

P93076Hu01
Carbonic Anhydrase IX (CA9)
Organism: Homo sapiens (Human)
Instruction manual

FOR IN VITRO USE AND RESEARCH USE ONLY
NOT FOR USE IN DIAGNOSTIC OR THERAPEUTIC PROCEDURES

4th Edition (Revised in August, 2012)

[DESCRIPTION]

Protein Names: Carbonic Anhydrase IX

Synonyms: CA9, G250, MN

Species: Human

Size: 100µg

Source: *Escherichia coli*-derived

Subcellular Location: Nucleus. Cell membrane; Single-pass type I membrane protein.
Cell projection, microvillus membrane.

[PROPERTIES]

Residues: Pro59~Asp414 (Accession # Q8N6T7), with two N-terminal Tags, His-tag and T7-tag.

Grade & Purity: >95%, 43 kDa as determined by SDS-PAGE reducing conditions.

Formulation: Supplied as liquid form in Phosphate buffered saline(PBS), pH 7.4.

Endotoxin Level: <1.0 EU per 1µg (determined by the LAL method).

Applications: SDS-PAGE; WB; ELISA; IP.

(May be suitable for use in other assays to be determined by the end user.)

Predicted Molecular Mass: 43.3 kDa

Predicted isoelectric point: 4.8

[PREPARATION]

Reconstitute in sterile PBS, pH7.2-pH7.4.



[STORAGE AND STABILITY]

Storage: Avoid repeated freeze/thaw cycles.

Store at 2-8°C for one month.

Aliquot and store at -80°C for 12 months.

Stability Test: The thermal stability is described by the loss rate of the target protein. The loss rate was determined by accelerated thermal degradation test, that is, incubate the protein at 37°C for 48h, and no obvious degradation and precipitation were observed. (Referring from China Biological Products Standard, which was calculated by the Arrhenius equation.) The loss of this protein is less than 5% within the expiration date under appropriate storage condition.

[SEQUENCES]

The target protein is fused with two N-terminal Tags, His-tag and T7-tag, its sequence is listed below.

MGSSHHHHHHSSGLVPRGSHMASMTGGQQMGRGSEFELRRQ-PL GEEDLPSEED SPREEDPPGE EDLPGEEDLP
GEEDLPEVKP KSEEEGSLKL EDLPTVEAPG DPQEPQNNAH RDKEGDDQSH WRYGGDPPWP RVSPACAGRF QSPVDIRPQL
AAFCPALRPL ELLGFQLPPL PELRLRNNGH SVQLTLPPGL EMALGPGREY RALQLHLHWG AAGRPGSEHT VEGHRFP AEI
HVVHLSTAF RVD EALGRPG GLAVLA AFLE EGPEENSAYE QLLSRLEEIA EEGSETQVPG LDISALLPSD FSR YFQYEGS
LTPPCAQGV IWT VFNQTV M LSAKQLHTLS DTLWGP GDSR LQLNFRATQP L NGRVIEASF PAGVDSSPRA AEPVQLNSCL AAGD

