

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion
Q8BIH0	SP130_MOUSE	Mus musculus	Histone deacetylase complex subunit SAP130	41.736082	S134;T135;S274;T278;T279;T280;S290;T293;S295;S301;T320;S428;S434	T329;S416;S439;S864;T865;S884	21606357;22517741;36852467;34678516	MSSSQFPRLGTPSPGLSQPPSQIASS GSAGLINQVATVND EAGRDADVGTR EHVGPSSSLPPREEKQEPVVVVRPYP QVQMLPAHHAVASATPVAVTAPPAH LTPAVPLSFSEGLMKPPKPTMPSRP IAPAPPSTMSLPPKVPQVTVTMESS IPQASAI PVATISGQQGHPSNLHHIM TTNVQMSIIRSNA PG PPLHIGASHLP RGA AAAAVMSSSKVTTVLRPTSQLP NAATAQPAVQH LIHQIQSRPPVTTTS STIPPAVVATVSATRAQSPVITTTAAH AADSTLSRPTLSIQHPPSAAISIQRPA QSRDVTTRITLPSHPALGTPKQQLHT MAQKTI FSTGTPVAAATVAPILATNT LPSTTTAGSVSHTQAPTSTIVTMTMP SHSSHATAVTTSNIPVAKVVPQQITH TSPRIQPDYPPERSLLIPISGHRASPN PVAMETRNDNRPSVVPVQFYFLPTY PPSAYPLAAHTYTPITSSVSTIRQYPV SAQAPNSTITAOTGVGVASTVHLNP MQLMTVDASHARHIQGIQPAPISTQ GIQPAPIGTSGIQPAPIGTPGIHSAPI NTQGLQPAAMANQQPQPEGKTSAV VLADGATIVANPISNPFSAAPAATTV VQTHSQSASTNTPAQGSSPRPSILRK KPATDGM AVRKTLLPPQPPDVATPR VESSMRSASGSPRPAGAKPKSEVHV SIATPVTV SLETISNQNAEOPTVAVP PTAQPPPTIPSMIAAASPPSQPAIAL STIPGAVPVTPPITIAATPTLSAPVGG TPSTVLGPPVPEIKVKEEAEPVDITRP VSTVPPLATNTVSPSLALLASNL SMP PSDLPPGASPRKPRKQQH VISTEEG DMMETNSTDDEKSAKSLLVKAEK RKSPPEKVIDEEGVRYVPVRRPPITL LRHYRNPWKAAYHHFQRYSDVRVK EKKAMLQEI ANQKGVSCRAQGWK VHLC AAQLQLTNLEHDVYERLTNL QEGII PKKKAATDDDLHRINELIQGN MQRCKLVMDQISEARDSMLKVL DH KDRV LKLLNKNGT VKVSKLKRKEK V	None	None	None	None	None