

Goat Anti-SEC61A1 Antibody

Peptide-affinity purified goat antibody Catalog # AF1965a

Specification

Goat Anti-SEC61A1 Antibody - Product Information

Application WB
Primary Accession P61619

Other Accession
Reactivity
Predicted
Host
Reactivity
Reactivity
Predicted
Fig. Dog, Cow
Goat

Clonality Polyclonal Concentration 100ug/200ul

Isotype IgG Calculated MW 52265

Goat Anti-SEC61A1 Antibody - Additional Information

Gene ID 29927

Other Names

Protein transport protein Sec61 subunit alpha isoform 1, Sec61 alpha-1, SEC61A1, SEC61A

Format

0.5 mg lgG/ml in Tris saline (20mM Tris pH7.3, 150mM NaCl), 0.02% sodium azide, with 0.5% bovine serum albumin

Storage

Maintain refrigerated at 2-8°C for up to 6 months. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.

Precautions

Goat Anti-SEC61A1 Antibody is for research use only and not for use in diagnostic or therapeutic procedures.

Goat Anti-SEC61A1 Antibody - Protein Information

Name SEC61A1

Synonyms SEC61A

Function

Component of SEC61 channel-forming translocon complex that mediates transport of signal peptide-containing precursor polypeptides across the endoplasmic reticulum (ER) (PubMed:12475939, PubMed:22375059, PubMed:<a



 $href="http://www.uniprot.org/citations/28782633" target="_blank">28782633, PubMed:29719251, PubMed:32814900). Forms a ribosome receptor and a gated pore in the ER membrane, both functions required for cotranslational translocation of nascent polypeptides (PubMed:32814900/a>).$

href="http://www.uniprot.org/citations/22375059" target="_blank">22375059, PubMed:28782633, PubMed:29719251). May cooperate with auxiliary protein SEC62, SEC63 and HSPA5/BiP to enable post-translational transport of small presecretory proteins (PubMed:22375059, PubMed:29719251). The SEC61 channel is also involved in ER membrane insertion of transmembrane proteins: it mediates membrane insertion of the first few transmembrane segments of proteins, while insertion of subsequent transmembrane regions of multi-pass membrane proteins is mediated by the multi-pass translocon (MPT) complex (PubMed:32820719, PubMed:36261522). The SEC61 channel cooperates with the translocating protein TRAM1 to import nascent proteins into the ER (PubMed:8616892). Controls the passive efflux of calcium ions from the ER lumen to the cytosol through SEC61 channel, contributing to the maintenance of cellular calcium homeostasis (PubMed:28782633, Plays a critical role in nephrogenesis, specifically at pronephros stage (By similarity).

Cellular Location

Endoplasmic reticulum membrane; Multi-pass membrane protein Note=Localizes exclusively in granular structures in the endoplasmic reticulum (ER).

Tissue Location

Expressed in proximal and distal tubules in kidney (at protein level).

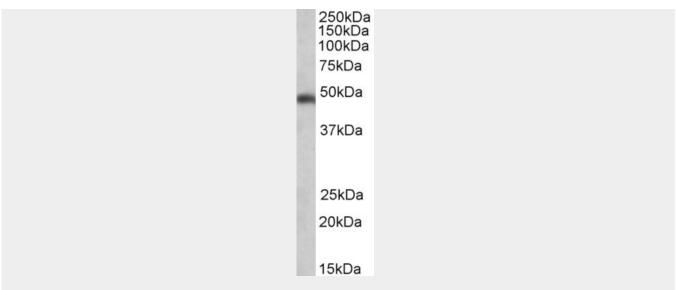
Goat Anti-SEC61A1 Antibody - Protocols

Provided below are standard protocols that you may find useful for product applications.

- Western Blot
- Blocking Peptides
- Dot Blot
- Immunohistochemistry
- Immunofluorescence
- <u>Immunoprecipitation</u>
- Flow Cytomety
- Cell Culture

Goat Anti-SEC61A1 Antibody - Images





AF1965a (0.3 μ g/ml) staining of Human Brain (Cerebellum) lysate (35 μ g protein in RIPA buffer). Primary incubation was 1 hour. Detected by chemiluminescence.

Goat Anti-SEC61A1 Antibody - Background

The protein encoded by this gene belongs to the SECY/SEC61- alpha family. It appears to play a crucial role in the insertion of secretory and membrane polypeptides into the endoplasmic reticulum. This protein found to be tightly associated with membrane-bound ribosomes, either directly or through adaptor proteins. This gene encodes an alpha subunit of the heteromeric SEC61 complex, which also contains beta and gamma subunits.

Goat Anti-SEC61A1 Antibody - References

Defining the human deubiquitinating enzyme interaction landscape. Sowa ME, et al. Cell, 2009 Jul 23. PMID 19615732.

Control of translocation through the Sec61 translocon by nascent polypeptide structure within the ribosome. Daniel CJ, et al. J Biol Chem, 2008 Jul 25. PMID 18480044.

Large-scale mapping of human protein-protein interactions by mass spectrometry. Ewing RM, et al. Mol Syst Biol, 2007. PMID 17353931.

Signal sequence and keyword trap in silico for selection of full-length human cDNAs encoding secretion or membrane proteins from oligo-capped cDNA libraries. Otsuki T, et al. DNA Res, 2005. PMID 16303743.

The status, quality, and expansion of the NIH full-length cDNA project: the Mammalian Gene Collection (MGC). Gerhard DS, et al. Genome Res, 2004 Oct. PMID 15489334.