

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	extracel region
P06756	ITAV_HUMAN	Homo sapiens	Integrin alpha-V	12.63613	T876	NaN	34725712;38665916;34019948;33214551;35289036;40596516	MAFPRRRLRLGPRGLPLLLSGLLLP LCRAFNLVDVSPAESGPEGSYFGF AVDFFVPSASSRMFLLVGAPKANTT QPGIVEGGQVLKCDWSSTRRCQPIE FDATGNRDYAKDDPLEFKSHQWFG ASVRSKQDKILACAPLYHWRTMKQ EREPVGTCTFLQDGTKTVEYAPCRSQ DIDADGQGFQCGGFSIDFTKADRVL LGGPGSFYWQQLISDQVAEIVSKY DPNVYSIKYNNQLATRTAQAFDDSY LGYSVAVGDFNGDGIDDFVSGVPRA ARTLGMVYIYDGKNMSSLYNFTGEQ MAAYFGFSVAATDINGDDYADVFIG APLFMDRGSQKLEQEVGOVSVSLQ RASGDFQTTKLNQFEVFAFSGSAIAP LGDLDQDGFNDIAIAAPYGGEDKKGI VYIFNGRSTGLNAVPSQILEGWAA RSMPPSFGYSMKGATDIDKNGYPDL IVGAFVDRAILYRARPVITVNAGLE VYPSILNQDNKTCSLPGTALKVSCFN VRFCKADGKGVLPKRLNFOVELLL DKLKQKGAIRRALFLYRSRPSHKN MTISRGLMQCEEIAYLRDESEFR DKLTPITIFMEYRLDYRTAADTTGLQ PILNQFTPANISRQAHILLDCGEDNV CKPKLEVSVSDQKKIYIGDDNPLTL IVKAQNQEGEGAYEAELIVSIPLQADFI GVVRNNEALARLSCAFKTENQTRQV VCDLGNPMKAGTQLLAGLRFVHQ QSEMDTSVKFDLQIQSSNLFDKVSP VVSHKVDLAVLAAVEIRGVSSPDHVF LPINWEHKENPETEEDVGPVQHI YELRNNGPSSFSKAMLHLQWPYKY NNNTLLYLHYDIDGPMNCTSDMEI NPLRIKISSLQTEKNDTVAGQGERD HLITKRDALSEGDIHTLGCQVAQCL KIVCQVRLDRGKSAILYVKSLLWTE TFMKNENQNHYSYLSKSSASFNVIEF PYKNLPIEDITNSTLVTTNVTWGIQF APMPVPVWVHLLAVLAGLLLAVLVF VMYRMGFFKRVPRPPQEEQEREQLQ PHENGEGNSET	True	True	4.364	False	0.375	0.328	False	5.0	4.131