

Datasheet for ABIN7317483 HDAC4 Protein



Overview

Quantity:	100 µg
Target:	HDAC4
Origin:	Human
Source:	Baculovirus infected Insect Cells
Protein Type:	Recombinant
Product Details	
Purpose:	Recombinant Human HDAC4 Protein (aa 612-1084)
Sequence:	Met612-Leu1084

Characteristics: A DNA sequence encoding the human HDAC4 (Met612-Leu1084) was expressed and purified

with two additional amino acids (Gly & Pro) at	at the N-terminus.
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> 90 % as determined by reducing SDS-PAGE.

Endotoxin Level:	< 1.0 EU per µg as determined by the LAL method

Target Details

Purity:

Target:	HDAC4
Alternative Name:	HDAC4 (HDAC4 Products)
Background:	Background: HDAC4 (histone deacetylase 4), belongs to class II of the histone
	deacetylase/acuc/apha family. Histone Deacetylases (HDACs) are a group of enzymes closely
	related to sirtuins. They catalyze the removal of acetyl groups from lysine residues in histones
	and non-histone proteins, resulting in transcriptional repression. In general, they do not act

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	autonomously but as components of large multiprotein complexes, such as pRb-E2F and
	mSin3A, that mediate important transcription regulatory pathways. There are three classes of
	HDACs, classes 1, 2 and 4, which are closely related Zn2+-dependent enzymes. HDACs are
	ubiquitously expressed and they can exist in the nucleus or cytosol. Their subcellular
	localization is effected by protein-protein interactions and by the class to which they belong.
	HDACs have a role in cell growth arrest, differentiation and death and this has led to substantial
	interest in HDAC inhibitors as possible antineoplastic agents. HDAC4 possesses histone
	deacetylase activity and represses transcription when tethered to a promoter. It does not bind
	DNA directly, but through transcription factors MEF2C and MEF2D. HDAC4 seems to interact in
	a multiprotein complex with RbAp48 and HDAC3.
	Synonym: AHO3,BDMR,HA6116,HD4,HDAC-4,HDAC-A,HDACA
Molecular Weight:	50.9 kDa
Molecular Weight: NCBI Accession:	50.9 kDa NP_006028
Molecular Weight: NCBI Accession: Pathways:	50.9 kDa NP_006028 Regulation of Muscle Cell Differentiation, Skeletal Muscle Fiber Development, Regulation of
Molecular Weight: NCBI Accession: Pathways:	50.9 kDa NP_006028 Regulation of Muscle Cell Differentiation, Skeletal Muscle Fiber Development, Regulation of Carbohydrate Metabolic Process
Molecular Weight: NCBI Accession: Pathways:	50.9 kDa NP_006028 Regulation of Muscle Cell Differentiation, Skeletal Muscle Fiber Development, Regulation of Carbohydrate Metabolic Process
Molecular Weight: NCBI Accession: Pathways: Application Details	50.9 kDa NP_006028 Regulation of Muscle Cell Differentiation, Skeletal Muscle Fiber Development, Regulation of Carbohydrate Metabolic Process
Molecular Weight: NCBI Accession: Pathways: Application Details Restrictions:	50.9 kDa NP_006028 Regulation of Muscle Cell Differentiation, Skeletal Muscle Fiber Development, Regulation of Carbohydrate Metabolic Process For Research Use only
Molecular Weight: NCBI Accession: Pathways: Application Details Restrictions:	50.9 kDa NP_006028 Regulation of Muscle Cell Differentiation, Skeletal Muscle Fiber Development, Regulation of Carbohydrate Metabolic Process For Research Use only

Format:	Lyophilized
Reconstitution:	Please refer to the printed manual for detailed information.
Buffer:	Lyophilized from sterile 20 mM Tris,500 mM NaCl, pH 7.4, 10 % gly
Storage:	4 °C,-20 °C,-80 °C
Storage Comment:	Generally, lyophilized proteins are stable for up to 12 months when stored at -20 to -80°C. Reconstituted protein solution can be stored at 4-8°C for 2-7 days. Aliquots of reconstituted
	samples are stable at < -20°C for 3 months.