

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
Q8BSQ9	PB1_MOUSE	Mus musculus	Protein polybromo-1	25.685855	NaN	S10;S39;S131;T134;S178;S319;S353;S355;S371;S375;S498;S509;S636;S648;S689;S948;S987;T1289;S1405	39627609	MGSKRRRATSPSSSVSGDFDDGHH SVPTPGSRKRRRLNSLPTVDPIAVC HELYNTIRDYKDEQGRLLCELFIRAP KRRNQPDYEVVVSQPIDLMKIQOKL KMEEYDDVNLTLADFQLLFFNNAKAY YKPDSPFYKAACKLWDLYLRLTRNEF VQKGEADDEDDDEGQDNQGTLD GSSPGYLKEILEQLEAIVVATNPSG RLISELFQKLPKVOYPDYAIKEPI DLKTIAQRIQNGSYKSIHAMAKDIDL LAKNAKTYNEPGSQVFKDANSIKKIF YMKKAIEIHHHEMTKSSLRIRTASNL AAARLTGPSHNKSSLGEERNPTSKY YRNKRAVQGGRLSAITMALQYGSSES EEDAALAAARYEEGESEAESITSFMD VSNPFHQLYDTRSCRNHQGLIAE PFFHLPKSKKYPDYYQIKMPSISLQQ IRTKLNQYETLDHLECDLNLMPFE NAKRYNVPNSAIYKRVLKLQOVVMA KKKELARRDDIEDGSMISSATSDT GSAKRKSKKIRKQRMKILFNVVLE AREPGSGRRLCDLFMVKPSKDYDP YYKIILEPMDLKIIEHNIRNDKYAGEE GMMEDMKLMFRNARHYNEEGSQV YNDAHILEKLLKDKRRELGPLPDDD DMASPKLKLRSKSGVSPKSKYMTM MQQKLEVEYEAVKNYTDKRGRRLS AIFLRLPSRSELPDYLTIKKPMDE KIRSHMMANKYQDIDSMVEDFVMM FNNACTYNEPESLIYKDALVLHKVLL ETRRDLEGDDESHVNVNLLIQELIH NLFVSVMSHQDDEGRCSYDSLAEIP AVDPNSPNKPLTFDIIRKNVESNRY RRLDLFQEHMFEVLERARRMNRD SEIYEDAVELQQFIRIRDELCKNGEI LLSPALSYTTKHLHNDVEKEKKEKL PKEIEEDKLRKREKREAEKSEDSG TTGLSGLHRTYSQDCSFKNSMYHVG DYVYVEPAEANLQPHIVCIERLWEDS AGEKWLYGCWFYRPNETFHLATRK FLEKEVFKSDYYNKVPVSKILGKCVV MFKVKEYFKLCPENFRDEDVFCESR YSAKTKSFKKIKLWTMPISSVRFVPR DVPLPVVRVASVFANADKGDDEKNT DNSDDNRAEDNFNLEKEKEDVPVE MSNGEPGCHYFEQLRYNDMWLKV GDCVFIKSHGLVPRVGRIEKVVWR DGAAYFYGPFIHPEETEHEPTKMFY KKEVFLSNLEETCPMSCILGKCAVLS FKDFLSCRPEIPEPENDILLCESRYNE SDKQMKKFKGLKRFSLSAKVVDDEI YYFRKPIIPQKPEPSLLEKKIQLLEAK FAELEGGDDDIEMGEEDSEVIEAP SLPQLQTPLANELDLMPTPPQSTPK SAKGSAKKESKRKINMSGYILFSSE MRAVIKAQHPDYSGELSLRVLGTWV RNLETAKAEYEGMMGGYPPGLPPL QGPVDGLVSMGSMQPLHPGGPPH HLPVGPVGLPGIPVGMNQGAVPM VGTAPGGSPYQQVGVLPVGGQA PPPYPCPHPAGPPVIQPTTMMFVAP PPKTORLLHSEAYLKYIEGLSAESNS ISKWDQTLAARRRDVHLSKEQESRL PSHWLKSKGAHTTMADALWRLRDL MLRDTLNRQAYNLENV	True	False	2.649	4.802	1.609	0.998	False	1.354	1.471