

MRC PPU Reagents and Services

Standard Operating Procedure

Preparation of WWP2 C838A [1 – 870]

Enzyme description:- WWP2 C838A [1 - 870]

Clone number:- DU 79146

Source:- Recombinant

Expression system:- *E.coli*

Tag:- N-terminal GST

Purification method:- GSH Sepharose

Calculated molecular mass:-

Monoisotopic 125, 624.79 daltons

Average Mass 125, 704.16 daltons

[cysteines reduced, methionines have not been oxidised

Theoretical pI:- 6.39

Purity:- >85 %

Enzyme storage buffer:-

50 mM HEPES-NaOH pH 7.5, 150 mM NaCl, 10 % Glycerol, 1 mM DTT

Storage temperature:- -70 °C

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Clone Data Sheet

WWP2 C838A [1 - 870]

<u>Protein</u>	WWP2 C838A [1 - 870]
<u>Clone number</u>	DU 79146
<u>Species</u>	Human
<u>Accession number</u>	O00308
<u>Tags</u>	N-terminal GST
<u>Bacterially expressed protein</u>	MSPILGYWKIKGLVQPTRLLLEYLEEKYEEHLYERDEGDKWRNKKFELG LEFPNLPIYYIDGDVKLTQSMAIIRYIADKHNMGGCPKERAESMLEGA VLDIIRGVSRAYSKDFETLKVDFLSKLPEMLKMFEDRLCHKTYLNGDH VTHPDFMLYDALDVLYMDPMCLDAFPKLVCFKRKIEAIPQIDKYLKSS KYIAWPLQGWQATFGGGDHPPKSDLEVLFQGPLGS MASASSSRAGVALP FEKSQTLKVVSAKPVKHNRQPRINSYVEAVDGLPSETKKTGKRIGSS ELLWNEIIILNVTAQSHLDLKWSCHTRLNELLGTASVNLNSVNLKNNGG KMENMQLTLNLQTENKGSVVSGGELTIFLDGPTVDLGNVPNGSALTDGSS QLPSRDSSGTAVAPENRHQPPSTNCGGRSRTHRHSAGSARTTPATGEQ SPGARSRHRQPVKNSGHSGLANGTVNDEPTTATDPEEPSVVGVTSPAA PLSVTPNPNTTSLPAPATPAEGERGSTSGTQQLPAAAQAPDALPAGWEQ RELPGNGRVYYVDHNTKTTWERPLPPGWEKRTDPRGRFYYVDHNTRTT WQRPTAEYVRNYEQWQSQRNQLQGAMQHFSQRFLYQSSASTHDPLGP LPPGWEKRQDNNGRYYVNHNTRTTQWEDPRTQGMIQEPALPPGWEMKYT SEGVRYFVDHNTRTTFKDPRPGFESGTKQGSPGAYDRSFRWKYHQFRF LCHSNALPSHVKISVSQTLFEDSFQQIMNMKPYDLRRRLYIIMRGEEG LDYGGIAREWFFLLSHEVILNPMYCLFEYAGKNNYCLQINPASSINPDHL TYFRFIGRFIAMALYHGKEIDTGFPLYKRMLNKRPTLKDESIDPEF YNSIVWIKENNLEECGLELYFIQDMEILGKVTTHELKEGGESIRVTEEN KEEYIMLLTDWRFTRGVEEQTKAFIDGFNEVAPLEWLRYFDEKELELML CGMQEIDMSDWQKSTIYRHYTKNSKQIQWFWQVVKEMDNEKRIRLLQFV TGTCRVPVGFAELIGSNGPQKFCIDKVGKETWLPRSHTAFNRLDLPPY KSYEQLREKLLYAIETEGFGQE
<u>Native sequence</u>	Amino acids M1 – E870 (end) of human WWP2. Residue M232 of fusion protein is equivalent to M1 of the native enzyme. The GST tag is located at residues 1 – 220.
	The enzyme has a C838A catalytic inactive mutation. Residue C838 is equivalent to A1069 of the fusion protein.
<u>Protease cleavage</u>	Prescission site (<u>LEVLFQGP</u>) at residues 221 – 228