 1mL LB to the competent cell, incubated under 37°C, 200rpm. Then spread the cell on LB agar plate with appropriate antibiotic.

**Plasmid extraction and sequencing:** Pick up a single colony of E.Coli Trelief  $5\alpha$  containing the plasmid from the agar plate and add into LB liquid medium with appropriate antibiotic. Then culture under the condition of  $37^{\circ}$ C, 250rpm for 12-16h.

Then use Sangon's SanPrep kit (only pACYC plasmids) or Vazyme's RapidLyse kit for plasmid extraction. The plasmid was then sent to Sangon Biotech for sequencing (Sanger Sequencing).

Colony PCR of bacteria: Pick up a single colony on the plate and add into the PCR reaction liquid (RapidTaq) as template. PCR Procedure: 95°C 10min – (95°C 30s – 55°C 15s – 72°C 60s) 30 cycles – 72°C 5min.

Culturing *V. natriegens* ATCC14048: *V. natriegens* ATCC14048 grows well in 2216E and LB medium. V2 salts can be added to LB to make LBv2 medium. To make competent cell and perform transformation, only use BHIv2 medium.

Preparation of antibiotic solution: For carbenicillin storage solution, dissolve 500mg carbenicillin disodium powder in water and volume to 10mL. For kanamycin storage solution, dissolve 300mg kanamycin sulfate powder in water and volume to 10mL. For chloramphenicol storage solution, dissolve 250mg chloramphenicol powder in anhydrate ethanol and volume to 10mL. All the storage solution was filtered with 0.22µm filter membrane and stored under -20°C. Add the storage solution to the medium under the ratio of 1:1000 when antibiotic is used.

Preparation of medium: LB medium: dissolve 5g yeast extract, 10g peptone, 10g sodium chloride in water and volume to 1L. To make solid medium, add another 20g agar. Then the medium is sterilized under 121°C for 20min. To make 2216E or BHI medium, follow the instructions. To make BHIv2 or BHIv2 sucrose medium, sterilize BHI medium, v2 salt solution and sucrose solution separately and mix together after cooling down to room temperature.

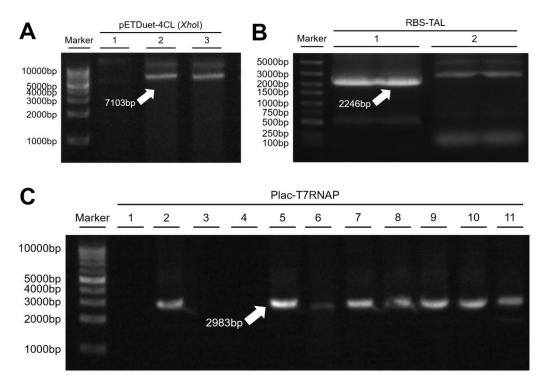


Figure S1. Electropherogram of fragments used when constructing pETDuet-T7RNAP-4CL-TAL. A. Digestion product of plasmid pETDuet-4CL; B. Amplified RBS-TAL fragments; C. Amplified Plac-T7RNAP fragments.

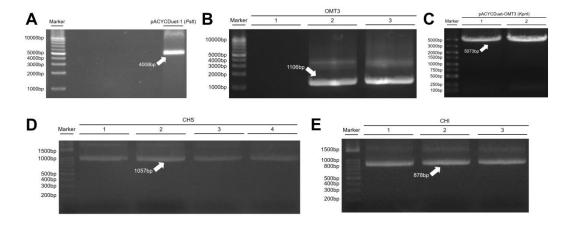


Figure S2. Electropherogram of fragments used when constructing pACYCDuet-OMT3-CHI-CHS. A. Digestion product of plasmid pACYCDuet-1; B. Amplified OMT3 fragments; C. Digestion product of plasmid pACYCDuet-OMT3; D, E. Amplified CHS or CHI fragments.

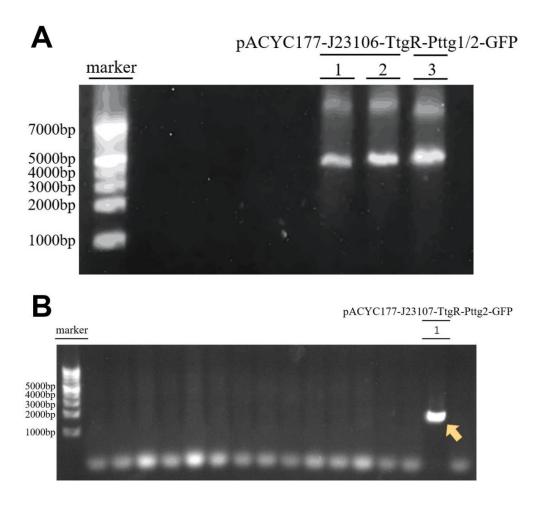


Figure S3. Electropherogram of colony PCR results when constructing pACYC177 biosensor plasmids. A. Colony PCR result of *E. coli* clones containing pACYC177-J23106-TtgR-Pttg1-GFP or pACYC177-J23106-TtgR-Pttg2-GFP; B. Colony PCR result of *E. coli* clone containing pACYC177-J23107-TtgR-Pttg2-GFP.