

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
A0A178UGM3	A0A178UGM3_ARATH	Arabidopsis thaliana	Bromo domain-containing protein	27.144759	NaN	NaN	28154133	MSQIARRRRKGRPSKADLAARRSSS SPRESGYELRRGQRQNRVRYNFD DYFDEEEDEVEEKKRQKLLKQVL KLNQSRARADPPVKSRRVRHASDY EEDEEDDEAEVEEVSEKRVKVK RKLNRQDEEEEEEKDYDVEEEEE EEEGHADSEEDDKERKRRSASGM LHFLCYFVFRALIIMTNLISQSALVV LVVLGNQCDHSETTPILDKKSLELI LDKLGKDIYGVYAEVDPPEELPDYH DMIEHPMDFSTVRKKLANGSYSTLE ELESVLLICSNAMQYNSSDTVYYK QARTIQEMGKRKFEKARLKIKRAEK ELKTDEKVKPDSSVKKQVRQPF SRN GLEAVGSDFFSSGANLASGGASQNEP VSTQIGGHEKHSYTDVLFEGNTSLV DSLEKAEDLSSGKGLFGKCGRKL SV VEEDRRATYEDSDQQGDRSESI FTT FESEIKQFVAVGLHAEHAYGRSLAR FAATLGPVAVKIASQRIEQALPADFK FGRGWVGEYEPLTPVLLFETCTPK EPPKASVFSKRKSNAAKTNETLF KTPLPKAEQQGSRPVRDGNHAF FFP ASIGALSEGSPSFVATQVGNLKSMS QHEYRNPSQLDFVKPQNRIPQV EL NLPPPAEQTNSSGSGCVLENQSF GKS DTVASYRSDMMRNMSSDSEHY KHQMTTNGIFPGGLRNGKVS PGVN NRMFDLSTDFANQMSRTATSSQ QP MRQSQSHEEQAIMRNFNERART QHNSTYNHPKADAPPKISSPQ SARS EDSGNASVAAARAWMSIGAGG NNK QTFENASNPKSSQISAE SLYNPSRE HFHQQAFKPRDAEETQFHPQ RNGF PFQTFVHOPVHGMMNGGYQ PFQ NRPIVFPQMAAPTSDFSRFHV QSQW RGGITPQVQLKQRQENFLPP DLNI GVHSPDSPAKQSSQVRVDSQ QPDLA LQL	None	None	None	None	None	None	None		

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