RPC239Hu01 100µg **Recombinant Thyroid Hormone Receptor Alpha (THRa) Organism Species: Homo sapiens (Human)** Instruction manual

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

10th Edition (Revised in Jan, 2014)

kDa 70

44

33

26

22

18

14

10

### [PROPERTIES]

Residues: Met1~Pro490 **Tags:** Two N-terminal Tags, His-tag and T7-tag Accession: P10827 Host: E. coli Subcellular Location: Nucleus. **Purity:** >95% Endotoxin Level: <1.0EU per 1µg (determined by the LAL method). Formulation: Supplied as lyophilized form in PBS, pH7.4, containing 5% trehalose, 0.01% sarcosyl. Predicted isoelectric point: 6.4 Predicted Molecular Mass: 58.5kDa

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

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

# [USAGE]

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

# 15% SDS-PAGE

### [ 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.

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

The sequence of the target protein is listed below.

MEQKPSKVEC GSDPEENSAR SPDGKRKRKN GQCSLKTSMS GYIPSYLDKD EQCVVCGDKA TGYHYRCITC EGCKGFFRRT IQKNLHPTYS CKYDSCCVID KITRNQCQLC RFKKCIAVGM AMDLVLDDSK RVAKRKLIEQ NRERRRKEEM IRSLQQRPEP TPEEWDLIHI ATEAHRSTNA QGSHWKQRRK FLPDDIGQSP IVSMPDGDKV DLEAFSEFTK IITPAITRVV DFAKKLPMFS ELPCEDQIIL LKGCCMEIMS LRAAVRYDPE SDTLTLSGEM AVKREQLKNG GLGVVSDAIF ELGKSLSAFN LDDTEVALLQ AVLLMSTDRS GLLCVDKIEK SQEAYLLAFE HYVNHRKHNI PHFWPKLLMK EREVQSSILY KGAAAEGRPG GSLGVHPEGQ QLLGMHVVQG PQVRQLEQQL GEAGSLQGPV LQHQSPKSPQ QRLLELLHRS GILHARAVCG EDDSSEADSP SSSEEEPEVC EDLAGNAASP