



#### P80761Hu01

# Aminoacyl tRNA Synthetase Complex Interacting Multifunctional Protein 1(AIMP1)

Organism: Homo sapiens (Human)

Instruction manual

FOR IN VITRO USE AND RESEARCH USE ONLY NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES

5th Edition (Revised in January, 2013)

### [ DESCRIPTION ]

**Protein Names:** Aminoacyl tRNA Synthetase Complex Interacting Multifunctional Protein 1

Synonyms: AIMP1, EMAP2, SCYE1

Species: Human

Size: 100µg

Source: Escherichia coli-derived

**Subcellular Location:** Nucleus. Cytoplasm, cytosol. Cytoplasmic vesicle, secretory vesicle. Secreted.

Endoplasmic reticulum. Golgi apparatus.

# [PROPERTIES]

Residues: Lys148~Lys312 (Accession # Q12904),

with N-terminal His-Tag.

Grade & Purity: >95%, 22kDa as determined by

SDS-PAGE reducing conditions.

Formulation: Supplied as lyophilized form in PBS, pH

7.4, containing 5% sucrose, 0.01% sarcosyl.

**Endotoxin Level:** <1.0 EU per  $1\mu g$  (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:** 19.6kDa

Predicted isoelectric point: 6.8

Human AIMP1

kDa
94
66.2
45
33
26
20

15% SDS-PAGE





## [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 N-terminal His-Tag, its sequence is listed below.

MGHHHHHHSGSEF- KPI DVSRLDLRIG CIITARKHPD ADSLYVEEVD VGEIAPRTVV

SGLVNHVPLE QMQNRMVILL CNLKPAKMRG VLSQAMVMCA SSPEKIEILA PPNGSVPGDR

ITFDAFPGEP DKELNPKKKI WEQIQPDLHT NDECVATYKG VPFEVKGKGV CRAQTMSNSG

IK