

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A2A5K6	ZN335_MOUSE	Mus musculus	Zinc finger protein 335	25.875327	S510	S976;S1007;S1149	30059200	MEENEVESSDAAPRGPQPEEPS GLGVCTSEAVSADSSDAATVPGLTEA DDSGVGQSSDGGNHSVEEVSEISIT DPLPHGCLPDSSSVSRGPVAEMPGG PPALVHSSVLPDPSMLVSDCTASSS DLGSAIDKIIESTIGPDLIQSCITVTSG EEGGAETTQYLILQGPDDGAPMASS MSTSTLANSLAIEALADGPTSTSAC LEPPEEQDPPSSVAQPPAPVTEEL DLQSLAMMEVVVVQQFKCKMCQY RSSTKATLLRHMRERHFRPALAAAA AATGKRGRVRKWTSTKTTEEDRPE EEEEDDDIVDAGAIIDLEEDSDYNP AEDEPRGRQLRQRPTSTPRRRR PGRPRKLPRLTSDLHDGVGQPLVS SQSTQSPPELQDLEAPSSSGLRALGK VGRGLVESGVSQSDAENAAPSCODE ADAPRRRRGRPSRRFLGKKYRKY KSPKLLRPLYCRICGSRFLSHEDLR FHVNSHEAGDPQLFRCLQCSYRSRR WSSLKEHMFNVHVGSKPKCDECSY TSVYRKDVIRHAAVHSQDRKKRDP TPKLSFPVPCVGRVYPMQKRLTQH MKTHSTKPHMCDKCGKSFKKRYT FKMHLLTHIQAVANRRFKCEFEFV CEDKKALLNHQLSHVSDKPFKCSFC PYRTFREDFLLSHVAVKHTGAKPFA CEYCHFSTRHKKNLRLHVRCHRAN SFEEWGRRRPPEPPSRRRPFFSLQ IEELKQQHSTAPGPPSSPGPEAPQE PAPFQSPETPPLCPDALGGTTIYQQ GAEESTAVATQTALDLLNMSAQR LGATALQVAVVKSEGIEAELTSTGGQ PSPEDTTPRVVTLHMAESGSSVAE SQLGPSDLQIQLPSGPFGGASYSVI TAPPVEGRTSASGPPYREPPGAAQ AVVSDTLKEAGTHYMAADGTLH HIELTADGSI SFPSPDTLAPGKWP LQCGGPPRDGSEVLSPTKTHHMG SQGSSTPPAASHTLGLVVPQSPSA AASSTKFKFCKVCSEAFPSRAEMES HKRAHAGPAAFKCPDPPFSARQWP EVRAHMAQHSSLRPHQCNCQSFAS KNKDLRRHMLTHTNEKPFCHVC GQRFNRNGHLKFHIIQLHSIDGRKT GTSTARAPAQTIILNSEEETLTLHTA FOSSHGVLGTERLQALSOEHIIVAQ EQTVTNQEEATYIQTADGQTVQHL VTSNQQVYIISQDGVQHLLPQYEVV VPDGHIIQVQEGQITHIQYEQGTFEL QESQIQYVPVSPSQQLVTQAOLEAAA HSAVTAVADAAMAQAQGLFGTEAV PEQIHQLQHOGIEYDVTILSDD	None	None	None	None	None	None	None	None	