

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A2A5K6	ZN335_MOUSE	Mus musculus	Zinc finger protein 335	11.956056	S510	S976;S1007;S1149	30059200	MEENEVESSSDAAPRPGQPEEPS GLGVCTSEAVSADSSDAATVPLGLE DDSGVGQSSDGGNHSVEEVSEIS DPLPHGCLPDSSSVSRGPVAEMPG PPALVHSSVLPDPSMLVSDCTASS DLGSAIDKIIESTIGPDLIQSCITV EEGGAETTQYLILQGPDDGAPMASS MSTSTLANSLAAIEALADGPTSTS LEPPEEPQGDPSVAQPPAPVTEEL DLQSLEAMMEVVVVQQFKCKMCQY RSSTKATLLRHMRRERHFRPALAAA AATGKRGRVRKWTSTKTTEEDRPE EEEEDDDIVDAGAIIDLEEDSDYNP AEDEPRGRQLRLQRPTSTPRPRRR PGRPRKLPRLSDLHDGVGQPLVS SQSTQSPPELQDLEAPSSSGLRALG VGRGLVESGVSQSDAENAAPSCQDE ADAPRRRRGRPSRRFLGKKYRKY KSPKPLLRPYLCRICSRFLSHEDLR FHVNSHEAGDPQLFRCLQCSYRSRR WSSLKEHMFNHVGSKPYKCDECSY TSVYRKDVIRHAAVHSQDRKKRPDP TPKLSSFPCPVCGRVYPMQKRLTQH MKTHSTEKPHMCDKCGKSFKKRYT FKMHLLTHIQAVANRRFKCEFCEFV CEDKKALLNHQLSHVSDKPFKCSFC PYRTFREDFLLSHVAVKHTGAKPFA CEYCHFSTRHKKNLRLHVRCRHAN SFEWGRRHPEEPPSRRRPFFSLQ IEELKQQHSTAPGPPPLSSPGPEAP PAPFQSPETPPLLCPDALGGTTIYQ GAEESTAVATQTALDLLLLNSAQRE LGATALQVAVKSEGIEAELTSTGGQ PSPEDTTPRVVTLHMAESGSSVAAE SQLGPSDLQQIALPSGPFGGASYSVI TAPPVEGRTSASGPPYREEPPGEEAQ AVVSDTLKEAGTHYIMAADGTQLH HIELTADGSISFSPDTLAPGTKWPL LQCGGPPRDGSEVLSPKTHHMGG SQGSSTPPPAASHTLGLVVPOSPPSA AASSTKKFSCKVCSEAFPSRAEMES HKRAHAGPAAFKCPDCPFSARQWP EVRAHMAQHSSLRPHQCNQCSFAS KNKKDLRRHMLTHTNEKPFSCHVC GQRFNRNGHLKFHIQRLHSIDGRKT GTSTARAPAQTIILNSEEETLATLHTA FQSSHGVLGTERLQQALSQEHIIVAQ EQVTNQEEATYIQEITADGQTVQHL VTSDNQVQYIISQDGVQHLLPQEYVV

							VPDGHHIQVQEGQITHIQYEQGTPFL QESQIQYVPVSPSQQLVTQAQLEAAA HSAVTAVADAAMAQAQGLFGTEEAV PEQIHLQHQGIEYDVITLSDD
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