OriGene Technologies, Inc.
9620 Medical Center Drive, Ste 200
Rockville, MD 20850, US
Phone: +1-888-267-4436
https://www.origene.com
techsupport@origene.com
EU: info-de@origene.com
CN: techsupport@origene.cn

Product datasheet for SC206680

CRYZL1 (NM_145858) Human 3' UTR Clone

Product data:

Product Type:	3' UTR Clones
Product Name:	CRYZL1 (NM_145858) Human 3' UTR Clone
Symbol:	CRYZL1
Synonyms:	4P11; QOH-1
Mammalian Cell Selection:	Neomycin
Vector:	pMirTarget (PS100062)
ACCN:	NM_145858
Insert Size:	511 bp
Insert Sequence:	>SC206680 3'UTR clone of NM_145858 The sequence shown below is from the reference sequence of NM_145858. The complete sequence of this clone may contain minor differences, such as SNPs. Blue=Stop Codon Red=Cloning site GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC GGAAGAAAAAAGCAAGTTGTTCAATTTTAATTTTCTTCTTTCTCAGACCTCAGTCGGATGAACATATTC CAGTATTTGAAGCCAGAATTTTCTTTGGAAATTGTTGAGAAAAACCAAGGAAGATAAAACAAGTTGCAT TTTTAAGCACGTTTCTCTGCTAAGACAAGATGCTCAGTTGACACATTTGAAAAGTGTTTGAAAAATTCT TGTGCAAATGATCAAGATAATTCTATAATTAACATCTTAAGGGAATTTTTCTAAAAACCTTTTCATTGT TTCTATATATTTTGCCCCTGCTATAAAATTCCTTCCATGAAGAAAACTGCTGCTTTCAGCAAAAGTCAC ACTACTCTTGATAAAAGCTGTTGCAGGCCTTTGCTAAGCTATCAAAGTAACGTATTAATTTTGTATCAA CTCCGTTCTCAACACCTTCCTTAAGTCTTTGCTGTCATAATTTAAGCATTTGAGTATATTTTGAAGTCT TAAAAGACTTAGCCCATAGGCACTTAAA ACGCGTAAGCGGCCGCGGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG
Restriction Sites:	SgfI-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).

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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:	NM_145858.3
Summary:	This gene encodes a protein that has sequence similarity to zeta crystallin, also known as quinone oxidoreductase. This zeta crystallin-like protein also contains an NAD(P)H binding site. Alternatively spliced transcript variants have been observed but their full-length nature has not been completely determined. [provided by RefSeq, Jul 2008]
Locus ID:	9946
MW:	19.7