K11 linked Di-Ubiquitin

Cat. No.	SBB-UP0063
Lot. No.	181940063

K11 Di-Ubiquitin

The array of cellular processes initiated and regulated by ubiquitin has been partially explained by the structural diversity of differently linked ubiquitin chains. In a ubiquitin chain, ubiquitin moieties can be conjugated through one of their lysine residues (K6, K11, K27, K29, K33, K48 and K63) or the N-terminal methionine residue (M1), offering countless possibilities to assemble a specific polymer. Ubiquitin molecules can also be modified by other post-translational modifications, including acetylation and phosphorylation, adding another layer of ubiquitin signal regulation and diversification.

The abundance of K11 linkages strongly increase when the metazoan anaphase-promoting complex APC/C is active during mitosis, and APC/C has been shown to assemble K11linked ubiquitin chains to drive proteasomal degradation and exit from mitosis. This K11-linked di-ubiquitin was enzymatically conjugated, and purified via liquid chromatography.



1) Dikic, I., Wakatsuki, S., & Walters, K. J. (2009). Ubiquitin-binding domains – from structures to functions. Nature Reviews Molecular Cell Biology, 10(10), 659-671. https://doi.org/10.1038/nrm2767

2) Akutsu, M., Dikic, I., & Bremm, A. (2016). Ubiquitin chain diversity at a glance. Journal of Cell Science, 129(5), 875-880. https://doi. org/10.1242/jcs.183954

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Product Information

Quantity: 25 µg Molecular Weight: 17 kDa

Concentration: 58 μ M, 1 mg/mL

Purity: >95% by SDS-PAGE

Storage Buffer: 50 mM HEPES pH 7.5

Storage: -80C, Avoid multiple freeze / thaw

Quality Control and Performance Data



K11-Linked Di-Ubiquitin SDS-PAGE. From left to right, increasing amounts of di-ubiquitin were loaded onto a 10-20% SDS-PAGE gel, stained with Coomassie brillant blue. Purity is > 95%.

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