

## 32-2153: AMPD2 Recombinant Protein

**Alternative Name :** Adenosine Monophosphate Deaminase 2, Adenosine Monophosphate Deaminase 2 (Isoform L), EC 3.5.4.6, SPG63, AMP Deaminase Isoform L, AMP Deaminase 2, AMPD Isoform L, AMPD, PCH9, AMP deaminase 2.

### Description

Source : Escherichia Coli. AMPD2 Human Recombinant produced in E.Coli is a single, non-glycosylated polypeptide chain containing 667 amino acids (236-879 a.a) and having a molecular mass of 77.0kDa. AMPD2 is fused to a 23 amino acid His-tag at N-terminus & purified by proprietary chromatographic techniques. Adenosine Monophosphate Deaminase 2, also known as AMPD2 is significant in purine metabolism by converting AMP to IMP. AMPD2 which functions as a homotetramer, is one of the three AMP deaminases shown in mammals. More than a few transcript variants encoding different isoforms have been discovered for AMPD2.

### Product Info

<b>Amount :</b>	10 µg
<b>Purification :</b>	Greater than 85% as determined by Analysis by SDS-PAGE.
<b>Content :</b>	AMPD2 protein solution (0.25mg/ml) containing Phosphate buffered saline (pH7.4) and 10% glycerol.
<b>Storage condition :</b>	Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods of time. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA). Avoid multiple freeze-thaw cycles.
<b>Amino Acid :</b>	MGSSHHHHHH SGLVPRGSH MGSDDLDAK SVVRALFIRE KYMALSLQSF CPTTRRYLQQ LAEKPLETRT YEQGPDTTPVSADAPVHPPAL EQHPYEHCEP STMPGDLGLG LRMVRGVVHV YTRREPDEHC SEVELPYDDL QEFVADVNVLMALIINGPIK SFCYRRLQYL SSKFQMHVLLNEMKELAAQK KVPHRDFYNI RKVDTHIHAS SCMNQKHLLR FIKRAMKRHLEEIVHVEQGR EQTLREVFES MNLTAYDLSV DTLDVHADRN TFHRFDKFNA KYNPIGESVL REIFIKTDNRVSGKYFAHII KEVMSDLEES KYQNAELRLS IYGRSRDEWD KLARWAVMHR VHSPNVRWLQVPRFLDVYR TKGQLANFQE MLENIFLPLF EATVHPASHP ELHLFLEHVDGFDSDVDESK PENHVFNLES PLPEAWVEED NPPYAYLYY TFANMAMLNH LRRQRGFHTF VLRPHCGEAGPIHHLVSAFM LAENISHGLL LRKAPVLQYL YYLAQIGIAM SPLSNNLSFL SYHRNPLPEYLSRGLMVLSL TDDPLQFHFT KEPLMEEYSI ATQVWKLSSC DMCELARNSVLMMSGFSHKVK SHWLGPNYTK EGPEGNDIRR TNVPDIRVG YRYETLCQELA LITQAVQSEM LETIPEEAGITMSPGPQ.

