

UniprotKB ID	Entry name	organism	full name	oglcnaescore	oglcnaesites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extrace region
A2AJT4	PNISR_MOUSE	Mus musculus	Arginine/serine-rich protein PNISR	25.685855	NaN	S204;S211;S290;S304;S313;S321;S465;S467;T485;S726	39627609	MWDQGGQPWQQWPLNQQQWMQS FQHQQDPSQIDWAALAAQAWIAQRE ASGQQSIVEQPPGMPNGQDMSA MESGPNNHGNGFQGDNSFNRMWQP EWGMHQQPPHPPPEQPWMPAPG PMDIVPPSEDSNSQDSGEFAPDNRH IFNQNNHNFQGGPPDNFAVGPVNQF DYQHGAAGFPQGGFHPPYWQPGP PGPPAPTQNRREPPSFRDRQRSPIA LPVKQEPPIQIDAVKRRRLPAWIREGL EKMEREKQKKLEKERMEQQRSQLS KKEKATEDAEGGDGPRLPQRSKFD SDEEDEDAENLEAVSSGKVTSPSP APQEEHSEPEMTEEEKEYOMLLT KMLLTEILLDVTDEEIYYAKDAHRK ATKAPAKQLAQSSALASLTGLGGLG GYGSGDSEDRSDRGSESDTDDEE LRHRIRQKQEAQFWRKEKEQQLLQD KQIEEEKQQTERTVKEMNEFIHREQ NSLSLEASEADRDVNDKRTNPNE APSVLEPKREHKGKKEKGRSRSGSS SSGSSSSGSRSSSSSSVSSSYSSS SGSSCTSSRSSPKRRKRPSRSRSP AKARRSRSRYSRRVKVDSSRTGK LRDRRRSNRSSIERRRRNRSPSRD RRRSRSRDRRTNRSSRSRDRR KIEDPRGNLSGNSHKHKGEAKEQD RKKEKQKREKDFKSSQDRLKRRK RESERTFCRSGSISVKVIRHDSRQDS KKNATKDSKRHSQDSSGRSSSESP GSSKEKAKPKHSRSRSLKRSQRS GKKASRKHKSRSR	True	False	3.333	4.365	1.105	0.657	False	2.675	1.046