

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P78347	GTF2I_HUMAN	Homo sapiens	General transcription factor II-I	6.665498	S32;T785	S19;S103;S207;S210;S214;T248;T398;S412;T503;S517;T556;T558;S668;S674;S722;S784;S823;S278;S298	34019948;25367160;28510447;30379171;33214551;32119511;23301498;34725712;35083852;33465208;35132862;35008409	MAQVAMSTLPVEDEESSES RMVVT FLMSALESMCKELAKSKAEVACIAV YETDVFVVGTERGRAFNTRKDFQK DFVKYCVVEEEKAAEMHKMKSTTQ ANRMSVDAVEIETLRKTVEDYFCFC YGKALGKSTVVPVPEKMLRDQSAV VVQGLPEGVAFKHPENYDLATLKIWI LENKAGISFIIKRPFLPKKHVGGRV MVTADRSILSPGGSCGPIKVKTEPT EDSGISLEMAAVTVKEESEDPDYYQ YNIQAGPSETDDVDEKQPLSKPLQG SHHSSEGNEGTEMEVPAEDSTQHV PSETSEDPEVEVTIEDDDYSPPSKRP KANELPQPPVPEPANAGKRKRVREFN FEKWNARITDLRKQVEELFERKYAQ AIKAKGPVTIPYPLFQSHVEDLYVEG LPEGIPFRPSTYGIPLERILLAKERI RFVIKKHELLNSTREDLQDKPASG VKEEWYARITKLRKMVDQLFCKKFA EALGSTEAKAVPYQKFEAHPNDLYV EGLPENIPFRSPSWYGIPRLEKIIQVG NRIKFVIKRPELLTHSTTEVTQPRTN TPVKEDWNVRITKLRKQVEEIFNLK FAQALGLTEAVKVPYPVFESNPEFLY VEGLPEGIPFRSPTWFGIPRLERIVR GSNKIKFVVKPELVISYLPFGMASK INTKALQSPKRPRSPGNSKVPPIEV TVEGPNNNPQTSAVRTPQTQNGS NVPFKPRGREFSFEAWNAKITDLKQ KVENLFNEKCGEALGLKQAVKVPFA LFESFPEDFYVEGLPEGVPFRPSTF GIPRLEKILRNKAKIKFIIKKPEMFET AIKESTSSKSPPRKINSSPNVNTTAS GVEDLNIIQVTIPDDDNERLSKVEKA RQLREQVNDLFSRKFGEAIGMGFPV KVPYRKITINPGCVVVDGMPPGVSFK APSYLEISSMRRILDSAEFIKFTVIRP FPGLVINNLVDQSESEGPVIQESAE PSQLEVPATEEIKETDGSSQIKQEPD PTW