	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	extracellular region
0	O35874	SATT_MOUSE	Mus musculus	Neutral amino acid transporter A	24.474581	NaN	S507;S527;S530	33300544	MEKSGETNGYLDGTQAEPAAGPRTP ETAMGKSQRCASFFRRHALVLLTVS GVLVGAGMGAALRGLQLTRTQITYL AFPGEMLLRMLRMIILPLVVCSLVSG AASLDASSLGRLGGIAVAYFGLTTLS ASALAVALAFIIKPGAGAQTLQSSSL GLENSGPPPVSKETVDSFLDLLRNL FPSNLVVAAFTTSATDYTVVTHNTSS GNVTKEKIPVVTDVEGMNILGLVLFA LVLGVALKKLGPEGEDLIRFFNSFNE ATMVLVSWIMWYVPIGIMFLIGSKIV EMKDIVMLVTSLGKYIFASMLGHVI HGGIVLPLVYFAFTRKNPFTFLLGLL TPFATAFATCSSSATLPSMMKCIEEN NGVDKRISRFILPIGATVNMDGAAIF QCVAAVFIAQLNNVDLNAGQIFTILV TATASSVGAAGVPAGGVLTIAIILEAI GLPTHDLSLILAVDWIVDRTTTVVNV EGDALGAGILNHLNQKVVKKGEQEL QEVKVEAIPNSKSEEETSPLVTHQN PAGPVAIAPELESKESVL	False	True	1.643	1.77	1.882	1.512	0.732	4.497	2.001