

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P55198	AF17_HUMAN	Homo sapiens	Protein AF-17	16.068151	T225;S401;S410;T411;T412;T413;S414;S415;S416;S429;T433;S438;T507;S512;T531;S535;S537;S540;S542;S545;T554;S555;S561;S563;S572;T573;S576;S577;S578;T585;S586	S258;S378;S423;T451	34019948;35132862;35254053;30397120;33214551;30059200;35289036;37340703;29351928;23301498;31492838;32574038;30379171;28411811	MKEMVGGCCVCSDERGWAENPLVY CDGHACSVAVHQACYGIVQVPTGPW FCRKCESQERAARVRCELCPHKDGA LKRTDNGGWAAHVVCALYIPEVQFAN VLTMEPIVLQYVPHDRFNKTCYCEE QGRESKAASGACMTCNRHGCRQAF HVTCAQMAGLLCEEEVLEVDNVKY CGYCKYHFSKMKTSRHSSGGGGGGG AGGGGGSMGGGGSGFISGRRSRAS PSTQQEKHPHTHHERGQKKS RKDKE RLKQKHKKRPESPPSILTPPVVPTAD KVSSASSSSHHEASTQETSESSRES KGGKSSSHSLSHKGGKLSGKGVSS FTSASSSSSSSSSSSGGPFQPAVSSL QSSPDFSAFPKLEQPEEDKYSKPTAP APSAPPSPSAPEPPKADLFEQKVVS GFGPIMRFSTTTSSSGRARAPSPGDY KSPHVTGSGASAGTHKRMPALSATP VPADETPETGLKEKKHKASKRSRHG PGRPKGSRNKEGTGGPAAPSLPSAQ LAGFTATAASPFSGSLVSSGLGGLS SRTFGPSGSLPSLSLESPLLAGIYTS NKDPISHSGGMLRAVCSTPLSSLL GPPGTSALPRLSRSPFTSTLPSASSI STTQVFLAGSTFSLPSTHIFGTPMG AVNPLLSQAESSHTEPDLEDSCFRC RGTSPQESLSSMSPISLPALFDQTA SAPCGGGQLDPAAPGTTNMEQLLEK QGDGEAGVNIVMLKALHALQKEN QRLQEQLSLTAKKERLQILNVQLSV PFPALPAALPAANGPVPGPYGLPPQA GSSDSLSTSKSPPGKSSLGLDNLST SSEDPHSGCPSRSSSSLSFHSTPPPL PLLQQSPATLPLALPGAPLPPQPQ NGLGRAPGAAGLGAMPMAEGLLGG LAGSGGLPLNGLLGGLNAAAPNPA SLSQAGGAPTLLPGCLNSLTEQQR HLLQQQEQLLQQLLQQLASPQLTPE HQTVVYQMIQIQKRELQRLQMA GGSQLPMASSLAGSSTPLLSAGTPG LLPTASAPPLLPAGALVAPSLGNNTS LMAAAAAA AVAAAGGPPVLT AQTN PFLSLSGAEGSGGPKGGTADKGAS ANQEKG

0