	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extra regio
0	P15105	GLNA_MOUSE	Mus musculus	Glutamine synthetase	38.701624	T142	T104;S343	37507081;36031420;37453647;33300544;36288343;39927985;22645316	MATSASSHLNKGIKQMYMSLPQGE KVQAMYIWVDGTGEGLRCKTRTLDC EPKCVEELPEWNFDGSSTFQSEGSN SDMYLHPVAMFRDPFRKDPNKLVL CEVFKYNRKPAETNLRHICKRIMDM VSNQHPWFGMEQEYTLMGTDGHP FGWPSNGFPGPQGPYYCGVGADKAY GRDIVEAHYRACLYAGVKITGTNAEV MPAQWEFQIGPCEGIRMGDHLWIA RFILHRVCEDFGVIATFDPKPIPGNW NGAGCHTNFSTKAMREENGLKCIE EAIDKLSKRHQYHIRAYDPKGGLDN ARRLTGFHETSNINDFSAGVANRGA SIRIPRTVGQEKKGYFEDRRPSANCD PYAVTEAIVRTCLLNETGDEPFQYKN	True	False	4.347	1.125	4.0	3.0	False	4.0	0.656