

## *Division of Signal Transduction Therapy*

### **Standard Operating Procedure**

#### **Preparation of active RSK3 [2 - 733]**

|                                           |                  |
|-------------------------------------------|------------------|
| <b><u>Enzyme description:-</u></b>        | RSK3 [2 - 733]   |
| <b><u>Clone number:-</u></b>              | DU 902           |
| <b><u>Source:-</u></b>                    | Recombinant      |
| <b><u>Expression system:-</u></b>         | <i>E.coli</i>    |
| <b><u>Tag:-</u></b>                       | N-terminal GST   |
| <b><u>Purification method:-</u></b>       | GSH Sepharose    |
| <b><u>Expression level:-</u></b>          | 1.5 mg/L         |
| <b><u>Calculated molecular mass:-</u></b> | 110, 248 daltons |
| <b><u>Purity:-</u></b>                    | 70 %             |

#### **Activation protocol:-**

RSK3 (2  $\mu$ M) is activated by incubation with 5 U/ml GST-p42MAPKinase [DU 650 or DU 1844] and 3.3  $\mu$ g/ml GST-PDK1 [DU 954] in 50 mM Tris-HCl pH 7.5, 0.1mM EGTA, 0.1 % 2-mercaptoethanol, 10 mM magnesium acetate, 0.1 mM ATP for 30 min at 30 °C.

At present we are unable to remove the upstream activators from the RSK3.

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

50 mM Tris-HCl pH 7.5, 270 mM Sucrose, 150 mM NaCl, 0.1 mM EGTA, 0.1 % 2-mercaptoethanol, 0.02 % Brij-35, 0.2 mM PMSF, 1 mM Benzamidine

**Storage temperature:-** -70 °C

**Assay:-** Standard filter binding assay

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

50 mM Tris-HCl pH 7.5, 0.1% 2-mercaptoethanol, 0.1 mM EGTA, 10 mM MgAc

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### **Substrate:-**

Long S6 [KEAKEKRQEIQAKRRRLSSLRASTSKSGGSQK]

Final concentration: 30  $\mu$ M

**Specific activity range:-** to be determined

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## **Clone Data Sheet**

### **RSK3 [2 - 733]**

|                                             |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   |
|---------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b><u>Protein</u></b>                       | RSK3 [2 - 733]                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| <b><u>Clone number</u></b>                  | DU 902                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
| <b><u>Species</u></b>                       | Human                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             |
| <b><u>Accession number</u></b>              | X85106                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
| <b><u>Tags</u></b>                          | N-terminal GST                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| <b><u>Bacterially expressed protein</u></b> | <p>MSPILGYWKIKGLVQPTRLLEYLEEKYEEHLYERDEGDKWRNKKFEL<br/>GLEFPNLPYYIDGDVKLTQSMAIIRYIADKHNLGGCPKERAIEISMLE<br/>GAVLDIRYGVSRIAYSKDFETLKVDFLSKLPEMLKMFEDRLCHKTYLN<br/>GDHVTHPDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAIPIQIDKY<br/>LKSSKYIAWPLQGWQATFGGGDHPPKSDLEVLFGQPLGSPEFDLSMKK<br/>FAVRRFFSVYLRRKSRSKSSSLRLEEEGVVKEIDISHHVKEGF EKAD<br/>PSQFELLKVLGQGSYGKVFLVRKVKGSDAGQLYAMKVLKKATLKVRDR<br/>VRSKMERDILAEVNHPFIVKLHYAFQTEGKLYLILDFLRGGDLFTRL<br/>KEVMFTEEDVKFYLAELALALDHLHSLGIIYRDLKPENILLDEEGHIK<br/>ITDFGLSKEAIDHDKRAYSFCGTIEYMAPEVVNRRGHTQSADWWSFGV<br/>LMFEMLTGSLPFQGKDRKETMALILKAKLGMPQFLSGEAQSLLRALFK<br/>RNPCNRLGAGIDGVVEIKRHPFFVTIDWNTLYRKEIKPPFKPALGRPE<br/>DTFHFDPFTARTPTDSPGVPPSANAHHLFRGFSFVASSLIQEPSQQD<br/>LHKVPVHPVIVQQLHGNNIHFTDGYEIKEDIGVGSYSVCKRCVHKATDT<br/>EYAVKIIDKSKRDPSEEIEILLRYGQHPNIIITLKDVYDDGKFVYLVME<br/>LMRGGELLDRI LRQRYFSEREASDVLCTITKTMDYLHSQGVVHRDLKP<br/>SNILYRDESGSPESIRVCDFGFAKQLRAGNGLLMTPCYTANFVAPEVL<br/>KRQGYDAACDIWSLGILLYTMLAGFTPANGPDDTPEEILARIGSGKY<br/>ALSGGNWDSISDAAKDVVSKMLHVDPHQRLTAMQVLKHPWVVNREYLS<br/>PNQLSRQDVHLVKGAMAATYFALNRTQPAPRLEPVLSSNLAQRRGMKR<br/>LTSTRL</p> |
| <b><u>Native sequence</u></b>               | <p>Amino acids D2 – L733 (end) of human RSK3.<br/>Residue D235 of the fusion protein is equivalent to D2 of the native enzyme. The GST tag is located at residues 1 – 220.</p>                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| <b><u>Protease cleavage</u></b>             | PreScission ( <u>LEVLFQGPL</u> ) at residues 221 – 229                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            |
| <b><u>Cloning sites</u></b>                 | <i>Eco</i> R1 and <i>Sal</i> I site of pGEX6P-1                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   |

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### Nucleotide sequence of insert

ATGGACCTGAGCATGAAGAAGTTCGCCGTGCGCAGGTTCTTCTCTGTG  
TACCTGCGCAGGAAGTCGCGCTCCAAGAGCTCCAGCCTGAGCCGGCTC  
GAGGAAGAAGGTGTCTGTAAGGAGATAGACATCAGCCATCATGTGAAG  
GAGGGCTTTGAGAAGGCAGATCCTTCCCAGTTTGAGCTGCTGAAGGTT  
TTAGGACAAGGATCCTATGGAAAGGTGTTCTGGTGAGGAAGGTGAAG  
GGGTCCGACGCTGGGCAGCTCTACGCCATGAAGGTCCTTAAGAAAGCC  
ACCCTAAAAGTTCGGGACCGAGTGAGATCGAAGATGGAGAGAGACATC  
TTGGCAGAAGTGAATCACCCCTTCATTGTGAAGCTTCATTATGCCTTT  
CAGACGGAAGGAAAGCTCTACCTGATCCTGGACTTCCTGCGGGGAGGG  
GACCTCTTCACCCGGCTCTCCAAAGAGGTCATGTTACGGAGGAGGAT  
GTCAAGTTCTACCTGGCTGAGCTGGCCTTGGCTTTAGACCATCTCCAC  
AGCCTGGGGATCATCTACAGAGATCTGAAGCCTGAGAACATCCTCCTG  
GATGAAGAGGGGACATTAAGATCACAGATTTTCGGCCTGAGTAAGGAG  
GCCATTGACCACGACAAGAGAGCGTACTCCTTCTGCGGGACGATCGAG  
TACATGGCACCCGAGGTGGTGAACCGGCGAGGACACACGCAGAGTGCC  
GACTGGTGGTCTTTCGGCGTGCTCATGTTTGAGATGCTCACGGGGTCC  
CTGCCGTTCCAGGGGAAGGACAGGAAGGAGACCATGGCTCTCATCCTC  
AAAGCCAAGCTGGGGATGCCGCAGTTCCTCAGTGGGGAGGCACAGAGT  
TTGCTGCGAGCTCTCTTCAAACGGAACCCCTGCAACCGGCTGGGTGCT  
GGCATTGACGGAGTGGAGGAAATTAAGCGCCATCCCTTCTTTGTGACC  
ATAGACTGGAACACGCTGTACCGGAAGGAGATCAAGCCACCGTTCAA  
CCAGCATTGGGCAGGCCTGAGGACACCTTCCACTTTGACCCCGAGTTC  
ACAGCGCGGACGCCCACAGACTCTCCTGGCGTCCCCCGAGTGCAAAC  
GCTCATCACCTGTTTAGAGGATTGAGCTTTGTGGCCTCAAGCCTGATC  
CAGGAGCCCTCACAGCAAGATCTGCACAAAGTCCCAGTTCACCCAATC  
GTGCAGCAGTTACACGGGAACAACATCCACTTCACCGATGGCTACGAG  
ATCAAGGAGGACATCGGGGTGGGCTCCTACTCAGTGTGCAAGCGATGT  
GTGCATAAAGCCACAGACACCGAGTATGCCGTGAAGATCATTGATAAG  
AGCAAGAGAGACCCCTCGGAAGAGATTGAGATCCTCCTGCGGTACGGC  
CAGCACCCGAACATCATCACCTCAAGGATGTCTATGATGATGGCAAG  
TTTGTGTACCTGGTAATGGAGCTGATGCGTGGTGGGGAGCTCCTGGAC  
CGCATCCTCCGGCAGAGATACTTCTCGGAGCGCGAAGCCAGTGACGTC  
CTGTGCACCATCACCAAGACCATGGACTACCTCCATTCCAGGGGGTT  
GTTTCATCGAGACCTGAAGCCGAGTAACATCCTGTACAGGGATGAGTCG  
GGGAGCCCAGAATCCATCCGAGTCTGCGACTTCGGCTTTGCCAAGCAG  
CTGCGCGCGGGGAACGGGCTGCTCATGACACCCTGCTACACGGCCAAT  
TTCGTGGCCCCGGAGGTCCTGAAGCGTCAAGGCTATGATGCGGCGTGT  
GACATCTGGAGTTTGGGGATCCTGTTGTACACCATGCTGGCAGGATTT  
ACCCCTTTTGCAAATGGGCCAGACGATAACCCCTGAGGAGATTCTGGCG  
CGGATCGGCAGTGGGAAGTATGCCCTTTCTGGGGGAACTGGGACTCG  
ATATCTGACGCAGCTAAAGACGTCGTGTCCAAGATGCTCCACGTGGAC  
CCTCATCAGCGCCTGACGGCGATGCAAGTGCTCAAACACCCGTGGGTG  
GTCAACAGAGAGTACCTGTCCCCAAACAGCTCAGCCGACAGGACGTG  
CACCTGGTGAAGGGCGCGATGGCCGCCACCTACTTTGCTCTAAACAGA  
ACACCTCAGGCCCCGCGGCTGGAGCCCGTGCTGTATCCAACCTGGCT  
CAGCGCAGAGGCATGAAGAGACTCACGTCCACGCGGCTGtag