

## *Division of Signal Transduction Therapy*

### **Standard Operation Procedure**

#### **Preparation of His-JOSD2**

<b><u>Enzyme description:-</u></b>	His-JOSD2
<b><u>Clone number:-</u></b>	DU20941
<b><u>Source:-</u></b>	BL21 Recombinant
<b><u>Tag:-</u></b>	His
<b><u>Purification method:-</u></b>	N-terminal His <sub>6</sub> -tag
<b><u>Expression level:-</u></b>	2 mg/L

#### **Calculated molecular mass:-**

Monoisotopic	23186 Da
Average Mass	23199 Da
[cysteines reduced, methionines have not been oxidised]	

**Theoretical pI:-** 7.35

**Purity:-** 60%

#### **Enzyme storage buffer:-**

50 mM HEPES pH 7.5, 10% glycerol, 150mM NaCl, 1mM DTT

**Storage temperature:-** -80°C

#### **Assay:-**

Ub-Rho110-Gly cleavage assay monitored by Ex/Em 485/535 nm

#### **Assay buffer:-**

40 mM Tris pH 7.5, 100 mM NaCl, 5 mM DTT, 0.01% Triton X-100, 0.005% Ovalbumin, 0.5 μM Ub-Rho110-Gly

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**Clone Data Sheet**

**His-JOSD2**

<b><u>Protein</u></b>	His-JOSD2
<b><u>Synonyms</u></b>	Josephin-2
<b><u>Clone Number</u></b>	DU20941
<b><u>Species</u></b>	Human
<b><u>Accession Number</u></b>	Protein: Q8TAC2 DNA: NM_138334.2
<b><u>Tags</u></b>	N-terminal His <sub>6</sub> -tag
<b><u>Amino acid sequence of expressed protein</u></b>	<b>MGSSHHHHHSSGLEVL<u>FQGP</u>GSMSQAPGAQPSPTVYHERQRLELCAVH ALNNVL<u>QQLF</u>SQEAADEICKRLAPDSRLNPHRSLLGTYDNNVIMAAL QGLGLAAVWDRRRPLSQLALPQVLGLILNLPSVSLGLLSLPLRRRHV ALRQVDGVYYNLDSKLRAPEALGDEDGVRAFLAALAQGLCEVLLVVTKE VEEKGSWLRTD</b>
<b><u>Native sequence</u></b>	in bold
<b><u>Protease cleavage</u></b>	Precision site underlined
<b><u>Cloning sites</u></b>	BamH1 / Not1

**DNA sequence of  
insert**

GGATCCATGTCCCAGGCCCGGGAGCACAGCCGAGCCCACCCACCGTGTA  
CCACGAACGGCAGCGCCTGGAGCTGTGTGCTGTCCACGCCCTCAACAACG  
TTCTGCAGCAGCAGCTCTTTAGCCAGGAGGCTGCCGATGAGATCTGCAAG  
AGGTTGGCCCCAGACTCCCGGCTGAACCCTCATCGCAGCCTCCTGGGCAC  
CGGCAACTATGATGTCAATGTGATCATGGCCGCTCTGCAGGGGCTGGGCC  
TGGCCGCCGTGTGGTGGGACAGGAGGAGGCCCTGTCCCAGCTGGCCCTG  
CCCCAGGTACTGGGGCTGATCCTGAACCTGCCCTCGCCCGTGTGCTGGG  
GCTGCTGTCACTGCCGCTGCGCCGGCGGCAC TGGGTGGCCCTGCGCCAGG  
TGGACGGTGTCTACTACAACCTGGACTCCAAGCTGCGGGCGCCCGAGGCC  
CTGGGGGATGAGGACGGAGTCAGGGCCTTCTGGCGGCTGCGCTGGCCCA  
GGGCCTGTGCGAGGTGCTGCTGGTAGTGACCAAGGAGGTGGAGGAGAAGG  
GCAGCTGGCTGCGGACAGACTGAGCGGCCGC