TEL:4006-871-227 Web:www.ybio.net Email:shybio@126.com

YB91880Mu01 100µg

Cyclin Dependent Kinase 7 (CDK7) Organism: Mus musculus (Mouse) Instruction manual

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

6th Edition (Revised in March, 2013)

## [ PROPERTIES ]

**Residues:** Tyr12<sup>~</sup>Phe295 (Accession # Q03147), 45 with N-terminal His-Tag. 33 Host: E. coli 26 Subcellular Location: Nucleus. Cytoplasm, 20 perinuclear region. 14.4 **Purity:** >95% **Endotoxin Level:** <1.0EU per 1µg (determined by the LAL method). Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% sucrose, 0.01% sarcosyl. Predicted isoelectric point: 6.3 Predicted Molecular Mass: 33.7kDa Applications: SDS-PAGE; WB; ELISA; IP. (May be suitable for use in other assays to be determined by the end user.)

[ USAGE ]



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Web:www.ybio.net Email:shybio@126.com Reconstitute in sterile PBS, pH7.2-pH7.4.



## [ <u>STORAGE AND STABILITY</u> ]

## Storage: Avoid repeated freeze/thaw cycles.

Store at  $2-8^{\circ}C$  for one month.

Aliquot and store at  $-80^{\circ}$ 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.

## [ <u>SEQUENCES</u> ]

The target protein is fused with N-terminal His-Tag, its sequence is listed below. MG H H H H H S G S E F - YE K L D F L G E GQ FAT V Y K A R DK N T N Q I VA I KK I K L G H R S E AKDGINRTAL REIKLLQELS HPNIIGLLDA FGHKSNISLV FDFMETDLEV IIKDNSLVLT PSHIKAYMLM TLQGLEYLHQ HWILHRDLKP NNLLLDENGV LKLADFGLAK SFGSPNRAYT HQVVTRWYRA PELLFGARMY GVGVDMWAVG CILAELLLRV PFLPGDSDLD QLTRIFETLG TPTEEQWPDM CSLPDYVTFK SFPGVPLQHI FIAAGDDLLE LIQGLFLFNP CTRTTASQAL KTKYF