

ARG55734 anti-HIF-1 alpha antibody

Package: 100 μg Store at: -20°C

Summary

Product DescriptionGoat Polyclonal antibody recognizes HIF-1 alphaTested ReactivityHu, Ms, RatTested ApplicationWBSpecificityDetects HIF1a in transfected cells and a band of approximately 120 kDa by Western blot in cell lysates.HostGoatClonalityPolyclonalIsotypeIgGTarget NameHIF-1 alphaSpeciesHumanImmunogenRecombinant peptide around aa. 780 (C-terminus) of Human HIF-1 alphaConjugationUn-conjugatedAlternate NamesClass E basic helix-loop-helix protein 78; Basic-helix-loop-helix-PAS protein MOP1; HIF01; HIF-1-alpha; PASD8; HIF-1alpha; Member of PAS protein 1; ARNT-interacting protein; bHLHe78		
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Application Instructions

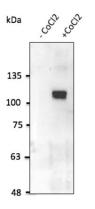
Application table	Application	Dilution
	WB	1:500-1:2000
Application Note	* The dilutions indicate recommended starting dilutions and the optimal dilutions or concentrations should be determined by the scientist.	
Observed Size	92-130 kDa	

Properties

Form	Liquid
Purification	Affinity purification with immunogen.
Buffer	PBS, 0.05% Sodium azide and 20% Glycerol
Preservative	0.05% Sodium azide
Stabilizer	20% Glycerol
Concentration	3 mg/ml
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. 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

Cono Symbol	HIF1A
Gene Symbol Gene Full Name	hypoxia inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor)
Background	This gene encodes the alpha subunit of transcription factor hypoxia-inducible factor-1 (HIF-1), which is a
	heterodimer composed of an alpha and a beta subunit. HIF-1 functions as a master regulator of cellular
	and systemic homeostatic response to hypoxia by activating transcription of many genes, including those
	involved in energy metabolism, angiogenesis, apoptosis, and other genes whose protein products
	increase oxygen delivery or facilitate metabolic adaptation to hypoxia. HIF-1 thus plays an essential role in
	embryonic vascularization, tumor angiogenesis and pathophysiology of ischemic disease. Alternatively
	spliced transcript variants encoding different isoforms have been identified for this gene. [provided by
	RefSeq, Jul 2011]
Function	Functions as a master transcriptional regulator of the adaptive response to hypoxia. Under hypoxic
	conditions, activates the transcription of over 40 genes, including erythropoietin, glucose transporters, glycolytic enzymes, vascular endothelial growth factor, HILPDA, and other genes whose protein products
	increase oxygen delivery or facilitate metabolic adaptation to hypoxia. Plays an essential role in
	embryonic vascularization, tumor angiogenesis and pathophysiology of ischemic disease. Binds to core
	DNA sequence 5'-[AG]CGTG-3' within the hypoxia response element (HRE) of target gene promoters.
	Activation requires recruitment of transcriptional coactivators such as CREBPB and EP300. Activity is
	enhanced by interaction with both, NCOA1 or NCOA2. Interaction with redox regulatory protein APEX
	seems to activate CTAD and potentiates activation by NCOA1 and CREBBP. Involved in the axonal
	distribution and transport of mitochondria in neurons during hypoxia. [UniProt]
Calculated Mw PTM	93 kDa In normoxia, is hydroxylated on Pro-402 and Pro-564 in the oxygen-dependent degradation domain (ODD)
F I IVI	by EGLN1/PHD2 and EGLN2/PHD1. EGLN3/PHD3 has also been shown to hydroxylate Pro-564. The
	hydroxylated prolines promote interaction with VHL, initiating rapid ubiquitination and subsequent
	proteasomal degradation. Deubiquitinated by USP20. Under hypoxia, proline hydroxylation is impaired
	and ubiquitination is attenuated, resulting in stabilization.
	In normoxia, is hydroxylated on Asn-803 by HIF1AN, thus abrogating interaction with CREBBP and EP300
	and preventing transcriptional activation. This hydroxylation is inhibited by the Cu/Zn-chelator, Clioquinol.
	S-nitrosylation of Cys-800 may be responsible for increased recruitment of p300 coactivator necessary for
	transcriptional activity of HIF-1 complex. Requires phosphorylation for DNA-binding. Phosphorylation at Ser-247 by CSNK1D/CK1 represses kinase
	activity and impairs ARNT binding. Phosphorylation by GSK3-beta and PLK3 promote degradation by the
	proteasome.
	Sumoylated; with SUMO1 under hypoxia. Sumoylation is enhanced through interaction with RWDD3.
	Both sumoylation and desumoylation seem to be involved in the regulation of its stability during hypoxia.
	Sumoylation can promote either its stabilization or its VHL-dependent degradation by promoting
	hydroxyproline-independent HIF1A-VHL complex binding, thus leading to HIF1A ubiquitination and
	proteasomal degradation. Desumoylation by SENP1 increases its stability amd transcriptional activity.
	There is a disaccord between various publications on the effect of sumoylation and desumoylation on its
	stability and transcriptional activity. Acetylation of Lys-532 by ARD1 increases interaction with VHL and stimulates subsequent proteasomal
	degradation (PubMed:12464182). Deacetylation of Lys-709 by SIRT2 increases its interaction with and
	hydroxylation by EGLN1 thereby inactivating HIF1A activity by inducing its proteasomal degradation
	(PubMed:24681946).
	Polyubiquitinated; in normoxia, following hydroxylation and interaction with VHL. Lys-532 appears to be
	the principal site of ubiquitination. Clioquinol, the Cu/Zn-chelator, inhibits ubiquitination through
	preventing hydroxylation at Asn-803. Ubiquitinated by a CUL2-based E3 ligase.
	The iron and 2-oxoglutarate dependent 3-hydroxylation of asparagine is (S) stereospecific within HIF
	CTAD domains.



ARG55734 anti-HIF-1 alpha antibody WB image

Western blot: 100 μg of HT1080 cell lysates in the absense/presence of CoCl2 stained with ARG55734 anti-HIF-1 alpha antibody at 1:2500 dilution.