	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P12080	ITA2_DROME	Drosophila melanogaster	Integrin alpha-PS2	30.404062	NaN	NaN	33925313;28604694	MSGDSIHRRRMALHCPITSLILLLIA MSAHGYNIDLPSYVRFRQSSNSMFG FSIAMHKGRSGFYGNQNNVSLIVGA PKFDTSRYQQGVTEAGGVFKCSLND DDCKLVPFDSKGNNRNVDKEVVDR KSYQWLGATVATGRDSDLVVACAPR YVFHTMTPSRAFRIDPVGTCFTSHN FEEFYEVSPCRTNNWGYHRQGSCQ AGFSAAINGNGSRLFIGAPGSWYWQ GQTYSIPPDAKFPFKPPLYQPFGTGG MASSHDVTRPENQVFSTSESASVND DSYLGYSMVTGDFDGDRSEDVAIGM PRGGNLVGRIVVNRWNMANIFNITG RQIGEYFGYSLATSDVDGDGLDDLLI GAPMYTDPDNVEGKYDVGRVYILLQ GGPTEEKRWTTEHIRDGYHSKGRFG LALTTLGDVNGDGYGDFAVGAPYDG PEGRGVVYIFHGSPMGPLAKPSQIIK SEQLVEGAPYPRTFGFALSGGLDMD GNTYPDLAVGAYSSDQVFIFKSRPVA AVNAETSFASNSKLISLDDRSCQLVR DHKKVPCMLLTTCWSYTGRYLPEQL DFDVSWLLDAKKLLNPRMFFLRDE GKNIRNQTIRLNYGQKYCLNETVYLL DKVQDKLTPLEVEARYNLRSSRPLD PMVRHRRSILEPVIDQNREIVLRDAI NIQKNCGPDNICEPDLKLKVSTVDK YLFGSPEPLVIEVFISNTNEDAFEAA FYMVTPPDLQFRKLQQLGEKKDTPI TCSPPTPENNHTLKCDIGNPLESGKI AHFKISLVPEEKYGSSSSYDFYWEAN STNLEKPGSEYDNKIRQSVGIWVDT DLDIKGTSLPDYQLYKADDYKELENA TKEDDIGPQVVHIYEIRNNRPSIIEEA EVFIHLPYETIVGDPLMYLLNQPETG GKIQCDDVAFNEYNLLLDEKLVKKS YLQAQGAIWNSAQVSGQSSSSSSSG GASVHIEKARGEGFVRGVLVSNSTD AGDKLSPKQVEQRRQEDTLEALGDA SFVHRDRASQAVQEPQVNQTSFTTY STSSSSSGSGAPSAQLRGHSTQGHI QMAGPVQHTSSSSSSNYRSWPAQQ QQQHQQLLLAGSGGSGLGSPVTFN DKSQFGGRNNNFHTGTLDLGTLNR GNVDNELYRSQGQYQNPSQSLGQS QGQFQANANQGHYQGQNQAQFQA RNPGFQGQTSYQGQTQYSGQPGGY QTHHVTYSSGSKPYYGRENEDFYDE DNLQQATPGHWSSSSSSSSSSGTRR LRRSNDKDGATEKPLQIDLNSPCQS ARCKSIRCVVTNLGTEDGDAAFVAIR ARMVAKTMEKLASNVPLNVSTLAVA NVTLLPFIGAPKDAIVKTHEIFYKAEP EPLQVPDVVPLWVVVLAACAGALIFL LLVWLLYKCGFFNRNRPTDHSQER QPLRNGYHGDEHL	None	None	None	None	None	None	None	None	None