

## Product datasheet for SC206538

## PON2 (NM\_001018161) Human 3' UTR Clone

## **Product data:**

## OriGene Technologies, Inc.

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Product Type:	3' UTR Clones
Product Name:	PON2 (NM_001018161) Human 3' UTR Clone
Vector:	pMirTarget (PS100062)
Symbol:	PON2
ACCN:	NM_001018161
Insert Size:	498 bp
Insert Sequence:	<pre>&gt;SC206538 3'UTR clone of NM_001018161 The sequence shown below is from the reference sequence of NM_001018161. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC</pre>
	TACCACAGAGCCTTGTATTGTGAACTCTAAATTGTACTTTTGGCATGAAAGTGCGATAACTTAACAATT AATTTTCTATGAATTGCTAATTCTGAGGGAATTTAACCAGCAACATTGACCCAGAAATGTATGGCATGT GTAGTTAATTTTATTCCAGTAAGGAACGGCCCTTTTAGTTCTTAGAGCACTTTTAACAAAAAAGGAAAA TGAACAGGTTCTTTAAAATGCCAAGGAACGGGACAGAAAAGAAAG
<b>Restriction Sites:</b>	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 001018161.2</u>



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	PON2 (NM_001018161) Human 3' UTR Clone – SC206538
Summary:	This gene encodes a member of the paraoxonase gene family, which includes three known members located adjacent to each other on the long arm of chromosome 7. The encoded protein is ubiquitously expressed in human tissues, membrane-bound, and may act as a cellular antioxidant, protecting cells from oxidative stress. Hydrolytic activity against acylhomoserine lactones, important bacterial quorum-sensing mediators, suggests the encoded protein may also play a role in defense responses to pathogenic bacteria. Mutations in this gene may be associated with vascular disease and a number of quantitative phenotypes related to diabetes. Alternatively spliced transcript variants encoding different isoforms have been described. [provided by RefSeq, Jul 2008]
Locus ID:	5445
MW:	19.1

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