

32-2522: MAT2A Recombinant Protein

Alternative Name : MATA2,MATII,SAMS2,MAT-2A,S-adenosylmethionine synthetase isoform type-2,AdoMet synthetase 2,Methionine adenosyltransferase 2,Methionine adenosyltransferase II,MAT2A,AMS2.

Description

Source : Escherichia Coli. MAT2A Human Recombinant fused with His tag (20 a.a.) at C-terminus produced in E.Coli is a single, non-glycosylated, polypeptide chain containing 415 amino acids and having a molecular mass of 45.8 kDa. The MAT2A is purified by proprietary chromatographic techniques. MAT2A is an important enzyme in cellular metabolism and catalyzes the formation of S-adenosylmethionine (SAME) from L-methionine and ATP. MAT2A is expressed in extrahepatic tissues. In liver, MAT2A expression associates with growth, dedifferentiation, and cancer. NF-kappa B and AP-1 are necessary for basal MAT2A expression in HepG2 cells and mediate the increase in MAT2A expression in response to TNF-alpha. Up-regulation of MAT2A provides growth improvement and s-adenosylmethionine and methylthioadenosine thus can block mitogenic signaling in colon cancer cells. Lower expression of both MAT2A and MAT2beta and interfere with leptin signaling in liver cancer cells.

Product Info

Amount : 20 µg
Purification : Greater than 95.0% as determined by SDS-PAGE.
Content : The MAT2A solution (1mg/ml) contains 20mM Tris pH-8 & 10% glycerol.
Storage condition : MAT2A although stable 4°C for 4 weeks, should be stored desiccated below -18°C. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA). Please prevent freeze-thaw cycles.
Amino Acid : MGSSHHHHHH SSGLVPRGSH MNGQLNGFHE AFIEEGTFLF TSESVGEGHP DKICDQISDA
VLDAHLQQDP DAKVACETVA KTG MILLAGE ITSRAAVDYQ KVVREAVKHI GYDDSSKGFY YKTCNVLVAL
EQQSPDIAQG VHLDRNEEDI GAGDQGLMFG YATDETEECM PLTIVLAHKL NAKLAELRRN GTLPWLRPDS
KTQVTVQYMQ DRGAVLPIRV HTIVISVQHD EEVCLDEM RD ALKEKVIKAV VPAKYLDEDT IYHLQPSGRF
VIGGPQGDAG LTGRKIIVDT YGGWGAHGGG AFSGKDYTKV DRSAAYAARW VAKSLVKGGL
CRRVLVQVSY AIGVSHPLSI SIFHYGTSQK SERELLEIVK KNFDLRPGVI VRDLDLKKPI YQRTAAYGHF
GRDSFPWEVP KKLKY.

