

Product datasheet

Recombinant Rabbit CTLA4 protein (Fc Chimera) ab219704

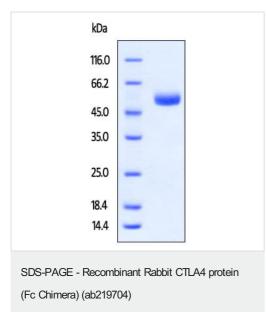
1 Image

Description

Product name	Recombinant Rabbit CTLA4 protein (Fc Chimera)		
Purity	> 95 % SDS-PAGE.		
Endotoxin level	< 1.000 Eu/µg		
Expression system	HEK 293 cells		
Accession	<u>P42072</u>		
Protein length	Protein fragment		
Animal free	No		
Nature	Recombinant		
Species	Rabbit		
Sequence		KALHVSQPAVVLASSRGVASFVCEYASSHKATEVRVTVL RQANSQMTEVC AMTYTVENELTFIDDSTCTGISHGNKVNLTIQGLSAMDTGLY ICKVELMY PPPYYVGMGNGTQIYVIEPEPCPDSD	
Predicted molecular weight	40 kDa including tags		
Amino acids	36 to 161		
Additional sequence information	Fused with a human IgG1 Fc tag at the C-terminus.		
Specifications			
Our Abpromise guarantee covers the use of ab219704 in the following tested applications.			
The application notes include recommended starting dilutions; optimal dilutions/concentrations should be determined by the end user.			
Applications	SDS-PAGE		
Form	Lyophilized		
Preparation and Storage			
Stability and Storage	Shipped at 4°C. Store at 4°C (sta thaw cycle.	ble for up to 12 months). Store at -20°C or -80°C. Avoid freeze /	

Reconstitution General Info	pH: 7.4 Constituents: 0.61% Tris, 0.75% Glycine, L-Arginine, Sodium chloride, 5% Trehalose Lyophilised from 0.22 µm filtered solution. Reconstitute with sterile deionized water to a concentration of 200 µg/ml.
Function	Inhibitory receptor acting as a major negative regulator of T-cell responses. The affinity of CTLA4 for its natural B7 family ligands, CD80 and CD86, is considerably stronger than the affinity of their cognate stimulatory coreceptor CD28.
Tissue specificity	Widely expressed with highest levels in lymphoid tissues. Detected in activated T-cells where expression levels are 30- to 50-fold less than CD28, the stimulatory coreceptor, on the cell surface following activation.
Involvement in disease	 Genetic variation in CTLA4 influences susceptibility to systemic lupus erythematosus (SLE) [MIM:152700]. SLE is a chronic, inflammatory and often febrile multisystemic disorder of connective tissue. It affects principally the skin, joints, kidneys and serosal membranes. SLE is thought to represent a failure of the regulatory mechanisms of the autoimmune system. Note=Genetic variations in CTLA4 may influence susceptibility to Graves disease, an autoimmune disorder associated with overactivity of the thyroid gland and hyperthyroidism. Genetic variation in CTLA4 is the cause of susceptibility to diabetes mellitus insulin-dependent type 12 (IDDM12) [MIM:601388]. A multifactorial disorder of glucose homeostasis that is characterized by susceptibility to ketoacidosis in the absence of insulin therapy. Clinical fetaures are polydipsia, polyphagia and polyuria which result from hyperglycemia-induced osmotic diuresis and secondary thirst. These derangements result in long-term complications that affect the eyes, kidneys, nerves, and blood vessels. Genetic variation in CTLA4 is the cause of susceptibility to celiac disease type 3 (CELIAC3) [MIM:609755]. It is a multifactorial disorder of the small intestine that is influenced by both environmental and genetic factors. It is characterized by malabsorption resulting from inflammatory injury to the mucosa of the small intestine after the ingestion of wheat gluten or related rye and barley proteins. In its classic form, celiac disease is characterized in children by malabsorption and failure to thrive.
Sequence similarities	Contains 1 lg-like V-type (immunoglobulin-like) domain.
Post-translational modifications	N-glycosylation is important for dimerization. Phosphorylation at Tyr-201 prevents binding to the AP-2 adapter complex, blocks endocytosis, and leads to retention of CTLA4 on the cell surface.
Cellular localization	Cell membrane. Exists primarily an intracellular antigen whose surface expression is tightly regulated by restricted trafficking to the cell surface and rapid internalisation and.

Images



SDS-PAGE analysis of ab219704 stained overnight with Coomassie Blue.

The reduced protein migrates as 50-55 kDa in SDS-PAGE due to glycosylation.

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