

Full-Length

SPG21 (Human) Recombinant Protein (P01)

Catalog # H00051324-P01

Size 10 ug, 25 ug

Applications



Specification	
Product Description	Human SPG21 full-length ORF (NP_057714.1, 1 a.a 308 a.a.) recombinant protein with GST-tag a t N-terminal.
Sequence	MGEIKVSPDYNWFRGTVPLKKIIVDDDDSKIWSLYDAGPRSIRCPLIFLPPVSGTADVFFRQILALTG WGYRVIALQYPVYWDHLEFCDGFRKLLDHLQLDKVHLFGASLGGFLAQKFAEYTHKSPRVHSLIL CNSFSDTSIFNQTWTANSFWLMPAFMLKKIVLGNFSSGPVDPMMADAIDFMVDRLESLGQSELA SRLTLNCQNSYVEPHKIRDIPVTIMDVFDQSALSTEAKEEMYKLYPNARRAHLKTGGNFPYLCRSA EVNLYVQIHLLQFHGTKYAAIDPSMVSAEELEVQKGSLGISQEEQ
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	61.4
Interspecies Antigen Sequence	Mouse (98); Rat (82)
Preparation Method	in vitro wheat germ expression system
Purification	Glutathione Sepharose 4 Fast Flow
Quality Control Testing	12.5% SDS-PAGE Stained with Coomassie Blue.
Storage Buffer	50 mM Tris-HCI, 10 mM reduced Glutathione, pH=8.0 in the elution buffer.

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Product Information

Storage Instruction

Store at -80°C. Aliquot to avoid repeated freezing and thawing.

Note

Best use within three months from the date of receipt of this protein.

Applications

- Enzyme-linked Immunoabsorbent Assay
- Western Blot (Recombinant protein)
- Antibody Production
- Protein Array

Gene Info — SPG21

Entrez GenelD	<u>51324</u>
GeneBank Accession#	<u>NM_016630.3</u>
Protein Accession#	<u>NP_057714.1</u>
Gene Name	SPG21
Gene Alias	ACP33, BM-019, GL010, MASPARDIN, MAST
Gene Description	spastic paraplegia 21 (autosomal recessive, Mast syndrome)
Omim ID	<u>248900 608181</u>
Gene Ontology	Hyperlink
Gene Summary	The protein encoded by this gene was identified by a two-hybrid screen using CD4 as the bait. It binds to the hydrophobic C-terminal amino acids of CD4 which are involved in repression of T cell activation. The interaction with CD4 is mediated by the noncatalytic alpha/beta hydrolase fold do main of this protein. It is thus proposed that this gene product modulates the stimulatory activity of CD4. At least three different transcript variants encoding two different isoforms have been found f or this gene. [provided by RefSeq
Other Designations	acid cluster protein 33 spastic paraplegia 21