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## P92175Hu02 <br> Left/Right Determination Factor 1 (LEFTY1) 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: Left/Right Determination Factor 1
Synonyms: LEFTY1, LEFTB, LEFTYB
Species: Human
Size: $100 \mu \mathrm{~g}$
Source: Escherichia coli-derived
Subcellular Location: Secreted. [ PROPERTIES]
Residues: Leu167~Glu332 (Accession \# O75610), 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 $1 \mu \mathrm{~g}$ (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: 19.9kDa
Predicted isoelectric point: 5.5
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## [ PREPARATION]

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

## [ STORAGE AND STABILITY ]

Storage: Avoid repeated freeze/thaw cycles.
Store at $2-8^{\circ} \mathrm{C}$ for one month.
Aliquot and store at $-80^{\circ} \mathrm{C}$ 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 $37^{\circ} \mathrm{C}$ for 48 h , 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- LVSV HESGWKAFDV TEAVNFWQQL SRPRQPLLLQ VSVQREHLGP

| LASGAHKLVR | FASQGAPAGL | GEPQLELHTL | DLGDYGAQGD | CDPEAPMTEG |
| :--- | :--- | :--- | :---: | :---: |
| TRCCRQEMYI | DLQGMKWAEN | WVLEPPGFLA | YECVGTCRQP | PEALAFKWPF | LGPRQCIASE TDSLPMIVSI KE

