

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
D3YXK2	SAFB1_MOUSE	Mus musculus	Scaffold attachment factor B1	35.97895	T582	S24;S55;S79;T188;S195;S197;S209;S405;S406;S437;S602;S604;S623;S626	40021952;28528544;34887587;25153642;39627609	MAETLSGLGDASAAGAAVSSAASE TGTRRLSDLRVIDLRAELKKNLDSS GNKSVLMERLKAIEDEGGNPDEIE VTSECNKKMPKRPSPKGRKPEDEGVE DNGLEENSGDGGQEDVETSLENLQD MDMMDISVLEADIDNGSVADCV E EEEEATLPEGLADSTELVEGDLKGLP EQLQEHAIDDKDTVNNVDTSSDFT MLQEMEEASLEPENEKILDILGETC KSEPVKEEGSELEQPPAQATSSVGP DRKLAEEEDLFESC GHPEEEEEEE EDQEEEQEEEGDLALASSKSESPS TRCQWSEADAPLAVVKREPADAPGG GTGMDREPVGLEEPVEQSSTAAQLP EATSQELVRAPTAALSPEPODSKEDV KKFAFDACNDVPAPPKESASSEGAD QKMSSVEEDSDTKRLSREEKGRSSC GRNFWVSGLSSTTRATDLKLNLF SRY GKVVGAKVVTNARSPGARCYGFVTM STAE EATKCI SHLHKTELHGKMISVE KAKSEPTGKRVPDRRDGDSKKEKAS TDRSANLKREEKGERKDDAKTDD GSTEKSKDADDQKPGPSERSRTTKS GSRGTERTVVMDKSKGVPVISVKTS GSKERASKSQDRKSASREKRSVVSF DKVKESRKS RDSERRE RERSEREQ RLQAQWEREERERLEIARERLAFHR HRLERERMERERLERERMHVQEER RREQERIHREEREELRRQELRYEQE RRPAVRRPYEVDGRRDDAYWPEAKR AALDDRYHSDFSRQDRFHDFDHRD RGRYPNHSDRREGSRSMMDGREG QHYP ERHGGPERHGRD SRD GWGYG SNKRLSEGRGLPPPPRRDWGEHGR RLEDDRAWQGTADGGMMERD HKR WQGGERSMSGHSGPGHMMNRGG MSGRGSFAPGGASRGHVIPRGGMQ AGFGGQSRGSRPSDARFTRRY	True	False	1.641	4.539	False	False	False