

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q3V0T4	ITAD_MOUSE	Mus musculus	Integrin alpha-D	12.769218	S1147	NaN	34418053;29187734	MVFKTIWIERYRKLINLFRALASCHG SNLDVEKPVVFKEDAASFGQTVVQF GGSRLVVGAPLEAVAVNQTGQLYDC APATGVCQPILLHIPLEAVNMSLGLS LVADTNNSQLLACGPTAQRACAKN MYAKGSCLLLGSSLQFIQAI PATMPE CPGQEMDIAFLIDGSGSIDQSDFTQ MKDFVKALMGQLASTSTSFSLMQY SNILKTHFTTFTEFKSSLSPOSLVDAIV QLQGLTYTASGIQKVKELFHSKNG ARKSAKKILIVITDGGKFRDPLEYRH VIPEAEKAGIIRYAIGVGD AFREPTAL QELNTIGSAPSQDHVFKVGNFVALR SIQRQIQEKIFAIEGTESSSSSFQHE MSQEGFSSALSMDGPVLGAVGSFS WSGGAFLYPSNMRSTFINMSQENE DMRDAYLGYSTALAFWKGVHSLILG APRHQHTGKVVIFTQESRHWRPKSE VRGTQIGSYFGASLCSVMDRDRGST DLVLIGVPHYEHTRGGQVSVC PMP GVRSRWHCGTTLHGEQHPWGRF GAALTVLGDVNGDSLADVAIGAPGE EENRGAVYIFHGASRQDIAPSPSQRV TGSQFLRLQYFGQSLSGGQDLTQD GLVDLAVGAQGHVLLLRSLPLLKVGI SIRFAPSEVAKTVYQCWGRTPTVLEA GEATVCLTVRKGSPDLLGDVQSSVR YDLALDPGRLISRAIFDETKNCTLTR RKTGLGDHCETMKLLLPDCVEDAV TPILRLNLSLAGDSAPSRNLRPVLAV GSQDHVTASFPFEKNCKQELLCEGN LGVSNFNSGLQVLEVGSSPELTVTVT VWNEGEDSYGTLIKFYPAELSYRRV TRAQQPHPYPLRLACEA EPTGQESL RSSSCSINHPIFREGAKATFMITFDV SYKAFLGDRLLLRASASSENK PETS KTAFLQLELPVKYTVYTVISRQEDSTK HFNFSSSHGERQKEAEHRYRVN NL SPLTLAISVNFVWPILLNGVAVWDVT LRSPAQGVSCVVSQREPPQHSDLLTQI QGRSVLDCAIADCLHLRCDIPSLGTL DELDFILKGNLSFGWISQTLQKKVLL LSEAEITFN TSVYSQLPGQEAFLRAQ VSTMLEEYVVYEPVFLMVFS SVGGL LLLALITVALYKLGFFKRQYKEMLDL PSADPD PAGQADSNHETPPHLTS