

ARG66938 anti-Bcl 6 antibody [SQab22268]

Package: 100 μl, 50 μl Store at: -20°C

Summary

Product Description	Recombinant Rabbit Monoclonal antibody [SQab22268] recognizes Bcl 6
Tested Reactivity	Hu
Tested Application	IHC-P
Host	Rabbit
Clonality	Monoclonal
Clone	SQab22268
Isotype	lgG
Target Name	Bcl 6
Species	Human
Immunogen	Recombinant fragment within Human Bcl 6.
Conjugation	Un-conjugated
Alternate Names	ZNF51; LAZ3; Zinc finger protein 51; Protein LAZ-3; Zinc finger and BTB domain-containing protein 27; B- cell lymphoma 6 protein; B-cell lymphoma 5 protein; BCL5; BCL6A; BCL-5; ZBTB27; BCL-6

Application Instructions

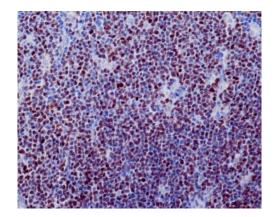
Application table	Application	Dilution
	IHC-P	1:100 - 1:200
Application Note	IHC-P: Antigen Retrieval: Heat mediated was performed using Tris/EDTA buffer (pH 9.0). Incubate the samples at RT (18-25°C) for 30 min. * The dilutions indicate recommended starting dilutions and the optimal dilutions or concentrations should be determined by the scientist.	
Positive Control	Human diffuse large B cell I	lymphoma tissue

Properties

Form	Liquid
Purification	Purification with Protein A.
Buffer	PBS, 0.01% Sodium azide, 40% Glycerol and 0.05%BSA.
Preservative	0.01% Sodium azide
Stabilizer	40% Glycerol and 0.05%BSA
Storage instruction	For continuous use, store undiluted antibody at 2-8°C for up to a week. For long-term storage, aliquot and store at -20°C or below. Storage in frost free freezers is not recommended. Avoid repeated freeze/thaw cycles. Suggest spin the vial prior to opening. The antibody solution should be gently mixed before use.

Bioinformation

Gene Symbol	BCL6
Gene Full Name	B-cell CLL/lymphoma 6
Background	The protein encoded by this gene is a zinc finger transcription factor and contains an N-terminal POZ domain. This protein acts as a sequence-specific repressor of transcription, and has been shown to modulate the transcription of STAT-dependent IL-4 responses of B cells. This protein can interact with a variety of POZ-containing proteins that function as transcription corepressors. This gene is found to be frequently translocated and hypermutated in diffuse large-cell lymphoma (DLCL), and may be involved in the pathogenesis of DLCL. Alternatively spliced transcript variants encoding different protein isoforms have been found for this gene. [provided by RefSeq, Aug 2015]
Function	Transcriptional repressor mainly required for germinal center (GC) formation and antibody affinity maturation which has different mechanisms of action specific to the lineage and biological functions. Forms complexes with different corepressors and histone deacetylases to repress the transcriptional expression of different subsets of target genes. Represses its target genes by binding directly to the DNA sequence 5'-TTCCTAGAA-3' (BCL6-binding site) or indirectly by repressing the transcriptional activity of transcription factors. In GC B-cells, represses genes that function in differentiation, inflammation, apoptosis and cell cycle control, also autoregulates its transcriptional expression and upregulates, indirectly, the expression of some genes important for GC reactions, such as AICDA, through the repression of microRNAs expression, like miR155. An important function is to allow GC B-cells to proliferate very rapidly in response to T-cell dependent antigens and tolerate the physiological DNA breaks required for immunglobulin class switch recombination and somatic hypermutation without inducing a p53/TP53-dependent apoptotic response. In follicular helper CD4(+) T-cells (T(FH) cells), promotes the expression of T(FH)-related genes but inhibits the differentiation of T(H)1, T(H)2 and T(H)17 cells. Also required for the establishment and maintenance of immunological memory for both T- and B-cells. Suppresses macrophage proliferation through competition with STAT5 for STAT-binding motifs binding on certain target genes, such as CCL2 and CCND2. In response to genotoxic stress, controls cell cycle arrest in GC B-cells in both p53/TP53-dependedent and -independent manners. Besides, also controls neurogenesis through the alteration of the composition of NOTCH-dependent transcriptional complexes at selective NOTCH targets, such as HES5, including the recruitment of the deacetylase SIRT1 and resulting in an epigenetic silencing leading to neuronal differentiation. [UniProt]
Calculated Mw	79 kDa
PTM	Phosphorylated by MAPK1 in response to antigen receptor activation at Ser-333 and Ser-343. Phosphorylated by ATM in response to genotoxic stress. Phosphorylation induces its degradation by ubiquitin/proteasome pathway.
	Polyubiquitinated. Polyubiquitinated by SCF(FBXO11), leading to its degradation by the proteasome.
	Acetylated at Lys-379 by EP300 which inhibits the interaction with NuRD complex and the transcriptional repressor function. Deacetylated by HDAC- and SIR2-dependent pathways. [UniProt]
Cellular Localization	Nucleus. [UniProt]



ARG66938 anti-Bcl 6 antibody [SQab22268] IHC-P image

Immunohistochemistry: Formalin/PFA-fixed and paraffin-embedded sections of Human diffuse large B cell lymphoma tissue stained with ARG66938 anti-Bcl 6 antibody [SQab22268]. Antigen Retrieval: Heat tissue section in Tris-EDTA buffer (pH 9.0).