

Produktinformation



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Datasheet

KLRC1 (Human) Recombinant Protein

Catalog Number: P10272

Regulation Status: For research use only (RUO)

Product Description: Human KLRC1 (P26715-1, Arg100-Leu233) partial recombinant protein with His-Avi tag at N-terminus expressed in HEK293 cells.

Sequence: Arg100-Leu233

Host: Human

Theoretical MW (kDa): 18.3

Protocols: See our web site at http://www.abnova.com/support/protocols.asp or product page for detailed protocols

Form: Lyophilized

Preparation Method: Mammalian cell (HEK293) expression system

Purity: > 95% as determined by Tris-Bis PAGE;> 95% as determined by HPLC

Endotoxin Level: < 1 EU per 1 ug of protein (determined by LAL method)

Recommend Usage: Biological Activity ELISA Tris-Bis PAGE The optimal working dilution should be determined by the end user.

Storage Buffer: Lyophilized from filtered solution in PBS, pH 7.4 (5% trehalose).

Storage Instruction: After reconstitution with deionized water to a final concentration more than 100 ug/ml, store at 4°C for 1 week. For long term storage, store at -80°C for 1 year.

Aliquot to avoid repeated freezing and thawing.

Entrez GenelD: 3821

Gene Symbol: KLRC1

Gene Alias: CD159A, MGC13374, MGC59791, NKG2, NKG2A

Gene Summary: Natural killer (NK) cells are

lymphocytes that can mediate lysis of certain tumor cells and virus-infected cells without previous activation. They can also regulate specific humoral and cell-mediated immunity. The protein encoded by this gene belongs to the killer cell lectin-like receptor family, also called NKG2 family, which is a group of transmembrane proteins preferentially expressed in NK cells. This family of proteins is characterized by the type II membrane orientation and the presence of a C-type lectin domain. This protein forms a complex with another family member, KLRD1/CD94, and has been implicated in the recognition of the MHC class I HLA-E molecules in NK cells. The genes of NKG2 family members form a killer cell lectin-like receptor gene cluster on chromosome 12. Four alternatively spliced transcript variants encoding two distinct isoforms have been observed. [provided by RefSeq]