

**RPH108Hu01 50µg**

**Recombinant Sulfatase 1 (SULF1)**

**Organism Species: Homo sapiens (Human)**

***Instruction manual***

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

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12th Edition (Revised in Aug, 2016)

**[ PROPERTIES ]**

**Source:** Prokaryotic expression

**Host:** *E.coli*

**Residuess:** Pro609~Gly871

**Tags:** N-terminal His and GST Tag

**Tissue Specificity:** Golgi apparatus

**Purity:** > 95%

**Traits:** Freeze-dried powder

**Buffer formulation:** 20mM Tris, 150mM NaCl, pH8.0, containing 1mM EDTA, 1mM DTT, 0.01% SKL, 5% Trehalose and Proclin300.

**Original Concentration:** 200µg/mL

**Applications:** Positive Control; Immunogen; SDS-PAGE; WB.

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

**Predicted isoelectric point:** 8.3

**Predicted Molecular Mass:** 61.1kDa

**Accurate Molecular Mass:** 60kDa as determined by SDS-PAGE reducing conditions.

**[ USAGE ]**

Reconstitute in 20mM Tris, 150mM NaCl (pH8.0) to a concentration of 0.1-1.0 mg/mL. Do not vortex.

**[ 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. 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. The loss rate is less than 5% within the expiration date under appropriate storage condition.

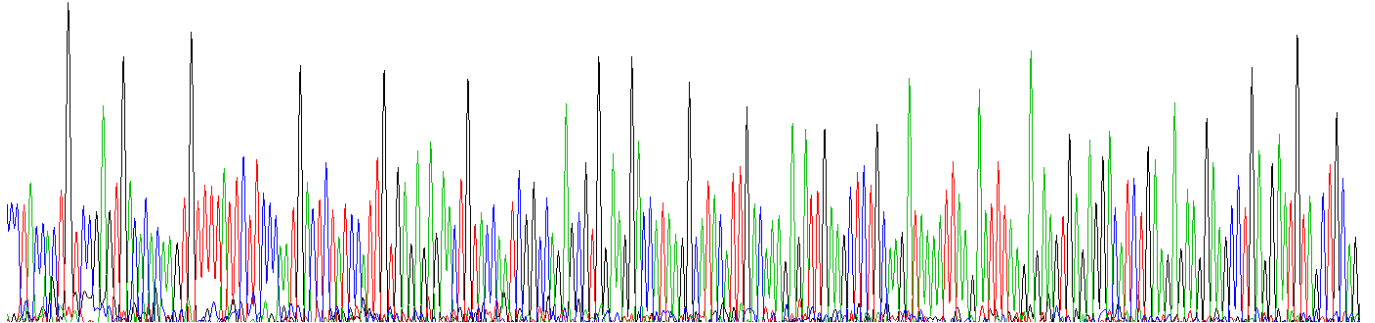
**[ SEQUENCE ]**

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PT TVRVTHKCFI LPNDSIH CER ELYQSARAWK DHKAYIDKEI
EALQDKIKNL REVRGHLKRR KPEECSCSKQ SYNKEKGVK KQEKLKSHLH
PFKEAAQEV D SKLQLFKENN RRRKKERKEK RRQRKGEECS LPGLTCFTHD
NNHWQTAPFW NLGSFCACTS SNNNTYWCLR TVNETHNFLF CEFATGFLEY
FDMNTDPYQL TNTVHTVERG ILNQLHVQLM ELRSCQGYKQ CNPRPKNL DV
GNKDGGSYDL HRGQLWDGWE G
    
```

**[ IDENTIFICATION ]**

CCTHCACTGTCCGGTGACGACAAGTGTTTATTCTCCATGACTCTATCATTTGTGAGAGACTGTACCAATCGGCAGAGCGTGGAGGACCATAAGGCATCATTTGCAAAGGATTTGAACCTCTCAAGATAAATTAAGAATTTAAGGAGGTGAGAGGCATCTGAAGAGAGGAGACCTGAGAGAGGACCTGAGGATGTAGCTGCAG  
PTTVRVTHKCFILPNDSIHCERELYQSARAWKDHKAYIDKEIEALQDKIKNLREVRGHLKRRKPEECSCS



**Figure. Gene Sequencing (Extract)**

