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CYP2C19 Pre-design Chimera RNAi

Catalog # : H00001557-R01 規格 : [10 nmol] [20 nmol]

List All

Specification

Product Description: Homo sapiens cytochrome P450, family 2, subfamily C, polypeptide 19 (CYP2C19), mRNA.

Reactivity: Human

Supplied Product: DEPC water

Target Refseq: NM_000769

Storage Instruction: Store at -20°C, do not exceed 4 - 5 freeze-thaw cycles to ensure product integrity.

Note: Position of the Chimera RNAi.



Publication Reference

1. dsCheck: highly sensitive off-target search software for double-stranded RNA-mediated RNA interference.
Naito Y, Yamada T, Matsumiya T, Ui-Tei K, Saigo K, Morishita S. Nucleic Acids Res. 2005 Jul 1;33(Web Server issue):W589-91.
2. Functional dissection of siRNA sequence by systematic DNA substitution: modified siRNA with a DNA seed arm is a powerful tool for mammalian gene silencing with significantly reduced off-target effect.
Ui-Tei K, Naito Y, Zenno S, Nishi K, Yamato K, Takahashi F, Juni A, Saigo K. Nucleic Acids Res. 2008 Apr;36(7):2136-51. Epub 2008 Feb 11.
3. Guidelines for the selection of highly effective siRNA sequences for mammalian and chick RNA interference.
Ui-Tei K, Naito Y, Takahashi F, Haraguchi T, Ohki-Hamazaki H, Juni A, Ueda R, Saigo K. Nucleic Acids Res. 2004 Feb 9;32(3):936-48. Print 2004.
4. siDirect: highly effective, target-specific siRNA design software for mammalian RNA interference.
Naito Y, Yamada T, Ui-Tei K, Morishita S, Saigo K. Nucleic Acids Res. 2004 Jul 1;32(Web Server issue):W124-9.

Applications

RNAi Knockdown

Gene Information

Entrez GeneID: [1557](#)

Gene Name: CYP2C19

Gene Alias: CPCJ,CYP2C,P450C2C,P450IIC19

Application Image

RNAi Knockdown

Gene cytochrome P450, family 2, subfamily C, polypeptide 19

Description:

Omim ID: [124020](#)

Gene Ontology: [Hyperlink](#)

Gene Summary: This gene encodes a member of the cytochrome P450 superfamily of enzymes. The cytochrome P450 proteins are monooxygenases which catalyze many reactions involved in drug metabolism and synthesis of cholesterol, steroids and other lipids. This protein localizes to the endoplasmic reticulum and is known to metabolize many xenobiotics, including the anticonvulsive drug mephenytoin, omeprazole, diazepam and some barbiturates. Polymorphism within this gene is associated with variable ability to metabolize mephenytoin, known as the poor metabolizer and extensive metabolizer phenotypes. The gene is located within a cluster of cytochrome P450 genes on chromosome 10q24.
[provided by RefSeq]

Other Designations: OTTHUMP00000020132,OTTHUMP00000059588,S-mephenytoin 4-hydroxylase,cytochrome P-450 II C,cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 19,flavoprotein-linked monooxygenase,mephenytoin 4'-hydroxylase,microsomal monooxygenase,xenobi

Gene Pathway

[Arachidonic acid metabolism](#) [Drug metabolism - cytochrome P450](#)

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[Metabolism of xenobiotics by cytochrome P450](#) [Monoterpene biosynthesis](#)

[Retinol metabolism](#)

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