

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q99PU7	BAP1_MOUSE	Mus musculus	Ubiquitin carboxyl-terminal hydrolase BAP1	25.810548	NaN	S292;S369;S395;T492;S520;S536;S584	29187734	MNKGWLELESDPGLFTLLVEDFGV KGVQVEIYDLQSKCQGPVYGFILF KWIEERRSRKYSTLVDDTSVIDDI VNNMFFAHQLIPNSCATHALLSVLL NCSNVDLGP T LSRMKDFTKGFSPES KGYAIGNAPELAKAHNSHARPEPRH LPEKQNGLSAVRTMEAFHFVSYVPI TGRLFELDGLKVYPIDHGPWGEDEE WTDKARRVIMERIGLATAGEPYHDIR FNLMAVVPDRRIKYETRLHVLKVN QTVLEALQQLIRVTQPELIQTHKSQE SQLPEESKPASSKSPGLEAGRT SECTQTDGAEVAVAGSCTTTHSPP SKCKLVVPPGSSLNGVPPNPAPIVQ RLPAFLDNHNYAKSPMQEEDLAA GVGRSRVPVRAPQYSEDEDDYEDE DEDVQNTNPAIRYKRKGTGKPGSLS NSSDGGQLSVLQPNNTINVLTEKLQES QKDLVPLSIKTSSGAGSPAVAVPTH SQPSPTPSNESTDTASEIGSAFNSPL RSPRSANPTRPSSPVTSHISKVLFGE DDSLLRVDCIRYNRAVRDLGPVISTG LLHLAEDGVLSPALTEGGKGSST RSSQGSQSSGLEEKVEVETESRD KPGLNRSSEPLSGEKYSPKELLALK CVAEAIANYEACLKEEVEKRFKID DQRRTHNYDEFICTFISMLAQEGML ANLVEQNISVRRRQVSGRLHKQR KPDRRKRSRPYAKAKRQ	True	False	3.649	4.719	1.753	1.492	0.705	1.598	1.535