

9853 Pacific Heights Blvd. Suite D. San Diego, CA 92121, USA Tel: 858-263-4982

Email: info@abeomics.com

32-3200: ANXA5 Recombinant Protein

Alternative Name:

PP4,ANX5,ENX2,ANXA5,Annexin A5,Annexin-5,Annexin V,Lipocortin V,Endonexin II,Calphobindin I,CBP-

 $I, Placental\ anticoagulant\ protein\ I, PAP-I, Placental\ anticoagulant\ protein\ 4, Thromboplastin$

inhibitor, Vascular anticoagulant-alpha, VAC-alph

Description

Source: Escherichia Coli. ANXA5 produced in E.Coli is a single, non-glycosylated polypeptide chain containing 320 amino acids (1-320 a.a.) and having a molecular mass of 35.9 kDa.ANXA5 is purified by proprietary chromatographic techniques. ANXA5 is a member of the annexin family of calcium-dependent phospholipid binding proteins which are involved in membrane-related activity along exocytotic and endocytotic pathways. ANXA5 is a phospholipase A2 and protein kinase C inhibitory protein with calcium channel properties and takes part in cellular signal transduction, inflammation, growth and differentiation. ANXA5 is an anticoagulant protein that acts as an indirect inhibitor of the thromboplastin-specific complex, which is involved in the blood coagulation cascade. ANXA5 regulates coagulability in the blood stream by binding to phosphatidylserine and sulfatide. ANXA5 protects sinsuoidal endothelial cells from ischemia reperfusion damage. ANXA5 is necessary for normal CFTR chloride channel activity.

Product Info

Amount : 20 μg

Purification: Greater than 95.0% as determined by SDS-PAGE.

Content: The ANXA5 protein solution contains 20mM Tris-HCl, pH-8, 1mM DTT and 10% glycerol.

Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods

Storage condition: 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: MAQVLRGTVT DFPGFDERAD AETLRKAMKG LGTDEESILT LLTSRSNAQR QEISAAFKTL FGRDLLDDLK

SELTGKFEKL IVALMKPSRL YDAYELKHAL KGAGTNEKVL TEIIASRTPE ELRAIKQVYE EEYGSSLEDD VVGDTSGYYQ RMLVVLLQAN RDPDAGIDEA QVEQDAQALF QAGELKWGTD EEKFITIFGT RSVSHLRKVF DKYMTISGFQ IEETIDRETS GNLEQLLLAV VKSIRSIPAY LAETLYYAMK GAGTDDHTLI RVMVSRSEID

LFNIRKEFRK NFATSLYSMI KGDTSGDYKK ALLLLCGEDD.

