

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9BZL6	KPCD2_HUMAN	Homo sapiens	Serine/threonine-protein kinase D2	20.261778	S395;S396	S30;T87;S197;S198;S200;S203;S206;S212;S214;S244;T407;T438;S518;S706;S710;T717;S876	22564745;35254053	MATAPSYAGLPGSPGPGSPPPPGGL ELQSPPLLQIPAPGSGVSFHIQIGL TREFVLLPAASELAHVKQLACSIVDQ KFPECGFYGLYDKILLFKHDPTSANL LQLVRSSGDIQEGDLVEVVLSASATF EDFQIRPHALTVHSYRAPAFCDHCG EMLFGLVRQGLKCDGCGLNYHKRC AFSIPNNCSGARKRRLSSTSLASGHS VRLGTSESLPCTAEELSRSTTELLPR RPPSSSSSSSASSYTGRPIELDKMLL SKVKVPHTFLIHSYTRPTVCQACKKL LKGLFRQGLQCKDCKFNCHKRCAT RVPNDCLGEALINGDVPMEEATDFS EADKSALMDESEDSGVIPGSHSENA LHASEEEEEGEGGKAQSSLGYIPLMR VVQSVRHTTRKSSTTLREGWVVHYS NKDTLRKRHYWRLDCKCITLFQNT TNRYYKEIPLSEILTVESAQNFSLVPP GTNPHCFEIVTANATYFVGEMPPGGT PGGPGSQGAEAAARGWETAIRQALMP VILQDAPSAPGHAPHRQASLSISVSN SQIQENVDIATVYQIFPDEVLGSGQF GVVYGGKHRKTGRDVAVKVIDKLRF PTKQESQLRNEVAILQSLRHPGIVNL ECMFETPEKVFVVMKLEKLGDMLEM ILSSEKGRLEPERLTKFLITQILVALRH LHFKNIVHCDLKPENVLLASADFPF QVKLCDFGFARIIGEKSFRRSVVGTP AYLAPEVLLNQGYNRSLDMWSVGI MYVSLSGTFFNEDEDINDQIQNAA FMYPASPWSHISAGAILINNLLQVK MRKRYSDKSLSHPWLQEYQTWLD LRELEGKMGERYITHESDDARWEQF AAEHPLPGSGLPTDRDLGGACPPQD HDMQGLAERISVL