

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q96JG6	VPS50_HUMAN	Homo sapiens	Syndetin	22.327667	S15	S15;S494;S498;S559;S561	37217939;39302247	MQKIKSLMTRQGLKSPQESLSDLGA IESLRVPGKEEFRELRQPSDPAEQ ELINSIEQVYFVSDSFDIVKYELEKLP PVLNLQELEAYRDKLKQQAQAVSKK VADLILEKQPAYVKELERVTSLQTGL QLAAVICTNGRRHLNIAKEGFTQASL GLLANQRKRQLLIGLLKSLRTIKTLQ RTDVRLSEMEEEDYPGAIQLCLEC QKAASFVKHYSCISELNSKLQDTLEQ IEEQDVALSKICKNFNDINHYTKVQQ AYRLLGKTQTAMDQLHMHFTQAIH NTVFQVVLGYVELCAGNTDTKFKQL QYKDLCTHVTPDSYIPCLADLCKAL WEVMLSYYRTMEWHEKHDNEDTA SASEGSNMIGTEETNFDRGYKKKL EHGLTRIWQDVOLKVKTYLLGTDLSI FKYDDFIFVLDIISRLMQVEEFCGS KSEVLQESIRKQSVNYFKNYHRTRL DELRMFLENETWELCPVKSNFSILQ LHEFKFMEQSRSPSVSPSKQPVSTS SKTVTLFEQYCSGGNPFQIQAHHKD EETEDVLSANGYESDEQEKSAYQEY DSDSDVPEELKRDYVDEQTGDGPVK SVSRETLKSRKSDYSLNKVNAPILT NTTLNVIRLVGKYMQMMNLIKPIAF DVIHFMSQLFDYLYAIYTFGRNDS LESTGLGLSSRLRRTLNRQESLIDL EVSADPTATLTAAEERKEKVPSPHLS HLVVLTSGDITLYGLAERVVATESLVF LAEQFEFLQPHLDAVMPAVKKPFLQ QFYSQTVSTASELRKPIYVIVAGKAL DYEQMLLLMANVKWDVKEIMSQH NIYVDALLKEFEQFNRRLENEVSKRV RIPLPVSNILWEHCIRLANRTIVEGYA NVKCCSNEGRALMQLDFQQFLMKL EKLTDIRPIPDKEFVETIKAYYLLEN DMERWIKEHREYSTKQLTNLVNVC LGSHINKKARQKLLAAIDDIRPKR	True	False	4.012	3.01	False	0.906	1.974	2.502	4.0