

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P11688	ITA5_MOUSE	Mus musculus	Integrin alpha-5	9.761602	NaN	S130	33300544	MGSWTPRSPRSPLHAVLLRWGPRR LPPLLPLLLLLWPPPLQVGGFNLD APAVLSGPPGSLFGFSVEFYRPG VSVLVGAPKANTSQPGVLQGGAVYV CPWGTSPIQCTTIQFDSKGSRIESS LYSAKGEEPVEYKSLQWFGATVRAH GSSILACAPLYSWRTEKDPQNDPVG TCYLSTENFTRILEYAPCRSDFGSAA GQGYCQGGFSAEFTKTGRVVLGGPG SYFWQGGILSATQEQISESYYPEYLI NPVQGGQLQTRQASSVYDDSYLGYSV AVGEFSGDDTEDFVAGVPKGNLTYG YVTVLNGSDIHSLYNVSGEQMASYF GYAVAATDTNGDGLDDLLVGAPLLM ERTADGRPQEVGRVYIYLQRPAGIDP TPTLTLTGQDEFSTRFGSSLTPLGDL QDGYNDVAIGAPFGGEAQGGVVFIF PGGPGGLSTKPSQVLQPLWAAGRTP DFFGSALRGGRLDGNYPDLIVGS FGVDKALVYRGRPIISASASLTIFPSM FNPEERSCSLEGNPVSCINLSFCLN ASGKHVPNSIGFEVELQLDWQKQK GGVRRALFLTQATLTQTLLIQNGA REDCREMKIYLRNESEFRDKLSPIHI ALNFSLDPKAPMDSHGLRPVLHYQS KSRIEDKAQILLDCGEDNICVPDLQL DVYGEKKHVYLGDKNALNLTFAHQ NLGEGGAYEAE LRVTAPLEAEYSGL VRHPGNFSSLSCDYFAVNQSRQLVC DLGNPMKAGTSLWGGLRFTVPHLQ DTKKTIQDFQILSKNLNNSQSNVV SFPLSVEAQAQVSLNGVSKPEAVIFP VSDWNPQDQPQKEEDLGPVHHVY ELINQGPSSISQGVLELSCPQALEGQ QLLYVTKVTGLSNCTSNYTPNSQGL ELDPETSPHHLQKREAPGRSSTASG TQVLKCPKCFRLRCEFGPLHRQE SRSLQLHFRVWAKTFLQREYQPFSL QCEAVYEALKMPYQILPRQLPQKKL QVATAVQWTKAEGSNGVPLWIIILAI LFGLLLLGLLIYVLYKLGFFKRSLPYG TAMEKAQLKPPATSDA