

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
P11584	ITBX_DROME	Drosophila melanogaster	Integrin beta-PS	25.496522	NaN	T831;T843	33925313	MILERNRRQLALLMIAILAAIAGQT DAQKAAKLTAVSTCASKEKCHTCIQT EGCAWCMQPDFKGGQSRCYQNTSSL CPEEFAYSPITVEQILVNNKLTNYK AELAAGGGGSAMSGSSSSSYSSSS SSSFYSQSSSGSSASGYEYSAGEI VQIQPQSMRLALRVNEKHNIKISYSQ AEGYPVDLYYLMDSLKSMEDDKAKL STLGDKLSETMKRITNNFHLGFGSF VDKVLMPYVSTIPKLEHPCENCKA PYGYQNHMPLNNNTESFSNEVKNA TVSGNLDAPEGGFDAIMQAIACRSQI GWREQARRLLVFSTDAGFHYAGDG KLGGVIAPNDDGECHLSPKGEYTHST LQDYPSISQINQKVKDNAINIIFAVTA SQLSVYEKLVHEHIQSSAAKLDNDS SNVVELVKVEYRKISSVEMKDNAT GDVKITYFSSCLSNQPEVQTSKCDN LKEGQVVSFTAQIQLLKCPEDPRDW TQTIHISPVGINEVMQIQLTMLCSCP CENPGSIGYQVQANSCSGHGTSMC GICNCDDSYFGNKCECSATDLTSKF ANDTSCRADSTSTTDCSGRGHCVCG ACECHKRPNPIEHSKHCEDNDFSC ERNRNQLCSGPDHGTCCEGRCKCK PGWTGSNCGCQESNDTCMPGGGE ICSGHGTCECGVCKCTVNDQGRFSG RHCEKCPCTSGRCQELKDCVQCQM YKTGELKNGDDCARNCTQFVPGVE KVEIDETKDEQMCKFFDEDDCKFM FKYSEQELHVYAQENKECPAKVFM LGIVMGVIAAIVLVGLAILLLWKLTT IHDRREFARFEKERMNAKWDGTGEN PIYKQATSTFKNPMYAGK	None	None	None	None	None	None	None		

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