

Product datasheet for TR500470

Cyp1a1 Mouse shRNA Plasmid (Locus ID 13076)

Product data:

OriGene Technologies, Inc.

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Product Type:	shRNA Plasmids
Product Name:	Cyp1a1 Mouse shRNA Plasmid (Locus ID 13076)
Locus ID:	13076
Synonyms:	AHH; AHRR; CP11; Cyp1a2; CYPIA1; P450-1
Vector:	pRS (TR20003)
E. coli Selection:	Ampicillin
Mammalian Cell Selection:	Puromycin
Format:	Retroviral plasmids
Components:	Cyp1a1 - Mouse, 4 unique 29mer shRNA constructs in retroviral untagged vector(Gene ID = 13076). 5µg purified plasmid DNA per construct 29-mer scrambled shRNA cassette in pRS Vector, TR30012, included for free.
RefSeq:	<u>BC125440, BC125444, NM 001136059, NM 009992, NM 001136059.1, NM 001136059.2, NM 009992.1, NM 009992.2, NM 009992.3, NM 009992.4</u>
UniProt ID:	<u>P00184</u>



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GRIGENE Cyp1a1 Mouse shRNA Plasmid (Locus ID 13076) – TR500470

Summary:	A cytochrome P450 monooxygenase involved in the metabolism of various endogenous
	substrates, including fatty acids, steroid hormones and vitamins. Mechanistically, uses
	molecular oxygen inserting one oxygen atom into a substrate, and reducing the second into a
	water molecule, with two electrons provided by NADPH via cytochrome P450 reductase (CPR;
	NADPH-ferrihemoprotein reductase). Catalyzes the hydroxylation of carbon-hydrogen bonds.
	Exhibits high catalytic activity for the formation of hydroxyestrogens from estrone (E1) and
	17beta-estradiol (E2), namely 2-hydroxy E1 and E2, as well as D-ring hydroxylated E1 and E2
	at the C15alpha and C16alpha positions. Displays different regioselectivities for
	polyunsaturated fatty acids (PUFA) hydroxylation. Catalyzes the epoxidation of double bonds of certain PUFA. Converts arachidonic acid toward epoxyeicosatrienoic acid (EET) regioisomers, 8,9-, 11,12-, and 14,15-EET, that function as lipid mediators in the vascular system. Displays an absolute stereoselectivity in the epoxidation of eicosapentaenoic acid (EPA) producing the 17(R),18(S) enantiomer. May play an important role in all-trans retinoic acid biosynthesis in extrahepatic tissues. Catalyzes two successive oxidative transformation of all-trans retinol to all-trans retinal and then to the active form all-trans retinoic acid. May also participate in eicosanoids metabolism by converting hydroperoxide species into oxo metabolites (lipoxygenase-like reaction, NADPH-independent).[UniProtKB/Swiss-Prot
	Function]
shRNA Design:	These shRNA constructs were designed against multiple splice variants at this gene locus. To be certain that your variant of interest is targeted, please contact <u>techsupport@origene.com</u> . If you need a special design or shRNA sequence, please utilize our <u>custom shRNA service</u> .
Performance Guaranteed:	OriGene guarantees that the sequences in the shRNA expression cassettes are verified to correspond to the target gene with 100% identity. One of the four constructs at minimum are guaranteed to produce 70% or more gene expression knock-down provided a minimum transfection efficiency of 80% is achieved. Western Blot data is recommended over qPCR to evaluate the silencing effect of the shRNA constructs 72 hrs post transfection. To properly assess knockdown, the gene expression level from the included scramble control vector must

For non-conforming shRNA, requests for replacement product must be made within ninety (90) days from the date of delivery of the shRNA kit. To arrange for a free replacement with newly designed constructs, please contact Technical Services at techsupport@origene.com. Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled shRNA control (Western Blot data preferred).

be used in comparison with the target-specific shRNA transfected samples.

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