

**PARP1 Blocking Peptide (N-term)**  
Synthetic peptide  
Catalog # BP19834a**Specification****PARP1 Blocking Peptide (N-term) - Product Information**

Primary Accession [P09874](#)  
Other Accession [P11103](#),  
[NP\\_001609.2](#)

**PARP1 Blocking Peptide (N-term) - Additional Information**

**Gene ID** 142

**Other Names**

Poly [ADP-ribose] polymerase 1, PARP-1, ADP-ribosyltransferase diphtheria toxin-like 1, ARTD1, NAD(+) ADP-ribosyltransferase 1, ADPRT 1, Poly[ADP-ribose] synthase 1, PARP1, ADPRT, PPOL

**Target/Specificity**

The synthetic peptide sequence is selected from aa 138-151 of HUMAN PARP1

**Format**

Peptides are lyophilized in a solid powder format. Peptides can be reconstituted in solution using the appropriate buffer as needed.

**Storage**

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C.

**Precautions**

This product is for research use only. Not for use in diagnostic or therapeutic procedures.

**PARP1 Blocking Peptide (N-term) - Protein Information**

**Name** PARP1 ([HGNC:270](#))

**Function**

Poly-ADP-ribosyltransferase that mediates

**PARP1 Blocking Peptide (N-term) - Background**

This gene encodes a chromatin-associated enzyme, poly(ADP-ribosyl)transferase, which modifies various nuclear proteins by poly(ADP-ribosyl)ation. The modification is dependent on DNA and is involved in the regulation of various important cellular processes such as differentiation, proliferation, and tumor transformation and also in the regulation of the molecular events involved in the recovery of cell from DNA damage. In addition, this enzyme may be the site of mutation in Fanconi anemia, and may participate in the pathophysiology of type I diabetes.

**PARP1 Blocking Peptide (N-term) - References**

Majewski, P.M., et al. J. Biol. Chem. 285(45):34828-34838(2010)  
Kim, M., et al. Cancer Sci. 101(11):2436-2442(2010)  
Dong, Y., et al. Cancer Res. 70(20):8088-8096(2010)  
Krishnakumar, R., et al. Mol. Cell 39(5):736-749(2010)  
Lee, K.A., et al. Rheumatol. Int. (2010) In press  
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poly-ADP- ribosylation of proteins and plays a key role in DNA repair (PubMed:<a href="http://www.uniprot.org/citations/17177976" target="\_blank">17177976</a>, PubMed:<a href="http://www.uniprot.org/citations/18172500" target="\_blank">18172500</a>, PubMed:<a href="http://www.uniprot.org/citations/19344625" target="\_blank">19344625</a>, PubMed:<a href="http://www.uniprot.org/citations/19661379" target="\_blank">19661379</a>, PubMed:<a href="http://www.uniprot.org/citations/23230272" target="\_blank">23230272</a>, PubMed:<a href="http://www.uniprot.org/citations/25043379" target="\_blank">25043379</a>, PubMed:<a href="http://www.uniprot.org/citations/33186521" target="\_blank">33186521</a>, PubMed:<a href="http://www.uniprot.org/citations/32028527" target="\_blank">32028527</a>, PubMed:<a href="http://www.uniprot.org/citations/26344098" target="\_blank">26344098</a>). Mediates glutamate, aspartate, serine or tyrosine ADP-ribosylation of proteins: the ADP-D-ribosyl group of NAD(+) is transferred to the acceptor carboxyl group of target residues and further ADP-ribosyl groups are transferred to the 2'-position of the terminal adenosine moiety, building up a polymer with an average chain length of 20-30 units (PubMed:<a href="http://www.uniprot.org/citations/7852410" target="\_blank">7852410</a>, PubMed:<a href="http://www.uniprot.org/citations/9315851" target="\_blank">9315851</a>, PubMed:<a href="http://www.uniprot.org/citations/19764761" target="\_blank">19764761</a>, PubMed:<a href="http://www.uniprot.org/citations/25043379" target="\_blank">25043379</a>, PubMed:<a href="http://www.uniprot.org/citations/28190768" target="\_blank">28190768</a>, PubMed:<a href="http://www.uniprot.org/citations/29954836" target="\_blank">29954836</a>). Serine ADP- ribosylation of proteins constitutes the primary form of ADP- ribosylation of proteins in response to DNA damage

(PubMed:<a href="http://www.uniprot.org/citations/33186521" target="\_blank">33186521</a>). Mainly mediates glutamate and aspartate ADP-ribosylation of target proteins in absence of HPF1 (PubMed:<a href="http://www.uniprot.org/citations/19764761" target="\_blank">19764761</a>, PubMed:<a href="http://www.uniprot.org/citations/25043379" target="\_blank">25043379</a>). Following interaction with HPF1, catalyzes serine ADP-ribosylation of target proteins; HPF1 conferring serine specificity by completing the PARP1 active site (PubMed:<a href="http://www.uniprot.org/citations/28190768" target="\_blank">28190768</a>, PubMed:<a href="http://www.uniprot.org/citations/29954836" target="\_blank">29954836</a>, PubMed:<a href="http://www.uniprot.org/citations/33186521" target="\_blank">33186521</a>, PubMed:<a href="http://www.uniprot.org/citations/32028527" target="\_blank">32028527</a>). Also catalyzes tyrosine ADP-ribosylation of target proteins following interaction with HPF1 (PubMed:<a href="http://www.uniprot.org/citations/30257210" target="\_blank">30257210</a>, PubMed:<a href="http://www.uniprot.org/citations/29954836" target="\_blank">29954836</a>). PARP1 initiates the repair of DNA breaks: recognizes and binds DNA breaks within chromatin and recruits HPF1, licensing serine ADP-ribosylation of target proteins, such as histones, thereby promoting decompaction of chromatin and the recruitment of repair factors leading to the reparation of DNA strand breaks (PubMed:<a href="http://www.uniprot.org/citations/17177976" target="\_blank">17177976</a>, PubMed:<a href="http://www.uniprot.org/citations/18172500" target="\_blank">18172500</a>, PubMed:<a href="http://www.uniprot.org/citations/19344625" target="\_blank">19344625</a>, PubMed:<a href="http://www.uniprot.org/citations/19661379" target="\_blank">19661379</a>, PubMed:<a href="http://www.uniprot.org/citations/23230272" target="\_blank">23230272</a>).

target="\_blank">23230272</a>,  
PubMed:<a href="http://www.uniprot.org/citations/27067600"  
target="\_blank">27067600</a>). In  
addition to base excision repair (BER)  
pathway, also involved in double-strand  
breaks (DSBs) repair: together with  
TIMELESS, accumulates at DNA damage  
sites and promotes homologous  
recombination repair by mediating  
poly-ADP-ribosylation (PubMed:<a href="http://www.uniprot.org/citations/26344098"  
target="\_blank">26344098</a>,  
PubMed:<a href="http://www.uniprot.org/citations/30356214"  
target="\_blank">30356214</a>). Mediates  
the poly(ADP-ribosyl)ation of a number of  
proteins, including itself, APLF and CHFR  
(PubMed:<a href="http://www.uniprot.org/citations/17396150"  
target="\_blank">17396150</a>,  
PubMed:<a href="http://www.uniprot.org/citations/19764761"  
target="\_blank">19764761</a>). In  
addition to proteins, also able to  
ADP-ribosylate DNA: catalyzes  
ADP-ribosylation of DNA strand break  
termini containing terminal phosphates and  
a 2'-OH group in single- and  
double-stranded DNA, respectively  
(PubMed:<a href="http://www.uniprot.org/citations/27471034"  
target="\_blank">27471034</a>). Required  
for PARP9 and DTX3L recruitment to DNA  
damage sites (PubMed:<a href="http://www.uniprot.org/citations/23230272"  
target="\_blank">23230272</a>). PARP1-  
dependent PARP9-DTX3L-mediated  
ubiquitination promotes the rapid and  
specific recruitment of 53BP1/TP53BP1,  
UIMC1/RAP80, and BRCA1 to DNA damage  
sites (PubMed:<a href="http://www.uniprot.org/citations/23230272"  
target="\_blank">23230272</a>). Acts as a  
regulator of transcription: positively  
regulates the transcription of MTUS1 and  
negatively regulates the transcription of  
MTUS2/TIP150 (PubMed:<a href="http://www.uniprot.org/citations/19344625"  
target="\_blank">19344625</a>). Plays a  
role in the positive regulation of IFNG  
transcription in T-helper 1 cells as part of an  
IFNG promoter-binding complex with TXK  
and EEF1A1 (PubMed:<a href="http://www.uniprot.org/citations/17177976"  
target="\_blank">17177976</a>). Involved  
in the synthesis of ATP in the nucleus,

together with NMNAT1, PARG and NUDT5 (PubMed:<a href="http://www.uniprot.org/citations/27257257" target="\_blank">27257257</a>). Nuclear ATP generation is required for extensive chromatin remodeling events that are energy-consuming (PubMed:<a href="http://www.uniprot.org/citations/27257257" target="\_blank">27257257</a>).

**Cellular Location**

Nucleus. Nucleus, nucleolus. Chromosome  
Note=Localizes to sites of DNA damage.

**PARP1 Blocking Peptide (N-term) -  
Protocols**

Provided below are standard protocols that you may find useful for product applications.

- [Blocking Peptides](#)