	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	P36543	VATE1_HUMAN	Homo sapiens	V-type proton ATPase subunit E 1	22.677653	S193	T56	40596516;30379171	MALSDADVQKQIKHMMAFIEQEAN EKAEEIDAKAEEEFNIEKGRLVQTQR LKIMEYYEKKEKQIEQQKKIQMSNL MNQARLKVLRARDDLITDLLNEAKQ RLSKVVKDTTRYQVLLDGLVLQGLY QLLEPRMIVRCRKQDFPLVKAAVQK AIPMYKIATKNDVDVQIDQESYLPED IAGGVEIYNGDRKIKVSNTLESRLDLI AQQMMPEVRGALFGANANRKFLD	True	False	4.459	2.406	False	False	False	4.0	4.0