

Electronic Supplementary Material (ESI) for RSC Advances.
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Supplementary data file for *RSC advance*

A Multifunctional Oxidosqualene Cyclase from *Tripterygium regelii* producing both α - and β -amyrin

Yun Lu^a, Jiawei Zhou^a, Tianyuan Hu^a, Yifeng Zhang^{ac}, Ping Su^{ac}, Jiadian Wang^{ac}, Wei Gao^{*ab}, and Luqi Huang^c

a. School of Traditional Chinese Medicine, Capital Medical University, Beijing 100069, China.

b. Beijing Key Lab of TCM Collateral Disease Theory Research, School of Traditional Chinese Medicine, Capital Medical University, Beijing, China.

c. State Key Laboratory of Dao-di Herbs, National Resource Center for Chinese Materia Medica, China Academy of Chinese Medical Sciences, Beijing, China.

*Corresponding author: Wei Gao; e-mail: weigao@ccmu.edu.cn; Tel.: +86 10 83916572; Fax: +86 10 83911627.

Fig S1.

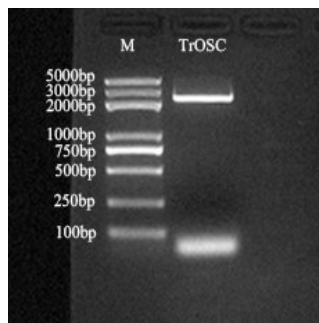


Fig S1. Agarose gel electrophoresis of the cDNA of *TrOSC*, which coded for a 2289 bp open reading frame. M: Trans 2K plus DNA Marker.

Fig S2.



Fig S2. The 3D structure prediction of *TrOSC* protein using homology-based modelling. 3D modelling of *TrOSC* was used protein 1w6j.1 (Lanosterol synthase) as a template (Modeling range: amino acids 54-748) and showed 41.54% sequence identity.

Table S1. List of primers used in this study.

Primer Name	Sequence (5' → 3')
OSC-F	AAATATCAAATATGTGGAGGCTTA
OSC-R	TGGACTGGGCATTGGTCTA
TrOSC-pYES2-F	GACTCACTATAGGGAATATTAATGTGGAGGCTTAAGGTGGG
TrOSC-pYES2-R	ACTAGATGCATGCTCGAGCTTACACGTCTGGAGCACTTTTCG
qTrOSC-F	TCTTATTGTCCCTACAGAGTGAAAACG
qTrOSC-R	ATGTCCAGGATATAGCCTTTTGAATAG
EF- α F	CCAAGGGTGAAAGCAAGGAGAGC
EF- α R	CACTGGTGGTTTTGAGGCTGGTATCT

Table S2. Other plant OSCs included in phylogenetic tree.

Gene name	Species	Type of Synthase	Genebank accession number
<i>PqBAS</i>	<i>Panax quinquefolius</i>	beta-amyrin synthase	JX262290
<i>PjBAS</i>	<i>Panax japonicus</i>	beta-amyrin synthase	KP658156
<i>BcBAS</i>	<i>Bupleurum chinense</i>	beta-amyrin synthase	EU400220
<i>SlBAS</i>	<i>Solanum lycopersicum</i>	beta-amyrin synthase	NM_001247675
<i>PgBAS</i>	<i>Platycodon grandiflorus</i>	beta-amyrin synthase	KY412556
<i>AeBAS</i>	<i>Aralia elata</i>	beta-amyrin synthase	HM219225
<i>PgBAS</i>	<i>Panax ginseng</i>	beta-amyrin synthase	O82140
<i>KsBAS</i>	<i>Kalopanax septemlobus</i>	beta-amyrin synthase	MF627963
<i>EsBAS</i>	<i>Eleutherococcus senticosus</i>	beta-amyrin synthase	KX378998
<i>RsMS</i>	<i>Rhizophora stylosa</i>	multifunctional triterpene synthase	AB263203
<i>BpBAS</i>	<i>Betula platyphylla</i>	beta-amyrin synthase	Q8W3Z1
<i>BpBAS</i>	<i>Betula platyphylla</i>	beta-amyrin synthase	AB055512
<i>LjMS</i>	<i>Lotus japonicus</i>	multifunctional triterpene synthase	AF478455
<i>PsMAS</i>	<i>Pisum sativum</i>	mixed-amyrin synthase	AB034803
<i>PsMAS</i>	<i>Pisum sativum</i>	mixed-amyrin synthase	Q9LRH7
<i>RcLUS</i>	<i>Ricinus communis</i>	lupeol synthase	NM_001323755
<i>KcMS</i>	<i>Kandelia candel</i>	multifunctional triterpene synthase	AB257507
<i>MdMAS</i>	<i>Malus domestica</i>	mixed-amyrin synthase	FJ032006
<i>EjMAS</i>	<i>Eriobotrya japonica</i>	mixed-amyrin synthase	JX173279
<i>RsMS</i>	<i>Rhizophora stylosa</i>	multifunctional triterpene synthase	BAF80442
<i>OeMAS</i>	<i>Olea europaea</i>	mixed-amyrin synthase	AB291240
<i>CrMAS</i>	<i>Catharanthus roseus</i>	mixed-amyrin synthase	JN991165
<i>LjLUS</i>	<i>Lotus japonicus</i>	lupeol synthase	AB181245
<i>VrLUS</i>	<i>Vigna radiata</i> var. <i>radiata</i>	lupeol synthase	XM_014649572
<i>AdLUS</i>	<i>Arachis duranensis</i>	lupeol synthase	XM_016110878
<i>BpLUS</i>	<i>Betula platyphylla</i>	lupeol synthase	AB055511
<i>WsLUS</i>	<i>Withania somnifera</i>	lupeol synthase	JQ728552
<i>OeLUS</i>	<i>Olea europaea</i>	lupeol synthase	AB025343
<i>ToLUS</i>	<i>araxacum officinale</i>	lupeol synthase	AB025345
<i>AmCAS</i>	<i>Abies magnifica</i>	cycloartenol synthase	AF216755
<i>DzCAS</i>	<i>Dioscorea zingiberensis</i>	cycloartenol synthase	AM697885
<i>DzCAS</i>	<i>Dioscorea zingiberensis</i>	cycloartenol synthase	FN547822

<i>AmCAS</i>	<i>Allium macrostemon</i>	cycloartenol synthase	AB025353
<i>LaCAS</i>	<i>Luffa aegyptiaca</i>	cycloartenol synthase	AB033334
<i>CpCAS</i>	<i>Cucurbita pepo</i>	cycloartenol synthase	AB116237
<i>SlCAS</i>	<i>Solanum lycopersicum</i>	cycloartenol synthase	NM_001246855
<i>AtCAS</i>	<i>Arabidopsis thaliana</i>	cycloartenol synthase	U02555
<i>BpCAS</i>	<i>Betula platyphylla</i>	cycloartenol synthase	AB055510
<i>LjCAS</i>	<i>Lotus japonicus</i>	cycloartenol synthase	AB181246
<i>PsCAS</i>	<i>Pisum sativum</i>	cycloartenol synthase	D89619
<i>KcCAS</i>	<i>Kandelia candel</i>	cycloartenol synthase	AB292609
<i>BkCAS</i>	<i>Bupleurum kaoi</i>	cycloartenol synthase	AY514456
<i>PgCAS</i>	<i>Panax ginseng</i>	cycloartenol synthase	AB009029
<i>CaCAS</i>	<i>Centella asiatica</i>	cycloartenol synthase	AY520819
