

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q14AX6	CDK12_MOUSE	Mus musculus	Cyclin-dependent kinase 12	35.285006	T588;S589;T1359	T57;T73;S235;S248;S264;S273;S275;S300;S302;S309;S311;S317;S322;S324;S331;S332;S333;S337;S340;S342;S344;S382;S384;S399;S419;S422;T511;S610;S640;S677;S681;T688;S885;T889;S1049;S1079;T1240;T1242;T1240;T1242	22645316;22517741;36852467;21606357	MPNSERHGGKKDGGSGGASGTSQPS SGGGSSNSRERHRLVSKHKRHKSK HSDKDVLVTPEAASLGTIHKPLVEYD DISSDSDTFSDDTAFKSHDRRENER RGTDNRDLRHRHRHQQHRRSRDLL KTKQTEKEKNQEVSKSGSMKDRVS GSSKRSVEGSDDYGAQLSKSGSKE SRSSKMHEKTRKERELKSGYKDRS KSHRKRETPKSYKTVASPKRRSRSP HRKWSDDSSKQDDSPSGASYGQDYD LSPPRSHTSSNYDSYKKSPGSTSRRO SISPPYKEPSAYQSSTRSPSPYSRRQR SVSPYSRRRSSSYERSGSYSGRSPSP YGRRRSSPFLSKRSLRSPLPSRKS MKSRSRSPAYSRHSSSHSKKRS RSRHSSISPVRLPLNSSLGAELSRKK KRAAAAAAAAAKMDGKESKSSPIILPK KEKLEVKESGLESKKLPRSIKSEKST PDELTVVAHSNPEVKHCLDTGKVR LDENLQKHPAKDLKAQGTKDVKVA PKEVIVTSKETETSEKETLPLPLTITS PPPLPATTPPPQTPPLPLPLPAIPL QPPLPPPQPPFSQVPVSSTSILPSSPH PRTSTLSSQTNSQPPVQVSMKTQVSI TAAIPHLKTSTLPLPLPPLPGDDD MDSPKETLPSKPAKKEKEQRTRHLL TDLPLPELPGGDPSPDSPEPKAITP PQQPYKKRPKICCPRYGERRQTESD WGKRCVDKFDIIGIIGEGTYGQVYKA KDKDTGELVALKKVRLDNEKEGFPI TAIREIKILRQLVHQSVVNMKEIVTD KQDALDFKKDKGAFYLVFEYMDHD LMGLLESGLVHFSEDHIKSFMKQL MEGLDYCHKKNFLHRDIKCSNILLN NSGQIKLADFLARLYNSESRPYT NKVITLWYRPELLLGEERYTPAIDV WSCGCILGELFTKKPIFQANLELAQL ELISRLCGSPCPAVWPDVVKLPYFNT MKPKKQYRRRLREFFSFIPSAALDLL DHMLTLDPSKRCTAEQTLQSDFLKD VELSKMAPPDLPHWQDCHELWSKK RRRQRQSGIVIEDPPPSKASRKETTS GTTAEPVKNNSPAPPQAPVKAEPG PGDAVGLGDITQQLNQSELAVLLNL LQSQTDLSIPQMAQLLNHNSNPEMQ QQLEALNQSISALTEASSQQQDSESI APEESLKEVPSVPVVLPPAEQTTPEA SNTPADMQNVLAVLLSQLMKTQEP AGNLEENTNDKNSGPGPRRTPTM PQEEAAACPPHILPPEKRPPEPPGPP PPPPPPPLVEGDLSSAPQELNPAVTA ALLQLLSQPEAEPPGHLPHHEHQALR PMEYSTRSHPNRTYGNTDGPETGFS SADTDERSSGPALTESLVQTPVKNR TFSGSVSHLGESNSYQGTGSVQFPG DQDLRFTRVPLALHSVVGQPFLKSE GNSNSVVHAETKLQNYGELGPGTTG

