

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
Q6PGL7	WASC2_MOUSE	Mus musculus	WASH complex subunit 2	36.83566	S496	S157;S159;S204;S205;S209;S284;T322;S388;S533;S538;S613;S614;S723;S747;S752;S783;S798;S870;S873;S1049;S1067;S1084;S1109;S1169;S1172;S1173;S1333	40885482	MNRTSPDSEPPASEPVWERPWSV EEIRRSSQNWSLAADAGLLQFLQEF SQQTISRTHEIKKQVDGLIQTETH CRLHNVFNDFLMLSNQFIENRVYD EEVEEQVLKAEAEKAEQEKREKQK IDLIPKQVEAVNYGLQVLDSAFEQLD IKAGNSDSEEDDANERVDLILEPKDL YIDRPLPYLIGSKLFMEQEDVGLGEL SSEEGSVGSDRGSIVDSEDEKEEEE SDEDFASHSDNDQNQHHTQISDEE EDDDGDLFADSEKEGDDIEDIEESA KSKRPTFADELAARIKGDINQRKE GQTDGKPKQTVKEKKERRTPADDEE DILFPPPTLTDDEFSPPGSRGGLFSN GQGLFDDDESDLFKEAPRARPAQA PVSEELPPSPKPGKKIPAGAVSVLLG HPDVSGSTAPSLEKQKHGQOPTPG KSSHLPAPAGLFDDDDNDNDEDDN NFFMPSSSKPSKTDKVKSTAIIFDDD EGDLFKEKAEALPAASVSQTHESKT RADKTIALPSSKNLKLVSQTKQKGL FSDEEDSDFSSQSSSKPKSASLPS SQPPTSVSLFGDEDEEDSLFGSAAA KKQTSLSLQPSQEKAKPSEQPSKKT SALLFSSDEEDQWNIADSHTKLASD NKSKGELWDSGATQGEAKAVKKT NLFEDDDDDVDLFAIAKDSQKKTQ RTSLLFEDDAESGSLFGLPPTSVPS ATTKESVPKVPLLFSDEEDSEVPSG VKPEDLKVDNARVSPEVGSADVASIA QKEGLLPASDQAEAGGSDIFSSSSPL DKGAKGRTRTVLSLFDDEDEKVEDE SSTCAPQDGREKGLKTDSPKSTGV FQDEELLFSHKLQKDNPDVDFLAG TKKIRSSVPSGGSLFGDDEDDDLFSS AKTQPVVPEKKGTLKDHVPVSLKNQ DPLDSTQGSKEKSTWKTEPAQDSSG LTPFKSREPSSRIGKIQLANLAINPAAL LPTVALQIPGTPVSSSELAFPSSEPG RSHILSVPTLPGSVEAGVSFDLPAQ ADTLHSANKSRVKVRGKRRPQTRAA RRLAAQESSEAEEDVTVDRGPVAQLS SSPVLPNGHQPLLQPRMASGOTSSE TATAPPWEGGPVLSAADRSFFVKSR PQTGNEADLFDSDGDFPKSRGQSQV EGAGVMAGEPPSHSSGGRKEKSLAF PDLSEGSSTEDLFSQVVKPRAAKNRN PFPLLEDEEDLFDPRGKKNERKPD SHQDSVSKTHDIFEDDIFATEAIKPF PKKREKGRTEPNLFDNDNIDFADLT VKPKESKKKVAAKSMFDDDDDDIF SSGLQAKASKPKSQSAEAAEQRSE HKVASIFDDPLNAFGSQ	False	False	2.781	3.41	1.096	0.961	1.881	3.679	0.934