

Product datasheet for **SC329183**

PARP8 (NM_001178055) Human Untagged Clone

Product data:

Product Type:	Expression Plasmids
Product Name:	PARP8 (NM_001178055) Human Untagged Clone
Tag:	Tag Free
Symbol:	PARP8
Synonyms:	ARTD16; pART16
Mammalian Cell Selection:	None
Vector:	<u>pCMV6-XL5</u>
E. coli Selection:	Ampicillin (100 ug/mL)



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Fully Sequenced ORF: >NCBI ORF sequence for NM_001178055, the custom clone sequence may differ by one or more nucleotides

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ATGGGGATGTGTTCAAGGCAAGAGCGAATTCAGAAGGATATCGACGTCGTGATCCAGAAG
TCCAGAGCTGAGAAGGACTGCCTGTTTGCAGATTCAGATACTCTGACTCCACCTTTACT
TTTACCTACGTTGGCGGCCCCAGAAGTGTATCCTACTCAGTACATGTATCTGAAGATTAC
CCAGATAATACATATGTGTCAAGTTCAGAGAATGATGAAGATGTGCTAGTTACTACAGAG
CCAATACCAGTAATTTTTCATAGAATAGCAACAGAATTAAGAAAAACAAATGACATTAAC
TGTTGCTTATCCATAAAATCCAAATTACAAAAGGAAAATGGGGAGGAATCAAGACAGAAT
AGTACAGTGGAGGAAGATTCTGAAGGTGACAATGATTCCGAAGAATTTTATTACGGAGGG
CAGGTGAACTATGATGGGGAAGTGCACAAGCACCCACAAGTGAAGCTGATTTGTCAGCA
GTTAGAGAGATATATGGGCCACATGCAGTTTCTCTCAGGGAATATGGAGCCATTGATGAT
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GAAGTAATTCGAACAGAACCTATAATTGTTGACTACACTGTTCACTTACACAGTATTTA
AATGGCCAGTGCCCACTGTTGATGTTTTCAGATTTCCACAAAAGAGCGATTTGGATTG
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GAAAAATCCAATTGCCTGCACAATAAAAAGTTGTGAGAGAAGAAAGTGAAGTCTCCCTG
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AAATCCAACTGAAATCTGAGCAGGACGGAATCTCCAAAACGCATAAGCTGCTGCGGAGG
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CGGATCCCTGTATTAATGAATATTGTGTGTTTGTGATGAGCCACATGTGTTTCAAAT
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CTGGGAGTAATGAATGAAGCTGCTGATGAAATAGCAACTGGAGCTCAGGTGGTAGATCTA
CTAGTATCCATGTGTAGGTCTGCGTTGGAATCTCCTAGAAAAGTTGTGATTTTCGAGCCA
TATCCTTCTGTGGTAGATCCTAATGATCCTCAGATGTTGGCTTCAACCCAGGAAAAAG
AACTATGATCGAGTAATGAAAGCACTGGATAGCATAAATTCTATCAGAGAATGACACAA
GCACCATATCTGGAAATCAAGAAGCAAATGGATAAACAGGACCCCTTGCTCATCCCTTA
CTGCAATGGGTTATATCAAGTAATAGATCACATATTGTGAAACTGCCAGTTAACAGGCAA
TTGAAGTTTATGCATACTCCACATCAGTTCCTTCTTCTCAGCAGTCCACCAGCCAAAGAA
TCCAATTTTAGAGCTGCTAAAAAATCTTTGGAAGCACCTTTGCATTTTATGGCTCACAC
ATTGAAAATGGCACTCCATCCTGAGGAATGGTCTGGTTGTTGCTTCTAATACACGATTG
CAGCTCCATGGTCAATGTATGGAAGTGAATCTATCTTAGTCCAATGTCAAGCATATCA
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CAAAGCCGTAACCTAAAAATGCATAGCCTTATGTGAAGTATCACCTCATCTGACCTGCAC
AAACATGGAGAGATATGGGTTGTCCCAATACTGACCATGTCTGCACACGATTCTTTTTTC
GTCTATGAAGACGGCCAAGTGGGAGATGCAAATATTAATACACAAGAAGGAGGCATTAC
AAAGAGATCCTCCGAGTAATTGGTAATCAAAGTCTACTGGTTAA
    
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Restriction Sites: Please inquire

ACCN: NM_001178055

OTI Disclaimer:	Our molecular clone sequence data has been matched to the reference identifier above as a point of reference. Note that the complete sequence of our molecular clones may differ from the sequence published for this corresponding reference, e.g., by representing an alternative RNA splicing form or single nucleotide polymorphism (SNP).
OTI Annotation:	This TrueClone is provided through our Custom Cloning Process that includes sub-cloning into OriGene's pCMV6 vector and full sequencing to provide a non-variant match to the expected reference without frameshifts, and is delivered as lyophilized plasmid DNA.
Components:	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
Reconstitution Method:	<ol style="list-style-type: none">1. Centrifuge at 5,000xg for 5min.2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.3. Close the tube and incubate for 10 minutes at room temperature.4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.
RefSeq:	<u>NM_001178055.1</u> , <u>NP_001171526.1</u>
RefSeq Size:	7201 bp
RefSeq ORF:	2565 bp
Locus ID:	79668
UniProt ID:	<u>Q8N3A8</u>
Cytogenetics:	5q11.1
Gene Summary:	<p>Mono-ADP-ribosyltransferase that mediates mono-ADP-ribosylation of target proteins. [UniProtKB/Swiss-Prot Function]</p> <p>Transcript Variant: This variant (1) represents the longest transcript and encodes the longest isoform (1). Variants 1 and 2 encode the same isoform. Sequence Note: This RefSeq record was created from transcript and genomic sequence data to make the sequence consistent with the reference genome assembly. The genomic coordinates used for the transcript record were based on transcript alignments.</p>