## 32-4842: Recombinant Human SH3-domain GRB2-like endophilin B1

Alternative Bif-1,CGI-61,dJ612B15.2,PPP1R70,Endophilin-B1,Bax-interacting factor 1,SH3 domain-containing GRB2Name : like protein B1,KIAA0491.

## Description

Source : E.coli. SH3GLB1 Human Recombinant produced in E. coli is a single polypeptide chain containing 373 amino acids ( $1-365$ ) and having a molecular mass of 41.9 kDa . SH3GLB1 is fused to 8 amino acid His-tag at C-terminus \& purified by proprietary chromatographic techniques. Endophilin-B1 (SH3GLB1) is a part of the endophilin family and highly expresses in heart, skeletal muscle, kidney and placenta. Endophilin B1 belongs to the B subgroup of the endophilin family which is needed for preservation of mitochondrial morphology and for the regulation of the outer mitochondrial membrane dynamics. SH3GLB1 is required for normal outer mitochondrial membrane dynamics. Furthermore, SH3GLB1 is required for coatomer-mediated retrograde transport in certain cells. SH3GLB1 interacts with SH3GLB2 and Bcl-2-associated X protein and involved in regulating apoptotic signaling pathways.

## Product Info

| Amount : | $20 \mu \mathrm{~g}$ |
| :---: | :---: |
| Purification : | Greater than $90 \%$ as determined by SDS-PAGE. |
| Content : | The SH3GLB1 solution ( $1 \mathrm{mg} / 1 \mathrm{ml}$ ) contains 20 mM Tris- HCl buffer ( pH 8.0 ), $0.1 \mathrm{M} \mathrm{NaCl}, 10 \%$ glycerol and 2 mM DTT. |
| Storage condition : | Store at $4^{\circ} \mathrm{C}$ if entire vial will be used within $2-4$ weeks. Store, frozen at $-20^{\circ} \mathrm{C}$ for longer periods of time. For long term storage it is recommended to add a carrier protein ( $0.1 \%$ HSA or BSA).Avoid multiple freeze-thaw cycles. |
| Amino Acid : | MNIMDFNVKK LAADAGTFLS RAVQFTEEKL GQAEKTELDA HLENLLSKAE CTKIWTEKIM |
|  | KQTEVLLQPN PNARIEEFVY EKLDRKAPSR INNPELLGQY MIDAGTEFGP GTAYGNALIK |
|  | CGETQKRIGT ADRELIQTSA LNFLTPLRNF IEGDYKTIAK ERKLLQNKRL DLDAAKTRLK |
|  | KAKAAETRNS SEQELRITQS EFDRQAEITR LLLEGISSTH AHHLRCLNDF VEAQMTYYAQ |
|  | CYQYMLDLQK QLGSFPSNYL SNNNQTSVTP VPSVLPNAIG SSAMASTSGL VITSPSNLSD |
|  | LKECSGSRKA RVLYDYDAAN STELSLLADE VITVFSVVGM DSDWLMGERG NQKGKVPITY |



