

MRCPPU Reagents and Services

Standard Operating Procedure

Preparation of CDKL5 [350 – 650]

Enzyme description:- CDKL5 [350 – 650]

Clone number:- DU 50406

Source:- Recombinant

Expression system:- *E.coli*

Tag:- N-terminal GST

Purification method:- GSH Agarose

Calculated molecular mass:-

Monoisotopic 59, 590.85 daltons

Average Mass 59, 628.20 daltons

[cysteines reduced, methionines have not been oxidised]

Theoretical pI:- 8.75

Purity:- >80 %

Enzyme storage buffer:-

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

Storage temperature:- -70 °C

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Clone Data Sheet

CDKL5 [350 - 650]

<u>Protein</u>	CDKL5 [350 - 650]
<u>Clone number</u>	DU 50406
<u>Species</u>	Human
<u>Accession number</u>	NM_003159.3
<u>Tags</u>	N-terminal GST
<u>Bacterially expressed protein</u>	<p>MSPILGYWKIKGLVQPTRLLEYLEEKYEEHLYERDEGDKWRNKKFELG LEFPNLPYYIDGDVKLTQSMAIIRYIADKHNMLGGCPKERAEISMLEGA VLDIRYGVSR IAYS KDFETLKVDFLSKLPEMLKMFEDRLCHKTYLNGDH VTHPDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAI PQIDKYLKSS KYIAWPLQGWQATFGGGDHPPKSDLEVLVLFQGPLGS SVGLPRADEGLPAN ESFLNGNLAGASLSPLHTKTYQASSQPGSTSKDLTNNNI PHLLSPKEAK SKTEFDENIDPKPSEGPGTKYLKSNRSRQONRHSFMESSQSKAGTLQPN EKQSRHSYIDTIPQSSRSPSYRTKAKSHGALS DSKSVSNLSEARAQIAE PSTSRYFPSSCLDLNSPTSPTPTRHSDTRTLLSPSGRNNRNEGTLDSRR TTTRHSKTMEELKLPEHMDSSSHSHLSAPHEFSYGLGYTSPFSSQORP HRHSMYVTRDKVRAKGLDGSLSIGQGMAARANSLQLLSPQPG</p>
<u>Native sequence</u>	Amino acids S350 – G650 (K1030 end residue) of human CDKL5. Residue S232 of the fusion protein is equivalent to S350 of the native enzyme. The GST tag is located at residues 1 – 220.
<u>Protease cleavage</u>	PreScission (<u>LEVLFQGP</u>) residues 221 - 228
<u>Cloning sites</u>	<i>Bam</i> H1 and <i>Not</i> 1 of pGEX6P-1