

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P78347	GTF2I_HUMAN	Homo sapiens	General transcription factor II-I	9.280177	S32;T785	S19;S103;S207;S210;S214;T248;T398;S412;T503;S517;T556;T558;S668;S674;S722;S784;S823;S278;S298	25367160;35132862;35008409;35083852;34019948;23301498;28510447;34725712;38665916;32119511;3214551;33465208;30379171	MAQVAMSTLPVEDEESSES RMVVT FLMSALESMCKELAKSKAEVACIAV YETDVFVVGTERGRAFNTRKDFQK DFVKYCVVEEEKAAEMHKMKSTTQ ANRMSVDAVEIETLRKTVEDYFCFC YGKALGKSTVVPVPEKMLRDQSAV VVQGLPEGVAFKHPENYDLATLKW LENKAGISFIIKRPFLPKKHVGGRV MVTADRSILSPGGSCGPIKVKTEPT EDSGISLEMAAVTVKEESEDPDYYQ YNIQAGPSETDDVDEKQPLSKPLQG SHHSSEGNEGTEMEVPAEDSTQHV PSETSEDPEVEVTIEDDDYSPPSKRP KANELPQPPVPEPANAGKRKRVREFN FEKWNARITDLRKQVEELFERKYAQ AIKAKGPVTIPYPLFQSHVEDLYVEG LPEGIPFRRPSTYGIPRLERILLAKERI RFVIKKHELLNSTREDLQDKPASG VKEEWYARITKLRKMVDQLFCKKFA EALGSTEAKAVPYQKFEAHPNDLYV EGLPENIPFRSPSWYGIPRLEKIIQVG NRIKFVIKRELLTHSTTEVTQPRTN TPVKEDWNVRITKLRKQVEEIFNLK FAQALGLTEAVKVPYPVFESNPEFLY VEGLPEGIPFRSPTWFGIPRLERIVR GSNKIKFVVKPELVISYLPFGMASK INTKALQSPKRPRSPGSNSKVPEIEV TVEGPNNNPQTSAVRTPQTQNGS NVPFKPRGREFSFEAWNAKITDLKQ KVENLFNEKCGEALGLKQAVKVPFA LFESFPEDFYVEGLPEGVPFRRPSTF GIPRLEKILRNKAKIKFIIKKPEMFET AIKESTSSKSPPRKINSSPNVNTTAS GVEDLNIIQVTIPDDDNERLSKVEKA RQLREQVNDLFSRKFGEAIGMGFPV KVPYRKITINPGCVVVDGMPPGVFSK APSYLEISSMRRILDSAEFIKFTVIRP FPGLVINNLVDQSESEGPVIQESAE PSQLEVPATEEIKETDGSSQIKQEPD PTW