

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P10493	NID1_MOUSE	Mus musculus	Nidogen-1	22.380232	NaN	NaN	37507081	MLDASGCSWAMWTWALLQLLLLVG PGGCLNRQELFFPGPGQGDLELEAG DDVVSPLSELIGELSFYDRTDITSVYV TTNGIHAMSEPPATEYHPGTFPPSFG SVAPFLADLDTTDGLGNVYREDLS PFIIQMAAEYVQRGFPEVSFQPTSVV VVTWESVAPYGGPSSSPAEEGKRNT FQAVLASSNSSSYAIFLYPEDGLQFF TTFSKKDESQVPAVVGFSKGLVGFL WKSNGAYNIFANDRESIENLAKSSN AGHQGVVWFIEIGSPATAKGVVSADV NLDLDDDGADYEDEYDLVTSHLGL EDVATPSPSHSPRRGYDPHNVPRIL SPGYEATERPRGVPTERTRSFQLPAE RFPQHHPQVIDVDEVEETGVVFSYN TGSQQTCCANNRHQCSVHAECRDYA TGFCRCVANYTGNRQCVAEGSP QRVNGKVKGRIFVGSQVVPVFENT DLHSYVVMNHGRSYTAISTIPETVGY SLLPLAPIGGIIGWMMFAVEQDGFKN GFSITGGEFTROAEVTFLGHPGKLV KQQFSGIDEHGHLTISTELEGRVPQI PYGASVHIEPYTELYHYSSVITSSST REYTMPEPDQDGAAPSHTHIYQWR QTITFQECAHDDARPALPSTQQLSV DSVFVLYNKEERILRYALSNSIGPVR DGSPDALQNPCYIGTHGCDNSAACR PGPGTQFTCECSIGFRGDGQTCYDID ECSEQPSRCGNHAVCNNLPGTFR ECVEGYHFSDRGTCAEEDQRPINY CETGLHNCIPQRAQCIYMGSSYT CSCLPGFSGDGRACRDVDECCQHSR CHPDAFCYNTPGSFTCQCKPGYQGD GFRCMPGEVSKTRCQLEREHILGAA GGADAQRPTLQGMFVPCDEYGHY VPTQCHHSTGYCWCVDRDGRELEG SRTPPGMRPPCLSTVAPPPIHQGPVVP TAVIPLPPGTHLLFAQTGKIERLPLER NTMKKTEAKAFLHIPAKVIIGLAFDC VDKVVYWTDISEPSIGRASLHGGEPT TIIRQDLGSPEGIALDHLGRTIFWTD SQLDRIEVAKMDGTQRRVLFDTGLV NPRGIVTDPVRGNLYWTDWNRDNP KIETSHMDGTNRRILAQDNLGLPNG LTFDAFSSQLCWVDAGTHRAECLNP AQPGRRKVLEGLQYPAVTSYGKNL YYTDWKTNSVIAMDLAISKEMDTFH PHKQTRLYGITALSQCPQGHNYCSV NNGGCTHLCLPTPGSRTCRCPDNTL GVDCIERK	False	True	1.978	2.09	1.822	1.805	1.397	3.352	4.538