USP27X [DAC-tagged]

Deconjugating enzyme: Deubiquitylase

Alternate Names: USP22L, X-linked ubiquitin carboxyl-terminal hydrolase 27, Ubiquitin thioesterase 27, Ubiquitin-specific-processing protease 27

Cat. No. 64-0046-050 Quantity: 50 µg Lot. No. 30156 Storage: -70°C

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Background

Deconjugating enzymes (DCEs) are proteases that process ubiquitin or ubiquitinlike gene products, reverse the modification of proteins by a single ubiquitin or ubiquitin-like protein (UBL) and remodel polyubiquitin (or poly-UBL) chains on target proteins (Reyes-Turcu et al., 2009). The deubiquitylating - or deubiquitinating - enzymes (DUBs) represent the largest family of DCEs and regulate ubiquitin dependent signalling pathways. The activities of the DUBs include the generation of free ubiquitin from precursor molecules, the recycling of ubiquitin following substrate degradation to maintain cellular ubiquitin homeostasis and the removal of ubiquitin or ubiquitin-like proteins (UBL) modifications through chain editing to rescue proteins from proteasomal degradation or to influence cell signalling events (Komander et al., 2009). There are two main classes of DUB, cysteine proteases and metalloproteases. Ubiquitin specific protease 27X (USP27X) is a member of the cysteine protease enzyme family and cloning of the gene was first described by Puente et al. (2003). Although closely related to USP22, which deubiquitylates histones, USP27X lacks the N-terminal ubiquitin binding protein zinc finger domain, suggesting it does not have the ability to deubiquitylate histones (Puente et al., 2003).

References:

Komander D, Clague MJ and Urbe S (2009) Breaking the chains: structure and function of the deubiquitinases. Nat Rev Mol Cell Biol 10, 550-563.

Lee DW, Peggie M, Deak M, Toth R, Gage ZO, Wood N, et al. (2012) The Dac-tag, an affinity tag based on penicillin-binding protein 5. Anal Biochem 428, 64-72.

Puente XS, Sanchez LM, Overall CM and Lopez-Otin C (2003) Human and mouse proteases: a comparative genomic approach. Nature reviews. *Genetics* **4**, 544-558.

Reves-Turcu FE, Ventii KH and Wilkinson KD (2009) Regulation and cellular roles of ubiquitin-specific deubiquitinating enzymes. Ann Rev Biochem 78, 363-397.

Physical Characteristics

Species: human Protein Sequence: Please see page 2

Source: E. coli Quantity: 50 µg

Concentration: 0.5 mg/ml

Formulation: 50 mM HEPES pH 7.5,

150 mM sodium chloride, 2 mM dithiothreitol, 10% glycerol

Molecular Weight: ~79.3 kDa

Purity: >67% by InstantBlue™ SDS-PAGE

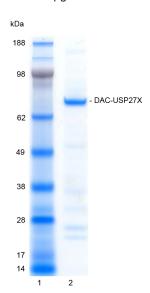
Stability/Storage: 12 months at -70°C;

aliquot as required

Quality Assurance

Purity:

4-12% gradient SDS-PAGE InstantBlue™ staining Lane 1: MW markers Lane 2: 1 µg DAC-USP27X



Protein Identification:

Confirmed by mass spectrometry.

Deubiquitylase Enzyme Assay:

The activity of DAC-USP27X was validated by the monitoring of mono-ubiquitin generation as a result of the enzyme catalysed cleavage of K63-linked di-ubiquitin. Incubation of the substrate in the presence or absence of DAC-USP27X was compared confirming the deubiquitylating activity of DAC-USP27X.



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Lot-specific COA version tracker: v1.0.0

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CERTIFICATE OF ANALYSIS Page 2 of 2

Physical Characteristics

Continued from page 1

MSAIPGVPQIDAESYILIDYNSGKV LAEQNADVRRDPASLTKMMTSYVIGQAMK **AGKFKETDLVTIGNDAWATGNPVFKGSSLM** FLKPGMQVPVSQLIRGINLQSGNDACVAMA **DFAAGSQDAFVGLMNSYVNALGLKNTHFQT** VHGLDADGQYSSARDMALIGQALIRDVPNEY SIYKEKEFTFNGIRQLNRNGLLWDNSLN **VDGIKTGHTDKAGYNLVASATEGOMRL ISAVMGGRTFKGREAESKKLLTWGFRFFET** VNPENLYFQGGSMCKDYVYDKDIEQIAKE EQGEALKLQASTSTEVSHQQCSVPGLGEK FPTWETTKPELELLGHNPRRRRITSSFTIGL RGLINLGNTCFMNCIVQALTHTPILRDF FLSDRHRCEMPSPELCLVCEMSSLFRELYS GNPSPHVPYKLLHLVWIHARHLAGYRQQDA HEFLIAALDVLHRHCKGDDVGKAANNPN HCNCIIDQIFTGGLQSDVTCQACHGVST TIDPCWDISLDLPGSCTSFWPMSPGRESS VNGESHIPGITTLTDCLRRFTRPEHLGS SAKIKCGSCQSYQESTKQLTMNKLPVVACF HFKRFEHSAKQRRKITTYISFPLELDMTP FMASSKESRMNGQLQLPTNSGNNENKYS LFAVVNHQGTLESGHYTSFIRHHKDQWFKCD DAVITKASIKDVLDSEGYLLFYHKQVLE HESEKVKEMNTQAY

Tag (**bold text**): N-terminal DAC (Lee *et al.* (2012))
Protease cleavage site: TEV™ (ENLYFQ▼G)
USP27X (regular text): Start *bold italics* (amino acid residues 1-438)

Accession number: NP_001138545



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