

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
P43406	ITAV_MOUSE	Mus musculus	Integrin alpha-V	37.975063	T76;S617	NaN	33300544;40885482	MAAPGRLLLRPRPGGLLLLPGLLLP LADAFNLDVESPAEYAGPEGSYFGF AVDFFPEPSTSSRMFLLVGAPKANTT QPGIVEGGQVLKCECSSRRRCQPIEF DSTGNRDYAKDDPLEFKSHQWFGA SVRSKQDKILACAPLYHWRTEMKQE REPVGTCFLQDGTKTVEYAPCRSKNI DADGGQFCQGGFSIDFTKADRVLG GPGSFYWQQLISDQVAEHSKYDPN VYSIKYNNQLATRTAQAFDDSYLGY SVAVGDFNGDGIEDFVSGVPRAART LGMVVIYDGKNMSSLHNFTGEQMA AYFGFSVAATDINGDDYADVFIGAPL FMDRGSQGLQEVGQVSVSLQRAV GDFQITKLNQFVFAVFGSAIAPLG DLDDQDGFNDIAIAAPYGGEDKKGLV YIFNGRSTGLNSVPSQILEGQWAAQ SMPPSFGYSMKGATDVRNGYPDL VVGAFGVDRAVLYRARPVVTNAGL EVYPSILNQDNKICPLPGTALKVSCF NVRFLKADGKGTLPKRLHFQVELL LDKLGKGAIRRALFLHNRSVPVHSK TMTVFRGGQMCEELVAYLRDESE FRDKLTPITIFMEYRLDQRTAADATG LQPLNQFTPANVSRQAHILLDCGED NVCKPKLEVSVNSDQKKIYIGDDNP LTLTVKAQNOGEGAYEELIVSIPPQ ADFIGVRRNNEALARLSCAFKTENQ TRQVVCDLGNPMKAGTQLLAGLRFS VHQSEMDSVVKFDLKIQSSNSFDN VSPVVSYSYKDLAVLAAVEIRGVSSPD HIFLPIPNWEYKENPETEEDVGPVQ HIYELRNNGPSSFSKAILNLQWPYKY NNNTLLYLHYDIDGPMNCTADTEI NPLRIKTPEKNDTAAAGQGERNHLI TKRDLTREGDVHTLGCGLAKCLQIT CQVGRDRGKSAILYVKSLLWTETF MNKENQNHYSYLSKSSASFNIIEFPY KNLPIEDLFNSTLVTTNITWGIQFAP MPVPVWVILAVLAGLLLLAVLVFVM YRMGFFKVRPPQEEQEREQLQPH ENEGNSSET	False	True	3.329	2.565	2.225	2.021	1.652	5.0	2.77

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