

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
P17301	ITA2_HUMAN	Homo sapiens	Integrin alpha-2	17.656361	NaN	NaN	34725712;38665916	MGPERTGAAPLPLLLVLALSQILNC CLAYNVGLPEAKIFSGPSSSEQGYAV QQFINPKGNWLLVSGPWSGFPENR MGDVYKCPVDLSTATCEKLNLOQTST SIPNVTEMKTNMSLGLILTRNMGTG GFLTCLGPLWAQQCGNQYTTGVC DISPDFQLSASFSPATQPCPSLIDVVV VCDESNISYPWDVAVKNFLEKFKVQGL DIGPTKTQVGLIQYANNPRVFNLNT YKTKEEMIVATSQTSQYGGDLTNTF GAIQYARKYAYSAAASGRRSATKVM VVVTDGESHDGSMMLKAVIDQCNDH NILRFGLAVLGYLNRNALDTKNLIKEI KAIASIPTRYFFNVSDAALLEKAG TLGEQIFSIEGTVOGGDNFQMEMSQ VGFSADYSSQNDILMLGAVGAFGWS GTIVQKTSHGHLIFPKQAFDQILQDR NHSSYLGYVAIISTGESTHFVAGAP RANYTGQIVLYSVNENGNITVIQHR GDQIGSYFGSVLCSVDVDKDITDVL LVGAPMYMSDLKKEEGRVYLFITKE GILGQHOFLEGPEGIENTRFGSAIAA LSDINMDGFNDVIVGSPLENQNSGA VYIYNGHQGTIRTKYSQKILGSDGAF RSHLQYFGRSLDGYGDLNGDSITDV SIGAFQOVVQLWSQSIADVAIEASFT PEKITLVNKNAQIILKLCFSAKFRPTK QNNQVAIVYNITLDADGFSSRVTSR GLFKENNERCLQKNMVMVNAQSCP EHIIYIQEPSDVVNSLDLRVDISLENP GTSPALEAYSETAKVFSIPFKDCGE DGLCISDLVLDVROIPAAQEQPFIVS NQNKRLTFSVTLKKNKRESAYNTGIV VDFSENLFASFSLPVDGTEVTCQV AASQKSVACDVGPALKREQQVTFTI NFDNFLQNLQNASLSFQALSESQ EENKADNLVNLKIPLLYDAEIHLTRS TNINFYEISSDGNVPSIVHSFEDVGP KFIFSLKVTTGSVPVSMATVIIHIPQY TKEKNPLMYLTGVQTDKAGDISCNA DINPLKIGQTSSSVSFKSEFRHTKE LNCRTASCNVTCWLKDVHMKGEY FVNVTIRWNGTFASSTFQTVQLTA AAEINTYNPEIYVIEDNTVTIPLMIMK PDEKAEVPTGVIIGSIIAGILLLLALVA ILWKLGFKKRYEKMTKNPDEIDET TELSS	False	True	2.174	2.567	2.069	2.036	1.55	5.0	2.72