

Datasheet for ABIN7555031  
**PMS2 Protein (AA 1-862) (His tag)**



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## Overview

Quantity:	1 mg
Target:	PMS2
Protein Characteristics:	AA 1-862
Origin:	Human
Source:	HEK-293 Cells
Protein Type:	Recombinant
Purification tag / Conjugate:	This PMS2 protein is labelled with His tag.
Application:	Western Blotting (WB), SDS-PAGE (SDS)

## Product Details

Purpose:	Custom-made recombinat PMS2 Protein expressed in mammalian cells.
Sequence:	<p>MERAESSSTE PAKAIKPIDR KSVHQICSGQ VVLSLSTAVK ELVENS LDAG ATNIDLK LKD</p> <p>YGVDLIEVSD NGCGVEEENF EGLTLKHHTS KIQEFADLTQ VETFGFRGEA LSSLCALSDV</p> <p>TISTCHASAK VGTRL MFDHN GKIIQKTPYP RPRGTTVSVQ QLFSTLPVRH KEFQRNIKKE</p> <p>YAKMVQVLHA YCIISAGIRV SCTNQLGQ GK RQPVVCTGGS PSIKENIGSV FGQKQLQSLI</p> <p>PFVQLPPSDS VCEEYGLSCS DALHNLFYIS GFISQCTHGV GRSSDRQFF FINRRPCDPA</p> <p>KVCRLVNEVY HMYNRHQYPF VVLNISVDSE CVDINVTDPK RQILLQEEKL LLAVLKTS LI</p> <p>GMFDSVDNKL NVSQQLLDV EGNLIKMHAA DLEKPMVEKQ DQSPSLRTGE EKKDVSISRL</p> <p>REAFSLRHTT ENKPHSPKTP EPRRSPLGQK RGMLSSSTSG AISDKGVLRP QKEAVSSSHG</p> <p>PSDPTDRAEV EKDSGHGSTS VDSEGF SIPD TGS HCSSEYA ASSPGDRGSQ EHVDSQEKAP</p> <p>KTDDSFSDVD CHSNQEDTGC KFRVLPQPTN LATPNTKRFK KEEILSSSDI CQKLVNTQDM</p> <p>SASQVDVAVK INKKVPLDF SMSSLAKRIK QLHHEAQQSE GEQNYRK FRA KICPGENQAA</p>

EDELKREISK TMFAEMEIG QFNLGFIITK LNEIDIVDQ HATDEKYNFE MLQHTVLQG  
QRLIAPQTLN LTAVNEAVLI ENLEIFRKNG FDFVIDENAP VTERAKLISL PTSKNWTFGP  
QDVDELIFML SDSPGVMCRP SRVKQMFASR ACRKSVMIGT ALNTSEMKKL ITHMGEMDHP  
WNCPHGRPTM RHIANLGVIS QN **Sequence without tag. The proposed Purification-Tag is based on experiences with the expression system, a different complexity of the protein could make another tag necessary. In case you have a special request, please contact us.**

Characteristics:	<p>Key Benefits:</p> <ul style="list-style-type: none"><li>• Made to order protein - from design to production - by highly experienced protein experts.</li><li>• Protein expressed in mammalian cells and purified in one-step affinity chromatography</li><li>• The optimized expression system ensures reliability for intracellular, secreted and transmembrane proteins.</li><li>• State-of-the-art algorithm used for plasmid design (Gene synthesis).</li></ul> <p>This protein is a made-to-order protein and will be made for the first time for your order. Our experts in the lab try to ensure that you receive soluble protein.</p> <p>If you are not interested in a full length protein, please contact us for individual protein fragments.</p> <p>The big advantage of ordering our made-to-order proteins in comparison to ordering custom made proteins from other companies is that there is no financial obligation in case the protein cannot be expressed or purified.</p>
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Purity:	> 90 % as determined by Bis-Tris Page, Western Blot
Grade:	custom-made

Target Details

Target:	PMS2
Alternative Name:	PMS2 ( <a href="#">PMS2 Products</a> )
Background:	Mismatch repair endonuclease PMS2 (EC 3.1.-.-) (DNA mismatch repair protein PMS2) (PMS1 protein homolog 2),FUNCTION: Component of the post-replicative DNA mismatch repair system (MMR) (PubMed:30653781, PubMed:35189042). Heterodimerizes with MLH1 to form MutL alpha. DNA repair is initiated by MutS alpha (MSH2-MSH6) or MutS beta (MSH2-MSH3) binding to a dsDNA mismatch, then MutL alpha is recruited to the heteroduplex. Assembly of the MutL-MutS-heteroduplex ternary complex in presence of RFC and PCNA is sufficient to activate endonuclease activity of PMS2. It introduces single-strand breaks near the mismatch

## Target Details

and thus generates new entry points for the exonuclease EXO1 to degrade the strand containing the mismatch. DNA methylation would prevent cleavage and therefore assure that only the newly mutated DNA strand is going to be corrected. MutL alpha (MLH1-PMS2) interacts physically with the clamp loader subunits of DNA polymerase III, suggesting that it may play a role to recruit the DNA polymerase III to the site of the MMR. Also implicated in DNA damage signaling, a process which induces cell cycle arrest and can lead to apoptosis in case of major DNA damages. Possesses an ATPase activity, but in the absence of gross structural changes, ATP hydrolysis may not be necessary for proficient mismatch repair (PubMed:35189042). {ECO:0000269|PubMed:16873062, ECO:0000269|PubMed:18206974, ECO:0000269|PubMed:23709753, ECO:0000269|PubMed:30653781, ECO:0000269|PubMed:35189042}.

Molecular Weight:	95.8 kDa
UniProt:	<a href="#">P54278</a>
Pathways:	<a href="#">DNA Damage Repair</a> , <a href="#">Production of Molecular Mediator of Immune Response</a>

## Application Details

Application Notes:	In addition to the applications listed above we expect the protein to work for functional studies as well. As the protein has not been tested for functional studies yet we cannot offer a guarantee though.
Restrictions:	For Research Use only

## Handling

Format:	Liquid
Buffer:	The buffer composition is at the discretion of the manufacturer.
Handling Advice:	Avoid repeated freeze-thaw cycles.
Storage:	-80 °C
Storage Comment:	Store at -80°C.
Expiry Date:	12 months