

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P70670	NACAM_MOUSE	Mus musculus	Nascent polypeptide-associated complex subunit alpha, muscle-specific form	29.680179	S29;S128;S129;S159;T179;S180;T181;S182;S186;T189;T227;S240;S392;S403;T408;T430;S432;S437;S522;T557;T558;T559;S560;T580;S615;S708;S752;T760;T774;T798;T826;T835;T886;T972;S1243;T1485;S1541;S1613;S1645;S1691;S1695;T1728;T1746;S1749;S1750;S1764	T590;S822;S1174;S1177;T1364;S1368;S1392;T1398;S1400;S1423;S1492;S2015;S2104;T2131;T2133;S2138;S2158;S2163;S2175	30016717;37507081;35822049;36288343;26192747;36852467;37453647	MPGEATETVPATEQELPQQAETAV LPMSSALKVAAVGQPGTPPSSLGP QQSPIVTAHQPSLPLPSSVSTPFVEP FAQPITAETALPSGTAPPTPTFLPHLI GPPISPAALALASPMIGLAQK GARSS SAPLSLVALAPHSVQKSSVCPHPLT SPPSAAGAELGALTASIPPLEPKTSTS QVPSQGTNLKGTAPCPD VVRAFP SHLENPLASVQPGLMSCPQTLNNTS PVKGVPISSAL TQSRLSLNLKGPVSP PARNTAAPSILAPSTSLGCHLPLLH HSSVDSPIQPPGQSGLAVSNPTSVG HSGIAASCPPERCVVPALPSRLLAVD SGAAPSDDKGSSAVTNELCSPPGSS NVAGTSLSPKASLVPKGSNVALQPL VTQVPASQKTGLKEIPVSCIGATHHA LDNPSAISVAPATHVPPPTSSGLVSS KDPASPVTSLVVPAAHKQFPAPPASA TLGVPVSPLPATEGLKNLPISALVNV GAPVSPAQAGLPTRKDTTLQPLAPIA LKESPPSSQSASSLEVLSEDVTVTKKTT GGPAPVVRPAIAGVATTTSLRADSPP AVIRADSCVSPNTVSOPLKRSVTDPA MAPRTAKNTAPSTTSLVPLASEGCP VASSMALSPQNASVSETALALSPEIP KSVFPDPPLAEISFSNARKVDVASH MESSGSSRQGHPDASVTAKGTVVCL ADSSLDTSV SASKGSALSGASSPLYP LEVSFLPEAGLAVQGP KGSNLKLSPT PPSSKGAPVPSTGAPPSPKGAPIVPT ESSISSKQVPAEILPSPQKTPEVTASR LISAVQSPKVDPIMSDVTPTSPKKT S ATAVPKDTSATLSLKSVPAVTSLSPP KAPVAPSNEATIVPTEIPTSLKNALAA ATPKETLATSIPKVTSPSPQKTPKSVS LKGAPAMTSKKATEIAASKDVSPSQF PKEVPLLPHVPPPTSPKSPVSDTLSG ALTSPPPKGPPATLAETPTYPKKSPK PAASKKTPATPSPEGVTAVPLEIPPCS KKAPKTAAPKESSATSSSKRAPKTAV SKEIPSKGVTAVPLEISLPLKETSKSA TPGEKSASSPKRSPKTAGPKETPPGG VTAVPPEISLPPKETPQNATPNESLA ASSQKRSPKTSVPKETPPGGVTAMP LEIPSAPQKAPKTAVPKQIPTPEDAVT ILAGSPLSPKKASKTAAPKEAPATPSV GVIAVSGEISPSPKKTSKTAAPKENS ATLPPKRS PKTAAPKETPATSSSEGV T AVPSEISPSPTPASKGVVTLTPKGA PNALAESPASPKKVPKTAAPPEETSTT PSPQKIPKVAGPKEASATPPSKKTPK TAVPKETSAPSEGV TAVPLEIPPSPRK APKTAAPKETPAPSPEGATTAPVQIP PSPRKGSKKAGSKETPTTSPPEGVTA APLEIPISSKKT SKMASPKETLVT PSS KKL SQT VGP KETSLEGATAVPLEIPP SHKKAPKTVDPKQVPLTPSPKDAPTT LAESPSPKKAPKTAAPP SERVTTPV PEKPATPQKASATTASKVPVPAETQE VAVSSRETVPVPAVPPVKNPSSHKKT SKTIELKEAPATLPPSPTKSPKIPSSK KAPRTSAPKEFPASPSIKPVTTSLAQT APPSLQKAPSTTIPKENLAAPAVLPV

