

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
A0A087WR50	A0A087WR50_MOUSE	Mus musculus	Fibronectin	22.380232	NaN	NaN	37507081	MLRGPGRLLLLAVLCLGTSVRC EAGKSKRQAQQIVQPSVAVSOSK PGCFDNGKHQINQWERTYLGNA LVCTCYGSSRGFNCEKPEPEETCF DKYTGNTYKVGDTYERPKDSMIWDC TCIGAGRGRISCTIANRCEGGQSYK IGDKWRRPHETGGYMLECLCLGNG KGEWTCPIAEKCFDHAAGTSYVVG ETWEKPYQGWMMVDCCLGEGNG RITCTSRNRCNDQDTRTSYRIGDTW SKKDNRRGNLLQCVCTGNRGEWKC ERHALQSASAGSGSFTDVRTAIYQPQ THPQPAPYGHCVTDSGVVYSVGMQ WLKSQGNKQMLCTCLGNGVSCQET AVTQTYGGNSNGEPCVLPFTYNGRT FYSCCTTEGRQDGHLCSTTSNYEQ DQKYSFCTDHAVLVQTRGGNSNGA LCHFPLYNNRNYTDCSEGRDN MKWCGTTQNYDADQKGFPCMAA HEEICTTNEGVMYRIGDQWQKQHD LGHMMRCTCVGNRGEWACIPYSQ LRDQCIVDDITYNVNDTFHKRHEEG HMLNCTCFGQGRGRWKCDPIDQCO DSETRTFYQIGDSWEKVFHGVRYQC YCYGRGIGEWHCQPLQTYPGTTGPV QVIITETPSQPNSHPIQWNAPEPSHI TKYLRRWRPKTSTGRWKEATIPGHL NSYTIKGLTPGVIEGQLISIQYGHR EVTRFDFTTSASTPVTSNTVTGETAP YSPVVATSESVTEITASSFVSVWSA SDTVSGFRVEYELSEEGDEPQYLDLP STATSVNIPDLLPGRKYVNVVYQISEE GKQSLILSTSQTAPDAPPDPTVDQV DDTSIVVRWSRQAPITGYRIVYSPSV EGSSTELNLPETANSVTLSDLQPGV QYNITTYAVEENQESTPVFIQQETT TPRSDNVPPPTDLQFVELTDVKVTIM WTPPDSVSVGYRVEVLPVSLPGEHG QRLPVNRNTFAEITGLSPGVTYLFKV FAVHQGRESNPLTAQQTIKLDAPTN LQFVNETDRTVLVWTPPRARIAGY RLTAGLTRGGQPKQYVNGPLASKYP LRNLQPGSEYTVTLVAVKGNQQSPK ATGVFTTLQPLRSIPPYNTEVTETIV ITWTPAPRIGFKLGVRSQGEAPRE VTSDSGSIVVSGLTPGVEYTYTIQVLR DGQERDAPVNRVVTPLSPPTNLHL EANPDTGVLTVSWERSTTPDITGYRI TTTTPTNGQQGTSLEEVHADQSSCT FENLNPGLYNSVYTVKDDKESAPI SDTVVPAVPPPTDLRFTNIGPDTMRV TWAPPSIELTNLLVRYSPVKNEEDV AELSISPSDNVVLNLLPGTEYLV VSSVYEQHESIPLRGRQKGLDPTG FDSSDITANSFTVHWVAPRAPITGYII RHHAHESVGRPRQDRVPPSRNSITL TNLNPGETEYVSVIANGREESPLIG QQATVSDIPRDLEVIASPTSLISWE PPAVSVRYRITYGETGGNSPVQEF VPGSKSTATINNIKPGADYITILYAVT GRGDSPASSKPVSYNYKTEIDKPSQM QVTDVQDNSISVRWLPSTSPVTGYR VTTTTKNGLPSKTKTASPDQTEMTI EGLQPTVEYVSVYANRNGESQPL VQTAVTNIDRPKGLAFTDVDVDSIKI AWESPGQVSRVRYTSSPEDGIRE LFPAPDGEDDTAELQGLRPGSEYTV SVVALHDDMESQPLIGQSTAIAPT NLKFSQVTPTSFTAQWIAPSVQLTGY RVRVNPKEKTGPMKEINLSPDSSVI VSGLMVATKYEVSVYALKDLTISRPA QGVITLENVSPRRARVTDATETI TISWRTKETITGFQVDAIPANGQTP VQRSISPDRVSYTITGLQPGTDYKHL YTLNDNARSSPVIIDASTAIDAPSNLR FLTTTPNLLVSWQAPRARITGYIKY EKPGSPPREVVPRPRGVTEATITGL EPGTEYTIYVIALKNNQKSEPLIGRKK	None	None	None	None	None	None	None		

