

Product datasheet for PH302062

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PEX5 (NM 000319) Human Mass Spec Standard

Product data:

Product Type: Mass Spec Standards

Description: PEX5 MS Standard C13 and N15-labeled recombinant protein (NP 000310)

Species: Human **HEK293 Expression Host: Expression cDNA Clone**

or AA Sequence:

RC202062

Predicted MW: 69.9 kDa

>RC202062 protein sequence **Protein Sequence:**

Red=Cloning site Green=Tags(s)

MAMREL VEAECGGANPLMKLAGHFTQDKALRQEGLRPGPWPPGAPASEAASKPLGVASEDEL VAEFLQDQ NAPLVSRAPQTFKMDDLLAEMQQIEQSNFRQAPQRAPGVADLALSENWAQEFLAAGDAVDVTQDYNETDW SQEFISEVTDPLSVSPARWAEEYLEQSEEKLWLGEPEGTATDRWYDEYHPEEDLQHTASDFVAKVDDPKL ANSEFLKFVRQIGEGQVSLESGAGSGRAQAEQWAAEFIQQQGTSDAWVDQFTRPVNTSALDMEFERAKSA IELQAELEEMAKRDAEAHPWLSDYDDLTSATYDKGYQFEEENPLRDHPQPFEEGLRRLQEGDLPNAVLLF EAAVQQDPKHMEAWQYLGTTQAENEQELLAISALRRCLELKPDNQTALMALAVSFTNESLQRQACETLRD WLRYTPAYAHLVTPAEEGAGGAGLGPSKRILGSLLSDSLFLEVKELFLAAVRLDPTSIDPDVQCGLGVLF NLSGEYDKAVDCFTAALSVRPNDYLLWNKLGATLANGNQSEEAVAAYRRALELQPGYIRSRYNLGISCIN LGAHREAVEHFLEALNMORKSRGPRGEGGAMSENIWSTLRLALSMLGQSDAYGAADARDLSTLLTMFGLP

Q

SGPTRTRPLEQKLISEEDLAANDILDYKDDDDK**V**

C-Myc/DDK Tag:

Purity: > 80% as determined by SDS-PAGE and Coomassie blue staining

Concentration: >0.05 µg/µL as determined by microplate BCA method

Labeling Method: Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine

Buffer: 25 mM Tris-HCl, 100 mM glycine, pH 7.3

Storage: Store at -80°C. Avoid repeated freeze-thaw cycles.

Stability: Stable for 3 months from receipt of products under proper storage and handling conditions.

RefSeq: NP 000310

RefSeg Size: 3190





RefSeq ORF: 1893

Synonyms: PBD2A; PBD2B; PTS1-BP; PTS1R; PXR1; RCDP5

Locus ID: 5830

UniProt ID: <u>P50542</u>, <u>A0A0S2Z480</u>

Cytogenetics: 12p13.31

Summary: The product of this gene binds to the C-terminal PTS1-type tripeptide peroxisomal targeting

signal (SKL-type) and plays an essential role in peroxisomal protein import. Peroxins (PEXs) are proteins that are essential for the assembly of functional peroxisomes. The peroxisome biogenesis disorders (PBDs) are a group of genetically heterogeneous autosomal recessive, lethal diseases characterized by multiple defects in peroxisome function. The peroxisomal biogenesis disorders are a heterogeneous group with at least 14 complementation groups

and with more than 1 phenotype being observed in cases falling into particular

complementation groups. Although the clinical features of PBD patients vary, cells from all PBD patients exhibit a defect in the import of one or more classes of peroxisomal matrix

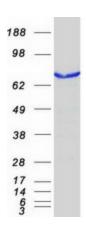
proteins into the organelle. Defects in this gene are a cause of neonatal

adrenoleukodystrophy (NALD), a cause of Zellweger syndrome (ZWS) as well as may be a cause of infantile Refsum disease (IRD). Alternatively spliced transcript variants encoding

different isoforms have been identified. [provided by RefSeq, Oct 2008]

Protein Families: Druggable Genome

Product images:



Coomassie blue staining of purified PEX5 protein (Cat# [TP302062]). The protein was produced from HEK293T cells transfected with PEX5 cDNA clone (Cat# [RC202062]) using MegaTran 2.0 (Cat# [TT210002]).