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**YBG609Hu01 100μg**

**Recombinant Steroid Sulfatase Isozyme S (STS)**

**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)

## **[ PROPERTIES ]**

**Residues: His235~Arg583**

**Tags: Two N-terminal Tags, His-tag and T7-tag**

**Accession: P08842**

**Host: *E. coli***

**Subcellular Location: Endoplasmic reticulum membrane; Multi-pass membrane protein.**

**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: 6.0**

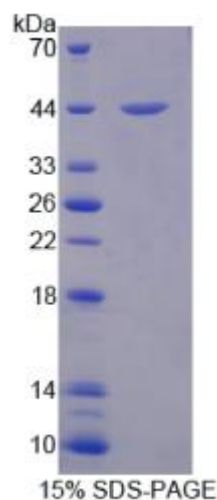
**Predicted Molecular Mass: 43.6kDa**

**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.**





## **[ 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 sequence of the target protein is listed below.**

**HYFRPL NCFMMRNYEI IQQPMSYDNL TQRLTVEAAQ FIQRNTETPF LLVLSYLHVH  
TALFSSKDFA GKSQHGVDYGD AVEEMDWSVG QILNLLDELRLANDTLIYFT SDQGAHVEEV  
SSKGEIHGGS NGIYKGGKAN NWEGGIRVPG ILRWPRVIQA GQKIDEPTSN MDIFPTVAKL  
AGAPLPEDRI IDGRDLMPLL EGKSQRSDEH FLFHYCNAYL NAVRWHPQNS TSIWKAFFFT  
PNFNPVGSNG CFATHVCFCF GSYVTHHDPPLLFDISKDPR ERNPLTPASE PRFYEILKVM  
QEADRHTQT LPEVPDQFSW NNFLWKPWLQ LCCPSTGLSC QCDREKQDKR LSR**