

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region	
0	P36543	VATE1_HUMAN	Homo sapiens	V-type proton ATPase subunit E 1	18.379595	S193	T56	30379171	MALSDADVQKQIKHMMAFIEQEAN EKAEEDAKAEEEFNIEKGRLVQTQR LKIMEYYEKKEKQIEQQKKIQMSNL MNQARLKVLRARDDLITDLLNEAKQ RLSKVVKDTTRYQVLLDGLVLQGLY QLLEPRMIVRCRKQDFPLVKA AVQK AIPMYKIATKNDVDVQIDQESYLPED IAGGVEIYNGDRKIKVSN TLESRLDI AQQMMPEVRGALFGANANRKF LD	True	False	4.618	3.139	1.766	1.345	1.386	4.32	4.286