

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P34216	EDE1_YEAST	Saccharomyces cerevisiae	EH domain-containing and endocytosis protein 1	9.365641	NaN	T238;S241;S244;T245;S248;S249;T251;S265;S419;T450;T477;T487;S495;S848;S931;S950;S964;S1008;S1012;S1020;T1046;S1069;S1087;S1093;S1095;S1096;S1100;T1111;S1181;S1187;T1307;S1343	33229814	MASITFRTPSSQEAFYNQKFHQL DTEDLGVVTGEAVRPLFASSGLPGQ LLSQVWATVDIDNKGFLNLFNSAA LRMIAQINQAPNPQPSAALYESTPTQ LAFSFINQNPAPMQSISATGNTNNT DIPALSSNDIAKFSQLFDRITAKGAQT VAGDKAKDIFLKARLPNQTIGEIWAL CDRDASGVLDKSEFIMAMYLIQLCM SHHPSMNTPPAVLPTQLWDSIRLEP VVVNQPNRTTPLSANSTGVSSLTRH STISRLSTGAFSNAASDWSLSFEKKQ QFDAIFDSLKQHAGSLSSAVLVPPF LSSRLNQETLATIWDLADIHNNAEF TKLEFAIAMFLIQKKNAGVELPDVIP NELLQSPALGLYPPNPLPQQQSAPQI AIPSRASKPSLQDMPHQVSAPAVNT QPTVPQVLPQNSNNGSLNDLLALNP SFSSPSPTKAQTVVQNNTNNSFSYD NNGGQATLQQQQPQQPPPLTHSSS GLKKFTPTSNTFGQSIIEEPEEQEQL RESSDTFSAQPPVPKHASSPVKRTA STTLQVQPNFVFSMPAGAATSAAT GAAVGAAVGAAALGASAFSRSSNNA FKNQDLFADGEASAQLSNATTEMA NLSNQVNSLSKQASITNDKKSRAQ ELKRVTEMKNSIQIKLNNLRSTHDQ NVKQTEQLEAQVLQVNKENETLAQ QLAVSEANYHAAESKLNELTTDLQE SQTNAELKEQITNLSMTASLQSQ LNEKQQQVKQERSMVDVNSKQLEL NQVTVANLQKEIDGLGKISVYLTKQ KELNDYQKTVEEQHAQLQAKYQDLS NKDTDLDREKQLEERNRQIEEQEN LYHQHVSKLQEMFDDLSQRKASFE KADQELKERNIEYANNVRELSERQM NLAMGQLPEDAKDIIAKSASNTDTT KEATSRGNVHEDTVSKFVETTENS NLNVNRVKDDEEKTERTESDVFDR DVPTLGSQSDSENANTNNGTQSGN ETANPNLTETLSDRFDGDLNEYGIP RSQSLTSSVANNAPQSVRDDVELPE TLEERDTINNTANRDNTGNLSHIPG EWEATPATASTDVLSNETTEVIEDGS TTKRANSNEDGESVSSIQESPKISAQ PKAKTINEEFPPIQELHIDESDSSSS DDDEFEDTREIPSATVKTLPYNAQ PTSSLEIHTEQVIKYPAPGTSPSHNE GNSKKASTNSILPVKDEFDDEFAGL EQAAVEEDNGADSESEFENVANAG

								SMEQFETIDHKDLLDELQMNAFTG TLTSSSNPTIPKPQVQQQSTSDPAQV SNDEWDEIFAGFGNSKAETKVATP SIPQQPIPLKNDPIVDASLSKGPVNR GVATTPKSLAVEELSGMGFTEEEEAH NALEKCNWDLEAATNFLLDSA
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