

# Nck, a melanoma cDNA encoding a cytoplasmic protein consisting of the src homology units SH2 and SH3

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A cDNA clone encoding a novel cytoplasmic protein, nck, has been isolated from a human melanoma library using monoclonal antibodies produced against the melanoma associated antigen MUC18 (1). The nck cDNA codes for a protein of 377 amino acids which lacks a signal peptide or transmembrane region. It identifies a mRNA of 2.1kb which is expressed in a wide range of human cell lines. Nck shows no homology with MUC18 and consists of one SH2 (or B+C box) and 3 SH3 (or A box) domains. These domains have been found in nonreceptor tyrosine kinases, phospholipase c-γ, the CT10 viral oncogene crk and a variety of proteins in unicellular eukaryotes and are thought to regulate enzymatic activity and/or submembrane localization (2, 3).

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## REFERENCES

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3. Katan, M., Parker, P.J. (1988) *Nature* **332**, 203.

1 AATTCCCGAATTCCGGGAGCAGGCCCTCGTCCGCTTACGGCATCACGGCGCCGCACTGGCGCTCTGG  
70 AGCCCTCTTCAGTGTGAGCTGAAAGATGGCAGAAGAAGTGGTAGCTGGCAAAATTGGATTAT  
MA E E V V V V A K F D Y  
139 GTGGCCCAACAAAGAACAGGGTTGGACATCAGAGAAGAATGAGATTATGGCTTCCTGGATGATTCTAAG  
V A Q Q E E E E E R H E E E E E E E E E E  
208 TCTTGTTGCGGAAGTTCGAAATTCCTGAGAAACAAACAGGTTTGCTCTCTGAACTGGAGGGAGAA  
P S V P D S A S F A D D S F V D E G E R L Y  
277 AACAGTGCTCGGAAAGCAGCTCATGGAAAGAAGCTTAAACACACCTTACGGCTGGAGAAA  
N S A R K C A I V N L K D T L G I G K V K R  
346 AAACCTAOTGPGCCAGAATCTGCACTCTCTGCTGATGATGATGATGATGATGATGATGATGATGATGAT  
K P S V P D S A S F A D D S F V D E G E R L Y  
415 GACCTCAACATGGCCGCTTATGTAATTAACTACATGGCTGAGAGAGAGGATGAAATTACATTGATA  
D L N M P A Y V R F I L E F E K G D V M P E L K  
484 AAGGGCAAAAGGTTGATCAGTGAGAGATGAGGGGGCTGGCTGGACTACATGGACAA  
K G T C A A C C D G W V R G S Y N G Q  
553 GTTGATGATGCTCCCTTCAACTAACTGGAGAGGGTGAGCATGCTTGGGTGACCATGATGGGTTCT  
V G W F P S E N D P E E R E K I G H V G L V F  
622 CTGTCAGGAAATTTAGCAGCAGCTGCTGTCATTAACCTTAATACTGGGCAAGGTGGTGGTACAGGGT  
L S E R K P S L T G K F A G N P X X G I L C C  
691 CTTTACCCATTTCAGCTCATCTAAATGATGAGACTCTAAATTGAGAAAGATGATGGATGATGGTATT  
Y V V T V N O H H P L T G K F A G N P X X G I L C C  
760 GAAAAACCTGAAAATGGCCCTGGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCT  
K Y Y T V N O H H P L T G K F A G N P X X G I L C C  
829 AAAAACATGTTACCCATTAGCTGCAACAGTCAGGGTTTGCTGGCAATTCCTGGTATTATGCCAAACTCACCAGG  
E T V Y C I G Q R E  
898 GATTACATTTAGGCTTCACCTACTGGAAAGTTTGCTGGCAATTCCTGGTATTATGCCAAACTCACCAGG  
D Y I R P S L T G K F A G N P X X G I L C C  
967 CATCAAGCAGAAAGGGCATTAAATGAGAACAGGGGATTGCCCATTCCTGGCTGGCTGGCTGGCT  
I Q Y V T V N O H H P L T G K F A G N P X X G I L C C  
1036 TCUCCAAATGATTTCTCACTAAACCTAC  
F E T V Y C I G Q R E  
1105 GAGACCTGCTCTGCTCTGGGGCGCTAACTTCAC  
E T V Y C I G Q R E  
1174 GGACCAATTTTACAGTGAAACAGGAGAAATTAATCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCT  
A P I F T S E Q G E R L Y L V K H L S  
1243 CCAGAAAGTGACTGCTGTTGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCT  
1312 CAAGCTCTGGATTGAGAAATTTCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCTGGCT  
1381 CAGCCATACATATATACTATCTATGAGATTGGACATAGTATTTATATAC

Figure 1. The nucleotide and deduced amino acid sequence of human nck cDNA (clone drop 6). The SH2 domain (B, C boxes double underline) and SH3 domains (single underline) are indicated.

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