

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P14859	PO2F1_HUMAN	Homo sapiens	POU domain, class 2, transcription factor 1	30.655673	T226;S232;S242;T255;T259;T263;S269;T270;S442;S535;T725;S728;S730;S734;T735;T736;T737;T738;S740	T270;T276;S283;S385;S448	29237092;27655845;23580612;21740066;31637018;38665916;37340703;30379171;34019948;29351928;30397120;35132862;35254053;20305658;31492838;28510447;35289036;32119511;33214551;30059200;23301498;35138101	MNNPSETSKPSMESGDGNTGTQTNGLDFQKQPVVGGAIATAQAQAFGLHLHQVQLAGTSLQAAAQSLNVQSKSNEESGDSQQPSQPSQQPSVQAAIPQTQLMLAGGQITGLTLTPAQQLLLQQAQAQAQLLAAAVVQHSASQQHSAAGATISASAATPMTQIPLSQPIQIAQDLQQLQQLQQQNLNLQQFVLVHPTTNLQPAQFIISQTPQGGQGLLQAQNLTLQLPQQSQANLLQSQPSITLTSQPA TPTRTIAATPIQTLPQSQSTPKRIDTPSLEEPSDLEELEQFAKTFKQRRIKLGFQGDVGLAMGKLYGNDFSQTTISRFEALNLSFKNMCKLKPILLEKWLND AENLSSDSSLSSPSALNSPGIEGLSR RRKKRTSIETNIRVALEKSFLENQKP TSEEITMIADQLNMEKEVIRVWFCN RRQKEKRINPPSSGGTSSSPIKAIFPS PLSLVATTPLSLVTSSAATTLTVSPVLP LPSAAVTNLSVTGTSDTTSNNTATVI STAPPASSAVTSPSLSPSPSASASTSE ASSASETSTTQTTSTPLSSPLGTSQV MVTASGLQTA AAAALQGAQLPANA SLAAMAAAAGLNPSLMAPSQFAAGG ALLSLNPGLSGALSPALMSNSTLAT IQALASGGSLPITSLDATGNLVFANA GGAPNIVTAPLFLNPQNLSLLTSPV SLVSAAAAASAGNSAPVASLHATSTSA ESQNSLFTVASASGAASTTTTASKA Q