# SANTA CRUZ BIOTECHNOLOGY, INC.

# C1INH (B-11): sc-377062



## BACKGROUND

The serine proteinase inhibitors (serpins) comprise a superfamily of proteins with a diverse set of functions, including the control of complement activation, blood coagulation, programmed cell death and cell development. Serpins are secreted glycoproteins that contain a stretch of peptide that mimics a true substrate for a corresponding serine protease. The most abundant serpins in human plasma are  $\alpha$ -1-antitrypsin (AAT) and  $\alpha$ -1-antichymotrypsin (AACT). Other serpin family members include pigment epithelium-derived growth factor (PEDF), human protease nexin 1 (PN-1), protease inhibitor 6 (PI-6), thyroxine-binding globulin precursor (TBG), protease inhibitor 9 (PI-9), serine protease inhibitor 3 (Spi3), plasma protease C1 inhibitor (C1INH), Headpin, SerpinB12, monocyte/neutrophil elastase inhibitor members 1a, 1b and 1c (M/NEI) and squamous cell carcinoma antigens 1 and 2 (SCCA1/2). Antithrombin-III (ATIII) is a crucial serine protease inhibitor that regulates the coagulation cascade in blood and inhibits Thrombin.

# REFERENCES

- Curd, J.G., et al. 1981. Purification and characterization of two functionally distinct forms of C1 inhibitor from a patient with angioedema. Clin. Exp. Immunol. 145: 261-270.
- 2. Pixley, R.A., et al. 1985. The regulation of human Factor XIIa by plasma proteinase inhibitors. J. Biol. Chem. 260: 1723-1729.

## **CHROMOSOMAL LOCATION**

Genetic locus: SERPING1 (human) mapping to 11q12.1; Serping1 (mouse) mapping to 2 D.

#### SOURCE

C1INH (B-11) is a mouse monoclonal antibody specific for an epitope mapping between amino acids 215-249 within an internal region of C1INH of human origin.

# PRODUCT

Each vial contains 200  $\mu g\, lgG_1$  kappa light chain in 1.0 ml of PBS with < 0.1% sodium azide and 0.1% gelatin.

C1INH (B-11) is available conjugated to agarose (sc-377062 AC), 500 µg/ 0.25 ml agarose in 1 ml, for IP; to HRP (sc-377062 HRP), 200 µg/ml, for WB, IHC(P) and ELISA; to either phycoerythrin (sc-377062 PE), fluorescein (sc-377062 FITC), Alexa Fluor<sup>®</sup> 488 (sc-377062 AF488), Alexa Fluor<sup>®</sup> 546 (sc-377062 AF546), Alexa Fluor<sup>®</sup> 594 (sc-377062 AF594) or Alexa Fluor<sup>®</sup> 647 (sc-377062 AF647), 200 µg/ml, for WB (RGB), IF, IHC(P) and FCM; and to either Alexa Fluor<sup>®</sup> 680 (sc-377062 AF680) or Alexa Fluor<sup>®</sup> 790 (sc-377062 AF790), 200 µg/ml, for Near-Infrared (NIR) WB, IF and FCM.

Blocking peptide available for competition studies, sc-377062 P, (100 µg peptide in 0.5 ml PBS containing < 0.1% sodium azide and 0.2% stabilizer protein).

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#### **RESEARCH USE**

For research use only, not for use in diagnostic procedures.

## **APPLICATIONS**

C1INH (B-11) is recommended for detection of C1INH of mouse, rat and human origin by Western Blotting (starting dilution 1:100, dilution range 1:100-1:1000), immunoprecipitation [1-2  $\mu$ g per 100-500  $\mu$ g of total protein (1 ml of cell lysate)], immunofluorescence (starting dilution 1:50, dilution range 1:50-1:500), immunohistochemistry (including paraffin-embedded sections) (starting dilution 1:50, dilution range 1:50-1:500) and solid phase ELISA (starting dilution 1:30, dilution range 1:30-1:3000).

Suitable for use as control antibody for C1INH siRNA (h): sc-45608, C1INH siRNA (m): sc-45609, C1INH shRNA Plasmid (h): sc-45608-SH, C1INH shRNA Plasmid (m): sc-45609-SH, C1INH shRNA (h) Lentiviral Particles: sc-45608-V and C1INH shRNA (m) Lentiviral Particles: sc-45609-V.

Molecular Weight of C1INH: 55 kDa.

Molecular Weight of glycosylated C1INH: 75-105 kDa.

#### DATA





C1INH (B-11): sc-377062. Western blot analysis of C1INH expression in Hep G2 ( $\mathbf{A}$ ), HEK293 ( $\mathbf{B}$ ) and THP-1 ( $\mathbf{C}$ ) whole cell lysates and mouse liver ( $\mathbf{D}$ ) and rat liver ( $\mathbf{E}$ ) tissue extracts

C1INH (B-11): sc-377062. Immunoperoxidase staining of formalin fixed, paraffin-embedded human adrenal gland tissue showing cytoplasmic staining of glandular cells

## SELECT PRODUCT CITATIONS

- 1. Yang, J., et al. 2016. iTRAQ-based proteomics identification of serum biomarkers of two chronic hepatitis B subtypes diagnosed by traditional chinese medicine. Biomed Res. Int. 2016: 3290260.
- García-Hernández, V., et al. 2018. A tandem mass tag (TMT) proteomic analysis during the early phase of experimental pancreatitis reveals new insights in the disease pathogenesis. J. Proteomics 181: 190-200.
- Mueller, S.K., et al. 2019. Tissue and exosomal serine protease inhibitors are significantly overexpressed in chronic rhinosinusitis with nasal polyps. Am. J. Rhinol. Allergy 33: 359-368.
- 4. van Heukelum, S., et al. 2021. A central role for anterior cingulate cortex in the control of pathological aggression. Curr. Biol. 31: 2321-2333.e5.
- Kim, H.N., et al. 2021. The thrombin receptor modulates astroglia-neuron trophic coupling and neural repair after spinal cord injury. Glia 69: 2111-2132.
- Heukelum, S.V., et al. 2021. Structural degradation in midcingulate cortex is associated with pathological aggression in mice. Brain Sci. 11: 868.

#### **STORAGE**

Store at 4° C, \*\*DO NOT FREEZE\*\*. Stable for one year from the date of shipment. Non-hazardous. No MSDS required.