

## Product datasheet for **SC202043**

### Cyclophilin E (PPIE) (NM\_203456) Human 3' UTR Clone

#### Product data:

Product Type:	3' UTR Clones
Product Name:	Cyclophilin E (PPIE) (NM_203456) Human 3' UTR Clone
Symbol:	Cyclophilin E
Synonyms:	CYP-33; CYP33
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_203456
Insert Size:	202 bp
Insert Sequence:	<p>&gt;SC202043 3'UTR clone of NM_203456 The sequence shown below is from the reference sequence of NM_203456. The complete sequence of this clone may contain minor differences, such as SNPs. <b>Blue</b>=Stop Codon <b>Red</b>=Cloning site</p> <pre> GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAA<b>GCGATCGCC</b> AGCCAGCCGAGGTCCTGGAAGCTGACGT<b>AG</b>AGCTCGTGCCGACGGCAGACCTGCCGGCCGTGGGAGCCG TGGACGTCATCTGCAGGGACAGAAGGGCAAGGTCTTTTCTGGGGTTCTACTGTGTGCAGCTACTATG GGGTACCAGGGTGGGGGATGCCCTGATGAGCACATTTGTCAAATAAATGAATGACAGGAAACCA <b>ACGCGT</b>AAGCGGCCGCGCATCTAGATTGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTCGATTCCACCGCCGCTTCTATGAAAGG </pre>
Restriction Sites:	Sgfl-MluI
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><a href="#">NM_203456.3</a></u>



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**Summary:**

The protein encoded by this gene is a member of the peptidyl-prolyl cis-trans isomerase (PPIase) family. PPIases catalyze the cis-trans isomerization of proline imidic peptide bonds in oligopeptides and accelerate the folding of proteins. This protein contains a highly conserved cyclophilin (CYP) domain as well as an RNA-binding domain. It was shown to possess PPIase and protein folding activities, and it also exhibits RNA-binding activity. Alternative splicing results in multiple transcript variants. A related pseudogene, which is also located on chromosome 1, has been identified. [provided by RefSeq, Aug 2010]

**Locus ID:**

10450

**MW:**

7.4