

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	extra
Q8CHH5	BICRL_MOUSE	Mus musculus	BRD4-interacting chromatin-remodeling complex-associated protein-like	37.097486	S559	S621;S976	37507081;34887587;40997131;36852467	MDDDDSDCLLDLIGDPQALNYFLHG PSSKSGSDVDTNAGYSAANSNSIFA NSTNADPKSALKGVSDQLGEGPSDG LPLASSLQFLEDELESSPLPDLSEDQ PFDILQKSLQEANITEQTLAEAYLD ASIGSSQQAQALHPSSASFTQAS NVSNYSGQTLQPIGVTHVPGASFA SNTVGVQHGFMOHVGISVPSOHL NSSQISGSGQIQIGSFCNQPSMMTI NNLDGSGIILKGGGQAPSNVSGGL LVHRQTPNGNSLFGNSTSSPVAQPV TVPFNSTNFQASLPVHNIIQRGLAP NSNKVPINIQPKPVQMGQSSAYNVN NLGIQQHHAQQGISFAPTSPOGVS VGPMSVNVNQNRKPVTSQAVS GTGGSIVHSPMGQPHTPQSQFLIPT SLSVSSNSVHHVQAINQQLLQTPS QLISGVASEHVMNLRNNSNMLRT NQPYSGQMLNNQNTAVQLVSGQTF ATSGSPVIVNHASPOIVGGQMPQQ ASPTVLHLSPGQSSVSGRPFATM PAVSGMAGPARFPVSSASTAHPTL GPTVQSGAPGSNFTGDQLTQANRTP APVSVSHRLPVSASKSPSTLSNTPGT QQQFFCQAQKKCLNQTSPIPTSKTT DGLRPSQIPGLLSTALPGQDSGSKIM PATLGATQAQPESSVSGSPSQAVQ VDSHPGQKRPAKQLTKGAFILQQL QRDQAHAVTPDKSQFRSLNDTVQRL LSYHVCQGSMPTEEDLRQVDNEFE EVATQLLKRTOAMLNKYRFLLEDA MRINPSAEMVMIDRMFNQEERASL SRDKRLALVDEGFQADFCCSFKLD EAVPETPLDRSDOHRSKTSSLHQVP RAQSRDRAKPGMAEATNHDQFHLV PNHIVVSAEGNISKKSEGHRTLKFD RGVLGQYRGPPEDKGRRDPKAVSR CSPGPEGHRKSLPRPDHGSESKLPG VLASSHMEMPCLDSFQDKALRNPS KNEVLHTDIMKSGSEPQDLQTKS LEKTFKNILELKNRPPSDPTASGA ADLDFPSFPMASQENCLEKFIPDH SEGVVETDSILEAAVNSILEC	True	False	1.547	5.0	False	False	0.188	False	0.188