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## dnaK Recombinant DNAK ATPase binding domain (aa 1-384)

CSI13249       50 µg         Alternate Names:       HSP-70, HSP70, DnaK, Chaperone protein dnaK, Heat shock protein 70, Heat shock 70 kDa protein, groP, grpF, seg, b0014, JW0013.         Description:       DnaK, originally identified for its DNA replication by bacteriophage I in <i>E. coli</i> is the bacterial HSP-70 chaperone. This protein is involved in the folding and assembly of newly synthesized polypeptide chains and in preventing the aggregation of stress-denatured proteins.         DnaK (amino acids 1-384) is N-terminal ATPase domain and ATP bound to the ATPase domain induces a conformational change in the substrate binding domain (residues 385 -638). The protein coding region of the ATPase domain of DNAK (amino acids 1-384) was amplified by PCR and cloned into an <i>E. coli</i> expression vector. The ATPase domain of DNAK was purified to apparent homogeneity by using conventional column chromatography techniques.         Recombinant DnaK Substrate Binding Domain produced in <i>E. Coli</i> is a single, non-glycosylated polypeptide chain containing 384 amino acids.         Physical Appearance:       Sterile filtered colorless solution.         Gene ID:       944750         Source: <i>E. coli</i> Molecular Mass:       48.1 kDa         Formulation:       The DnaK protein contains 25 mM Tris-HCl, pH 7.5 + 100 mM NaCl + 5 mM DTT and 10% Giycerol.         Purity:       Greater than 95.0% as determined by <ul> <li>(a) Analysis by SDS-PAGE.</li> <li>Amino Acid Sequence:</li> <li>MGKIIGIDLG TTNSCVAIMD GTTPRVLENA EGDRTTPSII AYTQDGETLV</li> <li>GOPAKROAVT NPQNTLFAIK RLIGRREQDE EVQRPOVSIMP FKIIAADNGD</li> <li>AWVEVKGQK</li></ul>	Catalog No.	CSI13248	<b>Quantity:</b> 10 μg	
CS113250       1.0 mg         Alternate Names:       HSP-70, HSP70, DnaK, Chaperone protein dnaK, Heat shock protein 70, Heat shock 70 KDa protein, groP, grpF, seg, b0014, JW0013.         Description:       DnaK, originally identified for its DNA replication by bacteriophage 1 in <i>E. coli</i> is the bacterial HSP-70 chaperone. This protein is involved in the folding and assembly of newly synthesized polypeptide chains and in preventing the aggregation of stress-denatured proteins.         DnaK (amino acids1-384) is N-terminal ATPase domain and ATP bound to the ATPase domain induces a conformational change in the substrate binding domain (residues 385 -638). The protein coding region of the ATPase domain of DNAK (amino acids 1-384) was amplified by PCR and cloned into an <i>E. coli</i> expression vector. The ATPase domain of DNAK was purified to apparent homogeneity by using conventional column chromatography techniques.         Recombinant DnaK Substrate Binding Domain produced in <i>E. Coli</i> is a single, non-glycosylated polypeptide chain containing 384 amino acids.         Physical Appearance:       Sterile filtered colorless solution.         Gene ID:       944750         Source: <i>E. coli</i> Molecular Mass:       48.1 kDa         Formulation:       The DnaK protein contains 25 mM Tris-HCl, pH 7.5 + 100 mM NaCl + 5 mM DTT and 10%Glycerol.         Purity:       Greater than 95.0% as determined by (a) Analysis by SDS-PAGE.         Amino Acid Sequence:       MGKIIGIDLG TTNSCVAIMD GTTR-VLENA EGDRTTPSII AYTODGETLV GAPKRQAVT NPQNTLFAIK RLIGRRFQDE EVQRDVSIMP FKIIAADNGD AVVEVKGQKM APPOLSAEVL KKMKKTAEDY LGEPVTEAVI TV		CSI13249	50 µg	
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Gene ID:       944 / 50         Source:       E. coli         Molecular Mass:       48.1 kDa         Formulation:       The DnaK protein contains 25 mM Tris-HCl, pH 7.5 + 100 mM NaCl + 5 mM DTT and 10%Glycerol.         Purity:       Greater than 95.0% as determined by (a) Analysis by RP-HPLC. (b) Analysis by SDS-PAGE.         Amino Acid Sequence:       McKHIGIDLG TTNSCVAIMD GTTPRVLENA EGDRTTPSII AYTQDGETLV GQPAKRQAVT NPQNTLFAIK RLIGRRFQDE EVQRDVSIMP FKIIAADNGD AWVEVKGQKM APPQISAEVL KKMKKTAEDY LGEPVTEAVI TVPAYFNDAQ RQATKDAGRI AGLEVKRIIN EPTAAALAYG LDKGTGNRTI AVYDLGGGTF DISIIEIDEV DGEKTFEVLA TNGDTHLGGE DFDSRLINYL VEEFKKDQGI DLRNDPLAMQ RLKEAAEKAK IELSSAQQTD VNLPYITADA TGPKHMNIKV TRAKLESLVE DLVNRSIEPL KVALQDAGLS VSDIDDVILV GGQTRMPMVQ KKVAEFFGKE PRKDVNPDEA VAIGAAVQGG VLTG.         Storage & Stability:       Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°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.	Physical Appearance:	Sterile Intered coloriess soluti	DII.	
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NOT FOR HUMAN USE FOR RESEARCH ONLY NOT FOR DIAGNOSTIC OR THERAPEUTIC USE	Storage & Stability:	Store at 4°C if entire vial will the Store, frozen at -20°C for long For long term storage it is recovered and the storage it is recovered.	be used within 2-4 weeks. ger periods of time. ommended to add a carrier protein (0.1% HSA or BSA).	

