



# P91350Hu01 Adipose Differentiation Related Protein (ADRP) 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: Adipose Differentiation Related Human	ADRP
Protein	kDa
Synonyms: ADRP, PLIN2, ADFP	66.2
Species: Human	45
Size: 100µg	33
Source: Escherichia coli-derived	26
Subcellular Location: Membrane; Peripheral	20
membrane protein.	20
[ PROPERTIES ]	
Residues: Ala5~Tyr232 (Accession # Q99541), with	14.4
N-terminal His-Tag.	
Grade & Purity: >95%, 28kDa as determined by 15% SDS-	PAGE
SDS-PAGE reducing conditions.	
Formulation: Supplied as lyophilized form in PBS, pH	
7.4, containing 5% sucrose, 0.01% sarcosyl.	
<b>Endotoxin Level:</b> <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: 25.9kDa	
Predicted isoelectric point: 7.1	

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### [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- AVDPQP SVVTRVVNLP LVSSTYDLMS SAYLSTKDQY PYLKSVCEMA ENGVKTITSV AMTSALPIIQ KLEPQIAVAN TYACKGLDRI EERLPILNQP STQIVANAKG AVTGAKDAVT TTVTGAKDSV ASTITGVMDK TKGAVTGSVE KTKSVVSGSI NTVLGSRMMQ LVSSGVENAL TKSELLVEQY LPLTEEELEK EAKKVEGFDL VQKPSYYVRL GSLSTKLHSR AY

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