

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
Q8BUH8	SEN7_MOUSE	Mus musculus	Sentrin-specific protease 7	25.385059	NaN	S12;S13;S25;S434;S435	28528544	MDRARPGRRRASSEIVTEGKRKSS PADLQKITKLLTVKSEDLAQSPK LRGSECWWTSLRNVICLDHKKPK AARGCPPKGLPKRHLRMLTNVLW TDLGRFRKTLPRKDALNCAPSKVQ SDSLPSTSVDSIETCQRDLPLHQLN LSERTPRVILTDIRQTELGKRYLKIP VTEASLSDTANLSEQLSSSSDGSLE SCQSVNHHKSFLESSESGPKPSRTGDV PAKEAACGGQKQGGDDGGVTPEMAA PHPKDFNTGNKGCYLEEGTSNKN TSYSYSEMDHTPVSRKRKRGRSNF HDSHNSKTSLDKPTHTKEEENDSS VSRKLEESGEDSHQDPAPPEGLAPE SLESEATNLRFAAGQEPDASAASG RASSPNKLESSASSEVSENSVAVK GEALTLKEASPPGGSSEESQLLISAE PIVSSDEEGPVEHKNSVILKLP EIMSENQGTSDPQLSELTGACESV QVTSELPYNDVENISCIKNSSEM DLKLDIFITCVYIGKIKGTPKGCVTFT KKYKIPFQVSTNEISLTVDTARLKRF GLWESKDEDHSKRSHAILFLWLSSD YLQDIQTQLENPMLSQQSKANEFIF LELNSSISQREELKLDIMMEISTKN GNLHLSCLPWWQALPLFQDLSPQE ISFLHYYSASALPTAAGADMKKKS VSQPSNSDTIKPTYTFLHKQSSGCYS LSITSSPEEEWQEVNRTGPVQKLIVY PPPPTKGGGLGVTNEDLECEEGEFL NDVIIDFYLYLLEKASDELVERSHI FSSFFYKCLTRKENNLTEDNPDLV AQRHRRVRTWTRHINIFNKDYIFV PVNESSHWYLAVICFPWLEAVYED CPQTVSQQFQQSQHDKMTDN DPHTTSTVSTSAEDSQSTEVNMSVP KKMCKRPCILILDSLKAASIQNTVQN LREYLEVEWEVKRKRTHREFSKTNM VDLCPKVPKQDNSSDCGVYLLQYVE SFFQDPIVNFELPIHLEKWFPRHVIK TKREDIRELILKHLQKGGSC	True	False	1.575	4.436	1.295	1.051	0.894	1.066	0.865

0