

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9H0E3	SP130_HUMAN	Homo sapiens	Histone deacetylase complex subunit SAP130	28.053026	S108;T122;T129;S300;T304;T305;T306;T311;S313;T319;S321;S327;S331;T346;S349;T355;S454;S460;S480;T491;S495;T503;T505;T508;S509;S510;S512;T513;S635;S639;T646;T657	T355;S442;S465;S855;T856;S875	32119511;34725712;32574038;22661428;30397120;29237092;33214551;30059200;35289036;30379171;34019948;35254053;37340703;27655845;28657654;31492838;34846842;35132862;29351928	MGPPRHPQAGEIEAGGAGGRRRLQ VEMSSQFPRLGAPSTGLSQAPSI ANSGSAGLINPAATVNDSEGRDSEV SAREHMSSSSSLQSREEKQEPVVVR PYPQVQMLSTHHAVASATPVAVTAP PAHLTPAVPLSFSEGLMKPPPKPTM PSRPIAPAPPSTLSLPPKVPQVTVT MESSIPQASAI PVATISGQQGHPSNL HHIMTTNVQMSIIRSNAPGPPLHIGA SHLPRGAAAAAVMSSSKVTTVLRPT SQLPNAATAQPAVQHIIHQPIQSRPP VTTSNAIPPAVVATVSATRAQSPVITT TAAHATDSALS RPTLSIQHPPSAAISI QRPAQSRDVTTRITLPSHPALGTPKQ QLHTMAQKTIFSTGTPVAAATVAPIL ATNTIPSATTAGSVSHTQAPTSTIVT MTVPSHSSHATAVTTSNIPVAKVVP QQITHTSPRIQPDYPAERSSLIPISGH RASPNPVAMETRSDNRPSVPVQFQY FLPTYPPSAYPLAHTYTPITSSVSTIR QYPVSAQAPNSAITAQTG VGVASTV HLNPMQLMTVDASHARHIQIQPAP ISTQGIQPAPIGTPGIQPAPLGTQGIH SATPINTQGLQPAPMGTQQPQPEGK TSAVV LADGATIVANPISNPFSAAPA ATTVVQTHSQSASTNAPAQGSSPRP SILRKKPATDGAKPKSEIHVSMATPV TVSMETVSNQNNNDQPTIAVPPTAQQ PPPTIPTMIAAASPPSQPAVALSTIPG AVPITPPITTTIAAAPPSTVVGGLSSV LGPPVPEIKVKEEVEPMDIMRPVSAV PPLATNTVSPSLALLANL SMP TSD LPPGASPRKKPRKQHVISTEEGDM METNSTDDEKSTAKSLLVKAERKS PPKEYIDEEGVRYVVRPRPPITLLRH YRNPWKAAYHHFQRYSDVRVKEEK KAMLQEIANQKGVSCRAQGKVVHL CAAQLLQLTNLEHDVYERLTNLQEG IIPKKAATDDDLHRINELIQNMQR CKLVMDQISEARDSMLKVL D HKDR VLKLLNKNGTVKKVSKLKRKEKV

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