

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
Q9H7N4	SFR19_HUMAN	Homo sapiens	Splicing factor, arginine/serine-rich 19	15.723012	T357;S534;S548	S239;T335;S448;S453;S498;S500;S526;S548;S612;S614;T706;S719;S725;T732;S734;S738;S872;S874;S929;S936;S963;S965;T976;T989;S992;T1001	23301498;31373491;34019948;29351928	MEEEDSRGKTEESGEDRGDGPDP RDPTLSPSAFILRAIQAVGSSLQGD LPNDKDGSRCHGLRWRRCSRPRSE PRSQESGGTDTATVLDMATDSFLAG LVSVLDPDPTWVPSRLDLRPGESD MLELVAEVRIGDRDPIPLPVPSLLPR LRAWRTGKTVSPQSNSSRPTCARHL TLGTGDGGPAPPAPSSASSPSPSP SSSSPSPPPPPPPAPPAPPAPRFDIY DPFHPTDEAYSPPPAPEQKYDPFEPT GSNPSSSAGTPSPEEEEEEEEEEEE EEEEEEEEEGLSQSSISRISETLAGIYD DNSLSQDFPGDESPRPDAQPTQPTP APGTPPQVDSTRADGAMRRRVFVVG TEAEACREGKVSVEVVTAGGAALPP PLLPPGDSEIEEGEIVQPEEPRLALS LFRPGGRAARPTPAASATPTAQLPQ PPAPRAPEGDDFSLHAESDGEGAL QVDLGEPAAPPAADSRWGGDLDRR KILTQRRERYRQRSPSPAPAPAAAA AGPPTRKKRRERKRSGEAKAASS SSGTQPAPPASPWDSKKHRSRDR KPGSHASSARRRSRSRSRSTRR RSRSTDRRRGGSRRRSREKRRRRR RSASPPPATSSSSSRERHRGKHR DGGGSKKKKRSRSRGEKRSRSGDGS EKAPAPAPPPSGSTSCGDRDRRRR AVPPSIQDLTDHDLFAIKRTITVGRLD KSDPRGSPAPASSPKREVLVDSEGL SGEERGGKSSQKDRRRSGAASSSSS SREKGSRRKALDGGDRDRDRDRDR DRDRSSKKARPPKESAPSSGPPPKPP VSSGSGSSSSSSCSSRKVKLQSKVA VLIREGVSTTPAKDAASAGLGSIGV KFSRDRESRSPFLKPERAPTEMAK AAPGSTKPKKTKVKAKAGAKTKGT KGKTKPSKTRKKVRSGGGSGGSGGQ VSLKSKADSCSQAAGTKGAEETSW SGEERAAPVSTPPPKAAPPPALTP DSQTVDSCKTPEVSFLPEEATEEAG VRGGAEEEEEEEEEEEEEEEEEQ PATTTATSTAAAAPSTAPSAGSTAGD SGAEDGPASRVSQLPTLPPMPWNL PAGVDCTTSGVLALTALLFKMEEAN LASRAKAQELIQATNQILSHRKPPSS LGMTAPVPTSGLPPGPSSYLLPGS LPLGGCGSTPPTPTGLAATSDKREGS SSSEGRGDTDKYLKLTQERAVEE VKLAIKPYQKKDITKEEYKDILRKAV HKICHSKSGEINPVKVSNLVRAVQ RYRYFRKHGRKPGDPPGPPRPPKEP GPPDKGGPGLPLPPL