

## Active Recombinant Human Baculoviral IAP Repeat-Containing 2, AVI-tagged

**Cat. No.** BIRC2-3104H    **Lot. No.** (See product label)

### SPECIFICATION

|                         |  |
|-------------------------|--|
| <b>Cat. No.</b>         | BIRC2-3104H  |
| <b>Product Overview</b> | Recombinant Human BIRC2 (Glu 144-Leu 356) with a AVI tag at C-terminus and additional two amino acids (Gly & Pro) at the N-terminus, was expressed in E.coli.  |
| <b>Description</b>      | <p>Baculoviral IAP repeat-containing protein 2, also known as Inhibitor of apoptosis protein 2, HIAP2, c-IAP1, RING finger protein 48, and BIRC2, is a member of the IAP family. The inhibitor of apoptosis (IAP) proteins are a family of anti-apoptotic regulators found in viruses and metazoans. The UBA (ubiquitin-associated) domain of IAPs is located between the BIR (baculovirus IAP repeat) domains and the CARD (caspase activation and recruitment domain) or the RING (really interesting new gene) domain of c-IAP1 and c-IAP2 or XIAP (X-linked IAP) respectively. c-IAP1 contains three BIR repeats, one CARD domain and one RING-type zinc finger. c-IAP1 and c-IAP2 are recruited to tumor necrosis factor receptor 1 (TNFR1) associated complexes where they can regulate receptor-mediated signaling. Both c-IAP1 and c-IAP2 have been implicated in TNFalpha-stimulated NF-kappaB activation. Treatment of cells with IAP antagonists leads to proteasomal degradation of c-IAP1 and c-IAP2. Deletion or mutation of the UBA domain decreases this degradation, probably by diminishing the interaction of the c-IAPs with the proteasome. Ubiquitin binding may be an important mechanism for rapid turnover of auto-ubiquitinated c-IAP1 and c-IAP2.</p> |
| <b>Formulation</b>      | Lyophilized from sterile 10mM Tris, 5% glycerol, 0.5mM EDTA, 5mM DTT, pH 7.5   |

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|                             |   |
|-----------------------------|---|
| <b>Purity</b>               | > 92 % as determined by SDS-PAGE  |
| <b>Endotoxin</b>            | < 1.0 EU per µg of the protein as determined by the LAL method.   |
| <b>Bio-activity</b>         | Measured by its ability to inhibit DEVD-AFC cleavage activity in cell extracts activated by addition of cytochrome c and dATP.  |
| <b>Stability</b>            | Samples are stable for up to twelve months from date of receipt at -70°C.   |
| <b>Predicted N terminal</b> | Gly   |
| <b>Molecular Mass</b>       | The recombinant human cIAP1 ( Glu 144-Leu 356 ) consists of 230 amino acids and has a calculated molecular mass of 26.5 kDa as estimated by SDS-PAGE under reducing conditions. |
| <b>Storage</b>              | Store it under sterile conditions at -70°C. It is recommended that the protein be aliquoted for optimal storage. Avoid repeated freeze-thaw cycles.                             |

## GENE INFORMATION


|                    |   |
|--------------------|---|
| <b>Gene Name</b>   | <a href="#">BIRC2 baculoviral IAP repeat-containing 2 [ Homo sapiens ]</a>  |
| <b>Synonyms</b>    | BIRC2; baculoviral IAP repeat-containing 2; IAP homolog B; IAP-2; NFR2-TRAF signalling complex protein; RING finger protein 48; TNFR2-TRAF-signaling complex protein 2; apoptosis inhibitor 1; baculoviral IAP repeat-containing protein 2; inhibitor of apoptosis protein 2; API1; MIHB; HIAP2; RNF48; cIAP1; Hiap-2; c-IAP1 |
| <b>Gene ID</b>     | <a href="#">329</a>   |
| <b>mRNA Refseq</b> | <a href="#">NM_001166</a>   |

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| <b>Protein Refseq</b>      | NP_001157   |
| <b>MIM</b>                 | 601712  |
| <b>UniProt ID</b>          | Q13490  |
| <b>Chromosome Location</b> | 11q22   |
| <b>Pathway</b>             | Apoptosis; Focal adhesion; NOD-like receptor signaling pathway; Pathways in cancer; Small cell lung cancer; Ubiquitin mediated proteolysis; Apoptosis |
| <b>Function</b>            | metal ion binding; protein N-terminus binding; protein binding; zinc ion binding  |

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