	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	extra regio
0	P56199	ITA1_HUMAN	Homo sapiens	Integrin alpha-1	20.5777	S425;T426;S1121	NaN	35289036;37340703;38665916	MAPRPRARPGVAVACCWLLTVVLRC CVSFNVDVKNSMTFSGPVEDMFGY TVQQYENEEGKWVLIGSPLVGQPKN RTGDVYKCPVGRGESLPCVKLDLPV NTSIPNVTEVKENMTFGSTLVTNPN GGFLACGPLYAYRCGHLHYTTGICS DVSPTFQVVNSIAPVQECSTQLDIVI VLDGSNSIYPWDSVTAFLNDLLERM DIGPKQTQVGIVQYGENVTHEFNLN KYSSTEEVLVAAKKIVQRGGRQTMT ALGIDTARKEAFTEARGARRGVKKV MVIVTDGESHDNHRLKKVIQDCEDE NIQRFSIAILGSYNRGNLSTEKFVEEI KSIASEPTEKHFFNVSDELALVTIVK TLGERIFALEATADQSAASFEMEMS QTGFSAHYSQDWVMLGAVGAYDW NGTVVMQKASQIIIPRNTTFNVESTK KNEPLASYLGYTVNSATASSGDVLYI AGQPRYNHTGQVIIYRMEDGNIKILQ TLSGEQIGSYFGSILTTTDIDKDSNTD ILLVGAPMYMGTEKEEQGKVYVYAL NQTRFEYQMSLEPIKQTCCSSRQHN SCTTENKNEPCGARFGTAIAAVKDL NLDGFNDIVIGAPLEDDHGGAVYIYH GSGKTIRKEYAQRIPSGGDGKTLKFF GQSIHGEMDLNGDGLTDVTIGGLGG AALFWSRDVAVVKVTMNFEPNKVNI QKKNCHMEGKETVCINATVCFDVK LKSKEDTIYEADLQYRVTLDSLRQIS RSFFSGTQERKVQRNITVRKSECTK HSFYMLDKHDFQDSVRITLDFNLTD PENGPVLDDSLPNSVHEYIPFAKDC GNKEKCISDLSLHVATTEKDLLIVRS QNDKFNVSLTVKNTKDSAYNTRTIV HYSPNLVFSGIEAIQKDSCESNHNIT CKVGYPFLRRGEMVTFKILFQFNTS YLMENVTIYLSATSDSEEPPETLSDN VVNISIPVKYEVGLQFYSSASEYHISI AANETVPEVINSTEDIGNEINIFYLIR KSGSFPMPELKLSISFPNMTSNGYP VLYPTGLSSSENANCRPHIFEDPFSI NSGKKMTTSTDHLKRGTILDCNTCK FATITCNLTSSDISQVNVSLILWKPTF IKSYFSSLNLTIRGELRSENASLVLSS SNQKRELAIQISKDGLPGRVPLWVIL LSAFAGLLLLMLLILALWKIGFFKRP LKKKMEK	False	True	1.766	2.276	2.001	1.924	1.306	5.0	4.616