



Full-Length

LILRA5 (Human) Recombinant Protein

Catalog # H00353514-G01 Size 10 ug

Specification	
Product Description	Human LILRA5 full-length ORF (NP_067073.1) recombinant protein without tag. This product is belong to Proteoliposome (PL).
Sequence	MAPWSHPSAQLQPVGGDAVSPALMVLLCLGLSLGPRTHVQAGNLSKATLWAEPGSVISRGNSV TIRCQGTLEAQEYRLVKEGSPEPWDTQNPLEPKNKARFSIPSMTEHHAGRYRCYYYSPAGWSEP SDPLELVVTGFYNKPTLSALPSPVVTSGENVTLQCGSRLRFDRFILTEEGDHKLSWTLDSQLTPS GQFQALFPVGPVTPSHRWMLRCYGSRRHILQVWSEPSDLLEIPVSGAADNLSPSQNKSDSGTAS HLQDYAVENLIRMGMAGLILVVLGILIFQDWHSQRSPQAAAGR
Host	Wheat Germ (in vitro)
Theoretical MW (kDa)	32.8
Interspecies Antigen Sequence	Mouse (57); Rat (58)
Form	Liquid
Preparation Method	in vitro wheat germ expression system with proprietary liposome technology
Purification	None
Recommend Usage	Heating may cause protein aggregation. Please do not heat this product before electrophoresis.
Storage Buffer	25 mM Tris-HCl of pH8.0 containing 2% glycerol.
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

Antibody Production



Product Information

Gene Info — LILRA5

Entrez GenelD	<u>353514</u>
GeneBank Accession#	NM_021250.2
Protein Accession#	NP_067073.1
Gene Name	LILRA5
Gene Alias	CD85, CD85F, ILT11, LILRB7, LIR9
Gene Description	leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 5
Omim ID	606047
Gene Ontology	<u>Hyperlink</u>
Gene Ontology Gene Summary	Hyperlink The protein encoded by this gene is a member of the leukocyte immunoglobulin-like receptor (LIR) family. LIR family members are known to have activating and inibitory functions in leukocytes. Cr osslink of this receptor protein on the surface of monocytes has been shown to induce calcium flux and secretion of several proinflammatory cytokines, which suggests the roles of this protein in trig gering innate immune responses. This gene is one of the leukocyte receptor genes that form a ge ne cluster on the chromosomal region 19q13.4. Four alternatively spliced transcript variants enco ding distinct isoforms have been described. [provided by RefSeq

Disease

Hepatitis C