

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrio
Q9JLV1	BAG3_MOUSE	Mus musculus	BAG family molecular chaperone regulator 3	51.491371	S2;T5;S7;S14;T160;T163;S179;S189;T320;S372;S373	S138;S179;S204;S280;S281;S285;T291;S297;S380;S382;S390	22645316;40885482;36852467;36288343;25153642;40021952;28528544;28135057;30059200;30016717	MSAATQSPMMQMASGNGASDRDP LPPGWEIKIDPQTGWPFVVDHNSRT TTWNDPRVPEGPKDTASSANGPSR DGSRLLPiREGHPiYQLRPGYiPIPV LHEGSENROPHLFHAYSQPGVQRF RTEAAAATPQRSQSPLRGGMTEAAQ TDKQCGQMPATATTAAGPPTAHGP ERSQSPAASDCSSSSSASLPSSGRS SLGSHQLPRGYiPIPVIEQNITRPAA QPSFHQAQKTHYPAAQGEYQPPQP VYHKIQGDDWEPRPLRAASPFRRSPV RGASSREGSPARSGTPVHCPSPIRVH TVVDRPQPMTHREPPVVTQPENKPE SKPGPAGDLPFGHIPIQVIRREADS KPVSQSPPPAEKVEVKVSSAPIPCP SPSPAPSAVPSPPKNVAAEQKAAPSP APAEPAAPKSGEATPPKHHPGLKVE AILEKVQGLEQAVDSFEGKKTDKKYL MIEEYLTKEALLALDSVDPEGRADVR QARRDGVKRVQITILEKLEQKVIDVPG QVQVYELQPSNLEAEQPLQEIMGAV VADKDKKGPENKDPQTESQLEAKA ATPPNPSNPADSAGNLVAP	True	False	4.684	4.603	3.359

0