

O位 N-乙酰葡萄糖胺 OGT 抗体

产品货号: mIR6996

英文名称: O-GlcNAc transferase

中文名称: O 位 N-乙酰葡萄糖胺 OGT 抗体

别名: OGT; HRNT1; MGC22921; O-GlcNAc transferase; O GlcNAc; O-Linked N-Acetylglucosamine Transferase; O GlcNAc transferase p110 subunit; O GlcNAc transferase subunit p110; O linked N acetylglucosamine (GlcNAc) transferase (UDP N acetylglucosamine:polypeptide N acetylglucosaminyl transferase); O linked N acetylglucosamine transferase 110 kDa subunit; O-GlcNAc transferase subunit p110; O-linked N-acetylglucosamine transferase 110 kDa subunit; OGT1_HUMAN; UDP N acetylglucosamine peptide N acetylglucosaminyltransferase 110 kDa subunit; UDP-N-acetylglucosamine:polypeptide beta N acetylglucosaminyltransferase 110 kDa subunit; Uridinediphospho N acetylglucosamine:polypeptide beta N acetylglucosaminyl transferase.

研究领域: 神经生物学 信号转导 转录调节因子 Alzheimer's

抗体来源: Rabbit

克隆类型: Polyclonal

交叉反应 : Human, Mouse, Rat, Cow,

产品应用 : WB=1:500-2000 ELISA=1:500-1000 IHC-P=1:400-800 IHC-F=1:400-800 IF=1:50-200 (石蜡切片需做

抗原修复)

not yet tested in other applications.

optimal dilutions/concentrations should be determined by the end user.

分子量: 117kDa

细胞定位: 细胞核 细胞浆 细胞膜

性 状: Lyophilized or Liquid

浓 度: 1mg/ml

免疫原: KLH conjugated synthetic peptide derived from human O-GlcNAc transferase:951-1046/1046



亚 型: IgG

纯化方法: affinity purified by Protein A

储存液: 0.01M TBS(pH7.4) with 1% BSA, 0.03% Proclin300 and 50% Glycerol.

保存条件: Store at -20 °C for one year. Avoid repeated freeze/thaw cycles. The lyophilized antibody is stable at room temperature for at least one month and for greater than a year when kept at -20 °C. When reconstituted in sterile pH 7.4 0.01M PBS or diluent of antibody the antibody is stable for at least two weeks at 2-4 °C.

PubMed: PubMed

产品介绍: Addition of nucleotide-activated sugars directly onto the polypeptide through O-glycosidic linkage with the hydroxyl of serine or threonine. Mediates the O-glycosylation of MLL5 and HCFC1. Promotes proteolytic maturation of HCFC1. Since both phosphorylation and glycosylation compete for similar serine or threonine residues, the two processes may compete for sites, or they may alter the substrate specificity of nearby sites by steric or electrostatic effects. O-GlcNAc transferase has been purified from rat liver. It exists as a heterotrimeric complex with two subunits of the same molecular mass and one shorter subunit. Both polypeptides are related; the short subunit band is either a proteolytic product of the polypeptide or the product of an alternative translation start site. O-GlcNAc transferase is expressed as multiple transcripts that are present in different amounts in various human tissues, with the highest levels of expression in pancreas. Immunofluorescence of human cells expressing rat O-GlcNAc transferase indicated that it is present in both the nucleus and cytosol. HeLa cells expressing O-GlcNAc transferase do not survive well during prolonged incubations, suggesting that this protein may be toxic to the cells.

Function:

Catalyzes the transfer of a single N-acetylglucosamine from UDP-GlcNAc to a serine or threonine residue in cytoplasmic and nuclear proteins resulting in their modification with a beta-linked N-acetylglucosamine (O-GlcNAc). Glycosylates a large and diverse number of proteins including histone H2B, AKT1, MLL5, MAPT/TAU and HCFC1. Can regulate their cellular processes via cross-talk between glycosylation and phosphorylation or by affecting proteolytic processing. Involved in insulin resistance in muscle and adipocyte cells via glycosylating insulin signaling components and inhibiting the 'Thr-308' phosphorylation of AKT1, enhancing IRS1 phosphorylation and attenuating insulin signaling. Component of a THAP1/THAP3-HCFC1-OGT complex that is required for the regulation of the transcriptional activity of RRM1. As part of the NSL complex it may be involved in acetylation of nucleosomal histone H4 on several lysine residues.

Isoform 2, the mitochondrial isoform (mOGT), is cytotoxic and triggers apoptosis in several cell types including INS1, an insulinoma cell line.

Subunit:

Heterotrimer; consists of one 78 kDa subunit and two 110 kDa subunits dimerized via TPR repeats 6 and 7. Interacts (via TPR repeats 6 and 7) with ATXN10 (By similarity). Component of the MLL5-L complex, at least



composed of MLL5, STK38, PPP1CA, PPP1CB, HCFC1, PPP1CC and ACTB. Component of a THAP1/THAP3-HCFC1-OGT complex. Component of the NSL complex at least composed of MOF/KAT8, KANSL1, KANSL2, KANSL3, MCRS1, PHF20, OGT1/OGT, WDR5 and HCFC1. Interacts directly with HCFC1; the interaction O-glycosylates HCFC1, regulates its proteolytic processing and transcriptional activity and, in turn, stabilizes OGT in the nucleus. Interacts (via TPRs 1-6) with SIN3A; the interaction mediates transcriptional repression in parallel with histone deacetylase.

Subcellular Location:

Isoform 2: Mitochondrion. Membrane. Note=Associates with the mitochondrial inner membrane. Isoform 3: Cytoplasm. Nucleus. Cell membrane. Note=Mostly in the nucleus. Retained in the nucleus via interaction with HCFC1. After insulin induction, translocated from the nucleus to the cell membrane via phophatidylinisotide binding. Colocalizes with AKT1 at the plasma membrane. Isoform 4: Cytoplasm. Nucleus.

Tissue Specificity:

Highly expressed in pancreas and to a lesser extent in skeletal muscle, heart, brain and placenta. Present in trace amounts in lung and liver.

Post-translational modifications:

Ubiquitinated, leading to its proteasomal degradation.

DISEASE:

Note=Regulation of OGT activity and altered O-GlcNAcylations are implicated in diabetes and Alzheimer disease. O-GlcNAcylation of AKT1 affects insulin signaling and, possibly diabetes. Reduced O-GlcNAcylations and resulting increased phosphorylations of MAPT/TAU are observed in Alzheimer disease (AD) brain cerebrum.

Similarity:

Belongs to the O-GlcNAc transferase family.

Contains 13 TPR repeats.

SWISS:

P35858

Gene ID:

8473

Important Note:

This product as supplied is intended for research use only, not for use in human, therapeutic or diagnostic applications.

产品图片



