

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6PJG2	MDEAS_HUMAN	Homo sapiens	Mitotic deacetylase-associated SANT domain protein	27.147534	S57;T58;S59;S69;T83;S117;S132;T133;S438;S530	S461;T655;S661;T704;S709;T715;S923	29237092;35254053;35132862;34019948;31492838;23301498;28657654;32574038;29351928;38665916;33214551;30059200	MNLQAQPKAQNKRRKRLFGGQPEA PKEQPPPLQPPQQSIRVKEEQYLGH EGPGGAVSTSQPVELPPSSLALLNS VVYGPERTSAAMLSQQVASVKWPNS VMAPGRGPERGGGGVSDSSWQQQ PGQPPPHSTWNCHSLSLYSATKGS HPGVGVPTYYNHPEALKREKAGGPQ LDRYVRPMPQKVQLEVGRPQAPL NSFHAACKPPNQSLPLQPFQAFGH QVNRQVFRQGPFPNPVAAFPPQKQ QQQQPQQQQQQQAALPQMPLFE NFYSMPQQPSQQPQDFGLQAPGL GQSHLAHHSMAYPFPNPDMPNPE LRKALLQDSAPQALPQVQIPFRRS RRLSKEGILPPSALDGAGTQPGQEAT GNLFLHHWPLQPPPGSLGQPHPE ALGFPLELRESQLLPDGERLAPNGR EREAPAMGSEGMRAVSTGDCGQV LRGGVIQSTRRRRASQEANLLTLA QKAVELASLQNAKDGGSGEEKRKS LASTTKCGVEFSEPSLATKRAREDSG MVPLIIPVSVVRTVDPTEAAQAGGL DEDGKGPEQNP AEHKPSVIVTRRRS TRIPGTDAQQAEDMNVKLEGEPSV RKPQRPRPEPLIPTKAGTFIAPPVY SNITPYQSHLRSVRLADHPSERSFE LPPYTPPILSPVREGSGLYFNAIIST TIPAPPITPKSAHRTLRLTNSAEVTP PVLSVMGEATPVSI EPRINVGSRFQA EIPLMRDRALAAADPHKADLVWQP WEDLESSREKQRQVEDLLTAACSSI FPGAGTNQELALHCLHESRGDILET LNKLLKKPLRPHNHPLATYHYTGS DQWKMAERKLFNKGIAIYKDFFLV QKLIQTKTVAQCVEFYTYKKQVKIG RNGTLTFGDVDTSD EKS AQEEVEVD IKTSQKFPRVPLPRRESPSEERLEPK REVKEPRKEGEEVPEIQEKEEQEE GRERSRRAAVKATQTLQANESASD ILILRSHESNAPGSAGGQASEKPREG TGKSRRALPFSEKKTETFTSKTQN QENTFPCKKCGR