

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A9QM73	SMG7_ARATH	Arabidopsis thaliana	Nonsense-mediated mRNA decay factor SMG7	27.144759	NaN	NaN	28154133	MMTLQMDKTTASSSWERAKSIYDEI AELANKRQKAGNPPDPNLLQLLREK YEAIILESHTFSEQHNIPLWQLHY KRIEYFRLHINRVLASSTSTAAQNVK GPSKAEQIAQLKQFRFTLSEATGFY HDMILKIRSKYGLPLGSFSEDQSQ NLSDKDGKELAEVQKALKSCHRCLI YLGDLARYKGYAEGDSRSRQYASA SSYYLQAAASLWPASGNPHHQLAIVA SYSRDEFVTTYRYFRSLAVEYPPFTA RDNLIVAFDKNRQSYEKLFPVSKDS SKRLTGKGRGKGADISLKDATLVAGP EKDKVTIANEMLKAFSIRFVHLNGIL FTRTSLETFFDVLASTSSSLREVISLG SAKELTLGIDTSDSALFIVRVVTMLIF SVHNSKKELEGQSYAEIVQRVEPAR NSLTASFELLGLVIEKCVQLGDPSSS YFLPGVLVFEWLACCPDIALGSDPD DRQTVARNSFWNQFVFFNQVLSL GPTFIDDVEDETCFSNMSLYDERET ENRLALWEDYELRGFLPLPAQTILN FSRKHSFGTEGPKEKKARIKRIFAAG KALTSVIKVDQNHVYFDSKSKKFLV GVKPADDFLDSHSSPPKACNALQDN QVMIDHNSPIMQLDQQIYMGEEDD DDEVIVFKPLVTEKRKEASDQIYVPS GGFRKSDQVTTMGDFKALSGSDVAF HENQILQARGNASIQVPASVGANLL GPLQPSTQSQAMHMQQVQVQVVP ASVGANLLGLLLTSTQSQAMHMQQ VQTQAVNPQPAQSLAASRLQPIQSQ VAQPLPSRVVHFQQTQAQVSHVSPA HSQSTSFGGGSKWSPEEAAASLASSL SGFAQLGNQGHVMRNEMQGNHGV YYPAHSLPVHQSYNGNGMGGMPYS QSRTEAVFPPKIDPVLSSGVVADGL GVQSSLAKKNPISRFRHLGPPPGF NSVPAKLQKEPAPGSELGNNHLPV DDYSWLDGYQAQSSRGVGLNSSLN YATSGKPEHLGSTGNLNGPANFPF PGKQVPTSQVQADFPYFQNPQKDNF VDKNHQSTQLPEQYQGQSTWSSRH FV	None	None	None	None	None	None	None		

0