# YB307Hu01 100 $\mu \mathrm{g}$ <br> Recombinant Cat Eye Syndrome Chromosome Region, Candidate 1 (CECR1) Organism Species: Homo sapiens (Human) <br> Instruction manual <br> FOR IN VITRO USE AND RESEARCH USE ONLY <br> NOT FOR USE IN CLINICAL DIAGNOSTIC PROCEDURES 

## [ PROPERTIES]

Residues: Ile30~Lys511

| Tags: Two N-terminal Tags, His-tag and T7-tag | $=\frac{k 0}{}$ |
| :--- | :--- |
| Accession: Q9NZK5 | -44 |
| Host: E. coli | -33 |
| Subcellular Location: Secreted. | 26 |
| Purity: $>95 \%$ | -22 |

$\begin{array}{ll}\text { Endotoxin Level: <1.0EU per 1 } \mu \mathrm{g} \text { (determined by the LAL } & \\ \begin{array}{ll}\text { method). } & 18 \\ \text { Formulation: Supplied as lyophilized form in PBS, pH7.4, } & 14 \\ \text { containing 5\% trehalose, } 0.01 \% \text { sarcosyl. } & 10 \\ \text { Predicted isoelectric point: 8.2 Predicted Molecular Mass: } & 15 \% \text { SDS-PAGE }\end{array}\end{array}$ 59.6kDa

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.

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## [ 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 sequence of the target protein is listed below.
I DETRAHLLLK EKMMRLGGRL VLNTKEELAN ERLMTLKIAE MKEAMRTLIF PPSMHFFQAK HLIERSQVFN ILRMMPKGAA LHLHDIGIVT MDWLVRNVTY RPHCHICFTP RGIMQFRFAH PTPRPSEKCS KWILLEDYRK RVQNVTEFDD SLLRNFTLVT QHPEVIYTNQ NVVWSKFETI FFTISGLIHY APVFRDYVFR SMQEFYEDNV LYMEIRARLL PVYELSGEHH DEEWSVKTYQ EVAQKFVETH PEFIGIKIIY SDHRSKDVAV IAESIRMAMG LRIKFPTVVA GFDLVGHEDT GHSLHDYKEA LMIPAKDGVK LPYFFHAGET DWQGTSIDRN ILDALMLNTT RIGHGFALSK HPAVRTYSWK KDIPIEVCPI SNQVLKLVSD LRNHPVATLM ATGHPMVISS DDPAMFGAKG LSYDFYEVFM GIGGMKADLR TLKQLAMNSI KYSTLLESEK NTFMEIWKKR WDKFIADVAT K

