



Anti-GCSH (full length) polyclonal antibody (DPAB-DC1311)

This product is for research use only and is not intended for diagnostic use.

PRODUCT INFORMATION

| | |
|----------------------------|---|
| Antigen Description | Degradation of glycine is brought about by the glycine cleavage system, which is composed of four mitochondrial protein components: P protein (a pyridoxal phosphate-dependent glycine decarboxylase), H protein (a lipoic acid-containing protein), T protein (a tetrahydrofolate-requiring enzyme), and L protein (a lipoamide dehydrogenase). The protein encoded by this gene is the H protein, which transfers the methylamine group of glycine from the P protein to the T protein. Defects in this gene are a cause of nonketotic hyperglycinemia (NKH). Two transcript variants, one protein-coding and the other probably not protein-coding, have been found for this gene. Also, several transcribed and non-transcribed pseudogenes of this gene exist throughout the genome.[provided by RefSeq, Jan 2010] |
| Immunogen | GCSH (AAH00790.1, 1 a.a. ~ 173 a.a) full-length recombinant protein with GST tag. The sequence is MALRVVRSVRALLCTLRVPLPAAPCPPRPWQLGVGAVRTLRTGPALLSVRKFTKEHEWV TTENGIGTVGISNFAQEALGDVVYCSLPEVGTKLNKQDEFGALESVKAASELYSPLSGEV TEINEALAENPGLV NKSCYEDGWLIKMTLSNPSELDELMSEEAYEKYIKSIEE |
| Source/Host | Mouse |
| Species Reactivity | Human |
| Conjugate | Unconjugated |
| Applications | WB (Recombinant protein), ELISA, |
| Size | 50 µl |
| Buffer | 50 % glycerol |
| Preservative | None |

Storage

Store at -20°C or lower. Aliquot to avoid repeated freezing and thawing.

GENE INFORMATION

| | |
|---------------------|---|
| Gene Name | GCSH glycine cleavage system protein H (aminomethyl carrier) [Homo sapiens (human)] |
| Official Symbol | GCSH |
| Synonyms | GCSH; glycine cleavage system protein H (aminomethyl carrier); GCE; NKH; glycine cleavage system H protein, mitochondrial; lipoic acid-containing protein; mitochondrial glycine cleavage system H-protein; |
| Entrez Gene ID | 2653 |
| Protein Refseq | NP_004474 |
| UniProt ID | P23434 |
| Chromosome Location | 16q23.2 |
| Pathway | Glycine, serine and threonine metabolism; Glyoxylate and dicarboxylate metabolism. |
| Function | aminomethyltransferase activity; enzyme binding; |