

## *Division of Signal Transduction Therapy*

### **Standard Operation Procedure**

### **Preparation of GST-GABARAP**

<b><u>Enzyme description:-</u></b>	GABARAP
<b><u>Clone number:-</u></b>	DU40011
<b><u>Source:-</u></b>	BL21 recombinant
<b><u>Tag:-</u></b>	N-terminal GST
<b><u>Purification method:-</u></b>	GSH-Sepharose
<b><u>Expression level:-</u></b>	6mg/L
<b><u>Calculated molecular mass:-</u></b>	
Monoisotopic	41232 Da
Average Mass	41257 Da
[cysteines reduced, methionines have not been oxidised]	
<b><u>Theoretical pI:-</u></b>	6.82
<b><u>Purity:-</u></b>	90%
<b><u>Enzyme storage buffer:-</u></b>	
50 mM HEPES pH 7.5, 10% glycerol, 150mM NaCl, 1mM DTT	
<b><u>Storage temperature:-</u></b>	-80°C
<b><u>Assay:-</u></b>	

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**Clone Data Sheet**

**GST-GABARAP**

<b><u>Protein</u></b>	GST-GABARAP
<b><u>Synonyms</u></b>	Gamma-aminobutyric acid receptor associated protein; ATG8A; FLJ25768; MGC120154; MGC120155; MM46
<b><u>Clone Number</u></b>	DU40011
<b><u>Species</u></b>	Human
<b><u>Accession Number</u></b>	Protein: NP_009209; DNA: NM_007278.1
<b><u>Tags</u></b>	GST-
Aminoacid sequence of the expressed protein	MSPILGYWKIKGLVQPTRLLEYLEEKYEEHLYERDEGDKWRNKKFE LGLFNPYPYIDGDVKLTQSMAIIRYIADKHNMLGGCPKERAIEISM LEGAVLDIRYGVSR IAYSKDFETLKVDFLSKLPEMLKMFEDRLCHKT YLNGDHVTHPDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAIPO IDKYLKSSKYIAWPLQGWQATFGGGDHPPKSDHPPKSDENLYFQGG <b>MKFVYKEEHPFEKRRSEGEKIRKKYPDRVPVIVEKAPKARIGDLDDK</b> <b>KYLVPSDLTVGQFYFLIRKRIHLRAEDALFFFVNNVIPPTSATMGQL</b> <b>YQEHHEEDFFLYIAYSDESIVYGL</b>
Native sequence	
Protease cleavage	TEV
Cloning sites	BamH1 / NotI

**DNA sequence of  
insert**

GGATCCATGAAGTTCGTGTACAAAGAAGAGCATCCGTTGAGAAGCG  
CCGCTCTGAGGGCGAGAAGATCCGAAAGAAATACCCGGACCGGGTGC  
CGGTGATAGTAGAAAAGGCTCCCAAAGCTCGGATAGGAGACCTGGAC  
AAAAAGAAATACCTGGTGCCTTCTGATCTCACAGTTGGTCAGTTCTA  
CTTCTTGATCCGGAAGCGAATTCATCTCCGAGCTGAGGATGCCTTGT  
TTTTCTTTGTCAACAATGTCATTCCACCCACCAGTGCCACAATGGGT  
CAGCTGTACCAGGAACACCATGAAGAAGACTTCTTTCTCTACATTGC  
CTACAGTGACGAAAGTGTCTACGGTCTGTGAGCGGCCGC