

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q13393	PLD1_HUMAN	Homo sapiens	Phospholipase D1	27.26403	S629;T632;T641	S499;S561;S629	31492838	MSLKNEPRVNTSALQKIAADMSNII ENLDTRELHFEGEEVDYDVSPSPDK IQEVYIPFSAIYNTQGFKEPNIQTYLS GCPKIAQVLEVERFTSTTRVPSINLYT IELTHGEFKWQKRKFKFHFQEFHRE LLKYKAFIRIPITRRHTFRRQNVREE PREMPSLPRSENMIREEQFLGRRK QLEDYLTKILKMPMYRNYHATTEFL DISQLSFIHDLGPKGIEGMIMKRSGG HRIPGLNCCGQGRACYRWSKRWLIV KDSFLLYMKPDSGAIAFVLLVDKEFK IKVGKKEETKYGIRIDNLSRTLILKC NSYRHARWWGGAIEEFIQKHGTNF LKDHRFGSYAAIQENALAKWYVNAK GYFEDVANAMEEANEIIFITDWWLS PEIFLKRVPVVEGNNRWRLDCILKRKA QQGVRFIMLYKEVELALGINSEYTK RTLMLRHPNIKVMRHPDHVSSTVYL WAHHEKLVIIDQSVAFVGGIDLAYGR WDDNEHRLTDVGSVKRVTSGPSLG SLPPAAMESMESLRLKDKNEPVQNL PIQKSIDDVDSKLGIGKPRKFSKFS LYKQLHRHHLHDADSISSIDSTSSYF NHYSHHNLIHGLKPHFKLFHPSSE SEQGLTRPHADTGSIRSLQTGVGEL HGETRFWHGKDYCNFVKDWWQL DKPFADFIDRYSTPRMPWHDIASAV HGKAARDVARHFIQRWNFTKIMKS KYRSLYPFLLPKSQTAAHELRYQVP GSVHANVQLLRSAADWSAGIKYHEE SIHAAYVHVIENSRYHYIENQFFISC ADDKVVFNKIGDAIAQRILKAHREN QKYRVYVVIPLLPGFEGDISTGGGNA LQAIMHFNYRTMCRGENSILGQLKA ELGNQWINYISFCGLRTHAELEGNL VTELIYVHSKLLIADDNTVIIGSANIN DRSMLGKRDEMAVIVQDTETVPSV MDGKEYQAGRFAAGLRLQCFRVVL GYLDDPSEDIQDPVSDKFFKEVWVS TAARNATIYDKVFRCLPNDEVHNLI QLRDFINKPVLAKEDPIRAEEELKKI RGFLVQFFFYFLSEESLLPSVGTKEA IVPMEVWT	True	True	3.627	4.029	1.681	4.347	5.0	5.0	1.767