

32-4714: Recombinant Human Related RAS Viral (r-ras) Oncogene Homolog 2

Alternative Name : Ras-related protein R-Ras2,Ras-like protein TC21,Teratocarcinoma oncogene,RRAS2,TC21.

Description

Source : Escherichia Coli. RRAS2 Human Recombinant fused with a 20 amino acid His tag at N-terminus produced in E.Coli is a single, non-glycosylated, polypeptide chain containing 221 amino acids (1-201 a.a.) and having a molecular mass of 25.2 kDa.The RRAS2 is purified by proprietary chromatographic techniques. RRAS2 (RAS viral (r-ras) oncogene homolog 2) is a member of the Ras like GTPase family that cycle between GTP-bound active and GDP-bound inactive states at the cytoplasmic face of the plasma membrane. RRAS2 is a plasma membrane-associated GTP-binding protein with GTPase activity. RRAS2 is implicated in the pathogenesis of human cancers. Hence, overexpression of active RRAS2 in EpH4 cells induces tumorigenicity through the phosphoinositide 3-kinase, p38 MAPK, and mTOR pathways, as these cells lose their sensitivity to the normal growth inhibitory role of TGF-beta. RRAS2 might transduce growth inhibitory signals across the cell membrane, exerting its effect via an effector shared with the Ras proteins but in an antagonistic manner.RRAS2 is ubiquitously present in all tissues examined, with the highest levels in the heart, placenta, and skeletal muscle. Moderate levels of RRAS2 are found in the lung and liver and low levels in the brain, kidney, and pancreas.RRAS2 deficiencies are linked to ovarian cancer.

Product Info

Amount :	50 µg
Purification :	Greater than 95.0% as determined by SDS-PAGE.
Content :	The RRAS2 solution contains 20mM Tris-HCl buffer (pH8.0), 1mM DTT, 10% glycerol and 2mM EDTA.
Storage condition :	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.
Amino Acid :	MGSSHHHHHH SSGLVPRGSH MAAAGWRDGS GQEKYRLVVV GGGVVGKSAL TIQFIQSYFV TDYDPTIEDS YTKQCVIDDR AARLDILDTA GQEEFGAMRE QYMRTGEGFL LVFSVTDRGS FEEIYKFQRQ ILRVKDRDEF PMILIGNKAD LDHQRQVTQE EGQQLARQLK VTYMEASAKI RMNVDQAFHE LRVIRKQFQE QECPPSPEPT RKEKDKKGCH C.