

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P11688	ITA5_MOUSE	Mus musculus	Integrin alpha-5	9.108089	NaN	S130	33300544	MGSWTPRSPRSPLHAVLLRWGPRR LPPLLPLLLLLWPPPLQVGGFNLD APAVLSGPPGSLFGFSVEFYRPGRDG VSVLVGAPKANTSQPGVLQGGAVYV CPWGTSPIQCTTIQFDSKGSRIESS LYSAKGEEPVEYKSLQWFGATVRAH GSSILACAPLYSWRTEKDPQNDPVG TCYLSTENFTRILEYAPCRSDFGSAA GQGYCQGGFSAEFTKTGRVVLGGPG SYFWQGGILSATQEQISESYYPEYLI NPVQGGQLQTRQASSVYDDSYLGYSV AVGEFSGDDTEDFVAGVPKGNLTYG YVTVLNGSDIHSLYNVSGEQMASYF GYAVAATDTNGDGLDDLLVGAPLLM ERTADGRPQEVGRVYIYLQRPAGIDP TPTLTLTGQDEFSTRFGSSLTPLGDLD QDGYNDVAIGAPFGGEAQGGVVFIF PGGPGGLSTKPSQVLQPLWAAGRTP DFFGSALRGGRLDGNYPDLIVGS FGVDKALVYRGRPIISASASLTIFPSM FNPEERSCSLEGNPVSCINLSFCLN ASGKHVPNSIGFEVELQLDWQKQK GGVRRALFLTQATLTQTLLIQNGA REDCREMKIYLRNESEFRDKLSPIHI ALNFSLDPKAPMDSHGLRPVLHYQS KSRIEDKAQILLDCGEDNICVPDLQL DVYGEKKHVYLGDKNALNLTFHAQ NLGEGGAYEAE LRVTAPLEAEYSGL VRHPGNFSSLSCDYFAVNQSRQLVC DLGNPMKAGTSLWGGLRFTVPHLQ DTKKTIQDFQILSKNLNNSQSNVV SFPLSVEAQAQVSLNGVSKPEAVIFP VSDWNPQDQPQKEDLGPVHHVY ELINQGPSSISQGVLELSCPQALEGQ QLLYVTKVTGLSNCTSNYTPNSQGL ELDPETSPHHLQKREAPGRSSTASG TQVLKCPKCFRLRCEFGPLHRQE SRSLQLHFRVWAKTFLQREYQPFSL QCEAVYEALKMPYQILPRQLPQKKL QVATAVQWTKAEGSNGVPLWIIILAI LFGLLLLGLLIYVLYKLGFFKRSLPYG TAMEKAQLKPPATSDA