	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	l
0	P22670	RFX1_HUMAN	Homo sapiens	MHC class II regulatory factor RFX1	15.960948	S85;T104;S150;S240;T244	S60;S978;S979	34725712;38665916;34019948;33214551;39302247;31492838;39531497;32119511;35138101;35132862;29351928;40596516;35254053	MATQAYTELQAAPPPSQPPQAPPQA QPQPPPPPPPAAPQPPQPPTAAATPQ PQYVTELQSPQPQAQPPGGQKQYVT ELPAVPAPSQPTGAPTPSPAPQQYIV VTVSEGAMRASETVSEASPGSTASQ TGVPTQVVQQVQGTQQRLLVQTSVQ AKPGHVSPLQLTNIQVPQQALPTQR LVVQSAAPGSKGGQVSLTVHGTQQV HSPPEQSPVQANSSSSKTAGAPTGT VPQQLQVHGVQQSVPVTQERSVVQA TPQAPKPGPVQPLTVQGLQPVHVAQ EVQQLQQVPVPHVYSSQVQYVEGGD ASYTASAIRSSTYSYPETPLYTQTAST SYYEAAGTATQVSTPATSQAVASSGS MPMYVSGSQVVASSTSTGAGASNSS GGGGSGGGGGGGGGGGGGGSGSTG GGGSGAGTYVIQGGYMLGSASQSYS HTTRASPATVQWLLDNYETAEGVSL PRSTLYCHYLLHCQEQKLEPVNAAS FGKLIRSVFMGLRTRRLGTRGNSKY HYYGLRIKASSPLLRLMEDQQHMA MRGQPFSQKQRLKPIQKMEGMTNG VAVGQQPSTGLSDISAQVQQYQQFL DASRSLPDFTELDLQGKVLPEGVGP GDIKAFQVLYREHCEAIVDVMVNLQ FTLVETLWKTFWRYNLSQPSEAPPL AVHDEAEKRLPKAILVLLSKFEPVLQ WTKHCDNVLYQGLVEILIPDVLRPIP SALTQAIRNFAKSLESWLTHAMVNI PEEMLRVKVAAAGAFAQTLRRYTSL NHLAQAARAVLQNTAQINQMLSDL NRVDFANVQEQASWVCRCEDRVVQ RLEQDFKVTLQQQNSLEQWAAWLD GVVSQVLKPYQGSAGFPKAAKLFLL KWSFYSSMVIRDLTLRSAASFGSFH LIRLLYDEYMYYLIEHRVAQAKGETPI AVMGEFANLATSLNPLDPDKDEEEE EEEESEDELPQDISLAAGGESPALGP ETLEPPAKLARTDARGLFVQALPSS	True	False	1.359	5.0	False	False	False	l