

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8CHH5	BICRL_MOUSE	Mus musculus	BRD4-interacting chromatin-remodeling complex-associated protein-like	15.999574	S559	S621;S976	34887587;36852467;37507081	MDDDDDSCLLDLIGDPQALNYFLHG PSSKSGSDDVTNAGYSAANSNSIFA NSTNADPKSALKGVSDQLGEGPSDG LPLASSLQFLEDELESSPLPDLSEDQ PFDILQKSLQEANITEQTLAEEAYLD ASIGSSQQFAQAQLHPSSASFTQAS NVSNYSGQTLQPIGVTHVPVGASFA SNTVGVQHGFMQHVGISVPSQHLP NSSQISGSGQIQLIGSFGNQPSMMTI NNLDGSQIILKSGSQQAPSNVSGGL LVHRQTPNGNSLFGNSTSSPVAQP TVPFNSTNFQASLPVHNIIIQRGLAP NSNKVPINIQPKPVQMGQQSAYNVN NLGIQQHHAQQGISFAPTSSPQGSV VGPHMSVNIVNQONTRKPVTSQAVS GTGGSIHSPMGQPHTPQSQFLIPT SLSVSSNSVHHVQAINQQLLTQPS QLISGQVASEHVMLNRNSSNMLRT NQPYSGQMLNNQNTAVQLVSGQTF ATSGSPVIVNHASQIVGGQMPLQQ ASPTVLHLSPGQSSVSQGRPGFATM PAVSGMAGPARFPAVSSASTAHPTL GPTVQSGAPGSNFTGDQLTQANRTP APVSVSHRLPVSASKSPSTLSNTPGT QQQFFCQAQKKCLNQTSP IPTSKTT DGLRPSQIPGLLSTALPGQDSGSKIM PATLGATQAQPESSVSGSSPSQTAVQ VDSHPGQKRPAAKQLTKGAFILQQL QRDQAHAVTPDKSQFRSLNDTVQRL LSYHVCQGSMPTEEDLRQVDNEFE EVATQLLKRTQAMLNKYRFLLEDA MRINPSAEMVMIDRMFNQEERASL SRDKRLALVDPEGFQADFCCSFKLD EAVPETPLDRSDQHRSKTSSLHQVP RAQSRDRAKPGMAEATNHDQFHLV PNHIVVSAEGNISKKSEGHSRTLKFD RGVLGQYRGPPEDKGGRRDPAKVSR CSPGPEGHRKSLPRPDHGSESKLPG VLASSHMEMPCLD SFQDKALRNSP KNEVLHTDIMKSGGEPQPDQLTKS LEKTFKNILELKNSGRPPSDPTASGA ADLDFPSFSPMASQENCLEKFIPDH SEGVVETDSILEAAVNSILEC