

## 32-2644: PARP1 Recombinant Protein

**Alternative Name :** ADPRT,ADPRT1,pADPRT,pADPRT-1,PARP,PARP-1,PPOL,Poly [ADP-ribose] polymerase 1,NAD(+) ADP-ribosyltransferase 1,Poly[ADP-ribose] synthase 1,PARP1.

### Description

Source : Escherichia Coli. PARP1 Human Recombinant produced in E.Coli is a single, non-glycosylated, polypeptide chain containing 354 amino acids (662-1014a.a.) and having a molecular mass of 39.6 kDa. PARP1 is purified by proprietary chromatographic techniques. PARP1 takes part in the base excision repair pathway, by catalyzing the poly ADP-ribosyl of a restricted number of acceptor proteins involved in chromatin architecture and in DNA metabolism.. PARP1 mediates the poly ADP-ribose of APLF and CHFR. PARP1 positively regulates the transcription of MTUS1 and negatively regulates the transcription of MTUS2/TIP150. PARP1 is a chromatin-associated enzyme, poly (ADP-ribosyl) transferase, which modifies various nuclear proteins by poly ADP-ribosyl. PARP1 takes part in the regulation of various significant 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. PARP1 is a site of mutation in Fanconi anemia, and is involved in the pathophysiology of type I diabetes.

### Product Info

**Amount :** 25 µg  
**Purification :** Greater than 95% as determined by SDS-PAGE.  
**Content :** PARP1 solution containing 20mM Tris pH-8, 1Mm DTT and 10% glycerol.  
**Storage condition :** PARP1 Human Recombinant although stable at 4°C for 1 week, should be stored below -18°C. Please prevent freeze thaw cycles.  
**Amino Acid :** MKSKLPKPVQ DLIKMFIDVE SMKKAMVEYE IDLQKMPGK LSKRQIQAAAY SILSEVQQAV  
SQGSSDSQIL DLSNRFYTLI PHDFGMKKPP LLNNADSVQA KAEMLDNLLD IEVAYSLLRG  
GSDDSSKDPI DVNYEKLKTD IKVVDRDSEE AEIIRKYVKN THATTHNAYD LEVIDIFKIE  
REGECQRYKP FKQLHNRLL WHGSRTTNFA GILSQGLRIA PPEAPVTGYM FGKGIYFADM  
VSKSANYCHT SQGDPIGLIL LGEVALGNMY ELKHASHISK LPKGKHSVKG LGKTTDPDSA  
NISLDGVDVP LGTGISSGVN DTSLLYNEYI VYDIAQVNLK YLLKLFNFK TSLW.

