## 32-2101: ACAD8 Recombinant Protein

# Alternative Name Acyl-CoA dehydrogenase family member 8 mitochondrial,ACAD-8,Isobutyryl-CoA 

: dehydrogenase,Activator-recruited cofactor 42 kDa component,ARC42,FLJ22590.

## Description

Source : Escherichia Coli. ACAD8 Human Recombinant produced in E.coli is a single, non-glycosylated polypeptide chain containing 416 amino acids (23-415) and having a molecular mass of 45.1 kDa .ACAD8 is fused to a 23 amino acid His-tag at N terminus \& purified by proprietary chromatographic techniques. Acyl CoA dehydrogenase is the enzymeused to catalyzethe first step of ?-oxidationin Fatty acid metabolism.Acyl-coenzyme A (CoA) dehydrogenases (ACADs) are a family of mitochondrial enzymes that catalyze the first dehydrogenation step in the bets-oxidation of fatty acyl-CoA derivatives. Several human ACADs exist and all ACADs catalyze the same initial dehydrogenation of the substrate at the beta-carbon atom and require electron transfer flavoprotein as an alectron acceptor. The predicted 415-amino acid ACAD8 protein contains many of the residues conserved in most other ACADs, including an active site glutamic acid residue and residues important for tetramer formation.

## Product Info

| Amount : | $20 \mu \mathrm{~g}$ |
| :---: | :---: |
| Purification : | Greater than 95\% as determined by SDS-PAGE. |
| Content : | The ACAD8 solution contains 20 mM Tris- HCl buffer ( pH 8.0 ), $0.15 \mathrm{M} \mathrm{NaCl}, 1 \mathrm{mM}$ DTT and $30 \%$ glycerol. |
| 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 : | MGSSHHHHHH SSGLVPRGSH MGSLVQTGHR SLTSCIDPSM GLNEEQKEFQ KVAFDFAARE |
|  | MAPNMAEWDQ KELFPVDVMR KAAQLGFGGV YIQTDVGGSG LSRLDTSVIF EALATGCTST |
|  | TAYISIHNMC AWMIDSFGNE EQRHKFCPPL CTMEKFASYC LTEPGSGSDA ASLLTSAKKQ |
|  | GDHYILNGSK AFISGAGESD IYVVMCRTGG PGPKGISCIV VEKGTPGLSF GKKEKKVGWN |
|  | SQPTRAVIFE DCAVPVANRI GSEGQGFLIA VRGLNGGRIN IASCSLGAAH ASVILTRDHL |
|  | NVRKQFGEPL ASNQYLQFTL ADMATRLVAA RLMVRNAAVA LQEERKDAVA LCSMAKLFAT |
|  | DECFAICNQA LQMHGGYGYL KDYAVQQYVR DSRVHQILEG SNEVMRILIS RSLLQE. |



