	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	ext reg
0	Q9Y2H2	SAC2_HUMAN	Homo sapiens	Phosphatidylinositide phosphatase SAC2	26.21016	S1103	S827;S830;S878;S881;S907;S910;S1103	29351928	MELFQAKDHYILQQGERALWCSRR DGGLQLRPATDLLLAWNPICLGLVE GVIGKIQLHSDLPWWLILIRQKALVG KLPGDHEVCKVTKIAVLSLSEMEPQ DLELELCKKHHFGINKPEKIIPSPDD SKFLLKTFTHIKSNVSAPNKKKVKES KEKEKLERRLLEELLKMFMDSESFY YSLTYDLTNSVQRQSTGERDGRPLW QKVDDRFFWNKYMIQDLTEIGTPDV DFWIIPMIQGFVQIEELVVNYTESSD DEKSSPETPPQESTCVDDIHPRFLVA LISRRSRHRAGMRYKRRGVDKNGN VANYVETEQLIHVHNHTLSFVQTRG SVPVFWSQVGYRYNPRPRLDRSEKE TVAYFCAHFEEQLNIYKKQVIINLVD QAGREKIIGDAYLKQVLLFNNSHLTY VSFDFHEHCRGMKFENVQTLTDAIY DIILDMKWCWVDEAGVICKQEGIFR VNCMDCLDRTNVVQAAIARVVMEQ QLKKLGVMPPEQPLPVKCNRIYQIM WANNGDSISRQYAGTAALKGDFTRT GERKLAGVMKDGVNSANRYYLNRF KDAYRQAVIDLMQGIPVTEDLYSIFT KEKEHEALHKENQRSHQELISQLLQ SYMKLLLPDDEKFHGGWALIDCDPS LIDATHRDVDVLLLLSNSAYYVAYYD DEVDKVNQYQRLSLENLEKIEIGPEP TLFGKPKFSCMRLHYRYKEASGYFH TLRAVMRNPEEDGKDTLQCIAEMLQ ITKQAMGSDLPIIEKKLERKSSKPHE DIIGIRSQNQGSLAQGKNFLMSKFSS LNQKVKQTKSNVNIGNLRKLGNFTK PEMKVNFLKPNLKVNLWKSDSSLE TMENTGVMDKVQAESDGDMSSDN DSYHSDEFLTNSKSDEDRQLANSLE SVGPIDYVLPSCGIIASAPRLGSRSQS LSSTDSSVHAPSEITVAHGSGLGKG QESPLKKSPSAGDVHILTGFAKPMDI YCHRFVQDAQNKVTHLSETRSVSQQ ASQERNQMTNQVSNETQSESTEQT PSRPSQLDVSLSATGPQFLSVEPAHS VASQKTPTSASSMLELETGLHVTPSP SESSSSRAVSPFAKIRSSMVQVASIT QAGLTHGINFAVSKVQKSPPEPEIIN QVQQNELKKMFIQCQTRIIQI	None	None	None	None	None	None	None	None	No