

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	extr regi
A2AIV2	VIR_MOUSE	Mus musculus	Protein virilizer homolog	25.573173	T994;S1372	S133;S138;S173;T184;S222;T913;S1578;T1707	33300544;34418053	MAVDSSMELLFLDTFKHPSAEQSSH IDVVRFPCCVYINEVRVIPGVAHRS GLPDNRAYGETSPHTFQLDLFFNNV SKPSAPVFDRLGSLEYDENTSIIFRP NSKVNTDGLVLRGWYNCLTLAIYGS VDRVISHDRDSPPPPPPPPPPPQPQP TLKRNLKHADGEKEQDFNGSPPRP QPRGPRTPPGPPPPDDEDDPMSLP VSGDKEEDVPHREDFEPISPDRNS VPQEGQYSDEGEVEEVEEVEEEDDE DDVDVEEVEEDEDDEDDCHTVDSIPDD EEEEEEVEEVEEVEEVEEVEEVEEVEE SDEDGIADLERETFKYPNFDVEYTP DLASVPPMTYDPYDRELAPLLYFSCP YKTTFEIEISRMKDQGPDKENSGAV EASVKLTELLDLYQEDRGAKWVTAL EEIPSLIIGLSYLQLKNTQDQSLGQL VDWTMQALNLQVAFRQPIALNVRQ LKAGTKLVTSLAECGAPGVTELLQAG VINVLFDLLFADHVSSSLKLNFAKAL DSVISMTEGMEAFLRSTQNEKSGYQ RLELELLDQTVRVVTAGSAILQKCH FYEILSEIKRLGDHIAEKTSAVPNHS EPDQDQDQDQDQDQDQDQDQDQDQDQ MDMDLLESSIIEGEIEKLTNLEEV FHVMTAPHTMTQPPVKSFTIARIT GPPERDDPYVLFVRYLHSHHFLV TLLSIPITSAHQVGLQATKDVLFKFLA QSQKGLFFMSEYEATNLLIRALCH LYDQDEEGLQSDGADDAFALWLQ DSTQTLQCITELFSHFQRCASEETD HSDLLGLTHNLNLYLITFNPVGRSAVG HVFSLDKNLQSLITLMEYYSKEALG DSKSKSVAYNYACVLTLVVAQSSS GVQMLEQHAASLLKLCADENNAK LQELGKWLEPLKNLRFINCIPNIE YVKQIDNLMTAEVGLTTLALRVLC NVACPPPPVEGQQKDLKWNLAVIQL FSAEGMDTFIRVLQKLSILTQPWR LHVNMGTTLHRVTTISMARCTLTL KTMLTELLRGGSFEFKDMRVPSALV TLHMLLCSIPLSGRLDSDEQKIQNDI IDILLTFTQGVNEKLTISEETLANNT WSLMLKEVLSLKVPEGFFSGLLILL SELLPLPLPMQTTQVIEPHDISVALN TRKLWSMHLHVQAKLLQEIERSFSG TTCQPIQHMLRRICVQLCDLASPTAL LMRTVLDLIVEDLQSTSEKKEKQYT SQTRLLALLDALASHKACKLAILHL INGTIKGDERYAEIFQDLLALVRS DSVTRQQCVYVTSILQSLCDQDIAL ILPSPSEGPASELEQLSNLPSKELM TAICDCLLATLANSESSYNCLLTCVR TMMFLAEHDYGLFHLKSSLRKNSS ALHSLLRVVSTFSKDTGELASASLD FMRQILNADAMGCCGDDSGLMEVE GAHPPRTMSLNAALKQLLQSKES PESLFLLEKLVLEHSDKDDLESLL DNVIGLQKMLESSGEPLPLSDQDVE PVLAPESLQNLFNRRAYVLADVM DDQLKSMWTFPFAEIEDTDLDLVK VDLIELSEKCCSDFDLHSELEERSFLS EPSSPGRSKTTKGFGLGKHKHETFIT SSGKSEYIEPAKRAHVPPPRGRGRG GFGQIRPHDIFRQRKQNTSRPPSM HVDDFVAESKEVVPQDGIPPKRPL KVSQKISSRGGFSGNRGGRGAFHSQ NRFFTTPPASKGNYSRREGTRGSSWS AQNTPRGNYNESRGGQSNFNRGPL PPLRPLSSTGYRPSRDRASRGRGGL GPSWASTNSGSGSRGKRVVSGGSG RGRHVRSFTR	None	None	None	None	None	None	None		