

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	en ret
P14859	PO2F1_HUMAN	Homo sapiens	POU domain, class 2, transcription factor 1	26.432285	T59;T226;S232;S242;T255;T259;T263;S269;T270;S442;S535;T725;S728;S730;S734;T735;T736;T737;T738;S740	T270;T276;S283;S385;S448	27655845;23301498;30379171;31492838;40914422;35138101;29237092;30397120;40596516;21740066;38665916;30059200;39534244;34019948;28510447;29351928;39531497;40307207;37340703;31637018;35289036;35132862;20305658;33214551;23580612;35254053;32119511	MNNPSETSKPSMESGDGNTGTQTNGLDFQKQPVVGGAISTAAQAQAFGLGHLHQVQLAGTSLQAAAQSLNVQSKSNEESGDSQQPSQPSQQPSVQAAIPQQTQLMLAGGQITGLTLTPAQQLLLQQAQAQAQLLAAAVQHSASQHSASAGATISASAATPMTQIPLSQPIQIAQDLQQLQQLQQQLNLNQFVLVHPTTNLQPAQFIISQTPGQQQLLQAQNLTLQLPQQSQANLLQSQPSITLTSQPA TPTRTIAATPIQTLPOSQSTPKRIDTPSLEEPSDLEEEQFAKTFKRRIKLGFQGDVGLAMGKLYGNDFSQTTISRFEALNLSFKNMCKLKPLEKWLND AENLSSDSSLSSPSALNSPGIEGLSR RRRKRTSIETNIRVALEKSFLENQKPTSEEITMIADQLNMEKEVIRVWFCNRRQKEKRINPPSSGGTSSSPIKAIFPSP TSLVATTPLSLVTSSAATTLTVSPVLP LTSAAVTNLSVTGTSDDTTSNNTATVI STAPPASSAVTSPSLSPSPASASTSE ASSASETSTTQTTSTPLSSPLGTSQV MVTASGLQTA AAAALOGAAQLPANA SLAAMAAAAGLNPSLMAPSQAAGG ALLSLNPGT LSGALSPALMSNSTLAT IQALASGGSLPITSLDAGNLVFNAGGAPNIVTAPLFLNPQNL SLLTSNPV SLVSA AAAASAGNSAPVASLHATSTSAESIQNLSLFTVASASGAASSTTTASKA Q	True	False	1.367	5.0	1.487	4.2