	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membr
0	P42227	STAT3_MOUSE	Mus musculus	Signal transducer and activator of transcription 3	67.856333	T714;T717;S719	T705;T714;S727	18519567;28280036;36852467;24788674;22645316;25153642;38437956;22517741;32848054;38191912;26866564;37058110;28135057;28528544;35822049	MAQWNQLQQLDTRYLEQLHQLYSD SFPMELRQFLAPWIESQDWAYAASK ESHATLVFHNLLGEIDQQYSRFLQE SNVLYQHNLRRIKQFLQSRYLEKPM EIARIVARCLWEESRLLQTAATAAQQ GGQANHPTAAVVTEKQQMLEQHLQ DVRKRVQDLEQKMKVVENLQDDFD FNYKTLKSQGDMQDLNGNNQSVTR QKMQQLEQMLTALDQMRRSIVSEL AGLLSAMEYVQKTLTDEELADWKRR QQIACIGGPPNICLDRLENWITSLAE SQLQTRQQIKKLEELQQKVSYKGDPI VQHRPMLEERIVELFRNLMKSAFVV ERQPCMPMHPDRPLVIKTGVQFTTK VRLLVKFPELNYQLKIKVCIDKDSGD VAALRGSRKFNILGTNTKVMNMEES NNGSLSAEFKHLTLREQRCGNGGR ANCDASLIVTEELHLITFETEVYHQG LKIDLETHSLPVVVISNICQMPNAWA SILWYNMLTNNPKNVNFFTKPPIGT WDQVAEVLSWQFSSTTKRGLSIEQL TTLAEKLLGPGVNYSGCQITWAKFC KENMAGKGFSFWVWLDNIIDLVKK YILALWNEGYIMGFISKERERAILST KPPGTFLLRFSESSKEGGVTFTWVE KDISGKTQIQSVEPYTKQQLNNMSF AEIIMGYKIMDATNILVSPLVYLYPDI PKEEAFGKYCRPESQEHPEADPGSA APYLKTKFICVTPTTCSNTIDLPMSP RTLDSLMQFGNNGEGAEPSAGGQF ESLTFDMDLTSECATSPM	None	None	None	None	None	None	None	None