

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A5PKW4	PSD1_HUMAN	Homo sapiens	PH and SEC7 domain-containing protein 1	19.994096	NaN	S126;S156;S720	37217939	MAQGAMRFCEGDCAISPPRCRR WLPEGPVQSPASMYGSTGSLRR VAGPGPRGRELGRVTAPCTPLRGPP SPRVAPSPWAPSSPTGQPPGAOSS VVIFRFVEKASVRPLNGLPAPGGLSR SWDLGGVSPRPPTALPGSNRKLK LEASTSDPLPARGGSALPGSRNLVH GPPAPPOVGADGLYSSLPNGLGGPP ERLATLFGGPA DTGFLNQD TWSSP REVSSHAQRIARAKWEFFYGSLDPP SSGAKPPEQAPSPPGVGRSQGSGV AVGRAAKYSETDLDTVPLRCYRETDI DEVLAEREEADSAIESQPSSEGGPGT AYPPAPRPGPLPGPHPSLGSNEDE DDDEAGGEEDVDDEVFEASEGARP GSRMPLKSPVPLPGTSPSADGPD FSCVFEAILESHRAKGTSTSLASLE ALASPGTQSPFFTFELPPQPPAPRP DPPAPAPLAPLEPDSGTSSAADGPW TQRGEEEEAEARAKLAPGREPPSPC HSEDSLGLGAAPLGSEPPLSQLVSD SDSELDSTERLALGSTD T L S N G O K A DLEAAQRLAKRLYRLDGRKADVAR HLGKNNDFSKLVAGEYLKFFVFTG MTLDQALRVFLKELALMGETQERER VLAHFSQRYFQCNPALSSSEDGAHT LTCALMLLNTDLHGHNIGKRMTCC DFIGNLEGLNDGGDFPRELLKALYS SIKNEKLQWAIDEEELRRSLELADP NPKVIKRISGGSGSSPFLDLTPEP GAAVYKHGALVRKVHADPDCRKT GKRGWKSPHGILKGMILYLQKEEYK PGKALSETELKNAISIHHALATRASD YSKRPHVFLRTADWRVFLFQAPSL EQMQSWITRINVVAA MFSAPFPAA VSSQKKFSRPLPSAATRLSQEEQVR THEAKLKAMASELREHRAAQLGKK GRGKEAEEQRQKEAYLEFEKSRYST YAALLRVKLGAGSEELDAVEAALAQA GSTEDGLPPSHSSPSLQPKPSSQPRA QRHSSEPRPGAGSGRRKP	None	None	None	None	None	None	None	None	

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