

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum
P07207	NOTCH_DROME	Drosophila melanogaster	Neurogenic locus Notch protein	47.949187	T171;T210;T248;T403;T481;T519;T557;T595;T632;S670;T708;T784;T822;T900;S1015;S1053;T1091;T1129	S2447	27268051;30018219;38359610;22158438;23671640;18948267	<p>MQSQRSRRRSRAPNTWICFWINKM  HAVASLPASLPLLLLTLFAFANLNPV  RGDTALVAASCTSVGCGNGGTCVT  QLNGKTYCACDSHYVGDYCEHRNP  CNSMRCQGGTCCQVTFRNRGRPGIS  CKCPLGFDESLCEIAPNACDHVTC  LNGGTCQLKTLLEEYTCACANGYTGE  RCETKNLNCASSPCRNGATCTALAGS  SSFTCSCPPGFTGDTCSYDIEECQSN  PCKYGGTCVNTHGSYQCMCPTGYT  GKDCDTKYKPCSPSPCQNGGICRSN  GLSYECKCPKGFEGKNCSEQNYDDC  LGHLCQNGGTCIDGISDYTCRCPPN  FTGRFCQDDVDECAQRDHPVCQNG  ATCTNTHGSYSICVNVGAGLDCSN  NTDDCKQAACFYGATCIDGVGSFYC  QCTKGTGLLCHLDDACTSNPCHA  DAICDTPINGSYACSCATGYKGVDC  SEDIDECQGGSPCEHNGICVNTPGS  YRCNCSQGFTGPRCETNINECESHP  CQNEGSCLDDPGTFRVCVMPGFTG  TQCEIDIDECQSNPCLNDGTCHDKI  NGFKCSCALGFTGARCQINIDDCQS  QPCRNRGICHDSIAGYSCECPPGYT  TSCEININDCDSNPCHRGKCIDDVN  SFKCLCDPGYTGVCQKQINECESNP  CQFDGHCQDRVGSYYCQCAQGTSG  KNCENVNVECHSNPCNNGATCIDG  INSYKQCQVPGFTGOHCEKNVDECI  SSPCANNGVCIDQVNGYKCECPRGF  YDAHCLSDVDECANPCVNEGRCE  DGINEFICHCPPGYTGKRCELDIDEC  SSNPCHGGTGYDKLNAFSCQCMPT  GYTGQKCEINIDCVNTPCGNGGTC  IDKVNKYKCVCKVPFTGRDCESKMD  PCASNRCKNEAKCTPSSNFLDFSCT  CKLGYTGRYCEDEDIDECSSPCRN  GASCLNVPGSYRCLCTKGYEGRDCA  INTDDCASFPQNGGTCCLDGIGDYS  CLCVDFGDFGKHCETDINECLSQPCQ  NGATCSQYVNSYTCTCPLGFSGINC  QTNDEDCTESSCLNGGSCIDGINGY  NCSCLAGYSGANCQYKLNKCDNSNP  CLNGATCHEQNNNEYTCHCPSGFTG  KQCSEYVDWCGQSPCENGATCSQM  KHQFSCKCSAGWTGKLCVQVITISQ  DAADRKGLSLRQLCNGNTCKDYGN  SHVCYCSQGYAGSYCQKEIDECSQ  PCQNGGTCRDLIGAYECQCRQGFQ  QNCELNIDDCAPNCPQNGGTCCHDR  VMNFSKSCPPGTMGIICEINKDDCK  PGACHNNGSCIDRVGGFECVCPQG  FVGARCEGDINECLSNPCSNAGTLD  CVQLVNNYHCNCRPGHMGRHCEH  KVDFAQSPQNGGNCNIRQSGHH  CICNNGFYGKNCESLQDCDSNPC  RVGNVVADEGFYRCECPRGTLGE  HCEIDLDECSNPQAQGAACEDLL  GDYEELCPKWKGRCDIYDANYPG  WNGGSGSNDRYAADLEQQRAMC  DKRGCTEKQNGICDSDCNTYACN  FDGNDCSLGINPWANCTANECWNK  FKNGKCNEECNNAACHYDGHDCER  KLKSCDSLFDAYCQKHYDGFCDYG  CNNAECSWDGLDCENKTQSPVLAE  GAMSVVMLMNVEAFREIQAOFLRN  MSHMLRRTTVRLKDALGHDHINWK  DNVVRPEIEDTDFARKNKILYTOQV  HQTGIQIYLEIDNRKCTECFTHAVEA  AEFLAATAAKHQLRNDFQIHSVRGI  KNPGEDEDNPPANVYVITGILVII  ALAFFGMVLSQQRKRAHGVTFWPE  GFRAPAAVMSRRRRDPHQEMRNL  NKQVAMQSQGVGQPGAHWSDDDES  DMPLPKRORSDPVSGVGLGNNGGY  ASDHTMVSEYEEADQRVWSQAHL  VVDVRAIMTPPAHQDGGKHDVDAR  GPCGLTPLMIAAVRGGGLDTGEDIA  NNEDSTAQVISDLLAQGAELNATMD  KTGETSLHLAARFARADAARLLDA</p>	None	None	None	None	None	None

GADANCQDNTGRTPHLHAAVAADAM  
GVFQILLRNRATNLNARMHDGTTPL  
ILAAARLAIEGMVEDLITADADINAAD  
NSGKTALHWAAAVNNTAVNILLM  
HHANRDAQDDKDETPFLAAREGSY  
EACKALLDNFANREITDHMDRLPRD  
VASERLHHDIVRLLDEHVPRSPQML  
SMTPOAMIGSPPPGQQQPQLITQPT  
VISAGNGGNNGNASGKQSNQTA  
KQKAACKAKLIEGSPDNGLDATGSL  
RRKASSKTSAAASKKAANLNLNPG  
QLTGGVSGVPGVPPTNSAAQAAAAA  
AAAVAAMSHELEGSFVGVGMGGNL  
PSPYDTSSMYSNAMAAPLANGNPN  
TGAKQPPSYEDCIKNAQSMQSLQGN  
GLDMIKLDNYAYSMGSPFQELLNG  
QGLGMNGNGORNGVGPVLPGGL  
CGMGGLSGAGNGNSHEQGLSPPYS  
NQSPPHSVQSSLALSPHAYLGSPPS  
AKSRPSLPTSPTHIQAMRHATQKQ  
FGGSNLNSLLGGANGGGVGGGGG  
GGGGVGGPQNSPVSLGIISPTGSD  
MGIMLAPPQSSKNSAIMQTISPQQ  
QQQQQQQQQHQQQQQQQQQQ  
QQQQQLGGLEFGSAGLDLNGFCG  
SPDSFHSGQMNPPSIQSSMSGSSPS  
TNMLSPSSQHNQAFYQLTPSSQ  
HSGGHTPQHLVQTLDSYPTSPESP  
GHWSSSPRSNSDWSEGVQSPAAN  
NLYISGGHQANKGSEAIYI