

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q80V94	AP4E1_MOUSE	Mus musculus	AP-4 complex subunit epsilon-1	25.590209	NaN	S699;S851	35822049	MSDMVERTLTALPGLFQNLQGGP AASRAPFFSRLGGLIRGVTALSSKHE EEKLIQQLSSLKATVSAPTTTTLTKTM KECMVRLIYCEMLGYDASFGYIHAIK LAQQGNLLEKRVGYLAVSLFLHESH ELLLLLVNTVVKDLQSTNLVEVCMA LTVVSQIFPREMIPAVLPLIEDKLOH SKEIIRRKAVLALYKFYLIAPNOVQHI HTKFRKALCDRDVGVMAASLHIYLR MIKENASGYKDLTESFVTLKQVVG KLPVEFSYHSVPAPWLQIQLLRILGL LGKDDERTSELMYDVLDESLRRAEL NHNVTYAILFECVHTIYSIYPKSELLE KAAKICGKFLVSPKINLKYGLKALT YVIQQDPSLALQHQITIECLDHPDPI IKRETLELLYRITNAQNVVIVQKML EYLHQSKEEHIIISLVGRIAEAEKYA PDNVVFIQTMNAVFSVGGDVMHPD ILSNFLRLLAEGFDDDETDQQLRLYA VOSYLLTLLDMENTFYPQRFLQVMS WVLGEYSYLLDKESPEEVITRLYKLL MSDSISSETKAWLFAAVTKLTPQAH SSPLVEKLIQEFTVSLNTCLRQHAFE LKHLHENTELMKSLQGAQNCEDIV ADASLSFLDGFVAEGLSQGAAPYKP HHQRQEEQLSQEKVLNFEYGLSFS SSGFTGRQSPAGISLGSDISGNSAET GLKETSSLKMEGIKLVGKEGYLPK KESGTGDKPEASHVPAEGATVENVD QATTRKDQAQGHIPSTEEKEKQLLA SSLFVGLGPENTVDLLGKADVSHK FRRKSKLVQAQDKTPSAPTAPCSAL SLGSDVAGGDEEDGLSAVDRGDGELS SELFRSESLSGPPSAEKLESVSLPVP SLFADNNMEVFNPSSSATSTVKEE TPECRHSGLVEICSNEAVSVSSYKV WRDDCLLVIWAVTSKTDSEFTDAQL EIFPVENFKIIEQPECSSPVIETERTK SFQYSVQMESPCIEGTLSGFIKYQM MDTHSVQLEFSMNLPLLDLFIPLKIS TEDFGKLWLSFANDVKQTIKISEPGV ALTSVLTQLQNLRLRVIDVIGNEGL LACKLLPSTPCVLHCRVHADAVLW FRSSSSVLSLDYLSCHCQKVMQTS	False	True	1.875	2.122	0.776	0.907	5.0	1.414	0.964

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