

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|------------------------------------------|--------------|---------------|-----------------------------------------------------------------------------------------------------------------|----------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Q6PFD9 | NUP98_MOUSE | Mus musculus | Nuclear pore complex protein Nup98-Nup96 | 23.969972 | S1033 | S524;S608;S612;S618;S623;S625;S653;T670;S673;S680;S681;S839;S888;S934;S1027;S1042;S1059;S1063;T1069;S1328;T1771 | 22645316;21606357;29187734 | MFNKSFGTTPFGGSTGGFGTTSTFGQ NTGFGTSSGAFGTSAFGSSNNTGG LFGNSQTKPGGLFGTSSFSQPATSTS TGFGFGTSTGTNSLFGTASTGTSLF SSQNNFAQNKPTGFGNFGTSTSSG GLFGTTNTTSPFGSTSGSLFGPSSF TAAPTGTTIKFNPPGTDTMVKAGVS TNISTKHQCITAMKEYESKSLEELRL EDYQANRKGPNQVGGGTTAGLFG SSPATSSATGLFSSSTNSAFSYGQN KTAFGTSTTGFGTNPGGFLGQQNQ TTSLFSKPFQATTTPTGFSFGNTS TLGQPSTNTMGLFGVTQASQPGGLF GTATNTSTGTAFTGTGLFGQPNTG FGAVGSTLFGNNKLTTFGTSTTSAPS FGTSSGGLFGNKPTLTLGTNTNTSN FGFGTNNSSGSIKPAAGTLGTGL GTGFGTALGAGQASLFGNNQPKIGG PLGTGAFGAPGFNTSTAILGFGAPQA PVALTDPNASAAQAVLQQHLSLT YSPFGDSPLFRNPMSDPKKKEERLK PTNPAAQKALTPPHYKLTTPRPA RPKALQTTGTAKSHLFDGLDDDEPS LANGAFMPKKSIIKLVKLNNSNL FSPVNHDSDELASPSEYPERGERS FLSKPVDENNOQDGEDDSLVSRYT NPIAKPIQTPESVGNKNNSSSNVE DTIVALNMRAALRNGLEGSSEETS HDESLQDDREEIENAYHIHPAGIV LTKVGYTIPSMDDLAKITNEKGE VSDFTIGRKGYSIYFEGDVNLTNLN LDDIVHIRRKEVIVVDDNQKPPVGE GLNRKAEVTLDGWVPTDKTSRCLIK SPDRLADINYEGRLEAVSRKQGAQF KEYRPETGSWVFKVSHFSKYGLQDS DEEEHPPKTTSKKLKTAPLPPAGQ ATTFQMTLNGKPAPPQSQSPEVEQ LGRVVELSDMVDITQEPVPSVLE ESVPEDQEPVSASTHIASSLGINPHV LQIMKASLLVDEEDVDAMDQRF PSKGETVQEICSPRLPISASHSSSR SIVGGLLQSKFASGTFLSPASVQEC RTPRTSSRMNIPSTSPWSVPLPLATV FTVPSPAEVQLKTVGIRROPGLVPL EKSITYGKGLLMDMALFMGRSFRV GWGPNWTLANSGEQLHGSHELEN HQVADSMYGFLLPNPVAVKSLSESP FKVHLEKLGRLQRKLEDLQLYQTP LELKLKSTVHVDELCPVLPNPGVS VIHDYADWVKDSPGDFLELPIVKHW SLTWTLCEALWGHLKELDGQLDEPS EYIQTLERRRAFSRWLSHTAAPQIEE |

