

| UniprotKB ID | Entry name  | organism     | full name                      | oglcnacscore | oglcnac sites                                                                                                                                                                                                                                                                                                                                                                     | phosphorylation sites                                                                                                                        | PMIDS                                                                                                                                                                                                 | sequence                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
|--------------|-------------|--------------|--------------------------------|--------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| O75376       | NCOR1_HUMAN | Homo sapiens | Nuclear receptor corepressor 1 | 36.80595     | S2;S3;S4;S14;S24;T28;T32;S45;S46;S51;S54;S123;T1026;T1027;T1030;S1039;S1040;T1042;T1043;S1050;T1064;S1073;T1075;S1082;S1114;S1263;S1266;T1449;S1450;S1453;S1454;S1457;T1479;S1480;T1484;S1487;T1491;S1493;S1545;S1806;S1812;S1857;S1861;S1877;T1883;S1884;S1885;S1889;S1913;T1925;T1926;T1928;S2033;S2034;S2069;S2070;T2078;S2079;T2080;S2084;S2086;S2219;T2233;T2242;S2246;T2299 | S172;S224;S999;S1111;S1195;S1196;S1249;S1263;S1281;S1322;T1367;S1450;S1472;S1592;S1977;S1981;S2102;S2120;S2136;S2151;S2184;T2399;S2436;S2438 | 32119511;35132862;31637018;23301498;26374642;30059200;20068230;35289036;33214551;29351928;29237092;30379171;34019948;35254053;37340703;27655845;28657654;35138101;34846842;30620550;31492838;32574038 | MSSSGYPPNQGAFASTEQSRYPHSHVQYTFPNTRHQQEFAVDPYRSSHLEVSQASQLLQQQQQLRRRPSLLSEFHPGSDRPQERRTSYEPFHGPGSPVDHDSLESKRPRLEQVSDSHFQRVSAALVPLVHPLPEGLRASADAKKDPAFGGKHEAPSSPISGQPCGDDQNASPSKLSKEELIQSMRDVREIAKVEQQILKLKKKQQQLEEEAAKPEPEKPVSPPPVEQKHRSIVQIYDENRKKAEAAHKIFEGLGPKVELPLYNQPSDTKVYHENIKTNQVMRKKLILFFKRRNHARKQREQKICQRYDQLMEAWEEKVDRIENNPRRKAKESKTREYYEKQFPEIRKQREQERFQRVGQRGAGLSATIARSEHEISEIIDGLSEQENNEKQMRQLSVIPMMFDAEQRRVKFINMNGLMEDPMMKVYKDRQFMNVVTDHEKEIFKDKFIQHPKNFGLIASYLERKSVPCVLYYYLTKKKNENYKALVRRNYGKRRGRNQQIARPSQEEKVEEKEDKAEKTEKKEEEKKDEEEKDEKEDSKENTKEKDKIDGTAEEETEEREQATPRGRKTANSQGRRKGRITRSMTNEAAAAASAAAAAETEPPPPPLPPPEPISTEPVETSRWTEEMEVAKKGLVEHGRNWAAIAKMVGTKSEAQCKNFYFNYKRRHNLNDNLQQHKQKTSRKPREERDVQCESVASTVSAQEDIEDIEASNEEENPEDSEVEAVKPSSEDSPENATSRGNTEPAVELEPTTETAPSTSPSLAVPSTKPAEDES VETQVNDSSISAETAEQMDVDQQEHSAEEGSVCDPPPATKADSVDVEVRVPENHASKVEGDNTKERDLDRASEKVEPRDEDLVVAQQINAQRPEPQSDNDSSATCSADEDVDGEPERQRMFPMDSKPSLLNPTGSILVSSPLKPNPLDLPQLQHRAAVIPPMVSVCTPCNIPGTPVSGYALYQRHIKAMHESALLEEQRQROEQIDLECRSSTSPCGTSKSPNREWEVLQPAPHQVITNLPEGVRLPTRPTRPPPLIPSSKTTVASEKPSFIMGGSISQGTPTGYLTSHNQASYTQETPKPSVGSISLGLPRQGESAKSATLPYIKQEEFSPRSQNSQPEGLLVRAQHEGVV RGTAGAIQEGSITRGTPTSKISVESIPSLRGSITQGTALPQTGIPTEALVKGSISRMPIEDSSPEKGREEAASKGHVYEGKSGHILSYDNIKNAREGTRSPRTAHEISLKRYSVESVEGNIKQGMSMRES PVSAPLEGLICRALPRGSPHSDLKER TVLSGSIMQGTPRATTESFEDGLKYPKQIKRESPPIRAFEGAITKGPYDGITTIKEMGRSIEIIPRQDILTQESRKTPEVVQSTRPIIEGSISQGTPIKFDNNSGQSAIKHNKSLITGPKLSRGMPPLEIVPENIKVVERGKYEDVKAGETVRSRHTSVVSSGPSVLRSTLHEAPKAQLSPGIYDDTSARRTPVSYQNTMSRGSMMNRSTDVTISSNKSTNHERKSTLTPQRESIPAKSPVPGVDPVVSHSPFDPHHRGSTAGEVYRSHLPTHLDPAMPFHRALDPAAAAAYLFQRQLSPTPGYPS |

QYQLYAMENTRQITLNDYITSQQMQ  
VNLRPDVARGLSPREQLGLPYPAT  
RGIIDLTNMPPTILVPHPGGTSTPPM  
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ALVSTPVRTKTSNRYPESQAQSVH  
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KLNSSGGGSDMAAAQPGTEIFNLP  
AVTTSGSVSSRGHSFADPASNLGLE  
DIIRKALMGSFDDKVEDHGVVMSQP  
MGVVPGTANTSVVTSGETRREEGDP  
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TPGWAWEDRPSSTGSTQFPYNPLT  
MRMLSSTPPTPIACAPSAVNQAAPH  
QQNRIWEREPAPLLSAQYETLSDSD  
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