

C1S (light chain, Cleaved-Ile438) rabbit pAb

Cat No.:ES19982

For research use only

Overview

Product Name	C1S (light chain, Cleaved-Ile438) rabbit pAb
Host species	Rabbit
Applications	WB; ELISA
Species Cross-Reactivity	Human;Rat;Mouse;
Recommended dilutions	WB 1:1000-2000 ELISA 1:5000-20000
Immunogen	Synthesized peptide derived from human C1S (light
	chain, Cleaved-Ile438)
Specificity	This antibody detects endogenous levels of Human
	C1S (light chain, Cleaved-Ile438, protein was cleaved
	amino acid sequence between 437-438)
Formulation	Liquid in PBS containing 50% glycerol, 0.5% BSA and
	0.02% sodium azide.
Storage	Store at -20 $^\circ\!\mathrm{C}$. Avoid repeated freeze-thaw cycles.
Protein Name	C1S (light chain, Cleaved-Ile438)
Gene Name	C1S
Cellular localization	
Purification	The antibody was affinity-purified from rabbit
	antiserum by affinity-chromatography using
	epitope-specific immunogen.
Clonality	Polyclonal
Concentration	1 mg/ml
Observed band	28kD
Human Gene ID	716
Human Swiss-Prot Number	P09871
Alternative Names	Complement C1s subcomponent (EC 3.4.21.42;C1
	esterase;Complement component 1 subcomponent
	s) [Cleaved into: Complement C1s subcomponent
	heavy chain; Complement C1s subcomponent light
	chain]
Background	catalytic activity:Cleavage of Arg- -Ala bond in
	complement component C4 to form C4a and C4b,
	and Lys(or Arg)- -Lys bond in complement
	component C2 to form C2a and C2b: the 'classical'
-	



+86-27-59760950

ELKbio@ELKbiotech.com

www.elkbiotech.com

23-2, No.388 Gaoxin 2nd Road, Wuhan East Lake Hi-tech Development Zone, Hubei , P.R.C



pathway C3 convertase., disease: Defects in C1S are the cause of selective C1s deficiency [MIM:120580]; that is associated with early onset multiple autoimmune diseases., enzyme regulation: Inhibited by SERPING1., function: C1s B chain is a serine protease that combines with C1q and C1s to form C1, the first component of the classical pathway of the complement system. C1r activates C1s so that it can, in turn, activate C2 and C4., online information:C1S mutation db,PTM:The iron and 2-oxoglutarate dependent 3-hydroxylation of aspartate and asparagine is (R) stereospecific within EGF domains., similarity: Belongs to the peptidase S1 family., similarity: Contains 1 EGF-like domain., similarity: Contains 1 peptidase S1 domain., similarity: Contains 2 CUB domains., similarity: Contains 2 Sushi (CCP/SCR) domains., subunit: C1 is a calcium-dependent trimolecular complex of C1q, C1r and C1s in the molar ration of 1:2:2. Activated C1s is an disulfide-linked heterodimer of a heavy chain and a light chain.,



+86-27-59760950

ELKbio@ELKbiotech.com

www.elkbiotech.com

23-2, No.388 Gaoxin 2nd Road, Wuhan East Lake Hi-tech Development Zone, Hubei , P.R.C