

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P12270	TPR_HUMAN	Homo sapiens	Nucleoprotein TPR	47.304208	S111;T1266;T1275;S1318;S1660;T1672;S1676;T1677;S1679;S1691;S1804;S2063;S2176;S2187;T2330	S379;S522;S523;S632;S1185;T1692;S1893;S2034;S2037;S2048;S2050;S2073;T2116;T2137;S2155	30379171;33214551;27655845;31492838;32119511;20068230;30059200;25367160;33465208;28657654;29351928;31373491;29237092;21158410;34019948;3626323;27114449;22661428;32574038;28510447	MAAVLQQVLERTELNLKPKSVQNKLEKFLADQQSEIDGLKGRHEKFKVES EQQYFEIEKRLSHSQERLVNETREC QSLRLELEKLNQLKALTEKNKELE IAQDRNIAIQSQFTRTKEELEAEKRD LIRTNERLSQELEYLTEDVKRLNEKL KESNTTKGELQKLDDELQASDVSVK YREKRLEQEKELLHSQNTWLNTELK TKTDELLALGREKGNIELELKC�LE NKKEEVSRLLEEQMNGLKTSNEHLO KHVEDLLTKLKEAKEQQASMEEFK HNELNAHIKLSNLYKSAADDSEAKS NELTRAHEELHKLLKEAGEANKAIQ DHLLVEVQSKDQMEKEMLEKIGRL EKELENANDLLSATKRKGAILSEEL AAMSPTAAAVAKIVKPGMKLTELIN AYVETQDQLLEKLENKRINKYLDEI VKEVEAKAPILKROREERYERAQKAVA SLSVKLEQAMKEIQRLQEDTDKANK QSSVLERDNRRMEIQVKDLSQQIRV LLMELEEARGNHVIRDEEVSSADISS SSEVISQHLVSYRNIEELQQQNQRLL VALRELGETREREQETTSSKITELQ LKLESALTELEQLRKSQRHQMQQLVD SIVRQRDMYRILLSQTTGVAIPLHAS SLDDVSLASTPKRPSTSQTVPAPV PVIESTEAIKAAALKQQLQEIFENYKK EKAENEKIQNEQLEKLQEQVTDLRS QNTKISTQLDFASKRYEMLQDNVEG YRREITSLHERNQKLTATTQKQEQII NTMTQDLRGANEKLAVAQVRAENL KKEKEMKLESEVRLSQQRESLLAEQ RGQNLLLTNLQTIQGILERSSETETKQ RLSSQIEKLEHEISHLKKLENEVEQ RHILTRNLVDVQLLDTKRQLDTETNL HLNTKELLKNAQKEIATLKQHLSNM EVQVASQSSQRTGKGQPSNKEDVD DLVSQLRQTEEQVNDLKERLKTSTS NVEQYQAMVTSLEESLNKEKQVTEE VRKNIEVRLKESAEFQTQLEKKLME VEKEKQELQDDKRAIESMEQQLSE LKKTLSSVQNEVQEQALQRASTALS EQQARRDCQEQAKIAVEAQNKYERE LMLHAADVEALQAAKEQVSKMASV RQHLEETTQKAESQLECKASWEER ERMLKDEVSKVCRCEDLEKQNRNL LHDQIEKLSDKVVASVKEGVQGPLN VSLSEEGKSQEQILEILRFIRREKEIA ETRFVQAQVESLRYRQRVLELREL QELQDSLNAEREKVQVTAQTMAQH EELMKKTEMTNVVMTNKMREEK ERLEQDLQMQAKVRKLELDILPLQ EANAELSEKSGMLQAEKLLLEEDVK RWKARNQHLSVQKDPDTEEYRKL LSEKEVHTKRIQQLTEEIGRLKAEIA RSNASLTNNQNLIQSLKEDLNKVRT EKETIQKDLDAKIIDIQEKVKTITQVK KIGRRYKTQYEELKAQQDKVMETSA QSSGDHQEQHVSQEMQELKETLN QAETKSLESQVENLQKTLSEKET EARNLQEQTVQLQSELSRLRQDLQD RTTQEEQLRQQITEKEEKTRKAIVAA KSKIAHLAGVKDQLTKENEELQORN

GALDQQKDEL DV RITALKS QYEGRIS
RLERELREHQERHLEQRDEPQEPS
NKVPEQQRQITLKTTTPASGERGIAS
SDPPTANIKPTPVVSTPSKVTAAMA
GNKSTPRASIRPMVTPATVTNPTTTP
TATVMPTTQVESQEAMQSEGPEH
VPVFGSTSGSVRSTSPNVQPSISQPIL
TVQQQTQATAFVQPTQQSHPOIEPA
NQELSSNIVEVQSSPVERPSTSTAV
FGTVSATPSSSLPKRTRREEEDSTIE
ASDQVSDDTVEMPLPKKLKSVTPVG
TEEEVMAEESTDGEVETQVYNQDS
QDSIGEGVTQGDYTPMEDSEETSQS
LQIDLGPLQSDQQTTTSSQDGGQKGG
DDVIVIDSDDEEEDDENDGEHEDY
EEDDEDDDEDDDTGMGDEGEDS
NEGTGSADGNDGYEADDAEGGDGT
DPGTETEESMGGEENHRAADSQN
SGEGNTGAAESSFSQEVSRQQPSS
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TPLQVAAPVTVFTESTTSDASEHASQ
SVPMTTSTGTLSSTNETATGDDGD
EVFV AESEGISSEAGLEIDSQEEEE
PVQASDES DLPSTS QDPPSSSSVDTS
SSQPKPFRRVRLQTTLRQGVRGRQF
NRQRGVSHAMGGRRGINRGNIN