

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

Email: info@abeomics.com

## 32-3196: ANXA2 Recombinant Protein

Alternative Name :

ANX2,ANX2L4,CAL1H,LIP2,LPC2,LPC2D,P36,PAP-IV,ANXA2,Annexin A2,Annexin-2,Annexin II,Lipocortin II,Calpactin-1 heavy chain,Calpactin I heavy chain,Chromobindin-8,p36,Protein I,Placental anticoagulant protein IV

## **Description**

Source: Escherichia Coli. ANXA2 Human Recombinant produced in E.Coli is a single, non-glycosylated, polypeptide chain containing 376 amino acids (1-339a.a.) and having a molecular mass of 42.8 kDa. ANXA2 is fused to a 37 amino acid His-Tag at N-Terminus and purified by proprietary chromatographic techniques. ANXA2 is part of the annexin family and is involved in the regulation of cellular growth and in signal transduction pathways. ANXA2 protein functions as an autocrine factor which increases osteoclast formation and bone resorption. ANXA2 is associated with sickle cell osteonecrosis. Reduced ANXA2 expression is associated with osteosarcoma metastases. ANXA2 is part of the putative cell surface vitamin D binding protein binding site complex and functions to mediate the chemotactic cofactor effect. ANXA2 is involved in dysferlin deficiency and in muscular dystrophies. Human colon adenocarcinoma cell differentiation is related with an up-regulation of ANXA2.

## **Product Info**

**Amount :** 20 μg

**Purification :** Greater than 95% as determined by SDS-PAGE.

**Content:** ANXA2 solution containing 20mM Tris pH-8, 0.1M NaCl 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: MRGSHHHHHH GMASMTGGOO MGRDLYDDDD KDRWGSHMST VHEILCKLSL EGDHSTPPSA

YGSVKAYTNF DAERDALNIE TAIKTKGVDE VTIVNILTNR SNAQRQDIAF AYQRRTKKEL ASALKSALSG HLETVILGLL KTPAQYDASE LKASMKGLGT DEDSLIEIIC SRTNQELQEI NRVYKEMYKT DLEKDIISDT SGDFRKLMVA LAKGRRAEDG SVIDYELIDQ DARDLYDAGV KRKGTDVPKW ISIMTERSVP HLQKVFDRYK SYSPYDMLES IRKEVKGDLE NAFLNLVQCI QNKPLYFADR LYDSMKGKGT RDKVLIRIMV SRSEVDMLKI

RSEFKRKYGK SLYYYIQQDT KGDYQKALLY LCGGDD.

