

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q60974	NCOR1_MOUSE	Mus musculus	Nuclear receptor corepressor 1	37.675474	T1038;T1039;S1052;T1493;S1496;S1502;S1873;S1877;S1893;T1899;S1900;S1901;T2247;T2256;T2257	S172;S224;S1011;S1122;S1206;S1207;S1274;S1292;S1333;T1378;S1459;S1481;S1598;S1993;S1997;S2116;S2134;S2150;S2165;S2198;T2412;S2449;S2451	36852467;21606357;19458039;35822049;30059200;22517741	MSSSGYPPNQGAFSTEQSRYPSSHV QYTFPSARHQQEFVDPYRSSHLEV SQASQLLQQQQQLRRRPSLLSEF HPGSDRPQERRSGYEQFHGPPSPVD HDSLESKRPRLEQVSDSHFQRISAA VLPLVHTLPEGLRSSANAKKDPAFG VKHEAPSSPLSGQPCGDDQNASPSK LSKEELIQSMDRVREIAKVEQQILK LKKKQQQLEEEAAKPEPEKPVSPPP VEQKHRSIVQIYDENRKKAEAAHKE FEGLGPKVELPLYNQPSDTKVYHENE KTNQVMRKKLILFFKRRNHARKQRE QKICQRYDQLMEAWEKKVDRNIENN PRRKAKESKTREYYEKQFPEIRKQRE QQERFQRVGQRGAGLSATIARSEHE ISEIIDGLSEQENNEKQMRQLSVIPP MMFDAEQRRVKFINMNGLMEDPMM KVYKDRQFMNVWTDHEKEIFKDKFI QHPKNFGLIASYLERKSVPCVLYYY LTKKNENYKALVRRNYGKRRGRNQ QIARPSQEEKVEEKEEDKAEKTEKK EEEKKDDEEKDDKEDSKETTKEKDR TEATAEPEEREQVTPRGRKTANSQ GRGKGRVTRSMTSEAAAAANAATAAT EPPPPPLPPPPEPISTEPVTSRWTE EEMEVAKKGLVEHGRNWAAIAKMV GTKSEAQCKNFYFNYKRRHNLNLDNL LQQHKQKASRKPREERDVSCESVA STVSAQEDEDIEASNEEENPEDSEG AENSSDTEAPSPVPEAAKSEDSS ENAASRGNTEPVAELEATTPAPCA SPSSAVPTTKPAERESVEAQVTDAS AETAEPMDVDHEECGAEGSSVLDPP APTKADSVDPQVPEMNTASKGEGD AKERDLESTSEKTEARDEDVVVAEQI ERPEPQSDDDSSATCSADEGVDGEP ERQRVFPMDAKPSLLTPPGSILISSPI KPNLLDLPQLQHRAAVIPPMVSCPT CNIPIGTPVSGYALYQRHIKAMHESA LLEEQRQRQEQVDLECRSSTSPCST SKSPNREWEVLQAPHQVITNLPEG VRLPTTRPTRPPPPLIPSSKTTVASEK PSFIMGGISISQGTPTGTYLSSHQAYP QEAPKPSVGSISLGLPRQQUESTKAAP LTYIKQEEFSPRSQNSQPEGLLVRAQ HEGVVRGTAGAVQEGSITRGTASKI SVETISSLRGSITQGTALPQAGIPTE ALVKGPSVRMPIEESPEKVVREAAAS KGHVYIEGKSGHILSYDNINAREGT RSPRTAHEMSLKRSEYEAEGSIKQG MSMRESPVSAPLEGLICRALPRGSP HSDLKERTVLSGSIMQGTPRATAES FEDGLKYPKQIKRESPPIRAFEGAITK GKPYDGITTIKEMGRSHEIPRQDILT QESRKTPEVVQSTRPIIEGSISQGTPI KFDNNSGQSAIKHNKSLITGPKLP RGMLEIVPENIKVVERGKYEDVKAG EPVRRARHTSVSSGSPVLRSTLHEAP KAQLSPGLYDDSSARRTPVSYQNTIS RGSPMMNRTSDVSSKSSASHERKS TLTPTQRESIPAKSPVPGVDPIVSHSP FDPHHRSSAAGEVYRSHLPHLDP MPFHRLDPAAYLLQRQLSPTPGY

PSQYQLYAMENTRQTILNDYITSQQ
MQVNLRPDVTRGLSPREQLGLPYP
ATRGIIDLTMPPPTILVPHAGGTSTP
PMDRITYIPGTQVTFPPRPYNAASLS
PGHPHTLAAAASAERERERERERER
ERERERERERERERIAAAPADLYLRP
GSEQPGRPGSHGYVRSPPSPSVRTQE
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QLRIMPLPSGGPSISQGLPASRYNTA
ADALAALVDAAASAPQMDVSKTKES
KHEAARLENLRSRAAVSEQQOLE
QKNLEVEKRSVQCVCTSSALPSGKA
QPHASVVYSEAGKDKGPPPKSRYEE
ELRTRGKTTTTAANFIDVIITRQIASD
KDARERGSQSSDSSSLSSHRYETA
SDAIEVISPASSAPPQEKPOAYQPD
MVKANQAENESTROYEGPLHHYRS
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PRTHRLITLADHICQIITQDFARNQV
PSQASTSTFQTSALSSTPVRTKTS
SRYSPESQSQTVLHPRPGPRVSPEN
LVDKSRGSRPGKSPERSHIPSEYEP
ISPPQGPVHEKQDSMLLSQRGVD
PAEQRSDSRSPGISYLP SFFTKLES
TSPMVKSKKQEIFRKLNSGGGSDSD
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HSFADPASNLGLEDIIRKALMG SFD
DKVEDHGVM SHPVGIMPGSASTSV
VTSSEARRDEGEPSPHAGVCKPKLI
NKNSNRKSKSPIPGQSYLGTERPSSV
SSVHSEGDYHRQTPGWAWEDRPSS
TGSTQFPYNPLTIRMLSSTPPTQIAC
APSAITQAAPHQQNRIWEREPAPLLS
AQYETLSDSDD