

Product datasheet for SC202319

Pirin (PIR) (NM_003662) Human 3' UTR Clone

Product data:

OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	Pirin (PIR) (NM_003662) Human 3' UTR Clone
Symbol:	Pirin
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_003662
Insert Size:	235 bp
Insert Sequence:	<pre>>SC202319 3'UTR clone of NM_003662 The sequence shown below is from the reference sequence of NM_003662. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC AAAACCTGGAAATCAAAGATTGGGAACTAGTGGAAAGCGGAAGAGCAGGTCTTGATGTGTCCTAGAATT TTGCCATTTCTGAGATTGAGCCATTGAAGGCATTCCATTTCTAAAGCTTATTTAGCCGGTGCTTCTAAA GAATTCCACACTAACGTGATAACATGGTTTTTGTAACAATAAATGTAGGATATTTCCTGGCACATGCAA ATAAACCTAATCATTGTTTCTTTAAAAA ACGCGTAAGCGGCGCGGC</pre>
Restriction Sites:	CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG Sgfl-Mlul
OTI Disclaimer:	Our molecular clone sequence data has been matched to the sequence identifier above as a point of reference. Note that the complete sequence of this clone is largely the same as the reference sequence but may contain minor differences , e.g., single nucleotide polymorphisms (SNPs).
Components:	The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The package also includes 100 pmols of both the corresponding 5' and 3' vector primers in separate vials.
RefSeq:	<u>NM 003662.4</u>



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Pirin (PIR) (NM_003662) Human 3' UTR Clone – SC202319
This gene encodes a member of the cupin superfamily. The encoded protein is an Fe(II)- containing nuclear protein expressed in all tissues of the body and concentrated within dot- like subnuclear structures. Interactions with nuclear factor I/CCAAT box transcription factor as well as B cell lymphoma 3-encoded oncoprotein suggest the encoded protein may act as a transcriptional cofactor and be involved in the regulation of DNA transcription and replication. Alternatively spliced transcript variants have been described. [provided by RefSeq, Jul 2008]
8544
9.2

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