

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6AI39	BICRL_HUMAN	Homo sapiens	BRD4-interacting chromatin-remodeling complex-associated protein-like	18.344998	T577;S582;S590;S591;S592;S660;T663	S623;S980	31492838;29351928;34019948;29237092;35254053;30059200;33214551;35289036;32574038	MDDDDDSCLLDLIGDPQALNYFLHG PSNKSSNDDLNTAGYSAANSNSIFA NSSNADPKSSLKGVSNQLGEGPSDG LPLSSSLQFLEDELESSPLDLTEDQ PFDILQKSLQEANITEQTLAEEAYLD ASIGSSQQAQAQLHPSSASFTQAS NVSNSYGQTLQPIGVTHVPVGASFA SNTVGVQHGFMQHVGISVPSQHLS NSSQISGSGQIQLIGSFGNHPSMMTI NNLDGSQLKKGSGQQAPSNVSGGL LVHRQTPNGNSLFGNSSSPVAQPV TVPFNSTNFQTSLPVHNIIIQRGLAP NSNKVPINIQPKPIQMGQQNTYNVN NLGIQQHHVQQGISFASASSPQGSV VGPHMSVNIVNQQNTRKPVTSQAVS STGGSIVIHSPMGQPHAPQSQFLIPT SLSVSSNSVHHVQTINGQLLQTQPS QLISGQVASEHVMLNRNSSNMLRT NQPYTGPMLNQNNTAVHLVSGQTF AASGSPVIANHASPQLVGGQMPLQQ ASPTVLHLSPGQSSVSQGRPGFATM PSVTSMSGPSRFPVSSASTAHPSLG SAVQSGSSGSNFTGDQLTQPNRTPV PVSVSHRLPVSSKSTSTFSTNTPGTG TQQQFFCQAQKCLNQTSPISAPKT TDGLRQAQIPGLLSTTLPGQDSGSKV ISASLGTAQPQEQKVVGSSPGHPAV QVESHSGGQKRPAKQLTKGAFILQ QLQRDQAHTVTPDKSHFRSLSDAVQ RLLSYHVCQGSMPTEEDLRKVDNEF ETVATQLLKRTQAMLNKYRCLLED AMRINPSAEMVMIDRMFNQEERAS LSRDKRLALVDPEGFQADFCCSFKL DKAAHETQFGRSDQHGSKASSLQP PAKAQGRDRAKTGVTEPMNHQDFH LVPNHIVSAEGNISKKTECLGRALK FDKVGLVQYQSTSEEKASRREPLKA SQCSPGPEGHRKTSSRSDHGTESKL SSILADSHLEMTCNNSFQDKSLRNS PKNEVLHTDIMKGSQEPQPDQLTK SLETTFKNILELKKAGRQPQSDPTVS GSVELDFPNFSPMASQENCLEKFIP DHSEGVVETDSILEAAVNSILEC