

*Division of Signal Transduction Therapy*

**Standard Operation Procedure**

**Preparation of NFE2L2 (1-605 = full length)**

<b><u>Enzyme description:-</u></b>	NFE2L2
<b><u>Clone number:-</u></b>	DU23113
<b><u>Source:-</u></b>	Recombinant
<b><u>Tag:-</u></b>	N-terminal GST
<b><u>Purification method:-</u></b>	GSH- Sepharose, protease treatment, SEC
<b><u>Expression level:-</u></b>	0.3 mg/L
<b><u>Calculated molecular mass:-</u></b>	
Monoisotopic	67784 Da
Average Mass	67824 Da
[cysteines reduced, methionines have not been oxidised]	
<b><u>Theoretical pI:-</u></b>	4.59
<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**

**Protein name NFE2L2 1- 605 (full length)**

<b><u>Protein</u></b>	NFE2L2 (full length)
<b><u>Synonyms</u></b>	NRF2
<b><u>Clone Number</u></b>	DU23113
<b><u>Species</u></b>	Human
<b><u>Accession Number</u></b>	Protein: Q16236 Gene: NM_006164.3
<b><u>Tags</u></b>	cleaved from N-terminal GST
Aminoacid sequence of the expressed protein	<b>MMDLELPPPGLPSQODMDLIDILWRQDIDLVGSREVFDFSORRKEYELEK QKLEKERQEQLQKEQEKAFFAQLQDEETGEFLPIQPAQHIQSETSGSA NYSQVAHIPKSDALYFDDCMQLLAQTFFVDDNEVSSATFQSLVPDIPGH IESPVFIATNQAQSPETSVAQVAPVDLDGMQODIEQVWEELLSIPELQCL NIENDKLVETTMVPSPEAKLTEVDNYHFYSSIPSMEKEVGNCSPHFLNAF EDSFSSILSTEDPNQLTVNSLNSDATVNTDFGDEFYSAFIAEPSISNSMP SPATLHSLSELLNGPIDVSDLSLCKAFNQNHPESTAEFNDSDSGISLNT SPSVASPEHSVSSYGD TLLGLSDSEVEELDSAPGSVKQNGPKTPVHSS GDMVQPLSPSQGSTHVHDAQCENTPEKELPVSPGHRKTPFTKDKHSSRL EAHLTRDELRAKALHIPFVVEKIINLPVVDFNEMMSKEQFNEAQLALIRD IRRRGKNKVAAQNCRKRKLENI VELEQDL DHLKDEKEKLLKEKGENDKSL HLLKKQLSTLYLEVFSMLRDEDGKPYSPSEYSLQOTRDNVFLVPKSKKP DVKKN</b>
Native sequence	in bold
Protease cleavage	TEV
Cloning sites	BamH1 / Not1

**DNA sequence of  
the expression  
cassette**

atgtcccctatactaggttattggaaaattaagggccttgtgcaaccact  
cgacttcttttggaaatcttgaagaaaaatgaagagcatttgtatgag  
cgcgatgaaggtgataaatggcgaaacaaaaagtttgaattgggttggag  
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tag