His₈- Avi-DDB1-biotinylated

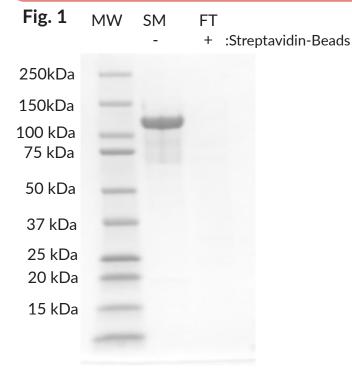
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DDB1 - DNA damage-binding protein 1, which is both involved in DNA repair and protein ubiquitination, as part of the UV-DDB complex and DCX (DDB1-CUL4-X-box) complexes, respectively. Core component of the UV-DDB complex (UVdamaged DNA-binding protein complex), a complex that recognizes UV-induced DNA damage and recruit proteins of the nucleotide excision repair pathway (the NER pathway) to initiate DNA repair. The UV-DDB complex preferentially binds to cyclobutane pyrimidine dimers (CPD), 6-4 photoproducts (6-4 PP), apurinic sites and short mismatches. Functions as a component of numerous distinct DCX (DDB1-CUL4-X-box) E3 ubiquitin-protein ligase complexes which mediate the ubiquitination and subsequent proteasomal degradation of target proteins. The functional specificity of the DCX E3 ubiquitin-protein ligase complex is determined by the variable substrate recognition component recruited by DDB1. The protein has an N-terminal 8x-His Tag followed by an Avi-Tag that was endogenous biotinylated during expression in Sf21 Cells.

Quality Control and Performance Data



Product Information

Quantity: 50 µg Molecular Weight: 130 kDa

Concentration: 17.7 µM, 2.3 mg/mL

Purity: >98% by SDS-PAGE

Biotinylation: >99% by Streptavidin-Bead pull down

Storage Buffer: 25 mM HEPES pH 7.5, 150 mM NaCl, 0.1

mM TCEP (no detergents or glycerol)

Storage: -80C, Avoid multiple freeze / thaw

Quality Control and Performance Data

Fig. 2	MW	1.0 μg	2.0 μg
250kDa			
150kDa	-		
100 kDa 75 kDa	=		-
50 kDa	-		
37 kDa	-		
25 kDa 20 kDa	=		
15 kDa			

Fig1. 6 μ g of His-Avi-DDB1 was incubated for 10 min w/ and w/o 20 μ L of packed Neutravidin Beads. Post incubation flow-through (FT) of unbound material was collected and analyzed side by side with starting material (SM). >99% of His-Avi-DDB1 is biotinylated.

Fig2. His-Avi-DDB1 SDS-PAGE. From left to right, increasing amounts of protein loaded onto a 10-20% SDS-PAGE gel, stained with Coomassie brillant blue. Purity is > 97%.

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