	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus
0	Q86UU1	PHLB1_HUMAN	Homo sapiens	Pleckstrin homology-like domain family B member 1	17.125884	NaN	S51;S192;S220;S223;S324;S334;S381;S404;S430;S443;S461;S470;S489;S501;S518;S520;T522;S533;S539;S551;S555;S563;S578;S583;S678;S971;S1017	34725712;29485866;37217939;38665916	MDALNRNQIGPGCQTQTMVQKGPL DLIETGKGLKVQTDKPHLVSLGSGRL STAITLLPLEEGRTVIGSAARDISLQG PGLAPEHCYIENLRGTLTLYPCGNAC TIDGLPVRQPTRLTQGCMLCLGQST FLRFNHPAEAKWMKSMIPAGGRAP GPPYSPVPAESESLVNGNHTPQTAT RGPSACASHSSLVSSIEKDLQEIMDS LVLEEPGAAGKKPAATSPLSPMANG GRYLLSPPTSPGAMSVGSSYENTSPA FSPLSSPASSGSCASHSPSGQEPGPS VPPLVPARSSSYHLALQPPQSRPSGA RSESPRLSRKGGHERPPSPGLRGLL TDSPAATVLAEARRATESPRLGGQLP VVAISLSEYPASGALSQPTSIPGSPKF QPPVPAPRNKIGTLQDRPPSPFREPP GSERVLTTSPSRQLVGRTFSDGLATR TLQPPESPRLGRRGLDSMRELPPLS PSLSRRALSPLPTRTTPDPKLNREVA ESPRPRRWAAHGASPEDFSLTLGAR GRRTRSPSPTLGESLAPHKGSFSGRL SPAYSLGSLTGASPCQSPCVQRKLSS GDLRVPVTRERKNSITEISDNEDDLL EYHRRQRQERLREQEMERLERQRL ETILNLCAEYSRADGGPEAGELPSIG EATAALALAGRRPSRGLAGASGRSS EEPGVATQRLWESMERSDEENLKE ECSSTESTQQEHEDAPSTKLQGEVL ALEEERAQVLGHVEQLKVRVKELEQ QLQESAREAEMERALLQGEREAERA LLQKEQKAVDQLQEKLVALETGIQK ERDKEAEALETETKLFEDLEFQQLE RESRVEEERELAGQGLLRSKAELLR SIAKRKERLAILDSQAGQIRAQAVQE SERLARDKNASLQLLQKEKEKLTVL ERRYHSLTGGRPFPKTTSTLKEMEK LLLPAVDLEQWYQELMAGLGTGPAA ASPHSSPPPLPAKASRQLQVYRSKM DGEATSPLPRTRSGPLPSSSGSSSSS SQLSVATLGRSPSPKSALLTQNGTGS LPRNLAATLQDIETKRQLALQQKGQ QVIEEQRRRLAELKQKAAAEAQCQW DALHGAAPFPAGPSGFPPLMHHSIL HHLPAGRERGEEGEHAYDTLSLESS DSMETSISTGGNSACSPDNMSSASG LDMGKIEEMEKMLKEAHAEKNRLM ESREREMELRRQALEEERRRREQVE RRLQSESARRQQLVEKEVKMREKQ FSQARPLTRYLPIRKEDFDLKTHIES SGHGVDTCLHVVLSSKVCRGYLVKM GGKIKSWKKRWFVFDRLKRTLSYYV DKHETKLKGVIYFQAIEEVYYDHLRS AAKKRFFRFTMVTESPNPALTFCVK THDRLYYMVAPSAEAMRIWMDVIVT GAEGYTQFMN	None	None	None	None	None	None	None