

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9H0E3	SP130_HUMAN	Homo sapiens	Histone deacetylase complex subunit SAP130	23.525536	S108;T122;T129;T305;T306;S327;S331;T346;S349;S460;S480;T491;S495;T503;T505;T508;S509;S510;S512;T513;S635;S639;T646;T657	T355;S442;S465;S855;T856;S875	30379171;33214551;30059200;27655845;31492838;32119511;22661428;28657654;32574038;29351928;29237092;34019948;30397120	MGPPRHPQAGEIEAGGAGGRRRLQ VEMSSQFPRLGAPSTGLSQAPSI ANSGSAGLINPAATVNDSEGRDSEV SAREHMSSSSSLQSREEKQEPVVVR PYPQVQMLSTHHAVASATPVAVTAP PAHLTPAVPLSFSEGLMKPPPKPTM PSRPIAPAPPSTLSLPPKVPQVTVT MESSIPQASAI PVATISGQQGHPSNL HHIMTTNVQMSIIRSNAPGPPLHIGA SHLPRGAAAAAVMSSSKVTTVLRPT SQLPNAATAQPAVQHIIHQPIQSRPP VTTSNAIPPAVVATVSATRAQSPVITT TAAHATDSALS SRPTLSIQHPPSAAISI QRPAQSRDVTTRITLPSHPALGTPKQ QLHTMAQKTIFSTGTPVAAATVAPIL ATNTIPSATTAGSVSHTQAPTSTIVT MTVPSHSSHATAVTTSNIPVAKVVP QQITHTSPIQPDYPAERSSLIPISGH RASPNPVAMETRSDNRPSVPVQFQY FLPTYPPSAYPLAHTYTPITSSVSTIR QYPVSAQAPNSAITAQTG VGVASTV HLNPMQLMTVDASHARHIQIQPAP ISTQGIQPAPIGTPGIQPAPLGTQGIH SATPINTQGLQPAPMGTQQPQPEGK TSAVV LADGATIVANPISNPFSAAPA ATTVVQTHSQSASTNAPAQGSSPRP SILRKKPATDGAKPKSEIHVSMATPV TVSMETVSNQNNNDQPTI AVPPTAQQ PPPTIPTMIAAASPPSQPAVALSTIPG AVPITPPITIAAAPP SVTVGGSLSSV LGPPVPEIKVKEEVEPMDIMRPVSAV PPLATNTVSPSLALLANL SMP TSD LPPGASPRKKPRKQ QHVISTEEGDM METNSTDDEKSTAKSLLVKAEKRKS PPKEYIDEEGVRYVVRPRPPITLLRH YRNPWKAAYHHFQRYSDVRVKEEK KAMLQEIANQKGVSCRAQGK VHL CAAQLLQLTNLEHDVYERLTNLQEG IIPKKAATDDDLHRINELIQNMQR CKLVMDQISEARDSMLKVL DHKDR VLKLLNKNGTVKKVSKLKRKEKV