



METH rabbit pAb

| Catalog No | BYab-08950 |
|--------------------|---|
| Isotype | lgG |
| Reactivity | Human; Mouse;Rat |
| Applications | WB |
| Gene Name | MTR |
| Protein Name | METH |
| Immunogen | Synthesized peptide derived from human METH AA range: 1110-1160 |
| Specificity | This antibody detects endogenous levels of METH at Human/Mouse/Rat |
| Formulation | Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide. |
| Source | Polyclonal, Rabbit,IgG |
| Purification | The antibody was affinity-purified from rabbit serum by affinity-chromatography using specific immunogen. |
| Dilution | WB 1: 500-2000 |
| Concentration | 1 mg/ml |
| Purity | ≥90% |
| Storage Stability | -20°C/1 year |
| Synonyms | |
| Observed Band | |
| Cell Pathway | Cytoplasm . |
| Tissue Specificity | Widely expressed. Expressed at the highest levels in pancreas, heart, brain, skeletal muscle and placenta (PubMed:8968737, PubMed:8968735). Expressed at lower levels in lung, liver and kidney (PubMed:8968737, PubMed:8968735). |
| Function | catalytic activity:5-methyltetrahydrofolate + L-homocysteine = tetrahydrofolate + L-methionine.,cofactor:Binds 1 zinc ion per subunit.,cofactor:Methylcobalamin (MeCBL).,disease:Defects in MTR are the cause of methylcobalamin deficiency type G (cbIG) [MIM:250940]; also known as homocystinuria-megaloblastic anemia complementation type G. It is an autosomal recessive inherited disease that causes mental retardation, macrocytic anemia, and homocystinuria. Mild deficiency in MS activity could be associated with mild hyperhomocysteinemia, a risk factor for cardiovascular disease and possibly neural tube defects. MS mutations could also be involved in tumorigenesis.,disease:Defects in MTR may be a cause of susceptibility to folate-sensitive neural tube defects (folate-sensitive NTD) [MIM:601634]. The most common NTDs are open spina bifida |
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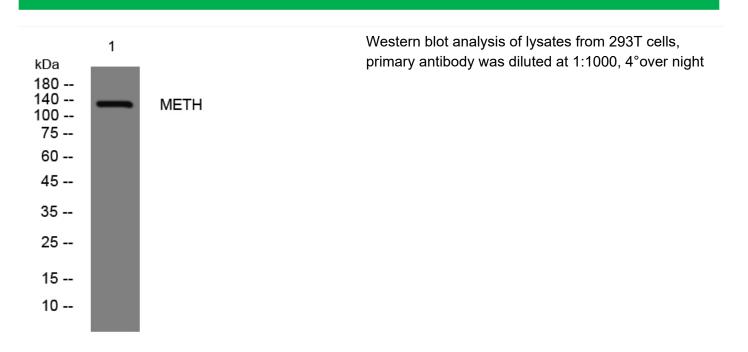




| Background | This gene encodes the 5-methyltetrahydrofolate-homocysteine methyltransferase. This enzyme, also known as cobalamin-dependent methionine synthase, catalyzes the final step in methionine biosynthesis. Mutations in MTR have been identified as the underlying cause of methylcobalamin deficiency complementation group G. Alternatively spliced transcript variants encoding distinct isoforms have been found for this gene. [provided by RefSeq, May 2014], |
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| matters needing attention | Avoid repeated freezing and thawing! |
| Usage suggestions | This product can be used in immunological reaction related experiments. For more information, please consult technical personnel. |

(myelomeningocele) and anencephaly. Genetic defects in M

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