

*Division of Signal Transduction Therapy*

**Standard Operation Procedure**

**Preparation of UBE3C 641-1083 (end)**

<b><u>Enzyme description:-</u></b>	UBE3C 641-1083 (end)
<b><u>Clone number:-</u></b>	DU45301
<b><u>Source:-</u></b>	insect recombinant
<b><u>Tag:-</u></b>	cleaved from N-terminal GST
<b><u>Purification method:-</u></b>	GSH-Sepharose, Prescission Protease
<b><u>Expression level:-</u></b>	3mg/L
<b><u>Calculated molecular mass:-</u></b>	
Monoisotopic	51053 Da
Average Mass	51084Da
[cysteines reduced, methionines have not been oxidised]	
<b><u>Theoretical pI:-</u></b>	5.65
<b><u>Purity:-</u></b>	80%
<b><u>Enzyme storage buffer:-</u></b>	
50mM HEPES pH 7.5, 150mM NaCl, 10% glycerol, 1mM DTT	
<b><u>Storage temperature:-</u></b>	-80°C
<b><u>Assay:-</u></b>	

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**Clone Data Sheet**

**UBE3C 641-1083**

<b><u>Protein</u></b>	UBE3C 641-1083
<b><u>Synonyms</u></b>	HECTH2
<b><u>Clone Number</u></b>	DU45301
<b><u>Species</u></b>	Human
<b><u>Accession Number</u></b>	Protein: Q15386 DNA NM_014671
<b><u>Tags</u></b>	cleaved from N-terminal GST
Aminoacid sequence of the expressed protein	<b>GPLGSPEFQLYVPASRHVWRFRRMGRIGPLQSTLDVGLESPPLSVSEERQLA VLTELPFVVPFEERVKIFORLIYADKQEVQDGPFLDGINVTIRRNIIYEDA YDKLSPENEPDLKKRIRVHLLNAHGLDEAGIDGGGIFREFLNELLKSGFNPN QGFFKTTNEGLLYPNPAAQMLVGDSFARHYFYLGRMLGKALYENMLVELPFA GFFLSKLLGTSADVDIHHLASLDPEVYKNLLFLKSYEDDVEELGLNFTVVNN DLGEAQVVELKFGGKDI PVTSANRIAY IHLVADYRLNRQIRQHCLAFRQGLA NVVSLEWLRMFDQOEIQVLI SGAQVP ISLEDLKSFTNYSGGYSADHPVIKVF WRVVEGFTDEEKRLLKFTVTSRPPLLGFKELYPAFCIHNGGSDLERLPTA STCMNLLKLPEFYDETLRSKLLYAIECAAGFELS</b>
Native sequence	in bold
Protease site	Precission Protease
Cloning sites	EcoR1/Not1

**DNA sequence of  
the expression  
cassette**

ATGTCCCCTATACTAGGTTATTGGAAAATTAAGGGCCTTGTGCAACCCAC  
TCGACTTCTTTTGGAAATATCTTGAAGAAAAATATGAAGAGCATTGTATG  
AGCGCGATGAAGGTGATAAATGGCGAAACAAAAAGTTTGAATTGGGTTTG  
GAGTTTCCAATCTTCCTTATTATATTGATGGTGATGTTAAATTAACACA  
GTCTATGGCCATCATACTTATATAGCTGACAAGCACAAACATGTTGGGTG  
GTTGTCCAAAAGAGCGTGCAGAGATTTCAATGCTTGAAGGAGCGGTTTTG  
GATATTAGATACGGTGTTCGAGAATTGCATATAGTAAAGACTTTGAAAC  
TCTCAAAGTTGATTTTCTTAGCAAGCTACCTGAAATGCTGAAAATGTTCCG  
AAGATCGTTTATGTCATAAAACATATTTAAATGGTGATCATGTAACCCAT  
CCTGACTTCATGTTGTATGACGCTCTTGATGTTGTTTTATACATGGACCC  
AATGTGCCTGGATGCGTTCCCAAATTAGTTTGTTTTTAAAAACGTATTG  
AAGCTATCCCACAAATTGATAAGTACTTGAAATCCAGCAAGTATATAGCA  
TGGCCTTTCAGGGCTGGCAAGCCACGTTTGGTGGTGGCGACCATCCTCC  
AAAATCGGATCTGGAAGTTCTGTTCCAGGGGCCCTGGGATCCCCGGAAT  
TCCAGCTCTATGTGCCAGCATCCAGACATGTGTGGAGGTTCCGGCGGATG  
GGGAGGATAGGCCCGCTGCAGTCCACCCTGGACGTGGGTTTGGAGTCCCC  
GCCGCTGTCTGTGTCTGAGGAAAAGACAGCTTGCTGTCTGACAGAGTTGC  
CTTTTGTGGTTCCATTTGAGGAACGAGTAAAGATCTTTCAGAGGTTGATT  
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CTCAATGCCCATGGCCTGGATGAAGCTGGCATTGATGGTGGTGGTATTTT  
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GGTCTTTAAGACTACTAATGAAGGGCTTCTGTACCCCAACCCGGCTGCT  
CAGATGCTTGTGGGAGATTCTTTTGCCAGACATTACTACTTCCTAGGCAG  
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CAGGCTTCTTTCTTTCCAAGTTGCTTGGAAACCAGTGCCGACGTGGACATT  
CACCACCTCGCTCCCTAGACCTGAGGTGTATAAGAATTTGCTCTTTCT  
GAAGAGCTACGAAGACGATGTGGAGGAGCTTGGGCTGAACTTCACTGTGG  
TGAACAATGACCTGGGAGAGGCGCAGGTAGTTGAACTAAAAATTCGGTGGG  
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GCCAGGGCCTTGCCAATGTCGTCAGCCTCGAGTGGCTCCGAATGTTTGAT  
CAGCAAGAAATTCAGGTATTAATTTCTGGTGCACAAGTTCCATAAGCCT  
AGAGGACCTAAAAATCCTTTACAACTATTTCAGGAGGCTATTCTGCAGACC  
ATCCTGTTATTAAGGTCTTCTGGAGAGTTGTGGAAGGGTTCACCTGATGAA  
GAAAAGCGCAAATGCTGAAGTTTGTAAACAAGCTGCTCTCGACCCCTCT  
CTTGGGGTTTTAAGGAGTTGTATCCCGCATTTTGTATTACACAACGGAGGCT  
CCGACCTTGAGCGGCTCCCCACAGCCAGCACCTGCATGAACCTGCTGAAG  
CTCCCCGAGTTCTATGACGAGACACTTTTTCGAAGTAAACTTCTCTATGC  
GATTGAATGTGCCGCTGGCTTTGAGCTGAGCTGA