

| UniprotKB ID | Entry name  | organism     | full name   | oglcnacscore | oglcnac sites   | phosphorylation sites      | PMIDS   | sequence   |
|--------------|-------------|--------------|---|--------------|---|----------------------------|---|--|
| Q6P4R8       | NFRKB_HUMAN | Homo sapiens | Nuclear factor related to kappa-B-binding protein | 37.51136     | S746;S749;T750;S752;S795;S803;T841;T843;T845;S881;S1027;S1030;S1067;S1080;S1116;S1130;T1141;S1150;T1151;S1154;S1156;T1161;S1168;T1170;S1173;T1174;S1175;S1189;T1225;T1263;S1267;T1273;S1275;S1278;T1287 | S228;S298;S351;S1022;S1291 | 30059200;31637018;30379171;20068230;33214551;27655845;31492838;32119511;22661428;20305658;28657654;32574038;29351928;21740066;29237092;34019948;30620550;23301498 | <p>MDSLDHMLTDPLELGPCGDGHGTR<br/> IMEDCLLGTRVSLPEDLLEDPEIFF<br/> DVVSLSTWQEVLSDSQREHLQOFLP<br/> QFPEDSAEQNELILALFSGENFRF<br/> GNPLHIAQKLFRDGHFNPEVVKYRQ<br/> LCFKSQYKRYLNSQQQYFHRLLKQI<br/> LASRDLLEMARRSGPALPFRQKRP<br/> SPSRTPEEREWRTOQRYLKVLRV<br/> EECGDTALSSDEEDLSSWLPSSPAR<br/> SPSPAVPLRVVPTLSTTDMKTADKVE<br/> LGDSLKIMLKKHHEKRKHQPDPH<br/> DLLTGDLTLDIMTRVNAGRKGLA<br/> ALYDLAVLKKKVKKEKEKKEKIKTI<br/> KSEAEDLAEPLSSTEGVAPLSQAPSP<br/> LAIPAICEEPLDLKPCLGINEISSSF<br/> FSLLEILLLESQASLPMLEERVLDW<br/> QSSPASSLNSWFSAPNWAELVLP<br/> LQYLAGESRAVPSSFPVFEKQKQ<br/> QWKLLGQSQDNEKELALFQLWLE<br/> TKDQAFCKQENEDSSDATTVPVRV<br/> TDYVVRPSTGEEKRVFQEQERYYS<br/> QPHKAFTRMHGFESVVGVPVKGVF<br/> DKETSLNKAREHLLRSDRPAYVTIL<br/> SLVRDAAARLPNGEGTRAEICELLKD<br/> SQFLAPDVTSTQVNTVVSGALDRLH<br/> YEKDPCKYDGRKLWIYLRDRSEE<br/> EFERIHQAQAAAARKALQKPKP<br/> PSKVKSSKESSIKVLSGSPSEQSQM<br/> SLSDSSMPPTPVPVPTPTTALPAIPI<br/> SPPPVSAVNKSGPSTVSEPAKSSSGV<br/> LLVSSPTMPLHGTMLSPASSQTAPS<br/> SQAARVSHSGSAGLSQVRVVAQP<br/> SLPAVPQQSGGPAQTLQMPAGPQI<br/> RVPATATQTKVVPQTMATVPVKAQ<br/> TTAATVQRPGPGQTGLTIVTSLPATAS<br/> PVSKPATSSPGTSAPSASTAAVIQNV<br/> TGQNIKQVAITGQLGVKQPTGNSIP<br/> LTATNFRIQGDVLRLLPSSITDAK<br/> GQTVLRITPDMMATLAKSQVTTVKL<br/> TQDLFGTGGNTTGKGISATLHVTSN<br/> PVHAADSPAKASSASAPSTPTGTTV<br/> VKVTPDLKPTASSAFRLMPALGVS<br/> VADQKKGSTVASSEAKPAATIRIVQG<br/> LGMVPPKAGQTITVATHAKQASVA<br/> SGSGTVHTSAVSLPSMNAAVSKTVA<br/> VASGAASTPISISTGAPTVRQVPVSTT<br/> VVSTSQAGKLPTRITVPLSVISQPMK<br/> GKSVVTAPIKGNLGNLSGLGRNII<br/> TTMPAGTKLIAGNKPVSFLLTAQQLQ<br/> QLQQQGQATQVRIQTVPASHLQQGT<br/> ASGSSKAVSTVVVTTAPSPKQAPEQQ</p> |